

## SPERM DONOR GENETIC TESTING SUMMARY

Donor # 7666

Fairfax Cryobank recommends reviewing this genetic testing summary with your healthcare provider to determine suitability.

Last Updated: 11/3/2025

Donor Reported Ancestry: African American, Brazilian

Jewish Ancestry: No

Genetic Test*	Result	Comments Donor's Residual Risk**
Chromosome analysis (karyotype)	Normal male karyotype	No evidence of clinically significant chromosome abnormalities
Hemoglobin evaluation	Normal hemoglobin fractionation and MCV, slightly decrease in MCH	Results consistent with DNA test results that indicate this donor is a silent carrier of alpha thalassemia.
Expanded Genetic Disease Carrier Screening Panel attached - 549 diseases by gene sequencing and del/dup analysis.	<p><b>Silent Carrier: Alpha - Thalassemia (aa/a - ) (HBA2)</b></p> <p><b>Carrier: Thyroid Dyshormonogenesis 6 (DUOX2)</b></p> <p>Negative for other genes tested.</p>	<p>Partner testing is recommended before using this donor.</p> <p>Most people with a variant in this gene are carriers of TDH6, but do not have the condition. Some people with a variant in this gene have symptoms of TDH6 as babies that go away as they age.</p>

\*No single test can screen for all genetic disorders. A negative screening result significantly reduces, but cannot eliminate, the risk for these conditions in a pregnancy.

\*\*Donor residual risk is the chance the donor is still a carrier after testing negative.

Patient Information	
Patient Name:	Donor 7666
Date Of Birth:	[REDACTED]
Gender:	Male
Ethnicity:	Other
Patient ID:	N/A
Medical Record #:	N/A
Collection Kit:	[REDACTED]
Accession ID:	N/A
Case File ID:	[REDACTED]

Test Information	
Ordering Physician:	[REDACTED]
Clinic Information:	Fairfax Cryobank
Phone:	
Report Date:	07/16/2025
Sample Collected:	07/01/2025
Sample Received:	07/03/2025
Sample Type:	Blood



## CARRIER SCREENING REPORT

**ABOUT THIS SCREEN:** Horizon™ is a carrier screen for specific autosomal recessive and X-linked diseases. This information can help patients learn their risk of having a child with specific genetic conditions.

**ORDER SELECTED:** The Horizon Custom panel was ordered for this patient. Males are not screened for X-linked diseases

### FINAL RESULTS SUMMARY:



#### SILENT CARRIER for Alpha-Thalassemia (aa/a-)

Positive for the pathogenic alpha 3.7 deletion of the HBA2 gene. Depending on the carrier status of the individual's partner, this couple may be at increased risk to have a child with Hemoglobin H Disease. Carrier screening for this individual's partner is suggested.

#### CARRIER for Thyroid Dyshormonogenesis 6

Positive for the likely pathogenic variant c.4133\_4146del (p.E1378Gfs\*30) in the DUOX2 gene. If this individual's partner is a carrier for THYROID DYSHORMONOGENESIS 6, their chance to have a child with this condition may be as high as 1 in 4 (25%). Carrier screening for this individual's partner is suggested.

#### Negative for 547 out of 549 diseases

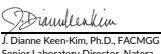
No other pathogenic variants were detected in the genes that were screened. The patient's remaining carrier risk after the negative screening results is listed for each disease/gene on the Horizon website at <https://www.natera.com/panel-option/h-all/>. Please see the following pages of this report for a comprehensive list of all conditions included on this individual's screen.

Carrier screening is not diagnostic and may not detect all possible pathogenic variants in a given gene.

#### RECOMMENDATIONS

Individuals who would like to review their Horizon report with a Natera Laboratory Genetic Counselor may schedule a telephone genetic information session by calling 650-249-9090 or visiting [naterasession.com](http://naterasession.com). Clinicians with questions may contact Natera at 650-249-9090 or email [support@natera.com](mailto:support@natera.com). Individuals with positive results may wish to discuss these results with family members to allow them the option to be screened. Comprehensive genetic counseling to discuss the implications of these test results and possible associated reproductive risk is recommended.

  
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**Patient Information**

Patient Name: Donor 7666

**Test Information**

Ordering Physician: [REDACTED]

Date Of Birth: [REDACTED]  
Case File ID: [REDACTED]

Clinic Information: Fairfax Cryobank

Report Date: 07/16/2025

**ALPHA-THALASSEMIA SILENT CARRIER****Understanding Your Horizon Carrier Screen Results****What is Alpha-Thalassemia?**

Alpha-Thalassemia refers to a group of inherited blood disorders that reduce the amount of hemoglobin, the protein in red blood cells that carries oxygen to cells throughout the body. A person with one of the Alpha-Thalassemia diseases has lifelong anemia. Mild anemia can lead to tiredness, irritability, dizziness, lightheadedness and a rapid heartbeat. Severe anemia can be life threatening and may require routine blood transfusions. In some cases, affected individuals have been treated with stem cell transplantation from cord blood or bone marrow. Couples at risk of having an affected child may consider cord blood banking, as siblings have a higher chance of being a match for stem cell transplantation than a non-related individual. More information can be found at: <https://parentsguidecordblood.org/en>. Clinical trials involving potential new treatments for these conditions may be available (see [www.clinicaltrials.gov](http://www.clinicaltrials.gov)).

**What causes Alpha-Thalassemia?**

Hemoglobin is made of both alpha globin and beta globin proteins. There are four HBA genes (also called alpha globin genes) that are responsible for making alpha globin. Alpha-Thalassemia occurs when three or more of these four alpha globin genes are missing or changed. The exact type of Alpha-Thalassemia a person has depends on how many of the alpha globin genes are not working. Hemoglobin H Disease (a/-): three missing or changed alpha globin genes. A person who has three missing or changed alpha globin genes has Hemoglobin H Disease. Hemoglobin H Disease can be mild or severe. People with severe disease may have chronic anemia, liver disease, and bone changes. Some people with Hemoglobin H Disease require frequent blood transfusions and other treatments. Alpha-Thalassemia Major, also known as Hemoglobin Bart's Disease (---): four missing or changed alpha globin genes. This results in severe fatal anemia. Affected babies develop symptoms before birth and without treatment typically do not survive the newborn period. Fetal blood transfusions during pregnancy may allow survival until after birth, at which time either lifelong transfusions or a stem cell transplantation will be necessary. Mothers who are pregnant with a fetus with Alpha-Thalassemia major can develop health problems during pregnancy. Alpha-Thalassemia is inherited in an autosomal recessive manner. Children typically inherit four copies of each alpha globin gene, two copies from the mother and two copies from the father. This means that both parents must be carriers of one or more missing or changed alpha globin genes to have a child who is affected with Hemoglobin H Disease or Alpha-Thalassemia Major.

**What do my carrier results mean?**

One missing or changed alpha globin gene was identified with your Horizon test. People with one missing or changed alpha globin gene are Alpha-Thalassemia silent carriers. People who are silent carriers for Alpha-Thalassemia usually have no health problems and have normal hemoglobin levels. Thalassemia can occur in people of any ethnicity. It is more common in people with Chinese, Southeast Asian, Indian, Middle Eastern, African, and Mediterranean ancestry.

If your partner is a carrier for Alpha-Thalassemia with two genes missing or changed on the same chromosome (in 'cis'), you would have a 1 in 4, or 25%, chance in each pregnancy of having a child with Hemoglobin H Disease. You are not at risk for having a baby with Alpha-Thalassemia Major. The majority of people of Asian ancestry who have two missing alpha globin genes have them on the same chromosome (in 'cis').

If your partner is a carrier for Alpha-Thalassemia with two genes missing or changed that are located on opposite chromosomes (in "trans"), each of your children would have a 50% chance of being carriers of Alpha-Thalassemia (with two genes missing or changed on opposite chromosomes), but you are not at risk to have a child with either Hemoglobin H Disease or Alpha-Thalassemia Major. The majority of people of African-American ancestry who have two missing alpha-globin genes have them on opposite chromosomes.

If your partner is an Alpha-Thalassemia Silent Carrier (with one gene missing or changed), each of your children would have a 25% chance of being carriers of Alpha-Thalassemia (with two genes missing or changed on opposite chromosomes) and a 50% chance of being Alpha-Thalassemia Silent carriers. You would not be at risk to have a child with either Hemoglobin H Disease or Alpha-Thalassemia Major.

**What can I do next?**

You may wish to speak with a local genetic counselor about your carrier test results. A genetic counselor in your area can be located on the National Society of Genetic Counselors website ([www.nscc.org](http://www.nscc.org)). Your siblings and other relatives are at increased risk to also have this mutation. You are encouraged to inform your family members of your test results as they may wish to consider being tested themselves. If you are pregnant, your partner can have carrier screening for Alpha-Thalassemia ordered by a health care professional. If your partner is not found to be a carrier for Alpha-Thalassemia, your risk of having a child with Hemoglobin H Disease is greatly reduced. Couples at risk of having a baby with Hemoglobin H Disease can opt to have prenatal diagnosis done through chorionic villus sampling (CVS) or amniocentesis during pregnancy or can choose to have the baby tested after birth. If you are not yet pregnant, your partner can have carrier screening for Alpha-Thalassemia ordered by a health care professional. If your partner is found to be a carrier for Alpha-Thalassemia (with two missing or non-working alpha globin genes on the same chromosome, in 'cis') you have several reproductive options to consider:

- Natural pregnancy with or without prenatal diagnostic testing of the fetus or testing the baby after birth for Hemoglobin H Disease
- Preimplantation genetic diagnosis (PGD) with in vitro fertilization (IVF) to test embryos for Hemoglobin H Disease
- Adoption or use of a sperm or egg donor who is not a carrier for Alpha-Thalassemia

**What resources are available?**

- March of Dimes: <http://www.marchofdimes.org/baby/thalassemia.aspx>
- Cooley's Anemia Foundation: [www.thalassemia.org](http://www.thalassemia.org)
- Prenatal diagnosis done by CVS: <http://www.marchofdimes.org/chorionic-villus-sampling>.

