



Aminomethylthiophene-2-carboxylic Acids as Dipeptide Mimetic in New Growth Hormone Secretagogues

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Abstract 3-Aminomethylbenzoic acid is a well established dipeptide mimetic. Herein, aminomethylthiophene-2-carboxylic acids¹⁾ have been synthesized as analogues of 3-aminomethylbenzoic acid. Their use as a dipeptide-mimetic at the *N*-terminal of novel growth hormone secretagogues is described. © 1997 Elsevier Science Ltd.

Introduction

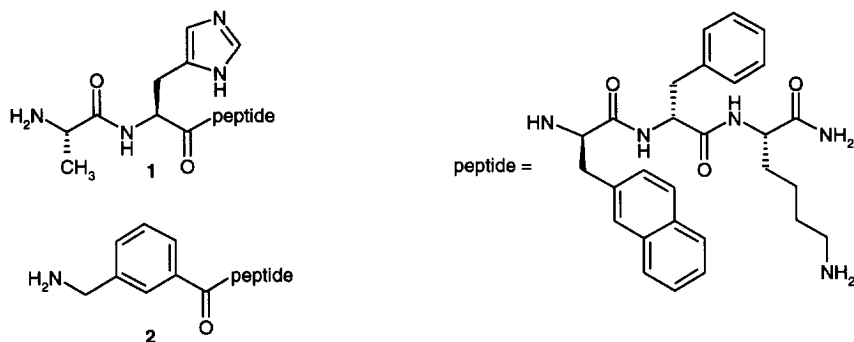
The discovery, that the hexapeptide His-*D*-Trp-Ala-Trp-*D*-Phe-Lys-NH₂ (GHRP-6)²⁾ releases growth hormone by a distinct mechanism, which is different from that of growth hormone releasing hormone (GHRH)³⁾, has resulted in a number of research programs, which lead to other peptidic growth hormone releasers, such as *D*-Ala-*D*-2-Nal-Ala-Trp-*D*-Phe-Lys-NH₂ (GHRP-2)⁴⁾ or non-peptidic releasers such as L-692,429⁵⁾ or MK-0677⁶⁾. The aim is to develop an orally available drug, that might advantageously substitute the direct administration of growth hormone in a number of therapeutical indications.

Discussion

During our screening for growth hormone secretagogues, we identified **1**⁷⁾ as lead compound. The utilization of the well established dipeptide mimetic 3-aminomethylbenzoic acid⁸⁾ at the *N*-terminal gave an equipotent compound **2**. We decided to focus on similar thiophene analogues¹⁾ and their ability to serve as dipeptide-mimetic.

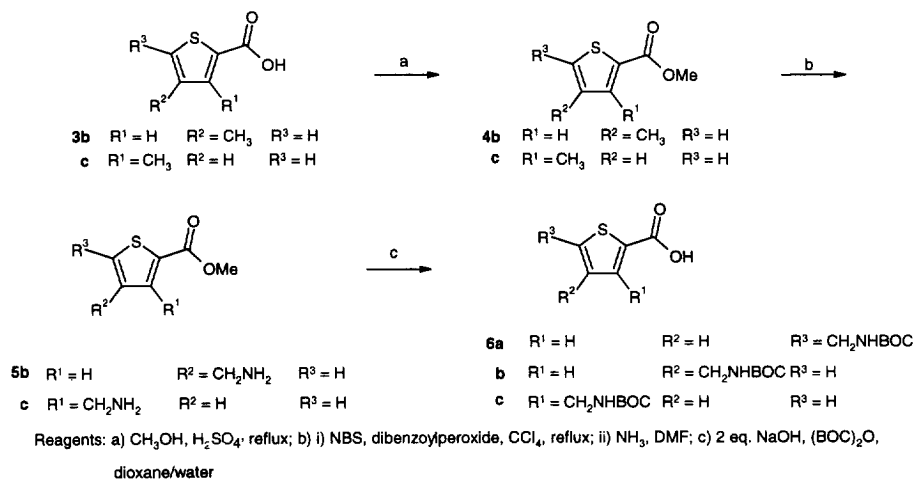
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Therefore the BOC-protected 5-aminomethyl-2-carboxylic acid (**6a**)⁸ and its isomers **6b** and **6c** were synthesized and incorporated into peptides of type **1**. Starting with methylthiophene-2-carboxylic acids **3b**⁹ and **3c**¹⁰, the corresponding esters **4b** and **4c** were prepared. NBS-bromination followed by treatment with ammonia in DMF furnished the aminoesters **5b** and **5c**¹¹. A one-pot saponification and BOC-protection afforded the protected amino acids **6b** and **6c**¹².

