

Supplementary table 1. Molecular Genetic test Report.

Name	***	Age	17 years old
Sampling Date	2020-05-08	Gender	Male
Report Date	2020-06-01	Sample Type	EDTA Whole Blood
Inspection department	The First Affiliated Hospital of China Medical University	Test Item	Focus-hepatopathy

Genetic test results

Genomic DNA was extracted from the samples submitted by the subjects, and the sequencing data were analyzed by constructing genomic DNA library, enriching target DNA fragments, Illumina sequencing and bioinformatics software.

No clear pathogenic changes were observed in this sample within the detection range, but the existence of undetected variation outside the detection range cannot be ruled out.

High-throughput sequencing parameters captured in the target area

Sample number	The coverage of the target area	Average depth of target area	The 20X coverage of the target area
QX2000346401	99.90%	117.310	99.60%

Other mutations of unknown clinical significance

Gene	Variation information	Normal person carrying frequency gnomAD/E xAC	Homozygous/Heterozygous	Sequencing depth Variation ratio	Transcription version Gene subregion	Genetic disease information
<i>AH11</i>	c.3173C>G chr6-13564 p.4455 .A1058G	0 0.000009	Het	48.0/56 0.54	NM_01765 1.4 exon24	1. Joubert syndrome type 3 (AR)
<i>AMACR</i>	c.888G>A chr5-33989 p.459 .P296P	0 0	Het	81.0/73 0.47	NM_01432 4.5 exon5	1. Congenital bile acid synthesis disorder type 4 (AR) 2. A deficiency of α -formyl-coA racemate (AR)

<i>ATP8B1</i>	c.2663C>A chr18-5532 p.8450 .T888K	0 0	Het	96.0/80 0.45	NM_00560 3.4 exon22	1. ICP-1(AD) 2. PFIC-1 (AR) 2. BRIC-1(AR)
<i>ERCC6</i>	c.2645A>G chr10-5068 p.1587 .Y882C	0.000065 0.000016	Het	34.0/19 0.36	NM_00012 4.2 exon14	1. Cockayne syndrome type B (AR) 2. Lung cancer (AD , SMu) 3. Brain-eye-facial skeletal syndrome type 1 (AR) 4. Uv sensitivity syndrome Type 1 (AR) 5. Age related macular degeneration type 5 (-) 6 、 De Sanctis-Cacchione syndrome(AR)
<i>FASTKD2</i>	c.-51+8G>T chr2-20763 0386 splice	0 0	Het	3.0/7 0.7	NM_01492 9.3 intron1	1. Defective mitochondrial complex IV (AR, Mi) 2. Combined oxidative phosphorylation defect type 44 (AR)
<i>GPC4</i>	c.1469-4C> T chrX-13243 7101 splice	0 0.000011	Hem	0.0/36 1	NM_00144 8.2 intron8	1. Keipert syndrome(XR)
<i>LRBA</i>	c.6047-3T> C chr4-15151 8270 splice	0 0.000117	Het	14.0/8 0.36	NM_00672 6.4 intron38	2. Common variant immune deficiency type 8 with autoimmune disease (AR)
<i>TPII</i>	c.631+3G> A chr12-6979 303 splice	0 0.000016	Het	32.0/25 0.44	NM_00036 5.5 intron6	1. Propanose phosphate isomerase deficiency (AR)
<i>UNC13D</i>	c.2588G>A chr17-7382 p.7216 .G863D	0.000194 0.000358	Het	74.0/47 0.39	NM_19924 2.2 exon27	1. Familial hemophagocytic histiocytosis type 3 (AR)

