	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	Q14999	CUL7_HUMAN	Homo sapiens	Cullin-7	18.696129	NaN	S339	30379171;37217939;32522876	MVGELRYREFRVPLGPGLHAYPDEL IRQRVGHDGHPEYQIRWLILRRGDE GDGGSGQVDCKAEHILLWMSKDEIY ANCHKMLGEDGQVIGPSQESAGEV GALDKSVLEEMETDVKSLIQRALRQ LEECVGTIPPAPLLHTVHVLSAYASIE PLTGVFKDPRVLDLLMHMLSSPDYQ IRWSAGRMIQALSSHDAGTRTQILLS LSQQEAIEKHLDFDSRCALLALFAQA TLSEHPMSFEGIQLPQVPGRVLFSLV KRYLHVTSLLDQLNDSAAEPGAQNT SAPEELSGERGQLELEFSMAMGTLI SELVQAMRWDQASDRPRSSARSPG SIFQPQLADVSPGLPAAQAQPSFRRS RRFRPRSEFASGNTYALYVRDTLQP GMRVRMLDDYEEISAGDEGEFRQS NNGVPPVQVFWESTGRTYWVHWH MLEILGFEEDIEDMVEADEYQGAVA SRVLGRALPAWRWRPMTELYAVPYV LPEDEDTEECEHLTLAEWWELLFFI KKLDGPDHQEVLQILQENLDGEILD DEILAELAVPIELAQDLLLTLPQRLN DSALRDLINCHVYKKYGPEALAGNQ AYPSLLEAQEDVLLLDAQAQAKDSE DAAKVEAKEPPSQSPNTPLQRLVEG YGPAGKILLDLEQALSSEGTQENKVK PLLLQLQRQPQPFLALMQSLDTPET NRTLHLTVLRILKQLVDFPEALLLPW HEAVDACMACLRSPNTDREVLQELI FFLHRLTSVSRDYAVVLNQLGARDAI SKALEKHLGKLELAQELRDMVFKCE KHAHLYRKLITNILGGCIQMVLGQIE DHRRTHQPINIPFFDVFLRYLCQGSS VEVKEDKCWEKVEVSSNPHRASKLT DHNPKTYWESNGSAGSHYITLHMR RGILIRQLTLLVASEDSSYMPARVVV CGGDSTSSLHTELNSVNVMPSASRV ILLENLTRFWPIIQIRIKRCQQGGIDT RIRGLEILGPKPTFWPVFREQLCRHT RLFYMVRAQAWSQDMAEDRRSLLH LSSRLNGALRQEQNFADRFLPDDEA AQALGKTCWEALVSPVVQNITSPDE DGISPLGWLLDQYLECQEAVFNPQS RGPAFFSRVRRLTHLLVHVEPCEAP PPVVATPRPKGRNRSHDWSSLATRG LPSSIMRNLTRCWRAVVEKQVNNFL TSSWRDDDFVPRYCEHFNILQNSSS ELFGPRAAFLLALQNGCAGALLKLP FLKAAHVSEQFARHIDQQIQGSRIGG AQEMERLAQLQQCLQAVLIFSGLEIA TTFEHYYQHYMADRLLGVVSSWLE GAVLEQIGPCFPNRLPQQMLQSLST SKELQRQFHVYQLQQLDQELLKLED TEKKIQVGLGASGKEHKSEKEEEAG AAAVVDVAEGEEEEEENEDLYYEGA MPEVSVLVLSRHSWPVASICHTLNP RTCLPSYLRGTLNRYSNFYNKSQSH PALERGSQRRLQWTWLGWAELQFG NQTLHVSTVQMWLLLYLNDLKAVS VESLLAFSGLSADMLNQAIGPLTSSR GPLDLHEQKDIPGGVLKIRDGSKEPR SRWDIVRLIPPQTYLQAEGEDGQNL EKRRNLLNCLIVRILKAHGDEGLHID QLVCLVLEAWQKGPCPPRGLVSSLG KGSACSSTDVLSCILHLLGKGTLRRH DDRPQVLSYAVPVTVMEPHTESLNP GSSGPNPPLTFHTLQIRSRGVPYASC TATQSFSTFR	None	None	None	None	None	None	None	None	None