# SCOTTISH GENETICS LABORATORY CONSORTIUM

GENOMIC TEST DIRECTORY

FOR

RARE AND INHERITED DISEASE





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## 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 work as a formal consortium arrangement, to deliver 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 was 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 Genetics Laboratory Consortium Genomic Test Directory for Rare and Inherited Disease contains a list of all services currently available in Scotland.

This document will be reviewed annually.

#### NHS SCOTLAND GENETIC LABORATORY CONTACT DETAILS

#### • Aberdeen (NHS Grampian)

Address: Genetics and Molecular Pathology Laboratory Services, Polwarth Building,

Foresterhill, Aberdeen AB25 2ZD

Email address: gram.molgen@nhs.scot

Website: <a href="https://www.nhsgrampian.org/service-hub/north-of-scotland-medical-genetics">https://www.nhsgrampian.org/service-hub/north-of-scotland-medical-genetics</a>

#### Dundee (NHS Tayside)

Address: East of Scotland Regional Genetics Service, Level 6, Ninewells Hospital, Dundee DD1

9SY

Email address: Tay.esrg@nhs.scot

Website: https://www.nhstayside.scot.nhs.uk/OurServicesA-Z/Genetics/PROD\_29554o/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/GeneticLaboratoryServices/Pa

qes/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

 $Website: \underline{https://www.nhsggc.org.uk/about-us/professional-support-sites/laboratory-professional-support-s$ 

medicine/laboratory-disciplines/laboratory-genetics/#

#### **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.

# **CARDIOLOGY**

## **ANDERSEN-TAWIL SYNDROME**

## **AVAILABLE TESTING**

Centre	Method		Scope and ran	ige of test	Targets	TAT
Aberdeen	Sanger	Whole	gene screen	SNVs, indels	KCNJ2, KCNJ5	56
Family me	mber testing	as indica	ated above			14
Proforma required?		YES	Cardiac Arrhy	thmia 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	Sc	ope and rar	nge of test	Targets	TAT
Aberdeen	NGS	Whole ge	ne screen	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 mem	ber testing	as indicate	ed above			14
Proforma re	quired?	YES	Cardiac A	rrhythmia Proforma	a (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 o	gene screen	SNVs, indels	PKP2, DSG2, DSC2, DSP, SCN5A, ABCC9, DES, HCN4, JUP, LMNA, PLN, RYR2, TGFB3, TMEM43	112
Family mem	nber testing		as indicated above			14
Proforma re	equired?	YES	Cardiac Arrh	nythmia Proforma (	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			Targets	TAT
Aberdeen	NGS	Whole gene screen		SNVs, indels	SCN5A, ABCC9, GJA1, GJA5, HCN4, KCNA5, KCNE5, NPPA, SCN2B, SCN4B	56
Family mem	nber testing		as indicated above			
Proforma re	equired?	YES	Cardiac Arrh	nythmia 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 ran	ige of test	Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	TAZ	56
Family me	mber testing		as indicated al	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	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole ge	ne screen	SNVs, indels	SCN5A, CACNA1C, CACNA2D1, CACNB2, GPD1L, HCN4, KCND3, KCNE3, KCNE5, KCNJ8, RANGRF, SCN1B, SCN2B, SCN3B, SCN10A, SLMAP, TRPM4	112
Family men	nber testing	as indic			ated above	14
Proforma re	equired?	YES	Cardiac Ar	rhythmia Proforma	a (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			Targets	TAT
Aberdeen	NGS	Whole gene screen		SNVs, indels	RYR2, CALM1, CALM2, CASQ2, DPP6, TRDN	56
Family mem	ber testing		as indicated above			
Proforma required?		YES	Cardiac A	rrhythmia 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 rar	ge of test	Targets	TAT		
Edinburgh	NGS	Whole gene screen	SNVs, indels	ACTC1, ACTN2, BAG3, CSRP3, DES, DMD, DSP, FLNC, LAMP2, LMNA, MYBPC3, MYH7, MYL2, MYL3, NKX2-5, PLN, RBM20, SCN5A, TNNC1, TNNI3, TNNI3K, TNNT2, TPM1, TTN (N2-B isoform), VCL	56		
Family member testing			as indicated above				
Proforma re	quired?	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, TRPM4	56
Family member testing				as indicated al	bove	14
Proforma required?		YES	Cardiac Arrhy	thmia 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**

Centre	Method	Scope and ran	ge of test	Targets	TAT	
Edinburgh	NGS	Whole gene screen	SNVs, indels	ACTC1, ACTN2, CSRP3, FHL1, FLNC, GLA, JPH2, LAMP2, MYBPC3, MYH7, MYL2, MYL3, PLN, PRKAG2, TNNC1, TNNI3, TNNT2, TPM1, TTR	56	
Family men	nber testing	as indicated above				
Proforma re	quired?	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 screen		SNVs, indels Exon level CNV*	KCNQ1*, KCNH2*, KCNE1*, KCNE2*, SCN5A, KCNJ2, ANK2, AKAP9, CACNA1C, CALM1, CALM2, CAV3, KCNJ5, NOS1AP, SCN4B, SNTA1, TRPM4	112
Family member testing		as indicated above			14	
Proforma required?		YES	Cardiac Arrhy	thmia 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 ran	ige of test	Targets	TAT
Dundee	NGS	Whole gene screen	SNVs, indels	FBN1, TGFBR1, TGFRB2	56
Family member testing			as indicated al	pove	14
Proforma required?		NO			

#### REFERRAL CRITERIA

- Clinical features of Marfan syndrome giving a Ghent systemic score of ≥ 7 in an adult over 18
  years
- 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**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gen screen	e SNVs, indels Exon level CNV	AARS2, ABCC9, ACAD9, ACADVL, ACTA1, ACTC1, 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, NDUFAF3, NDUFAF4, NDUFAF5, NDUFS1, NDUFS8, NDUFS1, 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, TNN13, TNN13K, TNNT2, TPM1, TSFM, TTN, TTR, VCL	112
Family member testing			·	as indicated above	14
Proforma re	Proforma required?				

#### 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. To be finalised with NICCs.

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

## **SHORT QT SYNDROME**

## **AVAILABLE TESTING**

Centre	Method		Scope and rar	nge of test	Targets	TAT
Aberdeen	NGS	Whole	gene screen	SNVs, indels	KCNQ1, KCNH2, KCNJ2	56
Family mem	Family member testing		as indicated above			
Proforma required?		YES	Cardiac Arrhy	Cardiac Arrhythmia Proforma (see centre website)		

## REFERRAL CRITERIA

- Abnormal ECG (QTc ≤36oms in males, ≤37oms 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	Whole gene screen	SNVs, indels	ACTA2, BGN, COL1A1, COL1A2, COL3A1, COL5A1, COL5A2, EFEMP2, ELN, FBN1, FBN2, FLCN, FLNA, LOX, MED12, MYH11, MYLK, NOTCH1, PLOD1, PRKG1, SKI, SLC2A10, SMAD2, SMAD3, SMAD4, SMAD6, TGFB2, TGFB3, TGFBR1, TGFBR2	112
Family member testing		as indicated above			
Proforma required?		40			

#### 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 ≥ 7 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 rar	nge 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			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 ra	nge of test	Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	APTX	56
Family member testing			as indicated	d above	14
Proforma required?		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 ra	nge of test	Targets	TAT
Aberdeen	Chromosome breakage analysis*	Whole genome screen	Aneuploidy	Genome wide	28
Aberdeen	NGS	Whole gene screen	SNVs, indels	BLM	56
Family member testing			as indicated	l 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, sun-sensitive, telangiectatic, hypo- and hyperpigmented skin
- Confirmed diagnosis from chromosome breakage analysis

- Clinical Genetics
- Haematology

## 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 men	nber testing		as indicated 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 screen	SNVs, indels	ERCC6, ERCC8	56
Family men	nber 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 men	nber testing		as indicated above		
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 breakage analysis*	Whole genome screen	Chromosome breakage	Whole genome	28
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV (limited genes*)	BRCA2, BRIP1, ERCC4, FANCA, FANCB, FANCC, FANCD2, FANCE, FANCF, FANCG, FANCI, FANCL, PALB2, RAD51C, SLX4	56
Family men	nber testing		as indica	ated above	14
Proforma re	equired?	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 gene screen	SNVs, indels Exon level CNVs	ACAN, ANKRD11, BLM, BRAF, BRCA2, BRIP1, CBL, CCDC8, CDKN1C, CUL7, ERCC4, FANCA, FANCB, FANCC, FANCD2, FANCE, FANCF, FANCG, FANCI, FANCL, FGFR3, HMGA2, HRAS, IGF1, IGF1R, IGF2, KRAS, LZTR1, MAP2K1, MAP2K2, NBN, NRAS, OBSL1, PALB2, PIK3R1, PLAG1, PPP1CB, PTPN11, RAF1, RIT1, SHOC2, SLX4, SOS1, SOS2, SRCAP, TOP3A, TRIM37, UBE2T	56
Family men	nber testing			as indicated above	14
Proforma re	quired?	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 men	nber testing		as indicated above		
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 1, 9 & 16	Chromosomes 1, 9 & 16	28
Aberdeen	NGS	Whole gene screen	SNVs, indels	DNMT <sub>3</sub> B	56
Family men	nber testing		as indicated above		14
Proforma required? NO		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 men	nber testing		as indicated a	14	
Proforma re	equired?	NO			

## REFERRAL CRITERIA

 Clinical phenotype suggestive of LIG<sub>4</sub> 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, CDC6	56
Family men	nber testing		as indicated above		14
Proforma re	quired?	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 men	nber testing		as indicated above		
Proforma re	quired?	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	mber testing		as indica	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	ESCO <sub>2</sub>	56
Family men	nber testing		as indicated above		14
Proforma required? NO		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 men	nber testing		as indicated above		
Proforma required? NO		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 men	nber testing	as indicated above			14
Proforma re	quired?	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 men	nber testing		as indicated above		
Proforma re	quired?	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 men	nber testing		above	14	
Proforma re	quired?	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, GTF2H5	56
Family men	nber testing		as indicated above		
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	TBX <sub>3</sub>	56
Family men	nber testing		as indic	ated above	14
Proforma re	quired?	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 men	nber testing as indica			ated above	14
Proforma required? NO		NO			

## REFERRAL CRITERIA

• Clinical phenotype suggestive of Warsaw Breakage Syndrome

## **REQUESTING SPECIALTIES**

## **WERNER SYNDROME**

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels	WRN	56
Family men	nber testing	as indicated above			14
Proforma required?		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	nge of test	Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels	XPA, XPC, ERCC1, ERCC3, ERCC4, ERCC5, DDB2, POLH	56
Family men	nber testing		as indicated above		
Proforma re	quired?	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 ra	nge of test	Targets	TAT
Edinburgh	NGS	Whole gene screen	SNVs, indels	ABCC6, ACTA2, ACVR1, ADAMTS2, ALPL, ATP6VoA2, B3GALT6, B4GALT7, BMP1, CBS, CHST14, COL11A1, COL1A1, COL1A2, COL2A1, COL3A1, COL5A1, COL5A2, CRTAP, ELN, FBLN5, FBN1, FBN2, FKBP10, FKBP14, IFITM5, LEPRE1 (P3H1), LRP5, MYLK, NOTCH1, NOTCH2, PKD2, PLOD1, PLOD2, PPIB, PRDM5, RIN2, SERPINF1, SERPINH1, SLC2A10, SLC39A13, SMAD3, SP7, TGFB2, TGFBR1, TGFBR2, TNXB, ZNF469	112
Family men	nber testing		as indicated above		14
Proforma required? NO				•	

## REFERRAL CRITERIA

- See criteria for Ehlers-Danlos Syndrome, Stickler Syndrome / Cleft Palate
- Please contact the laboratory to discuss indications not included by above criteria

# **REQUESTING SPECIALTIES**

## **EHLERS DANLOS SYNDROME**

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	NGS	Whole gene screen	SNVs, indels	ADAMTS2, B3GALT6, B4GALT7, COL1A1, COL1A2, COL3A1, COL5A1, COL5A2, CHST14, PLOD1, FKBP14, RIN2, PRDM5, ZNF469, SLC39A13, TNXB, C1R, C1S	112
Family mer	nber testing			as indicated above	
Proforma re	equired?	NO			

## 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**

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	NGS	Whole gene screen	SNVs, indels	ANKRD11, COL11A1, COL11A2, COL2A1, COL9A1, COL9A2, COL9A3, FLNA, FLNB, FOXE1, FOXC2, IRF6, IRF7, PLOD3, SATB2, SLC26A2, SOX9, TP63, TBX1, TBX22	112
Family mem	nber testing			as indicated above	
Proforma re	quired?	NO			

## REFERRAL CRITERIA

- Two or more of the following:
  - o Retinal detachment or: High myopia with onset before 6 years
  - Cleft palate
  - o Vitreous abnormality
  - o Joint hypermobility or premature joint degeneration
  - o 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	Targeted screen Aneuploidy		Chromosomes 13, 18, 21	7	
Proforma required?	YES	NIPT request form				

## 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 Dundee Edinburgh 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 Dundee Edinburgh 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 Dundee	QF- PCR	Targeted screen	STRs	Chromosomes 13, 18, 21, X/Y	5
Edinburgh Glasgow	Microarray	Whole genome screen	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 Dundee	QF- PCR	Targeted screen	STRs	Chromosomes 13, 18, 21, X/Y	28
Edinburgh Glasgow	Microarray	Whole genome screen CNV		Whole genome	25
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	Targete	d screen	CNV Methylation abnormalities	15q11-13 markers	28
Glasgow	Sanger	Whole ge	ne screen	SNVs, indels	UBE3A	56
Glasgow	PCR	Targete	argeted screen STRs		Microsatellite markers	28
Family men	nber testing			as indicated above		14
Proforma re	quired?	NO				

## REFERRAL CRITERIA

- Clinical features that include:
  - o 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**

Centre	Method	Scope and range of test			Targets	TAT
Glasgow	MLPA	Target	ed screen	CNV Methylation abnormalities	11p15 markers	28
Glasgow	PCR	Target	ed screen	STRs	Microsatellite markers	28
Proforma re	quired?	NO				

# REFERRAL CRITERIA

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

- Clinical Genetics
- Paediatrics

## **CHARGE SYNDROME**

## **AVAILABLE TESTING**

Centre	Method	Scope and ra	nge of test	Targets	TAT
Glasgow	Sanger	Whole gene screen MLPA	SNVs, indels Exon level CNV	CHD <sub>7</sub>	56
Family member testing			as indicated above		14
Proforma required?		NO			

## REFERRAL CRITERIA

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

## **REQUESTING SPECIALTIES**

# CONGENITAL ABNORMALITIES, MULTIPLE

# AVAILABLE TESTING

Centre	Method	Scope and r	ange of test	Targets	TAT
Aberdeen Dundee	Microarray	Whole genome	Structural variants	Whole genome	14
Edinburgh Glasgow	Edinburgh Glasgow Karyotype		screen CNV		.
Proforma required?		NO			

# REFERRAL CRITERIA

• Multiple abnormalities present at birth

- Clinical Genetics
- Paediatrics

# CORNELIA DE LANGE, ATYPICAL

# **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	NGS	Whole gene screen	SNVs, indels	NIPBL, SMC1A, SMC3, HDAC8, RAD21, ANKRD11, KMT2A, AFF4, NAA10	56
Family member	rtesting	as indicated above			
Proforma required	d?	NO			

# REFERRAL CRITERIA

- Normal Karyotype or array CGH
- Characteristic facial appearance (including long eyelashes, upturned nose, long philtrum)
- Post-natal growth delay
- Developmental delay

- Clinical Genetics
- Paediatrics

# **DEVELOPMENTAL DELAY**

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen					
Dundee	Microarray	Whole gene screen	CNV	Whole genome	28
Edinburgh Glasgow					
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

## **DEVELOPMENTAL DISORDERS**

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Edinburgh	NGS	W	hole gene screen	SNVs, indels	DDG <sub>2</sub> P	112
Family m	Family member testing as indicated above			as indicated above		14
Proforma re	quired?	NO	DECIPHER entry required			

## REFERRAL CRITERIA

- Unexplained intellectual disability or global developmental delay where clinical features are suggestive of an underlying monogenic disorder requiring sequencing and targeted genetic testing is not possible
- Microarray analysis has previously been performed on a sample from the proband
- Samples from a proband plus both parents (trio) are required

## **REQUESTING SPECIALTIES**

# Di GEORGE (22q11 DELETION) SYNDROME

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen Dundee Edinburgh Glasgow	Microarray / MLPA	Targeted screen	CNV	Di George Critical Region	28 Urgent 5 Prenatal 14
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 and 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 rang	ge of test	Targets	TAT
Aberdeen Dundee Edinburgh Glasgow	Microarray	Whole genome screen	CNV	Whole genome	28
Glasgow	NGS	Whole gene screen	SNVs, 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 me	Family member testing		as in	dicated above (Glasgow)	14
Proforma req	Proforma required?		al form (see c	entre 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 a	nd range of test	Targets	TAT
Aberdeen Edinburgh Glasgow	PCR & 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

