

MOLECULAR GENETICS TEST REQUISITION

All Information Must Be Completed Before Sample Can Be Processed

PATIENT INFORMATION

Patient Name: _____, _____, _____
Last First MI

Address: _____

Home Phone: _____

MR# _____ Date of Birth _____ / _____ / _____

Gender: Male Female

ETHNIC/RACIAL BACKGROUND (Choose All)

- European American (White) African-American (Black)
- Native American or Alaskan Asian-American
- Pacific Islander Ashkenazi Jewish ancestry
- Latino-Hispanic _____
(specify country/region of origin)
- Other _____
(specify country/region of origin)

BILLING INFORMATION (Choose ONE method of payment)

PATIENT BILLING

Check Enclosed Money Order Credit Card (Visa, MC, Amex., Disc.)

Credit Card Number: _____

Card Holder Name: _____

Expiration Date: _____

Signature: _____

REFERRING INSTITUTION

Institution: _____

Address: _____

City/State/Zip: _____

Accounts Payable Contact Name: _____

Phone: _____

Fax: _____

Email: _____

COMMERCIAL INSURANCE / POLICY HOLDER INFORMATION*

Name: _____

Gender: _____ Date of Birth _____ / _____ / _____

Authorization Number: _____

Insurance Name: _____

Insurance Address: _____

City/State/Zip: _____

Insurance ID Number: _____

Group Number: _____

Insurance Phone Number: _____

* PLEASE NOTE:

- Insurance can only be billed if requested at the time of service
- Please call 866-450-4198 with any billing questions.

SAMPLE/SPECIMEN INFORMATION

SPECIMEN TYPE: Amniotic fluid Blood Cytobrushes Saliva

Cord blood CVS Bone marrow Other _____

Tissue (specify): _____

Specimen Date: _____ / _____ / _____ Time: _____

Specimen Amount: _____

Each test requires 3 mL of whole blood in EDTA tube. Please call before sending alternate tissue samples, and for free cytobrush or saliva collection kits.

DRAWN BY: _____

*Phlebotomist must initial tube of specimen to confirm sample identity

Has patient received a bone marrow transplant? Yes No

If yes, date of bone marrow transplant _____

Percent engraftment _____

REFERRING PHYSICIAN

Physician Name (print): _____

Address: _____

Phone: (_____) _____ Fax: (_____) _____

Email: _____

Genetic Counselor/Lab Contact Name: _____

Phone: (_____) _____ Fax: (_____) _____

Email: _____

_____ Date: ____ / ____ / ____

Referring Physician Signature (REQUIRED)

Patient signed completed ABN

Medical Necessity Regulations: At the government's request, the Molecular Genetics Laboratories would like to remind all physicians that when ordering tests that will be paid under federal health care programs, including Medicare and Medicaid programs, that these programs will pay only for those tests the relevant program deems to be (1) included as covered services, (2) reasonable, (3) medically necessary for the treatment and diagnosis of the patient, and (4) not for screening purposes.

INDICATIONS/DIAGNOSIS/ICD-9 CODE

Reason for Testing:

- Diagnosis in symptomatic patient
- Asymptomatic infant with abnormal newborn screen
- Carrier (Heterozygote) testing
- Presymptomatic diagnosis of at-risk sibling
- Prenatal testing (by previous arrangement only)
- Family history of disease

SAMPLE/SPECIMEN INFORMATION
Has patient received a bone marrow transplant?

- Yes
- No

If yes, date of bone marrow transplant _____

Percent engraftment _____

PEDIGREE OR FAMILY HISTORY

 Parental Consanguinity Y N

TEST(S) REQUESTED
Custom Gene Testing

Sanger sequencing of any clinically relevant gene is available by prior arrangement. Custom single gene requisition available at: www.cincinnatichildrens.com/customtesting

Deletion/Duplication Assay

Custom deletion/duplication analysis available for many genes. List of available genes and test requisition available at: www.cincinnatichildrens.com/deldup

Whole Exome Sequencing

Exome sequencing test requisition available at: www.cincinnatichildrens.org/exome

Epidermolysis Bullosa

- EBSeq Epidermolysis Bullosa Panel
(*CD151, CDSN, CHST8, COL17A1, COL7A1, DSP, DST, EXPH5, FERMT1 (KIND1), ITGA3, ITGA6, ITGB4, JUP, KRT5, KRT14, LAMA3, LAMB3, LAMC2, PKP1, PLEC1, TGM5*)

