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Synthesis, molecular docking and antimicrobial evaluation of novel benzoxazole derivatives

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Abstract In this research, previously and newly synthesized 5-amino-2-(4-substitutedphenyl/benzyl)benzoxazoles (3a-3l) and 2-substituted-5-(4-nitro/aminophenylsulfonamido)benzoxazoles (5a-5l, 6a-6l) were evaluated for their antimicrobial activities against Pseudomonas aeruginosa ATCC 27853, Staphylococcus aureus ATCC 29213, Escherichia coli ATCC 25922, Enterococcus faecalis ATCC 29212 and Mycobacterium tuberculosis H37RV ATCC 27294 and their drug-resistant isolates Candida albicans ATCC 10231 and Candida krusei ATCC 6258. The chemical structures of the newly synthesized compounds were characterized by IR, ¹H NMR, ¹³C NMR, LC-MS and elemental analysis. Microbiological results indicated that the compounds possessed a broad spectrum of activity against the tested microorganisms at the minimum inhibitory concentration (MIC) values between 256 and 8 µg/mL. Compounds 3a, 3c and 3f exhibited significant antimycobacterial activity showing MIC value of 8 µg/mL against both M. tuberculosis and its drug-resistant isolate. InhA, the enoyl-acyl carrier protein reductase from M. tuberculosis, is one of the key enzymes in the FASII system involved in mycobacterial fatty acid elongation cycle, which has been validated as an effective antimicrobial target. Molecular docking into active site of InhA

was performed on **3FNE.PDB** file to understand ligandprotein interactions. The compounds obtained from this research can be used as scaffolds in the design of new potent drugs.

Keywords Antimicrobial activity · Benzoxazole · Enoyl-ACP reductase · Molecular docking · *Mycobacterium tuberculosis* · Sulfonamide

Introduction

In the past few decades, the dramatically rising prevalence of multidrug-resistant microbial infections has caused a serious healthcare problem. In particular, the emergence of multidrug-resistant strains of Gram-positive bacterial pathogens such as methicillin-resistant *Staphylococcus aureus* and *Staphylococcus epidermis* and vancomycin-resistant enterococcal infection is a problem of ever-increasing significance (Dalhoff, 1994; Lee and Hacker, 1999; Livermore, 2000; Poole, 2001; Abbanat *et al.*, 2003). In order to prevent these serious medical problems, there is still need for the new classes of antimicrobial agents.

Tuberculosis (TB), in the past also called phthisis, phthisis pulmonalis or consumption, is a widespread, infectious disease caused by various strains of mycobacteria, usually *Mycobacterium tuberculosis* (Kumar *et al.*, 2007). The disease has been the leading cause of morbidity and mortality among the infectious diseases. To address these issues, it is necessary to increase research and developmental studies to obtain novel and potent drugs.

The benzoxazole scaffold is a core structure found in a wide class of natural and synthetic compounds. Benzoxazole ring has provided the basis for the design of biologically relevant molecules with broad therapeutic importance

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(Prudhomme et al., 1986; Yalcin et al., 1992; Balani et al., 1992; Davey et al., 1993; Kim et al., 1996; Oren et al., 1997; Ueki and Taniguchi, 1997; Oren et al., 1998; Akbay et al., 2003; Plemper et al., 2004; Pinar et al., 2004; Yildiz-Oren et al., 2004a, b; Vinsova et al., 2005; Rida et al., 2005; Varga et al., 2005; Lage et al., 2006; Oksuzoglu et al., 2008; Ertan et al., 2009; Zilifdar et al., 2014). Besides, several benzoxazole derivatives have been reported to possess antitubercular activity (Ileana et al., 2006; Klimesova et al., 2009; Rana et al., 2014).

On the other hand, heterocycles possessing sulfonamido moieties have attracted obvious attention due to their role as pharmacophores (Andrighetti-Fröhner *et al.*, 2009; Akurathi *et al.*, 2010; Lu *et al.*, 2011; Luo *et al.*, 2011; Chandak *et al.*, 2013; Kamal *et al.*, 2013). Studies revealed that sulfonamide compounds had antibacterial effects (Gadad *et al.*, 2000; Ezabadi *et al.*, 2008; Kamal *et al.*, 2013; Azab *et al.*, 2013). In the last few years, we reported the synthesis of several 2,5-(and/or 6)-di(or tri)-substitutedbenzoxazole derivatives as the antimicrobial agents (Yalcin *et al.*, 1992; Oren *et al.*, 1997, 1998; Yildiz-Oren *et al.*, 2004a, b; Temiz-Arpaci *et al.*, 2005; Arisoy *et al.*, 2008; Alper-Hayta *et al.*, 2008; Ertan *et al.*, 2009).

InhA, the enoyl-acyl carrier protein reductase from M. tuberculosis, is one of the key enzymes in the FASII system involved in mycobacterial fatty acid elongation cycle, which has been validated as an effective antimicrobial target. Inhibition of mycolic acid biosynthesis is the first event detected in M. tuberculosis treated with isoniazid (INH), one of the most effective anti-TB agents (Takayama et al., 1972). At first INH must be activated by KatG, a catalase-peroxidase that oxidizes INH to an acyl-radical which then forms a covalent adduct with co-factor NAD⁺, the co-substrate for InhA. Then, the INH-NAD adduct functions as a potent inhibitor of InhA (Zhang et al., 1992; Johnsson and Schultz, 1994). The InhA is one of the best validated targets for the development of antitubercular agents. However, the majority of isoniazid (INH)-resistant clinical strains are observed mainly due to the emergence of KatG mutants that do not form an INH-NAD adduct. Thus, compounds that directly inhibit InhA avoiding activation by KatG would be promising candidates for combating MDR-TB (Lu et al., 2010).

Triclosan (5-chloro-2-(2,4-dichloro-phenoxy)phenol ether) is known to inhibit the synthesis of fatty acids targeting InhA directly whose inhibition leads to the lysis of *M. tuberculosis*. In an early study using a structure-based drug design approach, a series of 5-substituted derivatives of triclosan was developed. (5-(2-Pyridylmethyl)-2-(2,4dichloro-phenoxy)phenol ether) is one of the derivatives of triclosan which are found more potent than triclosan against purified InhA (Freundlich *et al.*, 2009). X-ray crystal structures of InhA in complex with this triclosan derivative become available in Protein Data Bank (3FNE). It is reported that triclosan and its derivatives bind similarly with catalytic residue Tyr158 and co-factor NAD⁺ (Lu *et al.*, 2010; Shrinivas *et al.*, 2015). Because of that reason binding to Tyr158 and co-factor NAD⁺ is important for direct inhibition of this enzyme.

In this study, new promising bioactive compounds were designed and synthesized by a simple and efficient method, followed by the evaluation of their biological activities. The synthesis emphasized a strategy that two pharmacologically compatible moieties in one molecule by attaching a sulfonamide group to a benzoxazoles combined. Based on the above considerations, we synthesized some novel 2-substituted-5-(4-nitro/aminophenylsulfonamido)benzoxazole derivatives by using 5-amino-2-(4-substitutedphenyl/ benzyl)benzoxazole as an starting compounds. In here, compounds 5b, 5d, 5g, 5j, 5k, 5l and 6a-6d, 6f-6l were synthesized for the first time. All of the previously (Ertan-Bolelli et al., 2014) and newly synthesized sulfonamidobenzoxazole and 5-aminobenzoxazole derivatives were evaluated for their in vitro antimicrobial activities against human pathogenic microbes. Molecular docking into active site of InhA was performed to understand ligand-protein interactions.

Materials and methods

Chemistry

All chemicals and solvents were purchased from commercial vendors and were used without purification. The progress of the reaction was monitored on ready-made silica gel plates (Merck). The melting points were measured with a capillary melting point apparatus (Buchi B540) and were uncorrected. Yields were calculated after recrystallization. The IR spectra were recorded on a Jasco FT/IR-420 spectrometer as KBr disk. The ¹H NMR spectra were recorded employing a VARIAN Mercury 400-MHz FT spectrometer, chemical shifts (d) were in parts per million relative to TMS, and coupling constants (J) were reported in hertz. Mass spectra were taken on a Waters Micromass ZQ using the ESI method. Elemental analyses were performed by Leco CHNS-932 CHNS-O analyzer. The result of the elemental analyses (C, H, N, S) were within ± 0.4 % of the calculated amounts.

General procedure for the preparation of 2-(4substitutedphenyl/benzyl)-5-aminobenzoxazoles (3a-3l)

The derivatives were synthesized by heating 2,4-diaminophenol dihydrochloride (1) (0.01 mol) with suitable acid (2a-2l) (0.01 mol) in polyphosphoric acid (PPA) (24 g) and stirring at 170–200 °C for 1.5–2.5 h. At the end of the reaction period, the residue was poured into an ice-water mixture and neutralized with an excess of NaOH (10 %) solution, and the residue was filtered and boiled with charcoal (200 mg) in ethanol and filtered. After the evaporation of solvent in vacuo, the crude product was obtained and recrystallized from ethanol–water (1:3) mixture (Yildiz-Oren *et al.*, 2004a; Sener *et al.*, 1987; Wynne *et al.*, 2009).

2-Phenyl-5-aminobenzoxazole (**3a**) (Sener *et al.*, 1987) This compound was prepared by using compounds **1** and **2a** at 180 °C for 2.5 h. It was obtained as creamy solid in 52 % yield; mp 152–154 °C; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 5.13$ (2H, s, NH₂), 6.68 (1H, dd, J = 8.8 Hz, J = 2.0 Hz, H-6), 6.88 (1H, d, J = 2.0 Hz, H-4), 7.43 (1H, d, J = 8.8 Hz, H-7), 7.57–7.60 (3H, m, H-3', H-4', H-5'), 8.13–8.15 (2H, m, H-2', H-6'); ESIMS *m*/*z* 211.7 [M + H]⁺ (75), 252.7 [M + H + 41(CH₃CN)]⁺ (100); Anal. Calcd. for C₁₃H₁₀N₂O: C, 74.27; H, 4.79; N, 13.33. Found: C, 74.08; H, 4.82; N, 13.03.

