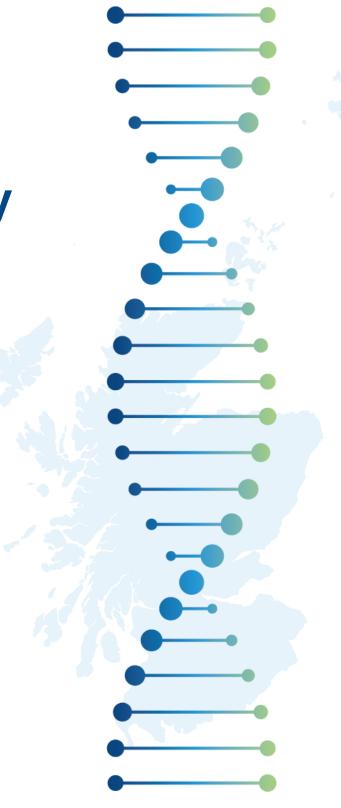


# **Scottish Strategic Network for Genomic Medicine**

# **Genomic Test Directory**

**Rare & Inherited Disease** 

November 2024



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NSD611-003.20 V5

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# **Scottish Strategic Network for Genomic Medicine**

# **CHANGE SUMMARY**

| Version<br>History | Type of change                                     | Summary of change  | Link        |
|--------------------|--|--|-------------|
| New                | Removal of text                                    | Removed clinician feedback that has been left in from a draft version  | <u>Link</u> |
| New                | Correction of removal of text                      | Correction of removal of referral criteria from Hereditary Breast Cancer.  | <u>Link</u> |
| New                | Correction   | Moved Hearing loss entry to pharmacogenomics section   | <u>Link</u> |
| New                | Addition of requesting specialty                   | Iron Regulation: Inclusion of Gastroenterology as a requesting specialty.  | Link        |
| New                | Addition to referral criteria                      | Addition of text: If there is a clinical suspicion of Congenital Central Hypoventilation Syndrome (CCHS) in association with HSCR, please discuss with clinical genetics to arrange the most appropriate testing | Link        |
| New                | Modification to existing entry-removal of an entry | Removed Medullary Thyroid Cancer as a test criteria from page 216.   | n/a         |
| New                | Addition of a gene to panel                        | Addition of FAN1 to gene targets.  | Link        |
| New                | Addition to referral criteria                      | Added text: Note: This test is currently available as pilot programme.   | <u>Link</u> |
| New                | Addition of a gene to panel                        | Addition of ASPH to gene targets.  | <u>Link</u> |
| V4                 | Addition   | Test Directory version control added.  | N/A         |
| V4                 | Addition   | SSNGM Demand Optimisation working group wrote and agreed national guidance on the application of referral criteria when assessing a requested test. This guidance has now been included.                         | LINK        |

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| V4 | Addition     | Pharmacogenomic testing for ASTHMA ß2-ADRENERGIC RECEPTOR (ADRB2) p.(Gly16Arg) GENOTYPING and AMINOGLYCOSIDE RELATED DEAFNESS MT-RNR1 M.1555A>G GENOTYPING added from cancer directory.   | <u>LINK</u> |
|----|--------------|---|-------------|
| V3 | Correction   | Inherited Cancer title included where previously omitted  | LINK        |
| V3 | Correction   | Hereditary Breast Cancer returned to directory and removed "• Combined pathology-adjusted Manchester score of ≥15 or single gene pathology-adjusted Manchester score of ≥10 or aCanRisk score of ≥10%" from the referral criteria as not applied in Scotland. | LINK        |
| V3 | Correction   | Familial Melanoma returned to directory   | LINK        |
| V3 | Correction   | CDH1 included for Aberdeen in Hereditary Diffuse Gastric Cancer Syndrome where previously omitted   | LINK        |
| V3 | Correction   | BRIP changed to BRIP1 in Hereditary Breast/Ovarian Cancer Syndrome  | <u>LINK</u> |
| V3 | Correction   | BRIP changed to BRIP1 in Hereditary Ovarian Cancer Syndrome   | LINK        |
| V2 | Modification | Sub-headings removed from table of contents as linked to same information as the heading.   |             |
| V2 | Amendment    | Referral criteria for Hereditary Breast/Ovarian Cancer Syndrome: Founder Variants Only changed to remove Italian reference and referral criteria updated to reflect current practice.   | LINK        |





# **Scottish Strategic Network** for Genomic Medicine

#### INTRODUCTION

# NHS SCOTLAND LABORATORY GENETIC SERVICES

NHS Scotland genetics services are delivered through four regional genetics centres in Aberdeen, Dundee, Edinburgh and Glasgow. Each centre offers a closely integrated laboratory and clinical service. NHS National Services Scotland commission the four genetics laboratories in Scotland to work as a formal network arrangement, to deliver an equitable, high quality genetic testing service for Scotland. All laboratories are accredited by United Kingdom Accreditation Service (UKAS) in accordance with the recognized ISO 15189:2012 standard.

Molecular genetics testing was nationally designated in 1985 and cytogenetics in 2009. Molecular pathology testing services were nationally commissioned as a single designated multi-site national specialist service from 1 April 2013.

Genetics and molecular pathology services are evolving with the workload increasing each year, as new advances increase the range of conditions which can be tested for. In molecular genetics there are a small number of 'core' tests performed in all four centres, with the majority of tests being performed in one laboratory for all of Scotland. The service undertakes testing for over 200 conditions.

# PURPOSE OF DOCUMENT

The Scottish Strategic Network for Genomic Medicine (SSNGM) Genomic Test Directory for Rare and Inherited Disease contains a list of all services currently available in Scotland.

This document will be reviewed annually.

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# NHS SCOTLAND GENETIC LABORATORY CONTACT DETAILS

#### Aberdeen (NHS Grampian)

Address: Genetics and Molecular Pathology Laboratory Services, Polwarth

Building, Foresterhill, Aberdeen AB25 2ZD Email address: <a href="mailto:gram.molgen@nhs.scot">gram.molgen@nhs.scot</a>

Website: https://www.nhsgrampian.org/service-hub/north-of-scotland-medical-

genetics

#### • Dundee (NHS Tayside)

Address: East of Scotland Regional Genetics Service, Level 6, Ninewells

Hospital, Dundee DD1 9SY Email address: <u>Tay.esrg@nhs.scot</u>

Website: https://www.nhstayside.scot.nhs.uk/OurServicesA-

Z/Genetics/PROD\_295540/index.htm

# Edinburgh (NHS Lothian)

Address: South East Scotland Genetic Service, Western General Hospital,

Crewe Road, Edinburgh, EH4 2XU

Email address: edinburgh.dna@nhslothian.scot.nhs.uk /

wgh.cytogenetics@nhslothian.scot.nhs.uk Phone: 0131 537 1116 / 0131 537 1940

Website: https://services.nhslothian.scot/clinicalgeneticsservice/GeneticLabora

toryServices/Pages/default.aspx

#### Glasgow (NHS Greater Glasgow & Clyde)

Address: West of Scotland Centre for Genomic Medicine, Laboratory Genetics, Level 2B Laboratory Medicine & FM Building, Queen Elizabeth

University Hospital, Glasgow G51 4TF

Email address: Genetic.Laboratories@ggc.scot.nhs.uk

Website:

<u>Laboratory Genetics - NHSGGC</u>





# **TEST REQUESTING**

Testing will be delivered either locally or nationally according to the test directory. However, samples should be taken and sent to your **LOCAL** genetics laboratory with the appropriate completed genetics referral form (or proforma if required). For local sample acceptance policies and referral forms, please see the local laboratory website.

Services are provided for the clinical indications listed when referred from the appropriate specialties.

# SAMPLE REQUIREMENTS

For most rare and inherited disease genomic tests with the exception of karyotyping, an EDTA blood sample is required. For karyotyping tests, a lithium heparin blood sample is required.

Other sample types may be required for some services including:

- Urine samples may be required for some mitochondrial tests.
- Appropriate fresh tissue samples from post mortems for various tests.

For specific sample requirements, please see the local laboratory website.





# TESTING METHODOLOGY

Different methods are utilised depending on the scope of testing. These methods include techniques to detect a single variant up to genome wide screens. The different methods include:

- PCR
- Sanger sequencing
- Next Generation Sequencing (NGS) panels vary in size from a small to large number of genes
- Fragment analysis
- Multiplex Ligation Probe Amplification (MLPA)
- Karyotype
- Microarray
- Chromosome breakage

# SCOPE AND RANGE OF TEST

The scope and range of testing refers to the extent of testing and the types of variant that will be detected.

The scope of testing includes:

- Targeted testing testing of specific region(s)
- Whole gene screen sequence of coding region of relevant gene(s)
- Whole gene screen and copy number sequence of coding region and assessment of exon level copy number
- Genome wide detection of large scale rearrangements

The types of variants detected includes:

- Small sequence variants
  - Single nucleotide variants (SNVs)
  - Insertions / deletions (indels)
- Copy number variants (CNVs)
  - o Exon level
  - o genome wide level
- Repeat expansions
- Aneuploidy
- Genome wide rearrangements

The targets tested refer to the genes / regions tested for the particular clinical indication.

Testing is provided for the affected individual only in most cases. If parental samples are required for Trio analysis, this will be stated in the test information.





# **REPORTING TIMES**

Reporting times are listed based on calendar days. These range from 3 to 112 days depending on urgency and complexity of testing. Where more urgent testing is required than what is stated for treatment decisions, please contact the laboratory providing testing to discuss.

# CLINICAL CONSENT AND COUNSELLING IMPLICATIONS

It is the referring clinician's responsibility to ensure that testing and /or storage of genetic material is discussed with the patient and that a summary of clinical consent is included in the patient's health record. Further information regarding consent can be found at https://www.bsgm.org.uk/healthcare-professionals/confidentiality-and-genetic-information. The patient should discuss and understand the following:

#### Family implications

The results of my test may have implications for other members of my family. I acknowledge that my results may sometimes be used to inform the appropriate healthcare of others. This could be done in discussion with me, or in such a way that I am not personally identified in this process.

#### Uncertainty

The results of my test may reveal genetic variation whose significance is not yet known. Deciding whether such variation is significant may require sharing of information about me including (inter)national comparisons with variation in others. I acknowledge that interpretation of my results may change over time as such evidence is gathered.

#### Unexpected information

The results of some tests may reveal a chance of a disease in the future, and nothing to do with why I am having this test. This may be found by chance, while focusing on the reason for my test, and I may then need further tests to understand what this means for me. If these additional findings are to be looked for, I will be given more information about this.

#### DNA storage

Normal laboratory practice is to store the DNA extracted from my sample even after the current testing is complete. My sample might be used as a 'quality control' for other testing, for example, that of family members.

#### Data storage

Data from my test will be stored to allow for possible future interpretations.

#### Health records

Results from my test and my test report will be part of my patient health record.

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# REFERRAL CRITERIA

The referral criteria outlined in this directory have been developed using national and international guidance appropriate to each individual test that has been commissioned nationally. They have been reviewed by specialists in each test field where appropriate, including NHS clinicians and scientists working in Scotland. All four genomic testing laboratories have been consulted and a unified referral criteria agreed based on the testing commissioned nationally.

The referral criteria should be used by referrers to inform appropriate test requesting and will be implemented on a national basis by all four centres. Where a request for a test does not meet the referral criteria, the appropriate testing centre laboratory will contact the referrer to advise on the outcome and provide guidance about what to do next.

On rare occasions it might be appropriate to offer testing to a patient who do not meet the referral criteria for a particular test. These referrals will be considered on a case by cases basis and must be discussed with a senior laboratory scientist/ manager before testing can be approved.

Referrers should consider the referral criteria and if further advice is required, please contact the testing centre via the <u>contact details in this directory</u>.





# **CARDIOLOGY**

#### **ANDERSEN-TAWIL SYNDROME**

# Available testing

| Centre                | Method                                   | Scope and range of test |  |              | Targets      | TAT |
|-----------------------|--|-------------------------|--|--------------|--------------|-----|
| Aberdeen              | Sanger                                   | Whole gene screen       |  | SNVs, indels | KCNJ2, KCNJ5 | 56  |
| Family me             | Family member testing as indicated above |                         |  |              |              | 14  |
| Proforma required? YE |  | YES                     | Cardiac Arrhythmia Proforma (see centre website) |              |              |     |

#### Referral criteria

- Ventricular arrhythmia and /or prolonged QTc
- Periodic paralysis
- Distinctive facial and skeletal features

- Cardiologist with expertise in ICC
- Clinical Genetics
- Neurology
- Paediatric Neurology





#### **ARRHYTHMIA PANEL**

# Available testing

| Centre             | Method                           | Sco | Scope and range of test |                    | Targets   | TAT |
|--------------------|----------------------------------|-----|-------------------------|--------------------|---|-----|
| Aberdeen           | NGS                              |     | e gene<br>een           | SNVs, indels       | KCNQ1, KCNH2, KCNE1, KCNE2, KCNJ2, SCN5A, RYR2, DSC2, DSG2, DSP, PKP2, ABCC9, AKAP9, ANK2, CACNA1C, CACNA2D1, CACNB2, CALM1, CALM2, CASQ2, CAV3, DES, DPP6, GJA1, GJA5, GPD1L, HCN4, JUP, KCNA5, KCND3, KCNE5, KCNE3, KCNJ5, KCNJ8, LMNA, NOS1AP, NPPA, PLN, RANGRF, SCN10A, SCN1B, SCN2B, SCN3B, SCN4B, SLMAP, SNTA1, TGFB3, TMEM43, TRDN, TRPM4 | 112 |
| Family mer testing | Family member as indicated above |     |                         |                    | 14  |     |
| Proforma re        | equired?                         | YES | Cardiac A               | Arrhythmia Profori | ma (see centre website)   |     |

#### Referral criteria

- Out of Hospital Cardiac Arrest with no known cause
- Sudden cardiac death with negative post mortem

- Cardiologist with expertise in ICC
- Clinical Genetics
- · Pathology in discussion with Clinical Genetics





#### ARRHYTHMOGENIC CARDIOMYOPATHY

#### Available testing

| Centre                    | Method       | Scope and range of test |           |                   | Targets  | TAT |
|---------------------------|--------------|-------------------------|-----------|-------------------|--|-----|
| Aberdeen                  | NGS          | Whole gene<br>screen    |           | SNVs, indels      | PKP2, DSG2, DSC2, DSP,<br>SCN5A, ABCC9, DES, HCN4,<br>JUP, LMNA, PLN, RYR2,<br>TGFB3, TMEM43 | 112 |
| Family mer                | nber testing |                         |           | as indicate       | ed above   | 14  |
| Proforma required? YES Ca |              |                         | Cardiac A | rrhythmia Proforn | na (see centre website)  | •   |

#### Referral criteria

- A possible, borderline or definite diagnosis according to 2010 modified Task Force criteria
- Fibrosis & fatty replacement of myocardium affecting one or both ventricles seen on Echocardiogram or Post mortem investigations
- Clinical phenotype considered to be compatible with ACM (e.g. dilated cardiomyopathy, arrhythmia, heart failure)

- Cardiologist with expertise in ICC
- Clinical Genetics
- Pathology





#### **ATRIAL FIBRILLATION**

# Available testing

| Centre      | Method       | Scope and range of test                              |  | nge of test  | Targets  | TAT |
|-------------|--------------|--|--|--------------|--|-----|
| Aberdeen    | NGS          | Whole gene<br>screen                                 |  | SNVs, indels | SCN5A, ABCC9, GJA1, GJA5,<br>HCN4, KCNA5, KCNE5, NPPA,<br>SCN2B, SCN4B | 56  |
| Family mer  | nber testing | as indicated above                                   |  |              | ed above   | 14  |
| Proforma re | equired?     | YES Cardiac Arrhythmia Proforma (see centre website) |  |              |  |     |

#### Referral criteria

 Atrial fibrillation detected at young age with family history of atrial fibrillation or sudden cardiac death

- · Cardiologist with expertise in ICC
- Clinical Genetics





#### **BARTH SYNDROME**

# Available testing

| Centre                | Method | Scope and range of test        |                | Targets        | TAT |
|-----------------------|--------|--------------------------------|----------------|----------------|-----|
| Aberdeen              | Sanger | Whole gene screen SNVs, indels |                | TAFAZZIN (TAZ) | 56  |
| Family member testing |        |                                | as indicated a | bove           | 14  |
| Proforma required?    |        | NO                             |                |                |     |

#### Referral criteria

- Cardiomyopathy
- Neutropenia
- Fatigue & general muscle weakness
- Growth / feeding issues

- Cardiology
- Clinical Genetics
- Paediatrics





#### BRUGADA SYNDROME AND SODIUM CHANNEL DISEASE

# Available testing

| Centre      | Method   | Sco                  | Scope and range of test                             |              | Targets  | TAT |
|-------------|----------|----------------------|---|--------------|--|-----|
| Aberdeen    | NGS      | Whole gene<br>screen |   | SNVs, indels | SCN5A, CACNA1C, CACNA2D1,<br>CACNB2, GPD1L, HCN4, KCND3,<br>KCNE3, KCNE5, KCNJ8,<br>RANGRF, SCN1B, SCN2B,<br>SCN3B, SCN10A, SLMAP, TRPM4 | 112 |
| Family r    |          |                      |   | as indica    | ited above   | 14  |
| Proforma re | equired? | YES                  | ES Cardiac Arrhythmia Proforma (see centre website) |              |  |     |

#### Referral criteria

- Cardiac arrest in the absence of secondary causes, most commonly at night
- Arrhythmia triggered by fever
- Type 1 Brugada ECG
- Atrial arrhythmia, sinus node dysfunction, or conduction disease, with or without QT prolongation predominantly in children and young people.

- Cardiologist with expertise in ICC
- Clinical Genetics





# CATECHOLAMINERGIC POLYMORPHIC VENTRICULAR TACHYCARDIA (CPVT)

#### Available testing

| Centre  | Method | Scope and range of test |  | nge of test  | Targets                                  | TAT |
|---|--------|-------------------------|--|--------------|--|-----|
| Aberdeen  | NGS    | Whole gene screen       |  | SNVs, indels | RYR2, CALM1, CALM2, CASQ2,<br>DPP6, TRDN | 56  |
| Family men testing  | nber   | as indicated above      |  |              |  | 14  |
| Proforma required? YES Cardiac Arrhythmia Proforma (see centre website) |        |                         |  |              |  |     |

#### Referral criteria

- Ventricular fibrillation or polymorphic VT.
- Bi-directional VT on exercise.
- Resuscitated from cardiac arrest, or syncope compatible with tachyarrhythmia especially related to physical activity, or acute emotion, in the presence of an unremarkable ECG (e.g. normal QT interval), and in the absence of structural heart or coronary artery disease.
- Family history of premature sudden cardiac death particularly due to physical activity or emotion.

- Cardiologist with expertise in ICC
- Clinical Genetics
- Pathology in discussion with Clinical Genetics





# **DILATED CARDIOMYOPATHY (DCM)**

#### Available testing

| Centre                | Method | Scope and ra         | nge of test  | Targets  | TAT |
|-----------------------|--------|----------------------|--------------|--|-----|
| Edinburgh             | NGS    | Whole gene<br>screen | SNVs, indels | ACTC1, ACTN2, BAG3, CSRP3,<br>DES, DMD, DSP, FLNC, LAMP2,<br>LMNA, MYBPC3, MYH7, MYL2,<br>MYL3, NKX2-5, PLN, RBM20,<br>SCN5A, TNNC1, TNNI3, TNNI3K,<br>TNNT2, TPM1, TTN (N2-B isoform),<br>VCL | 56  |
| Family member testing |        | as indicated above   |              |  |     |
| Proforma required?    |        | NO                   |              |  |     |

#### Referral criteria

- Left ventricular failure with echocardiographic/MRI evidence of dilated cardiomyopathy (REQUIRED)
- Patients with left ventricular dilatation due to coronary artery disease or haemochromatosis do not require genetic testing with this panel.
- If other potential precipitants are present hypertension, hypo / hyperthyroidism, myocarditis, peripartum, alcohol abuse, exposure to cardiotoxic drugs, then expert advice should be sought prior to genetic testing.
- Family history of skeletal myopathy, cardiomyopathy or related sudden death please provide details (including the diagnosis) of the affected relatives.
- Pathologically confirmed non-ischaemic dilated cardiomyopathy at post mortem.

- Cardiologist with expertise in ICC
- Clinical Genetics
- Pathology in discussion with Clinical Genetics





#### **HEART BLOCK**

# Available testing

| Centre                | Method | Scope and range of test |  |              | Targets                     | TAT |
|-----------------------|--------|-------------------------|--|--------------|-----------------------------|-----|
| Aberdeen              | NGS    | Whole gene screen       |  | SNVs, indels | SCN5A, HCN4, LMNA,<br>TRPM4 | 56  |
| Family member testing |        | as indicated above      |  |              | 14                          |     |
| Proforma required?    |        | YES                     | YES Cardiac Arrhythmia Proforma (see centre website) |              |                             |     |

#### Referral criteria

- Heart block (see also Brugada and sodium channel disease)
- Syncope associated with heart block

- Cardiologist with expertise in ICC
- Clinical Genetics





#### HYPERTROPHIC CARDIOMYOPATHY (HCM)

#### Available testing

| <u> </u>                 |        |                      |                 |   |     |  |  |  |
|--------------------------|--------|----------------------|-----------------|---|-----|--|--|--|
| Centre                   | Method | Scope and rar        | ige of test     | Targets   | TAT |  |  |  |
| Edinburgh                | NGS    | Whole gene<br>screen | SNVs,<br>indels | ACTC1, ACTN2, CSRP3, FHL1, FLNC,<br>GLA, JPH2, LAMP2, MYBPC3, MYH7,<br>MYL2, MYL3, PLN, PRKAG2, TNNC1,<br>TNNI3, TNNT2, TPM1, TTR | 56  |  |  |  |
| Family member<br>testing |        |                      | as in           | dicated above   | 14  |  |  |  |
| Proforma required?       |        | NO                   |                 |   |     |  |  |  |

| Centre             | Method | Scope and ran     | ige of test     | Targets                              | TAT |
|--------------------|--------|-------------------|-----------------|--------------------------------------|-----|
| Edinburgh          | Sanger | Whole gene screen | SNVs,<br>indels | Familial amyloid polyneuropathy: TTR | 56  |
| Proforma required? |        | NO                |                 |                                      |     |

#### Referral criteria

- ECG or echocardiographic/MRI evidence of hypertrophic cardiomyopathy (REQUIRED)
- No evidence of hypertensive or valvular heart disease sufficient to cause cardiac hypertrophy
- Family history of skeletal myopathy, cardiomyopathy or related sudden death
   please provide medical details of the affected relatives.
- Pathologically confirmed HCM at post-mortem with no history of hypertension or evidence of valvular heart disease sufficient to cause cardiac hypertrophy.

- Cardiologist with expertise in ICC
- Clinical Genetics
- Pathology in discussion with Clinical Genetics





#### LONG QT SYNDROME

#### Available testing

| Centre                | Method | Scope and range of test |  |                                    | Targets  | TAT |
|-----------------------|--------|-------------------------|--|------------------------------------|--|-----|
| Aberdeen              | NGS    | Whole gene<br>screen    |  | SNVs, indels<br>Exon level<br>CNV* | KCNQ1*, KCNH2*, KCNE1*,<br>KCNE2*, SCN5A, KCNJ2, ANK2,<br>AKAP9, CACNA1C, CALM1,<br>CALM2, CAV3, KCNJ5, NOS1AP,<br>SCN4B, SNTA1, TRPM4 | 112 |
| Family member testing |        |                         | as indicated above                               |                                    |  | 14  |
| Proforma required?    |        | YES                     | Cardiac Arrhythmia Proforma (see centre website) |                                    |  |     |

#### Referral criteria

- Abnormal ECG (QTc ≥440ms in males, ≥460ms in females)
- Syncope or apparent seizures compatible with ventricular tachyarrhythmia, especially relating to stress or high emotion, physical activity including swimming, sudden loud noise or at rest or in bed.
- Exclude other causes of QT prolongation (e.g. QT prolonging drugs, electrolyte or calcium disturbance, hypothyroidism, ischaemia, dilated cardiomyopathy)
- Family history of unexplained premature sudden cardiac death, syncope or seizures among immediate family members.

- Cardiologist with expertise in ICC
- Clinical Genetics
- Pathology in discussion with Clinical Genetics





#### **MARFAN SYNDROME**

#### Available testing

| Centre                | Method                     | Scope and range of test |              | Targets              | TAT |
|-----------------------|----------------------------|-------------------------|--------------|----------------------|-----|
| Dundee                | NGS<br>(targeted<br>panel) | Whole gene screen       | SNVs, indels | FBN1, TGFBR1, TGFRB2 | 56  |
| Family member testing |                            |                         | 14           |                      |     |
| Proforma required?    |                            | NO                      |              |                      |     |

#### Referral criteria

- Clinical features of Marfan syndrome giving a Ghent systemic score of ≥ 5 in an adult over 18
  vears
- In children, clinical features of Marfan syndrome giving a lower Ghent score following assessment in a clinical service with expertise in the diagnosis of Marfan syndrome.
- Clinical features suggestive of Loeys-Dietz syndrome
- Ectopia lentis if other causes such as homocystinuria (due to cystathionine beta-synthase deficiency) have been excluded.
- Aortic sinus dilatation, defined as z score >3 for body surface area in children, and > 2 for body surface area in adults. See also Thoracic Aortic Aneurysm and Dissection.
- Thoracic aortic aneurysm or dissection. See also Thoracic Aortic Aneurysm and Dissection.

- Cardiologist with expertise in ICC
- Clinical Genetics





#### PAEDIATRIC CARDIOMYOPATHY

#### Available testing

| AARS2, ABCC9, ACAD9, ACADVL, ACTA1, ACTC1, ACTN2, AGK, AGL, ALIMS1, ALIPK3, ARSB, ATP5D, ATPAF2, BAG3, BRAF, CACNA1C, CBL, CDH2, COA5, COA6, COX10, COX14, COX15, COX20, COX6B1, CPT2, CSRP3, DES, DMD, DNAJC19, DOLK, DSC2, DSC2, DSP, EMD, EPG5, FAH, FHL1, FHOD3, FKTN, FLNC, GAA, GLB1, GUSB, HADHA, HADHB, HCN4, HRAS, IDH2, IDS, IDUA, JPH2, JUP, KRAS, LAMP2, LIMNA, LRPPRC, LZTR1, MAP2K1, MAP2K2, MIB1, MLYCD, MRPL44, MUT, MYBPC3, MYH6, MYH7, MYL2, MYL3, MYPN, NDUFA1, NDUFA10, NDUFA11, NDUFA2, NDUFAF1, NDUFAF2, NDUFAF3, NDUFAF4, NDUFAF5, NDUFS1, NDUFS3, NDUFS1, NDUFS2, NDUFV1, NDUFS4, NDUFS6, NDUFS7, NDUFS8, NDUFV1, NDUFV2, NEXN, NF1, NKX2-5, NONO, NRAS, NUBPL, PCCA, PCCB, PDLIM3, PKP2, PLN, PNPLA2, PPA2, PPCS, PPP1CB, PPP1R13L, PRKAG2, PTPN11, RAF1, RBM20, RIT1, RYR2, SCN5A, SCO1, SCO2, SDHA, SDHAF1, SDHD, SGCD, SHOC2, SLC25A4, SOS1, SOS2, SURF1, TAZ, TBX5, TMEM126B, TMEM43, TMEM70, TNNC1, TNNI3, TNNI3K, TNNT2, TPM1, TSFM, TTN, TTR, VCL  as indicated above  14 | Centre   | Method   | Scope and | range of test        | Targets   | TAT |
|---|----------|----------|-----------|----------------------|---|-----|
| Family member as indicated above 14 testing   | Aberdeen | NGS      | gene      | indels<br>Exon level | ACTN2, AGK, AGL, ALMS1, ALPK3, ARSB, ATP5D, ATPAF2, BAG3, BRAF, CACNA1C, CBL, CDH2, COA5, COA6, COX10, COX14, COX15, COX20, COX6B1, CPT2, CSRP3, DES, DMD, DNAJC19, DOLK, DSC2, DSG2, DSP, EMD, EPG5, FAH, FHL1, FHOD3, FKTN, FLNC, GAA, GLB1, GUSB, HADHA, HADHB, HCN4, HRAS, IDH2, IDS, IDUA, JPH2, JUP, KRAS, LAMP2, LMNA, LRPPRC, LZTR1, MAP2K1, MAP2K2, MiB1, MLYCD, MRPL44, MUT, MYBPC3, MYH6, MYH7, MYL2, MYL3, MYPN, NDUFA1, NDUFA10, NDUFA11, NDUFA2, NDUFAF1, NDUFAF2, NDUFAF3, NDUFAF4, NDUFAF5, NDUFB11, NDUFB3, NDUFS1, NDUFS2, NDUFS3, NDUFS4, NDUFS6, NDUFS7, NDUFS8, NDUFV1, NDUFV2, NEXN, NF1, NKX2-5, NONO, NRAS, NUBPL, PCCA, PCCB, PDLIM3, PKP2, PLN, PNPLA2, PPA2, PPCS, PPP1CB, PPP1R13L, PRKAG2, PTPN11, RAF1, RBM20, RIT1, RYR2, SCN5A, SCO1, SCO2, SDHA, SDHAF1, SDHD, SGCD, SHOC2, SLC22A5, SLC25A20, SLC25A4, SOS1, SOS2, SURF1, TAZ, TBX5, TMEM126B, TMEM43, TMEM70, TNNC1, TNNI3, TNNI3K, TNNT2, | 112 |
|   | _        | mber     |           |                      | as indicated above  | 14  |
|   |          | equired? | NO        |                      |   | ı   |

#### Referral criteria

- Child (under 16) with cardiomyopathy where no other non-genetic cause has been found, and there is no family history of Adult Onset Cardiomyopathy.
- If there is a family history of "non-syndromic" adult onset cardiomyopathy (dilated, hypertrophic) then the relevant adult cardiomyopathy panel should be considered instead.
- If there are features of a specific "non-syndromic" cardiomyopathy such as Arrhythmogenic Cardiomyopathy, then the Arrhythmogenic Cardiomyopathy panel should be considered instead.
- If the cardiomyopathy is one of multiple features of a likely multisystem disorder suggestive of Noonan syndrome or a Rasopathy, then the Noonan/Rasopathy panel should be considered instead.
- If the cardiomyopathy is one of multiple features of a likely multisystem disorder not suggestive of Noonan syndrome or a Rasopathy, please seek expert advice as a broader spectrum test may be appropriate.

- Cardiologist with expertise in ICC
- **Clinical Genetics**
- Pathology in discussion with Clinical Genetics





#### SHORT QT SYNDROME

# Available testing

| Centre      | Method  |       | Scope and range of test |                    | Targets             | TAT |
|-------------|---|-------|-------------------------|--------------------|---------------------|-----|
| Aberdeen    | NGS   | Whole | gene screen             | SNVs, indels       | KCNQ1, KCNH2, KCNJ2 | 56  |
| Family men  | nber testing  |       |                         | as indicated above |                     |     |
| Proforma re | Proforma required? YES Cardiac Arrhythmia Proforma (see centre website) |       |                         | ee centre website) |                     |     |

#### Referral criteria

- Abnormal ECG (QTc ≤360ms in males, ≤370ms in females)
- Syncope compatible with tachyarrhythmia or cardiac arrest.
- A family history of SCD at age < 40 years

- Cardiologist with expertise in ICC
- Clinical Genetics





# THORACIC AORTIC ANEURYSM & DISSECTION (TAAD)

#### Available testing

| Centre                | Method                     | Scope and range of test |              | Targets  | TAT |
|-----------------------|----------------------------|-------------------------|--------------|--|-----|
| Dundee                | NGS<br>(targeted<br>panel) | Whole gene<br>screen    | SNVs, indels | ABL1, ACTA2, ARIH1, ASPH, BGN, CBS,<br>COL3A1, COL5A1, COL5A2, EFEMP2,<br>ELN, FBLN5, FBN1, FBN2, FKBP14, ,<br>FLNA, FOXE3, IPO8, LOX, MFAP5, ,<br>MYH11, MYLK, NOTCH1, PLOD1, PRKG1,<br>SKI, SLC2A10, SMAD2, SMAD3, SMAD4,<br>SMAD6, TGFB2, TGFB3, TGFBR1,<br>TGFBR2, THSD4 | 112 |
| Family mer<br>testing | mber                       | as                      |              | ndicated above   | 14  |
| Proforma r            | equired?                   | NO                      |              |  |     |

#### Referral criteria

- Thoracic aortic aneurysm\* or dissection with onset before age 60 and no classical cardiovascular risk factors
- Aneurysm or dissection of any part of the aorta during pregnancy
- Clinical features of Marfan syndrome giving a Ghent systemic score of ≥ 5 in an adult over 18 years
- Aortic sinus dilatation, defined as z score >3 for body surface area in children, and > 2 for body surface area in adults.
- Clinical features suggestive of Loeys-Dietz syndrome
- High clinical suspicion of a condition predisposing to aortic/arterial disease
   AND diagnostic testing for other conditions such as Ehlers Danlos syndrome
   (where indicated) has not identified a cause
- Any deceased individual with a thoracic aortic aneurysm\* or dissection detected at autopsy meeting one of the above criteria and who have relatives who will benefit from cascade testing using a genetic diagnosis
- \*Thoracic aortic aneurysm defined as:
- In children: z score >2 for body surface area
- In adults: z score > 2 for body surface area or dilatation >38 mm

- Cardiologist with expertise in ICC
- Clinical Genetics
- Pathology in discussion with Clinical Genetics
- Cardiothoracic surgery in discussion with clinical genetics





# **CHROMOSOME BREAKAGE**

## **ATAXIA TELANGIECTASIA (& AT-LIKE)**

# Available testing

| Centre                | Method    | Scope and range of test |                    | Targets            | TAT |
|-----------------------|-----------|-------------------------|--------------------|--------------------|-----|
| Aberdeen              | Karyotype | Whole genome screen     | Chromosomes 7 & 14 | Chromosomes 7 & 14 | 28  |
| Aberdeen              | NGS       | Whole gene screen       | SNVs, indels       | ATM, MRE11         | 56  |
| Family member testing |           |                         | as indicat         | ed above           | 14  |
| Proforma required?    |           | NO                      |                    |                    |     |

#### Referral criteria

- Clinical phenotype suggestive of ataxia telangiectasia elevated serum AFP levels and ≥1 of the following criteria:
  - Progressive gait and truncal ataxia with onset between 1-4 years old,
     Ocular motor apraxia, Ocular telangiectasia, Chorea and dysarthia,
     Frequent infections (Immunodeficiency), Malignancy

- Clinical Genetics
- Haematology
- Oncology in discussion with Clinical Genetics





## ATAXIA WITH OCULOMOTOR APRAXIA & HYPOALBUMINEMIA

#### Available testing

| Centre      | Method         | Scope and range of test |              | Targets | TAT |
|-------------|----------------|-------------------------|--------------|---------|-----|
| Aberdeen    | Sanger         | Whole gene screen       | SNVs, indels | APTX    | 56  |
| Family r    | member<br>ting |                         | as indicated | d above | 14  |
| Proforma re | equired?       | NO                      |              |         |     |

#### Referral criteria

 Clinical phenotype suggestive of ataxia with oculomotor apraxia & hypoalbuminemia

## Requesting specialties

- Clinical Genetics
- Haematology

#### **BLOOM SYNDROME**

## Available testing

| Centre             | Method                              | Scope and range of test |              | Targets     | TAT |
|--------------------|-------------------------------------|-------------------------|--------------|-------------|-----|
| Aberdeen           | Chromosome<br>breakage<br>analysis* | Whole genome screen     | Aneuploidy   | Genome wide | 28  |
| Aberdeen           | NGS                                 | Whole gene screen       | SNVs, indels | BLM         | 56  |
| Family me          | y member testing as indicated       |                         | d above      | 14          |     |
| Proforma required? |                                     | NO                      |              |             | •   |

<sup>\*5</sup>ml lithium heparin blood sample required. Send guaranteed next day delivery directly to Aberdeen laboratory, preferably on Monday-Wednesday. If possible, please also send an anonymised control blood (5ml lithium heparin) with completed control form (available on centre website).

#### Referral criteria

- Clinical phenotype suggestive of Bloom syndrome growth deficiency, sunsensitive, telangiectatic, hypo- and hyperpigmented skin
- Confirmed diagnosis from chromosome breakage analysis

## Requesting specialties

- Clinical Genetics
- Haematology

NSD611-003.20 V





#### CEREBRO-OCULO-FACIO-SKELETAL SYNDROME

## Available testing

| Centre      | Method   | Scope and range of test |              | Targets             | TAT |
|-------------|----------|-------------------------|--------------|---------------------|-----|
| Aberdeen    | NGS      | Whole gene screen       | SNVs, indels | ERCC1, ERCC2, ERCC6 | 56  |
| Family r    |          |                         | as indic     | ated above          | 14  |
| Proforma re | equired? | NO                      |              |                     |     |

#### Referral criteria

 Clinical phenotype suggestive of Cerebro oculo facio skeletal syndrome – microcephaly, congenital cataracts, severe mental retardation, facial dysmorphism, arthrogryposis

## Requesting specialties

- Clinical Genetics
- Haematology

#### **COCKAYNE SYNDROME**

#### Available testing

| Centre                | Method | Scope and range of test |              | Targets      | TAT |
|-----------------------|--------|-------------------------|--------------|--------------|-----|
| Aberdeen              | NGS    | Whole gene<br>screen    | SNVs, indels | ERCC6, ERCC8 | 56  |
| Family member testing |        |                         | as indic     | ated above   | 14  |
| Proforma required?    |        | NO                      |              |              |     |

#### Referral criteria

 Clinical diagnosis of Cockayne syndrome – mental retardation, microcephaly, progressive neurologic & retinal degeneration, skeletal abnormalities, gait defects, sun sensitivity

- Clinical Genetics
- Haematology





#### **DUANE-RADIAL RAY & IVIC SYNDROME**

## Available testing

| Centre      | Method   | Scope and range of test |              | Targets    | TAT |
|-------------|----------|-------------------------|--------------|------------|-----|
| Aberdeen    | Sanger   | Whole gene screen       | SNVs, indels | SALL4      | 56  |
| Family r    |          |                         | as indic     | ated above | 14  |
| Proforma re | equired? | NO                      |              |            |     |

#### Referral criteria

• Clinical phenotype suggestive of Duane-radial ray & IVIC syndrome – upper limb anomalies, ocular anomalies, renal anomalies

- Clinical Genetics
- Haematology





#### **FANCONI ANAEMIA**

## Available testing

| Centre                | Method                              | Scope and range of test |  | Targets  | TAT |
|-----------------------|-------------------------------------|-------------------------|--|--|-----|
| Aberdeen              | Chromosome<br>breakage<br>analysis* | Whole genome screen     | Chromosome<br>breakage                             | Whole genome   | 28  |
| Aberdeen              | NGS                                 | Whole gene<br>screen    | SNVs, indels<br>Exon level CNV<br>(limited genes*) | BRCA2, BRIP1, ERCC4, FANCA,<br>FANCB, FANCC, FANCD2, FANCE,<br>FANCF, FANCG, FANCI, FANCL,<br>PALB2, RAD51C, SLX4, TOP3A,<br>UBE2T | 112 |
| Family member testing |                                     |                         | as indica  | ated above   | 14  |
| Proforma required?    |                                     | NO                      |  |  |     |

<sup>\*5</sup>ml lithium heparin blood sample required. Send guaranteed next day delivery directly to Aberdeen laboratory, preferably on Monday-Wednesday. If possible, please also send an anonymised control blood (5ml lithium heparin) with completed control form (available on centre website).

#### Referral criteria

- Clinical phenotype suggestive of Fanconi anaemia persistent or recurrent pancytopenia, short stature, abnormal skin pigmentation, skeletal malformations of the upper and lower limbs, microcephaly, and ophthalmic and genitourinary tract anomalies.
- Confirmed diagnosis from chromosome breakage analysis

- Clinical Genetics
- Haematology
- Immunology





#### **GROWTH FAILURE IN EARLY CHILDHOOD**

## Available testing

| Centre      | Method         | Scope and               | range of test                      | Targets   | TAT |
|-------------|----------------|-------------------------|------------------------------------|---|-----|
| Aberdeen    | NGS            | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNVs | ACAN, ANKRD11, BLM, BRAF, BRCA2, BRIP1,<br>CBL, CCDC8, CDKN1C, CUL7, ERCC4, FANCA,<br>FANCB, FANCC, FANCD2, FANCE, FANCF,<br>FANCG, FANCI, FANCL, FGFR3, HMGA2, HRAS,<br>IGF1, IGF1R, IGF2, KRAS, LZTR1, MAP2K1,<br>MAP2K2, NBN, NRAS, OBSL1, PALB2, PIK3R1,<br>PLAG1, PPP1CB, PTPN11, RAF1, RIT1, SHOC2,<br>SLX4, SOS1, SOS2, SRCAP, TOP3A, TRIM37,<br>UBE2T | 112 |
| ,           | member<br>tina |                         |                                    | as indicated above  | 14  |
| Proforma re |                | NO                      |                                    |   |     |

#### Referral criteria

• Height/length more than 3 standard deviations below the mean at the age of at least 2 years.

# Requesting specialties

Clinical Genetics

#### **HOLT-ORAM SYNDROME**

# Available testing

| Centre           | Method   | Scope and range of test |              | Targets    | TAT |
|------------------|----------|-------------------------|--------------|------------|-----|
| Aberdeen         | Sanger   | Whole gene screen       | SNVs, indels | TBX5       | 56  |
| Family r<br>test |          |                         | as indic     | ated above | 14  |
| Proforma re      | equired? | NO                      |              |            |     |

## Referral criteria

• Clinical phenotype suggestive of Holt-Oram Syndrome – Congenital heart defect/cardiac conduction disease and upper limb malformation

# Requesting specialties





# IMMUNODEFICIENCY-CENTROMERIC INSTABILITY-FACIAL ANOMALIES SYNDROME

## Available testing

| Centre      | Method                             | Scope and range of test |                          | Targets                      | TAT |
|-------------|------------------------------------|-------------------------|--------------------------|------------------------------|-----|
| Aberdeen    | Karyotype                          | Whole genome screen     | Chromosomes<br>1, 9 & 16 | Chromosomes 1, 9 & 16        | 28  |
| Aberdeen    | NGS                                | Whole gene screen       | SNVs, indels             | DNMT3B, ZBTB24, CDCA7, HELLS | 56  |
|             | Family member as indicates testing |                         | ted above                | 14                           |     |
| Proforma re | equired?                           | NO                      |                          |                              |     |

#### Referral criteria

 Clinical phenotype suggestive of Immunodeficiency-Centromeric Instability-Facial Anomalies Syndrome

- Clinical Genetics
- Haematology
- Immunology





#### **LIG4 SYNDROME**

## Available testing

| Centre             | Method                             | Scope and range of test |              | Targets | TAT |
|--------------------|------------------------------------|-------------------------|--------------|---------|-----|
| Aberdeen           | Sanger                             | Whole gene screen       | SNVs, indels | LIG4    | 56  |
| ,                  | Family member as indicated testing |                         | above        | 14      |     |
| Proforma required? |                                    | NO                      |              |         |     |

#### Referral criteria

 Clinical phenotype suggestive of LIG4 syndrome – immunodeficiency, developmental delay, growth delay

# Requesting specialties

- Clinical Genetics
- Haematology
- Immunology

# **MEIER-GORLIN SYNDROME**

## Available testing

| Centre                | Method   | Scope and range of test |              | Targets                         | TAT |
|-----------------------|----------|-------------------------|--------------|---------------------------------|-----|
| Aberdeen              | NGS      | Whole gene screen       | SNVs, indels | ORC1, ORC4, ORC6, CDT1,<br>CDC6 | 56  |
| Family member testing |          | as indicated            | above        | 14                              |     |
| Proforma re           | equired? | NO                      |              |                                 |     |

#### Referral criteria

 Clinical phenotype suggestive of Meier-Gorlin syndrome – severe intrauterine & postnatal growth retardation, microcephaly, bilateral microtia, aplasia or hypoplasia of patellae

## Requesting specialties





# NATURAL KILLER CELL AND GLUCOCORTICOID DEFICIENCY WITH DNA REPAIR DEFECT

## Available testing

| Centre      | Method                 | Scope and range of test |              | Targets | TAT |
|-------------|------------------------|-------------------------|--------------|---------|-----|
| Aberdeen    | NGS                    | Whole gene screen       | SNVs, indels | MCM4    | 56  |
|             | Family member as indic |                         | ated above   | 14      |     |
| Proforma re | equired?               | NO                      |              |         |     |

#### Referral criteria

 Clinical phenotype suggestive of a Natural killer Cell & Glucocorticoid deficiency with DNA repair defect – growth retardation, microcephaly, decreased numbers of natural killer cells, recurrent infection, respiratory failure

- Clinical Genetics
- Haematology
- Immunology





# **NIJMEGEN BREAKAGE SYNDROME (& NBS-LIKE)**

## Available testing

| Centre             | Method        | Scope and range of test |                            | Targets            | TAT |
|--------------------|---------------|-------------------------|----------------------------|--------------------|-----|
| Aberdeen           | Karyotype     | Whole genome screen     | Chromosomes 7 & 14 studies | Chromosomes 7 & 14 | 28  |
| Aberdeen           | NGS           | Whole gene screen       | SNVs, indels               | NBN, RAD50         | 56  |
| Family me          | ember testing |                         | as indica                  | ted above          | 14  |
| Proforma required? |               | NO                      |                            |                    |     |

#### Referral criteria

- Clinical phenotype suggestive of Nijmegen Breakage Syndrome microcephaly, growth retardation, immunodeficiency
- Confirmed diagnosis from chromosome breakage analysis

- Clinical Genetics
- Haematology
- Immunology





#### **ROBERTS-SC PHOCOMELIA SYNDROME**

## Available testing

| Centre      | Method         | Scope and range of test |              | Targets     | TAT |
|-------------|----------------|-------------------------|--------------|-------------|-----|
| Aberdeen    | Karyotype      | Whole genome screen     | Aneuploidy   | Genome wide | 28  |
| Aberdeen    | Sanger         | Whole gene screen       | SNVs, indels | ESCO2       | 56  |
|             | member<br>ting | as indicated above      |              |             | 14  |
| Proforma re | equired?       | NO                      |              |             |     |

#### Referral criteria

 Clinical phenotype suggestive of Roberts / SC phocomelia syndrome – growth retardation, extremity malformations, craniofacial anomalies, developmental delay, cardiac anomalies, renal anomalies

# Requesting specialties

Clinical Genetics

#### ROTHMUND-THOMSON / RAPADILINO / BALLER-GEROLD

#### Available testing

| Centre             | Method                               | Scope and range of test |              | Targets | TAT |
|--------------------|--------------------------------------|-------------------------|--------------|---------|-----|
| Aberdeen           | NGS                                  | Whole gene screen       | SNVs, indels | RECQL4  | 56  |
|                    | Family member as indicated a testing |                         | above        | 14      |     |
| Proforma required? |                                      | NO                      |              |         |     |

#### Referral criteria

 Clinical phenotype suggestive of Rothmund Thomson / Rapadilino / Baller-Gerold

- Clinical Genetics
- Dermatology





#### **SECKEL SYNDROME**

## Available testing

| Centre      | Method                 | Scope and range of test |              | Targets                   | TAT |
|-------------|------------------------|-------------------------|--------------|---------------------------|-----|
| Aberdeen    | NGS                    | Whole gene screen       | SNVs, indels | ATR, RBBP8, CEP152, CENPJ | 56  |
|             | Family member as indic |                         | ated above   | 14                        |     |
| Proforma re | equired?               | NO                      |              |                           |     |

#### Referral criteria

• Clinical phenotype suggestive of Seckel Syndrome – growth retardation, microcephaly with mental retardation, characteristic facial appearance

# Requesting specialties

Clinical Genetics

#### TAR SYNDROME

# Available testing

| Centre             | Method                 | Scope and range of test |              | Targets | TAT |
|--------------------|------------------------|-------------------------|--------------|---------|-----|
| Aberdeen           | Sanger                 | Whole gene screen       | SNVs, indels | RBM8A   | 56  |
| _                  | Family member as indic |                         | ated above   | 14      |     |
| Proforma required? |                        | NO                      |              |         |     |

#### Referral criteria

Clinical phenotype suggestive of Thrombocytopenia-absent radius syndrome

- Clinical Genetics
- Haematology





#### **TOWNES-BROCKS SYNDROME**

## Available testing

| Centre                | Method   | Scope and range of test |              | Targets | TAT |
|-----------------------|----------|-------------------------|--------------|---------|-----|
| Aberdeen              | Sanger   | Whole gene screen       | SNVs, indels | SALL1   | 56  |
| Family member as indi |          | as indicated            | above        | 14      |     |
| Proforma re           | equired? | NO                      |              |         |     |

#### Referral criteria

 Clinical phenotype suggestive of Townes-Brocks Syndrome – triad of imperforate anus, dysplastic ears & thumb malformations

## Requesting specialties

Clinical Genetics

#### **TRICOTHIODYSTROPHY**

# Available testing

| Centre                               | Method | Scope and range of test |              | Targets                         | TAT |
|--------------------------------------|--------|-------------------------|--------------|---------------------------------|-----|
| Aberdeen                             | NGS    | Whole gene screen       | SNVs, indels | ERCC2, ERCC3, MPLKIP,<br>GTF2H5 | 56  |
| Family member as indicated a testing |        | above                   | 14           |                                 |     |
| Proforma required?                   |        | NO                      |              |                                 |     |

## Referral criteria

• Clinical diagnosis of Tricothiodystrophy – brittle, sulfur-deficient hair which displays a diagnostic alternating light and dark banding pattern

- Clinical Genetics
- Dermatology





#### **ULNAR-MAMMARY SYNDROME**

#### Available testing

| Centre      | Method   | Scope and range of test |              | Targets    | TAT |
|-------------|----------|-------------------------|--------------|------------|-----|
| Aberdeen    | Sanger   | Whole gene screen       | SNVs, indels | TBX3       | 56  |
| Family r    |          |                         | as indic     | ated above | 14  |
| Proforma re | equired? | NO                      |              |            |     |

#### Referral criteria

 Clinical phenotype suggestive of Ulnar-Mammary Syndrome – posterior limb deficiencies or duplications, mammary gland hypoplasia and / or dysfunction, abnormal dentition, delayed puberty in males, genital anomalies

## Requesting specialties

Clinical Genetics

#### WARSAW BREAKAGE SYNDROME

#### Available testing

| Centre                | Method | Scope and range of test |              | Targets    | TAT |
|-----------------------|--------|-------------------------|--------------|------------|-----|
| Aberdeen              | NGS    | Whole gene screen       | SNVs, indels | DDX11      | 56  |
| Family member testing |        |                         | as indic     | ated above | 14  |
| Proforma required?    |        | NO                      |              |            |     |

#### Referral criteria

Clinical phenotype suggestive of Warsaw Breakage Syndrome

## Requesting specialties





#### **WERNER SYNDROME**

## Available testing

| Centre      | Method         | Scope and ra      | ange of test | Targets | TAT |
|-------------|----------------|-------------------|--------------|---------|-----|
| Aberdeen    | NGS            | Whole gene screen | SNVs, indels | WRN     | 56  |
| Family r    | member<br>ting |                   | as indicated | d above | 14  |
| Proforma re | equired?       | NO                |              |         |     |

#### Referral criteria

• Clinical phenotype suggestive of Werner syndrome – accelerated aging, bilateral cataracts, diabetes mellitus, osteoporosis, premature arteriosclerosis

## Requesting specialties

- Clinical Genetics
- Dermatology

#### **XERODERMA PIGMENTOSUM**

## Available testing

| Centre                | Method   | Scope and ra         | ange of test | Targets   | TAT |  |  |
|-----------------------|----------|----------------------|--------------|---|-----|--|--|
| Aberdeen              | NGS      | Whole gene<br>screen | SNVs, indels | XPA, XPC, ERCC1, ERCC3, ERCC4,<br>ERCC5, DDB2, POLH | 56  |  |  |
| Family member testing |          | as indicated above   |              |   |     |  |  |
| Proforma re           | equired? | NO                   |              |   |     |  |  |

## Referral criteria

 Clinical diagnosis of Xeroderma Pigmentosum – XP-related features in eye, neurological systems or related cancer

