	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	Q8MYL1	FACD2_DROME	Drosophila melanogaster	Fanconi anemia group D2 protein homolog	28.06426	S489;S490	NaN	40245499	MYKQFKKRSKKPLNTIDENATIKVPR LAETTTNISVESSSGGSEENIPASQE HTQRFLSQHSVILAATLGRTQSSSR NIATLSRQPNNFFELVLVRAGVQLD QGDSLILACDHVPIVSKLREIFTSASS YTDKMETFKTGLNAAMAPGSKLVQ KLLTGCTVDAAGEEQIYQSQNSMFM NFLMIDFMRDACVEVLLNKIEEVAK SDRVIMGKAAIPLPLLPLMLTQLRYL TASHKVEIYSRIEVIFNRATESAKLDII ANAELILDASMHDEFVELLNINYSST EDLFHMTTVQTLGNLSLSDRTQAKL RVRILDFATSGQCSDAILPHLIRLLLN VLKIDTDDSVRDLIGSLRGIFNWRHT NETEKISASEDSKKSQLELFGFLELG LIRSKKFYQACQRSSASVPAAEFTSF DMILLLLIHVNEDNSLYIENILRRRI KLEHITVSILEEIRQHYRHILEQHITT LMNILHDFMREKNRIVSDFAKSSYSI LFKIFNSIQKNILKKLLELTCDKSSP HLTTMALELLRELQRKSAKDVQNCA TLLIPMLDRISDLSLTQTRVAMDLLC HVAFPDPNLSPCLQLQEQVDMVVK KQLINSIDNIKKQGIIGCVQLIDAMAR IANNGVDRDEFIASVENVDSLPDGR GKMAANLIIRTEASIGNSTESLALFF EELATVFNQRNEGTSGCELDNQFIA WACDLVTFRFQASFVTENVPETLKG IKLEYQLNINELDDTDANTESDVLNI GINISKLVLSPKVKACDSIYVLAPLFN YVRVLYKHRHQDSLESINALLGCAIV LPSFFEDDNYVSVFENFEAEQQKDI LSIYFHTVNWMRVSISAFASQRDPPT RRRVLSRLGELIRIEQRMKPLLARAP VDFVAPPYQFLTNVKLSNQNQKRPG PKPAAKLNATLPEPDLTGNQPSIADF TIKVGQCKTVKTKTDFEQMYGPRER YRPMEVEIIMLLVEQKFVLNHQLEE EQMGEFLGLLELRFLLEDVVQKLEA AVLRHHDSYDADSFRPHLAKPEDFI CDLLPCLHEVNNHLITLGEAIDNQL TEVSHVYSNLDLFKDQFCYIKSCFGL CVRLFALYFAWSEWSDKSQEQLLH KSLLKLQPKTQWKRLEKQSVPQLAN LTFQYFLKYEKSVLNLSTAVHLHRLL CNLLKLGSLQSDASQPRFQQAEDLR ILCGTLLRRKWFHYSGTLDKGGQCN IYLDELVKGFLKKSNAKSQTELLTEL VKQCSILNTKDKALTSFPNFKKANF PLLFRGLCEVLIHSLSGQVSVDSRGD KLKLWESAVDLLNGLLSIVQQVEQP RNFGLFLKHSLLFLKLLLQHGMSAL ESIVREDPERLTRFLHELQKVTRFLH QLCCHSKSIKNTAIISYIPSLRETIETL VFRVKALLAANNCHSAFHMGNMIN RDLHGDSIITPRSSFAGEENSDDELP ADDTSVDETVLGDDMGITAVSVSTR PSDGSRRSKSSSRSKCF	True	False	2.395	4.696	1.271	0.872	False	0.974	0.934