

Estimated residual risks after tested negative using Blueprint Genetics Carrier Test

Blueprint Genetics



This table below displays residual risks after a negative result for each of the genes and corresponding disorders and is not calculated separately for each tested individual. If a patient is reported to be a carrier of a disease, their residual risk equals “1” and this table does not apply for that disease.

Residual risk values were calculated based on published mutation or carrier frequencies for dominant and recessive disorders (e.g., Guo MH and Gregg AR, 2019) and estimated detection rates of Blueprint Genetics’ testing technologies. Detection rate is probability to detect disease-causing variants when accounting for sequence coverage, mapping quality and known mutation profile.

Key: **AD** = autosomal dominant, **AR** = autosomal recessive; **XL** = X-linked; **XLD** = X-linked dominant, **XLR** = X-linked recessive, **Worldwide** = global, non-specific populations

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (FMR1 / DUO) 461 genes	Core Reproductive Screen (FMR1 / DUO) 126 genes	Ashkenazi Jewish Reproductive Screen (FMR1 / DUO) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
AAAS	Achalasia-addisonianism-alacrimia syndrome	AR	Worldwide	1 in 416	99	1 in 41580	•		•	•	
AARS2	Progressive leukoencephalopathy with ovarian failure	AR	Worldwide	1 in 514	99	1 in 51390	•	•		•	•
ABCA3	Pulmonary surfactant metabolism dysfunction type 3	AR	Worldwide	1 in 270	99	1 in 27000	•	•		•	•
ABCB11	Progressive familial intrahepatic cholestasis type 2	AR	Worldwide	1 in 200	99	1 in 20000	•			•	
ABCC6 [#]	Pseudoxanthoma elasticum	AR	Worldwide	1 in 90	98	1 in 4500	•	•		•	•
ABCC8	Hyperinsulinemic hypoglycemia	AD/AR	Worldwide	1 in 180	99	1 in 18000	•	•	•	•	•
ABCD1 [#]	Adrenoleukodystrophy	XLR	Worldwide	1 in 8500	98	1 in 425000	•	•			
ACAD9	Acyl-CoA dehydrogenase family, deficiency	AR	Worldwide	1 in 600	99	1 in 60000	•			•	
ACADM	Acyl-CoA dehydrogenase, medium chain, deficiency	AR	Worldwide	1 in 110	99	1 in 11000	•	•		•	•
ACADS	Short-Chain Acyl-CoA Dehydrogenase Deficiency	AR	Worldwide	1 in 127	99	1 in 12690	•			•	
ACADSB	2-Methylbutyrylglycinuria	AR	Worldwide	1 in 199	99	1 in 19890	•			•	
ACADVL	Very long chain Acyl-CoA dehydrogenase deficiency	AR	Worldwide	1 in 200	99	1 in 20000	•	•		•	•
ACOX1	Peroxisomal acyl-CoA oxidase deficiency	AR	Worldwide	1 in 3254	99	1 in 325400	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>ACSF3</i>	Combined malonic and methylmalonic aciduria	AR	Worldwide	1 in 132	99	1 in 13200	•			•	
<i>ADA</i>	Severe combined immunodeficiency due to adenosine deaminase deficiency	AR	Worldwide	1 in 420	99	1 in 42000	•			•	
<i>ADA2</i>	ADA2 deficiency	AR	Worldwide	1 in 335	99	1 in 33480	•			•	
<i>ADAMTS2</i> Excluded from the analysis: NM_021599, exon 11	Ehlers-Danlos syndrome	AR	Worldwide	1 in 2200	98	1 in 110000	•		•	•	
<i>ADAMTSL4</i>	Isolated ectopia lentis	AR	Worldwide	1 in 50	99	1 in 4950	•			•	
<i>ADGRG1</i>	Bilateral polymicrogyria	AR	Worldwide	1 in 980	99	1 in 98000	•			•	
<i>ADGRV1</i>	Usher syndrome type IIC	AR	Worldwide	1 in 192	99	1 in 19170	•	•		•	•
<i>AGA</i>	Aspartylglucosaminuria	AR	Worldwide	1 in 970	99	1 in 97000	•	•		•	•
<i>AGL</i>	Glycogen storage disease	AR	Worldwide	1 in 620	99	1 in 62000	•		•	•	
<i>AGPS</i>	Rhizomelic chondrodysplasia punctata type 3	AR	Worldwide	1 in 158	99	1 in 15800	•			•	
<i>AGXT</i>	Primary hyperoxaluria type 1	AR	Worldwide	1 in 316	99	1 in 31600	•	•		•	•
<i>AHI1</i>	Joubert syndrome	AR	Worldwide	1 in 2300	99	1 in 230000	•	•		•	•
<i>AIFM1</i>	Severe X-linked mitochondrial encephalomyopathy, Cowchock syndrome and Deafness	XLR	Worldwide	1 in 450	99	1 in 45000	•				
<i>AIRE</i>	Autoimmune polyendocrinopathy syndrome	AR	Worldwide	1 in 210	99	1 in 21000	•	•		•	•
<i>ALDH3A2</i>	Sjogren-Larsson syndrome	AR	Worldwide	1 in 1200	99	1 in 120000	•			•	
<i>ALDH7A1</i>	Pyridoxine Dependent Epilepsy	AR	Worldwide	1 in 294	99	1 in 29430	•			•	
<i>ALDOB</i>	Fructose intolerance, hereditary	AR	Worldwide	1 in 160	99	1 in 16000	•	•		•	•
<i>ALG6</i>	Congenital disorder of glycosylation	AR	Worldwide	1 in 920	99	1 in 92000	•			•	
<i>ALMS1</i> #	Alström syndrome	AR	Worldwide	1 in 1920	98	1 in 96000	•	•		•	•
<i>ALPL</i>	Hypophosphatasia perinatal lethal, infantile, juvenile and adult forms	AD/AR	Worldwide	1 in 588	99	1 in 58800	•	•		•	•
<i>AMT</i>	Glycine encephalopathy	AR	Worldwide	1 in 1240	99	1 in 124000	•			•	
<i>ANO10</i>	Spinocerebellar ataxia	AR	Worldwide	1 in 520	99	1 in 52000	•	•		•	•

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>AP1S1</i>	MEDNIK syndrome	AR	Worldwide	1 in 1423	99	1 in 142290	•			•	
<i>AP3B1</i>	Hermansky-Pudlak syndrome	AR	Worldwide	1 in 212	95	1 in 4248	•			•	
<i>AR</i>	Androgen insensitivity and Hypospadias 1, X-linked	XLR	Worldwide	1 in 18000	97	1 in 600000	•				
<i>ARG1</i>	Hyperargininemia	AR	Worldwide	1 in 296	99	1 in 29600	•			•	
<i>ARL6</i>	Bardet-Biedl syndrome and Retinitis pigmentosa	AR	Worldwide	1 in 712	99	1 in 71190	•			•	
<i>ARSA</i>	Metachromatic leukodystrophy	AR	Worldwide	1 in 288	99	1 in 28800	•	•	•	•	•
<i>ARSB</i>	Mucopolysaccharidosis (Maroteaux-Lamy)	AR	Worldwide	1 in 613	99	1 in 61300	•			•	
<i>ARX</i>	Epileptic encephalopathy, Hydranencephaly with abnormal genitalia and Lissencephaly	XL	Worldwide	1 in 450000	90	1 in 4500000	•	•			
<i>ASL</i>	Argininosuccinic aciduria	AR	Worldwide	1 in 223	99	1 in 22300	•	•		•	•
<i>ASNS</i> [#]	Asparagine synthetase deficiency	AR	Worldwide	1 in 500	97	1 in 16667	•			•	
<i>ASPA</i>	Aspartoacylase deficiency (Canavan disease)	AR	Worldwide	1 in 427	99	1 in 42700	•	•	•	•	•
<i>ASS1</i>	Citrullinemia	AR	Worldwide	1 in 1343	99	1 in 134300	•	•		•	•
<i>ATM</i>	Ataxia-Telangiectasia	AR	Worldwide	1 in 423	99	1 in 42300	•		•	•	
<i>ATP13A2</i>	Kufor-Rakeb syndrome and Spastic paraplegia	AR	Worldwide	1 in 900	99	1 in 90000	•			•	
<i>ATP6V1B1</i>	Renal tubular acidosis with deafness	AR	Worldwide	1 in 866	99	1 in 86600	•			•	
<i>ATP7A</i>	Menkes disease	XLR	Worldwide	1 in 175000	99	1 in 17500000	•				
<i>ATP7B</i>	Wilson disease	AR	Worldwide	1 in 97	99	1 in 9700	•	•	•	•	•
<i>ATRX</i>	Alpha-thalassemia intellectual disability syndrome	XLR	Worldwide	1 in 100000	99	1 in 10000000	•				
<i>AVPR2</i>	Nephrogenic diabetes insipidus	XLR	Worldwide	1 in 144000	97	1 in 4800000	•				
<i>B9D1</i>	Meckel syndrome	AR	Worldwide	1 in 5400	99	1 in 540000	•			•	
<i>B9D2</i>	Meckel syndrome	AR	Worldwide	1 in 5400	99	1 in 540000	•			•	
<i>BBS1</i>	Bardet-Biedl syndrome	AR	Worldwide	1 in 520	99	1 in 52000	•	•		•	•

