

Synthesis, Antimicrobial, Antioxidant and Molecular Docking Study of Some Novel Bis-1, 2, 4-Triazolo [3, 4-b]-1, 3, 4-Thiadiazoles

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Abstract

A novel series of 1-aryl-3,4-bis-(3-alkyl/phenyl-[1,2,4] triazolo[3,4-b][1,3,4]thiadiazol)-1H-pyrazole (5a-i) are synthesized by the cyclocondensation of 1-(aryl)-1H-pyrazol-3,4-dicarboxylic acids with 3-alkyl/aryl-4-amino-5-mercapto-1,2,4-triazoles. Pyrazole dicarboxylic acids were prepared by the 1, 3-dipolar cyclo addition of 3-aryl sydnone with dimethylacetylenedicarboxylate (DMAD). The newly synthesized compounds were studied for their antibacterial, antifungal and antioxidant activities. Particularly compounds 5a and 5g showed considerable antibacterial activity against the standard drug, while all the tested compounds displayed poor inhibitory effect against fungi. Compound 5d exhibited good antioxidant activity. The docking study was performed with *Acinetobacter baumannii* penicillin-binding protein target using AutoDock 4.2, which proved H-bond interaction and strong binding affinity.

Keywords: Antimicrobial; Dipolar addition; Molecular docking; Pyrazole; Triazolothiadiazole

Introduction

The present scenario in synthetic chemistry has been focused on designing new molecules by the lead hybridization-based synthesis of different pharmacophore fragments in a single molecule with improved biological efficacy.

Pyrazole derivatives have gained immense importance due to the variety of biological activities associated with them such as, antibacterial [1], antiviral [2], anticancer [3], anti-inflammatory [4], antidiabetic [5], anti-depressant [6], antioxidant [7], antitubercular [8], antihypertensive [9] etc. On the other hand, triazolothiadiazoles are reported to exhibit a broad spectrum of biological profile. They were found to possess antibacterial [10], analgesic [11], antitubercular, anticancer, anti-inflammatory and antimicrobial [12] activities.

Previous studies suggested that the presence of pyrazole and triazolothiadiazole pharmacophore plays an important role in the enhancement of pharmacological activity. Encouraged by these findings and aiming at synthesizing new hybrid molecules having enhanced biological properties [13,14], we herein report the synthesis of novel series of bis-1,2,4-triazolo[3,4-b]-1,3,4-thiadiazoles starting from aryl sydnone and their antibacterial, antifungal and antioxidant activity. Molecular docking study was also performed.

Experimental

Materials and Methods

All the reagents and solvents were purchased from Sigma-Aldrich or Hi-Media and used after distillation/recrystallization. ¹H NMR spectra were recorded on Bruker Avance II NMR spectrometer operating at 400 MHz and all the chemical shift values were reported in parts per million (ppm) relative to tetramethylsilane (TMS). Mass spectra were acquired on a SHIMADZU LCMS-8030 mass spectrometer. Melting points of the synthesized compounds were determined in open capillary tubes in Innovative DTC-967A digital melting point apparatus. SHIMADZU FT-IR 157 spectrophotometer was used for recording IR spectra. C H N analysis was performed with Vario-EI Elementar-III model analyzer. *In-silico* study was done using Auto Dock 4.2.

General procedure for the synthesis of 1-(aryl)-1H-pyrazol-3, 4-dimethyl carboxylate (2a-c)

3-Arylsydnes **1a-c** (1 mmol) and DMAD (1 mmol) in 10 mL of dry xylene was refluxed in an oil bath at 120-125 °C for 1 hr. After the completion of reaction, solvent was removed using rotary evaporator. The solid obtained was recrystallized from ethanol [15,16].

General procedure for the synthesis of 1-(aryl)-1H-pyrazol-3, 4-dicarboxylic acid (3a-c)

1-(Aryl)-1H-pyrazol-3,4-dimethyl carboxylate (1 mmol) and sodium hydroxide (2 mmol) were taken in aqueous alcohol (50 mL) and refluxed in an oil bath for 2 hrs. After cooling, the reaction mixture was acidified using hydrochloric acid (pH=2). The solid separated was filtered off and washed thoroughly with water. The dried product was recrystallized from ethanol.

