	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	e r
0	P16383	GCFC2_HUMAN	Homo sapiens	Intron Large complex component GCFC2	30.518348	S602	S16;S17;S19;S40;S96;T97;S129;S174;S180;T213;S214;S217	23301498;40307207	MAHRPKRTFRQRAADSSDSDGAEE SPAEPGAPRELPVPGSAEEEPPSGGG RAQVAGLPHRVRGPRGRGRVWASS RRATKAAPRADEGSESRTLDVSTDE EDKIHHSSESKDDQGLSSDSSSSLG EKELSSTVKIPDAAFIQAARRKRELA RAQDDYISLDVQHTSSISGMKRESE DDPESEPDDHEKRIPFTLRPQTLRQ RMAEESISRNEETSEESQEDEKQDT WEQQQMRKAVKIIEERDIDLSCGNG SSKVKKFDTSISFPPVNLEIIKKQLNT RLTLLQETHRSHLREYEKYVQDVKS SKSTIQNLESSSNQALNCKFYKSMKI YVENLIDCLNEKIINIQEIESSMHALL LKQAMTFMKRRQDELKHESTYLQQ LSRKDETSTSGNFSVDEKTQWILEEI ESRRTKRRQARVLSGNCNHQEGTS SDDELPSAEMIDFQKSQGDILQKQK KVFEEVQDDFCNIQNILLKFQQWRE KFPDSYYEAFISLCIPKLLNPLIRVQLI DWNPLKLESTGLKEMPWFKSVEEF MDSSVEDSKKESSSDKKVLSAIINKT IIPRLTDFVEFLWDPLSTSQTTSLITH CRVILEEHSTCENEVSKSRQDLLKSI VSRMKKAVEDDVFIPLYPKSAVENK TSPHSKFQERQFWSGLKLFRNILLW NGLLTDDTLQELGLGKLLNRYLIIAL LNATPGPDVVKKCNQVAACLPEKWF ENSAMRTSIPQLENFIQFLLQSAHKL SRSEFRDEVEEIILILVKIKALNQAES FIGEHHLDHLKSLIKED	True	False	4.037	5.0	False	False	False	0.188	F