

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
P57080	UBP25_MOUSE	Mus musculus	Ubiquitin carboxyl-terminal hydrolase 25	25.685855	NaN	S85;T740	39627609	MTVEQNVLQSSAAQKHQQTFLNQL REITGINDAQILQQALKDSNGNLELA VAFLTAKNAKTPPQEETGYQTALPG NDRYISVGSQADANVIDLTGDDKDD LQRAIALSLAENRAFRETGITDEEQ AISRVLEASIAENKACLKRTPIEVWR DSRNPYDRKRQEKAPVGLKNVGN CWFSAVIQSLFNLLEFRRLVLYNYP SNAODLPRNQKEHRNLPFMRELRY LFALLVGTKRKYVDPSPRAVEILKDAF KSNDSQQQDVSEFTHKLLDWLEDA FQMKAEETDEEKPNMVELFYG RFLAMGVLEGGKFFENTEMFGQYPL QVNGFKDLHECLEAAMIEGEIESLH SDNSGKSGQEHWFTELPVLTFFELS RFEFNQALGRPEKIHNKLEFPQVLY LDRYMHRNREITRIKREIKRLKDY TVLQORLERYLSYSGPKRFPVLDVL QYALEFASSKPVCTSPVDDIDASSA SGPLPSQSLPSTTEQQGPCASDLPGS SSPASGAALPLRSVIHKPFTOSRIPPD LPMHPAPRHITTEELCVLESCLRHW RTEIENDTRDLQESISRHRTELMY DKSMIQVPYRLHAVLVHEGQANAG HYWAYIFDHRESRWMKYNDIAVTKS SWEELVRDSFGGYRNASAYCLMYID DKAQFLIQEEFNKETGQALVGMETL PPDLRDFVEEDNQRFEKELEEWDT QLAQRSLQEKLLAAPKLRREAEASAT AQAGGADYLEQPSRSDLSKHWKEET LRVIKASHDLEDKGPETVLSAIDL EYSRLVKLAQEDTPPETDYRLHHVL VYFIQNQAPKKIIEKTLLEQFGDRNL SFDERCHNIMKVAQAKLEMIPPEV NLEEYEEWHADYKFFRETTMYLITG LENFQRESYIDSLFLLCAYQNNKEL LSKGPYRGHDGELISHYRRECLLKL NEQAAELFESGEDGDVNNGLIMNE FIVPFLPLLVDMEEKDILAVEDMR NRWCSYLGQEMEANLQEKLTDFLP KLLDCSTEIKGFHEPPKLPYSAAHEL CERFARIMLSLRTPADGR	True	False	2.695	4.013	2.194	1.22	1.065	1.504	1.285