

Mitochondrial and Metabolic Test Requisition Form

Patient Information	Sample Information			
First name Gender	Medical record # Specimen ID Date sample obtained (mm/dd/yy) □ Blood in EDTA (5-6 mL in lavender top tube) □ Muscle (~50mg flash frozen; shipped on dry ice) □ DNA (>20 ug): Tissue source concentration (ug/ml) Vol(ul) □ Oral Rinse (At least 30 mL of Scope oral rinse in a 50 mL centrifuge tube) □ Dried Blood Spots (2 cards) - Not accepted for any testing with a del/dup component □ Buccal Swab			
Mailing address	☐ Other(Call lab) Patient has had a blood transfusion ☐ Yes ☐ No Date of last transfusion/_/_			
City State Zip code	(2-4 weeks of wait time is required for mtDNA testing only) Fibroblasts are recommended for patients who had an allogenic bone marrow transplant.			
Home phone Work phone	See www.genedx.com/specimen-requirements for details. Clinical Diagnosis: ICD-10 Codes:			
Email Patient's primary language if not English	Age at Initial Presentation: Add. ICD-10 Codes:			
Ordering Account Information	Statement of Medical Necessity			
Acct # Account Name Reporting Preference*. Care Evolve Fax Email *If unmarked, we will use the account's default preferences or fax to new clients.	This test is medically necessary for the diagnosis or detection of a disease, illness, impairment, symptom, syndrome or disorder. The results will determine my patient's medical management and treatment decisions. The person listed as the Ordering Provider is authorized by law to order the tests(s) requested herein. I confirm that I have provided genetic testing information to the patient and the patient has			
Physician NPI #	consented to genetic testing.			
Genetic Counselor	Signature of Physician or Other Authorized NPI Provider (required) Date			
Street address I	Patient Consent (sign here) I have read the attached Informed Consent document and I give permission to			
Street address 2	GeneDx to perform genetic testing as described. I also give permission for my specimen and clinical information to be used in de-identified studies at GeneDx			
City State Zip code	to improve genetic testing and for publication, if appropriate. My name or other personal identifying information will not be used in or linked to the results of any			
Phone Fax (important)	studies and publications. I also give GeneDx permission to inform me or my health care provider in the future about research opportunities, including treatments for			
Email Beeper the condition in my family. More information is available on our website: www.genedx.com				
Physician or GC/Acct # Fax#/Email/CE #				
Physician or GC/Acct # Fax#/Email/CE #	Patient/Guardian Signature Date			
PATIENT STATUS – ONE MUST BE CHECKED: Hospital Inpatient Hospital C	Outpatient 🛘 Not a Hospital Patient 🛮 Hospital Patient Date of Discharge:			
Payme	ent Options			
□ Insurance Bill	Referral/Prior Authorization # Please attach copy of Referral/authorization GeneDx Benefit Investigation #			
Insurance Carrier Policy Name	fit Investigation (only if OOP cost is >\$100)			
Insurance ID # Group # Name of Insured	Date of Birth Insurance Address City State Zip			
Secondary Insurance Insurance ID# Group # Name of Insur	Relationship to Insured			
Carrier Name Please include a copy of the front and back of the patient's insurance of				
Please include a copy of the front and back of the patient's insurance card (include secondary when applicable) I represent that I am covered by insurance and authorize GeneDx, Inc. to give my designated insurance carrier, health plan, or third party administrator (collectively "Plan") the information on this form and other information provided by my health care provider necessary for reimbursement. I authorize Plan benefits to be payable to GeneDx. I understand that GeneDx will attempt to contact me if my estimated out-of-pocket responsibility will be greater than \$100 per test (for any reason, including co-insurance and deductible, or non-covered services). If GeneDx is unsuccessful in its attempts to contact me, I understand that it will be my responsibility to contact GeneDx to determine my out-of-pocket cost and to pay my out-of-pocket responsibility. I will cooperate fully with GeneDx by providing all necessary documents needed for Plan billing and appeals. I understand that I am responsible for sending GeneDx any and all of the money that I receive directly from my Plan in payment for this test. Reasonable collection and/or attorney's fees, including filing and service fees, shall be assessed if the account is sent to collection but said fees shall not exceed those permitted by state law. I permit a copy of this authorization to be used in place of the original.				
Patient Signature (required)	Date			
☐ Institutional Bill GeneDx Account #	☐ Patient Bill If I have insurance coverage for this testing, I am electing to be treated as a self-pay patient for this testing. As such, I agree that neither GeneDx nor I will submit a claim to my insurance for this testing.			
Hospital/Lab Name	Please bill my credit card (all major cards accepted) ☐ MasterCard ☐ Visa ☐ Discover ☐ American Express			
Contact Name	Name as it appears on card			
Address	Account Number Expiration date CVC			
City State Zip Code	Signature Date			
Phone Fax	For GeneDx Use Only			



