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Identification of potent, selective and orally bioavailable phenyl ((R)-3-phenylpyrrolidin-3-yl)sulfone analogues as ROR γ t inverse agonists

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The retinoic acid-related orphan receptor (ROR) family of nuclear receptors comprises three members: $ROR\alpha$, $ROR\beta$ and ROR γ (ROR γ t being a splice variant of ROR γ).¹ The three RORs share a high degree of sequence similarity, but exhibit distinct tissue distribution patterns and distinct functional roles in the regulation of many physiological processes including development, immunity, circadian rhythm and cellular metabolism.² RORyt is the key transcription factor that drives differentiation of naïve CD4⁺ T helper cells to Th17 cells, and induces the transcription of IL-17A and IL-17F.³ There is abundant evidence that the IL-17/Th17 pathway plays an important role in the pathogenesis of psoriasis.⁴ Biologics known to inhibit the IL-17/Th17 pathway are clinically validated for the treatment of psoriasis.5 Targeting RORyt also provides a novel opportunity to treat other autoimmune diseases such as inflammatory bowel disease (IBD), rheumatoid arthritis (RA) and multiple sclerosis (MS).6 In addition, RORyt-deficient mice show significantly reduced Th17 cell populations and decreased susceptibility to experimental autoimmune encephalomyelitis (EAE) as well as intestinal and skin inflammation.^{1,7} Thus RORyt has garnered significant attention as a small molecule therapeutic target.8-12

ABSTRACT

An X-ray crystal of one of our previously discovered ROR γ t inverse agonists bound to the ROR γ t ligand binding domain revealed that the cyclohexane carboxylic acid group of compound **2** plays a significant role in ROR γ t binding, forming four hydrogen bonding and ionic interactions with ROR γ t. SAR studies centered around the cyclohexane carboxylic acid group led to identification of several structurally diverse and more potent compounds, including new carboxylic acid analogues **7** and **20**, and cyclic sulfone analogues **34** and **37**. Notably, compounds **7** and **20** were found to maintain the desirable pharmacokinetic profile of **2**.

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Our group recently reported the discovery of a series of bicyclic sulfonamides as RORyt inverse agonists.¹³ A representative compound 1 (Figure 1) showed an EC₅₀ of 39 nM in the Jurkat cell-based Gal4 reporter assay for RORyt. However, it suffered from agonist activity against pregnant X receptor (PXR) in vitro (EC₅₀ = 2000 nM, Y_{max} = 100%) and cytochrome P450 induction in vivo. We then described structure-based drug design efforts that led to a novel phenyl (3-phenylpyrrolidin-3yl)sulfone series as exemplified by compound 2 (Figure 1).¹⁴ Compound 2 exhibited an EC₅₀ of 119 nM in the ROR γ t Gal4 reporter assay, no detectable activity against PXR and desirable pharmacokinetic (PK) profiles in mouse. Furthermore, compound 2 displayed dose-dependent inhibition of IL-17 production in a mouse IL-2/IL-23-induced pharmacodynamic model and biologic-like efficacy in an IL-23-induced mouse acanthosis model. This letter describes subsequent structure-activity relationship (SAR) studies to improve the potency of compound 2.



Figure 1. Previously reported RORyt inverse agonists.

Previous SAR studies¹⁴ of the phenyl (3-phenylpyrrolidin-3yl)sulfone scaffold revealed that the (R)-enantiomer of the pyrrolidine was more potent than the (S)-enantiomer; the perfluoroisopropyl and para-fluorophenyl groups were optimal for balanced potency-selectivity profiles; and the transcyclohexane stereochemistry in compound 2 provided greater potency than the cis-isomer. To gain insights on how to improve potency, an X-ray crystal structure of compound 2 with the ligand binding domain (LBD) of RORyt was solved (Figure 2).15 Compared to a previously disclosed structure of a related analogue (compound 12 in ref. 14), the phenyl (3phenylpyrrolidin-3-yl)sulfone backbone of 2 adopted a similar binding mode in RORyt. The trans-cyclohexane ring adopted a chair conformation, placing the two carbonyl groups in a 1,4-diequatorial arrangement. The terminal carboxylic acid moiety fits nicely in a binding site surrounded by multiple polar amino acid residues - forming four polar (hydrogen bonding and ionic) interactions with backbone NH groups of Gln286 and Leu287 and guanidine side chains of Arg364 and Arg367.



Figure 2. X-ray crystal structure of compound 2 and the LBD of ROR γ t (PDB ID 6P9F). The carbons of the protein are colored in pink and those of 2 in green. Sulfurs are colored in yellow, nitrogens in blue, oxygens in red and fluorines in cyan. Hydrogen bonds between compound 2 and ROR γ t LBD are indicated by a dashed line.