2-(4-Chlorophenyl)-5-aminobenzoxazole (**3b**) (Chancellor *et al.*, 2011) This compound was prepared by using compounds **1** and **2b** at 200 °C for 1.5 h. It was obtained as creamy solid in 82 % yield; mp 193–195 °C; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 5.14$ (2H, s, NH₂), 6.67 (1H, dd, J = 8.8 Hz, J: 2.4 Hz, H-6), 6.86 (1H, d, J = 2.4 Hz, H-4), 7.40 (1H, d, J = 8.8 Hz, H-7), 7.63 (2H, d, J = 8.4 Hz, H-3', H-5'), 8.11 (2H, d, J = 8.8 Hz, H-2', H-6'); ESIMS *m*/*z* 245.6 [M + H]⁺ (40), 247.6 [M + H + 2]⁺ (13), 286.7 [M + H + 41(CH₃CN)]⁺ (100), 288.7 [M + H + 2 + 41 (CH₃CN)]⁺ (36); Anal. Calcd. for C₁₃H₉ClN₂O: C, 63.81; H, 3.71; N, 11.45. Found: C, 63.54; H, 3.89; N, 11.39.

2-(4-Fluorophenyl)-5-aminobenzoxazole (**3c**) (Sener *et al.*, 1987) This compound was prepared by using compounds **1** and **2c** at 200 °C for 2 h. It was obtained as creamy solid in 86 % yield; mp 157–159 °C; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 5.14$ (2H, s, NH₂), 6.67 (1H, dd, J = 8.8 Hz, J = 2.4 Hz, H-6), 6.87 (1H, d, J = 2.4 Hz, H-4), 7.41–7.45 (3H, m, H-7, H-3', H-5'), 8.16–8.20 (2H, m, H-2', H-6'); ESIMS *m*/*z* 229.7 [M + H]⁺ (55), 270.8 [M + H + 41(CH₃CN)]⁺ (100); Anal. Calcd. for C₁₃H₉ FN₂O: C, 68.42; H, 3.97; N, 12.27. Found: C, 67.95; H, 3.96; N, 11.98.

2-(4-Bromophenyl)-5-aminobenzoxazole (**3d**) (Sener *et al.*, 1987) This compound was prepared by using compounds **1** and **2d** at 200 °C for 2.5 h. It was obtained as creamy solid in 60 % yield; mp 197–200 °C; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 5.14$ (2H, s, NH₂), 6.67 (1H, dd, J = 8.8 Hz, J = 2.4 Hz, H-6), 6.85 (1H, d, J = 2.4 Hz, H-4), 7.40 (1H, d, J = 9.2 Hz, H-7), 7.77 (2H, d,

 $J = 8.4 \text{ Hz}, \text{ H-3'}, \text{ H-5'}, 8.04 (2\text{H}, \text{d}, J = 8.8 \text{ Hz}, \text{ H-2'}, \text{H-6'}; \text{ ESIMS } m/z 289.6 [M + \text{H}]^+ (58), 291.7 [M + \text{H} + 2]^+ (48), 330.7 [M + \text{H} + 41(\text{CH}_3\text{CN})]^+ (92), 332.7 [M + \text{H} + 2 + 41(\text{CH}_3\text{CN})]^+ (100); \text{ Anal. Calcd. for } \text{C}_{13}\text{H}_9\text{BrN}_2\text{O: C}, 54.00; \text{ H}, 3.14; \text{N}, 9.69. \text{ Found: C}, 53.70; \text{H}, 3.08; \text{N}, 9.37.$

2-(4-Ethylphenyl)-5-aminobenzoxazole (**3e**) (Sener *et al.*, 1987) This compound was prepared by using compounds **1** and **2e** at 180 °C for 2 h. It was obtained as creamy solid in 81 % yield; mp 127–128 °C; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 1.23$ (3H, t, CH₃), 2.70 (2H, q, CH₂), 5.11 (2H, s, NH₂), 6.66 (1H, dd, J = 8.4 Hz, J = 2.0 Hz, H-6), 6.86 (1H, d, J = 1.6 Hz, H-4), 7.41 (3H, t, H-7, H-3', H-5'), 8.05 (2H, d, J = 8.4 Hz, H-2', H-6'); ESIMS *m*/ *z* 239.7 [M + H]⁺ (75), 280.8 [M + H + 41(CH₃CN)]⁺ (100); Anal. Calcd. for C₁₅H₁₄N₂O: C, 75.61; H, 5.92; N, 11.76. Found: C, 75.39; H, 5.97; N, 11.56.

2-(4-Methylphenyl)-5-aminobenzoxazole (**3f**) (Wynne *et al.*, 2009) This compound was prepared by using compounds **1** and **2f** at 170 °C for 1.5 h. It was obtained as creamy solid in 92 % yield; mp 172–174 °C; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 2.40$ (3H, s, CH₃), 5.12 (2H, s, NH₂); 6.66 (1H, dd, J = 8.8 Hz, J = 2.4 Hz, H-6); 6.87 (1H, d, J = 2.4 Hz, H-4), 7.40–7.41 (3H, m, H-7, H-3', H-5'), 8.02 (2H, d, *J*: 8.0, H-2', H-6'); ESIMS *m*/*z* 225.7 [M + H]⁺ (98), 266.7 [M + H + 41(CH₃CN)]⁺ (100); Anal. Calcd. for C₁₄H₁₂N₂O: C, 74.98; H, 5.39; N, 12.49. Found: C, 75.15; H, 5.28; N, 12.02.

2-(4-Methoxyphenyl)-5-aminobenzoxazole (**3g**) (Chen *et al.*, 2011) This compound was prepared by using compounds **1** and **2g** at 120 °C for 2 h. It was obtained as creamy solid in 41 % yield; mp 156–158 °C; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 3.86$ (3H, s, CH₃), 5.09 (2H, s, NH₂), 6.64 (1H, dd, J = 8.8 Hz, J = 2.0 Hz, H-6), 6.85 (1H, d, J = 2.0 Hz, H-4), 7.13 (2H, d, J = 9.2 Hz, H-3', H-5'), 7.38 (1H, d, J = 8.8 Hz, H-7), 8.07 (2H, d, J = 8.4 Hz, H-2', H-6'); ESIMS *m*/*z* 241.6 [M + H]⁺ (100), 282.6 [M + H + 41(CH₃CN)]⁺ (100); Anal. Calcd. for C₁₄H₁₂N₂O₂: C, 69.99; H, 5.03; N, 11.66. Found: C, 69.73; H, 4.83; N, 11.45.

2-Benzyl-5-aminobenzoxazole (**3h**) (Yildiz-Oren *et al.*, 2004a) This compound was prepared by using compounds **1** and **2h** at 180 °C for 1.5 h. It was obtained as creamy solid in 61 % yield; mp 79–81 °C; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 4.22$ (2H, s, CH₂), 5.08 (2H, s, NH₂), 6.59 (1H, dd, J = 9.2 Hz, J = 2.4 Hz, H-6), 6.79 (1H, d, J = 2.4 Hz, H-4), 7.26–7.35 (6H, m, Ar); ESIMS m/z 225.6 [M + H]⁺ (68), 266.6 [M + H + 41(CH₃CN)]⁺ (100); Anal. Calcd. for C₁₄H₁₂N₂O. 0.75 H₂O: C, 70.72; H, 5.72; N, 11.78. Found: C, 70.50; H, 5.90; N, 11.20.

2-(4-Chlorobenzyl)-5-aminobenzoxazole (**3i**) (Yildiz-Oren *et al.*, 2004a) This compound was prepared by using compounds **1** and **2i** at 200 °C for 1.5 h. It was obtained as creamy solid in 75 % yield; mp 83–85 °C; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 4.24$ (2H, s, CH₂), 5.03 (2H, s, NH₂), 6.58 (1H, dd, J = 8.8 Hz, J = 2.0 Hz, H-6), 6.77 (1H, d, J = 2.0 Hz, H-4), 7.55 (1H, d, J = 8.8 Hz, H-7), 7.37–7.43 (4H, m, phenyl); ESIMS *m*/*z* 259.7 [M + H]⁺ (32), 261.7 [M + H + 2]⁺ (10), 300.8 [M + H + 41(CH₃CN)]⁺ (100), 302.7 [M + H + 2 + 41(CH₃CN)]⁺ (35); Anal. Calcd. for C₁₄H₁₁ClN₂O. 0.25 H₂O: C, 63.88; H, 4.40; N, 10.64. Found: C, 63.76; H, 4.42; N, 10.90.

2-(4-Fluorobenzyl)-5-aminobenzoxazole (**3j**) (Oksuzoglu *et al.*, 2007) This compound was prepared by using compounds **1** and **2j** at 200 °C for 1.5 h. It was obtained as creamy solid in 72 % yield; mp 76–77 °C; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 4.23$ (2H, s, CH₂), 5.02 (2H, s, NH₂), 6.57 (1H, dd, J = 8.4 Hz, J = 2.0 Hz, H-6), 6.77 (1H, d, J = 2.4 Hz, H-4), 7.15–7.20 (2H, m, H-3', H-5'); 7.27 (1H, d, J = 8.4 Hz, H-7), 7.37–7.41 (2H, m, H-2', H-6'); ESIMS m/z 243.7 [M + H]⁺ (58), 284.7 [M + H + 41(CH₃CN)]⁺ (100); Anal. Calcd. for C₁₄H₁₁ FN₂O: C, 69.41; H, 4.58; N, 11.56. Found: C, 69.23; H, 4.35; N, 11.48.

