ORIGINAL RESEARCH



Synthesis, anti-inflammatory evaluation, and docking studies of some new thiazole derivatives

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Abstract A series of new 2-substituted-*N*-(1,3-thiazole-2-yl) acetamide **3–7** and *N*-(benzo[*d*]thiazol-2-yl)-2-(substituted) acetamide **10–13** derivatives have been synthesized and evaluated in vivo (rat paw edema) for their anti-inflammatory activities and in silico(docking studies) to recognize the hypothetical binding motif of the title compounds with the cyclooxygenase isoenzyme (COX-2) employing GLIDE software (Schrodinger Inc.). The compounds, **10–13** were found to have good anti-inflammatory activities [around 84–93 % of the standard: indomethacin]. The binding mode of the title compounds has been proposed based on the docking studies. Further, the predicted ADME properties of all the tested compounds were found to be in the ranges as predicted by QikProp for 95 % of known oral drugs and alsosatisfy the Lipinski's rule of five.

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Introduction

Nonsteroidal anti-inflammatory drugs (NSAIDs) are important therapeutic agents for the treatment of pain and inflammation of everyday life. Through their anti-inflammatory, antipyretic, and analgesic activities they represent a choice of treatment in various inflammatory diseases such as arthritis and rheumatisms (Stefano et al., 2001). NSAIDs are competitive inhibitors of cyclooxygenase (COX), the enzyme that catalyzes the first step of the biosynthesis of prostaglandins (PGG₂) from arachidonic acid which serves as a precursor for the synthesis of PGs, prostacyclines, and thromboxanes such as TXA₂ that are collectively termed as prostanoids (Vane et al., 1998). It is well established that COX exists in two different isoforms, a constitutive form (COX-1) and an inducible form (COX-2) (Xie et al., 1991). The constitutive COX-1 isozyme is expressed in many tissues and appears to be important for the maintenance of various physiological functions such as cytoprotection of gastric mucosa, regulation of renal blood flow, and platelet aggregation (William and David, 1996). In contrast, COX-2 isozyme is not detected in most normal tissues, but its expression is rapidly induced by stimuli such as mitogenes and oncogenes, growth factors, hormones, and disorders of water-electrolyte homeostasis linking its involvement to pathological processes such as inflammation and various cancer types (Harvey, 1996; Kawamori et al., 1998; Shigeru et al., 2007). The classical NSAIDs act by reducing the production of proinflammatory PGs at the sites of injury via COX-2 inhibition. The side effects associated with these

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NSAIDs such as GI ulcer and renal function suppression are due to the inhibition of COX-1 pathway (Allison et al., 1992; Charles and Raymond, 1995). Thus the success of NSAIDs in treatment of various inflammatory disorders depends on the selective inhibition of COX-2 over COX-1 isoenzyme. Moreover, recent studies indicating the role of COX-2 inhibitors in cancer chemotherapy especially colon cancer (Taketo, 1998; Lisa and Dan, 2010) and neurological diseases such as Parkinson (Van Gool et al., 2003) and Alzheimer's (Paul et al., 2003) diseases still continues to attract investigations on development of COX-2 inhibitors. As a consequence, many selective COX-2 inhibitors have been developed (Balakumar et al., 2010). However, the market withdrawal of some COXIBs such as rofecoxib due to its adverse cardiovascular side effects (Jean et al., 2005) imposed a great challenge to the researchers to explore and evaluate alternative templates with selective COX-2 inhibitory activity (Sara and Afshin, 2010).

Recently, a series of 8/10-trifluoromethyl-substitutedimidazo[1,2-c]quinazolines (Balakumar et al., 2010), 3-alkoxy-4-methanesulfonamido acetophenone derivatives (Alka et al., 2012) cycloalkyl/aryl-3,4,5-trimethylgallates (Mamta et al., 2013) have been reported from our laboratory as potent anti-inflammatory agents. The literature reveals that thiazole derivatives are very potential for their wide range of pharmacological activities including antiinflammatory, antiallergic, antibacterial, antitumor, antihyperlipidemic activities (Afshin et al., 2007; Masakazu et al., 1998; Michel et al., 1997; Jeffery et al., 1999; Pawan and Sawhney, 1997; Rosaria et al., 2005; Franklin et al., 2008, Prakash et al., 2008; Bharti et al., 2010; Balladka et al., 2010). In continuation of our earlier effort, in the present study, we report the synthesis and evaluation of in vivo anti-inflammatory activity of a series of new 2-substituted-N-(1,3-thiazol-2-yl)acetamide (3-7) and N-(benzo[d]thiazol-2-yl)-2-(substituted)acetamide (10-13)derivatives. Further, we propose the molecular interactions and the binding mode of these target compounds using COX-2 isoenzyme based on in silico docking studies.

Experimental

Reactions were routinely monitored by thin layer chromatography (TLC) on silica gel (pre-coated F254 103 Merck plates) and visualized the products under UV light (254 nm). The ¹H was recorded on Bruker Avance II 400 MHz spectrometer for solutions in CDCl₃/DMSO- d_6 in parts per million (δ) downfield from tetramethylsilane as internal standard, and *J* values were given in Hz. The ¹³C NMR spectra were recorded at 100 MHz. The spin multiplicities are indicated by the symbols, s (singlet), d (doublet), t (triplet), dd (double doublet), m (multiplet), and br (broad). IR spectra were recorded on Perkin Elmer FT-IR Spectrometer (Spectrum RX I) using KBr pellet technique. Melting points were recorded in open capillaries on LABINDIA melting point apparatus and were uncorrected. Mass spectra (ESI) were recorded on Waters Micromass Q-TOF Micro.

General procedure for the preparation of 2-chloro-*N*-(1,3-thiazol-2-yl)acetamide (**2**) and *N*-(benzo[*d*]thiazol-2-yl)-2-chloroacetamide (**9**)

Method A

To a solution of 2-amino-1,3-thiazole 1/2-amino-1,3-benzothiazole **8** (0.10 mol) in freshly distilled THF, triethylamine (0.10 mol) was added followed by dropwise addition of chloroacetyl chloride (0.12 mol) at 0–5 °C and allowed to stir for 30 min–3 h (Table 1). The reaction mixture was poured into crushed ice and neutralized with 5 % HCl followed by extraction with EtOAc. The organic phase was dried on anhydrous Na₂SO₄ and concentrated under reduced pressure to get the pure compound 2-chloro-*N*-(1,3-thiazol-2-yl)acetamide **2** and *N*-(benzo[*d*]thiazol-2yl)-2-chloroacetamide **9**.