**Patient Information**

Patient Name: [REDACTED]

**Test Information**

Ordering Physician: [REDACTED]



Clinic Information: [REDACTED]

Date Of Birth: [REDACTED]

Case File ID: [REDACTED]

Report Date: [REDACTED]

**THYROID DYSHORMONOGENESIS 6****Understanding Your Horizon Carrier Screen Results****What does being a carrier mean?**

Your results show that you are a carrier of thyroid dyshormonogenesis 6 (TDH6). Most people with a variant in this gene are carriers of TDH6, but do not have the condition. Some people with a variant in this gene have symptoms of TDH6 as babies that go away as they age.

Your children are at risk for TDH6 or for short-term symptoms of this condition, but you are not certain to have a child with this condition. Further testing can be done to see if your partner or donor is a carrier.

**What is thyroid dyshormonogenesis 6 (TDH6)?**

TDH6 causes the body to not make enough thyroid hormones, resulting in congenital hypothyroidism (CH).<sup>1,2</sup> Some people with CH have no symptoms. Other people with CH can be less active, sleep more than normal, and have feeding problems or constipation. People with CH that is not treated can also have slow growth and intellectual disability.<sup>2</sup> With early treatment, people with TDH6 usually have normal development.<sup>3</sup> Newborn screening can detect over 90% of babies with CH.<sup>4</sup>

Carriers of TDH6 can have mild hypothyroidism as babies. Thyroid hormone levels can be lower than average at birth and increase with age.<sup>1,2</sup>

Clinical trials involving potential new treatments for this condition could be available (see [clinicaltrials.gov](https://clinicaltrials.gov)).

**What causes thyroid dyshormonogenesis 6 (TDH6)?**

TDH6 is caused by changes, or variants, in the DUOX2 gene. These changes make the gene not work properly. Genes are a set of instructions inside the cells of our bodies that tell our bodies how to grow and function. Everyone has two copies of the DUOX2 gene. Carriers of TDH6 have one working copy and one non-working copy of the gene. Some carriers have low levels of thyroid hormones as babies, but have normal thyroid function as they get older. People with TDH6 have no working copies of the gene.

TDH6 is usually passed down, or inherited, from both genetic parents. We inherit one copy of the DUOX2 gene from each of our genetic parents. When both genetic parents are carriers, each child has a 1 in 4 (25%) chance of inheriting two non-working genes and having TDH6. Each child also has a 1 in 2 (50%) chance of being a carrier of TDH6 and a 1 in 4 (25%) chance of inheriting two working copies of the gene. This type of inheritance is called autosomal recessive inheritance.

**Will my children have thyroid dyshormonogenesis 6 (TDH6)?**

If your partner or donor also has a non-working copy of the DUOX2 gene, your children could have TDH6. Each child you have together would have a 1 in 4 (25%) chance of having TDH6. Each child you have together would also have a 1 in 4 (25%) chance of not having any variants in the DUOX2 gene. Each child would have a 1 in 2 (50%) chance of being a carrier and could have symptoms of the condition as a baby.

If your partner or donor has DUOX2 carrier screening and no variants are found, the chance that your children would have two TDH6 variants is very low. In this situation, each child you have together would have a 1 in 2 (50%) chance of being a carrier and could have symptoms of TDH6 as a baby.

**What can I do next?**

If you want to know if your children are at risk for TDH6, your partner or donor would need to have DUOX2 carrier screening. If you have questions about this testing, please ask your healthcare provider or use the resources below. Many people find it helpful to speak with a genetic counselor.

If your partner or donor is found to be a TDH6 carrier, your children would be at risk for having TDH6. Your children are also at risk of being carriers who have low levels of thyroid hormones as babies.

If you or your partner or surrogate are currently pregnant, tests called CVS (chorionic villus sampling) and amniocentesis can be done during pregnancy to find out if a baby has TDH6. These tests both have a small risk of miscarriage. Babies can also be tested for TDH6 after birth instead.

If you or your partner or surrogate are not yet pregnant, you could have these options:

- natural pregnancy with CVS or amniocentesis to test for TDH6 during pregnancy;
- natural pregnancy and testing the baby after birth for TDH6;
- preimplantation genetic testing (PGT-M) with in vitro fertilization (IVF) to test embryos for TDH6;
- adoption; or
- use of a sperm or egg donor who had no variants found in DUOX2 carrier screening.

**Where can I find more information?**

- Pediatric Endocrine Society [pedsendo.org/patient-resource/congenital-hypothyroidism](https://pedsendo.org/patient-resource/congenital-hypothyroidism)
- American Thyroid Association [thyroid.org/professionals](https://thyroid.org/professionals)
- CVS [marchofdimes.org/chorionic-villus-sampling](https://marchofdimes.org/chorionic-villus-sampling)
- Amniocentesis [marchofdimes.org/pregnancy/amniocentesis](https://marchofdimes.org/pregnancy/amniocentesis)
- PGT-M [natera.com/womens-health/spectrum-preimplantation-genetics](https://natera.com/womens-health/spectrum-preimplantation-genetics)

**What does this mean for my family?**

**Patient Information**

Patient Name:

**Test Information**

Ordering Physician: [REDACTED]

Date Of Birth: [REDACTED]

Case File ID: [REDACTED]

Clinic Information:

Report Date:

You likely got (inherited) this non-working gene from one of your genetic parents. Your genetic siblings and other family members could also carry it. You should tell your family members about your test results so they can decide if they want carrier screening for TDH6.

**References**

1. Moreno JC et al. Inactivating mutations in the gene for thyroid oxidase 2 (THOX2) and congenital hypothyroidism. *New Eng. J. Med.* 347: 95-102, 2002.
2. Vigone MC et al. Persistent mild hypothyroidism associated with novel sequence variants of the DUOX2 gene in two siblings. *Hum. Mutat.* 26: 395, 2005.
3. MedlinePlus [Internet]. Bethesda (MD): National Library of Medicine (US). Congenital hypothyroidism; [updated 2015 Sep 1; cited 2024 March 3]. Available from: <https://medlineplus.gov/genetics/condition/congenital-hypothyroidism/>.
4. Büyükebiz A. Newborn screening for congenital hypothyroidism. *J Clin Res Pediatr Endocrinol.* 2013;5 Suppl 1(Suppl 1):8-12. doi: [10.4274/jcrpe.845](https://doi.org/10.4274/jcrpe.845). Epub 2012 Nov 15. PMID: 23154158; PMCID: PMC3608007.

**Patient Information**

Patient Name: [REDACTED]

**Test Information**

Ordering Physician: [REDACTED]

Date Of Birth: [REDACTED]

Clinic Information: [REDACTED]

Case File ID: [REDACTED]

Report Date: [REDACTED]

**VARIANT DETAILS****DUOX2, c.4133\_4146del (p.E1378Gfs\*30), likely pathogenic**

- The c.4133\_4146del (p.E1378Gfs\*30) variant in the DUOX2 gene has not been observed in the gnomAD v2.1.1 dataset.
- This premature termination variant is predicted to cause nonsense-mediated decay (NMD) in a gene where loss-of-function is a known mechanism of disease.
- This variant has been described in ClinVar [ID: 3618727].

**HBA1/HBA2, alpha 3.7 deletion, pathogenic**

- The alpha 3.7 or 4.2 deletion of the HBA1/HBA2 gene is a recombination deletion between the HBA1 and HBA2 gene, resulting in loss of one copy of the HBA1/HBA2 genes.
- Single allele deletion involving one of the four copies of the HBA1/HBA2 genes (alpha 3.7 deletion or alpha 4.2 deletion) has been reported in conjunction with deletions encompassing both HBA1 and HBA2 genes in individuals with HbH disease (PMID: 20301608, 7734346, 27492767, 29032940). Two single allele deletions in trans (alpha 3.7 deletion homozygous, alpha 4.2 deletion in trans, or alpha 3.7 deletion in trans with alpha 4.2 deletion) have been reported in individuals with alpha-thalassemia trait (PMID: 20301608, 29032940).
- This variant has been described in ClinVar [ID: 433555, 648517].

**Patient Information**

Patient Name: [REDACTED]

**Test Information**

Ordering Physician: [REDACTED]



Clinic Information: [REDACTED]

Date Of Birth: [REDACTED]

Case File ID: [REDACTED]

Report Date: [REDACTED]

**DISEASES SCREENED**

Below is a list of all diseases screened and the result. Certain conditions have unique patient-specific numerical values, therefore, results for those conditions are formatted differently.