- Clinical Genetics
- Paediatrics

# **INFERTILITY, MALES**

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen Dundee Edinburgh Glasgow	Karyotype	Whole genome screen	Structural variants, CNV	Whole genome	28
Aberdeen Dundee Edinburgh Glasgow	ARMS	Targeted screen	SNVs, indels	50 common mutations	28
Dundee Edinburgh Glasgow	PCR	Targeted screen	Y chromosome markers	AZFa, AZFb, AZFc	28
Proforma required?		NO			

## REFERRAL CRITERIA

- Karyotype 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
  - o CBAVD (or isolated CUAVD) diagnosed from expert clinical examination, OR
  - o CBAVD identified at incidental herniotomy

- Clinical Genetics
- Fertility clinics

# **INFERTILITY, FEMALES**

## **AVAILABLE TESTING**

Centre	Method	Scope ar	nd range of test	Targets	TAT
Aberdeen Dundee	Karyotype	Whole genome	Structural variants,	Whole genome	28
Edinburgh Glasgow	Karyotype	screen	CNV	whole genome	20
Aberdeen Edinburgh	PCR	Targeted screen	Triplet repeat expansion	FMR1	28
Glasgow 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 ovarian failure, AND
- Non genetic causes have been excluded including presence of thyroid and adrenal autoantibodies

- Clinical Genetics
- Fertility clinics

# KLINEFELTER SYNDROME

# AVAILABLE TESTING

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen Dundee	Karyotype	Whole general series	CNV	Whole general	0
Edinburgh Glasgow	Microarray	- Whole genome screen	CIVV	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 Dundee Edinburgh Glasgow	Microarray	Whole genome 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

## **MOLAR PREGNANCY**

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Dundee Edinburgh	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

- Molar pregnancy may be suspected during routine booking scan, or at emergency presentation in clinic. In the majority of cases, molar pregnancy is 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 molar pregnancy is 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	e and range of test	Targets	TAT
Glasgow	MLPA	Targeted screen	CNV Methylation abnormalities	15q11-13 region	28
Glasgow	PCR	Targeted screen	STRs	Microsatellite markers	28
Proforma required?		NO			

## REFERRAL CRITERIA

- Clinical features that include:
  - o Severe hypotonia and/or feeding difficulties in early infancy
  - o Global developmental delay
  - o 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 Dundee	QF- PCR	Targeted screen	STRs	Chromosomes 13, 18, 21, X/Y	28
Edinburgh 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

# SILVER-RUSSELL SYNDROME

#### **AVAILABLE TESTING**

Centre	Method	Scope a	nd range of test	Targets	TAT
Glasgow	MLPA	Targeted screen CNV Methylation abnormalitie		11p15 region	28
Glasgow	PCR	Targeted screen	Targeted screen STRs		28
Proforma required?		NO			

## REFERRAL CRITERIA

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

- Clinical Genetics
- Paediatrics

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

#### **AVAILABLE TESTING**

Centre	Method	Scope and ran	ge of test	Targets	TAT
Glasgow	Sanger	Whole gene screen	SNVs, indels	DHCR <sub>7</sub>	56
Family member testing			as ind	icated above	14
Proforma re	quired?	NO			

#### REFERRAL CRITERIA

- Clinical features that include:
  - o Prenatal and postnatal growth restriction
  - Microcephaly
  - o 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
  - o 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 Prenatal 3
Proforma red	quired?	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 test		Targets	TAT
Glasgow	PCR	Targeted screen	Targeted screen Methylation analysis		28
Proforma re	quired?	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	Who	le gene screen	SNVs, indels	GNAS	56
Family memb		as indicated above				
Proforma requir	red?	YES	Endocrine disor	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 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	SNVs, indels	AR	56
Family me	ember testing		as indicated abov	ve	14
Proforma 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 ra	nge of test	Targets	TAT
Dundee	Sanger	Whole gene screen	SNVs, indels	GCK	56
Family men	nber testing		as indicated ab	ove	14
Proforma required?		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 range		e of test	Targets	TAT
Dundee	NGS	Whol	e gene screen	SNVs, indels	PRKAR1A	56
Family member testing				as indica	ited above	14
Proforma requ	uired?	YES	Endocrine diso	rders proforma (s	ee 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)
  - o Myxoma (cutaneous and mucosal)
  - o 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
  - o 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
  - o Psammomatous melanotic schwannomas (PMS)
  - o Blue nevus, epithelioid blue nevus
  - o Breast ductal adenoma
  - Osteochondromyxoma

- Clinical Genetics
- Dermatology
- Endocrinology

#### **CONGENITAL HYPERINSULINISM**

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS	Whole gene screen	SNVs, indels exon level CNV (selected genes)	ABCC8, AKT2, CACNA1D, GCK, GLUD1, GPC3, HADH, HNF1A, HNF4A, INSR, KCNJ11, KDM6A, KMT2D, PMM2, SLC16A1, TRMT10A	112
Family me	Family member testing		as in	dicated above	14
Proforma required? NO		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
  - o Detectable serum insulin or c-peptide when the blood glucose is <3mmol/l, OR
  - 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	Sanger	Whole gene screen	SNVs, indels	NKX2-1 (TITF1)	56
Dundee	NGS	Whole gene screen	SNVs, indels	DUOX2, DUOXA2, FOXE1, GLIS3, GNAS, HESX1, IGSF1, IRS4, IYD, LHX3, LHX4 NKX2-1, OTX2, PAX8, POU1F1, PRKAR1A, PROP1, SECISBP2, SLC16A2, SLC26A4, SLC5A5, TBL1X, TG, THRB, THRA, TRHR, TPO, TSHR, TSHB	112
Family men	nber testing		as i	ndicated above	14
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 rar	nge of test	Targets	TAT
Dundee	Sanger	Whole gene screen	SNVs, indels	AQP2, AVPR2	56
Family men	nber testing		as ind	icated above	14
Proforma requ	uired?	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 range of test		Targets	TAT
Dundee	NGS	Whole gene screen	SNVs, indels	AKT2, BRWD3, CDKN1C, CHD8, DIS3L2, DNMT3A, EZH2, GPC3, MTOR, NFIB, NFIX, NSD1, OFD1, PDGFRB, PIK3CA, PTEN, RNF125, SETD2, SUZ12	112
Family mer	mber testing		as indicat	ed above	14
Proforma re	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	Whole ger	ne SNVs, indels	AP2S1, CASR, CDC73, CDKN1B, GCM2, GNA11, MEN1, RET (exons 5, 8, 10, 11, 13-16)	56
Family me	mber testing			as indicated above	14
Proforma re	quired?	YES I	Endocrine disorders p	roforma (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
  - o 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
  - o 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	Whole ge	Whole gene screen SNVs, inde		AP2S1, CASR, CDC73, CDKN1B, GCM2, GNA11, MEN1, RET (exons 5, 8, 10, 11, 13-16)	56
Family men	nber testing		as ind		icated above	14
Proforma re	quired?	YES	Endocrine	disorders proforma	(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			Targets	TAT
Dundee	NGS	Whole scre	•	SNVs, indels	AIRE, CASR, GATA3, GCM2, GNA11, PTH, TBCE	56
Family me	mber testing		as		indicated above	14
Proforma re	quired?	YES	Endocri	ne disorders prof	orma (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	Sanger	Whole ge	Whole gene screen SN		MEN1, CDKN1B, AIP	56
Family men	nber testing			as ind	icated above	14
Proforma re	quired?	YES	Endocrine	disorders proforma	(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
  - o 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	ember testing		as indicat	ed above	14
Proforma re	quired?	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	quired?	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	Whole ge	ne screen	SNVs, indels Exon level CNV* *Only available on request, not routinely offered	CDC <sub>73</sub>	56
Family men	nber testing			as indicated a	above	14
Proforma	required?	YES	Endocrine disorders proforma (see centre 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 range of test		Targets	TAT
Dundee	NGS	Whole gene screen	SNVs, indels	ALB, SECISBP2, SLC16A2, THRA, THRB, TSHR, TTR	112
Family men	nber testing		as indic	ated above	14
Proforma re	quired?	NO			

## REFERRAL CRITERIA

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

- Clinical Genetics
- Endocrinology

#### HYPOGONADOTROPHIC HYPOGONADISM

#### **AVAILABLE TESTING**

Centre	Method	9	Scope and rang	e of test	Targets	TAT
Glasgow	NGS	Whole	gene screen	SNVs, indels	ANOS1, CHD7, CUL4B, FEZF1, FGF8, FGFR1, FSHB, GNRH1, GNRHR, KISS1R, NRoB1, PROK2, PROK2R, SEMA3E, SOX2, SOX10, SPRY4, TAC3, TACR3, WDR11	112
Family me	mber testing			as indic	ated above	14
Proforma re	quired?	YES	DSD referral f	orm (see centre w	vebsite)	

#### REFERRAL CRITERIA

• Clinical history of Hypogonadism

#### **REQUESTING SPECIALTIES**

- Clinical Genetics
- Paediatrics

#### HYPOPHOSPHATEMIC RICKETS

#### **AVAILABLE TESTING**

Centre	Method	Scope and ran	ge of test	Targets	TAT
Dundee	NGS	Whole gene screen	SNVs, indels	CYP27B1, CYP2R1, DMP1, ENPP1, FAM2oC, FGF23, PHEX, SLC34A1, SLC34A3, VDR	112
Family me	mber 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	Whole ger screen	ne SNVs, indels	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
Family member testing		•	as indicated above	14	
Proforma re	equired?	YES	Monogenic diabete	s 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 MODY based on MODY calculator http://www.diabetesgenes.org/content/modyprobability-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.
  - o Post-pubertal children or adults with insulin resistance:
    - Severely elevated plasma insulin (typically greater than 150pmol/L in non-diabetic nonobese 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
  - o 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	Scope and range of test			Targets	TAT
Dundee	Sanger	Targete	d screen	SNVs, indels	MEN1, CDKN1B, AIP	56
Family men	nber testing			as ind	icated above	14
Proforma re	equired?	YES	Endocrine	disorders proforma	a (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:
  - o Parathyroid hyperplasia/multiglandular adenomas
  - Pituitary tumors
  - o Endocrine tumors of the gastro-entero-pancreatic (GEP) tract
  - Carcinoid tumors
  - o Adrenocortical tumors
- MEN1-related non-endocrine tumours include:
  - o facial angiofibromas
  - o collagenomas
  - o meningioma
- Overlapping clinical indications:
  - o Familial Hyperparathyroidism
  - o Familial Pituitary Adenoma (FIPA)

- Clinical Genetics
- Endocrinology

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

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Dundee	Sanger	Targete	d screen	SNVs, indels	RET (exons 5, 8, 10, 11, 13, 14, 15, 16)	56
Family men	nber testing			as ind	icated above	14
Proforma re	quired?	YES	Endocrine	disorders proforma	(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
  - o ≥2 MEN2-related endocrine abnormalities (any age), OR
  - ≥1 MEN2-related endocrine abnormality and a first degree relative with ≥1 MEN2related endocrine abnormality
- MEN2-related endocrine abnormalities include: Medullary Thyroid Carcinoma (MTC), Phaechromocytoma/paraganglioma, Parathyroid adenoma/hyperplasia, Hirschprungs disease
- Overlapping clinical indications:
  - o Phaeochromocytoma and paraganglioma panel

- Clinical Genetics
- Endocrinology

#### PHAEOCHROMOCYTOMA AND PARAGANGLIOMA

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Dundee	NGS	Whole ge screen		SNVs, indels Exon level CNV in relevant genes	SDHA, SDHB, SDHC; SDHD, SDHAF2, VHL, MAX, TMEM127, RET (exons 5, 8, 10, 11, 13 to 16), FH	56
Family men	nber testing			as	indicated above	14
Proforma re	quired?	YES	End	locrine disorders profor	ma (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 (<60 years), OR
  - o 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 *NF*1.
- Overlapping clinical indications:
  - o 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 rang	ge of test	Targets	TAT
Dundee	NGS	Whole gene screen	SNVs, indels	ARMC5, PDE11A, PDE8B, PRKAR1A	112
Family me	mber testing		as indica	ted 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 ran	ge of test	Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	KCNJ <sub>5</sub>	56
Family member testing			as ind	icated above	14
Proforma re	quired?	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	Whole gene screen SNVs, indels Exon level CNV		HNF1B	56
Family member testing			as indicated ab	pove	14
Proforma required?		NO			

#### REFERRAL CRITERIA

- Any individual with a clinical presentation consistent with the condition.
- Overlapping indication: Monogenic Diabetes. Full Monogenic diabetes panel will be added for all patients with diabetes with or without renal cysts unless requested otherwise.

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

## SEVERE EARLY ONSET OBESITY

#### **AVAILABLE TESTING**

Centre	Method		Scope and rang	ge of test	Targets	TAT
Dundee	NGS	Whole	gene screen	SNVs, indels	ALMS1, ARL6, BBS1, BBS10, BBS12, BBS2, BBS4, BBS5, BBS7, BBS9, CEP19, GNAS, LEP, LEPR, MC4R, MKKS, MKS1, MYT1L, NTRK2, PCSK1, PHF6, POMC, SDCCAG8, SIM1, TTC8, VPS13B	112
Family member testing		as indi			ated above	14
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	Whole ge	ne screen	SNVs, indels	THRB	56
Family men	nber testing			as ind	icated above	14
Proforma re	quired?	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	Whole ge	ne screen	SNVs, indels Exon level CNV	VHL	56
Family men	nber testing			as ind	icated above	14
Proforma re	quired?	YES	Endocrine	disorders proforma	a (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
  - o ≥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:
  - o Phaeochromocytoma and paraganglioma

- Clinical Genetics
- Endocrinology

# **EYES**

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

#### **AVAILABLE TESTING**

Centre	Method	Scope and rang	ge 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

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

# **ALBINISM & NYSTAGMUS**

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	AP3B1, CACNA1A, CACNA1F, CASK, FRMD7, GPR143, HPS1, HPS3, HPS4, HPS5, HPS6, LRMDA, LYST, OCA2, PAX6, RAB27A, SACS, SETX, SLC24A5, SLC38A8, SLC45A2, TYR, TYRP1	112
Family men	nber testing	•		as indicated above	14
Proforma re	quired?	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 screen	SNVs, indels, Exon level 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 indicated above			14
Proforma required?		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 gene screen	SNVs, indels, Exon level CNV	ARL6, BBS1, BBS2, BBS4, BBS5, BBS7, BBS9, BBS10, BBS12, LZTFL1, MKKS, MKS1, SDCCAG8, TMEM67,TTC8	112
Family member testing		as indicated above			14
Proforma required?		NO			

#### REFERRAL CRITERIA

- Clinical features suggestive of Bardet-Biedl Syndrome (≥4 primary features or 3 primary features & ≥2 secondary features
  - o 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 ra	inge of test	Targets	TAT
Aberdeen	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV	BEST1	56
Family member testing		as indicated above			14
Proforma required?		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

### **BRITTLE CORNEA SYNDROME**

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen SNVs, indels		PRDM5, ZNF469	56
Family member testing			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

- 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		СНМ	56
Family member testing			14		
Proforma required?		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

	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels, Exon level 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				as indicated above	14
testing 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 screen	SNVs, indels, Exon level CNV	ADAMTS18, ALDH18A1, B3GLCT, CHRDL1, CHST6, COL17A1, COL8A2, DCN, GJA1, GRHL2, GSN, HMX1, KERA, KRT12, KRT3, LCAT, MIR184, OVOL2, PIK3R1, PIKFYVE, PRDM5, PXDN, RAB18, RAB3GAP1, RAB3GAP2, SLC16A12, SLC4A11, STS, TACSTD2, TCF4, TGFBI, UBIAD1, ZEB1, ZNF469	112
Family men	nber testing		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 gene screen	SNVs, indels, Exon level CNV	CHST6, COL17A1, COL8A2, DCN, GRHL2, GSN, KERA, KRT12, KRT3, LCAT, MIR184, OVOL2, PIKFYVE, PRDM5, SLC4A11, STS, TACSTD2, TCF4, TGFBI, UBIAD1, ZEB1, ZNF469	112	
Family men	nber testing		as indicated above			
Proforma re	quired?	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 screen	SNVs, indels Exon level CNV	ADAMTS10, ADAMTS17, CYP1B1, FOXC1, FOXD3, FOXE3, JAG1, LTBP2, MYOC, PITX2	56
Family men	nber testing		as indicated above		
Proforma required? NO		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 rar	nge of test	Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	ATOH7, FZD4, LRP5, NDP, TSPAN12	56
Family men	Family member testing as indicate			d above	14
Proforma required?		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 gene screen	SNVs, indels, Exon level CNV	ADAMTS10, ADAMTS17, CPAMD8, CREBBP, CYP1B1, DDX58, FOXC1, FOXE3, IFIH1, LMX1B, LTBP2, MYOC, OCRL, PAX6, PITX2, SBF2, SH3PXD2B, TEK	112
Family men	Family member testing as indicated above		as indicated above	14	
Proforma required? NO					

#### REFERRAL CRITERIA

• Clinical features suggestive of a monogenic cause of Glaucoma

## **REQUESTING SPECIALTIES**

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

#### **MACULAR DYSTROPHY**

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger MLPA	Whole gene screen	SNVs, indels	PRPH2	56
Family men	nber testing		as inc	licated above	14
Proforma re	quired?	NO			

## REFERRAL CRITERIA

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

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

## **OCULAR MALFORMATIONS**

## **AVAILABLE TESTING**

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

## REFERRAL CRITERIA

- Microphthalmia or anophthalmia or uveoretinal coloboma where there is evidence to support a likely monogenic cause, for example bilateral disease, consanguinity or additional ocular and non-ocular features, OR
- 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 screen	SNVs, indels, Exon level CNV	GPR143, HPS1, HPS3, HPS4, HPS5, LRMDA, LYST, OCA2, SLC24A5, SLC45A2, TYR, TYRP1	112
Family men	nber testing		as indicated above		
Proforma re	Proforma required? NO				

## REFERRAL CRITERIA

• Clinical features suggestive of Oculocutaneous Albinism – very light skin and light coloured irises, decreased sharpness of vision, nystagmus, strabismum, photophobia

#### REQUESTING SPECIALTIES

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

#### OPTIC ATROPHY TYPE 1

#### **AVAILABLE TESTING**

Centre	Method	Scope and ra	nge of test	Targets	TAT
Aberdeen	NGS MLPA	Whole gene screen	SNVs, indels Exon level CNV	OPA1	56
Family men	member testing as indicated above			ited above	14
Proforma required?		NO			

#### REFERRAL CRITERIA

 Clinical features suggestive of Optic Atrophy type 1 – vision loss, difficulty distinguishing between colours, pallor of the optic nerve. May also have sensorineural hearing loss, ataxia, myopathy

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

## **OPTIC NEUROPATHY**

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT	
Aberdeen	NGS	Whole gene screen	SNVs, indels, Exon level CNV	ACO2, C12orf65, C19orf12, CISD2, DNM1L, MFF, MFN2, NR2F1, OPA1, OPA3, RTN4IP1, SLC25A46, SLC52A2, SPG7, SSBP1, TMEM126A, WFS1	112	
Family men	nber testing		as indicated above			
Proforma re	quired?	NO	as indicated above 14			

## REFERRAL CRITERIA

• Clinical features suggestive of an Optic Neuropathy

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

## **RETINAL DISORDERS**

#### **AVAILABLE TESTING**

Centre	Method	Scope and	range of test	Targets	TAT		
Aberdeen	NGS	Whole gene screen	SNVs, indels, Exon level 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 (C210rf2), 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, GPR143, 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 (c20rf71), 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		
Family member testing			as indicated above	14			
Family member testing			as indicated above 1.				

