

**MATRIX METALLOPROTEINASE ACTIVATION IN HEART
FAILURE DUE TO VOLUME-OVERLOAD AND THE EFFECT
OF AT₁ RECEPTOR BLOCKADE**

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In Partial Fulfillment of the Requirement
For the Degree Of:**

MASTERS OF SCIENCE IN PHYSIOLOGY

By

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**Matrix Metalloproteinase Activation in Heart Failure Due to Volume-Overload and
the Effect of AT1 Receptor Blockade**

BY

Baljit S. Walia

**A Practicum submitted to the Faculty of Graduate Studies of The University
of Manitoba in partial fulfillment of the requirements of the degree**

of

MASTER OF SCIENCE

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TO MY PARENTS

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I. ABSTRACT

Matrix remodeling is a tightly regulated process and represents a steady-state balance between extracellular matrix removal and matrix synthesis. The former is controlled by matrix degrading enzymes and their inhibitors and is involved in normal tissue function and development. A breakdown in regulation of these control processes may contribute to pathological degradation of connective tissue in various diseased conditions and the development of congestive heart failure subsequent to altered ventricular geometry. Zymogens that mediate the removal of connective tissue are the members of matrix metalloproteinase family.

In the present study, we have investigated the role of matrix metalloproteinases (MMP) during the development of volume overload induced moderate heart failure. Furthermore, we have tested whether the renin angiotensin system (RAS) or specifically the activation of AT₁ receptors is involved in the progression to cardiac hypertrophy and heart failure caused by volume overload. This was achieved by the use of losartan, an AT₁ receptor antagonist.

Surgically induced aorto-venous fistula (AV shunt: AVF) is the model of volume overloaded heart failure used in these investigations. Animals were divided into three groups; a sham-operated control group, an AV shunt group and an AV shunt group treated with losartan. Hemodynamic and anatomical studies were performed at 8 weeks after the shunt surgery while all other studies were performed at 1 and 8 week post AV shunt induction. To examine whether MMP and MMP inhibitory protein (TIMP) activities are altered in this model of heart failure, zymography and reverse zymography were performed respectively using ventricular tissue samples. Immunoreactive MMP

protein abundance was determined by Western blot analysis. To evaluate the role of angiotensin II, AT₁ antagonism was applied to experimental animals (losartan:40 mg/kg/day). The hallmarks of heart failure are significant increases in the LVEDP, right atrial pressure (RAP), and lung congestion. Losartan treatment for 8 weeks was associated with improved hemodynamic performance. Interstitial collagen deposition as determined by cardiac 4-hydroxyproline content, was significantly decreased in ventricular samples (LV and RV) from the experimental group (both 1 and 8 week), compared to respective control groups. Losartan treatment was associated with significant improvement of cardiac left ventricular (LV) collagen concentration at both time points studied. However, 1 week losartan treatment did not have a significant effect on right ventricular (RV) collagen content while 8 weeks of losartan treatment was associated with significant normalization of collagen content. Western blot analysis did not reveal any significant alterations in steady-state myocardial MMP (MMP-1, MMP-2, MMP-9) protein content at either 1 or 8 week in both left and right ventricles. Gelatinolytic analysis of left ventricular tissue samples revealed significant increases in MMP-1, MMP-2 and MMP-9 activities in AV shunt group compared to controls at 1 week after the creation of shunt while for 1 week RV, MMP-1 and MMP-9 activities were increased in shunt group vs controls. 8 week LV and RV exhibited a significant increase in MMP-1 activity whereas MMP-2 and MMP-9 activities were comparable in experimental vs. control group. Losartan treatment was associated with significant normalization of MMP-9 activity in 1 week LV sample but had no effect on LV MMP-1 or MMP-2 activities and RV MMP (MMP-1, MMP-9) activity. However, 8 week Losartan treatment was associated with improved MMP-1 activity in both right and left

ventricular sample. Reverse zymography did not reveal any detectable changes in myocardial TIMP-1 activity among the three groups studied. These results indicate that; i) AVF rats with volume overload exhibit both right- and left-sided heart failure, ii) AT₁ antagonism attenuates to a large extent, the development of heart failure and iii) these changes are paralleled by a normalization in collagen content as well as myocardial MMP activities. It is suggested that cardiac MMP activation may play a role in cardiac matrix remodeling associated with heart failure due to volume overload.

II. INTRODUCTION

Ventricular dilatation is an important event in the progression of congestive heart failure (CHF). Alterations in cardiac loading typically results in pressure-induced, concentric hypertrophic growth and/or volume-induced, eccentric hypertrophic enlargement of the affected ventricular chamber.^{1,2} The response to severe volume overload consists of three phases: i) induction of an acute dilatation with a modest increase in sarcomeric length along with rearrangement and slippage of myofibrils.³ The second phase consists of compensatory remodeling characterized by further chamber dilation and eccentric hypertrophy. This hypertrophic response (in which sarcomeric units are added primarily in series), permits ventricular volume to increase so that an increased stroke volume can be accomplished with normal shortening of each sarcomere, and is associated with normalized wall stress.³ In the third phase there is progression of compensatory hypertrophy towards heart failure, however, the ratio of myocyte length to myocyte width increases markedly. This introduces a disproportionate lengthening of the individual contractile cells resulting in a markedly dilated, relatively thin-walled chamber^{1,2}. In addition, the function of these elongated cells has been shown to be markedly depressed relative to myocytes from untreated control animals.⁴ Cardiac fibrillar collagen forms an intricate structural weave that interconnects muscle cells, musclebundles and blood vessels and provides structural integrity of adjoining myocytes.^{5,6} It is believed that collagen strands bear mechanical loading during systole and limit myocyte excursion during diastole. These actions belie the important role of collagen in functional properties of the ventricle.⁶⁻⁸ It has been demonstrated that the

structure and composition of the fibrillar collagen matrix is significantly altered in various models of CHF. Collagen degradation has been proposed to contribute to systolic contractile dysfunction in several cardiovascular abnormalities such as ischemia,^{9,10} stunned myocardium,^{11,12} and dilated cardiomyopathy.¹³ The cellular and molecular mechanisms converting external stimuli into extracellular and subsequently intracellular biologically active compounds are unknown, but abnormalities in myocardial extracellular matrix (ECM) and/or regulation of myocardial matrix components might play a role. The remodeling process involves a cascade of events that are initiated, directed and coordinated by biological signals that possibly involve cell surface receptor proteins, membrane bound enzymes, physical perturbations or cell shape change, altered ion concentrations, feedback pathways and cellularly released agents (chemotactic agents), and other biologically active compounds. These signals indirectly produce the necessary changes required to initiate and successfully complete tissue remodeling and restore short-lived homeostasis. Fibrillar collagens maintain cardiac muscle alignment and impart a tensile strength to the myocardium thereby governing tissue stiffness and ventricular shape and size. An anatomic requisite for ventricular remodeling may be a disruption and degradation of collagen fibers. The conformation of interstitial collagen is such that it is highly resistant to degradation by all proteinases except for specific collagenases. Matrix metalloproteinases (MMPs) constitute an important enzyme system that is known to contribute to the tissue remodeling process by affecting the breakdown of fibrillar collagen matrix network.^{14,15} They are a class of zinc-dependent enzymes that have a high specificity for matrix components.¹⁶ Altered expression and activity of MMPs have been identified in several pathological processes, including with the

development of severe CHF, such as in cardiomyopathy (CMP),^{17,18} myocardial infarction (MI)¹⁹ pressure overload²⁰ and pacing induced heart failure.²¹ However, the role of MMPS in volume overload induced heart failure has not been well defined. In this regard, a major goal of this study was to determine whether heart failure by volume overload is accompanied by any changes in myocardial MMP activity and/or expression. As a small increase or decrease in MMP activity may tip the balance in favor of matrix degradation or synthesis, respectively, this issue is relevant to the progression of heart failure.

The renin-angiotensin system (RAS) plays an important role in the control of cardiovascular homeostasis.²² Several studies have indicated the existence of local RAS in individual organs and tissues, such as the kidney, vascular smooth muscle, brain and heart.²³ Angiotensin II (angiotensin) is a key effector molecule of the RAS. Angiotensin increases protein synthesis in cardiac myocytes, DNA synthesis in cardiac fibroblasts^{24,25} and apoptosis in cultured cardiac myocytes²⁶. Angiotensin exerts its biological effect via binding to at least two different receptor subtypes, angiotensin type 1 and type 2 receptors (AT₁ and AT₂ receptors respectively).^{27,28} Previous studies have suggested the involvement of the cardiac RAS in the development of pressure overload hypertrophy. Upregulation of cardiac components, such as left ventricular angiotensinogen, angiotensin converting enzyme and angiotensin receptor mRNA have been described in pressure overloaded hypertrophy.²⁹⁻³¹ An increase in cardiac renin activity as well as in plasma and LV cardiac angiotensin I and angiotensin II have also been reported in volume overload due to AV shunt.³² Furthermore, atrial hypertrophy in rats with AV shunt is paralleled with significant increase in atrial AT₁ mRNA content.³³ There is

evidence that myocyte angiotensin is released by stretch of the cell and can mediate the effects of stretching on myocyte hypertrophy and gene expression.³⁴ AT₁ receptor regulation may be important in hypertrophy and heart failure since AT₁ transmits hypertrophic stimuli, controls proliferation, differentiation and neurohormonal activity.³⁵ We undertook to investigate whether RAS, via AT₁ receptor activation, is involved in the development of volume overload induced cardiac hypertrophy. To this end, we have used losartan a common non-peptide angiotensin (type I) receptor specific antagonist.

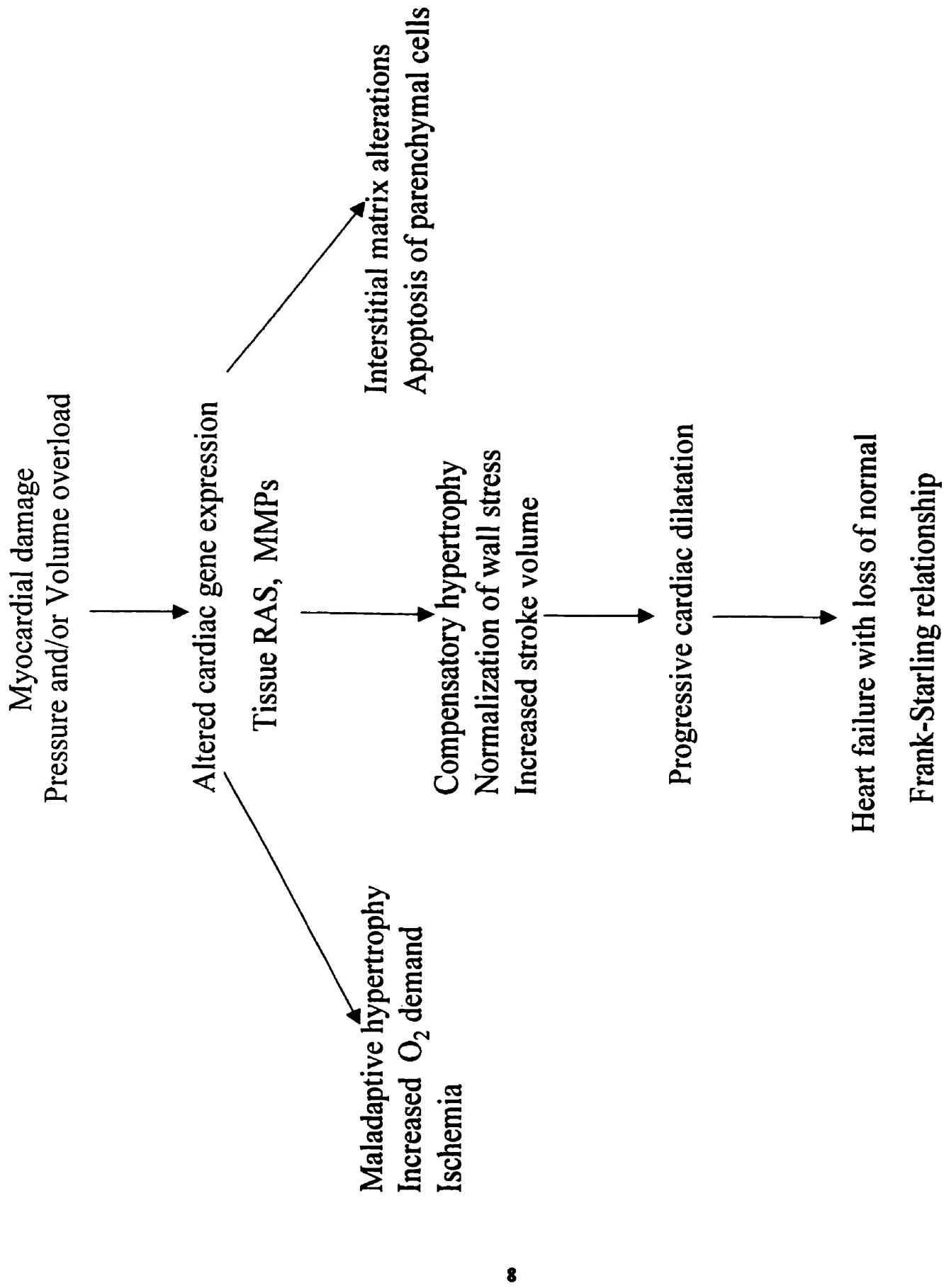


Figure 1. Pathogenetic mechanisms in myocardial remodeling due to pressure and/or volume overload.

III. REVIEW OF THE LITERATURE

1.1 Heart Failure: Incidence and Mortality Rate

Congestive heart failure (CHF) is a major contributor to the morbidity and mortality associated with cardiovascular diseases.³⁶ Currently, about three hundred thousand Canadians are estimated to be suffering from heart failure³⁷ and as the population ages, the incidence of CHF is expected to increase. In the age group 80 and above, the prevalence of CHF increases from 1% to 10% suggesting that heart failure incidence increases with advancing age.³⁶ Heart failure develops in both women and men, with less prevalence in women. In men, the 1- and 5- year survival rates are 55% and 38% respectively, whereas it is 64% and 38% respectively in women.³⁸ Aside from the loss of life and reduced quality of life, heart failure has a substantial economic impact. Approximately 35% of all patients with heart failure are hospitalized once a year and billions of dollars in health care resources are used directly and indirectly on the treatment of patients with heart failure. Thus remedies are needed that can not only attenuate the consequence of heart failure but also restore to normal, the pathological remodeling and attendant functional abnormalities of the affected heart.

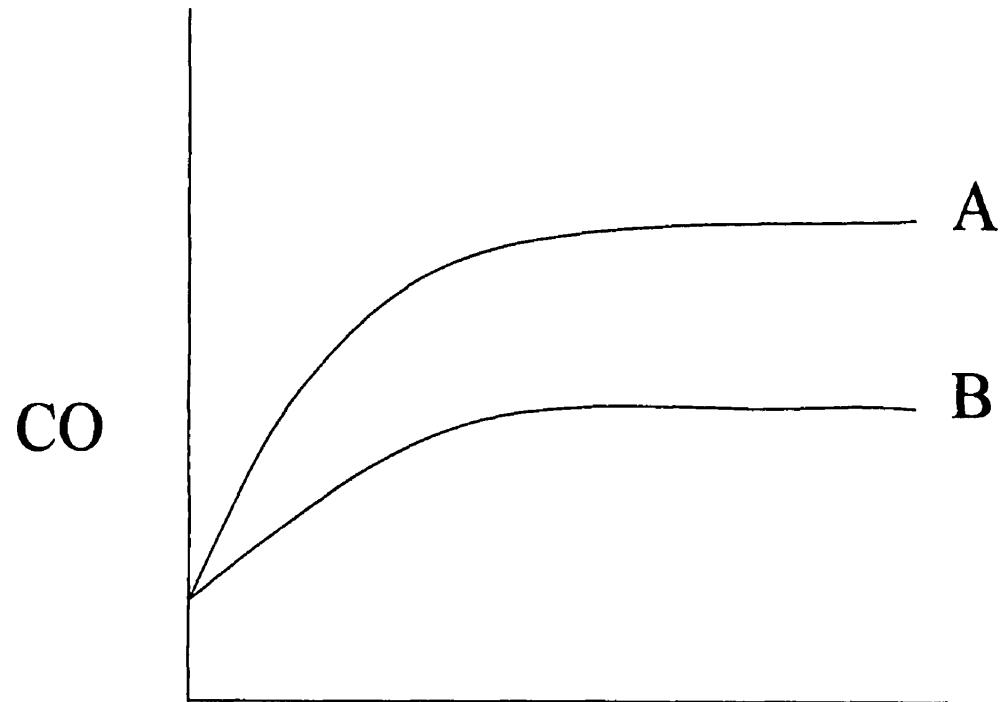
1.2 Heart Failure Defined

Heart failure has been defined as the mechanical failure of the heart to maintain systemic perfusion commensurate with the requirements of the metabolizing tissues³⁹ and is the common end point of many types of cardiovascular diseases. Many factors, alone or in combination, may lead to this pathological condition which include; loss of muscle (myocardial infarctions), decreased myocardial contractility (dilated cardiomyopathy),

pressure overload (aortic stenosis or hypertension), restricted filling (pericardial restriction, endomyocardial restriction, hypertrophy or scarring), and volume overload (anemia, chronic strenuous exercise, systemic arteriovenous fistula, left to right shunt due to ventricular septal defect and hyperthyroidism).

1.2 Pathophysiology of volume overload induced heart failure

Preload is the stretching force that determines end-diastolic muscle length. Ventricular preload is directly related to the ventricular end-diastolic pressure, which in turn influences ventricular end-diastolic volume. Therefore, preload increases as ventricular diastolic filling increases, imparting greater mechanical stretch on the myocardium.^{1,2} When myocardial function is normal and the individual is at rest, the amount of venous blood returning to the ventricles produces a moderate amount of muscle stretch. Normal activity above baseline produces a moderate increase in the volume of blood returning to the heart, further stretching the muscle fibers of each ventricle as the heart is filled during diastole. This degree of increased stretch improves stroke volume and cardiac output by increasing the force of contraction according to *Frank-Starling law* (Fig.1, curve A).⁴⁰ Stretch or changes in the resting sarcomere length effects factors such as calcium influx during excitation, calcium induced release of calcium from the sarcoplasmic reticulum, and sensitivity of the myofilaments to a given amount of calcium. Volume overload of the ventricles occur when abnormally increased diastolic filling is imposed on the ventricular chamber such that increase in filling volume lead to smaller changes in stroke volume and CO compared to normal heart (Fig.1, curve B). Clinically, this may occur in the left ventricle with *mitral valvular insufficiency*



Preload (LVEDV) or stretch

Figure.2 Frank-Starling curve showing the relationship between cardiac strech (preload) and cardiac output.

Curve A represents optimal strech of sacomeres, curve B represents diseased (volume overload) state.

and/or regurgitation and/or *aortic valvular regurgitation*. Problems of the corresponding right ventricular volume overload are *tricuspid valvular regurgitation* and *pulmonary valvular regurgitation*. Congenital and acquired cardiac lesions may also impose a volume overload on the left ventricle. A *ventricular septal defect* presents additional volume to the left atrium and consequently the left ventricle to the right ventricle and eventually back to the left atrium. A *patent ductus arteriosus* permits ejection of blood from aorta into the pulmonary arterial tree and consequently to the left ventricle. An *atrial septal defect* imposes a volume overload on the left atrium into the right atrium and right ventricle. Problems in volume distribution in the heart are of considerable importance due to their role in contribution to cardiac remodeling. Increased diastolic wall stress caused by volume overload results in general fiber elongation and chamber dilation (eccentric hypertrophy).¹ In mild heart failure, cardiac output (CO) is reduced because of limited myocardial contraction which may be due to loss of normal ventricular geometry. Adaptations by the body (Table 1), which results in expansion of blood volume and increased venous tone, lead to an increase in ventricular filling pressure. In the presence of mild ventricular dysfunction, this augmentation of filling pressure may cause a compensatory increase in CO. Nevertheless a plateau is reached rapidly, after which additional preload (filling) produces no improvement in the amount of blood ejected. As heart failure becomes more severe, deterioration in cardiac function may be accompanied by marked increase in filling pressure that are not accompanied by improved CO. In end-stage heart failure, CO may barely reach levels that are adequate for the maintainence of minimal perfusion of the body tissues at rest.

