

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P47035	NET1_YEAST	Saccharomyces cerevisiae	Nucleolar protein NET1	9.365641	NaN	S60;S166;S231;S252;S437;S439;S447;S452;S497;T676;S830;T1042;S1056;S1059	33229814	MYKLQVVLVPPSLQATMPIQFGYGP TIAESSQLLPNRTNMAQSAGDASLQ YANLRSANVSFTPSYFNQSRFRKFL LFTKPTNTLLNLSDEIIDKCEKMYP LQEDIEILSLQDNSGCDLDPDFLVKD VFNVNNIVRVILKNEIDLDDAPVSL YKSVKRSKLNNGSPQSVQPQQQIPS SSGVLRIAKKRPTGTTTTTIRSATN GSMRVSTPLARQIYPPSSKIVSNNS DDEDEDIGERSFLPPPTQPQSPPIRIS SGIDAGKKIKSSIVEEDIVRSATVDP DKTKQORLLSGTPIMSTMTPNRVTL TGQRVVSEHAHKNELVFSASASSSS FANGGTAAVTAQDINRKPPVTTT SGMLKIPEPRISEIEKELKEGPSSPAS ILPAKAAKIPMKKPYLENGENYESDD SSSENQETPETEPHASKASLQRSQS SIADNNGSPVKNSPLGDAMPNVH LAELPKASNTSITKSSNGESWGKQQ EHQPPRKSSLETIVEKKSQAEPGIV EPKRMTNFLDDNQVREKEDTNDKL LEKEILPTIPHNDQPILASSDKSNGT LKSLAGKVSSNNNASKEDGTIINGTI EDDGNDNDEVDTTVRIVPQDS FPKSDLFKMIEGDDTDLPQWFKGK NSRTSGNSKNSKPYTTVLNKDIDNS KPDPRNILPQRTPRSAAKRAAQLLAG AKKNEVPQKSTEDSSSAASTDDESE SGIETDFSSDDDFKRKNMSVPPNNGP KDISLHSLKGSVVPVKDSKIINKEVD EERNDKRDSQKKS AVSESSVTNSKI SEQMAKSFYPNSNKKQNEATKVET KPATQASSFPVVGGSVATKGTTSF NEEGNRKNVTKAKNESAQIDRQQ KETT SRVADLKSANIGGEDLNKKA GSKEPEKASANIQDANDKNN SKEKE DSKSKQVSQKCLKMTDHLKEGNVQ LPKPSANDKLDLAKFTNSKTLVPP GIISNEKNSSANDDDSSSSSGSSTE DESSSSSSSSDEETSTRKARRVVN TPREPVRSSSKIEAPSPSVNKKINAT PDKIPVTQLMDMSSPPSVKSKTTSN PSSILHDLPRKVRPSLSSLDLVS PDKVKEKTSKSNESQTKAPSSSDDE SSSDSDSNSSSDSVSDSSSDSKSES DSDSDGSDSDDGKSFISAKSASAAL GKKKKPSGGFASLIKDFKKK