

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q92585	MAML1_HUMAN	Homo sapiens	Mastermind-like protein 1	14.005457	S321;T325;S331;S335;T350;T595;S794;S801;T965;S975	S45;S120;S303;S314;S360;S1015	31492838;34019948;35254053;35132862;35138101;30059200;23301498;33214551	MVLPTCPMAEFALPRHSVAVMERLR RRIELCRRHHSTCEARYEAVSPERLE LERQHTFALHQRCIQAKAKRAGKHR QPPAATAPAPAAPAPRLDAADGPEH GRPATHLHDTVKRNLDSATSPQNG DQQNGYGDLFPGHKKTRREAPLGV AISSNGLPPASPLGQSDKPSGADALQ SSGKHSGLDSLNNKRLADSSSLHLN GGSNPSESFPLSLNKELKQEPVEDL PCMITGTVGSISQSNLMPDLNLNEQ EWKELIEELNRSVPDEDMKDLFNE DFEEKKDPESSGSATQTPLAQDINIK TEFSPAAFEQEQLGSPQVRAGSAGQ TFLGPSSAPVSTDSPSLGGSOTLFHT SGQPRADNPSPNLMPASQAQNAQ RALAGVVLPSQGGGASELSSAHQL QQIAAKQKREQMLQNPQATPAPAP QGMSTWQQTGPSHSSLDVVPYPMK PASPSYKQDFTNSKLLMMPSVNKS SPRPGGPYLQPSHVNLLSHQPPSNL NQNSANNQGSVLDYGNTKPLSHYK ADCGQSGSPGSGQSKPALMAYLPQQL SHISHEQNSLFLMKPKPGNMPFRSL VPPGQEQNPSSVPVQAQATSVGTQP PAVSVASSHNSSPYLSSQQQAAMK QHQLLDQKQREQQQKHLQQQQF LQRQHLLAEQEKQQFQRHLTRPPP QYQDPTQGSFPQVQVQFTGSSAAMP GMNTLGPSNSSCPRVFPQAGNLM MGPGHASVSSLPTNSGQDRGVAQ FPGSQNMPQSSLYGMASGITQIVAQ PPPQATNGHAHIPRQTNVGNQTSVS AAYGQNSLGSSGLSQHNKGTLP GLTKPPVPRVSPAMGGQNSSWQH GMPNLSSGQTPGNSNVSPFTAASSFH MQQQAHLKMSSPQFSQAVPNRPMA PMSSAAAVGSLLPVSAQRTSAPA PAPPPTAPQQGLPGLSPAGPELGAFS QSPASQMGGRAGLHCTQAYPVRTA GQELPFAYSGQPGGSLSSVAGHTD LIDSLKNTSEEWMSDLDDLGSQ