

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membran
Q61699	HS105_MOUSE	Mus musculus	Heat shock protein 105 kDa	23.373761	NaN	S509;S510;S558;T562;S810;T816	37453647;39627609;37507081	MSVVGLDVGSQSCYIAVARAGGIETI ANEFSDRCTPSVISFGSKNRTIGVAA KNQOITHANNVTSSFKRFHGRAFN DPFIQKEKENLSYDLVPMKNGGVGI KVMYMDEEHFFSVEQITAMLLTKLK ETAENNLKPVVTDVVISVPSFFTDAE RRSVLDAAQIVGLNCLRLMNDMTA VALNYGIYKQDLPNAEEKPRVVVFV DMGHSSFQVSACAFNKGKLVLT AFDPFLGGKNFDEKLVEHFCAEFKT KYKLDAKSKIRALLRLHQECEKLLKL MSSNSTDLPLNIECFMNDKDVSGK MNRSQFEELCAELLQKIEVPLHSLM AQTOLKAEDVSAIEIVGGATRIPAVK ERIAKFFGKDVSTTLNADEAVARGC ALQCAILSPAFAKREFSVTDVAVPPIS LVVNHSEETEVEGVHEVFSRNHAAP FSKVLTFLLRRGPFELEAFYSDPQGV YPEAKIGRFVVQVNSAQKDGKESRV KVKVRVNTHGIFTISTASMVEKVPT EEDGSSLEADMECPNQRPTSSD KNIQDDNSEACTQPQVQTDGQOTS QSPPELTSEESKTPDADKANEEK VDQPPEAKPKIKVVNVELPVEANLV WQLGRDLLNMYIETEGKMIMQDKL EKERNDAKNAVEECVYEFRDKLCGP YEKFICEQEHEKFLRLTETEDWLYE EGEDQAKQAYIDKLEELMKGMPVK VRFQEAERPKVLEELQRLQHYAK IAADFRGKDEKYNHIDSEMKKVEK SVNEVMEWMNVMNAQAKRSLDQ DPVVRTHEIRAKVKELNNVCEPVVT QPKKIESPKLERTPNPNDKEDL EGKNNLGAEAPHQNGECPNEKGS VNMDLD	True	False	4.242	5.0	2.132	2.024	1.16	1.616