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LOCAL RENIN-ANGIOTENSIN SYSTEMS



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LOCAL RENIN-ANGIOTENSIN SYSTEMS

(LOKALE RENINE-ANGIOTENSINE SYSTEMEN)

PROEFSCHRIFT

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Voor mijn ouders

Aan Romana

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1. Admiraal PJJ, Derkx FHM, Danser AHJ, Pieterman H, Schalekamp MADH. Metabolism and production of angiotensin I in different vascular beds in subjects with hypertension. *Hypertension* 15: 44-55, 1990.
2. Danser AHJ, Koning MMG, Admiraal PJJ, Derkx FHM, Verdouw PD, Schalekamp MADH. Metabolism of angiotensin I by different tissues in the intact animal. Provisionally accepted *Am J Physiol*.
3. Danser AHJ, Koning MMG, Admiraal PJJ, Sassen LMA, Derkx FHM, Verdouw PD, Schalekamp MADH. Production of angiotensins I and II at tissue sites in the intact pig. Provisionally accepted *Am J Physiol*.
4. Danser AHJ, van den Dorpel MA, Deinum J, Derkx FHM, Franken AAM, Peperkamp E, de Jong PTVM, Schalekamp MADH. Renin, prorenin, and immunoreactive renin in vitreous fluid from eyes with and without diabetic retinopathy. *J Clin Endocrinol Metab* 68: 160-167, 1989.
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7. Wagner J, Danser AHJ, Paul M, Derkx FHM, de Jong PTVM, Schalekamp MADH, Ganten D. Demonstration of renin-, angiotensinogen- and angiotensin converting enzyme-mRNA expression in human eyes by the polymerase chain reaction. In preparation.

1. GENERAL INTRODUCTION

1.1 History

In 1898 Tigerstedt and Bergman (1) found that crude saline extracts of rabbit kidneys contained a pressor substance, which they named *renin*. More than 30 years later, in 1934, Goldblatt and his colleagues recognized the significance of this finding when they discovered that it was possible to produce persistent hypertension in dogs by clamping the renal artery (2). In 1940, Braun-Menendez and his co-workers in Argentina (3) and Page and Helmer at the Cleveland Clinic in the United States (4) reported that renin was an enzyme that acted on a plasma protein substrate to catalyze the formation of the actual pressor material, a peptide, which was named *hypertensin* by the former group and *angiotonin* by the latter. These 2 terms were used for nearly 20 years, until it was agreed to rename the pressor substance *angiotensin* and to call the plasma protein substrate *angiotensinogen*. In 1954 two forms of angiotensin were recognized (5): the first a decapeptide (*angiotensin I*) and the second an octapeptide (*angiotensin II*), formed from angiotensin I by cleavage by another enzyme, termed *angiotensin converting enzyme*. Since these discoveries, ample evidence for the involvement of the renin-angiotensin system in the pathogenesis of various forms of clinical and experimental hypertension has been obtained by many investigators.

1.2 Circulating and tissue renin-angiotensin system(s)

Circulating renin-angiotensin system. The renin-angiotensin system has traditionally been viewed as an endocrine system. Circulating renin, released from the kidney by a variety of stimuli (of which the most important are hypotension and diminished delivery of sodium to the distal tubular macula densa sites), continuously produces angiotensin I from circulating liver-derived angiotensinogen (Figure 1). Angiotensin I is converted to angiotensin II as the blood passes through the lungs and other vascular beds where angiotensin converting enzyme (ACE) is present in high concentrations on vascular endothelial cells. Circulating angiotensin II controls blood pressure directly by arteriolar vasoconstriction and indirectly by regulating sodium and potassium homeostasis. The latter action of angiotensin II functions via a direct effect on renal sodium reabsorption (6) and via stimulation of adrenal aldosterone biosynthesis (7) and secretion (8), which, in turn, promotes renal sodium reabsorption and potassium secretion.

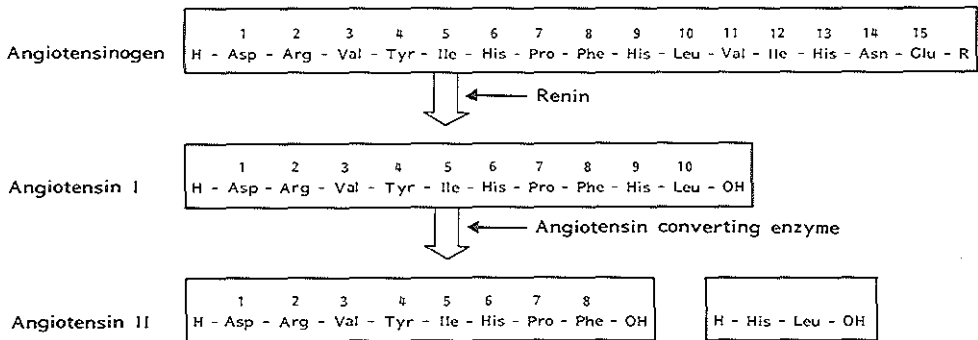


Figure 1. Outline of the biochemistry of the human renin-angiotensin system. Renin cleaves its substrate (angiotensinogen) between leucine 10 and valine 11 to form the decapeptide angiotensin I. Angiotensin converting enzyme removes residues histidine 9 and leucine 10 to form the octapeptide angiotensin II, the biologically active endproduct of the system.

Local renin-angiotensin systems. Recent evidence suggests that besides a circulating renin-angiotensin system of renal origin, there also exist tissue or local renin-angiotensin systems in different organ sites. Renin-like enzymatic activity, angiotensinogen, ACE, angiotensin I, angiotensin II and angiotensin receptors have been demonstrated in multiple tissues. In addition, molecular biologic techniques have clearly shown that both renin and angiotensinogen genes are expressed in organs involved in cardiovascular homeostasis (i.e. kidney, adrenal, heart, brain and blood vessels) (9,10) as well as in reproductive organs (testis and ovary) (10). Thus, an emerging concept is that angiotensin may be produced locally in many tissues, independent of the circulating system, and that locally generated angiotensin may play an important role in influencing the function of these target organs. Locally generated angiotensin may even reach the circulating blood and thus contribute to the angiotensin levels found in plasma (11).

Metabolism of angiotensins. Extensive metabolism of angiotensins is known to occur in peripheral tissues, due to the enzymatic activity of so-called 'angiotensinases' (Figure 2): aminopeptidases ('angiotensinase A': aminopeptidase A, aminopeptidase B, aminopeptidase M, leucine aminopeptidase, dipeptidyl peptidase) (12-14), endopeptidases ('angiotensinase B': prolyl-endopeptidase, neutral endopeptidase 24.11) (15) and

carboxypeptidases ('angiotensinase C': carboxypeptidase P, prolyl-carboxypeptidase, ACE, cathepsin A) (16,17). Yet, despite this high degree of metabolism, the concentrations of angiotensin I and angiotensin II in venous blood are similar to those in arterial blood (11,18). Angiotensin generation in blood alone seems to be too low to account for the levels of angiotensin I and angiotensin II in venous blood (11,18). Peripheral tissues may therefore indeed be a major site of angiotensin production. Possibly, the venous plasma levels of angiotensins I and II represent a spillover from the tissue sites of production (Figure 3).

However, plasma angiotensin levels in anephric subjects are usually low or undetectable (19), suggesting that it is mainly kidney-derived renin that is responsible for the presence of angiotensins in plasma. Tissue production and subsequent release of angiotensins into the circulation may therefore largely depend upon uptake of circulating kidney-derived renin by the tissues. The local tissue concentration of ACE and angiotensin degrading enzymes would then determine how much of the locally generated angiotensin I and angiotensin II is released into the circulation.

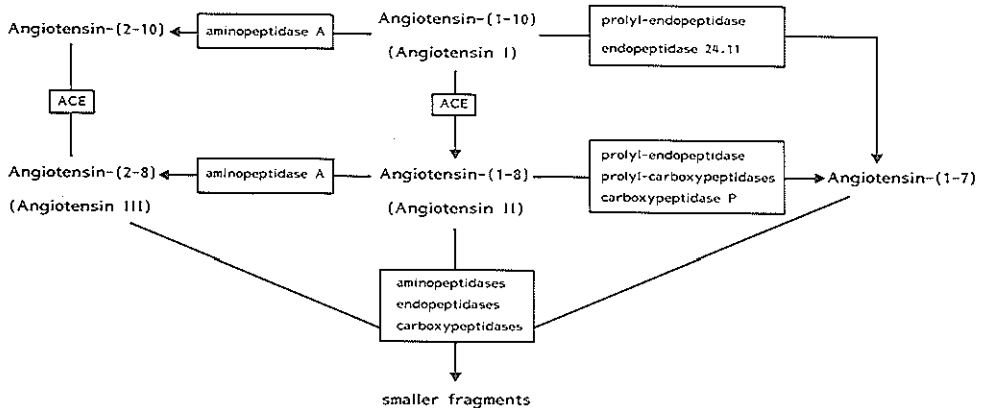


Figure 2. Schematic outline of alternate enzymatic cascades participating in the metabolism of angiotensins.

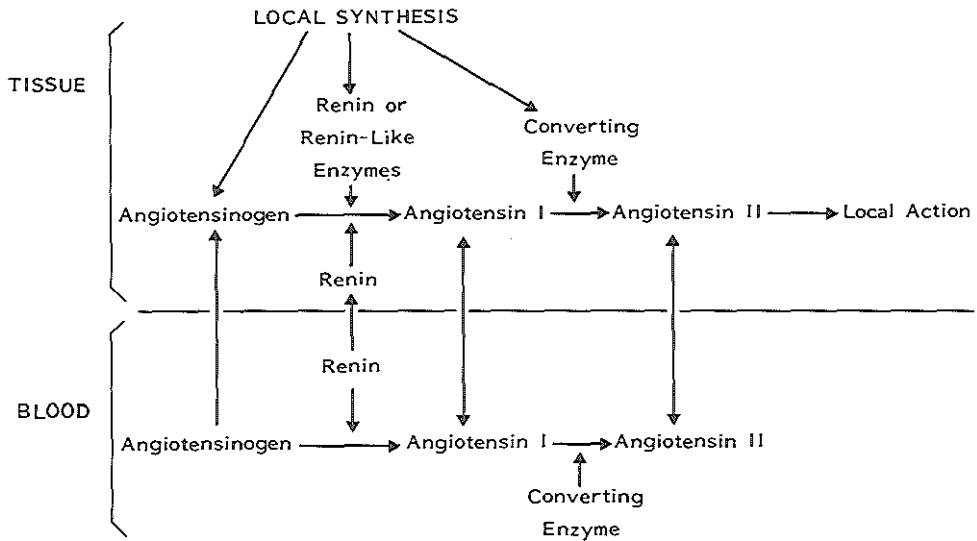


Figure 3. Circulating renin-angiotensin system vs. local renin-angiotensin system: interactions and possible angiotensin generation sites.

Prorenin. In contrast with the low levels of angiotensin (and renin) in plasma from nephrectomized subjects, the plasma concentrations of prorenin, the inactive precursor of renin, in these subjects are still in the (low) normal range (20). This shows that a substantial fraction of plasma prorenin is produced at extrarenal sites. Indeed the release of prorenin, but not renin, into the circulation has now been shown for the ovary, testis and adrenal (21,22). Cultured smooth muscle cells (23), theca cells (24) and adrenal glomerulosa cells (25) have also been reported to secrete prorenin, but not renin, into the medium.

Prorenin has always been considered to have no biological function except being the inactive precursor of renin. Several proteases have been shown to be capable of activating prorenin *in vitro* (26-29), but the enzyme responsible for *in vivo* processing of prorenin has not been identified. In man, under normal circumstances approximately 90 % of plasma renin is present in its inactive form (30). This percentage increases even further (up to 95-99 %) under certain conditions, for instance pregnancy (30) and diabetes mellitus complicated by microvascular complications (31). There is now good evidence

that the ovary is the main source of this increase in plasma prorenin in pregnant women (32). The elevated levels in diabetic subjects with microvascular complications are more difficult to explain. Again, an extrarenal source of prorenin may be involved. Renin is often within the normal range in these patients (31,33). In a recent study Franken and colleagues showed that the elevated levels of prorenin in diabetic subjects were related to the presence of retinopathy, particularly the proliferative type, and were independent of sex, age, duration of diabetes, blood pressure, neuropathy and blood levels of glucose and hemoglobin-A1c (33). Possibly the eye and other organs affected by microvascular complications are the unknown extrarenal sources of the increase in plasma prorenin in diabetic subjects. Elevated levels of plasma prorenin may perhaps be used as a marker for the activation of tissue renin-angiotensin systems (34).

In this view it is of interest to know that it has recently been shown that prorenin is able to become reversibly catalytically active, a process that is most likely due to a change in the conformation of the prorenin molecule and unfolding of the prosegment from a "closed" to an "open" position (35-37). Assuming that a dynamic equilibrium exists between the "open" and "closed" forms of prorenin (Figure 4), with the "open" form being catalytically active, one may hypothesize that under certain circumstances (receptor-binding, low pH), for instance at a local level, stabilization of the "open" form may occur, allowing prorenin to cleave angiotensin I from angiotensinogen. A (pro)renin receptor has so far not been identified.

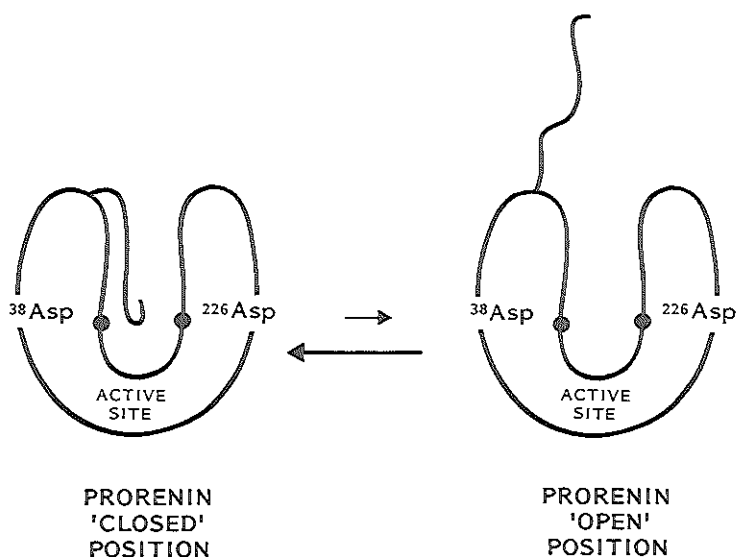


Figure 4. Hypothetical activation of prorenin by unfolding the prosegment.

The release of locally produced angiotensins into the circulation is, as discussed above, probably largely due to local uptake of kidney-derived renin. Local production of angiotensins not being released into the circulation (i.e. remaining local, for instance intracellular) may very well be due to locally synthesized renin or prorenin. Possibly kidney-derived renin is limited to the extracellular fluid (or bound to the cell-membrane) and cleaves angiotensinogen in plasma or interstitial fluid only, whereas locally synthesized prorenin or renin acts intracellularly only.

Function of local renin-angiotensin systems. The function of most, if not all, local renin-angiotensin systems is not yet clear. Locally produced angiotensin II may influence vascular tone, both through a direct effect via angiotensin II receptors and by enhancing norepinephrine release from noradrenergic nerve endings in blood vessels (38). It may also stimulate prostaglandin biosynthesis (39). In the heart, angiotensin II possesses positive inotropic (40) properties, while at the same time it may constrict coronary blood vessels. In the kidney, angiotensin II plays a role in regulating tubuloglomerular feedback, vasa recta blood flow and sodium reabsorption. In addition, mesangial cell contractility and glomerular filtration pressure may be influenced by local angiotensin production (41-43). An adrenal renin-angiotensin system probably regulates aldosterone biosynthesis (25). Similar to this effect of angiotensin II on adrenal steroid hormone synthesis, a local renin-angiotensin system in reproductive organs could be involved in the regulation of steroid hormone production (21,44). Brain angiotensin may contribute to the regulation of blood pressure, thirst and vasopressin as well as catecholamine release (45). Angiotensin II has recently been reported to have growth promoting effects (46-48); it stimulates neovascularization in the rabbit cornea model (49).

1.3 Blockers of the renin-angiotensin system

ACE inhibitors. ACE inhibitors, which block the formation of angiotensin II from angiotensin I, are now widely used in the treatment of hypertension and congestive heart failure. The design of captopril, the first commercially available ACE inhibitor, resulted from a rational and systemic study of the interaction of carboxypeptidase A and ACE with their substrates and inhibitors (50). Very productive research undertaken by several drug companies has led to the development of a whole range of new ACE inhibitors, some of which are now commercially available as well: enalapril, lisinopril, quinapril, perindopril, ramipril and cilazapril. Their hemodynamic profile is characterized by vasodilatation devoid of an accompanying tachycardia or a reduction in cardiac output (51). Not only were ACE inhibitors found to be effective for treating high renin-dependent forms of hypertension, but unexpectedly, they were also active in normal or low renin hypertension (52).

Long-term treatment by ACE inhibitors does not seem to completely suppress the circulating renin-angiotensin system, as plasma angiotensin II and aldosterone tend to return toward pretreatment values (53). This finding has led to speculations on the presence of local renin-angiotensin systems and their importance in the maintenance of high blood pressure. Tissue renin-angiotensin systems may explain why there is no correlation between plasma renin before treatment and the effect on blood pressure.

Renin inhibitors. ACE is not a specific enzyme: it can hydrolyze substrates other than angiotensin I, such as bradykinin, substance P, and enkephalins. The side-effects observed with all ACE inhibitors, such as angioneurotic edema, urticaria and cough, may be related to this lack of selectivity. The development of renin inhibitors was thus thought to be an interesting alternative to ACE inhibitors for 2 reasons: 1) the first step in the angiotensin II forming cascade, the hydrolysis of angiotensinogen by renin, is rate limiting; and 2) renin has a unique specificity for angiotensinogen as there is no other known substrate for this enzyme (54). Knowledge of the molecular structures of both renin and angiotensinogen and the molecular mechanisms of action of aspartyl proteases such as renin has speeded up the development of high-affinity inhibitors specific for renin. Most of the recently developed inhibitors are peptidic angiotensinogen analogues. Their oral availability is usually extremely low. The first reports showing that renin inhibitors are indeed effective blood pressure lowering agents in hypertensive subjects are now available (55,56). Their hypotensive effect was not accompanied by reflex tachycardia and outlasted the suppression of plasma angiotensin II levels. The latter finding again underlines the importance of tissue renin-angiotensin systems. However, much work, including improvement of oral availability and comparison of the blood pressure lowering effect of renin inhibitors with the blood pressure lowering effect of ACE inhibitors, still has to be done. The latter comparison could also shed light on the still unanswered question whether the hypotensive effect of ACE inhibitors is due solely to the blockade of angiotensin II formation or to the combination of a blockade of angiotensin II formation and a blockade of bradykinin degradation.

Angiotensin II antagonists. Even renin inhibitors may not totally eliminate the influence of the renin-angiotensin system, because angiotensin forming enzymes other than renin may cleave angiotensinogen as well (57). To get round this problem, recently attention has been drawn to the development of non-peptide angiotensin II receptor-antagonists. Since the early 1970s the angiotensin II receptor-antagonist saralasin, a peptide analogue of angiotensin II, has been available. Saralasin lowers blood pressure in conditions characterized by high renin levels (e.g. renovascular hypertension) (58), but this effect is only reached after intravenous administration. Saralasin has a short half life and displays partial agonist properties (59), thus complicating

the interpretation of results obtained with this peptide analogue. In the past few years several non-peptide angiotensin II antagonists without agonistic properties have been developed, and some were shown to lower blood pressure in spontaneously hypertensive rats after oral administration (60). At least 2 different subtypes of angiotensin II receptors, AT₁ (formally known as AII-1 or AII-B) and AT₂ (formally known as AII-2 or AII-A), have been characterized based upon their differential affinities for non-peptide angiotensin II antagonists currently available (61). The proto-typical antagonist of the AT₁ receptor is DuP 753 (62). The proto-typical antagonists of the AT₂ receptor are CGP 42112A, PD 123177, and PD 123319 (63-65). The functions of the AT₂ subtype receptors are presently not known (66), but those of the AT₁ receptor subtype have already been characterized extensively. AT₁ receptors mediate most of the well-known effects of angiotensin II, such as vasoconstriction (peripheral or centrally induced), smooth muscle contraction or hypertrophy, aldosterone secretion, and catecholamine release (67). Binding of angiotensin II to the AT₁ receptor activates phospholipase C, which hydrolyses phosphatidylinositol biphosphate to inositol 1,4,5-triphosphate (IP₃) and diacylglycerol (68-70). IP₃ mobilizes calcium from the endoplasmatic reticulum, while diacylglycerol activates protein kinase C. In hepatocytes angiotensin II was found not only to activate phospholipase C, but to inhibit adenylate cyclase as well (68). Both responses were mediated through the AT₁ receptor. Very recently two groups succeeded in cloning the adrenal (71) and vascular (72) AT₁ receptor subtype. Human studies with non-peptide angiotensin II receptor-antagonists, specifically those inhibiting the AT₁ receptor subtype (DuP 753), so far have been performed in normotensive subjects only (73). DuP 753 appeared to be a well-tolerated, orally active, potent and long-lasting antagonist of angiotensin II.

1.4 Aim of the thesis

The aim of this thesis was twofold. First, the regional metabolism and production of angiotensin I and angiotensin II were quantified *in vivo* in man and in anesthetized pigs. This was done by giving constant infusions of radiolabeled ¹²⁵I-angiotensin I into either a peripheral vein (man) or into the left cardiac ventricle (pig). Blood samples were taken under steady state conditions from various arterial and venous sampling sites. By measuring in each sample the levels of intact radiolabeled angiotensin I and angiotensin II and of endogenous angiotensin I and angiotensin II one can calculate both the degree of regional metabolism of angiotensin I and angiotensin II and the

amount of angiotensin I and angiotensin II that is generated in a certain vascular bed. Additional measurements of plasma renin activity (PRA) and calculations of the regional conversion of ^{125}I -angiotensin I to ^{125}I -angiotensin II make it possible to calculate the amount of locally generated angiotensin I that can be attributed to the action of circulating renin on circulating renin substrate and the amount of locally generated angiotensin II that can be accounted for by regional conversion of arterially delivered angiotensin I. Our data show clearly that part of angiotensin I in plasma is produced locally, probably in vascular tissue, and not in circulating plasma by PRA, and that most of the renin responsible for this local production is kidney-derived. Part of circulating angiotensin II also appeared to be produced locally, independent of plasma angiotensin I.

The second aim of this thesis was to investigate the existence of a local renin-angiotensin system in the eye. This was done based upon previous findings in diabetic subjects (33), which showed that a relationship existed between the high plasma prorenin levels and the presence of retinopathy, especially the proliferative type, in these subjects. Ocular fluid samples obtained from human subjects (both diabetic and non-diabetic) during eye surgery were examined for this purpose. Additionally, angiotensin and renin/prorenin measurements were performed in ocular tissue extracts from both cows and pigs. Human vitreous and subretinal fluid were found to contain prorenin levels too high to be explained by leakage from plasma only. This was not the case with angiotensinogen, angiotensin I or angiotensin II in these ocular fluids. Renin levels were in the low to undetectable range. In contrast, angiotensin I, angiotensin II, renin and prorenin levels in bovine and porcine ocular tissues were far too high to be explained by the presence of plasma in these tissues only. It seems therefore that indeed the eye contains its own renin-angiotensin system. The only component of the ocular renin-angiotensin system which is being released from the tissues into the vitreous fluid appears to be prorenin. Since prorenin was twofold higher in vitreous fluid obtained from eyes affected by proliferative diabetic retinopathy than in vitreous fluid from eyes of non-diabetic patients, one may hypothesize that the renin-angiotensin system is involved in the development of diabetic retinopathy. The final proof for the existence of an intraocular renin-angiotensin system was obtained by the demonstration of renin-, angiotensinogen- and ACE-mRNA expression in human eyes by the polymerase chain reaction.

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2. METABOLISM AND PRODUCTION OF ANGIOTENSIN I IN DIFFERENT VASCULAR BEDS IN SUBJECTS WITH HYPERTENSION

2.1 Summary

To study the metabolism and production of angiotensin I, highly purified mono-iodinated ^{125}I -angiotensin I was given by constant systemic intravenous infusion, either alone ($n = 7$) or combined with unlabeled angiotensin I ($n = 5$), to subjects with essential hypertension, who were treated with the angiotensin converting enzyme inhibitor captopril (50 mg b.i.d.). Blood samples were taken from the aorta and the renal, antecubital, femoral and hepatic veins. ^{125}I -angiotensin I and angiotensin I were extracted from plasma, separated by high performance liquid chromatography and quantitated by gamma counting and radioimmunoassay. Plasma renin activity was measured at pH 7.4. The plasma decay curves after discontinuation of the infusions of ^{125}I -angiotensin I and unlabeled angiotensin I were similar for the two peptides. The regional extraction ratio of ^{125}I -angiotensin I was $47 \pm 4\%$ (mean \pm SEM) across the forearm, $59 \pm 3\%$ across the leg, $81 \pm 1\%$ across the kidneys and $96 \pm 1\%$ across the hepatomesenteric vascular bed. These results were not different from those obtained for infused unlabeled angiotensin I. Despite the rapid removal of arterially delivered angiotensin I, no difference was found between the venous and arterial levels of endogenous angiotensin I across the various vascular beds, with the exception of the liver where angiotensin I in the vein was 50% lower than in the aorta. Thus, 50 - 90% of endogenous angiotensin I in the veins appeared to be derived from regional de novo production. The blood transit time is 0.1 - 0.2 min in the limbs and in the kidneys and 0.3 - 0.5 min in the hepatomesenteric vascular bed. This is too short for plasma renin activity to account for the measured de novo angiotensin I production. It was calculated that less than 20 - 30% in the limbs and in the kidneys and approximately 60% in the hepatomesenteric region of de novo produced angiotensin I could be accounted for by circulating renin. These results indicate that a high percentage of plasma angiotensin I may be produced locally (i.e., not in circulating plasma)

2.2 Introduction

Renin (EC 3.4.23.15), an aspartyl protease synthesized mainly by the kidneys,

acts on renin substrate (angiotensinogen) to form the decapeptide angiotensin I (ANG I), which is then converted to the octapeptide angiotensin II (ANG II) by angiotensin converting enzyme (ACE). All these components of the renin-angiotensin system are present in plasma. Plasma renin activity (PRA), which is the rate of ANG I formation *in vitro*, is considered to be a measure of the production of plasma ANG I *in vivo*. However, there is evidence from animal studies to suggest that the rate of angiotensin metabolism is too high for PRA to maintain the actual plasma levels of ANG I and II (1-3). Thus, part of plasma ANG I may be formed locally rather than in circulating plasma.

The source of renin involved in local ANG I production is not known. It is possible that the circulating renin-angiotensin system not only serves to deliver ANG I to the blood vessel wall for conversion to ANG II but that it also serves to deliver renin for local ANG I production. Through binding or uptake of plasma renin by vascular tissue or through local activation of plasma prorenin, the local concentration of renin may reach levels higher than in circulating plasma. Another possibility is that local production of ANG I depends on locally synthesized rather than systemically delivered renin.

Renin is synthesized by various tissues apart from the kidney and there is evidence for local angiotensin generation in some of these tissues (2,4). In culture, bovine and canine vascular smooth muscle cells and bovine endothelial cells synthesize ANG II and secrete this peptide into the medium (5,6). Isolated rat mesenteric arteries, when perfused with solutions free of renin and renin substrate, release ANG II into the perfusate (7), and in similar experiments, isolated rat hindlegs were found to release both ANG I and ANG II (8). Thus, part of ANG I and ANG II circulating in plasma may have been generated locally in the interstitium or in the cells of vascular tissue, and it has been postulated that a vascular renin-angiotensin system is important for the regulation of vascular tone and the maintenance of hypertension (9-12).

The production rate of angiotensins in different vascular beds cannot be simply derived from the regional blood flow and the arteriovenous concentration gradient, because a high rate of production may be matched by a high rate of metabolism. Studies using tracer amounts of radiolabeled angiotensins for measuring ANG I-II conversion and ANG I and II degradation are scarce because of the difficulty to measure the angiotensins separately from their metabolites and also because of the instability of some earlier radiolabeled iodine angiotensin preparations. More recent studies in dogs using highly purified mono-iodinated ^{125}I -ANG I have demonstrated that this preparation can be used for measuring intrarenal ANG I-II conversion and ANG I degradation (13,14). Mono-iodinated ^{125}I -ANG II is a potent, full agonist analog of ANG II (15,16).

The present study was undertaken to explore the possibility of measuring the rate of metabolism of ANG I in humans by intravenous infusion of mono-iodinated ^{125}I -ANG I. These infusions were given to hypertensive subjects at the time renal vein renin sampling was performed as part of the diagnostic

workup for renovascular hypertension while the subjects were treated with the ACE inhibitor captopril. ^{125}I -ANG I and ^{125}I -ANG II and endogenous ANG I and ANG II were isolated from aortic plasma and from renal, antecubital, femoral and hepatic venous plasma, and separated by high performance liquid chromatography (HPLC). The isolated radiolabeled angiotensins were quantitated in the gamma counter and the concentrations of endogenous angiotensins were measured by radioimmunoassay. Regional extraction of ANG I was calculated from measurements of the arterial and venous levels of ^{125}I -ANG I. From this and from measurements of the concentrations of endogenous ANG I, information could be obtained on ANG I production in different vascular beds. Parallel measurements of ^{125}I -ANG II provided information on ANG I-II conversion.

2.3 Materials and methods

Chemicals

[Ile⁵]-ANG-(1-10) decapeptide, [Ile⁵]-ANG-(1-8) octapeptide, [Ile⁵]-ANG-(2-8) heptapeptide and [Ile⁵]-ANG-(1-7) heptapeptide, which are denoted here as ANG I, ANG II, ANG III and ANG-(1-7) respectively, were obtained from Bachem, Bubendorf, Switzerland. [Ile⁵]-ANG-(2-10) nonapeptide was from Senn Chemicals, Dielsdorf, Switzerland. [Ile⁵]-ANG-(3-8) hexapeptide, [Ile⁵]-ANG-(4-8) pentapeptide and ANG-(1-4) tetrapeptide were from Peninsula Laboratories, Belmont, CA, USA. The World Health Organization (WHO) standards of ANG I and ANG II (86/536 and 86/538) were from the National Institute for Biological Standards and Control, London, UK. Methanol and ortho-phosphoric acid (both analytical grade) were from Merck, Darmstadt, Germany. Bovine serum albumin (BSA) was from Sigma, St. Louis, MO, USA. Human serum albumin (HSA) was from Merieux, Lyon, France. Water for HPLC was prepared with a Milli-Q system from Waters, Millford, MA, USA. The statine containing renin inhibitor CGP 29,287 was a kind gift of Dr. K. Hofbauer (Ciba-Geigy, Basel, Switzerland).

Preparation of angiotensin antisera

ANG I and ANG II antisera were produced in New Zealand White rabbits by intracutaneous injections of 330 μg of either ANG I or ANG II coupled to BSA with 1-ethyl-3-(3-dimethylamino)propyl-carbodiimide HCl and suspended in Freund's complete adjuvant. Booster injections of the conjugates suspended in Freund's incomplete adjuvant were given at 4 week intervals. The ANG I antiserum was used in the radioimmunoassay in a final dilution of 1 : 250.000. This antiserum gave a B/Bo ratio of 0.9 at 2 fmol ANG I and a B/Bo ratio of 0.5 at 20 fmol ANG I. The antiserum also reacted (100 %) with ANG-(2-10)

nonapeptide but did not react (less than 0.1 %) with ANG II, ANG III, ANG-(3-8) hexapeptide and ANG-(4-8) pentapeptide. The ANG II antiserum was used in a final dilution of 1 : 200.000. It gave a B/Bo ratio of 0.9 at 1.5 fmol ANG II and a B/Bo ratio of 0.5 at 16 fmol ANG II. This antiserum also reacted with ANG III (55 %), ANG-(3-8) hexapeptide (73 %) and ANG-(4-8) pentapeptide (100 %) but virtually not with ANG I (0.3 %) and ANG-(2-10) nonapeptide (0.2 %). The renin inhibitor CGP 29,287, in concentrations up to 100 μ M, did not react with either ANG I or ANG II antisera.

Blood sampling

Blood for angiotensin measurements was rapidly (7-13 sec) drawn by a plastic syringe containing the following inhibitors (0.5 mL inhibitor solution in 10 mL blood), 6.25 mM disodium EDTA, 1.25 mM 1,10-phenanthroline and 100 nM of the renin inhibitor CGP 29,287 (final concentrations in blood) and transferred into prechilled polystyrene tubes. These inhibitors were used in order to prevent ANG I generation, ANG I-II conversion and ANG I and II degradation during blood collection and handling of the samples. The blood samples were centrifuged at 3000 g for 10 min at 4 °C. Plasma was stored at -70 °C, extracted within two days and assayed within two weeks. The renin inhibitor CGP 29,287 is a renin substrate analog. At a concentration of approximately 1 nM it inhibits PRA by 50 % (17). We chose to use the inhibitor in a final concentration of 100 nM on the basis of experiments, in which we added the inhibitor in concentrations ranging from 0.1 to 1000 nM to a plasma sample with normal PRA (12 pmol/L · min) and to plasma samples with elevated PRA (77 and 260 pmol/L · min). ANG I formation during incubation was inhibited by 98 % at 37 °C and by 100 % at 20 °C, both at an inhibitor concentration of 100 nM.

Blood for PRA measurements was collected into polystyrene tubes containing EDTA (0.2 mL disodium EDTA in 10 mL blood, final concentration 6.25 mM). The samples were centrifuged at room temperature at 3000 g for 10 min, and plasma was stored at -20 °C.

Extraction of angiotensins

Solid phase extraction cartridges (SepPak C18, Waters, Millford, MA, USA) were conditioned with 4 mL methanol and then equilibrated with two times 4 mL of cold water. Thawed plasma samples (2 mL) were applied to the cartridges. After the bound peptides were washed two times with 4 mL cold water, they were eluted with 2 mL methanol and collected into conical polypropylene tubes. The methanol extracts were evaporated at 4 °C with a Savant Speed Vac concentrator (Savant Instruments, Farmingdale, NY, USA). Recovery of labeled and unlabeled ANG I and II from the SepPak cartridges was 95 - 98 %.

Separation of angiotensins by HPLC

Separation of the peptides in the methanol extracts was performed by HPLC using the method of Nussberger et al (18) with some modifications. We used a reversed phase Nucleosil C18 steel column of 250 x 4.6 mm and 10 μ m particle size equipped with a direct-connect guard column (Alltech, Eke, Belgium). The HPLC equipment consisted of a LKB 2150 pump, a 2152 controller, a 2122 fraction collector and a 2155 column oven (LKB, Bromma, Sweden). Samples were injected with a Rheodyne 7125 injection valve equipped with a 250 μ L loop (Rheodyne, Cotati, CA, USA). Mobile phase A was 0.085% orthophosphoric acid containing 0.02% sodium azide (pH 2.33). Mobile phase B was methanol. The flow was 1 mL/min and the working temperature was 45 °C. The column was calibrated with pure 125 I-labeled standards of ANG I and ANG II and with unlabeled ANG I, ANG II, ANG III and some of their metabolites. The vacuum dried plasma extracts were dissolved in 100 μ L HPLC solvent (65 % A/35 % B, vol/vol), centrifuged and injected with a 100 μ L Hamilton syringe (no. 1710, Hamilton, Bonaduz, Switzerland). Elution was performed as follows: 65 % A/35 % B from 0 to 9 min followed by a linear gradient to 45 % A/55 % B until 18 min. The eluate was collected in 20-sec fractions into polystyrene tubes coated with BSA.

The concentrations of 125 I-ANG I and 125 I-ANG II in the HPLC fractions were measured in the gamma counter for 20 min. The fractions containing ANG I and ANG II were pooled separately. They were neutralized with 0.5 M sodium hydroxide and evaporated in the Savant concentrator. The concentrations of ANG I and ANG II were measured by radioimmunoassay. Recoveries for the HPLC separation alone were 90 - 95 % for both labeled and unlabeled ANG I and II. The overall recoveries of ANG I and ANG II added to plasma samples were 85 ± 7 % and 84 ± 8 % (mean \pm SD, n = 6). Similar values were obtained for 125 I-ANG I and 125 I-ANG II. Results were not corrected for incomplete recovery.

Radioimmunoassay of angiotensins

The dried HPLC fractions were dissolved in 0.25 M Tris buffer, pH 7.4, containing 0.35 % BSA and 0.02 % sodium azide. They were then incubated with 125 I-ANG I or 125 I-ANG II and the appropriate antiserum at 4 °C for 3 days. Separation of bound and free angiotensin was achieved by charcoal adsorption, and both the supernatant and the pellet were counted to a counting error of less than 1 %. The concentrations of ANG I and ANG II (from Bachem) in the standard solutions of the radioimmunoassay were checked by comparison with the WHO ANG I and ANG II reference standards. The concentrations of angiotensins are expressed as picomoles per liter plasma. The lower limit of detection (2 x SD difference from Bo) was 1.2 fmol/tube for

ANG I and 0.7 fmol/tube for ANG II. The normal level of ANG I in antecubital venous plasma from healthy individuals in supine position was 20.9 pmol/L (geometric mean), range 12.3 - 42.4 pmol/L (n = 18). The normal level of ANG II in these individuals was 4.8 pmol/L (geometric mean), range 2.5 - 8.2 pmol/L (n = 18).

Measurement of plasma renin activity

PRA was measured by incubating of the plasma at 37 °C for 0, 15, 30 and 60 min. During the 60-min incubation pH varied from 7.1 at t = 0 min to 7.5 at t = 60 min. PRA measurements under these conditions gave results not different from measurements in the presence of 100 mM imidazol buffer, pH 7.4. With this buffer pH remained 7.4 during incubation. In order to prevent ANG I-II conversion, ANG I degradation, prorenin-renin conversion and bacterial growth, the following mixture of inhibitors was added to plasma before incubation (35 µL of inhibitor solution/mL plasma): 5 mM disodium EDTA, 3.4 mM 8-hydroxyquinoline sulfate, 2.4 mM phenylmethylsulfonylfluoride, 2.2 nM aprotinin and 1 mg/mL neomycin sulfate (final concentrations in plasma).

The inhibitors do not interfere with the reaction of renin with substrate. The kinetic constants, V_{\max} and K_m , of this reaction are not modified when these inhibitors are added to mixtures of highly purified human renin and human substrate. The measured levels of PRA are not different from those calculated from these kinetic constants and from the concentrations of renin and substrate measured in plasma (19). ANG I that was generated during incubation was quantitated by radioimmunoassay. ANG I generation in the PRA assay was linear in the first 30 min of incubation but in some samples the rate of ANG I generation was somewhat lower in the following 30 min. Only the first linear part of the ANG I generation curve was used for calculating PRA. The recovery of renin or ANG I added to plasma prior to assay was 98 - 100 %.

PRA is expressed as picomoles ANG I per liter plasma per minute of incubation (pmol/L · min). The normal level of PRA in antecubital venous plasma was 12.7 pmol/L · min (geometric mean), range 2.3 - 66.7 pmol/L · min (n = 74). This is in close agreement with the results obtained by the "antibody-trapping" technique (20,21).

Measurement of renal plasma flow

Renal plasma flow was determined by measurement of the clearance of ^{131}I -hippurate (Amersham Intl., Amersham, UK), which was given by constant intravenous infusion at the time the procedure of renal vein sampling was performed. Renal plasma flow was calculated by dividing the clearance of ^{131}I -hippurate by the renal extraction ratio (22).

Preparation of radiolabeled angiotensin I for intravenous infusion

ANG I was radiolabeled by the chloramine T method (23). All solutions were made in sterile pyrogen free water. The whole procedure was carried out under aseptic conditions. In short, 4 nmol ANG I in 20 μ L of 0.25 M sodium phosphate buffer, pH 7.5, was mixed with 1 mCi (37 MBq) 125 I-sodium iodide (Amersham Intl.). The reaction, which occurred at room temperature, was started by the addition of 20 μ L of 7.0 mM chloramine T. After 25-30 seconds the reaction was stopped with 20 μ L of 11.0 mM sodium metabisulfite. Free 125 I-iodide and labeled peptide were separated on a 5 x 50 mm Dowex-AG-I-X8 column (50-100 mesh, Bio-Rad, Richmond, CA, USA). Elution was carried out with 0.1 M acetic acid containing 0.1 % HSA. Labeled ANG I does not bind to the column under these conditions. Fractions of 0.5 mL were collected and the fractions containing labeled peptide were pooled.

To obtain mono-iodinated 125 I-ANG I, the labeled peptide was applied to a 1.6 x 100 cm Biogel P-4 column (200-400 mesh, Bio-Rad) and eluted with 0.05 M acetic acid containing 0.1 M NaCl and 0.1 % HSA. Fractions of 2.5 mL were collected and counted in a gamma counter. The 125 I-ANG I was eluted from the Biogel P-4 column in two separate peaks. The first peak represented mono-iodinated 125 I-ANG I and the second peak consisted of di-iodinated 125 I-ANG I. Fractions from the first peak were pooled, sterilized by filtration through a 0.22 μ m Millipore membrane filter (Waters, Millford, MA, USA) and stored at -20 °C until use. The specific radioactivity of the 125 I-ANG I preparation was approximately 3.6×10^6 cpm/pmol (74 kBq/pmol). To obtain information about the purity of the mono-iodinated 125 I-ANG I, a sample of the pooled fractions of the first peak was injected into the HPLC column. The HPLC elution profile of the radioactive material indicated that 98 % consisted of mono-iodinated 125 I-ANG I.

Subjects

Ten subjects with hypertension (four men, six women mean age 48, range 26-67 years) were studied at the time they were undergoing renal vein renin sampling followed by renal angiography for diagnostic purposes. The renal vein renin ratio was normal (< 1.5) as was the renal angiogram. Blood pressure in the outpatient clinic was repeatedly above 160/100 mm Hg despite antihypertensive medication. Routine urine analysis, serum creatinine and serum electrolytes were normal, and the subjects were considered to have essential hypertension. Seven subjects received an infusion of 125 I-ANG I, and three subjects received an infusion of both 125 I-ANG I and ANG I. Two additional subjects with essential hypertension (one woman 47 years old and one man 58 years old) were studied at the end of a 24-hour period of intra-arterial blood pressure recording. Renal vein renin sampling and renal angiography had been performed some months before and were normal. These

two subjects received an infusion of both $^{125}\text{I-ANG I}$ and ANG I .

All subjects received 5 mL Lugol's solution/day from 2 days before to 4 days after the angiotensin infusion. The subjects were treated with the ACE inhibitor captopril (50 mg p.o., b.i.d.). The infusion studies were performed 4-6 hours after administration of the morning dose of captopril.

Infusion protocol

$^{125}\text{I-ANG I}$ was infused at a rate of approximately 3.0×10^6 cpm/min, and ANG I at a rate of approximately 1.2 nmol/min for 20 min. An indwelling needle for blood sampling was placed in the antecubital vein of the left arm. After insertion of the catheters into the abdominal aorta and the inferior caval vein via the femoral artery and vein by the Seldinger technique, angiotensin was infused via the antecubital vein of the right arm. A blood sample was taken from the abdominal aorta before the infusion. Ten minutes after the infusion had been started, samples were taken from the aorta, the hepatic vein and the antecubital vein of the left arm. In the following 10 minutes, a second and third series of samples were taken simultaneously from the aorta, the antecubital vein and a renal vein, first from one kidney and then from the other. A sample from the femoral vein was taken shortly before discontinuation of the infusion. The arterial plasma levels of $^{125}\text{I-ANG I}$, $^{125}\text{I-ANG II}$, ANG I and ANG II remained constant between 8 and 20 min after the start of the infusion (coefficient of variation < 4 %). For measuring the elimination half life of $^{125}\text{I-ANG I}$ and infused ANG I , samples were taken from the antecubital vein or from the aorta at approximately 0.5, 1, 2, 3 and 4 min after the infusion had been stopped. After the last blood sample had been collected the radiocontrast injection for renal angiography was given.

The two subjects studied some months after renal vein catheterization received an i.v. infusion of $^{125}\text{I-ANG I}$ and ANG I in one arm, and blood samples were taken from the brachial artery and the antecubital vein of the other arm before the infusion and at 10, 15 and 20 min after the start of the infusion. For measuring the elimination half life of the infused peptides, samples were taken from the brachial artery at the above mentioned times after the infusion had been stopped.

Urinary excretion of radioactivity was followed over a period of 96 hours. Ninety percent of the administered radioactivity was excreted within 24 hours and 98 percent was recovered in the 96 hour period. The calculated exposure to radioactivity was 0.24 μGy (or 0.6 mRad) from the $^{125}\text{I-ANG I}$ infusion and 12 μGy (or 30 mRad) from the $^{131}\text{I-hippurate}$ infusion (24). The protocol was approved by the Hospital Ethical Review Committee.

Calculations

The regional extraction ratio of $^{125}\text{I-ANG I}$ was calculated as follows:

$$ER = 1 - [^{125}\text{I-ANG I}]_{\text{ven}} / [^{125}\text{I-ANG I}]_{\text{art}}$$

The same formula was used for the calculation of the extraction ratio of infused ANG I. In this case the plasma level of ANG I before infusion was subtracted from the level during infusion.

The venous plasma level of ANG I derived from de novo regional production was calculated as follows:

$$[\text{ANG I}]_{\text{ven}} \text{ from } \text{de novo production} = [\text{ANG I}]_{\text{ven}} - (1-ER) \cdot [\text{ANG I}]_{\text{art}}$$

For the renal vascular bed it was possible to calculate the net de novo production rate of ANG I because we determined not only the renal extraction ratio of $^{125}\text{I-ANG I}$ and the renal arteriovenous difference in endogenous ANG I but also the renal extraction ratio and clearance of $^{131}\text{I-hippurate}$. The renal extraction ratio of $^{131}\text{I-hippurate}$ was calculated as follows:

$$ER_{\text{hip}} = 1 - [^{131}\text{I-hippurate}]_{\text{ven}} / [^{131}\text{I-hippurate}]_{\text{art}}$$

ER_{hip} was 0.72 ± 0.02 for the left kidney and 0.75 ± 0.03 for the right kidney ($p > 0.05$). If it is assumed that the renal plasma flow is equal on both sides, the single kidney plasma flow can be calculated as follows:

$$\text{Single kidney plasma flow} = \text{clearance of } ^{131}\text{I-hippurate} / (ER_{\text{hip left}} + ER_{\text{hip right}})$$

The single kidney net de novo production of ANG I is then obtained by multiplication of single kidney plasma flow by the renal venous plasma level of intrarenally de novo produced ANG I.

