

## SUPPLEMENTARY MATERIAL TO:

### Diseases caused by mutations in mitochondrial carrier genes *SLC25*: a review

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**Table S1. Disease-causing mutations in mitochondrial carriers.** For each gene/carrier the mutations are listed in the following order: first the nonsense, deletion/insertion and splicing mutations, and afterwards the missense mutations.

Mitochondrial carrier	Disease-associated mutation	Mutation effect in the protein (or problem in splicing)	Homo- / Heterozygous patients	References
<b>SLC25A1, CIC</b>  <b>Position in Fig. 2</b>	c.18_24dup	p.Ala9Profs*82	2/1	(1) (2)
	c.517_526del	p.Arg173Glyfs*2	1/1	(1) (3)
	c.648_655del	p.Met218Serfs*25	0/1	(3)
	c.768C > G	p.Tyr256*	0/1	(1)
	c.821C > T	p.Ala274Ilefs*24	2/4	(1)
	10 c.82G > A	p.Ala28Thr	0/1	(3)
	22 c.119T > A	p.Ile40Asn	0/1	(3)
	27 c.134C > T	p.Pro45Leu	0/2	(1)(2)
	29 c.139G > A	p.Glu47Lys	1/0	(3)
	55 c.205G > T	p.Asp69Tyr	3/0	(4)
	79 c.278G > A	p.Gly93Asp	1/0	(3)
	120 c.389G > A	p.Gly130Asp	0/1	(5)
	134 c.430G > C	p.Glu144Gln	1/0	(1)
	159 c.499G > A	p.Gly167Arg	0/1	(1)
	185 c.578C > G	p.Ser193Trp	1/1	(1)(3)
	190 c.593G > A	p.Arg198His	1/0	(6)
	194 c.605T > C	p.Met202Thr	0/2	(1)
	227 c.713A > G	p.Asn238Ser	1/0	(7)
	236 c.740G > A	p.Arg247Gln	12/0	(4)(8)(9)
	256 c.784T > C	p.Cys262Arg	1/0	(3)

276	c.844C > T	p.Arg282Cys	2/1	(1)
276	c.844C > G	p.Arg282Gly	1/0	(1)
276	c.845G > A	p.Arg282His	0/1	(5)
291	c.890A > G	p.Tyr297Cys	0/1	(1)
<b>SLC25A3, PiC</b>	c.158-9A > G c.886-898delins7	intron splice site p.296_300Ins	4/0 0/1	(10) (11)
<b>Position in Fig. 2</b>				
14	c.215G > A	p.Gly72Glu	2/0	(12)
156	c.599T > G	p.Leu200Trp	0/1	(11)
<b>SLC25A4, AAC1</b>	c.111+1G > A c.116_137del c.390del	p.Gln39Leufs*14 no exon 1 p.Phe130Leufs*41	0/1 1/0 0/1	(13) <sup>1</sup> (14) <sup>1</sup> (15) <sup>1</sup>
<b>Position in Fig. 2</b>				
32	c.97A > T	p.Lys33Gln	0/1	(16) <sup>2</sup>
79	c.239G > A	p.Arg80His	1/0	(17) <sup>2</sup>
89	c.269C > A	p.Ala90Asp	0/3	(18) <sup>3</sup>
97	c.293T > C	p.Leu98Pro	0/2	(19) <sup>3</sup>
103	c.311A > G	p.Asp104Gly	0/4	(20) <sup>3</sup>
113	c.340C > G	p.Ala114Pro	?	(21) <sup>3</sup>
122	c.368C > A	p.Ala123Asp	1/1	(15) <sup>1</sup> (22) <sup>1</sup>
140	c.423G > C	p.Leu141Phe	4/0	(15) <sup>1</sup>
217	c.653A > C	p.Gln218Pro	1/0	(23)
234	c.703C > G	p.Arg235Gly	3/0	(17) <sup>2</sup>
235	c.707G > C	p.Arg236Pro	0/1	(13) <sup>1</sup>
288	c.865G > A	p.Val289Met	?	(21) <sup>3</sup>
<b>SLC25A10, DIC</b>	c.304A > T c.790-37G > A <sup>4</sup>	p.Lys102* intron	0/1 0/1	(24) (24)
	c.684C > T <sup>4</sup>	p.Pro228Pro	0/1	(24)
<b>SLC25A12, AGC1</b>				
<b>Position in Fig. 2</b>				
34	c.1058G > A	p.Arg353Gln	2/0	(25)
130	c.1331C > T	p.Thr444Ile	1/0	(26)
283	c.1769A > G	p.Gln590Arg	1/0	(27)
<b>SLC25A13, AGC2</b>	c.-3251_c.15+18443del2 1709bp c.2T > C c.15G > A g.16-2A > T r.16_212dup c.46G > T c.70-862_212+3527del453 2 c.70_215del c.72T > A c.127C > T c.172_173delIGT c.265delIG	gross deletion  missing start codon splicing splicing Ex2_3dup p.Glu16* deletion  deletion p.Tyr24* p.Arg43* p.Val58Gfs* p.Asp89fs*	?  0/3 0/8 0/1 ? 0/2 0/1  0/1 0/1 1/3 2/0 0/1	(28)  (29)(30)(31) (32)(33) (34) (35) (32) (36)  (36) (34) (37) (37) (34)