2-(4-Bromobenzyl)-5-aminobenzoxazole (3k) (Arisoy *et al.*, 2008) This compound was prepared by using compounds **1** and **2k** at 200 °C for 1.5 h. It was obtained as creamy solid in 89 % yield; mp 101–104 °C; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 4.22$ (2H, s, CH₂), 5.03 (2H, s, NH₂), 6.58 (1H, dd, J = 8.4 Hz, J = 2.0 Hz, H-6), 6.76 (1H, d, J = 2.0 Hz, H-4), 7.27 (1H, d, J = 8.4 Hz, H-7), 7.32 (2H, d, J = 8.0 Hz, H-3', H-5'), 7.55 (2H, d, J = 8.8 Hz, H-2', H-6'); ESIMS *m*/*z* 303.5 [M + H]⁺ (30), 305.5 [M + H + 2]⁺ (40), 346.7 [M + H + 2 + 41(CH₃CN)]⁺ (100); Anal. Calcd. for C₁₄H₁₁BrN₂O: C, 55.47; H, 3.66; N, 9.24. Found: C, 55.80; H, 3.83; N, 9.69.

2-(4-Methylbenzyl)-5-aminobenzoxazole (3I) (Ertan-Bolelli *et al.*, 2014) This compound was prepared by using compounds **1** and **2l** at 180 °C for 1.5 h. It was obtained as creamy solid in 61 % yield; mp 82–84 °C; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 2.27$ (3H, s, CH₃), 4.16 (2H, s, CH₂), 5.01 (2H, s, NH₂), 6.57 (1H, dd, J = 8.4 Hz, J = 2.0 Hz, H-6), 6.76 (1H, d, J = 2.4 Hz, H-4), 7.14 (2H, d, J = 8.0 Hz, H-3', H-5'), 7.22 (2H, d, J = 8.0 Hz, H-2', H-6'), 7.25 (1H, d, J = 8.8 Hz, H-7); ESIMS *m*/*z* 239.7 [M + H]⁺ (50), 280.7 [M + H + 41(CH₃CN)]⁺ (100); Anal. Calcd. for C₁₅H₁₄N₂O. 1 H₂O: C, 70.29; H, 6.29; N, 10.93. Found: C, 70.56; H, 5.99; N, 10.55.

General procedure for the preparation of 5-(4nitrophenylsulfonamido)benzoxazole derivatives (5a-5l)

Pyridine (0.95 mmol) and 4-nitrobenzenesulfonyl chloride (4) (0.52 mmol) were added to a solution of 2-(4-substitutedphenyl)-5-aminobenzoxazole (3a-3l) (0.048 mmol) in dichloromethane (2 mL). The reaction mixture was stirred at the room temperature for 16 h. At the end of the reaction, the residue was filtered and washed with saturated solution of CuSO₄ and NaHCO₃ in water (Wynne *et al.*, 2009) and then recrystallized with a mixture of ethyl acetate/*n*-hexan (1:4). The crystals were dried *in vacuo*. Only compounds **5b**, **5d**, **5g**, **5j**, **5h** and **5i** were published by us in 2014 (Ertan-Bolelli *et al.*, 2014).

2-Phenyl-5-(4-nitrophenylsulfonamido)benzoxazole (**5a**) (Ertan-Bolelli *et al.*, 2014) This compound was prepared by using **3a**. It was obtained as creamy solid in 57 % yield; mp 247–250 °C.

2-(4-Chlorophenyl)-5-(4-nitrophenylsulfonamido)benzoxazole (5b) This compound was prepared by using 3b. It was obtained as creamy solid in 65 % yield; mp 263–265 °C; IR (KBr) v_{max} 3275, 3131–3081, 1528, 1480, 1349, 1314, 1161, 1085 cm⁻¹; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 7.17$ (1H, dd, J = 8.8 Hz, J = 2.0 Hz, H-6), 7.53 (1H, d, J = 2.0 Hz, H-4), 7.66 (2H, d, J = 8.8 Hz, H-3', H-5'), 7.71 (1H, d, J = 9.2 Hz, H-7), 8.02 (2H, d, J = 8.4 Hz, H-2", H-6"), 8.14 (2H, d, J = 8.8 Hz, H-2', H-6'), 8.38 (2H, d, J = 9.2 Hz, H-3", H-5"), 10.75 (1H, s, NH); ¹³C NMR (DMSO-d₆, 400 MHz,): $\delta = 111.51, 112.51, 120.01, 124.58, 124.91,$ 128.29, 128.96, 129.43, 133.92, 136.86, 141.83, 144.43, 147.70, 149.78, 162.39 (15C, Ar-C); ESIMS m/z (%) 428.20 (100) $[M-H]^+$, 430.22 (40) $[M-H + 2]^+$; Anal. Calcd. for C₁₉H₁₂ClN₃O₅S: C 53.09, H 2.81, N 9.78, S 7.46. Found: C 52.92, H 3.08, N 9.84, S 7.52.

2-(4-Fluorophenyl)-5-(4-nitrophenylsulfonamido)benzoxazole (**5c**) (Ertan-Bolelli *et al.*, 2014) This compound was prepared by using **3c**. It was obtained as creamy solid in 46 % yield; mp 230–232 °C.

2-(4-Bromophenyl)-5-(4-nitrophenylsulfonamido)benzoxazole (**5d**) This compound was prepared by using **3d**. It was obtained as creamy solid in 67 % yield; mp 266–267 °C; IR (KBr) v_{max} 3275, 3128, 1526, 1477, 1348, 1312, 1161, 1071; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 7.19$ (1H, d, J = 8.0 Hz, H-6), 7.54 (1H, s, H-4), 7.70 (1H, d, J = 8.8 Hz, H-7), 7.79 (2H, d, J = 8.4 Hz, H-3', H-5'), 8.02–8.06 (4H, m, Ar), 8.39 (2H, d, J = 8.0 Hz, H-3", H-5"), 10.76 (1H, s, NH); ¹³C NMR (DMSO-d₆, 400 MHz,): $\delta = 111.57$, 112.55, 120.08, 124.64, 125.29, 125.87, 128.33, 129.13, 132.40, 133.99, 141.88, 144.48, 147.74, 149.82, 162.56 (15C, Ar–C); ESIMS *m/z* 472.13 [M-H]⁺ (100), 474.13 [M-H + 2]⁺ (95); Anal. Calcd. for C₁₉H₁₂BrN₃O₅S: C, 48.12; H, 2.55; N, 8.86; S, 6.76. Found: C, 47.85; H, 2.86; N, 8.91; S, 6.84.

2-(4-Ethylphenyl)-5-(4-nitrophenylsulfonamido)benzoxazole (**5e**) (Ertan-Bolelli *et al.*, 2014) This compound was prepared by using **3e**. It was obtained as creamy solid in 53 % yield; mp 217–219 °C.

2-(4-Methylphenyl)-5-(4-nitrophenylsulfonamido)benzoxazole (**5f**) (Ertan-Bolelli *et al.*, 2014) This compound was prepared by using **3f**. It was obtained as creamy solid in 43 % yield; mp 244–245 °C.

2-(4-Methoxyphenyl)-5-(4-nitrophenylsulfonamido)ben-

zoxazole (**5g**) This compound was prepared by using **3g**. It was obtained as yellowish solid in 71 % yield; mp 227–230 °C; IR (KBr) ν_{max} 3264, 1526, 1498, 1349, 1308, 1160; ¹H NMR (DMSO-d₆, 400 MHz,): δ = 3.86 (3H, s, CH₃), 7.08 (1H, dd, J = 8.8 Hz, J = 2.0 Hz, H-6), 7.15 (2H, d, J = 8.8 Hz, H-3', H-5'), 7.45 (1H, d, J = 2.0 Hz, H-4), 7.66 (1H, d, J = 8.4 Hz, H-7), 7.98 (2H, d, J = 8.8 Hz, H-2", H-6"), 8.09 (2H, d, J = 9.2 Hz, H-2', H-6'), 8.37 (2H, d, J = 9.2 Hz, H-3", H-5"), 10.68 (1H, s, NH); ¹³C NMR (DMSO-d₆, 400 MHz,): δ = 55.50 (1C, CH₃), 111.19, 112.29, 114.79, 118.39, 119.32, 124.62, 128.32, 129.18, 133.65, 142.18, 144.51, 147.65, 149.81, 162.31, 163.51 (15C, Ar–C); ESIMS *m*/*z* 426.80 [M + H]⁺ (100); Anal. Calcd. for C₂₀H₁₅N₃O₆S: C, 56.47; H, 3.55; N, 9.88; S, 7.54. Found: C, 56.56; H, 3.71; N, 10.09; S, 7.53.

2-Benzyl-5-(4-nitrophenylsulfonamido)benzoxazole (5h) (Ertan-Bolelli *et al.*, 2014) This compound was prepared by using **3h**. It was obtained as creamy solid in 55 % yield; mp 145–146 $^{\circ}$ C.

2-(4-Chlorobenzyl)-5-(4-nitrophenylsulfonamido)benzoxazole (5*i*) (Ertan-Bolelli *et al.*, 2014) This compound was prepared by using **3i**. It was obtained as creamy solid in 40 % yield; mp 179–181 °C.

