# Pharmacological characterization of angiotensin II AT<sub>2</sub> receptor subtype heterogeneity in the rat adrenal cortex and medulla

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Adrenal angiotensin II (AII) receptors have been pharmacologically and structurally divided into two main subtypes, AT<sub>1</sub> and AT<sub>2</sub>. Radioligand receptor binding assays with <sup>125</sup>Isarcosine<sup>1</sup>, isoleucine<sup>8</sup> angiotensin II (125I-SI AII) in the presence of losartan, an AT1 selective ligand, and PD123177 an AT2 selective ligand, indicated that the AT<sub>1</sub> subtype was predominant in membrane homogenates of the rat adrenal cortex (AT<sub>1</sub> Bmax =  $649 \pm 62$  fmol/mg protein; AT<sub>2</sub> Bmax =  $237 \pm 29$  fmol/ mg protein). In membrane homogenates of the adrenal medulla, the AT<sub>2</sub> subtype was predominant (AT<sub>1</sub> Bmax =  $55 \pm 5$  fmol/mg protein; AT<sub>2</sub> Bmax =  $109 \pm 29$  fmol/mg protein). Overall 58% of the 125I-SI All binding in the rat adrenal was to the AT1 subtypes, and 42% was to the AT2 subtypes. The outer cortex contained 59% of the All receptor binding sites in the adrenal, while the medulla accounted for the remaining 41%. The affinity of the AT<sub>1</sub> binding sites in membrane homogenates of the cortex and medulla ( $K_D=672\pm123~pM$  and  $573\pm85~pM$ , respectively) was not significantly different. The affinity for <sup>125</sup>I-SII All of AT<sub>2</sub> binding sites in membrane homogenates was higher than that of AT, binding sites. The affinity for 1251-SI All of AT, binding sites in membrane homogenates of the outer cortex ( $K_D = 265 \pm 35 \text{ pM}$ ) was significantly less than that in the medulla ( $K_D = 133 \pm$ 11 pm). In vitro receptor autoradiography also demonstrated that the AT2 subtype in frozen sections of the cortex had a lower affinity  $(K_D = 1512 \pm 191 \text{ pM})$  than that in the medulla  $(K_D =$ 867  $\pm$  72 pm). The heterogeneous affinity of adrenal AT<sub>2</sub> binding sites may indicate existence of multiple AT2 receptor subtypes in the rat adrenal.

**Keywords:** angiotensin II (All) receptors; AT<sub>1</sub>; AT<sub>2</sub>; binding affinity; rat; adrenal cortex; adrenal medulla; autoradiography

#### Introduction

Angiotensin II (AII) binds to specific receptors on cell surfaces to initiate a variety of physiological effects including arteriolar vasoconstriction (Braun-Menendez et al., 1940; Page & Helmer, 1940), aldosterone secretion (Laragh et al., 1960), catecholamine release (Zimmerman & Whitmore, 1967; Hughes & Roth, 1971), prolactin (Canonico & Mac-Leod, 1986) and vasopressin (Keil et al., 1975) secretion and induction of drinking behavior (Booth, 1968; Andersson & Westbye, 1970; Epstein et al., 1970). This multiplicity of actions suggests a complex interaction of AII with single or multiple receptors. Recently, two AII receptor subtypes were characterized in peripheral tissue and in the brain on the basis of their sensitivity to sulfhydryl reducing agents (Chiu et al., 1989b; Whitebread et al., 1989; Speth et al., 1991) and selective AII receptor antagonists (Chiu et al., 1989a; Whitebread et al., 1989; Rowe et al., 1990). AT<sub>1</sub> receptors are selectively blocked by losartan and a host of newer generation AT<sub>1</sub> selective antagonists (Timmermans et al., 1993). Ligands selective for AT<sub>2</sub> receptors include CGP42112 (Whitebread et al., 1989), PD123177 and its congeners (Chiu et al., 1989a; Dudley et al., 1990) and p-aminophenylalanine<sup>6</sup> AII (Speth & Kim, 1990). Both AII receptor subtypes were found by radioligand binding assays in the rat adrenal (Chiu et al., 1989a; Whitebread et al., 1989). The adrenal cortex contains mainly the AT<sub>1</sub> subtype, the adrenal medulla mainly the AT<sub>2</sub> subtype (Chiu et al., 1989a).

The AT<sub>1</sub> receptor was initially cloned from bovine adrenal glomerulosa and cultured rat vascular smooth muscle cells (Murphy et al., 1991; Sasaki et al., 1991). Subsequent genomic analysis and homology cloning revealed two AT<sub>1</sub> receptor isoforms in the rat and mouse, AT<sub>1a</sub> and AT<sub>1b</sub> (Elton et al., 1992; Iwai & Inagami, 1992; Kakar et al., 1992a; Sasamura et al., 1992). The isoforms have 95% identical amino acid sequence. Vascular smooth muscle and lung express primarily AT<sub>1a</sub> mRNA, whereas AT<sub>1b</sub> mRNA is primarily expressed in the adrenal and pituitary gland (Kakar et al., 1992b). The two AT<sub>1</sub> subtypes have similar ligand binding profiles (Chiu et al., 1993). The AT<sub>2</sub> receptor has also been cloned from rat fetus (Mukoyama et al., 1993) and the PC12W cell line (Kambayashi et al., 1993) with identical amino acid sequence and structure.

