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RESEARCH ARTICLE

Structure-based design, synthesis, molecular docking study and biological evaluation of 1,2,4-triazine derivatives acting as COX/15-LOX inhibitors with anti-oxidant activities

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Abstract

A set of 1,2,4-triazine derivatives were designed as cyclooxygenase-2 (COX-2) inhibitors. These compounds were synthesized and screened for inhibition of cyclooxygenases (COX-1 and COX-2) based on a cellular assay using human whole blood (HWB) and lipoxygenase (LOX-15) that are key enzymes in inflammation. The results showed that $3-(2-(benzo[d][1,3]dioxol-5-ylmethylene)hydrazinyl)-5,6-bis(4-methoxyphenyl)-1,2,4-triazine (G11) was identified as the most potent COX-2 inhibitor (78%) relative to COX-1 (50%). Ferric reducing anti-oxidant power (FRAP) assay revealed that compound G10 possesses the highest anti-oxidant activity. The compound G3 with IC₅₀ value of <math>124 \,\mu$ M was the most potent compound in LOX inhibitory assay. Molecular docking was performed and a good agreement was observed between computational and experimental results.

Introduction

Inflammation, a process whereby a huge number of mediators are released due to over-activation of different cascades, is associated with many diseases, such as rheumatoid arthritis (RA), osteoarthritis (OA), bronchospasm and psoriasis¹. Enzymatic action of phospholipases A2 (PLA2s) on phospholipid bilayer results in arachidonic acid release, which is a precursor in the production of eicosanoids including prostaglandins (PGs) and leukotrienes $(LTs)^2$. The transformation of arachidonic acid to PGs is mediated by cyclooxygenase (COX) enzyme family (COX-1, COX-2 and COX-3). The most important isoforms are COX-1 and COX-2, whereas, COX-3 is expressed in the brain and its biological role is still unknown.

COX-1 is the constitutive form of the enzyme and is present in the stomach, intestines, kidneys and platelets which causes gastrointestinal cytoprotection as well as renal effects. On the other hand, COX-2 is constitutively expressed only in low amounts in kidney, brain and ovaries, but its expression is considerably enhanced during inflammatory situations or mitogenic stimulation^{3,4}.

Keywords

Cyclooxygenase, FRAP assay, 15-lipoxygenase, molecular modeling, 1,2,4-triazine

History

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Non-steroidal anti-inflammatory drugs (NSAIDs) are amongst the most widely used drugs worldwide. Traditional NSAIDs (classical NSAIDs or nonselective COX inhibitors), such as ibuprofen, indomethacin and naproxen inhibit both COX-1 and COX-2. Several serious side effects, such as gastrointestinal irritation and hemorrhage and kidney failure are caused by these agents⁵.

The discovery of molecules that selectively inhibit COX-2, named coxibs (celecoxib, rofecoxib and valdecoxib) has been a response to the necessity of lowering these side effects. Indeed, the selectivity of this class of drugs resulted in improved gastrointestinal tolerance. Consecutive clinical studies revealed that long term use of several coxibs is accompanied by increased risk level of cardiovascular problems, such as myocardial infarction, stroke, systemic and pulmonary hypertension, and congestive heart failure⁶. Hence, finding effective and safe anti-inflammatory drugs seems to be an urgent necessity.

In previous investigations, we have reported some new compounds as anti-inflammatory agents. As a part of our ongoing effort to develop effective candidates for inhibition of inflammation, a set of 5,6-diaryl-1,2,4-triazine derivatives were designed⁷.

In contrast with the traditional NSAIDs which belong to different categories of chemical structures, selective COX-2 inhibitors can be classified into three groups (Figure 1): (1) diaryl ethers with methanesulfonamide at position-2 of one ring, Nemisulide and CGP-28237 can be considered as lead compounds in this class⁸; (2) trans-stilbenoids, such as resveratrol, a natural product found in grapes and other food products⁹; (3) heterocyclic

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Figure 1. Different classes of selective COX-2 inhibitors.



systems (such as 1,2,4 triazole^{10,11}, imidazole¹², pyrazol¹³, quinoline¹⁴, indol¹⁵, pyrimidine¹⁶, 1,2,4-triazine^{17,18}) with adjacent diaryl containing sulfamoyl (SO₂NH₂) or methylsulfonyl (SO₂Me) substitution at position-4 of one ring which exist in celecoxib, SC-558 and etoricoxib.

The compounds bearing 1,2,4-triazine core have been identified to exhibit a broad spectrum of biological effects, such as antifungal¹⁹, anti-HIV²⁰, adenosine A_2A receptor antagonist²¹, kinase inhibitor antitumor^{22,23}, analgesic¹⁷, anti-anxiety and anti-inflammation effects²⁴.

As a part of our research program toward the search of potent COX-2 inhibitors with lower side effects, improved affinity and efficacy and dual modes of action, molecular hybridization strategy was employed to design 1,2,4-triazine derivatives. After the synthesis of compounds, *in vitro* assays and molecular docking study were also performed.

Experimental protocols

Chemistry

NMR spectra were recorded on Brucker 500 spectrometer or Varian unity 400 spectrometer relative to tetramethylsilane (TMS) as an internal standard using $CDCl_3$ or dimethyl sulfoxide (DMSO)-d₆ as the solvents. The mass spectra were obtained on an Agilent 6410 instrument. The IR spectra were recorded on a FT-IR Perkin-Elmer spectrophotometer (KBr disks). Melting points were determined on a Kofler hot stage apparatus and are uncorrected. Elemental analyses were performed on a CHN–O–rapid elemental analyzer (Heraeus, Hanau, Germany).