- Clinical Genetics
- Dermatology





# **CONNECTIVE TISSUE DISORDERS**

#### **CONNECTIVE TISSUE**

#### Available testing

| Centre                | Method | Scope and range of test |                    | Targets   |     |
|-----------------------|--------|-------------------------|--------------------|---|-----|
| Edinburgh             | NGS    | Whole gene<br>screen    | SNVs,<br>indels    | ABCC6, ACTA2, ACVR1, ADAMTS2, ALPL,<br>ATP6V0A2, B3GALT6, B4GALT7, BMP1, CBS,<br>CHST14, COL11A1, COL1A1, COL1A2, COL2A1,<br>COL3A1, COL5A1, COL5A2, CRTAP, ELN, FBLN5,<br>FBN1, FBN2, FKBP10, FKBP14, IFITM5, LEPRE1<br>(P3H1), LRP5, MYLK, NOTCH1, NOTCH2, PKD2,<br>PLOD1, PLOD2, PPIB, PRDM5, RIN2, SERPINF1,<br>SERPINH1, SLC2A10, SLC39A13, SMAD3, SP7,<br>TGFB2, TGFBR1, TGFBR2, TNXB, ZNF469 | 112 |
| Family member testing |        |                         | as indicated above | 14  |     |
| Proforma required?    |        | NO                      |                    |   | I   |

| Centre                | Method   | Scope and range of test |                    | Targets                         | TAT |
|-----------------------|----------|-------------------------|--------------------|---------------------------------|-----|
| Edinburgh             | NGS      | Whole gene<br>screen    | SNVs,<br>indels    | Pseudoxanthoma elasticum: ABCC6 | 112 |
| Family member testing |          |                         | as indicated above | 14                              |     |
| Proforma re           | equired? | NO                      |                    |                                 |     |

| Centre                | Method                | Scope and range of test |                 | Targets                |     |
|-----------------------|-----------------------|-------------------------|-----------------|------------------------|-----|
| Edinburgh             | NGS                   | Whole gene<br>screen    | SNVs,<br>indels | Hypophosphatasia: ALPL | 112 |
| Family mer            | Family member testing |                         |                 | as indicated above     | 14  |
| Proforma required? NO |                       |                         |                 |                        |     |

#### Referral criteria

- See criteria for Ehlers-Danlos Syndrome
- Please contact the laboratory to discuss indications not included by above criteria

Individuals who have characteristic features of Pseudoxanthoma elasticum:

- Papules or plaques on the skin of the neck and/or flexural creases

   (antecubital fossae, axillae, groin, or popliteal fossae) and/or calcified
   dystrophic elastic fibres on biopsied skin using a von Kossa or similar stain)

   AND/OR
- Retinal finding (angioid streaks, peau d'orange, or choroidal vascularization)

Individuals who have characteristic features of Hypophosphotasia:

NSD611-003.20 V5





 clinical features of infantile hypophosphatasia (growth failure, craniotabes, craniosynostosis, blue sclerae, flail chest, costochondral enlargement, scoliosis, thickening of wrists, knees, and ankles, bowing of legs, lax ligaments, hypotonia), undermineralisation of growing / remodelling bone, pathologic fractures, premature loss of deciduous teeth with dental root remaining attached, bone pain

#### and

 biochemical abnormality showing reduced activity of serum alkaline phosphatase (ALP) or elevated urine phosphoethanolamine (PEA) \*\*\*Need referral criteria\*\*\*

- Clinical Genetics
- Rheumatology for hypophosphatasia
- Ophthalmology for Pseudoxanthoma elasticum





#### **EHLERS DANLOS SYNDROME**

#### Available testing

| Centre    | Method | Scope and range of test |                 | Targets   |     |
|-----------|--------|-------------------------|-----------------|---|-----|
| Edinburgh | NGS    | Whole<br>gene<br>screen | SNVs,<br>indels | ADAMTS2, AEBP1, ALDH18A1, ATP6V0A2, ATP6V1A, ATP7A, B3GALT6, B4GALT7, BGN, C1R, C1S, CBS, CHST14, COL12A1, COL1A1, COL1A2, COL3A1, COL5A1, COL5A2, COL6A1, COL6A2, COL6A3, DSE, EFEMP2, ELN, FBLN5, FBN1, FBN2, FKBP14, GORAB, LOX, LTBP4, PLOD1, PRDM5, PYCR1, RIN2, ROBO3, SKI, SLC39A13, SMAD2, SMAD3, TGFB2, TGFB3, TGFBR1, TGFBR2, TNXB and ZNF469 | 112 |

Gene list from panel app Ehlers Danlos syndromes panelv2.3 https://panelapp.genomicsengland.co.uk/panels/53/v2.3

#### Referral criteria

- Referral criteria as per Malfait et al (2017) Am J Med Genetics 175C:8-26
- Includes following subtypes:
  - Classic 1 and 2, classic-like 1, arthrochalasia 1 and 2, cardiac valvular, dermatosparaxis, kyphoscoliotic 1 and 2, musculocontractural 1, periodontal 1 and 2, spondylodysplastic 1, 2 and 3 and vascular
  - Combined osteogenesis imperfecta and Ehlers-Danlos syndrome 2, Macrocephaly, alopecia, cutis laxa, and scoliosis
  - o Brittle cornea syndrome 1 and 2
- Samples for Hypermobile EDS will not be accepted as the genetic basis is unknown

## Requesting specialties





#### STICKLER SYNDROME / CLEFT PALATE

## Available testing

#### STICKLER SYNDROME

| Centre    | Method | Scope and range of test |                 | Targets   |     |
|-----------|--------|-------------------------|-----------------|---|-----|
| Edinburgh | NGS    | Whole<br>gene<br>screen | SNVs,<br>indels | Stickler: COL11A1, COL11A2, COL2A1, COL9A1,<br>COL9A2, COL9A3, GZF2 | 112 |

Gene list from panel app Stickler syndrome panelv3.0 https://panelapp.genomicsengland.co.uk/panels/3/v3.0

# Cleft palate

|               |     |                         |                 |  | eft: ACTB, ACTG1, AMER1, ANKRD11,<br>GAP29, ARHGAP31, ASXL1, B3GLCT, BCOR.   |     |
|---------------|-----|-------------------------|-----------------|--|--|-----|
| Edinburgh     | NGS | Whole<br>gene<br>screen | SNVs,<br>indels | EFN ESC FLN GI IFT K MAF MSX NC F RE SC SF | BMP2, C2CD3, C5orf42, CC2D2A, CDH1, DKN1C, CHD7, CHRNG, CHST14, COL11A1, COL11A2, COL2A1, COL9A1, COLEC10, COLEC11, CTCF, CTNND1, DHCR7, DHODH, DLL4, DOCK6, DVL1, DVL3, DYNC2H1, DYNC2LI1, EBP, EDNRA, IB1, EFTUD2, EIF2S3, EIF4A3, EOGT, EPG5, CO2, EYA1, FAM20C, FGD1, FGFR1, FGFR2, A, FLNB, FOXC2, FRAS1, GJA1, GLI3, GPC3, RHL3, HDAC8, HYLS1, ICK, IFT140, IFT172, F80, IMPAD1, IRF6, KAT6A, KCNJ2, KDM6A, GIAA0586, KIF1BP, KIF7, KMT2D, MAP3K7, PRE2, MASP1, MBTPS2, MEIS2, MID1, MKS1, K1, MYMK, NECTIN1, NEDD4L, NEK1, NIPBL, DTCH1, OFD1, PAX3, PHF8, PIEZO2, PIGN, PIGV, POLR1C, POLR1D, PORCN, PTCH1, BM10, ROR2, RPL5, RPS26, SALL4, SATB2, ARF2, SEPT9 (SEPTIN9), SF3B4, SHH, SIX1, IX3, SIX5, SKI, SLC26A2, SMAD3, SMAD4, SMC1A, SMC3, SMS, SNRPB, SON, SOX9, PECC1L, STAMBP, TBX22, TCOF1, TCTN3, TELO2, TFAP2A, TGDS, TGFB3, TGFBR1, GFBR2, TMCO1, TP63, TRAPPC9, TRIM37, BB, TXNL4A, USP9X, WNT5A, XYLT1, ZEB2, ZIC2, ZIC3, ZSWIM6 | 112 |
| Family me     |     | •                       |                 | as   | s indicated above  | 14  |
| Proforma requ | ,   | NO                      |                 |  |  |     |

Gene list from panel app Clefting panelv2.2 https://nhsgms-panelapp.genomicsengland.co.uk/panels/81/v2.2

|         | Centre                | Method | Scope and               | range of te     | st Targets                   |                | TAT             |  |
|---------|-----------------------|--------|-------------------------|-----------------|------------------------------|----------------|-----------------|--|
| 77      | Edinburgh             | NGS    | Whole<br>gene<br>screen | SNVs,<br>indels | Van der Woude syndrome: IRF6 |                | 112             |  |
| / / / • | Family member testing |        | as indicated above      |                 |                              |                |                 |  |
|         |                       |        |                         | NSD             | 611-003.20 V5 P              | age <b>5</b> 8 | 3 of <b>347</b> |  |





#### Referral criteria

- Two or more of the following:
  - o Retinal detachment or: High myopia with onset before 6 years
  - o Cleft palate
  - Vitreous abnormality
  - o Joint hypermobility or premature joint degeneration
  - Sensorineural hearing loss
  - Facial features (flat midface with depressed nasal bridge, reduced nasal protrusion, anteverted nares and micrognathia)

# Requesting specialties





# **DEVELOPMENTAL DISORDERS**

## ANEUPLOIDY SCREENING - NON-INVASIVE PRENATAL TESTING (NIPT)

#### Available testing

| Centre             | Method                   | Scope and range of test |            | Targets                | TAT |
|--------------------|--------------------------|-------------------------|------------|------------------------|-----|
| Dundee             | NGS<br>(genome-<br>wide) | Targeted screen         | Aneuploidy | Chromosomes 13, 18, 21 | 7   |
| Proforma required? | YES                      | NIPT request form       | า          |                        |     |

#### Referral criteria

- Higher chance biochemical screen result (>1:150) OR
- Previous trisomy 13, 18 or 21
- Pregnancy must be >10 weeks gestation confirmed by ultrasound scan

## Exclusion criteria

NIPT is not an appropriate test if any of the following are not excluded:

- Fetal demise / vanishing twin
- Blood transfusion within 4 months
- Transplant surgery within 1 year
- Immuno / stem cell therapy within 1 year
- Maternal malignancy within 1 year
- Known maternal chromosome anomaly

- Obstetrics
- Clinical Genetics





# **ANEUPLOIDY TESTING - PRENATAL (AF / CVS)**

#### Available testing

| Centre                                     | Method | Scope and range of test |      | Targets                            | TAT |
|--|--------|-------------------------|------|------------------------------------|-----|
| Aberdeen<br>Dundee<br>Edinburgh<br>Glasgow | QF-PCR | Targeted screen         | STRs | Chromosome markers 13, 18, 21, X/Y | 3   |
| Proforma required?                         |        | NO                      |      |                                    |     |

#### Referral criteria

- Higher chance biochemical screen result (>1:150) OR
- High chance Non-Invasive Prenatal Test (NIPT) result OR
- Abnormalities detected on ultrasound scan OR
- Previous trisomy detected OR
- Family history of known single gene disorder (referral through Clinical Genetics only)
- Family history of known chromosomal rearrangement (referral through Clinical Genetics only)

- Obstetrics
- Clinical Genetics





# **ANEUPLOIDY TESTING - PRENATAL (AF / CVS)**

#### Available testing

| Centre                                     | Method     | Scope and range of test |     | Targets      | TAT |
|--|------------|-------------------------|-----|--------------|-----|
| Aberdeen<br>Dundee<br>Edinburgh<br>Glasgow | Microarray | Whole genome screen     | CNV | Whole genome | 14  |
| Proforma required?                         |            | NO                      |     |              |     |

#### Referral criteria

- One or more abnormalities detected on ultrasound scan e.g. structural heart malformations, possible tracheoesphageal fistula, possible duodenal atresia, cleft lip, structural renal malformations, bladder extrophy, absent radius unilateral or bilateral, pleural effusion OR
- An isolated nuchal translucency NT ≥ 3.5 mm when crown-rump length measures from 45 mm to 84 mm (at approximately 11 weeks 0 days to 13 weeks 6 days).

- Obstetrics
- Clinical Genetics





# ANEUPLOIDY / MICRODUPLICATION / MICRODELETION NEONATAL SCREENING (URGENT)

# Available testing

| Centre               | Method     | Scope and range of test |      | Targets                        | TAT |
|----------------------|------------|-------------------------|------|--------------------------------|-----|
| Aberdeen<br>Dundee   | QF- PCR    | Targeted screen         | STRs | Chromosomes 13, 18, 21,<br>X/Y | 5   |
| Edinburgh<br>Glasgow | Microarray | Whole genome CNV        |      | Whole genome                   | 14  |
| Proforma required?   |            | NO                      |      |                                |     |

#### Referral criteria

- Features suggestive of Trisomy 13, 18 or 21
- Congenital malformation/abnormalities
- Ambiguous genitalia
- Dysmorphic features
- Failure to thrive

- Neonatologists
- Clinical Genetics





# ANEUPLOIDY / MICRODUPLICATION / MICRODELETION POSTNATAL SCREENING (ROUTINE)

Available testing

| Centre               | Method     | Scope and range of test |      | Targets                     | TAT |
|----------------------|------------|-------------------------|------|-----------------------------|-----|
| Aberdeen<br>Dundee   | QF- PCR    | Targeted screen         | STRs | Chromosomes 13, 18, 21, X/Y | 28  |
| Edinburgh<br>Glasgow | Microarray | Whole genome screen     | CNV  | Whole genome                | 20  |
| Proforma required?   |            | NO                      |      |                             |     |

#### Referral criteria

- Clinical suspicion of mosaic Trisomy 13, 18 or 21
- Features of sex chromosome abnormality

- Clinical Genetics
- Obstetrics





#### **ANGELMAN SYNDROME**

#### Available testing

| Centre                | Method |                    | Scope a              | nd range of test                 | Targets                | TAT |
|-----------------------|--------|--------------------|----------------------|----------------------------------|------------------------|-----|
| Glasgow               | MLPA   | Targeted           | d screen             | CNV<br>Methylation abnormalities | 15q11-13 markers       | 28  |
| Glasgow               | Sanger | Whole              | e gene<br>een        | SNVs, indels                     | UBE3A                  | 56  |
| Glasgow               | PCR    | Targete            | Targeted screen STRs |                                  | Microsatellite markers | 28  |
| Family member testing |        | as indicated above |                      | 14                               |                        |     |
| Proforma required? NO |        |                    |                      |                                  |                        |     |

#### Referral criteria

- Clinical features that include:
  - Severe developmental delay and intellectual disability
  - o Seizures
  - Microcephaly
  - Severe speech impairment
  - o Gait ataxia and/or tremulousness of the limbs

- Clinical Genetics
- Paediatrics





#### **BECKWITH WIEDEMANN SYNDROME**

#### Available testing

|             | 3                  |         |           |                                  |                        |     |
|-------------|--------------------|---------|-----------|----------------------------------|------------------------|-----|
| Centre      | Method             |         | Scope a   | and range of test                | Targets                | TAT |
| Glasgow     | MLPA               | Targete | ed screen | CNV<br>Methylation abnormalities | 11p15 markers          | 28  |
| Glasgow     | PCR                | Targete | ed screen | STRs                             | Microsatellite markers | 28  |
| Proforma re | Proforma required? |         |           |                                  |                        | •   |

#### Referral criteria

- Clinical features that include:
  - o Macrosomia
  - o Hemihyperplasia and/or macroglossia
  - o Omphalocele (exomphalos) or umbilical hernia
  - Embryonal tumour (e.g. Wilms tumour, hepatoblastoma, neuroblastoma, rhabdomyosarcoma) in childhood

- Clinical Genetics
- Paediatrics





#### **CHARGE SYNDROME**

## Available testing

| Centre                | Method | Scope and range of test                            |                    | Targets | TAT |
|-----------------------|--------|--|--------------------|---------|-----|
| Glasgow               | Sanger | Whole gene screen SNVs, indels MLPA Exon level CNV |                    | CHD7    | 56  |
| Family member testing |        |  | as indicated above |         | 14  |
| Proforma required?    |        | NO   |                    |         |     |

#### Referral criteria

- Clinical features that include:
  - Coloboma
  - Choanal atresia or stenosis
  - Cleft palate with or without cleft lip
  - o Cranial nerve dysfunction or anomaly
  - Characteristic ear malformations
  - o Tracheoesophageal fistula or oesophageal atresia
  - o Cardiovascular malformation
  - Genital hypoplasia

## Requesting specialties





# **CONGENITAL ABNORMALITIES, MULTIPLE**

# Available testing

| Centre               | Method             | Scope and I  | range of test       | Targets      | TAT |  |
|----------------------|--------------------|--------------|---------------------|--------------|-----|--|
| Aberdeen<br>Dundee   | Microarray         | Whole genome | Structural variants | Whole genome | 14  |  |
| Edinburgh<br>Glasgow | Karyotype          | screen       | CNV                 | S            |     |  |
| Proforma require     | Proforma required? |              | NO                  |              |     |  |

#### Referral criteria

• Multiple congenital malformations

- Clinical Genetics
- Paediatrics





# CORNELIA DE LANGE SYNDROME (CdLS) and CdLS-LIKE DISORDERS Available testing

| Centre                | Method | Scope and range of test |              | Targets   | TAT |
|-----------------------|--------|-------------------------|--------------|---|-----|
| Edinburgh             | NGS    | Whole gene<br>screen    | SNVs, indels | NIPBL*, SMC1A, SMC3, HDAC8,<br>RAD21, ANKRD11, KMT2A,<br>AFF4, NAA10, BRD4, PUF60 | 56  |
| Family member testing |        |                         | as indicat   | ed above  | 14  |
| Proforma requir       | ed?    | NO                      |              |   |     |

<sup>\*</sup> MLPA analysis of NIPBL is available for patients presenting with "classical" CdLS. Please contact the laboratory for details.

#### Referral criteria

- Normal Karyotype or array CGH And
- developmental delay and
- clinical features suggestive of CdLS or CdLS like disorder, for example
  - synophrys and/or thick eyebrows
  - o short nose, concave nasal ridge and/or upturned nasal tip
  - o long and/or smooth philtrum
  - o thin upper lip vermillion and/or downturned corners of mouth
  - hand oligodactyly and/or adactyly
  - o congenital diaphragmatic hernia
  - o prenatal growth retardation
  - o postnatal growth retardation
  - microcephaly
  - o small hands
  - short fifth finger

hirsutismsee also Kline, A.D., Moss, J.F., Selicorni, A. et al. Diagnosis and management of Cornelia de Lange syndrome: first international consensus statement. Nat Rev Genet 19, 649–666 (2018). https://doi.org/10.1038/s41576-018-0031-0Requesting specialties





#### **DEVELOPMENTAL DELAY**

## Available testing

| Centre                                     | Method     | Scope and range of test |     | Targets      | TAT |
|--|------------|-------------------------|-----|--------------|-----|
| Aberdeen<br>Dundee<br>Edinburgh<br>Glasgow | Microarray | Whole gene screen       | CNV | Whole genome | 28  |
| Proforma required?                         |            | NO                      |     |              |     |

#### Referral criteria

- Significant delay in one or more of the following developmental areas
  - o Gross motor
  - Vision and fine motor
  - o Hearing, speech and language
  - o Social, emotional and behavioural

- Clinical Genetics
- Paediatrics
- Psychiatrists for Adults with Learning Disability





#### **DEVELOPMENTAL DISORDERS**

## Available testing

| Centre      | Method                | Scope and range of test |                 | Targets            | TAT   |          |
|-------------|-----------------------|-------------------------|-----------------|--------------------|---|----------|
| Edinburgh   | NGS                   | Who                     | ole gene screen | SNVs, indels       | DDG2P*  | 112      |
| Family m    | Family member testing |                         |                 | as indicated above | Э   | 14       |
| Proforma re | equired?              | NO                      |                 |                    | HPO terms and growth parthe referral documents. | ameters. |

<sup>\*</sup> Gene list available at https://www.ebi.ac.uk/gene2phenotype. Only genes in the moderate, strong, and definitive categories are analysed.

#### Referral criteria

- Severe neurodevelopmental disorder and
  - o congenital anomalies, or
  - o abnormal growth parameters, or
  - o dysmorphic features, or
  - o unusual behavioural phenotype.
- Local clinical genetics departmental MDT has assessed suitability for this test.
- Microarray analysis has previously been performed.
- Samples from proband and both parents are required (trio)

## Requesting specialties





# Di GEORGE (22q11 DELETION) SYNDROME

## Available testing

| Centre                                     | Method               | Scope and range of test |     | Targets                   | TAT   |
|--|----------------------|-------------------------|-----|---------------------------|---|
| Aberdeen<br>Dundee<br>Edinburgh<br>Glasgow | Microarray /<br>MLPA | Targeted screen         | CNV | Di George Critical Region | 28 Urgent 5* Prenatal 14  *Contact lab to discuss TAT |
| Proforma required?                         |                      | NO                      |     |                           | •   |

#### Referral criteria

- Heart abnormalities detected on ultrasound scan OR
- Congenital heart defect consistent with Di George syndrome (e.g. ventricular septal defect, tetralogy of Fallot, interrupted aortic arch or truncus arteriosus).
- Palatal anomalies (e.g. velopharyngeal incompetence, submucous cleft palate or bifid uvula).

- Cardiology
- Clinical Genetics
- Obstetrics
- Paediatrics





# **DISORDERS OF SEXUAL DEVELOPMENT (DSD)**

# Available testing

| Centre                        | Method     | Scope and ran        | ge of test                             | Targets   | TAT |
|-------------------------------|------------|----------------------|--|---|-----|
| Aberdeen<br>Dundee<br>Glasgow | Microarray | Whole genome screen  | CNV                                    | Whole genome  | 28  |
| Glasgow                       | NGS        | Whole gene<br>screen | SNVs,<br>indels                        | AMH, AMHR2, ANOS1, AR, ARX, ATRX, CBX2, CHD7, CUL4B, CYB5A, CYP11A1, CYP11B1, CYP17A1, CYP19A1, DHCR7, DHH, DMRT1, FEZF1, FGF8, FGFR1, FOXL2, FSHB, GATA4, GNRH1, GNRHR, HSD17B3, HSD3B2, INSL3, KISS1R, LHB, LHCGR, MAMLD1, MAP3K1, NR0B1, NR3C1, NR5A1, POR, PROK2, PROK2, RSPO1, RXFP2, SEMA3E, SOX2, SOX3, SOX9, SOX10, SPRY4, SRD5A2, SRY, STAR, TAC3, TACR3, TSPYL1, WDR11, WNT4, WT1 | 112 |
| Family member testing         |            | as ind               | icated above (Glasgow)                 | 14  |     |
| Proforma red                  | quired?    | YES DSD refer        | DSD referral form (see centre website) |   |     |

# Referral criteria

- Ambiguous genitalia and/or impalpable gonads at birth OR
- Delayed puberty in adolescence
- No chromosomal abnormalities detected by karyotype analysis.

- Clinical Genetics
- Endocrinology
- Paediatrics





#### **FRAGILE X**

# Available testing

| Centre                           | Method         | Scope ar        | nd range of test         | Targets | TAT |
|----------------------------------|----------------|-----------------|--------------------------|---------|-----|
| Aberdeen<br>Edinburgh<br>Glasgow | PCR &<br>TPPCR | Targeted screen | Triplet repeat expansion | FMR1    | 28  |
| Proforma required?               |                | NO              |                          |         |     |

## Referral criteria

- Clinical features characteristic of fragile X syndrome or other FMR1-related disorder
  - Typical fragile X syndrome manifestations in females: learning difficulty (usually mild, IQ often 80-85, but can be moderate or severe LD)
  - Typical fragile X syndrome manifestations in males: moderate to severe developmental delay / learning difficulty (IQ if measured would be 35-70)

Family history of Fragile X Requesting specialties

- Clinical Genetics
- Paediatrics





# **INFERTILITY, MALES**

# Available testing

| Centre                                     | Method    | Scope and           | range of test               | Targets                         | TAT |
|--|-----------|---------------------|-----------------------------|---------------------------------|-----|
| Aberdeen<br>Dundee<br>Edinburgh<br>Glasgow | Karyotype | Whole genome screen | Structural variants,<br>CNV | Whole genome                    | 28  |
| Aberdeen Dundee Edinburgh Glasgow          | ARMS      | Targeted screen     | SNVs, indels                | Common CFTR pathogenic variants | 28  |
| Dundee<br>Edinburgh<br>Glasgow             | PCR       | Targeted screen     | Y chromosome<br>markers     | AZFa, AZFb, AZFc                | 28  |
| Proforma required?                         |           | NO                  |                             |                                 |     |

#### Referral criteria

- Karyotype Patients with unexplained infertility who are going to undergo infertility treatment
- Y Chromosome microdeletions Patients with non-obstructive azoospermia or severe oligospermia where testicular sperm extraction (TESE)/microdissection TESE (mTESE) is considered and outcome of testing will inform eligibility for (m)TESE and success of sperm retrieval
- Cystic Fibrosis Male infertility associated with obstructive azoospermia, AND
  - CBAVD (or isolated CUAVD) diagnosed from expert clinical examination, OR
  - CBAVD identified at incidental herniotomy

- Clinical Genetics
- Fertility specialist





# **INFERTILITY, FEMALES**

# Available testing

| Centre                                     | Method    | Scope an            | d range of test             | Targets      | TAT |
|--|-----------|---------------------|-----------------------------|--------------|-----|
| Aberdeen<br>Dundee<br>Edinburgh<br>Glasgow | Karyotype | Whole genome screen | Structural variants,<br>CNV | Whole genome | 28  |
| Aberdeen<br>Edinburgh<br>Glasgow           | PCR       | Targeted screen     | Triplet repeat expansion    | FMR1         | 28  |
| Proforma required?                         |           | NO                  |                             |              |     |

#### Referral criteria

- Four consecutive months of unexplained amenorrhoea (primary or secondary), AND
- Elevated serum FSH of >30IU/L on two separate occasions at least 6 weeks apart, AND
- Age of onset is <30 years, AND
- Non genetic causes have been excluded including presence of thyroid and adrenal auto-antibodies

- Clinical Genetics
- Fertility specialist





#### KLINEFELTER SYNDROME

# Available testing

| Centre             | Method     | Scope and range of test |      | Targets      | TAT |  |
|--------------------|------------|-------------------------|------|--------------|-----|--|
| Aberdeen<br>Dundee | Karyotype  | Whole genome            | CNV  | Whole geneme | 20  |  |
| Glasgow            | Microarray | screen                  | CINV | Whole genome | 28  |  |
| Edinburgh          | Karyotype  | Whole genome screen     | CNV  | Whole genome | 28  |  |
| Proforma required? |            | NO                      |      |              |     |  |

#### Referral criteria

- Primary hypogonadism
- Cryptorchidism
- Gynaecomastia
- Infertility

- Clinical Genetics
- Endocrinology
- Fertility clinics





# **MICRODELETION / MICRODUPLICATION SYNDROMES**

## Available testing

| Centre                                     | Method     | Scope and range of test |     | Targets      | TAT |
|--|------------|-------------------------|-----|--------------|-----|
| Aberdeen<br>Dundee<br>Edinburgh<br>Glasgow | Microarray | Whole genome<br>screen  | CNV | Whole genome | 28  |
| Proforma required?                         |            | NO                      |     |              |     |

#### Referral criteria

 Clinical suspicion of a common microdeletion syndrome to include but not restricted to:

1p36 deletion syndrome, Wolf-Hirschhorn syndrome, Cri-du-Chat syndrome, Sotos syndrome, Saethre-Chotzen syndrome, Williams-Beuren syndrome, Williams-Beuren duplication syndrome, Langer-Giedion syndrome, Rubinstein-Taybi syndrome, Miller-Dieker syndrome, Smith-Magenis syndrome.

- Clinical Genetics
- Paediatrics





#### **HYDATIDIFORM MOLE**

#### Available testing

| Centre                         | Method | Scope and range of test |      | Targets                            | TAT |
|--------------------------------|--------|-------------------------|------|------------------------------------|-----|
| Dundee<br>Edinburgh<br>Glasgow | QFPCR  | Targeted screen         | STRs | Chromosome markers 13, 18, 21, X/Y | 28  |
| Dundee                         | FISH   | Targeted screen         | CNV  | CEPX, Y and 12 markers             | 2   |
| Proforma required? NO          |        |                         |      |                                    |     |

#### Referral criteria

- Hydatidiform Mole may be suspected during routine booking scan, or at emergency presentation in clinic. In the majority of cases, hydatidiform Moleis suspected after pathological analysis of products of conception (POC), initially reviewed at local regional pathology departments.
- Pathological suspicion of Hydatidiform Mole prompts referral to the Hydatidiform Mole Follow-Up Service (HMFUS), based within Ninewells Hospital, Dundee. HMFUS provides a national service for all women in Scotland.
- Diagnosis of a hydatidiform Moleis achieved by MDT which includes gynaecology, pathology and genetics, coordinated via HMFUS.
- For more information visit <a href="https://www.nss.nhs.scot/specialist-healthcare/specialist-services/hydatidiform-mole/">https://www.nss.nhs.scot/specialist-healthcare/specialist-services/hydatidiform-mole/</a>

#### Genetic testing in isolation

- In some cases, a complete homozygous mole can be identified solely by genotyping using QF-PCR in the absence of any parental samples. If maternal samples are provided, further molar genotypes such as complete heterozygous complements, associated with a complete hydatidiform mole, and diandric triploidy associated with partial moles, may be identified. Mosaic and chimeric moles may be harder to interpret.
- FISH testing will not differentially distinguish between normal diploid pregnancies and diandric diploid complements, associated with complete moles. Or differentiate between diandric and dygnic triploidy and therefore should not be offered as a sole test for diagnosis of molar pregnancy.

# Requesting specialties

Pathology





#### **PRADER-WILLI SYNDROME**

# Available testing

| Centre       | Method | Scope           | and range of test                | Targets                | TAT |
|--------------|--------|-----------------|----------------------------------|------------------------|-----|
| Glasgow      | MLPA   | Targeted screen | CNV<br>Methylation abnormalities | 15q11-13 region        | 28  |
| Glasgow      | PCR    | Targeted screen | STRs                             | Microsatellite markers | 28  |
| Proforma rec | uired? | NO              |                                  |                        |     |

#### Referral criteria

- Clinical features that include:
  - o Severe hypotonia and/or feeding difficulties in early infancy
  - Global developmental delay
  - Hypogonadism
  - o Excessive eating with central obesity if uncontrolled in childhood

- Clinical Genetics
- Paediatrics





#### RECURRENT MISCARRIAGE

# Available testing

| Centre               | Method     | Scope and range of test |      | Targets                     | TAT |
|----------------------|------------|-------------------------|------|-----------------------------|-----|
| Aberdeen<br>Dundee   | QF- PCR    | Targeted screen         | STRs | Chromosomes 13, 18, 21, X/Y | 28  |
| Edinburgh<br>Glasgow | Microarray | Whole genome screen     | CNV  | Whole genome                | 20  |
| Proforma required?   |            | NO                      |      |                             |     |

# Referral criteria

• Tissue from 3<sup>rd</sup> or subsequent consecutive miscarriage

- Fetal Medicine
- Pathology
- Gynaecology





#### SILVER-RUSSELL SYNDROME

# Available testing

| Centre                | Method | Scope and range of test |                                  | Targets                | TAT |
|-----------------------|--------|-------------------------|----------------------------------|------------------------|-----|
| Glasgow               | MLPA   | Targeted screen         | CNV<br>Methylation abnormalities | 11p15 region           | 28  |
| Glasgow               | PCR    | Targeted screen         | STRs                             | Microsatellite markers | 28  |
| Proforma required? NO |        | NO                      |                                  |                        |     |

#### Referral criteria

- Clinical features that include:
  - o Postnatal growth failure
  - o Small for gestational age
  - Characteristic facies
  - Limb asymmetry
  - o Feeding difficulties

- Clinical Genetics
- Paediatrics





#### **SMITH-LEMLI-OPITZ**

# Available testing

| Centre      | Method         | Scope and range of test |                    | Targets | TAT |  |
|-------------|----------------|-------------------------|--------------------|---------|-----|--|
| Glasgow     | Sanger         | Whole gene screen       | SNVs, indels       | DHCR7   | 56  |  |
|             | member<br>ting |                         | as indicated above |         |     |  |
| Proforma re | equired?       | NO                      |                    |         |     |  |

#### Referral criteria

- Clinical features that include:
  - o Prenatal and postnatal growth restriction
  - Microcephaly
  - Moderate-to-severe intellectual disability
  - Malformations that may include distinctive facial features, cleft palate, cardiac defects, underdeveloped external genitalia in males, postaxial polydactyly, syndactyly of the toes
  - Elevated serum concentration of 7-dehydrocholesterol (7-DHC)

# Requesting specialties

Clinical Genetics





# **UNIPARENTAL DISOMY, CHROMOSOME 14**

# Available testing

| Centre             | Method | Scope and range of test |      | Targets                              | TAT              |
|--------------------|--------|-------------------------|------|--------------------------------------|------------------|
| Glasgow            | PCR    | Targeted screen         | STRs | Chromosome 14 microsatellite markers | 28<br>Prenatal 3 |
| Proforma required? |        | NO                      |      |                                      |                  |

#### Referral criteria

- Prenatal testing is available for:
  - o Balanced carriers of Robertsonian translocations
  - Fetuses with a familial or de novo balanced Robertsonian translocation that contains chromosome 14
  - Fetuses with a normal karyotype where a parent is a carrier of a Robertsonian translocation that contains chromosome 14
- Postnatal testing is available in patients with a clinical suspicion of maternal uniparental disomy of chromosome 14.

- Clinical Genetics
- Paediatrics





#### X-INACTIVATION STUDIES

# Available testing

| Centre             | Method | Scope and range of te | st                   | Targets                     | TAT |
|--------------------|--------|-----------------------|----------------------|-----------------------------|-----|
| Glasgow            | PCR    | Targeted screen       | Methylation analysis | AR CAG trinucleotide repeat | 28  |
| Proforma required? |        | NO                    |                      |                             |     |

# Referral criteria

• Possible manifesting carrier of an X-linked recessive condition

- Clinical Genetics
- Paediatrics





# **ENDOCRINOLOGY**

# ALBRIGHT'S HEREDITARY, PSEUDOHYPOPARATHYROIDISM / PSEUDOPSEUDOHYPOPARATHYROIDISM

#### Available testing

| Centre                                  | Method                     |                          | Scope and r   | ange of test | Targets | TAT |
|---|----------------------------|--------------------------|---------------|--------------|---------|-----|
| Dundee                                  | NGS<br>(targeted<br>panel) | Whol                     | e gene screen | SNVs, indels | GNAS    | 56  |
| Family member testing                   |                            | as indicated ab          | ove           | 14           |         |     |
| Proforma required? YES Endocrine disord |                            | ders proforma (see centr | e website)    |              |         |     |

#### Referral criteria

- Individuals with a clear clinical diagnosis of Albright hereditary osteodystrophy, pseudohypoparathyroidism or pseudopseudohypoparathyroidism based on clinical and biochemical assessment
- Note: Imprinting defects and large deletions are not tested for.

- Clinical Genetics
- Endocrinology





#### ANDROGEN INSENSITIVITY SYNDROME

# Available testing

| Centre    | Method        | Scope and range of test |                                | Targets | TAT |
|-----------|---------------|-------------------------|--------------------------------|---------|-----|
| Glasgow   | NGS           | Whole gene screen       | Whole gene screen SNVs, indels |         | 56  |
| Family me | ember testing |                         | as indicated abo               | ove     | 14  |
| Proform   | a required?   | NO                      |                                |         |     |

#### Referral criteria

- Undermasculinisation of external genitalia at birth OR
- Abnormal secondary sexual development in puberty OR
- Infertility in individuals with a 46,XY karyotype.

# Requesting specialties

- Clinical Genetics
- Endocrinology

#### **ASYMPTOMATIC FASTING HYPERGLYCAEMIA**

## Available testing

| Centre                | Method           | Scope and range of test |                              | Targets | TAT |
|-----------------------|------------------|-------------------------|------------------------------|---------|-----|
| Dundee                | Sanger or<br>NGS | Whole gene screen       | SNVs, indels, exon level CNV | GCK     | 56  |
| Family member testing |                  |                         | as indicated at              | pove    | 14  |
| Proforma requ         | ired?            | NO                      |                              |         |     |

#### Referral criteria

Asymptomatic fasting hyperglycaemia: fasting glucose 5.5-8mmols/L

- Clinical Genetics
- Endocrinology
- Obstetrics
- Paediatrics





#### **CARNEY COMPLEX**

#### Available testing

| Centre             | Method                     | \$    | Scope and rang | ge of test                         | Targets       | TAT |
|--------------------|----------------------------|-------|----------------|------------------------------------|---------------|-----|
| Dundee             | NGS<br>(targeted<br>panel) | Whole | e gene screen  | SNVs,<br>indels, exon<br>level CNV | PRKAR1A       | 56  |
| Family men         | nber testing               |       |                | as indica                          | ated above    | 14  |
| Proforma required? |                            | YES   | Endocrine disc | orders proforma                    | (see website) |     |

#### Referral criteria

 Two or more of the features from the list below (with histological confirmation where relevant)

OR

- One feature from the list below (with histological confirmation where relevant) and an affected first degree relative:
  - Spotty skin pigmentation with typical distribution (lips, conjunctiva, vaginal and penile mucosa)
  - Myxoma (cutaneous and mucosal)
  - Cardiac myxomas
  - o Breast myxomatosis or fat-suppressed MRI suggestive of this finding
  - PPNAD or paradoxical positive response of urinary glucocorticosteroid excretion to dexamethasone administration during Liddles test
  - Acromegaly due to GH-producing adenoma
  - Large cell calcifying Sertoli cell tumour (LDDST) or characteristic calcification on testicular ultrasound
  - Thyroid carcinoma or multiple, hypoechoic nodules on thyroid ultrasound in a young patient
  - Psammomatous melanotic schwannomas (PMS)
  - o Blue nevus, epithelioid blue nevus
  - o Breast ductal adenoma
  - Osteochondromyxoma

#### Requesting specialties

- Clinical Genetics
- Dermatology
- Endocrinology

#### **CONGENITAL HYPERINSULINISM**

#### Available testing

| 4 | Centre | Method Scope and range of test |  | Targets | TAT |
|---|--------|--------------------------------|--|---------|-----|
| ۹ |        | 7 7                            |  |         |     |





| Dundee    | NGS<br>(clinical<br>exome) | Whole<br>gene<br>screen | SNVs, indels<br>exon level CNV<br>(selected genes) | ABCC8, AKT2, CACNA1D, GCK,<br>GLUD1, GPC3, HADH, HNF1A,<br>HNF4A, INSR, KCNJ11, KDM6A,<br>KMT2D, PMM2, SLC16A1, TRMT10A | 112 |
|-----------|----------------------------|-------------------------|--|---|-----|
| Family me | mber testing               |                         | as inc   | licated above   | 14  |
| Proforma  | a required?                | NO                      |  |   |     |

#### Referral criteria

- Hypoglycaemia accompanied by one of the following, with no identifiable cause:
  - During an episode of hypoglycaemia there is a requirement for the glucose infusion to be at a rate of >8mg/kg/min, OR
  - Detectable serum insulin or c-peptide when the blood glucose is <3mmol/l, OR</li>
  - o Suppressed or undetectable serum fatty acids and ketone bodies
- Urgent neonatal requests can be accommodated. Please contact the laboratory to discuss. Where possible, clinicians are asked to submit samples from the probands parents for the DNA to be stored to allow prompt follow-up of variants

- Clinical Genetics
- Endocrinology
- Paediatrics





#### **CONGENITAL HYPOTHYROIDISM**

# Available testing

| Centre                | Method                     | Scope and range of test |                 | Targets   | TAT |
|-----------------------|----------------------------|-------------------------|-----------------|---|-----|
| Dundee                | NGS<br>(clinical<br>exome) | Whole gene<br>screen    | SNVs,<br>indels | DUOX2, DUOXA2, FOXE1, GLIS3,<br>GNAS, HESX1, IGSF1, IRS4, IYD, LHX3,<br>LHX4 NKX2-1, OTX2, PAX8, POU1F1,<br>PRKAR1A, PROP1, SECISBP2,<br>SLC16A2, SLC26A4, SLC5A5, TBL1X,<br>TG, THRB, THRA, TRHR, TPO, TSHR,<br>TSHB | 112 |
| Family member testing |                            |                         | as in           | dicated above   | 14  |
| tes                   | ung                        |                         |                 |   |     |
| Proforma              | required?                  | NO                      |                 |   |     |

#### Referral criteria

- Congenital hypothyroidism, thyroid hypoplasia or agenesis with or without syndromic features, OR
- Thyroid dyshormonogenesis, OR
- Raised serum thyroid stimulating hormone (TSH) level:
  - o With enlarged thyroid gland, OR
  - o In the absence of thyroid autoantibodies

- Clinical Genetics
- Endocrinology





#### **CONGENITAL NEPHROGENIC DIABETES INSIPIDUS**

#### Available testing

| Centre                | Method | Scope and range of test |              | Targets      | TAT |
|-----------------------|--------|-------------------------|--------------|--------------|-----|
| Dundee                | Sanger | Whole gene screen       | SNVs, indels | AQP2, AVPR2  | 56  |
| Family member testing |        |                         | as ind       | icated above | 14  |
| Proforma required?    |        | NO                      |              |              |     |

#### Referral criteria

Any individual with a clinical presentation consistent with the condition.

# Requesting specialties

- Clinical Genetics
- Endocrinology

#### **CONGENITAL OVERGROWTH DISORDERS**

# Available testing

| Centre     | Method                     | Scope and r          | ange of test | Targets   | TAT |
|------------|----------------------------|----------------------|--------------|---|-----|
| Dundee     | NGS<br>(clinical<br>exome) | Whole gene<br>screen | SNVs, indels | AKT2, BRWD3, CDKN1C, CHD8,<br>DIS3L2, DNMT3A, EZH2, GPC3,<br>MTOR, NFIB, NFIX, NSD1, OFD1,<br>PDGFRB, PIK3CA, PTEN, RNF125,<br>SETD2, SUZ12 | 112 |
| ,          | member<br>ting             |                      | as indicat   | ed above  | 14  |
| Proforma r | equired?                   | NO                   |              |   |     |

#### Referral criteria

- Any individual with clinical features suggestive of:
  - Atypical Beckwith-Wiedemann syndrome, Classical Beckwith-Wiedemann syndrome, Simpson-Golabi-Behmel syndrome, Sotos syndrome, Weaver syndrome
- Overlapping investigations: Beckwith-Wiedemann syndrome, Microdeletion/Microduplication Syndromes

# Requesting specialties

Clinical Genetics





#### **FAMILIAL HYPERPARATHYROIDISM**

#### Available testing

| Centre             | Method                     | Scope and               | range of test   | Targets  | TAT |
|--------------------|----------------------------|-------------------------|---|--|-----|
| Dundee             | NGS<br>(targeted<br>panel) | Whole<br>gene<br>screen | SNVs,<br>indels, exon<br>level CNV<br>(selected<br>genes) | AP2S1, CASR, CDC73, CDKN1B, GCM2,<br>GNA11, MEN1, RET (exons 5, 8, 10, 11,<br>13-16) | 56  |
| Family me          | mber testing               |                         | â   | as indicated above   | 14  |
| Proforma required? |                            | YES En                  | docrine disorder  | s proforma (see centre website)  | •   |

#### Referral criteria

- Primary hyperparathyroidism (unexplained hypercalcaemia with PTH high or in the upper normal range, and calcium clearance: creatinine clearance ratio > 0.02) which meets ONE of the criteria below:
  - o Presenting before the age of 35, OR
  - Presenting at any age with ONE of:
    - Proven multi-glandular involvement, OR
    - Hyperplasia on histology, OR
    - Ossifying fibroma(s) of the maxilla and / or mandible, OR
    - At least one first degree relative with unexplained hyperparathyroidism
- Testing in other contexts e.g. where age of onset is not clear or with a later onset but strong family history is also appropriate.
- Overlapping indications:
  - Familial Hypocalciuric hypercalcaemia test should be used where there is hypercalcaemia (and inappropriately normal or raised PTH) with hypocalciuria (calcium clearance: creatinine clearance ratio < 0.02)</li>
  - Multiple Endorine Neoplasia Type 1 & Type 4
  - Multiple Endocrine Neoplasia Type 2A
  - O Hyperparathyroidism-Jaw Tumour Syndrome/Parathyroid carcinoma

- Clinical Genetics
- Endocrinology





#### FAMILIAL HYPOCALCIURIC HYPERCALCAEMIA

#### Available testing

| Centre             | Method               | Scope and range of test |          |  | Targets  | TAT |
|--------------------|----------------------|-------------------------|----------|--|--|-----|
| Dundee             | NGS(targeted panel)  | Whole                   | •        | SNVs, indels,<br>exon level<br>CNV (selected<br>genes) | AP2S1, CASR, CDC73, CDKN1B,<br>GCM2, GNA11, MEN1, RET (exons 5, 8,<br>10, 11, 13-16) | 56  |
| Family m           | amily member testing |                         |          | as indi  | cated above  | 14  |
| Proforma required? |                      | YES                     | Endocrin | ne disorders profo                                     | rma (see centre website)   |     |

#### Referral criteria

- Individuals with hypercalcaemia with hypocalciuria (calcium clearance: creatinine clearance ratio < 0.02), with normal and/or elevated PTH
- Overlapping indications:
  - Familial hyperparathyroidism test should be used for hypercalcaemia (with normal or raised PTH) with calcium clearance: creatinine clearance ratio > 0.02 in the presence of an appropriate clinical indication (see Familial Hyperparathyoridism panel)
- Note that the same gene panel is used for FHH and Familial Hyperparathyroidism referrals.

- Clinical Biochemistry
- Clinical Genetics
- Endocrinology
- Nephrology





#### FAMILIAL HYPOPARATHYROIDISM

# Available testing

| Centre             | Method                     | Scope and range of test |  | ange of test  | Targets                                      | TAT |
|--------------------|----------------------------|-------------------------|--|---|--|-----|
| Dundee             | NGS<br>(targeted<br>panel) | Whole<br>scre           | gene<br>een  | SNVs,<br>indels, exon<br>level CNV<br>(selected<br>genes) | AIRE, CASR, GATA3, GCM2, GNA11, PTH,<br>TBCE | 56  |
| Family me          | mber testing               |                         | as   |   | indicated above                              | 14  |
| Proforma required? |                            | YES                     | ES Endocrine disorders proforma (see centre website) |   |  |     |

#### Referral criteria

- Individuals with non-syndromic hypoparathyroidism with low calcium levels and low or inappropriately normal serum PTH, with no detectable cause.
- Any individual with clinical features suggestive of an AIRE disorder.
- Testing of patients who are normocalcaemic may occasionally be appropriate after consultation with an expert in calcium homeostasis

- Clinical Genetics
- Endocrinology





#### FAMILIAL ISOLATED PITUITARY ADENOMA

# Available testing

| Centre      | Method                     | Scope and range of test |               |  | Targets                 | TAT |
|-------------|----------------------------|-------------------------|---------------|--|-------------------------|-----|
| Dundee      | NGS<br>(targeted<br>panel) |                         | e gene<br>een | SNVs, indels,<br>exon level<br>CNV (selected<br>genes) | MEN1, CDKN1B, AIP       | 56  |
| ,           | member<br>ting             |                         |               | as ind   | icated above            | 14  |
| Proforma re | equired?                   | YES                     | Endocrin      | e disorders proforn                                    | na (see centre website) |     |

#### Referral criteria

- Individuals with one of the following:
  - Any pituitary adenoma <20 years</li>
  - o Any pituitary macroadenoma <30 years of age
  - Isolated pituitary adenoma developing under the age of 35, with at least one first degree relative with an isolated pituitary adenoma
- Overlapping clinical indications:
  - Multiple Endocrine Neoplasia Type 1 & Type 4 (included in this panel MEN1 and CDKN1B genes)

- Clinical Genetics
- Endocrinology





#### FAMILIAL NEUROHYPOPHYSEAL DIABETES INSIPIDUS

# Available testing

| Centre             | Method                   | Scope and         | d range of test | Targets | TAT |
|--------------------|--------------------------|-------------------|-----------------|---------|-----|
| Dundee             | Sanger                   | Whole gene screen | SNVs, indels    | AVP     | 56  |
| Family me          | member testing as indica |                   | ed above        | 14      |     |
| Proforma required? |                          | NO                |                 |         |     |

# Referral criteria

• Any individual with a clinical presentation consistent with the condition.

