

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
P12686	RT13_YEAST	Saccharomyces cerevisiae	37S ribosomal protein MRP13, mitochondrial	24.475439	NaN	NaN	33229814	MGTTTVVINEGPILLIRALHRATTNK KMFRSTVWRRFASTGEIAKAKLDEF LIYHKTDAKLKPFYRPKNAQILLTKD IRDPKTREPLQPRPPVKPLSKQTLND FIYSVEPNSTELLDWFKEWGTGSIRK RAIWYISPIHVQKMLTASFCKIGKYA HMGVLLYGIEHKFLKAQNPSVFDIE HFFNTNIMCALHRNRLKDYKDAEIA QRKLQVAWKVLRNKNNTGLANIL VATLGRQIGFTPPELTGLQPVDISLPDI PNSSSGAELKDLSKYEGYLIARTLL DIDQHNAQYLELQEFIRQYNALSE SSDPYDTHLKALGLETTPPPQESTEK EEK	None	None	None	None	None	None	None	None	None

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