



Synthesis, antimicrobial evaluation and docking studies of fluorinated imine linked 1,2,3-triazoles

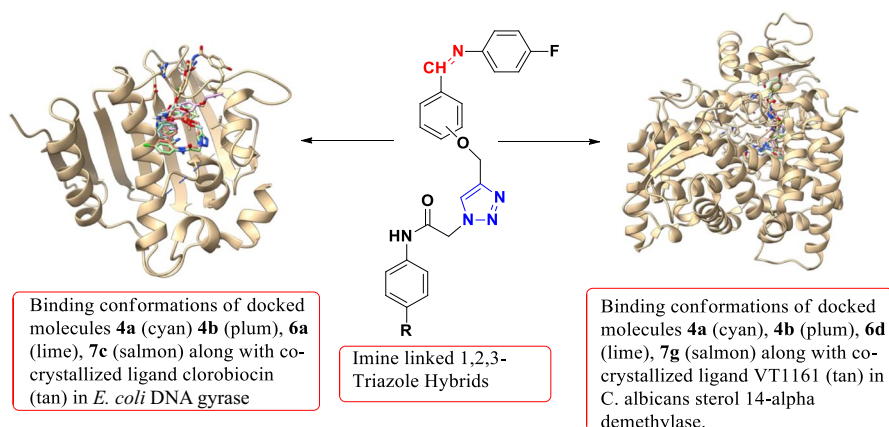
Aman Kumar¹ · Kashmiri Lal¹ · Nisha Poonia¹ · Ashwani Kumar² · Anil Kumar³

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Abstract

A diverse series of imine linked 1,2,3-triazole hybrids has been synthesized via Cu(I)-promoted click reaction, with an aim to develop some new antimicrobial molecules. The structural characterization of the synthesized triazole hybrids was accomplished using various spectral techniques like ¹H NMR, ¹³C NMR, FTIR and HRMS. Among the synthesized hybrids, **7d** exhibited highest antimicrobial efficacy against *R. oryzae* and *S. aureus* with MIC of 0.0123 μmol/mL and 0.0061 μmol/mL, respectively. Docking studies of the terminal alkynes (**4a**, **4b**) and triazole hybrids (**6a**, **6d**, **7c**, **7g**) showing high potency towards *E. coli* DNA gyrase and *C. albicans* sterol 14- α demethylase were also undertaken to understand the binding behaviour.

Graphical abstract



Keywords Imine · Click chemistry · 1,2,3-triazoles · Antimicrobial activity · Docking

✉ Kashmiri Lal
klal_iitd@yahoo.com

Extended author information available on the last page of the article

Introduction

Despite the availability of various antimicrobial drugs in the market, microbial infections are a foremost cause of deaths worldwide [1]. Just like Covid-19, microbial resistance is a threat to the health security, it may become next global health emergency, if necessary actions are not taken to prevent the microbial resistance [2, 3]. According to the annual report of GARDP, 1.2 million people die every year due to the infections caused by drug resistant microbes. Europe and US are facing a financial loss of 13.5 billion dollars every year due to the hospital associated infections and also by 2050, approximate 28.3 million people could be pushed into the extreme poverty (https://gardp.org/?gclid=Cj0KCQjwnoqLBhD4ARIsAL5JedK4wcgihTB_tiMNA5mWXcrz6y0nxXmDl-dJSHJWejvguu6L3cIG8EQaAmFbEALw_wcB (accessed on 31.01.2022)). The situation is worsened by the inappropriate use of the drugs available in the market and it results into the diminishing drug's efficacy [4]. The main cause of drug resistant development in case of antibiotics is the formation of biofilms by bacteria [5]. To overcome this worsened situation, the development of microbial drugs with novel mode of action is highly desired. Scientists are using various approaches to develop antimicrobial drugs for saving human life from the health emergencies of future and molecular hybridization approach is evolved as a most promising method to achieve this target [6, 7].

A large number of natural and synthetic pharmacophores have been reported as antimicrobial agents [8]. In the last two decades, 1,2,3-triazoles have emerged as the one of the most promising pharmacophores exhibiting important biological activities like antibacterial [9, 10], antifungal [11], anti-Alzheimer [12], anti-cancer [13], anti-human immunodeficiency virus (anti-HIV) [14], antidiabetic [15], antimalarial [16], antitubercular [17], anti-inflammatory activity [18]. The hybridization of 1,2,3-triazole unit with several other pharmacophores like Schiff bases [19], chalcones [20], indole [21], pyrazolines [22], semicarbazone [23], urea [24], oxazolones [25] has empowered to develop various molecular hybrid displaying excellent inhibition against a variety of microorganisms. The high stability of 1,2,3-triazoles against acidic and basic hydrolysis and inertness towards reducing and oxidizing agents motivated scientists to exploit 1,2,3-triazole hybrids for bioactivity evaluation [26].

