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Design, synthesis, biological evaluation, and molecular docking of

chalcone derivatives as anti-inflammatory agents

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Abstract: In this study, two series of 35 new chalcone derivatives containing aryl-piperazine or aryl-sulfonyl-piperazine fragment were synthesized and their structures were characterized by ¹H, ¹³C and ESI-MS. The *in vivo* and *in vitro* anti-inflammatory activities of target compounds were evaluated by using classical para-xylene-induced mice ear-swelling model and ELISA assays. Furthermore, docking studies were performed in COX-2 (4PH9). The *in vivo* anti-inflammatory assays indicated that most of the target compounds showed significant anti-inflammatory activities. Docking results revealed that the anti-inflammatory activities of compounds correlated with their docking results. Especially, compound **60** exhibited the most potent anti-inflammatory activity *in vivo* with the lowest docking score of -17.4 Kcal/mol and could significantly inhibit the release of LPS-induced IL-6 and TNF- α in a dose-dependent manner *in vitro*.

Keywords: inflammation, chalcone, derivatives, anti-inflammatory activity, COX-2

Inflammation is a complex biological process which seriously threatens human health. Exaggerated and prolonged inflammation may cause various diseases, such as arthritis, sepsis, and even cancer.¹ At present, the most widely used drugs in treating

inflammation are non-steroidal anti-inflammatory drugs (NSAIDs), which account for 35% of the global market for prescription of pain medications.²⁻³ Common NSAIDs, such as aspirin and indomethacin can inhibit both cyclooxygenase-1 (COX-1) and cyclooxygenase-2 (COX-2)⁴⁻⁵, displaying significant toxicity to gastrointestinal tract and kidney.⁶ For that reason, it is of great importance and urgent need to develop new anti-inflammation drugs.⁷

Chalcone which belongs to a flavonoid family is widely distributed in plants such as vegetables, fruits, tea and spices. It has been demonstrated that chalcone posses many important bioactivities including anti-oxidant⁸, anti-cancer⁹, anti-bacterial¹⁰, anti-fungal¹¹, and anti-inflammation¹². Recently, chalcone and its derivatives have been reported to show anti-inflammatory activities in acute lung injury¹³⁻¹⁴ and could protect liver from hepatic injury by inhibition of hepatic inflammation and fibrosis.¹⁵ Based on its favorable anti-inflammatory activity, chalcone framework has been used for chemical modification to find novel derivatives with better pharmacological or pharmacokinetic profiles.¹⁶

N-aryl piperazine moieties belong to an important class of organic compounds which is widely used in medicinal chemistry.¹⁷ Recently, *N-aryl* piperazine derivatives have been attracting considerable attentions for their versatile properties in pharmacology¹⁸⁻¹⁹ and anti-inflammatory activities in *vitro*²⁰ and in *vivo*²¹. Based on these data, we introduced *N-aryl* piperazine moiety into the chalcone skeleton to synthesize series I of chalcone derivatives. It is well recognized that selective COX-2 inhibitors (the coxibs) exhibit moderate gastrointestinal toxicity comparing with traditional NSAIDs.²² In view of the structures of coxibs, we discover that the coxibs bear an arylsulfonyl moiety (Fig. 1), implying that compounds bearing arylsulfonyl moiety might show selective COX-2 inhibition activity. Thus, to make COX-2 selective compounds, series II was synthesized with an arylsulfonyl moiety insertion based on the structures of compounds from series I. The anti-inflammatory activities of these two series were screened *in vivo* and *in vitro*. Molecular modeling with COX-2 was also studied.



Figure 1. Structures of selective COX-2 inhibitors (coxibs).

The synthesis of target compounds is outlined in Scheme 1. Condensation of 4-fluoro acetophenone (1) with substituted benzaldehyde 2a-2c in the presence of sodium hydroxide solution in enthanol at room temperature afforded the substituted chalcone 3a-3c.²³ For series I containing aryl-piperazine fragment, intermediates 3a-3c reacted with substituted aryl-piperazine in the presence of DMF solution at 120 °C for 12-16 h to give compounds 4a-4s under nitrogen protection. For series II containing aryl-sulfonyl-piperazine fragment, treatment of 3a-3c with piperazine by heating in the presence of potassium carbonate under nitrogen protection provided the intermediate products 5a-5c. Especially, the reaction time of this step must be controlled under 5 to 6 h otherwise unprecedented complicated products obtained. Then intermediates 5a-5c reacted with arenlsulfonyl to give compounds 6a-6p as reported.²⁴ The structures and characterization of target compounds were depicted in Table 1 and supplementary information, respectively.



Scheme 1. Synthetic routes of chalcone derivatives. Reagents and conditions: (i) 10% NaOH, EtOH, rt; (ii) aryl piperazine derivatives, K₂CO₃, DMF, N₂, 12~16 h; (iii) piperazine, K₂CO₃, 120 °C, N₂, 4 h; (IV) (**6a~6n**): arenelsulfonyl chloride, K₂CO₃, acetone, rt, 24h; (**6o~6p**): arenelsulfonyl chloride, K₂CO₃, DMF, 50 °C, 12 h.

