



Center for Personalized Diagnostics
Test Requisition (v1.1 12/12)

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Web: <http://www.pennmedicine.org/cpd>

Address:

205 John Morgan Building
3620 Hamilton Walk
Hospital of The University of Pennsylvania
Philadelphia, PA 19104

Space for Patient
Label

Received Date: _____

Time _____ Tech _____

Specimen volume: _____

Tube/container: _____

Accession Number: _____

CPD Case Number: _____

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Robert D. Daber, PH.D., DABMG, Technical Director of Clinical Genomics
Jennifer Morrisette, PH.D., FACMG, Clinical Director

PATIENT INFORMATION

NAME: _____
Last First MI

Medical Record # : _____

DOB: ____ / ____ / ____

Gender: ☐ M ☐ F

☐ Inpatient ☐ Outpatient

Reported Ethnicity: _____

SAMPLE INFORMATION

WBC count: _____ %Blasts: _____

Tumor %: _____

Date of Collection: _____

Time of Collection: _____ AM / PM

Tissue Source: _____

SAMPLE TYPE

☐ Bone Marrow in EDTA (purple top)

☐ Blood in EDTA (purple top)

☐ Lymph Node in Media or Saline

☐ FFPE Block/Slides

REPORTING INFORMATION

Physician: _____

☐ HUP ☐ PPMC ☐ Other: _____

Phone: _____

Email: _____

INDICATION FOR STUDY

Diagnosis: _____

ICD9 code(s): _____

Relevant Hx: _____

TESTING

Hematologic Malignancy Sequencing Panel: Sequence analysis of 33 genes

(ASXL1, ATM, BRAF, CBL, CDKN2A, DDX3X, DNMT3A, ETV6, EZH2, FBXW7, FLT3, GNAS, IDH1, IDH2, JAK2, KIT, KLHL6, KRAS, MAPK1, MYD88, NOTCH1, NPM1, NRAS, PTEN, PTPN11, PHF6, RUNX1, SF3B1, TET2, TP53, WT1, XPO1, ZMYM3).

Solid Tumor Sequencing Panel Sequence analysis of 47 genes (ABL1, AKT1, ALK, APC, ATM, BRAF, CDH1, CSF1R, CTNNB1, EGFR, ERBB2, ERBB4, FBXW7, FGFR1, FGFR2, FGFR3, FLT3, GNA11, GNAQ, GNAS, HNF1A, HRAS, IDH1, JAK2, JAK3, KDR, KIT, KRAS, MET, MLH1, MPL, NOTCH1, NPM1, NRAS, PDGFRA, PIK3CA, PTEN, PTPN11, RB1, RET, SMAD4, SMARCB1, SMO, SRC, STK11, TP53, VHL)

The design of each panel was based on the literature at the time of development, either the full length of a gene or the mutational hot spots of a gene are targeted. For a complete listing of genomic coordinates covered in this panel please contact the laboratory.