

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q92585	MAML1_HUMAN	Homo sapiens	Mastermind-like protein 1	16.556554	S321;T325;S331;S335;T350;S571;T595;S794;S801;T965;S975	S45;S120;S303;S314;S360;S1015	33214551;35254053;38253038;35132862;31492838;35138101;34019948;23301498;30059200	MVLPTCPMAEFALPRHSVAVMERLR RRIELCRRHHSTCEARYEAVSPERLE LERQHTFALHQRCIQAKAKRAGKHR QPPAATAPAPAAPAPRLDAADGPEH GRPATHLHDTVKRNLDSATSPQNG DQQNGYGDLFPGHKKTRREAPLGV AISSNGLPPASPLGQSDKPSGADALQ SSGKHSGLDSLNNKRLADSSSLHLN GGSNPSESFPLSLNKELKQEPVEDL PCMITGTVGSISQSNLMPDLNLNEQ EWKELIEELNRSVPDEDMKDLFNE DFEEKKDPESSGSATQTPLAQDINIK TEFSPAAFEQEQLGSPQVRAGSAGQ TFLGPSSAPVSTDSPSLGGSOTLFHT SGQPRADNPSPNLMPASQAQNAQ RALAGVVLPSQGGGASELSSAHQL QQIAAKQKREQMLQNPQATPAPAP QGMSTWQQTGPSHSSLDVVPYPMK PASPSYKQDFTNSKLLMMPSVNKS SPRPGPYLQPSHVNLLSHQPPSNL NQNSANNQGSVLDYGNTKPLSHYK ADCGQSGSPGSGQSKPALMAYLPQQL SHISHEQNSLFLMKPKPGNMPFRSL VPPGQEQNPSSVPVQAQATSVGTQP PAVSVASSHNSSPYLSSQQQAAMK QHQLLDQKQREQQQKHLQQQQF LQRQHLLAEQEQQFQRHLTRPPP QYQDPTQGSFPQVQVQFTGSSAAMP GMNTLGPSNSSCPRVFPQAGNLM MGPGHASVSSLPTNSGQDRGVAQ FPGSQNMPQSSLYGMASGITQIVAQ PPPQATNGHAHIPRQTNVGNQTSVS AAYGQNSLGSGLSQHNKGTLP GLTKPPVPRVSPAMGGQNSSWQH GMPNLSSGQTPGNSNVSPFTAASSFH MQQQAHLKMSSPQFSQAVPNRPMA PMSSAAVGSLLPPVSAQRTSAPA PAPPPTAPQQGLPGLSPAGPELGAFS QSPASQMGGRAGLHCTQAYPVRTA GQELPFAYSGQPGGSLSSVAGHTD LIDSLKNTSEEWMSDLDDLGSQ