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First Name	Last Nar	ne				Date of Birth (mm/dd/yy)
	Family His	story of	Disorde	r/Symptom	S	
□ No Known Family History □ Pedigree Attached □ Adopted	-	-	Paternal	Dis	order/Symptoms	
Other clinical history or testing (summarize or attach reports) Array CGH: Chromosomes/FISH: Other relevant results (clinical or research): The property of the pedigree and/or include additional information Draw/attach pedigree and/or include additional information					formation	
	Family	Membe	r/Carrie	r Testing		
Testing for known familial variant in a nuc ☐ 9011 Testing for ONE known familial varia ☐ 9012 Testing for TWO known familial var ☐ 905 Testing for ONE known familial exochromosomal microarray del/dup Testing for known mtDNA variant(s)	ant in a nuclear gene iants in a nuclear gene		Proband Proband Relations	Name: GeneDx Acc#:_ ship to proband: t/Carrier testing	Unaffected or asympto	omatic (Circle if applies)
□ 453 Testing for ONE to THREE mtDNA variant(s) (heteroplasmy detection range: 1.5%-100%) □ 9017 Testing for ONE mtDNA variant (heteroplasmy detection range: 25%-100%) □ 9020 Testing for TWO mtDNA variants (heteroplasmy detection range: 25%-100%)			Test of Positive caveate ☐ Family on the	was performed we control not avaluage on a row Member Test F	d at another lab. vailable. Please initial to legative report Report included - A cleate family member is reco	ir copy of the test report
	panded Family M	1ember	Testing (ı			
First Name: Last Name: Symptomatic Father:	sent later** DOB:		☐ Asym _l Other:	ne: otomatic	Last Name: Symptomatic later** Relationship to	o Proband: DOB: o Proband: DOB:
☐ Asymptomatic ☐ Symptomatic			** Additi	onal samples mu	ust be received within 3 and test selection	weeks
Single Gene Analysis/Write-in Test Selection						
☐ 906 Deletion/Duplication Analysis of ONE nuclear gene	Write in desired gene	:				
Test Code	Test Name					
Test Code	Test Name					
Please see our website (www.genedx.com/forms) or other requisitions for tests not included on this requisition.						



(seq & del/dup of 130 genes)

and del/dup testing

Mitochondrial and Metabolic Disorders Testing

Account # Account Name

First Name Last Name	Date of Birth (mm/dd/yy)

GeneDx Mitochondrial Genetic Testing Mitochondrial Disorders ☐ 577 Progressive External Ophthalmoplegia (PEO)/ Optic Atrophy □ J809 MitoXpanded Panel (~1800 genes)* ☐ 554 Concurrent full sequence analysis & deletion testing of the Nuclear Gene Panel (seq & del/dup of 44 genes) mito genome (not a trio based test) ☐ 578 Methylglutaconic Aciduria Nuclear Gene Panel (seq & del/dup of 14 genes) * "Xpanded Family Member Testing" portion must be completed from Page 2 ☐ 938 Congenital Sideroblastic Anemia Panel (ABCB7, ALAS2, GLRX5, PUSI, ☐ 615 Combined Mito Genome Plus Mito Focused Nuclear Gene Panel SLC19A2, SLC25A38, TRNT1, YARS2, Mitochondrial genome large deletion (seq & del/dup of mito genome and of 202 nuclear genes) ☐ 554 Full sequence analysis and deletion testing of the mitochondrial genome ☐ 704 65 mtDNA Point Variants Plus Large Deletions Panel ☐ 573 Mitochondrial Focused Nuclear Gene Panel (seq & del/dup 202 genes) ☐ 444 Deletion/duplication analysis of mito genome ☐ 575 Mitochondrial Encephalopathy/Leigh Syndrome Nuclear Gene Panel ☐ 394 POLG gene sequencing ☐ 557 PUSI Gene sequencing (seq & del/dup of 134 genes) ☐ 576 Lactic Acidosis/Pyruvate Metabolism Nuclear Gene Panel ☐ 582 SDHA Gene sequencing