1-(p-Anisyl)-1H-pyrazol-3, 4-dicarboxylic acid (3a): White solid, Yield: 95%. M.P.: 204 °C. Anal. calcd for C₁₂H₁₀N₂O₅ (%):C, 54.97; H, 3.84; N, 10.68; Found: C, 54.92; H,3.80; N,10.74. IR (KBr cm⁻¹) 3426 (O-H), 1717 (C=O), 1244 (C-O); ¹H NMR* (400 MHz, DMSO-*d*₆) δ 3.81 (s, 3H, OCH₃), 7.08-7.10 (d, 2H, J=7 Hz, o-protons of p-anisyl), 7.83-7.86 (d, 2H, J=7 Hz, m-protons of p-anisyl), 8.98 (s, 1H, H of pyrazole ring). MS: m/z: 260.90 [M⁺-1].

1-(Phenyl)-1H-pyrazol-3, 4-dicarboxylic acid (3b): Colorless needles, Yield: 71%. M.P.: 235-236 °C. Anal. calcd for C₁₁H₈N₂O₄ (%):C, 56.90; H, 3.47; N, 12.06; Found: C, 57.07; H, 3.58; N, 12.18. IR (KBr cm⁻¹) 3427 (O-H), 1717 (C=O), 1246 (C-O); ¹H NMR* (400 MHz, DMSO-*d*₆) δ 7.41-7.45 (m, 1H, p-proton of phenyl), 7.53-7.57 (dd, 2H, J=7.56 Hz, m-protons of anisyl), 7.93-7.95 (d, 2H, J=7.68 Hz, o-protons of anisyl), 9.1 (s, 1H, H of pyrazole ring). MS: m/z: 230.90 [M⁺-1].

1-(p-Tolyl)-1H-pyrazol-3, 4-dicarboxylic acid (3c): White solid, Yield: 75%. M.P.: 215 °C. Anal. calcd for C₁₂H₁₀N₂O₄ (%):C, 58.54; H, 4.09; N, 11.38; Found: C, 58.41; H, 4.16; N, 11.26. IR (KBr cm⁻¹) 3426 (O-H), 1716 (C=O), 1244 (C-O); ¹H NMR* (400 MHz, DMSO-*d*₆) δ 2.39 (s, 3H, CH₃), 7.13-7.16 (d, 2H, J=7.56Hz p-protons of p-tolyl), 7.75-7.79 (d, 2H, J=7.56 Hz, m-protons of p-tolyl), 9.0 (s, 1H, H of pyrazole ring). MS: m/z: 244.90 [M⁺-1].

*The signals due to carboxyl group protons are not seen, as may be due to rapid exchange with the protons of water impurity present in DMSO-*d*₆.

General procedure for the synthesis of 1-aryl-3,4-bis-(3-alkyl/phenyl-[1,2,4]triazolo[3,4-b][1,3,4]thiadiazol)-1H-pyrazole (5a-i)

Triazole (4a-c) (2 mmol), substituted acids (3a-c) (1 mmol), and phosphorus oxychloride (20 ml) was taken in an R.B. flask and heated in an oil bath for 6-8 hrs at 90 °C. After cooling the contents to room temperature, the resulting reaction mass was poured into a beaker having ice flakes. The solid obtained was

filtered, washed with sodium bicarbonate solution followed by water and recrystallized from ethanol.

1-(p-Anisyl)-3,4-bis-(3-methyl-[1,2,4]triazolo[3,4-b][1,3,4]thiadiazol)-1H-pyrazole (5a): Gray solid, Yield: 92%. M.P.: 292 °C. Anal. calcd for C₁₈H₁₄N₁₀O₂ (%):C, 47.99; H, 3.13; N, 31.09; Found: C, 47.96; H,3.12; N,31.05. IR (KBr cm⁻¹) 3065 (aromatic C-H), 1510 (C=N),1071 (C-S); ¹H NMR (400 MHz, DMSO-*d*₆) δ, 2.57 (s, 3H, CH₃), 2.82 (s, 3H, CH₃), 3.9 (s, 3H, OCH₃), 7.85-7.89 (m, 4H, Ar-H), 9.18 (s, 1H, H of pyrazole ring) MS : m/z: 451.05 [M⁺+1].

