

Table S2. Yeast strains used in this study

Strain name	Genotype	Origin
BY4741	<i>MATa ura3 his3 leu2 met15 TRP1</i>	Euroscarf
Arx1-TAP	<i>MATα his3 leu2 ura3 Arx1-TAP::TRP1</i>	[1]
Enp1-TAP	<i>MATa ura3 his3 leu2 TRP1 Enp1-TAP::HIS3MX</i>	Open biosystems
Enp1-TAP <i>mex67</i> shuffle	<i>MATα ura3 his3 leu2 ade2 TRP1 mex67::HIS3MX ENP1-TAP::HIS3MX <pRS316-MEX67></i>	this study
Enp1-TAP <i>mtr2</i> shuffle	<i>MATα ura3 his3 leu2 ade2 TRP1 mtr2::HIS3MX ENP1-TAP::HIS3MX <pRS316-MTR2></i>	this study
Enp1-TAP <i>slx9</i> Δ	<i>MATa ura3 his3 leu2 TRP1 Enp1-TAP::HIS3MX slx9::NATNT2</i>	this study
Hrr25-TAP	<i>MATa ura3 his3 leu2 TRP1 Hrr25-TAP::HIS3MX</i>	Open biosystems
Kre35-TAP	<i>MATα ura3 his3 leu2 trp1 met15 Kre35-TAP::TRP1</i>	[1]
<i>mex67</i> shuffle <i>slx9</i> Δ	<i>MATa ura3 his3 leu2 trp1 ade2 mex67::HIS3MX slx9::KANMX <pRS316-MEX67></i>	this study
<i>mex67</i> shuffle <i>yrb2</i> Δ	<i>MATa ura3 his3 leu2 trp1 ade2 mex67::HIS3MX <pRS316-MEX67> yrb2::KAN</i>	this study
<i>mex67</i> shuffle <i>rps15-1</i>	<i>MATa ura3 his3 leu2 trp1 ade2 mex67::HIS3MX <pRS316-MEX67> rps15::KAN <pRS316-MEX67> <Rps15-LEU></i>	this study
<i>mex67</i> shuffle <i>bud23</i> Δ	<i>MATa ura3 his3 leu2 ade2 mex67::HIS3MX bud23::KANMX <pRS316-MEX67></i>	this study
<i>mex67</i> shuffle <i>rps6a</i> Δ	<i>MATa ura3 his3 leu2 ade2 mex67::HIS3MX rps6a::KANMX <pRS316-MEX67></i>	this study
<i>mex67</i> shuffle <i>ltv1</i> Δ	<i>MATa ura3 his3 leu2 trp1 ade2 mex67::HIS3MX ltv1::KAN <pRS316-MEX67></i>	this study
<i>mtr2</i> shuffle <i>slx9</i> Δ	<i>MATa ura3 his3 leu2 trp1 ade2 mtr2::HIS3MX slx9::KANMX <pRS316-MTR2></i>	this study
<i>mtr2</i> shuffle <i>yrb2</i> Δ	<i>MATa ura3 his3 leu2 trp1 ade2 mtr2::HIS3MX yrb2::KAN <pRS316-MTR2></i>	this study
<i>mtr2</i> shuffle <i>rps15-1</i>	<i>MATa ura3 his3 leu2 trp1 ade2 mtr2::HIS3MX rps15::KAN <pRS316-MTR2> <Rps15-LEU></i>	this study
<i>mtr2</i> shuffle <i>bud23</i> Δ	<i>MATa ura3 his3 leu2 ade2 mtr2::HIS3MX bud23::KANMX <pRS316-MTR2></i>	this study

<i>mtr2</i> shuffle <i>rps6a</i> Δ	<i>MAT</i> α <i>ura3 his3 leu2 ade2 mtr2::HIS3MX rps6a::KANMX</i> <pRS316-MTR2>	this study
<i>mtr2</i> shuffle <i>ltv1</i> Δ	<i>MAT</i> α <i>ura3 his3 leu2 trp1 ade2 mtr2::HIS3MX ltv1::KAN</i> <pRS316-MTR2>	this study
<i>nmd3</i> shuffle <i>slx9</i> Δ	<i>MAT</i> α <i>his3 leu2 ura3 trp1 slx9::HIS3MX nmd3::KANMX4</i> <pRS316-NMD3>	this study
Noc4-TAP	<i>MAT</i> α <i>ura3 his3 leu2 TRP1 Noc4-TAP::HIS3MX</i>	Open biosystems
Rio2-TAP	<i>MAT</i> α <i>ura3 his3 leu2 TRP1 Rio2-TAP::HIS3MX</i>	Open biosystems
Rio2-TAP <i>mex67</i> shuffle	<i>MAT</i> α <i>ura3 his3 leu2 trp1 ade2 mex67::HIS3MX Rio2-TAP::KANMX</i> <pRS316-MEX67>	this study
Rio2-TAP <i>mtr2</i> shuffle	<i>MAT</i> α <i>ura3 his3 leu2 trp1 ade2 mtr2::HIS3MX Rio2-TAP::KANMX</i> <pRS316-MTR2>	this study
Rio2-TAP <i>yrb2</i> Δ	<i>MAT</i> α ; <i>his3D1; leu2D0; met15D0; yrb2::KANMX RIO2-TAP::URA</i>	this study
Rix1-TAP	<i>MAT</i> α <i>ura3 his3 leu2 trp1 met15 Rix1-TAP::TRP1</i>	[1]
<i>GAL-rps15-1</i>	<i>MAT</i> α <i>his3 leu2 ura3 rps15::KANMX</i> <pFL36-pGal-rps15-1>	[2]
<i>Slx9</i> -GFP <i>Gar1</i> -mCherry	<i>MAT</i> α / <i>MAT</i> α <i>ura3 his3 leu2 met15 TRP1 slx9GFP::HIS3MX Gar1mCherry::KANMX</i>	this study
<i>slx9</i> Δ	<i>MAT</i> α <i>ura3 his3 leu2 TRP1 slx9::KANMX</i>	this study
<i>slx9</i> Δ	<i>MAT</i> α <i>ura3 his3 leu2 TRP1 slx9::HIS3MX</i>	this study
<i>slx9</i> Δ <i>arx1</i> Δ	<i>MAT</i> α <i>ura3 his3 leu2 TRP1 slx9::NATNT2 arx1::KANMX</i>	this study
<i>slx9</i> Δ <i>ecm1</i> Δ	<i>MAT</i> α <i>ura3 his3 leu2 TRP1 slx9::NATNT2 ecm1::KANMX</i>	this study
<i>slx9</i> Δ <i>sub2</i> shuffle	<i>MAT</i> α <i>ura3 his3 leu2 TRP1 slx9::KANMX sub2::KAN</i> <pRS316-SUB2>	this study
<i>slx9</i> Δ <i>yra1</i> shuffle	<i>MAT</i> α <i>ura3 his3 leu2 TRP1 slx9::KANMX yra1::HIS3</i> <pRS316-YRA1>	this study
Ssf1-TAP	<i>MAT</i> α <i>his3 leu2 ura3 Ssf1-TAP ::TRP1</i>	[1]
<i>xpo1-1</i>	<i>MAT</i> α <i>ura3 his3 leu2 ade2 trp1 can1 xpo1::LEU2</i> <pKW457-xpo1-1-HIS3>	[3]
<i>yvh1</i> Δ	<i>MAT</i> α <i>ura3 his3 leu2 TRP1 yvh1::KANMX</i>	[4]

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