# Pharmacological profile of GR117289 in vitro: a novel, potent and specific non-peptide angiotensin  $AT_1$  receptor antagonist

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<sup>1</sup> This paper describes the effects of GRI17289 (1-[[3-bromo-2-[2-(lH-tetrazol-5-yl)phenyl]-5-benzofuranyl]methyl]-2-butyl-4-chloro-lH-imidazole-5-carboxylic acid) at angiotensin receptors and binding sites in rabbit aorta, rat liver and bovine cerebellum preparations in vitro.

2 In rabbit isolated aortic strips, GR117289 (0.3, 1 and 3 nM) caused a concentration-related, insurmountable suppression of the concentration-response curve to angiotensin II (All). When the contact time was increased, a greater degree of antagonism of All was observed, suggesting that GRi17289 is slow to reach equilibrium. A  $pK_B$  of 9.8  $\pm$  0.1 was calculated for GR117289 after 3h incubation. GR117289 (1 $\mu$ M) did not affect contractile responses to phenylephrine or 5-hydroxytryptamine (5-HT) in the rabbit aorta.

<sup>3</sup> GRI 17289 (1 nM) alone caused <sup>a</sup> marked suppression and a slight rightward displacement of the All concentration-response curve. Co-incubation with the competitive, surmountable AT, receptor antagonist, losartan (10 nM, 100 nM and  $1 \mu$ M), resulted in a concentration-related upward and rightward displacement of the concentration-response curve to subsequently administered All. In separate experiments in which preparations were pre-incubated with GR117289 (1 nM), subsequent addition of losartan (1 $\mu$ M) for 2, 15 or 45 min caused a further, but similar, rightward displacement of the concentration-response curve to subsequently administered All with a time-dependent increase in the maximum response.

4 Suppression of All-induced contractile responses, caused by superfusion with GRI17289 (0.3, <sup>1</sup> or <sup>3</sup> nM) was not reversed by continuously washing the tissues for <sup>3</sup> h; in fact, the potency of GRI 17289 was slightly enhanced after this period.

5 In rat liver membranes, GR117289 was a potent competitor with  $[3H]-AII$  for AT<sub>1</sub> binding sites  $(pK_i = 8.7 \pm 0.1)$  but in bovine cerebellum membranes, it was a very weak competitor for AT<sub>2</sub> binding sites ( $pK_i$  < 6). Pre-incubation of rat liver membranes with GR117289 had little effect on its affinity  $(pK<sub>i</sub> = 9.1 \pm 0.21)$ , but increasing the concentration of bovine serum albumen in the assay buffer from 0.001% to 0.1% w/v decreased affinity ( $pK_i = 7.5 \pm 0.1$ ).

In saturation binding experiments in rat liver membranes, GR117289 (12 nM) increased the  $K_d$  of [<sup>3</sup>H]-AII from  $0.28 \pm 0.06$  nM to  $0.37 \pm 0.02$  nM, and decreased B<sub>max</sub> from  $10.0 \pm 0.1$  to  $5.6 \pm 0.1$ 0.3 fmol mg<sup>-1</sup> tissue. In other experiments, GR117289 (1  $\mu$ M) did not alter the rate of dissociation of  $[^3H]$ -AII from AT<sub>1</sub> binding sites, following addition of excess unlabelled AII.

In rabbit aorta vascular smooth muscle membranes, GR117289 competed with [<sup>125</sup>I]-Sar<sup>1</sup>Ile8 AII for binding to  $AT_1$  binding sites. In the presence of 0.1% w/v bovine serum albumen, a pIC<sub>50</sub> of 7.6 ± 0.1 was calculated. Under the same conditions, but with rat liver membranes, a pIC<sub>50</sub> of 7.8  $\pm$  0.1 was determined.

8 Taken together, these results show that GRI17289 is a potent, specific, selective and insurmountable antagonist at angiotensin AT, receptors. Its profile in the rabbit aorta is consistent with the proposal that GRI17289 is a slowly reversible (pseudo-irreversible) antagonist at these receptors.

Keywords: GR117289; angiotensin II (AII);  $AT_1$  and  $AT_2$  receptors; rabbit aorta; rat liver; bovine cerebellum; insurmountable antagonist

#### Introduction

in cardiovascular homeostasis, and the therapeutic success of angiotensin converting enzyme (ACE) inhibitors, such as cap-<br>times inhibitors, is to prevent the action of the main effector of this<br>topril and enalapril, in the treatment of hypertension and<br>system, angiotensin II (AII), topril and enalapril, in the treatment of hypertension and heart failure has confirmed the involvement of this system in these disease states. Despite their clinical success, the use of in nature, and many have retained significant agonist activity.<br>ACE inhibitors is not without problems, dry cough and a This, coupled with their short durati ACE inhibitors is not without problems, dry cough and a propensity to induce functional renal failure being amongst propensity to induce functional renal failure being amongst administered systemically, and poor oral bioavailability, has the most common (Gavras & Gavras, 1988).

