

1    **Supplemental Tables and Figure.**

2    Table S1. *C. albicans* strains used in this work.

3    Table S2. Primers used in this work.

4    Table S3. List of genes differentially expressed during colonization relative to laboratory  
5    grown cells, in at least one strain in one organ.

6    Table S4. List of genes showing common differential expression in WT *C. albicans* cells  
7    colonizing the cecum and the ileum.

8    Table S5. Expression of selected genes in WT and *efg1*<sup>-</sup> null mutant cells colonizing the  
9    GI tract, determined by microarray.

10   Table S6. List of genes differentially expressed between *efg1*<sup>-</sup> null mutant and WT cells  
11   colonizing organs of the GI tract.

12   Table S7. List of genes differentially expressed between *efg1*<sup>-</sup> *efh1*<sup>-</sup> double null mutant  
13   and WT cells colonizing organs of the GI tract.

14   Table S8. List of genes differentially expressed between *efg1*<sup>-</sup> *cph1*<sup>-</sup> double null mutant  
15   and WT cells colonizing organs of the GI tract.

16   Table S9. List of genes differentially expressed in all *efg1*<sup>-</sup> strains tested relative to WT  
17   cells colonizing organs of the GI tract.

18   Figure S1. Colonization levels of strains that were co-inoculated into BALB/c mice.

19   For the experiments shown in Fig. 7, colonization levels of each individual strain were  
20   calculated and graphed. Brackets at top indicate the two strains that were co-inoculated  
21   into the same mouse. Each symbol represents CFU/gm in fecal pellets from an individual  
22   mouse; bars are the geometric means. Days of sample collection (post-inoculation) are  
23   indicated.

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

Strain	Genotype	Reference or Source
CAI-4	SC5314 $\Delta ura3::imm434/\Delta ura3::imm434$	(3)
CKY101	CAI-4 (pDB152)	(1)
HLC67	CAI-4 $efg1::hisG/efg1::hisG$	(5)
Can 36	CAI-4 $efg1::hisG/efg1::hisG cph1::hisG/cph1::hisG$	(5)
CKY136	CAI-4 $efg1::hisG/efg1::hisG$ (pDB152)	(4)
CKY138	Can 36 (pDB152)	(4)
H1.22-1	CAI-4 $efg1::hisG/efg1::hisG efh1::hisG/efh1::hisG$	(2)
JPY109	H1.22-1 (pRC3915)	This study
JPY 105	CAI-4 $ade2::pISAT$	(6)
JPY 106	HLC67 $ade2::pISAT$	This study
JPY 110	HLC67 $rps1::ACT1pr-SOD5$	This study

## References

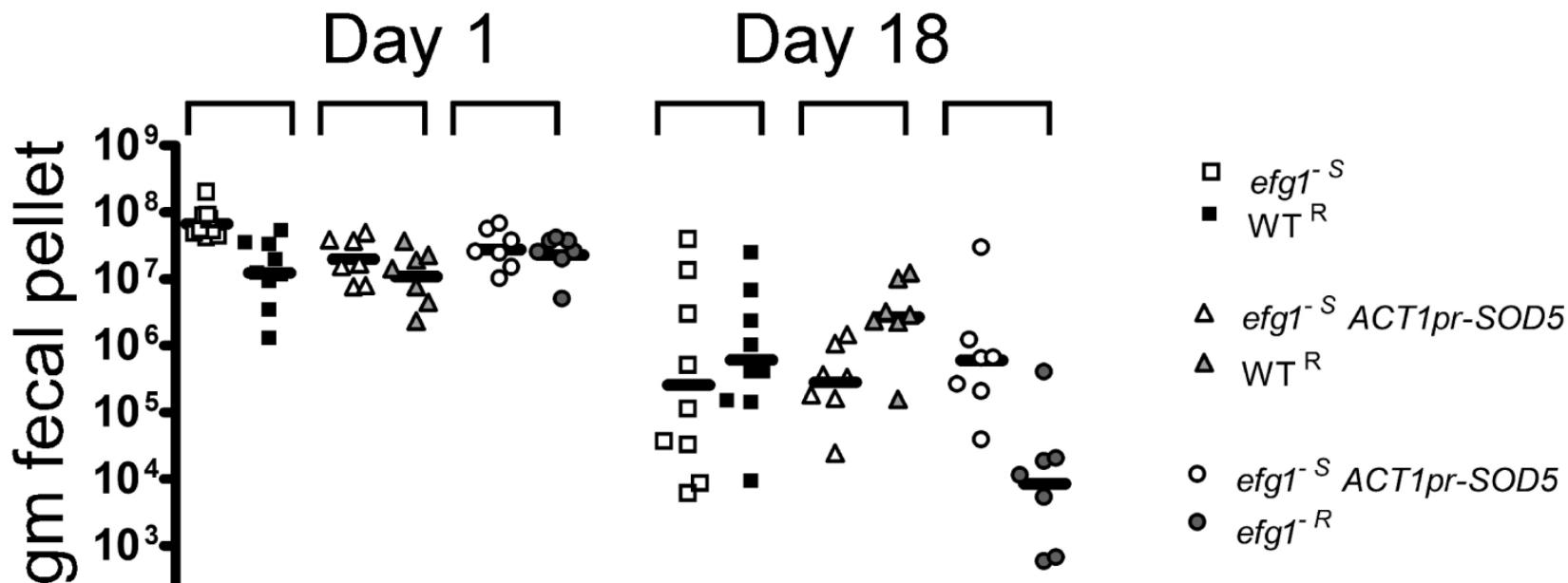
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Table S2. Primers used in this study.