GeneDx Metabolic Genetic Testing Inborn Errors of Metabolism Next-Generation Panels ☐ J979 Combined Lysosomal and Peroxisomal Disorders Panel ☐ T382 Fatty Acid Oxidation Disorders Panel (seq & del/dup of 15 genes) (seq & del/dup of 83 genes) ☐ T010 Hyperammonemia, Urea Cycle and Transporter Defects Panel ☐ T013 Lysosomal Disorders Panel (seq & del/dup of 58 genes) (seq & del/dup of 48 genes) ☐ J978 Peroxisomal Disorders Panel (seq & del/dup of 25 genes) ☐ T012 Metabolic Myopathy Panel (seq & del/dup of 30 genes) ☐ T011 Methylmalonic Acidemia, Disorders of Cobalamin Metabolism and ☐ J976 Creatine Deficiency Syndromes Panel (seq & del/dup of 3 genes) ☐ J977 Congenital Disorders of Glycosylation Panel Related Disorders Panel (seq & del/dup of 19 genes) (seq & del/dup of 108 genes) Riboflavin Transporter Deficiency and Related Disorders ☐ J980 Disorders Associated with C4 Elevation (seq & del/dup of 6 genes) (seq & del/dup of 9 genes) ☐ J995 Disorders of Hyperphenylalaninemia and Biopterin Metabolism Panel (seq & del/dup of 7 genes) **Inborn Errors of Metabolism Single Gene Tests** ■ 508 3-Hydroxyacyl-CoA dehydrogenase deficiency (HADH) ☐ 399 Glutaric aciduria type I (GCDH) ☐ 380 6-pyruvoyl--tetrahydropterin synthase deficiency (PTS) Glutaric aciduria II / Multiple acyl-CoA dehydrogenase deficiency (MADD) ☐ 354 ß-ketothiolase deficiency (ACATI) □ 280 ETFDH ☐ 279 ETFB ☐ 278 ETFA Glycerol kinase Deficiency (GK) ☐ 2631 Acid sphingomyelinase deficiency (SMPD1) ☐ J975 Adrenoleukodystrophy, X-linked (ABCD1 sequencing and deletion/ ☐ 438 GK sequencing ☐ 906 GK Exon-level deletion testing duplication testing) ■ 287 Glycogen storage disease II (Pompe disease) (GAA) ☐ 465 Arginase deficiency (ARGI) ☐ 649 Glycogen storage disease type V (GSD V) (PYGM) ☐ 426 Argininosuccinic Aciduria (ASL) ☐ 657 GMI-gangliosidosis (GLBI) ☐ 658 Aspartylglucosaminuria (AGA) ☐ 230 GTP cyclohydrolase I deficiency (GCHI) ■ 294 Biotinidase deficiency (BTD) ☐ 3211 HMG CoA lyase deficiency (HMGCL) ☐ 320 Holocarboxylase synthetase deficiency (HLCS) ☐ 564 Canavan disease (ASPA) sequencing and deletion/duplication testing ☐ 429 Carnitine-Acylcarnitine Translocase Deficiency (SLC25A20) ☐ 331 Homocystinuria (CBS) ☐ 425 Carnitine palmitoyltransferase IA deficiency (CPTIA) ☐ T386 Hurler syndrome/ mucopolysaccharidosis type I (IDUA) ☐ 334 Carnitine palmitoyltransferase deficiency type II (CPT2) ☐ 351 Isobutyryl CoA dehydrogenase deficiency (ACAD8) ☐ 500 Citrin Deficiency (SLC25A13) Isovaleric acidemia (IVD) ☐ 382 Classic Citrullinemia (ASSI) ☐ 3191 Full sequencing 3192 Sequence exon 9 only (includes common A282V mutation) ☐ 274 Cobalamin C deficiency (MMACHC) ☐ 659 Combined malonic and methylmalonic aciduria (ACSF3) ☐ 3193 Rest of IVD (if 3192 negative) ☐ 490 Dihydrolipoamide Dehydrogenase Deficiency (DLD) □ 507 Krabbe disease (GALC) ☐ 381 Dihydropteridine reductase (DHPR) deficiency (QDPR) LCHAD/trifunctional protein deficiency (HADHA/HADHA and HADHB) ☐ 558 Ethylmalonic Encephalopathy (ETHEI) sequencing and deletion/ □ 2711 HADHA Tier I (common mutation; c.1528G>C) duplication testing Reflex testing: HADHA (full), HADHB if necessary: 2712, 272 Fabry disease (GLA) □ 2712 HADHA Full sequencing □ 272 HADHB Full sequencing ☐ 2321 GLA sequencing Lowe syndrome (OCRL) 906 GLA deletion/duplication testing, females ☐ 335 Lowe syndrome, OCRL full sequencing ☐ 605 Free sialic storage disorders (SLC17A5) sequencing and deletion/ □ 906 OCRL deletion/duplication testing, females duplication testing ☐ 655 Lysosomal acid lipase deficiency (LIPA) ☐ 661 Fucosidosis (FUCAI) ☐ 404 Malonyl-CoA decarboxylase deficiency (MLYCD) ■ 2843 Fumarate hydratase deficiency (FH) Maple Syrup Urine Disease (MSUD) ☐ 499 Galactokinase Deficiency (GALKI) ☐ 4881 BCKDHA ☐ 4882 BCKDHB ☐ 4883 DBT ☐ 349E Galactosemia / Galactosyltransferase deficiency (GALT) sequencing ☐ 488 BCKDHA/ BCKDHB/ DBT All NOW



