



Identification of amino acid tethered triazoles as potential antifungal leads: appraisal of their mode of action

Parteek Prasher

Department of Chemistry, University of Petroleum and Energy Study, Dehradun, Uttranchal, (India)

ABSTRACT

A new set of compounds containing indole nucleus appended with carefully selected amino acids via a triazole linker are identified. These compounds showed substantial antifungal activities against the BWP17 strain of candida albicans. The preliminary screening of the compounds was done by docking in the active site of lanosterol 14- α demethylase enzyme (pdb ID - 3JUS) with ArgusLab.exe software. Subsequently the screened compounds were subjected to in-vivo antifungal assays. The probable mechanism of action of the test compounds was studied by GC analysis of the membrane metabolites. The compounds with tryptophan and histidine substitution were found to be most effective with MIC₈₀ of 50.2 μ g/ml and 78.81 μ g/ml respectively. In the fungal cells treated with compound **3b** the lanosterol levels were found to increase indicating that lanosterol 14- α demethylase enzyme could be the probable target. Additionally, the R6G influx/ efflux assay provided the evidence for the fungicidal properties of the test compounds.

Keywords: CA BWP 17 – candida albicans BWP 17, R6G – Rhodamine 6G, GC – gas chromatography, MIC₈₀ - minimum inhibitory concentration for 80% inhibition

INTRODUCTION

Diseases caused by fungus are one of the major global challenges today. Starting from local infections to systemic mycosis the fungi are causing more deaths compared to malaria and tuberculosis. Yeasts, primarily the Candida albicans species are the most noxious mediators of invasive mycoses predominantly among the immuno compromised patients [1-2]. The pathogenicity and infectiousness of the fungal diseases therefore needs to be checked and regulated.

Various classes of antifungal drugs have been developed till date to counter the menace created by fungal diseases. The polyene antifungals like amphotericin-B, natamycin primarily target ergosterol [3] which helps in maintaining the fungal membrane integrity. These amphiphilic polyenes on interaction with the fungal membrane make it leaky by creating the pores which eventually leads to the membrane disruption and fungal death. The azole antifungal drugs on the other hand target the CYP P450 14- α demethylase enzyme [4] which mediates the biosynthesis of ergosterol from lanosterol. This alters the morphology and physical properties of the fungal membrane by impeding the functioning of the membrane proteins. Another class of antifungals, allylamines [5] target the enzyme squalene epoxidase which mediates the conversion of squalene to squalene-2,3-epoxide. The allylamines like terbinafine leads to the build up of squalene causing a pH imbalance in the fungal membrane thereby limiting its growth. The newest class of antifungals, Echinocandins [6] impair the functioning of beta-1, 3-D-glucan synthase enzyme in fungi by noncompetitive inhibition eventually leading to the cell death. Despite all the efforts the antifungal therapy is still far off the target because of the efflux pumps operating in the fungal membranes [7]. The present work focuses on the design of the compounds chiefly to target the lanosterol 14- α demethylase enzyme present in the fungal membrane. The biological importance of these compounds is already been checked for their antiinflammatory

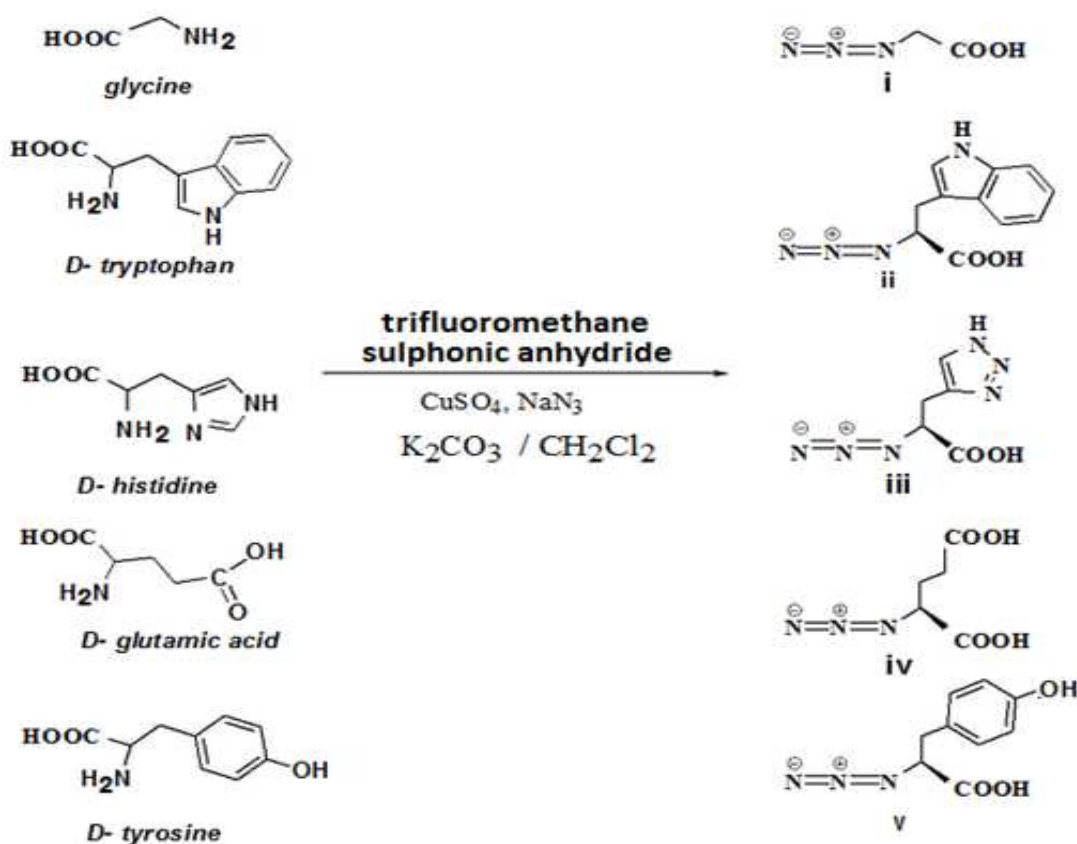
property by targeting the enzymes 5-LOX and COX-1/2 enzymes [8]. Interestingly, besides targeting the desired enzyme the compounds were also found to have membrane lysing properties.

The validation of indole moiety and azole nucleus in the antifungal therapeutics for targeting the fungal membrane has been widely acclaimed. The indole nuclei substituted with the carefully selected amino acids via azole linkage has been tested for its antifungal efficacy against the BWP17 strain of candida albicans. The screening of the appropriate amino acids was done by docking the test compounds in the lanosterol 14- α demethylase enzyme (pdb ID – 3JUS) with the help of ArgusLab.exe software. Apart from that, the Lipinski parameters were also calculated for the test compounds theoretically by molinspiration software. From the preliminary investigations, it was observed that the compounds having glycine, tryptophan, histidine, glutamic acid and tyrosine substitution displayed a better docking characteristics in the active site of the enzyme. The hydrophilicity index (ClogP value) for these compounds was also evaluated experimentally by determining the octanol/ water partitioning coefficient by the shake flask method and the concentration of analytes was found by quantitative HPLC method. The hydrophilicity index thus obtained was found to be quite similar to that required by the lipinski parameters.

