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## The Discovery of Anthranilic Acid-based MMP Inhibitors. Part 3: Incorporation of Basic Amines

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**Abstract**—Anthranilic acid derivatives bearing basic amines were prepared and evaluated in vitro and in vivo as inhibitors of MMP-1, MMP-9, MMP-13, and TACE. Piperazine **4u** has been identified as a potent, selective, orally active inhibitor of MMP-9 and MMP-13. © 2001 Elsevier Science Ltd. All rights reserved.

In an effort to find novel therapies for the treatment of diseases such as atherosclerosis,<sup>1</sup> rheumatoid arthritis, osteoarthritis,<sup>2</sup> and cancer<sup>3</sup> we have recently disclosed a series of sulfonamide hydroxamic acid inhibitors of MMP-1, MMP-9, MMP-13, and TACE, based on an anthranilic acid scaffold.<sup>4</sup> It has been postulated that the aberrant control of specific MMPs and TACE is a causative factor in the etiology of these pathologies. The SAR of the anthranilic acid 3- and 5-positions (**4**, R<sup>3</sup> and R<sup>5</sup>) as well as the P1' moiety (**4**, R<sup>1</sup>), leading to compounds exemplified by **1** (Fig. 1), with nanomolar level in vitro activity, and oral bioavailability, has been discussed.<sup>5</sup>

In the course of our work on the anthranilate hydroxamates, it became clear that oral activity was greatly enhanced by the incorporation of a basic amine moiety into the inhibitor molecule. In fact, few compounds in this series lacking a basic amine displayed any activity in an in vivo bioactivity model. A similar circumstance has been reported for the sulfonamide-hydroxamates, exemplified by the oncology clinical candidate CGS-27023A.<sup>6</sup>

We now report on the synthesis and evaluation of a series of anthranilate hydroxamates, of general structure

**4**, containing basic amine groups at R<sup>2</sup> or R<sup>3</sup>. Analysis of NMR and molecular modeling studies of anthranilate hydroxamates bound to the active site of MMP-13 suggested that polar functionality at these two positions would be solvent exposed.<sup>7</sup>

### Chemistry

In general, the desired sulfonamide hydroxamic acids were prepared as previously described (Scheme 1).<sup>4,8</sup> Sulfonamide formation was followed by attachment of the desired basic moieties and the fully functionalized anthranilate sulfonamides were then converted into the hydroxamic acids. Analogues **4a–4e** were obtained after alkylation of **3a** (R<sup>1</sup>=OMe; R<sup>2</sup>=H; R<sup>3</sup>, R<sup>4</sup>=Me; R<sup>5</sup>=H) or **3b** (R<sup>1</sup>=OMe; R<sup>2</sup>=H; R<sup>3</sup>, R<sup>4</sup>=Me; R<sup>5</sup>=Br) with the appropriate picolyl or benzylic chloride using potassium carbonate in DMF. The synthesis of benzylic amine **4f** began with the alkylation of **3a** with 4-carbomethoxybenzyl bromide, followed by selective ester hydrolysis (NaOH) and reduction of the

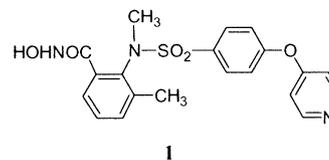
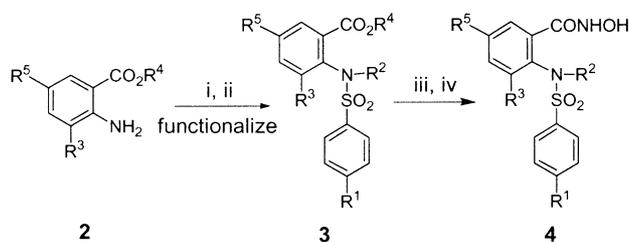


Figure 1. Sulfonated anthranilate hydroxamic acids.

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**Scheme 1.** (i) 4- $R^1$ PhSO<sub>2</sub>Cl, TEA; (ii)  $R^2$ X, NaH or K<sub>2</sub>CO<sub>3</sub>; (iii) NaOH; (iv) (COCl)<sub>2</sub>, DMF, NH<sub>2</sub>OH.

