	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q9H4Z2	ZN335_HUMAN	Homo sapiens	Zinc finger protein 335	25.875327	S997	S976;S992;S1007;S1153	30059200	MEENEVESSSDAAPGPGRPEEPSES GLGVGTSEAVSADSSDAAAAPGQAE ADDSGVGQSSDRGSRSQEEVSESSS SADPLPNSYLPDSSSVSHGPVAGVT GGPPALVHSSALPDPNMLVSDCTAS SSDLGSAIDKIIESTIGPDLIQNCITVT SAEDGGAETTRYLILQGPDDGAPMT SPMSSSTLAHSLAAIEALADGPTSTS TCLEAQGGPSSPVQLPPASGAEEPD LQSLEAMMEVVVVQQFKCKMCQYR SSTKATLLRHMRERHFRPVAAAAAA AGKKGRLRKWSTSTKSQEEEGPEEE DDDDIVDAGAIDDLEEDSDYNPAED EPRGRQLRLQRPTPSTPRPRRRPGR PRKLPRLEISDLPDGVEGEPLVSSQS GQSPPEPQDPEAPSSSGPGHLVAMG KVSRTPVEAGVSQSDAENAAPSCPD EHDTLPRRRGRPSRRFLGKKYRKYY YKSPKPLLRPFLCRICGSRFLSHEDL RFHVNSHEAGDPQLFKCLQCSYRSR RWSSLKEHMFNHVGSKPYKCDECS YTSVYRKDVIRHAAVHSRDRKKRPD PTPKLSSFPCPVCGRVYPMQKRLTQ HMKTHSTEKPHMCDKCGKSFKKRY TFKMHLLTHIQAVANRRFKCEFCEF VCEDKKALLNHQLSHVSDKPFKCSF CPYRTFREDFLLSHVAVKHTGAKPF ACEYCHFSTRHKKNLRLHVRCRHA SSFEEWGRRHPEEPPSRRRPFFSLQ QIEELKQQHSAAPGPPPSSPGPPEIP PEATTFQSSEAPSLLCSDTLGGATIIY QQGAEESTAMATQTALDLLLNMSA QRELGGTALQVAVVKSEDVEAGLAS PGGQPSPEGATPQVVTLHVAEPGGG AAAESQLGPPDLPQITLAPGPFGGTG YSVITAPPMEEGTSAPGTPYSEEPAG EAAQAVVVSDTLKEAGTHYIMATDG TQLHHIELTADGSISFPSPDALASGA KWPLLQCGGLPRDGPEPPSPAKTHC VGDSQSSASSPPATSKALGLAVPPSP PSAATAASKKFSCKICAEAFPGRAEM ESHKRAHAGPGAFKCPDCPFSARQ WPEVRAHMAQHSSLRPHQCSQCSF ASKNKKDLRRHMLTHTKEKPFACH LCGQRFNRNGHLKFHIQRLHSPDG RKSGTPTARAPTQTPTQTIILNSDDE TLATLHTALQSSHGVLGPERLQQAL SQEHIIVAQEQTVTNQEEAAYIQEITT ADGQTVQHLVTSDNQVQYIISQDGV QHLLPQEYVVVPEGHHIQVQEGQIT HIQYEQGAPFLQESQIQYVPVSPGQQ LVTQAQLEAAAHSAVTAVADAAMAQ AQGLFGTDETVPEHIQQLQHQGIEY DVITLADD	None	None	None	None	None	None	None	None	None