Tsutsumi and Saavedra (1992) reported heterogeneity of angiotensin II  $AT_2$  receptors in the rat brain on the basis of differential guanine nucleotide and pertussis toxin sensitivity. Speth (1993) also reported that <sup>125</sup>I-SI AII and <sup>125</sup>I CGP42112 binding to  $AT_2$  receptor binding sites in rat adrenal and brain displayed different responses to  $\beta$ -mercaptoethanol. These previous radioligand binding assays suggest the existence of an  $AT_2$  receptor heterogeneity. In this study, the binding characteristics of both major AII receptor subtypes in the rat adrenal cortex and medulla were compared using both homogenate binding assays and *in vitro* receptor autoradiography.

### Results

Saturation isotherm analyses and Rosenthal (Scatchard) transformations of 125I-SI AII binding to AT1 and AT2 AII receptor subtypes in membrane homogenate preparations of the outer cortex and inner/medullary regions are shown in Table 1 and Figures 1 and 2. As shown in Figure 1C, the concentration of AT<sub>1</sub> receptors in homogenates of the outer cortex is more than 10-fold higher than in homogenates of the inner/medullary region. The  $K_D$  of  $^{125}$ I-SI AII binding to AT<sub>1</sub> binding sites in these two regions of the rat adrenal is similar (Table 2). The concentration of AT<sub>2</sub> receptors in the outer cortex was also greater than that in the inner/medullary region by a factor of 2 (Figure 2). However, as shown in Figure 2C and Table 2, the K<sub>D</sub> of <sup>125</sup>I-SI AII for the cortical AT<sub>2</sub> receptors (265 ± 79 pM) was significantly greater (P < 0.01) than that for medullary AT<sub>2</sub> receptors  $(133 \pm 25 \text{ pM})$ . Analysis of variance of the  $K_D$  values revealed that 125I-AII bound to the adrenal AT2 receptors with a significantly higher affinity (F<sub>1,14</sub> = 118, P < 0.001) than to the adrenal AT<sub>1</sub> receptors.

The amounts of AII receptor subtypes and their proportion in different regions of the rat adrenal are described in Table 2. In the whole adrenal, AT<sub>1</sub> receptors, defined as maximal <sup>125</sup>I-SI AII binding capacity (Bmax) in the presence of 10<sup>-5</sup> M PD 123177, accounted for 59% of the total AII



Table 1 Comparison of 1251-SI AII binding to AT1 and AT2 receptors in adrenal cortex and medulla

	Adren	al cortex	Adrenal medulla <sup>a</sup>				
Membrane suspensions							
AT <sub>1</sub>	$K_D =$	$672 \pm 123$	$K_D = 573 \pm 85$				
•	Bmax =	$649 \pm 62$	$Bmax = 55 \pm 5$				
AT <sub>2</sub>	$K_D =$	$265 \pm 79 \dagger$	$K_D = 133 \pm 25*\dagger$				
		$237 \pm 29$	$Bmax = 109 \pm 29$				
	n = 5		n = 5				
Frozen sections							
AT,	$K_D =$	$1512 \pm 191$	$K_D = 867 \pm 72**$				
-	Bmax =	198 ± 13 <sup>6</sup>	$Bmax = 516 \pm 48^{b}$				
	n	· = 6	n=6				

Bmax is expressed as fmol/mg protein; KD is in units of pm. <sup>a</sup>Includes zona fasciculata and zona reticularis in membrane suspensions. The AT<sub>1</sub> subtype was assayed in the presence of 10<sup>-5</sup> M PD123177. The  $AT_2$  subtype was assayed in the presence of  $10^{-5}\,\mathrm{M}$ losartan. The values are expressed as mean  $\pm$  SE. \*P<0.05, \*\*P < 0.01, by paired t-test, significantly less than AT<sub>2</sub> in cortex.  $\dagger P < 0.001$ , by two-way ANOVA, significantly less than AT<sub>i</sub> bBmax values in frozen sections were derived from values of fmoles per g wet weight based on empirical determinations of the amount of protein per unit wet weight (178 mg kg<sup>-1</sup>) in whole adrenal homogenates

receptors. Seventy-five percent of these AT<sub>1</sub> receptors  $(286 \pm 22 \text{ fmoles})$  were located in the outer cortex, which encompasses the zona glomerulosa and the capsule, while 25% (96  $\pm$  6 fmoles) were present in the remaining regions of the adrenal. The AT<sub>2</sub> subtype of the AII receptors, defined as <sup>125</sup>I-SI AII binding in the presence of 10<sup>-5</sup> M losartan, accounted for 41% of the rat adrenal AII receptors. Thirtysix percent of these  $AT_2$  receptors (98  $\pm$  13 fmoles) were located in the outer cortex, while 64% (174  $\pm$  30 fmoles) were present in the inner regions of the adrenal. Nearly threefourths of the outer cortical AII receptors were of the AT<sub>1</sub> subtype, while nearly two-thirds of the inner/medullary AII receptors were of the AT2 subtype.