The renin-angiotensin-aldosterone system plays a pivotal role An alternative way of intervening in the renin-angiotensin-<br>in cardiovascular homeostasis, and the therapeutic success of aldostererone axis, that might avoid t recently, angiotensin receptor antagonists have been peptidic<br>in nature, and many have retained significant agonist activity. prevented these compounds from establishing a therapeutic role. However, in 1988, Timmermans and associates reported that a series of l-benzylimidazole-5-acetate derivatives Author for correspondence.<br>Present address: Department of Pharmacology, Fisons, p.l.c., isolated tissues, including rat adrenal cortical microsomes Present address: Department of Pharmacology, Fisons, p.l.c., isolated tissues, including rat adrenal cortical microsomes<br>Lesearch and Development Laboratories. Pharmaceuticals Division. and rabbit isolated aorta (Chiu et a known example of these compounds is losartan (DuP753;

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Chiu et al., 1990). However, comparison of the profiles of action of compounds such as losartan, with that shown by imidazopyridines such as PD123177 (see Wong et al., 1990a), has revealed the existence of (at least) two distinct populations of angiotensin binding sites. These two binding sites are generally assumed to be synonymous with different angiotensin receptors, and it has been proposed that these be termed  $AT<sub>1</sub>$  and  $AT<sub>2</sub>$  receptors (Bumpus et al., 1991). Losartan and PD123177 are archetypal  $AT_1$  and  $AT_2$  receptor antagonists, respectively. The  $AT_1$  receptors appear to mediate most, if not all, of the established effects of All (e.g. vasoconstriction, steroidogenesis, dipsogenesis) and the role, if any, of  $AT_2$ receptors remains unclear (e.g. see Zarahn et al., 1992).

Recently, Middlemiss et al. (1991) reported on a series of bromobenzofurans, a novel class of potent, non-peptide, angiotensin receptor antagonists. These compounds inhibit AII-induced contraction of rabbit isolated aortic strips and reduce blood pressure in renal artery-ligated, hypertensive rats. Among these compounds, GRI17289 (1-[[3-bromo-2-[2- ( 1H - tetrazol - 5 - yl)phenyl] - 5 - benzofuranyl]methyl] - 2 - butyl - 4-chloro-lH-imidazole-5-carboxylic acid) (Figure 1) exhibited potent All antagonist activity in vitro, and exerted marked and prolonged antihypertensive activity after oral administration in vivo. This paper presents a detailed account of the pharmacological profile of GR117289 in vitro. Preliminary accounts have been presented to the British Pharmacological Society (Marshall et al., 1991; Robertson et al., 1991).

#### **Methods**

#### Functional studies in rabbit aorta

A detailed account of the rabbit isolated aortic strip preparation, and the experimental protocol used have been given previously (Robertson et al., 1992). Briefly, helical strips (1.5-2 cm) of thoracic aorta from male, New Zealand White rabbits were suspended in glass tissue chambers under a resting tension of 0.5 g in a physiological salt solution at 37°C, containing indomethacin  $(30 \,\mu\text{M})$  and ascorbic acid (100  $\mu$ M), and gassed with 95% O<sub>2</sub>/5% CO<sub>2</sub>.

In each experiment, four strips of aorta from a single rabbit were used simultaneously. In one series of experiments, the tissues were suspended under a resting tension of 0.5 <sup>g</sup> in <sup>a</sup> superfusion apparatus (see Coleman & Nials, 1989) in which the preparations were constantly superfused with physiological salt solution at 37C.

# Experimental protocols

In 4 separate, matched preparations, three consecutive concentration-contractile response curves to All (0.1 nM-100 nM) were constructed until reproducible, by the cumulative addition of All to the bathing fluid. The last of these



Figure 1 Chemical structure of GR117289.

was termed the 'pretest' curve. Subsequently, three tissues were exposed to one of three concentrations of GRI17289 (0.3, <sup>1</sup> or <sup>3</sup> nM), for periods ranging from <sup>45</sup> min to <sup>4</sup> h. A fourth tissue was exposed to vehicle alone and served as a time-matched control. After the antagonist (or vehicle) incubation period, a final ('test') cumulative All concentration-response curve was constructed. In a separate series of experiments, losartan (10, 100 or 1000 nM) was coincubated for varying periods with GRI 17289 (1 nM) before the test All curve. In some experiments, 5-hydroxytryptamine (5-HT,  $30 \text{ nM} - 10 \mu\text{M}$ ) or phenylephrine (10 nM- $30 \mu$ M) was used in place of AII throughout.

In experiments using superfusion, cumulative All concentration-response curves were constructed by sequentially increasing the concentration of All in the superfusate. The flow rate over each tissue was  $2 \text{ ml min}^{-1}$ . AII was introduced into the superfusate in solution at a rate of 0.02 ml min-'. GRI 17289 (0.3, <sup>1</sup> or <sup>3</sup> nM), or vehicle, was superfused for 45 min over the tissues (again, the flow rate of solution containing GR117289 was  $0.02$  ml min<sup>-1</sup>). A final All concentration-response curve (test) was then constructed, either immediately following the GRI 17289 (or vehicle) contact period, or after a subsequent 3 h period during which time the tissues were superfused (washed) with physiological salt solution alone.

#### Expression of results: determination of antagonist potency

For any individual preparation, the contractile response to each concentration of All (or 5-HT, or phenylephrine) in the test curve was expressed as a percentage of the maximum contractile response to the agonist in the pretest curve. Values are shown as mean  $\pm$  s.e.mean. In some cases, the concentration of agonist required to elicit a half maximum response  $(EC_{50})$  was calculated; geometric mean  $(95\%$  confidence intervals)  $EC_{50}$  values were calculated from these data.

Incubation with GRl 17289 reduced the maximal All contractile response (insurmountable antagonism). For this reason, conventional Schild analysis could not be used to obtain a measure of potency of GRl 17289. Instead, an apparent  $pK_B$  for GR117289 was derived by use of a doublereciprocal regression plot (Kenakin, 1984). A graph of 1/A vs  $1/A'$  was plotted, where A and A' were the equieffective concentrations of All in the absence or presence of GRI17289, respectively. The gradient (G) of this plot was then used in the Gaddum equation to obtain an estimate of  $pK_B$ : thus,  $pK_B = -\log$  ([B]/G-1), where B is the antagonist concentration.

