

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P27546	MAP4_MOUSE	Mus musculus	Microtubule-associated protein 4	29.461493	S887	S5;S60;S99;S254;T260;T277;T349;S381;T410;T447;S475;T494;T503;S506;T511;S512;S517;S519;S598;S617;T658;S667;S684;S694;S760;S798;S826;S901;S914;T915;S973;S1046;S1118;S1124;S333;S334	22645316;22517741;28135057;25153642;29187734	MADLSLVDALTEPPPEIEGEIKRDFM AALEAEPYDDIVGETVEKTEFIPLLD GDEKTGNSESKKKPCLDTSQVEGIP SSKPTLLANGDHGMEGNNTAGSPT DFLEERVDPDYQSSQNWPEDASF FQPQQVLDTDQAEFPNEHRDDGLA DLLFVSSGPTNASAFTERDNPSEDS YGMLPCDSFASTAVVSQEWSVGAPN SPCSESCVSPEVTIETLQPATELSKA AEVESVKEQLPAKALETMAEQTTDV VHSPSTDTPGPDTEAALAKDIEEIT KPDVILANVTQPSTESDMFLAQDME LLTGTEAAHANNIILPTEPDESSTKD VAPPMEEEEIVPGNDTTPKETETTL IKMDLAPPEDVLLTKETELAPAKGM VSLSEIEEALAKNDESSAEIPVAQET VVSETEVVLATEVVLPSDPITTLTKDV TLPLEAERPLVTDMTPSLETEMTLG KETAPPTETNLGMAKDMSPLESEV TLGKDVVILPETKVAEFNNTPLSEE EVTSVKDMSPSAETEAPLAKNADLH SGTELIVDNSMAPASDLALPLETKVA TVPIKDKGTVQTEEKPREDSQLASM QHKGQSTVPPCTASPEPVKAAEQMS TLPIDAPSPLENLEQKETPGSQPSEP CSGVSQRQEEAKAAVGVGTGNDITPP NKEPPPSPEKKAKPLATTQPAKTSTS KAKTQPTSLPKQPAPTTSGGLNKKP MSLASGSVPAAPHKRPAATATARP STLPARDVKPKPITEAKVAEKRTSPS KPSSAPALKPGPKTTPVSKATSPST LVSTGPSSRSPATTLPKRPTSİKTEGK PADVKRMTAKSASADLSRSKTTTSAS SVKRNTTPTGAAPPAGMTSTRVKPM SAPSRSSGALSVDKKPTSTKPSSSAP RVSRLATTVSAPDLKSVRSKVGSTEN IKHQPGGGRAKVEKKTEAATTAGKP EPNAVTKAAGSIASAQKPPAGKVQIV SKKVSYSHIQSKCGSKDNIKHVPGG GNVQIQNKKVDISKVSSKCGSKANIK HKPGGGDVKIESQKLNFKKAQAKV GSLDNVGHLPAGGAVKTEGGGSEAL PCPGPPAGEEPVIPEAAPDAGAP TSA SGLSGHTTLLSGGGDQREPQTLD S Q I QETSI