	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	Q12432	EAF3_YEAST	Saccharomyces cerevisiae	Chromatin modification-related protein EAF3	24.475439	NaN	S201	33229814	MVDLEQEFALGGRCLAFHGPLMYE AKILKIWDPSSKMYTSIPNDKPGGSS QATKEIKPQKLGEDESIPEEIINGKCF FIHYQGWKSSWDEWVGYDRIRAYN EENIAMKKRLANEAKEAKKSLLEQQ KKKKLSTSLGGPSNGGKRKGDSRSN ASISKSTSQSFLTSSVSGRKSGRSSA NSLHPGSSLRSSSDQNGNDDRRRS SSLSPNMLHHIAGYPTPKISLQIPIKL KSVLVDDWEYVTKDKKICRLPADVT VEMVLNKYEHEVSQELESPGSQSQL SEYCAGLKLYFDKCLGNMLLYRLER LQYDELLKKSSKDQKPLVPIRIYGAIH LLRLISVLPELISSTTMDLQSCQLLIK QTEDFLVWLLMHVDEYFNDKDPNR SDDALYVNTSSQYEGVALGM	None	None	None	None	None	None	None	None	None