

# Schedule of Accreditation

issued by

## United Kingdom Accreditation Service

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



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ISO 15189:2012

### Great Ormond Street Hospital for Children NHS Foundation Trust

Issue No: 006 Issue date: 27 May 2021

North Thames GLH  
Barclay House (Levels 4, 5 & 6)  
37 Queen Square  
London  
WC1N 3BH  
United Kingdom

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Testing performed by the Organisation at the locations specified below

#### Locations covered by the organisation and their relevant activities

##### Laboratory locations:

| Location details  | Activity                              | Location code   |     |
|---|---------------------------------------|---|-----|
| <b>Address</b><br>Regional Genetics Laboratory<br>Barclay House (Levels 5 & 6)<br>37 Queen Square<br>London<br>WC1N 3BH   | <b>Local contact</b><br>Lucy Jenkins  | Rare and Inherited diseases<br>Molecular genetics<br>Cytogenetics                 | RGL |
| <b>Address</b><br>SIHMDS – Acquired Genomics<br>Level 2<br>Camelia Botnar Laboratories<br>Great Ormond Street Hospital for Children<br>NHS Foundation Trust<br>London<br>WC1N 3JH | <b>Local contact</b><br>Dariusz Ladon | Acquired genomics as part of SIHMDS service<br>Molecular Genetics<br>Cytogenetics | CBL |



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#### DETAIL OF ACCREDITATION

| Materials/Products tested  | Type of test/Properties measured/Range of measurement                                     | Standard specifications/ Equipment/Techniques used  | Location Code |
|--|---|---|---------------|
| HUMAN TISSUE AND FLUIDS  | <u>Molecular Genetics examination procedures for the purposes of clinical diagnosis</u>   | Documented methods for DNA and RNA extraction using one or a combination of the techniques below by in-house procedures, using commercial kits and manual extraction  |               |
| Blood / bone marrow / CSF/ fresh tissue / CSF / frozen tissue / body fluid | Extraction  | Using Maxwell 16 and Nanodrop one (spectrophotometer)<br>SOP: HSOP 782<br>And RGS MOT9035   | CBL           |
| FFPE section rolls   | Extraction  | Using Maxwell 16<br>Nanodrop one (spectrophotometer)<br>SOP: HSOP 782<br>And RGS MOT9038  | CBL           |
| Blood / bone marrow / fresh tissue / frozen tissue                         | Extraction  | Using Maxwell 16 and Nanodrop one (spectrophotometer)<br>SOP: HSOP 782<br>And RGS MOT9036   | CBL           |
| FFPE section rolls   | Extraction  | Using Maxwell 16 and Nanodrop one (spectrophotometer)<br>SOP: HSOP 782<br>And RGS MOT9039   | CBL           |
|  | DNA and RNA profiling for detection of abnormal sequences associated with acquired cancer | In-house documented methods and following manufacturers' instructions   |               |
| DNA and RNA. Extracted from primary samples as detailed above              |   | Allele specific Polymerase chain reaction (AS-PCR) and reverse transcription using 7500 real-time PCR system<br>Process SOP: RGS MOT9048<br>Analysis SOP: RGS ANA9049 | CBL           |
| DNA. Extracted from primary samples as detailed above                      |   | Fragment analysis using 3500 Genetic analyser<br>Tetrad thermal cycler<br>Process SOP FSOP 045 (RGS MOT 9045)<br>Anaysis SOP: FSOP 046 (RGS ANA9046)                  | CBL           |



