



Scottish Strategic Network for Genomic Medicine

Genomic Test Directory

Rare & 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 to work as a formal network arrangement, to deliver an equitable, high quality genetic testing service for Scotland. All laboratories are accredited by United Kingdom Accreditation Service (UKAS) in accordance with the recognized ISO 15189:2012 standard.

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

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

PURPOSE OF DOCUMENT

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

This document will be reviewed annually.





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

genetics

• Dundee (NHS Tayside)

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

Hospital, Dundee DD1 9SY Email address: Tay.esrg@nhs.scot

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

Z/Genetics/PROD_295540/index.htm

• Edinburgh (NHS Lothian)

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

Crewe Road, Edinburgh, EH4 2XU

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

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

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

toryServices/Pages/default.aspx

Glasgow (NHS Greater Glasgow & Clyde)

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

University Hospital, Glasgow G51 4TF

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

Website:

<u>Laboratory Genetics - NHSGGC</u>





TEST REQUESTING

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

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

SAMPLE REQUIREMENTS

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

Other sample types may be required for some services including:

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

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





TESTING METHODOLOGY

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

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

SCOPE AND RANGE OF TEST

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

The scope of testing includes:

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

The types of variants detected includes:

- Small sequence variants
 - Single nucleotide variants (SNVs)
 - Insertions / deletions (indels)
- Copy number variants (CNVs)
 - 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 range of test			Targets	TAT
Aberdeen	Sanger	Whole	gene screen	SNVs, indels	KCNJ2, KCNJ5	56
Family me	member testing as indicated above					
Proforma	required?	YES	YES Cardiac Arrhythmia Proforma (see centre website)			

Referral criteria

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

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





ARRHYTHMIA PANEL

Available testing

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

Referral criteria

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

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





ARRHYTHMOGENIC CARDIOMYOPATHY

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS		ole gene creen	SNVs, indels	PKP2, DSG2, DSC2, DSP, SCN5A, ABCC9, DES, HCN4, JUP, LMNA, PLN, RYR2, TGFB3, TMEM43	112
Family mer	nber testing		as indicate		ed above	14
Proforma re	equired?	YES	Cardiac Ar	rhythmia Proform	a (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		ole gene creen	SNVs, indels	SCN5A, ABCC9, GJA1, GJA5, HCN4, KCNA5, KCNE5, NPPA, SCN2B, SCN4B	56
Family mer	nber testing		as indicated above			
Proforma re	equired?	YES	YES Cardiac Arrhythmia Proforma (see centre website)			

Referral criteria

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

- Cardiologist with expertise in ICCClinical Genetics





BARTH SYNDROME

Available testing

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

Referral criteria

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

- Cardiology
- Clinical Genetics
- Paediatrics





BRUGADA SYNDROME AND SODIUM CHANNEL DISEASE

Available testing

Centre	Method	Sco	pe and ra	nge of test	Targets	TAT	
Aberdeen	NGS		e gene een	SNVs, indels	SCN5A, CACNA1C, CACNA2D1, CACNB2, GPD1L, HCN4, KCND3, KCNE3, KCNE5, KCNJ8, RANGRF, SCN1B, SCN2B, SCN3B, SCN10A, SLMAP, TRPM4	112	
Family r	member ting			as indica	ated above	14	
Proforma re	equired?	YES	ES Cardiac Arrhythmia Proforma (see centre website)				

Referral criteria

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

- Cardiologist with expertise in ICC
- Clinical Genetics





CATECHOLAMINERGIC POLYMORPHIC VENTRICULAR TACHYCARDIA (CPVT)

Available testing

Centre	Method	Scope and range of test			Targets	TAT	
Aberdeen	NGS	Whole gene screen SNVs, ind		SNVs, indels	RYR2, CALM1, CALM2, CASQ2, DPP6, TRDN	56	
Family men	Family member as indica			as indicat	ed above	14	
testing							
Proforma required? YES Cardiac Arrhythmia Proforma (see			ma (see centre website)				

Referral criteria

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

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





DILATED CARDIOMYOPATHY (DCM)

Available testing

Centre	Method	Scope and ra	nge of test	Targets	TAT
Edinburgh	NGS	Whole gene 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 r		as indic		ated above	14
Proforma re	equired?	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 me	mber testing			as indicated a	bove	14
Proforma re	equired?	YES	Cardiac Arrh	ythmia Proforma (s	ee centre website)	

Referral criteria

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

- Cardiologist with expertise in ICC
- Clinical Genetics





HYPERTROPHIC CARDIOMYOPATHY (HCM)

Available testing

	U				
Centre	Method	Scope and ran	nge 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 r			as ir	dicated above	14
Proforma re	equired?	NO			

Centre	Method	Scope and ran	ige of test	Targets	TAT
Edinburgh	Sanger	Whole gene screen	SNVs, indels	Familial amyloid polyneuropathy: TTR	56
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 Arrhythmia Proforma (see centre website)				

Referral criteria

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

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





MARFAN SYNDROME

Available testing

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

Referral criteria

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

- Cardiologist with expertise in ICC
- Clinical Genetics





PAEDIATRIC CARDIOMYOPATHY

Available testing

Centre	Method	Scope and	range of test	Targets	
Aberdeen	NGS	Whole gene screen	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, NDUFA5, NDUFA5, NDUFAF4, NDUFA5, NDUFS4, NDUFS6, NDUFS7, NDUFS8, NDUFS3, NDUFS4, NDUFS6, NDUFS7, NDUFS8, NDUFV1, NDUFV2, NEXN, NF1, NKX2-5, NONO, NRAS, NUBPL, PCCA, PCCB, PDLIM3, PKP2, PLN, PNPLA2, PPA2, PPCS, PPP1CB, PPP1R13L, PRKAG2, PTPN11, RAF1, RBM20, RIT1, RYR2, SCN5A, SCO1, SCO2, SDHA, SDHAF1, SDHD, SGCD, SHOC2, SLC22A5, SLC25A20, SLC25A4, SOS1, SOS2, SURF1, TAZ, TBX5, TMEM126B, TMEM43, TMEM70, TNNC1, TNNI3, TNNI3K, TNNT2, TPM1, TSFM, TTN, TTR, VCL	112
Family mer testing	Family member			as indicated above	14
Proforma required?		NO			

Referral criteria

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

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





SHORT QT SYNDROME

Available testing

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

Referral criteria

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

- Cardiologist with expertise in ICC
- Clinical Genetics





THORACIC AORTIC ANEURYSM & DISSECTION (TAAD)

Available testing

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

Referral criteria

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

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





CHROMOSOME BREAKAGE

ATAXIA TELANGIECTASIA (& AT-LIKE)

Available testing

Centre	Method	Scope and ra	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 range of test		Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	APTX	56
Family member as ind		as indicate	d above	14	
Proforma re	equired?	NO			

Referral criteria

Clinical phenotype suggestive of ataxia with oculomotor apraxia & hypoalbuminemia

Requesting specialties

- Clinical Genetics
- Haematology

BLOOM SYNDROME

Available testing

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

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

Referral criteria

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

- 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 i	member ting	as indicated a		ated above	14
Proforma re	equired?	NO			

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Haematology

COCKAYNE SYNDROME

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels	ERCC6, ERCC8	56
_	Family member as indicates testing		ated above	14	
Proforma re	equired?	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 member as i		as indic	ated above	14	
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Haematology





FANCONI ANAEMIA

Available testing

Centre	Method	Scope and range	of test	Targets	TAT
Aberdeen	Chromosome 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, TOP3A, UBE2T	112
Family member testing			as indica	ated above	14
Proforma required?		NO			

^{*5}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	112
,	member ting			as indicated above	14
Proforma re	equired?	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	I I I I I I I I I I I I I I I I I I I		56
	Family member as indicates testing		ated above	14	
Proforma required?		NO			

Referral criteria

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

Requesting specialties

Clinical Genetics





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	DNMT3B, ZBTB24, CDCA7, HELLS	56
		ted above	14		
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





LIG4 SYNDROME

Available testing

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

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Haematology
- Immunology

MEIER-GORLIN SYNDROME

Available testing

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

Referral criteria

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

Requesting specialties

Clinical Genetics





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 r		r as indic		ated above	14
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





NIJMEGEN BREAKAGE SYNDROME (& NBS-LIKE)

Available testing

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

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





ROBERTS-SC PHOCOMELIA SYNDROME

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Karyotype	Whole genome screen	Aneuploidy	Genome wide	28
Aberdeen	Sanger	Whole gene screen	SNVs, indels	ESCO2	56
	member ting	as indicated above			14
Proforma required?		NO			•

Referral criteria

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

Requesting specialties

Clinical Genetics

ROTHMUND-THOMSON / RAPADILINO / BALLER-GEROLD

Available testing

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

Referral criteria

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

- Clinical Genetics
- Dermatology





SECKEL SYNDROME

Available testing

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

Referral criteria

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

Requesting specialties

Clinical Genetics

TAR SYNDROME

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	RBM8A	56
Family r			as indic	ated above	14
Proforma re	equired?	NO			

Referral criteria

• Clinical phenotype suggestive of Thrombocytopenia-absent radius syndrome

- Clinical Genetics
- Haematology





TOWNES-BROCKS SYNDROME

Available testing

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

Referral criteria

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

Requesting specialties

Clinical Genetics

TRICOTHIODYSTROPHY

Available testing

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

Referral criteria

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

- Clinical Genetics
- Dermatology





ULNAR-MAMMARY SYNDROME

Available testing

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

Referral criteria

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

Requesting specialties

Clinical Genetics

WARSAW BREAKAGE SYNDROME

Available testing

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

Referral criteria

• Clinical phenotype suggestive of Warsaw Breakage Syndrome

Requesting specialties

Clinical Genetics





WERNER SYNDROME

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels	WRN	56
_	Family member as indicated testing		d above	14	
Proforma re	equired?	NO			

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Dermatology

XERODERMA PIGMENTOSUM

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels	XPA, XPC, ERCC1, ERCC3, ERCC4, ERCC5, DDB2, POLH	56
Family r	nember ting		as indica	ted above	14
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Dermatology





CONNECTIVE TISSUE DISORDERS

CONNECTIVE TISSUE

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	NGS	Whole gene screen	SNVs, indels	ABCC6, ACTA2, ACVR1, ADAMTS2, ALPL, ATP6V0A2, B3GALT6, B4GALT7, BMP1, CBS, CHST14, COL1A1, 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 r		as indicated above		14	
Proforma required? NO					

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

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	NGS	Whole gene screen	SNVs, indels	Hypophosphatasia: ALPL	112
Family member testing			as indicated above	14	
Proforma required? NO				•	

Referral criteria

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

Individuals who have characteristic features of Pseudoxanthoma elasticum:

- Papules or plaques on the skin of the neck and/or flexural creases (antecubital fossae, axillae, groin, or popliteal fossae) and/or calcified dystrophic elastic fibres on biopsied skin using a von Kossa or similar stain) AND/OR
- Retinal finding (angioid streaks, peau d'orange, or choroidal vascularization)





Individuals who have characteristic features of Hypophosphotasia:

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

and

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

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





EHLERS DANLOS SYNDROME

Available testing

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

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

Referral criteria

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

Requesting specialties

Clinical Genetics





STICKLER SYNDROME / CLEFT PALATE

Available testing

STICKLER SYNDROME

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

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

Cleft palate

Centre	Method	Scope ar	nd range of	test	Targets	TAT
Edinburgh	NGS	Whole gene screen	SNVs, indels	ARH BMF CI COL. COL. EB E FG GJ MAP MK NI PIG R SC SIX3, S STAI	Cleft: ACTB, ACTG1, AMER1, ANKRD11, HGAP29, ARHGAP31, ASXL1, B3GLCT, BCOR, P2, C2CD3, C5orf42, CC2D2A, CDH1, CDKN1C, HD7, CHRNG, CHST14, COL11A1, COL11A2, COL2A1, COLEC10, EC11, CTCF, CTNND1, DHCR7, DHODH, A, DOCK6, DVL1, DVL3, DYNC2H1, DYNC2LI1, P, EDNRA, EFNB1, EFTUD2, EIF2S3, EIF4A3, OGT, EPG5, ESCO2, EYA1, FAM20C, FGD1, GFR1, FGFR2, FLNA, FLNB, FOXC2, FRAS1, A1, GLI3, GPC3, GRHL3, HDAC8, HYLS1, ICK, ET140, IFT172, IFT80, IMPAD1, IRF6, KAT6A, NJ2, KDM6A, KIAA0586, KIF1BP, KIF7, KMT2D, GSK7, MAPRE2, MASP1, MBTPS2, MEIS2, MID1, GS1, MSX1, MYMK, NECTIN1, NEDD4L, NEK1, MPBL, NOTCH1, OFD1, PAX3, PHF8, PIEZO2, MN, PIGV, POLR1C, POLR1D, PORCN, PTCH1, BM10, ROR2, RPL5, RPS26, SALL4, SATB2, CARF2, SEPT9 (SEPTIN9), SF3B4, SHH, SIX1, SIX5, SKI, SLC26A2, SMAD3, SMAD4, SMC1A, MC3, SMS, SNRPB, SON, SOX9, SPECC1L, MBP, TBX22, TCOF1, TCTN3, TELO2, TFAP2A, DS, TGFB3, TGFBR1, TGFBR2, TMCO1, TP63, TRAPPC9, TRIM37, TUBB, TXNL4A, USP9X, WNT5A, XYLT1, ZEB2, ZIC2, ZIC3, ZSWIM6	112
Family n	nember				s indicated above	14
testing		NO				
Proforma re	equirea?	NO				

Gene list from panel app Clefting panelv2.2

https://nhsgms-panelapp.genomicsengland.co.uk/panels/81/v2.2

Centre	Method	Scope and range of test		est	Targets	TAT
Edinburgh	NGS	Whole gene screen	SNVs, indels		Van der Woude syndrome: IRF6	112
Family member testing				а	as indicated above	14
Proforma re	quired?	NO				





Referral criteria

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

Requesting specialties

• Clinical Genetics





DEVELOPMENTAL DISORDERS

ANEUPLOIDY SCREENING - NON-INVASIVE PRENATAL TESTING (NIPT)

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS (genome- wide)	Targeted screen	Aneuploidy	Chromosomes 13, 18, 21	7
Proforma required?	YES	NIPT request form	1		

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 CNV screen		Whole genome	20
Proforma required?		NO			

Referral criteria

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

- Clinical Genetics
- Obstetrics





ANGELMAN SYNDROME

Available testing

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

Referral criteria

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

- Clinical Genetics
- Paediatrics





BECKWITH WIEDEMANN SYNDROME

Available testing

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

Referral criteria

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

- Clinical Genetics
- Paediatrics





CHARGE SYNDROME

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	Sanger	Whole gene screen SNVs, indels Exon level CNV		CHD7	56
Family mem	ber testing	as indicated above			
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
 - Characteristic ear malformations
 - o Tracheoesophageal fistula or oesophageal atresia
 - o Cardiovascular malformation
 - o Genital hypoplasia

Requesting specialties

• Clinical Genetics





CONGENITAL ABNORMALITIES, MULTIPLE

Available testing

Centre	Method	Scope and range of test		Targets	TAT	
Aberdeen Dundee	Microarray	Whole genome	Structural variants	Whole genome	14	
Edinburgh Glasgow	Karyotype	screen	CNV	3		
Proforma required?		NO				

Referral criteria

• Multiple congenital malformations

- Clinical Genetics
- Paediatrics





CORNELIA DE LANGE SYNDROME (CdLS) and CdLS-LIKE DISORDERS

Available testing

Centre	Method	Scope and range of test		Targets	TAT		
Edinburgh	NGS	Whole gene screen	SNVs, indels	NIPBL*, SMC1A, SMC3, HDAC8, RAD21, ANKRD11, KMT2A, AFF4, NAA10, BRD4, PUF60	56		
Family membe	Family member testing		as indicated above				
Proforma require	ed?	NO					

^{*} MLPA analysis of NIPBL is available for patients presenting with "classical" CdLS. Please contact the laboratory for details.

Referral criteria

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

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

Clinical Genetics





DEVELOPMENTAL DELAY

Available testing

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

Referral criteria

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

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





DEVELOPMENTAL DISORDERS

Available testing

Centre	Method		Scope and rai	nge of test	Targets	TAT	
Edinburgh	NGS	Wh	ole gene screen	SNVs, indels	DDG2P*	112	
Family member testing			as indicated above				
Proforma required?		NO	DECIPHER entry required, including HPO terms and growth parameters. Please provide the DECIPHER ID in the referral documents.				

