

# New York State Hematopathology Requisition

**Client Information** 

Required Information           Account #:	Account Name:	
City, ST, ZIP:		
Phone:	Fax:	
Requisition Completed by:	Date:	
Ordering Physician:	NPI #:	
Treating Oncologist/Physician:	NPI #:	
By completing this section, the undersigned certifier such test(s) are medically necessary for the care/tre	es that he/she is licensed to order the test(s) listed below and that eatment of this patient.	
Authorized Signature:	Date:	

# **Billing Information**

Required: Please include face sheet and front/back of patient's insurance card.					
Patient Status (Must Choose 1): Hospital Patient (in) Hospital Patient (out) Non-Hospital Patient					
Bill to:  Client Bill	Insurance	Medicare	Medicaid	Patient/Self-Pay	
Split Billing - Client (TC) and Insurance (PC) OP Molecular to MCR, all other testing to Client					
Bill charges to other Hospital/Facility:					
Prior Authorization #			See neogenomics	.com/billing for more info.	

#### **Clinical Information**

I: Please attach patient's pathology report (required), clinical history, and other applicable report(s) □ ICD 10 (Diagnosis) Code/Narrative (Required):

Reason for Referral

New Diagnosis □ Relapse/Refractory □ Monitoring 

Bone Marrow Transplant (required information for Oncology Cytogenetics) None Autologous Allogeneic Sex Mismatch

#### Consultation

COMPASS® Comprehensive evaluation including morphology Blood and/or Bone Marrow Paraffin block for Morphology to follow A NeoGenomics pathologist will select medically necessary tests COMPASS® Select: (Without morphology) (with any exceptions noted or marked by the client) to provide Blood and/or Bone Marrow comprehensive analysis and professional interpretation for the Morphology performed by client (Morphology performed by client (Morphology report required. Please fax to avoid testing delays.) Lymph Node/Tissue for Lymphoma\* Difference in a power of the power of th materials submitted Please attach CBC for Blood and Bone Marrow (required) Split fresh specimens to RPMI and formalin Paraffin block for Morphology to follow Do not add NGS Profile without prior approval Flow Cytometry G - Global T - Tech-Only Follow-Up/Add-On panels are available in conjunction with, or after Attach CBC with all flow requests on blood or bone marrow. a Main Panel result has been reported by NeoGenomics or client. **Diagnostic/Prognostic Panels** Add-On Tubes Follow-Up Panels G T G T Add-on Tables G AML B-ALL CLL/Mantle CLL/Mantle Hairy Cell Hairy Cell Plasma Ce T-ALL TRBC1/L RBC1/L Ď Ò Standard L/L Panel (24 Markers) G Т Extended L/L Panel (31 Markers) High Sensitivity PNH Ă AML High Sensitivity First
 T&B Tissue Panel
 TRBC1/T-Cell Lymphoma Companion B-ALL CLL/Mantle Cell Companion N/A 🗖 B-ALL Hairy Cell Erythroid-Mega D Plasma Cell T-ALL □ NA B-ALL MRD Panel (Bone Marrow) Plasma Cell T-ALL NA B-ALL MRD Panel (Peripheral Blood) NA CLL MRD Panel TRBC1/LGL NA MM MRD Panel Specimen Hold Option: 
Refrigerate and Hold Tech-Only Opt Out Option: To avoid delay in patient care and as medically necessary for an individual patient, additional markers will be added by the flow lab when abnormal populations are detected. Please refer to NeoGenomics additional markets will be addee by the normal when an information populations are detected. These refer to redocenomic Flow Cytometry Guidelines for addition on tech-only add-on medical necessity criteria. Tech-only clients may instruct NeoGenomics to not follow this stated criteria by checking this box. Cytogenetics Oncology Chromosome Analysis □ Reflex to FISH if cytogenetics is normal (reflex FISH panel must be selected) Reflex to FISH if cytogenetics is incomplete (<20 metaphases) G G T MDS Standard FISH G G T MDS Extended FISH Follow-up Constitutional Chromosome Analysis (only if recommended by Oncology Chromosome Analysis) □ Other: Specimen Hold Option: 
Culture and Hold (liquid samples & lymph nodes; n/a for solid tissues) **Molecular Genetics** □ IgVH Mutation Analysis\* □ JAK2 V617F - Qualitative\* MPL Mutation Analysis ☐ ABL1 Kinase Domain\* MPN JAK2 V617F with If negative, reflex to JAK2 Exon 12-13
 If negative, reflex to CALR
 If negative, reflex to MPL
 JAK2 V617F - Quantitative B-Cell Gene Rearrangement Sequential Reflex to JAK2 BCR-ABL1 Standard p210, p190\* 12-13, CALR and MPL<sup>†</sup> MYD88 Mutation Analysis BRAF Mutation Analysis Calreticulin (CALR)† NPM1 Mutation Analysis FLT3 Mutation Analysis JAK2 Exon 12-13\* PML-RARA, t(15;17) □ IDH1/IDH2 by PCR KIT (c-KIT) Mutation Analysis T-Cell Receptor Beta