2-(4-Fluorobenzyl)-5-(4-nitrophenylsulfonamido)benzoxazole (**5j**) This compound was prepared by using **3j**. It was obtained as creamy solid in 53 % yield; mp 151–153 °C; IR (KBr) v_{max} 3252, 3128–3039, 2924, 1527, 1480, 1348, 1308, 1223, 1161; ¹H NMR (DMSO-d₆, 400 MHz,) $\delta = 4.30$ (2H, s, CH₂), 7.06 (1H, dd, J = 8.8 Hz, J = 2.0 Hz, H-6), 7.15–7.19 (2H, m, H-3', H-5'), 7.39–7.42 (3H, m, Ar), 7.58 (1H, d, J = 8.4 Hz, H-7), 7.96 (2H, d, J = 9.2 Hz, H-2", H-6"), 8.35 (2H, d, J = 8.8 Hz, H-3", H-5"), 10.64 (1H, s, NH); ¹³C NMR (DMSO-d₆, 400 MHz,): δ = 33.12 (1C, CH₂), 111.05, 112.46, 115.20, 115.41, 119.29, 124.54, 128.24, 130.94–131.07, 133.29, 141.25, 144.43, 147.82, 149.74, 160.03, 162.45 (15C, Ar–C); ESIMS *m*/*z* 428.48 [M + H]⁺ (100); Anal. Calcd. for C₂₀H₁₄FN₃O₅S: C, 56.20; H, 3.30; N, 9.83; S, 7.50. Found: C, 55.95; H, 3.52; N, 9.88; S, 7.57.

2-(4-Bromobenzyl)-5-(4-nitrophenylsulfonamido)benzoxazole (5k) This compound was prepared by using 3k. It was obtained as creamy solid in 41 % yield; mp 187–189 °C; IR (KBr) v_{max} 3115, 3057, 2856, 1525, 1488, 1348, 1312, 1165, 1068; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 4.30$ (2H, s, CH₂), 7.06 (1H, dd, J = 8.8 Hz, J = 2.0 Hz, H-6), 7.33 (2H, d, J = 8.0 Hz, H-3', H-5'), 7.39 (1H, d, J = 2.0 Hz, H-4), 7.54 (2H, d, J = 8.8 Hz, H-2', H-6'), 7.58 (1H, d, J = 8.8 Hz, H-7), 7.96 (2H, d, J = 9.2 Hz, H-2", H-6"), 8.35 (2H, d, J = 8.8 Hz, H-3", H-5"), 10.64 (1H, s, NH); ¹³C NMR (DMSO-d₆, 400 MHz,): $\delta = 33.31$ (1C, CH₂), 111.07, 112.45, 119.32, 120.26, 124.55, 128.24, 131.35, 131.42, 133.31, 134.24, 141.22, 144.43, 147.81, 149.74, 166.18 (15C, Ar-C) ESIMS m/z 488.37 [M + H]⁺ (85), 490.37 [M + H + 2]⁺ (100); Anal. Calcd. for C₂₀H₁₄BrN₃O₅S: C, 49.19; H, 2.89; N, 8.61; S, 6.57. Found: C, 49.24; H, 3.17; N, 8.65; S, 6.68.

2-(4-Methylbenzyl)-5-(4-nitrophenylsulfonamido)benzoxazole (51) This compound was prepared by using 31. It was obtained as creamy solid in 57 % yield; mp 158–160 °C; IR (KBr) v_{max} 3258, 3076, 2924–2790, 1527, 1472, 1349, 1310, 1166; ¹H NMR (DMSO-d₆, 400 MHz,) $\delta = 2.27$ (3H, s, CH₃), 4.23 (2H, s, CH₂), 7.03 (1H, dd, J = 8.4 Hz, J = 2.4 Hz, H-6), 7.14 (2H, d, J = 8.0 Hz, H-3', H-5', 7.22 (2H, d, J = 8.4 Hz, H-2', H-6'), 7.37 (1H, d, J = 2.0 Hz, H-4), 7.56 (1H, d, J = 8.8 Hz, H-7), 7.94 (2H, d, J = 8.8 Hz, H-2'', H-6''), 8.34 (2H, d, J = 8.8 Hz,H-3", H-5"), 10.63 (1H, s, NH); ¹³C NMR DMSO-d₆: $\delta = 20.59$ (1C, CH₃), 33.72 (1C, CH₂), 111.08, 112.51, 119.32, 124.61, 128.30, 128.92, 129.18, 131.80, 133.35, 136.22, 141.36, 144.52, 147.88, 149.80, 166.82 (15C, Ar-C); ESIMS m/z 424.70 [M + H]⁺ (100); Anal. Calcd. for C₂₁H₁₇N₃O₅S: C, 59.57; H, 4.05; N, 9.92; S, 7.57. Found: C, 59.44; H, 4.12; N, 9.98; S, 7.44.

General procedure for the preparation of 5-(4aminophenylsulfonamido)benzoxazole derivatives (6a, 6e-h, 6l)

0,5 mmol 2-(4-Substitutedphenyl/benzyl)-5-(4-nitrophenyl sulfonamido)benzoxazole (**5a**, **5e**–**h**, **5l**) in ethanol (50 mL) was reduced by hydrogenation using 40 psi of H₂ and 10 % Pd–C (40 mg) until uptake of H₂ ceased (Zheng *et al.*, 2007). The catalyst was filtered on a bed of celite and washed with ethanol, and the filtrate was concentrated in

vacuo. The crude product was purified by recrystallization with a mixture of ethyl acetate/*n*-hexan (1:4) to obtain **6a**, **6e–h**, **6l**. The crystals were dried in vacuo. All the compounds are new except **6e** (Ertan-Bolelli *et al.*, 2014).

2-Phenyl-5-(4-aminophenylsulfonamido)benzoxazole

(6a) This compound was prepared by using 5a. It was obtained as white solid in 41 % yield; mp 266–269 °C; IR (KBr) v_{max} 3343, 3247, 1486, 1307, 1142; ¹H NMR (DMSO-d₆, 400 MHz,) $\delta = 5.98$ (2H, s, NH₂), 6.51 (2H, d, J = 8.8 Hz, H-3", H-5"), 7.11 (1H, dd, J = 8.4 Hz, J = 2.0 Hz, H-6), 7.38 (2H, d, J = 8.8 Hz, H-2", H-6"), 7.43 (1H, d, J = 2.0 Hz, H-4), 7.60–7.66 (4H, m, Ar), 8.16 (2H, dd, J = 7.6 Hz, J = 1.6 Hz, H-2', H-6'), 9.97 (1H, s, NH); ¹³C NMR DMSO-d₆: $\delta = 110.87$, 110.97, 112.49, 118.70, 123.90, 126.20, 127.16, 128.68, 129.20, 131.91, 135.59, 141.72, 146.83, 152.78, 162.99 (15C, Ar–C); ESIMS m/z 366.40 [M + H]⁺ (100); Anal. Calcd. for C₁₉H₁₅N₃O₃S. 0.1 CH₃COOC₂H₅: C, 62.27; H, 4.26; N, 11.23; S, 8.57. Found: C, 61.96; H, 4.20; N, 11.43; S, 8.64.

2-(4-Ethylphenyl)-5-(4-aminophenylsulfonamido)benzoxazole (**6e**) (Ertan-Bolelli *et al.*, 2014) This compound was prepared by using **5e**. It was obtained as white solid in 60 % yield; mp 251–254 °C.

2-(4-Methylphenyl)-5-(4-aminophenylsulfonamido)ben-

zoxazole (**6f**) This compound was prepared by using **5f**. It was obtained as white solid in 33 % yield; mp 289–291 °C; IR (KBr) v_{max} 3406–3335, 3108, 1466, 1309, 1187; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 2.40$ (3H, s, CH₃), 5.97 (2H, s, NH₂), 6.52 (2H, d, J = 9.2 Hz, H-3", H-5"), 7.09 (1H, dd, J = 8.8 Hz, J = 2.0 Hz, H-6), 7.39 (5H, t, Ar), 7.62 (1H, d, J = 8.8 Hz, H-7), 8.04 (2H, d, J = 8.4 Hz, H-2', H-6'), 9.93 (1H, s, NH); ESIMS *m*/*z* 380.80 [M + H]⁺ (100); Anal. Calcd. for C₂₀H₁₇N₃O₃S. 0.1 CH₃COOC₂H₅: C, 63.11; H, 4.62; N, 10.82; S, 8.26. Found: C, 62.75; H, 4.61; N, 11.06; S, 8.23.

2-(4-Methoxyphenyl)-5-(4-aminophenylsulfonamido)benzoxazole (6g) This compound was prepared by using 5g. It was obtained as white solid in 43 % yield; mp 273–274 °C; IR (KBr) v_{max} 3396–3329, 3056, 2838, 1465, 1310, 1187; ¹H NMR (DMSO-d₆, 400 MHz,) $\delta = 3.86$ $(3H, s, CH_3)$, 5.97 $(2H, s, NH_2)$, 6.51 (2H, d, J = 8.8 Hz), H-3", H-5"), 7.06 (1H, dd, J = 8.4 Hz, J = 1.6 Hz, H-6), 7.14 (2H, d, J = 8.8 Hz, H-3', H-5'), 7.37 (3H, d, J = 8.8 Hz, H-2", H-6"), 7.60 (1H, d, J = 9.2 Hz, H-7), $8.09 (2H, d, J = 8.8 \text{ Hz}, \text{H-2'}, \text{H-6'}), 9.91 (1H, s, \text{NH}); {}^{13}\text{C}$ NMR DMSO-d₆: $\delta = 55.42$ (1C, CH₃), 110.66, 110.68, 112.47, 114.67, 118.16, 118.54, 123.95, 128.67, 129.05, 135.41, 141.92, 146.75, 152.74, 162.11, 163.13 (15C, Ar-C); ESIMS m/z 396.70 [M + H]⁺ (100); Anal. Calcd. for C₂₀H₁₇N₃O₄S: C, 60.75; H, 4.33; N, 10.63; S, 8.11. Found: C, 60.79; H, 4.31; N, 10.75; S, 8.01.