#### 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.

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

## **USHER SYNDROME**

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels, Exon level CNV	MYO7A, USH1C, CDH23, PCDH15, USH1G, ADGRV1, DFNB31 (WHRN), USH2A	56
Family men	nber testing		as indicated above		14
Proforma re	na required? 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 rang	e of test	Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	FRMD <sub>7</sub>	56
Family member testing as ind		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 men	nber testing	r testing as indica		ed above	14	
Proforma re	quired?	NO	10			

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 gene screen	SNVs, indels Exon level CNV	ABCB11, ABCB4, ABCC2, AKR1D1, ALDOB, AMACR, ATP8B1, BAAT, BCS1L, CLDN1, CYP27A1, CYP7A1, DCDC2, FAH, HSD3B7, JAG1, MYO5B, NOTCH2, NPC1, NPC2, NR1H4, PEX1, PEX12, PEX26, PEX6, SERPINA1, SLC25A13, TALDO1, TJP2, UGT1A1, VIPAS39, VPS33B	112
Family member testing			as indicated above	14	
Proforma re	quired?	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	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 r	ange 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

## HIRSCHSPRUNG DISEASE

## **AVAILABLE TESTING**

Centre	Method	Scope and ra	nge of test	Targets	TAT
Dundee	NGS	Whole gene screen	SNVs, indels	RET as a single gene, or 8 gene panel: EDN3, EDNRB, KIF1BP, L1CAM, PHOX2B, RET, SOX10,ZEB2	56 - RET 112 -panel
Family member testing		as indicate	d above	14	
Proforma required? NO					

## REFERRAL CRITERIA

- Diagnosis of Hirschsprung disease (HSCR) and at least one of the following:
  - o 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

- Clinical Genetics
- Gastroenterology
- Hepatology
- Paediatrics

## **PANCREATITIS**

## **AVAILABLE TESTING**

Centre	Method	Scope and rar	nge of test	Targets	TAT
Aberdeen	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV	SPINK1, PRSS1	56
Aberdeen Dundee Edinburgh Glasgow	ARMS	Targeted screen	SNVs, indels	CFTR common mutations	28
Family member testing			as indica	ted above	14
Proforma required?		NO			

## REFERRAL CRITERIA

- Recurrent acute pancreatitis
- Chronic pancreatitis
- 1<sup>st</sup> 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 Exon level CNV		ALAD, ALAS2, CPOX, FECH, HMBS, PPOX, UROD, UROS	56
Family member testing as		as indicat	ed above	14	
Proforma required? NO		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 ra	nge of test	Targets	TAT
Aberdeen	Sanger MLPA	Whole gene screen SNVs, indels Exon level CNV		ATP7B	56
Family men	Family member testing		as indicat	ed above	14
Proforma required? 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 MLPA	Whole ge	ne screen	SNVs, indels Exon level CNV	SERPINC1	56
Family men	nber testing			as indi	cated above	14
Proforma re	quired?	YES	Molecular	Haematology requ	est 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 ge	ne screen	SNVs	GP1BA, GP1BB, GP9	56
Family men	nber testing		as indicated above			14
Proforma re	quired?	YES	(ES Molecular Haematology request form (see centre website)			

## REFERRAL CRITERIA

• Platelet function testing suggestive of Bernard Soulier syndrome

- Clinical Genetics
- Haematology

## **COAGULATION & FIBRINOLYSIS PANEL**

## **AVAILABLE TESTING**

Centre	Method	Sc	ope and rar	nge of test	Targets	TAT
Edinburgh	NGS	Whole ge	ne screen	SNVs	F2, F5, F7, F8, F9, F10, F11, F12, F13A1, F13B, FGA, FGB, FGG, GGCX, KLKB1, KNG1, LMAN1, MCFD2, SERPINE1, SERPINF2, THBD, VKORC1, VWF	84
Family men	nber testing		as indicated above			14
Proforma re	quired?	YES	S Molecular Haematology request 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 a	nd range of test	Targets	TAT
Edinburgh	NGS	Whole gene screen		SNVs (plus exon level CNV for F8 where appropriate)	F5, F8, LMAN1, MCFD2	56
Family men	nember testing as indicated above			2	14	
Proforma required? YES Molecular			Molecular	Haematology request form (se	ee 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 screen SNVs, indels Exon level CNV		RPL5, RPS10, RPL11, RPL35A, RPS7, RPS19, RPS24, RPS26, GATA1, RPS17	56
Family men	Family member testing		as indi	cated above	14
Proforma required?		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 gene screen	SNVs, indels	EGLN1, EPAS1, EPO, EPOR, HBA1, HBA2, HBB, VHL	112
Family men	member testing			as indicated above	14
Proforma required? NO					

## REFERRAL CRITERIA

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

## **REQUESTING SPECIALTIES**

- Clinical Genetics
- Haematology

## **FACTOR VII DEFICIENCY**

#### **AVAILABLE TESTING**

Centre	Method		Scope and ra	ange of test	Targets	TAT
Edinburgh	Sanger MLPA	Whole o	gene screen	SNVs Exon level CNV	F7	56
Family men	nber testing	as indicated above				14
Proforma required? YES			Molecular Ha	ematology 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 range of test			Targets	TAT	
Edinburgh	Sanger MLPA	Whole	gene screen	SNVs Exon level CNV	F10	56	
Family me	ember testing	mber testing as indicated above				14	
Proforma required? YES Mole			Molecular Ha	Haematology 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 MLPA	Whole	e gene screen	SNVs Exon level CNV	F11	56
Family me	Family member testing			as indicated	above	14
Proforma required?		YES	Molecular Had	ematology request forn	n (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 ge	ne screen	SNVs	FGA, FGB, FGG	56
Family men	nber testing			as indi	cated above	14
Proforma re	quired?	YES	Molecular	Haematology request 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

## **GLANZMANN THROMBASTHENIA**

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Edinburgh	NGS	Whole ge	ne screen	SNVs	ITGA2B, ITGB3	56
Family men	nber testing		as indicated above			
Proforma re	quired?	YES	Molecular	cular Haematology request 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 Dundee Glasgow	ARMS (D & G) Sanger (A)	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	Whole gene screen	HBB, HBA	56	
Family member testing			as indicated ab	pove	14
Proforma required?		NO			

## REFERRAL CRITERIA

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

- Clinical Genetics
- Haematology

#### **HAEMOPHILIA A**

#### **AVAILABLE TESTING**

Centre	Method		Scope an	d range of test	Targets	TAT
Edinburgh	NGS MLPA Inversion PCR	Whole gene screen		SNVs Exon level CNV Inversions *	F8	56
Family me	Family member testing		as indicated above			
Proforma required?		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 r	ange of test	Targets	TAT
Edinburgh	Sanger MLPA	Wh	ole gene screen	SNVs, indels Exon level CNV	F9	56
Family me	Family member testing		as indicated above			
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 ra	nge of test	Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	BRCA2, BRIP1, CTC1, DKC1, ELANE, FANCA, FANCB, FANCC, FANCD2, FANCE, FANCF, FANCG, FANCI, FANCL, FANCM, G6PC3, GATA1, GATA2, GFI1, HAX1, MPL, NHP2, NOP10, PALB2*, RAD51C, RPL11, RPL35A, RPL5, RPS10, RPS17, RPS19, RPS24, RPS26, RPS7, RUNX1, SBDS, SLX4, SRP72, TERT, TINF2, WAS, WRAP53	112
Family men	nber testing		as indic	ated above	14
Proforma re	quired?	NO			

## REFERRAL CRITERIA

• Clinical phenotype suggestive of an inherited bone marrow failure disorder

- Clinical Genetics
- Haematology

#### **IRON REGULATION**

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	
Edinburgh	NGS	Whole gene screen	SNVs, indels	ABCB7, ALAS2, ATP7B, BMP6, CP, CYBRD1, FTL, GBA, GLRX5, HAMP, HFE, HFE2, SLC11A2, SLC25A38, SLC40A1, TF, TFR2, TMPRSS6	112
Family member testing			as i	ndicated above	14
Proforma re	quired?	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

## **MYELODYSPLASTIC SYNDROME**

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen SNVs, indels Exon level CNV		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		Scope and	range of test	Targets	TAT
Aberdeen	Sanger	Whole ge	ne screen	SNVs, indels	ELANE	56
Family mem	nber testing			as indicated	d above	14
Proforma re	quired?	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 gen screen	e SNVs, indels	ABCG5, ABCG8, ACTB, ACTN1, ANKRD26, ANO6, AP3B1, AP3D1, ARPC1B, BLOC1S3, BLOC1S6, CDC42, CYCS, DIAPH1, DTNBP1, ETV6, FERMT3, FLI1, FLNA, FYB1, GBA, GATA1, GFI1B, GNE, GP1BA, GP1BB, GP6, GP9, HOXA11, HPS1, HPS3, HPS4, HPS5, HPS6, ITGA2B, ITGB3, KDSR, LYST, MECOM, MPIG6B, MPL, MYH9, NBEA, NBEAL2, P2RY12, PLA2G4A, PLAU, RASGRP2, RBM8A, RNU4ATAC, RUNX1, SLFN14, SRC, STIM1, STXBP2, TBXA2R, TBXAS1, THPO, TUBB1, VIPAS39, VPS33B, VWF, WAS	84
Family member testing			as indicated above	14	
Proforma re	quired?	YES	Molecular Haema	atology 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 MLPA	Whole gene screen		SNVs, indels Exon level CNV	PROC	56
Family men	nber testing			as indicated a	bove	14
Proforma re	quired?	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 MLPA	Whole ge	ne screen	SNVs, indels Exon level CNV	PROS1	56
Family men	nber testing		as inc		ited above	14
Proforma re	quired?	YES	Molecular	Haematology reques	Haematology request form (see centre website)	

## REFERRAL CRITERIA

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

- Clinical Genetics
- Haematology

## **RED CELL ENZYMES**

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	NGS	Whole gene screen SNVs, indels		ALDOA, AK1, G6PD, GCLC, GPI, GSR, GSS, HK1, NT5C3A, PFKM, PKLR, TPI1, CYB5R3	112
Family member testing		as i	ndicated above	14	
Proforma required?		NO			

#### REFERRAL CRITERIA

• Clinical presentation or biochemical enzyme deficiency highly suggestive of a specific red cell enzyme deficiency

## **REQUESTING SPECIALTIES**

- Clinical Genetics
- Haematology

## **RED CELL MEMBRANOPATHY**

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	NGS	Whole gene screen SNVs, indels		AK1, ANK1, EPB41, EPB42, RhAG, SLC2A1, SLC4A1, SPTA1, SPTB, XK	112
Family me	mber testing		as inc	licated above	14
Proforma re	quired?	NO			

#### REFERRAL CRITERIA

• Phenotypic presentation highly suggestive of a gene included in the current panel.

- Clinical Genetics
- Haematology

## SCHWACHMAN-DIAMOND SYNDROME

## AVAILABLE TESTING

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole gene screen		SNVs, indels Exon level CNV	SBDS, DNAJC21	56
Family me	mber testing			as indicated	above	14
Proforma re	quired?	YES	GEN FORM	215 Primary Immunod	deficiency Request form (see centre we	bsite)

## 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	<i>HBB</i> p.(Glu⁊Val)	28 Prenatal 3
Edinburgh	Sanger	Targeted screen	SNVs	HBB p.(Glu7Val)	28 Prenatal 3
Glasgow	Sanger	Targeted screen (incl. newborn screening)	SNVs	<i>HBB</i> p.(Glu7Val)	28 Prenatal 3 Newborn screening 7
Proforma required?		NO	•		

## **REFERRAL CRITERIA**

- Sickle cell anaemia diagnosed by Haemaology 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 range	e of test	Targets	TAT
Aberdeen Dundee EdinburghMP*	Sanger (A) ARMS (D) Real time PCR (E)		Targeted screen	SNVs	F5 p.R534Q F2 c.*97G>A	28
Proforma require	Proforma required?					

<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 gene screen		SNVs	ADAMTS13, F2, F5, HRG, PIGA, PLG, PROC, PROS1, SERPINC1, SERPIND1, THBD	84
Family member testing			as indicated above			
Proforma required?		YES	Molecular Haematology request 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 rar	nge of test	Targets	TAT
Edinburgh	NGS MLPA	Who	ole gene screen	SNVs Exon level CNV	VWF	56
Family member testing		as indicated above				14
Proforma required?		YES Molecular Haematology request form (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

# **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:
  - o Individuals in whom aminoglycoside therapy may be required
  - o 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

## **BRANCHIOOTORENAL (BOR) SYNDROME**

## **AVAILABLE TESTING**

Centre	Method	Scope and	range of test	Targets	TAT	
Dundee	NGS	Whole gene screen	SNVs, indels, Exon level CNV ( <i>EYA</i> 1)	EYA1, SIX1, SIX5	56	
Family member testing		as indicated above				
Proforma required?		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	Whole gene screen	SNVs, 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 member testing				as indicated above	14
Proforma required?		NO			

## 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	Whole gene screen SNVs, indels (GJB2) Deletions (GJB6)		GJB2, GJB6	28
Family men	nber testing		14		
Proforma required?		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 rar	nge of test	Targets	TAT
Dundee	Sanger	Whole gene screen	SNVs, indels	SLC26A4, FOXI1	56
Family member testing			as indic	ated above	14
Proforma re	quired?	NO			

# REFERRAL CRITERIA

- Any individual with a clinical presentation consistent with the condition:
  - o Autosomal recessive deafness also associated with thyroid goiter
  - o 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	Whole gene screen	SNVs, indels	ADGRV1, CDH23, CIB2, CLRN1, MYO7A, PCDH15, PDZD7, USH1C, USH1G, USH2A, WHRN	112
Family me	mber testing			as indicated above	14
Proforma re	quired?	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	Whole gene screen	SNVs, indels	PAX3, MITF, SOX10, SNAI2, EDNRB, EDN3, KIT	112
Family me	member testing			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	Whole gene screen SNVs, indels		WFS1	56
Family me	mber testing		as inc	icated above	14
Proforma required?		NO			

### REFERRAL CRITERIA

- Any individual with a clinical presentation consistent with the condition.
- Available as single gene analysis via targeted Monogenic Diabetes NGS panel

- Clinical Genetics
- Endocrinology

# **IMMUNOLOGY**

# ADENOSINE DEAMINASE DEFICIENCY (ADAD)

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	Sanger MLPA	Whole ge	ne screen	SNVs, indels Exon level CNV	ADA2 (CECR1)	56
Family mem	ber testing			as indicated a	above	14
Proforma required? YES			GEN FORI	M 215 Primary Immunodef	iciency Request form (see centre website)	

### 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Immunology

# ANHYDROTIC ECTODERMODYSPLASIA WITH ID

### **AVAILABLE TESTING**

Centre	Method	!	Scope and i	ange of test	Targets	TAT
Aberdeen	NGS	Whole ge	ne screen	SNVs, indels Exon level CNV	IKBKG (NEMO), NFKBIA (IKBA)	56
Family men	nber testing			as indicate	ed above	14
Proforma re	equired? YES GEN FO		GEN FOR	M 215 Primary Immunod	eficiency Request form (see centre website)	

### 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 2017 (J Clin Immunol., 2018 38:129–143).

- 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 member testing			as indi	cated above	14
Proforma re	quired?	NO			

