

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracel region
Q7TMB8	CYFP1_MOUSE	Mus musculus	Cytoplasmic FMR1-interacting protein 1	28.385736	NaN	S583;T1234	29187734;28528544;33300544	MAAQTLEDALSNVDLLEELPLPDQ QPCIEPPSSLLYQPNFNTNFEDRN AFVTGIARYIEQATVHSSMNEEMLEE GQEYAVMLYTWRSRAIPQVKCNE QPNRVEIYEKTVLEPEVTKLMNF MYFQRNAIERFCGEVRRCLHAERRK DFVSEAYLITLGGKFINMFAVLDLKN MKCSVKNDHSAYKRAAQFLRKMAD PQSIQESQNLMSFLANHNKIQSLO QQLEVISGYEELLADIVNLCDVYEN RMYLTPSEKHMLLKVMGFGLYLM GSVSNYKLDAAKRNILSKIDKYFKQ LQVVPLFGDMQIELARYIKTSAHYEE NKSRTWCASSSSSPQYNICEQMIQI REDHMRFISELARYSNSEVVTGSGR QEAQKTDAEYRKLFDLALQGLQLLS QWSAHVMEVYSWKLVPHTDKYSNK DCPDNAEEYERATRYNTTEEFKAL VEVIAMIKGLQVLMGRMESVFNHAI RHTVYAALQDFSOVTLREPLRQAIKK KKNVQSVLQAIRKTVCDWETGHEP FNDPALRGEKDPKSGFDIKVPRRAV GPSSTQLYMVRTMLESLIADKSGSK KTLRSSLEGPTILDIEKPHRESFFYT HLINFSETLQCCDLSQLWFRFFFL ELTMGRRRIQFPIEMSMPWILTDHILE TKEASMMMEYVLYSLDLYNDSAHYAL TKFNKQFLYDEIEAEVNLCFDQFVY KLADQIFAYYKVMAGSLLDKRLRSE CKNQGATIHLPSPNRYETLLKQRHV QLLGRSIDLNRITQVSAAMYKSLE LAIGRFESEDLTSVVELDGLLEINRM THKLLSRYLTLDSFDAMFREANHNH SAPYGRITLHVFWELNYDFLPNYCY NGSTNRFVRTVLPFSQEFQDKQPN AQPYLHGSKALNLAYSSYGSYRNF VGPPHFQVICRLLGYQGIAMVMEELL KVVKSLQGTILQYVKTLMVMPKIC RLPRHEYGSPGILEFFHHQLKDIVEY AELKTVCFQNLREVGNVAVLFCLLIE QSLSLEEVCDLLHAAPFQNLPRIHV KEGERVDAKMKRLESKYAPLHLVPL IERLGTQQIAIAREGDLLTKERLCC GLSMFEVILTRIRTFDDPIWRGPLP SNGVMHVDECVEFHRLWSAMQFV YCIPVGTHEFTVEQCFCGDGLHWAGC MIIVLLGQORRFVAVLDFCYHLLKVQ KHDGKDEIKNVPLKMKMVERIRKFI LNDEIITLDKYLKSGDGESTPVEHV RCFQPPIHQSLASS	False	False	2.443	2.355	1.861	0.998	0.781	1.729	1.291