	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	Q08831	VTS1_YEAST	Saccharomyces cerevisiae	RNA-binding protein VTS1	24.475439	NaN	NaN	33229814	MKHPYEEFPTGSKSPYNMSRGAHP GAVLLSPQSSAINKNNPGSNSGNNQ GNSSVTANVLSPQSHSMSLNDMLD QQSFMLDTAGTRAQPLQQQQQQQQ QQQQASLPSLNIQTVSSTAAGSAIVS PMMQSPKALQSTLSSTSMYLDSFQR SPNNILGIPSQSGSIPLPQSRQSQQQ SQSQKNDPNMGTNFSQDINQLCSW ISMLNSSQQNTVMDNILSILNDDVL KYTKLKIETLTNTPFISPPLPAIASPIP NRDDTQILNIDSVFSSSPITNDPENT DNLLYQNWSPQPHSIPISQPIYDNIT DASQRSKSAEPHVNSSPNLIPVQKQ FNNGNSTKYKKLPSENPNYLSHSLS SSHSFFQPKKRSNMGNEYNSHHHH SLHHPLHNTTSYFSNTSRPSGTDLN KSNQNVFNNTITHPNAGPTSATSTS TSSNGNTPLSSNSSMNPKSLTDPKL LKNIPMWLKSLRLHKYSDALSGTPW IELIYLDDETLEKKGVLALGARRKLL KAFGIVIDYKERDLIDRSAY	None	None	None	None	None	None	None	None	None