Mitochondrial and Metabolic Disorders Testing

Account # Account Name

First Name	Last Name	Date of Birth (mm/dd/yy)

GeneDx Metabolic Genetic Testing			
Inborn Errors of Metabolism Single Gene Tests (cont.)			
☐ 565 Maroteaux-Lamy syndrome/mucopolysaccharidosis VI (ARSB)	Ornithine transcarbamylase deficiency (OTC)		
MCAD deficiency (ACADM)	☐ 313 OTC sequencing (males)		
☐ 2682 Full gene sequencing NOW	☐ 313E OTC sequencing and deletion/duplication testing (females)		
2681 Sequence exon 11 only (includes common K329E mutation)	☐ 273 Phenylalanine hydroxylase (PAH)		
☐ 2683 Rest of ACADM	☐ 287 Pompe disease/glycogen storage disease type II (GAA)		
☐ 649 McArdle disease (PYGM)	Propionic acidemia		
☐ 563 Metachromatic leukodystrophy (ARSA)	□ 2901 Tier 1: PCCB □ 2902 Tier 2: PCCA, if necessary		
473 Methionine adenosyltransferase I/III deficiency (MATIA)	☐ 365 Primary/systemic carnitine deficiency (SLC22A5)		
2-Methyl-3-hydroxybutyryl-CoA dehydrogenase deficiency (HSD17B10)	☐ 528 PSAP-related disorders (PSAP)		
463 HSD17B10 sequencing	☐ 540 Pyruvate carboxylase deficiency (PC)		
906 HSD17B10 deletion/duplication testing, females Machalana and Co.A. and analysis defision res	Pyruvate Dehydrogenase EI-Alpha Deficiency (PDHAI)		
3-Methylcrotonyl CoA carboxylase deficiency ☐ 2881 Tier 1: MCCC2 ☐ 2882 Tier 2: MCCC1, if necessary	☐ 461 PDHA1 sequencing☐ 906 PDHA1 deletion/duplication testing, females		
□ 501 3-Methylglutaconic aciduria type I (AUH)	☐ 462 Pyruvate Dehydrogenase EI-Beta Deficiency (PDHB)		
Methylmalonic acidemia (MUT, MMAA, MMAB)	□ 605 Salla disease (SLC17A5) sequencing and deletion/duplication testing		
□ 2752 MUT full sequencing □ 276 MMAA □ 277 MMAB	□ 515 Sandhoff disease (HEXB) sequencing and deletion/duplication testing		
☐ MUT, MMAA, MMAB all NOW: 2752, 276, 277	Sanfilippo syndrome/ Mucopolysaccharidosis III (MPS IIIA, IIIB, IIIC, and IIID)		
☐ 657 Morquio B disease (GLBI)	☐ 591 Sanfilippo A (SGSH sequencing)		
☐ 608 Morquio syndrome A/ Mucopolysaccharidosis IVA (GALNS)	☐ 592 Sanfilippo B (NAGLU sequencing)		
☐ 648 Mucolipidosis I (NEUI)	☐ 593 Sanfilippo C (HGSNAT sequencing)		
☐ 2432 Mucolipidosis type IV (MCOLNI) sequence analysis	☐ 609 Sanfilippo D (GNS sequencing and deletion/duplication testing)		
☐ 906 MCOLNI deletion/duplication testing, including common 6.45kb deletion	☐ 610 SGSH/ NAGLU/ HGSNAT/ GNS All NOW		
☐ T386 Mucopolysaccharidosis type I (Hurler syndrome) (IDUA)	☐ 528 Saposin deficiency (combined, SapA, SapB, and SapC) (PSAP)		
Mucopolysaccharidosis III (MPSIII)/Sanfilippo syndrome (Types A, B, C and D)	Short/branched chain acyl-CoA dehydrogenase deficiency (ACADSB)		
☐ 591 MPSIII A (SGSH sequencing)	□ 383 Full Sequencing		
☐ 592 MPSIII B (NAGLU sequencing)	☐ 529 M389V (common Hmong mutation)		
☐ 593 MPSIII C (HGSNAT sequencing)	☐ 269 Short-chain acyl-CoA dehydrogenase (SCAD) deficiency (ACADS)		
☐ 609 MPSIII D (GNS sequencing and deletion/duplication testing)	Get Sialidosis (NEUI)		
 ☐ 610 SGSH/NAGLU/HGSNAT/GNS All now ☐ 608 Mucopolysaccharidosis IVA/Morquio syndrome A (GALNS) 	Smith-Lemli-Opitz syndrome (DHCR7) ☐ 2502 DHCR7 sequencing		
☐ 565 Mucopolysaccharidosis VI (Maroteaux-Lamy syndrome) (ARSB)	☐ 519 Tay-Sachs disease (HEXA)		
☐ 657 Mucopolysaccharidosis type IVB (GLBI)	Tyrosinemia		
☐ 611 Multiple sulfatase deficiency (SUMF1)	☐ 3661 Type I (FAH)		
☐ 478 N-acetylglutamate synthase deficiency (NAGS)	☐ 494 Type II (TAT)		
☐ 607 Neuronal ceroid-lipofuscinosis 2 (TPPI)	495 Type III (HPD)		
Niemann-Pick disease (NPD), type C	☐ 270 Very long chain acyl-CoA dehydrogenase (VLCAD) deficiency		
☐ 246 NPD type CI (NPCI) ☐ 247 NPD type C2 (NPC2/HEI)	(ACADVL)		
☐ 581 NPC1 and NPC2 both NOW	☐ J975 X-linked adrenoleukodystrophy (ABCD1 sequencing and deletion/duplication testing)		



