

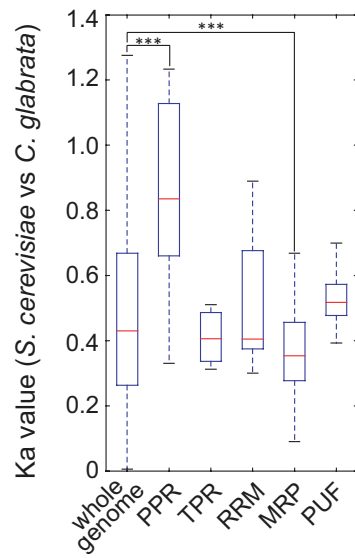
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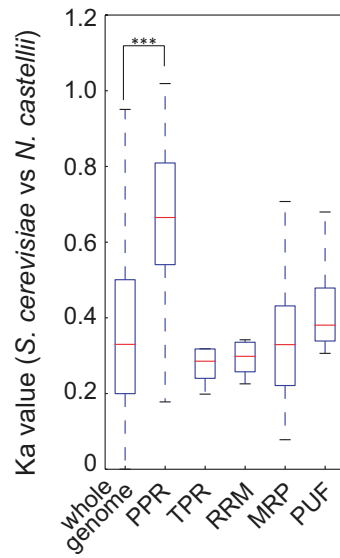
Supplementary Figures

Appendix Figure S1

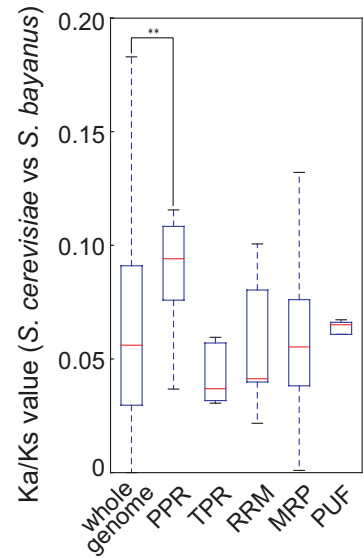
A



B



C



Supplementary Figure Legends

Appendix Figure S1. PPR genes show significantly increased evolutionary rates.

(A) Box plot of the evolutionary rates (Ka) of PPR, TPR, RRM, MRP, and PUF genes and the whole genome by comparing *S. cerevisiae* and *Candida glabrata* (a *Saccharomyces sensu lato* species). (B) Box plot of the evolutionary rates (Ka) by comparing *S. cerevisiae* and *N. castellii* (a *Saccharomyces sensu lato* species). (C) Box plot of the evolutionary rates (Ka/Ks) by comparing *S. cerevisiae* and *S. bayanus*. *p* values were calculated by two-sample Kolmogorov–Smirnov test. **, *p* value < 0.01. ***, *p* value < 0.001.

Supplementary Tables

Appendix Table S1. Summary of suppressor substitutions on Ccm1

Mutation	Position on the alignment	Position on PPR motif ^a	# of clones	PPR motif ^b
S→R (AGC→AGA)	242	20	1	a
F→L (TTC→TTA)	294	3	1	c
S→L (TCG→TTG)	305	15	1	c
E→K (GAA→AAA)	341	14	1	d
Q→H (CAA→CAC)	372	8	2	e
Q→K (CAA→AAA)	372	8	1	e
E→Q (GAA→CAA)	375	11	3	e
E→V (GAA→GUA)	375	11	1	e
E→K (GAA→AAA)	375	11	2	e
D→N (GAC→AAC)	400	1	1	f
D→Y (GAC→TAC)	400	1	1	f
A→G (GCT→GGT)	461	11	1	g
F→V (TTT→GTT)	476	26	1	g
F→I (TTT→ATT)	476	26	1	g
F→C (TTT→TGT)	476	26	1	g
K→R (AAG→AGG)	480	33	1	g

^a According to [1].

^b Referring to Fig 4D.

Appendix Table S2. Investigated MRP genes showed no evidence of incompatibility

Protein	Location	Function	Glucose medium	Glycerol medium
MRPL31	Chr 11	Large subunit of the mitochondrial ribosomal protein	Normal	Normal
MRPS17	Chr 13	Small subunit of the mitochondrial ribosomal protein	Normal	Normal
MRPL39	Chr 13	Large subunit of the mitochondrial ribosomal protein	Normal	Normal
MRPL24	Chr 12	Large subunit of the mitochondrial ribosomal protein	Normal	Normal
MRP2	Chr 16	Small subunit of the mitochondrial ribosomal protein	Normal	Normal
MRPS9	Chr 2	Large subunit of the mitochondrial ribosomal protein	Normal	Normal

Compatibility was tested between *S. cerevisiae* and three other species, *S. paradoxus*, *S. mikatae*, and *S. bayanus*.

