

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q3KQU3	MA7D1_HUMAN	Homo sapiens	MAP7 domain-containing protein 1	12.366532	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	23301498;30379171;31373491;35254053;28657654;34019948	MESGPRAELGAGAPPAVVARTPPPEP RPSPEGDPSPPPPPMSALVPDTPPDT PPAMKNATSSKQLPLEPESPSGQVG PRPAPPQEEESPSSEAKSRGTPPAM GPRDARPPRRSSQPSPTAVPASDSPP TKQEVKKAGERHKLAKERREERAKY LAAKKAVWLEKEEKAKALREKQLQE RRRRLEEQRLKAEQORRAALEERQEQ KLEKNKERYEAAIQRSVKKTWAEIR QQRWSWAGALHHSSPGHKTSGSRC SVSAVNLPKHVDSIINKRLSKSSATL WNSPSRNRSLQLSAWESSIVDRLMT PTLSFLARSRAVTLPRNGRDQGRG CDPGRGPTWGRAGASLARGPQPDR THPSAAVPVPCPRASASPLTPCSVTR SVHRCAPAGERGERRKPNAGGSPAP VRRRPEASPVQKKEKKDKERENEKE KSALARERSLKKRQSLPASPRARLSA STASELSPKSKARPSSTSWHRPAS PCPSPGPGHTLPPKPPSPRGTTASPK GRVRRKEEAKESPSAAGPEDKSQSK RRASNEKESAAPASPAPSPAPSPTPA PPQKEQPPAETPTDAAVLTSPAPAP PVTPSKPMAGTTDREEARLLAEKR RQAREQREEREQERRLQAERDKRM REEQLAREAEARAEREAEARRREEQ EAREKAQAEQEEQERLQKQKEEAEA RSREEAERQLEREREKHFQQEQER QERRKRLEEIMKRTRKSEVSETKQK QDSKEANANGSSPEPVKAVEARSPG LQKEAVQKEEPIQPQPQWLSKEL PASLVNGLQPLPAHQENGFSTNGPS GDKLSRTPETLLPFAEAEAFLLKAV VQSPQVTEVL