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>BBS10</i>	Bardet-Biedl syndrome	AR	Worldwide	1 in 562	99	1 in 56200	•	•		•	•
<i>BBS12</i>	Bardet-Biedl syndrome	AR	Worldwide	1 in 1230	99	1 in 123000	•	•		•	•
<i>BBS2</i>	Bardet-Biedl syndrome and Retinitis pigmentosa	AR	Worldwide	1 in 870	99	1 in 87000	•	•	•	•	•
<i>BBS4</i>	Bardet-Biedl syndrome	AR	Worldwide	1 in 768	99	1 in 76800	•			•	
<i>BBS5</i>	Bardet-Biedl syndrome	AR	Worldwide	1 in 919	99	1 in 91900	•			•	
<i>BBS7</i>	Bardet-Biedl syndrome	AR	Worldwide	1 in 873	99	1 in 87300	•			•	
<i>BBS9</i>	Bardet-Biedl syndrome	AR	Worldwide	1 in 959	99	1 in 95900	•			•	
<i>BCKDHA</i>	Maple syrup urine disease	AR	Worldwide	1 in 1002	99	1 in 100200	•		•	•	
<i>BCKDHB</i>	Maple syrup urine disease	AR	Worldwide	1 in 475	99	1 in 47500	•	•	•	•	•
<i>BCS1L</i>	GRACILE syndrome and other BCS1L related disorders	AR	Worldwide	1 in 720	99	1 in 72000	•			•	
<i>BLM</i>	Bloom syndrome	AR	Worldwide	1 in 632	99	1 in 63200	•	•	•	•	•
<i>BRAT1</i>	Lethal neonatal rigidity and multifocal seizure syndrome	AR	Worldwide	1 in 540	99	1 in 54000	•			•	
<i>BSND</i>	Bartter syndrome	AR	Worldwide	1 in 866	99	1 in 86600	•			•	
<i>BTB</i>	Biotinidase deficiency	AR	Worldwide	1 in 642	99	1 in 64200	•	•		•	•
<i>CANT1</i>	Desbuquois dysplasia and Epiphyseal dysplasia, multiple	AR	Worldwide	1 in 1170	99	1 in 117000	•			•	
<i>CAPN3</i>	Limb-girdle muscular dystrophy	AR	Worldwide	1 in 112	99	1 in 11200	•			•	
<i>CASQ2</i>	Catecholaminergic polymorphic ventricular tachycardia (CPVT)	AR	Worldwide	1 in 1661	99	1 in 166140	•	•		•	•
<i>CBS</i>	Homocystinuria due to cystathionine beta-synthase deficiency	AR	Worldwide	1 in	99	1 in 41100	•	•		•	•
<i>CC2D2A</i> Excluded from the analysis: NM_020785, exon 7	Joubert syndrome, Meckel syndrome and other CC2D2A related disorders	AR	Worldwide	1 in 389	99	1 in 38900	•	•		•	•
<i>CCDC88C</i>	Congenital hydrocephalus type 1	AR	Worldwide	1 in 2236	99	1 in 223600	•	•		•	•
<i>CD40</i>	Hyper-IgM syndrome	AR	Worldwide	1 in 21600	99	1 in 2160000	•			•	
<i>CD40LG</i>	Hyper-IgM syndrome	XLR	Worldwide	1 in 36000	97	1 in 1200000	•				
<i>CDH23</i>	Usher syndrome type 1D	AR	Worldwide	1 in 970	99	1 in 97000	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>CENPJ</i>	Primary microcephaly and Seckel syndrome	AR	Worldwide	1 in 540	99	1 in 54000	•			•	
<i>CEP290</i> [#]	Leber congenital amaurosis 10 and other CEP290 related ciliopathies	AR	Worldwide	1 in 84	99	1 in 8400	•	•		•	•
<i>CERKL</i>	Retinitis pigmentosa	AR	Worldwide	1 in 194	99	1 in 19400	•			•	
<i>CFTR</i>	Cystic fibrosis	AR	Worldwide	1 in 42	99	1 in 4200	•	•	•	•	•
<i>CHAT</i>	Congenital myasthenic syndrome	AR	Worldwide	1 in 939	99	1 in 93870	•			•	
<i>CHM</i> Excluded from the analysis: NM_001145414, exon 5	Choroideremia	XLR	Worldwide	1 in 37500	96	1 in 937500	•				
<i>CHRNE</i>	Myasthenic syndrome	AR	Worldwide	1 in 302	99	1 in 30200	•	•		•	•
<i>CIITA</i>	Bare lymphocyte syndrome	AR	Worldwide	1 in 1224	99	1 in 122400	•			•	
<i>CLCN1</i>	Myotonia congenita	AR	Worldwide	1 in 632	99	1 in 63200	•	•		•	•
<i>CLN3</i>	Neuronal ceroid lipofuscinosis type 3	AR	Worldwide	1 in 760	99	1 in 76000	•	•		•	•
<i>CLN5</i>	Neuronal ceroid lipofuscinosis type 5	AR	Worldwide	1 in 1021	99	1 in 102100	•			•	
<i>CLN6</i>	Neuronal ceroid lipofuscinosis type 6	AR	Worldwide	1 in 4355	99	1 in 435500	•			•	
<i>CLN8</i>	Neuronal ceroid lipofuscinosis type 8	AR	Worldwide	1 in 1896	99	1 in 189600	•			•	
<i>CLRN1</i>	Usher syndrome type 3A and Retinitis pigmentosa	AR	Worldwide	1 in 970	99	1 in 97000	•	•	•	•	•
<i>CNGA3</i>	Achromatopsia	AR	Worldwide	1 in 157	99	1 in 15660	•			•	
<i>CNGB3</i>	Achromatopsia and Juvenile macular degeneration	AR	Worldwide	1 in 978	99	1 in 97800	•	•		•	•
<i>COL27A1</i>	Steel syndrome	AR	Worldwide	1 in 500	99	1 in 50000	•			•	
<i>COL4A3</i>	Alport syndrome	AR	Worldwide	1 in 500	99	1 in 50000	•		•	•	
<i>COL4A4</i>	Alport syndrome	AR	Worldwide	1 in 707	99	1 in 70700	•			•	
<i>COL4A5</i>	Alport syndrome	XLD/XLR	Worldwide	1 in 30000	98	1 in 1500000	•			•	
<i>COL7A1</i>	Epidermolysis bullosa dystrophica	AR	Worldwide	1 in 152	99	1 in 15200	•	•		•	•
<i>COLQ</i>	Congenital myasthenic syndrome	AR	Worldwide	1 in 939	99	1 in 93870	•			•	
<i>CPS1</i>	Carbamoylphosphate synthetase I deficiency	AR	Worldwide	1 in 4877	99	1 in 487700	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>CPT1A</i>	Carnitine palmitoyltransferase deficiency	AR	Worldwide	1 in 3115	99	1 in 311500	•			•	
<i>CPT2</i>	Carnitine palmitoyltransferase II deficiency	AR	Worldwide	1 in 1157	99	1 in 115700	•	•	•	•	•
<i>CRB1</i>	Leber congenital amaurosis, Pigmented paravenous chorioretinal atrophy and Retinitis pigmentosa	AD/AR	Worldwide	1 in 112	99	1 in 11200	•			•	
<i>CRPPA</i>	Muscular dystrophy-dystroglycanopathy		Worldwide	1 in 2842	99	1 in 284220	•			•	
<i>CTNS</i>	Cystinosis	AR	Worldwide	1 in 1179	99	1 in 117900	•		•	•	
<i>CTSD</i>	Neuronal ceroid lipofuscinosis	AR	Worldwide	1 in 7200	99	1 in 720000	•			•	
<i>CTSF</i>	Neuronal ceroid lipofuscinosis	AR	Worldwide	1 in 1080	93	1 in 15429	•			•	
<i>CTSK</i>	Pycnodysostosis	AR	Worldwide	1 in 3382	99	1 in 338200	•			•	
<i>CYBA</i>	Chronic granulomatous disease	AR	Worldwide	1 in 866	99	1 in 86600	•			•	
<i>CYBB</i>	Chronic granulomatous disease	XLR	Worldwide	1 in 150000	99	1 in 15000000	•				
<i>CYP11B1</i> #	Congenital adrenal hyperplasia due to 11-Beta-Hydroxylase Deficiency	AR	Worldwide	1 in 224	92	1 in 2800	•			•	
<i>CYP11B2</i>	Corticosterone methyloxidase type I deficiency and Corticosterone methyloxidase type II deficiency	AR	Worldwide	1 in 1420	99	1 in 142000	•			•	
<i>CYP17A1</i>	Congenital adrenal hyperplasia due to 17-Alpha-hydroxylase deficiency	AR	Worldwide	1 in 112	99	1 in 11200	•			•	
<i>CYP19A1</i>	Aromatase deficiency	AR	Worldwide	1 in 707	99	1 in 70700	•			•	
<i>CYP1B1</i>	Primary congenital glaucoma	AR	Worldwide	1 in 235	97	1 in 7830	•			•	
<i>CYP21A2</i> #	Adrenal hyperplasia, congenital, due to 21-hydroxylase deficiency Hyperandrogenism, nonclassic , due to 21-hydroxylase deficiency	AR	Worldwide	1 in 61	75	1 in 244	•	•		•	•
<i>CYP27A1</i>	Cerebrotendinous xanthomatosis	AR	Worldwide	1 in 402	99	1 in 40200	•	•		•	•
<i>CYP27B1</i>	Vitamin D-dependent rickets	AR	Worldwide	1 in 578	99	1 in 57800	•	•		•	•
<i>CYP7B1</i>	Spastic paraplegia type 5A	AR	Worldwide	1 in 470	99	1 in 46980	•			•	
<i>DBT</i>	Maple syrup urine disease	AR	Worldwide	1 in 513	99	1 in 51300	•			•	
<i>DCLRE1C</i> #	Severe combined immunodeficiency with sensitivity to ionizing radiation	AR	Worldwide	1 in 1682	98	1 in 84100	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>DDX11</i>	Warsaw breakage syndrome	AR	Worldwide	1 in 1581	99	1 in 158100	•			•	
<i>DGAT1</i>	Diarrhea	AR	Worldwide	1 in 1350	95	1 in 27000	•			•	
<i>DHCR7</i>	Smith-Lemli-Opitz syndrome	AR	Worldwide	1 in 87	99	1 in 8700	•	•	•	•	•
<i>DHDDS</i>	Retinitis pigmentosa	AR	Worldwide	1 in 2344	99	1 in 234400	•	•	•	•	•
<i>DKC1</i>	Dyskeratosis congenita and Hoyeraal-Hreidarsson syndrome	XLR	Worldwide	1 in 2000000	99	1 in 200000000	•				
<i>DLD</i>	Dihydrolipoyl dehydrogenase deficiency	AR	Worldwide	1 in 1130	99	1 in 113000	•	•	•	•	•
<i>DMD</i>	Duchenne & Becker muscular dystrophy and Dilated cardiomyopathy	XLR/XLD	Worldwide	1 in 2000	99	1 in 200000	•	•			
<i>DNAH5</i>	Ciliary dyskinesia	AR	Worldwide	1 in 178	99	1 in 17800	•		•	•	
<i>DNAI1</i>	Ciliary dyskinesia	AR	Worldwide	1 in 720	99	1 in 72000	•		•	•	
<i>DNAI2</i>	Ciliary dyskinesia	AR	Worldwide	1 in 689	99	1 in 68900	•		•	•	
<i>DOK7</i>	Congenital myasthenic syndrome	AR	Worldwide	1 in 373	95	1 in 7452	•			•	
<i>DYNC2H1</i>	Short -rib thoracic dysplasia with or without polydactyly type 1	AR	Worldwide	1 in 273	99	1 in 27300	•	•		•	•
<i>DYSF</i>	Limb-girdle muscular dystrophy	AR	Worldwide	1 in 470	99	1 in 47000	•			•	
