



Published in final edited form as:

J Drug Target. 2014 July ; 22(6): 536–542. doi:10.3109/1061186X.2014.905948.

Enhancement of antifungal activity by integrin-targeting of branched histidine rich peptides

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Abstract

The treatment of invasive candidiasis associated with growing numbers of immunocompromised patients remains a major challenge complicated by increasing drug resistance. A novel class of branched histidine-lysine (bHK) peptides has promising antifungal activity, and exhibits a mechanism similar to natural histatins, and thus may avoid drug resistance. The present studies evaluate ligand targeting of bHK peptides to fungal surface integrins by determining whether a cyclic RGD (cRGD) peptide with a large PEG linker could enhance bHK peptide antifungal activity. Whereas conjugates containing only the PEG linker reduced bHK peptide activity, conjugates with the cRGD-PEG ligand resulted in marked enhancement of activity against *C. albicans*. This study provides the first demonstration of benefit from ligand targeting of antifungal agents to fungal surface receptors.

Keywords

targeting; peptide; copolymer; conjugate; integrin; fungal infections

INTRODUCTION

Treatment of invasive candidiasis and other life-threatening disseminated invasive fungal infections remains a challenge, with a persistently high rate of mortality. Moreover, the number of immuno-compromised and immunosuppressed patients is increasing and as a result, the numbers of opportunistic infections including both superficial and invasive fungal infections are increasing [1–3]. Current therapeutics for invasive fungal infections involve

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Declaration of Interest

PVS, YL, and MCW have financial interest and equity in Aparna Biosciences. PVS, YL, MCW, and AJM are interested in commercialization of the biomacromolecules in this manuscript. QL and S-TC declare no conflict of interest.

three classes: polyenes, primarily amphotericin B [4]; azoles, including recently introduced voriconazole [4,5]; and echinocandins, such as caspofungin [6]. Nonetheless, use of these agents is limited by emerging resistance and/or serious side effects [5,7,8]. Thus, development of new and more effective antifungal agents, preferably with new modes of action, is urgently needed, particularly since invasive fungal infections affect multiple organs. Thus a major challenge has been achieving adequate biodistribution to achieve therapeutic levels at the tissues where the fungal infection reside.

Recent studies of pathogenic fungi have suggested that surface integrin-like receptors may play a role in attachment to endothelial cells, potentially an important contributor to disseminated invasive infections [9,10]. Furthermore, studies have shown that, like mammalian integrins, peptides with an RGD sequence inhibit fungal binding to endothelial cells [9,10]. Also, use of RGD peptides to target these fungal surface integrins is a potential strategy for targeted delivery of antifungal agents to enhance interaction and uptake in fungal-infected tissues, leading to improved efficacy.

Recently, antifungal activity was identified for several branched histidine-lysine (bHK) peptides, originally developed as gene delivery agents [11,12]. This novel bHK peptide class of macromolecules with non-natural branching, and capacity to incorporate non-natural amino acids via solid phase synthesis, offers a large range in biochemical structure to optimize anti-microbial activity. An initial set of bHK peptides was identified that have widespread antifungal activity against a variety of pathogenic fungal species, including candida, aspergillus, and cryptococcus [12,13], with a mode of action that appears to be similar to that of naturally occurring salivary histidine-rich histatins, with which microbial resistance has not been reported [14–16]. Most studies indicate that histatins are internalized and target either mitochondria or Trk.1 potassium channel, although one recent study indicates that histatins may act at the fungal cell surface [14–16]. Despite uncertainty in the precise target, both bHK peptides and histatins have very specific antifungal activity, in contrast to most linear antimicrobial peptides that have widespread antibacterial activity [17]. Interestingly, bHK peptides studied to date have significantly greater activity, as well as low mammalian cell cytotoxicity, making them particularly attractive for development of new treatments for disseminated invasive fungal infections. In addition, the non-natural bHK peptide branching increases the potential for conjugated ligands to act multivalently in binding to fungal surface receptors as a means to achieve targeting to widespread infections. Thus, the novel bHK peptide class has broad structural versatility for optimization and site specific ligand attachment at a molecular level.

