

Whole Exome Sequencing

Gene package Neuronal migration disorders, version 3, 1-2-2018



Technical information

DNA was enriched using Agilent SureSelect Clinical Research Exome V2 capture and paired-end sequenced on the Illumina platform (outsourced). The aim is to obtain 8.1 Giga base pairs per exome with a mapped fraction of 0.99. The average coverage of the exome is ~50x. Duplicate reads are excluded. Data are demultiplexed with bcl2fastq Conversion Software from Illumina. Reads are mapped to the genome using the BWA-MEM algorithm (reference: <http://bio-bwa.sourceforge.net/>). Variant detection is performed by the Genome Analysis Toolkit HaplotypeCaller (reference: <http://www.broadinstitute.org/gatk/>). The detected variants are filtered and annotated with Cartagenia software and classified with Alamut Visual. Additionally, MPLA analysis was performed for several (fragments of) genes involved in neuronal migration disorders (SALSA P061 Lissencephaly; MRC Holland). It is not excluded that pathogenic mutations are being missed using this technology. At this moment, there is not enough information about the sensitivity of this technique with respect to the detection of deletions and duplications of more than 5 nucleotides and of somatic mosaic mutations (all



Dept. Clinical Genetics

| HGNC approved gene symbol | Phenotype description including OMIM phenotype ID(s) | OMIM gene ID | Transcript | median depth | % covered >10x | % covered >20x | % covered >30x |
|---------------------------|---|--------------|----------------|--------------|----------------|----------------|----------------|
| ACTB | Baraitser-Winter syndrome 1, 243310?Dystonia, juvenile-onset, 607371 | 102630 | NM_0011101.3 | 135 | 100 | 100 | 100 |
| ACTG1 | Baraitser-Winter syndrome 2, 614583 Deafness 20/26, 604717 | 102560 | NM_001199954.1 | 121 | 100 | 100 | 100 |
| ADA2 | Polyarteritis nodosa, childhood-onset, 615688 ?Sneddon syndrome, 182410 | 607575 | NM_001282225.1 | 62 | 100 | 99 | 87 |
| ADAR | Aicardi-Goutieres syndrome 6, 615010 Dyschromatosis symmetrica hereditaria, 127400 | 146920 | NM_0011111.4 | 67 | 100 | 100 | 98 |
| ADGRG1 | Polymicrogyria, bilateral frontoparietal, 606854 Polymicrogyria, bilateral perisylvian, 615752 | 604110 | NM_005682.6 | 67 | 100 | 100 | 98 |
| AKT1 | Breast cancer, somatic, 114480 Colorectal cancer, somatic, 114500 Cowden syndrome 6, 615109 Ovarian cancer, somatic, 167000 Proteus syndrome, somatic, 176920 {Schizophrenia, susceptibility to}, 181500 | 164730 | NM_005163.2 | 72 | 100 | 100 | 97 |
| AKT3 | Megalencephaly-polymicrogyria-polydactyly-hydrocephalus syndrome 2, 615937 | 611223 | NM_005465.4 | 49 | 100 | 94 | 75 |
| ANKLE2 | ?Microcephaly 16, primary, 616681 | 616062 | NM_015114.2 | 70 | 99 | 93 | 87 |
| AP1S2 | Mental retardation syndromic 5, 304340 | 300629 | NM_001272071.1 | 44 | 100 | 80 | 65 |
| AP3B2 | Epileptic encephalopathy, early infantile, 48, 617276 | 602166 | NM_001278512.1 | 72 | 100 | 99 | 95 |
| AP4B1 | Spastic paraplegia 47, 614066 | 607245 | NM_006594.3 | 60 | 100 | 100 | 96 |
| AP4E1 | Spastic paraplegia 51, 613744 Stuttering, familial persistent, 1, 184450 | 607244 | NM_007347.4 | 50 | 100 | 99 | 87 |

