

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P06105	SC160_YEAST	Saccharomyces cerevisiae	Protein SCP160	9.365641	NaN	T50;S54;S63;S85;S87;S89;S630;S112	33229814	MSEEQTAIDSPSPSTVEGSVETVTTID SPSTTASTIAATAEEHPQLEKKPTPLP SLKDLPSLGSNAAFANVKVSWGPN MKPAVSNSPSPSPSAPSLTTGLGAK RMRSKNIQEAFLLDLQSQLSITKPEL SRIVQSVKKNHDVSVESTLSKNART FLVSGVAANVHEAKRELVKKLTKPI NAVIEVPSKCKASIIGSGGRTIREISD AYEVKINVSKEVNENSYDEDMDDDTT SNVSLFGDFESVNLAKAKILAIVKEE TKNATIKLVVEDEKYLPIYIDVSEFAS DEGDDEEVKVQFYKKS GDIVILGPREK AKATKTSIQDYLLKKNLDEEKVKI PSKFQFLIDAEELKEKYNVIVTFPSTP DDELVSFVGLRDKVGEAITYARSSSK SYVVELDISKAHSKNLTHAKNLIMY FTKYSVLKGLEESHPNVKISLPSIQSL PTAETVTIHISAKSDEANDIKAVRKE LISFVNNIPPSETLVITDLDYELFGGS IKHCLLASESSVAFVQFGDYYPNDN SILLVALTEDEDFKPSIEEIQASLNKA NESLNSLRTKQNNMETKTYEFSEEV QDSLFPSSATWKLIMEDISEQEGH LQIKLHTPEENQLTVRGDEKAAKAA NKIFESILNSPSSKSKMTVNI PANSV ARLIGNKGSNLQQIREKFACQIDIPN EENNNASKDKTVEVTLTGLEYNLTH AKKYLA AEAKKWADIITKELIVPVKF HGSLIGPHGTYRNLQEKYNVFINF PRDNEIVTIRGPSRGVNKAHEELKAL LDFEMENGHKMVINVP AEHV PRIIG KNGDNINDIRA EYGVEMDFLQKSTD PKAQETGEVELEITGSRQNIKDAKR VESIVAEASDFVTEVLKIDHKYH KSI VGSGGHILREIISKAGGEEIRNKSVDI PNADSENKDITVQGPQKFVKKVVEE INKIVKDAENSVTKTIDIPAERKGALI GPGGIVRRQLESEFNINLFV PNKDD PSGKITITGAPENVEKA EKILNEIIR ENFDREVDVPASIEYV SERGAFIQK LRMDLSVNVRF GNTSKKANKLARA PIEIPLEKVC GSTEGENA EKTFTTIEE VGAPTSSEEGDITMRLTYEPIDLSSIL SDGEEKEVTKDTSNDSAKKEEALDT AVKLIKERIAKAPSATYAGYVWGADT RRFNMIVGPGGSNIKKIREAADVIIN VPRKSDKVN DVVYIRG TKAGVEKAG EMVLKSLRR