2.4 Results

HPLC separation of angiotensin peptides

Under the conditions specified in Methods, satisfactory separations were obtained between ANG I, ANG II and their metabolites as well as their radiolabeled counterparts (Fig. 1). Apart from the metabolites shown in Fig. 1, we also tested $^{125}\text{I-ANG-(1-4)}$ and $^{125}\text{I-ANG-(1-7)}$. These metabolites had a retention time of approximately 4 min. Assessment of injection to injection and day to day variability of the retention times of $^{125}\text{I-ANG I}$ and $^{125}\text{I-ANG II}$ demonstrated excellent stability of the chromatographic conditions. Separation patterns for radiolabeled angiotensins of plasma extracts from patients who had received an intravenous infusion of $^{125}\text{I-ANG I}$ showed four peaks with retention times of 4.2, 9.9, 13.0 and 16.2 min (Fig. 2). Until now, the material in the first peak has not been identified. The other peaks were $^{125}\text{I-ANG II}$, $^{125}\text{I-ANG (2-10)}$ and $^{125}\text{I-ANG I}$. The retention times of $^{125}\text{I-ANG I}$ and $^{125}\text{I-ANG II}$ differed from the retention times of any of the metabolites we tested, and the difference was sufficient to obtain adequate separations.

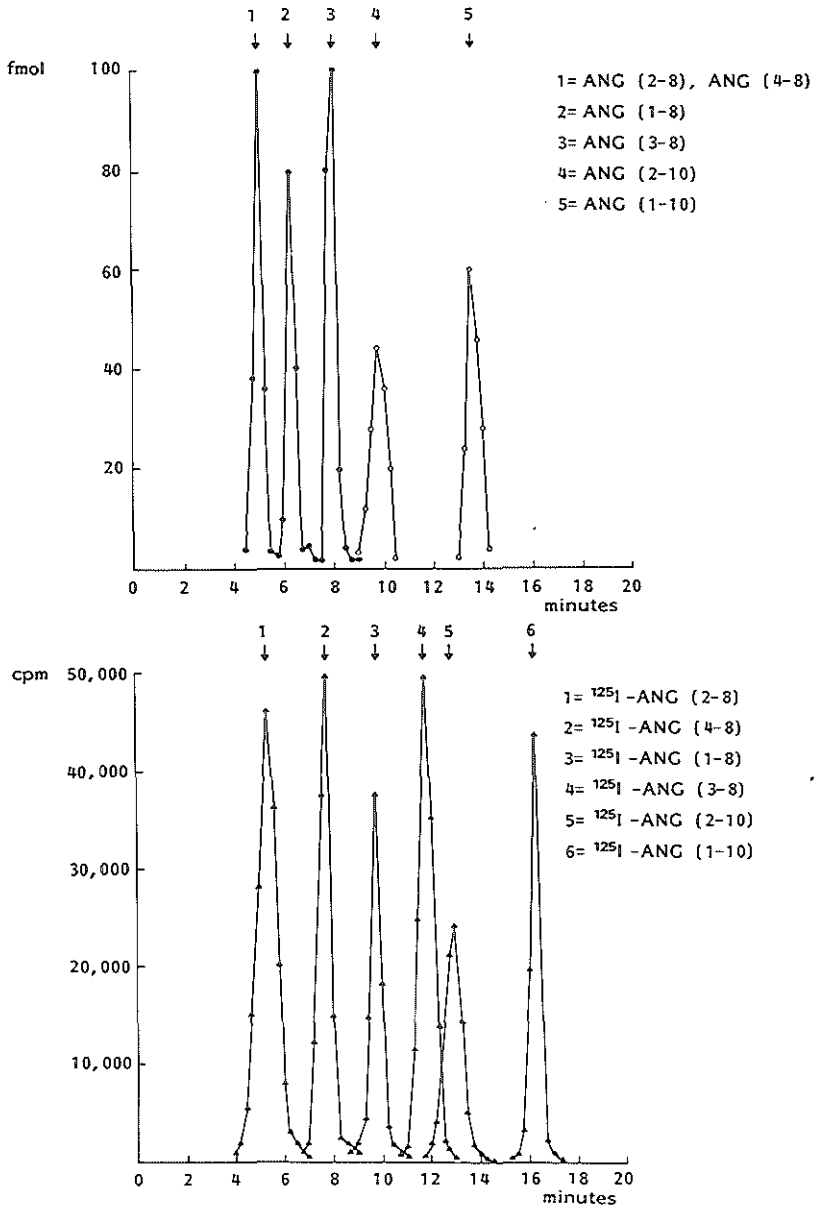


Figure 1. Line graph showing separation of standard angiotensin peptides (upper panel) and ¹²⁵I-labeled angiotensin peptides (lower panel) by reversed phase high performance liquid chromatography. Detection by radioimmunoassay with angiotensin I antiserum (peaks 4 and 5) or with angiotensin II antiserum (peaks 1-3) (upper panel) or by gamma counting (lower panel).

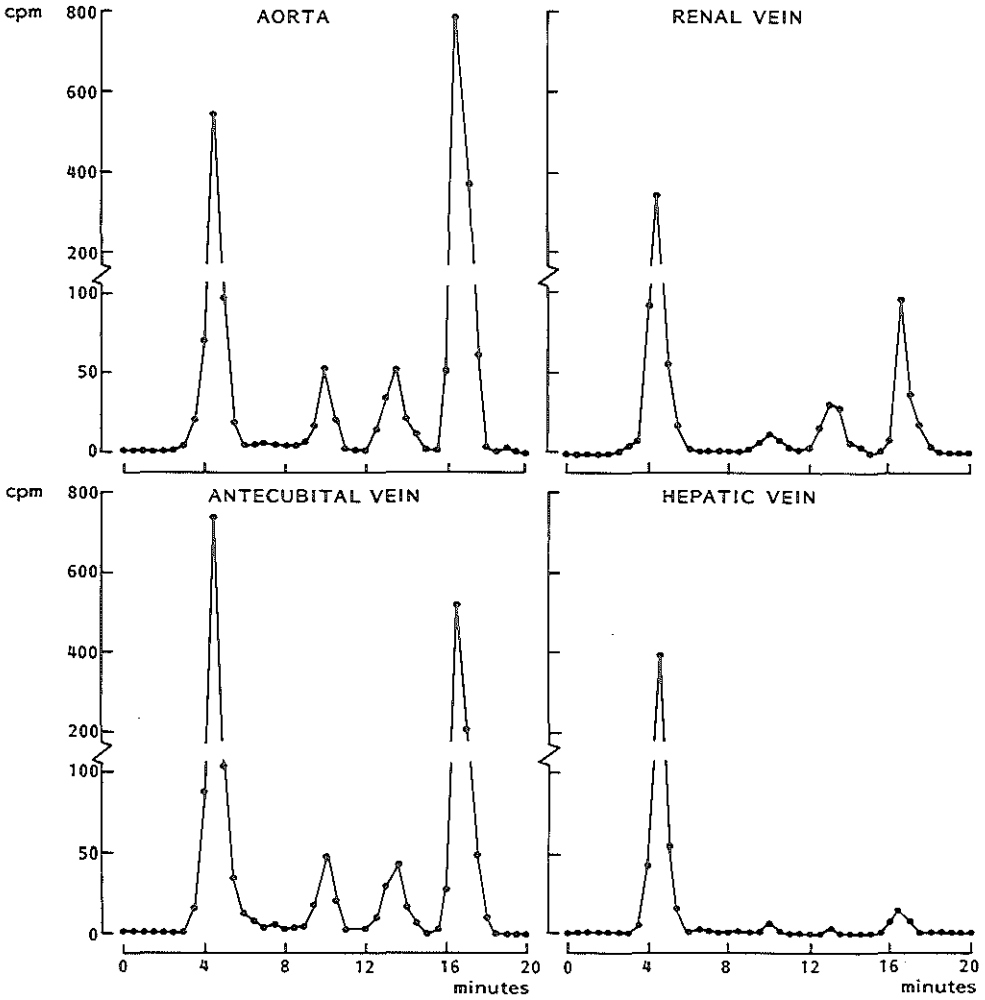


Figure 2. Line graph showing HPLC separation of ^{125}I -labeled angiotensins in plasma of subject who received a constant intravenous infusion of ^{125}I -ANG I. Stationary and mobile phases as described under Fig. 1. Detection by gamma counting. Peaks with retention times of 9.9, 13.0 and 16.2 min correspond with ^{125}I -ANG (1-8) = ^{125}I -ANG II, ^{125}I -ANG (2-10) and ^{125}I -ANG (1-10) = ^{125}I -ANG I respectively. Material in the first peak with a retention time of 4.2 min was not identified.

Table 1. Plasma levels of ^{125}I -ANG I, ANG I and plasma renin activity during constant intravenous infusion of both ^{125}I -ANG I and ANG I.

	PRA			Infusion Rates	
	pmol ANG I/ L · min	^{125}I -ANG I 10^3 cpm/L	ANG I pmol/L	^{125}I -ANG I 10^6 cpm/min	ANG I nmol/min
<u>Subject 1</u>				3.14	1.38
Before infusion					
Artery	7.6	-	5.1		
During infusion					
Artery	6.8	1536	723		
Hepatic Vein	-	96	52		
Left Renal Vein	-	260	105		
Right Renal Vein	-	258	106		
Femoral Vein	-	901	404		
<u>Subject 2</u>				2.70	1.59
Before infusion					
Artery	5.2	-	9.5		
During infusion					
Artery	4.3	1021	665		
Hepatic Vein	-	32	32		
Left Renal Vein	-	215	124		
Right Renal Vein	-	242	130		
Antecubital Vein	-	503	313		
Femoral Vein	-	518	319		
<u>Subject 3</u>				3.02	0.68
Before infusion					
Artery	3.8	-	6.8		
During infusion					
Artery	4.0	1254	307		
Hepatic Vein	-	53	11.9		
Left Renal Vein	-	235	58		
Right Renal Vein	-	325	79		
Antecubital Vein	-	528	130		
Femoral Vein	-	495	143		
<u>Subject 4</u>				3.05	1.54
Before infusion					
Artery	63.2	-	50		
During infusion					
Artery	57.3	696	613		
Antecubital Vein	-	401	439		
<u>Subject 5</u>				3.50	1.50
Before infusion					
Artery	62.2	-	76		
During infusion					
Artery	47.2	810	382		
Antecubital Vein	-	500	236		

In subjects 1-3, "Artery" is the abdominal aorta. In subjects 4-5, "Artery" is the brachial artery.

Regional extraction of angiotensin I

An important question is whether or not the rate of metabolism of ^{125}I -ANG I in our subjects differed from the metabolism of unlabeled ANG I. To answer this question unlabeled ANG I was infused together with ^{125}I -ANG I. Results obtained with these combined infusions are summarized in Table 1. Because of ACE inhibitor treatment, the infusions of ANG I did not result in a significant reduction of PRA. The preinfusion level of ANG I was therefore taken as a measure of the contribution of endogenous ANG I to the level of ANG I during infusion of this peptide. Thus, for calculation of the regional extraction of infused ANG I, the preinfusion level of ANG I was subtracted from the level during infusion. From measurements of the arterial and venous plasma levels of ^{125}I -ANG I and unlabeled ANG I it appeared that, during infusion of the two peptides, the regional extraction ratios for labeled and unlabeled ANG I were not different (Fig. 3). In most patients the plasma concentrations of ^{125}I -ANG I and unlabeled ANG I fell mono-exponentially after discontinuation of the infusion. Some curves, however, showed a delay. This delay can be explained, at least in part, by the passage time of blood from the infusion site to the sampling site but the delay could also represent the exit of ^{125}I -ANG I and ANG I from tissue sites. Estimates of the elimination half life of ^{125}I -ANG I and ANG I were therefore made from the steepest straight portion of the decay curves and the results were not different for the two peptides (Fig. 4). The estimated elimination half life of ^{125}I -ANG I, which was determined in all subjects, was 0.73 ± 0.06 min (mean \pm SEM).

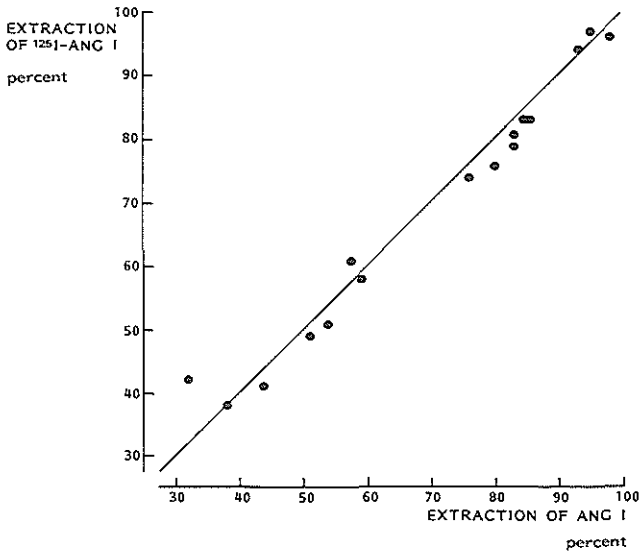


Figure 3. Scatter plot showing comparison of the regional extraction of ^{125}I -ANG I with that of ANG I during constant intravenous infusion of the two peptides in five subjects. Data are expressed as percentage of arterial plasma concentration.

Table 2. Arterial and venous plasma levels of angiotensins in seven hypertensive subjects during constant intravenous infusion of ^{125}I -ANG I.

Sampling site	^{125}I -ANG I 10 ³ cpm/L	ANG I pmol/L	^{125}I -ANG II 10 ³ cpm/L	ANG II pmol/L
Aorta	661 (439-1335)	34.0 (8.9-142)	133 (86-234)	2.1 (0.8-5.0)
Antecubital vein	341* (234-872)	34.1 (7.8-134)	114* (72-199)	2.3 (1.1-5.8)
Left renal vein	123* (62-225)	33.0 (9.0-124)	23* (14-33)	0.8* (0.4-1.5)
Right renal vein	129* (61-191)	31.7 (9.1-120)	21* (11-38)	0.8* (0.4-1.9)
Femoral vein	265* (158-617)	33.9 (7.8-134)	109* (62-199)	1.8 (0.8-4.3)
Hepatic vein (n=5)	31* (22-44)	17.2*(4.9-59)	5* (2-7)	0.4* (0.2-0.6)

^{125}I -ANG I infusion rate was $3.28 (2.46-4.31) \times 10^6$ cpm/min. Values are geometric mean (range). * significantly different from corresponding aortic levels ($p < 0.02$), Wilcoxon signed rank test.

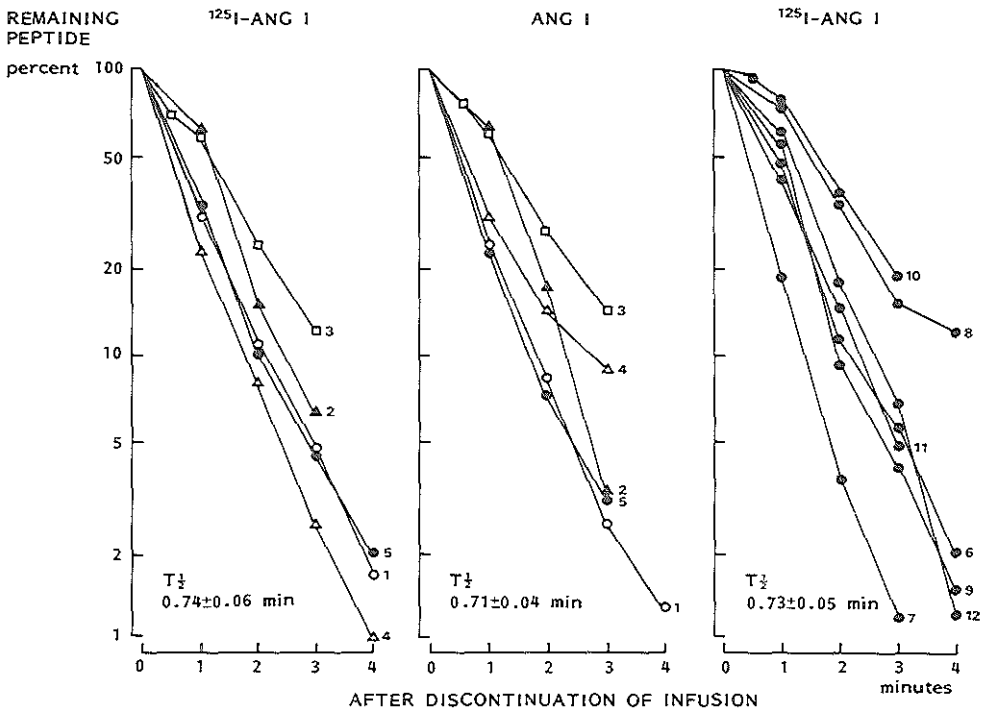


Figure 4. Line graphs showing in vivo elimination of ^{125}I -ANG I and ANG I after discontinuation of a constant intravenous infusion of both ^{125}I -ANG I and ANG I (left and middle panel) or a constant intravenous infusion of ^{125}I -ANG I only (right panel). Plasma ANG I levels in the middle panel were corrected for preinfusion ANG I levels. Each patient in left and middle panel is denoted by a different symbol and number. Numbers in these panels refer to subjects in Table 1. Blood was sampled from the aorta in subjects 1, 3, 6 and 7, from the brachial artery in subjects 4 and 5, or from the antecubital vein in subjects 2 and 8-12.

PRA and angiotensin levels vary widely among subjects with essential hypertension. This was also the case in our study. PRA in the aorta ranged from 3.8 - 122 pmol/L · min. The plasma levels of ^{125}I -ANG I and II and endogenous ANG I and II in the aorta and in the veins of the kidneys, forearm, leg and liver during infusion of ^{125}I -ANG I, are presented in Table 2.

Regional extraction of arterially delivered ^{125}I -ANG I was not influenced by the infusion of unlabeled ANG I (Fig. 5). The extraction ratio of ^{125}I -ANG I was $80 \pm 2\%$ for the right kidney, $82 \pm 1\%$ for the left kidney, $47 \pm 4\%$ for the forearm, $59 \pm 3\%$ for the leg and $96 \pm 1\%$ for the hepatomesenteric vascular bed. We did not measure the ^{125}I -ANG I levels in mixed venous plasma but an approximation was made by assuming an overall extraction of systemically delivered ^{125}I -ANG I of 60%. The mixed venous level upstream from the site where the infused ^{125}I -ANG I enters the central venous compartment can then be obtained by dividing the infusion rate by the cardiac output, which was taken to be 3 L plasma per minute per 1.73 m^2 body surface area, and by adding to this quotient 40% of the arterial level of ^{125}I -ANG I. In this way it was calculated that approximately 30 - 40% of the ^{125}I -ANG I delivered to the cardiopulmonary circulation was extracted by this vascular bed. From this and from a systemic extraction of 60% a whole body extraction of about 75% can be calculated. Because of this high extraction rate, the "true" half life of ANG I in the body must be shorter than the whole body transit time of blood (about 0.5 min). The "true" half life of ANG I must therefore be shorter than the half life (0.73 min) we estimated from the plasma ^{125}I -ANG I and ANG I decay curves after the infusion of these peptides had been stopped. This difference again may represent the exit of ^{125}I -ANG I and ANG I from tissue sites.

Regional angiotensin I-II conversion

We did not measure the ^{125}I -ANG II levels in mixed venous plasma, but they are likely to be lower than in aortic plasma, because the ^{125}I -ANG II levels we measured in the renal, antecubital, femoral and hepatic veins were lower than in the aorta (Table 2) and because observations in animals demonstrated extraction of ANG II in the circulation of the head (1). One may therefore assume that ^{125}I -ANG I-II conversion had occurred in the cardiopulmonary region, presumably the lungs, despite the fact that our subjects were on ACE inhibitor treatment. Since the extraction of ^{125}I -ANG II by the various vascular beds was not measured, it was not possible to calculate the degree of ^{125}I -ANG I-II conversion in these regions.

Regional production of angiotensin I

In spite of the rapid extraction of arterially delivered ANG I, the arterial and venous plasma levels of endogenous ANG I across the kidneys and the limbs

showed little difference (Fig. 5). The vein-to-artery ANG I concentration ratio was 0.98 ± 0.05 across the kidneys, 1.01 ± 0.03 across the forearm, and 1.00 ± 0.03 across the leg. ANG I concentration in the hepatic vein was lower than in the aorta. The vein-to-artery ANG I concentration ratio across the hepatomesenteric vascular bed was 0.52 ± 0.03 . From these data and from the regional extraction ratio of ANG I, the venous concentration of ANG I derived from regional de novo production could be calculated. Figure 6 clearly illustrates the point that a high proportion of venous ANG I originated from regional de novo production. For the renal vascular bed it was possible to calculate the net de novo production rate of ANG I because renal plasma flow was measured. Single kidney plasma flow was 0.283 ± 0.020 L/min and single kidney net de novo production of ANG I was calculated to be 7.1 pmol/min (geometric mean), range 2.4 - 29.5 pmol/min.

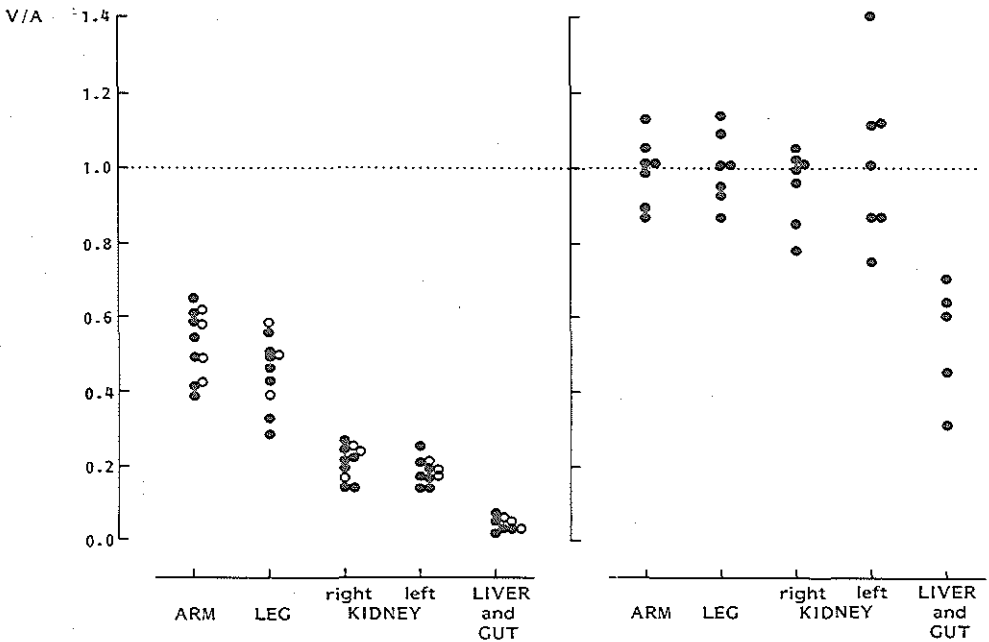


Figure 5. Scatter plots showing vein-to-artery ratios of $^{125}\text{I-ANG I}$ (left panel) and endogenous ANG I (right panel) across various vascular beds during constant i.v. infusion of $^{125}\text{I-ANG I}$. Open circles represent data obtained during infusion of both $^{125}\text{I-ANG I}$ and ANG I.

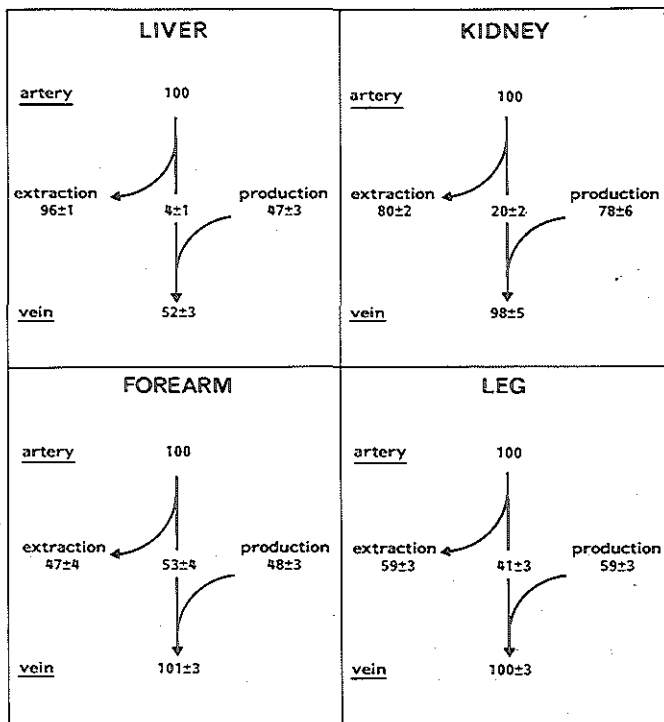


Figure 6. Schematic drawing showing regional venous plasma concentration of ANG I originating from *de novo* production of ANG I in various vascular beds. Data (mean \pm SEM) are expressed as a percentage of the arterial plasma concentration of ANG I.

Table 3. Contribution of plasma renin activity to venous levels of ANG I in seven hypertensive subjects.

	Kidney	Forearm	Leg	Liver and Gut*
Mean blood transit time (min)	0.17	0.17	0.17	0.42
Arterial PRA (pmol ANG I/L · min)	24.0 (7.3-117)	24.2 (6.7-122)	24.5 (8.9-119)	see under 'Forearm'
Venous PRA (pmol ANG I/L · min)	30.6 (7.3-211)	25.1 (7.2-119)	26.3 (8.1-135)	23.0 (6.2-82)
Venous ANG I originating from <i>de novo</i> production (pmol/L)	25.7 (7.4-102)	16.1 (4.3-63)	20.0 (4.9-71)	15.8 (4.5-53)
Venous ANG I produced by circulating PRA (pmol/L)	5.3 (1.3-33)	4.2 (1.2-20)	4.1 (1.1-18)	8.9 (2.5-42)
% of <i>de novo</i> produced ANG I	21 \pm 3	27 \pm 3	22 \pm 3	62 \pm 9

Values are geometric mean (range) or mean \pm SEM. Mean blood transit times are from the literature (25,26). Three arterial samples were collected from the abdominal aorta during the steady state phase: sample 1 simultaneously with the antecubital vein and hepatic vein samples, sample 2 simultaneously with a left or a right renal vein sample, and sample 3 simultaneously with a renal vein sample on the opposite site. The femoral vein sample was taken immediately after aortic sample 3. The left and right renal vein levels of PRA and ANG I were averaged as were the two corresponding aortic levels of PRA. * n=5.

Contribution of plasma renin activity to regional production of angiotensin I

The mean transit time of blood is 0.1 - 0.2 min in the kidneys and the limbs and 0.3 - 0.5 min in the hepatomesenteric vascular bed (25,26). From this and the measured levels of PRA, we can calculate the contribution of circulating PRA to the total regional production of ANG I. In the classical concept of the renin-angiotensin system circulating renin should fully account for this production. The calculations, however, showed that circulating PRA only accounted for less than 20 - 30 % of ANG I production in the kidneys, the forearm and the leg. In the hepatomesenteric vascular bed circulating PRA accounted for about 60 % of the ANG I production in this region (Table 3).

2.5 Discussion

Circulating ANG I and ANG II are rapidly metabolized by the action of peptidases. ANG I is converted to what is considered to be the most important biologically active endproduct of the renin-angiotensin system, ANG II. Both ANG I and ANG II are degraded into smaller inactive peptides. The term "metabolism" is used here to denote both conversion and degradation. There are very few reports on measurements of ANG I metabolism in intact animals (13,14,27), and we are not aware of any such data in humans.

The present study addressed the question whether in humans the rate at which ANG I is generated by the reaction of circulating renin with circulating renin substrate, is sufficiently high to maintain the actual plasma level of ANG I in the face of the rapid metabolism of this peptide. We studied ANG I metabolism in different vascular beds by measuring the extraction of arterially delivered ¹²⁵I-ANG I during constant intravenous infusion of this peptide. Again, the term "extraction" includes the processes of conversion and degradation. In our hypertensive subjects on captopril treatment the regional extraction of ¹²⁵I-ANG I by the kidneys, the limbs and the hepatomesenteric vascular bed was not different from the extraction of simultaneously infused unlabeled ANG I. Also, the plasma decay curves after discontinuation of the ¹²⁵I-ANG I and ANG I infusions, were very similar. This supports the validity of using our measurements of ¹²⁵I-ANG I metabolism for estimation of the metabolism and production of endogenous ANG I, at least in subjects on ACE inhibitor treatment. With such treatment, ANG I metabolism is mainly due to degradation into smaller inactive fragments. Apparently there is little or no difference in the rate of degradation between labeled and unlabeled ANG I. Whether the conversion rates of the two peptides are also similar has to be established by measurements in subjects not on ACE inhibitor treatment.

Extraction of arterially delivered ANG I by the vascular beds we studied ranged from 47 to 96 %. The plasma levels of ACE and angiotensinase activity are by far too low to account for the rapid removal of ANG I. Thus, it

appears that most of ANG I metabolism takes place at the surface of blood vessels or in the vascular tissue. In studies in dogs, ^{125}I -ANG I was infused into the renal artery and it was found that about 80 % was removed during a single passage of blood through the kidney (13). This was for the most part due to degradation rather than conversion. Our results in humans are in agreement with these data.

From the regional extraction ratios of ANG I and from the arteriovenous differences in ANG I it was calculated that approximately 50 - 90 % of ANG I in the regional veins was not derived from arterially delivered ANG I but from regional de novo production. In the dog it was found that 70 - 80 % of ANG I in the renal veins was derived from de novo production in the kidney (13), which is similar to what we found for the human kidney.

With the exception of the hepatomesenteric vascular bed, where the plasma level of ANG I in the hepatic vein was found to be much lower than in the aorta, there was little or no arteriovenous difference in ANG I. Thus, the rapid removal of plasma ANG I was matched by a high rate of production. By taking into consideration the blood transit time in the various vascular beds, one can easily see that circulating levels of PRA were not sufficient to account for this high rate of ANG I production. The mean transit time of blood in the kidneys and the limbs is 0.1 - 0.2 min or less (25,26). From this and from the measurements of PRA, the extraction of arterially delivered ANG I and the arteriovenous difference in the plasma levels of ANG I, it was calculated that less than 20-30 % of ANG I produced in the kidneys, the forearm and the leg could be accounted for by the generation of ANG I by circulating PRA (Table 3). For the renal vascular bed it was possible to calculate the net de novo production rate of ANG I, because renal plasma flow was known. The single kidney intravascular plasma volume is approximately 40 mL (25). From this and from the level of PRA in the renal vein it was calculated that less than 10 % of ANG I produced in the kidney could be accounted for by circulating PRA. The calculated contribution of PRA is a maximum value, because it is quite likely that part of regionally produced ANG I is metabolized before it can reach the veins. The blood transit time in the hepatomesenteric vascular bed is 0.3 - 0.5 min (26), and in this region the ANG I production could be largely accounted for by circulating PRA.

The conclusion of this study, that a high proportion of ANG I in plasma appears not to be generated by the action of circulating renin on circulating renin substrate, depends on the validity of using PRA in vitro as a precise measure of the generation of ANG I by plasma in vivo. We therefore performed the PRA assay at near physiological pH under conditions that did not cause denaturation of renin, activation of prorenin and conversion and degradation of ANG I and did not interfere with the reaction of renin with its substrate.

From the data presented in Table 2 it is evident that a discrepancy exists between the ^{125}I -ANG I / ^{125}I -ANG II and ANG I/ANG II ratios. One

possibility might be that the ^{125}I -ANG II peak is contaminated by some metabolite. We are confident, however, that the ^{125}I -ANG II peak is not contaminated by the possible aminopeptidase breakdown products, ^{125}I -ANG-(2-8), ^{125}I -ANG-(3-8), ^{125}I -ANG-(4-8) and ^{125}I -ANG-(2-10), nor by the prolyl- or neutral-endopeptidase breakdown products, ^{125}I -ANG-(1-4) and ^{125}I -ANG-(1-7). However, we did not measure the retention time of ^{125}I -ANG-(1-9), a possible product of carboxypeptidase attack on ^{125}I -ANG I. We did not observe any heterogeneity in the ^{125}I -ANG II peak when plasma samples were analyzed under different chromatographic conditions, also indicating that ^{125}I -ANG II was the only radiolabeled peptide in this peak. Because we did not compare the rates of metabolism of ^{125}I -ANG II and ANG II, our study only provides semi-quantitative information of ANG I-II conversion in the various vascular beds. At any rate our measurements of the arterial and venous plasma concentrations of ^{125}I -ANG I and ^{125}I -ANG II indicate that, despite ACE inhibitor treatment, ANG I-II conversion had occurred in the cardiopulmonary vascular bed, presumably the lungs.

The results of the present study add new evidence to previously published data supporting the hypothesis that a major fraction of ANG I in circulating plasma is produced locally rather than in circulating plasma itself. The local production of ANG I may depend on synthesis of renin *in situ*. Binding or uptake of plasma renin or prorenin and subsequent activation of prorenin and local production of ANG I are other possibilities. Whatever the exact mechanism, vascular production of ANG I may contribute to the control of vascular tone and this contribution may be independent, at least in part, of the circulating levels of renin and renin substrate.

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3. METABOLISM OF ANGIOTENSIN I BY DIFFERENT TISSUES IN THE INTACT ANIMAL

3.1 Summary

To quantify in the intact animal the regional conversion of angiotensin (ANG) I to ANG II and its degradation to peptides other than ANG II, mono-iodinated ^{125}I -ANG I (alone and in combination with unlabeled ANG I) was given to anesthetized pigs by constant infusion into the left cardiac ventricle. Blood samples were taken to measure the plasma levels of ^{125}I -ANG I, ^{125}I -ANG II, ANG I and ANG II. Sampling sites were: aorta, right and left atrium, pulmonary artery, great cardiac vein and femoral, renal, jugular and ear lobe veins. The metabolic degradation rate in the various vascular beds was the same for ^{125}I -ANG I and ANG I, but the conversion rate was 2 times higher for ^{125}I -ANG I. ANG I appeared to be distributed over a compartment with a size corresponding to 18-32 (mean 24) % of total body weight. Whereas the vein/artery ^{125}I -ANG I ratios across the systemic vascular beds were < 0.5 , the total radioactivity ratios were close to 1.0, indicating that elimination of ^{125}I -ANG I was due to rapid metabolism. After ACE inhibitor treatment, fractional ANG I metabolism (the fraction of arterially delivered ANG I that was metabolized during a single passage of blood) was 10 % in the lungs (conversion 4 %), as compared to 56 % in the combined systemic vascular beds (conversion 1 %). Fractional ANG I metabolism during ACE inhibition was 93 % in the kidney, 50-70 % in myocardium, skeletal muscle, head and skin, and 21 % in the left and 14 % in the right cardiac cavity. Without ACE inhibition, fractional ANG I metabolism was 29 % in the lungs (conversion 25 %), 49 % in the combined systemic vascular beds (conversion 10 %), 38 % in the left cardiac cavity (conversion 11 %) and 14 % in the right cardiac cavity (conversion 0 %). Thus, vascular beds other than the lungs make an important contribution to the conversion of circulating ANG I. ANG II formation *in vivo* appears to depend on ACE, but ANG I-II conversion does not seem to be a critical intermediary step in ANG I metabolism. Reduction of ANG I conversion by ACE inhibitor treatment does not cause significant ANG I accumulation. Our study shows that the rates of ANG I conversion and degradation in different tissues can be measured by using mono-iodinated ^{125}I -ANG I as a tracer. The present results can serve as a basis for future measurements of regional ANG I production rates in the intact animal.

3.2 Introduction

Angiotensin II (ANG II), the most important biologically active product of the renin-angiotensin system, is a powerful hypertensive agent. It acts on specific receptors in vascular smooth muscle, kidney, adrenal, brain and other tissues (7,12,13). The ANG II that reaches these receptors is derived from angiotensin I (ANG I), which is produced both in circulating blood and in the tissues (1,3). To better understand the workings of the tissue renin-angiotensin systems, as opposed to the circulating renin-angiotensin system, it is essential to determine how ANG I is handled by the tissues. Arterially delivered ANG I is rapidly extracted from the circulation by the tissues, due to the hydrolytic attack by peptidases that degrade ANG I into smaller biologically inactive peptides or convert ANG I to ANG II. To date however few studies have been performed in intact animals to address this issue. In one such study ¹²⁵I-labeled ANG I was infused into the renal artery of the dog to measure intrarenal ANG I degradation and ANG I-II conversion (14,19). We administered ¹²⁵I-ANG I, via systemic intravenous infusion, to subjects with essential hypertension who were being treated with the angiotensin converting enzyme (ACE) inhibitor captopril. We found that the extraction of ¹²⁵I-ANG I by the kidney, the leg, the forearm and the liver did not differ from the extraction of unlabeled native ANG I (1).

The aim of the present study was to examine further whether ¹²⁵I-ANG I is a suitable tracer to assess the distribution of endogenous ANG I and to quantify ANG I-II conversion and ANG I metabolic degradation in the intact organism. For this purpose we administered ¹²⁵I-ANG I, alone or in combination with unlabeled ANG I, via a constant infusion into the left cardiac ventricle of the pig and studied the whole body as well as the regional metabolism of the two peptides, both during ACE inhibitor treatment and without such treatment.

3.3 Materials and methods

Chemicals

[Ile⁵]-ANG-(1-10) decapeptide (ANG I), [Ile⁵]-ANG-(1-8) octapeptide (ANG II), and [Ile⁵]-ANG-(2-8) heptapeptide (ANG III) were obtained from Bachem, Bubendorf, Switzerland. [Ile⁵]-ANG-(2-10) nonapeptide (ANG-(2-10)) was from Senn Chemicals, Dielsdorf, Switzerland. [Ile⁵]-ANG-(3-8) hexapeptide (ANG-(3-8)), [Ile⁵]-ANG-(4-8) pentapeptide (ANG-(4-8)), and [Ile⁵]-ANG-(1-7) heptapeptide (ANG-(1-7)) were from Peninsula Laboratories, Belmont, CA, USA. Methanol, ortho-phosphoric acid (both analytical grade) and 1,10-phenantroline were from Merck, Darmstadt, Germany. Bovine serum albumin (BSA) was from Sigma, St. Louis, MO, USA. Water for high performance liquid chromatography (HPLC) was prepared with a Milli-Q system from

Waters, Millford, MA, USA. The renin inhibitor Ro 42,5892 was a kind gift of Dr. P. van Brummelen (Hoffmann-La Roche, Basel, Switzerland).

Preparation of radiolabeled angiotensins

Mono-iodinated ^{125}I -ANG I was prepared with the chloramine-T method and purified as described previously (5). ^{125}I -labeled preparations of ANG II, ANG III, ANG-(3-8), ANG-(4-8), ANG-(2-10), ANG-(1-7) and tyrosine were also made.

Separation of angiotensins by HPLC

Angiotensins and their metabolites were extracted from plasma by reversible adsorption to octadecylsilyl silica (SepPak C18, Waters, Millford, MA, USA) and separated by reversed phase HPLC, according to the method of Nussberger et al (15) with some modifications (1). Separations were performed on a reversed phase Nucleosil C18 steel column of 250 x 4.6 mm and 10 μm particle size. Mobile phase A was 0.085 % ortho-phosphoric acid containing 0.02 % sodium azide. Mobile phase B was methanol. The flow was 1.5 mL/min and the working temperature was 45 °C. SepPak plasma extracts were dissolved in 100 μL of HPLC solvent and injected. Elution was performed as follows: 65 % A/35 % B from 0 to 6 min followed by a linear gradient to 45 % A/55 % B until 12 min. The eluate was collected in 20-sec fractions into polystyrene tubes coated with BSA. The concentrations of ^{125}I -ANG I and its metabolites in the HPLC fractions were measured in the gamma counter. The fractions containing unlabeled ANG I and ANG II were neutralized with 0.5 M sodium hydroxide and vacuum dried at 4 °C. Recovery after HPLC separation alone was 90-95 % for both labeled and unlabeled ANG I and ANG II. The overall recovery of ^{125}I -ANG I and ^{125}I -ANG II added to plasma samples was 85 (7) % and 84 (8) % (mean and SD, n=6) respectively. Similar values were obtained for ANG I and ANG II. Results were not corrected for incomplete recovery.

Assay of angiotensins

ANG I and ANG II concentrations were measured by radioimmunoassay (1) after SepPak extraction and HPLC separation. The ANG I antiserum crossreacted with ANG-(2-10) (100 %) but not (less than 0.1 %) with ANG II, ANG III, ANG-(3-8), ANG-(4-8) and ANG-(1-7). The ANG II antiserum crossreacted with ANG III (55 %), ANG-(3-8) (73 %) and ANG-(4-8) (100 %), but not (less than 0.2 %) with ANG I, ANG-(2-10) or ANG-(1-7). These patterns of crossreactivity show that the antibodies in both antisera were directed against the C-terminal sequences of ANG I and ANG II.

In vivo studies on angiotensin I metabolism

Studies were carried out in twenty female pigs (crossbred Yorkshire x Landrace, Hedelse Varkens Combinatie, Hedel, The Netherlands) with a body weight of 23-31 kg. The pigs were kept on a normal sodium diet. Twelve pigs had been treated with captopril, 25 mg twice daily, for 3 days. The experiments in these pigs were performed 3-4 hours after the last captopril dose.

The pigs were sedated with azaperone (Stresnil, Janssen Pharmaceuticals, Beerse, Belgium), 5 mg/kg i.m., 15 min later followed by metomidate (Hypnodil, Janssen Pharmaceuticals), 3 mg/kg i.v.. Subsequently the pigs were intubated for artificial ventilation with a mixture of oxygen and nitrous oxide (1:2). Respiratory rate and tidal volume were adjusted to keep arterial blood gases within the normal range. In order to maintain an adequate anesthesia, a catheter was placed in the superior caval vein via the external jugular vein for the administration of 160 mg/kg α -chloralose (Merck, Darmstadt, Germany). Fluid losses were corrected with saline, also via this catheter. A peripheral ECG-lead was monitored throughout the experiment. An 8F catheter inserted into the descending aorta, via a femoral artery, was used to measure central aortic pressure (50 AD pressure transducer, Spectramed, Bilthoven, The Netherlands). A 7F catheter was manipulated into the left ventricle, via the carotid artery, under X-ray control, for the infusion of ^{125}I -ANG I and unlabeled ANG I. After the heart had been exposed by a midsternal split, electromagnetic flow probes (Skalar, Delft, The Netherlands) were placed around the ascending aorta and the left anterior descending coronary artery.

Blood for angiotensin measurements was collected from the aorta, pulmonary artery, left and right atrium and great cardiac vein in 8 captopril-treated and 8 untreated pigs. An 8F catheter was placed in the aorta, via a femoral artery. A 7F catheter was introduced into the pulmonary artery under radiographic control, via a femoral vein. The left and the right atrial appendages were catheterized each with an 8F catheter. The vein accompanying the left anterior descending coronary artery (great cardiac vein) was cannulated with a polyethylene catheter. In 4 additional captopril-treated pigs, blood was collected from the aorta and the femoral, renal, jugular vein and ear lobe veins. A 7F catheter was inserted into the right femoral vein and subsequently positioned in the right renal vein under fluoroscopy. An 7F catheter was placed in the left femoral vein. The left jugular vein was cannulated with a 7F catheter. The ear lobe vein was cannulated with a polyethylene catheter.

After a stabilization period of 45 min, following the completion of instrumentation, the pigs were given a constant infusion of ^{125}I -ANG I, 3.6×10^6 (0.5×10^6) cpm/min (mean and SD), either alone (9 captopril-treated pigs, 5 untreated pigs) or combined with unlabeled ANG I, 0.9 (0.3) nmol/min (3

captopril-treated pigs, 3 untreated pigs), into the left cardiac ventricle. During the infusion, which lasted 20 minutes, hematocrit measurements were performed in arterial blood to enable us to convert blood flow into plasma flow. Blood samples (5 mL) to measure ^{125}I -labeled and unlabeled angiotensins were taken from each sampling site at 2 min before and at 9, 14 and 19 min after the start of the infusion. To determine the half lives of ^{125}I -ANG I and unlabeled ANG I blood samples were taken in the first 2-3 min period after the infusion had been stopped.

The blood was rapidly drawn with a plastic syringe containing the following inhibitors (0.25 mL inhibitor solution in 5 mL blood), 0.01 mM Ro 42,5892, 6.25 mM disodium EDTA and 1.25 mM 1,10-phenantroline (final concentrations in blood). Previous measurements of plasma renin activity in porcine plasma had shown that a concentration of 0.01 mM of the renin inhibitor Ro 42,5892 was sufficient to cause complete inhibition of ANG I production (unpublished results). The blood samples were immediately transferred into prechilled polystyrene tubes and centrifuged at 3000 g for 10 min at 4 °C. Plasma was stored at -70 °C and assayed within two weeks.

In vitro studies on angiotensin I metabolism

Since the results of our *in vivo* studies raised the possibility that the rate of ^{125}I -ANG I-II conversion might be different from the rate of ANG I-II conversion, the *in vitro* elimination rate of ^{125}I -ANG I in porcine plasma was compared with that of unlabeled ANG I under standardized conditions, both in the presence and absence of captopril. In the presence of captopril the elimination of ^{125}I -ANG I and ANG I is due to degradation, whereas in the absence of captopril the elimination is due both to ^{125}I -ANG I and ANG I degradation and ^{125}I -ANG I-II and ANG I-II conversion.

Forty mL blood was collected from 4 female pigs, with a body weight of 24-29 kg. The blood was collected into polystyrene tubes containing trisodium citrate (0.2 mL in 10 mL blood, final concentration 0.013 M). The blood samples were immediately centrifuged at 3000 g for 10 min at room temperature. Plasma was stored at -20 °C.

Prior to the experiment, the renin inhibitor Ro 42,5892 was added to plasma (final concentration 0.01 mM), in order to prevent ANG I formation. Captopril (final concentration 0.4 mM) was added to study degradation separately from conversion.

After the plasma containing the inhibitors had been brought to a temperature of 37 °C in a waterbath, the experiment was started by adding at $t=0$ either 10^6 cpm ^{125}I -ANG I or 2 pmol ANG I (both in a volume of 100 μL) to 2 mL plasma. Plasma aliquots of 200 μL were taken at 0.25, 1, 2, 5, 10, 20 and 40 min and immediately mixed with an inhibitor solution containing the following inhibitors (20 μL inhibitor solution in 200 μL plasma) 1.25 mM 1,10-phenantroline and 6.25 mM disodium EDTA (final concentrations in plasma).