1.3 Experimental Models of Heart Failure

There are a plethora of different experimental models available to the investigator for the assessment of volume overload. These are as follows:

- i) Aortic insufficiency- in the rabbit or rat, this condition is induced by traumatic rupture of the aortic valve leaflets with a catheter. Volume overload of the left ventricle is produced with congestive heart failure in some animals.⁴¹
- ii) Tricuspid insufficiency- this method produces volume overload of the right ventricle. In addition to cutting cordae tendinae, pulmonary aortic banding increases the regurgitation and the degree of congestive failure.⁴²
- iii) Mitral insufficiency- This model involves cutting of the mitral valve chordae. Animals usually expire with acute cardiac failure.⁴³
- iv) Atrial septal defect- surgical production of a defect in the interstitial septum results in volume overload of the right ventricle,⁴⁴ but has not been a successful model for the production of congestive heart failure.
- V) Anemia- this model may be produced by withholding access to supplemental dietary iron from young animals, and results in marked cardiac hypertrophy leading to cardiac decompensation if combined with additional bleeding to reduce blood hematocrit to less than 3-5%. The rat and pig are most commonly used mammals for this particular purpose.⁴⁴
- vi) Creation of an arterio-venous Shunt

An arterio-venous shunt can be created in many different ways in experimental animals. These includes:

- a) Aorto-vena caval: this method is most commonly used in dog and

rat⁴⁵ and produces increased left ventricular end diastolic pressure

b) Aorto-Pulmonary: this method produces volume overload of the left ventricle with increased left ventricular end-diastolic pressure.⁴⁶

c) Aortic-left atrium: this method produces volume overload of the left ventricle with increased left ventricular end diastolic pressure.⁴⁶

Volume overload by AV shunt has the advantages of high reproducibility, low mortality, the rapidity with which it can be performed, and the option it provides the surgeon with to control the degree of volume overload by selecting the size of the shunt. Four weeks after the creation of a shunt, hemodynamic and neurohormonal changes are in progress and these are similar in many aspects to human heart failure depending on the shunt size.⁴⁷

Volume overload (as well as pressure overload) increases wall stress and activates the initiation of compensatory myocardial hypertrophy, which is initially an adaptive process, but which ultimately results in further deterioration of myocardial performance.

1.4 Compensatory adaptations in heart failure

Many compensatory mechanisms, which initially tend to restore normal cardiac function are activated in heart failure (Table 1). Activation of these compensatory mechanisms augment cardiac performance by altering four determinants of myocardial pump function: preload, afterload, contractility and heart rate.⁴⁸ However, as the factors responsible for heart failure persists, these compensatory mechanisms become abnormally augmented and maladaptive changes occur which lead to cardiac dysfunction; further changes in the peripheral circulation aggravate the condition.

Cardiac and Extracardiac Alterations leading to Heart Failure

Type of adaptation	Effect
Cardiac	Increased heart rate
	Maximized Frank-Starling mechanism
	Dilatation
	Hypertrophy (eccentric and concentric)
	Widening of AV oxygen difference
Extracardiac (systemic)	Increased sympathetic (SNS) stimulation
	Activation of RAS
	Release of atrial natriuretic peptide

Table 1. Compensatory mechanisms undergoing in heart subject to hemodynamic loads

Neurohormones, especially derived from the sympathetic nervous system (SNS) and those of the RAS, are important mediators of both cardiac and extra-cardiac adaptations.⁴⁹

The predominant cardiac compensatory mechanism in heart failure is myocardial remodeling,⁵⁰ a complex of anatomic, physiologic, histologic and molecular changes that the myocardium undergoes in response to trophic stimuli. In normal hearts, the majority of myocardial cells remain in the G phase of mitosis and do not routinely re-enter the cell cycle to increase their number, but they do commonly undergo remodeling by which they can increase in length (dilation) or volume (hypertrophy). Chronic abnormal myocyte hypertrophy is associated with intrinsic loss of contractility and furthermore dilation leading to impaired cardiac function (Frank-Starling mechanism). In addition, ventricular dilation increases wall stress, one of the determinants of myocardial oxygen demand. Hypertrophy, if severe and long standing, is usually associated with the loss of contractile force. Studies have shown that hypertrophied heart tends to outgrow its blood supply, which is manifest as decreased capillary density. This tends to limit coronary flow and coronary vascular reserve and thus may further worsen oxygen supply and the availability of high energy phosphates in hypertrophied muscle.⁵¹ Activation of SNS may reduce blood flow to the skin, viscera, and kidneys and raises peripheral vascular resistance by constricting blood vessels, thereby increasing afterload.^{52,53} These actions shunt blood to vital organs, increase preload, and stimulates retention of sodium and water by kidneys to compensate for perceived lack of blood volume; however, this compensatory action proves to be maladaptive.⁵³ Fluid retention increases venous return and RV and LV

filling pressures (volume overload), which contributes to congestive symptoms of heart failure. Similarly, increased afterload adds to the strain on heart. Falling CO stimulates other body systems to try to return blood flow to normal levels. The major neurohormonal systems that regulate these compensatory mechanisms are the sympathetic, RAS, and atrial natriuretic systems. Since the early 1980's, the RAS has been implicated in the pathophysiology of heart failure. Angiotensin II has many properties that are deleterious to the patient with heart failure.⁴⁹ It is a potent vasoconstrictor of the peripheral arterioles⁵⁴ and plays a major role in increasing afterload in these patients. The peptide also has direct and indirect effect on renal excretion of sodium and water.⁴² Both effects increase circulatory blood volume, further increasing preload.

The myocardium is composed of myocytes and nonmyocyte cells.³⁹ Therefore heterogeneity in myocardial structure is mediated by the involvement of different myocardial compartments. The hypertrophic growth of cardiac myocytes (parenchymal cells) leads to an increment in myocardial mass while nonmyocyte growth (stromal cells) is reflected in a structural remodeling of the cardiac interstitium. Disproportionate growth between myocyte and nonmyocyte cells set the stage for abnormal myocardial function. It is now recognized that the extracellular matrix in general, and fibrillar collagens in particular, are essential element of myocardial structure and function and dynamic participant in its remodeling. Some evidence exists to support the view that matrix components per se may be involved directly in modulating the growth of cardiac myocytes.⁸

1.5 CARDIAC EXTRACELLULAR MATRIX

The myocardium comprises of cardiac myocytes surrounded by cardiac interstitium that contains a complex fibrillar collagen and elastin network, and non myocyte cells including fibroblasts, macrophages, mast cells, vascular smooth muscle cells and endothelial cells. Two-thirds of all myocardial cells are nonmyocytes. It has been suggested that structural abnormalities within the cardiac interstitium serve as the primary mechanism for myocardial failure.⁸

The collagen matrix of the myocardium is a structural continuum which includes the valve leaflets, cordae tendinae, as well as the interstitial and perivascular fibrillar weave that serves to maintain the architecture of the myocardium and to transmit developed force in heart cycle.⁵⁵ The matrix is segregated into various components; the epimysium, the perimysium and endomysium. The endomysial struts maintain myocyte alignment and prevents their slippage, the epimysium fibres protects the muscle fibres from being overstretched while perimysial weaves facilitate force transmission through the myocardium.. The matrix is an important determinant of the physical characteristics of the myocardium including modulation of anisotropic passive stretching of these tissues. Components of the matrix include collagens, proteoglycans, glycoproteins (e.g. fibronectin), peptide growth factors and proteases.⁵⁶ The major structural proteins of the myocardial collagen matrix are fibrillar type I and III collagens.^{57,58} Type I collagen is usually present in the form of thick fibers, and possesses the greatest tensile strength of all myocardial matrix components,⁵⁹ while type III collagen forms fine reticular networks. The ratio of type I to type III collagens found in non-human primate myocardium is 7.4:1 and in normal rat hearts is 3.5:1.^{57,58} These geometrical features and

arrangements of collagen fibers, together with collagen phenotype, cross-linking and collagen protein concentration, determines the stiffness of intact myocardium. Other types of collagen proteins including type IV and type V collagens are represented in small amounts within the myocardium and are mainly associated with cell membranes. The myocardium also contains a small number of elastin fibres. The matrix and cytoskeletal components have been suggested to combine together to create a system that is mechanically self-stabilizing.⁶⁰ In addition to being a scaffolding that supports the cellular elements of various tissues, the matrix and its components serve as modulator of cell growth and tissue differentiation.⁶¹ Tissue differentiation results in changing dynamics between parenchymal and stromal matrix elements. There is an increasing appreciation that through changes in the nature (weave) and quantity of the matrix, nonmyocytes in the heart could play a primary role in determining the response of the myocardium to pathologic stimuli.⁶² Pathologic myocardial hypertrophy often involves interstitial fibrosis, which may both alter the mechanical properties of the myocardium and restrict the delivery of the nutrients to myocytes. Conversely, a reduction or alteration in the quantity of certain structural proteins could compromise the integrity of the extracellular skeleton and lead to chamber dilation. Changes in matrix composition requires the removal of previous extracellular components. This is accomplished by the action of selective proteases which selectively degrade matrix components and may alter cell-matrix attachments.^{14,16} Normally, both matrix removal and synthesis occurs simultaneously in an orderly and progressive fashion. The end-stage of differentiation is a steady-state balance between new matrix formation and matrix removal. Again, this occurs in a manner such that a preservation of matrix function, with no gross change in

matrix composition, is maintained. In the pathological state however, normal homeostasis between the parenchymal elements and the surrounding stroma is disrupted. This may cause alterations in the distribution, composition and function of the matrix that are evident in many pathological conditions including congestive heart failure. Matrix changes that occur in heart failure includes increase in fibronectin, laminin and vimentin proteins, as well as altered deposition of collagen fibers I, III, IV and VI.⁶³⁻⁶⁵ Significant increase in collagen tissue concentration has also been found in the ventricular free walls of rat, 13 weeks after myocardial infarction.⁶⁶ In an animal model of tachycardia induced heart failure, Zellner et al.⁶⁷ found significant reductions in myocyte attachment to the basement membrane proteins laminin, fibronectin and collagen IV. Marked alterations in content and isoform distribution of collagen in cardiac interstitium have also been reported in human CHF.⁶⁸ Collectively, these changes may result in the loss of force transmission through the ventricular free wall. Secondly, changes in myocyte scaffolding and alignment could result in fiber slippage, fiber realignment and ventricular wall thinning.⁷ Thus, alterations in the ECM may have a potentially large impact on ventricular function and this nonmyocyte remodeling process may, therefore play a significant role in the decline of cardiac pump function in CHF.

1.5.1 Collagen

Collagen, which is the most abundant structural protein of the body, is present in the myocardium in relatively small amounts (2-4% of total myocardial volume). Cardiac collagen is highly insoluble and difficult to extract. Collagen type I and III represents more than 90% of the total collagen in the myocardium.^{7,69} In the heart, fibrillar collagens

are produced exclusively by fibroblasts and myofibroblasts.^{70,71}

1.5.2 Collagen Function

Fibrillar collagens have multiple functions and are essential components of the myocardium. They impart a tensile strength to the myocardium that governs tissue stiffness and maintains myocardial ultrastructure as well as beat-to-beat cardiac function. Collagen fibers interconnect myocytes to one another and facilitate the transmission of myocyte generated force to the ventricular chambers.⁷² The energy stored in the coiled collagen fibers during myocyte contraction contributes to myocardial relaxation and re-lengthening.⁷³ Collagen weaves and struts support intramural coronary vessels within the cardiac interstitium and are known to contribute to the transduction of force generated by myocytes, prevention of muscle fiber slippage, and protection of myocytes from overstretching. Collagen type IV is localized in the basement membrane of cardiac myocytes and fibroblasts where it forms a crucial structural component and plays a regulatory role in molecular transport and cell adhesion. Collagen type V co-exists with type IV in basement membrane and is interspersed with types I and III collagens in the cardiac interstitium. Type VI collagen is found in the cardiac interstitium where it is associated with other fibrillar collagen types and coat the surface of collagen fibers. Collagen type IV, V, and VI constitute approximately 10 to 12% of total cardiac collagen.

1.5.3 Collagen Biosynthesis

In the ECM, collagen is continuously being turned over and being replaced by newly

secreted collagen. The amount of collagen secreted is the net result of intracellular collagen synthesis and intracellular collagen degradation. It has been reported that 10 to 90% of newly synthesized procollagen undergo degradation before or soon after their secretion.^{74,75} Although this apparent inefficiency is intriguing, it may be a pathway to allow for rapid adjustments in collagen production. In such a system, the activation or inhibition of existing metabolic pathways modulates the amount and type of collagen production without requiring a significant lag phase during which these pathways are upregulated de novo. Additionally, degradation may be a method of quality control, since it has been shown to recognize and selectively destroy structurally aberrant procollagen molecules.⁷⁶ The biosynthesis of a collagen molecule involves at least 8 distinct enzymes (5 intracellularly and 3 extracellularly). In general, collagen synthesis involves a process similar to that for other secretory proteins. However, it is unique in its post-translational modifications of the primary molecule.⁷⁷ Procollagen is synthesized by membrane bound ribosomes and the resultant polypeptide has a signal sequence at its amino terminus. In the cytoplasm, the collagen mRNA is translated by polysomes located on the external surface of the endoplasmic reticulum.⁷⁸ The result is a transcribed polypeptide chain which is translated within the lumen of the rough endoplasmic reticulum (RER) to the Golgi body where it is packaged for secretion into condensing granules and secretory vesicles.⁷⁹ During this phase, the polypeptide undergoes post-translational modification^{77,79} involving the removal of the signal polypeptide, hydroxylation of specific proline and lysine residues, glycosylation of specific hydroxylysine residues, pro-alpha chain association, intra- and inter-chain disulfide bond formation as well as the formation of an overall triple helical structure. The procollagen

molecule is then secreted into the extracellular space. The hydroxylation of proline and lysine is dependent on the presence of ascorbic acid.⁸⁰ In vitro, even in the absence of ascorbic acid, collagen is still produced and secreted by fibroblasts, however there is a significant decrease in hydroxylation.⁸¹ Procollagen precursors, of collagen phenotypes I, II and III, are converted to insoluble tropocollagen by procollagen peptidase mediated cleavage of the non-helical propeptides at the amino termini.⁸² This enzymatic cleavage of the procollagen peptide is thought to occur in a series of stages and that partial retention of the procollagen sequence temporarily distinguishes the immature collagen by preventing it from fully aligning in a manner typical of mature collagen. The conversion to mature collagen is time-dependent. As the tropocollagen molecule matures, covalent linkages form and serve to augment the early non-covalent and hydrogen bond cross-linkages. The covalent cross-links become more stable as collagen matures.⁸³ The cross-linking is important for optimal functioning and may also serve to regulate the rate of catabolism in vivo.⁸⁴ Selective gene expression, control over the process of polypeptide cleavage, varying the amount and nature of the non collagen matrix molecules secreted, and alterations in the amount of collagen cross-linking have all been proposed as mechanisms of cellular control over collagen fibril size and structure.⁸⁵

1.5.4 Collagen Phenotype as a Biological Signal

Collagen has been shown histologically and biochemically to be metabolically active. Collagen synthesis responds to physiologic and functional demands of the tissue and this response often produces changes in the proportion of collagen phenotypes, which is most prevalent for the types I and III interstitial collagens. Studies have shown that in several

pathological conditions, the ratio and cross-linking between type I and type III collagens is severely altered^{86,87} and this may have a bearing on the altered functioning in hypertrophied hearts.

There are three possible routes by which a tissue may alter the amount and proportion of its collagen phenotypes. Clonal selection of cells producing a particular collagen phenotype would produce a long-term shift in collagen phenotype production by changing the ratio of fibroblast subpopulations and would involve stimulation of progenitor cells to proliferate and eventually fully differentiate.⁸⁸ A change in the present cell population's synthetic pattern would produce a more rapid response and a more short term resolution.⁸⁹ The third method would be to alter the susceptibility of various collagen phenotypes to degradation. For example, in certain situations extracellular collagenase has been observed to be relatively selective in its digestion of collagen phenotypes.⁹⁰ The ability to modulate the overall synthetic pattern of various collagen phenotypes may be essential for normal tissue development and repair. Therefore changes in the proportion of collagen phenotypes (especially that of type I and III) may be a biologic indicator of the metabolic state of a connective tissue, the dynamic functional demands being experienced and the changing cellular activity.

1.6 Collagen Degradation

Although the accumulation of total collagen in various tissues is stable throughout life,⁹¹ collagen can be rapidly turned over in various organs depending on the tissue requirements. Collagen is degraded by several different mechanisms. There are two major modes of intra-cellular degradation; basal and enhanced.^{92,93} Basal degradation is

believed to occur between the ER and the Golgi apparatus, whereas enhanced degradation involves the lysosomal system.⁹⁴ The degradation of de novo synthesized collagen is presumed to be a mechanism by which the cell destroys abnormal molecules,^{76,95} selects for collagen types by controlling the ratio of interstitial collagen chains available⁹⁶ and rapidly modulates the amount of collagen being produced.

Once extracellular collagen is cross-linked, it is largely resistant to degradation by proteolytic enzymes. The peptide bonds of the collagen are protected from enzymatic attack, and subsequent denaturation, by the cross-linkages of the fibrils and triple helical arrangements of the collagen molecule. The second major method of collagen degradation is via phagocytosis.⁹⁶ It has been observed that fibroblasts can phagocytose enzymatically cleaved collagen fiber segments and subsequently degrade collagen within phagolysosomes.⁹⁷ Beertsen⁹⁸ believes that the collagen phagocytosis is related to the rate of collagen turnover. During hypofunction, there is a net loss of extracellular collagen fibrils due to two-fold increase in its cellular ingestion. It appears that an external stimulus may stimulate fibroblasts to become phagocytic.⁹⁹ Cytokines are known to influence the phagocytic process. IL-1 has been shown to inhibit phagocytosis while TGF-beta was shown to enhance this process.¹⁰⁰ Finally, collagenase, a metalloproteinase, is responsible for extracellular collagen homeostatic and remodelling turnover.^{101,102}

1.6.1 Matrix metalloproteinases (MMPs)

MMPs are a family of 15 now known zinc-dependent, neutral endopeptidases that can act together or in concert with other biological factors to degrade various components

of the extracellular connective tissue matrix including collagens, gelatins, fibronectin, laminin and proteoglycans. This family of enzymes attacks collagen at specific sites on the molecule thus splitting it into segments which denature⁹⁹ and the denatured chains are digested by proteolytic enzymes.¹⁰⁴ The myocardial collagenase system was first identified by Montfort and Perez-Tamayo in 1975.¹⁰⁴ The cardiac MMP is localized in endomyocardium, subendocardial space and in interstitial space. MMPs are produced by fibroblasts, polymorphonuclear leucocytes and macrophages¹⁰⁵ and are classified into three major groups: interstitial collagenase, gelatinase (type IV collagenases) and stromelysins (Table IV). Interstitial (type 1) collagen (MMP-1) is the only enzyme capable of cleaving the alpha chains of fibrillar collagens producing 3/4 and 1/4 size fragments.¹⁰² At physiologic temperatures, the cleaved fragments spontaneously denature into gelatin derivatives. The gelatinases (MMP-2 and MMP-9) degrade gelatin derivatives and the collagen present on the basement membrane, type IV collagen. The third class of MMPs, the stromelysins have relatively broad substrate specificity.