Imine unit also known as azomethine is also an important class of pharmacophores because of their ease of synthesis from carbonyl and amino derivatives as well as their abilities to exhibit various interactions with different biological targets. Compounds containing imine unit display a wide range of pharmacological properties including antibacterial [27], antifungal [28], anti-HIV [29], anti-cancer [30], anti-inflammatory [31], antimalarial [32], antitubercular [33]. Imines can easily be obtained by nucleophilic addition of carbonyl compounds and amino compounds [34, 35]. As reported in the literature, Gurjaspreet et al. synthesized schiff base linked triazole silatranes and evaluated their antimicrobial potential [36, 37]. Yuqin et al. derived paeonol schiff base containing 1,2,3-triazole and assessed their antilung adenocarcinoma potential [38]. Therefore, inspired from

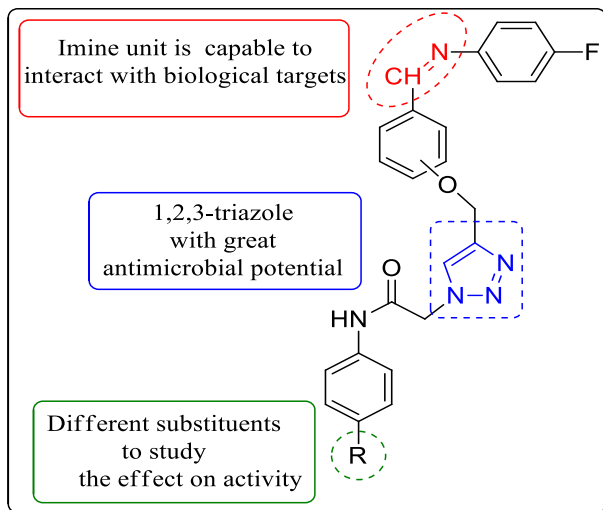


Fig. 1 Designed imine linked 1,2,3-triazole hybrids

bioactivity profile of imine, triazole and amide units it was envisioned to synthesize some derivatives containing these units for the antimicrobial screening (Fig. 1). Herein, we have outlined the synthesis of various imine linked 1,2,3-triazoles as antimicrobial compounds. The docking studies of potent triazole hybrids against *C. albicans* sterol 14- α demethylase and *E. coli* DNA gyrase was carried out and obtained results are in support with the *in vitro* antimicrobial screening results.

Materials and methods

General

Starting materials employed for the synthesis of triazole hybrids were obtained from the commercial sellers, open capillary tube method was employed to record the melting points of the compounds and are reported as such. TLC plates were used to visualize the progression of the reaction. IR spectral data was recorded on Shimadzu IR Affinity FTIR spectrophotometer. Bruker Avance III 400 nano bay spectrometer operating at 400 and 100 MHz and Bruker Avance NMR spectrometer operating at 500 and 125 MHz in deuterated (DMSO-*d*₆) using tetramethyl silane (TMS) as internal standard (chemical shift in ppm) was used to record ¹H and ¹³C NMR.

General procedure for the synthesis of imine linked alkynes (**4a**, **4b**)

To synthesize imines, an equimolar amount of *m/p*-hydroxy benzaldehydes (8.2 mmol) and *p*-fluoro aniline (8.2 mmol) was dissolved in 20 mL of dry ethanol and refluxed for 6 h at 80 °C in the presence of catalytic amount of HCl. When reaction was accomplished, obtained solid was washed with ethanol and then recrystallized from ethanol. Further, these synthesized imines (7.0 mmol) were propargylated with propargyl bromide (80% in toluene) (8.4 mmol) on refluxing in acetone in presence of K₂CO₃ (21 mmol). Propargylation was completed after continue refluxing for 7 h. Then ice-cold water was poured in the reaction mixture, precipitates were obtained in case of **4b** alkyne, which was further washed with ice-cold water and recrystallized in ethanol. But in case of **4a** alkyne on addition of water in reaction mixture, no precipitates obtained and it was extracted with ethyl acetate and **4a** was obtained from organic layer after evaporating solvent under vacuum.

General procedure for the synthesis of imine linked 1,2,3-triazoles (**6a–6g**, **7a–7g**)

