

# GENETICS LABORATORY WHOLE EXOME SEQUENCING (WES) FORM

Ship To: O'Donoghue Research Bldg 1122 NE 13 Street, Suite 1400

> Phone: 405-271-3589 Fax: 405-271-7117

Oklahoma City, OK 73104

www.genetics.ouhsc.edu

After hours phone: 405-496-9514

PLEASE COMPLETE ALL FORMS AND SEND WITH SAMPLE(S)

Page 1 of 4						
	REFERRING PHYSICIAN/FACILITY			PATIEN'	T INFORMATION	
Physician N	Name		Patient Name (la			
NPI			Parent Name (if patient is minor)			
Phone() Fax()			DOB SSN MRN			
Genetic Co	unselor Phone()		Sex:   Male   F	emale □ Ambiguo	us 🗆 Other 🗆 Inpa	atient   Outpatient
Laboratory	/Institution		Ethnicity of patient (check all that apply)  □ African-American □ Asian □Caucasian/NW European □ E. Indian			
Address		□ Hispanic □ Jewish-Ashkenazi □ Jewish-Sephardic □ Native American □ Native Hawaiian/Other Pacific Islander □ Other				
City	State Zip		Patient's Address			
Phone_(	) Fax ()				State	
		SPECIMEN	INFORMATION			
□ Peripher	al Blood 3-5 cc larged EDTA tube (purple top), mix w	vell keep at r	oom temperature	or cooler, do not f	reeze.	
□ Isolated	DNA qty 20ug Specim	en Collectio	n Date		_	
	materials via Fedex/UPS, packages can only be accepte or after hours. For courier service in Oklahoma City me					y personnel on
		TEST	OPTIONS			
□ WES-	Proband Only TAT 90 days	PT: 81415	□ WES- Tri	o TAT 90 days	СР	T: 81416x2
	FAMIL	Y MEMBER S	AMPLE INFORMA	TION		,
Biological Mother:		DOB:		☐ Symptomatic	☐ Asymptomatic	**All samples related
Biological Father:		DOB:		□ Symptomatic	☐ Asymptomatic	to proband must be
Other:		DOB:		☐ Symptomatic	☐ Asymptomatic	received within 10
	Relationship to proband:	Total # of sar	mples sent			days of proband
**A COMP	LETED PATIENT/PROBAND CLINICAL INFORMATION	FORM AND	CLINIC NOTES MI	JST ACCOMPANY 1	THE SAMPLES.	
	PATIENT/GUARDIAN CONSENT-	Please read	the Informed Co	nsent document be	efore signing	
sion for [m publication and publicaties, includ	I the Informed Consent document and I give OUHSC (y/my child's] specimen and clinical information to be now, when appropriate. [My/my child's] name or other pations. I also give OUHSC Genetics Lab permission to ing treatments for the condition in [my/my child's] for its box to opt-out of receiving secondary findings.	e used in de-i personal ider inform me o amily.	identified studies ntifying information r [my/my child's]	at OUHSC Genetics on will not be used health care provide	Lab to improve genet in or linked to the resuler in the future about in	ic testing and for ults of any studies
⊔ Cneck th	is box to opt- <b>out</b> of research.	ratient	t/Guardian Signat	ure	Date	
	ADDITIONAL REPORT			GENETICS	LABORATORY USE ONL	Υ
Physician/Facility			Laboratory Number			
Phone_() Fax_()			Date/Time/Location of Pick-Up or Delivery			
Address		Initials	Check-in	Previous Lab Number		



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CONSENT AND AUTHORIZATION			
I understand that my health care provider has ordered whole exome sequencing for myself/my child:			
Sign Here			

#### **INFORMED CONSENT**

#### What are potential results from this test?

There are three possible result outcomes:

- 1) Positive Result: A positive result indicates that a genetic change has been identified, and it explains the potential cause of [my/my child's] genetic condition or indicates that [I am/my child is] at increased risk to develop a disorder in the future. It is possible to test positive for more than one genetic change.
- 2) Negative Result: A negative result means that no disease-causing genetic change was identified in the test performed. It does NOT guarantee that [I/my child] will be healthy or free from genetic disorders or medical conditions. If [I/my child] test negative for a genetic change known to cause a specific condition in other members of the family, a diagnosis of the same genetic disorder in [me/my child] due to this specific change has been ruled out.
- Nariant of Uncertain Significance (VUS): A variant of uncertain significance indicates that a genetic change was detected, but it is currently unknown whether that change is associated with a genetic condition. A VUS is not the same as a positive result and does not specify whether [I am/my child is] at increased risk for a genetic condition. The variant may be a normal genetic change found in the human population, or it could be disease-causing. Further analysis may be warranted and include testing both parents (if not already tested) as well as other family members such as: siblings, grand parents, or aunts and uncles. Detailed medical records or additional information may be requested to re-classify this result.

#### **Additional Positive Results**

- 1) Secondary Findings: Secondary findings are genetic changes identified 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 whole exome sequencing. Refer to the latest version of the ACMG Recommendations for Reporting of Secondary Findings in Clinical Exome and Genome Sequencing for a complete list of the genes and associated genetic disorders.
- 2) Incidental Findings: In rare instances, this test may reveal an unexpected, yet important genetic change that is not directly related to the reason for ordering this test. This test may tell me about the risk for another genetic condition [I am/my child is] not aware of, or it may indicate differences in the number or rearrangements of sex chromosomes. Information may be disclosed to the ordering health care provider if it will likely impact [your/ your child's] medical care.

#### What will be reported

- 1) For the proband: All known and/or expected disease-causing genetic changes identified in the coding exons of the genes correlating with the submitted phenotype or included in the recommended gene list issued by the ACMG.
- 2) For relatives: The presence or absence of any secondary findings reported for the proband will be provided for all relatives.

#### **Limitations and Risks**

- 1) Disease-causing genetic changes 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 disease-causing genetic changes.
- 2) Pathogenic variants that may be present in a relative, but are not present in the proband, will not be identified or reported.
- 3) Only changes in the genetic sequence will be reported in secondary findings. Larger deletions/duplications, abnormal methylations, triplet repeats and other expansions, or other variants not routinely identified by whole exome sequencing will not be reported.
- 4) This test has the potential to identify non-paternity and consanguinity (relatedness) between the proband's parents. It may be necessary to report these results to the health care provider who ordered the test if it will affect medical management of the proband.
- 5) Inaccurate results may be reported due to mislabeled samples, inaccurate reporting of clinical/medical information, rare technical errors, or unusual circumstances such as bone marrow transplantation, or the presence of mosaicism (change not present in every cell).

#### Interpretation

- 1) Results will be interpreted with available information in the medical literature, research databases, and scientific databases. Due to the change in medical/scientific knowledge and published studies, new information that becomes available may replace or build upon the information utilized in the interpretation of [my/my child's] results.
- 2) Providers can contact the laboratory at any time to discuss the classification or re-classification of an identified genetic change.
- 3) [My/my child's] health care providers and I may monitor publicly available resources used by the medical community, such as ClinVar (www.clinvar.com), to find updated information about the interpretation of [my/my child's] genetic change(s).
- 4) Results will be released in a single report in the proband's name. Additional reports will NOT be released in the proband's family members' names if a trio is submitted.
- 5) Occasionally, an additional sample may be needed if the initial specimen is not adequate for analysis.

#### **Patient Confidentiality and Genetic Counseling**

- 1) It is strongly recommended that [I/my child] receive genetic counseling before and after having this test. I can find a genetic counselor in my area at www.nsgc.org.
- 2) Further testing or consultations may be necessary based on results.
- To maintain confidentiality, the test results will only be released to the referring health care provider, the ordering laboratory, me, other health care providers involved in [my/my child's] diagnosis and treatment, or to others as entitled by law.
- 4) The United States Federal Government has enacted several laws that prohibit discrimination based on genetic test results by health insurance companies and employers. In addition, these laws prohibit unauthorized disclosure of this information. For more information, I understand I can visit www.genome.gov/10002077.