Method B

A mixture of 2-amino-1,3-thiazole 1/2-amino-1,3-benzothiazole **8** (0.001 mol) in freshly distilled THF and triethylamine and chloroacetylchloride (0.001 mol) was heated under microwave irradiation at 560 W for 10–20 s (Table 1). The reaction mixture was worked up following the same process as mentioned in Method A to get the pure compound 2-chloro-N-(1,3-thiazol-2-yl)acetamide **2** and N-(benzo[d]thiazol-2-yl)-2-chloroacetamide **9**.

2-*Chloro-N-(1,3-thiazol-2-yl)acetamide* (2) White solid, Yield: 51 & 80 % (MW), M.P.: 161–162 °C, IR: 3,496, 3,089, 2,926, 1,676, 1,598 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 3.33 (*s*, 2H, -C<u>H</u>₂), 7.11 (*d*, *J* = 6.8, 1H, Ar-<u>H</u>), 7.23 (*d*, *J* = 6.8, 1H, Ar-<u>H</u>), 10.37 (*s*, 1H, -N<u>H</u>); ¹³C NMR (CDCl₃, 100 MHz) δ 45.44 (Cl-<u>C</u>H₂), 114.22 (<u>C</u>5-1,3-thiazole), 130.66 (<u>C</u>4-1,3-thiazole), 159.93 (<u>C</u>2-1,3thiazole), 166.72 (C=O); ESIMS *m*/*z* 177.68 (M⁺+1).

N-(*benzo[d]thiazol-2-yl*)-2-chloroacetamide (**9**) Slight yellow solid, Yield: 70 & 95 % (MW), M.P.: 162–163 °C, IR: 3,505, 3,048, 2,920, 1,691, 1,646, 1,265 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 4.37 (*s*, 2H, –C<u>H</u>₂–), 7.85 (*m*, 2H, Ar–<u>H</u>), 7.42 (*m*, 2H, Ar–<u>H</u>), 12.63 (*s*, 1H, –N<u>H</u>); ¹³C NMR (CDCl₃, 100 MHz) δ 47.68 (Cl–<u>CH</u>₂), 120.20 (benzo[*d*]thiazole), 122.54 (benzo[*d*]thiazole), 125.59 (benzo[*d*]thiazole), 127.67 (benzo[*d*]thiazole), 132.33 (benzo

Table 1 Comparative data of conventional (Method A) and microwave-assisted (Method B) synthesis of compounds 2-7 and 9-13



Compounds	R	M.P.	Conventional (Method A)		Microwave-assisted (Method B)	
		(°C)	Time (h)	% Yield	Time (s) at 560 W	% Yield
2	Cl	160-162	30 min	51	10	80
3	-N_O	140–142	8 h	35	50	75
4		148–151	8 h	49	50	80
5		160–162	8 h	57	50	80
6	H_3C $-HN$ $-CH_3$	162–164	8 h	60	45	85
7		158–160	8 h	57	50	85
9	Cl	161-162	3 h	70	20	95
10		210–212	2 h	70	30	85
11		190–193	2 h	75	30	87
12	-HN-	146–148	4 h	65	50	88
13		120–122	4 h	65	50	85

M.P. melting point

[*d*]thiazole), 149.94 (<u>benzo</u>[*d*]thiazole), 167.64 (<u>C</u>=O), 175. 55 (C2-benzo[*d*]thiazole); ESIMS *m*/*z* 227.93 (M⁺+1).

General procedure for the preparation of 2-(substituted)-*N*-(1,3-thiazol-2-yl)acetamide (**3**–**7**) and *N*-(benzo[*d*]thiazol-2-yl)-2-(substituted)acetamide (**10–13**)

Method A

A mixture of compound 2/9 (0.10 mol) in freshly distilled THF and triethylamine and appropriate amines (0.10 mol) was heated under reflux for 2–8 h (Table 1). The reaction

mixture was poured into crushed ice and neutralized with 5 % HCl followed by extraction with EtOAc. The organic phase was dried on anhydrous Na_2SO_4 and concentrated under reduced pressure. The residue was purified by column chromatography (EtOAc-Hexane 8:2) to afford the desired compound 2-(*substituted*)-N-(1,3-thiazol-2-yl)acet-amide (3–7) and N-(*benzo[d]thiazol-2-yl)-2-(substituted*) acetamide (10–13) as a solid.

Method B

A mixture of compound **2/9** (0.001 mol) in freshly distilled THF and triethylamine and appropriate amines (0.001 mol)

was heated under microwave irradiation at 560 W for 30-50 s (Table 1). The reaction mixture was worked up following the same process as mentioned in Method A to get the pure compound **3–7** and **10–13**.

2-Morpholino-N-(1,3-thiazol-2-yl)acetamide (3) White solid, Yield: 35 & 75 % (MW), M.P.: 140–142 °C, IR: 3,089, 2,928, 1,676, 1,598 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 2.55 (*t*, *J* = 4.6, 4H, -C<u>H</u>₂–), 3.19 (*s*, 2H, -C<u>H</u>₂), 3.70 (*t*, *J* = 4.4, 4H, -C<u>H</u>₂–), 6.93 (*d*, *J* = 6.8, 1H, Ar-<u>H</u>), 7.38 (*d*, *J* = 6.8, 1H, Ar-<u>H</u>), 10.37 (*s*, 1H, -N<u>H</u>); ¹³C NMR (CDCl₃, 100 MHz) δ 56.28 (CH₂), 61.39 (CH₂– N-CH₂), 64.23 (CH₂–O-CH₂), 114.53 (C5-1,3-thiazole), 130.91 (C4-1,3-thiazole), 160.84 (C2-1,3-thiazole), 166.33 (C=O); ESIMS *m*/z 228.2 (M⁺+1).

