

Whole Exome Sequencing Gene package Hearing impairment, version 6, 30-9-2019



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. 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 types of sequence changes).



Dept. Clinical Genetics

| HGNC approved gene symbol | Phenotype description including OMIM phenotype ID(s) | OMIM gene ID | median depth | % covered >10x | % covered >20x | % covered >30x |
|---------------------------|--|--------------|--------------|----------------|----------------|----------------|
| ACTB | Baraitser-Winter syndrome 1, 243310?Dystonia, juvenile-onset, 607371 | 102630 | 192 | 100 | 100 | 100 |
| ACTG1 | Baraitser-Winter syndrome 2, 614583 Deafness 20/26, 604717 | 102560 | 188 | 100 | 100 | 100 |
| ADCY1 | ?Deafness 44, 610154 | 103072 | 109 | 97 | 95 | 93 |
| ADGRV1 | ?Febrile seizures, familial, 4, 604352 Usher syndrome, type 2C, 605472 Usher syndrome, type 2C, GPR98/PDZD7 digenic, 605472 | 602851 | 80 | 100 | 100 | 97 |
| AIFM1 | Combined oxidative phosphorylation deficiency 6, 300816 Cowchock syndrome, 310490 Deafness 5, 300614 | 300169 | 55 | 100 | 96 | 82 |
| ATP1A3 | Alternating hemiplegia of childhood 2, 614820 CAPOS syndrome, 601338 Dystonia-12, 128235 | 182350 | 144 | 100 | 100 | 100 |
| ATP2B2 | {Deafness 12, modifier of}, 601386 | 108733 | 139 | 100 | 100 | 100 |
| ATP6V1B1 | Renal tubular acidosis with deafness, 267300 | 192132 | 136 | 100 | 100 | 100 |
| BCS1L | Bjornstad syndrome, 262000 GRACILE syndrome, 603358 Leigh syndrome, 256000 Mitochondrial complex III deficiency, nuclear type 1, 124000 | 603647 | 170 | 100 | 100 | 100 |
| BDP1 | ?Deafness 112, 618257 | 607012 | 60 | 99 | 96 | 85 |

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|---------------------------|---|--------------|------------------|----------------|----------------|----------------|
| BSND | Bartter syndrome, type 4a, 602522 Sensorineural deafness with mild renal dysfunction, 602522 | 606412 | 115 | 100 | 100 | 100 |
| CABP2 | Deafness 93, 614899 | 607314 | 94 | 100 | 100 | 96 |
| CACNA1D | Primary aldosteronism, seizures, and neurologic abnormalities, 615474 Sinoatrial node dysfunction and deafness, 614896 | 114206 | 102 | 100 | 100 | 98 |
| CCDC50 | ?Deafness 44, 607453 | 611051 | 73 | 100 | 100 | 98 |
| CD164 | ?Deafness 66, 616969 | 603356 | 71 | 100 | 100 | 93 |
| CDC14A | Deafness 32, with or without immotile sperm, 608653 | 603504 | 79 | 100 | 100 | 94 |
| CDH23 | Deafness 12, 601386 {Pituitary adenoma 5, multiple types}, 617540 Usher syndrome, type 1D, 601067 Usher syndrome, type 1D/F digenic, 601067 | 605516 | 135 | 100 | 100 | 100 |
| CEACAM16 | Deafness 4B, 614614 Deafness 113, 618410 | 614591 | 94 | 100 | 100 | 100 |
| CEP78 | Cone-rod dystrophy and hearing loss, 617236 | 617110 | 69 | 100 | 100 | 94 |
| CIB2 | Deafness 48, 609439 Usher syndrome, type II, 614869 | 605564 | 195 | 100 | 100 | 100 |
| CLDN14 | Deafness 29, 614035 | 605608 | 66 | 100 | 98 | 91 |
| CLIC5 | ?Deafness 103, 616042 | 607293 | 76 | 100 | 100 | 100 |
| CLPP | Perrault syndrome 3, 614129 | 601119 | 125 | 100 | 100 | 100 |
| CLRN1 | Retinitis pigmentosa 61, 614180 Usher syndrome, type 3A, 276902 | 606397 | 83 | 100 | 100 | 97 |
| COA8 | Mitochondrial complex IV deficiency, 220110 | 616003 | No coverage data | | | |
| COCH | Deafness 9, 601369 ?Deafness 110, 618094 | 603196 | 71 | 98 | 95 | 93 |
| COL11A1 | ?Deafness 37, 618533 Fibrochondrogenesis 1, 228520 {Lumbar disc herniation, susceptibility to}, 603932 Marshall syndrome, 154780 Stickler syndrome, type II, 604841 | 120280 | 62 | 100 | 99 | 93 |
| COL11A2 | Deafness 13, 601868 Deafness 53, 609706 Fibrochondrogenesis 2, 614524 Otospondylomegapiphyseal dysplasia, 184840 Otospondylomegapiphyseal dysplasia, 215150 | 120290 | 110 | 100 | 100 | 100 |