# Requesting specialties

- Clinical Genetics
- Endocrinology

# **GLUCOCORTICOID REMEDIABLE ALDOSTERONISM (GRA)**

# Available testing

| Centre      | Method   | Scope and range of test |             | Targets                                | TAT |
|-------------|----------|-------------------------|-------------|--|-----|
| Aberdeen    | PCR      | Targeted screen         | Fusion gene | CYP11B1, CYP11B2 fusion gene detection | 28  |
| Proforma re | equired? | NO                      |             |  |     |

#### Referral criteria

• Hypertension presenting in childhood to early adulthood

- Clinical Genetics
- Endocrinology





# HYPERPARATHYROIDISM-JAW TUMOUR SYNDROME / INHERITED PARATHYROID CARCINOMA

# Available testing

| Centre   | Method                     | Scope and range of test |               |                                | Targets        | TAT |
|----------|----------------------------|-------------------------|---------------|--------------------------------|----------------|-----|
| Dundee   | NGS<br>(targeted<br>panel) |                         | e gene<br>een | SNVs, indels<br>Exon level CNV | CDC73          | 56  |
| ,        | member<br>ting             |                         |               | as indicated a                 | above          | 14  |
| Proforma | required?                  | YES                     | Endocrine     | e disorders proforma (see co   | entre website) |     |

#### Referral criteria

- All Patients with parathyroid carcinoma
- Clinical phenotype of HPT-JT (i.e. primary hyperparathyroidism and ossifying fibroma or maxilla and mandible
- Or ≥1 HPT-JT manifestation and a first degree relative with ≥1 HPT-JT manifestation
- HPT-JT manifestations include primary hyperparathyroidism (including parathyroid adenoma and carcinoma) and ossifying fibroma of the mandible and maxilla

- Clinical Genetics
- Endocrinology





#### **HYPERTHYROIDISM**

# Available testing

| Centre             | Method                     | Scope and rang    | ge of test   | Targets  | TAT |
|--------------------|----------------------------|-------------------|--------------|--|-----|
| Dundee             | NGS<br>(clinical<br>exome) | Whole gene screen | SNVs, indels | ALB, SECISBP2, SLC16A2, THRA,<br>THRB, TSHR, TTR | 112 |
|                    | member<br>ting             |                   | as indic     | ated above                                       | 14  |
| Proforma required? |                            | NO                |              |  | •   |

#### Referral criteria

- Hyperthyroidism where common causes have been excluded:
  - Clinical exclusion of common causes such as toxic solitary nodules or multinodular goitre, AND
  - Graves disease excluded by negative TSH receptor autoantibodies when the patient is biochemically hyperthyroid, AND
  - Patient presenting below the age of 18 OR patient has a first degree relative with unexplained hyperthyroidism

- Clinical Genetics
- Endocrinology





#### HYPOGONADOTROPIC HYPOGONADISM

# Available testing

| Centre      | Method       | S     | cope and rang | ge of test      | Targets  | TAT |
|-------------|--------------|-------|---------------|-----------------|--|-----|
| Glasgow     | NGS          | Whole | gene screen   | SNVs, indels    | ANOS1, CHD7, CUL4B, FEZF1, FGF8,<br>FGFR1, FSHB, GNRH1, GNRHR,<br>KISS1R, NR0B1, PROK2, PROK2R,<br>SEMA3E, SOX2, SOX10, SPRY4,<br>TAC3, TACR3, WDR11 | 112 |
| Family me   | mber testing |       | as indi       |                 | ated above   | 14  |
| Proforma re | equired?     | YES   | Hypogonado    | trophic Hypogor | nadism referral form (see centre website)  |     |

# Referral criteria

Clinical history of Hypogonadism

# Requesting specialties

- Clinical Genetics
- Endocrinology
- Paediatrics

# **HYPOPHOSPHATEMIC RICKETS**

# Available testing

| Centre                | Method                     | Scope and range of test |              | Targets  | TAT |
|-----------------------|----------------------------|-------------------------|--------------|--|-----|
| Dundee                | NGS<br>(clinical<br>exome) | Whole gene<br>screen    | SNVs, indels | CYP27B1, CYP2R1, DMP1, ENPP1,<br>FAM20C, FGF23, PHEX, SLC34A1,<br>SLC34A3, VDR | 112 |
| Family member testing |                            |                         | as in        | dicated above  | 14  |
| Proforma re           | equired?                   | NO                      |              |  |     |

#### Referral criteria

 Hypophosphataemia with no identifiable cause, with evidence of decreased renal phosphate reabsorption, which has or could lead to presentation with rickets

- Clinical Genetics
- Endocrinology





#### **MONOGENIC DIABETES**

## Available testing

| Centre      | Method                     | Scope and range of test |   | Targets   | TAT |
|-------------|----------------------------|-------------------------|---|---|-----|
| Dundee      | NGS<br>(targeted<br>panel) | Whole<br>gene<br>screen | SNVs,<br>indels, exon<br>level CNV<br>(selected<br>genes) | ABCC8, AKT2, APPL1, CEL, CISD2, DCAF17, DNAJC3, DYRK1B, GATA4, GATA6, GCK, HNF1A, HNF4A, HNF1B, INS, INSR, KCNJ11, LMNA, NEUROD1, PAX6, PCBD1, PDX1, PIK3R1, PLIN1, POLD1, PPARG, PPP1R15B, RFX6, SLC29A3, TRMT10A, WFS1, ZBTB20, ZFP57, mitochondrial MIDD variant m.3243A>G | 112 |
| ,           | member<br>ting             |                         |   | as indicated above  | 14  |
| Proforma re | equired?                   | YES                     | Monogenic diabet  | es 33 gene NGS panel proforma (see centre website)  |     |

## Referral criteria

- Individuals meeting any one of the following criteria:
  - Minimum two generation family history of diabetes with at least one individual diagnosed under the age of 35 years with BMI less than 30, negative GAD and IA2 autoantibodies and detectable C-peptide, OR
  - High risk of Maturity onset diabetes of the young (MODY) based on MODY calculator http://www.diabetesgenes.org/content/mody-probability-calculator, OR
  - Diabetes in conjunction with cystic renal disease and/or congenital anomaly of the kidney or urinary tract (likely HNF1B), OR
  - Diabetes in conjunction with other extra-pancreatic features suggestive of monogenic diabetes. e.g. deafness, congenital heart disease, epilepsy, diabetes insipidus, developmental delay etc.
  - Post-pubertal children or adults with insulin resistance:
    - Severely elevated plasma insulin (typically greater than 150pmol/L in nondiabetic non-obese subject), AND
    - Clinical features consistent with severe insulin resistance, e.g. polycystic ovarian syndrome, acanthosis nigricans, diabetes with high insulin requirements, post-prandial hypoglycaemia, OR
    - Post-pubertal severe insulin resistance with plasma adiponectin >5mg/l, OR
  - Clinical features of lipodystrophy, including:
    - Abnormal fat distribution (with abdominal fat preservation), AND
    - Acanthosis nigricans and/or very high insulin requirement, AND
    - Impaired glucose tolerance/diabetes

- Clinical Genetics
- Endocrinology
- Paediatrics





# **MULTIPLE ENDOCRINE NEOPLASIA (TYPE 1, TYPE 4)**

#### Available testing

| Centre     | Method                     | Sco     | pe and ra                   | nge of test   | Targets                 | TAT |
|------------|----------------------------|---------|-----------------------------|---|-------------------------|-----|
| Dundee     | NGS<br>(targeted<br>panel) | Targete | d screen                    | SNVs, indels ,<br>exon level<br>CNV (selected<br>genes) | MEN1, CDKN1B, AIP       | 56  |
| _          | member<br>sting            |         |                             | as ind  | icated above            | 14  |
| Proforma r | equired?                   | YES     | Endocrine disorders proform |   | ma (see centre website) |     |

#### Referral criteria

- Testing of individual affected with endocrine abnormalities where the individual +/- family history meets one of the following criteria:
  - o Multiple endocrine neoplasia type 1 (MEN1). The proband has:
    - Parathyroid multiglandular disease (hyperplasia/ adenomas) (<35 years), OR</li>
    - Any pituitary adenoma or insulinoma (< 20years), OR</li>
    - Pituitary macroadenoma (<30 years), OR</li>
    - ≥2 MEN1-related endocrine abnormalities (any age), OR
    - ≥1 MEN1-related endocrine abnormality and ≥1 MEN1-related non-endocrine tumours (any age), OR
    - ≥1 MEN1-related endocrine abnormality and a first degree relative has ≥1 MEN1-related endocrine abnormality
- MEN1-related endocrine abnormalities include:
  - Parathyroid hyperplasia/multiglandular adenomas
  - o Pituitary tumors
  - Endocrine tumors of the gastro-entero-pancreatic (GEP) tract
  - Carcinoid tumors
  - Adrenocortical tumors
- MEN1-related non-endocrine tumours include:
  - o facial angiofibromas
  - o collagenomas
  - o meningioma
- Overlapping clinical indications:
  - Familial Hyperparathyroidism
  - Familial Pituitary Adenoma (FIPA)

## Requesting specialties

- Clinical Genetics
- Endocrinology

NSD611-003.20 V5





# MULTIPLE ENDOCRINE NEOPLASIA (TYPE 2a, TYPE 2B) AND MEDULLARY THYROID CARCINOMA

## Available testing

| Centre      | Method                     | Sco     | pe and ra | nge of test         | Targets                                  | TAT |
|-------------|----------------------------|---------|-----------|---------------------|--|-----|
| Dundee      | NGS<br>(targeted<br>panel) | Targete | d screen  | SNVs, indels        | RET (exons 5, 8, 10, 11, 13, 14, 15, 16) | 56  |
| ,           | member<br>ting             |         | as inc    |                     | icated above                             | 14  |
| Proforma re | equired?                   | YES     | Endocrin  | e disorders proforr | na (see centre website)                  |     |

#### Referral criteria

- Testing of individual (proband) affected with endocrine abnormalities where the individual +/- family history meets one of the following criteria. The proband has:
  - o MTC (any age), OR
  - ≥2 MEN2-related endocrine abnormalities (any age), OR
  - ≥1 MEN2-related endocrine abnormality and a first degree relative with
     ≥1 MEN2-related endocrine abnormality
- MEN2-related endocrine abnormalities include: Medullary Thyroid Carcinoma (MTC), Phaechromocytoma/paraganglioma, Parathyroid adenoma/hyperplasia, Hirschprungs disease
- Overlapping clinical indications:
  - Phaeochromocytoma and paraganglioma panel

- Clinical Genetics
- Endocrinology





#### PHAEOCHROMOCYTOMA AND PARAGANGLIOMA

#### Available testing

| Centre | Method                     | Scope      | and range of test                                     | Targets   | TAT |
|--------|----------------------------|------------|---|---|-----|
| Dundee | NGS<br>(targeted<br>panel) | Whole gene | e SNVs, indels<br>Exon level CNV in<br>relevant genes | SDHA, SDHB, SDHC; SDHD, SDHAF2, VHL,<br>MAX, TMEM127, RET (exons 5, 8, 10, 11, 13<br>to 16), FH | 56  |
|        | member<br>ting             |            | as  | indicated above   | 14  |
| Š .    |                            |            | Endocrine disorders prof                              | orma (see centre website)   |     |

#### Referral criteria

- Testing of individual (proband) affected with cancer where the individual +/-family history meets one of the following criteria. The proband has:
  - Unilateral phaeochromocytoma (<60 years), OR</li>
  - o Paraganglioma of the head and neck (at any age), OR
  - Sympathetic, metastatic or abdominal, thoracic, pelvic paraganglioma (any age), OR
  - o Bilateral phaeochromocytoma (any age), OR
  - o Phaeochromocytoma and renal cell carcinoma (any age), OR
  - Phaeochromocytoma / paraganglioma (any age) AND ≥1 relative (first / second / third degree relative) with phaeochromocytoma / paraganglioma / renal cell cancer (any age)
- Individuals with clinical features associated with Neurofibromatosis Type 1 can also be tested for variants in *NF1*.
- Overlapping clinical indications:
  - Multiple Endocrine Neoplasia Type 2 (tested for within this panel: RET gene)

- Clinical Genetics
- Endocrinology





## PIGMENTED NODULAR ADRENOCORTICAL DISEASE

# Available testing

| Centre                | Method                     | Scope and range of test |              | Targets                          | TAT |
|-----------------------|----------------------------|-------------------------|--------------|----------------------------------|-----|
| Dundee                | NGS<br>(clinical<br>exome) | Whole gene<br>screen    | SNVs, indels | ARMC5, PDE11A, PDE8B,<br>PRKAR1A | 112 |
| Family member testing |                            |                         | as indica    | ited above                       | 14  |
| Proforma required?    |                            | NO                      |              |                                  |     |

#### Referral criteria

- Primary pigmented nodular adrenocortical disease, OR
- Clinical diagnosis of ACTH-independent Cushing syndrome of unknown aetiology.

# Requesting specialties

- Clinical Genetics
- Endocrinology

#### PRIMARY HYPERALDOSTERONISM

# Available testing

| Centre                | Method | Scope and rai     | nge of test  | Targets      | TAT |
|-----------------------|--------|-------------------|--------------|--------------|-----|
| Aberdeen              | Sanger | Whole gene screen | SNVs, indels | KCNJ5        | 56  |
| Family member testing |        |                   | as ind       | icated above | 14  |
| Proforma required?    |        | NO                |              |              |     |

#### Referral criteria

Hypertension presenting in childhood (under 10 years of age)

- Clinical Genetics
- Endocrinology





#### **RENAL CYSTS & DIABETES**

# Available testing

| Centre                | Method                     | Scope and r        | ange of test                   | Targets | TAT |  |
|-----------------------|----------------------------|--------------------|--------------------------------|---------|-----|--|
| Dundee                | NGS<br>(targeted<br>panel) | Whole gene screen  | SNVs, indels<br>Exon level CNV | HNF1B   | 56  |  |
| Family member testing |                            | as indicated above |                                |         |     |  |
| Proforma required?    |                            | NO                 |                                |         |     |  |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition.
- Overlapping indications:
  - Monogenic Diabetes. Full Monogenic diabetes panel will be added for all patients with diabetes with or without renal cysts unless requested otherwise.
  - Cystic kidney panel note that this will be applied if the primary indication is kidney/renal cysts

- Clinical Genetics
- Endocrinology
- Fetal Medicine
- Nephrology
- Paediatrics
- Renal





#### **SEVERE EARLY ONSET OBESITY**

# Available testing

| Centre                | Method                     | Scope and range of test |                                       |              | Targets  | TAT |
|-----------------------|----------------------------|-------------------------|---------------------------------------|--------------|--|-----|
| Dundee                | NGS<br>(clinical<br>exome) | Whole gene screen       |                                       | SNVs, indels | ALMS1, ARL6, BBS1, BBS10, BBS12,<br>BBS2, BBS4, BBS5, BBS7, BBS9,<br>CEP19, GNAS, LEP, LEPR, MC4R,<br>MKKS, MKS1, MYT1L, NTRK2, PCSK1,<br>PHF6, POMC, SDCCAG8, SIM1, TTC8,<br>VPS13B | 112 |
| Family member testing |                            | as indicated above      |                                       |              |  |     |
| Proforma required?    |                            | YES                     | Obesity proforma (see centre website) |              |  |     |

# Referral criteria

- BMI >3.5 SDS
- Age of onset below 5 years
- No significant developmental delay or dysmorphic features (referral to Clinical Genetics required as other testing may be more appropriate)

- Clinical Genetics
- Endocrinology
- Obesity specialist





#### THYROID HORMONE RESISTANCE

# Available testing

| Centre                | Method                     | Scope and range of test |   |              | Targets | TAT |
|-----------------------|----------------------------|-------------------------|---|--------------|---------|-----|
| Dundee                | NGS<br>(targeted<br>panel) | Whole gene<br>screen    |   | SNVs, indels | THRB    | 56  |
| Family member testing |                            | as indicated above      |   |              |         |     |
| Proforma required?    |                            | YES                     | Endocrine disorders proforma (see centre website) |              |         |     |

# Referral criteria

• Clinical and biochemical picture consistent with thyroid hormone resistance with or without a relevant family history

- Clinical Genetics
- Endocrinology





#### **VON HIPPEL LINDAU SYNDROME**

#### Available testing

| Centre                   | Method                     | Scope and range of test |   |                                   | Targets | TAT |
|--------------------------|----------------------------|-------------------------|---|-----------------------------------|---------|-----|
| Dundee                   | NGS<br>(targeted<br>panel) | Whole gene<br>screen    |   | SNVs, indels<br>Exon level<br>CNV | VHL     | 56  |
| Family member<br>testing |                            |                         | as indicated above                                |                                   |         |     |
| Proforma required?       |                            | YES                     | Endocrine disorders proforma (see centre website) |                                   |         |     |

#### Referral criteria

- Testing of individual (proband) affected with VHL-related tumours where the individual/family history meets one of the following criteria:
  - o Retinal angioma, spinal or endolymphatic sac tumour (<40 years), OR
  - o Cerebellar haemangioblastoma (<60 years), OR
  - ≥2 VHL-related tumours (any age), OR
  - ≥1 VHL-related tumour and a first degree relative with ≥1 VHL-related tumour (where one of the tumours is retinal angioma / hemangioblastoma)
- Deceased affected individual (proband) where (i) the individual +/- family history meets one of the above criteria, (ii) appropriate tissue is available (tumour or normal), and (iii) no living affected individual is available for genetic testing
- VHL-related tumours comprise: Retinal angioma, Spinal or cerebellar hemangioblastoma, adrenal or extra-adrenal phaeochromocytoma, Renal cell carcinoma, multiple renal and/or pancreatic cysts, endolymphatic sac tumors, papillary cystadenomas of the epididymis or broad ligament, neuroendocrine tumour of the pancreas
- Overlapping clinical indications:
  - Phaeochromocytoma and paraganglioma

- Clinical Genetics
- Endocrinology
- Nephrology
- Ophthalmology
- Urology
- Neurosurgery





# **EYES**

# ABCA4 ASSOCIATED OPHTHALMIC CONDITIONS(incl. STARGARDT DISEASE, CONE-ROD DYSTROPHY, FUNDUS FLAVIMACULATUS)

# Available testing

| Centre                | Method | Scope and range of test |              | Targets | TAT |
|-----------------------|--------|-------------------------|--------------|---------|-----|
| Edinburgh             | NGS    | Whole gene screen       | SNVs, indels | ABCA4   | 56  |
| Family member testing |        |                         | as indicated | d above | 14  |
| Proforma required?    |        | NO                      |              |         |     |

#### Referral criteria

- Clinical features that indicate a likely diagnosis of *ABCA4* associated ophthalmic conditions i.e.
  - o Progressive loss of central vision
  - o Retinal flecks
  - Macular atrophysparing of peripapillary region

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





## **ALBINISM & NYSTAGMUS**

# Available testing

| Centre      | Method         | Scope and range of test |                                   | Targets  | TAT |
|-------------|----------------|-------------------------|-----------------------------------|--|-----|
| Aberdeen    | NGS            | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | AP3B1, CACNA1A, CACNA1F, CASK, FRMD7,<br>GPR143, HPS1, HPS3, HPS4, HPS5, HPS6,<br>LRMDA, LYST, OCA2, PAX6, RAB27A, SACS,<br>SETX, SLC24A5, SLC38A8, SLC45A2, TYR,<br>TYRP1 | 112 |
| ,           | member<br>ting |                         | as indicated above                |  | 14  |
| Proforma re | equired?       | NO                      |                                   |  |     |

## Referral criteria

• Clinical features suggestive of a monogenic cause of Albinism & Nystagmus

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





# **ANTERIOR SEGMENT DYSGENESIS (ASD)**

# Available testing

| Centre                | Method | Scope and range of test |                                    | Targets   | TAT |
|-----------------------|--------|-------------------------|------------------------------------|---|-----|
| Aberdeen              | NGS    | Whole gene<br>screen    | SNVs, indels,<br>Exon level<br>CNV | ADAMTS18, ALDH18A1, ATOH7, B3GLCT, BEST1, BMP7, CHRDL1, CHST6, COL4A1, COL8A2, CRYGC, CYP1B1, DCN, EYA1, FBN1, FOXC1, FOXE3, FOXL2, GJA1, GNPTG, GSN, KERA, KRT12, KRT3, LAMB2, LCAT, LMX1B, LTBP2, MYOC, NOTCH2, OPTN, PAX3, PEX2, PIKFYVE, PITX2, PITX3, PRDM5, PXDN, RAB18, RAB3GAP1, RAB3GAP2, SEC23A, SH3PXD2B, SIX3, SLC16A12, SLC4A11, SLC4A4, TACSTD2, TGFBI, UBIAD1, VSX1, WDR36, ZEB1 | 112 |
| Family member testing |        |                         | as inc                             | dicated above   | 14  |
| Proforma re           |        | NO                      |                                    |   |     |

## Referral criteria

 Clinical features suggestive of Anterior Segment Dysgenesis – glaucoma, iris hypoplasia, vascularization and opacity in the cornea, corectopia, polycoria, ectopia lentis, cataracts

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





## **BARDET-BIEDL SYNDROME**

# Available testing

| Centre      | Method                | Scope and range of test |                                 | Targets   | TAT |
|-------------|-----------------------|-------------------------|---------------------------------|---|-----|
| Aberdeen    | NGS                   | Whole<br>gene<br>screen | SNVs, indels,<br>Exon level CNV | ARL6, BBS1, BBS2, BBS4, BBS5, BBS7,<br>BBS9, BBS10, BBS12, LZTFL1, MKKS, MKS1,<br>SDCCAG8,TTC8, WDPCP | 112 |
| Family me   | nily member testing   |                         | а                               | as indicated above  | 14  |
| Proforma re | Proforma required? NO |                         |                                 |   |     |

## Referral criteria

- Clinical features suggestive of Bardet-Biedl Syndrome (≥4 primary features or 3 primary features & ≥2 secondary features
  - Primary features: Retinal dystrophy, Renal abnormalities, Obesity, Polydactyly, Learning difficulties, Hypogonadism in males
  - Secondary features: Speech disorder / delay, Strabismus / cataracts / astigmatism, Brachydactyly / syndactyly, developmental delay, Polyuria / polydipsia, Ataxia / poor coordination / imbalance

- Clinical Genetics
- Nephrology
- Ophthalmology in discussion with Clinical Genetics





# BEST DISEASE, VITELLIFORM MACULAR DYSTROPHY (VMD), AR BESTROPHINOPATHY (ARB)

## Available testing

| Centre      | Method                             | Scope and range of test |                                | Targets | TAT |
|-------------|------------------------------------|-------------------------|--------------------------------|---------|-----|
| Aberdeen    | Sanger<br>MLPA                     | Whole gene screen       | SNVs, indels<br>Exon level CNV | BEST1   | 56  |
| Family me   | Family member testing as indicated |                         |                                | above   | 14  |
| Proforma re | equired?                           | NO                      |                                |         |     |

#### Referral criteria

 Clinical features suggestive of Best disease, Vitelliform Macular dystrophy (VMD), AR bestrophinopathy - reduced vision and an early, significant reduction in electro-oculogram (EOG) light rise

# Requesting specialties

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

# **BLEPHAROPHIMOSIS, PTOSIS, AND EPICANTHUS INVERSUS (BPES)**

## Available testing

| Centre      | Method         | Scope and range of test |                                    | Targets | TAT |
|-------------|----------------|-------------------------|------------------------------------|---------|-----|
| Aberdeen    | Sanger<br>MLPA | Whole gene screen       | SNVs, indels<br>Exon level<br>CNVs | FOXL2   | 56  |
| _           | member<br>ting |                         | as indicated                       | d above | 14  |
| Proforma re | equired?       | NO                      |                                    |         |     |

#### Referral criteria

 Clinical features suggestive of BPES – blepharophimosis, ptosis and epicanthus inversus either with premature ovarian failure (BPES type 1) or without (BPED type II).

## Requesting specialties

Clinical Genetics

Ophthalmology in discussion with

#### **BRITTLE CORNEA SYNDROME**

# Available testing

|   | Centre | Method | Scope and range of test | Targets | TAT |
|---|--------|--------|-------------------------|---------|-----|
| П |        |        |                         |         |     |





| Aberdeen           | NGS          | Whole gene screen | SNVs, indels | PRDM5, ZNF469 | 56 |
|--------------------|--------------|-------------------|--------------|---------------|----|
| Family me          | mber testing |                   | as indica    | ated above    | 14 |
| Proforma required? |              | NO                |              |               |    |

## Referral criteria

 Clinical features suggestive of isolated Brittle Cornea Syndrome (can also be a feature in Ehlers-Danlos Syndrome, see Connective Tissue Disorders) – Thinning of the cornea, myopia, blue sclera, retinal detachment

# Requesting specialties

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

## **CHOROIDERAEMIA**

## Available testing

| Centre                | Method   | Scope and range of test |                                    | Targets | TAT |
|-----------------------|----------|-------------------------|------------------------------------|---------|-----|
| Aberdeen              | Sanger   | Whole gene screen       | SNVs, indels<br>Exon level<br>CNVs | СНМ     | 56  |
| Family member testing |          |                         | as indicated                       | d above | 14  |
| Proforma re           | equired? | NO                      |                                    |         |     |

#### Referral criteria

• Clinical features suggestive of Choroideraemia - consistent with X-linked ocular disorder, degeneration of choriocapillaris, retinal pigment epithelium and retinal photoreceptor

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





## **CONGENITAL CATARACTS**

# Available testing

| Centre                | Method                | Scope and               | I range of test                    | Targets  | TAT |
|-----------------------|-----------------------|-------------------------|------------------------------------|--|-----|
| Aberdeen              | NGS                   | Whole<br>gene<br>screen | SNVs, indels,<br>Exon level<br>CNV | ADAMTS10, AGK, AGPS, ALDH18A1, B3GLCT, BCOR, BFSP1, BFSP2, CHMP4B, COL11A1, COL18A1, COL2A1, COL4A1, CRYAA, CRYAB, CRYBA1, CRYBA4, CRYBB1, CRYBB2, CRYBB3, CRYGC, CRYGD, CRYGS, CYP27A1, CYP51A1, DHCR7, DNMBP, EED, EIF2B2, EPHA2, ERCC2, ERCC3, ERCC6, ERCC8, FAM126A, FOXE3, FTL, FYCO1, GALK1, GALT, GCNT2, GEMIN4, GJA3, GJA8, GNPAT, GTF2H5, HMX1, HSF4, HTRA2, INPP5K, JAM3, LCAT, LIM2, LONP1, LSS, MAF, MAN2B1, MIP, MSMO1, MYH9, NDP, NF2, NHS, OCRL, OPA3, P3H2, PAX6, PEX1, PEX10, PEX11B, PEX12, PEX13, PEX14, PEX16, PEX19, PEX2, PEX26, PEX3, PEX5, PEX6, PEX7, PITX3, PXDN, RAB18, RAB3GAP1, RAB3GAP2, SC5D, SIL1, SLC2A1, SLC33A1, SRD5A3, TDRD7, TFAP2A, VIM, VSX2, WFS1, WRN, XYLT2 | 112 |
| ,                     | Family member testing |                         |                                    | as indicated above   | 14  |
| Proforma required? NO |                       |                         |                                    |  |     |

## Referral criteria

• Clinical features suggestive of a monogenic cause of congenital cataracts

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





# **CORNEAL ABNORMALITIES (incl. CORNEAL DYSTROPHY & BCS)**

# Available testing

| Centre                | Method         | Scope and range of test |                                    | Targets  | TAT |
|-----------------------|----------------|-------------------------|------------------------------------|--|-----|
| Aberdeen              | NGS            | Whole gene<br>screen    | SNVs, indels,<br>Exon level<br>CNV | ADAMTS18, ALDH18A1, B3GLCT,<br>CHRDL1, CHST6, COL8A2, DCN, GJA1,<br>GSN, HMX1, KERA, KRT12, KRT3, LTBP2,<br>MAF, OVOL2, PIK3R1, PIKFYVE, PITX2,<br>PRDM5, RAB18, RAB3GAP1, RAB3GAP2,<br>SLC16A12, SLC4A11, TACSTD2, TGFBI,<br>UBIAD1, VSX1, ZEB1, ZNF469 | 112 |
| Family r              | member<br>ting | as indicated above      |                                    | 14   |     |
| Proforma required? NO |                |                         |                                    |  |     |

#### Referral criteria

Clinical features suggestive of a monogenic cause of corneal abnormalities

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





# **CORNEAL DYSTROPHY**

# Available testing

| Centre      | Method         | Scope and range of test |                                    | Targets  | TAT |  |
|-------------|----------------|-------------------------|------------------------------------|--|-----|--|
| Aberdeen    | NGS            | Whole<br>gene<br>screen | SNVs, indels,<br>Exon level<br>CNV | CHST6, COL17A1, COL8A2, DCN, GRHL2,<br>GSN, KERA, KRT12, KRT3, LCAT, OVOL2,<br>PIKFYVE, PRDM5, SLC4A11, STS, TACSTD2,<br>TCF4, TGFBI, UBIAD1, ZEB1, ZNF469 | 112 |  |
| ,           | member<br>ting |                         | as indicated above                 |  | 14  |  |
| Proforma re | equired?       | NO                      | NO                                 |  |     |  |

## Referral criteria

Clinical features suggestive of a monogenic cause of Corneal Dystrophy

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





## **EYE MOVEMENT DISORDER**

## Available testing

| Centre      | Method                       | Scope and range of test |                                   | Targets   | TAT |
|-------------|------------------------------|-------------------------|-----------------------------------|---|-----|
| Aberdeen    | NGS                          | Whole gene<br>screen    | SNVs, indels<br>Exon level<br>CNV | CHN1, COL25A1, DCC, FRMD7,<br>HOXA1, KIF21A, MAFB, PHOX2A,<br>ROBO3, SALL1, SALL4, TUBB2B,<br>TUBB3 | 56  |
| ,           | Family member as inditesting |                         | cated above                       | 14  |     |
| Proforma re | equired?                     | NO                      |                                   |   |     |

#### Referral criteria

 Clinical features suggestive of a monogenic cause of an eye movement disorder

# Requesting specialties

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

# FAMILIAL EXUDATIVE VITRORETINOPATHY (FEVR)

# Available testing

| Centre      | Method   | Scope and range of test |                                | Targets                            | TAT |
|-------------|----------|-------------------------|--------------------------------|------------------------------------|-----|
| Aberdeen    | NGS      | Whole gene screen       | SNVs, indels<br>Exon level CNV | ATOH7, FZD4, LRP5, NDP,<br>TSPAN12 | 56  |
| Family r    |          |                         | as indicated                   | d above                            | 14  |
| Proforma re | equired? | NO                      |                                |                                    |     |

#### Referral criteria

• Clinical features suggestive of Familial Execudative Vitroretinopathy – vision loss or blindness, retinal detachment, strabismus, leukocoria

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





## **GLAUCOMA**

# Available testing

| Centre      | Method   | Scope and range of test |                                    | Targets   | TAT |  |
|-------------|----------|-------------------------|------------------------------------|---|-----|--|
| Aberdeen    | NGS      | Whole<br>gene<br>screen | SNVs, indels,<br>Exon level<br>CNV | ADAMTS10, ADAMTS17, CPAMD8, CREBBP,<br>CYP1B1, DDX58, FOXC1, FOXE3, IFIH1,<br>LMX1B, LTBP2, MYOC, OCRL, PAX6, PITX2,<br>SBF2, SH3PXD2B, TEK | 112 |  |
| Family r    |          |                         | as indicated above                 |   | 14  |  |
| Proforma re | equired? | NO                      | 0                                  |   |     |  |

## Referral criteria

Clinical features suggestive of a monogenic cause of Glaucoma

# Requesting specialties

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

## VITELLIFORM MACULAR DYSTROPHY

## Available testing

| Centre             | Method             | Scope and range of test |                                | Targets      | TAT |
|--------------------|--------------------|-------------------------|--------------------------------|--------------|-----|
| Aberdeen           | Sanger<br>MLPA     | Whole gene screen       | SNVs, indels<br>Exon level CNV | BEST1, PRPH2 | 56  |
| ,                  | y member as indica |                         | ted above                      | 14           |     |
| Proforma required? |                    | NO                      |                                |              |     |

## Referral criteria

 Clinical features suggestive of monogenic Macular Dystrophy – loss of central vision

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





## **NORRIE DISEASE**

# Available testing

| Centre                | Method         | Scope and range of test |                                    | Targets | TAT |
|-----------------------|----------------|-------------------------|------------------------------------|---------|-----|
| Aberdeen              | Sanger<br>MLPA | Whole gene screen       | SNVs, indels<br>Exon level<br>CNVs | NDP     | 56  |
| Family member testing |                |                         | as indicated                       | d above | 14  |
| Proforma re           | equired?       | NO                      |                                    |         |     |

## Referral criteria

 Clinical features and ocular investigations suggestive of Norrie disease consistent with X-linked congenital blindness

- Clinical Genetics
- in discussion with Clinical Genetics





## **OCULAR MALFORMATIONS**

# Available testing

| Centre                | Method                           | Scope and range of test |                 | Targets  | TAT |
|-----------------------|----------------------------------|-------------------------|-----------------|--|-----|
| Edinburgh             | NGS                              | Whole gene<br>screen    | SNVs,<br>indels | ACTB, ACTG1, ALDH1A3, BCOR,<br>C12ORF57, CHD7, COL4A1, FOXC1,<br>FOXE3, CHD7, GJA8, ITPA, ITPR1,<br>MAB21L1, MAB21L2, NAA10, OTX2, PAX2,<br>PAX6, PITX2, PITX3, RAB18, RAB3GAP1,<br>RAB3GAP2, RARB, RAX, RBP4, SALL2,<br>SALL4, SHH, SIX3, SMCHD1, SMOC1,<br>SOX2, STRA6, TBC1D20, VAX1, VSX2,<br>YAP1, ZEB2, ZIC2 | 112 |
| Family men            | ember testing as indicated above |                         | ndicated above  | 14   |     |
| Proforma required? NO |                                  | NO                      |                 |  |     |

## Referral criteria

- Non-syndromic microphthalmia, anophthalmia, coloboma (MAC) spectrum
- Aniridia
- Microarray analysis is recommended prior to testing as copy number variants are frequently observed in both MAC spectrum disorders and aniridia

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics
- Paediatrics





## **OCULOCUTANEOUS ALBINISM**

## Available testing

| Centre      | Method         | Scope and range of test |                                    | Targets   | TAT |
|-------------|----------------|-------------------------|------------------------------------|---|-----|
| Aberdeen    | NGS            | Whole gene<br>screen    | SNVs, indels,<br>Exon level<br>CNV | GPR143, HPS1, HPS3, HPS4, HPS5,<br>LRMDA, LYST, OCA2, SLC24A5,<br>SLC45A2, TYR, TYRP1 | 112 |
| Family r    | member<br>ting |                         | as indi                            | cated above   | 14  |
| Proforma re | equired?       | NO                      |                                    |   |     |

#### Referral criteria

- Clinical features suggestive of Oculocutaneous Albinism very light skin and light coloured irises, decreased sharpness of vision, nystagmus, strabismum, photophobia
- Where X-linked Oculocutaneous Albinism is suspected, single gene testing for GPR143 can be requested (Sanger and MLPA)

# Requesting specialties

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

#### **OPTIC NEUROPATHY**

## Available testing

| Centre      | Method         | Scope and range of test |                                 | Targets  | TAT |
|-------------|----------------|-------------------------|---------------------------------|--|-----|
| Aberdeen    | NGS            | Whole<br>gene<br>screen | SNVs, indels,<br>Exon level CNV | ACO2, C12orf65, C19orf12, CISD2, DNM1L,<br>MFF, MFN2, NR2F1, OPA1, OPA3, RTN4IP1,<br>SLC25A46, SLC52A2, SPG7, SSBP1,<br>TMEM126A, WFS1 | 112 |
| Family r    | member<br>ting |                         | as indicated above              |  | 14  |
| Proforma re | equired?       | NO                      |                                 |  |     |

## Referral criteria

Clinical features suggestive of an Optic Neuropathy

## Requesting specialties

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

NSD611-003.20 V





## **RETINAL DISORDERS**

# Available testing

| Centre      | Method   | Scope and ra         | ange of test                          | Targets   | TAT |
|-------------|----------|----------------------|---------------------------------------|---|-----|
| Aberdeen    | NGS      | Whole gene<br>screen | SNVs,<br>indels,<br>Exon level<br>CNV | ABCA4, ABHD12, ACO2, ADAM9, ADAMTS18, ADGRV1, AGBL5, AHI1, AIPL1, AIRE, ALMS1, ARHGEF18, ARL2BP, ARL6, ATF6, ATOH7, BBS1, BBS10, BBS12, BBS2, BBS4, BBS5, BBS7, BBS9, BEST1, C1QTNF5, C8orf37, CABP4, CACNA1F, CACNA2D4, CAPN5, CC2D2A, CDH23, CDH3, CDHR1, CEP164, CEP290, CEP78, CERKL, CFAP410 (C21orf2), CFH, CHM, CIB2, CLN3, CLN5, CLN6, CLN8, CLRN1, CNGA1, CNGA3, CNGB1, CNGB3, CNNM4, COL18A1, COL4A1, CRB1, CRX, CSPP1, CTNNB1, CTSD, CWC27, CYP4V2, DHDDS, EFEMP1, ELOVL4, ERCC6, ERCC8, EYS, FAM161A, FLVCR1, FZD4, GNAT1, GNAT2, GNPTG, GPK143, GPR179, GRK1, GRM6, GUCA1A, GUCA1B, GUCY2D, HARS, HCCS, HGSNAT, HMX1, IDH3A, IDH3B, IFT140, IKBKG, IMPDH1, IMPG1, IMPG2, INPP5E, IQCB1, KCNJ13, KCNV2, KIAA1549, KIF11, KIZ, KLHL7, LCA5, LRAT, LRIT3, LRP2, LRP5, LZTFL1, MAK, MERTK, MFRP, MFSD8, MKKS, MKS1, MYO7A, NDP, NMNAT1, NPHP1, NPHP3, NPHP4, NR2E3, NRL, NYX, OAT, OFD1, OPN1LW, OPN1MW, OTX2, PANK2, PCARE (c2orf71), PCDH15, PCYT1A, PDE6A, PDE6B, PDE6C, PDE6G, PEX1, PEX2, PEX7, PHYH, PLA2G5, POC1B, PPT1, PRCD, PROM1, PRPF3, PRPF31, PRPF4, PRPF6, PRPF8, PRPH2, PRPS1, RAB28, RAX2, RBP3, RBP4, RCBTB1, RD3, RDH12, RDH5, REEP6, RGS9, RHO, RLBP1, RP1, RP1L1, RP2, RP9, RPE65, RPGR, RPGRIP1, RPGRIP1L, RS1, SAG, SCAPER, SDCCAG8, SLC24A1, SLC38A8, SNRNP200, SPATA7, SRD5A3, TIMM8A, TIMP3, TMEM237, TOPORS, TPP1, TRIM32, TRPM1, TSPAN12, TTC8, TTLL5, TUB, TULP1, USH1C, USH1G, USH2A, VCAN, VPS13B, WDPCP, WDR19, WHRN, ZNF408, ZNF423 | 112 |
|             | member   |                      |                                       | as indicated above  | 14  |
|             | ting     |                      |                                       |   |     |
| Proforma re | equired? | NO                   |                                       |   |     |

#### Referral criteria

- Clinical features suggestive of a monogenic Retinal disorder
- Where clinical testing indicates a subset of genes should be tested, please indicate this on the referral form and testing can be performed by either NGS or Sanger sequencing.
- Please note, ORF15 sequencing is not currently available in the Aberdeen laboratory. Where testing is required, please send to the Manchester laboratory.

# Requesting specialties

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

NSD611-003.20 V5





## **USHER SYNDROME**

# Available testing

| Centre      | Method         | Scope and range of test |                                    | Targets   | TAT |
|-------------|----------------|-------------------------|------------------------------------|---|-----|
| Aberdeen    | NGS            | Whole<br>gene<br>screen | SNVs, indels,<br>Exon level<br>CNV | MYO7A, USH1C, CDH23, PCDH15, USH1G,<br>ADGRV1, DFNB31 (WHRN), USH2A | 56  |
| Family r    | member<br>ting |                         | as indicated above                 |   | 14  |
| Proforma re | equired?       | NO                      |                                    |   |     |

#### Referral criteria

- Clinical features suggestive of Usher Syndrome retinitis pigmentosa and sensorineural hearing loss.
- If clinical presentation is mainly hearing loss, testing should be performed in Dundee

# Requesting specialties

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

## X-LINKED CONGENITAL NYSTAGMUS

# Available testing

| Centre             | Method       | Scope and range of test |              | Targets   | TAT |
|--------------------|--------------|-------------------------|--------------|-----------|-----|
| Aberdeen           | Sanger       | Whole gene screen       | SNVs, indels | FRMD7     | 56  |
| Family me          | mber testing |                         | as indica    | ted above | 14  |
| Proforma required? |              | NO                      |              |           |     |

## Referral criteria

 Clinical features suggestive of X-linked Congenital Nystagmus – nystagmus presenting within first 6 months of life

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





## X-LINKED JUVENILE RETINOSCHISIS

# Available testing

| Centre      | Method                   | Scope and range of test |              | Targets | TAT |
|-------------|--------------------------|-------------------------|--------------|---------|-----|
| Aberdeen    | Sanger                   | Whole gene screen       | SNVs, indels | RS1     | 56  |
| Family r    | Family member as indicat |                         | ed above     | 14      |     |
| testing     |                          |                         |              |         |     |
| Proforma re | equired?                 | NO                      |              |         |     |

Can be performed prior to Retinal Degeneration panel if required

## Referral criteria

• Clinical features suggestive of X-linked Juvenile Retinoschisis

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





# **GASTROHEPATOLOGY**

#### **CHOLESTASIS**

# Available testing

| Centre      | Method         | Scope and range of test |                                   | Targets  | TAT |
|-------------|----------------|-------------------------|-----------------------------------|--|-----|
| Aberdeen    | NGS            | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | ABCB11, ABCB4, ABCC2, AKR1D1, ALDOB,<br>AMACR, ATP8B1, BAAT, BCS1L, CLDN1,<br>CYP27A1, CYP7A1, DCDC2, FAH, HSD3B7,<br>JAG1, MYO5B, NOTCH2, NPC1, NPC2, NR1H4,<br>PEX1, PEX12, PEX26, PEX6, SERPINA1,<br>SLC25A13, TALDO1, TJP2, UGT1A1, VIPAS39,<br>VPS33B | 112 |
| ,           | member<br>ting |                         |                                   | as indicated above   | 14  |
| Proforma re | equired?       | NO                      |                                   |  |     |

## Referral criteria

- Neonatal conjugated hyperbilirubinaemia where multifactorial and infective causes have been excluded
- Unexplained cholestasis developing <18 years old
- Unexplained cholestasis >18 years old where other causes excluded

- Clinical Genetics
- Gastroenterology
- Hepatology
- Paediatrics





## **CRIGLER-NAJJAR SYNDROME, TYPE 1 AND 2**

# Available testing

| Centre             | Method                     | Scope and range of test            |   | Targets                        | TAT |
|--------------------|----------------------------|------------------------------------|---|--------------------------------|-----|
| Dundee             | Sanger & fragment analysis | Targeted screen   Promoter variant |   | UGT1A1, TA Allele7 (A[TA]7TAA) | 28  |
| Proforma required? |                            | NO                                 | _ |                                |     |

#### Referral criteria

- Individuals with unconjugated hyperbilirubinaemia in the absence of haemolysis, where a molecular diagnosis will contribute to management
- Urgent requests for neonates are processed in 5 days.

# Requesting specialties

- Clinical Genetics
- Gastroenterology
- Hepatology
- Paediatrics

# **GILBERT SYNDROME**

## Available testing

| Centre             | Method   | Scope and range of test          |  | Targets                | TAT |
|--------------------|----------|----------------------------------|--|------------------------|-----|
| Dundee             | Fragment | Targeted screen Promoter variant |  | TA Allele7 (A[TA]7TAA) | 28  |
| Proforma required? |          | NO                               |  |                        |     |

#### Referral criteria

 Individuals with mild unconjugated hyperbilirubinaemia in the absence of haemolysis, where a molecular diagnosis will contribute to management

- Clinical Genetics
- Gastrohepatology
- Paediatrics
- General Practice





## HIRSCHSPRUNG DISEASE

## Available testing

| Centre                | Method                     | Scope and range of test |              | Targets  | TAT |
|-----------------------|----------------------------|-------------------------|--------------|--|-----|
| Dundee                | NGS<br>(clinical<br>exome) | Whole gene<br>screen    | SNVs, indels | : EDN3, EDNRB, KIF1BP, L1CAM,<br>PHOX2B, RET, SOX10,ZEB2 | 112 |
| Family member testing |                            |                         | as indicate  | d above  | 14  |
| Proforma required? NO |                            | NO                      |              |  |     |

#### Referral criteria

- Diagnosis of Hirschsprung disease (HSCR) and at least one of the following:
  - Family history of HSCR, at least 1 affected first or second degree relative, OR
  - HSCR occurring as part of a syndrome or with other anomalies associated with the listed genes.
  - If there is a clinical suspicion of Congenital Central Hypoventilation Syndrome (CCHS) in association with HSCR, please discuss with clinical genetics to arrange the most appropriate testing.

- Clinical Genetics
- Gastroenterology
- Hepatology
- Paediatrics





## **PANCREATITIS**

# Available testing

| Centre                                     | Method         | Scope and range of test |                                   | Targets              | TAT |
|--|----------------|-------------------------|-----------------------------------|----------------------|-----|
| Aberdeen                                   | Sanger<br>MLPA | Whole gene<br>screen    | SNVs, indels<br>Exon level<br>CNV | SPINK1, PRSS1        | 56  |
| Aberdeen<br>Dundee<br>Edinburgh<br>Glasgow | ARMS           | Targeted screen         | SNVs, indels                      | CFTR common variants | 28  |
| Family member testing                      |                |                         | as indica                         | ited above           | 14  |
| Proforma required?                         |                | NO                      |                                   |                      |     |

## Referral criteria

- Recurrent acute pancreatitis
- Chronic pancreatitis
- 1st episode of acute pancreatitis <18 years old
- 1<sup>st</sup> episode of acute pancreatitis with a first degree relative who has also had pancreatitis
- Secondary causes excluded (e.g. excessive alcohol, gallstones)

- Clinical Genetics
- Gastroenterology
- Hepatology
- Lipidology
- Paediatrics





## **PORPHYRIAS**

# Available testing

| Centre                | Method   | Scope and range of test |                                   | Targets  | TAT |
|-----------------------|----------|-------------------------|-----------------------------------|--|-----|
| Aberdeen              | NGS      | Whole gene screen       | SNVs, indels<br>Exon level<br>CNV | ALAD, ALAS2, CPOX, FECH, HMBS,<br>PPOX, UROD, UROS | 56  |
| Family member testing |          |                         | as indicat                        | red above  | 14  |
| Proforma re           | equired? | NO                      |                                   |  |     |

#### Referral criteria

Clinical diagnosis of porphyria with suspected monogenic cause

# Requesting specialties

- Clinical Genetics
- Gastroenterology
- Hepatology

## **WILSON DISEASE**

# Available testing

| Centre      | Method         | Scope and range of test |                                | Targets  | TAT |
|-------------|----------------|-------------------------|--------------------------------|----------|-----|
| Aberdeen    | Sanger<br>MLPA | Whole gene screen       | SNVs, indels<br>Exon level CNV | АТР7В    | 56  |
| Family r    | member<br>ting |                         | as indicat                     | ed above | 14  |
| Proforma re | equired?       | NO                      |                                |          |     |

## Referral criteria

• Clinical phenotype suggestive of Wilson disease – high liver copper, high urinary copper, high free copper, low caeruloplasmin

- Clinical Genetics
- Gastroenterology
- Hepatology





# **HAEMATOLOGY**

## **ANTITHROMBIN DEFICIENCY**

# Available testing

| Centre      | Method         | Scope and range of test |          |                                   | Targets                        | TAT |
|-------------|----------------|-------------------------|----------|-----------------------------------|--------------------------------|-----|
| Edinburgh   | Sanger<br>MLPA | Whole gene<br>screen    |          | SNVs, indels<br>Exon level<br>CNV | SERPINC1                       | 56  |
| Family men  | nber testing   |                         |          | as indi                           | cated above                    | 14  |
| Proforma re | equired?       | YES                     | Molecula | r Haematology rec                 | uest form (see centre website) |     |

## Referral criteria

 Antithrombin activity and/or antigen below the normal range on at least two occasions

# Requesting specialties

- Clinical Genetics
- Haematology

#### **BERNARD-SOULIER SYNDROME**

## Available testing

| Centre      | Method       | Scope and range of test |          |                   | Targets                        | TAT |
|-------------|--------------|-------------------------|----------|-------------------|--------------------------------|-----|
| Edinburgh   | NGS          | Whole gene screen       |          | SNVs              | GP1BA, GP1BB, GP9              | 56  |
| Family men  | nber testing |                         |          | as indi           | cated above                    | 14  |
| Proforma re | equired?     | YES                     | Molecula | r Haematology red | uest form (see centre website) |     |

## Referral criteria

• Platelet function testing suggestive of Bernard Soulier syndrome

- Clinical Genetics
- Haematology





#### **COAGULATION & FIBRINOLYSIS PANEL**

## Available testing

| Centre      | Method       | Scope and range of test |                              | nge of test | Targets  | TAT |
|-------------|--------------|-------------------------|------------------------------|-------------|--|-----|
| Edinburgh   | NGS          |                         | e gene<br>een                | SNVs        | ACVRL1, CHST14, COL3A1, ENG, F2,<br>F5, F7, F8, F9, F10, F11, F12, F13A1,<br>F13B, FGA, FGB, FGG, GGCX, KLKB1,<br>KNG1, LMAN1, MCFD2, SERPINE1,<br>SERPINF2, THBD, VKORC1, VWF | 84  |
| Family mem  | nber testing |                         | as inc                       |             | cated above  | 14  |
| Proforma re | equired?     | YES                     | YES Molecular Haematology re |             | quest form (see centre website)  |     |

## Referral criteria

- · Suspected congenital unexplained bleeding disorder, meeting both of
  - o normal coagulation factors or deficiency of multiple coagulation factors
  - life long significant bleeding history (eg OBS >9), or personal bleeding history and family history of bleeding
- **Note:** specific genes are available as sub-panels where there is a highly suggestive phenotype such as Factor II, V or XIII deficiency

- Clinical Genetics
- Haematology





## **COMBINED FACTOR V AND VIII DEFICIENCY**

## Available testing

| Centre                                 | Method       | Scope and range of test    |                      |   | Targets              | TAT |
|--|--------------|----------------------------|----------------------|---|----------------------|-----|
| Edinburgh                              | NGS          | Whole gene<br>screen       |                      | SNVs<br>(plus exon level CNV for<br>F8 where appropriate) | F5, F8, LMAN1, MCFD2 | 56  |
| Family mem                             | nber testing |                            |                      | as indicated abov   | ve .                 | 14  |
| Proforma required? YES Molecular Haema |              | r Haematology request form | (see centre website) |   |                      |     |

#### Referral criteria

Factor V and factor VIII levels below the normal range on at least two occasions

# Requesting specialties

- Clinical Genetics
- Haematology

## **DIAMOND BLACKFAN ANAEMIA**

# Available testing

| Centre      | Method         | Scope and range of test |                                   | Targets  | TAT |
|-------------|----------------|-------------------------|-----------------------------------|--|-----|
| Aberdeen    | NGS            | Whole gene<br>screen    | SNVs, indels<br>Exon level<br>CNV | RPL5, RPS10, RPL11, RPL35A, RPS7,<br>RPS19, RPS24, RPS26, GATA1, RPS17 | 56  |
| _           | member<br>ting |                         | as indi                           | cated above  | 14  |
| Proforma re | equired?       | NO                      |                                   |  |     |

## Referral criteria

 Clinical phenotype suggestive of Diamond Blackfan Anaemia – Presenting in the 1<sup>st</sup> year of life. Normochromic macrocytic anaemia, reticulocytopenia and nearly absent erythroid progenitors in the bone marrow.

- Clinical Genetics
- Haematology





## **ERYTHROCYTOSIS**

# Available testing

| Centre                | Method                | Scope and range of test |              | Targets  | TAT |
|-----------------------|-----------------------|-------------------------|--------------|--|-----|
| Edinburgh             | NGS                   | Whole<br>gene<br>screen | SNVs, indels | EGLN1, EPAS1, EPO, EPOR, HBA1, HBA2,<br>HBB, VHL | 112 |
| Family mem            | Family member testing |                         |              | as indicated above                               | 14  |
| Proforma required? NO |                       |                         |              |  |     |

## Referral criteria

- Idiopathic erythrocytosis with:
  - No acquired JAK2 variants
  - Secondary causes excluded
  - Young onset and/or family history

# Requesting specialties

- Clinical Genetics
- Haematology

## **FACTOR VII DEFICIENCY**

# Available testing

| Centre                             | Method         | Scope and range of test |                    |   | Targets | TAT |  |
|------------------------------------|----------------|-------------------------|--------------------|---|---------|-----|--|
| Edinburgh                          | Sanger<br>MLPA | Whole gene screen       |                    | SNVs<br>Exon level CNV                        | F7      | 56  |  |
| Family mem                         | ber testing    |                         | as indicated above |   |         |     |  |
| Proforma required? YES Molecular F |                |                         | Molecular Ha       | Haematology request form (see centre website) |         |     |  |

## Referral criteria

Factor VII level below the normal range on at least two occasions

- Clinical Genetics
- Haematology





## **FACTOR X DEFICIENCY**

## Available testing

| Centre  | Method         |  | Scope and ra        | ange of test              | Targets | TAT |
|---|----------------|--|---------------------|---------------------------|---------|-----|
| Edinburgh   | Sanger<br>MLPA |  | nole gene<br>screen | SNVs<br>Exon level CNV    | F10     | 56  |
| Family me   | ember testing  |  |                     | ed above                  | 14      |     |
| Proforma required? YES   Molecular Haematology re |                |  | aematology request  | form (see centre website) |         |     |

#### Referral criteria

• Factor X level below the normal range on at least two occasions

# Requesting specialties

- Clinical Genetics
- Haematology

## **FACTOR XI DEFICIENCY**

## Available testing

| Centre   | Method         |  | Scope and r              | ange of test           | Targets | TAT |
|--|----------------|--|--------------------------|------------------------|---------|-----|
| Edinburgh  | Sanger<br>MLPA |  | nole gene<br>screen      | SNVs<br>Exon level CNV | F11     | 56  |
| Family me  | ember testing  |  |                          | as indicated           | above   | 14  |
| Proforma required? YES Molecular Haematology request f |                |  | orm (see centre website) |                        |         |     |

## Referral criteria

Factor XI level below the normal range on at least two occasions

- Clinical Genetics
- Haematology





## **FIBRINOGEN DEFICIENCY**

# Available testing

| Centre     | Method       | Scope and range of test |          |                   | Targets                         | TAT |
|------------|--------------|-------------------------|----------|-------------------|---------------------------------|-----|
| Edinburgh  | NGS          | Whole gene screen       |          | SNVs              | FGA, FGB, FGG                   | 56  |
| Family mer | nber testing |                         |          | cated above       | 14                              |     |
| Proforma r | equired?     | YES                     | Molecula | r Haematology red | quest form (see centre website) |     |

#### Referral criteria

• Diagnosis of hypo-, a- or dys- fibrinogenaemia with a reduced antigenic and/or functional fibrinogen level on at least two occasions

# Requesting specialties

- Clinical Genetics
- Haematology

# **G6PD Deficiency**

# Available testing

| Centre      | Method   | Scope and range of test |   |  |    | Targets      | TAT |
|-------------|----------|-------------------------|---|--|----|--------------|-----|
| Edinburgh   | NGS      | Whole gene<br>screen    |   |  | 's | G6PD         | 56  |
| Family r    |          |                         | · |  |    | icated above | 14  |
| Proforma re | equired? | NO                      |   |  |    |              |     |

#### Referral criteria

- Genetic test result will aid determination of carrier status in female at significant risk because of family history
  - Male with a clinical suspicion of G6PD deficiency and G6PD activity results are unavailable or uninformativeHaematology





## **GLANZMANN THROMBASTHENIA**

# Available testing

| Centre      | Method       | Scope and range of test |          |                   | Targets                         | TAT |
|-------------|--------------|-------------------------|----------|-------------------|---------------------------------|-----|
| Edinburgh   | NGS          | Whole gene screen       |          | SNVs              | ITGA2B, ITGB3                   | 56  |
| Family mem  | nber testing |                         |          | as indi           | cated above                     | 14  |
| Proforma re | equired?     | YES                     | Molecula | r Haematology red | quest form (see centre website) |     |

## Referral criteria

• Platelet function testing suggestive of Glanzmann thrombasthenia

- Clinical Genetics
- Haematology





## **HAEMOCHROMATOSIS**

# Available testing

| Centre                        | Method                                     | Scope and range of test |    | Targets                     | TAT |  |  |  |
|-------------------------------|--|-------------------------|----|-----------------------------|-----|--|--|--|
| Aberdeen<br>Dundee<br>Glasgow | ARMS (G)<br>Sanger (A)<br>Genotyping Assay | Targeted screen SNVs    |    | <i>HFE</i> p.C282Y & p.H63D | 28  |  |  |  |
| Proforma required?            |  | NO                      | NO |                             |     |  |  |  |

## Referral criteria

• Raised serum ferritin and transferrin saturation

# Requesting specialties

- Clinical Genetics
- GPs
- Haematology

# HAEMOGLOBINOPATHY (incl. SICKLE CELL DISEASE, ALPHA AND BETA THALASSAEMIAS)

# Available testing

| Centre      | Method         | Scope and r        | ange of test | Targets | TAT |  |
|-------------|----------------|--------------------|--------------|---------|-----|--|
| Edinburgh   | Sanger<br>MLPA | Whole gene screen  | SNVs, indels | HBB     | 56  |  |
| Edinburgh   | MLPA           |                    | Indels       | HBA     | 56  |  |
| Family me   | ember testing  | as indicated above |              |         |     |  |
| Proforma re | equired?       | NO                 |              |         | •   |  |

## Referral criteria

• Clinical features indicative of likely thalassaemia or other clinically significant haemoglobinopathy.