2-Benzyl-5-(4-aminophenylsulfonamido)benzoxazole

(6h) This compound was prepared by using 5h. It was obtained as creamy solid in 68 % yield; mp 154–156 °C; IR (KBr) v_{max} 3410–3344, 3154, 3031, 1455, 1311, 1186; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 4.25$ (2H, s, CH₂), 5.92 (2H, s, NH₂), 6.47 (2H, d, J = 8.4 Hz, H-3", H-5"), 6.99 (1H, dd, J = 8.8 Hz, J = 2.4 Hz, H-6), 7.25–7.32 (8H, m, Ar), 7.47 (1H, d, J = 8.4 Hz, H-7), 9.83 (1H, s, NH); ¹³C NMR DMSO-d₆: $\delta = 34.05$ (1C, CH₂), 110.55, 110.91, 112.45, 118.13, 123.92, 126.95, 128.55, 128.62, 128.96, 134.92, 135.06, 141.08, 146.99, 152.71, 166.16 (15C, Ar–C); ESIMS *m*/*z* 380.80 [M + H]⁺ (100); Anal. Calcd. for C₂₀H₁₇N₃O₃S. 0.1 CH₃COOC₂H₅: C, 63.11; H, 4.62; N, 10.82; S, 8.26. Found: C, 62.67; H, 4.43; N, 11.15; S, 8.40.

2-(4-Methylbenzyl)-5-(4-aminophenylsulfonamido)ben-

zoxazole (61) This compound was prepared by using 51. It was obtained as creamy solid in 34 % yield; mp 192–193 °C; IR (KBr) v_{max} 3463–3412, 3216, 1477, 1311, 1183; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 2.27$ (3H, s, CH₃), 4.21 (2H, s, CH₂), 5.95 (2H, s, NH₂), 6.49 (2H, d, J = 8.8 Hz, H-3", H-5"), 7.01 (1H, dd, J = 9.2 Hz, J = 2.4 Hz, H-6), 7.14 (2H, d, J = 8.0 Hz, H-3', H-5'), 7.22 (2H, d, J = 8.0 Hz, H-2', H-6'), 7.30 (1H, d, J = 2.0 Hz, H-4), 7.33 (2H, d, J = 9.2 Hz, H-2", H-6"), 7.49 (1H, d, J = 8.4 Hz, H-7), 9.85 (1H, s, NH); ¹³C NMR DMSO-d₆: $\delta = 20.61$ (1C, CH₃), 33.74 (1C, CH₂), 110.59, 110.99, 112.52, 118.19, 123.99, 128.69, 128.90, 129.17, 131.91, 135.11, 136.15, 141.16, 147.06, 152.77, 166.42 (15C, Ar–C); ESIMS m/z 394.80 [M + H]⁺ (100); Anal. Calcd. for C₂₁H₁₉N₃O₃S: C, 64.11; H, 4.87; N, 10.68; S, 8.15. Found: C, 63.98; H, 4.95; N, 10.67; S, 7.98.

General procedure for the preparation of 5-(4aminophenylsulfonamido)benzoxazole derivatives (6b–d, 6i–k)

The reduction of the nitro group for compounds bearing halogen atoms was carried out with iron powder and ammonium chloride. 2-(4-Substitutedphenyl/benzyl)-5-(4nitrophenylsulfonamido)benzoxazole (5b-d, 5i**k**) (4.65 mmol) was solved in methanol (20 mL), and then iron powder (0.78 g, 13.65 mmol) and an aqueous solution of ammonium chloride (1.3 g, 23.3 mmol, 20 mL) were added in it. The reaction mixture was stirred at 70 °C for 2.5 h. The reaction mixture was cooled and filtered to remove the inorganic residues and then evaporated (Habens et al., 2005). The precipitate was dissolved in ethyl acetate and extracted with water and then recrystallized with a mixture of ethyl acetate/*n*-hexan (1:4) to obtain 6b-d, 6i-k. The crystals were dried in vacuo. All the compounds are new.

2-(4-Chlorophenyl)-5-(4-aminophenylsulfonamido)ben-

zoxazole (**6b**) This compound was prepared by using **5b**. It was obtained as creamy solid in 31 % yield; mp 314–315 °C; IR (KBr) v_{max} 3419–3349, 3124, 1482, 1311, 1160, 1093; ¹H NMR (DMSO-d₆, 400 MHz,) $\delta = 5.92$ (2H, s, NH₂), 6.47 (2H, d, J = 8.4 Hz, H-3", H-5"), 7.08 (1H, dd, J = 8.8 Hz, J = 2.0 Hz, H-6), 7.34 (2H, d, J = 8.8 Hz, H-2", H-6"), 7.39 (1H, d, J = 1.6 Hz, H-4), 7.60–7.65 (3H, m, Ar), 8.11 (2H, d, J = 8.0 Hz, H-2', H-6'), 9.92 (1H, s, NH); ESIMS *m*/*z* 400.70 [M + H]⁺ (100), 402.70 [M + H + 2]⁺ (40); Anal. Calcd. for C₁₉ H₁₄ClN₃O₃S: C, 57.07; H, 3.53; N, 10.51; S, 8.02. Found: C, 57.10; H, 3.63; N, 10.78; S, 7.90.

2-(4-Fluorophenyl)-5-(4-aminophenylsulfonamido)ben-

zoxazole (**6c**) This compound was prepared by using **5c**. It was obtained as creamy solid in 33 % yield; mp 280–282 °C; IR (KBr) v_{max} 3408–3341, 3106, 1465, 1313, 1237, 1162; ¹H NMR (DMSO-d₆, 400 MHz,) $\delta = 5.96$ (2H, s, NH₂), 6.51 (2H, d, J = 8.8 Hz, H-3", H-5"), 7.11 (1H, dd, J = 8.8 Hz, J = 2.4 Hz, H-6), 7.38 (2H, d, J = 8.8 Hz, H-2", H-6"), 7.42–7.47 (3H, m, Ar), 7.64 (1H, d, J = 8.8 Hz, H-7), 8.19–8.22 (2H, m, H-2', H-6'), 9.95 (1H, s, NH); ¹³C NMR DMSO-d₆: $\delta = 110.82$, 110.96, 112.48, 116.33, 116.56, 118.67, 122.86, 122.89, 123.88, 128.67, 129.77, 129.87, 135.63, 141.68, 146.86, 152.77, 162.18, 162.91, 165.40 (15C, Ar–C); ESIMS *m/z* 384.80 [M + H]⁺ (100); Anal. Calcd. for C₁₉H₁₄FN₃O₃S: C, 59.52; H, 3.68; N, 10.96; S, 8.36. Found: C, 59.42; H, 3.76; N, 11.22; S, 8.32.

2-(4-Bromophenyl)-5-(4-aminophenylsulfonamido)ben-

zoxazole (**6d**) This compound was prepared by using **5d**. It was obtained as white solid in 46 % yield; mp 304–305 °C; IR (KBr) v_{max} 3349, 1477, 1310, 1160, 1071; ¹H NMR (DMSO-d₆, 400 MHz,) $\delta = 5.97$ (2H, s, NH₂), 6.51 (2H, d, J = 8.8 Hz, H-3", H-5"), 7.12 (1H, dd, J = 8.8 Hz, J = 2.0 Hz, H-6), 7.38 (2H, d, J = 9.2 Hz, H-2", H-6"), 7.43 (1H, d, J = 2.4 Hz, H-4), 7.65 (1H, d, J = 8.4 Hz, H-7), 7.81 (2H, d, J = 8.4 Hz, H-3', H-5'), 8.07 (2H, d, J = 8.4 Hz, H-2', H-6'), 9.97 (1H, s, NH); ¹³C NMR DMSO-d₆: $\delta = 111.55$, 111.62, 114.98, 119.40, 126.70, 127.70, 129.18, 129.75, 132.48, 135.90, 142.26, 147.47, 150.50, 163.56 (15C, Ar–C); ESIMS *m/z* 444.80 (100) [M + H]⁺; 446.80 (80) [M + H + 2]⁺; Anal. Calcd. for C₁₉H₁₄BrN₃O₃S: C 51.36, H 3.18, N 9.46, S 7.22. Found: C 51.19, H 3.28, N 9.71, S 7.06.

2-(4-Chlorobenzyl)-5-(4-aminophenylsulfonamido)ben-

zoxazole (**6**i) This compound was prepared by using **5**i. It was obtained as creamy solid in 32 % yield; mp 176–178 °C; IR (KBr) v_{max} 3464–3327, 3218, 1478, 1311, 1183, 1091; ¹H NMR (DMSO-d₆, 400 MHz,) δ = 4.30

(2H, s, CH₂), 5.95 (2H, s, NH₂), 6.50 (2H, d, J = 8.8 Hz, H-3", H-5"), 7.03 (1H, dd, J = 8.4 Hz, J = 2.0 Hz, H-6), 7.31–7.40 (7H, m, Ar), 7.51 (1H, d, J = 8.8 Hz, H-7), 9.87 (1H, s, NH); ¹³C NMR DMSO-d₆: $\delta = 33.25$ (1C, CH₂), 110.57, 110.86, 112.44, 118.15, 123.88, 128.46, 128.62, 130.95, 131.69, 133.92, 135.11, 141.01, 146.95, 152.70, 165.82 (15C, Ar–C); ESIMS m/z 414.70 [M + H]⁺ (100), 416.70 [M + H + 2]⁺ (40); Anal. Calcd. for C₂₀H₁₆ ClN₃O₃S: C, 58.04; H, 3.90; N, 10.15; S, 7.75. Found: C, 58.13; H, 3.89; N, 10.34; S, 7.51.