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|---------------------------|--|--------------|--------------|----------------|----------------|----------------|
| COL2A1 | Achondrogenesis, type II or hypochondrogenesis, 200610 Avascular necrosis of the femoral head, 608805 Czech dysplasia, 609162 Epiphyseal dysplasia, multiple, with myopia and deafness, 132450 Kniest dysplasia, 156550 Legg-Calve-Perthes disease, 150600 Osteoarthritis with mild chondrodysplasia, 604864 Platyspondylic skeletal dysplasia, Torrance type, 151210 SED congenita, 183900 SMED Strudwick type, 184250 Spondyloepiphyseal dysplasia, Stanescu type, 616583 Spondyloperipheral dysplasia, 271700 Stickler syndrome, type I, nonsyndromic ocular, 609508 Stickler syndrome, type I, 108300 Vitreoretinopathy with phalangeal epiphyseal dysplasia | 120140 | 97 | 100 | 100 | 99 |
| COL4A3 | Alport syndrome 2, 203780 Alport syndrome 3, 104200 Hematuria, benign familial, 141200 | 120070 | 63 | 100 | 97 | 90 |
| COL4A4 | Alport syndrome 2, 203780 Hematuria, familial benign, 141200 | 120131 | 70 | 100 | 100 | 96 |
| COL4A5 | Alport syndrome 1, 301050 | 303630 | 42 | 100 | 88 | 67 |
| COL4A6 | ?Deafness 6, 300914 | 303631 | 62 | 100 | 98 | 90 |
| COL9A1 | ?Epiphyseal dysplasia, multiple, 6, 614135 Stickler syndrome, type IV, 614134 | 120210 | 65 | 100 | 99 | 91 |
| COL9A2 | Epiphyseal dysplasia, multiple, 2, 600204 ?Stickler syndrome, type V, 614284 | 120260 | 100 | 100 | 99 | 96 |
| CRYM | Deafness 40, 616357 | 123740 | 76 | 100 | 100 | 99 |
| DCDC2 | ?Deafness 66, 610212 Nephronophthisis 19, 616217 Sclerosing cholangitis, neonatal, 617394 | 605755 | 96 | 100 | 100 | 97 |
| DIABLO | Deafness 64, 614152 | 605219 | 105 | 100 | 100 | 99 |
| DIAPH1 | Deafness 1, 124900 Seizures, cortical blindness, microcephaly syndrome, 616632 | 602121 | 84 | 100 | 99 | 93 |
| DIAPH3 | Auditory neuropathy, 1, 609129 | 614567 | 61 | 100 | 99 | 93 |
| DMXL2 | ?Deafness 71, 617605 ?Polyendocrine-polyneuropathy syndrome, 616113 | 612186 | 67 | 100 | 99 | 93 |