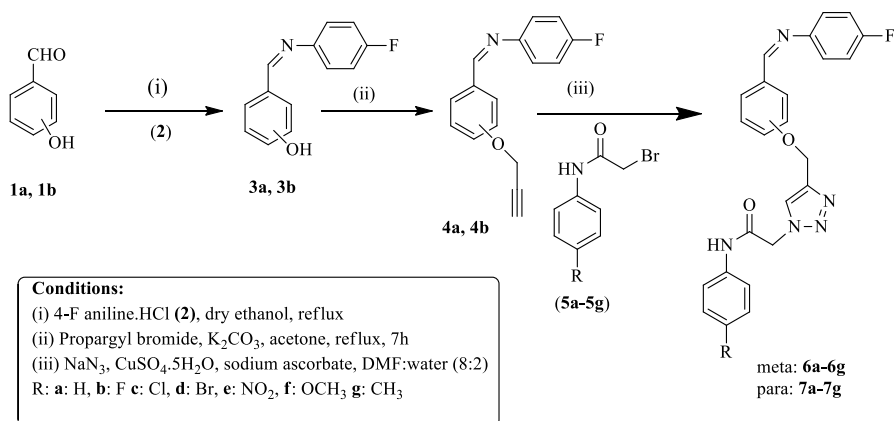
To synthesize imine linked 1,2,3-triazole hybrids, *N*-phenyl acetamide bromide derivatives **5a–5g** (1.0 mmol), imine linked alkynes **4a**, **4b** (1.0 mmol), sodium azide (3.0 mmol) in DMF: water (8:2) were stirred for 6 h at 50 °C, after the addition of copper sulphate pentahydrate (10 mol%) and sodium ascorbate (20 mol%). Reaction was monitored by using TLC under the UV lamp, on completion reaction mixture was cooled and diluted with ice-cold ammonium chloride (aq.): ammonia solution (9: 1 v/v). Solid thus obtained was filtered and washed with cold water to furnish imine linked 1,2,3-triazoles (**6a–6g**, **7a–7g**).

Pharmacology

The preliminary *in vitro* pharmacological potential of the synthesized imine linked 1,2,3-triazole hybrids (**6a–6g**, **7a–7g**) as well as their precursor terminal alkynes (**4a**, **4b**) were studied against several microbial strains using serial dilution method [39].

Docking details

The three-dimensional structures of molecules were drawn and developed with Marvin Sketch [40] and docking simulations were carried out using Autodock Vina software [41] in to protein (DNA gyrase of *E. coli* and sterol 14- α -demethylase of *C. albicans*) with PDB ID: 5TZ1 downloaded from protein data bank. The protein preparation was carried out in UCSF Chimera [42]. The protocols followed while



Scheme 1 Synthesis of imine linked 1, 2, 3-triazole hybrids (6a–6g, 7a–7g)

carrying out docking simulations were same as described in literature [43]. The visualization of the docking results was done using Discovery Studio Visualizer [44] and Chimera X [45].

Results and discussion

Chemistry

In continuation to our work on synthesis and bio-evaluation of 1,2,3-triazoles, a library of new imine linked 1,2,3-triazoles were synthesized. Synthetic protocol used to achieve imine appended 1,2,3-triazoles involved three steps. In first step *m/p*-hydroxy benzaldehydes (**1a–1b**) were condensed with *p*-fluoro aniline in dry ethanol with few drops of HCl as catalyst under refluxing to obtain imine derivatives (**3a–3b**). The propargylation of imine derivatives (**3a–3b**) was performed in acetone with propargyl bromide (80% in toluene) in presence of K₂CO₃ on refluxing. Finally, convenient click chemistry approach was used for one pot synthesis of 1,2,3-triazoles, where synthesized imines with a terminal alkyne (**4a–4b**) were reacted with *N*-bromoacetamide derivatives (**5a–5g**) using sodium azide, copper sulphate pentahydrate-sodium ascorbate catalytic system in DMF: water (8:2) as shown in scheme 1. Imine linked 1,2,3-triazole compounds were furnished in high yield i.e. 76–89% as shown in Table 1.

All the newly synthesized imine-based 1,2,3-triazole hybrids (**6a–6g**, **7a–7g**) were characterized by various spectral techniques like FTIR, ¹H, ¹³C NMR and HRMS data. The characteristics band/ peaks of two representative compounds **6a** and **7a** is discussed below:

FTIR analysis

In the IR spectra of synthesized hybrids **6a** and **7a** bands were observed at 3257 cm^{-1} and 3280 cm^{-1} respectively confirmed the presence of $-\text{NH}$ group of amide functionality [46]. Formation of triazole ring was confirmed by the appearance of a band at 3142 cm^{-1} in both **6a** and **7a** because of $\text{C}-\text{H}$ stretching vibration of triazole unit [47]. In compounds **6a** and **7a**, a band of strong intensity because of the amidic $\text{C}=\text{O}$ stretching was observed at 1669 cm^{-1} and 1681 cm^{-1} , respectively. While band due to the $\text{C}=\text{N}$ stretching was detected at 1600 cm^{-1} for **6a** and at 1607 cm^{-1} for **7a** [48, 49]. For the compound **6a** out of plane bending vibrations of benzene appeared at 696 cm^{-1} and 787 cm^{-1} , whereas in compound **7a** out of plane bending vibration of benzene appeared at 837 cm^{-1} [50]. In plane bending vibrations of benzene ring were recognized in the region of $1300\text{--}1000\text{ cm}^{-1}$ and are overlapped with the other stretching vibrations. In compounds **6a** and **7a** $\text{C}-\text{N}$ stretching vibrations of azole unit was spotted at 1313 cm^{-1} and 1312 cm^{-1} respectively, whereas $\text{N}=\text{N}$ stretching was detected at 1412 cm^{-1} in **6a**, whereas in **7a** this stretching vibration is observed at 1415 cm^{-1} [51].

