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Two promising natural lipopeptides from *Bacillus subtilis* effectively induced membrane permeabilization in *Candida glabrata*

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Candida glabrata is an important opportunistic human pathogen well known to develop resistance to antifungal drugs. Due to their numerous desirable qualities, antimicrobial lipopeptides have gained significant attention as promising candidates for antifungal drugs. In the present study, two bioactive lipopeptides (AF₄ and AF₅ m/z 1071.5 and 1085.5, respectively), coproduced and purified from *Bacillus subtilis* RLID12.1, consist of seven amino acid residues with lipid moieties. In our previous studies, the reversed phased-HPLC purified lipopeptides demonstrated broad-spectrum of antifungal activities against over 110 *Candida albicans, Candida* non-*albicans* and mycelial fungi. Two lipopeptides triggered membrane permeabilization of *C. glabrata* cells, as confirmed by propidium iodide-based flow cytometry, with PI uptake up to 99% demonstrating fungicidal effects. Metabolic inactivation in treated cells was confirmed by FUN-1-based confocal microscopy. Together, the results indicate that these lipopeptides have potentials to be developed into a new set of antifungals for combating fungal infections.

KEYWORDS

antifungal, bioactive, *Bacillus subtilis*, *Candida glabrata*, lipopeptides, membrane permeabilization

Introduction

After *Candida albicans*, the most prevalent yeast species reported to have been isolated from humans is *Candida glabrata* (*C. glabrata*). The past two and a half decades have witnessed *C. glabrata* becoming one of the most predominant *Candida* species, causing fungal infections (Chakrabarti et al., 2009; Lockhart et al., 2012; Benaducci et al., 2014; Cleveland et al., 2015; Pappas et al., 2018; Naicker et., 2023). The reason behind this steep upward trend of *C. glabrata* infections may be associated with geographical variation and the rampant overuse of azoles (Richardson and Lass-Flörl, 2008). Given the prevalence of *C. glabrata* as the second leading cause of candidiasis infections,

including oral candidiasis, candidemia, invasive candidiasis, and urinary candidiasis (Esfandiary et al., 2012; Nash et al., 2016), it is essential to understand the alarming 40% mortality rate associated with C. glabrata infection underscores the seriousness of this opportunistic human pathogen, especially in immunocompromised hosts (Rodrigues et al., 2014; Mota et al., 2015; Nagayoshi et al., 2017). It has been recently reported that C. glabrata has developed several strategies to survive and thrive in its host cells and found ways to overcome antifungal resistance to commonly used antifungals, which also contribute to increased virulence (Pappas et al., 2018; Salazar et al., 2018). Flow cytometry (FCM) has been one of the major techniques that has successfully demonstrated its utility in determining the effect of antimycotics on yeast cells, cell membranes, or mechanisms of action when used with appropriate dye or fluorescent probes and experimental conditions (Pore, 1994). In our previous study (Ramachandran et al., 2018b), the antifungal susceptibility test (AFST) analyses of the two lipopeptides were done according to the CLSI guidelines. In this current investigation, the two lipopeptide fractions (AF₄ and AF₅) coproduced by Bacillus subtilis RLID 12.1 were purified, and the main objective of the study was to determine their membrane-permeabilizing effects on the yeast cell membrane and metabolic activity at two different peptide concentrations using the membraneimpermeant propidium iodide (PI) dye and two-colour FUN-1 stain.

Materials and methods

Microorganisms

The test strain *C. glabrata* ATCC 2001 was obtained from the National Culture Collection of Pathogenic Fungi (NCCPF), Post Graduate Institute of Medical Education and Research (PGIMER) Chandigarh, India and maintained as 20% glycerol stock at -80° C. The producer strain was also maintained as glycerol stock at -80° C.