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| HUMAN TISSUE AND FLUIDS (cont'd)                      | <u>Molecular Genetics examination procedures for the purposes of clinical diagnosis (cont'd)</u>   | In-house documented methods and following manufacturers' instructions  |               |
| DNA. Extracted from primary samples as detailed above | DNA and RNA profiling for detection of abnormal sequences associated with acquired cancer (cont'd) | High resolution Melt curve analysis using 7500 real-time PCR system (ARMs based)<br>Process SOP: FSOP 032 (RGS MOT9032)<br>Analysis SOP: RGS ANA9034   | CBL           |
| PCR product   |  | Sanger sequence analysis using Tetrad thermal cycler<br>Exasapit clean up of PCR products<br>SOP: RGSMOT9009<br>Cycle sequencing<br>SOP: RGS MOT9005<br>Clean up of big dye products<br>SOP: RGS MOT9040<br>Using Mutation Surveyor Software to analyse DNA variants<br>Analysis SOP: RGS ANA9037<br>Sequence Analysis and Mutation Classification<br>Analysis SOP: FSOP 025 (RGS ANA9025) | CBL           |
| RNA   |  | Reverse transcription-PCR using Thermal cycler<br>Reverse transcription<br>SOP: RGS MOT9013<br>Reporting Agarose Gel Electrophoresis<br>Analysis SOP: RGS ANA9060<br>Electrophoresis gel tank and power pack<br>Agarose Gel Analysis of DNA and RNA<br>SOP: HSOP 646 (RGS MOT9646)   | CBL           |



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| HUMAN TISSUE AND FLUIDS (cont'd) | <u>Molecular Genetics examination procedures for the purposes of clinical diagnosis (cont'd)</u>   | In-house documented methods and following manufacturers' instructions   |               |
| DNA                              | DNA and RNA profiling for detection of abnormal sequences associated with acquired cancer (cont'd) | Methylation arrays using Bisulfite conversion and FFPE restore SOP: RGS MOT 9053<br>Methylation array analysis using bCHAPs pipeline, Analysis SOP:RGS ANA9055  | CBL           |
| DNA                              |  | Multiplex Ligation Probe Amplification (MLPA) using Tetrad thermal cycler SOP:RGS MOT9015<br>MLPA Analysis using GeneMarker Software Analysis SOP:RGS ANA9027   | CBL           |
| DNA                              | DNA profiling for detection of abnormal sequences associated with colo-rectal cancer               | Multi-ligation Probe Analysis (MLPA). Performed by manual in-house method or using the Biomek NK span-8 system.<br>Microsatellite instability (MSI) analysis: Manual in-house method<br>RGS MOT0401 Setting up MLPA using the Automated system<br>RGS MOT0406 Manual MLPA set up (molecular)<br>RGS MOT0409 MS-MLPA set-up<br>RGS LAB0102 ABI3730 XL General Use<br>RGS ANA0201 Capillary Electrophoresis of MLPA and QF-PCR products using the ABI 3730<br>RGS ANA0205 Analysis of MLPA using GeneMarker Software (Molecular)<br>RGP DCA0003 Colorectal cancer disease policy<br>RGS ANA0210 MLH1 methylation MS-MLPA Analysis<br>RGS MOT0410 Promega MSI kit - Set up and analysis<br>RGS EXT0371 Promega MSI kit | RGL           |



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| HUMAN TISSUE AND FLUIDS (cont'd)                       | <u>Molecular Genetics examination procedures for the purposes of clinical diagnosis (cont'd)</u>                               | Documented methods for DNA and RNA extraction using commercial kits and extraction  |               |
| Whole Blood<br>Amniotic Fluid<br>CVS<br>Tissue Samples | Extraction of DNA  | RGS MOT0101 – DNA extraction from cultured cells<br>RGS MOT0103 - DNA salting out method for CVS<br>RGS MOT0104 - DNA salting out method for tissue<br>RGS MOT0106 – Extraction of DNA from cells/blood spots by EZ1<br>RGS MOT0110 – DNA extraction from tissue using FUJIQuickGene-Mini80<br>RGS MOT0111 – DNA extraction from blood or saliva using FUJIQuickGene-610L<br>RGS MOT0112 - DNA extraction using the iGENatal Extraction kit<br>RGS MOT0128 – DNA extraction using the Chemagic Star | RGL           |
| Whole Blood  |  | DNA extraction from whole blood using Kurabo QuickGene Auto240L<br>RGS MOT0134 - DNA extraction using the QuickGene Auto240L  | RGL           |
| DNA. Extracted from primary samples as detailed above. | DNA and RNA profiling for detection of abnormal sequences for common and rare inherited/predisposing genetic disease condition | Sanger Sequencing techniques and targeted testing using ABI 3730XL, ABI 3130XL, Beckman NX Robotic Workstations, Thermal Cyclers in accordance with SOPs: -<br>RGS MOT0114 – Excel Instructions for streamlined robot PCR & Sequencing<br>RGS MOT0305 – Streamlined Robot PCR setup<br>RGS MOT0311 PCR purification, sequencing and clean up<br>RGS LAB0107 – DYAD/TETRAD PCR Machines<br>RGS LAB0102 - ABI3730 XL General Use<br>RGS LAB0723 - Equipment – Biomek NX mulitchannel robot            | RGL           |



