

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P43565	RIM15_YEAST	Saccharomyces cerevisiae	Serine/threonine-protein kinase RIM15	9.365641	NaN	S380;S476;T704;S709;S733;S736;S737;T747;S1044;S1048;S1064;T1075;S1421;S1531;S1538;S1542;S1565;S1764	33229814	MFNRSNTAGGSQAMKEGLGINKLS PISSNSNPSSLTSSNYEKYLQATEK NPCMILELELDGKVRYGSPQWNFTIT GVADDSGSSPTYIADLILGSDQDKGV FQKATDMLLMNDDTSTITFKIAA DYEGSAGCDESTITTLARGILIRD GHTQLPSHTMWIVKPRNDWSDFY ANEDAQDDMVIQLSDNCDDIDIQLP EEFAKTLGFGAKIFVQYLKRIRLEMII DEFNLPLPKMELCRVCENFVPVWW LETHSQSCVCEHRTESLIQLLHDNL LEQQAILANFTKDSEYKGSQIQVRSN NFLNQVLDLRELCQDAIDINPSEM VPDLYHSLSTFPQDNGNNNNNNNN NNNNNNNALLDQFPIQKDTVSLNS YFQFSPRTNHNINQVNTSWQSRFFLN DDQDPGLALLIHDTLDLARKKVDVAV LRLDNAMTYSLKIKNEVNNYVVQLI REQIEINKHAILTHPMNLRSSSIFHS PLPQIHSQQPEAENLIYSSSTPLQVQ HDQCASFEPKSHLEPIFPVSSIE ETPTANDIRHPSPLPRSCSNTVMKL PTPRKLDNGLFSDAYLNADIIPNP SIESTISIDRDNNTNSRGSSMKQYGI GEATDSRTSNSERPSSSSSRLGIRSR SITPROKIEYSHVDNDDRTNEMLSR DKDSLQPQPSVDTTITSSTQATTTGT KTNSNNSTNSVLPKLMTSISLTPRR GSPSFGNLASHSMQQTNSFKLIHDK SPISSPFTFSKDFLTPEQHPSNIART DSINNAML TSPNMPLSPLLATNQT VKSPTPSIKDYDILKPISKGAYGSVYL ARKKLTGDYFAIKVLRKSDMIAKNQ VTNVKSERAIMMVQSDKPYVARLFA SFQNKDNLFLVMEYLPGGDLATLIK MMGYLPDQWAKQYLTEIVVGVNDM HQNGIIHDLKPENLLIDNAGHVKL TDFGLSRAGLIRRHKFVPHKSSLSIS STLPIDNPANNFTMNNNNNSNHSQ STPDSFTSDHKQYNRSKKSSLGQQY EHSEYSSTNSHSMTPPTSTNTVVY PSYYRGKDRSHGSSNIDLPAASLRSE SQLSFLLDISRSSTPPLANPTNSNA NNIMRRKSLTENKSFSNDLLSSDAI AATNTNINSNNNISLSPAPSDALFY PDDSKQNKFFGTDPYLAPETIEGK GEDNKQCDWWSVGCIFPELLGYPP FHAETPDVAVFKKILSGVIQWPEFKNE EEREFLTPEAKDLIEKLLVVDPAKR

