Collaborative Genomics Core
UNIVERSITY OF WISCONSIN–MADISON

University of Wisconsin
Collaborative Genomics Core
465 Henry Mall
Madison, WI 53706-1578
http://www.med.wisc.edu/uwcgc

Neoplasia Diagnosis (11/2015) CGC# 132

(PLEASE PRINT USING CAPITALS- FIELDS IN RED ARE REQUIRED)

(1) Patient Last Name | First Name | Middle Name

(2) Name Change- Former Last Name

(3) Patient Address

(4) City | State | Zip | County of Residence

(5) Date of Birth | (6) Age | (7) Sex ☐ Male ☐ Female

(8) Ethnicity ☐Hispanic/Latino ☐Non-Hispanic/Latino

(9) ☐Amer Indian ☐Black/African Amer ☐White ☐Asian ☐Pacific Islander ☐Other _____

(10) Chart #/ Patient ID Number | (11) Submitter Specimen ID Number

(13) Additional Report Copies Needed?
Please check this box ☐ AND
Enter the clinician's name and address on the back of this form

(14) Ordering Provider

(15) NPI # _ _ _ _ _ _ _ _ _ _

(16) Attached copies of front and back of insurance card(s)?

(17) Medicare generally does not cover routine screening tests. ABN attached? ☐ YES ☐ NO

(18) ☐ MEDICAID# _____ ☐ MEDICARE# _____
☐ PRIVATE INSURANCE# _____ ☐ Bill to Submitter

(20) Please write the letter corresponding to the appropriate ICD-10 Code to the left of the test name below (where applicable)

(A) ICD-10 Code _____ (B) ICD-10 Code _____ (C) ICD-10 Code _____ (D) ICD-10 Code _____ (E) ICD-10 Code _____

(21) Date of collection | (22) Time of collection

Specimen Source ☐ Bone Marrow ☐ Blood ☐ Tumor/Lymph Node (tissue type _____)
☐ Paraffin Section (tissue type _____) Fixative Used: _____ Fixation Time: ☐ <6 hours ☐ 6 - 48hrs ☐ >48 hrs

Reason for Referral (please provide in addition to ICD-10 code):

Check All That Apply

CHROMOSOME ANALYSIS

- ☐ 811 Chromosome Analysis, Unstimulated Blood, for Hematologic Disorders
- ☐ 812 Chromosome Analysis, Bone Marrow, for Hematologic Disorders
- ☐ 836 Chromosome Analysis, Tumor/Pleural Effusion/ Ascites fluid, for Neoplastic Disorders

MOLECULAR ANALYSIS

- ☐ 887M31 RT-PCR for BCR/ABL1 Fusion Transcript, t(9;22)(q34;q11.2)
- ☐ 887M35 RT-PCR for PML/RARA Fusion Transcript, t(15;17)(q24;q21)
- ☐ 893M51 Quantitative PCR for BCR/ABL1, t(9;22)(q34;q11.2) Major breakpoint
- ☐ 893M52 Quantitative PCR for BCR/ABL1, t(9;22)(q34;q11.2) minor breakpoint
- ☐ 840C41 Qualitative BRAF V600E [BRAF]
- ☐ 840C42 Qualitative PCR for Calreticulin [CALR]
- ☐ 840C43 EGFR Polymorphism Analysis [EGFR]
- ☐ 840C44 Quantitative PCR for JAK2 (V617F) [JAK2]
- ☐ 840C45 KRAS Polymorphism Analysis [KRAS]
- ☐ 840C46 Microsatellite Instability Analysis [MSI]
- ☐ 840C47 STR Engraftment- Unseparated Lineages, Bone Marrow [STREBM]
- ☐ 840C48 STR Engraftment- Unseparated Lineages, Blood [STREWB]
- ☐ 840C49 STR Engraftment- Separated Lineages, (CD3 (STRCD3)/, CD33 (STRCD33)) [STRSEP]
- ☐ 840C40 Cancer Mutation Panel (50 Genes) [CANPNL]

FISH ANALYSIS PARAFFIN SECTIONS (FFPE)

- ☐ 883 ERBB2 (HER2) Gene Amplification Status
- ☐ 886F5 Deletion 1p36 for glial neoplasia
- ☐ 886F6 Deletion 19q13 for glial neoplasia

