

TABLE S1. Rare damaging variants identified in 15 Filipino children with hearing loss requiring cochlear implants

Bold=probable cause of hearing loss; Italics=probable cause of other phenotypes

ID	Age	Sex	AdditionalFeatures	FamilyHistory	Genotype	Variant	rsID	gnomAD_EAS(hg38)	gnomAD_highestMAF(hg38)	1000Genomes	GMEVariome	GenomeAsia100K_SEA	scaledCADD(hg38)	Damaging_dbNSFP41a	IMPC_gene/MouseHearingPhenotype	Literature on Known Phenotypes	Missed in previous analyses with hg19-refSeq/gnomAD or dbNSFP33a				
1	3.95	M	EVA, L; brain cysts		Heterozygous	DSPP:NM_014208:exon4:c.G730A:p.G244R	rs1044690454	0.001	24.3	FA/mLr/mSVM/MT/PP2/SI	NoEarPhenotype	PMID11175790: multiple Chinese families with hearing loss, one with EVA	No			
					Heterozygous	ANLN:NM_018685:exon11:c.G1921C:p.E641Q	rs748210737	.	0.0015	EAS:0.0015	20.9	MA/PP2	NoKoMouse	PMCG93648: heterozygous missense variant co-segregating with branchio-oto-renal syndrome in Chinese family	gnomAD MAF>0.001	
					Potential Compound Heterozygous or Haplotype	ZNF462:NM_021224:[exon3:c.G3916A:p.G1306S]; [exon3:c.G4129A:p.E1377K]	NA; rs768977077	NA; 0	NA; NFE:0.0001	28.9; 24.8	LRT/MT/PP2/SI; LRT/MA/MT/PP2/SI	Prewaning lethality	PMCG935050: hearing loss or inner ear malformation (not specified) in 50% of Weiss-Kruszka syndrome patients with heterozygous loss-of-function variants in exon 3 of ZNF462	No	
					Heterozygous	CEP290:NM_025114:exon39:c.C5284T:p.R1762C	rs373307908	.	0.0004	OTH:0.00096	0.013	29.6	LRT/MT/PP2/PR/SI	NoEarPhenotype	PMCG2597962: heterozygous missense mutation in patient with Joubert syndrome including cystic kidney disease and hearing loss	No
3	2.83	M	Malformed cochlea with incomplete cochlear turns, B; EVA, L; global developmental delay		Heterozygous	LMX1A:NM_177398:exon5:c.G606C:p.L202F	24.8	FA/LRT/mLr/mSVM/MT/PP2/PR/SI	PMCG511360: KO-hom but not het mice have cochleovestibular defects	PMCG5973959: heterozygous missense variants co-segregate with nonsyndromic hearing loss in two Dutch families; PMCG6094940: homozygous missense variant co-segregates with nonsyndromic hearing loss in Pakistani family; PMID32840933: functional heterozygous missense variant in Korean proband	No			
					Potential Compound Heterozygous or Haplotype	USH2A:NM_206933:[exon47:c.G9286A:p.V3096M]; [exon46:c.C9109T:p.R3037C]	rs147267500; rs770418427	0; 0.0002	AFR:0.0001; EAS:0.0002	0.007; 0.007	18.3; 23.8	MA/MT/SI/PP2; LRT/MA/MT/PR/SI	PMCI1838616: Ush2a-KO mice have progressive photoreceptor degeneration and moderate nonprogressive hearing loss	MIM276901: AR Usher syndrome type 2A	Variant V3096M was ruled out in previous CLHNS screening due to MAF>0.01 (Truong et al. 2019)	
					Heterozygous	ZFH4:NM_024721:exon10:c.G5119A:p.A1707T	rs764848741	0	FIN:0.0002	25.8	MT/PR/SI	NotPhenotyped	PMCG3155189: 8q21.11 deletion including ZFH4 in 8 patients with sensorineural hearing loss; developmental delay may be present	No
