

Table S1. *Candida* strains used in this study.

Strain	Source/Reference
<i>C. albicans</i> SC5314 Caf2.1-dTom-NATr	R. L. Gratacap, J. F. Rawls, and R. T. Wheeler, <i>Dis. Model. Mech.</i> 6: 1260–70, 2013, doi: 10.1242/dmm.012039
<i>C. albicans</i> WT-GFP (SC5314 (Peno1-yEGFP ₃ -NAT))	R. T. Wheeler, D. Kombe, S. D. Agarwala, and G. R. Fink, <i>PLoS Pathog.</i> 4: 1–12, 2008, doi: 10.1371/journal.ppat.1000227
<i>C. albicans</i> SC5314 Caf2:FR	A. C. Bergeron, B. G. Seman, J. H. Hammond, L. S. Archambault, D. A. Hogan, and R. T. Wheeler, <i>Infect. Immun.</i> , 2017, doi:10.1128/IAI.00475-17
<i>C. parapsilosis</i> clinical isolate 4175 (A010) CpURA3/CpURA3 ENO1/ENO1::GFP-NAT1	S. Gonja, B. Larson, and C. A. Gale, <i>Yeast</i> , 33(2): 63–9, 2016, doi: 10.1002/yea.3141
<i>C. parapsilosis</i> clinical isolate 4175 (A010) CpURA3/CpURA3 ENO1-mCherry-NAT1/ENO1	