

OUTLINED AREAS MUST BE COMPLETED

PATIENT IDENTIFIER	PATIENT IDENTIFIER		
	NAME, LAST (Please Print)		FIRST M.I.
	BIRTHDATE	M/F	DATE/TIME COLLECTED
	STREET		PHONE #
	CITY	STATE	ZIP

BILLING	INSURANCE CARRIER NAME		ADDRESS		
	INSURED NAME		INSURED ID#	PT. RELATIONSHIP TO INSURED: <input type="checkbox"/> Self <input type="checkbox"/> Spouse <input type="checkbox"/> Dependent	
	GROUP # or NAME		<input type="checkbox"/> INSURANCE CHANGE		
	<input type="checkbox"/> MEDICARE #		<input type="checkbox"/> MEDICAID #		<input type="checkbox"/> SELF-PAY

STAT **AFFIX TO SPECIMEN CONTAINER**
I attest that this patient has been informed about and has given consent for the test(s) I have ordered below under applicable law.

X _____
PHYSICIAN SIGNATURE

REPORT COPY TO: _____

Please fill out this form as completely as possible in order to prevent delays in processing the sample. Hours of Operation: M-F 7am to 5pm

DIAGNOSIS CODES (MUST BE PROVIDED)	

PATIENT STATUS

New Diagnosis Remission Post BMT/SCT: _____
 Post Treatment Relapse Male Donor Female Donor

Clinical History: _____
Diagnosis: _____
PB-WBC Count: _____ **Blast Count:** _____
Date and Time Specimen Collected: _____
Specimen Type: BM Aspirate BM-Bx BM Smear
 Blood PB Smear CSF Fluid, indicate type: _____
 Lymph Node Other Tissue, indicate type: _____
 Fresh Tissue FNA Touch Prep FFPE

FLOW CYTOMETRY (Send to CFAM)
Diagnosis/History Info REQUIRED

HLX Flow Cytometry CASE Leukemia/Lymphoma Immunophenotyping - Green (preferred) or Lavender top
Rule Out: AML ALL MDS NHL LGL HCL
 CLL Myeloma Other: _____

THS T-Cell Subset - Monitor (CD3, CD4, CD8)
 FTHS Full T-Cell Subset (CD3, CD4, CD8, CD19, CD16,56)
 PNH (Lavender top-PB Only)
 CD34 Stem Cell (CD34 ABS)
 CD3 Absolute

MOLECULAR ONCOLOGY (Send to NSUH Molecular Pathology)

HLX IGR B-Cell Gene Rearrangement
 HLX TCR T-Cell Gene Rearrangement
 HLX BCR Quantitative *BCR::ABL1* p210 p190
 HLX JAK2 JAK2 V617F Mutation Quantitative
 HLX JAK2RLX JAK2 Quantitative JAK2 V617F mutation with reflex to Exon 12-15

HLX CALR CALR Mutation Analysis
 HLX MPL MPL Mutation Analysis
 HLX FLT3 FLT3 ITD-TKD Mutation Analysis
 HLX WM/LPL MYD88 and CXCR4 Panel Mutation Analysis
 HLX MYD88 MYD88 L265P Mutation Analysis
 HLX CXCR4 CXCR4 C1013G Mutation Analysis
 DNA and HOLD BRLI Myeloid Panel
 Other: _____

CYTOGENETICS AND FISH (Send to NSUH Cytogenetics)

Test Requested: HLX Chromosome Oncology CASE Cytogenetics
 HLX FISH Oncology CASE

For FISH panels, check box against specific panel.
For individual probes check box next to specific probes.