					Heterozygous	NRP1:NM_003873:exon10:c.G1655A:p.R552Q	rs757990959	0	OTH:0.0005	29.2	FA/LRT/MA/mLr/mSVM/MT/PP2/SI	Het-KO mice with abnormal ABR	PMCG5695633: conditional-KO mice with disorganized outer spiral bundles and enlarged microvessels of stria vascularis, progressive hearing loss	No
					Heterozygous	COL2A1:NM_001844:exon50:c.G3569A:p.R1190H	rs748549541	27.3	FA/LRT/mLr/mSVM/MT/PP2/PR/SI	PMID10100048: Tg-mice with small misshapen otic capsule	MIM120140: multiple AD syndromes with hearing loss, including Stickler syndrome	No
					Hemizygous	ARHGAP4:NM_001164741:exon18:c.G2175C:p.Q725H	rs200637748	0.002	EAS:0.002	EAS:0.001	21.4	MT/PR/SI	NoEarPhenotype	PMID26707211: missense variant in child with intellectual disability; PMID2965914: deletion including ARHGAP4 in twins with intellectual disability	No; gnomAD MAF>0.001
5	3.84	F	HIB with dehiscence, L; neonatal infection; pervasive developmental delay		Potential Compound Heterozygous or Haplotype	CCDC186:NM_153249:[exon3:c.A461C:p.K154T]; [exon3:c.G391A:p.A131T]	rs994643573; rs147799223	0.0004; 0.0004	EAS:0.0004; 0.0004	NA; NA	22.4; 15.4	LRT/MT/PP2/PR/SI; SI	Prewaning lethality	PMCG7818090, PMCG502059: homozygous variants in patients with global developmental delay	Yes		
					Heterozygous	DMXL2:NM_015263:exon3:c.T257C:p.L86S	rs761692429	0.0002	OTH:0.0005	24.1	LRT/MT/SI/PP2	Hom-KO Prewaning lethality, het-KO decreased bone mineral content	PMID27657680, PMID33715530: heterozygous missense variants cosegregating with AD hearing loss in Chinese and Cameroonian families	No
					Heterozygous	ZFR2:NM_015174:exon19:c.C2653T:p.R885X	rs375337964	0.001	45	MT	NoKoMouse	PMID23610052: 19p13.3 deletions including ZFR2 in patients with intellectual disability or developmental delay, 40% with hearing loss	Yes
					Potential Compound Heterozygous or Haplotype	MCM3AP:NM_003906:[exon25:c.G5383A:p.A1795T]; [exon1:c.G998T:p.R333L]	rs17183290; rs17182552	0.004; 0.002	EAS:0.004; EAS:0.002	EAS:0.003; EAS:0.002	0.01; 0.01	26.7; 22.5	LRT/MA/MT/PP2/SI; MA	NoKoMouse	PMID32319184: AR Charcot-Marie-Tooth disease, including one Finnish patient with compound heterozygous variants and intellectual disability and hearing loss; PMCG477159: MCM3AP is required for inhibition of cellular DNA synthesis in viral infection	Yes; 1KG MAF>0.001	
6	10.81	M	PSCD + HIB, B; EVA, R; pneumonia, sinusitis, and progressive hearing loss	Hearing loss, paternal side	Heterozygous	DNAH14:NM_001367479:exon5:c.G8842T:p.A2948S	rs541854543	0.0004	EAS:0.0004	0.003	23.9	PP2	NoKoMouse	PMID30125339: heterozygous deletion in patient with hydrocephalus and cognitive dysfunction; PMID3357779: compound heterozygous DNAH14 variants in Chinese proband with bronchiectasis, chronic sinusitis and otitis media	No		