Collagenases

There are three distinct collagenases: MMP-1,¹⁰⁷ MMP-8¹⁰⁸ and MMP-13.¹⁰⁹ These enzymes have the unique ability of cleaving the alpha chains of fibrillar collagens at specific sites. Once the initial cleavage of collagens occurs, the two collagen fragments are no longer stable and the normal triple helical structure is lost leaving the remnant polypeptide chains open to further degradation by other proteinases. Substrates degraded by interstitial collagenase (MMP-1) are collagen types I-III, VII, X¹¹⁰ and gelatins and other proteins such as serum amyloid A.¹¹¹ MMP-8 degrades collagen types I-III. The

three collagenases mentioned above vary in their specificity for the fibrillar collagens. For example, the substrate with highest affinity for MMP-1 is type III collagen, MMP-8 has the greatest activity against type I collagen and MMP-13 is most efficient at degrading type II collagen. MMP-1 is synthesized by stimulated fibroblasts and macrophages,¹¹² MMP-8 is synthesized predominantly by monocyte cells and is stored after secretion in the specific granules that are released upon stimulation of the cell. MMP-13 is synthesized mainly by monocytes.

Gelatinases

These enzymes (MMP-2, MMP-9) have substrate specificity for denatured collagens, type IV (basement membrane) collagens, vitronectin, aggrecan, galectin, and elastin.^{113,114} Expression of MMP-2 is the most widely spread of all MMPs. This enzyme is produced by mesenchymal cells and is often secreted as a complex with TIMP-2.¹¹¹ MMP-9 is expressed in transformed and tumor derived cells, neutrophils, monocytes and alveolar macrophages and is often secreted from macrophages as a complex with TIMP-1.¹¹⁶

Stromelysins

Stromelysins have broad substrate specificity. MMP-3 can degrade aggrecan, link protein, fibronectin, vitronectin, laminin, gelatins, type IV collagen, collagen crosslinks, and procollagens and they also participate in the activation of other MMPs.¹¹⁷ MMP-3 is not normally widely expressed but can be readily induced by growth factors, cytokines (such as IL-1), and tumor promoters in cultured mesenchymal cells such as chondrocytes and connective tissue fibroblasts.¹¹⁸ MMP-7 has no C-terminus domain and is produced

by a limited number of human cell types, including glandular epithelium, mononuclear phagocytes, and renal mesangial cells.¹¹⁹ The substrates it degrades are similar in extent to stromelysin-1 with the addition of elastin but it cannot cleave type X collagen.¹²⁰ It can also activate procollagenases and can cleave α_1 -antitrypsin and entactin.^{121,122} It is secreted by human endomyrium during the reproductive cycle¹²³ and can efficiently inactivate Serpins, the serine proteinase inhibitors.^{121,122}

Membrane Type MMPs (MT-MMPs)

MT-MMPs are a group of enzymes that are present on the cell surface in an active form.¹²⁴ There are at least 4 members of this family, MT1-MMP (MMP-14) to MT4-MMP (MMP-17). All MT-MMPs have a similar domain structure to other MMPs but with three insertions. The first insertion is a furin recognition site located between the propeptide and the N-terminus and it is believed that the MT-MMP arrive at the cell surface in an active form. The second insertion is found in the N-terminus and it is thought to be involved in substrate specificity. The third insertion is a much larger (73 amino acid) and is located after the haemopexin domain in the C-terminus. This insertion contain the hydrophobic transmembrane domain. MMP-14 (MT1-MMP) is found on the surface of the cells.¹²⁵ It is found expressed in human colon, breast, head and neck carcinomas¹²⁶ and its principal substrate appears to be proGelatinase A (inactive MMP-2) for which it acts as a specific activator.¹²⁷

Table 2. The Matrix metalloproteinase family.

Enzyme	MMP No.	Matrix substrates or functions
Collagenases		
Interstitial collagenase	MMP- 1	Collagens I, II, III, VII, and X, gelatins, entactin, aggrecan, link protein
Neutrophil collagenase	MMP-8	Collagen I, II, and III, aggrecan, link protein
Collagenase 3	MMP-13	Collagens I, II, III
Collagenase 4 (xenopus)	MMP-18	Collagen I
Gelatinases		
Gelatinase A	MMP-2	Gelatins, collagens I, IV, V, VII, X, and XI, fibronectin, laminin, aggrecan, elastin, vitronectin
Gelatinase B	MMP-9	Gelatins, collagens IV, V, XIV, aggrecan, elastin, entactin, vitronectin
Stromelysins		
Stromelysin 1	MMP- 3	Aggrecan, gelatins, fibronectin, laminin, collagens III, IV, IX, and X, vitronectin
Stromelysin 2	MMP-10	Aggrecan, fibronectin, collagen IV
Membrane-type MMPs		
MT1-MMP	MMP- 14	Collagens I, II, III, fibronectin, laminin-1, vitronectin, dermatan sulfate proteoglycan, activates proMMP-2 and proMMP-13
MT2-MMP	MMP-15	Not known
MT-3MMP	MMP-16	Activates proMMP-2
MT-4-MMP	MMP- 17	Not known
Others		
Matrilysin	MMP- 7	Aggrecan, fibronectin, laminin, gelatins, collagen IV, elastin, entactin, small tenascin-C, vitronectin
Stromelysin 3	MMP-11	Weak activity on fibronectin, laminin, collagen IV, aggrecan, gelatins
Metalloelastase (Unnamed)	MMP- 12	Elastin
	MMP- 19	Not known

Metalloelastase

(MMP-12) is a 21 kDa MMP found at the surface of human skin fibroblasts along with MMP-2, degrades elastin and is secreted by macrophages.¹²⁸ The substrates for elastases are elastin and plasminogen.

A high percentage of sequence identity exists among the members of matrixin family. For example, human MMP-1 has 60% identity to human MMP-8 and human MMP-3 has 56% identity with human MMP-9. The MMPs share the following characteristics¹⁶: they contain common sequences of amino acids constituting distinct domains; they are secreted as inactive proenzymes; activation can be achieved proteolytically or by treatment with mercurial compounds; activation is accompanied by a fall in molecular weight; they contain zinc at the active centre; the active enzymes are inhibited by tissue inhibitors of metalloproteinases (TIMPs); the active enzymes operate best at neutral pH; and the active enzymes all cleave components of matrix. All three classes of MMPs are secreted by endothelial cells as inactive pro-enzymes, which are devoid of collagenolytic activity. Extracellular activation is required in order to produce functional enzyme. MMPs have a basic 5-domain structure, with the region of the enzyme typically referred to as a signal peptide, a propeptide, a catalytic site , a hinge region and a pixin-like domain.¹⁰⁶ Activation of the pro-enzyme can be achieved through a variety of agents and enzymes. The conformational changes necessary for activation can be brought about by chemical agents such as detergents (SDS), oxidants and other chaotropic agents. Proteases that are capable of activating these enzymes include bacterial collagenases and host-derived proteases such as the serine protease plasmin, stromelysin and the polymorphonuclear (PMN) derived cathepsins. There is

evidence that some of the MMPs are capable of autoactivation as well.

Interstitial or extracellular degradation of collagen in the normal heart is relatively slow. The synthesis rate of collagen for right and left ventricles in dog is 0.56% of total ventricular collagen per day.¹²⁹ This rate of turnover is in agreement with the small amount of active myocardial collagenase (1-2% of total myocardial collagenase) estimated by Tyagi et al.¹³⁰ (in vitro activation with trypsin and plasmin increased the amount of active enzyme to between 80 and 90% of total collagenase).¹³¹ As an unabated secretion or activation of proteinases is capable of destroying an organism's supportive framework, the collagenase system is normally highly regulated.

1.6.2 Regulation of MMP in the heart

The MMPs are regulated at three different levels, including transcriptional regulation, secretion in latent form, and the inhibition of the active enzymes.

Transcriptional regulation

Normal expression of MMPs is characterized by tightly controlled regulation to maintain normal tissue function. A number of cytokines and growth factors have been shown to either stimulate or inhibit the synthesis of MMPs. The pro-inflammatory cytokines interleukin1 (IL-1) and tumor necrosis factor (TNF) stimulate numerous cell types to secrete both collagenase and stromelysin.^{132,133} platelet derived growth factor (PDGF), epidermal growth factor (EGF), basic fibroblast growth factor (bFGF) and tumor necrosis factor-alpha (TNF- α) are all known to stimulate the synthesis and secretion of collagenase and stromelysin.¹³⁴⁻¹³⁶ (TGF- β_1) decreases steady state collagenase 3 (MMP-13) mRNA in osteoblast cultures and decreases levels of

immunoreactive procollagenase.¹³⁷ IL-4,¹³⁸ IFN γ (interferon),¹³⁹ and insulin like growth factors (IGF),¹⁴⁰ can down-regulate MMP biosynthesis while heparin may activate MMPs.¹⁴¹ Within one tissue, cytokines and growth factors may have different effects on individual cell types present. IL-1 induces the expression of collagenase and stromelysin in human fibroblasts but not in keratinocytes,¹¹⁰ and skin and synovial fibroblasts respond differentially to cytokine stimulation.¹⁴² In addition, synergistic effects occur with combinations of cytokines and growth factors.^{142,143} These differences make it difficult to predict the outcome of blocking the action of any individual cytokine in an attempt to modify MMP activation. In many instances where connective tissue is broken down, cooperation between cells occur.¹⁴⁴ Some studies have shown that direct contact between T lymphocytes and monocytes is a major pathway to up-regulate MMP biosynthesis and that factors produced by T-cells can also increase collagenase synthesis by dermal fibroblasts.¹⁴⁵

Secretion of latent matrix metalloproteinases

One characteristic of MMPs is that they are all secreted as inactive Zymogens (with the exception of the MT-MMPs), and they require activation in order to have any effect on the ECM. Plentiful collagenase is found in normal heart but a majority of it is in zymogen form. Tyagi et al.¹²⁹ estimated the activated collagenase component in rat myocardium to be between 1-2% of total myocardial collagenase. The propeptides that are lost during activation contain a common conserved sequence PRCGVPD. The cysteine residue (C) is known to be bound to the catalytic zinc molecule and disruption of this interaction initiates activation.^{146,147} All MMPs can be activated in vitro with organo-

mercurial compounds (eg. 4-aminophenylmercuric acetate), but no agent responsible for the physiological activation of all MMPs has been clearly defined. Complete activation *in vivo* is achieved by removal of the propeptide proteolytically either by other proteinases or auto-catalytically.¹⁴⁸ This allows the enzyme to bind to its substrate and the zinc at the active site can hydrolyze susceptible bonds in the protein substrates. Plasmin (derived from plasminogen by the action of plasminogen activators) has been strongly implicated in collagenase and stromelysin-1 activation in cell model systems.¹⁴⁹ It is not, however an efficient activator of gelatinase A. Mast cell proteinases can also activate proforms of the metalloproteinases.¹⁵⁰ In vitro, stromelysin-1 is an efficient activator of collagenase and gelatinase B (MMP-9).¹⁵¹ Numerous studies indicate that members of the MMP family have the ability to activate one another. This is well illustrated by MT-MMP, localized to the cell membrane of tumor cells and is known to activate MMP-2,¹²⁰ which is produced primarily by stromal fibroblasts. Some MMPs (MMP-11, MT-MMPs) have a 10 amino acid insert between the propeptide and the N-terminal domain. This sequence (GLSARNRQKR) in MMP-11 is recognized by a Golgi associated serine proteinase, furin. Furin activates these MMPs and thus the MMPs are secreted from the cell in an active form.¹¹⁸ Recently matrilysin have been found to activate both progelatinase A and B in the presence of their bound inhibitors TIMP-1 and TIMP-2 respectively.¹⁵² The activation of the metalloproteinases *in vivo* is likely to be a critical step in terms of their biological behavior, because it is this activation that will tip the balance in favor of matrix degradation.

Inhibition of active enzymes

The third level of MMP regulation occurs through the inhibition of their actions.

Inhibitors of MMPs include the broad spectrum α_2 -microglobulin and the specific tissue inhibitors of matrix metalloproteinases (TIMPs).

1.6.3 Tissue Inhibitors of MMPs (TIMPs)

TIMPs are a family of proteins that can form complexes with matrix MMPs and irreversibly inactivate them. TIMPs are proteins of about 200 amino-acid residues, 12 of which are cysteines involved in disulfide bonds. Four members of this family of proteins have been identified and includes TIMP-1, TIMP-2, TIMP-3 and TIMP-4.¹⁵³⁻¹⁵⁶ Of these TIMP-1, TIMP-2 and TIMP-4 are known to be expressed in the heart and are secreted as soluble proteins. TIMP-3 is present as insoluble protein in the matrix.¹⁵⁷ All active MMPs are inhibited by TIMPs^{168,159} and all connective tissues contain TIMPs.¹⁵⁶ The main form, TIMP-1 is a 30-kDa glycoprotein and is synthesized and secreted by most connective tissue cells as well as by macrophages. It also resides in platelet α -granules and can be identified in most body fluids. The amino acid sequence of human TIMP-1 is largely conserved across species. TIMP-1 has relatively high affinity for the active forms of collagenases, stromelysins and gelatinases and forms irreversible, non-covalent complexes with them. The inhibitory activity resides in the first three loops of a structure which is characterized by the presence of six disulfide bridges.¹⁶¹ The protein is remarkably stable to both temperature and low pH. TIMP-1 is very similar to the second member of the family, TIMP-2, a 21kDa unglycosylated protein secreted by human melanoma cells and which has inhibitory properties similar to TIMP-1. Human TIMP-2 is about 40% homologous with human TIMP-1 and the six disulphide bridges in TIMP-1 are conserved in TIMP-2. Both proteins bind tightly to the active form of the MMPs in a

1:1 ratio. TIMP-2 may bind to the proform of gelatinase A but not the progelatinase B which can bind to TIMP-1. The third member of the family, TIMP-3 is markedly different from TIMP-1 and TIMP-2, but it retains the six disulfide bridges. TIMP-2 has been suggested to be involved in growth promoting activities. Recently, it has been reported that TIMP-2 overexpression caused reduction in cell proliferation while TIMP-3 overexpression induced DNA synthesis and promoted smooth muscle cell death by apoptosis.¹⁶² TIMP-4 has been found to be highly expressed in heart indicating a role in cardiac collagen remodeling.¹⁶³

TIMP plays an important role in controlling tissue breakdown by blocking the action of the activated MMPs and also by preventing the activation of proenzymes.¹⁶⁴ Both C- and N-terminal domains of MMPs are involved in binding to TIMPs. TGF- β and retinoic acid are able to up-regulate the production of TIMP-1.¹⁶⁵ IL-6¹⁶⁶ and other members of this family have been shown to up-regulate TIMP-1 from fibroblasts of some connective tissues. Angiotensin II was also found to up-regulate TIMP-1 gene expression in rat heart endothelial cells.¹⁶⁷

TIMPS are present wherever MMPs are found as they are co-expressed. Thus, when MMPs are over-expressed, TIMPs are also over-expressed¹⁶⁸ and some studies have showed up-regulation of MMPs to be associated with a similar up-regulation of TIMPs.^{151,162} However, some studies have shown that MMPs and TIMPs can be regulated independently and sometimes reciprocally.¹⁷⁰⁻¹⁷³

Thus in some diseased states, certain factors may exist that can cause over expression of MMP or TIMP, thereby favoring matrix degradation or matrix production respectively.

Extracellular Matrix Degradation

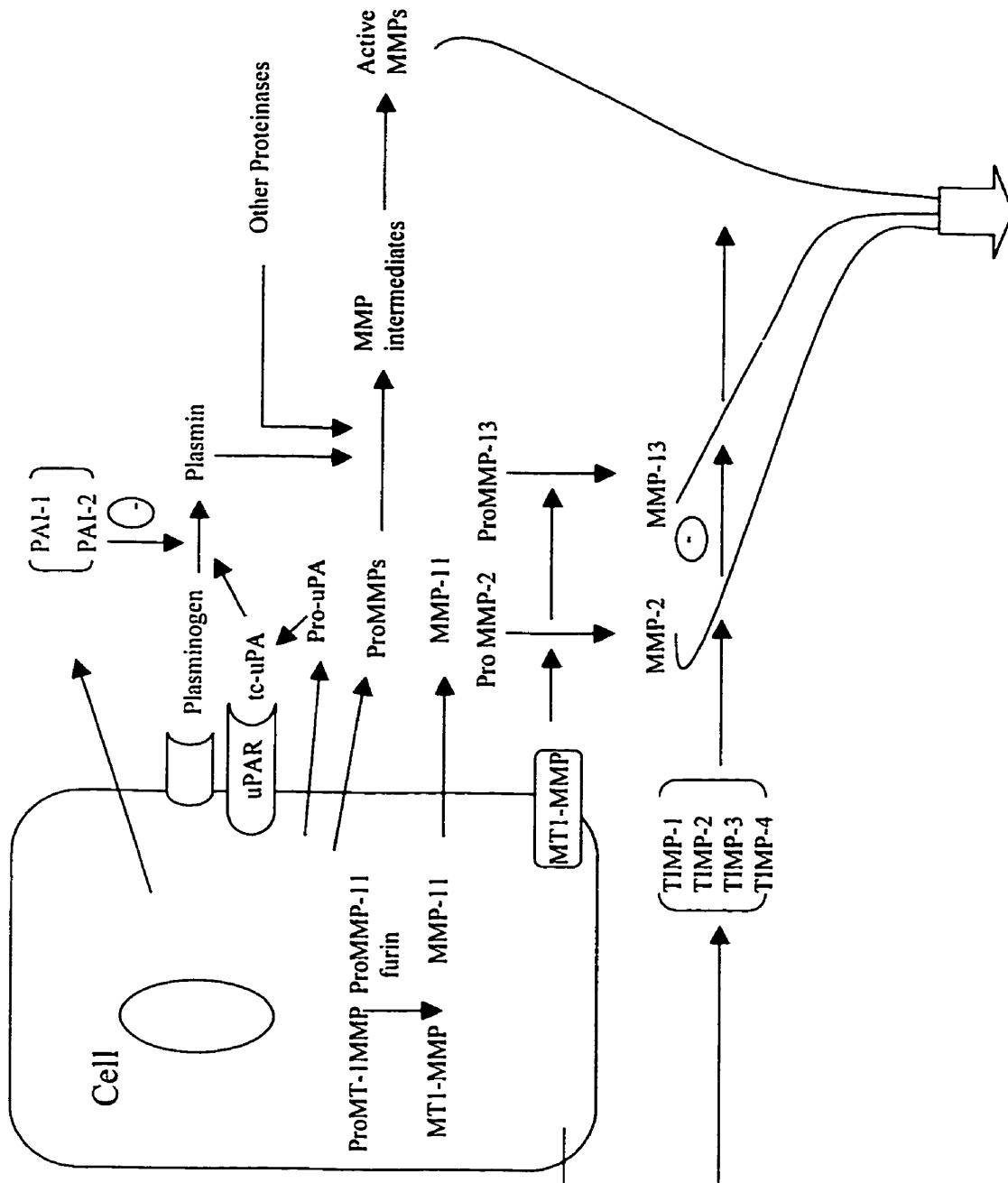


Figure 3. Proposed activation mechanisms of pro matrix metalloproteinases. uPAR, urokinase-type plasminogen activator receptor; tc-uPA, two chain urokinase type- plasminogen activator inhibitor. (Hidaeki Nagase. Activation Mechanisms of Matrix Metalloproteinases: A Review).