#### Radioligand binding studies

Livers were obtained from Lister hooded rats (300-400 g) which were killed by cervical dislocation, and aortic tissue was obtained from male New Zealand White rabbits. Bovine cerebellum was obtained from Glaxo Institute for Molecular Biology, Geneva. The tissue was homogenized in 10-20 volumes  $(w/v)$  of ice cold homogenizing buffer (50 mM Tris, <sup>5</sup> mM EDTA, pH 7.4 at 4°C) in <sup>a</sup> Polytron PlO homogenizer. The homogenate was centrifuged at  $48,000 g$  for  $12 \text{ min}$  at 4'C and the supernatant discarded. The pellet was resuspended in homogenizing buffer using the Polytron P1O and then centrifuged as before. The final pellet was resuspended in <sup>50</sup> mM Tris buffer (pH 7.4 at <sup>25</sup>'C) at <sup>a</sup> tissue concentration of 400 mg ml<sup>-1</sup> and stored at  $-70^{\circ}$ C until required.

# $[3H]$ -angiotensin II binding to rat liver and bovine cerebellum membranes

Binding assays were performed by incubating the membranes (5 mg tissue/tube) in 500  $\mu$ l of assay buffer (Tris 50 mM,

NaCl 100 mM,  $MgCl<sub>2</sub>$  10 mM, EDTA 1 mM, BSA 0.001% w/v and bacitracin  $1 \text{ mM}$ ) with approximately 0.3 nM  $[3H]$ -AII with or without competing compounds. Assay tubes were incubated at room temperature for 90 min, after which time, bound and free radioactivity were separated by rapid filtration through Whatman GF/B glass fibre filters, which were pretreated with 0.1% polyethyleneimine (PEI), using a Brandel cell harvester. The filters were washed with ice cold wash buffer  $(100 \text{ mM } \text{NaCl}, 5 \text{ mM } \text{MgCl}_2)$  and trapped radioactivity was determined by liquid scintillation counting using a Packard 2200Ca scintillation counter. Specific binding was defined as that displaceable by  $1 \mu M$  AII. In saturation experiments, rat liver membranes were incubated with increasing concentrations of [3H]-AII as described. Bound radioactivity, in the presence and absence of  $1 \mu M$  AII for each concentration of [3H]-AII, was determined by filtration as described above. To determine the effects of GR117289 on saturation binding, membranes were incubated with <sup>12</sup> nM GR117289 for 30 min prior to the addition of the  $[3H]$ -AII. In dissociation experiments,  $[{}^{3}H]$ -AII (0.3 nM) was incubated to steady state with rat liver membranes (45 min at room temperature) in magnesium-free assay buffer. AII  $(1 \mu M)$ , with or without GR117289 (final concentration,  $1 \mu M$ ), was then added and the amount bound after various time intervals between 0 and 240 min was determined by filtration as previously described. Membranes were incubated to steady state in the presence (total) or absence (nonspecific) of  $1 \mu$ M AII and the dissociation rates of specific binding (total minus nonspecific binding) were determined.

# $[1^{125}I]$ -Sar<sup>1</sup>-Ile<sup>8</sup>-angiotensin II binding

Membranes (0.1 mg tissue/tube for rat liver and <sup>10</sup> mg tissue/ tube for rabbit aorta) were incubated in  $150 \mu l$  of assay buffer (Tris 50 mM, NaCl 100 nM,  $MgCl<sub>2</sub>$  10 mM, EDTA <sup>1</sup> mM, BSA 0.1% w/v and bacitracin <sup>1</sup> mM) with approximately  $0.05$  nM  $[$ <sup>125</sup>I]-Sar<sup>1</sup>-Ile<sup>8</sup>-angiotensin II ( $[$ <sup>125</sup>I]-Sarile) with or without competing compounds. Assay tubes were incubated for 90 min at room temperature after which time, bound and free radioactivity was separated as described above. Trapped radioactivity was measured with an LKB 1282 Compugamma counter. Specific binding was defined as that displaceable by  $3 \mu M$  AII.

#### Data analysis

Competition binding data were analysed by use of iterative curve fitting techniques (Michel & Whiting, 1984). Data from [<sup>125</sup>I]-Sarile binding experiments are presented as the negative logarithm of the  $IC_{50}$  (pIC<sub>50</sub>). In the case of [3H]-AII experiments, the  $IC_{50}$  values were corrected for the presence of the radioligand using the Cheng-Prusoff approximation (Cheng & Prusoff, 1973) and are presented as  $pK_i$ . Saturation experiments are presented in a Scatchard plot.  $K_d$  and  $B_{\text{max}}$ values were calculated by use of LIGAND (Munson & Rodbard, 1980). In dissociation experiments, dissociation rates were calculated using ENZFITTER (Biosoft, Cambridge).

#### Drugs used

Angiotensin II (human sequence) was obtained from Nova biochem., U.K. Ltd., Bacitracin, bovine serum albumin (BSA), 5-hydroxytryptamine (creatinine sulphate complex), phenylephrine hydrochloride and indomethacin were obtained from Sigma. [3H]-angiotensin II ([3H]-AII) and [1251]-Sar<sup>1</sup>Ile<sup>8</sup> angiotensin II ([<sup>125</sup>I]-Sarile) were obtained from NEN, Du Pont (specific activities of 73.4 and 2200 Ci mMol<sup>-1</sup>). GR117289 and losartan (potassium salt) were synthesized in the Chemistry Division, Glaxo Group Research Ltd.

