

Table S1. *C. albicans* stains used in this study

Strain	Parent / background	Genotype	Source
SC5314		Wild type	(1)
CAI4	SC5314	<i>ura3::1 imm434/ura3::1 imm434</i>	(1)
BWP17	SC5314	<i>ura3::1 imm434/ura3::1 imm434 his1::hisG/his1::hisG arg4::hisG/arg4::hisG</i>	(2)
HLY4176	CAI4	<i>ura3::1 imm434/ura3::1 imm434 MET3/met3::MET3p-UME6_{C778/785S}-13MYC-URA3</i>	(3)
HLY3578	BWP17	<i>ura3::1 imm434/ura3::1 imm434 his1::hisG/his1::hisG arg4::hisG/arg4::hisG ADE2/ade2::MAL2p-MYC-HGC1-URA3</i>	(4)
YLC12	HLY4176	<i>ura3::1 imm434/ura3::1 imm434 MET3/met3::MET3p-UME6_{C778/785S},_{S437A}-13MYC-URA3</i>	This study
YLC13	SN250	<i>UME6_{S437A}::UME6_{S437A} ENO1::CAS9-SAT1</i>	This study
SN250	SC5314	<i>his1Δ/his1Δ, leu2Δ::C.d. HIS1/leu2Δ::C.m. LEU2, arg4Δ/arg4Δ, URA3/ura3Δ::imm434 IRO1/iro1Δ::imm434</i>	(5)
<i>ptc2</i>	SN250	<i>his1Δ/his1Δ, leu2Δ::C.d. HIS1/leu2Δ::C.m. LEU2, arg4Δ/arg4Δ, URA3/ura3Δ::imm434 IRO1/iro1Δ::imm434 ptc2Δ::C.d. HIS1/ptc2Δ::C.m. LEU2</i>	(5)
YLC15	SN250	<i>his1Δ/his1Δ, leu2Δ::C.d. HIS1/leu2Δ::C.m. LEU2, arg4Δ/arg4Δ, ura3Δ/ura3Δ::imm434 IRO1/iro1Δ::imm434 SSN3::SSN3-13MYC-URA3</i>	This study
HLY4080	BWP17	<i>ubr1::ARG4/ubr1::HIS1 ura3::1 imm434/ura3::1 imm434 his1::hisG/his1::hisG arg4::hisG/arg4::hisG</i>	This study
HLY3586	SN148	<i>ura3::imm434/ura3::imm434 iro1::imm434/iro1::imm434 grr1::C.d.HIS1/grr1::C.m.LEU2 arg4Δ/arg4Δ leu2Δ/leu2Δ his1Δ/his1Δ</i>	(4)

References

1. Fonzi, W. A. & Irwin, M. Y. (1993) *Genetics* **134**, 717-28.
2. Wilson, R. B., Davis, D. & Mitchell, A. P. (1999) *J Bacteriol* **181**, 1868-74.
3. Lu, Y., Su, C., Solis, N. V., Filler, S. G. & Liu, H. (2003) *Cell Host & Microbe* **14**, 499-509
4. Wang, A., Lane, S., Tian, Z., Sharon, A., Hazan, I. & Liu, H. (2007) *Eukaryot Cell* **6**, 253-61.
5. Noble, S, M., French, S., Kohn, L, A., Chen, V., & Johnson, A, D. (2010) *Nat Genet* **42**, 590-8