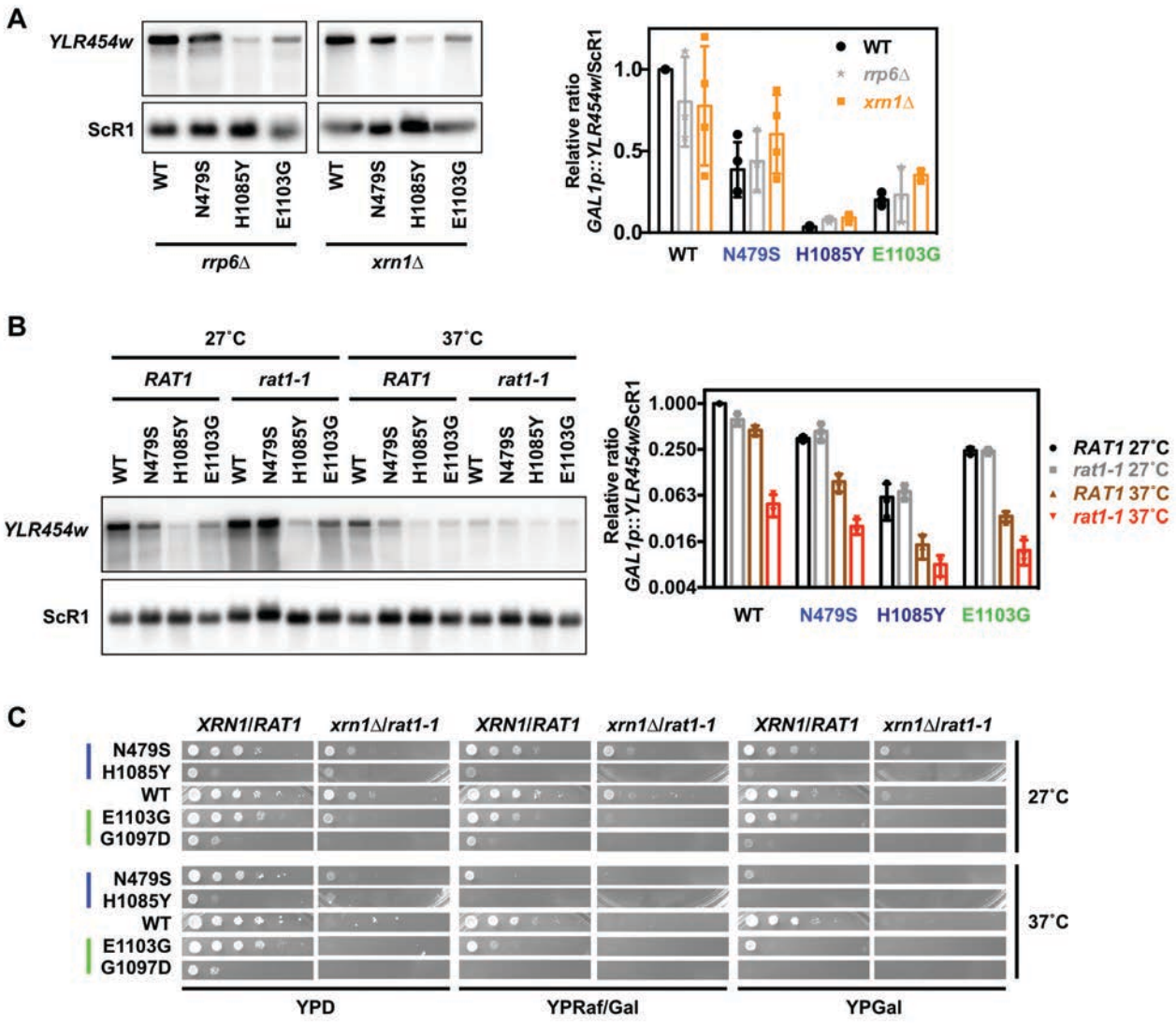
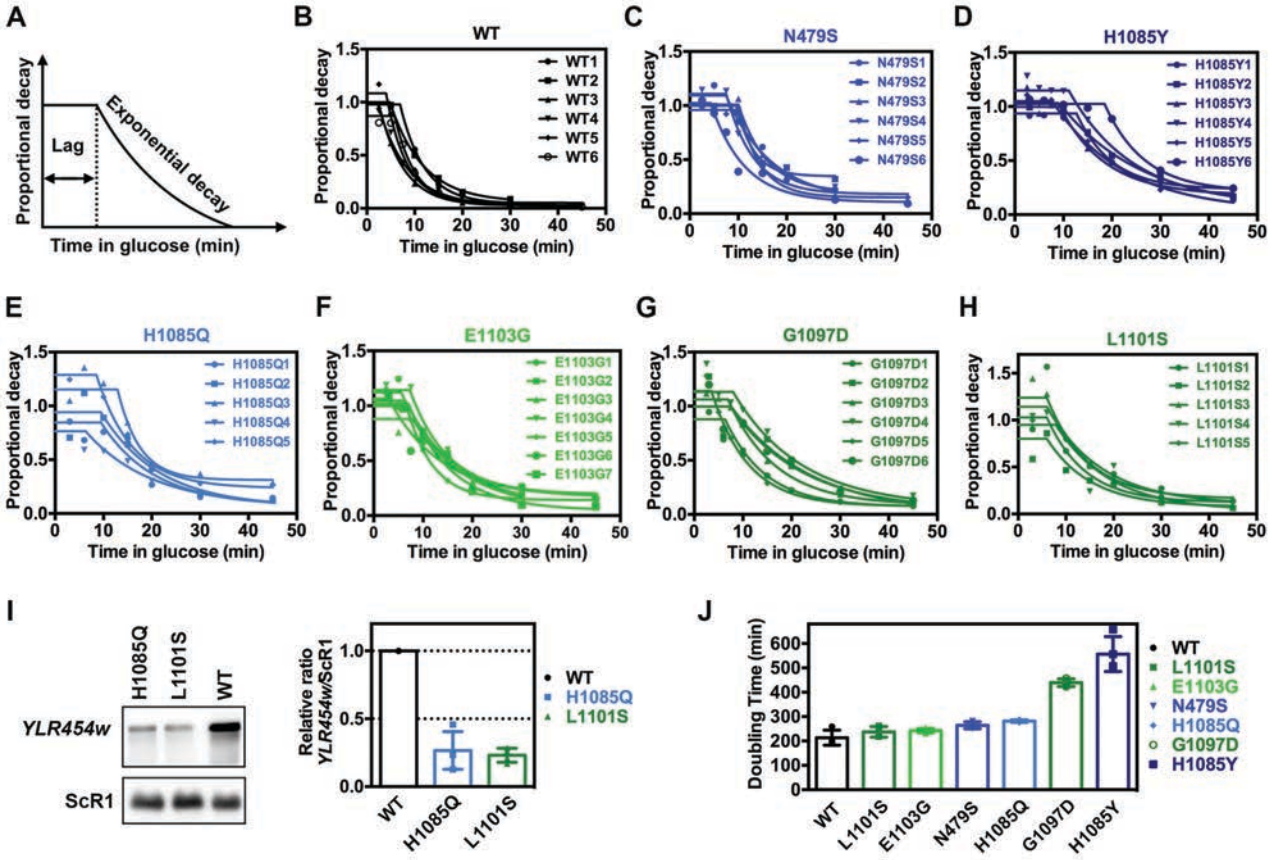
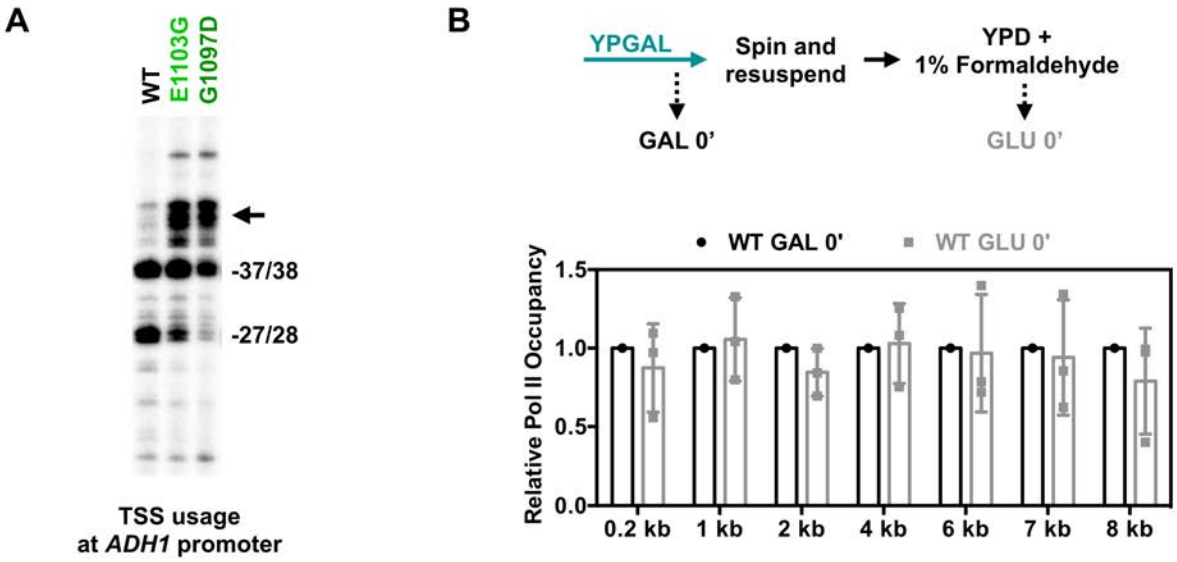


