



N-Methylbenzanilide Derivatives as a Novel Class of Selective V_{1A} Receptor Antagonists

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Abstract—During our efforts to develop a novel class of selective V_{1A} receptor antagonists, the *N*-methylbenzanilide structure was applied to a 4,4-difluoro-1-benzazepine derivative, **4**, which is a selective V_{1A} receptor antagonist. Further structural modifications gave **16a** with high V_{1A} affinity and V_2/V_{1A} selectivity ($K_i = 5.71$ nM, $V_2/V_{1A} = 140$) and potent V_{1A} receptor antagonist activity ($ID_{50} = 0.0080$ mg/kg iv). © 2002 Elsevier Science Ltd. All rights reserved.

Introduction

Arginine vasopressin (AVP) is a peptide hormone which is released from the posterior pituitary and exerts a variety of biological effects in mammals. So far, two AVP receptor subtypes (V_{1A} and V_2) have been identified, in periphery.¹ The V_{1A} receptor mediates phospholipase C activation and intracellular calcium mobilization, which causes various known AVP actions, such as vasoconstriction, platelet aggregation, and growth of vascular smooth muscle cell.¹ The V_2 receptor is positively coupled to adenylate cyclase and plays a predominant role in the antidiuretic response to AVP which promotes water reabsorption.²

AVP may play a role in several diseases and disorders, such as heart failure, hypertension, coronary renal vasospasm, hyponatremia, and dysmenorrhea.^{3,4} Therefore, the development of subtype selective AVP receptor antagonists is essential to investigate the pathophysiological roles of AVP and could lead to new therapeutic tools. Recently, two nonpeptide selective V_{1A} receptor antagonists, OPC-21268 (**1**)^{5,6} and SR49059 (**2**)⁷ have been reported (Fig. 1). They have undergone clinical testing.

In the course of the search for AVP antagonists at our laboratories, the 4,4-difluoro-5-methylene-2,3,4,5-tetra-

hydro-1*H*-1-benzazepine derivative, **3** (V_{1A} $K_i = 0.160$ nM, V_2 $K_i = 0.770$ nM), was discovered as a V_{1A} and V_2 receptor dual antagonist.⁸ We moved into the next research program which addressed the development of selective V_{1A} receptor antagonists. Extensive study of the 4,4-difluoro-1-benzazepine derivatives afforded a series of compounds with the selectivity for the V_{1A} receptor versus the V_2 receptor, such as the 2-methyl-3-furyl derivative, **4** (V_{1A} $K_i = 0.102$ nM, V_2 $K_i = 33.9$ nM, $V_2/V_{1A} = 330$).⁹ On the other hand, Ohkawa et al. reported the *N*-methylbenzanilide derivative, FR179544 (**5**), as a V_{1A} and V_2 receptor dual antagonist.¹⁰ This report prompted us to apply the *N*-methylbenzanilide structure to **4**, which might afford a novel class of selective V_{1A} receptor antagonists. Subsequently, we attempted modifications of the tether linking the *N*-methylbenzanilide template with the terminal amide part, and the 2-methyl-3-furyl part. In this paper, we describe the synthesis and pharmacological evaluation of *N*-methylbenzanilide derivatives.

Chemistry

The basic structure of the novel compounds was constructed by condensation of a *N*-methylaniline with a 4-acylaminobenzoyl chloride (Scheme 1) or condensation of a *N*-methyl-4'-aminobenzanilide with a substituted benzoyl chloride (Scheme 2). Scheme 1 shows the preparation of compounds **9a–c** and **12a–d**. The *N*-methylaniline (**6**) was coupled with a 4-[(2-methyl-3-furyl)carboxamido]benzoyl chloride to afford the *N*-

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Table 1. Binding affinities of *N*-methylbenzanilide derivatives for V_{1A} and V₂ receptors

Compd	R1	R2	R3	K_i^a (nM)		Selectivity K_i ratio V_2/V_{1A}
				V_{1A}	V_2	
9a		Me		8.77 ± 0.66	73.4 ± 11	8.4
9b		Me		4.97 ± 0.73	94.5 ± 20	19
12a		H		1140 ± 230	2250 ± 160	2.0
12b		H		491 ± 34	1360 ± 270	2.8
12c		H		78.6 ± 11	168 ± 22	2.1
12d		H		335 ± 93	2040 ± 420	6.1
9c		Me		2.93 ± 0.24	41.5 ± 9.3	14
16a		Me		5.71 ± 0.83	782 ± 110	140
16b		Me		7.59 ± 0.48	445 ± 36	59
1				23.5 ± 4.39	$> 10,000$	> 430
3				0.620 ± 0.23	1.19 ± 0.02	1.9
4				0.102^b	33.9^b	330

^a K_i values were obtained from two or more independent experiments. Each value indicates mean \pm SEM.

