

Potent Antagonists of the Antidiuretic Responses to Arginine-vasopressin Based on Modifications of [1-(β -Mercapto- β,β -cyclopentamethylenepropionic Acid),2-D-phenylalanine,4-valine]arginine-vasopressin at Position 4[†]

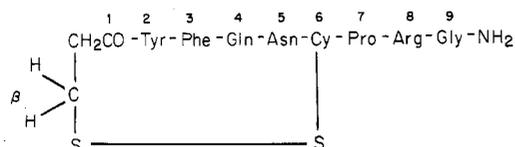
Maurice Manning,*[‡] Aleksandra Olma,^{‡,§} Wieslaw A. Klis,^{‡,||} Janny Seto,[§] and Wilbur H. Sawyer[§]

Department of Biochemistry, Medical College of Ohio at Toledo, Toledo, Ohio 43699, and Department of Pharmacology, College of Physicians and Surgeons of Columbia University, New York, New York 10032. Received January 7, 1983

As part of a program in which we are attempting (a) to delineate the structural features at positions 1-9 in our previously reported antidiuretic antagonists required for antidiuretic antagonism and (b) to obtain analogues with enhanced antidiuretic potency and/or selectivity, we have synthesized 14 new analogues of the antidiuretic antagonist [1-(β -mercapto- β,β -cyclopentamethylenepropionic acid),2-D-phenylalanine,4-valine]arginine-vasopressin [d-(CH₂)₅-D-Phe²VAVP], in which the valine residue at position 4 was replaced by the following L-amino acids and glycine: Ile, Abu, Thr, Ala, Gln, Lys, Cha, Nle, Nva, Phe, Leu, Gly, Tyr, and Pro. These analogues are 1, d-(CH₂)₅-D-Phe²Ile⁴AVP; 2, d-(CH₂)₅-D-Phe²Abu⁴AVP; 3, d-(CH₂)₅-D-Phe²Thr⁴AVP; 4, d-(CH₂)₅-D-Phe²Ala⁴AVP; 5, d-(CH₂)₅-D-Phe²AVP; 6, d-(CH₂)₅-D-Phe²Lys⁴AVP; 7, d-(CH₂)₅-D-Phe²Cha⁴AVP; 8, d-(CH₂)₅-D-Phe²Nle⁴AVP; 9, d-(CH₂)₅-D-Phe²Nva⁴AVP; 10, d-(CH₂)₅-D-Phe²Phe⁴AVP; 11, d-(CH₂)₅-D-Phe²Leu⁴AVP; 12, d-(CH₂)₅-D-Phe²Gly⁴AVP; 13, d-(CH₂)₅-D-Phe²Tyr⁴AVP; 14, d-(CH₂)₅-D-Phe²Pro⁴AVP. The protected intermediates required for the synthesis of all of these peptides were prepared by the solid-phase method and cleaved from the resin by ammonolysis. Following deblocking with Na in NH₃ and oxidizing with K₃[Fe(CN)₆], each peptide was purified on Sephadex G-15 in a two-step procedure using 50% HOAc and 0.2 M HOAc as eluents. Analogues 1-14 were tested for agonistic and antagonistic activities by antidiuretic, vasopressor, and oxytocic assays in rats. Analogues 1, 2, and 4-6 exhibit no detectable antidiuretic agonistic activity. All analogues, with the exception of the Pro⁴-containing analogue, are antidiuretic antagonists. Their antidiuretic pA₂ values are as follows: 1, 8.24 ± 0.08; 2, 7.96 ± 0.07; 3, 7.62 ± 0.09; 4, 7.52 ± 0.03; 5, 7.21 ± 0.07; 6, 7.22 ± 0.12; 7, 7.19 ± 0.08; 8, 7.12 ± 0.09; 9, 6.99 ± 0.06; 10, 6.07 ± 0.11; 11, 6.07 ± 0.11; 12, 5.85 ± 0.05; 13, ~5.57; 14, a weak agonist (0.004 U/mg). Analogues 1-14 also antagonize the vascular responses to arginine-vasopressin (AVP) and the in vitro oxytocic responses to oxytocin. Analogues 1, 2, 3, and 5 have also been shown to antagonize the in vivo oxytocic responses to oxytocin. Five of these analogues (1, 2, 3, 6, and 7) exhibit enhanced antidiuretic/antivasopressor selectivity. d-(CH₂)₅-D-Phe²Lys⁴AVP and other position-4 analogues with side-chain functional groups may be useful covalent ligands with which to probe the structural characteristics of AVP renal and vascular receptors. With an antidiuretic "effective dose" of 0.46 ± 0.07 nmol/kg and a pA₂ value of 8.24 ± 0.08, d-(CH₂)₅-D-Phe²Ile⁴AVP (1) appears to be the most potent antidiuretic antagonist reported to date. This and some of the other analogues reported here should prove to be useful pharmacological tools for studies on the role(s) of AVP in the etiology of water retention states and may also be of value as therapeutic agents for the treatment of such conditions in humans. These findings also provide valuable insights for the design of more potent and selective antidiuretic antagonists.