### **Genetics Laboratory**

### **Patient/Proband Clinical Information Form**

Last Name:	First	: MI: DOB:		
Primary Indications for Testing		Development & Cognition		
□ Multiple Congenital Anomali     □ Developmental Delays     □ Neurological/Muscular Disor      Previous Testing     □ Karyotype/FISH     □ CMA     □ Newborn Screen Result     □ Other Results      Family History (provide pedigree)     □ Consanguinity     □ Family History of Genetic Dis	der 	Autism Spectrum Fine Motor Delays Global Delay Gross Motor Delays Intellectual Delays Mild Moderate Severe Learning Delays Speech Delay		
Perinatal History	Cardiac	Skeletal	Neurological & Muscular	
☐ IUGR / SGA	☐ Arrhythmia	☐ Arthrogryposis	Ataxia	
Growth	☐ ASD ☐ Cardiomyopathy	☐ Club Foot/Feet☐ Contractures	<ul><li>Brain Anomaly</li><li>Cerebellar anomaly</li></ul>	
☐ Failure to thrive	☐ Coarctation of aorta	Joint Hypermobility	☐ Chorea/Dystonia	
<ul><li>Macrocephaly</li><li>Microcephaly</li></ul>	■ Dextrocardia	☐ Kyphosis	☐ Encephalopathy	
Overgrowth/Tall	☐ Tetralogy of fallot	Limb Anomaly	☐ Holoprosencephaly	
☐ Short stature	<ul><li>□ Ventriculomegaly</li><li>□ VSD</li></ul>	<ul><li>Osteopenia</li><li>Pes Planus</li></ul>	<ul><li>☐ Hydrocephalus</li><li>☐ Hypertonia</li></ul>	
□ Other:	□ Other:	Polydactyly	Hypotonia	
Craniofacial Anomalies		□ Scoliosis	Lissencephaly	
☐ Cleft Lip	GI	☐ Skeletal Dysplasia	☐ Leukodystrophy	
☐ Cleft Palate	<ul><li>Anal Atresia</li><li>Chronic Obstruction</li></ul>	<ul><li>Syndactyly</li><li>Vertebral anomaly</li></ul>	<ul><li>Muscle Weakness/Atrophy</li><li>Peripheral Neuropathy</li></ul>	
<ul><li>Craniosynostosis</li><li>Dysmorphic Facies</li></ul>	Dysphagia	Other:	☐ Vermis Hypoplasia	
☐ Ear Malformation	☐ Esophageal Atresia		Other:	
☐ Other:	☐ Gastroschisis	Endocrine		
	Hirschsprung Disease	Diabetes Insipidus	Cancer/Tumors  Tumor	
Ear / Hearing Loss (HL)  Conductive HL	☐ Liver Disease ☐ Omphalocele	<ul><li>Diabetes Mellitus</li><li>Hyperthyroidism</li></ul>	(describe)	
☐ Microtia	□ Polysplenia	☐ Hypothyroidism	(46561186)	
☐ Sensorineural HL	☐ Situs Inversus	☐ Hyperparathyroidsim	Age of Onset	
Other:	□ Other:	☐ Hypoparathyroidism	Chin Hair 9 Naile	
Eye Anomalies Genitourinary		Hematologic/Immuno	Skin, Hair & Nails  Abnormal Hair	
☐ Aniridia	☐ Ambiguous Genitals	☐ Anemia	☐ Abnormal Nails	
☐ Congenital Cataract	☐ Cryptochidism	☐ Immunodeficient	☐ Hyperpigmentation	
☐ Cortical Blindness/CVI	☐ Hydronephrosis	☐ Neutropenia	(describe)	
☐ Coloboma ☐ Glaucoma	<ul><li>Hypospadias</li><li>Kidney Malformation</li></ul>	☐ Pancytopenia☐ Thrombocytopenia	Hypopigmentation (describe)	
Optic Nerve Abnormality	Renal Agenesis	Other:	☐ Lipoma	
☐ Ptosis	☐ Renal Tubulopathy		Other:	
Retinitis Pigmentosa	□ Other:		Bankalaniin Alexaniin aliai	
□ Other:			Metabolic Abnormalities  Hyperammonemia	
Pulmonary			☐ Ketosis	
☐ Diaphragmatic Hernia			☐ Lactic Acidosis	
☐ TE Fistula			☐ Metabolic Acidemia	
Other:			Other:	