^1H NMR analysis

In the ^1H NMR spectrum of synthesized triazole hybrid **6a**, peak observed at δ 10.49 ppm is due to the $-\text{NH}$ proton of amide functionality. Singlets observed at δ 5.37 and δ 5.27 ppm were due to the methylene protons present adjacent to the amide functionality and oxygen atom. A singlet at δ 8.62 ppm assigned to the $-\text{C}=\text{NH}-$ proton [37]. Whereas, a confirmatory singlet was observed at δ 8.29 ppm is due to the triazolyl proton [52]. While, compound **7a** showed a peak due to $-\text{NH}$ proton of amide at δ 10.50 ppm, two singlets due to methylene protons were observed at δ 5.38 and δ 5.29 ppm. Singlets due to $-\text{C}=\text{NH}-$ and triazolyl proton were observed at δ 8.55 ppm and δ 8.31 ppm, respectively.

^{13}C NMR analysis

From the analysis of ^{13}C NMR spectrum of synthesized triazole hybrids it was found that in compound **6a**, two peaks at δ 61.64 and δ 52.68 ppm confirmed the existence of two methylenes, whereas in compound **7a** these two peaks observed at δ 61.67 and δ 52.70 ppm. C_4 carbon peak of triazole ring in **6a** appeared at δ 142.81 ppm, while in **7a** it is appeared at δ 142.58 ppm. The presence of all these peaks in synthesized compounds ^{13}C NMR spectra confirms the formation of triazoles [52].

Table 1 Synthesis of imine linked 1, 2, 3-triazole hybrids (**6a–6g**, **7a–7g**) via click chemistry

Entry	Compounds	R	Yield (%) ^[a]	M. P. (°C)
1	4a	–	85	35–37
2	4b	–	88	56–59
3	6a	H	82	160–163
4	6b	F	79	144–146
5	6c	Cl	86	162–165
6	6d	Br	82	174–176
7	6e	NO ₂	89	220–223
8	6f	OCH ₃	82	190–193
9	6g	CH ₃	77	187–190
10	7a	H	81	228–231
11	7b	F	77	219–221
12	7c	Cl	82	223–224
13	7d	Br	79	231–234
14	7e	NO ₂	82	220–222
15	7f	OCH ₃	76	226–229
16	7g	CH ₃	88	243–245

^[a] Yield of synthesized imine linked 1,2,3-triazole hybrids

Mass analysis

HRMS data of synthesized triazole hybrid **6a** was in good agreement with the calculated value. The molecular ion peak $[M+H]^+$ of **6a** C₂₄H₂₀FN₅O₂ was obtained at 430.1746 in HRMS which was in complete agreement with the calculated mass: 430.1674. When molecule get fragmented then a peak corresponding to C₁₈H₁₈N₅O₂ was obtained at 337.1354 which was in agreement with the calculated value 337.1533. Fragmentation of amide linkage produced molecular ion C₁₈H₁₄FN₄O₂ which gives a $[M+H]^+$ peak at 338.1385 which is closely resembles with the calculated value 338.1174.

Antimicrobial activity

The preliminary antimicrobial potency of the newly prepared imine linked 1,2,3-triazole hybrid compounds (**6a–6g**, **7a–7g**) as well as their precursor terminal alkynes (**4a**, **4b**) were examined against some bacterial and fungal strains using standard serial dilution method [39]. All samples were examined at five different concentrations of 50, 25, 12.5, 6.25, 3.125 µg/mL against *Bacillus subtilis* (MTCC 441), *Staphylococcus aureus* (MTCC 3160), *Pseudomonas aeruginosa* (MTCC 424), *Escherichia coli* (MTCC 16,521), *Salmonella enterica*, *Candida albicans* (MTCC 183), and *Rhizopus oryzae* microbial species.

The *in vitro* activity results evinced that most of the synthesized imine clubbed 1,2,3-triazoles (**6a–6g**, **7a–7g**) were more potent than their precursor terminal alkynes (**4a**, **4b**) against the tested microbial strains. It shows that the introduction of triazole scaffold augmented the antimicrobial potential of the synthesized hybrid molecules. Activity of all the synthesized hybrid molecules was recorded in terms of minimum inhibitory concentration and compared with their precursor terminal alkynes as well as with the ciprofloxacin (antibacterial) and fluconazole (antifungal) as shown in Table 2 and Table 3, respectively. From the antimicrobial assay results, it was observed that compound **6d** (MIC = 0.0123 $\mu\text{mol/mL}$) was found to be most potent against Gram-positive bacteria *B. subtilis*. Whereas, compound **7d** (MIC = 0.0061 $\mu\text{mol/mL}$) showed excellent and better efficacy than the ciprofloxacin (MIC = 0.0094 $\mu\text{mol/mL}$) against *S. aureus*. In case of Gram-negative bacteria, compound **6d** and **7d** showed most potent activity among synthesized compounds against two bacterial strains *S. enterica* and *P. aeruginosa*, whereas against bacterial strain *E. coli*, compounds **7c** (MIC = 0.0135 $\mu\text{mol/mL}$) and **7f** (MIC = 0.0136 $\mu\text{mol/mL}$) were reported as most active.

