



Olorofim Effectively Eradicates Dermatophytes *In Vitro* and *In Vivo*

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ABSTRACT Superficial fungal infections are prevalent worldwide, with dermatophytes as the most common cause. Various antifungal agents including azoles and allylamines are commonly used to treat dermatophytosis. However, their overuse has yielded drug-resistant strains, calling for the development of novel antimycotic compounds. Olorofim is a newly developed antifungal compound that targets pyrimidine biosynthesis in molds. The purpose of this study was to determine the *in vitro* and *in vivo* antifungal effects of olorofim against common dermatophytes. The *in vitro* activity of olorofim against dermatophytes was assessed by microtiter broth dilution method. Bioinformatic analysis of olorofim binding to dihydroorotate dehydrogenase (DHODH) of dermatophytes was also performed, using *Aspergillus fumigatus* DHODH as a template. The *in vivo* efficacy of the drug was investigated, using a guinea pig model, experimentally infected with *Microsporum gypseum*. Microtiter assays confirmed the high *in vitro* sensitivity of dermatophytes to olorofim (MIC = 0.015–0.06 mg/liter). Amino acid sequence analysis indicated that DHODH is highly conserved among dermatophytes. The critical residues, in dermatophytes, involved in olorofim binding were similar to their counterparts in *A. fumigatus* DHODH, which explains their susceptibility to olorofim. Typical skin lesions of dermatophyte infection were observed in the guinea pig model at 7 days postinoculation. Following 1 week of daily topical administration of olorofim, similar to the clotrimazole group, the skin lesions were resolved and normal hair growth patterns appeared. In light of the *in vitro* and *in vivo* activity of olorofim against dermatophytes, this novel agent may be considered as a treatment of choice against dermatophytosis.

KEYWORDS *Microsporum gypseum*, antifungals, dermatophytes, dihydroorotate dehydrogenase, olorofim

According to a recent estimate of the prevalence of fungal diseases, the burden of superficial fungal infections, including skin, hair and nail, approximates 750 million people worldwide (1). Superficial mycosis is mainly attributed to two divergent fungal groups: dermatophytes, as the most common causative agents, including the genera *Trichophyton*, *Microsporum*, and *Epidermophyton*; and non-dermatophytes yeasts from the genera *Candida*, *Trichosporon*, and *Malassezia* (2). In 2017, an estimated five million people in the US were diagnosed with dermatophytes, imposing a direct cost of ~\$821 million to the US health care system (<https://www.cdc.gov/fungal/cdc-and-fungal/burden.html>). Dermatophytes spread *via* direct contact with infected people, animals, or soil, as well as indirectly from fomites. These pathogens can colonize the keratinized structures present in skin, hair, and nails, causing superficial infections, known as dermatophytosis (3). Dermatophytoses are mainly treated by local application and/or systemic administration of azole-based drugs including clotrimazole,

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TABLE 1 *In vitro* susceptibility of aspergilli and dermatophytes to olorofim, posaconazole, voriconazole, and clotrimazole

Antifungal compound	MIC (mg/L)							
	<i>Aspergillus</i>		Dermatophytes					
	<i>Aspergillus fumigatus</i> PTCC5009	<i>Aspergillus flavus</i> PTCC5004	<i>Trichophyton mentagrophytes</i> NBRC5809	<i>Trichophyton tonsurans</i> CBS 130814	<i>Trichophyton rubrum</i> IR613	<i>Epidermophyton floccosum</i> CBS 130793	<i>Microsporum canis</i> PTCC5069	<i>Microsporum gypseum</i> PTCC5070
Olorofim	0.01	0.01	0.01	0.06	0.01	0.03	0.03	0.03
Posaconazole	0.15	0.3	0.04	0.6	0.08	0.12	0.3	0.6
Voriconazole	0.15	0.15	0.15	0.15	0.15	0.12	0.6	0.6
Clotrimazole	2	4	0.25	1	16	2	4	1

econazole, ketoconazole, miconazole, tioconazole, as well as terbinafine, amorolfine, tolnaftate, and griseofulvin (4, 5).

