

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P00815	HIS2_YEAST	Saccharomyces cerevisiae	Histidine biosynthesis trifunctional protein	9.365641	NaN	NaN	33229814	MVLPILPLIDDLASWNSKKEYVSLVG QVLLDGSSLSNEEILQFSKEEEVPLV ALSLPSGKFSDDDEIIAFLNNGVSSLFI ASQDAKTAEHLVEQLNVPKERVVVE ENGVFSNQFMVKQKFSQDKIVSIKK LSKDMLTKEVLGEVRTDRPDGLYTT LVVDQYERCLGLVYSSKKSIAKAIDL GRGVYYSRSRNEIWIKGETSGNGQK LLQISTDCDSALKFIVEQENVGFCH LETMSCFGEFKHGLVGLSLLKQRL QDAPEESYTRRLFND SALLDAKIKEE AEELTEAKGKELSWEAADLFYFAL AKLVANDVSLKDVENNLNMKHLKV TRRKGDAKPKFVGQPKAEEEEKLTGPI HLDVVKASDKVGVQKALSRPIQKTS EIMHLVNPPIENVRDKGNSALLEYTE KFDGVKLSNPVLNAPFPPEEYFEGLT EEMKEALDLSIENVRKFHAAQLPTE TLEVETQPGVLC SRFPRIEKVGLYIP GGTAILPSTALMLGVPAQVAQCKEIV FASPPRKSDGKVSPEVVYVAEKVGAS KIVLAGGAQAVAAMAYGTETIPKVDK ILGPGNQFVTA AKMYVQNDTQALCS IDMPAGPSEVLVIADEDADVDFVASD LLSQAHEHGIDSQVILVGVNLSEKKIQ EIQDAVHNQALQLPRVDIVRKCIAHS TIVLCDGYEEALEMSNQYAPEHLILQ IANANDYVKLVDNAGSVFVGAYTPE SCGDYSSGTNHTLPTYGYARQYSGA NTATFQKFITAQNITPEGLENIGRAV MCVAKKEGLDGH RNAVKIRMSKLG LIPKDFQ

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