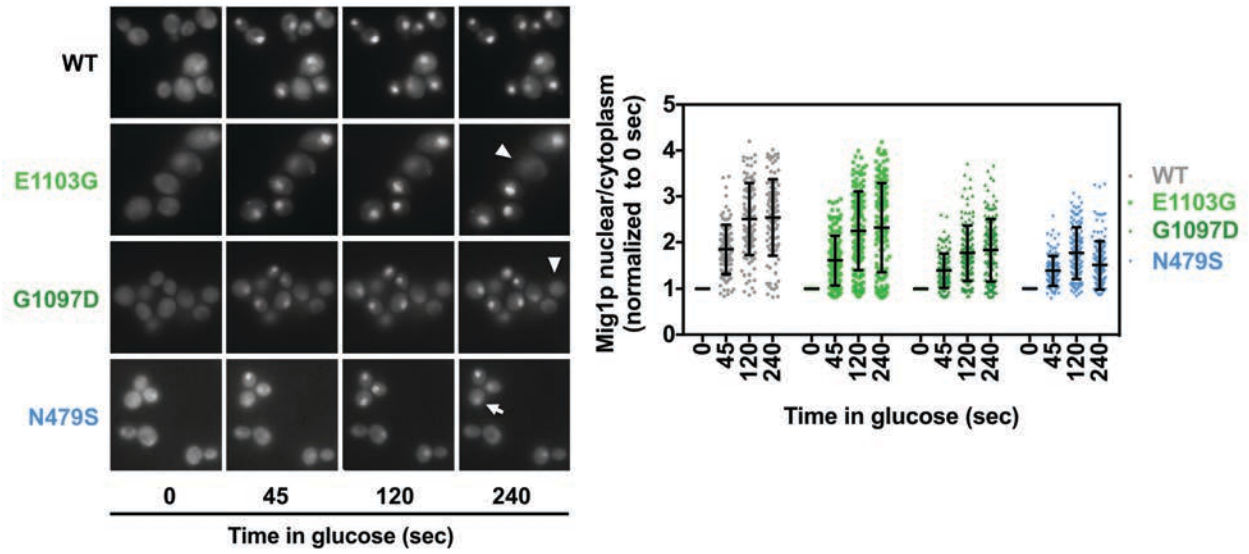
Supplementary Figure 1. Modulation of reporter gene expression defect in pre-mRNA processing factor mutants. (A) *GAL1p*::*YLR454w* reporter expression level in Pol II catalytic mutants in WT, *xrn1*Δ, or *rrp6*Δ backgrounds determined by Northern blotting. WT Pol II sample (*XRN1/RRP6*) was run in parallel with *xrn1*Δ or *rrp6*Δ mutant samples on each blot for normalization purposes (**Figure 2A**) (B) *GAL1p*::*YLR454w* reporter expression level in Pol II catalytic mutants in WT and in *rat1-1* background at permissive (27˚C) or restrictive (37˚C) temperatures. Overnight grown cells were inoculated in fresh YPGal media to amplify to mid-log growth at 27˚C, then shifted to 37˚C to inactivate Rat1p. RNAs were isolated from half of each culture prior to temperature shift (27˚C samples). Remaining cultures were washed and resuspended in pre-warmed YPGal media to grow for another 2 hrs at 37˚C prior to RNA isolation. Relative *GAL1p*::*YLR454w* expression levels were normalized to WT (*RAT1*) 27˚C value. Data shown are average of three biological repeats with error bars representing the standard deviation (SD) of the mean. Note: values were normalized to WT (*RAT1*) at 27˚C, and are presented on logarithmic scale. (C) 10-fold serial dilutions of saturated cultures of Pol II catalytic mutants alone or in combination with *xrn1*Δ/*rat1-1* mutant were plated for growth at 27˚C or 37˚C on different media as indicated. Blue and green bars indicate LOF and GOF mutants, respectively.



