

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: _____

aCGH+SNP study: _____

FISH studies (Please select from the list below)

- | | | |
|---|---|--|
| <input type="checkbox"/> t(1q25.2)- <i>ABL2</i> rearrangement | <input type="checkbox"/> t(11p15.4)-NUP98 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::PDGFRA</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 -CEP 7,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 [p16]</i> |
| <input type="checkbox"/> t(6;9)- <i>DEK::NUP214</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>RUNX1::RUNX1T1</i> fusion | <input type="checkbox"/> inv(16)- <i>CBFB::MYH11</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>KMT2A [MLL]</i> rearrangement | <input type="checkbox"/> del(Xp22.3/Yp11.3)- <i>P2RY8::CRLF2</i> fusion | <input type="checkbox"/> Trisomy for chromosome _____ |

AML Panel: *EGR1, D7S486, AML1/IETD, MLL, CBFB*

B-ALL Panel: *BCR/ABL1, MLL, ETV6/RUNX1, 4, 10, 17*

MDS Panel: *EGR1, D7S486, CEP 8, TP53, D20S108*

Ph-like B-ALL Follow-up Panel: *ABL1, ABL2, CRLF2, JAK2, PDGFRB*

MM Panel: *CKS1B, ATM, RB1/D13S25, IGH, TP53*

CLL Panel: *ATM, CEP 12, D13S319, 13q34, TP53*

IGH Panel: *IGH/FGFR3, IGH/CCND1, IGH/MAF*

Second Tier AML Panel: *MECOM, DEK::NUP214, NUP98*

Eosinophilia Panel: *LSI 4q12, PDGFRB, FGFR1, JAK2, ETV6*

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.