<i>EDA</i>	Hypohidrotic ectodermal dysplasia	XLR	Worldwide	1 in 5700	99	1 in 570000	•				
<i>EIF2AK4</i>	Pulmonary venoocclusive disease	AR	Worldwide	1 in 635	99	1 in 63540	•			•	
<i>EIF2B2</i>	Leukoencephalopathy with vanishing white matter	AR	Worldwide	1 in 783	99	1 in 78300	•			•	
<i>EIF2B5</i>	Leukoencephalopathy with vanishing white matter and Ovarioleukodystrophy	AR	Worldwide	1 in 1472	99	1 in 147200	•			•	
<i>ELP1</i>	Familial dysautonomia	AR	Worldwide	1 in 373	99	1 in 37260	•	•	•	•	•
<i>EMD</i>	Emery-Dreifuss muscular dystrophy	XLR	Worldwide	1 in 300000	99	1 in 30000000	•				
<i>EPG5</i>	Vici syndrome	AR	Worldwide	1 in 864	99	1 in 86400	•			•	
<i>ERCC2</i>	Xeroderma pigmentosum and Photo-sensitive trichothiodystrophy	AR	Worldwide	1 in 167	99	1 in 16700	•	•		•	•
<i>ERCC6#</i>	Cockayne syndrome type B and other ERCC6 related disorders	AR	Worldwide	1 in 723	98	1 in 36150	•			•	
<i>ERCC6L2</i>	Bone marrow failure syndrome 2	AR	Worldwide	1 in 1080	99	1 in 108000	•			•	
<i>ERCC8</i>	Cockayne syndrome	AR	Worldwide	1 in 4730	99	1 in 473000	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>ESCO2</i>	Roberts syndrome	AR	Worldwide	1 in 978	99	1 in 97800	•			•	
<i>ETFA</i>	Glutaric aciduria type II	AR	Worldwide	1 in 1997	99	1 in 199700	•			•	
<i>ETFB</i>	Glutaric aciduria	AR	Worldwide	1 in 3600	99	1 in 360000	•			•	
<i>ETFDH</i>	Glutaric aciduria	AR	Worldwide	1 in 1422	99	1 in 142200	•			•	
<i>ETHE1</i>	Ethylmalonic encephalopathy	AR	Worldwide	1 in 3214	99	1 in 321400	•			•	
<i>EVC</i>	Ellis-van Creveld syndrome	AR	Worldwide	1 in 1630	96	1 in 40750	•			•	
<i>EVC2</i>	Ellis-van Creveld syndrome	AR	Worldwide	1 in 1230	99	1 in 123000	•	•		•	•
<i>EXOSC3</i>	Pontocerebellar hypoplasia	AR	Worldwide	1 in 514	99	1 in 51390	•			•	
<i>EYS</i> #	Retinitis pigmentosa	AR	Worldwide	1 in 189	99	1 in 18900	•			•	
<i>F11</i>	Factor XI deficiency	AD/AR	Worldwide	1 in 219	99	1 in 21900	•		•	•	
<i>F9</i>	Factor IX deficiency	XLD/XLR	Worldwide	1 in 2000	99	1 in 200000	•	•			
<i>FAH</i>	Tyrosinemia	AR	Worldwide	1 in 371	99	1 in 37100	•	•	•	•	•
<i>FAM161A</i>	Retinitis pigmentosa	AR	Worldwide	1 in 534	99	1 in 53400	•		•	•	
<i>FANCA</i>	Fanconi anemia	AR	Worldwide	1 in 1756	99	1 in 175600	•	•		•	•
<i>FANCC</i>	Fanconi anemia	AR	Worldwide	1 in 623	99	1 in 62300	•	•	•	•	•
<i>FANCE</i>	Fanconi anemia	AR	Worldwide	1 in 1350	99	1 in 135000	•			•	
<i>FANCG</i>	Fanconi anemia	AR	Worldwide	1 in 1736	99	1 in 173600	•			•	
<i>FANCI</i>	Fanconi anemia	AR	Worldwide	1 in 569	99	1 in 56880	•			•	
<i>FANCL</i>	Fanconi anemia	AR	Worldwide	1 in 1800	99	1 in 180000	•			•	
<i>FH</i>	Fumarase deficiency	AR	Worldwide	1 in 970	99	1 in 97000	•			•	
<i>FHL1</i> #	Reducing bod myopathy and Hypertrophic and dilated cardiomyopathy	XL	Worldwide	1 in 1000000	99	1 in 100000000	•			•	
<i>FKRP</i>	Muscular dystrophy-dystroglycanopathy	AR	Worldwide	1 in 350	99	1 in 35000	•	•		•	•
<i>FKTN</i>	Muscular dystrophy-dystroglycanopathy	AR	Worldwide	1 in 724	99	1 in 72400	•	•	•	•	•
<i>FMO3</i>	Trimethylaminuria	AR	Worldwide	1 in 321	99	1 in 32100	•	•		•	•
<i>FMR1</i>	Fragile X syndrome	XLD	Worldwide	1 in 4000 full mutation	99	1 in 400000	•	•	•		
<i>FMR1</i>	Fragile X syndrome	XLD	Worldwide	1 in 201 premutation	99	1 in 20100	•	•	•		
<i>FRAS1</i>	Fraser syndrome	AR	Worldwide	1 in 540	99	1 in 54000	•			•	
<i>G6PC</i>	Glycogen storage disease	AR	Worldwide	1 in 354	99	1 in 35400	•	•	•	•	•
<i>GAA</i>	Glycogen storage disease	AR	Worldwide	1 in 134	99	1 in 13400	•	•	•	•	•
<i>GALC</i>	Krabbe disease	AR	Worldwide	1 in 784	99	1 in 78400	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>GALE</i>	Galactose epimerase deficiency	AR	Worldwide	1 in 3375	99	1 in 337500	•			•	
<i>GALK1</i>	Galactokinase deficiency	AR	Worldwide	1 in 780	99	1 in 78000	•			•	
<i>GALNS</i>	Mucopolysaccharidosis (Morquio syndrome)	AR	Worldwide	1 in 432	97	1 in 14400	•			•	
<i>GALNT3</i>	Familial hyperphosphatemic tumoral calcinosis	AR	Worldwide	1 in 794	99	1 in 79380	•			•	
<i>GALT</i>	Galactosemia	AR	Worldwide	1 in 183	99	1 in 18300	•	•	•	•	•
<i>GAMT</i>	Guanidinoacetate methyltransferase deficiency	AR	Worldwide	1 in 917	99	1 in 91700	•			•	
<i>GBA#</i>	Gaucher disease	AR	Worldwide	1 in 360	90	1 in 3600	•	•	•	•	•
<i>GBE1</i>	Glycogen storage disease	AR	Worldwide	1 in 316	99	1 in 31600	•	•		•	•
<i>GCDH</i>	Glutaric aciduria	AR	Worldwide	1 in 217	99	1 in 21700	•			•	
<i>GCH1</i>	BH4-deficient hyperphenylalaninemia B	AR	Worldwide	1 in 15428	99	1 in 1542780	•			•	
<i>GCSH</i>	Glycine encephalopathy	AR	Worldwide	1 in 3600	90	1 in 36000	•			•	
<i>GFM1</i>	Combined oxidative phosphorylation deficiency	AR	Worldwide	1 in 1223	99	1 in 122300	•			•	
<i>GFPT1</i>	Congenital myasthenic syndrome	AR	Worldwide	1 in 3086	99	1 in 308610	•			•	
<i>GJB1</i>	Charcot-Marie-Tooth neuropathy	XLD	Worldwide	1 in 5000	99	1 in 500000	•		•	•	
<i>GJB2</i>	Deafness	AD/AR	Worldwide	1 in 37	99	1 in 3700	•	•	•	•	•
<i>GJB6</i>	Deafness	AD/AR	Worldwide	1 in 238	99	1 in 23800	•	•		•	•
<i>GLA</i>	Fabry disease	XLR/XLD	Worldwide	1 in 40000	99	1 in 4000000	•	•		•	•
<i>GLB1</i>	GM1-gangliosidosis Mucopolysaccharidosis (Morquio syndrome)	AR	Worldwide	1 in 512	99	1 in 51200	•	•		•	•
<i>GLDC</i>	Glycine encephalopathy	AR	Worldwide	1 in 722	99	1 in 72200	•			•	
<i>GLE1</i>	Lethal congenital contracture syndrome	AR	Worldwide	1 in 2341	99	1 in 234100	•			•	
<i>GNE</i>	Nonaka myopathy	AR	Worldwide	1 in 236	99	1 in 23600	•			•	
<i>GNPAT</i>	Rhizomelic chondrodysplasia punctata	AR	Worldwide	1 in 3600	99	1 in 360000	•			•	
<i>GNPTAB</i>	Mucopolipidosis	AR	Worldwide	1 in 213	99	1 in 21300	•	•		•	•
<i>GNPTG</i>	Mucopolipidosis	AR	Worldwide	1 in 1258	99	1 in 125800	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>GNS</i>	Mucopolysaccharidosis (Sanfilippo syndrome)	AR	Worldwide	1 in 5368	99	1 in 536800	•			•	
<i>GORAB</i>	Geroderma osteodysplasticum	AR	Worldwide	1 in 2348	99	1 in 234810	•			•	
<i>GP9</i>	Bernard-Soulier syndrome	AR	Worldwide	1 in 2144	99	1 in 214400	•			•	
<i>GRHPR</i>	Primary hyperoxaluria type 2	AR	Worldwide	1 in 866	99	1 in 86600	•			•	
<i>GRIP1</i>	Fraser syndrome	AR	Worldwide	1 in 489	99	1 in 48900	•			•	
<i>HADH</i>	3-hydroxyacyl-CoA dehydrogenase deficiency	AR	Worldwide	1 in 6353	99	1 in 635310	•			•	
<i>HADHA</i>	Long chain 3-hydroxyacyl-CoA dehydrogenase deficiency	AR	Worldwide	1 in 416	99	1 in 41600	•			•	
<i>HADHB</i>	Trifunctional protein deficiency	AR	Worldwide	1 in 1113	99	1 in 111330	•			•	
<i>HAMP</i>	Juvenile hemochromatosis	AR	Worldwide	1 in 6750	99	1 in 675000	•			•	
<i>HAX1</i>	Severe congenital neutropenia	AR	Worldwide	1 in 3785	99	1 in 378500	•			•	
<i>HBA1</i> #	Alpha-thalassemia (Hemoglobin Bart syndrome / Hemoglobin H disease)	AR/Di-genic	Worldwide	1 in 50	90	1 in 500	•	•	•	•	•
<i>HBA2</i> #	Alpha-thalassemia (Hemoglobin Bart syndrome / Hemoglobin H disease)	AR/Di-genic	Worldwide	1 in 50	90	1 in 500	•	•	•	•	•
<i>HBB</i>	Beta-thalassemia and Other Thalassemias/Hemoglobinopathies	AD/AR/Digenic	Worldwide	1 in 116	99	1 in 11600	•	•	•	•	•
<i>HEXA</i>	Tay-Sachs disease and Hexosaminidase A deficiency	AR	Worldwide	1 in 423	99	1 in 42300	•	•	•	•	•
<i>HEXB</i>	Sandhoff disease	AR	Worldwide	1 in 587	99	1 in 58700	•			•	
<i>HGD</i>	Alkaptonuria	AR	Worldwide	1 in 500	99	1 in 50000	•			•	
<i>HGSNAT</i>	Mucopolysaccharidosis (Sanfilippo syndrome) and Retinitis pigmentosa	AR	Worldwide	1 in 1023	97	1 in 34100	•			•	
<i>HJV</i>	Juvenile hemochromatosis	AR	Worldwide	1 in 1080	99	1 in 108000	•			•	
<i>HLCS</i>	Holocarboxylase synthetase deficiency	AR	Worldwide	1 in 912	99	1 in 91200	•			•	
<i>HMGCL</i>	3-hydroxy-3-methylglutaryl-CoA lyase deficiency	AR	Worldwide	1 in 2240	99	1 in 224000	•			•	
<i>HOGA1</i>	Primary hyperoxaluria type 3	AR	Worldwide	1 in 866	99	1 in 86600	•		•	•	
<i>HPRT1</i>	Kelley-Seegmiller syndrome and Lesch-Nyhan syndrome	XLR	Worldwide	1 in 760000	97	1 in 25333333	•				

