



Tetrahydroquinoline derivatives as CRTH2 antagonists

Jiwen Liu^{*}, Yingcai Wang, Ying Sun, Derek Marshall, Shichang Miao, George Tonn, Penny Anders, Joel Tocker, H. Lucy Tang, Julio Medina

Amgen Inc., 1120 Veterans Boulevard, South San Francisco, CA 94080, USA

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ABSTRACT

A series of tetrahydroquinoline-derived inhibitors of the CRTH2 receptor was discovered by a high throughput screen. Optimization of these compounds for potency and pharmacokinetic properties led to the discovery of potent and orally bioavailable CRTH2 antagonists.

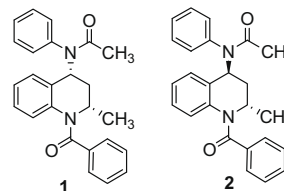
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CRTH2 (chemoattractant receptor-homologous molecule expressed on Th2 cells), also known as DP₂, is a G-protein coupled receptor related to the *N*-formyl peptide receptor (FPR) subfamily of chemoattractant receptors. Its endogenous ligand is prostaglandin D₂ (PGD₂). PGD₂ is the major cyclooxygenase product formed and secreted by activated mast cells during allergic reactions.^{1–3} PGD₂ also signals through prostanoid D (DP or DP₁) receptor. The DP receptor is primarily expressed on airway epithelium, smooth muscle and platelets, while CRTH2 is selectively expressed on Th2 cells, T cytotoxic type 2 (Tc2) cells, eosinophils, and basophils.^{4–6} Stimulation of CRTH2 by PGD₂ mediates multiple inflammatory responses, such as chemotaxis of eosinophils, basophils and Th2 cells, eosinophil activation and degranulation, cytokine production from Th2 T cells, and leukotriene production by mast cells.^{7–13} Therefore, blockade of CRTH2 is likely to be beneficial in the treatment of allergic diseases triggered by PGD₂.

Several research groups, including ours, discovered that tetrahydroquinoline derivatives are potent CRTH2 antagonists.^{14–19} These compounds were of special interest to us, because to our knowledge, it was the only series of CRTH2 antagonists devoid of a carboxylic acid moiety. Here we report the discovery, optimization and structure activity relationship (SAR) of the tetrahydroquinoline derivatives.

Tetrahydroquinoline **1** (Table 1), discovered in a high throughput screen, inhibited the binding of ³H-PGD₂ to hCRTH2 receptors on 293 cells with an IC₅₀ of 0.043 μM (Table 1).²⁰ Compound **1** also inhibited CRTH2 mediated cell migration in response to PGD₂ with an EC₅₀ of 11 nM using hCRTH2 stably transfected CEM cells.²¹ The

Table 1



Compd	Chiral Center	CRTH2 IC ₅₀ ^a in buffer (μM)	CRTH2 IC ₅₀ ^a in plasma (μM)
1	Racemic	0.043	1.05
2	Racemic	>10	
1a ^b	(2 <i>S</i> ,4 <i>R</i>)	0.017	0.44
1b ^b	(2 <i>R</i> ,4 <i>S</i>)	0.42	>10

^a Displacement of ³H-PGD₂ from the CRTH2 receptor expressed on 293 cells. Assay run in buffer containing 0.5% BSA or in 50% plasma. See Ref. 20 for assay protocol. Values are means of three experiments, standard deviation is ±30%.

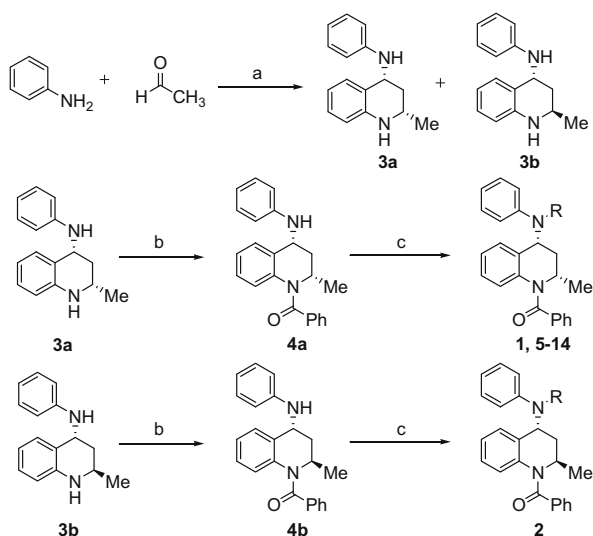
^b ee >99%.

^{*} Corresponding author.

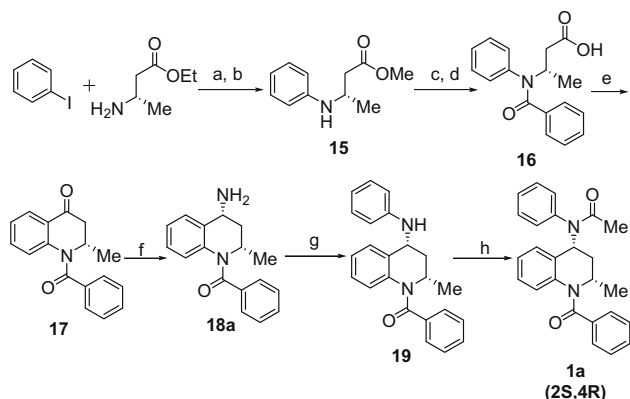
E-mail address: jiwenl@amgen.com (J. Liu).

structure of compound **1** was determined to be *cis* by synthesis²² (Scheme 1) and NMR studies.²³ Conversely, the *trans* isomer (**2**) had weak CRTH2 activity. Furthermore, the stereo-selective synthesis (Scheme 2) indicated the (2*S*,4*R*) enantiomer **1a** was responsible for the majority of the CRTH2 activity of racemic compound **1** (Table 1).