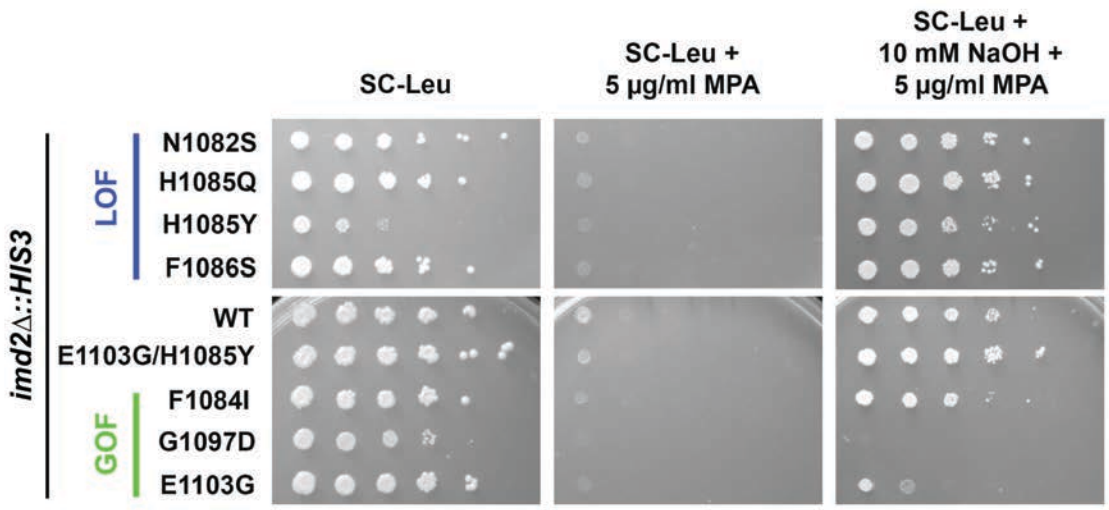
Supplementary Figure 2. Pol II catalytic mutants alter *GAL1*p::*YLR454w* mRNA decay and growth in YPGal media (A) Schematic of the mRNA decay curve fitting showing 'lag' and 'exponential decay' periods after transcriptional shut-off. (B-H) Individual decay curve for *GAL1*p::*YLR454w* decay in Pol II catalytic mutants plotted using non-linear regression using GraphPad prism. (I) *GAL1*p::*YLR454w* expression level in H1085Q and L1101S determined by Northern blotting. Values normalized to WT *GAL1*p::*YLR454w* expression level. Data shown are average of three biological repeats with error bars representing the standard deviation (SD) of the mean. (J) Doubling time of Pol II mutants in YPGal media determined using Tecan plate reader. Error bars represent average +/- SD of four replicate cultures.



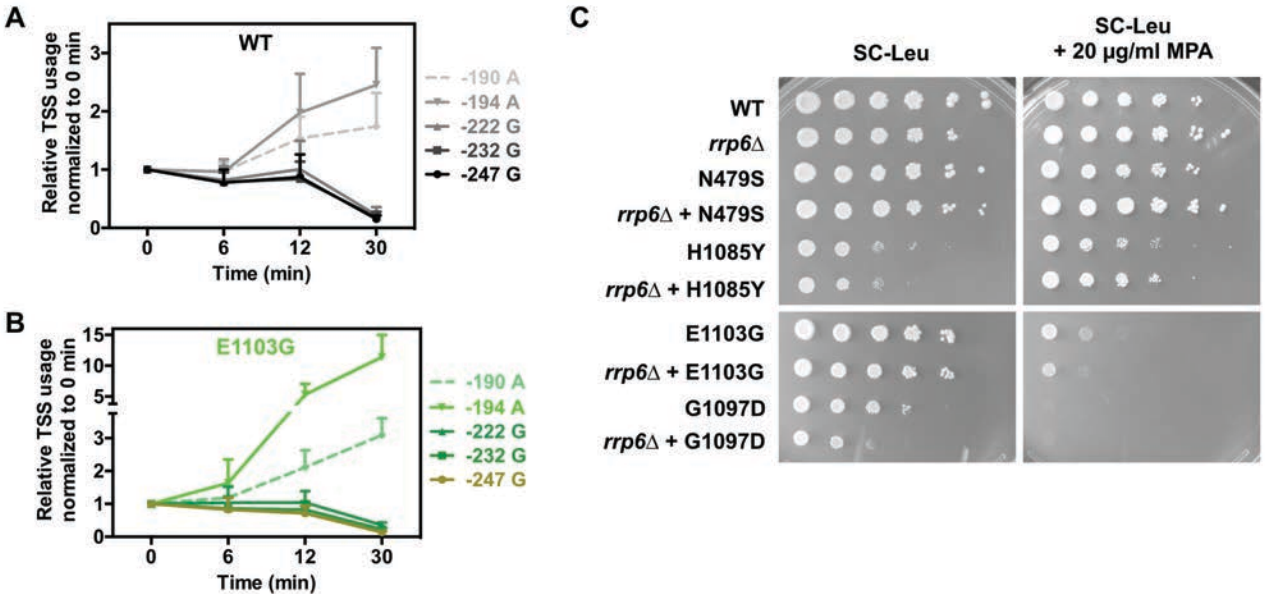
Supplementary Figure 3. Validation of GOF mutant phenotypes and formaldehyde crosslinking kinetics. (A) Primer extension analysis of start site usage at *ADH1* in WT and GOF catalytic mutants shows expected upstream shift (arrow) for GOF mutants. (B) Evaluating formaldehyde cross-linking efficiency during shut-off of *GAL1*p::*YLR454w* transcription by addition of 4% glucose. WT culture was grown to mid-log phase at 30˚C and half of the culture was isolated and fixed with 1% formaldehyde as time 0 in YPGal (GAL 0'). Then, the rest half was quickly centrifuged and resuspended in YPD (4% dextrose) containing 1% formaldehyde to obtain time 0 in YPD + formaldehyde (Glu 0'). ChIP was performed to determine Pol II occupancy over *GAL1*p::*YLR454w* (see Fig. 5A) in GAL 0' versus Glu 0'. This comparative analysis allows us to determine if there is any apparent repression during cross-linking in the presence of glucose to inhibit *GAL1*p::*YLR454w* transcription. Bar graphs show average of three biological repeats +/- SD.



Supplementary Figure 4. Impaired Mig1p nuclear import in G1097D. Representative images of nuclear localization of Mig1p-GFP in WT and catalytic mutants upon glucose (4% final) addition (left) and average Mig1p nuclear localization +/- SD (right). Data from multiple cells (WT, n=104 cells from 6 individual experiments; E1103G, n= 147 cells from 8 individual experiments; G1097D, n= 152 from 5 individual experiments; N479S, n=129 from 9 individual experiments) used to make the histogram in **Figure 7**. Arrowhead showing example of non-responding cell that does not show Mig1p-GFP foci accumulation upon glucose addition and arrow showing cell that decays Mig1p-GFP signal over time.



