

Design and synthesis of new 1,2-diaryl-4,5,6,7-tetrahydro-1H-benzo[d]imidazoles as selective cyclooxygenase (COX-2) inhibitors

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Abstract A new series of 1,2-diaryl-4,5,6,7-tetrahydro-1H-benzo[d]imidazoles, possessing a methylsulfonyl pharmacophore, were synthesized to evaluate their biological activities as selective cyclooxygenase-2 (COX-2) inhibitors. In vitro COX-1 and COX-2 isozyme inhibition studies were carried out to acquire structure–activity relationship data with respect to the point that molecular modeling studies showed that designed compounds bind in the primary binding site such that the SO₂Me substituent at *para*-position of C-2 phenyl ring inserts into the 2° pocket present in COX-2 enzyme. COX-1 and COX-2 inhibition studies showed that all compounds were selective inhibitors of the COX-2 isozyme with IC₅₀ values in the highly potent 0.34–0.69 μ M range, and COX-2 selectivity indexes in the 52.3–163.8 range. 1-(4-Fluorophenyl)-2-(4-(methylsulfonyl)phenyl)-4,5,6,7-tetrahydro-1H-benzo[d]imidazole was identified as the most potent (IC₅₀ = 0.34 μ M), and selective (SI = 163.8), COX-2 inhibitor among the synthesized compounds.

Keywords Imidazoles · COX-1 · COX-2

Introduction

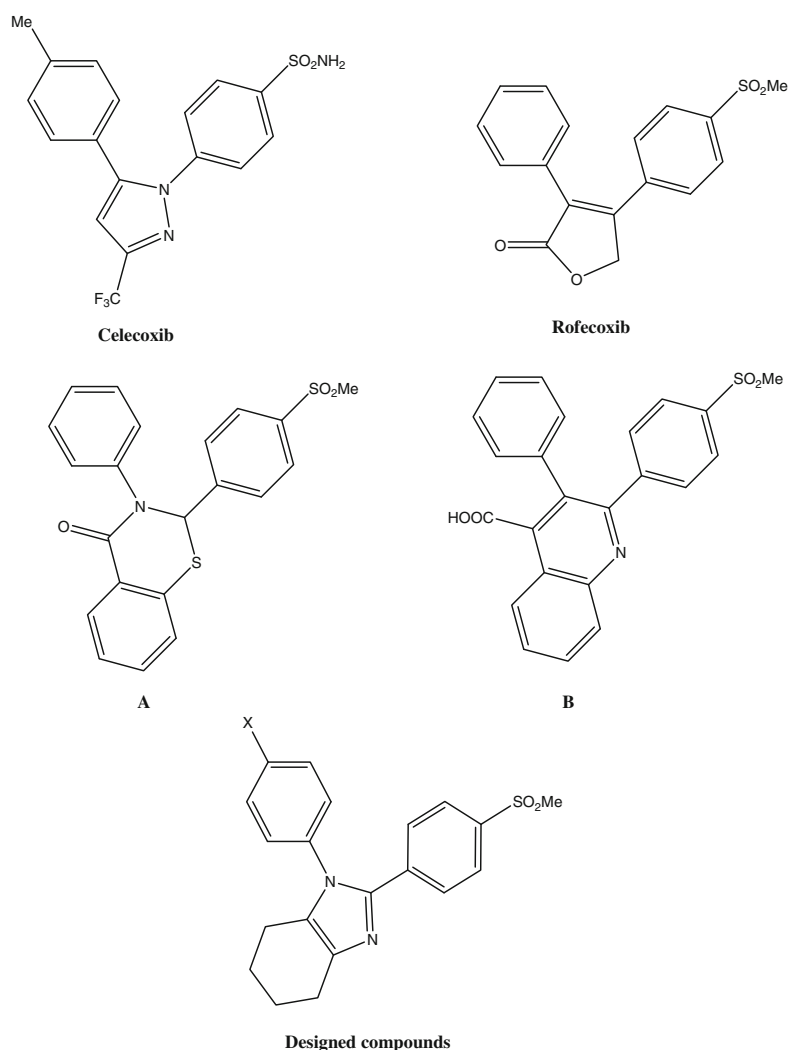
Prostaglandins act as potent mediators of pain, fever, and inflammation. Cyclooxygenase (COX) catalyzes the conversion of arachidonic acid to the prostaglandin endoperoxide PGG₂ then PGH₂, which serves as the precursor for the formation of PGs and thromboxanes (Fu *et al.*, 1990). Nowadays, it is well established that there are at least two COX isozymes, COX-1 and COX-2 (Vane and Botting, 1998). The constitutive COX-1 isozyme is produced in a variety of tissues and appears to be important to the maintenance of physiological functions such as gastro protection and vascular homeostasis (Smith and Dewitt, 1996). Alternatively, the COX-2 isozyme is induced by mitogenic and proinflammatory stimuli linking its involvement to inflammatory processes (Herschman, 1996). Thus, selective inhibition of COX-2 over COX-1 is useful for the treatment of inflammation and inflammation-associated disorders with reduced gastrointestinal toxicities. In addition to role of COX-2 in rheumatoid arthritis and osteoarthritis, it is also implicated in cancer and angiogenesis (Kawamori *et al.*, 1998; Katori and Majima, 2000; Srinath *et al.*, 2003; Takkouche *et al.*, 2008). Recent studies have also shown that selective COX-2 inhibitors may slow the progress of Alzheimer's disease without causing gastrointestinal damage (Nivsarkar *et al.*, 2008). 1,2-Diarylheterocycles have been extensively studied as selective COX-2 inhibitors (Ghodsai *et al.*, 2010; Penning *et al.*, 1997; Prasit *et al.*, 1999; Riendeau *et al.*, 2002; Navidpour *et al.*, 2006; Salimi *et al.*, 2007; Zarghi *et al.*, 2009). All these tricyclic molecules possess 1,2-diaryl substitution on a central hetero or carbocyclic ring system (Fig. 1). Celecoxib and rofecoxib are two typical COX-2 inhibitors in this class. However, the recent withdrawal of some diarylheterocyclic selective COX-2 inhibitors such as