2-(*Phenylamino*)-*N*-(1,3-thiazol-2-yl)acetamide (4) White solid, Yield: 49 & 80 % (MW), M.P.: 148–151 °C, IR: 3,350, 3,030, 2,965, 1,669, 1,610, 1,590 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 3.87 (*s*, 2H, -CH₂), 4.12 (*s*, 1H, -NH), 7.01(*d*, *J* = 6.9, 1H, Ar-<u>H</u>), 7.23 (*m*, 3H, Ar-<u>H</u>), 7.42 (*d*, *J* = 6.9, 1H, Ar-<u>H</u>), 7.51 (*d*, *J* = 7.2, 2H, Ar-<u>H</u>), 10.41 (*s*, 1H, -N<u>H</u>); ¹³C NMR (CDCl₃, 100 MHz) δ 54.28 (CH₂), 111.18 (<u>C</u>5-1,3-thiazole), 112.61 (<u>C</u>3'&<u>C</u>5'-phenyl), 118.76 (<u>C</u>4'-phenyl), 130.33 (<u>C</u>2'&C6'-phenyl), 131.89 (<u>C</u>4-1,3-thiazole), 145.22 (<u>C</u>1'-phenyl), 161.29 (<u>C</u>2-1,3-thiazole), 166.72 (<u>C</u>=O); ESIMS *m*/z 234.1 (M⁺+1).

2-(*o*-Toluidino)-N-(1,3-thiazol-2-yl)acetamide (5) White solid, Yield: 57 & 80 % (MW), M.P.: 160–162 °C, IR: 3,345, 3,029, 2,964, 1,670, 1,620, 1,514 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 2.2 (*s*, 3H, -C<u>H</u>₃), 3.90 (*s*, 2H, -C<u>H</u>₂), 4.01 (*s*, 1H, -N<u>H</u>), 6.49 (*d*, *J* = 7.2, 1H, Ar-<u>H</u>), 6.97 (*m*, 4H, Ar-<u>H</u>), 7.41 (*d*, *J* = 7.2, 1H, Ar-<u>H</u>), 11.62 (*s*, 1H, -N<u>H</u>); ¹³C NMR (CDCl₃, 100 MHz) δ 18.22 (CH₃), 55.11 (CH₂), 113.32 (C5-1,3-thiazole), 118.11 (C2'-phenyl), 119.22 (C3'-phenyl), 121.39 (C5'-phenyl), 125.66 (C4'-phenyl), 127.88 (C6'-phenyl), 131.56 (C4-1,3-thiazole), 144.29 (C1'-phenyl), 161.46 (C2-1,3-thiazole), 166.39 (C=O); ESIMS *m*/z 248.2 (M⁺+1).

2-(*p*-Toluidino)-*N*-(1,3-thiazol-2-yl)acetamide (**6**) White solid, Yield: 60 & 85 % (MW), M.P.: 162–164 °C, IR: 3,356, 3,123, 2,913, 1,664, 1,616, 1,521 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 2.24 (*s*, 3H, -C<u>H</u>₃), 4.02 (*s*, 2H, -C<u>H</u>₂), 4.13 (*s*, 1H, -N<u>H</u>), 6.56 (*d*, *J* = 6.8, 2H, Ar-<u>H</u>), 7.00 (*d*, *J* = 7.1, 1H, Ar-<u>H</u>), 7.02 (*d*, *J* = 6.8, 2H, Ar-<u>H</u>), 7.42 (*d*, *J* = 7.1, 1H, Ar-<u>H</u>), 10.89 (*s*, 1H, -N<u>H</u>); ¹³C NMR (CDCl₃, 100 MHz) δ 20.43 (<u>CH</u>₃), 55.34 (<u>CH</u>₂), 112.94 (<u>C</u>5-1,3-thiazole), 114.32 (<u>C</u>3'&<u>C</u>5'-phenyl), 129.11 (<u>C</u>4'-phenyl), 130.18 (<u>C</u>2'&<u>C</u>6'-phenyl), 131.11 (<u>C</u>4-1,3-

thiazole), 145.33 (<u>C</u>1'-phenyl), 161.29 (<u>C</u>2-1,3-thiazole), 167.22 (<u>C</u>=O); ESIMS m/z 248.3 (M⁺+1).

2-(4-Methoxyphenylamino)-N-(1,3-thiazol-2-yl)acetamide (7) White solid, Yield: 57 & 85 % (MW), M.P.: 158–160 °C, IR: 3,348, 3,041, 2,898, 1,661, 1,515, 1,238, 1,033 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 3.72 (*s*, 3H, –OCH₃), 3.95 (*s*, 2H, –CH₂), 4.13 (*s*, 1H, –NH), 6.60 (*d*, J = 7.0, 1H, Ar–H), 6.78 (*d*, J = 7.2, 2H, Ar–H), 7.00 (*d*, J = 7.2, 2H, Ar–H), 7.42(*d*, J = 7.0, 1H, Ar–H), 11.12 (*s*, 1H, –NH); ¹³C NMR (CDCl₃, 100 MHz) δ 55.26 (CH₂), 56.38 (OCH₃), 112.72 (C5-1,3-thiazole), 115.45 (C3'&C5'phenyl), 116.47 (C2'&C6'-phenyl), 131.66 (C4-1,3-thiazole), 138.76 (C4'-phenyl), 154.24 (C1'-phenyl), 162.11 (C2-1,3-thiazole), 167.46(C=O); ESIMS *m*/*z* 264.1 (M⁺+1).

N-(*benzo*[*d*]*thiazo*1-2-*y*]*)*-2-(*1H*-1,2,4-*triazo*1-1-*y*]*)acetamide* (*10*) White solid, Yield: 70 & 85 % (MW), M.P.: 240 °C, IR: 3,045, 2,934, 1,668, 1,625, 1,267 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 4.01 (*s*, 2H, CH₂), 7.39 (*m*, 2H, Ar–H), 7.71 (*d*, *J* = 6.8, 1H, Ar–H), 7.88 (*s*, 1H, N=CH), 7.98 (*s*, 1H, N = CH), 8.75 (*d*, *J* = 7.0, 1H, Ar–H), 12.43 (*s*, 1H, NH); ¹³C NMR (CDCl₃, 100 MHz) δ 56.57 (CH₂), 120.32 (<u>benzo</u>[*d*]thiazole), 122.63 (<u>benzo</u>[*d*]thiazole), 125.16 (<u>benzo</u>[*d*]thiazole), 127.39 (<u>benzo</u>[*d*]thiazole), 132.54 (<u>benzo</u> [*d*]thiazole), 144.14 (<u>C</u>5'-triazole), 148.11 (<u>C</u>3'-triazole), 149.81 (<u>benzo</u>[*d*]thiazole), 169.33 (<u>C</u>2-benzo[*d*]thiazole), 175.44 (<u>C</u>=O); ESIMS *m*/*z* 260.4 (M⁺+1).

