

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q3UHF7	ZEP2_MOUSE	Mus musculus	Transcription factor HIVEP2	47.70794	S1271	S811;S942;S947;S1040;S1431;S1435;S2102;S2281;S2285;S2413;S2415	19458039;22645316;22517741;34678516;35822049	MDTGDTALGQKATSRSGETDSVSGR WRQEQSAVLKMFSSQEGPRPQI DPEQIGNAASQALFGSGKLASPGEG LHQVTEKQYPPHRSPYPCQHSLSF PQHSLSQGMTHSHKPHQSLEGPWW LFPGLPSVASEDLFPPFMHGHSGG YPRKKISNLNPAYSQYSQKSIEQAED AHKKEHKPKKPGKYICPYCSRACAK PSVLKKHRSHTGERPYPCPCGFSF KTKSNLYKHRKSHAHAIKAGLVPFT ESSVSKLDLEAGFIDVEAEIHSDGEQ STDTDEESSLFAEASDKVSPGPPVPL DIASRGGYHGSLEESLGGPMKVPIII PKSGIPLASEGSQYLSSEMLPNPSLN AKADDSTVKQKLALRLSEKKGODS EPLNLLSPHSGSTDSGYFSRSESA EQQISPPNTNAKSYEEIIFGKYRCLSP RNTLSVTPTGQERTAMGRRGIMEPL PHLNTLRLEVKMFEDPISQLNPSKGE MDPGQINMLKTTKFNSECROPQAIP SSVRNEGKPYPGNFLGNSPMLLEAP VDSSPLIRSNMPTSSATNLSVPPSL RGSHSFDERMTGSDDVFPYPTVGIP PQRMLRRQAAPFELPSVQEGHMESE HPARVSKGLASPSLKEKLLPGDRP GYDYDVCRRPKYKWESETLQKQSYL GSFKQGGYFMDPSVPVQGVPTMF GTTCENRKRREKSVGDEEDVPMIC GGMGNAPVGMMSSEYDPLQDGG RSGFAMTAHESLAHGHSDRLDPA PQLPSRSPSLGSEDLPLAADPKMT DLGKKPPGNVISVIQHTNSLRPNSF ERSESTEMVACPODKTPSPAETCDS EVLEAPVSPEWAPPDGGESGSKPT PSQQVPHSYHAQPRLVQHQNIQVP EIRVTEEPDKPEKEEAPTKEPEKPV EEFQWPQRSETLSQLPAEKLPPKKK RLRLADLEHSSGESSFESTGTGLSRS PSQESNLSSHSSFSMSFDREETVKL TAPPKQDESGKHSEFLTVPAGSYSL VPGHHHQEMRRCSEQMPCPHPT EVPEIRSKSFDYGNLSHAPVAGTSPS TLPSPRERKKCFLVRQASFGSPEIA QGEAGVDPVSKQEHMEHLHAGLRA AWSSVLPPLPGDDPGKQVGTCCPLS SGPPLHLTQQQIMHMDSQESLRNP LIQPTSVMYTKSKHLEQPHLFPHQDA VFFSQNALFQFQYPTVCMVHLPA QQPPWWQTHFPHFAPHQNSYSK PPFQADLHSSYLEHVAEHTGKSA DYPHAKEQTYPCYSGTSLHSKNLP LKFPSPDGSKSTETPTEQLLREDFAS ENAGPLQSLPGTVVVRIOITHVPSYG SVMYTSISQILQNSPAIVICKVDEN MTORTLVTNAAMQIGLNLIAQVLGQ HTGLEKYPLWKVPQTLPLGLESIPL CLPSTSDNAASLGGSKRMLSPASSL ELFMETKQKRVKEEKMYGQIVEEL SAVELTNSDIKKGLSRPQKPLVRQ GCASEPKDGCFSRSSFSSLSLSPSS QDHPSASGPPPNREILPGSRAPRR KFSGPSESRESSELDMDDETSSDMS MSPQSSALPTGGGQEEEGKARKLP VSMLVHMASGPGGNVANSTLLFTD VADFQILQFPPLRRTTTVSWCFLNY TKPSFVQQATFKSSVYASWCISSCNP NPSGLNTKTTLALLRSKQITAEIYTL AAMHRPGAGKLTSSVVKQFAQMK PDAPFLFGNKLERKLAGNVLKERGK GEIHGDKDLGSKQTEPIRIKIFEGGY KSNEDYVYVGRGRGKYICEECGIR CKKPSMLKKHIRTHTDVRPYVCKLC NFAFKTKGNLTKHMKSAHMKKCL ELGVSMTSVDDTETEEAENMEELH KTSEKHSMSGISTDHOFSDAEESDG EDGDDNDDDDDDDDDDQDGLTP KTRSRSTSPQPPRFSSLPVNVGAVA HGVPDSSSLGHSSLSLYLVTLPISQVT QLMTPSDSCDDTQMTEYQRLFQSK STDSEPKDRLDIPSSMDEEAMLS	None	None	None	None	None	None	

EPSSSPRDFSPSSYRSPGYDSSPCR  
DNSPKRYLIPKGDLSRRHLSRRDL  
SPMRHLSRKEAALRRMSQGDAS  
PRRHLSRRPLSPGKDITARRDLSR  
RERRYMTTIRAPSPRRALYPNPLSM  
GQYLQTEPVLGPPNLRGIPQVPYF  
SLYGDQEGAYEHGSSLFPEGPTDY  
VFSHLPLHSQQVRAPIPMVPVGGI  
QMVHSLPPALSGLHPPPTLPLPTEG  
SEEKKGAPGEAFKDPYILSRRHEK  
QAPQVLQSSGLPSSPSPRLLMKQS  
TSEDSLNSTEREQENIQTCTKAIAS  
LRIATEEAALLGADPPTWVQESPOKP  
LESAHVSIRHFGGPEPGQCTSAAH  
PDLHDGEKDTFGTSQTAVAHPTFYS  
KSSVDEKRVDQSSKELSLSTEEGN  
EPSPEKNQLH