

Original article

In vitro synergy of pseudolaric acid B and fluconazole against clinical isolates of *Candida albicans*

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Summary

Candida albicans is the most common fungal pathogen in humans. The emergence of resistance to azole antifungals has raised the issue of using such antifungals in combination to optimise therapeutic outcome. The objective of this study was to evaluate in vitro synergy of pseudolaric acid B (PAB) and fluconazole (FLC) against clinical isolates of C. albicans. The in vitro antifungal activity of PAB, a diterpene acid from Pseudolarix kaempferi Gordon, was evaluated alone and in combination with FLC against 22 FLC-resistant (FLC-R) and 12 FLC-susceptible (FLC-S) C. albicans using the chequerboard microdilution method and time-killing test assays. Synergism was observed in all 22 (100%) FLC-R strains tested as determined by both fractional inhibitory concentration index (FICI) with values ranging from 0.02 to 0.13 and bliss independence (BI) models. Synergism was observed in two of 12 (17%) FLC-S strains as determined by FICI model with values ranging from 0.25 to 0.5 and in three of 12 (18%) FLC-S strains as determined by BI model. For FLC-R strains, the drug concentrations of FLC and PAB, where synergistic interactions were found, ranged from 0.06 to 4 μ g ml⁻¹ and 0.5 to 4 μ g ml⁻¹ respectively. For FLC-S strains, the drug concentrations of FLC and PAB were $1-8 \ \mu g \ ml^{-1}$ and $0.5-4 \ \mu g \ ml^{-1}$ respectively. The BI model gave results consistent with FICI, but no antagonistic activity was observed in any of the strains tested. These interactions between PAB and FLC were confirmed using the time-killing test for the selected strains. Fluconazole and PAB exhibited a good synergism against azole-R isolates of C. albicans.

Key words: Pseudolaric acid B, Candida albicans, antifungal susceptibility, synergy, chequerboard method, time-kill curves.

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Introduction

Candida albicans is a major human fungal pathogen causing various forms of candidiasis ranging from chronic superficial mycoses such as vaginitis to severe and life-threatening systemic infections, predominantly in patients with a compromised immune system.¹ *Candida albicans* has been reported to be the fourth leading cause of nosocomial infections.²

Over the years, the polyene fungicidal agent Amphotericin B has become the standard treatment for candidal infections, but the severe nephrotoxicity of its conventional form and the costs of its lipid forms limit its widespread use. Currently, the azole antifungal

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compounds such as fluconazole (FLC) have emerged as the principal and first-line drugs used in the treatment of candidal infections in non-neutropenic patients. Fluconazole has the advantage of being stable parenteral formulation, and having excellent oral bioavailability and efficacy–toxicity profiles.^{3,4} However, resistance to FLC is increasing in *C. albicans* and other species.⁵ The emergence of drug resistance can probably be ascribed to the fungistatic rather than fungicidal characteristics of FLC action.⁶ Combination therapy is one approach that can be used to improve the efficacy of antifungal therapy for difficult-to-treat infections.⁷

The root and trunk bark of *Pseudolarix kaempferi* Gordon (Pinaceae), known as 'Tu-Jin-Pi' in China, have been traditionally used as a remedy for fungal infections of the skin. Pseudolaric acid B (PAB), which is a diterpene acid, was isolated and identified as the main antifungal constituent of *P. kaempferi* Gordon.⁸ It has been reported that PAB was active against *Trichophyton mentagrophytes, Torulopsis petrophilum, Microsporum gypseum* and *Candida* spp.⁹ A previous report also showed that PAB reduced the number of recovered colony-forming units significantly at different dosages in a murine model of disseminated candidiasis, while infected mice treated intravenously with PAB had a longer survival time than those treated with the vehicle alone.⁹

In an attempt to improve the antifungal effect of FLC, we investigated the combined effects of FLC and PAB against clinical isolates of *C. albicans* using alternative methods.

