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> Synthesis and Evaluation of a New Series of 3,5-bis ((5-bromo-6-methyl-2-*t*aminopyrimidin-4-yl)thio)-4*H*-1,2,4-triazol-4-amines and their Cyclized Products "Pyrimidinylthio pyrimidotriazolothiadiazines" as 15- Lipoxygenase Inhibitors

Running tittle: 3,5-bis(substituted pyrimidinylthio) triazolamines

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Abstract

A series of new 3,5-bis((5-bromo-6-methyl-2-*t*-aminopyrimidin-4-yl)thio)-4*H*-1,2,4-triazol-4amines and their cyclized products "pyrimidinylthio pyrimidotriazolothiadiazines" were designed, synthesized, and evaluated as potential inhibitors of 15-lipoxygenase (15-LO). Their syntheses started by initial condensation of 2:1 equivalents of pyrimidine with triazole and subsequent nucleophilic displacement of the chlorine atoms with secondary amines and finally cyclocondensation in the presence of NaNH₂. The compounds **4d** and **4f** showed the best IC₅₀ of 15-LO inhibition (IC₅₀ = 9 and 12 μ M, respectively). Compounds **4a-g** were docked into 15-LO. We suggest that the hydrogen bonds in quaternary nitrogen of piperazine ring of compounds **4d** and **4f** appears to play major role in lipoxygenase inhibition by this set of synthesized analogs and hydrophobic nature of this protein's binding site should be considered in ongoing investigations.

Introduction

Triazoles and their derivatives play important roles in medicinal, agricultural and industrial fields (1-3). These compounds are known to exhibit antibacterial and antifungal (4), analgesic (5), anticancer and antimicrobial (6), anticonvulsant (7), anti-inflammatory properties (8). Some of the modern day drugs with triazole nucleus are Ribavirin (antiviral agent), Alprazolam (anxiolytic agent), Fluconazole, Itraconazole (antifungal agent) and Rizatriptan (antimigrane

agent). The commonly known systems are triazoles fused with pyridines (9), pyridazines (10), pyrazines (11), imidazoles (12) and triazines (13).

Triazolothiadiazines are a class of heterocyclic compounds that have been described as antiinflammatory (14), anticandidal (15), analgesic (16), antibacterial, anticancer (17) and antiviral agents (18). Moreover Pyrimidine-fused heterocycles often appear as the core of biologically active compounds (19-21) such as anticancer (22), antiviral (23), anti-inflammatory (24), RET Kinase Inhibitors (25). A literature survey reveals that there are not many examples of triazolethiadiazine fused with pyrimidine reported. Bside more recently, 15-lipoxygenase (15-LO) has emerged as an attractive target for therapeutic intervention (26). 15-LO has been implicated in the progression of certain cancers (27, 28) and chronic obstructive pulmonary disease (COPD) (27). Evidence for the inhibition of 15-LO in the treatment of vascular disease is, however, most compelling (29). Both transgenic and knockout studies implicate a role for 15-LO in atherogenesis (30,31). The enzyme is abundantly expressed in macrophages residing within the atherosclerotic lesion (26). Keeping this in mind, and in continuation of our recent studies (32, 33) on the enzyme inhibitory activity of heterocyclic compound against soybean 15lipoxygenase, we considered the synthesis of new derivatives of 3,5-bis((5-bromo-6-methyl-2-taminopyrimidin-4-yl)thio)-4H-1,2,4-triazol-4-amines and their cyclized products pyrimidinylthio pyrimidotriazolothiadiazines.

Method and material

Chemistry

Melting points were taken on an Electrothermal type 9100 melting point apparatus and are uncorrected. The IR spectra were obtained in KBr disks on an Avatar 370 FT-IR Thermo-Nicolet spectrometer (v_{max} in cm⁻¹). The ¹H NMR spectra were recorded on a Bruker AC at 100 MHz,

400 MHz using tetramethylsilane (TMS) as an internal reference. Chemical shift values were given in ppm, at room temperature using $CDCl_3$, DMSO- d_6 or CD_3COCD_3 as a solvent; chemical shifts were reported in parts per million (ppm) downfield from TMS; multiplicities were abbreviated as: s: singlet; d: doublet; q: quartet; m: multiplet; br s: broad singlet. The Mass spectra were obtained on a Varian Mat CH-7 and Agilent 5973 instruments at 70 eV. Elemental analysis was performed on a Thermo Finnigan Flash EA microanalyzer.

