

Medical Record # _____

PATIENT HISTORY AND REQUEST FORM FOR CANCER CYTOGENETIC ANALYSIS

Patient Name: _____

Date of Birth: _____

Address: _____

Race: _____

Phone: _____

Sex: _____

Requesting Physician: _____

Phone: _____

Billing Information: _____

Phone: _____

(Please attach additional documents as needed)

Reason for studies: _____

Sample: Bone marrow aspirate

Collected Date: _____

Peripheral blood

Time: _____

Bone marrow core biopsy

Studies requested Please check (✓) those that apply		
Chromosome studies: _____ FISH studies (Please select from the list below)	aCGH+SNP study: _____	
<input type="checkbox"/> t(1q25.2)- <i>ABL2</i> rearrangement	<input type="checkbox"/> t(11p15.4)- <i>NUP98</i> rearrangement	<input type="checkbox"/> 1q21.3 gain- <i>CKS1B</i>
<input type="checkbox"/> t(1;19)- <i>TCF3/PBX1</i> fusion	<input type="checkbox"/> t(11;14)- <i>IGH/CCND1</i> fusion	<input type="checkbox"/> del(4q12)- <i>FIP1L1/IPDGFRB</i> fusion
<input type="checkbox"/> inv/t(3q26.2)- <i>MECOM</i> rearrangement	<input type="checkbox"/> t(12p13)- <i>ETV6</i> rearrangement	<input type="checkbox"/> del(5q31)- <i>EGR1</i>
<input type="checkbox"/> t(4;14)- <i>IGH/FGFR3</i> fusion	<input type="checkbox"/> t(12;21)- <i>ETV6/RUNX1</i> fusion	<input type="checkbox"/> del(7q,-7 - <i>ELN</i> ,D7S486
<input type="checkbox"/> t(5q32)- <i>PDGFRB</i> rearrangement	<input type="checkbox"/> t(14q11.2)- <i>TCRA/D</i> rearrangement	<input type="checkbox"/> del(9p21)- <i>CDKN2A</i> [p16]
<input type="checkbox"/> t(6;9)- <i>DEK/INUP214</i> fusion	<input type="checkbox"/> t(14q32.3)- <i>IGH</i> rearrangement	<input type="checkbox"/> del(11q22.3)- <i>ATM</i>
<input type="checkbox"/> t(8p11.2)- <i>FGFR1</i> rearrangement	<input type="checkbox"/> t(14;16)- <i>IGH/MAF</i> fusion	<input type="checkbox"/> del(13q14.2) [MM] - <i>RB1</i>
<input type="checkbox"/> t(8q24)- <i>MYC</i> rearrangement	<input type="checkbox"/> t(14;18)- <i>IGH/BCL2</i> fusion	<input type="checkbox"/> del(13q14.3) [CLL] -D13S319
<input type="checkbox"/> t(8;14)- <i>IGH/MYC</i> fusion	<input type="checkbox"/> t(15;17)- <i>PML/RARA</i> fusion	<input type="checkbox"/> del(17p13.1)- <i>TP53</i>
<input type="checkbox"/> t(8;21)- <i>AML1/ETO</i> fusion	<input type="checkbox"/> inv(16)- <i>CBFB</i> rearrangement	<input type="checkbox"/> del(20q12)-D20S108
<input type="checkbox"/> t(9;22)- <i>BCR/ABL1</i> fusion	<input type="checkbox"/> t(22q12)- <i>EWSR1</i> rearrangement	<input type="checkbox"/> X/Y for BMT
<input type="checkbox"/> t(9p24.1)- <i>JAK2</i> rearrangement	<input type="checkbox"/> t(Xp22.33/Yp11.32)- <i>CRLF2</i> rearrangement	<input type="checkbox"/> Trisomy 8
<input type="checkbox"/> t(9q34.1)- <i>ABL1</i> rearrangement	<input type="checkbox"/> t(X;Y;14)- <i>IGH/CRLF2</i> fusion	<input type="checkbox"/> Trisomy 4/10/17
<input type="checkbox"/> t(11q23)- <i>MLL</i> [<i>KMT2A</i>] rearrangement	<input type="checkbox"/> del(Xp22.3/Yp11.3)- <i>P2RY8/CRLF2</i> fusion	<input type="checkbox"/> Trisomy for chromosome _____
<input type="checkbox"/> AML Panel: <i>EGR1</i> , D7S486, <i>AML1/ETO</i> , <i>MLL</i> , <i>CBFB</i>	<input type="checkbox"/> B-ALL Panel: <i>BCR/ABL1</i> , <i>MLL</i> , <i>ETV6/RUNX1</i> , 4,10,17	
<input type="checkbox"/> MDS Panel: <i>EGR1</i> , D7S486, CEP 8, <i>TP53</i> , D20S108	<input type="checkbox"/> Ph-like B-ALL Follow-up Panel: <i>ABL1</i> , <i>ABL2</i> , <i>CRLF2</i> , <i>JAK2</i> , <i>PDGFRB</i>	
<input type="checkbox"/> MM Panel: <i>CKS1B</i> , <i>ATM</i> , <i>RB1/D13S25</i> , <i>IGH</i> , <i>TP53</i>	<input type="checkbox"/> COG High Risk B-ALL Panel: <i>ABL1</i> , <i>ABL2</i> , <i>PDGFRB</i>	
<input type="checkbox"/> IGH Panel: <i>IGH/FGFR3</i> , <i>IGH/CCND1</i> , <i>IGH/MAF</i>	<input type="checkbox"/> CLL Panel: <i>ATM</i> , CEP 12, D13S319,13q34, <i>TP53</i>	

Bone marrow specimens should be sent in bone marrow transport medium. Please call 934-9555 to request a supply of this medium.

Blood specimens (5-10cc) should be shipped in a **Sodium heparinized** vacutainer. Specimens collected in lithium heparin or EDTA cannot be used for chromosome analysis. Specimens should be transported **as soon as possible** at room temperature.