	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	extracellul region
0	Q8C120	SH3R3_MOUSE	Mus musculus	E3 ubiquitin-protein ligase SH3RF3	47.784353	T600	S395;S792	22517741;34678516;40885482	MLLGASWLCASKAAATAARGEGED RQGEQQRGAQARTEEDMDESSLLD LLECSVCLERLDTTAKVLPCQHTFC RRCLESIVCSRHELRCPECRILVGCG VDELPANILLVRLLDGIRQRPRTGAS PGSSPPARPGPGTFSALAGGAGGAT GSPPCSPVFLSAAAGSSTSSLCDVAT NRSVPVAKTLSQLPYAKALYSYEGKE PGDLKFNKGDIIILRRKVDENWYHG ELQGMHGFLPASYIQCVRPLPQALP QGKALYDFEMKDRDQDKDCLTFTK DEVLTVIRRVDDNWAEGMLGDKIGI FPLLYVELNDSAKQLIEMDKLCPAAT TAYNYDALLSSDPSTVASVAPGPTLS SSGAVSAFQRRVDSKKNAKKRHSFT ALSVTHKSSQAASHRHSMEISAPVLI SSSDPRAAARIGELAHLSCTVPTQDS SSAGPVPTALPRAAAVAGEQGMSPK VQLPLNVYLALYAYKPQKNDELELR KGEMYRVLEKCQDGWFKGASLKTG VSGVFPGNYVTPVSRVPGGGAGLPW NNVLGGSPLAKGMATIMHPGGGSL SSPATAARSALPLTTLQDHMQHPAT SLPTGSCLRHSAQPTASQAGDTTIPT ATHASAQALDRPTATVSPLRTQTSPS RLPSTGLRPRSVASPQHGQQSPAQM CPRPAIPFTSAASAITPPNVSAANLSG EVGGTPISGLSTPSLINTGFKPDDKK NEKKEKKSGLLKLLAGASTKKKSRS PPSVSPTHDPQSAMDTSLQGAMGP EVSPLTVHGRAGSCPIESEMQGAIGL EPLHRKAGSLDLNFSLSPSRQATLS MASIRPEPKPLPRERYRVVVSYPPQS EAEIELKEGDIVFVHKKHEDGWFKG TLQRNGRTGLFPGSFVESF	False	False	2.245	0.834	0.658	False	False	False	False