Title: Mucolipidosis II *GeneReview* Table 2 Authors: Leroy JG, Cathey SS, Friez MJ

Updated: May 2012

Note: The following information is provided by the author(s) listed above and has not

been reviewed by *GeneReviews* staff.

Table 2. *GNPTAB* Mutation Summary Table

Exon	Mutation	Alternate Name	Protein	Reference
1	c.10A>C	174A>C	p.K4Q	Kudo [2006]
1	c.44C>A		p.S15Y	Cathey [2010]
2	c.118-2A>G		Splicing	Bargal [2006]
2	c.121delG		F.S.	Encarnacão [2009]
2	c.136C>T		p.R46X	Cathey [2010]
2	c.163_164insT		F.S.	Zarghooni [2009]
2	c.168T>A		p.Y56X	Cathey [2010]
2	c.171delA		F.S.	Cathey [2010]
2	Duplication exon 2		F.S.	Otomo [2009]
3	c.232delGTT		p.V78del	Cathey [2010]
3	c.242G>T		p.W81L	Encarnacão [2009]
3	c.310C>T		p.Q104X	Paik [2005]
4	c.342delCA		F.S.	Cathey [2010]
5	c.440delC		F.S.	Encarnacão [2009]
5	c.445delG		F.S.	Zarghooni [2009]
5	c.517insA		F.S.	Cathey [2010]
5	c.545T>A + c.614A>C		p.V182D + p.Q205P	Zarghooni [2009]
5	c.569A>T	733A>T	p.D190V	Kudo [2006]
5	c.571+3A>C		F.S.	Cathey [2010]
5	Alu insert		F.S.	Tappino [2008]
6	c.616del CAGA	773_776del CAGA	F.S.	Kudo [2006]
6	c.625delAGGGG		F.S.	Cathey [2010]
7	c.637-1G>A		Splicing	Cathey [2010]
7	c.648deIAGAA		F.S.	Cathey [2010]
7	c.749dupA		F.S.	Tappino [2009]
7	c.750+3A>C		Splicing	Zarghooni [2009]

Exon	Mutation	Alternate Name	Protein	Reference
7	c.755_759delCCTCT		F.S.	Tappino [2009]
7	c.771G>A		F.S. (Skipping of exon 7)	Steet [2005]
8	c.832C>T	996C>T	p.Q278X	Kudo [2006]
8	c.848delA	1012delA	F.S.	Kudo [2006]
8	c.857dupA		F.S.	Tappino [2009]
8	c.914_915insA		F.S.	Otomo [2009]
9	c.940C>T		p.Q314X	Tappino [2009]
9	c.1000C>T		p.R334X	Cathey [2010]
9	c.1001G>A		p.R334Q	Cathey [2010]
9	c.1001G>T		p.R334L	Otomo [2009]
9	c.1042A>C		p.l348L	Cathey [2010]
9	c.1090C>T		p.R364X	Cathey [2010]
10	c.1120T>C		p.F374L	Otomo [2009]
10	c.1123C>T		p.R375X	Cathey [2010]
10	c.1191_1194dupGCTG		F.S.	Tappino [2009]
10	c.1196C>T		p.S399F	Bargal [2006]
10	c.1206dupT		F.S.	Tappino [2009]
10	c.1208T>C		p.I403T	Tappino [2009]
10	c.1220A>C		p.D407A	Tiede [2005a]
11	c.1285-2A>G		Splicing	Cathey [2010]
11	c.1325G>A		p.C442Y	Tappino [2009]
11	c.1331dupG		F.S.	Tappino [2009]
11	c.1381T>G		p.C461G	Tappino [2009]
11	c.1385_1386insA		F.S.	Zarghooni [2009]
11	c.1399delG		F.S.	Cathey [2010]
11	c.1402T>A		p.C468S	Cathey [2010]
12	c.1514G>A		p.C505Y	Cathey [2010]
12	c.1519C>T		p.Q507X	Tappino [2009]
12	c.1580delC	1744delC	F.S.	Kudo [2006]
13	c.1625insC		F.S.	Tiede [2005b]
13	c.1738_1741 tripTATA	1902_1905 tripTATA	F.S.	Kudo [2006]
13	c.1759C>T		p.R587X	Cathey [2010]
13	c.1895C>G		p.A663G	Tiede [2005a]
13	c.1959_1962delTAGT		F.S.	Tappino [2009]