Fatty Acid Oxidation Disorders

- ACADM* (K329E) genotyping only
- ACADM* (K329E) genotyping, with reflex to full sequencing if indicated
- ACADM* full gene sequence analysis
- ACADM* full gene sequence analysis, with reflex to MetaboSeq® if indicated
- ACADVL* full gene sequence analysis
- ACADVL* full gene sequence analysis, with reflex to MetaboSeq® if indicated
- CPT2* full gene sequence analysis
- CPT2* full gene sequence analysis, with reflex to MetaboSeq® if indicated
- HADHA* (E510Q) genotype analysis
- HADHA* (E510Q) genotype analysis, with reflex to MetaboSeq® if indicated
- MetaboSeq® (19 gene next-generation sequencing panel)
- SLC22A5* full gene sequence analysis
- SLC22A5* full gene sequence analysis, with reflex to MetaboSeq® if indicated

Genetic Pharmacology Services

- Psychiatry Pharmacogenetics Expanded Panel
- TPMT Genotype Analysis
- Opioid CYP2D6 Pharmacogenetics Panel
- CYP2C19
- CYP2C9
- CYP2D6
- CYP2D6/CYP2C19
- Individual Drug
Specify drug: _____
(See www.cincinnatichildrens.org/service/g/genetic-pharmacology/default/ for list)

Hearing Loss Testing

(Please provide audiogram and MRI/CT report, if available).

Hearing Loss Panels

- Hearing Loss Panel Tier I (*GJB2, GJB6* and 8 mitochondrial mutations)
- OtoSeq® Hearing Loss Panel (next-generation sequencing of 23 genes)
- Hearing Loss Panel Tier I with reflex to OtoSeq® Hearing Loss Panel, if indicated
- Branchiootorenal Spectrum Disorder (BOR/BOS) Panel (sequencing of *EYA1, SIX1, SIX5*)
- Branchiootorenal Spectrum Disorder (BOR/BOS) Panel with reflex to OtoSeq® reanalysis, if indicated
- Hearing loss mtDNA panel (961, 1555, 1494, 3242, 3271, 7445, 7511, 8344)
- Pendred Syndrome Panel (*SLC26A4, FOXI1, KCNJ10*)
- Pendred Syndrome Panel with reflex to OtoSeq® reanalysis, if indicated
- Usher Syndrome Panel (sequencing of *CDH23, CLRN1, GPR98, MYO7A, PCDH15, USH1C, UHS1G, USH2A, WHRN*)
- Usher Syndrome Panel with reflex to OtoSeq® reanalysis, if indicated

Single Gene Tests

- CDH23* (*USH1D* and *DFNB12*)
- EYA1* (Branchiootorenal spectrum disorder type 1)
- GJB2* (connexin 26)
- GJB6* (connexin 30) deletion analysis by MLPA
- MYO7A* (*USH1B, DFNB2, DFNB11*)
- OTOF* (*AUNB1, DFNB9*)
- SLC26A4* (Pendred syndrome, *DFNB4*)

TEST(S) REQUESTED CONTINUED
Hematology Testing
Hemoglobin Testing

- Comprehensive globin gene analysis
- HBA1 and HBA2 (α -globin) sequence analysis
- HBA1 and HBA2 (α -globin) deletion analysis by MLPA
- HBB (β -globin) sequence analysis
- HBB (β -globin) deletion analysis by MLPA

Hemolytic Anemia

- Hemolytic Anemia Panel (includes sequence analysis of AK1, ALDOA, ANK1, CDAN1, EPB41, EPB42, G6PD, GCLC, GPI, GPX1, GSR, GSS, HK1, KLF1, NT5C3A, PFKM, PGK1, PIEZO1, PKLR, RHAG, SEC23B, SLC2A1 (GLUT1), SLC4A1, SPTA1, SPTB, TPI1, XK)
- Congenital Dyserythropoietic Anemia (CDA) Panel (includes sequence analysis of CDAN1, KLF1, SEC23B)
- CDA Panel with reflex to Hemolytic Anemia Panel reanalysis, if indicated
- RBC Membrane Disorders Panel (includes sequence analysis of ANK1, EPB41, EPB42, PIEZO1, RHAG, SLC2A1, SLC4A1, SPTA1, SPTB, XK)
- RBC Membrane Disorders Panel with reflex to Hemolytic Anemia Panel reanalysis, if indicated
- RBC Enzymopathy Panel (includes sequence analysis of AK1, ALDOA, G6PD, GCLC, GPI, GPX1, GSR, GSS, HK1, NT5C3A, PFKM, PGK1, PKLR, TPI1)
- RBC Enzymopathy Panel with reflex to Hemolytic Anemia Panel reanalysis, if indicated

Thrombophilia Testing

- Factor V (Leiden)
- MTHFR (677 C>T and 1298 A>C) genotype
- Prothrombin (Factor II) G20210A genotype
- Thrombophilic polymorphism panel (MTHFR 677C>T and 1298 A>C, Factor V- Leiden, Factor II-Prothrombin G20210A, PAI-1 4G/5G genotypes)