IVS4ins6kb	p.Glu110fs*	0/7	(31)(34)
c.329-1687_c.468+3865del	p.Glu110fs*	0/1	(31)(38)
c.329-154_c468+252del2646;c468+2394_c468ins23	p.Glu110fs*	0/1	(31)
c.448G > T	p.Glu150*	0/1	(34)
c.475C > T	p.Gln159*	0/2?	(34)(39)
c.478delC	p.Leu160Trpfs*	1/0	(40)
c.493C > T	p.Gln165*	0/1	(31)
c.495delA	p.Gln165*fs	0/1	(31)
c.550C > T	p.Arg184*	0/7	(31)(32)(41)(42)(43)
c.IVS6(1789bp)ins (c.615+1G > C)	p.Ala206fs*	1/1?	(32)
c.615+1G > A	splicing	2/5?	(37)(39)(42)(43)(44)
c.640C > T	p.Gln214*	0/1	(45)
IVS6+5G > A (c.615+5G > A)	splicing	0/53	(31)(34)(41)(42)(43)(46)(47)(48)(49)
c.650delT	p.Phe217fs*	?	(50)
c.674C > A	p.Ser225*	8/19?	(33)(51)(52)(53)(54)(55)
c.754G > A	p.Ala206fs*	0/3	(29)(34)(56)
c.754+6T>G(IVS7+6T>G)	splicing?	?	(45)
c.755-1G > C	p.252fs*	0/1	(31)
c.755-2A > G	splicing	0/1	(57)
c.775C > T	p.Gln259*	0/4	(31)(34)(42)(46)
c.845_848+1delG	p.Gly283fs*	0/2	(31)(58)
c.847G > T	p.Gly283*	0/1	(48)
c.848+3A > C	splicing	0/1	(37)
c.851_854del4 (c.852_855del4)	p.Arg284fs*	214/348?	(29)(30)(31)(33)(34)(37)(39)(42)(43)(45)(46)(47)(48)(51)(52)(53)(54)(59)(60)
c.933G > A	splicing ?	0/1	(33)
c.933_933ins+1GCA G	p.Ala312fs*	0/1	(31)
c.955C > T	p.Arg319*	0/4	(34)(42)(47)(48)(57)
c.985_986insT	p.Ala329fs*	0/1	(42)
c.1019_1177del (c.1177+1G>A)	p.340_392del	74/102	(30)(31)(33)(34)(43)(48)(51)(52)(53)(54)(59)(61)(62)
c.1019_c.1177+893del	p.340_392del	?	(63)
c.1063C > T	p.Arg355*	0/3	(37)
c.1078C > T	p.Arg360*	0/5?	(31)(32)(42)(43)(48)
c.1092_1095delT	p.Phe365fs*	0/8	(31)(34)(41)(43)
c.1146delA	p.Arg383fs*	0/1	(32)
c.1189C > T	p.Gln397*	0/1	(32)
c.1192_1193delT	p.Leu398fs*	?	(64)
c.1231_1311del (c.1311+1G>A)	p.411_437del	5/26?	(33)(51)(53)(54)
g.IVS13 + 2T >G	splicing	0/1	(32)
c.1375delG	p.Ala459fs*	0/1	(32)
c.1381G > T	p.Glu461*	0/1	(31)
c.1399C > T	p.Arg467*	0/7?	(31)(34)(48)(64)
c.1452+1G > A	splicing	0/1	(31)
c.1453_1591dup	p.Met532fs*	4/0?	(32)(65)
c.1453_1591del	p.Gly485fs*	0/1	(32)
c.1610_1612del2ins2	p.Leu537Tyrfs*	2/0	(66)(67)
c.1645C > T	p.Gln549*	0/1?	(34)(68)
c.1638_1660dup23	p.Ala554Glyfs*	17/66?	(29)(31)(34)(37)(39)(42)(43)

