

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

	PATIENT INI	FORMATION		
irs	t Name	Last Name		
Pati	Assigned at Birth: Male Female ent Karyotype (if known): der Identification (optional): il	Date of Birth (mm/dd/	⁽ уу)	
Add	lress			
City		State	Zip Code	
Pho	ne (mobile preferred)	Is this patient decease Deceased Date:	d? O Yes ONo	
) est	SAMPLE INF e Sample Collected (mm/dd/yy)	ORMATION Medical Record #		
Jul	e sumple collected (mm/dd/yy)	Medical Record #		
OE	Blood OBuccal Swab Other (specify s	source):		
	reatment-related RUSH (optional) son: OTransplantation OPregnancy (Surgery Other		
	ient has had a blood transfusion () Yes (nsfusion:	
2-4	weeks of wait time is required for some te	esting)		
ibr	ient has had an allogeneic bone marrow oblasts are required for patients who had owww.genedx.com/specimen-requirement	an allogeneic bone marı		
ati	ient has a personal history of a hematolo	gic malignancy or dise	ase	
	res (specify diagnosis)		ONo	
ye	s, please call the lab to discuss with a genet	tic counseior the most ap	ppropriate sample type.	
	ORDERING PROVI	DER ATTESTATIO	N	
By signing this form, the ordering provider attests that (i) he/she authorizes and directs GeneDx to perform the testing indicated; (ii) he/she is the ordering provider and is authorized by law to order the test(s) requested; (iii) any test(s) requested on this Test Requisition Form ("TRF") are reasonable and medically necessary for the diagnosis or treatment of a disease, illness, impairment, symptom, syndrome or disorder; (iv) the test results will determine the patient's medical management and treatment decisions of this patient's condition on this date of service; (v) the patient or the individual/family member authorized to make decisions for the patient (collectively, the "patient"), in addition to any relatives', when applicable, has been supplied with information regarding genetic testing, and has consented to undergo genetic testing; (vi) the full and appropriate diagnosis codes are indicated to the highest level of specificity; (vii) he/she will not seek reimbursement from any third party, including but not limited to federal healthcare programs if testing is covered by GeneDx and will inform the patient of the same; (viii) GeneDx may share contact information for the ordering provider and other healthcare providers listed on the this order with third parties regarding the requested genetic testing and potential clinical trial or study opportunities; and (ix) the patient or the individual/family member authorized to be contacted via the email address or mobile phone number provided for this and future testing.				
	New York Retention Opt-In. By checking t York State resident who gives permission longer than 60 days after testing has bee	for GeneDx to retain any		
	Patient Research Opt-Out. By checking the opt out of being contacted for research s		e patient wishes to	
	Health Information Exchange Opt-in. ChrFL, MA, NV, NY, RI, and VT and wishes to op Health Information Exchange participatio	t-in to having their infor		
	Health Information Exchange Opt-out. C other US state or territory and wishes to o Exchange.			

Л				Gene D
	ACCOUNT IN	IFORMAT	TION	
GeneDx Account Nu	mber	Account No	ame	
Phone		Fax		
Address				
City		State		Zip Code
Ordering Provider N	ame			Role/Title
NPI		Phone Num	ber	
Send Report Via: Fax #/Email:	Fax			
Additional Ordering	Provider Name (optiona	I)		Role/Title
NPI				
Send Report Via:	Fax Email Portal			
Fax #/Email:				
SEND ADDITIONAL R	EPORT COPIES TO (option	al)		
Provider Name		GeneDx Aco	ct#	
Fax #/Email:				
	ICD-10-C	M CODE	s	
ICD-10-CM Codes to	support all test(s) ordere	ed		
Clinical Diagnosis				Age of Onset
	PAYMENT OPTIO	NS (Sele	ect One,)
O INSURANCE BILL	Patient Status			
Select all that apply	Is this individual currently	a Hospital In	patient? (Yes O No
☐ Commercial ☐ Medicaid	Name of Insurance Carr	ier	Insurance	D#:
Medicare	Relationship to Insured			
□Tricare	OSelf OSpouse OCh	nild Onthe	r•	