Supplementary Figure 5. Addition of NaOH in the media abrogates MPA effects. 10-fold serial dilutions of saturated cultures of Pol II catalytic mutants were plated for growth at 30˚C on synthetic media containing NaOH. All strains contain endogenous *IMD2* deletion (*imd2*Δ::*HIS3*), rendering them highly sensitive to MPA. Only G1097D (strongest GOF) shows sensitivity to MPA in presence of NaOH in the medium.



Supplementary Figure 6. Stabilization of *IMD2* CUTs allows *IMD2* TSSs at intermediate downstream positions to be observed. Kinetics of upstream 'G' start site loss and subsequently gain of presumptive novel 'A' sites upon MPA (20 µg/ml) treatment in WT (A) and E1103G (B) cells containing *rrp6*Δ. Error bars represent average of three biological repeats +/- SD. Note difference in scale between (A) and (B). (C) Deletion of *RRP6* does not confer MPA resistance to MPA-sensitive mutants.10-fold serial dilution of saturated cultures of Pol II mutants strains alone and in combination with *rrp6*Δ mutant plated on synthetic complete medium lacking leucine (SC-Leu) and SC-Leu medium containing 20 µg/ml MPA (final) to determine MPA sensitivity.

Supplementary Table1. List of mutants with corresponding references that used glucose shut-off experiment to determine apparent in vivo elongation rate on *GAL1*p::*YLR454w* reporter.

Mutant/Condition tested	Apparent In vivo elongation	Reference
rpb2-10 *hpr1*, *thp2*, *mft1*, *cdc73*, *rtf1*, *spt4*, *ctk1*, *ctk2* and *ppr2*.	*rpb2-10* apparent slower than WT no detectable effect	(1)
asf1	no detectable effect	(2)
set2, pob3 and set2/pob3	no detectable effect	(3)
swi2	no detectable effect	(4)
gcn5	*gcn5* apparent slower than WT	(5)
esa1 *gcn5* *esa1/gcn5*	*esa1* apparent slower than WT *gcn5* no detectable effect *esa1*/ *gcn5* apparent slower than WT	(6)
chd1 *spt5-242*	*chd1* apparent slower than WT *spt5-242* apparent slower than WT	(7)
rpb1 N488D	*rpb1* N488D apparent slower than WT	(8)
dhh1 *ccr4* *not4*	*dhh1* apparent slower than WT *ccr4* apparent slower than WT *not4* apparent slower than WT	(9)
Temperature	apparent elongation rate increases with temperature	(10)
pfd1 *dst1*	*pfd1* apparent slower than WT *dst1* no detectable effect	(11)
rpb1 E1103G	*rpb1* E1103G apparent faster than WT	(12)
rat1-1	*rat1-1* apparent faster than WT	(13)
rpb2 K864G/K865G/∆866-871	no detectable effect	(14)

Supplementary Table 2.

Experiment/Phenotype	**LOF/Slow mutant phenotype**	**GOF/Fast mutant phenotype**
Steady state Pol II occupancy (over *GAL1*p::*YLR454w*)	Decreased overall Pol II occupancy	Decreased overall Pol II occupancy
Apparent Pol II processivity defect at the 3' end of *GAL1*p::*YLR454w*	Apparent 3' end processivity defect in *rpb1* H1085Y	Apparent 3' end processivity defect in *rpb1* G1097D
Steady state Pol II occupancy in glucose vs galactose media	Subtle increase in overall Pol II occupancy in galactose for *rpb1* H1085Y	Decrease in overall Pol II occupancy in galactose for *rpb1* G1097D
In vivo gene expression (Reporters- *GAL1p*::*YLR454w*, *TEF1p*::*YLR454w*, *TEF1* and *GAL1*)	Gene expression decreases; most robust effect for *rpb1* H1085Y	Gene expression decreases; most robust effect for *rpb1* G1097D
*GAL1*p induction kinetics	Induction is delayed	Induction is delayed
Genetic interaction with pre-mRNA processing factors	Suppresion of Ts- phenotype of *rat1-1* and *xrn1*Δ/*rat1-1* mutants	Synthetic sick interactions with *xrn1*Δ and *rat1-1*
mRNA decay rate of *GAL1p*::*YLR454w*	Decay rate decreases	Decay rate decreases
In vivo elongation rate over *GAL1*p::*YLR454w* template	Apparent elongation rate slower than WT on *GAL1*p::*YLR454w* (with caveats noted in main text)	Apparent elongation rate slower than WT on *GAL1*p::*YLR454w* (with caveats noted in main text, especially for G1097D)
snR33 termination window	Shorter termination window than WT	Longer termination window than WT
Ability of GTP-sensing in absence of endogenous *IMD2*	Retain ability to sense GTP; MPA sensitivity is suppressed by addition of guanine	Retain ability to sense GTP; MPA sensitivity poorly suppressed by addition of guanine
Response to GTP depletion in absence of endogenous *IMD2*	Constitutively use downstream functional 'A' start site at *IMD2* promoter; sense GTP depletion and can further shift TSS downstream to functional 'A' site upon MPA treatment	Sense GTP depletion but cannot shift TSS downstream to functional 'A' site; instead use novel TSSs that produce non-functional *IMD2* CUT

Supplementary Table 3. List of yeast strains used in this study.

Yeast strain	Relevant mutation	Genotype	Reference	Relevant Figure
CKY1124	*RPB1* WT	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w* [pRS315 *RPB1 CEN LEU2*]	This study	Figure 1-5
CKY1110	*rpb1* N479S	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w* [pRS315 *rpb1 N479S CEN LEU2*]	This study	Figure 1-5
CKY1111	*rpb1* H1085Q	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w* [pRS315 *rpb1 H1085Q CEN LEU2*]	This study	Figure 3, and 4
CKY1112	*rpb1* E1103G	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w* [pRS315 *rpb1 E1103G CEN LEU2*]	This study	Figure 1-5
CKY1113	*rpb1* M1079R	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w* [pRS315 *rpb1 M1079R CEN LEU2*]	This study	Figure 3
CKY1114	*rpb1* G1097D	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w* [pRS315 *rpb1 G1097D CEN LEU2*]	This study	Figure 1-5
CKY1116	*rpb1* H1085Y	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w* [pRS315 *rpb1 H1085Y CEN LEU2*]	This study	Figure 1-5
CKY2064	*rpb1* L1101S	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w* [pRS315 *rpb1 L1101S CEN LEU2*]	This study	Figure 3, and 4
CKY2065	*rpb1* F1086S	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w* [pRS315 *rpb1 F1086S CEN LEU2*]	This study	Figure 3
CKY1132	*RPB1* WT	*MATα ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ rpb1Δ::CLONATMX RPB3::3XFLAG::kanmX kanmx::TEFp::YLR454w* [pRS315 *RPB1 CEN LEU2*]	This study	Figure 1 and 2