The carboxylic acid moiety of **2** not only engages multiple hydrogen bonding interactions with the ROR γ t LBD, but also improves metabolic stability of the series. Therefore, part of our SAR optimization efforts was to keep the carboxylic acid group intact and explore additional substitution to the cyclohexane ring in order to improve ROR γ t potency. We also envisioned that rigidification of the cyclohexane ring with fused/bridged bicyclic scaffolds, while maintaining the two carbonyl groups close to the 1,4-di-equatorial orientation, could lead to potency improvements. In parallel, we also explored structural diversity by investigating alternative functional groups to the carboxylic acid, because of the potential liability of the carboxylic acid group toward glucuronidation.

The effects of additional substitution to 1- and 4-position of the cyclohexane ring (α to the carboxylic acid and the carboxamide, respectively) are summarized in Table 1. The trans-1-methyl cyclohexane carboxylic acid 3 was perhaps more active than compound 2 (EC₅₀ of 79 nM vs. 119 nM). However, the trans-1-ethyl and 1-fluoro analogues (4 and 5) lost activity approximately 2 and 5 fold, respectively, compared to 2. Substitution at 4-position of the cyclohexane ring turned out to be more fruitful. Alkyl-substituted trans-analogues (6-8) improved potency by 2-4 fold compared to 2, with the ethyl derivative 7 being the most potent at 31 nM. The two ether compounds (9 and 10) were less potent compared to the alkyl analogues. Fluorosubstitution (11) was tolerated, whereas the two hydroxy analogues (12 and 13, cis or trans not assigned) were slightly less potent than 2. Similar to previous observation¹⁴, *cis*-cyclohexane carboxylic acids 14 and 15 showed much weaker activity compared to their trans counterparts (5 and 6, respectively). This could be explained by the fact that the *cis*-cyclohexanes in 14 and 15 cannot project the two carbonyl groups in the 1,4-di-equatorial orientation as observed in the structure of compound $2/ROR\gamma t$ (Figure 2) and further demonstrated the importance of the diequatorial conformation for binding affinity. Overall, analogues in Table 1 showed good selectivity against PXR and good liver microsome stability. Additionally, all compounds in Table 1 (as well as Tables 2 and 3) exhibited excellent selectivity against ROR α (>40 μ M), ROR β (>40 μ M), LXR α (>7.5 μ M) and LXR β (>7.5 μ M).

SAR related to the replacement of the cyclohexane ring with bicyclic scaffolds is outlined in Table 2. Three fused bicyclo[3.1.0]hexane analogues (16-18) were synthesized and were found to lose RORyt activity compared to 2. Of the two bridged analogues, the bicyclo[2.2.2]octane compound 20 showed EC₅₀ of 33 nM in the RORyt Gal4 reporter assay, while the corresponding bicyclo[2.2.1]heptane analogue 19 was about 10-fold less potent (392 nM). Also noteworthy here is that compound 20 maintained high selectivity against PXR and displayed promising liver microsome stability. Two oxabicyclo[2.2.2]octane analogues (21 and 22) were also prepared. Unfortunately, both of them lost potency by several fold compared to 20. The enhanced potency of 20 compared to 2 could be explained by the ability of bicyclo[2.2.2]octane ring, a known bioisostere to the 1,4-trans-cyclohexane,16 to place the two carbonyl groups in 180° arrangement, reminiscent of the 1,4diequatorial binding conformation of 2 (figure 2). The added steric bulk of bicyclo[2.2.2]octane could also enhance binding to RORyt. In contrast, the bicyclo[2.2.1]heptane ring in 19 and the bicyclo[3.1.0]hexanes in 16-18 cannot effectively achieve the linear arrangement for the two carbonyl groups, thus leading to reduced RORyt potency.