					Heterozygous	ATAD2B:NM_017552:exon4:c.A475G:p.I159V	rs912934573	20.3	FA/mLr/MT	Hom-KO mice with abnormal ABR (het not tested)	No	
					Heterozygous	ECT2:NM_001258315:exon15:c.G1579A:p.D527N	rs182472799	0.0002	SAS:0.001	EAS:0.001	0.01	21.2	LRT/MT	Het-KO mice with abnormal ABR	No; gnomAD MAF>0.001	
					Heterozygous	TCOF1:NM_001135243:exon24:c.C3823T:p.R1275W	rs768512539	0	ASJ:0.0006	0 (SouthAsia=0.0007)	24.9	PP2/PR/SI	PMCG489999: het-mutant mice with severe craniofacial defects, middle ear cavitation and hearing loss	MIM154500: AD Treacher-Collins syndrome often with conductive hearing loss and cleft palate	Previously ruled out due to lack of craniofacial features in patient carrying the variant (Truong et al. 2019)	
					Heterozygous	PTPRQ:NM_001145026:exon40:c.T6179C:p.V2060A	rs375150180	0.00097	EAS:0.00097	EAS:0.003	0.017	27.8	MT/SI	PMCG740823: rapid postnatal deterioration in cochlear hair-bundle structure	PMCG599672: AD nonsyndromic hearing loss	Yes
					Heterozygous	LRRK1:NM_024652:exon2:c.C16T:p.Q6X	rs772698984	33	MT	PMCG4472125: Lrrk1-KO mice have severe osteopetrosis	PMCG5348726: LRRK1 deficiency in humans and mice lead to increased bone mineral density, reduced body length and bone marrow, and sclerosis of vertebral endplates and pelvis; PMCG5769692: reports of AR osteosclerotic metaphyseal dysplasia due to homozygous LRRK1 variants	No
					Heterozygous	CPS1:NM_001875:exon27:c.3337-1G>T	34	MT	Prewaning lethality	MIM237300: AR carbamoyl phosphate synthetase I deficiency; PMID26440671: Includes hypertonia at presentation but may also have hypotonia and mild motor delay later
7	8	F	HIB, L; OM, L; mild motor delay and hypotonia; history of urinary and upper respiratory tract infections	Older brother seen for giftedness; Down syndrome, maternal cousin; asthma, paternal side	Heterozygous	GMPPB:NM_013334:exon8:c.G1032C:p.E344D	23.9	MT/SI	NotPhenotyped	PMCG3710768, PMCG4843780: two patients with compound heterozygous GMPPB variants had muscle weakness, intellectual disability and sensorineural hearing loss	No		
					Heterozygous	TNXB:NM_019105:exon5:c.C2461T:p.R821X	rs749890642	0	NFE:0.00006	0 (Oceania=0.007)	36	LRT/MT	NoEarPhenotype	MIM600985: AR Ehlers-Danlos syndrome (EDS), AD vesicoureteral reflux; PMID27253074: conductive or sensorineural hearing loss common in EDS	No	
					Heterozygous	PLEC:NM_201380:exon20:c.2868+1G>A	rs368904034	0.001	34	MT	NotPhenotyped	MIM601282: AR/AD epidermolysis bullosa simplex, AR limb-girdle muscular dystrophy	No
					Heterozygous	ARHGAP21:NM_020824:exon7:c.G475T:p.D159V	rs1266442941	0.0002	EAS:0.0002	29	LRT/MT/PP2/PR/SI	Het-KO mice with abnormal ABR	PMCG406497: candidate gene for autism	No