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|---------------------------|---|--------------|----------------|--------------|----------------|----------------|----------------|
| AP4M1 | Spastic paraplegia 50, 612936 | 602296 | NM_004722.3 | 75 | 100 | 100 | 98 |
| AP4S1 | Spastic paraplegia 52, 614067 | 607243 | NM_007077.4 | 37 | 100 | 95 | 69 |
| ARFGEF2 | Periventricular heterotopia with microcephaly, 608097 | 605371 | NM_006420.2 | 55 | 100 | 98 | 85 |
| ARNT2 | ?Webb-Dattani syndrome, 615926 | 606036 | NM_014862.3 | 55 | 100 | 98 | 88 |
| ARX | Epileptic encephalopathy, early infantile, 1, 308350 Hydranencephaly with abnormal genitalia, 300215 Lissencephaly 2, 300215 Mental retardation 29 and others, 300419 Partington syndrome, 309510 Proud syndrome, 300004 | 300382 | NM_139058.2 | 42 | 89 | 80 | 65 |
| ASNS | Asparagine synthetase deficiency, 615574 | 108370 | NM_133436.3 | 59 | 100 | 100 | 92 |
| ASPM | Microcephaly 5, primary, 608716 | 605481 | NM_018136.4 | 62 | 100 | 100 | 96 |
| ASXL1 | Bohring-Opitz syndrome, 605039 Myelodysplastic syndrome, somatic, 614286 | 612990 | NM_015338.5 | 77 | 100 | 100 | 97 |
| ATAD3A | Harel-Yoon syndrome, 617183 | 612316 | NM_018188.3 | 89 | 100 | 98 | 97 |
| ATAD3B | No OMIM phenotype | 612317 | NM_031921.5 | 95 | 99 | 98 | 96 |
| ATP6V0A2 | Cutis laxa, type IIA, 219200 Wrinkly skin syndrome, 278250 | 611716 | NM_012463.3 | 56 | 100 | 98 | 87 |
| ATR | ?Cutaneous telangiectasia and cancer syndrome, familial, 614564 Seckel syndrome 1, 210600 | 601215 | NM_001184.3 | 58 | 100 | 96 | 78 |
| ATRIP | No OMIM phenotype | 606605 | NM_130384.2 | 73 | 100 | 98 | 92 |
| B3GALNT2 | Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies, type A, 11, 615181 | 610194 | NM_152490.4 | 42 | 100 | 97 | 76 |
| B4GAT1 | Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 13, 615287 | 605517 | NM_006876.2 | 114 | 100 | 100 | 100 |
| CASK | FG syndrome 4, 300422 Mental retardation and microcephaly with pontine and cerebellar hypoplasia, 300749 Mental retardation, with or without nystagmus, 300422 | 300172 | NM_003688.3 | 42 | 100 | 95 | 69 |
| CCND2 | Megalencephaly-polymicrogyria-polydactyly-hydrocephalus syndrome 3, 615938 | 123833 | NM_001759.3 | 68 | 100 | 100 | 98 |
| CDK5 | ?Lissencephaly 7 with cerebellar hypoplasia, 616342 | 123831 | NM_004935.3 | 78 | 100 | 100 | 100 |
| CDK5RAP2 | Microcephaly 3, primary, 604804 | 608201 | NM_018249.5 | 47 | 100 | 95 | 78 |
| CDK6 | ?Microcephaly 12, primary, 616080 | 603368 | NM_001145306.1 | 49 | 100 | 98 | 90 |
| CENPJ | Microcephaly 6, primary, 608393 ?Seckel syndrome 4, 613676 | 609279 | NM_018451.4 | 60 | 100 | 100 | 96 |
| CEP135 | Microcephaly 8, primary, 614673 | 611423 | NM_025009.4 | 61 | 100 | 96 | 80 |
| CEP152 | Microcephaly 9, primary, 614852 Seckel syndrome 5, 613823 | 613529 | NM_001194998.1 | 52 | 100 | 97 | 84 |
| CEP63 | ?Seckel syndrome 6, 614728 | 614724 | NM_025180.3 | 54 | 100 | 95 | 78 |
| CHMP1A | Pontocerebellar hypoplasia, type 8, 614961 | 164010 | NM_001083314.3 | 56 | 100 | 100 | 96 |
| CIT | Microcephaly 17, primary, 617090 | 605629 | NM_001206999.1 | 61 | 100 | 97 | 85 |
| CLP1 | Pontocerebellar hypoplasia, type 10, 615803 | 608757 | NM_006831.2 | 70 | 100 | 100 | 100 |
| COL18A1 | Knobloch syndrome, type 1, 267750 | 120328 | NM_130445.2 | 76 | 100 | 100 | 95 |

