GeneD

Age of Onset

All sections on this page are required unless otherwise specified. Incomplete information could result in a delay of testing.

PATIENT	INFORMATION
First Name	Last Name
Sex Assigned at Birth: O Male O Female Patient Karyotype (if known): Gender Identification (optional):	Date of Birth (mm/dd/yy)
Email	
Address	
City	State Zip Code
Phone (mobile preferred)	Is this patient deceased? O Yes ONo Deceased Date:
SAMPLE	INFORMATION
Date Sample Collected (mm/dd/yy)	Medical Record #
○Blood ○Buccal Swab ○Other (spec	cify source):
See www.genedx.com/specimen-requirem Patient has a personal history of a hema O Yes (specify diagnosis) If yes, please call the lab to discuss with a g	had an allogeneic bone marrow transplant. ments for details. atologic malignancy or disease ONo genetic counselor the most appropriate sample type
	DVIDER ATTESTATION er attests that (i) he/she authorizes and directs
GeneDx to perform the testing indicated; authorized by law to order the test(s) req Requisition Form ("TRF") are reasonable of treatment of a disease, illness, impairmer results will determine the patient's medic patient's condition on this date of service authorized to make decisions for the pati any relatives', when applicable, has been testing, and has consented to undergo gridagnosis codes are indicated to the high reimbursement from any third party, incluprograms if testing is covered by GeneDx GeneDx may share contact information for providers listed on the this order with third and potential clinical trial or study opport	; (ii) he/she is the ordering provider and is quested; (iii) any test(s) requested on this Test and medically necessary for the diagnosis or ent, symptom, syndrome or disorder; (iv) the test cal management and treatment decisions of this e; (v) the patient or the individual/family member icient (collectively, the "patient"), in addition to in supplied with information regarding genetic genetic testing; (vi) the full and appropriate hest level of specificity; (vii) he/she will not seek luding but not limited to federal healthcare is and will inform the patient of the same; (viii) for the ordering provider and other healthcare ard parties regarding the requested genetic testing trunities; and (ix) the patient or the individual/ted via the email address or mobile phone
	king this box, I confirm that the patient is a New ssion for GeneDx to retain any remaining sample is been completed.
Patient Research Opt-Out. By checking opt out of being contacted for research	ing this box, I confirm that the patient wishes to rch studies.
	 Check this box if your patient resides in CA, to opt-in to having their information shared for pation.
	ut. Check this box if your patient resides in any to opt-out of participation in Health Information

GeneDx Account Number	Account Name	
Phone	Fax	
Address		
City	State	Zip Code
Ordering Provider Name		Role/Title
NPI	Phone Number	
Send Report Via: ☐ Fax ☐ Email Fax #/Email:	Portal	
Additional Ordering Provider Name	e (optional)	Role/Title
NPI		
Send Report Via: ☐ Fax ☐ Email Fax #/Email:	Portal	
SEND ADDITIONAL REPORT COPIES T	O (optional)	
Provider Name	GeneDx Acct#	
Fax #/Email:		

ICD-10-CM CODES

ICD-10-CM Codes to support all test(s) ordered

Clinical Diagnosis

	PAYMENT O	PTIONS (Sele	ect One)			
O INSURANCE BILL	Patient Status					
Select all that apply	Is this individual currently a Hospital Inpatient? O Yes O No					
Commercial Medicaid	Name of Insurance Carrier		Insurance ID#:			
Medicare	Relationship to Ins	ured				
Tricare	OSelf OSpouse	OChild OOther	r:			
CHAMPVA	Policy Holder's Nar	me	Policy Holder's Date	of Birth		
FOR ALL INSURANCE PROVIDE FRONT AND BACK COPY OF	Referral/Prior Authorization # (please attach)		Hold test for cost estimate and contact patient if estimate is \$250 (for in-network/contracted commercial insurance only)			
CARD(S)	Secondary Insurance Type:					
	Insurance Carrier	Insurance ID #	Subscriber Name	Date of Birth		
	Relationship to Insured					
	OSelf OSpouse Ochild Oother:					
O PATIENT BILL	If Patient Bill is selected, I am electing to be treated as a self-pay patient for this testing. I agree that neither GeneDx nor I will submit a claim to my insurance for this testing, if I have insurance. GeneDx will send an invoice to the patient listed above.			ill submit a		
	Authorized Patient/Guardian Signature					
O INSTITUTIONAL BILL	GeneDx Account #	ŧ	51 01 10			
	Hospital/Lab Name		Place Sticker/Stamp Here			