General procedure

All 3-(2-arylidenehydrazinyl) 5,6-bis(4-methoxyphenyl)-1,2,4-triazine derivatives were prepared according to the procedure described⁷.

3-(2-Benzylidenehydrazinyl)-5,6-bis(4-methoxyphenyl)1,2,4-triazine (G1)⁷

Compound G1 was prepared by the described procedure using 106 mg (1 mmol) of benzaldehyde (yield 83%), m.p.: 245–247 °C; ¹H NMR (DMSO-d₆, 500 MHz), $\delta_{\rm H}$: 10.29 (s, 1H, NH), 8.24 (s, 1H, N=CH), 7.81 (d, J = 7.1 Hz, 2H, ArH), 7.66 (d, J = 8.8 Hz, 2H, ArH), 7.50 (d, J = 8.7 Hz, 2H, ArH), 7.43–7.38 (m, 3H, ArH), 6.93 (d, J = 8.7 Hz, 2H, ArH), 6.88 (d, J = 8.7 Hz, 2H, ArH), 3.86 ppm (s, 6H, $2 \times O$ –CH₃H); IR (KBr, ν): 3440, 3211, 1605, 1571, 831 cm⁻¹; MS (EI) m/z (%): 411.2 (M⁺, 29), 334 (20), 308 (14), 238 (100), 223 (49), 195 (12), 152 (11). Anal. Calcd. for C₂₄H₂₁N₅O: C, 70.06; H, 5.14; N, 17.02; found: C, 70.21; H, 4.86; N, 16.95.

3-(2-(4-Nitrobenzylidene)hydrazinyl) 5,6-bis(4-methoxyphenyl)-1,2,4-triazine (**G2**)⁷

The titled compound was synthesized by the described procedure from 4-nitrobenzaldehyde (151 mg, 1 mmol) as yellow solid (yield 88%), m.p.: 265–267 °C; ¹H NMR (DMSO-d₆, 500 MHz), $\delta_{\rm H}$: 12.23 (s, 1H, NH), 8.36 (s, 1H, N=CH), 8.31 (d, J = 8.6 Hz, 2H, ArH), 7.99 (d, J = 8.0 Hz, 2H, ArH), 7.52 (d, J = 8.3 Hz, 2H, ArH), 7.39 (d, J = 8.6 Hz, 2H, ArH), 6.98–6.96 (m, 4H, ArH), 3.79 ppm (s, 6H, 2 × O–CH₃); IR (KBr, ν): 3421, 3205, 1607, 1585, 838 cm⁻¹; MS (EI) m/z (%): 456 (M⁺, 21), 308 (16), 238 (100), 223 (25), 195(11). Anal. Calcd. for C₂₄H₂₀N₆: C, 63.15; H, 4.42; N, 18.41; found: C, 62.78; H, 4.12; N, 18.51.

3-(2-(4-Fluorobenzylidene)hydrazinyl)-5,6-bis(4methoxyphenyl)-1,2,4-triazine (**G3**)⁷

The titled compound G3 was synthesized by the general procedure using 4-fluorobenzaldehyde (124 mg, 1 mmol) (yield 72%), m.p.: 257–259 °C; ¹H NMR (DMSO-d₆, 500 MHz), $\delta_{\rm H}$: 11.64 (s, 1H, NH), 8.25 (s, 1H, N=CH), 8.06–8.02 (m, 2H, ArH), 7.49 (d, J = 8.5 Hz, 2H, ArH), 7.35 (d, J = 8.5 Hz, 2H, ArH),

7.30–7.26 (m, 2H, ArH), 6.96–6.93 (m, 4H, ArH), 3.78 ppm (s, 6H, 2 × O–CH₃); ¹³C NMR (DMSO-d₆, 125 MHz), $\delta_{\rm C}$ = 163.39, 161.45, 160.99, 159.39, 158.21, 155.45, 150.64, 141.87, 137.48, 131.19, 130.86, 130.80, 130.21, 128.63, 127.87, 122.86, 116.03, 115.86, 113.83, 112.42, 112.24, 55.27, 55.11 ppm; IR (KBr, ν): 3444, 3223, 2913, 1607, 1589, 1514, 1256, 836 cm⁻¹; MS (EI) *m*/*z* (%): 429 (M, 36), 334 (29), 308 (10), 238 (100), 223 (53), 195 (12), 152 (11). Anal. Calcd. for C₂₄H₂₀FN₅O₂: C, 67.12; H, 4.69; N, 16.31; found: C, 66.85; H, 4.81; N, 15.98.

5,6-Bis(4-methoxyphenyl)-3-(2-(thiophen-2-ylmethylene)hydrazi-nyl)-1,2,4-triazine $(\mathbf{G4})^7$

The yellow solid product G4 was prepared from thiophene-2carbaldehyde (112 mg, 1 mmol) in similar manner described in general procedure (yield: 91%), m.p.: 250–253 °C; ¹H NMR (DMSO-d₆, 500 MHz), $\delta_{\rm H}$: 11.79 (s, 1H, NH), 8.45 (s, 1H, N=CH), 7.61 (s, 1H, ArH), 7.44–7.46 (m, 2H, ArH), 7.39–7.35 (m, 3H, ArH), 7.12 (brs, 1H, ArH), 6.96–6.95 (m, 4H, ArH), 3.78 ppm (s, 6H, 2 × O–CH₃); ¹³C NMR (DMSO-d₆, 125 MHz), $\delta_{\rm C}$ = 160.94, 159.34, 158.03, 155.37, 151.23, 150.33, 139.65, 138.66, 131.13, 130.17, 129.28, 128.67, 127.90, 127.76, 55.27, 55.10 ppm. IR (KBr, ν): 3350, 3220, 1608, 1514, 1360, 831 cm⁻¹; MS (EI) *m*/*z* (%): 417(M⁺, 61), 308 (10), 283 (10), 252 (11), 238 (100), 223 (57), 195(20), 180 (13), 152 (23), 39 (16). Anal. Calcd. for C₂₂H₁₉N₅O₄S: C, 66.23; H, 4.59; N, 16.78; found: C, 65.94; H, 4.74; N, 16.94.