**Autosomal Recessive**

1	17-BETA HYDROXYSTEROID DEHYDROGENASE 3 DEFICIENCY (HSD17B3) negative	BIOTINIDASE DEFICIENCY (BTD) negative
3	3-BETA-HYDROXYSTEROID DEHYDROGENASE TYPE II DEFICIENCY (HSD3B2) negative	BIOTIN-THIAMINE-RESPONSIVE BASAL GANGLIA DISEASE (BTBGD) (SLC19A3) negative
	3-HYDROXY-3-METHYLGLUTARYL-COENZYME A LYASE DEFICIENCY (HMGCL) negative	BLOOM SYNDROME (BLM) negative
	3-HYDROXYACYL-COA DEHYDROGENASE DEFICIENCY (HADH) negative	BRITTLE CORNEA SYNDROME 1 (ZNF469) negative
	3-METHYLACRYLIC ACIDURIA (MAAA) negative	BRITTLE CORNEA SYNDROME 2 (PRDM5) negative
	3-PHOSPHOGLYCERATE DEHYDROGENASE DEFICIENCY (PHGDH) negative	
5	5-ALPHA-REDUCTASE DEFICIENCY (SRD5A2) negative	
6	6-PYRUVOYL-TETRAHYDROPTERIN SYNTHASE (PTPS) DEFICIENCY (PTS) negative	
A		C
	ABCA4-RELATED CONDITIONS (ABCA4) negative	CANAVAN DISEASE (ASPA) negative
	ABETALIPOPROTEINEMIA (MTPP) negative	CARBAMOYL PHOSPHATE SYNTHETASE I DEFICIENCY (CPS1) negative
	ACHONDROGENESIS, TYPE 1B (SLC2A2) negative	CARNITINE DEFICIENCY (SLC22A5) negative
	ACHROMATOPSY, CNGB3-RELATED (CNGB3) negative	CARNITINE PALMITOYLTRANSFERASE IA DEFICIENCY (CPT1A) negative
	ACRODERMATITIS ENTEROPATHICA (SLC39A4) negative	CARNITINE PALMITOYLTRANSFERASE II DEFICIENCY (CPT2) negative
	ACTION MYOCLONUS-RENAL FAILURE (AMRF) SYNDROME (SCARB2) negative	CARNITINE-ACYLCARNITINE TRANSLOCASE DEFICIENCY (SLC25A20) negative
	ACUTE INFANTILE LIVER FAILURE, TRMU-RELATED (TRMU) negative	CARPENTER SYNDROME (RAB23) negative
	ACYL-COA OXIDASE I DEFICIENCY (ACOX1) negative	CARTILAGE-HAIR HYPOPLASIA (RMRP) negative
	AICARDI-GOUTIERES SYNDROME (SAMHD1) negative	CATECHOLAMINERGIC POLYMORPHIC VENTRICULAR TACHYCARDIA (CASQ2) negative
	AICARDI-GOUTIERES SYNDROME, RNASEH2A-RELATED (RNASEH2A) negative	CD59-MEDIATED HEMOLYTIC ANEMIA (CD59) negative
	AICARDI-GOUTIERES SYNDROME, RNASEH2B-RELATED (RNASEH2B) negative	CEP152-RELATED MICROCEPHALY (CEP152) negative
	AICARDI-GOUTIERES SYNDROME, RNASEH2C-RELATED (RNASEH2C) negative	CEREBRAL DYSGENESIS, NEUROPATHY, ICHTHYOSIS, AND PALMOPLANTAR KERATODERMA (CEDNIK) SYNDROME (SNAP29) negative
	AICARDI-GOUTIERES SYNDROME, TREX1-RELATED (TREX1) negative	CEREBROTENDINOUS XANTHOMATOSIS (CYP27A1) negative
	ALPHA-MANNOSIDOSIS (MAN2B1) negative	CHARCOT-MARIE-TOOTH DISEASE, RECESSIVE INTERMEDIATE C (PLEKHG5) negative
	ALPHA-THALASSEMIA (HBA1/HBA2) see first page	CHARCOT-MARIE-TOOTH DISEASE, TYPE 4D (NDRG1) negative
	ALPORT SYNDROME, COL4A3-RELATED (COL4A3) negative	CHEDIAK-HIGASHI SYNDROME (LYST) negative
	ALPORT SYNDROME, COL4A4-RELATED (COL4A4) negative	CHOREOACANTHOCYTOSIS (VPS13A) negative
	ALSTROM SYNDROME (ALMS1) negative	CHRONIC GRANULOMATOUS DISEASE, CYBA-RELATED (CYBA) negative
	AMISH INFANTILE EPILEPSY SYNDROME (ST3GAL5) negative	CHRONIC GRANULOMATOUS DISEASE, NCF2-RELATED (NCF2) negative
	ANDERMANN SYNDROME (SLC12A6) negative	CILIOPATHIES, RPGRIP1L-RELATED (RPGRIP1L) negative
	ARGININE:GLYCINE AMIDINOTRANSFERASE DEFICIENCY (AGAT DEFICIENCY) (GATM) negative	CITRIN DEFICIENCY (SLC25A13) negative
	ARGININEMIA (ARG1) negative	CITRULLINEMIA, TYPE 1 (ASS1) negative
	ARGINOSUCCINATE LYASE DEFICIENCY (ASL) negative	CLN10 DISEASE (CTSD) negative
	AROMATASE DEFICIENCY (CYP19A1) negative	COHEN SYNDROME (VPS13B) negative
	ASPARAGINE SYNTHETASE DEFICIENCY (ASNS) negative	COL11A2-RELATED CONDITIONS (COL11A2) negative
	ASPARTYLGLYCOSAMINURIA (AGA) negative	COMBINED MALONIC AND METHYLMALONIC ACIDURIA (ACSF3) negative
	ATAXIA WITH VITAMIN E DEFICIENCY (TTPA) negative	COMBINED OXIDATIVE PHOSPHORYLATION DEFICIENCY 1 (GFM1) negative
	ATAXIA-TELANGIECTASIA (ATM) negative	COMBINED OXIDATIVE PHOSPHORYLATION DEFICIENCY 3 (TSFM) negative
	ATAXIA-TELANGIECTASIA-LIKE DISORDER 1 (MRE11) negative	COMBINED PITUITARY HORMONE DEFICIENCY 1 (POU1F1) negative
	ATRANSFERRINEMIA (TF) negative	COMBINED PITUITARY HORMONE DEFICIENCY-2 (PROP1) negative
	AUTISM SPECTRUM, EPILEPSY AND ARTHROGRYPOSIS (SLC35A3) negative	CONGENITAL ADRENAL HYPERPLASIA, 11-BETA-HYDROXYLASE DEFICIENCY (CYP11B1) negative
	AUTOIMMUNE POLYGLANDULAR SYNDROME, TYPE 1 (AIRE) negative	CONGENITAL ADRENAL HYPERPLASIA, 17-ALPHA-HYDROXYLASE DEFICIENCY (CYP17A1) negative
	AUTOSOMAL RECESSIVE CONGENITAL ICHTHYOSIS (ARCI), SLC27A4-RELATED (SLC27A4) negative	CONGENITAL ADRENAL HYPERPLASIA, 21-HYDROXYLASE DEFICIENCY (CYP21A2) negative
	AUTOSOMAL RECESSIVE SPASTIC ATAXIA OF CHARLEVOIX-SAGUENAY (SACS) negative	CONGENITAL ADRENAL INSUFFICIENCY, CYP11A1-RELATED (CYP11A1) negative
B		CONGENITAL AMEGAKARYOCYTIC THROMBOCYTOPENIA (MPL) negative
	BARDET-BIEDL SYNDROME, ARL6-RELATED (ARL6) negative	CONGENITAL CHRONIC DIARRHEA (DGAT1) negative
	BARDET-BIEDL SYNDROME, BBS10-RELATED (BBS10) negative	CONGENITAL DISORDER OF GLYCOSYLATION, TYPE 1, ALG1-RELATED (ALG1) negative
	BARDET-BIEDL SYNDROME, BBS12-RELATED (BBS12) negative	CONGENITAL DISORDER OF GLYCOSYLATION, TYPE 1A, PMM2-Related (PMM2) negative
	BARDET-BIEDL SYNDROME, BBS1-RELATED (BBS1) negative	CONGENITAL DISORDER OF GLYCOSYLATION, TYPE 1B (MPI) negative
	BARDET-BIEDL SYNDROME, BBS2-RELATED (BBS2) negative	CONGENITAL DISORDER OF GLYCOSYLATION, TYPE 1C (ALG6) negative
	BARDET-BIEDL SYNDROME, BBS4-RELATED (BBS4) negative	CONGENITAL DYSERYTHROPOIETIC ANEMIA TYPE 2 (SEC23B) negative
	BARDET-BIEDL SYNDROME, BBS5-RELATED (BBS5) negative	CONGENITAL FINNISH NEPHROSIS (NPHS1) negative
	BARDET-BIEDL SYNDROME, BBS7-RELATED (BBS7) negative	CONGENITAL HYDROCEPHALUS 1 (CCDC88C) negative
	BARDET-BIEDL SYNDROME, BBS9-RELATED (BBS9) negative	CONGENITAL HYPERINSULINISM, KCNJ11-Related (KCNJ11) negative
	BARDET-BIEDL SYNDROME, TTC8-RELATED (TTC8) negative	CONGENITAL INSENSITIVITY TO PAIN WITH ANHIDROSIS (CIPA) (NTRK1) negative
	BARE LYMPHOCYTE SYNDROME, CITA-RELATED (CITA) negative	CONGENITAL MYASTHENIC SYNDROME, CHAT-RELATED (CHAT) negative
	BARTTER SYNDROME, BSND-RELATED (BSND) negative	CONGENITAL MYASTHENIC SYNDROME, CHRN-RELATED (CHRN) negative
	BARTTER SYNDROME, KCNJ1-RELATED (KCNJ1) negative	CONGENITAL MYASTHENIC SYNDROME, COLO-RELATED (COLQ) negative
	BARTTER SYNDROME, SLC12A1-RELATED (SLC12A1) negative	CONGENITAL MYASTHENIC SYNDROME, DOK7-RELATED (DOK7) negative
	BATTEN DISEASE, CLN3-RELATED (CLN3) negative	CONGENITAL MYASTHENIC SYNDROME, RAPSN-RELATED (RAPSN) negative
	BETA-HEMOGLOBINOPATHIES (HBB) negative	CONGENITAL NEPHROTIC SYNDROME, PLCE1-RELATED (PLCE1) negative
	BETA-KETO THIOLASE DEFICIENCY (ACAT1) negative	CONGENITAL NEUTROPENIA, G6PC3-RELATED (G6PC3) negative
	BETA-MANNOSIDOSIS (MANBA) negative	CONGENITAL NEUTROPENIA, HAX1-RELATED (HAX1) negative
	BETA-UREIDOPROPIONASE DEFICIENCY (UPB1) negative	CONGENITAL NEUTROPENIA, VPS45-RELATED (VPS45) negative
	BILATERAL FRONTOPARIEL POLYMICROGYRIA (GPR56) negative	CONGENITAL SECRETORY CHLORIDE DIARRHEA 1 (SLC26A3) negative
		CORNEAL DYSTROPHY AND PERCEPTIVE DEAFNESS (SLC4A11) negative
		CORTICOSTERONE METHYLOXIDASE DEFICIENCY (CYP11B2) negative
		COSTEIFF SYNDROME (3-METHYGLUTAConIC ACIDURIA, TYPE 3) (OPA3) negative
		CRB1-RELATED RETINAL DYSTROPHIES (CRB1) negative
		CYSTIC FIBROSIS (CFTR) negative
		CYSTINOSIS (CTNS) negative
		CYTOCHROME C OXIDASE DEFICIENCY, PET100-RELATED (PET100) negative
		CYTOCHROME P450 OXIDOREDUCTASE DEFICIENCY (POR) negative
D		D-BIFUNCTIONAL PROTEIN DEFICIENCY (HSD17B4) negative

