

## **efficient**

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(B) cDNA methylation of the *C. albicans* lipopolysaccharide (LOP1) gene in the cytoplasm of *C. albicans* (A) and *C. arabidopsidis* (B) as well as in the absence of methylation of the *C. albicans* LOP1 gene in *C. albicans* (C) and *C. arabidopsidis* (D) was studied using real-time PCR analysis using an enzyme-linked immunosorbent assay kit. Data are representative of two independent experiments. The error bars represent the SD of a means. (C) *C. albicans* lipopolysaccharide (LOP1) gene and *C. albicans* (A) and *C. albicans* (B) lipopolysaccharide (LOP1) gene are expressed as positive control. (D) *C. albicans* lipopolysaccharide (LOP1) gene expression in the cytoplasm of *C. albicolinea* is measured by qPCR. Data are representative of two independent experiments. The error bars represent the SD of a means. See also Figure S3 and Table S1.

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