

**Supplemental Table S2**

<b>Strain</b>	<b>Relevant Genotype</b>	<b>Reference</b>
BWP17	<i>ura3Δ/Δ his1Δ/Δ arg4Δ/Δ</i>	1
GP1 (“Wild-type”)	<i>ura3Δ/Δ:URA3 his1Δ/Δ:HIS1 arg4Δ/Δ:ARG4</i>	2
<i>pmc1Δ/Δ</i>	<i>ura3Δ/Δ:URA3 his1Δ/Δ arg4Δ/Δ pmc1Δ:HIS1/pmc1Δ:ARG4</i>	This study
<i>pmc1Δ/Δ + PMC1</i>	<i>ura3Δ/Δ:URA3: PMC1 his1Δ/Δ arg4Δ/Δ pmc1Δ:HIS1/pmc1Δ:ARG4</i>	This study
<i>vcx1Δ/Δ</i>	<i>ura3Δ/Δ:URA3 his1Δ/Δ arg4Δ/Δ vcx1Δ:HIS1/vcx1Δ:ARG4</i>	This study
<i>vcx1Δ/Δ + VCX1</i>	<i>ura3Δ/Δ:URA3: VCX1 his1Δ/Δ arg4Δ/Δ vcx1Δ:HIS1/vcx1Δ:ARG4</i>	This study

1. Wilson RB, Davis D, Mitchell AP. Rapid hypothesis testing with *Candida albicans* through gene disruption with short homology regions. *Journal of Bacteriology*. 1996;181(6):1868 - 74.
2. Luna-Tapia A, Willems HME, Parker JE, Tournu H, Barker KS, Nishimoto AT, et al. Loss of Upc2p-Inducible *ERG3* Transcription Is Sufficient To Confer Niche-Specific Azole Resistance without Compromising *Candida albicans* Pathogenicity. *mBio*. 2018;9(3):e00225-18.