

Sup Table 1. Genotyping of flanked STR and SNP markers closely linked to the NM_002087.2(*GRN*):c.1179+104_1536delinsCTGA variant indicating a possible shared haplotype of a length of maximal ~14Mb.

GRCh37	Marker	Shared “haplotype”	Allel		Allel		Allel	
<i>Chromosome 17</i>			<i>Family 1, III:1</i>		<i>Family 2, III:1</i>		<i>Family 3, III:1</i>	
29705947	rs2525574	-	C	C	C	C	T	T
32366000	rs3744516	-	G	G	A	A	G	G
34951327	rs2285639	T	T	T	T	T	C	T
36491365	rs4795267	G	G	G	G	G	C	G
38346802	rs56304903	C	G	C	C	C	G	C
40459562	rs2272087	A	A	A	G	A	A	A
41753084	D17S1789	196	180	196	196	196	180	196
41820180	D17S951	205	212	205	205	205	205	205
42370474	D17S1860	194	190	194	188	194	194	194
42426940	rs9897526	G	G	G	G	G	A	G
42429267	c.1179+104_1536delinsCTGA	MUT	REF	MUT	REF	MUT	REF	MUT
42722874	rs7502041	G	A	G	A	G	G	G
42807066	D17S1861	101	101	101	107	101	89	101
43015415	D17S1804	232	237	232	230	232	232	232
43057745	D17S934	169	175	169	169	169	183	169
43260266	D17S950	177	192	177	177	177	179	177
44049329	rs75242405	C	C	C	C	C	T	C
46629593	rs2229304	T	G	T	T	T	T	T
47653804	rs2676791	-	T	T	A	A	T	T
48191818	rs77267183	-	C	C	A	A	A	C