2-(4-Fluorobenzyl)-5-(4-aminophenylsulfonamido)benzoxazole (**6j**) This compound was prepared by using **5j**. It was obtained as creamy solid in 77 % yield; mp 187–190 °C; IR (KBr) v_{max} 3464–3327, 3216, 1478, 1311, 1182, 1223; ¹H NMR (DMSO-d₆, 400 MHz,): δ = 4.28 (2H, s, CH₂), 5.95 (2H, s, NH₂), 6.50 (2H, d, J = 8.8 Hz, H-3", H-5"), 7.03 (1H, dd, J = 8.8 Hz, J = 2.4 Hz, H-6), 7.15–7.19 (2H, m, H-3', H-5'), 7.31–7.42 (5H, m, Ar), 7.50 (1H, d, J = 8.8 Hz, H-7), 9.86 (1H, s, NH); ¹³C NMR DMSO-d₆: δ = 33.14 (1C, CH₂), 110.56, 110.89, 112.45, 115.18, 115.39, 118.15, 123.90, 128.63, 130.95–131.09, 135.09, 141.06, 146.98, 152.71, 160.01, 162.42, 166.09 (15C, Ar–C); ESIMS m/z 398.70 [M + H]⁺ (100); Anal. Calcd. for C₂₀H₁₆FN₃O₃S: C, 60.44; H, 4.06; N, 10.57; S, 8.07. Found: C, 60.36; H, 3.98; N, 10.80; S, 7.95.

2-(4-Bromobenzyl)-5-(4-aminophenylsulfonamido)benzoxazole (6k) This compound was prepared by using 5k. It was obtained as creamy solid in 66 % yield; mp 170–172 °C; IR (KBr) v_{max} 3478–3380, 3308, 1490, 1302, 1187, 1074; ¹H NMR (DMSO-d₆, 400 MHz,): $\delta = 4.28$ $(2H, s, CH_2)$, 5.95 $(2H, s, NH_2)$, 6.50 (2H, d, J = 8.4 Hz)H-3", H-5"), 7.03 (1H, dd, J = 8.4 Hz, J = 2.0 Hz, H-6), 7.31-7.35 (5H, m, Ar), 7.51 (1H, d, J = 8.8 Hz, H-7), 7.54 (2H, d, J = 8.4 Hz, H-2', H-6'), 9.86 (1H, s, NH); ¹³C NMR DMSO-d₆: $\delta = 33.32$ (1C, CH₂), 110.57, 110.87, 112.45, 118.17, 120.20, 123.89, 128.63, 131.32, 131.40, 134.34, 135.11, 141.02, 146.97, 152.71, 165.75 (15C, Ar-C); ESIMS m/z 458.70 $[M + H]^+$ (100), 460.70 $[M + H + 2]^+$ (100); Anal. Calcd. for C₂₀H₁₆BrN₃O₃S: C, 52.41; H, 3.52; N, 9.17; S, 6.99. Found: C, 52.21; H, 3.62; N, 9.53; S, 6.86.

Antimicrobial evaluation

Standard strains of *P. aeruginosa* ATCC 27853, *S. aureus* ATCC 29213, *Escherichia coli* ATCC 25922, *Enterococcus faecalis* ATCC 29212 and *M. tuberculosis* H37RV ATCC 27294 and clinical isolates of these microorganisms, *Candida albicans* ATCC 10231 ve *Candida krusei* ATCC 6258, were included in the study.

Microdilution method

Bacterial susceptibility testing was performed according to the guidelines of CLSI M100-S181. MHB was added to each well of the microplates. The bacterial suspensions used for inoculation were prepared at 10^5 CFU/mL by diluting fresh cultures at McFarland 0.5 density (10^7 CFU/ mL). Suspensions of the bacteria at 10^5 CFU/mL concentration were inoculated to the twofold diluted solution of the compounds. A 10-µL bacteria inoculum was added to each well of the microplates. There were 10^4 CFU/mL bacteria in the wells after inoculations. Microplates were incubated at 37 °C overnight (CLSI, 2006a).

Fungal susceptibility testing was performed according to the guidelines of CLSI M27-A32. RPMI-1640 medium with L-glutamine (Sigma) buffered to pH 7 with MOPS was added to each well of the microplates. The colonies were suspended in sterile saline, and the resulting suspension was adjusted to McFarland 0.5 density (10^6 CFU/mL). A working suspension was prepared by a 1:100 dilution followed by a 1:20 dilution of the stock suspension. A 10 µL of this suspension at 10^3 CFU/mL was inoculated to the twofold diluted solution of the compounds. Microplates were incubated at 35 °C for 24–48 h (CLSI, 2006b).

After incubation, the lowest concentration of the compounds that completely inhibits macroscopic growth was determined and reported as minimum inhibitory concentrations (MIC). All solvents and diluents, pure microorganisms and pure media were used in control wells. All the experiments were done in 3 parallel series.

Microplate Alamar Blue Assay (MABA)

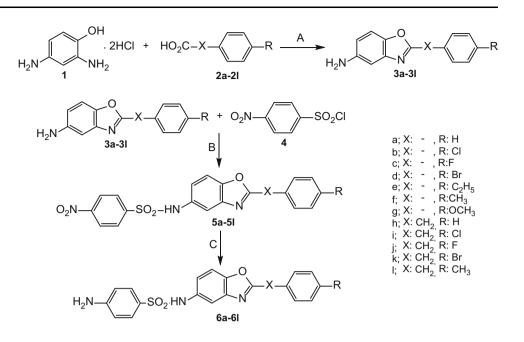
Mycobacterium tuberculosis H37Rv ATCC 27294 (American Type Culture Collection) was subcultured on Middlebrook 7H11 agar (Becton–Dickinson). Suspensions were prepared in 0.04 % (vol/vol) Tween 80–0.2 % bovine serum albumin so that their turbidities matched that of a McFarland no. 1 turbidity standard. Isoniazid (INH) and ethambutol (EMB) were obtained from Sigma. Stock solutions of INH and EMB were prepared in deionized water.

Sterile deionized water was added to all outer-perimeter wells of sterile 96-well plates to minimize evaporation of the medium in the test wells during incubation. One hundred microliters of six drug solutions were added to the wells in rows B to G in columns 2 and 3 by using a multichannel pipette, 100 μ L was transferred from column 3 to column 4, and the contents of the wells were mixed well. Identical serial 1:2 dilutions were continued through column 10, and 100 μ L of excess medium was discarded from the wells in column 10. Final drug concentration ranges were as follows: for INH, 0.031 to 8.0 μ g/mL and for

EMB, 0.5 to 128 µg/mL. One hundred microliters of M. tuberculosis inoculum was added to the wells in rows B to G in columns 2-11. The plates were sealed with Parafilm and were incubated at 37 °C for 5 days. Fifty microliters of a freshly prepared 1:1 mixture of 10X Alamar Blue reagent and 10 % Tween 80 was added to well B11. The plates were reincubated at 37 °C for 24 h. If well B11 turned pink, the reagent mixture was added to all wells in the microplate. The microplates were resealed with Parafilm and were incubated for an additional 24 h at 37 °C, and the colors of all wells were recorded. A blue color in the well was interpreted as no growth, and a pink color was scored as growth. A few wells appeared violet after 24 h of incubation, but they invariably changed to pink after another day of incubation and thus were scored as growth. The MIC was defined as the lowest drug concentration which prevented a color change from blue to pink (Franzblau et al., 1998).

Molecular docking

The crystal structure of the InhA was retrieved from the Protein Data Bank (PDB ID: 3FNE) (Freundlich et al., 2009). Accelrys Discovery Studio 3.5 (Accelrys Inc., 2012) software was used for preparation of protein and ligands. The target protein was taken, the ligand was extracted, hydrogens were added, and their positions were optimized using the all-atom CHARMm forcefield and the Adopted Basis set Newton-Raphson (ABNR) method available in Discovery Studio 3.5 protocol until the root mean deviation (RMS) gradient was <0.05 kcal/mol/Å². The minimized protein was defined as the receptor using the binding site module. The binding site was defined from the cavity finding method which was modified to accommodate all the important interacting residues in the active site of the enzyme. Binding sphere for 3FNE (27.49, 3.71, 10.20, 13.16) was selected from the active site using the binding site tools. The most antimycobacterial active compounds 3a, 3c, 3f and the triclosan derivative, which is the ligand of **3FNE.PDB** crystal structure, were sketched, and allatom CHARMm forcefield parameterization was assigned and then minimized using the ABNR method as described above. Conformational searches of the ligands were carried out using a simulated annealing molecular dynamics (MD) approach. The ligands were heated to a temperature of 700 K and then annealed to 200 K. CDOCKER (Wu et al., 2003) method was performed by using Discovery Studio 3.5 (Accelrys Inc., 2012). The protein is held rigid, while the ligands are allowed to be flexible during refinement. The docking and scoring methodology was first validated by docking of triclosan derivative. The docked position of triclosan derivative overlaps well with the crystal structure position, with an RMSD of 0.53 Å. Afterward, molecular Scheme 1 Synthetic pathway of 2-(substitutedbenzyl/phenyl)-5-aminobenzoxazole (**3a–3l**) and 5-(4-nitro/aminophenylsulfonamido)benzoxazole (**5a– 5l**, **6a–6l**) derivatives. *Reagents and conditions:* A) PPA, 170-200 °C, 1.5-2.5 h; B) Pyridine and dichloromethane, RT, 16 h; C) Pd–C/H₂, RT (for **6a, 6e–h, 6l**); methanol, iron powder and ammonium chloride, 70 °C, 2.5 h (for **6b–d**, **6i–k**)



docking studies were performed on the antimycobacterial active compounds (**3a**, **3c**, **3f**). Finally, all docked poses were scored by applying Analyze Ligand Poses subprotocol, and binding energies were calculated by applying Calculate Binding Energy subprotocol in Discovery Studio 3.5 by using in situ ligand minimization step (ABNR method) and implicit solvent model (GBMV). The lowest binding energy was taken as the best-docked conformation of the compounds for the macromolecule.