This study investigated whether a cyclic RGD peptide ligand can facilitate delivery of bHK peptides to candida, and enhance antifungal activity. The small cyclic RGD pentapeptide studied, c(RGDfK), has been clinically validated for targeting a pair of mammalian integrins associated with endothelial cells at sites of neovasculature such as tumor angiogenesis with low levels of nonspecific tissue uptake [18]. The studies here evaluated whether conjugation of this ligand to antifungal bHK peptides using a large PEG-5000 linker could enhance antifungal activity, overcoming any potential adverse effects of conjugation. This study is the first to show enhanced antimicrobial activity by ligand targeting.

MATERIALS AND METHODS

C. *Albicans* strains

C. albicans (ATCC 10231, MYA-576) were obtained from American Type Culture Collection (Manassas, VA).

Maintenance of *C. Albicans*—*C. albicans* were maintained in yeast-maltose (YM) medium (Becton Dickinson, Sparks, MD) containing 0.3% yeast extract, 0.3% malt extract, 0.5% peptone, and 1.0% glucose. Minimal inhibitory concentration (MIC), metabolic, and uptake experiments were done in RPMI medium (buffered by MOPS, pH-7.2).

Synthesis of HK Peptides

The biopolymer core facility at the University of Maryland synthesized the branched HK peptides on a Ranin Voyager solid phase synthesizer (PTI, Tucson, AZ) as previously described [11]. If the peptide purity was less than 95%, then the peptides were further purified on an HPLC column with System Gold operating software by using a Dynamax 21-4 × 250 mm C-18 reversed phase preparative column with a binary solvent system. Further analyses of the peptides were performed with a Voyager MALDI-TOF mass spectroscopy (Applied Biosystems, Foster City, CA) and amino acid analysis (AAA Laboratory Service, Boring, OR).

PEG-cRGD conjugates of HK peptides

The above HK peptides were then modified with the PEG and the cRGD targeting ligand as follows. cRGD with a sequence, cyclo (Arg-Gly-Asp-D-Phe-Lys), was obtained from Peptides International (Louisville, KY). Targeting ligand cRGD was conjugated to the bHK peptide through a PEG molecule. Ligand targeted bHK peptide conjugates were synthesized in a two-step procedure as described previously with the polyethyleneimine polymer [18]. In the first step, cRGD was conjugated to a 5-KD molecular weight polyethylene glycol (PEG) by using a heterobifunctional PEG, maleimide-PEG-succinimidyl carboxymethyl (SCM), obtained from Creative PEG Works (Salem, NC). Molar equivalents of cRGD and maleimide-PEG-SCM were reacted in the presence of 1.5 equivalents of N,N-diisopropylethylamine (DIPEA) in dimethyl sulfoxide (DMSO). The reaction was completed in 1 h and the product was precipitated with dry ether. The resulting conjugate was characterized by mass spectrometry. In the second step, the Mal-PEG-cRGD was reacted with bHK peptide in DMSO at 4:1 molar ratio and in presence of 2 equivalents of DIPEA. The reaction mixture was stirred at room temperature for 24 h and the product was dialyzed in 25-KD MWCO dialysis tubing against 0.05% TFA/water for 48 h and was lyophilized and characterized by amino acid analysis.

Solid-phase $\alpha_v\beta_3$ binding assay

The binding of cRGD, H2K4b-PEG-cRGD and H3K(H)4b-PEG-cRGD to $\alpha_v\beta_3$ integrin was determined using a solid-phase competitive binding assay [19]. The cRGDfK(biotin-PEG-PEG)] peptide (Peptides International), together with the HK or control peptides, was used to detect binding to $\alpha_v\beta_3$ integrin. Microtiter 96-well vinyl assay plates (Corning, NY) were coated overnight at room temperature with 100 μ l/well of a solution of purified human

integrin $\alpha_v\beta_3$ in Triton X-100 (Millipore, Billerica, MA) at a concentration of 100 ng/ml in coating buffer ($1 \times$ TBS, pH 7.4, 2 mM $MgCl_2$, 1.5 mM $MgCl_2$) overnight at room temperature. The plates were then washed three times with binding buffer (0.1% Tween-20, 1% BSA in coating buffer). The wells were blocked for 1 h with 250 μ l of blocking buffer (3% BSA in binding buffer). The plates were washed three times with binding buffer. After blocking, 50 μ l of the competitors (peptide concentrations range from 1×10^{-4} to 0.5 mM) was first added into wells, and then 50 μ l of c[RGDfK(biotin-PEG-PEG)] peptide were added into the wells with or without pre-added competitors. The plate was gently mixed and incubated for 1 h at 37°C. After washing away the unbound competitors and c[RGDfK(biotin-PEG-PEG)] peptide, the bound c[RGDfK(biotin-PEG-PEG)] peptide was detected with avidin-HRP (horseradish peroxidase) and TMB (3,3',5,5'-tetramethylbenzidine) substrate with absorbance readings at 450 nm by using a microplate reader. Background reading was subtracted from all measurements. The wells without competitors were used as control (100% binding).