EXPERIMENTAL SECTION

Synthesis

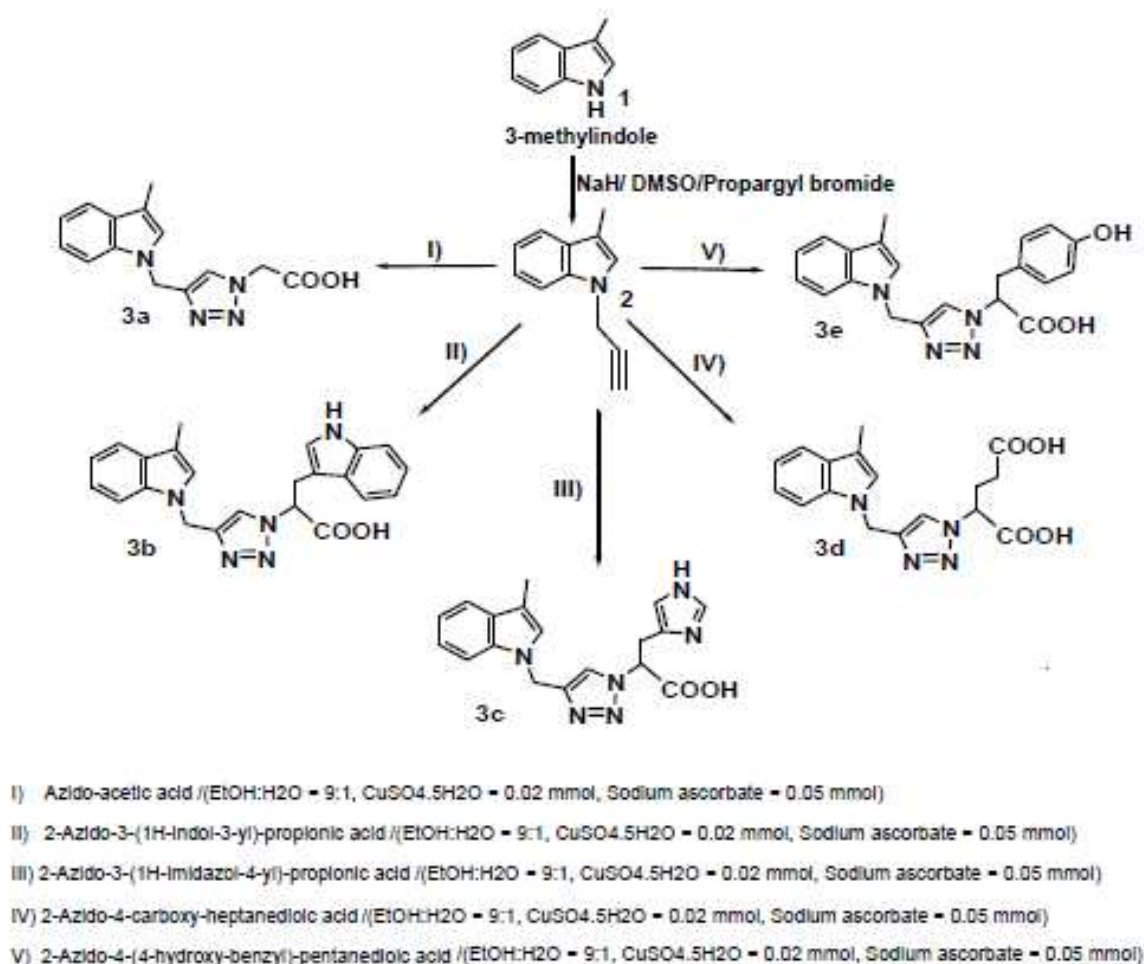
The L-amino acids (1 mmol) were dissolved in 20 ml of DCM in a round bottom flask kept on an ice bath. The mixture was stirred for a few minutes till the solution becomes clear. Activated K₂CO₃ (1.2 equivalent, activated by storing at 100 °C overnight) was then added to this solution. Afterwards trifluoromethanesulphonic anhydride (0.1 mmol) was very carefully added, dropwise to the reaction mixture. On addition, the colour of the reaction mixture immediately changes from colourless to light green. After stirring the contents of reaction mixture for 2-3 minutes was added CuSO₄.5H₂O (2.0 mmol) followed by the addition of sodium azide (1.0 mmol). The progress of the reaction was monitored by thin layer chromatography. The reaction completed in about 6-8 hours. After the reaction completion, the reaction mixture was washed with 15 ml of distilled water and the product was extracted by chloroform using a separator funnel. The final product was azide adduct of the amino acid which is generally procured in a very good yield (compounds i, ii, iii, iv and v, Scheme 1).



Scheme 1

3-methylindole (1) was dissolved in 5 ml DMSO and allowed to stir for a few seconds till a clear solution is

obtained. To this solution was added sodiumhydride (1.2 equivalents, washed with dry hexane to remove the paraffin coating). The reaction mixture was now stirred for 3-4 minutes till a visible change in the colour of the reaction mixture is observed. The change in colour corresponds to the generation of anion on the indole nucleus. Further, propargyl bromide (1.2 equivalents) was added carefully; drop wise to the reaction mixture. The completion of the reaction was monitored with the help of thin layer chromatography. Reaction completes in 25 minutes to give the product (2). The product (2) was dissolved in 9:1 solution of EtOH: H₂O to get a clear solution. To this solution was immediately added 0.02 mmol CuSO₄.5H₂O. After waiting for 5 minutes, 0.05 mmol sodium ascorbate was added to the reaction mixture and the contents were allowed to stir for 10 minutes. Finally, the azidoamino acid obtained in scheme 1 was added to the reaction mixture. The monitoring of the reaction was done by thin layer chromatography. The reaction generally completes in 10-12 hours to get the products 3a-3e (scheme 2) in very good yield. The further refining of the products was done by column chromatography using ethylacetate: hexane as eluent. All the reactions were performed in vaccum fume hood by using carefully cleaned oven-dried lobachemie glassware. The stirring of the reaction mixtures was done on REMI made magnetic stirrers using teflon coated magnetic beads. Drying of diethyl ether was done over activated anhydrous calcium chloride followed by passing the thin sodium wire through it. Refluxing of acetonitrile was done over anhydrous P₂O₅ followed by distillation over anhydrous K₂CO₃. The dried acetonitrile was finally stored over the activated molecular sieves of size 4 Å to prevent the infiltration of moisture. Reactions were monitored by TLC using silica gel GF254. The chromatograms thus developed were viewed in ultraviolet light and staining was done with iodine. Column chromatography was done to purify the compounds. Silica gel of 100-200 mesh size was used using hexane and ethyl acetate as preferred eluents.