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| HUMAN TISSUE AND FLUIDS (cont'd)                       | <u>Molecular Genetics examination procedures for the purposes of clinical diagnosis (cont'd)</u>   | In-house documented methods and following manufacturers' instructions   |               |
| DNA. Extracted from primary samples as detailed above. | DNA and RNA profiling for detection of abnormal sequences for common and rare inherited/predisposing genetic disease conditions (cont'd) | PCR using CFEU2 / CFEU4 kit<br>RGS LAB0106 – GeneAmp 9700PCR Machine SOP<br>RGS LAB0102 - ABI3730 XL General Use<br>RGS DGE0304 - Elucigene CF-EU2 Set Up and Analysis<br>RGS DGE0305 - Elucigene CF-HT4 - set up and analysis  | RGL           |
| DNA. Extracted from primary samples as detailed above. | DNA and RNA profiling for detection of repetitive sequences to exclude maternal cell contamination and identity testing                  | PCR using Power Plex16 HS assay   | RGL           |
| DNA extracted from primary samples as detailed above.  |  | Direct PCR and Amplidex FMR1 PCR Kits (P/N 76008 VH Bio) to amplify across the <i>FMR1</i> gene triplet repeat by Triplet Primed PCR. Detection is using fragment analysis on a capillary sequencer or ABI3730XL as documented in RGS DGE0501 Fragile X PCR<br>RGS DGE0506 Fragile X - Asuragen Amplidex FMR1 PCR<br>RGS LAB0102 - ABI3730 XL General Use | RGL           |
| DNA extracted from primary samples as detailed above   |  | Direct PCR and Triplet Primed PCR to amplify across the <i>DM1</i> gene triplet repeat. Detection is using fragment analysis on a capillary sequencer (ABI3730XL) as documented in RGS DGE1100 Myotonic dystrophy - TP-PCR testing  | RGL           |
| DNA. Extracted from primary samples as detailed above. |  | Restriction enzyme digest performed manually according to in-house method described in RGS MOT0307 – Manual PCR using KCl, NH4 or Megamix buffers<br>RGS MOT0310 – Agarose Gel preparation and Electrophoresis  | RGL           |



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| HUMAN TISSUE AND FLUIDS (cont'd)                  | <u>Molecular Genetics examination procedures for the purposes of clinical diagnosis (cont'd)</u> | In-house documented methods and following manufacturers' instructions  | RGL           |
| DNA extracted from Whole Blood Amniotic Fluid CVS |  | Prenatal Exome sequencing by Next Generation Sequencing supported by PCR amplification or hybridisation. Sequencing carried out using the Illumina NextSeq sequencer.<br>RGS MOT0118 Quantification of DNA using Qubit fluorometer<br>RGS MOT0129 Quantity and Quality assessment using Agilent 2200 TapeStation<br>RGS MOT0232 NextSeq for SureSelect<br>RGS MOT0132 Rapid low input DNA manual Agilent SureSelect XT library preparation<br>RGS ANA0512 GOSHG2P Analysis<br>RGS ANA0513 Genesis pipeline<br>RGP DEX002 Prenatal exomes                         |               |
| DNA extracted from Whole Blood Amniotic Fluid CVS |  | Custom Clinical Panel by Next Generation Sequencing supported by PCR amplification or hybridisation. Sequencing carried out using the Illumina NextSeq sequencer.<br>RGS MOT0118 Quantification of DNA using Qubit fluorometer<br>RGS MOT0129 Quantity and Quality assessment using Agilent 2200 TapeStation<br>RGS MOT0232 NextSeq for SureSelect<br>RGS MOT0226 SureSelect Library Preparation<br>RGS MOT0231 NGS BRAVO SURESELECT XT LIBRARY PREPARATION<br>RGS ANA0512 GOSHG2P Analysis<br>RGS ANA0513 Genesis pipeline<br>RGP DEX0003 Custom Clinical Panel |               |



