

<u>Gene</u>	<u>OMIM</u>	<u>Inheritance</u>	<u>Pt. Zygosity</u>	<u>Position</u>	<u>Nucleotide</u>	<u>Amino Acid</u>	<u>Parental Studies</u>	<u>Comment</u>
<i>DNM1L</i>	614388	AD	Het	Chr12:32875536	c.1048G>A	p.G350R	Mother 6-8% mosaicism	Novel variant
<i>ACADSB</i>	610006	AR	Het	Chr10:124812677	c.1228+1G>A	N/A	Father heterozygous	One variant detected (AR disorder)
<i>ACSF3</i>	614265	AR	Het	Chr16:89199607	c.1303G>T	p.E435X	Father heterozygous	One variant detected (AR disorder)
<i>CLN8</i>	600143	AR	Het	Chr8:1728557	c.685C>G	p.P229A	ND	One variant detected (AR disorder)
<i>COQ9</i>	614654	AR	Het	Chr16:57485113	c.235C>T	p.P79S	ND	One variant detected (AR disorder)
<i>AH11</i>	608629	AR	Het	Chr6:135778794	c.989A>G	p.D330G	ND	One variant detected (AR disorder)
<i>DHFR</i>	613839	AR	Het	Chr5:79950481	c.68C>T	p.S23F	ND	One variant detected (AR disorder)
<i>SLC25A3</i>	610773	AR	Het	Chr12:98987859	c.103G>C	p.G35R	ND	One variant detected (AR disorder)
<i>CHKB</i>	602541	AR	Het	Chr22:51021197	c.14C>T	p.A5V	ND	One variant detected (AR disorder)
<i>SLC6A3</i>	613135	AR	Het	Chr5:1403128	c.1676C>T	p.A559V	ND	One variant detected (AR disorder)
<i>PEX1</i>	214100	AR	Het	Chr8:92129015	c.2718+3G>A	N/A	ND	One variant detected (AR disorder)
<i>ALG13</i>	601110	XL	Hem	ChrX:110970229	c.1922A>G	p.H641R	Mother heterozygous	Patient had normal N-glycan analysis
<i>MECP2</i>	300672	XL	Hem	ChrX:153296146	c.1133C>T	p.A378V	Mother heterozygous	Most variants X-linked dominant; Male with same variant seen in ExAC data; "benign clinical significance in ClinVar; rs201314910