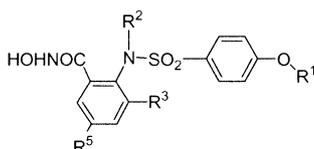
resulting acid to the alcohol with borane–THF. The alcohol was next converted into the corresponding bromide (PPh<sub>3</sub>/CBr<sub>4</sub>) and then displaced with dimethylamine. Aryl piperazine derivative **4g** resulted from alkylation of **3a** with 4-bromobenzyl bromide followed by Buchwald amination of the aryl bromide with 1-methylpiperazine. Propargylic amines **4h–4j** were obtained by alkylating **3a** or **3b** with propargyl bromide and subsequent Mannich alkylation. Variants **4k–4r** bearing amines at the anthranilate 3-position were prepared by amine displacement of the benzylic bromides resulting from NBS bromination of *N*-methyl sulfonamides **3c** ( $R^1 = \text{OMe}$ ;  $R^2 = \text{Me}$ ;  $R^3, R^4 = \text{Me}$ ;  $R^5 = \text{H}$ ) or **3d** ( $R^1 = \text{OMe}$ ;  $R^2 = \text{Me}$ ;  $R^3, R^4 = \text{Me}$ ;  $R^5 = \text{Br}$ ). The 5-aryl derivatives **4s–4t** were obtained by Suzuki couplings of the aryl boronic acids with **3e** ( $R^1 = \text{OMe}$ ;  $R^2 = \text{Me}$ ;  $R^3 = \text{CH}_2\text{N}[(\text{CH}_2)_2]_2\text{NMe}$ ;  $R^4 = \text{Me}$ ;  $R^5 = \text{Br}$ ). The biaryl ether analogues **4u** and **4v** arose from **3f** ( $R^1 = \text{OPh-4-Cl}$ ;  $R^2 = \text{Me}$ ;  $R^3, R^4 = \text{Me}$ ;  $R^5 = \text{H}$  or Br).

## Biology

All of the anthranilate hydroxamic acids were tested *in vitro*<sup>9</sup> for their ability to inhibit MMP-1, MMP-9, MMP-13, and TACE<sup>10</sup> (Table 1). Inhibitors of MMP-9 are potentially valuable as inhibitors of tumor metastasis,<sup>3</sup> while MMP-13 inhibitors may offer protection from the cartilage degradation associated with osteoarthritis.<sup>2</sup> Inhibitors of TACE are potentially valuable for the treatment of rheumatoid arthritis, Crohn's disease and other inflammatory diseases.<sup>11</sup> Selectivity over MMP-1 was sought in order to examine whether the inhibition of MMP-1 is a possible source of the musculoskeletal side effects seen in clinical trials of broad spectrum MMP inhibitors.<sup>12</sup>

The *in vitro* potencies against the MMPs and TACE for the series of anthranilate hydroxamic acid analogues in which a variety of basic amines have been linked to the sulfonamide nitrogen (**4a–4j**) are shown in Table 1. All of these compounds are potent inhibitors of MMP-9 and, with the exception of **4d**, MMP-13. They are comparable *in vitro* to CGS-27023A. The picolyl analogues, **4a–4c**, and the benzyl derivatives **4f–4g**, are essentially equipotent (3–20 nM) with similar selectivity profiles. Incorporation of a 5-bromo substituent into the gelatinase selective compound **4d** provides **4e**, a 3 nM MMP-13 inhibitor with enhanced potency against all four enzymes screened and no selectivity for MMP-9. The propargylic amines **4h–4j** are also excellent (6–19 nM) inhibitors of MMP-9 and MMP-13 with weak selectivity