### REFERRAL CRITERIA

- 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 generation	e SNVs, indels Exon level CNV	ADAM17, AICDA, AP3B1, B2M, BTK, CBL, CD4oLG, CORO1A, CTC1, CTPS1, CYBA, CYBB, DCLRE1C, DOCK8, FERMT1, FOXP3, GUCY2C, HPS1, HPS4, HPS6, ICOS, IFNGR1, IFNGR2, IKBKG, IL1oRA, IL2RA, ITGB2, MAGT1, NCF1, NCF2, NCF4, NF1, PIK3CD, PIK3R1, PTEN, PYCARD, SKIV2L, SLC37A4, STK4, TTC37, VPS13B, WAS	112
Family men	nber testing			as indicated above	14
Proforma re	quired?	YES	GEN FORM 215 Prima	ary Immunodeficiency Request form (see centre website)	

### REFERRAL CRITERIA

• Please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Rheumatology

### **AUTOINFLAMMATORY DISORDERS**

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gen screen	e SNVs, indels Exon level CNV	ACP5, ADA2 (CECR1), ADAM17, ADAR1, AP1S3, CARD14, COPA, IFIH1, IL1RN, IL36RN, LPIN2, MEFV, MVK, NOD2, NLCR4, NLRP1, NLRP3, NLRP12, OTULIN, PLCG2, POLA1, PSMB8, PSTPIP1, RNASEH2A, RNASEH2B, RNASEH2C, SAMHD1, SH3BP2, SLC29A3, TMEM173, TNF1IP3, TNFAIP3, TNFRSF1A, TREX1, USP18	112
Family men	nber testing		•	as indicated above	14
Proforma re	equired?	YES	GEN FORM 215 Prima	ary Immunodeficiency Request form (see centre website)	

### 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Immunology
- Rheumatology

### **BACTERIAL AND PARASITIC INFECTIONS**

### **AVAILABLE TESTING**

Centre	Method	Scope a	nd range of test	Targets	TAT
Aberdeen	NGS	Whole gene SNVs, indels screen Exon level CNV		ACT1 (TRAF3IP1), APOL1, CARD9, HMOX1, IRAK1, IRAK4, MYD88, NBAS, NCSTN, PSEN, PSENEN, RANBP2, RPSA, STAT1, IL17F, IL17RA, 1L17RC, TIRAP	112
Family men	nber testing			as indicated above	14
Proforma re	quired?	YES	GEN FORM 215 Prima	ary Immunodeficiency Request form (see centre website)	

#### 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# BACTERIAL INFECTIONS, AUTOINFLAMMATION, AMYLOPECTINOSIS

### **AVAILABLE TESTING**

Centre	Method	9	Scope and r	ange of test	Targets	TAT
Aberdeen	NGS	Whole ge	ne screen	SNVs, indels Exon level CNV	HOIL1 (RBCK1), HOIP1 (RNF31)	56
Family men	nber testing			as indicat	ed above	14
Proforma required? YES GEN FORM 215 Primary Immun			deficiency Request form (see centre website)			

# 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# **CALCIUM CHANNEL DEFECTS**

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole ger	ne screen	SNVs, indels Exon level CNV	ORAI1, STIM1	56
Family men	nber testing			as indicated ab	ove	14
Proforma re	quired?	YES	GEN FOR	GEN FORM 215 Primary Immunodeficiency Request form (see centre website)		

### REFERRAL CRITERIA

- Autoimmunity
- EDA
- Non-progressive myopathy
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2017 (J Clin Immunol., 2018 38:129–143).

- 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 Exon level CNV	CYBA, CYBB, NCF1, NCF2, NCF4	56
Family mem	ber testing			as indicated	above	14
Proforma re	quired?	YES	GEN FORM 21	RM 215 Primary Immunodeficiency Request form (see centre website)		

### 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Immunology

# **COMBINED IMMUNODEFICIENCIES (CVID)**

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT	
Aberdeen	NGS	Whole gene	e SNVs, indels Exon level CNV	B2M, BCL11B, CD3G, CD8A, CIITA, DOCK8, IL21,LAT, LCK, MAGT1, MAP3K14, MSN, OX40 (TNFRSF4), RFX5, RFXANK, RFXAP,RHOH, STK4, TAP1, TAP2, TAPBP, TRAC, UNC119, ZAP70	56	
Family men	nber testing			as indicated above		
Proforma re	quired?	YES	GEN FORM 215 Prima	ary Immunodeficiency Request form (see centre website)		

# REFERRAL CRITERIA

- Generally less profound than SCID.
- For specific CVID subpanels (Low CD4 / MHCII expression absent, Low CD4 / MHCII expression present, Low CD8, Low B cells, Ig often normal, Ig low, Normal Ig with poor specific antibody response), please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

### **COMPLEMENT DEFICIENCIES**

### **AVAILABLE TESTING**

Centre	Method	Scope a	and range of test	Targets	TAT
Aberdeen	NGS	Whole gen	e SNVs, indels Exon level CNV	C1QA, C1QB, C1QC, C1R, C1S, C2, C3, C4A, C4B, C5, C6, C7, C8A, C8B, C8G, C9, CD55, CD59, CFB, CFD, FCN3, MASP2, PFC (CFP), SERPING1	56
Family men	nber testing			as indicated above	14
Proforma re	quired?	YES	GEN FORM 215 Primar	y Immunodeficiency Request form (see centre website)	

### 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Immunology
- Rheumatology

### **CONGENITAL THROMBOCYTOPENIA**

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole ge	ne screen	SNVs, indels Exon level CNV	ARPC1B, WAS, WIPF1	56
Family men	nber testing			as ind	icated above	14
Proforma re	quired?	YES GEN F		M 215 Primary Immunodeficiency Request form (see centre website)		

### 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 2017 (J Clin Immunol., 2018 38:129–143).

- 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 ge	ne screen	SNVs, indels Exon level CNV	MTHFD1, SLC46A1, TCN2	56 or 112
Family men	nber testing			as ind	icated above	14
Proforma re	quired?	YES	YES GEN FORM 215 Primary		unodeficiency Request form (see centre website)	

### REFERRAL CRITERIA

- Megablastic anaemia.
- Ig decreased.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on website and IUIS 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

### DNA REPAIR DEFECTS

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole gene screen		SNVs, indels Exon level CNV	ATM, BLM, CDCA7, DNMT3B, GINS1, HELLS, LIG1, MCM4, NBS1 (NBN), PMS2, POLE1, POLE2, NSMCE3, ERCC6L2, RNF168, ZBTB24	56
Family men	nber testing		as inc		icated above	14
Proforma required? YES GEN		GEN FORI	ORM 215 Primary Immunodeficiency Request form (see centre website)			

# 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# DYSKERATOSIS CONGENITA (DKC)

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT	
Aberdeen	NGS	Whole gen screen	SNVs, indels Exon level CNV	CTC1, DKC1,PARN, NOLA2 (NHP2), NOLA3 (NOP10), RTEL1, SAMD9, SAMD9L, SNM1B / APOLLO (DCLRE1B), STN1, TERC, TERT, TINF2, TPP1, WRAP53	112	
Family men	nber testing			as indicated above		
Proforma re	quired?	YES	GEN FORM 215 Prima	nary Immunodeficiency Request form (see centre website)		

### 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# FAMILIAL HLH DUE TO PRF1 VARIANTS

### **AVAILABLE TESTING**

Centre	Method	Scope and	d range of t	est	Targets	TAT
Aberdeen	Sanger MLPA	Whole ge	ne screen	SNVs, indels Exon level CNV	PRF1	56
Family mem	nber testing			as indicated a	above	14
Proforma re	quired?	YES	GEN FOR	M 215 Primary Immunodef	iciency Request form (see centre website)	

### 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 gene	e SNVs, indels Exon level CNV	AP3B1, AP3D1, CD27, CD70, CTPS1, DNASE2, FAAP24, ITK, LYST, MAGT1, PRF1, PRKCD, RAB27A, RASGRP1, RLTPR (CARMIL2), SH2DIA, SLC29A3, STX11, STXBP2, UNC13D, XIAP	56	
Family men	nber testing			as indicated above		
Proforma re	quired?	YES	GEN FORM 215 Prima	ry Immunodeficiency Request form (see centre website)		

### 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology
- Rheumatology

# HENNEKAM-LYMPHANGIECTASIA-LYMPHEDEMA SYNDROME

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole ger	ne screen	SNVs, indels Exon level CNV	CCBE1	56
Family men	nber testing			as indicate	d above	14
Proforma re	quired?	YES	YES GEN FORM 215 Primary Immuno		eficiency Request form (see centre website)	

### 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 2017 (J Clin Immunol., 2018 38:129–143).

- 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 ge	ne screen	SNVs, indels Exon level CNV	SP110	56
Family men	nber testing		as indicated above			14
Proforma re	quired?	YES	GEN FOR	GEN FORM 215 Primary Immunodeficiency Request form (see centre website)		

### 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# **HEREDITARY AMYLOIDOSIS**

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole gene screen		SNVs, indels Exon level CNV	APOA1, APOA2, APOA4, APOC2, APOC3, APOE, FGA, GSN, IL31RA, LYZ, TTR, UNC13D	112
Family men	nber testing		as indicated above			14
Proforma required? YES G		GEN FORI	M 215 Primary Immi	unodeficiency Request form (see centre website)		

### 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Rheumatology

# HEREDITARY ANGIOEDEMA, TYPES I & II

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	Sanger MLPA	Whole gene screen		SNVs, indels Exon level CNV	SERPING1	56
Aberdeen	NGS	Whole ge	ne screen	SNVs, indels Exon level CNV	SERPING1, Factor XII, PLG, ANGPT1	56
Family men	nber testing		as indicated above			
Proforma re	quired?	YES	GEN FORI	M 215 Primary Immunod	eficiency Request form (see centre website)	

### REFERRAL CRITERIA

- Hereditary angioedema
- Spontaneous activation of the complement pathway with consumption of C<sub>4</sub>/C<sub>2</sub>.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Immunology

# **HYPER IGE SYNDROMES (HIES)**

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole ge	ne screen	SNVs, indels Exon level CNV	PGM3, SPINK5, STAT3	56
Family men	nber testing		as indicated above			
Proforma re	quired?	? YES GEN FO		1 215 Primary Immunodeficiency Request form (see centre website)		

### 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

### **HYPOGAMMAGLOBULINAEMIA**

### **AVAILABLE TESTING**

Centre	Method	Scope a	and range of test	Targets	TAT	
Aberdeen	NGS	Whole gend	e SNVs, indels Exon level CNV	ATP6AP1, BLNK, BTK, CD19, CD20 (MS4A1), CD79A, CD79B, CD81, IGHM, IGLL1, IKZF1 (IKAROS), IRF2BP2, MOGS, NFKB1, PIK3CD, PIK3R1, PTEN, TCF3, TNFRSF13B (TACI), TNFRSF13C (BAFFR), TRNT1, TTC37, TWEAK (TNFSF12)	56 or 112	
Family men	nber testing			as indicated above		
Proforma re	quired?	YES	GEN FORM 215 Prima	ary Immunodeficiency Request form (see centre website)		

### REFERRAL CRITERIA

- IgG, IgA and / or IgM decreased
- Exclude second causes: drugs [Hx], myeloma [bone marrow], lymphoma. Ig loss (not hypo-IgM) 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# **ID WITH MULTIPLE INTESTINAL ATRESIAS**

### **AVAILABLE TESTING**

Centre	Method		Scope and	range of test	Targets	TAT
Aberdeen	NGS	Whole ge	ne screen	SNVs, indels Exon level CNV	TTC7A	56
Family men	nber testing		as indicated above			
Proforma required? YES		GEN FORI	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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

### IMMUNO-OSSEOUS DYSPLASIAS

### **AVAILABLE TESTING**

Centre	Method	Scope a	and range of test	Targets	TAT
Aberdeen	NGS	Whole gene	SNVs, indels Exon level CNV	EXTL3, MYSM1, RMRP, RNU4ATAC, SMARCAL1	56
Family men	nber testing		as indicated above		
Proforma re	quired?	YES	GEN FORM 215 Primar	y 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Immunology

# INTERFERONOPATHY / SLS / AGS / COMPLEMENT

### **AVAILABLE TESTING**

Centre	Method	Scope and i	range of test	Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	ACP5, ADAM17, C1QA, C1QB, C1QC, C1R, C2, C3, C5, C6, C7, C8A, C8B, C9, CFH, CFHR5, CFI, CFP, DNASE1, DNASE1L3, IFIH1, IRF8, RASGRP1, RNASEH2A, RNASEH2B, RNASEH2C, SAMHD1, SNORD118, TREX1, USP18	112
Family mem	ber testing		as indicated above		
Proforma re	quired?	YES	GEN FORM 215 Prima	ary Immunodeficiency Request form (see centre website)	

### 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 ge	ne screen	SNVs, indels Exon level CNV	KDM6A, KMT2D (MLL2)	56	
Family men	nber testing		as indicated above				
Proforma required? YES			GEN FORI	M 215 Primary Immunod	eficiency Request form (see centre website)		

### REFERRAL CRITERIA

- Typical facial abnormalies
- Cleft or high arched palate
- Skeletal abnormalities
- Short stature
- Intellectual disability
- Congenital heart defects
- Recurrent infecons (otitis media, pneumonia) in 50% of patients
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

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

#### AVAILABLE TESTING

Centre	Method		Scope and r	ange of test	Targets	TAT		
Aberdeen	NGS	Whole	e gene screen	SNVs, indels Exon level CNV	CXCR4 (WHIM), CYBB, FCGR3A, IFIH1, IFNAR2, IFNGR1, IFNGR2, IL12B, IL12RB1, IRF3, IRF7, IRF8, ISG15, JAK1, RORC, STAT1, STAT2, TBK1, TICAM1 (TRIF), TLR3, TMC6, TMC8, TRAF3, TYK2, UNC93B1	56		
Family men	nber testing		as indicated above					
Proforma re	quired?	YES	GEN FORM 21	GEN FORM 215 Primary Immunodeficiency 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 suscebtibility 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# **MISCELLANEOUS AUTOINFLAMMATORY CONDITIONS**

# **AVAILABLE TESTING**

Centre	Method	Scope ar	nd range of test	Targets	TAT	
Aberdeen	NGS	Whole general screen	e SNVs, indels Exon level 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 mem	ber testing			as indicated above		
Proforma re	quired?	YES	GEN FORM 215 Pri	mary Immunodeficiency Request form (see centre website)		

# 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 ge screen		SNVs, indels Exon level CNV	C16ORF57 (USB1), CLPB, COH1 (VPS13B), CSF3R, DNAJC21, ELANE, G6PC3, G6PT1 (SLC37A4), GFl1, HAX1, HYOU1, JAGN1, LAMTOR2, MKL1 (MRTFA), SBDS, SMARCD2, TAZ, VPS45, WAS, WDR1	56
Family men	nber testing		as indicated above			14
Proforma re	quired?	YES	GEN	I FORM 215 Primar	y 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

### OTHER ANTIBODY DEFICIENCIES

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT	
Aberdeen	NGS	Whole ge	ne screen	SNVs, indels Exon level CNV	AICDA, CARD11, IGKC, INO80, MSH6, UNG	56	
Family men	nber testing			as indicated above			
Proforma required? YES		GEN FORI	M 215 Primary Immui	nodeficiency Request form (see centre website)			

# 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# PURINE NUCLEOSIDE PHOSPHORYLASE DEFICIENCY

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole gene screen		SNVs, indels Exon level CNV	PNP	56
Family member testing		as indicated above				14
Proforma required?		YES	GEN FORM 215 Primary Immunodeficiency Request form (see centre website)			

### REFERRAL CRITERIA

- Autoimmune hemolytic anaemia
- Neurological impairment.
- Hypouricemia.
- Please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# **SEVERE COMBINED IMMUNODEFICIENCY (SCID)**

# **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole ge screen		SNVs, indels Exon level CNV	ADA, AK2, CD3D, CD3E, CD247, CORO1A, DCLRE1C (ARTEMIS), FOXN1, IL2RG, IL7R, JAK3, LIG4, NHEJ1, PRKDC, PTPRC, RAG1, RAG2	56
Family men	nber testing				as indicated above	14
Proforma re	quired?	YES	GEN	FORM 215 Primar	y Immunodeficiency Request form (see centre website)	

# REFERRAL CRITERIA

- CD<sub>3</sub> T cell lymphopenia: CD<sub>3</sub>+ 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# STAT<sub>5</sub>B DEFICIENCY

# **AVAILABLE TESTING**

Centre	Method		Scope an	d range of test	Targets	TAT
Aberdeen	NGS	Whole gen	ne screen	SNVs, indels Exon level CNV	STAT <sub>5</sub> B	56
Family men	nber testing as indicated above			ove	14	
Proforma required? YES		YES	GEN FORI	M 215 Primary Immunodefici	ency Request form (see centre website)	

# 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# STROKE

# **AVAILABLE TESTING**

Centre	Method		Scope and ran	ge of test	Targets	TAT
Aberdeen	NGS	Whole o	gene screen	SNVs, indels Exon level CNV	CBS, CST3, GLA, HTRA1, NOTCH3, ADA2	56
Family mem	ber testing			as indicated	d above	14
Proforma required? YES			GEN FORM 2:	15 Primary Immunod	eficiency Request form (see centre website)	