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>HPS1</i> [#]	Hermansky-Pudlak syndrome	AR	Worldwide	1 in 500	98	1 in 25000	•	•		•	•
<i>HPS3</i>	Hermansky-Pudlak syndrome	AR	Worldwide	1 in 1581	99	1 in 158100	•	•	•	•	•
<i>HPS4</i>	Hermansky-Pudlak syndrome	AR	Worldwide	1 in 1367	99	1 in 136710	•			•	
<i>HPS5</i>	Hermansky-Pudlak syndrome	AR	Worldwide	1 in 1350	99	1 in 135000	•			•	
<i>HPS6</i>	Hermansky-Pudlak syndrome	AR	Worldwide	1 in 1636	99	1 in 163620	•			•	
<i>HSD17B3</i>	Neurodevelopmental disorder with hypotonia, neonatal respiratory insufficiency, and thermodysregulation	AR	Worldwide	1 in 450	99	1 in 45000	•			•	
<i>HSD17B4</i>	D-bifunctional protein deficiency and Perrault syndrome	AR	Worldwide	1 in 899	99	1 in 89900	•			•	
<i>HSD3B2</i>	3-beta-hydroxysteroid dehydrogenase II deficiency	AR	Worldwide	1 in 1922	99	1 in 192200	•			•	
<i>HYLS1</i>	Hydroletharus syndrome	AR	Worldwide	1 in 1197	99	1 in 119700	•			•	
<i>IDS</i> [#]	Mucopolysaccharidosis type 2	XLR	Worldwide	1 in 206	90	1 in 2060	•				
<i>IDUA</i>	Mucopolysaccharidosis	AR	Worldwide	1 in 158	99	1 in 15800	•	•		•	•
<i>IL2RG</i>	Combined immunodeficiency	XLR	Worldwide	1 in 100000	99	1 in 10000000	•				
<i>IL7R</i>	Severe combined immunodeficiency	AR	Worldwide	1 in 1440	99	1 in 144000	•			•	
<i>IQCB1</i>	Senior-Loken syndrome	AR	Worldwide	1 in 628	99	1 in 62820	•			•	
<i>IVD</i>	Isovaleric acidemia	AR	Worldwide	1 in 565	99	1 in 56500	•	•		•	•
<i>JAK3</i>	Severe combined immunodeficiency	AR	Worldwide	1 in 2160	99	1 in 216000	•			•	
<i>KCNJ1</i>	Bartter syndrome, antenatal	AR	Worldwide	1 in 720	99	1 in 72000	•			•	
<i>KCTD7</i>	Epilepsy, progressive myoclonic	AR	Worldwide	1 in 3600	99	1 in 360000	•			•	
<i>KIAA0586</i> Excluded from the analysis: NM_001244189, exons 6 and 33	Joubert syndrome and Short rib thoracic dysplasia with polydactyly	AR	Worldwide	1 in 124	95	1 in 2484	•	•		•	•
<i>KIF14</i>	Meckel syndrome 12	AR	Worldwide	1 in 1800	99	1 in 180000	•			•	
<i>L1CAM</i>	Hydrocephalus due to congenital stenosis of aqueduct of Sylvius and Other L1CAM related disorders	XLR	Worldwide	1 in 16000	99	1 in 1600000	•	•			
<i>LAMA2</i>	LAMA2 related muscular dystrophy	AR	Worldwide	1 in 857	99	1 in 85700	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>LAMA3</i>	Junctional epidermolysis bullosa	AR	Worldwide	1 in 1856	99	1 in 185600	•			•	
<i>LAMB3</i>	Junctional epidermolysis bullosa	AR	Worldwide	1 in 673	99	1 in 67300	•			•	
<i>LAMC2</i>	Junctional epidermolysis bullosa	AR	Worldwide	1 in 2112	99	1 in 211200	•			•	
<i>LARGE1</i>	Muscular dystrophy-dystroglycanopathy	AR	Worldwide	1 in 3724	85	1 in 24828	•			•	
<i>LCA5</i>	Leber congenital amaurosis	AR	Worldwide	1 in 1417	99	1 in 141700	•			•	
<i>LDLR</i>	Familial hypercholesterolemia	AD/AR	Worldwide	1 in 120	99	1 in 12000	•			•	
<i>LHX3</i>	Combined pituitary hormone deficiency	AR	Worldwide	1 in 3725	99	1 in 372500	•			•	
<i>LIFR</i>	Stuve-Wiedemann dysplasia	AR	Worldwide	1 in 2153	99	1 in 215300	•			•	
<i>LIG4</i>	LIG4 syndrome (Severe combined immunodeficiency with sensitivity to ionizing radiation)	AR	Worldwide	1 in 554	99	1 in 55350	•			•	
<i>LIPA</i>	Cholesterol ester storage disease and Wolman disease	AR	Worldwide	1 in 614	99	1 in 61400	•			•	
<i>LIPH</i>	Hypotrichosis 7	AR	Worldwide	1 in 720	99	1 in 72000	•			•	
<i>LOXHD1</i>	Deafness	AR	Worldwide	1 in 513	99	1 in 51300	•		•	•	
<i>LRP2</i>	Donnai-Barrow syndrome	AR	Worldwide	1 in 462	99	1 in 46200	•	•		•	•
<i>LRPPRC</i>	Leigh syndrome French-Canadian type	AR	Worldwide	1 in 1347	99	1 in 134700	•			•	
<i>LYST</i>	Chediak-Higashi syndrome	AR	Worldwide	1 in 1440	99	1 in 144000	•			•	
<i>MAN2B1</i>	Alpha mannosidosis	AR	Worldwide	1 in 734	99	1 in 73400	•			•	
<i>MCCC1</i>	3-Methylcrotonyl-CoA carboxylase 1 deficiency	AR	Worldwide	1 in 628	99	1 in 62800	•			•	
<i>MCCC2</i>	3-Methylcrotonyl-CoA carboxylase 2 deficiency	AR	Worldwide	1 in 754	99	1 in 75400	•	•		•	•
<i>MCOLN1</i>	Mucopolipidosis	AR	Worldwide	1 in 1785	99	1 in 178500	•	•	•	•	•
<i>MCPH1</i> Excluded from the analysis: NM_001322042, exon 14	Microcephaly	AR	Worldwide	1 in 623	97	1 in 20767	•	•		•	•
<i>MED17</i>	Postnatal progressive microcephaly with seizures and brain atrophy	AR	Worldwide	1 in 1988	99	1 in 198800	•			•	
<i>MEFV</i>	Familial Mediterranean fever	AR	Worldwide	1 in 145	97	1 in 4830	•		•	•	
<i>MESP2</i>	Spondylocostal dysostosis 2	AR	Worldwide	1 in 2247	99	1 in 224700	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>MFSD8</i>	Neuronal ceroid lipofuscinosis	AR	Worldwide	1 in 1784	99	1 in 178400	•	•		•	•
<i>MID1</i> [#]	Opitz GBBB syndrome	XLR	Worldwide	1 in 56250	96	1 in 1406250	•	•			
<i>MKKS</i>	Bardet-Biedl syndrome and McKusick-Kaufman syndrome	AR	Worldwide	1 in 1125	99	1 in 112500	•			•	
<i>MKS1</i>	Bardet-Biedl syndrome and Meckel syndrome	AR	Worldwide	1 in 632	97	1 in 21067	•			•	
<i>MLC1</i>	Megalencephalic leukoencephalopathy with subcortical cysts	AR	Worldwide	1 in 1021	99	1 in 102100	•			•	
<i>MMAA</i>	Methylmalonic acidemia	AR	Worldwide	1 in 1082	99	1 in 108200	•			•	
<i>MMAB</i>	Methylmalonic acidemia	AR	Worldwide	1 in 1866	99	1 in 186600	•			•	
<i>MMACHC</i>	Methylmalonic aciduria and homocystinuria	AR	Worldwide	1 in 207	99	1 in 20700	•	•		•	
<i>MMADHC</i>	Methylmalonic aciduria and homocystinuria	AR	Worldwide	1 in 3784	99	1 in 378400	•			•	•
<i>MMUT</i>	Methylmalonic acidemia due to methylmalonyl-CoA mutase deficiency	AR	Worldwide	1 in 257	99	1 in 25740	•			•	
<i>MPI</i>	Congenital disorder of glycosylation	AR	Worldwide	1 in 1368	99	1 in 136800	•			•	
<i>MPL</i>	Congenital megakaryocytic thrombocytopenia	AR	Worldwide	1 in 812	99	1 in 81200	•		•	•	
<i>MPV17</i>	Mitochondrial DNA depletion syndrome	AR	Worldwide	1 in 1216	99	1 in 121600	•			•	
<i>MRE11</i>	Ataxia-telangiectasia-like disorder-1	AR	Worldwide	1 in 1029	99	1 in 102870	•			•	
<i>MTHFR</i>	Homocystinuria due to MTHFR deficiency	AR	Worldwide	1 in 1261	99	1 in 126100	•			•	
<i>MTM1</i>	Centronuclear myopathy	XLR	Worldwide	1 in 25000	98	1 in 1250000	•				
<i>MTTP</i>	Abetalipoproteinemia	AR	Worldwide	1 in 2540	99	1 in 254000	•		•	•	
<i>MVK</i>	Mevalonic aciduria	AR	Worldwide	1 in 374	99	1 in 37400	•	•		•	•
<i>MYO15A</i>	Deafness	AR	Worldwide	1 in 163	99	1 in 16290	•			•	
<i>MYO7A</i>	Usher syndrome type I	AR	Worldwide	1 in 384	99	1 in 38400	•			•	
<i>NAGLU</i>	Mucopolysaccharidosis (Sanfilippo syndrome)	AR	Worldwide	1 in 286	99	1 in 28600	•			•	
<i>NAGS</i>	N-acetylglutamate synthase deficiency	AR	Worldwide	1 in 1837	99	1 in 183700	•			•	
<i>NBN</i>	Nijmegen breakage syndrome	AR	Worldwide	1 in 1623	99	1 in 162300	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>NCF2</i>	Chronic granulomatous disease	AR	Worldwide	1 in 2038	99	1 in 203760	•			•	
<i>NDRG1</i>	Charcot-Marie-Tooth disease	AR	Worldwide	1 in 7234	99	1 in 723400	•			•	
<i>NDUFAF5</i>	Mitochondrial complex I deficiency	AR	Worldwide	1 in 1423	99	1 in 142300	•			•	
<i>NDUFAF6</i>	Leigh syndrome	AR	Worldwide	1 in 1661	99	1 in 166140	•			•	
<i>NDUFS4</i>	Leigh syndrome due to mitochondrial complex I deficiency	AR	Worldwide	1 in 1589	99	1 in 158900	•			•	
<i>NDUFS6</i>	Mitochondrial complex I deficiency	AR	Worldwide	1 in 5468	99	1 in 546800	•		•	•	
<i>NEB</i> #	Nemaline myopathy	AR	Worldwide	1 in 1134	86	1 in 8100	•	•	•	•	•
<i>NGLY1</i>	Congenital disorder of deglycosylation	AR	Worldwide	1 in 939	99	1 in 93870	•			•	
<i>NPC1</i>	Niemann-Pick disease	AR	Worldwide	1 in 487	98	1 in 24350	•			•	
<i>NPC2</i>	Niemann-pick disease	AR	Worldwide	1 in 3855	99	1 in 385500	•			•	
<i>NPHP1</i>	Joubert syndrome, Nephronophthisis and Senior-Loken syndrome	AR	Worldwide	1 in 235	99	1 in 23490	•			•	
<i>NPHP3</i>	Meckel syndrome, Nephronophthisis and Renal-hepatic-pancreatic dysplasia	AR	Worldwide	1 in 376	99	1 in 37620	•			•	
<i>NPHS1</i>	Nephrotic syndrome	AR	Worldwide	1 in 152	99	1 in 15200	•	•		•	•
<i>NPHS2</i>	Nephrotic syndrome	AR	Worldwide	1 in 987	99	1 in 98700	•			•	
<i>NR0B1</i>	Congenital adrenal hypoplasia	XLR	Worldwide	1 in 60000	99	1 in 6000000	•	•			
<i>NR2E3</i>	Enhanced S-cone syndrome and Retinitis pigmentosa	AD/AR	Worldwide	1 in 894	99	1 in 89400	•		•	•	
<i>NTRK1</i>	Congenital insensitivity to pain with anhidrosis	AR	Worldwide	1 in 1196	99	1 in 119600	•			•	
<i>OAT</i>	Gyrate atrophy of choroid and retina	AR	Worldwide	1 in 892	99	1 in 89200	•			•	
<i>OCA2</i>	Albinism	AR	Worldwide	1 in 164	99	1 in 16400	•	•		•	•
<i>OPA3</i>	3-methylglutaconic aciduria and Optic atrophy	AD/AR	Worldwide	1 in 6258	99	1 in 625800	•			•	
<i>ORC4</i>	Meier-Gorlin syndrome (Ear-patella-short stature syndrome)	AR	Worldwide	1 in 1543	99	1 in 154260	•			•	
<i>OTC</i>	Ornithine transcarbamylase deficiency	XLR/XLD	Worldwide	1 in 30000	98	1 in 1500000	•	•			
<i>OTOF</i>	Deafness	AR	Worldwide	1 in 410	99	1 in 41040	•			•	
<i>PAH</i>	Phenylketonuria	AR	Worldwide	1 in 62	99	1 in 6200	•	•	•	•	•

