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: () Male () Female	Date of Birth (mm/dd/yy)				
Patient Karyotype (if known):					
Gender Identification (optional):					
Email					
Address					
City	State	Zip Code			
Phone (mobile preferred)	Is this patient deceased? O Yes O No Deceased Date:				

SAMPLE INFORMATION					
Date Sample Collected (mm/dd/yy) Medical Record #					
OBlood OBuccal Swab OOther (spec	ify source):				
Treatment-related RUSH (optional) Reason: O Transplantation O Pregnancy O Surgery O Other:					
Patient has had a blood transfusion () Yes () No Date of Last Transfusion: (2-4 weeks of wait time is required for some testing)					
Patient has had an allogeneic bone marr	ow transplant 🔿 Yes 🔿 No				
Fibroblasts are required for patients who had an allogeneic bone marrow transplant. See www.genedx.com/specimen-requirements for details.					
Patient has a personal history of a hemat	tologic malignancy or disease				
OYes (specify diggnosis)					

If yes, please call the lab to discuss with a genetic counselor the most appropriate sample type.

ORDERING PROVIDER ATTESTATION

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 this box, I confirm that the patient is a New York State resident who gives permission for GeneDx to retain any remaining sample longer than 60 days after testing has been completed.
- Patient Research Opt-Out. By checking this box, I confirm that the patient wishes to opt out of being contacted for research studies.
- Health Information Exchange Opt-in. Check this box if your patient resides in CA, FL, MA, NV, NY, RI, and VT and wishes to opt-in to having their information shared for Health Information Exchange participation.
- Health Information Exchange Opt-out. Check this box if your patient resides in any other US state or territory and wishes to opt-out of participation in Health Information Exchange.

Signature of Ordering Provider

	Date	
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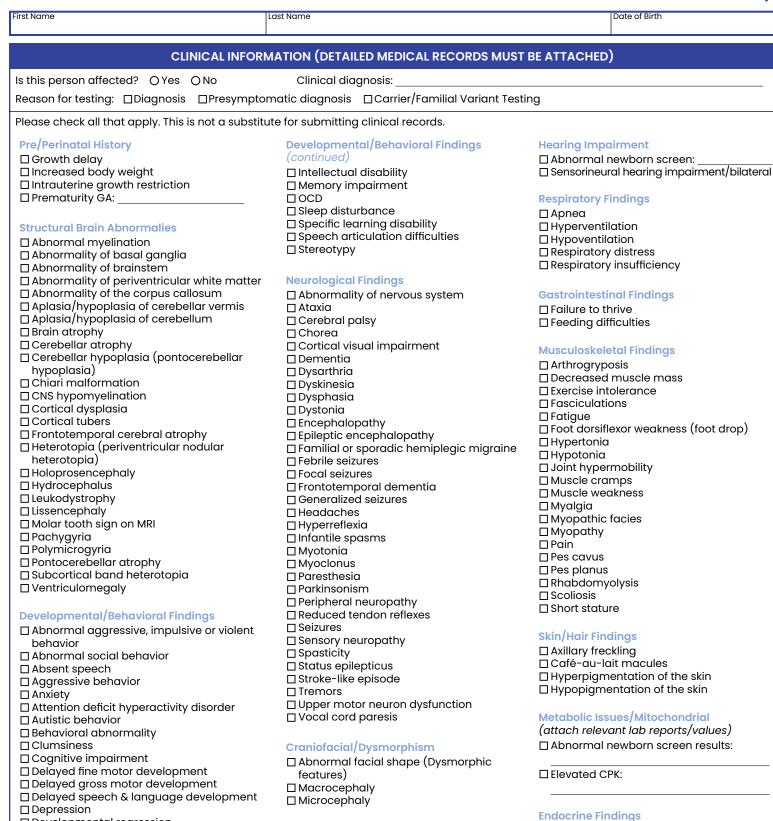
ACCOUNT INFORMATION							
GeneDx Account Number Account Name							
Phone	Fax						
Address							
City	State	Zip Code					
Ordering Provider Name		Role/Title					
NPI Phone Number							
Send Report Via: 🛛 Fax 🗍 Email 🗋 Portal							
Fax #/Email:							
Additional Ordering Provider Name (optional) Role/Title							
NPI							
Send Report Via: 🛛 Fax 🗋 Email 🗋 Portal							
Fax #/Email:							
SEND ADDITIONAL REPORT COPIES TO (optional)							
Provider Name GeneDx Acct#							

Fax #/Email:

ICD-10-CM CODES	
ICD-10-CM Codes to support all test(s) ordered	
Clinical Diagnosis	Age of Onset