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| HUMAN TISSUE AND FLUIDS (cont'd)                      | <u>Molecular Genetics examination procedures for the purposes of clinical diagnosis (cont'd)</u>  | In-house documented methods and following manufacturers' instructions   |               |
| DNA extracted from primary samples as detailed above. | DNA and RNA profiling for detection of abnormal sequences for common and rare inherited/predisposing genetic disease condition (cont'd) | Multi-ligation Probe Analysis (MLPA). Performed by manual in-house method or using the Biomek NK span-8 system.<br>RGS MOT0401 Setting up MLPA using the Automated system<br>RGS MOT0406 Manual MLPA set up (molecular)<br>RGS ANA0201 Capillary Electrophoresis of MLPA and QF-PCR products using the ABI 3730 Genetic Analyser<br>RGS ANA0205 Analysis of MLPA using GeneMarker Software (Molecular)  | RGL           |
| DNA extracted from primary samples as detailed above  |   | Next Generation Sequencing supported by PCR amplification or hybridisation. Sequencing is carried out using the Illumina MiSeq and NextSeq sequencers.<br>Quantification of DNA using Qubit fluorometer using RGS MOT0118<br>Quantity and Quality assessment using Agilent 2200 TapeStation RGS MOT0129<br>RGS MOT0216 MiSeq Sequencing for SureSelect Libraries<br>RGS MOT0232 NextSeq for SureSelect<br>RGS MOT0226 SureSelect Library Preparation<br>RGS MOT0231 NGS BRAVO SURESELECT XT Library preparation<br>RGS ANA0512 GOSHG2P Analysis<br>RGS ANA0513 Genesis pipeline | RGL           |





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| HUMAN TISSUE AND FLUIDS (cont'd)   | <u>Non-Invasive Prenatal examination procedures for the purposes of clinical diagnosis and antenatal screening</u> |   |               |
| Maternal Blood plasma              | Cell Free Fetal DNA (cffDNA) extraction from maternal plasma.  | RGS MOT0201 – ffDNA – Sample collection and separation of plasma DNA extraction based on Magnetic bead technology using the Qiagen Symphony SP DNA extractor<br>RGS MOT0224 for cffDNA extraction   | RGL           |
| DNA extracted from Maternal Plasma | Non-invasive prenatal diagnosis (NIPD)   | Real Time PCR analysis of cffDNA to determine fetal gender performed on the Applied Biosystems StepOne Plus Real-Time PCR Machine as documented in<br>RGS MOT0201 ffDNA Sample Collection and Separation of Plasma;<br>RGS DGE 0908 - ffDNA RQ-PCR Sexing<br>RGS LAB0133 Applied Biosystems StepOne Plus Real-Time PCR Machine (to replace RGS LAB0122 7300 Real Time PCR Machine   | RGL           |
| DNA extracted from Maternal Plasma | Non-invasive prenatal diagnosis (NIPD)   | Next Generation Sequencing performed on the MiSeq according to the in house method(s) to carry out non-invasive prenatal diagnosis for single gene disorders where the mutation is paternally derived, or <i>de novo</i> as documented in<br>RGS MOT0201 ffDNA Sample Collection and Separation of Plasma;<br>RGS MOT0215 - Using the QIAcube for MinElute® PCR Purification<br>RGS MOT0216 – MiSeq Sequencing for SureSlect Libraries<br>RGS DGE0911 Non-invasive prenatal diagnosis for de novo and paternal mutations (MiSeq)<br>RGS ANA0519 - NIPD and low level mosaicism analysis | RGL           |



