

**Supplementary Table S1. Variants identified by clinical exome sequencing in the proband with facial, digital, cardiac anomalies with developmental delay**

Gene	Reference transcript ID	Base change	Codon change	Variant type	OMIM phenotype	Inheritance
<i>ABCA7</i>	NM_019112.4	c.5961G>T	p.Glu1987Asp	missense	{Alzheimer disease 9, susceptibility to}	AD
<i>AKT2</i>	NM_001626	.6c.431G>A	p.Arg144Gln	missense	Diabetes mellitus, type II   Hypoinsulinemic hypoglycemia with hemihypertrophy	AD   AD
<i>ANK3</i>	NM_020987.	5c.8506C>T	p.His2836Tyr	missense	Intellectual developmental disorder, autosomal recessive 37	AR
<i>DNAH10</i>	NM_207437.3	c.10444C>T	p.Arg3482Cys	missense	Spermatogenic failure 56	AR
<i>FSIP2</i>	NM_173651.4	c.3351G>A	p.Met1117Ile	missense	Spermatogenic failure 34	AR
<i>HSPA9</i>	NM_004134.7	c.661A>C	p.Asn221His	missense	Anemia, sideroblastic, 4   Even-plus syndrome	AD   AR
<i>HYDIN</i>	NM_001270974.2	c.13080G>T	p.Glu4360Asp	missense	Ciliary dyskinesia, primary, 5	AR
<i>IGSF3</i>	NM_001542.4	c.1784G>A	p.Trp595Ter	nonsense	?Lacrimal duct defect	AR
<i>ITGA8</i>	NM_003638.3	c.3128G>T	p.Arg1043Ile	missense	Renal hypodysplasia/aplasia 1	AR
<i>LIPA</i>	NM_000235.4	c.10C>T	p.Arg4Trp	missense	Cholesteryl ester storage disease   Wolman disease	AR   AR
<i>RECQL</i>	NM_002907.4	c.1784A>G	p.Gln595Arg	missense	RECON progeroid syndrome	AR
<i>TPP2</i>	NM_003291.4	c.1067A>G	p.Tyr356Cys	missense	Immunodeficiency 78 with autoimmunity and developmental delay	AR
<i>TRAF7</i>	NM_032271.3	c.1964G>A	p.Arg655Gln	missense	Cardiac, facial, and digital anomalies with developmental delay	AD
<i>VANGL2</i>	NM_020335.3	c.1319G>A	p.Arg440Gln	missense	Neural tube defects	AD

OMIM, Online Mendelian Inheritance in Man; AD, autosomal dominant; AR, autosomal recessive