N-(*benzo*[*d*]*thiazol*-2-*yl*)-2-(*1H*-*imidazol*-1-*yl*)*acetamide* (*11*) White solid, Yield: 75 & 87 % (MW), M.P.: 230 °C, IR: 3,042, 2,920, 1,662, 1,646, 1,260 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 4.12 (s, 2H, CH₂), 7.01 (*d*, *J* = 6.6, 1H, N–CH=CH), 7.32 (*d*, *J* = 7.0, 1H, N–CH=CH–), 7.45 (*s*, 1H, N=CH), 7.56 (*m*, 2H, Ar–H), 8.01 (*d*, *J* = 7.0, 1H, Ar–H), 8.12 (*d*, *J* = 7.3, 1H, Ar–H), 11.98 (*s*, 1H, NH); ¹³C NMR (CDCl₃, 100 MHz) δ 39.47 (CH₂), 119.23 (<u>benzo</u>[*d*]thiazole), 121.18 (C4'-imidazole), 122.30 (<u>benzo</u>[*d*] thiazole), 128.49 (C5'-imidazole), 132.36 (<u>benzo</u>[*d*]thiazole), 168.91 (C2-benzo[*d*]thiazole), 175.12 (C=O); ESIMS *m*/z 259.3 (M⁺+1).

N-(*benzo*[*d*]*thiazo*1-2-*y*1)-2-(*4*-*fluorophenylamino*)*acetamide* (*12*) White solid, Yield: 64 & 88 % (MW), M.P.: 146–148 °C, IR: 3,398, 3,048, 2,930, 1,675, 1,648, 1,380, 1,252 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 4.01 (*s*, 2H, C<u>H</u>₂), 4.15 (*s*, 1H, N<u>H</u>), 7.02 (*d*, *J* = 6.9, 2H, Ar–<u>H</u>), 7.12 (*d*, *J* = 7.0, 2H, Ar–<u>H</u>), 7.45 (*m*, 2H, Ar–<u>H</u>), 7.73 (*d*, *J* = 7.1, 1H, Ar–<u>H</u>), 7.80 (d, J = 7.3, 1H, Ar–<u>H</u>), 11.12 (s, 1H, –N<u>H</u>); ¹³C NMR (CDCl₃, 100 MHz) δ 54.91 (<u>CH</u>₂), 115.34 (<u>C2'&C6'-4-fluorophenyl</u>), 118.96 (<u>benzo[</u>*d*]thiazole), 119.12 (<u>C3'&C5'-4-fluorophenyl</u>), 122.11 (<u>benzo[</u>*d*]thiazole), 124.84 (<u>benzo[</u>*d*]thiazole), 126.88 (<u>benzo[</u>*d*]thiazole), 132.29 (<u>benzo</u>[*d*]thiazole), 146.32 (<u>C1'-4-fluorophenyl</u>), 152.33 (<u>benzo[</u>*d*]thiazole), 154.11 (<u>C4'-4-fluorophenyl</u>), 169.21 (<u>C2-benzo</u>[*d*]thiazole), 174.93 (<u>C</u>=O); ESIMS *m*/*z* 302.1 (M⁺+1).

*N-(benzo[d]thiazol-2-yl)-2-(4-methoxyphenylamino)acet*amide (13) White solid, Yield: 65 & 85 % (MW), M.P.: 120-122 °C, IR: 3,389, 3,041, 2,920, 1,692, 1,646, 1.265 cm⁻¹; ¹H NMR (CDCl₃, 400 MHz) δ 3.82 (s, 3H, OCH_3 , 4.11 (s, 2H, CH₂), 5.50 (s, 1H, NH), 6.95 (d, J = 6.7, 2H, Ar–H), 7.51 (d, J = 6.8, 2H, Ar–H), 7.65 (m, 2H, Ar–H), 7.8 (d, J = 7.1, 1H, Ar-H), 7.91 (d, J = 7.3, 1H, Ar-H), 12.61(s, 1H, NH); ¹³C NMR (CDCl₃, 100 MHz) δ 54.77 (CH₂), 56.13 (OCH₃), 116.52 (C2'&C6'-4-methoxyphenyl), 117.33 119.14 (C3'&C5'-4-methoxyphenyl), (benzo[d]thiazole), 122.20 (benzo[d]thiazole), 123.11 (benzo[d]thiazole), 125.97 (benzo[d]thiazole), 132.67 (benzo[d]thiazole), 138.73 (C1'-4-methoxyphenyl), 152.92 (benzo[d]thiazole), 152.11 (C4'-4-methoxyphenyl), 169.33 (C2-benzo[d]thiazole), 174.74 (C=O); ESIMS m/z 314.2 (M⁺+1).

Pharmacological evaluation: anti-inflammatory activity-carrageenan-induced rat paw edema assay

The in vivo anti-inflammatory screening for the synthesized compounds was performed by using the functional model of carrageenan-induced rat paw edema and is presented as the percentage inhibition of edema at the right hind paw in comparison to the control (Table 2). Carrageenan-induced edema is a nonspecific inflammation, but is highly sensitive to NSAIDs. Indomethacin, a potent NSAID was used as a reference standard. COX-2-mediated increase in prostaglandin E₂ (PG-E₂) production contributes to the severity of the inflammatory and pain responses in this model. Wistar rats (female) weighing between 190 and 260 g, obtained from Central Animal House, Panjab University, Chandigarh, India and were used in the present study. Animals were kept in wire-mesh cages and maintained under constant environmental food and water (ad libitum), in a constant lightdark cycle. During the course of experiment, the general behavior of animals was normal. All the experimental protocols were approved by the Institutional Animal Ethical Committee (IACE), and experiments were conducted in accordance with the standard guidelines. Carrageenan and indomethacin were procured from Sigma Chemical, Co. For statistical analysis, we have used GraphPad Prism 3.0 version (GraphPad Software Inc (2007).

The synthesized compounds were tested for their antiinflammatory activity against carrageenan-induced paw edema at dose of 10 mg/kg. The percentage inhibition of edema was calculated using the formula given below:

$$\frac{V_{\rm c}-V_{\rm t}}{V_{\rm c}} imes 100$$

where V_c is the increase in paw volume of control (in the absence of test compound), and V_t is the increase in paw volume after administration of the test compound.