Following our recent discovery of the first effective antagonists of in vivo antidiuretic responses to arginine-vasopressin (AVP),^{1,2} we embarked on a program aimed at

[†]Symbols and abbreviations are in accordance with the recommendations of the IUPAC-IUB Commission on Biochemical Nomenclature (*J. Biol. Chem.* 1971, 247, 977). All amino acids are in the L configuration unless otherwise noted. Other abbreviations used are: Abu, α -aminobutyric acid; Cha, cyclohexylalanine; Nle, norleucine; Nva, norvaline; DMF, dimethylformamide; DCC, dicyclohexylcarbodiimide; BOC, *tert*-butyloxycarbonyl; Bzl, benzyl; Tos, tosyl; HOAc, acetic acid; HOBT, *N*-hydroxybenzotriazole, NPE, nitrophenyl ester; AVP, arginine-vasopressin; VAVP, [4-valine]arginine-vasopressin; d, 1-deamino; (CH₂)₅, cyclopentamethylene; d(CH₂)₅, 1-deaminocyclopentamethylene. The d(CH₂)₅ abbreviation, rather than the more cumbersome 1- β -Mpa(β -CH₂)₅¹² abbreviation, is used as the designation for the 1-(β -mercapto- β,β -cyclopentamethylenepropionic acid) residue at position 1 in analogues of [1-deamino]arginine-vasopressin (dAVP) and [1-deamino]oxytocin. dAVP has the following structure:



Thus, the d(CH₂)₅ abbreviation as employed in this and in many previous publications from these^{1-6,19,20,22} and numerous other laboratories (see, for example, ref 7-9 cited in Manning, M.; Lammek, B.; Kruszynski, M.; Seto, J.; Sawyer, W. H., *J. Med. Chem.* 1982, 25, 408) specifies that the two hydrogens on the β -carbon of the β -mercapto- β,β -cyclopentamethylenepropionic acid (β -Mpa) residue at position 1 in [1-deamino]arginine-vasopressin and analogues are replaced by a cyclopentamethylene, (CH₂)₅, substituent.

enhancing their antidiuretic potency and antidiuretic/antivasopressor selectivity and eliminating their weak transient antidiuretic agonistic activity. We selected the most potent of the original antagonists, [1-(β -mercapto- β,β -cyclopentamethylenepropionic acid),2-O-ethyltyrosine,4-valine]arginine-vasopressin [d-(CH₂)₅Tyr(Et)VAVP],^{1,2} as the lead compound and focused initially on modifications at position 2.³⁻⁵ These studies led in time to two new lead compounds, [1-(β -mercapto- β,β -cyclopentamethylenepropionic acid),2-D-phenylalanine,4-valine]arginine-vasopressin [d(CH₂)₅-D-Phe²VAVP] and its D-Ile² analogues, d(CH₂)₅-D-Ile²VAVP.⁵ d(CH₂)₅-D-Phe²-VAVP and d(CH₂)₅-D-Ile²VAVP are the most potent antidiuretic antagonists previously reported. They are almost 3 times more potent than d(CH₂)₅Tyr(Et)VAVP. Furthermore, they are devoid of evident antidiuretic agonism and also exhibit enhanced antidiuretic/antivasopressor selectivity relative to that of d-

[‡]Medical College of Ohio at Toledo.

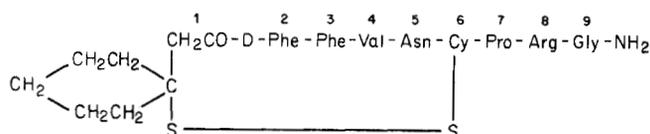
[§]College of Physicians and Surgeons of Columbia University.

^{||}Visiting investigator from Technical of Lodz, Poland.

^{||}Visiting investigator from University of Wroclaw, Poland.

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(CH₂)₅Tyr(Et)VAVP. Their ease of synthesis relative to d(CH₂)₅Tyr(Et)VAVP⁵ was another decisive reason in favor of using d(CH₂)₅-D-Phe²VAVP and d(CH₂)₅-D-Ile₂VAVP to explore modifications at other positions. d(CH₂)₅-D-Phe²VAVP has the following structure:



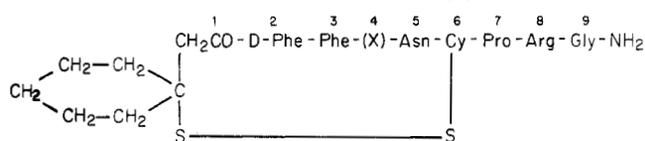
In this report we describe a series of analogues of d(CH₂)₅-D-Phe²VAVP in which the valine residue at position 4 has been replaced by a number of other amino acids. We will present our findings on a related series of position-4-modified analogues of d(CH₂)₅-D-Ile²VAVP separately.

Our reasons for selecting position 4 for study are as follows: We had clear evidence for a critical requirement for valine at position 4 in our original antidiuretic antagonists, i.e., those in which position 2 is occupied by an L-Tyr(alk) residue (where alk = Me, Et, *i*-Pr and *n*-Pr).^{1,2} Thus, for example, although d(CH₂)₅Tyr(Me)AVP⁶ and d(CH₂)₅Tyr(Et)AVP,⁷ both of which have glutamine at position 4, are potent vasopressor antagonists, they are also weak antidiuretic agonists. Valine was thus incorporated as a matter of course in all subsequently designed antidiuretic antagonists aimed at determining how chiral, steric, and electronic changes at the remaining positions would effect antidiuretic potency. In the light of our preliminary findings from these studies, we decided to reassess the requirement for valine and were particularly intrigued by the possibility that valine might not be an essential requirement in those antagonists that have a D-amino acid at position 2. In a preliminary investigation in which valine in d(CH₂)₅-D-Phe²VAVP was replaced by isoleucine, we were gratified to find that the resulting compound, d(CH₂)₅-D-Phe²Ile⁴AVP, was a highly potent antidiuretic antagonist. Encouraged by this finding and by the possibility of obtaining even more potent antidiuretic antagonists, we decided to further examine the structural features of the amino acid side chain at position 4 in d(CH₂)₅-D-Phe²X⁴AVP required for antidiuretic potency and selectivity. In addition to isoleucine, we selected the following L-amino acids as replacements for the valine residue at position 4 in d(CH₂)₅-D-Phe²VAVP: Abu, Thr, Ala, Gln, Lys, Cha, Nle, Nva, Phe, Leu, Gly, Tyr, and Pro.