Signature of Ordering Provider

Date



First Name | Last Name | Date of Birth |

CLINICAL INFORMATION (DETAILED MEDICAL RECORDS MUST BE ATTACHED)

CLINICAL INFORMATION (DETAILED MEDICAL RECORDS MUST BE ATTACHED)					
Is this person affected? O Yes O No	Clinical diagnosis:				
Reason for testing: Diagnosis Presymptomatic diagnosis Carrier/Familial Variant Testing					
Please check all that apply. This is not a substitute	e for submitting clinical records.				
Pre/Perinatal History	Developmental/Behavioral Findings	Hearing Impairment			
☐ Growth delay	(continued)	□ Abnormal newborn screen:			
☐ Increased body weight	☐ Intellectual disability	☐ Sensorineural hearing impairment/bilateral			
□ Intrauterine growth restriction □ Prematurity GA:	☐ Memory impairment ☐ OCD	Pospiratory Findings			
The maturity OA.	☐ Sleep disturbance	Respiratory Findings Apnea			
Structural Brain Abnormalies	☐ Specific learning disability	☐ Hyperventilation			
☐ Abnormal myelination	☐ Speech articulation difficulties	☐ Hypoventilation			
☐ Abnormality of basal ganglia	☐ Stereotypy	□ Respiratory distress			
☐ Abnormality of brainstem		□ Respiratory insufficiency			
☐ Abnormality of periventricular white matter	Neurological Findings				
☐ Abnormality of the corpus callosum ☐ Aplasia/hypoplasia of cerebellar vermis	☐ Abnormality of nervous system	Gastrointestinal Findings			
☐ Aplasia/hypoplasia of cerebellum	□ Ataxia □ Cerebral palsy	☐ Failure to thrive			
☐ Brain atrophy	☐ Chorea	☐ Feeding difficulties			
□ Cerebellar atrophy	☐ Cortical visual impairment	Musculoskeletal Findings			
□ Cerebellar hypoplasia (pontocerebellar	□ Dementia	☐ Arthrogryposis			
hypoplasia)	□ Dysarthria	☐ Decreased muscle mass			
☐ Chiari malformation ☐ CNS hypomyelination	□ Dyskinesia	☐ Exercise intolerance			
☐ Cortical dysplasia	□ Dysphasia □ Dystonia	□ Fasciculations			
☐ Cortical tubers	☐ Encephalopathy	☐ Fatigue			
☐ Frontotemporal cerebral atrophy	□ Epileptic encephalopathy	☐ Foot dorsiflexor weakness (foot drop) ☐ Hypertonia			
☐ Heterotopia (periventricular nodular	Familial or sporadic hemiplegic migraine	Hypotonia			
heterotopia) Holoprosencephaly	☐ Febrile seizures	☐ Joint hypermobility			
☐ Hydrocephalus	☐ Focal seizures ☐ Frontotemporal dementia	□ Muscle cramps			
Leukodystrophy	☐ Generalized seizures	Muscle weakness			
Lissencephaly	_ ☐ Headaches	☐ Myalgia			
☐ Molar tooth sign on MRI	□Hyperreflexia	☐ Myopathic facies ☐ Myopathy			
□ Pachygyria	☐ Infantile spasms	Pain			
□ Polymicrogyria □ Pontocerebellar atrophy	□ Myotonia □ Myoclonus	□ Pes cavus			
Subcortical band heterotopia	□ Paresthesia	□ Pes planus			
□ Ventriculomegaly	□ Parkinsonism	☐ Rhabdomyolysis			
	□ Peripheral neuropathy	□ Scoliosis □ Short stature			
Developmental/Behavioral Findings	☐ Reduced tendon reflexes	- Chort stature			
☐ Abnormal aggressive, impulsive or violent	Seizures	Skin/Hair Findings			
behavior Abnormal social behavior	☐ Sensory neuropathy ☐ Spasticity	☐ Axillary freckling			
Absent speech	☐ Status epilepticus	□ Café-au-lait macules			
Aggressive behavior	□ Stroke-like episode	☐ Hyperpigmentation of the skin			
□ Anxiety	Tremors	☐ Hypopigmentation of the skin			
☐ Attention deficit hyperactivity disorder	Upper motor neuron dysfunction	Metabolic Issues/Mitochondrial			
☐ Autistic behavior	☐ Vocal cord paresis	(attach relevant lab reports/values)			
☐ Behavioral abnormality ☐ Clumsiness	Craniofacial/Dysmorphism	Abnormal newborn screen results:			
Cognitive impairment	Abnormal facial shape (Dysmorphic	Abriornariewborn screen results.			
☐ Delayed fine motor development	features)	☐ Elevated CPK:			
□ Delayed gross motor development	☐ Macrocephaly				
Delayed speech & language development	□ Microcephaly´				
☐ Depression		Endocrine Findings			
☐ Developmental regression☐ Frequent falls	Eye Defects/Vision	□ Delayed puberty			
☐ Gait disturbance	☐ Abnormality of vision				
□ Global developmental delay	□ Cataracts	Vascular System			
□Hyperactivity	□ Nystagmus	☐ Arteriovenous malformation			
☐ Incoordination	□ Optic atrophy	□ Stroke			
	Cardiac Findings				
	□ Cardiac rhabdomyoma	☐ Other:			
	Cardiac defect:				