The samples were kept on ice and SepPak extraction of angiotensins was performed within one hour. The SepPak extracts were applied to the HPLC column and ^{125}I -ANG I and ANG I were measured as described before.

Captopril, in the concentration we used, caused complete blockade of conversion. Experiments in which ^{125}I -ANG II or ANG II was added to plasma, demonstrated that angiotensin degradation was not altered by captopril.

Calculations

Regional extraction of ANG I. The ANG I extraction ratio, E, is defined as follows:

$$E = 1 - \text{ANG } I_{\text{out}} / \text{ANG } I_{\text{in}}, \quad (1)$$

in which $\text{ANG } I_{\text{in}}$ is the steady state concentration of exogenous ANG I in inflowing (arterial) plasma and $\text{ANG } I_{\text{out}}$ the concentration in outflowing (venous) plasma during the infusion of ANG I into the left cardiac ventricle.

Extraction ratios of ^{125}I -ANG I and unlabeled ANG I were calculated for the following vascular beds: 1) left cardiac cavity (left atrium to aorta), 2) combined systemic vascular beds (aorta to right atrium), 3) myocardium (aorta to great cardiac vein), 4) right cardiac cavity (right atrium to pulmonary artery), 5) lungs (pulmonary artery to left atrium), 6) skeletal muscle (aorta to femoral vein), 7) kidney (aorta to renal vein), 8) head (aorta to jugular vein), and 9) skin (aorta to ear lobe vein).

The clearance of ANG I from the blood during its passage in a given vascular bed is given by the following equation:

$$\text{Plasma clearance} = Q \cdot E = Q \cdot (\text{ANG } I_{\text{in}} - \text{ANG } I_{\text{out}}) / \text{ANG } I_{\text{in}}, \quad (2)$$

in which Q is the plasma flow.

The clearance rate calculated in this way is not the total clearance by a given tissue because only part of the intracardially infused and arterially delivered ANG I reaches the tissue sites where it is metabolized. Therefore the term intrinsic clearance is introduced (18,20,21). Intrinsic clearance is a measure of the inherent ability of the tissues to metabolize ANG I, i.e. the maximal eliminating capacity in the absence of flow limitations. To calculate the intrinsic clearance, it is assumed that the ANG I concentration in the tissue compartment from which ANG I is eliminated equals its concentration in outflowing plasma, so that:

$$\text{Intrinsic clearance} = Q \cdot E / (1 - E) = Q \cdot (\text{ANG } I_{\text{in}} - \text{ANG } I_{\text{out}}) / \text{ANG } I_{\text{out}} \quad (3)$$

This equation is based upon the so-called 'venous equilibrium' model, in which the vascular bed between the arterial and venous sampling sites is considered to be a single, well-stirred compartment (18,21). For most vascular beds, $ANG I_{in}$ and $ANG I_{out}$ are the concentrations of exogenous ANG I in plasma of arterial and venous blood respectively. For the right cardiac cavity, $ANG I_{in}$ is the concentration of exogenous ANG I in the right atrium and $ANG I_{out}$ the concentration in the pulmonary artery. For the left cardiac cavity, $ANG I_{in}$ was calculated as follows:

$$ANG I_{in} = ANG I_{left\ atrium} + \text{Infusion rate/ Cardiac output of plasma.} \quad (4)$$

$ANG I_{out}$ for the left cardiac cavity is the concentration of exogenous ANG I in the aorta.

Regional degradation and conversion of ANG I. The regional ANG I extraction ratio, E , is determined by the metabolic rate constant, k , and the time, t , during which the arterially delivered ANG I is exposed to peptidases during the passage of the blood from the arterial to the venous side of the vascular bed. E can be written as a function of k and t , as follows:

$$E = 1 - e^{-kt}. \quad (5)$$

The metabolism of ANG I comprises both ANG I degradation and ANG I-II conversion, so that:

$$k = k_1 + k_2, \quad (6)$$

in which k_1 is the first order rate constant for ANG I degradation and k_2 the first order rate constant for conversion. Substitution of equation (6) into (5) gives:

$$E = 1 - e^{-(k_1t + k_2t)}, \quad (7)$$

in which t is the time during which ANG I is exposed to the degrading and converting enzymes as the blood flows from the arterial side to the venous side.

From equation (7) it follows that:

$$k_1t + k_2t = -\ln(1-E) = -\ln(ANG I_{out}/ANG I_{in}). \quad (8)$$

Assuming k_1 to be the same for ANG II as for ANG I, k_1t can be calculated as

follows:

$$k_1t = - \ln [(ANG I_{out}+ANG II_{out})/(ANG I_{in}+ANG II_{in})]. \quad (9)$$

Subsequently k_2t can be calculated by subtracting k_1t from $k_1t + k_2t$ as follows:

$$k_2t = \ln[(ANG I_{out}+ANG II_{out})/(ANG I_{in}+ANG II_{in})] - \ln(ANG I_{out}/ANG I_{in}). \quad (10)$$

The fractional degradation of arterially delivered ANG I was defined as follows:

$$\text{Fractional degradation} = E \cdot k_1t/(k_1t + k_2t). \quad (11)$$

The fractional conversion of arterially delivered ANG I was defined as follows:

$$\text{Fractional conversion} = E \cdot k_2t/(k_1t + k_2t). \quad (12)$$

Statistical evaluation. Differences in metabolism between ^{125}I -ANG I and ANG I were evaluated for statistical significance ($p < 0.05$) by using Student's t-test for paired observations. Differences in metabolism of ANG I between captopril-treated and untreated pigs were analysed with Student's t-test for unpaired observations.

3.4 Results

Hemodynamic effects of ^{125}I -ANG I and ANG I infusions

Heart rate and cardiac output of plasma did not change during the infusion experiments (Fig. 1). Coronary plasma flow, as judged from the left anterior descending coronary artery flow, also remained constant. The mean value of cardiac output of plasma was used to calculate ^{125}I -ANG I and ANG I clearance rates in the pulmonary and combined systemic vascular beds respectively. Arterial pressure showed a tendency to fall during the experiment but the change was not statistically significant. Coronary plasma flow tended to be higher with captopril treatment than without treatment, but again the difference did not reach statistical significance.

Identification of ^{125}I -ANG I, ^{125}I -ANG II and other ^{125}I -labeled peptides present in plasma during ^{125}I -ANG I infusion

HPLC separation patterns of ^{125}I -labeled peptides in plasma from captopril-

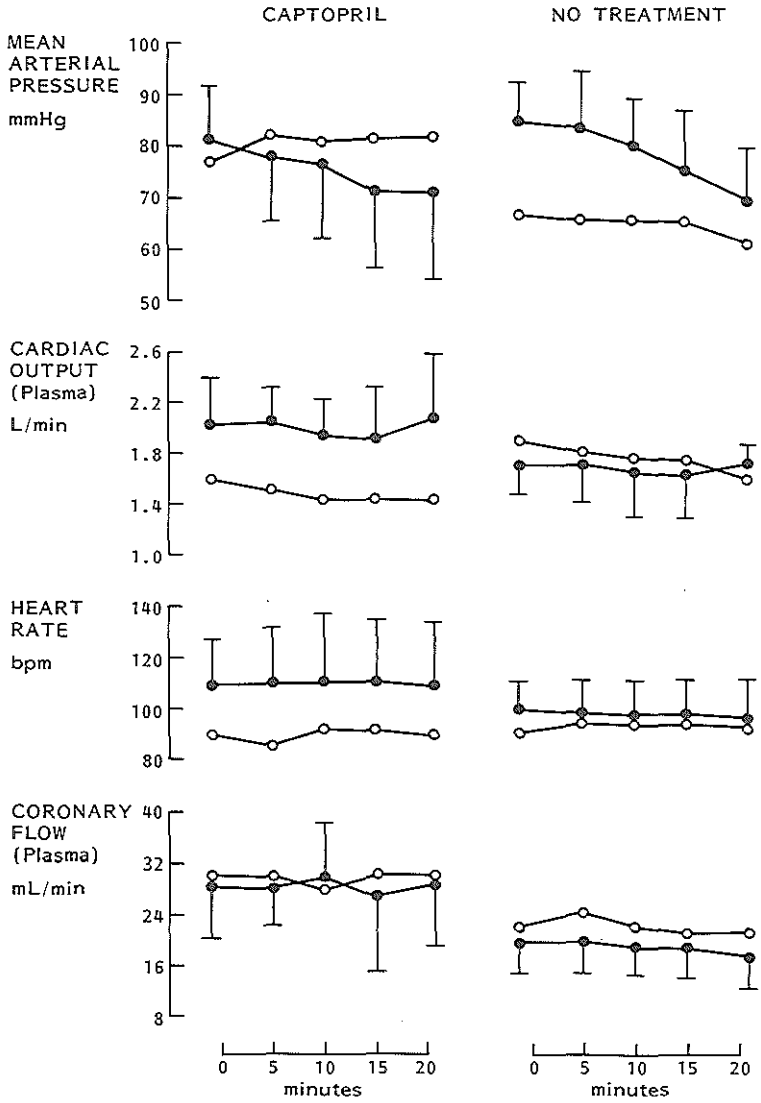


Figure 1. Hemodynamic parameters (mean and SD) in captopril-treated and untreated pigs before, during and after constant infusion into the left cardiac ventricle of either ^{125}I -ANG I alone (closed circles, $n=5$ in both groups) or ^{125}I -ANG I in combination with ANG I (open circles, $n=3$ in both groups). 'Coronary flow' means the flow in the left anterior descending coronary artery.

treated and untreated pigs are shown in Fig. 2. Satisfactory separations were obtained between ^{125}I -ANG I and ^{125}I -ANG II, and between these peptides and their ^{125}I -labeled metabolites. A comparison with the retention times of the various ^{125}I -labeled standards demonstrated that the ^{125}I -ANG I and ^{125}I -ANG II peaks were virtually free of ^{125}I -tyrosine, ^{125}I -ANG-(1-7), ^{125}I -ANG III, ^{125}I -ANG-(3-8), ^{125}I -ANG-(4-8) or ^{125}I -ANG-(2-10). In addition, the ANG I antiserum was able to bind 90-100 % of the radioactivity in the ^{125}I -ANG I peak (and less than 3 % in the ^{125}I -ANG II peak), whereas the ANG II antiserum bound more than 90 % of the radioactivity in the ^{125}I -ANG II peak (and less than 5 % in the ^{125}I -ANG I peak).

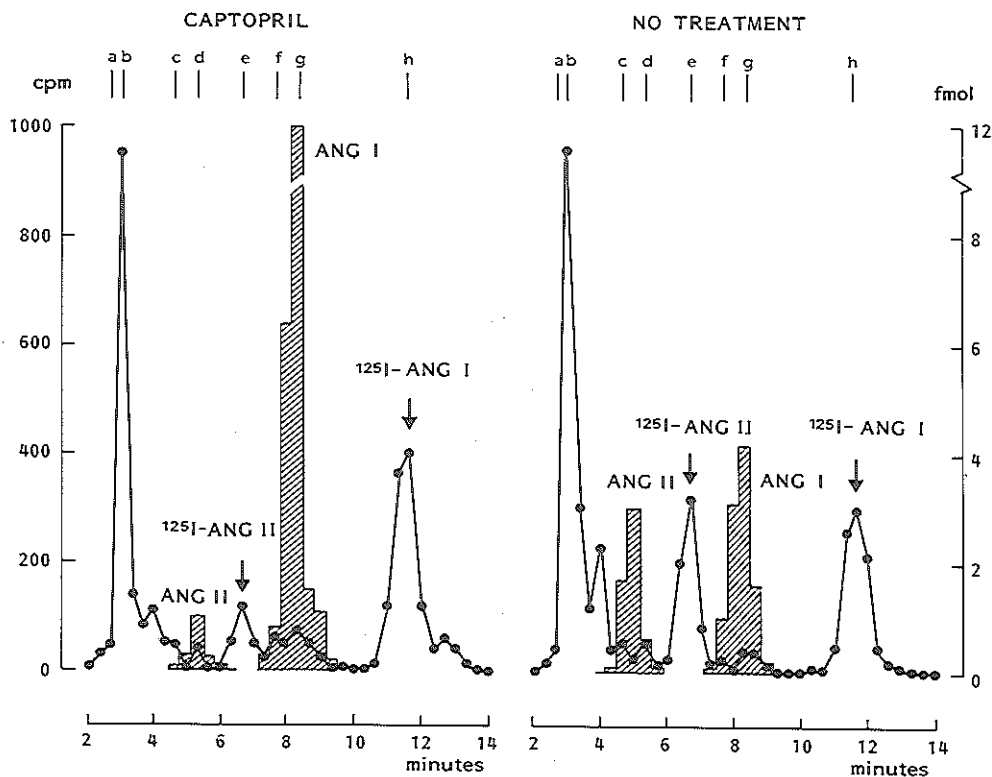


Figure 2. HPLC elution profile of ^{125}I -labeled and endogenous angiotensins in aortic plasma of a captopril-treated (left) and an untreated (right) pig, during constant infusion of ^{125}I -ANG I into the left cardiac ventricle. Radiolabeled peptides (line) were measured by gamma counting, and endogenous angiotensins (bars) were measured by radioimmunoassay. Details on the HPLC elution program are given in the text (see Methods section). a, ^{125}I -ANG-(1-7); b, ^{125}I -tyrosine; c, ^{125}I -ANG III; d, ^{125}I -ANG-(4-8); e, ^{125}I -ANG II; f, ^{125}I -ANG-(2-10); g, ^{125}I -ANG-(3-8); h, ^{125}I -ANG I.

In addition to the ^{125}I -ANG I and ^{125}I -ANG II peaks, separate peaks with retention times corresponding with ^{125}I -tyrosine (and possibly also ^{125}I -ANG-(1-7)) and ^{125}I -ANG-(2-10) could be readily identified in plasma. Peaks corresponding with other metabolites, such as ^{125}I -ANG III and ^{125}I -ANG-(4-8), were very low or absent.

Not only radiolabeled ANG I and ANG II, but also unlabeled ANG I and ANG II were satisfactorily separated from each other and from either tyrosine, ANG-(1-7), ANG III, ANG-(3-8), ANG-(4-8) and ANG-(2-10). The small amounts of radioactivity present in the HPLC fractions containing unlabeled ANG I and ANG II did not interfere with the radioimmunoassays of these peptides.

Plasma levels of ^{125}I -ANG I during and following ^{125}I -ANG I infusion

The arterial and venous plasma levels of ^{125}I -ANG I remained constant from the 9th minute of ^{125}I -ANG I infusion (the first time blood was sampled during the infusion) until the end of the infusion, which demonstrates that a steady state was reached. The steady state plasma levels of ^{125}I -ANG I across the various vascular beds (mean value of the three samples taken at 9, 14 and 19 minutes from each sampling site) were used to calculate the ^{125}I -ANG I extraction ratios of these beds. Tables 1 and 2 summarize the results obtained in 8 captopril-treated and 8 untreated pigs. The high extraction ratios of ^{125}I -ANG I document the rapid metabolism of ^{125}I -ANG I by the tissues. It should be noted that ^{125}I -ANG I was extracted not only in vascular beds where the arterially delivered blood has to pass the capillaries to reach the veins, but also in the right and left cardiac cavities (right atrium to pulmonary artery and left atrium to aorta). The ^{125}I -ANG I extraction ratios of the combined systemic vascular beds (aorta to right atrium) and the myocardium (aorta to great cardiac vein) were not significantly altered by captopril treatment. In contrast, the ^{125}I -ANG I extraction ratio of the lungs (pulmonary artery to left atrium) was greatly reduced by captopril ($p < 0.01$), confirming that ANG I-II conversion is the main pathway by which ANG I is eliminated by the lungs.

In contrast with ^{125}I -ANG I, the total radioactivity levels in plasma kept increasing during the infusion, at all sampling sites, indicating accumulation of one or more radiolabeled metabolites. At the end of the infusion more than 95 % of the total radioactivity of inflowing plasma was recovered in the outflowing plasma across the cardiac cavities and lungs. Across the other vascular beds we studied 90-95 % of arterially delivered radioactivity was recovered in venous plasma. These findings document that the extraction of ^{125}I -ANG I by the tissues we studied is mainly, if not wholly, due to its local metabolism. There was no indication of local accumulation of unmetabolized ^{125}I -ANG I.

Table 1. Steady state plasma levels of ^{125}I -ANG I and ^{125}I -ANG II in captopril-treated and untreated pigs during constant ^{125}I -ANG I infusion into the left cardiac ventricle.

SAMPLING SITE	^{125}I -ANG I cpm/mL		^{125}I -ANG II cpm/mL	
	CAPTOPRIL TREATMENT	NO TREATMENT	CAPTOPRIL TREATMENT	NO TREATMENT
Aorta	1861 (553)	1197 (267)	189 (166)	922 (145)
Right Atrium	769 (169)	534 (129)	128 (114)	710 (134)
Pulmonary Artery	656 (102)	466 (105)	107 (100)	595 (95)
Left Atrium	568 (114)	259 (102)	161 (152)	771 (121)
Great Cardiac Vein	959 (145)	585 (215)	166 (149)	993 (226)

Data are means and SD (n=8). ^{125}I -ANG I infusion rate was $3.08 (0.34) \times 10^6$ cpm/min in the captopril-treated pigs and $3.36 (0.60) \times 10^6$ cpm/min in the untreated pigs.

Table 2. Regional extraction ratios of ^{125}I -ANG I in captopril-treated and untreated pigs during constant ^{125}I -ANG I infusion into the left cardiac ventricle.

VASCULAR BED	CAPTOPRIL TREATMENT	NO TREATMENT
Left Cardiac Cavity	0.22 (0.16)	0.46 (0.13)*
Right Cardiac Cavity	0.14 (0.09)	0.14 (0.09)
Combined Systemic Vascular Beds	0.57 (0.11)	0.55 (0.06)
Lungs	0.14 (0.10)	0.47 (0.12)*
Myocardium	0.45 (0.11)	0.52 (0.13)
Head	0.58 (0.47-0.67)	ND
Skeletal Muscle	0.70 (0.58-0.84)	ND
Kidney	0.94 (0.86-0.98)	ND
Skin	0.52 (0.47-0.56)	ND

Data on the left and right cardiac cavities, the combined systemic vascular beds, lungs and myocardium are means and SD (n=8). Data on head, skeletal muscle, kidney and skin are means and ranges (n=4). ND, not done

* $p < 0.05$, captopril treatment vs. no treatment.

The disappearance of ^{125}I -ANG I from plasma after the ^{125}I -ANG I infusion had been stopped was followed in the aorta in all animals (Fig. 3). In the first 0.5 min the level fell rapidly by more than 50 %. Thereafter it fell more gradually, in a mono-exponential way, with a $t_{1/2}$ of 0.82 (0.12) min with captopril and 0.38 (0.17) min without captopril (means and SD, $n=8$ in both groups, $p < 0.01$).

In 2 animals the disappearance of ^{125}I -ANG I after the ^{125}I -ANG I infusion had been stopped was followed both in the pulmonary artery and in the aorta. At steady state during the infusion, the level of ^{125}I -ANG I in the aorta was higher than in the pulmonary artery but, within 30 sec after the infusion had been stopped, ^{125}I -ANG I in the aorta fell to a level close to the level in the pulmonary artery (Fig. 4). Thereafter ^{125}I -ANG I both in the aorta and in the pulmonary artery fell in a mono-exponential way, and the slopes of the two elimination curves were not different. At the moment the infusion was stopped, the level of ^{125}I -ANG I in the aorta that was derived by extrapolation of the mono-exponential curve, was not different from the level in the left atrium at steady state during the infusion. The pulmonary artery/aorta ^{125}I -ANG I concentration ratio in the elimination phase was not different from the steady state pulmonary artery/left atrium ratio during the infusion. Thus, in contrast with the arteriovenous ^{125}I -ANG I concentration ratio across the combined systemic vascular beds, the arteriovenous ratio across the lungs during the infusion was maintained after the infusion. These results are in agreement with the assumption that in the combined systemic vascular beds ANG I is cleared from a single well-stirred compartment that is larger than the blood compartment ('venous equilibrium' model, vide supra), whereas in the pulmonary vascular bed ANG I is mainly cleared from the blood compartment.

Plasma levels of ^{125}I -ANG I and ANG I during and following combined ^{125}I -ANG I and ANG I infusion

The plasma levels of ANG I at the various sampling sites (aorta, right atrium, pulmonary artery, left atrium, great cardiac vein) prior to the infusion were similar, both in the captopril-treated and untreated animals (Table 3). They were 20.1-23.3 pM in the former group, as compared to 2.5-4.9 pM in the latter group. The ANG II levels prior to the infusion were 2.1-3.3 and 1.9-3.0 pM respectively. These results show that reduced ANG I-II conversion after ACE inhibition is overcome by the elevated ANG I levels.

ANG I and ANG II rose by a factor of 7 or more during the infusion in the captopril-treated group, and by more than 100 in the untreated group. From the steady state plasma levels of ANG I during the infusion, the ANG I extraction ratios across the various vascular beds were calculated and compared with the ^{125}I -ANG I extraction ratios. The results are summarized in Table 4.

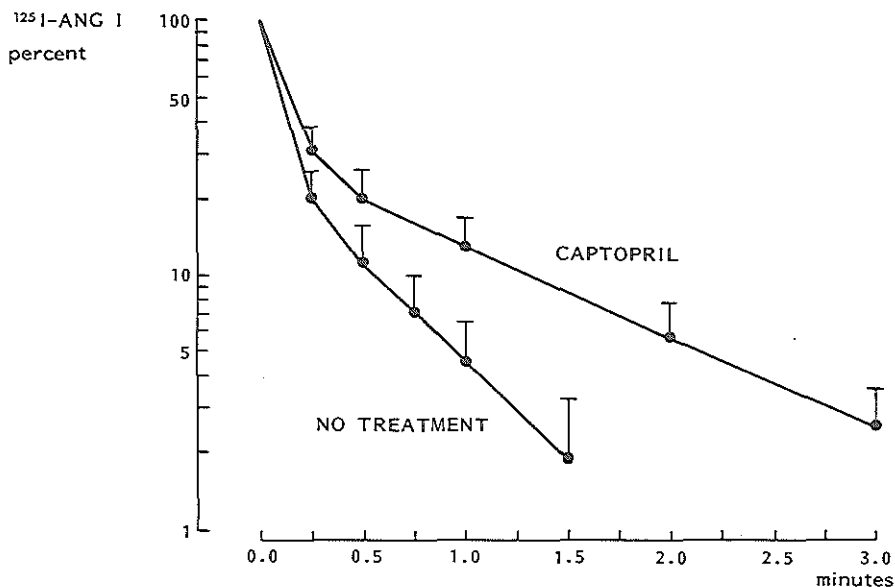


Figure 3. Elimination of ^{125}I -ANG I from aortic plasma in captopril-treated and untreated pigs (mean and SD, $n=8$ in both groups), after discontinuation of a constant infusion of ^{125}I -ANG I into the left cardiac ventricle.

Table 3. Steady state plasma levels of ANG I and ANG II in captopril-treated and untreated pigs before and during constant ANG I infusion into the left cardiac ventricle.

SAMPLING SITE	ANG I pM		ANG II pM	
	CAPTOPRIL TREATMENT	NO TREATMENT	CAPTOPRIL TREATMENT	NO TREATMENT
Aorta	21.4 -> 376	3.6 -> 541	3.3 -> 36.6	2.6 -> 229
Right Atrium	23.3 -> 201	4.9 -> 285	2.3 -> 24.8	2.2 -> 175
Pulmonary Artery	21.9 -> 170	3.6 -> 249	2.1 -> 21.5	1.9 -> 149
Left Atrium	20.1 -> 146	2.5 -> 187	3.0 -> 32.1	3.0 -> 218
Great Cardiac Vein	21.1 -> 175	3.4 -> 330	3.2 -> 29.5	2.3 -> 272

Data (means, $n=3$) before and during infusion are given. ANG I infusion rate was 0.77 nmol/min in the captopril-treated pigs and 1.07 nmol/min in the untreated pigs.

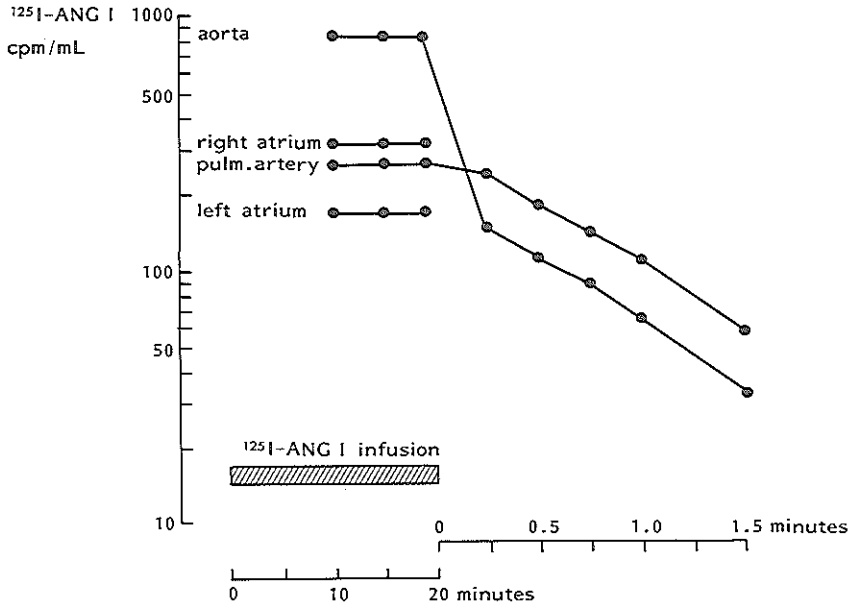


Figure 4. Plasma levels of ^{125}I -ANG I in the aorta, right atrium, pulmonary artery and left atrium at steady state during constant infusion of ^{125}I -ANG I into the left cardiac ventricle and in the elimination phase (aorta and pulmonary artery) after discontinuation of the infusion. Data are from a representative experiment in an untreated pig.

Table 4. Regional extraction ratios of ^{125}I -ANG I and ANG I in captopril-treated and untreated pigs during constant combined infusion of ^{125}I -ANG I and ANG I into the left cardiac ventricle.

VASCULAR BED	CAPTOPRIL TREATMENT		NO TREATMENT	
	^{125}I -ANG I	ANG I	^{125}I -ANG I	ANG I
Left Cardiac Cavity	0.35	0.37	0.45	0.33*
Right Cardiac Cavity	0.20	0.15	0.07	0.12
Combined Systemic Vascular Beds	0.49	0.46	0.55	0.46*
Lungs	0.17	0.14	0.38	0.27*
Myocardium	0.54	0.51	0.44	0.40

Data are means (n=3).

*The extraction ratio for ^{125}I -ANG I was higher than for ANG I in all three experiments

With captopril treatment no difference in extraction by the various vascular beds was observed between ^{125}I -ANG I and ANG I. Without captopril treatment however, the extraction by the lungs, the combined systemic vascular beds and the left cardiac cavity was higher for ^{125}I -ANG I than for ANG I. The elimination half lives of ^{125}I -ANG I and ANG I were not significantly different in the captopril-treated nor in the untreated animals (Fig. 5).

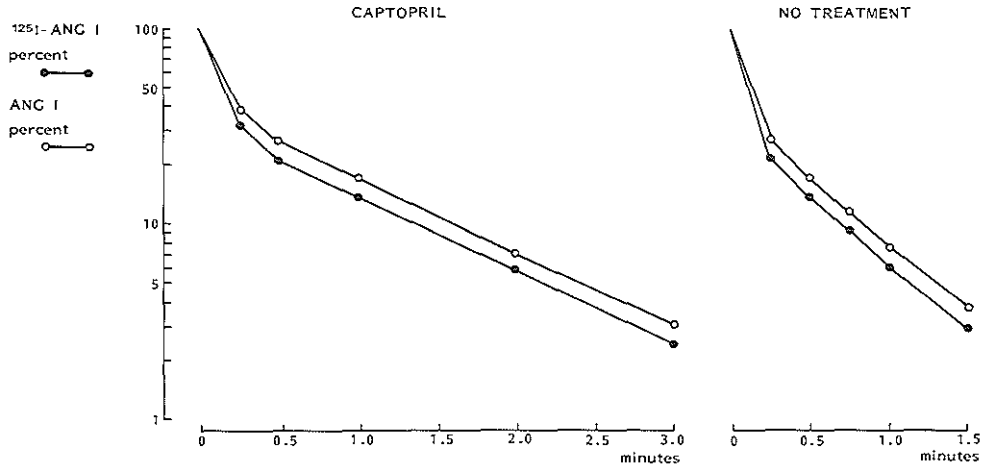


Figure 5. Plasma levels (means, n=3) of ^{125}I -ANG I (closed symbols) and ANG I (open symbols) in the aorta in the elimination phase after discontinuation of a constant combined infusion of ^{125}I -ANG I and ANG I into the left cardiac ventricle.

Regional ^{125}I -ANG I degradation and ^{125}I -ANG I-II conversion as compared with regional ANG I degradation and ANG I-II conversion

The metabolism of ANG I can be divided into two enzymatic processes, ANG I degradation and ANG I-II conversion. The reaction kinetics of these two processes are characterized by the first order rate constants k_1 and k_2 for degradation and conversion, respectively. As can be seen from Table 5, k_1 for ^{125}I -ANG I and k_1 for ANG I were similar. In contrast, k_2 for ^{125}I -ANG I was 1.3-2.7 times k_2 for ANG I, which indicates that the ^{125}I -ANG I-II conversion rate was approximately two times the ANG I-II conversion rate. These findings explain why in the untreated pigs the ^{125}I -ANG I extraction ratios were higher than the ANG I extraction ratios.

Table 5. Regional ^{125}I -ANG I and ANG I degradation and conversion constants, k_{1t} and k_{2t} respectively, in captopril-treated and untreated pigs during constant combined infusion of ^{125}I -ANG I and ANG I into the left cardiac ventricle.

VASCULAR BED	CAPTOPRIL TREATMENT				NO TREATMENT			
	^{125}I -ANG I		ANG I		^{125}I -ANG I		ANG I	
	k_{1t}	k_{2t}	k_{1t}	k_{2t}	k_{1t}	k_{2t}	k_{1t}	k_{2t}
Left Cardiac Cavity	0.36	0.09	0.47	0.05*	0.37	0.26	0.29	0.12*
Right Cardiac Cavity	0.23	0.01	0.17	0.00	0.11	0.00	0.15	0.01
Combined Systemic Vascular Beds	0.66	0.05	0.62	0.02*	0.53	0.29	0.51	0.12*
Lungs	0.02	0.17	0.07	0.08*	0.06	0.45	0.01	0.35*
Myocardium	0.72	0.11	0.74	0.04*	0.24	0.36	0.27	0.25*

Data are means (n=3)

* k_{2t} for ANG I was smaller than for ^{125}I -ANG I in all three experiments.

Table 6. Regional ANG I degradation and conversion constants, k_{1t} and k_{2t} respectively, calculated from the steady state plasma levels of ^{125}I -ANG I in captopril-treated and untreated pigs during constant infusion of ^{125}I -ANG I into the left cardiac ventricle.

VASCULAR BED	k_{1t}	k_{1t}	k_{2t}	k_{2t}
	CAPTOPRIL TREATMENT	NO TREATMENT	CAPTOPRIL TREATMENT	NO TREATMENT
Left Cardiac Cavity	0.23 (0.17)	0.35 (0.15)	0.02 (0.02)	0.15 (0.06)*
Right Cardiac Cavity	0.15 (0.11)	0.15 (0.12)	0.00 (0.00)	0.00 (0.01)
Combined Systemic Vascular Beds	0.84 (0.23)	0.54 (0.11)*	0.02 (0.01)	0.14 (0.03)*
Lungs	0.07 (0.10)	0.04 (0.04)	0.05 (0.04)	0.30 (0.10)*
Myocardium	0.59 (0.26)	0.32 (0.21)*	0.03 (0.02)	0.23 (0.05)*
Head	0.77 (0.56-1.07)		0.05 (0.02-0.10)	
Skeletal Muscle	1.04 (0.68-1.65)		0.12 (0.08-0.19)	
Kidney	2.58 (1.51-3.49)		0.24 (0.20-0.26)	
Skin	0.69 (0.64-0.74)		0.02 (0.00-0.04)	

Data on the left and right cardiac cavities, the combined systemic vascular beds, lungs and myocardium are means and SD (n=8). Data on head, skeletal muscle, kidney and skin are means and ranges (n=4).

* $p < 0.05$, captopril treatment vs. no treatment.

In vitro experiments, in which ^{125}I -ANG I and ANG I were added to plasma and the elimination of the two peptides was followed, also showed a higher conversion rate for ^{125}I -ANG I than for ANG I. Elimination followed first order kinetics, with a rate constant k of 0.68 (0.04) min^{-1} for ^{125}I -ANG I and 0.41 (0.02) min^{-1} for ANG I (means and SD, $n=5$). After complete blockade of the conversion by the addition of captopril, k was reduced to 0.17 (0.02) min^{-1} for ^{125}I -ANG I and 0.14 (0.01) min^{-1} for ANG I ($n=5$). The slow elimination of ^{125}I -ANG I and ANG I in plasma containing captopril as compared to plasma without captopril shows that the in vitro elimination of ^{125}I -ANG I and ANG I from plasma in the absence of captopril was mainly caused by ^{125}I -ANG I-II conversion and ANG I-II conversion respectively. In the presence of captopril k equals the first order rate constant for degradation, k_1 . By subtracting k_1 from the value of k measured in the absence of captopril, one obtains the first order rate constant for conversion, k_2 . For ^{125}I -ANG I, k_2 was 0.51 (0.03) min^{-1} , and for ANG I it was 0.27 (0.02) min^{-1} . It appeared that k_2 was 1.9 (0.3) times higher for ^{125}I -ANG I than for ANG I. These in vitro findings are in good agreement with the in vivo results described above.

Table 6 gives the calculated regional degradation and conversion constants, k_{1t} and k_{2t} respectively, for ANG I. Calculations were based on equations (8), (9) and (10), which give the k_{1t} and k_{2t} values for ^{125}I -ANG I. Because of the results of the above experiments in which the conversion rates of ^{125}I -ANG I and ANG I were compared, k_{2t} for ^{125}I -ANG I was multiplied by 0.5 to give k_{2t} for ANG I. In the combined systemic vascular beds but not in the lungs and the cardiac cavities, blockade of ANG I-II conversion by captopril treatment was associated with a rise in the ANG I degradation constant.

From the data presented in Table 6, the fractional degradation and conversion of arterially delivered ANG I in the various vascular beds can be calculated by using equations (11) and (12). The results are shown in Fig. 6. In the untreated animals, 25 (7) percent of the arterially delivered ANG I was converted to ANG II during a single passage of blood in the lungs and 10 (2) percent in the combined systemic vascular beds (means and SD, $n=8$). Fractional ANG I degradation in these animals was only 4 (3) percent in the lungs as compared to 39 (6) percent in the combined systemic vascular beds. In the captopril-treated animals fractional ANG I-II conversion was reduced to very low levels, 4 (4) percent in the lungs and 1 (1) percent in the combined systemic vascular beds. Fractional degradation during captopril treatment remained low in the lungs, 6 (8) percent, but rose to 55 (11) in the combined systemic vascular beds. Thus the captopril-induced reduction in ANG I-II conversion in the combined systemic vascular beds was associated with a compensatory rise in ANG I degradation.

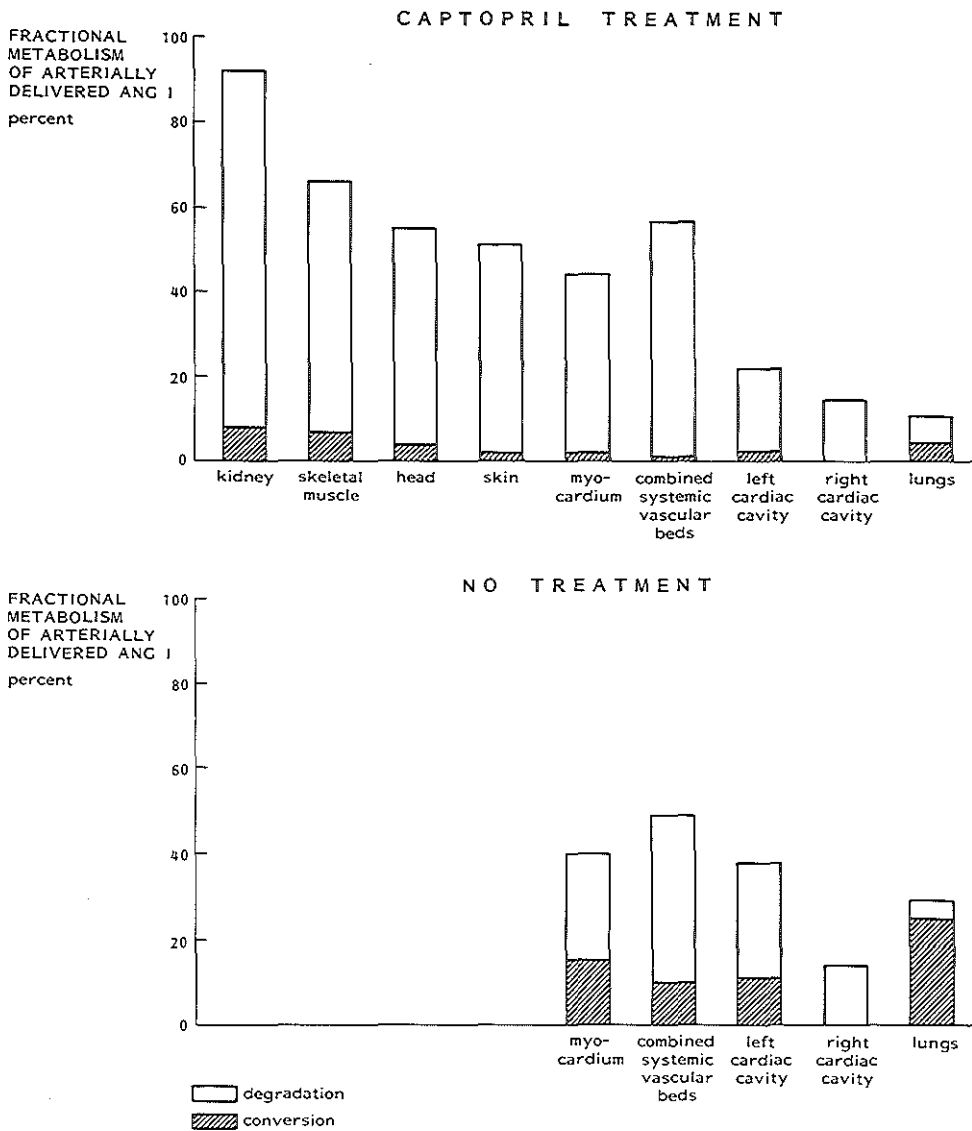


Figure 6. Fractional degradation (open bars) and fractional conversion (hatched bars) of arterially delivered ANG I in different vascular beds. Data (means) on myocardium, combined systemic vascular beds, lungs, and right and left cardiac cavity were obtained in 8 captopril-treated and 8 untreated pigs. Data on other vascular beds were obtained in 4 captopril-treated pigs.

ANG I clearance rates and distribution volume

Clearance calculations included a correction for the higher conversion rate of ^{125}I -ANG I as compared to ANG I (see above). Pulmonary clearance of ANG I was 0.15 (0.16) L/min in the captopril-treated animals and 0.49 (0.16) L/min in the untreated animals ($n=8$ in both groups, $p < 0.01$). Intrinsic clearance of ANG I in the combined systemic vascular beds was 2.51 (0.94) L/min with captopril and 1.70 (0.61) L/min without captopril ($n=8$ in both groups, difference not significant). Thus, with captopril treatment, the contribution of the lungs to the whole body clearance of ANG I appears to be negligible. Under these circumstances the whole body clearance of ANG I can be calculated, according to the single compartment model, as $(\text{Infusion rate of } ^{125}\text{I-ANG I})/({}^{125}\text{I-ANG I}_{\text{left atrium}})$, in which ${}^{125}\text{I-ANG I}_{\text{left atrium}}$ is the steady state concentration of $^{125}\text{I-ANG I}$ in the left atrium during the infusion of $^{125}\text{I-ANG I}$ into the left cardiac ventricle. Whole body clearance was 5.60 (1.07) L/min ($n=8$). Distribution volume, calculated as $(\text{whole body clearance} \cdot t_{1/2})/\ln 2$, was 6.47 (0.93) L. This value represents 18-32 (mean 24) % of total body weight. These results show that ANG I is not limited to the intravascular compartment, since the plasma volume is only 4-5 % of total body weight (6).

3.5 Discussion

Earlier reports have raised some doubt as to whether ^{125}I -labeled angiotensins can be used as a tracer for in vivo metabolic studies (9,11). The aim of the present work was to address this point specifically with regard to ANG I, because this peptide is the direct product of renin's enzymatic action and because little quantitative data are available on ANG I metabolism by the tissues in vivo. Such data are relevant particularly in the light of recent evidence of renin's action at tissue sites as opposed to renin in circulating plasma (1,3).

We used highly purified mono-iodinated ^{125}I -ANG I in our experiments. It was infused directly into the left cardiac ventricle to obtain immediate mixing of the tracer with the arterial blood delivered to the tissues. We performed our experiments in pigs, because in these animals it is possible to collect enough blood from multiple sampling sites and to collect these samples quickly enough to prevent metabolism ex vivo. The arterial and venous plasma levels of ^{125}I -ANG I at steady state were used to quantify the metabolism of endogenous ANG I by the various vascular beds. Total radioactivity, which includes ^{125}I -ANG I as well as its radiolabeled metabolites, was also measured in arterial and venous plasma. Our results demonstrated that the low ^{125}I -ANG I vein/artery ratios as measured across the various vascular beds are indeed due to rapid metabolism and not to local accumulation of ^{125}I -ANG I.

Our results in pigs in which ANG I was infused simultaneously with ^{125}I -ANG I demonstrate that the metabolism of ^{125}I -ANG I was different from that of ANG I. It appeared that the extraction ratios across the pulmonary and the combined extrapulmonary vascular beds were higher for ^{125}I -ANG I than for ANG I, a phenomenon that was observed in untreated pigs, but not in pigs treated with the ACE inhibitor captopril. The metabolism of ANG I comprises ANG I-II conversion and ANG I degradation. By the latter process peptides other than ANG II are formed. The above results indicate that the rate of ^{125}I -ANG I-II conversion was higher than the rate of ANG I-II conversion but that the rate of degradation was the same for ^{125}I -ANG I and ANG I. This was further confirmed by expressing the extraction ratios, E , as a function of k_1t and k_2t , k_1 and k_2 being the first order rate constants for degradation and conversion respectively, and t the time ^{125}I -ANG I or ANG I is exposed to the degrading and converting enzymes. By doing so it was found that k_2t for ^{125}I -ANG I-II conversion in the pulmonary and the combined extrapulmonary vascular beds was indeed higher than k_2t for ANG I-II conversion, and that k_1t for ^{125}I -ANG I degradation was not different from k_1t for ANG I degradation. As expected, it was conversion and not degradation that was reduced by captopril treatment. The first order rate constant for ^{125}I -ANG I-II conversion in vivo appeared to be approximately 2 times that of ANG I-II conversion. In vitro experiments with plasma demonstrated that the V_{max}/K_m ratio of angiotensin converting enzyme (ACE) with ^{125}I -ANG I as the substrate was also 2 times higher than with ANG I as the substrate. This is probably the explanation for our findings in vivo. To calculate ANG I extraction by the vascular beds, the measurements of ^{125}I -ANG I extraction were corrected for the difference between ^{125}I -ANG I-II and ANG I-II conversion rates. This was done by multiplying the first order rate constant for ^{125}I -ANG I-II conversion by a factor of 0.5. In these calculations it was assumed that k_1t for ^{125}I -ANG I degradation was not different from k_1t for ^{125}I -ANG II degradation. Preliminary experiments in which we infused ^{125}I -ANG II instead of ^{125}I -ANG I, showed indeed little difference in degradation rate between ^{125}I -ANG I and ^{125}I -ANG II, both across the pulmonary and the combined systemic vascular beds (unpublished results). In our calculations it was further assumed that k_1t for ANG I degradation was not different from k_1t for ANG II degradation.

The intrinsic clearance of ANG I in the combined systemic vascular beds was calculated to be approximately 1-2 times the cardiac output of plasma. This is much too high to be explained by metabolism in the circulating blood and elimination via glomerular filtration. The high clearance rate is caused by rapid metabolism in the tissues. It is interesting to note that, in the heart, the rapid metabolism of plasma ANG I does not seem to depend on the passage of blood through the capillaries. Probably the endocardium, like the vascular

endothelium, is a site of ANG I metabolism. The presence of ACE on the surface of the cardiac valves has indeed been demonstrated (8,22).

In the combined systemic vascular beds about 20 percent of the metabolism of arterially delivered ANG I was due to ANG I-II conversion, whereas in the pulmonary vascular bed about 90 percent of the metabolism was due to ANG I-II conversion. It is well known that the lungs are not the only site of ANG I-II conversion (4,10,16). Our study provides quantitative information on the extrapulmonary ANG I-II conversion and shows in fact that about half of the conversion of circulating ANG I occurs outside the lungs. There was not only considerable conversion in the combined systemic vascular beds, but also in the left cardiac cavity.

After ACE inhibitor treatment there was little ANG II generation from infused ANG I. The ACE inhibitor-induced decrease in k_2t for ANG I-II conversion was associated with an increase in k_1t for ANG I degradation in the combined systemic vascular beds, but not in the cardiac cavities and the lungs. The mechanism of this increase in k_1t is unknown. Our study indicates that systemically delivered ANG I is rapidly distributed over a compartment with a size corresponding to 24 % of total body weight, which is also the size of the extracellular fluid volume. There may be some degree of compartmentalization of the processes of conversion and degradation in the sense that it is mainly the blood-borne ANG I that is subject to conversion, whereas interstitial fluid ANG I is mainly subject to degradation. Our calculations of k_1t and k_2t do not account for this, but it is conceivable that such compartmentalization may explain why, in contrast with the combined systemic vascular beds, in the cardiac cavities and the lungs no rise in k_1t was observed after ACE inhibitor treatment. In the cardiac cavities and the lungs the ANG I delivered by the inflowing blood is primarily exposed to the endocardium and the endothelial surface of the lung vessels. The lung is known to contain relatively little interstitial fluid as compared to other tissues (17), so that in the lung the arterially delivered ANG I is mainly restricted to the blood compartment. Our results are supported by experiments, in which isolated rabbit aorta was incubated with ANG I. Also in these experiments inhibition of ACE was associated with increased degradation (2). Anyhow, our results show that decreased conversion during captopril treatment was associated with increased degradation. Thus, there was little or no ANG I accumulation during ACE inhibitor treatment. Renin stimulation is therefore by far the most important, if not the only, cause for the rise in plasma ANG I that is usually observed after ACE inhibitor treatment.