Comp	R ₁	R_2	Yields	Comp	R_1	R ₂	Yields
4 a	-	3,4-Cl	58%	4s	4'-Cl	3-OCH ₃	57%
4b	-	4-NO ₂	56%	6a	4'-Cl	4-C1	78%
4c	-	2-F	53%	6b	4'-Cl	4-CF ₃	77%
4d	-	1-Benzyl	56%	6c	-	4-CH ₃	81%
4 e	-	2,4-CH ₃	63%	6d		4-Cl	68%
4f	-	3-C1	51%	6e	-	3-(CH=CH)-4	71%
4g	-	2,3-Cl	58%	6f	-	4-Br	79%
4h	-	3-OCH ₃	52%	6g	-	4-OCH ₃	80%
4 i	-	4-CH ₃	55%	6h	-	2,4,6-CH ₃	81%
4j	-	2-Cl	51%	6i	-	-	75%
4k	-	3-CF ₃	57%	6j	-	4-OCF ₃	76%
41	-	-	60%	6k	-	2,4,6-isopropyl	78%
4m	3',4'-OCH ₃	3,4-Cl	57%	61	-	4-Butyl	69%
4n	3',4'-OCH ₃	2,4-CH ₃	61%	6m	-	4-CF ₃	66%
40	4'-Cl	2,4-CH ₃	56%	6n	-	3,4-OCH ₃	73%
4 p	4'-Cl	3-C1	54%	60	3',4'-OCH ₃	4-Br	76%
4q	4'-Cl	4-OCH ₃	58%	6р	3',4'-OCH ₃	4-OCH ₃	72%
4r	4'-Cl	2-OCH ₃	57%				

Table 1. Structures of synthesized chalcone derivatives.

The *in vivo* anti-inflammatory activities of target compounds were evaluated using *in vivo* para-xylene-induced mice ear-swelling model. All the target compounds were administered at a dose of 40 mg/kg and aspirin (100 mg/kg) and celecoxib (5 mg/kg) were used as reference drugs. The results showed that for series I, only compounds **4b**,

4m, **4n**, and **4q** exhibited significant anti-inflammatory activities (p < 0.05). While for series II, almost all the derivatives displayed significant anti-inflammatory activities except compound **61** (p < 0.05) (Table 2). Compound **60** displayed the most potent anti-inflammatory activity which was better than that of aspirin and even equal to that of celecoxib. Preliminary structure activity relationship (SAR) analysis indicated that: 1) In view of the structures of compounds 4i with 6c and 4l with 6i, we found compounds 6c and 6i contained an extra sulforyl moiety over compounds 4i and 4i, respectively. The *in vivo* anti-inflammatory activities of these compounds were 6c >>4i, 6i >> 4l. These results implied that sulforyl group could improve compounds anti-inflammatory activity. 2) Comparing compounds 4b and 4i, the in vivo anti-inflammatory activity was 4b >> 4i. Comparing compounds 6d, 6f, 6g, 6j, 6m and 6c, 6l, the *in vivo* anti-inflammatory activities were 6d, 6f, 6g, 6j, 6m, 6m > 6c, 6l. These results implied that substitution at 4 position of aryl-piperazine or aryl-sulfonyl-piperazine with electron-withdrawing group showed advantage over electron-donating group in improving compound's anti-inflammatory activity. 3) Comparing compounds 4f and 4p, 4h and 4s, the *in vivo* anti-inflammatory activities were 4f > 4p, 4h > 4s. These results indicated that substitution at 4' position of benzene ring could not benefit from improving compound's anti-inflammatory activity.

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	Compound	Dose	Swelling	Inhibition	Docking	p value	
		(mg/kg)	degree (mg)	(%)	Score		
					(Kcal/mol)		
	control		7.0 ± 1.6				
	celecoxib	5	1.6 ± 0.7	77.03	-14.53	p < 0.001	
	aspirin	100	2.5 ± 0.8	64.91	-9.47	p < 0.001	
	chalcone	40	5.5 ± 0.7	21.68	-9.25	p = 0.07	
	4 a	40	6.1 ± 1.5	13.41	-11.12	p = 0.36	
	4 b	40	3.1 ± 1.3	55.35	-13.22	p < 0.001	
	4 c	40	7.0 ± 3.4	0.57	-10.57	p = 0.98	

