	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	Q03337	TRS31_YEAST	Saccharomyces cerevisiae	Trafficking protein particle complex subunit 31	24.475439	NaN	NaN	33229814	MSQRIIQPSASDQQFPGKSDGYEYTV GPKQAITSEASTTYIPSRIYSESLLFK RQEASLSAMAFLFQEMISQLHRTCK TAGDFETKLSDYGHNIGIRLLELLNF RASVSPSSLPRASAFLSQNESSSKLS NASNSPGMLANSSTATSASANERLQ EKQTESLSNYITKMRRRDLKILDILQ FIHGTLWSYLFNHVSDDLVKSSERD NEYMIVDNFPTLTQFIPGENVSCEYF VCGIIKGFLFNAGFPCGVTAHRMPQ GGHSQRTVYLIQFDRQVLDREGLRF G	None	None	None	None	None	None	None	None	None