



RATIONAL DESIGN, SYNTHESIS, AND SERINE PROTEASE INHIBITORY ACTIVITY OF NOVEL P₁-ARGININOYL HETEROCYCLES

Susan Y. Tamura,* Brian M. Shamblin, Terence K. Brunck, and William C. Ripka

Department of Medicinal Chemistry, Corvas International, Inc., 3030 Science Park Rd., San Diego, CA 92121

Abstract: Peptidomimetic derivatives featuring a P₁-argininoyl heterocycle were designed. The preparation of two key building blocks containing benzoxazole or benzimidazole rings and their incorporation into thrombin and factor Xa specific sequences is described. The serine protease inhibitory activity of these targets was evaluated. Molecular modeling of two representative structures is presented. © 1997 Elsevier Science Ltd.

Thrombin, Factor VIIa, and Factor Xa are key members of the trypsin class of serine protease enzymes involved in the blood coagulation cascade. They play a vital role in the regulation of normal hemostasis and abnormal intravascular thrombus development.¹ Recent advances in the elucidation of the structure and function of human thrombin have led to an increased understanding of the pivotal role played by this multifunctional enzyme in the regulation of hemostatic processes and maintenance of vascular function.² Peptidomimetic inhibitors of this key enzyme are emerging as potential therapeutic agents for the prevention and treatment of thrombotic vascular disease.³

Peptidoyl heterocycles that inhibit various serine proteases such as human neutrophil elastase, prolyl endopeptidase, and thrombin are of current interest as therapeutic targets for the treatment of emphysema,^{4,5} amnesia,⁶ and thrombotic vascular disease,⁷ respectively. Prompted by the recent report of Maryanoff et al.⁷ on peptidoyl heterocycle containing thrombin inhibitors, we report herein our work in the area of antithrombotic drugs, which has led to the design of related argininoyl heterocycles.

We investigated the argininoyl heterocycles **3a–c** as a novel class of direct serine protease inhibitors (Figure 1). The new targets incorporate an arginine-based α -ketoheterocycle as the electrophilic transition-

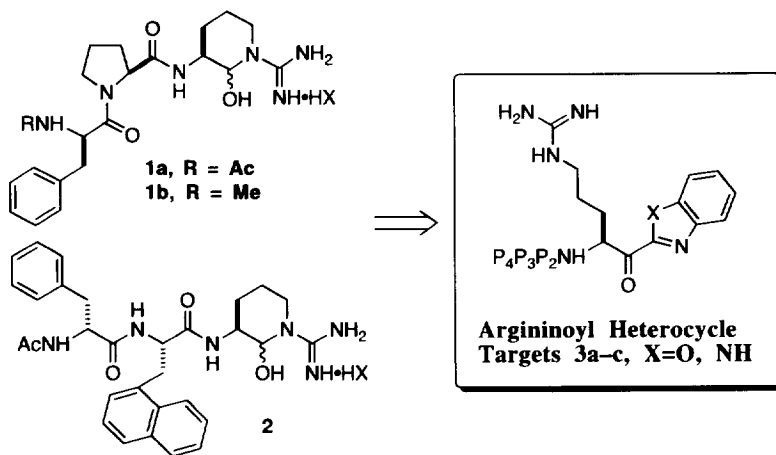
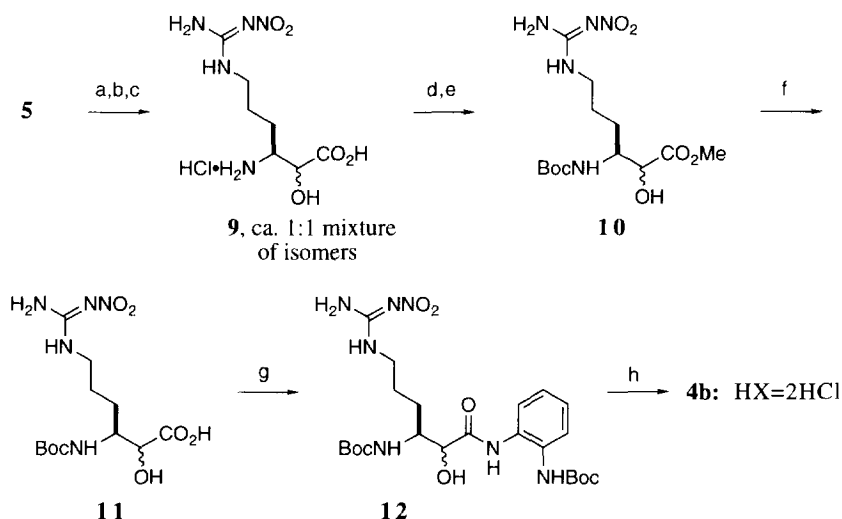


Figure 1. Structures of Prototypical Thrombin and Factor Xa Inhibitors and Argininoyl Heterocycles **3a–c**.