3,5-bis((5-bromo-2-chloro-6-methylpyrimidin-4-yl)thio)-4H-1,2,4-triazol-4-amine (3):

To a solution of 4-amino-1,2,4-triazolidine-3,5-dithione **2** (10 mmol, 1.48) and Et₃N (26 mmol, 3.2 ml) in (40 ml) DMF was added 5-Bromo-2,4-dichloro-6-methylpyrimidine **1** (20 mmol, 4.84 gr). The solution was stirred for 2h at room temperature and then was poured into water (150 mL). The resulting precipitate was filtered off and washed with water (2 × 50 mL) and diethyl ether (20 mL), respectively. After drying the solid, 5.50 gr of **3** were obtained as a crude product. Yield: 98%, m.p. 246-247°C; ¹H NMR (100 MHz, DMSO-d₆): δ 2.56 (s, 6H, 2CH₃), 6.12 ppm (s, 2H, NH₂, D₂O exchangeable); IR: v 3332, 3252, 2958, 1523, 755cm⁻¹; MS (*m/z*) 556 (M⁺), 558 (M⁺+2); Anal. Calcd. for C₁₂H₈Br₂Cl₂N₈S₂: C, 25.78; H, 1.44; N, 20.04; S, 11.47. Found: C, 25.97; H, 1.48; N, 20.35; S, 11.28.

The general procedure for preparation (4a–g) compounds

The appropriate secondary amine (30 mmol) was added to a stirred mixture of compound **3** (10 mmol, 5.59 g) in acetonitrile` (100 mL), and the solution was heated under reflux for 30-60 min. Progress of the reaction was monitored by TLC using chloroform:methanol (30:1). The solid obtained on cooling was filtered and recrystallized from ethanol.

3,5-bis((5-bromo-6-methyl-2-morpholinopyrimidin-4-yl)thio)-4H-1,2,4-triazol-4-amine(4a):

White powder; yield: 89%; m.p. 248-249 °C; ¹H NMR (100 MHz, CDCl₃): δ 2.46 (s, 6H, 2CH₃), 3.48-3.57 (m, 8H, 4CH₂N), 3.57-3.62 (m, 8H, 4CH₂O), 5.02 ppm (s, 2H, NH₂, D₂O exchangeable); IR (KBr disc): v 3328, 3271, 2973,2916,2847, 1543, 770cm⁻¹.; MS (*m/z*) 658

(M⁺), 660 (M⁺+2); Anal. Calcd. for C₂₀H₂₄Br₂N₁₀O₂S₂: C, 36.37; H, 3.66; N, 21.21; S, 9.71. Found: C, 36.55; H, 3.62; N, 21.51; S, 9.36%.

3,5-bis((5-bromo-6-methyl-2-(piperidin-1-yl)pyrimidin-4-yl)thio)-4H-1,2,4-triazol-4-amine (4b):

White powder; yield: 91%; m.p. 242 °C; ¹H NMR (100 MHz, CDCl₃): δ 1.20-1.82 (m, 12H, 6CH₂), 2.40 (S, 6H, 2CH₃), 3.21-3.60 (m, 8H, 4CH₂N), 5.02 ppm (s, 2H, NH₂, D₂O exchangeable); IR (KBr disc): v 3338, 3260, 2931, 2851, 1555, 768cm⁻¹.; MS (*m/z*) 654 (M⁺), 656 (M⁺+2); Anal. Calcd. for C₂₂H₂₈Br₂N₁₀S₂: C, 40.25; H, 4.30; N, 21.34; S, 9.77. Found: C, 40.48; H, 4.46; N, 21.18; S, 9.91%.