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Fig. 1 Some representative examples of COXIBs (celecoxib and rofecoxib), 1,3-benzthiazine-4-one (**a**), quinoline-4-carboxylic acid (**b**) lead compounds and our 4,5,6,7-tetrahydro-1H-benzo[d]imidazole scaffolds



rofecoxib due to its adverse cardiovascular side effects (Dogné *et al.*, 2005; Solomon, 2005) encourages the researchers to explore and evaluate new structural ring templates possessing COX inhibitory activity. In addition, recent studies have suggested that rofecoxib's adverse cardiac events may not be a class effect but rather related to its particular chemical structure (Dogne *et al.*, 2006). For this reason novel scaffolds with high COX-2 inhibitory activity need to be found and evaluated for their biological activities. Recently, we reported a group of new diaryl-heterocyclic COX-2 inhibitors possessing a bicyclic central ring which exhibited highly selectivity for COX-2 inhibition. (Ghodsi *et al.*, 2010; Zarghi *et al.*, 2009). As part of our ongoing program to design new types of tricyclic selective COX-2 inhibitors, we now report the design, synthesis, cyclooxygenase inhibitory, and some molecular modeling studies of a new group of 1,2-diarylheterocycles possessing a 4,5,6,7-tetrahydro-1H-benzo[d]imidazole central ring and a COX-2 SO₂Me pharmacophore at the *para*-position of C-2 phenyl ring.