CKY1133	*rpb1* N479S	*MATα ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ rpb1Δ::CLONATMX RPB3::3XFLAG::kanmX kanmx::TEFp::YLR454w* [pRS315 *rpb1 N479S CEN LEU2*]	This study	Figure 2
CKY1135	*rpb1* H1085Y	*MATα ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ rpb1Δ::CLONATMX RPB3::3XFLAG::kanmX kanmx::TEFp::YLR454w* [pRS315 *rpb1 H1085Y CEN LEU2*]	This study	Figure 1 and 2
CKY1136	*rpb1* E1103G	*MATα ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ rpb1Δ::CLONATMX RPB3::3XFLAG::kanmX kanmx::TEFp::YLR454w* [pRS315 *rpb1 E1103G CEN LEU2*]	This study	Figure 2
CKY1137	*rpb1* G1097D	*MATα ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ rpb1Δ::CLONATMX RPB3::3XFLAG::kanmX kanmx::TEFp::YLR454w* [pRS315 *rpb1 G1097D CEN LEU2*]	This study	Figure 1 and 2
CKY1492	*RPB1* WT	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rrp6Δ::kanmx* [pRS315 *RPB1 CEN LEU2*]	This study	Figure 3, 6 and 10
CKY1493	*rpb1* N479S	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rrp6Δ::kanmx* [pRS315 *rpb1 N479S CEN LEU2*]	This study	Figure 3 and 6
CKY1494	*rpb1* H1085Q	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rrp6Δ::kanmx* [pRS315 *rpb1 H1085Q CEN LEU2*]	This study	Figure 3
CKY1495	*rpb1* F1086S	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rrp6Δ::kanmx* [pRS315 *rpb1 F1086S CEN LEU2*]	This study	Figure 3
CKY1496	*rpb1* H1085Y	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rrp6Δ::kanmx* [pRS315 *rpb1 H1085Y CEN LEU2*]	This study	Figure 3 and 6
CKY1498	*rpb1* M1079R	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rrp6Δ::kanmx* [pRS315 *rpb1 M1079R CEN LEU2*]	This study	Figure 3
CKY1499	*rpb1* L1101S	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rrp6Δ::kanmx* [pRS315 *rpb1 L1101S CEN LEU2*]	This study	Figure 3
CKY1500	*rpb1* E1103G	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rrp6Δ::kanmx* [pRS315 *rpb1 E1103G CEN LEU2*]	This study	Figure 3, 6 and 10
CKY1501	*rpb1* G1097D	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX*	This study	Figure 3 and 6

		kanmx::GAL1p::YLR454w rrp6Δ::kanmx [pRS315 *rpb1 G1097D CEN LEU2*]		
CKY1503	*RPB1* WT	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rat1-1 xrn1Δ::kanmx* [pRS315 *RPB1 CEN LEU2*]	This study	Figure 3
CKY1504	*rpb1* N479S	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rat1-1 xrn1Δ::kanmx* [pRS315 *rpb1 N479S CEN LEU2*]	This study	Figure 3
CKY1505	*rpb1* H1085Q	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rat1-1 xrn1Δ::kanmx* [pRS315 *rpb1 H1085Q CEN LEU2*]	This study	Figure 3
CKY1506	*rpb1* F1086S	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rat1-1 xrn1Δ::kanmx* [pRS315 *rpb1 N479S CEN LEU2*]	This study	Figure 3
CKY1507	*rpb1* H1085Y	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rat1-1 xrn1Δ::kanmx* [pRS315 *rpb1 H1085Y CEN LEU2*]	This study	Figure 3
CKY1509	*rpb1* M1079R	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rat1-1 xrn1Δ::kanmx* [pRS315 *rpb1 M1079R CEN LEU2*]	This study	Figure 3
CKY1510	*rpb1* L1101S	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rat1-1 xrn1Δ::kanmx* [pRS315 *rpb1 L1101S CEN LEU2*]	This study	Figure 3
CKY1511	*rpb1* E1103G	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rat1-1 xrn1Δ::kanmx* [pRS315 *rpb1 E1103G CEN LEU2*]	This study	Figure 3
CKY1512	*rpb1* G1097D	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX kanmx::GAL1p::YLR454w rat1-1 xrn1Δ::kanmx* [pRS315 *rpb1 G1097D CEN LEU2*]	This study	Figure 3
CKY1519	*RPB1* WT	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w xrn1Δ::kanmx* [pRS315 *RPB1 CEN LEU2*]	This study	Figure 3
CKY1520	*rpb1* N479S	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w xrn1Δ::kanmx*	This study	Figure 3

		[pRS315 *rpb1 N479S CEN LEU2*]		
CKY1521	*rpb1* H1085Q	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w xrn1Δ::kanmx* [pRS315 *rpb1 H1085Q CEN LEU2*]	This study	Figure 3
CKY1522	*rpb1* F1086S	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w xrn1Δ::kanmx* [pRS315 *rpb1 F1086S CEN LEU2*]	This study	Figure 3
CKY1523	*rpb1* H1085Y	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w xrn1Δ::kanmx* [pRS315 *rpb1 H1085Y CEN LEU2*]	This study	Figure 3
CKY1525	*rpb1* M1079R	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w xrn1Δ::kanmx* [pRS315 *rpb1 M1079R CEN LEU2*]	This study	Figure 3
CKY1526	*rpb1* L1101S	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w xrn1Δ::kanmx* [pRS315 *rpb1 L1101S CEN LEU2*]	This study	Figure 3
CKY1527	*rpb1* E1103G	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w xrn1Δ::kanmx* [pRS315 *rpb1 E1103G CEN LEU2*]	This study	Figure 3
CKY1528	*rpb1* G1097D	*MATa lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w xrn1Δ::kanmx* [pRS315 *rpb1 G1097D CEN LEU2*]	This study	Figure 3
CKY1533	RPB1 WT	*MATa ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w rat1-1* [pRS315 *RPB1 CEN LEU2*]	This study	Figure 3
CKY1534	*rpb1* N479S	*MATa ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w rat1-1* [pRS315 *rpb1 N479S CEN LEU2*]	This study	Figure 3
CKY1535	*rpb1* H1085Q	*MATa ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w rat1-1* [pRS315 *rpb1 H1085Q CEN LEU2*]	This study	Figure 3
CKY1536	*rpb1* F1086S	*MATa ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w rat1-1* [pRS315	This study	Figure 3

