


Schedule of Accreditation

issued by

United Kingdom Accreditation Service

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

 <p>8413</p> <p>Accredited to ISO 15189:2012</p>	<h3>NHS Lothian</h3> <p>Issue No: 008 Issue date: 26 February 2024</p>	
	<p>Genetics Laboratory South East Scotland Genetic Service David Brock Building Western General Hospital Edinburgh EH4 2XU</p>	<p>Contact: Bernard Lawless Email:bernard.lawless@nhs.scot Website: http://tinyurl.com/Edinburghgenelab</p>
<p>Testing performed at the above address only</p>		

DETAIL OF ACCREDITATION

Materials/Products tested	Type of test/Properties measured/Range of measurement	Standard specifications/ Equipment/Techniques used
<p>HUMAN TISSUES & FLUIDS</p> <p>Amniotic Fluid.</p> <p>Peripheral Blood, Bone Marrow, Mouth wash (Oragene Collection) and Buccal scrapes.</p> <p>CVS, fresh tissue, frozen or fixed human tissue or cells, neonatal and Maternal peripheral bloods.</p> <p>Genomic DNA extracted in-house from the sample types listed above or received as primary sample type from an external source.</p>	<p><u>Genomics analysis for the purpose of clinical diagnosis of rare disease, inherited and acquired cancer</u></p> <p>DNA extraction, quantification and quality check for subsequent in-house analysis, referral to other specialist centres and long term storage.</p>	<p>Documented in house procedures incorporating manufacturer's instructions (where relevant)</p> <p>Manual, semi automated and Automated DNA extraction and quantification using:</p> <p>DNA extraction:</p> <p>Manual extraction processes:</p> <p>Igenatal Genomic DNA extraction kit SOP GENE-WC87.</p> <p>Semi-Automated and Automated extraction processes:</p> <p>Chemagic 360 DNA extractor and Janus Robotic Workstation SOP GENE-WM321 and SOPGENE-WM43.</p> <p>Qiagen EZ1 Advanced XL with DNA tissue kit and DNA blood kit SOP GENE-WC87.</p> <p>DNA Quantification for QC purposes:</p> <p>Using: NanoDrop One and Promega Quantus Fluorometer SOP's GENE-WM363 and GENE-WM17.</p>



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<p>Genomic DNA extracted in-house from the sample types listed above or received as primary sample type from an external source. (cont'd)</p>	<p>Detection of genetic variants of SNVs and small indels. [definitive list to be held by this laboratory]</p>	<p>Sanger Sequencing Standard primer design methodology</p> <p>SOP's GENE-WM166 GENE-WM177</p> <p>Semi- nested PCR design methodology</p> <p>SOP GENE-WM157</p> <p>PCR amplification using in-house methodology using:</p> <p>For manual process</p> <p>PCR blocks, ABI3500xl Capillary electrophoresis instruments.</p> <p>For automated process</p> <p>Beckman Biomek NXp Liquid handler, PCR blocks.</p> <p>Sequencing of products by ABI3500xl Capillary electrophoresis instruments.</p> <p>Analysis using SoftGenetics Mutation Surveyor™ software and interpretation of variants by Alamut software.</p> <p>Procedures:</p> <p>SOP's GENE-WM42, GENE-WM136 GENE-WM16, GENE-WM28, GENE-WM19, GENE-WM12, GENE-WM8, GENE-WM57</p>



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<p>Genomic DNA extracted in-house from the sample types listed above or received as primary sample type from an external source. (cont'd)</p>	<p>Targeted screening for the detection of SNVs and small insertions/deletions using custom-designed gene panels. [definitive list to be held by this laboratory]</p>	<p>Next Generation Sequencing</p> <p>Library amplification and hybridisation performed using Illumina DNA prep kit.</p> <p>Enrichment with Twist Bioscience probes.</p> <p>Library amplification and hybridisation performed using Illumina DNA prep kit. Enrichment with Twist Bioscience or Illumina TruSight Capture probes (set up performed using the Hamilton Microlab STAR Liquid Handling System) or manually</p> <p>Paired-end next-generation sequencing performed on Illumina MiSeq instruments.</p> <p>Using</p> <p>Applied Biosystems thermal cyclers, Promega Quantus fluorometer and Illumina MiSeq.</p> <p>Data analysis using SoftGenetics, NextGENe and Illumina Local Run Manager/ Variant Studio .</p> <p>SOP's GENE-WM342, GENE-WM167, GENE-WM2, GENE-WM158, GENE-WM57, GENE-WM16, GENE-WM383, GENE-WM162</p>