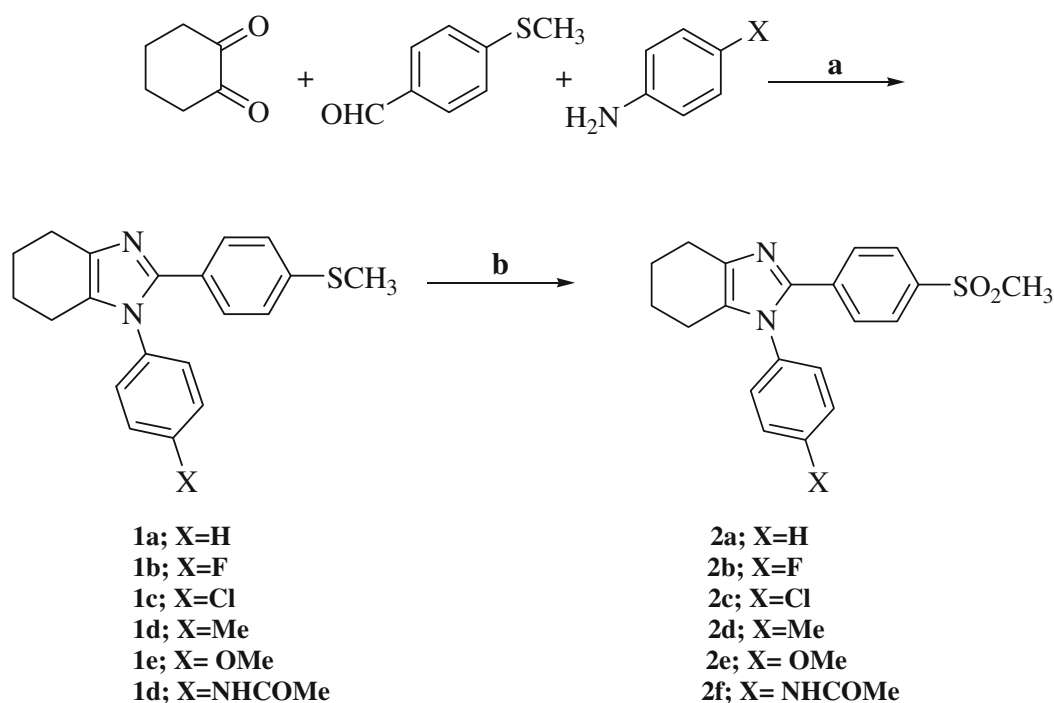
Results and discussions

Chemistry

The target 4,5,6,7-tetrahydro-1H-benzo[d]imidazole derivatives were synthesized via the route outlined in Scheme 1. Accordingly, 1,2-cyclohexanedione, 4-methylthiobenzaldehyde and aniline derivatives in the presence of excess amount of ammonium acetate were treated in microwave reactor at 180 watt power to obtain 1,2-diaryl-4,5,6,7-tetrahydro-1H-benzo[d]imidazoles (**1a–f**). Oxidation of the crude products **1a–f** using Oxone in THF/water afforded the 1-(4-substituted-phenyl)-2-(4-(methylsulfonyl)phenyl)-4,5,6,7-tetrahydro-1H-benzo[d]imidazoles **2a–f** (Yields: 12.7–15.5%).

Enzyme inhibitory activity

It is well established for the diarylheterocyclic class of COX-2 inhibitors that a *para*-methylsulfone substituent on



Scheme 1 Reagents and conditions: **a** NH_4OAc , ACOH , CHCl_3 , microwave 180 watt, 20 min. **b** Oxone, THF, water, 25°C , 2 h

one of the phenyl rings is a requirement for good COX-2 potency and selectivity. Accordingly, a group of 1,2-diaryl-4,5,6,7-tetrahydro-1H-benzo[d]imidazoles (**2a–f**) containing a variety of substituents (H, F, Cl, Me, OMe, and NHCOMe) at the *para*-position of the N-1 phenyl ring, were synthesized to investigate the effect of these substituents on COX-2 selectivity and potency. The ability of the synthesized compounds to inhibit the COX-1 and COX-2 isozymes was determined using chemiluminescent enzyme assays (Table 1). In vitro COX-1/COX-2 inhibition studies showed that all compounds were selective inhibitors of the COX-2 isozyme with IC_{50} values in the highly potent 0.34–0.69 μM range, and COX-2 selectivity indexes (SI) in the 61.2–163.8 range. SAR data (IC_{50} μM values) acquired by determination of the in vitro ability of the title compounds to inhibit the COX-1 and COX-2 isozymes showed that the nature and size of substituent attached to N-1 phenyl ring influenced both selectivity and potency for COX-2 inhibitory activity. Our results indicated that the introduction of suitable substituents such as F (**2b**) and OMe (**2e**) at the *para*-position of N-1 phenyl ring increased both selectivity and potency for COX-2 inhibitory activity. However, compounds having Cl or NHCOMe substituents (**2c** and **2f**) showed less potency for COX-2 isozyme that may be explained by steric parameter. Based upon, compound **2a** containing unsubstituted N-1 phenyl ring showed better COX-2 inhibitory activity compared to compounds **2c** and **2f**. According to these results, 1-(4-fluorophenyl)-2-(4-(methylsulfonyl)phenyl)-4,5,6,7-tetrahydro-1H-benzo[d]

imidazole (**2b**) was the most potent ($\text{IC}_{50} = 0.34 \mu\text{M}$), and selective ($\text{SI} = 163.8$), COX-2 inhibitor among the synthesized compounds. However, it was less selective than

