

Design, Synthesis and Biological Evaluation of 4-(Imidazolylmethyl)-2-(4-methylsulfonyl phenyl)-Quinoline Derivatives as Selective COX-2 Inhibitors and *In-vitro* Anti-breast Cancer Agents

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Abstract

A new group of 4-(Imidazolylmethyl)quinoline derivatives possessing a methylsulfonyl COX-2 pharmacophore at the *para* position of the C-2 phenyl ring were designed and synthesized as selective COX-2 inhibitors and *in-vitro* anti breast cancer agents.

In-vitro COX-1 and COX-2 inhibition studies showed that all the compounds were potent and selective inhibitors of the COX-2 isozyme with IC₅₀ values in the potent range 0.063-0.090 μM, and COX-2 selectivity indexes in the 179.9 to 547.6 range. Molecular modeling studies indicated that the methylsulfonyl substituent can be inserted into the secondary pocket of COX-2 active site for interactions with Arg⁵¹³. Cytotoxicity of quinolines 9a-e against human breast cancer MCF-7 and T47D cell lines were also evaluated. All the compounds 9a-e were more cytotoxic against MCF-7 cells in comparison with those of T47D which express aromatase mRNA less than MCF-7 cells. The data showed that the increase of lipophilic properties of substituents on the C-7 and C-8 quinoline ring increased their cytotoxicity on MCF-7 cells and COX-2 inhibitory activity. Among the quinolines 9a-e, 4-((1*H*-Imidazol-1-yl)methyl) 7,8,9,10-tetrahydro-2-(4-methylsulfonylphenyl)-benzo[h]quinoline (9d) was identified as the most potent and selective COX-2 inhibitor as well as the most cytotoxic agent against MCF-7 cells.

Keywords: Quinolines; COX-2 inhibitory; Aromatase inhibitory; Breast cancer.

Introduction

Breast cancer is one of the leading causes of cancer-related mortality among women worldwide (1). In most of cases, it is hormone-dependent because tumor progression is dependent on high levels of circulating estrogens, which help the cancer cells to proliferate. Moreover, in postmenopausal

women, biologically active estrogens are locally produced from circulating inactive steroids in an intracrine mechanism in breast cancer tissues and confer estrogenic activities to carcinoma cells (2) A series of enzymes are involved in this intratumoral or in situ production of estrogens in breast carcinoma tissues, aromatase, a member of the cytochrome P450 family, is the key enzyme in this process, promoting the aromatization of the A ring of androgen precursors (3) the other enzyme involved in breast cancer are COX-2. In addition to the role of COX-2 in

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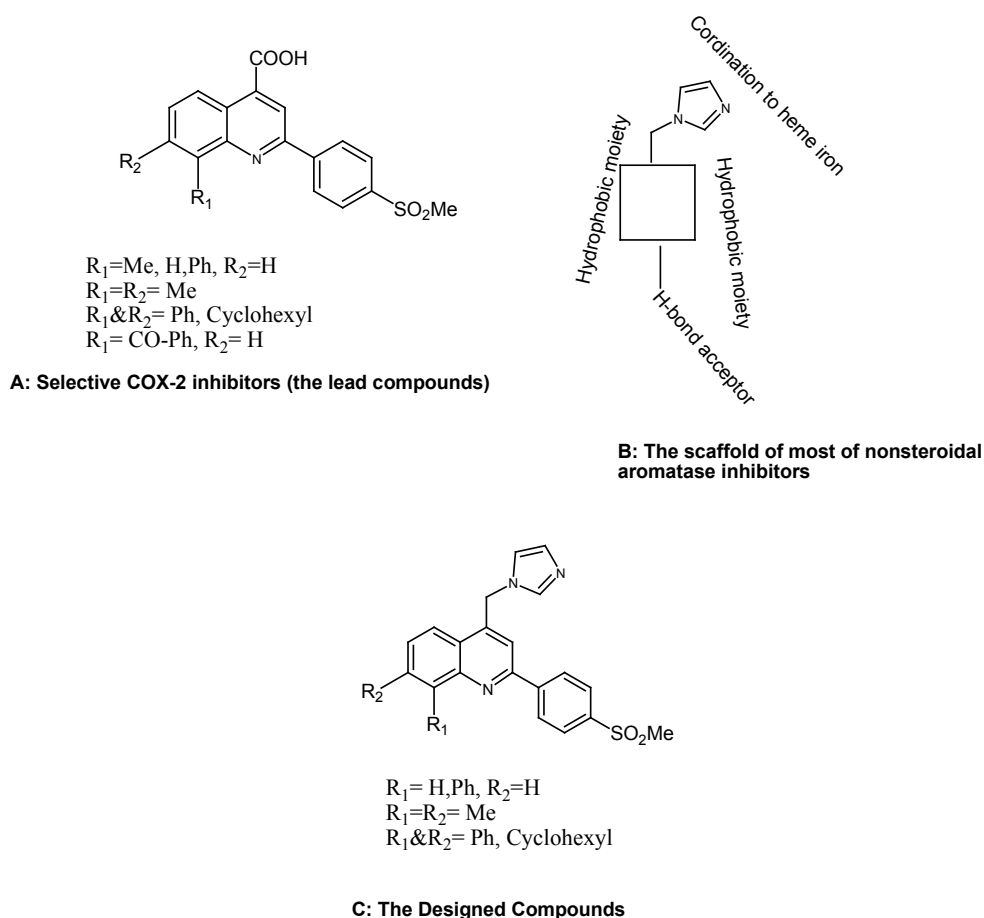