<input type="checkbox"/> S/P Sex Mismatched	<input type="checkbox"/> MDS Panel
<input type="checkbox"/> Bone Marrow Transplant	<input type="checkbox"/> -5 /deletion 5q
<input type="checkbox"/> X / Y Probes	<input type="checkbox"/> -7 /deletion 7q
<input type="checkbox"/> MPD Panel	<input type="checkbox"/> CEP8 (Trisomy 8)
<input type="checkbox"/> <i>PDGFRA::FIP1L1</i> fusion <i>CHIC2</i> del(4q12)	<input type="checkbox"/> Deletion 20q
<input type="checkbox"/> <i>PDGFRB</i> Break Apart (5q32)	<input type="checkbox"/> <i>TP53</i> (deletion 17p)
<input type="checkbox"/> <i>FGFR1</i> Break Apart (8p11.23)	<input type="checkbox"/> AML Panel #1
<input type="checkbox"/> CML - BCR::ABL1 t(9;22)	<input type="checkbox"/> -5/ deletion 5q
<input type="checkbox"/> CLL Panel	<input type="checkbox"/> -7/ deletion 7q
<input type="checkbox"/> <i>MYB</i> (deletion 6q)	<input type="checkbox"/> <i>RUNX1T1::RUNX1</i> t(8;21)
<input type="checkbox"/> <i>ATM</i> (deletion 11q)	<input type="checkbox"/> <i>KMT2A</i> Break Apart (11q23)
<input type="checkbox"/> <i>CCND1::IGH</i> t(11;14)	<input type="checkbox"/> <i>PML::RARA</i> t(15;17)
<input type="checkbox"/> CEP12 (Trisomy 12)	<input type="checkbox"/> <i>CBFB</i> inv(16)/t(16;16)
<input type="checkbox"/> D13S319/ <i>LAMP1</i> (deletion 13q)	<input type="checkbox"/> <i>TP53</i> (deletion 17p)
<input type="checkbox"/> <i>TP53</i> (deletion 17p)	<input type="checkbox"/> AML Panel #2
<input type="checkbox"/> Multiple Myeloma Panel	<input type="checkbox"/> <i>NUP98</i> Break Apart (11p15.4)
<input type="checkbox"/> 1p32(<i>CDKN2C</i>)/1q21(<i>CKS1B</i>)	<input type="checkbox"/> <i>KMT2A</i> Break Apart (11q23)
<input type="checkbox"/> <i>FGFR3::IGH</i> t(4;14)	<input type="checkbox"/> ALL Panel #1
<input type="checkbox"/> <i>CCND1::IGH</i> t(11;14)	<input type="checkbox"/> <i>MYC</i> Break Apart (8q24)
<input type="checkbox"/> <i>RB1</i> (deletion 13q)	<input type="checkbox"/> <i>BCR::ABL1</i> t(9;22)
<input type="checkbox"/> <i>IGH::MAF</i> t(14;16)	<input type="checkbox"/> <i>KMT2A</i> Break Apart (11q23)
<input type="checkbox"/> <i>IGH::MAFB</i> t(14;20)	<input type="checkbox"/> <i>ETV6::RUNX1</i> t(12;21)
<input type="checkbox"/> <i>TP53</i> (deletion 17p)	<input type="checkbox"/> ALL Panel #2
<input type="checkbox"/> Lymphoma Panel	<input type="checkbox"/> <i>ABL1</i> Break Apart (9q34)
(Surgical Pathology #): _____	<input type="checkbox"/> <i>BCL6</i> Break Apart (3q27)
<input type="checkbox"/> <i>BCL6</i> Break Apart (3q27)	<input type="checkbox"/> <i>MYC::IGH</i> t(8;14)
<input type="checkbox"/> <i>MYC</i> Break Apart (8q24)	<input type="checkbox"/> <i>CCND1::IGH</i> t(11;14)
<input type="checkbox"/> <i>IGH</i> Break Apart (14q32)	<input type="checkbox"/> <i>BIRC3::MALT1</i> t(11;18)
<input type="checkbox"/> <i>IGH::BCL2</i> t(14;18)	
<input type="checkbox"/> MDM2/CEP12 for Liposarcoma (Surgical Pathology #): _____	

NSUH Cytogenetics will reference to LIJ Cytogenetics

SPECIMEN PROCESSING DATA – FOR LAB USE ONLY

Flow Has _____
Cytogenetics Has _____
Molecular Has _____
Date Received: _____
Specimen Quality: _____
Specimen Quantity: _____
Culture Medium Used: MarrowMAX _____
Number of Cultures: 1 2 3 _____
Culture Duration: _____