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| HUMAN TISSUE AND FLUIDS (cont'd)<br><br>Blood | <u>Non-Invasive Prenatal examination procedures for the purposes of clinical diagnosis and antenatal screening (cont'd)</u><br><br>Relative Haplotype Dosage Analysis (RHDO) | Cell free foetal DNA (cffDNA) extracted from maternal blood samples for library preparation and Non-Invasive Prenatal Diagnosis (NIPD) using<br><br>QIASymphony SP for cffDNA extraction RGS MOT0224<br>DNA Extraction using the Chemagic Star RGS MOT0128<br>DNA Extraction From Blood or Saliva using FujiFilm/Kurabo QuickGene-610L RGS MOT0111<br>Quantification of DNA using Qubit fluorometer using RGS MOT0118<br>NextSeq for Sureselect RGS MOT0232<br>Quantity and Quality assessment using Agilent 2200 TapeStation RGS MOT0129<br>Covaris Operation RGS MOT0233<br>RGS ANA0514 Relative Haplotype Dosage Analysis<br>RGS MOT0131 Preparation of cffDNA samples for RHDO<br>RGS MOT0132 NIPD by Relative haplotype dosage analysis (RHDO) | RGL           |



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| <p>HUMAN TISSUE AND FLUIDS (cont'd)</p> <p>DNA extracted from:<br/>Blood, Saliva, Chorionic villus tissue, Uncultured amniocytes, Cultured cells from chorionic villus samples, Cultured cells from amniotic fluid, Saliva, Thymus</p> <p>Blood / Amniotic Fluid / Chorionic Villi / pregnancy loss tissue</p> | <p><u>Cytogenetics examination procedures for the purposes of clinical diagnosis</u></p> <p>Prenatal Diagnosis<br/>Reproductive Medicine Disorders<br/>Developmental Disorders</p> | <p>Documented methods for DNA extraction, quantification, labelling, hybridisation, identification, data analysis and reporting using one or a combination of the techniques below by in-house procedures and/or commercially verified procedures: -</p> <p>Microarray Analysis using the Illumina Beadchip CytoSNP- 850k array Platform DNA Extraction From Blood or Saliva using:</p> <p>FujiFilm/Kurabo QuickGene-610L RGS MOT0111<br/>DNA Extraction using the Chemagic Star RGS MOT0128<br/>RGS MOT0112 - DNA extraction using the iGENatal Extraction kit<br/>Quantification of DNA using Qubit fluorometer using RGS MOT0118<br/>Manual Processing of Illumina Beadchip CytoSNP-850K RGS MOT0607<br/>Automated Processing of Illumina Beadchip CytoSNP-850K RGS MOT0609</p> <p>Equipment includes:<br/>High Speed Microplate shaker, Hybridisation Oven, Vortex, pulse centrifuge, heat block, vacuum desiccator, TECAN Evo.<br/>Production and processing of Beadchip Data files using NextSeq550 and Beeline software RGS ANA0410<br/>Equipment includes:<br/>NextSeq 550, Beeline software<br/>Updated analysis document:<br/>Microarray Analysis and Checking using InfoQuant Fusion and cnTrack RGS ANA0408</p> | RGL           |



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| <p>HUMAN TISSUE AND FLUIDS (cont'd)</p> <p>DNA extracted from:<br/>Blood, Saliva, Chorionic villus tissue, Uncultured amniocytes, Cultured cells from chorionic villus samples, Cultured cells from amniotic fluid, Saliva, Thymus (Cont'd)</p> | <p><u>Cytogenetics examination procedures for the purposes of clinical diagnosis (cont'd)</u></p> | <p>Quantative PCR (qPCR) is performed using the StepOnePlus Real Time PCR system to confirm copy number variation detected by microarray and NGS.<br/>SOP(s):</p> <p>RGS MOT0701 Setting up a qPCR using StepOnePlus Real-Time PCR System</p> <p>Analysis by ABI StepOne software. Interpretation by manual in-house method documented in SOP(s):</p> <p>RGS MOT0702 Primer Design for qPCR<br/>RGS ANA0204 Real Time qPCR Data Processing, Analysis and Checking using StepOne Software</p> | <p>RGL</p>    |