	Ex16+74_IVS17-32del516	p.Gln556fs*	2/6?	(45)(46)(47)(48)(51)(53)(54)(32)(62)
	c.1706_1707delTA	p.Thr569fs*	0/1	(31)(34)
	c.1709_1710insA(1706_1707insA)	p.Ile570fs*	?	(69)
	c.1736G > A	p.Trp579*	0/1	(34)
	IVS16ins3kb	p.Ala584fs*	7/109?	(29)(30)(31)(32)(34)(42)(43)(45)(46)(47)(48)
	c.1799_1800insA	p.Tyr600*	0/20?	(32)(33)(53)(54)
	c.1801G > T	p.Glu601*	0/15?	(32)(42)(52)(54)
	c.1813C > T	p.Arg605*	1/8?	(32)(53)(54)
	c.1841+3_1841+4delAA	splicing	0/2	(58)(60)
	c.74C > A	p.Ala25Glu	0/1	(37)
	c.103A > G	p.Met35Val	0/1	(31)
	c.115G > T	p.Asp39Tyr	?	(70)
	c.221C > T	p.Ser74Phe	?	(68)
	c.254T > C	p.Leu85Pro	1/0	(42)(43)
	c.284C > A	p.Ala95Asp	0/1	(45)
	c.287T > C	p.Phe96Ser	0/2	(42)
	c.415G > A	p.Gly139Arg	0/1	(34)
	c.443A > G	p.Tyr148Cys	0/1	(34)
	c.527G > T	p.Gly176Val	0/1	(34)
	c.790G > A	p.Val264Ile	?	(71)
	c.869T > C	p.Ile290Thr	?	(50)
	c.998G > A	p.Gly333Asp	0/1	(47)(48)
	c.1046T > C	p.Ile349Thr	?	(72)
	c.1048G > A	p.Asp350Asn	0/5	(31)(34)
	c.1063C > G	p.Arg355Gly	0/2	(31)(34)
	c.1064G > A	p.Arg355Gln	0/2	(31)(49)
	c.1067T > A	p.Met356Lys	?	(72)
	c.1157G > T	p.Gly386Val	0/1	(63)
	c.1177G > A	p.Gly393Ser	0/1?	(31)(32)(39)
	c.1215G > T	p.Lys405Asn	0/2	(34)
	c.1231G > A	p.Val411Met	0/1	(48)
	c.1307_1308del2ins2	p.Gly436Glu	0/1	(73)
	c.1311C > T	p.Cys437Cys	1/0	(46)
	c.1336A > C	p.Thr446Pro	0/1	(32)
	c.1349A > G	p.Glu450Gly	0/1	(42)
	c.1354G > A	p.Val452Ile	0/1	(74)
	c.1357A > G	p.Lys453Glu	?	(75)
	c.1358A > G	p.Lys453Arg	0/1	(64)
	c.1364G > T	p.Arg455Leu	?	(34)
	c.1420G > A	p.Val474Met	?	(76)
	c.1430T > G	p.Leu477Arg	?	(77)
	c.1465T > C	p.Cys489Arg	0/1	(32)(66)
	c.1478A > G	p.Asp493Gly	0/2	(55)
	c.1498T > G	p.Tyr500Asp	0/1	(34)
	c.1505C > T	p.Pro502Leu NP	0/1	(46)(78)
	c.1592G > A	p.Gly531Asp	0/1	(34)
	c.1622C > A	p.Ala541Asp	0/11	(47)(48)(57)
	c.1637C > T	p.Thr546Met	0/4	(34)
	c.1637C > G	p.Thr546Arg	0/1	(37)
	c.1658G > A	p.Arg553Gln	0/1	(39)
	c.1754G > A	p.Arg585His	0/1?	(42)(43)
	c.1763G > A	p.Arg588Gln	0/2	(32)
	c.1763G > C	p.Arg588Pro	?	(68)

