

**Table S2.** *Candida albicans* strains used in this study

Strain	Genotype	Source
SN152	<i>ura3Δ::λimm434::URA3-IRO1</i> <i>arg4::hisG his1::hisG leu2::hisG</i> <i>ura3Δ::λimm434</i> <i>arg4::hisG his1::hisG leu2::hisG</i>	[1]
W0-1	wild-type isolate	[2]
SN250	<i>ura3Δ::λimm434::URA3-IRO1</i> <i>arg4::hisG his1::hisG leu2::hisG::CdHIS1</i> <i>ura3Δ::λimm434</i> <i>arg4::hisG his1::hisG leu2::hisG::CmLEU2</i>	[3]
TF120	<i>ura3Δ::λimm434::URA3-IRO1</i> <i>arg4::hisG his1::hisG leu2::hisG zfu2Δ::CdHIS1</i> <i>ura3Δ::λimm434</i> <i>arg4::hisG his1::hisG leu2::hisG zfu2Δ::CmLEU2</i>	[3]
TF141	<i>ura3Δ::λimm434::URA3-IRO1</i> <i>arg4::hisG his1::hisG leu2::hisG zcf8Δ::CdHIS1</i> <i>ura3Δ::λimm434</i> <i>arg4::hisG his1::hisG leu2::hisG zcf8Δ::CmLEU2</i>	[3]
TF55	<i>ura3Δ::λimm434::URA3-IRO1</i> <i>arg4::hisG his1::hisG leu2::hisG try4Δ::CdHIS1</i> <i>ura3Δ::λimm434</i> <i>arg4::hisG his1::hisG leu2::hisG try4Δ::CmLEU2</i>	[3]
TF116	<i>ura3Δ::λimm434::URA3-IRO1</i> <i>arg4::hisG his1::hisG leu2::hisG orf19.5910Δ::CdHIS1</i> <i>ura3Δ::λimm434</i> <i>arg4::hisG his1::hisG leu2::hisG orf19.5910Δ::CmLEU2</i>	[3]
TF114	<i>ura3Δ::λimm434::URA3-IRO1</i> <i>arg4::hisG his1::hisG leu2::hisG fgr17Δ::CdHIS1</i> <i>ura3Δ::λimm434</i> <i>arg4::hisG his1::hisG leu2::hisG fgr17Δ::CmLEU2</i>	[3]
JCP721	<i>ura3Δ::λimm434::URA3-IRO1</i> <i>arg4::hisG his1::hisG leu2::hisG zfu2Δ::CdHIS1 rps10Δ::ZFU2-SAT1</i> <i>ura3Δ::λimm434</i> <i>arg4::hisG his1::hisG leu2::hisG zfu2Δ::CmLEU2 RPS10</i>	This work
JCP758	<i>ura3Δ::λimm434::URA3-IRO1</i> <i>arg4::hisG his1::hisG leu2::hisG zcf8Δ::CdHIS1 rps10Δ::ZCF8-SAT1</i> <i>ura3Δ::λimm434</i> <i>arg4::hisG his1::hisG leu2::hisG zcf8Δ::CmLEU2 RPS10</i>	This work
JCP686	<i>ura3Δ::λimm434::URA3-IRO1</i> <i>arg4::hisG his1::hisG leu2::hisG try4Δ::CdHIS1 rps10Δ::TRY4-SAT1</i> <i>ura3Δ::λimm434</i> <i>arg4::hisG his1::hisG leu2::hisG try4Δ::CmLEU2 RPS10</i>	This work
JCP759	<i>ura3Δ::λimm434::URA3-IRO1</i> <i>arg4::hisG his1::hisG leu2::hisG orf19.5910Δ::CdHIS1 rps10Δ::ORF19.5910-SAT1</i> <i>ura3Δ::λimm434</i> <i>arg4::hisG his1::hisG leu2::hisG orf19.5910Δ::CmLEU2 RPS10</i>	This work
JCP837	<i>ura3Δ::λimm434::URA3-IRO1</i> <i>arg4::hisG his1::hisG leu2::hisG FGR17::AgTEF1p-NAT1-AgTEF1UTR-TDH3p-FGR17</i> <i>ura3Δ::λimm434</i> <i>arg4::hisG his1::hisG leu2::hisG FGR17</i>	This work
JCP906	<i>ura3Δ::λimm434::URA3-IRO1</i> <i>arg4::hisG his1::hisG leu2::hisG::CdHIS1 UME6::AgTEF1p-NAT1-AgTEF1UTR-TDH3p-UME6</i> <i>ura3Δ::λimm434</i> <i>arg4::hisG his1::hisG leu2::hisG::CmLEU2 UME6</i>	This work
JCP907	<i>ura3Δ::λimm434::URA3-IRO1</i> <i>arg4::hisG his1::hisG leu2::hisG::CdHIS1 WOR3::AgTEF1p-NAT1-AgTEF1UTR-TDH3p-WOR3</i> <i>ura3Δ::λimm434</i> <i>arg4::hisG his1::hisG leu2::hisG::CmLEU2 WOR3</i>	This work

1. Noble SM, Johnson AD (2005) Strains and strategies for large-scale gene deletion studies of the diploid human fungal pathogen *Candida albicans*. *Eukaryot Cell* 4:298-309.
2. Slutsky B, Staebell M, Anderson J, Risen L, Pfaller M, et al. (1987) "White-opaque transition": a second high-frequency switching system in *Candida albicans*. *J Bacteriol* 169:189-197.
3. Homann OR, Dea J, Noble SM, Johnson AD (2009) A phenotypic profile of the *Candida albicans* regulatory network. *PLoS Genet* 5:e1000783.