


Schedule of Accreditation

issued by

United Kingdom Accreditation Service

2 Pine Trees, Chertsey Lane, Staines-upon-Thames, TW18 3HR, UK

 8176 Accredited to ISO 15189:2012	Birmingham Women's and Children's NHS Foundation Trust	
	Issue No: 012 Issue date: 06 August 2024	
	Birmingham Women's Hospital Mindelsohn Way Edgbaston Birmingham B15 2TG	Contact: Jennie Bell Tel: +44 (0)1214 721377 E-Mail: jennie.bell1@nhs.net Website: www.bwc.nhs.uk

Testing performed by the Organisation at the locations specified below

Locations covered by the organisation and their relevant activities

Laboratory locations:

Location details	Activity
Birmingham Women's Hospital West Midlands Regional Genetics Laboratory Mindelsohn Way Edgbaston Birmingham B15 2TG Local contact Jennie Bell	Molecular Genetics

Site activities performed away from the locations listed above:

Location details	Activity
Birmingham Research Park Limited Institute of Research and Development Birmingham Research Park Vincent Drive Edgbaston Birmingham B15 2SQ Local contact Jennie Bell	Reporting



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DETAIL OF ACCREDITATION

Materials/Products tested	Type of test/Properties measured/Range of measurement	Standard specifications/ Equipment/Techniques used
HUMAN TISSUES AND BODY FLUIDS	Genomic analysis for the purpose of clinical diagnosis of rare disease and cancer	Documented in house procedures incorporating manufacturer's instructions (where relevant)
Whole Blood Bone Marrow	Sample processing, DNA/RNA extraction, quantification and quality check for subsequent in-house analysis (see below), referral to specialist centres and long term storage	Lineage specific cell separation Automated using Automacs Miltenyi Biotech autoMACS®Pro Separator SOP: PP 02.01.95 Manual and automated DNA extraction and quantification using: Qiagen QIASymphony SP platform with: QIASymphony DNA Midi Kit QIASymphony DSP virus kit SOPs: PP 03.01.21 PP 03.01.13 PP 03.01.48
Whole Blood Bone Marrow Saliva (Oragene self collection kit) Plasma		
Whole Blood Bone Marrow		Qiagen QIAcube with Qiagen QIAamp DNA Blood Mini Kit SOP: PP 03.01.15



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HUMAN TISSUES AND BODY FLUIDS	Genomic analysis for the purpose of clinical diagnosis of rare disease and cancer	Documented in house procedures incorporating manufacturer's instructions (where relevant)
FFPE Dried Blood Spots		Promega Maxwell 16 with Promega Maxwell 16 FFPE Plus LEV DNA Purification Kit and LEV Blood Kit PP 03.01.43 PP 03.01.46
MACS separated cell fractions Amniotic Fluid CVS		Qiagen EZ1 using EZ1Tissue Kit PP 03.01.08: PP 03.01.14: PP 03.01.10:
Fresh frozen solid tissue*		Qiagen EZ1 using EZ1Tissue Kit with manual homogenation PP 03.01.07
Fresh tissue Foetal Tissue POC		Qiagen EZ1 using EZ1Tissue Kit with tissue aubruption using Precellys 24 Homogeniser PP 03.01.19 Manual DNA Extraction:
Peripheral Blood Bone Marrow Foetal Blood Mouth Wash Cultured Cells		Qiagen Gentra Puregene Kit PP 03.01.02 PP 03.01.06 PP 03.01.05
Peripheral Blood		Biorad Instagene matrix PP 03.01.12
Peripheral Blood Bone Marrow, Foetal Blood CVS,Fixed Cells		Phenol Chloroform extraction PP 03.01.03, PP 03.01.04 PP 03.01.11



