

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P39083	RGA1_YEAST	Saccharomyces cerevisiae	Rho-type GTPase-activating protein 1	9.365641	NaN	T278;S291;T532	33229814	MASTAPNEQFPSCVRCKEFITGHA YELGCDRWHTHCFACYKCEKPLSCE SDFLVLGTGALICFDCSDSKNCCKG KIDDLAILSSSNEAYCSDCFKCKC GENIADLRYAKTRGLFCLSCHEKLL AKRKYYEKKRRLKKNLPSLPTPVID NGHTDEVSAVLEPKTFSRPASLV NEIPSGSEPSKDIETNSSDIVPHFITG YNDSDDNSGSSKFGSNVSIDVIGPE ENSTEHVNDVKEEAEAPSANMSL NVATDPTLSCKEPPSHSRNLLNKTP LRNSSGOYLAKSPSSYRQGIIVNDSL EESDQIDPPNNSRNASELLTSVLH SPVSVNMKNPKGSNTDIFNTGEISQ MDPSLSRKVLNNIVEETNALQRPVV E.VKEDRSVPDLAGVQQEQA EKYSY SNNSGKGRKISRSLRRSKDLMINL KSRATGKQDSNVKLS PASKVTSRRS QDLMRDND SHTGLDTPNSNSTSLDI LVNNQKSLNYKRFTDNGTLRVTSGK ETALEEQKNHSFKSPSPIDHLLQSPA TPSNVSMYRTPPLDSSLTFDRRNGS SYSNQNY SIPS WQKTPKTQLENSDN FEEQKETLYENSESRNDPSLDKEIVT AEHYLKQLKINLKELESQREELMKEI TEMKSMKEALRRHIESYNSEKNKLY LDSNELSNPPMINEISLGESPPVK HVATASSVARSSVKPKFWKFFSSAK PQTEQSIQGVSTNNTNSIVKSAPVLL SAPSSGSNSGRLEISPPVLQNPNEFS DVRLVPIENDANMGQSKDGEEYLD GSNLYGSSLVARCN YENNEIPMILS VCIDFIESDEENMRSEGIYRKSGSQL VIEEIEKQFSAWKVQNTETPNILTE QDLNVVTGVLKRYLRKLPNPIFTFQI YEPLMRLVKSKKMMENLPFVGGKL SLEAKNSDTYMSSKSALKNILEDLPR EHYRVLRLVSEHIEKVTRYSHWNRM TLYNLALVFAPGLIRDFSGEKDIIDM KERNYIVAFIFGNYKDILT