

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	g
Q7Z6J0	SH3R1_HUMAN	Homo sapiens	E3 ubiquitin-protein ligase SH3RF1	15.732757	T505;T509;S519;T520;S525;T529;S532;S551;S553;S561	S304;S532;S735	28657654;35132862;35254053;37217939;34725712;23301498;34019948;29237092;30059200;32574038;38665916;36240223;30620550;39534244;38253038;39531497;33214551;31492838;35138101	MDESALLDLEPCVCLERLDASAKV LPCQHTFCKRCLLGVGSRNELRCP ECRTLVGSGVEELPSNILLVRLLDGI KQRPWKPGPGGGSGTINCTNALRSQ SSTVANCSSKDLQSSQGGQQPRVQS WSPPVVRGIPQLPCAALYNYEGKEP GDLKFSKGDIIILRRQVDENWYHGE VNGIHGFFPTNFVQIHKPLPQPPQC KALYDFEVKDKKADKDCLPFAKDDV LTVIRRVNENWAEGLMADKIGIFPIS YVEFNAAKQLIEWDKPPVPGVDAG ECSSAAAQSSSTAPKHSDTKKNTKKR HSFTSLTMANKSSQASQNRHSMEIS PPVLISSSNPTAAARISELSGLSCSAP SQVHISTTGLIVTPPPSSPVTTGPSFT FPSDVPYQAAALGTLNPPPLPPPLAA TVLASTPPGATAAAAAAGMGPRPMA GSTDQIAHLRPQTRPSVYVAIYPYTP RKEDELELRKGMFLVFERCQDQGW FKGTSMHTSKIGVFPNGYVAPVTRA VTNASQAKVPMSTAGQTSRGVTMV SPSTAGGPAQKLOQNGVAGSPSVVP AAVVSAAHIQTSPOAKVLLHMTGQM TVNQARNAVRTVAAHNQERPTAAVT PIQVQNAAGLSPASVGLSHSLASP QPAPLMPGSATHTAASISRASAPLA CAAAAPLTSPSITSASLEAEPGRIVT VLPGLTPSPDSASSACGNSSATKPKD DSKKEKKGLLKLKSGASTKRRKRVSP PASPTLEVELGSAELPLQGAVGPELP PGGGHGRAGSCPVDGDGPVTTAVA GAALAQDAFHRKASSLDSAVPIAPPP RQACSSLGPVLNESRPVVCERHRVV VSYPPQSEAELELEKGDIVFVHKRE DGWFKGTLQRNGKTGLFPGSFVENI	True	False	5.0	3.67	1.215	1.167	3