

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
Q8CEF9	T132C_MOUSE	Mus musculus	Transmembrane protein 132C	24.474581	NaN	NaN	33300544	MRSEGAAPRRAARYGALSLVLATLL GQVTESRGVMNDNIQRFSSLPYLPV SFHVLRAETAFFLKEANPDPLRNAS LQSRVESFFIYKAQQPPVLNVSYGPY SAEKVIPLDLMLNPNFLGPTSKFPFD WRLKAYILQEKVYLSHPKVQVLFHIV GRDWDHHRDEKLPCLRVFAFRDSR EVRGSCRLGGPLGLCVAQLEMLPG WFSPPAVVSGRRRPAERPEGSPVEL YYAVQPGDERGDCGTGGDTRKDNAIR PGKDGQEGRTSHLQKIGTISLYRAQD SNQSELRLDGNVVIWLPSPVKQG DIVTASVTIANNSTVDHFILRAKVKK GVNILTVOITSEPROWDVROEVGNG GKHHTTSVACQRLGPGARNRSSLNF SEVMQMNFEIASFSSLSCTQPITWO VEYPRKGATDIAVSEIFISQKDLVAIV PLAMDTELLNTAILTGKTVAMPVRV VSVEENSTLRDISELVECKATDENVI KVS DHCDYVFNKGKIKGKMDSVV NFTYQHLSAPLHVTVVWVRLPLQIE VSDTELSQVKGWRVPIVASKRPTRD SEEEEEEQKGRGCTLQFQHATVRV LTQFVSEGAGPWGQLSHLLSPDWQ FDITHLVADFMKLESPHIATLQDSRV LVGREVGMTTIQVLSPLSDSILAET VTVLDDKVSVDLAVQVAVGLSVTL HPISENNKATSAVAMAEELLRAPKK EAIISTWLQFSDGSVTPLDIYDSKDF SLTAISLDEAVVISIPQLSPWWPTVV AEGEGQGPLLVRVMSIAEACQKSKR KSVLAVGIGHVGVKFGWDDADSSQT GEKDEEEIKNHASDRRQKIQLDERP GQDELYHGNFPGDREEGALSATTTT KSLLDNNVGKSGRRD GARLHSIPID FTNFPAHVDLPKAKTRGTLEENGLM QTAHGLSDLEIGMYALLGVFCLAILV FLINCATFAFKYRHKQVPLEGQASM THSHDWVWLGNEAELLENIGDLSP PQDEHTTIIDRGLGCENHLLLN GGSQKPTQSQVHRPPGSGGRQTRREP RQEPANSPTSKMKKVKFATFTIPPEE SCPTVNSILSGEDDIKWVCQDLVVG APKELRTYLEKFQDSV	None	None	None	None	None	None	None		