3-(2-(4-Hydroxy-3-methoxybenzylidene)hydrazinyl)-5,6-bis(4mthoxyphenyl)-1,2,4-triazine (G5)⁷

Compound G5 was obtained according to the described general procedure using 4-hydroxy-3-methoxybenzaldehyde (152 mg, 1 mmol) as yellow solid (yield 88%): m.p.: 245-248 °C; ¹H NMR (DMSO-d₆, 500 MHz), $\delta_{\rm H}$: 11.62 (s, 1H, NH), 9.46 (s, 1H, N=CH), 8.15 (s, 1H, ArOH), 7.49 (d, J = 8.5 Hz, 2H, ArH), 7.36 (d, J = 8.5 Hz, 2H, ArH), 7.31 (s, 1H, ArH), 7.11 (d, J = 8.1 Hz, 7.11 Hz)1H, ArH), 6.96–6.94 (m, 4H, ArH), 6.84 (d, *J* = 8.1 Hz, 1H, ArH), 3.84 (s, 3H, OCH₃), 3.78 ppm (s, 6H, $2 \times O$ -CH3); ¹³C NMR $(DMSO-d_6, 125 \text{ MHz}), \delta_C = 161.48, 159.29, 158.22, 155.28,$ 149.99, 148.27, 147.91, 144.08, 131.12, 130.93, 130.72, 130.14, 129.16, 128.78, 128.01, 126.29, 120.87, 115.51, 113.79, 113.70, 113.57, 109.18, 55.50, 55.25, 55.09 ppm; IR (KBr, ν): 3485, 3225, 1607, 1508, 1257, 843 cm⁻¹; MS (EI) *m/z* (%): 417 (M⁺, 61), 323 (10), 308 (37), 252 (14), 238 (100), 223 (53), 195 (14), 152 (12). Anal. Calcd. for C₂₅H₂₃N₅O₄: C, 65.63; H, 5.07; N, 15.31; found: C, 65.93; H, 5.36; N, 15.51.

3-(2-(2,4-Dichlorobenzylidene)hydrazinyl)-5,6-bis(4-methoxyphe-nyl)-1,2,4-triazine ($\mathbf{G6}$)⁷

The titled compound G6 was synthesized according to the general procedure using 2,4-dichlorobenzaldehyde (175 mg, 1 mmol) as pale yellow solid (yield 78%), m.p.: 248–251 °C; ¹H NMR (DMSO-d₆, 500 MHz), $\delta_{\rm H}$: 12.14 (s, 1H, NH), 8.61 (s, 1H, N=CH), 8.06 (d, J=8.6 Hz, 1H, ArH), 7.72 (d, J=2.0 Hz, 1H, ArH), 7.53 (dd, J=2.0 Hz, J=8.5 Hz, 1H, ArH), 7.50 (d, J=8.7 Hz, 2H, ArH), 7.38 (d, J=8.7 Hz, 2H, ArH), 6.97–6.95 (m, 4H, ArH), 3.79 ppm (s, 6H, 2 × O–CH₃); IR (KBr, ν): 3444, 3183, 1607, 1511, 1369.82 (M⁺², 5), 334 (16), 308 (15), 238 (100), 223 (45), 195 (13), 152 (10). Anal. Calcd. for C₂₄H₁₉Cl₂N₅O₂: C, 60.01; H, 3.99; N, 14.58; found: C, 59.84; H, 4.16; N, 14.41.

5,6-Bis(4-methoxyphenyl)-3-(2-((5-nitrofuran-2-yl)methylene) hydrazinyl)-1,2,4-triazine (G7)

The titled compound G7 was synthesized according to the general procedure using 5-nitrofuran-2-carbaldehyde (141 mg, 1 mmol) as

a pale yellow solid (yield 78%), m.p.: $248-251 \,^{\circ}\text{C}^{-1}\text{H}$ NMR (DMSO-d₆, 500 MHz), δ_{H} : 12.34 (s, 1H, NH), 8.74 (s, 1H, N=CH), 7.82 (d, J = 3.9 Hz, 1H, ArH), 7.49–7.47 (m, 2H, ArH), 7.39–7.37 (m, 2H, ArH), 7.21 (d, J = 3.9 Hz, 1H, ArH), 6.97–6.95 (m, 4H, ArH), 3.79 ppm (s, 6H, $2 \times \text{O-CH}_3$) IR (KBr, ν): 3443, 2839, 1608, 1514, 1353, 1254, 1071, 834 cm⁻¹ MS (EI) m/z (%): 446 (M⁺, 17), 400 (100), 267 (24), 238 (79), 223 (86), 195 (23), 180 (22), 152 (21), 134 (17), 119 (16), 79 (12). Anal. Calcd. for C₂₂H₁₈N₆O₅: C, 59.19; H, 4.06; N, 18.83; found: C, 59.64; H, 3.94; N, 18.11.

