

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0 Q99KR7	PPIF_MOUSE	Mus musculus	Peptidyl-prolyl cis-trans isomerase F, mitochondrial	24.088486	NaN	NaN	37507081;36288343	MLALRCGPRLGLLSGPRSA PLLLSA TRTCS DGGARGANSSSGNPLVYLDV GADGQPLGRVVLELKADVVPKTAEN FRALCTGEKGFYKGS TFHRVIPAF MCQAGDFTNHNGTGGRSIYGS RFP DENFTLKHVGPVLSMANAGPNTN GSQFFICTIKTDWLDGKHVVFVGHVK EGMDVVKKIESFGSKSGKTSKKIVIT DCGQLS	True	False	2.697	2.36	4.84	2.423	1.395	2.379	2.16