	PAYMENT O	PTIONS (Sele	ect One)		
O INSURANCE BILL Select all that apply	Patient Status Is this individual currently a Hospital Inpatient? O Yes O No				
Commercial	Name of Insuranc	ce Carrier	Insurance ID#:		
☐ Medicare ☐ Tricare	Relationship to Ins OSelf OSpouse	oured OChild OOthe	r:		
	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 #	Contact patient if	Hold test for cost estimate and contact patient if estimate	
CARD(S)	Secondary Insura	nce Type:	└─ is >\$250 (for in-network/ contracted commercial insurance only)		
	Insurance Carrier	Insurance ID #	Subscriber Name	Date of Birth	
	Relationship to Ins OSelf OSpouse	ured OChild OOthe	r:		
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.				
	Authorized Patient/Guardian Signature				
	GeneDx Account #	ŧ			
	Hospital/Lab Name		Place Sticker/Stamp Here		





- Developmental regression
- Frequent falls
- Gait disturbance
- Global developmental delay
- □ Hyperactivity .
- □ Incoordination

Eye Defects/Vision

Abnormality of vision
 Cataracts
 Nystagmus
 Optic atrophy

Cardiac Findings

Cardiac rhabdomyoma

Other: _

□ Stroke

Delayed puberty

Vascular System

Arteriovenous malformation

Gene

First Name

Last Name

Date of Birth

Gene

also provide their accession #:

FAMILY HISTORY					
🗆 No Known Family History	□ Pe	edigree Att	ached DAdopted		
Relationship	Maternal	Paternal	Relevant History	Age at Dx	
1	0	0			
2	0	0			
3	0	0			

		PREVIOUS GENETIC TESTING
Personal or family history of genetic testing	O No	O Yes (If yes, please complete all fields below)
Relation to patient (self, sibling, etc.), Genetic Test(s) o	and Resu	It (e.g. positive, negative, etc.). If relative was tested at GeneDx, please

If patient or relative(s) were found to have a positive or VUS result on prior testing, please provide details below. Indicate any Variants of Interest[‡] via the checkbox below.

Relation (self, sibling, etc.)	Gene	Transcript #	c./p. (SNV) 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 # <u>OR</u> build, coordinates					
Abnormal karyotype, FISH, or other results:					

‡ For certain tests, GeneDx **may** be able to specifically comment upon the presence or absence of previously identified variant(s) of interest in the report. Complete variant information must be provided in the table above at the time the test order is placed. If you do not complete the table above and check off that a previously identified variant is a variant of interest, it will not be possible to comment upon the presence or absence of the variant in the report retrospectively. This service is not applicable to targeted variant testing.

TARGETED VARIANT TESTING						
Individual to be tested: OAffected	d/Symptomo	atic OUnaffected/	Asymptomatic			
□ Known Familial Variant(s) in a Nuclea			t Identified in Research Lab	Targeted Mosaic Variant Testing*		
🛛 Known Familial Copy Number Varian	t(s)	□Known mtDNA Variant(s) Testing	*Insurance Billing NOT Accepted; Patient Bill or Institutional Bill MUST be selected on page 1		
Proband Name		Relationship to Proband		Proband GeneDx Accession #		
	Non-GeneDx Test: □ Family member test report included (recommended if previous test was performed at another lab) □ 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 f	ill out the belo	w information if family mer	nber 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 Number of Variants:						
Gene(s) Exon #			Coordinates	Genome Build		
Gene(s) Exon #			Coordinates	Genome Build		