Angiotensin II was dissolved in distilled water and stored frozen as aliquots (1 mM) which were thawed on the day of experimentation and diluted with distilled water. Phenyle-

phrine and 5-HT were dissolved and diluted with distilled water. Indomethacin was dissolved in NaOH (2 M) and added directly to the physiological salt solution to give a final concentration of  $30 \mu$ M. Bacitracin and BSA were dissolved directly in Tris buffer.

All other chemicals and constituents for the physiological salt solution were of Analar grade and obtained from BDH Ltd., England.

#### Results

#### The effect of GR117289 on angiotensin II-induced contraction in rabbit aortic strips

Cumulative addition of AII  $(0.3 \text{ nm} - 0.3 \mu\text{m})$  caused concentration-related contractions of rabbit isolated aortic strips. After a 45 min incubation period, GRl 17289 (0.3, <sup>1</sup> or 3 nM) produced a concentration-related, insurmountable antagonism of AII (Figure 2a). GR117289 (3 nM) caused approximately 65% suppression of the maximum response to All. Despite this profound, concentration-related suppression of the maximum response to All, GRI17289 did not significantly change the  $EC_{50}$  of AII. The geometric mean  $EC_{50}$  (95% confidence limits) values in the absence and presence of 0.3, 1 or 3 nm GR117289 were 5.4  $(3.4-8.7)$ , 5.3  $(3.2-8.3)$ , 5.85  $(4.7-7.1)$ , and 6.13  $(5.5-6.8)$ nM, respectively. For comparison, Figure 2b shows the effect of the sur-



Figure 2 The contractile effect of angiotensin II (All) in the rabbit isolated aorta, in the presence of (a) GR117289 ( $\bigcirc$  0.3,  $\bigcirc$  1 or  $\Box$ 3 nM) or (b) losartan ( $O$  30,  $\Delta$  100 or  $\Box$  1000 nM) or vehicle for losartan or GR117289 ( $\bullet$ ) previously incubated for 45 min (n = 4 for each). Results are shown as mean response (± s.e.mean, vertical bars) in the test curve, expressed as <sup>a</sup> % of the maximum response to AII in the pretest curve. Data for losartan are from Robertson et al. (1992) and republished with permission.

mountable, competitive angiotensin receptor antagonist, losartan (30, 100 and 300 nM) on All-induced contractions in the rabbit aorta: losartan did not affect the maximum response to AII. As found previously (Robertson et al., 1992), the  $pA_2$  of losartan was calculated to be  $8.2 \pm 0.1$ (slope =  $1.1 \pm 0.1$ ) by Schild analysis.

Like AII, both 5-HT (30 nM-10  $\mu$ M) and phenylephrine (10 nM-30  $\mu$ M) caused concentration-related contractions in the rabbit aorta. GR117289 (1 $\mu$ M) had no significant effect on the contractile response to either agonist (data not illustrated).

All concentration-response curves, before and after a <sup>3</sup> h incubation with the vehicle for GRI17289 (pretest and test curves) were highly reproducible. Compared with their respective time-matched vehicle controls, a 3 h incubation with GR117289 (0.3 nM) caused a more profound reduction (77% suppression, Figure 3a) of the maximum response to All than had been observed after only 45 min (25% suppression, Figure 2a) or 2 h incubation (36% suppression, data not shown). In tissues incubated with vehicle for 4 h, All responses became more variable (Figure 3b). However, GR1 17289 (0.3 nM), caused little further reduction (81% suppression, Figure 3b) of the maximum response to All under these conditions, compared with that seen after <sup>3</sup> h incubation. Thus, GR117289 seemed to have reached equilibrium after approximately <sup>3</sup> h incubation. Using the method of Kenakin (1984), the p $K_B$  of GR117289 was derived from 7 separate experiments in which GR117289 (0.3 nM) was incubated with aortic tissue for 3 h: a  $pK_B$  of 9.8  $\pm$  0.1 was



Figure 3 The contractile effect of angiotensin II (AII) in the rabbit isolated aorta, in the presence of GR1 17289 (O 0.3 nM) or vehicle for GR117289 ( $\bullet$ ) incubated for either (a), 3 h or (b), 4 h (n = 4 for each).

calculated. The geometric mean  $EC_{50}$  (95% confidence interval) of AII was significantly greater  $(P < 0.005$ , Student's unpaired t test) in the presence of GR117289 (8.2 (5.1-13.2) nM), than in its absence  $(2.7 (1.5-4.9)$  nM), indicating a small rightward displacement of the All concentration-response curve.

#### The effect of losartan on GRJI7289-induced antagonism of angiotensin II

In tissues incubated with GRI17289 (1 nM) for <sup>2</sup> <sup>h</sup> <sup>45</sup> min, the All concentration-response curve was markedly suppressed and displaced to the right, compared with timematched, vehicle-treated controls. In preparations which were co-incubated with losartan (10 nM, 100 nM or 1  $\mu$ M) for the final 45 min of the GR117289 incubation period, the subsequent All concentration-response curves were displaced upwards (with 10 nM, 100 nM or  $1 \mu$ M losartan) and to the right (with 100 nM or  $1 \mu$ M losartan), in a concentrationrelated manner (Figure 4).