Scheme 1. Reagents and conditions: (a) KCN, KHCO_3 , THF, H_2O ; (b) TBDMSCl, imidazole, 67%; (c) H_2S , pyridine, Et_3N , 100%; (d) MeI; (e) *o*-aminophenol, EtOH, AcOH, reflux, 63% for 2 steps; (f) 50% TFA/ CH_2Cl_2 , 90%.

sensitive functionalities, was developed (Scheme 1). Boc nitroargininal **5**^{12,13} was converted to the *t*-butyldimethylsilyl cyanohydrin **6**, which was isolated a 1:1 mixture of stereoisomers,¹⁴ in two steps by treatment with potassium cyanide followed by TBDMS chloride (67% overall yield). Using a classical procedure,^{15a} thioamide **7** was produced in quantitative yield. Formation of the thioimide, followed by condensation with *o*-aminophenol, produced the desired heterocycle **8** in 63% overall yield.^{15b,c} Deprotection with TFA proceeded smoothly to give intermediate **4a** in 90% yield.

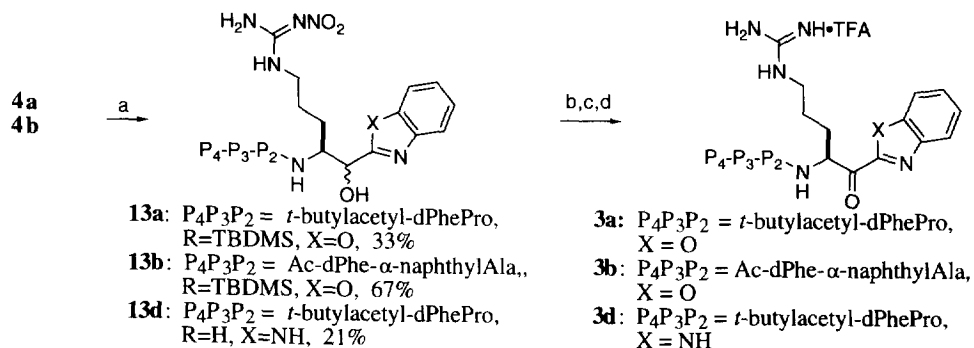
The thioimide methodology developed successfully to prepare **4a** was applied to the preparation of benzimidazole **4b**. Interestingly, the thioimide failed to react with 1,2-phenylenediamine under similar conditions. Alternatively (Scheme 2), α -hydroxy- β -amino acid **9**, which was isolated as a 1:1 mixture of stereoisomers,^{14,16} was prepared from Boc nitroargininal **5** in three steps: formation of the cyanohydrin, conversion to the imide, and acidic hydrolysis of the imide with concomitant removal of the Boc protection. Compound **10** was converted to the Boc hydroxy acid **11** by Fischer esterification, followed by reProtection of the β -amino group with Boc anhydride. Overall yield for the five step protocol was 55%. Hydrolysis of the ester efficiently afforded the Boc hydroxy acid in 95% yield. Coupling of the hydroxy acid **11** to the monoprotected 1,2-phenylenediamine¹⁷ proceeded in fair yield (41%, unoptimized) using HBTU/HOBt to give hydroxy amide **12**. Under the acidic conditions both Boc protecting groups were removed and cyclization occurred to produce the benzimidazole **4b** in 86% crude yield.



Scheme 2. Reagents and conditions: (a) KCN, KHCO₃, THF, H₂O; (b) MeOH, HCl, 0°C-rt; (c) 6 N HCl, reflux; (d) MeOH, HCl, reflux; (e) Boc₂O, NaHCO₃, THF, H₂O, 55% from **2**; (f) LiOH, aqueous MeOH; Dowex cation exchange resin, 95%; (g) mono-Boc-1,2-phenylenediamine, HBTU, HOBt, 4-methylmorpholine, 41%; (h) 4 N HCl, reflux, 86%.

The intermediates **4a** and **4b** were used in the syntheses of peptidoyl heterocycles **3a–c** (Scheme 3). Elaboration with *t*-butylacetyl-d-PhePro-OH¹⁷ or Ac-dPhe- α -naphthylAla-OH¹⁸ was carried out using HBTU/HOBt/NMM. Yields for the coupling were 33% (**13a**), 67% (**13b**), and 21% (**13c**). HF cleavage of intermediates **13a–c** went smoothly to give the hydroxyheterocycles in 50–70% yield. Oxidation of the hydroxybenzoxazole using a modified Moffatt protocol,⁴ followed by HPLC purification, gave the desired

argininoyl benzoxazoles **3a** and **3b** and argininoyl benzimidazole **3c**.



