Table S1. Prediction of variant pathogenicity by in silico tools.

Tool	Score	Interpretation	Confidence Level
SIFT	0	Pathogenic	High
Mutation Taster	1	Uncertain	Moderate
FATHMM	−6.99	Pathogenic	Moderate
DANN	0.9967	Uncertain	Moderate
M-CAP	0.9939	Pathogenic	Moderate
REVEL	0.96	Pathogenic	Strong