^{*} Gene list available at https://www.ebi.ac.uk/gene2phenotype. Only genes in the moderate, strong, and definitive categories are analysed.

Referral criteria

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

Requesting specialties

• Clinical Genetics





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 *Contact lab to discuss TAT
Proforma required?		NO			

Referral criteria

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

- Cardiology
- Clinical Genetics
- Obstetrics
- Paediatrics





DISORDERS OF SEXUAL DEVELOPMENT (DSD)

Available testing

Centre	Method	Scope and ran	ge of test	Targets	TAT	
Aberdeen Dundee 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 ind	icated above (Glasgow)	14	
Proforma required?		YES DSD referral form (see centre website)				

Referral criteria

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

- Clinical Genetics
- Endocrinology
- Paediatrics





FRAGILE X

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen Edinburgh Glasgow	PCR & TPPCR	Targeted screen	Triplet repeat expansion	FMR1	28
Proforma requ	ired?	NO			

Referral criteria

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

Family history of Fragile X Requesting specialties

- Clinical Genetics
- Paediatrics





INFERTILITY, MALES

Available testing

Centre	Method	Scope and r	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	Common CFTR pathogenic variants	28
Dundee Edinburgh Glasgow	PCR	Targeted screen	Y chromosome markers	AZFa, AZFb, AZFc	28
Proforma required?		NO			

Referral criteria

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

- Clinical Genetics
- Fertility specialist





INFERTILITY, FEMALES

Available testing

Centre	Method	Scope ar	nd range of test	Targets	TAT
Aberdeen Dundee Edinburgh Glasgow	Karyotype	Whole genome screen	Structural variants, CNV	Whole genome	28
Aberdeen Edinburgh Glasgow	PCR	Targeted screen	Triplet repeat expansion	FMR1	28
Proforma required?		NO			

Referral criteria

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

- Clinical Genetics
- Fertility specialist





KLINEFELTER SYNDROME

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen Dundee Glasgow	Karyotype	Whole genome	CNV	Whole geneme	20
	Microarray	screen	CINV	Whole genome	28
Edinburgh	Karyotype	Whole genome screen	CNV	Whole genome	28
Proforma requ	ired?	NO			

Referral criteria

- Primary hypogonadism
- Cryptorchidism
- Gynaecomastia
- Infertility

- Clinical Genetics
- Endocrinology
- Fertility clinics





MICRODELETION / MICRODUPLICATION SYNDROMES

Available testing

Centre	Method	Scope and ran	ge of test	Targets	TAT
Aberdeen Dundee Edinburgh Glasgow	Microarray	Whole genome screen	CNV	Whole genome	28
Proforma require	ed?	NO			

Referral criteria

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

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

- Clinical Genetics
- Paediatrics





HYDATIDIFORM MOLE

Available testing

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

Referral criteria

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

Genetic testing in isolation

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

Requesting specialties

Pathology





PRADER-WILLI SYNDROME

Available testing

Centre	Method	Scope	and range of test	Targets	TAT
Glasgow	MLPA	Targeted screen	CNV 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
 - 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	28	
Proforma requ	ired?	NO				

Referral criteria

• Tissue from 3rd or subsequent consecutive miscarriage

- Fetal Medicine
- Pathology
- Gynaecology





SILVER-RUSSELL SYNDROME

Available testing

Centre	Method	Scope a	nd range of test	Targets	TAT
Glasgow	MLPA	Targeted screen	CNV Methylation abnormalities	11p15 region	28
Glasgow	PCR	Targeted screen	STRs	Microsatellite markers	28
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Paediatrics





SMITH-LEMLI-OPITZ

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	Sanger	Whole gene screen	SNVs, indels	DHCR7	56
	member ting		as ind	icated above	14
Proforma re	equired?	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 re	equired?	NO			

Referral criteria

- Prenatal testing is available for:
 - 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	Methylation analysis	AR CAG trinucleotide repeat	28
Proforma re	equired?	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 (targeted panel)	Whol	e gene screen	SNVs, indels	GNAS	56
Family memb	er testing	sting as indicated above			pove	14
Proforma required? YES Endocrine disorders proforma (see centre we			e website)			

Referral criteria

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

- Clinical Genetics
- Endocrinology





ANDROGEN INSENSITIVITY SYNDROME

Available testing

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

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Endocrinology

ASYMPTOMATIC FASTING HYPERGLYCAEMIA

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	Sanger or NGS	Whole gene screen	SNVs, indels, exon level CNV	GCK	56
Family mer	nber testing		as indicated ab	oove	14
Proforma requ	ired?	NO			

Referral criteria

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

- Clinical Genetics
- Endocrinology
- Obstetrics
- Paediatrics





CARNEY COMPLEX

Available testing

Centre	Method	5	Scope and rang	ge of test	Targets	TAT
Dundee	NGS (targeted panel)	Whole	e gene screen	SNVs, indels, exon level CNV	PRKAR1A	56
Family member testing				as indica	ated above	14
Proforma required? YES		Endocrine disorders proforma ((see website)		

Referral criteria

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

- Clinical Genetics
- Dermatology
- Endocrinology





CONGENITAL HYPERINSULINISM

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS (clinical exome)	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 member testing		as in	dicated above	14	
Proforma	a required?	NO			

Referral criteria

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

- Clinical Genetics
- Endocrinology
- Paediatrics





CONGENITAL HYPOTHYROIDISM

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS (clinical exome)	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 member as in testing		dicated 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 range of test		Targets	TAT
Dundee	Sanger	Whole gene SNVs, indels		AQP2, AVPR2	56
Family member testing			as ind	icated above	14
Proforma red	Proforma required? NO				

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Endocrinology

CONGENITAL OVERGROWTH DISORDERS

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS (clinical exome)	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 member 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	Scope and range of test		Targets	TAT
Dundee	NGS (targeted panel)	Who gene scree	Э	SNVs, indels, exon level CNV (selected genes)	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	equired?	YES	Enc	locrine disorders	s proforma (see centre website)	

Referral criteria

- Primary hyperparathyroidism (unexplained hypercalcaemia with PTH high or in the upper normal range, and calcium clearance: creatinine clearance ratio > 0.02) which meets ONE of the criteria below:
 - o Presenting before the age of 35, OR
 - 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)
 - Multiple Endorine Neoplasia Type 1 & Type 4
 - Multiple Endocrine Neoplasia Type 2A
 - O Hyperparathyroidism-Jaw Tumour Syndrome/Parathyroid carcinoma

- Clinical Genetics
- Endocrinology





FAMILIAL HYPOCALCIURIC HYPERCALCAEMIA

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Dundee	NGS(targeted panel)	Whole	O	SNVs, indels, exon level CNV (selected genes)	AP2S1, CASR, CDC73, CDKN1B, GCM2, GNA11, MEN1, RET (exons 5, 8, 10, 11, 13-16)	56
Family member testing				as indi	cated above	14
Proforma required?		YES	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		ange of test	Targets	TAT
Dundee	NGS (targeted panel)		e gene een	SNVs, indels, exon level CNV (selected genes)	AIRE, CASR, GATA3, GCM2, GNA11, PTH, TBCE	56
Family me	mber testing			as	indicated above	14
Proforma re	equired?	YES	YES Endocrine disorders pr		oforma (see centre website)	

Referral criteria

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

- Clinical Genetics
- Endocrinology





FAMILIAL ISOLATED PITUITARY ADENOMA

Available testing

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

Referral criteria

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

- Clinical Genetics
- Endocrinology





FAMILIAL NEUROHYPOPHYSEAL DIABETES INSIPIDUS

Available testing

Centre	Method	Scope and	d range of test	Targets	TAT
Dundee	Sanger	Whole gene screen	SNVs, indels	AVP	56
Family member testing			as indicat	ed above	14
Proforma re	equired?	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	Proforma required? 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 (targeted panel)	Whole	e gene een	SNVs, indels Exon level CNV	CDC73	56
_	member ting			as indicated a	above	14
Proforma	required?	YES	/ES 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 (clinical exome)	Whole gene screen	SNVs, indels	ALB, SECISBP2, SLC16A2, THRA, THRB, TSHR, TTR	112
	member ting		as indic	ated above	14
Proforma required?		NO			

Referral criteria

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

- Clinical Genetics
- Endocrinology





HYPOGONADOTROPIC HYPOGONADISM

Available testing

Centre	Method	Scope and range of test		ge of test	Targets	TAT
Glasgow	NGS	Whole	gene screen	SNVs, indels	ANOS1, CHD7, CUL4B, FEZF1, FGF8, FGFR1, FSHB, GNRH1, GNRHR, KISS1R, NR0B1, PROK2, PROK2R, SEMA3E, SOX2, SOX10, SPRY4, TAC3, TACR3, WDR11	112
Family me	Family member testing		as indic	ated above	14	
Proforma required? YES Hypogona		Hypogonado	trophic Hypogor	nadism referral form (see centre website)		

Referral criteria

• Clinical history of Hypogonadism

Requesting specialties

- Clinical Genetics
- Endocrinology
- Paediatrics

HYPOPHOSPHATEMIC RICKETS

Available testing

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

Referral criteria

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

- Clinical Genetics
- Endocrinology





MONOGENIC DIABETES

Available testing

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

Referral criteria

- Individuals meeting any one of the following criteria:
 - Minimum two generation family history of diabetes with at least one individual diagnosed under the age of 35 years with BMI less than 30, negative GAD and IA2 autoantibodies and detectable C-peptide, OR
 - High risk of Maturity onset diabetes of the young (MODY) based on MODY calculator http://www.diabetesgenes.org/content/mody-probability-calculator, OR
 - Diabetes in conjunction with cystic renal disease and/or congenital anomaly of the kidney or urinary tract (likely HNF1B), OR
 - Diabetes in conjunction with other extra-pancreatic features suggestive of monogenic diabetes. e.g. deafness, congenital heart disease, epilepsy, diabetes insipidus, developmental delay etc.
 - Post-pubertal children or adults with insulin resistance:
 - Severely elevated plasma insulin (typically greater than 150pmol/L in nondiabetic non-obese subject), AND
 - Clinical features consistent with severe insulin resistance, e.g. polycystic ovarian syndrome, acanthosis nigricans, diabetes with high insulin requirements, post-prandial hypoglycaemia, OR
 - Post-pubertal severe insulin resistance with plasma adiponectin >5mg/l, OR
 - 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	Sco	Scope and range		Targets	TAT
Dundee	NGS (targeted panel)	Targete	d screen	SNVs, indels , exon level CNV (selected genes)	MEN1, CDKN1B, AIP	56
	member ting			as ind	icated above	14
Proforma re	equired?	YES	Endocrin	e disorders proforr	ma (see centre website)	

Referral criteria

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

- Clinical Genetics
- Endocrinology





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

Available testing

Centre	Method	Sco	Scope and range of test		Targets	TAT
Dundee	NGS (targeted panel)	Targete	d screen	SNVs, indels	RET (exons 5, 8, 10, 11, 13, 14, 15, 16)	56
_	member ting			as ind	icated above	14
Proforma r	equired?	YES	Endocrine	e disorders proforr	na (see centre website)	

Referral criteria

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

- Clinical Genetics
- Endocrinology





PHAEOCHROMOCYTOMA AND PARAGANGLIOMA

Available testing

Centre	Method	Scope a	and range of test	Targets	TAT	
Dundee	NGS (targeted panel)	Whole gene 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	
	member ting		as indicated above			
<u> </u>			ndocrine disorders prof	orma (see centre website)		

Referral criteria

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

- Clinical Genetics
- Endocrinology





PIGMENTED NODULAR ADRENOCORTICAL DISEASE

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS (clinical exome)	Whole gene SNVs, indels screen		ARMC5, PDE11A, PDE8B, PRKAR1A	112
Family member testing			as indica	ated above	14
Proforma re	equired?	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 range of test		Targets	TAT
Aberdeen	Sanger	Whole gene SNVs, indels screen		KCNJ5	56
Family member testing			as ind	icated above	14
Proforma required? NO			·		

Referral criteria

• Hypertension presenting in childhood (under 10 years of age)

- Clinical Genetics
- Endocrinology





RENAL CYSTS & DIABETES

Available testing

Centre	Method	Scope and r	ange of test	Targets	TAT
Dundee	NGS (targeted panel)	Whole gene screen SNVs, indels Exon level CNV		HNF1B	56
Family member testing			bove	14	
Proforma required?		NO			

Referral criteria

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

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





SEVERE EARLY ONSET OBESITY

Available testing

Centre	Method	S	Scope and range of test		Targets	TAT
Dundee	NGS (clinical exome)	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 indic		as indic	ated above	14
Proforma required?		YES	Obesity profe	orma (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		nge of test	Targets	TAT
Dundee	NGS (targeted panel)		e gene een	SNVs, indels	THRB	56
_	member ting			as ind	icated above	14
Proforma re	equired?	YES	Endocrine	disorders proforma	a (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 (targeted panel)	Whole gene screen		SNVs, indels Exon level CNV	VHL	56
Family member testing			as indicated above			14
Proforma required?		YES	Endocrin	e disorders proforr	na (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
 - Cerebellar haemangioblastoma (<60 years), OR
 - ≥2 VHL-related tumours (any age), OR
 - ≥1 VHL-related tumour and a first degree relative with ≥1 VHL-related tumour (where one of the tumours is retinal angioma / hemangioblastoma)
- Deceased affected individual (proband) where (i) the individual +/- family history meets one of the above criteria, (ii) appropriate tissue is available (tumour or normal), and (iii) no living affected individual is available for genetic testing
- VHL-related tumours comprise: Retinal angioma, Spinal or cerebellar hemangioblastoma, adrenal or extra-adrenal phaeochromocytoma, Renal cell carcinoma, multiple renal and/or pancreatic cysts, endolymphatic sac tumors, papillary cystadenomas of the epididymis or broad ligament, neuroendocrine tumour of the pancreas
- Overlapping clinical indications:
 - o Phaeochromocytoma and paraganglioma

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





EYES

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

Available testing

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

Referral criteria

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

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





ALBINISM & NYSTAGMUS

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole 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 member testing		as indicated above			
Proforma required?		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		as indicated above				
testing						
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 an	d 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,TTC8, WDPCP	112
Family member testing		as indicated above			
Proforma required?		NO			•

Referral criteria

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

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





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

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV	BEST1	56
Family me	mber testing		as indicated	above	14
Proforma re					

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

BLEPHAROPHIMOSIS, PTOSIS, AND EPICANTHUS INVERSUS (BPES)

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNVs	FOXL2	56
Family r	nember ting		as indicated	d above	14
Proforma required?		NO			

Referral criteria

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

Requesting specialties

Clinical Genetics

Ophthalmology in discussion with





BRITTLE CORNEA SYNDROME

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels	PRDM5, ZNF469	56
Family me	mber testing		as indica	ited above	14
Proforma re	Proforma required? NO				

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

CHOROIDERAEMIA

Available testing

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

Referral criteria

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

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





CONGENITAL CATARACTS

Available testing

Centre	Method	Scope and	I range of test	Targets	TAT
Aberdeen	NGS	Whole 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 m				as indicated above	14
Proforma r	equired?	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, COL8A2, DCN, GJA1, GSN, HMX1, KERA, KRT12, KRT3, LTBP2, MAF, OVOL2, PIK3R1, PIKFYVE, PITX2, PRDM5, RAB18, RAB3GAP1, RAB3GAP2, SLC16A12, SLC4A11, TACSTD2, TGFBI, UBIAD1, VSX1, ZEB1, ZNF469	112
_	member ting	as indicated above		indicated above	14
Proforma required? NO			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, OVOL2, PIKFYVE, PRDM5, SLC4A11, STS, TACSTD2, TCF4, TGFBI, UBIAD1, ZEB1, ZNF469	112
	member ting			as indicated above	14
Proforma re	equired?	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	CHN1, COL25A1, DCC, FRMD7, HOXA1, KIF21A, MAFB, PHOX2A, ROBO3, SALL1, SALL4, TUBB2B, TUBB3	56
	amily member as ind testing		cated above	14	
Proforma re	equired?	NO			

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

FAMILIAL EXUDATIVE VITRORETINOPATHY (FEVR)

Available testing

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

Referral criteria

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

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





GLAUCOMA

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole 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 r	member ting		as indicated above		14
Proforma re	equired?	NO			