Figure 1. Chemical structures of our lead compounds (A and B) and our designed compounds (C).

inflammatory disorders such as rheumatoid arthritis and osteoarthritis, it is also implicated in cancer and angiogenesis. In this regard, several epidemiologic studies have been reported that inhibitors of COX-2 enzyme reduce the risk of colorectal, breast, and lung cancer, and COX-2 is expressed in these cancers (4-6). So we decided to design quinoline derivatives as dual inhibitors of COX-2 and aromatase to find new approaches for the prevention and treatment of breast cancer.

We recently reported several investigations describing the design, synthesis, and a molecular modeling study for a group of 2-phenyl-4-carboxyl quinolones (Figure 1, A) possessing a methyl sulfonyl COX-2 pharmacophore at the *para* position C-2 phenyl ring (7, 8). Our results showed that quinoline ring is a suitable scaffold for COX-2 inhibitory activity. On the other hand, the structure of nonsteroidal aromatase inhibitors can be considered to consist of two parts. One part is theazole part with a nitrogen

atom which interacts with the heme iron atom of the cytochrome P450 of aromatase. The second part is the bulky aryl part, which mimics the steroid ring of the substrate (androstendione) (Figure 1, B) (8, 9).

So we changed the carboxyl group of our selective COX-2 inhibitors (possessing a methyl sulfonyl COX-2 pharmacophore at the *para* position C-2 phenyl ring) with the imidazole ring, the main pharmacophore for anti-aromatase activity (9) and as shown in Figure 1, our designed compounds (C) are possessing bulky aryl part, which are important for inhibiting both aromatase and COX-2 enzymes.

Experimental

Chemistry

All chemicals and solvents used in this study were purchased from Merck AG and Aldrich Chemical. Melting points were determined

with a Thomas-Hoover capillary apparatus. Infrared spectra were acquired using a Perkin Elmer Model 1420 spectrometer. A Bruker FT-500 MHz instrument (Bruker Biosciences, USA) was used to acquire ¹H NMR spectra with TMS as internal standard. Chloroform-D and DMSO-d₆ were used as solvents. 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 an 6410Agilent LCMS triple quadrupole mass spectrometer(LCMS) with an electrospray ionization (ESI) interface. Microanalyses, determined for C and H, were within ± 0.4% of theoretical values.

General procedure for preparation of 2-(4-(methylthio)phenyl)-7,8-substituted-quinoline-4-carboxylic acid (4a-4e)

These compounds were synthesized according to our pervious methods(7, 11).

General procedure for preparation of 2-(4-methylthio-phenyl)-7,8-substituted-quinoline-4-yl)methanol (7a-7e)

LiAlH₄ (0.45 g, 12 mmol) was suspended in dry THF (20 ml) under a nitrogen atmosphere. Under vigorous stirring a solution of appropriate acid(4a-4e) (5.67 mmol) in dry THF was added dropwise to keep the reaction mixture slightly boiling. After stirring for 2 h at room temperature, the suspension was carefully hydrolyzed with NaOH solution (10%) till no more hydrogen was produced. The solid was filtered off and washed thoroughly with chloroform(12). The filtrate was dried over Na₂SO₄ and the solvent was removed under reduced pressure to yield a yellow oil which was crystallized from ether/hexane (25:75 v/v) (yield: 62-79 %).

