	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	P46062	SIPA1_MOUSE	Mus musculus	Signal-induced proliferation-associated protein 1	25.643919	NaN	T62;S65;S178;S299;S309;S812;S834;S907	28135057	MWAGGVGSPRRGMAPAPTDDLFAR KLRQPARPPLTPNTFEPRPARGPLLR SGSDAGEVRPPTPASPRARAHSHED ASRPAATPTRLFTDPLALLGLPAEEP EPTFPPVLEPRWFAHYDVQSLLFDW APRPRGTGSHTEANSGTLAEGQTTT SDLLLGAPGFVSELGGEGELGLGGPI SPPVPPALPNAAVSVLEEPQTRTTTY SLEHADLGAGYYRKYFYGKEHQNFF GLDEALGPVAVSLRREEKEGSGGGT LHSYRVIVRTTQLRTLRGTISEDALPP GPPSVSPRKLLEHVAPRLSPTCLRLG SASPKVPRQLLTLDEQVLSFQRKGGI LYCRAGQGSEEEMYNNQEAGAAFM QFLTLLGDVVRLKGFESYRAQLDTK TDSTGTHSLYTTYQDHEIMFHVSTM LPYTPNNQQQLLRKRHIGNDIVTIVF QEPGSKPFCPTTIRSHFQHVFLVVRA HAPCTPHTSYRVAVSRTQDTPAFGP ALPEGGGPFAANADFRAFLLAKALN GEQAAGHARQFHAMATRTRQQYLQ DLATNEVTTTSLDSASRFGLPSLGGR RRATPRSPGADVQAAGALMWGVRA APGARVAAGAETSGPDDAEVPCLLGI SAETLVLVAPRDGRVVFNCACRDVL AWTFSEHQLDLYHGRGEAITLRLDG APGQAVGEVVARLQLVSRGCETREL ALPRDGQGRLGFEVDAEGFITHVER FTFAETTGLRPGARLLRVCGQTLPKL GPETAAQMLRSAPKVCVTVLPPDES GRPRRSFSELYMLSLKEPSRRGGPE PVQDETGKLVILPPTKQLLHFCLKDS SSPPGPGDLTEERTEFLRTHNSLSS GSSLSDEAPVLPNTTPDLLLVTTANP SAPGTDRETPPSQDQSGSPSSHEDT SDSGPELRASILPRTLSLRNSISKIMS EAGSETLEDEWQSISEIASTCNTILE SLSREGQPISESGDPKEALKCDSEPE PGSLSEKVSHLESMLWKLQEDLQR EKADRAALEEEVRSLRHNNQRLLAE SESAATRLLLASKHLGAPTTDLA	None	None	None	None	None	None	None	None	None