We conclude that mono-iodinated ^{125}I -ANG I is a suitable tracer for the type of metabolic studies we did, with the proviso that one has to account for the fact that ^{125}I -ANG I is converted by ACE at a higher rate than ANG I. Our study indicates that ANG I infused into the circulation is rapidly distributed over a compartment corresponding in size with the extracellular fluid volume.

The large arteriovenous concentration gradients of ^{125}I -ANG I that were observed across the various vascular beds during ^{125}I -ANG I infusion are caused by rapid metabolism in the tissues. ANG II formation depends on ACE, but ANG I conversion is not a critical intermediary step in ANG I metabolism. The present results may be used as a basis for further studies aimed at quantifying in the intact animal ANG I production at tissue sites versus ANG I production in the circulation.

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4. PRODUCTION OF ANGIOTENSINS I AND II AT TISSUE SITES IN THE INTACT PIG

4.1 Summary

Angiotensins (ANG) are known to be generated in circulating plasma and at tissue sites. It is not known to what extent generation of ANG I outside the blood stream and conversion of this locally generated ANG I to ANG II contribute to the circulating levels of ANG I and ANG II. To answer these questions, the plasma levels ANG I, ANG II, ^{125}I -ANG I, and ^{125}I -ANG II were measured in arterial and venous blood across a number of vascular beds, during constant infusion of mono-iodinated ^{125}I -ANG I into the left cardiac ventricle of anesthetized pigs with unstimulated and stimulated renin production. From the vein/artery ratios of ^{125}I -ANG I and II and the arteriovenous gradients of ANG I and II it could be calculated what fraction of ANG I and II in regional venous plasma was arterially delivered and what de novo produced, and what fraction of venous ANG II was derived from regional conversion of arterially delivered ANG I. By parallel measurements of plasma renin activity (PRA) and by taking into account the regional blood transit time, it was possible to estimate the contribution of circulating PRA to the de novo production of ANG I and II. De novo ANG I production in the combined systemic vascular beds was increased during treatment with the angiotensin converting enzyme (ACE) inhibitor captopril (n=12) or the diuretic furosemide (n=4), as compared with untreated animals (n=8). ANG II production in the combined systemic vascular beds was greater than in the lungs, demonstrating the importance of extrapulmonary ANG I-II conversion. In the lungs virtually no ANG I but 31 % of ANG II in venous plasma was derived from de novo production, and de novo produced ANG II could fully be accounted for by conversion of circulating ANG I. In myocardium, head, skin, skeletal muscle and kidney, respectively 40, 58, 55, 67, and 94 % of venous ANG I, and 32, 49, 40, 59 and 85 % of venous ANG II was derived from de novo production. In these vascular beds a substantial fraction of de novo produced ANG I could not be accounted for by circulating PRA (> 25 % in myocardium, > 80 % in the other regions, if blood transit time = 10 sec). In addition, de novo produced venous ANG II was too high to be attributed only to conversion of circulating ANG I. These results indicate that indeed production of ANG I at tissue sites contributes to its circulating level, and that some circulating ANG II may not be derived from circulating ANG I.

4.2 Introduction

The classical view of renin and angiotensins as circulating hormones has been challenged by a number of investigators in recent years. It is now recognized that many tissues contain a complete renin-angiotensin system (8), and there is increasing evidence that part of the angiotensin I (ANG I) in circulating plasma does not originate from the catalytic action of circulating renin on circulating angiotensinogen (1,3,10). Some plasma ANG I is probably produced by renin that is present in vascular tissue (19). Vascular renin may be plasma-derived (17,22) or is perhaps synthesized *in situ* (8,14). It is not known how much ANG I is produced outside the circulating blood and how much of this extracirculatory ANG I enters the circulation.

Angiotensin II (ANG II), the most important biologically active product of the renin-angiotensin system in circulating plasma, is commonly thought to be derived from the conversion of circulating ANG I by membrane-bound angiotensin converting enzyme (ACE) of the vascular endothelium. ACE however, is not only present in the vascular endothelium but also elsewhere in vascular tissue (2,9,18) and it is therefore conceivable that part of circulating ANG II is derived from the conversion of ANG I that is produced at tissue sites outside the circulating blood.

Both ANG I and ANG II are rapidly eliminated by the tissues, due to degradation by peptidases into peptides with little or no biological activity. ANG I is also eliminated by its conversion to ANG II. In a previous study we measured degradation and conversion in different vascular beds in intact pigs (5). This was done by administering ^{125}I -ANG I via constant infusion into the left cardiac ventricle and by measuring the steady state plasma levels of ^{125}I -ANG I and ^{125}I -ANG II in the inflowing (arterial) and outflowing (venous) blood of different vascular beds. Here we report on the results of simultaneous measurements of the plasma levels of endogenous ANG I and ANG II and radiolabeled ANG I and ANG II, during the constant infusion of ^{125}I -ANG I. From these results combined with data presented in the previous paper we were able to estimate what proportion of ANG I is produced in circulating plasma and what at tissue sites. It was also possible to estimate what proportion of ANG II is produced by the conversion of arterially delivered ANG I and what not.

4.3 Materials and methods

Assay of angiotensins

The materials that were used and the methods for measuring the plasma levels of labeled and unlabeled ANG I and ANG II are described in the previous paper (5).

Measurement of plasma renin activity

Plasma renin activity (PRA) was measured by incubating plasma for 10, 20, 30 and 60 min at pH 7.4 and 37 °C in the presence of a mixture of inhibitors to block ANG I degradation, ANG I-II conversion and prorenin-renin conversion and to prevent bacterial growth (6). Preliminary experiments showed that ANG I generation was linear in the first 30 min of incubation. Thereafter ANG I generation tended to decrease slightly. Only the first linear part of ANG I generation was used for measuring PRA.

Infusion Protocol

Twentyfour female pigs (crossbred Yorkshire x Landrace, Hedelse Varkens Combinatie, Hedel, The Netherlands) with a body weight of 23-31 kg were included in the study. The pigs were kept on a normal sodium diet. In 12 pigs the renin production was stimulated by treatment with captopril (25 mg twice daily, for 3 days), and in 4 pigs by treatment with furosemide (40 mg twice daily, for 3 days). The experiments in the captopril-treated and furosemide-treated pigs were performed 3-4 hours after dosing.

The pigs were anesthetized, and arterial and venous catheters were installed as described before (5). Blood pressure, heart rate, cardiac output and left anterior descending coronary artery blood flow were monitored continuously during the infusion. Hematocrit measurements were performed in arterial blood to enable us to convert blood flow measurements into plasma flow values. Sampling sites were: aorta, pulmonary artery, left and right atrium, great cardiac vein, and renal, femoral, jugular and ear lobe veins.

After a stabilization period of 45 min following the completion of instrumentation, the pigs were given a constant infusion of ^{125}I -ANG I, 3.6×10^6 (0.9×10^6) cpm/min (mean and SD), into the left cardiac ventricle during 20 min. The infusion did not influence blood pressure, heart rate, cardiac output or coronary blood flow.

Blood samples for measuring the plasma levels of ^{125}I -labeled and unlabeled angiotensins were taken from the various sampling sites at 2 min before and at steady state during the infusion of ^{125}I -ANG I as described before (5).

Blood samples for measuring PRA were also taken during the infusion. The blood was collected into polystyrene tubes containing disodium EDTA (0.1 mL EDTA solution in 3 mL blood, final concentration in blood 10 mM). The samples were centrifuged at 3000 g for 10 min at room temperature. Plasma was stored at -20 °C.

Calculations

Regional production of angiotensin I. Regional ANG I production was

calculated for the following vascular beds: 1) lung (pulmonary artery to left atrium), 2) combined systemic vascular beds (aorta to right atrium), 3) myocardium (aorta to great cardiac vein), 4) head (aorta to jugular vein), 5) skin (aorta to ear lobe vein), 6) skeletal muscle (aorta to femoral vein) and 7) kidney (aorta to renal vein).

Regional ANG I production will be referred to in two ways: 1) total regional ANG I production, and 2) regional ANG I production by circulating PRA. Total regional ANG I production in the systemic vascular beds was calculated as the product of the regional intrinsic clearance and the plasma level of ANG I in outflowing (venous) blood:

$$\text{Total regional ANG I production} = Q \cdot [E_I / (1 - E_I)] \cdot \text{ANG I}_{\text{out}} \quad (1)$$

in which $\text{ANG I}_{\text{out}}$ is the concentration of endogenous ANG I in outflowing (venous) plasma. Q is the regional plasma flow. For the combined systemic vascular beds Q equals the cardiac output of plasma. E_I is the extraction ratio of ANG I, which according to our previous paper (5) is given by the following formula:

$$E_I = 1 - e^{-(k_1 t + 0.5 \cdot k_2 t)}, \quad (2)$$

in which k_1 and k_2 are the first order rate constants of ^{125}I -ANG I degradation (to peptides other than ^{125}I -ANG II) and ^{125}I -ANG I-II conversion respectively, and t is the time during which ^{125}I -ANG I is exposed to the degrading and converting enzymes as the blood flows from the arterial to the venous side. Equation (2) accounts for the difference in conversion rate between labeled and unlabeled ANG I (5).

Equation (1) is based on the 'venous equilibrium' model. In this model the vascular bed between the arterial and venous sampling sites is considered to be a single, well-stirred compartment with an ANG I concentration equal to that in outflowing (venous) plasma (25).

As shown previously (5), the 'venous equilibrium' model can be applied to the systemic vascular beds but not to the pulmonary vascular bed. For the lungs it can be assumed that the clearance of ANG I equals its clearance from plasma, so that:

$$\text{Total pulmonary ANG I production} = Q \cdot [\text{ANG I}_{\text{out}} - (1 - E_I) \cdot \text{ANG I}_{\text{in}}], \quad (3)$$

in which $\text{ANG I}_{\text{out}}$ and ANG I_{in} are the plasma concentrations of endogenous ANG I in the left atrium and the pulmonary artery respectively. Q equals the cardiac output of plasma.

The regional ANG I production by PRA was calculated as follows:

$$\text{Regional ANG I production by PRA} = Q \cdot \text{PRA} \cdot \text{transit time}, \quad (4)$$

in which transit time is the transit time of plasma as it flows from the arterial to the venous side of the vascular bed.

Regional venous ANG I originates in part from arterially delivered ANG I and is in part derived from *de novo* production. Regional venous ANG I from arterial delivery is given by the following equation:

$$\text{ANG I}_{\text{out}} \text{ from ANG I}_{\text{in}} = (1 - E_I) \cdot \text{ANG I}_{\text{in}}. \quad (5)$$

Regional production of angiotensin II. Regional production of ANG II was calculated for the same vascular beds as for ANG I. In analogy with regional ANG I production, the regional production of ANG II will be referred to as 1) total regional ANG II production, and 2) regional ANG II production by conversion of arterially delivered ANG I. Total regional ANG II production in the systemic vascular beds was calculated as the product of the regional intrinsic clearance of ANG II and the plasma level of ANG II in outflowing (venous) blood:

$$\text{Total regional ANG II production} = Q \cdot [E_{II}/(1-E_{II})] \cdot \text{ANG II}_{\text{out}}, \quad (6)$$

in which $\text{ANG II}_{\text{out}}$ is the concentration of endogenous ANG II in outflowing (venous) plasma. E_{II} is the extraction ratio of ANG II, which is given by the following formula:

$$E_{II} = 1 - e^{-k_1 t}. \quad (7)$$

This equation is analogous with equation (2) for E_I , the extraction ratio of ANG I. The difference between the two equations is related to the fact that ANG I is subject to both conversion and degradation, whereas ANG II is only subject to degradation. Equation (7) is based upon the assumption that the first order rate constants for $^{125}\text{I-ANG II}$ degradation and $^{125}\text{I-ANG I}$ degradation are equal. It further accounts for the fact that the degradation rates for labeled and unlabeled ANG I are not different (5).

In analogy with pulmonary ANG I production, the total pulmonary ANG II production was calculated as follows:

$$\text{Total pulmonary ANG II production} = Q \cdot [\text{ANG II}_{\text{out}} - (1-E_{II}) \cdot \text{ANG II}_{\text{in}}]. \quad (8)$$

The regional ANG II production from arterially delivered ANG I was calculated as the product of regional plasma flow, the plasma level of ANG I in inflowing (arterial) blood, and the fractional conversion of arterially delivered ANG I. Fractional conversion of arterially delivered ANG I is given, according to our previous paper (5), by the following equation:

$$\text{Fractional conversion} = E_I \cdot [0.5 \cdot *k_{2t}/(*k_{1t} + 0.5 \cdot *k_{2t})], \quad (9)$$

so that:

Regional ANG II production from arterially delivered ANG I =

$$Q \cdot \text{ANG I}_{in} \cdot E_I \cdot [0.5 \cdot *k_{2t}/(*k_{1t} + 0.5 \cdot *k_{2t})]. \quad (10)$$

Regional venous ANG II originates in part from arterially delivered ANG II and is in part derived from regional de novo production. Regional conversion of arterially delivered ANG I is, at least partly, responsible for this de novo production. Regional venous ANG II from arterial delivery is given by the following equation:

$$\text{ANG II}_{out} \text{ from ANG II}_{in} = (1 - E_{II}) \cdot \text{ANG II}_{in}. \quad (11)$$

If, as mentioned above, it is assumed that the first order rate constants for ¹²⁵I-ANG II degradation and ¹²⁵I-ANG I degradation are equal, and that the degradation rates for labeled and unlabeled ANG I are also equal (5), the regional venous ANG II produced by regional conversion of arterially delivered ANG I is given by the following equation:

$$\text{ANG II}_{out} \text{ from ANG I}_{in} = (E_I - E_{II}) \cdot \text{ANG I}_{in}. \quad (12)$$

Statistical evaluation. Differences in ANG I and ANG II concentration between the various sampling sites were evaluated for statistical significance ($p < 0.05$) by using Scheffé's test for multiple comparison. Differences between total regional ANG I production and regional ANG I production by PRA, and differences between total regional ANG II production and regional ANG II production from arterially delivered ANG I were evaluated for statistical significance ($p < 0.05$) by using Student's t-test for paired observations. Differences between the various groups of pigs were evaluated for statistical significance ($p < 0.05$) by using either Student's t-test or Mann-Whitney's U-test for unpaired observations.

4.4 Results

Plasma levels of angiotensin I, angiotensin II and PRA

The plasma levels of ANG I, ANG II and PRA were stable over the experimental period the blood samples were collected. As expected, the levels of PRA and ANG I were increased in the captopril-treated and furosemide-treated pigs as compared to the untreated animals (Table 1). The levels of ANG I and PRA were linearly correlated, and the regression line in the captopril-treated group was not significantly different from that in the other two groups (Fig. 1). The levels of ANG I and ANG II were also correlated, but for a given level of ANG I, ANG II was lower in the captopril-treated group than in the other two groups (Fig. 2).

Plasma ANG I in the untreated and the furosemide-treated pigs was higher in the right atrium than in the aorta, and it was lower in the left atrium than in the pulmonary artery (Table 1). Thus, there was net ANG I delivery to the systemic circulation by the combined systemic vascular beds, and net ANG I extraction from the systemic circulation by the lungs. These arteriovenous differences were not seen in the captopril-treated pigs.

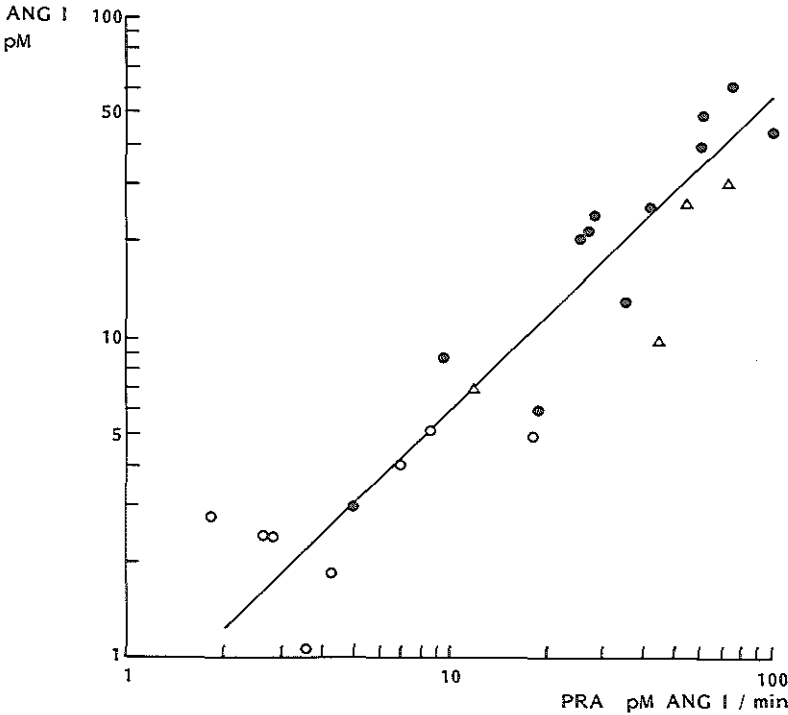


Figure 1. Relationship between the aortic plasma levels of PRA and ANG I in untreated (○), captopril-treated (●) and furosemide-treated (△) pigs ($\log \text{ANG I} = 0.89 \cdot \log \text{PRA} - 0.12$, $r = 0.92$).

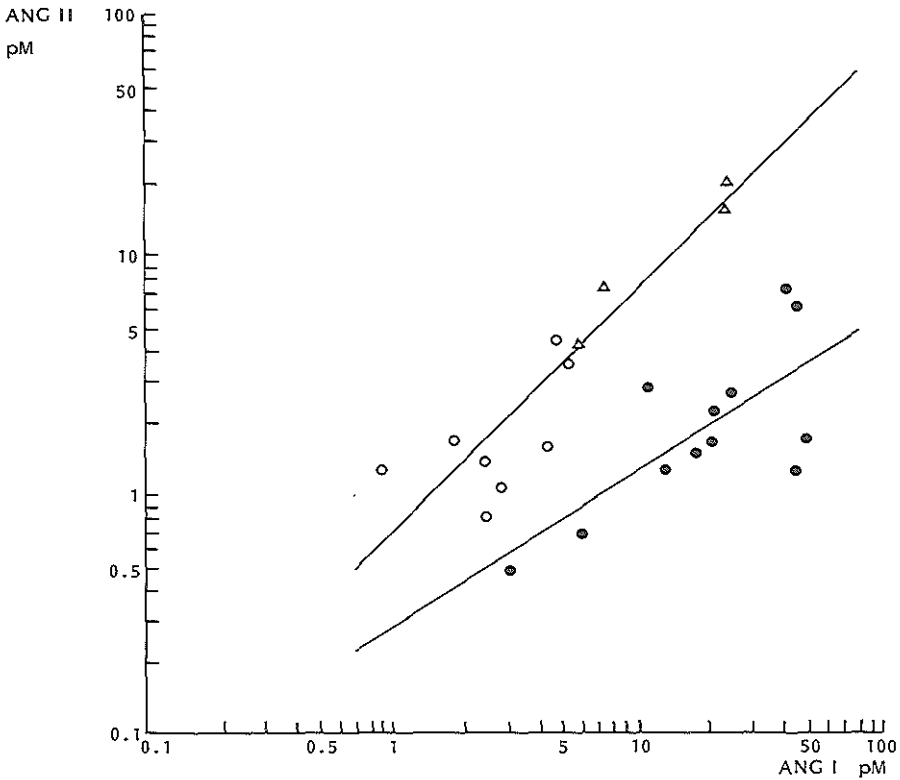


Figure 2. Relationship between the aortic plasma levels of ANG I and ANG II in untreated (o) and furosemide-treated (Δ) pigs (upper line; $\log \text{ANG II} = 1.03 \cdot \log \text{ANG I} - 0.17$, $r=0.91$) and in captopril-treated pigs (\bullet) (lower line; $\log \text{ANG II} = 0.63 \cdot \log \text{ANG I} - 0.53$, $r=0.70$).

In contrast with ANG I, plasma ANG II was lower in the right atrium than in the aorta, and it was higher in the left atrium than in the pulmonary artery (Table 1). Again, these arteriovenous differences were seen in the untreated and furosemide-treated animals but not in the captopril-treated pigs. Thus, in the absence of ACE inhibition, there was net removal of ANG II from the systemic circulation by the combined systemic vascular beds, and net delivery of ANG II to the systemic circulation by the lungs. Across the lungs, the veno-arterial difference in ANG II was close to the arteriovenous difference in ANG I. This is in accordance with our previous work, showing little degradation of both ANG I and ANG II in the lungs (5), so that most of the arterial ANG I delivered to the lungs and converted in this organ is recovered as ANG II in the blood of the left atrium.

Table 1. Plasma levels of ANG I , ANG II and PRA.

	NO TREATMENT n=8	CAPTOPRIL GROUP 1 n=8	CAPTOPRIL GROUP 2 n=4	FUROSEMIDE n=4
<u>ANG I pM</u>				
Aorta	2.9 (1.4)	21.5 (14.7)	30.4 (16.6)	14.2 (7.9)
Right Atrium	3.5 (2.0)*	23.2 (14.9)	-	18.8 (10.6)*
Pulmonary Artery	3.4 (1.7)*	22.2 (15.0)	-	18.0 (11.0)*
Left Atrium	2.1 (0.9)	20.9 (15.5)	-	13.4 (7.9)
Great Cardiac Vein	2.9 (1.2)	20.8 (13.9)	-	12.8 (7.1)
Jugular Vein	-	-	29.8 (14.0)	18.4 (11.3)**
Ear Lobe Vein	-	-	33.7 (18.2)	-
Femoral Vein	-	-	31.5 (18.0)	-
Renal Vein	-	-	45.6 (28.5)**	-
<u>ANG II pM</u>				
Aorta	2.0 (1.3)	2.2 (2.2)	3.1 (1.9)	12.4 (6.5)
Right Atrium	1.7 (1.1)*	1.8 (1.3)	-	10.3 (5.2)*
Pulmonary Artery	1.4 (0.9)*	1.5 (1.4)	-	8.9 (4.7)*
Left Atrium	2.2 (1.8)	2.2 (2.0)	-	13.3 (7.6)
Great Cardiac Vein	1.8 (0.8)	2.3 (2.1)	-	13.0 (6.9)
Jugular Vein	-	-	3.3 (2.5)	12.0 (6.7)
Ear Lobe Vein	-	-	2.8 (1.2)	-
Femoral Vein	-	-	2.8 (1.5)	-
Renal Vein	-	-	2.0 (1.2)***	-
<u>PRA pM ANG I/min</u>				
Aorta	7.4 (5.7)	48.8 (39.8)	37.6 (18.8)	59.8 (31.5)
Right Atrium	7.7 (6.5)	49.9 (39.3)	-	55.8 (26.0)
Pulmonary Artery	7.8 (6.7)	50.4 (37.7)	-	59.8 (29.7)
Left Atrium	7.4 (5.8)	46.5 (32.9)	-	56.0 (26.6)
Great Cardiac Vein	7.9 (7.5)	49.5 (37.8)	-	60.0 (30.7)
Jugular Vein	-	-	37.5 (19.7)	51.1 (26.3)
Ear Lobe Vein	-	-	38.8 (17.5)	-
Femoral Vein	-	-	33.6 (16.7)	-
Renal Vein	-	-	75.8 (50.2)**	-

Data are means and SD.

* in untreated and furosemide-treated pigs, angiotensin plasma levels in right atrium and pulmonary artery were significantly different from plasma levels in aorta and left atrium ($p < 0.05$, Scheffé's test for multiple comparison).

** levels were higher than in aorta in all 4 experiments.

*** levels were lower than in aorta in all 4 experiments.

Regional production of angiotensin I

As demonstrated by the data in Fig. 3, the regional production rates in the systemic vascular beds were high enough to compensate for the rapid elimination of this peptide. In the lungs all ANG I in venous plasma appears to be derived from arterial delivery and not from de novo production. In the combined systemic vascular beds, and in myocardium, head, skin, skeletal muscle and kidney, respectively 59 (14), 40 (13), 58 (6), 55 (11), 67 (11) and 94 (5) % of venous ANG I was derived from de novo production (mean and SD). In Table 2, the ANG I production rates in the combined systemic vascular beds, the lungs and the myocardium are given. ANG I production was increased about 10-fold in the captopril- and furosemide-treated pigs as compared with the untreated pigs.

Table 2. ANG I and ANG II production rates in the combined systemic vascular beds, the lungs and the myocardium.

		NO TREATMENT	CAPTOPRIL	FUROSEMIDE
<u>Combined Systemic Vascular Beds</u>				
ANG I production	(pmol/min)	3.5 (2.2-12.8)	43.0 (8.6-180)*	50.1 (27.2-85.0)*
ANG II production	(pmol/min)	1.6 (0.8-5.1)	3.6 (0.5-7.3)	21.7 (11.6-30.6)*
<u>Lungs</u>				
ANG I production	(pmol/min)	0.2 (0.0-0.6)	0.3 (0.0-3.6)	0.7 (0.0-1.5)
ANG II production	(pmol/min)	0.8 (0.4-4.7)	1.0 (0.2-3.4)	7.4 (2.2-16.5)*
<u>Myocardium</u>				
ANG I production	(fmol/min)	39.4 (12.2-71.5)	319 (95.4-1035)*	252 (78.8-741)*
ANG II production	(fmol/min)	9.1 (2.9-23.1)	18.7 (7.3-157)	151 (49.5-397)*

Data are medians and ranges.

Calculations for the combined systemic vascular beds and the myocardium were based on equations (1) and (6), and for the lungs on equations (3) and (8). The cardiac output of plasma was 1.71 (0.28) L/min in the no treatment group, 1.79 (0.43) L/min in the captopril group, and 2.30 (0.52) L/min in the furosemide group (means and SD). Left anterior descending coronary artery plasma flow, which was used for the calculations of ANG I and ANG II production by the myocardium, was 20.3 (5.7) mL/min in the no treatment group, 29.2 (8.5) mL/min in the captopril group, and 24.7 (8.9) mL/min in the furosemide group. Since total coronary plasma flow is larger than the left anterior descending coronary artery plasma flow, the actual myocardial production rates will be larger than the figures presented here.

* significantly different from no treatment group ($p < 0.005$, Mann-Whitney U-test for unpaired observations).

The ANG I extraction ratios across the combined systemic vascular beds in these groups of animals were not significantly different (Fig. 3). As mentioned above, the arterial ANG I/PRA ratios in the various animal groups were also not different. The ANG I production rates were therefore linearly correlated with the circulating levels of PRA, and for a given level of PRA, the regional ANG I production rates in the untreated and furosemide-treated groups were not different from the production rate in the captopril-treated group.

When the ANG I production in the systemic vascular beds by circulating PRA (equation 4) is expressed as a percentage of the total regional ANG I production (equation 1), one does not need to know the regional plasma flow but one needs to know the blood transit time. Table 3 gives for a number of systemic vascular beds the percent contribution of circulating PRA to total regional ANG I production, assuming a blood transit time of 10 sec (13,15,23). Probably, the capillaries are the main site where ANG I is removed from the circulation by diffusion into the interstitium and peptidase attack. This is also the site where ANG I from the interstitium equilibrates with circulating ANG I. The blood transit times in the capillary beds is 1-3 sec (11). Our estimate of 10 sec is therefore probably too high, but this overestimation strengthens the conclusion that a substantial part of ANG I production in the systemic vascular beds we studied could not be accounted for by circulating PRA. There was little ANG I production in the pulmonary vascular bed; the small amount that was produced could be accounted for by PRA assuming a pulmonary blood transit time of 3 sec.

Table 3. Percentage of the total regional ANG I production in different beds that can be attributed to circulating PRA.

VASCULAR BED	ANG I production by circulating PRA (% of total regional production)	
	NO CAPTOPRIL	CAPTOPRIL
Lungs	119 (70)	109 (67)
Combined Systemic Vascular Beds	38 (18)*	31 (17)*
Myocardium	76 (24)*	64 (40)*
Head	44 (14)*	18 (7)*
Skin	-	16 (4)*
Skeletal Muscle	-	11 (5)*
Kidney	-	2 (2)*

Data are means and SD.

The 'no captopril' group comprises both the untreated and the furosemide-treated pigs. Calculations were based on equations (1), (3) and (4), assuming a blood transit time of 10 sec in the systemic vascular beds and 3 sec in the lungs (see Results section).

* significantly different from 100 ($p < 0.05$, Student's t-test for paired observations).

$(\text{ANG I}_{\text{ven}} \text{ derived from } \text{ANG I}_{\text{art}}) / (\text{ANG I}_{\text{art}})$

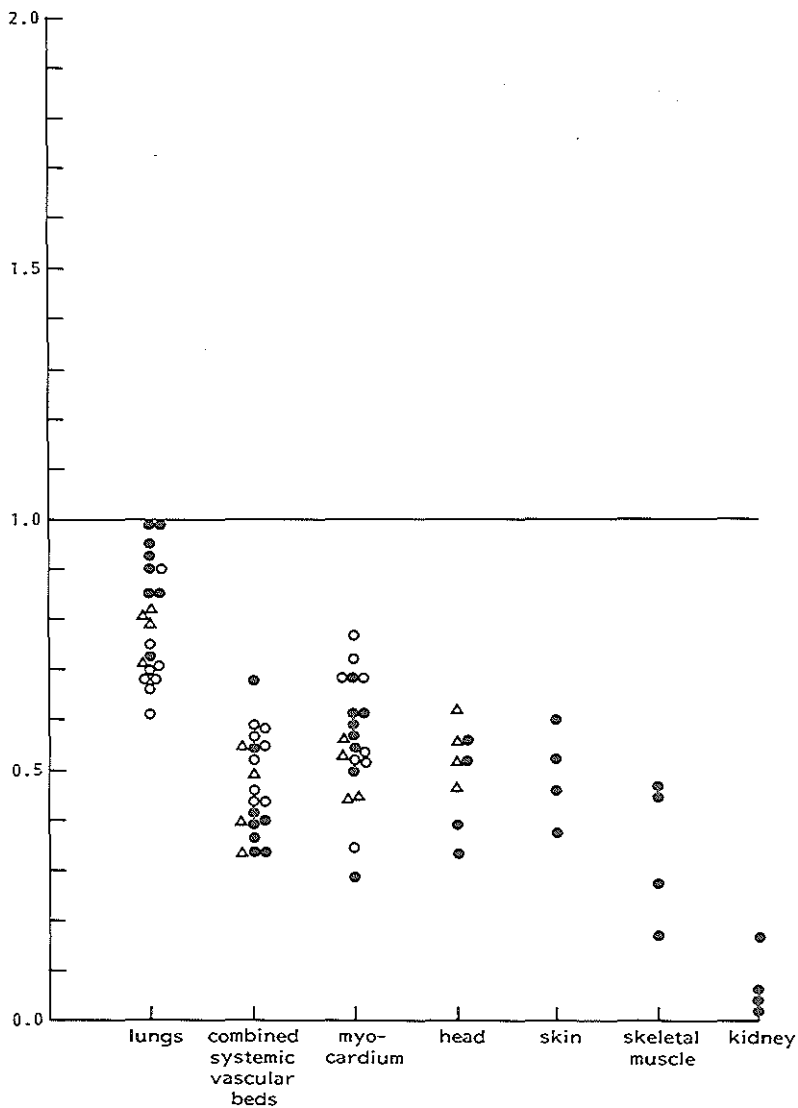
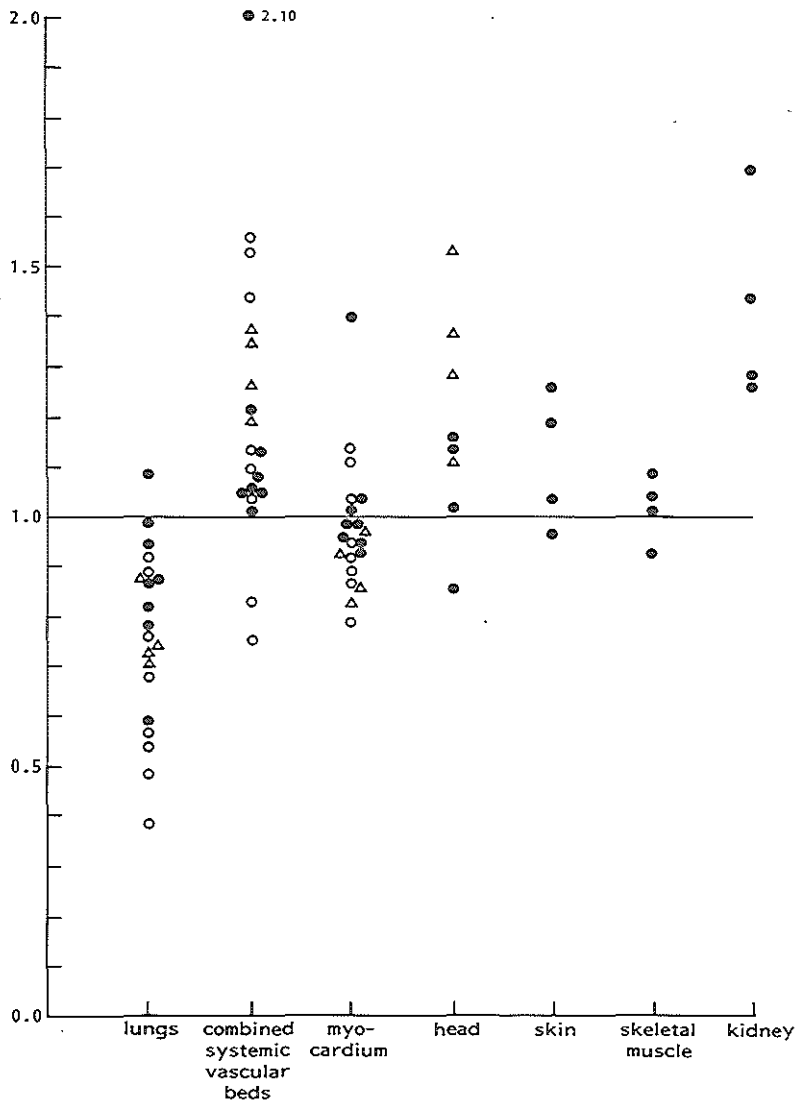


Figure 3. Calculated venous plasma levels of ANG I derived from arterially delivered ANG I (left panel) vs. the actually measured venous plasma levels of ANG I (right

$(\text{ANG I}_{\text{ven}}) / (\text{ANG I}_{\text{art}})$



panel) in different vascular beds in untreated (o), captopril-treated (●) and furosemide-treated (Δ) pigs. Calculations were made on the basis of equation (5). The venous ANG I levels are expressed as a fraction of the arterial level of ANG I.

Regional production of angiotensin II

The rapid elimination of ANG II in the systemic vascular beds was compensated for by regional production of this peptide, with the kidney as an exception (Fig. 4). The ANG II level in the renal vein was 60-70 percent of that in the artery. In the present study the renal extraction of ANG II and the arterial and venous levels of ANG II across the kidney were only measured in pigs on captopril treatment, but studies in humans also showed that the levels of ANG II in the renal vein are lower than those in the renal artery, both during captopril treatment and without such treatment (1,21). There is also net ANG II extraction from the systemic circulation by the liver (1). Since about 40 percent of the cardiac output flows through kidneys and liver, the net extraction by these organs of systemically delivered ANG II explains why the ANG II level in the right atrium was lower than in the aorta, despite the absence of an arteriovenous ANG II gradient across the other vascular beds.

In the lungs, in the combined systemic vascular beds, and in myocardium, head, skin, skeletal muscle and kidney, respectively 31 (22), 39 (19), 32 (20), 49 (8), 40 (11), 59 (15) and 85 (8) % of ANG II in venous plasma was derived from *de novo* production (mean and SD). In Table 2 the ANG II production rates in the combined systemic vascular beds, the lungs and the myocardium are given. The ANG II production rate in the combined systemic vascular beds was about 2 times higher than that in the lungs, which may illustrate the important contribution of extrapulmonary ANG I-II conversion to the whole body ANG II production.

Table 4. Fractional conversion of arterially delivered ANG I in different vascular beds.

VASCULAR BED	NO TREATMENT	CAPTOPRIL	FUROSEMIDE
Lungs	0.25 (0.07)	0.04 (0.04)*	0.22 (0.04)
Combined Systemic Vascular Beds	0.10 (0.02)	0.01 (0.01)*	0.12 (0.04)
Myocardium	0.17 (0.02)	0.02 (0.02)*	0.18 (0.02)
Head	-	0.04 (0.02)**	0.16 (0.01)
Skin	-	0.02 (0.02)	-
Skeletal Muscle	-	0.07 (0.02)	-
Kidney	-	0.08 (0.02)	-

Data are means and SD.

Fractional conversion was calculated by using equation (9).

* significantly different from 'no treatment' group ($p < 0.01$, Student's t-test for unpaired observations).

** significantly different from 'furosemide' group ($p < 0.01$, Student's t-test for unpaired observations).

Data on fractional ANG I-II conversion in the various systemic vascular beds we studied are presented in Table 4. On the basis of these data, the percent contribution of conversion of arterially delivered ANG I to the total regional production of ANG II was calculated. The results are shown in Table 5. They indicate that a substantial part of ANG II production in the systemic vascular beds could not be accounted for by conversion of arterially delivered ANG I. ANG II production in the lungs could fully be accounted for by the conversion of ANG I from the circulation.

Table 5. Percentage of the total regional ANG II production in different vascular beds that can be attributed to conversion of arterially delivered ANG I.

VASCULAR BED	ANG II production by conversion of circulating ANG I (% of total regional production)			
	<u>NO CAPTOPRIL</u>		<u>CAPTOPRIL</u>	
	production by conversion of arterially delivered ANG I	production by conversion of ANG I from PRA	production by conversion of arterially delivered ANG I	production by conversion of ANG I from PRA
Lungs	116 (79)	13 (13)*	117 (73)	10 (16)*
Combined Systemic Vascular Beds	26 (18)*	10 (4)*	16 (14)*	6 (5)*
Myocardium	57 (27)*	17 (22)*	41 (30)*	18 (13)*
Head	35 (3)*	18 (2)*	33 (7)*	7 (2)*
Skin	-	-	38 (27)*	8 (6)*
Skeletal Muscle	-	-	42 (16)*	9 (5)*
Kidney	-	-	12 (4)*	4 (1)*

Data are means and SD.

The 'no captopril' group comprises both the untreated and the furosemide-treated pigs. ANG II production by conversion of arterially delivered ANG I was calculated by using equation (10). ANG II production by conversion of ANG I from PRA was calculated by multiplying the regional ANG I production by PRA (equation 4) with the fractional conversion of arterially delivered ANG I (data presented in Table 4). Blood transit times were assumed to be 10 sec in the systemic vascular beds and 3 sec in the lungs (see Results section).

* significantly different from 100 ($p < 0.05$, Student's t-test for paired observations).

$(\text{ANG II}_{\text{ven}} \text{ derived from } \text{ANG II}_{\text{art}} \text{ and from regional conversion of } \text{ANG I}_{\text{art}}) / (\text{ANG II}_{\text{art}})$

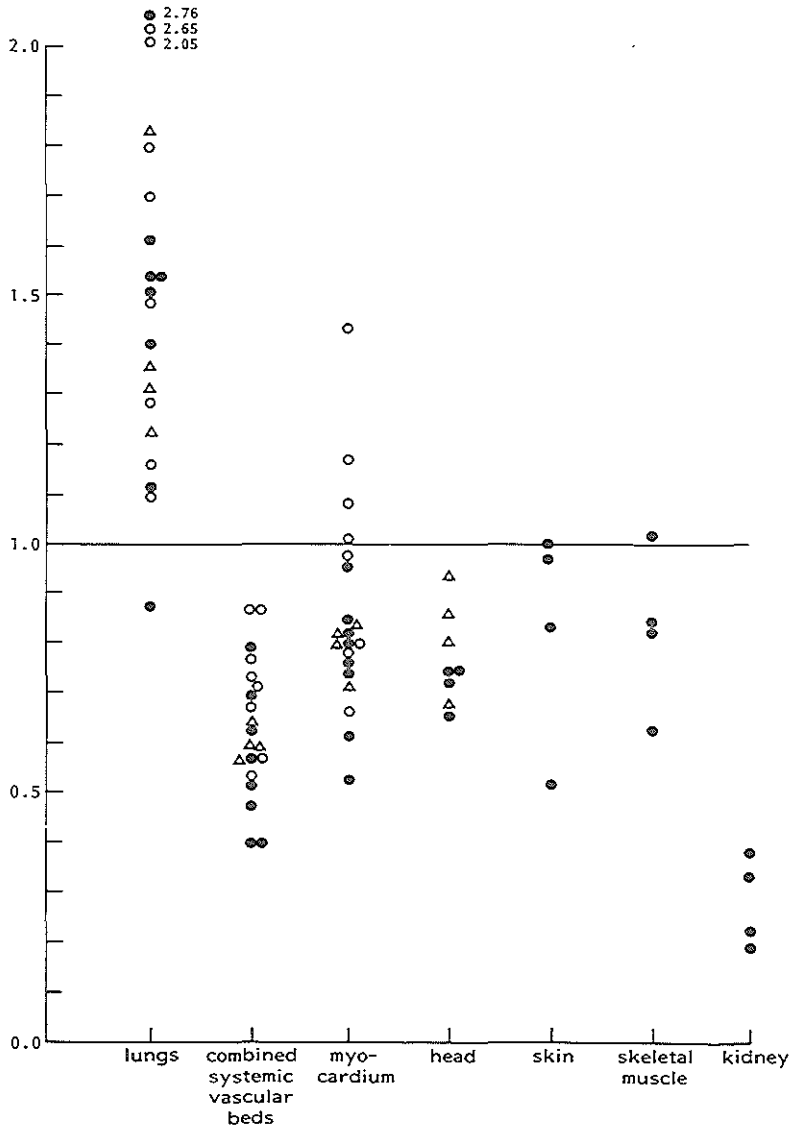
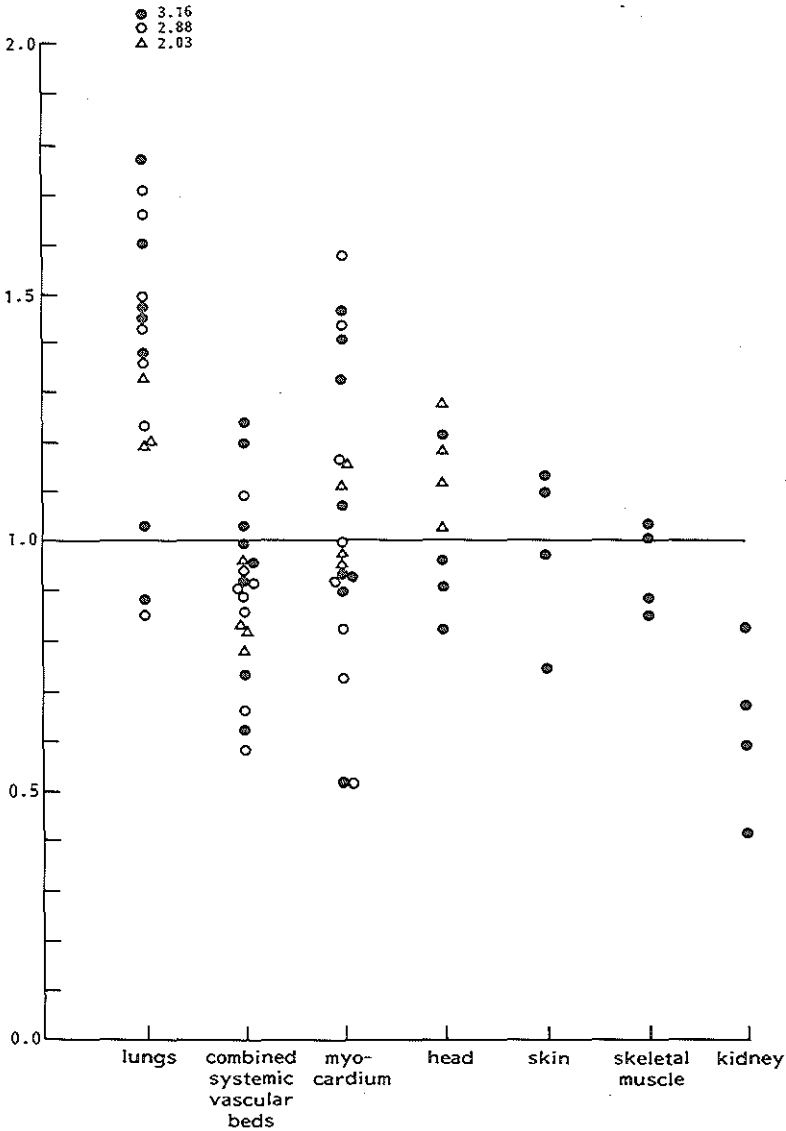


Figure 4. Calculated regional venous plasma levels of ANG II derived from arterially delivered ANG II and from conversion of arterially delivered ANG I (left panel) vs. actually measured venous plasma levels of ANG II (right panel) in different vascular beds in

$(\text{ANG II}_{\text{ven}}) / (\text{ANG II}_{\text{art}})$



untreated (o), captopril-treated (●) and furosemide-treated (Δ) pigs. Venous ANG II derived from arterially delivered ANG II and from regional conversion of arterially delivered ANG I was calculated on the basis of equations (11) and (12). The venous ANG II levels are expressed as a fraction of the arterial level of ANG II.

Conversion in a vascular bed of ANG I generated by PRA during the transit of blood through that bed would also contribute to the regional de novo production of ANG II. Assuming that in a vascular bed the fractional conversion of this regionally produced ANG I is the same as the fractional conversion of ANG I that is arterially delivered to that bed, one can calculate the contribution of PRA to the total regional production of ANG II. As shown in Table 5, this contribution of PRA is small and does not make up for the difference between total regional ANG II production and regional ANG II production from arterially delivered ANG I. Thus, part of ANG II production in the circulation does not appear to originate from ANG I in the circulation; it may originate from the conversion of ANG I produced at tissue sites.