To confirm the distribution patterns and differential binding characteristics of 125 I-SI AII to AT2 receptors in the outer cortex and inner/medullary regions, in vitro receptor autoradiographic analyses were performed. As shown in Figure 3, 125I-SI AII binding to AT<sub>2</sub> receptors was localized to two regions: the zona glomerulosa, and the medulla. The concentration of AT2 receptors in the medulla only, in contrast to membrane homogenates which included the zona reticularis and zona fasciculata with the medulla, was more than twice that of the outer adrenal cortex (Table 2 and Figure 4). As seen in membrane homogenates, the K<sub>D</sub> of the AT2 receptors of the cortex for 125I-SI AII was again significantly greater (P < 0.01) than that for the AT<sub>2</sub> receptors in the medulla (Table 2 and Figure 4).

#### Discussion

To date, almost all known effects of AII in adult tissues are attributable to the AT<sub>1</sub> receptor through its ability to activate phospholipase C which results in mobilization of intracellular calcium and the activation of protein kinase C (PKC) or via the inhibition of adenylate cyclase activity (Murphy et al., 1991; Sasaki et al., 1991; Iwai & Inagami, 1992; Sasamura et al., 1992). On the other hand, no consistent physiological function has been demonstrated for the AT2 receptor. The AT2 receptor is abundantly and widely expressed in various fetal tissues (Grady et al., 1991), immature brain (Cook et al., 1991; Millan et al., 1991; Tsutsumi & Saavedra, 1991) and PC12W cell line (Speth & Kim, 1990; Webb et al., 1992; Brechler et al., 1993). However, this receptor is only sparsely expressed in the normal adult animal, primarily in the adrenal medulla (Chiu et al., 1989a,b), the uterine myometrium (Whitebread et al., 1989), the ovarian granulosa (Pucell

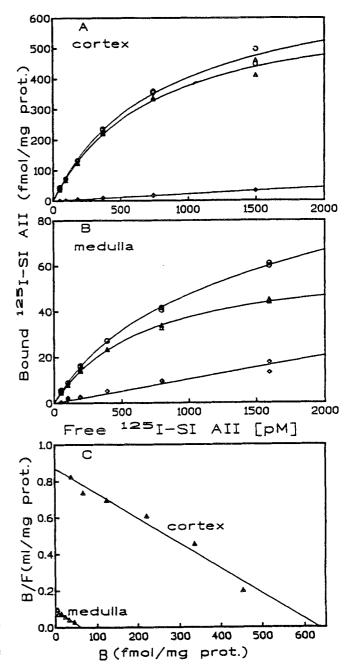
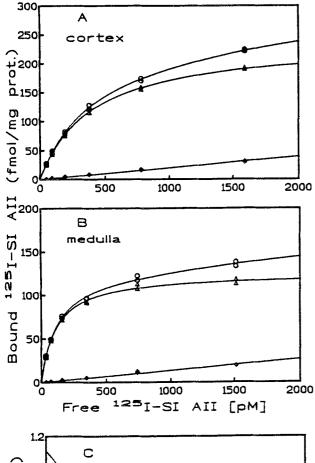


Figure 1 Saturation isotherm of <sup>125</sup>I-SI AII binding and linear transformation of specific <sup>125</sup>I-SI AII binding to AT<sub>1</sub> receptors in rat adrenal membrane suspensions in the presence of 10<sup>-5</sup> M PD123177. Samples were run in triplicate. Nonspecific binding was determined in the presence of  $3\,\mu\text{M}$  AII. Specific binding ( $\Delta$ ) was derived from the difference between total binding (O) and nonspecific binding (\$). The curves for total and specific binding were generated from the equation:  $Y = Bmax \cdot X/(K_D + X) + C \cdot X$  where Y = total bound,  $X = {}^{125}I$ -SI AII concentration, and C is the ratio of nonspecific binding. The derived values for this example were 646 fmol mg-1 protein for Bmax, 692 pm for KD in adrenal cortex (A) and 62 fmol/ mg protein for Bmax. 657 рм for KD in adrenal medulla (В). (С) shows Rosenthal (Scatchard) plots of the data derived from specific binding to the adrenal cortex in (A) (A) and the adrenal medulla in  $(\mathbf{B})(\Delta)$ 

et al., 1991) and central nervous system (Chang et al., 1990; Gehlert et al., 1990; Rowe et al., 1990; Millan et al., 1991). These patterns of AT<sub>2</sub> expression suggest that the AT<sub>2</sub> receptor may play a role in growth and development. However, this receptor does not appear to involve any known classical intracellular signalling pathways.