First Name

Last Name

Date of Birth

Gene

TEST INAME TEST NAME TEST NAME 190 Chromosomal Microarroy (Microarroy/A) D52 Fragile X Syndrome (<i>FMRI</i> repect analysis) NEURODEVELOPMENTAL DISORDERS AND EPILEPSY Image: A syndrome (<i>FMRI</i> repect analysis) 1305 Autism/D Ponel (eq & del/dup of 103 genes) [729 1305 Autism/D Ponel (eq & del/dup of 104 genes) [720 1305 Autism/D Ponel (eq & del/dup of 144 genes) [720 1400 Hempilegic Migraine Panel (500+ genes, trics preferred) [7127 Angeiman Syndrome/Proder-Will Syndrame Methylation MPA (UR). delation) 1400 Hempilegic Migraine Panel (500+ genes, trics preferred) [1351 Microcepholay Xpandade* Panel (500+ genes, trics preferred) 1516 Comprehenative Enjoin Micromations Rand (seq & del/dup of 103 J.IS1 Microcepholay Xpandade* Panel (500+ genes, trics preferred) 1526 Cerebral Covenous Malformations (MIII, CCM2, PDCDI/D eq & [J.IS3 Lekicolystrophy Xpandad** Panel (500+ genes, trics preferred) 1781 Dementite Panel (1300 * genes, trics preferred) [1482 1782 Anala Xpandad** Panel (1300 * genes, trics preferred) [1483 1784 Dementite Panel (1300 * genes, trics preferred) [1484 <							
CODE LEST NAME CODE TEST NAME 910 Chromosonial Microarray (Microarray (A 352 Proglie X Syndrome (<i>Ir,Mi</i> repeat analysis) 1395 Autism/ID Pranel (seq & del/dup of 130 genes) 2729 Retr/Angelman Related Disorders Panel (seq & del/dup of 25 genes & methydion MIRA) 1395 Comprehensive Epilepay Panel (seq & del/dup of 14 genes) 1720 Interview Steries Panel (SCI & TSC2 et a & del/dup of 25 genes & methydion MIRA) 1400 Hemiplegic Migraine Panel (seq & del/dup of 4 genes) 1721 Angeiman Syndrome (Prader-Willi Syndrome Methylation MLPA (UPD, deletion) 1401 Hemiplegic Migraine Panel (seq & del/dup of 103 genes) 1531 Microcephaly Xpanded* Panel (800+ genes, trics preferred) 1538 Comprehensive Brain Malformations Panel (seq & del/dup of 103 genes) 1545 Heinoted Hydrocephalus X-iniked Spassic Parapisagi (ANSA/CRASH MOVEMENT DISORDERS 2 J853 leukodystrophy Xpanded* Panel (300+ genes, trics preferred) 1742 17420 Natio Xpandad* Ponel (soq or hy of 11 genes, for patients 18 years and tespes 1585 Spinocerebellar Atakia Repeat Expansion Analysis (ATAN, ATXNZ, AT		TES	T MENU				
NEURODEVELOPMENTAL DISORDERS AND EPILEPSY Rett/Angeiman Releted Disorders Panel (seq & del/dup of 103 genes) 729 Rett/Angeiman Releted Disorders Panel (seq & del/dup of 25 genes & mothylotion MLPA) 1523 Comprehensive Epilepsy Panel (seq & del/dup of 144 genes) 730 Tuberous Sciencis Panel (TSC) & TSC2 seq & del/dup) 1403 Hemiplegic Migraine Panel (seq & del/dup of 4 genes) Interous Sciencis Panel (R00+ genes, trics preferred) Interous Sciencis Panel (R00+ genes, trics preferred) 1403 Hemiplegic Migraine Panel (seq & del/dup of 4 genes) Comprehensive Brain Motiformations (RRT, CCM2, POCDID seq & IJBS Microcepholy Xpanded* Panel (800+ genes, trics preferred) 1544 Dementia Panel (seq only of 11 genes, for patients 18 years and genes) IS52 X-iniked hydrocepholy Xpanded* Panel (800+ genes, trics preferred) 1762 Ataxia Xpanded* Panel (seq & del/dup of 200 yeas, trics preferred) IS52 X-iniked hydrocepholy Xpanded* Panel (800+ genes, trics preferred) 1762 Ataxia Xpanded* Panel (seq & del/dup of 200 yeas, trics preferred) IS52 X-iniked hydrocepholy Xpanded* Panel (800+ genes, trics preferred) 1762 Ataxia Xpanded* Panel (seq & del/dup of 200 yeas, trics preferred) ITH8 Y=intoarchiter Ataxia Report Analysis (ATXN, ATXN2, A		TEST NAME		TEST NAME			
1395 Autism/ID Panel (seq & del/dup of 103 genes) [729] Rett/Angelman Related Disorders Panel (seq & del/dup of 25 genes & methylaction MLPA) 523 Comprehensive Epilepsy Panel (seq & del/dup of 144 genes) [720] Tubercous Sciences Panel (75C1 & TSC2 seq & del/dup) 100 Hemplegic Migraine Panel (seq & del/dup of 4 genes) [7127] Angelman Syndrome/Prader-Willi Syndrome Methylation MLPA 100 Hemplegic Migraine Panel (seq & del/dup of 4 genes) [7127] Angelman Syndrome/Prader-Willi Syndrome Methylation MLPA 1010 Hemplegic Migraine Panel (seq & del/dup of 4 genes) [7127] Angelman Syndrome/Prader-Willi Syndrome Methylation MLPA 1010 Hemplegic Migraine Panel (seq & del/dup of 103 genes) [1518] Microcepholy Xpanded* Panel (800+ genes, trios preferred) 1614 Comprehentice Panel (seq only of 11 genes, for patients 18 years and older) [152] X-linked Hydrocephalus/X-linked Spostic Paraplegia/MASA/CRASH 1742 Ataxia Xpanded* Panel (1300+ genes, trios preferred) [1184] [1184] Spinocerebelar Ataxia Repeat Expansion Analysis (ATXNI atAXNI, ATXNI,	910	Chromosomal Microarray (MicroarrayDx)	□ 522	Fragile X Syndrome (FMR1 repeat analysis)			