	PAYMENT OPTIONS (Select One)				
O INSURANCE BILL Select all that apply	Patient Status Is this individual cu	rrently a Hospital In	patient? () Yes) No	
☐ Commercial ☐ Medicaid	Name of Insurance	e Carrier	Insurance ID#:		
☐ Medicare ☐ Tricare	Relationship to Insured OSelf Ospouse Ochild Oother:				
☐ CHAMPVA	Policy Holder's Na	me	Policy Holder's Date	of Birth	
FOR ALL INSURANCE PROVIDE FRONT AND BACK COPY OF	Referral/Prior Auth (please attach)	norization #	Hold test for cost estimate a contact patient if estimate		
CARD(S)	Secondary Insura	псе Туре:	is >\$250 (for in-network/ contracted commercial insurance only)		
	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 se			ill submit a	
	Authorized Patient/Guardian Signature				
O INSTITUTIONAL BILL	GeneDx Account #	#	Place Sticker/St	amp Horo	
	Hospital/Lab Nam	е	riace Sticker/St	апр пеге	

Signature of Ordering Provider

Date



First Name Last Name Date of Birth

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 r	not a substitute for submitting clinica	al records.				
Pre/Perinatal History	□Hypertelorism	☐ Hepatomegaly/Splenomegaly	☐ Renal agenesis			
☐ Cystic hygroma	□ Low set ears	☐ Hepatic fibrosis	☐ Renal insufficiency			
☐ Growth delay	☐ Macrocephaly	☐ Inflammatory bowel disease	□ Renal tubular dysfunction/acidosis			
☐ Increased nuchal translucency	☐ Microcephaly	☐ Intestinal perforation				
☐ Intrauterine growth retardation	☐ Micrognathia	☐ Intrahepatic biliary atresia	Metabolic/Mitochondrial			
☐ Nonimmune hydrops fetalis	Retrognathia	Laryngomalacia	(Attach relevant lab reports/values)			
☐ Multiple prenatal fractures	☐ Short neck	Nausea	Abnormal LDL-Cholesterol levels			
Oligohydramnios	Synophrys	Pancreatitis	Abnormal newborn screen result:			
□Polyhydramnios	☐ Wide nasal bridge	Pyloric stenosis	Alternative and a leaves at AA are south			
Structural Drain Abnormalities	Eva Visian Almannariitias	☐ Tracheoesophageal fistula	Abnormal plasma AA result:			
Structural Brain Abnormalities	Eye/Vision Abnormalities	☐ Vomiting	Abnormal urine OA result:			
☐ Abnormal myelination ☐ Abnormality of basal ganglia	☐ Abnormality of vision ☐ Anophthalmia	Musculoskeletal	☐ Elevated CPK: ☐ Elevated hepatic transaminases			
Abnormality of the corpus callosum	☐ Blue sclerae	Abnormal connective tissue	☐ Hyperglycemia			
Aplasia/hypoplasia of cerebellum	☐ Cataracts	Abnormality of bone mineral density	☐ Hypoglycemia			
Apidsid/hypopidsid of cerebellari	Coloboma	Abnormality of the ribs	☐ Hypokalemia			
Holoprosencephaly	☐ Ectopia lentis	Abnormality of the upper limb	☐ Increased serum pyruvate			
Hydrocephalus	External ophthalmoplegia	Bowing of the long bones	☐ Lactic acidosis			
Lissencephaly	☐ Microphthalmia	☐ Bruising susceptibility	☐ Vitamin D deficiency			
☐ Molar tooth sign on MRI	☐ Myopia	☐ Clinodactyly	- Vicariii i B delicione)			
☐ Ventriculomegaly	Nystagmus	☐ Ectrodactyly	Endocrine			
_ remandation again,	Photophobia	Fractures of the long bones	☐ Amenorrhea			
Developmental/Behavioral	Ptosis	Hyperostosis	□ BMI:			
☐ Absent speech	☐ Strabismus	☐ Hypertonia	Delayed puberty			
Attention deficit hyperactivity disorder	_	☐ Hypotonia	☐ Diabetes insipidus			
Autistic behavior	Hearing Impairment	Limb joint contracture	☐ Diabetes mellitus			
☐ Behavioral abnormality	☐ Conductive hearing impairment	Overgrowth %ile:	☐ Ectopic calcification			
