

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	extracell region
Q14703	MBTP1_HUMAN	Homo sapiens	Membrane-bound transcription factor site-1 protease	24.902576	S574;S589;T875	S168	40596516;38253038	MKLVNIWLLLLLVLLCGKKHLGDRL EKKSFEKAPCPGCSHLLTKVEFSSTV VEYEVIVAFNGYFTAKARNSFISSAL KSSEVDNWRRIIPRNNPSSDYPSDFE VIQIKEKQKAGLLTLEDHPNIKRVTP QRKVFRLSLKYAESDPTVPCNETRWS QKWQSSRPLRRASLSLGSQGFWHAT GRHSSRRLRLRAIPRQVAOTLQADV WQMGYTGANVRVAVFDTGLSEKHP HFKNVKERTNWTNERTLDDGLGHG TFVAGVIASMRECOGFAPDAELHIFR VFTNNQVSYTSWFLDAFYAILKKID VLNLSIGGPDFMDHPFVDKVVWELTA NNVIMVSAIGNDGPLYGLTNNPADQ MDVICVGGIDFEDNIARFSSRGM WELPGGYGRMKPDIVTYGAGVRGSG VKGGCRALSGTSVASPVVAGAVILLV STVQKRELVNPASMKQALIASARRL PGVNMFEQGHGKLDLLRAYQILNSY KPQASLSPSYIDLTECPYMWPYCSQP IYYGGMPTVVNVTILNGMGVTRIV DKPDWQPYPQNGDNIEVAFSYSSV LWPWSGYLAISVTKKAASWEGIAQ GHVMTVASPAETESKNGAEQTSTV KLPIKVKIIPTPRSKRVLWDQYHNL RYPFGYFRDNLRMKNDPLDWN HIHTNFRDMYQHLSMGGYFVEVLG APFTCFDASQYGTLLMVDSEEEYFP EEIAKLRRVDNGLSLVIFSDWYNTS VMRKVKFYDENTRQWWMPDTGGA NIPALNELLSVWNMGFSDGLYE TLANHDMMYYASGCSIAKFPEDGVVIT QTFKDQGLEVLKQETAVVENVPILG LYQIPAEGGGRIVLYGDSNCLDDSH RQKDCFWLLDALLQYTSYGVTPPSL SHSGNRQRPPSGAGSVTPERMEN HLHRYSKVLEAHLGDPKPRPLPAC RLSWAKQPLNETAPSNLWKHQKL LSIDLKVVLPNFRSNRPQVRPLSPG ESGAWDIPGGIMPGRYNQEVEGQ VFAFLGAMVLAFFVQINKAKSRP KRRKPRVKRQPLMQQVHPPKTPSV	False	False	False	False	False	4.131	4.0	1.969	1.172