

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P33332	SEC3_YEAST	Saccharomyces cerevisiae	Exocyst complex component SEC3	9.365641	NaN	T290;S606	33229814	MRSSKSPFKRKSHSRETSHDENTSF FHKRTISGSSAHHSRNVSQGAVPSS APPVSGGNYSHKRNVSRASNSSQTS NFLAEQYERDRKAIINCCFSRPDHK TGEPPNNYITHVRIIEDSKFPSSRPPP DSKLENKKRLLLSAKPNNAKLIQI HKARENSDGSFQIGRTWQLTELVRV EKDLEISEGFILTMASKYYWETNSAK ERTVFIKSLITLYIQTFEGHVPVLN WDLSLFYLDERSYQRAVITNRPGSV SPIKSPTSNFTTNTTQSVGSPFSAP TERTRSETESVNPVSTPASVEYHAG MKSLNKAPYSSNSTLNEVNKRYELE QQQQQEEAELRRLEEQRQLQLQKE NEMKRLEEERIKQEERKRQMELE HQRQLEEEERKRQMELEAKQMEL KRQRQFEQQRLKKERELLEIQRKQ REQETAERLKKEEQEALAKKEEEK SKRNKVDNESYQEQINGKVDNLLD LNAVLAETETPTMONGTYVPERS TARAHQQLKPLNIAKVESLGGSDL NDSISLSDEIAGLNTSNLSGEDQDE KNDLSFEKGDEVRYSNFEGEAPH VYHEVSIIQEEAPAVSQKLILPEENN ESEALIESKEEIKTMENIDDEVILLEIL TDINWSIEDDADSMIERIDLRLAETE YLFNQNLQKIGPNIRPYEDKVND ECHRIIPTLSLFLMEMSNFSNDIENV ESQDNLQVESANKLLWNTLDEL LKTVSLDEISLNQLECEPIREKNLPW MENQLNLLLKAFQAIGSDGNEVEY NLREISGLKQRLQFYEKVTKIFLNRI VEEMQKFSNIRGQDISHDQMIRIL TLLIFSPILFCKEISQKSYQAIVEN WNVSIQPVYMELWTKKISQLQGIDT NDEKMNLSLSQLLNEWDTFRKER KTNDINPVFKNSFLLTECLQTMRO ECIVYQNFVEVFFHISSKHNFEYIK HFNDPDAPPILLDTVKVMQSDREAA VIETQLVSRIFQPIVTRLSSYFVELVK AEPTVAPALTFYLENEIKSLESSNHE FLLSAVTRMYTQIKQVWSDNVEEQV LHFERISNATTNGEILPGILDLPVGL KNSDLFQFAKRSMDIKDTDEGYES IELMNSSFRKLSIAATRSITHKEVNS SINPNLSDTAALNNDYMETISLLVNS NWLTEMLSMLNFNKDGIFDTSLOQ VKKVFDVEKESYASFLLRDTMPKLT AFVYGVSNIIENTNNVNMTPSRWA