Referral criteria

• Clinical features suggestive of a monogenic cause of Glaucoma

Requesting specialties

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

VITELLIFORM MACULAR DYSTROPHY

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV	BEST1, PRPH2	56
Family r		30.33		ted above	14
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





NORRIE DISEASE

Available testing

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

Referral criteria

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

- Clinical Genetics
- in discussion with Clinical Genetics





OCULAR MALFORMATIONS

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	NGS	Whole gene 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		14
Proforma required? NO					

Referral criteria

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

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





OCULOCUTANEOUS ALBINISM

Available testing

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

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

OPTIC NEUROPATHY

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole 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 r	member ting		6	as indicated above	14
Proforma re	equired?	NO			

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 ra	inge 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 (C21orf2), CFH, CHM, CIB2, CLN3, CLN5, CLN6, CLN8, CLRN1, CNGA1, CNGA3, CNGB1, CNGB3, CNNM4, COL18A1, COL4A1, CRB1, CRX, CSPP1, CTNNB1, CTSD, CWC27, CYP4V2, DHDDS, EFEMP1, ELOVL4, ERCC6, ERCC8, EYS, FAM161A, FLVCR1, FZD4, GNAT1, GNAT2, GNPTG, 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 (c2orf71), PCDH15, PCYT1A, PDE6A, PDE6B, PDE6C, PDE6G, PEX1, PEX2, PEX7, PHYH, PLA2G5, POC1B, PPT1, PRCD, PROM1, PRPF3, PRPF31, PRPF4, PRPF6, PRPF8, PRPH2, PRPS1, RAB28, RAX2, RBP3, RBP4, RCBTB1, RD3, RDH12, RDH5, REEP6, RGS9, RHO, RLBP1, RP1, RP1L1, RP2, RP9, RPE65, RPGR, RPGRIP1, RPGRIP1L, RS1, SAG, SCAPER, SDCCAG8, SLC24A1, SLC38A8, SNRNP200, SPATA7, SRD5A3, TIMM8A, TIMP3, TMEM237, TOPORS, TPP1, TRIM32, TRPM1, TSPAN12, TTC8, TTLL5, TUB, TULP1, USH1C, USH1G, USH2A, VCAN, VPS13B, WDPCP, WDR19, WHRN, ZNF408, ZNF423	112
	member			as indicated above	14
tes	ting				
Proforma re	equired?	NO			

Referral criteria

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

- 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
,	member ting			as indicated above	14
Proforma re	equired?	NO			

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics

X-LINKED CONGENITAL NYSTAGMUS

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	FRMD7	56
Family me	amily member testing as indica			ited above	14
Proforma re	roforma 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 member as indicated testing		ed above	14	
Proforma re	Proforma required? NO				

Can be performed prior to Retinal Degeneration panel if required

Referral criteria

• Clinical features suggestive of X-linked Juvenile Retinoschisis

- Clinical Genetics
- Ophthalmology in discussion with Clinical Genetics





GASTROHEPATOLOGY

CHOLESTASIS

Available testing

Centre	Method	Scope and	d 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
,	member ting		as indicated above		14
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Gastroenterology
- Hepatology
- Paediatrics





CRIGLER-NAJJAR SYNDROME, TYPE 1 AND 2

Available testing

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

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Gastroenterology
- Hepatology
- Paediatrics

GILBERT SYNDROME

Available testing

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

Referral criteria

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

- Clinical Genetics
- Gastrohepatology
- Paediatrics
- General Practice





HIRSCHSPRUNG DISEASE

Available testing

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

Referral criteria

- Diagnosis of Hirschsprung disease (HSCR) and at least one of the following:
 - Family history of HSCR, at least 1 affected first or second degree relative, OR
 - HSCR occurring as part of a syndrome or with other anomalies associated with the listed genes

- Clinical Genetics
- Gastroenterology
- Hepatology
- Paediatrics





PANCREATITIS

Available testing

Centre	Method	Scope and range 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 variants	28
Family member testing			as indica	ited above	14
Proforma required?		NO			

Referral criteria

- Recurrent acute pancreatitis
- Chronic pancreatitis
- 1st episode of acute pancreatitis <18 years old
- 1st 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 ran	ge 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 indicat	ed above	14
Proforma required? 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	ange of test	Targets	TAT
Aberdeen	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV	АТР7В	56
Family r			as indicat	ed above	14
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Gastroenterology
- Hepatology





HAEMATOLOGY

ANTITHROMBIN DEFICIENCY

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Edinburgh	Sanger MLPA		e gene een	SNVs, indels Exon level CNV	SERPINC1	56
Family mem	nber testing			as indi	cated above	14
Proforma re	required? YES Molec			r Haematology red	quest 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 scr	e gene een	SNVs	GP1BA, GP1BB, GP9	56
Family mem	nber testing		as indicated above			14
Proforma re	quired?	YES	Molecula	r Haematology red	uest form (see centre website)	

Referral criteria

• Platelet function testing suggestive of Bernard Soulier syndrome

- Clinical Genetics
- Haematology





COAGULATION & FIBRINOLYSIS PANEL

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Edinburgh	NGS	Whole scre	J	SNVs	ACVRL1, CHST14, COL3A1, ENG, 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	equired?	YES	S Molecular Haematology reque		quest form (see centre website)	

Referral criteria

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

- Clinical Genetics
- Haematology





COMBINED FACTOR V AND VIII DEFICIENCY

Available testing

Centre	Method		Scope 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 mem	nber testing			as indicated abov	re	14
Proforma required? YES Molecular			Molecula	r Haematology request form	(see centre website)	

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Haematology

DIAMOND BLACKFAN ANAEMIA

Available testing

Centre	Method	Scope and ra	nge 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
,	member ting		as indi	cated above	14
Proforma re	equired?	NO			

Referral criteria

 Clinical phenotype suggestive of Diamond Blackfan Anaemia – Presenting in the 1st 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 mem	nber testing	sting		as indicated above	14
Proforma required? NO					

Referral criteria

- Idiopathic erythrocytosis with:
 - 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 range of test			Targets	TAT
Edinburgh	Sanger MLPA	Whole gene screen		SNVs Exon level CNV	F7	56
Family mem	nber testing		as indicated above			
Proforma re	quired?	YES Molecular Haematology request form			n (see centre website)	

Referral criteria

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

- Clinical Genetics
- Haematology





FACTOR X DEFICIENCY

Available testing

Centre	Method		Scope and ra	ange of test	Targets	TAT
Edinburgh	Sanger MLPA		nole gene screen	SNVs Exon level CNV	F10	56
Family me	ember testing			as indicate	ed above	14
Proforma re	quired?	YES	Molecular H	aematology request	form (see centre website)	

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Haematology

FACTOR XI DEFICIENCY

Available testing

Centre	Method		Scope and	range of test	Targets	TAT
Edinburgh	Sanger MLPA		nole gene screen	SNVs Exon level CNV	F11	56
Family me	ember testing			as indicated	above	14
Proforma re	quired?	YES	Molecular H	aematology request for	orm (see centre website)	

Referral criteria

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

- Clinical Genetics
- Haematology





FIBRINOGEN DEFICIENCY

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Edinburgh	NGS	Whole scre	e gene een	SNVs	FGA, FGB, FGG	56
Family mem	nber testing		as indicated above			
Proforma required? YES Molecular			Molecula	r Haematology red	quest form (see centre website)	

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Haematology

G6PD Deficiency

Available testing

Centre	Method	Scope and range of test			Targets			TAT
Edinburgh	NGS		Whole gene SNV screen		s G6PD			56
Family n						icated above		14
Proforma re	quired?	NO						

Referral criteria

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





GLANZMANN THROMBASTHENIA

Available testing

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

Referral criteria

• Platelet function testing suggestive of Glanzmann thrombasthenia

- Clinical Genetics
- Haematology





HAEMOCHROMATOSIS

Available testing

Centre	Method	Scope and rang	ge of test	Targets	TAT
Aberdeen Dundee Glasgow	ARMS (G) Sanger (A) Genotyping Assay	Targeted screen	SNVs	<i>HFE</i> p.C282Y & p.H63D	28
Proforma required?		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 MLPA	Whole gene screen	SNVs, indels	HBB	56	
Edinburgh	MLPA		Indels	HBA	56	
Family member testing		as indicated above				
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 and range of test			Targets	TAT
Edinburgh	NGS MLPA Inversion PCR		hole gene screen	SNVs Exon level CNV Inversions *	F8	56
Family member testing		as indicated above				14
Proforma required?		YES	Molecular Haematology request form (see centre website)			

^{*} 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	Whol	e gene screen	SNVs, indels Exon level CNV	F9	56
Family me	ember testing	as indicated above				14
Proforma rec	quired?	YES	S Molecular Haematology request form (see centre website)			

Referral criteria

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

- Clinical Genetics
- Haematology





INHERITED BONE MARROW FAILURE

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene 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
,	member tina		as indic	cated above	14
Proforma re		NO			l

Referral criteria

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

- Clinical Genetics
- Haematology





IRON REGULATION

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	NGS	Whole gene 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 ir	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 ra	ange of test	Targets	TAT	
Aberdeen	NGS	Whole gene screen SNVs, indels Exon level CNV		SRP72, GATA2	56	
Family m	ember testing		oove	14		
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	Sco	pe and	range of test	Targets	TAT
Aberdeen	Sanger	_	Whole gene SNVs, indels		ELANE	56
Family member as indicate testing		as indicated	d above	14		
Proforma re	equired?	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	TAT
Edinburgh	NGS	Whole gene screen	SNVs, indels	ABCG5, ABCG8, ACTB, ACTN1, ADAMTS13, ANKRD26, ANO6, AP3B1, AP3D1, ARPC1B, BLOC1S3, BLOC1S6, CDC42, CYCS, DIAPH1, DTNBP1, ETV6, FERMT3, FLI1, FLNA, FYB1, GATA1, GBA, GF11B, GNE, GP1BA, GP1BA, GP1BB, GP6, GP9, HOXA11, HPS1, HPS3, HPS4, HPS5, HPS6, IKZF5, ITGA2B, ITGA2B, ITGB3, ITGB3, KDSR, LYST, MECOM, MPIG6B, MPL, MYH9, NBEA, NBEAL2, P2RY12, PLA2G4A, PLAU, PTGS1, RASGRP2, RBM8A, RNU4ATAC, RUNX1, SLFN14, SRC, STIM1, STXBP2, TBXA2R, TBXAS1, THPO, TUBB1, TPM4, VIPAS39, VPS33B, VWF, WAS	84
Family member testing			as indicated above	14	
Proforma re	equired?	YES N	Nolecular Haem	natology 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	I range of test	Targets	TAT
Edinburgh	Sanger MLPA	Whole	e gene een	SNVs, indels Exon level CNV	PROC	56
Family mem	nber testing			as indicated a	above	14
Proforma re	quired?	YES	Molecula	r Haematology request for	orm (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 gene screen		SNVs, indels Exon level CNV	PROS1	56	
Family member testing		as indicated above				14	
Proforma required?		YES	YES Molecular Haematology request form (see centre website)				

Referral criteria

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

- Clinical Genetics
- Haematology





RARE ANAEMIA PANEL (Panel app R92)

Available testing

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

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

Referral criteria

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

- Clinical Genetics
- Haematology





SCHWACHMAN-DIAMOND SYNDROME

Available testing

Centre	Method	5	Scope and ra	ange of test	Targets	TAT
Aberdeen	NGS	Whole gene screen		SNVs, indels Exon level CNV	SBDS, DNAJC21	56
Family member testing			14			
Proforma required?		YES	ES GEN FORM 215 Primary Immunodeficiency Request form (see centre website)			

Referral criteria

• Clinical phenotype suggestive of Schwachman-Diamond Syndrome

- Clinical Genetics
- Haematology





SICKLE CELL ANAEMIA

Available testing

Centre	Method	Scope and range of	test	Targets	TAT
Aberdeen	Sanger	Targeted screen	SNVs	HBB p.(Glu7Val)	28 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 Haematology test
- For prenatal testing, both parents to be confirmed as carrier by genetics prior to offering invasive prenatal test. Please contact the laboratory to discuss
- Newborn screening (Glasgow)
 - Newborns who have undergone a blood transfusion prior to the blood spot sample being taken.

- Clinical Genetics
- Haematology
- Obstetrics





THROMBOPHILIA (FACTOR V LEIDEN & PROTHROMBIN)

Available testing

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

^{*}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	/ES 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 range of test			Targets	TAT
Edinburgh	NGS MLPA	Whol	e gene screen	SNVs Exon level CNV	VWF	56
Family m	ember testing	er testing as indicate			above	14
Proforma required? YES Molecular Hae			Molecular Hae	matology request fo	rm (see centre website)	

Referral criteria

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

- Clinical Genetics
- Haematology





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 re	equired?	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
 - 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 (clinical exome)	Whole gene screen	SNVs, indels, Exon level CNV (EYA1)	EYA1, SIX1, SIX5	56
Family member as indicated a testing		above	14		
Proforma re	equired?	NO			

Referral criteria

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

- Audiology
- Clinical Genetics
- Nephrology





HEARING LOSS, SYNDROMIC & NON SYNDROMIC

Available testing

Centre	Method	Scope and	range of test	Targets	TAT
Dundee	NGS (clinical exome)	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			as indicated above	14	
testing Proforma required? NO		NO			<u> </u>

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 range of test		Targets	TAT		
Dundee	Sanger and fragment analysis	Whole gene screen	SNVs, indels (GJB2) Deletions (GJB6)	GJB2, GJB6	28		
	Family member as indicated above testing			/e	14		
Proforma re	equired?	NO	IO				

Referral criteria

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

- Audiology
- Clinical Genetics
- Paediatrics





PENDRED SYNDROME

Available testing

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

Referral criteria

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

- Audiology
- Clinical Genetics
- Paediatrics





USHER SYNDROME

Available testing

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

Referral criteria

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

Requesting specialties

• Clinical Genetics

WAARDENBURG SYNDROME

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS (clinical exome)	Whole gene screen	SNVs, indels	PAX3, MITF, SOX10, SNAI2, EDNRB, EDN3, KIT	112
Family me	mily member testing			as indicated above	14
Proforma re	Proforma required? NO				

Referral criteria

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

Requesting specialties

• Clinical Genetics





WOLFRAM SYNDROME

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS or Sanger	Whole gene screen SNVs, indels		WFS1	56
Family member testing			as inc	icated above	14
Proforma required?		NO			

Referral criteria

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

- Clinical Genetics
- Endocrinology





IMMUNOLOGY

ADENOSINE DEAMINASE DEFICIENCY (ADAD)

Available testing

Centre	Method	,	Scope and	range of test	Targets	TAT
Aberdeen	Sanger MLPA		e gene een	SNVs, indels Exon level CNV	ADA2 (CECR1)	56
Family mer testing	nber		as indicated above			14
Proforma re	a required? YES GEN FORM 215 Primary Immunodeficiency Request form (see centre			ebsite)		

Referral criteria

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

- Clinical Genetics
- Immunology





ANHYDROTIC ECTODERMODYSPLASIA WITH ID

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole scr	e gene een	SNVs, indels Exon level CNV	IKBKG (NEMO), NFKBIA (IKBA)	56
Family r				as indicate	ed above	14
Proforma re	equired?	YES	GEN FOR	RM 215 Primary Immur	nodeficiency Request form (see centre we	bsite)

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





ALPHA 1 ANTITRYPSIN DEFICIENCY

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	Sanger	Whole gene screen	SNVs, indels	SERPINA1	28
Family member testing			as indi	cated above	14
Proforma required?		NO			

Referral criteria

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

- Clinical Genetics
- Gastroenterology
- Hepatology
- Respiratory Medicine





ASSOCIATION WITH GI INFLAMMATION

Available testing

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

Referral criteria

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

- Clinical Genetics
- Rheumatology





AUTOINFLAMMATORY DISORDERS

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	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 r				as indicated above	14
Proforma re	equired?	YES	GEN FORM 215 Pr	imary Immunodeficiency Request form (see centre web	site)

Referral criteria

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

- Clinical Genetics
- Immunology
- Rheumatology





BACTERIAL AND PARASITIC INFECTIONS

Available testing

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

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





BACTERIAL INFECTIONS, AUTOINFLAMMATION, AMYLOPECTINOSIS

Available testing

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

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





CALCIUM CHANNEL DEFECTS

Available testing

Centre	Method		Scope an	d range of test	Targets	TAT
Aberdeen	NGS	Whole scr	e gene een	SNVs, indels Exon level CNV	ORAI1, STIM1	56
Family r				as indicated ab	ove	14
Proforma re	equired?	YES	GEN FOR	RM 215 Primary Immunode	ficiency Request form (see centre we	bsite)