| HGNC approved gene symbol | Phenotype description including OMIM phenotype ID(s) | OMIM gene ID | median depth | % covered >10x | % covered >20x | % covered >30x |
|---------------------------|---|--------------|--------------|----------------|----------------|----------------|
| DSPP | Deafness 39, with dentinogenesis, 605594 Dentin dysplasia, type II, 125420 Dentinogenesis imperfecta, Shields type II, 125490 Dentinogenesis imperfecta, Shields type III, 125500 | 125485 | 43 | 95 | 70 | 55 |
| EDN3 | Central hypoventilation syndrome, congenital, 209880 {Hirschsprung disease, susceptibility to, 4}, 613712 Waardenburg syndrome, type 4B, 613265 | 131242 | 90 | 100 | 100 | 98 |
| EDNRB | ABCD syndrome, 600501 {Hirschsprung disease, susceptibility to, 2}, 600155 Waardenburg syndrome, type 4A, 277580 | 131244 | 96 | 100 | 100 | 100 |
| ELMOD3 | ?Deafness 88, 615429 | 615427 | 96 | 100 | 100 | 97 |
| EPS8 | ?Deafness 102, 615974 | 600206 | 68 | 100 | 100 | 94 |
| EPS8L2 | Deafness autosomal recessive 106, 617637 | 614988 | 132 | 100 | 97 | 95 |
| ERAL1 | Perrault syndrome 6, 617565 | 607435 | 132 | 100 | 100 | 100 |
| ESPN | Deafness 36, 609006 Deafness, neurosensory, without vestibular involvement, 609006 | 606351 | 101 | 97 | 88 | 81 |
| ESRP1 | ?Deafness 109, 618013 | 612959 | 67 | 100 | 98 | 93 |
| ESRRB | Deafness 35, 608565 | 602167 | 142 | 100 | 100 | 100 |
| EXOSC2 | Short stature, hearing loss, retinitis pigmentosa, and distinctive facies, 617763 | 602238 | 69 | 100 | 100 | 97 |
| EYA1 | Anterior segment anomalies with or without cataract, 602588 Branchiootic syndrome 1, 602588 Branchiootorenal syndrome 1, with or without cataracts, 113650 ?Otofaciocervical syndrome, 166780 | 601653 | 86 | 100 | 100 | 98 |
| EYA4 | ?Cardiomyopathy, dilated, 1J, 605362 Deafness 10, 601316 | 603550 | 62 | 100 | 100 | 94 |
| FGF3 | Deafness, congenital with inner ear agenesis, microtia, and microdontia, 610706 | 164950 | 122 | 100 | 100 | 100 |
| FOXF2 | No OMIM phenotype | 603250 | 109 | 95 | 91 | 89 |
| FOXI1 | Enlarged vestibular aqueduct, 600791 | 601093 | 161 | 100 | 100 | 100 |
| GAB1 | ?Deafness 26, 605428 | 604439 | 72 | 100 | 99 | 97 |
| GATA3 | Hypoparathyroidism, sensorineural deafness, and renal dysplasia, 146255 | 131320 | 203 | 100 | 100 | 100 |
| GIPC3 | Deafness 15, 601869 | 608792 | 115 | 100 | 95 | 93 |
| GJB2 | Bart-Pumphrey syndrome, 149200 Deafness 3A, 601544 Deafness 1A, 220290 Hystrix-like ichthyosis with deafness, 602540 Keratitis-ichthyosis-deafness syndrome, 148210 Keratoderma, palmoplantar, with deafness, 148350 Vohwinkel syndrome, 124500 | 121011 | 150 | 100 | 100 | 100 |