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|---------------------------|---|--------------|----------------|--------------|----------------|----------------|----------------|
| COL4A1 | Angiopathy, hereditary, with nephropathy, aneurysms, and muscle cramps, 611773 Brain small vessel disease with or without ocular anomalies, 607595 {Hemorrhage, intracerebral, susceptibility to}, 614519 Porencephaly 1, 175780 ?Retinal arteries, tortuosity of, 180000 | 120130 | NM_001845.4 | 61 | 100 | 99 | 91 |
| COL4A2 | {Hemorrhage, intracerebral, susceptibility to}, 614519 Porencephaly 2, 614483 | 120090 | NM_001846.2 | 67 | 100 | 100 | 96 |
| CRADD | Mental retardation 34, with variant lissencephaly, 614499 | 603454 | NM_003805.3 | 96 | 100 | 100 | 100 |
| CRB2 | Focal segmental glomerulosclerosis 9, 616220 Ventriculomegaly with cystic kidney disease, 219730 | 609720 | NM_173689.6 | 76 | 100 | 100 | 99 |
| CSTB | Epilepsy, progressive myoclonic 1A (Unverricht and Lundborg), 254800 | 601145 | NM_000100.3 | 62 | 100 | 100 | 98 |
| CTC1 | Cerebroretinal microangiopathy with calcifications and cysts, 612199 | 613129 | NM_025099.5 | 72 | 100 | 100 | 98 |
| CTNND2 | No OMIM phenotype | 604275 | NM_001332.3 | 66 | 99 | 95 | 87 |
| DAB1 | No OMIM phenotype | 603448 | NM_021080.3 | 59 | 100 | 100 | 93 |
| DAG1 | Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 9, 616538 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 9, 613818 | 128239 | NM_001165928.3 | 103 | 100 | 100 | 100 |
| DCHS1 | Mitral valve prolapse 2, 607829 Van Maldergem syndrome 1, 601390 | 603057 | NM_003737.2 | 94 | 100 | 100 | 100 |
| DCX | Lissencephaly, 300067 Subcortical laminar heteropia, 300067 | 300121 | NM_178153.2 | 47 | 100 | 96 | 83 |
| DEPDC5 | Epilepsy, familial focal, with variable foci 1, 604364 | 614191 | NM_001242896.1 | 56 | 100 | 98 | 90 |
| DKC1 | Dyskeratosis congenita, 305000 | 300126 | NM_001363.4 | 45 | 100 | 96 | 83 |
| DNMT3A | Tatton-Brown-Rahman syndrome, 615879 | 602769 | NM_175629.2 | 67 | 100 | 100 | 96 |
| DYNC1H1 | Charcot-Marie-Tooth disease, axonal, type 20, 614228 Mental retardation 13, 614563 Spinal muscular atrophy, lower extremity-predominant 1, AD, 158600 | 600112 | NM_001376.4 | 69 | 100 | 99 | 95 |
| EIF2AK3 | Wolcott-Rallison syndrome, 226980 | 604032 | NM_004836.5 | 53 | 100 | 95 | 83 |
| EMG1 | Bowen-Conradi syndrome, 211180 | 611531 | NM_006331.7 | 57 | 100 | 100 | 100 |
| EML1 | Band heterotopia, 600348 | 602033 | NM_001008707.1 | 50 | 100 | 97 | 85 |
| EOMES | No OMIM phenotype | 604615 | NM_005442.3 | 63 | 100 | 100 | 98 |
| ERCC1 | Cerebrooculofacioskeletal syndrome 4, 610758 | 126380 | NM_001983.3 | 52 | 100 | 100 | 95 |
| ERCC2 | Cerebrooculofacioskeletal syndrome 2, 610756 Trichothiodystrophy 1, photosensitive, 601675 Xeroderma pigmentosum, group D, 278730 | 126340 | NM_000400.3 | 75 | 100 | 100 | 99 |
| ERCC5 | Cerebrooculofacioskeletal syndrome 3, 616570 Xeroderma pigmentosum, group G, 278780 Xeroderma pigmentosum, group G/Cockayne syndrome, 278780 | 133530 | NM_000123.3 | 65 | 100 | 99 | 94 |

