UGT8

215 missense variants predicted as deleterious by at least one method

MAF of missense variants predicted as deleterious

Variant ID (MAF>0.5)
■ 4−115589302−A−G

MutPred	CADD	PROVEAN	DANN	MutationAssessor	SIFT	PolyPhen−2 HDIV	ClinPred	PolyPhen−2 HVAR	FATHMM−MKL	ADME	LRT	Eigen−PC	VEST	M−CAP	MetaLR	MetaSVM	MVP	REVEL	FATHMM	PrimateAI
126	192	60	181	80	103	83	104	82	194	109	138	141	93	88	10	12	52	25	0	13