^bMean from two experiments.

derivatives revealed that the 4-aminopiperidine ring as the terminal basic part is superior to the 4-alkylpiperazine ring with respect to the V_2/V_{1A} selectivity.¹⁴ Thus, we utilized these results to **9a** having a 4-methylpiperazine to obtain the 4-piperidinopiperidine derivative (**9b**). As a result, compound **9b** exhibited improved V_{1A} affinity and V_2/V_{1A} selectivity ($K_i = 4.97$ nM, $V_2/V_{1A} = 19$) in comparison to **9a**. This result was similar to the SAR of **3** and its derivatives.

We then modified the tether linking the *N*-methylbenzanilide template with the terminal amide of **9b**. The comparison of the binding results of the phenoxy derivatives (**12a–c**) demonstrated that the rank of order of potency in their V_{1A} affinities was 1,4- > 1,3- > 1,2-

substitution pattern. Therefore, we examined another aromatic tethers with 1,4-substitution pattern. The phenyl derivative (**12d**) showed considerably weak V_{1A} affinity. In contrast, the benzyloxy derivative (**9c**) exhibited high V_{1A} affinity ($K_i = 2.93$ nM), being better than that of **9b**. These results suggested that the distance between the terminal basic moiety and the *N*-methylbenzanilide template might be important to exert high V_{1A} affinity and that the 4-benzyloxy moiety might be suitable for the tether.

Next, we converted the 2-methyl-3-furyl part of **9** into another heterocycle utilizing the SAR of **3**.¹⁴ The 2-(2-ethyl-1-imidazolyl)phenyl derivatives (**16a–b**) showed similar V_{1A} affinities, but reduced V_2 affinities compared

Table 2. In vivo activities of *N*-methylbenzanilide derivatives

Compd	ID ₅₀ ^a (mg/kg iv)	<i>n</i>
9a	0.11 ± 0.01	3
9c	0.038 ± 0.002	4
16a	0.0080 ± 0.002	4
16b	0.015 ± 0.005	4
1	0.34 ± 0.05	6

^aID₅₀ represents the dose of the compounds required to inhibit AVP-induced pressor response in pithed rats by 50%. Each value indicates mean ± SEM.

to **9a**, which resulted in an improvement of V_2/V_{1A} selectivity. Especially, compound **16a** had high V_2/V_{1A} selectivity ($V_2/V_{1A} = 140$).

Thus, application of *N*-methylbenzanilide structure to **4** and further structural modifications utilizing the SAR of **3** led to novel compounds which had high V_{1A} affinity and V_2/V_{1A} selectivity.

From the results of the binding assay, some compounds were selected for in vivo evaluation¹³ (Table 2). All the tested compounds dose-dependently antagonized an increase in DBP induced by AVP (30 mU/kg iv) via the V_{1A} receptor. In particular, compound **16a**, which had the highest V_2/V_{1A} selectivity in this series, exhibited potent V_{1A} receptor antagonist activity with an ID₅₀ value of 0.0080 mg/kg, which was 43-fold more potent than that of **1**.

Conclusion

Application of the *N*-methylbenzanilide structure to a selective V_{1A} receptor antagonist, **4**, gave the novel 2-methyl-3-furyl derivative (**9a**) which showed high V_{1A} affinity and moderate V_2/V_{1A} selectivity. Further modifications utilizing the SAR of **3** gave novel compounds which exhibit high V_{1A} affinity and high V_2/V_{1A} selectivity, such as the 2-(2-ethyl-1-imidazolyl)phenyl derivative (**16a**). Compound **16a** showed potent V_{1A} receptor antagonist activity in vivo and was 43-fold more potent than **1**. This study would provide a novel approach to selective V_{1A} receptor antagonists, which might be useful in the treatment of diseases in which AVP is involved via the V_{1A} receptor.

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- Satisfactory analytical data were obtained for all the target compounds. For example, **16a**: colorless amorphous powder: ¹H NMR (400 MHz, DMSO-*d*₆) δ 1.16 (3H, t, *J* = 7.6 Hz), 1.43–1.47 (2H, m), 1.53–1.61 (2H, m), 1.69–1.76 (2H, m), 2.22 (3H, s), 2.39 (2H, t, *J* = 7.6 Hz), 2.73 (5H, br s), 3.02–3.51 (8H, m), 3.15 (3H, s), 3.87–3.93 (2H, m), 6.63 (1H, d, *J* = 8.0 Hz), 6.79 (1H, s), 6.99 (1H, d, *J* = 8.0 Hz), 7.19 (2H, d, *J* = 8.0 Hz), 7.42 (2H, d, *J* = 8.0 Hz), 7.71–7.81 (5H, m), 7.93–7.95 (1H, m), 10.81 (1H, s), 11.41 (1H, br s), 15.05 (1H, br s); MS (FAB) *m/z* 651 (MH⁺). Anal. calcd for C₃₈H₄₆N₆O₄·2HCl·1.8H₂O: C, 60.36, H, 6.88, N, 11.11; Cl, 9.38. Found: C, 60.32, H, 7.15, N, 11.00, Cl, 9.45.
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