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|---------------------------|--|--------------|----------------|--------------|----------------|----------------|----------------|
| ERCC6 | Cerebrooculofacioskeletal syndrome 1, 214150 Cockayne syndrome, type B, 133540 De Sanctis-Cacchione syndrome, 278800 {Lung cancer, susceptibility to}, 211980 {Macular degeneration, age-related, susceptibility to, 5}, 613761 Premature ovarian failure 11, 616946 UV-sensitive syndrome 1, 600630 | 609413 | NM_000124.3 | 60 | 100 | 98 | 92 |
| ERMARD | ?Periventricular nodular heterotopia 6, 615544 | 615532 | NM_018341.2 | 51 | 100 | 98 | 87 |
| FAT4 | Hennekam lymphangiectasia-lymphedema syndrome 2, 616006 Van Maldergem syndrome 2, 615546 | 612411 | NM_024582.4 | 69 | 100 | 100 | 98 |
| FIG4 | Amyotrophic lateral sclerosis 11, 612577 Charcot-Marie-Tooth disease, type 4J, 611228 ?Polymicrogyria, bilateral temporooccipital, 612691 Yunis-Varon syndrome, 216340 | 609390 | NM_014845.5 | 40 | 100 | 96 | 77 |
| FKRP | Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 5, 613153 Muscular dystrophy-dystroglycanopathy (congenital with or without mental retardation), type B, 5, 606612 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 5, 607155 | 606596 | NM_001039885.2 | 99 | 100 | 100 | 100 |
| FKTN | Cardiomyopathy, dilated, 1X, 611615 Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 4, 253800 Muscular dystrophy-dystroglycanopathy (congenital without mental retardation), type B, 4, 613152 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 4, 611588 | 607440 | NM_006731.2 | 65 | 100 | 100 | 97 |
| FLNA | Cardiac valvular dysplasia, 314400 Congenital short bowel syndrome, 300048 FG syndrome 2, 300321 Frontometaphyseal dysplasia 1, 305620 Heterotopia, periventricular, 300049 Intestinal pseudoobstruction, neuronal, 300048 Melnick-Needles syndrome, 309350 Otopalatodigital syndrome, type I, 311300 Otopalatodigital syndrome, type II, 304120 Terminal osseous dysplasia, 300244 | 300017 | NM_001110556.1 | 79 | 100 | 100 | 100 |
| FLVCR2 | Proliferative vasculopathy and hydraencephaly-hydrocephaly syndrome, 225790 | 610865 | NM_017791.2 | 81 | 100 | 100 | 87 |
| FRMD4A | ?Corpus callosum, agenesis of, with facial anomalies and cerebellar ataxia, 616819 | 616305 | NM_018027.3 | 55 | 100 | 95 | 82 |
| GNAQ | Capillary malformations, congenital, 1, somatic, mosaic, 163000 Sturge-Weber syndrome, somatic, mosaic, 185300 | 600998 | NM_002072.4 | 68 | 100 | 100 | 96 |
| HNRNP | Au-Kline syndrome, 616580 | 600712 | NM_002140.3 | 35 | 95 | 77 | 52 |
| IBA57 | ?Multiple mitochondrial dysfunctions syndrome 3, 615330 ?Spastic paraplegia 74, 616451 | 615316 | NM_001010867.2 | 89 | 100 | 100 | 98 |
| IER3IP1 | Microcephaly, epilepsy, and diabetes syndrome, 614231 | 609382 | NM_016097.4 | 43 | 100 | 86 | 75 |
| IFIH1 | Aicardi-Goutieres syndrome 7, 615846 Singleton-Merten syndrome 1, 182250 | 606951 | NM_022168.3 | 65 | 100 | 99 | 90 |

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|---------------------------|---|--------------|----------------|--------------|----------------|----------------|----------------|
| INTS8 | No OMIM phenotype | 611351 | NM_017864.3 | 38 | 100 | 93 | 62 |
| ISPD | Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 7, 614643 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 7, 616052 | 614631 | NM_001101426.3 | 58 | 100 | 99 | 84 |
| ITSN1 | No OMIM phenotype | 602442 | NM_003024.2 | 44 | 99 | 94 | 77 |
| JAM3 | Hemorrhagic destruction of the brain, subependymal calcification, and cataracts, 613730 | 606871 | NM_032801.4 | 49 | 100 | 97 | 83 |
| KATNB1 | Lissencephaly 6, with microcephaly, 616212 | 602703 | NM_005886.2 | 80 | 100 | 100 | 100 |
| KIF11 | Microcephaly with or without chorioretinopathy, lymphedema, or mental retardation, 152950 | 148760 | NM_004523.3 | 58 | 100 | 99 | 87 |
| KIF1BP | Goldberg-Shprintzen megacolon syndrome, 609460 | 609367 | NM_015634.3 | 66 | 100 | 99 | 91 |
| KIF2A | Cortical dysplasia, complex, with other brain malformations 3, 615411 | 602591 | NM_001098511.2 | 59 | 100 | 94 | 75 |
| KIF5C | Cortical dysplasia, complex, with other brain malformations 2, 615282 | 604593 | NM_004522.2 | 45 | 100 | 95 | 76 |
| KIF7 | Acrocallosal syndrome, 200990 ?Al-Gazali-Bakalnova syndrome, 607131 ?Hydroletharus syndrome 2, 614120 Joubert syndrome 12, 200990 | 611254 | NM_198525.2 | 66 | 99 | 95 | 89 |
| KNL1 | Microcephaly 4, primary, 604321 | 609173 | NM_170589.4 | 51 | 100 | 98 | 91 |
| KPTN | Mental retardation 41, 615637 | 615620 | NM_007059.3 | 73 | 100 | 100 | 94 |
| L1CAM | CRASH syndrome, 303350 Corpus callosum, partial agenesis of, 304100 Hydrocephalus due to aqueductal stenosis, 307000 Hydrocephalus with Hirschsprung disease, 307000 Hydrocephalus with congenital idiopathic intestinal pseudoobstruction, 307000 MASA syndrome, 303350 | 308840 | NM_000425.4 | 78 | 100 | 100 | 99 |
| LAMA1 | Poretti-Boltshauser syndrome, 615960 | 150320 | NM_005559.3 | 57 | 100 | 98 | 89 |
| LAMA2 | Muscular dystrophy, congenital merosin-deficient, 607855 Muscular dystrophy, congenital, due to partial LAMA2 deficiency, 607855 | 156225 | NM_000426.3 | 50 | 100 | 98 | 86 |
| LAMB1 | Lissencephaly 5, 615191 | 150240 | NM_002291.2 | 65 | 100 | 98 | 93 |
| LAMC1 | No OMIM phenotype | 150290 | NM_002293.3 | 54 | 100 | 98 | 88 |
| LAMC3 | Cortical malformations, occipital, 614115 | 604349 | NM_006059.3 | 67 | 100 | 100 | 98 |
| LARGE1 | Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 6, 613154 Muscular dystrophy-dystroglycanopathy (congenital with mental retardation), type B, 6, 608840 | 603590 | NM_004737.4 | 63 | 100 | 100 | 94 |
| LARP7 | Alazami syndrome, 615071 | 612026 | NM_001267039.1 | 50 | 100 | 98 | 85 |
| MCPH1 | Microcephaly 1, primary, 251200 | 607117 | NM_024596.3 | 59 | 94 | 94 | 85 |
| MED17 | Microcephaly, postnatal progressive, with seizures and brain atrophy, 613668 | 603810 | NM_004268.4 | 69 | 100 | 100 | 94 |
| MPDZ | Hydrocephalus, nonsyndromic 2, 615219 | 603785 | NM_003829.4 | 49 | 100 | 97 | 85 |
| MYCN | Feingold syndrome 1, 164280 | 164840 | NM_005378.4 | 79 | 100 | 97 | 93 |
| NBN | Aplastic anemia, 609135 Leukemia, acute lymphoblastic, 613065 Nijmegen breakage syndrome, 251260 | 602667 | NM_002485.4 | 52 | 100 | 95 | 76 |
| NDE1 | Lissencephaly 4 (with microcephaly), 614019 ?Microhydranencephaly, 605013 | 609449 | NM_001143979.1 | 61 | 100 | 99 | 92 |
| NID1 | No OMIM phenotype | 131390 | NM_002508.2 | 74 | 100 | 100 | 99 |