Our efforts to identify non-carboxylic acid alternatives are summarized in Table 3. Compounds 23-31 were derived from direct replacement of the carboxylic acid group with known bioisosteres, while keeping the cyclohexane ring intact. Compared to the acid 2, the tetrazole compound 23 showed significant loss of potency (~10 fold). Both isomers of hydantoin analogues (24 and 25) were also less potent than 2. The potency loss for both tetrazole and hydantoins could be due to the fact that the increase steric bulk of these groups compared to the carboxylic acid cannot be accommodated by the small binding pocket of RORyt LBD. The amide 26 had comparable potency while the ester 27 showed much weaker activity as anticipated. Compound 28. one of the two

Table 1. In vitro data of substituted cyclohexane carboxylic acid analogues

			$R^2 OH$ 4 1 0 F		$\frac{R^{1}}{4} \xrightarrow{\text{cis}} \frac{R^{2}}{1} \xrightarrow{\text{OH}} F$	
Cmpd #	Structure	R ¹	R ²	RORγt EC ₅₀ nM ^{a,b}	PXR EC ₅₀ nM (% max.) ^c	LM (h, m) (% remaining) ^d
2	Α	Н	Н	119 ± 28	>50000 (18)	97, 86
3	Α	Н	Me	79 ± 20	>50000 (18)	97, 93
4	Α	Н	Et	266 ± 119	>50000 (6)	100, - ^e
5	Α	Н	F	508 ± 152	>17000 (62)	100, 70
6	Α	Me	Н	56 ± 12	>50000 (7)	80, 95
7	Α	Et	Н	31 ± 24	>50000 (6)	89, 83
8	Α	<i>n</i> -Pr	Н	68 ± 37	_e	_ ^e , _ ^e
9	Α	MeO	Н	594 ± 12	>18000 (94)	82, 95
10	Α	MeOCH ₂	Н	109 ± 53	>50000 (7)	100, 88
11	Α	F	Н	74 ± 36	10600 (31)	- ^e , - ^e
12	Α	OH (isomer 1)	Н	183 ± 67	>50000 (9)	100, 99
13	Α	OH (isomer 2)	Н	269 ± 40	>50000 (19)	95, 98
14	В	Н	F	4177 ^c	>22000 (69)	82, 94
15	В	Me	Н	776 ^c	>50000 (18)	85, 100

^a Values are means of two or more experiments performed in duplicate unless otherwise noted. ^b RORyt reporter assay was performed using a Jurkat cell line. ^c Value from a single experiment performed in duplicate. ^dMetabolic stability in human and mouse liver microsome; values are percentage remaining after 10 min of incubation. ^eNot tested.

Table 2. In vitro data of bicyclic carboxylic acid analogues



^a Values are means of two or more experiments performed in duplicate unless otherwise noted. ^bRORyt reporter assay was performed using a Jurkat cell line. ^c Value from a single experiment performed in duplicate. ^d Metabolic stability in human and mouse liver microsome; values are percentage remaining after 10 min of incubation. "Not tested.

A LR

methyl sulfone isomers (*cis* or *trans* not assigned), exhibited similar potency to compound **2**. The primary alcohol **30** had respectful potency of 200 nM in Gal4 assay while the secondary alcohol **31** was notably less potent. Although several compounds showed comparable activity to **2** from this effort, no noticeable potency improvement was observed. Next set of compounds (**32**-**40**) represented analogues where the cyclohexane carboxylic acid moiety was replaced with cyclic sulfones. The five-membered cyclic sulfone isomers (**32** and **33**, absolute stereochemistry not assigned) had similar activity to **2**. It was exciting to find that the six-membered cyclic sulfone **34** improved ROR γ t potency to 33 nM. To our delight, it also displayed excellent metabolic stability and PXR selectivity. Thus substitution to the six-membered cyclic sulfone was pursued. Both methyl and ethyl analogues (**35** and **36**) lost potency by 2-3 fold compared to **34**. On the other hand, the fluoro analogue **37** maintained ROR γ t potency at 36 nM, good PXR selectivity and metabolic stability. Additional analogues with more polar hydroxy, hydroxymethyl and methoxy groups (**38-40**, respectively) resulted in reduced activity.

Table 3. In vitro data of non-carboxylic acid analogues

		$F_3C \xrightarrow{CF_3}_{D \leq S} \xrightarrow{N}_{O \leq S}_{O \leq S}$	¯}_F	R
Cmpd #	R	RORγt EC ₅₀ nM ^{a,b}	PXR EC ₅₀ nM (% max.) ^c	LM (h, m) $(\% \text{ remaining})^d$
2	€-<>>Ko	119 ± 28	>50000 (18)	97, 86
23		1260 ± 499	>50000 (11)	100, 73
24	NH H H Somer 1	1900 ^c	>14000 (28)	84, 81
25	NH NH Isomer 2	572°	>50000 (3)	93, 82
26	€NH2	197 ± 140	>50000 (5)	100, 100
27		1167 ^c	<u>_</u> e	<u>-</u> ^e , - ^e
28	€ S ⁰ Isomer 1	128 ± 56	1800 (29)	88, 69
29	€ S S S S S S S S S S S S S	336 ± 106	>50000 (6)	100, - ^e
30	€	200 ± 107	>50000 (8)	75, 63
31	}OH	749 ± 290	>50000 (13)	86, 58
32	s s=0 Isomer 1	160 ± 22	>13000 (49)	100, 78
33	S=O Isomer 2	129 ± 94	7200 (37)	100, 96
34	ş–∕_s≶o	33 ± 11	>50000 (5)	93, 96
35	ş s s o	102 ± 46	>50000 (10)	_ ^e , _ ^e
36	₹ ₹ S S O	83 ^c	5200 (32)	3, 0
37	₹ S S	36 ± 8	>50000 (16)	100, 90
38	HOS_O	123 ± 60	>50000 (6)	100, 90
39	€ CH S S O CO CO CO CO CO CO CO CO CO	216 ± 120	>50000 (8)	75, 31
40	€ S	146 ± 24	>50000 (3)	_ ^e , _ ^e