The given data are of five animals (female wistar rats) per group and divided into eleven (11) groups. Group 1 referred as control (animals received carrageenan along with the vehicle [0.5 % carboxymethylcellulose (CMC)]) and Group 2 received standard reference (indomethacin) along with the vehicle prior to the administration of carrageenan. Group 3, 4, 5, 6, 7, 8, 9, 10 and 11 received the test compounds (3-7 and 10-13), respectively 1 h prior to the administration of carrageenan. All the test compounds were suspended in 0.5 % of CMC and administered orally (10 mg/kg) 60 min prior to the injection of 0.1 ml of freshly prepared carrageenan (1 %) in physiological solution (154 mM NaCl) into the subplanter tissue of hind paw of each rat. The equivalent volume of carrageenan (1 %) in physiological solution was injected into hind paw of the control. The volume was measured three times using water plethysmometer prior to the administration of carrageenan, 2 and 4 h after the injection. The increase in volume of the paw was adopted as a measure of edema (Winter et al., 1962). The antiedematous effects of the compounds were estimated as percentage inhibition of the induced inflammation in comparison with control. Statistical analysis was carried out using a one-way analysis of variance (ANOVA). In all cases, post hoc comparisons of the means of individual groups were performed using Tukey's test. A significance level of p < 0.001 denoted significance in all cases.

Molecular modeling study

Molecular modeling investigations were carried out using Dell Precision work station T3400 running Intel Core2 Duo Processor, 4 GB RAM, 250 GB hard disk, and NVidia Quodro FX 4500 graphics card. Maestro 9.4, GLIDE v5.9 XP docking program, Schrodinger Inc. (Maestro version and 9.4, 2013; Richard *et al.*, 2004) was employed for the docking studies.

Preparation of protein

PDB structure (www.rcsb.org) **1CX2** (Crystal structure of COX-2) was downloaded, refined, and prepared using Schrodinger protein preparation wizard tool (Glide), which performs the following steps: assigning of bond orders, addition of hydrogens, optimization of hydrogen bonds by

Table 2 Anti-inflammatory data of compounds 3-7 and 10-13 on carrageenan-induced paw edema in rats



Treatment	R	Volume of paw edema (ml	Volume of paw edema (ml)		
		2 h	4 h		
Control (carrageenan treated)	_	0.93 ± 0.012	0.90 ± 0.020		
Indomethacin ^b	_	$0.21 \pm 0.023^{\rm a} \ (77.42)$	$0.19 \pm 0.019^{\rm a} \ (78.89)$		
3	—NO	0.36 ± 0.015^{a} (61.29)	$0.32 \pm 0.021^{a} (64.44)$		
4		$0.41 \pm 0.020^{a} \ (55.91)$	$0.40 \pm 0.035^{\mathrm{a}} (55.55)$		
5		0.39 ± 0.042^{a} (58.06)	$0.37 \pm 0.030^{\mathrm{a}} \ (58.88)$		
6		$0.37 \pm 0.036^{a} \ (60.22)$	0.34 ± 0.025^{a} (62.22)		
7		$0.36 \pm 0.017^{a} \ (61.29)$	$0.31 \pm 0.030^{\mathrm{a}} \ (65.55)$		
10		$0.32 \pm 0.019^{a} \ (65.59)$	$0.29 \pm 0.021^{\mathrm{a}} \ (67.77)$		
11		0.29 ± 0.016^{a} (68.81)	$0.27 \pm 0.070^{a} \ (70.00)$		
12	-HN-	0.26 ± 0.019^{a} (72.04)	0.23 ± 0.090^{a} (74.44)		
13		$0.27 \pm 0.023^{\mathrm{a}}$ (70.97)	0.25 ± 0.013^{a} (72.22)		

Values are expressed as mean \pm S.E.M (n = 5) and analyzed by ANOVA

Values in parenthesis (percentage inhibition of edema), 10 mg/kg

^a Different from carragenan group (p < 0.001)

^b Reference standard

flipping amino side chains, correction of charges, and minimization of the protein complex. All the bound water molecules, ligands and cofactors, were removed (preprocess) from the proteins which were taken in.mae format. The tool neutralized the side chains that are not close to the binding cavity and do not participate in salt bridges. This step is then followed by restrained minimization of cocrystallized complex, which reorients side chain hydroxyl groups and alleviates potential steric clashes. The complex obtained was minimized using OPLS_2005 force field (George and Richard, 2001) with Polack-Ribiere Conjugate Gradient (PRCG) algorithm. The minimization was terminated either completion of 5,000 steps or after the energy gradient converged below 0.05 kcal/mol.

Preparation of ligands

Structures of the ligands (3–7 and 10–13) were sketched using built panel of Maestro and taken in.*mae* format. Lig-Prep is a utility of Schrodinger software suit that combines tools for generating 3D structures from 1D (Smiles) and 2D (SDF) representation, searching for tautomers, steric isomers, and perform a geometry minimization of the ligands. Molecular Mechanics Force Fields (OPLS_2005) with default settings were employed for the ligand minimization.

Docking studies

Docking studies were carried out using the above mentioned prepared protein (PDB: **1CX2**) and ligands (**3–7** and **10–13**), by employing Glide XP docking program (Schrodinger Inc.) following the reported procedure (Maestro version 9.4, and Glide v5.9, 2013; Richard *et al.*, 2004).

Prediction of ADME properties

The QikProp module of Schrodinger is a quick, accurate, easyto-use absorption, distribution, metabolism, and excretion (ADME) prediction program design to produce certain descriptors related to ADME. QikProp predicts physically significant descriptors and pharmaceutically relevant properties of organic molecules, either individually or in batches. QikProp has two modes: normal mode and fast mode. In fast mode, certain time-consuming calculations are omitted; some properties are not predicted, and some have different values. In the present study, QikProp was run in normal processing mode with default options (QikProp, version 3.6, 2013). To set up the calculation, pose viewer file (generated after docking with Glide) was used to consider the receptor and source of ligands. After choosing the receptor and ligands, by using pose viewer file (*pv.maegz*), the program QikProp that generates the descriptors were run with default options that were chosen to produce reasonable descriptors. The selected properties that are known to influence metabolism, cell permeation, and bioavailability are presented in Table 3.