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





CHRONIC GRANULOMATOUS DISEASE

Available testing

Centre	Method		Scope and ra	ange of test	Targets	TAT
Aberdeen	NGS	Whole	gene screen	SNVs, indels Exon level CNV	CYBA, CYBB, NCF1, NCF2, NCF4	56
Family men	nber testing			as indicated	above	14
Proforma required? YES GEN			GEN FORM	GEN FORM 215 Primary Immunodeficiency Request form (see centre websi		

Referral criteria

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

- Clinical Genetics
- Immunology





COMBINED IMMUNODEFICIENCIES (CVID)

Available testing

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

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





COMPLEMENT DEFICIENCIES

Available testing

Centre	Method	Scope	and range of test	Targets	TAT
Aberdeen	NGS	Whole ge screen	i Exon ievei	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
,	member ting			as indicated above	14
Proforma re	equired?	YES	GEN FORM 215 Prir	mary Immunodeficiency Request form (see centre web	site)

Referral criteria

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

- Clinical Genetics
- Immunology
- Rheumatology





CONGENITAL THROMBOCYTOPENIA

Available testing

Centre	Method	Sco	pe and ra	nge of test	Targets	TAT
Aberdeen	NGS		e gene een	SNVs, indels Exon level CNV	ARPC1B, WAS, WIPF1	56
Family r	member ting			as ind	icated above	14
Proforma re	equired?	YES	GEN FOR	RM 215 Primary In	nmunodeficiency Request form (see centre we	bsite)

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





DEFECTS OF VITAMIN B12 AND FOLATE METABOLISM

Available testing

Centre	Method	Scope and range of test			Targets	TAT		
Aberdeen	NGS	Whole gene screen		SNVs, indels Exon level CNV	MTHFD1, SLC46A1, TCN2	56 or 112		
Family r				as ind	icated above	14		
Proforma re	oforma required? YES			GEN FORM 215 Primary Immunodeficiency 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 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





DNA REPAIR DEFECTS

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole scre	gene een	SNVs, indels Exon level CNV	ATM, BLM, CDCA7, DNMT3B, GINS1, HELLS, LIG1, MCM4, NBS1 (NBN), PMS2, POLE1, POLE2, NSMCE3, ERCC6L2, RNF168, ZBTB24	56
Family r	nember ting	as i		as ind	icated above	14
Proforma re	Proforma required? YES G		GEN FOR	RM 215 Primary In	nmunodeficiency Request form (see centre web	osite)

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





DYSKERATOSIS CONGENITA (DKC)

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene 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 r				as indicated above	14
Proforma re	equired?	YES	GEN FORM 215 Pr	imary Immunodeficiency Request form (see centre we	bsite)

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





FAMILIAL HLH DUE TO PRF1 VARIANTS

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	Sanger MLPA	Whole scr	e gene een	SNVs, indels Exon level CNV	PRF1	56
Family mer testing	nber	as indicated			above	14
			deficiency Request form (see centre we	ebsite)		

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





HAEMOPHAGOCYTIC LYMPHOHISTOPCYTOSIS (HLH) & EBV SUSCEPTIBILITY

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	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 r	member ting			as indicated above	14
Proforma re				mary Immunodeficiency Request form (see centre web	site)

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology
- Rheumatology





HENNEKAM-LYMPHANGIECTASIA-LYMPHEDEMA SYNDROME

Available testing

Centre	Method	S	cope and	range of test	Targets	TAT
Aberdeen	NGS		Whole gene SNVs, inde screen Exon level C		CCBE1, FAT4	56
Family r				as indicate	d above	14
Proforma re	Proforma required? YES GEN FORM 215 Primary Immu			nodeficiency Request form (see centre we	bsite)	

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





HEPATIC VENO-OCCLUSIVE DISEASE WITH IMMUNODEFICIENCY (VODI)

Available testing

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

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





HEREDITARY AMYLOIDOSIS

Available testing

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

Referral criteria

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

- Clinical Genetics
- Rheumatology





HEREDITARY ANGIOEDEMA, TYPES I & II

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	Sanger MLPA	Whole gene screen		SNVs, indels Exon level CNV	SERPING1	56
Aberdeen	NGS	Whole gene screen		SNVs, indels Exon level CNV	SERPING1, Factor XII, PLG, ANGPT1	56
Family r	nember ting		as indicated above			
Proforma required? YES GEN FORM 215 Primary Immu			nodeficiency Request form (see centre we	bsite)		

Referral criteria

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

- Clinical Genetics
- Immunology





HYPER IGE SYNDROMES (HIES)

Available testing

Centre	Method		Scope an	d range of test	Targets	TAT
Aberdeen	NGS		e gene een	SNVs, indels Exon level CNV	PGM3, SPINK5, STAT3	56
Family member as indicated above testing			ove	14		
Proforma required? YES			GEN FOR	RM 215 Primary Immunode	ficiency Request form (see centre web	site)

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





HYPOGAMMAGLOBULINAEMIA

Available testing

Centre	Method	Scope ar	nd range of test	Targets	TAT
Aberdeen	NGS	Whole gene screen	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 r		ber as indicated above		as indicated above	14
Proforma re	equired?	YES	GEN FORM 215 Pri	imary Immunodeficiency Request form (see centre we	ebsite)

Referral criteria

- IgG, IgA and / or IgM decreased
- Exclude second causes: drugs [Hx], myeloma [bone marrow], lymphoma. Ig loss (not hypo-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 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





ID WITH MULTIPLE INTESTINAL ATRESIAS

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole scr	e gene een	SNVs, indels Exon level CNV	TTC7A	56
Family r				as indicated	above	14
Proforma required? YES GEN FORM 215 Primary Imr			RM 215 Primary Immuno	deficiency Request form (see centre we	ebsite)	

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





IMMUNO-OSSEOUS DYSPLASIAS

Available testing

Centre	Method	Scope a	and range of test	Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	EXTL3, MYSM1, RMRP, RNU4ATAC, SMARCAL1	56
Family r				as indicated above	14
Proforma required? YES GEN FORM 215 Prim			GEN FORM 215 Prim	nary Immunodeficiency Request form (see centre we	bsite)

Referral criteria

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

- Clinical Genetics
- Immunology





INTERFERONOPATHY / SLS / AGS / COMPLEMENT

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole 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 mer testing	nber		as indicated above		
Proforma re	equired?	YES	GEN FORM 215 Pri	mary Immunodeficiency Request form (see centre we	ebsite)

Referral criteria

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

- Clinical Genetics
- Immunology
- Rheumatology





KABUKI SYNDROME

Available testing

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

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





MENDELIAN SUSCEPTIBILITY TO MYCOBACTERIAL DISEASE (MSMD) AND VIRAL INFECTION

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdee n	NGS	Whole 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 member testing		as indicated above			14
Proforma required?		YE GEN FOR	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 susceptibility to viral infection Herpes simplex Encephalitis, Predisposition to severe viral infection), please refer to GEN FORM 215 Primary Immunodeficiency Request form on centre website and IUIS 2022 (J Clin Immunol., 2022 42:1473-1507).

- Clinical Genetics
- Haematology
- Immunology





MISCELLANEOUS AUTOINFLAMMATORY CONDITIONS

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	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 mer testing	nber		as indicated above		
Proforma re	equired?	YES (GEN FORM 215 I	Primary Immunodeficiency Request form (see centre web	site)

Referral criteria

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

- Clinical Genetics
- Immunology
- Rheumatology





NEUTROPENIA

Available testing

Centre	Method	Scope	and range of test	Targets	TAT	
Aberdeen	NGS	Whole ge screen	i Exon ievei	C16ORF57 (USB1), CLPB, COH1 (VPS13B), CSF3R, DNAJC21, ELANE, G6PC3, G6PT1 (SLC37A4), GFI1, HAX1, HYOU1, JAGN1, LAMTOR2, MKL1 (MRTFA), SBDS, SMARCD2, TAZ, VPS45, WAS, WDR1	56	
_	nember ting		as indicated above			
Proforma re	equired?	YES	GEN FORM 215 Primary Immunodeficiency Request form (see centre website)			

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





OTHER ANTIBODY DEFICIENCIES

Available testing

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

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





PURINE NUCLEOSIDE PHOSPHORYLASE DEFICIENCY

Available testing

Centre	Method	Scope and range of test			Targets	TAT	
Aberdeen	NGS		e gene een	SNVs, indels Exon level CNV	PNP	56	
Family r				as indicated ab	oove	14	
Proforma re	equired?	YES	ES GEN FORM 215 Primary Immunodeficiency Request form (see centre web				

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





SEVERE COMBINED IMMUNODEFICIENCY (SCID)

Available testing

Centre	Method	Scope 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 r	member ting		as indicated above			14
Proforma re	equired?	YES	GEN FORM 215 Primary Immunodeficiency Request form (see centre webs			

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





STAT5B DEFICIENCY

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS		e gene een	SNVs, indels Exon level CNV	STAT5B	56
Family r			as indicated above			
Proforma required? YES GEN FORM 215 Primary Immunodeficiency Request form (see centre				ebsite)		

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





STROKE

Available testing

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

Referral criteria

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

- Clinical Genetics
- Immunology
- Rheumatology





SYNDROMES ASSOCIATED WITH AUTOIMMUNITY AND OTHERS

Available testing

Centre	Method	Scope and range of test		Targets	TAT	
Aberdeen	NGS	Whole gene screen	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 r			as indicated above			
Proforma re	equired?	YES	GEN FORM 215 Primary Immunodeficiency Request form (see centre we			

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





SYNDROMES ASSOCIATED WITH CONGENITAL DEFECTS OF PHAGOCYTES

Available testing

Centre	Method	Scope	and range of to	est Targets	TAT	
Aberdeen	NGS	Whole ge	I EXONIE	evel FERMT3 (LADIII), FPR1, GATA2, G6PD, ILGB2	56	
Family r			as indicated above			
Proforma re	equired?	YES	GEN FORM 215 Primary Immunodeficiency Request form (see centre we			

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





THYMIC DEFECTS WITH CONGENITAL ANOMALIES

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	NGS	Whole gene screen		SNVs, indels Exon level CNV	CHD7, SEMA3E, TBX1, FOXN1	56
Family r				as indicat	ed above	14
Proforma required? YES GEN FORM 215 Primary Immu			unodeficiency Request form (see centre w	ebsite)		

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





VASCULOPATHY

Available testing

Centre	Method	Scope and range of test		Targets	TAT	
Aberdeen	NGS	Whole gene screen	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 mer testing	nber		as indicated above			
Proforma re	equired?	YES	GEN FORM 215 Primary Immunodeficiency Request form (see centre webs			

Referral criteria

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

- Clinical Genetics
- Immunology
- Rheumatology





VERY EARLY ONSET INFLAMMATORY BOWEL DISEASE (VEO-IBD)

Available testing

Centre	Method	Scope	and range of test	Targets	TAT		
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	ADAM17, AICDA, CD40LG, BTK, CD3G, ZAP70, WAS, CYBA, CYBB, NCF1, NCF2, NCF4, DOCK8, EPCAM (Sanger only), FOXP3, GUCY2C, HPS1, HPS4, HPS6, ADA, IL2RG, LIG4, DCLRE1C, RAG2, IL10, II10RA, IL10RB, ITGB2, LRBA, ICOS, PIK3R1, PLCG2, RET, SH2D1A, XIAP, SKIV2L, TTC37, SLC37A4, SKIV2L, STAT1, STAT3, STXBP2	112		
Family men	nber		as indicated above				
testing		VEO	VEO OFN FORM 045 B:				
Proforma re	equired?	YES	ES GEN FORM 215 Primary Immunodeficiency Request form (see centre webs				

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 gene screen		SNVs, indels Exon level CNV	EPG5	56
Family r			as indicated		d above	14
Proforma required? YES GEN FORM		RM 215 Primary Immun	odeficiency Request form (see centre w	ebsite)		

Referral criteria

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

- Clinical Genetics
- Haematology
- Immunology





X-LINKED CGD

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen	Sanger MLPA	Whole ge	ene screen	SNVs, indels Exon level CNV	CYBB	56
Family r			as indica		d above	14
Proforma re	<u> </u>	YES	GEN FORI	M 215 Primary Immur	nodeficiency Request form (see centre we	ebsite)

Referral criteria

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

- Clinical Genetics
- Immunology





COWDEN SYNDROME / PTEN HAMARTOMA TUMOUR SYNDROME (PHTS)

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Aberdeen Glasgow	NGS	Whole g	_	SNVs, indels Exon level CNV	PTEN	56
Family r			as indicated above		ve	14
Proforma re	equired?	NO				

Referral criteria

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

- Clinical Genetics
- Dermatology
- Neurology
- Paediatrics





DICER1 SYNDROME

Available testing

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

Referral criteria

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

Requesting specialties





FAMILIAL MELANOMA

Available testing

Centre	Method	S	Scope and	range of test	Targets	TAT
Glasgow	NGS		e gene een	SNVs, indels	BAP1, BRCA2, CDKN2A, CDK4, POT1	56
	member ting			as indicated	dabove	14
Proforma re	equired?	YES	Inherited	cancer proforma (see c	entre website)	

Referral criteria

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

- Clinical Genetics
- Oncology





GORLIN SYNDROME (BASAL CELL NEVUS SYNDROME)

Available testing

Centre	Method	Scope and ra	ange of test	Targets	TAT
Glasgow	NGS MLPA	Whole gene screen	SNVs, indels Exon level CNV	PTCH1, SUFU	56
,	member ting		as indica	ated above	14
Proforma re	equired?	NO			

Referral criteria

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

Requesting specialties





HEREDITARY BREAST / 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 r				as indicated above			
Proforma re	equired?	YES	Glasgow laboratory	only (see centre website)			

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

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

Requesting specialties

- Clinical Genetics
- Oncology

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

Living unaffected individual who meets ONE of the following criteria:

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

Requesting specialties





HEREDITARY BREAST / OVARIAN CANCER SYNDROME:

Founder Variants ONLY

Available testing

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

Referral criteria

Living affected individual with:

- Breast Cancer <50 years OR
- Manchester Score ≥10 or ≥5% mutation probability on CanRisk*
- AND is from one of the following founder populations
 - o Ashkenazi Jewish
 - o Poland
 - Orkney specific variant
 - Italian specific family (Edinburgh)

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

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

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

Requesting specialties





HEREDITARY BREAST / OVARIAN CANCER SYNDROME: deceased TESTING

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels	BRCA1, BRCA2	56
Family r			•	as indicated above	14
Proforma re	equired?	YES			

Referral criteria

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

Requesting specialties





HEREDITARY COLORECTAL CANCER, LYNCH SYNDROME AND POLYPOSIS

Available testing

Centre	Method		Scope and range of test	Targets	TAT
Edinburgh	NGS	Whole gene screen	SNVs, indels CNV analysis (MLPA) – Polyposis referrals: APC, MUTYH (selected exons), and GREM1 (upstream region) Lynch referrals and patient dx CRC <45yrs: MLH1, MSH2, MSH6 and EPCAM (selected exons)	APC, BMPR1A, MBD4, MLH1, MSH2, MSH3, MSH6, MUTYH, NTHL1, PMS2, POLD1 (exons 4-12), POLE (exons 3-13), PTEN, RNF43, SMAD4, STK11	56
Family m			as indicated above		14
Proforma re	quired?	YES	Colorectal cancer gene panel proforma (see centre web	site)	

Referral criteria

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





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





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

Requesting specialties





HEREDITARY DIFFUSE GASTRIC CANCER SYNDROME

Available testing

Centre	Method		Scope and ra	ange of test	Targets	TAT
Edinburgh	NGS	Whole	gene screen	SNVs, indels	CDH1, CTNNA1	56
Family men	nber testing			as indicated abo	ove	
Proforma red	quired?	NO				

Referral criteria

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

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

Note: At least one cancer should be histologically confirmed

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

- Clinical Genetics
- Gastroenterology





HEREDITARY LEIOMYOMATOSIS AND RENAL CELL CANCER SYNDROME

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Dundee	NGS (targeted panel)	Whole	gene screen	SNVs, indels, exon level CNVs	FH	56
Family mer	mber testing			as indicated abo	ove	14
Proforma red	quired?	NO				

