

Supplementary Information

Table S1: Strains used in this study

Name	Genotype	Reference
<i>C. albicans</i> SC5314	wild-type	[1]
<i>C. albicans</i> CA-MT14	<i>hat1</i> Δ/Δ	[2]
<i>C. albicans</i> CA-MT569	<i>gcn5</i> Δ/Δ	[3]
<i>C. dubliniensis</i> CD36	wild-type	[4]
<i>C. parapsilosis</i> ATCC22019	wild-type	www.attc.org
<i>C. krusei</i> ATCC6258	wild-type	www.attc.org
<i>C. tropicalis</i> AKH2143	wild-type	clinical isolate
<i>C. guilliermondii</i> AKH2119	wild-type	clinical isolate
<i>C. kefyr</i> AKH2057	wild-type	clinical isolate
<i>C. lipolytica</i> AKH2269	wild-type	clinical isolate
<i>C. lusitaniae</i> AKH2124	wild-type	clinical isolate
<i>C. glabrata</i> ATCC2001	wild-type	www.attc.org
<i>S. cerevisiae</i> BY4741	MATa <i>his3</i> Δ1 <i>leu2</i> Δ0 <i>LYS2</i> <i>met15</i> Δ0 <i>ura3</i> Δ0	[5]

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