In a related series of experiments, GR117289 (1 nM) incubated with rabbit aorta for 2 h, again caused a marked suppression and rightward displacement of the All concentration-response curve, although this was not as marked as that seen when the incubation time was 2 h 45 min (compare Figure <sup>5</sup> with Figure 4). In the same series of experiments, GR1 17289 (1 nM) was incubated for <sup>2</sup> h and then losartan  $(1 \mu M)$  was also added for either 2, 15 or 45 min, before construction of the test All curve. Coincubation of losartan with GR1 17289 caused <sup>a</sup> larger rightward displacement of the All concentration-response curve



Figure 4 The contractile effect of angiotensin II (All) in rabbit isolated aorta, in the absence (@) or presence of GRI 17289 (A 1 nm, 2 h 45 min incubation) alone, or losartan ( $\square$  10 nm,  $\bigcirc$  100 nm, or  $\nabla$  1  $\mu$ m) co-incubated with GR117289 for the final 45 min of the 2 h 45 min incubation period  $(n = 4)$ .



Figure 5 The contractile effect of angiotensin II (All) in the rabbit isolated aorta, in the absence  $(\bullet)$  or presence of GR117289 ( $\blacksquare$  1 nm, 2 h incubation) alone, or co-incubated with losartan  $(1 \mu M)$  for a further 2 min  $(\nabla)$ , 15 min (O) or 45 min ( $\Delta$ ) (n = 5).

than incubation with GR117289 alone. The concentrations of AII required to produce 50% of the maximum response in the test curve, in preparations co-incubated with losartan and GRI17289 were similar, regardless of the duration of the losartan incubation (geometric mean  $EC_{50}$  values (95%) confidence limits) after 2, 15 and 45 min losartan coincubation were 731 (224-2381), 621 (260-1483) and 625 (209-1875)nM, respectively). In contrast, co-incubation with losartan caused a time-related increase of the maximum response to AII compared with that observed in preparations treated with GRI17289 alone (Figure 5).

## The effect of continuous washing on GR117289 antagonism of angiotensin II in superfused rabbit aortic strips

In superfused rabbit aortic strips, increasing concentrations of AII  $(1-30 \text{ nm})$  caused concentration-related increases in the contractile tension. The sensitivity of the preparations to AII was similar to that seen in the experiments carried out in a fixed volume of bathing fluid. As before, GRI 17289 (0.3, <sup>1</sup> or 3 nM) perfused over the tissues for 45 min, caused a concentration-related, insurmountable suppression of the contractile response to AI. The highest concentration of GR1 17289 used (3 nM), reduced the maximum response to AII by approximately 92%. When separate tissues from the same rabbits were treated in the same way with GR117289, but then superfused with drug-free physiological salt solution



Figure 6 The contractile effect of angiotensin II (All) in rabbit superfused isolated aorta, (a) in the absence of GR117289  $(\bullet)$  or after GR117289 ( $\bigcirc$  0.3,  $\bigcirc$  1 or  $\Box$  3 nm) was superfused for 45 min or, (b) in the absence of GR117289 ( $\bullet$ ) or after GR117289 ( $\circ$  0.3,  $\Delta$  1 or  $\Box$  3 nM) were superfused for 45 min and followed by drugfree physiological salt solution (prolonged washing) for 3 h, respectively  $(n = 3)$ .

alone for another 3 h, subsequent All responses were not restored; rather they were reduced further (see Figure 6b).

# Characteristics of GR117289 binding in rat liver and bovine cerebellum

In competition binding studies using  $[3H]$ -AII at angiotensin AT, receptors in rat liver membranes, GR1<sup>17289</sup> was potent with a p $K_i$  of 8.7  $\pm$  0.1 (n = 12) and a Hill coefficient (nH) close to unity (1.11). In contrast, in bovine cerebellum, GR117289 had negligible affinity for angiotensin  $AT_2$  sites  $(pK_i \leq 6, n = 2)$ .

The  $AT_1$  receptor has been shown to exist in high and low affinity states depending on its G-protein coupling. The competition studies described above were conducted in the presence of  $Mg^{2+}$ , which would result in the majority of the receptors being in the high affinity state with respect to agonist binding. In order to determine whether GRI17289 could discriminate between high and low affinity states of the  $AT<sub>1</sub>$  receptor, competition studies were carried out in the presence and absence of  $Mg^{2+}$ . Under these conditions, the pIC<sub>50</sub> values were found to be  $8.5 \pm 0.1$  ( $n = 3$ , nH = 0.99) and  $8.7 \pm 0.2$  ( $n = 3$ , nH = 1.31), respectively.

In order to determine whether GR117289 attained equilibrium during the 90 min incubation period of the [<sup>3</sup>H]-AII binding assay, the liver membranes were preincubated with the antagonist for <sup>1</sup> h before addition of the radioligand. A <sup>1</sup> h preincubation resulted in a small, but not significant, increase in the p $K_i$  from  $8.7 \pm 0.3$  ( $n = 3$ , nH = 1.06) to 9.1  $\pm$  0.21 (*n* = 3, nH = 1.4).

To establish the nature of the antagonism exerted by GR117289, its effect on the saturation curve of [<sup>3</sup>H]-AII binding was determined. The results for GR117289 (12 nM) are shown in Figure 7 in the form of a Scatchard analysis. GR117289 (12 nM) decreased the  $B_{\text{max}}$  from 10.0  $\pm$  0.1 fmol mg<sup>-1</sup> tissue to  $5.6 \pm 0.3$  fmol mg<sup>-1</sup> tissue and increased the  $K_d$  of [<sup>3</sup>H]-AII-binding from  $0.28 \pm 0.06$  nM to  $0.37 \pm 0.02$  nM  $(n = 3)$ .