<b>Cloning primers</b>		
<i>SOD5</i>	SOD5F1 cloning	5'-CCCAAGCTTGGGATCATGAAGTATTGTCCAT-3'
<i>SOD5</i>	SOD5R1 cloning	5'-CTAAGCTAGAGCCTCTGGATTCAATTACT-3'
<b>RT-PCR primers</b>		
<i>ACT1</i>	ABAf3	5'-TATGAAAGTTAACGATTATTGCTCCACCAGAAA-3'
<i>ACT1</i>	ABAR3	5'-GGAAAGTAGACAATGAAGCCAAGATAGAAC-3'
<i>CIP1</i>	CIP1F	5'-AAGTTGATGAATATGCTCCAGGGTTTTAG-3'
<i>CIP1</i>	CIP1R	5'-TCTGGTAATTCTCTGGATTAGGATTAGTAGCA-3'
<i>CTN1</i>	CTN1F	5'-GTTTCTGCTAATGGGTTGGTATTGGTTATA-3'
<i>CTN1</i>	CTN1R	5'-GTGTTTGGCTAGCTTGTTCCTCTTCTG-3'
<i>HYR1</i>	HYR1F1	5'-GTTGCTAATCCAGTTACTACTTCCACAGA-3'
<i>HYR1</i>	HYR1R1	5'-GGTTAGGTTCCATCAAAGCCAGTCATA-3'
<i>ICL1</i>	ICL1F	5'-AGCATTGGCTGTTGATGACTTCTCTA-3'
<i>ICL1</i>	ICL1R	5'-GGTAACACTGACCATCTTC-3'
<i>INO1</i>	INO1F	5'-GTTGGGTGGTCACAACAAAATAAGTAT-3'
<i>INO1</i>	INO1R	5'-TTCAGTAGGCCACCAAGTCAATA-3'

<i>PHO89</i>	PHO89F	5'-TGTGTTACGCTGCCGGTGCTTAGTTG-3'
<i>PHO89</i>	PHO89R	5'-CATTCTCCAGTTGAGGGATTCCATTCTCT-3'
<i>RBT5</i>	RBT5F	5'-AATCTTGGCCACTCCATTGTTCATCAT-3'
<i>RBT5</i>	RBT5R	5'-AGCGGCAGTAGATTGGCAGTG-3'
<i>RIM101</i>	RIM101F1	5'-AGCAATACCACGACCATCACGGATACTACTCA-3'
<i>RIM101</i>	RIM101R1	5'-TGCTTGGCCTCTTTGTTGTTGTTCATCA-3'
<i>SOD5</i>	SOD5F1	5'-GAGGGACACGGCAATGCTAACACAACAT-3'
<i>SOD5</i>	SOD5R1	5'-ACCTGCTGCCAAGAGACCATTACTACTG-3'
<i>WH11</i>	WH11F	5'-CTCAAAAATCCACTCCAGAACAAAT-3'
<i>WH11</i>	WH11R	5'-TGGAGTCAAAAATAGCATCA-3'
<i>WOR1</i>	WOR1F1	5'-AACAACGGTAGCAATGCCCACTG-3'
<i>WOR1</i>	WOR1R1	5'-ATTGCCATTACCACCAACTAACACTCCACTG-3'



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