Methods of Detecting MMPs

Many techniques are employed to detect MMPs. These include immunohistochemistry, enzyme-linked immunosorbent assays (ELISA), western blotting, assays of mRNA levels and gelatin zymography. Gelatin zymography has the advantages of measuring enzymatic activity qualitatively and of distinguishing active from inactive enzymes.

1.8 Renin Angiotensin System (RAS)

Both the healthy and diseased heart use several neuroendocrine mechanisms (including RAS) to stabilize or enhance cardiac performance.¹⁷⁴ However, sustained activation leads to progressive myocyte dysfunction and myocyte hypertrophy, and seems to perpetuate hemodynamic abnormalities.¹⁷⁵ The renin angiotensin system consists of a cascade of enzymatic reactions that lead to the formation of angiotensin II (Fig.3), a potent vasoconstrictor important in the pathophysiology of cardiovascular diseases. Angiotensin II is formed primarily through the hydrolysis of angiotensin I by angiotensin converting enzyme (ACE). Angiotensin II can also be produced from angiotensin I through the action of non-ACE enzymes, such as chymostatin-sensitive angiotensin II generating enzyme, chymase, and cathepsin G.¹⁷⁶ Also, angiotensin can be produced directly from angiotensinogen with the aid of non renin enzymes such as tissue plasminogen activator (t-Pa) cathepsin G and tonin. The effects of angiotensin include: stimulation of adrenal medulla to secrete aldosterone, enhanced sodium and water retention, arterial vasoconstriction, and presynaptic release of norepinephrine from sympathetic nerve endings.¹⁷⁷ More important, there is evidence that angiotensin is a

growth factor and potent stimulator of myocyte hypertrophy and fibroblast proliferation.¹⁷⁸ Bienlich et al.¹⁷⁹ have demonstrated that angiotensin is required for the growth of the new born pig heart. Myocyte hypertrophy may initially augment cardiac function, but eventually the hypertrophied myocardium outgrows its blood supply and leads to further myocardial decompensation. Excessive fibroblast proliferation contributes to remodeling of the cardiac interstitium and changes in compliance. Recently, it has been shown that angiotensin stimulate collagen gel contraction in isolated heart fibroblasts in a dose dependent manner and that this response is inhibited by AT₁ receptor antagonist losartan.¹⁸⁰ Angiotensin has also been shown to stimulate increases in angiotensinogen¹⁸¹ and TGF- β ¹⁸² in the heart. The transcripts and corresponding translation products of endogenous angiotensinogen and renin have been identified in the myocardium, cultured cardiac myocytes and fibroblasts.¹⁸³ Angiotensin I, angiotensin II and ACE have been detected in cultured cardiac myocytes and fibroblasts.¹⁸⁴ Angiotensinogen mRNA levels are higher in cardiac atria than ventricles. ACE has been localized to the subendocardial tissues and is also found within sinoatrial and atrioventricular nodal regions. Using Langendorff perfused rat hearts, Lindpainter et al.¹⁸⁵ showed angiotensin I to angiotensin II conversion within these. They also demonstrated the synthesis of angiotensinogen in the hearts, its capability to respond to modulatory stimuli, and its local, specific enzymatic activation to biologically active peptides.¹⁸⁶ In the myocardium, abnormalities of angiotensin have been linked to progressive fibrosis, endothelial cell dysfunction, thrombosis and atherogenesis.^{187,188} Angiotensin is suggested to be a principal determinant of cardiac fibroblast hypertrophy/hyperplasia as well as enhanced collagen, fibronectin and integrin synthesis and deposition.¹⁸⁹

1.8.1 Evidence For the Putative Role of RAS in Cardiac Hypertrophy

Several lines of evidence support the role of angiotensin as a primary modulator involved in cardiac remodeling and pathological hypertrophy. Angiotensin peptides exert a number of physiologic effects on the heart, many of which may be directly or indirectly involved in the stimulation of cardiac growth. Among the direct actions of angiotensin peptides are positive inotropic¹⁹⁰ and chronotropic¹⁹¹ influences, coronary vasoconstriction,¹⁹² effect on myocardial metabolism,¹⁹³ proarrhythmic^{194,195} and potential proischemic^{194,196} actions, and growth promoting effects. Terminally differentiated cells, such as cardiac myocytes, have been demonstrated to respond with hypertrophic changes to angiotensin.¹⁹⁷ In a recent study, Mazzolami et al.¹⁹⁸ showed that transgenic mice, overexpressing the rat angiotensinogen gene in the heart developed myocardial hypertrophy without signs of fibrosis independently from the presence of hypertension. Experiments have also demonstrated that exposure to mechanical deformation, such as stretch or shear forces, induces the synthesis of angiotensin that is secreted into the culture medium and that subsequently exerts growth promoting effects in an autocrine fashion on the cells.³⁴ Angiotensin appears to mediate this hypertrophic response because the angiotensin receptor blocker prevents it. Several secondary pathways have been implicated in the trophic and proliferative properties of angiotensin:¹⁹⁹ expression of the proto-oncogenes c-myc, c-fos, and c-jun is induced by angiotensin, as is expression of other growth factors, such as the early growth response gene (egr-1), and cytokines such as TGF-β. These observations are in line with previously documented effects of angiotensin on DNA turnover, RNA turnover and content, and protein synthesis.²⁰⁰ Marked increases in angiotensinogen gene expression confined to the non infarcted,

acutely hypertrophic portion of the LV occurs in the early post MI phase.²⁰⁰ Similarly, ACE gene expression has been shown to increase during post MI-remodeling.²⁰² Angiotensin mRNA increases in LV exposed to pressure overload induced by abdominal aortic banding²⁰³ and the percent of myocytes containing renin, angiotensin I and angiotensin II was significantly increased in hypertrophied hearts.²⁰⁴ Both experimental and clinical studies have documented high efficacy of ACE inhibitors and AT₁ receptor antagonists to block cardiac hypertrophy and remodeling in several models of CHF.²⁰⁵⁻²⁰⁸ A long term study also showed that an ACE inhibitor was effective not only in improving survival but also in preventing²⁰⁹ heart failure suggesting that the RAS plays a key role in the pathophysiology of HF. Cardiac hypertrophy induced by pressure overload in the mouse is prevented by administration of an AT₁ receptor antagonist²¹⁰ indicating that angiotensin has a direct effect in mediating cardiac growth in this model. In aortic insufficiency model of volume overload, chronic captopril treatment was associated with reduced development of LV dilation and hypertrophy²¹¹ suggesting the role of RAS in this model. Angiotensin is shown to be involved in the development of cardiac hypertrophy in volume overload model (aortocaval shunt) by both hemodynamic and cardiac tropic effects. Ruzicka et al.²¹² showed an increase in plasma angiotensin I and angiotensin II as well as cardiac renin activity subsequent to induction of AV shunt in rats. The hypertrophic response in this model was blunted by using an AT₁ receptor antagonist.

1.8.2 Angiotensin Receptors in the Myocardium

Cellular responses induced by angiotensin are mediated by at least two types of

membrane receptors, AT₁ and AT₂ receptors,^{27,28,213} which are the only two human angiotensin receptors to have been cloned. The AT₁ subtype mediates the physiological actions of angiotensin, including hypertrophy, vasoconstriction and liberation of catecholamines from sympathetic nerve endings.²¹⁴ The AT₁ receptor is a member of G protein superfamily and is divided into AT_{1A}, AT_{1B} and AT_{1c} subclasses. Among them, the AT_{1A} and AT_{1B} isoforms contain 18 to 22 different aminoacids yet maintaining similar binding profiles for angiotensin and nonpeptide antagonists (including losartan) as well as for AT₁ receptor antagonists. The AT₁ receptors predominate in the cerebral cortex, vasculature, kidney, and liver while the AT₂ receptor is present primarily in adrenal medula, uterus, ovary, brain and developing foetus. It has been shown that the AT₁ receptor contributes to the regulation of blood pressure in AT_{1A} receptor deficient mice.²¹⁵ Further, AT₂ receptor knockout mice exhibited significantly increased blood pressure, increased sensitivity to pressor action of angiotensin and altered exploratory behavior, suggesting a role of AT₂ in brain function and blood pressure control.^{216,217} Cardiac AT₁ receptor density has been found to be increased in hypertrophy models²¹⁸ or after myocardial infarction.²¹⁹ Similarly, atrial hypertrophy in cardiac volume overload showed a parallel increase in AT₁ mRNA content.²²⁰ Booz and Baker²²¹ have demonstrated that the AT₁ receptor has a growth stimulatory effect that depends on receptor number, whereas the AT₂ receptor suppresses growth of cultured cardiomyocytes. Because AT₁ receptor mediated signals induce myocyte hypertrophy and proliferation of fibroblasts, it is suggested that increased AT₁ receptor density in such pathological states enhances the action of angiotensin. The AT₂ receptor has an antiproliferative effect on neointima formation²²² and on coronary endothelial cells²²³ and

are involved in apoptosis.²²⁴ Accumulated evidence analyzing human hearts has reported that both AT₁ and AT₂ receptors are expressed in human atria and ventricles. One study has shown that AT₂ receptor sites were increased by 150% during heart failure while AT₁ receptor sites were increased in the hypertrophic state and then decreased to the control level during heart failure.²²⁵ Also, both AT₁ and AT₂ receptors are localized at higher densities in the fibroblasts present in fibrous regions.²²⁶ AT₂ receptor suppressed synthesis and secretion of major matrix components such as fibronectin and collagen type I. In fibroblasts isolated from normal adult rat hearts, Brilla et al²²⁷ reported that high concentrations of angiotensin stimulated collagen synthesis by both AT₁ and AT₂ receptors and that AT₂ receptor decreased collagenase activity resulting in increased collagen synthesis. One recent study using both AT₁ and AT₂ receptor antagonists showed that AT₁ receptor antagonist and not AT₂ receptor antagonist prevented progression of heart failure in cardiomyopathic hamsters.²²⁸

1.9 Blockade Of The Renin Angiotensin System

1.9.1 Blockade of Angiotensin Formation: Renin and ACE inhibitors - ACE inhibitors act by blocking the conversion of angiotensin I to the metabolically active angiotensin II. In recent years, ACE blockade has been the predominant mechanism for blocking the actions of angiotensin II.^{229,230} However, inhibition at this site is not optimal; in addition to acting on angiotensin I, ACE also degrades bradykinin and other peptides that are substrates for the enzyme. Therefore, ACE inhibition results in bradykinin accumulation, which can then stimulate the production of nitric oxide and prostacyclin. Many of the adverse effects associated with ACE inhibitors, including cough and angioneurotic edema

are believed to be a consequence of interrupting the metabolism of peptides such as bradykinin, substance P, or tachykinin that accumulate as a result of inhibiting their breakdown. Furthermore, plasma levels of angiotensin after chronic administration of ACE inhibitors tend to return towards normal.²³¹ The rise in angiotensin levels may be caused by angiotensin produced by systems other than the RAS, which are unaffected by ACE inhibition (Fig.4). Renin inhibition and hence, inhibition of the conversion of angiotensinogen to Ang I, is a potential method of blockade of angiotensin formation. However, to date renin inhibitors are not clinically available.

1.9.2 Angiotensin receptor blockers- these agents block the action of angiotensin at the receptor level and conceivably could block the effects of angiotensin not only through classical ACE pathway but also by the chymase pathway. These antagonists compete with angiotensin for its specific receptors, AT₁ or AT₂. Unlike the situation with ACE inhibitors, bradykinin metabolism is not affected, and therefore the actions of bradykinin are not potentiated. Losartan (DUP 753, MK-954, or Cozaar; Merck Research laboratories, West Point, PA), a non peptide, biphenylimidazole potassium salt is the first agent in this class of angiotensin receptor blockers. The Elderly (Elite)²³² trial compared the effectiveness of losartan with captopril (ACE inhibitor). Patients experiencing adverse effects were fewer in losartan treated group (12%) as compared to captopril receiving group (29%). Also, death, hospital admission for heart failure, or both occurred in 9.4% patients treated with losartan and 13.2% of patients treated with captopril. AT₁ receptor could therefore provide the same beneficial effects as ACE with fewer side effects.

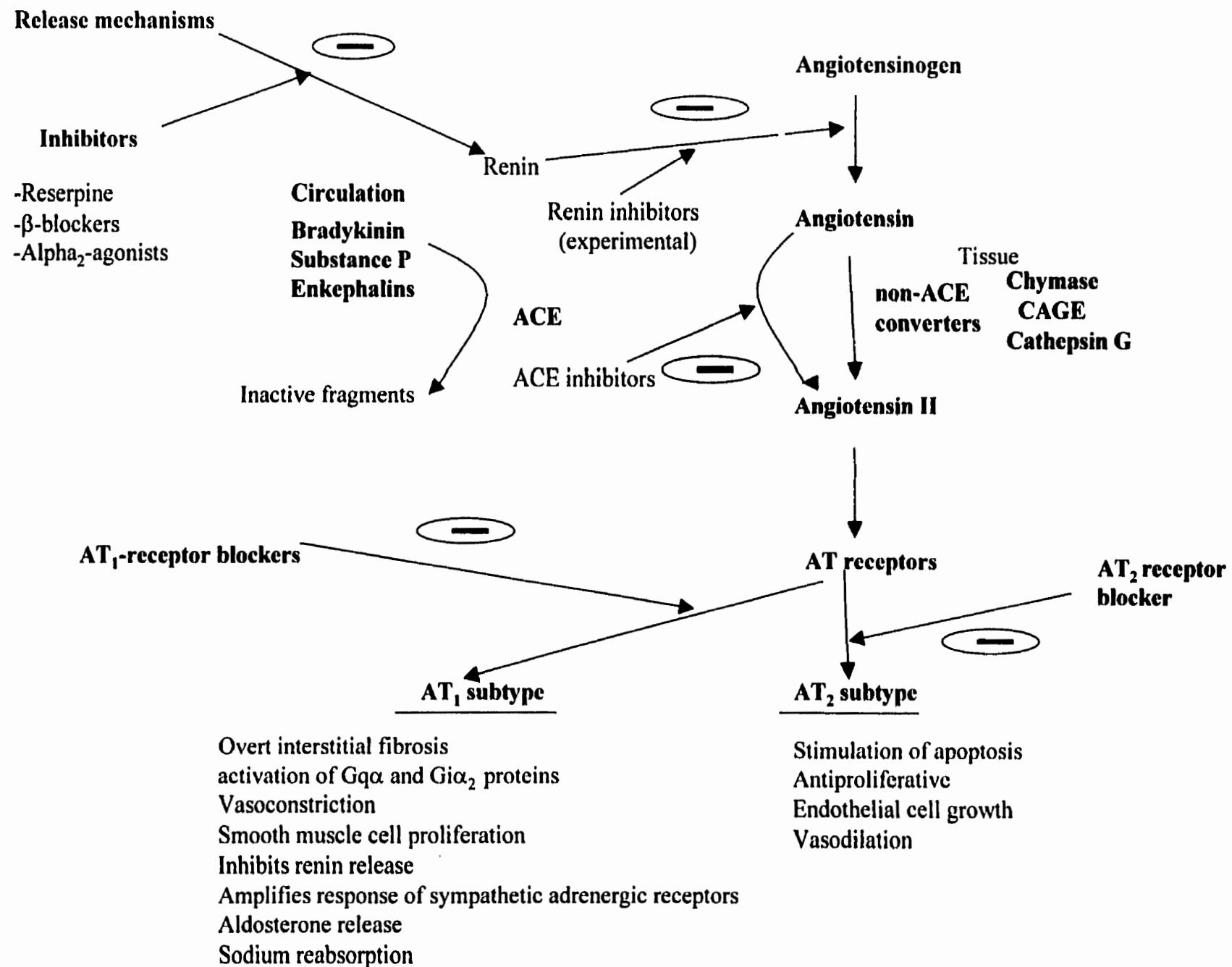


Figure 4. Diagram of the renin-angiotensin system, showing putative physiologic and pharmacologic interactions.

IV. RATIONALE, HYPOTHESIS, AND OBJECTIVES

Elevated expression of interstitial collagens in response to pressure overload and in post-MI heart have been extensively studied and are well documented.²³³⁻²³⁶ These changes are associated with increased myocardial stiffness and may contribute to ventricular dysfunction. These studies have also suggested the involvement of trophic stimuli such as angiotensin and aldosterone in the regulation of connective tissue growth. On the other hand, minimal data are available to address the effect of cardiac volume overload on the composition of the myocardial interstitium. In a study of the canine heart, cardiac volume overload by aortocaval (AC) shunting for 2 months resulted in no significant changes in the left ventricular (LV) collagen fraction.²³⁷ Similar results were reported by Michel *et al.*²³⁸ who also found no change in LV collagen density after 1 or 3 months of cardiac volume overload by AC shunt in rats. The collagen concentration was reported to be increased in a cat model of right ventricular volume overload due to atrial septal defect.²²⁹ Similarly, in a pig model of AV shunt, Harper *et al.*²⁴⁰ found a slight increase in collagen concentration, 4-6 weeks after induction of shunt. However, Ruzicka *et al.*²⁴¹ reported decreases in LV collagen after 4 to 10 weeks of volume overload by AC shunting in rats. Therefore, the data concerning the alterations in collagen proteins are controversial. Furthermore, although MMP activation has been demonstrated to play a significant role in tissue remodeling in a number of pathological processes such as tumor metastasis,²⁴² pacing induced HF,²¹ CMP,^{17,18} post-MI (scar),¹⁹ there is as yet no information as to whether altered MMP activity and abundance are involved in the volume overload induced CHF process.

We hypothesized that ventricular dilation in experimental volume overload is due to alteration of the cardiac collagen weave via the activation of ventricular MMPs. As ventricular dilatation may contribute to the pathogenesis of heart failure in volume overloaded hearts, these events are implicated in the progression of heart failure in this experimental model. In view of this, the main objective of this thesis was to determine whether any alteration occurs in the cardiac collagen in experimental volume overload heart failure and whether cardiac MMP activity and/or protein abundance is altered in these experimental hearts. We also tested whether altered MMP activity or expression in AV overloaded hearts is regulated by AT₁ receptor activation. Thus a relationship between collagen concentration in AVF hearts, collagen removal by MMPs, and their reliance of this system on AT₁ activation was sought.

V. MATERIALS AND METHODS

1. Experimental Animals

The experimental protocols for animal studies were approved by the Animal Care Committee of the University of Manitoba, following guidelines established by the Medical Research Council of Canada and the Canadian Council on Animal Care. Male Sprague-Dawley rats weighing 200-250 gms were used for the study. Animals were housed in solid bottom polycarbonate cages, were fed laboratory grade rat chow and sterilized water ad libitum, and were subjected to 12 hours of light and darkness throughout the housing period.

Volume overload was produced by surgical creation of aortocaval shunt as described previously by Garcia and Diebold.⁴⁴ The skin was sterilized with povidone-iodine solution before a mid-line celiotomy was performed. After isofluorane anesthesia, the vena-cava and abdominal aorta were exposed by opening the abdominal cavity via a mid-line incision. The intestines were displaced laterally using sterile sponges. Blunt dissection was used to remove the overlying adventitia and expose the vessels, taking care not to disrupt the tissue connecting the vessels. Both vessels were then occluded proximal and distal to the intended puncture site. An 18 gauge needle was inserted into the exposed abdominal aorta and advanced through the medial wall into the vena-cava to create the fistula. The needle was then removed and the puncture point sealed by a drop of cyanoacrylate glue. The abdominal musculature and skin incisions were closed by standard technique with absorbable suture and autoclips. Throughout the surgery, ventilation of the lung was maintained by positive pressure inhalation of 95% oxygen and

5% carbon dioxide mixed with isofluorane. Sham-operated rats were treated similarly except that no puncture was made. The animals were divided into three groups: sham operated animals, shunt animals and shunt animals treated with losartan (40 mg/kg/day, gastric gavage). Drug treatment started 1 day after the shunt surgery. The animals were sacrificed by decapitation at either 1 week or 8 week subsequent to the surgery. The hearts were snap-frozen in liquid nitrogen and stored at -70°C for further analysis.