Appendix Table S3. Strain list

Strain Name	Species	Genotype
JLD01	<i>S. cerevisiae</i>	<i>MATa/α ura3-1/ura3-1 ade2-1/ade2-1 his3-11,15/his3-11,15 leu2-3,112/leu2-3,112 trp1-1/trp1-1 can1-100/can1-100</i>
JYL1476	<i>S. cerevisiae</i>	<i>MATa ura3-1 ade2-1 his3Δ::3xHA leu2Δ::3xHA trp1-1 can1::MFap-HIS3 MFap-LEU2 ρ⁰</i>
JYL1477	<i>S. cerevisiae</i>	<i>MATa ura3-1 ade2-1 his3Δ::3xHA leu2Δ::3xHA trp1-1 can1::MFap-HIS3 MFap-LEU2 ρ⁰</i>
JYL1026	<i>S. bayanus</i>	<i>MATα ho::KAN ura3 his3Δ::3xHA leu2Δ::3xHA</i>
JYL1027	<i>S. bayanus</i>	<i>MATa ho::KAN ura3 his3Δ::3xHA leu2Δ::3xHA</i>
JYL1479	<i>S. bayanus</i>	<i>MATa ho::HPH ura3 his3Δ::3xHA leu2Δ::3xHA can1::MFap-HIS3 MFap-LEU2 ρ⁰</i>
JYL1478	<i>S. bayanus</i>	<i>MATα ho::HPH ura3 his3Δ::3xHA leu2Δ::3xHA can1::MFap-HIS3 MFap-LEU2 ρ⁰</i>
JYL1137	<i>S. paradoxus</i>	<i>MATa ho::HPH ura3 lys2 his3Δ::3xHA leu2Δ::3xHA</i>
JYL1251	<i>S. mikatae</i>	<i>MATa ho::HPH ura3 lys2</i>
JLD47	<i>S. kudriavzevii</i>	<i>MATa/α HO/HO ura3/ura3 lys2/lys2</i>
Sb-chr7 x Sc-ρ ⁰	<i>S. cerevisiae</i>	<i>MATa/α ura3-1/ura3-1 ade2-1/ade2-1 his3-11,15/his3Δ::3xHA leu2-3,112/ leu2Δ::3xHA trp1-1/trp1-1 can1-100/can1::MFap-HIS3 MFap-LEU2 Sc-chr7/Sb-chr7 Sc-mt</i>
Sb-chr7 x Sb-ρ ⁰	<i>S. cerevisiae x S. bayanus</i>	<i>MATa/α ura3-1/ura3 ade2-1/ade2-1 his3-11,15/his3Δ::3xHA leu2-3,112/ leu2Δ::3xHA trp1-1/TRP1 can1-100/can1::MFap-HIS3 MFap-LEU2 ho::HPH Sb-chr7/Sb-chr7 Sc-mt</i>
Sb-chr7-ρ ⁰ x Sb	<i>S. cerevisiae x S. bayanus</i>	<i>MATa/α ura3-1/ura3 his3-11,15/his3Δ::3xHA leu2-3,112/leu2Δ::3xHA ade2-1/ADE2 trp1-1/TRP1 can1-100/CAN1 ho::KAN Sb-chr7/Sb-chr7 Sb-mt</i>
Sc+Sb-CCM1 x Sc-ρ ⁰	<i>S. cerevisiae</i>	<i>MATa/α ura3-1/ura3-1 ade2-1/ade2-1 his3-11,15/his3Δ::3xHA leu2-3,112/ leu2Δ::3xHA trp1-1/trp1-1 can1-100/can1::MFap-HIS3 MFap-LEU2 ccm1Δ::SbCCM1-HIS3/CCM1 Sc-mt</i>
Sc+Sb-CCM1 x Sb-ρ ⁰	<i>S. cerevisiae x S. bayanus</i>	<i>MATa/α ura3-1/ura3 ade2-1/ade2-1 his3-11,15/his3Δ::3xHA leu2-3,112/ leu2Δ::3xHA trp1-1/TRP1 can1-100/can1::MFap-HIS3 MFap-LEU2 ho::HPH ccm1Δ::SbCCM1-HIS3/SbCCM1 Sc-mt</i>
Sc+Sb-CCM1-ρ ⁰ x Sb	<i>S. cerevisiae x S. bayanus</i>	<i>MATa/α ura3-1/ura3 his3-11,15/his3Δ::3xHA leu2-3,112/leu2Δ::3xHA ade2-1/ADE2 trp1-1/TRP1 can1-100/CAN1 ho::KAN ccm1Δ::SbCCM1-HIS3/SbCCM1 Sb-mt</i>
Sb+Sc-CCM1 x Sc-ρ ⁰	<i>S. cerevisiae x S. bayanus</i>	<i>MATa/α ura3-1/ura3 his3-11,15/his3Δ::3xHA leu2-3,112/leu2Δ::3xHA ade2-1/ADE2 trp1-1/TRP1 can1-100/CAN1 ho::KAN Sbccm1Δ::ScCCM1-HIS3/ScCCM1 Sb-mt</i>