Thrombotic Microangiopathy (aHUS and TTP)

- aHUS Genetic Susceptibility Panel (includes sequence analysis of C3, CFB, CFH, CFHR1, CFHR3, CFHR5, CFI, DGKE, MCP, THBD) and MLPA analysis for CFHR1/CFHR3 deletion)
- ADAMTS13 gene sequencing

Hereditary Immunodeficiency Testing
Autoimmune Lymphoproliferative Syndrome

- Autoimmune Lymphoproliferative Syndrome (ALPS) Panel by next generation sequencing (NGS) (CASP8, CASP10, FADD, FAS, FASLG, ITK, MAGT1, NRAS, KRAS)
- FAS (TNFRSF6)
- FASLG (TNFSF6)
- CASP10

- Somatic FAS sequence analysis of sorted double-negative T cell (DNTEC) You MUST call 513-636-2731 and 513-636-4685 in advance for specimen requirements and to schedule this test)

 Bone Marrow Failure Syndromes Panel by next-generation sequencing (NGS)

(AP3B1, BRCA2 (FANCD1), BRIP1 (FANCJ), CSF3R, CXCR4, DKC1, ELANE, ERCC4 (FANCC), FANCA, FANCB, FANCC, FANCD2, FANCE, FANCF, FANCG, FANCI, FANCL, FANCM, G6PC3, GATA1, GATA2, GFI1, HAX1, LAMTOR2 (ROBLD3), LYST, MPL, NHP2 (NOLA2), NOP10 (NOLA3), PALB2 (FANCN), RAB27A, RAC2, RAD51C (FANCO), RBM8A, RMRP, RPL5, RPL11, RPL15, RPL26, RPL35A, RPS7, RPS10, RPS17, RPS19, RPS24, RPS26, RTEL1, SBDS, SLC37A4, SLX4 (FANCP), SRP72, TAZ, TERC (hTR), TERT, TINF2, USB1, VPS13B, VPS45, WAS, WRAP53 (TCAB1, WDR79))

 Chromosome Breakage Syndrome Panel

(ATM, BLM, LIG4, NBN, NHEJ1)

 Diamond-Blackfan anemia panel

(GATA1, RPL5, RPL11, RPL15, RPL26, RPL35A, RPS7, RPS10, RPS17, RPS19, RPS24, RPS26)

 Dyskeratosis congenita Panel

(DKC1, NHP2 (NOLA2), NOP10 (NOLA3), RTEL1, TERC (hTR), TERT, TINF2, WRAP53 (WDR79, TCAB1))

Familial hemophagocytic lymphohistiocytosis

- Hemophagocytic Lymphohistiocytosis (HLH) Panel (AP3B1, BLOC1S6, CD27, ITK, LYST, MAGT1, PRF1, RAB27A, SH2D1A, SLC7A7, STX11, STXBP2, UNC13D (MUNC13-4), XIAP (BIRC4))

If inadequate DNA is present, we will prioritize testing according to our FHL testing algorithm, unless you indicate a different order of prioritization below.

- ___ UNC13D (MUNC13-4)
- ___ PRF1
- ___ RAB27A (Griscelli syndrome)
- ___ STXBP2
- ___ STX11

Fanconi anemia

- Fanconi Anemia Panel by next-generation sequencing (NGS) (BRCA2 (FANCD1), BRIP1 (FANCJ), ERCC4 (FANCC), FANCA, FANCB, FANCC, FANCD2, FANCE, FANCF, FANCG, FANCI, FANCL, FANCM, PALB2 (FANCN), RAD51C (FANCO), SLX4 (FANCP))
- FANCA
- FANCC
- FANCC IVS4+4 A>T (common Ashkenazi mutation) only
- FANCG

Lymphoproliferative disorders (Including EBV-Related)

- SH2D1A
- XIAP (BIRC4)
- ITK
- MAGT1

Severe Combined Immunodeficiencies

- Severe Combined Immunodeficiency Panel by next-generation sequencing (NGS) (ADA, CD3D, CD3E, DCLRE1C, FOXP1 (WHN), IL2RG, IL7R, JAK3, LIG4, NHEJ1, ORAI1, PNP, PTPRC, RAG1, RAG2, RMRP, STAT5B, STIM1, TBX1, ZAP70)
- X-linked severe combined immunodeficiency (IL2RG)

TEST(S) REQUESTED CONTINUED
Severe congenital neutropenia

- Inherited Neutropenia Panel by next-generation sequencing (NGS)
(*AP3B1, C16orf57, CSF3R, CXCR4, ELANE (ELA2), G6PC3, GATA1, GATA2, GFI1, HAX1, LAMTOR2 (ROBLD3), LYST, RAB27A, RAC2, SBDS, SLC37A4, TAZ, VPS13B, VPS45, WAS, WIPF1*)
- ELANE (ELA2)*
- HAX1*
- WAS* (males only)