1-(p-Anisyl)-3,4-bis-(3-ethyl-[1,2,4]triazolo[3,4-b][1,3,4]thiadiazol)-1H-pyrazole (5b): Green solid, Yield: 87%. M.P.: 276 °C. Anal. calcd for C₂₀H₁₈N₁₀O₂ (%):C, 50.20; H, 3.79; N, 29.27; Found: C, 50.14; H,3.76; N,29.24. IR (KBr cm⁻¹) 3061 (aromatic C-H), 1516 (C=N),1072 (C-S); ¹H NMR (400 MHz, DMSO-*d*₆) δ 1.44-1.49 (m, 6H, CH₃), 3.10-3.18 (m, 4H, CH₂), 3.85 (s, 3H, OCH₃), 7.08 (d, 2H, J= 9 Hz, o-protons of anisyl), 7.91 (d, 2H, J= 9Hz, m-protons of anisyl), 9.32 (s, 1H, H of pyrazole ring) MS : m/z: 479.10 [M⁺+1].

1-(p-Anisyl)-3,4-bis-(3-phenyl-[1,2,4]triazolo[3,4-b][1,3,4]thiadiazol)-1H-pyrazole (5c): Green solid, Yield: 87 %. M.P.: 276 °C. Anal. calcd for C₂₈H₁₈N₁₀O₂ (%):C, 58.52; H, 3.16; N, 24.37. Found: C, 58.48; H, 3.14; N, 24.32. IR (KBr cm⁻¹) 3059 (aromatic C-H), 1516 (C=N), 1080 (C-S); ¹H NMR (400 MHz, DMSO-*d*₆) δ 3.89 (s, 3H, OCH₃), 7.2-8.3 (m, 14H, Ar-H), 9.6 (s, 1H, H of pyrazole ring) MS: m/z: 575.00 [M⁺+1].

1-Phenyl-3,4-bis-(3-methyl-[1,2,4]triazolo[3,4-b][1,3,4]thiadiazol)-1H-pyrazole (5d): Gray solid, Yield: 73 %. M.P.: 293 °C. Anal. calcd for C₁₇H₁₂N₁₀S₂ (%):C, 48.56; H, 2.88; N, 33.31. Found: C, 48.54; H,2.86; N,33.28. IR (KBr cm⁻¹) 3076 (aromatic C-H), 1516 (C=N),1082 (C-S); ¹H NMR (400 MHz, DMSO-*d*₆) 2.61 (s, 3H, CH₃), 2.82 (s, 3H, CH₃), 7.51-8.01 (m, 5H, Ar-H), 9.56 (s, 1H, H of pyrazole ring) MS : m/z: 421.05 [M⁺+1].

1-Phenyl-3,4-bis-(3-ethyl-[1,2,4]triazolo[3,4-b][1,3,4]thiadiazol)-1H-pyrazole (5e): White solid, Yield: 73%. M.P.: 268 °C. Anal. calcd for C₂₀H₁₆N₁₀S₂ (%):C, 50.88; H, 3.60; N, 31.23. Found: C, 50.84; H,3.58; N, 31.20. IR (KBr cm⁻¹) 3071 (aromatic C-H), 1510 (C=N),1070 (C-S); ¹H NMR (400 MHz, DMSO-*d*₆) δ 1.36-1.39 (m, 6H, CH₃), 3.05-3.17 (m, 4H, CH₂), 7.62-7.66 (d, 2H, J= 7.56 Hz, o-protons of phenyl), 8.04 (d, 2H, J= 7.64Hz, m-protons of phenyl), 7.5-7.54 (dd, 1H, J=7.4 Hz, p-proton of phenyl), 9.58 (s, 1H, H of pyrazole ring) MS : m/z: 449.05 [M⁺+1].

1-Phenyl-3,4-bis-(3-phenyl-[1,2,4]triazolo[3,4-b][1,3,4]thiadiazol)-1H-pyrazole (5f): White solid, Yield: 91%. M.P.: 294 °C. Anal. calcd for C₂₇H₁₆N₁₀S₂ (%):C, 59.54; H, 2.96; N, 25.72. Found: C, 59.52; H,2.92; N,25.68. IR (KBr cm⁻¹) 3076 (aromatic C-H), 1514 (C=N),1072 (C-S); ¹H NMR (400 MHz, DMSO-*d*₆) δ 7.44 (m, 3H, protons of phenyl), 7.48 (m, 3H, protons of phenyl), 7.53 (d, 1H, J= 7.6Hz, proton of phenyl),

7.61-7.64 (t, 2H, J= 7.2Hz, protons of phenyl), 7.862 (t, 2H, J= 7.2Hz, protons of phenyl), 8.20-8.22 (dd, 2H, J= 7.6Hz, protons of phenyl), 8.34-8.36 (m, 2H, protons of phenyl), 8.7 (s, 1H, H of pyrazole ring). MS: m/z: 545.05 [M⁺+1].