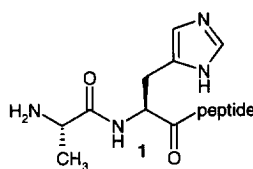
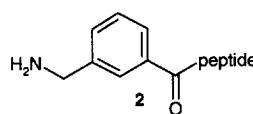
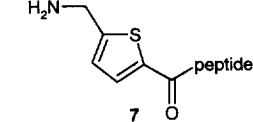
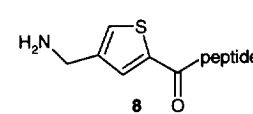
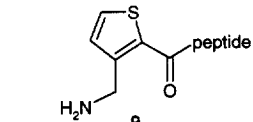
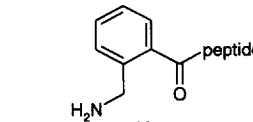
Scheme 1

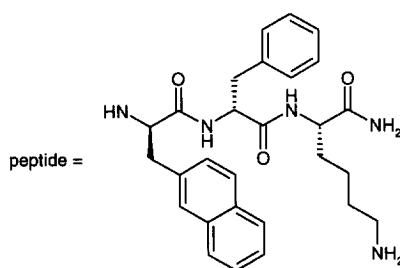


The amino acids **6a** - **c** were incorporated into the peptide by standard peptide synthesis on *Rink*-resin, and cleaved with trifluoroacetic acid¹³. The resulting peptides were tested in a functional *in vitro* rat pituitary assay^{14,15}. The test results are shown in Table 1 as mean values

of two experiments. The best dipeptide mimetics were 3-aminomethylbenzoic acid and its 2,3-substituted thiophene analogue **6c**, leading to peptides **2** and **9**, respectively.

Table 1

	EC ₅₀ [nM]
	6
	7
	210
	675
	27
	85



Conclusion

In growth hormone secretagogues of type **1**, the *N*-terminal Ala-His-moiety can be replaced by the known dipeptide-mimetic 3-aminomethylbenzoic acid without loss of activity. When the thiophene-analogue **6c** is used at the *N*-terminal, the resulting peptide **9** shows very similar activity. Surprisingly, **6a** and **6b** as *N*-terminal cause a sharp drop in activity of the peptides **7** and **8**, even though the orientation of substituents on the thiophene-rings is similar to that of 3-aminomethylbenzoic acid. Peptide **10**, with the *ortho*-substituted 2-aminomethylbenzoic

acid at the *N*-terminal, shows decreased activity, compared to **2**. From this one can conclude that, despite different orientation of substituents, 3-aminomethylbenzoic acid and 3-aminomethylthiophene-2-carboxylic acid (**6c**) are dipeptide mimetics at the *N*-terminal of growth hormone secretagogues. The low activities of **7** and **8** may be caused by different conformation of the amino acids **6a** or **6c** at the *N*-terminal due to enhanced sterical demand by the sulfur atom. Another hypothesis is hydrogen-bonding of the sulfur atom to groups at the binding site, that does not occur with benzene and results in a different orientation of the thiophene moiety.

References and Notes

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- 12) ¹H-NMR (DMSO d₆): **6b**: δ 1.40 (s, 9 H); 4.10 (d, 2 H); 7.40 (t, 1 H); 7.56 (s, 1 H); 7.61 (s, 1 H); 13.05 (br, 1 H). MS: calc: 257.07218, found: 257.071944.
¹H-NMR (DMSO d₆): **6c**: δ 1.40 (s, 9 H); 4.43 (d, 2 H); 7.05 (d, 1 H); 7.37 (t, 1 H); 7.75 (d, 1 H); 13.05 (br, 1 H). MS: 257.07215, found: 257.071864.
- 13) All peptides were characterized by MS, 2 independent HPLC-systems, and except for **10**, amino acid analysis.
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