Referral criteria

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

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

Requesting specialties





HEREDITARY OVARIAN CANCER SYNDROME

Available testing

Centre	Method	Scope 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 mer testing	nber			as indicated above	14
Proforma required? YES Glasgow laboratory			Glasgow laboratory	only (see centre website)	

Referral criteria

• High-grade non-mucinous ovarian cancer, any age

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

- Clinical Genetics
- Oncology





INHERITED PANCREATIC CANCER

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Glasgow	NGS MLPA	Whole	•	SNVs, indels Exon level CNV	BRCA2, CDK4, CDKN2A, MLH1, MSH2, MSH6, PALB2, STK11, TP53	56
Family r			as indicated above			
Proforma re	equired?	YES	Heredi	itary cancer profoi	rma (see centre website)	

Referral criteria

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

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

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

- Clinical Genetics
- Oncology in discussion with Clinical Genetics





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 me	mber testing	bove	14		
Proforma required?		YES Colored	ctal cancer gene panel profo	orma (see centre website)	

Referral criteria

- Juvenile polyposis syndrome:
 - $\circ\quad a.\geq 5$ juvenile polyps of the colorectum, OR
 - \circ b. \geq 2 juvenile polyps throughout the GI tract, OR
 - \circ c. \geq 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	d range of test	Targets	TAT	
Aberdeen Glasgow	NGS	Whole gene screen	SNVs, indels Exon level CNV	TP53	56	
Family mem	ber testing	as indicated above				
Proforma required?		NO				

Referral criteria

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

- Clinical Genetics
- Oncology





MEDULLARY THYROID CANCER

Available testing

Centre	Method	Scope and range of test			Targets	TAT	
Dundee	NGS	Targeted screen		SNVs, indels	RET (exons 5, 8, 10, 11, 13, 14, 15, 16)	56	
Family r			as indicated above			14	
Proforma required? YES			Endocrin	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 range of test			Targets	TAT
Edinburgh	NGS MLPA	Whole gene screen		SNVs, indels Exon level CNV	STK11	56
Family mer	mber testing	ting as indicated above				14
Proforma re	equired?	YES	Colorectal can	cer gene panel proform	na (see centre website)	•

Referral criteria

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

Requesting specialties





RENAL CANCER

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS	Whole gene screen	SNVs, indels, Exon level CNV for selected genes	BAP1, FH, FLCN, MET, PTEN, SDHB, VHL	56
	Family member as in-		dicated above	14	
Proforma re	equired?	NO			

Referral criteria

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

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

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

Requesting specialties





RHABDOID TUMOUR

Available testing

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

Referral criteria

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

Requesting specialties





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
,	member ting		as indicated above		14
Proforma re	equired?	NO			-

Referral criteria

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

- Clinical Genetics
- Metabolic





BATTEN DISEASE

Available testing

Centre	Method	Scope a	and range of test	Targets	TAT
Aberdeen	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV	TPP1	56
Family me	Family member testing as indicated a			ove	14
Proforma re	equired?	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 indicated above			14
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Metabolic





BIOTIN-THIAMINE-RESPONSIVE BASAL GANGLIA DISEASE (BTBGD)

Available testing

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

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Metabolic

BROWN VIALETTO VAN LAERE SYNDROME (BVVLS)

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	SLC52A2, SLC52A3	56
	member ting		as indic	ated above	14
Proforma required?		NO			

Referral criteria

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

- Clinical Genetics
- Metabolic





CARNITINE PALMITOYLTRANSFERASE II DEFICIENCY

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	CPT2	56
Family r			as indic	ated above	14
Proforma re	equired?	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 r			as indicated above		14
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 r			as indica	ted above	14
Proforma required?		NO			

Referral criteria

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

- Clinical Genetics
- Metabolic





COBALAMIN C DEFICIENCY

Available testing

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

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Metabolic

CREATINE DEFICIENCY SYNDROME

Available testing

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

Referral criteria

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

- Clinical Genetics
- Metabolic





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

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole 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, LPIN1, MMAA, MMAB, MMACHC, MMADHC, MUT, NAGS, OAT, OTC, OXCT1, PCCA, PCCB, SLC16A1, SLC22A5, SLC25A13, SLC25A15, SLC25A20, SLC7A7, SLC52A2, SLC52A3	112
Family member testing			as indicated above	14	
Proforma re	<u> </u>	NO			

Referral criteria

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

- Clinical Genetics
- Metabolic





DISORDERS OF CARBOHYDRATE METABOLISM (incl. GLYCOGEN STORAGE DISORDERS)

Available testing

Centre	Method	Scope and	I range of test	Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	AGL, ALDOA, ALDOB, 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, SLC16A1, SLC37A4	112
Family me	mber testing	nber testing as indicated above		14	
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Metabolic





FABRY DISEASE

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	Sanger	Whole gene screen	SNVs, indels	GLA	56
Family n			as indicated above		
Proforma re	quired?	NO			

Referral criteria

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

- Clinical Genetics
- Metabolic





FAMILIAL HYPERCHOLESTEROLAEMIA

Available testing

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

Referral criteria

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

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 must complete proforma or discuss with Lipid consultant prior to referral
- Lipidology
- Metabolic





FANCONI-BICKEL SYNDROME

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	SLC2A2	56
Family r			as indi	cated above	14
Proforma re	equired?	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	I 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 r	member ting		as indicated above		14
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Metabolic
- Neurology





GALACTOSAEMIA

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV	GALT	56
,	Family member as indicated a testing			bove	14
Proforma re	equired?	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 r			as indica	ated above	14
Proforma re	equired?	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 as in testing		dicated above	14		
Proforma re	equired?	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 as in-		dicated above	14	
Proforma required?		NO			

Referral criteria

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

- Clinical Genetics
- Metabolic





GLYCOGEN STORAGE DISEASE

Available testing

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

Referral criteria

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

GLYCOGEN STORAGE DISEASE 1A

Available testing

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

Referral criteria

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

- Clinical Genetics
- Metabolic





HOMOCYSTINURIA

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	CBS, MMADHC, MTHR, MTR, MTRR	56
Family r			as indica	ted above	14
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Metabolic





HYPERLIPIDAEMIA, TYPE III

Available testing

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

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Lipidology

HYPERTRIGLYCERIDAEMIA / FAMILIAL CHYLOMICRONAEMIA SYNDROME / LIPOPROTEIN LIPASE DEFICIENCY

Available testing

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

Referral criteria

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

- 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 member as indic		cated above	14		
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Lipidology





LYSOSOMAL STORAGE DISORDERS

Available testing

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

Referral criteria

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

- Clinical Genetics
- Metabolic





MAPLE SYRUP URINE DISEASE (MSUD)

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	BCKDHA, BCKDHB, DBT	56
_	Family member as indicated testing		d above	14	
Proforma re	equired?	NO			

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Metabolic

MEDIUM CHAIN ACYL-COA DEHYDROGENASE DEFICIENCY (MCADD)

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	Sanger	Whole gene screen	SNVs, indels	ACADM	56
	Family member as indicated testing			above	14
Proforma re		NO			

Referral criteria

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

- Clinical Genetics
- Metabolic
- Paediatrics





METACHROMATIC LEUKODYSTROPHY

Available testing

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

Referral criteria

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

- Clinical Genetics
- Metabolic





MUCOPOLYSACCHARIDOSIS (mps) PANEL

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Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	ARSB, ARSK, GALNS, GLB1, GNS, GUSB, HGSNAT, HYAL1, IDS, IDUA, NAGLU, SGSH	112
Proforma req	uired?	NO			

Referral criteria

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





MUCOPOLYSACCHARIDOSIS TYPE 1 (HURLER / SCHEIE SYNDROME)

Available testing

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

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Metabolic

MUCOLIPIDOSIS II & III ALPHA / BETA

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	GNPTAB	56
Family r		as ind		cated above	14
Proforma re	Proforma required? NO				

Referral criteria

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

- Clinical Genetics
- Metabolic





MULTIPLE ACYL-CoA DEHYDROGENASE DEFICIENCY (MADD)

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	ETFDH, ETFA, ETFB, SLC52A2, SLC52A3	56
_	member ting		as in	dicated above	14
Proforma re	equired?	NO			

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Metabolic
- Neurology

NEURONAL CEROID LIPOFUSCINOSIS (NCL)

Available testing

Centre	Method	Scope and range of test		Targets	TAT		
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	ATP13A3, CLN3, CLN5, CLN6, CLN8, CTSD, DNAJC5, KCTD7, MFSD8, PPT1, TPP1	56		
Family r			as indi	cated above	14		
test							
Available g	Available genes: See website						
Proforma re	equired?	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 as indi		cated above	14	
Proforma re	equired?	NO			

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Metabolic

NIEMANN-PICK DISEASE TYPES A & B

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	SMPD1	56
Family r		er as inc		icated above	14
Proforma re	equired?	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 member as indicated testing		ed above	14		
Proforma re	equired?	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 r	member ting		as indic	ated above	14
Proforma re	equired?	NO			•

Referral criteria

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

- Clinical Genetics
- Metabolic





ORGANIC ACIDAEMIAS & COFACTOR / VITAMIN DISORDERS

Available testing

Centre	Method	Scope and	d 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, PC, PCCA, PCCB, PDHA1, PDHB, PDHX, PRDX1, SLC19A3, SLC46A1, SLC52A3, SUCLA2, SUCLG1, TAZ, TCN2, TMEM70	112
,	member	·		as indicated above	14
tes	ting				
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Metabolic





ORGANIC ACIDURIA

Available testing

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

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Metabolic

ORNITHINE AMINOTRANSFERASE DEFICIENCY

Available testing

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

Referral criteria

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

- Clinical Genetics
- Metabolic





ORNITHINE TRANSCARBAMULASE DEFICIENCY

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	NGS MLPA	Whole gene screen	SNVs, indels Exon level CNV	ОТС	56
Family mem	nber testing		as indic	ated above	14
Proforma re	quired?	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, ACBD5, ACOX1, ACK, AGPS, AGXT, AMACR, ARSE, CAT, DNM1L, DYM, EBP, FAR1, GNPAT, GRHPR, HOGA1, HSD17B4, NDHL, PEX1, PEX2, PEX3, PEX5, PEX6, PEX7, PEX10, PEX11B, PEX12, PEX13, PEX14, PEX16, PEX19, PEX26, PHYH, SCP2, TRIM37	112
Family member testing			as indicated above	14	
Proforma re	equired?	Proforma required? NO			

Referral criteria

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

- Clinical Genetics
- Metabolic





PHENYLKETONURIA

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV	PAH	56
	member ting	as inc		licated above	14
Proforma re	equired?	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	56
	Family member as indi		cated above	14	
Proforma re	equired?	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	56
,	Family member as indica		ted above	14	
testing					
Proforma re	Proforma required? NO				

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Metabolic

REFSUM DISEASE

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	NGS	Whole gene screen	SNVs, indels Exon level CNV	PEX7, PHYH	56
	Family member as indic		ited above	14	
Proforma re	equired?	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			as indicated above			
testing						
Proforma re	equired?	NO				

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Metabolic

TANGO2-RELATED METABOLIC ENCEPHALOPATHY & ARRHYTHMIAS

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger	Whole gene screen Long Range PCR	SNVs, indels Ex3-9 deletion	TANGO2	56
Family r			as indicated abo	ove	14
Proforma re	equired?	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	SNVs, indels	HEXA	56
Family r	member	screen	as ind	dicated above	14
Family member as included as i		200104 45010			
Proforma re	equired?	NO			

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Metabolic

TRIMETHYLAMMINURIA

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger	Whole gene screen	SNVs, indels	FMO3	56
Family member testing		as indicated 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 indicated above			14
Proforma required?		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.3460G>A, m.11778G>A, m.14484T>C)	28
Family member testing		as 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 and pyrosequencing	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 member 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
 - myoclonic epilepsy with ragged red fibres (MERRF)
 - o neuropathy, ataxia and retinitis pigmentosa (NARP)
 - maternally inherited Leigh syndrome (MILS)

- Clinical Genetics
- Endocrinology
- Metabolic
- Neurology
- Ophthalmology





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

Available testing

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

Referral criteria

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

- Clinical Genetics
- Endocrinology
- Metabolic
- Neurology





MITOCHONDRIAL INHERITED DIABETES AND DEAFNESS (MIDD)

Available testing

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

Referral criteria

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

- Clinical Genetics
- Endocrinology
- Metabolic





MUSCULOSKELETAL

BECKER MUSCULAR DYSTROPHY (BMD)

Available testing

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

Referral criteria

- Clinical features that include:
 - o Progressive symmetric muscle weakness
 - o Increase in serum concentration of creatine kinase (CK)
 - Calf hypertrophy
 - o 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 me	mber testing		as indica	ated above	14
Proforma required?		NO			

Referral criteria

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

Requesting specialties

Clinical Genetics





DUCHENNE MUSCULAR DYSTROPHY (DMD)

Available testing

Centre	Method	Scope and range of test		Targets	TAT
	MLPA	Targeted screen	Exon level CNV	DMD	28
Glasgow	Sanger	Whole gene screen	SNVs, indels	DMD	56
Family me testing	mber	as indicated above			14
Proforma r	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Paediatrics
- Neurology





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

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	Sanger	Targeted screen	SNVs, indels	FGFR3 (exons 7, 10, 13, 15, 19)	28
Family mem	Family member testing as inc		as ind	cated above	14
Proforma re	quired?	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	<i>ACVR1</i> (p.R206H)	28
,	Family member as inc		icated above	14	
Proforma re	equired?	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	Targets	TAT	
Glasgow	NGS	Whole gene screen	Whole gene screen SNVs, indels		
	member as indicated above				
Proforma re	equired?	NO			

Referral criteria

• Growths of multiple osteochondromas

- Clinical Genetics
- Orthopaedics
- Paediatrics





LIMB GIRDLE MUSCULAR DYSTROPHY (LGMD)

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	Sanger	Whole gene screen	SNVs, indels	DES, FKRP, LMNA	56
Family member as ind testing		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 myotonic dystrophy type 1.
- Suggestive features include
 - o Hypotonic infant with or without joint contractures
 - o Muscle myotonia
 - o Muscle weakness
 - o Presenile cataracts
 - o Temporal muscle wasting and / or frontal balding
 - o Adverse anaesthetic reaction
 - o Family history of Myotonic Dystrophy
 - Unexplained excessive somnolence or cardiac conduction system abnormalities with additional features as above.