International methylation MPA 623 Comprehensive tpilepsy Panel (sag & del/dup of 14 genes) 730 Tuberous Sciencesis Panel (TSCL & TSC2 seg & del/dup) 921 tpi Xpanel (sig a de del/dup of 4 genes) 730 Tuberous Sciencesis Panel (TSCL & TSC2 seg & del/dup) 1400 Herriplegic Migraine Panel (seg & del/dup of 4 genes) 730 Tuberous Sciencesis Panel (SCL & TSC2 seg & del/dup) 1400 Herriplegic Migraine Panel (seg & del/dup of 103 genes) 1272 Angeimon Syndrome/Praded* Panel (800+ genes, trics preferred) 1536 Comprehensive Brain Kennel (seg end of the panes) 1351 Microcopholus/X-linked Spacia Panes, trics preferred) 1740 Herriplegic Migraine Panel (seg end) of 11 genes, for patients 18 years and edder) 1482 Spinocerebelar Atasia Repect Expansion Analysis (ATXNI, ATXNZ, ATXNI, ATXNI, ATXNZ, ATXNI, ATXNZ, ATXNI, ATXNZ, ATXNI, ATXNZ, ATXNI, ATXNZ, ATXNI, ATXNI, ATXNZ, ATXNI, ATXNZ, ATXNI, ATXNZ, ATXNI, ATXNZ, ATXNI, ATXNZ, ATXNI, ATXNI, ATXNZ, ATXNI, ATXNI, ATXNZ, ATXNI, ATXNZ, ATXNI, ATXNI, ATXNZ, ATXNI, ATXNZ, ATXNI, ATXNZ, ATXNI, ATXNI, ATXNZ, ATXNI, ATXNI, ATXNI, ATXNZ, ATXNI, ATXNI, ATXNZ, ATXNI, ATXNZ, ATXNI, ATXNZ, ATXNI, ATXN	NEUROD	EVELOPMENTAL DISORDERS AND EPILEPSY					
B21 Epi Xpanded* Panel (1300+ genes, trios preferred) T122 Angeiman Syndrome/Prader-Willi Syndrome Methylation MLPA (VPD, deletion) T400 Herniplegic Migraine Panel (seq & del/dup of 4 genes) Comprehensive Brain Malformatione Panel (seq & del/dup of 103 ganes) J511 Microcephaly Xpanded* Panel (800+ genes, trios preferred) B526 Cemptehensive Brain Malformatione Renel (seq & del/dup of 103 ganes) J553 Leukodystrophy Xpanded* Panel (800+ genes, trios preferred) B526 Cemptehensive Brain Malformations (KRITI, CCM2, PDCDI0 seq & del/dup) J1553 Leukodystrophy Xpanded* Panel (300+ genes, trios preferred) B526 Cemptehensive Brain Malformations (KRITI, CCM2, PDCDI0 seq & del/dup) Spinocerebeliar Ataxia Repect Expansion Analysis (ATXMI, ATXM2, ATXM3, ATXM7, ATXM2, CAMMA Repect) B1762 Moxix Apanded* Panel (1300+ genes, trios preferred) B484 Spinocerebeliar Ataxia Repect Expansion Analysis (ATXMI, aTXM2, ATXM3, ATXM7, ATXM2, CAMMA Repect) B1762 Moxix Apanded* Panel (1300+ genes, trios preferred) B1483 Spinocerebeliar Ataxia Type 3 Repect Analysis (ATXMI, aTXM2, ATXM3, ATXM7, ATXM3, ATXM2, ATXM2	☐ T395	Autism/ID Panel (seq & del/dup of 103 genes)	□ 729				
Interview (UPb, deletion) Interview (UPb, deletion) Interview (UPb, deletion) CNS MALFORMATIONS AND DISORDERS (UPb, deletion) Is81 Comprehensive Brain Multormations Panel (seq & del/dup of 103 genes) J511 Microcephaly Xpanded* Panel (800+ genes, trios preferred) Is52 Comprehensive Brain Multormations (<i>KRIT, CCM2, PDCDI</i>) seq & IJ853 Leukodystrophy Xpanded* Panel (800+ genes, trios preferred) IT844 Dementia Panel (seq only of 11 genes, for potients 18 years and del/dup) IS52 Spinocereballar Ataxia Repect Diporsion Analysis (ATXNI, ATXN2, ATXN3, ATXN7, ATXN8, CACNAIA repect) I/762 Ataxia Xpanded* Panel (1300+ genes, trios preferred) ITH83 Spinocereballar Ataxia Repect Analysis (ATXNI, ATXN2, ATXN3, ATXN7, ATXN8, CACNAIA repect) I/762 Ataxia Xpanded* Panel (1300+ genes, trios preferred) ITH83 Spinocereballar Ataxia Type Repect Analysis (ATXNI, epect) IH97 Demetarianson Branel (seq & del/dup of 83 genes) ITH84 Spinocereballar Ataxia Type Repect Analysis (ATXNI repect) IH98 Friedreich Ataxia Repect Analysis (FXN repect) IH98 Spinocereballar Ataxia Type Repect Analysis (ATXNI repect) IH94 Friedreich Ataxia Repect Analysis (FXN repect) IH98 Spinocereballar	□ 523	Comprehensive Epilepsy Panel (seq & del/dup of 144 genes)	□ 730	Tuberous Sclerosis Panel (TSC1 & TSC2 seq & del/dup)			