□ Delayed fine motor development	□ bilateral □ unilateral	☐ Pectus carinatum	☐ Elevated hemoglobin A1c			
☐ Delayed gross motor development	☐ Sensorineural hearing impairment	☐ Pectus excavatum	Goiter			
□ Delayed speech & language	☐ bilateral ☐ unilateral	□ Polydactyly	☐ Hypercalcemia			
development	☐ Hearing impairment, mixed or	☐ Short stature	☐ Hyperthyroidism			
□ Developmental regression	unknown	☐ Skeletal dysplasia	☐ Hypophosphatemia			
☐ Global developmental delay	□ bilateral □ unilateral	☐ Small chest circumference	☐ Hypothyroidism			
Hyperactivity		☐ Syndactyly	Low alkaline phosphatase			
☐ Intellectual disability	Cardiac	TC ratio:	MODY: age of onset			
Obsessive compulsive disorder	Abnormal heart morphology	☐ Thoracic hypoplasia	Pancreatic islet autoantibody			
☐ Specific learning disability	Aortic root dilation	□ Vertebral abnormalities	negativity			
☐ Stereotypy	Arrhythmia	art his	□Rickets			
Manualantani	Atrial septal defect	Skin/Hair	the control of our or the control of our			
Neurological	Cardiomyopathy	Abnormal blistering of the skin	Hematological or Immunological			
Abnormality of nervous system	DCM HCM	Abnormality of hair	☐ Anemia			
Anosmia, congenital	Coarctation of aorta	Abnormality of nail	☐ Bone marrow hypocellularity			
Ataxia	Heart murmur	Angieleratema	☐ Immunodeficiency			
Cerebral palsy	☐ Heterotaxy ☐ Hypertension	☐ Angiokeratoma ☐ Café-au-lait macules				
□ Dystonia □ Encephalopathy	Patent ductus arteriosus	☐ Dry skin	☐ Pancytopenia☐ Recurrent infections			
Epileptic encephalopathy	☐ Tetralogy of Fallot	□ Eczema	Recurrent otitis media			
Familial or sporadic hemiplegic	☐ Ventricular septal defect	☐ Hyperextensible skin	☐ Thrombocytopenia			
migraine	- Voltatodiai ooptaraoloot	Hyperpigmentation of the skin	☐ Thromboembolism			
☐ Focal seizures	Respiratory	☐ Hypertrichosis				
Headaches	Asthma	☐ Hypopigmentation of the skin	Vascular System			
☐ Hyperreflexia	Bronchiectasis	☐ Ichthyosis	Aneurysm			
☐ Infantile spasms	☐ Pneumothorax	Recurrent skin infections	Arterial calcification			
Peripheral neuropathy	Pulmonary fibrosis	☐ Velvety skin (soft skin)	Arterial dissection			
Reduced tendon reflexes	Recurrent upper respiratory infections	☐ Xanthomatosis	Arteriovenous malformation			
☐ Seizures	☐ Respiratory distress		□ Lymphedema			
☐ Sensory neuropathy	Respiratory insufficiency	Genitourinary	☐ Stroke			
□ Spasticity		☐ Abnormal renal biopsy:				
☐ Stroke-like episode(s)	Gastrointestinal		Additional Clinical Findings:			
□Tremors	☐ Abnormality of the liver	☐ Abnormal urine analysis:				
,	☐ Aganglionic megacolon					
Craniofacial/Dysmorphism	☐ Cholestasis	☐ Ambiguous genitalia				
Abnormal facial shape	Congenital diaphragmatic hernia	Chronic kidney disease				
☐ Cleft lip	Constipation	Cryptorchidism				
☐ Cleft palate	Diarrhea	Cystic renal dysplasia				
☐ Craniosynostosis	☐ Duodenal stenosis/atresia	Hydronephrosis				
Downslanted palpebral fissures	Exocrine pancreatic insufficiency	Hypospadias				
	Failure to thrive	☐ Micropenis				
External ear malformation	Feeding difficulties	□ Nephrocalcinosis				
Facial asymmetry	Gastroesophageal reflux	☐ Nephrotic syndrome				
☐ Frontal bossing	Gastrointestinal dysmotility	☐ Nephrolithiasis				
☐ High palate	Gastroschisis	☐ Polycystic kidney disease				