The *in vitro* antifungal assay of the synthesized compounds displayed that most of the imine based 1,2,3-triazole hybrids were more efficient than the fluconazole (MIC = 0.0408 $\mu\text{mol/mL}$). 1,2,3-Triazole hybrids prepared from **4b** displayed more potency than the alkyne against both the tested fungal strains. Compound **7g** displayed excellent activity against *C. albicans* having MIC (0.0141 $\mu\text{mol/mL}$), whereas compound **6d** (MIC = 0.0246 $\mu\text{mol/mL}$) and **7d** (MIC = 0.0246 $\mu\text{mol/mL}$) also displayed better activity than the fluconazole (MIC = 0.0408 $\mu\text{mol/mL}$) against *C. albicans*. All newly synthesized triazoles except **6d**, **6f** exhibited better potency than the fluconazole (MIC = 0.0408 $\mu\text{mol/mL}$) against *R. oryzae*.

Structure-activity relationship

Following SAR was established after analysing the results obtained from the antimicrobial assay of the synthesized imine linked 1,2,3-triazole hybrids:

- (i) As most of the synthesized imine-based 1,2,3-triazole hybrids displayed more efficacy than their corresponding terminal alkynes which led to the concept that molecular hybridization enhanced the activity.
- (ii) Triazoles bearing bromine atom on the phenyl ring found to be more potent against *S. enterica* and *P. aeruginosa* and there is no general trend of activity was observed for other compounds.
- (iii) Most of the triazole hybrids synthesized from meta substituted alkyne **4a** were found to be equally or more potent than the triazoles synthesized from para substituted alkyne **4b** against *B. subtilis*. Compound **6d** bearing bromine atom and **7e** bearing nitro group on the phenyl ring were found to be most active.
- (iv) Compound **7d** with bromine atom on the phenyl ring was found to be the most active against *S. aureus*. Most of the imine linked 1,2,3-triazole hybrids synthesized from alkyne **4b** were found to be more potent than the correspond-

Table 2 Antibacterial screening data of the synthesized derivatives (**4a**, **4b**, **6a–6g**, **7a–7g**)

Minimum inhibitory concentration (MIC, $\mu\text{mol/mL}$)					
Entry	<i>S. enterica</i>	<i>B. subtilis</i>	<i>S. aureus</i>	<i>P. aeruginosa</i>	<i>E. coli</i>
4a	0.0987	0.0987	0.0493	0.0987	0.0987
4b	0.0493	0.0987	0.0493	0.0987	0.0987
6a	0.0291	0.0145	0.0073	0.0291	0.0145
6b	0.0559	0.0279	0.0140	0.0279	0.0279
6c	0.0539	0.0135	0.0269	0.0135	0.0269
6d	0.0246	0.0123	0.0246	0.0123	0.0246
6e	0.0263	0.0132	0.0263	0.0132	0.0263
6f	0.0272	0.0136	0.0136	0.0272	0.0272
6g	0.0282	0.0141	0.0282	0.0141	0.0282
7a	0.0291	0.0145	0.0073	0.0145	0.0582
7b	0.0279	0.0279	0.0070	0.0140	0.0279
7c	0.0539	0.0539	0.0067	0.0135	0.0135
7d	0.0246	0.0246	0.0061	0.0123	0.0492
7e	0.0263	0.0132	0.0066	0.0132	0.0263
7f	0.0544	0.0136	0.0068	0.0136	0.0136
7g	0.0282	0.0564	0.0141	0.0282	0.0282
Ciprofloxacin	0.0094	0.0094	0.0094	0.0094	0.0094

Table 3 Antifungal screening results of the synthesized hybrids (**4a**, **4b**, **6a–6g**, **7a–7g**)

Minimum inhibitory concentration (MIC, $\mu\text{mol/mL}$)		
Compounds	<i>C. albicans</i>	<i>R. oryzae</i>
4a	0.0493	0.0247
4b	0.0987	0.0987
6a	0.0582	0.0291
6b	0.0559	0.0279
6c	0.0539	0.0135
6d	0.0246	0.0492
6e	0.0527	0.0132
6f	0.0272	0.0544
6g	0.0564	0.0282
7a	0.0291	0.0291
7b	0.0279	0.0140
7c	0.0539	0.0270
7d	0.0246	0.0123
7e	0.0527	0.0263
7f	0.0272	0.0272
7g	0.0141	0.0282
Fluconazole	0.0408	0.0408

ing imine linked 1,2,3-triazole hybrids obtained from the **4a** alkyne against *S. aureus*.