1-(p-Tolyl)-3,4-bis-(3-methyl-[1,2,4]triazolo[3,4-b][1,3,4]thiadiazol)-1H-pyrazole (5g): Green solid, Yield: 85%. M.P.: 297 °C. Anal. calcd for C₁₈H₁₄N₁₀S₂ (%):C, 49.76; H, 3.25; N, 32.24. Found: C, 49.74; H, 3.22 N, 32.21. IR (KBr cm⁻¹) 3068 (aromatic C-H), 1506 (C=N), 1072 (C-S); ¹H NMR (400 MHz, DMSO-*d*₆) δ 2.42 (s, 3H, CH₃), 2.71 (s, 3H, CH₃), 2.74 (s, 3H, CH₃), 7.39 (d, 2H, J= 8.24Hz, o-protons of tolyl), 7.90 (d, 2H, J= 8.48Hz, m-protons of tolyl) 9.46 (s, 1H, H of pyrazole ring) MS : m/z: 435.05 [M⁺+1].

1-(p-Tolyl)-3,4-bis-(3-ethyl-[1,2,4]triazolo[3,4-b][1,3,4]thiadiazol)-1H-pyrazole (5h): Greenish blue solid, Yield: 76%. M.P.: 241 °C. Anal. calcd for C₂₀H₁₈N₁₀S₂ (%):C, 51.93; H, 3.92; N, 30.28. Found: C, 51.88; H, 3.89; N, 30.26. IR (KBr cm⁻¹) 3061 (aromatic C-H), 1513 (C=N), 1071 (C-S); ¹H NMR (400 MHz, DMSO-*d*₆) δ 1.28-1.36 (m, 6H, CH₃), δ 2.39 (s, 3H, CH₃), 3.01-3.15 (m, 4H, CH₂), 7.28 (d, 2H, J= 8.42 Hz, o-protons of tolyl), 7.85 (d, 2H, J= 8.48Hz, m-protons of tolyl), 9.57 (s, 1H, H of pyrazole ring) MS : m/z: 463.10 [M⁺+1].

1-(p-Tolyl)-3,4-bis-(3-phenyl-[1,2,4]triazolo[3,4-b][1,3,4]thiadiazol)-1H-pyrazole (5i): Green solid, Yield: 91 %. M.P.: >303 °C. Anal. calcd for C₂₈H₁₈N₁₀S₂ (%):C, 60.20; H, 3.25; N, 25.07. Found: C, 60.18; H, 3.23; N, 25.02. IR (KBr cm⁻¹) 3072 (aromatic C-H), 1516 (C=N), 1072 (C-S); ¹H NMR (400 MHz, DMSO-*d*₆) δ 2.50 (s, 3H, CH₃), 7.29-8.3(m, 14H, Ar-H), 9.6 (s, 1H, H of pyrazole ring) MS: m/z: 559.05 [M⁺+1].

Assay of *in vitro* antibacterial activity

Bacterial and fungal strains were purchased from National collection of industrial microorganisms, Pune, India. Antibacterial activity was tested against Gram-positive bacteria *Staphylococcus aureus* (NCIM - 5021), *Bacillus subtilis* (NCIM 2197) and Gram-negative bacteria *Escherichia coli* (NCIM-2931), *Pseudomonas aeruginosa* (NCIM-2036) using Ciprofloxacin as the reference drug. Antifungal activity of the newly synthesized compounds was tested against two fungi namely *Candida albicans* (NCIM 3471) and *Aspergillus niger* (NCIM 3452) using Fluconazole as the reference drug.

The sterilized nutrient agar medium was distributed 100 mL each in two 250 mL conical flasks and allowed to cool to room temperature. To these media, 18-24 h grown bacterial/fungal sub-cultures were added and shaken thoroughly to ensure uniform distribution of organisms throughout the medium. Then, agar medium was distributed in equal portions, in sterilized Petri dishes, ensuring that each Petri dish contains about 45-50 mL of the medium. The medium was then allowed for solidification. The cups were made with the help of a sterile cork borer (6 mm diameter) punching into the set of agar media. The solutions of required concentrations (100 µg/mL) of test compounds were prepared by dissolving the compounds

in DMSO were filled into the cups with 1 mL of respective solution. Then, the Petri dishes were kept for incubation in an inverted position for 24-48 h at 37 °C in an incubator. When growth inhibition zones were developed surrounding each cup, their diameter in mm was measured and compared with that of the standard drugs [17,18]. Each experiment was made in triplicate using DMSO as a control.