- Clinical Genetics
- Neurology
- Ophthalmology
- Paediatrics





MYOTONIC DYSTROPHY TYPE 2 (DM2)

Available testing

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

Referral criteria

Clinical phenotype consistent with a diagnosis of Myotonic Dystrophy Type 2

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

Requesting specialties

- Clinical Genetics
- Neurology

OCULOPHARYNGEAL MUSCULAR DYSTROPHY

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	PCR and Sanger	Targeted screen	Repeat expansion, SNV	PABPN1 – GCN repeat expansion and c.35G>C	28
Proforma re	equired?	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 CNVs*	BMP1, COL1A1*, COL1A2*, CREB3L1, CRTAP, FAM46A, FKBP10, IFITM5, KDELR2, P3H1 (LEPRE1), PLOD2, PLS3, PPIB, SERPINF1, SERPINH1, SP7, SPARC, TMEM38B, WNT1	112
Family me	mber testing		as indicated above		14
Proforma required? NO					

Referral criteria

- Multiple fractures of long bones without significant trauma AND at least two of the following:
 - Wormian bones
 - o Blue / grey sclera
 - Hearing loss
 - o Ribs, broad and breaded, thin & irregular
 - Short stature
 - 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, FAM20C, FERMT3, LEMD3, LRP5, OSTM1, PTH1R, RASGRP2, SNX10, SOST, TCIRG1, TGFB1, TNFRSF11A, TNFSF11, TYROBP	112
	member ting			as indicated above	14
Proforma re	equired?	NO			

Referral criteria

• Characteristic radiographic changes

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





PRIMORDIAL DWARFISM, MICROCEPHALY

Available testing

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

Referral criteria

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

Requesting specialties

Clinical Genetics

PROXIMAL SYMPHALANGISM

Available testing

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

Referral criteria

• Clinical features strongly suggestive of proximal symphalangism

Requesting specialties

• Clinical Genetics





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, RRAS2, RIT1, SHOC2, SOS1, SOS2, SPRED1, SPRED2	112
Family men	nber testing		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
 - Short stature
 - Developmental disability
- At least 1 of:
 - Cardiomyopathy
 - o Congenital heart disease
 - o Arrhythmia
 - Suggestive malignancy (bladder carcinoma, Rhabdomyosarcoma, Leukaemia, phaeochromocytoma)
 - Skin abnormalities (hyperkeratosis, café au lait patches, ulerythema oophorogenes, keratosis pilaris, excess palmar skin)

- Clinical Genetics
- Paediatrics





SHORT STATURE, INCLUDING TURNER SYNDROME

Available testing

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

Referral criteria

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

Other specific features may include

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

- Clinical Genetics
- Paediatrics





SKELETAL DYSPLASIA

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

Available testing

Centre	Method	Scope and range 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, COL2A1, COL9A1, COL9A2, 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 member			as i	ndicated above	14
testii	ng				
Proforma red	quired?	NO			

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	NGS	Whole gene screen	SNVs, indels	Nail-patella syndrome: LMX1B	112
Family member as in testing		ndicated above	14		
Proforma red	quired?	NO			

Referral criteria

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





- 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	Proforma required? NO				

Referral criteria

• Clinical features that indicate a likely diagnosis of SBMA

- Clinical Genetics
- Neurology





NEUROLOGY

AICARDI-GOUTIERES SYNDROME

Available testing

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

Referral criteria

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

- Clinical Genetics
- Neurology





NEUROMUSCULAR ARTHROGRYPOSIS

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	NGS	Whole gene screen	SNVs, indels	ACTA1, ADAMTS10, ANTXR2, ASCC1, ASXL1, B3GALNT2, B4GAT1, BICD2, CHAT, CHRNA1, CHRNB1, CHRND, CHRNE, CHRNG, CHST14, CNTNAP1, COL12A1, COL6A1, COL6A2, COL6A3, COLQ, DAG1, DNM2, DOK7, DPAGT1, DYNC1H1, ECEL1, ERCC6, ERCC8, EXOSC3, FAM20C, FBN2, FGFR2, FKBP10, FKRP, FKTN, GBA, GBE1, GLDN, GLE1, GMPPB, ADGRG6, HSPG2, ISPD, KLHL40, KLHL41, LAMA2, LARGE1, LMOD3, MAGEL2, MPZ, MTM1, MUSK, MYBPC1, MYH2, MYH3, MYH7, MYH8, NALCN, NEB, PEX1, PEX10, PEX11B, PEX12, PEX13, PEX14, PEX16, PEX19, PEX2, PEX26, PEX3, PEX5, PEX6, PEX7, PFKM, PIEZO2, PLOD1, PLOD2, POMGNT1, POMGNT2, POMK, POMT1, POMT2, POR, PRG4, RAPSN, RYR1, SCARF2, SCN4A, SKI, SLC5A7, SMAD4, STAC3, SYNE1, TMEM5, TNNI2, TNNT1, TNNT3, TPM2, TPM3, TRPV4, TSEN54, UBA1, VAMP1, VIPAS39, VPS33B, ZC4H2	112
Family member testing				as indicated above	14
Proforma r	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.

- Clinical Genetics
- Neurology





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 in	dicated above	14
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Neurology





CAPILLARY MALFORMATIONS

Available testing

Centre	Method	Scope and ra	ange of test	Targets	TAT
Dundee	Sanger MLPA	Whole gene screen SNVs, indels Exon level CNV		RASA1	56
_	member ting		as indicated	above	14
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Dermatology
- Neurology





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

Available testing

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

Referral criteria

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

- Clinical Genetics
- Neurology





COMMON CRANIOSYNOSTOSIS SYNDROMES

Available testing

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

Referral criteria

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

Requesting specialties

• Clinical Genetics





CORTICAL BRAIN MALFORMATIONS

Available testing

Centre	Method	Scope and rang	ge of test	Targets	TAT
Dundee	NGS (clinical exome)	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, WDR62	112
_	member ting		as indicated above		14
Proforma r		NO			I

Referral criteria

• Cortical brain malformation with features suggestive of a monogenic cause

- Clinical Genetics
- Neurology





CREUTZFELDT-JAKOB DISEASE (CJD)

Available testing

Centre	Method	Scop	pe and range of test	Targets	TAT
Edinburgh	PCR	Repeat-primed PCR	Octapeptide repeat expansion	PRNP	28
Edinburgh	Sanger	Whole gene screen	SNVs, indels	PRNP	56
Family n		as indicated above			14
Proforma required?		NO			

Referral criteria

• Clinical features that indicate a likely diagnosis of CJD

- Clinical Genetics
- Neurology





DEMENTIA

Available testing

Centre	Method	Scope and	range of test	Targets	TAT
Edinburgh	PCR	Targeted screen	Hexanucleotide repeat expansion	c90RF72	28
Dundee	NGS (clinical exome)	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
,	amily member as indicated above testing		d above	14	
Proforma re	equired?	NO			

Referral criteria

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

- Clinical Genetics
- Neurology





DENTATORUBRAL PALLIODOLUYSIAN ATROPHY (DRPLA)

Available testing

Centre	Method	Scope a	nd 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 (clinical exome)	Whole gene screen	SNVs, indels	ACTB, AFG3L2, ANO3, APTX, ATM, ATP1A2, ATP7B, C19orf12, CACNA1A, CHMP2B, CP, CSF1R, CYP27A1, DCAF17, FBXO7, FTL, GFAP, GNAL, HPCA, LYST, NKX2-1, PANK2, PDE10A, PDGFB, PDGFRB, PNKD, PRKRA, PRNP, PRRT2, RNF216, SGCE, SLC19A3, SLC20A2, SLC2A1, SPR, TBK1, THAP1, TIMM8A, TOR1A, TUBB4A, WDR45	112
,	member tina		as indicated above		14
Proforma r		NO			

Referral criteria

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





EPILEPSY

Available testing

Centre	Method	Scope and ran	ge of test	Targets	TAT
Glasgow	Microarray	Whole genome screen	CNV	Whole genome	28
Glasgow	NGS	Whole gene 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 me	mber testing			as indicated above	14
Proforma re	equired?	YES See epileps	y referral form	(see centre website)	•

Referral criteria

• Unexplained epilepsy with clinical suspicion of a monogenic cause.

- Clinical Genetics
- Neurology
- Paediatrics





EPISODIC ATAXIA

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	Sanger	Whole gene screen	SNVs, indels	KCNA1, CACNA1A	56
,			dicated above	14	
testing					
Proforma re	Proforma required? NO				

Referral criteria

• Paroxysmal attacks of ataxia and vertigo and/or nausea

Requesting specialties

- Clinical Genetics
- Neurology

EPISODIC MOVEMENT, MIGRAINE & EPILEPTIC DISORDERS (BRAIN CHANNELOPATHIES)

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS (clinical exome)	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 member testing			as indicated above	14
Proforma r		NO			

Referral criteria

 Unexplained clinical phenotype associated with a brain channelopathy 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 (targeted panel)	Whole gene screen	SNVs, indels, exon level CNVs	KRIT1, CCM2, CCM3	56
Family member testing			as indicate	ed above	14
Proforma re	equired?	NO			

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Neurology

FAMILIAL HEMIPLEGIC MIGRAINE

Available testing

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

Referral criteria

• Hereditary ataxia with onset in adulthood

Requesting specialties

- Neurology
- Clinical Genetics

FRIEDRICH ATAXIA (FRDA)

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	PCR & 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

Comtra	Mathad	C	l versus of toot	Townsto	TAT
Centre	Method	Scope and	range of test	Targets	TAT
Edinburgh	PCR & TP-PCR	Targeted screen	Triplet repeat expansions	SCA1, SCA2, SCA3, SCA6, SCA7, FRDA, FMR1	28
Edinburgh *	NGS	Whole gene screen	SNVs, indels	AAAS, ABCBT, ABCD1, ADAR, AFG3L2, ALS2, ANO10, APTX, ATL1, ATM, ATP1A3, ATP7B, BSCL2, CACNA1A, CACNA1G, CAPN1, COQ8A, CYP27A1, CYP7B1, DDHD2, FA2H, FGF14, FTL, FXN, GBA2, GCH1, GRID2, HSPD1, IFIH1, ITPR1, KCNA1, KCNC3, KCND3, KIF1A, KIF5A, L1CAM, NIPA1, OPA3, PDYN, PLP1, PNPLA6, POLG, PRKCG, PRNP, PRRT2, REEP1, RTN2, RNaseH2B, SACS, SETX, SIL1, SLC1A3, SLC2A1, SPART, SPAST, SPG11, SPG21, SPG7, SPTBN2, STUB1, SYNE1, TGM6, TMEM240, TTBK2, TTPA, TWNK, UBAP1, VPS13D, WASHC5, ZFYVE26	112
Glasgow*	NGS	Whole 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 m				as indicated above	14
Proforma re	<u> </u>	NO			l .
Proforma required?		110			

^{*}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
 - Normal/routine neurological bloods, and vitamin E testing
 - Negative spinocerebellar ataxia repeat expansion panel, including FXTAS and FRDA
 - o MRI neuroimaging normal, or isolated cerebellar atrophy
 - Family history of ataxia, or young age of onset (<50)

- Clinical Genetics
- Neurology

^{*}For patients referred from West of Scotland, testing performed in Glasgow





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

Available testing

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

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Neurology

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

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Aberdeen	Sanger MLPA	Whole Gene screen SNVs, indels Exon level CNV		PMP22	56
Family member testing			as indicat	red above	14
Proforma required?		NO			

Referral criteria

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

- Clinical Genetics
- Neurology





HEREDITARY SPASTIC PARAPLEGIA (HSP)

Available testing

Centre	Method	Scope and	range of test	Targets	TAT		
Edinburgh*	NGS MLPA	Whole gene screen	SNVs, indels CNV**	AAAS, ABCBT, ABCD1, ADAR, AFG3L2, ALS2, ANO10, APTX, ATL1**, ATM, ATP1A3, ATP7B, BSCL2, CACNA1A, CACNA1G, CAPN1, COQ8A, CYP27A1, CYP7B1, DDHD2, FA2H, FGF14, FTL, FXN, GBA2, GCH1, GRID2, HSPD1, IFIH1, ITPR1, KCNA1, KCNC3, KCND3, KIF1A, KIF5A, L1CAM, NIPA1, OPA3, PDYN, PLP1, PNPLA6, POLG, PRKCG, PRNP, PRRT2, REEP1**, RNaseH2B, RTN2, SACS, SETX, SIL1, SLC1A3, SLC2A1, SPART, SPAST**, SPG11, SPG21, SPG7**, SPTBN2, STUB1, SYNE1, TGM6, TMEM240, TTBK2, TTPA, TWNK, UBAP1, VPS13D, WASHC5 and ZFYVE26	112		
Glasgow*	NGS 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 member			I	as indicated above	14		
Proforma red	J	YES Ed	inburah onlv – H	n only – HSP referral proforma (see centre website)			

^{*} For patients referred from East of Scotland, testing performed in Edinburgh

Referral criteria

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

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

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

- Clinical Genetics
- Neurology

^{*} 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 (clinical exome)	Whole gene screen	SNVs, indels	CDON, DHCR7, DISP1, FGF8, FGFR1, GLI2, PTCH1, SHH, SIX3, TGIF1, ZIC2	112
Family member testing			as in	dicated above	14
Proforma required?		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		HTT	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)





HUNTINGTON DISEASE-LIKE disorders

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	Flanking PCR	Targeted screen	Triplet repeat expansion	JPH3	28
Edinburgh	Sanger* Flanking PCR and TP- PCR	Targeted screen	SNVs and Indels Triplet repeat expansion	FTL*, C9orf72, PRNP, TBP, ATN1, JPH3	28
Proforma re	quired?	NO		<u> </u>	

Referral criteria

- Clinical features that indicate a likely diagnosis of Huntington disease-like 2 (for *JPH3* testing)
- HD testing has been completed
- For patients with a HD-like phenotype, a screen including testing for *C9orf72*, *PRNP*, SCA17 (*TBP*), DRPLA (*ATN1*) and HDL2 (*JPH3*) repeat expansions, and sequencing of FTL, is also available

Requesting specialties

Clinical Genetics





LESCH-NYHAN SYNDROME

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	Sanger MLPA	Whole Gene screen	SNVs, indels Exon level CNV	HPRT1	56
Family me	Family member testing as indicated a			bove	14
Proforma re	quired?	NO			

Referral criteria

- Hyperuricaemia
- Psychomotor delay
- Mild to moderate intellectual disability
- Self-injurious behavior

Requesting specialties

- Clinical Genetics
- Neurology

LEUKOENCEPHALOPATHY WITH VANISHING WHITE MATTER

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS (clinical exome)	Whole gene screen	SNVs, indels	EIF2B1, EIF2B2, EIF2B4, EIF2B5, EIF2B3	112
Family member as in testing		dicated above	14		
Proforma re	equired?	NO			

Referral criteria

 Individuals with unexplained leukodystrophy on neuroimaging with onset in adulthood

- Clinical Genetics
- Neurology





NEURODEGENERATION WITH BRAIN IRON ACCUMULATION (NBIA)

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	NGS	Whole 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 member testing			as indicated above	14
Proforma re	equired?	NO			

Referral criteria

• Suspected clinical diagnosis in patients with hallmark findings of NBIA, or further assessment of patients with clinical diagnosis of idiopathic NBIA who have had mutations ruled out in other genes.

- Clinical Genetics
- Neurology





NEUROFIBROMATOSIS TYPE 1 (NF1)

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS (targeted panel)	Whole gene screen	SNVs, indels Exon level CNV	NF1	56
Family r			as indicat	ed above	14
Proforma re	equired?	NO			

Referral criteria

- Clinical diagnosis of NF1, as defined below, AND molecular diagnosis is required for management of the proband or for reproductive planning
- Diagnosis requires two of:
 - At least 6 café au lait macules (at least 0.5cm in a child and 1.5cm in an adult)
 - o At least 2 subcutaneous or cutaneous neurofibromas
 - o Plexiform neurofibroma
 - o Optic glioma
 - o At least 2 Lisch nodules
 - 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 (clinical exome)	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
,	nily member testing			as indicated above	14
Proforma re	equired?	NO			

Referral criteria

- This includes the disorders:
 - o Congenital insensitivity to pain
 - o Inherited erythromelalgia
 - o Paroxysmal extreme pain disorder
 - Small fibre neuropathy
 - o Familial episodic pain syndromes
 - o Hereditary sensory and autonomic neuropathies
 - o Forms of Hereditary sensory neuropathy with prominent sensory loss
- Individuals with a disorder of pain perception, including insensitivity to pain or increased pain perception that is likely to be monogenic in aetiology

- Clinical Genetics
- Neurology





PARKINSON'S DISEASE

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS (clinical exome)	Whole gene screen	SNVs, indels, Exon level CNV	ATP13A2, ATP1A3, DCTN1, DNAJC13, DNAJC6, FBXO7, FTL, , 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				as indicated above	14
tes	ting				
Proforma r	equired?	No			

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 test	Targets	TAT
Dundee	Sanger MLPA	Whole gene screen	SNVs, indels Exon level CNV (<i>PLP1</i>)	PLP1, GJC2(GJA12)	56
Family member as indicated testing			above	14	
Proforma re	quired?	NO			

Referral criteria

- Any individual with clinical or imaging features suggestive of a *PLP1* disorder
- Pathogenic variants in *GJC2* are associated with Pelizaeus-Merzbacher-like disease, an autosomal recessive disorder.

