

Full name:	Cataract - UGent
Type of panel:	<u>in-house</u>
Version number:	v3
Laboratory:	<u>Centrum Medische Genetica - UZ Gent</u>
Created:	17 Jun 2019 - 11:49
Changed:	16 Dec 2022 - 10:38

Related Diseases

- 3-methylglutaconic aciduria type 3
- Autosomal dominant cutis laxa
- Autosomal dominant optic atrophy and cataract
- Autosomal dominant spastic paraplegia type 9A
- Autosomal dominant spastic paraplegia type 9B
- Autosomal dominant vitreoretinchoroidopathy
- Autosomal recessive bestrophinopathy
- Autosomal recessive spastic paraplegia type 9B
- Axenfeld-Rieger syndrome
- BOR syndrome
- Branchiootic syndrome
- Cataract-microcornea syndrome
- Cerebrotendinous xanthomatosis
- Cerulean cataract
- Classic galactosemia
- Cockayne syndrome type 1
- Cockayne syndrome type 2
- Cockayne syndrome type 3
- Coloboma of choroid and retina
- Congenital cataracts-facial dysmorphism-neuropathy syndrome
- Coralliform cataract

- Craniometaphyseal dysplasia
- Dent disease type 2
- Early-onset anterior polar cataract
- Early-onset lamellar cataract
- Early-onset non-syndromic cataract
- Early-onset nuclear cataract
- Early-onset posterior polar cataract
- Early-onset posterior subcapsular cataract
- Early-onset sutural cataract
- Erythrokeratoderma variabilis
- Familial exudative vitreoretinopathy
- Familial isolated dilated cardiomyopathy
- Familial multiple meningioma
- Familial thoracic aortic aneurysm and aortic dissection
- Full schwannomatosis
- Galactokinase deficiency
- Hereditary hyperferritinemia-cataract syndrome
- Hypochondrogenesis
- Hypomyelination-congenital cataract syndrome
- Isolated aniridia
- Isolated congenital sclerocornea
- Isolated optic nerve hypoplasia/aplasia
- Kniest dysplasia
- Knobloch syndrome
- Legg-Calvé-Perthes disease
- Microphthalmia, Lenz type
- Morning glory disc anomaly
- Nance-Horan syndrome
- Oculocerebrorenal syndrome of Lowe
- Oculodentodigital dysplasia
- Otofaciocervical syndrome
- Platyspondylic dysplasia, Torrance type
- Pulverulent cataract
- Rare isolated myopia
- Retinitis pigmentosa
- Retinopathy of prematurity

- Stickler syndrome type 1
- Syndactyly type 3
- Total early-onset cataract
- UV-sensitive syndrome

Related Analytes

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
<u>ABHD12</u>	99.98	1	
<u>ADAMTSL4</u>	99.66	1	
<u>AGK</u>	99.99	1	
<u>ALDH18A1</u>	99.96	1	
<u>B3GLCT</u>	99.90	1	
<u>BCOR</u>	99.97	1	
<u>BEST1</u>	99.86	1	
<u>BFSP1</u>	100.00	1	
<u>BFSP2</u>	99.09	1	
<u>CHMP4B</u>	99.96	1	
<u>COL11A1</u>	90.72	1	
<u>COL18A1</u>	99.99	1	
<u>COL2A1</u>	99.87	1	
<u>CRYAA</u>	19.49	1	
<u>CRYAB</u>	100.00	1	
<u>CRYBA1</u>	99.99	1	

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
<u>CRYBA2</u>	100.00	1	
<u>CRYBA4</u>	100.00	1	
<u>CRYBB1</u>	99.46	1	
<u>CRYBB2</u>	99.94	1	
<u>CRYBB3</u>	99.99	1	
<u>CRYGB</u>	99.99	1	
<u>CRYGC</u>	100.00	1	
<u>CRYGD</u>	100.00	1	
<u>CRYGS</u>	100.00	1	
<u>CTDP1</u>	99.97	1	
<u>CYP27A1</u>	100.00	1	
<u>CYP51A1</u>	97.82	1	
<u>DNMBP</u>	99.94	1	
<u>EPG5</u>	99.95	1	
<u>EPHA2</u>	99.99	1	
<u>EYA1</u>	99.81	1	
<u>FBN1</u>	99.85	1	
<u>FOXE3</u>	99.29	1	
<u>FTL</u>	99.99	1	
<u>FYCO1</u>	100.00	1	

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
<u>FZD4</u>	100.00	1	
<u>GALK1</u>	100.00	1	
<u>GALT</u>	100.00	1	
<u>GCNT2</u>	100.00	1	
<u>GJA1</u>	100.00	1	
<u>GJA3</u>	100.00	1	
<u>GJA8</u>	99.99	1	
<u>HMX1</u>	100.00	1	
<u>HSF4</u>	99.98	1	
<u>HYCC1</u>	99.81	1	
<u>INPP5K</u>	99.94	1	
<u>INTS1</u>	100.00	1	
<u>JAM3</u>	100.00	1	
<u>LCAT</u>	99.97	1	
<u>LEMD2</u>	100.00	1	
<u>LIM2</u>	100.00	1	
<u>LSS</u>	99.98	1	
<u>MAF</u>	99.73	1	
<u>MIPEP</u>	99.98	1	
<u>MIR184</u>	100.00	1	

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
<u>MYH9</u>	99.95	1	
<u>NDP</u>	99.98	1	
<u>NF2</u>	100.00	1	
<u>NHS</u>	99.96	1	
<u>OCRL</u>	99.89	1	
<u>OPA3</u>	100.00	1	
<u>P3H2</u>	99.93	1	
<u>PANK4</u>	100.00	1	
<u>PAX6</u>	99.95	1	
<u>PITX3</u>	100.00	1	
<u>PXDN</u>	100.00	1	
<u>RRAGA</u>	100.00	1	
<u>SIL1</u>	99.95	1	
<u>SIPA1L3</u>	99.97	1	
<u>SLC16A12</u>	100.00	1	
<u>SLC33A1</u>	99.67	1	
<u>TDRD7</u>	99.87	1	
<u>UNC45B</u>	100.00	1	
<u>VIM</u>	100.00	1	
<u>VSX2</u>	99.99	1	

GENE	% OF CODING SEQUENCE SUFFICIENTLY COVERED TO DETECT HETEROZYGOUS MUTATIONS	COPY NUMBER VARIATION	COMMENTS
<u>WFS1</u>	99.99	1	