

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P38321	YB75_YEAST	Saccharomyces cerevisiae	Uncharacterized protein YBR225W	24.475439	NaN	S105	33229814	MGSNKEAKNIDSKNDRGLTSITSNK ISNLKAHDNHTSSMITEHKNADEK GKQEKESRNGTTQSSSSVESHSPQV SHHSDKLSSFDSPLHLPNFRLADDL FNSNSRRSSDSAASSSVSKLKAQLS KIGLHHHTSNNKHSHRSGTPTSEV KANYSPDPSAPRFIVSNMVGNGRGG GGLHGATSNVVKLHRSRKKWDWNT LPASDSSLLIKTVSGNHNLINICIDGE FKQIMYDPNHNELFNRMDLFLSFN MDSPPEDSLIFAKRRLRSYIDFLTKY LESRKYAFECYPFNIENIINIETEVKC FPSFDPLKDYSEIESLIQLWLAQSQK FLLQSNFFFSSEVVEELIKRKPTR QHSNPTISSTSNKISDPTLYIQQLDIE ANSRPFVISDPLDEIDILLIRPLHKTL GGWQLAYDEPSLNIADFALDLSPW MIDSSDNDNAQKNASEIAPEYLTNL QNYLPRKGSRAKIVSDEQEVIENSS NASEYMYDCMNRKFFDDAKERIS RNNFNQGVIEDPLNDQFASRSLSL PSSGADAVKRKKSPTKATKKS GFVN FFKRKHSQLASTSHTTSPSVSPSIS SSSPKIQQSHISSPPRTEKAPHVKS ANQAHQNEWLENFFCRTLNNYKEI DLPTQFILPKEVKRSSNAQLQPEDEP PLSSPISNSDNFSPNEGLDRAKSA IYGKEYLKLRLPFASDTIPAVICPWV WTSLSYYKWKALLREIYRSIIPGGYAL AIVPDLRISNTYYTGILGNADAEKAN NSSEEFLLTKERDKTFDAMAIDAINK GLHIHPTKHLTRTFKDVGFTEGKSSV LSLKTGDFKTDMGFLNEFNLSLDMW DYMLRRQLPDSSCPPKDTDPTTLFK RYVEEHIGKIDDNAGCFRTLYVVAQK PKLPYTK	None	None	None	None	None	None	None	None	

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