



Sample: 16012132564

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a description goes here

Clinical note

No clinical note is added



Gene	undefined
Mode of inheritance	recessive
Consequence	stop gained
HGVSp	ENSP00000326018.6:p.Arg953Ter
HGVSc	ENST00000324001.7:c.2857C>T
rsld	rs104894714
gnomAD allele frequency	3.183e-05
ExAC allele frequency	
REVEL	
Impact	high
Location	chr19: 40901402-40901403
Variant type	snp
Variant quality	1933.95

■ Notes:

— 2019-8-30 15:16 Tonya Lee Disera

asdf

Genes

Phenotypes searched	Genes (100)			
GTR: • undefined	MPZ, EGR2, PMP22, PRX, GDAP1, FGD4, DNM2, HSPB1, FIG4, DYNC1H1, NEFL, MFN2, LMNA, MTMR2, NDRG1, SH3TC2, SBF2, RAB7A, LITAF,			
Phenolyzer: • undefined	HSPB8, TRPV4, MED25, LRSAM1, BSCL2, SBF1, SPG11, HK1, PEX7, PHYH, SPAST, GAN, GJB1, AARS1, GARS1, GNB4, YARS1, AIFM1, COX6A1, PRPS1, PDK3, DHTKD1, PLEKHG5, KARS1, KIF1B, TRIM2, INF2, C12orf65, ATP1A1, HADHB, IGHMBP2, TFG, GLA, HINT1, DNAJB2, MARS1, HARS1, KIF5A, TTR, CTDP1, FBLN5, SPTLC1, MORC2, DNMT1, HOXD10, SCN9A, SPTLC2, SLC12A6, ATL1, REEP1, TSEN54, ALMS1, RARS2, FA2H, L1CAM, SEPT9, SMCHD1, PLP1, ALS2, DUX4, ETS1, SPG7, FGF3, ISCU, MARS2, GARS, TSEN2, TSEN34, VRK1, GRN, SPG20, KIAA0196, AARS, EXOSC3, OPA3, GATA2, SEPSECS, FRG1, PRKCA, YBX1, AKT1			
HPO:				

Disclaimers

Genetic testing information has caveats and should not be considered a definitive diagnosis.

References/Methodology

DNA sequencing was performed in accordance with established Utah Genome Project (UGP) methodologies including sample preparation, sequencing and data analysis.