

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
A0AVI2	FR1L5_HUMAN	Homo sapiens	Fer-1-like protein 5	19.994096	NaN	NaN	37217939	MLRLVVQSAKIDPPLAPLPRPCMSID FRDIKKRTRVVEGNDPVVWNETLIWH LWNRPLENDSFLQVTLQDMGSQKK ERFIGLATVLLKPLLKQPSEVLFVKD LTLNHSMKPTDCTVTLQVAHMSN QDIEKTGAEDHLGITAREAASQKLM VPGSTAHRALSSKQPHFQVRVKVFE ARQLMGNNIKPVVKVSIAGQQHQTR IKMGNNPFFNEIFFQNFHEVPAKFF DETLIQVVNSSAMRYKAEIGRFQTD IGFIYHSPGHTLLRKLWGLCQPNNP GSGVTGYLKVTIYALGVGDQALIDQK LLYCTDDTDIQIFKSAVVPINMAYLQ LFIYCAEDLHLKHHQSVNPQLEVELI GEKLRTHMQTQTDNPIWNQILTFRI QLPCLSSYKFRVLDCKRKDCPDEIG TASLSLNQISSTGEEIEGVYSGFLPCF GPSFLTLLHGGKKAPFRIQEEGACIPD SVRDGLAYRGRVFLELITQIKSYQDS TIKDLSEVTRIEKHQNRQKYGLCVI FLSCTMMPNFKELIHFEVSIGHYGN KMDLNYKPLVSSTPYSPVIYDGNIIYH YVPWYNTKPVVAVTSNWEDVSFRM NCLNLLHFTRDRLKANLDTLTKSTRN PKDPALLYQWEKLLRELAEDCKRPL PCMTYQPKATSLDRKRWQLRSLLLQ ELAQKAKQAKPKDMVATAEDWLRYR LNTVLPPEPQMGLPDVMIWLVAKEQ RVAYAQVPAHSVLFSPAGALHSGRL CGKIQTLFLOYPEGEGQKDVLPABL RVCMWLGNVTDSDKLQLLRQGDTA VYAEMYENQAKYKQWGOQGLYHC PNFSDVMGNKTLPMDFQPPLGWH WQDSWTVPEQRLLLDIDINKSQVL EEVYENQGRDTRGAWGPAIAPNTDV NGQPMEARENKCPQGWHFKKDW VVELNHAVDSKQWEYGVGIPPSGLP QVWSPVEKTYHSCRERRRWARVRF NHGELSHEQETLSFLQLGLARKEE GWEYDTFGSKFHLNPPQSRFRFR CWRRRLAPNKDKGIAPIFLEGLA MDLKYHAGKEEDSKTWPWGLDRQF RDPQRQDTRPPNLPFIYCTFNKPHY YQLFCYIYQARNLVSNIQILTFQGPFI RVVFLNHSQCTQTLRSSAGPTWAQT LIFQHLLLYENPQDTKESPLVLEL WORDFWGKESLWGRSVWPPMVWL DLQDRILPPMRWHPLVKELGKEE ILASCELILQTEKLGKQLPILSVPWK NGAYTLPKSIQPTIKRMAIEILAWGL RNMKKASSQLLVEFGEESLRTEPI RDFQTNPNFPESESVLVTVMPT EAYALPLVVVVDNWFQQTVTGQ ANIDFLQPYFCDPWAQDYMHPKLP LSEKQHDFLGYLYRKFWFKSSKAE DEYEHEVDWWSKLFWATDEHKS LKYKYDYHTLKVYECELEAVPAFQGL QDFCQTFKLYQEQPKLDSVVGFEK GLFRIYFPFENPEAPKPLQFLVWPE REDFPQPCLVRVYMVRAINLQPDY NGLCDPYVILKLGKTELGNRDMYQP NTLDPIFGMMFELTCNIPLEKDLEIQ LYDFDLFSPDDKIGTTVIDLENRLS GFGAHCGLSKSYCQSGPFRWRDQ PPSYLLERYAKRKGLPPPLFSPEEDA VFYNGKFKLQSFEPKPTPTVHGLGP KKERLALYLLHTQGLVPEHVETRTLY SHSQPGIDQGKVMWVDIFPKKLG PGPQVNIINPRPKRYELRCHWKTAN VDLVDDNLSREKTSDIYIKGWLYGLE KDMQKTDIHYHSLTGEADFNWR FIMDYLAARTCVQSQKDYIWSLDAT SMKFFPARLIQVVDNDFSPDDFLG VLELDLSDMPLPARHAKQCSIRMM	None	None	None	None	None	None	None		