4.5 Discussion

Estimates of ANG I and ANG II production rates were made from measurements of the arterial and venous plasma levels of labeled ANG I and ANG II as well as endogenous ANG I and ANG II during constant infusion of ^{125}I -ANG I. ^{125}I -ANG I was infused directly into the left cardiac ventricle. We assumed that by this way of administration the route followed by exogenous ANG I resembled that of endogenous ANG I as closely as possible. ^{125}I -ANG I was extensively metabolized in all systemic vascular beds we studied (5). However, the venous levels of endogenous ANG I were similar to or even higher than the arterial levels of ANG I. Venous ANG I is therefore largely derived from regional de novo ANG I production. By measuring the circulating levels of plasma renin activity (PRA) in conjunction with those of radiolabeled and endogenous ANG I we were able to calculate how much of the regional de novo production of ANG I was due to circulating PRA and how much was not. Our results indicated that a major part of the de novo production of ANG I in the systemic vascular beds we studied did not originate from the action of circulating renin on circulating angiotensinogen. These calculations were based on a blood transit time of 10 sec. As mentioned under 'Results', this is probably an underestimation, but it only strengthens our conclusion that PRA could not account for the de novo ANG I production. This appeared to be the case not only in the kidneys but also in a number of other vascular beds.

The evidence for local ANG I production, i.e. ANG I production not in circulating blood, which was obtained in pigs treated with the ACE inhibitor captopril as well as in untreated animals, confirms our earlier observations in humans, all on captopril (1). Some untreated pigs had plasma ANG I levels as low as 2-3 times the detection limit of the assay. We therefore also studied a group of furosemide-treated animals. Results in these animals confirmed the conclusions drawn from the measurements in the untreated group.

The local production of ANG I in extrarenal tissues, i.e. ANG I production

not due to circulating PRA, may still depend on the presence of kidney-derived renin in vascular tissue rather than on in situ synthesized renin (17,22). In bilaterally nephrectomized patients the plasma levels of renin and ANG I are known to be very low (4,7,20). The production of ANG I in plasma and at tissue sites in these patients may therefore be expected to be low as well. Thus, kidney-derived renin may indeed be responsible for most of the extrarenal production of ANG I at tissue sites, at least for that fraction of ANG I in the tissues that equilibrates with circulating ANG I. In accordance with this conclusion is the fact that the spontaneous release of angiotensins by isolated perfused organs is very low (rat hindquarter) (12) or even undetectable (rat heart) (16). Perfusion of rat hindquarters and heart with renin results in a dose-dependent release of ANG I and ANG II (12,16).

Our calculations are based on the so-called 'venous equilibrium' model for the intrinsic clearance of exogenous and endogenous substances by tissues (25). This model considers the distribution volume of these substances to be a single, well-stirred compartment; their concentration in this compartment equals that in the venous effluent. The volume of blood present in the conduit vessels is not taken into consideration. The implicit assumption in our calculations is therefore that the arterial and venous plasma levels of the radiolabeled and endogenous angiotensins we measured are representative for the levels at the arterial and venous side of the microcirculation. This assumption is probably valid, because there is little metabolism of angiotensins in the large vessels and because PRA may be expected to cause only small gradients in ANG I and ANG II along the large vessels. Indeed, in a study in humans we found no differences in the levels of ^{125}I -ANG I, ^{125}I -ANG II, and ANG I and ANG II between blood samples taken deeply from the femoral vein and from the iliac vein and inferior caval vein just under the entrance of the renal veins (Admiraal, unpublished results). We also found no differences in plasma levels between the ascending aorta and the femoral artery in captopril-treated pigs (data not shown).

From our data it could be calculated that not only ^{125}I -ANG I but also ^{125}I -ANG II was rapidly metabolized in the systemic vascular beds we studied. Again, as for ANG I, the venous levels of ANG II were similar to the arterial levels, indicating that ANG II is regionally produced. The lungs are an important site of ANG II production, but it is of some interest that the rate of de novo production of ANG II in the combined systemic vascular beds was higher than in the lungs. ANG II production in the pulmonary and the combined systemic vascular beds in the captopril-treated pigs was not lower than in the untreated pigs, despite the fact that ANG I-II conversion was greatly reduced. Thus, the inhibition of ANG I-II conversion by captopril was overcome by increased ANG I production.

Regional de novo production of ANG II may be due to conversion of circulating ANG I, i.e. ANG I that is arterially delivered or ANG I that is regionally produced by PRA. De novo production of ANG II in the lungs could

be fully attributed to conversion of circulating ANG I. In contrast, a substantial part of the de novo production of ANG II in the systemic vascular beds we studied did not originate from the conversion of circulating ANG I. In the kidneys by far the most of the de novo produced ANG II does not seem to originate from circulating ANG I. ANG I-II conversion is probably the main pathway of ANG II production. In the systemic vascular beds some of the ANG II production, and in the kidneys probably most of it, may therefore originate from the conversion of ANG I produced at tissue sites. However, alternative pathways for ANG II generation (e.g. direct cleavage from angiotensinogen) have been described (24).

Whereas our results do provide evidence for the local formation of ANG I and ANG II in a number of tissues and for the release of these locally formed angiotensins into the circulation, the results do not militate against the fact that most of the circulating ANG I is formed by the action of circulating renin on circulating angiotensinogen and that most of the circulating ANG II is derived from circulating ANG I. Our calculations of ANG I and ANG II production at tissue sites did not include the production of these peptides in the blood of the large arteries and veins. The volume of blood in the microcirculation of the tissues is only a small fraction of the total blood volume in the body. The total amount of circulating renin in the large arteries and veins is therefore much greater than in the microcirculation, and its contribution to the whole body production of ANG I is therefore also much greater. The local formation of both ANG I and II in tissues may have important physiological implications, in that it may serve to generate sufficiently high angiotensin concentrations at tissue receptor sites, concentrations that could well be higher than in the circulation.

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5. RENIN, PRORENIN AND IMMUNOREACTIVE RENIN IN VITREOUS FLUID FROM EYES WITH AND WITHOUT DIABETIC RETINOPATHY

5.1 Summary

Renin, prorenin and immunoreactive renin were present in vitreous and subretinal fluid of eyes from subjects with and without diabetic retinopathy. Renin substrate, albumin, transferrin and immunoglobulin G were also found in these ocular fluids. In many samples renin levels were close to the detection limit of the assay. The levels of renin substrate, albumin, transferrin and immunoglobulin G varied widely among ocular fluid samples, but in each individual sample the levels were, relative to each other, similar to those in plasma. In contrast, the prorenin level in ocular fluid was up to 100 times higher than expected on the basis of the plasma protein content of ocular fluid. Moreover, there was little difference in prorenin concentrations between samples with a low and a high plasma protein content. Prorenin, relative to albumin and other plasma proteins, was higher in vitreous fluid from eyes with proliferative diabetic retinopathy complicated by traction retinal detachment than in eyes of non-diabetic subjects with spontaneous retinal detachment. It appears that prorenin (and possibly renin) in ocular fluid is controlled by an active and specific process, possibly local synthesis within the eye. In view of the vascular actions of angiotensin II, an intraocular renin-angiotensin system may play a role in diabetic retinopathy.

5.2 Introduction

The kidney secretes both renin and prorenin, an inactive precursor of renin, into the circulation. Plasma of nephrectomized patients contains little or no renin but it does contain prorenin (1,2), in concentrations sometimes as high as those in normal individuals. It thus appears that extrarenal production can make a major contribution to the level of prorenin in plasma, whereas most, if not all, renin in plasma is secreted by the kidneys. Synthesis of renin or prorenin and other components of the renin-angiotensin system is known to occur at various extrarenal sites, for instance adrenal (3,4) pituitary (3,4), testis (3,4), brain (5), and ovary (6-8). Cultured human chorionic cells (9)

and ovarian thecal cells (8) release prorenin into the medium and there is good evidence that in women with hyperstimulated cycles and during pregnancy, the ovary, probably the corpus luteum, releases prorenin into plasma (7,10).

A common feature of the organs in which synthesis of renin or prorenin occurs is their extensive vascularization (11). The eye, particularly the retina and uveal tract, is such a highly vascularized organ. Angiotensin II-binding sites have been found in retinal blood vessels (12), and transvitreal infusion of angiotensin I and II produces constriction of the retinal arteries (13). The retina contains angiotensin converting enzyme activity (14) and this enzyme is also found in aqueous fluid (15). Here we report measurements of renin, prorenin, immunoreactive renin, renin substrate and various plasma proteins in aqueous, vitreous and subretinal fluid. The ocular fluid samples were obtained at the time of cataract extraction or vitrectomy and the protein concentrations in these samples were compared with those in simultaneously obtained plasma. Our study included eyes affected by proliferative diabetic retinopathy because the renin-angiotensin system has been implicated in neovascularization (16).

5.3 Subjects and methods

Non-diabetic subjects

Aqueous fluid was collected at the time of cataract extraction from 21 subjects (15 women and 6 men; mean age, 68 yr; range, 26-86 yr). Four subjects were receiving a diuretic and six a β -adrenergic antagonist.

Vitreous fluid aspirates were obtained from 16 subjects (8 women and 8 men; mean age, 52 yr; range, 20-82 yr). The samples were collected at the time of pars plana vitrectomy, which was performed because of recurrent retinal detachment due to proliferative vitreoretinopathy. Four subjects were receiving a diuretic, in 3 of them combined with a β -adrenergic antagonist.

Subretinal fluid was obtained from 18 subjects (8 women and 10 men; mean age, 59 yr; range, 8-76 yr), with rhegmatogenous retinal detachment, which is a type of retinal separation precipitated by a hole or a tear in the retina. In this type of detachment fluid accumulates between the retinal pigment epithelial layer and the neural retina. The retinal detachments had occurred between 1 day and 3 months (median, one week) before subretinal fluid collection. Three subjects were receiving a diuretic, in 2 of them combined with a β -adrenergic antagonist.

Diabetic subjects

Vitreous fluid was obtained from 15 diabetic subjects with proliferative diabetic retinopathy (8 women and 7 men; mean age, 51 yr; range, 28-71 yr). Vitrectomy was performed because of traction retinal detachment. The

duration of diabetes ranged from 6-32 yr. Twelve subjects were receiving insulin, 5 were receiving a diuretic and 1 was receiving a β -adrenergic antagonist.

Aqueous fluid can only be collected at the time of cataract extraction. In diabetic subjects, however, this procedure may stimulate proliferative retinopathy. Cataract extraction is therefore not performed in eyes affected by proliferative diabetic retinopathy. Consequently, aqueous fluid could not be collected from such eyes. We also were unable to collect subretinal fluid from diabetic subjects with traction retinal detachment because drainage of subretinal fluid is rarely performed in these subjects and, if it is performed, the approach is via the transvitreal route, so that the sample is heavily contaminated with material from the vitreous. In subjects with a rhegmatogenous retinal detachment, subretinal fluid is removed via the transscleral route, where no such contamination occurs.

Collection of ocular fluid samples

Approximately 0.1 mL aqueous fluid was collected with a tuberculin syringe and a 25-gauge needle. The needle was introduced at the limbus of the cornea through the groove of the cataract incision. A 0.3-1.0 mL sample of vitreous fluid was aspirated before substitution fluid was infused into the vitreous. Subretinal fluid was aspirated transsclerally, after local diathermic coagulation of the choroid.

The ocular fluid samples were free of macroscopically visible blood and were frozen at -70°C immediately after collection. A peripheral venous blood sample was drawn simultaneously with the collection of ocular fluid. Blood for determination of renin, prorenin, immunoreactive renin, renin substrate, albumin, transferrin and immunoglobulin G (IgG) was collected in tubes containing 0.1 volume of 0.13 mol/L trisodium citrate. The blood was immediately centrifuged at $3000 \times g$ for 10 minutes at room temperature, and 1-mL aliquots of plasma were stored at -70°C . Blood for determination of angiotensin II was collected in prechilled tubes containing 0.1 volume of 0.06 mmol/L pepstatin-A, 0.125 mol/L disodium EDTA and 0.025 mol/L phenantroline in order to block renin, angiotensin converting enzyme and angiotensinases, respectively. The blood samples were immediately centrifuged at $3000 g$ for 10 minutes at 4°C , and 2-mL aliquots of plasma were stored at -70°C .

Analytical methods

Renin was measured in duplicate by enzyme-kinetic assay, in which the samples were incubated at 37°C and pH 7.5 with saturating amounts of sheep renin substrate in the presence of inhibitors of angiotensinases and angiotensin converting enzyme. The generated angiotensin I was quantitated by RIA (17).

For measuring prorenin in plasma, prorenin was converted into renin by incubation with Sepharose-bound trypsin (0.25 mg/mL) for 48 hours at 4 °C. Previous studies, including measurements of total immunoreactive renin (renin plus prorenin), indicated that the prorenin to renin conversion in plasma is complete after incubation with the immobilized trypsin under these circumstances and that destruction of renin or prorenin does not occur (18). Experiments in which known quantities of purified human kidney renin were added to ocular fluid demonstrated that in some samples destruction of renin did occur with this method. This destruction might be due to the low content of serine protease inhibitors in ocular fluids as compared to plasma. Therefore, in ocular fluid we chose to use plasmin to convert prorenin into renin (17,19). For this purpose the sample was incubated with plasmin at a final concentration of 0.5 $\mu\text{mol/L}$ for 48 hours at 4 °C before the assay.

Comparison of the results of the enzyme-kinetic assay in plasmin-activated ocular fluid samples with the results of the assay of total immunoreactive renin in non-activated samples demonstrated that the conversion by plasmin was complete without any loss of prorenin or renin; the specific enzymatic activity of plasmin-activated prorenin in ocular fluid samples was not different from the specific activity of purified kidney renin and plasma renin (see *Results*). Plasmin at the concentration mentioned above cannot be used to activate prorenin in native whole plasma because of its high content of plasmin inhibitors.

The concentrations of renin and prorenin measured by the enzyme-kinetic assay were expressed as milliunits per L using the WHO human kidney renin standard 68/356 (WHO International Laboratory for Biological Standards, London, United Kingdom) as reference standard. The lower limit of detection was 0.5 mU/L and the interassay variability at low concentrations of renin or prorenin (2-5 mU/L) was 11 % for both renin and prorenin. Immunoreactive renin was measured in duplicate with a sandwich assay (18,20) using the monoclonal antibodies R 3-27-6 and R 3-36-16 (Ciba-Geigy, Basel, Switzerland). The two monoclonal antibodies recognize different epitopes of the renin molecule and react equally well with human kidney renin and chorionic cell culture prorenin. The assay was carried out in polystyrene tubes (Star Tubes, code 4-70319; Nunc, Roskilde, Denmark). The inner surface of these tubes was coated with antibody R 3-27-6 (21). Immunoreactive renin in the assay sample is quantitatively bound to this antibody. The amount of solid phase-bound immunoreactive renin was measured with antibody R 3-36-16, which had been radiolabeled with ^{125}I . The results of this assay were expressed as nanograms per L using highly purified human kidney renin (Ciba-Geigy) as a standard. One milliunit of the WHO human kidney renin standard corresponded to 1.41 ng of the Ciba-Geigy standard. The lower limit of detection was 5 ng/L, and the interassay variability was 8 %.

The concentration of renin substrate was determined as the maximum quantity of angiotensin I that was generated during incubation at 37 °C and pH

7.5 with an excess of purified active human kidney renin in the presence of inhibitors of angiotensinases and angiotensin converting enzyme (18). The lower limit of detection was 1 nmol/L and the interassay variability was 10 %.

Immunoreactive angiotensin II was measured by radioimmunoassay after SepPak (Waters, Milford, MA, USA) extraction of the sample (22). The lower limit of detection was 2 pmol/L and the interassay variability was 15%.

Albumin, transferrin and IgG were measured by single radial immunodiffusion (LC and NOR-Partigen plates, Behringwerke, Marburg, Germany) according to the method of Mancini et al (23).

Data analysis

Plasma proteins enter the vitreous mainly by diffusion. One of the reasons why the concentrations of these proteins are low in vitreous fluid is that they have to cross a relatively impermeable barrier. Breakdown of this so-called blood-retinal barrier leads to increased diffusion of plasma proteins into the eye. The rate of diffusion of a given protein is related to its molecular size and plasma concentration. In accordance with this is the fact that the concentrations of the different proteins relative to each other are similar in plasma and vitreous fluid (24,25).

Thus, unless certain specific uptake processes exist, for which in the eye no evidence is available with regard to any of the proteins mentioned in this paper, one would expect a relatively high intraocular albumin concentration (due to partial breakdown of the blood-retinal barrier) to be accompanied by a proportionally high concentration of plasma proteins of comparable size. Therefore, we chose to take the vitreous fluid/plasma concentration ratio of albumin as an index of the integrity of the blood-retinal barrier, an abnormally high ratio being an indication of breakdown of this barrier. By multiplying this ratio with the level of a given protein in plasma, the level of this protein in ocular fluid can be estimated, assuming that, as mentioned above, this protein is transferred from the blood into the vitreous and *vice versa* by mechanisms that are qualitatively and quantitatively the same as those for the transfer of albumin. For example, for renin substrate the calculation would be as follows:

$$[RS_{oc}] = [RS_{pl}] \times [ALB_{oc}]/[ALB_{pl}],$$

in which RS is renin substrate, ALB is albumin, oc is ocular fluid, pl is plasma, and brackets denote the concentration.

If our assumptions are correct, the calculated concentrations should be equal or at least closely correlated to the actually measured concentrations. Therefore, the two sets of data were analyzed by linear regression.

For analyzing differences between diabetic and nondiabetic subjects unpaired t-tests were performed after logarithmic transformation of the data.

Values were considered significant if $p < 0.05$.

5.4 Results

Non-diabetic subjects

The levels of renin in many vitreous and aqueous fluid samples were at or below the detection limit of the assay (0.5 mU/L), which is less than 5 % of the level in plasma. In subretinal fluid the renin level was about 20 % of that in plasma (Table 1). Prorenin was detectable in all samples of vitreous and aqueous fluid; its level in vitreous fluid was about 20 % and in aqueous fluid about 5 % of that in plasma. In subretinal fluid the prorenin level was as high as in plasma. Renin and prorenin concentrations in the fluid compartments of the eye were in the order: subretinal fluid > vitreous fluid > aqueous fluid. The levels of renin substrate in subretinal, vitreous and aqueous fluid were 10, 5 and 0.5 % of those in plasma, respectively. Thus, they too were in the order: subretinal fluid > vitreous fluid > aqueous fluid. This was also true for the levels of albumin, transferrin and IgG (Table 2). There was no correlation between the levels in ocular fluid and those in plasma for any of the proteins.

As described under *Data analysis* above, a theoretical concentration in ocular fluid for each protein was predicted based on the albumin content of the sample. For renin substrate, transferrin and IgG the calculated and measured values were linearly correlated in both vitreous and subretinal fluid, and the slopes of these correlation lines were not significantly different from 1.0 (Tables 3 and 4). Renin substrate and transferrin concentrations in ocular fluid could, in fact, be accurately predicted by these calculations. The IgG level measured was systematically about 2 times lower than that calculated, which may be due, at least in part, to its larger molecular size as compared to that of albumin and the other proteins.

For prorenin the findings were different. The prorenin levels in both vitreous and subretinal fluid varied much less than the levels of the other proteins (Tables 1 and 2). Furthermore, calculating prorenin concentrations on the basis of the albumin content of the sample yielded much lower (down to 1/100th) values than those actually measured, particularly in samples with a low plasma protein content (intact blood-retinal barrier). The prorenin level in subretinal fluid was higher than that in vitreous fluid, even when corrections were made for the higher plasma protein content in subretinal fluid samples (Fig. 1). In both vitreous and subretinal fluid the slopes of the regression lines describing the correlation between the measured and calculated prorenin concentrations were significantly different from 1.0, thereby indicating the different behavior of prorenin as compared to albumin and other plasma proteins.

The data on prorenin shown in Tables 1, 3 and 4 and Figs. 1-3 were obtained

Table 1. Levels of prorenin, renin, and renin substrate in ocular fluids.

	n	Prorenin (mU/L)		Renin (mU/L)		Renin substrate (nmol/L)	
		Eye	Plasma	Eye	Plasma	Eye	Plasma
<u>Non-diabetic subjects</u>							
Aqueous vs. plasma	21	4.4 2.0-8.7	163 36.7-453	< 0.5 ND-0.5	9.8 1.5-62.9	5.1 1.6-15.4	1080 898-1430
Vitreous vs. plasma	16	34.5 17.4-61.9	174 67.0-396	< 1.0 ND-2.8	17.3 3.7-51.0	54.5 3.0-630	1120 791-3030
Subretinal vs. plasma	18	132 36.8-305	128 65.1-251	2.4 0.9-6.4	14.3 4.3-105	107 10-1430	1230 841-2250
<u>Diabetic subjects</u>							
Vitreous vs. plasma	15	61.0 19.0-172	357 121-679	< 2.0 ND-3.5	17.7 4.4-127	61.0 7.0-1000	1030 628-2040

Shown are the geometric mean and range. ND, Not detectable. In vitreous fluid and plasma the levels of prorenin, but not those of renin substrate, were higher in diabetic than in non-diabetic subjects ($p < 0.01$).

Table 2. Levels of albumin, IgG, and transferrin in ocular fluids.

	n	Albumin (g/L)		IgG (g/L)		Transferrin (g/L)	
		Eye	Plasma	Eye	Plasma	Eye	Plasma
<u>Non-diabetic subjects</u>							
Aqueous vs. plasma	21	0.19 0.06-0.45	33.1 28.3-39.4	ND	ND	ND	ND
Vitreous vs. plasma	16	1.55 0.11-20.2	33.0 22.9-42.5	0.19 0.03-0.70	11.0 6.61-14.4	0.18 0.02-1.74	2.44 1.60-2.99
Subretinal vs. plasma	18	3.03 0.39-28.5	35.4 29.2-41.9	0.48 0.07-5.10	11.2 7.0-18.2	0.45 0.06-3.97	2.57 2.08-3.44
<u>Diabetic subjects</u>							
Vitreous vs. plasma	15	1.51 0.32-17.1	29.2 21.7-36.8	0.19 0.04-0.96	9.2 3.88-16.7	0.14 0.06-0.37	2.06 1.50-2.70

Shown are the geometric mean and range. The levels of albumin, IgG, and transferrin in both vitreous and plasma did not differ between diabetic and non-diabetic subjects. ND, Not done.

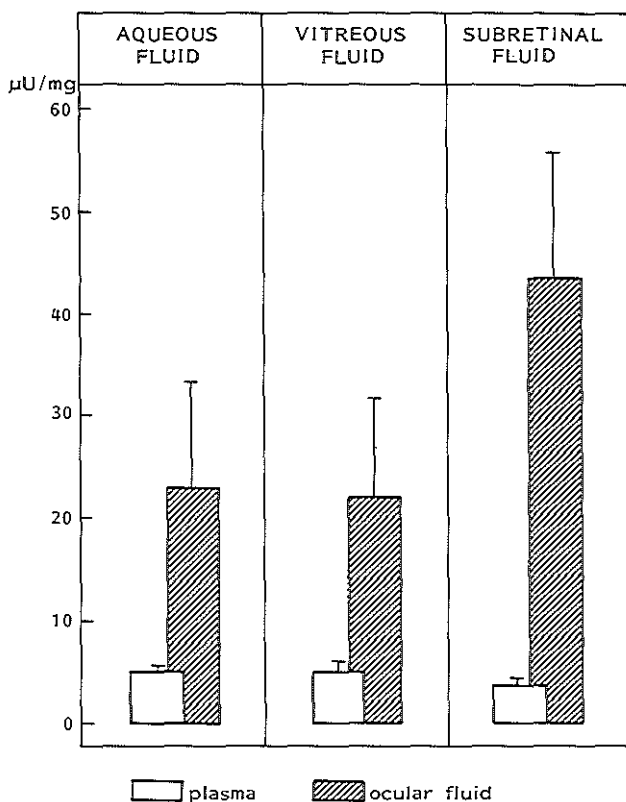


Figure 1. Mean (\pm SE) prorenin/albumin concentration ratio in ocular fluids and plasma of nondiabetic patients.

by the enzyme-kinetic assay. That prorenin measured by this assay is, in fact, prorenin is supported by the excellent agreement with the measurements of immunoreactive renin (Fig. 4). The mean specific enzymatic activity of in vitro activated prorenin was 0.7 ± 0.2 (\pm SD) mU/ng ($n=9$) in vitreous fluid and 0.6 ± 0.2 mU/ng ($n=10$) in subretinal fluid. These values are not different from the specific activity of renin from plasma and kidney (18).

Immunoreactive angiotensin II was 11.1 ± 1.8 pmol/L in vitreous fluid ($n=12$) compared to 17.5 ± 1.3 pmol/L in plasma. In subretinal fluid ($n=15$) it was 14.8 ± 1.6 pmol/L compared to 23.9 ± 2.0 pmol/L in plasma.

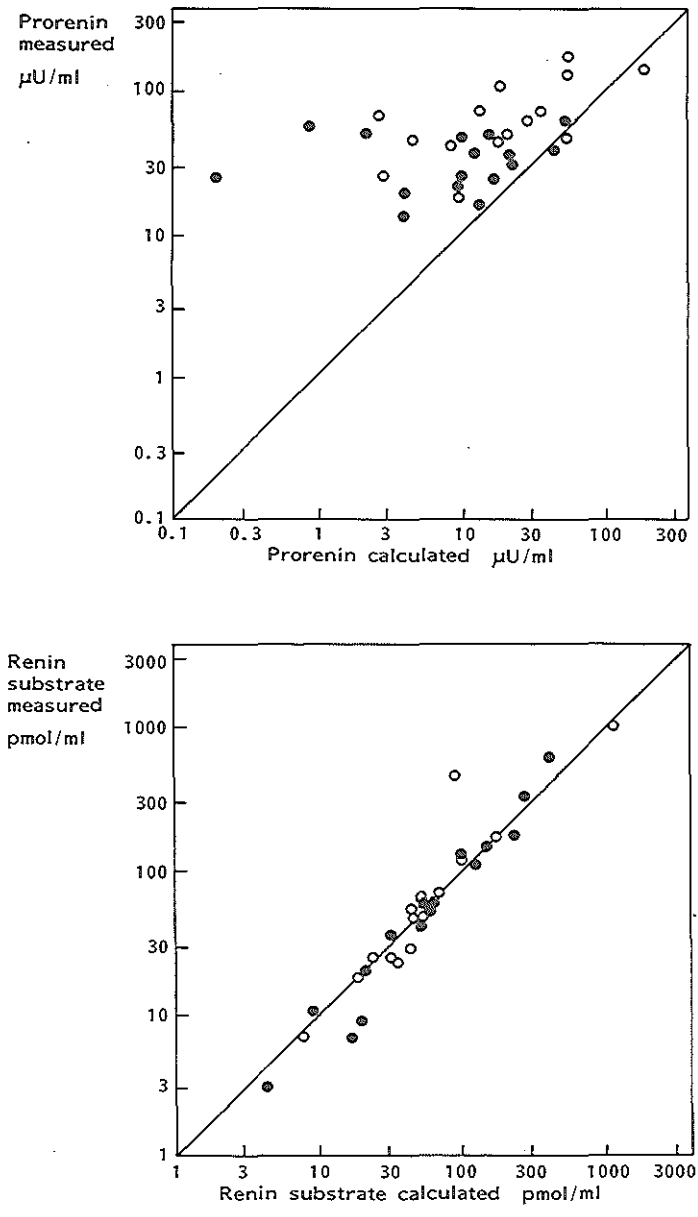


Figure 2. Measured versus calculated concentrations of prorenin (top panel) and renin substrate (bottom panel) in vitreous fluid. For an explanation of the calculation see text (*Data analysis*). The slopes and significance levels of the correlations are given in Table 3. \circ diabetic, \bullet non-diabetic.

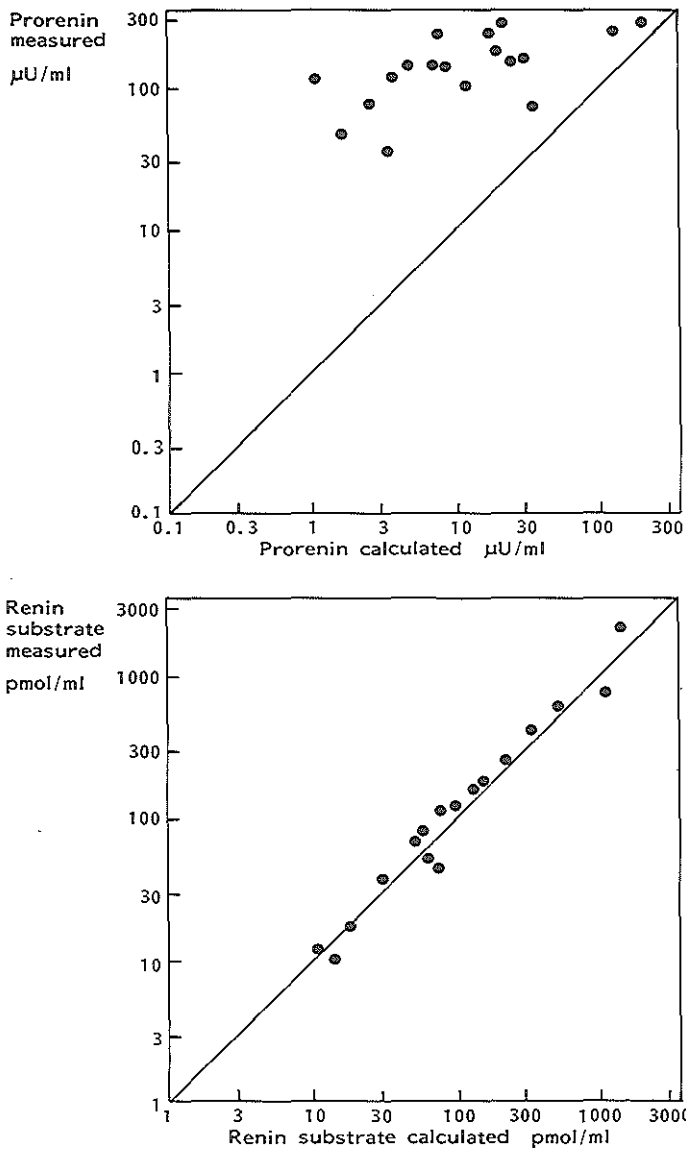


Figure 3. Measured versus calculated concentrations of prorenin (top panel) and renin substrate (bottom panel) in subretinal fluid. For an explanation of the calculation see text (*Data analysis*). The slopes and significance levels of the correlations are given in Table 4.

Table 3. Correlations between the measured and calculated concentrations of proteins in vitreous fluid.

Protein	Regression Line	r	p
Prorenin			
Non-diabetic subjects	$y = 0.015^a x + 1.523$	0.05	NS
Diabetic subjects	$y = 0.328^a x + 1.370$	0.64	< 0.05
Renin substrate	$y = 1.024x - 0.028$	0.97	< 0.0001
IgG	$y = 1.063x - 0.344$	0.91	< 0.0001
Transferrin	$y = 0.899x + 0.066$	0.89	< 0.0001

y is the log (measured concentration); x is the log (calculated concentration). For renin substrate, IgG, and transferrin, data from diabetic and nondiabetic subjects were combined because no differences were found for these proteins between the two groups (Tables 1 and 2).

^a Slope different from 1.0, $p < 0.0001$.

Table 4. Correlations between the measured and calculated concentrations of proteins in subretinal fluid.

Protein	Regression Line	r	p
Prorenin	$y = 0.271^a x + 1.839$	0.61	< 0.01
Renin substrate	$y = 0.975x + 0.064$	0.98	< 0.0001
IgG	$y = 0.941x - 0.302$	0.99	< 0.0001
Transferrin	$y = 0.934x + 0.262$	0.94	< 0.0001

y is the log (measured concentration); x is the log (calculated concentration).

^a Slope different from 1.0, $p < 0.0001$.

Diabetic subjects

The results of the renin substrate, albumin, transferrin, and IgG measurements in vitreous fluid and plasma of the diabetic subjects were similar to those in the non-diabetic subjects (Tables 1 and 2). The levels of renin and prorenin in vitreous fluid were higher in the diabetic than in the non-diabetic subjects. Prorenin was also higher when allowance was made for differences in plasma protein content of the samples. In the diabetic subjects the prorenin concentration of vitreous fluid correlated with the plasma prorenin concentration ($r=0.78$, $n=15$, $p < 0.001$). In the non-diabetic subjects there was no significant correlation between the levels of prorenin in vitreous fluid and plasma.

As in the non-diabetic subjects, prorenin in vitreous samples with low plasma protein content was much higher (up to 25 times) than expected on the basis of

the albumin content of the samples.

Immunoreactive angiotensin II was 9.0 ± 2.5 pmol/L in vitreous fluid (n=15) compared to 9.9 ± 2.9 pmol/L in plasma.

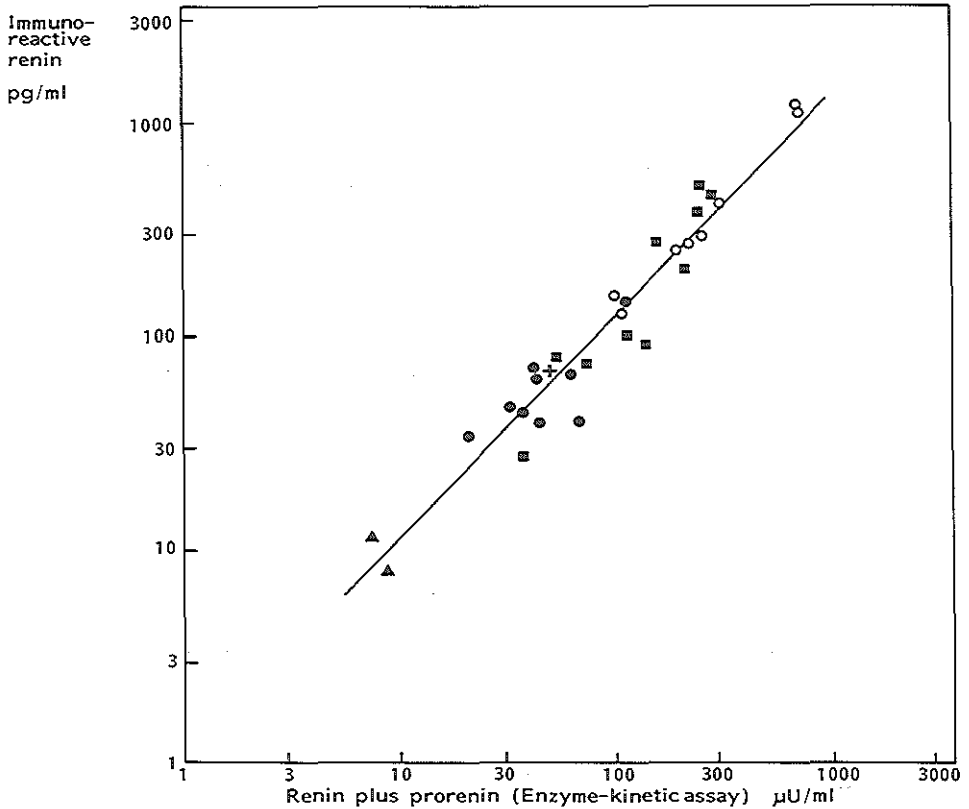


Figure 4. Total renin (prorenin plus renin) measured by enzyme-kinetic assay versus immunoreactive renin ($r=0.97$, $p < 0.0001$). ○ plasma, ■ subretinal fluid, ● vitreous fluid, ▲ aqueous fluid, + WHO human kidney renin standard.

5.5 Discussion

The levels of albumin (mol wt, 69 K), transferrin (mol wt, 90 K), IgG (mol wt, 150 K) and renin substrate (mol wt, 65 K) in vitreous fluid differed widely from sample to sample, but in each individual sample the levels were, relative

to each other, comparable to those in plasma. The IgG level in vitreous fluid, relative to that of albumin, was systematically somewhat lower than in plasma, probably due to its larger molecular size (25). These results are in agreement with earlier findings that most soluble protein in the vitreous is derived from plasma (24-26).

The plasma protein content of normal vitreous fluid has been estimated to be 0.5-2 % of that in plasma (25,26). In our study vitreous fluid from eyes with recurrent retinal detachment due to proliferative vitreoretinopathy contained higher levels of plasma proteins. These higher values probably reflect partial breakdown of the blood-retinal barrier in such eyes (27).

The blood-retinal barrier is formed by the tight junctions between the endothelial cells of the retinal capillaries and the tight junctions between the retinal pigment epithelial cells, the latter restricting the transfer of plasma proteins escaping from the capillaries of the choroid (28). Proteins from plasma may enter the vitreous as a result of focal cellular necrosis, opening of the intercellular junctions, or vesicular transport and formation of transcellular channels. The rate of diffusion of proteins through such discontinuities in the blood-retinal barrier depends upon the concentration gradient across this barrier, the molecular size of the proteins and the number and area of discontinuities. Our results are in accordance with the contention that diffusion through these pores is the main mechanism of transfer of plasma proteins to the vitreous.

This process, however, does not appear to hold true for prorenin (mol wt, 54 K). The concentration of prorenin, relative to that of albumin, was much higher in vitreous fluid than in plasma, and the prorenin level in vitreous fluid also was little influenced by its plasma protein content. Thus, prorenin may enter the vitreous by a mechanism that is different from that of albumin and other plasma proteins. This mechanism is selective for prorenin and may involve an active process. Receptor-mediated transcellular transport is such a selective mechanism, but as yet there is no evidence for the existence of cell-membrane receptors of prorenin. Our findings raise the possibility that not all the prorenin in the vitreous is derived from plasma but that some of it is produced in the eye.

As described under *Data analysis* above, the vitreous level of prorenin that has crossed the blood-retinal barrier by passive diffusion in the same way as albumin can be estimated by multiplying the vitreous/plasma concentration ratio of albumin by the plasma prorenin level. By subtracting this calculated level of plasma-derived prorenin from the level actually measured, we estimated the level of prorenin that entered the vitreous by some process that is different from diffusion out of the circulation. In most samples of vitreous fluid the estimated level of prorenin that had entered the vitreous by such a diffusion-independent process was more than 4 times higher than the estimated level of prorenin that had entered the vitreous by passive diffusion from blood. Thus, relative to the total amount of prorenin in vitreous fluid, the

contribution of plasma-derived prorenin crossing the blood-retinal barrier merely by diffusion appears to be small. The implicit assumption underlying these calculations is that albumin and prorenin leave the vitreous in the same way, that is by free diffusion into the aqueous fluid (29), where the concentrations of these proteins were much lower than in the vitreous.

The concentrations of albumin, transferrin, IgG and renin substrate were 2-3 times higher in subretinal fluid than in vitreous fluid. This was to be expected, since subretinal fluid from eyes with retinal detachment is more or less a concentrate of vitreous fluid (27). Vitreous fluid enters the subretinal space through the hole(s) of the retina, and water is actively absorbed from the subretinal space by the retinal pigment epithelium. Again, the findings for prorenin were different. The concentration of prorenin, relative to albumin, was much higher in subretinal fluid than in plasma, particularly in subretinal samples with low plasma protein concentrations. Moreover, relative to albumin and other plasma proteins, prorenin was 2 times higher in subretinal than in vitreous fluid. If it is assumed, on the basis of the evidence discussed above, that most of the prorenin in the vitreous is not derived from plasma but is produced in the eye, the difference in prorenin content between subretinal and vitreous fluid may suggest that the subretinal compartment is closer to the site of prorenin production.

Not only were the prorenin concentrations of vitreous and subretinal fluid higher than expected, so too were the renin (mol wt, 48 K) concentrations. The data on renin, however, are more difficult to interpret than those on prorenin because in many samples renin was at or below the detection limit of the assay and because some prorenin to renin conversion may have occurred during storage and handling of the samples. Even as little as 1 % conversion will result in a large percentage increase of renin in these samples.

Immunoreactive angiotensin II also was found in samples of vitreous fluid, in concentrations comparable to those in plasma. Further work is needed to answer the question of its origin.

That the levels of albumin and IgG in vitreous fluid from eyes affected by proliferative diabetic retinopathy were higher than the levels in normal eyes can be explained by the increased permeability of the blood-retinal barrier in this condition (30). The higher vitreous level of prorenin, relative to those of albumin and other plasma proteins, in the diabetic subjects as compared to non-diabetic subjects is more difficult to explain. The same arguments in favor of the hypothesis that, generally, diffusion from the blood contributes little to the total amount of prorenin in the vitreous, apply to both diabetic and non-diabetic subjects. It seems, therefore, unlikely that the higher level of prorenin in vitreous fluid of the diabetic subjects (2 times that in non-diabetic subjects) was caused by the higher level in plasma (also 2 times that in non-diabetic subjects). It might be the other way around; increased release or leakage of prorenin from the eye affected by proliferative diabetic retinopathy may contribute to the increased prorenin level in plasma.

This possibility is further supported by the finding that, in contrast with other proteins, the plasma concentration of prorenin in diabetic subjects correlated significantly with the concomitant vitreous prorenin concentration. Considering the fact that in some diabetic subjects the blood-retinal barrier for plasma proteins was still relatively intact (low vitreous/plasma albumin concentration ratio), whereas in others it was extremely leaky, no such correlation was to be expected, if diffusion from plasma into the vitreous was the main mechanism of transfer of prorenin.

An elevated plasma prorenin level in diabetic subjects has been found to be associated with microvascular complications, including retinopathy (31). Evidence is accumulating that neovascularization is initiated by diffusible chemical factors arising from ischemic areas of the retina (32). Renin and angiotensin have been found in cultured neuronal and glial cells from rat brain (33,34); both cell types are abundantly present in the retina. Angiotensin II acts on vascular tone and has mitogenic and trophic actions on vascular smooth muscle and other cells (35). In fact, it has been reported to promote neovascularization (16). An intraocular renin-angiotensin system may, therefore, play a role in proliferative diabetic retinopathy.

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6. IDENTIFICATION AND QUANTIFICATION OF RENIN AND PRORENIN IN THE BOVINE EYE

6.1 Summary

Angiotensin II, the most important biologically active product of the renin-angiotensin system, has been reported to play a role in neovascularization, and prorenin has been found in the vitreous of human eyes, particularly in those affected by proliferative diabetic retinopathy, a disease characterized by neovascularization. The prorenin level in these eyes was, relative to that of plasma albumin, higher than in eyes without neovascularization. These findings suggested that an intraocular renin-angiotensin system exists, which might be involved in the development of retinal neovascularization in diabetes mellitus. In this study angiotensin I-generating activity was measured in bovine aqueous humor and vitreous and in extracts of bovine retina, pigment epithelium-choroid and anterior uveal tract, before and after subjecting these extracts to procedures known to convert prorenin to renin. The measurements were made by incubation at 37 °C with plasma from nephrectomized rats at pH ranging from 5.0-8.5. True renin in the ocular samples could be separated from nonrenin acid protease by α -casein-Sepharose affinity column chromatography at pH 3.5; true renin did not bind to the column, whereas acid protease did. True renin was further identified by its relatively high pH optimum (6.5-7.0) for angiotensin I generation, its complete inhibition with specific renin antiserum, and its high affinity for specific renin inhibitors. More than 75 % of angiotensin I-generating activity of the ocular samples consisted of true renin. Approximately 90 % or more of total renin (renin plus prorenin) in aqueous humor, vitreous, and ocular tissues could not be explained by trapped plasma. Total renin in aqueous humor and renin in vitreous were near the detection limit of the assay of angiotensin I-generating activity. In vitreous prorenin comprised 99 % of total renin, in retina 81 %, and in pigment epithelium-choroid and anterior uveal tract less than 50 %. Prorenin in ocular fluids showed a concentration gradient, posterior vitreous > anterior vitreous > aqueous humor, suggesting that the main source of extracellular prorenin was in the posterior eye. These data support the contention of local renin and/or prorenin synthesis in the eye and are in accordance with the observations in other tissues that extrarenal synthesis of renin is often associated with the release of mainly, or exclusively, prorenin into extracellular fluid.

6.2 Introduction

The aspartyl protease renin is the key enzyme in the formation of angiotensin II, an octapeptide with well known functions in blood pressure regulation and fluid and electrolyte homeostasis. Classically, renin is considered a blood-borne enzyme synthesized and secreted by the kidney together with its inactive precursor, prorenin. Plasma from nephrectomized subjects predictably contains little or no renin but it does contain prorenin, in some cases in concentrations as high as in normal individuals (1). Apparently most, if not all, renin in plasma originates from the kidney, whereas a large proportion of plasma prorenin is produced at extrarenal sites.

Synthesis of renin or prorenin and other components of the renin-angiotensin system (RAS) is known to occur in many organs besides the kidney, for instance adrenal (2,3), pituitary (2,3), testis (2,3), brain (4) and ovary (5-7). In the luteal phase of the menstrual cycle and during pregnancy the ovary, probably the corpus luteum, secretes prorenin into the circulation (5). Certain renal and extrarenal tumors secrete prorenin in large amounts (8), and in some patients carrying these tumors plasma renin is elevated as well. Finally, cultured smooth muscle cells (9), chorionic cells (10) and ovarian follicular theca cells (6) have been shown to secrete prorenin into the medium.

It has been postulated that a correlation exists between the degree of vascularization of tissues and their renin content (11), and there is evidence to suggest that angiotensin II has a role in angiogenesis (12). Because neovascularization is a hallmark of progressed diabetic retinopathy, we wondered whether renin might also be synthesized in the eye. Recently we described the presence of prorenin in the human eye (13). Levels of prorenin in vitreous and subretinal fluid obtained during eye surgery were, relative to albumin and other plasma proteins, up to 100 times higher than in plasma. These results lend support to the hypothesis that synthesis of prorenin may occur in the eye. Presumably synthesis takes place in one or more of the surrounding tissues of the vitreous, since vitreous itself is virtually acellular. In the present study we attempted to identify and to measure renin and prorenin in extracts of different parts of freshly obtained bovine eyes and in simultaneously taken plasma.