5,6-Bis(4-methoxyphenyl)-3-(2-[(1H-indol-3-yl)methylene]hydrazinyl)-1,2,4-triazine (**G8**)⁷

The titled compound G8 was synthesized according to the general described procedure using 1H-indole-3-carbaldehyde (145 mg, 1 mmol) as yellow solid (yield: 74%), m.p.: 258–262 °C; ¹H NMR (DMSO-d₆, 500 MHz), $\delta_{\rm H}$: 11.48 (s, 1H, NH), 8.51 (d, J = 7.2 Hz, 1H, ArH), 8.45 (s, 1H, N=CH), 7.77 (brs, 1H, NH), 7.56 (d, J = 7.7 Hz, 2H, ArH), 7.44 (d, J = 7.7 Hz, 2H, ArH), 7.39 (d, J = 8.4 Hz, 2H, ArH), 7.21–7.18 (m, 2H, ArH), 6.99–6.96 (m, 4H, ArH), 3.80 ppm (s, 6H, $2 \times O-CH_3$); ¹³C NMR (DMSO-d₆, 125 MHz), $\delta_{\rm C} = 160.95$, 159.24, 158.44, 155.01, 149.39, 140.71, 136.98, 131.82, 131.12, 130.13, 129.01, 128.92, 128.26, 124.44, 122.50, 122.32, 120.23, 113.83, 113.74, 112.35, 111.68, 55.32, 55.12 ppm; IR (KBr, ν): 3414, 3010, 1609, 1510, 1249, 1065, 839 cm^{-1} ; MS (EI) m/z (%): 450 (M⁺, 4), 308 (14), 238 (100), 223(47), 195 (15), 142 (18), 116 (16). Anal. Calcd. for C₂₆H₂₂N₆O₂: C, 69.32; H, 4.92; N, 18.66; found: C, 68.97; H, 4.76; N, 18.34.

3-(2-(2-Chloro-6-nitrobenzylidene)hydrazinyl)-5,6-bis(4-methox-yphenyl)-1,2,4-triazine $(\mathbf{G9})^7$

The yellow solid product G9 was prepared from 2-chloro-6nitrobenzaldehyde (185 mg, 1 mmol) according to the general procedure (yield: 71%), m.p.: 258–262 °C; ¹H NMR (DMSO-d₆, 500 MHz), $\delta_{\rm H}$: 12.33 (s, 1H, NH), 8.68 (s, 1H, N=CH), 8.14– 8.13(m, 2H, ArH), 7.70 (dd, J = 2.45 Hz, J = 8.8 Hz, 1H, ArH), 7.51 (d, J = 8.8 Hz, 2H, ArH), 7.39 (d, J = 8.7 Hz, 2H, ArH), 6.98–6.96 (m, 4H, ArH), 3.79 ppm (s, 6H, 2 × O–CH3); IR (KBr, ν): 3424, 3121, 1602, 1519, 1125, 831 cm⁻¹; MS (EI) *m/z* (%): 490 (M⁺, 21), 492 (M⁺², 6), 334 (16), 308 (15), 238 (100), 223 (45), 195 (13), 152(10). Anal. Calcd. for C₂₄H₁₉CIN₆O₄: C, 58.72; H, 3.90; N, 17.12; found: C, 58.97; H, 3.72; N, 17.16.

3-(2-(1,3-Dihydroxybenzylidene)hydrazinyl)-5,6-bis(4-methoxyphenyl)-1,2,4-triazine (G10)⁷

The titled compound G10 was synthesized according to the general procedure from 2,4-dihydroxybenzaldehyde (138 mg, 1 mmol) as yellow solid (yield: 86%), m.p.: 262–266 °C; ¹H NMR (DMSO-d₆, 500 MHz), $\delta_{\rm H}$: 11.93 (s, 1H, NH), 11.71 (s, 1H, ArOH), 9.89 (s, 1H, ArOH) 8.31 (s, 1H, N=CH), 7.77 (s, 1H, ArH), 7.50 (d, J = 8.7 Hz, 2H, ArH), 7.38 (d, J = 8.7 Hz, 2H, ArH), 7.25 (d, J = 8.4 Hz, 1H, ArH), 6.97–6.95 (m, 4H, ArH), 6.37–6.35 (dd, J = 2.2 Hz, J = 8.3 Hz, 1H, ArH), 3.79 ppm (s, 6H, 2 × O–CH3); IR (KBr, ν): 3352, 2837, 1608, 1517, 1253, 834 cm⁻¹; MS (m/z, %): 443 (M⁺, 22), 308 (11), 238 (100), 195 (11). Anal. Calcd. for C₂₄H₂₁N₅O₄: C, 65.01; H, 4.77; N, 15.79; found: C, 64.91; H, 4.83; N, 16.01.

3-(2-(Benzo[d][1,3]dioxol-5-ylmethylene)hydrazinyl)-5,6-bis(4methoxyphenyl)-1,2,4-triazine (G11)⁷

The titled compound G11 was synthesized by the described general procedure using 1,3-benzodioxole-5-carbaldehyde (piper-onal) (150 mg, 1 mmol) (yield 88%), m.p.: 268–272 °C; ¹H NMR

(DMSO-d₆, 400 MHz), $\delta_{\rm H}$: 11.62 (s, 1H, NH), 8.16 (s, 1H, N=CH), 7.49 (d, J = 7.6 Hz, 2H, ArH), 7.40–7.34 (m, 2H, ArH), 7.31 (s, 1H, ArH), 14 (d, J = 7.6 Hz, 2H, ArH), 6.98–6.94 (m, 4H, ArH), 6.08 (s, 2H, ArH), 3.78 ppm (s, 6H, 2 × O–CH₃); IR (KBr, ν): 3429, 2962, 1606, 1515, 1175, 836 cm⁻¹; MS (EI) *m*/*z* (%): 455 (M⁺, 18), 334 (10), 308 (16), 238 (100), 223 (38), 195 (11). Anal. Calcd. for C₂₅H₂₁N₅O₄: C, 65.93; H, 4.65; N, 15.38; found: C, 65.71; H, 5.01; N, 15.65.

