

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)		
* Tumour tissue: estimation of tumour content is essential; macrodissection to maximise tumour content is desirable, where possible				
Test(s) requested (Please tick)				
Please check the SSNGM Cancer Test Directory for available testing and referral criteria https://www.genomics.nhs.scot/test-directories/				
Targeted Tests			Lymphoid NGS sub-panel [DNA sequence variants only, see page 2 for genes]	NGS to assist with:
Clonality	Ig rearrangement		Chronic Lymphoid Neoplasms	Diagnosis
	TCR rearrangement			
CLL / SLL	TP53 only		B-cell Lymphoma	Prognosis / risk
LPL	MYD88		T-cell Lymphoma	Treatment
Systemic mastocytosis	KIT		Plasma Cell Myeloma	
Hairy cell leukaemia	BRAF		Histiocytic Neoplasms	Discussed at MDT?
Additional information:			Clinical justification for lymphoid NGS testing [essential for test activation]:	
Other tests - please state tumour type, analysis required and reason:				
* PLEASE PROVIDE 3 x 5µm PER TEST, WHERE POSSIBLE *				
Pathology Audit trail (2 staff members must check that the correct block is cut and slides labelled correctly) ** please use Genetics microtome **				
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>