The 14 new analogues designed following the above reasoning are as follows: 1, [1-(β-mercapto-β,β-cyclopentamethylenepropionic acid),2-D-phenylalanine,4-isoleucine]arginine-vasopressin [d(CH₂)₅-D-Phe²Ile⁴AVP]; 2, [1-(β-mercapto-β,β-cyclopentamethylenepropionic acid),2-D-phenylalanine,4-(α-aminobutyric acid)]arginine-vasopressin [d(CH₂)₅-D-Phe²Abu⁴AVP]; 3, [1-(β-mercapto-β,β-cyclopentamethylenepropionic acid),2-D-phenylalanine,4-threonine]arginine-vasopressin [d(CH₂)₅-D-Phe²Thr⁴AVP]; 4, [1-(β-mercapto-β,β-cyclopentamethylenepropionic acid),2-D-phenylalanine,4-alanine]arginine-vasopressin [d(CH₂)₅-D-Phe²Ala⁴AVP]; 5, [1-(β-mercapto-β,β-cyclopentamethylenepropionic acid),2-D-phenylalanine]arginine-vasopressin [d(CH₂)₅-D-Phe²AVP]; 6, [1-(β-mercapto-β,β-cyclopentamethylenepropionic acid),2-D-phenylalanine,4-lysine]arginine-vasopressin [d(CH₂)₅-D-Phe²Lys⁴AVP]; 7, [1-(β-mercapto-β,β-cyclopentamethylenepropionic acid),2-D-phenylalanine,4-cyclohexylalanine]arginine-vasopressin [d(CH₂)₅-D-

Phe²Cha⁴AVP]; 8, [1-(β-mercapto-β,β-cyclopentamethylenepropionic acid),2-D-phenylalanine,4-norleucine]arginine-vasopressin [d(CH₂)₅-D-Nle⁴AVP]; 9, [1-(β-mercapto-β,β-cyclopentamethylenepropionic acid),2-D-phenylalanine,4-norvaline]arginine-vasopressin [d(CH₂)₅-D-Phe²Nva⁴AVP]; 10, [1-(β-mercapto-β,β-cyclopentamethylenepropionic acid),2-D-phenylalanine,4-phenylalanine]arginine-vasopressin [d(CH₂)₅-D-Phe²Phe⁴AVP]; 11, [1-(β-mercapto-β,β-cyclopentamethylenepropionic acid),2-D-phenylalanine,4-leucine]arginine-vasopressin [d(CH₂)₅-D-Phe²Leu⁴AVP]; 12, [1-(β-mercapto-β,β-cyclopentamethylenepropionic acid),2-D-phenylalanine,4-glycine]arginine-vasopressin [d(CH₂)₅-D-Phe²Gly⁴AVP]; 13, [1-(β-mercapto-β,β-cyclopentamethylenepropionic acid),2-D-phenylalanine,4-tyrosine]arginine-vasopressin [d(CH₂)₅-D-Phe²Tyr⁴AVP]; 14, [1-(β-mercapto-β,β-cyclopentamethylenepropionic acid),2-D-phenylalanine,4-proline]arginine-vasopressin [d(CH₂)₅-D-Phe²Pro⁴AVP].

These 14 analogues have the following general structure:



X = Ile (1), Abu (2), Thr (3), Ala (4), Gln (5), Lys (6), Cha (7), Nle (8), Nva (9), Phe (10), Leu (11), Gly (12), Tyr (13), Pro (14).

We now present the synthesis and some pharmacological properties of these 14 analogues.

Peptide Synthesis. The protected peptide precursors required for the synthesis of the 14 analogues of d(CH₂)₅-D-Phe²VAVP were prepared by the solid-phase method of peptide synthesis^{8,9} using previously described modifications.^{6,10,11} *p*-Nitrophenyl β-(benzylmercapto)-β,β-cyclopentamethylenepropionate¹² was used in each final coupling step. All active ester couplings^{13,14} were facilitated by the addition of *N*-hydroxybenzotriazole (HOBT).¹⁵ The 14 protected peptide amides were obtained by ammonolytic cleavage from the respective acyl octapeptide resins.^{10,11} Each protected precursor was deprotected with Na in NH₃¹⁶ by a previously described yield-enhancing modification of the standard workup procedure.² The deprotected disulfhydryl compounds were oxidatively cyclized with dilute potassium ferricyanide.¹⁷ The analogues were desalted and purified by gel filtration on Sephadex G-15 as previously described.^{2,5,6,18}

Bioassay Methods. The agonistic and antagonistic potencies of these analogues were measured by previously described methods.^{1,2,6,19,20} These included intravenous

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vasopressor assays in phenoxybenzamine-treated rats under urethane anesthesia, antidiuretic assays in rats under ethanol anesthesia, and oxytocic assays on isolated rat uteri and rat uteri in situ. The USP posterior pituitary reference standard was used in assays for agonistic and antagonistic activities, except for the rat uterus in situ. Synthetic oxytocin (Syntocinon, Sandoz) was used as the agonist in these assays. Agonistic activities are expressed in units per milligram. Antagonistic potencies were determined and expressed as "effective doses" and as pA_2 values.²¹ The effective dose is defined as the dose (in nanomoles per kilogram) that reduces the response seen from $2x$ units of agonist to the response with $1x$ unit of agonist. Estimated in vivo " pA_2 " values represent the negative logarithms of the effective doses divided by the estimated volume of distribution (67 mL/kg). Inhibition of antidiuretic responses was tested by injecting the standard 20 min after injecting the antagonist, to allow for recovery from the initial antidiuretic responses to some of the antagonists. For those analogues showing no agonistic activity (compounds 1, 2, and 4-6), the standard could be injected 10 min after the antagonist. Each peptide was administered in two doses, a high dose, which reduced the response to $2x$ units of agonist to less than the response to $1x$ unit of agonist, and a low dose, which did not fully reduce the response to that given by $1x$ unit of agonist. The effective dose in each case was obtained by interpolation on a logarithmic scale between the two doses of antagonist.²¹

