

MOLECULAR HAEMATOLOGY – FFPE SAMPLE GENETIC TEST REQUEST FORM

Contact details: North East Scotland Genetics Service, NHS Grampian, Polwarth Building, Foresterhill, Aberdeen AB25 2ZD.

Tel: 01224 553893 / 553820. Email: gram.molgen@nhs.scot web: www.nhsgrampian.org/medicalgenetics

Essential Patient Demographics (Patient label can be used)																																							
Forename:		Surname:																																					
CHI No.:		Date of Birth:		Male / Female (Circle as appropriate)																																			
Address (must include postcode):			Postcode:																																				
Essential Sample Information																																							
Reason for Referral: (Please include histological diagnosis and clinical information)																																							
Referring Clinician(s):		Ward / Clinic:		Referring Pathologist(s):																																			
External Pathology No. (If appropriate):	Aberdeen Pathology No. (Please include PB/PD):	Block / Part No.:	Estimation of Tumour %*:	Scrape all / Marked area* (Circle as appropriate)																																			
Tumour Sample Type (circle as appropriate): Biopsy / Resection / Cytology / Other (please state):																																							
Tumour Sample Site:			Primary or Metastasis (circle as appropriate)																																				
<p style="color: red;">* Tumour tissue: estimation of tumour content is essential; macrodissection to maximise tumour content is desirable, where possible</p> <p style="text-align: center;">Test(s) requested (Please tick)</p> <p>Please check the SSNGM Cancer Test Directory for available testing and referral criteria https://www.genomics.nhs.scot/test-directories/</p>																																							
<p style="text-align: center;">Targeted Tests</p> <table border="1"> <tr> <td rowspan="2">Clonality</td> <td><i>Ig</i> rearrangement</td> <td></td> </tr> <tr> <td><i>TCR</i> rearrangement</td> <td></td> </tr> <tr> <td>CLL / SLL</td> <td><i>TP53</i> only</td> <td></td> </tr> <tr> <td>LPL</td> <td><i>MYD88</i></td> <td></td> </tr> <tr> <td>Systemic mastocytosis</td> <td><i>KIT</i></td> <td></td> </tr> <tr> <td>Hairy cell leukaemia</td> <td><i>BRAF</i></td> <td></td> </tr> </table>			Clonality	<i>Ig</i> rearrangement		<i>TCR</i> rearrangement		CLL / SLL	<i>TP53</i> only		LPL	<i>MYD88</i>		Systemic mastocytosis	<i>KIT</i>		Hairy cell leukaemia	<i>BRAF</i>		<p style="text-align: center;">Lymphoid NGS sub-panel [DNA sequence variants only, see page 2 for genes]</p> <table border="1"> <tr> <td>Chronic Lymphoid Neoplasms</td> <td></td> </tr> <tr> <td>B-cell Lymphoma</td> <td></td> </tr> <tr> <td>T-cell Lymphoma</td> <td></td> </tr> <tr> <td>Plasma Cell Myeloma</td> <td></td> </tr> <tr> <td>Histiocytic Neoplasms</td> <td></td> </tr> </table>	Chronic Lymphoid Neoplasms		B-cell Lymphoma		T-cell Lymphoma		Plasma Cell Myeloma		Histiocytic Neoplasms		<p style="text-align: center;">NGS to assist with:</p> <table border="1"> <tr> <td>Diagnosis</td> <td></td> </tr> <tr> <td>Prognosis / risk</td> <td></td> </tr> <tr> <td>Treatment</td> <td></td> </tr> <tr> <td>Discussed at MDT?</td> <td></td> </tr> </table>	Diagnosis		Prognosis / risk		Treatment		Discussed at MDT?	
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Additional information:			Clinical justification for lymphoid NGS testing [essential for test activation]:																																				
<p style="color: red;">* PLEASE PROVIDE 3 x 5µm PER TEST, WHERE POSSIBLE *</p> <p style="color: red;">Pathology Audit trail (2 staff members must check that the correct block is cut and slides labelled correctly) ** please use Genetics microtome **</p>																																							
Date request received:		Block collected/Apex checked by:	Block cut by:	Block / slide label check: QA1:- QA2:-																																			
Date given to Pathologist for marking:			Date to Genetics:																																				
Reports will be made available on SCI Store																																							
Incomplete or illegible referral forms may lead to sample rejection and a delay in testing																																							

Table 1. Lymphoid NGS subpanel content

Lymphoid DNA subpanel	Genes
Chronic Lymphoid Neoplasms [CLN]	<i>ATM, BCL2, BIRC3, BRAF, BTK, CARD11, CCND1, CD79B, CDKN2A, CXCR4, EZH2, FBXW7, KLF2, KLHL6, KRAS, MAP2K1, MYD88, NOTCH1, NOTCH2, NRAS, PIK3CA, PLCG2, POT1, SF3B1, TNFAIP3, TP53, XPO1</i>
B-cell Lymphoma [BCL]	<i>ARID1A, ATM, B2M, BCL2, BIRC3, BRAF, BTK, CARD11, CCND1, CCND3, CCR6, CD58, CD79A, CD79B, CDKN2A, CREBBP, CRLF2, CXCR4, EP300, ETV6, EZH2, FBXW7, FOXO1, GNA13, GPR34, ID3, JAK2, KIT, KLF2, KLHL6, KMT2D, KRAS, MAP2K1, MYC, MYD88, NOTCH1, NOTCH2, NRAS, PIK3CA, PIM1, PLCG2, POT1, PTEN, RHOA, SF3B1, SOCS1, STAT6, SYK, TCF3, TNFAIP3, TNFRSF14, TP53, XPO1</i>
T-cell Lymphoma [TCL]	<i>ARID1A, CARD11, CD28, CDKN2A, CXCR4, DNMT3A, EP300, ETV6, EZH2, FOXO1, IDH1, IDH2, JAK1, JAK3, KIT, KMT2D, KRAS, NOTCH1, NRAS, PLCG1, RHOA, STAT3, STAT5B, TET2, TP53</i>
Plasma cell myeloma [PCM]	<i>BIRC3, BRAF, CCND1, CDKN2A, CXCR4, IDH1, IDH2, IKZF1, KRAS, MYC, MYD88, NRAS, PIK3CA, PIM1, STAT3, TP53</i>
Histiocytic Neoplasms [HIS]	<i>BRAF, CDKN2A, KRAS, MAP2K1, NRAS, PIK3CA, PIK3CD, PTEN, PTPN1</i>