Table 1 In vitro COX-1 and COX-2 enzyme inhibition data

Compound	X	IC_{50} (μM) ^a		COX-2 SI ^b
		COX-1	COX-2	
2a	H	56.5	0.55	102.7
2b	F	55.7	0.34	163.8
2c	Cl	61.6	0.67	91.9
2d	Me	40.1	0.62	64.6
2e	OMe	42.2	0.37	114.0
2f	NHCOMe	36.1	0.69	52.3
Celecoxib		24.3	0.06	405

^a Values are mean values of two determinations acquired using an ovine COX-1/COX-2 assay kit, where the deviation from the mean is <10% of the mean value

^b In vitro COX-2 selectivity index ($\text{COX-1 IC}_{50}/\text{COX-2 IC}_{50}$)

celecoxib ($IC_{50} = 0.06 \mu M$; $SI = 405$) in terms of COX-2 inhibitory activity. These data suggest that the compound **2b** should inhibit the synthesis of inflammatory prostaglandins via the cyclooxygenase pathway at sites of inflammation.

Docking study

The orientation of the highly potent and selective COX-2 inhibitors, 1-(4-fluorophenyl)-2-(4-(methylsulfonyl)phenyl)-4,5,6,7-tetrahydro-1H-benzo[d]imidazole **2b** in the COX-2 active site was examined by a docking experiment (Fig. 2). This molecular modeling showed that compound **2b** binds in the primary binding site such that the C-2 *para*-SO₂Me substituent inserts into the 2° pocket present in COX-2. One of the O-atoms of the SO₂Me moiety forms a H-bond with the NH of Arg⁵¹³ (distance = 2.6 Å), whereas the other O-atom is close to the NH of His⁹⁰ (distance = 3.8 Å). The N-3 atom of the central ring is also close to the NH of Arg¹²⁰ and can form hydrogen bond (distance = 2.9 Å) with this amino acid. It was interesting to note that, the *para*-fluoro substituent of N-1 phenyl ring was forming a hydrogen bond with hydroxyl group (OH) of Tyr³⁸⁵ (distance < 5 Å which may explain the higher potency of compound **2b** compared

with other analogs. These data also suggest that the compound **2b** should inhibit the synthesis of inflammatory prostaglandins via the cyclooxygenase pathway at sites of inflammation.

Conclusion

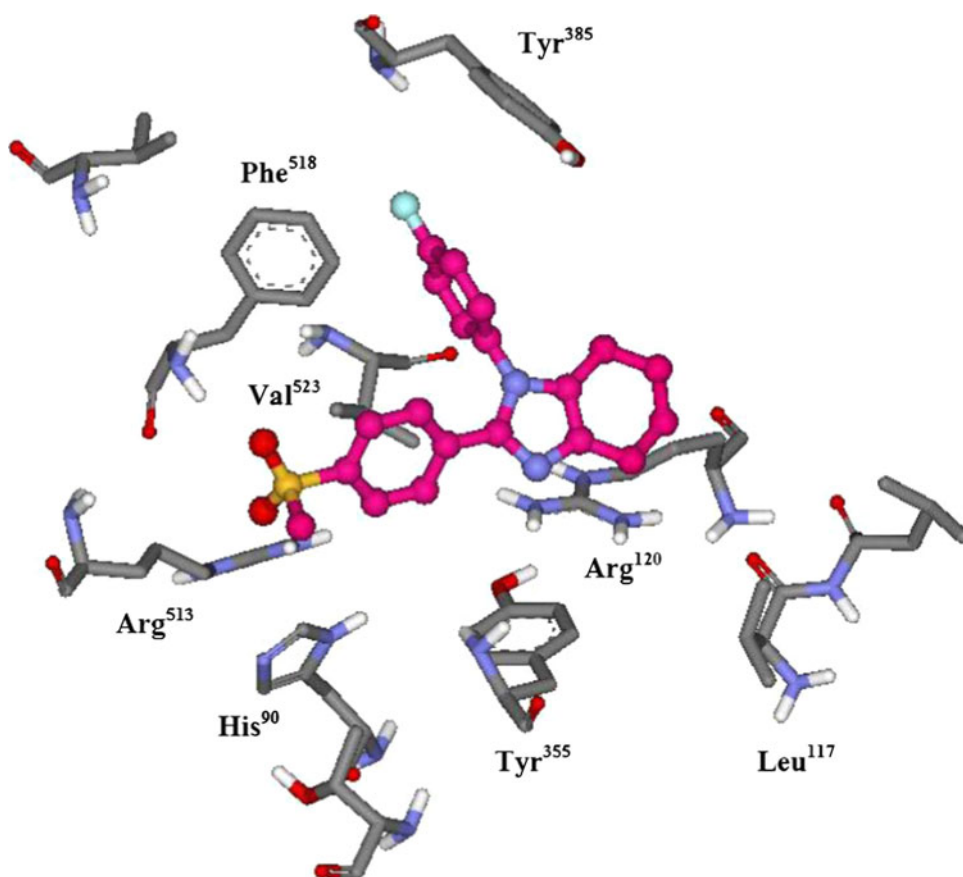
This study indicates that (i) the 4,5,6,7-tetrahydro-1H-benzo[d]imidazole moiety is a suitable scaffold (template) to design COX-1/-2 inhibitors, (ii) COX-1/-2 inhibition is sensitive to the type of substituent at C-2 phenyl ring, and (iii) 1-(4-fluorophenyl)-2-(4-(methylsulfonyl)phenyl)-4,5,6,7-tetrahydro-1H-benzo[d]imidazole (**2b**) exhibited good COX-2 inhibitory potency and selectivity.

