

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
P11499	HS90B_MOUSE	Mus musculus	Heat shock protein HSP 90-beta	40.528221	S434;S452;S462	S226;S255;S261;T297;T301;T305;S307;S445;T479;T484;S669;S718	22556278;33300544;22645316;29187734;34887587;24788674;37453647;36288343;39627609;37507081	MPPEVHHGEEEVETFAFQAEIAQLM SLIINTFYSNKEIFLRELISNASDALD KIRYESLTDPSKLDGKELKIDIIPNP QERTLTLVDTGIGMTKADLNNLGTI AKSGTKAFMEALQAGADISMIGQFG VGFYSAYLVAEKVVVITKHNDDEQY AWESSAGGSFTVRADHGEPIGRGTK VILHLKEDQTEYLEERRVKEVVKKH SQFIGYPITLYLEKEREKEISDDEAEE EKGEKEEEDKEDEEKPIEDVGSDE EDDSGKDKKKTKKIKEKYIDQEEL NKTKPIWTRNPDDITQEEYGEFYKSL TNDWEDHLAVKHFSVEGQLEFRAL LFIPRRAPFDLFENKKKKNIKLYVR RVFIMDSCELIPEYLNFIKRVVDSE DLPLNISREMLQQSKILKVIKRNIVK KCLELFSLEAEDKENYKFFYAFSK NLKLGIHEDSTNRRRLSELLRYHTS QSGDEMTSLSEYVSRMKETQKSIYYI TGESKEQVANSFAFVERVRKRGFEVV YMTEPIDEYCVQQLKEFDGKSLVSV TKEGLELPEDEEEKKKMEESKAKFE NLCKLMKEILDKKVEKVTISNRLVSS PCCIVTSTYGWTANMERIMKAQALR DNSTMGYMMAKKHLEINPDHPIVE TLRQKAEADKNDKAVKDLVLLFET ALLSSGFSLEDPQTHSNRIYRMKLG LGIDDEVTAEPSSAAMPDEIPPLEG DEDASRMEEVD	True	True	4.374	4.687	4.653	3.373	2.507