Compound **1** was synthesized according to [Scheme 1](#). Reaction of aniline with acetaldehyde in ethanol at room temperature afforded a mixture of *cis*-*trans* isomers **3a** and **3b** in >95% yield. Separation of the *cis* isomers **3a** was achieved in 35% yield by recrystallization from 10% EtOAc/Hex. The *trans* isomer **3b** was obtained in 30% yield from the purification of the mother liquor using silica column chromatography. Selective acylation of **3a** and **3b** with benzoyl chloride at 1-*N* position afforded amides **4a** and **4b**, respectively, in 90% yield. Reaction of **4a** and **4b** with acetyl chloride afforded **1** and **2**, respectively, in 85% yield. Compounds **5–14** ([Table 2](#)) were synthesized from *cis* intermediate **4a** using reductive amination, sulfonylation or acylation.

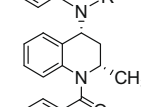


Scheme 1. Racemic synthesis of **1**, **2** and **5–14**. Reagents and conditions: (a) EtOH, rt, 24 h, 35% for compound **6** after recrystallization in 10% EtOAc/hexanes; (b) PhCOCl, triethylamine, DCM, rt, 24 h, 90%; (c) for amines: aldehydes, Na(OAc)₃BH, ClCH₂CH₂Cl, rt, 20 h, ~80%; for sulfonamides: sulfonyl chloride, DMAP, pyridine, rt, 3 h, 60%; for amides, acid chlorides, NaH, THF, rt, 20 h, 85%.



Scheme 2. Stereo-selective synthesis of **1a**. Reagents and conditions: (a) CuI, potassium carbonate, DMF, water, 90 °C, 48 h, 70%; (b) SOCl₂, MeOH, rt, 12 h, 90%; (c) PhCOCl, triethylamine, DCM, rt, 24 h, 90%; (d) LiOH, THF/MeOH/water, rt 4 h, 95%; (e) oxalyl chloride, DMF, DCM, 0 °C–rt, 3 h, then AlCl₃, DCM, 0 °C–rt, 12 h, 60%; (f) ammonium acetate, sodium cyanoborohydride, MeOH, 70 °C, 2 days, 85%; (g) phenyl boronic acid, pyridine, DMF, copper(II) acetate, air, 60 °C, overnight, 20%; (h) acetyl bromide, NaH, THF, 0 °C–rt, 3 h, 85%.

Table 2



Compd ^a	R	CRTH2 IC ₅₀ ^b in buffer (μM)
1	–COMe	0.043
4a	–H	>50
5	–CH ₂ CH ₃	3.17
6	–SO ₂ Ph	5.50
7	–COPh	0.105
8	–CO(CH ₂) ₃ CH ₃	0.064
9	–CO(CH ₂) ₂ CO ₂ H	0.005
10	–CO(CH ₂) ₂ CO ₂ NH ₂	0.029
11	–COCH ₂ CO ₂ H	0.54
12	–CO(CH ₂) ₃ CO ₂ H	0.022
13	–COCH=CHCO ₂ H	2.28
14	–CO(1,3-Ph)CO ₂ H	40.3

^a Mixture of racemic mixture of (2*S*,4*R*) and (2*R*,4*S*) enantiomers.

^b Displacement of ³H-labeled PGD₂ from the CRTH2 receptor expressed on 293 cells. See Ref. 20 for assay protocol. Values are means of three experiments, standard deviation is ±30%.

The stereo-selective synthesis (Scheme 2) of **1a** began with a CuI catalyzed coupling of iodobenzene with the (*S*)- β -amino acid ester.²⁴ Amide formation of the coupling product (**15**) with benzoyl chloride followed by ester hydrolysis yielded acid **16**. Conversion of the carboxylic acid to the acid chloride followed by an intramolecular Friedel-Crafts acylation provided ketone **17**.²⁵ Reductive amination with ammonium hydroxide produced primary amine **18a**, which was coupled with phenyl boronic acid in the presence of copper acetate to give compound **19**.¹⁴ Reaction of **19** with acetyl bromide afforded compound **1a** ((2*S*,4*R*)-enantiomer) with >99% ee. Compound **1b** ((2*R*,4*S*)-enantiomer) was obtained from chiral HPLC separation of racemate **1**.²⁶

Compounds **22**, **24** and **25** (Table 3) were synthesized according to Scheme 3. Reductive amination of ethyl acetoacetate with aniline afforded ester **20**. Saponification of **20** followed by intramolecular Friedel–Crafts acylation afforded ketone **21**.²⁷ Amide formation with benzoyl chloride followed by a reductive amination yielded compound **18**. Finally, compound **22** was obtained from acetamide formation of **18** with acetyl bromide. Compounds **24**

Table 3

Chemical structure of the CRTH2 antagonist scaffold, showing a 1-methyl-2-phenyl-1,2,3,4-tetrahydroquinoline core with a substituent R at the 3-position and a benzoyl group at the 4-position.

Compd ^a	R	CRTH2 IC ₅₀ ^b in buffer (μM)
1	Ph	0.043
22	H	3.62
24	Et	0.25
25	Bn	0.43

^a Mixture of racemic mixture of (2*S*,4*R*) and (2*R*,4*S*) enantiomers.

^b Displacement of ³H-labeled PGD₂ from the CRTH2 receptor expressed on 293 cells. See Ref. 20 for assay protocol. Values are means of three experiments, standard deviation is ±30%.

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