2-(4-Methylthio-phenyl)-8-phenylquinoline-4-yl)methanol (7a)

Yield: 77%; yellow crystalline powder; mp=131-133 °C; IR (KBr): ν (cm⁻¹), 3745-2371(OH); LCMS (ESI): 380.6 (M+23)⁺100.

2-(4-Methylthio-phenyl)-7,8-dimethylquinoline-4-yl)methanol (7b)

Yield: 62%; pale yellow crystalline powder;

mp=140-141 °C; IR (KBr): ν (cm⁻¹), 3406(OH); LCMS (ESI): 310.2 (M+1)⁺100.

2-(4-Methylthio-phenyl)-benzo[h]quinoline-4-yl)methanol (7c)

Yield: 68%; yellow crystalline powder; mp=111-113 °C; IR (KBr): ν (cm⁻¹), 3290(OH); LCMS (ESI): 332.9 (M+1)⁺100.

7,8,9,10-Tetrahydro-2-(4-methylthio-phenyl)-benzo[h]quinoline-4-yl)methanol (7d)

Yield: 79%; pale yellow crystalline powder; mp=139-140°C; IR (KBr): ν (cm⁻¹), 3234(OH); LCMS (ESI): 336.9 (M+1)⁺100.

2-(4-Methylthio)phenyl-quinoline-4-yl)methanol (7e)

Yield: 63%; yellow crystalline powder; mp=120-122 °C; IR (KBr): ν (cm⁻¹), 3224(OH); LCMS (ESI): 282.1 (M+1)⁺100.

General procedure for preparation of 2-(4-methylsulfonyl-phenyl)-7,8-substituted-quinoline-4-yl)methanol (8a-8e)

One gram of 2-(4-methylthio)phenyl-7,8-substituted-quinoline-4-yl)methanol (7a-7e) was dissolved in 10 ml THF and 5 g oxone in THF/water was added. The mixture was stirred at room temperature for 3-5 h, after evaporation of THF, the residue was extracted with chloroform and dried with sodium sulfate and then evaporated(13), the product was recrystallized in chloroform/hexane (yields: 67-70%).

2-(4-Methylsulfonyl-phenyl)-8-phenylquinoline-4-yl)methanol (8a)

Yield: 48%; yellow crystalline powder; mp=178-179 °C; IR (KBr): ν (cm⁻¹), 3442(OH) 1312, 1157(SO₂); LCMS (ESI): 390.8 (M+1)⁺100.

2-(4-Methylsulfonyl-phenyl)-7,8-dimethylquinoline-4-yl)methanol (8b)

Yield 42%; orange crystalline powder; mp=148-150 °C; IR (KBr): ν (cm⁻¹), 3241(OH), 1303, 1149(SO₂); LCMS (ESI): 342.1(M+1)⁺100.

2-(4-Methylsulfonyl-phenyl)-benzo[h]quinoline-4-yl)methanol (8c)

Yield: 65%; yellow crystalline powder;

mp=210-211 °C; IR (KBr): ν (cm⁻¹), 3485 (OH), 1296, 1145(SO₂); ¹HNMR (DMSO-d₆): δ (ppm) 3.28(s, 3H, SO₂Me), 5.18 (s, 2H, CH₂), 5.71 (s, 1H, OH), 7.74-7.80 (m, 2H, benzoquinoline H₈&H₉), 7.96-7.80(m, 2H,benzoquinoline H₆&H₅), 7.99 (d, 1H,benzoquinoline H₇) 8.11 (d, 2H, 4-methylsulfonylphenyl H₂&H₆, J=8.46Hz), 8.37 (s, 1H,benzoquinoline H₃), 8.63 (d, 2H, 4-methylsulfonylphenyl H₃&H₅, J=8.46Hz), 9.4 (d, 1H, quinoline H₁₀, J=7.34Hz); LCMS(ESI): 364.1 (M+1)⁺100.

7,8,9,10-Tetrahydro-2-(4-methylsulfonylphenyl)-benzo[h]quinoline-4-yl)methanol (8d)

Yield: 57%; yellow crystalline powder; mp=162-163°C; IR (KBr): ν (cm⁻¹), 3365 (OH), 1309, 1159(SO₂);LCMS (ESI): 368.9 (M+1)⁺100.

2-(4-Methylsulfonyl)phenyl-quinoline-4-yl)methanol (8e)

Yield: 75%; yellow crystalline powder; mp=164-166°C IR (KBr): ν (cm⁻¹), 3340(OH), 1301, 1151(SO₂);LCMS (ESI): 314.9 (M+1)⁺100.