- Clinical Genetics
- Haematology





## **HAEMOPHILIA A**

## Available testing

| Centre                             | Method                       |     | Scope and   | d range of test                        | Targets | TAT |
|------------------------------------|------------------------------|-----|---|--|---------|-----|
| Edinburgh                          | NGS<br>MLPA<br>Inversion PCR |     | hole gene<br>screen                                     | SNVs<br>Exon level CNV<br>Inversions * | F8      | 56  |
| Family member testing as indicated |                              |     | as indicated above                                      | )                                      | 14      |     |
| Proforma red                       | quired?                      | YES | Molecular Haematology request form (see centre website) |  |         | •   |

<sup>\*</sup> Inversion testing includes recurrent inversions with breakpoints within F8 intron 1 and 22 and is only included for severe haemophilia A, or moderate haemophilia A where no other causative variant is identified

## Referral criteria

Factor VIII level below the normal range on at least two occasions

# Requesting specialties

- Clinical Genetics
- Haematology

## **HAEMOPHILIA B**

## Available testing

| Centre                                   | Method         | Scope and range of test |   |                                | Targets | TAT |
|--|----------------|-------------------------|---|--------------------------------|---------|-----|
| Edinburgh                                | Sanger<br>MLPA | Who                     | e gene screen   | SNVs, indels<br>Exon level CNV | F9      | 56  |
| Family member testing as indicated above |                |                         |   | as indicated above             |         | 14  |
| Proforma required?                       |                |                         | YES   Molecular Haematology request form (see centre website) |                                |         |     |

#### Referral criteria

Factor IX level below the normal range on at least two occasions

- Clinical Genetics
- Haematology





## **INHERITED BONE MARROW FAILURE**

# Available testing

| Centre      | Method         | Scope and range of test |                                | Targets   | TAT |  |
|-------------|----------------|-------------------------|--------------------------------|---|-----|--|
| Aberdeen    | NGS            | Whole gene<br>screen    | SNVs, indels<br>Exon level CNV | BRCA2, BRIP1, CTC1, DKC1, ELANE,<br>FANCA, FANCB, FANCC, FANCD2,<br>FANCE, FANCF, FANCG, FANCI,<br>FANCL, FANCM, G6PC3, GATA1,<br>GATA2, GFI1, HAX1, MPL, NHP2,<br>NOP10, PALB2, RAD51C, RPL11,<br>RPL35A, RPL5, RPS10, RPS17, RPS19,<br>RPS24, RPS26, RPS7, RUNX1, SBDS,<br>SLX4, SRP72, TERT, TINF2, WAS,<br>WRAP53 | 112 |  |
| ,           | member<br>tina |                         | as indic                       | ated above  | 14  |  |
| Proforma re | equired?       |                         |                                |   |     |  |

## Referral criteria

- Clinical phenotype suggestive of an inherited bone marrow failure disorder
- Please note, the content of this panel is currently under review and will be discussed with users to ensure correct content and refine referral criteria.

- Clinical Genetics
- Haematology





#### **IRON REGULATION**

## Available testing

| Centre                | Method | Scope and range of test |              | Targets   | TAT |
|-----------------------|--------|-------------------------|--------------|---|-----|
| Edinburgh             | NGS    | Whole gene<br>screen    | SNVs, indels | ABCB7, ALAS2, ATP7B, BMP6, CP,<br>CYBRD1, FTL, GBA, GLRX5, HAMP, HFE,<br>HFE2, SLC11A2, SLC25A38, SLC40A1, TF,<br>TFR2, TMPRSS6 | 112 |
| Family member testing |        |                         | as ir        | ndicated above  | 14  |
| Proforma required?    |        | NO                      |              |   |     |

#### Referral criteria

- Juvenile Haemochromatosis (<30years) with severe iron overload in liver AND/OR heart. Raised serum ferritin >1000ug/L and transferrin saturation >90%
- Juvenile Haemochromatosis >30 years with unexplained severe haemochromatosis and HFE negative
- Ferroportin disease: raised serum ferritin with normal transferrin saturation and evidence of reticuloendothelial iron staining on liver biopsy or splenic iron overload on MRI and HFE mutations negative
- Haemochromatosis: raised serum ferritin and transferrin saturation C282Y negative
- Hereditary Hyperferritinemia cataract syndrome: High and constant levels of serum ferritin unresponsive to iron depletion and no signs of iron overload and no relevant clinical symptoms apart from visual impairment by cataract
- Biochemical evidence of unexplained iron overload and lack of homozygous/compound homozygous HFE mutations
- Iron Refractory Iron Deficiency Anaemia (IRIDA): Very low mean corpuscular volume (MCV) and low serum iron and low transferrin saturation, normal ferritin or ferritin levels in the lower limits of normal, no response to oral iron treatment

- Clinical Genetics
- Haematology
- Gastroenterology





## **MYELODYSPLASTIC SYNDROME**

## Available testing

| Centre                | Method | Scope and ra                                | ange of test | Targets      | TAT |  |
|-----------------------|--------|---|--------------|--------------|-----|--|
| Aberdeen              | NGS    | Whole gene screen SNVs, indel Exon level CN |              | SRP72, GATA2 | 56  |  |
| Family member testing |        | as indicated above                          |              |              |     |  |
| Proforma required?    |        | NO  |              |              | ·   |  |

## Referral criteria

• Clinical phenotype suggestive of monogenic Myelodysplastic syndrome

## Requesting specialties

- Clinical Genetics
- Haematology

## **NEUTROPENIA CONSISTENT WITH ELANE MUTATIONS**

## Available testing

| Centre             | Method                | So                   | cope and | range of test | Targets | TAT |
|--------------------|-----------------------|----------------------|----------|---------------|---------|-----|
| Aberdeen           | Sanger                | Whole gene<br>screen |          | SNVs, indels  | ELANE   | 56  |
| _                  | Family member testing |                      |          | as indicated  | d above | 14  |
| Proforma required? |                       | NO                   |          |               |         |     |

# Referral criteria

• Isolated neutropenia suggestive of ELANE pathogenic variants.

- Clinical Genetics
- Haematology





## **PLATELET PANEL**

# Available testing

| Centre                | Method | Scope and range of test |   | Targets   |    |  |
|-----------------------|--------|-------------------------|---|---|----|--|
| Edinburgh             | NGS    | Whole<br>gene<br>screen | SNVs,<br>indels   | ABCG5, ABCG8, ACTB, ACTN1, ADAMTS13, ANKRD26, ANO6, AP3B1, AP3D1, ARPC1B, BLOC1S3, BLOC1S6, CDC42, CYCS, DIAPH1, DTNBP1, ETV6, FERMT3, FLI1, FLNA, FYB1, GATA1, GBA, GFI1B, GNE, GP1BA, GP1BA, GP1BB, GP1BB, GP6, GP9, HOXA11, HPS1, HPS3, HPS4, HPS5, HPS6, IKZF5, ITGA2B, ITGA2B, ITGB3, ITGB3, KDSR, LYST, MECOM, MPIG6B, MPL, MYH9, NBEA, NBEAL2, P2RY12, PLA2G4A, PLAU, PTGS1, RASGRP2, RBM8A, RNU4ATAC, RUNX1, SLFN14, SRC, STIM1, STXBP2, TBXA2R, TBXAS1, THPO, TUBB1, TPM4, VIPAS39, VPS33B, VWF, WAS | 84 |  |
| Family member testing |        |                         |   | as indicated above  | 14 |  |
| Proforma required?    |        | YES                     | Molecular Haematology request form (see centre website) |   |    |  |

#### Referral criteria

- Suspected congenital (macro)thrombocytopenia or thrombocytopathy
- Confirmed platelet function defect (other than Glanzmann Thrombasthenia or Bernard Soulier syndrome pattern)
- Life long significant bleeding history (eg OBS >9), or personal bleeding history and family history of bleeding

- Clinical Genetics
- Haematology





## **PROTEIN C DEFICIENCY**

# Available testing

| Centre                | Method         | ,                    | Scope and   | range of test                  | Targets | TAT |
|-----------------------|----------------|----------------------|---|--------------------------------|---------|-----|
| Edinburgh             | Sanger<br>MLPA | Whole gene<br>screen |   | SNVs, indels<br>Exon level CNV | PROC    | 56  |
| Family member testing |                | as indicated above   |   |                                |         | 14  |
| Proforma required?    |                | YES                  | YES Molecular Haematology request form (see centre website) |                                |         |     |

## Referral criteria

Protein C level below the normal range on at least two occasions

# Requesting specialties

- Clinical Genetics
- Haematology

## **PROTEIN S DEFICIENCY**

# Available testing

| Centre                | Method         | Scope and range of test |  |                                | Targets | TAT |  |
|-----------------------|----------------|-------------------------|--|--------------------------------|---------|-----|--|
| Edinburgh             | Sanger<br>MLPA | Whole gene screen       |  | SNVs, indels<br>Exon level CNV | PROS1   | 56  |  |
| Family member testing |                | as indicated above      |  |                                |         | 14  |  |
| Proforma required?    |                | YES                     | ES Molecular Haematology request form (see centre website) |                                |         |     |  |

#### Referral criteria

• Protein S level below the normal range on at least two occasions

- Clinical Genetics
- Haematology





# RARE ANAEMIA PANEL (Panel app R92)

# Available testing

| Centre      | Method | Scope and rar        | ge of test   | Targets  | TAT |
|-------------|--------|----------------------|--------------|--|-----|
| Edinburgh   | NGS    | Whole gene<br>screen | SNVs, indels | ABCB7, ABCG5, ABCG8, ADA2, AK1, ALAS2, ALDOA, AMN, ANK1, C15orf41, CD59, CDAN1, COX4I2, CUBN, CYB5R3, DHFR, EPB41, EPB42, G6PD, GATA1, GCLC, GIF, GLRX5, GPI, GSR, GSS, HBA1, HBA2, HBB, HBD, HBG1, HBG2, HK1, HSPA9, KCNN4, KIF23, KLF1, LPIN2, MTR, MTRR, NT5C3A, PFKM, PIEZO1, PKLR, PUS1, RHAG, RPL11, RPL15, RPL26, RPL27, RPL31, RPL35A, RPL5, RPL9, RPS10, RPS17, RPS19, RPS24, RPS26, RPS27, RPS29, RPS7, SBDS, SEC23B, SLC11A2, SLC19A2, SLC25A38, SLC2A1, SLC4A1, SPTA1, SPTB, TCN2, TF, TMPRSS6, TPI1, TRNT1, UMPS, XK, YARS2 | 112 |
| Family r    |        |                      | as ii        | ndicated above   | 14  |
| Proforma re |        | NO                   |              |  | I   |

Gene list from panel app R92 rare anaemia panel v1.2 https://nhsgms-panelapp.genomicsengland.co.uk/panels/518/v1.2/

### Referral criteria

- Clinical presentation or biochemical enzyme deficiency highly suggestive of a specific monogenic red cell enzyme deficiency
- Clinical presentation highly suggestive of a specific monogenic red membrane disorderNon-immune haemolytic anaemia of likely monogenic cause with Haemoglobinopathies excluded

- Clinical Genetics
- Haematology





# **SCHWACHMAN-DIAMOND SYNDROME**

# Available testing

| Centre  | Method       | S                 | Scope and ra                       | ange of test                   | Targets       | TAT |
|---|--------------|-------------------|------------------------------------|--------------------------------|---------------|-----|
| Aberdeen  | NGS          | Whole gene screen |                                    | SNVs, indels<br>Exon level CNV | SBDS, DNAJC21 | 56  |
| Family me                                       | mber testing |                   |                                    | as indicated                   | above         | 14  |
| Proforma required? YES GEN FORM 215 Primary Imm |              |                   | nodeficiency Request form (see cen | tre website)                   |               |     |

## Referral criteria

• Clinical phenotype suggestive of Schwachman-Diamond Syndrome

- Clinical Genetics
- Haematology





#### **SICKLE CELL ANAEMIA**

# Available testing

| Centre             | Method | Scope and range of test                          |      | Targets                | TAT  |
|--------------------|--------|--|------|------------------------|--|
| Aberdeen           | Sanger | Targeted screen                                  | SNVs | HBB p.(Glu7Val)        | 28<br>Prenatal 3                           |
| Edinburgh          | Sanger | Targeted screen                                  | SNVs | HBB p.(Glu7Val)        | 28<br>Prenatal 3                           |
| Glasgow            | Sanger | Sanger Targeted screen (incl. newborn screening) |      | <i>HBB</i> p.(Glu7Val) | 28<br>Prenatal 3<br>Newborn screening<br>7 |
| Proforma required? |        | NO   |      |                        |  |

#### Referral criteria

- Sickle cell anaemia diagnosed by Haematology test
- For prenatal testing, both parents to be confirmed as carrier by genetics prior to offering invasive prenatal test. Please contact the laboratory to discuss
- Newborn screening (Glasgow)
  - Newborns who have undergone a blood transfusion prior to the blood spot sample being taken.

- Clinical Genetics
- Haematology
- Obstetrics





# THROMBOPHILIA (FACTOR V LEIDEN & PROTHROMBIN)

# Available testing

| Centre  | Method  | Scope and rang  | e of test | Targets                                  | TAT |
|---|---|-----------------|-----------|--|-----|
| Aberdeen<br>Dundee<br>EdinburghMP*<br>GlasgowRI | Sanger (A) Real time PCR(D) Real time PCR (E) | Targeted screen | SNVs      | <i>F5</i> p.R534Q<br><i>F</i> 2 c.*97G>A | 28  |
| Proforma required?                              |   | NO              |           |  |     |

<sup>\*</sup>Performed by Edinburgh Molecular pathology, see https://edinburghlabmed.co.uk/node/1728

#### Referral criteria

- Venous thromboembolic event less than 40 years, with no apparent secondary causes
- Family history of venous thromboembolic events

## Requesting specialties

- Clinical Genetics
- Haematology

#### **THROMBOSIS PANEL**

## Available testing

| Centre                              | Method       | Scope and range of test |               |                    | Targets   | TAT |
|-------------------------------------|--------------|-------------------------|---------------|--------------------|---|-----|
| Edinburgh                           | NGS          | Whole                   | e gene<br>een | SNVs               | ADAMTS13, F2, F5, HRG, PIGA, PLG,<br>PROC, PROS1, SERPINC1,<br>SERPIND1, THBD | 84  |
| Family mem                          | nber testing |                         |               | as indica          | ated above  | 14  |
| Proforma required? YES Molecular Ha |              |                         | Molecula      | r Haematology requ | est form (see centre website)   |     |

#### Referral criteria

- Significant personal and family history of thrombosis
- Normal protein C, protein S and antithrombin levels

- Clinical Genetics
- Haematology





# **VON WILLEBRAND DISEASE (VWD)**

# Available testing

| Centre                               | Method        |      | Scope and rai | nge of test            | Targets                 | TAT |
|--------------------------------------|---------------|------|---------------|------------------------|-------------------------|-----|
| Edinburgh                            | NGS<br>MLPA   | Whol | e gene screen | SNVs<br>Exon level CNV | VWF                     | 56  |
| Family m                             | ember testing |      |               | as indicated           | above                   | 14  |
| Proforma required? YES Molecular Hae |               |      | Molecular Hae | matology request fo    | rm (see centre website) |     |

#### Referral criteria

- Type 1/3 VWD: VWF antigen and/or activity below 30 IU/dL on at least two occasions
- Type 2 VWD: VWF antigenic or activity levels suggestive of type 2 VWD, with or without suggestive platelet function or multimer results.

- Clinical Genetics
- Haematology





# **BRANCHIOOTORENAL (BOR) SYNDROME**

# Available testing

| Centre     | Method                     | Scope and range of test |   | Targets          | TAT |
|------------|----------------------------|-------------------------|---|------------------|-----|
| Dundee     | NGS<br>(clinical<br>exome) | Whole gene<br>screen    | SNVs, indels,<br>Exon level CNV<br>(EYA1) | EYA1, SIX1, SIX5 | 56  |
|            | amily member testing       |                         | as indicated a                            | bove             | 14  |
| Proforma r | equired?                   | NO                      |   |                  |     |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition.
- Referrals should be discussed with Clinical Genetics.

- Audiology
- Clinical Genetics
- Nephrology





# **HEARING LOSS, SYNDROMIC & NON SYNDROMIC**

## Available testing

| Centre     | Method                     | Scope and               | range of test   | Targets  | TAT |
|------------|----------------------------|-------------------------|-----------------|--|-----|
| Dundee     | NGS<br>(clinical<br>exome) | Whole<br>gene<br>screen | SNVs,<br>indels | ABHD12, ACTG1, ADGRV1 (GPR98), ALMS1, ATP6V1B1, BCS1L, BSND, CABP2, CCDC50, CDH23, CEACAM16, CHD7, CIB2, CLDN14, CLPP, CLRN1, COCH, COL11A2, COL4A5, COL4A6, DIAPH1, DNMT1, DSPP, EDN3, EDNRB, EPS8, ESPN, ESRRB, EYA1, EYA4, FGF3, GATA3, GIPC3, GJB2, GJB3, GJB6, GPSM2, GRHL2, GRXCR1, GSDME (DFNA5), HOXA2, HSD17B4, ILDR1, KARS, KCNE1, KCNJ10, KCNQ1, KCNQ4, KIT, LARS2, LHFPL5, LOXHD1, LRTOMT, MARVELD2, MASP1, MITF, MSRB3, MYH14, MYH9, MYO15A, MYO3A, MYO6, MYO7A, OPA1, OSBPL2, OTOA, OTOF, OTOG, OTOGL, P2RX2, PAX2, PAX3, PCDH15, PDZD7, PJVK (DFNB59), PNPT1, POU3F4, POU4F3, PRPS1, PTPRQ, RDX, SALL1, SALL4, SERAC1, SERPINB6, SIX1, SIX5, SLC17A8, SLC26A4, SLC26A5, SLC4A11, SMPX, SNAI2, SOX10, SOX2, STRC, SYNE4, TBC1D24, TECTA, TIMM8A, TMC1, TMIE, TMPRSS3, TPRN, TRIOBP, USH1C, USH1G, USH2A, WFS1, WHRN (DFNB31) | 112 |
| Family men | mber                       |                         |                 | as indicated above   | 14  |
| Proforma r | equired?                   | NO                      |                 |  | ı   |

#### Referral criteria

- Discussion with Clinical Genetics is required before testing.
- Referrals for testing will be triaged by the Genomic Laboratory; testing should be targeted at those where a genetic or genomic diagnosis will guide management for the proband or family

- Audiology (with Clinical Genetics approval)
- Clinical Genetics





#### **NON-SYNDROMIC HEARING LOSS – DFNB1**

## Available testing

| Centre     | Method                                | Scope and            | d range of test                         | Targets    | TAT |
|------------|---------------------------------------|----------------------|---|------------|-----|
| Dundee     | Sanger<br>and<br>fragment<br>analysis | Whole gene<br>screen | SNVs, indels (GJB2)<br>Deletions (GJB6) | GJB2, GJB6 | 28  |
| Family tes | nember as indicated above             |                      | ve                                      | 14         |     |
| Proforma r | equired?                              | NO                   |   |            |     |

#### Referral criteria

• Any individual with congenital, sensorineural hearing loss which is confirmed, bilateral and has no syndromic features.

- Audiology
- Clinical Genetics
- Paediatrics





#### **PENDRED SYNDROME**

# Available testing

| Centre                | Method | Scope and range of test |              | Targets        | TAT |
|-----------------------|--------|-------------------------|--------------|----------------|-----|
| Dundee                | Sanger | Whole gene screen       | SNVs, indels | SLC26A4, FOXI1 | 56  |
| Family member testing |        |                         | as indic     | ated above     | 14  |
| Proforma required? NO |        |                         |              |                |     |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition:
  - o Autosomal recessive deafness also associated with thyroid goiter
  - Abnormal cochlea or enlarged vestibular aqueduct is considered the most likely presentation of Pendred Syndrome
- Note that *FOXI1* is analysed if a single heterozygous variant is detected in *SLC26A4*.

- Audiology
- Clinical Genetics
- Paediatrics





#### **USHER SYNDROME**

## Available testing

| Centre     | Method                     | Scope and range of test |              | Targets   | TAT |
|------------|----------------------------|-------------------------|--------------|---|-----|
| Dundee     | NGS<br>(clinical<br>exome) | Whole<br>gene<br>screen | SNVs, indels | ADGRV1, CDH23, CIB2, CLRN1, MYO7A,<br>PCDH15, PDZD7, USH1C, USH1G, USH2A,<br>WHRN | 112 |
| Family me  | mber testing               |                         |              | as indicated above  | 14  |
| Proforma r | equired?                   | NO                      |              |   |     |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition.
   Referrals should be discussed with Clinical Genetics.
- If clinical presentation is mainly ophthalmic, testing should be performed in Aberdeen

# Requesting specialties

Clinical Genetics

#### WAARDENBURG SYNDROME

## Available testing

| Centre                | Method                     | Scope and range of test |              | Targets                                       | TAT |
|-----------------------|----------------------------|-------------------------|--------------|---|-----|
| Dundee                | NGS<br>(clinical<br>exome) | Whole gene screen       | SNVs, indels | PAX3, MITF, SOX10, SNAI2, EDNRB, EDN3,<br>KIT | 112 |
| Family me             | mber testing               | <u>.</u>                |              | as indicated above                            | 14  |
| Proforma required? NO |                            |                         |              |   |     |

#### Referral criteria

• Any individual with a clinical presentation consistent with the condition.

## Requesting specialties

Clinical Genetics





#### **WOLFRAM SYNDROME**

# Available testing

| Centre      | Method           | Scope and range of test |              | Targets      | TAT |
|-------------|------------------|-------------------------|--------------|--------------|-----|
| Dundee      | NGS or<br>Sanger | Whole gene screen       | SNVs, indels | WFS1         | 56  |
| Family me   | mber testing     |                         | as ind       | icated above | 14  |
| Proforma re | equired?         | NO                      |              |              |     |

#### Referral criteria

• Any individual with a clinical presentation consistent with the condition.

- Clinical Genetics
- Endocrinology





# **IMMUNOLOGY**

## ADENOSINE DEAMINASE DEFICIENCY (ADAD)

# Available testing

| Centre  | Method         | Scope and range of test |  |                                | Targets      | TAT |
|---|----------------|-------------------------|--|--------------------------------|--------------|-----|
| Aberdeen  | Sanger<br>MLPA | Whole<br>scr            | e gene<br>een                          | SNVs, indels<br>Exon level CNV | ADA2 (CECR1) | 56  |
| Family mer testing                                  | nber           |                         |  | as indicated                   | above        | 14  |
| Proforma required? YES GEN FORM 215 Primary Immunoo |                |                         | deficiency Request form (see centre we | ebsite)                        |              |     |

#### Referral criteria

- Polyarteritis nodosa, childhood onset
- Early-onset recurrent ischemic stroke and fever
- Livedo racemosa
- Low IgM
- Hypogammaglobulinaemia
- Lymphopenia
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Immunology





#### ANHYDROTIC ECTODERMODYSPLASIA WITH ID

# Available testing

| Centre      | Method         | Scope and range of test |         |                                | Targets                                  | TAT    |
|-------------|----------------|-------------------------|---------|--------------------------------|--|--------|
| Aberdeen    | NGS            | Whole gene screen       |         | SNVs, indels<br>Exon level CNV | IKBKG (NEMO), NFKBIA (IKBA)              | 56     |
| _           | nember<br>ting |                         |         | as indicate                    | d above                                  | 14     |
| Proforma re | equired?       | YES                     | GEN FOR | RM 215 Primary Immur           | nodeficiency Request form (see centre we | osite) |

#### Referral criteria

- Anhidrotic ectodermal dysplasia
- Various infections (bacteria, mycobacteria viruses & fungi)
- Colitis
- Variable defects of skin, hair & teeth.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





#### **ALPHA 1 ANTITRYPSIN DEFICIENCY**

#### Available testing

| Centre      | Method       | Scope and range of test |              | Targets     | TAT |
|-------------|--------------|-------------------------|--------------|-------------|-----|
| Edinburgh   | Sanger       | Whole gene screen       | SNVs, indels | SERPINA1    | 28  |
| Family me   | mber testing |                         | as indi      | cated above | 14  |
| Proforma re | equired?     | NO                      |              |             |     |

#### Referral criteria

- A1AT quantification AND phenotyping should be requested first (Biochemistry)
- Plasma concentration of alpha-1-antitrypsin below normal range, AND
  - Prolonged neonatal jaundice with an inconclusive alpha-1-antitrypsin phenotyping result, OR 2. Mutation analysis will inform reproductive choice, OR
  - Adult with cirrhosis or emphysema where a genetic diagnosis would influence management following an inconclusive alpha-1-antitrypsin phenotyping result
- Referrals for testing will be triaged by the Genomic Laboratory; testing should be targeted at those where a genetic or genomic diagnosis will guide management for the proband or family.

- Clinical Genetics
- Gastroenterology
- Hepatology
- Respiratory Medicine





#### **ASSOCIATION WITH GI INFLAMMATION**

# Available testing

| Centre      | Method   | Scope and range of test |                                   | Targets  | TAT   |
|-------------|----------|-------------------------|-----------------------------------|--|-------|
| Aberdeen    | NGS      | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | ADAM17, AICDA, AP3B1, B2M, BTK, CBL,<br>CD40LG, CORO1A, CTC1, CTPS1, CYBA, CYBB,<br>DCLRE1C, DOCK8, FERMT1, FOXP3, GUCY2C,<br>HPS1, HPS4, HPS6, ICOS, IFNGR1, IFNGR2,<br>IKBKG, IL10RA, IL2RA, ITGB2, MAGT1, NCF1,<br>NCF2, NCF4, NF1, PIK3CD, PIK3R1, PTEN,<br>PYCARD, SKIV2L, SLC37A4, STK4, TTC37,<br>VPS13B, WAS | 112   |
| _           | member   |                         |                                   | as indicated above   | 14    |
|             | ting     | VEC I                   | CEN FORM 245 Dr                   | imanulmmunadafiaianay Daguaat farm (aaa aantra wah   | oito) |
| Proforma re | equired? | YES                     | GEN FORM 215 Pr                   | imary Immunodeficiency Request form (see centre web  | site) |

#### Referral criteria

 Please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Rheumatology





#### **AUTOINFLAMMATORY DISORDERS**

## Available testing

| Centre      | Method | Scope and range of test |  | Targets   | TAT |
|-------------|--------|-------------------------|--|---|-----|
| Aberdeen    | NGS    | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV                              | ACP5, ADA2 (CECR1), ADAM17, ADAR1, AP1S3,<br>CARD14, COPA, IFIH1, IL1RN, IL36RN, LPIN2,<br>MEFV, MVK, NOD2, NLCR4, NLRP1, NLRP3,<br>NLRP12, OTULIN, PLCG2, POLA1, PSMB8,<br>PSTPIP1, RNASEH2A, RNASEH2B, RNASEH2C,<br>SAMHD1, SH3BP2, SLC29A3, TMEM173,<br>TNF1IP3, TNFAIP3, TNFRSF1A, TREX1, USP18 | 112 |
| Family r    |        |                         | ·  | as indicated above  | 14  |
| Proforma re |        | YES                     | GEN FORM 215 Primary Immunodeficiency Request form (see centre |   |     |

#### Referral criteria

- Clinical features suggestive of a monogenic cause of Autoinflammatory disorders.
- For specific Autoinflammatory disorders subpanels (Monogenic Autoinflammatory diseases, Recurrent inflammation, Systemic inflammation with urticarial rash, Others, Sterile inflammation predominant on the bone / joints, Sterile inflammation predominant on the skin, Type 1 interferonopathies), please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Immunology
- Rheumatology





#### **BACTERIAL AND PARASITIC INFECTIONS**

# Available testing

| Centre      | Method         | Scope and range of test |                                   | Targets  | TAT   |
|-------------|----------------|-------------------------|-----------------------------------|--|-------|
| Aberdeen    | NGS            | Whole gene screen       | SNVs, indels<br>Exon level<br>CNV | ACT1 (TRAF3IP1), APOL1, CARD9, HMOX1,<br>IRAK1, IRAK4, MYD88, NBAS, NCSTN, PSEN,<br>PSENEN, RANBP2, RPSA, STAT1, IL17F, IL17RA,<br>1L17RC, TIRAP | 112   |
|             | member<br>ting |                         |                                   | as indicated above   | 14    |
| Proforma re | equired?       | YES                     | GEN FORM 215 Pr                   | imary Immunodeficiency Request form (see centre web  | site) |

#### Referral criteria

- Clinical features suggestive of a monogenic cause of Bacterial and Parastic infections.
- For specific Bacterial and parasitic infections subpanels (Predisposition to invasive bacterial infections, Predisposition to parasitic and fungal infections, Hydradenitis suppurativa, Acute liver failure due to NBAS deficiency, Acute necrotising encephalopathy), please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





# BACTERIAL INFECTIONS, AUTOINFLAMMATION, AMYLOPECTINOSIS

# Available testing

| Centre      | Method   | Scope and range of test |         |                                | Targets                                   | TAT     |
|-------------|----------|-------------------------|---------|--------------------------------|---|---------|
| Aberdeen    | NGS      | Whole<br>scre           | U       | SNVs, indels<br>Exon level CNV | HOIL1 (RBCK1), HOIP1 (RNF31)              | 56      |
| Family r    |          |                         |         | as indicat                     | ed above                                  | 14      |
| Proforma re | equired? | YES                     | GEN FOR | RM 215 Primary Immu            | unodeficiency Request form (see centre we | ebsite) |

#### Referral criteria

- Clinical features suggestive of a monogenic cause of Bacterical infections, Autoinflammation, Amylopectinosis.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





#### **CALCIUM CHANNEL DEFECTS**

# Available testing

| Centre      | Method   | Scope and range of test |         |                                | Targets                              | TAT    |
|-------------|----------|-------------------------|---------|--------------------------------|--------------------------------------|--------|
| Aberdeen    | NGS      | Whole gene screen       |         | SNVs, indels<br>Exon level CNV | ORAI1, STIM1                         | 56     |
| Family r    |          |                         |         | as indicated ab                | oove                                 | 14     |
| Proforma re | equired? | YES                     | GEN FOR | RM 215 Primary Immunode        | ficiency Request form (see centre we | bsite) |

#### Referral criteria

- Autoimmunity
- EDA
- Non-progressive myopathy
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





#### **CHRONIC GRANULOMATOUS DISEASE**

## Available testing

| Centre                              | Method       | Scope and range of test |             |                                | Targets                                | TAT     |
|-------------------------------------|--------------|-------------------------|-------------|--------------------------------|--|---------|
| Aberdeen                            | NGS          | Whole                   | gene screen | SNVs, indels<br>Exon level CNV | CYBA, CYBB, NCF1, NCF2, NCF4           | 56      |
| Family men                          | nber testing |                         |             | as indicated                   | above                                  | 14      |
| Proforma required? YES GEN FORM 215 |              |                         | GEN FORM    | 215 Primary Immunod            | deficiency Request form (see centre we | ebsite) |

#### Referral criteria

- Early onset of severe and recurrent infections affecting initially the natural barriers of the organism (lungs, lymph nodes, skin), and eventually inner structures (liver, spleen, bones, brain, and +++ hepatic abscess).
- Autoinflammatory phenotype.
- IBD.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Immunology





# **COMBINED IMMUNODEFICIENCIES (CVID)**

# Available testing

| Centre                | Method   | Scope and range of test |                                   | Targets   | TAT   |
|-----------------------|----------|-------------------------|-----------------------------------|---|-------|
| Aberdeen              | NGS      | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | CD40LG, CD40, ICOS, CD3G, CD8A, ZAP70,<br>ZAP70, TAP1, TAP2, TAPBP, B2M, CIITA,<br>RFXANK, RFX5, RFXAP, IKZF1 (AB deficiency –<br>hypogammaglobulinemia), DOCK8, DOCK2,<br>RHOH, STK4, TRAC, LCK, ITK (EBV susceptibility),<br>MALT1, CARD11 (AR LOF), BCL10, IL21, IL21R,<br>TNFRSF4, IKBKB, MAP3K14, RELB, RELA, MSN,<br>TFRC | 56    |
| Family member testing |          |                         | as indicated above                | 14  |       |
| Proforma re           | equired? | YES                     | GEN FORM 215 Pr                   | imary Immunodeficiency Request form (see centre web   | site) |

#### Referral criteria

- Generally less profound than SCID.
- please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





#### **COMPLEMENT DEFICIENCIES**

# Available testing

| Centre      | Method         | Scope and            | I range of test                   | Targets  | TAT   |
|-------------|----------------|----------------------|-----------------------------------|--|-------|
| Aberdeen    | NGS            | Whole gene<br>screen | SNVs, indels<br>Exon level<br>CNV | C1QA, C1QB, C1QC, C1R, C1S, C2, C3, C4A,<br>C4B, C5, C6, C7, C8A, C8B, C8G, C9, CD55,<br>CD59, CFB, CFD, FCN3, MASP2, PFC (CFP),<br>SERPING1 | 56    |
| Family r    | member<br>ting |                      |                                   | as indicated above   | 14    |
| Proforma re | equired?       | YES GE               | N FORM 215 Prin                   | nary Immunodeficiency Request form (see centre web   | site) |

#### Referral criteria

- Clinical features suggestive of a monogenic cause of Complement deficiencies.
- For specific Complement deficiencies subpanels (Disseminated Neisserial infections, Recurrent pyogenic infections, SLE-like syndrome, Low susceptibility to infection), please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Immunology
- Rheumatology





#### **CONGENITAL THROMBOCYTOPENIA**

## Available testing

| Centre      | Method               | Scope and range of test |                   |   | Targets            | TAT |
|-------------|----------------------|-------------------------|-------------------|---|--------------------|-----|
| Aberdeen    | NGS                  | Whole gene<br>screen    |                   | SNVs, indels<br>Exon level<br>CNV           | ARPC1B, WAS, WIPF1 | 56  |
| _           | Family member as inc |                         |                   | as ind                                      | icated above       | 14  |
| Proforma re | equired? YES GEN FOR |                         | RM 215 Primary In | nmunodeficiency Request form (see centre we | ebsite)            |     |

#### Referral criteria

- · Recurrent bacterial and viral infections
- Bloody diarrhoea
- Excema
- Vasculitis
- For specific Congenital thrombocytopenia subpanels (Wiskott Aldrich Syndrome, WIP deficiency, ARPC1B deficiency), please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





#### **DEFECTS OF VITAMIN B12 AND FOLATE METABOLISM**

# Available testing

| Centre                   | Method               | Scope and range of test |        |                                   | Targets                                     | TAT          |
|--------------------------|----------------------|-------------------------|--------|-----------------------------------|---|--------------|
| Aberdeen                 | NGS                  | Whole gene<br>screen    |        | SNVs, indels<br>Exon level<br>CNV | MTHFD1, SLC46A1, TCN2                       | 56 or<br>112 |
|                          | Family member as inc |                         |        | icated above                      | 14  |              |
| Proforma required? YES G |                      |                         | GEN FO | RM 215 Primary In                 | nmunodeficiency Request form (see centre we | bsite)       |

#### Referral criteria

- Megablastic anaemia.
- Ig decreased.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





#### **DNA REPAIR DEFECTS**

# Available testing

| Centre             | Method               | Scope and range of test |               |                                   | Targets   | TAT    |
|--------------------|----------------------|-------------------------|---------------|-----------------------------------|---|--------|
| Aberdeen           | NGS                  |                         | e gene<br>een | SNVs, indels<br>Exon level<br>CNV | ATM, BLM, CDCA7, DNMT3B, GINS1,<br>HELLS, LIG1, MCM4, NBS1 (NBN), PMS2,<br>POLE1, POLE2, NSMCE3, ERCC6L2,<br>RNF168, ZBTB24 | 56     |
| ,                  | Family member as inc |                         | icated above  | 14                                |   |        |
| Proforma required? |                      | YES                     | GEN FOR       | RM 215 Primary In                 | nmunodeficiency Request form (see centre web  | osite) |

#### Referral criteria

- Clinical features suggestive of a monogenic cause of DNA repair defects
- For specific DNA repair defects subpanels (Ataxia telangiectasia, Nijmegen breakage syndrome, Bloom syndrome, PMS2 deficiency, Immunodeficiency with centromeric instability & facial anomalies, MCM4 deficiency, RNF168 deficiency, POLE1 deficiency, POLE2 deficiency, NSME3 deficiency, ERCC6L2 (Hebo) deficiency, Ligase 1 deficiency. GINS1 deficiency), please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





# **DYSKERATOSIS CONGENITA (DKC)**

# Available testing

| Centre      | Method   | Scope and range of test |                                   | Targets  | TAT    |
|-------------|----------|-------------------------|-----------------------------------|--|--------|
| Aberdeen    | NGS      | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | CTC1, DKC1,PARN, NOLA2 (NHP2), NOLA3<br>(NOP10), RTEL1, SAMD9, SAMD9L, SNM1B /<br>APOLLO (DCLRE1B), STN1, TERC, TERT,<br>TINF2, TPP1, WRAP53 | 112    |
| Family r    |          |                         | as indicated above                |  |        |
| Proforma re | equired? | YES                     | GEN FORM 215 Pr                   | imary Immunodeficiency Request form (see centre we   | bsite) |

#### Referral criteria

- Myelodysplasia
- Defective telomere maintenance
- Exclude other causes: Fanconi Anaemia, Diamond-Blackfan
- For specific Dyskeratosis congenita panels (Dyskeratosis congenital, Coats plus syndrome, Others), please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





#### **FAMILIAL HLH DUE TO PRF1 VARIANTS**

## Available testing

| Centre             | Method                             | Scope ar     | nd range o    | f test                         | Targets                                | TAT     |
|--------------------|------------------------------------|--------------|---------------|--------------------------------|--|---------|
| Aberdeen           | Sanger<br>MLPA                     | Whole<br>scr | e gene<br>een | SNVs, indels<br>Exon level CNV | PRF1                                   | 56      |
| Family mer testing | Family member as indicated testing |              |               |                                | above                                  | 14      |
| Proforma required? |                                    | YES          | GEN FOR       | RM 215 Primary Immuno          | deficiency Request form (see centre we | ebsite) |

### Referral criteria

- Fever
- Cytopenias
- Increased activated Tc
- Decreased to absent NK and CTL activities cytotoxicity.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website.

- Clinical Genetics
- Haematology
- Immunology





# HAEMOPHAGOCYTIC LYMPHOHISTOPCYTOSIS (HLH) & EBV SUSCEPTIBILITY

# Available testing

| Centre      | Method         | Scope                   | and range of test                 | Targets   | TAT   |
|-------------|----------------|-------------------------|-----------------------------------|---|-------|
| Aberdeen    | NGS            | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | AP3B1, AP3D1, CD27, CD70, CTPS1, DNASE2,<br>FAAP24, ITK, LYST, MAGT1, PRF1, PRKCD,<br>RAB27A, RASGRP1, RLTPR (CARMIL2), SH2DIA,<br>SLC29A3, STX11, STXBP2, UNC13D, XIAP | 56    |
| _           | member<br>ting |                         |                                   | as indicated above  | 14    |
| Proforma re | equired?       | YES                     | GEN FORM 215 Pri                  | imary Immunodeficiency Request form (see centre web   | site) |

#### Referral criteria

- Clinical features suggestive of a monogenic cause of HLH & EBV susceptibility.
- For specific Hemophagocytic Lymphohistocytosis HLH & EBV susceptibility panels (Chediak Higashi syndrome, Griscelli syndrome type 2, Hermansky Pudiak Syndrome type 10, H ermansky Pudiak Syndrome type 2, Familial HLH Syndromes, Susceptibility to EBV, EBV associated HLH), please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology
- Rheumatology





#### HENNEKAM-LYMPHANGIECTASIA-LYMPHEDEMA SYNDROME

# **Available testing**

| Centre                | Method                            | Scope and range of test |         |                                | Targets                                  | TAT    |
|-----------------------|-----------------------------------|-------------------------|---------|--------------------------------|--|--------|
| Aberdeen              | NGS                               | Whole gene screen       |         | SNVs, indels<br>Exon level CNV | CCBE1, FAT4                              | 56     |
|                       | Family member as indicate testing |                         |         | d above                        | 14                                       |        |
| Proforma required? YE |                                   | YES                     | GEN FOR | RM 215 Primary Immur           | nodeficiency Request form (see centre we | bsite) |

#### Referral criteria

- Lymphangiectasia and lymphedema with facial abnormalies and other dysmorphic features.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





# HEPATIC VENO-OCCLUSIVE DISEASE WITH IMMUNODEFICIENCY (VODI)

# Available testing

| Centre                             | Method | Scope and range of test |         |                                | Targets                                | TAT    |
|------------------------------------|--------|-------------------------|---------|--------------------------------|--|--------|
| Aberdeen                           | NGS    | Whole gene screen       |         | SNVs, indels<br>Exon level CNV | SP110                                  | 56     |
| Family member as indicates testing |        |                         |         | as indicated                   | above                                  | 14     |
| Proforma required? YES             |        | YES                     | GEN FOR | RM 215 Primary Immuno          | deficiency Request form (see centre we | bsite) |

#### Referral criteria

- Hepatic veno-occlusive disease.
- Pneumocystis jirovecii pneumonia
- CMV
- Candida
- Thrombocytopenia
- Hepatosplenomegaly
- Cerebrospinal leukodystrophy
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





#### **HEREDITARY AMYLOIDOSIS**

# Available testing

| Centre                | Method   | Scope and range of test |  |                                   | Targets   | TAT    |
|-----------------------|----------|-------------------------|--|-----------------------------------|---|--------|
| Aberdeen              | NGS      | Whole gene<br>screen    |  | SNVs, indels<br>Exon level<br>CNV | APOA1, APOA2, APOA4, APOC2,<br>APOC3, APOE, FGA, GSN, IL31RA, LYZ,<br>TTR, UNC13D | 112    |
| Family member testing |          |                         |  | as ind                            | icated above  | 14     |
| Proforma re           | equired? | YES GEN FORM 215 Prim   |  | RM 215 Primary In                 | nmunodeficiency Request form (see centre we                                       | bsite) |

## Referral criteria

- Clinical features suggestive of a monogenic cause of Hereditary Amyloidosis.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Rheumatology





# HEREDITARY ANGIOEDEMA, TYPES I & II

# Available testing

| Centre             | Method         | Scope and range of test |         |                                | Targets                                  | TAT    |
|--------------------|----------------|-------------------------|---------|--------------------------------|--|--------|
| Aberdeen           | Sanger<br>MLPA | Whole gene screen       |         | SNVs, indels<br>Exon level CNV | SERPING1                                 | 56     |
| Aberdeen           | NGS            | Whole gene<br>screen    |         | SNVs, indels<br>Exon level CNV | SERPING1, Factor XII, PLG, ANGPT1        | 56     |
| Family i           | nember<br>ting |                         |         |                                | d above                                  | 14     |
| Proforma required? |                | YES                     | GEN FOR | RM 215 Primary Immur           | nodeficiency Request form (see centre we | bsite) |

#### Referral criteria

- Hereditary angioedema
- Spontaneous activation of the complement pathway with consumption of C4/C2.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Immunology





# **HYPER IGE SYNDROMES (HIES)**

# Available testing

| Centre                                   | Method | Scope and range of test |         |                                | Targets                               | TAT    |
|--|--------|-------------------------|---------|--------------------------------|---------------------------------------|--------|
| Aberdeen                                 | NGS    | Whole gene<br>screen    |         | SNVs, indels<br>Exon level CNV | PGM3, SPINK5, STAT3                   | 56     |
| Family member as indicated above testing |        |                         |         |                                | pove                                  | 14     |
| Proforma required? YES                   |        | YES                     | GEN FOR | RM 215 Primary Immunode        | ficiency Request form (see centre web | osite) |

#### Referral criteria

- Clinical features suggestive of a monogenic cause of Hyper IgE syndromes (HIES).
- For specific Hyper IgE syndromes (AD-HIES / Job syndrome, Comel Netherton syndrome, PGM3 deficiency), please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





#### **HYPOGAMMAGLOBULINAEMIA**

# Available testing

| Centre                                  | Method         | Scope a                 | and range of test                 | Targets  | TAT          |
|---|----------------|-------------------------|-----------------------------------|--|--------------|
| Aberdeen                                | NGS            | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | ATP6AP1, BLNK, BTK, CD19, CD20 (MS4A1),<br>CD79A, CD79B, CD81, IGHM, IGLL1, IKZF1<br>(IKAROS), IRF2BP2, MOGS, NFKB1, PIK3CD,<br>PIK3R1, PTEN, TCF3, TNFRSF13B (TACI),<br>TNFRSF13C (BAFFR), TRNT1, TTC37, TWEAK<br>(TNFSF12) | 56 or<br>112 |
| Family r                                | member<br>ting |                         |                                   | as indicated above   | 14           |
| Proforma required? YES GEN FORM 215 Pri |                |                         | GEN FORM 215 Pri                  | imary Immunodeficiency Request form (see centre we   | ebsite)      |

#### Referral criteria

- IgG, IgA and / or IgM decreased
- Exclude second causes: drugs [Hx], myeloma [bone marrow], lymphoma. Ig loss (not hypo-lgM) in urine, gastro-intestinal or skin.
- For specific Hypogammaglobulinaemia subpanels (B absent, B>1% Common Variable Immunodeficiency phenotype), please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





#### **ID WITH MULTIPLE INTESTINAL ATRESIAS**

## Available testing

| Centre                | Method | Scope and range of test |   |                                | Targets | TAT |  |  |
|-----------------------|--------|-------------------------|---|--------------------------------|---------|-----|--|--|
| Aberdeen              | NGS    | Whole gene<br>screen    |   | SNVs, indels<br>Exon level CNV | TTC7A   | 56  |  |  |
| Family member testing |        | as indicated above      |   |                                |         |     |  |  |
| Proforma required?    |        | YES                     | GEN FORM 215 Primary Immunodeficiency Request form (see centre website) |                                |         |     |  |  |

#### Referral criteria

- Bacterial (sepsis), fungal, viral infections
- Multiple intestinal atresias, often with intrauterine polyhydramnios and early demise, some with SCIDphenotype.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





#### **IMMUNO-OSSEOUS DYSPLASIAS**

# Available testing

| Centre                | Method | Scope a            | and range of test   | Targets                                   | TAT |  |  |
|-----------------------|--------|--------------------|---|---|-----|--|--|
| Aberdeen              | NGS    | Whole gene screen  | SNVs, indels<br>Exon level CNV  | EXTL3, MYSM1, RMRP, RNU4ATAC,<br>SMARCAL1 | 56  |  |  |
| Family member testing |        | as indicated above |   |   |     |  |  |
| Proforma required?    |        | YES                | GEN FORM 215 Primary Immunodeficiency Request form (see centre website) |   |     |  |  |

#### Referral criteria

- Clinical features suggestive of a monogenic cause of Immuno-Osseous Dysplasias
- For specific Immuno-osseous dysplasias subpanels (Cartilage Hair Hypoplasia, Schimke syndrome, MYSM1 deficiency, MOPD1 deficiency, EXLT3 deficiency), please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Immunology





## INTERFERONOPATHY / SLS / AGS / COMPLEMENT

## Available testing

| Centre             | Method   | Scope and range of test |                                   | Targets  | TAT    |
|--------------------|----------|-------------------------|-----------------------------------|--|--------|
| Aberdeen           | NGS      | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | ACP5, ADAM17, C1QA, C1QB, C1QC, C1R, C2,<br>C3, C5, C6, C7, C8A, C8B, C9, CFH, CFHR5,<br>CFI, CFP, DNASE1, DNASE1L3, IFIH1, IRF8,<br>RASGRP1, RNASEH2A, RNASEH2B,<br>RNASEH2C, SAMHD1, SNORD118, TREX1,<br>USP18 | 112    |
| Family mer testing | nber     |                         | •                                 | as indicated above   | 14     |
| Proforma re        | equired? | YES                     | GEN FORM 215 Pri                  | mary Immunodeficiency Request form (see centre we  | bsite) |

## Referral criteria

- Clinical features suggestive of a monogenic cause of Interferonopathy/ SLS / AGS / Complement disorders.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website.

- Clinical Genetics
- Immunology
- Rheumatology





## **KABUKI SYNDROME**

## Available testing

| Centre   | Method | Scope and range of test |                      |  | Targets             | TAT |
|--|--------|-------------------------|----------------------|--|---------------------|-----|
| Aberdeen                                       | NGS    | Whole gene screen       |                      | SNVs, indels<br>Exon level CNV           | KDM6A, KMT2D (MLL2) | 56  |
| Family r                                       |        |                         |                      | as indicate                              | d above             | 14  |
| Proforma required? YES GEN FORM 215 Primary Im |        |                         | RM 215 Primary Immur | nodeficiency Request form (see centre we | bsite)              |     |

## Referral criteria

- Typical facial abnormalies
- Cleft or high arched palate
- Skeletal abnormalities
- Short stature
- Intellectual disability
- Congenital heart defects
- Recurrent infections (otitis media, pneumonia) in 50% of patients
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





# MENDELIAN SUSCEPTIBILITY TO MYCOBACTERIAL DISEASE (MSMD) AND VIRAL INFECTION

## Available testing

| Centre       | Method                             | Scope and range of test |            |                                | Targets  | TAT |
|--------------|------------------------------------|-------------------------|------------|--------------------------------|--|-----|
| Aberdee<br>n | NGS                                | Whole ge<br>screer      |            | SNVs, indels<br>Exon level CNV | CXCR4 (WHIM), CYBB, FCGR3A, IFIH1,<br>IFNAR2, IFNGR1, IFNGR2, IL12B,<br>IL12RB1, IRF3, IRF7, IRF8, ISG15,<br>JAK1, RORC, STAT1, STAT2, TBK1,<br>TICAM1 (TRIF), TLR3, TMC6, TMC8,<br>TRAF3, TYK2, UNC93B1 | 56  |
| ,            | Family member as indicates testing |                         | ated above | 14                             |  |     |
| Proforma re  | equired?                           | YE GEN<br>S             | FORM 2     | 215 Primary Immuno             | deficiency Request form (see centre website)   | )   |

#### Referral criteria

- Clinical features suggestive of a monogenic cause of Mendelian Susceptibility to Mycobacterial Disease (MSMD) and Viral infection.
- For specific Mendelian Susceptilibility to Mycobacterial disease (MSMD) and viral infection subpanels (MSMD sever phenotypes, MSMD moderate phenotypes, Epidermodysplasia verruciformis (HPV), Predominant susceptibility to viral infection Herpes simplex Encephalitis, Predisposition to severe viral infection), please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





## **MISCELLANEOUS AUTOINFLAMMATORY CONDITIONS**

## Available testing

| Centre                | Method   | Scope and range of test |                                   | Targets  | TAT   |
|-----------------------|----------|-------------------------|-----------------------------------|--|-------|
| Aberdeen              | NGS      | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | ADA2, AIRE, AP1S3, CASP10, CASP8, COPA, COL7A1, CPT2, FAS, FASLG, FLNA, HTR1A, IL10, IL10RB, IL12B, IL12RB1, IL1RN, IL36RN, ISG15, LACC1, LPIN2, LRBA, LYN, MASP2, MAT2A, MBL2, MEFV, MVK, MYD88, NLRC4, NLRP1, NLRP12, NLRP3, NLRP6, NLRP7, NOD2, NRAS, OTULIN, PRKCD, PLCG2, POMP, PRG4, PSMA3, PSMB4, PSMB8, PSMB9, PSTPIP1, RAG1, RANBP2, SCN9A, SERPING1, SH2D1A, SH3BP2, TMEM173, TNFAIP3, TNFRSF11A, TNFRSF1A, TRAP1, TRNT1, USB1, WDR1 | 112   |
| Family member testing |          |                         | as indicated above                | 14   |       |
| Proforma re           | equired? | YES                     | GEN FORM 215 I                    | Primary Immunodeficiency Request form (see centre web  | site) |

## Referral criteria

• Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website.