Antifungal susceptibility testing is not routinely undertaken in cases of dermatophytosis, as the infecting organism is rarely identified. Resistance to commonly administered topical azoles has been increasingly reported (6). Moreover, a number of reports indicate that the terbinafine-resistant *T. mentagrophytes* are emerging and spreading globally (7–11). Although dermatophytosis is non-fatal, it can be disfiguring and contagious, requiring immediate treatment. Long-term treatment is often required, and compliance can be poor, and this can bring about the emergence of drug resistance strains (12, 13).

This fact emphasizes the urgent need for introducing new classes of antifungals, with novel mechanisms of action. A recently developed antifungal compound, olorofim (F2G Ltd, UK), has demonstrated high efficiency against *Aspergillus* species and some other molds (14, 15). Belonging to a new class of antifungals titled the orotomides, it specifically targets fungal dihydroorotate dehydrogenase (DHODH) (14), an essential enzyme in the *de novo* pyrimidine biosynthesis pathway (16). Olorofim does not inhibit human DHODH and is currently in a phase IIb open-label study, focusing on rare and resistant, life-threatening, invasive fungal infections (www.F2g.com).

Considering the strong antifungal activity of olorofim against different pathogenic molds (9, 14, 17), here we have investigated the *in vitro* and *in vivo* effects of this compound against dermatophytes, using an animal model of dermatophytosis.

RESULTS

***In vitro* sensitivity of dermatophytes to olorofim.** MICs of olorofim were determined for different dermatophyte and *Aspergillus* strains, in comparison to posaconazole and voriconazole. As demonstrated by the MICs listed in Table 1, both groups (dermatophyte and *Aspergillus* strains) were far more sensitive to olorofim, compared to posaconazole and voriconazole. *A. fumigatus* and *A. flavus* showed identical MIC value of 0.01 mg/liter; however, the susceptibility range for the dermatophyte isolates was 0.01–0.06 mg/liter. The highest MIC was observed in *T. tonsurans*.

DHODH is highly conserved among dermatophytes. The above MIC results led us to analyze the DHODH sequences of the dermatophytes tested, except *Epidermophyton floccosum*, for which no genome was available, and compare them to other fungal and human DHODH. The amino acid sequences revealed a significant conservation, among the aligned sequences (Fig. 1). The DHODH sequence similarity and identity among different strains of dermatophytes ranged from 74.3 to 95.5% and 73 to 90%, respectively. The overall similarity between DHODH of the investigated dermatophytes and *A. fumigatus* was about 63%. In previous reports, the key residues for olorofim binding were identified as His₁₁₆, Val₂₀₀, Arg₂₀₂, Met₂₀₉, Tyr₂₁₃, Tyr₅₀₇, and Tyr₅₁₂ (14, 22). Our results indicated that dermatophytes share six out of these seven critical residues with *A. fumigatus*. The only differing residue was Met₂₀₉, which was replaced by Leu in *Trichophyton* and *Microsporum*, and Val in *Candida* strains.

***In vivo* sensitivity of dermatophytes to olorofim.** All infected guinea pigs showed multiple signs of superficial fungal infection, including erythema, ulceration, mild shedding,