# 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 gen screen	e SNVs, indels Exon level CNV	AIRE, BACH2, CASP8, CASP10, CTLA4, FADD, FOXP3 (IPEX), IL2RA, IL10, IL10RA, IL10RB, ITCH, JAK1, LRBA, NFAT5, PEPD, STAT3, TNFRSF6 (FAS), TNFSF6 (FASLG), TPP2, ZAP70	56
Family men	nber testing		as indicated above		
Proforma re	quired?	YES	GEN FORM 215 Primary	y Immunodeficiency Request form (see centre website)	

# 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# SYNDROMES ASSOCIATED WITH CONGENITAL DEFECTS OF PHAGOCYTES

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gei screen	ne SNVs, indels Exon level CNV	ACTB, CEBPE, CSFR2A, CSFR2B, CTSC, FERMT3 (LADIII), FPR1, GATA2, G6PD, ILGB2 (LAD1), RAX2, SLC35C1 (LADII)	56
Family men	nber testing		as indicated above		
Proforma re	quired?	YES	GEN FORM 215 Primar	y Immunodeficiency Request form (see centre website)	

# 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# THYMIC DEFECTS WITH CONGENITAL ANOMALIES

# **AVAILABLE TESTING**

Centre	Method	9	Scope and r	ange of test	Targets	TAT
Aberdeen	NGS	Whole ge	ne screen	SNVs, indels Exon level CNV	CDH7, SEMA3E, TBX1	56
Family men	nber testing			as indicat	ed above	14
Proforma required? YES		YES	GEN FOR	M 215 Primary Immuno	deficiency Request form (see centre website)	

# 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, Charge syndrome), please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# **VASCULOPATHY**

# **AVAILABLE TESTING**

Centre	Method	Scope a	nd range of test	Targets	TAT
Aberdeen	NGS	Whole gene screen	e SNVs, indels Exon level CNV	ACTA2, BMPR2, COL3A1, COL4A1, COL5A1, COL5A2, EFEMP2, ELN, FBN1, FBN2, FOXE3, GUCA1B, LMNA, LOX, MFAP5, MYH11, MYLK, NOTCH1, PLOD1, PRKG1, RHOD, RNF213, SKI, SLC2A10, SMAD2, SMAD3, SMAD4, STX11, STXBP2, TGFB2, TGFB3, TGFBI, TGFBR1, YY1AP1	56
Family mem	ber testing			as indicated above	14
Proforma re	quired?	YES	GEN FORM 215 Prima	ary 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 a	nd range of test	Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	ADAM17, AICDA, CD4oLG, BTK, CD3G, ZAP7o, WAS, CYBA, CYBB, NCF1, NCF2, NCF4, DOCK8, EPCAM (Sanger only), FOXP3, GUCY2C, HPS1, HPS4, HPS6, ADA, IL2RG, LIG4, DCLRE1C, RAG2, IL1o, Il1oRA, IL1oRB, ITGB2, LRBA, ICOS, PIK3R1, PLCG2, RET, SH2D1A, XIAP, SKIV2L, TTC37, SLC37A4, STAT1, STAT3, STXBP2	112
Family mem	nber testing			as indicated above	14
Proforma re	quired?	YES	GEN FORM 215 Prima	ary Immunodeficiency Request form (see centre website)	

# 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	!	Scope and	range of test	Targets	TAT
Aberdeen	NGS	Whole ger	ne screen	SNVs, indels Exon level CNV	EPG <sub>5</sub>	56
Family men	nber testing			as indicated	l above	14
Proforma required? YES			GEN FOR	M 215 Primary Immunode	eficiency Request form (see centre website)	

# 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Haematology
- Immunology

# X-LINKED CGD

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	Sanger MLPA	Whole g	ene screen	SNVs, indels Exon level CNV	СҮВВ	56
Family men	nber testing			as indicated	d above	14
Proforma required? YES		GEN FORM	215 Primary Immunodo	eficiency Request form (see centre website)		

### 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 2017 (J Clin Immunol., 2018 38:129–143).

- Clinical Genetics
- Immunology

# **INHERITED CANCER**

# **COLORECTAL CANCER**

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Edinburgh	NGS	Whole ge screen		SNVs, indels	APC, BMPR1A, MBD4, MLH1, MSH2, MSH3, MSH6, MUTYH, NTHL1, PMS2, POLD1 (exons 4-12), POLE (exons 3-13), PTEN, RNF43, SMAD4, STK11	56
Family mem	nber testing			as indicated above 14		
Proforma re	quired?	YES	Colc	Colorectal cancer gene panel proforma (see centre website)		

### REFERRAL CRITERIA

- Living affected individual (proband) with colorectal cancer with:
  - o a. Diagnosed aged <30 OR
  - o b. Personal/family history of colorectal cancers reaching Amsterdam Criteria ( $\geqslant_3$  cases over  $\geqslant_2$
- 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.
- The proband's cancer and majority of reported cancers in the family should have been confirmed

### REQUESTING SPECIALTIES

# COWDEN SYNDROME / PTEN HAMARTOMA TUMOUR SYNDROME (PHTS)

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen Glasgow	NGS	Whole gene screen		SNVs, indels Exon level CNV	PTEN	56
Family member testing		as ind		as indicated abov	/e	14
Proforma required?		NO				

### REFERRAL CRITERIA

- Proband and / or family history meets one of the following criteria:
  - o 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
  - o 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
  - o ≥ 4 minor criteria
  - $\circ$  ≥ 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	NGS	Whole gene screen	SNVs, indels	DICER1	56
Family member testing			as indicate	d above	14
Proforma re	quired?	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; Childhoodonset 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 tumor; Multinodular goiter or differentiated thyroid cancer; Embryonal rhabdomyosarcoma other than thoracic or gynaecologic; Poorly differentiated neuroendocrine tumour; Undifferentiated sarcoma; Macrocephaly

#### REQUESTING SPECIALTIES

# **FAMILIAL MELANOMA**

### **AVAILABLE TESTING**

Centre	Method		Scope and	range of test	Targets	TAT
Glasgow	NGS	Whole ge	ne screen	SNVs, indels	BAP1, BRCA2, CDKN2A, CDK4, POT1	56
Family men	nber testing			as indicated	above	14
Proforma re	quired?	YES	Inherited o	cancer proforma (see cen	tre 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 melanomas in situ age < 30 OR
  - Melanoma in situ AND >/= 2 relatives (first / second / third degree) with melanoma in situ OR
  - Melanoma in situ AND >/= 1 first degree relative with melanoma in situ; one individual has multiple melanomas in situ OR
  - 1 Melanoma in situ OR melanoma in situ 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 in situ

- Clinical Genetics
- Oncology

# GORLIN SYNDROME (BASAL CELL NEVUS SYNDROME)

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	NGS	Whole gene screen	SNVs, indels	PTCH1, SUFU	56
Family member testing			as indica	ted above	14
Proforma re	quired?	NO			

#### REFERRAL CRITERIA

- Living individual affected (proband) where the individual history meets:
  - o ≥ 2 major and ≥ 1 minor criteria OR
  - ≥ 1 major and ≥ 3 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
  - o Palmar/plantar pits (two or more)
  - o SHH medulloblastoma, confirmed on tumour testing
  - o 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])
  - o 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	nd range of test	Targets	TAT
Aberdeen Glasgow	NGS	Whole gene	SNVs, indels Exon level CNV	BRCA1, BRCA2, TP53, PTEN, PALB2, STK11, CHEK2, ATM, RAD51C, RAD51D	56
Family member testing		•	as indicated above	14	
Proforma required? YES Glasgow laboratory o			Glasgow laboratory o	nly (see centre website)	

# REFERRAL CRITERIA

- Breast Cancer diagnosed <40 years
- Bilateral breast cancer, both <60 years
- Triple negative breast cancer, <60 years
- Male breast cancer, any age
- Two 1<sup>st</sup> degree relatives both diagnosed before the age of 45 years
- Manchester score of 15 / CanRisk score of 10%

- Breast Surgeons
- Clinical Genetics
- Oncology

# **HEREDITARY BREAST / OVARIAN CANCER SYNDROME**

# **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen Glasgow	NGS	Whole gene screen	SNVs, indels Exon level CNV	BRCA1, BRCA2, TP53, PTEN, PALB2, STK11, RAD51C, RAD51D, BRIP, MSH2, MSH6, MLH1, CHEK2, ATM	56
Family men	nber testing			as indicated above	14
Proforma required? YES Glasgow laboratory or		lasgow laboratory o	nly (see centre website)		

# REFERRAL 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
- Manchester score of 15 / CanRisk score of 10%

- Clinical Genetics
- Oncology

# HEREDITARY OVARIAN CANCER SYNDROME

# **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen Glasgow	NGS	Whole gene screen	SNVs, indels Exon level CNV	BRCA1, BRCA2, RAD51C, RAD51D, BRIP, MSH2, MSH6, MLH1, PALB2	56
Family mem	ber testing		•	as indicated above	14
Proforma required? YES Glasgow laboratory o			alasgow laboratory o	nly (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

### HEREDITARY PROSTATE CANCER

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		I range of test	Targets	ТАТ
Aberdeen	NGS <i>HOXB13</i> Sanger		e gene reen	SNVs, indels Exon level CNV	*BRCA1, BRCA2, ATM, HOXB13  **BRCA1, BRCA2, CHEK2, ATM, TP53, MLH1, MSH2, MSH6, RAD51D, PMS2, EPCAM, HOXB13.  PALB2 included where there is a family history of breast cancer	56
Family member testing			as indicated above	14		
Proforma required? YES   Prostate cancer proforma (see centre webs		(see centre website)				

# 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 MLPA	Whole gene screen		SNVs, indels Exon level CNV	SMAD4, BMPR1A	56	
Family member testing				as indicated a	bove	14	
Proforma required?		YES Co	ES Colorectal cancer gene panel proforma (see centre website)				

# REFERRAL CRITERIA

- Juvenile polyposis syndrome:
  - o a.  $\geq$  5 juvenile polyps of the colorectum, OR
  - $\circ$  b.  $\geqslant$  2 juvenile polyps throughout the GI tract, OR
  - o 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 Glasgow	NGS	Whole	gene screen	SNVs, indels Exon level CNV  TP <sub>53</sub>		56
Family mem	Family member testing			as indicated above		14
Proforma required?		NO				

#### REFERRAL CRITERIA

- Proband and / or family history meets one of the following criteria:
  - o Rhabdomyosarcoma (≤5 years)
  - Adrenocortical cancer (any age)
  - Choroid plexus cancer (any age)
  - Breast cancer (≤30 years)
  - Triple positive breast cancer (≤35 years)
  - ≥2 LFS-related cancers\* (both occurring ≤46 years; 2 breast cancers not eligible)
  - $\geq$ 1 LFS-related cancer\* with  $\geq$ 1 1<sup>st</sup> / 2<sup>nd</sup> degree relative with  $\geq$ 1 LFS-related cancer\* (one case  $\leq$ 46 years, the other  $\leq$ 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)

- Clinical Genetics
- Oncology

<sup>\*</sup> Sarcoma of bone/soft tissue, breast cancer, brain cancer, adrenocortical cancer or any childhood cancer (occurring ≤ 18 years)

# LYNCH SYNDROME (HNPCC)

#### **AVAILABLE TESTING**

Centre	Method		Scope and ra	inge of test	Targets	TAT
Edinburgh	NGS MLPA	Whole	gene screen	SNVs, indels Exon level CNV	MLH1, MSH2, MSH6, PMS2, EPCAM (del exons 8-9), POLD1 (exons 4-12), POLE (exons 3-13)	56
Family me	mber testing	]		as indicated above		14
Proforma re	quired?	d? YES Colored		ectal cancer gene panel proforma (see centre website)		

### REFERRAL CRITERIA

- Clinical Criteria for germline testing in an affected individual
  - a. The proband has a dMMR tumour where results of BRAF and/or MLH1 hypermethylation testing suggest Lynch syndrome
  - o b. The proband is diagnosed with colorectal cancer  $\le$  40 , irrespective of the dMMR status of the tumour
  - o c. The affected proband comes from a modified Amsterdam criteria positive family irrespective of the dMMR status of the tumour
  - d. (Wimmer score =>3)
- Clinical Criteria for germline testing in an unaffected individual
  - o a. First degree relative affected with Lynch-related cancer, AND
  - b. Family history of colorectal cancer/Lynch-related cancers reaches Amsterdam
     Criteria (≥3 cases over ≥2 generations with ≥1 case affected ≤50 years) AND
  - o c. 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
  - o d. Somatic sequencing is not possible, or failed, AND
  - o e. No living affected individual is available for genetic testing

#### REQUESTING SPECIALTIES

# **MEDULLARY THYROID CANCER**

# **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Dundee	NGS	Targete	d screen	SNVs, indels	RET (exons 5, 8, 10, 11, 13, 14, 15, 16)	56
Family mem	nber testing			as ind	icated above	14
Proforma re	quired?	YES	Endocrine	disorders proforma	(see centre website)	

# REFERRAL CRITERIA

- Medullary thyroid cancer (MTC) at any age.
- See entry for Multiple Endocrine Neoplasia Type 2A, Type 2B and Medullary Thyroid Cancer

- Clinical Genetics
- Endocrinology

### PEUTZ-JEGHERS SYNDROME

#### **AVAILABLE TESTING**

Centre	Method		Scope and ra	inge of test	Targets	TAT
Edinburgh	NGS MLPA	Whol	e gene screen	SNVs, indels Exon level CNV	STK11	56
Family me	Family member testing			as indicated above		14
Proforma required? YES			Colorectal canc	er gene panel proforma (s	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
  - o 3. Characteristic mucocutaneous pigmentation age
  - o 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
  - o 7. Characteristic mucocutaneous pigmentation, 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)
- 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

# **POLYPOSIS**

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT	
Edinburgh	NGS	Whole gene screen	SNVs, indels	APC, BMPR1A, MBD4, MSH3, MUTYH, NTHL1, POLD1 (exons 4-12), POLE (exons 3-13), PTEN, RNF43, SMAD4, STK11, GREM1 (upstream dup)	56	
Family men	nber testing		as indicated above 14			
Proforma re	quired?	YES	Colorectal cancer gene panel proforma (see centre website)			

### REFERRAL CRITERIA

- ≥5 polyps and colorectal cancer (<60 years) OR
- ≥10 polyps (age <60 years), OR
- ≥20 polyps (age ≥ 60 years), OR
- ≥5 polyps AND first degree relative with ≥5 polyps OR CRC (age <60years), OR
- ≥10 polyps (age ≥ 60 years) AND first degree relative with ≥5 polyps OR CRC (age <60 years).

Please note small hyperplastic rectal polyps can be ignored/not included in the count.

# Additional testing for:

- Desmoid tumour (APC and MUTYH)
- CHRPEs (APC only)
- Mosaic FAP: indicated if phenotype consistent with a diagnosis of FAP or attenuated FAP and result will impact on clinical management of relatives.

# REQUESTING SPECIALTIES

# **RENAL CANCER**

# **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS	Whole gene screen	SNVs, indels, Exon level CNV for FLCN, VHL, SDHB	BAP1, FH, FLCN, MET, PTEN, SDHB, VHL	56
Family men	nber testing		as indicated above		14
Proforma re	roforma required? 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
  - o Renal cancer AND first / second degree relative with renal cancer
  - o Renal cancer and features of an inherited renal cancer syndrome

# REQUESTING SPECIALTIES

# RHABDOID TUMOUR

# **AVAILABLE TESTING**

Centre	Method		Scope and ra	nge of test	Targets	TAT
Glasgow	NGS	Whole	Whole gene screen SNVs, i		SMARCA4, SMARCB1	56
Family men	nber testing			as indicated abo	ve	14
Proforma requ	uired?	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

# **METABOLIC**

# AMINO ACID DISORDERS & DISORDERS OF NEUROTRANSMISSION

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT	
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	ABAT, ALDH18A1, ALDH5A1, ALDH7A1, AMT, ASPA, CBS, CTH, D2HGDH, DBH, DDC, FAH, GABRG2, GCDH, GCH1, GLDC, GLRA1, HGD, L2HGDH, MAT1A, OAT, PAH, PCBD1, PNPO, QDPR, SLC25A22, SLC6A19, SLC7A7, SUOX	112	
Family men	nber testing		as indicated above		14	
Proforma re	quired?	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.

- Clinical Genetics
- Metabolic

# **BATTEN DISEASE**

# **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger MLPA	Whole gene SNVs, indels screen Exon level CNV		TPP1	56
Family me	ember testing		as indicated above		
Proforma re	na 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 re	quired?	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 men	nber testing		as indicated above		
Proforma required?		NO	_		

### REFERRAL CRITERIA

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

### **REQUESTING SPECIALTIES**

- Clinical Genetics
- Metabolic

# CARNITINE PALMITOYLTRANSFERASE II DEFICIENCY

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	CPT2	56
Family men	nber testing		as indic	ated above	14
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, indels	FOLR1	56
Family men	nber testing		as indicated above		
Proforma required? 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 men	nber testing		as indicated above			
Proforma re	quired?	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	ММАСНС	56	
Family men	nber testing		as indicated above			
Proforma re	quired?	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 men	nber testing		as indicated above		14
Proforma re	quired?	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 gene screen	SNVs, indels Exon level CNV	ACADM, ACADS, ACADVL, ARG1, ASL, ASS1, CPS1, CPT1A, CPT2, ETFA, ETFB, ETFDH, GLUD1, HADHA, HADHB, HMGCL, HMGCS2, IVD, MMAA, MMAB, MMACHC, MMADHC, MUT, NAGS, OAT, OTC, OXCT1, PCCA, PCCB, SLC22A5, SLC25A13, SLC25A15, SLC25A20, SLC7A7, SLC52A2, SLC52A3	112
Family member testing		as indicated above			
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.