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|---------------------------|---|--------------|----------------|--------------|----------------|----------------|----------------|
| NIN | ?Seckel syndrome 7, 614851 | 608684 | NM_020921.3 | 56 | 100 | 98 | 84 |
| NPRL3 | Epilepsy, familial focal, with variable foci 3, 617118 | 600928 | NM_001077350.2 | 54 | 100 | 97 | 89 |
| NSDHL | CHILD syndrome, 308050 CK syndrome, 300831 | 300275 | NM_015922.2 | 55 | 100 | 96 | 80 |
| OCLN | Pseudo-TORCH syndrome 1, 251290 | 602876 | NM_002538.3 | 65 | 100 | 100 | 97 |
| PAFAH1B1 | Lissencephaly 1, 607432 Subcortical laminar heterotopia, 607432 | 601545 | NM_000430.3 | 62 | 100 | 98 | 88 |
| PAX6 | Aniridia, 106210 Anterior segment dysgenesis 5, multiple subtypes, 604229 Cataract with late-onset corneal dystrophy, 106210 ?Coloboma of optic nerve, 120430 ?Coloboma, ocular, 120200 Foveal hypoplasia 1, 136520 Keratitis, 148190 ?Morning glory disc anomaly, 120430 Optic nerve hypoplasia, 165550 | 607108 | NM_001604.5 | 50 | 100 | 98 | 84 |
| PCNT | Microcephalic osteodysplastic primordial dwarfism, type II, 210720 | 605925 | NM_006031.5 | 84 | 100 | 99 | 95 |
| PHC1 | ?Microcephaly 11, primary, 615414 | 602978 | NM_004426.2 | 127 | 100 | 100 | 99 |
| PI4KA | Polymicrogyria, perisylvian, with cerebellar hypoplasia and arthrogryposis, 616531 | 600286 | NM_058004.3 | 76 | 100 | 99 | 91 |
| PIK3CA | Breast cancer, somatic, 114480 CLOVE syndrome, somatic, 612918 Colorectal cancer, somatic, 114500 Cowden syndrome 5, 615108 Gastric cancer, somatic, 613659 Hepatocellular carcinoma, somatic, 114550 Keratosis, seborrheic, somatic, 182000 Megalencephaly-capillary malformation-polymicrogyria syndrome, somatic, 602501 Nevus, epidermal, somatic, 162900 Non-small cell lung cancer, somatic, 211980 Ovarian cancer, somatic, 167000 | 171834 | NM_006218.2 | 65 | 100 | 99 | 92 |
| PIK3R2 | Megalencephaly-polymicrogyria-polydactyly-hydrocephalus syndrome 1, 603387 | 603157 | NM_005027.3 | 59 | 96 | 93 | 89 |
| PLK4 | Microcephaly and chorioretinopathy, 2, 616171 | 605031 | NM_014264.4 | 49 | 100 | 96 | 81 |
| PNKP | Ataxia-oculomotor apraxia 4, 616267 Microcephaly, seizures, and developmental delay, 613402 | 605610 | NM_007254.3 | 72 | 100 | 100 | 97 |
| POLR3B | Leukodystrophy, hypomyelinating, 8, with or without oligodontia and/or hypogonadotropic hypogonadism, 614381 | 614366 | NM_018082.5 | 53 | 100 | 97 | 81 |
| POMGNT1 | Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 3, 253280 Muscular dystrophy-dystroglycanopathy (congenital with mental retardation), type B, 3, 613151 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 3, 613157 Retinitis pigmentosa 76, 617123 | 606822 | NM_017739.3 | 63 | 100 | 100 | 95 |