**Patient Information**

Patient Name:

**Test Information**

Ordering Physician: [REDACTED]



Clinic Information:

Date Of Birth: [REDACTED]

Case File ID: [REDACTED]

Report Date:

**D**

DEAFNESS, AUTOSOMAL RECESSIVE 77 (LOXHD1) negative  
 DIHYDROPTERIDINE REDUCTASE (DHPR) DEFICIENCY (QDPR) negative  
 DONNAI-BARROW SYNDROME (LRP2) negative  
 DUBIN-JOHNSON SYNDROME (ABCC2) negative  
 DYSKERATOSIS CONGENITA SPECTRUM DISORDERS (TERT) negative  
 DYSKERATOSIS CONGENITA, RTEL1-RELATED (RTEL1) negative  
 DYSTROPHIC EPIDERMOLYSIS BULLOSA, COL7A1-Related (COL7A1) negative

**E**

EARLY INFANTILE EPILEPTIC ENCEPHALOPATHY, CAD-RELATED (CAD) negative  
 EHlers-DANLOS SYNDROME TYPE VI (PLOD1) negative  
 EHlers-DANLOS SYNDROME, CLASSIC-LIKE, TNXB-RELATED (TNXB) negative  
 EHlers-DANLOS SYNDROME, TYPE VII C (ADAMTS2) negative  
 ELLIS-VAN CREVELD SYNDROME, EVC2-RELATED (EVC2) negative  
 ELLIS-VAN CREVELD SYNDROME, EVC-RELATED (EVC) negative  
 ENHANCED S-CONE SYNDROME (NR2E3) negative  
 EPIMERASE DEFICIENCY (GALACTOSEMIA TYPE III) (GALE) negative  
 EPIPHYSEAL DYSPLASIA, MULTIPLE, 7/DESBUQUOIS DYSPLASIA 1 (CANT1) negative  
 ERCC6-RELATED DISORDERS (ERCC6) negative  
 ERCC8-RELATED DISORDERS (ERCC8) negative  
 ETHYLMALONIC ENCEPHALOPATHY (ETHE1) negative

**F**

FACTOR XI DEFICIENCY (F11) negative  
 FAMILIAL DYSAUTONOMIA (IKBKA) negative  
 FAMILIAL HEMOPHAGOCYTIC LYMPHOHISTIOCYTOSIS, PRF1-RELATED (PRF1) negative  
 FAMILIAL HEMOPHAGOCYTIC LYMPHOHISTIOCYTOSIS, STX11-RELATED (STX11) negative  
 FAMILIAL HEMOPHAGOCYTIC LYMPHOHISTIOCYTOSIS, STXB2-RELATED (STXB2) negative  
 FAMILIAL HEMOPHAGOCYTIC LYMPHOHISTIOCYTOSIS, UNC13D-RELATED (UNC13D) negative  
 FAMILIAL HYPERCHOLESTEROLEMIA, LDLRAP1-RELATED (LDLRAP1) negative  
 FAMILIAL HYPERCHOLESTEROLEMIA, LDLR-RELATED (LDLR) negative  
 FAMILIAL HYPERINSULINISM, ABCC8-RELATED (ABCC8) negative  
 FAMILIAL NEPHROGENIC DIABETES INSIPIDUS, AQP2-RELATED (AQP2) negative  
 FANCONI ANEMIA, GROUP A (FANCA) negative  
 FANCONI ANEMIA, GROUP C (FANCC) negative  
 FANCONI ANEMIA, GROUP D2 (FANCD2) negative  
 FANCONI ANEMIA, GROUP E (FANCE) negative  
 FANCONI ANEMIA, GROUP F (FANCF) negative  
 FANCONI ANEMIA, GROUP G (FANCG) negative  
 FANCONI ANEMIA, GROUP I (FANCI) negative  
 FANCONI ANEMIA, GROUP J (BRIP1) negative  
 FANCONI ANEMIA, GROUP L (FANCL) negative  
 FARBER LIPOGRANULOMATOSIS (ASA1) negative  
 FOVEAL HYPOPLASIA (SLC38A8) negative  
 FRASER SYNDROME 3, GRIP1-RELATED (GRIP1) negative  
 FRASER SYNDROME, FRAS1-RELATED (FRAS1) negative  
 FRASER SYNDROME, FREM2-RELATED (FREM2) negative  
 FRIEDREICH ATAXIA (FXN) negative  
 FRUCTOSE-1,6-BISPHOSPHATASE DEFICIENCY (FBP1) negative  
 FUCOSIDOSIS, FUCA1-RELATED (FUCA1) negative  
 FUMARASE DEFICIENCY (FH) negative

**G**

GABA-TRANSAMINASE DEFICIENCY (ABAT) negative  
 GALACTOKINASE DEFICIENCY (GALACTOSEMIA, TYPE II) (GALK1) negative  
 GALACTOSEMIA (GALT) negative  
 GALACTOSIALIDOSIS (CTSA) negative  
 GAUCHER DISEASE (GBA) negative  
 GCH1-RELATED CONDITIONS (GCH1) negative  
 GDF5-RELATED CONDITIONS (GDF5) negative  
 GERODERMA OSTEODYSPLASTICA (GORAB) negative  
 GITELMAN SYNDROME (SLC12A3) negative  
 GLANZMANN THROMBASTHENIA (ITGB3) negative  
 GLUTARIC ACIDEMIA, TYPE 1 (GCDH) negative  
 GLUTARIC ACIDEMIA, TYPE 2A (ETFA) negative  
 GLUTARIC ACIDEMIA, TYPE 2B (ETFB) negative  
 GLUTARIC ACIDEMIA, TYPE 2C (ETFDH) negative  
 GLUTATHIONE SYNTHETASE DEFICIENCY (GSS) negative  
 GLYCINE ENCEPHALOPATHY, AMT-RELATED (AMT) negative  
 GLYCINE ENCEPHALOPATHY, GLDC-RELATED (GLDC) negative  
 GLYCOGEN STORAGE DISEASE TYPE 5 (McArdle Disease) (PYGM) negative  
 GLYCOGEN STORAGE DISEASE TYPE IXB (PHKB) negative  
 GLYCOGEN STORAGE DISEASE TYPE IXC (PHKG2) negative  
 GLYCOGEN STORAGE DISEASE, TYPE 1a (G6PC) negative  
 GLYCOGEN STORAGE DISEASE, TYPE 1b (SLC37A4) negative  
 GLYCOGEN STORAGE DISEASE, TYPE 2 (POMPE DISEASE) (GAA) negative  
 GLYCOGEN STORAGE DISEASE, TYPE 3 (AGL) negative  
 GLYCOGEN STORAGE DISEASE, TYPE 4 (GBE1) negative  
 GLYCOGEN STORAGE DISEASE, TYPE 7 (PFKM) negative

GRACILE SYNDROME (BCS1L) negative

GUANIDINOACETATE METHYLTRANSFERASE DEFICIENCY (GAMT) negative

**H**

HARLEQUIN ICHTHYOSIS (ABCA12) negative  
 HEME OXYGENASE 1 DEFICIENCY (HMOX1) negative  
 HEMOCHROMATOSIS TYPE 2A (HFE2) negative  
 HEMOCHROMATOSIS, TYPE 3, TFR2-Related (TFR2) negative  
 HEPATOCEREBRAL MITOCHONDRIAL DNA DEPLETION SYNDROME, MPV17-RELATED (MPV17) negative  
 HEREDITARY FRUCTOSE INTOLERANCE (ALDOB) negative  
 HEREDITARY HEMOCHROMATOSIS TYPE 2B (HAMP) negative  
 HEREDITARY SPASTIC PARAPARESIS, TYPE 49 (TECPR2) negative  
 HEREDITARY SPASTIC PARAPLEGIA, CYP7B1-RELATED (CYP7B1) negative  
 HERMANSKY-PUDLAK SYNDROME, AP3B1-RELATED (AP3B1) negative  
 HERMANSKY-PUDLAK SYNDROME, BLOC1S3-RELATED (BLOC1S3) negative  
 HERMANSKY-PUDLAK SYNDROME, BLOC1S6-RELATED (BLOC1S6) negative  
 HERMANSKY-PUDLAK SYNDROME, HPS1-RELATED (HPS1) negative  
 HERMANSKY-PUDLAK SYNDROME, HPS3-RELATED (HPS3) negative  
 HERMANSKY-PUDLAK SYNDROME, HPS4-RELATED (HPS4) negative  
 HERMANSKY-PUDLAK SYNDROME, HPS5-RELATED (HPS5) negative  
 HERMANSKY-PUDLAK SYNDROME, HPS6-RELATED (HPS6) negative  
 HOLOCARBOXYLASE SYNTHETASE DEFICIENCY (HLCs) negative  
 HOMOCYSTINURIA AND MEGALOBLASTIC ANEMIA TYPE CBLG (MTR) negative  
 HOMOCYSTINURIA DUE TO DEFICIENCY OF MTHFR (MTHFR) negative  
 HOMOCYSTINURIA, CBS-RELATED (CBS) negative  
 HOMOCYSTINURIA, cbsE (MTRR) negative  
 HYDROLETHALUS SYNDROME (HYLS1) negative  
 HYPER-IGM IMMUNODEFICIENCY (CD40) negative  
 HYPERORNITHINEMIA-HYPERAMMONEMIA-HOMOCITRULLINURIA (HHH SYNDROME) (SLC2A15) negative  
 HYPERPHOSPHATEMIC FAMILIAL TUMORAL CALCIOSIS, GALNT3-RELATED (GALNT3) negative  
 HYPOMYELINATING LEUKODYSTROPHY 12 (VPS11) negative  
 HYPOPHOSPHATASIA, ALPL-RELATED (ALPL) negative