☐ Cyclic vomiting

Mitochondrial and Metabolic Clinical Information

Account # Account Name

Date of Birth (mm/dd/yy) First Name Last Name **DETAILED MEDICAL RECORDS MUST BE ATTACHED** _ Age at Initial Presentation:___ Parent/Carrier testing-unaffected/no symptoms Perinatal History: Seizures/Epilepsy: **Skeletal/Limb Abnormalities:** ☐ Fetal hydrops
☐ IUGR ☐ Seizure disorder □ Contractures ☐ Scoliosis/kyphosis ☐ Abnormal EEG Oligohydramnios/polyhydramnios (circle if ☐ Epileptic encephalopathy Dysostosis multiplex Focal seizures 🗖 Chondrodysplasia punctata applies) ☐ Generalized seizures – checkboxes below:
☐ Absence ☐ Clonic ☐ Prematurity Gibbus deformity Osteopenia **Growth:** ☐ Myoclonic □ Tonic-clonic 🗖 Other skeletal anomalies (type:__ Failure to thrive ■ Vertebral anomalies ☐ Microcephaly
☐ Macrocephaly **Brain Malformations/Abnormal Imaging:** ☐ Abnormal brain imaging (type: **Genitourinary Abnormalities:** ☐ Short staturé ☐ White matter abnormalities ☐ Steroid-resistant nephrotic syndrome Abnormal renal function (type:_ ☐ Abnormalities of the basal ganglia **Physical/Cognitive Development:** ☐ Agenesis of the corpus callosum ☐ Renal tubulopathy Developmental regression ☐ Global developmental delay Brain atrophy **Laboratory Abnormalities:**☐ Abnormal NBS______ ☐ Hydrocephalus ☐ Intellectual disability/MR ☐ Lissencephaly ☐ Learning disability ☐ Abnormal enzymes_ Pontocerebellar hypoplasia ☐ Motor ďelay ☐ Neopterin/biopterin - check boxes
☐ CSF ☐ Other ☐ Speech delay Muscular: Myopathy **Behavioral:** ☐ Aminoaciduria ☐ Rhabdomyolysis/myoglobinuria Autism spectrum disorder ☐ Urine glycosaminoglycans Abnormal electromyography (EMG) Autistic features ☐ Acylcarnitines ■ Easy fatigue ☐ Behavioral/psychiatric abnormalities (circle if ☐ Transferrin ☐ Exercise intolerance ☐ Creatine/guanidinoacetate applies) Hypertonia ☐ Very long chain fatty acids ☐ Early onset dementia ☐ Hypotonia ☐ Pipécolic acid **Craniofacial/Dysmorphic Features:** ☐ Joint hypermobility