- 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 gene screen	SNVs, indels Exon level CNV	AGL, ALDOA, ENO3, EPM2A, FBP1, G6PC, G6PC3, GAA, GALE, GALK1, GALT, GBE1, GYG1, GYS1, GYS2, LAMP2, LDHA, NHLRC1, PFKM, PGAM2, PGK1, PGM1, PHKA1, PHKA2, PHKB, PHKG2, PRKAG2, PYGL, PYGM, SLC2A2, 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.

- Clinical Genetics
- Metabolic

# **FABRY DISEASE**

# **AVAILABLE TESTING**

Centre	Method	Scope and ra	nge of test	Targets	TAT	
Edinburgh	Sanger	Whole gene screen	SNVs, indels	GLA	56	
Family member testing		as indicated above				
Proforma required?		NO				

### REFERRAL CRITERIA

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

- Clinical Genetics
- Metabolic

### FAMILIAL HYPERCHOLESTEROLAEMIA

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole gene screen		SNVs, indels Exon level CNV* (*LDLR only)	LDLR,APOE, PCSK9, APOB, LDLRAP1	56
Family member testing			as indicated above			14
Proforma required?		NO	Optional FH proforma on centre website			

#### 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)

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
- . 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

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
- Lipidology
- Metabolic

# **FANCONI-BICKEL SYNDROME**

# **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	SLC <sub>2</sub> A <sub>2</sub>	56
Family men	Family member testing as inc			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 gene screen	SNVs, indels Exon level CNV	ACADM, ACADS, ACADVL, CPT1A, CPT2, ETFA, ETFB, ETFDH, HADHA, HADHB, HMGCL, HMGCS2, IVD, MMAA, MMAB, MMACHC, MMADHC, OXCT1, SLC22A5, SLC25A20, SLC52A2, SLC52A3	112
Family men	nber testing		as indicated above		14
Proforma re	roforma 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	I range of test	Targets	TAT
Aberdeen	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV	GALT	56
Family men	Family member testing as indicated a			pove	14
Proforma required?		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 indic			ated above	14	
Proforma required? N		NO			

#### REFERRAL CRITERIA

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

- Clinical Genetics
- Metabolic

#### **GLUTARIC ACIDAEMIA TYPE 1**

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	GCDH	56
Family member testing as ir			as in	dicated above	14
Proforma re	quired?	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 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	quired?	NO			

#### REFERRAL CRITERIA

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

#### **REQUESTING SPECIALTIES**

- Clinical Genetics
- Metabolic

#### **HOMOCYSTINURIA**

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	CBS, MMADHC, MTHR, MTR, MTRR	56
Family men	nber testing	as indica		ted above	14
Proforma required? NO		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 ran	ge of test	Targets	TAT
Aberdeen	Sanger	Targeted screen	SNVs	APOE (Codons p.130 and p.176)	28
Proforma re	quired?	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 MLPA	Whole gene screen	SNVs, indels Exon level CNV* (* <i>LPL</i> only)	LPL, LMF1, APOC2, APOA5, GPI-HBP1	56
Family men	Family member testing as indicated above		ated above	14	
Proforma required? NO					

#### REFERRAL CRITERIA

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

- Clinical Genetics
- Gastrohepatology
- Lipidology

# **HYPOBETALIPOPROTEINAEMIA**

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	ANGPTL3, APOB, MTTP, PCSK9, SAR1B	56
Family men	nber testing	r testing as indicated above		cated above	14
Proforma required?		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 gene screen	SNVs, indels Exon level CNV	AGA, ARSA, ARSB, 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, IDS, IDUA, LIPA, MAN2B1, MANBA, MCOLN1, MFSD8, NAGA, NAGLU, NEU1, NPC1, NPC2, PPT1, PSAP, SGSH, SLC17A5, SMPD1, SUMF1, TPP1	112
Family men	Family member testing as indicated above		as indicated above	14	
Proforma required? 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.

- 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 Exon level CNV	BCKDHA, BCKDHB, DBT	56
Family men	Family member testing as indicate		d above	14	
Proforma required? NO		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	en SNVs, indels ACADM Exon level CNV		56
Family men	Family member testing as indicated		above	14	
Proforma required? NO		NO			

#### REFERRAL CRITERIA

- Clinical features suggestive of MADD
- 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 men	ly member testing as inc		icated above	14	
Proforma required?		NO			

# REFERRAL CRITERIA

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

#### **REQUESTING SPECIALTIES**

- Clinical Genetics
- 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 men	nber testing	er testing as indicated		ated above	14
Proforma required?		NO			

#### REFERRAL CRITERIA

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

- Clinical Genetics
- Metabolic

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

# **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	GNPTAB	
Family men	Family member testing as indi		cated above	14	
Proforma required?		NO			

#### REFERRAL CRITERIA

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

#### **REQUESTING SPECIALTIES**

- 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 Exon level CNV	ETFDH, ETFA, ETFB, SLC52A2, SLC52A3	56
Family men	nber testing		as in	dicated above	14
Proforma re	Proforma required? NO				

#### REFERRAL CRITERIA

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

- Clinical Genetics
- Metabolic
- Neurology

# **NEURONAL CEROID LIPOFUSCINOSIS (NCL)**

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT			
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	CLN3, CLN5, CLN6, CLN8, CTSD, DNAJC5, KCTD7, MFSD8, PPT1, TPP1	56			
Family men	Family member testing as indicated above				14			
Available ge	Available genes: See website							
Proforma re	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.

- Clinical Genetics
- Metabolic
- Neurology

#### **NIEMANN-PICK DISEASE**

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	NPC1, NPC2, SMPD1	56
Family member testing as indicated above		cated above	14		
Proforma required? NO		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 men	nber testing	er testing as inc		icated above	14
Proforma required? NO		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 screen	SNVs, indels Exon level CNV	NPC1, NPC2	56
Family men	nber testing		as indicated 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 screen	SNVs, indels Exon level CNV	ALDH7A1, AMT, GLDC, PPT1, TPP1	56
Family men	nber testing		as indicated above		14
Proforma required? 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 gene screen	SNVs, indels Exon level 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, PCCA, PCCB, PRDX1, SLC19A3, SLC46A1, SLC52A3, SUCLA2, SUCLG1, TAZ, TCN2, TMEM70	112
Family member testing			as indicated above	14	
Proforma re	quired?	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

#### 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 inc		icated above	14		
Proforma required?		NO			

#### REFERRAL CRITERIA

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

#### **REQUESTING SPECIALTIES**

- Clinical Genetics
- Metabolic

#### ORNITHINE TRANSCARBAMULASE DEFICIENCY

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	NGS MLPA	Whole gene screen	SNVs, indels Exon level CNV	ОТС	56
Family men	Family member testing as indicated above		ated above	14	
Proforma required? NO		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 gene screen	SNVs, indels Exon level CNV	ABCD1, ACOX1, AGPS, AGXT, AMACR, CAT, DNM1L, GNPAT, HSD17B4, PEX1, PEX2, PEX3, PEX5, PEX6, PEX7, PEX10, PEX11B, PEX12, PEX13, PEX14, PEX16, PEX19, PEX26, PHYH, SCP2	112
Family mem	nber testing	er 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.

- Clinical Genetics
- Metabolic

#### **PHENYLKETONURIA**

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV	PAH	
Family men	nber testing		as indicated above		14
Proforma re	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 screen	SNVs, indels Exon level CNV	GAA	
Family men	nber testing	as indicated above		14	
Proforma required? NO		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 Exon level CNV	PCCA, PCCB	
Family member testing as indicated above		ited above	14		
Proforma required? NO		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	ole gene screen SNVs, indels Exon level CNV PEX7, PHYH		56
Family member testing as indicated above		ited above	14		
Proforma required? NO		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 screen	SNVs, indels	ALDH5A1	56
Family member testing as indicate			ed above	14	
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	e of test	Targets	TAT
Aberdeen	Sanger	Whole gene screen Long Range PCR	SNVs, indels Ex3-9 deletion	$TANGO_2$	
Family men	Family member testing as indicated abo			ve	14
Proforma re	quired?	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, indels	HEXA	56
Family member testing as in		dicated 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	FMO <sub>3</sub>	56
Family men	ember testing as inc		as ind	icated above	14
Proforma required?		NO			

#### REFERRAL CRITERIA

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

- Clinical Genetics
- Metabolic

# **VLCAD DEFICIENCY**

# **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	ACADVL	56
Family member testing		as indic	ated above	14	
Proforma required? NO		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, indels	Common LHON mitochondrial DNA variants (m.346oG>A, m.11778G>A, m.14484T>C)	28
Family men	Family member testing as		s indicated above	14	
Proforma required?		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	Targeted screen	SNVs, indels	Common mitochondrial DNA variants MT-TL1:m.3243A>G MT-TK:m.8344A>G MT-ATP6:m.8993T>G/C Plus others relevant to phenotype	28
Family men	nber testing		as indicated above		14
Proforma required? 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
  - o myoclonic epilepsy with ragged red fibres (MERRF)
  - o neuropathy, ataxia and retinitis pigmentosa (NARP)
  - o 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	Sanger	Targeted screen	SNVs, indels	MTTL1 m. 3243A>G	28
Family member testing as in		dicated 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	Sanger	Targeted screen	SNV	MTTL1 m. 3243A>G	28
Family men	nber testing		as indicated 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
Classion	MLPA	Targeted screen	Exon level CNV	DMD	28
Glasgow	Sanger	Whole gene screen	SNVs, indels	DMD	56
Family men	nber testing	sting as indicated above		14	
Proforma required?		NO			

#### REFERRAL CRITERIA

- Clinical features that include:
  - o Progressive symmetric muscle weakness
  - o 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, indels	AGPS, ARSE, EBP, GNPAT, PEX7	56
Family member testing			as indica	ted above	14
Proforma re	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
Classicu	MLPA	Targeted screen	Exon level CNV	DMD	28
Glasgow	Sanger	Whole gene screen	SNVs, indels	DMD	56
Family men	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 men	nber testing		as indicated above		
Proforma required?		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 member testing as ind		icated above	14		
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	SNVs, indels	EXT1, EXT2	56
Family men	nber testing	as inc	licated above		14
Proforma required?		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	56
Family men	Family member testing as inc		icated above	14	
Proforma required?		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 Edinburgh Glasgow	PCR & TP-PCR	Targeted screen	Triplet repeat expansion	DMPK	28 Prenatal 3
Proforma required?		NO			

#### REFERRAL CRITERIA

- Clinical phenotype that could be consistent with (or "that raises suspicion of") myotonic dystrophy type 1. Suggestive features include
  - Hypotonic infant with or without joint contractures
  - o 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 an	d range of test	Targets	TAT
Aberdeen	PCR & QP-PCR	Targeted screen	4bp repeat expansion	ZNF9	28
Proforma required?		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 ra	nge of test	Targets	TAT
Dundee	PCR and Sanger	Targeted screen	Repeat expansion, SNV	PABPN1 – GCN repeat expansion and c.35G>C	28
Proforma required?		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 gene screen	SNVs, indels	BMP1, COL1A1, COL1A2, CREB3L1, CRTAP, FAM46A, FKBP10, IFITM5, P3H1 (LEPRE1), PLOD2, PLS3, PPIB, SERPINF1, SERPINH1, SP7, SPARC, TMEM38B, WNT1	112
Family member 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
  - o Short stature
  - o Dentinogenesis imperfect
  - o 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 gene screen	SNVs, indels	AMER1, ANKH, CA2, CLCN7, CTSK, FAM2oC, FERMT3, LEMD3, LRP5, OSTM1, PTH1R, RASGRP2, SNX1o, SOST, TCIRG1, TGFB1, TNFRSF11A, TNFSF11, TYROBP	112
Family member testing		as indicated above			14
Proforma required?		NO			

# REFERRAL CRITERIA

• Characteristic radiographic changes

- Adult Orthapaedics
- Clinical Genetics
- Paediatricis specialising in bone marrow transplantation, haematology, metabolic disease or orthapaedics

# PRIMORDIAL DWARFISM, MICROCEPHALY

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	NGS	Whole ger screen	SNVs, indels	NKRD11, ASPM, ATR; ATRIP, ATRX, BLM, CASK, CDC45, CDC6, CDKN1C, CDK5RAP2, CDT1, CENPE, CENPF, CENPJ, CEP135, CEP152, CEP63, CREBBP, DNA2, DNMT3A, DONSON, DPP6, DYRK1A, EP300, GMNN, IGF1, IGF1R, KIF11, KMT2A, KNL1, LARP7, LIG4, MCPH1, MRE11, NBN, NDE1, ORC1, ORC4, ORC6, PCNT, PHC1, PLK4, PNKP, POC1A, POLE, RAD50, RBBP8, RNU4ATAC, SMARCAL1, SRCAP, STIL, TCF4, TOP3A, TRAIP, TUBGCP6, VPS13B, WDR4, WDR62, XRCC4	112
Family men	nber testing			as indicated above	14
Proforma required?		YES	Primordial Dwarfish website)	n and Microcephaly gene panel referral proforma (see centr	e

## 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	GDF <sub>5</sub> , NOG1	56
Family men	Family member testing as indice		ated above	14	
Proforma required? NO					

### REFERRAL CRITERIA

• Clinical features strongly suggestive of proximal symphalangism

## **REQUESTING SPECIALTIES**

Clinical Genetics

# 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 screen	SNVs, indels	BRAF, CBL, HRAS, KRAS, LZTR1, MAP2K1, MAP2K2, MRAS, NF1, NRAS, PPP1CB, PTP11, RAF1, RIT1, SHOC2, SOS1, SPRED1	112
Family men	Family member testing as indicated above		as indicated above	14	
Proforma required? NO				·	

## REFERRAL CRITERIA

- At least 2 of the suggestive clinical features:
  - o Early feeding difficulty / failure to thrive
  - o Relative macrocephaly
  - o 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 a	nd range of test	Targets	TAT
Aberdeen Dundee	Karyotype	Whole genome screen	Structural rearrangements CNV	Whole genome	28
Edinburgh Glasgow	Microarray	Whole genome screen	CNV	Whole genome	28
Glasgow	Sanger	Whole gene screen	SNVs, indels	SHOX	56
Proforma required?		NO			

## REFERRAL CRITERIA

• Disproportionate short stature

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 rang	ge of test	Targets	TAT
Edinburgh	NGS	Whole gene screen	SNVs, indels	ACAN, ACP5, ADAMTS10, ADAMTSL2, AGPS, ALPL, ANKH, ARSE, B3GALT6, BMP1, BMPR1B, CA2, CANT1, CDC6, CDKN1C, CDT1, CHST3, CLCN7, COL10A1, COL11A1, COL11A2, COL1A1, COL1A2, COL9A3, 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 men	nber testing		as i	ndicated above	14
Proforma re	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
- 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 & TP-PCR	Targeted screen	Triplet repeat expansion	AR	28
Proforma re	quired?	NO			

## REFERRAL CRITERIA

• Clinical features that indicate a likely diagnosis of SBMA

## REQUESTING SPECIALTIES

Clinical Genetics

# **NEUROLOGY**

## **AICARDI-GOUTIERES SYNDROME**

### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS	Whole gene screen	SNVs, indels	ADAR, IFIH1, RNASEH2A, RNASEH2B, RNASEH2C, SAMHD1, TREX1	112
Family men	Family member testing			as indicated above	14
Proforma required? 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'
  - o 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	ТАТ
Glasgow	NGS	Whole gene screen	SNVs, indels	ACTA1, ADAMTS10, ANTXR2, ASCC1, ASXL1, B3GALNT2, B4GAT1, BICD2, CHAT, CHRNA1, CHRNB1, CHRND, CHRNE, CHRNG, CHST14, CNTNAP1, COL12A1, COL6A1, COL6A2, COL6A3, COLO, DAG1, DNM2, DOK7, DPAGT1, DYNC1H1, ECEL1, ERCC6, ERCC8, EXOSC3, FAM2OC, 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 men	nber testing			as indicated above	14
Proforma re	equired?	NO			

## 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.

## **REQUESTING SPECIALTIES**

• Clinical Genetics

# CADASIL

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	Sanger	Whole gene screen	SNVs, indels	NOTCH3	56
Family member testing as ir			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	inge of test	Targets	TAT
Dundee	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV	RASA1	56
Family men	Family member testing as indicated		above	14	
Proforma re	equired?	NO			

## REFERRAL CRITERIA

- Capillary malformations are the hallmark of capillary malformation-arteriovenous malformation (CM-AVM) syndrome.
- CV-AVM should be suspected in an individual with
  - o CM, generally multifocal, small, composed of dilated capillaries, localised on face and limbs
  - o 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	cgORF72	28
Edinburgh	NGS	Whole gene screen	SNVs, indels	ALS2, ANG, ANXA11, APP, CHMP2B, CSF1R, DCTN1, FIG4, FUS, GRN, ITM2B, MAPT, NEK1, OPTN, PFN1, PRNP, PSEN1, PSEN2, SETX, SOD1, SQSTM1, TARDBP, TBK1, UBQLN2, VAPB, VCP	112
Family member testing		as indicated	l 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

