	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	P41696	AZF1_YEAST	Saccharomyces cerevisiae	Asparagine-rich zinc finger protein AZF1	24.475439	NaN	S61;S286;S325	33229814	MPPPTAQFMGPTQAGQNESQNQSS GEAGEQNQEHGQGPTPILNQSQPAS SQPQHQQQRNESISYYTNFNQPRYS TDASINSFLNISDNVPVTSTGGPSSG GAYSNLPRLSTSSTHQPPDLSQIGRG FSIVNNLFPQQQQLQNQHRQQQQQ QQQQSHQQPPFKTPSFSTGLTGSSS QYQFLPRNDNTSQPPSKRNSVYLGP NDGPDFEFFSMQQSQQPQFQPSSR RESNSMRPPLLIPAATTKSQSNGTN NSGNMNTNADYESFFNTGTNNSNS NQNPYFLSSRNNSLKFNPEDFDFQF KRRNSFVRGTLDHSSQNAFIPESRL NSLSVNNKANGDPVADNVTNNMK GKSNEVDNDDGNDSSNNNNNNNN NNNNENNNDNNNDNNDNSINSAT STNIPNQEDHSLASTDTTSNSRKDL KEIEQRLRKHLNDEDNYSSAISRPLD KNDVIEGSEGLNKHIDESGMQPNII KKRKKDDSTVYVKNEMPRTDPPMS KDNSTSAEGAAMANFSGKEPPIPDIS SVSDDATNLIGATKVDQLMLIIQARK KGFTEKVNTTQDGDLLFNQTMDILP PKSELVGGVEKPKGTQNTRAVKKHE CPYCHRLFSQATHLEVHVRSHIGYK PFVCDYCGKRFTQGGNLRTHERLHT GEKPYSCDICDKKFSRKGNLAAHLV THQKLKPFVCKLENCNKTFTQLGN MKAHQNRFHKETLNALTAKLAEMN PSENIPLEERQLLEYFASIYKNSNRGI KGRGKGVGTKKSTISSPENHPASTIL NPNTNANNAIANDSENNGNPEGNI DSSSNSNPGSHSMISPTQKDMGTL QSQFIQNNFNNSVNSSNPSNQPIIN YNYTTLPHSRLGSSSSSNTNNNNSN FSVGAAPGVLMAPTTNNDFSFNLDQ SNDNERSQQEQVRFKNINYKS	None	None	None	None	None	None	None	None	None