	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	P00815	HIS2_YEAST	Saccharomyces cerevisiae	Histidine biosynthesis trifunctional protein	24.475439	NaN	NaN	33229814	MVLPILPLIDDLASWNSKKEYVSLVG QVLLDGSSLSNEEILQFSKEEEVPLV ALSLPSGKFSDDEIIAFLNNGVSSLFI ASQDAKTAEHLVEQLNVPKERVVVE ENGVFSNQFMVKQKFSQDKIVSIKK LSKDMLTKEVLGEVRTDRPDGLYTT LVVDQYERCLGLVYSSKKSIAKAIDL GRGVYYSRSRNEIWIKGETSGNGQK LLQISTDCDSDALKFIVEQENVGFCH LETMSCFGEFKHGLVGLESLLKQRL QDAPEESYTRRLFNDSALLDAKIKEE AEELTEAKGKKELSWEAADLFYFAL AKLVANDVSLKDVENNLNMKHLKV TRRKGDAKPKFVGQPKAEEEKLTGPI HLDVVKASDKVGVQKALSRPIQKTS EIMHLVNPIIENVRDKGNSALLEYTE KFDGVKLSNPVLNAPFPEEYFEGLT EEMKEALDLSIENVRKFHAAQLPTE TLEVETQPGVLCSRFPRPIEKVGLYIP GGTAILPSTALMLGVPAQVAQCKEIV FASPPRKSDGKVSPEVVYVAEKVGAS KIVLAGGAQAVAAMAYGTETIPKVDK ILGPGNQFVTAAKMYVQNDTQALCS IDMPAGPSEVLVIADEDADVDFVASD LLSQAEHGIDSQVILVGVNLSEKKIQ EIQDAVHNQALQLPRVDIVRKCIAHS TIVLCDGYEEALEMSNQYAPEHLILQ IANANDYVKLVDNAGSVFVGAYTPE SCGDYSSGTNHTLPTYGYARQYSGA NTATFQKFITAQNITPEGLENIGRAV MCVAKKEGLDGHRNAVKIRMSKLG LIPKDFQ	None	None	None	None	None	None	None	None	None