Results and Discussion

The antiantidiuretic, antivasopressor, and antioxytocic (in vitro for all analogues, in vivo only for some) potencies of the 14 $d(CH_2)_5$ -D-Phe²X⁴AVP analogues, where X = Ile, Abu, Thr, Ala, Gln, Lys, Cha, Nle, Nva, Phe, Leu, Gly, Tyr, and Pro, together with their antivasopressor/antiantidiuretic and antioxytocic/antiantidiuretic "effective dose" ratios, are presented in Tables I and II. With the exception of the Pro⁴ analogue, all of these peptides antagonize in vivo antidiuretic responses to AVP. However, the range of antiantidiuretic potencies as reflected by their effective doses and by their pA_2 values is very broad; from a high of 0.46 nmol/kg and a $pA_2 = 8.24$ for $d(CH_2)_5$ -D-Phe²Ile⁴AVP to a low of ~178 nmol/kg and a $pA_2 = \sim 5.57$ for $d(CH_2)_5$ -D-Phe²Tyr⁴AVP. $d(CH_2)_5$ -D-Phe²Ile⁴AVP appears to be more potent as an antiantidiuretic than its parent analogue, $d(CH_2)_5$ -D-Phe²VAVP ($pA_2 = 8.06$) and, thus, the most potent antidiuretic antagonist reported to date.

Val⁴ Requirement for Antidiuretic Antagonism Is No Longer Valid. On the basis of earlier findings^{1,2,6,7} we had assumed that the presence of a valine residue at position 4 is critical for antidiuretic antagonism. Thus, although $d(CH_2)_5$ Tyr(alk)AVP (where alk = Me, Et), which have Gln at position 4, are potent vasopressor antagonists, they are also weak antidiuretic agonists.^{6,7} Replacement of Gln by Val at position 4 converted these into potent antidiuretic antagonists.^{1,2} The surprising findings reported here illustrate very vividly the need for caution in drawing conclusions regarding structural requirements on the basis of single changes at a given position, for it is clear that when position 2 in $d(CH_2)_5$ Y²X⁴AVP is occupied by D-Phe, position 4 can tolerate a wide variety of aliphatic amino acids with retention of high antiantidiuretic potency.

Antagonists That Lack Apparent Agonistic Activity. As indicated earlier, $d(CH_2)_5$ -D-Phe²VAVP appears to be a pure in vivo antidiuretic antagonist; i.e., it has no evident agonistic activity.⁵ Remarkably, of the 13 new antagonists presented here, only 5 have retained this feature. These analogues are $d(CH_2)_5$ -D-Phe²Ile⁴-AVP; $d(CH_2)_5$ -D-Phe²Abu⁴-AVP; $d(CH_2)_5$ -D-Phe²Ala⁴AVP; $d(CH_2)_4$ -D-Phe²-AVP, and $d(CH_2)_5$ -D-Phe²Lys⁴-AVP. We cannot explain why all the remaining analogues have lost this characteristic.

Effects of Aliphatic vs. Aromatic Character at Position 4 on Antiantidiuretic Potencies. Aliphatic amino acids are clearly more effective than aromatic amino acids in endowing these position-4-modified peptides with potent antidiuretic antagonism. Thus, the Phe⁴ and Tyr⁴ analogues are much weaker antidiuretic antagonists than nearly all of the antagonists that have aliphatic amino acids at position 4 (Table I). This could be due to a combination of steric and electronic, i.e., aromaticity, factors. The possibility that aromatic character may play a more deleterious role than steric bulk may be ascertained by the fact that the Cha⁴-containing antagonist is over 10 times more potent than the Phe⁴ antagonist, although both amino acid side chains are of approximately equal size. These findings are in dramatic contrast to our previous observations on the structural requirements for aliphatic vs. aromatic D-amino acid substitutions at position 2 in antidiuretic antagonists.⁵ Thus, $d(CH_2)_5$ -D-Phe²VAVP and its D-Ile² analogue have virtually identical antiantidiuretic potencies, indicating virtually no differences between aliphatic and aromatic side chains at position 2 in these analogues.⁵

Effects of β -Branching, Steric Factors, and Ionization at Position 4 on Antiantidiuretic Potencies. The potencies of these new antidiuretic antagonists correlate reasonably well with the presence or absence of branching on the β -carbon of the side chain at position 4. Branching on the γ -carbon is clearly most deleterious. Thus, the Ile⁴ analogue is 20 times more potent than the Nle⁴ analogue, which in turn is 10 times more potent than the Leu⁴ analogue. Similarly, the Val⁴ analogue is over 10 times more potent than the Nva⁴ analogue. The need for a minimum aliphatic side chain at position 4 is illustrated by the fact that the Ala⁴ antagonist is over 600 times more potent than the Gly⁴ antagonist. The presence of a positively charged side chain at position 4 has been well tolerated. Thus, with an antiantidiuretic $pA_2 = 7.22$, the Lys⁴ analogue is equipotent with the Gln⁴ and Nle⁴-containing analogues. $d(CH_2)_5$ -D-Phe²Lys⁴AVP and related analogues, e.g., Orn⁴ or one containing some other functional or photoreactive group, could be useful tools with which to explore renal or vascular AVP receptors by covalent attachment via the δ - or ϵ -amino group at position 4.

Effects of Position-4 Modifications on Antivasopressor Potencies. The structural features at position 4 in $d(CH_2)_5$ -D-Phe²X⁴AVP required for vasopressor antagonism appear to be even less rigid than those required for antidiuretic antagonism. Analysis of the data in Table I reveals some interesting points. (1) The Phe⁴, Tyr⁴, Leu⁴, and Gly⁴ substitutions are not nearly as deleterious to antivasopressor potencies as they are to antiantidiuretic potencies. In fact, with an antivasopressor $pA_2 = 7.70$, the Leu⁴-containing analogue is one of the more potent vasopressor antagonists in the entire series. (2) In this series, aliphatic amino acids are clearly superior to aromatic amino acids at position 4 for preserving antivasopressor potency. (3) With an antivasopressor $pA_2 = 8.35$, the

