

| 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 |
|--------------|--------------|--------------|------------------------------------|--------------|---------------|-----------------------|----------|--|---------------|---------------|---------|---------|---------------|-----------------------|-----------------|-----------------|----------------------|
| E9QQ33 | E9QQ33_MOUSE | Mus musculus | E3 ubiquitin-protein ligase SH3RF1 | 26.088944 | S523 | NaN | 30016717 | MDESALLDLLECPVCLERLDASAKV LPCQHTFCRCLLGIVGSRNELRCP ECRTLVGSGVDELPSNILLVRLLDGI KQRPWKPGGGGGTTCTNTRAQ GSTVVNCGSKDLQSSQCGQPRVQ AWSPPVRGIPQLPCAALYNYEGKE PGDLKFSKGDIIILRRQVDENWYHG EVSGVHGFFPTNFVQIHKLPQPPPQ CKALYDFEVKDKKADKCLPFAKDD VLTVIRRVNENWAEGMLADKIGIFPI SYVEFNAAKQLIEWDKPPVPGVDT AECPSATAQSTSASKHPDTKKNTRK RHSFTSLTMANKSSQGSQNRHSME ISPPVLISSNPTAAARISELSGLSCS APSQVHIITGLIVTPPPSPVTTGPA FTFPSPVYQAAALGSMNPLPPPPLL AATVLAATPSGATAAVAAAAAAAAAAG MGRPVMGSSEQIAHLRPQTRPSVY VAIYPYTPRKEDELELRKGMFLVFE RCQDGWYKGTSMHTSKIIVFPNGY VAPVTRAVTNASQAKVSMSTAGQAS RGVTMVSPSTAGGPTQKPQNGVA GNPSVVPTAVVSAHIQTSPOAKVLL HMSGQMTVNQARNAVRTVAHNSQ ERPTAAVTPIQVQNAACLGPASVGLP HHSLASQPLPPMAGPAAHGAASVIS RTNAPMACAAGASLASPNMNTSAVLE TEPSGRTVTILPGLPTSPESAASACG NSSAGKPKDKSKKEKGLLKLKLSGA STKRKPRVSPASPTLDVELGAGEAP LQGA VGPELPLGGSHGRVGSCTDG DGPVAAGTAALAQDAFHRKTSLLDS AVPIAPPPRQACSSLGPMNEARPV VCERHRVVVSYPPQSEAELELKEGDI VFVHKKREDGWFKGTLQRNGKTGL FPGSFVENI | True | False | 4.278 | 3.608 | 1.102 | 1.041 | 3.437 | 1.447 | 1.093 |