* see marked tests in column at right for other FFPE tests

FISH ANALYSIS (PANELS)

- ☐ MYELOMA PANEL: includes 881F62, 870F40, 870F43, 870F47, 870F55, 882F82, 881F69, 881F70, 881F72
- ☐ MYELOMA HYPERDIPLOIDY PANEL: includes 870F45, 870F42
- ☐ MDS PANEL: includes 870F45, 870F42, 870F41, 870F44
- ☐ AML PANEL: includes 870F45, 870F42, 881F63, 882F80, 882F84
- ☐ CLL PANEL: includes 870F43, 870F51, 870F40, 870F47
- ☐ PEDIATRIC ALL PANEL: includes 881F67, 881F61, 870F48, 870F49, 870F50, 882F84

FISH ANALYSIS

- ☐ 870F40 11q22 Deletion/Duplication, ATM gene
- ☐ 870F41 Trisomy 8, D8Z2
- ☐ 870F42 Deletion 7q31, D7S522
- ☐ 870F43 Deletion 13q14, D13S319
- ☐ 870F44 Deletion 20q, D20S108
- ☐ 870F45 Deletion 5q31, EGR1
- ☐ 870F46 Deletion 9p21, P16
- ☐ 870F47 Deletion 17p13.1, TP53
- ☐ 870F48 Trisomy 4, CEP4
- ☐ 870F49 Trisomy 10, CEP10
- ☐ 870F50 Trisomy 17, D17Z1
- ☐ 870F51 Trisomy 12, D12Z3
- ☐ 870F52 X and Y sex chromosomes
- ☐ 870F55 Gain 1q21 for myeloma
- ☐ 881F60 AP12/MALT1 Fusion, t(11;18)(q21;q21)
- ☐ 881F61 BCR/ABL1 Fusion, t(9;22)(q34;q11.2)
- ☐ 881F62 CCND1/IGH Fusion, t(11;14)(q13;q32)
- ☐ 881F63 RUNX1T1/RUNX1 Fusion, t(8;21)(q22;q22)
- ☐ 881F64 IGH/BCL2 Fusion, t(14;18)(q32;q21)*
- ☐ 881F65 MYC/IGH Fusion, t(8;14)(q24;q32)*
- ☐ 881F66 PML/RARA Fusion, t(15;17)(q24;q21)
- ☐ 881F67 ETV6/RUNX1 Fusion, t(12;21)(p13;q22)
- ☐ 881F68 IGH/MALT1 Fusion, t(14;18)(q32;q21)
- ☐ 881F69 FGFR3/IGH Fusion, t(4;14)(p16;q32)
- ☐ 881F70 IGH/MAF Fusion, t(14;16)(q32;q23)
- ☐ 881F71 BCR/ABL1+ASS Tri-color Fusion, t(9;22)
- ☐ 881F72 IGH/MAFB t(14;20)(q32;q12)
- ☐ 882F80 CBFB Gene Rearrangement, 16q22
- ☐ 882F81 EWSR1 Gene Rearrangement, 22q12.2 *
- ☐ 882F82 IGH Gene Rearrangement, 14q32 *
- ☐ 882F83 MALT1 Gene Rearrangement, 18q21 *
- ☐ 882F84 KMT2A (MLL) Gene Rearrangement, 11q23
- ☐ 882F85 MYC Gene Rearrangement, 8q24 *
- ☐ 882F86 SS18 Gene Rearrangement, 18q11.2 *
- ☐ 882F87 ALK Gene Rearrangement, 2p23 *
- ☐ 882F88 BCL6 Gene Rearrangement, 3q27 *
- ☐ 882F89 FIP1L1-CHIC2-PDGFRa Gene Rearr., 4q12
- ☐ 882F90 PDGFRb Gene Rearrangement, 5q33
- ☐ 882F91 TRA/D Gene Rearrangement, 14q11.2
- ☐ 882F92 RARA Gene Rearrangement, 17q21
- ☐ 882F93 FGFR1 Gene Rearrangement, 8p11
- ☐ 882F94 DDIT3 (CHOP) Gene Rearr., 12q13 *
- ☐ 882F95 FUS Gene Rearrangement, 16p11 *
- ☐ Other _____

A sample processing fee is charged for blood and bone marrow samples that have only FISH analysis