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Table II. In Vitro and Some in Vivo Antioxytotic Potencies of AVP Antagonists Modified at Position 4

no.	peptide ^f	antioxytotic activities (rat uterus)				
		in vitro		in vivo		
		no Mg ²⁺ : pA ₂ ^{b,c}	0.5 mM Mg ²⁺ : pA ₂ ^{b,c}	ED, ^{a,c} nmol/kg	pA ₂	ED ratio ^d
1	d(CH ₂) ₅ -D-Phe ² VAVP ^e	7.74 ± 0.06	8.29 ± 0.05	8.9 ± 1.9	6.92 ± 0.10 (5)	13
2	d(CH ₂) ₅ -D-Phe ² Ile ⁴ AVP	8.04 ± 0.09	8.19 ± 0.06	5.8 ± 1.7	7.11 ± 0.12	13
3	d(CH ₂) ₅ -D-Phe ² Abu ⁴ AVP	7.73 ± 0.07	7.76 ± 0.08	2.4 ± 0.6	7.52 ± 0.10	3.1
4	d(CH ₂) ₅ -D-Phe ² Thr ⁴ AVP	8.04 ± 0.08	8.01 ± 0.11	3.2 ± 0.4	7.35 ± 0.07	1.9
5	d(CH ₂) ₅ -D-Phe ² Ala ⁴ AVP	7.68 ± 0.12	7.73 ± 0.05			
6	d(CH ₂) ₅ -D-Phe ² AVP	8.59 ± 0.08	8.43 ± 0.07	4.0 ± 0.6	7.25 ± 0.06	0.9
7	d(CH ₂) ₅ -D-Phe ² Lys ⁴ AVP	7.60 ± 0.06	7.47 ± 0.07			
8	d(CH ₂) ₅ -D-Phe ² Cha ⁴ AVP	7.78 ± 0.09	7.59 ± 0.07			
9	d(CH ₂) ₅ -D-Phe ² Nle ⁴ AVP	7.54 ± 0.07	7.78 ± 0.10			
10	d(CH ₂) ₅ -D-Phe ² Nva ⁴ AVP	8.03 ± 0.08	7.72 ± 0.05			
11	d(CH ₂) ₅ -D-Phe ² Phe ⁴ AVP	7.57 ± 0.07	7.67 ± 0.06			
12	d(CH ₂) ₅ -D-Phe ² Leu ⁴ AVP	7.76 ± 0.05	7.88 ± 0.03			
13	d(CH ₂) ₅ -D-Phe ² Gly ⁴ AVP	7.76 ± 0.08	7.33 ± 0.09			
14	d(CH ₂) ₅ -D-Phe ² Tyr ⁴ AVP	7.68 ± 0.10	7.43 ± 0.06			
	d(CH ₂) ₅ -D-Phe ² Pro ⁴ AVP	5.76 ± 0.07	5.43 ± 0.05			

^{a,b,c,e} See Table I. ^d ED ratio = antioxytotic ED/antiantidiuretic ED. ^f Abbreviation as in Table I.

Gln⁴-containing analogue is the most potent of the entire series.

Enhancement of Antiantidiuretic/Antivasopressor Selectivity. Only five of these analogues exhibit enhanced antiantidiuretic/antivasopressor selectivity, as reflected by their effective dose ratios. These are the Ile⁴, Abu⁴, Thr⁴, Lys⁴, and Cha⁴ analogues. With an antiantidiuretic/antivasopressor effective dose ratio of 2.2, d(CH₂)₅-D-Phe²Ile⁴AVP is the most selective of the series. It may be recalled that three of our earlier analogues, d(CH₂)₅-D-Leu²VAVP, d(CH₂)₅-D-Ile²VAVP, and d(CH₂)₅-D-Val²VAVP, exhibited effective dose ratios of 22, 12, and 12, respectively.⁵ Thus, the incorporation of Ile at position 4 in each of these three peptides might be expected to result in further enhancements of their antiantidiuretic/antivasopressor selectivities.

Effects of Position-4 Substituents on in Vitro and in Vivo Antioxytotic Potencies. All of these peptides inhibit in vitro responses to oxytocin on the rat uterus. Four of the most potent of the in vitro antagonists were also tested in vivo. These are d(CH₂)₅-D-Phe²Ile⁴AVP; d(CH₂)₅-D-Phe²Abu⁴AVP; d(CH₂)₅-D-Phe²Thr⁴AVP, and d(CH₂)₅-D-Phe²AVP. All four are potent antioxytotics in vivo. With a pA₂ of 7.52, d(CH₂)₅-D-Phe²Abu⁴AVP compares very favorably to some of our previously reported potent in vivo oxytotic antagonists²² and is, in fact, the most potent in vivo antioxytotic compound reported to date.

Conclusion

We have shown that replacement of the Val⁴ residue in the potent antidiuretic antagonist d(CH₂)₅-D-Phe²VAVP⁵ by a series of 14 different amino acids leads to retention of potent antidiuretic antagonism in a majority of the resulting analogues. This demonstrates that when D-Phe is substituted in position 2 in our original antagonists d(CH₂)₅Tyr(alk)VAVP (where alk = Me, Et),^{1,2} position 4 can tolerate a wide variety of structural changes without loss of antidiuretic antagonism. This is all the more surprising in view of our earlier finding that Val⁴ plays a critical role in endowing d(CH₂)₅Tyr(Et)VAVP with antiantidiuretic properties.^{1,2,6,7} Short-chain and β-branched aliphatic amino acids at position 4 in d(CH₂)₅-D-Phe²X⁴AVP are clearly superior to either long-chain or