Scheme 2

¹H and ¹³C NMR spectra and DEPT-135

The spectra were recorded at a frequency 500 MHz and 125 MHz for ¹H and ¹³C spectra respectively. CDCl₃ and DMSO were used as preferred solvents with TMS as an internal standard. The ¹H NMR spectral data were reported in form of chemical shifts (δ ppm), multiplicity (s = singlet, br s = broad singlet, d = doublet, dd = double doublet, t

= triplet, m = multiplet) and the coupling constant (expressed as J in Hertz). The ^{13}C and DEPT-135 NMR spectral data were represented in terms of chemical shift with +ve signals corresponding to CH_3 and CH carbons and the -ve signals corresponding to CH_2 carbons while an absent signal points towards a quaternary carbon. SP 300 PYE UNI CAM Infrared Spectrophotometer was used to record the Infrared spectra with anhydrous KBr pellets.

Mass spectra and LCMS procedure

Bruker MicroTOF QII mass spectrometer (Bruker Daltonik, Bremen, Germany) machine calibrated with sodium formate (internal calibrant) was used to record the mass spectra. KdScientific automated pump with flow rate 180 $\mu\text{L}/\text{h}$ was used to inject the sample to electrospray ionization source at a concentration 50 μM in acetonitrile-water-formic acid (7:2.9:0.1). Dry N_2 gas heated at 180 $^\circ\text{C}$ was used for the desolvation of the injected solution. Optimisation of the various parameters of the mass spectrometer was done for obtaining a maximum ion abundance. Specifically, the capillary voltage was adjusted at 4500 V and the vacuum was sustained at $3\text{-}4 \times 10^{-7}$ mbar. For LC-MS, Dionex Ultimate 3000 system was coupled to the mass spectrometer. Chirobiotic[®] T 10 μm chiral HPLC column (25 cm x 4.6 mm) was used for the analysis. A solution of Acetonitrile in water (1:1) was used as eluent. 2 μL of sample (injection volume) was loaded to the column. The flow rate was maintained 0.2 ml and the absorbance was set at 200, 220 and 254 nanometers. Sodium formate was used as internal calibrant.

$^1\text{H}/^{13}\text{C}/\text{DEPT-135}$ NMR data of the test compounds:

[4-(3-Methyl-indol-1-ylmethyl)-[1,2,3]triazol-1-yl]-acetic acid (3a). The compound (2) was dissolved in 9:1 solution of EtOH:H₂O and allowed to stir till a clear solution is obtained. To this clear solution was immediately added 0.02 mmol of $\text{CuSO}_4 \cdot 5\text{H}_2\text{O}$. After waiting for another 5 minutes, 0.05 mmol sodium ascorbate was added to the reaction mixture and the contents were allowed to stir for 10 minutes. Finally, the azidoacetic acid (i) synthesized in scheme 1, was added to the reaction mixture. The reaction was monitored by thin layer chromatography technique. The reaction generally completes in 8 hours to give a yellow coloured powdery solid, mp 90-95 $^\circ\text{C}$, 80% yield, IR (KBr) ν_{max} 3380 (NH), 1610 ($\text{C}=\text{O}$) cm^{-1} ; ^1H NMR (CDCl_3 , 500 MHz): δ = 9.42 (s, 1H, COOH) 2.30 7.83-8.401(m, 3H, ArH), 7.39-7.43 (m, 2H, ArH), 4.20 (s, 2H, CH_2), 4.01 (s, 2H, CH_2), 2.25 (s, 3H, CH_3), (^{13}C NMR normal/DEPT-135) (125 MHz, CDCl_3): 181.27 ($\text{C}=\text{O}$), 138.66 (C), 134.24 (C), 128.11(CH), 127.93 (CH), 125.74 (C), 122.87(ArCH), 115.79 (ArCH), 113.30 (ArCH), 27.48 (-ve, CH_2), 21.98 (-ve, CH_2), 21.04 (CH_3); ESI-MS (HRMS) calculated for $\text{C}_{14}\text{H}_{14}\text{N}_2\text{O}_4$ 271.1187. Found m/z 271.1181([M+H]⁺).

3-(1H-Indol-3-yl)-2-[4-(3-methyl-indol-1-ylmethyl)-[1,2,3]triazol-1-yl]-propionic acid (3b). The compound (2) was dissolved in 9:1 solution of EtOH:H₂O and allowed to stir till a clear solution is obtained. To this clear solution was immediately added 0.02 mmol of $\text{CuSO}_4 \cdot 5\text{H}_2\text{O}$. After waiting for another 5 minutes, 0.05 mmol sodium ascorbate was added to the reaction mixture and the contents were allowed to stir for 10 minutes. Lastly, the 2-Azido-3-(1H-indol-3-yl)-propionic acid (ii) obtained in scheme 2, was added to the reaction mixture. The reaction was monitored by thin layer chromatography technique. The reaction generally completes in 8 hours to give a bright yellow coloured solid, mp 137-142 $^\circ\text{C}$, 69% yield, $[\alpha]_{\text{D}}^{25} = +124$ (c 0.01, MeOH), IR (KBr) ν_{max} 3390 (NH), 3191 (NH), 1632 ($\text{C}=\text{O}$) cm^{-1} ; ^1H NMR (500 MHz, DMSO d_6): δ = 8.16 (s, 1H, COOH), 6.70-7.50 (m, 9H, ArH), 6.70 (s, 1H, ArH), 4.51(t, J = 10 Hz, 1H, ArH, 1H, CH), 4.20 (s, 2H, CH_2), 3.45-3.63 (m, 2H, CH_2), 2.26 (s, 3H, CH_3); (^{13}C NMR normal/DEPT- 135) (125 MHz, CDCl_3): 179.98 ($\text{C}=\text{O}$), 136.70 (C), 134.40 (C), 130.92 (ArCH), 127.47 (ArCH), 125.41 (ArCH), 123.83 (ArCH), 121.60 (ArCH), 119.44 (ArCH), 118.00 (ArCH), 114.60 (ArCH), 111.98 (ArCH), 109.85 (C), 107.07 (C), 53.15 (CH), 26.61 (CH_2), 25.45 (CH_2), 21.58 (CH_3); ESI-MS (HRMS) calculated for $\text{C}_{23}\text{H}_{21}\text{N}_5\text{O}_2$ 400.1775. Found m/z 400.1700 [M+H]⁺.