In order to investigate further the nature of the antagonism, the effect of GR117289 (1.0  $\mu$ M) on the dissociation of [3H]-AII, caused by addition of an excess of unlabelled All, was measured. There was no significant difference between the dissociation rate in the absence  $(t_1 = 6.5 \pm 1.1 \text{ min})$  or presence  $(t_1 = 7.4 \pm 1.9 \text{ min})$  of GR117289 (1.0  $\mu$ M,  $n=3$ ).

The affinity of GR117289 in binding assays was found to be markedly dependent on the concentration of BSA in the assay buffer. Increasing the BSA from 0.001% to 0.1% w/v resulted in a large rightward displacement of the inhibition curve which described the competition of  $[3H]$ -AII with GR117289. In the presence of  $0.1\%$  w/v BSA, the pK<sub>i</sub> of GR117289 in the rat liver decreased to  $7.5 \pm 0.1$  (nH = 1.11,  $n = 5$ ).



Figure 7 Scatchard analysis of specific binding data for [<sup>3</sup>H]angiotensin II ([ $^3$ H]-AII) in the absence ( $\bigcirc$ ) or presence ( $\bigcirc$ ) of GR117289 (12 nM). Data are taken from a representative experiment.

## Characteristics of GR117289 binding in rabbit aorta

In an attempt to correlate findings in functional and radioligand binding sites, the affinity of GRI 17289 for angiotensin binding sites in rabbit aorta smooth muscle membranes was determined. Because of the low density of receptors in this tissue, a higher specific activity ligand, ['251]-Sarile was used. In addition, to increase specific binding, the assay buffer contained 0.1% w/v BSA. Under these conditions, GR117289 competed with  $[1^{25}I]$ -Sarile for binding to angiotensin receptors and yielded a mean pIC<sub>50</sub> of  $7.6 \pm 0.1$  (nH = 1.13,  $n = 3$ ). Under the same conditions, but with rat liver membranes, the pIC<sub>50</sub> for GR117289 was  $7.8 \pm 0.1$  (nH = 1.08,  $n = 5$ ).

## **Discussion**

GR117289 is a potent antagonist against AII in rabbit aortic strips, and competes with [3H]-AII for binding sites in rat liver and with ['H]-AII or ['<sup>25</sup>I]-Sarile on rabbit aorta vascular smooth muscle membranes. These preparations were chosen because it is well established that the angiotensin receptors present belong to the  $AT_1$  subtype (Wong et al., 1990a; de Gasparo et al., 1990; Chang & Lotti, 1991) and  $AT<sub>1</sub>$  receptors are the only subtype that, thus far, are known to mediate the major physiological effects of All. In contrast, GR117289 had negligible affinity for  $AT_2$  sites in bovine cerebellum, thus establishing its high selectivity for  $AT_1$ receptors.

We have previously confirmed functionally that losartan is a potent, competitive, surmountable antagonist ( $pA_2 = 8.2$ ) of All in the rabbit aorta, using a tissue contact time of 45 min (Robertson et al., 1992). However, the most striking difference between GRI 17289 and losartan in this preparation, is the way in which the two compounds displace the concentration-response curves to All. Losartan caused a concentration-dependent, parallel, rightward displacement, without significantly affecting the maximum contractile response to AII. In contrast, GR117289 caused a concentration-related, insurmountable suppression of the All concentration-response-curve. This profile of action could be explained by a number of different mechanisms: GR117289 may be (1) a 'non-specific' antagonist, (2) an allosteric antagonist, (3) an irreversible antagonist, or (4) a slowly reversible (pseudo-irreversible) antagonist, of All.

#### Is GR117289 a 'non-specific' antagonist?

In this context, the term 'non-specific' is used to describe an antagonist which blocks, at some point, the chain of events leading to the production of a response by All. Alternatively, the compound may antagonize at a receptor unrelated to angiotensin.

Even at the relatively high concentration of  $1 \mu M$ , GR1 17289 did not affect contractile responses induced by 5-HT or phenylephrine in the rabbit aorta. This indicates not only that GR117289 has a low affinity ( $pK_B < 6$ ) for both 5-HT<sub>2</sub> receptors and  $\alpha_1$ -adrenoceptors in this tissue, but also that GR117289 does not interfere with the signal-transduction process that mediates the contraction to All since AII (Griendling et al., 1989), 5-HT (Roth et al., 1986) and phenylephrine (Homcy & Graham, 1985), all induce smooth muscle contraction by receptor mediated hydrolysis of phosphatidylinositol leading to increased  $[Ca^{2+}]$ . Thus, these data suggest that the site of action of GR117289 is the  $AT_1$ receptor macromolecule or a closely related (e.g. allosteric) site.

Further evaluation in other functional and radioligand binding studies has shown that, at  $\geq 1 \mu M$ , GR117289 has no detectable affinity for a wide range of receptors/binding<br>sites/enzymes including adrenoceptors  $(\alpha_2 - \beta_1 - \beta_2 - \beta_3 - \beta_4)$ . sites/enzymes including adrenoceptors  $(\alpha_2, \beta_1,$ adenosine  $(A_1, A_2)$ , bradykinin, dopamine  $(D_1, D_2)$ , 5hydroxytryptamine  $(5-HT_1, 5-HT_3)$ ,  $\gamma$ -aminobutyric acid (GABA<sub>A</sub>, GABA<sub>B</sub>), glycine, muscarinic  $(M_1, M_2)$ , opioid ( $\mu$ -, 6-, K-), neuropeptide Y and neurotensin receptors, ACE, renin or adenylate cyclase (unpublished observations).