γ-branched aliphatic amino acids in leading to enhancement of antiantidiuretic potencies. The substitution of aromatic amino acids at position 4 leads to drastic losses of antiantidiuretic antagonism. Five of the new antagonists, d(CH₂)₅-D-Phe²Ile⁴AVP, d(CH₂)₅-D-Phe²Abu⁴AVP, d(CH₂)₅-D-Phe²Ala⁴AVP, d(CH₂)₅-D-Phe²AVP, and d(CH₂)₅-D-Phe²Lys⁴AVP, exhibit no evident antidiuretic agonistic activity. With an antiantidiuretic pA₂ = 8.24, d(CH₂)₅-D-Phe²Ile⁴AVP appears to be the most potent antidiuretic antagonist reported to date. It also exhibits enhanced antiantidiuretic/antivasopressor selectivity relative to d(CH₂)₅-D-Phe²VAVP. The findings reported here on the structural flexibility at position 4 in d(CH₂)₅-D-Phe²X⁴AVP analogues, together with our earlier findings that position 2 in d(CH₂)₅X²VAVP can tolerate a wide variety of D-amino acids with retention of antidiuretic antagonism,³⁻⁵ offer very valuable clues for the design of even more potent and/or selective antidiuretic antagonists. Some of the highly potent antidiuretic antagonists reported here, in particular, d(CH₂)₅-D-Phe²Ile⁴AVP and d(CH₂)₅-D-Phe²Abu⁴AVP, are potentially useful as pharmacological tools for studies on AVP-induced water-retention states and might also be of therapeutic value for the treatment of AVP-induced hyponatremias of diverse etiologies, as, for example, in the Schwartz-Bartter syndrome.²³ The prospect of probing the structural characteristics of AVP renal and vascular receptors using d(CH₂)₅-D-Phe²Lys⁴AVP, and related position-4-modified analogues possessing other active groups, as covalent ligands is now also a distinct possibility.

Experimental Section

The procedure of "solid-phase" synthesis followed that previously published.⁸⁻¹¹ Chloromethylated resin (Chemalog, 1% cross-linked S-DVB, 200-400 mesh, 0.75-1.00 mequiv/g) was esterified with Boc-Gly by the cesium salt method²⁴ to an incorporation of 0.5 mmol/g. Amino acid derivatives were supplied by Bachem Inc. or Chemalog Inc. *p*-Nitrophenyl β-(benzylmercapto)-β,β-cyclopentamethylene propionate¹² was synthesized. Dimethylformamide (DMF) was distilled under reduced pressure. Other solvents and reagents were analytical grade. Thin-layer chromatography (TLC) was on silica gel (0.25 mm, Brinkmann Silplate). The following solvent systems were used: (A) butan-1-ol-acetic acid-water (4:1:1, v/v); (B) chloroform-methanol (7:3, v/v); (C) butan-1-ol-acetic acid-water-pyridine (15:3:3:10, v/v); (D) butan-1-ol-acetic acid-water (4:1:5, v/v, upper phase). Loads

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Table III. Physicochemical Properties of the Protected Peptides:
 β -(Benzylmercapto)- β -cyclopentamethylteneproylyl-D-Phe-Phe-X-Asn-Cys(Bzl)-Pro-Arg(Tos)-Gly-NH₂ (I-XIV)

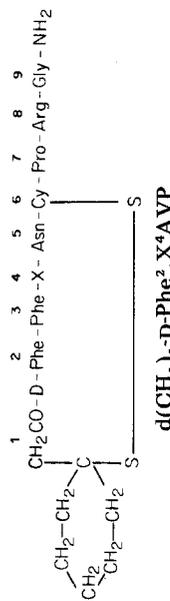
no.	X	formula	wt, mg	yield, %	mp, °C	[α] _D ²⁵ , deg	TLC				amino acid analyses ratios						
							R _f (A)	R _f (B)	R _f (C)	Phe	Asp	Cys(Bzl)	Pro	Arg	Gly	NH ₃	X
I	Ile	C ₇₃ H ₉₅ N ₁₃ O ₁₂ S ₃ ·2H ₂ O	380	65.9	211-213	-27.1 ^a	0.52	0.77	0.80	2.11	1.07	1.08	0.95	0.96	1.00	1.89	0.96
II	Abu	C ₇₁ H ₉₁ N ₁₃ O ₁₂ S ₃ ·2H ₂ O	510	72.0	205-206	-12.6 ^b	0.50	0.94	0.81	2.03	1.07	1.02	1.03	0.87	1.00	1.92	0.95
III	Thr(Bzl)	C ₇₈ H ₉₇ N ₁₃ O ₁₃ S ₃ ·2H ₂ O	340	57.5	170-173	-9.7 ^b	0.60	0.98	0.85	1.86	1.06	0.98	0.88	1.08	1.00	1.87	1.19
IV	Ala	C ₇₀ H ₈₉ N ₁₃ O ₁₂ S ₃ ·3H ₂ O	350	62.0	186-188	-13.3 ^b	0.52	0.68	0.80	2.01	0.96	1.07	1.03	0.98	1.00	2.06	0.98
V	Gln	C ₇₂ H ₉₂ N ₁₄ O ₁₃ S ₃ ·H ₂ O	430	59.1	185-187	-17.6 ^a	0.34	0.67	0.77	1.98	1.02	1.07	0.97	1.09	1.00	2.88	1.01
VI	Lys(Tos)	C ₈₀ H ₁₀₂ N ₁₄ O ₁₄ S ₃ ·2H ₂ O	410	63.0	163-165	-19.5 ^b	0.54	0.83	0.79	1.96	0.98	1.02	0.89	0.99	1.00	2.05	1.05
VII	Cha	C ₇₉ H ₉₈ N ₁₃ O ₁₂ S ₃ ·2H ₂ O	420	71.0	194-195	-19.2 ^b	0.51	0.84	0.91	2.01	1.02	1.13	0.87	1.12	1.00	1.96	1.17
VIII	Nle	C ₇₃ H ₉₅ N ₁₃ O ₁₂ S ₃ ·2H ₂ O	390	67.5	207-208	-19.4 ^b	0.49	0.67	0.84	1.98	1.03	1.11	0.98	1.04	1.00	1.96	0.87
IX	Nva	C ₇₁ H ₉₅ N ₁₃ O ₁₂ S ₃ ·2H ₂ O	340	60.2	206-207	-17.6 ^a	0.55	0.90	0.87	2.01	1.07	0.98	1.02	0.97	1.00	1.82	0.98
X	Phe	C ₇₆ H ₉₅ N ₁₃ O ₁₂ S ₃	370	62.0	200-202	-27.1 ^a	0.55	0.70	0.84	2.02	1.07	1.14	0.89	1.06	1.00	1.82	1.01
XI	Leu	C ₇₃ H ₉₅ N ₁₃ O ₁₂ S ₃	380	66.7	210-211	-24.4 ^b	0.56	0.69	0.79	2.01	0.96	0.98	0.90	1.07	1.00	1.93	1.08
XII	Gly	C ₆₉ H ₈₇ N ₁₃ O ₁₂ S ₃ ·2H ₂ O	433	77.5	134-137	-20.3 ^b	0.47	0.50	0.80	1.94	0.97	0.98	0.96	0.95	1.00	1.89	1.00
XIII	Tyr(Bzl)	C ₈₃ H ₉₉ N ₁₃ O ₁₃ S ₃ ·2H ₂ O	410	64.7	199-202	-22.5 ^b	0.59	0.92	0.85	1.98	1.10	1.03	0.98	0.93	1.00	2.03	0.96
XIV	Pro	C ₇₂ H ₉₁ N ₁₃ O ₁₂ S ₃ ·2H ₂ O	380	67.5	134 dec	-30.1 ^b	0.44	0.89	0.81	1.93	1.03	1.03	0.99	0.97	1.00	1.87	1.00

