

## Lack of Catheter Infection by the *efg1/efg1 cph1/cph1* Double-Null Mutant, a *Candida albicans* Strain That Is Defective in Filamentous Growth

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**The molecular controls regulating the successful colonization of *Candida albicans* on foreign materials are not known. Here we show that a mutant *C. albicans* strain defective in filamentous growth and lacking the transcription factors Efg1p and Cph1p has a profoundly deficient potential for colonizing on polyurethane catheters.**

The central venous catheter is known to be a major reservoir and source for *Candida* bloodstream infections in hospitalized patients (5). In addition, the ability of *Candida* species to adhere to foreign materials such as catheters has been well demonstrated (7). However, the molecular controls and environmental cues that regulate successful colonization of *Candida* spp. on such materials are not fully known. A number of pathogenicity and virulence factors that may contribute to successful colonization of *Candida albicans* have been proposed; of these, the greatest interest has been paid to the ability of *Candida* spp. to switch from yeast to filamentous growth. Dimorphism in *C. albicans* is regulated by at least two pathways, a conserved mitogen-activated protein kinase pathway that modulates the transcription factor Cph1p and a cyclic AMP-protein kinase A pathway that regulates the transcription factor Efg1p (2). A mutant *C. albicans* strain lacking both the transcription factors Efg1p (9) and Cph1p (3) is profoundly defective in filamentous growth (4), even though an Efg1p- and Cph1p-independent pathway for filamentous growth exists (6). We examined the abilities of the *efg1/efg1 cph1/cph1* double-null mutant and the *efg1/efg1 cph1/cph1* double mutant with the wild-type *EFG1* gene in the genome, as well as those of the *cph1/cph1* and *efg1/efg1* single mutants and isogenic wild-type *C. albicans* strains, to colonize on foreign materials (polyurethane catheters).

The strains used in this study and their genotypes were as follows: CAI4, *ura3::1 imm434/ura3::1 imm434*; JKC18, *ura3::1 im434/ura3::1 imm434 cph1::hisG/cph1::hisG-URA3-hisG*; HLC67, *ura3::1 imm434/ura3::1 imm434 efg1::hisG/efg1::hisG-URA3-hisG*; HLC69, *ura3::1 imm434/ura3::1 imm434 cph1::hisG/cph1::hisG efg1::hisG/efg1::hisG-URA3-hisG*; and HLC84, *ura3::1 imm434/ura3::1 imm434 cph1::hisG/cph1::hisG efg1::hisG/efg1::hisG (EFG1-URA3)* (see reference 1 for strain CAI4, reference 3 for strain JKC18, and reference 4 for the remaining

strains). Human plasma (10 ml) was injected through the lumen ports of single-lumen 7-French polyurethane catheters (Cook Catheters, Bloomington, Ind.), which were then incubated at 35°C for 24 h. Next, the catheters were removed from the plasma and placed in a culture medium (RPMI 1640 plus 0.165 M MOPS [morpholinepropanesulfonic acid]; pH 7.0) containing standardized suspensions (10<sup>5</sup> CFU/ml) of the *C. albicans* strains tested that were prepared from a 24-h culture plate. After incubation (35°C) in this medium for an additional 24 h, the catheters were carefully removed and flushed via the lumen with 100 ml of sterile saline. The catheters were then placed in a separate sterile culture tube containing 10 ml of sterile saline and gently vortexed to remove nonadherent organisms from outside the catheter lumen. The catheters were subsequently removed from the saline, aseptically cut at the 6-cm mark (on the catheter lumen), and placed in 10 ml of fresh sterile culture medium. The culture tubes were then sonicated (55,000 Hz) for 5 min to remove adherent organisms (8). After an additional 15 s of vortexing, a 50- $\mu$ l sample of the culture medium was removed to a potato dextrose agar (Remel, Lenexa, Kans.) plate using a spiral-gradient plating system (Spiral Biotech Inc., Bethesda, Md.). The numbers of viable CFU per milliliter of medium were determined according to the colony counts after 24- and 48-h incubations at 35°C. To further characterize the phenotype of catheter attachments by the wild-type and mutant *C. albicans* strains, experiments were repeated as described above except that the 6-cm catheter segment was placed in 4% glutaraldehyde in a 0.1 M sodium cacodylate buffer, pH 7.2. Catheter sections (2 cm) were then fixed in osmium tetroxide and dehydrated using a graded series of ethanol washings followed by immersion in hexamethyldisilazane. Next, the sections were air dried, mounted onto aluminum stubs, and sputter-coated with Pd and Au. A Hitachi S4000 scanning electron microscope (Hitachi Scientific Instruments, Mountain View, Calif.) was used for visualization of the samples. All *C. albicans* strains were tested simultaneously, and experiments were performed in triplicate. Mean data (CFU per milliliter) were compared using analysis of variance along with the Tukey's test for post hoc comparisons.

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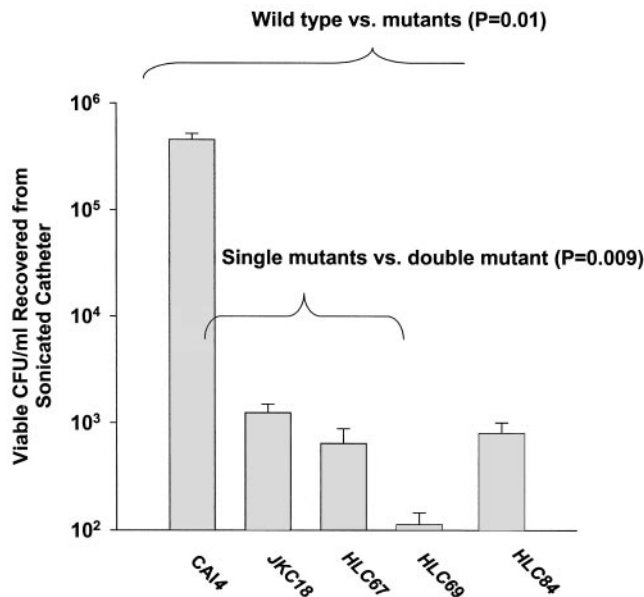