					Potential Compound Heterozygous or Haplotype	PCDH15:NM_001354411:exon29:c.C3787T:p.P1263S]; [CDH23:NM_022124:exon27:c.G3262A:p.V1088M]	rs775954124; rs200632520	0.004; 0.002	EAS:0.004; EAS:0.002	NA; 0.003	24.9; 24.3	MA/MT/PP2/PR/SI; LRT/MA/mLr/mSVM/MT/PP2/SI	PMCG2858222: double-heterozygous mice with progressive hearing loss	PMCG2858222: compound heterozygous probands with congenital profound hearing loss	No; gnomAD MAF>0.001	

					Heterozygous	MYO7A:NM_000260:exon36:c.G4921A:p.E1641K	rs767975012	0.0002	EAS:0.0002			0.003	26.2	LRT/MT/PP2/PR	AbsentPinnaReflex/AbnormalBone Mineralization(Hom); PMID15389316: mice heterozygous for missense variant have abnormal stereocilia bundles and low-frequency hearing loss	MIM276903: DFNA11; DFNB2; Usher syndrome IB	No	
					Heterozygous	DNAJC17:NM_018163:exon11:c.A871G:p.I291V							18.96	MA/MT/SI	Het-KO mice with abnormal ABR	PMID26355662: homozygous variant in family with retinitis pigmentosa and hypogammaglobulinemia	No	
					Potential Compound Heterozygous or Haplotype	NEO1:NM_02499:[exon3:c.G586C:p.D196H]; [exon24:c.C3443T:p.S1148F]	NA; rs1048863959	NA; 0	NA; LAT:0.0007				22.5; 25.9	MT; LRT/MT/PP2/PR/SI	NoKoMouse	PMID31953991: included in critical 15q24 microdeletion associated with autism and variable hearing loss, recurrent infections	No	
					Potential Compound Heterozygous or Haplotype	PIEZO2:NM_022068:[exon38:c.G5888A:p.R1963Q]; [exon37:c.C557T:p.R1853C]; [exon9:c.A1159G:p.S387G]	rs1480686209; rs569343097; rs371514584	0; 0.0002; 0	AFR:0.00002; ASJ:0.0003; SAS:0.0002	NA; EUR:0.001; NA	NA; 0.001 (Oceania=0.03); 0.01 (Oceania=0.10)	25.6; 22.6; 24.6	MA/MT/PR/SI; SI; LRT	Prewaning lethality	PMC5097934: biallelic variants in probands with arthrogyposis, respiratory insufficiency, muscular atrophy, scoliosis and mild distal sensory loss	No		
8	3.03	M	SSCD, L		Heterozygous	COL11A1:NM_080629:exon58:c.A4364C:p.K1455T	rs769350133	0.0004	EAS:0.0004				28.6	FA/LRT/mLr/mSVM/MT/PP2/PR/SI	DecreasedStartleReflex	MIM120280: DFNA37; AD Marshall Syndrome; AD Stickler	No	
					Heterozygous	ZFH4:NM_024721:exon3:c.G3007A:p.A1003T	rs369411377	0.002	EAS:0.002			0.009	27.8	MT	NotPhenotyped	PMC3155189: 8q21.11 deletion including ZFH4 in 8 patients with sensorineural hearing loss	No; gnomAD MAF>0.001	
					Heterozygous	TECTA:NM_005422:exon10:c.C2967A:p.H989Q	rs200821009	0.003	EAS:0.003	EAS:0.003			0.001	20.4	FA/LRT/mLr/mSVM/MT/PP2/PR/SI	PMc2869304: mice heterozygous for missense mutation have non-functional outer hair cells	MIM602754: DFNA8/12; DFNB21	No; gnomAD MAF>0.001
					Heterozygous	CEP290:NM_025114:exon48:c.G6629A:p.R2210H	rs371833544	0.0002	ME:0.003				23.2	LRT/MA/MT/PP2/PR/SI	NoEarPhenotype	PMC2597962: heterozygous missense mutation in patient with Joubert syndrome including cystic kidney disease and hearing loss	No	
9	8.19	F	EVA, L		Heterozygous	IST1:NM_001270976:exon8:c.C737G:p.P246R	rs774343604	0.0002	EAS:0.0002				24	LRT/MT/PP2/PR/SI	Het-KO mice with abnormal ABR		No	