Antifungal activity of bHKP and conjugates

Turbidity—Antifungal efficacy of bHK peptides was determined by measuring the viability of the cells (see next section) or by determining the growth of yeast cells in 96-well microtiter plates. Yeast cells were diluted between 2.5×10^3 and 5×10^3 cells/ml in RPMI1640-MOPS medium; 55 μ l of the cell suspension were then added to each well of a 96-well plate containing 45 μ l of bHK peptides with final concentrations of 0.1, 1, 2, 5, 10, 15, 20, and 25 μ M. Peptide-free controls were also included. The microtiter plates were then incubated at room temperature or 37°C for 24–48 h. After incubation, the microplates were shaken vigorously on a plate shaker to disperse the cells into a uniform fungal suspension. Turbidity of the suspension, a measure of fungal growth, was measured at 595 nm with a microplate reader with each data point representing the mean and standard deviation. MIC was defined as the lowest concentration of the peptide that resulted in at least 95% inhibition from untreated control and no visible growth.

WST-1 Cell viability Assay—Fungal growth was also assayed using the WST-1 assay which determines the mitochondrial activity of the treated and control fungal cells. *C. albicans* cells were cultured in a micro-plate with 100 μ l of RPMI-1640-MOPS containing increasing concentrations of bHK peptides (0.1, 1, 2, 5, 10, 15, 20 and 25 μ M) for 48 h at 37°C. At the end of the treatment, 10 μ l of WST-1 assay solution were added to each well. After gentle shaking, the plate was incubated for 30 min., and then shaken again for 1 min. WST-1 tetrazolium salt (Dojindi, Rockville, MD) was reduced to formazan by cellular dehydrogenases, generating a deep yellow colored formazan that was measured at 450 nm (650 nm as reference) in a Microplate reader. The color is directly correlated to cell number. Cells without bHK treatment were used as a control. Readings were then converted to percent absorbance, with control set at 100% and medium (negative control) at 0%.

Fluorescent Binding and Uptake of H2K4b and H2K4b-PEG-cRGD

C. albicans (ATCC 10231; 1×10^5 cells/ml) were added to each well of a four-chamber culture slide (BD Sciences, Bedford, MA) containing RPMI1640 with 20 mM MOPS (300 μ l). After 4 h, fluorescently-labeled (CF750) peptides, H2K4b and H2K4b-PEG-cRGD,

(final concentration, 100 ug/ml) were incubated with the cells for an additional 4 h at room temperature. The cells were then fixed with 4% paraformaldehyde and counterstained with cyto 9 (Invitrogen, Grand island, NY). After the coverslips were placed on the slides and sealed with cyto seal-60 (Richard-Allan scientific, Kalamazoo, MI), cell images were captured with a fluorescent Olympus IX81 microscope, fitted with a Hamamatsu Photonics C9100-02 EMCCD camera.

RESULTS

H2K4b-PEG-cRGD conjugates

Previous studies had shown that antifungal activity of bHK peptide was greater with increasing branching, up to the maximum of four branches. One of the most active species identified had a branch sequence that contained five repeats of a tripeptide with two histidines and one lysine, called H2K4b. The structures and sequences of H2K4b and H3K(H)4b peptides are shown in Figure 1. While these two anti-fungal bHK peptides have branches with several lysine residues, the vastly greater histidine content results in a macromolecule with physicochemical characteristics unlike the cationic anti-microbial peptides which bind and destabilize cell membranes with low specificity. However, the lysine content provides numerous primary amines for random conjugation by use of methodology we developed to generate cyclic RGD (cRGD) peptide ligand conjugates via large PEG linkers of polycation nanoparticle carriers of nucleic acids [18]. Conjugates of H2K4b and H3K(H)4b were prepared using this random attachment method with approximately four PEG-cRGD per macromolecule, labeled H2K4b-PEG-cRGD and H3K(H)4b-PEG-cRGD, respectively. Also, a control conjugate was prepared with only a PEG linker, labeled H2K4b-PEG.

