

### **Donor 4933**

## **Genetic Testing Summary**

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

Last Updated: 08/17/23

Donor Reported Ancestry: Indian Jewish Ancestry: No

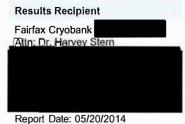
Genetic Test*	Result	Comments/Donor's Residual Risk**
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Chromosome analysis (karyotype)	Normal male karyotype	No evidence of clinically significant chromosome abnormalities
Hemoglobin evaluation	Normal hemoglobin fractionation and MCV/MCH results	Reduced risk to be a carrier for sickle cell anemia, beta thalassemia, alpha thalassemia trait (aa/ and a-/a-) and other hemoglobinopathies
Cystic Fibrosis (CF) carrier screening	Negative by genotyping of 99 mutations in the CFTR gene	1/190
Spinal Muscular Atrophy (SMA) carrier screening	Negative for deletions of exon 7 in the SMN1 gene	1/860
Hb Beta Chain-Related Hemoglobinopathy (including Beta Thalassemia and Sickle Cell Disease) by genotyping	Negative for 28 mutations tested in the HBB gene	1/180

<sup>\*</sup>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.

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





Name: 4933 4933 DOB

Ethnicity: Southeast Asian Sample Type: EDTA Blood Date of Collection: 05/12/2014 Date Rece 4 Barcode:

Indication: onor Test Type: The Counsyl Test Female

Not tested

### Counsyl Test Results Summary (Egg or Sperm Donor)

The Counsyl test (Fairfax Cryobank Fundamental Panel) uses targeted genotyping and copy number analysis as described in the methods section on page 2 to determine carrier status associated with 3 diseases. Please refer to page 3 for a complete list of diseases and genes included in this panel.



4933 4933



4933 4933's DNA test shows that he is not a carrier of any disease-causing mutation tested.



### Partner

The reproductive risk presented is based on a hypothetical pairing with a partner of the same ethnic group.

# Reproductive Risk Summary

No increased reproductive risks to highlight. Please refer to the following pages for detailed information about the results.

#### Clinical Notes

• If necessary, patients can discuss residual risks with their physician or a genetic counselor. To schedule a complimentary appointment to speak with a clinical expert about these results, please visit <a href="mailto:counsyl.com/my/consults/">counsyl.com/my/consults/</a>.





Name: 4933 4933 DOB Female

Not tested

#### **Methods and Limitations**

4933 4933: The Counsyl Test - targeted genotyping and copy number analysis.

Targeted genotyping: Targeted DNA mutation analysis is used to simultaneously determine the genotype of 127 variants associated with 2 diseases. The test is not validated for detection of homozygous mutations, and although rare, asymptomatic individuals affected by the disease may not be genotyped accurately.

Copy number analysis: Targeted copy number analysis is used to determine the copy number of exon 7 of the SMN1 gene relative to other genes. Other mutations may interfere with this analysis. Some individuals with two copies of SMN1 are carriers with two SMN1 genes on one chromosome and a SMN1 deletion on the other chromosome. In addition, a small percentage of SMA cases are caused by nondeletion mutations in the SMN1 gene. Thus, a test result of two SMN1 copies significantly reduces the risk of being a carrier; however, there is still a residual risk of being a carrier and subsequently a small risk of future affected offspring for individuals with two or more SMN1 gene copies. Some SMA cases arise as the result of de novo mutation events which will not be detected by carrier testing.

Limitations: In an unknown number of cases, nearby genetic variants may interfere with mutation detection. Other possible sources of diagnostic error include sample mix-up, trace contamination, bone marrow transplantation, blood transfusions and technical errors. If more than one variant is detected in a gene, additional studies may be necessary to determine if those variants lie on the same chromosome or different chromosomes. The Counsyl test does not fully address all inherited forms of intellectual disability, birth defects and genetic disease. A family history of any of these conditions may warrant additional evaluation. Furthermore, not all mutations will be identified in the genes analyzed and additional testing may be beneficial for some patients. For example, individuals of African, Southeast Asian, and Mediterranean ancestry are at increased risk for being carriers for hemoglobinopathies, which can be identified by CBC and hemoglobin electrophoresis or HPLC (ACOG Practice Bulletin No. 78. Obstet. Gynecol. 2007;109:229-37).