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<p>HUMAN TISSUES AND BODY FLUIDS</p> <p>Peripheral Blood Bone Marrow Foetal Blood</p> <p>Saliva (Oragene self collection kit)</p> <p>Genomic DNA extracted inhouse from the sample types listed above or received as primary sample type from an external source</p> <p>FFPE</p>	<p>Genomic analysis for the purpose of clinical diagnosis of rare disease and cancer</p> <p>Sample processing, DNA/RNA extraction, quantification and quality check for subsequent in-house analysis (see below), referral to specialist centres and long term storage</p>	<p>Documented in house procedures incorporating manufacturer's instructions (where relevant)</p> <p>Phenol Chloroform extraction – High Risk protocol</p> <p>PP 03.01.24</p> <p>DNA extraction using Genotek Prep-IT kit</p> <p>PP.03.01.13</p> <p>DNA Quantification for QC purposes Nanodrop 8000 Qubit 2.0/4.0 fluorometerPicoGreen quantification process for DNA samples using Fluroskan</p> <p>SOPs: PP 03.01.27 PP 03.01.36 PP 03.01.50</p> <p>Automated dual DNA and RNA extraction and quantification using:</p> <p>Dual DNA and RNA extraction using Promega Maxwell 16 LEV RNA FFPE Purification Kit and Promega Maxwell IVD</p> <p>MP 01.01.55</p>



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HUMAN TISSUES AND BODY FLUIDS	Genomic analysis for the purpose of clinical diagnosis of rare disease and cancer	Documented in house procedures incorporating manufacturer's instructions (where relevant)
Whole Blood Bone Marrow	Sample processing, DNA/RNA extraction, quantification and quality check for subsequent in-house analysis (see below), referral to specialist centres and long term storage	Automated RNA extraction using: Extraction of total RNA using the Promega Maxwell IVD and Promega Maxwell 16 LEV simply RNA Blood Kit PP 03.01.47
RNA extracted inhouse from the sample types listed above or received as primary sample type from an external source		Reverse Transcription Manual -High Capacity cDNA Reverse Transcriptase kit (Applied Biosystems) PP 03.01.26:
RNA extracted inhouse from FFPE		RNA Quantification for QC purposes Qubit PP 03.01.36
Genomic DNA extracted in-house from the sample types listed and received as primary samples from external sources	Detection of SNVs and Small indels [definitive list QA 01.02.60]	Fluorescence based ARMS PCR Multiplexing Luminex commercial CF XTAG kit: Equipment: Luminex 200 Analysis using: Luminex integrated software SOP: FRAG 01.01.05



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<p>HUMAN TISSUES AND BODY FLUIDS</p> <p>Genomic DNA and RNA extracted in-house from the sample types listed and received as primary samples from external sources</p>	<p>Genomic analysis for the purpose of clinical diagnosis of rare disease and cancer</p> <p>Detection of nuclei acid sequence variants – SNVs, small indels and splice site mutations</p> <p>[definitive list QA 01.02.60]</p>	<p>Documented in house procedures incorporating manufacturer's instructions (where relevant)</p> <p>Sanger Sequencing</p> <p>Using</p> <p>Standard primer design methodology and PCR amplification (where relevant to internal samples and confirmatory processes)</p> <p>SOP: SEQ 01.01.26 PP 03.01.35</p> <p>And:</p> <p>PCR blocks, ABI 3730 Capillary electrophoresis instruments</p> <p>Analysis and interpretation of variants by Mutation Surveyor software.</p> <p>SOP: 01.01.10 SEQ 01.01.25</p> <p>Fragment Length Analysis</p> <p>Fragment Length Analysis using Capillary Electrophoresis</p> <p>PCR, methylation specific PCR, Triplet Repeat PCR using in-house methods</p> <p>Equipment: PCR blocks and resolution using an ABI 3500XL capillary sequencer:</p> <p>Analysis using GeneMapper and GeneMarker</p> <p>SOPs: FRAG 01.01.07 FRAG 01.01.13</p>
<p>Genomic DNA extracted in-house from the sample types listed and received as primary samples from external sources</p>	<p>Detection of fragment length size, deletions, known SNVs and small indels, gene rearrangements, repeat expansions, linkage makers, short tandem repeats, microsatellites and methylation status</p> <p>[definitive list QA 01.02.60]</p>	