6.3 Materials and methods

Buffers and reagents

Bovine serum albumin (BSA), 8-hydroxyquinoline, bovine α -casein, bovine cathepsin D and bovine hemoglobin were obtained from Sigma (St. Louis, MO). Trichloroacetic acid (TCA), phenylmethylsulfonylfluoride (PMSF) and

sodium azide (NaN_3) were purchased from Merck (Darmstadt, Germany). EDTA, glycine and polyvinylpyrrolidone (PVP) were purchased from Riedel de Haën (Scelze, Germany), trypsin from Serva (Heidelberg, Germany), N-ethylmaleimide (NEM) from Aldrich (Milwaukee, WI, USA), sodium tetrathionate from Fluka (Buchs, Switzerland), plasmin from Kabi Vitrum (Stockholm, Sweden) and CNBr-activated Sepharose 4B from Pharmacia (Uppsala, Sweden). Aprotinin was purchased from Bayer (Leverkusen, Germany). Antiserum against BSA was obtained from Dakopatts (Glostrup, Denmark). Rabbit antiserum against bovine pituitary renin was a gift by Dr. Murakami, University of Tsukuba, Japan (14). The renin inhibitor Ro 42,5892 was a gift from Dr. van Brummelen (Hoffmann-La Roche, Basel, Switzerland). The statin-containing renin inhibitor CGP 29,287 was a gift from Dr. Hofbauer (Ciba-Geigy, Basel, Switzerland). The World Health Organization (WHO) human kidney renin standard 68/356 was obtained from the WHO International Laboratory for Biological Standards (London, United Kingdom). For studying the optimal conditions for *in vitro* activation of prorenin the following buffers were used: buffer A, pH 3.3, consisting of 0.05 M glycine, 0.001 M EDTA and 0.095 M NaCl, and buffer B, pH 7.4, consisting of 0.1 M phosphate, 0.001 M EDTA and 0.075 M NaCl. Buffer C, pH 7.4 was used for washing tissues and for homogenization; it consisted of 0.01 M phosphate and 0.15 M NaCl.

Collection of blood and ocular tissues

Blood was obtained from cows at the local slaughterhouse immediately after death. Blood was collected in tubes containing sodium citrate (final concentration, 0.013 M). At the same time both eyes were enucleated. Eyes and blood were kept on ice and were processed within 60 min. Blood was centrifuged at 4 °C for 10 min at 3000 g, and plasma was frozen at -20 °C. Eyes were dissected as described below. Dissection and extraction procedures were performed at 4 °C.

Aqueous humor was drawn using a tuberculin syringe with a fine needle. The eye was cut equatorially at the ora serrata and the anterior segment was lifted off. The vitreous body was isolated by gently shaking it out of the eye cup. Care was taken to remove all vitreous. The neural retina was cautiously teased away from the pigment epithelium with a thin glass rod and isolated by cutting it at the optic nerve. The choroid with adhering pigment epithelium layer was isolated by dissecting it from the sclera with a pair of fine scissors. The anterior uveal tract, consisting of iris and ciliary body, was isolated by removing the lens from the anterior eye cup, then gently pulling the anterior uveal tract loose from the sclera and blotting it on dry paper to remove any adhering vitreous. Cornea, lens and sclera were discarded.

For the chemical and immunological identification of renin and prorenin in the ocular tissue extracts and in plasma, either the retina, the pigment epi-

thelium-choroid or the anterior uveal tract of 60 bovine eyes, the vitreous bodies of 4 eyes or 70 mL of bovine plasma were pooled. Tissues were minced with scissors into small pieces and washed three times in two volume weights of ice-cold buffer C in order to remove as much plasma as possible. Tissues were homogenized in one volume weight of buffer C with a Polytron PT10/35 (Kinematica, Luzern, Switzerland) with 2 15-sec bursts at level 10. The Polytron was also used to break the gel structure of the vitreous. Homogenates were rapidly frozen and thawed three times and then centrifuged at 4 °C for 1 h at 13000 g. Supernatants were decanted and stored at -20 °C. Pellets were discarded. Preliminary experiments had shown that there was no difference in renin content in extracts that had been made in buffer C alone and those made in buffer C containing the protease inhibitors PMSF (0.0005 M), EDTA (0.001 M) and sodium tetrathionate (0.00025 M; final concentrations) (15). Apparently, inadvertent activation of prorenin during the extraction procedures did not take place in these extracts.

The extracts of retina, pigment epithelium-choroid and anterior uveal tract were divided into two portions. One portion of each extract was dialyzed against buffer A to pH 3.3 for 48 h and subsequently against buffer B to pH 7.4 for 48 h, both at 4 °C, after which denatured protein was removed by centrifugation. The resulting supernatants were dialyzed for 6 h in distilled water, lyophilized and stored at -20 °C. The acidification step removed effectively much of the angiotensinase activity, which in preliminary experiments was found to interfere with the renin assay in non-acidified lyophilized extracts despite the use of angiotensinase inhibitors. These lyophilisates were used for the experiments with renin antiserum or the renin inhibitor Ro 42,5892 or for α -casein-Sepharose affinity chromatography, as described below. The other portion of each homogenate was dialyzed against buffer B, pH 7.4, only. The dialyzed homogenates were centrifuged and used to establish the optimal method for activating prorenin.

For quantification of renin and prorenin in eyes and simultaneously taken plasma, eyes from eight cows were studied. Isolated parts of one eye were pooled with the corresponding parts of the other eye of the same animal, aqueous and vitreous in equal volumes. No washing procedure to remove plasma was included so as not to lose tissue renin. Tissues were weighed and one volume weight of buffer C was added. Homogenization was performed as described above.

To examine whether a concentration gradient exists for prorenin in the vitreous, the vitreous bodies of five eyes were rapidly frozen in liquid nitrogen immediately after enucleation and then divided into posterior, central and anterior parts after the surrounding tissues had been peeled off. Aqueous of these eyes was collected just before freezing.

Bovine kidney was homogenized in one volume weight of buffer C and submitted to pH 3.3 dialysis against buffer A for 48 h; pH was restored to pH 7.4 by dialysis against buffer B for 48 h. Denatured protein was removed by

centrifugation. The resulting supernatant contained renin at a concentration of about 20.000 ng angiotensin I/mL · h, as assessed by incubation at pH 7.4 with plasma from nephrectomized rats as a source of renin substrate. In most experiments this extract was diluted 300-fold in 0.1 M phosphate buffer, pH 7.4. No acid protease activity was detectable at this dilution.

Identification of renin

Affinity chromatography on α -casein-Sepharose. Aspartyl proteases bind to α -casein at low pH. Renin, however, is an exception because of its relatively high pH optimum. This property of renin can be used for separating renin from other aspartyl proteases by affinity chromatography on α -casein-Sepharose (16). α -casein was coupled to CNBr-activated Sepharose 4B (16), and a column of 10x1 cm was prepared. The column was equilibrated with 0.05 M HAc-Ac⁻ buffer, pH 3.5, containing 0.075 M NaCl. A lyophilized extract of pigment epithelium-choroid was dissolved in this buffer and 1 mL of this extract was applied to the column and left to stand for 5 h. Elution was started with adsorption buffer at a flow rate of approximately 10 mL/h. After a washout volume of 90 mL, a 0.1 M Tris-HAc buffer, pH 8.6, containing 1.0 M NaCl, was applied and elution was continued at a flow rate of 10 mL/h. Fractions of 3 mL were collected. All procedures were carried out at 4 °C. All fractions were assayed for angiotensin I-generating activity at pH 7.4 and for acid protease activity at pH 3.5.

pH optimum study. Plasma from nephrectomized rats, which served as a source of substrate for renin and activated prorenin, and the samples to be tested for angiotensin I-generating activity were brought to appropriate pH by overnight dialysis in separate dialysis bags at 4 °C in 0.1 M phosphate buffers ranging from pH 5.0-8.5. Angiotensin I-generating activity was assessed at these pH values; incubation conditions were the same as in the renin assay described below, except for pH. The pH of the incubation mixture did not change more than 0.05 pH unit during incubation. The following samples were tested: 1) pool of bovine plasma, which was treated with trypsin-Sepharose for activating prorenin (see below); 2) dilute bovine kidney extract; 3) bovine vitreous fluid, which was treated with acid and plasmin for activating prorenin (see below); 4) renin from bovine pigment epithelium-choroid purified by α -casein-Sepharose chromatography (see below); and 5) purified bovine cathepsin D.

Inhibition of renin by renin antiserum or renin inhibitors. The rabbit antiserum against bovine pituitary renin at 1:320 dilution caused 50 % inhibition of the angiotensin I-generating activity at pH 7.4 of a bovine kidney extract with a renin concentration of 70 ng angiotensin I/mL · h.

A 1:40 dilution of antiserum was made in 0.1 M Tris-HCl, pH 7.4,

containing 0.1% BSA and 0.02 % NaN_3 . Equal volumes of test sample and dilute antiserum were mixed and left to stand for 20 h at 4 °C. The following samples were tested: 1) bovine plasma, after treatment with trypsin-Sepharose for activating prorenin (see below); 2) vitreous fluid, which was treated with acid and plasmin for activating prorenin (see below); 3) lyophilized extracts of retina, pigment epithelium-choroid and anterior uveal tract diluted in distilled water; 4) bovine kidney extract; and 5) renin from pigment epithelium-choroid, purified by α -casein-Sepharose affinity chromatography (see below).

To assess the specificity of the antiserum, inhibition experiments were also conducted with bovine cathepsin D. Renin assays that followed preincubation with the antiserum were performed not only at pH 7.4, but also at pH 5.5 because cathepsin D has little angiotensin I-generating activity at neutral pH, whereas renin does have angiotensin I-generating activity at neutral pH as well as at pH 5.5. Controls were samples incubated with preimmune rabbit serum. Results were corrected for the presence of renin in the antiserum and preimmune serum (approximately 3 ng angiotensin I/mL · h).

Inhibition of angiotensin I-generating activity by two renin inhibitors was assessed by performing the renin assay in the presence of different inhibitor concentrations, ranging from 10^{-11} to 10^{-5} mol per liter (final concentration). One inhibitor was Ro 42,5892 from Hoffmann-La Roche, a substrate analogue, which is claimed to be enzyme specific for renin. The concentration at which it causes 50 % inhibition (IC_{50}) of human renin is 0.5×10^{-9} M in the plasma renin activity assay, in which plasma renin is allowed to react with endogenous renin substrate at pH 7.4. The IC_{50} of this inhibitor for bovine cathepsin D is 3.7×10^{-5} M, when assessed by proteolysis of denatured hemoglobin at pH 3.5 (information provided by the manufacturer). The other renin inhibitor was CGP 29,287 from Ciba-Geigy, also a substrate analogue, which has been described previously (17). CGP 29,287 is a potent inhibitor of human renin but appears to be less effective against dog or rat plasma renin, with an IC_{50} of 1×10^{-9} , 2×10^{-7} and 3×10^{-5} M respectively, in the plasma renin activity renin assay (17). Its IC_{50} for bovine cathepsin D is 4×10^{-5} M, again assessed by proteolysis of denatured hemoglobin at pH 3.5. The samples tested with the renin inhibitors were the same as those used for the antiserum inhibition experiments, except for renin purified from ocular tissues by α -casein-Sepharose chromatography, which was only used in the antiserum experiments.

To assess the specificity of the inhibitors for bovine renin, the effect of the inhibitors on angiotensin I-generating activity of bovine kidney renin was compared with their effect on the angiotensin I-generating activity of bovine cathepsin D. As in the antiserum inhibition experiments, this was done at pH 5.5. The effects of the inhibitors on the angiotensin I-generating activity of human kidney renin in our renin assay were also tested and this was done both at pH 5.5 and 7.4.

Assays of renin and activated prorenin. Different procedures were attempted for activating prorenin in bovine plasma, vitreous and extracts of retina, pigment epithelium-choroid and anterior uveal tract. We used three methods that were proven to be successful in activating human prorenin. One can either make use of endogenous proteolytic activators of prorenin, such as factor XII/kallikrein, plasmin, or pepsin-like proteases, or one can add activators, for instance plasmin or trypsin. In two of our methods we used an acidification step. Acidification has several effects: 1) prorenin undergoes a reversible conformational change of the molecule by which it becomes more susceptible to limited proteolysis necessary for irreversible activation (18); 2) acidification destroys inhibitors of proteases that activate prorenin at neutral pH, like kallikrein or plasmin (18); and 3) pepsin-like acid proteases, capable of activating prorenin, may become active at low pH, like for example in human amniotic fluid (18). The first method of activation we tested in this study consisted of acidification followed by neutralization; no exogenous activator was added. Two mL of sample were dialyzed in buffer A for 48 h at 4 °C. Then the dialysis buffer was changed to buffer B, which after 36 h was replaced by buffer B containing 6 % PVP to prevent dilution due to colloid-osmosis. Dialysis in the presence of PVP proceeded for another 12 h after which the content of dialysis bags was collected by rinsing the interior with buffer B. Volumes were adjusted to 2 mL with the same buffer. The second method involved incubation at 4 °C with immobilized trypsin as previously described (19). Trypsin had been coupled to CNBr-activated Sepharose 4B. The final concentration of trypsin was 4.4×10^3 Na-benzoyl-L-arginine ethyl ester units/500 μ L sample. Activation of prorenin in plasma with this method does not require prior acidification. After 48 h incubation trypsin was removed by centrifugation. This method is similar to the often used trypsin-soybean trypsin inhibitor combination (15), with the advantage that destructive proteolysis of prorenin and renin is less likely to occur. The third method was an acidification step followed by incubation with plasmin at pH 7.4. After plasmin inhibitors have been removed at low pH, exogenous plasmin will cause activation of prorenin at neutral pH without destruction of prorenin or renin, a problem which is sometimes encountered with trypsin and acid proteases. After dialysis for 48 h at 4 °C in buffer A, samples were rapidly adjusted to pH 7.4 with 1 M NaOH. Then 0.1 volume of plasmin solution in 0.15 M NaCl was added to a final concentration of 1 casein unit per mL. Incubation with plasmin was for 48 h at 4 °C. With each sample all three procedures were performed in duplicate.

Renin and activated prorenin were assayed in duplicate by measuring the rate of angiotensin I generation at pH 7.4 and 37 °C in the presence of the protease inhibitors PMSF (0.0024 M), EDTA (0.005 M), 8-hydroxyquinoline (0.0034 M), aprotinin (2.2×10^{-9} M) and N-ethylmaleimide (0.005 M; final concentrations) (15,19). The incubation mixture consisted of one volume of 0.1 M phosphate buffer, one volume sample, two volumes of renin substrate

and 0.14 volume of inhibitor solution. Plasma from nephrectomized rats was the source of renin substrate. The renin activity of this plasma was less than 0.1 ng angiotensin I/mL · h. The final substrate concentration in the incubate was 2.9 µg angiotensin I equivalents/mL, which was about four times the K_m . The mixtures were incubated for 1, 2, 3 and, in some cases, 6 h at 37 °C. Angiotensin I was measured by radioimmunoassay (19). Only incubations in which angiotensin I generation was linear with time were used for renin assay. Recovery of angiotensin I added to the incubation mixture was 98 % after 6 h incubation, demonstrating effective inhibition of angiotensinase activity.

Assays of serum albumin and acid protease. Albumin, as a measure of trapped plasma in the ocular tissue extracts, was determined with a single radial immunodiffusion technique, as described by Mancini et al. (20). One mL of the BSA antiserum precipitated 900 mg of BSA. Dilutions of samples were made in 0.07 M barbital, 0.003 M calcium lactate and 0.3 % Tween 20, pH 8.6. Standard BSA solutions ranged from 5 to 100 mg/L. Diffusion into the agar gel plates was allowed to proceed for two days in humidified air at room temperature. The agar plates were then stained with Coomassie blue. Recovery of added albumin ranged from 90-100 % in the ocular tissue extracts.

The acid protease assay was a modification of the method by Anson (21). Briefly, 200 µL sample, which had been brought to pH 3.5 with 1 M HCl, was added to 1 mL hemoglobin (1 %) in 0.1 M glycine/HAc, pH 3.5, at 37 °C. At $t=0$ and $t=20$ min 500 µL of this mixture was mixed with 500 µL ice-cold 10 % TCA. Denatured protein was removed by centrifugation and the absorption at 280 nm of the supernatant was read in an Uvikon 810 spectrophotometer (Kontron, Zurich, Switzerland). The difference in absorption at 280 nm between 20 and 0 min at 37 °C was taken as a measure of acid protease activity. Acid protease activity is expressed in units; one unit is the proteolytic activity that causes a change in extinction of 1000/min at 280 nm by TCA-soluble products at pH 3.5 and a temperature of 37 °C, with denatured hemoglobin as a substrate.

6.4 Results

Affinity chromatography

The protein concentration (absorption at 280 nm), the acid protease activity at pH 3.5 and the angiotensin I-generating activity at pH 7.4 of the eluted fractions from the α -casein-Sepharose column are depicted in Fig. 1. Renin-like activity was eluted later than the bulk of protein. Most angiotensin I-generating activity was found in a broad peak eluted at pH 3.5 (peak I).

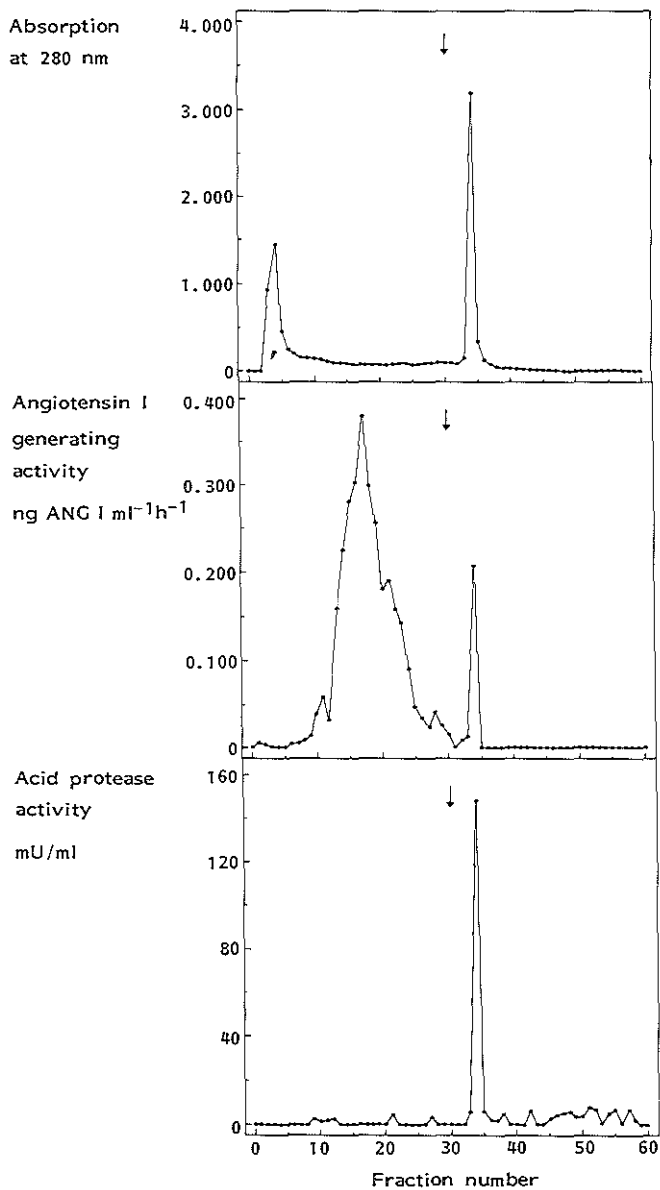


Figure 1. Separation of renin from non-renin acid protease in extract of bovine pigment epithelium-choroid by affinity chromatography on α -casein-Sepharose. Arrow indicates change of 0.05 M HAc-Ac⁻ buffer, containing 0.05 M NaCl, pH 3.5, to 0.1 M Tris-HAc buffer, containing 1 M NaCl, pH 8.6.

A small amount of angiotensin I-generating activity (peak II) was eluted after changing the pH to 8.6. Peak I contained renin activity but no acid protease activity. Peak II contained acid protease activity. Fractions 13-24 (peak I) were dialyzed against distilled water for 6 h and lyophilized. This preparation was used for determining the pH optimum of angiotensin I generation and for inhibition with renin antiserum. Peak II was not further evaluated because of its small volume and low renin-like activity.

Identification of renin

For preparing the extracts that were used for identification studies the tissues were washed to remove trapped plasma. Measurement of albumin in these extracts showed that washing with buffer C had removed 70-85 % of plasma. More than 90 % of the angiotensin I-generating activity of these extracts could not be accounted for by contamination with plasma or by simple diffusion of renin from plasma.

Angiotensin I
generating activity
(% of maximum)

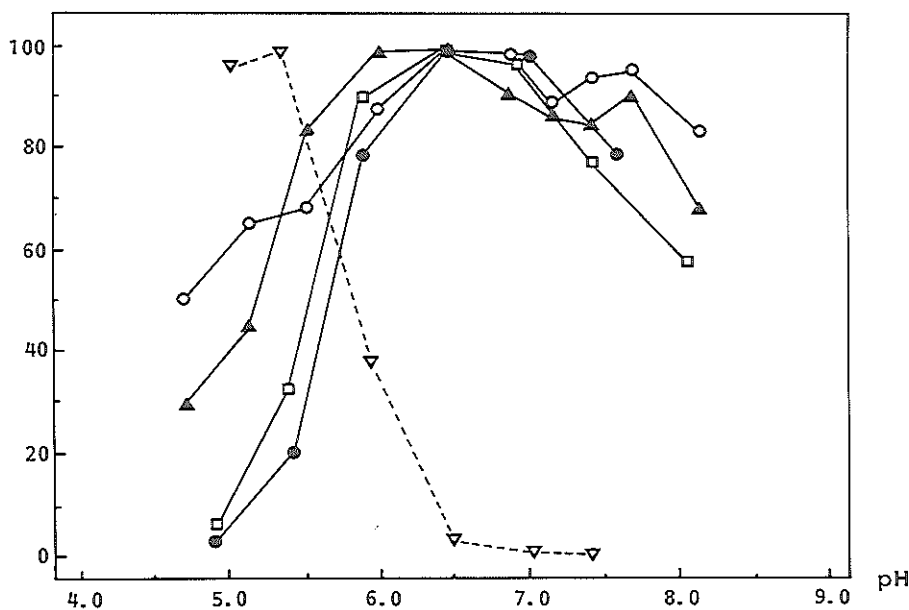


Figure 2. pH optimum curves for angiotensin I-generating activity of purified renin from bovine pigment epithelium-choroid (□), activated prorenin from bovine vitreous (▲), activated prorenin from bovine plasma (○), bovine kidney extract (●) and bovine cathepsin D (▽).

Table 1. Angiotensin I-generating activity of various bovine specimens and its inhibition by renin antiserum.

Sample	ANG I generation (ng ANG I/mL · h)		% Inhibition	
	pH7.4	pH5.5	pH7.4	pH5.5
Plasma	5.8		71	
Vitreous	6.4		70	
Retina	23.4		83	
Pigment epithelium-choroid	40.4		83	
Anterior uveal tract	76.7		93	
Kidney	70.1	20.1	97	94
Purified ocular tissue renin	10.6	2.9	83	95
Cathepsin D	ND	3.8		3

Plasma was treated with trypsin-Sepharose to activate prorenin. Vitreous was treated with plasmin after an acidification step in order to activate prorenin. Extracts of pools of bovine retina, pigment epithelium-choroid and anterior uveal tract were dialyzed to pH 3.3 for 48 h and then to pH 7.4 for 48 h, before being lyophilized. Bovine kidney extract was treated in a similar manner as the eye tissue extracts, but was not lyophilized. Purified ocular tissue renin was from pigment epithelium-choroid (peak I of α -casein column). For details on preparation of samples, see text. Renin assays were performed at pH 7.4 or pH 5.5 with and without preincubation with antiserum at pH 7.4. Experiments were carried out in duplicate. ND, not detectable.

Table 2. Renin in bovine plasma and various ocular tissue extracts after different procedures for activating prorenin.

Sample	Renin activity (ng ANG I/mL · h)			
	No activation	pH3.3->pH7.4	Trypsin-Sepharose	pH3.3->pH7.4 + plasmin
Plasma	1.6	3.0	5.9	4.1
Vitreous	0.04	4.7	4.8	6.1
Retina	0.3	1.6	1.2	2.1
Pigment epithelium-choroid	3.0	3.1	3.2	4.4
Anterior uveal tract	1.5	2.9	2.2	3.0

For details on the activation procedures of prorenin, see text. Experiments were carried out in duplicate.

The pH optimum curves of angiotensin I-generating activity for kidney extract, activated prorenin of vitreous and plasma, and purified pigment epithelium-choroid renin (peak I of α -casein-Sepharose column) are shown in Fig. 2. During incubation pH did not shift more than 0.05 pH-unit. Results are expressed as a percentage of maximal angiotensin I-generating activity. Curves were similar, with a pH optimum of 6.5-7.0. Activated prorenin in plasma and in vitreous appears to have a shoulder in the curve at pH 8.0. pH optimum curves were clearly different from the curve of cathepsin D angiotensin I-generating activity.

Angiotensin I-generating activity in kidney, retina, pigment epithelium-choroid and anterior uveal tract extracts and in vitreous as well as purified pigment epithelium-choroid renin was inhibited by 97 %, 83 %, 83 %, 93 %, 70 % and 83 %, respectively, by the antiserum against bovine pituitary renin (Table 1). No inhibition was observed when samples were incubated with preimmune rabbit serum. Cathepsin D was not inhibited by the antiserum at pH 5.5, whereas purified pigment epithelium-choroid renin was. This demonstrates that the inhibitory activity of the antiserum was retained at pH 5.5 and that the antiserum was specific for bovine renin.

For the sake of clarity only the inhibition curves for kidney renin, anterior uveal tract extract and cathepsin D for each of the two renin inhibitors at pH 5.5 are depicted in Fig. 3. Samples of the other ocular tissues and vitreous showed identical curves. The IC_{50} of Ro 42,5892 for bovine renin was approximately 8.3×10^{-9} M, whereas for bovine cathepsin D the IC_{50} appeared to be well over 10^{-5} M. There was no difference in inhibitory activity of Ro 42,5892 on human kidney renin at pH 5.5 nor at pH 7.4, proving that in this range pH did not influence the inhibitory activity of Ro 42,5892. CGP 29,287 was clearly less potent than Ro 42,5892, with an IC_{50} of 2.6×10^{-7} M, which is still much lower than the IC_{50} for bovine cathepsin D (4×10^{-5} M). Apparently, CGP 29,287 is more species-specific than Ro 42,5892.

Activation of prorenin

Acidification followed by rapid neutralization and subsequent plasmin treatment resulted in the highest renin activity in the ocular tissues and in vitreous (Table 2). Not so, however, in plasma, in which treatment with trypsin-Sepharose was the more effective to activate prorenin. Activation with trypsin-Sepharose at pH 7.4 (without prior acid treatment) and activation with plasmin at pH 7.4 (after acid-treatment) reached a plateau after 48 h at 4 °C in all samples tested (results not shown). Hence for quantification of total renin, i.e. renin plus prorenin, in the various compartments of the eye, we chose the acidification-plasmin method for the eye tissue extracts and for plasma we used the trypsin-Sepharose method. From these maximal activation values a

percentage of prorenin can be calculated. In vitreous prorenin comprised 99 % of total renin, in retina 81 %, whereas in pigment epithelium-choroid and anterior uveal tract less than 50 % of total renin was prorenin.

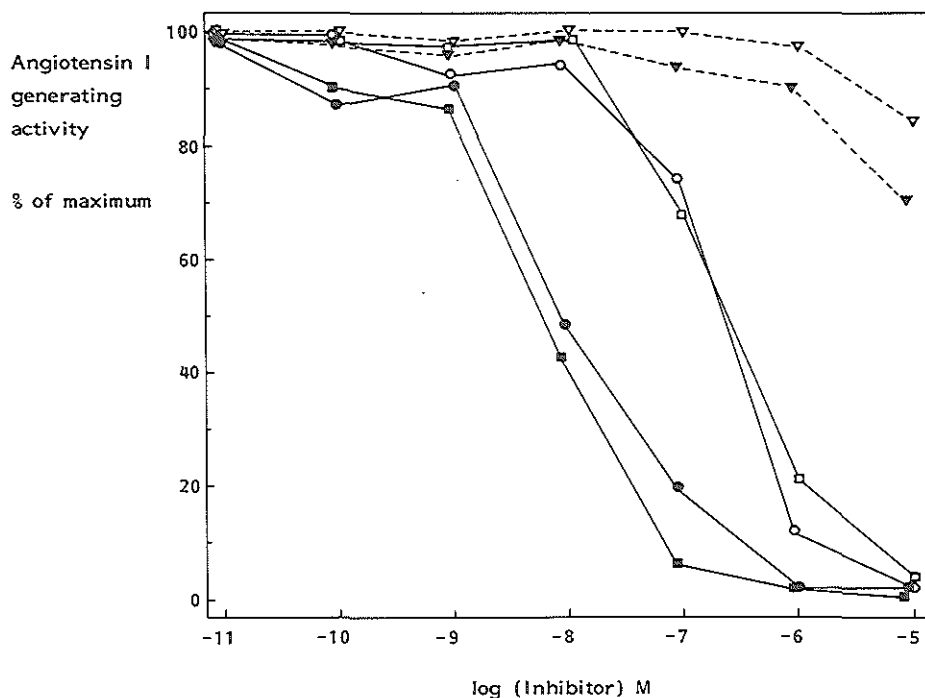


Figure 3. Inhibition of angiotensin I-generating activity of extract of bovine anterior uveal tract (■ and □), bovine kidney extract (● and ○) and bovine cathepsin D (▼ and ▽) by the renin inhibitors Ro 42,5892 (closed symbols) and CGP 29,287 (open symbols).

Renin and prorenin levels in different parts of the eye

Table 3 shows the total renin (renin plus prorenin) levels of plasma, aqueous humor, vitreous, retina, pigment epithelium-choroid and anterior uveal tract of eight cows. In none of the eye samples tested, renin could be accounted for by trapped plasma or by simple diffusion from plasma, since the renin/albumin ratio was much higher in the eye compartments than in plasma. In pigment epithelium-choroid 10 % of renin could be explained by contamination with plasma. In retina and anterior uveal this was only 1 % and 4 % respectively.

Total renin levels, expressed as enzymatic activity per mL of fluid, were approximately equal in vitreous and plasma. In pigment epithelium-choroid and anterior uvea the total renin concentration per gram tissue surpassed even that in plasma.

From Fig. 4 it can be seen that a concentration gradient exists for prorenin in the vitreous, with levels being highest in the posterior vitreous. There was a large difference in prorenin between anterior vitreous and aqueous, which was not apparent for total protein.

Table 3. Total renin (renin plus prorenin) and serum albumin in bovine plasma and different parts of the bovine eye.

Sample	Total renin (ng ANG I/g tissue · h)	Total renin per albumin (ng ANG I/mg albumin · h)	Total renin attributable to plasma (%)
Plasma	6.3 ± 0.5	0.21 ± 0.02	100
Aqueous	0.3 ± 0.04	2.36 ± 0.39	11±2
Vitreous	5.7 ± 0.4	26.8 ± 3.6	1±0.2
Retina	5.1 ± 0.7	16.3 ± 2.0	1±0.2
Pigment epithelium-choroid	14.5 ± 0.8	2.26 ± 0.19	10±1
Anterior uveal tract	10.1 ± 0.7	6.18 ± 0.50	4±0.3

The percentage of total renin attributable to contamination with or diffusion from plasma was calculated as follows: $(\text{alb}_{\text{tissue}}/\text{alb}_{\text{plasma}}) \times (\text{renin}_{\text{plasma}}/\text{renin}_{\text{tissue}}) \times 100$ %. Results were obtained from 8 cows and are means ± SEM.

6.5 Discussion

A critical problem in studying renin in tissue extracts is the distinction between renin and other aspartyl proteases that are present in the tissues. These pepsin-like acid proteases act optimally at a pH lower than the pH optimum of true renin. Nevertheless, they can influence the renin assay in which renin-like activity is detected by angiotensin I generation from renin substrate. The presence of acid proteases other than renin can cause an overestimation of renin content, even when the reaction is performed at pH 7.4 (22). That the renin-like activity we detected at pH 7.4 in ocular tissues was due to true renin is demonstrated by the following findings. We separated ocular tissue renin from acid protease by α -casein-Sepharose affinity chromatography. The peak that contained renin activity but not acid protease activity, showed a pH optimum of 6.5-7, which was similar to that of activated prorenin from vitreous and also similar to the pH optimum of kidney or plasma renin and very different from the pH optimum of purified cathepsin D, which is the most

important of the angiotensin I-generating acid aspartyl proteases in tissues. This almost neutral pH optimum already indicates that we are dealing here with true renin, since renin is the sole angiotensin I-generating aspartyl protease known to have a near-neutral pH optimum. Definitive proof that the renin-like activity in the eye extracts is true renin, was obtained by inhibition of enzymatic activity with a specific antiserum and with two experimental renin inhibitors, Ro 42,5892 and CGP 29,287, whereas purified bovine cathepsin D was inhibited by neither antiserum nor the renin inhibitors.

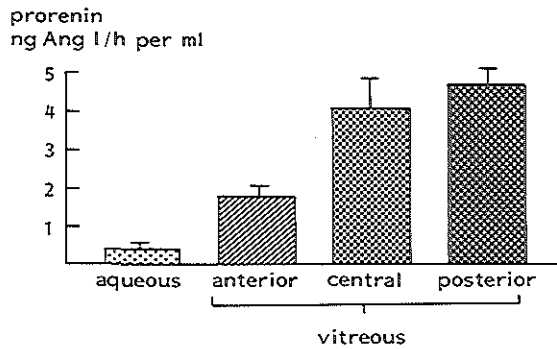
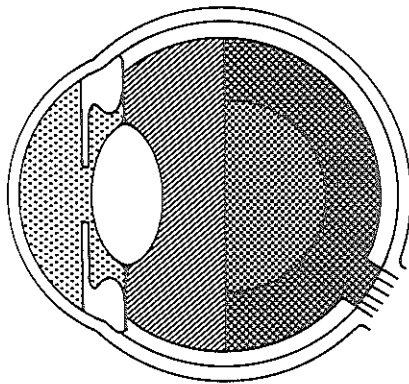


Figure 4. Prorenin in aqueous and in anterior, central and posterior vitreous of the bovine eye.

Another problem in the present study is that the eye is a highly vascularized organ, so that ocular tissue extracts are heavily contaminated with plasma. Renin-like activity in tissue extracts may, therefore, be due to kidney-derived renin present in plasma. Obviously, we could not perform nephrectomy to eliminate this source of renin, so we had to correct for plasma renin in the tissues. We did not wash the tissues for quantification studies so as not to lose any tissue renin. Determination of albumin content of the ocular specimens and of simultaneously taken plasma showed that plasma renin was of no or minor importance, since the renin/albumin ratio in eye structures was much higher than in plasma. In the experiments aimed at the identification of renin in the ocular tissues we first washed the tissues gently in buffer, and in this way we were able to remove some 80 % of the plasma, as estimated by measurement the albumin concentration. Consequently some 80 % of plasma renin was removed. The remainder of plasma renin in the resulting extracts was negligible relative to tissue renin, i.e. extravascular renin, since in the extracts that had not been washed, tissue renin already exceeded plasma renin by far.

Little is known about activation of bovine prorenin. Activation is possible with trypsin (23) but in tissues this has never been proven to be the optimal method. Moreover, we were dealing with two kinds of specimens, namely plasma and tissue extracts, which were very different in composition and may require different procedures for activating prorenin. We examined different methods in order to establish optimal activation conditions. Together they cover the activation mechanisms that are known for human prorenin. From our results it appears that in bovine materials endogenous activators play a less important role in *in vitro* activation of prorenin than in human plasma. In bovine plasma it was not possible to activate prorenin maximally by acidification followed by a neutral phase. In human plasma this occurs via the factor XII/kallikrein pathway (24). In tissue extracts acidification is supposed to activate acid proteases that, in turn, will activate prorenin (18). However in the bovine ocular tissue extracts acidification alone did not yield optimal results. These were obtained when plasmin was added during the neutral phase after the acidification step. Plasmin is known to be an effective activator also of human prorenin. Acid-pretreatment was a prerequisite for the activation of prorenin by plasmin; plasmin alone did not give activation in the ocular tissue extracts. In vitreous, however, acidification was not necessary (results not shown), probably because of the low content of plasmin inhibitors.

Some remarks can be made on the observed renin/prorenin ratio's. Little activation of prorenin seems to occur during the preparation of the ocular tissue extracts; the addition of inhibitors acting on different groups of proteases during preparation of these extracts had no effect on the renin/prorenin ratio. Our finding of bovine vitreous containing almost exclusively prorenin is consistent with our measurements in human vitreous, in that also human vitreous contains much more prorenin than renin (13). Furthermore the

renin/prorenin ratio in the various parts of the eye differs from that in plasma. This difference suggests that a separate regulation mechanism exists for renin in plasma and in the eye and, possibly, that renin in the eye is produced locally. A striking analogy exists with the situation in the bovine ovary (23). In follicular fluid the prorenin/renin ratio is greater than 30, as in vitreous, whereas in the surrounding theca cells this ratio is approximately 1, as compared to 0.7-4.0 in retina and uveal tract.

From our results it cannot be said whether renin in the eye is located intra- or extracellularly, nor is it possible to conclude whether renin is synthesized locally in the eye or sequestered from plasma. Sequestration is not very likely, since that would imply a specific uptake process. Until now cell receptors for plasma renin or prorenin have not been described. The distribution of renin and prorenin at various sites in the eye suggests local synthesis. Vitreous, being the large extracellular space in the eye, has a high percentage of prorenin, which is in agreement with findings of others that extrarenal renin synthesis is accompanied by secretion of exclusively prorenin into the extracellular fluid (6,7,8,15). The higher concentration of prorenin in the posterior segment of the vitreous suggests that the origin of vitreous prorenin is located either in the posterior retina or in the posterior uveal tract. A comment must be made on this observation. It has been shown that rapidly freezing and thawing of a balloon filled with a homogenous solution of albumin can cause uneven displacement of solute (25). The concentration differences amounted to no more than 15 %, with the outer onionskin-like fractions having the highest concentration. If a similar displacement of prorenin had occurred in our experiment, then the concentration in the anterior vitreous would be expected to be equal to that in the posterior, which clearly was not the case. Although our results, thus, may have been influenced to some extent by the freezing procedure, we think that our finding of a prorenin gradient is real and that indeed prorenin originates from the posterior pole of the eye.

One problem with the uveal tract as a source of vitreous prorenin is that, in order to explain the high levels in vitreous, prorenin has to be transferred across the retinal pigment epithelium, which, by virtue of its tight junctions, is part of the blood-retinal barrier. This barrier is not very permeable to proteins. Still, the high renin concentrations in the anterior and posterior uvea make this tissue a candidate for production of renin. Production in the retina is also conceivable. In the brain the same cell types that are present in the retina, both neuronal and glial cells, have been shown to synthesize renin (26). Since vitreous can be considered the sink of products originating from the retina (27), its high level of prorenin might well be explained by production in and secretion from the retina. The renin/prorenin ratio is quite different in uveal tissues, on the one hand, and in retina and vitreous, on the other hand, suggesting a third option that two distinct renin-producing systems exist in uvea and in retina separated by the blood-retinal barrier.

Reports have been published on components of the renin-angiotensin system

in the eye. Angiotensin converting enzyme has been found in bovine retina, in both vascular and neural fractions (28). Binding sites of angiotensin II are present in eye-vessels (29), apparently also on the abluminal side, not in reach of systemic angiotensin II, since intravitreal administration of this peptide causes vasoconstriction (30). Recently prorenin has been found by immunohistochemistry in human ciliary body (31). This raises the possibility that prorenin produced in the ciliary body is secreted into the aqueous. From there it may diffuse into the vitreous. However, we found prorenin in aqueous to be much lower than in the vitreous, and it is difficult to conceive how prorenin could be transported from a lower concentration in aqueous to a higher concentration in vitreous.

The function of a renin-angiotensin system in the eye is not yet clear. The higher concentration of renin in the uveal tract suggests that its major function is exerted there. Since the uveal tract is extremely vascular, and angiotensin II is a vasoactive peptide, it may serve a role in local blood supply. Topical administration of an angiotensin converting enzyme inhibitor to the eye lowers intraocular pressure (32). The renin-angiotensin system may, therefore, play a role in the control of intraocular pressure, possibly, in view of the presence of prorenin in the ciliary body (31), through an effect on the production of aqueous humor. Angiotensin II has trophic and mitogenic actions on vascular smooth muscle and other cells (33). We found prorenin to be increased in human vitreous from patients with proliferative diabetic retinopathy (13). In view of possible angiogenic effects of angiotensin II (12,34) the renin-angiotensin system may be a causative factor in proliferative retinopathy. In the ovary such a role of renin-angiotensin II has been proposed for the rapid vascularization of the developing corpus luteum which occurs after ovulation (35). Finally, inhibition of angiotensin converting enzyme in the isolated perfused feline eye causes changes in the electrophysiological response of the retina to light (36), suggesting that angiotensin II may have some neuromodulatory role in the retina.

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7. ANGIOTENSIN LEVELS IN THE EYE

7.1 Summary

Prorenin, the inactive precursor of renin, has been found in human vitreous fluid in concentrations that were too high to be explained by contamination with plasma. Moreover, its levels were twofold higher in vitreous fluid from eyes affected by proliferative diabetic retinopathy than in vitreous fluid from eyes without neovascularization, suggesting a role for the renin-angiotensin system in the development of retinopathy. Angiotensin (ANG) II has been reported to play a role in neovascularization. In the present study we measured angiotensin levels in ocular fluids obtained from diabetic and non-diabetic subjects, and in ocular tissues obtained from pigs. ANG I and ANG II were present in aqueous, vitreous and subretinal fluid of eyes from subjects with and without diabetic retinopathy. Large differences (by a factor of 10) in angiotensin concentrations were found between the various ocular fluid types, but after correction for leakage from plasma by expressing the angiotensin concentrations relative to ocular fluid albumin concentrations, all angiotensin concentrations became equal, indicating that ANG I and ANG II in human ocular fluid samples are largely plasma-derived. No differences were found between diabetic and non-diabetic subjects in either plasma or ocular fluid angiotensin concentrations. During a 2-hour i.v. infusion of ^{125}I -ANG I in the rabbit, less than 1 % of plasma ^{125}I -ANG I reached the vitreous fluid compartment, showing that, under normal conditions, angiotensins pass the blood-retinal barrier (BRB) only in very limited amounts. The relatively high angiotensin levels in human ocular fluid obtained during eye surgery are therefore probably the result of a partial breakdown of the BRB, leading to increased diffusion from plasma. Metabolism of ANG I in human ocular fluid occurred at a very low rate and was mainly due to conversion by ACE. Measurement of angiotensins in porcine ocular tissues showed that both ANG I and ANG II were present in retina, pigment epithelium-choroid and anterior uveal tract in concentrations much too high to be explained by contamination with plasma only. Low to undetectable angiotensin levels were found in porcine vitreous fluid. The ANG II/ANG I ratio was 3-5 fold higher in the ocular tissues than in plasma or vitreous fluid. These data strongly support the presence of a local renin-angiotensin system in the eye. In contrast with prorenin, little or no intraocularly synthesized ANG I and ANG II seems to be released into the ocular fluids, suggesting an autocrine or paracrine function of the intraocular renin-angiotensin system.

7.2 Introduction

According to the classical concept, angiotensin I (ANG I) and angiotensin II (ANG II) are generated within the circulation by sequential cleavage of liver-derived renin substrate. Renin, synthesized in the kidney, cleaves this substrate, forming ANG I, after which angiotensin converting enzyme (ACE) converts ANG I to ANG II, a potent vasoconstrictor and stimulant of the release of aldosterone from the adrenal. Recent evidence suggests that besides this circulating renin-angiotensin system, there also exist tissue or local renin-angiotensin systems in different organ sites, for instance in the adrenal, kidney, brain, testis and ovary (1-7). Local angiotensin production may therefore occur independent of circulating renin and circulating renin substrate.

Recently, we reported that high levels of prorenin, the inactive precursor of renin, are present in ocular fluids obtained from both diabetic and non-diabetic subjects during eye surgery (8). The prorenin concentrations were up to 100 times higher than expected based upon the plasma protein content of these samples. In vitreous fluid from eyes affected by proliferative diabetic retinopathy the prorenin concentrations were on average 2 times higher than in vitreous fluid from eyes of non-diabetic subjects, also when allowance was made for the differences in plasma protein content. The renin levels in most ocular fluid samples were close to the detection limit of the assay. These findings suggested that intraocular production of prorenin (and possibly renin) may occur, and that this production is increased in the presence of diabetic retinopathy.

Further studies in bovine eyes (9) showed indeed that both renin and prorenin were present in virtually all segments of the eye in concentrations much too high to be explained by the plasma content of the various ocular tissues only. The presence of ACE in the eye has been shown as well, not only in the retina, choroid and ciliary body (10), but also in aqueous fluid (11). Taken together, these findings suggest that the eye has its own renin-angiotensin system (RAS).

The function of an ocular RAS is not yet known. Its activation in the eyes of diabetic subjects with proliferative retinopathy may suggest that ANG II is involved in the development of neovascularization, as has been shown by others (12,13). As ANG II receptors have been detected in retinal blood vessels (14), a role for the RAS in the regulation of retinal vascular tone is also conceivable (15). Finally, the fact that both renin inhibitors (16) and ACE inhibitors (17) lower intraocular pressure would suggest that an intraocular RAS may play a role in the regulation of aqueous humor dynamics.

To our knowledge, no measurements of angiotensins in the eye have been reported so far. In this study, we present data on angiotensin levels in ocular fluid samples from diabetic and non-diabetic subjects. We also investigated the presence of ACE in human ocular fluid by measuring the amount of ^{125}I -ANG II formed in subretinal or vitreous fluid after the addition of ^{125}I -ANG I to

these fluids. In order to study whether plasma ANG I reaches the vitreous or aqueous fluid compartments, constant infusions of ^{125}I -ANG I were given to rabbits. The presence of radiolabeled angiotensins in the eye was studied at various time intervals during these infusions. Finally, an attempt was made to measure angiotensin levels in tissue extracts from porcine eyes.

7.3 Subjects and methods

Chemicals

[Ile⁵]-ANG-(1-10) decapeptide (ANG I), [Ile⁵]-ANG-(1-8) octapeptide (ANG II), and [Ile⁵]-ANG-(2-8) heptapeptide (ANG III) were obtained from Bachem, Bubendorf, Switzerland. [Ile⁵]-ANG-(2-10) nonapeptide (ANG-(2-10)) was from Senn Chemicals, Dielsdorf, Switzerland. [Ile⁵]-ANG-(3-8) hexapeptide (ANG-(3-8)), [Ile⁵]-ANG-(4-8) pentapeptide (ANG-(4-8)), and [Ile⁵]-ANG-(1-7) heptapeptide (ANG-(1-7)) were from Peninsula Laboratories, Belmont, CA, USA. Methanol, ethanol, ortho-phosphoric acid (all analytical grade), trisodium citrate and 1,10-phenanthroline were from Merck, Darmstadt, Germany. Bovine serum albumin (BSA) was from Sigma, St. Louis, MO, USA. Disodium EDTA was from Riedel de Haën, Scelze, Germany. Water for high performance liquid chromatography (HPLC) was prepared with a Milli-Q system from Waters, Millford, MA, USA. The renin inhibitor CGP 29,287 was a gift of Dr. K. Hofbauer (Ciba-Geigy, Basel, Switzerland). The renin inhibitor Ro 42,5892 was a gift of Dr. P. van Brummelen (Hoffmann-La Roche, Basel, Switzerland).