Materials and methods

Fungal strains and materials

Twenty-two clinical isolates of the FLC-R C. albicans strains¹⁰ were kindly provided by Jiang Y.Y. and 12 clinical isolates of the FLC-S C. albicans were obtained from the First Hospital of Jilin University and used in this study. In addition, three ATCC type *Candida* strains (C. albicans ATCC 10231, Candida parapsilosis ATCC 90018 and Candida krusei ATCC 6258) were acquired from the American Type Culture Collection (ATCC, Gaithersburg, MD, USA). The components of YPD broth (1% yeast extract, 2% peptone, and 2% glucose) were purchased from BD Biosciences, Inc. (Sparks, MD, USA). Pseudolaric acid B was purchased from the National Institute for the Control of Pharmaceutical and Biological Products (Beijing, China), FLC was obtained from Pfizer Inc. (New York, NY, USA) and stock solutions of varying concentrations were prepared in dimethyl sulphoxide (DMSO). Pseudolaric acid B and FLC were prepared using RPMI 1640 with glutamine broth medium, buffered to pH 7.0 with $0.165 \text{ mmol l}^{-1}$ morpholinepropanesulphonic acid.¹¹

Antifungal susceptibility testing

The minimum inhibitory concentrations (MICs) of PAB and FLC against the *Candida* strains mentioned above were determined by broth microdilution using twofold serial dilutions in RPMI 1640 medium as described by the Clinical and Laboratory Standards Institute (CLSI, formerly NCCLS) method M27-A.¹² The quality control (QC) strain, *C. krusei* ATCC 6258, and the reference strain, *C. parapsilosis* ATCC 90018, were included in each batch of susceptibility tests to ensure QC. The test was carried out in 96-well flat-bottomed microtitration plates. After agitation for 15 s, the plates were incubated at 35 °C without shaking, and readings were performed after 48 h of incubation by both visual reading and optical density (OD) determination.

For the visual reading, the MICs of FLC were determined according to CLSI procedure. For PAB, the MIC was defined as the lowest concentration showing 100% growth inhibition.⁹

For the OD determination, the MICs was defined as the lowest concentration of antifungal which resulted in 80% inhibition of growth compared with that of the drug-free control for FLC, and the MICs was defined as the lowest concentration showing 100% growth inhibition for PAB.⁹

Chequerboard method

The interaction between PAB and FLC against the 34 clinical isolates and the C. albicans ATCC 10231 strains mentioned above was assayed using a microdilution chequerboard technique.^{13,14} Drug dilutions were prepared to obtain four times the final concentration. A total of 50 µl of each FLC concentration was added to columns 2 to 12, and then 50 µl of PAB was added to rows B to H. To column 1, 50 µl of the medium containing the PAB solvent was added, and to row A, 50 µl of the medium containing the FLC solvent was added. The solvent DMSO in the medium comprised <1% of the total test volume. Thus, row A and column 1 contained only the azole and PAB, respectively, and the well at the intersection of row A and column 1 (well A1) was the drug-free well that served as the growth control. The final concentrations ranged from 0.0078 to $8 \ \mu g \ ml^{-1}$ for FLC, 0.125 to $8 \ \mu g \ ml^{-1}$ or 0.5 to $32 \ \mu g \ ml^{-1}$ for PAB, and the final inoculum size was 2.5×10^3 cfu ml⁻¹. After incubation, visual analysis of the MICs was performed and the OD values were measured at 595 nm. The percentage of growth in each well was calculated as the OD of each well. The background OD was subtracted from the OD of each well. Each isolate was tested in triplicate on different days. The background OD to be subtracted from the growth after incubation includes that of all the inoculated wells taken at time zero before incubation. This has to be distinct from the difference between the growth inhibition and the growth control after incubation.