^a c 0.9, DMF. ^b c 1, DMF.

Table IV. Physicochemical Properties of d(CH₂)₅-D-Phe²-X⁴-AVP Antagonists 1-14

no.	peptide ^a	X ⁴	wt, mg	yield, %	[α] _D ²⁵ , deg	TLC				amino acid analyses ratios						
						R _f (A)	R _f (C)	R _f (D)	Phe	Asp	1/2Cys	Pro	Arg	Gly	NH ₃	X ⁴
1	d(CH ₂) ₅ -D-Phe ² -Ile ⁴ -AVP	Ile	35	38.1	-116.4 ^b	0.33	0.58	0.39	2.10	1.06	0.92	0.92	0.94	1.00	1.94	0.92
2	d(CH ₂) ₅ -D-Phe ² -Abu ⁴ -AVP	Abu	40.2	43.1	-106.8 ^f	0.28	0.53	0.35	2.08	0.99	1.03	1.10	1.00	1.00	1.86	0.99
3	d(CH ₂) ₅ -D-Phe ² -Thr ⁴ -AVP	Thr	46.1	53.3	-118.3 ^c	0.22	0.48	0.34	1.94	0.94	0.92	0.98	0.99	1.00	1.86	0.92
4	d(CH ₂) ₅ -D-Phe ² -Ala ⁴ -AVP	Ala	46.5	51.1	-118.2 ^b	0.21	0.50	0.35	1.97	0.94	0.93	1.02	0.91	1.00	1.92	1.01
5	d(CH ₂) ₅ -D-Phe ² -Gln ⁴ -AVP	Gln	62.7	66.4	-136.4 ^c	0.13	0.40	0.32	2.01	1.03	1.11	0.98	1.09	1.00	1.90	1.00
6	d(CH ₂) ₅ -D-Phe ² -Lys ⁴ -AVP	Lys	41	49.4	-102.5 ^b	0.02	0.19	0.28	1.95	0.93	0.97	1.10	0.94	1.00	1.83	0.98
7	d(CH ₂) ₅ -D-Phe ² -Cha ⁴ -AVP	Cha	8.9	9.6	-70.9 ^d	0.36	0.62	0.41	1.89	1.01	0.92	0.89	0.97	1.00	2.00	1.18
8	d(CH ₂) ₅ -D-Phe ² -Nle ⁴ -AVP	Nle	24.2	26.4	-121.0 ^c	0.34	0.58	0.39	1.95	0.95	0.90	0.90	0.99	1.00	1.91	0.96
9	d(CH ₂) ₅ -D-Phe ² -Nva ⁴ -AVP	Nva	44	50.3	-101.8 ^c	0.32	0.57	0.38	1.94	0.94	1.03	0.91	0.98	1.00	1.95	0.96
10	d(CH ₂) ₅ -D-Phe ² -Phe ⁴ -AVP	Phe	21	22.8	-106.6 ^c	0.33	0.55	0.38	1.90	0.95	0.99	0.95	0.98	1.00	1.99	0.97
11	d(CH ₂) ₅ -D-Phe ² -Leu ⁴ -AVP	Leu	20.1	21.9	-114.0 ^f	0.34	0.59	0.39	2.13	0.93	0.94	0.87	1.09	1.00	1.95	1.00
12	d(CH ₂) ₅ -D-Phe ² -Gly ⁴ -AVP	Gly	46.2	51.2	-100.0 ^g	0.19	0.47	0.34	1.81	0.87	0.96	0.98	0.96	1.00	1.88	1.00
13	d(CH ₂) ₅ -D-Phe ² -Tyr ⁴ -AVP	Tyr	11.6	13.2	-58.8 ^f	0.31	0.53	0.37	1.82	0.89	1.12	0.92	0.98	1.00	1.84	0.93
14	d(CH ₂) ₅ -D-Phe ² -Pro ⁴ -AVP	Pro	34.4	38.9	-125.6 ^c	0.16	0.46	0.33	1.87	0.92	0.88	0.96	0.95	1.00	2.10	0.92

^a Abbreviations as in Table I. ^b c 0.4, 1 N AcOH. ^c c 0.3, 1 N AcOH. ^d c 0.11, 1 N AcOH. ^e c 0.15, 1 N AcOH. ^f c 0.25, 1 N AcOH. ^g c 0.5, 1 N AcOH.



of 10–50 μg were applied, and chromatograms were a minimum length of 10 cm. Iodine vapor was used for detection. For amino acid analysis,²⁵ peptides (~ 0.7 mg) were hydrolyzed with constant-boiling hydrochloric acid (400 μL) containing phenol (10 μL) in evacuated and sealed ampules for 18 h at 118 $^{\circ}\text{C}$. The analyses were performed on Models 121M and 119CL Beckman automatic amino acid analyzers. Molar ratios were referred to Gly (= 1.00). The cysteine content of the free peptides was estimated as $^{1/2}$ -cystine. Elemental analyses were performed by Integral Microanalytical Laboratories, Inc, Raleigh, NC. The analytical results for elements indicated by their symbols were within 0.4% of theoretical values. Optical rotations were measured with a Rudolph polarimeter Model 80.

