

**issued by**

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



**Cambridge University Hospitals NHS Foundation Trust**

**Issue No: 012   Issue date: 27 November 2025**

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### Materials/Products tested

Type of test/Properties  
measured/Range of measurement

Standard specifications/  
Equipment/Techniques used

## Genomic analysis for the purpose of clinical diagnosis of rare and inherited disease and cancer

Documented in house procedures  
incorporating manufacturer's  
instructions (where relevant)

**Sample preparation:**

Sample preparation, DNA extraction, quantification and quality check for subsequent in-house analysis (see below), referral to specialist centres and long term storage

Red Cell lysis and storage in  
Qiagen Nucleon B, RLT or Tri-  
Reagent  
CU-EX-SOP-34, CU-EX-SOP-23,  
CU-EX-SOP-5 and CU-EX-SOP-8

## Separation of mononuclear cells from peripheral blood or bone marrow using Lymphoprep CU-EX-SOP-31

### Manual and automated DNA extraction and quantification using:

**Automated Extraction:**  
Automated DNA extraction using  
Flex Star

CU-EX-SOP-13



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<p><b>HUMAN TISSUES AND BODY FLUIDS (cont'd)</b></p> <p>Whole Blood,Buffy Coat CVS/Amniocentesis Tissues Guthrie Spots</p> <p>Whole Blood Bone Marrow Body Fluid Tissue Lineage specific fractions</p> <p>Body fluid and human tissue (fresh, frozen or fixed) Formalin fixed paraffin embedded tissue Bone Marrow</p> <p>Bone Marrow</p> <p>Saliva and Buccal swabs</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>Genomic analysis for the purpose of clinical diagnosis of rare and inherited disease and cancer</u> (cont'd)</p>	<p>Documented in house procedures incorporating manufacturer's instructions (where relevant)</p> <p>Automated DNA extraction using Qiagen EZ1</p> <p>CU-EX-SOP-9</p> <p>Automated DNA extraction using QIAcube</p> <p>CU-EX-SOP-3 CU-EX-SOP-5</p> <p><b>Manual extraction:</b></p> <p>Manual DNA extraction using Qiagen DNEASY kit CU-EX-SOP-15 CU-EX-SOP-16 CU-EX-SOP-23</p> <p>Extraction of DNA from DNA lysis buffer (Q-AL or nucleon B) CU-EX-SOP-22</p> <p>Manual DNA extraction using- Oragene Purifyer CU-EX-SOP-11</p> <p>DNA Quantification for QC purposes:</p> <p>Nanodrop ND-1000 spectrophotometer and Qubit fluorometer</p> <p>CU-EQ-SOP-9 CU-EX-SOP-21</p>



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<p>HUMAN TISSUES AND BODY FLUIDS (cont'd)</p> <p>DNA Clean up</p> <p>Whole Blood, bone marrow, body fluid and human tissue (fresh, frozen or fixed)</p> <p>Whole Blood, bone marrow, body fluid and human tissue (fresh, frozen or fixed)</p>	<p><u>Genomic analysis for the purpose of clinical diagnosis of rare and inherited disease and cancer</u> (cont'd)</p> <p>RNA extraction and preparation of cDNA, 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>Qiagen QiaAmp DNA micro kit CU-EX-SOP12</p> <p><b>Automated Extraction:</b></p> <p>Automated RNA extraction and conversion to cDNA using QIAcube, NJ Research (Biorad) thermal cycler and Syngene Genius Bioimaging System</p> <p>CU-EX-SOP-3</p> <p>CU-EX-SOP-19</p> <p>CU-EX-SOP-32</p> <p><b>Manual and automated RNA extraction, conversion to cDNA and quantification using:</b></p> <p><b>Manual Extraction</b></p> <p><b>Extraction of RNA from RLT Buffer</b> CU-EX-SOP-24</p>



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<p><b>HUMAN TISSUES AND BODY FLUIDS (cont'd)</b></p> <p>Genomic DNA and cDNA extracted in-house from the sample types listed above or received as primary sample type from an external source</p> <p>Genomic DNA and cDNA received as primary sample type from an external source</p> <p>Illumina TriSight One data previously generated in-house</p> <p>Genomic DNA received as primary sample type from an external source</p>	<p><u>Genomic analysis for the purpose of clinical diagnosis of rare and inherited disease and cancer</u> (cont'd)</p> <p>Detection of nucleic acid sequence variants - SNVs and small indels-</p> <p>Detection of nucleic acid sequence variants - SNVs and small indels-</p> <p>See CU-UK-DTA-3</p> <p>SNVs and small indels See CU-UK-DTA-7</p> <p>Next Generation Sequencing for Small Variants</p>	<p>Documented in house procedures incorporating manufacturer's instructions (where relevant)</p> <p><b>Sanger Sequencing</b></p> <p>Sanger Sequencing Analysis using Mutation Surveyor by SoftGenetics</p> <p>CU-TS-SOP-29</p> <p><b>Sanger Sequencing</b> Using:</p> <p>Standard Primer Design methodology: CU-BI-SOP-2</p> <p>PCR/ Thermal cyclers, ABI 3730, Biomek robotics</p> <p>CU-CM-SOP-5 CU-CM-SOP-13 CU-TS-SOP-28 CU-CM-SOP-31 CU-TS-SOP-45 CU-TS-SOP-65 CU-TS-SOP-58, CU-TS-SOP-62</p> <p>Analysis using Dias Bioinformatics software CU-BI-MAN-1</p> <p>TWIST WES Comprehensive exome panel CU-NG-SOP-8 CU-BI-MAN-2 CU-BI-MAN-1 CU-BI-MAN-6 CU-NG-SOP-14 CU-NG-SOP-10 CU-RP-SOP-9</p>