3,5-bis((5-bromo-6-methyl-2-(4-methylpiperidin-1-yl)pyrimidin-4-yl)thio)-4H-1,2,4-triazol-4-amine (4c):

White powder; yield: 89%; m.p. 179 °C; ¹H NMR (100 MHz, CDCl₃): δ 0.91 (d, J = 6 Hz, 6H, 2CH₃), 1.10-1.30 (m, 2H, 2CH) 1.36-1.89 (m, 8H, 4CH₂), 2.41 (s, 6H, 2CH₃), 2.51-2.92 (m, 4H, equatorial hydrogens of 2CH₂N), 3.92-4.41 (m, 4H, axial hydrogens of 2CH₂N), 5.03 (s, 2H, NH₂, D₂O exchangeable); IR (KBr disc): v 3329, 3256, 2947, 2922 , 1547, 1515 , 772cm⁻¹; MS (*m/z*) 682 (M⁺), 684 (M⁺+2); Anal. Calcd. for C₂₄H₃₂Br₂N₁₀S₂: C, 42.11; H, 4.71; N, 20.46; S, 9.37. Found: C, 42.53; H, 4.78; N, 20.47; S, 9.51%.

3,5-bis((5-bromo-6-methyl-2-(4-methylpiperazin-1-yl)pyrimidin-4-yl)thio)-4H-1,2,4-triazol-4-amine (4d):

White powder; yield: 88%; m.p. 267-268°C; ¹H NMR (400 MHz, CDCl₃): δ 2.27 (s, 6H, 2CH₃) 2.32 (app. t, 8H, 4CH₂N), 2.38 (s, 6H, 2CH₃), 3.46 (app. t, 8H, 4CH₂N), 5.13 ppm (s, 2H, NH₂, D₂O exchangeable); IR (KBr disc): v 3354, 3268, 2966, 2937, 1547, 775cm⁻¹.; MS (*m/z*) 684

(M⁺), 686 (M⁺ + 2); Anal. Calcd. for C₂₂H₃₀Br₂N₁₂S₂: C, 38.49; H, 4.40; N, 24.48; S, 9.34 Found: C, 38.59; H, 4.31; N, 24.44; S, 9.05%.

3,5-bis((5-bromo-6-methyl-2-(4-phenylpiperazin-1-yl)pyrimidin-4-yl)thio)-4H-1,2,4-triazol-4-amine (4e):

White powder; yield: 87%; m.p. 199-201 °C; ¹H NMR (100 MHz, CDCl₃): δ 2.50 (s, 6H, 2CH₃), 3.01 (app. t, 8H, 4CH₂N), 3.60 (app. t, 8H, 4CH₂N), 5.10 (s, 2H, NH₂, D₂O exchangeable), 6.75-7.02 (m, 6H, aromatic), 7.12-7.41 ppm (m, 4H, aromatic); IR (KBr disc): v 3333, 3252, 2953, 2908, 1598, 1548, 758cm⁻¹.; MS (*m/z*) 808 (M⁺), 810 (M⁺+2); Anal. Calcd. for C₃₂H₃₄Br₂N₁₂S₂: C, 47.41; H, 4.23; N, 20.73; S, 7.91. Found: C, 47.63; H, 4.34; N, 20.75; S, 7.77%.

3,5-bis((5-bromo-6-methyl-2-(4-ethylpiperazin-1-yl)pyrimidin-4-yl)thio)-4H-1,2,4-triazol-4amine (4f):

White powder; yield: 91%; m.p. 220 °C; ¹H NMR (100 MHz, CDCl₃): δ 1.16 (t, J=7Hz, 6H, 2CH₃), 2.44 (s, 6H, 2CH₃), 2.50-2.57 (m, 12H, 4CH₂N,2CH₂), 3.56 (app. t, 8H, 4CH₂N), 5.12 ppm (s, 2H, NH₂, D₂O exchangeable); IR (KBr disc): v 3358, 3280, 2966, 2945, 2810, 1545, 767cm⁻¹.; MS (*m*/*z*) 712 (M⁺), 714 (M⁺+2); Anal. Calcd. for C₂₄H₃₄Br₂N₁₂S₂: C, 40.34; H, 4.80; N, 23.52; S, 8.97 Found: C, 40.73; H, 4.77; N, 23.10; S, 8.37%.