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>PC</i>	Pyruvate carboxylase deficiency	AR	Worldwide	1 in 5966	99	1 in 596600	•			•	
<i>PCCA</i>	Propionic acidemia	AR	Worldwide	1 in 1322	99	1 in 132200	•	•		•	•
<i>PCCB</i> Excluded from the analysis: NM_001178014, exon 4	Propionic acidemia	AR	Worldwide	1 in 972	99	1 in 97200	•	•		•	•
<i>PCDH15</i>	Usher syndrome type 1D	AR	Worldwide	1 in 874	99	1 in 87400	•	•	•	•	•
<i>PCNT</i>	Microcephalic osteodysplastic primordial dwarfism	AR	Worldwide	1 in 360	99	1 in 36000	•			•	
<i>PDHA1</i>	Pyruvate dehydrogenase E1-alpha deficiency	XLD	Worldwide	1 in 50000	99	1 in 5000000	•			•	
<i>PDHB</i>	Pyruvate dehydrogenase E1-beta deficiency	AR	Worldwide	1 in 3874	99	1 in 387400	•			•	
<i>PEPD</i>	Prolidase deficiency	AR	Worldwide	1 in 857	99	1 in 85680	•			•	
<i>PET100</i>	Mitochondrial complex IV deficiency	AR	Worldwide	1 in 9000	99	1 in 900000	•			•	
<i>PEX1</i>	Zellweger spectrum disorder	AR	Worldwide	1 in 673	99	1 in 67300	•	•		•	•
<i>PEX10</i>	Zellweger spectrum disorder	AR	Worldwide	1 in 4213	99	1 in 421300	•			•	
<i>PEX12</i>	Zellweger spectrum disorder	AR	Worldwide	1 in 964	99	1 in 96400	•			•	
<i>PEX2</i>	Zellweger spectrum disorder	AR	Worldwide	1 in 2145	99	1 in 214500	•		•	•	
<i>PEX26</i>	Peroxisome biogenesis disorder	AR	Worldwide	1 in 1688	93	1 in 24107	•			•	
<i>PEX6</i>	Zellweger spectrum disorder	AR	Worldwide	1 in 412	99	1 in 41200	•	•	•	•	•
<i>PEX7</i>	Refsum disease and Rhizomelic CDP type 1	AR	Worldwide	1 in 563	99	1 in 56300	•	•		•	•
<i>PFKM</i>	Glycogen storage disease	AR	Worldwide	1 in 1358	99	1 in 135800	•		•	•	
<i>PHGDH</i>	Neu-Laxova syndrome 1	AR	Worldwide	1 in 1463	99	1 in 146300	•		•	•	
<i>PHKB</i>	Glycogen storage disease	AR	Worldwide	1 in 590	99	1 in 59040	•			•	
<i>PHYH</i>	Refsum disease	AR	Worldwide	1 in 500	99	1 in 50000	•			•	
<i>PKHD1</i>	Polycystic kidney disease	AR	Worldwide	1 in 697	99	1 in 69700	•	•	•	•	•
<i>PLP1</i>	Pelizaeus-Merzbacher disease and Spastic paraplegia	XLR	Worldwide	1 in 200000	99	1 in 20000000	•	•			
<i>PMM2</i>	Congenital disorder of glycosylation	AR	Worldwide	1 in 114	99	1 in 11400	•	•	•	•	•