### Genetics Laboratory Billing Information Form

Patient Name LAST\_\_\_\_\_\_ FIRST\_\_\_\_\_\_ MI\_\_\_\_

# YOU MUST CHOOSE ONE OF THE THREE BILLING OPTIONS LISTED BELOW. PLEASE FORWARD ALL BILLING QUESTIONS TO DANIELLE OTIS AT DOTIS@OUHSC.EDU OR CALL 405-271-3589 OPT 4 AT THIS TIME WE DO NOT ACCEPT OUT-OF-STATE MEDICAID

PAYMENT OPTION 1-INSTITUTION	
INSTITUTION NAME	
BILLING ADDRESS	
CITY, STATE, ZIP	CONTACT NAME
PHONE NUMBER FAX NUMBE	ERCONTACT EMAIL ADDRESS
PAYMENT OPTION 2-SELF PAY (PAYMENT MUST BE S	ENT WITH SAMPLE)
☐ CREDIT CARD (CIRCLE ONE) AMEX DISCOVER V	ISA MASTERCARD AMOUNT TO CHARGE
VALID CARD #	EXP DATE
CVV CODE CARDHOLDER PRINTED NA	ME
	CITY, STATE, ZIP
CARDHOLDER SIGNATURE	
	NT ENCLOSED
PAYMENT OPTION 3-INSURANCE PROVIDE A LEGIBLE PLEASE NOTE: OUR FACILITY WILL CONFIRM COVERAGE OUR OFFICE CAN ALSO OBTAIN PRE-AUTHORIZATION	GE AND VERIFY WHETHER OR NOT THE TEST(S) ORDERED ARE COVERED BY YOUR PLAN.
PRIMARY INSURANCE POLICYHOLDER NAME	POLICYHOLDER DOB
PRIMARY POLICYHOLDER SS#	GENDER: M F EMPLOYER
RELATIONSHIP TO PATIENT	POLICY #
GROUP#	INSURANCE CO. NAME
PHONE	CLAIMS ADDRESS
CITY, STATE, ZIP	INSURANCE AUTH #
SECONDARY INSURANCE POLICYHOLDER NAME	POLICYHOLDER DOB
SECONDARY POLICYHOLDER SS#	GENDER: M F EMPLOYER
RELATIONSHIP TO PATIENT	POLICY #
GROUP#	INSURANCE CO. NAME
PHONE	CLAIMS ADDRESS
CITY, STATE, ZIP	INSURANCE AUTH #
ANY MEDICAL INFORMATION REQUESTED ON MYSELF, OR MY FITS OF INSURANCE TO UNIVERSITY OF OKLAHOMA HSC GENE AUTHORIZED SERVICES AND REMAINING BALANCES AFTER INS	GE PERFORMED. I AUTHORIZE THE UNIVERSITY OF OKLAHOMA HSC GENETICS LABORATORY TO FURNISH COVERED DEPENDENTS. IN CONSIDERATION OF SERVICES RENDERED, I TRANSFER AND ASSIGN ANY BENETICS LABORATORY. I UNDERSTAND I AM RESPONSIBLE FOR ANY CO-PAY, DEDUCTIBLES, OR NON-BURANCE REIMBURSEMENT. I UNDERSTAND I AM FULLY RESPONSIBLE FOR PAYMENT OF MY ACCOUNT IF S NOT A PARTICIPANT WITH MY HEALTH PLAN OR MY HEALTH PLAN DOES NOT FULLY REIMBURSE MY MEDINECESSITY.
PRINTED NAME	SIGNATURE