3-(2-(Furan-2-ylmethylene)hydrazinyl)-5,6-bis(4-methoxyphenyl)-1,2,4-triazine (G12)⁷

The yellow solid product G12 was prepared from furan-2carbaldehyde (96 mg, 1 mmol) according to the general procedure (yield: 71%), m.p.: 258–262 °C; ¹H NMR (DMSO-d₆, 500 MHz), $\delta_{\rm H}$: 11.79 (s, 1H, NH), 8.15 (s, 1H, N=CH), 7.82 (s, 1H, ArH), 7.47 (d, J = 8.6 Hz, 2H, ArH), 7.36 (d, J = 8.6 Hz, 2H, ArH), 6.96–6.94 (m, 4H, ArH), 6.85 (d, J = 3.2, 1H, ArH), 6.63 (brs, 1H, ArH), 3.78 ppm (s, 6H, 2 × O–CH3); ¹³C NMR (DMSO-d₆, 125 MHz), $\delta_{\rm C}$ = 161.57, 161.42, 159.99, 158.72, 156.17, 151.82, 150.92, 150.33, 145.73, 144.97, 134.10, 131.66, 130.71, 129.16, 128.44, 116.15, 114.30, 114.22, 112.57, 55.77, 55.61 ppm; IR (KBr, ν): 3443, 2839, 1608, 1254, 834 cm⁻¹; MS (EI) *m/z* (%): 401 (M⁺, 66), 372 (23), 308 (15), 267 (47), 252 (14), 238 (100), 223 (89), 195 (22), 152 (21), 135 (37). Anal. Calcd. for C₂₂H₁₉N₅O₃: C, 65.83; H, 4.77; N, 17.45; found: C, 65.51; H, 5.03; N, 17.12.

$3-(2-(4-Methoxybenzylidene)hydrazinyl)-5,6-bis(4-methoxyphe-nyl)-1,2,4-triazine (G13)^7$

The titled compound G13 was synthesized according to the general procedure from 4-methoxybenzaldehyde (136 mg, 1 mmol) as yellow solid (yield: 91%), m.p.: 254–256 °C; ¹H NMR (DMSO-d₆, 500 MHz), $\delta_{\rm H}$: 11.72 (s, 1H, NH), 8.21 (s, 1H, N=CH), 7.67 (d, J = 8.0 Hz, 2H, ArH), 7.49 (d, J = 8.0 Hz, 2H, ArH), 7.36 (d, J = 8.0 Hz, 2H, ArH), 7.02–6.94 (m, 6H, ArH), 3.80 (s, 3H, OCH3), 3.78 ppm (s, 6H, 2 × O–CH3); IR (KBr, ν): 3444, 3215, 1608, 1577, 1248, 834 cm⁻¹; MS (EI) *m*/*z* (%): 441 (M⁺, 14), 308 (31), 238 (100), 223 (49), 195 (12), 152 (10). Anal. Calcd. for C₂₅H₂₃N₅O₃: C, 68.01; H, 5.25; N, 15.86; found: C, 68.28; H, 4.93; N, 15.67.

5,6-Bis(4-methoxyphenyl)-3-(2-(pyridin-4-ylmethylene)-hydrazinyl)-1,2,4-triazine (G14)⁷

The yellow solid product G14 was prepared from pyridine-4carbaldehyde (107 mg, 1 mmol) according to the general procedure (yield: 81%), m.p.: 255–258 °C; ¹H NMR (DMSO-d₆, 500 MHz), $\delta_{\rm H}$: 12.17 (s, 1H, NH), 8.63 (d, J = 4.4 Hz, 2H, ArH), 8.23 (s, 1H, N=CH), 7.67 (d, J = 5.7 Hz, 2H, ArH), 7.52 (d, J = 8.8 Hz, 2H, ArH), 7.39 (d, J = 8.6 Hz, 2H, ArH), 6.98–6.96 (m, 4H, ArH), 3.79 ppm (s, 6H, 2 × O–CH3); IR (KBr, ν): 3434, 3146, 1609, 1569, 1507, 1237, 841 cm⁻¹; MS (EI) *m/z* (%): 412 (M⁺, 32), 334 (16), 308 (12), 238 (100), 223 (25), 195 (10). Anal. Calcd. for C₂₃H₂₀N₆O₂: C, 66.98; H, 4.89; N, 20.38; found: C, 66.75; H, 5.71;N, 19.93.

3-(2-(4-Bromobenzylidene)hydrazinyl)-5,6-bis(4-methoxyphenyl)-1,2,4-triazine (G15)

The yellow solid product G15 was prepared from 4-bromobenzaldehyde (185 mg, 1 mmol) according to the general procedure (yield: 82%), m.p.: 254–256 °C. ¹H NMR (DMSO-d₆, 500 MHz), $\delta_{\rm H}$: 11.92 (s, 1H, NH), 8.24 (s, 1H, N=CH), 7.69–7.64 (m, 4H, ArH), 7.50 (d, J = 8.2, 2H, ArH), 7.37 (d, J = 8.2 Hz, 2H, ArH), 6.97–6.95 (m, 4H, ArH), 3.79 ppm (s, 6H, 2 × O–CH3). IR (KBr, ν): 3422, 3223, 2907, 1607, 1592, 1510, 1254, 837 cm⁻¹; MS (m/z, %): 489.1 (M⁺, 11.5), 491.1 (M⁺², 11.4). Anal. Calcd. for C₂₄H₂₀BrN₅O₂: C, 58.79; H, 4.11; N, 14.28; found: C, 59.35; H, 4.91; N, 13.24.