Results and discussion

Chemistry

As illustrated in Scheme 1, target compounds were synthesized from 2-amino-1,3-thiazole **1** which on reaction with chloroacetvl chloride in the presence of tetrahydrofuran (THF) using triehtylamine as base maintaining the temperature between 0 and 5 °C with constant stirring resulted in the formation of 2-chloro-N-(1,3-thiazol-2vl)acetamide 2 in quantitative yield (Method A) (Narendra et al., 2010). Similarly, the compound 2 was also synthesized in good yields by irradiating 2-amino-1,3-thiazole 1 with chloroacetvl chloride in the presence of THF using triehtylamine at 560 W hitherto unreported in the literature (Method B). The compound 2 was characterized based on a typical NH absorption at $3,496 \text{ cm}^{-1}$, a strong signal at 1,676 cm⁻¹ of the carbonyl (C=O) group, H-Ar absorption at $3,089 \text{ cm}^{-1}$, and $-CH_2$ - absorption at 2.926 cm⁻¹. The ¹H NMR spectrum showed characteristic singlet at δ 3.33 (2H, -CH₂) ppm assignable to the two methylene protons (attached to chlorine atom). A broad singlet at δ 10.37 ppm corresponding to amine proton (–NH), and two doublets at δ 7.11 and 7.23 ppm integrated to one proton each and assignable to thiazole protons confirmed the formation of compound 2. The ¹³C NMR signal at δ 45.44 and 166.72 ppm indicated the presence of methylene carbon (attached to chlorine atom) and carbonyl carbon (C=O), respectively. The thiazole carbons appeared at δ 114.22 ppm (C5, thiazole), δ 130.66 ppm (C4, thiazole), and δ 159.93 ppm (C2, thiazole) with good intensity further confirmed the formation of the key intermediate compound 2. The compound 2 on reaction with various aryl amines in basic condition under reflux for 8 h resulted in the formation of target compounds 3-7 in quantitative vields. They were also synthesized in good vields by irradiating compound 2 with aryl amines under similar basic condition at 560 W. Compounds 3-7 were characterized based on a typical NH absorption at $3,240 \text{ cm}^{-1}$, a strong signal at 1,702 cm^{-1} of the carbonyl group and H–Ar absorption at $3,030 \text{ cm}^{-1}$ (except in compound 3 with $-CH_2$ absorption at 2,928 cm⁻¹). Moreover, the appearance of an additional new broad singlet of NH signal around δ 5 ppm (except in compound 3) along with the NH signal of amide bond around δ 10 ppm and aromatic protons of various aryl amines at around 7.5 ppm confirmed the formation of target compounds (3-7). The downfield shift of ¹³C NMR signal around δ 55 ppm indicated the attachment of aromatic substituents (except compound 3 with morpholine substituent) to methylene carbon. The appearance of downfield shift of 13 C NMR signal at δ 61.39 and 64.23 ppm, respectively assignable to methylene carbons of morpholine moiety confirmed the formation of compound 3. Moreover, the appearance of aromatic carbons around δ 111–140 ppm confirmed the formation of target compounds (4–7). The ¹³C NMR signal at δ 18.22, 20.43, and 56.38 ppm, respectively, assignable to methyl carbon (-CH₃) attached to ortho and para position of aromatic moiety of compound 5 and 6 as well as methoxy

Table 3 Properties of tested compounds calculated by QikProp

Comp No.	QPlogPo/w ^a	QPlogS ^b	QPPCaco ^c	#metab ^d	Percent human oral absorption ^e	Lipinski's rule of five ^f
3	0.753	-2.507	317.043	3	76.122	0
4	0.488	-2.223	297.565	3	74.078	0
5	0.384	-2.045	319.565	3	74.019	0
6	1.58	-3.654	408.183	3	82.927	0
7	3.016	-5.036	603.636	3	94.375	0
10	2.376	-4.548	527.28	3	89.575	0
11	0.736	-3.049	182.19	3	71.714	0
12	2.337	-4.656	267.979	4	84.089	0
13	1.3	-3.228	390.352	3	80.941	0

^a Predicted log of the octanol/water partition coefficient, range 95 % of drugs (-2 to 6.5)

^b Predicted log of aqueous solubility S (mol/l), range 95 % of drugs (-6.5 to 0.5)

^c Caco2 cell permeability in nm/s, range 95 % of drugs (<25 poor, >500 great). Caco-2 cells are a model for the gut-blood barrier. QikProp predictions are for nonactive transport.

^d Number of likely metabolic reactions. Range 95 % of drugs (1-8).

^e Predicted human oral absorption on 0-100 % scale (>80 % is high, and <25 % is poor)

^f Number of violations of Lipinski's rule of five. The rules are: mol_MW < 500, QPlogPo/w < 5, donorHB \leq 5, accptHB \leq 10. Compounds that satisfy these rules are considered druglike. (The "five" refers to the limits, which are multiples of 5.)

carbon $(-OCH_3)$ attached to *para* position of compound 7. Further, molecular ion peak (ESI–MS) corresponding to its molecular formula revealed the formation of all the products.

Similarly, the target compounds 10-13 were synthesized following scheme-2, starting from 2-amino-benzo[d]thiazole 8 which on reaction with chloroacetyl chloride in the presence of tetrahydrofuran (THF) using triehtylamine as a base maintaining the temperature between 0 and 5 °C with constant stirring resulted in the formation of N-(benzo[d]thiazol-2-yl)-2-chloroacetamide 9 in quantitative yield (Method A) (Guangfu et al., 2001). Similarly, the compound 9 was also synthesized in good yields by irradiating 2-aminobenzo[d]thiazole 8 with chloroacetyl chloride under similar basic condition at 560 W hitherto unreported in the literature (Method B). The compound 9 was characterized based on a typical NH absorption at 3,505 cm⁻¹, a strong signal at 1,691 cm⁻¹ of the carbonyl (C=O) group, H–Ar absorption at 3,048 cm⁻¹, and –CH₂– absorption at 2,920 cm⁻¹. The ¹H NMR spectrum showed characteristic singlet at δ 4.37 (2H, -CH₂) ppm assignable to the two methylene protons (attached to chlorine atom). A broad singlet at δ 12.63 ppm corresponding to amine proton (–NH) and two multiplets at δ 7.85 and 7.42 ppm, respectively, integrated to two protons each and assignable to aromatic protons confirmed the formation of compound 9. The 13 C NMR signal at δ 47.68 and 167.64 ppm indicated the presence of methylene carbon (attached to chlorine atom) and carbonyl carbon (C=O), respectively. The benzo[d]thiazole carbons appeared at δ 120.20 (benzo[d]thiazole), 122.54 (benzo[d]thiazole), 125.59 (benzo[d]thiazole),

