

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9H4Z2	ZN335_HUMAN	Homo sapiens	Zinc finger protein 335	11.956056	S997	S976;S992;S1007;S1153	30059200	MEENEVESSSDAAPGPRPEEPSES GLGVGTSEAVSADSSDAAAAPGQAE ADDSGVGQSSDRGSRSQEEVSESS SADPLPNSYLPDSSSVSHGPVAGVT GGPPALVHSSALPDPNMLVSDCTAS SSDLGSAIDKIIESTIGPDLIQNCITVT SAEDGGAETTRYLILQGPDDGAPMT SPMSSSTLAHSLAAIEALADGPTSTS TCLEAQGGPSSPVQLPPASGAEED LQSLEAMMEVVVVQQFKCKMCQYR SSTKATLLRHMRERHFRPVAAAAA AGKKGRLRKWSTSTKSQEEEGPEEE DDDDIVDAGAIDDLEEDSDYNPAED EPRGRQLRLQRPTPSTPRRRRPGR PRKLPRLAISDLDPGVEGEPLVSSQS GQSPPEPQDPEAPSSSGPGLVAMG KVSRTPEAGVSQSDAENAAPSCPD EHDTLPRRRGRPSRRFLGKKYRKY YKSPKPLLRPFLCRICGSRFLSHEDL RFHVNSHEAGDPQLFKCLQCSYRSR RWSSLKEHMFNHVGSKPYKCDECS 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