Table 2. Anti-inflammatory acitivites of chalcone derivatives

4d	40	5.8 ± 1.0	17.69	-10.37	p = 0.15
4 e	40	6.0 ± 0.7	14.98	-11.27	p = 0.19
4 f	40	6.0 ± 1.4	14.12	-11.18	p = 0.29
4 g	40	4.9 ± 2.4	30.67	-11.12	p = 0.10
4h	40	5.9 ± 0.8	15.98	-10.75	p = 0.17
4 i	40	6.8 ± 1.3	2.85	-10.81	p = 0.82
4j	40	6.7 ± 2.8	4.71	-10.20	p = 0.80
4k	40	5.3 ± 3.1	24.96	-12.03	p = 0.21
41	40	6.8 ± 2.0	3.71	-10.31	p = 0.81
4 m	40	2.6 ± 0.8	63.48	-14.99	p < 0.001
4n	40	3.6 ± 1.9	48.50	-13.33	p = 0.008
40	40	5.2 ± 2.6	25.68	-12.35	p = 0.17
4p	40	6.6 ± 2.8	5.85	-10.90	p = 0.77
4 q	40	3.2 ± 1.5	55.06	-13.44	p < 0.001
4 r	40	6.6 ± 1.2	6.13	-11.31	p = 0.61
4 s	40	6.9 ± 2.2	2.28	-12.08	p = 0.88
ба	40	3.0 ± 1.4	56.63	-13.04	p < 0.001
6b	40	2.8 ± 1.0	59.63	-13.56	p < 0.001
6с	40	4.9 ± 0.9	30.81	-11.50	p = 0.01
6d	40	3.7 ± 2.1	46.79	-11.33	p = 0.005
бе	40	4.6 ± 0.9	34.95	-12.52	p = 0.007
6f	40	3.7 ± 0.9	47.36	-12.64	p = 0.02
бg	40	3.8 ± 0.9	45.22	-12.57	p = 0.02
6h	40	3.1 ± 1.1	55.49	-13.01	p < 0.001
6i	40	4.9 ± 0.7	29.96	-11.20	p = 0.01
бј	40	3.7 ± 2.6	47.36	-12.86	p = 0.02
6k	40	2.8 ± 0.7	60.06	-13.92	p < 0.001
61	40	5.3 ± 3.0	23.97	-12.68	p = 0.25
6m	40	3.7 ± 0.8	47.93	-12.98	p < 0.001

6n	40	3.1 ± 1.1	55.49	-12.99	p < 0.001
60	40	1.7 ± 0.8	75.46	-17.40	p < 0.001
6р	40	2.5 ± 1.0	64.05	-15.98	p < 0.001

The statistical significance was calculated between each compound group and control group by using student's t-test of unpaired data. P values < 0.05 were considered statistically significant.

The structure and favorable *in vivo* anti-inflammatory activities of compounds led us to study the molecular docking with COX-2 enzyme. The results indicated that the anti-inflammatory activities of compounds might correlate with their docking results with COX-2 (Table 2). Compound **60**, **6p**, **4m**, and **6k** which had a lower docking score than others displayed better anti-inflammatory activities than other compounds. Especially, compound **60** had the lowest docking score of -17.4 Kcal/mol and the best anti-inflammatory activity (inhibition rate, 75.46%) which showed two hydrogen bonds (H-bonds) and two chemical bonds of arene-cation with COX-2 (Fig.2 and Fig. 3). H-bonds were formed between oxygen atom of methoxyl group with amino acid Tyr386 and Ser531 at distances of 2.46 and 2.78 Å, respectively. The intensities of H-bonds between compound **60** with COX-2 were 98% and 67%, respectively. Chemical bonds of arene-cation were formed between benzene ring of compound **60** and amino acid Arg121 of COX-2. To validate the results of molecular docking, further enzyme studies on COX-2 are needed.



Figure 2. 2D docking model of compound 60 with COX-2.



Figure 3. 3D docking model of compound 60 with COX-2.

Considering the favorable *in vivo* anti-inflammatory activity and docking results in silico, we chose compound **60** for further studies. To reveal the cytotoxity of compound **60**, MTT assay was conducted in mouse RAW264.7 macrophages. The results indicated that compound **60** showed no significant cytotoxity with a concentration less than 50 μ M (Fig. 4a). To further detect the effects of compound **60** on pro-inflammatory factors (TNF- α and IL-6) secretion, RAW264.7 macrophages were pretreated with different doses (10, 20, 40 μ M) of compound **60** for 2 h and then exposed to LPS (1 μ g/ml) for additional 22 h. The levels of TNF- α and IL-6 in the media were determined. The results displayed that compound **60** significantly inhibited TNF- α and IL-6 release by RAW264.7 macrophages in a dose-dependent manner (Fig. 4b).



Figure 4. Cytotoxicity and *in vitro* anti-inflammatory activity of compound **60**. (a) MTT assays of compound **60** in mouse RAW264.7 macrophages. (b) Compound **60** inhibited LPS-induced TNF- α and IL-6 release by RAW264.7 macrophages in a dose-dependent manner in mouse RAW264.7 macrophages.

In summary, two series of 35 new chalcone derivatives containing aryl-piperazine or aryl-sulfonyl-piperazine fragment were synthesized. The *in vivo* anti-inflammatory results revealed that most of the target compounds exhibited potent anti-inflammatory activity. Especially, compound **60** exhibited the most potent anti-inflammatory activity *in vivo* and could significantly inhibit the release of LPS-induced IL-6 and TNF- α by RAW264.7 macrophages in a dose-dependent manner *in vitro*. Furthermore, docking results indicated that the anti-inflammatory activities of compounds correlated with their docking results. Our finding might provide information on developing potentially new and safe anti-inflammatory agents.

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