bHK-PEG-cRGD binding to $\alpha_v\beta_3$ integrins

Competitive binding assays were used to assess the effect of the cRGD ligand conjugate on binding to immobilized $\alpha_v\beta_3$ integrin, shown in Figure 2. Whereas the non-ligand bHK peptides (H2K4b and H3K(H)4b) did not bind immobilized integrin, the ligand-conjugated H2K4b-PEG-cRGD and H3K(H)4b-PEG-cRGD showed higher affinity than free cRGD pentapeptide. Multivalency of cRGD ligands on the bHK peptides likely accounts for higher affinity for integrin compared to the free cRGD. Moreover, on the basis of data obtained from untargeted bHK peptides that show no competitive binding, cationic charge associated with the bHK-PEG-cRGD peptides had little contribution to integrin binding with this assay.

H2K4b-PEG-cRGD antifungal activity

The antifungal activity of these ligand-conjugates against two strains of *C. albicans* was determined and compared with that of the H2K4b parent macromolecule. In conditions favoring cell growth of candida shown in Figures 3 and 4, PEGylation of H2K4b resulted in complete loss of antifungal activity, whereas the cRGD-PEG conjugate showed increased potency (MIC: H2K4b-PEG-cRGD, 1.5×10^{-5} M; H2K4b $> 2.5 \times 10^{-5}$ M). These effects were observed by both turbidity and WST-1 metabolic assays. Compared to unmodified H2K4b, the ligand conjugate exhibited an altered dose response curve for metabolic inhibition, with a sigmoid shape expected for a binding mediated effect, shown in Figure 4. These effects of

ligand conjugation on fungal growth inhibition were the same for both fluconazole-sensitive (10231) and refractory (MYA-576) strains. Notably, the cRGD peptide had no inhibitory effect on the growth of *C. albicans* (data not shown).

In addition to the antifungal assays, we examined the antifungal effects of H2K4b-PEG-cRGD versus H2K4b with light microscopy. After overnight incubation of *C. albicans* at 37°C, which enables both hyphae and cell growth, there were marked differences observed between the H2K4b and H2K4b-PEG-cRGD treatment groups. Representative results, shown in Figure 5, illustrate the increased potency by ligand-targeting of H2K4b, as well as a shift in the effects on cells vs. hyphae. Whereas H2K4b significantly reduced the number of cells and hyphae, their growth in “nests” was still apparent. In contrast, there were only a few scattered hyphae in the H2K4b-PEG-cRGD treated group.

Comparison of binding and uptake into *C. albicans* of targeted and non-targeted H2K4b

We compared the binding of fluorescently-labeled H2K4b and H2K4b-PEG-cRGD to the *C. albicans*. As shown in Figure 6(B) and (E), binding and uptake of H2K4b-PEG-cRGD to *C. albicans* were much greater than binding and uptake of H2K4b, consistent with greater potency of the targeted bHKP at this concentration.

H3K(H)4b-PEG-cRGD

Given that the cRGD-ligand enhanced the potency of H2K4b, we examined H3K(H)4b, which has a similar structure but higher content of histidine and has been used in studies with RNAi nanoparticles [20]. The results also showed an increase in antifungal activity of H3K(H)4b by ligand conjugation with the cRGD pentapeptide in both fluconazole-sensitive and resistant strains, shown in supplementary Figure S1.

Discussion

The present study was designed to determine whether the biological antifungal activity of an emerging class of branched antimicrobial macromolecules composed of histidines and lysines could be enhanced by ligand-mediated targeting. Previous investigations have found $\alpha_v\beta_3$ integrin-like receptors on the surface of several pathogenic fungi, including *C. albicans* [9,10]. These integrins are up-regulated on cell surfaces and hyphae of candida species and appeared to have an important role in binding of candida to endothelial cells and invading tissues. We found that the ligand conjugated bHK peptides that targeted the integrins on the surface of *C. albicans* resulted in marked growth inhibition of the fungus. Although in this report we did not examine whether our ligand-targeted peptides reduced binding of the fungus to activated endothelial cells, it will be of interest to determine if the H2K4b-PEG-cRGD has a dual mechanism of action in reducing growth of *C. albicans in vivo*: a direct inhibitory or killing action as described here; and/or an indirect action, preventing *C. albicans* from interaction with vitronectin [10].