13	5.95	M	Global developmental delay	Hearing loss, 3rd-degree cousin	Potential Compound Heterozygous or Haplotype	GDAP2:NM_017686:[exon14:c.A1463G:p.Y488C]; [exon2:c.C44T:p.T151]	NA; rs1272520325	NA; 0	NA; AFR:0.00002				22.8; 19.8	MT/SI; LRT	AbsentPinnaReflex(Hom)	PMC7534050: AR adult-onset spinocerebellar ataxia	No	
					Heterozygous	SLC12A2:NM_001046:exon21:c.G2977T:p.E993X							60	MT	PMID10369265: Slc12a2-mice are deaf	MIM600840: DFNA78; AD Delpire-McNeill syndrome (global developmental delay and sensorineural hearing loss); AR Klippel syndrome	No	
					Homozygous	ARMC4:NM_018076:exon5:c.A647G:p.N216S	rs981525890	0.0002	EAS:0.0002		0 (Oceania=0.03)		13.99	SI	NoEarPhenotype	PMC3788828: AR primary ciliary dyskinesia, may include hearing loss due to primary ciliary dyskinesia	No	
18	2.77	M			Heterozygous	RERE:NM_012102:exon10:c.T910C:p.S304P						0.001	27.6	MT/PP2/SI	PMc3581587: heterozygous mice had decreased startle response and elevated ABR thresholds	PMc5903952: AD sensorineural hearing loss with eye, heart and renal defects	No	
					Heterozygous	KRIT1:NM_194456:exon5:c.C74T:p.S25F	rs779446044	0	OTH:0.0005			0.003	28.6	FA/LRT/MA/mLr/mSVM/MT/PP2/PR/SI	Het-/hom-KO have abnormal otic vesicle morphology	MIM604214: AD cavernous malformations	No	
					Potential Compound Heterozygous or Haplotype	CDH23:NM_022124:[exon11:c.G982A:p.A328T]; [exon20:c.G2236A:p.V746]	rs374545987; rs550384315	0.00096; 0.001	EAS:0.00096; EAS:0.001	AMR: 0.0014; AMR: 0.0014	0.02; 0.01 (Oceania=0.06)	28.4; 22.9	LRT/MT/PP2/SI; LRT/PP2/MT	Hom-KO mice with absent pinna reflex	MIM605516: DFNB12; AR Usher 1D	V746I was excluded due to CLHNS MAF=0.19 (Truong et al. 2019)	No	
					Heterozygous	MYO7A:NM_000260:exon36:c.G4921A:p.E1641K	rs767975012	0.0002	EAS:0.0002				0.003	26.2	LRT/MT/PP2/PR	AbsentPinnaReflex/AbnormalBone Mineralization(Hom); PMID15389316: mice heterozygous for missense variant have abnormal stereocilia bundles and low-frequency hearing loss	MIM276903: DFNA11; DFNB2; Usher syndrome IB	No
					Potential Compound Heterozygous or Haplotype	NFRKB:NM_006165:[exon22:c.C3191G:p.T1064S]; [NM_006165:exon20:c.C2408T:p.P803L]	rs559022552; rs557840039	0.003; 0.003	EAS:0.003; EAS:0.003	EAS:0.002; EAS:0.002			19.7; 24.4	LRT; LRT/MT/PP2/SI	NoKoMouse	PMC2648978: NFRKB included in submicroscopic deletion in 11q24-25 in patients with hearing loss and multiple anomalies	No	
19	5.66	F	Malformed cochlea, vestibules and semicircular canals, B; absent cochlear and inferior vestibular nerves, R		Heterozygous	PPOX:NM_000309:exon5:c.A394T:p.R132W	rs771307571	0.0002	EAS:0.0002				24.5	FA/mLr/mSVM/MT/PP2/PR/SI	Het-KO mice with abnormal ABR	MIM600923: AD porphyria variegata (1 case report of sudden hearing loss in a porphyria variegata patient)	No	