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|---------------------------|---|--------------|----------------|--------------|----------------|----------------|----------------|
| POMT1 | Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 1, 236670 Muscular dystrophy-dystroglycanopathy (congenital with mental retardation), type B, 1, 613155 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 1, 609308 | 607423 | NM_007171.3 | 69 | 100 | 100 | 98 |
| POMT2 | Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 2, 613150 Muscular dystrophy-dystroglycanopathy (congenital with mental retardation), type B, 2, 613156 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 2, 613158 | 607439 | NM_013382.5 | 52 | 100 | 99 | 89 |
| PTEN | Bannayan-Riley-Ruvalcaba syndrome, 153480 Cowden syndrome 1, 158350 Endometrial carcinoma, somatic, 608089 {Glioma susceptibility 2}, 613028 Lhermitte-Duclos syndrome, 158350 Macrocephaly/autism syndrome, 605309 Malignant melanoma, somatic, 155600 {Meningioma}, 607174 PTEN hamartoma tumor syndrome {Prostate cancer, somatic}, 176807 Squamous cell carcinoma, head and neck, somatic, 275355 VATER association with macrocephaly and ventriculomegaly, 276950 | 601728 | NM_000314.4 | 85 | 92 | 81 | 77 |
| PTF1A | Pancreatic agenesis 2, 615935 Pancreatic and cerebellar agenesis, 609069 | 607194 | NM_178161.2 | 78 | 100 | 100 | 97 |
| PYCR2 | Leukodystrophy, hypomyelinating, 10, 616420 | 616406 | NM_013328.3 | 74 | 100 | 100 | 100 |
| QARS | Microcephaly, progressive, seizures, and cerebral and cerebellar atrophy, 615760 | 603727 | NM_005051.2 | 75 | 100 | 100 | 96 |
| RAB18 | Warburg micro syndrome 3, 614222 | 602207 | NM_021252.4 | 71 | 100 | 100 | 88 |
| RAB3GAP1 | Warburg micro syndrome 1, 600118 | 602536 | NM_001172435.1 | 49 | 100 | 99 | 89 |
| RAB3GAP2 | Martsof syndrome, 212720 Warburg micro syndrome 2, 614225 | 609275 | NM_012414.3 | 49 | 100 | 96 | 79 |
| RAD50 | Nijmegen breakage syndrome-like disorder, 613078 | 604040 | NM_005732.3 | 71 | 100 | 100 | 95 |
| RARS2 | Pontocerebellar hypoplasia, type 6, 611523 | 611524 | NM_020320.3 | 47 | 100 | 97 | 76 |
| RBBP8 | Jawad syndrome, 251255 Pancreatic carcinoma, somatic Seckel syndrome 2, 606744 | 604124 | NM_002894.2 | 44 | 100 | 98 | 80 |
| RBM10 | TARP syndrome, 311900 | 300080 | NM_001204468.1 | 61 | 100 | 95 | 91 |
| RELN | {Epilepsy, familial temporal lobe, 7}, 616436 Lissencephaly 2 (Norman-Roberts type), 257320 | 600514 | NM_173054.2 | 52 | 100 | 98 | 85 |
| RNASEH2A | Aicardi-Goutieres syndrome 4, 610333 | 606034 | NM_006397.2 | 73 | 100 | 100 | 99 |
| RNASEH2B | Aicardi-Goutieres syndrome 2, 610181 | 610326 | NM_024570.3 | 48 | 100 | 97 | 81 |
| RNASEH2C | Aicardi-Goutieres syndrome 3, 610329 | 610330 | NM_032193.3 | 139 | 100 | 100 | 100 |
| RNASET2 | Leukoencephalopathy, cystic, without megalencephaly, 612951 | 612944 | NM_003730.4 | 68 | 100 | 100 | 84 |
| RNU4ATAC | Microcephalic osteodysplastic primordial dwarfism, type I, 210710 Roifman syndrome, 616651 | 601428 | NR_023343.1 | No coverage | 0 | 0 | 0 |