We also examined whether the random PEGylation conjugation method used to couple the peptide ligand to the antifungal bHK peptides affected the anti-fungal activity of this molecule observed in our previous study [12], by interfering with interactions between bHK peptides and the surfaces of fungi. PEGylation of proteins is well established to enhance

pharmacology for therapeutic application, such as minimizing cytokine induction, reducing interaction with cell surfaces such as the cells of the reticuloendothelial system, thereby increasing the circulating half-life of the PEGylated proteins in the blood stream. Since it is possible that PEGylation might interfere with the activity of H2K4b, we synthesized and tested two different conjugates of this peptide, one with PEG alone (H2K4b-PEG) and the other with cRGD attached to H2K4b through the PEG (H2K4b-PEG-cRGD). Indeed, we observed that the PEG-5000 linker had a PEGylation effect when no cRGD targeting ligand was present, significantly diminishing the observed anti-fungal activity of H2K4b. Interestingly, the addition of ligand to the end of PEG linker enhanced its antifungal activity: the H2K4b-PEG-cRGD peptide exhibited more potent and more consistent anti-fungal activity than did the unmodified form. Differences in activity and binding suggest quantitative and perhaps qualitative differences in the inhibition of fungi, most likely through the integrin-like receptors reported in previous studies [9,10]. Moreover, we think that the most likely explanation for the opposing results is that the PEGylation interferes with the binding mechanism of unmodified bHK peptide interaction with the fungi whereas the cRGD ligand conjugate provides a new mechanism of interaction through the $\alpha_v\beta_3$ integrins. In support of this interpretation is the observation that fluorescently labelled cRGD conjugate exhibited far higher levels of uptake into fungal cells throughout the region stained with a nucleic acid selective dye. Importantly, another antifungal bHK peptide also showed a similar increase in potency and shift in antifungal activity by the integrin binding ligand conjugation. Thus, the judicious addition of cRGD-PEG conjugates to bHK peptides augmented their antifungal activity, and indicates that targeting to surface expressed fungal integrin homologs when PEG linkers are utilized is promising for enhanced pharmacology including enhanced localization at sites of fungal infection that is expected to enhance efficacy and therapeutic index. Although our studies demonstrated that the ligand-mediated antifungal activity was enhanced when using a relatively large PEG-5000 linker, future studies are required to determine whether PEG linkers of different molecular weights will further augment activity *in vitro* and *in vivo*.

In addition to the cRGD peptide, a sufficient cationic charge on the HK peptides is likely essential to maintain its antifungal activity. Random addition of –PEG-cRGD with the conjugation method used reduces the effective cationic charge on HK, and creates a steric barrier, which was found to reduce antifungal activity. Therefore, alternative conjugation approaches may provide better retention of antifungal activity, such as site-specific addition of the –PEG-cRGD conjugate or intracellular cleavage. The addition of a PEG-ligand to specific locations could be done by adding reactive amino acids (e.g., cysteine) within the peptide that would not be likely to affect the antifungal activity (e.g., cysteine attached to the C-terminal end of the lysine core and PEG is conjugated to the sulfhydryl group of cysteine). Our group modified several bHK peptides by this method so that technical problems are unlikely [20]. Alternatively, the fungal binding by conjugated bHK peptides could be maintained by adding more lysines at the N-terminal ends of the branches. Exploring several methods to add ligands to different bHK peptides may be important because of differences in their activity, lysine content, and safety profile.

Conclusions

This study investigated whether a small cyclic RGD peptide ligand selective for mammalian $\alpha_v\beta_3$ and $\alpha_v\beta_5$ integrins can facilitate delivery and antifungal activity of bHK peptides via integrins on the surface of *C. albicans*. The studies demonstrated conjugation of this ligand to bHK peptides could enhance antifungal activity, overcoming any adverse effects of the conjugation on the bHK peptide antifungal activity. This study is the first to show enhanced antimicrobial activity by ligand targeting, and opens the door to what may be an important new direction for addressing difficult to treat infections. Specifically for bHK peptide antifungal therapeutic candidates, on the basis of these and our previous results we anticipate that ligand-mediated targeting will provide a potent broad spectrum therapeutic for treatment of life threatening invasive candidiasis [12,13], and potentially many other invasive fungal infections that require parenteral treatment. Nonetheless further studies of this advance are needed to explore more fully the therapeutic potential. In addition to ligands targeting candida integrin, different ligands may be considered to enhance targeting and activity toward a broad range of pathogenic fungi [21], adding to the versatility of this approach.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

The authors thank P. Talalay for careful reading and invaluable editing assistance of this manuscript. This work was supported by the National Institutes of Health (R01-CA136938).