^{*a*} Values are means of two or more experiments performed in duplicate unless otherwise noted. ^{*b*} RORγt reporter assay was performed using a Jurkat cell line. ^{*c*} Value from a single experiment performed in duplicate. ^{*d*} Metabolic stability in human and mouse liver microsome; values are percentage remaining after 10 min of incubation. ^{*e*} Not tested.

Compounds 7 and 20 were selected for further evaluation. Both compounds showed half-life of >120 min in the *in vitro* human, rat and mouse liver microsome stability assay. In a Caco-2 assay, both compounds 7 and 20 exhibited good permeability

(153 and 150 nm/s, respectively) and no detectable efflux (efflux ratio of 0.7 and 0.9, respectively). Compounds 7 and 20 were also found to have no significant CYP inhibition liabilities, with IC₅₀ values greater than 10 μ M against CYP1A2, 2B6, 2C8, 2C9, 2C19, 2D6 and 3A4. Because of their similar *in vitro* profiles, both compounds 7 and 20 were taken into mouse PK studies and the results were compared to compound 2 (Table 4). At an oral dose of 10 mg/kg, compound 7 displayed C_{max} of 3.7 μ M and AUC (area under the curve) of 16.6 μ M*h; and compound 20 achieved C_{max} of 6.7 μ M and AUC of 24 μ M*h which was comparable to compound 2.

Table 4. PK profile of 7, 20 and 2 in balb/c mice^a

Cmpd #	7	20	2
C_{max} (μM)	3.7	6.7	6.6
T _{max} (h)	0.5	1.0	2
AUC (µM*h)	16.6	24	36.6

^a Oral dose: 10 mg/kg; vehicle: 5% *N*-methyl pyrrolidinone (NMP), 76% PEG 300, 19% TPGS.

Schemes 1-4 outline several representative synthesis. Scheme 1 illustrates the synthesis of compound **3**. Methylation of diester **41** gave the *cis*-isomer **42** as the only product. Epimerization of **42** with LDA deprotonation and methanol quenching followed by saponification provided the diacid **43** as a 1:1 mixture of *cis*- and *trans*-isomers. To facilitate separation of isomers, compound **43** was treated with oxalyl chloride followed by benzyl alcohol to yield the monobenzyl ester **44**. Separation of two isomers of **44** by chiral HPLC followed by hydrogenation gave the desired *trans*-diacid **45**. BOP-mediated coupling of **45** with intermediate **46**¹⁴ completed the synthesis of **3**.



Scheme 1. Synthesis of 3. Reagents and reaction conditions: (a) LDA, MeI, THF (78%); (b) LDA, THF, MeOH (83%, 1:1 mixture of *cis* and *trans* isomers); (c) LiOH, THF (94%); (d) (COCl)₂, DMF, BnOH, DCM (69%); (e) chiral OJ-H separation; (f) Pd/C, H₂, ethyl acetate (100%); (g) 46, BOP, *i*-Pr₂NEt, DMF (54%).

Scheme 2 depicts the synthesis of compound 7. *In situ* selfdimerization of 47 by treatment of triethylamine and mesyl chloride yielded 48.¹⁷ Ester-directed stereoselective hydrogenation of 48 with Crabtree's catalyst provided 49 as a single isomer. Treatment of 49 with DIBAL gave diol 50. Oxidation of 50 using freshly prepared chromic acid solution followed by mono esterification produced 51. BOP-mediated coupling of 51 with 46 followed by deprotection provided 7.