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|---------------------------|---|--------------|----------------|--------------|----------------|----------------|----------------|
| RTEL1 | Dyskeratosis congenita 4, 615190 Dyskeratosis congenita 5, 615190 Pulmonary fibrosis and/or bone marrow failure, telomere-related, 3, 616373 | 608833 | NM_001283009.1 | 80 | 100 | 100 | 98 |
| RTTN | Microcephaly, short stature, and polymicrogyria with seizures, 614833 | 610436 | NM_173630.3 | 49 | 100 | 95 | 79 |
| SAMHD1 | Aicardi-Goutieres syndrome 5, 612952 ?Chilblain lupus 2, 614415 | 606754 | NM_015474.3 | 49 | 100 | 93 | 69 |
| SHOC2 | Noonan-like syndrome with loose anagen hair, 607721 | 602775 | NM_001324336.1 | 48 | 100 | 98 | 85 |
| SLC25A19 | Microcephaly, Amish type, 607196 Thiamine metabolism dysfunction syndrome 4 (progressive polyneuropathy type), 613710 | 606521 | NM_001126122.1 | 70 | 100 | 100 | 97 |
| SMPD4 | No OMIM phenotype | 610457 | NM_017951.4 | 80 | 100 | 100 | 96 |
| SNAP29 | Cerebral dysgenesis, neuropathy, ichthyosis, and palmoplantar keratoderma syndrome, 609528 | 604202 | NM_004782.3 | 91 | 100 | 100 | 96 |
| SRPX2 | ?Rolandic epilepsy, mental retardation, and speech dyspraxia, 300643 | 300642 | NM_014467.2 | 47 | 100 | 97 | 85 |
| STAMBP | Microcephaly-capillary malformation syndrome, 614261 | 606247 | NM_006463.4 | 51 | 100 | 100 | 90 |
| STIL | Microcephaly 7, primary, 612703 | 181590 | NM_001048166.1 | 53 | 100 | 99 | 90 |
| STRADA | Polyhydramnios, megalencephaly, and symptomatic epilepsy, 611087 | 608626 | NM_001003787.2 | 67 | 100 | 100 | 92 |
| TBC1D20 | Warburg micro syndrome 4, 615663 | 611663 | NM_144628.3 | 53 | 98 | 93 | 93 |
| TBC1D24 | DOOR syndrome, 220500 Deafness 86, 614617 Deafness 65, 616044 Epileptic encephalopathy, early infantile, 16, 615338 Myoclonic epilepsy, infantile, familial, 605021 | 613577 | NM_001199107.1 | 87 | 100 | 100 | 98 |
| TBC1D7 | Macrocephaly/megalencephaly syndrome, 248000 | 612655 | NM_016495.5 | 39 | 100 | 93 | 69 |
| TMEM5 | Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 10, 615041 | 605862 | NM_014254.2 | 58 | 100 | 98 | 78 |
| TMTC3 | Lissencephaly 8, 617255 | 617218 | NM_181783.3 | 58 | 100 | 98 | 87 |
| TMX2 | No OMIM phenotype | 616715 | NM_015959.3 | 48 | 100 | 93 | 79 |
| TRAIP | Seckel syndrome 9, 616777 | 605958 | NM_005879.2 | 61 | 100 | 100 | 97 |
| TREX1 | Aicardi-Goutieres syndrome 1, dominant and recessive, 225750 Chilblain lupus, 610448 {Systemic lupus erythematosus, susceptibility to}, 152700 Vasculopathy, retinal, with cerebral leukodystrophy, 192315 | 606609 | NM_016381.5 | 116 | 100 | 100 | 100 |
| TSC1 | Focal cortical dysplasia, type II, somatic, 607341 Lymphangioliomyomatosis, 606690 Tuberous sclerosis-1, 191100 | 605284 | NM_000368.4 | 105 | 100 | 100 | 100 |
| TSC2 | ?Focal cortical dysplasia, type II, somatic, 607341 Lymphangioliomyomatosis, somatic, 606690 Tuberous sclerosis-2, 613254 | 191092 | NM_000548.3 | 104 | 100 | 100 | 100 |
| TSEN54 | Pontocerebellar hypoplasia type 2A, 277470 Pontocerebellar hypoplasia type 4, 225753 ?Pontocerebellar hypoplasia type 5, 610204 | 608755 | NM_207346.2 | 67 | 100 | 96 | 95 |
| TUBA1A | Lissencephaly 3, 611603 | 602529 | NM_006009.3 | 89 | 100 | 100 | 100 |
| TUBA8 | Polymicrogyria with optic nerve hypoplasia, 613180 | 605742 | NM_018943.2 | 79 | 100 | 100 | 100 |