Boc-Asn-Cys(Bzl)-Pro-Arg(Tos)-Gly-resin (A). The protected pentapeptide resin A (27.15 g, 10 mM) was prepared from 20 g (10 mM) of Boc-Gly-resin by the solid-phase peptide synthesis methodology,^{8–11} i.e., four cycles of deprotection with 1 M HCl/HOAc, neutralization with 10% Et_3N in MeCl_2 , and coupling with Boc-Arg(Tos), Boc-Pro, Boc-Cys(Bzl) mediated by DCC and Boc-Asn NPE (with HOBT as additive), respectively.

β -(Benzylmercapto)- β,β -cyclopentamethylenepropionyl-D-Phe-Phe-Ile-Asn-Cys(Bzl)-Pro-Arg(Tos)-Gly-NH₂ (I). The pentapeptidyl resin (A; 1.1 g, 0.4 mmol) was converted to protected acyl octapeptidyl resin in four cycles of solid-phase peptide synthesis by using as the carboxy component Boc-L-Ile, Boc-L-Phe, Boc-D-Phe, and *p*-nitrophenyl β -(benzylmercapto)- β,β -cyclopentamethylenepropionate,¹² respectively, the latter coupling being facilitated by the addition of *N*-hydroxybenzotriazole (HOBT).¹⁵ The protected acyl octapeptidyl resin was ammonolyzed.¹⁰ The crude product was extracted with hot (60 $^{\circ}\text{C}$) DMF, precipitated by the addition of water, collected, and dried. It was taken up in hot DMF and reprecipitated with ethanol-ethyl ether to give the required protected peptide amide I (Table III).

β -(Benzylmercapto)- β,β -cyclopentamethylenepropionyl-D-Phe-Phe-X-Asn-Cys(Bzl)-Pro-Arg(Tos)-Gly-NH₂ [II–XIV, X = Abu (II), Thr(Bzl) (III), Ala (IV), Gln (V), Lys(Tos) (VI), Cha (VII), Nle (VII), Nva (IX), Phe (X), Leu (XI), Gly (XII), Tyr(Bzl) (XIII), and Pro (XIV)]. Each of the protected peptides II–XIV was synthesized from the pentapeptidyl resin (A; 1.1 g, 0.4 mmol) in the same manner as for I, except that in the first coupling, Boc-L-Ile was replaced by the Boc derivative of X in each case. The physicochemical properties of all of the protected peptides I–XIV are given in Table III.

[1-(β -Mercapto- β,β -cyclopentamethylenepropionic acid)-2-D-phenylalanine,4-isoleucine]arginine-vasopressin [$\text{d}(\text{CH}_2)_5\text{-D-Phe}^2,\text{Ile}^4\text{AVP}$] (1). A solution of the protected acyl octapeptide amide I (120 mg, 0.083 mmol) in sodium-dried and redistilled ammonia (500 mL) was treated at the boiling point and with stirring with sodium¹⁶ from a stick of the metal contained in a small-bore glass tube until a light-blue color persisted in the solution for 30 s. Dry acetic acid (0.4 mL) was added to discharge the color. The ammonia was evaporated, and nitrogen was passed

through the flask. After 5 min, the residue was dissolved in degassed aqueous acetic acid (20%, 50 mL) and quickly poured into ice-cold water (~ 1500 mL).² The pH was adjusted to ~ 7.0 with concentrated ammonium hydroxide. Following the neutralization, an excess of a solution of potassium ferricyanide (0.01 M, 15 mL)¹⁷ was added gradually with stirring. The yellow solution was stirred for an additional 20 min and for 10 min with anion-exchange resin (Bio-Rad AG-3, Cl^- forms, 40-g damp weight). The suspension was slowly filtered through a bed of resin (40 g damp weight). The bed was washed with water (100 mL), and the combined filtrate and washings were lyophilized. The resulting power (1.9 g) was desalted on a Sephadex G-15 column (110 \times 2.7 cm) eluting with aqueous acetic acid (50%)¹⁸ with a flow rate of 5 mL/h. The eluate was fractioned and monitored for absorbance at 254 nm. The fractions comprising the major peak were checked by TLC (A), pooled, and lyophilized, and the residue was further subjected to gel filtration on a Sephadex G-15 column (100 \times 1.5 cm) eluting with aqueous acetic acid (0.2 M)¹⁸ with a flow rate of 4 mL/h. The peptide was eluted in a single peak (absorbance 254 nm). Lyophilization of the pertinent fractions gave the vasopressin analogue 1 as a white powder (Table IV).

[1-(β -Mercapto- β,β -cyclopentamethylenepropionic acid), 2-D-phenylalanine,4-amino acid X]arginine-vasopressins [$\text{d}(\text{CH}_2)_5\text{-D-Phe}^2,\text{X}^4,\text{AVP}$ (2–14), X = Abu (2), Thr (3), Ala (4), Gln (5), Lys (6), Cha (7), Nle (8), Nva (9), Phe (10), Leu (11), Gly (12), Tyr (13), Pro (14)]. Each of these cyclized free peptides (2–14) was obtained from their respective protected precursors II–XIV by the procedure followed in obtaining the free peptide 1 from the protected intermediate I. Their physicochemical properties are given in Table IV. The pharmacological properties of peptides 1–14 are presented in Tables I and II.

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