List of Genetic Tests

Congenital hepatic fibrosis						49
<i>AHI1</i>	<i>ALMS1</i>	<i>ARL13B</i>	<i>ARL6</i>	<i>B9D1</i>	<i>B9D2</i>	<i>BBIP1</i>
<i>BBS1</i>	<i>BBS10</i>	<i>BBS12</i>	<i>BBS2</i>	<i>BBS4</i>	<i>BBS5</i>	<i>BBS7</i>
<i>BBS9</i>	<i>CC2D2A</i>	<i>CEP290</i>	<i>CEP41</i>	<i>CPLANE1</i>	<i>CSPP1</i>	<i>DNAJB11</i>
<i>GANAB</i>	<i>IFT27</i>	<i>INPP5E</i>	<i>IQCB1</i>	<i>KIAA0586</i>	<i>LZTFL1</i>	<i>MKS1</i>
<i>NPHP1</i>	<i>NPHP3</i>	<i>OFD1</i>	<i>PKD1</i>	<i>PKD2</i>	<i>PKHD1</i>	<i>RPGRIP1L</i>
<i>SDCCAG8</i>	<i>TCTN1</i>	<i>TCTN2</i>	<i>TMEM107</i>	<i>TMEM138</i>	<i>TMEM216</i>	<i>TMEM231</i>
<i>TMEM237</i>	<i>TMEM67</i>	<i>TRAF3IP1</i>	<i>TRIM32</i>	<i>TTC8</i>	<i>WDPCP</i>	<i>WDR19</i>
Wilson disease						1
<i>ATP7B</i>						
Cholestatic liver disease						16
<i>ABCB11</i>	<i>ABCB4</i>	<i>ABCD3</i>	<i>ACOX2</i>	<i>AKR1D1</i>	<i>AMACR</i>	<i>ATP8B1</i>
<i>CYP7B1</i>	<i>HSD3B7</i>	<i>JAG1</i>	<i>NOTCH2</i>	<i>NR1H4</i>	<i>SLC10A2</i>	<i>TJP2</i>
<i>VIPAS39</i>	<i>VPS33B</i>					
Mitochondrial liver disease						70
<i>ATP5F1A</i>	<i>ATP5F1E</i>	<i>ATPAF2</i>	<i>BCS1L</i>	<i>BOLA3</i>	<i>COA8</i>	<i>COX10</i>
<i>COX14</i>	<i>COX20</i>	<i>COX6B1</i>	<i>COX8A</i>	<i>CYC1</i>	<i>DGUOK</i>	<i>ECHS1</i>
<i>FASTKD2</i>	<i>FBXL4</i>	<i>FOXRED1</i>	<i>ISCA2</i>	<i>LYRM7</i>	<i>MGME1</i>	<i>MPC1</i>
<i>MPV17</i>	<i>MRM2</i>	<i>NDUFA1</i>	<i>NDUFA10</i>	<i>NDUFA11</i>	<i>NDUFA12</i>	<i>NDUFA2</i>
<i>NDUFA4</i>	<i>NDUFA9</i>	<i>NDUFAF1</i>	<i>NDUFAF2</i>	<i>NDUFAF3</i>	<i>NDUFAF4</i>	<i>NDUFAF5</i>
<i>NDUFAF6</i>	<i>NDUFB11</i>	<i>NDUFB3</i>	<i>NDUFB9</i>	<i>NDUFS1</i>	<i>NDUFS2</i>	<i>NDUFS3</i>
<i>NDUFS4</i>	<i>NDUFS6</i>	<i>NDUFS7</i>	<i>NDUFS8</i>	<i>NDUFV1</i>	<i>NDUFV2</i>	<i>NFS1</i>
<i>NUBPL</i>	<i>PCK2</i>	<i>PET100</i>	<i>POLG</i>	<i>POLG2</i>	<i>RRM2B</i>	<i>SCO1</i>
<i>SLC25A4</i>	<i>SUCLA2</i>	<i>SUCLG1</i>	<i>TACO1</i>	<i>TFAM</i>	<i>TK2</i>	<i>TMEM126B</i>
<i>TTC19</i>	<i>TWNK</i>	<i>UQCC2</i>	<i>UQCC3</i>	<i>UQCRB</i>	<i>UQCRC2</i>	<i>UQCRQ</i>

