

NOTE: PLEASE REFER TO THE BACK OF THIS REQUISITION FOR SPECIMEN HANDLING INSTRUCTIONS

MICHIGAN MEDICINE

DEPARTMENT OF PATHOLOGY

Testing / Diagnostic / Screening Requisition - Molecular Diagnostics Laboratory Requisition

RESULTS REPORTING LOCATION CODE:

MRN

BIRTHDATE

NAME

CSN

ICD-10 Code/Diagnosis:	Ordering Clinician to receive report: <input type="checkbox"/> See label above	
Collected by:		UMHS Dr. #: _____
Collected Date: ____ / ____ / ____	Collection Time: ____ : ____ am/pm	Attending Physician: (if different from above) UMHS Dr. #: _____

MOLECULAR DIAGNOSTIC LABORATORY

This request to order tests from the Molecular Diagnostics Laboratory certifies to the laboratory that (1) the ordering physician has obtained informed consent from the patient as required by applicable state or federal laws for each test ordered and (2) the ordering physician has authorization from the patient permitting the Molecular Diagnostics Laboratory to report results for each test ordered to the ordering physician.

For general information, call the Laboratory at 936-0565, M - F 8:00 - 4:30

TESTING WILL BE DELAYED OR NOT PERFORMED IF REQUISITION IS NOT COMPLETE!

SPECIMEN TYPE

SURG PATH ID#

PATIENT HISTORY/DIAGNOSIS:

HEMATOLOGY/ONCOLOGY

ACUTE MYELOID LEUKEMIA	GASTROINTESTINAL AND HEPATOBILARY	THYROID CANCER
<input type="checkbox"/> Myeloid NGS Panel	<input type="checkbox"/> Colorectal Cancer NGS Panel	<input type="checkbox"/> Solid Tumor NGS Panel
<input type="checkbox"/> TP53 Mutation in Malignancy	<input type="checkbox"/> KRAS Mutation	<input type="checkbox"/> BRAF V600E/V600K Mutation
<input type="checkbox"/> NPM1 Mutation	<input type="checkbox"/> NRAS Mutation	<input type="checkbox"/> BRAF (7q34) Rearrangement FISH
<input type="checkbox"/> FLT3 Mutation	<input type="checkbox"/> BRAF V600E/V600K Mutation	<input type="checkbox"/> TERT Promoter Mutation
<input type="checkbox"/> CEPBA Mutation	<input type="checkbox"/> Microsatellite Instability Analysis	<input type="checkbox"/> RET Mutation
<input type="checkbox"/> IDH1 and IDH2 Mutations	If MSI-H, perform <input type="checkbox"/> BRAF V600E <input type="checkbox"/> MLH1 Promoter Methylation	<input type="checkbox"/> SARCOMA
<input type="checkbox"/> KIT D816V Mutation	<input type="checkbox"/> MLH1 Promoter Methylation	<input type="checkbox"/> SYT/SSX Translocation (PCR)
<input type="checkbox"/> KIT Mutation for AML - Exons 8, 17	<input type="checkbox"/> Germiline MLH1 Promoter Methylation	<input type="checkbox"/> PAX/FOXO1 Translocation (PCR)
<input type="checkbox"/> PML/RARA t(15;17) Translocation, Quantitative	<input type="checkbox"/> UGT1A1 Promoter Genotype	<input type="checkbox"/> EWSR1/WT1 Translocation (PCR)
MYELOID NEOPLASMS	<input type="checkbox"/> HER2 (FISH)	<input type="checkbox"/> EWSR1/ATF1 Translocation (PCR)
<input type="checkbox"/> Myeloid NGS Panel	<input type="checkbox"/> FGFR2 (10q26) Rearrangement by FISH	<input type="checkbox"/> EWSR1/FL1 & EWSR1/ERG Translocation (PCR)
<input type="checkbox"/> TP53 Mutation in Malignancy	<input type="checkbox"/> Biliary Tract Malignancy (FISH)	<input type="checkbox"/> EWSR1 (22q12) Rearrangement (FISH)
<input type="checkbox"/> JAK2 V617F Mutation	<input type="checkbox"/> KIT Mutation - Exons 9,11,13,17	<input type="checkbox"/> MDM2 Amplification (FISH)
<input type="checkbox"/> JAK2 Exon 12 Mutation	If KIT is negative, perform: <input type="checkbox"/> PDGFRα <input type="checkbox"/> BRAF V600E	<input type="checkbox"/> CIC (19q13) Rearrangement (FISH)
<input type="checkbox"/> CALR Mutation	<input type="checkbox"/> PDGFRA Mutation for GIST	<input type="checkbox"/> PDGFβ (22q13) Rearrangement (FISH)
<input type="checkbox"/> MPL Mutation	<input type="checkbox"/> CENTRAL NERVOUS SYSTEM TUMORS	<input type="checkbox"/> NR4A3 (9q22-9q31) Rearrangement by (FISH)
<input type="checkbox"/> KIT D816V Mutation	<input type="checkbox"/> Cancer Cytogenomic Array-FFPE Tissue	<input type="checkbox"/> SOD3 (12q13) Rearrangement (FISH)
<input type="checkbox"/> BCR/ABL1 Analysis, Quantitative	<input type="checkbox"/> Solid Tumor NGS Panel	BREAST CANCER
<input type="checkbox"/> BCR/ABL1 Kinase Domain Mutation	<input type="checkbox"/> IDH1 and IDH2 Mutations for Glioma	<input type="checkbox"/> Solid Tumor NGS Panel
LYMPHOMA	<input type="checkbox"/> 1p/1q Deletion (FISH)	<input type="checkbox"/> HER2 (FISH)
<input type="checkbox"/> B Cell Clonality (IGH Gene Rearrangement)	<input type="checkbox"/> BRAF (7q34) Rearrangement (FISH)	<input type="checkbox"/> PIK3CA Mutation
<input type="checkbox"/> B Cell Clonality (IGK Gene Rearrangement)	<input type="checkbox"/> BRAF V600E/V600K Mutations	GENITOURINARY CANCER
<input type="checkbox"/> T Cell Clonality (TRG Gene Rearrangement)	<input type="checkbox"/> MGMT Promoter Methylation	<input type="checkbox"/> (1p2p) assessment by Cancer Cytogenomic Array-FFPE Tissue
<input type="checkbox"/> T Cell Clonality (TRB Gene Rearrangement)	<input type="checkbox"/> TERT Promoter Mutation	<input type="checkbox"/> TFE3-FISH (Cytogenetics)
<input type="checkbox"/> TP53 Mutation in Malignancy	<input type="checkbox"/> LUNG CANCER	<input type="checkbox"/> TFE6-FISH (Cytogenetics)
<input type="checkbox"/> IGH/BCL2 t(14;18) Translocation (PCR)	<input type="checkbox"/> Lung Cancer NGS Panel	<input type="checkbox"/> ERG-FISH (Cytogenetics)
<input type="checkbox"/> IGH/BCL2 t(14;18) Translocation (FISH)	<input type="checkbox"/> EGFR Mutation	<input type="checkbox"/> Solid Tumor NGS Panel
<input type="checkbox"/> BCL6 (3q27) Rearrangement (FISH)	<input type="checkbox"/> BRAF V600E/V600K Mutations	<input type="checkbox"/> FGFR Translocation/Mutation
<input type="checkbox"/> MYC (8q24) Rearrangement (FISH)	<input type="checkbox"/> KRAS Mutation	<input type="checkbox"/> TERT Promoter Mutation
If MYC is positive, perform: <input type="checkbox"/> IGH/BCL2 (FISH) <input type="checkbox"/> BCL6 (FISH)	<input type="checkbox"/> ALK Rearrangement for NSCLC (FISH)	<input type="checkbox"/> UroVysionTM (FISH)
<input type="checkbox"/> MALT1 (18q21) Rearrangement (FISH)	<input type="checkbox"/> ROS1 (6q22) Rearrangement (FISH)	MISCELLANEOUS
<input type="checkbox"/> MYD88 (L265P) Mutation	<input type="checkbox"/> RET (10q11) Rearrangement (FISH)	<input type="checkbox"/> Cancer Cytogenomic Array-FFPE Tissue
<input type="checkbox"/> BRAF V600E/V600K Mutations	<input type="checkbox"/> MET Amplification by FISH	<input type="checkbox"/> Solid Tumor NGS Panel
	<input type="checkbox"/> Mesothelioma FISH	<input type="checkbox"/> HER2 (FISH)
	MELANOMA	<input type="checkbox"/> Mesothelioma FISH
	<input type="checkbox"/> Cancer Cytogenomic Array-FFPE Tissue	<input type="checkbox"/> Microsatellite Instability
	<input type="checkbox"/> Melanoma NGS Panel	<input type="checkbox"/> If MSI-H perform MLH1 Promoter Methylation
	<input type="checkbox"/> BRAF V600E/V600K Mutations	<input type="checkbox"/> MLH1 Promoter Methylation
	<input type="checkbox"/> KIT Mutation for Melanoma - Exons 11,13,17	<input type="checkbox"/> Germiline MLH1 Promoter Methylation
	<input type="checkbox"/> NRAS Mutation	<input type="checkbox"/> Specimen Identity (discuss with Taulbee/Kunju/Pantanowitz)
	<input type="checkbox"/> BRAF (7q34) Rearrangement (FISH)	
	<input type="checkbox"/> TERT Promoter Mutation	

ENDOMETRIAL CANCER

- POLE Mutation
- Microsatellite Instability
- If MSI-H perform MLH1 Promoter Methylation
MLH1 Promoter Methylation

BONE MARROW TRANSPLANT ENGRAFTMENT ASSESSMENT

S,T	<input type="checkbox"/> Pre-BMT RECIPIENT, Engraftment Analysis
S,T	<input type="checkbox"/> Pre-BMT DONOR, Engraftment Analysis
S,T	DONOR FOR: Name: _____ MRN: _____
	<input type="checkbox"/> Post-BMT Engraftment Analysis (Pre-BMT MUST have been previously performed)
	Non-myeloablative transplant? <input type="checkbox"/> Yes <input type="checkbox"/> No
	Fractionation? <input type="checkbox"/> Yes <input type="checkbox"/> No
	Days post-transplant: _____