Molecular Docking studies

The binding interaction between macromolecule and ligands was done using AutoDock 4.2. Lamarckian genetic algorithm was used to study the docking calculation generated few poses for ligand molecules with the protein target [19]. Polar hydrogen bond network was optimized and the systematic Kollaman charges were added by means of a cluster-based approach. The grid map which was centered was predicted from the ligplot. In all the cases, we have used grid maps with a grid box size of 60×60×60 Å³ points with a grid-point spacing of 0.375 Å. During docking, centre grid parameters were specified for x, y and z axis as -55.15, -35.444 and 42.741, respectively. The Lamarckian genetic algorithm, the pseudo-Solis and Wets methods were applied for minimization using default parameters. Binding energy, torsional energy, intermolecular energy, number of H-bonds and RMS value were recorded in each ligand bound.

Assay of *in vitro* antioxidant activity

The free radical scavenging activity of test sample was measured by DPPH scavenging assay. Free radical scavenging activity of the test compounds was carried based on the scavenging activity of stable DPPH. 100 µg/mL of each test sample and standard BHA was taken in different test tubes and the volume was adjusted to 1mL using DMSO. Freshly prepared 1mL of 0.1 mM DPPH solution was mixed and vortexed thoroughly and left in dark for 30 min. The absorbance of stable DPPH radical was measured at 517 nm. The DPPH control was prepared using the same procedure. Radical scavenging activity was expressed as the inhibition percentage and was calculated using the equation [20].

$$\text{DPPH radical scavenging activity (\%)} = \frac{(A_{\text{Control}} - A_{\text{Sample}})}{(A_{\text{Control}})} \times 100$$

Where A_{Control} is the absorbance of DPPH radical+methanol; A_{Sample} is the absorbance of DPPH radical+test sample/standard BHA.

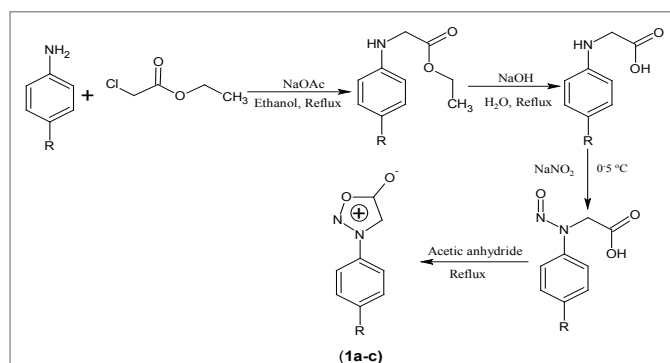
Results and Discussion

Synthesis

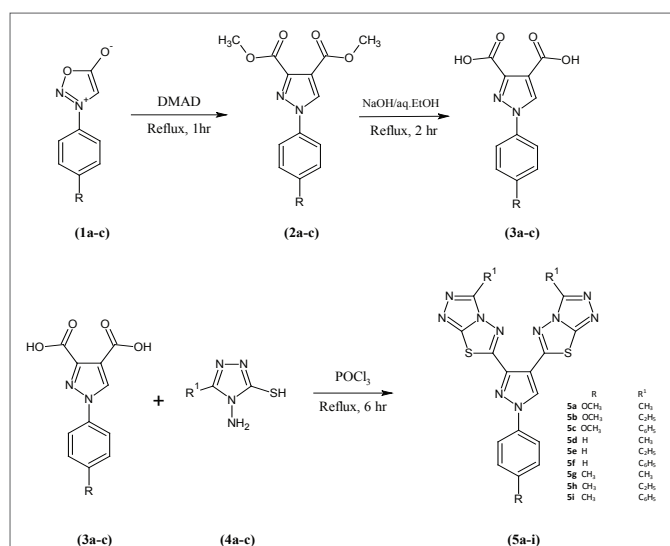
3-Aryl substituted sydnonones **1** were obtained by the reaction of appropriately substituted aniline with ethyl chloroacetate followed by hydrolysis, nitrosation and cyclization with acetic anhydride [21] (Scheme 1). These sydnonones **1** when treated with DMAD underwent 1, 3-dipolar cycloaddition reaction to give 1-aryl-1H-pyrazole-3, 4-dimethylcarboxylate **2**. Hydrolysis of