Interpretation of drug interaction

To assay the *in vitro* interactions between FLC and PAB against each strain, the data obtained by the spectrophotometric method were analysed using two models, FICI and BI, both of which have been used previously to characterise antifungal drug interactions.¹⁵ Fractional inhibitory concentration index and BI are non-parametric models based on the Loewe additivity (LA) and BI theories respectively.¹³

Fractional inhibitory concentration index

The FICI method was defined by the following equation: FICI = FICA + FICB = $C_A^{comb}/MIC_A^{alone} + C_B^{comb}/MIC_B^{alone}$, where MIC_A^{alone} and MIC_B^{alone} are the MIC values of drugs A and B when acting alone and C_A^{comb} and C_B^{comb} are concentrations of drugs A and B at isoeffective combinations respectively.¹³ Low off-scale MIC values were converted to the lowest tested doubling concentration. Among all the FICIs calculated for each data set, the FICI_{max} was reported as the FICI in all cases unless the FICI_{max} was >4, in which case the FICI_{max} was reported as the FICI was as follows: an FICI value of ≤ 0.5 represented synergy, an FICI value between 1 and 4 represented indifference and an FICI value >4 represented antagonism.¹⁶

Bliss independence analysis

Bliss independence model is described by the equation $I_i = (I_A + I_B) - (I_A \times I_B)$, where I_i is the predicted percentage of inhibition of the theoretical combination of drugs A and B, and I_A and I_B are the experimental percentages of inhibition for each drug acting alone. As I = 1 - E, where E is the percentage of growth, by substitution into the former equation, the following equation is derived: $E_i = E_A \times E_B$, where E_i is the predicted percentage of growth of the theoretical com-

bination of drugs A and B, respectively, and $E_{\rm A}$ and $E_{\rm B}$ are the experimental percentages of growth of each drug alone. An interaction is described by the difference (ΔE) between the predicted and measured percentages of growth at various concentrations ($\Delta E = E_{\text{predicted}} - E_{\text{predicted}}$ E_{measured}). Using the non-parametric approach described by Prichard *et al.* [17], E_A and E_B are obtained directly from the experimental data. Because of the nature of the interaction, testing with microtitre plates and a twofold dilution of either drug results in a ΔE for each drug combination. In each of the three independent experiments, the observed percentages of growth obtained from the experimental data were subtracted from the predicted percentages after which the average difference of three experiments was calculated. When the average difference and the 95% confidence interval for the three replicates were positive, statistically significant synergy was claimed. When the difference and the 95% confidence interval were negative, significant antagonism was claimed. In any other case, BI was concluded. The BI model was derived by calculating the sum of the percentages of all statistically significant synergistic $(\sum SYN)$ and antagonistic $(\sum ANT)$ interactions. Interactions with <100% statistically significant interactions were considered weak, interactions with 100-200% statistically significant interactions were considered moderate and interactions with >200% statistically significant interactions were considered strong, as described previously.¹⁵ In addition, the numbers of statistically significant SYN and ANT combinations among the 77 combinations of drug concentrations tested were calculated for each strain.

Time-kill curves

Candida albicans in RPMI 1640 medium was prepared at the starting inoculum density of 10⁵ cfu ml⁻¹.¹⁸ For one chosen clinical isolate FLC-R C. albicans YL313, the concentrations used were 16 μ g ml⁻¹ (1/2 × MIC) for PAB and 256 μ g ml⁻¹ (1/2 × MIC) for FLC.¹⁹ For clinical isolate FLC-S C. albicans YL381, the concentrations used were 256 μ g ml⁻¹ (1/2 × MIC) for PAB and 4 μ g ml⁻¹ (1/2 × MIC) for FLC. Dimethyl sulphoxide comprised <1% of the total test volume. At various predetermined time points (0, 12, 24, 36 and 48 h after incubation with agitation at 35 °C), 100 µl of aliquot was removed from each solution and serially diluted 10fold in sterile water. Subsequently, 100 µl of each dilution was streaked on a Sabouraud dextrose agar plate. Colony counts were determined after incubation at 35 °C for 48 h. The experiment was performed in triplicate. Synergism and antagonism were defined as

