

Donor 4409

Genetic Testing Summary

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

Last Updated: 08/20/18

Donor Reported Ancestry: Chinese

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/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 108 mutations in the CFTR gene	1/200
Spinal Muscular Atrophy (SMA) carrier screening	Negative for deletions of exon 7 in the SMN1 gene	<1/500
Hb Beta Chain-Related Hemoglobinopathy (including Beta Thalassemia and Sickle Cell Disease) by genotyping	Negative for 37 mutations tested in the HBB gene	1/160 for Beta-Thalassemia <1/500 for Sickle Cell

*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.

)(Counsyl				
Results Recipient	Ordering Healthcare Professional	Male Details	Female Details	
Fairfax Cryobank -	Fairfax Cryobank -	Name: Donor 4409 DOB: Ethnicity: East Asian Sample Type: Saliva (OG-300) Date of Collection: 03/24/2011 Barcode: Indication: Egg or Sperm Donor	Not tested	

Universal Genetic Test (Egg or Sperm Donor)

The Universal Genetic Test uses targeted DNA mutation analysis to simultaneously determine the carrier status of an individual for a number of Mendelian diseases. This report indicates which mutations, if any, were detected for each mutation panel. Because only select mutations are tested, the percentage of carriers detected varies by ethnicity. A negative test result does not eliminate the possibility that the individual is a carrier. Interpretation is given as an estimate of the risk of conceiving a child affected with a disease, which is based on reported ethnicity, the test results, and an assumption of no family history.



Donor 4409

Donor 4409's DNA test shows that he is not a carrier of any disease-causing mutation tested.



Partner

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



Child Risk Summary

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

Note on hemoglobinopathies:

Individuals of African, Southeast Asian, and Mediterranean ancestry are at increased risk for being carriers for hemoglobinopathies and may also benefit from carrier testing by CBC and hemoglobin electrophoresis or HPLC.



* 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, and technical errors. The child risk summary is provided as an aid to genetic counseling. Inaccurate reporting of ethnicity may cause errors in risk calculation.

This test was developed and its performance cha improvement Amendments of 1988 (CLIA) as qu be regarded as investigational or for research. Th	Laboratory Diractor: Jassica Jacobson, MD CLIA Number: 05D1102604	
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Male Name: Donor 4409 DOB:

Female Not tested

Full Results

Below are the full test results for all diseases on the panel. Noted are the specific genetic mutations for which the patient tested positive or negative. If there was insufficient data to determine the genotype for any variant, this will be noted as "no call." Also listed in this section is the patient's post-test risk of being a carrier of each disease as well as the odds that his future children could inherit each disease.