	Patient Information
	Last Name: 🗆 Male 🛛 Female
	First Name: M.I Medical Record #:
	Date of Birth: mm / dd / yyyy Other Pt ID/Acct #:
	Client represents it has obtained informed consent from patient to perform the services described herein.
j	Specimen Information
	Specimen ID: Block ID: Fixative/Preservative:
	Collection Date: mm / dd / yyyy Collection Time: DAM DPM
	Retrieved Date: mm / dd / yyyy
	Hospital Discharge Date: mm / dd / yyyy Body Site:
	Primary      Metastasis – If Metastasis, list Primary:
	Bone Marrow [must provide CBC and Path Report]:
	Green Top(s) Purple Top(s) Core Biopsy Clot
	Peripheral Blood: Green Top(s) Purple Top(s) Other      Fresh Tissue (Media Type required):
	Fluid: CSF Pleural Other
	□ Smears: Air Dried Fixed Stained (type of stain)     □ Slides # Unstained Stained □ H&E
	□ Paraffin Block(s) #:
	Choose best block (for global molecular/NGS testing only)
	Submit ≤4 blocks. Blocks will be combined for molecular testing when necessary. For all other testing, specify which block to use for each if sending multiple blocks. See back for details.
	Comments:
_	
	FISH G - Global T - Tech-Only (Reflex options are available with Global cases only.)
	HemeFISH® Hematologic FISH Panels G T G T
	Image: Construction of the state of the
	□ □ ALL - Pediatric □ Reflex to BCL6/MYC, IGK/MYC,
	□ □ ∧AM Standard □ N/A High-Grade B-Cell Lymphoma Reflex
	Alle Stational Version Sq5, 7q/-7, DEK/NUP214     Ic(6.9), p53 (17p13.1)/NF1 (17q11) and     MDS Extended     MDS Extended
	NUP98 (11p15) if negative III III MDS Standard III Reflex to ETV6 (12p13), MLL (11q23)
1	AML Favorable-Risk     and +19 if negative     AML Non-Favorable Risk     MPN
	CLL (fresh specimens only)
	Plasma Cell Myeloma Panels - Plasma Cell Enrichment will be performed on all bone marrow samples having G T plasma cell FISH tests.
	<ul> <li>Plasma Cell Myeloma</li> <li>Do not reflex to IqH Complex (applies to global only; tech-only will not reflex)</li> </ul>
	Plasma Cell Myeloma IgH Complex
	Plasma Cell Myeloma Prognostic Panel Individual Probes
	G T
	Image: Interpretation in NHL       Image: CDKN2A (p16) Deletion for ALL       Image: MYC/lgH/CEN8 t(8;14)         Image: Interpretation in NHL       Image: CDKN2A (p16) Deletion for ALL       Image: MYC/lgH/CEN8 t(8;14)         Image: Interpretation in NHL       Image: CDKN2A (p16) Deletion for ALL       Image: MYC/lgH/CEN8 t(8;14)         Image: Interpretation in NHL       Image: CDKN2A (p16) Deletion for ALL       Image: MYC/lgH/CEN8 t(8;14)         Image: Interpretation in NHL       Image: CDKN2A (p16) Deletion for ALL       Image: MYC/lgH/CEN8 t(8;14)         Image: Interpretation in NHL       Image: CDKN2A (p16) Deletion for ALL       Image: MYC/lgH/CEN8 t(8;14)         Image: Interpretation in NHL       Image: CDKN2A (p16) Deletion for ALL       Image: MYC/lgH/CEN8 t(8;14)         Image: Interpretation in NHL       Image: CDKN2A (p16) Deletion for ALL       Image: MYC/lgH/CEN8 t(8;14)         Image: Interpretation in NHL       Image: CDKN2A (p16) Deletion for ALL       Image: MYC/lgH/CEN8 t(8;14)         Image: Interpretation in NHL       Image: CDKN2A (p16) Deletion for ALL       Image: MYC/lgH/CEN8 t(8;14)         Image: Interpretation in NHL       Image: CDKN2A (p16) Deletion for ALL       Image: MYC/lgH/CEN8 t(8;14)         Image: Imag
	□ □ ALK for Lymphoma □ □ IGK/MYC t(2;8) □ □ TCL1 (14q32.1)t
	□ BCR/ABL1/ASS1 t(9:22) □ □ IGL/MYC t(8:22) □ □ PML/RARA t(15:17)
	□ BCL6/MYC t(3;8) □ JAK2 (9p24.1) □ Other
-	Specimen Hold Option: Direct Harvest and Hold Plasma Cell Enrichment and Hold
	NeoTYPE® & Neo Comprehensive® Cancer Profiles G - Global T - Tech-Only FISH G T G T
	ALL Profile     N/A AML Prognostic Profile     N/A AML Prognostic Profile     N/A ML Prognostic Profile     N/A ML Prognostic Profile     N/A ML Prognostic Profile
	N/A AML Prognostic Profile + FLT3 by PCR* IN/A Lymphoid Disorders Profile
	CLL Profile     Add IgVH Mutation Analysis
	<ul> <li>N/A Neo Comprehensive - Heme Cancers</li> <li>N/A Neo Comprehensive - Heme Cancers + FLT3 by PCR*</li> </ul>
	N/A Neo Comprehensive - Myeloid Disorders
-1	Specimen Hold Option: D Extract & Hold - TNA (all tests are TNA-based)

\*Note: The FLT3 Mutation Analysis test is available as client-bill only when ordered with NeoTYPE and Neo Comprehensive. The Molecular case reports separately from the NeoTYPE or Neo Comprehensive Profile (which also includes FLT3 gene by NGS) for the purpose of prompt therapy selection in patients with a new diagnosis of AML.