First Name							
First Name Last Name					Date of Birth		
FAMILY HISTORY							
□ No Known Family History	□Р	edigree Atto	ached	☐ Adopted			
Relationship	Materna	l Paternal		Relevant i	History		Age at Dx
1	0	0					
2	0	0					
3	0	0					
			PREVIOUS GE	NETIC TESTING			
Personal or family history a	of genetic tes	ting ON		lease complete all field	de below)		
· · ·			- , , , ,	· · · · · · · · · · · · · · · · · · ·			
Relation to patient (self, sibling,	, etc.), Genetic	Test(s) and R	esult (e.g. positive, neg	ative, etc.). If relative was	tested at GeneD	x, please also provide their	accession #:
If patient or relative(s) were for Indicate any Variants of Interes			S result on prior testing	g, please provide details b	elow.		
Relation (self, sibling, etc.)	Gene	Transcrip	t# c./p. (SN	IV) or exon # (CNV)	Build,	coordinates (CNV)	Variant of Interest‡?
1							
2							
3							
Required for sequence variants: gene, c./p., transcript # Required for CNVs: gene, transcript #, exon # OR build, coordinates							
Abnormal karyotype, FISH, or of	-	iia, coordii lates					
,							
‡ For certain tests, GeneDx may be a must be provided in the table above not be possible to comment upon the	e at the time the	test order is pla	ced. If you do not complet	e the table above and check (off that a previously	/ identified variant is a variant o	
			TARCETER	DIANT TECTING			
Individual to be tested: O	Affacted/Syr	motomatic		Asymptomatic	<u> </u>		
☐ Known Familial Variant(s) in		•		t Identified in Research La	h DTaraeted	Mosaic Variant Testing*	
☐ Known Familial Copy Numb			Known mtDNA Variant(*Insuranc	e Billing NOT Accepted; Pa	
Proband Name		Relat	tionship to Proband		Institution Proband Genel	nal Bill MUST be selected or Ox Accession #	n page 1
□Posit	ive control incl	uded/will be	sent - Positive control	previous test was perform is recommended if previon included on a negative re	ous test was per		
VARIANT INFORMATION		-			-	Number of Variants:	
Gene	Codi	ng DNA (c./m.)		Amino Acid (p.)		Transcript (NM#)	
Gene	Codi	ng DNA (c./m.)		Amino Acid (p.)		Transcript (NM#)	
COPY NUMBER VARIANT				1		Number of Variants:	
Gene(s)	Exon	#		Coordinates		Genome Build	
Gene(s)	Fyon	#		Coordinates		Genome Ruild	



