

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	ex re
Q5Y5T5	ZDHC8_MOUSE	Mus musculus	Palmitoyltransferase ZDHC8	29.185005	NaN	S335;S603;S624;S672;S679;S722;S740	22645316	MPRSPGTRLKPAKYIPVATAAALLVG SSTLFFVFTCPWLTRAVSPAIPVYNG ILFLFVLANFNSMATFMDPGVFPFRAD EDEDKEDDFRAPLYKNVDVRGIQVR MKWCATCHFYP RPCSHCSVCDCNC VEDFDHHCWPVNNCIGRRNYRYFF LFLLSLAHMGVAVFGLLYVLNHS EGLGAAHTTITMAVMCVAGLFFIPVI GLTGFHVVLVTRGRITNEQVTGKFR GGVNPFFTRGCGNVEHVLCSPLAPR YVVEPPRMPLSVSLKPPFLRPELLER AVPLKVKLSDNGLKAGRSKSKGSLD QLDEKPLDLGPPLPKIEAGTFGRDL KTPRPGSAESALSVQRTSPPTPAMYK FRPAFSTGPKTPFCGPNEQVPGPDS LTLADDSTHSLDFVSEPSLDLDPDHG PGGLRPPYPSPPLNTTDAFSGALRS LSLKAASRRGGDHMTLQPLRSEGGP PTPHRSLFAPHALPNRNGSLSYDSL LNPGSPSGHACPTHPSVGIASYHSPY LHPGPSDPPRPPRSFVLPGRPRE PSPVRYDNLRSRTIMASIQERKDREER ERLLRSQTDLSFGDSGVYDTPSSYSL QQASVLTGPRGSVLRYSRDDLVA GPGFGGARNPALQTSLSLSSMSR APRTSSSLQADQANNAPGPRPGS GSHRSPARQGLPSPPGTSPSPSYTGS KAVAFIHTLDRQPSLAMQRDHPQ LKTTPSKLNGQSPGMARLGAASPM GPNASPARHTLVKKVSGVGGTTYEIS V	True	False	1.379	1.153	5.0	1.896	3.836	2.53	1.6