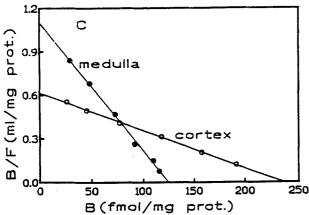


Figure 2 Saturation isotherm of 125I-SI AII binding and linear transformation of specific <sup>125</sup>I-SI AII binding to rat adrenal membrane suspensions in the presence of  $10^{-5}$  M losartan. Other details are as described in Figure 1. The derived values for this example were 238 fmol mg<sup>-1</sup> protein for Bmax and 393 pM for K<sub>D</sub> in adrenal cortex (A) and 129 fmol mg<sup>-1</sup> protein for Bmax and 115 pm for  $K_D$ in adrenal medulla (B). (C) shows Rosenthal (Scatchard) plots of the data derived from specific binding shown in (A) and (B)

Studies of AT<sub>2</sub> signal transduction mechanisms are controversial regarding whether or not the stimulation of AT<sub>2</sub> receptor by AII results in a reduction in intracellular cGMP and an increase in phosphotyrosine phosphatase (PTP) activity (Dudley et al., 1991; Pucell et al., 1991; Bottari et al., 1992; Webb et al., 1992; Brechler et al., 1993). Bottari and his associates (Bottari et al., 1992) postulated a dephosphorylation of phosphotyrosine in particulate guanylate cyclase by AT<sub>2</sub> receptor activated PTP as the cause of decline in cGMP. Sumners et al. (1991) demonstrated a reduction in cGMP in AII-treated rat neuronal cells that was antagonized by CGP 42112. However, subsequent studies by Brechler et

Table 2 The proportions and amounts of AII receptor subtypes in the rat adrenal

	Cortex		Мес	dulla	Total			
	Binding (fmol)	Percentage (%)	Binding (fmol)	Percentage (%)	Binding (fmol)	Percentage (%)		
AT <sub>1</sub>	286 ± 22	74	96 ± 6	36	382	58		
$AT_2$	$98 \pm 13$	26	174 ± 30	64	272	42		
Total	384	100	270	100	654	100		

The values for binding above are expressed as means ± s.e.m. (fmoles per adrenal) of five experiments from 39 rats. The adrenal cortex preparations used in this study contained an average of 37 mg membrane protein per g wet weight tissue, the adrenal medulla preparations contained an average of 57 mg membrane protein per g wet weight tissue. When multiplied by the average wet weight of the subdivisions, there was 0.41 mg membrane protein-adrenal cortex and 1.68 mg membrane protein per adrenal medulla. aincludes zona fasciculata and zona reticularis in membrane suspensions. The AT<sub>1</sub> subtype was assayed in the presence of 10<sup>-5</sup> M PD123177. The AT<sub>2</sub> subtype was assayed in the presence of 10-5 M losartan

al. (1993) indicate that CGP 42112 may be an agonist. Kambayashi et al. (1993) reported that CGP42112 inhibited PTP activity in the plasma membranes of PC12W cells, rat fetal skin, and COS-7 cells stably transfected with the AT<sub>2</sub> cDNA, respectively. These discrepancies make is difficult to resolve the biochemical mechanism(s) of AT<sub>2</sub> receptors at this

A sampling of previous studies of rat adrenal AII receptor binding are summarized in Table 3. Early studies (Glossman et al., 1974; Devynck et al., 1977) suggested the presence of subtypes of All receptors based upon the presence of different binding affinities for the radioligand. A subsequent study (Singh et al., 1986) revealed a small, but statistically significant higher  $K_D$  for <sup>125</sup>I-AII in the rat adrenal cortex compared to the medulla. It was not until relatively recently however, that AII receptors were formally differentiated into AT<sub>1</sub> and AT<sub>2</sub> subtypes based upon pharmacological studies with subtype selective antagonists and agonists (Chiu et al., 1989a; Whitebread et al., 1989; Chang & Lotti, 1990; Dudley et al., 1990; Balla et al., 1991). In view of the observations that the heptapeptide AIII has a higher affinity for the AT<sub>2</sub> receptor (Timmermans et al., 1991), it seems likely that the high affinity <sup>3</sup>H-AIII binding site in the rat adrenal observed by Devynck et al. (1977) was the AT<sub>2</sub> subtype, and that the lower affinity <sup>3</sup>H-AIII binding site was the AT<sub>1</sub> receptor. However, some studies of adrenal AII receptors using the agonist radioligands <sup>125</sup>I-AII and <sup>125</sup>I-AIII (Himeno et al., 1988) and the antagonist radioligand 125I-SI AII have not revealed differential binding affinities (Balla et al., 1991; Aguilera, 1992) except in the presence of a sulfhydrylreducing agent (Aguilera, 1992).