First Name Last Name Date of Birth

	TEST MENU					
TEST CODE	TEST NAME	TEST CODE	TEST NAME			
□ 910	Chromosomal Microarray (MicroarrayDx)	□ 522	Fragile X Syndrome (FMR1 repeat analysis)			
NEUROD	EVELOPMENTAL DISORDERS AND EPILEPSY					
□ ⊤395	Autism/ID Panel (seq & del/dup of 103 genes)	□ 729	Rett/Angelman Related Disorders Panel (seq & del/dup of 25 genes & methylation MLPA)			
□ 523	Comprehensive Epilepsy Panel (seq & del/dup of 144 genes)	□ 730	Tuberous Sclerosis Panel (TSC1 & TSC2 seq & del/dup)			
□ 921	Epi <i>Xpanded®</i> Panel (1300+ genes, trios preferred)	☐ TJ27	Angelman Syndrome/Prader-Willi Syndrome Methylation MLPA (UPD, deletion)			
□ 953	Epilepsy Del/Dup Panel (128 genes) (not a trio based test)	□ T400	Hemiplegic Migraine Panel (seq & del/dup of 4 genes)			
CNS MA	LFORMATIONS AND DISORDERS					
□ 691	Comprehensive Brain Malformations Panel (seq & del/dup of 103 genes)	☐ J511	Microcephaly Xpanded® Panel (800+ genes, trios preferred)			
□ 526	Cerebral Cavernous Malformations (<i>KRIT1, CCM2, PDCD10</i> seq & del/dup)	□ J853	Leukodystrophy Xpanded® Panel (300+ genes, trios preferred)			
☐ TB51	Comprehensive Holoprosencephaly Panel (seq & del/dup of 17 genes)	□ 552	X-linked Hydrocephalus/X-linked Spastic Paraplegia/MASA/CRASH Syndrome (LICAM seq & del/dup)			
□ T844	Dementia Panel (seq only of 11 genes, for patients 18 years and olde	er)				
MOVEMI	ENT DISORDERS					
☐ J762	Ataxia Xpanded® Panel (1300+ genes, trios preferred)	☐ TH83	Spinocerebellar Ataxia Repeat Expansion Analysis (ATXNI, ATXN2, ATXN3, ATXN7, ATXN8, CACNAIA repeat)			
☐ TH97	Dentatorubral-Pallidoluysian Atrophy Repeat Analysis (ATNI repeat)		☐ TH84 Spinocerebellar Ataxia Type 1 Repeat Analysis (<i>ATXN1</i> repeat)			
☐ T402	Dystonia and Parkinsonism Panel (seq & del/dup of 103 genes)		☐ TH85 Spinocerebellar Ataxia Type 2 Repeat Analysis (ATXN2 repeat) ☐ TH86 Spinocerebellar Ataxia Type 3 Repeat Analysis (ATXN3 repeat)			
	☐ T403 Dystonia Panel (seq & del/dup of 83 genes)		☐ TH87 Spinocerebellar Ataxia Type & Repeat Analysis (CACNAIA repeat)			
	☐ T401 Parkinson Disease Panel (seq & del/dup of 44 genes)		☐ TH88 Spinocerebellar Ataxia Type 7 Repeat Analysis (ATXN7 repeat)			
☐ TH95	Friedreich Ataxia Repeat Analysis (FXN repeat)		☐ TH89 Spinocerebellar Ataxia Type 8 Repeat Analysis (ATXN8 repeat)at)			
☐ TH94	Friedreich Ataxia Sequencing & Del/Dup (FXN seq & del/dup)					
☐ TL12	Spinocerebellar Ataxia and Related Disorders Panel (seq & del/dup of 56 genes)	☐ TK79	Xpanded® Adult Movement Disorders Panel (500+ genes, trio preferred)			
NEURON	MUSCULAR DISORDERS					
□ J805	Amyotrophic Lateral Sclerosis/Frontotemporal Lobar Degeneration (C9orf72 repeat analysis, for patients 18 years and older)	□ 737	Hereditary Neuropathy Panel (seq & del/dup of 89 genes)			
□ T404	Amyotrophic Lateral Sclerosis/Frontotemporal Lobar	□ 818	Myotonic Dystrophy 1 (DM1) (DMPK repeat analysis)			
	Degeneration Panel (seq & del/dup of 24 genes, for patients 18 years and older)	□ 819	Myotonic Dystrophy 2 (DM2) (CNBP repeat analysis)			
	Order of Reflex Testing:					
	☐ Activate J805, if non-diagnostic activate T404					
☐ TG80	Arthrogryposis Panel (seq & del/dup of 90 genes)	☐ TG82	Myotonia Panel (<i>CNBP</i> and <i>DMPK</i> repeat analysis, seq & del/dup of 8 genes)			
☐ TG78	Congenital Hypotonia Evaluation (SMN1, SMN2, DMPK, 15q11.2-q13.1)	□ 743	Oculopharyngeal Muscular Dystrophy (<i>PABPN1</i> repeat analysis)			
∏ Т G 77	Congenital Hypotonia <i>Xpanded®</i> Panel (1400+ genes; trios preferred)	□ 889	Neuromuscular Disorders Panel (115 genes) 390 Limb-Girdle Muscular Dystrophy Panel			
□ 742	CMTIA/HNPP (<i>PMP22</i> del/dup)	☐ TG81	Periodic Paralysis Panel (seq & del/dup of 9 genes)			
☐ GD1007	Duchenne/Becker MD (<i>DMD</i> seq & del/dup)	☐ T789	SMN1/2 Dosage Analysis			
□ 820	Spinal & Bulbar Muscular Atrophy (AR repeat analysis)	<u> </u>	1			

