

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q04183	TR120_YEAST	Saccharomyces cerevisiae	Trafficking protein particle complex II-specific subunit 120	9.365641	NaN	S379;S387	33229814	MNILKHFPSYVGPSKIRTLVIPIGHW TRKEFNNAVQKLSEFNEIHLSDVTP DSPIFTPQGFPHGKLFDFLTIDHDD ALELFLYDFEFPRKTFVIIGLVNDYS DPLTNLNFMKKEYPTLISPNIYVASS TPTKELEQTIDTMENVFASSPDMQK NIETIMCDIARNFLTALNSYSSYKH VTLRSPGAIGGNAVLKTTLIRQNSYT SSSSSTPMSAVQSSVSSSSKAGSVTT ASKRLSSFEMTTNSLKRSASLKLATT LSTSENRSQQKSLGRQMKILGNFQL LAGRYVDALNSFVDAITTYKVRDYL WLGSALDGISICFLLLSYLGLSYQIPQ IVSLICPVEKLNFESSSTGISPVDSNS KATASTTASSTPRNSISIAAMQSPRN SIMLSAPALNIDVENINLPLIKCIS DKVLYYYDLSLMHNSEYAPQVVYCE FLLKTLTFMTSCYKSSEFSKDVLDNI VKNQHRALSDIPNSPMFPRFEVYFY SNKLFELQLKEMQVEAQIKIYSTMA EVYRLLGYKRKQLFVLRLLMVALLAT PNKIAWHPDYRTLIDTIELLNINESE AKINVDDPSQSTWLILQKKILQLCIK VSRKINDFEYVAKFSSILITKYTHLLN QSEQDALFKEYIQPSITNESITSYWD PFILREVVINRILSDPTSNEIPLES VSSLESLENRQKTQDINPQEVFNPF KRVQPTSFVSNNSTKVPILVFLVGDK AEFTCRVQNPFKFDFTINDIQLDEEI SEFCEIDRKAVSYSGPYNVKAESIRSI TLPLIHKPTYKKIYEISCLKISILKLPL QKFDIINDSRRSNPVEEEAEYSKCIY GCLKIKILPEQPQLELLSTSKMTRNS WMMLDGTCTDFHITVRNKSLSAIN HIKIIPMNNIEQMLKPDYWKMPDP DLYIMEKQLDWLSKSCVRIKLPVI KPNETITFDLELDNTAVPFNFTGFDL LIEYGMSATDESCIYLKKSIPYEVTL RRTIEVPSMDIIPNELFSSQVENVD WIEYVMSKIRAESNLHSRDFILLLLD FRNSWIDGIKLVQFEDFTSNEYHV EASHTSRIIVPIKKIDYKYNFENTPI PRIYPGRQFIQSGLNEEQTIEMRQKF WCREHIISKLKCNWKLTTDQSVTGS VDFNKFIEKFDHKMVYTIYPGRLFYG VQLLLDEPKVKVGEIINLKIITEPTST CRRKQNSTVNFLDIVIFDSKTSKILP RSNRRILYNGSLTKPISTTKVSEINLE IPIEKGRYEFVCSKSKSNNQDGIIQF