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| <p>HUMAN TISSUE AND FLUIDS (cont'd)</p> <p>Whole Blood<br/>Amniotic Fluid<br/>CVS<br/>Tissue Samples</p> | <p><u>Cytogenetics examination procedures for the purposes of clinical diagnosis (cont'd)</u></p> <p>Chromosome analysis for<br/><br/>Prenatal Diagnosis<br/>Reproductive Medicine Disorders<br/>Developmental Disorders</p> | <p>Chromosome analysis - samples are processed either by the Hanabi automated system or by manual in-house method documented in SOP(s):</p> <p>RGS CUT0101 Preparation of blood culture medium and tubes<br/>RGS CUT0102 Setting up blood cultures<br/>RGS CUT0104 Blocking, releasing, and adding Colcemid to blood cultures<br/>RGS CUT0106 Harvesting Blood Cultures Manually<br/>RGS CUT0107 Manual Slidemaking from blood culture cell suspensions<br/>RGS CUT0108 G-banding slides from blood cultures<br/>RGS CUT0110 Harvesting blood cultures using the Hanabi-PIII metaphase harvester<br/>RGS CUT0111 Slide Making using the Hanabi-PIV Metaphase Auto Spreader<br/>RGS CUT0112 Preparation of blood samples for rapid FISH aneuploidy screening and sex determination<br/>RGS CUT0201 Preparation of amniotic fluid samples for culture, QF-PCR and other tests<br/>RGS CUT0202 Preparation of chorionic villus samples for culture, QF-PCR and other tests</p> | RGL           |



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| <p>HUMAN TISSUE AND FLUIDS (cont'd)</p> <p>Whole Blood<br/>Amniotic Fluid<br/>CVS<br/>Tissue Samples</p> | <p><u>Cytogenetics examination procedures for the purposes of clinical diagnosis (cont'd)</u></p> <p>Chromosome analysis for</p> <p>Prenatal Diagnosis<br/>Reproductive Medicine Disorders<br/>Developmental Disorders (cont'd)</p> | <p>RGS CUT0203 Processing amniocytes, chorionic villus and tissue culture cells in culture for DNA extraction or interphase FISH</p> <p>RGS CUT0204 Assessing amniotic fluid, chorionic villus and tissue cultures</p> <p>RGS CUT0205 Medium-changing, topping up and re-seeding amniotic fluid, chorionic villus and tissue cultures</p> <p>RGS CUT0206 Subculture and redistribution of amniotic fluid, chorionic villus and tissue cultures</p> <p>RGS CUT0207 Blocking amniotic fluid, chorionic villus and tissue cultures</p> <p>RGS CUT0208 Harvesting amniotic fluid, chorionic villus and tissue cultures</p> <p>RGS CUT0209 Slide-making from amniotic fluid, chorionic villus and tissue culture cell suspensions</p> <p>RGS CUT0210 G-banding slides from amniotic fluid, chorionic villus and tissue cultures</p> <p>RGS CUT0211 Completing culture sheets</p> <p>RGS CUT0212 Culture Media in Current Use and the Preparation of Complete Culture Media</p> <p>RGS CUT0301 Preparation of tissue samples for interphase FISH, touch preparations and setting up in culture</p> | <p>RGL</p>    |