**I**

IMERSLUND-GRÄSBECK SYNDROME 2 (AMN) negative  
 IMMUNODEFICIENCY-CENTROMERIC INSTABILITY-FACIAL ANOMALIES (ICF) SYNDROME, DNMT3B-RELATED (DNMT3B) negative  
 IMMUNODEFICIENCY-CENTROMERIC INSTABILITY-FACIAL ANOMALIES (ICF) SYNDROME, ZBTB24-RELATED (ZBTB24) negative  
 INCLUSION BODY MYOPATHY 2 (GNE) negative  
 INFANTILE CEREBRAL AND CEREBELLAR ATROPHY (MED17) negative  
 INFANTILE NEPHRONOPHTHISIS (INV5) negative  
 INFANTILE NEUROAXONAL DYSTROPHY (PLA2G6) negative  
 ISOLATED ECTOPIA LENTIS (ADAMTS4) negative  
 ISOLATED SULFITE OXIDASE DEFICIENCY (SUOX) negative  
 ISOLATED THYROID-STIMULATING HORMONE DEFICIENCY (TSHB) negative  
 ISOVALERIC ACIDEMIA (IVD) negative

**J**

JOHANSON-BLIZZARD SYNDROME (UBR1) negative  
 JOUBERT SYNDROME 2 / MECKEL SYNDROME 2 (TMEM216) negative  
 JOUBERT SYNDROME AND RELATED DISORDERS (JSRD), TMEM67-RELATED (TMEM67) negative  
 JOUBERT SYNDROME, AHI1-RELATED (AHI1) negative  
 JOUBERT SYNDROME, ARL13B-RELATED (ARL13B) negative  
 JOUBERT SYNDROME, B9D1-RELATED (B9D1) negative  
 JOUBERT SYNDROME, B9D2-RELATED (B9D2) negative  
 JOUBERT SYNDROME, C2CD3-RELATED / OROFACIODIGITAL SYNDROME 14 (C2CD3) negative  
 JOUBERT SYNDROME, CC2D2A-RELATED / COACH SYNDROME (CC2D2A) negative  
 JOUBERT SYNDROME, CEP104-RELATED (CEP104) negative  
 JOUBERT SYNDROME, CEP120-RELATED / SHORT-RIB THORACIC DYSPLASIA 13 WITH OR WITHOUT POLYDACTYLY (CEP120) negative  
 JOUBERT SYNDROME, CEP41-RELATED (CEP41) negative  
 JOUBERT SYNDROME, CPLANE1-RELATED / OROFACIODIGITAL SYNDROME 6 (CPLANE1) negative  
 JOUBERT SYNDROME, CSPP1-RELATED (CSPP1) negative  
 JOUBERT SYNDROME, INPP5E-RELATED (INPP5E) negative

JUNCTIONAL EPIDERMOLYSIS BULLOSA, COL17A1-RELATED (COL17A1) negative  
 JUNCTIONAL EPIDERMOLYSIS BULLOSA, ITGA6-RELATED (ITGA6) negative  
 JUNCTIONAL EPIDERMOLYSIS BULLOSA, ITGB4-RELATED (ITGB4) negative  
 JUNCTIONAL EPIDERMOLYSIS BULLOSA, LAMB3-RELATED (LAMB3) negative  
 JUNCTIONAL EPIDERMOLYSIS BULLOSA, LAMC2-RELATED (LAMC2) negative  
 JUNCTIONAL EPIDERMOLYSIS BULLOSA/LARYNGOONYCHOCUTANEOUS SYNDROME, LAMA3-RELATED (LAMA3) negative

**K**

KRABBE DISEASE (GALC) negative

**L**

LAMELLAR ICHTHYOSIS, TYPE 1 (TGM1) negative

**Patient Information**

Patient Name: [REDACTED]

**Test Information**

Ordering Physician: [REDACTED]



Clinic Information: [REDACTED]

Date Of Birth: [REDACTED]

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Report Date: [REDACTED]

**L**

LERON SYNDROME (GHR) **negative**  
 LEBER CONGENITAL AMAUROSIS 2 (RPE65) **negative**  
 LEBER CONGENITAL AMAUROSIS TYPE AIPL1 (AIPL1) **negative**  
 LEBER CONGENITAL AMAUROSIS TYPE GUCY2D (GUCY2D) **negative**  
 LEBER CONGENITAL AMAUROSIS TYPE TULP1 (TULP1) **negative**  
 LEBER CONGENITAL AMAUROSIS, IQCB1-RELATED/SENIOR-LOKEN SYNDROME 5 (IQCB1) **negative**  
 LEBER CONGENITAL AMAUROSIS, TYPE CEP290 (CEP290) **negative**  
 LEBER CONGENITAL AMAUROSIS, TYPE LCA5 (LCA5) **negative**  
 LEBER CONGENITAL AMAUROSIS, TYPE RDH12 (RDH12) **negative**  
 LEIGH SYNDROME, FRENCH-CANADIAN TYPE (LRPPRC) **negative**  
 LETHAL CONGENITAL CONTRACTURE SYNDROME 1 (GLE1) **negative**  
 LEUKOENCEPHALOPATHY WITH VANISHING WHITE MATTER (EIF2B5) **negative**  
 LEUKOENCEPHALOPATHY WITH VANISHING WHITE MATTER, EIF2B1-RELATED (EIF2B1) **negative**  
 LEUKOENCEPHALOPATHY WITH VANISHING WHITE MATTER, EIF2B2-RELATED (EIF2B2) **negative**  
 LEUKOENCEPHALOPATHY WITH VANISHING WHITE MATTER, EIF2B3-RELATED (EIF2B3) **negative**  
 LEUKOENCEPHALOPATHY WITH VANISHING WHITE MATTER, EIF2B4-RELATED (EIF2B4) **negative**  
 LIG4 SYNDROME (LIG4) **negative**  
 LIMB-GIRDLE MUSCULAR DYSTROPHY TYPE 8 (TRIM32) **negative**  
 LIMB-GIRDLE MUSCULAR DYSTROPHY, TYPE 2A (CAPN3) **negative**  
 LIMB-GIRDLE MUSCULAR DYSTROPHY, TYPE 2B (DYSF) **negative**  
 LIMB-GIRDLE MUSCULAR DYSTROPHY, TYPE 2C (SGCG) **negative**  
 LIMB-GIRDLE MUSCULAR DYSTROPHY, TYPE 2D (SGCA) **negative**  
 LIMB-GIRDLE MUSCULAR DYSTROPHY, TYPE 2E (SGCB) **negative**  
 LIMB-GIRDLE MUSCULAR DYSTROPHY, TYPE 2F (SGCD) **negative**  
 LIMB-GIRDLE MUSCULAR DYSTROPHY, TYPE 2I (FKRP) **negative**  
 LIPOAMIDE DEHYDROGENASE DEFICIENCY (DIHYDROLIPOAMIDE DEHYDROGENASE DEFICIENCY) (DLD) **negative**  
 LIPOID ADRENAL HYPERPLASIA (STAR) **negative**  
 LIPOPROTEIN LIPASE DEFICIENCY (LPL) **negative**  
 LONG CHAIN 3-HYDROXYACYL-COA DEHYDROGENASE DEFICIENCY (HADHA) **negative**  
 LRAT-RELATED CONDITIONS (LRAT) **negative**  
 LUNG DISEASE, IMMUNODEFICIENCY, AND CHROMOSOME BREAKAGE SYNDROME (LICS) (NSMCE3) **negative**  
 LYSINURIC PROTEIN INTOLERANCE (SLC7A7) **negative**

**M**

MALONYL-COA DECARBOXYLASE DEFICIENCY (MLYCD) **negative**  
 MAPLE SYRUP URINE DISEASE, TYPE 1A (BCKDHA) **negative**  
 MAPLE SYRUP URINE DISEASE, TYPE 1B (BCKDHB) **negative**  
 MAPLE SYRUP URINE DISEASE, TYPE 2 (DBT) **negative**  
 MCKUSICK-KAUFMAN SYNDROME (MKKS) **negative**  
 MECKEL SYNDROME 7/NEPHRONOPHTHISIS 3 (NPHP3) **negative**  
 MECKEL-GRUBER SYNDROME, TYPE 1 (MKS1) **negative**  
 MECR-RELATED NEUROLOGIC DISORDER (MECR) **negative**  
 MEDIUM CHAIN ACYL-CoA DEHYDROGENASE DEFICIENCY (ACADM) **negative**  
 MEDNIK SYNDROME (AP1S1) **negative**  
 MEGALENCEPHALIC LEUKOENCEPHALOPATHY WITH SUBCORTICAL CYSTS (MLC1) **negative**  
 MEROSIN-DEFICIENT MUSCULAR DYSTROPHY (LAMA2) **negative**  
 METABOLIC ENCEPHALOPATHY AND ARRHYTHMIAS, TANGO2-RELATED (TANGO2) **negative**  
 METACHROMATIC LEUKODYSTROPHY, ARSA-RELATED (ARSA) **negative**  
 METACHROMATIC LEUKODYSTROPHY, PSAP-RELATED (PSAP) **negative**  
 METHYLMALONIC ACIDEMIA AND HOMOCYSTINURIA TYPE CBL (LMBRD1) **negative**  
 METHYLMALONIC ACIDEMIA, MCEE-RELATED (MCEE) **negative**  
 METHYLMALONIC ACIDURIA AND HOMOCYSTINURIA, TYPE CBLC (MMACHC) **negative**  
 METHYLMALONIC ACIDURIA AND HOMOCYSTINURIA, TYPE CblD (MMADHC) **negative**  
 METHYLMALONIC ACIDURIA, MMAA-RELATED (MMAA) **negative**  
 METHYLMALONIC ACIDURIA, MMAB-RELATED (MMAB) **negative**  
 METHYLMALONIC ACIDURIA, TYPE MUT(0) (MUT) **negative**  
 MEVALONIC KINASE DEFICIENCY (MVK) **negative**  
 MICROCEPHALIC OSTEODYSPLASTIC PRIMORDIAL DWARFISM TYPE II (PCNT) **negative**  
 MICROPHTHALMIA / ANOPHTHALMIA, VSX2-RELATED (VSX2) **negative**  
 MITOCHONDRIAL COMPLEX 1 DEFICIENCY, ACAD9-RELATED (ACAD9) **negative**  
 MITOCHONDRIAL COMPLEX 1 DEFICIENCY, NDUFAF5-RELATED (NDUFAF5) **negative**  
 MITOCHONDRIAL COMPLEX 1 DEFICIENCY, NDUFS6-RELATED (NDUFS6) **negative**  
 MITOCHONDRIAL COMPLEX I DEFICIENCY, NUCLEAR TYPE 1 (NDUFS4) **negative**  
 MITOCHONDRIAL COMPLEX I DEFICIENCY, NUCLEAR TYPE 10 (NDUFAF2) **negative**  
 MITOCHONDRIAL COMPLEX I DEFICIENCY, NUCLEAR TYPE 17 (NDUFAF6) **negative**  
 MITOCHONDRIAL COMPLEX I DEFICIENCY, NUCLEAR TYPE 19 (FOXRED1) **negative**  
 MITOCHONDRIAL COMPLEX I DEFICIENCY, NUCLEAR TYPE 3 (NDUFS7) **negative**  
 MITOCHONDRIAL COMPLEX I DEFICIENCY, NUCLEAR TYPE 4 (NDUFS1) **negative**  
 MITOCHONDRIAL COMPLEX IV DEFICIENCY, NUCLEAR TYPE 2, SCO2-RELATED (SCO2) **negative**  
 MITOCHONDRIAL COMPLEX IV DEFICIENCY, NUCLEAR TYPE 6 (COX15) **negative**  
 MITOCHONDRIAL DNA DEPLETION SYNDROME 2 (TK2) **negative**