2006260601 (12/5/24) 1.2

Instructions for Specimen Collection, Transportation and/or Storage
Please label all tubes with patient name, date of birth and date of collection

Flow Cytometry

Leukemia/Lymphoma Specimens	Tube Type	Handling and Storage Conditions
Peripheral Blood (5ml)	(2) Green Top (Sodium Heparin) (2) Lavender Top (EDTA)	Room Temperature 18-25°C; deliver to Lab within 12 hours
Bone Marrow (0.5 ml min)	(2) Green Top (Sodium Heparin) (2) Lavender Top (EDTA)	Room Temperature 18-25°C; deliver to Lab within 12 hours
Fine Needle Aspirates/FNA	RPMI/15 ml Conical	Refrigerate at 2-8°C; deliver to Lab immediately
Tissues (0.5 cm ³)	RPMI/15 ml Conical	Refrigerate at 2-8°C; deliver to Lab immediately
CSF (3ml)	Sterile Container	Refrigerate at 2-8°C; deliver to Lab immediately
Bronchial Lavage/ BAL (3-5ml)	Sterile Container , Lavender (EDTA) or Green Top (Sodium Heparin)	Refrigerate at 2-8°C; deliver to Lab immediately
Other Body Fluids (3-5ml)	Sterile Container, Lavender (EDTA) or Green Top (Sodium Heparin)	Refrigerate at 2-8°C; deliver to Lab immediately
Subsets/T-cell – Peripheral blood	(1) Lavender Top (EDTA)	Room Temperature 18-25°C; deliver to Lab within 12 hours
CD34 Stem Cells / CD3 Absolute	(1) Lavender Top (EDTA)	Refrigerate at 2-8°C; deliver immediately
PNH – Peripheral blood	(2) Lavender Top (EDTA)	Room Temperature 18-25°C (Lay tube flat); deliver to Lab immediately

Chromosome Analysis and FISH

Specimen Type	Tube Type	Amount	Handling and Storage Conditions
Peripheral Blood	Green Top (Sodium Heparin)	5-10 ml	Send to the laboratory ASAP at room temperature 20-25°C. If there is a delay in transportation, store the sample in a refrigerator at 4°C / DO NOT FREEZE
Bone Marrow	Green Top (Sodium Heparin)	Minimum 2 ml with sufficient spicules	
Lymph Node	Sterile specimen container with sterile saline or RPMI 1640.	At least 0.5cm ³ piece separated aseptically	
Tumor tissue	Sterile specimen container with sterile saline or RPMI 1640.	At least 0.5cm ³ piece separated aseptically	
Touch Prep, BM/PB smear for FISH analysis	Prepare six smears on a clean positively charged glass slide. 1 smear per probe required. Immediately air dry or send it fresh to the laboratory ASAP.		Send to the laboratory ASAP at room temperature
Tissue block/Slides	4 unstained slides of FFPE tissue section specimen 4-5 micron thick with two unique identifiers, block ID and cut number clearly labeled on the positively charged slides are required. An additional Hematoxylin and Eosin (H&E) slide must be submitted along with the sample clearly marked by a pathologist to denote the area of interest for the FISH study.		Send to the laboratory ASAP at room temperature

Molecular Oncology

Molecular Specimens	Tube Type	Handling and Storage Conditions
Peripheral Blood (at least 2ml)	Lavender Top (EDTA)	Room Temperature 20-25°C or refrigerate at 4°C / DO NOT FREEZE
Bone Marrow	Lavender Top (EDTA)	Room Temperature 20-25°C or refrigerate at 4°C / DO NOT FREEZE
Tissues	Cut 10 sections at 10 micron thick, place in a 1.5ml microfuge tube or on slides.	Room Temperature 20-25°C
BCR::ABL1– Peripheral blood/Bone Marrow – (at least 3 ml)	Lavender Top (EDTA)	Refrigerate at 4°C, deliver to lab within 72 hours/ DO NOT FREEZE
Pleural Fluid, submit at least 0.5x10 ⁷ cells	collected in sterile tube	Transport at 4°C
FNA, submit at least 0.5x10 ⁷ cells	collected in sterile tube	Transport at 4°C