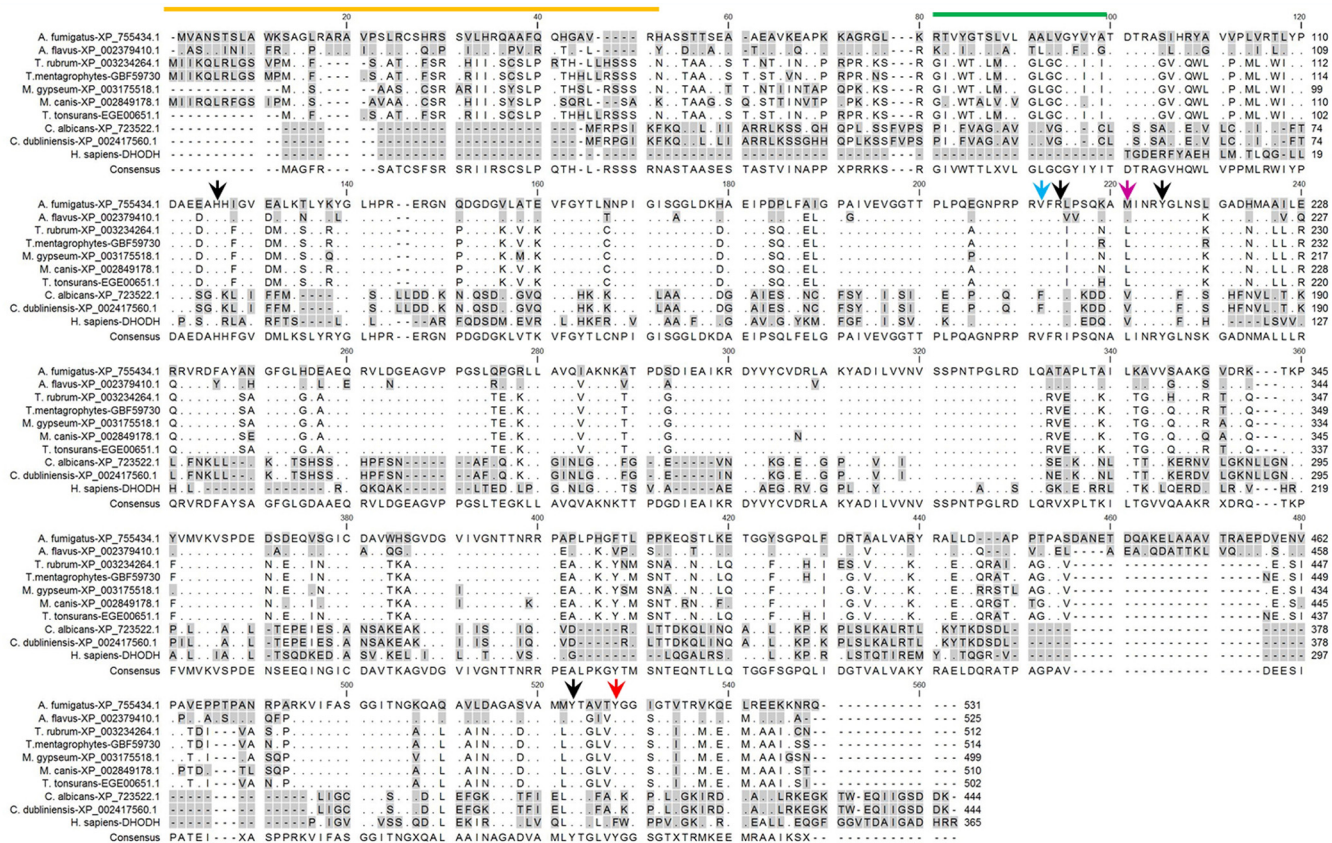


FIG 1 Alignment of different dermatophytes, *A. fumigatus*, *C. albicans*, and human (*H. sapiens*) DHODH amino acid sequences. Identical residues are distinguished by dots, and similar residues are highlighted in gray. The predicted mitochondrial targeting sequences are indicated by the orange line, and the predicted transmembrane domains are shown by the green line. The seven residues predicted to be important for orlofim binding in *A. fumigatus* DHODH are depicted by arrows, whose colors illustrate conservation status—black, purple, and blue for conserved residues unique to *A. fumigatus*, *Candida*, and humans, respectively.

and scaly skin, in the infected area, 7 days postinfection (Fig. 2). Fungal infection was validated by the presence of fungal elements in microscopic examination of scrapings (data not shown).

Following topical antifungal treatment, the skin samples were collected by scraping the edge of the healed lesions and examined by direct microscopy. In the control group (c: PEG-treated), the patches of hair loss and the readily visible ulcerated or scaly skin were still present, 1 week after PEG administration (Fig. 2 cl & cil). In this group, the lesions self-healed after 3–4 weeks of infection. However, in the orlofim-treated group (Group a), similar to the clotrimazole-treated group (Group b), no fungal elements were observed, at 7 days post treatment. Furthermore, orlofim treatment significantly reduced the redness and the skin lesions, followed by the appearance of normal pattern of hair growth and no sign of scaly skin after 1 week (Fig. 2 all and bil).