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<p>Genomic DNA extracted in-house from the sample types listed above or received as primary sample type from an external source. (cont'd)</p>	<p>Detection of fragment length size, deletions, known mutations, repeat expansions, short tandem repeats. [definitive list to be held by this laboratory]</p> <p>Determination of repeat size expansions. Detection of fragment length size, deletions, known mutations, repeat expansions, short tandem repeats. [definitive list to be held by this laboratory]</p> <p>Detection of Cystic fibrosis (CFTR) variants.</p> <p>Rapid detection of common trisomies.</p>	<p>Fragment Length Analysis</p> <p>Determination of repeat size using the Asuragen AmplideX kit with: Manual set up using Thermal cyclers, and ABI3500xl Capillary electrophoresis instrument. Analysis using SoftGenetics GeneMarker™ SOP's GENE-WM27, GENE-WM6, GENE-WM117, GENE-WM116.</p> <p>Detection of Cystic fibrosis (CFTR) variants using the YourGene Cystic Fibrosis Base kit</p> <p>Using thermal cyler and 3500XL Genetic Analysers. GENE-WM42 GENE-WM136</p> <p>Analysis using SoftGenetics GeneMarker™ software.</p> <p>SOP's GENE-WM24, GENE-WM5</p> <p>Quantitative Fluorescence Polymerase Chain Reaction (QF-PCR)</p> <p>PCR amplification using YourGene QST*R Base kit, YourGene QST*R plus XY kit, YourGene QST*R plus 13, 18, 21 kits, and YourGene Male Infertility Base kit and thermal cyclers with ABI 3500XL Genetic Analyser.</p> <p>Analysis and interpretation of aneuploidy results using GeneMarker software.</p>



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<p>Genomic DNA extracted in-house from the sample types listed above or received as primary sample type from an external source. (cont'd)</p>	<p>Determination of copy number changes (deletions and duplications).</p> <p>Detection of fragment length size, deletions, known mutations, repeat expansions, short tandem repeats. [definitive list to be held by this laboratory]</p> <p>Determination of repeat size expansions.</p>	<p>SOP's GENE-WC246, GENE-WC85, GENE-WC86, GENE-WC245.</p> <p>Multiplex Ligation-dependent Probe Amplification (MLPA)</p> <p>Using</p> <p>In-house methods or MRC Holland kits, thermocyclers and ABI3500xl Genetic Analyser.</p> <p>Analysis using GeneMarker SOP's GENE-WM45, GENE-WM7.</p> <p>In-house designed</p> <p>Flanking or repeat Primer PCR using in-house methods:</p> <p>Automated set up using Thermal cyclers, BiomekNXp liquid handling robot and ABI3500xl Capillary electrophoresis instrument.</p> <p>Analysis using SoftGenetics GeneMarker™</p> <p>SOP's GENE-WM152, GENE-WM187, GENE-WM156, GENE-WM179, GENE-WM29, GENE-WM118, GENE-WM153, GENE-WM154, GENE-WM151, GENE-WM194, GENE-WM294.</p> <p>Standard in-house primer design methodology SOP GENE-WM204.</p>



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<p>HUMAN TISSUES & FLUIDS (cont'd)</p> <p>Genomic DNA extracted in-house from the sample types listed above or received as primary sample type from an external source. (cont'd)</p>	<p><u>Genomics analysis for the purpose of clinical diagnosis of rare disease, inherited and acquired cancer</u> (cont'd)</p> <p>Manual multiplex short tandem repeat (STR) PCR.</p> <p>Detection of F8 intron 22 inversions by inverse PCR.</p> <p>Detection of F8 intron 1 inversions by long range PCR.</p>	<p>Documented in house procedures incorporating manufacturer's instructions (where relevant)</p> <p>Manual multiplex short tandem repeat (STR) PCR.</p> <p>Using thermal cyclers and ABI3500XL genetic analyser.</p> <p>Analysis of marker multiplexes and haplotyping by SoftGenetics GeneMarker™.</p> <p>SOP's GENE-WM114 and GENE-WM70</p> <p>In-house designed primers, thermocyclers and ABI3500xl Genetic Analyser. Analysis using GeneMarker</p> <p>GENE- WM377. GENE-WM378,</p> <p>In-house designed primers, thermocyclers and Gel electrophoresis using E-Gel Power Snap electrophoresis device.</p> <p>GENE- WM377, GENE-WM379</p>



