



Donor 2503

Genetic Testing Summary

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

Last Updated: 03/13/24

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 carrier screening (CFTR) | Negative by genotyping of 87 mutations in the CFTR gene | |
| Alpha 1 Antitrypsin Deficiency carrier screening (SERPINA1) | Negative by genotyping for 2 mutations in the SERPINA1 gene | |

*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 Name: Donor, 2503

Referring Physician: [REDACTED]

Specimen #: [REDACTED]

Patient ID: [REDACTED]

Client #: [REDACTED]

Case #: [REDACTED]

DOB: Not Given

Sex: M

SSN: [REDACTED]

Date Collected: 03/19/2002

Date Received: 04/30/2003

Lab ID: [REDACTED]

Hospital ID: [REDACTED]

Specimen Type: DNA

Fairfax Cryobank

Ethnicity: Asian

Indication: Carrier test / Gamete donor

RESULTS: Negative for the mutations analyzed

INTERPRETATION

The sample provided is negative for the mutations analyzed.

COMMENTS:

This analysis was performed on extracted DNA provided by Genetics and IVF Institute and identified as above.

| Mutation Detection Rates among Ethnic Groups | | Detection rates are based on mutation frequencies in patients affected with cystic fibrosis. Among individuals with an atypical or mild presentation (e.g. congenital absence of the vas deferens, pancreatitis) detection rates may vary from those provided here. | |
|--|---|---|--|
| Ethnicity | Carrier risk reduction when no family history | CF87 Detection rate | References |
| Caucasian | 1/25 to 1/325 | 92.6% | Genet in Med 3:168, 2001 in conjunction with Genet in Med 4:90, 2002 |
| African American | 1/65 to 1/338 | 81% | Genet in Med 3:168, 2001 |
| Hispanic | 1/46 to 1/162 | 72% | Genet in Med 3:168, 2001 |
| Ashkenazi Jewish | 1/26 to 1/834 | 97% | Am J Hum Genet 51:951, 1994 |
| Jewish, non-Ashkenazi | | Varies by country of origin | Genet Testing 5:47, 2001, Genet Testing, 1:35, 1997 |
| Asian | | Not Provided | Insufficient data |
| Other or Mixed Ethnicity | | Not Provided | Detection rate not determined and varies with ethnicity |

This interpretation is based on the clinical information provided and the current understanding of the molecular genetics of this condition. Although DNA-based testing is highly accurate, rare diagnostic errors may occur. Examples include misinterpretation because of genetic variants, blood transfusion, bone marrow transplantation, or erroneous representation of family relationships or contamination of a fetal sample with maternal cells.

METHOD

DNA is isolated from the sample and tested for the 87 CF mutations listed. Regions of the CFTR gene are amplified enzymatically and hybridized to specific CF mutation oligonucleotide probes. Results are characterized as positive or negative, and specimens with positive results are tested for specific mutation identity. The assay discriminates between $\Delta F508$ and the following polymorphisms: F508C, I506V, I506M and I507V.

This test was developed and its performance characteristics determined by Genzyme Genetics. It has not been cleared or approved by the U.S. Food and Drug Administration. The FDA has determined that such clearance or approval is not necessary. This test is used for clinical purposes. It should not be regarded as investigational or for research. The laboratory is regulated under the Clinical Laboratory Improvement Amendments of 1988 (CLIA) as qualified to perform high complexity clinical testing.

Under the direction of:



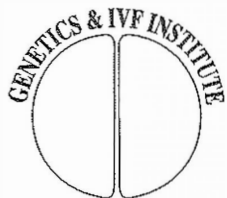
Lynne Rosenblum-Vos
Lynne Rosenblum-Vos, Ph. D.

Date: 05/08/2003

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Cystic fibrosis mutations analyzed

| | | |
|--------------------|---------------------|------------------------|
| ΔF508 | R1162X | 1898+1G>A |
| ΔI507 | R117C | 1898+5G>T |
| ΔF311 | R117H | 1949del84 |
| A455E | R1283M | 2043delG |
| A559T | R334W | 2143delT |
| C524X | R347H | 2183delAA>G |
| D1152H | R347P | 2184delA |
| D1270N | R352Q | 2307insA |
| E60X | R553X | 2789+5G>A |
| G178R | R560T | 2869insG |
| G330X | S1196X | 3120+1G>A |
| G480C | S1251N | 3120G>A |
| G542X | S1255X | 3659delC |
| G551D | S364P | 3662delA |
| G85E | S549I | 3791delC |
| G91R | S549N | 3821delT |
| I148T | S549R | 3849+10kbC>T |
| K710X | T338I | 3849+4A>G |
| L206W | V520F | 3876delA |
| M1101K | W1089X | 3905insT |
| N1303K | W1282X | 394delTT |
| P574H | Y1092X | 405+1G>A |
| Q1238X | Y563D | 405+3A>C |
| Q359K/T360K | 1078delT | 444delA |
| Q493X | 1161delC | 574delA |
| Q552X | 1609delCA | 621+1G>T |
| Q890X | 1677delTA | 711+1G>T |
| R1066C | 1717-1G>A | 711+5G>A |
| R1158X | 1812-1G>A | 712-1G>T |



GENETICS & IVF INSTITUTE
3022 Javier Road, Fairfax, Virginia 22031 (800) 654-GENE

Joseph D. Schulman, M.D., *Medical Director*

Name: Donor 2503
ID No.: [REDACTED]
Specimen: Peripheral blood
Referred By: Megan Taylor

Family No.: [REDACTED]
Sample No.: [REDACTED]
Date Drawn: 3/19/2002
Received: 3/19/2002

Test: α_1 -Antitrypsin S and Z mutations.

