

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q8CH02	SUGP1_MOUSE	Mus musculus	SURP and G-patch domain-containing protein 1	24.992371	T349	T128;S252;S322;S407;S409;S412;S483	36852467	MSLKMDNRDVAGKANRWFQMAQP KSGKMNMNLIHQEELIAQKKREIEA RMEQKARQSHVPSPPHPGEIADA HNSCISNKFANDGSFLQFLKLQKA QTSTD SAPRAPPSPMTPSSLKKPLVL SKRTGLGLSSPTGPVKNYSHAKQLP VAHRPSVFQSPDDDEEEDYEQWLEI KVSPPEGAETRRVIEKLARFVAEGGP ELEKVAMEDYKDNPAFTFLHDKNS REFLYRRKVAEIRKEAQKQAATQK VSPPEDEEAKNLAEKLARFIADGGPE VETIALQNNRENQAFSFLYDPNSQG YRYRQKLDEFKAKAGSTGSPFAPA PNPSLRKSAPEALSGAVPPITACPT PVAPAPAVNPTPSIPGKPTATAAVKR KRKSRWGPEEDKVELPPAELAQRDI DASPSPLSVQDLKGLGYEKGPVGL VGVTELSDAQKKQLKEQQEMQMY DMIMQHKRAMQDMQLLWEKALQQ HQHGYDSDEEVDSELGTWEHQLRR MEMDKTREWAEQLTQMGRGKHF GDFLPPDELEKFMETFKALKEG DYSEYKEFKLTVENIGYQMLMKMG WKEGEGLTGEGQIKNPVNGATTI DGAGFGIDRPAELSKEDDEYEAFRK RMMLAYRFRPNPLNPRRPPY	True	False	2.235	4.429	1.299	0.77	0.438	1.154	1.325