

Variant	TP53(NM_000546.6):c.1012T>C, p.Phe338Leu	
Frequency		
gnomAD Genomes	not found	
gnomAD exomes	not found	
Conservation		
Scores	Value	
PhastCons100way	0.897	
PhyloP100way	8.183	
Pathogenicity Scores		
Meta scores		
Engine	Calibrated Prediction	Score
BayesDel addAF	Pathogenic Moderate	0.2279
MetaLR	Pathogenic Supporting	0.8532
MetaRNN	Pathogenic Supporting	0.7701
MetaSVM	Pathogenic Supporting	0.8849
BayesDel noAF	Uncertain	0.08959
REVEL	Uncertain	0.681
Individual Predictions		
DEOGEN2	Pathogenic Moderate	0.9628
EVE	Pathogenic Supporting	0.6595
FATHMM-MKL	Pathogenic Supporting	0.9935
FATHMM-XF	Pathogenic Supporting	0.9229
LIST-S2	Pathogenic Supporting	0.9725
M-CAP	Pathogenic Supporting	0.4196
MutPred	Pathogenic Supporting	0.706
LRT	Benign Supporting	0.005234
BLOSUM	Uncertain	0
CADD	Uncertain	25.3999
DANN	Uncertain	0.9961
EIGEN	Uncertain	0.3318
FATHMM	Uncertain	-3.47
Mutation assessor	Uncertain	2.77
MutationTaster	Uncertain	0.9985
MVP	Uncertain	0.9348
Polyphen2 HDIV	Uncertain	0.787
Polyphen2 HVAR	Uncertain	0.595
PrimateAI	Uncertain	0.5514
PROVEAN	Uncertain	-3.77
SIFT	Uncertain	0.002
SIFT4G	Uncertain	0.005

[illegible]