| HGNC approved gene symbol | Phenotype description including OMIM phenotype ID(s) | OMIM gene ID | median depth | % covered >10x | % covered >20x | % covered >30x |
|---------------------------|--|--------------|--------------|----------------|----------------|----------------|
| GJB3 | Deafness 2B, 612644 Deafness, with peripheral neuropathy Deafness Deafness, digenic, GJB2/GJB3, 220290 Erythrokeratoderma variabilis et progressiva 1, 133200 | 603324 | 182 | 100 | 100 | 100 |
| GJB6 | Deafness 3B, 612643 Deafness 1B, 612645 Deafness, digenic GJB2/GJB6, 220290 Ectodermal dysplasia 2, Clouston type, 129500 | 604418 | 75 | 100 | 100 | 99 |
| GPSM2 | Chudley-McCullough syndrome, 604213 | 609245 | 93 | 100 | 100 | 96 |
| GRAP | Deafness 114, 618456 | 604330 | 52 | 68 | 68 | 65 |
| GREB1L | Renal hypodysplasia/aplasia 3, 617805 | 617782 | 72 | 100 | 98 | 93 |
| GRHL2 | Corneal dystrophy, posterior polymorphous, 4, 618031 Deafness 28, 608641 Ectodermal dysplasia/short stature syndrome, 616029 | 608576 | 81 | 100 | 100 | 99 |
| GRXCR1 | Deafness 25, 613285 | 613283 | 100 | 100 | 100 | 99 |
| GRXCR2 | ?Deafness 101, 615837 | 615762 | 109 | 100 | 100 | 100 |
| GSDME | Deafness 5, 600994 | 608798 | 94 | 100 | 100 | 100 |
| HARS1 | Charcot-Marie-Tooth disease, axonal, type 2W, 616625 Usher syndrome type 3B, 614504 | 142810 | 104 | 100 | 100 | 99 |
| HARS2 | ?Perrault syndrome 2, 614926 | 600783 | 76 | 100 | 100 | 99 |
| HGF | Deafness 39, 608265 | 142409 | 73 | 100 | 100 | 96 |
| HOMER2 | ?Deafness 68, 616707 | 604799 | 121 | 99 | 98 | 98 |
| HSD17B4 | D-bifunctional protein deficiency, 261515 Perrault syndrome 1, 233400 | 601860 | 62 | 100 | 98 | 93 |
| IFNLR1 | No OMIM phenotype | 607404 | 92 | 100 | 97 | 95 |
| ILDR1 | Deafness 42, 609646 | 609739 | 95 | 100 | 100 | 97 |
| KARS1 | ?Charcot-Marie-Tooth disease, recessive intermediate, B, 613641 Deafness 89, 613916 | 601421 | 126 | 100 | 100 | 97 |
| KCNE1 | Jervell and Lange-Nielsen syndrome 2, 612347 Long QT syndrome 5, 613695 | 176261 | 252 | 100 | 100 | 100 |
| KCNJ10 | Enlarged vestibular aqueduct, digenic, 600791 SESAME syndrome, 612780 | 602208 | 171 | 100 | 100 | 100 |
| KCNQ1 | Atrial fibrillation, familial, 3, 607554 Jervell and Lange-Nielsen syndrome, 220400 Long QT syndrome 1, 192500 {Long QT syndrome 1, acquired, susceptibility to}, 192500 Short QT syndrome 2, 609621 | 607542 | 160 | 97 | 94 | 93 |
| KCNQ4 | Deafness 2A, 600101 | 603537 | 147 | 98 | 95 | 91 |

| HGNC approved gene symbol | Phenotype description including OMIM phenotype ID(s) | OMIM gene ID | median depth | % covered >10x | % covered >20x | % covered >30x |
|---------------------------|--|--------------|------------------|----------------|----------------|----------------|
| KITLG | Deafness 69, unilateral or asymmetric, 616697 Hyperpigmentation with or without hypopigmentation, 145250 [Skin/hair/eye pigmentation 7, blond/brown hair], 611664 | 184745 | 78 | 100 | 100 | 97 |
| LARS2 | ?Hydrops, lactic acidosis, and sideroblastic anemia, 617021 Perrault syndrome 4, 615300 | 604544 | 79 | 100 | 100 | 99 |
| LHFPL5 | Deafness 67, 610265 | 609427 | 175 | 100 | 100 | 100 |
| LMX1A | No OMIM phenotype | 600298 | 81 | 100 | 100 | 99 |
| LOXHD1 | Deafness 77, 613079 | 613072 | 118 | 100 | 100 | 99 |
| LRTOMT | Deafness 63, 611451 | 612414 | 111 | 100 | 100 | 100 |
| MARVELD2 | Deafness 49, 610153 | 610572 | 92 | 100 | 100 | 97 |
| MCM2 | ?Deafness 70, 616968 | 116945 | 122 | 100 | 100 | 100 |
| MET | ?Deafness 97, 616705 Hepatocellular carcinoma, childhood type, somatic, 114550 {Osteofibrous dysplasia, susceptibility to}, 607278 Renal cell carcinoma, papillary, 1, familial and somatic, 605074 | 164860 | 64 | 100 | 100 | 96 |
| MIR96 | Deafness 50, 613074 | 611606 | No coverage data | | | |
| MITF | COMMAD syndrome, 617306 {Melanoma, cutaneous malignant, susceptibility to, 8}, 614456 Tietz albinism-deafness syndrome, 103500 Waardenburg syndrome, type 2A, 193510 Waardenburg syndrome/ocular albinism, digenic, 103470 | 156845 | 87 | 100 | 100 | 99 |
| MPZL2 | Deafness 111, 618145 | 604873 | 64 | 100 | 100 | 98 |
| MSRB3 | Deafness 74, 613718 | 613719 | 72 | 100 | 100 | 93 |
| MYH14 | Deafness 4A, 600652 ?Peripheral neuropathy, myopathy, hoarseness, and hearing loss, 614369 | 608568 | 114 | 100 | 99 | 97 |
| MYH9 | Deafness 17, 603622 Macrothrombocytopenia and granulocyte inclusions with or without nephritis or sensorineural hearing loss, 155100 | 160775 | 125 | 100 | 100 | 99 |
| MYO15A | Deafness 3, 600316 | 602666 | 134 | 100 | 99 | 98 |
| MYO3A | Deafness 30, 607101 | 606808 | 73 | 100 | 99 | 95 |
| MYO6 | Deafness 22, 606346 Deafness 22, with hypertrophic cardiomyopathy, 606346 Deafness 37, 607821 | 600970 | 82 | 100 | 99 | 94 |
| MYO7A | Deafness 11, 601317 Deafness 2, 600060 Usher syndrome, type 1B, 276900 | 276903 | 109 | 100 | 100 | 99 |
| NARS2 | Combined oxidative phosphorylation deficiency 24, 616239 ?Deafness 94, 618434 | 612803 | 55 | 99 | 97 | 92 |