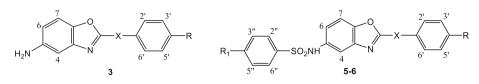
Results and discussion

Chemistry

In this study, 17 new 2-(4-substitutedphenyl)-5-(4-nitro/ aminophenylsulfonamido)benzoxazole derivatives (5b, 5d, 5g, 5j, 5k, 5l and 6a-6d, 6f-6l) were synthesized for the first time. For preparing them at first, 5-amino-2-(4-substitutedphenyl/benzyl)benzoxazoles (3a-3l) were obtained by heating appropriate acid with 2,4-diaminophenol in PPA (Yildiz-Oren et al., 2004b; Sener et al., 1987; Wynne et al., 2009). After, 5-amino-2-(4-substitutedphenyl/benzyl)benzoxazole derivatives (3a-3l) and 4-nitrobenzenesulfonyl chloride (4) were treated in pyridine and dichloromethane to prepare 5-(4-nitrophenyl sulfonamido)benzoxazole derivatives (5a-5l) (Wynne et al., 2009). Afterward, 5-(aminophenyl/benzylsulfonamido)benzoxazoles (6a-6l) were obtained by reduction using Pd-C/H₂ (Zheng et al., 2007) or iron powder/ammonium chloride (Habens et al., 2005) of compounds 5a-5l as given in Scheme 1 (Ertan-Bolelli et al., 2014). The newly synthesized structures were supported by spectral data. The IR, ¹H NMR, ¹³C NMR, mass spectra and elemental analysis results are in agreement with the proposed structures. According to the spectroscopic data of the compounds 5b, 5d, 5g, 5j, 5k, 5l and 6a-6d, 6f-6l, the IR spectra showed that characteristic NH (SO₂NH) stretching bands were in the 3106-3308 cm⁻¹ region. In the ¹H NMR spectra of the compounds 6a-6d, 6f-6l, the signals of NH₂ protons were observed at 5.92–5.98 ppm as a singlet band; the signal of NH (SO₂ NH) proton of the compounds 5b, 5d, 5g, 5j, 5k, 5l and 6a-6d, 6f-6l was observed at 9.83-10.76 ppm. Besides, benzylic CH₂ and aromatic CH₃ protons appeared at 4.21-4.30 ppm and 2.27-2.40 ppm as a singlet, respectively. Moreover, all the aromatic protons were observed at 6.47-8.39 ppm. ¹³C NMR spectra of the compounds were appropriate to their formulas. Mass spectra of the compounds showed $M^+ + H$ peaks in accordance with their formulas, since the electrospray ionization method was employed. On the other hand, the results of the elemental analyses (C, H, N, S) were within ±0.4 % of the calculated amounts.

In vitro antimicrobial evaluation

All of the previously and newly synthesized 5-amino-2-(4-substitutedphenyl/benzyl)benzoxazoles (**3a–3l**), 2-(4-substitutedphenyl)-5-(4-nitrophenylsulfonamido)benzoxazoles (**5a–5l**), 2-(4-substitutedphenyl)-5-(4-aminophenylsulfonamido)benzoxazoles (**6a–6l**) were tested for their in vitro antimicrobial activity against *S. aureus* ATCC 29213 and *E. faecalis* ATCC 29212 as Gram-positive bacteria, *E. coli* ATCC 25922 and *Pseudomonas aeruginosa* ATCC 27853 as GramTable 1 Structures of the tested benzoxazoles and their in vitro antimicrobial activities as MIC values ($\mu g/mL$)



Compo	unds			Microorganisms												
Code	Х	R	R ₁	Gram-negative				Gram	Gram-positive				Fungi		Mycobacteria	
				E.c.	E.c.*	<i>P.a.</i>	<i>P.a.</i> *	S.a.	S.a.*	E.f.	<i>E.f.</i> *	<i>C.a.</i>	<i>C.k.</i>	M.t.	<i>M.t.</i> ²	
3a	_	Н	_	64	64	32	32	64	64	64	32	64	64	8	8	
3b	-	Cl	-	64	64	32	64	64	64	64	32	32	64	16	8	
Bc	_	F	_	64	64	32	64	64	64	64	32	32	64	8	8	
3d	_	Br	_	64	64	32	64	64	64	64	32	32	64	8	32	
Be	_	C_2H_5	-	64	64	32	64	64	64	64	32	32	64	16	8	
3f	_	CH ₃	-	64	64	32	64	64	64	64	32	64	64	8	8	
Bg	_	OCH ₃	_	64	64	32	64	64	64	64	32	64	64	64	8	
3h	CH_2	Н	_	64	64	32	64	64	64	64	32	64	64	32	32	
Bi	CH_2	Cl	-	64	64	32	64	64	64	64	32	64	64	16	8	
3j	CH_2	F	_	64	64	32	64	64	64	64	32	64	64	16	32	
3k	CH_2	Br	_	64	64	32	64	64	64	64	32	64	64	16	16	
31	CH_2	CH ₃	_	64	64	32	64	64	64	64	32	64	64	32	64	
5a	_	Н	NO_2	128	128	128	32	256	128	128	128	128	64	64	32	
5b	_	Cl	NO_2	128	128	128	32	256	128	128	128	128	64	64	32	
5c	_	F	NO_2	128	128	128	64	256	128	128	128	128	64	64	64	
5d	_	Br	NO_2	128	128	128	64	256	128	128	128	128	64	64	32	
5e	_	C_2H_5	NO ₂	128	128	128	64	256	128	128	128	128	64	64	16	
5f	_	CH ₃	NO_2	64	128	128	32	128	128	128	128	128	64	64	64	
g	_	OCH ₃	NO_2	128	128	128	64	256	128	128	128	64	64	64	16	
5h	CH_2	Н	NO ₂	128	128	128	64	128	128	128	128	64	64	32	16	
5i	CH ₂	Cl	NO ₂	128	128	128	64	128	128	128	128	32	64	32	16	
5j	CH ₂	F	NO ₂	128	128	128	64	128	128	128	128	256	64	32	16	
5k	CH ₂	Br	NO_2	128	128	128	64	128	128	128	128	128	128	32	16	
51	CH ₂	CH ₃	NO ₂	128	128	128	64	256	128	128	128	64	64	32	16	
ba	_ 2	Н	NH ₂	128	128	128	32	128	128	128	128	128	64	64	32	
ób	_	Cl	NH ₂	128	128	128	32	256	128	128	128	128	64	64	64	
óc	_	F	NH ₂	128	128	128	32	256	128	128	64	128	64	64	64	
ód	_	Br	NH ₂	128	128	128	64	256	128	128	128	128	64	64	64	
ie i	_	C_2H_5	NH ₂	128	128	128	64	256	128	128	128	128	64	64	16	
óf	_	CH ₃	NH ₂	128	128	128	64	256	128	128	128	128	64	64	64	
óg	_	OCH ₃	NH ₂	128	128	128	64	64	128	128	128	64	64	64	64	
b b	CH_2	Н	NH ₂	128	128	128	64	256	128	128	128	64	64	32	16	
ii ii	CH ₂	Cl	NH ₂	128	128	128	64	256	128	128	128	64	64	16	16	
j	CH ₂	F	NH ₂	128	128	128	64	256	128	128	128	128	128	32	16	
y ók	CH ₂ CH ₂	Br	NH ₂	128	128	128	64	256 256	128	128	128	64	128	16	16	
51	CH ₂ CH ₂	CH ₃	NH ₂	128	128	128	64	256 256	128	128	128	128	128	16	16	
		C113	1112	<2	<2	<2	8	<2	<2	8	8	-	-	-		
Meropenem Ampicillin			<2 8	<2 64	<2	o _	<2	<2 32	°<2	о 4	_	_	_	_		
Ceftriaxone <2 128					- 16	- 32	<2 8	52 64	<2 -	4	_	_	_	-		

Table 1 continued

Compour	nds	Microorganisms													
Code	Х	R	R_1	Gram-negative				Gram-positive				Fungi		Mycobacteria	
				E.c.	E.c.*	<i>P.a.</i>	<i>P.a.</i> *	<i>S.a.</i>	S.a.*	E.f.	E.f.*	C.a.	<i>C.k.</i>	M.t.	<i>M.t.</i> *
Gentamicin				<2	128	<2	32	<2	64	4	4	_	_	-	_
Tetracycl	line	<2	128	16	32	1	32	2	64	_	_	-	_		
Trimetho	prim-Sulfa	<2	128	32	32	<2	32	<2	<2	_	_	-	_		
Sulfamethoxazole				16	128	_	_	<2	8	64	64	_	_	8	8
Ofloxacin				<2	16	<2	32	<2	<2	<2	<2	_	_	-	_
Ciprofloxacin				<2	16	<2	16	<2	<2	<2	<2	_	_	-	_
Fluconazole				_	_	_	_	_	_	_	_	1	64	-	_
Amphotericin B				_	_	_	_	_	_	_	_	< 0.25	0.5	-	_
Isoniazide				_	_	_	_	_	_	_	_	_	_	< 0.25	< 0.25
Ethambut	Ethambutol				_	_	_	_	_	_	_	_	_	2	2

E.c., E. coli ATCC 25922; E.c.*, E. coli isolate (include ESBL); P.a., Pseudomonas aeruginosa ATCC 27853; P.a.*, P. aeruginosa isolate (rezistant of gentamicin); S.a., Staphylococcus aureus ATCC 29213; S.a.*, S. aureus isolate (MRSA); E.f., Enterococcus faecalis ATCC 29212, E.f.*, E. faecalis isolate (rezistant of vancomycin); C.a., Candida albicans ATCC 10231; C.k., C. krusei ATCC 6258; M.t., M. tuberculosis H37RV ATCC 27294; M.t.*, M. tuberculosis isolate

negative bacteria strains, M. tuberculosis H37RV ATCC 27294 as Mycobacteria and their drug-resistant isolates C. albicans ATCC 10231 and C. krusei ATCC 6258 as fungus. The standard drugs meropenem, ampicillin, ceftriaxone, gentamicin, tetracycline, trimethoprim-sulfamethoxazole, sulfamethoxazole, ofloxacin and ciprofloxacin for antibacterial activity, isoniazid and ethambutol for antimycobacterial activity, and fluconazole and amphotericin B for antifungal activity were also screened under identical conditions for quality control and comparison. The minimum inhibitory concentration (MIC) values were determined by twofold serial dilution technique in Mueller-Hinton Broth (MHB) and Sabouraud Dextrose Agar (SDA) for the antibacterial and antifungal assays, respectively. Microplate Alamar Blue Assay (MABA) in Middlebrook 7H11 Agar was used in order to obtain the MIC values for antimycobacterial effect. All of the biological results are given in Table 1.

