Table S1. Yeast strains used in this study.

Strain	Genotype	Reference
W1588-4C	*MAT***a** *ade2-1 can1-100 his3-11,15 leu2-3,112 trp1-1 ura3-1*	Thomas and Rothstein 1989, Zhao *et al* 1998
YTT3320	W1588-4C; *isw2Δ::NatMX*	Au *et al* 2011
YTT6809	W1588-4C; *isw2Δ::NatMX*	this study
YTT3333	W1588-4C; *nhp10Δ::Hyg*	Au *et al* 2011
YTT2060	W1588-4C; *nhp10Δ::Hyg*	Vincent *et al* 2008
YTT3337	W1588-4C; isw2Δ::NatMX nhp10Δ::HYG	Au *et al* 2011
YTT2109	W1588-4C; *isw2Δ::NatMX nhp10Δ::HYG*	Vincent *et al* 2008
YTT1996	W1588-4C; *ISW2-K215R-3FLAG-KanMX*	Gelbart *et al* 2005
YTT1997	W1588-4C; *ISW2-K215R-3FLAG-KanMX*	Gelbart *et al* 2005
YTT3426	W1588-4C; *NHP10-3FLAG-KanMX*	Vincent *et al* 2008
YTT3427	W1588-4C; *NHP10-3FLAG-KanMX*	Vincent *et al* 2008
YTT6639	W1588-4C; *rpa49Δ::KanMX*	this study
YTT6673	W1588-4C; *isw2Δ::NatMX nhp10Δ::Hyg RPA190-2L-3FLAG::KanMX*	this study
YTT6679	W1588-4C; *RPA190-2L-3FLAG::KanMX*	this study
YTT6686	W1588-4C; *RPO31-2L-3FLAG::KanMX*	this study
YTT6693	W1588-4C; *isw2Δ::NatMX nhp10Δ::Hyg RPO31-2L-3FLAG::KanMX*	this study
YTT6915	W1588-4C; *Pol2-2L-3FLAG::KanMX*	this study
YTT6916	W1588-4C; *Pol2-2L-3FLAG::KanMX*	this study
YTT6917	W1588-4C; *isw2Δ::NatMX Pol2-2L-3FLAG::KanMX*	this study
YTT6918	W1588-4C; *isw2Δ::NatMX Pol2-2L-3FLAG::KanMX*	this study
YTT6919	W1588-4C; *nhp10Δ::Hyg Pol2-2L-3FLAG::KanMX*	this study
YTT6920	W1588-4C; *nhp10Δ::Hyg Pol2-2L-3FLAG::KanMX*	this study
YTT6921	W1588-4C; *isw2Δ::NatMX nhp10Δ::Hyg Pol2-2L-3FLAG::KanMX*	this study
YTT6922	W1588-4C; *isw2Δ::NatMX nhp10Δ::Hyg Pol2-2L-3FLAG::KanMX*	this study
YTT7009	W1588-4C; *Fob1-2L-3FLAG::KanMX*	this study
YTT7010	W1588-4C; *Fob1-2L-3FLAG::KanMX*	this study
YTT7011	W1588-4C; *isw2Δ::NatMX Fob1-2L-3FLAG::KanMX*	this study
YTT7012	W1588-4C; *isw2Δ::NatMX Fob1-2L-3FLAG::KanMX*	this study
YTT7013	W1588-4C; *nhp10Δ::Hyg Fob1-2L-3FLAG::KanMX*	this study
YTT7014	W1588-4C; *nhp10Δ::Hyg Fob1-2L-3FLAG::KanMX*	this study
YTT7015	W1588-4C; *isw2Δ::NatMX nhp10Δ::Hyg Fob1-2L-3FLAG::KanMX*	this study
YTT7016	W1588-4C; *isw2Δ::NatMX nhp10Δ::Hyg Fob1-2L-3FLAG::KanMX*	this study
YSI101	*MAT***a** *ade2-1 can1-100 his3-11,15 leu2-3,112 trp1-1 ura3-1 fob1::LEU2*	Ide *et al* 2010
YSI102	YSI101; 20 copies rDNA	Ide *et al* 2010
YSI103	YSI101; 40 copies rDNA	Ide *et al* 2010
YSI104	YSI101; 80 copies rDNA	Ide *et al* 2010
YTT6294	YSI102; isw2Δ::NatMX	this study
YTT6865	YSI102; nhp10Δ::HYG	this study
YTT6311	YSI102; isw2Δ::NatMX nhp10Δ::Hyg	this study
YTT6312	YSI102; isw2Δ::NatMX nhp10Δ::Hyg	this study

YTT3383	*MAT***a** ade2-1::pRS402 *can1-100 his3-11,15 leu2-3,112 trp1-1* ura3-1::pRS406	
YTT3385	*MAT***a** ade2-1::pRS402 *can1-100 his3-11,15 leu2-3,112 trp1-1* ura3-1::pRS406 isw2Δ::NatMX	
YTT3387	*MAT***a** ade2-1::pRS402 *can1-100 his3-11,15 leu2-3,112 trp1-1* ura3-1::pRS406 nhp10Δ::HYG	
YTT3388	*MAT***a** ade2-1::pRS402 *can1-100 his3-11,15 leu2-3,112 trp1-1* ura3-1::pRS406 isw2Δ::NatMX nhp10Δ::HYG	

Table S2. Primers used in this study.

Name	Comment	Sequence
ETS1-1	5' ETS1 qPCR	TGGGTTGATGCGTATTGAGA
ETS1-2	3' ETS1 qPCR	TCGCTGATTTGAGAGGAGGT
ALG9-1	5' ALG9 qPCR	CACGGATAGTGGCTTTGGTGAACAATTAC (TESTE *et al.* 2009)
ALG9-2	3' ALG9 qPCR	TATGATTATCTGGCAGCAGGAAAGAACTTGGG (TESTE *et al.* 2009)
ITS1-6	5' ITS1 qPCR	TGTTTTGGCAAGAGCATGAG
ITS1-7	3' ITS1 qPCR	TCGAATGCCCAAAGAAAAAG
RFB-1	5' RFB qPCR, probe	gcggggtctagaCCACTGTTCACTGTTCACTGTTCA
RFB-2	3' RFB qPCR, probe	cccggcgctagcAGAGAAGGGCTTTCACAAAGCT
rDNA_ETS1-1	5' ETS1 probe	CCATTCCGTGAAACACC
rDNA_ETS1-2	3' ETS1 probe	AAGAAAGAAACCGAAATCTC
AG_Fob1_1	5' Fob1 Gibson cloning (insert)	ctcactatagggcgaattgggtaccgggccTTAATAATGTACTTTGCAGATGTTTGTTCC
AG_Fob1_3	3' Fob1 Gibson cloning (insert)	cgcggtggcggccgctctagaactagtggaCTAATGATAATGGCTTTCTATTTGTTTTGC
AG_Fob1_2	5' Fob1 Gibson cloning (vector)	GGAACAAACATCTGCAAAGTACATTATTAAggcccggtacccaattcgccctatagtgag
AG_Fob1_4	3' Fob1 Gibson cloning (vector)	GCAAAACAAATAGAAAGCCATTATCATTAGtccactagttctagagcggccgccaccgcg

Teste, M. A., M. Duquenne, J. M. Francois and J. L. Parrou, 2009 Validation of reference genes for quantitative expression analysis by real-time RT-PCR in Saccharomyces cerevisiae. BMC Mol Biol 10**:** 99.

Table S3. Plasmids used in this study.

Name	Description
pRS426	URA3, 2μ
pRS426-Fob1	pRS426 with *FOB1* promoter and coding region