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>PNPO</i>	Pyridoxamine 5'-phosphate oxidase deficiency	AR	Worldwide	1 in 1200	99	1 in 119970	•			•	
<i>POLG</i>	POLG-related ataxia neuropathy spectrum disorders	AR	Worldwide	1 in 365	99	1 in 36500	•	•		•	•
<i>POMGNT1</i>	Muscular dystrophy-dystroglycanopathy	AR	Worldwide	1 in 785	99	1 in 78500	•			•	
<i>POMT1</i>	Muscular dystrophy-dystroglycanopathy	AR	Worldwide	1 in 581	99	1 in 58050	•			•	
<i>POMT2</i>	Muscular dystrophy-dystroglycanopathy	AR	Worldwide	1 in 1080	99	1 in 108000	•			•	
<i>POR</i>	POR deficiency	AR	Worldwide	1 in 450	99	1 in 45000	•			•	
<i>PPT1</i>	Neuronal ceroid lipofuscinosis	AR	Worldwide	1 in 346	99	1 in 34600	•	•		•	•
<i>PRF1</i>	Hemophagocytic lymphohistiocytosis	AR	Worldwide	1 in 1657	99	1 in 165700	•	•		•	•
<i>PROP1</i>	Combined pituitary hormone deficiency	AR	Worldwide	1 in 2868	99	1 in 286800	•			•	
<i>PRPS1</i> [#]	PRPS1 related disorders	XLR/XLD	Worldwide	1 in 50000	99	1 in 5000000	•			•	
<i>PSAP</i>	PSAP related disorders	AR	Worldwide	1 in 5687	99	1 in 568700	•			•	
<i>PTS</i>	Hyperphenylalaninemia, BH4-deficient	AR	Worldwide	1 in 917	99	1 in 91700	•			•	
<i>PUS1</i>	Mitochondrial myopathy and sideroblastic anemia	AR	Worldwide	1 in 6712	99	1 in 671200	•			•	
<i>PYGL</i>	Glycogen storage disease	AR	Worldwide	1 in 662	99	1 in 66240	•			•	
<i>PYGM</i>	Glycogen storage disease	AR	Worldwide	1 in 354	99	1 in 35400	•			•	
<i>QDPR</i>	BH4-deficient hyperphenylalaninemia C	AR	Worldwide	1 in 3600	97	1 in 120000	•			•	
<i>RAB23</i>	Carpenter syndrome 1	AR	Worldwide	1 in 1366	99	1 in 136600	•			•	
<i>RAG1</i>	Severe combined immunodeficiency	AR	Worldwide	1 in 392	99	1 in 39240	•	•		•	•
<i>RAG2</i>	Combined cellular and humoral immune defects with granulomas and Omenn syndrome	AR	Worldwide	1 in 1825	99	1 in 182500	•			•	
<i>RAPSN</i>	Congenital myasthenic syndrome	AR	Worldwide	1 in 2389	99	1 in 238900	•			•	
<i>RARS2</i>	Pontocerebellar hypoplasia	AR	Worldwide	1 in 812	99	1 in 81200	•			•	
<i>RDH12</i>	Leber congenital amaurosis and Retinitis pigmentosa	AD/AR	Worldwide	1 in 623	99	1 in 62300	•			•	
<i>RECQL4</i>	Rothmund-Thomson syndrome	AR	Worldwide	1 in 207	98	1 in 10350	•			•	
<i>RMRP</i>	Cartilage-hair hypoplasia	AR	Worldwide	1 in 812	98	1 in 40600	•	•		•	•