Sb+Sc-CCMI x Sb-ρ ⁰	<i>S. bayanus</i>	<i>MATa/α ho::KAN/ ho::KAN ura3/ura3 his3Δ::3xHA/his3Δ::3xHA leu2Δ::3xHa/leu2Δ::3xHa CAN1/can1::MFap-HIS3 MFap-LEU2 Sbccm1Δ::ScCCMI-HIS3/SbCCMI Sb-mt</i>
Haploid <i>MATa</i> & <i>MATα</i> cells and isogenic diploid <i>MATa/α</i> cells are available for the following strains		
Sb-chr7	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 Sb-chr7</i>
Sb-chr7-ρ ⁰	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 Sb-chr7 ρ⁰</i>
Sc+Sc-CCMI	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::ScCCMI-HIS3</i>
Sc+Sc-CCMI	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::ScCCMI-URA3</i>
Sc+Sp-CCMI	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SpCCMI-URA3</i>
Sc+Sm-CCMI	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SmCCMI-URA3</i>
Sc+Sk-CCMI	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SkCCMI-URA3</i>
Sc+Sb-CCMI	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI-HIS3</i>
Sc+Sb-CCMI	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI-URA3</i>
Sc+Sb-CCMI-ρ ⁰	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI-HIS3 ρ⁰</i>
Sb+Sc-CCMI	<i>S. bayanus</i>	<i>ho::KAN ura3 his3Δ::3xHA leu2Δ::3xHA Sbccm1Δ::ScCCMI-HIS3</i>
Sb+Sb-CCMI	<i>S. bayanus</i>	<i>ho::KAN ura3 his3Δ::3xHA leu2Δ::3xHA Sbccm1Δ::SbCCMI-HIS3</i>
Sc+Sc-CCMI (Tom20-TAP)	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::ScCCMI-HIS3 tom20Δ::TOM20-TAP</i>
Sc+Sb-CCMI (Tom20-TAP)	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI-HIS3 tom20Δ::TOM20-TAP</i>
Sc+Sb-CCMI ^{S242R}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{S242R}-HIS3</i>
Sc+Sb-CCMI ^{F294L}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{F294L}-HIS3</i>
Sc+Sb-CCMI ^{S305L}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{S305L}-HIS3</i>
Sc+Sb-CCMI ^{E341K}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{E341K}-HIS3</i>
Sc+Sb-CCMI ^{Q372H}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{Q372H}-HIS3</i>
Sc+Sb-CCMI ^{Q372K}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{Q372K}-HIS3</i>
Sc+Sb-CCMI ^{E375Q}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{E375Q}-HIS3</i>
Sc+Sb-CCMI ^{E375K}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{E375K}-HIS3</i>