2-[4-(3-Methyl-indol-1-ylmethyl)-[1,2,3]triazol-1-yl]-3-(1H-[1,2,3]triazol-4-yl)- propionic acid (3c). The compound (2) was dissolved in 9:1 solution of EtOH:H₂O and allowed to stir till a clear solution is obtained. To this clear solution was immediately added 0.02 mmol of $\text{CuSO}_4 \cdot 5\text{H}_2\text{O}$. After waiting for another 5 minutes, 0.05 mmol sodium ascorbate was added to the reaction mixture and the contents were allowed to stir for 10 minutes. Finally, the 2-Azido-3-(1H-imidazol-4-yl)-propionic acid (iii) procured in scheme 2, was added to the reaction mixture. The reaction was monitored by thin layer chromatography technique. The reaction generally completes in 8 hours too get a grey coloured solid, mp 121-126 $^\circ\text{C}$, 65% yield, $[\alpha]_{\text{D}}^{25} = +154$ (c0.01, MeOH), IR (KBr) ν_{max} 3299 (NH), 1600 ($\text{C}=\text{O}$) cm^{-1} ; ^1H NMR (CDCl_3 , 500 MHz): δ = 9.21 (s, 1H, COOH), 7.45-7.89 (m, 2H, ArH), 6.83-7.39 (m, 5H, ArH), 6.82 (s, 1H, ArH), 4.41 (t, J= 15 Hz, 1H, CH), 4.20 (s, 2H, CH_2), 3.33 (d, J= 10 Hz, 2H, CH_2), 2.27 (s, 3H, CH_3); (^{13}C NMR normal/DEPT- 135) (125 MHz, CDCl_3): δ C = 171.97 (COOH), 148.95 (ArCH), 138.64 (ArCH), 136.73 (ArCH), 127.49 (C), 125.96 (C), 121.59 (ArCH), 119.05 (ArCH), 116.64 (ArCH), 111.04 (ArCH), 107.08 (C), 54.27 (CH), 28.61 (CH_2), 25.46 (CH_2), 21.24 (CH_3); ESI-MS (HRMS) calculated for $\text{C}_{17}\text{H}_{17}\text{N}_7\text{O}_2$ 352.1524 Found m/z 352.1523 [M+H]⁺.

2-[4-(3-Methyl-indol-1-ylmethyl)-[1,2,3]triazol-1-yl]-pentanedioic acid (3d). The compound (2) was dissolved in 9:1 solution of EtOH:H₂O and allowed to stir till a clear solution is obtained. To this clear solution was immediately

added 0.02 mmol of CuSO₄.5H₂O. After waiting for another 5 minutes, 0.05 mmol sodium ascorbate was added to the reaction mixture and the contents were allowed to stir for 10 minutes. Finally, the 2-Azido-4-carboxyheptanedioic acid (iv) obtained in scheme 2, was added to the reaction mixture. The reaction was monitored by thin layer chromatography technique. The reaction generally completes in 8 hours to give a dark brown solid, mp 111-116 °C, 85% yield, $[\alpha]_D^{25} = +23$ (c 0.01, MeOH); IR (KBr) ν_{\max} 3412 (NH), 1574 (C=O) cm⁻¹; ¹H NMR (DMSO d₆, 500 MHz): δ = 10.41 (s, 1H, COOH), 9.81 (s, 1H, COOH), 7.91 (s, 1H, ArH), 7.53-7.82 (m, 2H, ArH), 7.22-7.39 (m, 3H, ArH), 4.39 (t, J = 5 Hz, 1H, CH), 4.20 (s, 2H, CH₂), 2.36-2.72 (m, 2H x CH₂), 2.27 (s, 3H, CH₃); (¹³C NMR normal/DEPT- 135) (125 MHz, DMSO d₆): δ C = 180.03 (COOH), 179.55 (COOH), 145.55 (ArCH), 134.66 (ArCH), 130.93 (C), 127.14 (C), 123.86 (C), 122.03 (ArCH), 116.85 (C), 114.55 (ArCH), 113.53 (ArCH), 109.82 (C), 53.78 (CH), 29.57 (CH₂, -ve), 25.78 (CH₂, -ve), 25.06 (CH₂, -ve), 21.53 (CH₃); ESI-MS (HRMS) calculated for C₁₇H₁₈N₄O₄ 342.1408 Found m/z 352.1410 [M+H]⁺.

3-(4-Hydroxy-phenyl)-2-[4-(3-methyl-indol-1-ylmethyl)-[1,2,3]triazol-1-yl]- propionic acid (3e). The compound (2) was dissolved in 9:1 solution of EtOH:H₂O and allowed to stir till a clear solution is obtained. To this clear solution was immediately added 0.02 mmol of CuSO₄.5H₂O. After waiting for another 5 minutes, 0.05 mmol sodium ascorbate was added to the reaction mixture and the contents were allowed to stir for 10 minutes. Finally, the 2-Azido-4-(4-hydroxy-benzyl)-pentanedioic acid (v) was added to the reaction mixture. The reaction was monitored by thin layer chromatography technique. The reaction generally completes in 8 hours to get a brown solid, mp 145-153 °C, 78% yield, $[\alpha]_D^{25} = +35$ (c 0.01, MeOH); IR (KBr) ν_{\max} ; 3315 (NH), 1701 (C=O) cm⁻¹; ¹H NMR (CDCl₃, 500 MHz): δ = 9.15 (s, 1H, COOH), 7.73-7.94 (m, 2H, ArH), 7.53-7.63 (m, 1H, ArH), 6.87-7.52 (m, 3H, ArH), 4.42 (t, J = 15 Hz, 1H, CH), 4.35 (s, 2H, CH₂), 3.33 (d, J = 3 Hz, 2H, CH₂), 2.20 (s, 3H, CH₃); (¹³C NMR normal/DEPT- 135) (125 MHz, CDCl₃): δ C = 173.53 (COOH), 145.30 (C), 144.82 (C), 135.27 (ArCH), 134.94 (ArCH), 130.75 (C), 129.87 (ArCH), 126.81 (ArCH), 126.33 (ArCH), 124.55 (ArCH), 123.30 (ArCH), 121.36 (ArCH), 113.30 (ArCH), 109.04 (C), 107.24 (C), 58.75 (CH), 29.29 (CH₂, -ve), 25.19 (CH₂, -ve), 21.53 (CH₃); ESI-MS (HRMS) calculated for C₂₁H₂₀N₄O₃ 377.1615 Found m/z 377.1615 [M+H]⁺.