# **CORTICAL BRAIN MALFORMATIONS**

# AVAILABLE TESTING

Centre	Method	Scope and rang	je of test	Targets	TAT		
Dundee	NGS	Whole gene screen	SNVs, 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, VLDLR, WDR6	112		
Family mer	nber testing		a	s indicated above	14		
Proforma required?		NO					

# REFERRAL CRITERIA

• Cortical brain malformation with features suggestive of a monogenic cause

- Clinical Genetics
- Neurology

# CREUTZFELD-JAKOB DISEASE (CJD)

# AVAILABLE TESTING

Centre	Method	Sco	pe and range of test	Targets	TAT
Edinburgh	PCR	Repeat-primed PCR	Octapeptide repeat expansion	PRN	28
Edinburgh	Sanger	Whole gene screen	SNVs, indels	PRN	56
Family men	Family member testing		as indicated above		14
Proforma required? NO		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	c9ORF72	28
Dundee	NGS	Whole gene screen	SNVs, indels	APP, CHMP2B, CSF1R, DNAJC5,DNMT1,EPM2A, GRN, ITM2B, MAPT, NHLRC1, NOTCH3, PSEN1, PSEN2, PRNP, TBK1, TARDBP, TYROBP, UBQLN2, VCP	112
Family member testing as indi		as indicated	d above	14	
Proforma re	quired?	NO			

## REFERRAL CRITERIA

- Unexplained dementia with:
  - Age at onset <55 years where acquired causes (e.g. stroke, tumour) have been excluded, OR
  - o 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 and range of test		Targets	TAT
Edinburgh	PCR & TP-PCR	Targeted screen Triplet repeat expansion		ATN1	28
Proforma re	quired?	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	Whole gene screen	SNVs, indels	ACTB, AFG3L2, ANO3, APTX, ATM, ATP1A2, ATP7B, C190rf12, 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 mer	mber testing		as in	dicated above	14
Proforma re	equired?	NO			

## REFERRAL CRITERIA

- Unexplained dystonia, chorea or related movement disorder with onset in adulthood with a likely monogenic cause
- Overlapping indications: Parkinson's Disease

- Clinical Genetics
- Neurology

# **EPILEPSY**

# AVAILABLE TESTING

Centre	Method	Scope and rang	ge of test	Targets	TAT
Glasgow	Microarray	Whole genome screen	CNV	Whole genome	28
Glasgow	NGS	Whole gene screen	SNVs, 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 or 112
Family mer	nber testing			as indicated above	14
Proforma re	quired?	Proforma required? NO			

# 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, indels	KCNA1, CACNA1A	56
Family men	Family member testing as in		dicated above	14	
Proforma re	equired?	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	TAT
Dundee	NGS	Whole gene screen	SNVs, indels	ADCY5, ATP1A2, ATP1A3, ATP7B, CACNA1A, CACNB4, GLRA1, GLRB, KCNA1, KCNJ2, KCNMA1, KCNQ2, KCNQ3, PNKD, PRRT2, SCN1A, SCN8A, SLC1A3, SLC2A1, SLC6A5, SPR	112
Family men	Family member testing as indicated above		as indicated above	14	
Proforma re	equired?	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	Whole gene screen	SNVs, indels	KRIT1, CCM2, CCM3	56
Family member testing as indicate			ed 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 screen	SNVs, indels	ATP1A2, CACNA1A, PRRT2, SCN1A, SLC2A1	56
Family men	Family member testing				14
Proforma re	equired?	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 Edinburgh Glasgow	PCR & TP-PCR	Targeted screen	Triplet repeat expansion	FMR1	28
Proforma requ	Proforma required?				

### 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 & 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	I range of test	Targets	TAT
Edinburgh	PCR & TP-PCR	Targeted screen	Triplet repeat expansions	SCA1, SCA2, SCA3, SCA6, SCA7, SCA8, SCA17, FRDA, FMR1, FRDA	28
Edinburgh*	NGS	Whole gene screen	SNVs, indels	BCB7, ABCD1, AFG3L2, ALS2, ANO10, APTX, ATL1, ATM, ATP1A3, ATP7B, BSCL2, CACNA1A, CACNA1G, CAPN1, COQ8A, CYP27A1, CYP7B1, DDHD2, FA2H, FGF14, FTL, FXN, GBA2, GCH1, GRID2, HSPD1, ITPR1, KCNA1, KCNC3, KCND3, KIF1A, KIF5A, L1CAM, NIPA1, PDYN, PLP1, PNPLA6, POLG, PRKCG, PRNP, PRRT2, REEP1, RTN2, 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 gene 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 meml	per testing			as indicated above	14
Proforma req	uired?	NO			

<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
  - o 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>

- Clinical Genetics
- Neurolology

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

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

### **AVAILABLE TESTING**

Centre	Method	Scope and ra	nge of test	Targets	TAT
Aberdeen	Sanger MLPA	Whole Gene screen	SNVs, indels Exon level CNV*	PMP22*, MPZ, GJB1, MFN2	56
Family me	ember testing		as indicated a	bove	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 rar	nge of test	Targets	TAT
Aberdeen	Sanger MLPA	Whole Gene screen	SNVs, indels Exon level CNV	PMP22	56
Family me	ember testing		as indicat	ed above	14
Proforma red	juired?	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 and	range of test	Targets	TAT
Edinburgh*	NGS MLPA	Whole gene screen	SNVs, indels CNV**	ABCB7, ABCD1, AFG3L2, ALS2, ANO10, APTX, ATL1**, ATM, ATP1A3, ATP7B, BSCL2, CACNA1A, CACNA1G, CAPN1, COQ8A, CYP27A1, CYP7B1, DDHD2, FA2H, FGF14, FTL, FXN, GBA2, GCH1, GRID2, HSPD1, ITPR1, KCNA1, KCNC3, KCND3, KIF1A, KIF5A, L1CAM, NIPA1, PDYN, PLP1, PNPLA6, POLG, PRKCG, PRNP, PRRT2, REEP1**, 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 MLPA	Whole gene screen	SNVs, indels, 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 meml	per testing		•	as indicated above	14
Proforma req	uired?	YES Ed	inburgh only – HS	P referral proforma (see centre website)	

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

### REFERRAL CRITERIA

- Exclusion of metabolic, neoplastic, alcohol, and drug-related causes
- Normal/routine neurological bloods, and vitamin E testing
- Negative spinocerebellar ataxia repeat expansion panel, including FXTAS and FA testing
- MRI neuroimaging normal, or isolated cerebellar atrophy
- Family history of ataxia, or young age of onset (<50)

- Clinical Genetics
- Neurology

<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	Whole gene screen	SNVs, indels	CDON, DHCR7, DISP1, FGF8, FGFR1, GLI2, PTCH1, SHH, SIX3, TGIF1, ZIC2	112
Family men	amily member testing as in		dicated above	14	
Proforma re	quired?	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 an	d range of test	Targets	TAT
Edinburgh	TPPCR	Targeted screen	Triplet repeat expansion	НТТ	14 Prenatal 3
Edinburgh	Linkage	Targeted Screen	Exclusion testing	НТТ	14 Prenatal 3
Proforma required?		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)

## **LESCH-NYHAN SYNDROME**

## **AVAILABLE TESTING**

Centre	Method	Scope and ra	nge of test	Targets	TAT
Glasgow	Sanger MLPA	Whole Gene screen	SNVs, indels Exon level CNV	HPRT1	56
Family me	ember testing		as indicated a	bove	14
Proforma required?		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	Whole gene screen	SNVs, indels	EIF2B1, EIF2B2, EIF2B4, EIF2B5, EIF2B3	112
Family men	nber testing		as indicated above		
Proforma re	quired?	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 gene screen	SNVs, indels	ATP13A2, C19ORF12, COASY, CP, DCAF17, FA2H, FTL, FUCA1, KIF1A, KMT2B, MECR, PANK2, PLA2G6, PSEN1, SCP2, SLC39A14, SQSTM1, TRIM32, UBTF, VPS13A, WDR45	112
Family men	nber testing			as indicated above	
Proforma required? 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	Whole gene screen	SNVs, indels Exon level CNV	NF1	56
Family men	nber testing		as indicated above		
Proforma re	quired?	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:
  - O At least 6 café au lait macules (at least 0.5cm in a child and 1.5cm in an adult)
  - o At least 2 subcutaneous or cutaneous neurofibromas
  - Plexiform neurofibroma
  - Optic glioma
  - o At least 2 Lisch nodules
  - o Bony dysplasia (sphenoid wing, long bone bowing, pseudarthrosis)
  - o Family history of NF1

- Clinical Genetics
- Paediatrics

## **PAIN DISORDERS**

## AVAILABLE TESTING

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS	Whole gene screen	SNVs, indels	ATL1, ATL3, ELP1, GLA, KIF1A, NGF, NTRK1, PRNP, RAB7A, RETREG1, SCN10A, SCN11A, SCN9A, SEPT9, SPTLC1, SPTLC2, TRPA1, TTR, WNK1	112
Family men	nber testing			as indicated above	
Proforma re	equired? NO				

### REFERRAL CRITERIA

- This includes the disorders:
  - o Congenital insensitivity to pain
  - o Inherited erythromelalgia
  - o Paroxysmal extreme pain disorder
  - o 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	Whole ger screen	SNVs, indels, Exon level CNV	ATP13A2, ATP1A3, DCTN1, DNAJC13, DNAJC6, FBXO7, FTL, GBA, GCH1, GRN, LRRK2, MAPT, PARK7 (DJ-1), PINK1, PLA2G6, PRKN (Parkin), RAB39B, SLC30A10, SNCA, SPG11, SYNJ1, TH, VPS35 CNV in SNCA, PARK2, PINK1, PARK7, ATP13A2, LRRK2, GCH1 and UCHL1	112
Family member testing			as indicated above	14	
Proforma required? YES Parkinson's Disease proforma		proforma			

# REFERRAL CRITERIA

- Parkinson's disease or complex Parkinsonism
  - o Age at onset <50 years, OR
  - o First degree relative affected at <50 years, OR
  - 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 t	est	Targets	TAT
Dundee	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV ( <i>PLP</i> 1)	PLP1, GJC2(GJA12)	56
Family member testing as ir		as indicated a	bove	14	
Proforma required?		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> (p.Leu689lle, p.lle693Thr, p.Thr704Met and p.Met1592Val )	56	
Family member testing		as indicated above				
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) SCN4A (codons 669, 672)	56	
Family member testing		as indicated above				
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</li>
- Three or more of the following six clinical/laboratory features:
  - Onset in the first or second decade
  - o 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
  - o A family history of the condition or genetically confirmed skeletal calcium or sodium channel mutation
  - o 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 a	nd range of test	Targets	TAT		
Glasgow	NGS	Whole gen screen	e 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 required?		YES	YES Peripheral neuropathy proforma (see centre website)				

## REFERRAL CRITERIA

- Firm clinical diagnosis of CMT
- Length dependent neuropathy on neurophysiology
- No pathogenic variant on first tier CMT testing (performed in Aberdeen)

- Clinical Genetics
- Neurology approved by Clinical Genetics

# **PORENCEPHALY**

## **AVAILABLE TESTING**

Centre	Method		Scope and range of to	Targets	TAT	
Dundee	NGS		Whole gene screen	SNVs, indels	COL4A1, COL4A2	112
Family member testing		as indicated above				14
Proforma required?		NO				

# REFERRAL CRITERIA

• Any individual with clinical features consistent with the condition

- Clinical Genetics
- Neurology

# **RETT (& RETT-LIKE) SYNDROME**

## **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Glasgow	Sanger MLPA	Whole gene screen		SNVs, indels Exon level CNV	MECP2, CDKL5	56
Family member testing		as indicat			icated above	14
Proforma required?		NO				

## REFERRAL CRITERIA

- Clinical features that include:
  - o Rapid developmental regression in infancy
  - o Seizures
  - o Severe intellectual disability
  - o Stereotypic hand movements
  - o Deceleration of head growth

- Clinical Genetics
- Paediatrics

### RHABDOMYOLOSIS & METABOLIC MYOPATHIES

### **AVAILABLE TESTING**

Centre	Method	Scope a	and range of test	Targets	TAT
Glasgow	NGS	Whole gen	SNVs, indels	ACADVL, AGL, ALDOA, ANO5, CACNA1S, CAPN3, CAV3, CPT2, DMD, DYSF, ENO3, ETFA, ETFB, ETFDH, FKRP, GAA, GBE1, GMPPB, GYG1, GYS1, HADHA, HADHB, ISCU, LDHA, LPIN1, PFKM, PGAM2, PGK1, PGM1, PHKA1, PNPLA2, PYGM, RBCK1, RYR1, SLC22A5, TANGO2	112
Family member testing				as indicated above	14
Proforma required?		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
  - o Mitochondrial myopathy/PEO considered and excluded where appropriate
- IN ADDITION PATIENTS AGED >10 years must fulfil at least one of the following three criteria:
  - o 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 K>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
  - o Moderate to profound XS lipid or glycogen on biopsy
  - o Cores/minicores on biopsy
- Muscle MRI characteristic of RYR1

- Clinical Genetics
- Metabolic
- Neurology

## SPINAL MUSCULAR ATROPHY

### **AVAILABLE TESTING**

Centre	Method	Sco	pe and ran	ge of test	Targets	TAT
Edinburgh	MLPA	Targete	d screen	CNV	SMN1	28
Glasgow	NGS	Whole ge	ne screen	SNVs, 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, SYT2, TRPV4, UBA1, VAPB, VCP, VRK1	112
Family member testing				as i	ndicated above	14
Proforma required?		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)
  - o 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
  - Carrier testing for partners of confirmed *SMN*<sub>1</sub> carriers.
- Whole gene screen
  - o dHMN/SMA clinical phenotype AND
  - o Compatible neurophysiology (not required in infants) AND
  - o 5q linked SMA excluded (not required in infants)

- Clinical Genetics
- Neurology

# SPINOCEREBELLAR ATAXIA 17 (SCA17)

# **AVAILABLE TESTING**

Centre	Method	Scope an	d range of test	Targets	TAT
Edinburgh	PCR & TP-PCR	Targeted screen	Triplet repeat expansion	TBP, AR	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 rang	e of test	Targets	TAT
Aberdeen Dundee	PCR	Targeted screen Deletion		<i>DYT1</i> (c.907_909del) Gene known as <i>TOR1A</i>	28
Proforma re	quired?	NO			

# REFERRAL CRITERIA

- DYT1 early-onset isolated dystonia should be suspected in individuals with
  - Onset of dystonia before the age of 26
  - o Isolated dystonia with no other abnormalities on neurologic examination, normal routine neuroimaging, no known cause of acquired dystonia
  - o 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 rar	nge of test	Targets	TAT
Dundee	NGS	Whole gene screen SNVs, indels		TSC1, TSC2	56
Family men	nber testing		as indicated	d above	14
Proforma required?		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		indicated above	14
Proforma requir	red?	NO	_		

#### REFERRAL CRITERIA

- 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
- Clinical features of Alport syndrome (high tone sensorineural hearing loss or characteristic ophthalmic signs such as perimacular flecks or anterior lenticonus)
- Haematuria / proteinuria

- Clinical Genetics
- Nephrology

# **BARTTER SYNDROME & GITELMAN SYNDROME**

#### **AVAILABLE TESTING**

Centre	Method		Scope and	d range of test	Targets	TAT
Dundee	NGS	Whole gene screen		SNVs, indels Exon level CNV ( <i>CLCNKB</i> ) if appropriate	BSND, CLCNKB, KCNJ1, SLC12A1, SLC12A3	56
Family men	nber testing			2	14	
Proforma required? YES Renal Genet		tics Proforma (see centre web	osite)			

#### REFERRAL CRITERIA

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

#### **REQUESTING SPECIALTIES**

- Clinical Genetics
- Nephrology

# **CYSTINURIA**

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Dundee	NGS	Whole g	ene screen	SNVs, indels	SLC3A1, SLC7A9	56
Family men	nber testing		as indicated above			
Proforma re	quired?	YES	Renal Genet	tics Proforma (see centre web	site)	

#### 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	Whole ge screen		SNVs, indels	AGXT, APRT, ATP6VoA4, ATP6V1B1, BSND, CA2, CASR, CLCN5, CLCNKB, CLDN16, CLDN19, CYP24A1, FAM20A, GRHPR, HOGA1, HPRT1, KCNJ1, OCRL, PHEX, SLC12A1, SLC22A12, SLC2A9, SLC34A1, SLC34A3, SLC3A1, SLC4A1, SLC7A9, SLC9A3R1, STRADA, XDH	112
Family mer	nber testing		•		as indicated above	14
Proforma re	equired?	YES	Rena	I Genetics Pro	forma (see centre website)	·

# REFERRAL CRITERIA

• Nephrocalcinosis or nephrolithiasis where acquired causes have been excluded

- Clinical Genetics
- Nephrology

# POLYCYSTIC KIDNEY DISEASE, AUTOSOMAL DOMINANT

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Dundee	NGS	Whole ge screer		SNVs, indels Exon level CNV	PKD1, PKD2	56
Dundee	NGS	Whole go		SNVs, indels	AGT, ALG8, ALG9, ANKS6, CEP164, CEP83, COL4A1, DNAJB11, DZIP1L, GANAB, HNF1B, INVS, LRP5, MAPKBP1, NPHP1, NPHP3, NPHP4, PKD1, PKD2, PKHD1, PRKCSH, REN, SEC61B, SEC61A1, SEC63, TMEM67, TSC1, TSC2, TTC21B, UMOD, VHL, WDR19	112
Family member testing			as indicated above	14		
Proforma re	quired?	Yes	Ren	al Genetics Proform	na (see centre website)	

#### 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.
  - o Initial analysis of *PKD1* and *PKD2* then further analysis of the full cystic kidney panel if appropriate.
- The full cystic kidney disease full panel is recommended for individuals that meet the following criteria:
  - o 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
  - o 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 KIDNEY DISEASE, AUTOSOMAL RECESSIVE

#### **AVAILABLE TESTING**

Centre	Method	Sc	ope and rar	ige of test	Targets	TAT
Dundee	NGS	Whole ge	ne screen	SNVs, indels	PKHD1	56
Family men	nber testing			as ind	icated above	14
Proforma re	quired?	Yes	Renal Gen	etics Proforma (see	centre website)	

#### REFERRAL CRITERIA

- 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

- Clinical Genetics
- Fetal Medicine
- Nephrology
- Paediatrics

# POLYCYSTIC LIVER DISEASE

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS	Whole gen screen	se SNVs, indels	AGT, ALG8, ALG9, ANKS6, CEP164, CEP83, COL4A1, DNAJB11, DZIP1L, GANAB, HNF1B, INVS, LRP5, MAPKBP1, NPHP1, NPHP3, NPHP4, PKD1, PKD2, PKHD1, PRKCSH, REN, SEC61B, SEC61A1, SEC63, TMEM67, TSC1, TSC2, TTC21B, UMOD, VHL, WDR19	112
Family men	nber testing	·		as indicated above	14
Proforma re	quired?	Yes	Renal Genetics Prof	forma (see centre website)	

# REFERRAL CRITERIA

• Individuals with a suspected or established diagnosis of Polycystic Liver Disease based on imaging or pathology.

