

TEST REQUISITION FORM

[Bone Marrow, Flow cytometry, Molecular haematooncology, Cytogenetics]

Patient Details

Patient's Name: _____ **Date:** _____
Age: _____ **Sex:** M / F / Others **Sample Type:** PB EDTA Heparin
 BM EDTA Heparin
Contact No: _____ **Others:** _____

Referring Clinician

Referred by: _____ **Contact No:** _____

Suspected Diagnosis

- | | | | | |
|--|-----------------------------------|--|--------------------------------|---|
| <input type="radio"/> Acute leukaemia | <input type="radio"/> JMML | <input type="radio"/> ETP-ALL | <input type="radio"/> MDS | <input type="radio"/> CML |
| <input type="radio"/> CMML | <input type="radio"/> B-NHL | <input type="radio"/> CLL | <input type="radio"/> AML | <input type="radio"/> T-NHL |
| <input type="radio"/> Acute leukaemia of ambiguous lineage | <input type="radio"/> MPAL | <input type="radio"/> Lymphoma | <input type="radio"/> MDS/MPN | <input type="radio"/> Follicular lymphoma |
| <input type="radio"/> Chronic leukaemia | <input type="radio"/> CML with BC | <input type="radio"/> ?atypical cells | <input type="radio"/> ?Relapse | <input type="radio"/> MPN |
| <input type="radio"/> Unknown | <input type="radio"/> ALL | <input type="radio"/> Others: please specify _____ | | |

Tests to be performed

Bone Marrow

- BM procedure aspiraton and reporting BM aspirate reporting BM biopsy reporting/review

Flow Cytometry

- | | | |
|--|--|--|
| <input type="checkbox"/> Acute leukaemia panel | <input type="checkbox"/> Chronic leukaemia panel | <input type="checkbox"/> Lymphocyte subset analysis (T, B, NK) |
| <input type="checkbox"/> CD34 enumeration | <input type="checkbox"/> MRD Panel | <input type="checkbox"/> CD19/CD20 (B) |
| <input type="checkbox"/> Acute/chronic leukaemia panel | <input type="checkbox"/> PNH testing | <input type="checkbox"/> CD16+56 (NK) |

Molecular haematooncology

- | | |
|--|--|
| <input type="checkbox"/> BCR-ABL1 (IS) Quantitative Analysis (p210) | <input type="checkbox"/> PML-RARA detection (Quantitative Analysis) |
| <input type="checkbox"/> BCR-ABL1 multiplex for detection of transcripts Qualitative | <input type="checkbox"/> PML-RARA detection (Qualitative Analysis) |
| <input type="checkbox"/> Minor BCR-ABL1 p190 Qualitative | <input type="checkbox"/> Chimerism Study |
| <input type="checkbox"/> Minor BCR-ABL1 p190 Quantitative | <input type="checkbox"/> IgVH mutation analysis for CLL |
| <input type="checkbox"/> Imatinib resistance mutation analysis (IRMA) | <input type="checkbox"/> Onco Haem panel by NGS DNA + RNA (AML, MPN, MDS/MPN, MDS, CMML, JMML, aCML, ET, PMF, PV, CEL, MDS/MPN-RS-T) |
| <input type="checkbox"/> JAK2 Panel : Exons 12 to 15 (includes V617F) | <input type="checkbox"/> Onco Haem - RNA ONLY by NGS |
| <input type="checkbox"/> JAK2 mutation study (V617F only) | <input type="checkbox"/> Onco Haem - DNA ONLY by NGS |
- Sample preservation for DNA RNA

DNA		RNA
HOTSPOT GENES	FULL GENES	FUSION DRIVER GENES
ABL1, BRAF, CBL, CSF3R, DNMT3A, FLT3, GATA2, HRAS, IDH1, IDH2, JAK2, KIT, KRAS, MPL, MYD88, NPM1, NRAS, PTPN11, SETBP1, SF3B1, SRSF2, U2AF1, WT1	ASXL1, BCOR, CALR, ETV6, EZH2, IKZF1, Nf1, PHF6, PRPF8, Rb1, RUNX1, SH2B3, STAG2, TET2, Tp53, ZRSR2	ABL1, ALK, BCL2, BRAF, CCND1, CREBBP, EGFR, ETV6(TEL), FGFR1, FGFR2, FUS, HMGA2, JAK2, KMTA2 A(MLL), MECOM, MLLT10, MLLT3, MYBL1, MYH11, NTRK3, NUP214, PDGFRA, PDGFRB, RARA, RBM15, RUNX 1(AML1) TCF3(E2A), TFE3

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Cytogenetics

- | | | |
|--|--|--|
| <input type="checkbox"/> Bone Marrow Karyotyping | <input type="checkbox"/> FISH for CLL panel (del(17p)(TP53), del(13q), del(11q), Trisomy 12) | <input type="checkbox"/> FISH for 11q (ATM) |
| <input type="checkbox"/> FISH for IGH/BCL2 [t(14;18)] | <input type="checkbox"/> FISH for del(5q) [5q31/5q33/5p15] | <input type="checkbox"/> FISH for 17p (p53) |
| <input type="checkbox"/> Fish for MYEOV/IGH [t(11;14)] | <input type="checkbox"/> FISH for del(7q) [7q22/7q36] | <input type="checkbox"/> FISH for E2A/TCF3 detection |
| <input type="checkbox"/> FISH for BCL6 | <input type="checkbox"/> FISH for BCR-ABL1 [t(9;22)] | <input type="checkbox"/> FISH for IgH |
| <input type="checkbox"/> FISH for AML panel [inv(16), MLL/KMT2A, PML-RARA, AML/ETO(RUNX1-RUNX1T1)] | <input type="checkbox"/> FISH for AML1/ ETO(RUNX1-RUNX1T1) [t(8;21)] | <input type="checkbox"/> FISH for inv(16) [CBFB-MYH11] |
| <input type="checkbox"/> FISH for ALL panel[E2A/TCF3, MLL/KMT2A, BCR-ABL1, TEL-AML1](ETV6-RUNX1) | <input type="checkbox"/> FISH for MDS [del(5q), del(7q), del(20q)] | <input type="checkbox"/> FISH for MM [IGH, del(17p)(TP53), del(13q), del(11q), Trisomy 12] |
| <input type="checkbox"/> FISH for Trisomy 8 | <input type="checkbox"/> FISH for MLL/KMT2A (breakapart) | <input type="checkbox"/> FISH for PDGFR ALPHA |
| <input type="checkbox"/> FISH for PML-RARA detection [t(15;17)] | <input type="checkbox"/> FISH for TEL- AML1 [t(12;21), ETV6/RUNX1] | <input type="checkbox"/> FISH for PDGFR BETA |
| <input type="checkbox"/> Preserve sample till pellet stage | | |
| <input type="checkbox"/> Other: please specify _____ | | |

TIME POINT

At Diagnosis: Yes / No

If under treatment, mention the time point with date of diagnosis

Presenting Complaints

Organomegaly

Liver:

Spleen:

LN:

If LN present, specify:

Treatment History

Transfusion history (Yes/No/If yes, date of last BT): _____

Family History: _____

Other investigations done elsewhere

(CBC/BM/IPT/Cytogenetics/FISH/Molecular/Biochemistry& Serology):

Signature of Clinician

*Please note: The samples must reach the lab within 12-24 hours of collection

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