

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q96BD5	PF21A_HUMAN	Homo sapiens	PHD finger protein 21A	21.838376	S109;T117;T122;T124;T125;S127;T142;T155;T270;T272;T273;T276;S277;S280	T350;T444;S447	28657654;35132862;35254053;34846842;23301498;34019948;29237092;30059200;32574038;30379171;37340703;22661428;38665916;36240223;39534244;35289036;39302247;29351928;39531497;31492838;33214551	MELQTLQEALKVEIQVHQKLV AQM KQDPQNA DLKQLHELOAKITALSE KQKR VVEQLRKNLIVKQEQPKFQI QPLPQSENKLQTAQQPLQLQQQ QQYHHHHAQSSAAASP NLTASQKT VTTASMITTKTLPVLVKAATATMPAS VVGQRPTIAMVTAINS QKAVLSTDVQ NTPVNLQTSSKVTGPGAEAVQIVAK NTVTLVQATPPQPIKVPQFIPPPRLTP RPNFLPQVRPKPVAQNNIPIAPAPP MLAAPQLIQRPVMLTKFTPTTLPTSQ NSIHPV RVVNGQTATI AKTFPMAQL TSIVIATPGTRLAGPQTVQLSKPSLEK QTVKSH TETDEKQTESRTITPPAAPK PKREENPQKLA FMVSLGLVTHDHLE EIQSKRQERKRRTTANPVYSGAVFEP ERKKS AVTYL NSTMHPGTRKRGRPP KYNAVLGFGALTPTSPQSSHPDSPE NEKTETTTFFPAPVQPVSLPSTSD GDIHEDFCSVCRKSGQLLMCDTCSR VYHLDCLDPPLKTIPKGMWICPRCQ DQMLKKEEAIPWPGT LAIVHSYIAYK AAKEEEKQKLLKWSSDLKQEREQLE QKVKQLSNSISKMEMKNTILARQK EMHSSLEKVKQLIRLIHGIDLSKPVD SEATVGAISNGPDCTPPANAATSTPA PSPSSQSTANCNQGEETK	True	False	1.851	4.736	1.054	0.578	False