


# Schedule of Accreditation

issued by

## United Kingdom Accreditation Service

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

 <p><b>UKAS</b> MEDICAL 21163</p> <p>Accredited to ISO 15189:2012</p>	<h3>Juno Genetics Limited</h3> <p><b>Issue No:</b> 001    <b>Issue date:</b> 12 October 2021</p>	
	<p><b>Juno Genetics Limited</b> Winchester House Heatley Road Oxford Science Park Oxford OX4 4GE</p>	<p><b>Contact:</b> Dagan Wells <b>Tel:</b> +44 (0)203 743 19944 <b>E-Mail:</b> Dagan.Wells@junogenetics.com <b>Website:</b> www.junogenetics.com</p>
<p><b>Testing performed at the above address only</b></p>		

### DETAIL OF ACCREDITATION

Materials/Products tested	Type of test/Properties measured/Range of measurement	Standard specifications/ Equipment/Techniques used
HUMAN TISSUES AND FLUIDS	<u>Molecular Genetics</u>	
Peripheral Blood, Saliva samples (in Oragene Media) and Products of conception	Extraction procedures: Automated extraction	Qiagen QIAamp kit and Qiacube
Peripheral Blood, Saliva samples (in Oragene Media) and Products of conception	Manual extraction	Qiagen QIAamp kit and Qiacube
Plasma	Extraction of cf DNA	Hamilton Liquid handler and VeriSeq workflow. Quantification using Spectramax Spectrophotometer
Trophectoderm cells	Preimplantation Genetic Testing for Monogenic Disorders (PGT-M)	Using: Repli-g SC or GenomiPi kit for DNA amplification by Multiple Displacement Amplification (MDA)  Infinium SNP array/Karyomapping kit for genotyping the SNPs. Using Illumina NextSeq 550 or Illumina NextSeq 550 DX as a scanner to "read" the genotyped SNP. Analysis using BlueFuse multi software v5.5 (Illumina) EXAM_PGT-M_Primer Design_234 EXAM_PGT_M_SOP_197
DNA extracted from Trophectoderm biopsy, Peripheral Blood or Saliva (also Genomic DNA received from an external source (mother/ father/ reference)	Direct Mutation Testing (for direct mutation site confirmation testing in PGT-M) using in house SOPs and which includes the below and analysis for all using Genemapper  Fragment analysis Minisequencing Expansion analysis Fragile X	In house and kit based methodology and analysis using ABI 3500DX



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<p>HUMAN TISSUES AND FLUIDS (Cont'd)</p> <p>DNA extracted from Trophoctoderm biopsy</p> <p>DNA extracted from Trophoctoderm biopsy</p> <p>DNA extracted from Peripheral Blood and Products of Conception</p>	<p><u>Molecular Genetics</u> (Cont'd)</p> <p>Preimplantation Testing for Aneuploidies (PGT-A) – Chromosome copy number assessment:</p> <p>Preimplantation Genetic Testing for Structural Rearrangements (PGT-SR)</p> <p>Products of Conception (POC) Testing – Chromosome Copy Number assessment Including Maternal Cell Contamination Determination</p>	<p>Hot Master-taq PCR and ABI 3500XL Hot Master-taq clean-up of PCR products using ExoSAP-IT, and Primer extension using SNaPshot and ABI 3500XL</p> <p>FailSafe™ one step PCR and ABI 3500XL</p> <p>AmplideX® PCR/CE FMR1 AmplideX PCR kit</p> <p>PGT-SEQ kit using FEC Library Preparation, Sequencing using Illumina NextSeq 550 and Illumina NextSeq 550 DX, Analysis using NexCCsv2 analysis software EXAM_PGT-Seq SOP_149</p> <p>SurePlex amplification and VeriSeq Library Preparation kits, Sequencing using Illumina NextSeq 550, Illumina NextSeq 550 DX and MiSeq, Analysis using BlueFuse Multi software (Illumina) EXAM_VeriSeqSOP_188</p> <p>PGT-Seq Kit using FEC library Preparation, Sequencing using Illumina NextSeq 550 and Illumina NextSeq 550 DX, Analysis using NexCCsv2 analysis software EXAM_PGT-Seq SOP_149 EXAM_POCSOP_187</p>



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Plasma	Non-invasive prenatal testing (NIPT)	Non Invasive Prenatal Testing (NIPT) for aneuploidy and sexing using VeriSeq automated workflow using VeriSeq Solution v 2.– Automated Cell free DNA extraction from plasma and Library, Preparation using the Hamilton Star Liquid Handler and Paralleled sequencing on a NextSeq 550 DX. Analysis with VeriSeq VeriSeq NIPT Assay Software v2 (for 5 and 24 chromosomes) SOP EXAM_Neo and Neo24 test_SOP_269
END		