Biological activities

In vitro COX inhibition assays

In vitro bioactivities of all the synthesized compounds were evaluated using celecoxib and indomethacin as the reference drugs. The HWB assay was employed to determine the activity of COX isoenzymes³⁶. Five-hundred microliters of fresh blood samples obtained from human volunteers were incubated with 1 µl of DMSO as a vehicle or 1 µl of 10 µM solution of synthesized compounds or indomethacin and celecoxib as the reference drugs. The plasma was separated after 1 h incubation at 37 °C. COX-1 level was measured by the production of thromboxane B2 (TxB2) using a commercially available enzyme immunoassay (EIA) kit.

In order to determine COX-2 activity, 500 μ l of fresh blood samples obtained from human volunteers were collected in EDTA-containing tubes, incubated with 1 μ l of DMSO as a vehicle or 1 μ l of 10 μ M solution of synthesized compounds or indomethacin and celecoxib as the reference drugs. The samples were incubated at 37 °C in the presence of 10 μ g/ml LPS for 24 h, after which the plasma was separated. COX-2 level was measured by the production of PGE2 using a commercially available EIA kit.

Inhibition of 15-LOX

The assay was performed according to the previously reported method³⁷. The compounds were dissolved in DMSO and phosphate buffer. The solution was added to the assay mixture that contained 9 ml of 0.1 M sodium phosphate buffer, pH 8 and enzyme (final concentration: 167 U/ml). The mixture was incubated for four minutes and then linoleic acid as substrate was added to reach the final concentration of 134 μ M. Quercetin, a well-known inhibitor of 15-LOX was used as a positive control. Absorbance was recorded by spectrophotometer at 234 nm during the measurement period. Enzyme activity experiments were tested in triplicate and the values are expressed as means ± SD.

Determination of the anti-oxidant activity by FRAP assay

FRAP assay³³ was employed to evaluate the anti-oxidant activities of the target compounds. For this purpose, FRAP solution was prepared by adding 10 ml acetate buffer 300 mM at pH 3.6 to 1 ml of aqueous solution of FeCl₃·6H₂O and 1 ml (10 mM) acidic solution of 2,4,6-tris(2-pyridyl)-s-triazine (TPTZ). Twenty five microliters of 100 μ M of compounds was added to FRAP solution. After incubation at 37 °C for 15 min, the absorbance at 585 nm was monitored relative to ferrous sulfate (FeSO₄) as the standard. Results are reported as mmol Fe(II) per gram of dry mass of compounds. Ascorbic acid was selected as a reference. The data are mean of three independent experiments.

Molecular docking study

Molecular docking study was performed using Autodock 4.2 and Auto Dock Tools 1.5.4 (ADT). For this purpose, the X-ray crystal structure of COX-1 carrying diclofenac as cognate ligand in binding site (PDB code: 3N8Y) and X-ray crystal structure of COX-2 in complex with SC-558 (PDB code: 1CX2) were downloaded from protein data bank (http://www.rcsb.org). Water molecules and co-crystalized ligand were excluded. The PDB file of protein was regenerated by adding polar hydrogens and computing Gasteiger charges.



Figure 2. Design of COX-2 inhibitors using molecular hybridization method.



Scheme 1. Synthesis pathway for the preparation of triazine derivatives. Reagents and conditions: (a) thiamine hydrochloride, NaOH, EtOH, reflux, 3 h; (b) HNO₃, 60 °C; (c) thiosemicarbazide, CH₃COOH, reflux, 3 h; (d) CH₃I, NaOH, EtOH, 2 h, rt; (e) NH₂NH₂·H₂O, EtOH, reflux, 12 h; (f) ArCHO, EtOH, 1–10 h, rt.

The 3D structures of ligands were created using Avogadro software. Each ligand structure was separately submitted to a molecular mechanics conformational search (MM⁺ force field) and the geometry of the most stable conformer was then optimized by semi-empirical method (AM1) in Hyperchem software. The Gasteiger charges were added and the degrees of torsions were defined to generate PDBQT format of the ligands.

The grid parameter file was generated using AutoGrid. Numbers of points in x, y, z dimensions were set to $60 \times 60 \times 60$ with 0.375 Å grid spacing. Rigid macromolecule was chosen to define the docking parameter file using Lamarckian Genetic Algorithm (LGA), with 200 GA runs and 25 000 000 energy evaluations while the other parameters were left as default. The results revealed that all tested compounds fit much well in the COX-2 cavity rather than COX-1 cavity.

Results and discussion

Fragment-based design

The target 1,2,4-triazine derivatives were designed by the molecular hybridization approach based on selective COX-2 inhibitor etoricoxib (1) and a number of potent anti-inflammatory agents that are reported in the literature (compounds 2–7) (Figure 2). Etoricoxib (1), commonly used for pain relief after surgery, is approved in more than 80 countries worldwide which is a selective COX-2 inhibitor, containing pyridine with bearing adjacent diaryls.

It has been reported that 1,2,4-triazine heterocyclic compounds with diaryl substituent (2) are effective COX-2 inhibitors^{17,18}. Some experiments have reported structures in which the sulfonyl moiety on the aromatic ring is missing $(3)^{25-27}$. Niedballa et al. showed that sulfonyl moiety can be replaced with alkoxy groups²⁸.