Median MIC (range) of drug alone (µg ml ⁻¹) Strains FLC PAB Clinical isolates of FLC-R <i>Candida albicans</i> YL 313 512 (512) 32 (16–32) YL 313 512 (512) 32 (16–32) YL 313 512 (512) 32 (16–32) YL 319 512 (512) 32 (16–32) YL 331 512 (512) 32 (16–32) YL 333 512 (512) 32 (16–32) YL 333 512 (512) 32 (16–32) YL 333 512 (512) 32 (16–32) YL 334 512 (512) 32 (16–32) YL 335 512 (512) 32 (32) YL 334 512 (512) 32 (32) YL 335 512 (52–512) 32 (16–32) YL 345 512 (256–512) 32 (32) YL 346 5512 (526–512) 32 (16–32) YL 349 5512 (256–512) 32 (32) YL 349 5512 (256–512) 32 (32) YL 353 512 (256–512) 32 (32) YL 353 512 (256–512) <th>LA Median MIC (range) in combination (μg ml⁻¹) FLC PAB 0.125 (0.06–0.25) 1 (0.5– 1 (0.5–1) 1 (0.5– 1 (0.5–1) 1 (0.5– 1 (0.5–1) 1 (0.5– 1 (0.5–1) 1 (1) 0.25 (0.25–0.5) 1 (0.5– 1 (0.5– 1 (1) 0.5– 2 (1–4) 2 (1–4) 0.5 (0.25–1) 1 (1–2) 2 (1–4) 2 (1–4) 0.5 (0.25–1) 1 (1–2) 0.5 (0.25–1) 1 (1–2) 0.5 (1–4) 2 (1–4) 0.5 (1–2) 2 (1–4) 0.5 (1–2) 2 (1–4) 0.5 (1–2) 2 (1–4) 0.5 (1–2) 2 (1–4) 0.5 (1–4) 2 (1–4) 0.5 (1–4) 2 (1–2) 0.5 (1–4) 2 (1–2) 0.5 (1–2) 2 (1–2) 2 (1–4) 0.5 (1–2) 2 (1–2) 2 (1–2) 0.5 (1–2) 2 (1–2) 2 (1–2) 2 (1–2) 0.5 (1–2) 2 (</th> <th>combination PAB 1 (0.5-1) 1 (1) 1 (0.5-1)</th> <th>Result</th> <th></th> <th></th> <th></th> <th></th>	LA Median MIC (range) in combination (μg ml ⁻¹) FLC PAB 0.125 (0.06–0.25) 1 (0.5– 1 (0.5–1) 1 (0.5– 1 (0.5–1) 1 (0.5– 1 (0.5–1) 1 (0.5– 1 (0.5–1) 1 (1) 0.25 (0.25–0.5) 1 (0.5– 1 (0.5– 1 (1) 0.5– 2 (1–4) 2 (1–4) 0.5 (0.25–1) 1 (1–2) 2 (1–4) 2 (1–4) 0.5 (0.25–1) 1 (1–2) 0.5 (0.25–1) 1 (1–2) 0.5 (1–4) 2 (1–4) 0.5 (1–2) 2 (1–4) 0.5 (1–2) 2 (1–4) 0.5 (1–2) 2 (1–4) 0.5 (1–2) 2 (1–4) 0.5 (1–4) 2 (1–4) 0.5 (1–4) 2 (1–2) 0.5 (1–4) 2 (1–2) 0.5 (1–2) 2 (1–2) 2 (1–4) 0.5 (1–2) 2 (1–2) 2 (1–2) 0.5 (1–2) 2 (1–2) 2 (1–2) 2 (1–2) 0.5 (1–2) 2 (combination PAB 1 (0.5-1) 1 (1) 1 (0.5-1)	Result				
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512 (512) 512 (128–512) 512 (128–512) 512 (256–512) 512 (256–512) 512 (256–512) 512 (256–512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 513 (512) 513 (512) 513 (512) 514 (512) 514 (512) 515 (512) 515 (512) 512 (512) 512 (512) 512 (512) 513 (512) 51	0.125 (0.06–0.5) 2 (1–4) 1 (1–2) 0.5 (0.25–1) 2 (1–4) 2 (1–4)	1 (1)	0.03 (0.03–0.06)	SYN	856.8 (38)	-5.8 (2)	SΥN