					Potential Compound Heterozygous or Haplotype	GOLGB1:NM_001256486:[exon14:c.G7472A:p.R2491Q]; [exon14:c.A7367C:p.E2456A]; [exon14:c.C7112G:p.S2371C]	rs767375937; rs751018606; rs919070773	0; 0.0002; NA	AFR:0.00002; EAS:0.0002; NA				24.8; 25.9; 26.2	LRT/MA/MT/PP2/SI; LRT/MA/MT/PP2/PR/SI; LRT/MA/MT/PP2/SI	PMID2185186: osteochondrodysplasia; PMc4958322: palatal shelf defects		No	
					Heterozygous	SLC36A1:NM_078483:exon7:c.C721T:p.Q241X							47	LRT/MT	KO-het abnormal craniofacial morphology		No	
					Heterozygous	TFEB:NM_001167827:exon2:c.C136G:p.H46D							22.6	MT/PR/SI	Het-KO mice with abnormal ABR; decreased BMD	PMCID: attenuates spiral ganglion neuron degeneration	No	
					Heterozygous	GDPD5:NM_030792:exon7:c.C404T:p.T135M	rs373413383	0	AFR:0.00002		0.01 (Oceania=0.07)		24.8	LRT/MA/MT/PP2	Hom-KO mice with abnormal ABR (het not tested)		No	
					Heterozygous	MYO18B:NM_032608:exon13:c.C255T:p.A852V							26.1	FA/LRT/mLr/mSVM/MA/MT/PP2/PR/SI	Het-KO mice with abnormal ABR	MIM616549: AR Klippel-Feil syndrome with nemaline myopathy and facial dysmorphism; PMc5885878: hearing loss common in KFS; PMID18722888: 60% of Klippel-Feil HL patients with ear anomalies including inner ear dysplasia and deformed internal acoustic canal	No	
20	14.59	F	Fluctuating hearing loss with steeply sloping audiogram prior to C1; turbinate hypertrophy, allergic rhinitis, nasopharyngeal nodule		Heterozygous	ARHGAP29:NM_004815:exon17:c.G1847A:p.R616H	rs374991521	0.001	OTH:0.001				25.3	FA/MA/mLr/mSVM/MT/PP2/PR/SI	Het-KO mice with abnormal ABR	PMC3501616: cleft lip and palate	No; gnomAD MAF>0.001	
					Heterozygous	CACNA1S:NM_000069:exon12:c.G1745C:p.G582A	rs377459546	0.0008	LAT:0.001			0.004	26	FA/LRT/MA/mLr/mSVM/MT/PP2/PR/SI	Het-KO mice with abnormal ABR	MIM170400: AD hypokalemic periodic paralysis	No; gnomAD MAF>0.001	
					Heterozygous	DSPP:NM_014208:exon5:c.G1577A:p.S526N	rs201485801	0.001	EAS:0.001	EAS:0.003		0.02	13	FA/mLr/mSVM/PP2/SI	NoEarPhenotype	PMID1175790: multiple Chinese families with hearing loss, one with EVA	No; gnomAD MAF>0.001	
					Heterozygous	CEP290:NM_025114:exon8:c.G503A:p.R168H	rs200063017	0.003	EAS:0.003		NEA:0.001	0.003	22.7	FA/SI	NoEarPhenotype	PMC2597962: heterozygous missense mutation in patient with Joubert syndrome including cystic kidney disease and hearing loss	No; gnomAD MAF>0.001	
					Heterozygous	BCL2L10:NM_020396:exon1:c.C414A:p.C138X	rs79743288	0.0006	EAS:0.0006				36	MT	NoKoMouse	PMc6817556: GWAS locus for age-related hearing loss	No	
					Homozygous	CLDN9:NM_020982:exon1:c.C75G:p.C25W	rs368045321	0.0004	OTH:0.0005			0.004	20.6	FA/LRT/MA/mLr/mSVM/MT/PP2/PR/SI	PMc2720454: AR deafness with loss of sensory cells	PMc6745279: AR nonsyndromic hearing loss	No	
					Heterozygous	ANKRD11:NM_013275:exon9:c.A3830G:p.E1277G							24.5	LRT/MA/MT/PP2/PR/SI	Het-KO mice with abnormal ABR	MIM148050: AD KBG syndrome, may include hearing loss in 1/3, also craniofacial dysmorphisms, susceptibility to nasopharyngeal carcinoma	No	