Beta Thalassemia	Your child's risk: 1 in 20,000	Risk before testing: 1 in 3,900	Reduced risk
Donor 4409: No mutations detected. This does not rule out the assuming a negative family history, is 1 in 160. 80% detection		itations. The post-test risk of bein	ig a carrier,
Gene: HBB. Variants (35): K17X, C39X, 619 bp deletion, Phe41fs, Ser9fs, Lys8fs, Phe71fs, IVS-II-849(A>C), IVS-II-849(A>G), Gly24 T>A, -87C>G, Hb -30T>A, CAP+1 A>C, Hb E, Hb C-Arab.			
Cystic Fibrosis	Your child's risk: 1 in 69,000	Risk before testing: 1 in 30,000	Reduced ris
Donor 4409: No mutations detected. No call for 3199del6. The being a carrier, assuming a negative family history, is 1 in 20		carrier of untested mutations. The	e post-test risk of
1717-1G>A, 1898+1G>À, 2789+5G>A, 3120+1G>A, 3849+10kbC>T, E60X, 3876delA, 3905insT, 1812-1G>A, 3272-26A>G, 2183AA>G, S549R(A>C), C A559T, G622D, R709X, K710X, Q890X, R1066C, R1070Q, W1089X, Y1092 574delA, 663delT, 935delA, 936delTA, 1161delC, 1609delCA, 1677delTA, 1	2X, G551D, R553X, R560T, R1162X, W1282X, N1303K, , R75X, E92X, Y122X, G178R, R347H, Q493X, V520F, S 391R, R117C, 1148T, L206W, G330X, T338I, R352Q, S3 2X, R1158X, S1196X, W1204X(c.3611G>A), Q1238X, S1 949del84, 2043delG, 2055del9>A, 2105-2117del13insA4	549N, P574H, M1101K, D1152H, S1235 54P, G480C, I506V. F508C, C524X, S54 251N, S1255X, R1283M, dele2-3 21kb, 3 GAAA, 3171delC, 3667del4, 3821delT, 12	R, 394delTT, 1078delT, 9I, S549R(T>G), Q552> 3199del6, F311del, 288insTA, 2184insA,
Gene: CFTR. Variants (108): G85E, R117H, R334W, R347P, A455E, G542 1717-1G-A, 1898+1G>A, 2789+5G>A, 3120+1G>A, 3849+10kbC>T, E60X 3876delA, 3905insT, 1812-1G>A, 3272-26A>G, 2183AA>G, S549R(A>C), C A559T, G622D, R709X, K710X, Q890X, R1066C, R1070Q, W1089X, V1092 574delA, 663delT, 935delA, 936delTA, 1161delC, 1609delCA, 1677delTA, 1 2307insA, 2869insG, 296+12T>C, 405+1G>A, 405+3A>C, 406-1G>A, 711+ Sickle Cell Disease	2X, G551D, R553X, R560T, R1162X, W1282X, N1303K, , R75X, E92X, Y122X, G178R, R347H, Q493X, V520F, S 391R, R117C, 1148T, L206W, G330X, T338I, R352Q, S3 2X, R1158X, S1196X, W1204X(c.3611G>A), Q1238X, S1 949del84, 2043delG, 2055del9>A, 2105-2117del13insA4	549N, P574H, M1101K, D1152H, S1235 54P, G480C, I506V. F508C, C524X, S54 251N, S1255X, R1283M, dele2-3 21kb, 3 GAAA, 3171delC, 3667del4, 3821delT, 12	R, 394delTT, 1078delT, 91, S549R(T>G), O5522 8199del6, F311del, 88insTA, 2184insA, 3612G>A).
1717-1G-A, 1898+1G-Å, 2769+5G-A, 3120+1G-A, 3849+10kbC-7, E60X 3876delA, 3905insT, 1812-1G-A, 3272-26A>G, 2183AA>G, S549R(A>C), CA A5597, G622D, R709X, K710X, 0890X, R1066C, R1070Q, W1089X, V1092 574delA, 663delT, 935delA, 936delTA, 1161delC, 1609delCA, 1677delTA, 1 2307insA, 2869insG, 296+12T>C, 405+1G>A, 405+3A>C, 406-1G>A, 711+ Sickle Cell Disease Donor 4409: No mutations detected. This does not rule out t	2X, G551D, R553X, R560T, R1162X, W1282X, N1303K, R75X, E92X, Y122X, G178R, R347H, Q493X, V520F, S 931R, R117C, I148T, L206W, G330X, T338I, R352Q, S3 2X, R1155X, S1196X, W1204X(c, 3611G-A), Q1238X, S1 1949del64, 2043delG, 2055del9-A, 2105-2117del13insAf 5G>A, 712-1G>T, 1811+1.6kbA>G, 1898+1G>T, 1898+5 Your child's risk: Less than 1 in 1,000,000 he possibility of being a carrier of untested mL	549N, P574H, M1101K, D1152H, S1235 34P, G480C, IS06V. F508C, C524X, S543 251N, S125X, R1233M, dele2-3 21kb, 3 3AAA, 3171delC, 3667del4, 3821delT, 12 G>T, 3120G>A, 457TAT>G, W1204X(c.3 Risk before testing: less than 1 in 1,000,000	R, 394delTT, 1078delT, JI, S549R(T>G), 05529 3199del6, F311del, 886insTA, 2184insA, 3612G>A). Reduced risi
1717-1G>A, 1898+1G>À, 2789+5G>A, 3120+1G>A, 3849+10kbC>T, E60X, 3876delA, 3905insT, 1812-1G>A, 3272-26A>G, 2183AA>G, S549R(A>C), C A559T, G622D, R709X, K710X, Q890X, R1066C, R1070Q, W1089X, Y1092 574delA, 663delT, 935delA, 936delTA, 1161delC, 1609delCA, 1677delTA, 1	2X, G551D, R553X, R560T, R1162X, W1282X, N1303K, R75X, E92X, Y122X, G178R, R347H, Q493X, V520F, S 591R, R117C, I148T, L206W, G330X, T338I, R352Q, S3 2X, R1158X, S1196X, W1204X(c, 3611G-A), Q1238X, S1 1949del84, 2043delG, 2055del9-A, 2105-2117del13insAf 5G>A, 712-1G>T, 1811+1.6kbA>G, 1898+1G>T, 1898+5 Your child's risk: Less than 1 in 1,000,000 he possibility of being a carrier of untested mu ction rate. er9fs, IVS-II-654, IVS-II-745, IVS-II-850, IVS-I-6, IVS-I-11	549N, P574H, M1101K, D1152H, S1235 54P, G480C, I506V, F508C, C524X, S543 251N, S1255X, R1283M, dele2-3 21kb, 3 3AAA, 3171delC, 3667del4, 3821delT, 12 (G>T, 3120G>A, 457TAT>G, W1204X(c.3 Risk before testing: less than 1 in 1,000,000 tations. The post-test risk of bein 0, IVS-I-5, IVS-I-1(G>A), IVS-I-1(G>T), -5	R, 394delTT, 1078delT, 9I, S549R(T>G), O552X 3199del6, F311del, 886insTA, 2184insA, 3612G>A). Reduced risi g a carrier, 38C>T, -28A>G, -29A>C

Donor 4409: No mutations detected. This does not rule out the possibility of being a carrier of untested mutations. The post-test risk of being a carrier, assuming a negative family history, is < 1 in 500. 93% detection rate. Gene: SMN1. Variants (1): Exon 7 deletion.



