

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q09666	AHNK_HUMAN	Homo sapiens	Neuroblast differentiation-associated protein AHNAK	32.823372	T146;S148;S230;S232;T247;S283;T308;T318;S332;S337;T348;S442;S501;S521;S572;S613;T638;S698;T788;S819;S947;S983;S1007;S1023;S1068;S1135;S1158;S1263;S1391;S1519;S1580;S1674;S1782;T1885;S1888;T1952;S1955;S2150;S2185;S2313;S2387;S2397;S2454;S2580;S2647;S2775;T2845;S3220;S3240;S3294;S3337;S3360;S3412;S3416;S3483;T3625;S3672;S3692;S4325;S4360;S4386;S4516;S4520;S4522;S4803;S4850;T4917;S5195;S5298;S5310;S5318;S5377;S5386;S5397;S5448;S5552;S5555;S5573;S5593;S5603;S5604;T5610;S5611;S5641;S5643;S5851	S41;S93;T101;S115;S135;T158;S177;S210;S212;S216;T218;S220;S256;S332;S337;S379;S470;T490;S511;T551;T553;S559;S570;S572;S658;S793;S819;S1010;S1042;S1068;S1170;T1192;S1196;S1286;S1298;S1580;S1654;S1856;S1923;T1986;S1990;S2092;S2118;S2138;T2181;S2287;T2309;S2397;S2580;S2600;S2670;S2708;S2728;S2798;T2832;T2845;S3054;S3092;S3182;S3220;S3362;S3409;S3412;S3426;S3544;T3716;S3746;S3766;S3836;S3874;S3964;S4002;S4022;S4092;T4100;S4150;S4220;S4258;S4278;S4360;S4406;S4425;T4430;S4460;S4480;S4486;S4516;S4520;T4564;S4684;S4722;T4766;S4812;S4900;S4903;S4908;S4953;S4960;S4986;S4993;T5009;S5077;S5099;S5110;S5125;S5261;S5318;S5332;S5369;S5386;S5393;S5400;T5415;S5448;S5519;S5530;S5552;S5577;S5620;S5641;S5731;S5739;S5749;S5752;S5762;S5763;S5780;S5782;S5790;S5793;T5794;T5824;S5830;S5841;T5845;S5851;S5857;S5863	22661428;31373491;37340703;38253038;33465208;30444036;30059200;35289036;33214551;28657654;32574038;29351928;20068230;37217939;21158410;34019948;34846842;35254053;30379171;29237092;34725712;35083852;35132862;30620550;31492838;32119511;27655845;23301498	MEKEETTRELLLPNWQSGSHGLTI AQRDDGVFVQEVTONSPAARTGVVK EGDQIVGATITYFDNLQSGEVTQLLNT MGHHTVGLKLRHKGDRSPEPGQTW TREVFSSCSSEVLSGDDEEYQRIYT TIKVPRLKSSEVGLDGETQSRITT VTRRVTYATVDVGTREGAKDIDISSP EFKIKIPRHELTEISNVVDVETQSGKT VIRLPESGGAASPTQSAVDIRAGAISA SGPQLQGGAGHSKLVQVTMPGKIVGGS GVNVAKGLDLGGRGVQVPAVDIS SSLGGRAVEVQGPSLES GDH GKIKF PTMKVPKFGVSTGREGQTPKAGLRV SAPEVSVGHKGGKPLGTIQAPQLEVS VPSANIEGLEGLKKGPOITGPSLEGD LGLKGAKPQGHIGVDASAPQIGCSIT GPSVEVQAPDIDVQGPSKLNVPKM KVPKFSVSGAKGEETGIDVTLPTGEV TVPGVSGDVSLPEIATGGLEGKMMKG TKVKTPEMIIQPKISMQDVDLSLGS PKLKGDIKVSAPGVQGDVKGQVAL KGSRVDIETPNLEGLTGPRLGSPSG KTGTCRISMSEVDLNAAPKVKGGV DVTLPRVEGKVKVPEVDVVRGPKVD SAPDVEAHGPEWNLKMPKMKMPTF STPGAKGEGPDVHMTLPKGDISISGP KVNVEAPDVNLEGLGGKLGKPDVVKL PDMSVKTPKISMVDVLDLHVKGTKVK GEYDVTVPKLEGLKGPVKDIDAPDV DVHGPDWHLKMPKMKMPKFSVPG FKAEGPEVDVNLPKADVDISGPKIDV TAPDVSIEEPEGKLGPKFKMPPEMN IKVPKISMVDVLDLHKGPNVKG EYD VTPKVESEIKVPDVELKSAKMDID VPDVEVQGPDWHLKMPKMKMPKF SMPGFKAEGPEVDVNLPKADVDISG PKVGVEVDVNIIEGPEGKLGPKFK MPPEMNIKAPKISMVDVLDLHMKGPK VKGEYDMTVPKLEGLKGPVKDVSA PDVEMQGPDWNLKMPKIKMPKFS MPKSLKGGEGPEFDVNLKANVDISAP KVDTNAPDLSLEGPEGKLGPKFKM PEMHFRAPKMSLPDVLDLKGPKM KGNVDISAPKIEGEMQVPDVIDIRGPK VDIKAPDVEGQGLDWSLKIPKMKMP KFSMPKLGEGPEVDVNLKADVVV SGPKVDIEAPDVSLEGPEGKLGPKF KMPPEMFKTPKISMVDVLDLHKGPKV KVGDDVSVPKVEGEMKVPDVEIK GPKMDIDAPDVEVQGPDWHLKMPK MKMPKFSMPGFKGEGREVDVNLK ADIDVSGPKVDVEVDVSLGPEGKL KGPKFKMPPEMFKAPKISMVDVLD NLKGPKLKGDVDVSLPEVEGEMKVP DVDIKGPKVDISAPDVDVHGPDWHL KMPKVKMPKFSMPGFKGEGPEVDV KLPKADVDVSGPKMDAEPDVIIEG PDAKLGPKFKMPPEMSIKPQKISIPD VGLHLKGPKMKGDYDVTVPKVEGEI KAPDVIDKGPVDINAPDVEVHGP DWHLKMPKVKMPKFSMPGFKGEGPE VDMNLPKADLVSGPKVDIDVDPVN LEAPEGLKLGPKFKMPSMNIQTHKI SMPDVG LNLKAPKLTDDVDSL PKV

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