PI*S Result: Negative.

PI*Z Result: Negative.

Conclusion: This individual is not a carrier of the S or Z α_1 -antitrypsin mutations.

Comment: Deficiency in the protease inhibitor α_1 -antitrypsin can cause chronic obstructive pulmonary disease (emphysema). Deficiencies in this enzyme occur through a variety of different mutations in the α_1 -antitrypsin gene. Two, called *PI*Z* and *PI*S*, are particularly common. Individuals who inherit two *PI*Z* alleles have a high risk of developing emphysema. They also may experience transient hepatitis or permanent liver damage in childhood or later in life. Individuals who inherit one *PI*Z* and one *PI*S* allele also have a somewhat increased risk for emphysema and liver disease. Persons who have one α_1 -antitrypsin allele that is intact and one that has the *PI*Z* mutation may have some increased risk of emphysema, especially with smoking. Since about 1 person in 20 in the U.S. is a carrier of a *PI*S* or *PI*Z* allele, healthy adults may want screening to determine if they and their partner are at risk of having a child with two deficient alleles. If results are positive, genetic counseling is indicated.

Note: This test examines the α_1 -antitrypsin gene at the specific positions associated with the common S and Z mutations. Mutations other than S and Z would not be detected. This method differs from PI Typing, in which the protein itself is examined and classified as S, Z, M (normal), or another variant.

March 27, 2002
Date

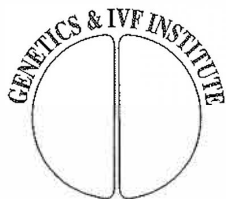
A. Maddalena
Anne Maddalena, PhD, ABMG

Laboratory Director

W. Christine Spence, PhD, ABMG

Associate Director

This test was developed and its performance characteristics determined by Genetics & IVF Institute. It has not been cleared or approved by the U.S. FDA. The FDA has determined that such clearance or approval is not necessary. Pursuant to the requirements of CLIA '88 this laboratory has established the test's accuracy and precision.



GENETICS & IVF INSTITUTE
3022 Javier Road, Fairfax, Virginia 22031 (800) 654-GENE

Joseph D. Schulman, M.D., Medical Director

Patient: DONOR #2503

PB Lab. No.: [REDACTED]

Hospital/Chart No: Fairfax

D.O.B./Age: [REDACTED]

Physician Name: [REDACTED]

Preliminary: --

Source No.: [REDACTED]

Final: 04-30-02

Date Received: 04-16-02

Collected: 04-15-02

Specimen: Blood

Test: Chromosome Analysis

Band Resolution:

400

550

850

Number of Cells Counted: 20

Modal Chromosome Count: 46

Number of Cells Karyotyped: 2

Karyotype: 46,XY

Impression: Normal Male Karyotype

Comment:

Clinical Cytogeneticists

Wayne S. Stanley, Ph.D.

Lillian D. Killos, Ph.D.

Julie Leana-Cox, Ph.D.

Denise A. Batista, Ph.D.

Most chromosome variants of no clinical significance, if present, are not reported. This analysis does not rule out the possibility of subtle structural chromosome abnormalities, low frequency chromosome mosaicism, or defects of non-chromosomal etiology.