Extraction and purification of antifungal compounds

For the purification of antifungal lipopeptides, cell-free supernatant (CFS) was obtained after 60 h at 30°C of incubation under stirring conditions of cells of the strain *B. subtilis* RLID 12.1. The CFS was subjected to a three-step purification that included HCl precipitation, solvent extraction using n-butanol, and silica gel (mesh size 230–400) based adsorption chromatography using proper ratios of methanol and chloroform. After the adsorption chromatography, all the collected fractions were tested for antifungal activity by spot-on-lawn assay against freshly grown *C. glabrata* cells spreading on Sabourad dextrose agar (SDA) plates. The 5 μ L

aliquots of fractions which showed the clear zones of inhibition after 24 h of incubation were considered as bioactive fractions. These bioactive fractions AF₄ and AF₅ were fractionated using the reversed phased-HPLC system (Agilent Technologies, United States) at a semi-preparative scale equipped with a variable wavelength detector and an Agilent C18 column (10 mm × 250 mm, 5 μ m) (Ramachandran et al., 2018a; Ramachandran et al., 2018b). The analytical scale HPLC profiles of AF₄ and AF₅ and their m/z values are shown in Supplementary Figure S1.

Antifungal susceptibility testing (AFST) against *Candida glabrata*

The minimum inhibitory concentrations (MICs) of the purified antifungal lipopeptides AF_4 and AF_5 were tested against *C. glabrata* ATCC 2001, according to CLSI guidelines (M27-A2) (Clinical and Laboratory Standards Institute, 2008). For antifungal assays, two-fold serial dilutions of each drug were prepared in RPMI-1640 (pH 7.0 \pm 0.1) (with L-glutamine and phenol red without sodium bicarbonate) was buffered with 0.165 M MOPS (morpholinepropanesulfonic acid) (Himedia) and supplemented with 0.2% glucose. As reference yeast strain for the AFST, *C. albicans* ATCC 24433 and the positive control amphotericin B (AMB) (HiMedia, India) were used.

PI uptake assay by flow cytometry (FCM)

To study the impact of novel antifungal lipopeptides on membrane integrity, lipopeptides AF_4 and AF_5 (8 and 16 mg/L), and AMB (×1 and ×2 MIC) were added to *Candida* cell suspension (1 × 10⁶ CFU/mL) in RPMI 1640 and incubated at 37°C for 18 and 5 h, respectively, in shaking conditions (Gokahmetoglu et al., 2003; Seyedjavadi et al., 2020). Two types of controls were used in the study: negative controls consisting of untreated cells, and PI positive controls, treated with 70% ethanol for 30 min. Post-incubation, the cells were harvested, washed with 1× phosphate buffered saline (PBS), and stained with PI at a concentration of 7.5 µg/mL for 20 min. Unstained and untreated cells were sampled and analysed in the beginning. Flow cytometry analysis was performed on 30,000 events using a FACSMelody flow cytometer with a 488 nm laser line and a 586 nm filter with PI detection.

Viability (CFU) assays

Prior PI staining, aliquots of yeast cultures (both untreated and treated) were collected, serially diluted in sterile 1×PBS, and plated in duplicate on SDA plates (Chaturvedi et al., 2004; Ramesh et al., 2023). The plates were incubated for 24 h at 37°C, colonies were



FIGURE 1

The effect of the lipopeptide AF_4 on the cell membrane permeability of *C. glabrata* 2001, measured by PI uptake. Experiments recorded forward scatter, side scatter, and the percentage of PI-stained cells, and the data was analyzed using the software FlowJo version 10.8.1. The histograms related to PI fluorescence intensity were split into two quadrants, i.e., one with PI negative cells (viable cells), and another with PI-positive cells (membrane compromised cells). (A) Flow cytometry analysis of membrane permeabilization assay by PI uptake. (B) The side scatter versus PI plot shows the shift of population of PI-positive cells along the x-axis. (C,D) Confocal images of *C. glabrata* cell membrane integrity treated with AF_4 , AF_5 and AMB monitored by PI uptake (Left panel, magnification 60 x 2 and, right panel 100 x 2, respectively). A few cells in the field as indicated by white arrows are non-viable.

counted, and findings were represented as the percentage reduction in CFU/mL compared to the growth in the untreated sample.