Pharmacology

MIC80

The antifungal activity of the test compounds **3 (a-e)** was evaluated by performing the MIC80 assay. Ten successive dilutions of the test compounds were prepared alongwith the solvent control with yeast extract peptone dextrose (YEPD) medium in a 96-well plate. The overnight grown cells of the BWP 17 strain of the fungus *Candida Albicans* were resuspended in a 0.9% saline solution to give a resultant optical density of 0.1 at a wavelength of 570 nm (OD570) These cells were further diluted 100 times in SDM medium. 100 μ l of these diluted cells were added to each of the wells in the 96 well plate except the column 11 which was taken as media control to examine any kind of contamination while performing the assay. Column 12 which was taken as positive control contained the fungal cells only in the growth medium. The assay was performed in duplicates. After preparing the plate it was incubated at a temperature of 30 °C for a duration of 48 hours. Afterwards, the monitoring of OD570 of the cells was done using a micro plate reader. The percent inhibition of the fungal growth in the presence of test compounds **3 (a-e)** was determined with respect to the positive control by deducting the percent inhibition due to the solvent control. The MIC80 values were thus calculated by plotting a graph between percent inhibition versus concentration of the test compound using the Graphpad Prism software. The percent inhibition of the fungi was determined at eight different concentrations of the test compounds **3 (a-e)**. The Compounds **3b**, **3c**, and **3d** containing tryptophan, histidine and glutamic acid substituents appended to the indole through a triazole linker were found to be most effective against CABWP-17 with MIC80 of 50.2, 78.81, and 188.05 μ g/ml, respectively. Compounds **3a** and **3e** with glycine and tyrosine substituents appended to indole via azole linker were found to be least effective with MIC80 values in the range of 617.90-752.78 μ g/ml.

Table 1. MIC80 values and experimentally determined hydrophilicity index of compounds 3 a-e for CA BWP17 strain


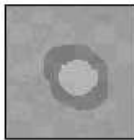






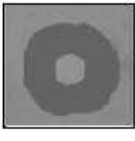

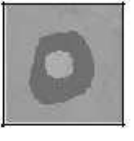


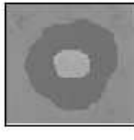






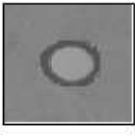

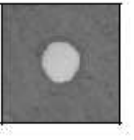







Compound	MIC80 (μ g/ml)	ClogP (experimental)
3a	617.90	3.45
3b	50.2	1.89
3c	78.81	1.78
3d	188.05	-1.01
3e	752.78	1.99

Disc diffusion assay

The disc diffusion assay was performed to further explore the antifungal efficacy of the test compounds. Taking YEPD as the culture medium the cells of the BWP17 strain of *Candida albicans* were grown overnight at 30°C. A 90 mm diameter YEPD-agar plate was then prepared and 10⁷ cells were spread on it. This plate was then dried for around 10-15 minutes. Small filter paper discs (5 mm diameter) were prepared, sterilized and placed on this plate with the help of clean forceps. A known concentration of each test compound dissolved in an appropriate solvent

was spotted on the paper disc. Similar procedure was repeated with the solvent controls. After incubating the assay plates for 48 hours at 30 °C the growth of the fungal cells at the periphery of the disc was checked. The diameter of the circle around the disc with no fungal growth corresponds to inhibition zone. In order to maintain an effective concentration of the test compounds in the culture medium the disc diffusion assay was performed at a slightly superior concentration than the MIC80 values. The images of inhibition zone (mm) of fungal culture in the presence of test compounds compared with the corresponding solvent are shown in table 2. The zone of inhibition was highest for the Compound **3b** followed by the compound **3c** at concentrations 100 µg/ml and 50 µg/ml. Whereas, the zone of inhibition was much narrower in case of the compounds **3a**, **3d** and **3e**.

Table 2. Diameter (in mm) of the zone of inhibition in the disc diffusion assay of highly effective compounds against CA BWP17

	Concentration of test compounds					
	solvent	100 µg/ml	50 µg/ml	25 µg/ml	12.5 µg/ml	6.125 µg/ml
3a						
3b						
3c						
3d						
3e						

Spot assay

The results of MIC80 evaluation and the disc diffusion assay led us to select the compounds **3b** and **3c** for performing the spot assay. A known concentration of the compounds **3b** and **3c** was added to the medium containing YEPD and agar. Solvent control plates were also prepared. The plates with naked YEPD-agar medium served as positive controls. The fungal cells growing overnight on the YEPD medium at a temperature of 30 °C served as primary culture medium. The secondary culture was prepared from a 4% inoculum of the primary culture medium. From the secondary culture were prepared five serial dilutions of cell suspension with 0.9% saline solution to give a resultant OD of 0.2, 0.04, 0.008, 0.00016 and 0.00032 at 570 nm. A 5 µl volume from each cell suspension was added to all the plates. After incubating the plates at 30 °C their images were recorded after 24 hours and 36 hours respectively. The spot assay indicates an impaired fungal growth in the presence of the compounds **3b** and **3c**. The results were quite comparable with the commercially used drug ketoconazole which showed a similar growth inhibition pattern after 24 hours and 36 hours of treatment.

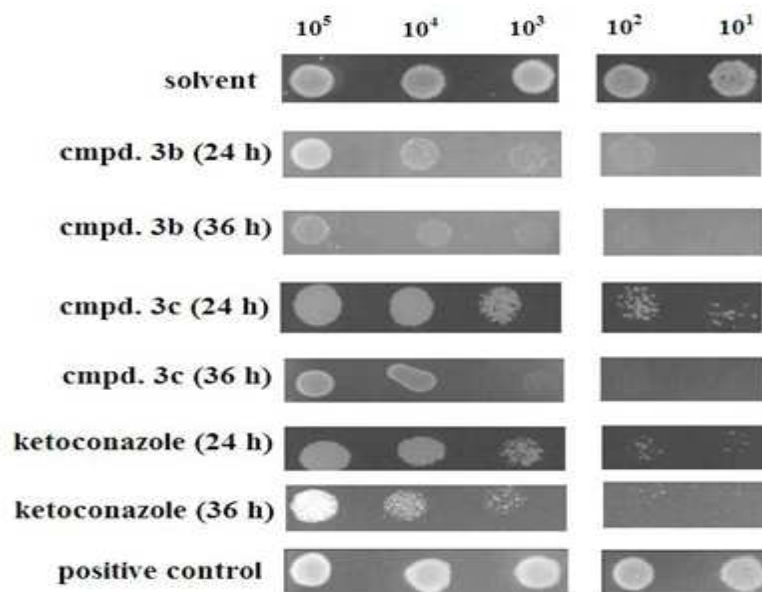


Figure 1. Spot assay for CA BWP17 in the presence of active compounds 3b and 3c

SEM analysis

The morphological changes occurring in the fungal cells in the presence of test compounds was monitored through the Scanning Electron Microscopy. The CA BWP17 cells were grown in the presence of the most active compound **3b**. The cells were pelleted, washed and then fixed into 2.5% glutaraldehyde and 2% Paraformaldehyde in 0.1M phosphate buffer at 4 °C. In order to get a superior staining of the intracellular membranes, the secondary fixing of the cells was done in Osmium tetroxide (OsO₄). The dehydration of the sample was done by sequentially treating the cells with increased concentrations of acetone from 50% to 95%. After drying the sample with critical point drier it was mounted on the stub SCM. Spatter coater was used for the gold coating of cells. The cells were viewed under the Carl Zeiss Microscope EVO40 at 30 KV. The morphological changes were noticeable on the fungal membrane in the presence of compound **3b** which got augmented with the passage of time. The SEM investigations clearly indicate that the compound **3b** targets the fungal membrane (figure 2).