Scheme 3 shows the synthesis of compounds 16 and 17. Treatment of the acid 52 with $(Boc)_2O$ and DMAP in *tert*-butanol gave the *tert*-butyl ester 53. Cyclopropanation of 53 with ethyl diazoacetate was achieved using rhodium (II) acetate dimer as a

catalyst in dichloromethane yielded a 1:5 mixture of **54** and **55**.¹⁸ After separation of two isomers, deprotection of each compound with TFA followed by HATU-mediated coupling with **46** and saponification provided **16** and **17**.



Scheme 2. Synthesis of 7. Reagents and reaction conditions: (a) TEA, MsCl, DCM (40%); (b) H₂, Crabtree's catalyst, DCM (98%); (c) DIBAL, toluene, -78 °C (99%); (d) H₂CrO₄, acetone, water (67%); (e) (Boc)₂O, DMAP, *t*-BuOH (40%); (f) **46**, BOP, *i*-Pr₂NEt, DMF (69%); (g) TFA, DCM (83%).



Scheme 3. Synthesis of 16 and 17. Reagents and reaction conditions: (a) (Boc)₂O, DMAP, *t*-BuOH (67%); (b) ethyl diazoacetate, Rh₂(OAc)₄, DCM (50%); (c) TFA, DCM; (d) 46, HATU, *i*-Pr₂NEt, DMF; (e) LiOH, THF.

Scheme 4 describes the synthesis of compounds 21 and 22. Treatment of ethyl acrylate (58) and two equivalents of diethyl malonate (59) with sodium hydride in THF provided compound 60. Decarboxylation of 60 followed by protection of the ketone yielded compound 61. LAH reduction of 61 gave the diol 62. Conversion of the diol to *bis*-tosylate followed by ketal deprotection delivered 63. Grignard addition to the ketone 63 followed by cyclization produced 64. Conversion of the tosylate to the acetate followed by hydrolysis provided the alcohol 65. Oxidation of 65 followed by esterification gave 66. Ozonolysis of 66 followed by oxidation yielded the acid 67. HATU-mediated coupling of 67 with 46 followed by hydrogenation completed the synthesis of 21. Alternatively, esterification of 68 with 46 followed by hydrolysis completed the synthesis of 22.

The rest of the compounds exemplified in this manuscript were prepared by BOP or HATU-mediated coupling of the common intermediate **46** with the corresponding carboxylic acid side chain.

In summary, detailed SAR studies around the cyclohexane carboxylic acid moiety of compound **2** were carried out. Compounds **7**, **20**, **34** and **37** all improved ROR γ t inverse agonist potency by several fold compared to **2**. Furthermore, compounds **7** and **20** were found to essentially maintain the desirable pharmacokinetic profile of **2**. Compounds **34** and **37** represent non-carboxylic acid alternatives with improved potency. The combination of improved potency, good PK profile and structure diversity of these new side chains represents significant progress in the SAR development of the phenyl ((*R*)-3-phenylpyrrolidin-3-yl)sulfone series of ROR γ t inverse agonists. Application of these findings with new core modifications to further improve potency will be reported in the future.



Scheme 4. Synthesis of 21 and 22. Reagents and reaction conditions: (a) NaH, THF (51%); (b) NaCl, DMSO, H₂O, 160 °C (88%); (c) ethylene glycol, TsOH, toluene, reflux (85%); (d) LAH, Et₂O, 0 °C (74%); (e) TsCl, pyridine (90%); (f) HCl, THF, reflux (94%); (g) vinylmagnesium bromide, THF, -78 °C; (h) NaH, DME, reflux (55% for two steps); (i) CsOAe, DMF, 100 °C (97%); (j) K₂CO₃, MeOH, H₂O (94%); (k) PDC, DMF; (l) BnBr, K₂CO₃, DMF (72% for two steps); (m) O₃, DCM, Me₂S, -78 °C; (n) NaClO₂ 2-methylbut-2-ene, NaH₂PO₄, H₂O, *t*-BuOH (85% for two steps); (o) 46, HATU, *i*-Pr₂NEt, DMF; (p) H₂, MeOH; (q) TMSCH₂N₂, MeOH, toluene; (r) LiOH, THF.

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- 15. Attempts to obtain a structure of compound **2** with ROR γ t LBD using the same protein as reported in references 13 and 14 were unsuccessful. Therefore, a new ROR γ t LBD construct was used for the co-crystal structure of compound **2**. The new construct differs from the previous one in that the protein was covalently linked to a SRC1 peptide which was found to help co-crystallization. Details about the modified protein can be found in the deposited structure (PDB ID 6P9F).
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, iEc, bioavail RORyt EC₅₀ = 31 nM Orally bioavailable in mouse

20 ROR γ t EC₅₀ = 33 nM Orally bioavailable in mouse

F₃C

RO