General procedure for preparation of 4-((1H-imidazol-1-yl)methyl)-2-(4-methylsulfonyl-phenyl)-7,8-substituted-quinoline

To a solution of the corresponding alcohol (1.59 mmol) in 15 ml NMP (N-methyl-2-pyrrolidone), CDI (carbonyl 1, 1-diimidazole) (1.29 g, 7.9 mmol) was added. Then the solution was heated to reflux for 20 h at 170°C. After cooling to ambient temperature, it was diluted with water (50 ml) and extracted with ethyl acetate. The combined organic phases was washed with brine and water, dried over Na₂SO₄ and evaporated under reduced pressure (14). Then the desired product was purified by flash chromatography on silica gel (dichloromethane/methanol, 90:10 v/v), (yield: 13-62%).

4-((1H-Imidazol-1-yl)methyl)-2-(4-methylsulfonyl phenyl)-8-phenyl-quinoline (9a)

Yield: 25%; cream crystalline powder; mp=196-198 °C;IR (KBr): ν (cm⁻¹), 1310, 1150 (SO₂);¹HNMR (CDCl₃): δ (ppm) 3.03 (s, 3H, SO₂Me) 5.73 (s, 2H, CH₂), 7.04 (s, 1H, imidazole H₅), 7.23(s, 1H, imidazole H₂), 7.34(s, 1H,

imidazole H₄), 7.45 (t, 1H, phenyl H₄, J=7.15Hz), 7.52 (t, 2H, phenyl H₃&H₅, J=7.31Hz), 7.69-7.76(m, 4H, quinoline H₆&H₇&phenyl H₂&H₆), 7.86 (d, 1H, quinoline H₅, J=6.88 Hz), 7.92 (s, 1H, quinoline H₃), 7.95 (d, 2H, 4-methoxysulfonylphenyl H₂&H₆, J=8.13Hz), 8.14 (d, 2H, 4-methoxysulfonylphenyl H₃&H₅, J=8.13Hz); LCMS(ESI): 440.6 (M+1)⁺100.

4-((1H-Imidazol-1-yl)methyl)-2-(4-methylsulfonyl phenyl)-7,8-dimethyl-quinoline (9b)

Yield: 61%; yellow crystalline powder; mp=218-219 °C; IR (KBr): ν (cm⁻¹), 1301, 1150 (SO₂); ¹HNMR (CDCl₃): δ (ppm) 2.58(s, 3H, Me), 2.91(s, 3H, Me), 3.12(s, 3H, SO₂Me), 5.76 (s, 2H, CH₂), 7.05 (s, 1H, imidazole H₅), 7.26(s, 1H, imidazole H₄), 7.39(s, 1H, imidazole H₂), 7.50 (d, 1H, quinoline H₆, J=8.53Hz), 7.73 (d, 1H, quinoline H₅, J=8.53Hz), 8.01 (s, 1H, quinoline H₃), 8.08 (d, 2H, 4-methylsulfonylphenyl H₂&H₆, J=8.53Hz), 8.26 (d, 2H, 4-methylsulfonylphenyl H₃&H₅, J=8.53Hz); LCMS(ESI): 392.1 (M+1)⁺100.

4-((1H-Imidazol-1-yl)methyl)-2-(4-methylsulfonyl phenyl)-benzo[h]-quinoline (9c)

Yield: 25%; cream crystalline powder; mp=216-217 °C; IR (KBr): ν (cm⁻¹) 1303, 1148 (SO₂);¹HNMR (DMSO-d₆): δ (ppm) 3.26(s, 3H, SO₂Me), 5.90 (s, 2H, CH₂), 7.01 (s, 1H, imidazole H₅), 7.36(s, 1H, imidazole H₂), 7.79 (m, 2H, benzoquinoline H₈&H₉), 7.93(s, 1H, imidazole H₄), 8.03-8.05(m, 3H,benzoquinoline H₃&H₆&H₇), 8.10 (d, 2H, 4-methylsulfonylphenyl H₂&H₆, J=8.32Hz), 8.14 (d, 1H,benzoquinoline H₁₀), 8.50 (d, 2H, 4-methylsulfonylphenyl H₃&H₅, J=8.32Hz), 9.4 (d, 1H, quinoline H₅); LCMS(ESI): 414.6 (M+1)⁺100.