2. Hemodynamic Measurements

Mean arterial pressure (MAP) and left ventricular function of sham-operated control, shunt and shunt treated with losartan groups were measured following induction of shunt, as described previously. Briefly, rats were anesthetized by intra-peritoneal injection of a Ketamine:Xylazine mixture (100mg/kg:10mg/kg). A micromanometer tipped catheter (2-0) (Millar SPR - 249) was advanced into the aorta to determine mean arterial pressure (MAP), and then further advanced to the left ventricular chamber to record left ventricular systolic pressure (LVSP), left ventricular end diastolic pressure (LVEDP), the maximum rate of isovolumic pressure development ($+dp/dt_{max}$) and the maximum rate of isovolumic pressure decay ($-dp/dt_{max}$). Hemodynamic data was computed instantaneously and displayed using a computer data acquisition work station (Biopac, Harvard Apparatus Canada). In another series of experiments, left ventricular function and blood pressure of sham, shunt and losartan treated rats were measured 8 weeks following induction of volume overload.

3. Protein Assay

Total protein concentration in cardiac samples was determined using the Bicinchoninic acid solution (BCA) kit (Sigma, St. Louis USA).

4. Zymography: Detection of Cardiac Matrix Metalloproteinase Activity

The tissue samples were homogenized in 100 mM Tris pH 7.4 containing 1mM PMSF, 4 µM leupeptin, 1 µM pepstatin A, and 0.3 µM aprotinin and were then sonicated for 3 X 5 seconds. Crude membrane and cytosolic fractions were isolated according to the methods of Gettys et al.²⁴³ Briefly, ventricular (left- and right) samples were centrifuged for 3000 x g at 4°C for 10 min. to remove broken cells and nuclei. The supernatant was further subjected to centrifugation for 48,000 x g for 20 min. at 4°C. The cytosolic fraction (Supernatant) was separated from the crude membrane pellet. Total protein concentration of cytosolic fractions was measured using a standardized colorimetric assay (BCA assay). These extracts were aliquoted, flash - frozen using liquid nitrogen, and stored at -80°C until the time of assay.

Cardiac MMP activity was determined by zymography.²⁴⁴ Gelatin (final concentration 1mg/ml) was added to the gel made with standard 7.5% SDS polyacrylamide and this gel was subjected to electrophoresis. 30µg of protein was loaded per lane without reduction or boiling (to maintain MMP activity). Samples were run at 200 volts, maintaining a running buffer temperature of 4°C. After electrophoresis, gels were washed twice for 30 min. each in 25 mM glycine (pH 8.3) containing 2.5% Triton X-100 with gentle shaking at 4°C in order to eliminate SDS. After incubating in incubation solution (50mM Tris pH 8.0,10mM CaCl₂) for 18 hours, gels were stained in

0.1% Coomassie Brilliant blue R- 250, for 30 minutes and then destained in acetic acid and methanol. Gels were dried and scanned using a CCD camera densitometer (Bio-Rad imaging densitometer GS 670) for determining relative lytic activity.

5. Determination of Total Cardiac Collagen

Right and left ventricular samples from different groups were ground into powder in liquid nitrogen. 100 mg (wet weight) cardiac tissue was then dried to constant weight. Tissue samples were digested in 6N HCl (6ml/100mg dry wt.) for 16 hrs at 105°C. Hydroxyproline was measured according to the method described by Chiariello et al.²⁴² A stock solution containing 40mM of 4-hydroxyproline in 1mM HCl was used as standard. Collagen concentration was calculated by multiplying hydroxyproline levels by a factor of 7.46, assuming that interstitial collagen contains an average of 13.4% hydroxyproline.²⁴⁵ The data was expressed as µg collagen per mg dry tissue.

6. Western Blot Analysis: Quantification of Cardiac Matrix Metalloproteinases

Western blot analysis was performed on tissue samples. Crude membrane and cytosolic fractions were isolated according to the methods of Gettys *et al.*²⁴³ Briefly, left and right ventricular samples were centrifuged for 3000 x g at 4°C for 10 min. to remove broken cells and nuclei. The supernatant was further subjected to centrifugation for 48,000 x g for 20 min. at 4°C. The cytosolic fraction (Supernatant) was separated from the crude membrane pellet. Total protein concentration of cytosolic fractions was measured using the BCA assay. Prestained low molecular weight marker and 30µg of protein from samples was separated on 10% SDS-PAGE, after reduction (100°C, 5min.).

Separated proteins (cytosolic) were transferred to 0.45 μ m polyvinylidene difluoride (PVDF) membrane. The membrane was blocked overnight at 4°C or at room temp. for 1 hour in Tris-buffered saline with 0.1% Tween-20 (TBS-T) containing 6% skim milk. After washing with TBS-T solution, membranes were probed with primary antibodies for 1 hour at room temperature. Primary antibodies used were; MMP-1 antibody (1:1000 dilution, The Binding Site, Birmingham, UK), MMP-2 antibody (1:500 dilution, Biogenesis, Poole, England) and MMP-9 antibody (1:500 dilution, Biogenesis, Poole, England). After washing, membranes were incubated with horseradish peroxidase (HRP) labeled secondary antibodies (horseradish peroxidase labeled Goat Anti-Rabbit IgG, 1:10,000 dilution) for 1 hour at room temperature. The targeted proteins were detected and visualized using enhanced chemiluminescence (ECL) "plus" according to manufacturer's instructions (Amersham Life Science Inc., Canada). Autoradiographs from western blot were quantified using a CCD camera imaging densitometer (Bio-Rad GS670, CA, USA).

7. Trichrome staining for total cardiac matrix

Serial cryostat cardiac sections from ventricular apexes were mounted on paraffin fixed, saline coated slides. Slides were deparaffinized and hydrated in distilled water. Fixing was done by immersing slides in Bouin's solution (saturated Picric acid & supernatant, 37-40% formaldehyde, Glacial acetic acid) for 1 hr. in a 60°C water bath. Once fixing was accomplished, slides were cooled and washed in running tap water for 5 min and then rinsed with distilled water. The slides were then placed in working hematoxilin solution (Hematoxylin, 95% ethanol, 29% aqueous Ferric Chloride, HCl) for

10 min and then rinsed in distilled water, washed in running tap water for 5 min and rinsed again. Slides were immersed in Biebrich-Acid Fuchsin solution (1% Biebrich Scalet, 1% Acid Fuchsin, Glacial Acetic acid) for 15 minutes. Slides were rinsed and placed in Phosphomylbdic-Phosphotungstic acid solution (Phosphomylbdic Acid, Phosphotungstic Acid) for 10 min, transferred to Aniline blue solution (Analine Blue, Glacial Acetic Acid) for 5 min and rinsed. Slides were placed in aqueous acetic acid solution for 3 min. followed by dehydration and mounting in permount. As a result, nuclei were stained black; cytoplasm, keratin, muscle and intracellular fibers stained red; collagen and mucus stained blue.

8. Reverse Zymography: Detection of MMP inhibitory (TIMP) activity

MMP inhibitory activity was assayed in 12% polyacrylamide gels including 1mg/ml gelatin. Prestained low molecular weight marker and 30 µg of protein from samples was loaded per lane without reduction. After electrophoresis, gels were washed twice (30 min each wash) in 25 mM Glycine (pH 8.3) containing 2.5% Triton X-100 and incubated at 37°C for 48 hours in a preparation of 10U/ml gelatinase (*Clostridium histolyticum*; Worthington Biochemical Corp., NJ, USA). After incubation, gels were stained using 0.1% Coomassie blue R-250 in a mixture of acetic acid:methanol:water (1:3:6) and destained in the same mixture without the dye. Gels were dried and scanned using a CCD Camera densitometer (Bio-Rad imaging densitometer GS 670).

9. Statistical Analysis

Assays were conducted in a completely randomized fashion in accordance with

methods in parametric statistics. All values were expressed as mean (S.E.M.) Differences between groups were assessed by one way analysis of variance (ANOVA) followed by Student-Newman-keuls test (SigmaStat) for significance of differences between data sets. Significant differences among groups were defined by a probability of less than 0.05.

VI. RESULTS

1. Hemodynamic Changes

Animals were assessed for left ventricular function at 8 weeks post AV shunt surgery. The data revealed an increase in left ventricular end diastolic pressure (LVEDP) and a decrease in $\pm dP/dt_{max}$ relative to their controls (Table 3). Rats with shunt had significantly lower left ventricular systolic (LVSP) pressure at 8 weeks after creation of shunt. Systolic and diastolic pressures corresponded with changes in LVSP and were significantly lower in rats with 8 week shunt compared with controls. Heart rate (HR) showed no significant changes in the two groups (Table 4). Lung congestion was noted by the increase in lung weight in shunt animals compared to controls.

2. General characteristics and cardiac anatomy

Hearts of experimental animals were significantly hypertrophied, as reflected by increased LV mass and also by the increased values for LVW/BW and RVW/BW ratios in 8 week experimental animals compared to controls (Table 3). Liver and kidney weights remained unchanged in experimental hearts vs. control values.

3. Effect of losartan treatment

Hemodynamic changes

Losartan treatment was initiated 1 day after shunt surgery and was associated with reduced LVEDP in shunt rats. LVSP was significantly elevated by 8 week losartan

Table 3. General characteristics of volume-overloaded rats with or without treatment of losartan at 8 weeks after aortocaval shunt surgery

	Sham	Shunt	Shunt+Los
BW (g)	501.1 ± 13.7	493 ± 12.6	483.9± 17.9
HW (mg)	1307 ± 89	1898 ± 145*	1493 ± 105**
LVW (mg)	1053 ± 77	1490 ± 102*	1205 ± 104*#
RVW (mg)	248 ± 10	408 ± 388*	289 ± 21*#
HW/BW (mg/100g)	261 ± 15	385 ± 34*	309 ± 258*#
LVW/BW (mg/100g)	211 ± 10	302 ± 23*	250 ± 16*#
RVW/BW (mg/100g)	50 ± 3.5	84 ± 7.5*	61 ± 5.2*#
Lung Weight (g)	1.78 ± 0.08	2.04 ± 0.06*	1.82 ± 0.13#
Liver Weight (g)	23.1 ±2.09	21.6 ± 1.91	22.5 ± 1.56
Kidney Weight (g)	1.8 ± 0.13	1.76 ± 0.15	1.79 ± 0.12

Values are mean±SE (n=6-8per group). BW: body weight; HW: heart weight; LVW: left ventricular weight; RVW: right ventricular weight; Los: Losartan. *P<0.05 compared with sham control; #P<0.05 compared with shunt group.

Table 4. Hemodynamic characteristics of volume-overloaded rats with or without treatment of losartan at 8 weeks after aortocaval shunt surgery

	Sham	Shunt	Shunt+Los
LVEDP (mmHg)	6.1 ± 0.57	13.9 ± 1.08*	9.1 ± 0.58
LVSP (mgHg)	134.8 ± 4.3	118.1 ± 3.8*	124.9 ± 5.5*
HEART RATE	297 ± 15	321 ± 20	308 ± 14
+dP/dT (mmHg/sec)	5746 ± 329	4711 ± 130*	5360 ± 317#
-dP/dT (mmHg/sec)	5862 ± 356	4664 ± 134*	5793 ± 358#
ASP	125 ± 5.9	101 ± 2.6*	110 ± 6.5*#
ADP	97.8 ± 3.8	65.2 ± 3.1*	78.3 ± 5.2*#
APP	29.3 ± 2.1	48.6 ± 4.2*	33.0 ± 2.7#

95

Values are mean±SE (n=6-8per group). LVEDP: left ventricular end diastolic pressure; LVSP: left ventricular systolic pressure; +dP/dt: maximum rate of isovolumic pressure development; -dp/dt: maximum rate of relaxation; ASP: arterial systolic pressure; ADP: arterial diastolic pressure; APP: arterial pulse pressure. *P<0.05 compared with sham control values;#P<0.05 compared with shunt group values.

treatment, but was still significantly lower than control values (Table 4). Losartan treatment was associated with modest normalization of systolic and diastolic pressures; nonetheless these values still remained significantly lower than the control group values. Losartan treatment for 8 weeks was also associated with significantly increased $\pm dp/dt_{max}$ vs. untreated shunt animals.

General characteristics and cardiac anatomy

Losartan treatment for 8 weeks was characterized by attenuation of the HW, LVW and RVW. However, these values remained significantly higher vs. control values. The ratios of HW/BW, LVW/BW and RVW/BW were also significantly reduced in the treatment group but were still significantly higher vs. controls. Lung weight was normalized by 8 week losartan treatment while losartan treatment was observed to have no effect on heart rate (Table 3).

4. Myocardial Zymographic activity

In order to examine whether changes in ventricular myocardial collagen degradative processes occurred during the progression of AV shunt induced moderate HF, MMP zymographic activity was determined from LV and RV myocardial extracts at 1 and 8 weeks of AV shunt A representative Zymogram for 1 and 8 weeks after creation of AV fistula using gelatin as a proteolytic substrate is shown in fig 5.

MMP-1, MMP-2 and MMP-9 Zymographic activities could be identified on the basis of calibrated molecular weight markers, which were included in each zymogram. The Zymograms were subjected to densitometric analysis in order to determine total proteolytic activity.

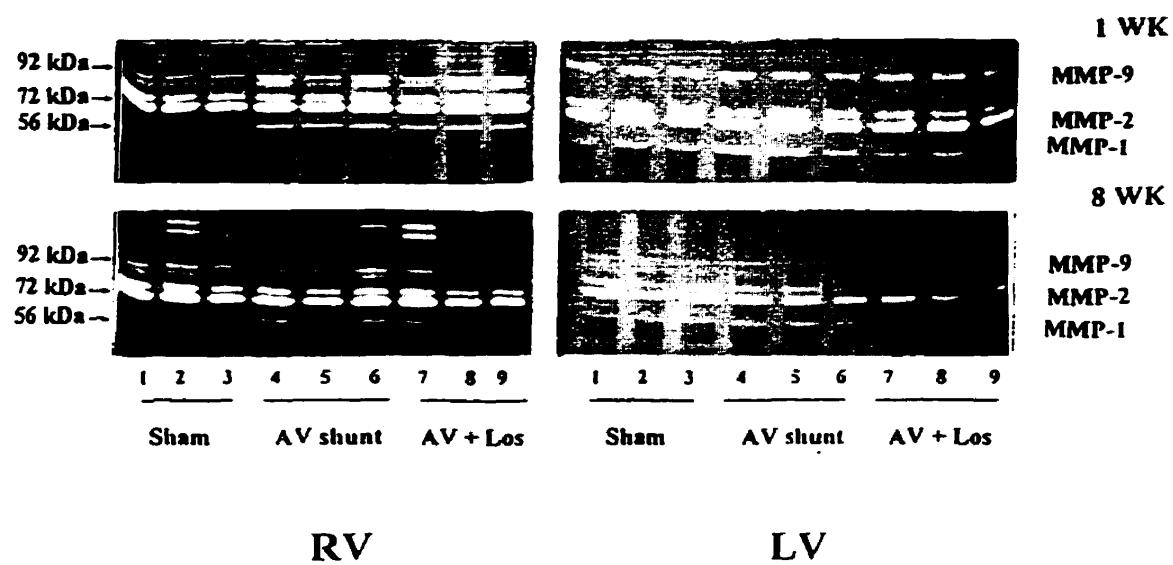


Figure 5. Representative zymography of cardiac matrix metalloproteinase (MMP) activity in 1 and 8 week ventricular (LV and RV) tissue samples from AV shunt animals, AV shunt animals treated with losartan (AV + Los) and age matched sham-operated controls (Sham). MMP-1 (56 kDa), MMP-2 (72 kDa) and MMP-9 (92 kDa) bands are indicated on the basis of calibrated molecular weight markers included in each zymogram.

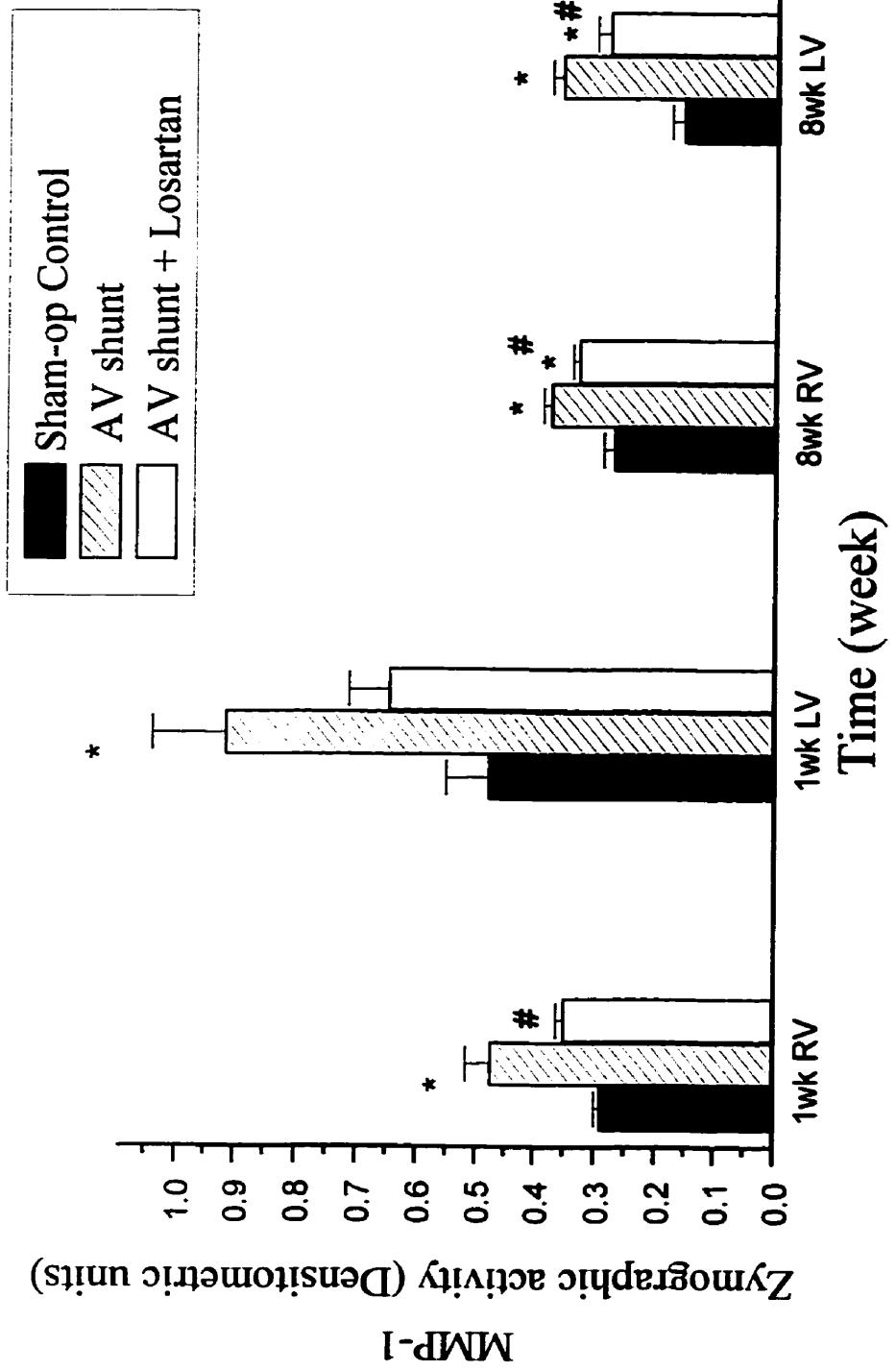


Figure 6. Estimation of the relative activity of left and right ventricular matrix metalloproteinase 1 (MMP-1) in 1 and 8 week myocardial samples taken from experimental animals (AV shunt) or AV shunt animals treated with losartan (AV shunt + Losartan) compared to age matched sham-operated controls. The data depicted is the mean \pm S.E.M of 4-7 experiments. *P<0.05 vs. control value. #P<0.05 vs. AV shunt value.