- Clinical Genetics
- Immunology
- Rheumatology





## **NEUTROPENIA**

## Available testing

| Centre      | Method         | Scope and range of test |   | Targets  | TAT |  |  |
|-------------|----------------|-------------------------|---|--|-----|--|--|
| Aberdeen    | NGS            | Whole genderscreen      | e SNVs, indels<br>Exon level<br>CNV                                       | C160RF57 (USB1), CLPB, COH1 (VPS13B),<br>CSF3R, DNAJC21, ELANE, G6PC3, G6PT1<br>(SLC37A4), GFI1, HAX1, HYOU1, JAGN1,<br>LAMTOR2, MKL1 (MRTFA), SBDS, SMARCD2,<br>TAZ, VPS45, WAS, WDR1 | 56  |  |  |
| Family r    | nember<br>ting |                         | as indicated above  |  |     |  |  |
| Proforma re | equired?       | YES (                   | S GEN FORM 215 Primary Immunodeficiency Request form (see centre website) |  |     |  |  |

#### Referral criteria

- Clinical features suggestive of a monogenic cause of Neutropenia.
- For specific Neutropenia subpanels (Schwachman-Diamond Syndrome, G6PC3 deficiency, Glycogen storage diasease type 1b, Cohen syndrome, Barth Syndrome, Clericuzio syndrome, VPS45 deficiency, P14/LAMTOR2 deficiency, JAGN1 deficiency, 2-Methylglutaconic aciduria, SMARCD2 deficiency, WDR1 deficiency, HYOU1 deficiency, No syndrome associated), please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





## **OTHER ANTIBODY DEFICIENCIES**

## Available testing

| Centre      | Method         | Scope and range of test |         |                                | Targets                                    | TAT     |
|-------------|----------------|-------------------------|---------|--------------------------------|--|---------|
| Aberdeen    | NGS            | Whole gene screen       |         | SNVs, indels<br>Exon level CNV | AICDA, CARD11, IGKC, INO80, MSH6,<br>UNG   | 56      |
| Family r    | member<br>ting |                         |         | as indic                       | ated above                                 | 14      |
| Proforma re | equired?       | YES                     | GEN FOI | RM 215 Primary Imr             | nunodeficiency Request form (see centre we | ebsite) |

## Referral criteria

- Clinical features suggestive of a monogenic cause of Other Antibody Deficiencies.
- For specific Other antibody deficiencies subpanels (Hyper IgM Syndromes; Isotype, Light Chain, or Functional Deficiencies; High Bc), please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





## PURINE NUCLEOSIDE PHOSPHORYLASE DEFICIENCY

## Available testing

| Centre   | Method | Scope and range of test |                                       |                                | Targets | TAT |
|--|--------|-------------------------|---------------------------------------|--------------------------------|---------|-----|
| Aberdeen   | NGS    |                         | e gene<br>een                         | SNVs, indels<br>Exon level CNV | PNP     | 56  |
| Family r   |        |                         |                                       | as indicated ab                | pove    | 14  |
| Proforma required? YES GEN FORM 215 Primary Immunode |        |                         | eficiency Request form (see centre we | ebsite)                        |         |     |

#### Referral criteria

- Autoimmune hemolytic anaemia
- Neurological impairment.
- Hypouricemia.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





# **SEVERE COMBINED IMMUNODEFICIENCY (SCID)**

## Available testing

| Centre      | Method         | Scope a             | and range of test  | Targets  | TAT |  |
|-------------|----------------|---------------------|--|--|-----|--|
| Aberdeen    | NGS            | Whole ger<br>screen | SNVs, indels<br>Exon level<br>CNV                                      | ADA, AK2, CD3D, CD3E, CD247, COR01A,<br>DCLRE1C (ARTEMIS), FOXN1, IL2RG, IL7R,<br>JAK3, LIG4, NHEJ1, PRKDC, PTPRC, RAG1,<br>RAG2 | 56  |  |
| Family r    | member<br>ting |                     | as indicated above   |  |     |  |
| Proforma re | equired?       | YES                 | S GEN FORM 215 Primary Immunodeficiency Request form (see centre websi |  |     |  |

## Referral criteria

- CD3 T cell lymphopenia: CD3+ T cells <300/µl.
- For specific SCID subpanels (SCID T-B+ CD19 normal, SCID T-B- CD19 low), please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





## **STAT5B DEFICIENCY**

## Available testing

| Centre      | Method   | Scope and range of test |         |                                | Targets                               | TAT     |
|-------------|----------|-------------------------|---------|--------------------------------|---------------------------------------|---------|
| Aberdeen    | NGS      | Whole gene screen       |         | SNVs, indels<br>Exon level CNV | STAT5B                                | 56      |
| Family r    |          |                         |         | as indicated ab                | oove                                  | 14      |
| Proforma re | equired? | YES                     | GEN FOI | RM 215 Primary Immunode        | eficiency Request form (see centre we | ebsite) |

#### Referral criteria

- Growth-hormone insensitive dwarfism
- Dysmorphic features
- Eczema
- Lymphocytic interstitial pneumonitis
- Autoimmunity
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





## **STROKE**

# Available testing

| Centre   | Method | Scope and range of test |  |                                | Targets                                | TAT |
|--|--------|-------------------------|--|--------------------------------|--|-----|
| Aberdeen   | NGS    | Whole g                 | jene screen                            | SNVs, indels<br>Exon level CNV | CBS, CST3, GLA, HTRA1, NOTCH3,<br>ADA2 | 56  |
| Family mer testing                                   | nber   |                         | as indicated above                     |                                |  |     |
| Proforma required? YES GEN FORM 215 Primary Immunode |        |                         | odeficiency Request form (see centre w | ebsite)                        |  |     |

## Referral criteria

- Clinical features suggestive of a monogenic cause of Stroke.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website.

- Clinical Genetics
- Immunology
- Rheumatology





## SYNDROMES ASSOCIATED WITH AUTOIMMUNITY AND OTHERS

#### Available testing

| Centre      | Method         | Scope and range of test |  | Targets  | TAT |  |
|-------------|----------------|-------------------------|--|--|-----|--|
| Aberdeen    | NGS            | Whole<br>gene<br>screen | SNVs, indels<br>Exon level CNV   | AIRE, BACH2, CASP8, CASP10, CTLA4, FADD,<br>FOXP3 (IPEX), IL2RA, IL10, IL10RA, IL10RB,<br>ITCH, JAK1, LRBA, NFAT5, PEPD, STAT3,<br>TNFRSF6 (FAS), TNFSF6 (FASLG), TPP2,<br>ZAP70 | 56  |  |
| Family r    | nember<br>ting |                         | •  | as indicated above   | 14  |  |
| Proforma re | equired?       | YES (                   | ES GEN FORM 215 Primary Immunodeficiency Request form (see centre well |  |     |  |

#### Referral criteria

- Clinical features suggestive of a monogenic cause of Syndromes associated with autoimmunity and others.
- For specific Syndrome associated with Autoimmunity and others subpanels (Syndromes with autoimmunity with increased CD4-CD8-TCRα/β ALPS, Syndromes with autoimmunity with occasionally increased CD4-CD8-TCRα/β, Syndromes with autoimmunity without increased CD4-CD8-TCRα/β and without regulatory T Cell defects, Syndromes with autoimmunity without increased CD4-CD8-TCRα/β and with regulatory T Cell defects, Immune dysregulation with Colitis), please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





# SYNDROMES ASSOCIATED WITH CONGENITAL DEFECTS OF PHAGOCYTES Available testing

| Centre      | Method   | Scope ar   | nd range of test   | Targets  | TAT |  |
|-------------|----------|------------|--|--|-----|--|
| Aberdeen    | NGS      | Whole gene | SNVs, indels<br>Exon level<br>CNV                                | ACTB, CEBPE, CSFR2A, CSFR2B, CTSC,<br>FERMT3 (LADIII), FPR1, GATA2, G6PD, ILGB2<br>(LAD1), RAX2, SLC35C1 (LADII) | 56  |  |
| Family r    |          |            | ·  | as indicated above   | 14  |  |
| Proforma re | equired? | YES G      | GEN FORM 215 Primary Immunodeficiency Request form (see centre v |  |     |  |

#### Referral criteria

- Clinical features suggestive of a monogenic cause of Syndromes associated with Congenital Defects of Phagocytes.
- For specific Syndrome associated with congenital defects of phagocytes subpanels (Papillion-Lefevre, Localised juvenile periodontitis, β-Actin, Leukocyte adhesion deficiency / LAD, MonMac syndrome, Specific granule deficiency, Pulmonary alveolar proteinosis, RAC2 deficiency, G6PD deficiency Class 1), please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





## THYMIC DEFECTS WITH CONGENITAL ANOMALIES

## Available testing

| Centre      | Method                       | Scope and range of test |               |                                | Targets                                  | TAT     |
|-------------|------------------------------|-------------------------|---------------|--------------------------------|--|---------|
| Aberdeen    | NGS                          |                         | e gene<br>een | SNVs, indels<br>Exon level CNV | CHD7, SEMA3E, TBX1, FOXN1                | 56      |
| Family r    |                              |                         |               | as indicat                     | ed above                                 | 14      |
| Proforma re | Proforma required? YES GEN F |                         | GEN FO        | RM 215 Primary Immu            | unodeficiency Request form (see centre w | ebsite) |

#### Referral criteria

- Clinical features suggestive of a monogenic cause of Thymic Defects with Congenital Anomalies.
- For specific Thymic defects with Congenital anomalies subpanels (TBX1 deficiency, Winged Helix nude FOXN1 deficiency), please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507). Please note, Charge syndrome (CHD7) screening specifically is performed in Glasgow laboratory.

- Clinical Genetics
- Haematology
- Immunology





## **VASCULOPATHY**

# Available testing

| Centre      | Method   | Scope and range of test |                                   | Targets  | TAT |  |  |
|-------------|----------|-------------------------|-----------------------------------|--|-----|--|--|
| Aberdeen    | NGS      | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | ACTA2, BMPR2, COL3A1, COL4A1, COL5A1,<br>COL5A2, EFEMP2, ELN, FBN1, FBN2, FOXE3,<br>GUCA1B, LMNA, LOX, MFAP5, MYH11, MYLK,<br>NOTCH1, PLOD1, PRKG1, RHOD, RNF213, SKI,<br>SLC2A10, SMAD2, SMAD3, SMAD4, STX11,<br>STXBP2, TGFB2, TGFB3, TGFBI, TGFBR1,<br>YY1AP1 | 56  |  |  |
| Family mer  | nber     |                         | as indicated above                |  |     |  |  |
| testing     |          |                         |                                   |  |     |  |  |
| Proforma re | equired? | YES                     | GEN FORM 215 Pr                   | EN FORM 215 Primary Immunodeficiency Request form (see centre website  |     |  |  |

## Referral criteria

- Clinical features suggestive of a monogenic cause of Vasculopathy.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website.

- Clinical Genetics
- Immunology
- Rheumatology





# **VERY EARLY ONSET INFLAMMATORY BOWEL DISEASE (VEO-IBD)**

## Available testing

| Centre      | Method   | Scope and range of test |                                   | Targets   | TAT   |  |
|-------------|----------|-------------------------|-----------------------------------|---|-------|--|
| Aberdeen    | NGS      | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | ADAM17, AICDA, CD40LG, BTK, CD3G, ZAP70, WAS, CYBA, CYBB, NCF1, NCF2, NCF4, DOCK8, EPCAM (Sanger only), FOXP3, GUCY2C, HPS1, HPS4, HPS6, ADA, IL2RG, LIG4, DCLRE1C, RAG2, IL10, II10RA, IL10RB, ITGB2, LRBA, ICOS, PIK3R1, PLCG2, RET, SH2D1A, XIAP, SKIV2L, TTC37, SLC37A4, SKIV2L, STAT1, STAT3, STXBP2 | 112   |  |
| Family men  | nber     |                         | as indicated above                |   |       |  |
| Proforma re | equired? | YES                     | GEN FORM 215 Pr                   | imary Immunodeficiency Request form (see centre web   | site) |  |

## Referral criteria

- Clinical features suggestive of a monogenic cause of Very Early Onset Inflammatory Bowel Disease (VEO-IBD).
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website.

- Clinical Genetics
- Immunology
- Rheumatology





## **VICI SYNDROME**

# Available testing

| Centre      | Method   | S   | cope and      | range of test                  | Targets                                | TAT     |
|-------------|----------|-----|---------------|--------------------------------|--|---------|
| Aberdeen    | NGS      |     | e gene<br>een | SNVs, indels<br>Exon level CNV | EPG5                                   | 56      |
| Family r    |          |     |               | as indicated                   | d above                                | 14      |
| Proforma re | equired? | YES | GEN FOR       | RM 215 Primary Immun           | odeficiency Request form (see centre w | ebsite) |

#### Referral criteria

- Agenesis of the corpus callosum
- Cataracts
- Cardiomyopathy
- Skin hypopigmentaon
- Intellectual disability
- Microcephaly
- CMC
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





## X-LINKED CGD

## Available testing

| Centre      | Method         | Scope and range of test |            |                                | Targets                                 | TAT     |
|-------------|----------------|-------------------------|------------|--------------------------------|---|---------|
| Aberdeen    | Sanger<br>MLPA | Whole go                | ene screen | SNVs, indels<br>Exon level CNV | CYBB                                    | 56      |
| Family r    |                |                         |            | as indicated                   | d above                                 | 14      |
| Proforma re | equired?       | YES                     | GEN FORI   | M 215 Primary Immun            | odeficiency Request form (see centre we | ebsite) |

#### Referral criteria

- Suggestive of X linked transmission.
- Early onset of severe and recurrent infections affecting initially the natural barriers of the organism (lungs, lymph nodes, skin), and eventually inner structures (liver, spleen, bones, brain, and +++ hepatic abscess).
- Autoinflammatory phenotype.
- IBD.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Immunology





# INHERITED CANCER

## COWDEN SYNDROME / PTEN HAMARTOMA TUMOUR SYNDROME (PHTS)

# Available testing

| Centre              | Method   | Scope and range of test |                   |                                | Targets | TAT |
|---------------------|----------|-------------------------|-------------------|--------------------------------|---------|-----|
| Aberdeen<br>Glasgow | NGS      | Whole o                 | -                 | SNVs, indels<br>Exon level CNV | PTEN    | 56  |
| Family r            |          |                         | as indicated abov |                                | ve      | 14  |
| Proforma re         | equired? | NO                      |                   |                                |         |     |

#### Referral criteria

- Proband and / or family history meets one of the following criteria:
  - Mucocutaneous lesions comprising
    - ≥ 6 facial papules, of which ≥ 3 are trichilemmoma
    - Cutaneous facial papules AND oral mucosal papillomatosis
    - Oral mucosal papillomatosis AND acral keratosis
    - ≥6 palmoplantar keratosis
  - Cerebellar dysplastic gangliocytoma (Adult Lhermitte-Duclos disease)
  - ≥2 major criteria of which should be macrocephaly
  - o ≥1 major criteria and ≥ 1 PTEN-HTS-related mucocutaneous lesion
  - o ≥1 major and ≥ 3 minor criteria
  - o Macrocephaly ≥99th centile AND ≥ 1 minor criteria
  - ≥ 1 PHTS-related mucocutaneous lesion
  - ≥ 4 minor criteria
  - ≥ 1 major criteria, AND ≥ 2 first / second degree relatives each with:
    - ≥ 1 major criteria, OR ≥ 1 PHTS-related mucocutaneous lesion,
       OR
    - ≥ 2 minor criteria (multiple cases of breast cancer are not eligible for inclusion)

- Clinical Genetics
- Dermatology
- Neurology
- Paediatrics





#### **DICER1 SYNDROME**

## Available testing

| Centre                | Method   | Scope and range of test |              | Targets | TAT |
|-----------------------|----------|-------------------------|--------------|---------|-----|
| Glasgow               | Sanger   | Whole gene screen       | SNVs, indels | DICER1  | 56  |
| Family member testing |          |                         | as indicate  | d above | 14  |
| Proforma re           | equired? | NO                      |              |         |     |

#### Referral criteria

- Testing of affected individual (proband) where the individual has one of the following diagnoses:
  - Pleuropulmonary blastoma or Lung cyst(s) in childhood, especially if multi-septated, multiple or bilateral; Thoracic, uterine, cervical or ovarian embryonal rhabdomyosarcoma; Cystic nephroma; Genitourinary sarcoma including undifferentiated sarcoma in childhood; Ovarian Sertoli Leydig tumour; Gynandroblastoma; Genitourinary/gynaecologic neuroendocrine tumors; Childhood-onset multinodular goitre or differentiated thyroid cancer (papillary or follicular); Ciliary body medulloepithelioma; Nasal chondromesenchymal hamartoma; Pineoblastoma; Pituitary blastoma, OR
- Testing of affected individual where there is a combination of two of the following diagnoses, either both in one affected individual or in two affected first degree relatives;
  - Lung cyst(s) in adults; Wilms tumour; Multinodular goitre or differentiated thyroid cancer; Embryonal rhabdomyosarcoma other than thoracic or gynaecologic; Poorly differentiated neuroendocrine tumour; Undifferentiated sarcoma; Macrocephaly
- NOTE: Where testing is being performed for hereditary colorectal cancer test criteria, the Edinburgh laboratory can perform testing for DICER1 using the same panel. Please contact the laboratory to discuss if required.

## Requesting specialties





## **FAMILIAL MELANOMA**

## Available testing

| Centre                 | Method         | Scope and range of test |               |                        | Targets  | TAT |  |  |
|------------------------|----------------|-------------------------|---------------|------------------------|--|-----|--|--|
| Glasgow                | NGS            |                         | e gene<br>een | SNVs, indels           | BAP1, BRCA2, CDKN2A, CDK4,<br>POT1             | 56  |  |  |
| Family r               | nember<br>ting |                         |               | as indicated           | above  | 14  |  |  |
| Proforma required? YES |                |                         | Inherited     | cancer proforma (see c | Inherited cancer proforma (see centre website) |     |  |  |

#### Referral criteria

- Testing of phenotypically affected individual (proband) where the individual +/-family history meets one of the following criteria. The proband has:
  - o ≥2 melanoma\* age < 30 OR
  - Melanoma\* AND >/= 2 relatives (first / second / third degree) with melanoma and/or melanoma in situ OR
  - Melanoma AND >/= 1 first degree relative with melanoma; one individual has multiple melanomas in situ OR
  - 1 Melanoma OR melanoma and atypical moles AND >/=1 first degree relative with pancreatic cancer < 60 OR</li>
  - Atypical moles AND >/= 2 relatives (first / second degree relatives) with melanoma
- NOTE: Melanoma includes melanoma in situ

- Clinical Genetics
- Oncology





# GORLIN SYNDROME (BASAL CELL NEVUS SYNDROME)

## Available testing

| Centre      | Method      | Scope and range of test |                                | Targets     | TAT |
|-------------|-------------|-------------------------|--------------------------------|-------------|-----|
| Glasgow     | NGS<br>MLPA | Whole gene screen       | SNVs, indels<br>Exon level CNV | PTCH1, SUFU | 56  |
| Family i    |             |                         | as indica                      | ited above  | 14  |
| Proforma re | equired?    | NO                      |                                |             |     |

#### Referral criteria

- Living individual affected (proband) where the individual history meets:
  - o ≥ 1 major criteria OR
  - o ≥ 2 minor criteria
- Major criteria:
  - Lamellar (sheet-like) calcification of the falx or clear evidence of calcification in an individual younger than age 20 years
  - o Jaw keratocyst: odontogenic keratocyst histologically
  - Palmar/plantar pits (two or more)
  - o SHH medulloblastoma, confirmed on tumour testing
  - Multiple basal cell carcinomas (BCCs) (>5 in a lifetime) or BCC before age 30 years
- Minor criteria:
  - Childhood medulloblastoma where SHH pathway in tumour has not been investigated (also called primitive neuroectodermal tumor [PNET])
  - Lympho-mesenteric or pleural cysts
  - Macrocephaly (OFC >97th centile)
  - Cleft lip/palate
  - Vertebral/rib anomalies observed on chest x-ray and/or spinal x-ray;
     bifid/splayed/extra ribs; bifid vertebrae
  - Preaxial or postaxial polydactyly
  - Ovarian/cardiac fibromas
  - Ocular anomalies (cataract, developmental defects, and pigmentary changes of the retinal epithelium)

## Requesting specialties





## HEREDITARY BREAST CANCER SYNDROME

## Available testing

| Centre              | Method         | Scope a                 | and range of test                 | Targets   | TAT |  |  |
|---------------------|----------------|-------------------------|-----------------------------------|---|-----|--|--|
| Aberdeen<br>Glasgow | NGS            | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | BRCA1, BRCA2, TP53, PTEN, PALB2, STK11,<br>CHEK2, ATM, RAD51C, RAD51D | 56  |  |  |
| ,                   | member<br>ting |                         |                                   | as indicated above  |     |  |  |
| Proforma re         | equired?       | YES                     | Glasgow laboratory                | only (see centre website)   |     |  |  |

## Referral criteria

Living affected individual with breast cancer who meets ONE of the following criteria:

- Breast Cancer diagnosed <40 years</li>
- Bilateral breast cancer, both <60 years
- Triple negative breast cancer, <60 years
- Male breast cancer, any age
- Breast cancer and a first degree relative with breast cancer, both diagnosed before the age of 45 years

- Breast Surgeons
- Clinical Genetics
- Oncology





## HEREDITARY BREAST / OVARIAN CANCER SYNDROME

#### Available testing

| Centre              | Method         | Scope and range of test |                                   | Targets   | TAT |
|---------------------|----------------|-------------------------|-----------------------------------|---|-----|
| Aberdeen<br>Glasgow | NGS            | Whole gene screen       | SNVs, indels<br>Exon level<br>CNV | BRCA1, BRCA2, TP53, PTEN, PALB2, STK11,<br>RAD51C, RAD51D, BRIP1, MSH2, MSH6,<br>MLH1, CHEK2, ATM | 56  |
| Family r            | nember<br>ting |                         |                                   | as indicated above  | 14  |
| Proforma re         | equired?       | YES                     | Glasgow laboratory                | only (see centre website)   |     |

Referral criteria for affected individualLiving affected individual who meets ONE of the following criteria:

- Breast and Ovarian cancer, any age
- Breast cancer (meeting breast panel criteria) with family history of ovarian cancer
- High-grade epithelial ovarian cancer, any age with a family history of breast cancer

## Requesting specialties

- Clinical Genetics
- Oncology

# Referral criteria FOR UNAFFECTED INDIVIDUAL WITH A FAMILY HISTORY OF BREAST CANCER

Living unaffected individual who meets ONE of the following criteria:

- Manchester score is ≥19, or their probability of germline pathogenic/likely pathogenic variant on CanRisk is ≥10% AND
- No affected family member or tumour sample available to test
- Combined pathology-adjusted Manchester score of ≥15 or single gene pathology-adjusted Manchester score of ≥10 or a CanRisk score of ≥10%

## Requesting specialties





## **HEREDITARY BREAST / OVARIAN CANCER SYNDROME:**

#### **Founder Variants ONLY**

## Available testing

| Centre      | Method         | Scope and range of test |                                   | Targets   | TAT |
|-------------|----------------|-------------------------|-----------------------------------|---|-----|
| Aberdeen    | NGS            | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | Ashkenazi Jewish: <i>BRCA1</i> c.68_69del,<br>c.5266dup & <i>BRCA2</i> c.5946del<br>Polish: <i>BRCA1</i> c.68_69del, c.181T>G,<br>c.4035del, c.5266dup<br>Orkney: <i>BRCA1</i> c.5207T>C p.(Val1736Ala) | 14  |
| Family r    | member<br>ting |                         |                                   | as indicated above  | 14  |
| Proforma re | equired?       | YES                     | I                                 |   |     |

#### Referral criteria

Living affected individual with:

- Breast Cancer <50 years OR</li>
- Manchester Score ≥10 or ≥5% mutation probability on CanRisk\*
- AND is from one of the following founder populations
  - Ashkenazi Jewish
  - o Poland
  - Orkney specific variant

Living unaffected individual who meets ONE of the following criteria and where no living affected is available to test:

- FDR with Breast Cancer <50 years OR
- Manchester Score ≥15 or ≥10% mutation probability on CanRisk\*
- AND is from one of the following founder populations
  - o Ashkenazi Jewish
  - Poland
  - Orkney specific variant

\*Undertake full panel test if meet criteria for breast and/or ovarian panel (see referral criteria)

## Requesting specialties





# **HEREDITARY BREAST / OVARIAN CANCER SYNDROME: deceased TESTING**Available testing

| Centre             | Method                | Scope an          | d range of test    | Targets      | TAT |  |  |
|--------------------|-----------------------|-------------------|--------------------|--------------|-----|--|--|
| Aberdeen           | NGS                   | Whole gene screen | SNVs, indels       | BRCA1, BRCA2 | 56  |  |  |
|                    | Family member testing |                   | as indicated above |              |     |  |  |
| Proforma required? |                       | YES               |                    |              |     |  |  |

## Referral criteria

- Germline testing can be carried out in a deceased relative affected with breast or ovarian cancer, if there is
  - o A tissue sample available for DNA extraction, AND
  - Pathology-adjusted Manchester score ≥17 or <u>CanRisk score</u> ≥15%, AND
  - No living affected individual is available for genetic testing

# Requesting specialties





## HEREDITARY COLORECTAL CANCER, LYNCH SYNDROME AND POLYPOSIS

## Available testing

| Centre                | Method |                         | Scope and range of test Targets   |   |    |  |  |
|-----------------------|--------|-------------------------|---|---|----|--|--|
| Edinburgh             | NGS    | Whole<br>gene<br>screen | SNVs, indels CNV analysis (MLPA) – Polyposis referrals: APC, MUTYH (selected exons), and GREM1 (upstream region) Lynch referrals and patient dx CRC <45yrs: MLH1, MSH2, MSH6 and EPCAM (selected exons) | APC, BMPR1A, MBD4, MLH1, MSH2, MSH6, MUTYH, NTHL1, PMS2, POLD1 (exons 4-12), POLE (exons 3-13), PTEN, RNF43, SMAD4, STK11 | 56 |  |  |
| Family member testing |        |                         | as indicated above  |   |    |  |  |
| Proforma re           | J      | YES                     | Colorectal cancer gene panel proforma (see centre wel   | bsite)  | ı  |  |  |

#### Referral criteria

- Clinical Criteria for germline testing in a living individual affected by
  - Diagnosed with colorectal cancer aged <45, irrespective of the dMMR</li> status of the tumour OR
  - Diagnosed with a dMMR tumour under age 70 where results of BRAF and/or MLH1 hypermethylation testing suggest Lynch syndrome\* OR
  - Diagnosed with a Lynch-related cancer\*\* and comes from a modified Amsterdam criteria (≥ 3 cases of Lynch-related cancer over ≥2 generations with ≥1 case diagnosed ≤50 years) positive family irrespective of the dMMR status of the tumour OR
  - Wimmer score =>3\*\*\*
  - Diagnosed with colorectal cancer <60 with ≥5 polyps</li>
- Clinical criteria for germline testing in a deceased individual affected by cancer:
  - o The individual +/- family history meets one of the above criteria, AND
  - Appropriate tissue is available (tumour or normal), AND
  - No living affected individual is available for genetic testing.
- Clinical Criteria for germline testing in an unaffected individual:
- First degree relative affected with Lynch-related cancer, AND





- Family history of colorectal cancer/Lynch-related cancers reaches Modified Amsterdam Criteria (≥3 cases over ≥2 generations with ≥1 case affected ≤50 years), AND
- Tumour sample analysis from affected family member has been attempted and is not possible, failed, indeterminate or indicates MMR deficiency (via IHC or MSI), AND
- Somatic sequencing is not possible, or failed, AND
- No living affected individual is available for genetic testing
- NOTE: The majority of reported cancers in the family, including that of the patient being tested if relevant, should have been confirmed where possible
- \* Where MLH1 promoter hypermethylation has been identified in tumour, testing of normal tissue or blood for constitutional MLH1 promoter hypermethylation can be offered in families where MLH1 promotor methylation has been identified in >1 affected individual with colorectal cancer ≤ 60. (Performed in Aberdeen, Dundee, Edinburgh MP and Glasgow laboratories).
- \*\*Lynch-related cancers include but are not restricted to: Colorectal, Endometrial, Endocervical, Epithelial ovarian, Urothelial (urethra, bladder TCC, ureters, renal pelvis), Pancreatic, Bile duct (cholangiocarcinoma), Prostate, Small bowel, Brain (Glioblastoma), Skin (Multiple sebaceous tumours).
- \*\*\* Wimmer score –Scoring system for Congenital Mismatch Repair Deficiency. Further information can be requested from a Regional Genetic Clinic.
  - Clinical Criteria for germline testing in a living individual affected by colorectal polyps:
    - ≥5 adenomatous polyps and colorectal cancer (<60 years) OR</li>
    - → ≥5 adenomatous polyps (age <40 years), OR</p>
    - ≥10 adenomatous polyps (age <60 years), OR</li>
    - ≥20 adenomatous polyps (age ≥ 60 years), OR
    - ≥5 adenomatous polyps (age <60 years) AND first degree relative with</li>
       ≥5 adenomatous polyps OR CRC (age <60 years), OR</li>
    - ≥10 adenomatous polyps (age ≥ 60 years) AND first degree relative with ≥5 adenomatous polyps OR CRC (age <60years).</li>
    - ≥5 serrated lesions/polyps proximal to the rectum, all being ≥5 mm in size, with ≥2 being ≥10 mm in size
    - >20 serrated lesions/polyps of any size distributed throughout the large bowel, with ≥5 being proximal to the rectum.
  - NOTE: Polyps should be histologically confirmed where possible.
     Testing may also be considered for unusual/large polyps occurring at a young age.
  - For Juvenile Polyps, see test criteria for Juvenile Polyposis Syndrome





- For Hamartomatous Polyps, see test criteria for Peutz Jegher Syndrome
- Clinical Criteria for germline testing in a living individual affected by an extra-colonic manifestation of Familial Adenomatous Polyposis where they are too young to have developed bowel polyps or colonoscopy surveillance has not yet been undertaken (APC only):
  - Aggressive fibromatosis/Desmoid tumour (CTNNB1 WT where testing performed) (and MUTYH tested if abdominal desmoid) OR
  - Multiple CHRPEs that are either (1) bilateral, (2) occur in multiple quadrants, (3) have pisiform shape OR (4) irregular borders AND NOT bear track (Please note that bear track lesions clumped in a single quadrant are not a risk factor for FAP) OR
  - Cribriform-morular variant of papillary thyroid cancer OR
  - Hepatoblastoma OR
  - Multiple osteomas of skull and mandible or multiple dental abnormalities (unerupted teeth, supernumerary teeth with dentigerous cysts or odontomas) in children/young adults
- Clinical criteria for mosaic FAP testing in a living individual on >1 polyps:
  - Negative germline testing for APC, AND
  - Fulfils clinical diagnosis of FAP or attenuated FAP, AND
  - Testing will impact on the management of the patient and/or their relatives

## Requesting specialties





#### HEREDITARY DIFFUSE GASTRIC CANCER SYNDROME

## Available testing

| Centre             | Method       | Scope a        | and ra | nge of test  | Targets      | TAT |
|--------------------|--------------|----------------|--------|--------------|--------------|-----|
| Edinburgh          | NGS          | Whole gene sci | reen   | SNVs, indels | CDH1, CTNNA1 | 56  |
| Aberdeen           |              | Whole gene sci | reen   | SNVs, indels | CDH1         |     |
| Family men         | nber testing |                |        |              |              |     |
| Proforma required? |              | NO             |        |              |              | 1   |

#### Referral criteria

Living affected individual (proband) where the individual +/- family history meets one of the criteria. The proband has:

- a. Diffuse gastric cancer (<50 years), OR
- Gastric in situ signet ring cells or pagetoid spread of signet ring cells under 50 years OR
- c. Diffuse gastric cancer at any age with a personal history or first degree relative with cleft lip or cleft palate OR
- d. Double primary diffuse gastric cancer and lobular breast cancer (both <70 years)</li>
- e.. Diffuse gastric cancer and ≥1 first / second degree relative has diffuse gastric cancer any age, OR
- f.. Diffuse gastric cancer at any age and ≥1 first / second degree relative has lobular breast cancer <70 years, OR
- g. Lobular breast cancer and ≥1 first / second degree relative has diffuse gastric cancer (≥1 case occurred at <70 years)
- h. 2 cases of lobular breast cancer <50 years e.g. bilateral or multiple ipsilateral tumours

Note: At least one cancer should be histologically confirmed

NOTE: Where testing is being performed for breast panel test criteria, CDH1
can be added and reported using the same panel (Glasgow and Aberdeen
laboratories). Please contact the relevant laboratory to discuss if required.

#### Requesting specialties

- Clinical Genetics
- Gastroenterology

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# HEREDITARY LEIOMYOMATOSIS AND RENAL CELL CANCER SYNDROME

## Available testing

| Centre       | Method                     | Scope and range of test |             |                                  | Targets | TAT |
|--------------|----------------------------|-------------------------|-------------|----------------------------------|---------|-----|
| Dundee       | NGS<br>(targeted<br>panel) | Whole                   | gene screen | SNVs, indels,<br>exon level CNVs | FH      | 56  |
| Family men   | nber testing               |                         | 14          |                                  |         |     |
| Proforma red | quired?                    | NO                      |             |                                  |         |     |

## Referral criteria

Testing of affected individual (proband) with hereditary leiomyomatosis and renal cell cancer (HLRCC) or other FH deficiency disorder where the individual +/- family history meets one of the following criteria. The proband has:

- a. Type 2 papillary, HLRCC associated RCC (WHO pathology definition) OR tubulo-papillary renal tumour at any age, OR
- b. Two of: cutaneous leiomyomata, renal tumour (any histology), OR uterine leiomyomata with classic histological features < 40 years OR
- c. Cutaneous leiomyomata AND one first / second / third degree relative with renal tumour, OR
- d. Cutaneous leiomyomata AND two first / second / third degree relatives with cutaneous leiomyomata OR uterine leiomyomata with classic histological features < 40 years, OR
- e. Uterine leiomyomata with classic histological features (age <40) OR
- f. Multiple cutaneous leiomyomata

# Requesting specialties





## HEREDITARY OVARIAN CANCER SYNDROME

## Available testing

| Centre                         | Method | Scope a           | nd range of test                  | Targets   | TAT |
|--------------------------------|--------|-------------------|-----------------------------------|---|-----|
| Aberdeen<br>Glasgow            | NGS    | Whole gene screen | SNVs, indels<br>Exon level<br>CNV | BRCA1, BRCA2, RAD51C, RAD51D, BRIP1,<br>MSH2, MSH6, MLH1, PALB2 | 56  |
| Family member testing          |        |                   | ·                                 | as indicated above  | 14  |
| Proforma required? YES Glasgow |        |                   | Glasgow laboratory                | only (see centre website)                                       |     |

## Referral criteria

High-grade non-mucinous ovarian cancer, any age

**N.B.** *BRCA1* and *BRCA2* testing in the tumour is also available, specifically for platinum sensitive high grade serous epithelial ovarian, fallopian tube or primary peritoneal cancer (FIGO stage III or stage IV). Please refer to the Scottish Molecular Pathology Laboratory Consortium Genomic Test Directory.

- Clinical Genetics
- Oncology





## **INHERITED PANCREATIC CANCER**

## Available testing

| Centre                          | Method      | Scop | e and ra      | ange of test                      | Targets   | TAT |
|---------------------------------|-------------|------|---------------|-----------------------------------|---|-----|
| Glasgow                         | NGS<br>MLPA |      | e gene<br>een | SNVs, indels<br>Exon level<br>CNV | BRCA2, CDK4, CDKN2A, MLH1,<br>MSH2, MSH6, PALB2, STK11,<br>TP53 | 56  |
| Family member as ir testing     |             |      |               | ndicated above                    |   |     |
| Proforma required? YES Heredita |             |      |               | itary cancer profor               | rma (see centre website)  |     |

#### Referral criteria

Testing of affected individual (proband) where the individual +/- family history meets one of the following criteria. The proband has:

- 1. Pancreatic cancer age <50, OR
- 2. Pancreatic cancer age <70, AND
- a. Breast cancer age <60, melanoma age <60, OR ovarian cancer, OR
- b. One first / second degree relative with pancreatic cancer age <60, OR
- c. Two first / second degree relatives with any of breast cancer age <60, melanoma age <60, OR ovarian cancer

NOTE: If there is a family history of BRCA-related cancers (breast, ovarian, prostate, pancreatic) or history of melanoma and the patient does not meet the above criteria, please consider if they meet testing criteria for the hereditary breast, ovarian or melanoma panels.

- Clinical Genetics
- Oncology in discussion with Clinical Genetics





#### HEREDITARY PROSTATE CANCER

## Available testing

| Centre                         | Met              | hod                | Scope and range of test |        |  | Targets   | TAT |  |
|--------------------------------|------------------|--------------------|-------------------------|--------|--|---|-----|--|
| Aberdeen                       | NO<br>HOX<br>San | (B13               | Whole<br>gene<br>scree  | Э      | SNVs, indels<br>Exon level<br>CNV        | BRCA1, BRCA2, CHEK2, ATM, TP53, MLH1, MSH2, MSH6, RAD51D, PMS2, EPCAM, PALB2*, HOXB13**. *PALB2 only included where there is a family history of breast cancer **Please note CNV analysis is not currently performed for this gene. | 56  |  |
| Family member testing          |                  | as indicated above |                         |        |  |   |     |  |
| Proforma YES Prostat required? |                  |                    |                         | Prosta | ate cancer proforma (see centre website) |   |     |  |

Note: This test is currently available as pilot programme.

#### Referral criteria

- A man with prostate cancer diagnosed below the age of 50 years
- A man with metastatic prostate cancer diagnosed below 60 years with one first degree relative (a brother or a father) diagnosed with prostate cancer below 60 years
- A man diagnosed with metastatic prostate cancer with two first degree relatives (or one first and one second degree relative who are all first degree relatives of each other) with prostate cancer (patient and two brothers/ patient + 1 brother and father/ patient, father and father's brother/ patient, father & father's father)
- A man with prostate cancer who has a family history of cancer with a Manchester score greater than or equal to 15

- Clinical Genetics
- Oncology in discussion with Clinical Genetics





#### **JUVENILE POLYPOSIS**

# Available testing

| Centre             | Method                | Scope and range of test |  |                                | Targets       | TAT |  |
|--------------------|-----------------------|-------------------------|--|--------------------------------|---------------|-----|--|
| Edinburgh          | NGS<br>MLPA           | Whole gene screen       |  | SNVs, indels<br>Exon level CNV | SMAD4, BMPR1A | 56  |  |
| Family me          | Family member testing |                         | as indicated above   |                                |               |     |  |
| Proforma required? |                       | YES                     | Colorectal cancer gene panel proforma (see centre website) |                                |               |     |  |

## Referral criteria

- Juvenile polyposis syndrome:
  - o a. ≥ 5 juvenile polyps of the colorectum, OR
  - $\circ$  b.  $\geq$  2 juvenile polyps throughout the GI tract, OR
  - c. ≥ 1 juvenile polyp and a first / second degree relative has juvenile polyp, OR criteria

- Clinical Genetics
- Oncology





#### LI-FRAUMENI SYNDROME

## Available testing

| Centre              | Method      |                    | Scope and   | I range of test                | Targets | TAT |
|---------------------|-------------|--------------------|-------------|--------------------------------|---------|-----|
| Aberdeen<br>Glasgow | NGS         | Whole              | gene screen | SNVs, indels<br>Exon level CNV | TP53    | 56  |
| Family mem          | ber testing | as indicated above |             |                                |         |     |
| Proforma requ       | ired?       | NO                 |             |                                |         |     |

#### Referral criteria

- Proband and / or family history meets one of the following criteria:
  - Any sarcoma (<18 years)</li>
  - o Rhabdomyosarcoma of embryonal anaplastic subtype (any age)
  - Adrenocortical cancer (any age)
  - Choroid plexus cancer (any age)
  - Breast cancer (≤40 years) eligible for full hereditary breast cancer panel Hypodiploid acute lymphoblastic leukaemia (<18 years)</li>
  - SHH medulloblastoma (<18 years)</li>
  - ≥2 LFS-related cancers\* (both occurring ≤46 years; 2 breast cancers not eligible)
  - ≥1 LFS-related cancer\* with ≥1 1<sup>st</sup> / 2<sup>nd</sup> degree relative with ≥1 LFS-related cancer\* (one case ≤46 years, the other ≤56 years; 2 breast cancers not eligible)
  - Cancer with ≥2 1<sup>st</sup> / 2<sup>nd</sup> degree relatives with cancer (sarcoma ≤45 years, any cancer ≤45 years and sarcoma or any cancer ≤45 years)
  - \* Sarcoma of bone/soft tissue, breast cancer, central nervous system tumours, adrenocortical cancer or any childhood cancer (occurring ≤ 18 years)

- Clinical Genetics
- Oncology





#### **PEUTZ-JEGHERs SYNDROME**

## Available testing

| Centre      | Method                                   |       | Scope and ra   | ange of test   | Targets | TAT |  |  |
|-------------|--|-------|----------------|--|---------|-----|--|--|
| Edinburgh   | NGS<br>MLPA                              | Whole | gene screen    | SNVs, indels<br>Exon level CNV                             | STK11   | 56  |  |  |
| Family mer  | Family member testing as indicated above |       |                |  |         | 14  |  |  |
| Proforma re | equired?                                 | YES   | Colorectal can | Colorectal cancer gene panel proforma (see centre website) |         |     |  |  |

#### Referral criteria

- Living affected individual (proband) where the individual +/- family history meets one of the criteria.
  - o 1. ≥2 PJS-type hamartomatous polyps, OR
  - 2. ≥1 PJS-type hamartomatous polyp and characteristic mucocutaneous pigmentation, OR
  - 3. Characteristic mucocutaneous pigmentation age
  - 4. Sex cord tumours with annular tubules (SCAT) at any age
  - o 5. Adenoma malignum of the cervix at any age
  - 6. ≥1 PJS-type hamartomatous polyp, AND ≥1 first / second degree relative with: a. ≥1 PJS-like feature, OR b. ≥2 PJS-related cancers (the two cancers can be in the same or different relatives), OR
  - 7. Characteristic mucocutaneous pigmentation (<10), AND ≥1 first / second degree relative with: a≥1 PJS-like feature, OR b. ≥2 PJSrelated cancers (the two cancers can be in the same or different relatives)
- Deceased affected individual (proband) where (i) the individual +/- family history meets one of the above criteria, (ii) appropriate tissue is available (tumour or normal), and (iii) no living affected individual is available for genetic testing PJS-like features: characteristic mucocutaneous pigmentation, PJS-type hamartomatous polyps PJS-related cancers: epithelial colorectal, gastric, pancreatic, breast, and ovarian cancers, sex cord tumors with annular tubules (SCTAT), adenoma malignum of the cervix, and Sertoli cell tumors (LCST) of the testes
- The majority of polyps should be histologically confirmed

## Requesting specialties





#### **RENAL CANCER**

#### Available testing

| Centre     | Method         | Scope and range of test |   | Targets                                 | TAT |
|------------|----------------|-------------------------|---|---|-----|
| Dundee     | NGS            | Whole<br>gene<br>screen | SNVs, indels,<br>Exon level CNV for<br>selected genes | BAP1, FH, FLCN, MET, PTEN, SDHB,<br>VHL | 56  |
| _          | member<br>ting |                         | as indicated above                                    |   | 14  |
| Proforma r | equired?       | NO                      |   |   |     |

#### Referral criteria

- Individuals with:
  - o Renal cancer (≤ 40 years), OR
  - o Type 2 papillary renal cancer (≤50 years), OR
  - o Bilateral/multifocal or unusual pathology renal cancer (any age), OR
  - Renal cancer AND first / second degree relative with renal cancer, both cases diagnosed under 50
  - Single gene testing can be requested where specific features are present.

Renal cancer and features of an inherited renal cancer syndrome such as:

- Cerebellar/spinal haemangioblastoma
- Retinal angioma
- Phaeochromocytoma/paraganglioma
- Spontaneous pneumothorax
- Fibrofolliculomas
- Trichodiscomas
- Cutaneous Leiomyomata
- Uterine leiomyomas (under 40 years of age with pathology suggesting FH mutation)
- Mesothelioma
- Uveal melanoma

# Requesting specialties

Clinical Genetics





## **RHABDOID TUMOUR**

# Available testing

| Centre                | Method         |       | Scope and range of test |                                | Targets          | TAT |
|-----------------------|----------------|-------|-------------------------|--------------------------------|------------------|-----|
| Glasgow               | Sanger<br>MLPA | Whole | gene screen             | SNVs, indels<br>Exon level CNV | SMARCA4, SMARCB1 | 56  |
| Family men            | nber testing   |       |                         | as indicated abo               | ove              | 14  |
| Proforma required? NO |                |       |                         |                                |                  |     |

## Referral criteria

- Child with atypical teratoid / rhabdoid tumouor (ATRT) or malignant rhabdoid tumour (MRT) showing loss of SMARCB1 on immunohistochemistry OR
- Small cell carcinoma of the ovary, hypercalcaemic type (SCCOHT) (any age)

# Requesting specialties

Clinical Genetics





# **METABOLIC**

#### AMINO ACID DISORDERS & DISORDERS OF NEUROTRANSMISSION

# Available testing

| Centre      | Method   | Scope and range of test |                                   | Targets   | TAT |
|-------------|----------|-------------------------|-----------------------------------|---|-----|
| Aberdeen    | NGS      | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | ABAT, ALDH18A1, ALDH5A1, ALDH7A1, AMT,<br>ASPA, CBS, CTH, D2HGDH, DBH, DDC, FAH,<br>GABRG2, GCDH, GCH1, GLDC, GLRA1, HGD,<br>L2HGDH, MAT1A, OAT, PAH, PCBD1, PNPO,<br>QDPR, SLC25A22, SLC6A19, SLC7A7, SUOX | 112 |
| Family r    |          |                         | as indicated above                |   | 14  |
| Proforma re | equired? | NO                      |                                   |   |     |

#### Referral criteria

- Clinical phenotype suggests an amino acid disorder or disorder of neurotransmission
- Biochemical testing supportive (abnormal urine or plasma amino acid profile, abnormal urine organic amino acid profile)
- Where biochemical testing indicates testing of a single gene, please indicate
  this on the referral form and testing can be performed by either NGS or
  Sanger sequencing. Single gene indications so far: ASPA (Sanger)

- Clinical Genetics
- Metabolic





## **BATTEN DISEASE**

# Available testing

| Centre                | Method         | Scope and range of test |                                | Targets | TAT |
|-----------------------|----------------|-------------------------|--------------------------------|---------|-----|
| Aberdeen              | Sanger<br>MLPA | Whole gene<br>screen    | SNVs, indels<br>Exon level CNV | TPP1    | 56  |
| Family me             | ember testing  |                         | as indicated above             |         |     |
| Proforma required? NO |                |                         |                                |         |     |

## Referral criteria

- Clinical features suggestive of Batten disease
- · Biochemical tests supportive of diagnosis

# Requesting specialties

- Clinical Genetics
- Metabolic

## **BIOTINIDASE DEFICIENCY**

# Available testing

| Centre                | Method       | Scope and range of test |              | Targets    | TAT |
|-----------------------|--------------|-------------------------|--------------|------------|-----|
| Dundee                | Sanger       | Whole gene screen       | SNVs, indels | BTD        | 56  |
| Family me             | mber testing |                         | as indic     | ated above | 14  |
| Proforma required? NO |              |                         |              |            |     |

## Referral criteria

 Individuals where newborn screening or biochemical findings indicate multiple carboxylase deficiency.