3,5-bis((5-bromo-6-methyl-2-(pyrrolidin-1-yl)pyrimidin-4-yl)thio)-4H-1,2,4-triazol-4-amine (4g):

White powder; yield: 90%; m.p. 245-246 °C; ¹H NMR (400 MHz, CDCl₃): δ 1.75–1.86 (m, 8H, 4CH2), 2.47 (s, 6H, 2CH3), 3.00 (m, 4H, 2CH₂N), 3.47 (m, 4H, 2CH₂N), 5.20 (s, 2H, NH₂, D2O exchangeable); IR (KBr disc): v 3330, 3268, 2966, 2870, 1557, 769cm–1; MS (m/z) 626 (M⁺), 628 (M⁺+2). Anal. Calcd for C₂₀H₂₄Br₂N₁₀S₂: C, 38.23; H, 3.85; N, 22.29; S, 10.21. Found: C, 38.18; H, 3.87; N, 21.97; S, 10.01%.

A mixture of each of compounds **4a-g** (10 mmol), NaNH₂ (30mmol) in dry acetonitrile (100 mL) was heated under reflux for 15-60 min. Progress of the reaction was monitored by TLC using chloroform:methanol (15:1) or ethylacetate:n-hexane (15:9). The mixture was cooled and the solvent was removed under reduced pressure. Then, a solution of acetic acid (1 mL) in water (20 mL) was added to the residue and the resulting precipitant was filtered off and recrystallized from ethanol.

4-(5-bromo-4-methyl-6-((6-methyl-8-morpholino-5H-pyrimido[5,4-e][1,2,4]triazolo[3,4b][1,3,4]thiadiazin-3-yl)thio)pyrimidin-2-yl)morpholine (5a):

White powder; yield: 75%; m.p.> 300 °C (dec.); ¹H NMR (100 MHz, CD₃COCD₃); δ 2.20 (s, 3H, CH₃), 2.40 (s, 3H, CH₃) 3.35-3.50 (m, 8H, 4CH₂N), 3.50-3.70 (m, 8H, 4CH₂O), 7.85 ppm (br s, 1H, NH, D₂O exchangeable); IR (KBr disc): v 3260, 3166, 2958, 1611cm⁻¹; MS (*m/z*) 578 (M⁺), 580 (M⁺+2) Anal. Calcd. for C₂₀H₂₃BrN₁₀O₂S₂: C, 41.45; H, 4.00; N, 24.17; S, 11.07 Found: C, 41.54; H, 4.10; N, 23.93; S, 10.89%.

3-((5-bromo-6-methyl-2-(piperidin-1-yl)pyrimidin-4-yl)thio)-6-methyl-8-(piperidin-1-yl)-5H-pyrimido[5,4-e][1,2,4]triazolo[3,4-b][1,3,4]thiadiazine (5b):

White powder; yield: 71%; m.p. 270 °C (dec.); ¹H NMR (400 MHz, CDCl₃): δ 1.30-1.70 (m, 12H, 6CH₂), 2.32 (s, 3H, CH₃), 2.43 (s, 3H, CH₃), 3.38 (app. t, 4H, 2CH₂N), 3.70 (t, J=5.2Hz, 4H, 2CH₂N) 7.80 ppm (br s, 1H, NH, D₂O exchangeable); IR (KBr disc): v 3260, 3170, 2931, 1631cm⁻¹.; MS (*m*/*z*) 574 (M⁺), 576 (M⁺+2); Anal. Calcd. for: C₂₂H₂₇BrN₁₀S₂: C, 45.91; H, 4.73; N, 24.34; S, 11.14. Found: C, 45.33; H, 4.81; N, 24.06; S, 10.88%.