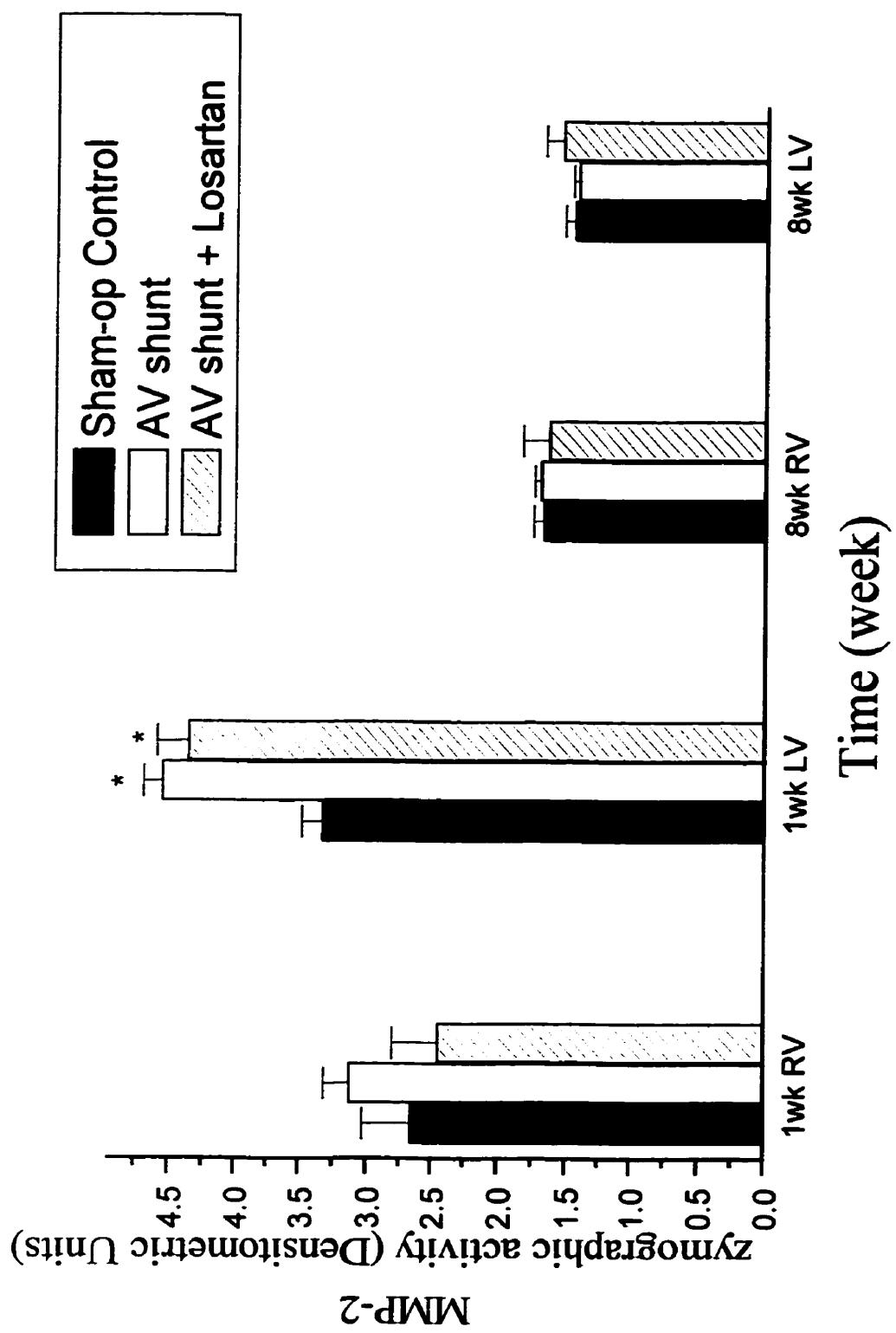


Figure 7. Estimation of the relative activity of left and right ventricular matrix metalloproteinase 2 (MMP-2) in 1 and 8 week myocardial samples taken from experimental animals (AV shunt) or AV shunt animals treated with losartan (AV shunt + Losartan) compared to age matched sham-operated controls. The data depicted is the mean \pm S.E.M of 4-7 experiments. *P<0.05 vs. control value. #P<0.05 vs. AV shunt value.

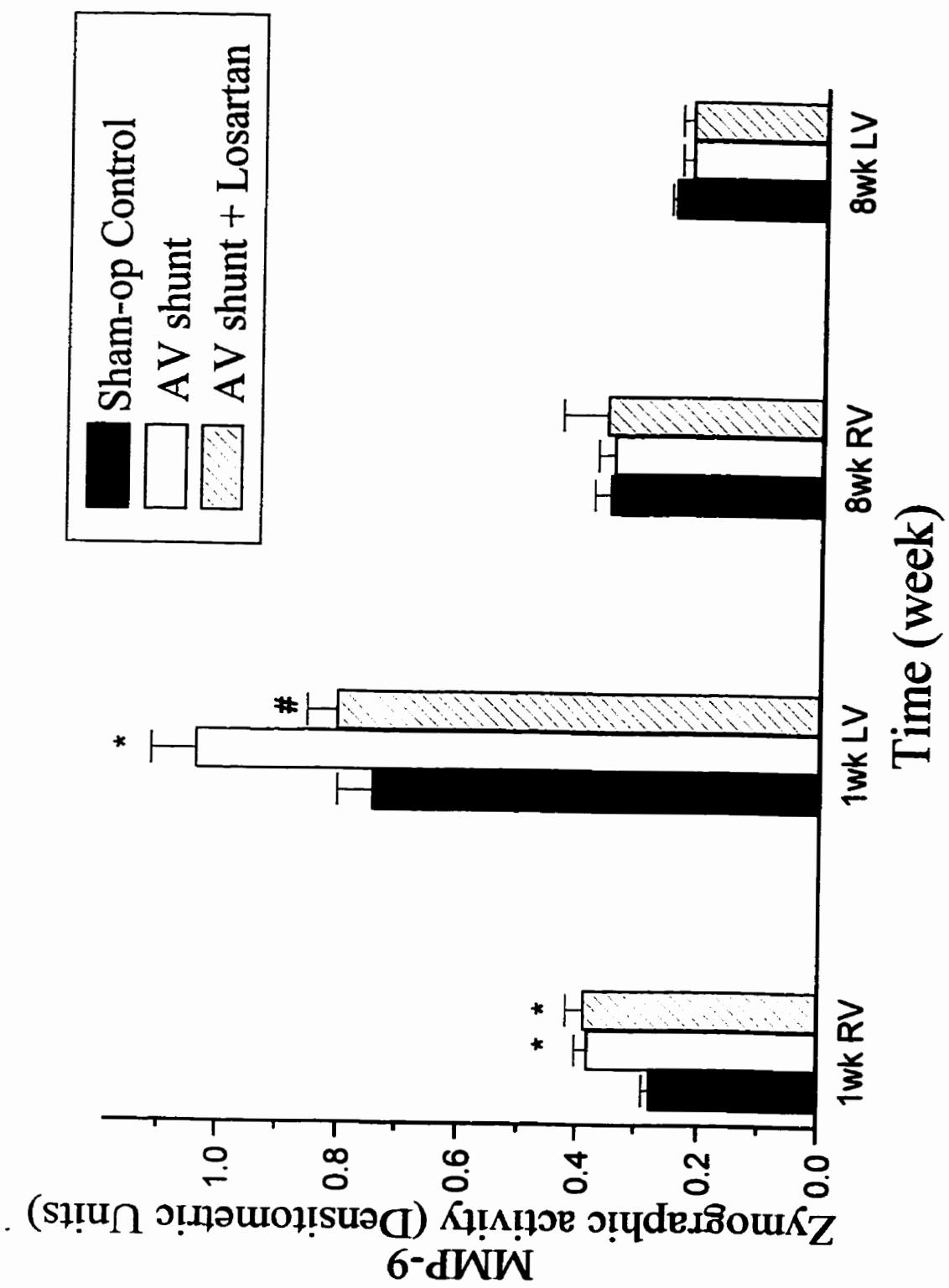


Figure 8. Estimation of the relative activity of left and right ventricular matrix metalloproteinase 9 (MMP-9) in 1 and 8 week myocardial samples taken from experimental animals (AV shunt) or AV shunt animals treated with losartan (AV shunt + Losartan) compared to age matched sham-operated controls. The data depicted is the mean \pm S.E.M of 4-7 experiments. *P<0.05 vs. control value.
#P<0.05 vs. AV shunt value.

1 week group

For left ventricles, MMP (MMP-1, MMP-2 and MMP-9) gelatinolytic activity showed significant increases in AV shunt group compared to sham operated control group. Losartan treatment for 1 week was associated with significant improvement in MMP-9 activity but had no effect on MMP-1 and MMP-2 activities. For right ventricles, there was no significant change in MMP-2 activity while MMP-1 and MMP-9 activities were significantly increased in AV shunt group compared to sham controls. 1 week losartan treatment had no effect on MMP-1, MMP-2 or MMP-9 activities (Figs. 5-8).

8 week group

At 8 weeks after the creation of AV shunt, the LV and RV myocardial MMP-1 activity was significantly increased compared to control group while MMP-2 and MMP-9 activities showed no significant differences between the two groups. Losartan treatment for 8 weeks was associated with significant attenuation in both left and right ventricular MMP-1 activity compared to AV shunt group which however still remained significantly higher than control values (Figs. 5-8).

5. Immunoreactive Myocardial MMP abundance

In order to determine whether the progression of AV shunt induced CHF is accompanied by change in MMP abundance, immunoblotting was performed for interstitial collagenase (MMP-1), the 72-kDa gelatinase (MMP-2) and the 92 kDa gelatinase (MMP-9). A representative immunoblot for these specific MMP species at both time groups studied, is shown in figs. 9, 11 and 13. In all LV and RV myocardial samples, a distinct immunoreactive band could be localized to the appropriate molecular

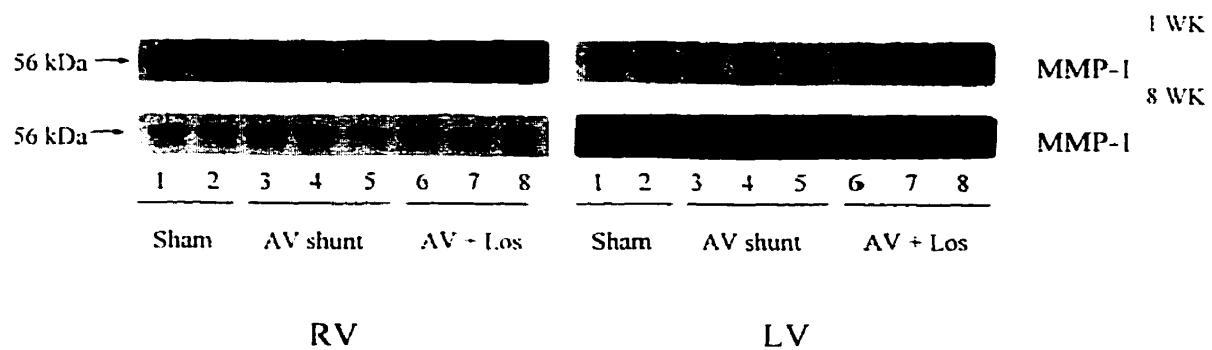


Figure 9. Representative Western autoradiograph showing bands for immunoreactive cardiac MMP-1 (56 kDa) in 1 and 8 week ventricular (LV and RV) tissue samples from experimental (AV shunt) animals, experimental animals treated with losartan (AV + Los) and respective age-matched sham-operated control (sham) animals.

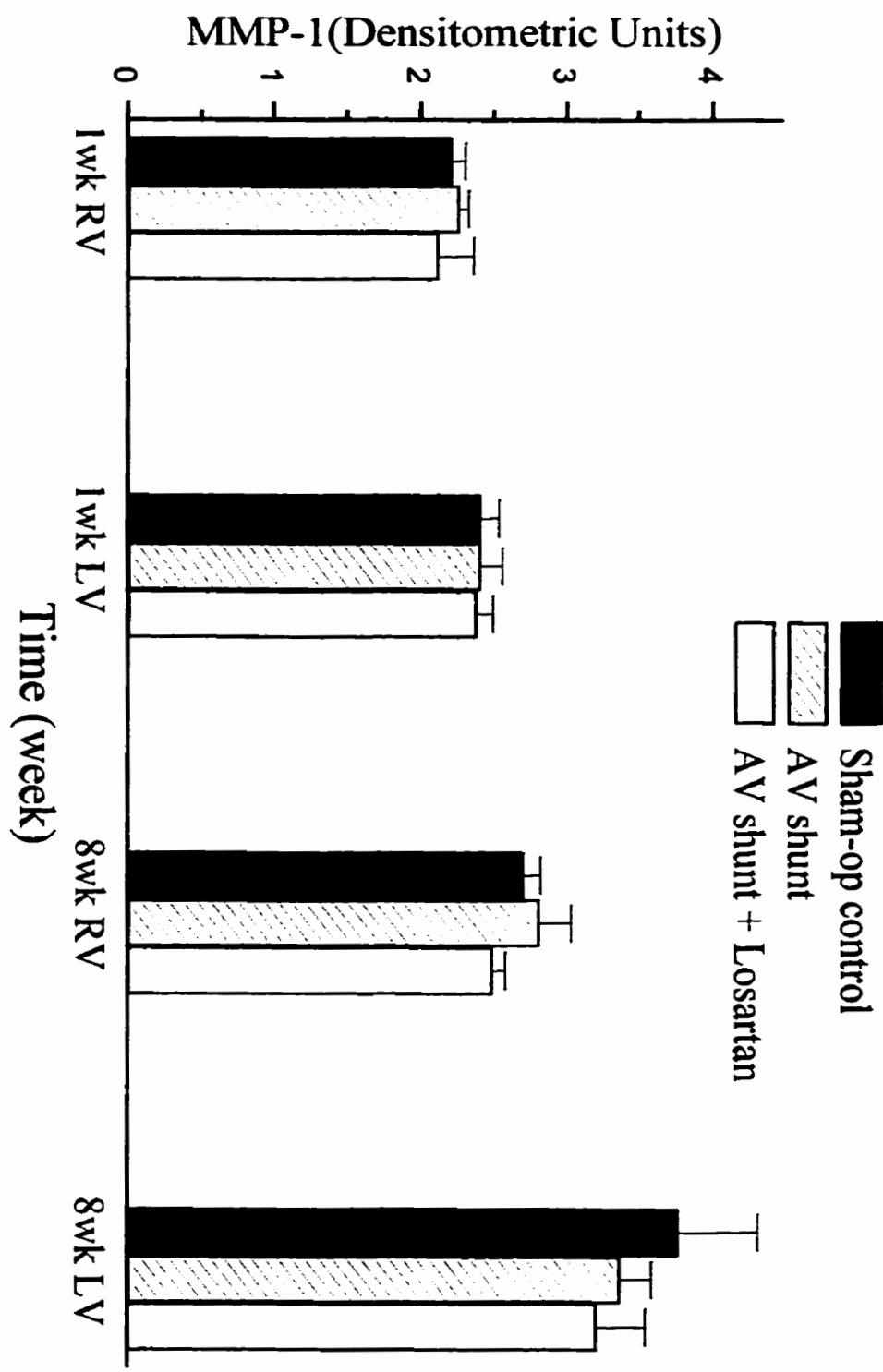


Figure 10. Western blot analysis for immunoreactive cardiac MMP-1 in left and right experimental animals (AV shunt), AV shunt animals treated with losartan (AV + Losartan) and sham-operated controls, 1 and 8 weeks after AV shunt induction. The quantified data depicted is the mean \pm S.E.M. of 3-4 experiments.
 * $P<0.05$ vs. control value. # $P<0.05$ vs. AV shunt value.

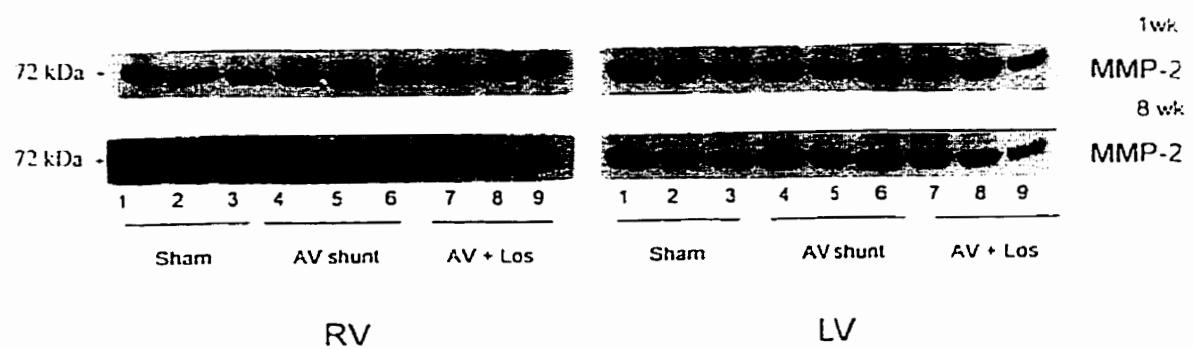


Figure 11. Representative Western blot showing bands for immunoreactive cardiac MMP-2 (72 kDa) in 1 and 8 week ventricular (LV and RV) tissue samples from experimental (AV shunt) animals, experimental animals treated with losartan (AV + Los) and respective age-matched sham-operated control (Sham) animals.