- (v) Triazole hybrids **7c** and **7f** bearing chlorine and methoxy substituent, respectively, on phenyl ring exhibited excellent activity against *E. coli*.
- (vi) Compound **7g** bearing methyl group on phenyl ring found to be most active against *C. albicans*.
- (vii) Compounds **6d** and **7d** having bromine substituent on phenyl ring displayed good to better antimicrobial efficacy against all the microbial strains.

Docking studies

In order to theorize the plausible mechanism responsible for the antimicrobial activity and to understand the effect of triazole ring insertion into the molecular structure. In silico studies were carried out. As reported in the literature, organic molecules having 1,2,3-triazole scaffold act as a good inhibitor of *E. coli* DNA gyrase, *C. albicans* 14- α sterol demethylase, α -glucosidase, etc. [15, 22]. Therefore, compounds **4a**, **4b**, **6a** and **7c** having good inhibition potential against *E. coli* were docked in the active site of *E. coli* DNA Gyrase. While, *C. albicans* enzyme sterol 14-alpha demethylase was chosen as a target for docking of compounds **4a**, **4b**, **6d** and **7g** as the latter molecules are better inhibitor of *C. albicans* growth.

Terminal alkynes **4a** and **4b** mainly interacted through hydrophobic interactions with in the active sites of DNA gyrase. Middle phenyl ring of **4a** showed one electrostatic interaction (π -anion) with GLU50. Residues ALA47, ILE78, PRO79, ALA86, ILE90 and VAL167 were involved in interactions. Triazole hybrids **6a** and **7c** made hydrogen bond with GLU42 via amide hydrogen. Nitrogen atom of triazole ring of compound **7c** also created an additional carbon-hydrogen bond with GLY119. These hydrogen bonds might be the cause of higher potency of triazole hybrids against *E. coli* as compared to their terminal alkyne precursors as the other interactions shown by these triazole hybrids are hydrophobic and with the same residues as observed in terminal alkyne interactions. The docking score of **4a**, **4b**, **6a** and **7c** was -7.0 , -6.4 , -8.0 and -8.1 kcal/mol, respectively. The value of correlation coefficient between the observed activity and the calculated binding energy was 0.95 kcal/mol which shows very good relationship between these two studies. The diagrammatic representation of docked conformation along with interacting residues is shown in Fig. 2. Also, the cartoon representation of compounds **4a**, **4b**, **6a** and **7c** with *E. coli* DNA gyrase is highlighted in Fig. 3.

Further, in sterol alpha demethylase, oxygen atom of terminal alkyne **4a** made carbon-hydrogen bond with GLY307 and middle phenyl ring was tangled in π - π stacked interactions with HEME molecule and an electrostatic π -cation interaction with HEME iron. ILE131, ILE304 and LEU376 helped through hydrophobic interactions. **4b** interacted with HEME molecule via carbon hydrogen bond, halogen bond, π - π stacked interactions and π -cation interactions. Phenoxy ring displayed π - π stacked interactions with TYR118. Triazole derivatives **6d** and **7g** created hydrogen bonds with TYR64 through amide hydrogen atom. Phenoxy ring was situated against TYR118 through T shaped π - π stacked interactions. Fluorophenyl