☐ Muscle fasciculations Phytanic acid ☐ Abnormal fat distribution Other: □ Coarse facies ☐ Muscle stiffness ☐ CPK abnormalities: (value:_ ☐ Facial dysmorphism – please describe ☐ Muscle wasting ☐ Elevated alanine ☐ Muscle weakness: proximal/distal/upper limb/ Elevated pyruvate lower limb (circle if applies) Hyperammonemia ☐ Inverted nipples ☐ Myotonia ☐ Hypoglycemia **Ophthalmalogic/Auditory:** ☐ Réspiratory insufficiency ☐ Kétosis ☐ Cataracts ☐ Lactic acidemia/high CSF lactate (circle if **Neurological:** Cherry red spot on macula applies) Dysphagia ☐ Corneal opacities ☐ Dysarthria☐ Myoclonus Low plasma carnitine ☐ CPEO (Ophthalmoplegia) Organic aciduria ☐ Eye movement disorder ■ Nérve conduction studies: Skin Abnormalities: ☐ Iris coloboma ☐ Ataxia Optic atrophy ☐ Scaly skin ■ Bulbar signs Ototoxicity (aminoglycoside-induced) ☐ Angiokeratoma ☐ Chorea ☐ Congenital neuropathy ☐ Ptosis Skin abnormality (type:_ ☐ Retinitis pigmentosa Distal motor neuropathy **Biopsy Abnormalities:** ☐ Sensorineural hearing loss Dystonia ☐ Muscle biopsy ☐ Vertical gaze palsy ☐ Episodic apnea (sudden) COX deficiency ☐ Nystagmus ☐ Hypomyelination ☐ Histology ☐ Strabismus ☐ Lower extremity weakness☐ Motor neuron dysfunction - check boxes: ☐ Large mitochondria (mt)/mt proliferation ☐ Retinal degeneration ☐ Ragged red fibers ☐ Pigmentary retinopathy Respiratory enzymes:
Ultrastructure (EM): ☐ Upper ☐ Lower ☐ Blindness/vision loss ☐ Pes cavus ☐ Recurrent ear infections Recurrent headache/migraine mtDNA depletion/multiple mtDNA deletions Other visual abnormality: ☐ Reduced/absent deep tendon reflexes Abnormal nerve biopsy ☐ Sensory neuropathy – check boxes:
☐ Hyperesthesia ☐ Paresthesia Cardiacvascular: ☐ Histochemical study ☐ Arrhythmia ☐ ASD/VSD (circle if applies) ☐ Hypéresthesia Glycogen storage ☐ Sleep apnea ☐ Other storage (type:_ ☐ Cardiomegaly ☐ Spasticity Other clinical history or testing: Cardiomyopáthy ☐ Stroke/stroke-like episodes ☐ Sideroblastic anemia Valvular heart disease ☐ Tremor/Parkinsonism (circle if applies) Pancytopenia ☐ Deep vein thrombosis ☐ Arraý CĠH: ☐ Abnormal sweating **Gastrointestinal:** Chromosomes/FISH: ☐ Chronic diarrhea☐ Constipation ☐ Abnormal temperature regulation Other relevant results (clinical or research): **Endocrine:** Draw/attach pedigree and/or include ☐ Delayed gastric emptying ☐ Pancreatic failure additional clinical information: Gastrointestinal reflux ☐ Adrenal insufficiency ☐ Nausea ☐ Diabetes mellitus – check boxes: ☐ Recurrent vomiting ☐ Type I ☐ Type II Hepatic dysfunction Hypoparathyroidism ☐ Hepatomégaly/splenomegaly (circle if applies) ☐ Hypothyroidism ☐ Cachexia ☐ Cholestasis