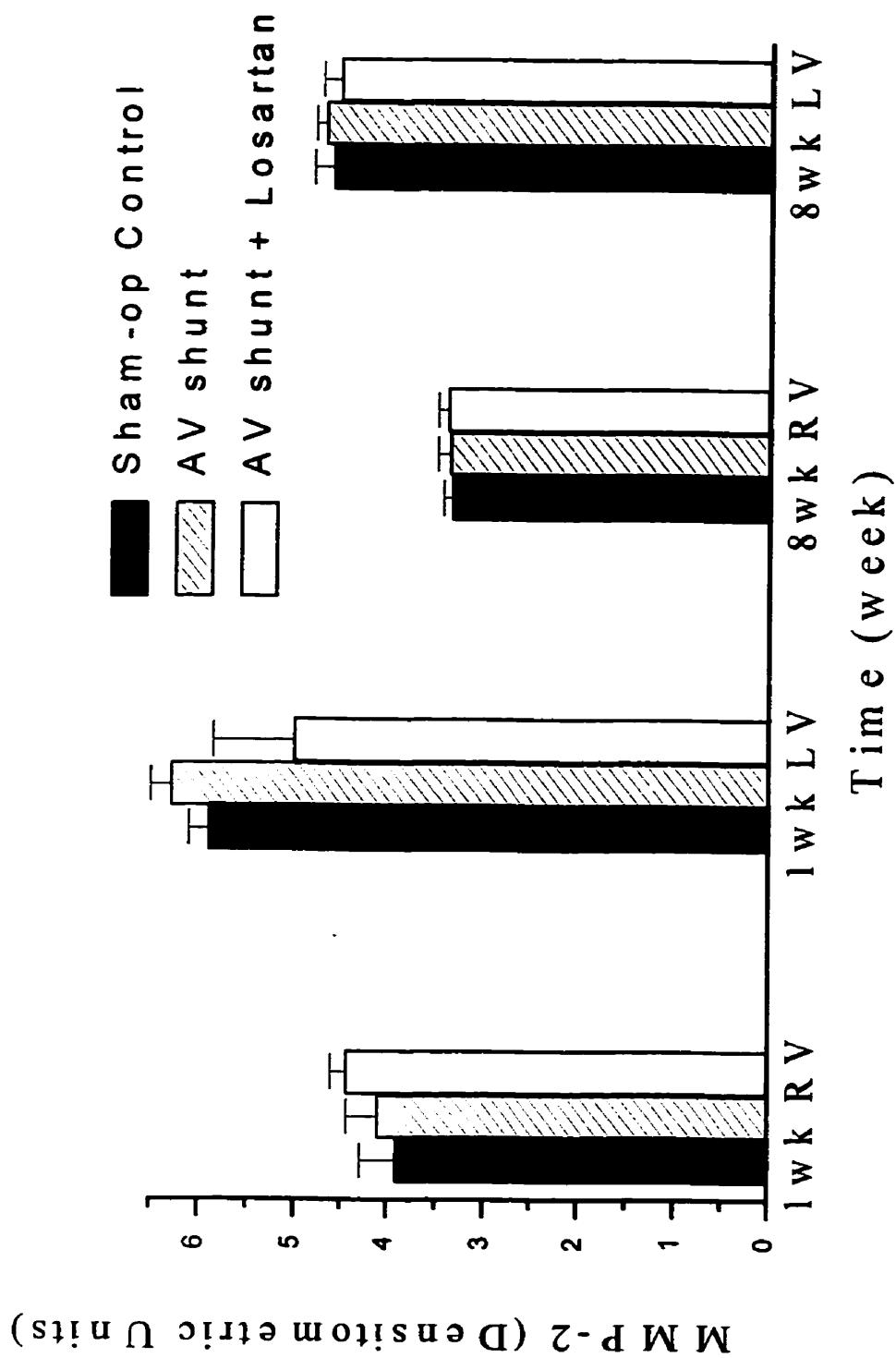


Figure 12. Western blot analysis for immunoreactive cardiac MMP-2 in left and right experimental animals (AV shunt), AV shunt animals treated with losartan (AV shunt + Losartan) and sham-operated controls, 1 and 8 weeks after AV shunt. The quantified data depicted is the mean \pm S.E.M. of 3-4 experiments.

*P<0.05 vs. control value. #P<0.05 vs. AV shunt value.

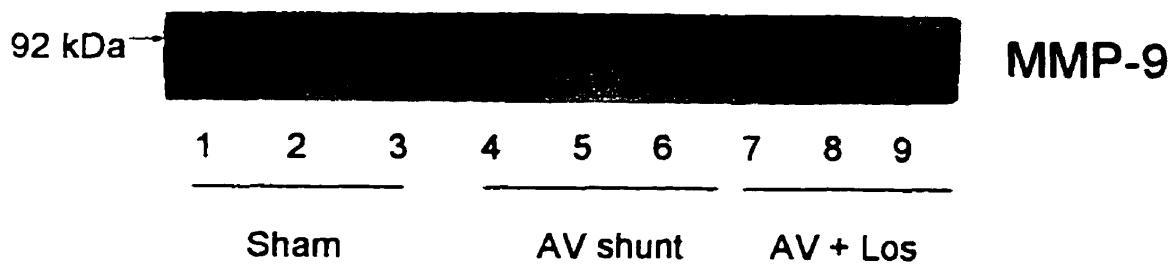


Figure 13. Representative Western autoradiograph showing bands for immunoreactive cardiac MMP-9 (92 kDa) in 8 week left ventricular tissue samples from experimental (AV shunt) animals, experimental animals treated with losartan (AV + Los) and respective age-matched sham-operated control (Sham) animals.

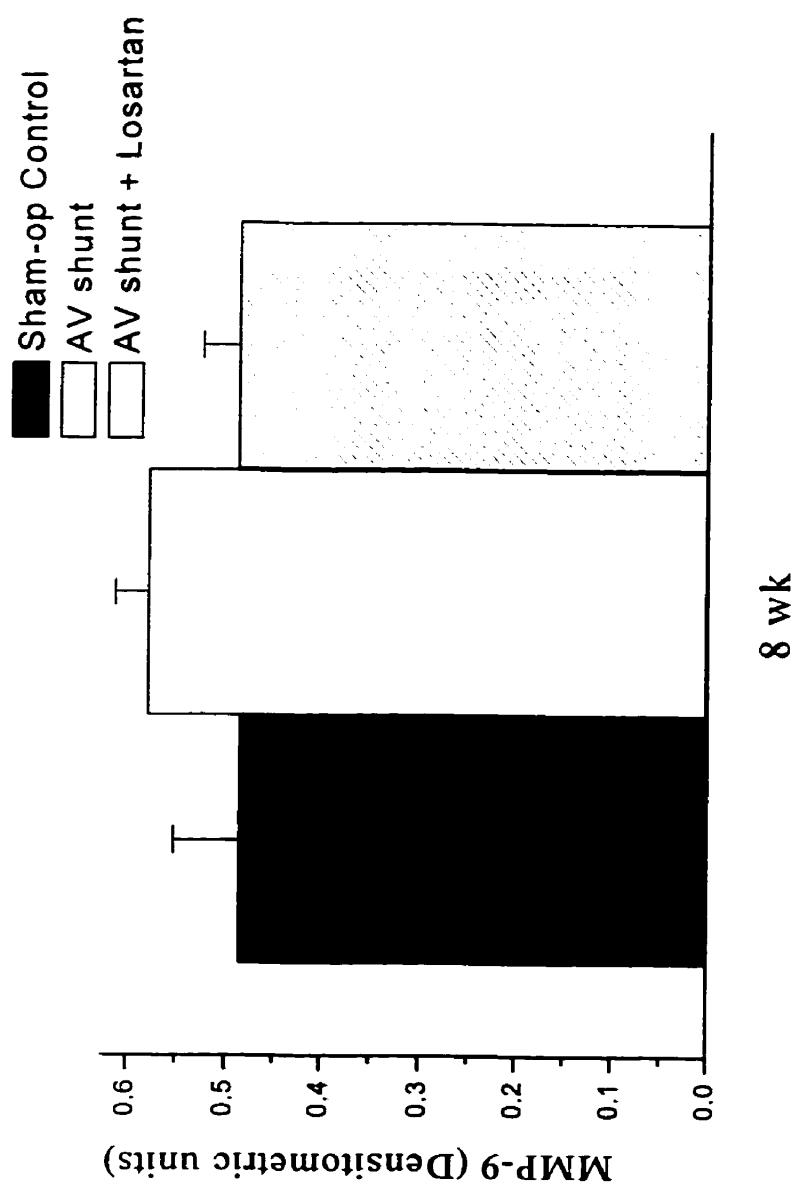


Figure 14. Western blot analysis for immunoreactive cardiac MMP-9 in left ventricles of experimental animals (AV shunt), AV shunt animals treated with losartan (AV shunt + Losartan) and sham-operated controls, 1 and 8 weeks after AV shunt. The quantified data depicted is the mean \pm S.E.M. of 3-4 experiments. * $P<0.05$ vs. control value. # $P<0.05$ vs. AV shunt value.

weight that corresponded to the specific MMP of interest.

1 week group

After 1 week of AV shunt, the immunoreactive signals for MMP-1, MMP-2 were unchanged from control levels. Densitometric analysis of the immunoblots was performed. The results obtained showed no significant differences between control, AV shunt and losartan treated groups (Figs. 9-12).

8 week group

At 8 week after the creation of AV shunt, the LV and RV myocardial MMP (MMP-1, MMP-2 and MMP-9) protein abundances showed no significant differences between the two groups. Losartan treatment had no significant effect on myocardial MMP (MMP-1, MMP-2, MMP-9) protein abundances at 8 week after shunt surgery (Figs 9-14).

6. Trichrome staining for total cardiac matrix

To localize and determine the specific distribution of secreted fibrillar collagens, Masson's trichrome staining was performed on heart sections taken from 8 wk AV shunt rats, 8 wk AV shunt group treated with losartan (40 mg/kg/day) and age matched sham operated controls. Representative cardiac sections from each group are shown in fig.15. Fibrillar collagens were stained blue. Collagen staining was greatly reduced in AV shunt operated group compared to sham controls. Losartan treatment was associated with modest increase in cardiac collagen as revealed by this technique. Furthermore, hypertrophy and dilation of the heart was clearly evident in cardiac sections from AV shunt operated animals as compared to those from sham operated controls. In sections from losartan treated AV shunt group, there was a marked decrease in hypertrophy and

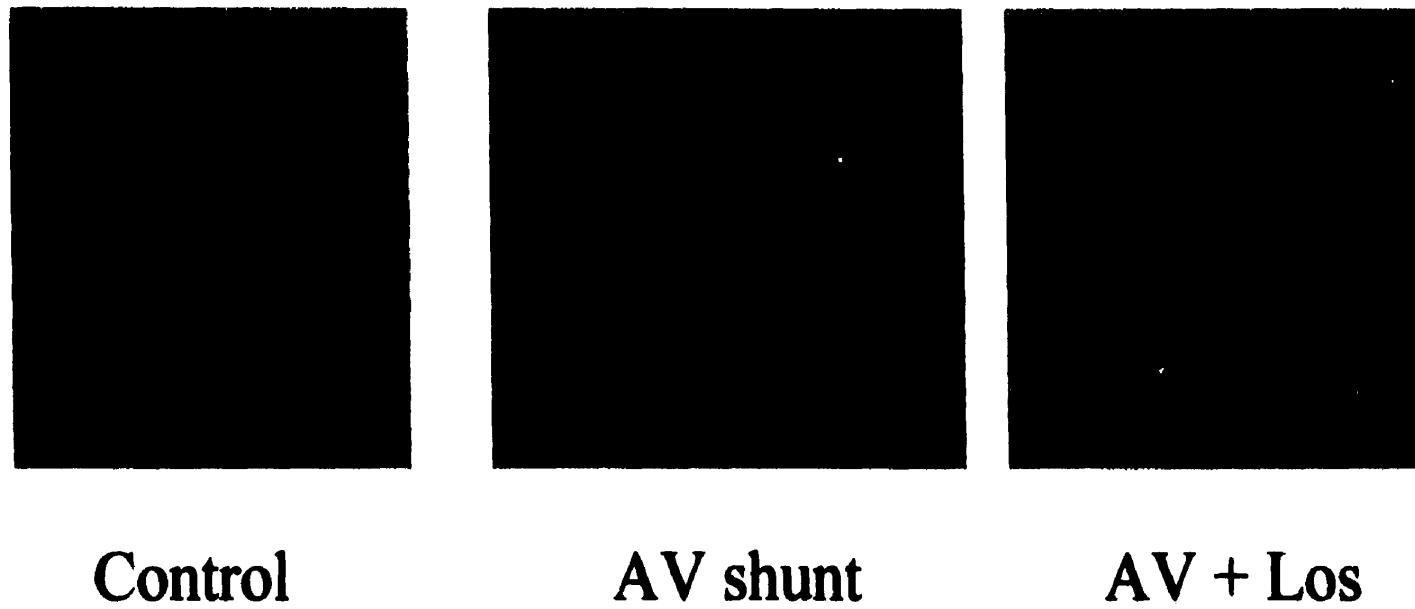


Figure 15. Trichrome staining for cardiac extracellular matrix in cross-sections of 8 week hearts from sham-operated control animals (no mixing of AV blood), experimental AV shunt animals or experimental animals treated with losartan for 8 weeks (AV + Los: 40 mg/kg/day). Fibrillar collagens in these tissue slices stain bright blue.

dilation (fig. 15) compared to AV shunt group.

7. Quantitative assessment of cardiac collagen concentration

To assess changes in collagen concentrations, 4-hydroxyproline assay was performed on tissue samples taken from right and left ventricles. Results are shown as representative histograms in figure 16 and 17.

1 week group

Cardiac collagen content, as determined by 4-hydroxyproline conc. showed significant decrease in AV shunt operated group as compared to sham operated controls in both right and left ventricles. Losartan treatment for 1 week was associated with a significant attenuation of this condition in left ventricular samples, while for right ventricle, losartan treatment had no significant effect.

8 week group

At 8 weeks after the creation of shunt, both LV and RV collagen content was significantly lower than that in controls. Losartan treatment for 8 weeks significantly improved this condition in LV while for RV, although the collagen content was significantly increased compared to AV shunt group, it still remained significantly lower than control group values.

8. Myocardial TIMP activity (reverse zymography)

TIMPs were identified using reverse zymography (by their ability to inhibit gelatinase activity) as more intensely stained bands at 29 kDa in the three groups studied. These bands corresponded well to the reported molecular weight of this inhibitor. The bands

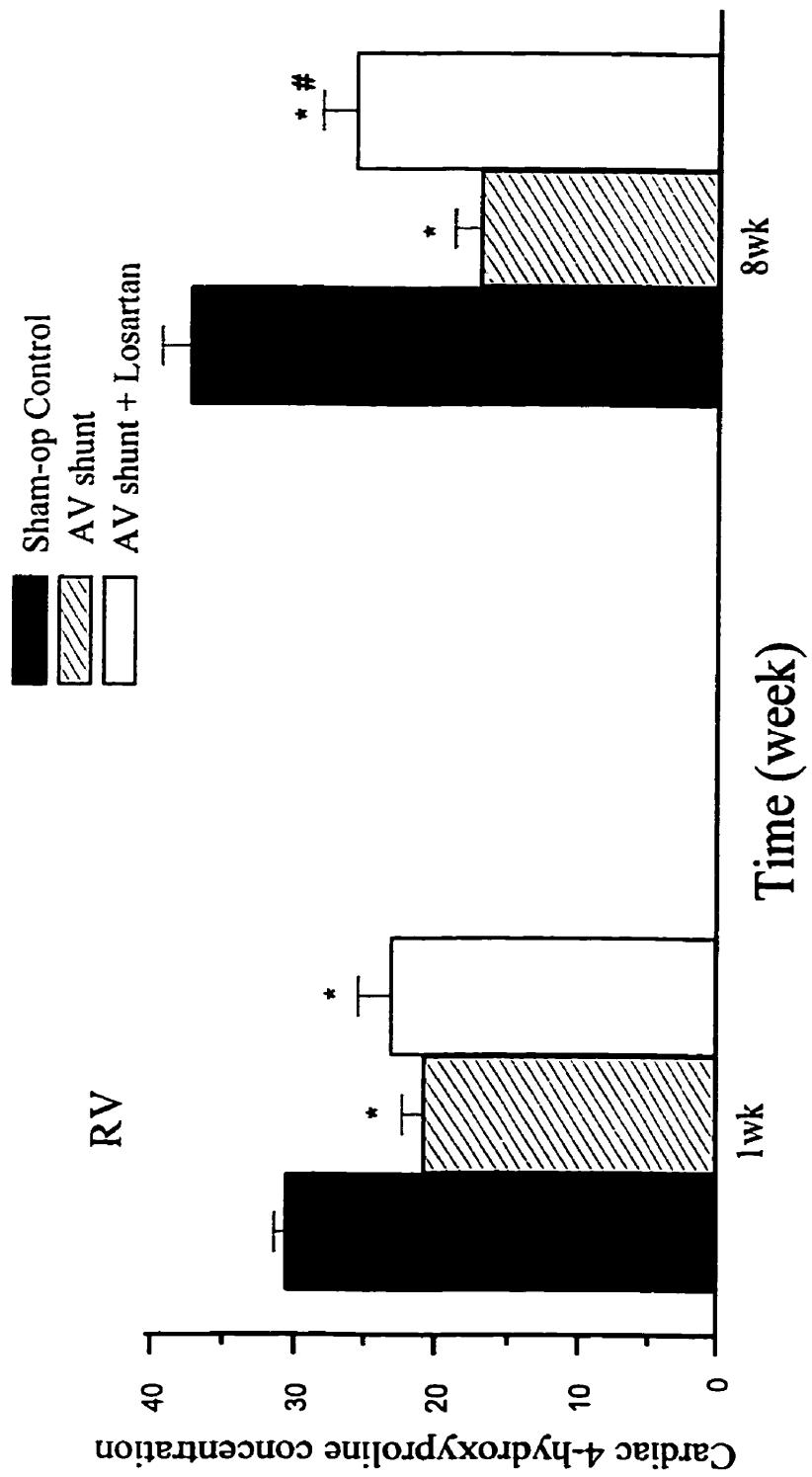


Figure 16. Myocardial 4-hydroxyproline concentration (relative collagen concentration) in right ventricular samples from 1 and 8 week experimental animals (AV shunt), AV shunt animals treated with losartan (AV shunt + Losartan), and age matched sham-operated controls. The data is expressed as mean \pm S.E.M from a total sample size of 4-6. *P<0.05 vs. Control values. #P<0.05 vs. AV shunt value.

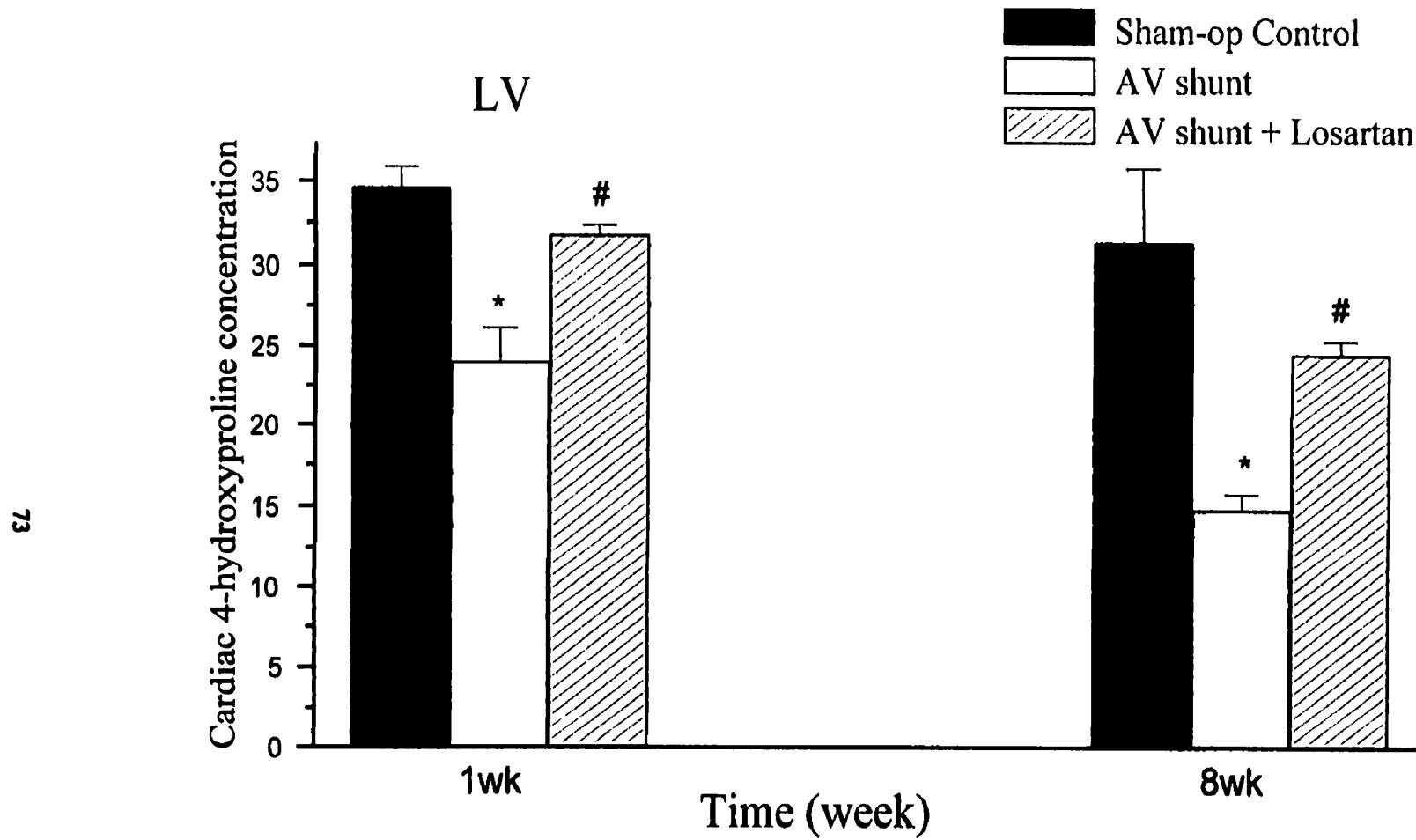


Figure 17. Myocardial 4-hydroxyproline concentration (relative collagen concentration) in left ventricular samples from 1 and 8 week experimental animals (AV shunt), AV shunt animals treated with losartan (AV + Losartan), and age matched sham-operated controls. The data is expressed as mean \pm S.E.M from a total sample size of 4-6. *P<0.05vs. Control values. #P<0.05 vs. AV shunt value.

were subjected to densitometric analysis in order to compare their relative inhibitory activities (Fig. 18).

