	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
0	Q96JM2	ZN462_HUMAN	Homo sapiens	Zinc finger protein 462	12.327822	S268;S292;T1525;S1784	S350;S354;S688;S1090;S1166;S2172;S2177	30379171;35254053;22826440	MEVLQCDGCDFRAPSYEDLKAHIQD VHTAFLQPTDVAEDNVNELRCGSVN ASNQTEVEFSSIKDEFAIAEDLSGQN ATSLGTGGYYGHSPGYYGQHIAANP KPTNKFFQCKFCVRYFRSKNLLIEH TRKVHGAQAEGSSSGPPVPGSLNYN IMMHEGFGKVFSCQFCTYKSPRRAR IIKHQKMYHKNNLKETTAPPPAPAP MPDPVVPPVSLQDPCKELPAEVVER SILESMVKPLTKSRGNFCCEWCSYQ TPRRERWCDHMMKKHRSMVKILSS LRQQQEGTNLPDVPNKSAPSPTSNS TYLTMNAASREIPNTTVSNFRGSMG NSIMRPNSSASKFSPMSYPQMKPKS PHNSGLVNLTERSRYGMTDMTNSS ADLETNSMLNDSSSDEELNEIDSEN GLSAMDHQTSGLSAEQLMGSDGNK LLETKGIPFRRFMNRFQCPFCPFLT MHRRSISRHIENIHLSGKTAVYKCD ECPFTCKSSLKLGAHKQCHTGTTSD WDAVNSQSESISSSLNEGVVSYESS SINGRKSGVMLDPLQQQQPPQPPPP PPPPPPSQPQPLQQPQPPQLQPPHQ VPPQPQTQPPPTQQPQPPTQAAPLH PYKCTMCNYSTTTLKGLRVHQQHK HSFCDNLPKFEGQPSSLPLENETDS HPSSSNTVKKSQTSILGLSSKNNFVA KASRKLANDFPLDLSPVKKRTRIDEI ASNLQSKINQTKQQEDAVINVEDDE EEEEDNEVEIEVELDREEEPTEPIIEV PTSFSAQQIWVRDTSEPQKEPNFRN ITHDYNATNGAEIELTLSEDEEDYYG SSTNLKDHQVSNTALLNTQTPIYGT EHNSENTDFGDSGRLYYCKHCDFN NKSARSVSTHYQRMHPYIKFSFRYIL DPNDHSAVYRCLECYIDYTNFEDLQ QHYGEHHPEAMNVLNFDHSDLIYR CRFCSYTSPNVRSLMPHYQRMHPT VKINNAMIFSSYVVEQQEGLNTESQ TLREILNSAPKNMATSTPVARGGGL PATFNKNTPKTFTPECENQKDPLVN TVVVYDCDVCSFASPNMHSVLVHYQ KKHPEEKASYFRIQKTMRMVSVDRG SALSQLSFEVGAPMSPKMSNMGSP PPPQPPPPDLSTELYYCKHCSYSNRS VVGVLVHYQKRHPEIKVTAKYIRQAP PTAAMMRGVEGPQGSPRPPAPIQQL NRSSSERDGPPVENEMFFCQHCDY GNRTVKGVLIHYQKKHRDFKANADV IRQHTATIRSLCDRNQKKPASCVLVS PSNLERDKTKLRALKCRQCSYTSPYF YALRKHIKKDHPALKATVTSIMRWA FLDGLIEAGYHCEWCIYSHTEPNGL LLHYQRRHPEHYVDYTYMATKLWA GPDPSPPSLTMPAEAKTYRCRDCVF EAVSIWDITNHYQAFHPWAMNGDE

									SVLLDIIKEKDAVEKPILSSEELAGPV NCENSIPTPFPEQEAECPEDARLSPE KSLQLASANPAISSTPYQCTVCQSEY NNLHGLLTHYGKKHPGMKVKAADF AQDIDINPGAVYKCRHCPYINTRIHG VLTHYQKRHPSIKVTAEDFVHDVEQ SADISQNDVEETSRIFKQGYGAYRCK LCPYTHGTLEKLKIHYEKYHNQPEF DVFSQSPPKLPVPLEPEMTTEVSPSQ VSITEEEVGEEPVSTSHFSTSHLVSH TVFRCQLCKYFCSTRKGIARHYRIKH NNVRAQPEGKNNLFKCALCAYTNPI RKGLAAHYQKRHDIDAYYTHCLAAS RTISDKPNKVIIPSPPKDDSPQLSEEL RRAVEKKKCSLCSFQSFSKKGIVSHY MKRHPGVFPKKQHASKLGGYFTAVY ADEHEKPTLMEEEERGNFEKAEVE GEAQEIEWLPFRCIKCFKLSFSTAEL LCMHYTDHHSRDLKRDFIILGNGPR LQNSTYQCKHCDSKLQSTAELTSHL NIHNEEFQKRAKRQERRKQLLSKQK YADGAFADFKQERPFGHLEEVPKIK ERKVVGYKCKFCVEVHPTLRAICNH LRKHVQYGNVPAVSAAVKGLRSHER SHLALAMFTREDKYSCQYCSFVSAF RHNLDRHMQTHHGHHKPFRCKLC SFKSSYNSRLKTHILKAHAGEHAYK CSWCSFSTMTISQLKEHSLKVHGKA LTLPRPRIVSLLSSHSHHSSQKATPA EEVEDSNDSSYSEPPDVQQQLNHYQ SAALARNNSRVSPVPLSGAAAGTEQ KTEAVLHCEFCEFSSGYIQSIRRHYR DKHGGKKLFKCKDCSFYTGFKSAFT MHVEAGHSAVPEEGPKDLRCPLCLY HTKYKRNMIDHIVLHREERVVPIEV CRSKLSKYLQGVVFRCDKCTFTCSS DESLQQHIEKHNELKPYKCQLCYYE TKHTEELDSHLRDEHKVSRNFELVG RVNLDQLEQMKEKMESSSSDDEDK EEEMNSKAEDRELMRFSDHGAALN TEKRFPCEFCGRAFSQGSEWERHVL RHGMALNDTKQVSREEIHPKEIME NSVKMPSIEEKEDDEAIGIDFSLKNE TVAICVVTADKSLLENAEAKKE