Table 1. COX-1 and COX-2 inhibitory activities of synthesized 1,2,4-triazine compounds.



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Compound	Ar	COX-2 (10 µM)	COX-1 (10 µM)	COX-2/COX-1
G1	\bigcap	71.85 ± 8.039	67.3 ± 6.057	1.07
G2	NO ₂	72.32 ± 9.023	76.86±13.098	0.94
G3	↓ ↓ F	72.09 ± 11.027	69.1 ± 6.088	1.04
G4	s	87.45 ± 11.008	72.03 ± 7.109	1.21
G5	OMe	63.58 ± 8.004	59 ± 6.098	1.08
G6	CI CI	61.2 ± 7.006	57.2 ± 3.073	1.07
G7		65.85 ± 4.010	59.8 ± 5.099	1.10
G8		60.5 ± 9.006	38.76 ± 6.054	1.56
G9		65.7 ± 7.009	50.76 ± 10.108	1.29
G10	но он	68.9 ± 5.007	63.73 ± 2.119	1.08
G11		78.00±.8.012	50.33 ± 3.085	1.55
G12		78.71 ± 9.023	66.43 ± 4.204	1.18
G13		71.85 ± 10.012	53.66 ± 6.102	1.34
G14	N	73.74 ± 9.011	64.26 ± 7.123	1.15
G15	Br	73.27 ± 12.006	59.36 ± 6.097	1.23
Indomethacin at 500 nM Celecoxib at 500 nM		84.32 ± 0.018 70.91 ± 0.031	$73.96 \pm 0.032 \\ 32.4 \pm 0.024$	1.14 2.19

The results are summarized for inhibition% of COX isoenzymes. The data are expressed as the mean \pm SEM from three different experiments (n = 3).

The natural product (4) resveratrol, has been introduced as a COX-2 inhibitor²⁹. Lin et al. demonstrated that bioisosteric replacement of the central -N=N- linkage of (5) with -N=C-double bond leads to 114-fold more COX-2 inhibitory effect than COX-1³⁰.

2-Benzylidene-1-phenylhydrazine derivative (6) has been proposed as a COX-2 inhibitor and hydrazine derivative (7) is described as a dual COX/5-LOX inhibitor³¹. With respect to these structural features, we designed a set of COX-2 inhibitors. The target compounds were synthesized via the same route that we had previously reported⁷.

Chemistry

The synthetic reactions used for the synthesis of the designed fifteen 1,2,4-triazine derivatives (G1–G15) are outlined in Scheme 1. 2-Hydroxy-1,2-bis(4-methoxyphenyl) ethanone (B) was prepared by thiamine hydrochloride-mediated coupling of 4-methoxy benzaldehyde (A). Oxidation of compound B afforded the corresponding 1,2-bis(4-methoxyphenyl)ethane-1,2-dione (C). 5,6-Diaryl-1,2,4-triazine-3-thiol (D) was acquired through cyclization of compound (C) with thiosemicarbazide. Compound (D) was treated with methyl iodide to afford corresponding methylthio compound (E). In the presence of excess amount of hydrazine hydrate, compound (F) was prepared. The final compounds (G1–G15) were prepared by treatment of (F) with equimolar amounts of the appropriate aldehydes. All the steps were carried out according to our previously reported procedures⁷.

Biological activity

In vitro COX inhibition assay

The human whole blood (HWB) assay, initially developed by Patrignani et al.³², was performed to assess the ability of synthesized compounds to inhibit COX-2 and COX-1 isoenzymes. Celecoxib and indomethacin were used as the reference drugs. The results are summarized in Table 1. Celecoxib as a selective COX-2 inhibitor exhibited 70.9%, 32.4% inhibition for COX-2 and COX-1, respectively, while in the case of indomethacin which is a nonselective COX inhibitor exhibited 84.3% inhibition for COX-2 and 73.9% inhibition for COX-1. These 15 compounds inhibited COX isoenzymes to various degrees, in a range of 60-78% at 10 µM. Compounds G9, G4, G13, G8 and G11 represented the highest selectivity for COX-2 in comparison with COX-1. The compound G11, namely 1,3-benzodioxole-substituted compound, showed the highest selectivity for COX-2 at 10 µM. A brief structure-activity relationship (SAR) can be distinguished as follows according to the data presented in Table 1:

- The introduction of electron-withdrawing moieties, such as nitro or fluoro on phenyl ring resulted in compounds with the lowest activities against COX-2 in this set. This effect can be seen in compounds G2 and G3.
- Para-substituted phenyl rings by both electron-donating and electron-withdrawing groups represented reduced activities.
 G13 and G15 showed a lower affinity toward COX-1 and hence represented higher COX-2/COX-1 ratios.
- Compounds with five membered heterocyclic rings, such as G4 and G12 displayed COX-2 selectivity. G11 and G8 with heterocyclic fused ring aldehydes were the most selective ones among the tested compounds.

In vitro 15-lipoxygenase (LOX) inhibitory activity

In vitro 15-LOX inhibitory activities of synthesized compounds were tested. The results are reported as IC_{50} values in Table 2.

Table 2. The 15-lipoxygenase inhibitory and anti-oxidant activities of the synthesized 1,2,4-triazine compounds.