MITOCHONDRIAL DNA DEPLETION SYNDROME 3 (DGUOK) **negative**MITOCHONDRIAL MYOPATHY AND SIDEROBLASTIC ANEMIA (MLASA1) (PUS1) **negative**  
 MITOCHONDRIAL TRIFUNCTIONAL PROTEIN DEFICIENCY, HADHB-RELATED (HADHB) **negative**MOLYBDENUM COFACTOR DEFICIENCY TYPE B (MOC52) **negative**MOLYBDENUM COFACTOR DEFICIENCY, TYPE A (MOC51) **negative**MUCOLIPIDOSIS II/III A (GNPTAB) **negative**MUCOLIPIDOSIS III GAMMA (GNPTG) **negative**MUCOLIPIDOSIS, TYPE IV (MCOLN1) **negative**MUCOPOLYSACCHARIDOSIS, TYPE I ( HURLER SYNDROME ) (IDUA) **negative**MUCOPOLYSACCHARIDOSIS, TYPE III A ( SANFILIPPO A ) (SGSH) **negative**MUCOPOLYSACCHARIDOSIS, TYPE III B ( SANFILIPPO B ) (NAGLU) **negative**MUCOPOLYSACCHARIDOSIS, TYPE III C ( SANFILIPPO C ) (HGSNAT) **negative**MUCOPOLYSACCHARIDOSIS, TYPE III D ( SANFILIPPO D ) (GNS) **negative**MUCOPOLYSACCHARIDOSIS, TYPE IV A ( MORQUO SYNDROME ) (GALNS) **negative**MUCOPOLYSACCHARIDOSIS, TYPE IV B/GM1 GANGLIOSIDOSIS (GLB1) **negative**MUCOPOLYSACCHARIDOSIS, TYPE IX (HYAL1) **negative**MUCOPOLYSACCHARIDOSIS, TYPE VI ( MAROTEAUX-LAMY ) (ARSB) **negative**MUCOPOLYSACCHARIDOSIS, TYPE VII (GUSB) **negative**MULIBREY NANISM (TRIM37) **negative**MULTIPLE PTERYGIUM SYNDROME, CHRNG-RELATED/ESCOBAR SYNDROME (CHRNG) **negative**MULTIPLE SULFATASE DEFICIENCY (SUMF1) **negative**MUSCLE-EYE-BRAIN DISEASE, POMGNT1-RELATED (POMGNT1) **negative**MUSCULAR DYSTROPHY-DYSTROGLYCANOPATHY (RXYLT1) **negative**MUSK-RELATED CONGENITAL MYASTHENIC SYNDROME (MUSK) **negative**MYONEUROGASTROINTESTINAL ENCEPHALOPATHY (MNGIE) (TYMP) **negative**MYOTONIA CONGENITA (CLCN1) **negative****N**N-ACETYLGlutamate synthase deficiency (NAGS) **negative**NEMALINE MYOPATHY, NEB-RELATED (NEB) **negative**NEPHRONOPHTHISIS 1 (NPHP1) **negative**NEURONAL CEROID LIPOFUSCINOSIS, CLN5-RELATED (CLN5) **negative**NEURONAL CEROID LIPOFUSCINOSIS, CLN6-RELATED (CLN6) **negative**NEURONAL CEROID LIPOFUSCINOSIS, CLN8-RELATED (CLN8) **negative**NEURONAL CEROID LIPOFUSCINOSIS, MFSD8-RELATED (MFSD8) **negative**NEURONAL CEROID LIPOFUSCINOSIS, PPT1-RELATED (PPT1) **negative**NEURONAL CEROID LIPOFUSCINOSIS, TPP1-RELATED (TPP1) **negative**NGLY1-CONGENITAL DISORDER OF GLYCOSYLATION (NGLY1) **negative**NIEMANN-PICK DISEASE, TYPE C1 / D (NPC1) **negative**NIEMANN-PICK DISEASE, TYPE C2 (NPC2) **negative**NIEMANN-PICK DISEASE, TYPES A / B (SMPD1) **negative**NIJMEGEN BREAKAGE SYNDROME (NBN) **negative**NON-SYNDROMIC HEARING LOSS, GJB2-RELATED (GJB2) **negative**NON-SYNDROMIC HEARING LOSS, MYO15A-RELATED (MYO15A) **negative**NONSYNDROMIC HEARING LOSS, OTOA-RELATED (OTOA) **negative**NONSYNDROMIC HEARING LOSS, OTOF-RELATED (OTOF) **negative**NONSYNDROMIC HEARING LOSS, PJVK-RELATED (PJVK) **negative**NONSYNDROMIC HEARING LOSS, SYNE4-RELATED (SYNE4) **negative**NONSYNDROMIC HEARING LOSS, TMC1-RELATED (TMC1) **negative**NONSYNDROMIC HEARING LOSS, TMPRSS3-RELATED (TMPRSS3) **negative**NONSYNDROMIC INTELLECTUAL DISABILITY (CC2D1A) **negative**NONMOPHOSPHATEMIC TUMORAL CALCINOSIS (SAMD9) **negative****O**OCULOCUTANEOUS ALBINISM TYPE III (TYRP1) **negative**OCULOCUTANEOUS ALBINISM TYPE IV (SLC45A2) **negative**OCULOCUTANEOUS ALBINISM, OCA2-RELATED (OCA2) **negative**OCULOCUTANEOUS ALBINISM, TYPES 1A AND 1B (TYR) **negative**ODONTO-ONYCHO-DERMAL DYSPLASIA / SCHOPF-SCHULZ-PASSARGE SYNDROME (WNT10A) **negative**OMENN SYNDROME, RAG2-RELATED (RAG2) **negative**ORNITHINE AMINOTRANSFERASE DEFICIENCY (OAT) **negative**OSTEOGENESIS IMPERFECTA TYPE VII (CRTAP) **negative**OSTEOGENESIS IMPERFECTA TYPE VIII (P3H1) **negative**OSTEOGENESIS IMPERFECTA TYPE XI (FKBP10) **negative**OSTEOGENESIS IMPERFECTA TYPE XIII (BMP1) **negative**OSTEOPETROSIS, INFANTILE MALIGNANT, TCIRG1-RELATED (TCIRG1) **negative**OSTEOPETROSIS, OSTM1-RELATED (OSTM1) **negative****P**PANTOTHENATE KINASE-ASSOCIATED NEURODEGENERATION (PANK2) **negative**PAPILLON LEFEVRE SYNDROME (CTSC) **negative**PARKINSON DISEASE 15 (FBXO7) **negative**PENDRED SYNDROME (SLC26A4) **negative**PERLMAN SYNDROME (DIS3L2) **negative**PGM3-CONGENITAL DISORDER OF GLYCOSYLATION (PGM3) **negative**PHENYLKETONURIA (PAH) **negative**PIGN-CONGENITAL DISORDER OF GLYCOSYLATION (PIGN) **negative**PITUITARY HORMONE DEFICIENCY, COMBINED 3 (LHX3) **negative**POLG-RELATED DISORDERS (POLG) **negative**

**Patient Information**

Patient Name: [REDACTED]

**Test Information**

Ordering Physician: [REDACTED]



Clinic Information: [REDACTED]

Date Of Birth: [REDACTED]

Case File ID: [REDACTED]

Report Date: [REDACTED]