		rpb1 F1086S CEN LEU2]		
CKY1537	*rpb1* H1085Y	*MATa ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w rat1-1* [pRS315 *rpb1 H1085Y CEN LEU2*]	This study	Figure 3
CKY1539	*rpb1* M1079R	*MATa ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w rat1-1* [pRS315 *rpb1 M1079R CEN LEU2*]	This study	Figure 3
CKY1540	*rpb1* L1101S	*MATa ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w rat1-1* [pRS315 *rpb1 L1101S CEN LEU2*]	This study	Figure 3
CKY1541	*rpb1* E1103G	*MATa ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w rat1-1* [pRS315 *rpb1 E1103G CEN LEU2*]	This study	Figure 3
CKY1542	*rpb1* G1097D	*MATa ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w rat1-1* [pRS315 *rpb1 G1097D CEN LEU2*]	This study	Figure 3
CKY2066	*RPB1* WT	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w hph(hyg)::dxo1Δ* [pRS315 *RPB1 CEN LEU2*]	This study	Figure 3
CKY2067	*rpb1* N479S	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w hph(hyg)::dxo1Δ* [pRS315 *rpb1 N479S CEN LEU2*]	This study	Figure 3
CKY2068	*rpb1* H1085Q	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w hph(hyg)::dxo1Δ* [pRS315 *rpb1 H1085Q CEN LEU2*]	This study	Figure 3
CKY2069	*rpb1* F1086S	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w hph(hyg)::dxo1Δ* [pRS315 *rpb1 F1086S CEN LEU2*]	This study	Figure 3
CKY2071	*rpb1* M1079R	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w hph(hyg)::dxo1Δ* [pRS315 *rpb1 M1079R CEN LEU2*]	This study	Figure 3
CKY2072	*rpb1* L1101S	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w hph(hyg)::dxo1Δ*	This study	Figure 3

		[pRS315 *rpb1 L1101S CEN LEU2*]		
CKY2073	*rpb1* E1103G	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w hph(hyg)::dxo1Δ* [pRS315 *rpb1 E1103G CEN LEU2*]	This study	Figure 3
CKY2074	*rpb1* G1097D	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w hph(hyg)::dxo1Δ* [pRS315 *rpb1 G1097D CEN LEU2*]	This study	Figure 3
CKY2075	*rpb1* H1085Y	*MATα lys2-128∂ ura3-52 his3Δ200 leu2Δ0 or Δ1 trp1Δ63 met15Δ0 rpb1Δ::CLONATMX RPB3::3XFLAG::kanmx kanmx::GAL1p::YLR454w hph(hyg)::dxo1Δ* [pRS315 *rpb1 H1085Y CEN LEU2*]	This study	Figure 3
CKY2076	*RPB1* WT	*MATa ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ gal10Δ56 rpb1Δ::CLONATMX RPB3::TAP::KlacTRP1 rai1Δ::kanmx* [pRS315 *RPB1 CEN LEU2*]	This study	Figure 3
CKY2077	*rpb1* N479S	*MATa ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ gal10Δ56 rpb1Δ::CLONATMX RPB3::TAP::KlacTRP1 rai1Δ::kanmx* [pRS315 *rpb1 N479S CEN LEU2*]	This study	Figure 3
CKY2078	*rpb1* H1085Q	*MATa ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ gal10Δ56 rpb1Δ::CLONATMX RPB3::TAP::KlacTRP1 rai1Δ::kanmx* [pRS315 *rpb1 H1085Q CEN LEU2*]	This study	Figure 3
CKY2079	*rpb1* F1086S	*MATa ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ gal10Δ56 rpb1Δ::CLONATMX RPB3::TAP::KlacTRP1 rai1Δ::kanmx* [pRS315 *rpb1 F1086S CEN LEU2*]	This study	Figure 3
CKY2081	*rpb1* M1079R	*MATa ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ gal10Δ56 rpb1Δ::CLONATMX RPB3::TAP::KlacTRP1 rai1Δ::kanmx* [pRS315 *rpb1 M1079R CEN LEU2*]	This study	Figure 3
CKY2082	*rpb1* L1101S	*MATa ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ gal10Δ56 rpb1Δ::CLONATMX RPB3::TAP::KlacTRP1 rai1Δ::kanmx* [pRS315 *rpb1 L1101S CEN LEU2*]	This study	Figure 3
CKY2083	*rpb1* E1103G	*MATa ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ gal10Δ56 rpb1Δ::CLONATMX RPB3::TAP::KlacTRP1 rai1Δ::kanmx* [pRS315 *rpb1 E1103G CEN LEU2*]	This study	Figure 3
CKY2084	*rpb1* G1097D	*MATa ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ gal10Δ56 rpb1Δ::CLONATMX RPB3::TAP::KlacTRP1 rai1Δ::kanmx* [pRS315 *rpb1 G1097D CEN LEU2*]	This study	Figure 3

CKY2085	*rpb1* H1085Y	*MATa ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ gal10Δ56 rpb1Δ::CLONATMX RPB3::TAP::KlacTRP1 rai1Δ::kanmx* [pRS315 *rpb1 H1085Y CEN LEU2*]	This study	Figure 3
CKY283	*RPB1* WT	*MATa ura3-52 his3Δ200 leu2Δ1 or Δ0 trp1Δ63 met15Δ0 lys2-128∂ gal10Δ56 rpb1Δ::CLONATMX RPB3::TAP::KlacTRP1* [pRP112 *RPB1 CEN URA3*]	(15)	Figure 8
CKY874	*RPB1* WT	*MATa leu2Δ0 or Δ1 ura3-52 his3Δ200 met15Δ0 trp1Δ63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1Δ::CLONAT imd2Δ::HIS3* [pRS315 *RPB1 CEN LEU2*]	This study	Figure 8 and 10
CKY876	*rpb1* N1082S	*MATa leu2Δ0 or Δ1 ura3-52 his3Δ200 met15Δ0 trp1Δ63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1Δ::CLONAT imd2Δ::HIS3* [pRS315 *rpb1 N1082S CEN LEU2*]	This study	Figure 8
CKY877	*rpb1* H1085Q	*MATa leu2Δ0 or Δ1 ura3-52 his3Δ200 met15Δ0 trp1Δ63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1Δ::CLONAT imd2Δ::HIS3* [pRS315 *rpb1 H1085Q CEN LEU2*]	This study	Figure 8
CKY878	*rpb1* H1085Y	*MATa leu2Δ0 or Δ1 ura3-52 his3Δ200 met15Δ0 trp1Δ63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1Δ::CLONAT imd2Δ::HIS3* [pRS315 *rpb1 H1085Y CEN LEU2*]	This study	Figure 8 and 10
CKY879	*rpb1* F1086S	*MATa leu2Δ0 or Δ1 ura3-52 his3Δ200 met15Δ0 trp1Δ63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1Δ::CLONAT imd2Δ::HIS3* [pRS315 *rpb1 F1086S CEN LEU2*]	This study	Figure 8
CKY882	*rpb1* H1085Y /E1103G	*MATa leu2Δ0 or Δ1 ura3-52 his3Δ200 met15Δ0 trp1Δ63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1Δ::CLONAT imd2Δ::HIS3* [pRS315 *rpb1 H1085Y/E1103G CEN LEU2*]	This study	Figure 8
CKY883	*rpb1* F1084I	*MATa leu2Δ0 or Δ1 ura3-52 his3Δ200 met15Δ0 trp1Δ63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1Δ::CLONAT imd2Δ::HIS3* [pRS315 *rpb1 F1084I CEN LEU2*]	This study	Figure 8
CKY884	*rpb1* G1097D	*MATa leu2Δ0 or Δ1 ura3-52 his3Δ200 met15Δ0 trp1Δ63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1Δ::CLONAT imd2Δ::HIS3* [pRS315 *rpb1 G1097D CEN LEU2*]	This study	Figure 8
CKY885	*rpb1* E1103G	*MATa leu2Δ0 or Δ1 ura3-52 his3Δ200 met15Δ0 trp1Δ63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1Δ::CLONAT imd2Δ::HIS3* [pRS315 *rpb1 E1103G CEN LEU2*]	This study	Figure 8 and 10
CKY717/718	*dst1Δ*	*MATa ura3-52 his3Δ200 leu2Δ1 or Δ0*	This	Figure 9