CNS MALFORMATIONS AND DISORDERS B81 Comprehensive Brain Malformations Panel (seq & del/dup of 103 genes) J511 Microcepholy Xpanded* Panel (800+ genes, trios preferred) J526 Cerebral Covernous Malformations (KNIT, CCM2, PDCDI0 seq & del/dup) J553 Leukodystrophy Xpanded* Panel (800+ genes, trios preferred) IT844 Dementia Panel (seq only of 1] genes, for patients 18 years and del/dup) IS52	921	Epi Xpanded® Panel (1300+ genes, trios preferred)	□ TJ27				
Genes Comprehensive Brain Malformations Panel (seq & del/dup of 103 genes) J511 Microcephaly Xpanded* Panel (800+ genes, trios preferred) del/dup) E326 Comprehensive Brain Malformations (KRIT), CCM2, PDCDID seq & del/dup) J453 Leukodystrophy Xpanded* Panel (800+ genes, trios preferred) T1844 Dementia Panel (seq only of 11 genes, for patients 18 years and older) T52 X-linked Hydrocephalus/X-linked Spastic Paraplegia/MASA/CRASH syntame J722 Ataxia Xpanded* Panel (1300+ genes, trios preferred) TH85 Spinocerebellar Ataxia Repeat Expansion Analysis (ATXM, ATXM2, ATXM3, ATXM7, ATXM3, CACMAIA repeat) TH84 Spinocerebellar Ataxia Type 1 Repeat Analysis (ATXM repeat) TH84 Spinocerebellar Ataxia Type 1 Repeat Analysis (ATXM repeat) TH84 TH84 Spinocerebellar Ataxia Type 1 Repeat Analysis (ATXM repeat) TH85 Filedreich Ataxia Repeat Analysis (ATXM repeat) TH85 Filedreich Ataxia Type 3 Repeat Analysis (ATXM repeat) TH85 Filedreich Ataxia Type 6 Repeat Analysis (ATXM repeat) TH85 Filedreich Ataxia Type 6 Repeat Analysis (ATXM repeat) TH85 Filedreich Ataxia Sequencing & 61/dup (FXM repeat)	□ T400	Hemiplegic Migraine Panel (seq & del/dup of 4 genes)					
genes) Labert Laber Labert Labert Labert Labert Labert Labert	CNS MA	LFORMATIONS AND DISORDERS					
del/dup) del/dup) del/dup) TH44 Dementia Panel (seq only of II genes, for patients IB years and older) 552 X-linked Hydrocephalus/X-linked Spastic Paraplegia/MASA/CRASH syndrome (LICAM seq & del/dup) MOVEMENT DISORDERS atxaia Xpanded* Panel (1300+ genes, trios preferred) TH83 Spinocerebellar Ataxia Repeat Xpansion Analysis (ATXNI, ATXNZ, ATXNZ, CACNAIA repeat) TH97 Dentatorubral-Pallidoluysian Atrophy Repeat Analysis (ATXNI repeat) TH83 Spinocerebellar Ataxia Type 1 Repeat Analysis (ATXNI repeat) TH402 Dystonia and Parkinsonism Panel (seq & del/dup of 103 genes) TH85 Spinocerebellar Ataxia Type 3 Repeat Analysis (ATXN7 repeat) TH402 Dystonia and Parkinsonism Panel (seq & del/dup of 44 genes) TH86 Spinocerebellar Ataxia Type 8 Repeat Analysis (ATXN7 repeat) TH84 Friedreich Ataxia Repeat Analysis (FXN repeat) TH88 Spinocerebellar Ataxia Type 8 Repeat Analysis (ATXN7 repeat) TH84 Priedreich Ataxia Repeat Analysis (FXN repeat) TH89 Spinocerebellar Ataxia Type 8 Repeat Analysis (ATXN7 repeat) TH84 Priedreich Ataxia Repeat Analysis (FXN repeat) TH89 Spinocerebellar Ataxia Type 8 Repeat Analysis (ATXN7 repeat) TH84 Priedreich Ataxia Repeat Analysis (FXN repeat) TH89 Spinocerebellar Ataxia Type 8 Repeat Analysis (ATXN7 repeat) T	691		🗆 J511	Microcephaly Xpanded® Panel (800+ genes, trios preferred)			
older) Syndrome (UCAM seq & del/dup) MOVEMENT DISORDERS J762 Ataxia Xpanded* Panel (1300+ genes, trios preferred) TH97 Dentatorubral-Pallidoluysian Atrophy Repeat Analysis (ATNI repeat) TH92 Dystonia and Parkinsonism Panel (seq & del/dup of 103 genes) T402 Dystonia Panel (seq & del/dup of 83 genes) T401 Parkinson Disease Panel (seq & del/dup of 43 genes) TH85 Friedreich Ataxia Repeat Analysis (ATNN repeat) TH85 TH95 Friedreich Ataxia Sepeat Analysis (ATNN repeat) TH94 Friedreich Ataxia Sepeat Analysis (ATNN repeat) TH95 Friedreich Ataxia Sequencing & Del/Dup (FXN seq & del/dup) T112 Spinocerebellar Ataxia Type 8 Repeat Analysis (ATNN repeat) TH85 Spinocerebellar Ataxia Type 8 Repeat Analysis (ATNN repeat) T112 Spinocerebellar Ataxia Type 8 Repeat Analysis (ATNN repeat) T112 Spinocerebellar Ataxia Type 8 Repeat Analysis (ATNN repeat) D1143 Arriver Analysis (ATNN repeat) T112 Spinocerebellar Ataxia Type 8 Repeat Analysis (ATNN repeat) D1189 Spinocerebellar Ataxia Type 8 Repeat Analysis (ATNN repeat) D1199 Arrystanght Lateral Sciences/Frontotemporal Lobar Degeneration (CBorT2 repeat an	526		□ J853	Leukodystrophy Xpanded® Panel (300+ genes, trios preferred)			