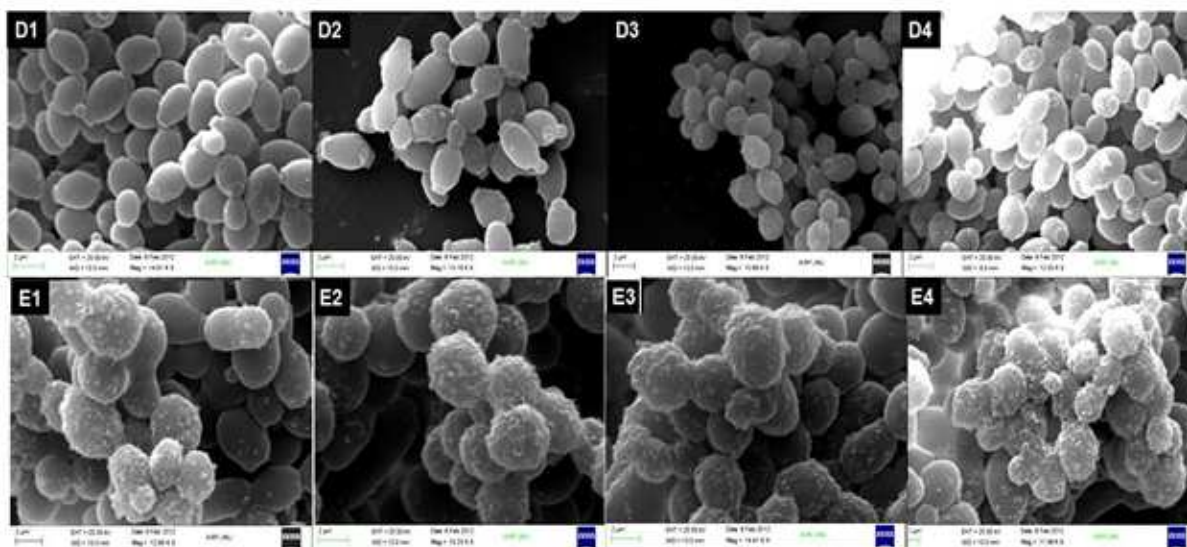


Figure 2. SEM images of CA BWP17 culture grown in the presence and absence of compound

3b at different time intervals. D1, D2, D3 and D4 indicate the growth of cells under the optimum conditions in the absence of the compound after 2 h, 6 h, 10 h and 16 h respectively. E1, E2, E3 and E4 indicate the growth of the cells in the presence of compound **3b** after 2 h, 6 h, 10 h and 16 h respectively.

GC analysis

To explore the inhibition of the enzyme lanosterol 14- α demethylase in the presence of the test compounds, the levels of ergosterol and lanosterol in the BWP17 cells treated with the compound 3b were assessed using GC-MS. The cells treated with compound 3b showed a depleted ergosterol content compared with the untreated cells. The levels of lanosterol on the other hand were found to increase. This clearly indicates an impaired functioning of the lanosterol 14- α demethylase enzyme in the presence of the compound 3b due to which much of the substrate lanosterol is not metabolized to ergosterol and hence it starts building up in the cells treated with the test compound.

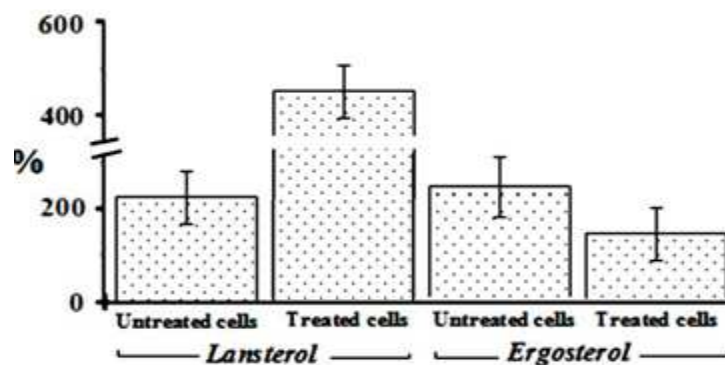


Figure 3. The figure indicates depleted levels of ergosterol in the cells treated with compound 3b

The level of lanosterol in the same cells was found to increase. The bars represent averages of three independent assays and the standard deviations are represented by error bars.

Docking Analysis

The inhibition of the enzyme activity by the test compounds was justified by docking experiments wherein the test compounds were docked in the active site of the enzyme lanosterol 14- α demethylase by using ArgusLab.exe software. Both the stereoisomers of the test compounds were docked and it was found that the 'R' stereoisomer showed a better docking profile in the enzyme active site. This was in accordance with the experimental data where the D-amino acids were selected for synthesizing the compounds. The numerous hydrogen bonding interactions were observed for the compounds 3b and 3c which interact with the active site residues principally through N atom of the triazole ring. The 'R' stereoisomer of compound 3b interacts with the residues Y576, Y590 and F597 via COOH and N atom of the nitrogen heterocycle by hydrogen bonding interactions of length 1.23 Å, 1.22 Å, 1.77 Å and 1.69 Å respectively. Similarly, the R' stereoisomer of the compound 3c interacts with the residues Y576, F597, G752 and A756 via N atom of the triazole ring and COOH functional group with hydrogen bonding interactions of length 2.22 Å, 1.88 Å, 1.72 Å and 1.61 Å respectively.