GeneDx tests are frequently updated and improved based upon the most recent scientific evidence. The test codes, genes, and gene quantities listed on this test requisition are subject to change by GeneDx at any time. The most current test menu, list of genes, and technical limitations included for a specific test panel may be found on our website, genedx.com. Please note that GeneDx reserves the right to modify and upgrade any ordered panel to the version currently listed on our website.



NEUR	OLOGY TEST REC	QUISITION FORM		Genelx				
First Name		Last Name		Date of Birth				
	TEST MENU (continued)							
TEST CODE	TEST	NAME	TEST	TEST NAME				
	MITOCHONDRIAL DISORDERS							
☐ 615	Combined Mito Genome Plus Mi	to Focused Nuclear Gene Panel	☐ TH12	Leber Hereditary Optic Neuropathy (LHON) Panel				
□ 554	Full sequence analysis and deletic	on testing of the mitochondrial gen	ome (not a tric	b based test)				
NEURO	METABOLIC DISORDERS							
□ J976	Creatine Deficiency Syndromes	Panel (seq & del/dup of 3 genes)	□ TH08	Pompe Disease/Glycogen Storage Disease Type II (GAA seq and del/dup)				
☐ TG94	Gaucher Disease (GBA seq)		☐ TG92	Wilson Disease (ATP7B seq & del/dup)				
☐ T012	Metabolic Myopathy Panel (seq	& del/dup of 30 genes)	□ J975	X-linked Adrenoleukodystrophy (ABCD1 seq & del/dup)				
NEURO	FIBROMATOSIS							
□ 961	Comprehensive NF Panel: NF1, SF	RED1, NF2 and SMARCB1 sequenci	ng and deletio	on/duplication testing				
□ 962	NF1 Panel: NF1 and SPRED1 sequer	ncing and deletion/duplication te	sting					
□ 963	NF2 Panel: LZTR1, NF2 and SMARC	B1 sequencing and deletion/dupl	ication testing	3				
☐ TA06	Reflex to Noonan Syndrome and	RASopathies panel (sequencing	of 25 genes) i	f 962 is non-diagnostic				
CUSTO	M DEL/DUP TESTING							
□ 906	Deletion/Duplication Analysis of	ONE Nuclear Gene	□ 703	Deletion/Duplication Analysis of 2-20 Nuclear Genes				
Write-in D	esired Gene(s) to be Tested:							
WRITE-	IN TEST SELECTION							
☐ Test	Code:	Test Name:						
		FAMILY MEMBER F	OR PANEL	TESTING OPTION				
NO SEPAR	ATE REPORT, ADDITIONAL SAMPLE	S MUST BE RECEIVED WITHIN 3 WE	EKS OF PROB	AND SAMPLE. See Test Menu page for proband test selection.				
☐ J767 ☐ TG86 ☐ 923 ☐ 725	Ataxia <i>Xpanded</i> ®, Family mer Congenital Hypotonia <i>Xpana</i> Epi <i>Xpanded</i> ®, Family membe Chromosomal Microarray Pa	led®, Family member testing er testing	☐ J854 ☐ J513 ☐ TK80	Leukodystrophy <i>Xpanded</i> ®, Family member testing Microcephaly <i>Xpanded</i> ®, Family member testing <i>Xpanded</i> ® Adult Movement Disorders Panel, Family member testing				
	First Name	Last Name	DOB	O Asymptomatic O Symptomatic				
Biologico Mother	1			O At GeneDx (Accession #:)				
	First Name	Last Name	DOB	O Not available O To be sent within 3 weeks				
Biologico				O Asymptomatic O Symptomatic O At GeneDx (Accession #:				
Father				O Not available O To be sent within 3 weeks				
	Relationship to Proband							
Other	First Name	Last Name	DOB	O Asymptomatic O Symptomatic				
Biologico Relative				O At GeneDx (Accession #:) O Not available O To be sent within 3 weeks				

