Title: Carnitine Palmitoyltransferase 1A Deficiency GeneReview Table 2

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Note: The following information is provided by the authors listed above and has not been reviewed by *GeneReviews* staff.

Table 2. Mutations in CPT1A

cDNA	Protein	Exon	
Homozygous missense and nonsense mutations			
298C>T	Q100X	4	
367C>T	R123C	4	
986C>T	T314I	9	
1069C>T	R357W	10	
1361A>G	D454G	12	
1436C>T	P479L	12	
2129G>A	G710E	17	
2126G>A	G709E	17	
478C>T	R160X	5	
1027T>G	F343V	10	
1393G>T	G465W	12	
1737C>A	Y579X	14	
Heterozygous missense and nonsense mutations			
96T>G	Y32X	2	
1079A>G	E360G	10	
823G>A	A275T	8	
912C>G	C304W	9	
823G>A	A275T	8	
1241C>T	A414V	11	
1493A>G	Y498C	13	
1069C>T	R357W	10	
1451T>C	L484P	12	
1425G>A	W475X	12	
1494T>G	Y498X	13	
946C>G?	R316G?	9	
1339C>T	R446X	11	
2156G>A	G719D	18	

cDNA	Protein	Exon
Deletions and insertion		
Homozygous IVS 14+3kb	del 581-702	15
E525 Ins FLPYHELSX	Trunc 240 amino acids del 626-676	16
1876-1G>A		
2027-2028 +2del AAGT		17
948delG	R316fsX328	9
1600delC	L534fsX	13

Mutations are matched to the protein sequence (GI 32879925) in the PubMed protein bank [Brown et al 2001, Gobin et al 2002, Bennett et al 2004, Stoler et al 2004, Korman et al 2005, Tsuburaya et al 2010].

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