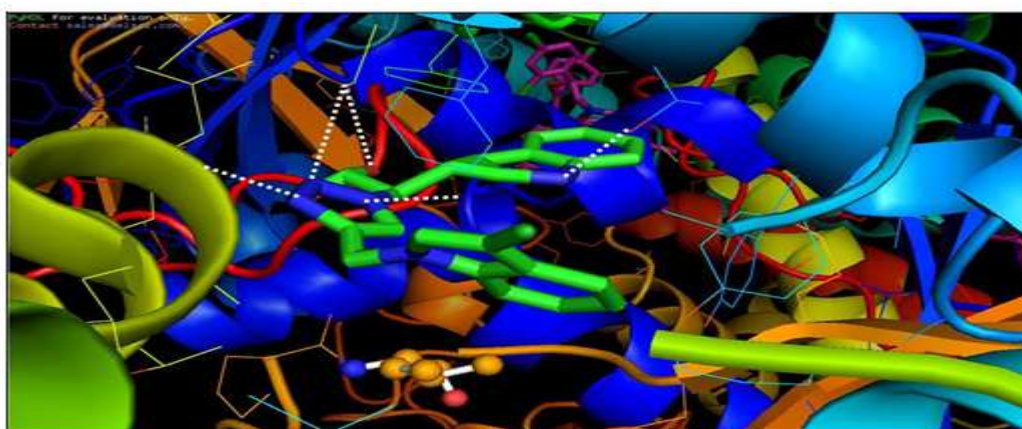
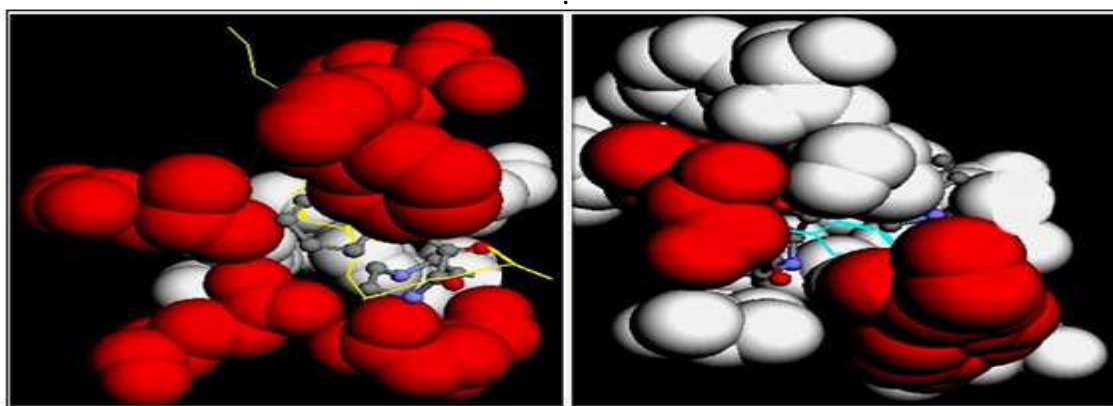


Figure 4. The crystal coordinates of the enzyme lanosterol 14- α demethylase (pdb ID 3jus) in association with the compound 3b. The white dotted lines represent the H-bonding interactions between compound and the amino acid residues in the active site of the enzyme. The interaction are principally through the triazole nucleus and COOH functional group of the compound

Table 3. The docking characteristics of the different stereoisomers of the compounds in the active site of the enzyme lanosterol 14- α demethylase

Compd.	Docking Score (Kcal/mol)	Hydrogen bond interactions			
3a	-10.45	Y576 (2.44 Å), COOH	-	-	-
3b (R)	-20.21	Y576 (1.23 Å), COOH	Y576 (1.22 Å), N (azole)	Y590 (1.77 Å), N (azole)	F597 (1.69 Å), NH (indole)
3b (S)	-12.34	G752 (3.11 Å), COOH	H372 (2.61 Å), COOH	A756 (1.99 Å), N (azole)	-
3c (R)	-18.70	Y576 (2.22 Å), N (azole)	F597 (1.88 Å), N (imidazole)	G752 (1.72 Å), N (azole)	A756 (1.61 Å), COOH
3c (S)	-10.78	G752 (3.61 Å), COOH	Q557 (2.91 Å), COOH	-	-
3d (R)	-15.41	Y576 (2.45 Å), COOH	A756 (2.22 Å), COOH	Y590 (1.99 Å), CONH2	-
3d (S)	-8.98	F597 (2.99 Å), COOH	Y576 (3.59 Å), COOH	-	-
3e (R)	-14.01	A756 (2.10 Å), CONH2	F597 (2.36 Å), COOH	Y590 (1.89 Å), COOH	-
3e (S)	-8.29	F597 (3.14 Å), COOH	-	-	-
ketoconazole	-16.81	Y576 (1.45 Å), N (imidazole)	F597 (1.66 Å), N (imidazole)	-	-

Figure 5. CPK model for showing the placement of the compound 3b in the active site of the enzyme lanosterol 14- α demethylase in the presence of the natural inhibitor shown in yellow and blue wireframe

Compound	Energy minimized structure	Nviolations	Volume	ClogP
3a		0	243.74	3.45
3b		1	343.56	1.89
3c		0	324.56	1.78
3d		1	332.45	-1.01
3e		0	299.90	1.99

Figure 6. Lipinski Parameters for the compounds. ClogP – hydrophilicity index, Nviolations – number of violations from Lipinski parameters

Study of Influx – Efflux of R6G in CA BWP17 in presence of compounds:

Several of the commercially available antifungal drugs are subjected to the cellular efflux pumps. To check the vulnerability of the test compounds **3b** and **3c** towards the fungal efflux pumps, the R6G influx-efflux assay was performed. R6G is a substrate of fungal efflux pumps. It was envisioned that if there is a competition between test compounds and R6G to interact with efflux pump, the transportation of R6G out of the cell will be debarred. Conversely, if there is efflux of R6G in presence of compounds it would indicate that there is no interaction between compound and efflux pump protein. Influx-efflux of R6G in fungal cells was monitored by recording the absorbance and fluorescence of extracellular fluid. CA BWP17 primary/ secondary culture was grown as described earlier. After centrifugation the cells were taken in PBS solution or non-growing medium. The PBS solution was now divided into three portions. One part was taken positive control, second part was considered as solvent control and the third part was treated with the test compounds. R6G was added to all the three portions. The assay was performed in triplicates. An increase in R6G influx was observed in positive control and solvent control experiments while in presence of compound **3b**, most of R6G entered the cell immediately and no change in R6G influx was observed with time. Similarly, the initial efflux of R6G in the presence of compound **3b** was found to be much faster compared to positive control and solvent control. Both influx as well as efflux assays were pointing towards rupture of the fungal membrane in presence of compound **3b** making free passage for influx and efflux of R6G. Similar trends were observed during the efflux of R6G in presence of compound **3b**.

