HEREDITARY CANCER TEST REQUISITION FORM



Age of Onset

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	
ati	Assigned at Birth: Male Female ent Karyotype (if known): der Identification (optional):	Date of Birth (mm/dd,	/yy)
Add	Iress		
City	,	State	Zip Code
ho	ne (mobile preferred)	Is this patient decease Deceased Date:	ed? O Yes O No
at	SAMPLE INF e Sample Collected (mm/dd/yy)	FORMATION Medical Record #	
	e sample soliceted (minidally)	medical Record #	
_	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	4 weeks of wait time is required for some te	esting)	
ibr	ient has had an allogeneic bone marrow to oblasts are required for patients who had a www.genedx.com/specimen-requirement	an allogeneic bone man	
_	ient has a personal history of a hematolo	gic malignancy or dise	_
	/es (specify diagnosis) s, please call the lab to discuss with a genet	tic counselor the most a	ONo
, -	o, produce can are laz to allocate mar a geriot	ao coursolor are rricot ap	ppropriate earripie type.
	ORDERING PROVI	DER ATTESTATIC)N
Ge aut Rec tres part any tes dia reir pro Ge pro and fan	signing this form, the ordering provider att neDx to perform the testing indicated; (ii) I thorized by law to order the test(s) request quisition Form ("TRF") are reasonable and a tatment of a disease, illness, impairment, sy ults will determine the patient's medical matient's condition on this date of service; (v) thorized to make decisions for the patient by relatives', when applicable, has been supting, and has consented to undergo genet gnosis codes are indicated to the highest matient in the service of the properties of the	ne/she is the ordering p ted; (iii) any test(s) required; (iii) any test(s) required; or an addition and treatrathe the patient or the individual of the patient of specificity; (vii) go but not limited to fed did will inform the patient he ordering provider and trites regarding the requires; and (ix) the patient or the patient or the patient of the patient or the patient of the patient or the	rovider and is uested on this Test the diagnosis or lisorder; (iv) the test nent decisions of this idual/family member nt"), in addition to regarding genetic nd appropriate he/she will not seek eral healthcare of the same; (viii) d other healthcare uested genetic testing or the individual/
	New York Retention Opt-In. By checking to York State resident who gives permission longer than 60 days after testing has been	for GeneDx to retain an	
	Patient Research Opt-Out. By checking the opt out of being contacted for research s		e patient wishes to
	Health Information Exchange Opt-in. Ch FL, MA, NV, NY, RI, and VT and wishes to op Health Information Exchange participatio	t-in to having their info	

ACCO	OUNT INFORMATIO	N .
GeneDx Account Number	Account Name	•
Phone	Fax	
Address	<u> </u>	
City	State	Zip Code
Ordering Provider Name	l	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 TO	O (optional)	
Provider Name	GeneDx Acct#	
Fax #/Email:	1	
ICI	D-10-CM CODES	
ICD-10-CM Codes to support all test	t(s) ordered	

	PAYMENT O	PTIONS (Sele	ect One)			
O INSURANCE BILL Select all that apply	Patient Status Is this individual cu	ırrently a Hospital In	patient? () Yes) No		
☐ Commercial ☐ Medicaid	Name of Insurance Carrier		Insurance ID#:			
☐ Medicare ☐ Tricare	· '	Relationship to Insured Oself Ospouse Ochild Oother:				
CHAMPVA FOR ALL INSURANCE	Policy Holder's Na	me	Policy Holder's Date	of Birth		
PROVIDE FRONT AND BACK COPY OF	Referral/Prior Authorization # (please attach)		Hold test for cost estimate and contact patient if estimate			
CARD(S)	Secondary Insura	nce Type:	is >\$250 (for in-ne contracted comm 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					
	Authorized Patient/Guardian Signature					
O INSTITUTIONAL BILL	GeneDx Account	#	Place Sticker/Stamp Here			
	Hospital/Lab Name		Fide Sticker/St	ar/stamp Here		