- Clinical Genetics
- Metabolic





# BIOTIN-THIAMINE-RESPONSIVE BASAL GANGLIA DISEASE (BTBGD)

# Available testing

| Centre      | Method   | Scope and range of test |              | Targets    | TAT |
|-------------|----------|-------------------------|--------------|------------|-----|
| Aberdeen    | Sanger   | Whole gene screen       | SNVs, indels | SLC19A3    | 56  |
| Family r    |          |                         | as indic     | ated above | 14  |
| Proforma re | equired? | NO                      |              |            |     |

## Referral criteria

- Clinical features suggestive of BTBGD
- Biochemical tests supportive of diagnosis

# Requesting specialties

- Clinical Genetics
- Metabolic

# **BROWN VIALETTO VAN LAERE SYNDROME (BVVLS)**

# Available testing

| Centre             | Method | Scope and range of test |              | Targets          | TAT |
|--------------------|--------|-------------------------|--------------|------------------|-----|
| Aberdeen           | Sanger | Whole gene<br>screen    | SNVs, indels | SLC52A2, SLC52A3 | 56  |
|                    |        | ated above              | 14           |                  |     |
| Proforma required? |        | NO                      |              |                  |     |

# Referral criteria

- Clinical features suggestive of BVVLS
- Biochemical tests supportive of diagnosis

- Clinical Genetics
- Metabolic





## CARNITINE PALMITOYLTRANSFERASE ii DEFICIENCY

# Available testing

| Centre      | Method                | Scope and range of test |              | Targets | TAT |
|-------------|-----------------------|-------------------------|--------------|---------|-----|
| Aberdeen    | Sanger                | Whole gene<br>screen    | SNVs, indels | CPT2    | 56  |
|             |                       |                         | ated above   | 14      |     |
| Proforma re | Proforma required? NO |                         |              |         |     |

## Referral criteria

- Clinical features suggestive of Carnitine Palmityltransferase II deficiency
- Biochemical tests supportive of diagnosis (Hypoketotic hypoglycaemia)

- Clinical Genetics
- Metabolic





## **CEREBRAL FOLATE TRANSPORT DEFICIENCY**

## Available testing

| Centre      | Method   | Scope and range of test |                 | Targets     | TAT |
|-------------|----------|-------------------------|-----------------|-------------|-----|
| Aberdeen    | Sanger   | Whole gene screen       | SNVs,<br>indels | FOLR1       | 56  |
| Family r    |          |                         | as indi         | cated above | 14  |
| Proforma re | equired? | NO                      |                 |             |     |

## Referral criteria

- Clinical features suggestive of Cerebral Folate Transport Deficiency
- Biochemical tests supportive of diagnosis (Vitamin B9 deficiency)

# Requesting specialties

- Clinical Genetics
- Metabolic

#### **CITRULLINAEMIA TYPE 1**

# Available testing

| Centre                   | Method   | Scope and range of test |              | Targets | TAT |
|--------------------------|----------|-------------------------|--------------|---------|-----|
| Aberdeen                 | Sanger   | Whole gene screen       | SNVs, indels | ASS1    | 56  |
| Family member as indicat |          | ted above               | 14           |         |     |
| Proforma re              | equired? | NO                      |              |         |     |

## Referral criteria

- Clinical features suggestive of Citrullinaemia Type 1
- Biochemical tests supportive of diagnosis (Abnormal plasma amino acid profile)

- Clinical Genetics
- Metabolic





## **COBALAMIN C DEFICIENCY**

# Available testing

| Centre      | Method                | Scope and range of test |              | Targets        | TAT |
|-------------|-----------------------|-------------------------|--------------|----------------|-----|
| Aberdeen    | Sanger                | Whole gene screen       | SNVs, indels | MMACHC         | 56  |
| Family r    |                       |                         | as ir        | ndicated above | 14  |
| Proforma re | Proforma required? NO |                         |              |                |     |

#### Referral criteria

- Clinical features suggestive of Cobalamin C Deficiency
- Biochemical tests supportive of diagnosis (Vitamin B12 deficiency)

# Requesting specialties

- Clinical Genetics
- Metabolic

## **CREATINE DEFICIENCY SYNDROME**

# Available testing

| Centre      | Method   | Scope and range of test |              | Targets            | TAT |
|-------------|----------|-------------------------|--------------|--------------------|-----|
| Aberdeen    | NGS      | Whole gene screen       | SNVs, indels | GATM, GAMT, SLC6A8 | 56  |
| Family r    |          |                         | as ind       | icated above       | 14  |
| Proforma re | equired? | NO                      |              |                    |     |

# Referral criteria

- Clinical features suggestive of Creatine Deficiency Syndrome
- Biochemical tests supportive of diagnosis

- Clinical Genetics
- Metabolic





# DISORDERS ASSOCIATED WITH HYPERAMMONAEMIA / FATTY ACID OXIDATION / KETOGENESIS / KETOLYSIS

# Available testing

| Centre                | Method | Scope and range of test |                                   | Targets   | TAT |
|-----------------------|--------|-------------------------|-----------------------------------|---|-----|
| Aberdeen              | NGS    | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | ACADM, ACADS, ACADVL, ARG1, ASL, ASS1, CPS1, CPT1A, CPT2, ETFA, ETFB, ETFDH, GLUD1, HADHA, HADHB, HMGCL, HMGCS2, IVD, LPIN1, MMAA, MMAB, MMACHC, MMADHC, MUT, NAGS, OAT, OTC, OXCT1, PCCA, PCCB, SLC16A1, SLC22A5, SLC25A13, SLC25A15, SLC25A20, SLC7A7, SLC52A2, SLC52A3 | 112 |
| Family member         |        |                         | as indicated above                | 14  |     |
| testing               |        |                         |                                   |   |     |
| Proforma required? NO |        |                         |                                   |   |     |

#### Referral criteria

- Clinical features suggestive of Disorders associated with Hyperammonaemia / Fatty Acid Oxidation / Ketogenesis / Ketolysis (e.g. encephalopathy, severe vomiting or loss of consciousness)
- Biochemical tests supportive of diagnosis (Plasma ammonia >150umol/L or Hypoketotic hypoglycaemia or severe ketoacidosis)
- Where biochemical testing indicates testing of a single gene, please indicate
  this on the referral form and testing can be performed by either NGS or
  Sanger sequencing. Single gene indications so far: HMGCL (Sanger),
  HADHA (Sanger), ARG1 (Sanger)

- Clinical Genetics
- Metabolic





# DISORDERS OF CARBOHYDRATE METABOLISM (incl. GLYCOGEN STORAGE DISORDERS)

## Available testing

| Centre                | Method | Scope and range of test |                                   | Targets  | TAT |
|-----------------------|--------|-------------------------|-----------------------------------|--|-----|
| Aberdeen              | NGS    | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | AGL, ALDOA, ALDOB, ENO3, EPM2A,<br>FBP1, G6PC, G6PC3, GAA, GALE, GALK1,<br>GALT, GBE1, GYG1, GYS1, GYS2, LAMP2,<br>LDHA, NHLRC1, PFKM, PGAM2, PGK1,<br>PGM1, PHKA1, PHKA2, PHKB, PHKG2,<br>PRKAG2, PYGL, PYGM, SLC2A2, SLC16A1,<br>SLC37A4 | 112 |
| Family member testing |        |                         | ;                                 | as indicated above   | 14  |
| Proforma required?    |        | NO                      |                                   |  |     |

#### Referral criteria

- Clinical features suggestive of a disorder of carbohydrate metabolism
- Biochemical or haematological tests supportive of diagnosis (e.g. Abnormal liver function, abnormal muscle physiology, hypoglycaemia, hypobilirubinaemia, presence of urinary reducing substances, reduced GALT, GALE activity in blood, abnormal CSF:blood glucome ratio)
- Where biochemical testing indicates testing of a single gene, please indicate
  this on the referral form and testing can be performed by either NGS or
  Sanger sequencing. Single gene indications so far: ALDOB (Sanger)

- Clinical Genetics
- Metabolic





## **FABRY DISEASE**

# Available testing

| Centre                | Method | Scope and range of test |              | Targets    | TAT |
|-----------------------|--------|-------------------------|--------------|------------|-----|
| Edinburgh             | Sanger | Whole gene screen       | SNVs, indels | GLA        | 56  |
| Family member testing |        |                         | as indic     | ated above | 14  |
| Proforma required?    |        | NO                      |              |            |     |

## Referral criteria

- In males: clinical and laboratory features characteristic of Fabry disease following alpha-galactosidase A enzyme testing
- In females: clinical features characteristic of Fabry disease

- Clinical Genetics
- Metabolic





## FAMILIAL HYPERCHOLESTEROLAEMIA

## Available testing

| Centre                | Method | Sc    | Scope and range of test   |   | Targets                         | TAT |  |
|-----------------------|--------|-------|---|---|---------------------------------|-----|--|
| Aberdeen              | NGS    | Whole | e gene<br>een   | SNVs, indels<br>Exon level CNV*<br>(*LDLR only) | LDLR,APOE, PCSK9, APOB, LDLRAP1 | 56  |  |
| Family member testing |        |       | as indicated above  |   |                                 |     |  |
| Proforma required? NO |        |       | Optional FH proforma on centre website. GPs should complete this form of discuss with lipid consultant prior to referral. |   |                                 | or  |  |

## Referral criteria

 Total cholesterol >7.5 mmol/l (>6.7mmol/l in a child < 16 years) or LDL cholesterol >4.9 mmol/l (>4 mmol/l in a child < 16 yrs)</li>

AND one or more of the following:

- Tendon xanthomas in the index individual or Tendon xanthomas in a 1st or 2nd degree relative
- Family history of myocardial infarction: in 2nd degree relative <50 yrs or in 1st degree relative < 60 yrs</li>
- Family history of raised total cholesterol: >7.5mmol/l in an adult 1st or 2nd degree relative or >6.7 mmol/l in a child or sibling < 16 yrs</li>

Secondary causes of hypercholesterolaemia should be excluded (diabetes, thyroid disease, abnormal LFTs). If in doubt, please seek advice from your local lipid clinic.

- Cardiologists
- Clinical Genetics
- GPs must complete proforma or discuss with Lipid consultant prior to referral
- Lipidology
- Metabolic





## **FANCONI-BICKEL SYNDROME**

# Available testing

| Centre                | Method | Scope and range of test |              | Targets     | TAT |
|-----------------------|--------|-------------------------|--------------|-------------|-----|
| Aberdeen              | Sanger | Whole gene screen       | SNVs, indels | SLC2A2      | 56  |
| Family member testing |        |                         | as indi      | cated above | 14  |
| Proforma required?    |        | NO                      |              |             |     |

# Referral criteria

- Clinical features suggestive of Fanconi Bickel Syndrome
- Biochemical tests supportive of diagnosis

- Clinical Genetics
- Metabolic





## **FATTY ACID OXIDATION**

# Available testing

| Centre                | Method | Scope and range of test |                                   | Targets  | TAT |
|-----------------------|--------|-------------------------|-----------------------------------|--|-----|
| Aberdeen              | NGS    | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | ACADM, ACADS, ACADVL, CPT1A, CPT2,<br>ETFA, ETFB, ETFDH, HADHA, HADHB, HMGCL,<br>HMGCS2, IVD, MMAA, MMAB, MMACHC,<br>MMADHC, OXCT1, SLC22A5, SLC25A20,<br>SLC52A2, SLC52A3 | 112 |
| Family member testing |        |                         | as indicated above                | 14   |     |
| Proforma required? NO |        |                         |                                   |  |     |

## Referral criteria

- Clinical features suggestive of a Fatty Acid Oxidation disorder
- Biochemical tests supportive of diagnosis
- Where biochemical testing indicates testing of a single gene, please indicate
  this on the referral form and testing can be performed by either NGS or
  Sanger sequencing.

- Clinical Genetics
- Metabolic
- Neurology





## **GALACTOSAEMIA**

# Available testing

| Centre      | Method         | Scope and         | range of test                  | Targets | TAT |
|-------------|----------------|-------------------|--------------------------------|---------|-----|
| Aberdeen    | Sanger<br>MLPA | Whole gene screen | SNVs, indels<br>Exon level CNV | GALT    | 56  |
| Family r    |                |                   | as indicated a                 | bove    | 14  |
| Proforma re | <u> </u>       | NO                |                                |         |     |

#### Referral criteria

- Clinical features suggestive of Galactosaemia
- Biochemical tests supportive of diagnosis (Increase galactose in blood)

# Requesting specialties

- Clinical Genetics
- Metabolic

# GAUCHER DISEASE (B-GLUCOCEREBROSIDASE DEFICIENCY)

# Available testing

| Centre                | Method | Scope and range of test |              | Targets    | TAT |
|-----------------------|--------|-------------------------|--------------|------------|-----|
| Aberdeen              | Sanger | Whole gene screen       | SNVs, indels | GBA        | 56  |
| Family member testing |        |                         | as indica    | ated above | 14  |
| Proforma required?    |        | NO                      |              |            |     |

## Referral criteria

- Clinical features suggestive of Gaucher disease
- Biochemical tests supportive of diagnosis (Decreased glucocerebrosidase enzyme levels)

# Requesting specialties

- Clinical Genetics
- Metabolic

# **GLUTARIC ACIDAEMIA TYPE 1**

# Available testing

| Centre | Method | Scope and range of test | Targets | TAT |
|--------|--------|-------------------------|---------|-----|
|        |        |                         |         |     |





| Aberdeen           | Sanger | Whole gene | SNVs, indels       | GCDH | 56 |  |  |
|--------------------|--------|------------|--------------------|------|----|--|--|
|                    |        | screen     |                    |      |    |  |  |
| ,                  | member |            | as indicated above |      |    |  |  |
| testing            |        |            |                    |      |    |  |  |
| Proforma required? |        | NO         |                    |      |    |  |  |

#### Referral criteria

- Clinical features suggestive of Glutaric Acidaemia Type 1
- Biochemical / newborn screen test supportive of diagnosis

# Requesting specialties

- Clinical Genetics
- Metabolic

## **GLYCEROL KINASE DEFICIENCY**

# Available testing

| Centre                | Method | Scope and range of test |              | Targets       | TAT |
|-----------------------|--------|-------------------------|--------------|---------------|-----|
| Aberdeen              | Sanger | Whole gene screen       | SNVs, indels | GK            | 56  |
| Family member testing |        |                         | as in        | dicated above | 14  |
| Proforma required?    |        | NO                      |              |               |     |

## Referral criteria

- Clinical features suggestive of Glycerol Kinase Deficiency
- Biochemical tests supportive of diagnosis (Glycerol peak in urine sample)

- Clinical Genetics
- Metabolic





## **GLYCOGEN STORAGE DISEASE**

# Available testing

| Aberdeen           | NGS | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | AGL, ALDOA, ALDOB, ENO3, EPM2A,<br>FBP1, G6PC, GAA, GBE1, GYG1, GYS1,<br>GYS2, LAMP2, LDHA, NHLRC1, PFKM,<br>PGAM2, PGK1, PGM1, PHKA1, PHKA2,<br>PHKB, PHKG2, PRKAG2, PYGL, PYGM,<br>SLC2A2, SLC37A4 | 112 |
|--------------------|-----|-------------------------|-----------------------------------|--|-----|
| Proforma required? |     | NO                      |                                   |  |     |

## Referral criteria

- Clinical features suggestive of a Glycogen storage disorder
- Biochemical or haematological tests supportive of diagnosis
- Where biochemical testing indicates testing of a single gene, please indicate
  this on the referral form testing can be performed by either NGS or Sanger
  sequencing.
- Metabolic

#### **GLYCOGEN STORAGE DISEASE 1A**

# Available testing

| Centre                | Method   | Scope and range of test |              | Targets   | TAT |
|-----------------------|----------|-------------------------|--------------|-----------|-----|
| Aberdeen              | Sanger   | Whole gene screen       | SNVs, indels | G6PC      | 56  |
| Family member testing |          |                         | as indica    | ted above | 14  |
| Proforma re           | equired? | NO                      |              |           |     |

## Referral criteria

- Clinical features suggestive of Glycogen Storage Disease 1A
- Biochemical tests supportive of diagnosis

- Clinical Genetics
- Metabolic





## **HOMOCYSTINURIA**

# Available testing

| Centre             | Method                             | Scope and range of test |                                | Targets                         | TAT |
|--------------------|------------------------------------|-------------------------|--------------------------------|---------------------------------|-----|
| Aberdeen           | NGS                                | Whole gene screen       | SNVs, indels<br>Exon level CNV | CBS, MMADHC, MTHR, MTR,<br>MTRR | 56  |
|                    | Family member as indicates testing |                         |                                | ted above                       | 14  |
| Proforma required? |                                    | NO                      |                                |                                 |     |

## Referral criteria

- Clinical features suggestive of Homocysteinuria
- Biochemical tests supportive of diagnosis (High homocysteine levels in blood)

- Clinical Genetics
- Metabolic





## HYPERLIPIDAEMIA, TYPE III

## Available testing

| Centre      | Method   | Scope and range of test |      | Targets                       | TAT |
|-------------|----------|-------------------------|------|-------------------------------|-----|
| Aberdeen    | Sanger   | Targeted screen         | SNVs | APOE (Codons p.130 and p.176) | 28  |
| Proforma re | equired? | NO                      |      |                               |     |

## Referral criteria

- Clinical features suggestive of Hyperlipidaemia Type III, e.g. accelerated atherosclerosis
- Biochemical tests supportive of diagnosis (Elevated cholesterol and triglycerides)

# Requesting specialties

- Clinical Genetics
- Lipidology

# HYPERTRIGLYCERIDAEMIA / FAMILIAL CHYLOMICRONAEMIA SYNDROME / LIPOPROTEIN LIPASE DEFICIENCY

# Available testing

| Centre      | Method                 | Scope and range of test |  | Targets                           | TAT |
|-------------|------------------------|-------------------------|--|-----------------------------------|-----|
| Aberdeen    | Sanger<br>MLPA         | Whole gene<br>screen    | SNVs, indels<br>Exon level CNV*<br>(*LPL only) | LPL, LMF1, APOC2, APOA5, GPI-HBP1 | 56  |
| ,           | Family member as indic |                         | ated above                                     | 14                                |     |
| Proforma re | Proforma required? NO  |                         |  |                                   |     |

#### Referral criteria

- Clinical features suggestive of hypertriglyceridaemia, e.g. recurrent pancreatitis, eruptive xanthomas, lipaemia retinalis.
- Secondary causes excluded.
- Biochemical tests supportive of diagnosis (Elevated triglycerides >20mmol/L)

# Requesting specialties

- Clinical Genetics
- Gastrohepatology
- Lipidology





## **HYPOBETALIPOPROTEINAEMIA**

# Available testing

| Centre      | Method                      | Scope and range of test |                                   | Targets                              | TAT |
|-------------|-----------------------------|-------------------------|-----------------------------------|--------------------------------------|-----|
| Aberdeen    | NGS                         | Whole gene<br>screen    | SNVs, indels<br>Exon level<br>CNV | ANGPTL3, APOB, MTTP, PCSK9,<br>SAR1B | 56  |
| _           | amily member as inditesting |                         | cated above                       | 14                                   |     |
| Proforma re | equired?                    | NO                      |                                   |                                      |     |

#### Referral criteria

- Clinical features suggestive of Hypobetalipoproteinaemia
- Biochemical tests supportive of diagnosis (Undetectable / low levels of ApoB)

- Clinical Genetics
- Lipidology





## LYSOSOMAL STORAGE DISORDERS

# Available testing

| Centre                | Method   | Scope and range of test |                                   | Targets   | TAT |
|-----------------------|----------|-------------------------|-----------------------------------|---|-----|
| Aberdeen              | NGS      | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | AGA, ARSA, ARSB, ARSK, ASAH1, CLN3, CLN5, CLN6, CLN8, CTNS, CTSA, CTSD, CTSK, DNAJC5, FUCA1, GAA, GALC, GALNS, GBA, GLA, GLB1, GM2A, GNE, GNPTAB, GNPTG, GNS, GUSB, HEXA, HEXB, HGSNAT, HYAL1, IDS, IDUA, LAMP2, LIPA, MAN2B1, MANBA, MCOLN1, MFSD8, NAGA, NAGLU, NEU1, NPC1, NPC2, PPT1, PSAP, SGSH, SLC17A5, SMPD1, SUMF1, TPP1 | 112 |
| Family member testing |          |                         | as indicated above                | 14  |     |
|                       |          | NO                      |                                   |   |     |
| Proforma re           | equired? | NO                      |                                   |   |     |

## Referral criteria

- Clinical features suggestive of a Lysosomal storage disorder
- Biochemical tests supportive of diagnosis (Abnormal urine MPS, oligosaccharide screen, white cell enzyme analysis)
- Where biochemical testing indicates testing of a single gene, please indicate
  this on the referral form and testing can be performed by either NGS or
  Sanger sequencing. Single gene indications so far: IDS (Sanger), SGSH
  (Sanger), SUMF1 (Sanger)

- Clinical Genetics
- Metabolic





# MAPLE SYRUP URINE DISEASE (MSUD)

# Available testing

| Centre                             | Method | Scope and range of test |                                | Targets             | TAT |
|------------------------------------|--------|-------------------------|--------------------------------|---------------------|-----|
| Aberdeen                           | NGS    | Whole gene screen       | SNVs, indels<br>Exon level CNV | BCKDHA, BCKDHB, DBT | 56  |
| Family member as indicated testing |        | d above                 | 14                             |                     |     |
| Proforma required? NO              |        |                         |                                |                     |     |

## Referral criteria

- Clinical features suggestive of Maple Syrup Urine Disease
- Biochemical tests supportive of diagnosis

# Requesting specialties

- Clinical Genetics
- Metabolic

# MEDIUM CHAIN ACYL-COA DEHYDROGENASE DEFICIENCY (MCADD)

# Available testing

| Centre             | Method         | Scope and range of test |              | Targets | TAT |
|--------------------|----------------|-------------------------|--------------|---------|-----|
| Glasgow            | Sanger         | Whole gene screen       | SNVs, indels | ACADM   | 56  |
|                    | member<br>ting |                         | as indicated | above   | 14  |
| Proforma required? |                | NO                      |              |         |     |

## Referral criteria

- Clinical features suggestive of MCADD
- · Biochemical tests supportive of diagnosis

- Clinical Genetics
- Metabolic
- Paediatrics





## **METACHROMATIC LEUKODYSTROPHY**

# Available testing

| Centre               | Method | Scope and range of test |              | Targets | TAT |
|----------------------|--------|-------------------------|--------------|---------|-----|
| Aberdeen             | Sanger | Whole gene screen       | SNVs, indels | ARSA    | 56  |
| Family member as inc |        | icated above            | 14           |         |     |
| Proforma required?   |        | NO                      |              |         |     |

# Referral criteria

- Clinical features suggestive of Metachromatic Leukodystrophy
- · Biochemical tests supportive of diagnosis

- Clinical Genetics
- Metabolic





**MUCOPOLYSACCHARIDOSIS (mps) PANEL** 

| Aberdeen              | NGS | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | ARSB, ARSK, GALNS, GLB1, GNS, GUSB,<br>HGSNAT, HYAL1, IDS, IDUA, NAGLU, SGSH | 112 |
|-----------------------|-----|-------------------------|-----------------------------------|--|-----|
| Proforma required? NO |     |                         |                                   |  |     |

## Referral criteria

- Clinical features suggestive of a Mucopolysaccharidosis disorder
- Biochemical tests supportive of diagnosis (Abnormal urine MPS)
- Where biochemical testing indicates testing of a single gene, please indicate
  this on the referral form and testing can be performed by either NGS or
  Sanger sequencing.
- Metabolic





# **MUCOPOLYSACCHARIDOSIS TYPE 1 (HURLER / SCHEIE SYNDROME)**

# Available testing

| Centre      | Method                | Scope and range of test |              | Targets    | TAT |
|-------------|-----------------------|-------------------------|--------------|------------|-----|
| Aberdeen    | Sanger                | Whole gene screen       | SNVs, indels | IDUA       | 56  |
|             | Family member testing |                         | as indica    | ated above | 14  |
| Proforma re | equired?              | NO                      |              |            |     |

#### Referral criteria

- Clinical features suggestive of Mucopolysaccharidosis Type 1
- · Biochemical tests supportive of diagnosis

# Requesting specialties

- Clinical Genetics
- Metabolic

## **MUCOLIPIDOSIS II & III ALPHA / BETA**

# Available testing

| Centre                | Method   | Scope and range of test |                                   | Targets | TAT |
|-----------------------|----------|-------------------------|-----------------------------------|---------|-----|
| Aberdeen              | NGS      | Whole gene<br>screen    | SNVs, indels<br>Exon level<br>CNV | GNPTAB  | 56  |
| Family member as indi |          | cated above             | 14                                |         |     |
| Proforma re           | equired? | NO                      |                                   |         |     |

# Referral criteria

- Clinical features suggestive of Mucolipidosis II & III Alpha / Beta.
- Biochemical tests supportive of diagnosis

- Clinical Genetics
- Metabolic





# MULTIPLE ACYL-Coa DEHYDROGENASE DEFICIENCY (MADD)

## Available testing

| Centre      | Method                                   | Scope and range of test |                                | Targets                                | TAT |
|-------------|--|-------------------------|--------------------------------|--|-----|
| Aberdeen    | NGS                                      | Whole gene screen       | SNVs, indels<br>Exon level CNV | ETFDH, ETFA, ETFB, SLC52A2,<br>SLC52A3 | 56  |
|             | Family member as indicated above testing |                         |                                |  | 14  |
| Proforma re | equired?                                 | NO                      |                                |  |     |

#### Referral criteria

- Clinical features suggestive of MADD
- Biochemical tests supportive of diagnosis

# Requesting specialties

- Clinical Genetics
- Metabolic
- Neurology

# **NEURONAL CEROID LIPOFUSCINOSIS (NCL)**

# Available testing

| Centre                | Method                                   | Scope and range of test |                                   | Targets   | TAT |  |  |
|-----------------------|--|-------------------------|-----------------------------------|---|-----|--|--|
| Aberdeen              | NGS                                      | Whole gene<br>screen    | SNVs, indels<br>Exon level<br>CNV | ATP13A3, CLN3, CLN5, CLN6, CLN8,<br>CTSD, DNAJC5, KCTD7, MFSD8, PPT1,<br>TPP1 | 56  |  |  |
| ,                     | Family member as indicated above testing |                         |                                   |   |     |  |  |
| Available g           | Available genes: See website             |                         |                                   |   |     |  |  |
| Proforma required? NO |  |                         |                                   |   |     |  |  |

#### Referral criteria

- Clinical features suggestive of Neuronal Ceroid Lipofuscinosis
- Haematological / Biochemical tests supportive of diagnosis (Demonstration of vacuolated lymphocytes, presence of pathological inclusions on tissue biopsies, deficient enzyme activity)
- Where biochemical testing indicates testing of a single gene, please indicate
  this on the referral form and testing can be performed by either NGS or
  Sanger sequencing.

# Requesting specialties

- Clinical Genetics
- Metabolic
- Neurology

NSD611-003.20 V5





## **NIEMANN-PICK DISEASE**

# Available testing

| Centre                | Method   | Scope and range of test |                                   | Targets           | TAT |
|-----------------------|----------|-------------------------|-----------------------------------|-------------------|-----|
| Aberdeen              | NGS      | Whole gene<br>screen    | SNVs, indels<br>Exon level<br>CNV | NPC1, NPC2, SMPD1 | 56  |
| Family member testing |          |                         | as indi                           | cated above       | 14  |
| Proforma re           | equired? | NO                      |                                   |                   |     |

#### Referral criteria

- Clinical features suggestive of Niemann Pick Disease
- · Biochemical tests supportive of diagnosis

# Requesting specialties

- Clinical Genetics
- Metabolic

#### **NIEMANN-PICK DISEASE TYPES A & B**

# Available testing

| Centre                   | Method | Scope and range of test |              | Targets | TAT |
|--------------------------|--------|-------------------------|--------------|---------|-----|
| Aberdeen                 | Sanger | Whole gene screen       | SNVs, indels | SMPD1   | 56  |
| Family member as testing |        | as ind                  | icated above | 14      |     |
| Proforma required?       |        | NO                      |              |         |     |

## Referral criteria

- Clinical features suggestive of Niemann Pick Disease Types A & B
- Biochemical tests supportive of diagnosis

- Clinical Genetics
- Metabolic





## **NIEMANN-PICK DISEASE TYPES C1 & C2**

# Available testing

| Centre                          | Method | Scope and range of test                       |          | Targets    | TAT |
|---------------------------------|--------|---|----------|------------|-----|
| Aberdeen                        | NGS    | Whole gene SNVs, indels screen Exon level CNV |          | NPC1, NPC2 | 56  |
| Family member as indica testing |        |   | ed above | 14         |     |
| Proforma required? NO           |        |   |          |            |     |

## Referral criteria

- Clinical features suggestive of Niemann Pick Disease Type C
- · Biochemical tests supportive of diagnosis

# Requesting specialties

- Clinical Genetics
- Metabolic

## NON KETOTIC HYPERGLYCINAEMIA

# Available testing

| Centre                 | Method   | Scope and range of test                       |    | Targets                        | TAT |
|------------------------|----------|---|----|--------------------------------|-----|
| Aberdeen               | NGS      | Whole gene SNVs, indels screen Exon level CNV |    | ALDH7A1, AMT, GLDC, PPT1, TPP1 | 56  |
| Family member as indic |          | ated above                                    | 14 |                                |     |
| Proforma re            | equired? | NO  |    |                                |     |

#### Referral criteria

- Clinical features suggestive of Non ketotic hyperglycinaemia
- Biochemical tests supportive of diagnosis

- Clinical Genetics
- Metabolic





## **ORGANIC ACIDAEMIAS & COFACTOR / VITAMIN DISORDERS**

# Available testing

| Centre        | Method   | Scope and range of test |                                   | Targets  | TAT |
|---------------|----------|-------------------------|-----------------------------------|--|-----|
| Aberdeen      | NGS      | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | ABCD4, ACSF3, AMN, AUH, BCKDHA, BCKDHB, BTD, CUBN, DBT, DHFR, DNAJC19, FOLR1, GIF, HCFC1, HLCS, IVD, LMBRD1, LPIN1, MCCC1, MCCC2, MCEE, MLYCD, MMAA, MMAB, MMACHC, MMADHC, MTHFD1, MTHFR, MTR, MTRR, MUT, OPA3, PC, PCCA, PCCB, PDHA1, PDHB, PDHX, PRDX1, SLC19A3, SLC46A1, SLC52A3, SUCLA2, SUCLG1, TAZ, TCN2, TMEM70 | 112 |
| Family member |          |                         | as indicated above                | 14   |     |
| tes           | ting     |                         |                                   |  |     |
| Proforma re   | equired? | NO                      |                                   |  |     |

## Referral criteria

- Clinical features suggestive of an organic acidaemia or cofactor / vitamin disorder
- Biochemical tests supportive of diagnosis (abnormal results of urine organic acid or amino acid screen, anaemia, unexplained deficiency of a specific vitamin)
- Where biochemical testing indicates testing of a single gene, please indicate
  this on the referral form and testing can be performed by either NGS or
  Sanger sequencing.

- Clinical Genetics
- Metabolic





## **ORGANIC ACIDURIA**

# Available testing

| Centre   | Method | Scope and range of test |              | Targets | TAT |
|----------|--------|-------------------------|--------------|---------|-----|
| Aberdeen | Sanger | Whole gene              | SNVs, indels | UMPS    | 56  |
|          |        | screen                  |              |         |     |

# Referral criteria

- Clinical features suggestive of Organic Aciduria
- Biochemical tests supportive of diagnosis

# Requesting specialties

- Clinical Genetics
- Metabolic

## ORNITHINE AMINOTRANSFERASE DEFICIENCY

# Available testing

| Centre                | Method | Scope and range of test |              | Targets      | TAT |
|-----------------------|--------|-------------------------|--------------|--------------|-----|
| Aberdeen              | Sanger | Whole gene screen       | SNVs, indels | OAT          | 56  |
| Family member testing |        |                         | as ind       | icated above | 14  |
| Proforma required? NO |        | NO                      |              |              |     |

# Referral criteria

- Clinical features suggestive of Ornithine Aminotransferase Deficiency
- · Biochemical tests supportive of diagnosis

- Clinical Genetics
- Metabolic





## ORNITHINE TRANSCARBAMULASE DEFICIENCY

# Available testing

| Centre      | Method                | Scope and range of test |                                | Targets    | TAT |
|-------------|-----------------------|-------------------------|--------------------------------|------------|-----|
| Edinburgh   | NGS<br>MLPA           | Whole gene screen       | SNVs, indels<br>Exon level CNV | ОТС        | 56  |
| Family mem  | nber testing          |                         | as indic                       | ated above | 14  |
| Proforma re | Proforma required? NO |                         |                                |            |     |

# Referral criteria

• Clinical features that indicate a likely diagnosis of Ornithine Transcarbamulase Deficiency

- Clinical Genetics
- Metabolic





## **PEROXISOMAL DISORDERS**

# Available testing

| Centre                | Method | Scope and range of test |                                   | Targets   | TAT |
|-----------------------|--------|-------------------------|-----------------------------------|---|-----|
| Aberdeen              | NGS    | Whole<br>gene<br>screen | SNVs, indels<br>Exon level<br>CNV | ABCD1, ACBD5, ACOX1, ACK, AGPS, AGXT,<br>AMACR, ARSE, CAT, DNM1L, DYM, EBP, FAR1,<br>GNPAT, GRHPR, HOGA1, HSD17B4, NDHL,<br>PEX1, PEX2, PEX3, PEX5, PEX6, PEX7, PEX10,<br>PEX11B, PEX12, PEX13, PEX14, PEX16, PEX19,<br>PEX26, PHYH, SCP2, TRIM37 | 112 |
| Family member testing |        |                         | as indicated above                | 14  |     |
| Proforma required? NO |        |                         |                                   |   |     |

## Referral criteria

- Clinical features suggestive of a Peroxisomal disorder
  - At least 2 of the following: Hypoptonia / developmental delay, Characteristic facial dysmorphism, Characteristic X-ray findings (e.g. stippling), Retinal dystrophy / sensorineural hearing loss, Liver dysfunction
- Biochemical tests supportive of diagnosis (Increased plasma very long chain fatty acids +/- erythrocyte membrane plasmalogens)
- Where biochemical testing indicates testing of a single gene, please indicate
  this on the referral form and testing can be performed by either NGS or
  Sanger sequencing. Single gene indications so far: ABCD1 (Sanger)

- Clinical Genetics
- Metabolic





## **PHENYLKETONURIA**

# Available testing

| Centre                | Method                | Scope and range of test |                                   | Targets      | TAT |
|-----------------------|-----------------------|-------------------------|-----------------------------------|--------------|-----|
| Glasgow               | Sanger<br>MLPA        | Whole gene<br>screen    | SNVs, indels<br>Exon level<br>CNV | РАН          | 56  |
| Family member testing |                       |                         | as ind                            | icated above | 14  |
| Proforma r            | Proforma required? NO |                         |                                   |              |     |

#### Referral criteria

- Elevated blood phenylalanine and low levels or absence of phenylalanine hydroxylase enzyme.
- Diagnosis of Phenylketonuria by Newborn screening.

# Requesting specialties

- Clinical Genetics
- Metabolic

# POMPE DISEASE / GLYCOGEN STORAGE DISEASE TYPE 2

# Available testing

| Centre                | Method | Scope and range of test |                                   | Targets | TAT |
|-----------------------|--------|-------------------------|-----------------------------------|---------|-----|
| Aberdeen              | NGS    | Whole gene<br>screen    | SNVs, indels<br>Exon level<br>CNV | GAA     | 56  |
| Family member testing |        | as indicated above      |                                   | 14      |     |
| Proforma required?    |        | NO                      |                                   |         |     |

#### Referral criteria

- Clinical features suggestive of Pompe disease
- · Biochemical tests supportive of diagnosis

- Clinical Genetics
- Metabolic





## **PROPRIONIC ANAEMIA**

# Available testing

| Centre                | Method | Scope and range of test |                                | Targets    | TAT |
|-----------------------|--------|-------------------------|--------------------------------|------------|-----|
| Aberdeen              | NGS    | Whole gene screen       | SNVs, indels<br>Exon level CNV | PCCA, PCCB | 56  |
| Family member testing |        |                         | as indica                      | ted above  | 14  |
| Proforma required?    |        | NO                      |                                |            |     |

## Referral criteria

- Clinical features suggestive of Proprionic Anaemia
- Biochemical tests supportive of diagnosis

# Requesting specialties

- Clinical Genetics
- Metabolic

#### **REFSUM DISEASE**

# Available testing

| Centre                | Method | Scope and range of test |                                | Targets    | TAT |
|-----------------------|--------|-------------------------|--------------------------------|------------|-----|
| Aberdeen              | NGS    | Whole gene screen       | SNVs, indels<br>Exon level CNV | PEX7, PHYH | 56  |
| Family member testing |        |                         | as indica                      | ited above | 14  |
| Proforma required?    |        | NO                      |                                |            |     |

## Referral criteria

- Clinical features suggestive of Refsum disease
- Biochemical tests supportive of diagnosis

- Clinical Genetics
- Metabolic





# SUCCINIC SEMIALDEHYDE DEHYRDOGENASE DEFICIENCY (SSADH)

# Available testing

| Centre             | Method | Scope and range of test |              | Targets | TAT |
|--------------------|--------|-------------------------|--------------|---------|-----|
| Aberdeen           | Sanger | Whole gene              | SNVs, indels | ALDH5A1 | 56  |
|                    |        | screen                  |              |         |     |
| Family r           | nember | as indicated above      |              |         | 14  |
| testing            |        |                         |              |         |     |
| Proforma required? |        | NO                      |              |         |     |

## Referral criteria

- Clinical features suggestive of Succinic Semialdehyde Dehydrogenase Deficiency (SSADH)
- Biochemical tests supportive of diagnosis

# Requesting specialties

- Clinical Genetics
- Metabolic

# **TANGO2-RELATED METABOLIC ENCEPHALOPATHY & ARRHYTHMIAS**

# Available testing

| Centre                | Method | Scope and range of test                                      |                 | Targets | TAT |
|-----------------------|--------|--|-----------------|---------|-----|
| Aberdeen              | Sanger | Whole gene screen SNVs, indels Long Range PCR Ex3-9 deletion |                 | TANGO2  | 56  |
| Family member testing |        |  | as indicated ab | ove     | 14  |
| Proforma required?    |        | NO   |                 |         |     |

## Referral criteria

Clinical features suggestive of TANGO2-related metabolic encephalopathy & arrhythmias

- Clinical Genetics
- Metabolic





## **TAY-SACHS DISEASE**

# Available testing

| Centre             | Method | Scope and range of test |                 | Targets       | TAT |
|--------------------|--------|-------------------------|-----------------|---------------|-----|
| Aberdeen           | Sanger | Whole gene screen       | SNVs,<br>indels | HEXA          | 56  |
| Family r           |        |                         | as inc          | licated above | 14  |
| Proforma required? |        | NO                      |                 |               |     |

## Referral criteria

- · Clinical features suggestive of Tay-Sachs Disease
- Biochemical tests supportive of diagnosis

# Requesting specialties

- Clinical Genetics
- Metabolic

#### **TRIMETHYLAMMINURIA**

# Available testing

| Centre                | Method | Scope and range of test |              | Targets      | TAT      |
|-----------------------|--------|-------------------------|--------------|--------------|----------|
| Aberdeen              | Sanger | Whole gene screen       | SNVs, indels | FMO3         | 56       |
| Family member testing |        |                         | as ind       | icated above | 14       |
| Proforma required?    |        | NO                      |              |              | <u> </u> |

## Referral criteria

- Clinical features suggestive of Trimethyamminuria
- Biochemical tests supportive of diagnosis

# Requesting specialties

- Clinical Genetics
- Metabolic

## **VLCAD DEFICIENCY**

# Available testing

| Centre | Method | Scope and range of test | Targets | TAT |
|--------|--------|-------------------------|---------|-----|
|        |        |                         |         |     |





| Aberdeen    | NGS          | Whole gene<br>screen | SNVs, indels<br>Exon level<br>CNV | ACADVL     | 56 |
|-------------|--------------|----------------------|-----------------------------------|------------|----|
| Family me   | mber testing |                      | as indica                         | ated above | 14 |
| Proforma re | equired?     | NO                   |                                   |            |    |

#### Referral criteria

- Clinical features suggestive of VLCAD Deficiency
- · Biochemical tests supportive of diagnosis

- Clinical Genetics
- Metabolic





# **MITOCHONDRIAL**

#### LEBER HEREDITARY OPTIC NEUROPATHY

#### Available testing

| Centre     | Method         | Scope and range of test |                 | Targets  | TAT |
|------------|----------------|-------------------------|-----------------|--|-----|
| Dundee     | Sanger         | Targeted screen         | SNVs,<br>indels | Common LHON mitochondrial DNA variants (m.3460G>A, m.11778G>A, m.14484T>C) | 28  |
| ,          | member<br>ting |                         | as              | indicated above  | 14  |
| Proforma r | equired?       | NO                      |                 |  |     |

#### Referral criteria

- Any individual suspected clinical diagnosis of Leber hereditary optic neuropathy
  - o Bilateral painless subacute visual failure at a young age
  - Optic disk atrophy
  - o Optic nerve dysfunction and absence of other retinal diseases

- Clinical Genetics
- Metabolic
- Neurology
- Ophthalmology





# MITOCHONDRIAL DISORDERS (MERRF, NARP, DEAFNESS AND CARDIOMYOPATHY)

#### Available testing

| Centre                   | Method                    | Scope and range of test |                 | Targets  | TAT |
|--------------------------|---------------------------|-------------------------|-----------------|--|-----|
| Dundee                   | Sanger and pyrosequencing | Targeted<br>screen      | SNVs,<br>indels | Common mitochondrial DNA variants<br>MT-TL1:m.3243A>G<br>MT-TK:m.8344A>G<br>MT-ATP6:m.8993T>G/C<br>Plus others relevant to phenotype | 28  |
| Family member testing as |                           | indicated above         | 14              |  |     |
| Proforma required? NO    |                           | NO                      |                 |  |     |

#### Referral criteria

- Possible mitochondrial disorder caused by mitochondrial DNA variants including individuals with clinical features suggestive of:
  - o chronic progressive external ophthalmoplegia (CPEO)
  - o Kearns-Sayre syndrome
  - myoclonic epilepsy with ragged red fibres (MERRF)
  - o neuropathy, ataxia and retinitis pigmentosa (NARP)
  - maternally inherited Leigh syndrome (MILS)

- Clinical Genetics
- Endocrinology
- Metabolic
- Neurology
- Ophthalmology





# MITOCHONDRIAL ENCEPHALOPATHY, LACTIC ACIDOSIS, AND STROKE-LIKE EPISODERS (MELAS)

#### Available testing

| Centre             | Method                       | Scope and range of test |      | Targets           | TAT |
|--------------------|------------------------------|-------------------------|------|-------------------|-----|
| Dundee             | Pyrosequencing               | Targeted screen         | SNVs | MT-TL1 m. 3243A>G | 28  |
| Family r           | Family member testing as inc |                         |      | icated above      | 14  |
| Proforma required? |                              | NO                      |      |                   |     |

#### Referral criteria

- The most common initial symptoms are seizures, recurrent headaches, stroke-like episodes, cortical vision loss, muscle weakness, recurrent vomiting, and short stature
- Please send a urine sample for adults.

- Clinical Genetics
- Endocrinology
- Metabolic
- Neurology





#### MITOCHONDRIAL INHERITED DIABETES AND DEAFNESS (MIDD)

#### Available testing

| Centre             | Method         | Scope and range of test |        | Targets           | TAT |
|--------------------|----------------|-------------------------|--------|-------------------|-----|
| Dundee             | Pyrosequencing | Targeted screen         | SNV    | MT-TL1 m. 3243A>G | 28  |
| Family r           | nember testing |                         | as ind | icated above      | 14  |
| Proforma required? |                | NO                      |        |                   |     |

#### Referral criteria

- Adult onset sensorineural hearing loss and diabetes or family history suggestive of a diagnosis of maternally inherited diabetes and deafness.
- Please send a urine sample for adults.

- Clinical Genetics
- Endocrinology
- Metabolic





# **MUSCULOSKELETAL**

# **BECKER MUSCULAR DYSTROPHY (BMD)**

### Available testing

| Centre                 | Method | Scope and range of test |                | Targets | TAT |
|------------------------|--------|-------------------------|----------------|---------|-----|
|                        | MLPA   | Targeted screen         | Exon level CNV | DMD     | 28  |
| Glasgow                | Sanger | Whole gene screen       | SNVs, indels   | DMD     | 56  |
| Family member as indic |        | ated above              | 14             |         |     |
| Proforma required?     |        | NO                      |                |         |     |

#### Referral criteria

- Clinical features that include:
  - o Progressive symmetric muscle weakness
  - Increase in serum concentration of creatine kinase (CK)
  - Calf hypertrophy
  - Cardiomyopathy

- Clinical Genetics
- Paediatrics
- Neurology





#### **CHONDRODYSPLASIA PUNCTATA**

#### Available testing

| Centre     | Method       | Scope and range of test |                 | Targets                         | TAT |
|------------|--------------|-------------------------|-----------------|---------------------------------|-----|
| Glasgow    | NGS          | Whole gene screen       | SNVs,<br>indels | AGPS, ARSE, EBP, GNPAT,<br>PEX7 | 56  |
| Family me  | mber testing |                         | as indica       | ted above                       | 14  |
| Proforma r | equired?     | NO                      |                 |                                 |     |

#### Referral criteria

- Stippling involving the epiphyses of the long bones and vertebrae, the trachea and distal ends of the ribs seen on x-ray OR rhizomelia with stippling involving the epiphyses knee, hip, elbow, and shoulder
- OR biochemical evidence of Chondrodysplasia punctata

### Requesting specialties

Clinical Genetics





### **DUCHENNE MUSCULAR DYSTROPHY (DMD)**

#### Available testing

| Centre                | Method | Scope and range of test |                | Targets | TAT |
|-----------------------|--------|-------------------------|----------------|---------|-----|
|                       | MLPA   | Targeted screen         | Exon level CNV | DMD     | 28  |
| Glasgow               | Sanger | Whole gene SNVs, indels |                | DMD     | 56  |
| Family member testing |        | as indicated above      |                |         | 14  |
| Proforma required?    |        | NO                      |                |         | ·   |

#### Referral criteria

- Clinical features that include:
  - Highly elevated serum concentration of creatine kinase (CK)
  - o Delay in motor milestones/frequent falls.
  - o Positive Gowers' sign
  - o Progressive symmetric muscle weakness

- Clinical Genetics
- Paediatrics
- Neurology





# FGFR3 RELATED SKELETAL DYSPLASIA (incl. ACHONDROPLASIA, HYPOCHONDROPLASIA, THANATOPHORIC DYSPLASIA, MUENKE SYNDROME)

### Available testing

| Centre      | Method       | Scope and range of test |              | Targets                         | TAT |
|-------------|--------------|-------------------------|--------------|---------------------------------|-----|
| Edinburgh   | Sanger       | Targeted screen         | SNVs, indels | FGFR3 (exons 7, 10, 13, 15, 19) | 28  |
| Family mem  | nber testing |                         | as ind       | icated above                    | 14  |
| Proforma re | equired?     | NO                      |              |                                 |     |

#### Referral criteria

Clinical features strongly suggestive of FGFR3-related skeletal dysplasias

- Clinical Genetics
- Neonatology
- Orthopaedics
- Paediatrics





#### FIBRODYSPLASIA OSSIFICANS PROGRESSIVA

# Available testing

| Centre             | Method | Scope and range of test |        | Targets         | TAT |
|--------------------|--------|-------------------------|--------|-----------------|-----|
| Glasgow            | Sanger | Targeted screen         | SNVs   | ACVR1 (p.R206H) | 28  |
| Family i           | member |                         | as ind | icated above    | 14  |
| tes                | ting   |                         |        |                 |     |
| Proforma required? |        | NO                      |        |                 |     |

#### Referral criteria

- Congenital malformations of the great toes i.e.hallux valgus, malformed first metatarsal, and/or monophalangism.
- Progressive heterotopic ossification

# Requesting specialties

- Clinical Genetics
- Orthopaedics
- Paediatrics

#### HEREDITARY MULTIPLE OSTEOCHONDROMAS / MULTIPLE EXOSTOSES

### Available testing

| Centre      | Method         | Scope and range of | Scope and range of test        |  | TAT |  |
|-------------|----------------|--------------------|--------------------------------|--|-----|--|
| Glasgow     | NGS            | Whole gene screen  | Whole gene screen SNVs, indels |  | 56  |  |
|             | member<br>ting | as indic           | as indicated above             |  |     |  |
| Proforma re | equired?       | NO                 |                                |  |     |  |

#### Referral criteria

Growths of multiple osteochondromas

- Clinical Genetics
- Orthopaedics
- Paediatrics





# LIMB GIRDLE MUSCULAR DYSTROPHY (LGMD)

#### Available testing

| Centre                | Method         | Scope and range of test |              | Targets         | TAT |
|-----------------------|----------------|-------------------------|--------------|-----------------|-----|
| Glasgow               | Sanger         | Whole gene screen       | SNVs, indels | DES, FKRP, LMNA | 56  |
| Family i              | member<br>ting |                         | as ind       | icated above    | 14  |
| Proforma required? NO |                | NO                      |              |                 |     |

#### Referral criteria

- Progressive weakness and atrophy of the Limb-Girdle muscles AND/OR
- Cardiomyopathy

- Clinical Genetics
- Neurology
- Paediatrics





### **MYOTONIC DYSTROPHY TYPE 1 (DM1)**

#### Available testing

| Centre                           | Method              | Scope and range of test |                          | Targets | TAT              |
|----------------------------------|---------------------|-------------------------|--------------------------|---------|------------------|
| Aberdeen<br>Edinburgh<br>Glasgow | PCR &<br>TP-<br>PCR | Targeted screen         | Triplet repeat expansion | DMPK    | 28<br>Prenatal 3 |
| Proforma required?               |                     | NO                      |                          |         |                  |

#### Referral criteria

- Clinical phenotype that could be consistent with myotonic dystrophy type 1.
- Suggestive features include
  - Hypotonic infant with or without joint contractures
  - Muscle myotonia
  - Muscle weakness
  - Presenile cataracts
  - o Temporal muscle wasting and / or frontal balding
  - o Adverse anaesthetic reaction
  - o Family history of Myotonic Dystrophy
  - Unexplained excessive somnolence or cardiac conduction system abnormalities with additional features as above.

- Clinical Genetics
- Neurology
- Ophthalmology
- Paediatrics





### **MYOTONIC DYSTROPHY TYPE 2 (DM2)**

#### Available testing

| Centre      | Method          | Scope and range of test |                      | Targets | TAT |
|-------------|-----------------|-------------------------|----------------------|---------|-----|
| Aberdeen    | PCR &<br>QP-PCR | Targeted screen         | 4bp repeat expansion | ZNF9    | 28  |
| Proforma re | equired?        | NO                      |                      |         |     |

#### Referral criteria

Clinical phenotype consistent with a diagnosis of Myotonic Dystrophy Type 2

 muscle pain and stiffness, progressive muscle weakness (predominantly proximal and axial), myotonia

# Requesting specialties

- Clinical Genetics
- Neurology

#### **OCULOPHARYNGEAL MUSCULAR DYSTROPHY**

# Available testing

| Centre               | Method            | Scope and range of test |                       | Targets                                   | TAT |
|----------------------|-------------------|-------------------------|-----------------------|---|-----|
| Dundee               | PCR and<br>Sanger | Targeted screen         | Repeat expansion, SNV | PABPN1 – GCN repeat expansion and c.35G>C | 28  |
| Proforma required? N |                   | NO                      |                       |   |     |

#### Referral criteria

• Clinical features strongly suggestive of oculopharyngeal muscular dystrophy.