This test was developed and its performance characteristics determined by Counsyl, Inc. It has not been cleared or approved by the US Food and Drug Administration (FDA). The FDA does not require this test to go through premarket review. This test is used for clinical purposes. It should not be regarded as investigational or for research. This laboratory is certified under the Clinical Laboratory Improvement Amendments of 1988 (CLIA) as qualified to perform high-complexity clinical testing. These results are adjunctive to the ordering physician's workup. CLIA Number: #05D1102604.

Lab Directors:

H. Peter Kang, MD, MS, FCAP

Jeleux Brext

Jelena Brezo, PhD, FACMG



Name: 4933 4933 DOE Female

Not tested

#### **Diseases Tested**

Cystic Fibrosis - Gene: CFTR. Variants (99): G85E, R117H, R334W, R347P, A455E, G542\*, G551D, R553\*, R560T, R1162\*, W1282\*, N1303K, c.1521\_1523delCTT, c.1519\_1521delATC, c.2052delA, c.3528delC, c.489+1G>T, c.579+1G>T, c.1585-1G>A, c.1766+1G>A, 2789+5G>A, c.2988+1G>A, 3849+10kbC>T, E60\*, R75\*, E92\*, Y122\*, G178R, R347H, Q493\*, V520F, S549N, P574H, M1101K, D1152H, c.2012delT, c.262\_263delTT, c.313delA, c.948delT, c.3744delA, c.3773dupT, c.1680-1G>A, 3272-26A>G, c.2051\_2052delAAinsG, S549R, R117C, L206W, G330\*, T338I, R352Q, S364P, G480C, C524\*, S549R, Q552\*, A559T, G622D, R709\*, K710\*, R764\*, Q890\*, R1066C, W1089\*, Y1092X, R1158\*, S1196\*, W1204\*, Q1238\*, S1255\*, c.3067\_3072del6, c.442delA, c.531delT, c.803delA, c.805\_806delAT, c.1545\_1546delTA, 1949del84, c.1911delG, c.1923\_1931del9ins1, c.1976delA, c.3039delC, c.3536\_3539delCCAA, c.3659delC, c.1155\_1156dupTA, c.2052dupA, c.2175dupA, c.2738insG, 296\*12T>C, c.273+1G>A, 405+3A>C, c.274-1G>A, 711+5G>A, c.580-1G>T, c.1766+1G>T, 1898+5G>T, Q996, c.325\_327delTATinsG, 3849+4A>G, c.1075\_1079del5ins5. IVS8-5T allele analysis is only reported in the presence of the R117H mutation. Detection rate: Southeagt Asian 54%.

Hb Beta Chain-Related Hemoglobinopathy (Including Beta Thalassemia and Sickle Cell Disease) - Gene: HBB. Variants (28): E7V, K18\*, Q40\*, c.126\_129delCTTT, c.27dupG, IVS-II-654, IVS-II-745, c.315+1G>A, IVS-I-6, IVS-I-10, IVS-I-5, c.92+1G>A, -88C>T, -28A>G, -29A>G, c.25\_26delAA, c.217dupA, c.316-2A>C, c.316-2A>G, G25, -87C>G, E7K, W16\*, c.51deJC, c.20delA, E27K, E122Q, E122K. Detection rate: Southeast Asian 86%.

Spinal Muscular Atrophy (copy number analysis only) - Gene: SMN1. Variant (1): SMN1 copy number. Detection rate: Southeast Asian 94%.



Name: 4933 4933 DOB Female

Not tested

#### **Risk Calculations**

Below are the risk calculations for all diseases tested. Since negative results do not completely rule out the possibility of being a carrier, the residual risk represents the patient's post-test likelihood of being a carrier and the reproductive risk represents the likelihood the patient's future children could inherit each disease. These risks are inherent to all carrier screening tests, may vary by ethnicity, are predicated on a negative family history and are present even after a negative test result. Inaccurate reporting of ethnicity may cause errors in risk calculation.