Other Primary Immunodeficiencies

- IPEX syndrome (*FOXP3*)
- Shwachman Diamond syndrome (*SBDS*)
- Wiskott-Aldrich syndrome (*WAS*)
- X-linked hyper IgM immunodeficiency (*CD40LG*)

Hereditary Liver Disease Testing

- JaundiceChip re-sequencing array (*ABCB4, ABCB11, ATP8B1, JAG1, SERPINA1*)

Single Gene Tests

- ABCB4* (PFIC3/MDR3 deficiency)
- ABCB11* (PFIC2/BSEP deficiency)
- ATP8B1* (PFIC1/FIC1 deficiency)
- BAAT* (Familial Hypercholanemia)
- JAG1* (Alagille syndrome)
- SERPINA1* genotyping assay for PI*Z and PI*S alleles
- SERPINA1* (alpha-1-antitrypsin deficiency)
- TJP2* (PFIC4/Familial Hypercholanemia)

Intellectual Disabilities Testing
Creatine deficiency syndromes

- SLC6A8* (creatine transporter defect)
- GATM* (AGAT deficiency)
- GAMT*
- Fragile X syndrome (*FMR1* – By PCR and Southern)
- MECP2* (Rett syndrome)
- Prader-Willi/Angelman syndrome (by methylation specific PCR)

Lysosomal Storage Disease Testing
Cystinosis

- CTNS* full gene sequencing only
- CTNS* 57 kb deletion analysis by MLPA only
- CTNS* 57 kb deletion analysis with reflex to full gene sequencing

Fabry Disease

- GLA* full gene sequencing

Gaucher Disease

- GBA* full gene sequencing
- Ashkenazi Panel

(TESTS ONLY: N370S, L444P, 84GG, IVS2+1G>A)

MPS II - Hunter syndrome

- IDS* full gene sequencing

Pompe Disease

- GAA* full gene sequencing for Infantile Pompe (by prior arrangement only)
- GAA* full gene sequencing

Mitochondrial Disorders
Mitochondrial DNA

- Whole mitochondrial genome (mtDNA) sequencing **and** deletion/duplication analysis by next generation sequencing (NGS)
 - Whole mitochondrial genome (mtDNA) sequencing by NGS
 - Reflex to deletion/duplication analysis by NGS, if warranted
- mtDNA panels**
- Common mutations mtDNA panel (mtDNA 1555, 3243, 3271, 3460, 8344, 8993, 11778, 14459, 14484)
 - Common mutations mtDNA panel with reflex to mtDNA whole genome sequencing
 - Neuromuscular disorders mtDNA panel (MELAS/MERRF: mtDNA 3243, 3271, 8344)
 - Neuromuscular disorders mtDNA panel with reflex to mtDNA whole genome sequencing
 - Leber Hereditary Optic Neuropathy mtDNA panel (mtDNA 3460, 11788, 14459, 14484)
 - Leber Hereditary Optic Neuropathy mtDNA panel with reflex to mtDNA whole genome sequencing
 - Hearing loss mtDNA panel (mtDNA 1555, 1494, 961, 7445, 7511)

POLG-related disorders: AD-PEO, SANDO, MIRAS

- POLG* full gene sequencing

IBMPFD

- VCP* full gene sequencing

Oncology Molecular Testing

Samples must be received within 24 hours of drawing.

- BCR/ABL t(9;22) Qualitative
- BCR/ABL t(9;22) Quantitative (p210)
- BCR/ABL t(9;22) Quantitative (p190)
- JAK2*
- PML-RAR α t(15;17)—(qualitative only)
- NPM1*

Targeted (family specific) mutation analysis

Gene of interest _____

Proband's name _____

Proband's DOB _____

Proband's mutation _____

Relationship to proband _____

Please call 513-636-4474 to discuss any family-specific mutation analysis with genetic counselor prior to shipment.

Whole Exome Sequencing

Exome sequencing test requisition available at www.cincinnatichildrens.org/exome

- Bone marrow engraftment by STR - same sex donor and recipient
Please specify:
 - Donor
 - Host (pre-transplant)
 - Host (post-transplant)
- Sorted marrow engraftment (WBC sub-populations)
You must call 513-803-2567 to schedule this test prior to sample submission.
Please specify: STR FISH
- Hereditary hemochromatosis (HFE- C282Y and H63D)
- DNA/RNA processing and storage (Call (513) 636-4474 to arrange)
- Maternal cell contamination
(by STR; required for all prenatal diagnostic testing)
- Twin zygosity (by STR)