3-((5-bromo-6-methyl-2-(4-methylpiperidin-1-yl)pyrimidin-4-yl)thio)-6-methyl-8-(4-

methylpiperidin-1-yl)-5H-pyrimido[5,4-e][1,2,4]triazolo[3,4-b][1,3,4]thiadiazine (5c):

White powder; yield: 74%; m.p. 220 °C; ¹H NMR (100 MHz, CDCl₃): δ 0.80- 1.10 (m, 6H, 2CH₃), 1.31-1.85 (m, 10H, 2CH and 4CH₂), 2.30 (s, 3H, CH₃), 2.42 (s, 3H, CH₃), 2.50-2.90 (m, 4H, equatorial hydrogens of 2CH₂N), 4.10-4.70 (m, 4H, axial hydrogens of 2CH₂N), 7.80 ppm (br. s, 1H, NH, D₂O exchangeable); IR (KBr disc): v 3280, 2946, 2923, 1635cm⁻¹.; MS (*m/z*) 602 (M⁺), 604 (M⁺+2); Anal. Calcd. for C₂₄H₃₁BrN₁₀S₂: C, 47.76; H, 5.18; N, 23.21; S, 10.62. Found: C, 47.37; H, 5.09; N, 23.04; S, 10.36%.

3-((**5**-bromo-6-methyl-2-(**4**-methylpiperazin-1-yl)pyrimidin-4-yl)thio)-6-methyl-8-(**4**methylpiperazin-1-yl)-5H-pyrimido[5,4-e][1,2,4]triazolo[3,4-b][1,3,4]thiadiazine (5d): White powder; yield: 75%; m.p.> 300 °C (dec.); ¹H NMR (100 MHz, CDCl₃); δ 2.20-280 (m, 20H, 4CH₃, 4CH₂N,), 3.40-3.90 (m, 8H, 4CH₂N), 8.12 ppm (br s, 1H, NH, D₂O exchangeable); IR (KBr disc): v 3284, 2934, 1585 cm⁻¹.; MS (*m/z*) 604 (M⁺), 606 (M⁺+2) Anal. Calcd. for C₂₂H₂₉BrN₁₂S₂: C, 43.63; H, 4.83; N, 27.76; S, 10.59. Found: C, 43.31; H, 4.93; N, 27.45; S, 10.61%.

3-((5-bromo-6-methyl-2-(4-phenylpiperazin-1-yl)pyrimidin-4-yl)thio)-6-methyl-8-(4phenylpiperazin-1-yl)-5H-pyrimido[5,4-e][1,2,4]triazolo[3,4-b][1,3,4]thiadiazine (5e):

White powder; yield: 73%; m.p. 240 °C (dec);¹H NMR (100 MHz, CDCl₃): δ 2.36 (s, 3H, CH₃), 2.47 (s, 3H, CH₃), 2.85-3.31 (m, 8H, 4CH₂N), 3.45-4.01 (m, 8H, 4CH₂N), 6.69-7.08 (m, 6H, aromatic), 7.08-7.41(m, 4H, aromatic), 8.11 ppm (br s, 1H, NH, D₂O exchangeable); IR (KBr disc): v 3245, 2953, 2913, 1598cm⁻¹.; MS (*m/z*) 728 (M⁺), 730 (M⁺+2); Anal. Calcd. for C₃₂H₃₃BrN₁₂S₂: C, 52.67; H, 4.56; N, 23.03; S, 8.79. Found: C, 52.39; H, 4.91; N, 23.18; S, 8.43%.

3-((5-bromo-2-(4-ethylpiperazin-1-yl)-6-methylpyrimidin-4-yl)thio)-8-(4-ethylpiperazin-1yl)-6-methyl-5H-pyrimido[5,4-e][1,2,4]triazolo[3,4-b][1,3,4]thiadiazine (5f):

White powder; yield: 70%; m.p. 260 °C (dec); ¹H NMR (100 MHz, CDCl₃): δ 0.92- 1.27 (m, 6H, 2CH₃), 2.06-2.69 (m, 18H, 2CH₃, 4CH₂N, 2CH₂), 3.29-3.98 (m, 8H, 4CH₂N) 8.50 ppm (br. s, 1H, NH, D₂O exchangeable); IR (KBr disc): v 3248, 2966, 1642 cm⁻¹.; MS (*m/z*) 632 (M⁺), 634 (M⁺+2) Anal. Calcd. for C₂₄H₃₃BrN₁₂S₂: C, 45.49; H, 5.25; N, 26.53; S, 10.12. Found: C, 45.23; H, 5.40; N, 26.66; S, 10.05%.