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HUMAN TISSUES AND BODY FLUIDS	Genomic analysis for the purpose of clinical diagnosis of rare disease and cancer	Documented in house procedures incorporating manufacturer's instructions (where relevant)
Genomic DNA extracted in-house from the sample types listed	Determination of copy number changes [definitive list QA 01.02.60]	Quantitative Fluorescence PCR (QF-PCR) Using: In-house methods for trisomy screen And: Thermocyclers. Resolution by capillary electrophoresis using ABI 3500XL. Analysis using Genemarker and Genemapper SOP: FRAG 01.01.16 PN 01.01.18
Genomic DNA and RNA extracted in house from the sample types listed above and received as primary samples from external sources (cont'd)	Qualitative Genotype analysis for SNVs, indels and fusion transcripts [definitive list QA 01.02.60]	Qualitative Reverse Transcriptase PCR (RTPCR) (Including nested) Using: Agarose gel electrophoresis SOPs:TP 01.01.67 TP 01.01.61 TP 01.01.60
RNA extracted in house from the sample types listed above and received as primary samples from external sources (cont'd)	For the quantitative detection of common fusion transcripts [definitive list QA 01.02.60]	Quantitative Real Time PCR (RQ-PCR) Using: In house methodology ABI 7500 Real time PCR system SOPs: QPCR 01.01.11 HOA 01.01.57 QPCR 01.01.27 HOA 01.01.61



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HUMAN TISSUES AND BODY FLUIDS	Genomic analysis for the purpose of clinical diagnosis of rare disease and cancer	Documented in house procedures incorporating manufacturer's instructions (where relevant)
Genomic DNA extracted in-house from the sample types listed and received as primary samples from external sources	Determination of methylation status and copy number – deletions and duplications [definitive list QA 01.02.60]	Multiplex Ligation-dependent Probe Amplification (MLPA) and methylation specific (MS) MLPA. Using Commercial MRC Hollandkits, thermocyclers and ABI3500xl Analysis using GeneMarker and Genemapper SOPs:FRAG 01.01.04 FRAG 01.01.10
Genomic DNA extracted in-house from the sample types listed above	Post-transplant Chimaerism analysis - % donor engraftment	Microsatellite analysis using in house PCR methodology and ABI 3500 XL. Analysis using; ChimerMarker Software SOPs: TP 01.01.63, HOA 01.01.14
Genomic DNA extracted in-house from the sample types listed above	Detection of known SNVs and indels at high sensitivity [definitive list QA 01.02.60]	Droplet Digital PCR Using: BioRAD commercial Kits and in-house methodology BioRAD QX200 Droplet Reader BioRAD AutoDroplet Generator BioRAD PCR Plate Reader Analysis using integrated software SOPs: QPCR 01.01.01, QPCR 01.01.31



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HUMAN TISSUES AND BODY FLUIDS	Genomic analysis for the purpose of clinical diagnosis of rare disease and cancer	Documented in house procedures incorporating manufacturer's instructions (where relevant)
Cell-Free DNA extracted in-house from plasma	CNV/(Rapid screening for aneuploidy (T13,18,21)	Next Generation Sequencing VeriSeq Solution v 2.0 For Non-invasive Prenatal Testing (NIPT) using With Automated cfDNA extraction and Library Preparation using Hamilton Star Liquid Handler and with massively paralleled sequencing on a NextSeq 550 with VeriSeq NIPT LRM module for analysis. Receipting, Processing and Reporting of Lucina NIPT referrals PN 01.01.43 And SEQ 01.01.69 NIPT Veriseq Technical protocol
Genomic DNA and RNA extracted in house from the sample types listed and samples received as primary samples from external sources (cont'd)	Gene screening of large gene panels for genetic variants – [definitive list QA 01.02.60] SNVs/indels SNVs/indels	Next Generation Sequencing: Sequencing by MiSeq KAPA Hyper plus/ NimbleGen (Roche) SOP:PN 01.02.10 PN 01.02.04 Multiplex PCR (QIAGEN) SOP:PN 01.02.02 PN 01.02.03