Signature of Ordering Provider

Date

Clinical Diagnosis

HEREDITARY CANCER TEST REQUISITION FORM



First Name Last Name Date of Birth

CLINICAL INFORMATION (DETAILED MEDICAL RECORDS MUST BE ATTACHED)								
	□ No relevant personal history							
DIAGNOSIS	DIAGNOSIS AGE AT DX PATHOLOGY							
□ Breast Cancer		ER PR HER2/neu ☐ Triple Negative ☐ Invasive Lobular ☐ Bilateral ☐ Two Primaries ☐ Invasive Ductal ☐ DCIS ☐ Other Pathology:						
☐ Colorectal Cancer		Location: Right Left Transverse Rectum Pathology:						
□ Ovarian Cancer		□ Epithelial □ Non-epithelial □ Other Pathology:						
☐ GI Polyps		☐ Adenomatous - total #: Location: ☐ Other - total #: ☐ Other Pathology:						
☐ Endometrial Cancer		Pathology:						
□ Hematologic Disease								
□ Prostate Cancer		Gleason Score:						
□ Skin Cancer		□ Melanoma □ Other Pathology:						
☐ Gastric Cancer/Tumor		Pathology:						
□ Endocrine Cancer/ Disease		Pathology:						
□ Renal Cancer/Tumor		Pathology:						
□ Brain Cancer/Tumor		Pathology:						
□ Pancreatic Cancer		Pathology:						
□ Pancreatitis								
☐ Other								
Comments								

(Continue to the next page)

Page 2 of 6

HEREDITARY CANCER TEST REQUISITION FORM



First Name			Last Name					Dat	e of Birth		
			·					·			
				FAMILY	HISTORY						
□ No Known Family	History		Pedigree Atto	ıched	☐ Adopt	ted					
Relationship	Paternal		R	elevant F	listory				Age at Dx		
			0								
	()	0								
	()	0								
			PR	EVIOUS GEI	NETIC TESTING						
Personal or family his	story of genetic	c testing	O No O Ye	es (If yes, ple	ase complete al	II fields be	elow)				
Relation to patient (self, If patient or relative(s) w	vere found to ha	ve a positiv	e or VUS result o					, please c	also provide	e their acc	ession #:
Relationship (self, sibling, etc.)	Interest‡ via the Gene		nscript#	c./p	o. (SNV) or n # (CNV)	Build, c	oordinates (C	·NIV	Variant of Interest?	Туре о	of Variant
(seii, sibiirig, etc.)				exo	II# (CNV)					□Germlin	ne Somatic
											ne Somatic
										□Germlin	ne Somatic
Required for sequence vari	ants: gene, c./p., tro	anscript# R	equired for CNVs	gene, transcript	#, exon # OR build, co	oordinates				I.	
☐ Lynch Screening MSI: ☐ Not Done IHC: ☐ Not Done Other Results		nor Type:] Stable/Lov Abser		HI □ MSH2 □]MSH6 □ PMS2						
‡For certain tests, GeneDx m be provided <u>in the table abo</u> be possible to comment up	ove at the time the	test order is p	olaced. If you do no	ot complete the t	able above and chec	k off that a	oreviously identi	ified variar	nt is a variant		
			TA	RGETED VA	RIANT TESTING	;					
Individual to be teste	d: OAffected	d/Symptor	natic O	Unaffected/	Asymptomatic						
☐ Known Familial Variant(s) in a Nuclear Gene ☐ Confirmation of Variant Identified in Research Lab ☐ Targeted Mosaic Variant Testing* ☐ Known Familial Copy Number Variant(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 GeneD	x Accessio	n#		
I	Positive contro	ol included/	will be sent - Po	sitive control i	previous test was p is recommended it included on a nego	f previous	test was perfo		t another la	ıb.	
VARIANT INFORMA	TION (please fi			n if family men		included)		Number	of Variants	3:	
Gene		Coding DNA	(c./m.)		Amino Acid (p.)			Transcript	(NM#)		
Gene	(c./m.)	/m.) Amino Acid (p.)			Transcript (NM#)						

Gene(s)

Gene(s)

COPY NUMBER VARIANT

Exon #

Exon #

Coordinates

Coordinates

Number of Variants:

Genome Build

Genome Build



HEREDII	ARY CANCER TES	ST REQU	ISITION	IFORM			G	enelx		
First Name		Last Name				Da	te of Birth			
			TEST	MENU						
TEST CODE	TEST N	AME		TEST CODE		TEST N	NAME			
BREAST/GYNECOLOGIC CANCER PANELS										
☐ B362	B362 BRCAT/2 Sequencing and Deletion/Duplication Analysis 🔲 J055 Breast Cancer Management Panel									
☐ B363	Reflex to Rest of Comprehensive Common Cancer Panel 🔲 B363 Reflex to Rest of Comprehensive Common Cancer Panel									
□ B273 Breast/Gyn Cancer Panel										
☐ B363	☐ B363 Reflex to Rest of Comprehensive Common Cancer Panel									
COLORECT	AL CANCER PANELS									
☐ B274	Colorectal Cancer Panel			□ B522	Lynch/Colore	ectal High Risk Pane	I			
☐ B363	Reflex to Rest of Comprehensive	Common Cance	er Panel	□ B363	Reflex to Resi	t of Comprehensive	Common Canc	er Panel		
MULTIPLE C	ANCER PANELS									
☐ B275	Comprehensive Common Cance	er Panel		☐ B751	Common Ca	incer Management	Panel			
				□ B363	Reflex to Resi	t of Comprehensive	Common Canc	er Panel		
OTHER CAN	CER SPECIFIC PANELS									
☐ B343	Pancreatic Cancer Panel			□ J665	Hereditary Pr	ostate Cancer Pane	el			
□ B363	Reflex to Rest of Comprehensive	Common Cance	er Panel	□ B363	Reflex to Rest	t of Comprehensive	Common Canc	er Panel		
☐ B394	Renal Cancer Panel									
☐ B363	Reflex to Rest of Comprehensive	Common Cance	er Panel							
SPECIALTY	PANELS									
□ T830	Hereditary MDS/Leukemia Panel ¹									
WRITE-IN T	EST SELECTION									
☐ Test Code		Test	Name:							
	ehensive Common Cancer Panel is should be ordered at the time of c									
'										
			NO MOTEUS	COLOGY PAN	EI.					
0	Duranta and Daniel		OSTOM ON	COLOGIFAN	EL					
	Ox Custom Panel									
□ B749	OncoGeneDx Custom Panel									
	ne or more genes to create a custo es from test code(s)					Panel is Negative, r	eflex to test code	e		
☐ AIP ☐ ALK ☐ ANKRD26 ☐ APC ☐ ATM ☐ AXIN2 ☐ BAP1 ☐ BARD1 ☐ BMPRIA	□ BRCA2 □ CTNNA1 □ BRIP1 □ DDX41 □ CDC73 □ DICER1 □ CDH1 □ EPCAM* □ CDK4 □ ETV6 □ CDKN1B □ FANCC □ CDKN2A □ FANCM	☐ GALNT12 ☐ GATA2 ☐ HOXB13 ☐ IKZF1 ☐ KIT ☐ LZTR1 ☐ MAX	☐ MET ☐ MITF ☐ MLHI ☐ MSH2 ☐ MSH3 ☐ MSH6 ☐ MUTYH ☐ NBN ☐ NFI	☐ NF2 ☐ NTHL1 ☐ PALB2 ☐ PAX5 ☐ PDGFRA ☐ PHOX2B* ☐ PMS2 ☐ POLD1 ☐ POLE	☐ POTI ☐ PRKARIA ☐ PTCHI ☐ PTEN ☐ RAD5IC ☐ RAD5ID ☐ RBI ☐ RECQL ☐ RET*	☐ RNF43 ☐ RPS20 ☐ RTEL1 ☐ RUNX1 ☐ SAMD9 ☐ SAMD9L ☐ SCG5/GREMI* ☐ SDHA* ☐ SDHAF2	☐ SDHB ☐ SDHC ☐ SDHD ☐ SMAD4 ☐ SMARCA4 ☐ SMARCBI ☐ SMARCEI ☐ SRP72 ☐ STKII	□ SUFU □ TERC □ TERT □ TINF2 □ TMEM127 □ TP53 □ TSC1 □ TSC2 □ VHL		
*Testing includes (seq only).	sequencing and deletion/duplication for	r all genes except: (CEBPA (seq only),	EPCAM (del/dup o	nly), <i>PHOX2B</i> (seq	only), RET (seq only), So	CG5/GREM1 (del/du	p only), SDHA		
			DID YOU REA	MEMBER TO	?					
□ Ledest :	and the latest and th		DID TOO KE	EMBER 10						
⊥abel speci	men tube appropriately with TWO	iaentifiers								

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.

 $\hfill \Box$ Get a signature for medical necessity and patient consent

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.

régo to p fam any	signing this form: (i) I acknowledge that I have read or have had read to me the arding genetic testing; (ii) I have had the opportunity to ask questions about serform genetic testing as ordered; (iv) I understand that, for tests that evaluated in the series of the se	the testing, the procedure, the risks, and the alternative ate data from multiple family members concurrently, te nade available to all tested individuals and their health which I may be contacted, I consent to receiving email	es; (iii) I authorize GeneDx est results from these care providers; (v) if at					
	Secondary Findings Opt-out. Check this box if you do not wish to receive AC ONLY; not for <i>Xpanded®</i> or Slice tests).	:MG 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 Y sample longer than 60 days after the completion of testing, and to be used validation, quality assurance, and training purposes. Otherwise, New York law for test development studies.	as a de-identified sample for test development and im	provement, internal					
	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.							
igno	gnature of Patient/Legal Guardian (required) Date							
igno	ature of Relative A/Legal Guardian	Relative A Relationship to Patient	Date					
igno	nature of Relative B/Legal Guardian Relative B Relationship to Patient Date							