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>RNASEH2A</i>	Aicardi-Goutières syndrome	AR	Worldwide	1 in 3086	99	1 in 308610	•			•	
<i>RNASEH2B</i>	Aicardi-Goutières syndrome	AR	Worldwide	1 in 416	99	1 in 41600	•	•		•	•
<i>RNASEH2C</i>	Aicardi-Goutières syndrome	AR	Worldwide	1 in 4320	95	1 in 86400	•			•	
<i>RPE65</i>	Leber congenital amaurosis and Retinitis pigmentosa	AD/AR	Worldwide	1 in 589	99	1 in 58900	•			•	
<i>RPGR</i>	Retinitis pigmentosa	XL	Worldwide	1 in 11250	99	1 in 1125000	•	•		•	•
<i>RPGRIP1</i>	Leber congenital amaurosis	AR	Worldwide	1 in 675	99	1 in 67500	•			•	
<i>RPGRIP1L</i> Excluded from the analysis: NM_015272, exon 23	Joubert syndrome, Meckel syndrome and Other ciliopathy syndromes	AR	Worldwide	1 in 947	98	1 in 47350	•			•	
<i>RS1</i>	Retinoschisis	XLR	Worldwide	1 in 10000	99	1 in 1000000	•	•			
<i>RTEL1</i>	Dyskeratosis congenita and Pulmonary fibrosis	AD/AR	Worldwide	1 in 1258	99	1 in 125800	•		•	•	
<i>SACS</i>	Spastic ataxia, Charlevoix-Saguenay	AR	Worldwide	1 in 1473	99	1 in 147300	•			•	
<i>SAMHD1</i>	Aicardi-Goutières syndrome	AR	Worldwide	1 in 1469	99	1 in 146900	•			•	
<i>SCO2</i>	Fatal infantile cardioencephalomyopathy due to cytochrome c oxidase deficiency	AR	Worldwide	1 in 728	99	1 in 72800	•	•		•	•
<i>SEPSECS</i>	Pontocerebellar hypoplasia type 2D	AR	Worldwide	1 in 975	99	1 in 97500	•			•	
<i>SGCA</i>	Limb-girdle muscular dystrophy	AR	Worldwide	1 in 517	99	1 in 51700	•			•	
<i>SGCB</i>	Limb-girdle muscular dystrophy	AR	Worldwide	1 in 2069	99	1 in 206900	•			•	
<i>SGCD</i>	Limb-girdle muscular dystrophy	AR	Worldwide	1 in 4269	99	1 in 426900	•			•	
<i>SGCG</i>	Limb-girdle muscular dystrophy	AR	Worldwide	1 in 2781	99	1 in 278100	•			•	
<i>SGSH</i>	Mucopolysaccharidosis (Sanfilippo syndrome)	AR	Worldwide	1 in 791	99	1 in 79100	•			•	
<i>SLC12A1</i>	Antenatal Bartter syndrome	AR	Worldwide	1 in 900	99	1 in 90000	•			•	
<i>SLC12A3</i>	Gitelman syndrome	AR	Worldwide	1 in 364	99	1 in 36400	•			•	
<i>SLC12A6</i>	Agenesis of the corpus callosum with peripheral neuropathy (Andermann syndrome)	AR	Worldwide	1 in 4476	99	1 in 447600	•			•	
<i>SLC17A5</i>	Salla disease	AR	Worldwide	1 in 899	99	1 in 89900	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>SLC19A2</i>	Thiamine-responsive megaloblastic anemia syndrome	AR	Worldwide	1 in 2700	99	1 in 270000	•			•	
<i>SLC19A3</i>	Thiamine metabolism dysfunction syndrome	AR	Worldwide	1 in 1165	99	1 in 116500	•	•		•	•
<i>SLC1A4</i>	Spastic tetraplegia, thin corpus callosum, and progressive microcephaly	AR	Worldwide	1 in 2785	99	1 in 278500	•			•	
<i>SLC22A5</i>	Primary carnitine deficiency	AR	Worldwide	1 in 365	98	1 in 18250	•			•	
<i>SLC25A13</i>	Citrin deficiency	AR	Worldwide	1 in 712	99	1 in 71200	•			•	
<i>SLC25A15[#]</i>	Hyperornithinemia-hyperammonemia-homocitrullinemia syndrome	AR	Worldwide	1 in 2144	99	1 in 214400	•			•	
<i>SLC25A20</i>	Carnitine-acylcarnitine translocase deficiency	AR	Worldwide	1 in 1137	99	1 in 113670	•			•	
<i>SLC26A2</i>	SLC26A2 related skeletal dysplasias	AR	Worldwide	1 in 356	99	1 in 35600	•	•		•	•
<i>SLC26A3</i>	Congenital Secretory Chloride Diarrhea 1	AR	Worldwide	1 in 491	99	1 in 49050	•			•	
<i>SLC26A4</i>	Pendred syndrome	AR	Worldwide	1 in 128	99	1 in 12800	•	•		•	•
<i>SLC35A2</i>	Congenital disorder of glycosylation	XL	Worldwide	1 in 72000	99	1 in 7200000	•			•	
<i>SLC35A3</i>	Arthrogryposis, impaired intellectual development, and seizures	AR	Worldwide	1 in 4922	99	1 in 492200	•		•	•	
<i>SLC37A4</i>	Glycogen storage disease	AR	Worldwide	1 in 1088	98	1 in 54400	•	•		•	•
<i>SLC39A4</i>	Acrodermatitis enteropathica	AR	Worldwide	1 in 755	99	1 in 75500	•			•	
<i>SLC6A8[#]</i>	Creatine deficiency syndrome	XLR	Worldwide	1 in 50000	97	1 in 1666667	•	•			
<i>SLC7A7</i>	Lysinuric protein intolerance	AR	Worldwide	1 in 827	98	1 in 41350	•			•	
<i>SMARCAL1</i>	Schimke immuno-osseous dysplasia	AR	Worldwide	1 in 1127	99	1 in 112700	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>SMN1</i> # *Analysis includes only SMN1 copy number analysis, SNV's are not tested in this gene. "Silent" carriers of SMA cannot be detected, i.e., individuals with two copies of SMN1 in one allele, and zero copies in the other allele.	Spinal muscular atrophy	AR	Worldwide	1 in 55	95	1 in 1100	•	•	•	•	•
<i>SMPD1</i>	Niemann-Pick disease	AR	Worldwide	1 in 416	99	1 in 41600	•	•	•	•	•
<i>SPG11</i>	Spastic paraplegia and Amyotrophic lateral sclerosis	AR	Worldwide	1 in 270	97	1 in 9000	•			•	
<i>SPG7</i>	Spastic paraplegia	AR	Worldwide	1 in 232	99	1 in 23220	•			•	
<i>SPINK5</i>	Netherton syndrome	AR	Worldwide	1 in 1581	99	1 in 158100	•			•	
<i>SPR</i>	Sepiapterin reductase deficiency (Dopa-responsive dystonia)	AR	Worldwide	1 in 1256	95	1 in 25110	•			•	
<i>SRD5A2</i>	Steroid 5-alpha-reductase 2 deficiency	AR	Worldwide	1 in 282	99	1 in 28170	•			•	
<i>ST3GAL5</i>	Ganglioside GM3 synthase deficiency	AR	Worldwide	1 in 1137	92	1 in 14209	•			•	
<i>STAR</i>	Lipoid adrenal hyperplasia	AR	Worldwide	1 in 2899	99	1 in 289900	•			•	
<i>STS</i>	Steroid sulfatase deficiency	XLR	Worldwide	1 in 4500	99	1 in 450000	•				
<i>STXBP2</i>	Familial hemophagocytic lymphohistiocytosis	AR	Worldwide	1 in 720	99	1 in 72000	•			•	
<i>SUMF1</i>	Multiple sulfatase deficiency	AR	Worldwide	1 in 1694	99	1 in 169400	•			•	
<i>SURF1</i>	Charcot-Marie-Tooth disease Leigh syndrome	AR	Worldwide	1 in 635	99	1 in 63540	•		•	•	
<i>SYNE4</i>	Deafness	AR	Worldwide	1 in 2400	99	1 in 240030	•			•	
<i>TANGO2</i>	Metabolic encephalomyopathic crises, recurrent, with rhabdomyolysis, cardiac arrhythmias, and neurodegeneration (MECRCN)	AR	Worldwide	1 in 1440	99	1 in 144000	•			•	
<i>TAT</i>	Tyrosinemia type II	AR	Worldwide	1 in 2354	99	1 in 235400	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>TBCE</i>	Progressive encephalopathy with amyotrophy and optic atrophy (PEAMO)	AR	Worldwide	1 in 2400	99	1 in 240030	•			•	
<i>TCIRG1</i>	Osteopetrosis, severe neonatal or infantile forms (OPTB1)	AR	Worldwide	1 in 1285	99	1 in 128500	•		•	•	
<i>TCTN2</i>	Joubert syndrome and Meckel syndrome	AR	Worldwide	1 in 3652	99	1 in 365200	•			•	
<i>TECPR2</i>	Spastic paraplegia 49	AR	Worldwide	1 in 2844	99	1 in 284400	•			•	
<i>TECRL</i>	Catecholaminergic polymorphic ventricular tachycardia (CPVT)	AR	Worldwide	1 in 1661	99	1 in 166140	•	•		•	•
<i>TF</i>	Atransferrinemia	AR	Worldwide	1 in 398	99	1 in 39800	•			•	
<i>TFR2</i>	Hemochromatosis	AR	Worldwide	1 in 1047	99	1 in 104700	•			•	
<i>TG</i>	Thyroid dysmorphogenesis	AR	Worldwide	1 in 360	97	1 in 12000	•			•	
<i>TGM1</i>	Ichthyosis, congenital	AR	Worldwide	1 in 315	99	1 in 31500	•			•	
<i>TH</i>	Segawa syndrome	AR	Worldwide	1 in 1688	99	1 in 168800	•			•	
<i>TMEM107</i>	Joubert syndrome	AR	Worldwide	1 in 1800	99	1 in 180000	•			•	
<i>TMEM216</i>	Joubert syndrome and Meckel syndrome	AR	Worldwide	1 in 1843	99	1 in 184300	•	•	•	•	•
<i>TMEM231</i>	Joubert syndrome and Meckel syndrome	AR	Worldwide	1 in 831	89	1 in 7552	•			•	
<i>TMEM67</i>	Joubert syndrome, Meckel syndrome, Nephronophthisis and COACH syndrome	AR	Worldwide	1 in 288	96	1 in 7200	•			•	
<i>TMPRSS3</i>	Deafness	AR	Worldwide	1 in 122	95	1 in 2448	•			•	
<i>TPO</i>	Thyroid dysmorphogenesis	AR	Worldwide	1 in 293	99	1 in 29340	•			•	
<i>TPP1</i>	Neuronal ceroid lipofuscinosis type 2	AR	Worldwide	1 in 516	99	1 in 51600	•	•		•	•
<i>TRAF3IP1</i>	Senior-Loken syndrome 9	AR	Worldwide	1 in 1186	99	1 in 118620	•			•	
<i>TREX1</i>	Aicardi-Goutières syndrome	AR	Worldwide	1 in 831	99	1 in 83070	•			•	
<i>TRIM37</i>	Mulibrey nanism	AR	Worldwide	1 in 360	99	1 in 36000	•			•	
<i>TSEN2</i> Excluded from the analysis: NM_001321278, exon 12	Pontocerebellar hypoplasia	AR	Worldwide	1 in 6353	87	1 in 48870	•			•	
<i>TSEN34</i>	Pontocerebellar hypoplasia	AR	Worldwide	1 in 13500	99	1 in 1350000	•			•	
<i>TSEN54</i>	Pontocerebellar hypoplasia	AR	Worldwide	1 in 416	95	1 in 8316	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>TSFM</i> Excluded from the analysis: NM_001172696, exon 5	Combined oxidative phosphorylation deficiency	AR	Worldwide	1 in 772	92	1 in 9650	•			•	
<i>TTC37</i>	Trichohepatoenteric syndrome	AR	Worldwide	1 in 720	99	1 in 72000	•			•	
<i>TTC8</i>	Bardet-Biedl syndrome and Retinitis pigmentosa	AR	Worldwide	1 in 3600	99	1 in 360000	•			•	
<i>TTPA</i>	Ataxia with isolated vitamin E deficiency	AR	Worldwide	1 in 869	99	1 in 86900	•			•	
<i>TWNK</i>	Mitochondrial DNA depletion syndrome and Perrault syndrome	AR	Worldwide	1 in 831	99	1 in 83069	•			•	
<i>TYMP</i>	Mitochondrial DNA depletion syndrome	AR	Worldwide	1 in 1286	99	1 in 128600	•			•	
<i>TYR#</i>	Oculocutaneous albinism	AR	Worldwide	1 in 90	98	1 in 4500	•	•		•	•
<i>UBR1</i>	Johanson-Blizzard syndrome	AR	Worldwide	1 in 5400	99	1 in 540000	•			•	
<i>UNC13D</i>	Familial hemophagocytic lymphohistiocytosis	AR	Worldwide	1 in 600	90	1 in 6003	•			•	
<i>USH1C</i>	Deafness Usher syndrome type IC	AR	Worldwide	1 in 981	97	1 in 32700	•			•	
<i>USH2A</i>	Retinitis pigmentosa 39 and Usher syndrome type 2A	AR	Worldwide	1 in 127	99	1 in 12700	•	•		•	•
<i>VPS13A</i>	Choreoacanthocytosis	AR	Worldwide	1 in 812	97	1 in 27067	•		•	•	
<i>VPS13B</i>	Cohen syndrome	AR	Worldwide	1 in 1006	99	1 in 100600	•			•	
<i>VPS45</i> Excluded from the analysis: NM_001279353, exon 13	Severe congenital neutropenia type 5	AR	Worldwide	1 in 4227	97	1 in 140900	•			•	
<i>VPS53</i>	Pontocerebellar hypoplasia type 2E	AR	Worldwide	1 in 3869	99	1 in 386900	•			•	
<i>VRK1</i>	Pontocerebellar hypoplasia	AR	Worldwide	1 in 4215	99	1 in 421500	•		•	•	
<i>XPA</i>	Xeroderma pigmentosum	AR	Worldwide	1 in 4687	99	1 in 468700	•			•	
<i>XPC</i>	Xeroderma pigmentosum	AR	Worldwide	1 in 952	99	1 in 95200	•	•		•	•
<i>ZFYVE26</i>	Spastic paraplegia 15	AR	Worldwide	1 in 854	99	1 in 85400	•			•	
<i>ZNF469</i>	Brittle cornea syndrome	AR	Worldwide	1 in 3085	99	1 in 308520	•			•	

