	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	Q0VG06	FP100_HUMAN	Homo sapiens	Fanconi anemia core complex-associated protein 100	21.607386	S466	S667	30379171;36240223	MAGAAPRVRYLAGFCCPLGGLAAGK PRVLCHEAEVFLSTGSELVYVYDQE GGLLTAAFRFPDQVWHLELLAPRRL LYALCARRGLYCLSLDHPGRSRSTS QDDRDSEDGDQPSPVIPVDPDACILP DAALCAFTLLDSVLVTLVQGPARWK MQLFEQPCPGEDPRPGGQIGEVELS SYTPPAGVPGKPAAPHFLPVLCSVSP SGSRVPHDLLGGSGGFTLEDALFGL LFGADATLLQSPVVLCGLPDGQLCC VILKALVTSRSAPGDPNALVKILHHL EEPVIFIGALKTEPQAAEAAENFLPD EDVHCDCLVAFGHHGRMLAIKASW DESGKLVPELREYCLPGPVLCAACG GGGRVYHSTPSDLCVVDLSRGSTPL GPEQPEEGPGGLPPMLCPASLNICS VVSLSASPRTHEGGTKLLALSAKGRL MTCSLDLDSEMPGPARMTTESAGQ KIKELLSGIGNISERVSFLKKAVDQR NKALTSLNEAMNVSCALLSSGTGPR PISCTTSTTWSRLQTQDVLMATCVL ENSSSFSLDQGWTLCIQVLTSSCAL DLDSACSAITYTIPVDQLGPGARREV TLPLGPGENGGLDLPVTVSCTLFYSL REVVGGALAPSDSEDPFLDECPSDV LPEQEGVCLPLSRHTVDMLQCLRFP GLAPPHTRAPSPLGPTRDPVATFLET CREPGSQPAGPASLRAEYLPPSVASI KVSAELLRAALKDGHSGVPLCCATL QWLLAENAAVDVVRARALSSIQGVA PDGANVHLIVREVAMTDLCPAGPIQ AVEIQVESSSLADICRAHHAVVGRM QTMVTEQATQGSSAPDLRVQYLRQI HANHETLLREVQTLRDRLCTEDEAS SCATAQRLLQVYRQLRHPSLILL	None	None	None	None	None	None	None	None	None