

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	extracellular region
0 Q9JLJ5	ELOV1_MOUSE	Mus musculus	Very long chain fatty acid elongase 1	44.377775	NaN	NaN	40997131;25153642	MEAVVNLYHELMKHADPRIQSYPL MGSPLLITSILLTYVYFILSLGPRIMA NRKPFQLRGFMIVYVNFSLVILSLYIV YEFLMSGWLSTYTWRCDPIDFSNSP EALRMVVRVWLFMLSKEIEMDTVI FILRKKDQVTFHVFHHSVLPWSW WWGIKAPGGMGSFHAMINSSVHV VMYLYYGLSALGPVAQPYLWKKH MTAIQLIQFVLVSLHISQYFMPSCN YQYPIIIHLIWMYGTIFILFSNFWYH SYTKGKRLPRAVQQNGAPATTKVKA N	False	False	1.727	1.707	1.872	3.907	1.168	3.577	1.655