4-((1H-Imidazol-1-yl)methyl)-7,8,9,10-tetrahydro-2-(4-methylsulfonylphenyl)-benzo[h]-quinoline (9d)

Yield: 35%; cream crystalline powder; mp=247-248 °C; IR (KBr): ν (cm⁻¹) 1318, 1163(SO₂);¹HNMR (CDCl₃): δ (ppm) 1.9-2.01(m, 4H, CH₂), 2.96 (m, 2H, CH₂), 3.08 (s, 3H, SO₂Me), 3.46 (m, 2H, CH₂), 5.67 (s, 2H, CH₂), 7.0 (s, 1H, imidazole H₅), 7.2(s, 1H, imidazole H₂), 7.29 (s, 1H, imidazole

H₄), 7.35 (d, 1H, quinoline H₆, J=8.57 Hz), 7.66 (d, 1H, quinoline H₅, J=8.58 Hz), 7.70 (s, 1H, quinoline H₃), 8.03 (d, 2H, 4-methoxysulfonylphenyl H₂&H₆, J=8.46Hz), 8.29 (d, 2H, 4-methoxysulfonylphenyl H₃&H₅, J=8.46Hz); LCMS(ESI): 418.7 (M+1)⁺100.

4-((1H-Imidazol-1-yl)methyl)-2-(4-methylsulfonyl phenyl)-quinoline (9e)

Yield: 15%; cream crystalline powder; mp=189-190 °C; IR (KBr): ν (cm⁻¹) 1300, 1150(SO₂); ¹H NMR (CDCl₃): δ (ppm) 3.1(s, 3H, SO₂Me), 5.76 (s, 2H, CH₂), 7.07 (s, 1H, imidazole H₃), 7.26(s, 1H, imidazole H₄), 7.29(s, 1H, imidazole H₂), 7.69-7.72 (m, 2H, quinoline H₆&H₃), 7.87 (t, 1H, quinoline H₇, J=7.27Hz), 7.98 (d, 1H, quinoline H₅, J=8.28Hz), 8.09 (d, 2H, 4-methylsulfonylphenyl H₂&H₆, J=8.37Hz), 8.26 (d, 2H, 4-methylsulfonylphenyl H₃&H₅, J=8.37Hz), 8.3 (d, 1H, quinoline H₈, J=8.40Hz); LCMS(ESI): 364.8 (M+1)⁺100.

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 1000 iterations reaching a convergence of 0.01 kcal/mol. 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. 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 (15, 16).

In-vitro cyclooxygenase (COX) inhibition assays

The assay was performed using an enzyme

chemiluminescent kit (Cayman Chemical, MI, USA) according to our previously reported method (17). The Cayman chemical chemiluminescent COX

(ovine) inhibitor screening assay utilizes the heme-catalyzed hydroperoxidase activity of ovine cyclooxygenases to generate luminescence in the presence of a cyclic naphthalene hydrazide and the substrate arachidonic acid. Arachidonate-induced luminescence was shown to be an index of real-time catalytic activity and demonstrated the turnover inactivation of the enzyme. Inhibition

of COX activity, measured by luminescence, by a variety of selective and non-selective inhibitors showed potencies similar to those observed with other *in-vitro* and whole cell methods.