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>512 5512 5512 5512 5512 5512 5512 5512	1 (1-2) 0.5 (0.25-1) 2 (1-4) 2 (1-4)	2 (1–4)	0.06 (0.03-0.04)	SYN	1180.6 (38)	-4.6 (2)	SYN
>512 512 (256-512) >512 (256-512) >512 (256) 512 (256-512) >512 (55-512) >512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 513 (512) 513 (512) 513 (512) 514 (512) 513 (512) 513 (512) 513 (512) 513 (512) 514 (512) 513 (512) 514 (512) 512 (512) 513 (512) 513 (512) 513 (512) 513 (512) 514 (512) 514 (512) 514 (512) 512 (512) 512 (512) 512 (512) 512 (512) 513 (512) 513 (512) 513 (512) 513 (512) 514 (512) 514 (512) 514 (512) 514 (512) 512 (512) 51	0.5 (0.25–1) 2 (1–4) 2 (1–4)	2 (1–4)	0.06 (0.03–0.13)	SYN	750.2 (32)	-25.4 (13)	SYN
512 (256–512) >512 (256–512) >512 (256) 512 (256–512) 512 (256–512) >512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 512 (512) 513 (512) 513 (512) 514 (512) 514 (512) 515 (512) 517 (512) 518	2 (1-4) 2 (1-4)	1 (1–2)	0.03 (0.03–0.06)	SYN	980.6 (31)	-55.8 (4)	SYN
>512 >512 256 (256) 512 (256–512) 512 (256–512) >512 256 (264–256) 512 (512) 512 (512) 512 (512) 512 (512) 512 (4–256) 512 (4–8) s (4–8) s (4–8)	2 (1-4)	1 (1–2)	0.06 (0.03–0.04)	SYN	518.8 (28)	-35.8 (6)	SYN
>512 256 (256) 512 (256-512) 512 (256-512) 512 (256-512) 512 (512) 512 (512) 512 (566) 5112 (256-512) 512 (256-512) 512 (266-512) 512 (264-256) 512 (266-212) 512 (266-212		2 (1–4)	0.07 (0.03–0.13)	SYN	582.1 (29)	-43.1 (7)	SYN
256 (256) 512 (256–512) 5512 (256–512) 5512 (256–512) 512 (512) 512 (512) 512 (56–512) 5112 (256–512) 5112 (256–512) 5112 (256–512) 512 (26–256) 512 (26–256) 512 (26–26) 513 (4–8) 5 (4–8)	1 (1)	2 (1–4)	0.06 (0.03-0.13)	SYN	630.8 (35)	-40.8 (5)	SYN
512 (256–512) >512 >512 256 (256–512) 512 (512) 256 (64–256) >512 (556–512) 512 (256–512) 512 (256–512) solates of FLC-S Candida albicans 8 (4–8) 8 (4–8)	0.25 (0.25–0.5)	2 (2–4)	0.06 (0.06–0.13)	SYN	250.5 (14)	-15.3 (4)	SYN