J762 Ataxia Xpanded* Panel (1300+ genes, trios preferred) IH87 J762 Ataxia Xpanded* Panel (1300+ genes, trios preferred) IH87 ITH97 Dentatorubral-Pallidoluysian Atrophy Repeat Analysis (ATNI repeat) IH84 IT402 Dystonia and Parkinsonism Panel (seq & del/dup of 103 genes) IH85 IT403 TA01 Parkinson Disease Panel (seq & del/dup of 44 genes) IH85 IT4197 Friedreich Ataxia Repeat Analysis (ATNN repeat) IH85 IT4198 Spinocerebellar Ataxia Type 3 Repeat Analysis (ATNN repeat) IT419 Friedreich Ataxia Repeat Analysis (FXN repeat) IH85 IT419 Friedreich Ataxia and Related Disorders Panel (seq & del/dup) ITK79 Amyotrophic Lateral Sclerosis/Frontotemporal Lobar Degeneration (C90rf72 repeat analysis, for patients 18 years and older) ITK79 J805 Arthrogryposis Panel (seq & del/dup of 90 genes) IT89 IT404 Amyotrophic Lateral Sclerosis/Frontotemporal Lobar Degeneration (C90rf72 repeat analysis, for patients 18 years and older) IT89 J805 Arthrogryposis Panel (seq & del/dup of 90 genes) IT88 J818 Myotonic Dystrophy 1 (DMI) (DMPK repeat analysis) IT89 J819 Order of Reliex Testing: IT819 Myotonic Dystrophy 2 (DM	□ T844		□ 552	X-linked Hydrocephalus/X-linked Spastic Paraplegia/MASA/CRASH Syndrome (LICAM seq & del/dup)			
Image: Hard Control - Pollidoluysian Atrophy Repeat Analysis (ATNI repeat) Image: ATXN3,	MOVEM	ENT DISORDERS					
□ H97 Dentatorubral-Pallidoluysian Atrophy Repeat Analysis (ATNI repeat) □ H84 Spinocerebellar Ataxia Type 1 Repeat Analysis (ATXNI repeat) □ T402 Dystonia and Parkinsonism Panel (seq & del/dup of 103 genes) □ T403 Dystonia Panel (seq & del/dup of 83 genes) □ H86 Spinocerebellar Ataxia Type 2 Repeat Analysis (ATXN3 repeat) □ TH95 Friedreich Ataxia Repeat Analysis (FXN repeat) □ H86 Spinocerebellar Ataxia Type 7 Repeat Analysis (ATXN3 repeat) □ TH94 Friedreich Ataxia Repeat Analysis (FXN repeat) □ H86 Spinocerebellar Ataxia Type 7 Repeat Analysis (ATXN3 repeat) □ TH94 Friedreich Ataxia and Related Disorders Panel (seq & del/dup) □ TK79 Xpanded* Adult Movement Disorders Panel (500+ genes, trio preferred) □ H86 Spinocerebellar Ataxia Type 7 Repeat Analysis (ATXN3 repeat) □ TH89 Spinocerebellar Ataxia Type 7 Repeat Analysis (ATXN3 repeat) □ TH94 Friedreich Ataxia and Related Disorders Panel (seq & del/dup) □ TK79 Xpanded* Adult Movement Disorders Panel (500+ genes, trio preferred) □ H89 Amyotrophic Lateral Sclerosis/Frontotemporal Lobar Degeneration C9orf72 repeat analysis, for patients 18 years and older) □ TK79 Motonic Dystrophy 1 (DM1) (DMPK repeat analysis) 0 □ T404 Amyotrophic Lateral Sclerosis/Frontotemporal Lobar Degeneration Panel (seq & del/dup of 24 genes, for patients 18 years and) older) □ 818 <td< td=""><td>🔲 J762</td><td>Ataxia Xpanded® Panel (1300+ genes, trios preferred)</td><td>□ ТН83</td><td></td></td<>	🔲 J762	Ataxia Xpanded® Panel (1300+ genes, trios preferred)	□ ТН83				
T402 Dystonia and Parkinsonism Panel (seq & del/dup of 103 genes)	☐ TH97			TH84 Spinocerebellar Ataxia Type 1 Repeat Analysis (ATXN1 repeat)			
Image:	🗖 T402						
TH95 Friedreich Ataxia Repeat Analysis (F/N repeat) TH98 Spinocerebeliar Ataxia type / Repeat Analysis (ATXN repeat) TH94 Friedreich Ataxia Sequencing & Del/Dup (FXN seq & del/dup) TH98 Spinocerebeliar Ataxia Type 8 Repeat Analysis (ATXN repeat) T112 Spinocerebeliar Ataxia and Related Disorders Panel (seq & del/dup) TK79 Xpanded* Adult Movement Disorders Panel (500+ genes, trio preferred) NEUROMUSCULAR DISORDERS TK9 Xpanded* Adult Movement Disorders Panel (500+ genes, trio preferred) J805 Amyotrophic Lateral Sclerosis/Frontotemporal Lobar Degeneration (C9orf72 repeat analysis, for patients 18 years and older) T37 T404 Amyotrophic Lateral Sclerosis/Frontotemporal Lobar Degeneration Panel (seq & del/dup of 24 genes, for patients 18 years and older) B18 Myotonic Dystrophy 1 (DM1) (DMPK repeat analysis) Order of Reflex Testing: B18 Myotonic Dystrophy 2 (DM2) (CNBP repeat analysis) T404 Arthrogryposis Panel (seq & del/dup of 90 genes) TG82 Myotonia Panel (CNBP and DMPK repeat analysis, seq & del/dup of 8 genes) T42 CMTIA/HNPP (PMP22 del/dup) T404 Strongenetation The panel (seq & del/dup of 90 genes) T682 Neuromuscular Disorders Panel (I15 genes) Seq & del/dup of 8 genes) T42 CMTIA/HNPP (PMP22 del/dup) T433 Oculopharyngeal Muscular Dystrophy (PA							