Compound	$SLO^a \ IC_{50} \ (\mu M)^b$	FRAP value
G1	143	0.309
G2	225	0.29
G3	124	0.309
G4	420	60.7
G5	NA ^c	269.9
G6	NA	0.264
G7	NA	0.1639
G8	NA	147.38
G9	370	0.198
G10	205	487
G11	NA	0.2781
G12	NA	0.3704
G13	NA	0.327
G14	190	42.8
G15	NA	0.293
Quercetin	16	1903.6
Vitamin C	-	674.78

^aSoybean lipoxygenase.

^bData are mean of three independent experiments.

^cNo activity.

The compound G3 containing flourophenyl moiety with IC_{50} value of $124 \,\mu$ M was the most potent one. As reported in Table 2, the phenyl substituted compound G1 showed IC_{50} value of $143 \,\mu$ M and insertion of different substituents resulted in decreased activity. Compounds G1, G14 and G10 (phenyl, pyridyl and di-hydroxy derivatives, respectively) were active against 15-LOX to some extent and compounds G2, G9, G4 (nitrophenyl, 2-chloro-6-nitrophenyl and thiophenyl, respectively) showed less activity while the other compounds were inactive.

Ferric reducing anti-oxidant power (FRAP) assay

The anti-oxidant potential of target compounds were determined using FRAP $assay^{33}$. This method is based on the ability of compounds in the reduction of ferric tripyridyl triazine complex to ferrous (Fe²⁺) form. The result is production of an intense blue color. The change in the absorbance at 585 nm is a tool for monitoring the reaction progress. Higher increase in the absorbance indicates that the potential of reduction is higher. The results are presented in Table 2. Ascorbic acid (vitamin C) was considered as a reference compound. Compound G10 showed the highest anti-oxidant activity.

Molecular modeling

In order to clarify the mechanism of COX inhibition and virtual interactions between synthetic compounds and active sites of COX-1 (PDB code: 3N8Y)³⁴ and COX-2 (PDB code: 1CX2)³⁵, the molecular docking was performed. Substitution of Ile523 in COX-1 relative to Val523 in COX-2 which is a less bulky amino acid and hence the conformational change at Tyr355 results in the presence of an accessible 2°-pocket in COX-2 active site. While the similar pocket in COX-1 is inaccessible. The other reason for effective COX-2 inhibition is the replacement of His513 in COX-1 by Arg513 in COX-2 which enables the ligands to form more H-bonds in the COX-2 active site. Therefore, because of the differences between the active sites of enzymes, it enables us to design selective COX-2 inhibitors.

The docking study was performed using Autodock4.2. All the ligands showed a similar orientation in COX-2 active site (Figure 3). Compound G11 docked in COX-2 is shown (Figure 4). In general, it is observed that the oxygen atom of methoxy phenyl moiety is placed close to the side chain of Arg-120 (guanidine

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Figure 3. Superimposition of ligands in the COX-2 binding pocket.



Figure 4. 3D orientation model of G11 (ball and stick) in COX-2 active pocket (blue: hydrophobic pocket, red: Tyr353, yellow: responsible for the COX-2 selectivity, purple: selectivity pocket).



H₂N...OCH₃, 2.5 Å) while the other methoxy phenyl is accommodated in hydrophobic pocket formed by Tyr385, Gly526, Ser530, Ala527. 1,2,4-Triazine and hydrazine moieties were inserted in the selectivity pocket that was created as a result of Val523 presence in COX-2 by Ile523 in COX-1, then they got involved in H-bond with Leu352 (N=N...C=O, 2.2 Å), (triazine N...C=O, 2.9 Å), Ser353 (triazine N...C=O, 3.1 Å), Tyr355 (triazine N...O-H, 2.8 Å). The other hydrogen bond was formed by benzo[1,3]dioxole fragment (O...N, 3.2 Å).

A similar method was used to compare ligand interactions in COX-1 (PDB code: 3N8Y) enzyme active site. All compounds showed a different conformation in COX-1 active site from their conformation in COX-2 active site.

The compound G11 as a representative example is shown in Figure 5. As it was mentioned before, the presence of an inaccessible pocket in COX-1 resulted in different conformation and hence different H-bonds with Ser530 (triazine N...C=O, 2.8 Å, 2.9 Å), Tyr385 (triazine N...OH, 2.8 Å) and Tyr355 (CH₃-O...OH, 3.5 Å).

Conclusion

In summary, fragment based design strategy was utilized to design 1,2,4-triazine derivatives as selective COX-2 inhibitors. The designed compounds were synthesized and their biological

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Figure 5. 3D model of the orientation of G11 (ball and stick) in COX-1 active site (blue: hydrophobic pocket, red: Tyr353, yellow: responsible for the COX-2 selectivity, purple: selectivity pocket).



activities were evaluated. The results demonstrated that compounds G9, G4, G13, G8 and G11 exhibited stronger effect against COX-2 in comparison with their activity against COX-1 at 10 μ M. *In vitro* 15-LOX inhibitory activity of synthesized compounds was also tested. Compound G3 with an IC₅₀ value of 124 μ M was the most potent derivative. The anti-oxidant potential of synthesized compounds was determined by FRAP assay and compound G10 showed the highest anti-oxidant activity. The binding patterns of compounds inside the COX-2/ COX-1 active sites were predicted using a docking technique. Considering the docking results, it can be suggested that 1,2,4triazine and hydrazine moieties are positioned in the selectivity pocket in COX-2 active site, which provides a useful scaffold for novel compounds with improved inhibitory activities.

Declaration of interest

The authors report no conflicts of interest. The authors alone are responsible for the content and writing of this article. The authors acknowledge the financial support of Shiraz University of Medical Sciences, Vice Chancellor of Research (Grant: 7442).

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