Gene	Phenotypes	Inheritance model	Ethnicity	Carrier frequency	Detection rate	Residual risk	Comprehensive Reproductive Screen (DUO*) 461 genes	Core Reproductive Screen (DUO*) 126 genes	Ashkenazi Jewish Reproductive Screen (DUO*) 67 genes	Comprehensive Reproductive Male Screen 433 genes	Core Reproductive Male Screen 114 genes
<i>ZNHIT3</i> Excluded from the analysis: ZNHIT3 NM_001281432, exon 5	PEHO syndrome	AR	Worldwide	1 in 771	77	1 in 3353	•			•	

Some genes in the Comprehensive Reproductive Screen have exons that are excluded from our target regions due to suboptimal coverage:

ADAMTS2 (NM_021599, exon 11)
 CC2D2A (NM_020785, exon 7)
 CHM (NM_001145414, exon 5)
 KIAA0586 (NM_001244189, exons 6 and 33)
 MCPH1 (NM_001322042, exon 14)
 PCCB (NM_001178014, exon 4)
 RPGRIP1L (NM_015272, exon 23)
 TSEN2 (NM_001321278, exon 12)
 TSFM (NM_001172696, exon 5)
 VPS45 (NM_001279353, exon 13)
 ZNHIT3 (NM_001281432, exon 5)

Special reporting policies:

CFTR: The 5T and associated TG repeat variants are not reported as their relevance in relation to classical cystic fibrosis is unclear and this test is not intended to screen for risk of congenital bilateral absence of the vas deferens (CBAVD) or CFTR-related pancreatitis risk.

For the CYP21A2 gene we report only specific variants listed below:

c.955C>T	p.(Gln319*)
c.844G>T	p.(Val282Leu)
c.293-13C>G	
c.1360C>T	p.(Pro454Ser)
c.518T>A	p.(Ile173Asn)
c.1447C>T	p.(Pro483Ser)
c.92C>T	p.(Pro31Leu)

TYR: The c.1205G>A, p.(Arg402Gln) (NM_000372.5) hypomorphic variant is typically associated with mild skin/hair/eye pigmentation changes and is therefore not reported as this test is intended to identify variants that cause severe TYR-related oculocutaneous albinism.

NGS doesn't usually detect inversions, gene conversions, balanced translocations, repeat expansions or, non-coding variants deeper than +/- 20 bps from exon-intron boundary unless otherwise indicated.

#Some, or all, of the gene is duplicated in the genome.

More information at www.blueprintgenetics.com.

Contact us

support@blueprintgenetics.com

Phone: +358 40 2511 372

Fax: +358 9 8565 7177