DISCUSSION

Superficial mycosis, the most prevalent fungal infection, affects 20–25% of the world population (23, 24). Dermatophytes are known as the main causative agents of the disease and are often treated by antifungal drugs, targeting the ergosterol biosynthetic pathway, including azoles and allylamines. Unfortunately, the widespread and unattended use of over-the-counter azoles, especially in combination with topical steroids, has led to the development of drug resistance in dermatophytes (5, 6). Accordingly, reports indicate an estimated 19% azole resistance rate among dermatophytes, particularly in certain tropical areas (6). In this sense, novel antifungal compounds with new targets are under investigation. Although the majority of antifungal drug discovery research is driven by the need for better treatment of invasive and systemic fungal diseases, novel compounds may be applicable in treatment of superficial mycosis. Olorofim, a highly potent and selective fungal DHODH

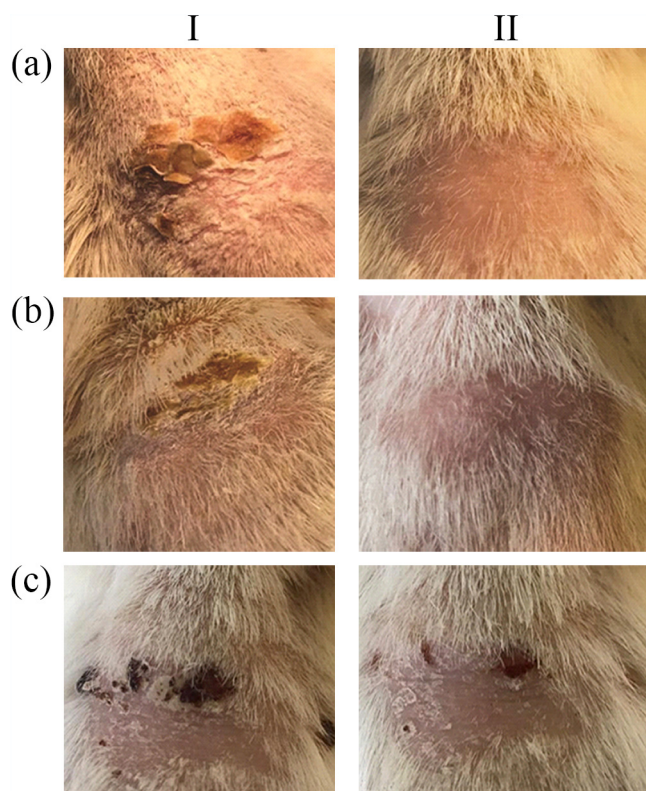


FIG 2 The effect of different treatments on skin lesions in guinea pigs Infected with *M. gypseum*. aI, before treatment with orlofim; all, After 1 week treatment with orlofim (10 $\mu\text{g}/\text{lesion}$); bI, before treatment by clotrimazole; bII, after 1 week treatment with clotrimazole (1%); cI, before treatment by PEG 300; cII, after 1 week treatment with PEG 300.

inhibitor, is active against clinically relevant molds, except *Mucorales*. Due to its novel mechanism of action, it is unlikely for orlofim to cause or be affected by cross-resistance to other antifungal classes (14, 17). The orlofim-susceptible fungal species are closely related in their DHODH amino acid sequences, compared to non-susceptible fungi, such as *Candida* spp, *Mucorales* spp, and *C. neoformans*.

Here, we have investigated the antifungal activity of orlofim against three different genera of dermatophytes, *Trichophyton*, *Microsporum*, and *Epidermophyton*. Our results confirmed the high sensitivity of these organisms to the drug, which is in agreement with previous reports (14, 25). The MIC levels (0.015 to 0.062 $\mu\text{g}/\text{ml}$) were in a range similar to those reported for *Trichophyton* and *Microsporum* by Astvad et al. (25). In our study, the variations observed in MIC levels may re-emphasize the species-specific differences among dermatophytes (8). The small number of fungal strains used for MIC determinations can be considered as a limitation of the present investigation. The high susceptibility of dermatophytes to orlofim can be attributed to the significant sequence similarity (~63%) of their DHODH amino acid sequences with that of *Aspergillus* spp. Based on the report by Oliver et al. ⁶, Val₂₀₀ and Met₂₀₉ are identified as key residues in *A. fumigatus* sensitivity to orlofim, such that mutation of these two amino acids occupying the same position in *Candida albicans* DHODH (Phe₁₆₂ and Val₁₇₁) to Val₁₆₂ and Met₁₇₁ resulted in a recombinant DHODH that was susceptible to orlofim inhibition in *in vitro* assays, while the original *C. albicans* DHODH was not. In the case of Met₂₀₉ in the *A. fumigatus* sequence, it seems that a conservative replacement of Leu in the dermatophytes studied is tolerated by orlofim, judging by the low MICs observed in these organisms. The remaining six residues, expected to influence orlofim binding of DHODH, are identically conserved between *A. fumigatus* and the dermatophytes, consistent with the low MIC values observed.