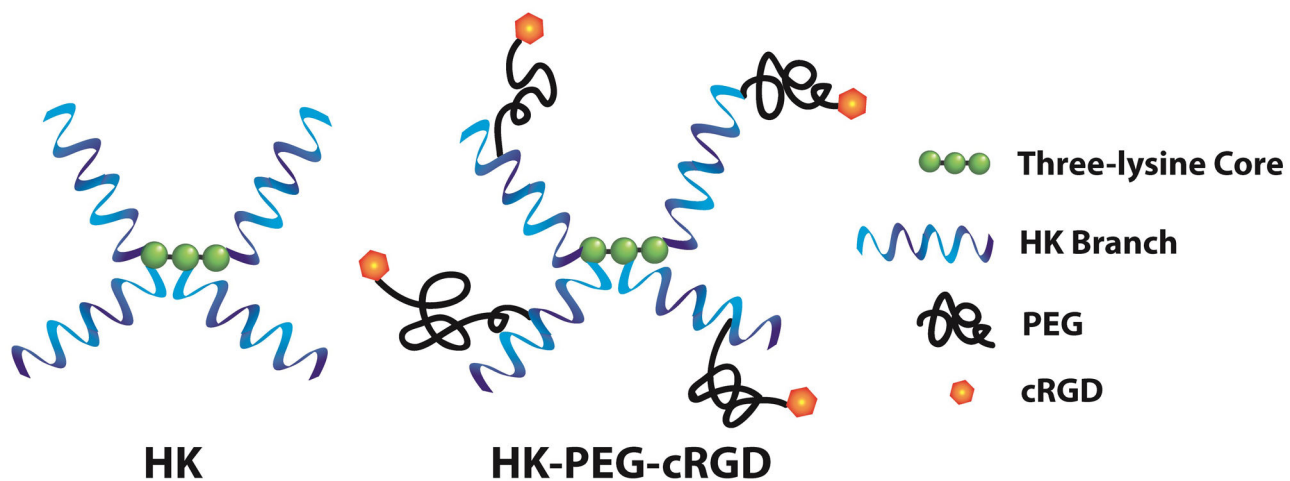
ABBREVIATIONS

bHK	branched histidine-lysine
cRGD	cyclic peptide containing arginine-glycine-aspartic acid motif
DIPEA	N,N-diisopropylethylamine
DMSO	dimethyl sulfoxide
HK	generic term for histidine-lysine peptides
H3K(H)4b and H2K4b	two unmodified four-branched peptides that differ in their histidine and lysine content
HRP	horseradish peroxidase
Mal-PEG-SCM	maleimide-PEG-succinimidyl carboxymethyl
PEG	polyethylene glycol
PEG-cRGD	conjugate of PEG and cRGD attached to a branched HK peptide
TFA	trifluoroacetate
TMB	3,3',5,5'-tetramethylbenzidine

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**Figure 1.**

Schematic structure of unmodified and modified bHKP. The unmodified peptides consist of four histidine-lysine branches originating from the lysine core. The HK branches contained KHKHHKHHKHHKHHKHHKHK for H2K4b or KHHHKHHHKHHHKHHHK for H3K(H)4b. Modified HK were similar except that PEG-cRGD conjugates were added randomly to the branches.