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<p>HUMAN TISSUES & FLUIDS (cont'd)</p> <p>DNA extracted from Blastocyst or trophoctoderm cells.</p>	<p><u>Genomics analysis for the purpose of clinical diagnosis of rare disease, inherited and acquired cancer</u> (cont'd)</p> <p>Haplotyping using multiplex short tandem repeat (STR) loci for allele size determination. [definitive list to be held by this laboratory]</p>	<p>Documented in house procedures incorporating manufacturer's instructions (where relevant)</p> <p>Preimplantation Genetic Testing</p> <p>Standard in-house primer design methodology SOP's GENE-WM204.</p> <p>Whole genome amplification (WGA) of DNA from biospies using Qiagen REPLI-g kit and thermal cyclers.</p> <p>Manual multiplex short tandem repeat (STR) PCR using PCR using thermal cyclers and ABI3500XL genetic analyser.</p> <p>Analysis of loci and haplotyping by SoftGenetics GeneMarker™.</p> <p>SOP's GENE-WM72, GENE-WM351, GENE-WM114, GENE-WM70.</p>



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Whole blood Amniotic fluid CVS Bone marrow Lymph node Fresh tissue samples	<p>G-banding/Karyotyping:</p> <p>Detection of chromosomal rearrangements or aberrations arising from: (e.g) Prenatally detected disorders Developmental disorders Reproductive medicine disorders Haematological/Oncology disorders. [definitive list to be held by this laboratory]</p> <p>(Preparative Pre-examination steps listed first)</p>	<p>Manual culturing and processing of human tissue to provide metaphase cells: using commercial media and:</p> <p>GENE-WC63: GENE-WC56: GENE-WC62: GENE-WC49: SOPs.</p> <p>Preparation of Myeloma cells using CD138+ve selection GENE-WC132:</p> <p>Cell Culture protocols</p> <p>GENE-WC1, GENE-WC95, GENE-WC96, GENE-WC57, GENE-WC58, GENE-WC64</p> <p>Cell Harvest Manual in-house Protocols GENE-WC2, GENE-WC99, GENE-WC66</p> <p>Slide Preparation:</p> <p>Slide preparation and banding techniques using Thermotron Humidity Chamber and Gemini Automated Stainer SOP's GENE-WC108, GENE-WC37 and GENE-WC38</p> <p>Analysis: G:banding analysis using the Bioview imaging system</p> <p>SOP's GENE-WC23, GENE-WC105, GENE-WC311, GENE-WC312 GENE-WC315, GENE-WC313</p>



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<p>HUMAN TISSUES & FLUIDS (cont'd)</p> <p>Genomic DNA extracted in-house from the sample types listed above or received as primary sample type from an external source.</p>	<p><u>Genomics analysis for the purpose of clinical diagnosis of rare disease, inherited and acquired cancer</u> (cont'd)</p> <p>Cytogenetic examinations for diagnosing postnatal disorders, prenatal diagnosis, neoplastic genetics including haemato-oncology and solid tumours, and loss of pregnancy.</p> <p>By detection of microscopic chromosomal imbalance (gains and losses) expressed as changes to copy number. [definitive list to be held by this laboratory]</p>	<p>Documented in house procedures incorporating manufacturer's instructions (where relevant)</p> <p>Microarray</p> <p>Processing using the Affymetrix Cytoscan 750K and HD array chips on a GeneChips Fluidics workstation. Analysis of array data using ChAS software.</p> <p>Processing using Cytoscan 750K and HD array chips on a GeneChips Fluidics workstation SOP's GENE- WC70 to WC104.</p> <p>Analysis of array data using ChAS software for postnatal, prenatal and oncology samples. GENE-WC107, GENE-WC123, GENE-WC243.</p>



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<p>HUMAN TISSUES & FLUIDS (cont'd)</p> <p>Cultured & uncultured cells Paraffin embedded tissues (PETs) Tumour imprints Purified plasma cells</p>	<p><u>Genomics analysis for the purpose of clinical diagnosis of rare disease, inherited and acquired cancer</u> (cont'd)</p> <p>Detection of chromosomal aberrations in the diagnosis of haematological malignancy, bone marrow failure syndromes, non-haematological malignancies and constitutional disorders and solid tumours.</p> <p>Break-apart probes Fusion products Deletion Insertion Copy Number / Amplification</p> <p>[definitive list to be held by this laboratory]</p>	<p>Documented in house procedures incorporating manufacturer's instructions (where relevant)</p> <p>Fluorescent in-situ hybridisation (FISH)</p> <p>Preparation of PETS slides by Fluorescent in situ hybridisation for: (FISH) using VIP2000 processor GENE-WC15:</p> <p>Documented in house methods using commercial probes.</p> <p>FISH protocols for processing slides SOP's GENE-WC6, GENE-WC13, GENE-WC14, GENE-WC130</p>
END		