Scheme 3. Reagents and conditions: (a) *t*-BuAc-dPhePro-OH, or Ac-dPhe- α -naphthylAla-OH, HBTU, HOBT, 4-methylmorpholine; (b) HF, anisole; (c) DMSO, EDC, dichloroacetic acid; (d) reverse-phase HPLC purification.

Biological Activity

The target argininoyl heterocycles were evaluated for their ability to inhibit a range of important serine proteases, including thrombin (FIIa), factor Xa (FXa), trypsin and plasmin (Table 1).¹⁹ In general, the SAR of the benzoxazoles parallels that of the corresponding aldehydes.

Table 1. In vitro IC₅₀ values (μM) of argininoyl heterocycles **3a–3c** and reference standards **1a** and **2** against a range of important serine proteases

cmpd	$P_4P_3P_2$	P_1' moiety	FIIa	FXa	trypsin	plasmin
3a	<i>t</i> -BuAc-dPhePro	benzoxazole	0.068	0.11	0.0023	1.32
3b	Ac-dPhe- α -naphthylAla	benzoxazole	>2.5	0.029	0.0064	0.0142
3c	<i>t</i> -BuAc-dPhePro	benzimidazole	0.0921	>2.5	1.01	>2.5
<i>reference standards:</i>						
1a	Ac-dPhePro	H	0.284	>2.5	0.156	>2.5
2	Ac-dPhe- α -naphthylAla	H	>2.5	0.025	0.025	<0.025

The most intriguing comparison in the in vitro assays is that of the benzoxazole **3a** to the benzimidazole **3c**. Although both possess the same $P_4P_3P_2$ segment, *t*-butylacetyl-dPhePro, the enzyme selectivities of these two inhibitors are dramatically different. Relative to **3a**, **3c** appears to have similar potency against thrombin, but lost most of its potency against both Factor Xa and trypsin, trends that parallel those of related argininoyl heterocycles.⁷

Models of the compounds **3a** and **3c** complexed with thrombin (Figures 2–4) have been constructed from a crystal structure of thrombin.²⁰ Placement of **3a** in the active site of thrombin in a fashion analogous to the peptidyl α -ketobenzoxazoles in elastase⁴ produced favorable thrombin interactions of the benzoxazole NH and O with His 57 and Gly 193, respectively, which would also be possible in trypsin and factor Xa. Placement of **3c** in the active site of thrombin in the same fashion presents the favorable His 57 interaction, but the NH of the benzimidazole and its sphere of hydration has two unfavorable steric interactions between the conserved Gly 193 and the conserved H α of residue 192 of the trypsin class of serine proteases. The benzimidazole is

postulated to rotate to fit into the enzyme, similar to the rotation of the heterocycle in the benzothiazole:thrombin X-ray structure reported by Costanzo et al.⁷ According to our model, this twist may result in a favorable interactions of the 60 insertion loop (Trp 60d, Lys 60f) unique to thrombin. Thus, **3c** retains the overall binding energy in the S₁' site of thrombin, but not in other trypsin-like serine proteases.



Figure 2. Inhibitor **3a** docked into the active site of thrombin.



Figure 3. Inhibitor **3c** docked into the active site of thrombin.



Figure 4. Superposition of inhibitors **3a** and **3c** in the active site of thrombin.

Conclusion

A convenient synthesis of new argininoyl benzoxazoles and argininoyl benzimidazoles was developed and utilized for the preparation of a series of related argininoyl heterocycle targets as potential inhibitors of thrombin and factor Xa. These compounds expressed interesting levels of biological activity with varying selectivities towards related serine proteases. Further investigations of related argininoyl heterocycles will be reported at a later date.

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10. All new compounds were characterized by full spectroscopic (NMR, IR, low/high-resolution MS) data. Yields refer to spectroscopically and chromatographically homogeneous ($\geq 95\%$ by ^1H NMR, HPLC, TLC) materials.
11. Attempts to convert *t*-butylacetyl-dPhePro-nitroargininal cyanohydrin to the corresponding peptidoyl benzoxazole via the imide employing the standard protocol⁴ were unsuccessful in our laboratory.
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18. *t*-Butylacetyl-dPheProOH and Ac-dPheProOH, prepared using standard solution phase peptide coupling techniques, were spectroscopically and chromatographically homogeneous ($\geq 95\%$ by ^1H NMR, and TLC).
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