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|---------------------------|--|--------------|--------------|----------------|----------------|----------------|
| NLRP3 | CINCA syndrome, 607115 Deafness 34, with or without inflammation, 617772 Familial cold inflammatory syndrome 1, 120100 Keratoendothelitis fugax hereditaria, 148200 Muckle-Wells syndrome, 191900 | 606416 | 125 | 100 | 100 | 100 |
| OPA1 | Behr syndrome, 210000 {Glaucoma, normal tension, susceptibility to}, 606657 ?Mitochondrial DNA depletion syndrome 14 (encephalocardiomyopathic type), 616896 Optic atrophy 1, 165500 Optic atrophy plus syndrome, 125250 | 605290 | 59 | 100 | 98 | 89 |
| OSBPL2 | Deafness 67, 616340 | 606731 | 102 | 100 | 100 | 100 |
| OTOA | Deafness 22, 607039 | 607038 | 78 | 77 | 74 | 72 |
| OTOF | Auditory neuropathy, 1, 601071 Deafness 9, 601071 | 603681 | 113 | 100 | 100 | 100 |
| OTOG | Deafness 18B, 614945 | 604487 | 121 | 100 | 100 | 99 |
| OTOGL | Deafness 84B, 614944 | 614925 | 76 | 100 | 100 | 95 |
| P2RX2 | Deafness 41, 608224 | 600844 | 136 | 100 | 100 | 99 |
| PAX3 | Craniofacial-deafness-hand syndrome, 122880 Rhabdomyosarcoma 2, alveolar, 268220 Waardenburg syndrome, type 1, 193500 Waardenburg syndrome, type 3, 148820 | 606597 | 104 | 100 | 100 | 100 |
| PCDH15 | Deafness 23, 609533 Usher syndrome, type 1D/F digenic, 601067 Usher syndrome, type 1F, 602083 | 605514 | 72 | 100 | 100 | 95 |
| PDE1C | ?Deafness 74, 618140 | 602987 | 63 | 100 | 97 | 90 |
| PDZD7 | Deafness 57, 618003 {Retinal disease in Usher syndrome type IIA, modifier of}, 276901 Usher syndrome, type IIC, GPR98/PDZD7 digenic, 605472 | 612971 | 84 | 100 | 100 | 99 |
| PET100 | Mitochondrial complex IV deficiency, 220110 | 614770 | 85 | 92 | 66 | 66 |
| PEX1 | Heimler syndrome 1, 234580 Peroxisome biogenesis disorder 1A (Zellweger), 214100 Peroxisome biogenesis disorder 1B (NALD/IRD), 601539 | 602136 | 62 | 100 | 99 | 94 |
| PEX26 | Peroxisome biogenesis disorder 7A (Zellweger), 614872 Peroxisome biogenesis disorder 7B, 614873 | 608666 | 118 | 100 | 100 | 100 |
| PEX6 | Heimler syndrome 2, 616617 Peroxisome biogenesis disorder 4A (Zellweger), 614862 Peroxisome biogenesis disorder 4B, 614863 | 601498 | 107 | 100 | 99 | 95 |
| PJKK | Deafness 59, 610220 | 610219 | 93 | 100 | 100 | 99 |

