

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q6PAL7	AHDC1_MOUSE	Mus musculus	Transcription factor Gibbin	28.070315	S800;S892;T923;S932;T1070;S1073;S1593	S267;S593;S825;S842;S892;S1060;S1180;S1315;S1317;S1392;T1394;S1396;S1498;S1540	22645316;34678516;35822049;36852467	MRVKPQGLVVTSSAVCSSPDYLR KYYPGGPPTPRLLPTRPPASPPDKA FSTHTFSENPRPPRRDPSSRRPPVL AKGDDLLPPRAARPVSAHCPSPAP DNSSLRHWDNGRVNLRPVVQLIDI MKDLTRLSQLQHSGVHLDCCGGLR LSRPPAPPPGDLQYSFFSSPSLANSI RSPEERANPHTKSERPSHPLYEPEP EPRDSPQPGQGHGPGAAATATGLPP EPEPDGPDYSELADADILSELASLTC PEAQLLEAQALEPPSPQPEPQLLDPO PRFLDPQALEPLGEGLELPLQPLAD PLGLPSLTQALDTLPDSLESQLLDP QALDPLPKLLDVPGRRLPEQQSLGH CQLAEPLRLDLCSPHGPPGPEGHPK YALRRTRDRPKLCRRRKAGRGRKAD SGPEGRLPLPMPTGLAAALAEPPPL PPPPPTLSGPGVPPELEPESSTQTPM VPTRKKGCRGVRRMVVKMAKIPVSL GRRNKTTYKVSSLSLSSVEGKELGL RVSSEPTLLKMKNNGRNVVVFPP GEMPIILKRKRGRPPKNLLGPGKPK EPTVVAEEAATVTAATMAMPEVKKR RRRKQKLASFQPSYAADANDSKAEY SDVLAKLAFLNRSQCAGRCSPPRC WTPSEPEVHQAPDTQSSISQFLHRV QGFRRRGKTKGGFGGRGGGHAACA ARCSFSDFFEGIGKKKKVVAVAAPGL VGPGLTELGHPKRKRGRGEVDAVTGK PKRKRRSRKNGLTFPEQVPSGPGFG EAGAEWVGDKGGGWAPHHGHPGG QAGRNCGFQGTARAFASTGLESQA SGRGSYYAGAPSGQTELSQERQNLF TGYFRLLDSDSSDLDLDFALSASRP ESRKASGTYAGPPSSALPAQRGLATF PSRGAKASPVAVGSSGAGADPSFQP VLPSRQTFPPGRATSYGITPATSDCR AAETFPKLAPPPSAVARSPTHPPAN TYPYQGGYGAGQSVFASAKPFSGQ DCANSKDCSFAYGSGNSLPASPSSA HSAGYAPPPTGGPCLPPSKASFFNSS EGGPFSGSAPTPLRCDSTRASTVSPG GYMVPKGTASAASVASSSSSSSFQPS PENCRQFVGASQWPFQGGGLDW ASEAFSQLYNPNFDCHGSEPNVILDI SNYTPQVKQQTAVSETFSESSDST QFSQPVGGGFRANSEASSEGGQS SLSSLEKLMMDWNEASSAPGYNWN QSVLFQSSSKPGRGRRKKVDLFEAS HLGFSTSTSATASGYPSKRSTGPRQP RGGRGSGACSAKKERGGTAAKAKFI PKQPVNPLFQDSPDLGLDYSGDS SMSPLPSQSRAFGVGERDPCDFMG PYSMNPSTPSDGTGQGFHCDSPSL GAAELDGKHFPLAHPPTVFDAGLQ KAYSPTCSPTLGFKEELRPPPSKLT CEPLKHGLQGASLSHAAQHLSCRD LPLGQPHYDSPCKGTAYWYPPGSA ARSPPYEGKVGSGLLADFLGRTEAV CLSAPHLASPPATPKADKEPLEMAR PPGPPRGPAATAGYGCPLLSDLTSL PVPRDSSLPLQDTAYRYPGFMPQAH PGLGGPKSGFLGPMAEPHPEDTFT VTSL