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HUMAN TISSUES AND BODY FLUIDS (cont'd)	<u>Genomic analysis for the purpose of clinical diagnosis of rare and inherited disease and cancer</u> (cont'd)	Documented in house procedures incorporating manufacturer's instructions (where relevant)
Genomic DNA received as primary sample type from an external source	Next Generation Sequencing	<b>TWIST Core, Endocrine, Neurology (CEN)</b> CU-NG-SOP-14 CU-BI-MAN-1 CU-NG-SOP-8 CU-BI-MAN-2 CU-RP-SOP-9
Whole Blood	Nucleic acid extraction	<b>Manual DNA extraction using Tissue Lyser and Qiagen Mini kit</b> CU-EX-SOP-17 CU-TS-MAN-6
Genomic DNA received pre extracted from an external source	Detection of SNVs and small indels  See CU-UK-DTA-8	<b>Real Time PCR</b>  <b>Using Applied biosystems Realtime 7500 analyser</b>  CU-CM-SOP-21  Using Rotogene 6000 series – QIAgen CU-CM-SOP-16 CU-CM-SOP-26 CU-CM-SOP-22
Formalin fixed paraffin embedded (FFPE) tissue	Sample preparation, DNA and RNA extraction, quantification and quality check for subsequent in-house analysis, referral to specialist centres and long term storage.	<b>Manual or Automated Extraction of nucleic acids from formalin fixed paraffin embedded (FFPE) tissue using MagMax</b> CU-EX-SOP-33



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<p>HUMAN TISSUES AND BODY FLUIDS (cont'd)</p> <p>Genomic DNA received as primary sample type from an external source</p>	<p><u>Genomic analysis for the purpose of clinical diagnosis of rare and inherited disease and cancer</u> (cont'd)</p> <p>Determination of fragment length size and detection of deletions, fusion deletions, known SNVs and indels, microsatellites, repeat expansion, linkage markers for the purpose of genotyping at specific known gene loci</p> <p>See CU-UK-DTA-8</p>	<p>Documented in house procedures incorporating manufacturer's instructions (where relevant)</p> <p><b>Fluorescent Based Fragment Analysis</b></p> <p><b>Using:</b></p> <p>Fluorescent Fragment Analysis using ABI 3730 and ABI 3500 xl genetic analysers and GeneMarker software</p> <p>CU-TS-SOP-34</p> <p>Commercial Methods: Fluorescent Fragment Assays:</p> <p>Fluorescent ARMS using the Elucigene kits CU-TS-SOP-17 and CU-TS-SOP-9</p> <p>QF-PCR kit (QF-PCR) - Devyser Kits CU-TS-SOP-44 CU-TS-SOP-43</p> <p>And:</p> <p>In house Fluorescent Fragment Assays using PCR and allele specific PCR and PCR of linked markers</p> <p>CU-CM-SOP-14 CU-TS-SOP-21 CU-TS-SOP-23 CU-TS-SOP-30 CU-TS-SOP-36 CU-TS-SOP-41</p> <p>Fluorescent Fragment Analysis using and 3730, 3500 genetic analysers and GeneMarker or Genemapper software</p> <p>CU-TS-SOP-34 and CU-RP-SOP-29</p>



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<b>HUMAN TISSUES AND BODY FLUIDS (cont'd)</b>	<u>Genomic analysis for the purpose of clinical diagnosis of rare and inherited disease and cancer</u> (cont'd)	Documented in house procedures incorporating manufacturer's instructions (where relevant)
Genomic DNA received as primary sample type from an external source	Determination of fragment size length, Detection of deletions, fusion deletions, known SNVs, repeat expansions and inversions for the purpose of genotyping at specific known gene loci -  See CU-UK-DTA-4	<b>Gel electrophoresis based Fragment Analysis</b>  In house methods for:  Long range PCR, MS-PCR, inverse shifiting PCR, multiplex PCR followed by gel electrophoresis.  SOP: CU-CM-SOP-15 CU-CM-SOP-44 CU-TS-SOP-23
Genomic DNA received as primary sample type from an external source	Detection of large repeat expansions  See CU-UK-DTA-6	<b>In house methods for:</b> Southern blot hybridization analysis using labelled probes.  CU-TS-SOP-20 CU-TS-SOP-7
Genomic DNA extracted in-house from human body fluids and tissues or received as primary sample type from an external source	Single Nucleotide Polymorphism (SNP) genotyping (C9ORF72 gene)	Nimagen SNP profile kit for identity checking on ABI 3500 and ABI 3730 genetic instruments  CU-WG-SOP-14
Genomic DNA extracted in-house from human body fluids and tissues or received as primary sample type from an external source	Determination of fragment length size and detection of deletions, fusion deletions, known SNVs and indels, microsatellites, repeat expansion, linkage markers for the purpose of genotyping at specific known gene loci in gene C9ORF72.	Fluorescent fragment analysis using ABI 3500 and ABI 3730 genetic instruments: Frontotemporal Dementia (FTD) and/or Amyotrophic Lateral Sclerosis (ALS) (C9ORF72 gene)  CU-TS-SOP-34 CU-RP-SOP-33