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<p>HUMAN TISSUES AND BODY FLUIDS</p> <p>Genomic DNA and RNA extracted in house from the sample types listed and samples received as primary samples from external sources (cont'd)</p>	<p>Genomic analysis for the purpose of clinical diagnosis of rare disease and cancer</p>	<p>Documented in house procedures incorporating manufacturer's instructions (where relevant)</p> <p>Next Generation Sequencing (cont):</p> <p><u>Analysis using:</u></p> <p>Agilent 2200 TapeStation, Qubit 2.0 Fluorometer, thermal cycler and 2.0 Fluorometer, Illumina MiSEQ</p> <p>Analysis using:</p> <p>BI 01.01.08: NIPD RHDO Bioinformatics Pipeline BI 01.01.07: NIPD Bespoke Bioinformatics Pipeline</p> <p>SEQ 01.01.22</p>
<p>DNA obtained from whole blood</p>	<p>DYPD testing</p>	<p>PCR with Maldi-TOF Agena MassArray PG 01.01.03</p>



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HUMAN TISSUES AND BODY FLUIDS Whole Blood Amniotic Fluid CVS Foetal blood Products of Conception Bone Marrow Tissues/Skin Biopsy	Genomic analysis for the purpose of clinical diagnosis of rare disease and cancer G-banding/Karyotyping: Detection of chromosomal rearrangements or aberrations arising from: (e.g) Prenatal Diagnosis Reproductive Medicine Disorders Developmental Disorders Haemto/Oncology Disorders Chromosome Breakage Disorders (preparative pre-examination steps listed first)	Documented in house procedures incorporating manufacturer's instructions (where relevant) Culturing and processing of human tissue/cells using in-house and commercial media to provide interphase cells: Cell Separation using autoMACS (see processing section) Cell Harvesting Automated process using Multiprep Cell Sprint Robotic Harvester SOP TP 01.01.15 Manual Process SOP: PP 02.01.40 Chromosome analysis, Microscopic and Macroscopic analysis of G banded chromosomes using Manual Processing and Varistain Banding Instrument and microscopes Analysis using Metasystems Icaris SOPs: PP 02.01.92 PP 02.01.93 PP 02.01.94 KARYO 01.01.06 GL 01.01.19 HOA 01.01.05



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HUMAN TISSUES AND BODY FLUIDS Formalin fixed paraffin embedded tissue (FFPE) Peripheral Blood Bone Marrow Fixed culture cells (more specific – cultured, uncultured PB, Marrow, AFs etc) Amniotic fluid CVS- Chronic villus samples	Genomic analysis for the purpose of clinical diagnosis of rare disease and cancer Detection of chromosomal aberrations in the diagnosis of haematological malignancy, bone marrow failure syndromes, non-haematological malignancies and constitutional disorders, solid tumours and companion testing Break-apart probes Fusion products Deletion Insertion Copy Number / Amplification	Documented in house procedures incorporating manufacturer's instructions (where relevant) Fluorescence in situ hybridisation (FISH) Culturing and processing of human tissue/cells using in-house and commercial media to provide interphase/metaphase cells: Preparation and harvesting as for G-banding Manual Process PP 02.01.40 And commercial and in house developed probes. Hybridisation using Hybrite/Thermobrite Analysis Fluorescence microscope and metasystems ISIS SOPs FISH 01.01.04, FISH 01.01.95, FISH 01.01.97, GI 01.01.06
END		