

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q3UHF7	ZEP2_MOUSE	Mus musculus	Transcription factor HIVEP2	35.437889	S1271	S811;S942;S947;S1040;S1431;S1435;S2102;S2281;S2285;S2413;S2415	19458039;22517741;35822049;22645316;34678516	<p> MDTGDTALGQKATSRSGETDSVSGR WRQEQSAVLKMSTFSSQEGPRQPQI DPEQIGNAASAQLFGSGKLASPGEG LHQVTEKQYPPHRPSPYPCQHLSLF PQHLSLQGMTHSHKPHQSLEGPPW LFPGLPSVASEDLFFPFMHGHS GG YPRKKISNLNPAYSQYSQKSIEQAED AHKKEHKPKKPGKYICPYCSRACAK PSVLKKHRSHTGERPYPCIPCGFSF KTKSNLYKHRKSHAHAIKAGLVPFT ESSVSKLDLEAGFIDVEAEIHSDGEQ STDTDEESSLFAEASDKVSPGPPVPL DIASRGGYHGSLEESLGGPMKVPILII PKSGIPLASEGSQYLSSEMLPNPSLN AKADDSHTVKQKLALRLSEKKGQDS EPSLNLLSPHSGSTDSGYFSRSESA EQQISPPNTNAKSYEEIIFGKYCRLSP RNTLSVTPGTQERTAMGRRGIMEPL PHLNTRLEVKMFEDPISQLNPSKGE MDPGQINMLKTTKFNSECRQPQAIP SSVRNEGKPYPGNFLGSPMLLEAP VDSSPLIRSNMPTSSATNLSVPPSL RGSFSFDERMTGSDDVFYPGTVGIP PQRMLRRQA AFELPSVQEGHMESE HPARVSKGLASPSLKEKLLPGDRP GYDYDVCRKPKKWEDESETLKQSYL GSFKQGG EYFMDPSVPVQGVPTMF GTTCENRKRREKESVGEEDVPMIC GGMGNAPVGMMSSEYDPKLQDGG RSGFAMTAHESLAHGHS DRLD PAR PQLPSRSPSLGSEDLPLAADPKMT DLGKKPPGNVISVIQHTNSLRPN SF ERSESTEMVACPQDKTPSPAETCDS EVLEAPVSP EWAPPGDGGESGSKPT PSQQVPQHSYHAQ PRLV RQHNIQVP EIRVTEEPDKPEKEKEAPTKEPEKPV EEFQWPQRSETLSQLPAEKLPPKKK RLRLADLEHSSGESSFESTGTGLSRS PSQESNL SHSSSFMSFDREETVKL TAPPKQDESGKHSEFLTVPAGSYSLS VPGHHHQEMRRCSSEQMPCPHPT EVPEIRSKSFDYGNLSHAPVAGTSPS T LSPSRERKKCF LVRQASFGSPEIA QGEAGVDPSVKQEHMEHLHAGLRA AWSSVLPPLPGDDPGKQVGTGGLS SGPPLHLTQQQIMHMDSQESLRNP LIQPTSYMTSKHLPEQPHLFPHQDA VPFSPIONALFQFYPTVCMVHLPA QQPPWWQTHFPHPFAPHPQNSYSK PPFQADLHSSYPLEHVAEHTGKKS A DYPHAKEQTYPCYSGTSGLHSKNLP LKFPSDPGSKSTETPTEQLLREDFAS ENAGPLQSLPGTVVPVRIQTHVPSYG SVMYTSISQILGQNSPAIVICKVDEN </p>

MTQRTLVTNAAMQGGIGLNIAQVLGQ
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PRRHLSRRPLSPGKDITARRDLSR
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