

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
Q96FS4	SIPA1_HUMAN	Homo sapiens	Signal-induced proliferation-associated protein 1	24.745947	NaN	T64;S67;S182;S304;S314;S817;S839;S912	26853435;37217939	MPMWAGGVGSPRRGMAPASTDDL ARKLRQPARPPLTPHTFEPRPVRG LRSQSDAGEARPPTPASPRARAH EEASRPAATSTRLFTDPLALLGLPA EPEPAFPVLEPRWFVFAHYDVQSL DWAPRSQGMGSHSEASSGTLASAE DQAASSDLLHGAPGFVCELGEGE GLGGPASPPVPPALPNAAVSILEEP NRTSAYSLEHADLGAGYRKYFYGK EHQNFPGMDESLGPAVAVSLRREE EGSGGGTLHSYRVIVRTTQLRTRG ISEDALPPGPPRGLSPRKLLEHVA LSPSCLRLGSASPVKVPTLLTLDEQ LSFQRKVGILYCRAGQGSEEMYN QEAGPAFMQFLTLLGDVVRKGFES YRAQLDTKTDSTGTHSLYTTYQDHE MFHVSTMLPYTPNNQQQLLRKRI GNDIVTIVFQEPGSKPFCPTTIRSH QHVFVVRHAHTPCTPHTTYRVAVSR TQDTPAFGPALPAGGGPFAANADFR AFLAKALNGEQAAGHARQFHAMA TRTRQQYLQDLATNEVTTTSLDSAS RFGLPSLGGRRRAAPRPGAELOAA GSLVWGVRAAPGARVAAGAQAQSGP GIEVPCLLGISAEALVLVAPRDGRV FNCACRDVLAWTFSEQLDLYHGR GEAITLRFDGSPGQAVGEVVARLQL VSRGCETRELALPRDQGRLGFEVD AEGFVTHVERFTFAETAGLRPGARL LRVCGQTLPSLRPEAAAQLLRSAPKV CVTVLPPDESGRPRRSFSELYTSLQ EPSRRGAPDPVQDEVQGVTLPTTK QLLHLCLQDGGSPPGDLAEERTE FLHSQNSLSPRSSLSDEAPVLPNTTP DLLLATTAKPSVPSADSETPLTQDRP GSPSGSEDKGNPAPELRASFLPRTLS LRNSISRIMSEAGSGTLEDEWQAISE IASTCNTILESLSREGQPIPEGDPKG TPKSDAEPEPGNLSEKVSHLESMLR KIQEDLQKEKADRAALEEVRSLRH NNRRLQAESESAATRLLLASKQLGS PTADLA	True	False	4.629	4.233	1.004	0.689	3.769	4.083