

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9H4Z2	ZN335_HUMAN	Homo sapiens	Zinc finger protein 335	10.544825	S997	S976;S992;S1007;S1153	30059200	MEENEVESSSDAAPGPRPEEPSES GLGVGTSEAVSADSSDAAAAPGQAE ADDSGVGQSSDRGSRSQEEVSESS SADPLPNSYLPDSSSVSHGPVAGVT GGPPALVHSSALPDPNMLVSDCTAS SSDLGSAIDKIIESTIGPDLIQNCITVT SAEDGGAETTRYLILQGPDDGAPMT SPMSSSTLAHSLAAIEALADGPTSTS TCLEAQGGPSSPVQLPPASGAEED LQSLEAMMEVVVVQQFKCKMCQYR SSTKATLLRHRERHFRPVAAAAA AGKKGRLRKWSTSTKSQEEEGPEEE DDDDIVDAGAIDDLEEDSDYNPAED EPRGRQLRLQRPTPSTPRRRRPGR PRKLPRLAISDLPDGVEGEPLVSSQS GQSPPEPQDPEAPSSSGPGLVAMG KVSRTPEAGVSQSDAENAAPSCPD EHDTLPRRRGRPSRRFLGKKYRKY YKSPKPLLRPFLCRICGSRFLSHEDL RFHVNSHEAGDPQLFKCLQCSYRSR RWSSLKEHMFNHVSGSKPYKCDECS YTSVYRKDVIRHAAVHSRDRKKRPD PTPKLSSFPCPVCGRVYPMQKRLTQ HMKTHSTEKPHMCDKCGKSFKKRY TFKMHLTHIQAVANRRFKCFCEFCF VCEDKKALLNHQLSHVSDKPFKCSF CPYRTFREDFLLSHVAVKHTGAKPF ACEYCHFSTRHKKNLRLHVRCRHA SSFEEWGRRHPEEPPSRRRPPFFSLQ QIEELKQQHSAAPGPPSSPGPPEIP PEATTFQSSEAPSLLCSDTLGGATIY QQGAEESTAMATQTALDLLLLNSA QRELGGTALQVAVVKSEDVEAGLAS PGGQPSPEGATPQVVTLHVAEPGGG AAAESQLGPPDLQITLAPGPFGGTG YSVITAPPMEEGTSAPGTPYSEEPAG EAAQAVVSDTLKEAGTHYIMATDG TQLHHIELTADGSISFSPDALASGA KWPLLQCGGLPRDGPEPPSPAKTHC VGDSQSSASSPPATSKALGLAVPPSP PSAATAASKKFCKICAEAFPGAEM ESHKRAHAGPGAFKCPDCPFSARQ WPEVRAHMAQHSSLRPHQCSQCSF ASKNKKDLRRHMLTHTKEKPFACH LCGQRFNRNGHLKFHIQRLHSPDG RKSGTPTARAPTQTPTQTIILNSDDE TLATLHTALQSSHGVLGPERLQQAL SQEHIIVAQEQTVTNQEEAAYIQEITT ADGQTVQHLVTSNDQVQYIISQDGV

QHLLPQEYVVVPEGGHHIQVQEGQIT  
HIQYEQGAPFLQESQIQYVPVSPGQQ  
LVTQAQLEAAAHSAVTAVADAAMAQ  
AQGLFGTDETVPHEHIQQLQHOGIEY  
DVITLADD