127.67 (benzo[d]thiazole), 132.33 (benzo[d]thiazole), 149.94 (benzo[d]thiazole), 175.55 (C2-benzo[d]thiazole) ppm, respectively, with good intensity further confirmed the formation of the key intermediate compound 9. The compound 9 on reaction with various aryl amines in basic condition under reflux for 2-4 h resulted in the formation of target compounds 10-13 in quantitative yields. They were also synthesized in good yields by irradiating 9 with various aryl amines in the presence of THF and triethylamine at 560 W. Compounds 10-13 were characterized based on a typical NH absorption at 3,505 cm⁻¹, a strong signal at 1,702 cm⁻¹ indicating carbonyl group and H–Ar absorption at 3,030 cm $^{-1}$. Further, the appearance of an additional new broad singlet of NH signal around δ 6.5–7 ppm (compounds 12 & 13) and typical triazole and imidazole proton peak at δ 7–8 ppm (compounds 10 & 11) along with the NH signal of amide bond around δ 12 ppm and aromatic protons of various aryl amines at around 7.5 ppm confirmed the formation of target compounds 10–13. The downfield shift of 13 C NMR signal around δ 55 ppm indicated the replacement of chlorine atom and attachment of aromatic substituents to methylene carbon. The appearance of downfield shift of ¹³C NMR signal at δ 144.14 (C5'-triazole), 148.11 (C3'-triazole) ppm, respectively assignable to triazole carbons confirmed the formation of compound 10. Similarly, the appearance of ¹³C NMR signal at δ 121.18 (C4'-imidazole), 128.49 (C5'-imidazole), 137.70 (C2'-imidazole) ppm, respectively assignable to imidazole carbons confirmed the formation of compound 11. Moreover, the appearance of additional aromatic carbon signals around δ 115–150 ppm confirmed the formation of target compounds 12 and 13. The Scheme 1 Synthesis of 2-(substituted)-*N*-(thiazol-2yl)acetamide derivatives. Reagents and conditions: (*i*) Method A, ClCH₂COCl, THF, triethyl amine, 0–5 °C, 30 min; (*ii*) Method B, ClCH₂COCl, THF, triethyl amine, MW 560 W, 10 s; (*iii*) Method A, aryl amines, THF, triethyl amine, reflux, 8 h; (*iv*) Method B, Aryl amines, THF, Triethyl amine, MW 560 W, 45–50 s

Scheme 2 Synthesis of *N*-(benzo[*d*]thiazol-2-yl)-2-(substituted)acetamide derivatives. Reagents and conditions: (*i*) Method A, ClCH₂COCl, THF, Triethyl amine, $0-5 \,^{\circ}C$, 3 h; (*ii*) Method B, ClCH₂COCl, THF, Triethyl amine, MW 560 W, 20 s; (*iii*) Method A, Aryl amines, THF, Triethyl amine, reflux, 2–4 h; (*iv*) Method B, Aryl amines, THF, Triethyl amine, MW 560 W, 30–50 s



¹³C NMR signal at δ 56.13 ppm assignable to methoxy carbon (–OCH₃) attached to *para* position of compound **13**. Further, molecular ion peak (ESI–MS) corresponding to its molecular formula revealed the formation of the product.

Under classical heating conditions, these reactions have certain disadvantages like long reaction times (30 min–8 h), high energy consumption and the need for large amounts of solvents for work up and purification. The MW-assisted reactions were carried out using a Catalyst Microwave Reactor, under constant irradiation power and by varying the temperature (the so-called "power control"). The best results were obtained when we used 80 % of the full power of the magnetron (560 W). The details of the optimized conditions employed, under MW irradiation as well as under classical heating are presented in Table 1.

Anti-inflammatory activity: Carrageenan-induced rat paw edema assay

All the title compounds were screened for their in vivo anti-inflammatory activity using the carrageenan-induced rat paw edema model and exhibited protection against carrageenan-induced edema (Table 2; Fig. 1a, b). The protection ranged up to 72 %, while the reference drug (indomethacin) showed 77 % at an equivalent dose. Among all the tested compounds, N-(benzo[d]thiazol-2yl)-2-(substituted)acetamides **10–13** showed remarkable anti-inflammatory activities compared to other derivatives. In particular, compounds **12** and **13** with 4-fluoroaniline and 4-methoxyaniline substituent showed better activity compare to analogs **10** and **11** with triazole and imidazole



(B) 1 Volume of paw edema at 4h 0.9 Paw edema (ml) 0.8 0.7 0.6 0.5 0.4 0.3 0.1 Indomethatin Control 3 5 6 1 ,0 1 2 3 Treatment

Fig. 1 Treated groups versus paw edema (ml). a after 2 h, b after 4 h

moieties. It indicates that the presence of benzene ring attached with thiazole moiety along with the presence of NH group attached to the aromatic ring is essential for showing the enhanced activity. Further it has been observed that a fluoro group at a strategic position on the heterocyclic ring enhances the activity of a molecule due to its enhanced lipophilicity (William, 2008) as it can be evident from the most active compound **12** (with 4-fluoroaniline group). The SAR of the title compounds was drawn based on docking studies and their binding mode analyses.

Molecular modeling studies

The final compounds **3**–7 and **10–13** were evaluated in silico (docking) to recognize their hypothetical binding mode using the X-ray crystal structure of COX-2 (PDB ID: 1CX2) and also to rationalize their structure activity relationships. To investigate the ability of molecular docking to reproduce an experimentally observed ligand-binding mode, the cocrystallized ligand **SC-558** (a selective COX-2 inhibitor) has been used as reference ligand (Fig. 2a). It was docked back into its binding site (Fig. 2a) of the crystal structure of the COX-2 using GLIDE docking program (Schrodinger Inc.). The top docked conformations (poses) closely resembled the cocrystallized conformation with a root-mean-square deviation (RMSD) 0.50–0.90 of nonhydrogen atomic positions of the ligand (**SC-558**).

The experimental binding mode of **SC-558** was reproduced and as shown in the Fig. 2a; the trifluoromethyl group (–CF₃) formed strong hydrogen bonding interactions with the exocyclic NH₂ group of Arg120 at a distance of 2.087 Å. The same group (–CF₃) was also found to be located in the vicinity of Leu531, Val116, Val349, Leu359, and Tyr355. The Val523 permits the location of the phenylsufonamide group of the ligand (**SC-558**) in a cavity formed by Phe518, Leu352, Val523, the backbone of Ser353, and Tyr355. Further sulphonamido group interacted with His90, Arg513, and Gln192 by forming a hydrogen bond which is the key point of interaction for COX-2 selectivity. In particular, sulfonamide -NH2 was found to form hydrogen bonding with the nitrogen atom of imidazole ring of His90 and oxygen atom of Ser353 at a distance of 2.505 and 3.19485 Å, respectively, whereas sulfonamide oxygen (=O) formed hydrogen bond with the NH of Arg513. Pyrazole ring of SC-558 is found to be located within the vicinity of Tyr355 and Val349. Sulphonyl group of the ligand was located around the residues of Gln192, Ser353, Phe518, Arg513, and Val523. Bromophenyl moiety was surrounded by Gly526, Val523, and Ala527. These residues were identified as crucial and involved in the binding which perhaps contributing to the selectivity of the ligand.