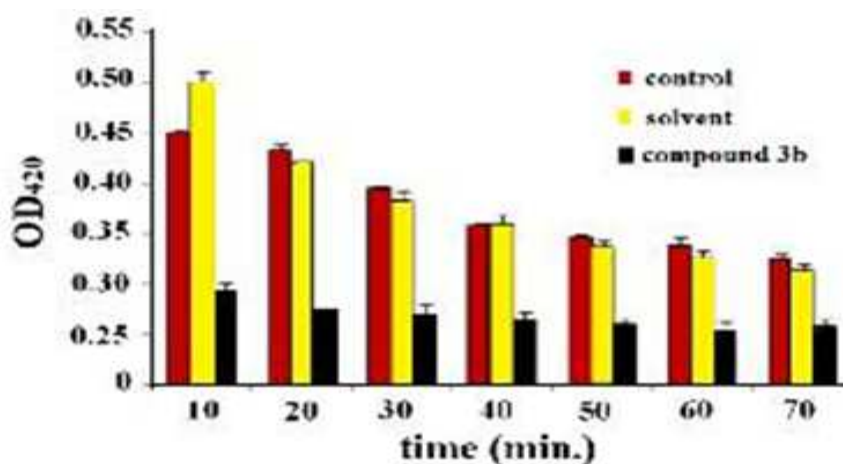


Figure 7. R6G influx in the presence of compound 3b

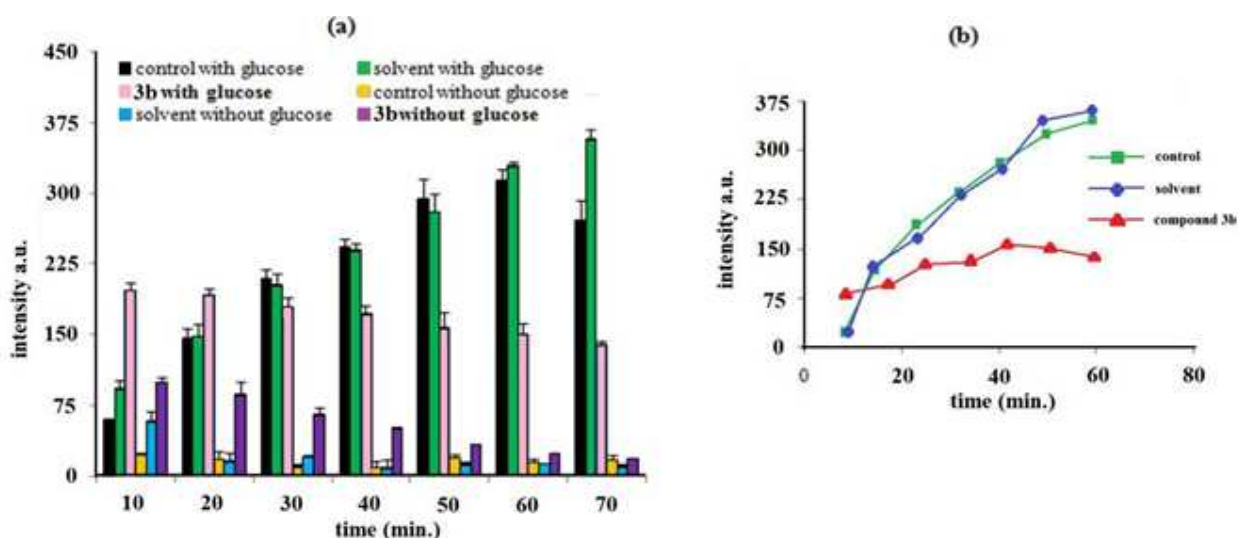


Figure 8. (a) R6G efflux in the presence of compound 3b. (b) Differentiation between active and passive efflux

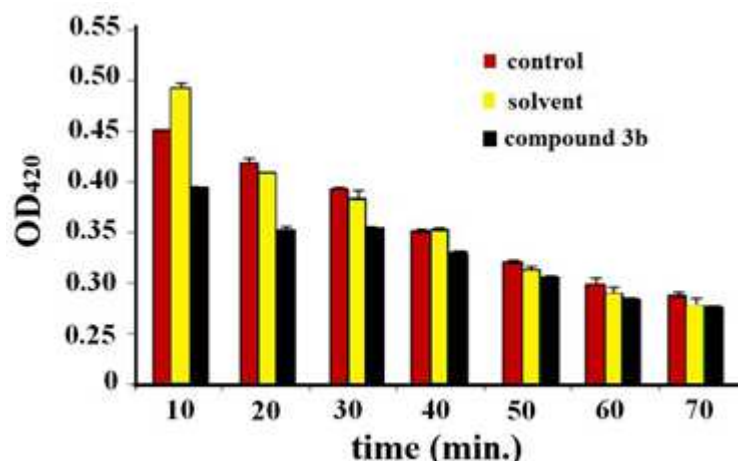


Figure 9. Influx of R6G in the presence of compound 3c

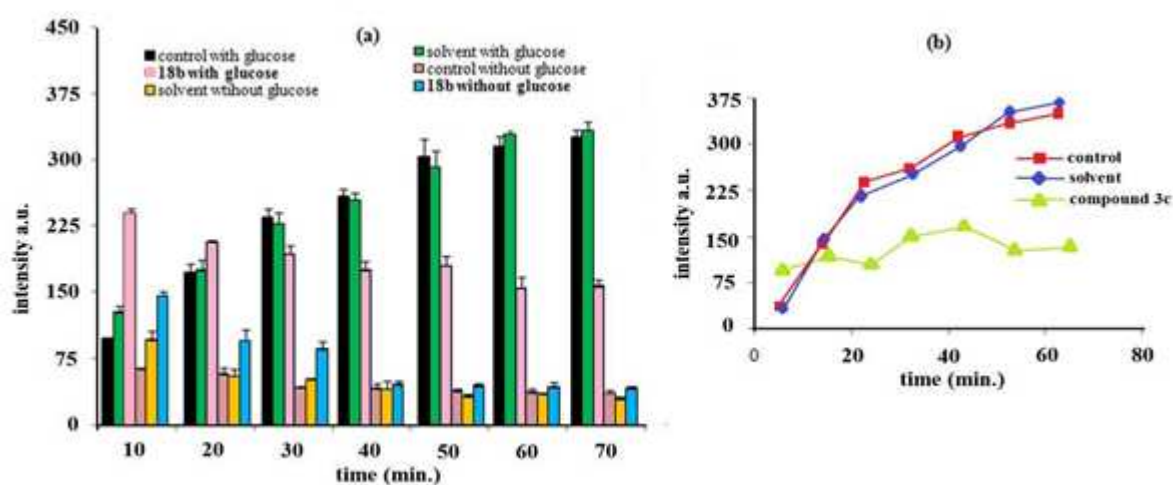


Figure 10. (a) R6G efflux in the presence of compound 3c. (b) differentiation between active and passive efflux

CONCLUSION

As conclusion, the amino acid tethered triazoles were identified with momentous antifungal activities. The minimum inhibitory concentration assay revealed that the compounds **3b** and **3c** with tryptophan and histidine substituents were most active inhibitors of the BWP 17 strain of fungi candida albicans. The observation was further confirmed by performing the spot assay and disc diffusion assays. The GC analysis of the metabolites of the fungal membrane revealed an elevated concentrations of the ergosterol which led us to conclude that the test compounds actively target the enzyme lanosterol 14- α demethylase. The docking of the test compounds in the active site of the enzyme further strengthened this observation. The study of the R6G efflux/ influx pumps in the fungal membrane revealed that the test compounds adversely influence the membrane integrity leading to the creation of a free passage for the entry and exit of metabolites across the membrane. Overall, highly efficient antifungal leads with membrane lysing properties were recognized.

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