3-((5-bromo-6-methyl-2-(pyrrolidin-1-yl)pyrimidin-4-yl)thio)-6-methyl-8-(pyrrolidin-1-yl)-5H-pyrimido[5,4-e][1,2,4]triazolo[3,4-b][1,3,4]thiadiazine (5g):

White powder; yield: 77%; m.p.> 300 °C (dec.); ¹H NMR (100 MHz, CDCl₃): δ 1.68–1.92 (m, 8H, 4CH₂), 2.10 (s, 3H, CH₃), 2.35 (s, 3H, CH₃), 2.98–3.42 (m, 8H, 4CH₂N), 8.12 (s, 1H, NH, D2O exchangeable). IR (KBr disc): v 3268, 3190, 2968, 1615cm–1; MS (*m/z*) 546 (M+), 548 (M⁺+2); Anal. Calcd for C₂₀H₂₃BrN₁₀S₂: C, 43.87; H, 4.23; N, 25.58; S, 11.71. Found: C, 43.52; H, 4.26; N, 25.60; S, 11.39%.

Biology

Linoleic acid and two assay solutions (A and B) were prepared in advance. Solution A was 50 mM 3-(dimethylamino) benzoic acid (DMAB) in a 100 mM phosphate buffer (pH 7.0). Solution B was a mixture of 10 mM 3-methyl-2-benzothiazolinone (MBTH) 3 mL, hemoglobin (5 mg/mL, 3 mL) in 50 mM phosphate buffer at pH 5.0 (25 mL). A linoleic acid solution was prepared by mixing 5 mg of linoleic acid with 50 mg Tween 20 in 3 mL water and then diluting with KOH 100 mM to a final volume of 5 mL. In the standard assay, the sample in ethanol (25 mL), soybean 15-lipoxygenase (SLO) (4000 units/mL in 50 mM phosphate buffer pH 7.0; 25 mL) and phosphate buffer pH 7.0 (50 mM; 900 mL) were mixed in a test tube and preincubation was carried out for 5 min at room temperature. A control test was done with the same volume of ethanol. After the preincubation, linoleic acid solution (50 mL) was added to start the

peroxidation reaction, and, 7 min later, solution A (270 mL) and then solution B (130 mL) was added to start the color formation. Further 5 min later, 200 mL of a 2% SDS solution was added to terminate the reaction. The absorbance at 598 nm was compared with control test. SLO and other chemicals were purchased from Sigma, Aldrich and Merck Co. respectively.

Docking

The mode of interaction between synthesized ligands and lipoxygenase was investigated by docking. 2D structure of chemicals was drowned in ChemDraw ultera 8.0 software and 3D structures were prepared by Hyperchem 7 software (http://www.hyper.com/), using molecular mechanic force filed pre-optimization followed by AM1 semiempirical calculation. The X-ray crystal structure of lipoxygenase (PDB ID: 1IK3) was downloaded from the Protein Data Bank at the Research Collaboratory for Structural Bioinformatics (www.rcsb.org). Further preparations such as polar hydrogen addition and water molecules elimination were done by MOE software. Synthesized chemicals were docked into the binding site of protein by MOE software. All atoms within a 5 Å around the co-crystallized ligand in crystal coordinates of lipoxygenase was selected as binding site. The docking simulations were performed using Triangle matcher placement algorithm in combination with London dG scoring function and forcefield as refinement method. For each compound, the top-score docking poses were selected for final ligand-target interaction analysis using LigX module in MOE Software. Validation of docking methodology was first evaluated by docking of co-crystalized ligand into the lipoxygenase binding site.