>512 >512 256 (256-512) 512 (512) 256 (64-256) >512 (256-512) 512 (256-512) 512 (256-512) 8 (4-8) 8 (4-8)	1 (0.5–2)	1 (1–2)	0.03 (0.03-0.07)	SYN	960.7 (32)	-30.4 (7)	SYN
>512 256 (256–512) 512 (512) 256 (64–256) >512 (256–512) 512 (256–512) 512 (256–512) 8 (4–8) 8 (4–8)	2 (1–4)	2 (1–4)	0.06 (0.03-0.13)	SYN	660.8 (36)	-42.1 (8)	SYN
256 (256–512) 512 (512) 256 (64–256) >512 (256–512) 512 (256–512) 512 (256–512) 8 (4–8) 8 (4–8)	1 (1–2)	2 (1–4)	0.06 (0.03–0.13)	SYN	1288.2 (51)	-26.6 (8)	SYN
512 (512) 256 (64–256) >512 512 (256–512) 512 (256–512) solates of FLC-S Candida albicans 8 (4–8) 8 (4–8)	2 (1–4)	2 (1–4)	0.07 (0.04–0.13)	SYN	883.1 (37)	-12.2 (5)	SYN
256 (64–256) >512 512 (256–512) 512 (256–512) 8 (4–8) 8 (4–8) 8 (4–8)	0.25 (0.13-0.5)	2 (1–4)	0.06 (0.03-0.13)	SYN	910.3 (28)	-27.6 (9)	SYN
>512 512 (256–512) 512 (256–512) 512 (256–512) 5126–512) 8 (4–8) 8 (4–8)	2 (1–4)	2 (1–4)	0.07 (0.04–0.13)	SYN	790.6 (33)	-28.5 (11)	SYN
	1 (1)	2 (1–4)	0.06 (0.03-0.13)	SYN	922.7 (43)	-16.4 (6)	SYN
	2 (1–4)	1 (1–2)	0.03 (0.03–0.06)	SΥΝ	1026.5 (58)	-6.6 (2)	SYN
8 (4–8) 8 (1–8)							
8 (1-8)	8 (2–8)	2 (1–4)	1 (0.5–1)	IND	108.8 (17)	-67 (9)	SYN
	8 (2–8)	1 (0.5–2)	1 (0.5–1)	IND	38.2 (21)	-85.5 (15)	IND
8 (4–8)	8 (2–8)	1 (0.5–2)	1 (0.5–1)	IND	15.6 (5)	-36.2 (14)	DNI
8 (2–8)	8 (2–8)	4 (2–4)	1.02 (0.52–1.02)	IND	40.8.2 (14)	-55.6 (20)	DNI
4 (2–8)	4 (2-4)	2 (1–4)	1 (0.5–1)	IND	29.2 (14)	-45.2 (33)	QNI
8 (2–8)	4 (2–8)	1 (2–4)	0.25 (0.25–0.5)	SYN	92.2 (50)	-52.4 (9)	QNI
8 (4–8)	8 (2–8)	0.5 (0.5–1)	1 (0.5–1)	IND	112.6 (24)	-42.5 (10)	SYN
	4 (2-4)	4 (2–8)	1.03 (0.50–1.03)	DNI	15.6 (4)	-86.2 (7)	DNI
YL 398 2 (2–4) 128 (64–128)	2 (1–4)	0.5 (0.5–1)	1 (0.5–1)	IND	75.8 (20)	-80.3 (8)	IND
2 (2–4)	1 (1–2)	1 (0.5–2)	0.25 (0.25–0.5)	SYN	121.6 (10)	-18.2 (3)	SYN
YL 423 4 (2–8) 128 (64–128)	4 (2-4)	1 (0.5–2)	1 (0.5–1.02)	DNI	55.4 (4)	-90.8 (9)	ONI
YL 433 8 (4–8) 256 (128–256)	8 (2–8)	1 (0.5–2)	1 (0.5–1.01)	IND	38.6 (5)	-7.6 (3)	QNI
ATCC 10231 2 (1–4) 32	0.25 (0.125–0.5)	32 (16–32)	1.13 (0.75–1.25)	IND	62.8 (6)	-17 (5)	DNI