Glucose metabolic disease						38
<i>AGL</i>	<i>ALDOA</i>	<i>ARSB</i>	<i>CTSA</i>	<i>ENO3</i>	<i>FBP1</i>	<i>G6PC</i>
<i>GAA</i>	<i>GALE</i>	<i>GALK1</i>	<i>GALNS</i>	<i>GALT</i>	<i>GBE1</i>	<i>GLB1</i>
<i>GNS</i>	<i>GUSB</i>	<i>GYG1</i>	<i>GYS1</i>	<i>GYS2</i>	<i>HGSNAT</i>	<i>HYAL1</i>
<i>IDS</i>	<i>IDUA</i>	<i>LDHA</i>	<i>NAGLU</i>	<i>PFKM</i>	<i>PGAM2</i>	<i>PGM1</i>
<i>PHKA1</i>	<i>PHKA2</i>	<i>PHKB</i>	<i>PHKG2</i>	<i>PRKAG2</i>	<i>PYGL</i>	<i>PYGM</i>
<i>SGSH</i>	<i>SLC37A4</i>	<i>TPI1</i>				
Defective glycoprotein synthesis						40
<i>ALG1</i>	<i>ALG11</i>	<i>ALG12</i>	<i>ALG13</i>	<i>ALG2</i>	<i>ALG3</i>	<i>ALG6</i>
<i>ALG8</i>	<i>ALG9</i>	<i>B4GALT1</i>	<i>CCDC115</i>	<i>COG1</i>	<i>COG2</i>	<i>COG4</i>
<i>COG5</i>	<i>COG6</i>	<i>COG7</i>	<i>COG8</i>	<i>DDOST</i>	<i>DOLK</i>	<i>DPAGT1</i>
<i>DPM1</i>	<i>DPM2</i>	<i>DPM3</i>	<i>MGAT2</i>	<i>MOGS</i>	<i>MPDU1</i>	<i>MPI</i>
<i>NGLY1</i>	<i>PMM2</i>	<i>RFT1</i>	<i>SLC35A1</i>	<i>SLC35A2</i>	<i>SLC35C1</i>	<i>SRD5A3</i>
<i>SSR4</i>	<i>STT3A</i>	<i>STT3B</i>	<i>TMEM165</i>	<i>TMEM199</i>		
Aminoacidopathy						53
<i>ABAT</i>	<i>ABCG5</i>	<i>ADA</i>	<i>ADK</i>	<i>AGA</i>	<i>AHCY</i>	<i>AMT</i>
<i>ASL</i>	<i>ASS1</i>	<i>CBS</i>	<i>COQ2</i>	<i>CPS1</i>	<i>CPT1A</i>	<i>CPT2</i>
<i>CTNS</i>	<i>CYP7A1</i>	<i>DNAJC19</i>	<i>ETFA</i>	<i>ETFB</i>	<i>ETFDH</i>	<i>FAH</i>
<i>GCDH</i>	<i>GCH1</i>	<i>GCSH</i>	<i>GLDC</i>	<i>GNMT</i>	<i>GSTZ1</i>	<i>HADH</i>
<i>HMGCL</i>	<i>HMGCS2</i>	<i>HPD</i>	<i>MAT1A</i>	<i>MCCCI</i>	<i>MCCC2</i>	<i>MMAA</i>
<i>MMAB</i>	<i>MMACHC</i>	<i>MMUT</i>	<i>NAGA</i>	<i>OTC</i>	<i>PC</i>	<i>PCBD1</i>
<i>PCCA</i>	<i>PCCB</i>	<i>PDSS2</i>	<i>PEPD</i>	<i>PKLR</i>	<i>SLC22A5</i>	<i>SLC25A13</i>
<i>SLC25A15</i>	<i>SLC25A20</i>	<i>SLC7A7</i>	<i>TAT</i>			
Hemochromatosis						7
<i>BMP2</i>	<i>FTH1</i>	<i>HAMP</i>	<i>HFE</i>	<i>HJV</i>	<i>SLC40A1</i>	<i>TFR2</i>
Porphyrinopathy						8
<i>ALAD</i>	<i>ALAS2</i>	<i>CPOX</i>	<i>FECH</i>	<i>HFE</i>	<i>HMBS</i>	<i>UROD</i>
<i>UROS</i>						
Abnormal metabolism of cholechrome						4
<i>ABCC2</i>	<i>SLCO1B1</i>	<i>SLCO1B3</i>	<i>UGT1A1</i>			

Peroxisomal disorders						17
<i>ABCD1</i>	<i>ACOX1</i>	<i>HSD17B4</i>	<i>PEX1</i>	<i>PEX10</i>	<i>PEX11B</i>	<i>PEX12</i>
<i>PEX13</i>	<i>PEX14</i>	<i>PEX16</i>	<i>PEX19</i>	<i>PEX2</i>	<i>PEX26</i>	<i>PEX3</i>
<i>PEX5</i>	<i>PEX6</i>	<i>PEX7</i>				
Polycystic liver disease						4
<i>ALG8</i>	<i>LRP5</i>	<i>PRKCSH</i>	<i>SEC63</i>			
Complex oxidative phosphorylation deficiency						23
<i>AARS2</i>	<i>AIFM1</i>	<i>ATP5F1A</i>	<i>C12orf65</i>	<i>C1QBP</i>	<i>ELAC2</i>	<i>FARS2</i>
<i>GFM1</i>	<i>LYRM4</i>	<i>MRPL3</i>	<i>MRPL44</i>	<i>MRPS16</i>	<i>MRPS22</i>	<i>MRPS7</i>
<i>MTFMT</i>	<i>PNPT1</i>	<i>RMND1</i>	<i>SFXN4</i>	<i>TARS2</i>	<i>TRMT10C</i>	<i>TSFM</i>
<i>TUFM</i>	<i>VAR2</i>					
Glucolipid metabolic disease						4
<i>GBA</i>	<i>NPC1</i>	<i>NPC2</i>	<i>SMPD1</i>			