| HGNC approved gene symbol | Phenotype description including OMIM phenotype ID(s) | OMIM gene ID | Transcript | median depth | % covered >10x | % covered >20x | % covered >30x |
|---------------------------|--|--------------|----------------|--------------|----------------|----------------|----------------|
| TUBB | Cortical dysplasia, complex, with other brain malformations 6, 615771 Symmetric circumferential skin creases, congenital, 1, 156610 | 191130 | NM_178014.2 | 135 | 100 | 100 | 99 |
| TUBB2A | Cortical dysplasia, complex, with other brain malformations 5, 615763 | 615101 | NM_001069.2 | 170 | 100 | 97 | 95 |
| TUBB2B | Polymicrogyria, symmetric or asymmetric, 610031 | 612850 | NM_178012.4 | 170 | 100 | 100 | 100 |
| TUBB3 | Cortical dysplasia, complex, with other brain malformations 1, 614039 Fibrosis of extraocular muscles, congenital, 3A, 600638 | 602661 | NM_006086.3 | 168 | 100 | 95 | 95 |
| TUBB4A | Dystonia 4, torsion, 128101 Leukodystrophy, hypomyelinating, 6, 612438 | 602662 | NM_006087.3 | 148 | 100 | 100 | 97 |
| TUBG1 | Cortical dysplasia, complex, with other brain malformations 4, 615412 | 191135 | NM_001070.4 | 99 | 100 | 100 | 100 |
| TUBGCP4 | Microcephaly and chorioretinopathy, 3, 616335 | 609610 | NM_001286414.1 | 51 | 100 | 96 | 80 |
| TUBGCP6 | Microcephaly and chorioretinopathy, 1, 251270 | 610053 | NM_020461.3 | 109 | 100 | 100 | 99 |
| VLDLR | Cerebellar hypoplasia and mental retardation with or without quadrupedal locomotion 1, 224050 | 192977 | NM_003383.3 | 48 | 100 | 99 | 87 |
| VPS13B | Cohen syndrome, 216550 | 607817 | NM_017890.4 | 54 | 100 | 98 | 87 |
| VRK1 | Pontocerebellar hypoplasia type 1A, 607596 | 602168 | NM_003384.2 | 42 | 100 | 98 | 81 |
| WASHC5 | Ritscher-Schinzel syndrome 1, 220210 Spastic paraplegia 8, 603563 | 610657 | NM_014846.3 | 45 | 100 | 97 | 78 |
| WDR4 | No OMIM phenotype | 605924 | NM_033661.4 | 63 | 100 | 100 | 94 |
| WDR62 | Microcephaly 2, primary, with or without cortical malformations, 604317 | 613583 | NM_001083961.1 | 79 | 100 | 100 | 98 |
| WDR73 | Galloway-Mowat syndrome, 251300 | 616144 | NM_032856.2 | 83 | 100 | 97 | 91 |
| WDR81 | Cerebellar ataxia, mental retardation, and dysequilibrium syndrome 2, 610185 | 614218 | NM_001163809.1 | 96 | 100 | 100 | 100 |
| YWHAE | No OMIM phenotype | 605066 | NM_006761.4 | 50 | 100 | 99 | 89 |
| ZIC1 | Craniosynostosis 6, 616602 | 600470 | NM_003412.3 | 110 | 100 | 100 | 100 |
| ZIC2 | Holoprosencephaly 5, 609637 | 603073 | NM_007129.3 | 82 | 95 | 90 | 87 |
| ZIC4 | No OMIM phenotype | 608948 | NM_001168378.1 | 92 | 100 | 100 | 97 |

- Gene symbols according HGNC
- OMIM release used: 2-6-2017
- "No OMIM phenotypes" indicates a gene without a current OMIM association
- OMIM phenotypes between "[]", indicate "nondiseases," mainly genetic variations that lead to apparently abnormal laboratory test values
- OMIM phenotypes between "{}", indicate risk factors
- OMIM phenotypes with a question mark, "?", before the disease name indicates an unconfirmed or possibly spurious mapping
- The statistics above are based on a set of 96 samples
- Median depth is the median of the mean sequence depth over the protein coding exons (± 10 bp flanking introns) of the longest transcript
- % Covered 10x describes the percentage of a gene's coding sequence (± 10 bp flanking introns) that is covered at least 10x
- % Covered 20x describes the percentage of a gene's coding sequence (± 10 bp flanking introns) that is covered at least 20x