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Synergy of PAB and FLC against Candida albicans

respective increases or decreases of $\geq 2 \log_{10}$ cfu ml⁻¹ in antifungal activity produced by the combination treatment compared with the more active agent alone after 24 h. A change of $< 2 \log_{10}$ cfu ml⁻¹ was considered indifferent.²⁰ Any decrease in the viable counts of the starting inoculum was considered 'killing'. Killing of >99.9% (3 logs) of the starting inoculum was defined as a fungicidal effect.¹⁹ Statistical analysis was performed using one-way analysis of variance (ANOVA), in SPSS 13.0 (SPSS Inc., Chicago, IL, USA) for Windows.

Results

Antifungal activities and interactions of drugs

The antifungal activities of the drug alone was assessed. For 22 clinical isolates of FLC-R *C. albicans*, the MIC values were ranged from 128 to \geq 512 µg ml⁻¹ for FLC and 16 to 64 µg ml⁻¹ for PAB treatment. For 12 clinical isolates of FLC-S *C. albicans*, the MIC values ranged from 2 to 8 µg ml⁻¹ for FLC and 128 to 512 µg ml⁻¹ for PAB treatment (Table 1). In this experiment, the MIC value of FLC against ATCC 10231 was 2 µg ml⁻¹ while the MIC of PAB against ATCC 10231 was 32 µg ml⁻¹. These results showed that PAB have better potential *in vitro* antifungal activity against FLC-R clinical isolates than against FLC-S clinical isolates.

The results of in vitro interaction between FLC and PAB against the C. albicans strains are shown in Table 1. There was good agreement between the FICI and BI models for the FLC/PAB combination treatment. For the 22 FLC-R strains tested, the interaction between FLC and PAB was SYN in all FLC-R strains using the FICI method, with FICI values ranging from 0.02 to 0.13. Using the BI method, all FLC-R strains showed very high percentages of SYN interactions, ranging from 250.5% to 1336.4% (Table 1). For 13 of the FLC-S strains tested, including C. albicans ATCC 10231, the FLC/PAB combination treatment displayed SYN or indifference with FICI values ranging from 0.25 to 1.25 using the FICI method, while the BI method indicated that the strain also showed either SYN or indifference interactions. We did not observe ANT interactions between FLC and PAB in either FLC-R or FLC-S C. albicans. The results above showed that there was good SYN antifungal effects against FLC-R clinical isolates when PAB was combined with FLC.

Time-kill curves

Further time–kill studies were conducted using FLC and PAB against one chosen clinical isolate FLC-R *C. albicans*

YL313 and one FLC-S C. albicans YL381. For FLC-R strain tested, time-kill curves verified synergism for the FLC/PAB combination [Fig. 1, resistant strain (R)]. The antifungal effect of PAB at 16 µg ml⁻¹ was more marked against C. albicans YL313 than FLC at 256 μ g ml⁻¹. Given an initial inoculum density of 10^5 cfu ml⁻¹, combination therapy yielded a 2.06 \log_{10} cfu ml⁻¹ decrease compared with 16 µg ml⁻¹ PAB after 24 h of incubation. The fungistatic activity of FLC was dramatically enhanced by the addition of PAB. In C. albicans YL313, the combination of FLC and PAB was fungicidal after 36 h of incubation (>99.9% decrease in viable counts). However, for FLC-S strain tested, an increases of 0.83 \log_{10} cfu ml⁻¹ in antifungal activity was produced by FLC/PAB combination treatment compared with the more active agent alone after 24 h. According to the result, indifference was observed for drug combinations against the FLC-S strain YL381 [Fig. 1, susceptible strain (S)].