		trp1∆63 met15∆0 lys2-128∂ gal10∆56 rpb1∆::CLONATMX RPB3::TAP::KlacTRP1 dst1∆::KANMX [pRP112 *RPB1 CEN URA3*]	study	
CKY982/983	*spt3∆*	*MATa ura3-52 his3∆200 leu2∆1 or ∆0 trp1∆63 met15∆0 lys2-128∂ gal10∆56 rpb1∆::CLONATMX RPB3::TAP::KlacTRP1 spt3∆::kanmx* [pRP112 *RPB1 CEN URA3*]	This study	Figure 9
CKY1164/1165	*spt4∆*	*MATa ura3-52 his3∆200 leu2∆1 or ∆0 trp1∆63 met15∆0 lys2-128∂ gal10∆56 rpb1∆::CLONATMX RPB3::TAP::KlacTRP1 spt4∆::kanmx* [pRP112 *RPB1 CEN URA3*]	This study	Figure 9
CKY986/987	*gal11∆*	*MATa ura3-52 his3∆200 leu2∆1 or ∆0 trp1∆63 met15∆0 lys2-128∂ gal10∆56 rpb1∆::CLONATMX RPB3::TAP::KlacTRP1 gal11∆::kanmx* [pRP112 *RPB1 CEN URA3*]	This study	Figure 9
CKY736/737	*sgf73∆*	*MATa ura3-52 his3∆200 leu2∆1 or ∆0 trp1∆63 met15∆0 lys2-128∂ gal10∆56 rpb1∆::CLONATMX RPB3::TAP::KlacTRP1 sgf73∆::KANMX* [pRP112 *RPB1 CEN URA3*]	This study	Figure 9
CKY970/971	*dst1∆*	*MATa leu2∆0 or ∆1 ura3-52 his3∆200 met15∆0 trp1∆63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1∆::CLONAT imd2∆::HIS3 dst1∆::kanmx* [pRP112 *RPB1 CEN URA3*]	This study	Figure 9
CKY966/967	*spt3∆*	*MATa leu2∆0 or ∆1 ura3-52 his3∆200 met15∆0 trp1∆63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1∆::CLONAT imd2∆::HIS3 spt3∆::kanmx* [pRP112 *RPB1 CEN URA3*]	This study	Figure 9
CKY968/969	*spt4∆*	*MATa leu2∆0 or ∆1 ura3-52 his3∆200 met15∆0 trp1∆63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1∆::CLONAT imd2∆::HIS3 spt4∆::kanmx* [pRP112 *RPB1 CEN URA3*]	This study	Figure 9
CKY974/975	*gal11∆*	*MATa leu2∆0 or ∆1 ura3-52 his3∆200 met15∆0 trp1∆63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1∆::CLONAT imd2∆::HIS3 gal11∆::kanmx* [pRP112 *RPB1 CEN URA3*]	This study	Figure 9
CKY972/973	*sgf73∆*	*MATa leu2∆0 or ∆1 ura3-52 his3∆200 met15∆0 trp1∆63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1∆::CLONAT imd2∆::HIS3 sgf73∆::kanmx* [pRP112 *RPB1 CEN URA3*]	This study	Figure 9
CKY728/729	*paf1∆*	*MATa ura3-52 his3∆200 leu2∆1 or ∆0 trp1∆63 met15∆0 lys2-128∂ gal10∆56 rpb1∆::CLONATMX RPB3::TAP::KlacTRP1 paf1∆::KANMX* [pRP112 *RPB1 CEN URA3*]	This study	Figure 9
CKY990/991	*bur2∆*	*MATa ura3-52 his3∆200 leu2∆1 or ∆0 trp1∆63 met15∆0 lys2-128∂ gal10∆56 rpb1∆::CLONATMX RPB3::TAP::KlacTRP1 bur2∆::kanmx* [pRP112 *RPB1 CEN URA3*]	This study	Figure 9
CKY988/989	*pop2∆*	*MATa ura3-52 his3∆200 leu2∆1 or ∆0*	This	Figure 9

		trp1∆63 met15∆0 lys2-128∂ gal10∆56 rpb1∆::CLONATMX RPB3::TAP::KlacTRP1 pop2∆::kanmx [pRP112 *RPB1 CEN URA3*]	study	
CKY976/977	*paf1∆*	*MATa leu2∆0 or ∆1 ura3-52 his3∆200 met15∆0 trp1∆63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1∆::CLONAT imd2∆::HIS3 paf1∆::kanmx* [pRP112 *RPB1 CEN URA3*]	This study	Figure 9
CKY980/981	*bur2∆*	*MATa leu2∆0 or ∆1 ura3-52 his3∆200 met15∆0 trp1∆63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1∆::CLONAT imd2∆::HIS3 bur2∆::kanmx* [pRP112 *RPB1 CEN URA3*]	This study	Figure 9
CKY978/979	*pop2∆*	*MATa leu2∆0 or ∆1 ura3-52 his3∆200 met15∆0 trp1∆63 lys2-128∂ RPB3::TAP::KlacTRP1 rpb1∆::CLONAT imd2∆::HIS3 pop2∆::kanmx* [pRP112 *RPB1 CEN URA3*]	This study	Figure 9
CKY2430/2431	*RPB1* WT	*MATα his3∆200 leu2∆0 or 1 ura3-52 lys2-128∂ met15∆0 trp1∆63 RPB3::TAP::KlacTRP1 rpb1∆::CLONATMX MIG1::EGFP::kanmx* [pRS315 *RPB1 CEN LEU2*]	This study	Figure 7
CKY2432/2433	*rpb1* E1103G	*MATα his3∆200 leu2∆0 or 1 ura3-52 lys2-128∂ met15∆0 trp1∆63 RPB3::TAP::KlacTRP1 rpb1∆::CLONATMX MIG1:: EGFP:: kanmx* [pRS315 *rpb1 E1103G CEN LEU2*]	This study	Figure 7
CKY2434/2435	*rpb1* G1097D	*MATα his3∆200 leu2∆0 or 1 ura3-52 lys2-128∂ met15∆0 trp1∆63 RPB3::TAP::KlacTRP1 rpb1∆::CLONATMX MIG1:: EGFP:: kanmx* [pRS315 *rpb1 G1097D CEN LEU2*]	This study	Figure 7
CKY2467/2468	*rpb1* N479S	*MATα his3∆200 leu2∆0 or 1 ura3-52 lys2-128∂ met15∆0 trp1∆63 RPB3::TAP::KlacTRP1 rpb1∆::CLONATMX MIG1:: EGFP:: kanmx* [pRS315 *rpb1 N479S CEN LEU2*]	This study	Figure 7

