

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P97363	SPTC2_MOUSE	Mus musculus	Serine palmitoyltransferase 2	26.066638	NaN	NaN	28135057;28528544	MRPEPGGCCRRPMPRANGCVKNGE VRNGYLRSSTATVAAAGQIHHVTEN GGLYKRPFNEAFEETPMLVAVLTYV GYGVLTFLFGYLRDFLRHWRIEKCHH ATEREEQKDFVSLYQDFENFYTRNL YMRIRDNWNRPICSVPGAKVDIMER KSHDYNWSFKYTGNIKGVINMGSY NYLGFARN TGSCQEA AAEVLKEYGA GVCSTRQEIGNLDKHEELEKLVARF LGVEAAMTYGMGFATNSMNIPALV GKGCLLSDELNHA SLV LGARLSGAT IRIFKHNNMQSLEKLLKDAIVYGQPR TRRPWKKILILVEGIYSMEGSIVRLPE VIALKKKYKAYLYLDEAHSIGALGPS GRGVVDYFGLDPEDVDVMMGTFTK SFGASGGYIGGKELIDYL RTHSHSA VYATSMSPVMEQIITSMKCMIGQD GTSLGKECIQQLAENTRYFRRLKE MGFIYGNEDSPVPLMLYMPAKIGA FGREMLKRNI GVVVVGFPATPIIESR ARFCLSAHTKEILDALKEIDEVGD LLQLKYSRHR LVP LLD RPFDETTYEE TED	True	True	2.931	2.599	4.744	4.537	2.401	2.48	2.188