

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P27816	MAP4_HUMAN	Homo sapiens	Microtubule-associated protein 4	53.858126	S329;T349;S358;S580;S659;T681;T707;S742;S760;S787;S789;S793;S797;T804;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	23301498;34105348;31373491;22661428;29351928;33214551;26853435;30059200;21158410;30620550;29237092;34065225;35254053;28657654;31637018;32870666;20305658;34725712;20068230;32574038;30379171;16408927;31492838;34846842;33465208;32119511;35138101;28510447;27655845;35132862;34019948	MADLSLADALTEPSPDIEGEIKRDFI ATLEAEAFDDVVGETVVGKTDYIPLLD VDEKGTGNSESKKKPCSETSQIEDTPS SKPTLLANGGHGVEGSDTTGSPTEF LEEKMAYQEYPNSQNWPEDTNFCF QPEQVVDPIQTDPPFKMYHDDDLADL VFPSSATADTTSIFAGQNDPLKDSYG MSPCNTAVVPQGSVEALNSPHSE SFVSPAEVAEPPQPTAVPLELAKEIE MASEERPPAQALEIMMGLKTTDMA PSKETEMALAKDMALATKTEVALAK DMESPTKLDVTLAKDMQPSMESDM ALVKDMELPTEKEVALVKDVRWPTE TDVSSAKNVVLPTETEVAPAKDVTL KETERASPIKMDLAPSKDMGPPKEN KKETERASPIKMDLAPSKDMGPPKE NKIVPAKDLVLLSEIEVAQANDIISST EISSAEKVALSSETEVALARDMTLPP ETNVILTKDKALPLEAEVAPVKDMA QLPETEIAPAKDVAPSTVKEVGLLD MSPLSETEMALGKDVTTPPETEVVLI KNVCLPPEMEVALTEDQVPALKTEA PLAKDGVTLANNVTPAKDVPPLSET EATPVPIKDMEIAQTKGISEDHLE SLQDVGQSAAPTFFMISPETVTGTGK KCSLPAEEDSVLEKLGGERKPCNSQP SELSSETSGIARPEEGRPVVSGTGND ITTPPNKELPPSPEKTKPLATTQPA KTSTSKAKTQPTSLPKQPAPTITIGGL NKKPMSLASGLVPAAPPKRPVAVASA RPSILPSKDVKPKPIADAKAPEKRASP SKPASAPASRSGSKSTQTVAKTTTAA AVASTGPSSRSPSTLLPKKPTAIKTEG KPAEVKMTAKSVPADLSRPKSTST SSMKKTTTTLSGTAPAAGVVPSRVKAT PMPSRPSTTPFIDKKPTSAPKSSTTP RLSRLATNTSAPDLKNVRSKVGSTE NIKHQPGGGRKVEKTEAAATTRK PESNAVTKTAGPIASAQKQPAGKVQI VSKKVSYSHIQSKCGSKDNIKHVPG GGNVQIQNKKVDISKVSSKCGSKAN IKHKPGGGDVKIESQKLNFKKAQA KVGSLDNVGHLPAGGAVKTEGGGSE APLCPGPPAGEEPAISEAAPEAGAPT SASGLNGHPTLSGGGDQREAOQLDS QIQETSI