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<p>HUMAN TISSUES AND BODY FLUIDS (cont'd)</p> <p>Sequencing data received from internal and external sources</p> <p>Whole blood Amniotic fluid Chorionic villus samples Bone Marrow Biopsy Material Body Fluids</p>	<p><u>Cytogenetic examinations, neoplastic genetics including haemato-oncology, - detection of sub microscopic chromosomal imbalance (gains and losses) expressed as changes to copy number and copy neutral loss/absence of heterozygosity</u></p> <p>Detection of genomic variation with potential clinical utility, including small variants (SNV &amp; indel) and structural variants (CNV, insertion, inversions and translocations resulting in fusion, ITD and PTD)</p> <p>G-Banding/Karyotyping</p> <p>Detection of chromosomal rearrangements or aberrations arising from:</p> <p>Prenatally detected Disorders Developmental disorders Reproductive medicine disorders Haematological/Oncology disorders</p> <p>(Preparative Pre-examination steps listed first)</p>	<p>Documented in house procedures incorporating manufacturer's instructions (where relevant)</p> <p>Detection of variants for clinical reporting of patients with rare disease and cancer: alignment and variant calling. GH-BI-POL-1, GH-BI-POL-2, GH-BI-POL-</p> <p><b>Manual culturing and processing of human tissue to provide metaphase cells:</b> using commercial media and:</p> <p>CU-CT-SOP-2, CU-CT-SOP-22, CU-CT-SOP-8, CU-CT-SOP-5, CU-CT-SOP-17, CU-CT-SOP-14, CU-CT-SOP-20</p> <p>Manual harvesting using in house methods : CU-CT-SOP-3 (whole blood and prenatal and tissue cultures CU-CT-SOP-7). CU-CT-SOP-13</p> <p>Slide Preparation, coverslipping and storage : CU-CT-SOP-4 and CU-CT-SOP-19</p> <p><b>Microscopic Analysis and reporting of G banded chromosomes:</b> Using: Brightfield microscopes and GenASIs image capture and analysis software</p> <p>CU-CT-SOP-25, CU-CT-SOP-26, CU-CT-SOP-27, CU-RP-SOP-38, CU-CT-SOP-15 and CU-RP-SOP-21, CU-RP-SOP-27</p>





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<p><b>HUMAN TISSUES AND BODY FLUIDS (cont'd)</b></p> <p>Genomic DNA received as primary samples from external sources</p> <p>Genomic DNA received as primary sample type from an external source</p> <p>Fixed cell slides from human tissues and body fluids:</p> <p>Amniotic fluid CVS Stemcells Wholeblood Bone Marrow CSF Biopsy Material Body fluids FFPE Tissue</p>	<p><u>Cytogenetic examinations</u> (cont'd)</p> <p>Detection of pathogenic and likely pathogenic DNA copy number variants and copy number neutral loss of heterozygosity (in diagnostic samples)</p> <p>Detection of whole exon deletions/duplications and specific methylation abnormalities</p> <p>See CU-UK-DTA-5</p> <p>Detection and analysis of genomic rearrangements and imbalances. Confirmation of genomic rearrangements - using locus specific probes:</p> <p>Deletion Copy Number/Amplification</p> <p>See CU-UK-DTA-2</p>	<p>Documented in house procedures incorporating manufacturer's instructions (where relevant)</p> <p><b>SNP Array</b> Using Infinium Global Screening Array (GSA) v 3 kit with Illumina Infinium HTS Assay and Tecan Robot Analysis using Illumina iScanner with Illumina Genome Studio software and NxClinical software CU-MA-SOP-1 CU-RP-SOP-4</p> <p><b>Multiplex Ligation Probe Analysis (MLPA) and methylation specific MS-MLPA</b>  using: in house or commercial MRC Holland kits  Thermal cyclers and 3500 / 3730 Genetic Analysers  Analysis using GeneMarker  CU-TS-SOP-37 CU-TS-SOP-16 CU-TS-SOP-42</p> <p><b>Fluorescent in-situ hybridisation (FISH)</b>  Using:  Cytocell and Vysis commercial kits, Vysis HYBrite hotplate and applied imaging (Leica)  Analysis using: Cytovision analysis software and and fluorescent microscope system  CU-FI-SOP-6, CU-FI-SOP-5, CU-FI-SOP-12, CU-FI-SOP-11, CU-FI-SOP-4, and CU-FI-SOP-8</p>
END		