Preparation of radiolabeled angiotensins

Mono-iodinated ^{125}I -ANG I was prepared with the chloramine-T method and purified as described previously (19). ^{125}I -labeled preparations of ANG II, ANG III, ANG-(3-8), ANG-(4-8), ANG-(2-10), ANG-(1-7) and tyrosine were also made.

Separation of angiotensins by HPLC

Angiotensins and their metabolites were extracted from plasma, ocular fluid or tissue homogenates by reversible adsorption to octadecylsilyl silica (SepPak C18, Waters, Millford, MA, USA) and separated by reversed phase HPLC, according to the method of Nussberger et al (20) with some modifications (19). The SepPak cartridges were conditioned with 4 mL methanol and equilibrated with 2 times 4 mL of cold water. Samples were passed through the cartridge followed by a wash with 2 times 4 mL of cold water. Adsorbed angiotensins were eluted with 3 mL methanol into polypropylene tubes and the

methanol was evaporated under vacuum rotation, using a Savant Speed Vac concentrator (Savant Instruments, Farmingdale, NY, USA). Separations were performed on a reversed phase Nucleosil C18 steel column of 250 x 4.6 mm and 10 μ m particle size. Mobile phase A was 0.085 % ortho-phosphoric acid containing 0.02 % sodium azide. Mobile phase B was methanol. The flow was 1.5 mL/min and the working temperature was 45 °C. SepPak methanol extracts were dissolved in 100 μ L of HPLC solvent and injected. Elution was performed as follows: 65 % A/35 % B from 0 to 9 min followed by a linear gradient to 45 % A/55 % B until 18 min. The eluate was collected in 20-sec fractions into polystyrene tubes coated with BSA. The concentrations of 125 I-ANG I and its metabolites in the HPLC fractions were measured in the gamma counter. The fractions containing unlabeled ANG I and ANG II were neutralized with 0.5 M sodium hydroxide and vacuum dried at 4 °C. The concentrations of unlabeled ANG I and ANG II were measured by radioimmunoassay (19).

Assay of angiotensins

ANG I and ANG II concentrations were measured by radioimmunoassay after SepPak extraction and HPLC separation. The ANG I antiserum crossreacted with ANG-(2-10) (100 %) but not (less than 0.1 %) with ANG II, ANG III, ANG-(3-8), ANG-(4-8) and ANG-(1-7). The ANG II antiserum crossreacted with ANG III (55 %), ANG-(3-8) (73 %) and ANG-(4-8) (100 %), but not (less than 0.2 %) with ANG I, ANG-(2-10) or ANG-(1-7). The lower limit of detection was 0.5 fmol/fraction for the ANG II assay, and 1.0 fmol/fraction for the ANG I assay.

Measurement of albumin and renin substrate

Human albumin was measured by single radial immunodiffusion (LC and NOR-Partigen plates, Behringwerke, Marburg, Germany) according to the method of Mancini et al (21). The concentration of renin substrate was determined as the maximum quantity of ANG I that was generated during incubation at 37 °C and pH 7.5 with an excess of active purified human kidney renin in the presence of inhibitors of angiotensinases and ACE (22). The lower limit of detection was 1 nmol/L.

Human studies

Non-diabetic subjects. Aqueous fluid was collected at the time of cataract extraction from 26 subjects (20 women, 6 men; mean age 71 yr, range 38-92 yr). Nine subjects were receiving a diuretic and 3 a β -adrenergic antagonist.

Vitreous fluid aspirates were obtained from 18 subjects (11 women, 7 men; mean age 57 yr, range 10-80 yr). The samples were collected at the time of

pars plana vitrectomy, which was performed because of recurrent retinal detachment due to proliferative vitreoretinopathy (n=12), macula pucker (n=3) or vitreous hemorrhage (n=3). Three subjects were receiving a diuretic and 1 a β -adrenergic antagonist.

Subretinal fluid was obtained from 21 subjects (11 women, 10 men; mean age 55 yr, range 13-79 yr), with a rhegmatogenous (n=20) or traumatic (n=1) retinal detachment. Three subjects were receiving a diuretic.

Diabetic subjects. Vitreous fluid was obtained from 24 diabetic subjects (6 women, 18 men; mean age 51 yr, range 21-75 yr). The duration of diabetes ranged from 1 to 55 years. The eyes were affected by proliferative diabetic retinopathy and vitrectomy was performed because of traction retinal detachment (n=15) or vitreous hemorrhage (n=9). Seventeen subjects were taking insulin, 6 were receiving a diuretic, in 4 of them combined with a β -adrenergic antagonist. No subretinal nor aqueous fluid was collected from diabetic subjects.

Collection of plasma and ocular fluid samples. Approximately 0.1 mL aqueous fluid was collected with a tuberculin syringe and a 27 gauge needle. The needle was introduced at the limbus of the cornea through the groove of the cataract incision. Vitreous fluid was aspirated before substitution fluid was infused into the vitreous. Subretinal fluid was aspirated transsclerally, after local diathermic coagulation of the choroid.

The ocular fluid samples were free of macroscopically visible blood and were collected, after informed consent was obtained, in prechilled plastic tubes containing 25 μ L of inhibitor solution (containing 2 μ M CGP 29,287, 125 mM disodium EDTA and 25 mM 1,10-phenantroline) in order to block renin, ACE and angiotensinases. The samples were frozen immediately after collection at -70 °C. Some ocular fluid was frozen separately without inhibitor solution for studies on the metabolism of 125 I-ANG I and for the measurement of renin substrate and albumin.

A peripheral venous blood sample was drawn simultaneously with the collection of ocular fluid in all patients. Blood for angiotensin measurements (10 mL) was collected in prechilled tubes containing 0.5 mL of the above inhibitor solution and centrifuged at 3000 g for 10 minutes at 4 °C. Plasma was stored at -70 °C and assayed within two weeks. Blood for the measurement of renin substrate and albumin and for studies on the metabolism of 125 I-ANG I (10 mL) was collected in tubes containing 0.1 vol 0.13 M trisodium citrate. The blood was immediately centrifuged at 3000 g for 10 min at room temperature and plasma was stored at -70 °C.

Because of the relatively low angiotensin levels and the small size of most ocular fluid samples (less than 0.5 mL), angiotensin measurements in ocular fluid were only performed in pools consisting of several mL of ocular fluid. The recovery of angiotensins added to subretinal and vitreous fluid pools was

96 and 85 % respectively (n=2 for each). Plasma levels of ANG I and ANG II were measured in individual patients. Renin substrate and albumin were measured in all individual samples.

Metabolism of ¹²⁵I-ANG I in human ocular fluid and plasma. The metabolism of ¹²⁵I-ANG I was studied in pools (2 mL) of subretinal fluid, vitreous fluid and plasma. After the ocular fluid and plasma pools had been brought to a temperature of 37 °C in a waterbath, the experiment was started by adding at t=0 50.000 counts per minute (cpm) of ¹²⁵I-ANG I (in 50 µL of TRIS buffer). Experiments were done in the presence or absence of the ACE inhibitor captopril (final concentration in the sample 0.4 mM). Samples of 400 µL were taken at various times (at 0, 1, 2 and 5 min for plasma, at 0, 5, 10 and 20 min for subretinal fluid and at 0, 30 and 60 min for vitreous fluid) and immediately mixed with 50 µL inhibitor solution. The samples were kept on ice and SepPak extraction was performed within one hour. The SepPak extracts were applied to the HPLC column and ¹²⁵I-ANG I and ¹²⁵I-ANG II were measured by gamma counting as described before. Results are expressed as a percentage of total cpm recovered after SepPak extraction.

¹²⁵I-ANG I infusions in rabbits

All procedures were in accord with the ARVO Resolution on the Use of Animals in Research. Three rabbits, weighing 2.9 - 3.3 kg, were anesthetized using intramuscular xylazine (Bayer, Leverkusen, Germany; 2 mg/kg) and ketamine HCl (Aesculaap BV, Boxtel, The Netherlands; 20 mg/kg). Additionally, they received topical oxybuprocaine HCl (M.S.D., Haarlem, The Netherlands) for local anesthesia. The ACE inhibitor ramipril was injected intraperitoneally (1 mg/kg) shortly before the experiment. This was performed in order to slow down the fast breakdown of infused ¹²⁵I-ANG I. ¹²⁵I-ANG I (mean concentration 3.4×10^6 cpm/mL) was infused at a rate of 0.1 mL per minute in the ear lobe vein during two hours. Arterial samples (1-2 mL) were taken from the carotid artery at 15, 30, 45, 60, 90 and 120 min after the start of the infusion. The blood was collected in prechilled tubes containing 0.1 mL of the above inhibitor solution and centrifuged at 3000 g for 10 minutes at 4 °C. One mL of plasma was counted in a gamma counter for measuring total radioactivity. After 1 hour, an aqueous and a vitreous sample were removed from one eye with an injection needle with a 25 and a 19 gauge respectively, and total radioactivity was counted in both samples. At the end of the infusion period the animals were killed and immediately aqueous and vitreous samples were taken from the second eye and counted in the gamma counter.

After the experiment, the ocular fluid and plasma samples were passed through SepPak columns and applied to the HPLC, as described before. Radiolabeled angiotensins were measured by gamma counting.

Angiotensin levels in porcine eyes

Eyes were taken from 20 anesthetized pigs (weight 25-50 kg, crossbred Yorkshire x Landrace, H.V.C., Hedel, The Netherlands). The pigs had been used for various pharmacological studies not involving the eye. An adequate depth of anesthesia had been maintained by the administration of 160 mg/kg α -chloralose (Merck, Darmstadt, Germany) in the superior caval vein, followed by continuous i.v. infusion of low dose sodium pentobarbitone (5 mg/kg per h). The eyes were dissected immediately following enucleation. Each eye was cut equatorially at the ora serrata, and the anterior segment was lifted off. The vitreous body was isolated by gently shaking it out of the eye cup. The neural retina was cautiously removed from the pigment epithelium with a thin glass rod and isolated by cutting it at the optic nerve. The choroid with adhering retinal pigment epithelium layer was isolated by dissecting it from the sclera with a pair of fine scissors. The anterior uveal tract, consisting of iris and ciliary body, was isolated by removing the lens from the anterior eye cup, then gently pulling the anterior uveal tract loose from the sclera and blotting it on dry paper to remove any adhering vitreous. Cornea, lens and sclera were discarded, whereas retina, pigment epithelium-choroid, anterior uveal tract and vitreous were snap frozen on dry ice and stored at -70°C . The whole procedure took 2-3 min.

Simultaneously with the removal of the eyes a peripheral blood sample was taken, after which the animals were killed. The blood (10 mL) was collected in a chilled polystyrene tube containing 0.5 mL inhibitor solution (containing 0.2 mM Ro 42,5892, 125 mM disodium EDTA and 25 mM 1,10-phenantroline). The blood was centrifuged at 3000 g for 10 min at 4°C . Plasma was stored at -70°C and assayed as described above within 2 weeks.

Pools of tissues obtained from 10 pigs (20 eyes) were homogenized with a Polytron PT10/35 (Kinematica, Luzern, Switzerland) in an iced solution of 0.1 M HCl/80 % ethanol according to the method of Chappell et al (23) with some modifications. In short, homogenates were centrifuged at 4°C for 20 min at 20,000 g and the supernatant was stored for 12 hrs at -20°C . The supernatant obtained after a second centrifugation step (4°C , 20,000 g, 20 min) was diluted 1:1 (v:v) with 1 % ortho-phosphoric acid, stored for 4-6 hrs at 4°C , and again centrifuged at 20,000 g. The final supernatant was further diluted 1:1 with 0.2 % ortho-phosphoric acid and concentrated on a SepPak column. Vitreous was concentrated on SepPak columns without further additions. SepPak extracts were applied to the HPLC as described above and angiotensins were measured by RIA.

Metabolism of angiotensins during storage at -70°C was studied by adding ^{125}I -ANG I to frozen tissues one week before homogenization. The efficiency of the extraction was determined by the addition of 200,000 CPM ^{125}I -ANG I immediately before homogenization. Overall recovery was 59 ± 9 % (mean and SD, n=6).

Statistical evaluation

Angiotensin levels in ocular fluid and plasma are expressed as mean and range. Because of non-linear distribution, albumin and renin substrate in ocular fluid and plasma are expressed as geometric mean and range. Unpaired t-tests were performed to analyze differences between diabetic and non-diabetic subjects. Values were considered significant if $p < 0.05$.

7.4 Results

ANG I, ANG II, renin substrate and albumin levels in human ocular fluid and plasma

Plasma angiotensin levels were similar in diabetic and non-diabetic subjects (Table 1). In non-diabetic subjects, the highest ocular angiotensin levels were found in subretinal fluid and the lowest (close to the detection limit of the assay) in aqueous fluid. ANG I and ANG II levels in vitreous fluid of diabetic subjects were similar to those in vitreous of non-diabetic subjects, although relative to ANG I, ANG II tended to be slightly higher ($p > 0.05$) in vitreous from diabetic subjects than in vitreous from non-diabetic subjects. The ANG I and ANG II concentrations in subretinal, vitreous and aqueous fluid were approximately 100, 50 and 10 % of those in plasma.

No differences in plasma or vitreous fluid albumin and renin substrate levels were found between diabetic and non-diabetic subjects (Table 2). As for ANG I and ANG II, the highest concentrations of albumin and renin substrate were found in subretinal fluid and the lowest in aqueous fluid. The concentrations of albumin and renin substrate in subretinal, vitreous and aqueous fluid were approximately 10, 5 and 1 percent of those in plasma.

The ocular fluid/plasma concentration ratio of renin substrate equalled that of albumin and both ratios were strongly correlated ($r=0.99$, $p < 0.001$), suggesting that both proteins enter the ocular fluid to the same degree, presumably by diffusion from plasma. This confirms earlier findings (8). Expressing the angiotensin levels relative to albumin (Figure 1) results in similar angiotensin levels in all ocular fluids. Apparently, the differences in angiotensin levels found between subretinal, vitreous and aqueous fluid parallel the differences in albumin (and renin substrate) levels, and therefore angiotensins in ocular fluid are most likely plasma-derived.

Table 1. Levels of ANG I and ANG II in individual plasma samples and ocular fluid pools of diabetic and non-diabetic subjects.

	n	ANG I (pM)	ANG II (pM)
<u>Non-diabetic subjects</u>			
plasma	39	27.0 (1.9-86.6)	8.0 (0.7-20.7)
subretinal fluid	3	24.5 (14.5-37.8)	5.4 (5.2-5.8)
vitreous fluid	3	12.7 (7.0-17.2)	2.3 (1.3-3.0)
aqueous fluid	2	2.0 (1.7-2.2)	0.5 (0.5-0.6)
<u>Diabetic subjects</u>			
plasma	16	23.6 (3.9-94.7)	6.7 (0.9-24.9)
vitreous fluid	4	8.7 (2.9-21.2)	4.4 (2.5-6.7)

Values are given as mean and range.

Table 2. Levels of albumin and renin substrate in ocular fluid and plasma of diabetic and non-diabetic subjects.

	n	Albumin (g/L)	Renin Substrate (nM)
<u>Non-diabetic subjects</u>			
plasma	39	32.5 (24.1-40.9)	1038 (572-1667)
subretinal fluid	21	3.05 (0.35-29.6)	99 (21-963)
vitreous fluid	18	1.25 (0.12-20.3)	42 (5-643)
aqueous fluid	26	0.24 (0.06-0.45)	6 (2-15)
<u>Diabetic subjects</u>			
plasma	16	31.1 (22.9-38.4)	937 (557-1280)
vitreous fluid	24	1.87 (0.84-14.6)	58 (24-635)

Values are given as geometric mean and range.

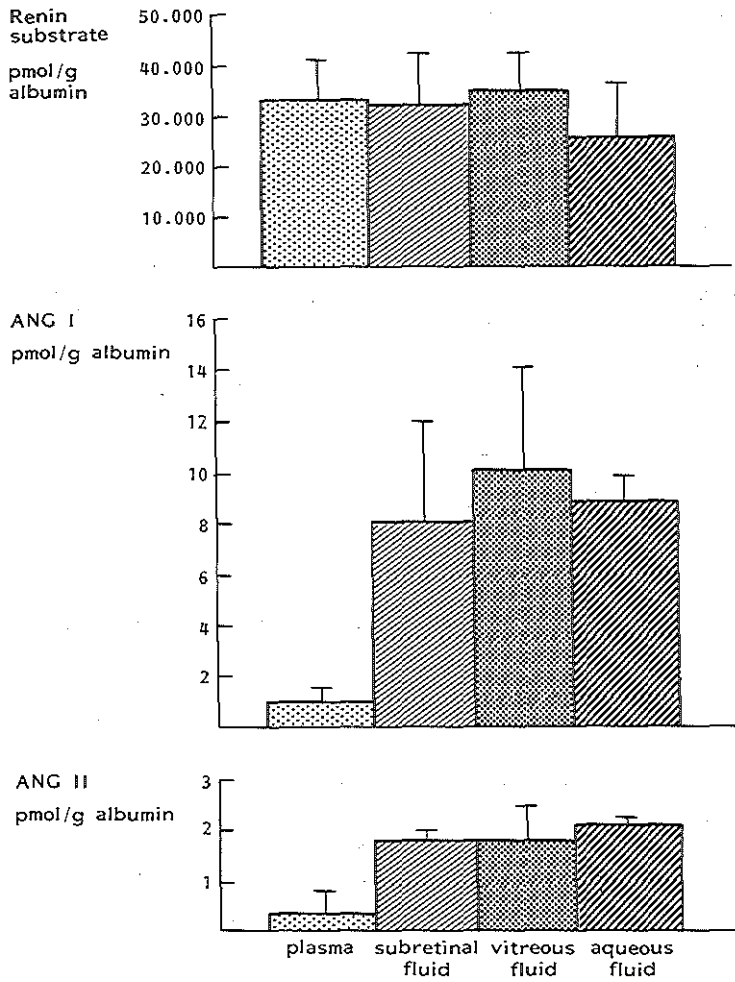


Figure 1. Levels (mean and SD) of renin substrate, ANG I and ANG II relative to those of albumin in plasma, subretinal fluid, vitreous fluid and aqueous fluid of non-diabetic subjects.

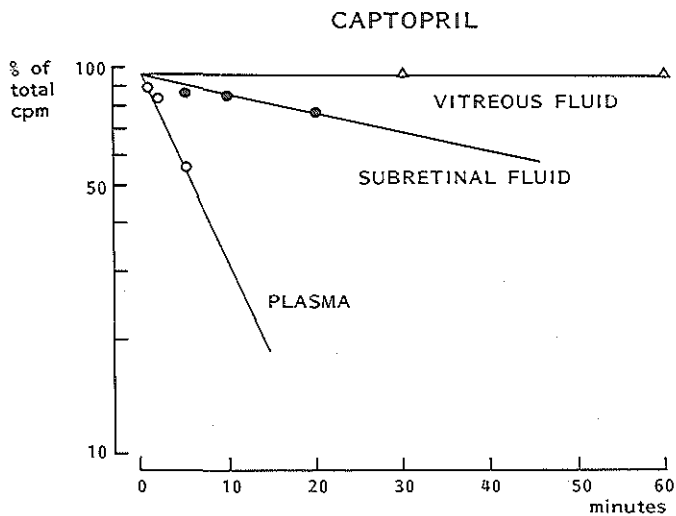
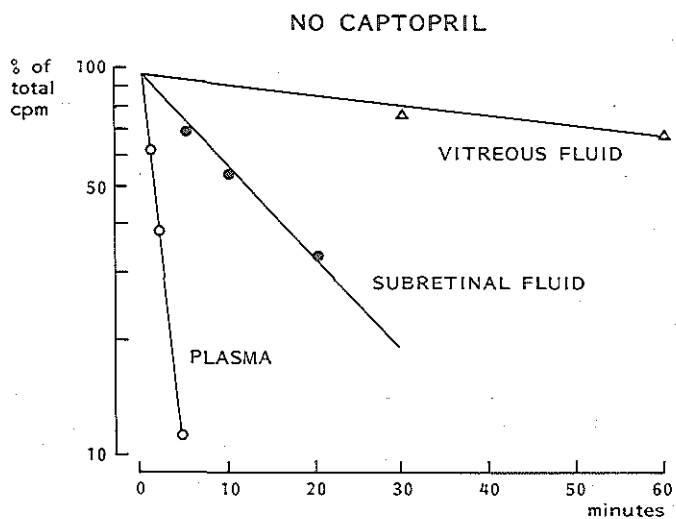


Figure 2. Metabolism of ^{125}I -ANG I in plasma, subretinal fluid and vitreous fluid during incubation at 37°C in the absence (top panel) or presence (bottom panel) of captopril. Data are means of 2 experiments each.

Metabolism of ¹²⁵I-ANG I in human ocular fluid and plasma

Incubating pools of plasma, pools of subretinal fluid and pools of vitreous fluid with ¹²⁵I-ANG I at 37 °C resulted in the prompt breakdown of the peptide (Figure 2). The half lives of ¹²⁵I-ANG I in plasma, subretinal fluid and vitreous fluid were 1.5, 14 and 134 minutes respectively (mean of 2 experiments each). HPLC separation of the various metabolites clearly showed that formation of ¹²⁵I-ANG II occurred (Figure 3). In vitreous only ¹²⁵I-ANG II and no other metabolites were formed.

In the presence of the ACE inhibitor captopril the breakdown of ¹²⁵I-ANG I was markedly reduced in subretinal fluid (half life 87 min, n=2) and plasma (half life 5.4 min, n=2) and completely inhibited in vitreous fluid (n=2) (Figure 2). ¹²⁵I-ANG II was no longer present among the metabolites, indicating that it is indeed ACE that is responsible for the formation of ¹²⁵I-ANG II in both ocular fluid and plasma. Based on the difference in half life with and without captopril, it could be calculated that 70-80 % of ANG I metabolism in plasma and subretinal fluid, and 100 % of ANG I metabolism in vitreous fluid is due to ANG I-II conversion by ACE.

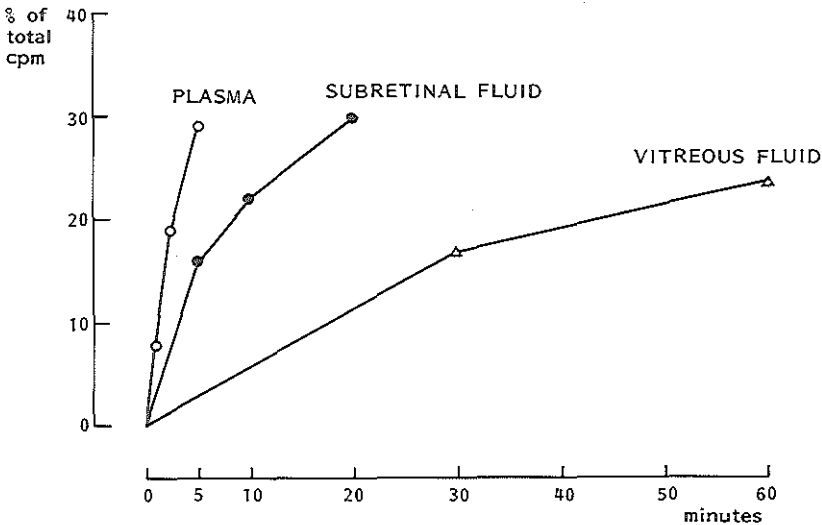


Figure 3. Formation of ¹²⁵I-ANG II during incubation of ¹²⁵I-ANG I at 37 °C in plasma, subretinal fluid and vitreous fluid. Data are means of 2 experiments each.

¹²⁵I-ANG I infusions in rabbits

Total radioactivity in plasma increased with time, but the ¹²⁵I-ANG I levels reached a plateau within 15 minutes which did not change during the infusion. The mean plateau concentration in the carotid artery (n=3) was 550 (range 165-1060) cpm ¹²⁵I-ANG I/mL plasma. Expressed as a percentage of total plasma radioactivity, the total radioactivity in aqueous and vitreous fluid did not differ between samples from the first (t=1 h) and the second (t=2 h) eye. The radioactivity levels were in aqueous fluid 25.8 (range 22.9-27.3) % and in vitreous fluid 2.4 (range 1.6-3.5) % of concomitant plasma values. HPLC separation of the radioactive components in ocular fluid revealed that virtually all radioactivity concentrated in the first few HPLC fractions. These fractions are known to contain small radiolabeled metabolites (predominantly ¹²⁵I-tyrosine). Only very limited amounts of ¹²⁵I-ANG I were found in aqueous and vitreous fluid: 21 ± 24 cpm/mL and 6 ± 2 cpm/mL, respectively. The aqueous, but not the vitreous fluid ¹²⁵I-ANG I concentrations correlated significantly (r=0.98; p < 0.001) with the plasma levels of ¹²⁵I-ANG I.

These findings demonstrate that, in normal healthy rabbits with presumably intact blood-retinal barriers, only approximately 1 % of plasma ANG I reaches the vitreous compartment and 4 % the aqueous compartment.

Angiotensin levels in porcine eyes

The levels of ANG I and ANG II in porcine plasma, vitreous and ocular tissues are given in Table 3. Figure 4 shows the results in a pigment epithelium-choroid extract. Not only ANG I and ANG II were detected, but small amounts of ANG III were present as well (Fig. 4). In absolute terms (expressed per g tissue, plasma or vitreous fluid) the levels of ANG I at all ocular sites were lower than those in plasma. The ANG II levels in the retina and anterior uveal tract were similar to or slightly higher than those in plasma, whereas in the pigment epithelium-choroid ANG II was threefold higher than in plasma. The relatively high tissue levels of ANG II were not the result of *in vitro* conversion of ANG I, nor was ANG I metabolized by other degrading enzymes during storage or purification, since adding ¹²⁵I-ANG I during storage or immediately before homogenization did not result in the formation of any angiotensin metabolite other than ¹²⁵I-ANG I. Previous studies have documented that ¹²⁵I-labeled angiotensins have similar recoveries to unlabeled peptides as determined by RIA after extraction and HPLC separation (24,25). In vitreous fluid ANG I and ANG II were frequently close to or below the detection limit of the assay. Vitreous angiotensin levels were less than 10 percent of concomitant plasma angiotensin levels. The vitreous ANG II/ANG I ratio was similar to that in plasma, suggesting that ANG I and ANG II in vitreous fluid are probably largely plasma-derived. The ANG II/ANG I ratios in ocular tissues were 3-5 fold higher than the plasma ANG II/ANG I ratio.

Table 3. Levels of ANG I and ANG II in porcine plasma, vitreous and ocular tissues.

	ANG I (fmol/g)	ANG II (fmol/g)	ANG II/ANG I ratio
plasma	9.7	7.5	0.8
vitreous fluid	< 1.1	< 0.9	0.8
retina	3.1	11.2	3.6
pigment epithelium-choroid	8.0	23.2	2.9
anterior uveal tract	3.1	12.4	4.1

Values are means of individual samples (plasma and vitreous fluid) or means of 2 measurements in tissue pools (taken from 20 eyes each).

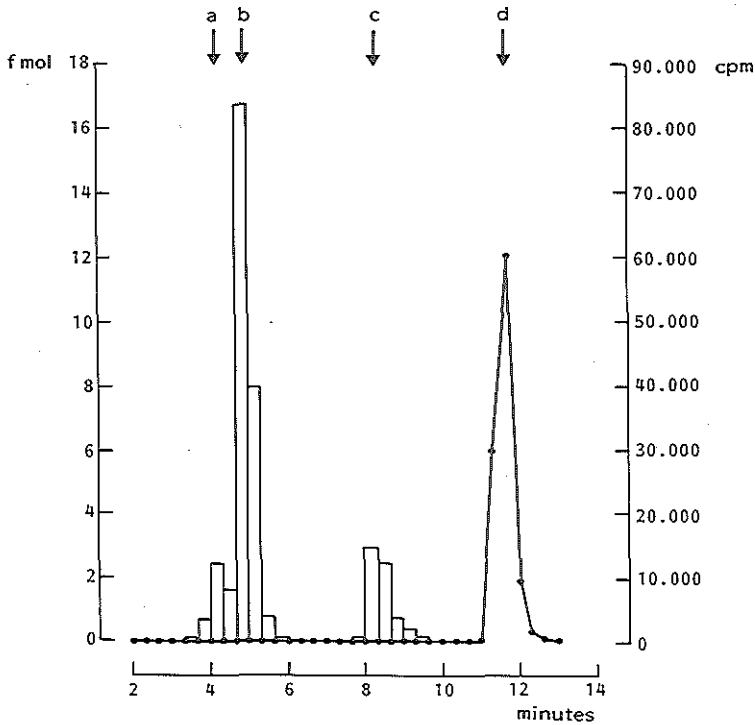


Figure 4. HPLC elution profile of endogenous angiotensins (bars) in a porcine pigment epithelium-choroid tissue extract. ¹²⁵I-ANG I (closed circles) was added to study metabolism during storage and purification (see *Subjects and methods* for details). The retention times of standard angiotensins are given as a (ANG III), b (ANG II), c (ANG I) and d (¹²⁵I-ANG I).

7.5 Discussion

ANG I and ANG II are present in ocular fluid samples taken from patients undergoing eye surgery. No differences in plasma or vitreous fluid angiotensin content were found between diabetic and non-diabetic subjects. In non-diabetic subjects, the highest angiotensin levels were present in subretinal fluid (equalling those in plasma) and the lowest (close to the detection limit of the assay) in aqueous fluid. Similar findings were done for albumin and renin substrate, confirming earlier data (8). Relative to concomitant plasma levels, the concentrations of albumin and renin substrate were approximately 10-fold lower than those of ANG I and ANG II. The ocular fluid/plasma concentration ratio of renin substrate equalled that of albumin in each individual sample, suggesting that both proteins enter the ocular fluid to the same degree, presumably by diffusion from plasma. The vitreous concentration of plasma proteins in normal, healthy eyes has been described to be only 1-2 % of concomitant plasma concentrations (26,27). The higher levels (5 % of plasma levels) we found in vitreous fluid for both albumin and renin substrate are probably due to partial breakdown of the blood-retinal barrier (BRB), leading to increased diffusion. Since no differences were found in the albumin and renin substrate content of vitreous fluid taken from diabetic and non-diabetic subjects, the degree of breakdown of the BRB must have been comparable in both groups of subjects.

As shown by our experiments in the rabbit, approximately 1 % of arterially present ^{125}I -ANG I enters the vitreous fluid compartment of healthy eyes during a 2-hour infusion. The levels reached in aqueous fluid were in a similar low range (4 % of arterial plasma levels), and correlated significantly with the arterial ^{125}I -ANG I levels. Apparently, the plasma ANG I levels determine the aqueous levels of ANG I. That no correlation was found with the concentrations in vitreous fluid is probably due to their extremely low levels. Following intravenous infusion, fluorescein (which has a molecular weight 4-5 fold lower than that of angiotensins) has also been reported to reach very low levels in vitreous and aqueous fluid (28-30). Fluorescein levels in vitreous were usually below 1 % of concomitant plasma levels. Others have shown (31,32) that ANG I nor ANG II is able to pass the blood-brain barrier, which is very comparable, if not equal to the BRB (33). Microperoxidase, which has a molecular weight in the same range as ANG I and ANG II, does not pass the BRB in normal rats (34). Thus, we may assume that in normal, healthy eyes with an intact BRB plasma ANG I (and ANG II) cross the BRB, if at all, only in very limited amounts. Partial breakdown of the BRB however, may lead to increased diffusion of angiotensins from the blood into the eye. Leakage of fluorescein into the eye for example is known to be increased 2-3 times in diabetic rats as compared with normal rats (35), and a direct correlation has been described between fluorescein leakage and breakdown of the BRB (36). ANG I and ANG II in ocular fluid may therefore, like albumin and renin

substrate, also be derived from plasma. Local generation of ANG I in ocular fluid seems unlikely because of the low levels of renin substrate and renin (8) reported in the various ocular fluids. The present study shows that ACE is present in subretinal and vitreous fluid, but the low rate at which ANG I-II conversion in these fluids occurs as compared with ANG I-II conversion in plasma, suggests that ANG II formation in ocular fluid is also of minor importance. An alternative for leakage from plasma or generation in ocular fluid would be that angiotensins are released from an intraocular angiotensin producing source, for instance the retina or the ciliary body. One would then expect a concentration gradient for ANG I and ANG II to exist independent of the plasma protein content of the samples. We have shown this for prorenin (8,9). However, after correction for leakage from plasma by expressing the angiotensin levels relative to those of albumin no more concentration differences were found between the various ocular fluids. This suggests that ANG I and ANG II in ocular fluid are indeed largely plasma-derived. The angiotensin ocular fluid/plasma concentration ratio being higher than the albumin and renin substrate ocular fluid/plasma concentration ratio is probably due to the much smaller size of ANG I (mol wt, 1.3 K) and ANG II (mol wt, 1.0 K) as compared with albumin (mol wt, 69 K) and renin substrate (mol wt, 65 K), leading them to diffuse far more easier from the circulation into the various ocular fluids. This is in accordance with our earlier observation (8) that the concentration of IgG (mol wt, 150 K) in vitreous fluid, relative to that of albumin, was somewhat lower than that in plasma due to its larger molecular size.

The high levels of prorenin, renin and ACE found by us and others in retina, choroid and ciliary body (9,10,37) favour the possibility that ANG I and ANG II are produced within ocular tissue(s) rather than in ocular fluid. The present study shows that both renin substrate and angiotensins in ocular fluid are largely plasma-derived, suggesting that if angiotensins are produced intraocularly they remain in the tissues where they are produced rather than that they enter the vitreous or aqueous fluid compartment. Possibly the rate at which angiotensins are metabolized in ocular tissues is too high to allow overflow into ocular fluids. Alternatively, angiotensin production may occur in compartment which does not equilibrate with the vitreous or aqueous fluid. Prorenin seems to be the only component of the intraocular RAS that does leak into ocular fluid. Both human and bovine vitreous contained virtually only prorenin and no renin (8,9). Cultured bovine theca cells (38) and rat adrenal glomerulosa cells (39) have also been reported to release only prorenin and no renin into the medium. In these cells, both prorenin and renin were found, suggesting that angiotensin production occurs presumably intracellularly only.

To test our hypothesis that angiotensins are produced locally within the eye, we measured angiotensin levels in plasma, vitreous and various ocular tissues of the pig. Pigs that had been used for various pharmacological studies were enucleated shortly before they were killed. In this way it was possible to

remove the eyes under standardized conditions, reducing the time between enucleation and freezing of the various tissues to a minimum of 2-3 min, a situation which proved to be impossible in the local slaughterhouse. Angiotensin levels in porcine vitreous fluid were extremely low, close to or below the detection limit of the assay. Plasma angiotensin levels were at least 10 times higher, confirming our finding in the rabbit that virtually no plasma ANG I nor plasma ANG II reaches the vitreous fluid compartment in normal eyes. In contrast, the angiotensin levels in the retina, pigment epithelium-choroid and anterior uveal tract (iris + ciliary body) were too high to be explained by contamination with plasma only. Expressed per g of tissue weight ocular ANG II levels were similar to or higher than plasma ANG II levels, whereas ocular ANG I levels were 30-80 % of plasma ANG I levels. Previous studies in bovine eyes (9) have shown that the plasma content of the retina, anterior uveal tract and pigment epithelium-choroid is only 1, 5 and 20 % of total tissue weight respectively, and one may assume that this is similar in porcine eyes. Thus, the high angiotensin levels in ocular tissues can only be explained by assuming either that angiotensins are taken up selectively from the circulation, or that they are synthesized locally. In disagreement with the former explanation is the fact that the tissue ANG II/ANG I ratio was 3-5 fold higher than the plasma ANG II/ANG I ratio. In vitro formation or metabolism of angiotensins is unlikely to be the explanation of this finding, for it was shown that no angiotensin metabolism occurred during either storage or purification of the samples. No metabolites of ¹²⁵I-ANG I were found after purification and HPLC separation, no matter whether ¹²⁵I-ANG I was added shortly before the homogenization or already during storage at -70 °C. Our findings therefore suggest that ANG I and ANG II in ocular tissues are synthesized locally rather than taken up from plasma.

In conclusion, we have shown that angiotensins are present in human ocular fluid samples as well as in porcine ocular tissues. Angiotensins in human ocular fluids do not seem to be generated within these fluids because of the low levels of renin, renin substrate and ACE found in these fluids, nor are they the result of angiotensin release from intraocular tissue sites. Comparison with ocular albumin levels suggests that angiotensins in human ocular fluid are probably largely plasma-derived. Prorenin, the inactive precursor of renin, is the only component of the intraocular RAS that is released into ocular fluid. The relatively high levels of ANG I and ANG II in porcine retina, pigment epithelium-choroid and anterior uveal tract, which cannot be explained based upon the presence of plasma in these tissues only, strongly support the idea of intraocular angiotensin generation. The virtual absence of angiotensins in porcine vitreous fluid and our finding that angiotensins in human ocular fluid samples are mainly plasma-derived suggest that the rate at which angiotensins are metabolized in ocular tissues is too high to allow overflow into ocular fluids, or that angiotensin production occurs in compartment which does not equilibrate with ocular fluids. Most likely, ANG II generated in ocular tissues

exerts a autocrine or paracrine function. An intraocular RAS may play a role in the development of diabetic retinopathy, in the regulation of retinal vascular tone or in the regulation of aqueous humor dynamics.

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8. DEMONSTRATION OF RENIN-, ANGIOTENSINOGEN- AND ANGIOTENSIN CONVERTING ENZYME-mRNA EXPRESSION IN HUMAN EYES BY THE POLYMERASE CHAIN REACTION

8.1 Introduction

The presence of all components of the renin-angiotensin system has been demonstrated in the eye, not only in ocular fluids obtained from human eyes (1,2) during eye surgery, but also in ocular tissues from bovine and porcine eyes (2,3). The levels were too high to be explained by contamination with plasma. The levels of prorenin were higher in vitreous fluid from eyes with proliferative diabetic retinopathy than in vitreous fluid from eyes of non-diabetic subjects (1). This suggests that an intra-ocular renin-angiotensin system may be involved in the pathogenesis of diabetic proliferative retinopathy. Angiotensin II has been implicated in the development of neovascularization (4).

However, definite proof of intra-ocular production of renin, angiotensinogen and angiotensin converting enzyme (ACE) in human eyes, as opposed to specific uptake from plasma, can only be obtained by demonstrating ocular renin-, angiotensinogen- and ACE-mRNA expression. In the present study we employed the RNase protection assay to detect renin-mRNA expression in fresh human ocular tissues. Since the ocular total mRNA content was at the limit of detection of the RNase protection assay, we also applied the polymerase chain reaction (PCR) to improve sensitivity. For synthesis of double-stranded cDNA, ocular mRNA was reversely transcribed and subsequently renin-cDNA was amplified by PCR using two primers flanking 376 basepairs (bp) of human renin-cDNA and spanning the second and third exon. Angiotensinogen- and ACE-mRNA were determined by primers spanning a 240 bp or 380 bp fragment, respectively. The amplified cDNAs were processed by Southern blotting and hybridization with the corresponding randomly labeled probes.

8.2 Subjects and methods

Subjects. Ocular tissues were removed from enucleated eyes within 1-2 min after enucleation and frozen in liquid nitrogen. The eyes were obtained from 18 subjects (12 men and 6 women; mean age 54 yr, range 26-78 yr). The indications for enucleation were: choroidal melanoma (n=11), ciliary body melanoma (n=1), phtisis bulbi (n=2), neovascular glaucoma (n=2), bullous keratopathy in a blind eye (n=1) and an inflamed blind eye after trauma (n=1). Depending on what part of the eye had to be sent to the pathology department, retina, retinal pigment epithelium-choroid, anterior uveal tract and sclera were isolated as described before (2,3). Sometimes it was not possible to separate retina and pigment epithelium-choroid from each other. The ocular tissues were stored at -70 °C.

RNAse protection assay for renin. ³²P-labeled RNA-transcripts were prepared by transcription of a 291 nucleotide antisense RNA from a SacI/EcoRV fragment of the human renin cDNA subcloned into pGEM4 vector using T7 RNA polymerase. This transcript comprised 225 nucleotides of human renin antisense RNA and 66 nucleotides of vector encoded sequence.

Total RNA was isolated from combined human ocular tissues by lithium chloride precipitation (5). RNAse protection assays were performed according to Goedert et al (6). Samples of dried RNA were dissolved in 30 µL 80 % formamide, containing 40 mM PIPES (pH 6.8), 400 mM NaCl, 1 mM EDTA and 200.000 cpm of the gel-purified transcript. After denaturation at 95 °C for 60 sec and incubation at 50 °C for 20 h, RNAse digestion was performed in 300 µL buffer containing 40 µg/mL RNAse A (Sigma, St. Louis, MI, USA) and 2 µg/mL RNAse T1 (Calbiochem, USA) for 45 min at 37 °C. After digestion with proteinase K, samples were electrophoresed on denaturing 5 % polyacrylamide gels.

Polymerase chain reaction. Total RNA was isolated from individual human ocular tissue samples by a modification of the lithium chloride method (5). After isolation, total RNA samples were checked by gel electrophoresis in an 1 % agarose gel stained with ethidium-bromide after denaturation with 6 M glyoxal, 0.25 M DMSO and 0.1 M NaH₂PO₄ (pH 7.0) at 50 °C for 60 min.

For use in the polymerase chain reaction, total RNA was reversely transcribed to cDNA according to Wang et al (7). 1 µg of total RNA was dissolved in 20 µL of a reaction mixture containing 1 mM of dATP, dCTP, dTTP and dGTP, 1 U of RNAsin (Boehringer Mannheim, Germany), 100 pmol random hexamers (Boehringer Mannheim), 50 mM KCl, 20 mM Tris-HCl (pH 8.4), 2.5 mM MgCl₂, 10 µg/µL nuclease-free bovine serum albumin and 200 U of Murine leukemia virus reverse transcriptase (MULV-RT, Gibco BRL, Germany). After incubation for 45 min at 42 °C, temperature was raised to 95

°C and then quickly lowered on ice. For amplification of the resulting cDNA, the sample volume was increased to 100 µL by a solution containing 50 mM KCl, 20 mM Tris-HCl (pH 8.4) and 2.5 mM MgCl₂ and 25 pmol of up- and downstream primers as well as 3 U of Taq polymerase (Perkin-Elmer, Cetus, USA). The thermal profile used on a Perkin Elmer/Cetus thermal cycler consisted of denaturation at 95 °C for 1 min, annealing at 60 °C for renin and at 55 °C for angiotensinogen and angiotensin converting enzyme, respectively, for 1 min, and an extension temperature of 72 °C for 1 min for 26 cycles. After PCR, 10 µL loading buffer (50 % glycerol, 10 mM Tris-HCl (pH 8.0) and 0.25 % bromphenol blue and xylene cyanol) were added to each sample and the amplification products were checked for the predicted sizes by agarose gel electrophoresis and then submitted to Southern blot analysis. Primers were selected with the computer program developed by Lowe et al (8), which was licensed to M. Paul. Human renin-cDNA was amplified by 21-mer oligonucleotides with the following sequences: AAATGAAGGGGGTGTCTGTGG as sense primer (bases 851-872) and AAGCCAATGCGTTGTTACGC (bases 1206-1227) as antisense primer. This yielded an amplification product of 376 bp in length spanning the second and third exon of renin-cDNA. The sense primer for the detection of ACE-cDNA spanned oligonucleotide bases 492-512 (GCCTCCCCAACAAGACTGCCA), and the antisense primer spanned base 860-880 (CCACATGTCTCCAGCCAGATG) of the human ACE-cDNA. Human angiotensinogen primers were situated over the fourth and fifth exon with the sense primer (bases 1209-1231) CTGCAAGGATCTTATGACCTGC and the antisense primer (bases 1404-1426) TACACAGCAAACAGGAATGGGC.

Southern blotting was performed as described (9). The amplified cDNA sequences were transferred from 1.3 % agarose gels to nylon membrane (Pall, Great Britain) in a LKB-vacuum blot chamber using 0.25 N HCl for precipitation for 30 min and subsequent neutralization on 0.5 N NaOH and 1.5 M NaCl. Blotting was terminated after 2 h on 20 x SSC (1 x SSC: 0.15 M NaCl, 0.015 M sodium citrate). cDNA was crosslinked to the nylon membrane in a UV-linker (Stratagene 1800). Membranes were prehybridized with 0.5 % deionized formamide, Denhard's solution, 25 µm/mL herring sperm DNA for 4 h. Hybridization was done in the same buffer overnight at 60 °C adding the corresponding probes, which were randomly labeled using ³²P-dCTP. Labeled probes were purified on a Sephadex G-50 column. Renin-cDNA was hybridized to a 1.3 kb long BamHI/HindIII rat renin-cDNA fragment obtained from the complete rat renin-cDNA cloned into a pGEM4 vector. The restriction fragment was separated from the vector on an agarose gel and isolated by Qiaex (Quiagen, Germany). A plasmid vector (Bluescript KS, pB 35-19) containing 3334 bp of human ACE-cDNA was cut with EcoRI and Bgl II to yield 1.7 and 1.6 kb inserts of ACE-cDNA, which were isolated from the 3.0 kb as probes. A 1 kb StuI fragment of a pGEM5 vector containing the

human angiotensinogen-cDNA allowed detection of angiotensinogen-cDNA. Nylon membranes were washed after hybridization at room temperature in 0.2 x SSC and 0.1 % sodiumdodecylsulphate for 30 min and 3 times at 56 °C for 30 min. Blots were exposed for 18-36 hrs at -80 °C to XAR x-ray films (Kodak).

8.3 Results

Using the RNase protection assay it was possible to demonstrate renin-mRNA expression in human pigment epithelium-choroid/retina, but not in retina alone or in sclera (Figure 1). These results were obtained by pooling tissues from several eyes, since the amount of total RNA recovered from individual eye samples was below the detection limit of the assay. Compared to the kidney, where 20 µg of RNA was used, the ocular renin mRNA levels seemed to be low, since 45 µg of RNA from pigment epithelium-choroid/retina did, but 25 µg of retinal RNA and 35 µg of scleral RNA did not reveal any renin-mRNA expression by this method.