2 with aqueous alcoholic sodium hydroxide gave 1-aryl-1H-pyrazole-3,4-dicarboxylic acid **3**. 3-Alkyl/phenyl-4-amino-5-mercapto-1,2,4-triazoles **4** were prepared as per the procedures reported in the literature [22,23]. Condensation of 3-alkyl/phenyl-4-amino-5-mercapto-1,2,4-triazoles **4** with 1-aryl-1H-pyrazole-3,4-dicarboxylic acid **3** in presence of phosphorous oxychloride as condensing agent gave the corresponding 1-aryl-3,4-bis-(3-alkyl/phenyl)-[1,2,4]-triazolo[3,4-b]-[1,3,4]-thiadiazol-1H-pyrazole (**5a-i**) is as shown in Scheme 2.

The structure of newly synthesized compounds was confirmed by ¹H-NMR, IR, LCMS and C, H, N analysis. The absence of S-H and NH₂ absorption bands confirmed the formation of product. The IR spectra of compound **5a-5i** showed absorption peak at 1506-1516 cm⁻¹ which is attributed to the stretching vibration of C=N. The characteristic absorption band due to C-S stretching was observed at 1070-1082 cm⁻¹, whereas C-H stretching bands at 3059-3076 cm⁻¹ associated with the aromatic rings were observed in all the molecules. ¹H-NMR spectrum of compound **5e** showed multiplet of methyl protons at δ, 1.36-1.39 ppm integrating for six protons. A multiplet due to four methylene protons was observed at δ, 3.05-3.17 ppm.



Scheme 1: The synthesis of sydnone **1a-1c**.



Scheme 2: Synthesis of bis-1,2,4-triazolo [3,4-b]-1,3,4-thiadiazoles **5a-5i**.

The peaks due to aromatic protons were seen at δ, 7.62-7.66, 8.04 and 7.5-7.54 ppm pertain to ortho, meta and para protons of phenyl ring of pyrazole. While the proton of pyrazole ring displayed as a singlet at δ, 9.58 ppm. Further evidence for the formation of triazolo-thiadiazoles (**5a-i**) was obtained by recording mass spectra, where molecular ion peaks obtained were in consistence with their molecular formula.

Antimicrobial Studies

All the newly synthesized triazolo-thiadiazoles (**5a-i**) were investigated for antibacterial and antifungal activity. Compound **5a** showed good activity against *Bacillus subtilis* and compound **5g** showed good activity against *Pseudomonas aeruginosa*. None of the compounds showed any considerable antifungal activity (Table 1).

Molecular Docking Studies

All the compounds (**5a-i**) were found to have minimum binding energy ranging from -5.19 to -8.99 kJ/mol with antimicrobial Acinetobacter baumannii penicillin-binding protein target (PDB Code: 3UDI). Among the molecules tested for docking study, 1-p-tolyl-3,4-bis-(3-ethyl-[1,2,4] triazolo[3,4-b] [1,3,4]thiadiazol)-1H-pyrazole **5h** showed minimum binding energy of -8.99 kJ/mol with ligand efficiency of -0.28. In the selected protein target maximum numbers of residues are nearer to the drug molecule and are hydrophobic in nature [24].

The ligand molecules, **5a**, **5b** and **5g** revealed binding energy of -8.32, -8.13 and -7.83 kJ/mol, with ligand efficiency of -0.27, -0.25 and -0.26, respectively. The completely wrapping of the molecules by amino acid residues at the active site pocket region as displayed in Figure 1. In **5a**, the oxygen atom of methoxy group displayed H-bonding interaction with the hydrogen atom of Ser434 at a distance of (2.174) Å, while the sulphur atom present in thiadiazolotriazole ring of the compound **5d**, **5e** and **5g** was involved in the H-bonding with the active site of amino acid residue Tyr485 at a distance of (2.667), (2.872) and (2.816) Å, respectively as depicted in Figure 2. The docking study results showed that the molecules **5a-5h** has good inhibition constant, vdW + H-bond + desolv energy with best RMSD value. The details of docked score results of the molecules are given in Table 2.

Antioxidant Studies

The synthesized compounds showed DPPH scavenging activity varying from 76.38±0.32% to 18.01±0.15%, whereas standard drug BHA showed 88.33±0.33% inhibition. Compound **5d**, **5e** and **5h** displayed 76.38±0.32%, 70.37±0.20% and 54.74±0.29% of activity closer to the standard employed. The percentage radical scavenging activity of the bis triazolothiadiazole has been described in Figure 3.