**Table 1.** *In vitro* potency of substituted anthranilate hydroxamic acids



Compd	$R^1$	$R^2$	$R^3$	$R^5$	IC <sub>50</sub> , (nM)			
					MMP-1	MMP-9	MMP-13	TACE
<b>4a</b>	Me	CH <sub>2</sub> -3-Py	Me	H	143	5	8	231
<b>4b</b>	Me	CH <sub>2</sub> -3-Py	Me	H	329	7	18	356
<b>4c</b>	Me	CH <sub>2</sub> -3-Py	Me	H	391	8	18	645
<b>4d</b>	Me	CH <sub>2</sub> Ph-4-O(CH <sub>2</sub> ) <sub>2</sub> NC <sub>5</sub> H <sub>10</sub>	Me	H	176	7	56	277
<b>4e</b>	Me	CH <sub>2</sub> Ph-4-O(CH <sub>2</sub> ) <sub>2</sub> NC <sub>5</sub> H <sub>10</sub>	Me	Br	35	2	3	108
<b>4f</b>	Me	CH <sub>2</sub> Ph-4-CH <sub>2</sub> NMe <sub>2</sub>	Me	H	213	3	11	243
<b>4g</b>	Me	CH <sub>2</sub> Ph-4-N[(CH <sub>2</sub> ) <sub>2</sub> ] <sub>2</sub> NMe	Me	H	142	7	20	146
<b>4h</b>	Me	CH <sub>2</sub> CCCH <sub>2</sub> NEt <sub>2</sub>	Me	H	470	11	19	218
<b>4i</b>	Me	CH <sub>2</sub> CCCH <sub>2</sub> NEt <sub>2</sub>	Me	Br	150	14	7	36
<b>4j</b>	Me	CH <sub>2</sub> CCCH <sub>2</sub> N[(CH <sub>2</sub> ) <sub>2</sub> ] <sub>2</sub> NMe	Me	Br	129	10	6	26
<b>4k</b>	Me	Me	CH <sub>2</sub> NEt <sub>2</sub>	H	744	8	84	210
<b>4l</b>	Me	Me	CH <sub>2</sub> N[(CH <sub>2</sub> ) <sub>2</sub> ] <sub>2</sub> O	H	608	5	14	174
<b>4m</b>	Me	Me	CH <sub>2</sub> ProMe	H	517	5	6	200
<b>4n</b>	Me	Me	CH <sub>2</sub> Im	H	781	10	43	157
<b>4o</b>	Me	Me	CH <sub>2</sub> N[(CH <sub>2</sub> ) <sub>2</sub> ] <sub>2</sub> NMe	H	306	1	12	154
<b>4p</b>	Me	Me	CH <sub>2</sub> N[(CH <sub>2</sub> ) <sub>2</sub> ] <sub>2</sub> NMe	Br	194	2	5	26
<b>4q</b>	Me	Me	CH <sub>2</sub> N[(CH <sub>2</sub> ) <sub>2</sub> ] <sub>2</sub> NPh	Br	444	3	3	87
<b>4r</b>	Me	Me	CH <sub>2</sub> N[(CH <sub>2</sub> ) <sub>2</sub> ] <sub>2</sub> NBoc	Br	674	7	6	63
<b>4s</b>	Me	Me	CH <sub>2</sub> N[(CH <sub>2</sub> ) <sub>2</sub> ] <sub>2</sub> NMe	Ph-4-OCF <sub>3</sub>	745	2	4	210
<b>4t</b>	Me	Me	CH <sub>2</sub> N[(CH <sub>2</sub> ) <sub>2</sub> ] <sub>2</sub> NMe	2-Naphthyl	343	2	3	49
<b>4u</b>	Ph-4-Cl	Me	CH <sub>2</sub> N[(CH <sub>2</sub> ) <sub>2</sub> ] <sub>2</sub> NMe	H	155	1	0.8	122
<b>4v</b>	Ph-4-Cl	Me	CH <sub>2</sub> N[(CH <sub>2</sub> ) <sub>2</sub> ] <sub>2</sub> NMe	Br	82	0.5	0.7	80
CGS-27023A	—	—	—	—	15	9	8	231

over MMP-1. Thus, compound **4h**, unsubstituted at the anthranilate 5-position, is a 19 nM MMP-13 inhibitor. Its 5-bromo analogue, **4i**, is a slightly more potent (7 nM) MMP-13 inhibitor and a substantially more potent TACE inhibitor (36 nM). Excellent TACE activity is also displayed by propargylic piperazine **4j** (26 nM), which retains activity against MMP-9 and MMP-13. However, as we had seen before,<sup>5</sup> despite their potency against isolated enzyme, compounds **4i** and **4j** did not display significant TNF- $\alpha$  inhibitory activity in a THP-1 cellular assay at 3  $\mu$ M.<sup>13</sup>

The feasibility of incorporating amines at the 3-position of the anthranilic acid ring was explored with benzylic amines **4k–4v** (Table 1). The diethylamine **4k** and imidazole **4n** were the least active members of this series against MMP-1 and MMP-13, but remained potent inhibitors of MMP-9 (8–10 nM). The proline methyl ester **4m** displayed selectivity approaching 100-fold for MMP-13 over MMP-1.

The piperazines **4o–4v** all proved to be potent inhibitors of MMP-13. The initial analogue, **4o**, is an excellent MMP-9 (1 nM) and MMP-13 (12 nM) inhibitor. Addition of a 5-bromo substituent gives **4p** with good activity versus MMP-13 (5 nM) and TACE (26 nM), but reduced MMP-1 selectivity. Larger substituents on the piperazine, as in the *N*-phenyl and *N*-Boc piperazines, **4q** and **4r**, retained MMP-13 activity and provided the desired selectivity over MMP-1. Similarly, substitution of 5-aryl groups (**4s–4t**) for the 5-bromo moiety of **4p** also provided potent MMP-13 inhibitors (3–4 nM) with more than 100-fold selectivity over MMP-1. Extension of the P1' group with the biaryl ethers **4u** and **4v** boosted potency against all four enzymes relative to their shorter P1' analogues **4o** and **4p**, but the greater increase for MMP-13 made these compounds more than 100-fold selective over MMP-1.