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|---------------------------|---|--------------|--------------|----------------|----------------|----------------|
| PNPT1 | Combined oxidative phosphorylation deficiency 13, 614932 Deafness 70, 614934 | 610316 | 57 | 100 | 97 | 83 |
| POU3F4 | Deafness 2, 304400 | 300039 | 120 | 100 | 100 | 100 |
| POU4F3 | Deafness 15, 602459 | 602460 | 247 | 100 | 100 | 100 |
| PIIP5K2 | Deafness 100, 618422 | 611648 | 48 | 99 | 93 | 80 |
| PRKCB | No OMIM phenotype | 176970 | 73 | 100 | 100 | 99 |
| PRPS1 | Arts syndrome, 301835 Charcot-Marie-Tooth disease recessive, 5, 311070 Deafness 1, 304500 Gout, PRPS-related, 300661 Phosphoribosylpyrophosphate synthetase superactivity, 300661 | 311850 | 50 | 100 | 99 | 90 |
| PTPRQ | Deafness 73, 617663 Deafness 84A, 613391 | 603317 | 80 | 100 | 99 | 95 |
| RAI1 | Smith-Magenis syndrome, 182290 | 607642 | 187 | 100 | 100 | 100 |
| RDX | Deafness 24, 611022 | 179410 | 54 | 100 | 93 | 76 |
| RIPOR2 | ?Deafness 104, 616515 | 611410 | 77 | 100 | 100 | 97 |
| ROR1 | ?Deafness 108, 617654 | 602336 | 87 | 97 | 96 | 96 |
| S1PR2 | Deafness 68, 610419 | 605111 | 153 | 100 | 100 | 100 |
| SERPINB6 | ?Deafness 91, 613453 | 173321 | 85 | 100 | 100 | 98 |
| SIX1 | Branchiootic syndrome 3, 608389 Deafness 23, 605192 | 601205 | 119 | 100 | 100 | 100 |
| SIX5 | Branchiootorenal syndrome 2, 610896 | 600963 | 75 | 100 | 95 | 85 |
| SLC17A8 | Deafness 25, 605583 | 607557 | 66 | 100 | 100 | 99 |
| SLC22A4 | {Rheumatoid arthritis, susceptibility to}, 180300 | 604190 | 103 | 100 | 100 | 96 |
| SLC26A4 | Deafness 4, with enlarged vestibular aqueduct, 600791 Pendred syndrome, 274600 | 605646 | 68 | 100 | 100 | 94 |
| SLC26A5 | ?Deafness 61, 613865 | 604943 | 61 | 100 | 100 | 96 |
| SLC29A3 | Histiocytosis-lymphadenopathy plus syndrome, 602782 | 612373 | 152 | 100 | 99 | 99 |
| SLC33A1 | Congenital cataracts, hearing loss, and neurodegeneration, 614482 Spastic paraplegia 42, 612539 | 603690 | 68 | 100 | 98 | 86 |
| SLC44A4 | ?Deafness 72, 617606 | 606107 | 94 | 100 | 100 | 100 |
| SLC52A2 | Brown-Vialetto-Van Laere syndrome 2, 614707 | 607882 | 173 | 100 | 100 | 100 |
| SLITRK6 | Deafness and myopia, 221200 | 609681 | 75 | 100 | 100 | 99 |
| SMPX | Deafness 4, 300066 | 300226 | 51 | 100 | 100 | 83 |
| SNAI2 | Piebaldism, 172800 Waardenburg syndrome, type 2D, 608890 | 602150 | 66 | 100 | 100 | 100 |
| SOX10 | PCWH syndrome, 609136 Waardenburg syndrome, type 2E, with or without neurologic involvement, 611584 Waardenburg syndrome, type 4C, 613266 | 602229 | 72 | 100 | 97 | 89 |