- Clinical Genetics
- Neurology





PERIODIC PARALYSIS, HYPERKALAEMIC

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	Sanger	Targeted screen	SNVs	SCN4A (p.Leu689lle, p.lle693Thr, p.Thr704Met and p.Met1592Val)	56
Family member testing			as indicated above	14	
Proforma re	quired?	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	14	
Proforma required? NO					

Referral criteria

Two or more attacks of muscle weakness with documented serum potassium
 3.5 mmol/L

OR

- One attack of muscle weakness and one attack of weakness in one relative with documented serum potassium <3.5 mmol/L OR
- Three or more of the following six clinical/laboratory features:
 - o Onset in the first or second decade
 - Duration of attack (muscle weakness involving ≥1 limbs) longer than two hours
 - The presence of triggers (previous carbohydrate rich meal, symptom onset during rest after exercise, stress)
 - o Improvement in symptoms with potassium intake
 - A family history of the condition or genetically confirmed skeletal calcium or sodium channel mutation
 - Positive long exercise test

AND

• Exclusion of other causes of hypokalemia (renal, adrenal, thyroid dysfunction; renal tubular acidosis; diuretic and laxative abuse)

- Clinical Genetics
- Neurology





PERIPHERAL NEUROPATHY

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	NGS	Whole gene screen	SNVs, indels	AARS, ATL1, ATP7A, BICD2, BSCL2, CCT5, DCTN1, DNM2, DNMT1, DYNC1H1, EGR2, FAM134B, FGD4, FIG4, GARS, GDAP1, HINT1, HSPB1, HSPB3, HSPB8, IGHMBP2, IKBKAP, INF2, KIF1A, LITAF, LMNA, LRSAM1, MARS, MTMR2, NDRG1, NEFL, NGF, NTRK1, PLEKHG5, PRPS1, PRX, RAB7A, REEP1, SBF2, SCN9A, SETX, SH3TC2, SLC52A1, SLC52A2, SLC52A3, SORD, SPTLC1, SPTLC2, TRPV4, VCP, WNK1, YARS	112
Family member testing			as indicated above	14	
Proforma r	equired?	NO			

Referral criteria

- Length dependent neuropathy on neurophysiology AND
- No pathogenic variant on first tier CMT testing (performed in Aberdeen)

AND one of the following-

- Genetic diagnosis will alter clinical management
- Genetic diagnosis will influence reproductive decisions

- Clinical Genetics
- Neurology





PORENCEPHALY

Available testing

Centre	Method		Scope and range of	Targets	TAT	
Dundee	NGS (clinical exome)	١	Whole gene screen	SNVs, indels	COL4A1, COL4A2	112
Family me	Family member testing		as	indicated above		14
Proforma re	Proforma required?					

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
	member ting			as ind	icated above	14
Proforma re	equired?	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





RHABDOMYOLYSIS & METABOLIC MYOPATHIES

Available testing

Centre	Method	Scope an	d range of test	Targets	TAT
Glasgow	NGS	Whole gene screen	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
_	member ting			as indicated above	14
Proforma r	equired?	NO			

This panel is intended for patients with isolated skeletal muscle symptoms. Patients with multisystem disease may be more appropriately tested on alternative panels

Referral criteria

Single episode rhabdomyolysis

- ALL MUST FULFIL 2 essential criteria:
 - o CK documented >10,000IU/L associated with muscle pain
 - 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:
 - No environmental cause AND Accustomed exercise (NOT too much, too fast, too soon)
 - High risk features- exercise intolerance preceding rhabdo +/OR weakness on examination >4mths after event +/OR family history documented rhabdo +/OR biochemistry classical of VLCAD, MADD, or CPT2 +/OR cardiomyopathy
 - o CK>500 IU/L >6 months after rhabdo episode

Recurrent rhabdomyolysis

- All must fulfil 3 essential criteria:
 - o CK documented >10,000IU/L associated with muscle pain on at least one occasion
 - At least one further episode of acute muscle pain associated with documented CK rise or pigmenturia
 - Mitochondrial myopathy/PEO considered and excluded where appropriate

Other criteria for rhabdo panel testing

- · Clinical suspicion metabolic myopathy AND any of
 - Moderate to profound XS lipid or glycogen on biopsy
 - o Cores/minicores on biopsy
- Muscle MRI characteristic of RYR1

- Clinical Genetics
- Metabolic
- Neurology





SPINAL MUSCULAR ATROPHY

Available testing

Centre	Method	Scope and rai	nge of test	Targets	TAT
Edinburgh	MLPA	Targeted screen	CNV	SMN1	28
Glasgow	NGS	Whole gene 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, SORD, SYT2, TRPV4, UBA1, VAPB, VCP, VRK1	112
Family member			as ir	ndicated above	14
testing					
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)
- clinical features point to a peripheral cause, i.e. particularly where the baby is alert and responsive and the floppiness appears static over a period of days
- o Carrier testing for partners of confirmed SMN1 carriers.
- Whole gene screen
 - dHMN/SMA clinical phenotype AND
 - o Compatible neurophysiology (not required in infants) AND
 - 5q linked SMA excluded (not required in infants)

- Clinical Genetics
- Neurology





SPINOCEREBELLAR ATAXIA 8 (SCA8)

Available testing

Centre	Method	Scope an	d range of test	Targets	TAT
Edinburgh	PCR & TP-PCR	Targeted screen Triplet repeat expansion		ATXN8OS	28
Proforma re	equired?	NO			

Referral criteria

• Testing only available to patients with a family history of SCA8 where the expansion has been shown to segregate with disease in the family

Requesting specialties

Clinical Genetics

Here for SMA, the two different labs are combined into one with two lists of indications. But for CMT, the two tests are listed separately and in fact one is called peripheral neuropathy. Periodic paralysis also listed separately. I think its better having them separately since they are separate tests but with the same name so they are listed one after the other. I know this format will change but still easier for everyone to understand.

^{****}Some feedback from clinicians





SPINOCEREBELLAR ATAXIA 17 (SCA17)

Available testing

Centre	Method	Scope an	d range of test	Targets	TAT
Edinburgh	PCR	Targeted screen	Triplet repeat expansion	TBP	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	ge of test	Targets	TAT
Aberdeen Dundee	PCR	Targeted screen	Deletion	<i>DYT1</i> (c.907_909del) Gene known as <i>TOR1A</i>	28
Proforma required?		NO			

Referral criteria

- DYT1 early-onset isolated dystonia should be suspected in individuals with
 - Onset of dystonia before the age of 26
 - Isolated dystonia with no other abnormalities on neurologic examination, normal routine neuroimaging, no known cause of acquired dystonia
 - 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 range of test		Targets	TAT
Dundee	NGS (targeted panel)	Whole gene screen	SNVs, indels, exon level deletions/duplications	TSC1, TSC2	56
	member ting		as indicated a	bove	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 ra	ange of test	Targets	TAT
Edinburgh	NGS	Whole gene screen	SNVs, indels	COL4A3, COL4A4, COL4A5	56
Family member testing		as	ndicated above	14	
Proforma required? NO					

Referral criteria

- Proband with haematuria and ONE of:
- 1. A first degree relative with haematuria or unexplained chronic renal failure, OR
- 2. Histological evidence following electron microscopy on renal biopsy of EITHER Alport syndrome (thickening and splitting of glomerular basement membrane +/- electron lucent areas) OR thin basement membrane disease (TBMD), OR
- 3. Clinical features of Alport syndrome (high tone sensorineural hearing loss or characteristic ophthalmic signs such as perimacular flecks or anterior lenticonus)

- Clinical Genetics
- Nephrology





BARTTER SYNDROME & GITELMAN SYNDROME

Available testing

Centre	Method		Scope and	I range of test	Targets	TAT
Dundee	NGS (clinical exome)	Whole go	ene screen	SNVs, indels Exon level CNV (CLCNKB) if appropriate	BSND, CLCNKB, KCNJ1, SLC12A1, SLC12A3	56
,	member ting		as indicated above			
Proforma re	equired?	YES	Renal Genetics Proforma (see centre website)			

Referral criteria

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

Requesting specialties

- Clinical Genetics
- Nephrology

CYSTINURIA

Available testing

Centre	Method		Scope and	I range of test	Targets	TAT	
Dundee	NGS (clinical exome)	Whole go	ene screen	SNVs, indels	SLC3A1, SLC7A9	56	
	member ting		as indicated above				
Proforma re	equired?	YES	YES Renal Genetics Proforma (see centre website)				

Referral criteria

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

- Clinical Genetics
- Nephrology





NEPHROCALCINOSIS OR NEPHROLITHIASIS

Available testing

Centre	Method	Scope and range of test		st Targets	TAT	
Dundee	NGS (clinical exome)	Whole ge screer		AGXT, APRT, ATP6V0A4, 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	
,	member ting			as indicated above	14	
Proforma r	equired?	YES	Renal Genetic	Renal Genetics Proforma (see centre website)		

Referral criteria

Nephrocalcinosis or nephrolithiasis where acquired causes have been excluded

- Clinical Genetics
- Nephrology
- Endocrinology





POLYCYSTIC KIDNEY DISEASE, AUTOSOMAL DOMINANT AND RECESSIVE

Available testing

Centre	Method	Scope	Scope and range of test		Targets	TAT
Dundee	NGS (targeted panel)	Whole ge		SNVs, indels Exon level CNV	PKD1, PKD2	56
Dundee	NGS (targeted panel)	Whole ge screen		SNVs, indels	PKHD1	56
Dundee	NGS (targeted panel)	Whole ge screer		SNVs, indels	AGT, ALG8, ALG9, ANKS6, CEP164, CEP83, COL4A1, DNAJB11, DZIP1L, GANAB, HNF1B, IFT140, 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 required? Yes		Yes		al Genetics Profo KHD1 only.	rma (see centre website). No proforma needed for Pk	(D1/2

Referral criteria

- For Autosomal Dominant Polycystic Kidney Disease: Individuals with a suspected or established diagnosis of Autosomal Dominant Polycystic Kidney Disease based on renal imaging.
 - Initial analysis of PKD1 and PKD2 then further analysis of the full cystic kidney panel if appropriate.
- Individuals with a suspected or established diagnosis of Autosomal Recessive Polycystic Kidney Disease based on renal imaging or pathology.
- Onset is typically prenatal, in infancy or early childhood/young adulthood
- The full cystic kidney disease full panel is recommended for individuals that meet the following criteria:
 - Patients with non-syndromic cystic renal disease (excluding acquired cystic disease due to chronic or end stage kidney disease) which is EITHER
 - Clinically not characteristic of ADPKD and underlying diagnosis is required for management purposes, OR
 - 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 LIVER DISEASE

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS (targeted panel)	Whole gene screen	SNVs, indels	AGT, ALG8, ALG9, ANKS6, CEP164, CEP83, COL4A1, DNAJB11, DZIP1L, GANAB, HNF1B, IFT140, 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	equired?	Yes	Renal Genetics Pr	roforma (see centre website)	

Referral criteria

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

Note this is the same panel as the full polycystic kidney disease panel Requesting specialties

- Clinical Genetics
- Fetal Medicine
- Nephrology
- Paediatrics





PRIMARY HYPEROXALURIA

Available testing

Centre	Method	Scope and range of test		nge of test	Targets	TAT
Dundee	NGS (clinical exome)		e gene een	SNVs, indels	AGXT, GPHPR, HOGA1	56
,	member ting			as indi	cated above	14
Proforma r	equired?	YES	Renal Ge	enetics Proforma (s	see centre website)	

Referral criteria

- Any individual with clinical and biochemical features consistent with the condition.
- Overlapping conditions: Nephrocalcinosis or nephrolithiasis

Requesting specialties

- Clinical Genetics
- Nephrology

PSEUDOHYPOALDOSTERONISM type 1

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS (clinical exome)	Whole gene screen	SNVs, indels	NR3C2, SCNN1A, SCNN1B, SCNN1G	112
Family member testing			as indi	cated above	14
Proforma re	equired?	NO			

Referral criteria

 Any individual with clinical and biochemical features consistent with the condition.

- Clinical Genetics
- Paediatrics





RENAL CILIOPATHY

Available testing

Centre	Method	Scope and range of test		Targets	TAT	
Dundee	NGS (clinical exome)	Whole gene screen	SNVs, indels,	AHI1, ALMS1, ANKS6, ARL13B, ARL6, B9D2, BBS1, BBS10, BBS12, BBS2, BBS4, BBS5, BBS7, BBS9, C2CD3, C5orf42 (CPLANE1), CC2D2A, CEP164, CEP290, CEP41, CEP83, CRB2, CSPP1, DDX59, DHCR7, DYNC2H1, HNF1B, HYLS1,ICK, IFT122, IFT43, INVS, IQCB1, KIF7, LZTFL1, MKKS, MKS1, NEK8, NPHP1, NPHP3, NPHP4, OFD1, PKD1, PKD2, PKHD1 PMM2, RPGRIP1L, SDCCAG8, TCTN1, TCTN2, TCTN3, TMEM138, TMEM216, TMEM231, TMEM237, TMEM67, TRAF3IP1, TTC21B, TTC8, WDPCP, WDR19, WDR35, WDR60	112	
Family member testing			as indicated above	14		
Proforma r	equired?	YES	ES Renal Genetics Proforma (see centre website)			

Referral criteria

- Individuals with a suspected clinical diagnosis associated with the above genes
- Relevant medical conditions:
 - o Joubert syndrome
 - o Alstrom syndrome
 - o Bardet-Biedl syndrome
 - o Meckel syndrome
 - o Nephronophthisis
 - o Smith-Lemli-Opitz syndrome
 - o Short rib thoracis dysplasia with or without polydactyly
 - o McKusick-Kaufman syndrome
 - o Senior-Loken syndrome

- Clinical Genetics
- Nephrology





RENAL TUBULOPATHIES, RENAL TUBULAR ACIDOSIS

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS (clinical exome)	Whole gene screen	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 member testing		1	as indicated above	14	
3			Renal Genetics Pr	roforma (see centre website)	

Referral criteria

- Patients with a primary renal tubulopathy presenting as one of the following conditions:
 - Hypokalaemic alkalosis with normal or low blood pressure (e.g. Bartter/Gitelman syndromes), OR
 - Hypokalaemic alkalosis with elevated blood pressure (e.g. Liddle syndrome), OR
 - o Hyperkalaemic acidosis with low/normal BP (PHA type 1), OR
 - o Hyperkalaemic acidosis with elevated BP (PHA type 2), OR
 - o Hypokalaemic acidosis (pRTA and renal Fanconi syndromes), OR
 - o Hypomagnesaemia, OR
 - Nephrogenic diabetes insipidus, OR
 - Other rare types of renal tubulopathy seen in an expert center
- Overlapping conditions: Nephrogenic diabetes insipidus, Bartter/Gitelman syndromes and Nephrocalcinosis or nephrolithiasis

- Clinical Genetics
- Nephrology





STEROID RESISTANT NEPHROTIC SYNDROME (SRNS) AND PROTEINURIC RENAL DISEASE

Available testing

Centre	Method	Scope at	nd range of test	Targets	TAT
Dundee	NGS (clinical exome)	Whole gene screen	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, TNS2, TP53RK, TRPC6, WDR73, WT1	112
Family member			as indicated above	14	
tes	ting				
Proforma r	equired?	YES	Renal Genetics P	roforma (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	Sc	Scope and range of test		Targets	TAT
Dundee	NGS (clinical exome)		ole gene creen	SNVs, indels	ANKS6, CEP164, CEP83, GATM, HNF1B, INVS, MUC1, NPHP1, NPHP3, NPHP4, REN, TMEM67, TTC21B, UMOD, WDR19	112
Family member testing		as ind	icated above	14		
Proforma required?		YES	Renal Gen	etics Proforma (see centre website)	

Referral criteria

- Previously known as hyperuricemic nephropathy, familial juvenile, type 1 & 2 and only UMOD and REN tested. Includes both dominant and recessive TKD.
- Renal impairment caused by tubulointerstitial fibrosis with no glomerular lesion, with no identifiable cause, often associated with medullary cysts, hyperuricaemia or gout, AND
- A first degree relative with TKD or unexplained end-stage renal disease
- Testing note: the majority of pathogenic variants in the MUC1 gene are within a Variable Nucleotide Tandem Repeat (VNTR) region, these are not detectable by this method

- Clinical Genetics
- Nephrology





RESPIRATORY

ASTHMA

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	Real time PCR	Targeted screen	SNV	ADRB2 p.(Gly16Arg)	28
L		NO			

Referral criteria

- Asthma patient who may be using or about to be prescribed long acting B2 agonist therapy.
- Some evidence to suggest that homozygotes for arginine at codon 16 (ADRB2 p.(Arg16Arg)) may not benefit from long acting B2 agonist therapy

- Clinical Genetics
- Respiratory





CYSTIC FIBROSIS

Available testing

Centre	Method	Scope and ra	ange of test	Targets	TAT
Aberdeen Dundee Edinburgh Glasgow	ARMS	Targeted screen	SNVs, indels	Common variants	28 Prenatal 3
Glasgow	ARMS	Targeted screen	SNVs, indels	CFTR newborn screening (p.508del, p.G542*, p.G551D, c.469+1G>T common variants)	7
Edinburgh	NGS Sanger	Whole gene screen	SNVs, indels Exon level CNV	CFTR	56
Family member testing		as indicated above			14
Proforma required?		NO			