Informed Consent

Account # Account Name

First Name Last Name Date of Birth (mm/dd/yy)

I understand that my health care provider has ordered the following genetic testing for {me/my child}:

General Information About Genetic Testing

What is genetic testing?

DNA provides instructions for our body's growth and development. Genes are distinct sequences of DNA, and are arranged on chromosomes. The DNA in a gene contains instructions for making proteins, which determine things like growth and metabolism as well as traits like eye color and blood type. Genetic disorders are caused by certain changes in DNA affecting the structure or number of chromosomes. Genetic testing is a laboratory test that tries to identify these changes in chromosomes or the DNA. Genetic testing can be a diagnostic test, which is used to identify or rule out a specific genetic condition. Genetic screening tests are used to assess the chance for a person to develop or have a child with a genetic condition. Genetic screening tests are not typically diagnostic and results may require additional testing.

The purpose of this test is to see if I, or my child, may have a genetic variant or chromosome rearrangement causing a genetic disorder or to determine the chance that I, or my child, will develop or pass on a genetic disorder in the future. 'My child' can also mean my unborn child, for the purposes of this consent.

If I/my child already know the specific gene variant(s) or chromosome rearrangement that causes the genetic disorder in my family, I will inform the laboratory of this information.

What could I learn from this genetic test?

The following describes the possible results from the test:

- I) Positive: A positive result indicates that a genetic variant has been identified that explains the cause of my/my child's genetic disorder or indicates that I/my child am at increased risk to develop the disorder in the future. It is possible to test positive for more than one genetic variant.
- 2) Negative: A negative result indicates that no disease-causing genetic variant was identified by the test performed. It does not guarantee that I/my child will be healthy or free from genetic disorders or medical conditions. If I/my child test negative for a variant known to cause the genetic disorder in other members of my/my child's family, this result rules out a diagnosis of the same genetic disorder in me/my child due to this specific change.
- 3) Inconclusive/Variant of Uncertain Significance (VUS): A finding of a variant of uncertain significance indicates that a genetic change was detected, but it is currently unknown whether that change is associated with a genetic disorder either now or in the future. A variant of uncertain significance is not the same as a positive result and does not clarify whether I/my child is at increased risk to develop a genetic disorder. The change could be a normal genetic variant or it could be disease-causing. Further analysis may be recommended, including testing parents and other family members. Detailed medical records or information from other family members also may be needed to help clarify results.
- 4) Unexpected results: In rare instances, this test may reveal an important genetic change that is not directly related to the reason for ordering this test. For example, this test may tell me about the risk for another genetic condition I/my child is not aware of or it may indicate differences in the number or rearrangement of sex chromosomes. This information may be disclosed to the ordering health care provider if it likely impacts medical care.

Result interpretation is based on currently available information in the medical literature, research and scientific databases. Because the literature, medical and scientific knowledge are constantly changing, new information that becomes available in the future may replace or add to the information GeneDx used to interpret my/my child's results. Providers can contact GeneDx at any time to discuss the classification of an identified variant. In addition, I or my/my child's health care providers may monitor publicly available resources used by the medical community, such as ClinVar (www.clinvar.com), to find current information about the clinical interpretation of my/my child's variant(s).

For tests that evaluate data from multiple family members, my spouse, or partner concurrently, results may be included in a single comprehensive report.

What are the risks and limitations of this genetic test?