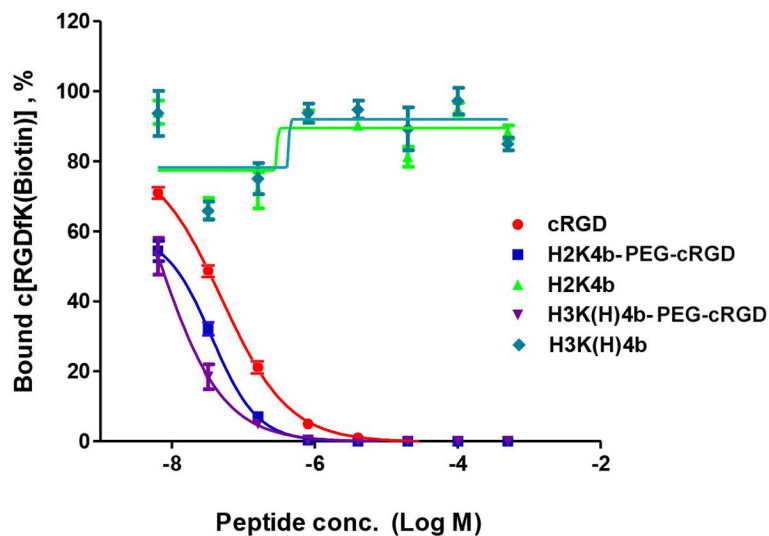


Figure 2. Competitive inhibition of biotinylated RGD peptide binding to immobilized integrin $\alpha_v\beta_3$ by cRGD-conjugated and unconjugated HK peptides. Competitive binding curves were obtained from the interactions between immobilized $\alpha_v\beta_3$ and the commercially available c[RGDfK(Biotin-PEG-PEG)] peptide in the presence of the cRGD conjugates. Binding in the absence of competitor was set at 100%. Bound c[RGDfK(Biotin-PEG-PEG)] was detected by avidin-HRP and TMB substrate.

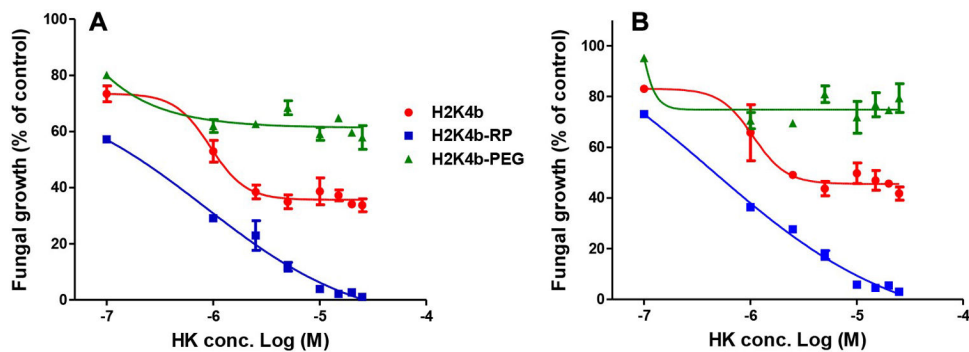


Figure 3.

In vitro antifungal activities of H2K4b and conjugates against two different strains of *C. albicans* (A, 10231; B, MYA-576). *C. albicans* cells were cultured in a microplate with 100 μ l of RPMI-1640 containing increasing concentrations of bHK peptides (0.1–25 μ M) for 48h at room temperature. At the end of treatment, fungal growth was measured at 595 nm. Cells without peptide treatment were used as a control. H2K4b-RP represents H2K4b-PEG-cRGD.

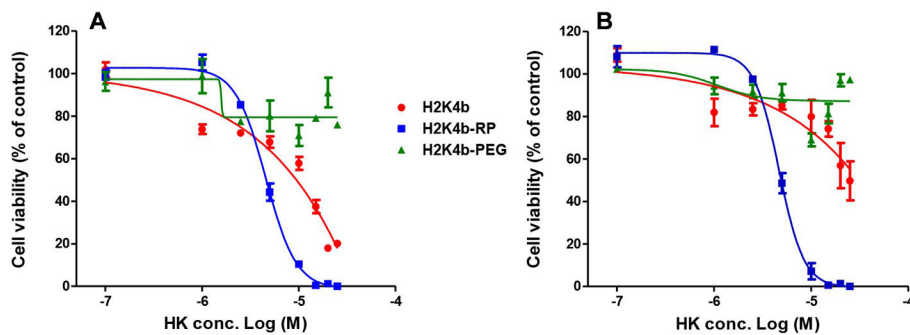


Figure 4. WST-1 cell viability assay. *C. albicans* cells (A, 10231; B, MYA-576) were cultured in a microplate with 100 μ l of RPMI-1640 containing varying concentrations of bHK peptides for 48h at room temperature. At the end of the treatment, 10 μ l of WST-1 assay solution were added to each well, and after 30 mins, the reduced formazan was measured at 450 nm (650nm as reference). Cells without peptide were used as control. H2K4b-RP represents H2K4b-PEG-cRGD.