Experimental

Materials

All reagents purchased from the Aldrich (USA) or Merck (Germany) Chemical Company and were used without further purifications.

Fig. 2 Compound **2b**: 1-(4-fluorophenyl)-2-(4-(methylsulfonyl)phenyl)-4,5,6,7-tetrahydro-1H-benzo[d]imidazole docked in the active site of murine COX-2 isozyme. Hydrogen atoms of the amino acid residues have been removed to improve clarity



General

Melting points (mp) were determined using a Thomas Hoover capillary apparatus (Philadelphia, USA). Infrared spectra were acquired on a Perkin-Elmer 1420 ratio recording spectrometer. A Bruker FT-500 MHz instrument (Bruker Biosciences, Germany) was used to acquire ^1H NMR spectra; chloroform- D used as solvent. Coupling constant (J) values are estimated in hertz (Hz) and spin multiples are given as s (singlet), d (double), t (triplet), q (quartet), m (multiplet), and br (broad). The mass spectral measurements were performed on a 6410Agilent LCMS triple quadrupole mass spectrometer (LCMS) with an electrospray ionization (ESI) interface. Elemental analyses were carried out with a Perkin-Elmer Model 240-C apparatus (Perkin-Elmer, Norwalk, CT, USA). The results of the elemental analyses (C, H, N) were within $\pm 0.4\%$ of the calculated amounts.

General procedure for preparation of 1-(4-substituted-phenyl)-2-(4-(methyl sulfonyl) phenyl)-4,5,6,7-tetrahydro-1H-benzo[d] imidazoles (**2a–f**)

2 mmol 4-methylthiobenzaldehyde and 2.4 mmol of aniline derivatives were dissolved in 4 ml CHCl_3 and 6 ml glacial acetic acid and placed in microwave reactor with 180 watt power for 5 min. 2 mmol of 1,2-cyclohexanedi-one and excess amount of ammonium acetate were added subsequently and placed in microwave reactor for 15 min with 180 watt power. After cooling, aqueous ammonia solution was added to obtain neutral pH and then extraction with CHCl_3 was performed. The organic phase was washed three times with saturated sodium bicarbonate solution, dried with sodium sulfate, and evaporated in vacuo (Gelens *et al.*, 2006). The crude product was dissolved in THF and excess amount of Oxone in THF/water was added and stirred at room temperature overnight. THF was evaporated and the residue was extracted with chloroform. The organic phase was washed three times with saturated sodium bicarbonate solution and dried with sodium sulfate and concentrated in vacuo. The oily residue was purified by plate chromatography using ethyl acetate–hexane (5:1) as mobile phase (Yields: 12.7–15.5%).