Conclusions

A novel series of bis-[1,2,4]triazolo[3,4-b][1,3,4]thiadiazol-1H-pyrazole were prepared by the cyclocondensation of

Table 1: Antibacterial and antifungal activity data of compounds **5a-5i**.

Compounds	Diameter of zone of inhibition (in mm) at 100 µg/mL					
	Antibacterial activities				Antifungal activities	
	Gram positive bacteria		Gram negative bacteria		<i>C. albicans</i>	<i>A. niger</i>
<i>S. aureus</i>	<i>B. subtilis</i>	<i>E. coli</i>	<i>P. aeruginosa</i>			
5a	11.5 ± 0.70	19.5 ± 0.70	12.5 ± 0.70	12 ± 1.4	10.5 ± 0.70	5 ± 0.00
5b	10.5 ± 0.70	17.5 ± 0.70	14.5 ± 0.70	11 ± 1.4	9 ± 1.4	4.5 ± 0.70
5c	15 ± 1.4	14 ± 0.0	9.5 ± 0.70	13 ± 1.4	9.5 ± 0.70	5.5 ± 0.70
5d	13.5 ± 0.70	12.5 ± 0.70	14.5 ± 0.70	11 ± 0.0	10 ± 0.0	6 ± 0.0
5e	16 ± 0.0	15.5 ± 2.1	12.5 ± 0.70	12.5 ± 0.70	10.5 ± 2.1	5.5 ± 0.70
5f	14.5 ± 0.70	13.5 ± 0.70	15 ± 1.4	13.5 ± 0.70	9 ± 1.4	4.5 ± 0.70
5g	12.5 ± 0.70	12 ± 0.0	14 ± 1.4	19.5 ± 0.70	9 ± 1.4	7 ± 0.0
5h	15.5 ± 0.70	17 ± 1.4	14 ± 0.0	18.5 ± 0.70	5.5 ± 0.70	5 ± 0.0
5i	14.5 ± 0.70	14.5 ± 0.70	10.5 ± 0.70	14 ± 1.4	9.5 ± 0.70	5.5 ± 0.70
Ciprofloxacin	23.5 ± 0.70	23.5 ± 0.70	22.5 ± 0.70	22.5 ± 0.70	-	-
Fluconazole	-	-	-	-	19.5 ± 0.70	22 ± 1.4

Table 2: The dock score results of triazolo-thiadiazoles (**5a-i**) with *Acinetobacter baumannii* penicillin-binding protein target (**PDB Code: 3UDI**).

Compounds	Binding Energy (kJ mol ⁻¹)	Ligand Efficiency	Inhibition Constant uM	vdW+H-bond+ desolv energy kcal/mol	No. of H- bonds	Bonding residues	Bond Length (Å)
5a	-8.32	-0.27	793.777	-9.33	2	3UDI:A: SER434:HG	2.174
5b	-8.13	-0.25	1.11	-9.16	-	-	-
5c	-7.19	-0.18	5.41	-8.83	-	-	-
5d	-7.39	-0.25	3.81	-7.88	1	3UDI: A: TYR485: O	2.667
5e	-7.2	-0.23	5.29	-8.53	1	3UDI: A: TYR485: O	2.872
5f	-6.35	-0.16	22.06	-7.96	-	-	-
5g	-7.83	-0.26	1.81	-8.29	1	3UDI: A: TYR485: O	2.816
5h	-8.99	-0.28	256.32	-9.86	-	-	-
5i	-5.19	-0.13	157.21	-6.87	-	-	-