Referral criteria

- Test in an individual clinically likely to be affected with cystic fibrosis:
- Child with clinical suspicion of CF (e.g. recurrent chest infections, failure to thrive, fat malabsorption, neonatal history of meconium ileus), AND
 - A not normal sweat test performed in a recognised experienced test centre/laboratory (i.e. sweat chloride >40mM with sufficient sweat obtained; >30mM in infants), OR
 - An additional urgent prenatal situation for the parents or for a close relative, but urgent sweat testing not accessible
- Adult with CT-proven bronchiectasis, AND
 - A not normal sweat test performed in a recognised experienced test centre/laboratory
 (i.e. sweat chloride >40mM with sufficient sweat obtained), OR
 - Chronic suppurative chest infection with colonisation by Pseudomonas and Staph aureus, OR
 - Additional exocrine pancreatic dysfunction
- Idiopathic chronic pancreatitis with exocrine dysfunction (fat malabsorption) with other obvious and acquired causes excluded, AND
 - A not normal sweat test performed in a recognised experienced test centre/laboratory
 (i.e. sweat chloride >40mM with sufficient sweat obtained), OR
 - Sweat testing not practical, and all other causes excluded
- Male infertility associated with obstructive azoospermia, AND
 - o CBAVD (or isolated CUAVD) diagnosed from expert clinical examination, OR
 - o CBAVD identified at incidental herniotomy
- Fetal echogenic bowel as bright as bone on 2nd trimester scan, AND
 - Both parents not available for carrier testing [if both parents are available, Cystic fibrosis carrier testing should be used instead of an invasive prenatal test], AND
 - Isolated anomaly or <2 other common fetal markers, AND
 - Other more common causes excluded (e.g. IUGR, placental failure, earlier bleeding, infection, raised aneuploidy markers)

- Clinical Genetics
- GP
- Obstetrics
- Paediatrics
- Respiratory





HEREDITARY HAEMORRHAGIC TELANGIECTASIA, PRIMARY PULMONARY HYPERTENSION

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Edinburgh	HHT- 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 member testing			as ir	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
 - 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
- ENT





PRIMARY CILIARY DYSKINESIA

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	NGS	Whole gene screen	SNVs, indels	ARMC4, C210RF59, 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 i	Family member a		a	as indicated above	14
tes	ting				
Proforma re	equired?	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
 - Persistent rhinorrhoea and cough distress where other causes have been excluded
 - o Sibling with PCD
- Childhood at least one of the following:
 - Persistent lifelong wet cough (cystic fibrosis excluded)
 - Unexplained bronchiectasis (cystic fibrosis excluded)
 - Serious otitis media in association with recurrent lower and upper airway symptoms
- Adults
 - Symptoms as above since, often associated with infertility or subfertility

- Clinical Genetics
- Paediatrics
- Respiratory Medicine





SURFACTANT METABOLISM DYSFUNCTION

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Glasgow	NGS	Whole gene screen	SNVs, indels	ABCA3, NKX2-1, SFTPB, SFTPC	56
		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	Sco	pe and rang	ge of test	Targets	TAT	
Dundee	Sanger	Whole ge	ene screen	SNVs, indels	TGM5	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 Painless peeling of the epidermis
 - o Itchy and red skin
 - o Blisters

- Clinical Genetics
- Dermatology





AUTOSOMAL RECESSIVE CONGENITAL ICHTHYOSIS (ARCI)

Available testing

Centre	Method	Scope and range of test			Targets	TAT	
Dundee	NGS (clinical exome)	Whole ge screen		SNVs, indels	ABCA12, ALDH3A2, ALOX12B, ALOXE3, CERS3, CYP4F22, NIPAL4, PNPLA1, SLC27A4, ST14, STS, SULT2B1, TGM1	56	
Family member testing		as indicated above			as indicated above	28	
Proforma required?		YES	Skin disorders proforma (see centre website)				

Referral criteria

- Any individual with a clinical presentation consistent with the condition
 - o Born with collodion membrane
 - Thick, hyperkeratotic skin
 - o The later development of at least one of the following:
 - classic lamellar ichthyosis (LI)
 - (nonbullous) congenital ichthyosiform erythroderma (CIE)
 - intermediate forms with some features of both LI and CIE, or nonLI/nonCIE form with mild hyperkeratosis
 - o Excludes Harlequin ichthyosis

- Dermatology
- Clinical Genetics





BIRT-HOGG-DUBE SYNDROME

Available testing

Centre	Method	Scope and range of test			Targets	TAT	
Dundee	NGS (targeted panel)	Whole gene screen		SNVs, indels Exon level CNV	FLCN	56	
Family member testing		as indicated above					
Proforma required?		NO					

Referral criteria

- Individuals with either:
 - five or more facial or truncal papules with at least one histologically confirmed fibrofolliculoma
- or two of:
 - o early-onset [age <50 years] or multifocal/bilateral renal cell cancer
 - o renal cell cancer with mixed chromophobe/oncocytic histology
 - o multiple lung cysts with or without spontaneous pneumothorax
 - o first degree relative with BHDS

- Clinical Genetics
- Dermatology
- Respiratory





BULLOUS CONGENITAL ICHTHYOSIFORM ERYTHRODERMA

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Dundee	Sanger		e gene een	SNVs, indels	KRT1, KRT10	56
_	member ting			as in	dicated above	28
Proforma re	equired?	YES	Skin diso	rders proforma (see centre website)	

Referral criteria

- Also known as Epidermolytic hyperkeratosis (EHK) or Epidermolytic ichthyosis (EI)
- Any individual with a clinical presentation consistent with the condition:
 - Hyperkeratotic scaliness
 - Severe blistering
 - o Hyperproliferation in the basal cells
 - o Thickened, granular layer of the epidermis
 - Skin biopsy recommended if mosaic form suspected (epidermolytic epidermal naevus)

- Clinical Genetics
- Dermatology





ECTODERMAL DYSPLASIA

Available testing

Centre	Method	Scope an	d range of test	Targets	TAT	
Dundee	NGS (clinical exome)	Whole gen screen	e 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	
	member ting			as indicated above	28	
Proforma r		YES S	YES Skin disorders proforma (see centre website)			

Referral criteria

- Any individual with a clinical diagnosis of ectodermal dysplasia with one or more of the following:
 - abnormality of hair (hypotrichosis, sparse hair, sparse/missing eyebrows)
 - abnormality of teeth (hypodontia, conical incisors)
 - o abnormality of skin (hypohidrosis, episodes of hyperthermia)
- Includes Hypohidrotic X-linked Ectodermal Dysplasia (XHED), Anhidrotic (autosomal dominant and recessive) Ectodermal Dysplasia, Odontoonychodermal Dysplasia (OODD), Clouston syndrome, Witkop syndrome, and Ectrodactyly, Ectodermal Dysplasia and Cleft Lip/Palate syndrome (EEC3)

- Clinical Genetics
- Dermatology





EPIDERMOLYSIS BULLOSA

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	Sanger	Whole gene screen	SNVs, indels	KRT5, KRT14	56
Dundee	NGS (clinical exome)	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 member testing				as indicated above	28
Proforma required? YES		Skin disorders	proforma (see centre website)		

Referral criteria

- Includes common types of Epidermolysis bullosa simplex (EBS): localized (EBS-loc, previously known as Weber-Cockayne type), generalized intermediate (EBS-gen intermed, previously known as Koebner type), motteled (EBS-MP) and generalized severe (EBS-gen sev, previously known as Dowling-Meara type)
 - Sanger sequencing for KRT5 and KRT14 for EBS
 - Dowling-Degos Syndrome Sanger sequencing for KRT5
 - Naegeli-Franceschetti-Jadassohn Syndrome Sanger sequencing for KRT14 exon 1
 - NGS test for other rarer forms of EB
- Genetically heterogeneous disorder of skin fragility, manifested by blistering and/or erosions with little or no trauma

- Clinical Genetics
- Dermatology





EPIDERMOLYTIC PALMOPLANTAR KERATODERMA (EPPK)

Available testing

Centre	Method	Scope and range of test			Targets	TAT
Dundee	Sanger		e gene een	SNVs, indels	KRT1, KRT9	56
_	Family member testing		as in	dicated above	28	
Proforma re	equired?	YES	Skin Disc	orders Proforma	(see centre website)	

Referral criteria

- Any individual with a clinical presentation consistent with the condition:
 - Yellow and diffuse thickening of the skin on the palms and soles (palmoplantar keratoderma)
 - o Erythema
 - o Localised epidermolytic hyperkeratosis
 - Onset in infancy

- Clinical Genetics
- Dermatology





FERGUSON-SMITH DISEASE

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	Sanger	Whole gene screen	SNVs, indels	TGFBR1	56
•	Family member as testing		as ind	icated above	28
Proforma re	equired?	NO			

Referral criteria

- Any individual with a clinical presentation consistent with the condition:
 - Squamous cell carcinomas or keratoacanthoma which heal spontaneously leaving pitted scars

Requesting specialties

- Clinical Genetics
- Dermatology

FOCAL PALMOPLANTAR KERATODERMA

Available testing

Centre	Method	Scope and rang	ge of test	Targets	TAT
Dundee	Sanger	Targeted screen	SNVs, indels	KRT6C (ex1&7), KRT16 (ex1,6,7,8)	56
Family men	nber testing		as in	dicated above	28
Proforma required? 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 rai	nge of test	Targets	TAT
Dundee	Sanger	Targeted screen	SNVs, indels	GLMN (exons 2, 3, 6, 8, 12, 13)	56
Family member testing			as ind	icated above	28
Proforma re	equired?	NO			

Referral criteria

- A clinical diagnosis of glomuvenous malformations (GVM) based on the International Society for the Study of Vascular Anomalies (ISSVA) classification
- Two or more combined malformations consisting of capillary and venous malformations found in one lesion

- Clinical Genetics
- Dermatology





HAIR DISORDERS

Available testing

Centre	Method	Scope and range of test		Targets	TAT
Dundee	NGS (clinical exome)	Whole gene 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
,	member ting			as indicated above	28
Proforma r	equired?	YES	Skin Disorders Pro	oforma (see centre website)	

Referral criteria

- Includes Hypotrichosis Simplex, Marine Unna Hypotrichosis, Familial Woolly Hair (WFH), Hypotrichosis with Juvenile Macular Dystrophy, Netherton Syndrome, Monilethrix, Clouston Syndrome, Menkes Syndrome, Hypohidrotic Ectodermal Dysplasia (HED), Trichothiodystrophy (TTD), Ectodermal Dysplasia-9 (ECTD9), Alopecia Universalis Congenita (ALUNC), Naxos Syndrome, CHAND Syndrome, and Atrichia with papular lesions (APL).
- Individuals with a hair disorder with a likely monogenic cause

- Clinical Genetics
- Dermatology





ICHTHYOSIS & ERYTHROKERATODERMA

Available testing

Centre	Method	Scope a	and ra	ange of test	Targets	TAT
Dundee	NGS (clinical exome)	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
_	member tina				as indicated above	28
Proforma re		YES	YES Skin disorders proforma (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
 - 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	Targeted screen SNVs, indels		<i>FLG</i> (p.Arg501*; c.2282_2285delCAGT, p.Arg2447*; p.Ser3247*)	28
Family member as ind testing		as indicate	ed above			28
Proforma required? Y		YES	Skin dis	sorders proforma	(see centre website)	

Referral criteria

- Any individual with a clinical presentation consistent with the condition
 - o 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 r	ange of test	Targets	TAT
Dundee	Sanger MLPA	Whole gene SNVs, indels screen Exon level CNV		SPRED1	56
Family member testing		as indicated above			28
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 inguinal freckling
 - o No other NF1-related criteria

- Clinical Genetics
- Dermatology





MULTIPLE CUTANEOUS AND MUCOSAL VENOUS MALFORMATIONS

Available testing

Centre	Method	Scope and ran	ge of test	Targets	TAT
Dundee	Sanger	Targeted screen	SNVs, indels	TIE2 exon 15, exon 17	28
Family member testing			as in	dicated above	28
Proforma re	equired?	NO			

Referral criteria

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

- Dermatology
- Clinical Genetics





PACHYONYCHIA CONGENITA

Available testing

Centre	Method	Scope and range of test		Targets	TAT	
Dundee	Sanger	Targeted screen	SNVs, indels	KT6A (ex1&7) KRT6B (ex1&7) KRT6C (ex1&7) KRT16 (ex1,6,7&8) KRT17 (ex1,6&7)	56	
Family member testing		as indicated above				
Proforma required?		Yes – SKIN DISORDERS PROFORMA				

Referral criteria

- Any individual with a clinical presentation consistent with the condition:
 - o Plantar keratoderma including callus with underlying blisters
 - o Plantar pain
 - Hypertrophic nail dystrophy, often present within the first few months of life
 - Oral leukokeratosis

- Dermatology
- Clinical Genetics





PALMOPLANTAR KERATODERMAS

Available testing

Centre	Method	Scope and range of test		Targets	TAT	
Dundee	Sanger	Targeted screen	SNVs, indels	KRT1, KRT5, KRT9, KRT10	56	
Dundee	NGS(clinical exome)	Whole gene screen	SNVs, indels	AAGAB, ABCA12, ABHD5, ADAM17, ALDH3A2, ALOX12B, ALOXE3, AP1S1, AQP5, ARSE, CAST, CDSN, CERS3, CLDN1, CSTA, CTSC, CYP4F22, DSC2, DSC3, DSG1, DSG4, DSP, EBP, ELOVL4, ENPP1, FLG, GJA1, GJB2, JUP, KANK2, KDSR, KRT1, KRT10, KRT2, KRT6C*, KRT9, LIPN, MBTPS2, MVK, LOR, NIPAL4, NSDHL, PEX7, PHYH, PKP1, PNPLA1, POMP, RHBDF2, RSPO1, SERPINB7, SLC27A4, SLURP1, SNAP29, SPINK5, ST14, STK11, STS, SULT2B1, TGM1, TRPV3, VPS33B	112	
Family member testing				as indicated above	28	
Proforma required?		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
 - Striate keratoderma with woolly hair
 - Keratoderma with deafness
 - Unusual/unique rare keratodermas occurring alone or as part of syndromes
 - o Erythrokeratoderma

- Clinical Genetics
- Dermatology





RARE GENETICS INFLAMMATORY SKIN DISORDERS

Available testing

Centre	Method	Scope and range of test			Targets	TAT	
Dundee	NGS(clinical exome)	Wh ge scre	ne	SNVs, indels	, ADA2, , , AIRE, , , , , CARD11, CARD14, CARD9, , , , , , , , , DOCK8, , EGFR, , , , , , , , , , GJA1, GJB3, , IL1RN, GJB4, IL36RN, KIT, , , , , NCSTN, , NLRP3, NOD2, NSDHL, OSMR, , PSENEN, RAG1, RAG2, SAMHD1, SH3PXD2B, SLC39A4, STAT3, TMEM173, TREX1	112	
Family member testing					as indicated above	28	
Proforma required?		YES	YES Skin disorders proforma (see centre website)				

Referral criteria

- Any individual with a clinical diagnosis of a rare inflammatory skin disorder of a likely germline genetic cause
 - Includes autoinflammatory disease (e.g. early onset urticaria, recurrent febrile erythemas), infantile pustular psoriasis, likely genetic forms of pityriasis rubra pilaris
- Primary lymphoedema FLT4 analysis. This informs treatment

- Clinical Genetics
- Dermatology
- Rheumatology





SUPERFICIAL EPIDERMOLYTIC ICHTHYOSIS (SEI)

(previously known as ICHTHYOSIS BULLOSA of SIEMENS)

Available testing

Centre	Method	Scope and rang	je of test	Targets	TAT
Dundee	Sanger	Whole gene screen	SNVs, indels	KRT2	56
Family member testing			as indi	cated above	28
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		e gene een	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				28	
Proforma required?		YES	YES Skin disorders proforma (see centre website)				

Referral criteria

- Any individual with a vascular skin disorder with a likely germline genetic cause
- Note this method is not optimised to detect mosaic variants

- Clinical Genetics
- Dermatology





X-LINKED ICHTHYOSIS

Available testing

Centre	Method	,	Scope and I	range of test	Targets	TAT	
Dundee	Sanger MLPA	Whole gene screen		SNVs, indels Exon level CNV	STS	56	
Family member testing			as indicated above			28	
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









END