This study used the peptide antagonist radioligand 125I-SI AII to identify and characterize AII receptor subtype binding sites in the rat adrenal cortex and medulla. Consistent with previous studies (Chiu et al., 1989a; Chang & Lotti, 1990; Balla et al., 1991; Wiest et al., 1991; Lu et al., In press), both AT<sub>1</sub> and AT<sub>2</sub> receptors were observed in the adrenal. However, previous studies of rat adrenal AII receptors have not examined the density and binding affinity of the individual All receptor subtypes in both the cortex and medulla. For the purposes of this study, AT<sub>1</sub> receptor binding was defined as specific (3 µM AII displaceable) 125 I-SI AII binding that occurred in the presence of the AT<sub>2</sub> selective, putative antagonist, PD 123177, at a concentration (10 µm) sufficient to saturate AT<sub>2</sub> receptors, with a negligible effect on adrenal AT<sub>1</sub> receptors (Lu et al., in press). AT<sub>2</sub> receptor binding was defined as specific (3 µM AII displaceable) 125I-SI AII binding that occurred in the presence of the AT1 selective antagonist, losartan, at a concentration (10  $\mu M$ ) sufficient to saturate  $AT_1$ receptors, with a negligible effect on adrenal AT<sub>2</sub> receptors (Lu et al., in press).

Estimates of the percentage of AII receptors in the rat

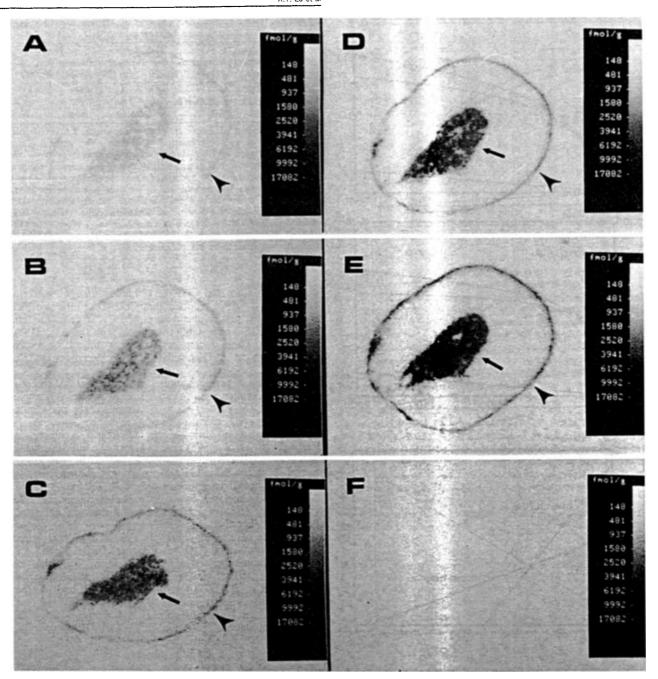


Figure 3 Computer analysed images of autoradiograms of <sup>125</sup>I-SI AII binding to adjacent adrenal sections incubated with five different concentrations; 91.8, 241, 485, 943, 2351 pm of <sup>125</sup>I-SI AII in the presence of 10<sup>-5</sup> M losartan, (A-E) of I-SI AII, respectively. (F) represents non-specific <sup>125</sup>I-SI AII binding at a concentration of 485 pm in the presence of 3 μm AII. The standard values in the graphs are for <sup>125</sup>I in units of fmoles/g wet weight. Since nonradioactive <sup>127</sup>I-SI AII was added at a ratio of 9:1 relative to <sup>125</sup>I-SI AII, the standard values on the right side of each panel should be multiplied by 10, the isotopic dilution factor, to indicate fm per g wet weight of I-SI AII binding. To derive fmoles of I-SI AII binding per mg protein this value should be divided by the average protein concentration per gram wet weight (178 mg protein per g wet weight). For this experiment, K<sub>D</sub> and Bmax were 1056 pm and 226 fmoles mg<sup>-1</sup> protein in the adrenal cortex and 639 pm and 417 fmoles mg<sup>-1</sup> protein in the adrenal medulla. The arrow heads point to the adrenal cortex, the arrows point to the adrenal medulla

adrenal cortex and medulla, based on the density of receptors per unit tissue, are about 90% and 10%, respectively (Catt et al., 1984; Singh et al., 1986). However, such estimates do not take into account the relative proportion of the adrenal that is cortex and that which is medulla. By multiplying the density of the receptor binding for each AII receptor subtype in the adrenal cortex and medulla (Table 1) by the amount of each tissue present in the adrenal, a value for the total number of binding sites for each AII receptor subtype binding sites per adrenal subdivision was obtained (Table 2). When compared in this manner, the proportion of total AII

receptors in the adrenal cortex and medulla was much closer, 59% in cortex and 41% in the medulla.