To investigate the *in vivo* efficacy of orlofim against dermatophytes, a guinea pig model of dermatophytosis was created. Guinea pigs have been widely used as a model

for establishment of dermatophytosis as the clinical signs of dermal infection in this animal are comparable to those seen in humans. The guinea pig model has also been used in predicting the efficacy of antifungal preparations, which makes it a valuable tool for the preclinical assessment of new antifungal compounds (26). We used an animal isolate of *M. gypseum* for infection in this model. Experimental infection of guinea pigs with *M. gypseum* has been previously described (21, 27, 28). Here, we first made several attempts to infect animals via skin abrasion, however with limited success. Hence, we decided to use corticosteroids to temporally suppress the immune system of the animals and to enhance the chance of infection. The administration of corticosteroids for successful establishment of dermatophytosis has also been previously reported (29). Despite the similarly used method for fungal inoculation, some variation in the clinical signs of infection was observed between the animals. This can be attributed to host factors, such as the varying strength of the immune system, in outbred animals (27, 30). However, clear differences in clinical outcomes in treated animals versus controls were observed (e.g., compare olorofim-treated, Fig. 2 all; versus control-treated, c; and immediately prior to olorofim treatment, a).

In most studies, the topical or oral treatment of dermatophytosis was undertaken 3–5 days following the infection (31). However, in our study, olorofim (0.1 mg/ml in PEG300) was topically administered daily at a dosage of 10 $\mu\text{g}/\text{lesion}$ from the eighth day, postinfection. This starting time of treatment was chosen based on the time at which the skin lesions were clearly visible (i.e., 7 days postinfection). Topical treatment of olorofim was carried out every day for 7 days, at which time the results were compared with the positive and negative controls. The selection of such a short time period was to avoid the self-healing time course of cutaneous dermatophytosis in guinea pigs (32). The drug cured the skin lesions during the first week of treatment, and the skin looked healthy and smooth with no redness, swelling, or scarring. These results indicated that olorofim was highly effective in the treatment of dermatophytosis in the guinea pig model. This may be due to the very low MICs observed for this compound against dermatophytes. We also found that the olorofim dosage of 10 $\mu\text{g}/\text{lesion}$ (2.5 $\mu\text{g}/\text{cm}^2$) mimics the therapeutic dose of clotrimazole, although other concentrations remain to be examined. The data for clotrimazole indicate that following the application of topical clotrimazole 1% cream, the concentration of clotrimazole would be around 100 $\mu\text{g}/\text{cm}^2$ in the stratum corneum, which is much higher than that used for olorofim in our study (33).

In conclusion, we demonstrated the efficacy of olorofim as a novel anti-dermatophytosis agent against various dermatophyte species, *in vitro* and also against *M. gypseum* infection *in vivo*. However, more detailed studies are required to elucidate its efficacy against other dermatophytes, as well as clarifying the drug pharmacokinetics, upon topical administration.

MATERIALS AND METHODS

Ethics. This study was conducted in accordance with institutional standards and approved by the Ethical Committee of the Pasteur Institute of Iran. (Ethical code: IR.PII.REC.1397.021).

Strains, culture conditions, and antifungal agents. Dermatophyte strains ($n = 6$, Table 1) were grown on Sabouraud dextrose agar (SDA, Merck, Germany) slants, supplemented with chloramphenicol (0.005%) and cycloheximide (0.04%), at 28°C for 10–14 days. *Aspergillus* strains ($n = 2$) were cultivated on SDA plates, at 37°C, for 3–5 days. To collect fresh spores, fungal colonies were gently washed with PBS-Tween 80, and the resulting suspension was filtered through a thin layer of sterilized glass wool to remove the hyphal fragments. The spores were then separated by centrifugation at 1000 $\times g$, for 10 min, and their concentrations were determined using hemocytometer counts. The final concentrations of spores were adjusted at 10⁶/ml to use in MIC assays.