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	Get a signature for medical necessity and patient consent

GeneDx tests are frequently updated and improved based upon the most recent scientific evidence. The test codes, genes, and gene quantities listed on this test requisition are subject to change by GeneDx at any time. The most current test menu, list of genes, and technical limitations included for a specific test panel may be found on our website, genedx.com. Please note that GeneDx reserves the right to modify and upgrade any ordered panel to the version currently listed on our website.

INFORMED CONSENT



First Name Last Name Date of Birth

For the purposes of this consent, "I", "my", and "your" will refer to me or to my child, including my unborn child, if my child is the person for whom the healthcare provider has ordered testing.

PURPOSE OF THIS TEST

The purpose of this test is (a) to see if I may have a genetic variant or chromosome rearrangement causing a genetic disorder; or (b) to evaluate the chance that I will develop or pass on a genetic disorder in the future. If I already know the specific gene variant(s) or chromosome rearrangement that causes the genetic disorder in my family, I agree to inform the laboratory of this information.

WHAT TYPE OF TEST RESULTS CAN I EXPECT FROM GENETIC TESTING?

- 1. <u>Positive</u>: A change in your DNA was found, which is very likely the cause of your features/symptoms. This is the most straightforward test result, which can be used as the basis to test other family members to determine their chances of having either the disease or a child with the disease.
- 2. <u>Negative</u>: No variants were found to explain your symptoms. This does not mean that you do not have a genetic condition. It is still possible that there is a genetic variant not found by the test that was ordered. Your healthcare provider or genetic counselor may discuss more testing either now or in the future.
- 3. <u>Variant of Uncertain Significance (VUS)</u>: A change in a gene was found. However, we are not sure whether this variant is the cause of your symptoms/features. More information is needed. We may suggest testing other family members to help figure out the meaning of the test result.
- 4. <u>Unexpected Results (ACMG Secondary Findings)</u>: 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 find you are at risk for another genetic condition I am not aware of or it may indicate differences in the number or rearrangement of sex chromosomes. We may disclose this information to the ordering healthcare provider if it likely affects medical care.

Because medical and scientific knowledge is constantly changing, new information that becomes available may supplement the information GeneDx used to interpret my results. Healthcare providers can contact GeneDx at any time to discuss the classification of an identified variant.

WHAT IS TRIO/DUO-BASED GENETIC TESTING?

For some genetic tests, including samples from the biological parents and/or other biological relatives along with the patient's sample can help with the interpretation of the test results. These tests are often referred to as "trio tests" since they typically include samples from the patient and both parents.

Samples from relatives should be submitted with the patient's sample. Clinical information must be provided for the patient and any relative who submits a sample.

I understand that GeneDx will use the relative sample(s) when needed for the interpretation of my test results and that my test report may include clinical and genetic information about a relative when it is relevant to the interpretation of the test results. I further understand that relatives will not receive an independent analysis of data nor a separate report.