Examination of adrenal AII receptor binding using *in vitro* receptor autoradiography revealed the explanation for the apparent low density of AII receptors in the medulla. The adrenal dissection separated the capsule and outer cortex of the adrenal from the medulla and the two inner layers of the cortex, the zona reticularis and the zona fasciculata. However as can be seen in Figure 3, negligible amounts of specific <sup>125</sup>I-SI AII binding were present in the two inner layers of the cortex in the rat. Only the medulla and the outer cortex

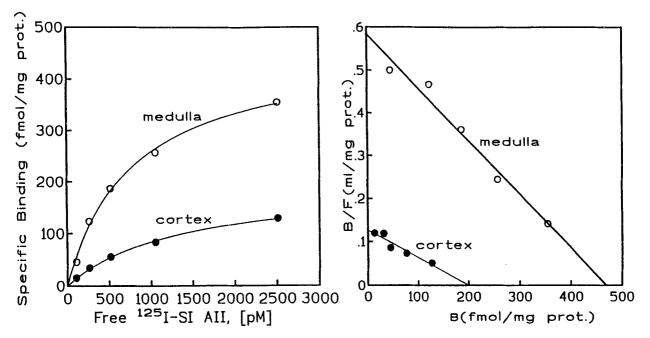


Figure 4 Saturation isotherm and linear transformation of specific 125I-SI AII binding to adrenal sections in the presence of 10-5 M losartan. The curves shown in (A) were generated from the equation:  $Y = Bmax X/(K_D + X)$  where Y = specific bound and  $X = {}^{125}I$ -SI AII concentration. (B) shows Rosenthal (Scatchard) plot of the data from (A). For this experiment,  $K_D$  and Bmax were 1600 pm and 198 fmoles mg<sup>-1</sup> protein in the adrenal cortex and 830 pm and 471 fmoles mg<sup>-1</sup> protein in the adrenal medulla

Table 3 Comparison of affinity of the binding sites for AII and its analogs

Rat tissue	Radioligand	$K_{DI}$	$K_{D2}$	Reference
Adrenal fraction (5 mm DTT)	125I-AII	0.2-0.5 пм	6-7 пм	Glossmann et al., 1974
Adrenal fraction (no DTT)	<sup>3</sup> H-AII	2.8 nm	25 nm	Devynck et al., 1977
	³H-AIII	0.15 пм	3.6 пм	
Adrenal cortex (1 mm DTT)	<sup>125</sup> I-AII	393 рм		Singh et al., 1986
medulla (1 mm DTT)	<sup>125</sup> I-AII	228 рм		
Adrenal cortex (no DTT)	<sup>125</sup> I-AII	860 рм		Himeno et al., 1988
	<sup>125</sup> I-AIII	900 рм		
medulla (no DTT)	<sup>125</sup> I-AII	1460 рм		
	<sup>125</sup> I-AIII	1160 рм		
Adrenal cortex (no DTT)	<sup>125</sup> I-SI AII	2.8 nm (IC <sub>50</sub> 1	for SI AII)	Balla et al., 1991
medulla (no DTT)	<sup>125</sup> I-SI AII	2.6 nm (IC <sub>50</sub> 1	for SI AII)	
Adrenal glomerulosa (no DTT)	<sup>125</sup> I-SI AII	1.6 nm (AT <sub>1</sub> )	1.7 nm $(AT_2)$	Aguilera, 1992
(2 mm DTT)	125I-SI AII	1.8 nm (AT <sub>1</sub> )		

displayed 125I-SI AII binding. Thus the estimates of 125I-SI All binding density in membrane homogenates of the medulla were decreased due to dilution with these non-AII receptor-containing regions.

Consistent with previous observations (Chiu et al., 1989a), the percentage of the AT<sub>1</sub> receptor subtype in the adrenal cortex (74%) was greater than that of the AT<sub>2</sub> receptor subtype. And the percentage of AT<sub>1</sub> receptors in the medulla (36%) was less than that of the AT<sub>2</sub> receptors. For the whole adrenal, the percentage of AT<sub>1</sub> receptors (58%) was only slightly greater than the percentage of AT2 receptors (42%).