- Clinical Genetics
- Neurology
- Ophthalmology





#### **OSTEOGENESIS IMPERFECTA**

#### Available testing

| Centre                | Method       | Scope and               | range of test         | Targets  | TAT |
|-----------------------|--------------|-------------------------|-----------------------|--|-----|
| Edinburgh             | NGS          | Whole<br>gene<br>screen | SNVs, indels<br>CNVs* | BMP1, COL1A1*, COL1A2*, CREB3L1, CRTAP,<br>FAM46A, FKBP10, IFITM5, KDELR2, P3H1<br>(LEPRE1), PLOD2, PLS3, PPIB, SERPINF1,<br>SERPINH1, SP7, SPARC, TMEM38B, WNT1 | 112 |
| Family me             | mber testing |                         | as indicated above    |  | 14  |
| Proforma required? NO |              |                         |                       |  |     |

#### Referral criteria

- Multiple fractures of long bones without significant trauma AND at least two of the following:
  - Wormian bones
  - o Blue / grey sclera
  - Hearing loss
  - o Ribs, broad and breaded, thin & irregular
  - Short stature
  - Dentinogenesis imperfect
  - Triangular face & narrow thorax
  - o Round faces & short barrel-shaped chest

# Requesting specialties

Clinical Genetics





#### **OSTEOPETROSIS**

### Available testing

| Centre      | Method         | Scope and range of test |                    | Targets  | TAT |
|-------------|----------------|-------------------------|--------------------|--|-----|
| Glasgow     | NGS            | Whole<br>gene<br>screen | SNVs, indels       | AMER1, ANKH, CA2, CLCN7, CTSK, FAM20C,<br>FERMT3, LEMD3, LRP5, OSTM1, PTH1R,<br>RASGRP2, SNX10, SOST, TCIRG1, TGFB1,<br>TNFRSF11A, TNFSF11, TYROBP | 112 |
| ,           | member<br>ting |                         | as indicated above |  | 14  |
| Proforma re | equired?       | NO                      |                    |  |     |

#### Referral criteria

• Characteristic radiographic changes

- Adult Orthopaedics
- Clinical Genetics
- Paediatrics specialising in bone marrow transplantation, haematology, metabolic disease or orthopaedics





#### PRIMORDIAL DWARFISM, MICROCEPHALY

#### Available testing

| Centre                | Method   | Scope and range of test |              | Targets   | TAT |
|-----------------------|----------|-------------------------|--------------|---|-----|
| Edinburgh             | NGS      | Whole<br>gene<br>screen | SNVs, indels | ANKRD11, ASPM, ATR, ATRX, BLM, CASK, CDC45, CDC6, CDKN1C, CDK5RAP2, CDT1, CENPF, CENPJ, CEP135, CEP152, CEP63, CREBBP, DNA2, DNMT3A (PWWP domain only), DONSON, DPP6, DYRK1A, EP300, GMNN, IGF1, IGF1R, KIF11, KMT2A, KNL1, LARP7, LIG4, MCPH1, MRE11, NBN, NDE1, ORC1, ORC4, ORC6, PCNT, PLK4, PNKP, POC1A, POLE, RAD50, RBBP8, RNU4ATAC, SMARCAL1, SRCAP, STIL, TCF4, TOP3A, TRAIP, TUBGCP6, VPS13B, WDR4, WDR62, XRCC4 | 112 |
| Family member testing |          |                         |              | as indicated above  | 14  |
| Proforma re           | equired? | NO                      |              |   |     |

#### Referral criteria

- Normal microarray
- No history of intrauterine infection, birth hypoxia, teratogens
- OFC smaller than -3SD

#### Requesting specialties

Clinical Genetics

#### PROXIMAL SYMPHALANGISM

### Available testing

| Centre      | Method                             | Scope and range of test |              | Targets    | TAT |
|-------------|------------------------------------|-------------------------|--------------|------------|-----|
| Dundee      | Sanger                             | Whole gene screen       | SNVs, indels | GDF5, NOG1 | 56  |
| ,           | Family member as indicates testing |                         | ated above   | 14         |     |
| Proforma re | equired?                           | NO                      |              |            |     |

#### Referral criteria

• Clinical features strongly suggestive of proximal symphalangism

# Requesting specialties

Clinical Genetics

NSD611-003.20 V





# RASOPATHIES (incl. NOONAN, COSTELLO, CFC, LEGIUS SYNDROMES, NF1 AND NSML) Available testing

| Centre      | Method                | Scope and range of test |                    | Targets   | TAT |
|-------------|-----------------------|-------------------------|--------------------|---|-----|
| Edinburgh   | NGS                   | Whole gene<br>screen    | SNVs, indels       | BRAF, CBL, HRAS, KRAS, LZTR1, MAP2K1,<br>MAP2K2, MRAS, NF1, NRAS, PPP1CB, PTP11,<br>RAF1, RRAS2, RIT1, SHOC2, SOS1, SOS2,<br>SPRED1, SPRED2 | 112 |
| Family men  | nber testing          |                         | as indicated above |   | 14  |
| Proforma re | Proforma required? NO |                         |                    |   |     |

#### Referral criteria

- At least 2 of the suggestive clinical features:
  - o Early feeding difficulty / failure to thrive
  - o Relative macrocephaly
  - Short stature
  - Developmental disability
- At least 1 of:
  - Cardiomyopathy
  - o Congenital heart disease
  - o Arrhythmia
  - Suggestive malignancy (bladder carcinoma, Rhabdomyosarcoma, Leukaemia, phaeochromocytoma)
  - Skin abnormalities (hyperkeratosis, café au lait patches, ulerythema oophorogenes, keratosis pilaris, excess palmar skin)

- Clinical Genetics
- Paediatrics





### SHORT STATURE, INCLUDING TURNER SYNDROME

#### Available testing

| Centre               | Method          | Scope and range of test |                                 | Targets      | TAT |
|----------------------|-----------------|-------------------------|---------------------------------|--------------|-----|
| Aberdeen<br>Dundee   | Karyotype       | Whole genome screen     | Structural rearrangements CNV   | Whole genome | 28  |
| Edinburgh<br>Glasgow | Microarray      | Whole genome screen     | CNV                             | Whole genome | 28  |
| Glasgow              | Sanger,<br>MLPA | Whole gene screen       | SNVs, indels,<br>Exon level CNV | SHOX         | 56  |
| Proforma required?   |                 | NO                      |                                 |              |     |

#### Referral criteria

- Disproportionate short stature
- Idiopathic short stature (males & females)

Other specific features may include

- Premature Ovarian Failure (Turner syndrome)
- Mesomelia and/or Madelung deformity (SHOX-deficiency disorders)

- Clinical Genetics
- Paediatrics





#### SKELETAL DYSPLASIA

(incl. KNIEST DYSPLASIA, CZECH DYSPLASIA, SPONDYLOPERIPHERAL DYSPLASIA, SPONDYLOENCHONDRODYSPLASIA, ACHONDROGENESIS, TYPE II OR HYPOCHONDROGENESIS, SPONDYLOEPIMETAPHYSEAL DYSPLASIA, WEILL-MARCHESANI SYNDROME 1)

#### Available testing

| Centre       | Method   | Scope and ran        | ge of test   | Targets   | TAT |
|--------------|----------|----------------------|--------------|---|-----|
| Edinburgh    | NGS      | Whole gene<br>screen | SNVs, indels | ACAN, ACP5, ADAMTS10, ADAMTSL2, AGPS, ALPL, ANKH, ARSE, B3GALT6, BMP1, BMPR1B, CA2, CANT1, CDC6, CDKN1C, CDT1, CHST3, CLCN7, COL10A1, COL11A1, COL11A2, COL2A1, COL9A1, COL9A2, COMP, CRTAP, CTSK, CUL7, CYP27B1, DHCR24, DLL3, DYM, DYNC2H1, EBP, EIF2AK3, ENPP1, ESCO2, EVC, EVC2, FAM20C, FGF23, FGFR1, FGFR2, FGFR3, FKBP10, FLNA, FLNB, GDF5, GNPAT, GPC6, HSPG2, IFT122, IFT140, IFT43, IFT80, IHH, KAT6B, LBR, LEPRE1, LIFR, LMX1B, LRP5, LTBP2, MATN3, MMP9, NEK1, NPR2, OBSL1, ORC1, ORC4, ORC6, OSTM1, PAPSS2, PCNT, PEX7, PHEX, PLOD2, PPIB, PTH1R, RMRP, RNU4ATAC, ROR2, RUNX2, SBDS, SERPINF1, SERPINH1, SHOX, SLC26A2, SLC34A3, SLC35D1, SLC39A13, SMAD4, SMARCAL1, SNX10, SOX9, TCIRG1, TGFB1, TNFRSF11A, TNFRSF11B, TNFSF11, TRAPPC2, TRIP11, TRPV4, TTC21B, VDR, WDR19, WDR35, WISP3, WNT5A, XYLT1 | 112 |
| Family m     |          |                      | as i         | ndicated above  | 14  |
| Proforma red | <u> </u> | NO                   |              |   |     |

| Centre          | Method  | Scope and range of test |              | Targets                      | TAT |
|-----------------|---------|-------------------------|--------------|------------------------------|-----|
| Edinburgh       | NGS     | Whole gene<br>screen    | SNVs, indels | Nail-patella syndrome: LMX1B | 112 |
| Family m testii |         |                         | as ii        | ndicated above               | 14  |
| Proforma red    | quired? | NO                      |              |                              |     |

#### Referral criteria

- Antenatal evidence (Ultrasound or other imaging modality) or Postnatal evidence of skeletal dysplasia (X ray and clinical examination)
- Multiple joint involvements (e.g. ephyseal or metaphyseal abnormalities)
- Short limbs (Long bone length-3SD below mean or serial measurement at or below 5th centile)
- Narrow thorax
- Poly and/or Oligodactyly
- Syndactyly
- Limb reduction defects
  - Fractures of long bones

NSD611-003.20 V5





- Poor mineralisation of calvarium or spine Requesting specialties
  - Clinical Genetics





### SPINAL AND BULBAR MUSCULAR ATROPHY (SBMA)

# Available testing

| Centre             | Method          | Scope and range of test |                          | Targets | TAT |
|--------------------|-----------------|-------------------------|--------------------------|---------|-----|
| Edinburgh          | PCR &<br>TP-PCR | Targeted screen         | Triplet repeat expansion | AR      | 28  |
| Proforma required? |                 | NO                      |                          |         |     |

#### Referral criteria

• Clinical features that indicate a likely diagnosis of SBMA

- Clinical Genetics
- Neurology





#### **NEUROLOGY**

#### **AICARDI-GOUTIERES SYNDROME**

#### Available testing

| Centre      | Method                     | Scope and range of test |              | Targets   | TAT |
|-------------|----------------------------|-------------------------|--------------|---|-----|
| Dundee      | NGS<br>(clinical<br>exome) | Whole<br>gene<br>screen | SNVs, indels | ADAR, IFIH1, RNASEH2A, RNASEH2B,<br>RNASEH2C, SAMHD1, TREX1 | 112 |
| ,           | Family member testing      |                         |              | as indicated above  | 14  |
| Proforma re | equired?                   | NO                      |              |   |     |

#### Referral criteria

- Individuals with a clinical presentation of the condition:
  - Newborns with a combination of features including enlarged liver and spleen (hepatosplenomegaly), elevated blood levels of liver enzymes, decreased platelets and neurological abnormalities. No evidence of viral infection
  - Children with encephalopathy, sterile pyrexias and seizures, developmental regression, microcephaly, white blood cells in CSF, calcification of the brain, spasticity, dystonia and hypotonia
  - o Isolated 'spastic paraparesis'
  - Singleton Merten syndrome
  - Bilateral striatal necrosis
  - o Familial chilblain lupus

- Clinical Genetics
- Neurology





#### **NEUROMUSCULAR ARTHROGRYPOSIS**

# Available testing

| Centre                | Method | Scope and range of test |                    | Targets  | TAT |
|-----------------------|--------|-------------------------|--------------------|--|-----|
| Glasgow               | NGS    | Whole<br>gene<br>screen | SNVs, indels       | ACTA1, ADAMTS10, ANTXR2, ASCC1, ASXL1, B3GALNT2, B4GAT1, BICD2, CHAT, CHRNA1, CHRNB1, CHRND, CHRNE, CHRNG, CHST14, CNTNAP1, COL12A1, COL6A1, COL6A2, COL6A3, COLQ, DAG1, DNM2, DOK7, DPAGT1, DYNC1H1, ECEL1, ERCC6, ERCC8, EXOSC3, FAM20C, FBN2, FGFR2, FKBP10, FKRP, FKTN, GBA, GBE1, GLDN, GLE1, GMPPB, ADGRG6, HSPG2, ISPD, KLHL40, KLHL41, LAMA2, LARGE1, LMOD3, MAGEL2, MPZ, MTM1, MUSK, MYBPC1, MYH2, MYH3, MYH7, MYH8, NALCN, NEB, PEX1, PEX10, PEX11B, PEX12, PEX13, PEX14, PEX16, PEX19, PEX2, PEX26, PEX3, PEX5, PEX6, PEX7, PFKM, PIEZO2, PLOD1, PLOD2, POMGNT1, POMGNT2, POMK, POMT1, POMT2, POR, PRG4, RAPSN, RYR1, SCARF2, SCN4A, SKI, SLC5A7, SMAD4, STAC3, SYNE1, TMEM5, TNNI2, TNNT1, TNNT3, TPM2, TPM3, TRPV4, TSEN54, UBA1, VAMP1, VIPAS39, VPS33B, ZC4H2 | 112 |
| Family member testing |        |                         | as indicated above | 14   |     |
| Proforma required?    |        | NO                      |                    |  | 1   |

#### Referral criteria

- Antenatally detected joint contractures of more than two different joints OR Born with joint contractures of more than two different joints.
- All cases should have DM1 testing before panel testing.
- Exclusion: Isolated talipes. Finger contractures/camptodactyly with no other joint contractures

Please consider alternative appropriate panels in children with definite cognitive involvement, particularly those where arthrogryposis is mild or additional clinical features are present.

- Clinical Genetics
- Neurology





#### **CADASIL**

### Available testing

| Centre             | Method         | Scope and range of test |              | Targets       | TAT |
|--------------------|----------------|-------------------------|--------------|---------------|-----|
| Glasgow            | Sanger         | Whole gene screen       | SNVs, indels | NOTCH3        | 56  |
|                    | member<br>ting |                         | as in        | dicated above | 14  |
| Proforma required? |                | NO                      |              |               |     |

#### Referral criteria

- · Mid-adult onset of recurrent ischemic stroke
- Cognitive decline progressing to dementia
- A history of migraine with aura
- Diffuse white matter lesions and subcortical infarcts on neuroimaging

- Clinical Genetics
- Neurology





#### **CAPILLARY MALFORMATIONS**

#### Available testing

| Centre             | Method                 | Scope and ra      | ange of test                   | Targets | TAT |
|--------------------|------------------------|-------------------|--------------------------------|---------|-----|
| Dundee             | Sanger<br>MLPA         | Whole gene screen | SNVs, indels<br>Exon level CNV | RASA1   | 56  |
| ,                  | Family member as indic |                   | as indicated                   | above   | 14  |
| Proforma required? |                        | NO                |                                |         |     |

#### Referral criteria

- Capillary malformations are the hallmark of capillary malformationarteriovenous malformation (CM-AVM) syndrome.
- CV-AVM should be suspected in an individual with
  - CM, generally multifocal, small, composed of dilated capillaries, localised on face and limbs
  - AVMs in soft tissue, bone and brain and may be associated with overgrowth
  - o Parkes Weber syndrome phenotype

- Clinical Genetics
- Dermatology
- Neurology





# COGNITIVE CONDITIONS (incl. ALS, FRONTOTEMPORAL DEMENTIA, MOTOR NEURONE DISEASE) Available testing

| Centre                | Method          | Scope and            | range of test                   | Targets   | TAT |
|-----------------------|-----------------|----------------------|---------------------------------|---|-----|
| Edinburgh             | Targeted screen | Repeat-primed PCR    | Hexanucleotide repeat expansion | c90RF72   | 28  |
| Edinburgh             | NGS             | Whole gene<br>screen | SNVs, indels                    | ALS2, ANG, ANXA11, APP,<br>CHCHD10, CHMP2B, CSF1R,<br>DCTN1, FIG4, FUS, GRN, ITM2B,<br>MAPT, NEK1, OPTN, PFN1, PRNP,<br>PSEN1, PSEN2, SETX, SOD1,<br>SQSTM1, TARDBP, TBK1,<br>UBQLN2, VAPB, VCP | 112 |
| Family member testing |                 |                      | as indicated                    | I above   | 14  |
| Proforma required?    |                 | NO                   |                                 |   |     |

#### Referral criteria

- Young onset or familial neurodegeneration starting in adulthood with a likely monogenic cause, including:
  - o 1. Unexplained dementia
    - a. Age at onset <55 years where acquired causes (e.g. stroke, tumour) have been excluded, OR
    - b. Family history of dementia of the same type in a first / second degree relative
- Amyotrophic lateral sclerosis (ALS) with or without frontotemporal dementia
  - a. Evidence of lower motor neuron (LMN) degeneration by clinical, electrophysiologic or neuropathologic examination, AND
  - b. Evidence of upper motor neuron (UMN) degeneration by clinical examination, AND c. Progressive course, AND
  - o d. Age of onset

- Clinical Genetics
- Neurology





#### **COMMON CRANIOSYNOSTOSIS SYNDROMES**

#### Available testing

| Centre                | Method   | Scope and range of test |              | Targets   | TAT |
|-----------------------|----------|-------------------------|--------------|---|-----|
| Edinburgh             | NGS      | Whole gene<br>screen    | SNVs, indels | EFNB1, ERF, FGFR1, FGFR2,<br>FGFR3 TCF12,TWIST1 | 112 |
| Family member testing |          |                         | as indicated | l above   | 14  |
| Proforma re           | equired? | NO                      |              |   |     |

#### Referral criteria

 Recognisable multisuture craniosynostosis syndromes consistent with pathogenic variants in EFNB1, ERF, FGFR1 common hot spots, FGFR2 common hot spots, FGFR3 common hot spots, TCF12 or TWIST1 or with unicoronal or bicoronal craniosynostosis.

# Requesting specialties

• Clinical Genetics





#### **CORTICAL BRAIN MALFORMATIONS**

## Available testing

| Centre                  | Method                     | Scope and rang       | ge of test        | Targets  | TAT |
|-------------------------|----------------------------|----------------------|-------------------|--|-----|
| Dundee                  | NGS<br>(clinical<br>exome) | Whole gene<br>screen | SNVs,<br>indels   | ACTB, ACTG1, ADGRG1 (GPR56), AKT3, ARFGEF2, ARX, ASPM, B3GALNT2, CASK, CCND2, DAG1, DCX, DYNC1H1, EMX2,FKRP, FKTN, FLNA, GPSM2, GRIN1, ISPD, KATNB1, KIF1BP (KIAA1279), KIF2A, KIF5C, LAMA2, LAMB1, LAMC3, LARGE1 (LARGE), MACF1, MTOR, NDE1, NEDD4L, OCLN, PAFAH1B1, PEX1, PEX10, PEX11B, PEX12, PEX13, PEX14, PEX16, PEX19, PEX2, PEX26, PEX3, PEX5, PEX6, PEX7, PIK3CA, PIK3R2, POMGNT1 (GTDC2), POMGNT2, POMT1, POMT2, RELN, RTTN, SMO, TMEM5 (now called RXYLT1), TUBA1A, TUBB, TUBB2A, TUBB2B, TUBB3, TUBG1, | 112 |
| Family member a testing |                            | as                   | s indicated above | 14   |     |
| Proforma r              |                            | NO                   |                   |  | I   |

#### Referral criteria

• Cortical brain malformation with features suggestive of a monogenic cause

- Clinical Genetics
- Neurology





### **CREUTZFELDT-JAKOB DISEASE (CJD)**

### Available testing

| Centre             | Method                | Scor              | pe and range of test         | Targets | TAT |
|--------------------|-----------------------|-------------------|------------------------------|---------|-----|
| Edinburgh          | PCR                   | Repeat-primed PCR | Octapeptide repeat expansion | PRNP    | 28  |
| Edinburgh          | Sanger                | Whole gene screen | SNVs, indels                 | PRNP    | 56  |
|                    | Family member testing |                   | as indicated above           |         |     |
| Proforma required? |                       | NO                |                              |         |     |

#### Referral criteria

• Clinical features that indicate a likely diagnosis of CJD

- Clinical Genetics
- Neurology





#### **DEMENTIA**

### Available testing

| Centre                | Method                     | Scope and range of test |                                 | Targets   | TAT |
|-----------------------|----------------------------|-------------------------|---------------------------------|---|-----|
| Edinburgh             | PCR                        | Targeted screen         | Hexanucleotide repeat expansion | c90RF72   | 28  |
| Dundee                | NGS<br>(clinical<br>exome) | Whole gene<br>screen    | SNVs, indels                    | APP, CHMP2B, CSF1R, DNAJC5,<br>DNMT1,EPM2A, GRN, ITM2B,<br>MAPT, NHLRC1, NOTCH3,<br>PSEN1, PSEN2, PRNP, TBK1,<br>TARDBP, TYROBP, UBQLN2,<br>VCP | 112 |
| Family member testing |                            |                         | as indicated                    | dabove  | 14  |
| Proforma required? NO |                            |                         |                                 |   |     |

#### Referral criteria

- Unexplained dementia with:
  - Age at onset <55 years where acquired causes (e.g. stroke, tumour) have been excluded, OR
  - Family history of dementia of the same type in a first / second degree relative

- Clinical Genetics
- Neurology





# DENTATORUBRAL PALLIODOLUYSIAN ATROPHY (DRPLA)

### Available testing

| Centre             | Method          | Scope ar        | nd range of test         | Targets | TAT |
|--------------------|-----------------|-----------------|--------------------------|---------|-----|
| Edinburgh          | PCR &<br>TP-PCR | Targeted screen | Triplet repeat expansion | ATN1    | 28  |
| Proforma required? |                 | NO              |                          |         |     |

#### Referral criteria

· Clinical features that indicate a likely diagnosis of DRPLA

- Clinical Genetics
- Neurology





#### **DYSTONIA**

### Available testing

| Centre             | Method                     | Scope and range of test |               | Targets   | TAT |
|--------------------|----------------------------|-------------------------|---------------|---|-----|
| Dundee             | NGS<br>(clinical<br>exome) | Whole gene<br>screen    | SNVs, indels  | ACTB, AFG3L2, ANO3, APTX, ATM, ATP1A2, ATP7B, C19orf12, CACNA1A, CHMP2B, CP, CSF1R, CYP27A1, DCAF17, FBXO7, FTL, GFAP, GNAL, HPCA, LYST, NKX2-1, PANK2, PDE10A, PDGFB, PDGFRB, PNKD, PRKRA, PRNP, PRRT2, RNF216, SGCE, SLC19A3, SLC20A2, SLC2A1, SPR, TBK1, THAP1, TIMM8A, TOR1A, TUBB4A, WDR45 | 112 |
| Family member      |                            | as in                   | dicated above | 14  |     |
| testing            |                            |                         |               |   |     |
| Proforma required? |                            | NO                      |               |   |     |

#### Referral criteria

- Unexplained dystonia, chorea or related movement disorder with onset in adulthood with a likely monogenic cause
- Overlapping indications: Parkinson's Disease
- Also can perform a Dopa responsive dystonia panel: GCH1, SPR and TH
- Neurology





#### **EPILEPSY**

# Available testing

| Centre     | Method       | Scope and ran        | ge of test      | Targets  | TAT             |
|------------|--------------|----------------------|-----------------|--|-----------------|
| Glasgow    | Microarray   | Whole genome screen  | CNV             | Whole genome   | 28              |
| Glasgow    | NGS          | Whole gene<br>screen | SNVs,<br>indels | ADSL, AFG3L2, AGAT, ALDH7A1, ARHGEF9, ARX, ATP1A2, ATP1A3, CACNA1A, CASK, CDKL5, CHD2, CHRNA2, CHRNA4, CHRNB2, CLCN4, CLN3, CLN5, CLN6, CLN8, CRH, CSTB, CTSD, DCX, DEPDC5, DNAJC5, DNM1, DOCK7, DYNC1H1, EEF1A2, EFHC1, EPM2A, FLNA, FOXG1, GABRA1, GABRG2, GABRB3, GABRD, GABRG2, GAMT, GLRA1, GLRB, GNAO1, GOSR2, GPHN, GRIN1, GRIN2A, GRIN2B, HCN1, KCNA1, KCNA2, KCNB1, KCNC1, KCNJ10, KCNMA1, KCNQ2, KCNQ3, KCNT1, KCTD7, LGI1, LIS1, MECP2, MEF2C, MFSD8, MOCS1, MOCS2, NEU1, NHLRC1, PCDH19, PIGA, PIK3R2, PLCB1, PNKP, PNPO, POLG, PPT1, PRICKLE1, PRICKLE2, PRRT2, RELN, SCARB2, SCN1A, SCN1B, SCN2A, SCN3A, SCN8A, SCN9A, SLC2A1, SLC6A1, SLC6A5, SLC6A8, SLC9A6, SLC12A5, SLC25A22, SPTAN1, SRPX2, STX1B, STXBP1, SUOX, SYNGAP1, TBC1D24 | 56<br>or<br>112 |
| Family me  | mber testing | 1                    |                 | as indicated above   | 14              |
| Proforma r | equired?     | YES See epilepsy     | referral form   | (see centre website)   |                 |

#### Referral criteria

• Unexplained epilepsy with clinical suspicion of a monogenic cause.

- Clinical Genetics
- Neurology
- Paediatrics





#### **EPISODIC ATAXIA**

#### Available testing

| Centre                | Method | Scope and range of test |                 | Targets        | TAT |
|-----------------------|--------|-------------------------|-----------------|----------------|-----|
| Glasgow               | Sanger | Whole gene screen       | SNVs,<br>indels | KCNA1, CACNA1A | 56  |
| Family member testing |        |                         | as inc          | licated above  | 14  |
| Proforma required?    |        | NO                      |                 |                |     |

#### Referral criteria

Paroxysmal attacks of ataxia and vertigo and/or nausea

# Requesting specialties

- Clinical Genetics
- Neurology

# EPISODIC MOVEMENT, MIGRAINE & EPILEPTIC DISORDERS (BRAIN CHANNELOPATHIES)

### Available testing

| Centre                   | Method                     | Scope and range of test |                 | Targets  |     |
|--------------------------|----------------------------|-------------------------|-----------------|--|-----|
| Dundee                   | NGS<br>(clinical<br>exome) | Whole<br>gene<br>screen | SNVs,<br>indels | ADCY5, ATP1A2, ATP1A3, ATP7B,<br>CACNA1A, CACNB4, GLRA1, GLRB, KCNA1,<br>KCNJ2, KCNMA1, KCNQ2, KCNQ3, PNKD,<br>PRRT2, SCN1A, SCN8A, SLC1A3, SLC2A1,<br>SLC6A5, SPR | 112 |
| Family member<br>testing |                            | as indicated above      |                 |  | 14  |
| Proforma required? No    |                            | NO                      |                 |  |     |

#### Referral criteria

• Unexplained clinical phenotype associated with a brain channel pathy and likely to have a monogenic cause

- Clinical Genetics
- Neurology





#### FAMILIAL CEREBRAL CAVERNOUS MALFORMATIONS (CCM)

#### Available testing

| Centre                | Method                     | Scope and range of test |                                  | Targets           | TAT |
|-----------------------|----------------------------|-------------------------|----------------------------------|-------------------|-----|
| Dundee                | NGS<br>(targeted<br>panel) | Whole gene<br>screen    | SNVs, indels,<br>exon level CNVs | KRIT1, CCM2, CCM3 | 56  |
| Family member testing |                            | as indicated above      |                                  |                   | 14  |
| Proforma required?    |                            | NO                      |                                  |                   |     |

#### Referral criteria

 Individuals with multiple CCMs, or one CCM and at least one other family member with one or more CCMs

### Requesting specialties

- Clinical Genetics
- Neurology

#### **FAMILIAL HEMIPLEGIC MIGRAINE**

#### Available testing

| Centre             | Method | Scope and range of test |        | Targets                        | TAT |
|--------------------|--------|-------------------------|--------|--------------------------------|-----|
| Glasgow            | NGS    | Whole gene              | SNVs,  | ATP1A2, CACNA1A, PRRT2, SCN1A, | 56  |
|                    |        | screen                  | indels | SLC2A1                         |     |
| Family member      |        |                         |        |                                | 14  |
| testing            |        |                         |        |                                |     |
| Proforma required? |        | NO                      |        |                                |     |

#### Referral criteria

- Migraine with aura characterized by the presence of a motor weakness during the aura
- Family history of migraines with aura

- Clinical Genetics
- Neurology





### FRAGILE X TREMOR ATAXIA SYNDROME (FXTAS)

# Available testing

| Centre                           | Method          | Scope and range of test |                          | Targets | TAT |
|----------------------------------|-----------------|-------------------------|--------------------------|---------|-----|
| Aberdeen<br>Edinburgh<br>Glasgow | PCR &<br>TP-PCR | Targeted screen         | Triplet repeat expansion | FMR1    | 28  |
| Proforma required?               |                 | NO                      |                          |         |     |

#### Referral criteria

· Hereditary ataxia with onset in adulthood

# Requesting specialties

- Neurology
- Clinical Genetics

#### FRIEDRICH ATAXIA (FRDA)

### Available testing

| Centre             | Method          | Scope and range of test |                          | Targets | TAT |
|--------------------|-----------------|-------------------------|--------------------------|---------|-----|
| Edinburgh          | PCR &<br>TP-PCR | Targeted screen         | Triplet repeat expansion | FXN     | 28  |
| Proforma required? |                 | NO                      |                          |         |     |

#### Referral criteria

· Clinical features that indicate a likely diagnosis of FRDA

- Clinical Genetics
- Neurology





## **HEREDITARY ATAXIA**

## Available testing

| Centre          | Method          | Scope and               | d range of test           | Targets  | TAT |
|-----------------|-----------------|-------------------------|---------------------------|--|-----|
| Edinburgh       | PCR &<br>TP-PCR | Targeted screen         | Triplet repeat expansions | SCA1, SCA2, SCA3, SCA6, SCA7, FRDA, FMR1   | 28  |
| Edinburgh<br>*  | NGS             | Whole<br>gene<br>screen | SNVs, indels              | AAAS, ABCBT, ABCD1, ADAR, AFG3L2, ALS2, ANO10, APTX, ATL1, ATM, ATP1A3, ATP7B, BSCL2, CACNA1A, CACNA1G, CAPN1, COQ8A, CYP27A1, CYP7B1, DDHD2, FA2H, FGF14, FTL, FXN, GBA2, GCH1, GRID2, HSPD1, IFIH1, ITPR1, KCNA1, KCNC3, KCND3, KIF1A, KIF5A, L1CAM, NIPA1, OPA3, PDYN, PLP1, PNPLA6, POLG, PRKCG, PRNP, PRRT2, REEP1, RTN2, RNaseH2B, SACS, SETX, SIL1, SLC1A3, SLC2A1, SPART, SPAST, SPG11, SPG21, SPG7, SPTBN2, STUB1, SYNE1, TGM6, TMEM240, TTBK2, TTPA, TWNK, UBAP1, VPS13D, WASHC5, ZFYVE26  | 112 |
| Glasgow*        | NGS             | Whole<br>gene<br>screen | SNVs, indels              | AAAS, ABCB7, ABHD12, AFG3L2, AMPD2, ANO10, AP1S2, APTX, ARSA, ATCAY, ATM, ATP1A3, CA8, CACNA1A, CACNA1G, CAMTA1, CASK, CHMP1A, CLN6, COQ8A, COX20, CP, CWF19L1, CYP27A1, CYP2U1, DARS2, DDHD2, DNAJC5, DNMT1, EIF2B1, EIF2B2, EIF2B3, EIF2B4, EIF2B5, ELOVL4, EPM2A, EXOSC3, FGF14, FLVCR1, FOLR1, FXN, GBA2, GJC2, GOSR2, GRID2, GRM1, HEXA, HEXB, ITPR1, KCNA1, KCNC3, KCND3, KCNJ10, KIF1C, MARS2, MMACHC, MRE11A, MTTP, NHLRC1, NPC1, NPC2, OPHN1, PAX6, PDYN, PEX16, PLA2G6, PMPCA, PNKP, PNPLA6, POLG, POLR3A, PRKCG, PRNP, PRRT2, RARS2, RNF170, RNF216, SACS, SAR1B, SEPSECS, SETX, SIL1, SLC1A3, SLC2A1, SLC9A6, SNX14, SPG7, SPTBN2, SRD5A3, STUB1, SYNE1, TGM6, TMEM240, TPP1, TSEN2, TSEN54, TTBK2, TTC19, TTPA, TUBB4A, TWNK, VLDLR, VRK1, WDR73, WDR81, WFS1, WWOX | 112 |
| Family metestin |                 |                         |                           | as indicated above   | 14  |
| Proforma red    | <u> </u>        | NO                      |                           |  | l   |

<sup>\*</sup>For patients referred from East of Scotland, testing performed in Edinburgh

#### Referral criteria

- Targeted screen:
  - Unexplained ataxia with onset in adulthood including where differential diagnosis encompasses STR loci
- NGS panels:
  - o Exclusion of metabolic, neoplastic, alcohol, and drug-related causes
  - o Normal/routine neurological bloods, and vitamin E testing
  - Negative spinocerebellar ataxia repeat expansion panel, including FXTAS and FRDA
  - o MRI neuroimaging normal, or isolated cerebellar atrophy
  - Family history of ataxia, or young age of onset (<50)</li>

## Requesting specialties

- Clinical Genetics
- Neurology

NSD611-003.20 V5

<sup>\*</sup>For patients referred from West of Scotland, testing performed in Glasgow





# HEREDITARY MOTOR AND SENSORY NEUROPATHY (HMSN) / CHARCOT MARIE TOOTH (CMT)

## Available testing

| Centre             | Method         | Scope and range of test |                                    | Targets                    | TAT |
|--------------------|----------------|-------------------------|------------------------------------|----------------------------|-----|
| Aberdeen           | Sanger<br>MLPA | Whole Gene<br>screen    | SNVs, indels<br>Exon level<br>CNV* | PMP22*, MPZ, GJB1,<br>MFN2 | 56  |
| Family me          | mber testing   | as indicated above      |                                    |                            | 14  |
| Proforma required? |                | NO                      |                                    |                            |     |

### Referral criteria

 Clinical suggestive of a hereditary neuropathy – distal muscle weakness and atrophy, clawing of hands, pes cavus

## Requesting specialties

- Clinical Genetics
- Neurology

# HEREDITARY NEUROPATHY WITH LIABILITY TO PRESSURE PALSIES (HLPP / HNPP)

## Available testing

| Centre             | Method         | Scope and range of test |                                   | Targets | TAT |
|--------------------|----------------|-------------------------|-----------------------------------|---------|-----|
| Aberdeen           | Sanger<br>MLPA | Whole Gene<br>screen    | SNVs, indels<br>Exon level<br>CNV | PMP22   | 56  |
| Family me          | mber testing   | as indicated above      |                                   |         |     |
| Proforma required? |                | NO                      |                                   |         |     |

#### Referral criteria

 Clinical suggestive of a hereditary neuropathy - periodic episodes of numbness and palsies following nerve compression or trauma

- Clinical Genetics
- Neurology





## HEREDITARY SPASTIC PARAPLEGIA (HSP)

## Available testing

| Centre       | Method      | Scope ar            | nd range of test            | Targets  | TAT |
|--------------|-------------|---------------------|-----------------------------|--|-----|
| Edinburgh*   | NGS<br>MLPA | Whole gen<br>screen | e SNVs, indels<br>CNV**     | AAAS, ABCB7, ABCD1, ADAR, AFG3L2, ALS2, ANO10, APTX, ATL1**, ATM, ATP1A3, ATP7B, BSCL2, CACNA1A, CACNA1G, CAPN1, COQ8A, CYP27A1, CYP7B1, DDHD2, FA2H, FGF14, FTL, FXN, GBA2, GCH1, GRID2, HSPD1, IFIH1, ITPR1, KCNA1, KCNC3, KCND3, KIF1A, KIF5A, L1CAM, NIPA1, OPA3, PDYN, PLP1, PNPLA6, POLG, PRKCG, PRNP, PRRT2, REEP1**, RNaseH2B, RTN2, SACS, SETX, SIL1, SLC1A3, SLC2A1, SPART, SPAST**, SPG11, SPG21, SPG7**, SPTBN2, STUB1, SYNE1, TGM6, TMEM240, TTBK2, TTPA, TWNK, UBAP1, VPS13D, WASHC5 and ZFYVE26 | 112 |
| Glasgow*     | NGS<br>MLPA | Whole gen<br>screen | e SNVs,<br>indels,<br>CNV** | ABCD1, ADAR, AFG3L2, AIMP1, ALDH18A1, ALS2, AP4B1, AP4E1, AP4M1, AP4S1, ARG1, ATP13A2, ATL1**, BSCL2, B4GALNT1, C12orf65, C19orf12, CAPN1, CYP27A1, CYP2U1, CYP7B1, DDHD1, DDHD2, ERLIN1, ERLIN2, FA2H, FARS2, GBA2, GJC2, HACE1, HSPD1, KIAA0196 (WASHC5), KIDINS220, KIF1A, KIF5A, L1CAM, NIPA1, NT5C2, OPA3, PLP1, PNPLA6, POLR3A, REEP1, RTN2, SACS, SERAC1, SLC16A2, SLC1A4, SLC25A46, SLC2A1, SLC33A1, SPAST**,SPG7, SPG11, SPG20 (SPART), SPG21, TUBB4A, WDR45B, ZEB2ZFYVE26, ZFYVE27                   | 112 |
| Family m     | ember       |                     | I                           | as indicated above   | 14  |
| testi        | <u> </u>    | \/F0                |                             | 100 ( )  |     |
| Proforma red | quirea?     | YES                 | Eainburgh only – F          | ISP referral proforma (see centre website)   |     |

<sup>\*</sup> For patients referred from East of Scotland, testing performed in Edinburgh

## Referral criteria

- Spastic diplegia with upper motor neurone signs
- Aside from bladder or bowel urgency, no other neurological defects\*
- Normal MRI imaging of head and spinal cord
- Normal CSF
- Routine neurological bloods normal

\*If additional neurological defects, please discuss with neurogenetics specialist before requesting

 Unexplained spastic paraplegia of likely monogenic aetiology, where genetic diagnosis will guide management

## Requesting specialties

- Clinical Genetics
- Neurology

NSD611-003.20 V5

<sup>\*</sup> For patients referred from West of Scotland, testing performed in Glasgow





### **HOLOPROSENCEPHALY DISORDERS**

## Available testing

| Centre      | Method                     | Scope and range of test |                    | Targets   | TAT |
|-------------|----------------------------|-------------------------|--------------------|---|-----|
| Dundee      | NGS<br>(clinical<br>exome) | Whole gene<br>screen    | SNVs, indels       | CDON, DHCR7, DISP1, FGF8, FGFR1,<br>GLI2, PTCH1, SHH, SIX3, TGIF1, ZIC2 | 112 |
| ,           | member<br>ting             |                         | as indicated above |   | 14  |
| Proforma re | equired?                   | NO                      |                    |   |     |

### Referral criteria

 Liveborn individual with unexplained holoprosencephaly in whom a chromosomal cause has been excluded by microarray or equivalent

## Requesting specialties

- Clinical Genetics
- Neurology

## **HUNTINGTON DISEASE (HD)**

## Available testing

| Centre       | Method  | Scope and range of test |                          | Targets | TAT              |
|--------------|---------|-------------------------|--------------------------|---------|------------------|
| Edinburgh    | TPPCR   | Targeted screen         | Triplet repeat expansion | HTT     | 14<br>Prenatal 3 |
| Edinburgh    | Linkage | Targeted Screen         | Exclusion testing        | НТТ     | 14<br>Prenatal 3 |
| Proforma red | quired? | NO                      |                          |         |                  |

## Referral criteria

- Clinical features that indicate a likely diagnosis of Huntington disease
- Exclusion testing only where confirmed diagnosis of Huntington disease in the family.

- Clinical Genetics
- Neurology (in consultation with Clinical Genetics)





## **HUNTINGTON DISEASE-LIKE disorders**

## Available testing

| Centre             | Method                                    | Scope and range of test |                          | Targets | TAT |
|--------------------|---|-------------------------|--------------------------|---------|-----|
| Edinburgh          | Flanking<br>PCR                           | Targeted screen         | Triplet repeat expansion | JPH3    | 28  |
| Edinburgh          | Sanger*<br>Flanking<br>PCR and TP-<br>PCR | Targeted screen         | SNVs and Indels          |         | 28  |
| Proforma required? |   | NO                      |                          |         |     |

#### Referral criteria

- Clinical features that indicate a likely diagnosis of Huntington disease-like 2 (for JPH3 testing)
- HD testing has been completed
- For patients with a HD-like phenotype, a screen including testing for *C9orf72*, *PRNP*, SCA17 (*TBP*), DRPLA (*ATN1*) and HDL2 (*JPH3*) repeat expansions, and sequencing of FTL, is also available

Requesting specialties

**Clinical Genetics** 





### **LESCH-NYHAN SYNDROME**

## Available testing

| Centre      | Method         | Scope and range of test |                                | Targets | TAT |
|-------------|----------------|-------------------------|--------------------------------|---------|-----|
| Glasgow     | Sanger<br>MLPA | Whole Gene screen       | SNVs, indels<br>Exon level CNV | HPRT1   | 56  |
| Family me   | mber testing   |                         | as indicated a                 | bove    | 14  |
| Proforma re | quired?        | NO                      |                                |         |     |

### Referral criteria

- Hyperuricaemia
- Psychomotor delay
- · Mild to moderate intellectual disability
- Self-injurious behavior

## Requesting specialties

- Clinical Genetics
- Neurology

## LEUKOENCEPHALOPATHY WITH VANISHING WHITE MATTER

## Available testing

| Centre                | Method                     | Scope and range of test |              | Targets                                | TAT |
|-----------------------|----------------------------|-------------------------|--------------|--|-----|
| Dundee                | NGS<br>(clinical<br>exome) | Whole gene<br>screen    | SNVs, indels | EIF2B1, EIF2B2, EIF2B4, EIF2B5, EIF2B3 | 112 |
| Family member testing |                            |                         | as in        | dicated above                          | 14  |
| Proforma re           | equired?                   | NO                      |              |  |     |

## Referral criteria

 Individuals with unexplained leukodystrophy on neuroimaging with onset in adulthood

- Clinical Genetics
- Neurology





## **NEURODEGENERATION WITH BRAIN IRON ACCUMULATION (NBIA)**

## Available testing

| Centre     | Method         | Scope and range of test |              | Targets  | TAT |
|------------|----------------|-------------------------|--------------|--|-----|
| Glasgow    | NGS            | Whole<br>gene<br>screen | SNVs, indels | ATP13A2, C19ORF12, COASY, CP, DCAF17,<br>FA2H, FTL, FUCA1, KIF1A, KMT2B, MECR, PANK2,<br>PLA2G6, PSEN1, SCP2, SLC39A14, SQSTM1,<br>TRIM32, UBTF, VPS13A, WDR45 | 112 |
| ,          | member<br>ting |                         |              | as indicated above   | 14  |
| Proforma r | equired?       | NO                      |              |  |     |

#### Referral criteria

• Suspected clinical diagnosis in patients with hallmark findings of NBIA, or further assessment of patients with clinical diagnosis of idiopathic NBIA who have had mutations ruled out in other genes.

- Clinical Genetics
- Neurology





## **NEUROFIBROMATOSIS TYPE 1 (NF1)**

## Available testing

| Centre      | Method                     | Scope and range of test |                                | Targets  | TAT |
|-------------|----------------------------|-------------------------|--------------------------------|----------|-----|
| Dundee      | NGS<br>(targeted<br>panel) | Whole gene<br>screen    | SNVs, indels<br>Exon level CNV | NF1      | 56  |
|             | member<br>ting             |                         | as indicat                     | ed above | 14  |
| Proforma re | equired?                   | NO                      |                                |          |     |

#### Referral criteria

- Clinical diagnosis of NF1, as defined below, AND molecular diagnosis is required for management of the proband or for reproductive planning
- Diagnosis requires two of:
  - At least 6 café au lait macules (at least 0.5cm in a child and 1.5cm in an adult)
  - At least 2 subcutaneous or cutaneous neurofibromas
  - Plexiform neurofibroma
  - o Optic glioma
  - At least 2 Lisch nodules
  - o Bony dysplasia (sphenoid wing, long bone bowing, pseudarthrosis)
  - Family history of NF1

- Clinical Genetics
- Paediatrics





### **PAIN DISORDERS**

## Available testing

| Centre      | Method                     | Scope and range of test |                    | Targets   | TAT |
|-------------|----------------------------|-------------------------|--------------------|---|-----|
| Dundee      | NGS<br>(clinical<br>exome) | Whole<br>gene<br>screen | SNVs, indels       | ATL1, ATL3, ELP1, GLA, KIF1A, NGF, NTRK1,<br>PRNP, RAB7A, RETREG1, SCN10A, SCN11A,<br>SCN9A, SEPT9, SPTLC1, SPTLC2, TRPA1, TTR,<br>WNK1 | 112 |
| ,           | member<br>ting             |                         | as indicated above |   | 14  |
| Proforma re | equired?                   | NO                      |                    |   |     |

### Referral criteria

- This includes the disorders:
  - Congenital insensitivity to pain
  - o Inherited erythromelalgia
  - o Paroxysmal extreme pain disorder
  - Small fibre neuropathy
  - o Familial episodic pain syndromes
  - o Hereditary sensory and autonomic neuropathies
  - o Forms of Hereditary sensory neuropathy with prominent sensory loss
- Individuals with a disorder of pain perception, including insensitivity to pain or increased pain perception that is likely to be monogenic in aetiology

- Clinical Genetics
- Neurology





### **PARKINSON'S DISEASE**

## Available testing

| Centre                | Method                     | Scope and range of test |                                    | Targets   | TAT |
|-----------------------|----------------------------|-------------------------|------------------------------------|---|-----|
| Dundee                | NGS<br>(clinical<br>exome) | Whole<br>gene<br>screen | SNVs,<br>indels, Exon<br>level CNV | ATP13A2, ATP1A3, DCTN1, DNAJC13,<br>DNAJC6, FBXO7, FTL, , GCH1, GRN, LRRK2,<br>MAPT, PARK7 (DJ-1), PINK1, PLA2G6, PRKN<br>(Parkin), RAB39B, SLC30A10, SNCA, SPG11,<br>SYNJ1, TH, VPS35<br>CNV in SNCA, PARK2, PINK1, PARK7,<br>ATP13A2, LRRK2, GCH1 and UCHL1 | 112 |
| Family member testing |                            |                         |                                    | as indicated above  | 14  |
| Proforma r            |                            | No                      |                                    |   |     |

### Referral criteria

- Parkinson's disease or complex Parkinsonism
  - o Age at onset <50 years, OR
  - First degree relative affected at <50 years, OR</li>
  - Complex features such as spasticity, gaze palsy, early dementia, early bulbar failure, dyspraxia, ataxia, postural hypotension, cortical sensory loss, brain iron accumulation on MRI brain

- Clinical Genetics
- Neurology





### PELIZAEUS-MERZBACHER DISEASE

## Available testing

| Centre      | Method         | Scope and range of test |   | Targets           | TAT |
|-------------|----------------|-------------------------|---|-------------------|-----|
| Dundee      | Sanger<br>MLPA | Whole gene<br>screen    | SNVs, indels<br>Exon level CNV<br>( <i>PLP1</i> ) | PLP1, GJC2(GJA12) | 56  |
| Family mem  | nber           |                         | as indicated a                                    | above             | 14  |
| testing     |                |                         |   |                   |     |
| Proforma re | quired?        | NO                      |   |                   |     |

## Referral criteria

- Any individual with clinical or imaging features suggestive of a *PLP1* disorder
- Pathogenic variants in *GJC2* are associated with Pelizaeus-Merzbacher-like disease, an autosomal recessive disorder.

- Clinical Genetics
- Neurology





## PERIODIC PARALYSIS, HYPERKALAEMIC

## Available testing

| Centre                | Method | Scope and range of test |      | Targets  | TAT |
|-----------------------|--------|-------------------------|------|--|-----|
| Dundee                | Sanger | Targeted screen         | SNVs | <i>SCN4A</i><br>(p.Leu689lle, p.lle693Thr, p.Thr704Met and<br>p.Met1592Val ) | 56  |
| Family member testing |        |                         |      | as indicated above   | 14  |
| Proforma required? NO |        |                         |      |  |     |

### Referral criteria

- Hyperkalemia (serum potassium concentration >5 mmol/L) or an increase of serum potassium concentration of at least 1.5 mmol/L during an attack of weakness and/or provoking/worsening of an attack by oral potassium intake
- Normal serum potassium between attacks
- Onset before age 20 years.

- Clinical Genetics
- Neurology





## PERIODIC PARALYSIS, HYPOKALAEMIC

## Available testing

| Centre                | Method | Scope and range of test |      | Targets  | TAT |
|-----------------------|--------|-------------------------|------|--|-----|
| Dundee                | Sanger | Targeted screen         | SNVs | CACNA1S (codons 528, 897, 1239)<br>SCN4A (codons 669, 672) | 56  |
| Family member testing |        |                         |      | as indicated above   | 14  |
| Proforma required? NO |        |                         |      |  |     |

#### Referral criteria

 Two or more attacks of muscle weakness with documented serum potassium <3.5 mmol/L</li>

OR

- One attack of muscle weakness and one attack of weakness in one relative with documented serum potassium <3.5 mmol/L OR
- Three or more of the following six clinical/laboratory features:
  - Onset in the first or second decade
  - Duration of attack (muscle weakness involving ≥1 limbs) longer than two hours
  - The presence of triggers (previous carbohydrate rich meal, symptom onset during rest after exercise, stress)
  - o Improvement in symptoms with potassium intake
  - A family history of the condition or genetically confirmed skeletal calcium or sodium channel mutation
  - Positive long exercise test

#### AND

• Exclusion of other causes of hypokalemia (renal, adrenal, thyroid dysfunction; renal tubular acidosis; diuretic and laxative abuse)

- Clinical Genetics
- Neurology





### PERIPHERAL NEUROPATHY

## Available testing

| Centre                | Method | Scope and range of test |              | Targets  | TAT |
|-----------------------|--------|-------------------------|--------------|--|-----|
| Glasgow               | NGS    | Whole<br>gene<br>screen | SNVs, indels | AARS, ATL1, ATP7A, BICD2, BSCL2, CCT5, DCTN1, DNM2, DNMT1, DYNC1H1, EGR2, FAM134B, FGD4, FIG4, GARS, GDAP1, HINT1, HSPB1, HSPB3, HSPB8, IGHMBP2, IKBKAP, INF2, KIF1A, LITAF, LMNA, LRSAM1, MARS, MTMR2, NDRG1, NEFL, NGF, NTRK1, PLEKHG5, PRPS1, PRX, RAB7A, REEP1, SBF2, SCN9A, SETX, SH3TC2, SLC52A1, SLC52A2, SLC52A3, SORD, SPTLC1, SPTLC2, TRPV4, VCP, WNK1, YARS | 112 |
| Family member testing |        |                         |              | as indicated above   | 14  |
| Proforma r            |        | NO                      |              |  |     |

### Referral criteria

- Length dependent neuropathy on neurophysiology AND
- No pathogenic variant on first tier CMT testing (performed in Aberdeen)

## AND one of the following-

- · Genetic diagnosis will alter clinical management
- Genetic diagnosis will influence reproductive decisions

- Clinical Genetics
- Neurology





### **PORENCEPHALY**

## Available testing

| Centre                | Method                     |    | Scope and range of | Targets         | TAT            |     |
|-----------------------|----------------------------|----|--------------------|-----------------|----------------|-----|
| Dundee                | NGS<br>(clinical<br>exome) | \  | Whole gene screen  | SNVs, indels    | COL4A1, COL4A2 | 112 |
| Family member testing |                            |    | as                 | indicated above |                | 14  |
| Proforma re           | equired?                   | NO |                    |                 |                |     |

## Referral criteria

• Any individual with clinical features consistent with the condition

- Clinical Genetics
- Neurology





## **RETT (& RETT-LIKE) SYNDROME**

## Available testing

| Centre                | Method         | Scope an      | e and range of test |                                   | Targets      | TAT |
|-----------------------|----------------|---------------|---------------------|-----------------------------------|--------------|-----|
| Glasgow               | Sanger<br>MLPA | Whole<br>scre | U                   | SNVs, indels<br>Exon level<br>CNV | MECP2, CDKL5 | 56  |
| Family member testing |                | as ind        | icated above        | 14                                |              |     |
| Proforma required?    |                | NO            |                     |                                   |              |     |

## Referral criteria

- Clinical features that include:
  - o Rapid developmental regression in infancy
  - Seizures
  - Severe intellectual disability
  - o Stereotypic hand movements
  - Deceleration of head growth

- Clinical Genetics
- Paediatrics





#### RHABDOMYOLYSIS & METABOLIC MYOPATHIES

#### Available testing

| Centre                | Method   | Scope and range of test |                    | Targets  | TAT |
|-----------------------|----------|-------------------------|--------------------|--|-----|
| Glasgow               | NGS      | Whole<br>gene<br>screen | SNVs, indels       | ACADVL, AGL, ALDOA, ANO5, CACNA1S, CAPN3,<br>CAV3, CPT2, DMD, DYSF, ENO3, ETFA, ETFB,<br>ETFDH, FKRP, GAA, GBE1, GMPPB, GYG1, GYS1,<br>HADHA, HADHB, ISCU, LDHA, LPIN1, PFKM,<br>PGAM2, PGK1, PGM1, PHKA1, PNPLA2, PYGM,<br>RBCK1, RYR1, SLC22A5, TANGO2 | 112 |
| Family member testing |          |                         | as indicated above | 14   |     |
| Proforma r            | equired? | NO                      |                    |  |     |

This panel is intended for patients with isolated skeletal muscle symptoms. Patients with multisystem disease may be more appropriately tested on alternative panels

#### Referral criteria

## Single episode rhabdomyolysis

- ALL MUST FULFIL 2 essential criteria:
  - o CK documented >10,000IU/L associated with muscle pain
  - Mitochondrial myopathy/PEO considered and excluded where appropriate
- IN ADDITION PATIENTS AGED >10 years must fulfil at least one of the following three criteria:
  - No environmental cause AND Accustomed exercise (NOT too much, too fast, too soon)
  - High risk features- exercise intolerance preceding rhabdo +/OR weakness on examination >4mths after event +/OR family history documented rhabdo +/OR biochemistry classical of VLCAD, MADD, or CPT2 +/OR cardiomyopathy
  - o CK>500 IU/L >6 months after rhabdo episode

### Recurrent rhabdomyolysis

- All must fulfil 3 essential criteria:
  - o CK documented >10,000IU/L associated with muscle pain on at least one occasion
  - At least one further episode of acute muscle pain associated with documented CK rise or pigmenturia
  - o Mitochondrial myopathy/PEO considered and excluded where appropriate

### Other criteria for rhabdo panel testing

- Clinical suspicion metabolic myopathy AND any of
  - Moderate to profound XS lipid or glycogen on biopsy
  - Cores/minicores on biopsy
- Muscle MRI characteristic of RYR1

## Requesting specialties

- Clinical Genetics
- Metabolic
- Neurology

NSD611-003.20 V





### SPINAL MUSCULAR ATROPHY

## Available testing

| Centre        | Method   | Scope and ran        | ge of test      | Targets  | TAT |
|---------------|----------|----------------------|-----------------|--|-----|
| Edinburgh     | MLPA     | Targeted screen      | CNV             | SMN1   | 28  |
| Glasgow       | NGS      | Whole gene<br>screen | SNVs,<br>indels | AARS, ASAH1, ATP7A, BICD2, BSCL2, CHCHD10, DCTN1, DNAJB2, DYNC1H1, EXOSC3, EXOSC8, FBXO38, FIG4, GARS, HEXA, HSPB1, HSPB3, HSPB8, IGHMBP2, LAS1L, MATR3, MFN2, PLEKHG5, REEP1, SCO2, SETX, SIGMAR1, SLC52A2, SLC52A3, SLC5A7, SOD1, SORD, SYT2, TRPV4, UBA1, VAPB, VCP, VRK1 | 112 |
| Family member |          |                      | as in           | dicated above  | 14  |
| test          | ing      |                      |                 |  |     |
| Proforma re   | equired? | NO                   |                 |  |     |

#### Referral criteria

- Targeted screen
  - Neonates or infants with unexplained hypotonia where the clinical picture is suggestive of a central cause, i.e. particularly where the baby is not alert, but lethargic or sleepy (part of hypotonic infant screen)
  - clinical features point to a peripheral cause, i.e. particularly where the baby is alert and responsive and the floppiness appears static over a period of days
  - o Carrier testing for partners of confirmed SMN1 carriers.
- Whole gene screen
  - dHMN/SMA clinical phenotype AND
  - Compatible neurophysiology (not required in infants) AND
  - 5q linked SMA excluded (not required in infants)

- Clinical Genetics
- Neurology





## **SPINOCEREBELLAR ATAXIA 8 (SCA8)**

## Available testing

| Centre      | Method          | Scope an        | d range of test                          | Targets | TAT |
|-------------|-----------------|-----------------|--|---------|-----|
| Edinburgh   | PCR &<br>TP-PCR | Targeted screen | Targeted screen Triplet repeat expansion |         | 28  |
| Proforma re | equired?        | NO              |  |         |     |

#### Referral criteria

 Testing only available to patients with a family history of SCA8 where the expansion has been shown to segregate with disease in the family

## Requesting specialties

Clinical Genetics

Here for SMA, the two different labs are combined into one with two lists of indications. But for CMT, the two tests are listed separately and in fact one is called peripheral neuropathy. Periodic paralysis also listed separately. I think its better having them separately since they are separate tests but with the same name so they are listed one after the other. I know this format will change but still easier for everyone to understand.