# Is GR117289 an allosteric antagonist?

An allosteric antagonist is one that binds to a site on the receptor macromolecule or cell membrane, close to, but not at, the site at which the agonist binds. Nevertheless, occupation of the allosteric site by the antagonist impairs the ability of the agonist-receptor complex to generate a response. Such <sup>a</sup> phenomenon has been proposed by Kaumann & Frenken (1985) to explain the insurmountable suppression, by methysergide, of vasoconstrictor responses to 5-HT in calf coronary arteries. The functional experiments described in this report cannot exclude the possibility that GRI17289 behaves as an allosteric antagonist of AIl. However, in radioligand binding studies (see below), GRi17289 was shown not to affect the rate of dissociation of  $[{}^{3}H]$ -AII from AT<sub>1</sub> receptors in rat liver membranes, as measured in isotope dilution experiments. Therefore, provided that GRI17289 binds to  $AT<sub>1</sub>$  receptors in the rabbit aorta and rat liver membranes in a comparable manner, an allosteric mechanism of action in the aorta is unlikely.

## Is GR117289 an irreversible antagonist?

Insurmountable antagonism can also occur when an antagonist forms an irreversible covalent bond with the receptor, such that the receptor number is effectively reduced to the point where a full agonist response cannot be achieved. However, it is unlikely that GRI17289 binds irreversibly to  $AT_1$  receptors. Evidence for this view comes from two main observations. Firstly, although the extent of the blocking action of GRI 17289 is progressive over <sup>3</sup> h of incubation, increasing the incubation time to 4h did not appear to result in any greater degree of antagonism. Thus,  $G\dot{R}$ 117289 seemed eventually to reach equilibrium with  $AT_1$ receptors in the rabbit aorta. Secondly, the degree of suppression, induced by GRI 17289, of the maximal response to AII, was reduced by co-incubation with losartan. Increasing concentrations of losartan displaced the concentration-response curve to AII, in the presence of GR1 17289, upward and to the right (Figure 4). If GR117289 had bound covalently to the  $AT_1$  receptor, then co-incubation with a competitive antagonist would not have been expected to have any effect on the AII maximum.

#### Is GRJ17289 a slowly reversible (pseudo-irreversible) antagonist?

A further explanation of insurmountable antagonism is that slow dissociation of the antagonist from the receptor occurs. In this situation, the agonist cannot reach equilibrium with the antagonist/receptor complex under the time constraints of the experiments (see Craig et al., 1990). In other words, the antagonist, once bound to the receptor, may dissociate from it so slowly as to appear irreversibly bound (pseudoirreversible antagonism). Thus, a simple explanation for the findings of the experiments described above in which GR1 17289 and losartan were co-incubated, is that GR1 17289 dissociates slowly from the  $AT_1$  receptors but, as it does, vacated receptors become occupied by losartan. Subsequently administered All then competes for receptors occupied by losartan, as well as those still occupied by GRI 17289. As the concentration of losartan is increased, proportionally more receptors are occupied by losartan than by GR117289. Because All can surmount the antagonism exerted by losartan, the maximal response to All increases in relation to the proportion of receptors occupied by losartan rather than by GR117289. If GR117289 had bound irreversibly to the  $AT_1$ receptors, co-incubation with losartan would merely have produced further rightward displacement of the already suppressed All curves. Similar to the present data, Wienen et al. (1990) and Wong & Timmermans (1991), demonstrated that the suppression of the AII curve produced by Sar'Ile8-AII and EXP3892, respectively, could also be reversed by losartan in rabbit aorta. More recent data (Entzeroth et al., 1991) have confirmed and expanded these observations.

For this explanation of the interaction between GRI 17289 and losartan to be valid, it is necessary that losartan reaches equilibrium with the  $AT_1$  receptor more rapidly than GR1 17289. In the present study, no experiments were conducted to determine whether losartan had reached equilibrium within the 45 min incubation period. However, other published evidence suggests that equilibrium is achieved very quickly. Using incubation times of less than 45 min, other groups have reported  $pA_2$  values for losartan against AII in the rabbit aorta, similar to the value (8.2) obtained by Robertson *et al.* (1992) (e.g. Rhaleb *et al.*, 1991, 10 min incubation time,  $pA_2 8.27$ ; Wong *et al.*, 1990b, 15 min incubation time,  $pA_2$  8.48). It would therefore appear that, unlike GR117289, losartan reaches equilibrium with the  $AT_1$ receptor rapidly, and certainly within 45 min. Further evidence in support of this concept is provided by the results shown in Figure 5, in which a single concentration of losartan was co-incubated for varying periods of time after pretreatment with GR117289. In these experiments, similar rightward displacements of the All curve were seen after 2 min or 45 min incubation with losartan, again suggesting a rapid equilibration of losartan with the receptors. Figure 5 also shows that the increase in the maximum response to All, in the presence of both antagonists, was progressive as the duration of losartan incubation was increased. This may be a measure of the slow rate of dissociation of GR117289 from the receptor.

The results obtained from the radioligand binding studies broadly support those from functional studies. In rat liver homogenates, the p $K_i$  estimate for GR117289 was  $8.7 \pm 0.1$ . This is lower than was obtained in the rabbit aorta  $(pK_B = 9.8)$  and the Hill coefficient suggested that GR117289 and [3H]-AII competed reversibly for the binding sites. However, it should be remembered that the competing agents were administered simultaneously in the binding studies, whereas the aorta was exposed to GR117289 for 3 h before challenge with All. Pre-incubation of liver membranes with GR117289 for 1 h before addition of  $[3H]$ -AII had little effect on the affinity estimate for GRl 17289. It was impractical to use longer pre-incubation times to determine whether the affinity of GRI 17289 would increase further.

