

Bobilev et al.

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<i>FID</i>	<i>Person ID</i>	<i>Sex</i>	<i>Inheritance</i>	<i>Variant</i>	<i>Variant Sequence</i>
100	100A151/s-100	F	Sporadic	c.357+1G>A	CATACCAAGC/ <u>a</u> taagttcattg
101	101A111/s-101	F	Sporadic	c.112-116delCGGCC	GCGGGGCC(CGGCC)GTGCGACATTTCCGAATTCTGCAG/
102	102A127/s-102	F	Sporadic	c.607C>T	CAA ATG <u>T</u> GA CTT CAG
103	103A025/s-103	M	Sporadic	c.802_806delGAAGA	CGA AGG GCC AAA TGG AGA AGA (GAA GA)A AAA CTG
104	104A098/s-104	M	Sporadic	c.454C>T	AAC GGG <u>T</u> AG ACC GGA
105	105A046/u-105	F	Parents Unknown	c.357+1G>T	CATACCAAGC/ <u>t</u> taagttcattg
108	108A128/f-108	M	Familial	c.63-70delGCCGGACT	GGG CGG CCA CT(G CCG GAC T)CC ACC CGG CAG
110	110A005/s-110	F	Sporadic	c.794G>A	GCC AAA <u>T</u> AG AGA AGA
112	112A047/f-113	F	Familial	c.199A>T	TCC ATC <u>T</u> GA CCC AGG
112	112A065/f-112	F	Familial	c.199A>T	TCC ATC <u>T</u> GA CCC AGG
112	112A068/f-114	M	Familial	c.199A>T	TCC ATC <u>T</u> GA CCC AGG
112	112A147/f-128	F	Familial	c.199A>T	TCC ATC <u>T</u> GA CCC AGG
114	114A157/s-172	F	Sporadic	c.482delG	CGC CCT G(G)T TGG TAT
114	114A159/f-173	M	Familial	c.482delG	CGC CCT G(G)T TGG TAT
114	114A158/f-174	F	Familial	c.482delG	CGC CCT G(G)T TGG TAT
115	115A062/s-115	F	Sporadic	c.1268A>T	TTA CAG <u>T</u> TA] AAAAAAAAAAAAA
117	117A040/s-117	F	Sporadic	c.524-101_534del112 bp	ggtgaggctg(tcgggatataatgctcttgaggttaagacta caccagcccccttttgaggctccaagttaatccaaatttctctaccatcctattctttgtccag/ATGGCTGCCAG) CAACAGGAAGG
118	118A051/s-118	M	Sporadic	c.57delG	GTC AAC GGG CG(G) CCA CTG CCG
119	119A107/s-119	M	Sporadic	c.1183+1G>A	ACTTCAACAG/ <u>a</u> tgagccactgc
120	120A131/f-120	M	Familial	c.949C>T	TTG GGC <u>T</u> GA ACA GAC
122	122A023/s-122	F	Sporadic	c.1033-2A>G	ctctcctcac <u>g</u> /CCCCCAGTCC
122	122A221/f-197	F	Familial	c.1033-2A>G	ctctcctcac <u>g</u> /CCCCCAGTCC

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124	124A141/s-124	F	Sporadic	c.771delG	CAG GTA TG(G) TTT TCT AAT
126	126A092/s-126	M	Sporadic	c.467G>A	GGA AGC T <u>A</u> G GGC ACC
127	127A123/s-127	F	Sporadic	c.480delT	ACC CGC CC(T) GGT TGG TAT CCG
128	128A200/s-190	F	Sporadic	c.141+18_+20delGCC	AATTCTGCAG/gtgatcctccccggcgcc(gcc)cca c
129	129A024/s-129	F	Sporadic	c.357+1G>A	CATACCAAGC/ <u>a</u> taagttcattg
130	130A081/f-130	M	Familial	c.1032+6T>G	CTGCCTATGCAA/gtaagggcgcgct
130	130A135/f-170	F	Familial	c.1032+6T>G	CTGCCTATGCAA/gtaagggcgcgct
133	133A039/s-133	F	Sporadic	c.112delC	GGG GCC (C)GG CCG TGC
135	135A035/s-135	F	Sporadic	c.122_123insGCGG	CCG TGC GA(GCGG)C ATT TCC
138	138A028/s-138	M	Sporadic	c.141+1G>T	AATTCTGCAG/ <u>t</u> gatcctccc
140	140A195/s-185	M	Sporadic	c.1268A>T	TTA CAG T <u>T</u> A] AAAAAAAAAAAAA
141	141A009/s-141	F	Sporadic	c.401delA	GCT AGC GAA A(A)G CAA CAG ATG
141	141A010/f-140	M	Familial	c.401delA	GCT AGC GAA A(A)G CAA CAG ATG
143	143A077/s-143	F	Sporadic	c.1174_+6delACTTCAACAGG TGAGC	GGGCACCTCGGGCACC(ACTTCAACA G/gtgagc)cactg
144	144A083/f-144	F	Familial	c.781C>T	TCT AAT <u>T</u> G A AGG GCC
144	144A089/f-167	M	Familial	c.781C>T	TCT AAT <u>T</u> G A AGG GCC
144	144A300/f-224	M	Familial	c.781C>T	TCT AAT <u>T</u> G A AGG GCC
147	147A017/f-147	M	Familial	c.204delC	ATC AGA CC(C) AGG GCA
147	147A059/f-146	M	Familial	c.204delC	ATC AGA CC(C) AGG GCA
147	147A061/f-149	M	Familial	c.204delC	ATC AGA CC(C) AGG GCA
147	147A064/f-148	F	Familial	c.204delC	ATC AGA CC(C) AGG GCA
150	150A018/s-150	F	Sporadic	c.179- 185delATTACGAinsCTGAT	GGGCAGGT(CTGAT)GACTGGCTCCAT
150	150A106/f-151	M	Familial	c.179- 185delATTACGAinsCTGAT	GGGCAGGT(CTGAT)GACTGGCTCCAT
152	152A114/s-152	F	Sporadic	c.141+2_+30delTGATCCTCCC GGCGCCGCCCCACTCGCCG	AATTCTGCAG/g(tgatcctccccggcgccccca ctcgccg)cccccgcgccc
153	153A075/s-153	F	Sporadic	c.949C>T	TTG GGC <u>T</u> G A ACA GAC