					Heterozygous	FLNA:NM_00110556:exon39:c.A6350G:p.N21175	rs375205247	0.002	EAS:0.002				20.2	FA/LRT/MT/PR	NoKoMouse	MIM311300: otopalatodigital syndrome inc. hearing loss	No; gnomAD MAF>0.001	
					Potential Compound Heterozygous or Haplotype	PATJ:NM_176877:[exon11:c.C1351G:p.R451G]; [exon34:c.4461+1G>A]	rs117405402; rs19981703	0.003; 0.003	EAS:0.003; EAS:0.003	EAS:0.003; EAS:0.004		0.007; 0.01	25.3; 32	LRT/MA/MT/PP2/PR/SI; MT	Hom-KO mice with abnormal startle reflex (late adult)		Yes; gnomAD MAF>0.001	
					Heterozygous	ARID1B:NM_001346813:exon20:c.G5648A:p.R1883H	rs758748419	0	FIN:0.00009		0 (SouthAsia=0.0007)		23.2	LRT/MA/MT/PP2/PR/SI	Prewaning lethality	MIM614556: AD Coffin-Siris syndrome	No	
					Heterozygous	MYO3A:NM_017433:exon32:c.C4483T:p.R1495X	rs371876274	0.0002	SAS:0.001	SAS:0.002			41	LRT/MT	Hom-KO mice with abnormal ABR (het not tested)	MIM606808: DFNB30	No; gnomAD MAF>0.001; previously ruled out due to lack of second allele explaining AR hearing loss	
					Heterozygous	CBLN3:NM_001039771:exon3:c.C550T:p.R184C	rs562291434	0.0002	EAS:0.0002				32	LRT/MT/PP2/PR/SI	Het-KO mice with abnormal ABR		No	

					Heterozygous	HOXB6:NM_018952:exon3:c.T405A:p.N135K	rs758723592							25.6	FA/LRT/MA/mLR/mSVM/MT/PP2/PR/SI	PMID1358998: gain-of-function mutant with early postnatal lethality, craniofacial/axial skeletal anomalies, cleft palate, microtia, skull bone defects etc		Yes	
					Heterozygous	GREB1L:NM_001142966:exon22:c.C3798G:p.S1266R	rs954005555	0.0006	EAS:0.0006					0.003	16.6	LRT/MA/MT/PR/SI	Prewaning lethality	PMC7349314: AD profound sensorineural hearing loss, variable genitourinary findings	No
					Heterozygous	TRMT1:NM_017722:exon4:c.C619T:p.Q207X									36	MT	Prewaning lethality; PMID33499731: subcellular relocation of RNA modification enzymes may play a role in neuronal plasticity and transmission of information	PMID30289604: intellectual disability, microcephaly, brain anomalies; PMC7981843: includes hearing loss	No
					Heterozygous	SALL4:NM_020436:exon4:c.C2782T:p.R928C	rs201682235	0	ASJ:0.0003	AMR:0.001					27.7	LRT/MA/MT/PP2/PR/SI	PMID16790473: SALL4 haploinsufficiency led to anorectal/heart anomalies, exencephaly and kidney agenesis	MIM607343: AD sensorineural hearing loss due to Duane-radial ray syndrome or IVIC syndrome	No; 1KG MAF>0.001
23	4.61	F		Hearing loss, uncle of grandfather	Potential Compound Heterozygous or Haplotype	CDH23:NM_022124:[exon7:c.C437T:p.P146L]; [exon27:c.G3262A:p.V1088M]; [exon48:c.G6911A:p.R2304Q]	rs765103490; rs200632520; rs201434373	NA; 0.002; 0.0015	NA; EAS:0.002; EAS:0.0015	NA; NA; EAS:0.001			0.001; 0.003; 0.007	24.7; 24.3; 22.7	LRT/MT/PP2/PR/SI; LRT/MA/mLR/mSVM/MT/PP2/SI; MT/SI	KO-hom abnormal righting response, abnormal gait, absent pinna reflex; PMID11138008: disrupted stereocilia organization	MIM605516: DFNB12; AR Usher 1D	No; gnomAD MAF>0.001; previously ruled out due to lack of second allele explaining AR hearing loss	