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T112 Spinocerebellar Ataxia and Related Disorders Panel (seq & del/ dup of 56 genes) TK79 Xpanded* Adult Movement Disorders Panel (500+ genes, trio preferred) NEUROMUSCULAR DISORDERS 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 Degeneration Panel (seq & del/dup of 24 genes, for patients 18 years and older) 818 Myotonic Dystrophy 1 (DM1) (DMPK repeat analysis) Order of Reflex Testing: 818 Myotonic Dystrophy 2 (DM2) (CNBP repeat analysis) Order of Reflex Testing: Myotonia Panel (CNBP and DMPK repeat analysis, seq & del/dup of 90 genes) T464 Arthrogryposis Panel (seq & del/dup of 90 genes) TG82 Myotonia Panel (CNBP and DMPK repeat analysis, seq & del/dup of 8 genes) Seq ens) T469 Arthrogryposis Panel (seq & del/dup of 90 genes) TG82 Myotonia Panel (CNBP and DMPK repeat analysis, seq & del/dup of 8 genes) Seq ens) T677 Congenital Hypotonia Xpanded* Panel (1400+ genes; trios preferred) B89 Neuromuscular Disorders Panel (115 genes) B80 Limb-Girdle Muscular Dystrophy Panel GD1007 Duchenne/Becker MD (DMD seq & del/dup) TG81 Periodic Paralysis Panel (seq & del/dup of 9 genes)	☐ TH95			☐ TH89 Spinocerebellar Ataxia Type 8 Repeat Analysis (<i>ATXN8</i> repeat)at)			
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Degeneration (C9orf72 repeat analysis, for patients 18 years and older) Image: Construction of the c	NEURON	IUSCULAR DISORDERS					
Degeneration Panel (seq & del/dup of 24 genes, for patients 18 Image: Constraint of the second o	☐ J805	Degeneration (C9orf72 repeat analysis, for patients 18 years and	737	Hereditary Neuropathy Panel (seq & del/dup of 89 genes)			
years and older) Image: Sand	□ T404	Amyotrophic Lateral Sclerosis/Frontotemporal Lobar	818	Myotonic Dystrophy 1 (DM1) (<i>DMPK</i> repeat analysis)			
Image: Activate J805, if non-diagnostic activate T404 Image: Activate J805, if non-diagnostic activate T404 Image: T680 Arthrogryposis Panel (seq & del/dup of 90 genes) Image: T682 Myotonia Panel (CNBP and DMPK repeat analysis, seq & del/dup of 8 genes) Image: T742 CMTIA/HNPP (PMP22 del/dup) Image: T743 Oculopharyngeal Muscular Dystrophy (PABPNI repeat analysis) Image: T677 Congenital Hypotonia Xpanded® Panel (1400+ genes; trios preferred) Image: T688 Neuromuscular Disorders Panel (115 genes) Image: T677 Duchenne/Becker MD (DMD seq & del/dup) Image: T688 Periodic Paralysis Panel (seq & del/dup of 9 genes)			□ 819	Myotonic Dystrophy 2 (DM2) (<i>CNBP</i> repeat analysis)			
TG80 Arthrogryposis Panel (seq & del/dup of 90 genes) TG82 Myotonia Panel (CNBP and DMPK repeat analysis, seq & del/dup of 8 genes) 742 CMTIA/HNPP (PMP22 del/dup) 743 Oculopharyngeal Muscular Dystrophy (PABPNI repeat analysis) TG77 Congenital Hypotonia Xpanded® Panel (1400+ genes; trios preferred) 889 Neuromuscular Disorders Panel (115 genes) G01007 Duchenne/Becker MD (DMD seq & del/dup) TG81 Periodic Paralysis Panel (seq & del/dup of 9 genes)		Order of Reflex Testing:					
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preferred) □ 890 Limb-Girdle Muscular Dystrophy Panel □ GD1007 Duchenne/Becker MD (DMD seq & del/dup) □ TG81 Periodic Paralysis Panel (seq & del/dup of 9 genes)	□ 742	CMTIA/HNPP (<i>PMP22</i> del/dup)	□ 743	Oculopharyngeal Muscular Dystrophy (PABPN1 repeat analysis)			
	☐ TG77		889				
820 Spinal & Bulbar Muscular Atrophy (AR repeat analysis) T789 SMN1/2 Dosage Analysis	GD1007	Duchenne/Becker MD (<i>DMD</i> seq & del/dup)	□ TG81	Periodic Paralysis Panel (seq & del/dup of 9 genes)			
	□ 820	Spinal & Bulbar Muscular Atrophy (AR repeat analysis)	□ T789	SMN1/2 Dosage Analysis			