First Name Last Name Date of Birth								
			FAMILY H	HISTORY				
□ No Known Family History	□ No Known Family History □ Pedigree Attached □ Adopted							
Relationship	Maternal	Paternal		Relevant i	History		Age at Dx	
1	0	0						
2	0	0						
3	0	0						
			PDEVIOUS OF	IETIO TEOTINO				
			PREVIOUS GEN					
Personal or family history of g			· · ·	ease complete all field				
Relation to patient (self, sibling, etc	c.), Genetic T	est(s) and R	Result (e.g. positive, nego	ative, etc.). If relative was	tested at GeneDx	, please also provide their a	ccession #:	
If patient or relative(s) were found	to have a n	ooitiyo or \/I	IIC requit on prior tecting	places provide details b	olow.			
Indicate any Variants of Interest‡ v				, piease provide details b	elow.			
Relation (self, sibling, etc.)	Gene	Transcrip	pt# c./p. (SN	V) or exon # (CNV)	Build, c	oordinates (CNV)	Variant of Interest‡?	
1								
2								
3								
Required for sequence variants: gene, c./p., transcript #								
Required for CNVs: gene, transcript #, e		d, coordinates	S					
Abnormal karyotype, FISH, or other	results:							
					.() ()			
‡ For certain tests, GeneDx may be able must be provided <u>in the table above</u> at not be possible to comment upon the p	the time the te	est order is plo	aced. If you do not complete	the table above and check of	off that a previously	identified variant is a variant of	rmation interest, it will	
not be possible to comment apon the p	reserice of up:	serice of the v	variant in the report retrospe	etively. This service is not up	plicable to targeted	variant testing.		
			TARGETED VAR	RIANT TESTING				
Individual to be tested: O Aff		•	OUnaffected/	Asymptomatic				
☐ Known Familial Variant(s) in a N ☐ Known Familial Copy Number V]Confirmation of Variant]Known mtDNA Variant(s	Identified in Research Lai	•	Mosaic Variant Testing* e Billing NOT Accepted; Pati	ont Pill or	
Пкложит ғатпінаг сору матпрег у	ununt(s)		JKHOWITTIIDNA VAHAIIL(S	s) resurig	Institution	al Bill MUST be selected on		
Proband Name		Relo	ationship to Proband		Proband GeneD	x Accession #		
Non-GeneDx Test:								
□Positive control included/will be sent - Positive control is recommended if previous test was performed at another lab. □Positive control not available (caveat language will be included on a negative report)								
VARIANT INFORMATION (please fill out the below information if family member report is not included) Number of Variants:								
Gene Coding DNA (c./m.) Amino Acid (p.) Transcript (NM#)								
Gene Coding DNA (c./m.) Amino Acid (p.) Transcript (NM#)								
COPY NUMBER VARIANT		,	l		<u> </u>	Number of Variants:		
Gene(s)	Exon #	:		Coordinates		Genome Build		
Gene(s)	Exon #	:		Coordinates		Genome Build		



