	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plas men
0	Q76LL6	FHOD3_MOUSE	Mus musculus	FH1/FH2 domain-containing protein 3	30.010553	T307;S345;T351	S345;S376;T413;S921;T933	36288343;30016717;34418053	MATLACRVQFLDDTDPFNSTNFPEP SRPPLFTFREDLALGTQLAGVHRLLR APHKLDDCTLQLSHNGAYLDLEATL AEQRDELEGFQDDTGRGKKNSIILR TQLSVRVHACIEKLYNSSGRDLRRAL FSLKQIFQDDKDLVHEFVIAEGLTCL IKVGAEADQNYQNYILRALGQIMLYV DGMNGVINHSETIQWLYTLVGSKFR LVVKTALKLLLVFVEYSESNAPLLIQ AVSAVDTKRGVKPWSNIMEILEEKD GVDTELLVYAMTLVNKTLAGLPDQD TFYDVVDCLEELGIAAVSQRHLNKK GTDLDLLEQFNIYEVALRHEDGDET AEPPPSGHRDRRRASMCSGGTVGE QQGLDRRRSRRHSIQNIKSPLSAPTS PCSPSVPAFKPSQVRDLCEKDEEEE EEEEQPITEPNSEEEREDDAQCQGK DSKASSASGQSSPGKDAAPESSALH TTSSPTSQGRWLSASTAARSPVLGG TSGPEASRPAARLLPPSPGLATRPST APKVSPTIDKLPYVPHSPFHLFSYDF EDSPLLTKDKGGDSQTENRYSNFSS NSFQSSRPSPGPSGSPSYASSFSSPQ DTRSSPSGLLTSSFRQHQESLAAER ERRRQEREERLQRIEREERNKFNRE YLDKREEQRQARGERYKYLEQLAAE TQEKEPRSQSVSRGRADLSLDLSLP AAPAPPSPSSQSPSADSQEALPVPSS PPTLQCPQVSGKDHEPELEAEAGQG ADEASQDIASAHRGAESQEEPVLEL EPEERASLSEKERQNEEVNERDNCS ASSISSSSSTLEREEKEDKLSEDRAT GLWSTSLQDVGVNGQCGDILTSKRF MLDMLYAHNRKSTEDEEKDDGEPG RSAQEVEAVASLATRISTLQANSQAP EESIKRVDIGCLDNRGSVKAFAEKFN SGEVGRGAISPDVESQDKVPDTPPA QLKTESDYIWDQLMANPRELRIQDM DFTDLGEEDDIDVLDVDLGHREAPG PPPPPPPTFLGLPPPPPPPLLDSVPPP PVPGNLLASPVFNTPQGLGWSQVPR GQPAFTKKKKTIRLFWNEVRPFEWP SKNNRRCREFLWSKLEPIKVDTSRL EHLFESKSKELSVTKKTAADGKRQEI IVLDSKRSNAINIGLTVLPPPRTIKIAI LNFDEYALNKEGIEKILTMIPTEEEK QKIQEAQLANPEVPLGSAEQFLLTLS SISELSARLHLWAFKMDYETTEKEV AEPLLDLKEGIDQLENNKTLGFILST LLAIGNFLNGTNAKAFELSYLEKVPE VKDTVHKQSLLHHVCTMVVENFPD SSDLYSEIGAITRSAKVDFDQLQDNL CQMERRCKASWDHLKAIAKHEMKP VLKQRMSEFLKDCAERIIILKIVHRRI INRFHSFLLFMGHPPYAIREVNINKF CRIISEFALEYRTTRERVLQQKQKRA NHRERNKTRGKMITDSGKFSGSSPA APSQPQGLSYAEDAAEHENMKAVLK TSSPALEDATPVLGVRTRSRASRGST SSWTMGTEESPSVTDDAADEIMDRI VKSATQVPSQRVVPRERKRSRANRK SLRRTLKSGLTPEEARALGLVGTSEL QL	None	None	None	None	None	None	None	None