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

Gene

	TEST MENU (continued)							
TEST CODE	TEST NAME	TEST CODE	TEST NAME					
ΜΙΤΟΟ	HONDRIAL DISORDERS							
615	Combined Mito Genome Plus Mito Focused Nuclear Gene Panel	□ TH12	Leber Hereditary Optic Neuropathy (LHON) Panel					
□ 554	Full sequence analysis and deletion testing of the mitochondrial genome	e (not a tric	based test)					
NEURC	METABOLIC DISORDERS							
🗆 J976	Creatine Deficiency Syndromes Panel (seq & del/dup of 3 genes)	□ тно8	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)					
NEURC	FIBROMATOSIS							
961 Comprehensive NF Panel: NF1, SPRED1, NF2 and SMARCB1 sequencing and deletion/duplication testing								
962	NFI Panel: NFI and SPREDI sequencing and deletion/duplication testin	g						
963	NF2 Panel: LZTR1, NF2 and SMARCB1 sequencing and deletion/duplicat	ion testing						
TA06 Reflex to Noonan Syndrome and RASopathies panel (sequencing of 25 genes) if 962 is non-diagnostic								
CUSTOM DEL/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	Test Code: Test Name:							

FAMILY MEMBER FOR PANEL TESTING OPTION

NO SEPARATE REPORT, ADDITIONAL SAMPLES MUST BE RECEIVED WITHIN 3 WEEKS OF PROBAND SAMPLE. See Test Menu page for proband test selection.							
☐ J767 ☐ TG86 ☐ 923 ☐ 725	Ataxia <i>Xpanded®</i> , Family member testing Congenital Hypotonia <i>Xpanded®</i> , Family member testing Epi <i>Xpanded®</i> , Family member testing Chromosomal Microarray Parental Testing		□ J854 □ J513 □ ТК80	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			
Biological Mother				O At GeneDx (Accession #:)			
				O Not available O To be sent within 3 weeks			
	First Name	Last Name	DOB	O Asymptomatic O Symptomatic			
Biological Father				O At GeneDx (Accession #:)			
				O Not available O To be sent within 3 weeks			
	Relationship to Proband						
Other	First Name	Last Name	DOB	O Asymptomatic O Symptomatic			
Biological Relative				O At GeneDx (Accession #:)			
				O Not available O To be sent within 3 weeks			

DID YOU REMEMBER TO...?

□ Label specimen tube appropriately with TWO identifiers

 $\hfill\square$ 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. Variant of Uncertain Significance (VUS): 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 deidentified 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.

Gene

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), 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.

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 ACMG secondary findings (Full Exome Sequencing and Genome Sequencing Tests ONLY; not for *Xpanded®* or Slice 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 contacted 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.

Signature of Patient/Legal Guardian (required)	Date	
Signature of Relative A/Legal Guardian	Relative A Relationship to Patient	Date
Signature of Relative B/Legal Guardian	Relative B Relationship to Patient	Date

Gene