	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	P38313	CENPU_YEAST	Saccharomyces cerevisiae	Inner kinetochore subunit AME1	24.475439	NaN	S41;S45;S53;S59;S101	33229814	MDRDTKLAFRLRGSHSRRTDDIDDD VIVFKTPNAVYREENSPIQSPVQPILS SPKLANSFEFPITTNNVNAQDRHEH GYQPLDAEDYPMIDSENKSLISESPQ NVRNDEDLTTRYNFDDIPIRQLSSSI TSVTTIDVLSSLFINLFENDLIPQALK DFNKSDDDQFRKLLYKLDLRLFQTI SDQMTRDLKDILDINVSNNELCYQL KQVLARKEDLNQQIISVRNEIQELKA GKDWHDLQNEQAKLNDKVKLNKRL NDLTSTLLGKYEGDRKIMSQDSEDD SIRDDSNILDIAHFVDLMDPYNGLLK KINKINENLSNELQPSL	None	None	None	None	None	None	None	None	None