The failure of GR117289 to increase the rate of dissociation of [3H]-AII from liver membranes, following isotope dilution with excess, unlabelled All argues against the binding of GR117289 to an allosteric site on the  $AT_1$  receptor macromolecule. However, examination of the data shown in Figure 7 shows that the nature of the binding of GR117289 to liver membranes is characteristic of neither a simple, competitive antagonist, nor of a wholly irreversible antagonist. GR117289 both reduced the  $B_{\text{max}}$  and increased the  $K_d$  of [H]-AII binding. These findings are, therefore, consistent with the view that GRl 17289 may be a slowly reversible antagonist at  $AT_1$  receptors.

Direct evidence for occupation of  $AT_1$  receptors on rabbit aorta by GR117289, was obtained from binding studies carried out on smooth muscle membranes from this tissue. However, the low density of receptors present in these preparations meant that the experimental conditions were different from those previously used with rat liver membranes. In particular, the BSA concentration was  $0.1\%$  w/v in these experiments. This almost certainly accounts for the low pIC<sub>50</sub> value  $(7.6 \pm 0.1)$  obtained for GR117289 in this preparation, a view supported by the finding of a similarly low pIC<sub>50</sub>  $(7.8 \pm 0.1)$  in rat liver membranes used under identical conditions. Chiu *et al.* (1991) have previously reported that BSA inhibits the binding of several di-acid

analogues of losartan. Thus, it is highly likely that GR117289 occupies <sup>a</sup> common binding site (i.e. receptor) in the two tissues, and that the measurements of potency obtained under these experimental conditions are underestimates of its true affinity.

The accuracy of the estimate of the dissociation constant made for GR117289, after 3 h incubation in the rabbit aorta, is debatable. It assumes that GRI17289 simply occupies the same receptor site as All and that their interaction reflects this phenomenon. However, the potency estimate made from functional studies is approximately 5 times higher than that determined from binding studies. This may be attributable to differences in experimental protocols (e.g. temperature, pretreatment time, whole tissue or membranes, the presence or absence of BSA). Although the results obtained in experiments in which rabbit aortic strips were continuously superfused with drug-free physiological salt solution for 3 h, after 45 min exposure to GR117289, are consistent with its being a very slowly dissociating antagonist, other factors may contribute to the long-lasting inhibition of responses to subsequently administered All. In particular, GRI17289 is a highly lipophilic agent (cLog  $P = 7.5$ ). Thus, it is conceivable that the profile of activity of GRI17289 is attributable, at least in part, to retention or, perhaps, even concentration within the membrane lipid. Apart from making it difficult to remove GR1 17289 by washing, this property might influence the estimation of its affinity for the angiotensin receptor in the rabbit aorta. For example, GRI17289 might modulate the interaction between All and its active site, not by changing its affinity (see above), but by reducing its efficacy as a result of altering the balance between receptor internalisation and expression, as has been suggested for some peptide antagonists of All (Liu et al., 1992). Thus, a smaller stimulus would be generated for any given agonist concentration. This would be reflected by a suppression, with little displacement, of the All concentration-response curve. For this reason, the  $pK_B$  of 9.8 for GR117289 in this preparation should be regarded as no more than an approximation of its true affinity.

Taken as a whole, the data suggest that the most likely explanation for the insurmountable antagonism of All by GRI17289 in the rabbit aorta is attributable to its slow association with, and dissociation from, the  $AT_1$  receptor (pseudo-irreversible antagonism). However, other explanations of the data are possible. For example, de Chaffoy de Courcelles et al. (1986) have described a receptor-transducer coupling model to explain insurmountable antagonism at  $5-\text{HT}_2$  receptor sites. In this model, an insurmountable antagonist produces its effect by binding to the receptor in such a way as to induce a conformational change in the receptor, which subsequently results in a decrease in the efficiency of stimulus-response coupling. Wong & Timmermans (1991) have tentatively applied this model to explain the insurmountable antagonism produced by EXP3892 (2'propyl-4'trifluoromethyl-5'-carboxylic acid derivative of losartan) at angiotensin  $AT_1$  receptors in the rabbit aorta. In this model, competitive, surmountable antagonists can take the place of insurmountable antagonists, and this would explain the interaction we have observed between losartan and GR117289. However, it is not possible to distinguish between this type of interaction and pseudo-irreversible antagonism.

There are several precedents in the angiotensin literature which suggest that insurmountable antagonism is not unique to GR117289. For example, in rabbit aortic strips, suppression of the All curve has been also reported for sarcosine substituted peptide angiotensin receptor antagonists such as Sar<sup>1</sup>-Cys(Me)<sup>8</sup>-AII (70% suppression; Freer et al., 1980), Sar<sup>1</sup>Ile<sup>8</sup>-AII (40-90% suppression; Wienen et al., 1990) or Sar<sup>1</sup>-Ala<sup>8</sup>-AII (saralasin, 50% suppression; Freer et al., 1980; Chiu et al., 1990). Similar findings have been made with non-peptide antagonists such as EXP3174 (5' carboxylic acid metabolite of losartan, 40% suppression; Wong et al., 1990c) and EXP3892, (40% suppression; Wong & Timmermans, 1991).

In conclusion, the present studies suggest that GRI 17289 is a potent, non-peptide, specific, selective and insurmountable, angiotensin AT, receptor antagonist. The mechanism of the insurmountable antagonism is debatable, but the data are

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consistent with GRI 17289 being a slowly reversible (pseudoirreversible) antagonist.

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