This test was developed and its performance characteristics determined by Counsyl, Inc. The laboratory is regulated under the Clinical Laboratory Improvement Amendments of 1988 (CLIA) as qualified to perform high-complexity clinical testing. This test is used for clinical purposes. It should not be regended as investigational or for research. These results are adjunctive to the ordering physician's workup.

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Cytogenetic Report

Client H	Fair	fax Cryobank						
Address								
Reporting Phone #						Ema	ail N/A	
Patient name/Donor A	lias	Donor # 4409				Patient DOB	N/A	
Done	or #					Specimen type	Peripheral	Blood
Collection D	ate	03/24/2011				Accession #		
Date Receiv	ved	03/25/2011						
				RESU	LTS			
СҮТ	00	GENETIC AN	ALYS	SIS			FISH	
Cells counted		20		Type of banding	GTG		Probe(s)	N/A
Cells analyzed		5		Band resolution	500	Nu	clei scored	N/A
Cells karyotyped		2						
Modal chromosome #	4	46						

KARYOTYPE 46,XY

INTERPRETATION

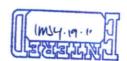
Normal male karyotype

No 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

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Wayne S. Stanley, Ph.D., FACMG Clinical Cytogeneticist



4/5/1 Date

Genetics and IVF Preimplantation Genetics Laboratory

Patient name: DONOR 4409

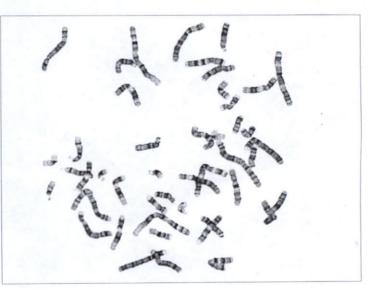
Case name:

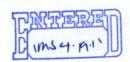
46,XY



Case:

Slide: B1 Cell: 6





Quest Diagnostics QUEST DIAGNOSTICS INCORPORATED CLIENT SERVICE 866.697.8378 SPECIMEN INFORMATION SPECIMEN : REQUISITION :	PATIENT INFORMATION ID,4409 DOB: AGE: GENDER: M FASTING: U ID: PHONE:	REPORT STATUS FINAL ORDERING PHYSICIAN STERN, HARVEY J CLIENT INFORMATION FAIRFAX CRYO BANK
COLLECTED: 03/30/2011 07:30 ET RECEIVED: 03/30/2011 23:37 ET REPORTED: 04/04/2011 15:10 ET		
Test Name	In Range Out of Range	Reference Range Lab
HEMOGLOBINOPATHY EVALUATION RED BLOOD CELL COUNT HEMOGLOBIN HEMATOCRIT MCV MCH	5.14 15.9 47.1 91.5 30.9	4.20-5.80 Million/uL QHO 13.2-17.1 g/dL 38.5-50.0 % 80.0-100.0 fL 27.0-33.0 pg
RDW HEMOGLOBIN A HEMOGLOBIN F HEMOGLOBIN A2 (QUANT) INTERPRETATION	12.6 97.6 <1.0 2.4	11.0-15.0 % >96.0 % QHO <2.0 % 1.8-3.5 %
	Normal phenotype.	
CBC (INCLUDES DIFF/PLT) WHITE BLOOD CELL COUNT RED BLOOD CELL COUNT HEMOGLOBIN HEMATOCRIT	6.2 5.14 15.9 47.1	QHO 3.8-10.8 Thousand/uL 4.20-5.80 Million/uL 13.2-17.1 g/dL 38.5-50.0 %
MCV MCH MCHC RDW PLATELET COUNT	91.5 30.9 33.8 12.6 221 3410	80.0–100.0 fL 27.0–33.0 pg 32.0–36.0 g/dL 11.0–15.0 % 140–400 Thousand/uL 1500–7800 cells/uL
ABSOLUTE NEUTROPHILS ABSOLUTE LYMPHOCYTES ABSOLUTE MONOCYTES ABSOLUTE EOSINOPHILS ABSOLUTE BASOPHILS NEUTROPHILS LYMPHOCYTES	2480 248 62 0 55 40	850-3900 cells/uL 200-950 cells/uL 15-500 cells/uL 0-200 cells/uL %
MONOCYTES EOSINOPHILS BASOPHILS	4 1 0	Z CIMS422.11

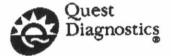
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TK-FAN 84/84/11 15:25 #12661368 1/3



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PATIENT INFORMATION ID,4409

AGE :

REPORT STATUS FINAL

ORDERING PHYSICIAN STERN, HARVEY J

QUEST DIAGNOSTICS INCORPORATED

COLLECTED: 03/30/2011 07:30 ET REPORTED: 04/04/2011 15:10 ET

GENDER: M FASTING: U

PERFORMING LABORATORY INFORMATION

QHO QUEST DIAGNOSTICS-HORSHAM, 900 BUSINESS CENTER DRIVE, HORSHAM, PA 19044-3408 Laboratory Director: HERMAN HURWITZ, MD, FCAP, CLIA: 39D0204404

DOB:

LIST OF RESULTS PRINTED IN THE OUT OF RANGE COLUMN:

ID,4409 -

Page 2 - End of Report

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