

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	e
Q10713	MPPA_HUMAN	Homo sapiens	Mitochondrial-processing peptidase subunit alpha	10.675769	T467	NaN	30379171;28510447;34725712;38665916	MAAVVLAATRLLRSGSWGCSRLRF GPPAYRRFSSGGAYPNILSSPLPGV PKPVFATVDGQEKFETKVTTLDNGL RVASQNKFGQFCTVGLINSGSRYEA KYLSGIAHFLEKLAFSSTARFDSKDE ILLTLEKHGGICDCQTSRDTTMYAVS ADSKGLDTVVALLADVVLQPRLTDE EVEMTRMAVQFELEDLNLRPDPEPL LTEMIEAAYRENTVGLHRFCPTEN VAKINREVLHSYLRNYTPDRMVLA GVGVEHEHLVDCARKYLLGVQPAW GSAEAVDIDRSVAQYTTGGIAKLERD MSNVSLGPTPIPELTHIMVGLESCSF LEEDFIPFVFLNMMMGGSFSAG GPGKGMFSRLYLNVLNRHHWMYN ATSYHHSYEDTGLLCIHASADPRQV REMVIEITKEFILMGGTVDTVELERA KTQLTSMMLMNLESRPVIFEDVGR QVLATRSRKLPHLCTLRNVKPEDV KRVASKMLRGKPAVAALGDLTDLPT YEHQ TALSSKDGRLPRTYRLFR	True	False	2.399	1.627	5.0	1.162	0.633	1.223	4