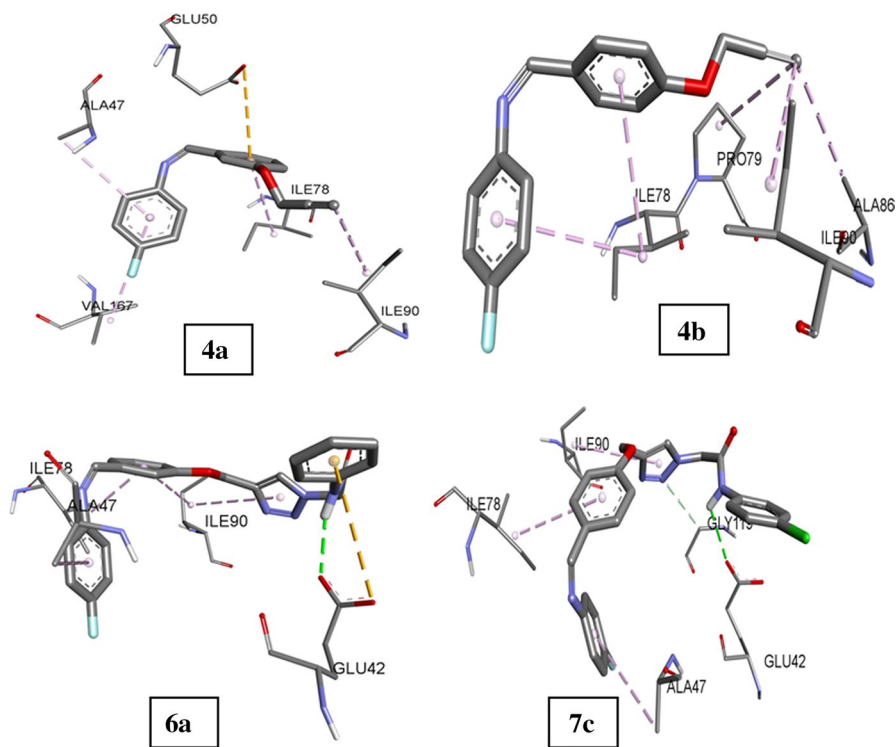


Fig. 2 Binding interactions of compound **4a**, **4b**, **6a** and **7c** in active site of *E. coli* DNA gyrase

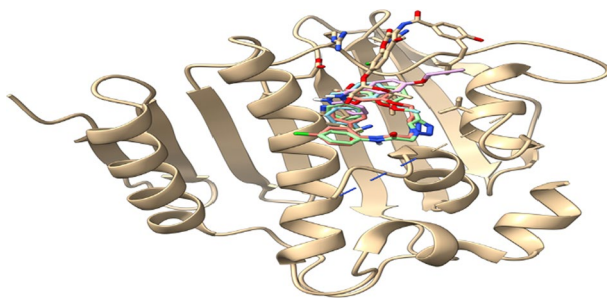


Fig. 3 Cartoon diagram showing binding conformations of docked molecules **4a** (cyan) **4b** (plum), **6a** (lime), **7c** (salmon) along with co-crystallized ligand clorobiocin (tan) in *E. coli* DNA gyrase

ring was stacked near HEME via π - π stacked interactions. These interactions along with other hydrophobic interactions involved in anchorage of these molecules in the active site are pictured in Fig. 4. The binding affinity of **4a**, **4b**, **6d** and **7g** in 5TZ1 was -10.1 , -9.4 , -11.6 and -11.8 kcal/mol. The correlation coefficient between observed antifungal activity against *C. albicans* and calculated binding affinity was 0.95 which shows good agreement between experimental and theoretical results.

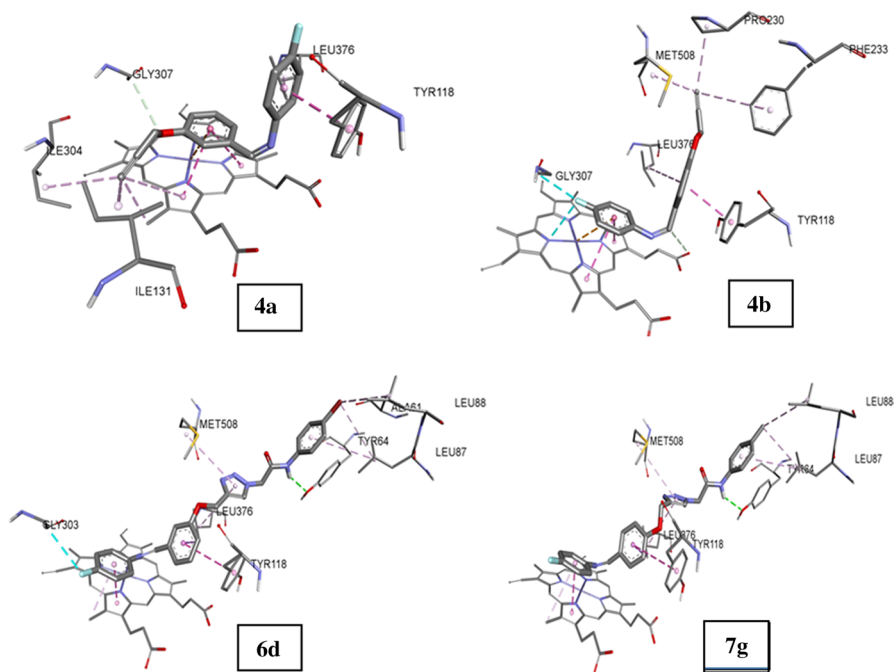


Fig. 4 Binding interactions of compound **4a**, **4b**, **6d** and **7g** in active site of *C. albicans* sterol 14- α demethylase

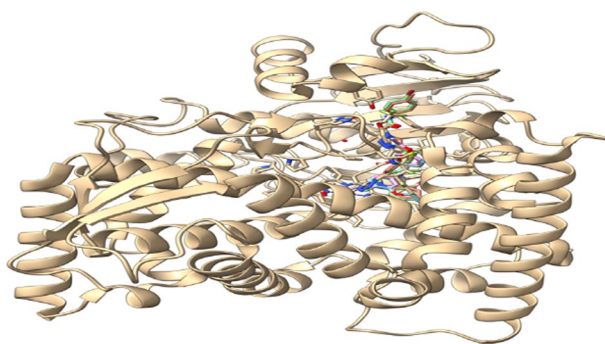


Fig. 5 Cartoon diagram showing binding conformations of docked molecules **4a** (cyan), **4b** (plum), **6d** (lime), **7g** (salmon) along with co-crystallized ligand VT1161 (tan) in *C. albicans* sterol 14- α demethylase

The docked conformations of these molecules along with co-crystallized ligand are shown in Fig. 4. Cartoon representation of compounds **4a**, **4b**, **6d** and **7g** with *C. albicans* sterol 14- α demethylase is highlighted in Fig. 5.