**Position in  
Fig. 2**

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283	c.1775A > C	p.Gln592Pro	1/0?	(34)
289	c.1793T > G	p.Leu598Arg	0/1	(32)
292	c.1801G > A	p.Glu601Lys	2/0?	(32)(43)(54)
296	c.1814G > A	p.Arg605Gln NP	0/5	(29)(78)
	c.1895C > T	p.Pro632Leu NP	0/1	(32)(78)
	c.1915G > A	p.Gly639Ser	?	(77)
<b>SLC25A15, ORC1</b>	c.56+1G > T	intron/fs	1/0	(79)
	c.96_97insCA	p.Met33Glnfs*	2/0?	(80)(81)
	c.164insA	p.Tyr55*	0/1	(82)
	c.265C > T	p.Gln89*	1/0	(82)
	c.446delG	p.S149Tfs*	2/0	(83)
	c.525insC	p.Ser175fs*	0/1	(79)
	c.535C > T	p.Arg179*	9/2	(79)(81)(82)(84)(85)(86)
	c.553_564del	p.Phe188del	32/7?	(80)(81)(87)(88)(89)
	c.684_685insAAC	p.228_229insAsn	1/0	(84)
	c.733A > T	p.Lys245*	2/0	(79)
	g.823C > T	p.Arg275*	2/1	(90)(91)
	c.861insG	p.Ser290*	1/0	(82)
	13q14del	large del	0/1	(87)
	IVS5+1G > A	exon 5 skipping	1/0	(82)
<b>Position in Fig. 2</b>				
13	c.44C > A	p.Ala15Glu	1/0	(80)
13	c.44C > T	p.Ala15Val	1/0	(92)
25	c.79G > A	p.Gly27Arg	2/3	(79)(82)
25	c.80G > A	p.Gly27Glu	1/0	(84)
28	c.88T > G	p.Phe30Val	0/1	(93)
30	c.95C > G	p.Thr32Arg	5/0	(94)
35	c.110T > G	p.Met37Arg	1/0	(79)
36	c.113A > C	p.Gln38Pro	0/1	(93)
76	c.208_209delCAinsTT	p.Ala70Leu	2/0	(79)
77	c.212T > A	p.Leu71Gln	1/0	(79)
119	c.337G > T	p.Gly113Cys	0/1	(95)
132	c.377C > G	p.Pro126Arg	1/0	(96)
183	c.538G > A	p.Glu180Lys	0/1	(87)
191	c.564C > G	p.Phe188Leu	1/0	(79)
193	c.572G > A	p.Gly190Asp	0/1	(82)
196	c.568T > C	p.Leu193Pro	0/1	(89)
216	c.646G > A	p.Gly216Ser	1/0	(79)
220	c.658G > A	p.Gly220Arg	3/1	(91)(97)
268	c.790G > C	p.Ala264Pro	0/2	(81)
276	c.815C > T	p.Thr272Ile	1/2	(79)(86)
277	c.818T > A	p.Met273Lys	0/1	(95)
279	c.824G > A	p.Arg275Gln	1/0	(82)
279	?	p.Arg275Gly	0/2	?
287	c.847C > T	p.Leu283Phe	0/1	(79)
<b>SLC25A16</b>				
<b>Position in Fig. 2</b>				
2	c.92G > T	p.Arg31Leu	9/0	(98)
<b>SLC25A19, TPC</b>				
<b>Position in Fig. 2</b>				
119	c.373G > A	p.Gly125Ser	4/0	(99)
175	c.530G > C	p.Gly177Ala	?	(100)