**P**

POLYCYSTIC KIDNEY DISEASE, AUTOSOMAL RECESSIVE (PKHD1) negative  
 PONTOCEREBELLAR HYPOPLASIA, EXOSC3-RELATED (EXOSC3) negative  
 PONTOCEREBELLAR HYPOPLASIA, RARS2-RELATED (RARS2) negative  
 PONTOCEREBELLAR HYPOPLASIA, TSEN2-RELATED (TSEN2) negative  
 PONTOCEREBELLAR HYPOPLASIA, TSEN54-RELATED (TSEN54) negative  
 PONTOCEREBELLAR HYPOPLASIA, TYPE 1A (VRK1) negative  
 PONTOCEREBELLAR HYPOPLASIA, TYPE 2D (SEPSECS) negative  
 PONTOCEREBELLAR HYPOPLASIA, VPS53-RELATED (VPS53) negative  
 PRIMARY CILIARY DYSKINESIA, CCDC103-RELATED (CCDC103) negative  
 PRIMARY CILIARY DYSKINESIA, CCDC39-RELATED (CCDC39) negative  
 PRIMARY CILIARY DYSKINESIA, DNAH11-RELATED (DNAH11) negative  
 PRIMARY CILIARY DYSKINESIA, DNAH5-RELATED (DNAH5) negative  
 PRIMARY CILIARY DYSKINESIA, DNA11-RELATED (DNA11) negative  
 PRIMARY CILIARY DYSKINESIA, DNA12-RELATED (DNA12) negative  
 PRIMARY CONGENITAL GLAUCOMA/PETERS ANOMALY (CYP1B1) negative  
 PRIMARY HYPEROXALURIA, TYPE 1 (AGXT) negative  
 PRIMARY HYPEROXALURIA, TYPE 2 (GRHPR) negative  
 PRIMARY HYPEROXALURIA, TYPE 3 (HOGA1) negative  
 PRIMARY MICROCEPHALY 1, AUTOSOMAL RECESSIVE (MCPH1) negative  
 PROGRESSIVE EARLY-ONSET ENCEPHALOPATHY WITH BRAIN ATROPHY AND THIN CORPUS CALLOSUM (TBCD) negative  
 PROGRESSIVE FAMILIAL INTRAHEPATIC CHOLESTASIS, ABCB4-RELATED (ABCB4) negative  
 PROGRESSIVE FAMILIAL INTRAHEPATIC CHOLESTASIS, TYPE 1 (PFIC1) (ATP8B1) negative  
 PROGRESSIVE FAMILIAL INTRAHEPATIC CHOLESTASIS, TYPE 2 (ABCB11) negative  
 PROGRESSIVE FAMILIAL INTRAHEPATIC CHOLESTASIS, TYPE 4 (PFIC4) (TJP2) negative  
 PROGRESSIVE PSEUDORHEUMATOID DYSPLASIA (CCN6) negative  
 PROLIDASE DEFICIENCY (PEPD) negative  
 PROPIONIC ACIDEMIA, PCCA-RELATED (PCCA) negative  
 PROPIONIC ACIDEMIA, PCCB-RELATED (PCCB) negative  
 PSEUDOXANTHOMA ELASTICUM (ABCC6) negative  
 PTERIN-4 ALPHA-CARBINOLAMINE DEHYDRATASE (PCD) DEFICIENCY (PCBD1) negative  
 PYCNOYDYSOSTOSIS (CTSK) negative  
 PYRIDOXAL 5'-PHOSPHATE-DEPENDENT EPILEPSY (PNPO) negative  
 PYRIDOXINE-DEPENDENT EPILEPSY (ALDH7A1) negative  
 PYRUVATE CARBOXYLASE DEFICIENCY (PC) negative  
 PYRUVATE DEHYDROGENASE DEFICIENCY, PDHB-RELATED (PDHB) negative

**R**

REFSUM DISEASE, PHYH-RELATED (PHYH) negative  
 RENAL TUBULAR ACIDOSIS AND DEAFNESS, ATP6V1B1-RELATED (ATP6V1B1) negative  
 RENAL TUBULAR ACIDOSIS, PROXIMAL, WITH OCULAR ABNORMALITIES AND MENTAL RETARDATION (SLC4A4) negative  
 RETINITIS PIGMENTOSA 25 (EYS) negative  
 RETINITIS PIGMENTOSA 26 (CERKL) negative  
 RETINITIS PIGMENTOSA 28 (FAM161A) negative  
 RETINITIS PIGMENTOSA 36 (PRCD) negative  
 RETINITIS PIGMENTOSA 59 (DHDDS) negative  
 RETINITIS PIGMENTOSA 62 (MAK) negative  
 RHIZOMELIC CHONDRODYSPLASIA PUNCTATA, TYPE 1 (PEX7) negative  
 RHIZOMELIC CHONDRODYSPLASIA PUNCTATA, TYPE 2 (GNPAT) negative  
 RHIZOMELIC CHONDRODYSPLASIA PUNCTATA, TYPE 3 (AGPS) negative  
 RLRP1-RELATED RETINOPATHY (RLRP1) negative  
 ROBERTS SYNDROME (ESCO2) negative  
 RYR1-RELATED CONDITIONS (RYR1) negative

**S**

SALLA DISEASE (SLC17A5) negative  
 SANDHOFF DISEASE (HEXB) negative  
 SCHIMKE IMMUNOSSEOUS DYSPLASIA (SMARCAL1) negative  
 SCHINDLER DISEASE (NAGA) negative  
 SEGAWA SYNDROME, TH-RELATED (TH) negative  
 SENIOR-LOKEN SYNDROME 4/NEPHRONOPHTHISIS 4 (NPHP4) negative  
 SEPIAFTERIN REDUCTASE DEFICIENCY (SPR) negative  
 SEVERE COMBINED IMMUNODEFICIENCY (SCID), CD3D-RELATED (CD3D) negative  
 SEVERE COMBINED IMMUNODEFICIENCY (SCID), CD3E-RELATED (CD3E) negative  
 SEVERE COMBINED IMMUNODEFICIENCY (SCID), FOXN1-RELATED (FOXN1) negative  
 SEVERE COMBINED IMMUNODEFICIENCY (SCID), IKBKB-RELATED (IKBKB) negative  
 SEVERE COMBINED IMMUNODEFICIENCY (SCID), IL7R-RELATED (IL7R) negative  
 SEVERE COMBINED IMMUNODEFICIENCY (SCID), JAK3-RELATED (JAK3) negative  
 SEVERE COMBINED IMMUNODEFICIENCY (SCID), PTPRC-RELATED (PTPRC) negative  
 SEVERE COMBINED IMMUNODEFICIENCY (SCID), RAG1-RELATED (RAG1) negative  
 SEVERE COMBINED IMMUNODEFICIENCY, ADA-Related (ADA) negative  
 SEVERE COMBINED IMMUNODEFICIENCY, TYPE ATHABASKAN (DCLRE1C) negative  
 (DYNCH2H1) negative  
 SHWACHMAN-DIAMOND SYNDROME, SBDS-RELATED (SBDS) negative  
 SIALIDOSIS (NEU1) negative  
 SJÖGREN-LARSSON SYNDROME (ALDH3A2) negative  
 SMITH-LEMLI-OPITZ SYNDROME (DHCR7) negative  
 SPASTIC PARAPLEGIA, TYPE 15 (ZFYVE26) negative

SPASTIC TETRAPLEGIA, THIN CORPUS CALLOSUM, AND PROGRESSIVE MICROCEPHALY (SPATCCM) (SLC1A4) negative  
 SPG11-RELATED CONDITIONS (SPG11) negative  
 SPINAL MUSCULAR ATROPHY (SMN1) negative SMN1: Two copies; g.27134T>G: absent; the absence of the g.27134T>G variant decreases the chance to be a silent (2+0) carrier.  
 SPINAL MUSCULAR ATROPHY WITH RESPIRATORY DISTRESS TYPE 1 (IGHMBP2) negative  
 SPINOCEREBELLAR ATAXIA, AUTOSOMAL RECESSIVE 10 (ANO10) negative  
 SPINOCEREBELLAR ATAXIA, AUTOSOMAL RECESSIVE 12 (WWOX) negative  
 SPONDYLOCOLSTAL DYSOSTOSIS 1 (DLL3) negative  
 SPONDYLOTHORACIC DYSOSTOSIS, MESP2-Related (MESP2) negative  
 STEEL SYNDROME (COL27A1) negative  
 STEROID-RESISTANT NEPHROTIC SYNDROME (NPHS2) negative  
 STUVE-WIEDEMANN SYNDROME (LIFR) negative  
 SURF1-RELATED CONDITIONS (SURF1) negative  
 SURFACTANT DYSFUNCTION, ABCA3-RELATED (ABCA3) negative

**T**

TAY-SACHS DISEASE (HEXA) negative  
 TBCE-RELATED CONDITIONS (TBCE) negative  
 THIAMINE-RESPONSIVE MEGALOBLASTIC ANEMIA SYNDROME (SLC19A2) negative  
 THYROID DYSHORMONOGENESIS 1 (SLC5A5) negative  
 THYROID DYSHORMONOGENESIS 2A (TPO) negative  
 THYROID DYSHORMONOGENESIS 3 (TG) negative  
 THYROID DYSHORMONOGENESIS 6 (DUOX2) see first page  
 TRANSCOBALAMIN II DEFICIENCY (TCN2) negative  
 TRICHOHEPATOENTERIC SYNDROME, SKIC2-RELATED (SKIC2) negative  
 TRICHOHEPATOENTERIC SYNDROME, TTC37-RELATED (TTC37) negative  
 TRICHOHYDROSTROPHY 1/XERODERMA PIGMENTOSUM, GROUP D (ERCC2) negative  
 TRIMETHYLMALINURIA (FMO3) negative  
 TRIPLE A SYNDROME (AAAS) negative  
 TSHR-RELATED CONDITIONS (TSHR) negative  
 TYROSINEMIA TYPE III (HPD) negative  
 TYROSINEMIA, TYPE 1 (FAH) negative  
 TYROSINEMIA, TYPE 2 (TAT) negative

**U**

USHER SYNDROME, TYPE 1B (MYO7A) negative  
 USHER SYNDROME, TYPE 1C (USH1C) negative  
 USHER SYNDROME, TYPE 1D (CDH23) negative  
 USHER SYNDROME, TYPE 1F (PCDH15) negative  
 USHER SYNDROME, TYPE 1J/DEAFNESS, AUTOSOMAL RECESSIVE, 48 (CIB2) negative  
 USHER SYNDROME, TYPE 2A (USH2A) negative  
 USHER SYNDROME, TYPE 2C (ADGRV1) negative  
 USHER SYNDROME, TYPE 3 (CLRN1) negative

**V**

VERY LONG-CHAIN ACYL-CoA DEHYDROGENASE DEFICIENCY (ACADVL) negative  
 VICI SYNDROME (EPG5) negative  
 VITAMIN D-DEPENDENT RICKETS, TYPE 1A (CYP27B1) negative  
 VITAMIN D-RESISTANT RICKETS TYPE 2A (VDR) negative  
 VLDLR-ASSOCIATED CEREBELLAR HYPOPLASIA (VLDLR) negative