Confocal laser scanning microscopy (CLSM)

The CLSM analysis was performed using PI (ThermoFisher, United States) and [2-chloro-4-(2, 3-dihydro-3-methyl-(benzol-

1, 3-thiazol-2-yl)-methylidene)-1-phenylquinolininum iodide] FUN-1 (ThermoFisher, United States) (Chan et al., 2011; Zhang et al., 2018). Cells were treated with AF_4/AF_5 (8 and 16 mg/L) for 18 h, AMB (1 mg/L) for 5 h, and untreated cells were used as control. Post-treatments, cells were harvested and resuspended in 1×PBS, following which staining with PI (7.0 µg/ mL) for 20 min in the dark and washing with 1×PBS were done. FUN-1 at a final concentration of 5 µM was prepared in 10 mM glucose-HEPES (GH) buffer, and cells were incubated in (GH)



FIGURE 2

The effect of the lipopeptide AF_5 on the cell membrane permeability of *C. glabrata* 2001, determined by Pl uptake. (A) Flow cytometry analysis of membrane permeabilization assay. Cells were treated with AF_5 (8) and (16) mg/L compared with the control (untreated). (B) The side scatter versus Pl plot shows the shift of population of Pl-positive cells along the x-axis. Density plot of Pl uptake by *C. glabrata* cell populations of a) unstained, b) control, and c) and d) AF_5 (8) and (16) mg/L treated cells, respectively, along with e) 70% ethanol. Ethanol-treated cells exhibited a high permeability to Pl (99.99% cells stained). (C) Confocal image (×200 magnification) showing cells stained with FUN-1, the AF_4 , AF_5 and AMB treated cells are without discernible and red fluorescent CIVS and show yellow-green fluorescence in the merged channel, representing metabolically inactive cells compared to untreated cells. The inset images show non-viable cells from another field.

buffer with FUN-1 stain for 30 min (Pina-Vaz et al., 2001; Kwolek-Mirek and Zadrag-Tecza, 2014; Yan et al., 2019). Treated and untreated cells were imaged with an Olympus FV3000 (Japan) microscope at 60 \times 2 and 100 \times 2 magnifications for PI-stained cells and 100 \times 2 for FUN-1 stained cells.

Statistical analysis

Experiments were performed in three individual times with two technical replicates. Data are presented as the means \pm SD values and, One-way ANOVA statistical analysis was done in the Graph-pad prism Software version 9.3.1.

Results and discussion

The MICs of two lipopeptides and AMB (used as the standard antifungal) against C. glabrata were determined to be 4 mg/L and 1 mg/L, respectively. The results agree with the previously determined MIC values (Ramachandran et al., 2018b; Ramchandran et al., 2020). To gain more insights into lipopeptides' membrane permeabilizing potential, we performed PI uptake assays using 8 mg/L (×2) and 16 mg/L $(\times 4)$ MICs as determined by the AFST for C. glabrata ATCC 2001. FC results were expressed as the fluorescence intensity (FI) of PI-stained yeasts. AF₄ and AF₅ treatments caused cell membrane disruption in C. glabrata cells, as seen by the enhanced fluorescence that resulted from PI uptake. In cells treated with AMB, AF₄, AF₅, and 70% ethanol, the increase in PI-fluorescence is manifested as a distinct shift of the peak along the x-axis (Figures 1A, B) and (Figures 2A, B). The histograms pertaining to all flow cytometry assays where C. glabrata ATCC 2001 cells were stained with PI after being exposed to AMB for 5 h (Gokahmetoglu et al., 2003) at 1 and 2 mg/L and AF_4/AF_5 (8 and 16 mg/L) for 18 h have been shown in Figures 1A, 2A. All the events of FC population density plots have been shown in Figures 1B, 2B.

