

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q14686	NCOA6_HUMAN	Homo sapiens	Nuclear receptor coactivator 6	21.209411	S967;S1216;T1224;S1231;T1232;S1235;S1641;S1645;S1649;T1685;S1916;T1920;T1932;T1933	S884;S2018	34019948;37217939;35132862;35254053;35138101;26960196;29237092;33214551;30059200;30620550;29351928;28657654;37340703;28604694;23301498;31492838;32574038;30379171	MVLDDLPLNLEDIYTS LCSSTMEDSE MDFDSGLEDDDDTKSDSILEDSTIFVA FKGNIDDKDFKWKLDAILKNVFNLL HMESSKLKVKVEPWNSVRVTFNIP REAAERLRILAQSNQQLRDLGLISV QIEGEGAINLALAQNRSQDVRMNGP MGAGNSVRMEAGFP MASGPGIIRM NNPATVMIPPGGNVSSSMMAPGPN PELQPRTPRPASQSDAMDPLLSGLH IQQQSHPSGSLAPPHHPMQPVSVNR QMN PANFPQLQQQQQQQQQQQQQ QQQQQQQQQQQLQARPPQHQHQ QQPQGIRPQFTAPTQVPVPPGWNQL PSGALQPPPAQGS LGTMTANQGWK KAPLP GPMQQQLQARPSLATVQTPS HPPPPYPFGSQASQAHTNFPQMS NPGQFTAPQMKSLQGGPSRVPTPLQ QPHLTNKSPASSPSSFQGGSPASSPT VNQTQQQMGRPPQNNPLPQGFQQ PVSSPGRNPMVQQGNVPPNFMVM QQQPPNQG PQSLHPGLGGMPKRLP PGFSAGQANPNFMQGV PSTTATTP GNSGAPQLQANQNVQHAGGQGAGP PQNQM QVSHGPPNMMQPSLMGIH GNMNNQAGTSGVPQVNL SNMQG QPQQGPPS QLMGMHQI VPSQGM VQQQGT LNPQNPMILSRAQLMPQG QMMVNPPSQNLG P SPQRMT PPKQ MLSQQG PQMMAPHNQMMGPQGO VLLQQNPMIEQIMTNQM QGNKQQF NTQNQSNVMGPAQIMRGPTPNMQ GNMVQFTGQMSGQMLPQQGVPNN SPSQVMGIQQVLRPPGSPHMAQ QHGD PATTANNDVLSQMMPDVSI QQTNMVPPHVQAMQNSASGNHF SGHGMSFNAPFSGAPNGNQMSCG QNPGFV NKDVTLTSPLLVNL LQSDI SAGHFGVNNKQNTNANKPKKKKP PRKKKNSQDLNTPDTRPAGLEAD QPPLPGEQGINLDNSGPKLPEFSNR PPGYPSQPVEQRPLQQMPPQLMQH VAPPPQPQQQPQLPQQQPPPP SQPQSQQQQQQQQMMMLMMQ QDPKSVRLPVSQNVHPPRGPLN PDS QRMPMQQSGSVPVMVSLQGPASVP PSPDKQRMMPVNTPLGNSNRKMV YQESPQNSSSPLAEMASLPEASGS EAPSVPGGPNMPSHVVL PQNQLM MTGPKPGSPLSATQGATPQQPPVN SLPSSHGHHPNVAAPTQTSRPKTP NRASPRPYYPQTPNRRPPSTEPSEIS LSPERLNASIALGFPQINIP LPPRPN LNRGFDQQGLNPTTLKAIGQAPS NL TMNPSNFATPQTHKLD SVVNSGK QNSGATKRASPSNSRRSSPGSSRK TTPSPGRQNSKAPKLT LASQTNAALL QNVELPRNVLSPTPLANPPVPGSF

PNNSSLNPQNSTVSVAAVGGVVED
NKESLNVPQSDSDCONSQSRKEQVN
IELKAVPAQEVKMOVVPEDQSKKDGQ
PSDPNKLPSVEENKNLVSPAMREAP
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EVSSNVAPSIPPVMSRPVSSSSISTPL
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PCTSSPVVPSHPPVQQVKELNPDEA
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LISAVPTTKSNHGGIASESLAGGLVE
EKVGSHPPELLPSIAPSQNLVSKETST
TALQASVARPELEVNAIVSGQSSEP
KEIVEKSKIPGRRNSRTEEPTVASES
VENGHRKRSSRPASASSSTKDITS
AVQSKRRKSK