<sup>\*\*\*\*</sup>Some feedback from clinicians





## **SPINOCEREBELLAR ATAXIA 17 (SCA17)**

## Available testing

| Centre      | Method  | Scope an        | d range of test                          | Targets | TAT |
|-------------|---------|-----------------|--|---------|-----|
| Edinburgh   | PCR     | Targeted screen | Targeted screen Triplet repeat expansion |         | 28  |
| Proforma re | quired? | NO              |  |         | •   |

#### Referral criteria

• Clinical features that indicate a likely diagnosis of SCA17

- Clinical Genetics
- Neurology





## **TORSION DYSTONIA**

## Available testing

| Centre             | Method | Scope and range of test  |  | Targets  | TAT |
|--------------------|--------|--------------------------|--|--|-----|
| Aberdeen<br>Dundee | PCR    | Targeted screen Deletion |  | <i>DYT1</i> (c.907_909del)<br>Gene known as <i>TOR1A</i> | 28  |
| Proforma required? |        | NO                       |  |  |     |

### Referral criteria

- DYT1 early-onset isolated dystonia should be suspected in individuals with
  - o Onset of dystonia before the age of 26
  - Isolated dystonia with no other abnormalities on neurologic examination, normal routine neuroimaging, no known cause of acquired dystonia
  - Family history of early onset dystonia
  - Factors specific to DYT1 early onset isolated dystonia e.g. Ashkanazi
     Jewish ancestry, 2 or more affected limbs.

- Clinical Genetics
- Neurology





## **TUBEROUS SCLEROSIS**

## Available testing

| Centre     | Method                     | Scope and range of test |   | Targets    | TAT |
|------------|----------------------------|-------------------------|---|------------|-----|
| Dundee     | NGS<br>(targeted<br>panel) | Whole gene<br>screen    | SNVs, indels, exon<br>level<br>deletions/duplications | TSC1, TSC2 | 56  |
| _          | member<br>ting             |                         | as indicated al                                       | bove       | 14  |
| Proforma r | equired?                   | NO                      |   |            |     |

#### Referral criteria

- Clinical features suggestive of tuberous sclerosis requiring molecular testing
- Testing should be typically be targeted at those with one or more major features or two or more minor features:
  - Major features:
    - Hypomelanotic macules (at least 3 of at least 5 mm in diameter)
    - Angiofibromas (at least three) or fibrous cephalic plaque
    - Ungual fibromas (at least two)
    - Shagreen patch
    - Multiple retinal hamartomas
    - Cortical dysplasias characteristic of tuberous sclerosis such as tubers and cerebral white matter radial migration lines
    - Subependymal nodules
    - Subependymal giant cell astrocytoma
    - Cardiac rhabdomyomas
    - Lymphangioleiomyomatosis (LAM)
    - Angiomyolipomas (at least two)
  - Minor features:
    - Confetti skin lesions
    - Dental enamel pits (>3)
    - Intraoral fibromas (at least two)
    - Retinal achromic patch
    - Multiple renal cysts
    - Non- renal hamartomas

- **Clinical Genetics**
- Neurology
- Nephrology
- Fetal medicine
- Respiratory medicine





## RENAL

### **ALPORT SYNDROME**

## Available testing

| Centre        | Method     | Scope and range of test |              | Targets                | TAT |
|---------------|------------|-------------------------|--------------|------------------------|-----|
| Edinburgh     | NGS        | Whole gene screen       | SNVs, indels | COL4A3, COL4A4, COL4A5 | 56  |
| Family memb   | er testing |                         | as           | ndicated above         | 14  |
| Proforma requ | ired?      | NO                      |              |                        |     |

### Referral criteria

- Proband with haematuria and ONE of:
- 1. A first degree relative with haematuria or unexplained chronic renal failure, OR
- 2. Histological evidence following electron microscopy on renal biopsy of EITHER Alport syndrome (thickening and splitting of glomerular basement membrane +/- electron lucent areas) OR thin basement membrane disease (TBMD), OR
- 3. Clinical features of Alport syndrome (high tone sensorineural hearing loss or characteristic ophthalmic signs such as perimacular flecks or anterior lenticonus)

- Clinical Genetics
- Nephrology





## **BARTTER SYNDROME & GITELMAN SYNDROME**

## Available testing

| Centre      | Method                     | Scope and range of test |                 |   | Targets                                  | TAT |
|-------------|----------------------------|-------------------------|-----------------|---|--|-----|
| Dundee      | NGS<br>(clinical<br>exome) | Whole g                 | ene screen      | SNVs, indels Exon level CNV (CLCNKB) if appropriate | BSND, CLCNKB, KCNJ1,<br>SLC12A1, SLC12A3 | 56  |
|             | member<br>ting             |                         | as indicated ab |   | е  | 14  |
| Proforma re | equired?                   | YES                     | Renal Gen       | etics Proforma (see centre                          | website)                                 |     |

## Referral criteria

• Any individual with a clinical presentation consistent with either condition.

## Requesting specialties

- Clinical Genetics
- Nephrology

## **CYSTINURIA**

## Available testing

| Centre      | Method                     |          | Scope and                                       | d range of test | Targets        | TAT |
|-------------|----------------------------|----------|---|-----------------|----------------|-----|
| Dundee      | NGS<br>(clinical<br>exome) | Whole go | ene screen                                      | SNVs, indels    | SLC3A1, SLC7A9 | 56  |
|             | member<br>ting             |          | as indicated above                              |                 |                |     |
| Proforma re | equired?                   | YES      | ES Renal Genetics Proforma (see centre website) |                 |                |     |

#### Referral criteria

Any individual with a clinical presentation consistent with the condition.

- Clinical Genetics
- Nephrology





### **NEPHROCALCINOSIS OR NEPHROLITHIASIS**

## Available testing

| Centre      | Method                     | Scope and range of test |     | nge of test     | Targets  | TAT |
|-------------|----------------------------|-------------------------|-----|-----------------|--|-----|
| Dundee      | NGS<br>(clinical<br>exome) | Whole ge<br>screer      |     | SNVs,<br>indels | AGXT, APRT, ATP6V0A4, ATP6V1B1, BSND, CA2,<br>CASR, CLCN5, CLCNKB, CLDN16, CLDN19,<br>CYP24A1, FAN1, FAM20A, GRHPR, HOGA1,<br>HPRT1, KCNJ1, OCRL, PHEX, SLC12A1,<br>SLC22A12, SLC2A9, SLC34A1, SLC34A3, SLC3A1,<br>SLC4A1, SLC7A9, SLC9A3R1, STRADA, XDH | 112 |
| _           | member<br>ting             |                         | ·   |                 | as indicated above   | 14  |
| Proforma re | equired?                   | YES                     | Ren | al Genetics P   | roforma (see centre website)   |     |

## Referral criteria

 Nephrocalcinosis or nephrolithiasis where acquired causes have been excluded.

- Clinical Genetics
- Nephrology
- Endocrinology





# POLYCYSTIC KIDNEY DISEASE, AUTOSOMAL DOMINANT AND RECESSIVE Available testing

| Centre | Method                     | Scope and            | d range of test                                     | Targets  | TAT |
|--------|----------------------------|----------------------|---|--|-----|
| Dundee | NGS<br>(targeted<br>panel) | Whole gene screen    | SNVs, indels<br>Exon level<br>CNV                   | PKD1, PKD2   | 56  |
| Dundee | NGS<br>(targeted<br>panel) | Whole gene<br>screen | SNVs, indels  | PKHD1  | 56  |
| Dundee | NGS<br>(targeted<br>panel) | Whole gene<br>screen | SNVs, indels  | AGT, ALG8, ALG9, ANKS6, CEP164, CEP83,<br>COL4A1, DNAJB11, DZIP1L, GANAB, HNF1B,<br>IFT140, INVS, LRP5, MAPKBP1, NPHP1,<br>NPHP3, NPHP4, PKD1, PKD2, PKHD1,<br>PRKCSH, REN, SEC61B, SEC61A1, SEC63,<br>TMEM67, TSC1, TSC2, TTC21B, UMOD, VHL,<br>WDR19 | 112 |
| _      | member<br>ting             |                      | •   | as indicated above   | 14  |
| J .    |                            |                      | rma (see centre website). No proforma needed for Ph | (D1/2  |     |

#### Referral criteria

- For Autosomal Dominant Polycystic Kidney Disease: Individuals with a suspected or established diagnosis of Autosomal Dominant Polycystic Kidney Disease based on renal imaging.
  - Initial analysis of PKD1 and PKD2 then further analysis of the full cystic kidney panel if appropriate.
- Individuals with a suspected or established diagnosis of Autosomal Recessive Polycystic Kidney Disease based on renal imaging or pathology.
- Onset is typically prenatal, in infancy or early childhood/young adulthood
- The full cystic kidney disease full panel is recommended for individuals that meet the following criteria:
  - Patients with non-syndromic cystic renal disease (excluding acquired cystic disease due to chronic or end stage kidney disease) which is EITHER
  - Clinically not characteristic of ADPKD and underlying diagnosis is required for management purposes, OR
  - Clinically symptomatic disease presenting before the age of 18, OR
  - Clinical diagnosis of ADPKD where a genetic diagnosis is required to influence management

- Clinical Genetics
- Fetal Medicine
- Nephrology
- Paediatrics





### **POLYCYSTIC LIVER DISEASE**

## Available testing

| Centre      | Method                     | Scope and range of test |                   | Targets   | TAT |
|-------------|----------------------------|-------------------------|-------------------|---|-----|
| Dundee      | NGS<br>(targeted<br>panel) | Whole<br>gene<br>screen | SNVs, indels      | AGT, ALG8, ALG9, ANKS6, CEP164, CEP83,<br>COL4A1, DNAJB11, DZIP1L, GANAB, HNF1B,<br>IFT140, INVS, LRP5, MAPKBP1, NPHP1, NPHP3,<br>NPHP4, PKD1, PKD2, PKHD1, PRKCSH, REN,<br>SEC61B, SEC61A1, SEC63, TMEM67, TSC1,<br>TSC2, TTC21B, UMOD, VHL, WDR19 | 112 |
| ,           | member<br>ting             |                         |                   | as indicated above  | 14  |
| Proforma re | equired?                   | Yes                     | Renal Genetics Pr | roforma (see centre website)  |     |

### Referral criteria

 Individuals with a suspected or established diagnosis of Polycystic Liver Disease based on imaging or pathology.

Note this is the same panel as the full polycystic kidney disease panel Requesting specialties

- Clinical Genetics
- Fetal Medicine
- Nephrology
- Paediatrics





### PRIMARY HYPEROXALURIA

## Available testing

| Centre      | Method                     | Scope and range of test |               |                     | Targets             | TAT |
|-------------|----------------------------|-------------------------|---------------|---------------------|---------------------|-----|
| Dundee      | NGS<br>(clinical<br>exome) |                         | e gene<br>een | SNVs, indels        | AGXT, GPHPR, HOGA1  | 56  |
| Family r    | nember<br>ting             |                         |               | as indi             | cated above         | 14  |
| Proforma re | equired?                   | YES                     | Renal Ge      | enetics Proforma (s | see centre website) |     |

### Referral criteria

- Any individual with clinical and biochemical features consistent with the condition.
- Overlapping conditions: Nephrocalcinosis or nephrolithiasis

## Requesting specialties

- Clinical Genetics
- Nephrology

## **PSEUDOHYPOALDOSTERONISM type 1**

## Available testing

| Centre      | Method                     | Scope and range of test |              | Targets                       | TAT |
|-------------|----------------------------|-------------------------|--------------|-------------------------------|-----|
| Dundee      | NGS<br>(clinical<br>exome) | Whole gene<br>screen    | SNVs, indels | NR3C2, SCNN1A, SCNN1B, SCNN1G | 112 |
| Family i    | member<br>ting             |                         | as indi      | cated above                   | 14  |
| Proforma re | equired?                   | NO                      |              |                               |     |

#### Referral criteria

 Any individual with clinical and biochemical features consistent with the condition.

- Clinical Genetics
- Paediatrics





### **RENAL CILIOPATHY**

## Available testing

| Centre     | Method                     | Scope and range of test |                     | Targets   | TAT |
|------------|----------------------------|-------------------------|---------------------|---|-----|
| Dundee     | NGS<br>(clinical<br>exome) | Whole<br>gene<br>screen | SNVs, indels,       | AHI1, ALMS1, ANKS6, ARL13B, ARL6, B9D2, BBS1, BBS10, BBS12, BBS2, BBS4, BBS5, BBS7, BBS9, C2CD3, C5orf42 (CPLANE1), CC2D2A, CEP164, CEP290, CEP41, CEP83, CRB2, CSPP1, DDX59, DHCR7, DYNC2H1, HNF1B, HYLS1,ICK, IFT122, IFT43, INVS, IQCB1, KIF7, LZTFL1, MKKS, MKS1, NEK8, NPHP1, NPHP3, NPHP4, OFD1, PKD1, PKD2, PKHD1 PMM2, RPGRIP1L, SDCCAG8, TCTN1, TCTN2, TCTN3, TMEM138, TMEM216, TMEM231, TMEM237, TMEM67, TRAF3IP1, TTC21B, TTC8, WDPCP, WDR19, WDR35, WDR60 | 112 |
| ,          | member<br>ting             |                         |                     | as indicated above  | 14  |
| Proforma r |                            | YES                     | Renal Genetics Prof | forma (see centre website)  | l   |

## Referral criteria

- Individuals with a suspected clinical diagnosis associated with the above genes
- Relevant medical conditions:
  - Joubert syndrome
  - Alstrom syndrome
  - o Bardet-Biedl syndrome
  - Meckel syndrome
  - o Nephronophthisis
  - o Smith-Lemli-Opitz syndrome
  - o Short rib thoracis dysplasia with or without polydactyly
  - o McKusick-Kaufman syndrome
  - Senior-Loken syndrome

- Clinical Genetics
- Nephrology





## RENAL TUBULOPATHIES, RENAL TUBULAR ACIDOSIS

## Available testing

| Centre      | Method                     | Scope and range of test |                   | Targets  | TAT |
|-------------|----------------------------|-------------------------|-------------------|--|-----|
| Dundee      | NGS<br>(clinical<br>exome) | Whole<br>gene<br>screen | SNVs, indels      | AP2S1, AQP2, ATP1A1, ATP6V0A4, ATP6V1B1,<br>AVPR2, BSND, CA2, CASR, CLCNKB, CLDN16,<br>CLDN19, CTNS, CUL3, CYP24A1, FAH, GATM,<br>GNA11, HNF1B, KCNJ1, KCNJ10, KLHL3, NR3C2,<br>REN, SCNN1A, SCNN1B, SCNN1G, SLC12A1,<br>SLC12A3, SLC22A12, SLC2A9, SLC4A1, SLC4A4,<br>SLC5A2, TRPM6, UMOD, WNK4 | 112 |
| ,           | member<br>ting             |                         |                   | as indicated above   | 14  |
| Proforma re | equired?                   | YES                     | Renal Genetics Pr | roforma (see centre website)   |     |

#### Referral criteria

- Patients with a primary renal tubulopathy presenting as one of the following conditions:
  - Hypokalaemic alkalosis with normal or low blood pressure (e.g. Bartter/Gitelman syndromes), OR
  - Hypokalaemic alkalosis with elevated blood pressure (e.g. Liddle syndrome), OR
  - o Hyperkalaemic acidosis with low/normal BP (PHA type 1), OR
  - o Hyperkalaemic acidosis with elevated BP (PHA type 2), OR
  - o Hypokalaemic acidosis (pRTA and renal Fanconi syndromes), OR
  - o Hypomagnesaemia, OR
  - o Nephrogenic diabetes insipidus, OR
  - Other rare types of renal tubulopathy seen in an expert center
- Overlapping conditions: Nephrogenic diabetes insipidus, Bartter/Gitelman syndromes and Nephrocalcinosis or nephrolithiasis

- Clinical Genetics
- Nephrology





## STEROID RESISTANT NEPHROTIC SYNDROME (SRNS) AND PROTEINURIC RENAL DISEASE

## Available testing

| Centre                | Method                     | Scope and               | d range of test    | Targets  | TAT |
|-----------------------|----------------------------|-------------------------|--------------------|--|-----|
| Dundee                | NGS<br>(clinical<br>exome) | Whole<br>gene<br>screen | SNVs,<br>indels    | ACTN4, ARHGDIA, CLCN5, COL4A3, COL4A4, COL4A5, COQ2, COQ6, COQ8B, CRB2, CUBN, DLC1, EMP2, FAT1, INF2, ITGA3, ITSN1, LAMB2, LMX1B, MAGI2, MYH9, MYO1E, NPHS1, NPHS2, NUP107, OCRL, PAX2, PDSS2, PLCE1, PODXL, SCARB2, SMARCAL1, TNS2, TP53RK, TRPC6, WDR73, WT1 | 112 |
| Family member testing |                            |                         | as indicated above | 14   |     |
| Proforma r            | <u> </u>                   | YES                     | Renal Genetics F   | Proforma (see centre website)  |     |

#### Referral criteria

- Steroid-resistant nephrotic syndrome presenting at any age, OR
- Proteinuria with a histological picture of focal segmental glomerulosclerosis (FSGS) or diffuse mesangial sclerosis (DMS) on biopsy, with no identifiable cause, where a transplant or immunosuppression is planned

- Clinical Genetics
- Nephrology





### **TUBULOINTERSTITIAL KIDNEY DISEASE**

## Available testing

| Centre                | Method                     | Scope and range of test |           |                 | Targets   | TAT |
|-----------------------|----------------------------|-------------------------|-----------|-----------------|---|-----|
| Dundee                | NGS<br>(clinical<br>exome) | Whole gene<br>screen    |           | SNVs,<br>indels | ANKS6, CEP164, CEP83, GATM,<br>HNF1B, INVS, MUC1, NPHP1,<br>NPHP3, NPHP4, REN, TMEM67,<br>TTC21B, UMOD, WDR19 | 112 |
| Family member testing |                            |                         |           | as indi         | icated above  | 14  |
| Proforma required?    |                            | YES                     | Renal Ger | netics Proforma | (see centre website)  |     |

## Referral criteria

- Previously known as hyperuricemic nephropathy, familial juvenile, type 1 & 2 and only UMOD and REN tested. Includes both dominant and recessive TKD.
- Renal impairment caused by tubulointerstitial fibrosis with no glomerular lesion, with no identifiable cause, often associated with medullary cysts, hyperuricaemia or gout, AND
- A first degree relative with TKD or unexplained end-stage renal disease
- Testing note: the majority of pathogenic variants in the MUC1 gene are within a Variable Nucleotide Tandem Repeat (VNTR) region, these are not detectable by this method

- Clinical Genetics
- Nephrology





## **RESPIRATORY**

## **ASTHMA**

## Available testing

| Centre             | Method           | Scope and range of test |     | Targets            | TAT |
|--------------------|------------------|-------------------------|-----|--------------------|-----|
| Dundee             | Real time<br>PCR | Targeted screen         | SNV | ADRB2 p.(Gly16Arg) | 28  |
| Proforma required? |                  | NO                      |     |                    |     |

### Referral criteria

- Asthma patient who may be using or about to be prescribed long acting B2 agonist therapy.
- Some evidence to suggest that homozygotes for arginine at codon 16 (ADRB2 p.(Arg16Arg)) may not benefit from long acting B2 agonist therapy

- Clinical Genetics
- Respiratory





#### **CYSTIC FIBROSIS**

#### Available testing

| Centre                            | Method        | Scope and ra       | ange of test                   | Targets   | TAT                 |  |  |
|-----------------------------------|---------------|--------------------|--------------------------------|---|---------------------|--|--|
| Aberdeen Dundee Edinburgh Glasgow | ARMS          | Targeted screen    | SNVs, indels                   | Common variants   | 28<br>Prenatal<br>3 |  |  |
| Glasgow                           | ARMS          | Targeted screen    | SNVs, indels                   | CFTR newborn screening<br>(p.508del, p.G542*, p.G551D,<br>c.469+1G>T common variants) | 7                   |  |  |
| Edinburgh                         | NGS<br>Sanger | Whole gene screen  | SNVs, indels<br>Exon level CNV | CFTR  | 56                  |  |  |
| Family member testing             |               | as indicated above |                                |   | 14                  |  |  |
| Proforma required?                |               | NO                 |                                |   |                     |  |  |

#### Referral criteria

- Test in an individual clinically likely to be affected with cystic fibrosis:
- Child with clinical suspicion of CF (e.g. recurrent chest infections, failure to thrive, fat malabsorption, neonatal history of meconium ileus), AND
  - A not normal sweat test performed in a recognised experienced test centre/laboratory
     (i.e. sweat chloride >40mM with sufficient sweat obtained; >30mM in infants), OR
  - An additional urgent prenatal situation for the parents or for a close relative, but urgent sweat testing not accessible
- Adult with CT-proven bronchiectasis, AND
  - A not normal sweat test performed in a recognised experienced test centre/laboratory
     (i.e. sweat chloride >40mM with sufficient sweat obtained), OR
  - Chronic suppurative chest infection with colonisation by Pseudomonas and Staph aureus, OR
  - Additional exocrine pancreatic dysfunction
- Idiopathic chronic pancreatitis with exocrine dysfunction (fat malabsorption) with other obvious and acquired causes excluded, AND
  - A not normal sweat test performed in a recognised experienced test centre/laboratory (i.e. sweat chloride >40mM with sufficient sweat obtained), OR
  - Sweat testing not practical, and all other causes excluded
- Male infertility associated with obstructive azoospermia, AND
  - o CBAVD (or isolated CUAVD) diagnosed from expert clinical examination, OR
  - o CBAVD identified at incidental herniotomy
- Fetal echogenic bowel as bright as bone on 2nd trimester scan, AND
  - Both parents not available for carrier testing [if both parents are available, Cystic fibrosis carrier testing should be used instead of an invasive prenatal test], AND
  - Isolated anomaly or <2 other common fetal markers, AND</li>
  - Other more common causes excluded (e.g. IUGR, placental failure, earlier bleeding, infection, raised aneuploidy markers)

- Clinical Genetics
- GP
- Obstetrics
- Paediatrics
- Respiratory





## HEREDITARY HAEMORRHAGIC TELANGIECTASIA, PRIMARY PULMONARY HYPERTENSION

## Available testing

| Centre                | Method      | Scope and range of test |                           | Targets   | TAT |
|-----------------------|-------------|-------------------------|---------------------------|---|-----|
| Edinburgh             | HHT-<br>NGS | Whole gene<br>screen    | SNVs,<br>indels,<br>CNV** | ACVRL1**, ENG**, EPHB4, GDF2, RASA1,<br>SMAD4**   | 56  |
| Edinburgh             | PPH-<br>NGS | Whole gene<br>screen    | SNVs,<br>indels,<br>CNV** | ACVRL1**, ATP13A3, BMPR2**, CAV1,<br>GDF2, EIF2AK4, ENG**, KCNK3, SMAD9,<br>SOX17, TBX4 | 56  |
| Family member testing |             | as indicated above      |                           |   | 14  |
| Proforma required?    |             | NO                      |                           |   | •   |

#### Referral criteria

- HHT: Test where any THREE of the following criteria are met:
  - o 1. Epistaxis: spontaneous, recurrent nose bleeds
  - 2. Telangiectases: multiple, at characteristic sites (lips, oral cavity, fingers, nose)
  - 3. Visceral lesions such as gastrointestinal telangiectasia (with or without bleeding), pulmonary arteriovenous malformation (AVM), hepatic AVM, cerebral AVMs, spinal AVM
  - 4. Family history: a first degree relative with HHT according to these criteria (as above) or an autosomal dominant family history of nosebleeds or first degree relative with cerebral AVM / cerebral haemorrhage / pulmonary or hepatic AVM.
- Alternatively, test where any ONE of the following criteria are met:
  - o A) Personal history of at least one pulmonary AVM\*
  - B) Personal history of two or more AVMs at one or more characteristic sites (pulmonary\*, cerebral, hepatic or spinal)
  - C) Personal history of at least one AVM and severe epistaxis or characteristic telangiectasia or family history
  - D) Personal history of telangiectasia, and refractory or severe epistaxis (e.g. requiring recurrent transfusions)
- \* \*Pulmonary AVM only if confirmed by cross sectional imaging (usually thoracic CT scan), and/or later therapeutic angiography/surgery. Do not diagnose if only supported by a positive right-to-left shunt study ("bubble echo") or chest x-ray
  - Clinical features that indicate a likely diagnosis of PPH.

## Requesting specialties

- Clinical Genetics
- Respiratory
- FNT

NSD611-003.20 V5





### PRIMARY CILIARY DYSKINESIA

## Available testing

| Centre             | Method | Scope and range of test |              | Targets   | TAT |
|--------------------|--------|-------------------------|--------------|---|-----|
| Glasgow            | NGS    | Whole gene<br>screen    | SNVs, indels | ARMC4, C210RF59, CCDC39, CCDC40,<br>CCDC65, CCDC103, CCDC114, CCDC151,<br>CCN0, DNAAF1, DNAAF2, DNAAF3,<br>DNAAF5, DNAH5, DNAH11, DNAI1, DNAI2,<br>DNAL1, DRC1, DYX1C1, GAS8, LRRC6,<br>MCIDAS, RPGR, RSPH1, RSPH4A, RSPH9,<br>SPAG1, ZMYND10 | 112 |
| Family member      |        |                         | а            | as indicated above  | 14  |
| testing            |        |                         |              |   |     |
| Proforma required? |        | NO                      |              |   |     |

#### Referral criteria

- Neonate at least one of the following:
  - Situs inversus plus lower airway or nasal symptoms
  - o Persistent respiratory distress where other causes have been excluded
  - Persistent rhinorrhoea and cough distress where other causes have been excluded
  - o Sibling with PCD
- Childhood at least one of the following:
  - Persistent lifelong wet cough (cystic fibrosis excluded)
  - Unexplained bronchiectasis (cystic fibrosis excluded)
  - Serious otitis media in association with recurrent lower and upper airway symptoms
- Adults
  - Symptoms as above since, often associated with infertility or subfertility

- Clinical Genetics
- Paediatrics
- Respiratory Medicine





#### SURFACTANT METABOLISM DYSFUNCTION

### Available testing

| Centre                | Method   | Scope and ran     | ige of test  | Targets                     | TAT |
|-----------------------|----------|-------------------|--------------|-----------------------------|-----|
| Glasgow               | NGS      | Whole gene screen | SNVs, indels | ABCA3, NKX2-1, SFTPB, SFTPC | 56  |
| Family member testing |          |                   | as in        | dicated above               | 14  |
| Proforma re           | equired? | NO                |              |                             |     |

#### Referral criteria

 Neonatal respiratory insufficiency of disproportionate severity for advanced gestation, with clinical and X-ray features consistent with pulmonary surfactant deficiency AND no other obvious cause for respiratory distress e.g. no difficult delivery, no infection, not premature

- Clinical Genetics
- Intensivists





# **SKIN**

### **ACRAL PEELING SKIN SYNDROME**

# Available testing

| Centre      | Method         | Scope and range of test |             |                  | Targets            | TAT |
|-------------|----------------|-------------------------|-------------|------------------|--------------------|-----|
| Dundee      | Sanger         | Whole ge                | ene screen  | SNVs, indels     | TGM5               | 56  |
|             | member<br>ting |                         |             | as indi          | cated above        | 28  |
| Proforma re | equired?       | YES                     | Skin disord | ers proforma (se | ee centre website) |     |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition:
  - o Painless peeling of the epidermis
  - o Itchy and red skin
  - o Blisters

- Clinical Genetics
- Dermatology





# **AUTOSOMAL RECESSIVE CONGENITAL ICHTHYOSIS (ARCI)**

### Available testing

| Centre                                      | Method                     | Scope and range of test |                    | range of test     | Targets   | TAT |
|---|----------------------------|-------------------------|--------------------|-------------------|---|-----|
| Dundee                                      | NGS<br>(clinical<br>exome) | Whole ge<br>screer      |                    | SNVs, indels      | ABCA12, ALDH3A2, ALOX12B, ALOXE3, CERS3,<br>CYP4F22, NIPAL4, PNPLA1, SLC27A4, ST14,<br>STS, SULT2B1, TGM1 | 56  |
| Family member testing                       |                            |                         | as indicated above | 28                |   |     |
| Proforma required? YES Skin disorders profo |                            |                         | Skir               | n disorders profe | orma (see centre website)   |     |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition
  - o Born with collodion membrane
  - o Thick, hyperkeratotic skin
  - o The later development of at least one of the following:
    - classic lamellar ichthyosis (LI)
    - (nonbullous) congenital ichthyosiform erythroderma (CIE)
    - intermediate forms with some features of both LI and CIE, or nonLI/nonCIE form with mild hyperkeratosis
  - o Excludes Harlequin ichthyosis

- Dermatology
- Clinical Genetics





#### **BIRT-HOGG-DUBE SYNDROME**

# Available testing

| Centre     | Method                     | Scope and range of test |   |                                | Targets   | TAT |
|------------|----------------------------|-------------------------|---|--------------------------------|-----------|-----|
| Dundee     | NGS<br>(targeted<br>panel) | Whole<br>scre           | U | SNVs, indels<br>Exon level CNV | FLCN      | 56  |
|            | member<br>ting             |                         |   | as indica                      | ted above | 14  |
| Proforma r | equired?                   | NO                      |   |                                |           |     |

#### Referral criteria

- Individuals with either:
  - five or more facial or truncal papules with at least one histologically confirmed fibrofolliculoma
- or two of:
  - o early-onset [age <50 years] or multifocal/bilateral renal cell cancer
  - o renal cell cancer with mixed chromophobe/oncocytic histology
  - o multiple lung cysts with or without spontaneous pneumothorax
  - o first degree relative with BHDS

- Clinical Genetics
- Dermatology
- Respiratory





#### **BULLOUS CONGENITAL ICHTHYOSIFORM ERYTHRODERMA**

### Available testing

| Centre      | Method         | Sco | pe and ran    | ge of test       | Targets             | TAT |
|-------------|----------------|-----|---------------|------------------|---------------------|-----|
| Dundee      | Sanger         |     | e gene<br>een | SNVs, indels     | KRT1, KRT10         | 56  |
| Family i    | member<br>ting |     |               | as in            | dicated above       | 28  |
| Proforma re | equired?       | YES | Skin diso     | rders proforma ( | see centre website) |     |

#### Referral criteria

- Also known as Epidermolytic hyperkeratosis (EHK) or Epidermolytic ichthyosis (EI)
- Any individual with a clinical presentation consistent with the condition:
  - Hyperkeratotic scaliness
  - Severe blistering
  - o Hyperproliferation in the basal cells
  - o Thickened, granular layer of the epidermis
  - Skin biopsy recommended if mosaic form suspected (epidermolytic epidermal naevus)

- Clinical Genetics
- Dermatology





#### **ECTODERMAL DYSPLASIA**

# Available testing

| Centre     | Method                     | Scope ar            | nd range of test                             | Targets  | TAT |  |
|------------|----------------------------|---------------------|--|--|-----|--|
| Dundee     | NGS<br>(clinical<br>exome) | Whole ger<br>screen | ne SNVs,<br>indels                           | APCDD1, CDH3, CDSN, DSG4, EDA, EDAR, EDARADD, GJB2, GJB6, GRHL2, HLA-DRA, HOXC13, HR, IKBKG, KRT14, KRT71, KRT74, KRT81, KRT83, KRT85, LIPH, LPAR6, MBTPS2, MSX1, NECTIN1, NECTIN4, NFKB2, NFKBIA, PKP1, PORCN, RSPO4, SNRPE, TP63, TSPEAR, WNT10A | 112 |  |
| ,          | member<br>ting             |                     | ;  | as indicated above   | 28  |  |
| Proforma r | equired?                   | YES                 | Skin disorders proforma (see centre website) |  |     |  |

#### Referral criteria

- Any individual with a clinical diagnosis of ectodermal dysplasia with one or more of the following:
  - abnormality of hair (hypotrichosis, sparse hair, sparse/missing eyebrows)
  - abnormality of teeth (hypodontia, conical incisors)
  - o abnormality of skin (hypohidrosis, episodes of hyperthermia)
- Includes Hypohidrotic X-linked Ectodermal Dysplasia (XHED), Anhidrotic (autosomal dominant and recessive) Ectodermal Dysplasia, Odontoonychodermal Dysplasia (OODD), Clouston syndrome, Witkop syndrome, and Ectrodactyly, Ectodermal Dysplasia and Cleft Lip/Palate syndrome (EEC3)

- Clinical Genetics
- Dermatology





#### **EPIDERMOLYSIS BULLOSA**

#### Available testing

| Centre                 | Method                     | Scope and range of test                      |                 | Targets  | TAT |
|------------------------|----------------------------|--|-----------------|--|-----|
| Dundee                 | Sanger                     | Whole<br>gene<br>screen                      | SNVs,<br>indels | KRT5, KRT14  | 56  |
| Dundee                 | NGS<br>(clinical<br>exome) | Whole gene screen                            | SNVs, indels    | COL17A1, COL7A1, DSP, DST, EXPH5, FERMT1,<br>ITGA3, ITGA6, ITGB4, JUP, KRT14, KRT5, LAMA3,<br>LAMB3, LAMC2, PKP1, PLEC, TGM5 | 112 |
| Family member testing  |                            | as indicated above                           |                 | 28   |     |
| Proforma required? YES |                            | Skin disorders proforma (see centre website) |                 |  |     |

#### Referral criteria

- Includes common types of Epidermolysis bullosa simplex (EBS): localized (EBS-loc, previously known as Weber-Cockayne type), generalized intermediate (EBS-gen intermed, previously known as Koebner type), motteled (EBS-MP) and generalized severe (EBS-gen sev, previously known as Dowling-Meara type)
  - Sanger sequencing for KRT5 and KRT14 for EBS
  - Dowling-Degos Syndrome Sanger sequencing for KRT5
  - Naegeli-Franceschetti-Jadassohn Syndrome Sanger sequencing for KRT14 exon 1
  - NGS test for other rarer forms of EB
- Genetically heterogeneous disorder of skin fragility, manifested by blistering and/or erosions with little or no trauma

- Clinical Genetics
- Dermatology





# **EPIDERMOLYTIC PALMOPLANTAR KERATODERMA (EPPK)**

# Available testing

| Cent   | re Mo               | ethod | Scop | e and ran                      | ge of test      | Targets              | TAT |
|--------|---------------------|-------|------|--------------------------------|-----------------|----------------------|-----|
| Dund   | ee Sa               | anger |      | Whole gene SNVs, indels screen |                 | KRT1, KRT9           | 56  |
| Fai    | nily mem<br>testing | ber   |      |                                | as in           | dicated above        | 28  |
| Profor | ma requir           | red?  | YES  | Skin Disc                      | orders Proforma | (see centre website) |     |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition:
  - Yellow and diffuse thickening of the skin on the palms and soles (palmoplantar keratoderma)
  - o Erythema
  - o Localised epidermolytic hyperkeratosis
  - Onset in infancy

- Clinical Genetics
- Dermatology





#### **FERGUSON-SMITH DISEASE**

# Available testing

| Centre                | Method                | Scope and range of test |              | Targets      | TAT |
|-----------------------|-----------------------|-------------------------|--------------|--------------|-----|
| Dundee                | Sanger                | Whole gene screen       | SNVs, indels | TGFBR1       | 56  |
| Family member testing |                       |                         | as ind       | icated above | 28  |
| Proforma re           | Proforma required? NO |                         |              |              |     |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition:
  - Squamous cell carcinomas or keratoacanthoma which heal spontaneously leaving pitted scars

# Requesting specialties

- Clinical Genetics
- Dermatology

#### FOCAL PALMOPLANTAR KERATODERMA

# Available testing

| Centre      | Method       | Scope and rang  | ge of test   | Targets                          | TAT |
|-------------|--------------|-----------------|--------------|----------------------------------|-----|
| Dundee      | Sanger       | Targeted screen | SNVs, indels | KRT6C (ex1&7), KRT16 (ex1,6,7,8) | 56  |
| Family men  | nber testing |                 | as in        | dicated above                    | 28  |
| Proforma re | quired?      | NO              |              |                                  |     |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition:
  - Focal palmoplantar hyperkeratosis
  - Palmoplantar keratoderma
  - Autosomal dominant

- Clinical Genetics
- Dermatology





#### **GLOMUVENOUS MALFORMATIONS**

# Available testing

| Centre                | Method   | Scope and range of test |              | Targets                         | TAT |
|-----------------------|----------|-------------------------|--------------|---------------------------------|-----|
| Dundee                | Sanger   | Targeted screen         | SNVs, indels | GLMN (exons 2, 3, 6, 8, 12, 13) | 56  |
| Family member testing |          |                         | as ind       | icated above                    | 28  |
| Proforma re           | equired? | NO                      |              |                                 |     |

#### Referral criteria

- A clinical diagnosis of glomuvenous malformations (GVM) based on the International Society for the Study of Vascular Anomalies (ISSVA) classification
- Two or more combined malformations consisting of capillary and venous malformations found in one lesion

- Clinical Genetics
- Dermatology





#### HAIR DISORDERS

# Available testing

| Centre                | Method                     | Scope and range of test |                    | Targets   | TAT |
|-----------------------|----------------------------|-------------------------|--------------------|---|-----|
| Dundee                | NGS<br>(clinical<br>exome) | Whole<br>gene<br>screen | SNVs, indels       | APCDD1, ATP7A, CDH3, CDSN, DSC3, DSG4,<br>EDAR, ERCC2, GJB2, GJB6, HOXC13, HR, JUP,<br>KRT71, KRT74, KRT81, KRT83, KRT85, KRT86,<br>LIPH, LPAR6, MBTPS2, RIPK4, SNRPE, SPINK5,<br>VDR | 112 |
| Family member testing |                            |                         | as indicated above | 28  |     |
| Proforma re           | equired?                   | YES                     | Skin Disorders Pro | oforma (see centre website)   |     |

#### Referral criteria

- Includes Hypotrichosis Simplex, Marine Unna Hypotrichosis, Familial Woolly Hair (WFH), Hypotrichosis with Juvenile Macular Dystrophy, Netherton Syndrome, Monilethrix, Clouston Syndrome, Menkes Syndrome, Hypohidrotic Ectodermal Dysplasia (HED), Trichothiodystrophy (TTD), Ectodermal Dysplasia-9 (ECTD9), Alopecia Universalis Congenita (ALUNC), Naxos Syndrome, CHAND Syndrome, and Atrichia with papular lesions (APL).
- Individuals with a hair disorder with a likely monogenic cause

- Clinical Genetics
- Dermatology





#### **ICHTHYOSIS & ERYTHROKERATODERMA**

#### Available testing

| Centre                | Method                     | Scope and range of test |         | ge of test    | Targets   | TAT |
|-----------------------|----------------------------|-------------------------|---------|---------------|---|-----|
| Dundee                | NGS<br>(clinical<br>exome) | Whole ge<br>screer      | 1 5     | NVs, indels   | AAGAB, ABCA12, ALOX12B, ALOXE3, AQP5,<br>CARD14, CAST, CERS3, CLDN1, CYP4F22,<br>DSC2, DSG1, DSP, ENPP1, FLG, GJA1, GJB2,<br>GJB3, GJB4, GJB6, JUP, KDSR, KRT1, KRT10,<br>KRT14, KRT16, KRT17, KRT2, KRT6A, KRT6B,<br>KRT6C, KRT9, LOR, MSMO1, NIPAL4, PIGL,<br>PNPLA1, RSPO1, RHBDF2, SERPINB7,<br>SLC27A4, SLURP1, SMARCAD1, SNAP29,<br>SPINK5, ST14, STS, SULT2B1, TAT, TGM1,<br>TRPV3 | 112 |
| Family member testing |                            |                         |         |               | as indicated above  | 28  |
| Proforma re           |                            | YES                     | Skin di | sorders profe | orma (see centre website)   |     |

#### Referral criteria

- Clinical presentation with at least two of the following features:
  - o born with collodion membrane
  - o erythroderma
  - o dark plate-like scales or fine white scaling
  - o ectropium/eclabium
  - o hyperkeratosis
- First line testing for punctuate PPK is Sanger sequencing of AAGAB; proceeding to the full panel if negative.
- For ARCI referrals, ARCI panel will be applied in the first instance; proceeding to the full panel if negative and appropriate.

- Clinical Genetics
- Dermatology





#### **ICHTHYOSIS VULGARIS**

# Available testing

| Centre             | Method | Scope and range of test |  |              | Targets  | TAT |  |  |
|--------------------|--------|-------------------------|--|--------------|--|-----|--|--|
| Dundee             | Sanger | Targeted screen         |  | SNVs, indels | <i>FLG</i><br>(p.Arg501*; c.2282_2285delCAGT,<br>p.Arg2447*; p.Ser3247*) | 28  |  |  |
| Family mer testing | mber   | as indicate             | as indicated above                           |              |  |     |  |  |
| Proforma required? |        | YES                     | Skin disorders proforma (see centre website) |              |  |     |  |  |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition
  - Early onset (usually before 1 year old)
  - o Mild ichthyosis/xerosis
  - Keratosis pilaris
  - Hyperlinear pals and soles
  - o Atopic eczema

- Dermatology
- Clinical Genetics





#### **LEGIUS SYNDROME**

# Available testing

| Centre             | Method         | Scope and range of test |                                | Targets | TAT |
|--------------------|----------------|-------------------------|--------------------------------|---------|-----|
| Dundee             | Sanger<br>MLPA | Whole gene<br>screen    | SNVs, indels<br>Exon level CNV | SPRED1  | 56  |
| Family mer testing | mber           | as indicated above      |                                |         | 28  |
| Proforma required? |                | NO                      |                                |         |     |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition:
  - o Five or more café au lait macules which are bilaterally distributed
  - Axillary or inguinal freckling
  - o No other NF1-related criteria

- Clinical Genetics
- Dermatology





#### **MULTIPLE CUTANEOUS AND MUCOSAL VENOUS MALFORMATIONS**

### Available testing

| Centre                | Method   | Scope and ran   | ge of test   | Targets               | TAT |
|-----------------------|----------|-----------------|--------------|-----------------------|-----|
| Dundee                | Sanger   | Targeted screen | SNVs, indels | TIE2 exon 15, exon 17 | 28  |
| Family member testing |          |                 | as in        | dicated above         | 28  |
| Proforma r            | equired? | NO              |              |                       |     |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition:
  - Small, multifocal cutaneous and/or mucosal bluish-purple vascular malformations
  - Early onset (mostly at birth)
  - Slow blood flow on Doppler ultrasound
  - Elevated D-dimer concentration

- Dermatology
- Clinical Genetics





#### **PACHYONYCHIA CONGENITA**

# Available testing

| Centre                | Method   | Scope and range of test |                               | Targets  | TAT |  |  |
|-----------------------|----------|-------------------------|-------------------------------|--|-----|--|--|
| Dundee                | Sanger   | Targeted screen         | SNVs, indels                  | KT6A (ex1&7)<br>KRT6B (ex1&7)<br>KRT6C (ex1&7)<br>KRT16 (ex1,6,7&8)<br>KRT17 (ex1,6&7) | 56  |  |  |
| Family member testing |          | as indicated above      |                               |  |     |  |  |
| Proforma r            | equired? | Yes - SKIN DISO         | Yes – SKIN DISORDERS PROFORMA |  |     |  |  |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition:
  - o Plantar keratoderma including callus with underlying blisters
  - Plantar pain
  - Hypertrophic nail dystrophy, often present within the first few months of life
  - Oral leukokeratosis

- Dermatology
- Clinical Genetics





#### PALMOPLANTAR KERATODERMAS

# Available testing

| Centre     | Method              | Scope and range of test |                      | Targets  | TAT |
|------------|---------------------|-------------------------|----------------------|--|-----|
| Dundee     | Sanger              | Targeted screen         | SNVs, indels         | KRT1, KRT5, KRT9, KRT10  | 56  |
| Dundee     | NGS(clinical exome) | Whole<br>gene<br>screen | SNVs, indels         | AAGAB, ABCA12, ABHD5, ADAM17, ALDH3A2, ALOX12B, ALOXE3, AP1S1, AQP5, ARSE, CAST, CDSN, CERS3, CLDN1, CSTA, CTSC, CYP4F22, DSC2, DSC3, DSG1, DSG4, DSP, EBP, ELOVL4, ENPP1, FLG, GJA1, GJB2, JUP, KANK2, KDSR, KRT1, KRT10, KRT2, KRT6C*, KRT9, LIPN, MBTPS2, MVK, LOR, NIPAL4, NSDHL, PEX7, PHYH, PKP1, PNPLA1, POMP, RHBDF2, RSPO1, SERPINB7, SLC27A4, SLURP1, SNAP29, SPINK5, ST14, STK11, STS, SULT2B1, TGM1, TRPV3, VPS33B | 112 |
| Family me  | ember testing       |                         |                      | as indicated above   | 28  |
| Proforma r | equired?            | YES                     | Skin disorders profe | orma (see centre website)  |     |

#### Referral criteria

- Initial testing by Sanger sequencing for KRT1 and KRT9 (epidermolytic PPK), KRT6c and KRT16 (focal PPK), and KRT6a/b/c, KRT16 and KRT17 (PC) before proceeding to full panel.
- Any individual with a clinical diagnosis of one of the following:
  - o Diffuse palmoplantar keratoderma
  - o Focal keratoderma with or without nail involvement
  - Pachyonychia congenital phenotype
  - o Punctate keratoderma
  - Striate keratoderma with woolly hair
  - Keratoderma with deafness
  - Unusual/unique rare keratodermas occurring alone or as part of syndromes
  - o Erythrokeratoderma

- Clinical Genetics
- Dermatology





#### RARE GENETICS INFLAMMATORY SKIN DISORDERS

#### Available testing

| Centre                | Method              | Scope and range of test |  |              | Targets  | TAT |  |
|-----------------------|---------------------|-------------------------|--|--------------|--|-----|--|
| Dundee                | NGS(clinical exome) | Wh<br>ge<br>scre        | ne   | SNVs, indels | , ADA2, , , AIRE, , , , , CARD11, CARD14,<br>CARD9, , , , , , , , , DOCK8, , EGFR, , , , , ,<br>, , , GJA1, GJB3, , IL1RN, GJB4, IL36RN, KIT, ,<br>, , NCSTN, , NLRP3, NOD2, NSDHL, OSMR, ,<br>PSENEN, RAG1, RAG2, SAMHD1, SH3PXD2B,<br>SLC39A4, STAT3, TMEM173, TREX1 | 112 |  |
| Family member testing |                     |                         |  |              | as indicated above   | 28  |  |
| Proforma required?    |                     | YES                     | YES   Skin disorders proforma (see centre website) |              |  |     |  |

#### Referral criteria

- Any individual with a clinical diagnosis of a rare inflammatory skin disorder of a likely germline genetic cause
  - Includes autoinflammatory disease (e.g. early onset urticaria, recurrent febrile erythemas), infantile pustular psoriasis, likely genetic forms of pityriasis rubra pilaris
- Primary lymphoedema FLT4 analysis. This informs treatment

- Clinical Genetics
- Dermatology
- Rheumatology





# SUPERFICIAL EPIDERMOLYTIC ICHTHYOSIS (SEI)

# (previously known as ICHTHYOSIS BULLOSA of SIEMENS)

### Available testing

| Centre      | Method       | Scope and rang    | ge of test   | Targets     | TAT |
|-------------|--------------|-------------------|--------------|-------------|-----|
| Dundee      | Sanger       | Whole gene screen | SNVs, indels | KRT2        | 56  |
| Family me   | mber testing |                   | as indi      | cated above | 28  |
| Proforma re | equired?     | NO                |              |             |     |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition:
- Erythroderma, widespread blistering, hyperkeratosis with onset at birth

# Requesting specialties

- Dermatology
- Clinical Genetics

#### **VASCULAR SKIN DISORDERS**

# Available testing

| Centre             | Method | Scope and range of test |               |                  | Targets   | TAT |
|--------------------|--------|-------------------------|---------------|------------------|---|-----|
| Dundee             | NGS    |                         | e gene<br>een | SNVs, indels     | ACVRL1, ADAMTS13, ALAS2, ATM, ATR,<br>CCBE1, ENG, EPHB4, F12, FECH, FLT4,<br>FOXC2, GLMN, KRIT1, PIK3CA, PIK3R2,<br>PTEN, RASA1, SCN9A, SMAD4, SOX18,<br>TEK, TMEM173 | 112 |
| Family mentesting  | ,      |                         |               | as i             | ndicated above  | 28  |
| Proforma required? |        | YES                     | Skin diso     | rders proforma ( | (see centre website)  |     |

#### Referral criteria

- Any individual with a vascular skin disorder with a likely germline genetic cause
- Note this method is not optimised to detect mosaic variants

- Clinical Genetics
- Dermatology





#### X-LINKED ICHTHYOSIS

# Available testing

| Centre      | Method         | Scope and range of test |             |                                | Targets     | TAT |
|-------------|----------------|-------------------------|-------------|--------------------------------|-------------|-----|
| Dundee      | Sanger<br>MLPA | Whole gene screen       |             | SNVs, indels<br>Exon level CNV | STS         | 56  |
| Family r    | member<br>ting |                         |             | as indicated a                 | bove        | 28  |
| Proforma re | equired?       | YES                     | Skin disord | lers proforma (see centr       | re website) |     |

#### Referral criteria

- Any individual with a clinical presentation consistent with the condition:
  - Steroid sulfatase (STS) enzyme deficiency
  - o Dry skin
  - o Hyperkeratosis
  - o Hypohidrosis
  - o Ichthyosis

- Dermatology
- Clinical Genetics





# PHARMACOGENOMIC TESTING

# ASTHMA & 2-ADRENERGIC RECEPTOR (ADRB2) p.(Gly16Arg) GENOTYPING

- Asthma patient who may be using or about to be prescribed long acting B2 agonist therapy.
- Some evidence to suggest that homozygotes for arginine at codon 16
   (ADRB2 p.(Arg16Arg)) may not benefit from long acting B2 agonist therapy

# Requesting specialties

- Clinical Genetics
- Respiratory

# AMINOGLYCOSIDE RELATED DEAFNESS MT-RNR1 M.1555A>G GENOTYPING Available testing

| Centre | Method | Scope and ra    | nge of test | Targets           | TAT |
|--------|--------|-----------------|-------------|-------------------|-----|
| Dundee | Sanger | Targeted screen | SNV         | MT-RNR1 m.1555A>G | 5   |

#### Referral criteria

Significant exposure to aminoglycosides posing risk of ototoxicity.

This indication would be relevant to:

- 1. Individuals in whom aminoglycoside therapy may be required
- 2. Individuals who have been exposed to aminoglycosides in whom mt.1555A>G status needs to be determined because of concern regarding hearing loss

### Requesting specialties

Clinical Genetics

Any specialty considering aminoglycoside treatment

NSD611-003.20 V5





# **HEARING LOSS**

#### **AMINOGLYCOSIDE RELATED DEAFNESS**

# Available testing

| Centre             | Method | Scope and range of test |     | Targets   | TAT     |
|--------------------|--------|-------------------------|-----|-----------|---------|
| Dundee             | Sanger | Targeted screen         | SNV | m.1555A>G | 28 or 5 |
| Proforma required? |        | NO                      |     |           |         |

#### Referral criteria

- Significant exposure to aminoglycosides posing risk of ototoxicity
- This indication would be relevant to:
  - Individuals in whom aminoglycoside therapy may be required
  - Individuals who have been exposed to aminoglycosides in whom mt.1555A>G status needs to be determined because of concern regarding hearing loss
- Note TAT is quicker for imminent treatment decisions

- Clinical Genetics
- Any specialty considering aminoglycoside treatment





# **Scottish Strategic Network for Genomic Medicine**

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