Results and discussion

Chemistry

The starting material 5-bromo-2,4-dichloro-6-methyl-pyrimidine **1** was prepared according to published method (34). Treatment of compound **1** with 4-amino-1,2,4-triazolidine-3,5-dithione **2**

at room temperature afforded intermediate **3** in which the C-4 chlorine atoms in the pyrimidine rings were replaced selectively by two sulfhydryl groups of ditiol **2**. The foregoing compound **2** was prepared according to the published procedure (35). Subsequent reaction of compound **3** with various secondary amines led to the selective replacement of the C-2 chlorine atoms in the pyrimidine rings to produce the corresponding diheteroaryl sulfide intermediates **4a-g**. The latter compounds subsequently underwent an intramolecular S_NAr reaction in the presence of NaNH₂ in boiling acetonitrile to give the desired tricyclic pyrimidinylthio pyrimido[5,4-e]triazolo[5,1b][1,3,4]thiadiazines **5a-g.** (Scheme 1)

Our attempt for replacement of Br atom of second pyrimidine ring in different solvents and temperature was failed. Structural preference of **5a-g** may be attributed to ring strain of two fused pyrimidothiadiazine rings with triazol. The yield of compounds **4a-g** were increased between 30-40% by changing ethanol to acetonitrile as a solvent. The structure of newly synthesized compounds **5a-g** was confirmed by recording their IR, ¹H NMR, elemental analysis and mass spectra. For example, the IR spectrum of compound **4b** exhibited the stretching vibration bands at 3338 and 3260cm⁻¹ due to symetrical and asymetrical vibrations of NH₂ group while the IR spectrum of compound **5b** showed a band at 3260cm⁻¹ for NH vibration. The ¹H NMR spectrum of the asymetrical cyclized product **5b** showed two singlet at δ 2.32 and 2.44 ppm in CDCl₃ belonging to methyl groups of the pyrimidine moieties whereas, the ¹H NMR spectrum of the symmetrical precursor **4b** showed a singlet at δ 2.46 ppm in CDCl₃ for methyl groups of the pyrimidine moieties. The ¹H NMR spectrum of the precursor **4b** showed the NH₂ signal at δ 5.02 ppm in CD₃Cl which was removed on adding D₂O. However, the ¹H NMR spectrum of the cyclized product **5b** did not show this signal and instead an exchangeable broad singlet peak at δ

7.80 ppm confirming the occurrence of heterocyclisation to **5b**. The mass spectrum of **5b** showed a molecular ion peak at m/z 574 (M^+) corresponding to the molecular formula $C_{22}H_{27}BrN_{10}S_2$.

Biology

The compounds **5a-g** and **4a-g** were evaluated in vitro to determine their activity on 15-LO inhibition, and their inhibitory potencies were compared to 4-methyl-2-(4-methyl piperazinyl)pyrimido[4,5-b]benzothiazine (4-MMPB) with IC₅₀ 18 μ M (31). Compounds **4d** and **4f** were the most potent inhibitors of 15-LO at a concentration of 9 and 12 μ M respectively. It was interesting to see that pyrrolidine, morpholine, piperidine, methylpiperidine, and phenylpiperazine analogs didn't show inhibitory activity on 15-LO (IC₅₀ > 500 μ M) (Table 1). Also Log Dose-response 15-LO activity curves for biologically active compounds **4d**, **4f** and reference **4-MMPB** has been shown in Figure 1.

Docking

In order to investigate the mode of interactions between the lipoxygenase and its inhibitors in a 3D fashion, the compounds were docked into the binding site of lipoxygenase. The validity and quality of docking procedure was evaluated by docking of co-crystalized ligand into the binding site of lipoxygenase. The top binding pose from docking studies shows a similar orientation in the binding pocket to the co-crystallized ligand found in crystal structure (PDB ID 11K3). The *root mean square deviation* (RMSD) between ligand docked into the binding site and co crystallized ligand in the crystal structure for lipoxygenase was 1.4 Å indicating good ability to reproduce the ligand binding mode observed in the experimental data (Figure 2). Figure 3 is 3D structure of docked chemicals in protein, in this picture red compound is co-crystalysed ligand

