

UniprotKB ID	Entry name	organism	full name	oglcnaScore	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi appara
Q3KQU3	MA7D1_HUMAN	Homo sapiens	MAP7 domain-containing protein 1	10.051975	S460;T838	T47;T51;S70;S86;S93;T97;S113;S116;T118;S123;S125;S254;S273;S313;S366;S399;S442;S446;S452;S454;S460;S479;S496;S544;S548;S552;T554;S742;S753;T813;T816;S834	38665916;34019948;28657654;30379171;31373491;23301498;37217939;40596516;35254053	MESGPRAELGAGAPPAVVARTPPEP RPSPEGDPSPPPPMSALVDPDTPPDT PPAMKNATSSKQLPLEPESPSGQVG PRPAPPQEESSSEAKSRGPTTPAM GPRDARPPRRSSQPSPTAVPASDSPP TKQEVKKAGERHKLAKERREERAKY LAAKAVVWLEKEEKAKALREKQLQE RRRLLEQRLKAEQRRRAALEERORO KLEKNKERYEAAIORSVKKTWAEIR QQRWSWAGALHHSSPGHKTSGSRC SVSAVNLPKHVDSIINKRLSKSSATL WNSPSRRNSLQLSAWESSIVDRLMT PTLSFLARSRSAVTLPRNGRDQGRG CDPGRGPTWGRAGASLARGQPDR THPSAAVPVCPRSASASPLTPCSVTR SVHRCAPAGERGERRKPNAGGSPAP VRRRPEASPVQKKEKDKERENEKE KSALARERSLKKRQSLPASPRARLSA STASELSPKSKARPPSPSTSWHRPAS PCPSPGPGHTLPPKPPSPRGTTASPK GRVRRKEEAKESPSAAGPEDKSQSK RRASNEKESAAPASPAPSPAPSPTPA PPQKEQPPAETPTDAAVLTSPPAPAP PVTPSKPMAGTTDREEAIRLLAEKR RQAREQREEREQERRLQAERDKRM REEQLAREAEARAEREAEARRREEQ EAREKAQAEQEEQERLQKQKEEAEA RSREAEERQRLEREKHFQQEQER QERRKRLEIIMKRTRKSEVSETKQK QDSKEANANGSSPEPVKAVEARSPG LQKEAVQKEEPIQEPQWLSLPSKEL PASLVNGLQPLPAHQENGFSTNGPS GDKLSRTPETLLPFAEAEAFLLKAV VQSPQVTEVL	False	False	False	2.812	False	False	False