Evaluation of the binding affinity for 125 I-SI AII of the two different AII receptor subtypes in the adrenal revealed signifiant differences. The binding affinity of AT<sub>1</sub> receptors in adrenal membrane homogenates was less than that of the AT<sub>2</sub> receptors. This observation contrasts with previous observations of <sup>125</sup>I-SI AII selectivity for AII receptor subtypes. In the rat adrenal glomerulosa, no difference in 125I-SI All binding affinity to AT<sub>1</sub> and AT<sub>2</sub> receptors was observed in the absence of sulfhydryl-reducing agents (Aguilera, 1992). In the rat brain (Rowe et al., 1992), and from comparison of rat liver (AT<sub>1</sub>) and PC12W cells (AT<sub>2</sub>) (Speth & Kim, 1990), <sup>125</sup>I-SI AII binds with lower affinity to the AT<sub>2</sub> subtype than to the AT<sub>1</sub> subtype. The sulfhydryl-reducing agent βmercaptoethanol, which increases 125I-SI AII binding to AT2 receptors in the brain (Speth et al., 1991), did not alter 125I-SI AII binding affinity in the whole adrenal (Speth, 1993). But dithiothreitol, another sulfhydryl-reducing agent, increases <sup>125</sup>I-SI AII binding affinity to AT<sub>2</sub> receptors in the adrenal glomerulosa (Aguilera, 1992).

The binding affinity of AT<sub>2</sub> receptors for <sup>125</sup>I-SI AII in both membrane homogenates and slide-mounted frozen sections of the adrenal medulla was greater (lower K<sub>D</sub>) than that of the AT<sub>2</sub> receptors in the cortex. The observation of higher binding affinity for 125I-SI AII in the medulla vs the cortex is consistent with the observation by Singh et al. (1986) of a higher binding affinity for <sup>125</sup>I-AII in the adrenal medulla. That study used a low (1 mm) concentration of dithiothreitol, which diminishes ligand binding to AT<sub>1</sub> receptors (Chiu et al., 1989b; Whitebread et al., 1989). Thus it is likely that the adrenal preparation used by Singh et al. (1986) expressed primarily AT<sub>2</sub> receptor-binding. The difference in binding affinity for 125I-SI AII between homogenates and tissue sections, shown in Table 1, is consistent with previous observations (Lu et al., in press) indicating that ligand binding affinity of AII receptors is decreased by the procedures required to prepare sections for in vitro receptor autoradiography.

The observation of different binding affinities for 125I-SI

AII at putative  $AT_2$  receptor binding sites in the adrenal cortex and medulla may be suggestive of AT<sub>2</sub> receptor heterogeneity. Previous radioligand binding studies of AT<sub>2</sub> receptors in the rat brain (Tsutsumi & Saavedra, 1992; Speth, 1993) also suggest the presence of AT<sub>2</sub> subtype heterogeneity. Of note, Balla et al. (1991) observed a twofold difference in PD123177 and a threefold difference in CGP42112 binding affinity for AT<sub>2</sub> receptors in the rat adrenal cortex and medulla. In contrast, comparisons of PD123177 binding affinity of AT2 receptors in the adrenal cortex and medulla carried out in this laboratory (Lu et al., in press) did not reveal substantial differences. The existence of two  $AT_1$  receptor subtypes,  $AT_{1a}$  and  $AT_{1b}$  in the rat has been confirmed by expression cloning studies (Elton et al., 1992; Iwai & Inagami, 1992). However, expression cloning studies of rat AT<sub>2</sub> receptors from different laboratories (Kambayashi et al., 1993; Mukoyama et al., 1993) have not revealed differences in the genes encoding the amino acid sequence of the receptor from two divergent types of tissue. The gene for the human AT<sub>2</sub> receptor shows a high degree of similarity to the rat and mouse AT, receptor gene Tsuzuki et al., 1994). In addition, the fetal mouse appears to possess only one AT<sub>2</sub> receptor gene (Nakajima et al., 1993). It is possible that the AT<sub>2</sub> receptors in different tissues follow different routes of post-translational processing, leading to subtle changes in the ligand binding characteristics. This scenario, which is well characterized for different neuropeptide transmitters (Mains et al., 1977; Amara et al., 1982), has not been established as a cause of receptor subtyping. However, Barker et al. (1993) observed four different isoforms of AII receptors in the rat adrenal using isoelectric focusing, suggesting that the differences could reflect different post-translational processing. Recently, Servant et al. (1994) reported that the marked disparity between the sizes of AT2 receptors from different tissues was related to different degrees of N-glycosylation. This could also cause moderate differences in ligand binding characteristics. Another possibility is that specific physical characteristics of the cells lead to subtle variations in the ability of the AT<sub>2</sub> receptors to bind ligands, e.g. the relative state of reduction or oxidation of the sulfhydryl bonds between cysteine residues in the extracellular domain of the receptors. Further studies are needed to determine if the differences observed in this study have important physiological or pharmacological significance.

## Materials and methods

Tissues were obtained from male Sprague-Dawley rats obtained from the Laboratory Animal Resource Center at Washington State University. For homogenate binding studies, rats were sacrificed by decapitation. The tissues were removed and further dissected. The adrenal was freed of adjacent adipose tissue and the adrenal cortex, primarily the outer zona glomerulosa, was separated from the medulla which also included portions of the zona fasciculata and zona reticularis. The adrenals from 7-9 rats were pooled for each comparison and weighed.