RISKS AND LIMITATIONS OF GENETIC TESTING

- 1. 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.
- 2. Accurate interpretation of test results may require knowing the true biological relationships in a family. I understand that if I fail to accurately state the biological relationships in my family, it could lead to incorrect interpretation of the test results, incorrect diagnoses, and/or inconclusive test results. If genetic testing reveals that the true biological relationships in a family are not as I reported them, including non-paternity (the reported father is not the biological father) and consanguinity (the parents are related by blood), I agree to have these findings reported to the healthcare provider who ordered the test.
- 3. Although genetic testing is highly accurate, inaccurate results may occur. These reasons include, but are not limited to mislabeled samples, inaccurate reporting of clinical/medical information, rare technical errors, or other reasons.
- 4. I understand that this test may not detect all of the long-term medical risks that I might experience. The result of this test does not guarantee my health and that additional diagnostic tests may still need to be done.
- 5. I agree to provide an additional sample if the initial sample 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 at www.nsgc.org. Further testing or additional consultations with a healthcare provider may be necessary.

To maintain confidentiality, test results will only be released to the referring healthcare provider, the ordering laboratory, to me, to other healthcare providers involved in my care, diagnosis and treatment, or to others with my consent or as permitted or required by law. Federal laws prohibit unauthorized disclosure of this information. More information can be found at: www.genome.gov/10002077

SAMPLE RETENTION

After testing is complete, my sample may be de-identified and be used for test development and improvement, internal validation, quality assurance, and training purposes. GeneDx will not return DNA samples to you or to referring healthcare providers, unless specific prior arrangements have been made.

I understand that samples from residents of New York State will not be included in the de-identified research studies described in this authorization and GeneDx will not retain them for more than 60 days after test completion, unless specifically authorized by my selection. The authorization is optional, and testing will be unaffected if I do not check the box for the New York authorization language. GeneDx will not perform any tests on the biological sample other than those specifically authorized.

DATABASE PARTICIPATION

De-identified health history and genetic information can help healthcare providers and scientists understand how genes affect human health. Sharing this de-identified information helps healthcare providers to provide better care for their patients and researchers to make new discoveries. GeneDx shares this type of information with healthcare providers, scientists, and healthcare databases. GeneDx will not share any personally identifying information and will replace the identifying information with a unique code not derived from any personally identifying information. Even with a unique code, there is a risk that I could be identified based on the genetic and health information that is shared. GeneDx believes that this is unlikely, though the risk is greater if I have already shared my genetic or health information with public resources, such as genealogy websites.

EPILEPSY PARTNERSHIP PROGRAM PARTICIPATION

I understand that GeneDx will send de-identified test results data, excluding ACMG secondary findings, to third parties for research or commercial purposes and that GeneDx is compensated for the provision of testing services and for data sharing with third parties that is compliant with applicable law. At no time will GeneDx share any patient personally identifiable information. GeneDx may share contact information for providers listed on the Test Requisition Form with third parties.

INFORMED CONSENT



First Name	Last Name	Date of Birth

PATIENT RECONTACT FOR RESEARCH PARTICIPATION

GeneDx may collaborate with other scientists, researchers and drug developers to advance knowledge of genetic diseases and to develop new treatments. If there are opportunities to participate in research relevant to the disorder in (my/my child's) family, GeneDx may contact my healthcare provider for research purposes, such as the development of new testing, drug development, or other treatment modalities. In some situations, such as if my healthcare provider is not available, I may be contacted directly. I can opt out of being contacted directly regarding any of the above activities by having my healthcare provider check the box for Patient Research Opt-Out. Any research that results in medical advances, including new products, tests or discoveries, may have potential commercial value and may be developed and owned by GeneDx or the collaborating researchers. If any individuals or corporations benefit financially from these studies, no compensation will be provided to (me/my child) or to (my/my child's) heirs.

EXOME/GENOME SEQUENCING SECONDARY FINDINGS

- · Applicable only for full exome sequencing and genome sequencing tests
- Does not pertain to Xpanded® or Slice tests

As many different genes and conditions are analyzed in an exome or genome sequencing test, these tests may reveal some findings not directly related to the reason for ordering the test. Such findings are called "incidental" or "secondary" and can provide information that was not anticipated.

Secondary findings are variants, identified by an exome or genome sequencing test, in genes that are unrelated to the individual's reported clinical features.