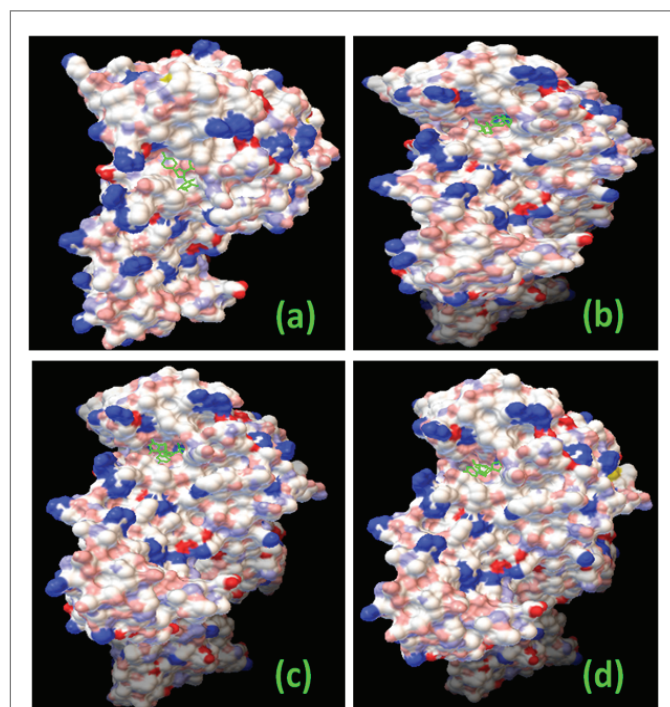


Figure 1: Figure (a), (b), (c) and (d) represents the enfolding of molecules **5a**, **5b**, **5g** and **5h** in the active site pocket of antimicrobial protein *Acinetobacter baumannii* penicillin-binding protein target (**PDB Code: 3UDI**).

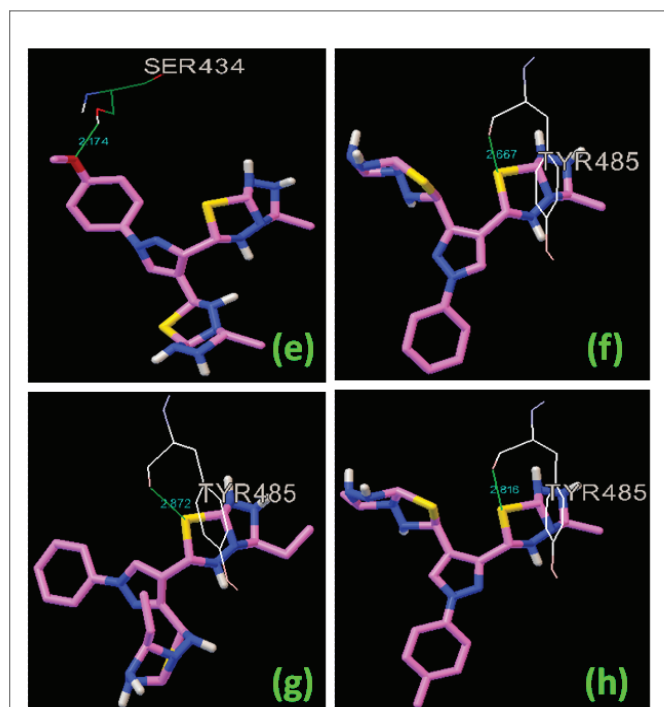
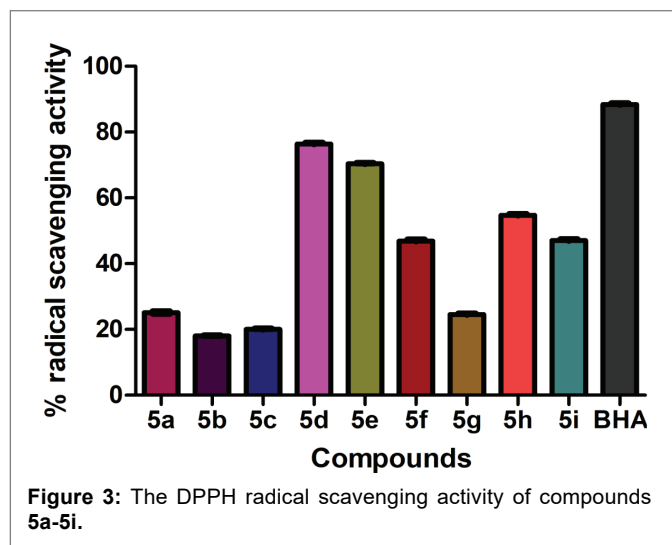


Figure 2: Figure (e), (f), (g) and (h) represents the H-bond interaction of ligand molecules **5a**, **5d**, **5e** and **5g** *Acinetobacter baumannii* penicillin-binding protein target (**PDB Code: 3UDI**).



3-substituted-4-amino-5-mercapto-1,2,4-triazoles with 1-(aryl)-1H-pyrazol-3,4-dicarboxylic acids. The newly synthesized compounds were characterized by spectral and analytical methods. Further molecular docking, antimicrobial and antioxidant studies were carried out. Compounds **5a** and **5g** showed significant antibacterial activity against *Staphylococcus aureus* and *Pseudomonas aeruginosa*. But none of the compound exhibited good antifungal activity. Compounds **5d**, **5e** and **5h** showed good radical scavenging property.

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