

**Supplemental Table I: Yeast strains used in this study.**

Strain	Genotype
MYY61	<i>MATa, ade2-1, leu2-3, can1-100, trp1-1, ura3-52, his3-11,15, Brr2-13myc::KANMX6, hmt1::HIS3</i>
MYY92	<i>MATa, ade2-1, leu2-3, can1-100, trp1-1, ura3-52, his3-11,15, Brr2-13myc::KANMX6,</i>
MYY72	<i>MATa, ade2-1, leu2-3, can1-100, trp1-1, ura3-52, his3-11,15, Prp11-9myc::TRP1, hmt1::HIS3</i>
MYY79	<i>MATa, ade2-1, leu2-3, can1-100, trp1-1, ura3-52, his3-11,15, Mud2-13myc::KANMX6, hmt1::HIS3</i>
MYY90	<i>MATa, ade2-1, leu2-3, can1-100, trp1-1, ura3-52, his3-11,15, Prp2-9myc::TRP1</i>
MYY104	<i>MATa, ade2-1, leu2-3, can1-100, trp1-1, ura3-52, his3-11,15, Prp2-9myc::TRP1, hmt1::HIS3</i>
MYY379	<i>MATa, ade2-1, leu2-3, can1-100, trp1-1, ura3-52, his3-11,15, Prp5-9myc::TRP1</i>
MYY386	<i>MATa, ade2-1, leu2-3, can1-100, trp1-1, ura3-52, his3-11,15, Prp5-9myc::TRP1, hmt1::HIS3</i>
MYY389	<i>MATa, ade2-1, leu2-3, can1-100, trp1-1, ura3-52, his3-11,15, Prp40-9myc::TRP1, hmt1::KANMX6</i>
MYY477	<i>MATa, his3<math>\Delta</math>1, leu2<math>\Delta</math>0, met15<math>\Delta</math>0, ura3<math>\Delta</math>0, SNP1-TAP::HIS3</i>
MYY478	<i>MATa, his3<math>\Delta</math>1, leu2<math>\Delta</math>0, met15<math>\Delta</math>0, ura3<math>\Delta</math>0, SNP1-GFP::HIS3</i>
MYY496	<i>MATa, ade2-1, leu2-3, can1-100, trp1-1, ura3-52, his3-11,15, Prp11-9myc::TRP1, hmt1-G68R::HIS3::LEU2</i>
MYY498	<i>MATa, ade2-1, leu2-3, can1-100, trp1-1, ura3-52, his3-11,15, Mud2-13myc::KANMX6, hmt1-G68R::HIS3::LEU2</i>
MYY512	<i>MATa, his3<math>\Delta</math>1, leu2<math>\Delta</math>0, met15<math>\Delta</math>0, ura3<math>\Delta</math>0, SNP1-GFP::HIS3, hmt1::KANMX6</i>
MYY514	<i>MATa, ade2-1, leu2-3, can1-100, trp1-1, ura3-52, his3-11,15, SNP1-9myc::TRP1</i>
MYY554	<i>MATa, his3<math>\Delta</math>1, leu2<math>\Delta</math>0, met15<math>\Delta</math>0, ura3<math>\Delta</math>0, SNP1-TAP::HIS3, hmt1::KANMX6</i>
MYY580	<i>MATa, his3<math>\Delta</math>1, leu2<math>\Delta</math>0, met15<math>\Delta</math>0, ura3<math>\Delta</math>0, SNP1-TAP::HIS3, hmt1- G68R::KANMX6::LEU2</i>
MYY582	<i>MATa,ade2-1, leu2-3, can1-100, trp1-1, ura3-52, his3-11,15, SNP1-9myc::TRP1, hmt1::KANMX6</i>
PSY3347	<i>MATa, ade2-1, leu2-3, can1-100, trp1-1, ura3-52, his3-11,15, Prp40-9myc::TRP1</i>

**Supplemental Table 2: Oligonucleotides used in this study.**

<u>Name</u>	<u>Description</u>	<u>Sequence</u>
MCY978	<i>RPS13</i> ChIP FWD	TGCTCCAGCTTGGTTCAAGTTGTC
MCY979	<i>RPS13</i> ChIP REV	TTGGGTAACACCGTGAGCATCTCT
MCY994	<i>ARP9</i> ChIP FWD	GGCTCCATTTAGGCAGGACAGTAT
MCY995	<i>ARP9</i> ChIP REV	AACCATCTTGTCTCGTGGTCCGAT
MCY998	<i>HOP2</i> ChIP FWD	AGAAATCAAACGATAGAGCCATTCAGG
MCY999	<i>HOP2</i> ChIP REV	GTTCTTTAGTTACAGAGTCAAGAGCGTC
MCY1016	<i>REC107</i> ChIP FWD	ACTGGAATGGGCAGGTAAACTGGA
MCY1017	<i>REC107</i> ChIP REV	ACTTGAGGGTATGCACTGTGTTGG