	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	P46944	TRS85_YEAST	Saccharomyces cerevisiae	Trafficking protein particle complex III-specific subunit 85	24.475439	NaN	NaN	33229814	MVFSYEHYMNLLFHLDNSKETVPP EIAKRIISNAIAPVITVTSTPLFDKHIQ ETYKVDSLYMLLRFFGGCVSDRDQA NEAKVGQHEHEVCDASDSTDSIPKN KNLEVPNLSKKGSRSRSNSLFQRDS TQSQYIRFTRPLGDLIETRDANDMLF NYHSLEVFLDNYLKLVAANTDEMVP HNLLKKSIYHSFFSLAISSTNNLSPY ETFNHPILSLIALDISNGEVYEDARD LLVNFKNLNHNTENFPIFMNTNEM LPVFLLCYNDDSQEEFEKCQALAKK LKKQLFVESILLALWKDSFIYDENSV IQLHQPVMSSLEEILFFLQAPTQTTL SLALINSIYDMLDYLVYDLMIPFMKR KVSFWEETILQPRKSLFNGAKFFKK FMNKNPVNGNHQHNSLTRDSQGN EYFASSSSEFLMRKLADWSMMLSD FKTAYSTYESLMDDLDAFPKYLASCI EWCAVSLLMGAQSIVTVKMIKNDIN PLIERALATYENCSRIQRGKGKESNS LDVTEPVRSYETRCMILASELFLSLS NTWTSTPYAIQYLETILDECKLGPCS QIMVWERLSDCYNLRVDPRIKHRVG AMKKDAKDTEDLRGEHKYSTDHFT DEDILSEGLTRRRKAAFFRLIAAKKW AEQKQWRQVSWCLKDIESTYSEIKF LHGNGLILSKLKNQLNLKDVDSAPR PSEKNLTRTSVSFIG	None	None	None	None	None	None	None	None	None