Discussion

Previous reports revealed that the aqueous extract of P. kaempferi Gord significantly thickened the hypha of Trichophyton rubrum when observed under a transmission electron microscope, and in addition, the internal substances of the cytoplasm and organelles were degraded, the empty cavity appeared and an irregular membraneous structure resided in the fungal cell.²¹ It was also shown that PAB is effective at inhibiting tumour growth targeting microtubules.²² From the data presented above in this study, we found that PAB alone has moderate and high MIC values against the FLC-R and FLC-S C. albicans strains tested respectively. In addition, PAB showed more SYN activity when administered with FLC against FLC-R strains than against FLC-S strains. To our surprise, treatment with PAB alone is more effective against FLC-R strains than against FLC-S strains, while the phenomenon is different from the synergist compounds to FLC such as tacrolimus (FK506), cyclosporin A, amiodarone, ibuprofen and retigeric acid B reported previously.^{13,14,23–25} Notably, the synergistic effect of FLC in combination with PAB is better than FLC in combination with other reported chemicals.^{13,14,23–25} In our studies, compared with the FICI, the BI model not only allows for objective statistical criterion, but also fits all the experimental concentrations to construct a 3D graph in order to visualise the nature and intensity of drug combinations without arbitrarily choosing an end point.¹³ Compared with the fully parametric and semi-parametric response surface approaches,²⁶ the BI model is not dependent on

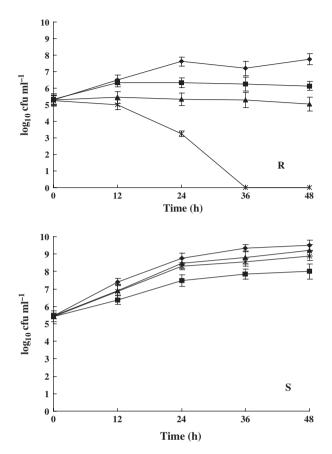


Figure 1 Time–kill curves of fluconazole (FLC) and pseudolaric acid B (PAB) alone and in combination against clinical azoleresistant strain (R) YL 313 and clinical azole-susceptible strain (S) YL 381. The strains at a starting inoculum density of 10^5 cfu ml⁻¹ were exposed to *in vivo*-achievable concentrations of 256 µg ml⁻¹ FLC, 16 µg ml⁻¹ PAB and 256 µg ml⁻¹ FLC + 16 µg ml⁻¹ PAB for R strain and 4 µg ml⁻¹ FLC, 256 µg ml⁻¹ PAB and 4 µg ml⁻¹ FLC + 256 µg ml⁻¹ PAB for S strain respectively. At 0, 12, 24, 36, and 48 h, aliquots were removed from each test tube to examine the cell viability. \blacklozenge , Growth control; \blacksquare , FLC; \blacktriangle , PAB; *, FLC + PAB. The experiments were performed three times. Data are expressed as mean ± standard deviation.

the data analysis program or the sigmoid dose–response, and thus does not fail to fit the data. Furthermore, the BI model showed an excellent reproducibility for the 77 combinations calculated for each strain. We also suggest the use of the BI model to assess the interaction between two drugs for its peerless advantages.²³ In general, our results indicated that there was a good agreement between the FICI and the BI models.

We verified the interactions in chequerboard microdilution using the time-killing test. Time-kill curves can provide growth kinetic information and give a more detailed picture of the effect of drug combinations on cell viability. This method is able to detect differences in the rate and extent of antifungal activity over time.⁷ Our results showed that the combination of PAB and FLC exhibited synergism or indifference against FLC-R or FLC-S *C. albicans* respectively. In this experiment, there was a good agreement between the conclusions drawn from the FICI method and the time–kill curves for the strains tested.

In conclusion, results in this study showed that the combination of FLC and PAB exhibited a good synergism against azole-R isolates of *C. albicans* using two non-parametric model approaches and that these results were verified by the time-killing test in some strains. However, the potential of using this combination therapy *in vivo* requires further investigation and further analysis is necessary to determine the underlying mechanism of this SYN interaction between FLC and PAB.

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