190	c.576G > C	p.Gln192His	1/0	(101)
192	c.580T > C	p.Ser194Pro	1/0	(102)
278	c.869T > A	p.Leu290Gln	1/0	(103)
292	c.910G > A	p.Glu304Lys	1/0	(103)
<b>SLC25A20, CAC</b>	c.65_69insTGTGC	p.Leu24Cysfs*	0/1	(104)
	c.84delT	p.His29Thrfs*	0/2	(105)(106)
	c.106-2A>T	splicing	0/1	(107)
	c.160_163del4ins4	p.54_55delGT insWA	1/2	(108)(109)
	c.168delT	p.Phe56Leufs*	0/1	(108)
	c.180delG	p.Lys61Argfs*	1/0	(104)
	c.199 - 10T > G	splicing	6/4	(104)(105)(106)(107)(110)
	c.261_388del1128		0/1	(111)
	c.270delC	p.Phe91Leufs*	1/0	(105)
	c.326+1delG	splicing	1/2	(105)(112)(113)
	c.362delG	p.Gly121Alafs*	0/1	(109)
	c.496C > T	p.Arg166*	1/1	(104)(108)(114)
	c.528delT	p.Met177Cysfs*	0/1	(108)
	c.532C > T	p.Arg178*	1/3	(104)(109)(115)
	c.516T > A	p.Trp192*	0/2	(107)
	c.609 - 1G > A	splicing	1/2	(104)
	c.609-3C>G	splicing	0/1	(113)
	c.671_780del110		0/1	(111)
	c.718+1G>C	splicing	0/1	(109)
	c.752_761del10	p.Asp251Gfs*	0/1	(108)
	c.779_781delAAG	p.Glu260del	0/1	(108)
	c.804delG	p.Phe269Serfs*	1/1	(104)(108)
	c.823C > T	p.Arg275*	0/1	(108)
	c.843+4_843+50del47	splicing	1/0	(109)
	c.897insC	p.Asn300Glnfs*	1/0	(116)
	26kdel	Ex5-9del	0/1	(108)
<b>Position in Fig. 2</b>				
20	c.67T > C	p.Cys23Arg	1/0	(108)
25	c.82G > T	p.Gly28Cys	1/0	(104)
29	c.94G > A	p.Asp32Asn	0/1	(104)
53	c.164C > T	p.Thr55Asn	0/1	(108)
79	c.241G > A	p.Gly81Arg	2 alleles?	(117)
85	c.260C > T	p.Ala87Val	0/1	(108)
135	c.397C > T	p.Arg133Trp	0/3	(108)(109)
182	c.533G > A	p.Arg178Gln	1/1	(104)
230	c.689C > G	p.Pro230Arg	0/1	(104)
231	c.691G > C	p.Asp231His	0/2	(104)(109)
238	c.713A > G	p.Gln238Arg	2/0	(118)(119)
285	c.842C > T	p.Ala281Val	1/0	(109)
<b>SLC25A21, ODC</b>				
<b>Position in Fig. 2</b>				
234	c.695A > G	p.Lys232Arg	1/0	(120)
<b>SLC25A22, GC1</b>	c.813_814delTG	p.Ala272Glnfs*	0/2	(121)
<b>Position in Fig. 2</b>				
60	c.166A > C	p.Thr56Pro	3/0	(122)
83	c.235G > A	p.Glu79Lys	0/1	(122)
83	c.235G > C	p.Glu79Gln	0/1	(123)
119	c.328G > C	p.Gly110Arg	2/0	(124)
192	c.617C > T	p.Pro206Leu	5/0	(125)(126)

