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| **Supplementary TABLE 1 |** Incidental findings of insufficient pathogenic evidence genes detected for the three probands in this study by trio-WES |
|  | Gene | Chromosome /exon location | Transcript number/nucleotide variantion/amnio acid change | Pathogenic classification | Zygotic state | Inheritedpattern | Related disease |
|  |  |  |  |  |  |  |  |
| Proband of c.1159C>T | *ADCY1* | Chr7/Exon9 | NM\_021116/c.1789C>T/p.R597W | LP | Het | AR | Autosomal recessive hereditary hearing loss Type 44 |
|  | *ASPM* | Chr1/Exon16 | NM\_018136/c.3812G>A/p.R1271Q | LP | Het | AR | Autosomal recessive hereditary microcephaly Type 5 |
|  | *CASC5* | Chr15/Exon21 | NM\_170589/c.6326C>A/p.T2109N | VUS | Het | AR | Autosomal recessive hereditary microcephaly Type 4 |
|  | *IGF1* | Chr12/Exon12 | NM\_001111283/c.\*4463C>A/- | VUS | Hom | AR | Growth retardation with hearing loss and intellectual disability casused by IGF1 deficiency |
|  | *LOXHD1* | Chr18/Exon32 | NM\_144612/c.4955T>C/p.I1652T | LP | Het | AR | Autosomal recessive hereditary hearing loss Type 77 |
|  | *MOCS2* | Chr5/Exon5 | NM\_004531/c.229A>G/p.T77A | LP | Het | AR | Deficiency of Complementary cofactor Group B |
|  | *PAFAHIB1* | Chr17/Exon2 | NM\_000430/c.-157G>A/- | VUS | Het | - | Anencephalic malformation type 1/ Ectopic subcortical banded gray matter |
|  | *PNPT1* | Chr2/Exon28 | NM\_033109/c.\*152delA/- | VUS | Het | AR | Autosomal recessive hereditary hearing loss Type 70 |
|  |  | NM\_033109/c.\*229\_\*232delAAGT/- |  | Hom |
|  |  | NM\_033109/c.\*258\_\*261delCATA/- |  | Het |
|  | *SMPD1* | Chr11/Exon2 | NM\_000543/c.899C>T/p.T300I | LP | Het | AR | Niemann-Pick Disease Type A |
|  |  | Chr11/Exon4 | NM\_000543/c.1307G>A/p.S436N |
|  | *TUBB3* | Chr16/Exon1 | NM\_006086/c.-115C>G/- | LP | Het | AD | Compound cortical dysplasia with other brain developmental abnormalities Type 1 |
|  |  | Chr16/Exon4 | NM\_001197181/c.\*286A>T/- |
|  |  |  |  |  |  |  |  |
| Proband of c.1309C>T | *NALCN* | Chr13/Exon3 | NM\_052867.2/c.194C>T/p.P65L | VUS | Het | AD | Congenital limb facial contracture, hypotonia and developmental delay |
|  |  |  |  |  |  | AR | Infantile hypotonia with impaired psychomotor development and special facial features Type 1 |
|  | *NALCN* | Chr13/Exon8 | NM\_052867.2/c.874A>G/p.S292G | VUS | Het | AD | Congenital limb facial contracture, hypotonia and developmental delay |
|  |  |  |  |  |  | AR | Infantile hypotonia with impaired psychomotor development and special facial features Type 1 |
|  |  |  |  |  |  |  |  |
| Proband of exon3\_exon4del | *ARX* | ChrX/- | NM-139058.2/c.592G>A/p.V198I | VUS | Het | XL | 1.X-linked anencephalic malformation Type 2 |
|  |  |  |  |  |  |  | 2.Brain edema with genital abnormalities |
|  |  |  |  |  |  |  | 3.Proud syndrome |
|  |  |  |  |  |  |  | 4.Partington syndrome |
|  |  |  |  |  |  |  | 5.Early infantile epileptic encephalopahty |
|  |  |  |  |  |  |  | 6.X linked mental retardation |
|  | *1.C16orf52**2.CDR2**3.EEF2K**4.POLR3E**5.VWA3A* | Chr16:g.21812007\_22547861(GRCh37/hg19)/- | - | Pathogenic | Het | - | 1.Developmental delay2.Mental retardation 3.Language backwardness4.Facial abnormalities5.Microcephaly6.Congenital heart defects7.Schizophrenia |
|  | *UQCRC2* | Chr16:g.21812007\_22547861(GRCh37/hg19)/- | - | Pathogenic | Het | - | 1.Mitochondrial encephalopathy2.Psychomotor delay3.Ataxia4.Liver dysfunction5.Metabolic acidosis6.Renal tubular lesions7.Muscle weakness8.Exercise intolerance |
| Chr: chromosome, Het: heterozygote, Hom: homozygote, LP: likely-pathogenic, VUS: variant of uncertain significance. |