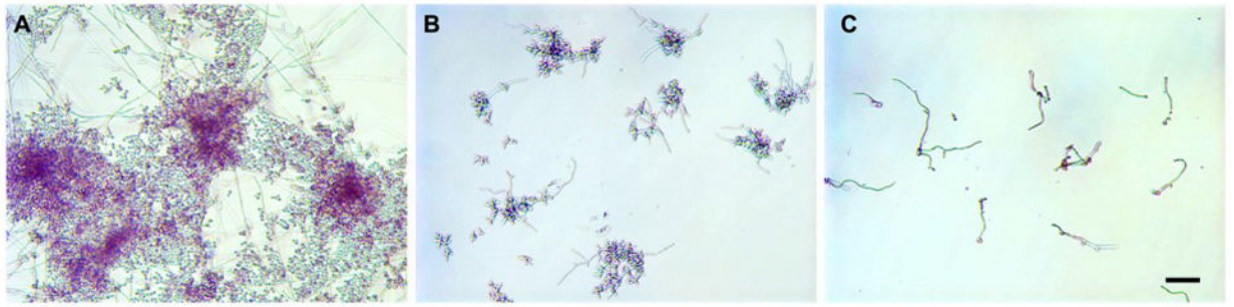


Figure 5.

Effects of bHK peptides and conjugates on the growth of MYA576, a fluconazole-resistant *C. albicans*. Fungal cells were incubated with 20 μM of H2K4b or conjugates at 37°C for overnight. (A) untreated; (B) 25 μM of H2K4b; and (C) 25 μM of H2K4b-PEG-cRGD. Bar represents 20 μm.

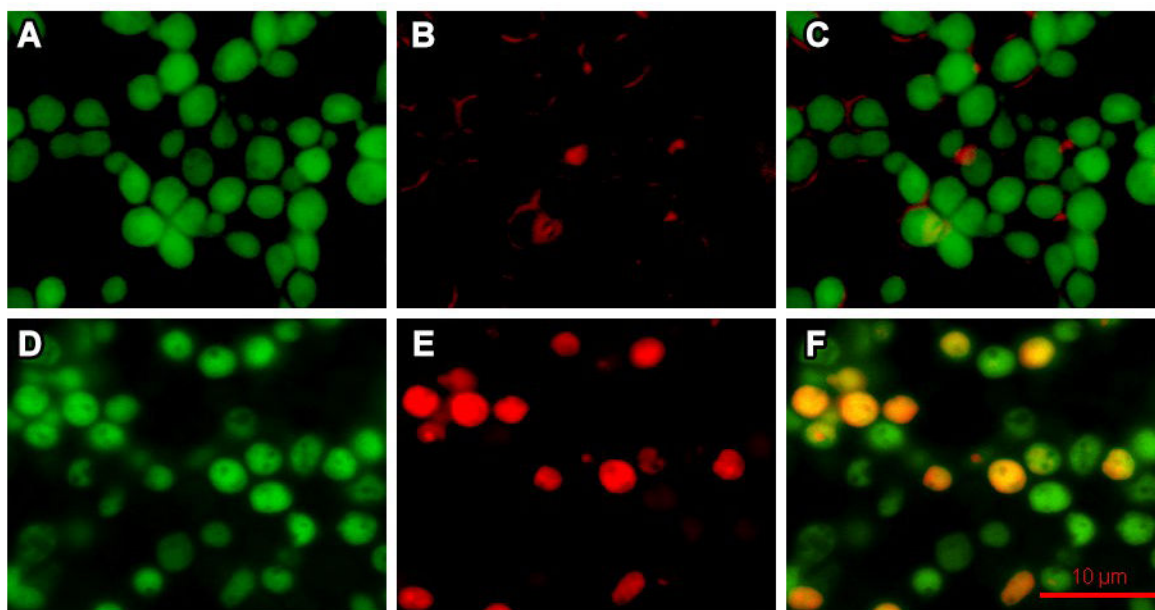


Figure 6. Comparison of binding and uptake into *C. albicans* with targeted and untargeted H2K4b. After fluorescent-labeled (CF750 - red) peptides, H2K4b and H2K4b-PEG-cRGD, (final concentration, 10 μ M) were incubated with *C. albicans* for 4h at RT, the cells were fixed and counterstained with cyto 9. Cell images were then captured with a fluorescent Olympus IX81 microscope. (A, B, C), H2K4b; (D, E, F), H2K4b-PEG-cRGD; (A, D), cyto 9 counterstain; (B, E), labeled H2K4b and H2K4b-PEG-cRGD peptides; (C, F), merged images of labeled peptides with counterstain. The bar represents 10 μ m.