220	c.706G > T	p.Gly236Trp	1/0	(127)
230	c.736T > C	p.Cys246Arg	0/1	(123)
233	c.746T > A	p.Val249Glu	0/1	(122)
258	c.818G > A	p.Arg273Lys	0/2	(121)
281	c.886G > A	p.Ala296Thr	2/0	(122)
<b>SLC25A24, APC1</b> <u>Position in Fig. 2</u>				
30	c.649C > T	p.Arg217Cys	0/2	(128)(129)
30	c.650G > A	p.Arg217His	0/9	(128)(129)(130)(131)
<b>SLC25A26, SAMC</b> <u>Position in Fig. 2</u>				
	c.33+1G > A	splice site	1/0	(132)
126	c.305C > T	p.Ala102Val	0/1	(132)
180	c.443T > G	p.Val148Gly	1/0	(132)
229	c.596C > T	p.Pro199Leu	0/1	(132)
<b>SLC25A32</b> <u>Position in Fig. 2</u>				
	c.425G > A	p.Trp142*	0/1	(133)
	c.-264_31delins14	missing start codon	1/0	(134)
139	c.440G > A	p.Arg147His	0/1	(133)
<b>SLC25A38</b> <u>Position in Fig. 2</u>				
	c.175C > T	p.Gln59*	0/1	(135)
	c.324_325delCT	p.Tyr109Leufs*	2/5	(136)(137)
	c.324_330del7bp	p.Leu108fs*	1/0	(138)
	c.336_347del11bp	p.Lys112fs	0/1	(136)
	c.349C > T	p.Arg117*	7/2	(136)(137)
	c.790A > T	p.Lys264*	2/0	(136)
	IVS3-1G > A	splice site	1/0	(136)
	c.858delA	p.Ala286fs*	1/0	(138)
	c.879T > G	p.Tyr293*	0/1	(136)
	c.912C > T	p.Arg305*	0/1	(136)
36	c.166C > A	p.Gln56Lys	1/0	(137)
77	c.281T > A	p.Ile94Asn	0/1	(139)
119	c.389G > A	p.Gly130Glu	0/1	(136)
123	c.400C > T	p.Arg134Cys	1/1	(135)(137)
123	c.401G > A	p.Arg134His	0/1	(136)
136	c.440T > A	p.Ile147Asn	1/0	(137)
146	c.469G > C	p.Gly157Arg	0/1	(139)
182	c.560G > C	p.Arg187Pro	0/3	(136)
182	c.560G > A	p.Arg187Gln	1/2	(135)(137)
185	c.569C > G	p.Pro190Arg	1/0	(137)
191	c.587T > C	p.Leu196Pro	1/0	(140)
204	c.625G > C	p.Asp209His	0/4	(136)(137)
220	c.683G > T	p.Gly228Val	4/0	(137)(138)
222	c.689T > C	p.Leu230Pro	1/0	(139)
276	c.832C > G	p.Arg278Gly	1/0	(137)
<b>SLC25A42, CoA and PAP carrier</b> <u>Position in Fig. 2</u>				
	c.380 +2T > A	splice site	1/0	(141)
276	c.871A > G	p.Asn291Asp	14/0	(141)(142)(143)

<b>SLC25A46</b>	1.897-KB DEL	missing start codon	2/0	(144)	
	c.42 > G	p.Tyr14*	0/2	(145)	
	c.165_166insC	p.His56fs*94	0/1	(146)	
	c.283+3G > T	p.Ser32Thrfs*	1/0	(147)	
	c.462+1G > A	intron	0/2	(145)	
	c.736A > T	p.Arg246*	2/0	(145)	
	c.882_885dupTTAC	p.Asn296fs*297	0/1	(146)	
	<b>Position in Fig. 2</b>				
	48	c.413T > G	p.Leu138Arg	2/0	(148)
	52	c.425C > T	p.Thr142Ile	1/0	(149)
	70	c.479G > C	p.Trp160Ser	1/0	(150)
	157	c.746G > A	p.Gly249Asp	0/1	(146)
	165	c.770G > A	p.Arg257Gln	1/0	(151)
	229	c.998C > T	p.Pro333Leu	0/1	(146)
231	c.1005A > T	p.Glu335Asp	3/0	(146)	
236	c.1018C > T	p.Arg340Cys	6/0	(146)(150)(152)	
237	c.1022T > C	p.Leu341Pro	2/0	(144)	

1) Mitochondrial DNA depletion syndrome 12B (cardiomyopathic type); 2) Mitochondrial DNA depletion syndrome 12A (cardiomyopathic type); 3) Autosomal dominant progressive external ophthalmoplegia with mitochondrial DNA deletions 2; 4) Mutations found on the same allele; NP, non-pathogenic.

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