The American College of Medical Genetics and Genomics (ACMG) has recommended that secondary findings identified in a specific subset of medically actionable genes associated with various inherited disorders be reported for all probands undergoing exome or genome sequencing. Please refer to the latest version of the ACMG recommendations for reporting of secondary findings in clinical exome and genome sequencing for complete details of the genes and associated genetic disorders. Reportable secondary findings will be confirmed by an alternate test method when needed.

WHAT WILL BE REPORTED FOR THE PATIENT?

All pathogenic and likely pathogenic variants associated with specific genotypes identified in the genes (for which a minimum of 10X coverage was achieved by exome sequencing or a minimum of 15X coverage was achieved by genome sequencing), as recommended by the ACMG.

WHAT WILL BE REPORTED FOR RELATIVES?

The presence or absence of any secondary finding(s) reported for the proband will be provided for all relatives analyzed by an exome or genome sequencing test.

LIMITATIONS

Pathogenic and/or likely pathogenic variants may be present in a portion of the gene not covered by this test and therefore are not reported. The absence of reportable secondary findings for any particular gene does not mean there are no pathogenic and/or likely pathogenic variants in that gene. Pathogenic variants and/or likely pathogenic variants that may be present in a relative, but are not present in the proband, will not be identified nor reported. Only changes at the sequence level will be reported in the secondary findings report. Larger deletions/duplications, abnormal methylation, triplet repeat or other expansion variants, or other variants not routinely identified by clinical exome and genome sequencing will not be reported.

FINANCIAL AGREEMENT AND GUARANTEE

For insurance billing, I understand and authorize GeneDx to bill my health insurance plan on my behalf, to release any information required for billing, and to be my designated representative for purposes of appealing any denial of benefits. I irrevocably assign to and direct that payment be made directly to GeneDx.

I understand that my out-of-pocket costs may be different than the estimated amount indicated to me by GeneDx as part of a benefit investigation. I agree to be financially responsible for any and all amounts as indicated on the explanation of benefits issued by my health insurance plan. If my insurance provider sends a payment directly to me for services performed by GeneDx on my behalf, I agree to endorse the insurance check and forward it to GeneDx within 30 days of receipt as payment towards GeneDx's claim for services rendered.

rego to p fam any	by signing this form: (i) I acknowledge that I have read or have had read to me the GeneDx Informed Consent document, and understand the information egarding genetic testing; (ii) I have had the opportunity to ask questions about the testing, the procedure, the risks, and the alternatives; (iii) I authorize GeneDx o perform genetic testing as ordered; (iv) I understand that, for tests that evaluate data from multiple family members concurrently, test results from these amily members may be included in a single comprehensive report that will be made available to all tested individuals and their healthcare providers; (v) if at any time I or my provider provide an email address or mobile phone number at which I may be contacted, I consent to receiving email or text messages from GeneDx; and (vi) I understand that this consent applies to all future communications unless I request a change in writing.				
	Secondary Findings Opt-out. Check this box if you do not wish to receive AC ONLY; not for $\textit{Xpanded}^{\circledast}$ or Slice tests).	CMG secondary findings (Full Exome Sequencing and Ge	enome Sequencing Tests		
	New York Retention Opt-in. By checking this box, I confirm that I am a New York State resident, and I give permission for GeneDx to retain any remaining sample longer than 60 days after the completion of testing, and to be used as a de-identified sample for test development and improvement, internal validation, quality assurance, and training purposes. Otherwise, New York law requires GeneDx to destroy my sample within 60 days, and it cannot be used for test development studies.				
	Patient Research Opt-out. Check this box if you wish to opt out of being con	tacted for research studies.			
	Health Information Exchange Opt-in. Check this box if you reside in CA, FL, MA, NV, NY, RI, and VT and wish to opt-in to my health information to be shared for Health Information Exchange participation.				
	Health Information Exchange Opt-out. Check this box if you reside in any other US state or territory and wish to opt-out of participation in Health Information Exchange.				
ignature of Patient/Legal Guardian (required) Date					
igna	gnature of Relative A/Legal Guardian Relative A Relationship to Patient Date				
Signa	nature of Relative B/Legal Guardian Relative B Relationship to Patient Date				