

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|------------|--------------|----------------------------------|--------------|---------------|--|---|---|
| P27546 | MAP4_MOUSE | Mus musculus | Microtubule-associated protein 4 | 29.004026 | 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 | 28135057;22517741;36288343;22645316;25153642;29187734 | MADLSLVDALTEPPPEIEGEIKRDFM AALEAEPYDDIVGETVEKTEFIPLLD GDEKTGNSESKKKPCLDTSQVEGIP SSKPTLLANGDHGMEGNNTAGSPT DFLEERVDPDYQSSQNWPEDASF FQPQQVLDTDQAEFPNEHRDDGLA DLLFVSSGPTNASAFTERDNPSEDS YGMLPCDSFASTAVVSQEWSVGAPN SPCSESCVSPEVTIETLQPATELSKA AEVESVKEQLPAKALETMAEQTTDV VHSPSTDTTPGPDTEAALAKDIEEIT KPDVILANVTQPSTESDMFLAQDME LLTGTEAAHANNIILPTEPDESSTKD VAPPMEEEEIVPGNDTTSPKETETTL IKMDLAPPEDVLLTKETELAPAKGM VSLSEIEEALAKNDESSAEIPVAQET VVSETEVVLATEVVLPSDPITTLTKDV TLPLEAERPLVTDMTPSLETEMTLG KETAPPTETNLGMAKDMSPLESEV TLGKDVVILPETKVAEFNNVTPLSEE EVTSVKDMSPSAETEAPLAKNADLH SGTELIVDNSMAPASDLALPLETKVA TVPIKDKGTVQTEEKPREDSQLASM QHKGQSTVPPCTASPEPVKAAEQMS TLPIDAPSPLENLEQKETPGSQPSEP CSGVSQRQEEAKAAVGVGTGNDITPP NKEPPPSPEKKAKPLATTQPAKTSTS KAKTQPTSLPKQPAPTTSGGLNKKP MSLASGSVPAAPHKRPAATATARP STLPARDVKPKPITEAKVAEKRTSPS KPSSAPALKPGPKTTPVSKATSPST LVSTGPSSRSPATTLPKRPTSİKTEGK PADVKRMTAKSASADLSRSKTTTSAS SVKRNTTPTGAAPPAGMTSTRVKPM SAPSRSSGALSVDKKPTSTKPSSSAP RVSRLATTVSAPDLKSVRSKVGSTEN IKHQPGGGRAKVEKKTEAATTAGKP EPNAVTKAAGSIASAQKPPAGKVQIV SKKVSYSHIQSKCGSKDNIHVPGG GNVQIQNKKVDISKVSSKCGSKANIK HKPGGGDVKIESQKLNFKKAQAKV GSLDNVGHLPAGGAVKTEGGGSEAL PCPGPPAGEEPVIPEAAPDAGAP TSA SGLSGHTTLLSGGGDQREPQTLD S Q I QETSI |