- Genetic testing is an important part of the diagnostic process.
 However, genetic tests may not always give a definitive answer. In some cases, testing may not identify a genetic variant even though one exists. This may be due to limitations in current medical knowledge or testing technology.
- Accurate interpretation of test results may require knowing the
 true biological relationships in a family. Failing to accurately state the
 biological relationships in my/my child's family may result in incorrect
 interpretation of results, incorrect diagnoses, and/or inconclusive
 test results. In some cases, genetic testing can reveal that the true
 biological relationships in a family are not as they were reported.
 This includes non-paternity (the stated father of an individual is not
 the biological father) and consanguinity (the parents of an individual
 are related by blood). It may be necessary to report these findings to
 the health care provider who ordered the test.
- Genetic testing is highly accurate. Rarely, inaccurate results may
 occur for various reasons. These reasons include, but are not limited
 to: mislabeled samples, inaccurate reporting of clinical/medical
 information, rare technical errors, or unusual circumstances such as
 bone marrow transplantation, or the presence of change(s) in such a
 small percentage of cells that the change(s) may not be detectable by
 the test (mosaicism).
- This test does not have the ability to detect all of the long-term
 medical risks that I/my child might experience. The result of this test
 does not guarantee my health or the health of my child/fetus. Other
 diagnostic tests may still need to be done, especially when only a
 genetic screening test has been performed previously.
- Occasionally, an additional sample may be needed if the initial specimen is not adequate.

Patient Confidentiality and Genetic Counseling

It is recommended that I receive genetic counseling before and after having this genetic test. I can find a genetic counselor in my area here: www.nsgc.org. Further testing or additional consultations with a health care provider may be necessary.

To maintain confidentiality, the test results will only be released to the referring health care provider, to the ordering laboratory, to me, to other health care providers involved in my/my child's diagnosis and treatment, or to others as entitled by law. The United States Federal Government has enacted several laws that prohibit discrimination based on genetic test results by health insurance companies and employers. In addition, these laws prohibit unauthorized disclosure of this information. For more information, I understand that I can visit www.genome.gov/10002077.

International Specimens

If I/my child reside outside the United States, I attest that by providing a sample for testing, I am not knowingly violating any export ban or other legal restriction in the country of my/my child's residence.

Additional information about the specific test being ordered is available from my health care provider or I can go to the GeneDx website, www.genedx.com. This information includes the specific types of genetic disorders that can be identified by the genetic test, the likelihood of a positive result, the limitations of genetic testing, as well as information about how specimens and information are stored and used.



A. Notifier:			
B. Patient Name:	C. Identification Number:		
Advance Beneficiary Notice of Noncoverage (ABN)			
NOTE: If Medicare doesn't pay for D.	below, you may have to pa	ay.	
Medicare does not pay for everything, e	ven some care that you or your health cal	re provider have	
good reason to think you need. We expe	ect Medicare may not pay for the D	below.	
D.	E. Reason Medicare May Not Pay:	F. Estimated Cost	
Note: If you choose Option 1 of that you might have, but	whether to receive the D. r 2, we may help you to use any other insomedicare cannot require us to do this.		
G. OPTIONS: Check only one bo	x. We cannot choose a box for you.		
also want Medicare billed for an official Summary Notice (MSN). I understand payment, but I can appeal to Medical does pay, you will refund any paymen OPTION 2. I want the Dask to be paid now as I am responsibl OPTION 3. I don't want the D	listed above. You may ask to be paral decision on payment, which is sent to me that if Medicare doesn't pay, I am response by following the directions on the MSN at I made to you, less co-pays or deductibusted above, but do not bill Medicate for payment. I cannot appeal if Medicate listed above. I understand with I cannot appeal to see if Medicare would be above.	e on a Medicare sible for . If Medicare les. are. You may the is not billed. this choice I	
H. Additional Information:			
	official Medicare decision. If you have	•	
• • • • • • • • • • • • • • • • • • •	D-MEDICARE (1-800-633-4227/TTY: 1-87	,	
ligning below means that you have rec	eived and understand this notice. You als	o receive a copy.	
g	5. 24.5.		
	programs and activities. To request this pull 0-MEDICARE or email: AltFormatReques		

According to the Paperwork Reduction Act of 1995, no persons are required to respond to a collection of information unless it displays a valid OMB control number. The valid OMB control number for this information collection is 0938-0566. The time required to complete this information collection is estimated to average 7 minutes per response, including the time to review instructions, search existing data resources, gather the data needed, and complete and review the information collection. If you have comments concerning the accuracy of the time estimate or suggestions for improving this form, please write to: CMS, 7500 Security Boulevard, Attn: PRA Reports Clearance Officer, Baltimore, Maryland 21244-1850.

Form CMS-R-131 (Exp. 03/2020)

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