

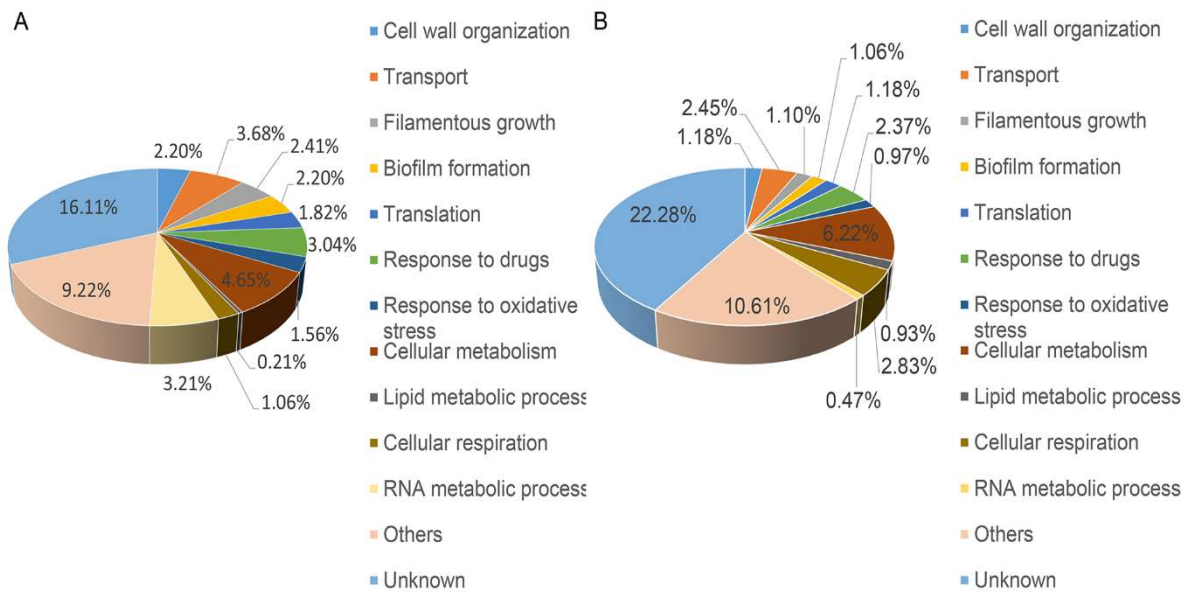
**Table S1.** The *C. albicans* strains used in this study.

Strain name	Genotype	Source <sup>#</sup>
SC5314	Wild type	[1]
<i>SFP1</i> mutants		
<i>sfp1</i> Δ/Δ	<i>sfp1</i> Δ::FRT/ <i>sfp1</i> Δ::FRT	[2]
<i>SFP1</i> RD	<i>sfp1</i> Δ::SFP1-FRT/ <i>sfp1</i> Δ::SFP1-FRT	[2]

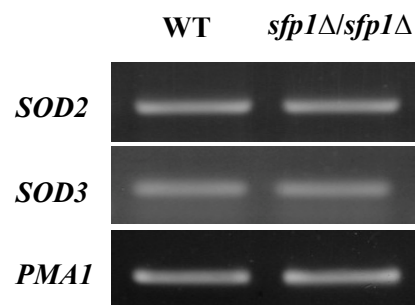
1. Gillum, A.M.; Tsay, E.Y.; Kirsch, D.R. Isolation of the *Candida albicans* gene for orotidine-5'-phosphate decarboxylase by complementation of *S. cerevisiae* *ura3* and *E. coli* *pyrF* mutations. *Mol. Gen. Genet.* **1984**, *198*, 179-182.
2. Chen, H.-F.; Lan, C.-Y. Role of *SFP1* in the regulation of *Candida albicans* biofilm formation. *PLoS One* **2015**, *10*, doi:10.1371/journal.pone.0129903.

**Table. S2.** Oligonucleotides used in this study.

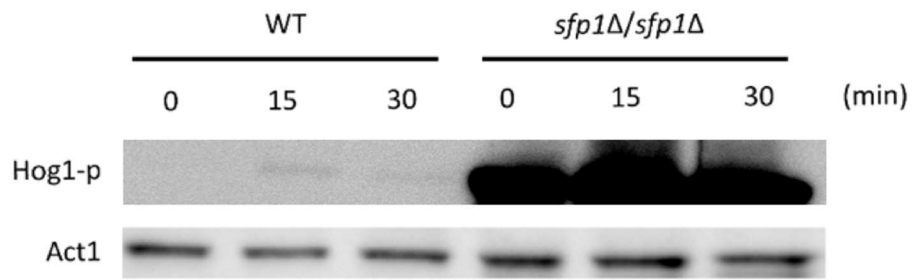
Name	Sequence (5' to 3')
SOD1-F	TGTTGTCAGAGGTGATTCAAAAGTC
SOD1-R	GTTGGAGCGGATTCGGATT
SOD4-F	TTTGAGCCAGCAAACAATGG
SOD4-R	CACCTGAAGGCAATCCAGTTAAA
SOD5-F	AAGGATTGCCCTCTGATATTGG
SOD5-R	GATGCTGGCACTGGTTTTTCA
CAT1-F	ATTTCATCCACACCCAAAGAGA
CAT1-R	CAAGTAATCCCAAACATGTTAGCA
GPX2-F	TTGGTGTGACTTTCCCCGTATT
GPX2-R	CCGGGCTTTTGAGACTTCAA
GCS1-F	CCACAAGCATTAAACAATTCAACTACA
GCS1-R	ACCCAATGTGCCGTGGTT
GTT11-F	AATATTTTCCAAAACAAACGTGGGG CTGA
GTT11-R	CCAATTCTTGAGCCTTGATGTATTGGGTTC
SSK1-F	TAAATGGAAAAAGGGAGGGTTTC
SSK1-R	AATCCCTGATTTCACTGGCAAT
SHO1-F	ACTGGTGCCATCATTAAACCCTAA
SHO1-R	TGATGAGCTGATCCACCAATAGA
CAP1-F	ACCGTGAACGTAAAGAACG
CAP1-R	GCTACCACCAGTATATTAGCC



**Figure S1.** Gene ontology (GO) distribution of *C. albicans* genes regulated by *SFP1*. The 1145 and 1220 genes upregulated (A) or downregulated (B) in the *sfp1Δ/sfp1Δ* mutant were classified by biological processes, separately. Because several GO terms could be assigned to one single gene, the most represented GO terms were calculated over 100 %.



**Figure S2.** The *SOD2* and *SOD3* gene expression in the *sfp1Δ/sfp1Δ* mutant. The expression levels of *SOD2* and *SOD3* were detected by RT-PCR. The *PMA1* transcript was used as a loading control. WT: the wild-type strain.



**Figure S3.** Hog1 phosphorylation by Western blot with a longer exposure. After cell treatment with 10 mM H<sub>2</sub>O<sub>2</sub> for 0, 15, 30 min, Hog1 phosphorylation was assayed by Western blotting. Act1 was used as a loading control. Anti-phospho-p38 (Thr180/Tyr182) antibody (Cell Signaling, Inc.) was used to detect phosphorylated Hog1. Rabbit polyclone anti-β-actin antibody (GeneTex, Inc.) was used to detect Act1. The blot was exposure for a longer time compared to that shown in Figure 7B.