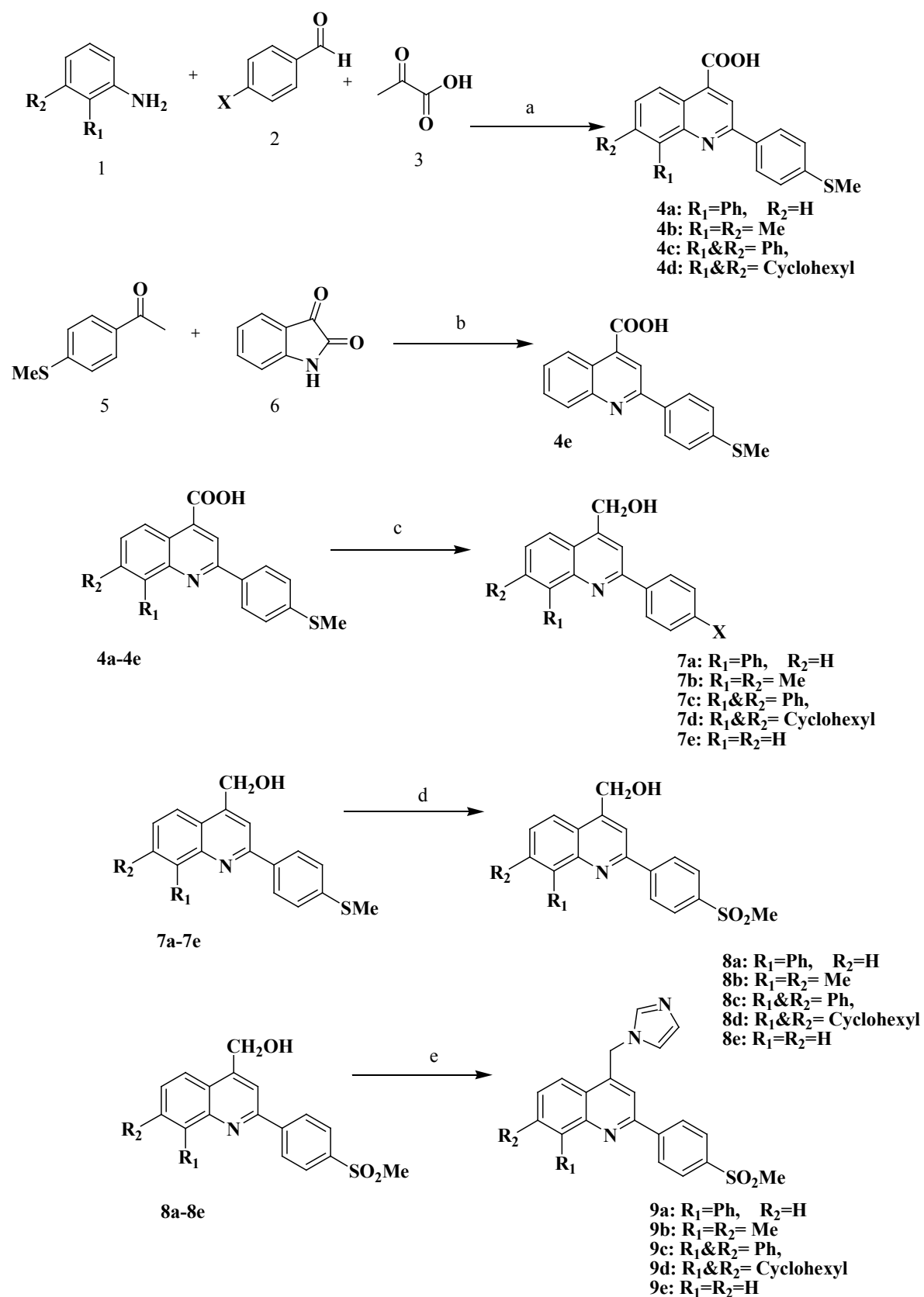
Cytotoxicity assay

Cell line and culture conditions

The human breast cancer T47D and MCF-7 cell lines were obtained from Pasteur Institute Cell Bank of IRAN (Tehran, IRAN). Cells were maintained in RPMI-1640 (Gibco, UK) culture medium supplemented with 10% fetal bovine serum (Gibco, UK) and 100 U MI⁻¹ of penicillin and 100 ng MI⁻¹ of streptomycin (Gibco, UK) at 37 °C in 5% CO₂ incubator. All reagent and chemicals used in this experiment were of cell culture or molecular biology grade, purchased from different international sources.

General procedure

The MTT (3-[4, 5-dimethylthiazol-2-yl]-2,5-diphenyl tetrazolium bromide) based assay was performed by seeding 5000 cells (T47D and MCF-7) per 180 μ L RPMI complete culture medium in each well of 96-well culture plates. The day after seeding, culture medium was changed with medium containing standard anti-tumor drug Doxorubicin as well as different concentrations of newly synthesized compounds and RPMI control (no drug). Cells were then incubated at 37 °C in 5% CO₂ incubator for 48h and 72h. Then 25 μ L of MTT solution (4mg MI⁻¹) were added to each well and further incubated at 37 °C for 3h. At the end of incubation, formazan crystals were dissolved in 100 μ L of DMSO and plates were read in a plate reader



Scheme 1. Reagents and conditions: (a) ethanol, reflux, 1-5 h (b) ethanol/KOH, reflux, 48 h (c) LiAlH₄/THF, 2 h (d) oxone/THF, 2-5 h (e) CDI/NMP, 170°C, 20 h.

Table 1. *In-vitro* COX-1 and COX-2 enzyme inhibition assay data for 4-(Imidazolylmethyl)-2-(4- methylsulfonyl phenyl)-Quinolinederivatives(9a-9e).