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3	153	153A048/f-154	M	Familial	c.949C>T	TTG GGC <u>T</u> G A CA G AC
4	155	155A102/s-155	M	Sporadic	c.332insG	GAG GGG <u>G</u> GTC TGT ACC
5	157	157A121/s-157	F	Sporadic	c.28C>T	GTG AAT <u>T</u> AG CTC GGT
6	160	160A091/s-160	F	Sporadic	c.607C>T	CAA ATG <u>T</u> G A CTT CAG
7	163	163A041/s-163	F	Sporadic	c.352-357+2delCCAAGCGT	CATA(CCAAGC/gt)aagttcattg
8	164	164A060/s-164	F	Sporadic	c.781C>T	TCT AAT <u>T</u> G A AGG GCC
9	168	168A014/s-168	F	Sporadic	c.365C>A	GTG TCA <u>T</u> A A ATA AAC
10	169	169A080/s-169	F	Sporadic	c.358-3_361delCAGGTGT	tttgattg(cag/GTGT)CATCAA
11	170	170A174/s-184	F	Sporadic	c.781C>T	TCT AAT <u>T</u> G A AGG GCC
12	173	173A291/s-220	F	Sporadic	c.766-3C>G	cctgattc <u>g</u> ag/GTATGGTTTT
13	185	185A164/s-180	F	Sporadic	c.1256delC	TGG C(C)A AGA TTA CAG TAA
14	189	189A257/s-209	M	Sporadic	c.631C>T	AAG CTG <u>T</u> A A AGA AAT
15	191	191A297/s-222	F	Sporadic	c.799A>T	TGG AGA <u>T</u> G A GAA GAA
16	227	227A302/s-227	M	Sporadic	c.795G>A	GCC AAA <u>T</u> G A AGA AGA
17	230	230A309/s-230	M	Sporadic	c.357+1G>A	CATACCAAGC/ <u>a</u> taagttcattg
18	232	232A315/s-211	F	Sporadic	c.781C>T	TCT AAT <u>T</u> G A AGG GCC
19	1001	1001A316/sUGA	M	Sporadic	c.949C>T	TTG GGC <u>T</u> G A ACA GAC
20	1003	1003A318/sUGA	F	Sporadic	c.343delG	TGT ACC AAC (G)AT AAC ATA
21	1004	1004A321/sUGA	M	Sporadic	c.1183+1G>A	CAACAG/ <u>a</u> tgagcc
22	228	228A305/s-228	M	Sporadic	Deletion P1 promoter to exon 4	
23	1002	1002A317/sUGA	F	Sporadic	Deletion exons 6 and 7	
24	161	161A100/s-161	F	Sporadic	WT1-PAX6; 3' extent unknown	
25	174	174A185/sw-191	F	Sporadic	WT1-PAX6; 3' extent unknown	
26	180	180A182/sw-190	F	Sporadic	WT1-PAX6; 3' extent unknown	
27	182	182A239/sw-201	F	Sporadic	WT1-PAX6; 3' extent unknown	
28	214	214A285/s060158	M	Parents Unknown	WT1-PAX6; 3' extent unknown	
29	219	219A261/s121996	F	Sporadic	WT1-PAX6; 3' extent unknown	
30	225	225A280/s72581	F	Sporadic	WT1-PAX6; 3' extent unknown	
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3	501	501A320/sUGA	F	Sporadic	Deletion of PAX6
4	121	121A049/s-121	F	Sporadic	No changes detected
5	162	162A122/s-162	F	Sporadic	c.766-12C>T
6	166	166A069/s-166	F	Sporadic	c.766-12C>T
7	190	190A305/501F	F	Sporadic	c.766-12C>T
8	231	231A312/s-231	F	Sporadic	No changes detected
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