2-(4-(Methylsulfonyl)phenyl)-1-phenyl-4,5,6,7-tetrahydro-1H-benzo[d]imidazole (**2a**)

Yield, 13.5%; white crystalline powder; mp 102–103°C; IR (KBr disk): ν (cm^{-1}) 1150, 1310 (SO_2); ^1H NMR (CDCl_3 , 500 MHz): δ 1.87–1.97 (m, 4H, CH_2), 2.40–2.43 (t, 2H, CH_2), 2.77–2.79 (t, 2H, CH_2), 3.06 (s, 3H, SO_2CH_3), 7.20–7.22 (m, 5H, phenyl), 7.58 (d, 2H, 4-methylsulfonylphenyl H_2 & H_6 , $J = 8.6$ Hz), 7.80 (d, 2H,

4-methylsulfonyl phenyl H_3 & H_5 , $J = 8.6$ Hz); LC–MS (ESI) m/z : 353 ($\text{M} + 1$, 100); Anal. Calcd. for $\text{C}_{20}\text{H}_{20}\text{N}_2\text{O}_2\text{S}$: C, 68.16; H, 5.72; N, 7.95. Found: C, 68.25; H, 5.92; N, 8.01.

1-(4-Fluorophenyl)-2-(4-(methylsulfonyl)phenyl)-4,5,6,7-tetrahydro-1H-benzo[d] imidazole (**2b**)

Yield, 13.5%; white crystalline powder; mp 117–118°C; IR (KBr disk): ν (cm^{-1}) 1150, 1310 (SO_2); ^1H NMR (CDCl_3 , 500 MHz): δ 1.87–1.95 (m, 4H, CH_2), 2.43 (t, 2H, CH_2), 2.78 (t, 2H, CH_2), 3.06 (s, 3H, SO_2CH_3), 7.19–7.22 (m, 4H, 4-fluorophenyl H_2 – H_6), 7.58 (d, 2H, 4-methylsulfonylphenyl H_2 & H_6 , $J = 8.6$ Hz), 7.80 (d, 2H, 4-methylsulfonylphenyl H_3 & H_5 , $J = 8.6$ Hz); LC–MS (ESI) m/z : 371.1 ($\text{M} + 1$, 100); Anal. Calcd. for $\text{C}_{20}\text{H}_{19}\text{FN}_2\text{O}_2\text{S}$: C, 64.85; H, 5.17; N, 7.56. Found: C, 64.99; H, 5.01; N, 7.66.

1-(4-Chlorophenyl)-2-(4-(methylsulfonyl)phenyl)-4,5,6,7-tetrahydro-1H-benzo[d] imidazole (**2c**)

Yield, 14.7%; white crystalline powder; mp 135–136°C; IR (KBr disk): ν (cm^{-1}) 1160, 1320 (SO_2); ^1H NMR (CDCl_3 , 500 MHz): δ 1.87–1.93 (m, 4H, CH_2), 2.42 (t, 2H, CH_2), 2.77 (t, 2H, CH_2), 3.06 (s, 3H, SO_2CH_3), 7.16 (d, 2H, 4-chlorophenyl H_2 & H_6 , $J = 8.6$ Hz), 7.47 (d, 2H, 4-chlorophenyl H_3 & H_5 , $J = 8.6$ Hz), 7.57 (d, 2H, 4-methylsulfonylphenyl H_2 & H_6 , $J = 8.6$ Hz), 7.81 (d, 2H, 4-methylsulfonylphenyl H_3 & H_5 , $J = 8.6$ Hz); LC–MS (ESI) m/z : 387.4 ($\text{M} + 1$, 100); Anal. Calcd. for $\text{C}_{20}\text{H}_{19}\text{ClN}_2\text{O}_2\text{S}$: C, 62.09; H, 4.95; N, 7.24. Found: C, 62.25; H, 5.16; N, 7.22.

1-(4-Methylphenyl)-2-(4-(methylsulfonyl)phenyl)-4,5,6,7-tetrahydro-1H-benzo[d] imidazole (**2d**)

Yield, 15.5%; white crystalline powder; mp 146–147°C; IR (KBr disk): ν (cm^{-1}) 1150, 1310 (SO_2); ^1H NMR (CDCl_3 , 500 MHz): δ 1.85–1.93 (m, 4H, CH_2), 2.42 (t, 2H, CH_2), 2.77 (t, 2H, CH_2), 3.05 (s, 3H, SO_2CH_3), 7.09 (d, 2H, 4-methylphenyl H_3 & H_5 , $J = 8.3$ Hz), 7.29 (d, 2H, 4-methylphenyl H_2 & H_6 , $J = 8.3$ Hz), 7.58 (d, 2H, 4-methylsulfonylphenyl H_2 & H_6 , $J = 8.6$ Hz), 7.78 (d, 2H, 4-methylsulfonylphenyl H_3 & H_5 , $J = 8.6$ Hz); LC–MS (ESI) m/z : 367.1 ($\text{M} + 1$, 100); Anal. Calcd. for $\text{C}_{21}\text{H}_{22}\text{N}_2\text{O}_2\text{S}$: C, 68.82; H, 6.05; N, 7.64. Found: C, 68.55; H, 6.12; N, 7.49.