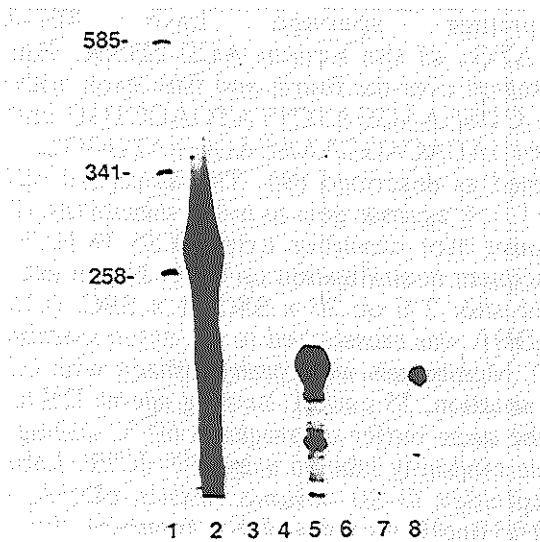


Figure 1. Detection of renin mRNA by RNase protection assay. Autoradiograph of 5 % polyacrylamide gel. Results in human ocular samples were obtained by pooling tissues from several patients. 1. pUC/Sau3a (length marker); 2. human renin probe (positive control); 3. tRNA (negative control); 4. rat kidney; 5. human kidney (20 µg); 6. sclera (35 µg); 7. retina (25 µg); 8. pigment epithelium-choroid/retina (45 µg).

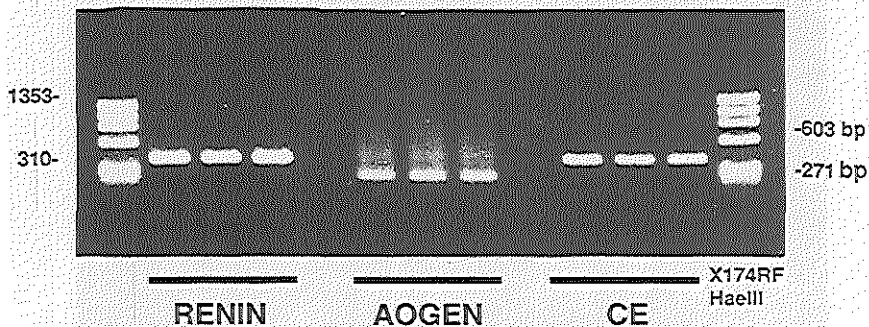


Figure 2. Detection of renin-, angiotensinogen- and ACE-mRNA expression by the polymerase chain reaction. Amplification products of renin (376 bp), angiotensinogen (217 bp) and ACE (388 bp) as obtained from renal total RNA (3 samples each). Ethidium-bromide stained 1.3 % agarose gel. X174RF vector HaeIII-fragments are used as length markers.

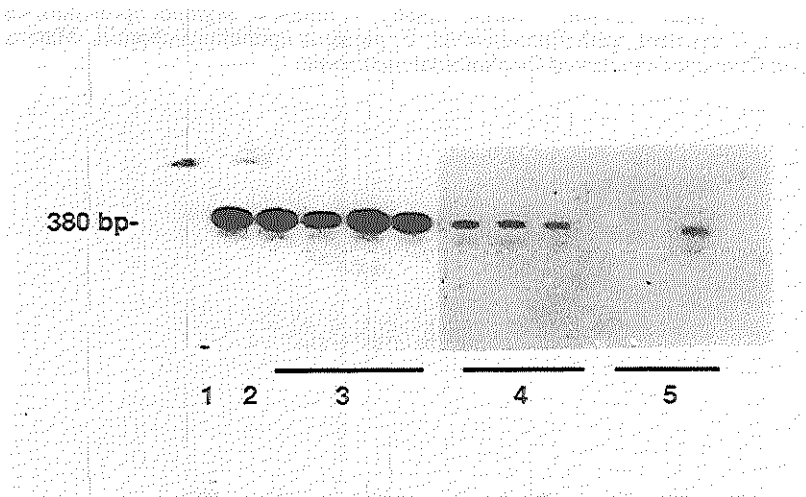


Figure 3. Demonstration of renin-mRNA expression in human eyes by the polymerase chain reaction. Southern blot of amplification products. 1. negative control; 2. human kidney; 3. pigment epithelium-choroid; 4. retina; 5. sclera (3-5: 3 samples each). Material was obtained from eyes enucleated for choroidal melanoma.

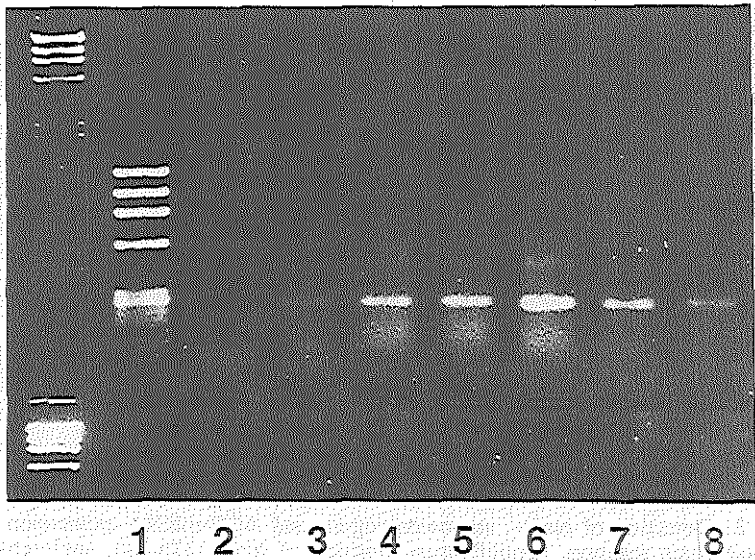


Figure 4. Demonstration of angiotensinogen-mRNA in human eyes by the polymerase chain reaction. Ethidium-bromide stained 1.3 % agarose gel. 1. X174RF HaeIII (length marker); 2. negative control; 3. human kidney; 4. retina; 5. pigment epithelium-choroid; 6. retina; 7. pigment epithelium-choroid; 8. pigment epithelium-choroid. Material was obtained from eyes enucleated for choroidal melanoma.

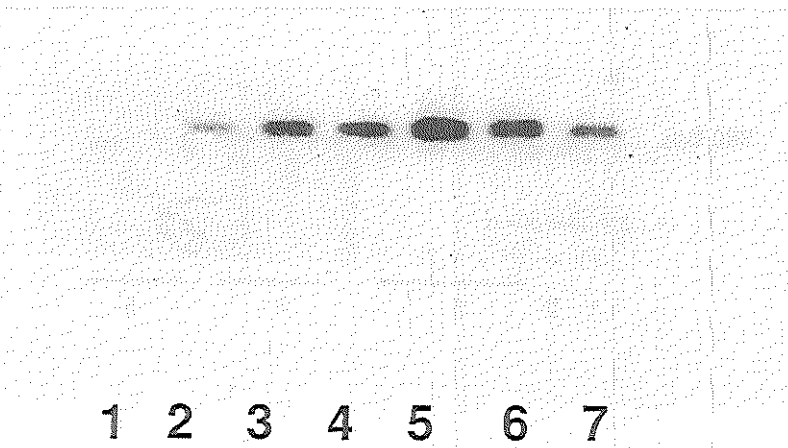


Figure 5. Demonstration of angiotensinogen-mRNA in human eyes by the polymerase chain reaction. Southern blot of amplification products. 1. negative control; 2. human kidney; 3. retina; 4. pigment epithelium-choroid; 5. retina; 6. pigment epithelium-choroid; 7. pigment epithelium-choroid. Material was obtained from eyes enucleated for choroidal melanoma.

Sensitivity could be improved by using the polymerase chain reaction. Using this method we were able to confirm that all components of the renin-angiotensin system are expressed in the kidney (Figure 2). Human kidney could therefore be used as a positive control in each assay. After amplification of cDNA by PCR all components of the renin-angiotensin system could be detected by Southern blotting in both retina and pigment epithelium-choroid (Figures 3, 5 and 6). The amplification products were checked for the predicted size by agarose gel electrophoresis (Figure 4). The retinal renin-mRNA levels were low but detectable (Figure 3), whereas in scleral tissue no positive signal for renin-mRNA could be found. The latter finding confirms the results from the RNase protection assay. In anterior uveal tract no amplification products could be detected. This may have been due to the fact that the mRNA in these samples was largely degraded, as observed when checking total RNA by gel electrophoresis.

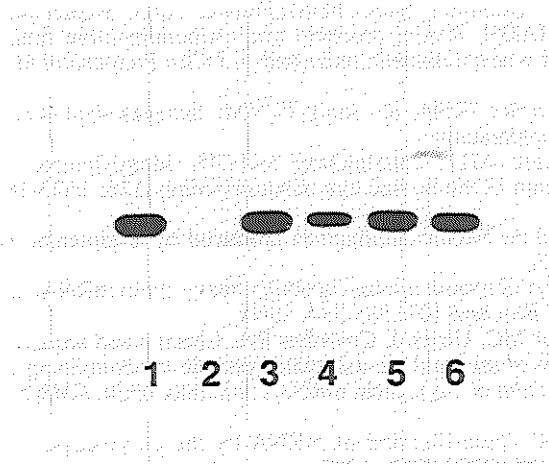


Figure 6. Demonstration of ACE-mRNA in human eyes by the polymerase chain reaction. Southern blot of amplification products. 1. human kidney; 2. negative control; 3. retina; 4. pigment epithelium-choroid; 5. pigment epithelium-choroid; 6. retina. Material was obtained from eyes enucleated for choroidal melanoma.

8.4 Discussion

The polymerase chain reaction is a powerful tool to amplify small amounts of DNA or mRNA for various molecular analyses (10). Using PCR we were able to demonstrate renin-, angiotensinogen- and ACE-mRNA expression in both retina and pigment epithelium-choroid, but not in sclera, of human eyes.

Renin mRNA could also be detected by the RNase protection assay. The pigment epithelium-choroid was found to contain the highest concentration of renin mRNA. Pooled samples of retinal RNA alone remained below the detection limits of this method.

These results strongly support the existence of an intra-ocular renin-angiotensin system. As gene-expression is highest in ocular parts, which are highly vascularized, the renin-angiotensin system may influence local blood supply and play a role in the process of neovascularization in such diseases as diabetic retinopathy. However, further work is needed to study whether changes in renin-mRNA expression do indeed occur in eyes with neovascularization.

From our results it can not be said in which cells the mRNA of the various components of the renin-angiotensin system is situated. *In situ* hybridization studies may provide an answer as to where exactly in the eye renin, angiotensinogen and ACE are synthesized.

8.5 References

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9. SUMMARY AND CONCLUSIONS

9.1 Aim of the thesis

Chapter 1

The renin-angiotensin system (RAS) is considered to be a major component in the maintenance of cardiovascular homeostasis in both the normal and the diseased state. Traditionally, the RAS has been viewed as an endocrine system. Circulating renin, released from the kidney, cleaves circulating liver-derived angiotensinogen (also known as renin substrate) to form angiotensin I (ANG I). ANG I is converted by endothelial-bound angiotensin converting enzyme (ACE) to angiotensin II (ANG II), a highly potent vasoconstrictor and aldosterone-releasing hormone. Initially, it was thought that blockers of the formation of ANG II (e.g. the ACE inhibitors, which were developed in the 1970s) would be effective for treating high renin-dependent forms of hypertension only. It was therefore an unexpected finding that ACE inhibitors were also effective in hypertensive patients with low or normal renin levels. Long-term treatment with ACE inhibitors does not seem to completely suppress the circulating RAS, as plasma ANG II and aldosterone tend to return to pretreatment levels. These findings have led to many speculations on the presence of so-called 'tissue' or 'local' renin-angiotensin systems, as opposed to the circulating RAS, and their role in the maintenance of high blood pressure. ACE inhibitors would then primarily exert their effect through blockade of these local systems.

Indeed, molecular biologic techniques have now convincingly shown that both renin and angiotensinogen genes are expressed in multiple tissues, including the heart, adrenal, brain, vessel wall, ovary and testis. An emerging concept is therefore that angiotensin may be produced locally in many tissues, independent of the circulating system, and that locally generated angiotensin plays an important role in influencing the function of these target organs. Locally generated angiotensin may even reach the circulating blood and thus contribute to the angiotensin levels found in plasma. To what extent generation of ANG I outside the blood stream and conversion of this locally generated ANG I to ANG II contribute to the circulating levels of ANG I and ANG II has not been investigated so far.

It was the first aim of the thesis to quantify the metabolism and production of ANG I and ANG II in various vascular beds in both humans and pigs. This was done by administrating constant infusions of ^{125}I -ANG I. Under steady state conditions the levels of ^{125}I -labeled and endogenous ANG I and ANG II were measured at both arterial and venous sampling sites. In addition, the plasma renin activity (PRA), the rate of ANG I formation in vitro, was

measured at each sampling site. The regional metabolism (including ANG I-II conversion) of ANG I and ANG II could be estimated from the arterial and venous levels of the radiolabeled peptides. The regional production could be calculated by comparing the regional extraction of the radiolabeled peptides with the actual levels of the endogenous ANG I and ANG II in arterial and venous plasma. In order to estimate the generation of ANG I outside the circulating blood, the contribution of PRA to the total regional ANG I production was calculated. The contribution of conversion of arterially delivered ANG I to the total regional ANG II production was also calculated.

The second aim of the thesis was to investigate the existence of a local RAS in the eye. Previously, a close correlation was found between the high plasma levels of prorenin, the inactive precursor of renin, and the presence of retinopathy, especially the proliferative type, in diabetic subjects. Prorenin may be involved in the local generation of angiotensin, whereas ANG II has been reported to have growth-promoting effects. The levels of the various components of the RAS were measured in human ocular fluids (both from eyes affected by proliferative diabetic retinopathy and from eyes of non-diabetic subjects) and in bovine and porcine ocular tissues. An attempt was also made to demonstrate the presence of renin-, angiotensinogen- and ACE-mRNA expression in human ocular tissues by the polymerase chain reaction.

9.2 Local metabolism and production of ANG I and ANG II in humans and pigs

Chapters 2, 3 and 4

During constant infusion of ^{125}I -ANG I extensive metabolism of the radiolabeled peptide occurred in all vascular beds that were studied, both in man and in the pig. In captopril-treated human subjects with essential hypertension, the regional extraction ratio of ^{125}I -ANG I was 47 ± 4 % (mean \pm SEM) across the forearm, 59 ± 3 % across the leg, 81 ± 1 % across the kidneys, and 96 ± 1 % across the hepatomesenteric vascular bed. These results were not different from those obtained for simultaneously infused unlabeled ANG I. Despite the rapid removal of arterially delivered ANG I, no difference was found between the venous and arterial levels of endogenous ANG I across the various vascular beds, with the exception of the liver where ANG I in the vein was 50 % lower than in the aorta. Thus, 50-90 % of endogenous ANG I in the veins appeared to be derived from regional de novo production. The blood transit time is 0.1-0.2 min in the limbs and in the kidneys and 0.3-0.5 min in the hepatomesenteric vascular bed. This is too short for PRA to account for the measured de novo ANG I production. Less than 20-30 % in the limbs and kidneys and approximately 60 % in the hepatomesenteric region of de novo produced venous ANG I could be accounted for by PRA.

These findings in humans could be confirmed in captopril-treated pigs and

were extended to untreated and furosemide-treated pigs. A kinetic model was introduced to estimate regional degradation and conversion of ANG I in the 3 groups of pigs. It was found that the metabolic degradation rate in the various vascular beds that were studied was the same for ^{125}I -ANG I and ANG I, but that the conversion rate was 2 times higher for ^{125}I -ANG I than for ANG I. ANG I appeared to be distributed over a compartment with a size corresponding to that of the extracellular fluid compartment (25 % of total body weight). After ACE inhibitor treatment, fractional ANG I metabolism (the fraction of arterially delivered ANG I that was metabolized during a single passage of blood) was 10 % in the lungs (conversion 4%), as compared to 56 % in the combined systemic vascular beds (conversion 1 %). Fractional ANG I metabolism during ACE inhibition was 93 % in the kidney, 50-70 % in myocardium, skeletal muscle, head and skin, and 21 % in the left and 14 % in the right cardiac cavity. Without ACE inhibition, fractional ANG I metabolism was 29 % in the lungs (conversion 25 %), 49 % in the combined systemic vascular beds (conversion 10 %), 38 % in the left cardiac cavity (conversion 11 %) and 14 % in the right cardiac cavity (conversion 0 %). These data show that in the systemic vascular beds, but not in the lungs or cardiac cavities, decreased conversion is accompanied by increased degradation.

Total regional ANG I and ANG II production rates in the various porcine systemic vascular beds were calculated on the basis of the so-called 'venous equilibrium' model. *De novo* ANG I production in the combined systemic vascular beds was increased during treatment with captopril or the diuretic furosemide, as compared to untreated animals. ANG II production in the combined systemic vascular beds was greater than in the lungs, demonstrating the importance of extrapulmonary ANG I-II conversion. ACE inhibition did not alter the ANG II production in the lungs or the combined systemic vascular beds. Apparently, reduced ANG I-II conversion was compensated for by increased ANG I production. Furosemide increased the ANG II production in all vascular beds that were studied.

In the lungs, ANG I production could fully be accounted for by circulating PRA, and ANG II production could be attributed to the intrapulmonary conversion of circulating ANG I. In contrast, in the other vascular beds we studied a substantial fraction of ANG I production could not be accounted for by circulating PRA, assuming a transit time of 10 sec (myocardium > 25 %, head, skin and skeletal muscle > 80 %, kidney > 95 %). The local ANG I production in all vascular beds correlated with the level of PRA. Part of the local ANG II production in these regions could not be attributed to regional conversion of circulating ANG I (myocardium > 30 %, head, skin and skeletal muscle > 50 %, kidney > 80 %). These results indicate that indeed production of ANG I at tissue sites contributes to its circulating level, and that some circulating ANG II may not be derived from circulating ANG I.

9.3 The ocular renin-angiotensin system

Chapters 5, 6, 7 and 8

Renin, prorenin, angiotensinogen, ANG I and ANG II were present in aqueous, vitreous and subretinal fluid obtained from human eyes during eye surgery. In many samples the renin levels were close to the detection limit of the assay. Large differences (up to factor of 10) in angiotensinogen, ANG I and ANG II concentrations were found between the various ocular fluid types, but these were reflected by similarly large differences in ocular fluid albumin concentrations, indicating that angiotensinogen, ANG I and ANG II in human ocular fluid samples are most likely plasma-derived. In contrast, the prorenin levels in human ocular fluid were up to 100 times higher than expected on the basis of the plasma protein content of these fluids. Moreover, there was little difference in prorenin concentrations between samples with a low and high plasma protein content. Relative to albumin, prorenin was higher in vitreous fluid from eyes with proliferative diabetic retinopathy complicated by traction retinal detachment than in vitreous fluid from eyes of non-diabetic subjects with spontaneous retinal detachment. These findings suggest that an intraocular RAS does exist and that it may play a role in the development of retinal neovascularization in diabetes mellitus.

To further substantiate the evidence for an intraocular RAS, the ANG I-generating activity was measured in bovine aqueous and vitreous fluid and in extracts of bovine retina, pigment epithelium-chorioid and anterior uveal tract before and after subjecting these ocular fluids and extracts to procedures known to convert prorenin to renin. Renin in the ocular samples could be separated from nonrenin acid protease by α -casein-Sepharose affinity column chromatography at pH 3.5; renin did not bind to the column, whereas acid protease did. Renin was further identified by its relatively high pH optimum (6.5-7.0) for ANG I generation, its complete inhibition with specific renin antiserum, and its high affinity for specific renin inhibitors. More than 75 % of ANG I-generating activity of the bovine ocular samples consisted of renin. Approximately 90 % or more of total renin (renin + prorenin) in aqueous humor, vitreous fluid, and ocular tissue could not be explained by trapped plasma. In vitreous fluid, prorenin comprised 99 % of the total renin, in retina 81 %, and in pigment epithelium-choroid and anterior uveal tract less than 50 %. Prorenin in ocular fluid showed a concentration gradient, posterior vitreous > anterior vitreous > aqueous humor, suggesting that the main source of prorenin in ocular fluid was in the posterior eye.

During a 2-hour intravenous infusion of ^{125}I -ANG I in the rabbit less than 1 % of intact plasma ^{125}I -ANG I reached the vitreous fluid compartment, showing that, under normal conditions, plasma angiotensins pass the blood-retinal-barrier only in very limited amounts. Measurements of angiotensins in porcine ocular tissues showed that both ANG I and ANG II were present in retina, pigment epithelium-choroid and anterior uveal tract in concentrations

much too high to be explained by contamination with plasma only. Low to undetectable angiotensin levels were found in porcine vitreous fluid. The ANG II/ANG I ratio was 3-5 fold higher in the ocular tissues than in plasma or vitreous fluid. Finally, using the polymerase chain reaction it was possible to demonstrate the expression of renin-, angiotensinogen- and ACE-mRNA in human eyes. The highest expression was found in the pigment epithelium-choroid and retina.

9.4 General conclusions

In all systemic vascular beds that were studied, both in man and in the pig, venous ANG I could not be accounted for by arterial delivery of ANG I, by the action of circulating renin on circulating angiotensinogen ('PRA') or by the sum of these two. A high percentage of plasma ANG I must therefore have been produced locally, i.e. not in circulating plasma. Our calculations of the local production rates of ANG I show indeed that in myocardium, head, skin, skeletal muscle and kidneys, but not in lungs, the contribution of PRA (assuming a transit time of 10 sec in the various systemic vascular beds and 3 sec in the lungs) to total ANG I production is considerably less than 100 %. For instance, in the kidneys only 2 % of the total ANG I production could be attributed to PRA. These calculations were based on the local extraction of ANG I and the arterial and venous ANG I levels; the venous ANG I levels were considered to be representative for the tissue levels of ANG I. In fact, the tissue levels may be higher than the ANG I levels in the venous effluent. Probably, the capillaries are the main site where ANG I is removed from the circulation by diffusion into the interstitium and peptidase attack. This is also the site where ANG I from the interstitium equilibrates with circulating ANG I. The blood transit time in the capillary beds is 1-3 sec. Thus, our estimated transit time of 10 sec in most systemic vascular beds is probably too high. Therefore our results may have underestimated the tissue ANG I production and at the same time they may have overestimated the contribution of PRA. However, this only strengthens our conclusion that ANG I is produced locally at tissue sites. The ANG I production rates in the various tissues correlated with PRA, suggesting that most of the renin responsible for local ANG I production is kidney-derived. Indeed, the plasma ANG I levels in nephrectomized subjects are known to be low and, in addition, the release of ANG I by isolated tissues perfused with non-renin containing perfusion fluids is also low or even undetectable.

Following captopril, the regional conversion of ANG I was markedly decreased. In the systemic vascular beds, but not in the lungs or the cardiac cavities, this decrease in conversion was accompanied by an increase in degradation. These findings suggest that there may be some degree of compartmentalization of the processes of conversion and degradation in the

sense that it is mainly the blood-borne ANG I that is subject to conversion whereas interstitial ANG I is mainly subject to degradation. The lungs are known to contain relatively little interstitial fluid. The size of the distribution volume of ANG I equalled the size of the extracellular fluid volume, suggesting that circulating ANG I is limited to the circulation and the interstitial fluid only. In other words: plasma ANG I may equilibrate with interstitial ANG I. This does not exclude ANG I production at tissue sites not equilibrating with circulating plasma, for instance intracellular ANG I production sites. Possibly, kidney-derived renin is limited to the extracellular fluid and cleaves angiotensinogen in plasma or interstitial fluid only, whereas locally synthesized (pro)renin cleaves (locally synthesized) angiotensinogen intracellularly.

Not only ANG I, but also ANG II appeared to be produced locally. Regional conversion of arterially delivered ANG I was not sufficient to account for the total regional production of ANG II. ANG II may originate from ANG I produced at tissue sites, although alternative pathways for ANG II generation (e.g. direct cleavage from angiotensinogen) have also been described.

The above assumptions concerning tissue ANG I and ANG II production could be tested by examining an as yet unexplored RAS, in the eye. The ocular tissue (pro)renin and angiotensin levels were found to be far too high to be explained by trapped plasma only. This confirms that plasma renin is not the only contributor to tissue angiotensin production, at least in the eye. In contrast, the levels of angiotensinogen, ANG I and ANG II in the various ocular fluids that could be obtained (both from humans and from animals) largely correlated with the albumin levels in these fluids. Angiotensinogen, ANG I and ANG II in ocular fluid are therefore most likely derived from plasma. This was not the case with prorenin. Our finding that prorenin was higher in vitreous from eyes affected by proliferative diabetic retinopathy than in vitreous from eyes of non-diabetic patients suggests a role for the ocular RAS in the development of retinal neovascularization. This would be compatible with our measurements of renin-, angiotensinogen- and ACE-mRNA in human ocular tissues, demonstrating the highest expression in the most vascularized parts of the eye. It seems therefore that both ANG I and ANG II are synthesized in ocular tissues, but that, in contrast with prorenin, little or no intraocularly synthesized ANG I and ANG II is released into the ocular fluids. This suggests an autocrine or paracrine function for the intraocular RAS. It cannot be concluded from our data if there is any diffusion of ocular angiotensins or prorenin into the plasma compartment. However, other organs known to contain a local RAS (such as ovary and testis) have been reported to secrete prorenin into the circulation.

Based upon differences in vitreous prorenin levels found between eyes with and without proliferative retinopathy long-term clinical studies should be performed to examine whether ACE inhibitor treatment has a favourable effect on the development and progression of retinal neovascularization in diabetic subjects.

10. SAMENVATTING EN CONCLUSIES

10.1 Doel van het onderzoek

Hoofdstuk 1

Het renine-angiotensine systeem (RAS) speelt een belangrijke rol bij de handhaving van de cardiovasculaire homeostasis. Traditioneel werd het RAS altijd gezien als een endocrien systeem. Circulerend renine, aan het bloed afgegeven door de nieren, reageert met angiotensinogeen (ook wel aangeduid als renine substraat) uit de lever, waarbij angiotensine I (ANG I) gevormd wordt. ANG I wordt door endotheel-gebonden angiotensin convertende enzyme (ACE) omgezet in angiotensine II (ANG II), een zeer potente vasoconstrictor die tevens de vrijzetting van aldosteron stimuleert. Op grond hiervan leek het logisch om aan te nemen dat remmers van de vorming van ANG II (zoals bijvoorbeeld de ACE remmers, die ontwikkeld werden in de zeventiger jaren) vooral werkzaam zouden zijn bij die vormen van hypertensie waarbij hoge plasma renine spiegels de oorzaak zijn van de hoge bloeddruk. Onverwacht bleken ACE remmers echter ook bij hypertensieve patienten met een lage of normale renine spiegel de bloeddruk te verlagen. Recent is zelfs gebleken dat langdurige behandeling met ACE remmers het circulerende RAS niet volledig onderdrukt, aangezien de plasma ANG II en aldosteron spiegels weer terugkomen op hun uitgangswaarden (terwijl de bloeddruk verlaagd blijft). Deze bevindingen hebben geleid tot talloze speculaties over de aanwezigheid van zgn. 'weefsel' of 'lokale' renine-angiotensine systemen, in tegenstelling dus tot het circulerende RAS, welke een rol zouden spelen bij de handhaving van de hoge bloeddruk. ACE remmers zouden dan met name hun effect bewerkstelligen via blokkade van deze lokale systemen. Gebruik makend van moleculair-biologische technieken kon men inderdaad aantonen dat renine- en angiotensinogeen-mRNA aanwezig zijn in een groot aantal weefsels en organen, o.m. het hart, de bijnier, de hersenen, de vaatwand, het ovarium en de testis. Men begint daarom meer en meer aan te nemen dat angiotensine lokaal geproduceerd wordt op vele plaatsen in het lichaam, onafhankelijk van het circulerende systeem, en dat dit lokaal geproduceerde angiotensine ook lokaal een functie heeft. Wellicht zou zelfs lokaal geproduceerd angiotensine de circulatie kunnen bereiken en zo bijdragen aan de in plasma gevonden angiotensine spiegels. Of produktie van ANG I buiten de circulatie en omzetting van dit lokaal gevormde ANG I in ANG II daadwerkelijk een rol spelen bij de handhaving van de plasma spiegels van ANG I en ANG II is nooit uitgezocht.

Het eerste doel van het onderzoek was om in diverse vaatbedden, zowel bij mensen als bij varkens, kwantitatief het metabolisme en de produktie van ANG I en ANG II vast te stellen. Hiertoe werd gebruik gemaakt van constante infusen van ^{125}I -ANG I. Na het bereiken van een steady state werden de spiegels van ^{125}I -gelabeld en endogeen ANG I en ANG II gemeten in diverse arteriële en veneuze monsters. Tevens werd de plasma renine activiteit (PRA), een maat voor de vorming van ANG I *in vitro*, op elke bloedafnameplaats gemeten. Het regionale metabolisme (inclusief ANG I-II conversie) van ANG I en ANG II kon worden afgeleid uit de arteriële en veneuze spiegels van de gelabelde peptiden. De regionale produktie kon worden berekend door de regionale extractie van de gelabelde peptiden te vergelijken met de daadwerkelijk gemeten endogene spiegels van ANG I en ANG II in arterieel en veneus plasma. Om een schatting te kunnen maken van de ANG I generatie buiten het circulerende plasma werd de bijdrage van de PRA aan de totale ANG I produktie uitgerekend. Tevens werd de bijdrage van conversie van arterieel aangeleverd ANG I aan de totale ANG II produktie berekend.

Het tweede doel van het onderzoek was om het bestaan van een lokaal RAS in het oog aannemelijk te maken. Eerder werd namelijk een sterke samenhang gevonden tussen de hoge plasma spiegels van prorenine, de inactieve precursor van renine, en de aanwezigheid van retinopathie, met name het proliferatieve type, bij patiënten met diabetes mellitus. Prorenine zou een rol kunnen spelen bij de lokale angiotensine vorming, terwijl van ANG II wordt verondersteld dat het ook groei-stimulerend zou kunnen werken. De spiegels van de diverse componenten van het RAS werden gemeten in humane oogvloeistoffen (zowel van ogen van patiënten met proliferatieve diabetische retinopathie als van ogen van niet-diabetische patiënten) en in oogweefsels van koeien en varkens. Tevens werd, met behulp van de polymerase ketting reactie, getracht de aanwezigheid van renine-, angiotensinogeen- en ACE-mRNA in humaan oogweefsel aan te tonen.

10.2 Lokaal metabolisme en lokale produktie van ANG I en ANG II bij de mens en bij het varken

Hoofdstukken 2, 3 en 4

Tijdens constante infusie van ^{125}I -ANG I bleek het gelabelde peptide in alle vaatbedden die werden bestudeerd, zowel bij de mens als bij het varken, voor een belangrijk deel te worden afgebroken. In patiënten met essentiële hypertensie die met captopril behandeld werden, bedroeg de regionale extractie ratio van ^{125}I -ANG I $47 \pm 4\%$ (gemiddelde \pm SEM) in de onderarm, $59 \pm 3\%$ in het been, $81 \pm 1\%$ in de nieren, en $96 \pm 1\%$ in het hepatomesenteriale vaatbed. De resultaten waren niet verschillend van die voor gelijktijdig geïnfuseerd ongelabeld ANG I. Ondanks de hoge mate van extractie van arterieel aangevoerd ANG I, waren er geen verschillen tussen de veneuze en arteriële spiegels van endogeen ANG I in de onderzochte vaatbedden, met

uitzondering van de lever, waar veneus ANG I 50 % lager was dan arterieel ANG I. Kennelijk is 50-90 % van veneus endogeen ANG I afkomstig van regionale de novo productie. De bloed doorstromingstijd is 0.1-0.2 min in arm, been en nieren en 0.3-0.5 min in het hepatomesenteriale vaatbed. Deze tijd is veel te kort om de volledige de novo productie van ANG I aan de PRA toe te kunnen schrijven. Minder dan 20-30 % in de ledematen en nieren en rond de 60 % in het hepatomesenteriale vaatbed van de novo geproduceerd veneus ANG I kon worden toegeschreven aan de PRA.

Deze bevindingen bij de mens konden worden bevestigd in captopril-behandelde varkens, en werden uitgebreid met metingen in onbehandelde en furosemide-behandelde varkens. Een farmacokinetisch model werd geïntroduceerd om de regionale degradatie en conversie van ANG I in de 3 groepen varkens te schatten. De degradatie van ¹²⁵I-ANG I en ANG I bleek niet te verschillen, maar de conversie van ¹²⁵I-ANG I verliep ongeveer 2 keer zo snel als die van ANG I. ANG I verdeelde zich over een volume dat wat grootte betreft overeen kwam met het extracellulaire vloeistof compartiment (25 % van het totale lichaamsgewicht). Na voorbehandeling met captopril, bedroeg het fractionele metabolisme van ANG I (de fractie van arterieel aangevoerd ANG I die tijdens de passage van een vaatbed afgebroken wordt) 10 % in de longen (waarvan 4 % door conversie) en 56 % in de gecombineerde systemische vaatbedden (1 % door conversie). Gedurende ACE remming was het fractionele metabolisme van ANG I 93 % in de nier, 50-70 % in myocard, skeletspierweefsel, hoofd en huid, en 21 % in de linkerhartholte en 14 % in de rechter hartholte. Zonder ACE remming bedroeg het fractionele metabolisme van ANG I 29 % in de longen (25 % door conversie), 49 % in de gecombineerde systemische vaatbedden (10 % door conversie), 38 % in de linker hartholte (11 % door conversie) en 14 % in de rechter hartholte (0 % door conversie). Deze getallen tonen aan dat in de systemische vaatbedden, maar niet in de longen of hartholtes, afgenomen conversie gepaard gaat met toegenomen degradatie.

De totale regionale ANG I en ANG II productie in diverse systemische vaatbedden bij het varken werd berekend op grond van het zogenaamde 'veneuze equilibrium' model. De novo ANG I productie in de gecombineerde systemische vaatbedden was toegenomen tijdens behandeling met de ACE remmer captopril en het diureticum furosemide, in vergelijking met de onbehandelde varkens. De ANG II productie was groter in de gecombineerde systemische vaatbedden dan in de longen, hetgeen wijst op het belang van ANG I-II conversie buiten de longen. De ANG II productie in de longen en de gecombineerde systemische vaatbedden veranderde niet significant na ACE remming. Kennelijk werd de afgenomen ANG I-II conversie gecompenseerd door toegenomen ANG I productie. Onder furosemide nam de ANG II productie in alle bestudeerde vaatbedden toe.

In de longen kon de ANG I productie volledig worden toegeschreven aan de PRA in het circulerende plasma. Tevens bleek de ANG II productie in de

longen volledig verklaard te kunnen worden uit intrapulmonale conversie van arterieel aangevoerd ANG I. In alle andere vaatbedden die werden bestudeerd was dit niet het geval: een belangrijk deel van de ANG I produktie kon daar niet worden verklaard door aan te nemen dat alleen de PRA in het circulerende plasma (uitgaande van een doorstromingstijd van 10 sec) verantwoordelijk is voor de lokale ANG I produktie (hart > 25 %, hoofd, huid en skeletspierweefsel > 80 %, nier > 95 %). Wel correleerde de lokale ANG I produktie in alle bestudeerde vaatbedden met de gemeten PRA. Een deel van de ANG II produktie in deze vaatbedden kon tevens niet worden verklaard uit de lokale conversie van circulerend ANG I (hart > 30 %, hoofd, huid en skeletspierweefsel > 50 %, nier > 80 %). Deze resultaten tonen aan dat lokale produktie van ANG I inderdaad bijdraagt aan de spiegels van circulerend ANG I, en dat een deel van het circulerende ANG II niet van circulerend ANG I afkomstig is.

10.3 Het renine-angiotensine systeem in het oog

Hoofdstukken 5, 6, 7 en 8

Renine, prorenine, angiotensinogeen, ANG I en ANG II waren aanwezig in kamerwater, glasvocht en subretinaalvocht verkregen uit menselijke ogen tijdens oogoperaties. In veel monsters lagen de renine spiegels dicht bij de detectie limiet van de meetmethode. Grote verschillen (soms wel 10-voudig) in de angiotensinogeen, ANG I en ANG II concentraties werden gevonden tussen de verschillende types oogvloeistof, maar deze gingen vergezeld van even grote verschillen in de albumine concentraties in deze oogvloeistoffen, hetgeen er op wijst dat angiotensinogeen, ANG I en ANG II in kamerwater, glasvocht en subretinaal vocht waarschijnlijk afkomstig zijn uit plasma. In tegenstelling hiermee, bleken de prorenine spiegels in humane oogvloeistoffen soms wel 100 keer hoger te liggen dan je zou verwachten op grond van de plasma eiwit concentraties in deze vloeistoffen. Bovendien verschilden de prorenine concentraties nauwelijks tussen monsters met een hoog gehalte aan plasma eiwitten en monsters met een laag gehalte aan plasma eiwitten. Uitgedrukt per g albumine was de prorenine concentratie in glasvocht afkomstig van ogen van patiënten met proliferatieve diabetische retinopathie en een tractie retinaloslating hoger dan in glasvocht van ogen van niet-diabetische patiënten met een spontane retinaloslating. Deze bevindingen suggereren dat er inderdaad sprake is van een intraoculair RAS, dat wellicht een rol speelt bij de ontwikkeling van retinale vaatnieuwvorming in patiënten met diabetes mellitus.

Om het bestaan van een intraoculair RAS verder te substantiëren, werd de ANG I-vormende activiteit in kamerwater en glasvocht en in extracten van retina, pigment epithelium-choroidea en iris-corpora ciliare van runderen gemeten, zowel voor als na blootstelling van deze oogvochten en extracten aan procedures die kunnen leiden tot de omzetting van prorenine in renine. Renine in de oogmonsters kon worden gescheiden van zure proteasen door gebruik te

maken van α -caseine-Sepharose affiniteitskolomchromatografie bij pH 3.5; renine werd niet door deze kolom gebonden, maar zure proteasen wel. Renine kon verder worden geïdentificeerd door het relatief hoge pH optimum (6.5-7.0) voor de ANG I vorming, en het feit dat de ANG I-vormende activiteit volledig kon worden geremd door gebruik te maken van hetzij een specifiek renine antilichaam, hetzij specifieke renine remmers. Meer dan 75 % van de ANG I-vormende activiteit van de runderoogmonsters was te wijten aan renine. Ongeveer 90 % van de totale hoeveelheid renine (renine + prorenine) in kamerwater, glasvocht en oogweefsels kon niet worden verklaard door het hierin aanwezige plasma. In glasvocht bestond 99 % van de totale hoeveelheid renine uit prorenine, in de retina 81 %, en in pigment epithelium-choroidea en iris-corpora ciliaria minder dan 50 %. Er was een concentratiegradiënt voor prorenine aanwezig in de onderzochte oogvloeistoffen: glasvocht_{achterin oog} > glasvocht_{voorin oog} > kamerwater. Dit lijkt erop te wijzen dat de belangrijkste bron van prorenine in oogvloeistoffen zich achterin het oog bevindt.

Gedurende een 2-uur durend infuus van ^{125}I -ANG I bij het konijn, bereikte minder dan 1 % van intact plasma ^{125}I -ANG I het glasvocht, suggererend dat er, onder normale condities, nauwelijks lekkage van plasma angiotensines door de bloed-retina-barrière plaatsvindt. Metingen van angiotensines in oogweefsels van varkens toonden aan dat zowel ANG I als ANG II aanwezig waren in retina, pigment epithelium-choroidea en iris-corpora ciliaria in concentraties die veel te hoog waren om verklaard te kunnen worden uit contaminatie met plasma. Lage tot onmeetbare angiotensine concentraties werden gevonden in varkensglasvocht. De ANG II/ANG I ratio was 3-5 keer zo hoog in de oogweefsels als in plasma of glasvocht. Tenslotte bleek het, gebruik makend van de polymerase ketting reactie, mogelijk om de expressie van renine-, angiotensinogeen- en ACE-mRNA in humaan oogweefsel aan te tonen. De hoogste mate van expressie werd aangetroffen in pigment epithelium-choroidea en retina.

10.4 Conclusies

In alle systemische vaatbedden die werden onderzocht, zowel bij mensen als bij varkens, kon een zeer groot deel van veneus ANG I niet worden verklaard door hetzij arteriële aanvoer van ANG I, hetzij de reactie van circulerend renine met circulerend angiotensinogeen ('PRA'), noch door de combinatie van deze twee bronnen. Een groot gedeelte van plasma ANG I wordt dus waarschijnlijk lokaal (d.w.z. niet in circulerend plasma) geproduceerd. Inderdaad bleek uit berekeningen van de lokale produktie van ANG I dat in hart, hoofd, huid, skeletspierweefsel en nieren, maar niet in longen, de bijdrage van de PRA (gebaseerd op een doorstromingstijd van 10 sec in de verschillende systemische vaatbedden en 3 sec in de longen) aan de totale produktie van ANG I beduidend minder is dan 100 %. In de nieren kon

bijvoorbeeld maar 2 % van de totale ANG I productie geweten worden aan de PRA. Bovenstaande berekeningen zijn gebaseerd op de lokale extractie van ANG I en de arteriële en veneuze ANG I spiegels; aangenomen werd dat de veneuze ANG I spiegels een afspiegeling zijn van de spiegels op weefselniveau. In werkelijkheid zijn de weefsel ANG I spiegels misschien hoger dan de ANG I spiegels in het veneuze effluent. Waarschijnlijk zijn het met name de capillairen waar ANG I door diffusie naar het interstitium en door afbraak door peptidases uit de circulatie verwijderd wordt. Hier ook staat ANG I uit het interstitium in evenwicht met plasma ANG I. De doorstromingstijd van de capillaire vaatbedden is slechts 1-3 sec. Daarom leidt het gebruik van een doorstromingstijd van 10 sec voor de berekening van de bijdrage van de PRA in de meeste systemische vaatbedden waarschijnlijk tot een te hoge bijdrage van de PRA. Onze resultaten zullen dus tegelijkertijd de weefsel ANG I productie iets onderschatten en het aandeel van de PRA iets overschatten. Zodoende wordt echter onze conclusie dat ANG I lokaal geproduceerd des te meer ondersteund. De lokale ANG I productie correleerde met de PRA, wat er op zou kunnen wijzen dat het renine dat verantwoordelijk is voor deze lokale productie afkomstig is uit de nier. Deze theorie wordt ondersteund door recente bevindingen dat a) de ANG I plasma spiegels in genefrectomeerde patiënten zeer laag zijn en b) de vrijzetting van ANG I uit geïsoleerde weefsels die geperfuseerd worden met een vloeistof die geen renine bevat ook zeer laag (vaak zelfs niet aantoonbaar) is.

Na voorbehandeling met captopril nam de lokale conversie van ANG I significant af. In de systemische vaatbedden, maar niet in de longen of hartholtes, ging deze afname in conversie gepaard met een toename in degradatie. Deze bevindingen suggereren dat er een zekere mate van compartimentalisatie bestaat voor conversie en degradatie: ANG I uit de circulatie staat vooral bloot aan conversie, terwijl ANG I in het interstitium met name door degraderende enzymen wordt afgebroken. De longen bevatten relatief weinig interstitieel vocht. De grootte van het verdelingsvolume van ANG I kwam overeen met de grootte van het extracellulaire vloeistof volume. Kennelijk is circulerend ANG I beperkt tot de bloedbaan en het interstitium. Met andere woorden: plasma ANG I equilibreert met interstitieel ANG I. Dit sluit niet uit dat er ook nog elders ANG I productie plaatsvindt, in een compartiment dat niet uitwisselt met het circulerende plasma (bijvoorbeeld een intracellulair compartiment). Mogelijk beperkt renaal renine zich tot extracellulair vocht en reageert het alleen met angiotensinogeen in plasma of in het interstitium, terwijl lokaal gesynthetiseerd (pro)renine slechts intracellulair ANG I vormt uit (lokaal gesynthetiseerd) angiotensinogeen.

Niet alleen ANG I, maar ook ANG II bleek lokaal geproduceerd te worden. De regionale conversie van arterieel aangevoerd ANG I was niet voldoende om de totale regionale productie van ANG II te verklaren. ANG II kan afkomstig zijn van in de weefsels geproduceerd ANG I, maar er zijn ook andere manieren waarop ANG II gevormd kan worden (bijvoorbeeld directe

afsplitsing van angiotensinogeen).

Bovenstaande aannames betreffende lokale ANG I en ANG II productie konden worden onderzocht op hun juistheid tijdens een onderzoek naar een tot dusverre onbekend lokaal RAS, in het oog. De spiegels van (pro)renine en angiotensine in oogweefsels waren veel te hoog om verklaard te worden uit hierin aanwezig plasma. Dit bevestigt dat plasma renine niet als enige bijdraagt aan de lokale productie van angiotensines, althans in het oog. In tegenstelling hiermee, bleek dat de spiegels van angiotensinogeen, ANG I en ANG II in diverse oogvochten (subretinaal vocht, glasvocht, kamerwater) grotendeels correleerden met de albumine spiegels in deze oogvochten. Angiotensinogeen, ANG I en ANG II in oogvocht zijn dus waarschijnlijk afkomstig uit plasma. Dit gold echter niet voor prorenine. Onze bevinding dat prorenine hoger was in glasvocht uit ogen van diabetische patiënten met proliferatieve retinopathie dan in glasvocht uit ogen van niet-diabetische patiënten suggereert dat het oculaire RAS een rol speelt bij de ontwikkeling van retinale neovascularisatie. De metingen van renine-, angiotensinogeen, en ACE-mRNA in humane oogweefsels ondersteunden deze theorie, aangezien de hoogste expressie gevonden werd in die delen van het oog die het meest doorbloed zijn. Kennelijk worden ANG I en ANG II wel gesynthetiseerd in oogweefsel, maar lekt er hiervan weinig naar glasvocht en/of kamerwater. Onze bevindingen suggereren daarom een autocriene of paracriene functie voor het RAS in het oog. Of er angiotensines of prorenine vanuit het oog diffunderen naar plasma kan uit onze gegevens niet worden vastgesteld. Prorenine lijkt in ieder geval wél naar oogvocht weg te lekken. Ook in het ovarium is gebleken dat lokaal geproduceerd prorenine niet alleen terechtkomt in (onder meer) follikelvocht, maar ook wordt afgegeven aan de bloedbaan.

Gebaseerd op de verschillen in prorenine concentratie in glasvocht tussen ogen met en zonder proliferatieve retinopathie zou het aanbeveling verdienen om de effecten van ACE remmers op het ontstaan respectievelijk de progressie van retinale neovascularisatie bij patiënten met diabetes mellitus nader te onderzoeken.

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