Microbiological results indicated that previously and newly synthesized compounds possessed a broad spectrum of activity against the tested microorganisms at MIC values between 8 and 256 μ g/mL.

According to the obtained data, all of the 2-(4-substitutedbenzyl/phenyl)-5-aminobenzoxazole derivatives (**3a**– **3l**) showed better antibacterial activity against *E. coli*, *S. aureus* and *E. faecalis* with their drug-resistant isolates and *P. aeruginosa* than their sulfonamide derivatives except compounds **5f**, **6g** and **6c**. The tested compounds indicated significant inhibitory effect against the Gramnegative enterobacter drug-resistant isolate of *P. aeruginosa* with MIC values of 32–64 µg/mL. Among the compounds, derivatives **3a**, **5a**, **5b**, **5f** and **6a–6c** showed good activities against *P. aeruginosa* clinical isolate, which is effective in nosocomial infections, comparable to that of ceftriaxone, gentamicin, tetracycline, trimethoprim-sulfamethoxazole and ofloxacin.

Moreover, **3a–31** compounds were found to be more potent than the other compounds against drug-resistant isolate of *E. faecalis* with MIC value of 32 μ g/mL. Even they were more effective than the standard drugs tetracycline and sulfamethoxazole. It could be noticed that an amine group at the 5th position of benzoxazole ring was very important for improving the activity.

On the other hand, although compounds **3b–3e** and **5i** indicated significant activity against *C. albicans* with MIC value of 32 μ g/mL, they exhibited less effect than the standard drugs. Surprisingly, antifungal activities against *C. krusei* of all derivatives except **5k** and **6j–6l** are comparable to fluconazole.

Furthermore, most of the compounds exhibited very important activity against *M. tuberculosis* and its clinical isolate with MIC values between 8 and 64 μ g/mL comparing with other human pathogenic microbes. Surprisingly, compounds **3a**, **3c**, **3d** and **3f** against *M. tuberculosis* and compounds **3a–3c**, **3e–3g** and **3i** against drug-resistant *M. tuberculosis* showed the same activity with standard drug sulfamethoxazole with a MIC value of 8 μ g/mL.

Molecular docking studies

The NADH-dependent InhA has been validated as the primary molecular target of the frontline antitubercular drugs. As shown in Table 1, compounds **3a**, **3c** and **3f** were found to be significantly active against both *M. tuberculosis* H37RV ATCC 27294 and its clinical isolate at the MIC value of 8 μ g/mL. Here, these compounds (**3a**, **3c** and **3f**)

Table 2 Docking results

Comp. code	Binding energy (kcal/mol)	Interacted residues*
3a	-16.5561	Gly96 (2.25 Å), Phe97, Met98, Met103, Phe149, Tyr158, Met161, Pro193, Ala198, Met199, Ile215, Leu218, NAD ⁺ (2.75 Å)
3c	-18.0974	Gly96 (2.45 Å), Phe97, Met98, Met103, Phe149, Tyr158, Met161, Pro193, Thr196, Ala198, Met199, Ile215, Leu218, NAD ⁺ (2.63 Å)
3f	-15.7779	Gly96 (2.44 Å), Phe97, Met103, Phe149, Tyr158 ^[b] , Met161, Pro193, Ala198, Met199, Ile215, Leu218, Glu219, NAD ⁺ (2.58 Å)
Triclosan derivative	-11.9466	Gly96, Phe97, Met98, Met103, Phe149 ^[b] , Met155, Pro156, Ala157, Tyr158 ^[a] (1.86 Å), Met161, Lys165 ^[b] , Pro193, Try196, Ala198, Met199, Ile215, Leu218, Glu219, NAD ⁺ (2.15 Å)

* Van der Waals contact distance <4 Å

Bold: H bonds; ^[a] $\pi - \pi$ interactions; ^[b] $\pi - \sigma$ interactions

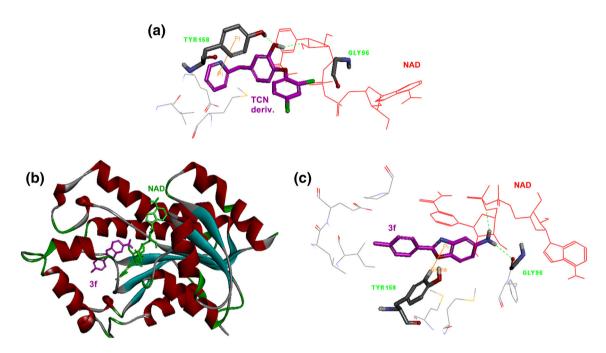


Fig. 1 a Docked position of triclosan (TCN) derivative (C atoms are purple): Hydroxyl group has H bonds with Tyr158 and NAD⁺ (red), and pyridine ring has pi–pi interaction with Tyr158. b): Docked position of **3f** (purple) with NAD⁺ (green) in 3FNE c): Docked

position of 3f (C atoms are purple): Amino group has H bonds with Gly96 and NAD⁺ (red), and oxazole ring has pi–sigma interaction with Tyr158

were docked into active site of InhA (**PDB ID: 3FNE**) by using CDOCKER method (Wu *et al.*, 2003; Accelrys Inc., 2012) in order to understand ligand–protein interactions. Firstly, one of the triclosan derivatives (**3FNE.PDB** ligand), which are the known specific inhibitor of InhA, was docked. After then, molecular docking studies were performed for the compounds **3a**, **3c** and **3f**. The molecular docking studies of the examined compounds showed that their binding to InhA active site with the position and orientation was found to be very close to the result of the crystal structure of triclosan derivative complex in InhA. All the docking scores are given in Table 2. According to the docking studies, triclosan derivative revealed H bonds with Tyr158 and co-factor NAD⁺, which are in accordance with the X-ray structure binding features (**3FNE**). As for the most active benzoxazole derivatives **3a**, **3c** and **3f**, the amino group at the 5th position of their benzoxazole ring interacted with an active site residue Gln96 and co-factor NAD⁺. Unlike other compounds, derivative **3f** showed also pi–sigma interaction between Tyr158 and the oxazole ring of it as shown in Fig. 1. Binding energies of these compounds **3a**, **3c**, **3f** and triclosan derivative are -16.5561, -18.0974, -15.7779 and -11.9466, respectively. *In vitro* studies demonstrated that **3a**, **3c** and **3f** exhibited promising antitubercular activity, and the docking results were also correlated with the experimental data.

Conclusion

In this study, the synthesis of 17 novel 2-substituted-5-(4nitro/aminophenylsulfonamido)benzoxazole derivatives (5b, 5d, 5g, 5j, 5k, 5l and 6a-6d, 6f-6l) were described by using 2-(4-substitutedphenyl)-5-aminobenzoxazole (3a-31). In addition, in vitro antimicrobial activities against human pathogenic microbes of all previously and newly synthesized benzoxazole derivatives (3a-3l, 5a-5l, 6a-6l) were investigated. Microbiological results indicated that all the tested compounds possessed a broad spectrum of activity. Generally, 2-(4-substitutedphenyl)-5-aminobenzoxazoles (3a-3l) were found to be more potent than their sulfonamido derivatives for all tested microorganisms. In particular, compounds 3a, 3b, 3c, 3e, 3f, 3g and 3i revealed very important effect against drug-resistant isolate of M. tuberculosis with MIC value of 8 µg/mL. The structure-activity relationship showed that amino group at the 5th position together with phenyl ring instead of benzyl ring on the 2nd position of benzoxazole played very important role for improving antituberculosis activity.

According to in vitro microbiological studies, 2-phenyl-5aminobenzoxazole (3a), 2-(4-fluorophenyl)-5-aminobenzoxazole (3c) and 2-(4-methylphenyl)-5-aminobenzoxazole (3f) exhibited significantly antitubercular activity against both M. tuberculosis and its drug-resistant isolate. For that reason, molecular docking of compounds 3a, 3c and 3f to the active site of InhA, which has been validated as the primary molecular target of the frontline antitubercular drugs, was carried out using CDOCKER method (Accelrys Inc., 2012) in order to understand ligand-protein interactions. Previous reports revealed that the catalytic residue Tyr158 and cofactor NAD⁺ played important role for the activity of InhA (Lu et al., 2010; Shrinivas et al., 2015). Molecular docking studies showed that promising antitubercular compounds **3a**, 3c and 3f directly interacted with InhA enzyme by binding to Tyr158 and/or co-factor NAD⁺, and these results were correlated with the experimental data. It can be concluded that compounds 3a, 3c and 3f could be considered as scaffolds for designing new potent antitubercular drugs.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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