#### WRITE-IN

Any testing ordered in the Write-In section will be sent to a reference laboratory that has NY licensure to perform requested testing. NeoGenomics will facilitate logistics on behalf of the client. NeoGenomics testing will be prioritized. □ Other:

#### FlexREPORT<sup>®</sup>

T-Cell Receptor Gamma

FlexREPORT: Please add summary report option to this case

Test is RNA-based. † Test is both DNA- and RNA-based

Specimen Hold Options: D Extract & Hold - DNA DExtract & Hold - RNA

# **Specimen Requirements**

Refrigerate specimen if not shipping immediately and use cool pack during transport. Please call the Client Services team with any questions regarding specimen requirements or shipping instructions at 866.776.5907 option 3. Please refer to the website for specific details on each specimen.

### Additional Billing Information

Any organization referring specimens for testing services pursuant to this Requisition Form ("Client") expressly agrees to the following terms and conditions.

- 1. Binding Service Order. This Requisition Form is a contractually binding order for the services ordered hereunder ("Services") and Client agrees that it is financially responsible for all tests billable to Client hereunder.
- 2. Third Party Billing by NeoGenomics and Right to Bill Client. Client agrees to accurately indicate on the front of the Requisition Form that either Client should be billed (e.g., Client receives reimbursement pursuant to a non-fee-for-service basis, including, but not limited to, a capitated, diagnostic related group ("DRG"), per diem, all-inclusive, or other such bundled or consolidated billing arrangement) or NeoGenomics should bill the applicable federal, state or commercial health insurer or other third party payer (collectively, "Payers") for all Services ordered pursuant to this Requisition Form. For all such Services billable to Payers, Client agrees to provide all billing information necessary for NeoGenomics to bill such payer. In the event NeoGenomics: (i) does not receive the billing information required for it to bill any Payers within ten days of the date that any Services are reported by NeoGenomics; (ii) the Services were performed for patients who have no Payer coverage arrangements; or (iii) the Payer identified by Client denies financial responsibility for the Services and indicates that Client is financially responsible, NeoGenomics shall have the right to bill such Services to Client.

### **Additional Specimen Information**

If submitting multiple blocks, clients must indicate either "Choose best block (global molecular/NGS testing only)", or assign the selection of blocks to individual tests. If multiple blocks are sent without a selection, they will be held until clarification is provided. Please call Client Services team with any questions regarding specimen information.

### **Specimen Hold Option Descriptions**

To preserve the integrity of samples and avoid unnecessary testing, NeoGenomics Laboratories offers the option of processing samples to maintain specimen integrity for extended periods, without a test order. Any hold order will result in billed charges to the ordering client if testing is not ordered/performed. Specimen Hold Options include:

FISH: <u>Direct Harvest and Hold</u>: FISH specimens will be minimally processed and directly harvested while the cells are still viable. Analysis is not performed until the client test order is received. Processed samples will be retained for 28 days. <u>Plasma Cell Enrichment and Hold</u>: Plasma cells will be isolated for bone marrow specimens. Sample should be received at NeoGenomics Laboratories within 72 hours of collection. Analysis is not performed until the client test order is received. Processed samples will be retained for 28 days.

Flow Cytometry: Refrigerate and Hold: Flow cytometry samples will be refrigerated and retained for 28 days, however, optimal stability is within 72 hours of collection.

Molecular Testing: Extract Nucleic Acid and Hold: Nucleic acid (DNA or RNA or TNA) will be isolated from viable cells and stored in a freezer. Use this option when it is known which test(s) may be added. Analysis is not performed until the client test order is received. Processed samples will be retained for 28 days.

### **Test Descriptions**

Please see complete test descriptions and all available tests at our website, www.neogenomics.com/test-menu.

# **Test Notations**

#### Specimen Usage

NeoGenomics makes every effort to preserve and not exhaust tissue, but in small and thin specimens, there is a possibility of exhausting the specimen in order to ensure adequate material and reliable results.

### FlexREPORT<sup>®</sup>

FlexREPORT can be ordered on any global or tech-only testing referred to NeoGenomics. This report template can be used to import data and images collected from testing performed outside of NeoGenomics, and incorporated into a one page summary report. Client logo and contact information will be in the header of the FlexREPORT.

### FISH

Plasma cell myeloma FISH panels: Plasma cell enrichment will be performed on bone marrow samples having plasma cell FISH. Sample should be received at NeoGenomics Laboratories within 72 hours of collection.