Earlier from the docking studies of the indomethacin, a nonselective COX-1/2 inhibitor, it has been observed that the carboxylate ion formed a salt bridge with the Arg120 in COX-2 isoenzyme which gives a more generalized anchoring point for all the classical NSAIDs, thus, limiting their selectivity due to curtailed freedom of movement of the ligand (Balakumar *et al.*, 2010). The present docking study also revealed the similar observation (Fig. 2b). The acetyl CH₂ of indomethacin formed a hydrogen bond with the phenolic OH group of Tyr355 at a distance of 3.256 Å, whereas the chlorophenyl group showed significant π - π stacking interaction with the phenolir of Phe518.

Interestingly, all the thiazole as well as benzothiazole derivatives **3–7** and **10–13** docked well into the binding pocket of COX-2 (Fig. 2c–h). The NH group and fluoro group at 4th position of the 4-fluoroaniline moiety of the most active compound **12** were found to form moderate H-bond with the phenolic OH group of Tyr355 (3.108 Å) and exocyclic NH₂ group of Arg120 (3.810 Å), respectively (Fig. 2d, e), as compared to **SC-558**. Similarly, the



Fig. 2 a Superimposition of redocked **SC-558** (*cyan* color) with its original position (*yellow* color) as cocrystal in the binding site of the crystal structure of COX-2 (PDB ID: 1CX2) isoenzyme (rmsd-0.50–0.90) showing H-bond interactions with the amino acid residues His90, Arg120, Tyr355, and Arg513, respectively. **b** Binding orientation of indomethacin within the binding site of COX-2 showing salt bridge formation between the carboxylate ion and Arg120. **c** Superimposition of compounds **3–7, 10–13**, and **SC-558** original position (yellow color) and orientation in the binding site of the crystal structure of COX-2. **d** Superimposition of compound **12** and **SC-558** with their orientations in the binding site of the crystal structure of COX-2. **e** Hypothetical binding

motif of compound **12** showing H-bonding interaction between F atom and N–H of fluoroaniline moiety with N–H of Arg120 and OH of Tyr355 at a distance 3.810 and 3.108 Å, respectively. **f** Hypothetical binding orientation of compound **13** showing H-bonding interaction between OH and N–H of *p*-anisidine moiety with N–H of Arg120 and OH of Tyr355 at a distance 3.846 and 2.288 Å, respectively. **g** Binding orientation of compound **10** in the binding site of the crystal structure of COX-2. **h** Superimposition of compounds **3**, **7**, and **SC-558** original position (*yellow* color) and orientation in the binding site of the crystal structure of COX-2 (Color figure online)

compound 13 showed significant H-bond between the H-atom of N-H group and O-atom at 4th position of the 4-methoxyaniline group with the phenolic OH group of Tyr355 and exocyclic NH₂ group of Arg120 at a distance 2.288 and 3.846 Å, respectively (Fig. 2f). The thiazole and benzothiazole moieties of all the target compounds were found to be aligned similar to the bromophenyl ring of SC558, and the benzene ring of the benzothiazole moiety showed significant π - π stacking interaction with Phe518 (Fig. 2c-g) similar to indomethacin which probably account for their better anti-inflammatory efficacy compared to other analogs 3-7 of the series. Whereas in case of compounds 10 and 11, the N-atom of triazole and imidazole moiety formed significant H-bond with the exocyclic NH₂ group of Arg120 at a distance 2.245 and 1.959 Å, respectively, (Fig. 2g) but there was no H-bond formation with Tyr355 due to lack of aromatic NH group which probably causes decrease in their activities compared to other benzothiazole derivatives.

The thiazole derivatives (4-7, except compound 3), on the other hand, were found to have different orientations compared to benzothiazole derivatives where the aromatic ring attached to amino moiety was found to be aligned with phenylsulfonamido group of SC558. Hence they were unable to form any hydrogen bond with Arg120, but showed significant hydrogen bond formation with other important amino acid residues such as His90 and Arg513 (Fig. 2h). Moreover, these derivatives could not form any π - π stacking interaction with Phe518 which might be the reason for their overall decrease in activity as compared to benzothiazole derivatives 10-13. Further, none of the compounds showed any kind of salt bridge formation like nonselective indomethacin. Hence it can be deduced that in silico docking, studies can enable us to get an insight and further guide us to rationally design novel and potent antiinflammatory agents in future.

Prediction of ADME properties

The expected ADME properties of the tested compounds were evaluated with QikProp module of Schrodinger (Table 3). The selected properties are known to influence metabolism, cell permeation, and bioavailability. Almost all the predicted properties of the tested compounds were in the ranges as predicted by QikProp for 95 % of known oral drugs and also satisfy the Lipinski's rule of five to be considered as druglike potential.

In the present study, a series of new 2-substituted-N-(1,3-

thiazole-2-yl)acetamide 3-7 and N-(benzo[d]thiazol-2-yl)-

Conclusion

2-(substituted)acetamide 10-13 derivatives have been synthesized and evaluated in vivo (rat paw edema) for their antiinflammatory activities and in silico (docking studies) to recognize the hypothetical binding motif of the title compounds with the cyclooxygenase isoenzyme (COX-2) employing GLIDE software (Schrodinger Inc.). The microwave-assisted synthesis of these compounds was found be economic, less time consuming with better yield as compared to classical heating conditions. Among all the tested compounds, N-(benzo[d]thiazol-2-yl)-2-(substituted)acetamides 10-13 showed remarkable anti-inflammatory activity compared to other derivatives (around 84-93 % of the standard: indomethacin). In particular, compounds 12 and 13 with 4-fluoroaniline and 4-methoxyaniline substituent showed better activity compared to analogs 10 and 11 with triazole and imidazole moieties. The binding mode of the title compounds has been proposed based on the docking studies. The predicted ADME properties of all the tested compounds were in the ranges as predicted by QikProp for 95 % of known oral drugs and also satisfy the Lipinski's rule of five which signifies a good absorption and hence, good bioavailability so that the observed differences in bioactivity of these compounds may be attributable to the differences in their chemical structures. Further, evaluation of these active compounds for their in vitro COX-2 binding studies is under progress.

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