It should be noted that at room temperature biaryl ether **4u** exists as a 1:1 mixture of atropisomers due to restricted rotation about the sulfonamide N–C bond. The racemate can be separated by chiral HPLC on a Chiralpak AD column eluting with ethanol. A 9:1 mixture of atropisomers is converted into a 3:2 mixture on heating for 0.5 h in refluxing ethanol. We have been unable to resolve these atropisomers in large quantity by either resolution or chemical synthesis. Compound **4u** was the only anthranilate hydroxamic acid that we investigated by chiral HPLC.

A single crystal X-ray structure of **4v** bound to the active site of MMP-13 was determined and is shown in Figure 2. As expected from our previous NMR structure determinations,<sup>6</sup> the anthranilate phenyl ring lies near the S2' subsite, the biaryl ether is buried in the S1' pocket and the piperazine ring is solvent exposed. The potency of **4v** versus MMP-1, despite its lengthy P1' group, is consistent with previous data from this series<sup>5</sup> and indicates that the arginine residue that forms the bottom of the MMP-1 S1' pocket is pushed aside to some degree by the P1' substituent, extending the depth of the pocket.

The in vivo bioactivity after oral dosing of some of the anthranilate-hydroxamates (**4a**, **4d**, **4g–4j**, **4l**, **4n–4q**, and **4s–4v**) against MMP-13 was initially assessed through the use of a dialysis tubing implant assay.<sup>14</sup> All of the compounds tested were compared to Novartis' sulfonamide-hydroxamate clinical lead, CGS-27023A, in the same experiment. Of these, **4g** (50 mpk), **4p** (50 mpk), **4u** (25 mpk), and **4v** (25 mpk) are 10–30% more potent than an equal dose of CGS-27023A, with **4u** and **4v** being the best.

Anthranilate-hydroxamates **4g**, **4p**, **4u**, and **4v** were also tested side-by-side with CGS-27023A in a bovine articular cartilage explant assay.<sup>15</sup> At a concentration of 1  $\mu$ M, compounds **4p** and **4v** provided a level of inhibition of cartilage degradation 60 and 20% higher than CGS-27023A, respectively. Compounds **4g** and **4u** were roughly equivalent to CGS-27023A at 1  $\mu$ M in this assay.

Hydroxamates **4p**, **4u**, and **4v** were next evaluated on oral dosing in an in vivo rat sponge-wrapped cartilage model.<sup>16</sup> All three compounds provided significant inhibition of collagen degradation in this model. However, while **4p** and **4v** at 50 mg/kg/bid were equipotent with a 50 mg/kg/bid dose of CGS-27023A, **4u** surpassed CGS-27023A. Oral dosing of **4u** at 50 mg/kg/bid provided a 73% inhibition of collagen degradation compared to a 55% inhibition by CGS-27023A at the same dose. Even a 25 mg/kg/bid dose of **4u** provided better protection from cartilage degradation than a 50 mg/kg/bid dose of CGS-27023A (38 vs 30%).

In summary, we have synthesized a series of anthranilate-hydroxamic acid MMP and TACE inhibitors bearing basic amine moieties. Many of these compounds are potent inhibitors of MMP-9 and MMP-13 in vitro. Four of these compounds were evaluated in vivo to assess their potential for treating osteoarthritis. Compounds **4g**, **4p**, **4u**, and **4v** are active in an in vitro cartilage degradation assay, display oral activity in an in

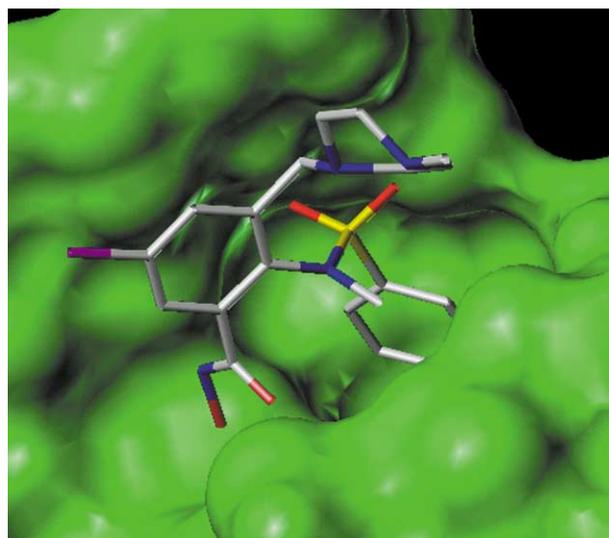


Figure 2. X-ray structure of compound **4v** in the active site of MMP-13.

vivo mouse bioactivity model and also demonstrated oral activity in a rat sponge-wrapped cartilage model. In particular, we have identified piperazine **4u** as a 0.8 nM MMP-13 inhibitor, with almost 200-fold selectivity over MMP-1, that is twice as potent as CGS-27023A in this efficacy model.

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