Exon	Mutation	Alternate Name	Protein	Reference
13	c.1965delC		F.S.	Tappino [2009]
13	c.1999G>T		p.E66X	Encarnacão [2009]
13	c.1999insT		F.S.	Cathey [2010]
13	c.2051_2055delACTCA	2215_2219delACTCA	F.S.	Kudo [2006]
13	c.2089_2090insC		F.S.	Otomo [2009]
13	c.2188T>AAA	2352T>AAA	F.S.	Kudo [2006]
13	c.2189delT		F.S.	Cathey [2010]
13	c.2196G>T		p.K732N	Zarghooni [2009]
13	c.2220_2221dupGA		F.S.	Tappino [2009]
13	c.2249_2250insA		F.S.	Encarnacão [2009]
13	c.2275delAA		F.S.	Cathey [2010]
13	c.2422delC		F.S.	Ma [2011]
13	c.2427delC		F.S.	Cathey [2010]
13	c.2533C>T		p.Q845X	Tiede [2005b]
13	c.2544delA		F.S.	Otomo [2009]
13	c.2550_2554delGAAAA		F.S.	Tappino [2009]
13	c.2574_2575delGA		F.S.	Paik [2005]
13	c.2591insG		F.S.	Cathey [2010]
13	c.2659dupA		F.S.	Tappino [2008]
13	c.2664C>G		p.Y888X	Cathey [2010]
13	c.2681G>A		p.W894X	Paik [2005]
13	c.2693insA		F.S.	Cathey [2010]
13	c.2693delA		F.S.	Otomo [2009]
13	c.2715+1G>A		Splicing	Paik [2005]
13	c.2715+2T>G		Splicing	Cathey [2010]
14	c.2777A>C		p.Q926P	Tappino [2009]
14	c.2783A>G + c.2864C>T		p.K928 + p.A955V	Zarghooni [2009]
14	c.2866C>T		p.H956Y	Otomo [2009]
14	c.2867A>G		p.H956R	Cathey [2010]
15	c.2916insT		F.S.	Bargal [2006]
15	c.3002T>C		p.L1001P	Tappino [2009]
15	c.3053A>G		p.D1018G	Cathey [2010]
15	c.3061C>T		p.Q1021X	Cathey [2010]

Exon	Mutation	Alternate Name	Protein	Reference
15	c.3091C>T		p.R1031X	Cathey [2010]
16	c.3145insC		F.S.	Tiede [2005b]
16	c.3160T>G		p.L1054V	Zarghooni [2009]
16	c.3173C>G		p.S1058X	Paik [2005]
16	c.3231_3234dupCTAC	3395_3398dupCTAC	F.S.	Kudo [2006]
16	c.3232delT		F.S.	Cathey [2010]
16	c.3249+1G>A		Splicing	Bargal [2006]
16	c.3249+1G>C		Splicing	Cathey [2010]
17	c.3252delA		F.S.	Tiede [2005b]
17	c.3310delG		F.S.	Otomo [2009]
17	c.3327insA		F.S.	Cathey [2010]
17	c.3335+1G>A		Splicing	Tiede [2005b]
17	c.3335+6T>G	Kudo IVS17+6T>G	Splicing	Kudo [2006]
18	c.3336-1G>C		Splicing	Cathey [2010]
18	c.3388_3389insC+c.3392C> T		Splicing	Otomo [2009]
18	c.3410T>A		p.L1137X	Cathey [2010]
18	c.3428_3429insA		F.S.	Otomo [2009]
18	c.3434+1G>A		Splicing	Bargal [2006]
19	c.3435-1G>A	IVS19-1G>A	Splicing	Kudo [2006]
19	c.3443_3446delTTTG		F.S.	Tappino [2009]
19	c.3458A>G		p.N1153S	Otomo [2009]
19	c.3474_3437delTA		F.S.	Paik [2005]
19	c.3487_3490delACAG		F.S.	Tappino [2009]
19	c.3503_3504delTC	3665_3666delTC	F.S.	Kudo [2006]
19	c.3523_3529delATGTTCC		F.S.	Tappino [2009]
19	c.3565C>T		p.R1189X	Paik [2005]
19	c.3566insA		F.S.	Tiede [2005b]
20	c.3613C>T		p.1205X	Tiede [2005b]
21	c.3741_3744delAGAA		F.S.	Otomo [2009]

^{1.} The alternate name column has been added to clarify the nomenclature of Kudo et al [2006]. The reference nucleotide sequence used by Kudo et al [2006] is GenBank AY687932 where the first nucleotide of the cDNA sequence is numbered 1. There are 164 nucleotides of untranslated sequence included in this numbering system. The mutation designations in the first column are numbered with the first nucleotide at the A of the ATG start codon.

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