

Gene	SNP	Gnomad AF	most_severe_consequence	Sift prediction	Polyphen	HGVSP	Gender	Age	Outcome
HECW1	chr7:43550491	<0.01	missense_variant	deleterious	possibly_damaging	T1432M	M	77	Survived
HECW1	chr7:43456393	<0.01	missense_variant	tolerated	benign	P866L	F	48	Survived
NEDD4	chr15:55916758	<0.01	missense_variant	deleterious_low_confidence	possibly_damaging	D25G	M	74	Survived
NEDD4	chr15:55833109	<0.01	splice_region_variant			c.3688-5T>C	N.A.	N.A.	N.A.
NEDD4	chr15:55951530	<0.01	missense_variant	tolerated	benign	Q60R	F	77	Survived
NEDD4L	chr18:58333891	0	missense_variant	tolerated	benign	P355L	M	63	Survived
NEDD4L	chr18:58349550	0	missense_variant	tolerated	benign	L530S	F	62	Survived
WWP2	chr16:69842059	<0.01	missense_variant	tolerated	benign	A172P	F	72	Survived
WWP2	chr16:69842128	<0.01	splice_region_variant			c.575+8C>G	M	24	Survived