

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P27816	MAP4_HUMAN	Homo sapiens	Microtubule-associated protein 4	56.923246	S329;T349;S358;S580;T707;S742;S760;S787;S793;S797;T810;T811;S853;T864;S914;S921;S928;T966;T975	S5;S14;S60;S99;S253;S280;T282;T354;S358;T380;S384;S440;T442;S507;S510;T521;T526;T571;S580;T585;S624;S636;S643;T687;S696;S713;S723;S787;S797;S825;S827;S853;S928;S941;T942;S1000;S1073;S1145;S1151;S337;S338;T28;S269;S803	30379171;33214551;27655845;31492838;32119511;30620550;16408927;20068230;31637018;30059200;26853435;33465208;28657654;29351928;31373491;29237092;21158410;34019948;34105348;3626323;20305658;22661428;32574038;34065225;32870666;23301498;28510447	MADLSLADALTEPSPDIEGEIKRDFI ATLEAEAFDDVVGETVVGKTDYIPLLD VDEKGTGNSESKKKPCSETSQIEDTPS SKPTLLANGGHGVEGSDTTGSPTEF LEEKMAYQEYPNSQNWPEDTNFCF QPEQVVDPIQTDPPFKMYHDDDLADL VFPSSATADTSIFAGQNDPLKDSYG MSPCNTAVVPQGSVEALNSPHSE SFVSPPEVAEPPQPTAVPLELAKEIE MASEERPPAQALEIMMGLKTTDMA PSKETEMALAKDMALATKTEVALAK DME SPTKLDVTLAKDMQPSMESDM ALVKDMELPTEKEVALVKDVRWPTE TDVSSAKNVVLPTETEVAPAKDVTL KETERASPIKMDLAPSKDMGPPKEN KKETERASPIKMDLAPSKDMGPPKE NKIVPAKDLVLLSEIEVAQANDIISST EISSAEKVALSSETEVALARDMTLPP ETNVILTKDKALPLEAEVAPVKDMA QLPETEIAPAKDVAPSTVKEVGLLD MSPLSETEMALGKDVTTPPETEVVLI KNVCLPPEMEVALTEDQVPALKTEA PLAKDGLVTLANNVTPAKDVPPLSET EATPVPIKDMEIAQTQKGISEDSHLE SLQDVGQSAAPTFFMISPETVTGTGK KCSLPAEEDSVLEKLGGERKPCNSQP SELSSETSGIARPEEGRPVVSGTGND ITTPPNKELPPSPEKKTPLATTQPA KTSTSKAKTQPTSLPKQPAPTIGGL NKKPMSLASGLVPAAPPKRPVAVASA RPSILPSKDVKPKPIADAKAPEKRASP SKPASAPASRSGSKSTQTVAKTTTAA AVASTGPSSRSPSTLLPKKPTAIKTEG KPAEVKMTAKSVPADLSRPKSTST SSMKKTTTTLSGTAPAAGVVPSRVKAT PMPSPSTTPPFIDKKPTSAKPSSTTP RLSRLATNTSAPDLKNVRSKVGSTE NIKHQPGGGRAKVEKTEAAATTRK PESNAVTKTAGPIASAQKQPAGKVQI VSKKVSYSHIQSKCGSKDNIKHVPG GGNVQIQNKKVDISKVSSKCGSKAN IKHKPGGGDVKIESQKLNFKKAQA KVGSLDNVGHLPAGGAVKTEGGGSE APLCPGPPAGEEPAISEAAPEAGAPT SASGLNGHPTLSGGGDQREAOQLDS QIQETSI