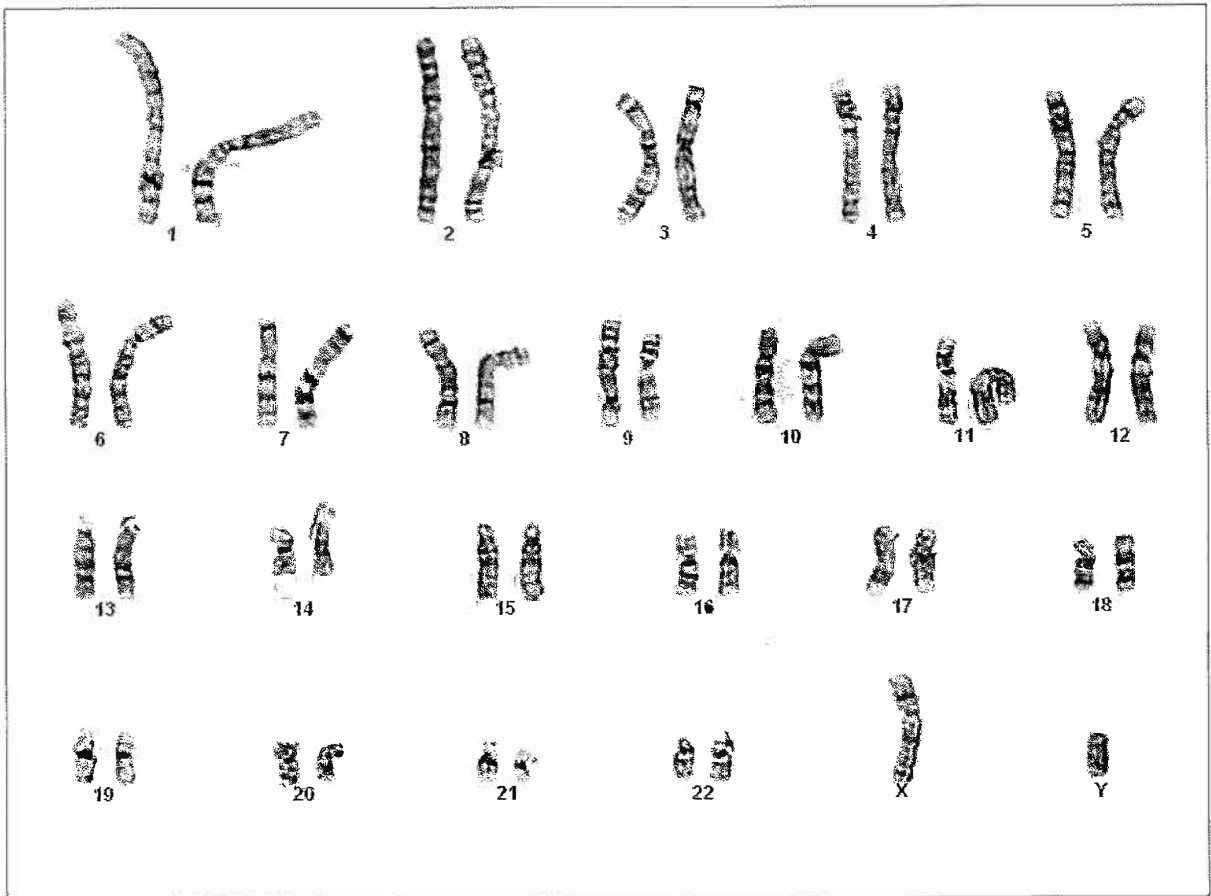
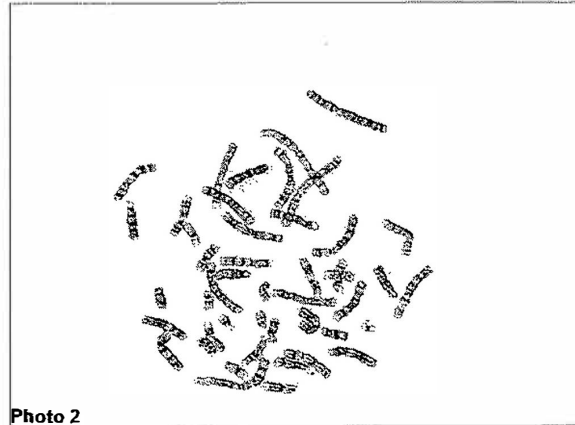
**Genetics & IVF Institute
Cytogenetics Laboratory**

Case name: [REDACTED]

Patient name: Donor #2503

Result:

Case comment:



Technologist: YSG



SPECIALTY LABORATORIES

2211 Michigan Avenue

310-828-6543

Santa Monica, CA 90404-3900

800-421-4449

SPECIALTY #

CLIENT #

PATIENT NAME: DONOR #2503,

PHYSICIAN:

NOTES:

PATIENT ID:

2503-

SPECIMEN ID:

FAIRFAX CRYOBANK

ATTN: STEVE POOL, PHD

3015 WILLIAMS DR STE 110

DOB: N/A

AGE: unknown

SEX: Unknown

FAIRFAX

VA 22031

DRAWN:

03/19/02 05:00

RECEIVED:

03/21/02 09:21

PRINTED:

03/26/02 13:23

FINAL REPORT:

03/26/02 13:23

FINAL

| TEST NAME | RESULTS | REFERENCE RANGE |
|--------------------|-----------|-----------------|
| ALT (SGPT) | | |
| ALT (SGPT) | 52 H U, L | (< 41) |
| AST (SGOT) | | |
| AST (SGOT) | 26 U, L | (< 41) |
| CHOLESTEROL, TOTAL | | |
| Cholesterol, Total | 183 mg/dL | (< 200) |

REFERENCE RANGES for Cholesterol, Total:

| Age | mg/dL | Risk Level |
|--------------|----------------------|-----------------|
| Less than 2 | No range established | |
| 2 - 18 years | < 170 | Desirable |
| | 170 - 199 | Borderline High |
| | > 200 | High |
| Adult | < 200 | Desirable |
| | 200 - 239 | Borderline High |
| | > 240 | High |

HEMOGLOBIN VARIANT SCREEN REFLEX TO ELECTROPHORESIS

| | | | |
|---------------|------|-----|-------------|
| Hemoglobin A | 96.8 | % | (93.5-98.3) |
| Hemoglobin A2 | 3.2 | H % | (2.1-3.0) |
| Hemoglobin F | 0.0 | % | (< 2.1) |