Supplementary Table 4. List of plasmids used in this study.

Plasmid	Description	Genotype	Reference
pCK518	pRS316 *RPB1*	ampr ColE1 ori *URA3 CEN ARS*	(15)
pCK859	pRS315 *RPB1*	ampr ColE1 ori *LEU2 CEN ARS*	(15)
pCK856	pRS315 *rpb1* N479S	ampr ColE1 ori *LEU2 CEN ARS*	(15)
pCK864	pRS315 *rpb1* L1101S	ampr ColE1 ori *LEU2 CEN ARS*	(15)
pCK867	pRS315 *rpb1* G1097D	ampr ColE1 ori *LEU2 CEN ARS*	(15)
pCK870	pRS315 *rpb1* H1085Y	ampr ColE1 ori *LEU2 CEN ARS*	(15)
pCK871	pRS315 *rpb1* F1086S	ampr ColE1 ori *LEU2 CEN ARS*	(15)
pCK886	pRS315 *rpb1* N1082S	ampr ColE1 ori *LEU2 CEN ARS*	(15)
pCK887	pRS315 *rpb1* H1085Q	ampr ColE1 ori *LEU2 CEN ARS*	(15)
pCK890	pRS315 *rpb1* H1085Y/E1103G	ampr ColE1 ori *LEU2 CEN ARS*	(15)
pCK955	pRS315 *rpb1* F1084I	ampr ColE1 ori *LEU2 CEN ARS*	(15)
pCK960	pRS315 *rpb1* E1103G	ampr ColE1 ori *LEU2 CEN ARS*	(15)

Supplementary Table 5. List of primers used in this study.

Primer	Description / Relevant gene	Sequence	Function	Reference
CKO667	KANMX - GAL1p/TEF1p F	TAACGCCGCCATCCAGTGTC	ChIP/RT-PCR	This study
CKO668	KANMX -TEF1p/R	GCGCGGAGTCCGAGAAAATC	ChIP/RT-PCR	This study
CKO669	KANMX -GAL1p/R	GCGAGGCACATCTGCGTTTC	ChIP/RT-PCR	This study
CKO672	YLR454w 0.2 kb F	GTACCGTCAGGCTAAAATCCGTTCG	ChIP/RT-PCR	This study
CKO673	YLR454w 0.2 kb R	GACCCCATTGAGCCAGTATTGTGA	ChIP/RT-PCR	This study
CKO653	YLR454w 1 kb F	ACAGGTTCAGAAATGAGATGCCAG	ChIP/RT-PCR Northern probe amplification	This study
CKO654	YLR454w 1 kb R	TTGGCGTGGCTTTGATGTTTTCG	ChIP/RT-PCR	This study
CKO655	YLR454w 2 kb F	AGCGACTTCATGTTCCAGCAACT	ChIP/RT-PCR	This study
CKO656	YLR454w 2 kb R	CTTGGCATAAAAACCGACCTAGCAC	ChIP/RT-PCR Northern probe amplification	This study
CKO657	YLR454w 3 kb F	TGGGGCCAACTAAAGGAGTTAC	ChIP/RT-PCR	This study
CKO658	YLR454w 3 kb R	CCAAAAGTTTGGCTGCGTTG	ChIP/RT-PCR	This study
CKO659	YLR454w 4 kb F	ACCTCCACTAAGCTCTACACAAAGT	ChIP/RT-PCR	This study
CKO660	YLR454w 4 kb R	TTCTGGGCACGAACAACGAG	ChIP/RT-PCR	This study
CKO661	YLR454w 5 kb F	GTCCCAACGGGTTCAAGGCATCC	ChIP/RT-PCR	This study
CKO662	YLR454w 5 kb R	ATCGGCATCAGCGTTGTGGT	ChIP/RT-PCR	This study
CKO663	YLR454w 6 kb F	ACTGTTGAAATGGAACGAGGACGC	ChIP/RT-PCR	This study
CKO664	YLR454w 6 kb R	CCTTCTGGTATCGCTTCCATACTCG	ChIP/RT-PCR	This study
CKO665	YLR454w 7 kb F	ACACAGTCGGTTTGGCGAG	ChIP/RT-PCR	This study
CKO666	YLR454w 7 kb R	ACTCGACAAAGTGGTCTCAACG	ChIP/RT-PCR	This study
CKO1780	YLR454w 8 kb F	GAGGGTCACAGATCTATTACTTGCCC	ChIP/RT-PCR	(12)
CKO1781	YLR454w 8 kb R	GTTGTGAGTTGCTTCAGTGGTGAAGTG	ChIP/RT-PCR	(12)
CKO946	TEL-VI 1	GCGTAACAAAGCCATAATGCCTCC	ChIP/RT-PCR	(12)
CKO947	TEF-VI 2	CTCGTTAGGATCACGTTCGAATCC	ChIP/RT-PCR	(12)
CKO1268	TEF1 F	ATGGGTAAAGAGAAGTCTC	Northern probe	This study

		AC	amplification	
CKO1269	TEF1 R	CAGCCTTTTGAGCAGCCTTGGTA	Northern probe amplification	This study
CKO1172	SCR1 F	AGGCTGTAATGGCTTTCTGGTGGGATGGGA	Northern probe amplification	(16)
CKO1173	SCR1 R	GATATGTGCTATCCCGGCCGCCTCCATCAC	Northern probe amplification	(16)
CKO1692	GAL1 F	GTGCCCGAGCATAATTAAGAAAT	Northern probe amplification	This study
CKO1693	GAL1 R	TGTAGTGACTTCTACCACTCTTA	Northern probe amplification	This study
CKO1782	snR33 +12 Top Strand	CTCTTTGTACGATGGTGTCACTC	Northern probe amplification	(12)
CKO1783	snR33 3'UTR low +347	AATTGTTAAATGCATTGGCTCG	Northern probe amplification/labeling	(12)

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