

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondr
Q9Y6Q9	NCOA3_HUMAN	Homo sapiens	Nuclear receptor coactivator 3	11.645293	S90;S652;T654;S844;S847;S857;S860;S863;T910;T912;T914;S917;S925	S214;S551;S569;S601;S694;S728;S857;S860;S867;S1033;S1330	29351928;38253038;39531497;30379171;23301498;38665916;34019948;31492838;30620550;35132862;39302247;35254053	MSG LGENLDPLASDSRKRKLPDTP GQGLTCSGKEKRRREQSKYIEELAEI ISANLSIDIDFNFKPKCAILKETVR QIRQIKEQGKTISNDDVQKADVSS GQGVIDKDSLGPLLLQALDGFLLV NRDGNIVFVSENVYQYQYKQEDLV NTSVYNILHEEDRKDFLKNLPKSTV NGVSWTNETQRQKSHTFNCRMLM KTPHDILEDINASPERRORYETMQC FALSQPRAMMEEGEDLQSCMICVA RRITTGERTFSPNPESFITRHLDSGK VVNIDTNSLRSSMRPGFEDIIRRCIQ RFFSLNDGQSWQKRHYQEAYLNG HAETPVYRFLADGTIVTAQTKSKLF RNPVTNDRHGFVSTHFLQREQNGY RPNPNPVGGQIRPPMAGCNSSVGG MSMSPNQGLQMPSSRAYGLADPST TGQMSGARYGGSSNIASLTPGPGMQ SPSSYQNNNYGLNMSSPPHGGSPGL APNQQNMISPRNRGSPKIASHQFS PVAGVHSPMASSGNTGNHFSFSSSL SALQAISEGVGTSLSTLSSPGPKLD NSPNMNTQPSKVSNOQSKSPLGFY CDQNPVSSMCQSNRDLHSDKES KESSVEGAENQRGPLESKGHKLLQ LLTCSDDRGHSSLTNSPLDSSCKE SSVSVTSPSGVSSSTSGGVSSSTNM HGSLLEKHRILHKLQNGNSPAEV AKITAEATGKDTSSITCGDGNVVKQ EQLSPKKKENNALLRYLLDRDDPSD ALSKELOPQVEGVDNKMSQCTSSSTI PSSSQEKDPKIKTETSEEGSGDLNLD DAILGDLTSSDFYNNSSISNGSHLGT KQOVFOGTNSLGLKSSQSVQIRPP YNRAVSLDSPVSVGSSPPVKNISAFP MLPKQPMGGNPRMMDSQENYGS SMGGPNRNVTVTQTPSSGDWGLPN SKAGRMEPMNSNSMGRPGDYNT SLPRPALGGSIPTPLRSNSIPGARPV LQQQQMLQMRPGEIPMGMGANP YGQAAASNQLGSWPDGMLSMEQVS HGTQNRPLLRNSLDDLVPSPNLEG QSDERALLDQLHTLNTDATGLEEI DRALGIPELVNQQALEPKQDAFQG QEAAVMMDQKAGLYGQTYPAQGGP MQGGFHLQGGQSPFSNMMNQMN QQGNFPLQGMHPRANIMRPTNT KQLRMQLQRLQGGQFLNQRQAL ELKMENTAGGAAMRPMMPQV SSQQGFLNAQMVARSRELLSHHF RQORVAMMMQQQQQQQQQQQQ QQQQQQQQQQQQQQQTQAFSPPP NVTASPSMDGLLAGPTMPQAPPQ FPYQPNYGMQQPDPAFGRVSSPPN AMMSSRMGPSQNPMMQHPQAASI YQSSEMKGWPSGNLARNSSFSQQQ FAHQGNPAVYSMVHMNGSSGHMG QMNMPMPMSGMPMPGDQKYC	True	True	4.619	5.0	2.074