


# Schedule of Accreditation

issued by

## United Kingdom Accreditation Service

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

 <p>Accredited to ISO 15189:2012</p>	<b>NewGene Ltd</b>	
	<b>Issue No: 003    Issue date: 31 October 2017</b>	
	<b>Biomedicine West Wing</b> <b>International Centre for Life</b> <b>Times Square</b> <b>Newcastle upon Tyne</b> <b>NE1 4EP</b>	<b>Contact: Jane Llsgo</b> <b>Tel: +44 (0) 0191 242 1923</b> <b>Fax: +44 (0) 0191 241 8799</b> <b>E-Mail: info@newgene.org.uk</b> <b>Website: www.newgene.org.uk</b>
<b>Testing performed at the above address only</b>		

### DETAIL OF ACCREDITATION

Materials/Products tested	Type of test/Properties measured/Range of measurement	Standard specifications/ Equipment/Techniques used
Formalin fixed paraffin-embedded tissue Slide sections Fixed cells DNA	<p><b><u>Molecular pathology</u></b></p> <p>Targeted mutation detection in specific genes, listed below, using DNA</p> <p>BRAF targeted mutation detection (colorectal cancer, hairy cell leukaemia and melanoma)</p> <p>EGFR targeted mutation detection (non-small-cell lung cancer)</p> <p>RAS targeted mutation detection in colorectal tumours including KRAS and NRAS genes</p>	<p>DNA extraction and quantification using in-house procedures or commercial kits (Promega Maxwell or Qiagen) according to the following procedures:            NG-PREEXAM-26            NG-PREEXAM-30            NG-EQUIP-07</p> <p>PCR amplification of DNA and mutation detection by MALDI-TOF mass spectrometry using a Sequenom MA4 system according to the following procedures:            NG-EXAM-SEQ-01            NG-EXAM-SEQ-02            NG-EXAM-SEQ-03            NG-EXAM-SEQ-05            NG-EXAM-SEQ-18            NG-POST-SEQ-01</p> <p>Fragment analysis (EGFR deletions and insertions) using PCR amplification, detection on an ABI 3139xl capillary analyser and data analysis using GeneMarker software according to the following procedureS:            NG-EXAM-SEQ-04            NG-IT-07            NG-POST-03</p>



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Whole Blood	<p><b><u>Molecular haematology</u></b></p> <p>Analysis of DNA and RNA</p> <p>BCR-ABL monitoring in chronic myeloid leukaemia</p> <p>JAK2, MPL and calreticulin targeted mutation detection in myeloproliferative neoplasms</p> <p>TPMT genotyping</p>	<p>DNA and RNA extraction and quantification using in-house procedures or commercial kits (Promega Maxwell or Qiagen) according to the following procedures:            NG-PREEXAM-07            NG-PREEXAM-10            NG-PREEXAM-34            NG-PREEXAM-35            NG-PREEXAM-36            NG-EQUIP-07</p> <p>Amplification of RNA by real time PCR on a Roche LC480 light cycler using a commercial kit (Ipsogen) and data analysis using Roche software according to procedures:            NG-EXAM-BCR-01            NG-EXAM-BCR-01            NG-EXAM-BCR-01</p> <p>PCR amplification of DNA. Mutation detection by MALDI-TOF mass spectrometry using a Sequenom MA4 system NG-EXAM-SEQ-01            NG-EXAM-SEQ-02</p> <p>NG-EXAM-FLX-14            NG-EXAM-SEQ-13            NG-POST-08            NG-POST-SEQ-01</p> <p>Fragment analysis (calreticulin) using PCR amplification and detection on an ABI 3130xl capillary analyser according to the following procedures:            NG-EXAM-SEQ-16</p>



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<p>Whole Blood Mouth washes DNA</p>	<p><b><u>Molecular Genetics</u></b></p> <p>Analysis of DNA and RNA for mutations in genes underlying hereditary diseases listed below.</p> <p>Breast cancer: Full gene sequencing of the BRCA1 and BRCA2 genes</p> <p>Noonan spectrum disorders: Full gene sequencing for the following genes - PTPN11, SPRED1, KRAS, SOS1, RAF1, NRAS, BRAF, SHOC2 (exon 1 only), MAP2K1, MAP2K2, HRAS, CBL, NF1, RIT1 and A2ML1</p> <p>Familial hypercholesterolaemia targeted mutation detection in genes APOB, PCSK9 and LDLR</p>	<p>DNA and RNA extraction and quantification using in-house procedures or commercial kits (Promega Maxwell or Qiagen) according to the following procedures: NG-PREEXAM-07 NG-PREEXAM-10 NG-PREEXAM-18 NG-PREEXAM-34 NG-PREEXAM-35 NG-PREEXAM-36 NG-EQUIP-04 NG-EQUIP-07</p> <p>PCR amplification using a Multiplicom system (breast cancer) or sequence capture using Haloplex (Noonans) and next generation sequencing using an Illumina MiSeq. Data analysis using commercial softwares (NextGENe) and in-house methods. Procedures: NG-EXAM-MISEQ-01 NG-EXAM-MISEQ-05 NG-EXAM-MISEQ-06 NG-EXAM-MISEQ-09 NG-EXAM-FLX-14 NG-POST-05 NG-POST-MISEQ-01 NG-POST-MISEQ-05 NG-POST-MISEQ-06</p> <p>PCR amplification of DNA and mutation detection by MALDI-TOF mass spectrometry using a Sequenom MA4 system according to the following procedures: NG-EXAM-SEQ-01 NG-EXAM-SEQ-02 NG-EXAM-SEQ-14</p>

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