Significant congruence was found when comparing the proportion of PI uptake with the drop in CFU/mL among AF4 treatments. A substantial association was observed between the percentage PI uptake estimated in flow cytometry and the percentage reduction in plate counts, presented in tabular form (Table 1) to summarize the findings. Cells unexposed to lipopeptides showed very negligible fluorescence, as evidenced by the PI uptake percentage graph (Figure 3). At 8 mg/L, the AF₄-treated C. glabrata showed an average of 81.92% PI-positive cells and 99.90% CFU reduction, better was the observation in AF₄-(16 mg/L)-treated C. glabrata cells, where 99.51% PI-positive cells were noted and a CFU reduction of 99.97% recorded (Figure 3; Table 1). At 8 mg/L, the AF5-treated C. glabrata showed an average of 92.14% PIpositive cells and 99.4% CFU reduction. However, AF5-(16 mg/L)-treated C. glabrata cells showed 81.35% PIpositive cells, and the CFU reduction of 99.92% was recorded, indicating that AF₅ even at a 2-fold concentration, did not show more membrane permeabilization and increased fungicidal effects. This may be due to the hydrophobic aggregation of AF₅ lipopeptide at higher concentrations (Ramchandran et al., 2020). In AMB (1 and 2 mg/L)-treated cells for 5 h, only 20.02% and 26.06% PI-positive cells were detected, respectively. The increase in the fluorescence intensity percentage observed in yeast cells treated with higher AF₄ and AF_5 (8 mg/L) concentrations agreed with the positive control values, indicating potential antifungal activity. In comparison, untreated cells showed negligible fluorescence.

Candida cells exposed to AF_4/AF_5 (8 mg/L) displayed PI uptake as evident from red fluorescence, indicating cell membrane damage and hence permeabilization [Figure 1C (×120) and Figure 1D (×200)] magnifications. Since the formation of cylindrical intravacuolar structures (CIVS) needs metabolically active cells with undamaged plasma membrane, the presence of yellow-green fluorescence (Figure 2C) in FUN-1 stained cells instead of red

TABLE 1 Correlation between	I Pl uptake	percentage	from flow	cytometry	and plate count.
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Drug (concentration)	PI uptake (%)	Plate	count
		Log reduction	Reduction (%)
AF ₄ (8 mg/L)	81.92 ± 8.7 SD	3.03	99.90 ± 0.79 SD
AF ₄ (16 mg/L)	99.51 ± 0.07 SD	3.64	99.97 ± 0.59 SD
AF ₅ (8 mg/L)	92.14 ± 6.8 SD	2.25	99.44 ± 0.82 SD
AF ₅ (16 mg/L)	81.35 ± 12.8 SD	3.10	99.92 ± 2.1 SD
AMB (1 mg/L)	20.02 ± 6.7 SD	4.12	99.99 ± 0.53 SD
AMB (2 mg/L)	26.06 ± 6.02 SD	4.43	99.99 ± 0.56 SD



fluorescence in lipopeptide-treated cells clearly indicates the loss of membrane integrity, metabolic activity, and cell death (Millard et al., 1997). The cell deaths are quite evident from the log reductions of 3.03 and 3.64 CFU/mL by AF₄ (8 and 16 mg/ L). In the lipopeptide-treated and AMB-treated cells CIVS were not observed. Collectively, it may be surmised that the prime targets of AF4 and AF5 might be the components of the cell membrane (Ramesh et al., 2023). The presence of a long acyl chain in antifungal cyclic lipopeptides such as iturin and fenzycin enables oligomer formation and subsequent insertion into the plasma membrane (Malina and Shai, 2005) and the organisation of aggregates in biological membranes triggering early membrane damage and membrane disintegration was also reported by Horn et al. (2013). The preliminary mechanism of action on the yeast cell membrane has been schematically depicted in the Supplementary Figure S2. Taken together, the results revealed that two lipopeptides demonstrated remarkable fungicidal effects on C. glabrata cells with enhanced cell membrane permeability, and damage. Consequently, the findings positively indicate that the lipopeptides AF4 and AF5 hold considerable promise as prospective antifungal agents.

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Data availability statement

The raw data supporting the conclusion of this article will be made available by the authors, without undue reservation.

Author contributions

MM was exclusively involved in the experimental works and preparing the preliminary draft. SMR oversaw the progress of the work and has been the co-principal investigator of the project. UR was involved in the logistics part, designing the experiments and revising the manuscript. All authors contributed to the article and approved the submitted version.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontierspartnerships.org/articles/10.3389/ abp.2024.11999/full#supplementary-material

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