- Clinical Genetics
- Fetal Medicine
- Nephrology
- Paediatrics

#### PRIMARY HYPEROXALURIA

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Dundee	NGS	Whole ge	ne screen	SNVs, indels	AGXT, GPHPR, HOGA1	56
Family men	nber testing			as indi	cated above	14
Proforma re	na required? YES Renal Genetics Proforma (se		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 ran	ge of test	Targets	TAT
Dundee	NGS	Whole gene screen	SNVs, indels	NR <sub>3</sub> C <sub>2</sub> , SCNN <sub>1</sub> A, SCNN <sub>1</sub> B, SCNN <sub>1</sub> G	112
Family men	nber testing		as indi	cated above	14
Proforma re	quired?	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	Whole ge screen		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
Family men	nber testing			as indicated above	14
Proforma re	equired?	quired? YES Renal Genetics Proforma (see centre website)			

# REFERRAL CRITERIA

• Individuals with a suspected clinical diagnosis associated with the above genes

- Clinical Genetics
- Nephrology

# RENAL TUBULOPATHIES, RENAL TUBULAR ACIDOSIS

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS	Whole ger screen	ne SNVs, indels	AP2S1, AQP2, ATP1A1, ATP6V0A4, ATP6V1B1, AVPR2, BSND, CA2, CASR, CLCNKB, CLDN16, CLDN19, CTNS, CUL3, CYP24A1, FAH, GATM, GNA11, HNF1B, KCNJ1, KCNJ10, KLHL3, NR3C2, REN, SCNN1A, SCNN1B, SCNN1G, SLC12A1, SLC12A3, SLC22A12, SLC2A9, SLC4A1, SLC4A4, SLC5A2, TRPM6, UMOD, WNK4	112
Family men	nber testing			as indicated above	14
Proforma re	quired?	YES	Renal Genetics Prof	forma (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
  - o 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
  - Nephrogenic diabetes insipidus, OR
  - o 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 range of test		Targets	TAT	
Dundee	NGS	Whole ger screen	ne SNVs, 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, TNS1, TP53RK, TRPC6,, WDR73, WT1	112	
Family men	nber testing			as indicated above	14	
Proforma re	equired?	YES	Renal Genetics 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	Whole gene screen		SNVs, indels	ANKS6, CEP164, CEP83, GATM, HNF1B, INVS, MUC1, NPHP1, NPHP3, NPHP4, REN, TMEM67, TTC21B, UMO, WDR19	112
Family me	mber testing			as inc	licated above	14
Proforma re	equired?	YES	Renal Gene	tics Proforma (see	e 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	Sanger	Targeted screen	SNV	ADRB2 p.(Gly16Arg)	28
Proforma re	quired?	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	nge of test	Targets	TAT
Aberdeen Dundee Edinburgh Glasgow	ARMS	Targeted screen	SNVs, indels	Common mutations	28 Prenatal 3
Glasgow	ARMS	Targeted screen	SNVs, indels	CFTR newborn screening (p.508del, p.G542*, p.G551D, c.469+1G>T common mutations)	7
Edinburgh	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV	CFTR	56
Family mem	ber 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 >4omM with sufficient sweat obtained; >3omM 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
  - o 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
  - o 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 ran	ge of test	Targets	TAT
Edinburgh	HHT-NGS	Whole gene screen	SNVs, indels, CNV**	ACVRL1**, ENG**, EPHB4, GDF2, RASA1, SMAD4**	56
Edinburgh	PPH-NGS	Whole gene screen	SNVs, indels, CNV**	ACVRL1**, ATP13A3, BMPR2**, CAV1, GDF2, EIF2AK4, ENG**, KCNK3, SMAD9, SOX17, TBX4	56
Family men	nber testing		as ii	ndicated 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
  - o 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:
  - 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.

- Clinical Genetics
- Respiratory

# PRIMARY CILIARY DYSKINESIA

#### **AVAILABLE TESTING**

Centre	Method	Scope and	range of test	Targets	TAT
Glasgow	NGS	Whole gene screen	SNVs, indels	ARMC4, C21ORF59, CCDC39, CCDC40, CCDC65, CCDC103, CCDC114, CCDC151, CCNO, DNAAF1, DNAAF2, DNAAF3, DNAAF5, DNAH5, DNAH11, DNAI1, DNAI2, DNAL1, DRC1, DYX1C1, GAS8, LRRC6, MCIDAS, RPGR, RSPH1, RSPH4A, RSPH9, SPAG1, ZMYND10	112
Family men	nber testing			as indicated above	14
Proforma re	quired?	NO			

#### REFERRAL CRITERIA

- Neonate at least one of the following:
  - o Situs inversus plus lower airway or nasal symptoms
  - o Persistent respiratory distress where other causes have been excluded
  - o Persistent rhinorrhoea and cough distress where other causes have been excluded
  - Sibling with PCD
- Childhood at least one of the following:
  - Persistent lifelong wet cough (cystic fibrosis excluded)
  - Unexplained bronchiectasis (cystic fibrosis excluded)
  - o Serious otitis media in association with recurrent lower and upper airway symptoms
- Adults
  - o Symptoms as above since, often associated with infertility or subfertility

- Clinical Genetics
- Paediatrics
- Respiratory Medicine

# SURFACTANT METABOLISM DYSFUNCTION

#### **AVAILABLE TESTING**

Centre	Method	Scope and rang	ge of test	Targets	TAT
Glasgow	NGS	Whole gene screen	SNVs, indels	ABCA3, NKX2-1, SFTPB, SFTPC	56
Family men	nber 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	Sc	ope and rang	e of test	Targets	TAT
Dundee	Sanger	Whole g	ene screen	SNVs, indels	TGM5	56
Family men	nber testing			as indi	cated above	14
Proforma re	quired?	YES	Skin disorde	ers proforma (see	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	Scope and range of test		Targets	TAT
Dundee	NGS	Whole ge screen		SNVs, indels	ABCA12, ALDH3A2, ALOX12B, ALOXE3, CERS3, CYP4F22, NIPAL4, PNPLA1, SLC27A4, ST14, STS, SULT2B1, TGM	56
Family mem	ber testing				as indicated above	14
Proforma re	quired?	YES	Skin disorders proforr		ma (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	S	cope and ra	nge of test	Targets	TAT
Dundee	Sanger MLPA	Whole ge	ne screen	SNVs, indels Exon level CNV	FLCN	56
Family men	nber testing			as indica	ited above	14
Proforma re	quired?	NO				

#### REFERRAL CRITERIA

- Individuals with either:
  - o 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	Whole ge	ne screen	SNVs, indels	KRT1, KRT10	56
Family men	nber testing			as in	dicated above	14
Proforma re	quired?	YES	Skin disor	ders proforma (se	e 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
  - o Skin biopsy recommended if mosaic form suspected (epidermolytic epidermal naevus)

- Clinical Genetics
- Dermatology

#### **ECTODERMAL DYSPLASIA**

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test		ange of test	Targets	TAT
Dundee	NGS	Whole ge screer		SNVs, 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
Family men	nber testing			as indicated above		14
Proforma re	equired?	YES	Skin	disorders proform	na (see centre website)	

#### REFERRAL CRITERIA

- Any individual with a clinical diagnosis of ectodermal dysplasia with one or more of the following:
  - o abnormality of hair (hypotrichosis, sparse hair, sparse/missing eyebrows)
  - o 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's disease, 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 gene screen	SNVs, indels	KRT <sub>5</sub> , KRT <sub>14</sub>	56
Dundee	NGS	Whole gene screen	SNVs, indels	COL17A1, COL7A1, DSP, DST, EXPH5, FERMT1, ITGA3, ITGA6, ITGB4, JUP, KRT14, KRT5, LAMA3, LAMB3, LAMC2, PKP1, PLEC, TGM5	112
Family men	nber testing			as indicated above	14
Proforma required? YES Skin disorders		Skin disorders p	roforma (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
  - o Dowling-Degos Syndrome Sanger sequencing for *KRT* 5
  - o 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**

Centre	Method	Scope and range of test			Targets	TAT
Dundee	Sanger	Whole ge	ne screen	SNVs, indels	KRT1, KRT9	56
Family men	nber testing			as in	dicated above	14
Proforma re	quired?	YES	Skin Disor	ders Proforma (se	ee 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 ran	ge of test	Targets	TAT
Dundee	Sanger	Whole gene screen	SNVs, indels	TGFBR1	56
Family men	nber testing		as ind	icated above	14
Proforma re	quired?	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 ran	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	14
Proforma re	quired?	NO			

#### REFERRAL CRITERIA

- Any individual with a clinical presentation consistent with the condition:
  - o Focal palmoplantar hyperkeratosis
  - o Palmoplantar keratoderma
  - Autosomal dominant

- Clinical Genetics
- Dermatology

# **GLOMUVENOUS MALFORMATIONS**

#### **AVAILABLE TESTING**

Centre	Method	Scope and rar	ige of test	Targets	TAT
Dundee	NGS	Targeted screen	SNVs, indels	GLMN (exons 2, 3, 6, 8, 12, 13)	56
Family mer	nber testing		as ind	icated above	14
Proforma re	quired?	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	Whole ger screen	SNVs, indels	APCDD1, ATP7A, CDH3, CDSN, DSC3, DSG4, EDAR, ERCC2, GJB2, GJB6, HOXC13, HR, JUP, KRT71, KRT74, KRT81, KRT83, KRT85, KRT86, LIPH, LPAR6, MBTPS2, RIPK4, SNRPE, SPINK5, VDR	112
Family men	nber testing			as indicated above	14
Proforma re	quired?	YES	Skin Disorders Prof	orma (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		inge of test	Targets	TAT
Dundee	NGS	Whole ge screen		SNVs, indels	AAGAB, ABCA12, ALOX12B, ALOXE3, AQP5, CARD14, CAST, CERS3, CLDN1, CYP4F22, DSC2, DSG1, DSP, ENPP1, FLG, GJA1, GJB2, GJB3, GJB4, GJB6, JUP, KDSR, KRT1, KRT10, KRT14, KRT16, KRT17, KRT2, KRT6A, KRT6B, KRT6C, KRT9, LOR, MSMO1, NIPAL4, PIGL, PNPLA1, RSPO1, RHBDF2, SERPINB7, SLC27A4, SLURP1, SMARCAD1, SNAP29, SPINK5, ST14, STS, SULT2B1, TAT, TGM1, TRPV3	112
Family men	nber testing				as indicated above	14
Proforma re	quired?	YES	Skin	disorders profor	ma (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> (p.Arg501*; c.2282_2285delCAGT, p.Arg2447*; p.Ser3247*)	28
Proforma re	quired?	YES	Skin disc	orders 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
  - o Keratosis pilaris
  - o Hyperlinear pals and soles
  - o Atopic eczema

- Dermatology
- Clinical Genetics

# **LEGIUS SYNDROME**

#### **AVAILABLE TESTING**

Centre	Method	Scope and ra	ange of test	Targets	TAT
Dundee	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV	SPRED1	56
Proforma re	equired?	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
  - o Axillary or inquinal 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 men	nber testing		as in	dicated above	14
Proforma re	equired?	NO			

#### REFERRAL CRITERIA

- Any individual with a clinical presentation consistent with the condition:
  - o Small, multifocal cutaneous and/or mucosal bluish-purple vascular malformations
  - Early onset (mostly at birth)
  - o Slow blood flow on Doppler ultrasound
  - Elevated D-dimer concentration

- Dermatology
- Clinical Genetics

# **PACHYONYCHIA CONGENITA**

#### **AVAILABLE TESTING**

Centre	Method	Scope and rar	nge of test	Targets	TAT
Dundee	Sanger	Targeted screen	SNVs, indels	KT6A (ex1&7)  KRT6B (ex1&7)  KRT6C (ex1&7)  KRT16 (ex1,6,7&8)  KRT17 (ex1,6&7)	56
Family member testing			as	indicated above	14
Proforma required?		NO			

# REFERRAL CRITERIA

- Any individual with a clinical presentation consistent with the condition:
  - o Plantar keratoderma including callus with underlying blisters
  - o Plantar pain
  - o 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	Whole ger screen	ne 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 member testing				as indicated above	14		
Proforma required?		YES	YES Skin disorders proforma (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
  - o Pachyonychia congenital phenotype
  - o Punctate keratoderma
  - o Striate keratoderma with woolly hair
  - o Keratoderma with deafness
  - o 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		range of test	Targets	TAT
Dundee	NGS	Whole scre	_	SNVs, indels	ABCC6, ADA2, ADAMTS2, AGPAT2, AIRE, ANTXR2, ATP6VoA2, ATP7A, ATP7B, CARD11, CARD14, CARD9, COL1A1, COL1A2, COL3A1, COL4A3, COL4A4, COL4A5, COL5A1, COL5A2, CSTA, CYBB, DCLRE1C, DOCK8, EFEMP2, EGFR, ELN, FBLN5, FDPS, FGF23, FLG, FLT4, FMO3, FOXC2, GALNT3, GGCX, GJA1, GJB3, IKBKG, IL1RN, GJB4, IL36RN, KIT, KRT1, KRT10, LYST, MVD, NCSTN, NLRP1, NLRP3, NOD2, NSDHL, OSMR, PSEN1, PSENEN, RAG1, RAG2, SAMHD1, SH3PXD2B, SLC39A4, STAT3, TMEM173, TREX1	112
Family member testing					as indicated above	14
Proforma required?		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
  - o Includes autoinflammatory disease (e.g. early onset urticaria, recurrent febrile erythemas), infantile pustular psoriasis, likely genetic forms of pityriasis rubra pilaris

- Clinical Genetics
- Dermatology
- Rheumatology

# SUPERFICIAL EPIDERMOLYTIC ICHTHYOSIS (SEI) (previously known as ICHTHYOSIS BULLOSA of SIEMENS)

#### **AVAILABLE TESTING**

Centre	Method	Scope and rang	e of test	Targets	TAT
Dundee	Sanger	Whole gene screen	SNVs, indels	KRT2	56
Family member testing			as indi	cated above	14
Proforma required?		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	Whole gene screen		SNVs, indels	ACVRL1, ADAMTS13, ALAS2, ATM, ATR, CCBE1, ENG, EPHB4, F12, FECH, FLT4, FOXC2, GLMN, KRIT1, PIK3CA, PIK3R2, PTEN, RASA1, SCN9A, SMAD4, SOX18, TEK, TMEM173	112	
Family member testing			as indicated above			14	
Proforma required?		YES	Skin disorders proforma (see centre website)				

#### REFERRAL CRITERIA

• Any individual with a vascular skin disorder with a likely germline genetic cause

- Clinical Genetics
- Dermatology

# X-LINKED ICHTHYOSIS

#### **AVAILABLE TESTING**

Centre	Method	Scope and range of test			Targets	TAT
Dundee	Sanger MLPA	Whole gene screen		SNVs, indels Exon level CNV	STS	56
Family member testing			as indicated above			
Proforma required?		YES	Skin disorders proforma (see centre website)			

# REFERRAL CRITERIA

- Any individual with a clinical presentation consistent with the condition:
  - o Steroid sulfatase (STS) enzyme deficiency
  - o Dry skin
  - o Hyperkeratosis
  - o Hypohidrosis
  - o Ichthyosis

- Dermatology
- Clinical Genetics