FIG. 1. Comparison of the colonization intensities (polyurethane central venous catheters) of *C. albicans* strains with the following genotypes: isogenic wild-type (CAI4), *efg1/efg1* (HLC67) and *cph1/cph1* (JKC18) single mutants, *efg1/efg1 cph1/cph1* double mutant (HLC69), and *efg1/efg1 cph1/cph1* with the wild-type *EFG1* gene (HLC84). Counts were analyzed by nonparametric Kruskal-Wallis one-way analysis of variance on ranks by using Dunn's pairwise multiple-comparison technique (Sigmastat statistical software, version 2.0; SPSS Science, Chicago, Ill.).

The *efg1/efg1 cph1/cph1* double-mutant strain was defective in its ability to colonize on the polyurethane catheters. Specifically, the mean number of viable CFU per milliliter recovered from the sonicated catheter segments was 3 to 4 log<sub>10</sub> lower for the double-mutant strain versus the wild-type strain ( $P = 0.01$ ) (Fig. 1). The *efg1/efg1* and *cph1/cph1* single-mutant strains had an intermediate phenotype, having a mean number of viable

CFU per milliliter recovered from the segments that was about 1 log<sub>10</sub> higher than that for the double-mutant strain (Fig. 1). When catheters were examined by scanning electron microscopy, decreased colonization was noted by the *efg1/efg1 cph1/cph1* double mutant, with occasional blastoconidia but no hyphal forms (Fig. 2).

Of the two major signaling pathways controlling hyphal growth, the Efg1p pathway appears more important since *efg1/efg1* mutants demonstrate a more severe morphogenetic block under most inducing conditions (4). Furthermore, *efg1/efg1* mutants are much less virulent than the *cph1/cph1* mutants in the mouse model of candidiasis (4). Our observation that the *efg1/efg1* mutant had more-severe adherence defects than the *cph1/cph1* mutant is consistent with these prior observations. Nevertheless, the *efg1/efg1 cph1/cph1* double mutant has the most severe morphogenetic and virulence defects (4). Such morphogenetic defects were also observed in our study. The presence of the wild-type *EFG1* gene integrated in the genome of the *efg1/efg1 cph1/cph1* double mutant (HLC84) caused the formation of germ tubes and pseudohyphae (4). The ability of the *efg1/efg1 cph1/cph1* double mutant to colonize was restored to the level of the *cph1/cph1* single mutant by integrating the wild-type *EFG1* gene in the genome (Fig. 1). Therefore, our observation is consistent with the hypothesis that Cph1p and Efg1p govern catheter colonization ability.

One of the limitations of our approach is that we did not evaluate fully the possibility of another, secondary mutation giving the phenotype observed. However, we think that it is unlikely that another mutation could affect the phenotypic characteristics in two genotypically different strains such as *efg1/efg1* and *cph1/cph1*. In addition, our preliminary data do not address which of the rather-complex and interrelated early events of catheter colonization, such as adherence and biofilm formation, are regulated predominantly by *EFG1* and *CPH1*.

In conclusion, both Efg1p and Cph1p appear to be central regulators of the developmental program that allows *C. albicans* to successfully colonize on foreign materials such as poly-

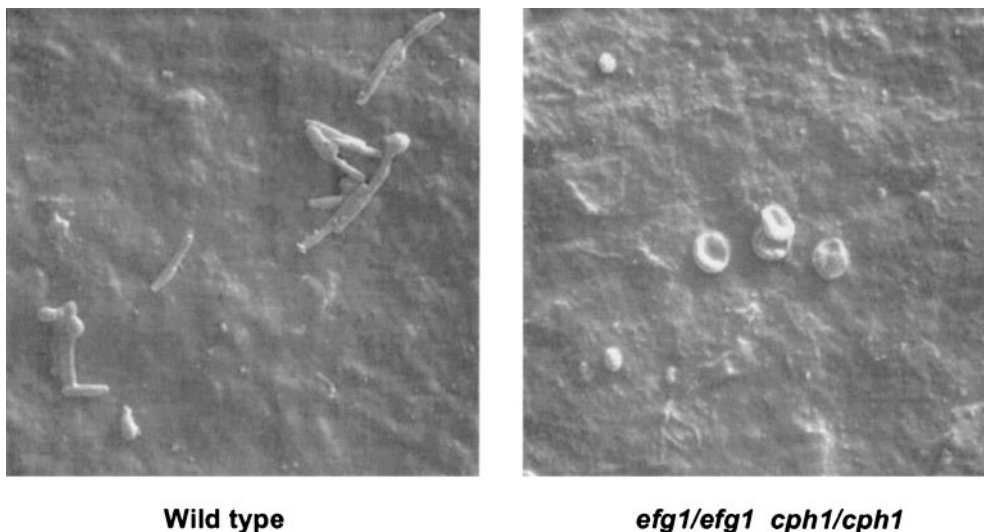


FIG. 2. Scanning electron micrographs (magnification, ×1,000) of exterior catheter lumens exposed to the wild-type (CAI4) and *efg1/efg1 cph1/cph1* double-mutant (HLC69) *C. albicans* strains.

urethane central venous catheters. It is possible that interruption of morphogenetic events induced by Efg1p and Cph1p through pharmacological or other means could affect the fitness of *C. albicans* to successfully colonize prosthetic materials or central venous catheters.

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