	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	P02827	HSP70_XENLA	Xenopus laevis	Heat shock 70 kDa protein	26.606588	NaN	NaN	17329255	MATKGVAVGIDLGTTYSCVGVFQHG KVEIIANDQGNRTTPSYVAFTDTERL IGDAAKNQVAMNPQNTVFDAKRLIG RKFNDPVVQCDLKHWPFQVVSDEG KPKVKVEYKGEEKSFFPEEISSMVLT KMKETAEAYLGHPVTNAVITVPAYF NDSQRQATKDAGVLAGLNILRIINEP TAAAIAYGLDKGARGEQNVLIFDLGG GTFDVSILTIDDGIFEVKATAGDTHL GGEDFDNRMVNHFVEEFKRKHKKD IGQNKRALRRLRTACDRAKRTLSSSS QASIEIDSLFEGIDFYTAITRARFEEL CSDLFRGTLEPVEKALRDAKLDKSQI HEIVLVGGSTRIPKVQKLLQDFFNGR ELNKSINPDEAVAYGAAVQAAILMG DKSENVQDLLLLDVAPLSLGLETAG GVMTVLIKRNTTIPTKQTQSFTTYSD NQPGVLIQVFEGERAMTKDNNLLG KFELSGIPPAPRGVPQIEVTFDIDAN GILNVSAVEKSSGKQNKITITNDKGR LSKEDIEKMVQEAEKYKADDDAQRE RVDAKNALESYAFNLKSMVEDENV KGKISDEDKRTISEKCTQVISWLENN QLAEKEEYAFQQKDLEKVCQPIITKL YQGGVPGGVPGGMPGSSCGAQARQ GGNSGPTIEEVD	None	None	None	None	None	None	None	None	None