

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
Q8CDI6	CD158_MOUSE	Mus musculus	Coiled-coil domain-containing protein 158	25.019509	S180;T927	NaN	34418053	MESKACESKNEDLLPSGITSKGGSS SPFFVTSTHGTHIENTSSGTLTQMP FFPKYEVELDSPRKSTPYPGKEHIER VLEEYSHQVKDLQRRLNESNELHEK QKFYLRQSVIDLQTKLQEMQMERDA MADIRRRESQSQEESRNQLQNTVR ELEAAKCLKEDMLKDSSTOIEQLRK MMLSHHEGLQEIRSLVDFEESAGK KICEHDSMSTMHFRSLGSAISKILRE LDTEISFLKGRIFPVEDQLETLSKES QNKIELLLQQHQDRIEQLISEHEVEI TGLTEKASSARSQANSVQSLEIIEQ QARNQNSMYMRQLSDLESTVSQLR SELRESKRMVEDKIEELEKQLVLAN SELTEARTERDQFSQESGNLDDQLQ KLLADLHKREKELSLEKEQNKRLWD RDTGNSITIDHLRRELDDRNMEVQR LEALLKAMKSECQGMERQMAAIQ GKNESLEKVSSTAQLESTKEMLRK VVEELTAKKMNLESSERTVSDLTAS LQEKERAIEATNAEITKLSRVDLKL QELQHLKNEGDLRNVQTECEALK LQMAEKDKVIEILRQIENMTQLVG QHGRTAGAMQVEKAQLEKEINDRK LELQEFKILKDKKDAKIRELEARVSD LELEKVLV NAGSERLRAVKDIRHE RDQLLNEVKTSRTELNLHLSSEYEV KRNFRNKSEEMESTTNRLKMQLKS AQSELEQTRNTLKTMEGSDGHAMK VAMGMQKQITAKRGQIDALQSKVQF LEEAVTSANKERHFLKEEKSLSQE LSTVATEKNKMAGELEVLRSQERRL KEKVANMEVALDKFAECODIIQRQE QESVRLKQHTLDVKELQGPYTSN SSVKPRLQPASVTRSHSNIPSSQST TSFLSHHSIKTNTPKEDPTRDLKQLL QELRTVINEEPAMALSKTEEDGRTP SLGALEDRVRDCITESLRAELCHRS NNSLRESTEGSKSSETLSREPVPLH PGDLEDPSSCFTFTSTASPSGKMSA SRSFSSSPKSPVHSLTSSAEESVN STPQYRSTKPIHSPTS AKDSQSPSLE TTGKTCQKLQNRLESLOTLVEDLQL KNQAMSSMIRNQEKRQKVKDQEK MLLK	None	None	None	None	None	None	None		