

1 **Table S1. List of primers used in this study.**

Primer	Sequence (5' to 3')	Description
pUC19_IF_Fw	aggatccccgggtaccga	Construction for pUC18-ΔCAGL0F04279g-natNT2, and pUC18-ΔCAGL0F04279g-kanMX6
pUC19_IF_Rv	ctagagtcgacctgcaggca	
pUC19_dCgRIB1_IF1_Fw	gcaggctcgactctagctgtataactcactctatacgaacc	
pUC19_dCgRIB1_IF1_Rv	gtacccggggatccttttgactctaacacagataaattg	
dCgRIB1_IF2_vecFw	tacagtaaagatctttacacaga	
dCgRIB1_IF2_vecRv	ttgttgattgaatttgaatttga	
dCgRIB1_IF2_insFw	aattcaatacaacaacgtacgctgcaggctcgac	
dCgRIB1_IF2_insRv	aagatctttactgtaatcgatgaattcgagctcgttt	
pUC19_dCgYHB1_IF1_Fw	gcaggctcgactctagtagaatgggcttttagtactaac	Construction for pUC18-YHB1-natNT2
pUC19_dCgYHB1_IF1_Rv	gcaggctcgactctagtagaatgggcttttagtactaac	
dCgYHB1_IF2_vecFw	gaaagatggatcatgatattctga	
dCgYHB1_IF2_vecRv	cttttggtgtgtgtgtttcttgggt	
dCgYHB1_IF2_insFw	aacaaacaacaaaagcgtacgctgcaggctcgac	
dCgYHB1_IF2_insRv	catgatccatctttcatcgatgaattcgagctcgttt	
CgRIB1_gateway_Fw	ggggacaagtttgtaaaaaaagcaggcttaattgtcccttcagttgtgc	Construction for pET53-CAGL0F04279g
CgRIB1_gateway_NoStop_Rv	ggggaccactttgtacaagaaagctgggtgattgtttcttgccttgaggg	
pUC19_dCgACT_IF1_Fw	aattcgtaatcatggctcatagctgtttcctgt	Construction for pUC18-CAGL0F04279g- <i>ACT1</i>
pUC19_dCgACT_IF1_Rv	cgagctcggtacccggggatc	
dCgACT_IF2_Fw	cgggtaccgagctcgatggattctggtatgttcga	
dCgACT_IF2_Rv	ccatgattacgaattttagaaacacttgtggtgaac	

3 **Table S1. Continued.**

pET53-CgRIB1-Fw	aacattaacatctagatggcacatcaccaccacc	Construction for pCU-PDC1- <i>ScRIB1</i> , and pCU-PDC1-CAGL0F04279g
pET53-CgRIB1-Rv	gggttggtgtctcgattacttttcgaactgcgggtgg	
dCgYHB1_Up500_Fw	gtagaatgggcttttagtact	Gene deletion
dCgYHB1_Down500_Fw	cgtattgttgaatttatgttg	
dCgRIB1_Up500_Fw	ctgtataactcactctatacgaaccattacac	Gene deletion, PCR analysis
dCgRIB1_Down500_Fw	tttggactctaacacagataaattgag	
clonNAT_R	cgatgtcctcgacggtcag	PCR analysis
kanmX4R	gtattgatgttgacgagtcgg	
CgRIB1_ORF522_Rv	aagccttcagcttctcgtc	
CgRIB1_Fw_RT-PCR	ctgggcatcaagaacgtcag	qPCR
CgRIB1_Rv_RT-PCR	gtccagtggatcgggatcat	
CgACT1_Fw_RT-PCR	tccatcgctcggtagacca ag	
CgACT1_Rv_RT-PCR	aaacggccatggtgttatcg	

5 **Table S2. List of yeast strains used in this study.**

Strain	Genotype	Description
KUE100-1	<i>his3 ura3</i>	WT strain
<i>rib1</i> Δ	<i>his3 ura3 rib1::natNT2</i> pCU-PDC1 pRS313	<i>rib1</i> single disruptant
<i>rib1</i> ΔΔ	<i>his3 ura3 rib1::natNT2 rib1::kanMX6</i> pCU-PDC1 pRS313	<i>rib1</i> double disruptant
<i>yhb1</i> Δ	<i>his3 ura3 yhb1::natNT2</i> pCU-PDC1 pRS313	<i>yhb1</i> disruptant
RIB1OE	KUE100-1 pCU-PDC1-CAGL0F04279g pRS313	<i>CgRIB1</i> overexpression
<i>rib1</i> ΔΔ + <i>ScRIB1</i>	<i>rib1</i> ΔΔ pCU-PDC1- <i>ScRIB1</i> pRS313	Complemented strain
<i>rib1</i> ΔΔ + <i>CgRIB1</i>	<i>rib1</i> ΔΔ pCU-PDC1-CAGL0F04279g pRS313	Complemented strain

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7 **Table S3. List of plasmids used in this study.**

Plasmid	Description
pUC18-ΔCAGL0F04279g-natNT2	Template to amplify the DNA fragment for CAGL0F04279g deletion by natNT2
pUC18-ΔCAGL0F04279g-kanMX6	Template to amplify the DNA fragment for CAGL0F04279g deletion by kanMX6
pUC18- <i>YHB1</i> -natNT2	Template to amplify the DNA fragment for <i>YHB1</i> deletion by natNT2
pUC18-CAGL0F04279g- <i>ACT1</i>	Template to draw the standard curve for quantitative PCR analysis
pCU-PDC1	Empty vector for <i>C. glabrata</i> harboring <i>URA3</i>
pRS313	Empty vector for <i>C. glabrata</i> harboring <i>HIS3</i>
pCU-PDC1-CAGL0F04279g	To express CAGL0F04279g under the control of the <i>PDC1</i> promoter
pCU-PDC1- <i>ScRIB1</i>	To express <i>ScRIB1</i> under the control of the PDC1 promoter
pET53-CAGL0F04279g	To express the recombinant protein encoded by CAGL0F04279g in <i>E. coli</i>