					Potential Compound Heterozygous or Haplotype	ERC4:NM_005236:[exon8:c.A1483G:p.T495A]; [exon11:c.C2545G:p.Q849E]	NA; rs374186605	NA; 0.001	NA; EAS:0.001				NA; 0.01	21.9; 22	LRT/MA/MT; LRT/MA/MT/PP2	Prewaning lethality	MIM278760: AR Cockayne syndrome with hearing loss	No; gnomAD MAF>0.001	
					Heterozygous	MYO18B:NM_032608:exon8:c.G1982A:p.W661X	rs372939044	0	AFR:0.0005						44	LRT/MT	Het-KO mice with abnormal ABR	MIM616549: AR Klippel-Feil syndrome with nemaline myopathy and facial dysmorphism; PMCS588578: hearing loss common in KFS; PMID18722888: 60% of Klippel-Feil HL patients with ear anomalies including inner ear dysplasia and deformed internal acoustic canal	No
24	6.1	M	EVA, B; fever, jaundice, foul umbilical discharge and apneic episodes with antibiotics and phototherapy in neonatal period		Hemizygous	FLNA:NM_00110556:exon39:c.A6350G:p.N21175	rs375205247	0.002	EAS:0.002						20.2	FA/LRT/MT/PR	NoKOMouse	MIM311300: otopalatodigital syndrome inc. hearing loss; PMC2957847; PMID15654694: Melnick-Needles syndrome includes skull base sclerosis	No; gnomAD MAF>0.001
					Hemizygous	G6PD:NM_000402:exon11:c.C1450T:p.R484C	rs398123546	0.0006	SAS:0.0007	SAS:0.001					31	FA/LRT/MA/mLR/mSVM/MT/PP2/PR/SI	PMCT744953: G6pd overexpression protects from hearing loss progression	MIM300908: hemolytic anemia	No; 1KG MAF>0.001
27	7.72	F	EVA, B; OM, L	Hearing loss, 2nd-degree cousin, paternal side	Heterozygous	MUC1:NM_001204288:exon6:c.G453A:p.W151X	rs369649894	0	LAT:0.00007				0.009	8.35	PP2/PR/SI	NoKOMouse	MIM158340: AD kidney disease; PMID12560150: Muc1 expressed in MEE from patients with otitis media	No	
					Heterozygous	ADAM28:NM_014265:exon7:c.577-1G>A	rs370669549	0.0002	EAS:0.0002				0.006	32	MT	NoKOMouse	PMID23640157: ADAM28 was screened but was negative for AD progressive bone disorder of skull which may include nerve entrapment leading to hearing loss	No	
					Potential Compound Heterozygous or Haplotype	MADD[NM_130470:exon14:c.T2458G:p.S820A]; [exon26:c.T4046A:p.V1349E]	rs778218513; rs756795627	0.0002; 0.0002	EAS:0.0002; EAS:0.0002			NA; 0.006		27.2; 23.4	LRT/MT/PP2/SI; MT/PP2/PR/SI	NoKOMouse	MIM603584: AR neurodevelopmental disorder with dysmorphic facies, impaired speech, hypotonia	No	
					Potential Compound Heterozygous or Haplotype	GDPD5:NM_030792:[exon8:c.G554A:p.R185H]; [exon7:c.C404T:p.T135M]	rs745585758; rs373413383	0.002; 0	ME:0.003; AFR:0.00002			0 (SouthAsia=0.0007); NA		23.1; 24.8	LRT/MT/PP2; LRT/MA/MT/PP2	Hom-KO mice with abnormal ABR (het not tested)		No; gnomAD MAF>0.001	
					Heterozygous	SNAP29:NM_004782:exon5:c.C766T:p.R256X	rs148156702	0.001	EAS:0.001	EAS:0.001					38	LRT/MT	Het-KO mice with increased BMD	MIM609528: AR cerebral dysgenesis, neuropathy, ichthyosis, palmoplantar keratoderma syndrome	No; gnomAD MAF>0.001