Sc+Sb-CCMI ^{E375V}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{E375V}-HIS3</i>
Sc+Sb-CCMI ^{D400N}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{D400N}-HIS3</i>
Sc+Sb-CCMI ^{D400Y}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{D400Y}-HIS3</i>
Sc+Sb-CCMI ^{A461G}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{A461G}-HIS3</i>
Sc+Sb-CCMI ^{F476V}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{F476V}-HIS3</i>
Sc+Sb-CCMI ^{F476I}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{F476I}-HIS3</i>
Sc+Sb-CCMI ^{F476C}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{F476C}-HIS3</i>
Sc+Sb-CCMI ^{F480R}	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ccm1Δ::SbCCMI^{F480R}-HIS3</i>
Sb+Sb-CCMI ^{E375Q}	<i>S. bayanus</i>	<i>ho::KAN ura3 his3Δ::3xHA leu2Δ::3xHA Sbccm1Δ::SbCCMI^{E375Q}-HSI3</i>
Sb+Sb-CCMI ^{D400N}	<i>S. bayanus</i>	<i>ho::KAN ura3 his3Δ::3xHA leu2Δ::3xHA Sbccm1Δ::SbCCMI^{D400N}-HSI3</i>
Sc+Sc-AEP1	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 aep1Δ pRS416-ScAEP1</i>
Sc+Sp-AEP1	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 aep1Δ pRS416-SpAEP1</i>
Sc+Sm-AEP1	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 aep1Δ pRS416-SmAEP1</i>
Sc+Sk-AEP1	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 aep1Δ pRS416-SkAEP1</i>
Sc+Sb-AEP1	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 aep1Δ pRS416-SbAEP1</i>
Sc+Sc-AEP3	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 AEP3Δ pRS416-ScAEP3</i>
Sc+Sp-AEP3	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 AEP3Δ pRS416-SpAEP3</i>
Sc+Sm-AEP3	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 AEP3Δ pRS416-SmAEP3</i>
Sc+Sk-AEP3	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 AEP3Δ pRS416-SkAEP3</i>
Sc+Sb-AEP3	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 AEP3Δ pRS416-SbAEP3</i>
Sc+Sc-ATP22	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ATP22Δ pRS416-ScATP22</i>
Sc+Sp-ATP22	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ATP22Δ pRS416-SpATP22</i>
Sc+Sm-ATP22	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ATP22Δ pRS416-SmAATP22</i>
Sc+Sk-ATP22	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ATP22Δ pRS416-SkATP22</i>
Sc+Sb-ATP22	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 ATP22Δ pRS416-SbATP22</i>

Sc+Sc-PET309	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 PET309Δ pRS416-ScPET309</i>
Sc+Sp-PET309	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 PET309Δ pRS416-SpPET309</i>
Sc+Sm-PET309	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 PET309Δ pRS416-SmPET309</i>
Sc+Sk-PET309	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 PET309Δ pRS416-SkPET309</i>
Sc+Sb-PET309	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 PET309Δ pRS416-SbPET309</i>
Sc+Sc-CBP1	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 CBP1Δ pRS416-ScCBP1</i>
Sc+Sm-CBP1	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 CBP1Δ pRS416-SmCBP1</i>
Sc+Sc-MRX1	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 MRX1Δ pRS416-ScMRX1</i>
Sc+Sm-MRX1	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 MRX1Δ pRS416-SmMRX1</i>
Sc+Sc-PET111	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 PET111Δ pRS416-ScPET111</i>
Sc+Sm-PET111	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 PET111Δ pRS416-SmPET111</i>
Sc+Sc-RMD9	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 RMD9Δ pRS416-ScRMD9</i>
Sc+Sp-RMD9	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 RMD9Δ pRS416-SpRMD9</i>
Sc+Sm-RMD9	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 RMD9Δ pRS416-SmRMD9</i>
Sc+Sk-RMD9	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 RMD9Δ pRS416-SkRMD9</i>
Sc+Sb-RMD9	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 RMD9Δ pRS416-SbRMD9</i>
Sc+Sc-RPM2	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 RPM2Δ pRS416-ScRPM2</i>
Sc+Sp-RPM2	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 RPM2Δ pRS416-SpRPM2</i>
Sc+Sm-RPM2	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 RPM2Δ pRS416-SmRPM2</i>
Sc+Sk-RPM2	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 RPM2Δ pRS416-SkRPM2</i>
Sc+Sb-RPM2	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 RPM2Δ pRS416-SbRPM2</i>
Sc+Sc-RPO41	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 RPO41Δ pRS416-ScRPO41</i>
Sc+Sp-RPO41	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 RPO41Δ pRS416-SpRPO41</i>
Sc+Sm-RPO41	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 RPO41Δ pRS416-SmRPO41</i>
Sc+Sk-RPO41	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100 RPO41Δ pRS416-SkRPO41</i>

Sc+Sb-RPO41	<i>S. cerevisiae</i>	<i>ura3-1 ade2-1 his3-11,15 leu2-3,112 trp1-1 can1-100</i> <i>RPO41Δ pRS416-SbRPO41</i>
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Supplementary References

1. Lipinski KA, Puchta O, Surendranath V, Kudla M, Golik P (2011) Revisiting the yeast PPR proteins--application of an Iterative Hidden Markov Model algorithm reveals new members of the rapidly evolving family. *Mol Biol Evol* **28**: 2935-48