1-(4-Methoxyphenyl)-2-(4-(methylsulfonyl)phenyl)-4,5,6,7-tetrahydro-1H-benzo[d] imidazole (**2e**)

Yield, 13.3%; white crystalline powder; mp 108–109°C; IR (KBr disk): ν (cm^{-1}) 1150, 1300 (SO_2); ^1H NMR (CDCl_3 , 500 MHz): δ 1.86–1.94 (m, 4H, CH_2), 2.42 (t, 2H, CH_2),

2.83 (t, 2H, CH₂), 3.02 (s, 3H, SO₂CH₃), 3.91 (s, 3H, OCH₃), 7.01 (d, 2H, 4-methoxyphenyl H₃ & H₅, *J* = 8.7 Hz), 7.14 (d, 2H, 4-methoxyphenyl H₂ & H₆, *J* = 8.7 Hz), 7.66 (d, 2H, 4-methylsulfonylphenyl H₂ & H₆, *J* = 8.6 Hz), 7.82 (d, 2H, 4-methylsulfonylphenyl H₃ & H₅, *J* = 8.6 Hz); LC–MS (ESI) *m/z*: 383.1 (*M* + 1, 100); Anal. Calcd. for C₂₁H₂₂N₂O₃S: C, 65.94; H, 5.80; N, 7.32. Found: C, 66.15; H, 5.72; N, 7.65.

1-(4-Acetamidophenyl)-2-(4-(methylsulfonyl)phenyl)-4,5,6,7-tetrahydro-1H-benzo[d]imidazole (2f)

Yield, 12.7%; white crystalline powder; mp 100–101°C; IR (KBr disk): ν (cm⁻¹) 1150, 1320 (SO₂); ¹HNMR (CDCl₃, 500 MHz): δ 1.88–1.91 (m, 4H, CH₂), 2.22 (s, 3H, NHC–OCH₃), 2.41 (t, 2H, CH₂), 2.80 (t, 2H, CH₂), 3.06 (s, 3H, SO₂CH₃), 7.16 (d, 2H, 4-acetamidophenyl H₃ & H₅, *J* = 8.6 Hz), 7.60 (d, 2H, 4-acetamidophenyl H₂ & H₆, *J* = 8.6 Hz), 7.68 (d, 2H, 4-methylsulfonylphenyl H₂ & H₆, *J* = 8.4 Hz), 7.79 (d, 2H, 4-methylsulfonylphenyl H₃ & H₅, *J* = 8.4 Hz); LC–MS (ESI) *m/z*: 410.1 (*M* + 1, 100); Anal. Calcd. for C₂₂H₂₃N₃O₃S: C, 64.53; H, 5.66; N, 10.26. Found: C, 64.75; H, 5.32; N, 10.19.

In vitro COX inhibition assays

The ability of the test compounds listed in Table 1 to inhibit ovine COX-1 and COX-2 (IC₅₀ value, μ M) was determined using chemiluminescent enzyme assays kit (catalog number 560101, Cayman Chemical, Ann Arbor, MI, USA) according to our previously reported method (Zarghi *et al.*, 2007).

Molecular modeling (docking) studies

Docking studies were performed using Autodock software Version 3.0. The coordinates of the X-ray crystal structure of the selective COX-2 inhibitor SC-558 bound to the murine COX-2 enzyme was obtained from the RCSB Protein Data Bank (1cx2) and hydrogens were added. The ligand molecules were constructed using the Builder module and were energy minimized for 1,000 iterations reaching a convergence of 0.01 kcal/mol Å. The energy minimized ligands were superimposed on SC-558 in the PDB file 1cx2 after which SC-558 was deleted. The purpose of docking is to search for favorable binding configuration between the small flexible ligands and the rigid protein. Protein residues with atoms greater than 7.5 Å from the docking box were removed for efficiency. Searching is conducted within a specified 3D docking box using annealing based on the Monte Carlo method and MMFF94 molecular mechanics force field for 8,000 iterations. These docked structures were very similar to the

minimized structures obtained initially. The quality of the docked structures was evaluated by measuring the intermolecular energy of the ligand–enzyme assembly (Kurumbail *et al.*, 1996).

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