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| <p>HUMAN TISSUE AND FLUIDS (cont'd)</p> <p>Whole Blood<br/>Amniotic Fluid<br/>CVS<br/>Tissue Samples</p> | <p><u>Cytogenetics examination procedures for the purposes of clinical diagnosis</u> (cont'd)</p> <p>Chromosome analysis for; -<br/>Prenatal Diagnosis<br/>Reproductive Medicine Disorders<br/>Developmental Disorders (cont'd)</p> | <p>Karyotyping analysis is performed manually using microscopes.<br/>SOP(s):<br/>RGS ANA0402 Chromosome Analysis - General Procedures</p> <p>Fluorescence in-situ Hybridisation (FISH) analysis. Probes hybridised manually and used to specific guidelines as documented in SOP(s):</p> <p>RGS MOT0801 Slide-making for FISH and preparation of the FISH process sheet<br/>RGS MOT0802 FISH setting up and hybridisation<br/>RGS MOT0803 Post hybridisation wash procedures, detection of indirectly labelled probes and counterstaining<br/>RGS MOT0804 FISH paperwork preparation and data recording<br/>RGS MOT0806 Molecular cytogenetic reagents and solutions - procedures for preparation</p> <p>Analysis is performed using fluorescence microscopes.<br/>SOP(s):<br/>RGS ANA0407 FISH analysis</p> | <p>RGL</p> <p>RGL</p> <p>RGL</p> |



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## Schedule of Accreditation

issued by

### United Kingdom Accreditation Service

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

### Great Ormond Street Hospital for Children NHS Foundation Trust

Issue No: 006 Issue date: 27 May 2021

Testing performed at main address only

| Materials/Products tested        | Type of test/Properties measured/Range of measurement   | Standard specifications/ Equipment/Techniques used   | Location Code |
|----------------------------------|---|--|---------------|
| HUMAN TISSUE AND FLUIDS (cont'd) | <u>Cytogenetics examination procedures for the purposes of clinical diagnosis</u> (cont'd)                              |  |               |
| Tissue Samples                   | Chromosome analysis for; -<br>Prenatal Diagnosis<br>Reproductive Medicine Disorders<br>Developmental Disorders (cont'd) | Multi-ligation Probe Analysis (MLPA). Performed by manual in-house method or using the Biomek NK span-8 system.<br>SOP(s):<br><br>RGS MOT0401 Setting up MLPA using the Automated system<br>RGS MOT0402 Manually Setting Up an MLPA reaction using MRC Holland SALSA kits (cytogenetics)<br>RGS ANA0201 Capillary Electrophoresis of MLPA and QF-PCR products using the ABI 3730 Genetic Analyser<br>RGS ANA0208 MLPA data processing and analysis using GeneMarker Analysis Software (Cytogenetics) | RGL           |
| Tissue Samples                   |   | Quantitative Fluorescence Polymerase Chain Reaction (QF-PCR). Analysis performed by manual in-house method documented in SOP(s):<br><br>RGS MOT0404 Manually Setting Up a QF-PCR Reaction<br>RGS ANA0201 Capillary Electrophoresis of MLPA and QF-PCR products using the ABI 3730 Genetic Analyser<br>RGS ANA0203 QF-PCR data processing and analysis using Genemarker   | RGL           |





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| <p>HUMAN TISSUE AND FLUIDS (cont'd)</p> <p>Bone Marrow, Peripheral Blood, CSF, Tissue sections</p> | <p><u>Cytogenetics examination procedures for the purposes of clinical diagnosis (cont'd)</u></p> <p>Chromosome analysis for: -<br/>Haematological malignancies including ALL, AML, CML, MPN, MDS, JMML, Lymphoma, solid tumours</p> | <p>G banding and G-band analysis using Leica Cytovision Karyotyping station and in house procedures:</p> <p>HSOP 632<br/>RGS LAB9609<br/>HSOP 614<br/>RGS LAB9604<br/>RGS LAB9601<br/>HSOP 605</p> <p>FISH analysis using Digital Scientific<br/>FISH analysis using system ISIS (Metasystem)</p> <p>Following in house documented procedures:<br/>RGS LAB9630<br/>HSOP 632<br/>HSOP 614<br/>RGS LAB9604<br/>RGS LAB9601</p> | CBL           |
| END  |  |  |               |