

Sample Report

Cystic Fibrosis Mutation Analysis

Any Hospital

USA

100 Main Street

Anytown, ST 12345

Patient Name:

Referring Physician: Ordering Doctor, MD

 Specimen #: 12345678
 Client #: 123456

 Patient ID: 12345678-9
 Case #: 1234567

DOB: /1978 Date Collected: 05/28/2014 Sex: F Date Received: 05/29/2014

SSN: LAB ID: Hospital ID:

Specimen Type: BLDPER

Ethnicity: Caucasian

Indication: Carrier Test / No family history

RESULTS: Negative for the 97 mutations analyzed

INTERPRETATION:

This individual is negative for the mutations analyzed. This result reduces but does not eliminate the risk to be a CF carrier. See Comments for ethnic-specific risk reductions based on a negative family history.

COMMENTS:

Mutations 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. congential absence of the vas deferens, pancreatitis) detection rates may vary from those provided here.				
Ethnicity	Carrier risk reduction when no family history	Detection rate	References		
African American	1/61 to 1/316	81%	ACOG Committee Opinion 486 PMID: 21422883; Heim PMID: 11388756		
Ashkenazi Jewish	1/24 to 1/767	97%	ACOG Committee Opinion 486 PMID: 21422883		
Asian American	1/94 to <1/183	49-55%	ACOG Committee Opinion 486 PMID: 21422883; Watson PMID: 1384328		
Caucasian	1/25 to 1/343	93%	ACOG Committee Opinion 486 PMID: 21422883; Heim PMID: 11388756; Palomaki PMID: 11882786		
Hispanic	1/58 to 1/260	78%	ACOG Committee Opinion 486 PMID: 21422883; Heim PMID: 11388756; California Database: (http://www.cdph.ca.gov/programs/GDSP/Documents/CFTabelCurrent.pdf)		
Jewish, non-Ashkenazi		Varies by country of origin	Orgad PMID: 11336401; Kerem PMID:10464623		
Mixed or Other		Not Provided	For counseling, consider using the ethnic background with the most conservative risk estimates.		

This interpretation is based on the clinical and family relationship information provided and the current understanding of the molecular genetics of this condition.

METHOD / LIMITATIONS:

CFTR gene regions are amplified enzymatically. The 97 CF mutations are tested by multiplex allele-specific primer extension, bead array hybridization, and fluorescence detection. The test discriminates between p.F508del and three polymorphisms (p.I506V, p.I507V and p.F508C). Numbering and nomenclature follow Human Genome Variation Society recommendations. Mutations and their legacy names are listed at www.integratedgenetics.com/CFplus. The DNA reference sequence is NG_016465.1. False positive or negative results may occur for reasons that include genetic variants, blood transfusions, bone marrow transplantation, erroneous representation of family relationships, or maternal contamination of a fetal sample.

Integrated Genetics is a business unit of Esoterix Genetic Laboratories, LLC, a wholly-owned subsidiary of Laboratory Corporation of America Holdings.

Electronically Signed By:	PhD_FACMG on 06/02/2014

MUTATIONS ANALYZED					
c.54-5940_273+10250del21kb	c.1973_1985del13insAGAAA	p.R117H	p.R553*		
c.262_263delTT	c.1976delA	p.Y122*	p.A559T		
c.273+1G>A	c.2012delT	p.G178R	p.R560T		
c.273+3A>C	c.2051_2052delAAinsG	p.L206W	p.P574H		
c.274-1G>A	c.2052delA	p.F312del	p.R709*		
c.313delA	c.2052dupA	p.G330*	p.K710*		
c.325_327delTATinsG	c.2175dupA	p.R334W	p.R764*		
c.442delA	c.2657+5G>A	p.T338I	p.Q890*		
c.489+1G>T	c.2737_2738insG	p.R347H	p.R1066C		
c.531delT	c.2988G>A	p.R347P	p.W1089*		
c.579+1G>T	c.2988+1G>A	p.R352Q	p.Y1092*		
c.579+5G>A	c.3039delC	p.[Q359K;T360K]	p.M1101K		
c.580-1G>T	c.3067_3072delATAGTG	p.S364P	p.D1152H		
c.803delA	c.3528delC	p.A455E	p.R1158*		
c.805_806delAT	c.3536_3539delCCAA	p.G480C	p.R1162*		
c.948delT	c.3659delC	p.Q493*	p.S1196*		
c.1155_1156dupTA	c.3717+12191C>T	p.I507del	p.W1204*		
c.1545_1546delTA	c.3744delA	p.F508del	p.Q1238*		
c.1585-1G>A	c.3773dupT	p.V520F	p.S1251N		
c.1680-1G>A	c.3889dupT	p.C524*	p.S1255*		
c.1766+1G>A	p.E60*	p.G542*	p.W1282*		
c.1766+5G>T	p.R75*	p.S549N	p.N1303K		
c.1820_1903del84 p.G85E c.1911delG p.E92* c.1923_1931del9insA p.R117C		p.S549R			
		p.G551D			
		p.Q552*			

This test was developed and its performance characteristics determined by Esoterix Genetic Laboratories, LLC. It has not been cleared or approved by the Food and Drug Administration. The FDA has determined that such clearance or approval is not necessary.