The tissues were mechanically homogenized in hypotonic buffer (20 mm sodium phosphate, pH 7.1-7.2) using a mechanical homogenizer (Tissuemizer, Tekmar, Cincinnati, OH). The homogenates were then centrifuged at 48 000 g for 20 min at 4°C. The precipitated membranes were resuspended in assay buffer (50 mm sodium phosphate, 150 mm NaCl, 1 mm EDTA, 0.1 mm bacitracin, pH 7.1-7.2) and centrifuged again. The membranes were resuspended at a concentration of approximately 10 mg initial wet weight per ml.

For analysis of AII receptor density and radioligand binding affinity, six concentrations of <sup>125</sup>I-SI AII ranging from 0.1 to 2 nm were used. Losartan and PD123177 were added at 10 µM concentrations to determine AT<sub>2</sub> and AT<sub>1</sub> receptor binding, respectively. Based on previous studies (Chiu et al., 1989a; Speth & Kim, 1990, Lu et al., in press), this concentration of these highly AII receptor subtype selective nonpeptide ligands gives near complete inhibition of the one subtype, with negligible inhibition of the other subtype. Nonspecific binding of <sup>125</sup>I-SI AII was determined in the presence of 3  $\mu$ M AII. Incubations were terminated by filtration through glass fibre filters (no. 32, Schleicher and Schuell, Keene, NH), prerinsed with 0.1% bovine albumin. Filters were rinsed three times with 50 mm sodium-potassium phosphate buffer, pH 7.4, at 22°C. Filter bound radioligand was assayed by sodium iodide crystal gamma scintillation counting. Membrane protein was determined by the method of Lowry et al. (1951).

For receptor autoradiographic studies of adrenal AII receptor subtypes, the rats were anesthetized with Equithesin (Jensen-Salsbury, Kansas City, MO) at 0.4 ml per 100g body weight and perfused intracardially with chilled phosphate buffered saline (20 mm sodium phosphate, pH 7.2). The adrenal was cleared of adjacent adipose tissue and frozen in a -20°C freezer for less than 1 week. The adrenals were sectioned at a thickness of 20 microns in a cryostat, air-dried and again stored at  $-20^{\circ}$ C. On the day of assay, the slidemounted adrenal sections were thawed at 37°C for about 15 s and preincubated for 30 min in assay buffer. The sections were then incubated in fresh assay buffer with a mixture of <sup>125</sup>I-SI AII and <sup>127</sup>I-SI AII yielding a final concentration of I-SI AII ranging from 0.1 to 2.5 nM in the presence of 10<sup>-5</sup> M losartan for 2 h. Nonspecific binding of I-SI AII was determined in the presence of  $3 \mu M$  AII.

The ratio of <sup>125</sup>I-SI AII to <sup>127</sup>I-SI AII was 1 to 9. The use of non-radioisotopic I-SI AII was necessitated by the extremely high amount of binding that occurred at high radioligand concentrations, exceeding the concentrations of 125I contained in the standards used for densitometric calibrations. <sup>127</sup>I-SI AII was prepared as described for <sup>125</sup>I-SI AII, except that a trace of <sup>125</sup>I-SI AII was added to monitor the peak of mono 127I-SI AII eluting from the HPLC column. The concentration of 127I-SI AII was determined by competition assay with 125I-SI AII.

Upon completion of the incubation, the sections were then dipped in two changes of distilled water, rinsed for 1 min in five changes of assay medium and finally dipped in another two changes of distilled water. The sections were then dried under a cool stream of air, mounted on cardboard and exposed to sheet film (SB-5, Kodak) in X-ray cassettes for an appropriate duration of time to insure sufficient film exposure at the lowest radioligand concentration. A videobased image analysis system (MCID, Imaging Research) was used to quantitate film exposure in units of fmol of 125 I-SI All bound per g tissue based on a standard curve generated from known amounts of 125I (Microscales, Amersham Inc. Arlington Heights, IL).

Binding constants were calculated using a computer program (Inplot 4.0, GraphPad Software, San Diego, CA). The dissociation constant (KD) and receptor density (Bmax) for specific binding were determined by the equation:  $Y = A \cdot X/$  $(B + X) + C \cdot X$ , where A = Bmax,  $B = K_D$ , C is the proportion of added <sup>125</sup>I-SI AII that was bound nonspecifically. X is the free radioligand concentration, and Y is total binding. The value of C·X was subtracted from this equation to yield the specific binding curve. For some assays the data was analysed by the linear transformation of Rosenthal (1967):  $B/F = (-1/K_D)B + Bmax/K_D$ , where B = specific binding, F = free radioligand concentration (commonly referred to as the Scatchard plot).

Data are expressed as mean ± standard error of the mean. Comparison of specific binding and binding constants in the different tissues was performed using a two-way ANOVA and a two-tailed Student's t-test. The null hypothesis was rejected when P < 0.05.



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