First Name	Last Name	Date of Birth

TEST MENU						
TEST CODE	TEST NAME	TEST CODE	TEST NAME			
DERMATOLOGIC DISORDERS						
□ 714	Birt-Hogg-Dube syndrome (FLCN)	□ 708	Slice — Congenital Ichthyosis			
□ 553	Incontinentia Pigmenti – Full Gene Sequencing and Common Deletion (IKBKG/NEMO)	□ 707	Slice – Epidermolysis Bullosa (EB)			
☐ 713	Hereditary Leiomyomatosis and Renal Cell Cancer (FH sequencing and del/dup analysis)	□ 188	Hermansky-Pudlak Syndrome: Puerto Rican Mutations (HPS1, HPS3 sequencing and del/dup analysis)			
DYSMORPH	OLOGY AND MULTIPLE CONGENITAL ANOMALIES					
□ 962	Neurofibromatosis Type 1 Panel	□ 963	Neurofibromatosis Type 2 Panel			
□ J660	Neurofibromatosis Type 1 (<i>NF1</i> single gene sequencing and del/dup analysis)	□ TA06	Noonan and Comprehensive RASopathies Panel			
ENDOCRINE	DISORDERS					
□ 676	Hypogonadotropic Hypogonadism Panel	□ 674	Maturity-Onset Diabetes of the Young (MODY) Panel			
HEMATOLO	GIC DISORDERS					
☐ TB47	Dyskeratosis Congenita Panel	□ 109	Shwachman-Diamond Syndrome (SBDS gene sequencing)			
□ 107	Dyskeratosis Congenita, Autosomal Dominant (TERC gene seque	encing)				
IMMUNOLO	GIC DISORDERS					
□ т990	Autoimmune Lymphoproliferative Syndrome (ALPS) Panel	☐ TA48	Severe Congenital Neutropenia, Autosomal Dominant (ELANE/ELA2)			
□ ⊤989	☐ T989 Chronic Granulomatous Disease (CGD) Panel					
NEUROLOG	C DISORDERS					
□ 526	Cerebral Cavernous Malformations Panel	□ 552	X-linked Hydrocephalus, X-linked Spastic Paraplegia, Masa, Crash Syndrome (<i>LICAM</i>)			
☐ TB51	Comprehensive Holoprosencephaly Panel					
PULMONAR	Y DISORDERS					
□ т829	Cystic Fibrosis/Congenital Bilateral Absence of The Vas Deferens (<i>CFTR</i> sequencing and del/dup analysis)	☐ TB46	Primary Ciliary Dyskinesia Panel			
RENAL AND	GASTROINTESTINAL DISORDERS					
☐ TG21	Alport Syndrome Panel	☐ TH01	Nephrolithiasis and Nephrocalcinosis Panel			
☐ TG23	Cystic Kidney and Liver Diseases Panel	☐ TG22	Polycystic Kidney Disease Panel			
☐ TG98	Hypokalemia and Related Disorders Panel	☐ TG90	Primary Hyperoxaluria Panel			
☐ TG99	Nephrotic Syndrome/Focal Segmental Glomerulosclerosis Panel					
REPRODUC	TIVE DISORDERS					
□ 522	FMR1-associated Premature Ovarian Failure, CGG Repeat Analysis Only (FMR1)	□ 677	Premature Ovarian Failure Panel			
RHEUMATO	LOGIC DISORDERS					
□ 367	Periodic Fever Syndromes Panel: Familial Hibernian fever/TRAPS; urticaria, NOMID; Cyclic neutropenia; PAPA syndrome; Majeed syn		anean fever; Hyper-IgD syndrome; Muckle Wells/familial cold			

(Continue on the next page)

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.



First Name	Last Name		Date of Birth		
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	TEST MENU	(continued)			
TEST CODE	TEST NAME	TEST CODE	TEST NAME		
SKELETAL D	ISORDERS				
☐ T992	Autosomal Dominant Osteogenesis Imperfecta Panel	☐ TA42	Limb Abnormalities and Reduction Defects Panel		
☐ TA40	Craniosynostosis Panel	☐ J797	Osteogenesis Imperfecta Panel		
☐ TA41	TA41 Ectrodactyly/Split Hand-Split Foot Malformation Panel		Skeletal Dysplasia Panel		
CUSTOM DI	EL/DUP TESTING				
□ 906	Deletion/Duplication Analysis of ONE Nuclear Gene	□ 703	Deletion/Duplication Analysis of 2-20 Nuclear Genes		
Write-in Desired Gene(s) to be Tested:					
WRITE-IN TEST SELECTION					
☐ Test Code	: Test Name:				

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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.



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.

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 regarding 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 to perform genetic testing as ordered; (iv) I understand that, for tests that evaluate data from multiple family members concurrently, test results from these family 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 $\textit{Xpanded}^{\$}$ or Slice tests).	tMG 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						
Signa	gnature of Relative A/Legal Guardian Relative A Relationship to Patient Date					
Signa	ature of Relative B/Legal Guardian Relative B Relationship to Patient Date					