Disease	4933 4933 Residual Risk	Reproductive Risk	
Cystic Fibrosis	1 in 190	1 in 66,000	
Hb Beta Chain-Related Hemoglobinopathy (Including Beta Thalassemia and Sickle Cell Disease)	1 in 180	1 in 17,000	
Spinal Muscular Atrophy	SMN1: 2 copies 1 in 860	1 in 180,000	



## Cytogenetic Report

Client Fa	irfax Cryobank -						
Address							
Reporting Phone #		Fax #		Email			
Patient name/Donor Alia	ns Donor # 4933			Patient DOB N	I/A		
Donor	# 4933-140512	40)		Specimen type P	eripheral	Blood	
Collection Dat	e 05/12/2014			Accession # 1	4-083CG		
Date Receive	d 05/13/2014						
		RESUI	TS				
CYTO	GENETIC ANAI	LYSIS			FISH		
Cells counted	20	Type of banding	GTG	I	Probe(s)	N/A	
Cells analyzed	5	Band resolution	550	Nucle	i scored	N/A	
Cells karyotyped	2						
Modal chromosome#	46						
KARYOTYPE 46,XY							

#### INTERPRETATION

Normal male karyotype

No clonal numerical or structural abnormalities were identified. This normal cytogenetic result does not exclude the possibility of the presence of subtle rearrangements beyond the technical limits of detection with this test.

Comments

Wayne S. Stanley, Ph.D., FACMG

Clinical Cytogeneticist

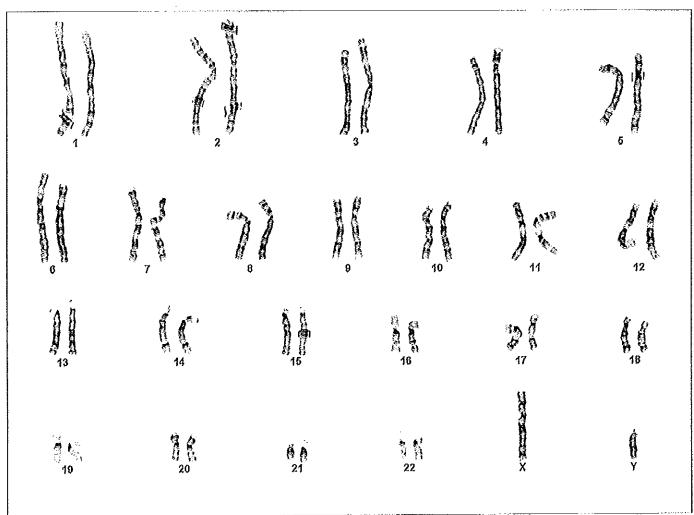
5/86/14 Date



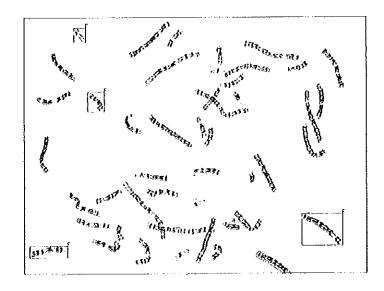
## Genetics and IVF Preimplantation Genetics Laboratory

Patient name: DONOR # 4933

Case name: 14-083CG



Case: 14-083CG Slide: B2 Cell: 6



ID, 4933

Patient Information	Specimen Information	Client Information		
ID, 4933	Specimen: Requisition:	Client #: 19104437 HO130000		
DOB: Not Given Gender: M Fasting: N Phone: NG Patient ID: NG	Collected: 05/12/2014 / 11:15 EDT Received: 05/13/2014 / 00:17 EDT Reported: 05/13/2014 / 14:49 EDT	FAIRFAX CRYO BANK		

Test Name HEMOGLOBINOPATHY EVALUATION	In Range	Out Of Range	Reference Range	Lab
RED BLOOD CELL COUNT	4.89		4.20-5.80 Million/uL	QHO
HEMOGLOBIN	14.5		13.2-17.1 g/dL	
HEMATOCRIT	42.3		38.5-50.0 %	
MCV	86.6		80.0-100.0 fL	
MCH	29.7		27.0-33.0 pg	
RDW	13.5		11.0-15.0 %	
HEMOGLOBIN A	96.5		>96.0 %	QHO
HEMOGLOBIN F	<1.0		<2.0 %	102-41
HEMOGLOBIN A2 (QUANT)	2.5		1.8-3.5 %	
INTERPRETATION				

Normal phenotype.

#### PERFORMING SITE:

Quest

Diagnostics

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QHO QUEST DIAGNOSTICS HORSHAM, 900 BUSINESS CENTER DRIVE, HORSHAM, PA 19044-3432 Laboratory Director: ANDREW S EDELMAN, MD PHD, CLIA: 39D0204404