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|---------------------------|---|--------------|--------------|----------------|----------------|----------------|
| SPATA5 | Epilepsy, hearing loss, and mental retardation syndrome, 616577 | 613940 | 68 | 100 | 100 | 97 |
| STRC | Deafness 16, 603720 | 606440 | 66 | 53 | 48 | 46 |
| SYNE4 | Deafness 76, 615540 | 615535 | 107 | 100 | 100 | 100 |
| TBC1D24 | DOORS syndrome, 220500 Deafness 86, 614617 Deafness 65, 616044 Epilepsy, rolandic, with proxysmal exercise-induce dystonia and writer's cramp, 608105 Epileptic encephalopathy, early infantile, 16, 615338 Myoclonic epilepsy, infantile, familial, 605021 | 613577 | 158 | 100 | 100 | 100 |
| TECTA | Deafness 8/12, 601543 Deafness 21, 603629 | 602574 | 123 | 100 | 100 | 98 |
| TIMM8A | Mohr-Tranebjaerg syndrome, 304700 | 300356 | 138 | 100 | 100 | 100 |
| TJP2 | Cholestasis, progressive familial intrahepatic 4, 615878 Hypercholanemia, familial, 607748 | 607709 | 83 | 100 | 100 | 99 |
| TMC1 | Deafness 36, 606705 Deafness 7, 600974 | 606706 | 61 | 100 | 100 | 94 |
| TMEM132E | ?Deafness 99, 618481 | 616178 | 117 | 100 | 99 | 96 |
| TMIE | Deafness 6, 600971 | 607237 | 85 | 100 | 100 | 99 |
| TMPRSS3 | Deafness 8/10, 601072 | 605511 | 74 | 100 | 100 | 99 |
| TMTC2 | No OMIM phenotype | 615856 | 79 | 100 | 100 | 98 |
| TNC | Deafness 56, 615629 | 187380 | 110 | 100 | 100 | 98 |
| TPRN | Deafness 79, 613307 | 613354 | 94 | 92 | 86 | 79 |
| TRIOBP | Deafness 28, 609823 | 609761 | 182 | 100 | 99 | 97 |
| TSPEAR | ?Deafness 98, 614861 Ectodermal dysplasia 14, hair/tooth type with or without hypohidrosis, 618180 | 612920 | 105 | 100 | 100 | 100 |
| TWNK | Mitochondrial DNA depletion syndrome 7 (hepatocerebral type), 271245 Perrault syndrome 5, 616138 Progressive external ophthalmoplegia with mitochondrial DNA deletions 3, 609286 | 606075 | 145 | 100 | 100 | 100 |
| TYR | Albinism, oculocutaneous, type IA, 203100 Albinism, oculocutaneous, type IB, 606952 {Melanoma, cutaneous malignant, susceptibility to, 8}, 601800 [Skin/hair/eye pigmentation 3, blue/green eyes], 601800 [Skin/hair/eye pigmentation 3, light/dark/freckling skin], 601800 Waardenburg syndrome/albinism, digenic, 103470 | 606933 | 83 | 100 | 100 | 98 |
| USH1C | Deafness 18A, 602092 Usher syndrome, type 1C, 276904 | 605242 | 80 | 100 | 98 | 92 |
| USH1G | Usher syndrome, type 1G, 606943 | 607696 | 180 | 100 | 100 | 100 |
| USH2A | Retinitis pigmentosa 39, 613809 Usher syndrome, type 2A, 276901 | 608400 | 82 | 100 | 100 | 98 |

| HGNC approved gene symbol | Phenotype description including OMIM phenotype ID(s) | OMIM gene ID | median depth | % covered >10x | % covered >20x | % covered >30x |
|---------------------------|--|--------------|--------------|----------------|----------------|----------------|
| WBP2 | Deafness 107, 617639 | 606962 | 60 | 100 | 96 | 89 |
| WFS1 | ?Cataract 41, 116400 Deafness 6/14/38, 600965 {Diabetes mellitus, noninsulin-dependent, association with}, 125853 Wolfram syndrome 1, 222300 Wolfram-like syndrome, 614296 | 606201 | 196 | 100 | 100 | 100 |
| WHRN | Deafness 31, 607084 Usher syndrome, type 2D, 611383 | 607928 | 119 | 100 | 100 | 100 |
| YAP1 | Coloboma, ocular, with or without hearing impairment, cleft lip/palate, and/or mental retardation, 120433 | 606608 | 73 | 100 | 99 | 94 |

- Gene symbols according HGNC
- OMIM release used: 8-9-2019
- "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 100 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 , 20x and 30 x describes the percentage of a gene's coding sequence (± 10 bp flanking introns) that is covered at least 10x, 20x or 30x