

Supplemental Table S3: Reference number and description of yeast strains used in this study.

Strain #	Derived from	Description / Phenotype	Mating type	Source or reference
YHL9632		Tf1-neo Integration (in the intergenic region <i>ssa1-upf3</i>), <i>ura4-D18 leu1-32 ade6-M210</i>	h+	Feng et al, <i>Nucleic Acids Res</i> 41, 775-789 (2013)
YHL9631		Tf1-neo Integration (in the intergenic region SPAC19A8.06- <i>sst4</i>), <i>ura4-D18 leu1-32 ade6-M210</i>	h+	Feng et al, <i>Nucleic Acids Res</i> 41, 775-789 (2013)
YHL9629		Tf1-neo Integration (in the intergenic region SPAC24B11.08c- <i>mpc2</i>), <i>ura4-D18 leu1-32 ade6-M210</i>	h+	Feng et al, <i>Nucleic Acids Res</i> 41, 775-789 (2013)
YHL9638		Tf1-neo Integration (in the intergenic region <i>sim4-rpl1001</i>), <i>ura4-D18 leu1-32 ade6-M210</i>	h+	Feng et al, <i>Nucleic Acids Res</i> 41, 775-789 (2013)
YHL10157	YHL9632	Tf1-neoAI Integration #1 (in the intergenic region <i>ssa1-upf3</i>), <i>ura4-D18 leu1-32 ade6-M210</i>	h+	This study
YHL10277	YHL9631	Tf1-neoAI Integration #2 (in the intergenic region SPAC19A8.06- <i>sst4</i>), <i>ura4-D18 leu1-32 ade6-M210</i>	h+	This study
YHL11123	YHL9629	Tf1-neoAI Integration #3 (in the intergenic region SPAC24B11.08c- <i>mpc2</i>), <i>ura4-D18 leu1-32 ade6-M210</i>	h+	This study
YHL11124	YHL9638	Tf1-neoAI Integration #4 (in the intergenic region <i>sim4-rpl1001</i>), <i>ura4-D18 leu1-32 ade6-M210</i>	h+	This study
YHL5661		WT diploid lacking Tf1, <i>ura4-D18/ura4-D18 leu1-32/leu1-32 ade6-M210/ade6-M216</i>	stable diploid h+	Chatterjee et al, <i>Nucleic Acids Res</i> 42, 8449-8460 (2014)
YHL1605		WT haploid lacking Tf1, <i>ura4-D18/ura4-D18 leu1-32/leu1-32 ade6-M210/ade6-M216</i>	h+	This study
YHL11396	YHL1605	Tf1-neo Integration #7 at Chr. II 3858489 (+) (<i>sat1-ssn6</i> intergenic region)	h+	This study
YHL11450		Tf1-neo Integration #5 at Chr. II 3607268 (-) (<i>mok13-zrt1</i> intergenic region)	h+	This study
YHL11550		Tf1-neo Integration #6 at Chr. I 678437 (-) (<i>tsc1-rgs1</i> intergenic region)	h+	This study
V2-33-H11		WT H11 <i>ade6-210</i> of same background than Bioneer deletions	h+	Bioneer
V2-33-H12		WT H12 <i>ade6-216</i> of same background than Bioneer deletions	h+	Bioneer
V2-23-H05		deletion of the GTPase <i>Ryh1</i>	h+	Bioneer
V2-12-G08		deletion of the phosphatidylinositol kinase <i>Tor1</i>	h+	Bioneer
V2-14-H10		deletion of the stress activated MAP kinase interacting protein <i>Sin1</i>	h+	Bioneer
V2-22-H06		deletion of the hamartin <i>Tsc1</i>	h+	Bioneer
V2-12-G04		deletion of the tuberlin <i>Tsc2</i>	h+	Bioneer
V2-21-H09		deletion of the TORC1 subunit <i>Tco89</i>	h+	Bioneer
YBN174		deletion of <i>gtr1</i>	h+	Bioneer
V2-05-B11		deletion of <i>npr2</i>	h+	Bioneer
V2-07-G09		deletion of <i>sat1</i>		