Conclusion

and docked analogs represented as line structures colored by atoms. It indicates synthetic ligands are in suitable position between active site of protein. As mentioned in Table 1 compounds 4d and 4f derivatives demonstrated the most potent inhibitory effects. Investigating ligand interaction mode of docked analogs by LigX module of MOE software revealed that quaternary nitrogen of piperazine ring in these two compounds have made appropriate site for hydrogen binding. As shown in figure 4,5 there is a hydrogen bond between quaternary nitrogen of piperazine ring and ser-510 in active compounds 4d, 4f. Inactive compounds 4a, b, c, g do not have piperazine ring and quaternary nitrogen and so they cannot have hydrogen bonding to ser -510. In inactive compound **4e** which has a piperazine ring, there is a bulk, electron withdrawing N-phenyl substituent, bigger than N-methyl (4d) and N-ethyl (4f), which prevents quaternization of nitrogen of piperazine. So this compound also cannot have hydrogen bond with ser-510 (Figure 6,7). It should be mentioned that presence of hydrophobic amino acids in binding site of lipoxygenase is remarkable (green area in Figure 2 and 3 and lots of hydrophobic amino acids colored as green in Figure 4 and 6 demonstrate this point) and can be considered as opportunity to develop lipoxygenase inhibitors.

In conclusion hydrogen bonds in quaternary nitrogen of piperazine ring of compounds **4d** and **4f** appears to play major role in lipoxygenase inhibition by this set of synthesized analogs and hydrophobic nature of this protein's binding site should be considered in ongoing investigation.

A series of 3,5-bis((5-bromo-6-methyl-2-*t*-aminopyrimidin-4-yl)thio)-4*H*-1,2,4-triazol-4-amines and their cyclized products pyrimidinylthio pyrimidotriazolothiadiazines were synthesized and

evaluated as potential 15-LO inhibitors. Among all these synthesized compounds only two derivatives **4d** and **4f** with IC_{50} 9 and 12 μ M respectively showed strong potency for 15-LO inhibition. The mode of interactions between the lipoxygenase and its inhibitors **4a-g** in a 3D fashion showed that hydrogen bonds in quaternary nitrogen of piperazine ring of compounds **4d** and **4f** appears to play a major role in lipoxygenase inhibition.

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Conflict of Interest

The authors confirm there is no conflict of interest.

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Figures and Table Legends

Table 1: IC₅₀ values for the **4a-g** and **5a-g** compounds.

Scheme 1: Synthesis of tricyclic pyrimidinylthio pyrimido[5,4-e]triazolo[5,1-b][1,3,4]thiadiazines 5a-g.
Figure 1: Log Dose-response 15-LO activity curves for biologically active compounds 4d (panel A), 4f (panel B) and reference 4-MMPB (panel C).

Figure 2: 3D representation of docked ligand (red color) into binding site of lipoxygenase using MOE program as well as co-crystallized one (blue color) in the crystal structure of lipoxygenase (PDB ID: 1IK3).

Figure 3: Map surface of docked analogs in active site of enzyme. Crystallized ligand is indicated by red and stick lines. Green: hydrophobic; violet: H bonding; blue: Mild polar.

Figure 4: The 2D representation of the interaction between compound **4d** (panel A) and compound **4f** (panel B) in the crystal structure of lipoxygenase (PDB ID: 1IK3) using LigX in MOE.

Figure 5: The 3D position of docked compound **4d** (panel A) and compound **4f** (panel B) in the active site of lipoxygenase (PDB ID: 1IK3), hydrogen bonds are in purple and all hydrogens were removed for clarification, compounds represented as stick and amino acid residues as lines colored by atoms.

Figure 6: The 2D representation of the interaction between compound **4b** (panel A), **4a** (panel B) and **4e** (panel C) and lipoxygenase.

Figure 7: The 3D position of docked compound **4b** (panel A), compound **4a** (panel B) and **4e**(panel C) in the active site of lipoxygenase (PDB ID: 11K3), hydrogen bonds are in purple and all hydrogens were removed for clarification, compounds represented as stick and amino acid residues as lines colored by atoms.

Entry	$-NR_2$	$IC_{50}(\mu M)$
4 a		>500
5a		>500
4 b	N—	>500
5b		>500
4 c	N	>500
5c		>500
4d		9
5d		>500
4e		>500
5e		>500
4 f		12
5f		>500
4 g	N—	>500
5g		>500

Table 1: IC₅₀ values for the **4a-g** and **5a-g** compounds









(W#) 172