Compound	R ₁	R ₂	COX-1 IC ₅₀ ^a (μ M)	COX-2 IC ₅₀ ^a (μ M)	Selectivity Index ^b
9d	Cyclohexyl		34.5	0.063	547.6
9c	Phenyl		35.9	0.068	527.9
9a	Phenyl	H	30.1	0.071	423.9
9b	Me	Me	26.0	0.090	288.9
9e	H	H	12.95	0.072	179.9
Celecoxib			24.3	0.060	405

^aValues are means of two determinations acquired using an ovine COX-1/COX-2 assay kit and the deviation from the mean is < 10% of the mean value.

^b*In-vitro* COX-2 selectivity index (COX-1 IC₅₀/COX-2 IC₅₀).

(TECAN, Austria) at 540 nm. This experiment was performed in triplicate determination each time (18,19).

Result and discussion

The synthesis of target compounds was carried out according to Scheme 1. All the synthesized compounds were assayed for their COX-2 inhibitory and also cytotoxic activities against breast cancer cell lines (MCF7 and T47D). The results are presented in Table 2.

SAR data (IC₅₀ 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 COX inhibition was sensitive to the lipophilic nature of substituents. As shown in Table 1, our results showed that the increase of lipophilic properties of substituents on the C-7 and C-8 quinoline ring increased COX-2 inhibitory potency and selectivity. The relative COX-2 potency, and COX-2 selectivity profiles for the 4-imidazolylmethylquinoline derivatives, with respect to the C-7 and C-8 substituents was 9d > 9c > 9a > 9b > 9e. However, among the 4-imidazolylmethylquinoline derivatives, compound 9d possessing an unsaturated cyclohexyl ring attached to C-7 and C-8 quinoline ring exhibited highest COX-2 inhibitory potency and selectivity (COX-2 IC₅₀ = 0.063 μ M; SI = 547.6) that was as potent as the reference drug celecoxib and more selective COX-2 inhibitor than celecoxib (COX-2 IC₅₀ = 0.060 μ M; SI = 405).

SAR data (IC₅₀ values) also showed that the COX inhibition was sensitive to the nature of

substituents on the C-4 quinoline ring. All of the 4-imidazolylmethylquinoline derivatives were less potent but more selective COX-2 inhibitors than their corresponding 4-carboxyl derivatives. Our molecular modeling studies showed that the carboxyl group can interact with Arg¹²⁰ in COX-2, so replacement of carboxyl group with imidazolylmethyl may decrease COX-2 inhibitory activity. In addition, carboxyl group can also interact with Arg¹²⁰ as a key amino acid in COX-1 enzyme, so 4-imidazolylmethylquinoline derivatives have less affinity to bind to COX-1 than the 4-carboxyl derivatives and as a consequence are more selective COX-2 inhibitors.

The binding interactions of the three most potent and selective COX-2 inhibitor compound (9a, 9c and 9d) within the COX-2 binding site were investigated. They all were docked well in the COX-2 binding site. The most stable enzyme-ligand complex of (9a, 9c and 9d) which possessing a MeSO₂ COX-2 pharmacophore at *para* position of C-2 phenyl ring within the COX-2 binding site (Figure 2) shows that the *p*-MeSO₂-phenyl moiety is oriented towards the COX-2 secondary pocket (Val⁵²³, Phe⁵¹⁸ and Arg⁵¹³). These observations together with experimental results provide a good explanation for design of potent and selective COX-2 inhibitors possessing 4-((1*H*-imidazol-1-yl)methyl)-2-(4-methylsulfonylphenyl)quinoline framework.

The cytotoxicity of quinolines 9a-e against human breast cancer MCF-7 and T47D cells by MTT assay after 2 and 3 days of exposure was also evaluated. After some initial evaluations