ADME analysis

The bioactivity score of the synthesized triazole hybrids and terminal alkynes was predicted by using an online software Molinspiration (<https://www.molinspiration.com/>) (Accessed on 24.01.2022)). As per Lipinski's rule of five, an orally active drug must have to follow at least four criteria out of the five which is as follows; (i) Molecular weight ≤ 500 (ii) hydrogen bond acceptors ≤ 10 (iii) hydrogen bond donors ≤ 5 , (iv) $\text{miLogP} \leq 5$ (v) rotatable bonds ≤ 10 . All synthesized compounds except **6d** and **7d** showed drug likeness as predicted by the bioactivity score. Presence of the rotatable bonds in synthesized hybrids makes them flexible. The higher values of TPSA for synthesized triazole hybrids as compared to their precursor terminal alkynes clearly showed that they have better transport properties and hydrogen bond formation capacity with the targeted enzymes. Percentage absorption was obtained by following the formula $\% \text{ ABS} = 109(0.345 \times \text{TPSA})$ [53].

Entry	Compounds	%ABS	MW	miLogP	TPSA	Natoms	nON	nOHNH	Nrotb	Volume
1	4a	101.55	253.28	3.83	21.60	19	2	0	4	231.69
2	4b	101.55	253.28	3.85	21.60	19	2	0	4	231.69
3	6a	80.91	429.45	4.22	81.41	32	7	1	8	377.49
4	6b	80.91	447.44	4.38	81.41	33	7	1	8	382.42
5	6c	80.91	463.90	4.90	81.41	33	7	1	8	391.02
6	6d	80.91	508.35	5.03	81.41	33	7	1	8	395.38
7	6e	65.10	474.45	4.18	127.24	35	10	1	9	400.82
8	6f	77.72	459.48	4.28	90.65	34	8	1	9	403.04
9	6g	80.91	443.48	4.67	81.41	33	7	1	8	394.05
10	7a	80.91	429.45	4.24	81.41	32	7	1	8	377.49
11	7b	80.91	447.44	4.41	81.41	33	7	1	8	382.42
12	7c	80.91	463.90	4.92	81.41	33	7	1	8	391.02
13	7d	80.91	508.35	5.05	81.41	33	7	1	8	395.38
14	7e	65.10	474.45	4.20	127.24	35	10	1	9	400.82
15	7f	77.72	459.48	4.30	90.65	34	8	1	9	403.04
16	7g	80.91	443.48	4.69	81.41	33	7	1	8	394.05

Conclusion

In summary, various new imine-based 1,2,3-triazole derivatives was synthesized via Cu(I)-catalyzed 1,3-dipolar cycloaddition reaction and their *in vitro* antimicrobial efficacy was also evaluated against various microbial strains. The antimicrobial screening results indicated that, the presence of triazole scaffold with imine led to augmentation in inhibition potential of the synthesized compounds. Triazole hybrid **7d** displayed most potent antibacterial potency against *S. aureus* with MIC of 0.0061 $\mu\text{mol/mL}$, also compound **7d** displayed effective antifungal efficacy against

R. oryzae with MIC of 0.0123 $\mu\text{mol/mL}$. The docking simulations of selected triazole hybrids with *E. coli* DNA Gyrase and *C. albicans* sterol 14- α demethylase were also in agreement with the observed antimicrobial screening data. Compound **7c** binds within *E. coli* DNA gyrase active site with binding energy of -8.1 kcal/mol, whereas compound **7g** displayed binding affinity of -11.8 kcal/mol for the sterol 14- α demethylase binding sites. Further, the value of correlation coefficient between the observed activity and the calculated binding energy was 0.95 kcal/mol which shows very good relationship between these two studies. We expect that imine linked 1,2,3-triazole hybrids may open new opportunities for the medicinal chemists as antimicrobial agents.

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Declarations

Conflict of interest No conflict of interest.

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Authors and Affiliations

Aman Kumar¹ · Kashmiri Lal¹  · Nisha Poonia¹ · Ashwani Kumar² · Anil Kumar³

¹ Department of Chemistry, Guru Jambheshwar University of Science & Technology, Hisar, Haryana 125001, India

² Department of Pharmaceutical Sciences, Guru Jambheshwar University of Science & Technology, Hisar, Haryana 125001, India

³ Department of Bio & Nano Technology, Guru Jambheshwar University of Science & Technology, Hisar, Haryana 125001, India