

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
Q14703	MBTP1_HUMAN	Homo sapiens	Membrane-bound transcription factor site-1 protease	22.903221	S574;S589;T875	S168	38253038	MKLVNIWLLLLVLLCGKKHLGDRL EKKSFEKAPCGCSHLTLKVEFSSTV VEYEVAFNGYFTAKARNSFISAL KSSEVDNWRRIIPRNNPSSDYPSDFE VIQIKEKQKAGLLTLEDHPNKRKRVTP QRKVFRLSKYAESDPTVPCNETRWS QKWQSSRPLRRASLSLGGFWHAT GRHSSRLLRAIPRQVAQTLQADVL WQMGYTGANVRVAVFDTGLSEKHP HFKNVKERTNWTNERTLDDGLGHG TFVAGVIASMRECCGFAPDAELHIFR VFTNNQVSYTSWFLDAFNAILKKID VLNLSIGGPDFMDHPFVDKVVWELTA NNVIMVSAIGNDGPLYGTLNPNADQ MDVIGVGGIDFEDNIARFSSRGMTT WELPGGYGRMKPDIVTYGAGVRGSG VKGGCRALSGTASVSPVAVGAVTLV STVQKRELVPNASMKQALIASARRL PCVNMFEQGHGKLDLLRAYQILNSY KPQASLSPSYDLTECPYMWYPCSQP IYYGGMPTVVNVTILNGMGVTGRIV DKPDWQPYLPQNGDNIEVAFYSYSSV LWPWSGYLAISISVTKKAASWEGIAQ GHVMITVASPAETESKNGAEQTSTV KLPIKVKIIPPRSKRVLWDQYHNL RYPPGYFPRDNLRMKNPDLWDNGD HIHTNFRDMYQHLRSMGYFVEVLG APFTCFDASQYGTLLMVDSEEEYFP EEIAKLRRVDNGLSLVIFSDWYNTS VMRKVKFYDENTRQWWMPDTGGA NIPALNELLSVWNMGFSDGLYEGEF TLANHDMYYASGCCIAKFPEDGVVIT QTFKDQGLEVLKQETAVVENVPILG LYQIPAEGGGRIVLYGDSNCLDDSH ROKDCFWLLDALLQYTSYGVTPPSL SHSGNRQRPPSGAGSVTPERMEN HLHRYSKVLEAHLGDPKPRPLPACP RLSWAKQPLNETAPSNLWKHQKL LSIDLKVVLPNFRSNRPQVRPLSPG ESGAWDIPGGIMPGRYNQEVGQTIP VFAFLGAMVLAFFVVQINKAKSRP KRRKPRVKRPQLMQQVHPPKTPSV	False	True	2.231	2.352	1.805	4.62	4.574	3.211	2.696