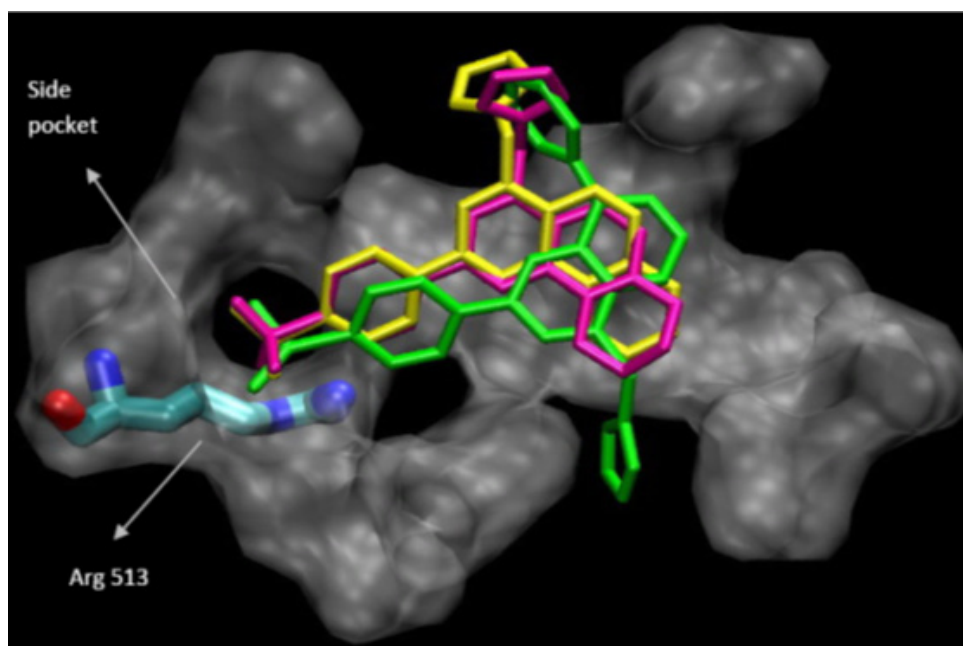


Figure 2. Docking 9a (in green), 9c (in yellow) and 9d (in pink) in the active site of murine COX-2.

concentrations of 10 and 25 μM of quinolines were used for evaluation and comparison of cytotoxicity of these compounds with doxorubicin at concentration of 250 nM against MCF-7 cells and concentrations of 25 and 50 μM of quinolines were used for evaluation and comparison of cytotoxicity of these compounds with doxorubicin at concentration of 100 nM against T47D cells. Results for each compound as the percentage of growth of the treated cells in comparison to untreated cells are shown in Table 1. The most active compound against MCF-7 cells was 9d, which exhibited 4% of survival (after 2 days exposure) ($\text{IC}_{50} < 5\mu\text{M}$). The IC_{50} of 9a was less than 10 μM and the IC_{50} of 9c was

10 μM but the IC_{50} of 9b and 9e were more than 10 μM which were the least potent compounds. The most potent compound against T47D cells was 9a which exhibited 33.8% of survival (after 2 days exposure) ($\text{IC}_{50} \leq 25\mu\text{M}$). Compounds 9c and 9e were the least potent compounds with $\text{IC}_{50} = 50\mu\text{M}$. The calculated IC_{50} values of all tested compounds after two days exposure showed that the order of the cytotoxicity against MCF-7 cells were $9d > 9a > 9c > 9b = 9e$ and in T47D cells were $9a > 9d > 9e > 9b = 9c$. These data showed that the increase of lipophilic properties of substituents on the C-7 and C-8 quinoline ring increased their cytotoxicity on MCF-7. This may be due to their ability to penetrate the cell membrane or their

Table 2. *In-vitro* cytotoxicity of quinolines (9a-9e).

Compound	R ¹	R ²	IC ₅₀ (μM)	Survival ^a (%) (10 μM)		IC ₅₀ (μM)	Survival ^b (%) (25 μM)	
			MCF-7	48h	72h	T47D	48h	72h
9d	cyclohexyl		<5	4	0.7	25	49.1	26.1
9a	Ph	H	<10	38.3	26.7	<25	33.8	25.4
9c		Phenyl	10	48.4	41.3	50	75.8	68.9
9b	Me	Me	25	61	47.2	50	66.2	69.5
9e	H	H	>25	67.2	58.2	25	47.4	47.5
Doxorubicin	-	-	0.25	50.2	31.7	0.1	53.2	27.1

^aPercent survival of MCF-7 cells following exposure to 10 μM concentration of compounds was determined after 48 h and 72 h exposure using MTT assay.

^bPercent survival of T47D cells following exposure to 25 μM concentration of compounds was determined after 48 h and 72 h exposure using MTT assay.

ability to inhibit or suppress some factors and enzymes such as COX-2. As our result showed that the order of the cytotoxicity of quinolines 9a-9e against MCF-7 cells were the same as that of their COX-2 inhibitory so one of their cytotoxic mechanism of these compounds may be through their COX-2 inhibitory activity. In addition, these quinolines 9a-9e were more cytotoxic against MCF-7 cells in comparison with those of T47D. So this may be due to their different mechanism of action in these two breast cancer cells such as more anti-aromatase activity in MCF-7 cells in comparison with T47D cells which express aromatase mRNA less than MCF-7 cells.

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