**W**

WALKER-WARBURG SYNDROME, CRPPA-RELATED (CRPPA) negative  
 WALKER-WARBURG SYNDROME, FKTN-RELATED (FKTN) negative  
 WALKER-WARBURG SYNDROME, LARGE1-RELATED (LARGE1) negative  
 WALKER-WARBURG SYNDROME, POMT1-RELATED (POMT1) negative  
 WALKER-WARBURG SYNDROME, POMT2-RELATED (POMT2) negative  
 WARSAW BREAKAGE SYNDROME (DDX11) negative  
 WERNER SYNDROME (WRN) negative  
 WILSON DISEASE (ATP7B) negative  
 WOLCOTT-RALLISON SYNDROME (EIF2AK3) negative  
 WOLMAN DISEASE (LIPA) negative  
 WOODHOUSE-SAKATI SYNDROME (DCAF17) negative

**X**

XERODERMA PIGMENTOSUM VARIANT TYPE (POLH) negative  
 XERODERMA PIGMENTOSUM, GROUP A (XPA) negative  
 XERODERMA PIGMENTOSUM, GROUP C (XPC) negative

**Z**

ZELLWEGER SPECTRUM DISORDER, PEX13-RELATED (PEX13) negative  
 ZELLWEGER SPECTRUM DISORDER, PEX16-RELATED (PEX16) negative  
 ZELLWEGER SPECTRUM DISORDER, PEX5-RELATED (PEX5) negative  
 ZELLWEGER SPECTRUM DISORDERS, PEX10-RELATED (PEX10) negative  
 ZELLWEGER SPECTRUM DISORDERS, PEX12-RELATED (PEX12) negative  
 ZELLWEGER SPECTRUM DISORDERS, PEX1-RELATED (PEX1) negative  
 ZELLWEGER SPECTRUM DISORDERS, PEX26-RELATED (PEX26) negative  
 ZELLWEGER SPECTRUM DISORDERS, PEX2-RELATED (PEX2) negative

**Patient Information**

Patient Name:

**Test Information**

Ordering Physician: [REDACTED]

Date Of Birth: [REDACTED]

Clinic Information:

Case File ID: [REDACTED]

Report Date:

Z

ZELLWEGER SPECTRUM DISORDERS, PEX6-RELATED (PEX6) negative



**Patient Information**

Patient Name: [REDACTED]

**Test Information**

Ordering Physician: [REDACTED]



Clinic Information: [REDACTED]

Date Of Birth: [REDACTED]

Case File ID: [REDACTED]

Report Date: [REDACTED]

**Testing Methodology, Limitations, and Comments:****Next-generation sequencing (NGS)**

Sequencing library prepared from genomic DNA isolated from a patient sample is enriched for targets of interest using standard hybridization capture protocols and PCR amplification (for targets specified below). NGS is then performed to achieve the standards of quality control metrics, including a minimum coverage of 99% of targeted regions at 20X sequencing depth. Sequencing data is aligned to human reference sequence, followed by deduplication, metric collection and variant calling (coding region +/- 20bp). Variants are then classified according to ACMGG/AMP standards of interpretation using publicly available databases including but not limited to ENSEMBL, HGMD Pro, ClinGen, ClinVar, 1000G, ESP and gnomAD. Variants predicted to be pathogenic or likely pathogenic for the specified diseases are reported. It should be noted that the data interpretation is based on our current understanding of the genes and variants at the time of reporting. Putative positive sequencing variants that do not meet internal quality standards or are within highly homologous regions are confirmed by Sanger sequencing or gene-specific long-range PCR as needed prior to reporting.

Copy Number Variant (CNV) analysis is limited to deletions involving two or more exons for all genes on the panel, in addition to specific known recurrent single-exon deletions. CNVs of small size may have reduced detection rate. This method does not detect gene inversions, single-exonic and sub-exonic deletions (unless otherwise specified), and duplications of all sizes (unless otherwise specified). Additionally, this method does not define the exact breakpoints of detected CNV events. Confirmation testing for copy number variation is performed by specific PCR, Multiplex Ligation-dependent Probe Amplification (MLPA), next generation sequencing, or other methodology.

This test may not detect certain variants due to local sequence characteristics, high/low genomic complexity, homologous sequence, or allele dropout (PCR-based assays). Variants within noncoding regions (promoter, 5'UTR, 3'UTR, deep intronic regions, unless otherwise specified), small deletions or insertions larger than 25bp, low-level mosaic variants, structural variants such as inversions, and/or balanced translocations may not be detected with this technology.

**SPECIAL NOTES**

For ABCC6, sequencing variants in exons 1-7 are not detected due to the presence of regions of high homology.

For CFTR, when the CFTR R117H variant is detected, reflex analysis of the polythymidine variations (5T, 7T and 9T) at the intron 9 branch/acceptor site of the CFTR gene will be performed. Multi-exon duplication analysis is included.

For CYP21A2, targets were enriched using long-range PCR amplification, followed by next generation sequencing. Duplication analysis will only be performed and reported when c.955C>T (p.Q319\*) is detected. Sequencing and CNV analysis may have reduced sensitivity, if variants result from complex rearrangements, in trans with a gene deletion, or CYP21A2 gene duplication on one chromosome and deletion on the other chromosome. This analysis cannot detect sequencing variants located on the CYP21A2 duplicated copy.

For DDX11, sequencing variants in exons 7-11 and CNV for the entire gene are not analyzed due to high sequence homology.

For GJB2, CNV analysis of upstream deletions of GJB6-D13S1830 (309kb deletion) and GJB6-D13S1854 (232kb deletion) is included.

For HBA1/HBA2, CNV analysis is offered to detect common deletions of -alpha3.7, -alpha4.2, --MED, --SEA, --FIL, --THAI, --alpha20.5, and/or HS-40.

For OTOA, sequencing variants in exons 25-29 and CNV in exons 21-29 are not analyzed due to high sequence homology.

For RPGRIP1L, variants in exon 23 are not detected due to assay limitation.

For SAMD9, only p.K1495E variant will be analyzed and reported.

**Friedreich Ataxia (FXN)**

The GAA repeat region of the FXN gene is assessed by trinucleotide PCR assay and capillary electrophoresis. Variances of +/-1 repeat for normal alleles and up to +/-3 repeats for premutation alleles may occur. For fully penetrant expanded alleles, the precise repeat size cannot be determined, therefore the approximate allele size is reported. Sequencing and copy number variants are analyzed by next-generation sequencing analysis.

**Friedreich Ataxia Repeat Categories**

Categories	GAA Repeat Sizes
Normal	<34
Premutation	34 - 65
Full	>65

**Patient Information**

Patient Name: [REDACTED]

**Test Information**

Ordering Physician: [REDACTED]



Clinic Information: [REDACTED]

Date Of Birth: [REDACTED]

Case File ID: [REDACTED]

Report Date: [REDACTED]

**Spinal Muscular Atrophy (SMN1)**

The total combined copy number of SMN1 and SMN2 exon 7 is quantified based on NGS read depth. The ratio of SMN1 to SMN2 is calculated based on the read depth of a single nucleotide that distinguishes these two genes in exon 7. In addition to copy number analysis, testing for the presence or absence of a single nucleotide polymorphism (g.27134T>G in intron 7 of SMN1) associated with the presence of a SMN1 duplication allele is performed using NGS.

Ethnicity	Two SMN1 copies carrier risk before g.27134T>G testing	Carrier risk after g.27134T>G testing	
		g.27134T>G ABSENT	g.27134T>G PRESENT
Caucasian	1 in 632	1 in 769	1 in 29
Ashkenazi Jewish	1 in 350	1 in 580	LIKELY CARRIER
Asian	1 in 628	1 in 702	LIKELY CARRIER
African-American	1 in 121	1 in 396	1 in 34
Hispanic	1 in 1061	1 in 1762	1 in 140

**Variant Classification**

Only pathogenic or likely pathogenic variants are reported. Other variants including benign variants, likely benign variants, variants of uncertain significance, or inconclusive variants identified during this analysis may be reported in certain circumstances. Our laboratory's variant classification criteria are based on the ACMG and internal guidelines and our current understanding of the specific genes. This interpretation may change over time as more information about a gene and/or variant becomes available. Natera and its lab partner(s) may reclassify variants at certain intervals but may not release updated reports without a specific request made to Natera by the ordering provider. Natera may disclose incidental findings if deemed clinically pertinent to the test performed.

**Negative Results**

A negative carrier screening result reduces the risk for a patient to be a carrier of a specific disease but does not completely rule out carrier status. Please visit <https://www.natera.com/panel-option/h-all/> for a table of carrier rates, detection rates, residual risks and promised variants/exons per gene. Carrier rates before and after testing vary by ethnicity and assume a negative family history for each disease screened and the absence of clinical symptoms in the patient. Any patient with a family history for a specific genetic disease will have a higher carrier risk prior to testing and, if the disease-causing mutation in their family is not included on the test, their carrier risk would remain unchanged. Genetic counseling is recommended for patients with a family history of genetic disease so that risk figures based on actual family history can be determined and discussed along with potential implications for reproduction. Horizon carrier screening has been developed to identify the reproductive risks for monogenic inherited conditions. Even when one or both members of a couple screen negative for pathogenic variants in a specific gene, the disease risk for their offspring is not zero. There is still a low risk for the condition in their offspring due to a number of different mechanisms that are not detected by Horizon including, but not limited to, pathogenic variant(s) in the tested gene or in a different gene not included on Horizon, pathogenic variant(s) in an upstream regulator, uniparental disomy, de novo mutation(s), or digenic or polygenic inheritance.

**Additional Comments**

These analyses generally provide highly accurate information regarding the patient's carrier status. Despite this high level of accuracy, it should be kept in mind that there are many potential sources of diagnostic error, including misidentification of samples, polymorphisms, or other rare genetic variants that interfere with analysis. Families should understand that rare diagnostic errors may occur for these reasons.