Azole compounds including voriconazole, posaconazole, and clotrimazole were purchased from Sigma-Aldrich, UK. Olorofim was kindly provided by F2g Limited, UK. All compounds were prepared and stored as 5 mg/ml stocks in DMSO at 4°C. Clotrimazole topical cream was provided by Emad Darman Pars pharmaceutical company, Iran.

Antifungal susceptibility testing. The antifungal susceptibility of dermatophytes was assessed by determining the MIC, based on the M38-E3rd protocol of the Clinical and Laboratory Standards Institute (CLSI), with some modifications (18). Briefly, a total of 10⁴ spores were suspended in 80 μl of RPMI 1640 medium, buffered to pH 7.0 (with MOPS) and seeded onto a 96-well microtiter plate. Then 20 μl of serial 2-fold dilutions of each test compound was added to each well and the plate was incubated at 28°C. The MICs were assessed after 96 h of incubation, at a final compound concentration range of 0.001–10 $\mu\text{g}/\text{ml}$ for voriconazole and posaconazole, 0.03–64 $\mu\text{g}/\text{ml}$ for clotrimazole, and 0.0001–1 $\mu\text{g}/\text{ml}$ for olorofim, MIC endpoints were defined as the lowest

concentration of each test compound that resulted in inhibition of growth (80% or more) by visual inspection, compared to the controls.

Sequence analysis. In order to identify DHODH homologues in different fungi, the amino acid sequence of *A. fumigatus* DHODH was retrieved from the KEGG database (E.C.1.3.5.2, AFUA_2G11010, XP_755434) and used as a template in a tBLASTn search against available dermatophytes, *Candida*, and human genomes. The sequences were aligned and formatted by CLC Main Workbench (<https://digitalinsights.qiagen.com>). Mitochondrial targeting sequences and transmembrane domains of the enzyme were predicted by MitoFates and Phobius servers, respectively (19, 20).

Animal model of infection. Nine albino female guinea pigs (300–350 g each) were purchased from the Laboratory of Animal Sciences, Pasteur Institute of Iran. Animals were housed in groups of three, kept under standard laboratory conditions (room temperature of 18–22°C, relative humidity of 40–50%, and 12h light/day cycle) and provided with food and water *ad libitum*. The guinea pigs were acclimated for 1 week prior to experimental treatments. To establish dermatophytosis, animals were immunosuppressed via intramuscular injection of prednisolone (10 mg/kg) and subcutaneous injection of hydrocortisone (5 mg/kg), 1 day prior to infection and 3 days thereafter. All guinea pigs were anesthetized, using intraperitoneal injection of ketamine (50 mg/kg), xylazine (5 mg/kg), and acepromazine (1 mg/kg). Once the animals were fully anesthetized, the posterior dorsal areas were gently shaved (~ 2 × 2 cm) and then abraded with the back of a sterile scalpel blade. The inoculation of fungi was carried out according to previous protocols, with some modifications (21). Briefly, an animal isolate of *Microsporium gypseum* was grown on SAB agar for 14 days and checked for the presence of micro- and macroconidia. One hundred microliters of this inoculum (10⁸ conidia in PBS/Tween 0.01%) were spread on the abraded area using a sterile pipette tip and left to dry. The inoculated area was surrounded by a thin layer of Vaseline and then dressed with sterile pads and bandaged with non-woven tape (TGMED), for dressing fixation. Each animal was placed on a hot water blanket until full recovery from anesthesia was achieved. For mycological evaluations, surface scrapings were collected from the inoculation sites at 7 days postinfection, and the obtained specimens underwent direct microscopic examination using routine KOH wet mount.

Eight days postinfection, the animals were randomly divided into 3 groups (a–c) and received topical treatments of the following compounds, once daily for 7 days: Group a, Olorofim (100 μl of 0.1 mg/ml in PEG300); Group b, 1% clotrimazole (as positive control, topical cream); and Group c, PEG 300 (100 μl, as negative control).

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We declare no conflict of interest.

E.M.A. designed the animal model study, performed all the related experiments, and participated in preparation of the first draft of the manuscript. A.S. performed the bioinformatic analyses and assisted in preparation of the first draft of the manuscript. N.P. performed MIC assays. M.N. and M.Z. helped in preparation of fungal cultures and spore inoculations. S.E. assisted in animal study. M.R.-A. was involved in interpretation of MICs. V.K. designed and supervised the whole study and critically revised and finalized the manuscript.

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