

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
076039	CDKL5_HUMAN	Homo sapiens	Cyclin-dependent kinase-like 5	18.379595	S479;S543	S407;S479;S720;S761	30379171	MKIPNIGNVMNKFEILGVVGEAYG VVLKCRHKETH EIVAIKKFKDSEENE EVKETTLLRELKMLRTLKQENIVELKE AFRRRGKLYLVFEYVEKNMLELLEE MPNGVPPEKVKSYIYQLIKAIHWCH KNDIVHRDIKPENLLISHNDVLKLCD FGFARNLSEGNNANYTEYVATRWY RPELLLGAPYGKSVDMWMSGCILG ELSDGQPLFPGESEIDQLFTIQKVLG PLPSEQMCLFYSNPRFHGLRFPVAVN HPQSLERRYLGILNSVLLDLMKNLL KLDPADRYLTEQCLNHPTFQTQRLL DRSPRS AKRKP YHVESSTLSNRNQ AGKSTALQSHHRSNSKDIQNLSVGL PRADEGLPANESFLNGNLGASLSP LHTKTYQASSQPGSTSKDLTNNNIP HLLSPKEAKSKTEFDNFIDPKPSEGP GTKYLKSNRSRQQNRHSFMESSQS KAGTLQPNEKQSRHSYIDTIPQSSRS PSYRTKAKSHGALSDSKSVNLSEA RAQIAEPSTSRYPSSCLDLNSPTSP TPTRHSDTRTLLSPSGRNNRNEGTL DSRRTTTRHSKTMEELKLP EHMDS HSHLSAPHEFSYGLGYTSPFSSQ QRPHRHSMYVTRDKVRAKGLDGSL SIGQGM AARANS LQLLSPQPGQLP PEMTVARSSVKETSREGTSSFHTRQ KSEGGVYHDPHSDDG TAPKENRHL YNDPVPRRVGSFYRVSPRPDNSFH ENNVSTRVSSLPSESSSGTNHNSKRO PAFDPWKSPENISHSEQLKEKEKQG FFRSMKKKKKKSQTVPNSDSPDLLT LQKSIHSASTPSSRPKEWRPEKISDL QTQSQPLKSLRLLHLSSASNHPAS SDPRFQPLTAQQTKNFS EIRIHP QASGGSSNIRQEPAPKGRPALQLPG QMDPGWHVSSVTRSA TEGPSYSEQ LGAKSGPNGHPYNRTRSRMPNLN DLKETAL	True	False	3.447	4.779	1.828	1.013	0.676	1.691	1.566