1 week group

The RV and LV TIMP activity showed no significant changes at 1 week after the creation of AV shunt compared with control. Losartan treatment had no effect on myocardial TIMP activity (Fig. 18 and 19).

8 week group

The left and right ventricular TIMP activity was unchanged at 8 weeks post AV shunt compared to control. Losartan treatment was not associated with any significant change (Fig. 18 and 19).

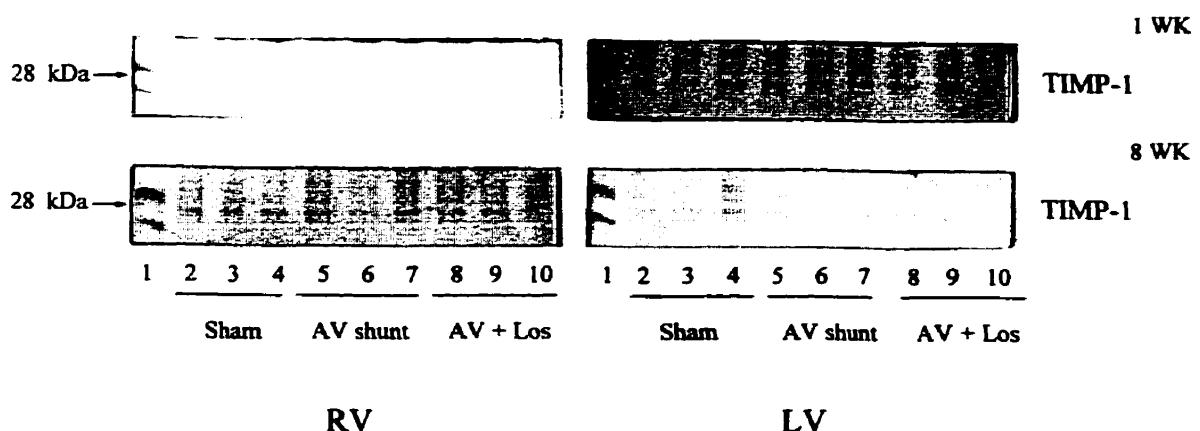


Figure 18. Representative gel showing reverse zymography for cardiac tissue inhibitor of matrix metalloproteinase-1 (TIMP-1) activity in 1 and 8 week ventricular (LV and RV) tissue samples taken AV shunt animals, AV shunt animals treated with losartan (AV + Los) and age matched sham-operated controls (sham). TMP-1 (28 kDa) band is indicated on the basis of calibrated molecular weight marker included in each zymogram (Lane 1).

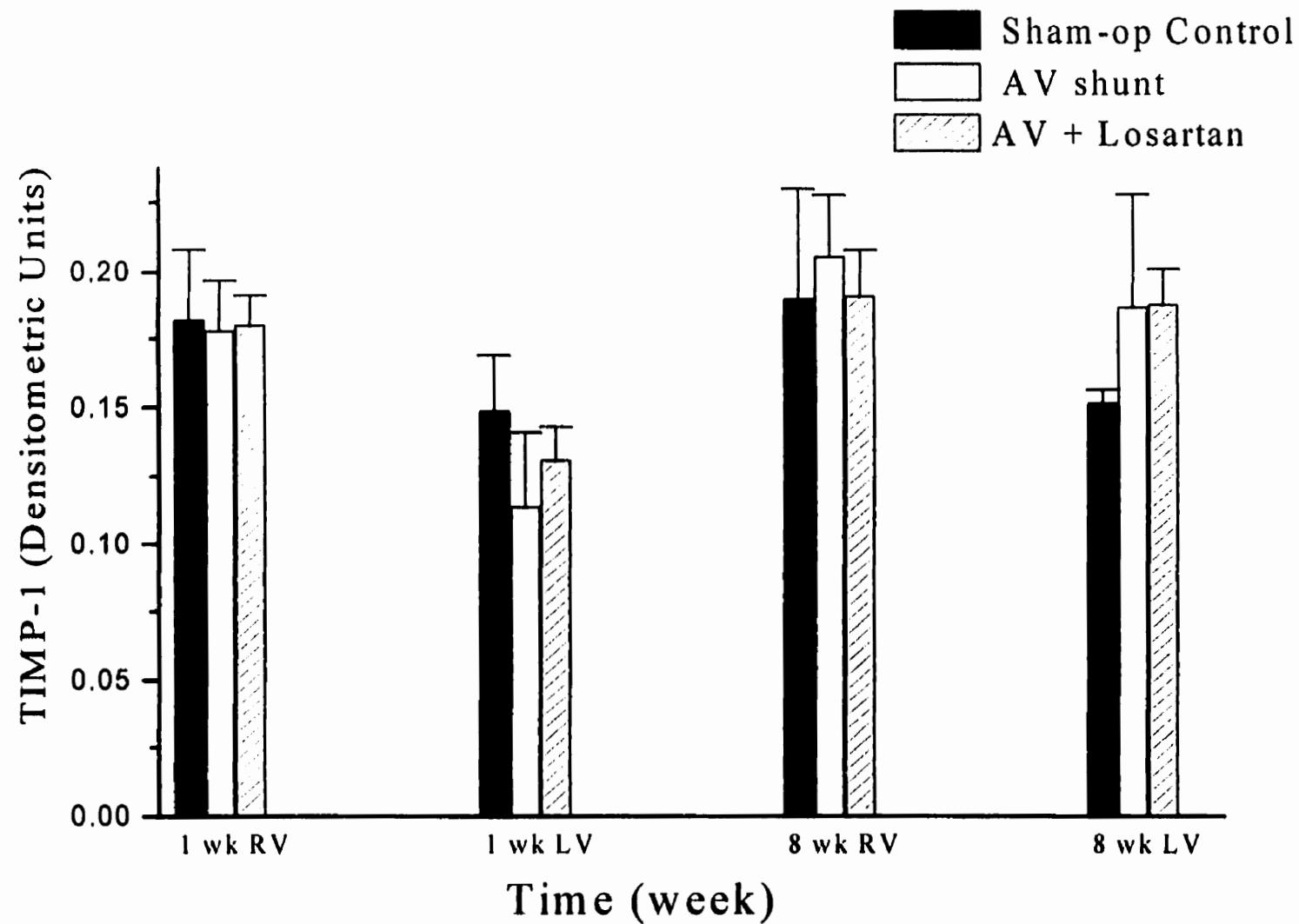


Figure 19. Estimation of relative activity of left and right ventricular tissue inhibitor of metalloproteinase-1 (TIMP-1) in 1 and 8 week myocardial samples taken from experimental animals (AV shunt) or AV shunt animals treated with losartan (AV + Losartan) compared to age matched sham-operated controls. The data depicted is the mean \pm S.E.M. of 4-6 experiments. * $P<0.05$ vs. control value. # $P<0.05$ vs. AV shunt value.

VII. DISCUSSION

Despite recent advances in understanding the mechanisms involved in the pathogenesis of heart failure, volume overload continues to be a substantial contributor to the morbidity and mortality of patients with this syndrome. Experimental models commonly employed to study volume overload have been mitral valve insufficiency²⁴⁶⁻²⁴⁸, aortic valve insufficiency^{249,250} and AV fistula.^{251,252,237,238,241} Each of these models is characterized by the production of volume overload and marked eccentric hypertrophy, but with notable differences in the hemodynamic variables. Mitral valve insufficiency results in a volume overload condition with low afterload due to retrograde flow of blood into the compliant left atrium. Aortic valve insufficiency results in a volume overloaded ventricle with high afterload due to the pressure gradient which causes retrograde flow into the LV. The AV shunt model produces a high output, bi-ventricular volume overload with normal systolic conditions, no retrograde flow, and a normal or slightly reduced afterload. Thus despite numerous studies that have been performed to understand the pathological abnormalities subsequent to volume overload, for comparative purposes, the data collected from each model require careful and cautious interpretation. We evaluated the role of MMPs in ventricular remodeling secondary to a sustained experimental volume overload using the rat model of AV shunt.⁴⁵ This model has the advantages of high reproducibility, low mortality and the ability to select the desired degree of volume overload. The results of this study confirms a positive correlation between increased end-diastolic loading of ventricles (volume overload) and myocardial remodeling, resulting in marked ventricular dilatation and hypertrophy. Our findings are: i) volume overload by an aortocaval shunt

was associated with decreased cardiac collagen content in both RV and LV at 1 and 8 week subsequent to shunt induction. ii) myocardial LV and RV MMP activity was significantly increased at 1 week after the opening of the shunt and remained elevated at 8 weeks after shunt creation iii) losartan administration to experimental animals was associated with attenuation of trends towards decreased collagen accumulation and increased myocardial MMP activity.

Changes in cardiac anatomy and cardiac peripheral hemodynamics by aorto-caval shunt

The AV shunt intervention used in this study provided a highly reproducible model of volume overload with a consistent degree of cardiac hypertrophy. Eccentric LV hypertrophy has been shown to develop in response to an increase in LVEDP and wall stress.²⁴⁶ In our model, LVEDP was significantly elevated at 8 weeks after the creation of AV shunt. Grossman *et al.*² found that end-diastolic wall stress was consistently increased in volume overloaded ventricles independent of the degree of compensation or stage of ventricular remodeling. They proposed that increased end-diastolic wall stress is the stimulus for the eccentric hypertrophy and ventricular dilatation associated with an increase in LVEDP secondary to bi-ventricular volume overload. The increases in body weight, LV and RV weight as well as lung weight seen in 8 week shunt rats were consistent with those heralding the development of congestive heart failure. Our results confirm that ventricular remodeling secondary to chronic volume overload in the AV fistula model is associated with the gradual progression to heart failure. We also determined that losartan treatment of experimental animals was associated with a

decrease in LVEDP. This is possibly caused by the prevention of sodium retention and volume expansion²⁵³, venodilation²⁵⁴ as well as modest decreases in afterload and modest increases in HR. On the other hand, LVSP and systolic and diastolic pressures all reduced in shunt rats, and were not affected with losartan treatment. These results are consistent with earlier studies showing that angiotensin only plays a role in the maintenance of blood pressure and total peripheral resistance shortly after induction of an aortocaval shunt.

Collagen protein concentration in volume overloaded hypertrophy

The present study demonstrated that interstitial collagen protein concentration as indicated by cardiac 4-hydroxyproline concentration was decreased in response to volume overload induced by AV shunt. This is consistent with an earlier report using a similar model in which Ruzika *et al.*²⁴¹ reported a decrease in left ventricular collagen concentration in response to volume overload. Similar results have been demonstrated in thyroxine induced cardiac hypertrophy where 8 weeks of daily thyroxine administration to juvenile rats resulted in the decrease in cardiac tissue collagen concentration.²⁵⁵ Weber *et al.*²²⁹ found a modest decrease in collagen volume fraction after 2 months in a dog model of AV shunt. Michel *et al.*²³⁸ also found a small decrease in LV collagen density in rats in a volume overload model of AV shunt. However in one other study, upregulation of steady state mRNA levels of collagens type I and III was noted even though collagen protein concentration remained unchanged.²⁵⁶ The inconsistencies among these results could be due to differences among the type and stage of heart failure associated with volume overload. Nevertheless, the majority of published data support the hypothesis that

decreased cardiac collagen concentration is a component of the pathology of ventricular dilatation. Since fibrillar collagens are recognized as structures that are essential for the maintenance of myocardial structural integrity, the decreased collagen accumulation may lead to dysfunction of the collagen weave with attendant loss of cardiac geometry and function. These changes may contribute, at least in part, to the progressive left ventricular dilation observed during volume overload.

Myocardial MMP and MMP Tissue inhibitory protein (TIMP) activity

Collagen removal in cardiac tissue is mediated through the activation of MMPs, which have high selectivity and affinity for protein components of the matrix.¹⁰⁶ One approach for determining relative MMP activity in tissue extracts is through the use of zymographic assays. In the present study, we found significantly increased MMP activity in both LV and RV tissue at 1 week after the induction of volume overload. Elevated MMP activity in experimental hearts was sustained through 8 weeks (the end point chosen for this study) and was accompanied by decreased cardiac collagen content in cardiac tissues. Several past reports have provided evidence to support the concept that increased MMP activity may contribute to the development of ventricular remodeling.¹⁷⁻

²¹ In one such report in which AV shunt model of volume overload was utilized, collagenase activity was found to be significantly elevated by 54% relative to control activity levels within 12 hours of volume overload induction which increased to ~86% above control activity at later time point (5 days to 8 weeks). These changes were paralleled by a decrease (~40%) in ventricular collagen concentration. Furthermore, reduced cardiac collagen concentration was accompanied by significant dilatation of

LV.²⁵⁷ It has been observed that similar ventricular remodeling preceded by increased collagenase activity occurs in myocardial infarction (MI). Within 3 hours post-MI, there was a 2-3 fold increase in collagenase activity and a 50% reduction in collagen content in the infarcted region.²⁶¹ In another investigation of stunned myocardium, rapid loss of ultrastructural collagen was observed to be preceded by a 74% increase in collagenase activity when compared to control values.²⁵⁹ Reddy *et al.*,²⁶⁰ found an eight-fold increase in collagenase activity in endomyocardial biopsy tissue obtained from patients with end-stage dilated CMP (DCM) compared with post-transplant patients with relatively normal hearts. In CMP hamsters, collagenase activity is progressively augmented with advancing age such that collagen degradation eventually exceeds synthesis resulting in an inadequate supportive interstitial collagen matrix.²⁶¹ That these changes occur at the age where significant ventricular dilatation and wall thinning occurs is indicative of a strong positive correlation between elevated collagenase activity and ventricular remodeling. Cleutjens *et al.*²⁶² have demonstrated the activation of collagenolytic system in the rat heart after coronary ligation. Tyagi *et al.*,²⁶³ reported increased collagenolytic activity in DCM and in infarcted ventricular tissues. Recently, in a pig model of pacing induced heart failure, Spinale *et al.*,²⁶⁴ reported a >25% decrease (vs control) in cardiac collagen content and an 80% increase in myocardial MMP activity after 7 days of supraventricular tachycardia (SVT). MMP activity increased to >100% after 14 days. These findings suggest that the increased activities of collagenases may thus be responsible for the degradation of collagen in the heart tissue. In the present study, we found persistent upregulation in MMP-1 activity in both LV and RV, while MMP-2 activity was found to be significantly increased only in 1 week LV. This result is in agreement with a report by

Armstrong *et al.*²⁶⁵ in which they suggested that the level of MMP-2 did not change significantly during any experimental condition. Furthermore, in the present study we observed significant increase in MMP-9 (vs control) in 1 week tissue (both LV and RV) samples. This again is consistent with the aforementioned observation by Armstrong *et al.*²⁶⁵ who reported significantly increased MMP-9 activity after 1 wk of pacing induced CHF. To determine whether increased zymographic activity during the progression of volume overload induced CHF was associated with changes in the relative steady state abundance of specific MMPs, we employed the western blotting technique. The results obtained showed no significant changes in MMP protein abundance at either 1 or 8 week after the induction of volume overload condition. This indicates that it is the increased activation, and not the increase in total cardiac steady state MMP protein, that might be responsible for the collagen loss seen in this study. TIMP-1 has been proposed to be an inhibitor of MMP-1 activity in heart. As we did not observe any change in cardiac TIMP-1 activity at any time point after the induction of volume overload, we suggest that TIMP does not play a major role in modulation of MMP activation in these experimental hearts. Although altered net collagen synthesis and extracellular degradation by other proteinases cannot be ruled out in contributing to reduced collagen concentration in volume overloaded hearts, the results from this study would favour the hypothesis that increased net MMP activity is responsible for enhanced collagen removal and net matrix degradation. The exact mechanism of an increase in MMP activity in this model is not clear. A number of cytokines and growth factors have been shown to either stimulate or inhibit collagen turnover including regulation of the synthesis of MMPs. The pro-inflammatory cytokines interleukin1 (IL-1) and tumor necrosis factor (TNF) stimulate

numerous cell types to secrete both collagenase and stromelysin.^{132,133} platelet derived growth factor (PDGF), epidermal growth factor (EGF), basic fibroblast growth factor (bFGF) and tumor necrosis factor-alpha (TNF- α) are all known to stimulate the synthesis and secretion of collagenase and stromelysin.¹³⁴⁻¹³⁶ Conversely, TGF- β_1 stimulation is associated with decreased steady state collagenase 3 (MMP-13) mRNA abundance in osteoblast cultures as well as decreased levels of immunoreactive procollagenase.¹³⁷ IL-4,¹³⁸ IFN γ (interferon),¹³⁹ and insulin like growth factors (IGF),¹⁴⁰ may downregulate MMP biosynthesis while heparin may activate MMPs.¹⁴¹ The circulatory levels of at least two inflammatory cytokines, TNF- α and IL-6, are elevated in patients with heart failure.^{266,267} Inflammatory cell proteinases (i.e neutrophil elastase, trypsin, plasmin) have been shown to activate the latent collagenase.²⁶⁸ Thus it is possible that altered signaling by one or more of these cytokines /growth factors leads to increased MMP activity in this model of volume overload.

Effect of AT₁ blockade on cardiac collagen concentration and MMP activity in volume overloaded hearts

Previous studies have suggested that the RAS is involved in the remodeling of the myocardial collagen matrix.^{241,269} Many of the cardiac RAS components including renin, angiotensin I, and angiotensin II expression have been shown to be elevated in volume overload due to AV shunt.³² Furthermore, losartan treatment has been shown to be effective in angiotensin II and attendant ventricular hypertrophy associated with volume overload.^{32,269,272} In the present study, we found decreased cardiac ventricular collagen concentration in the untreated animals at 1 and 8 weeks after the induction of volume overload. In both of these groups, losartan treatment was associated with restoration of

normal collagen concentration in ventricular samples. Ruzicka *et al.*²⁴¹ also found that 4-10 week losartan treatment was effective for attenuating decreased concentration of cardiac ventricular collagen in a rat model of volume overload and our results are therefore confirmatory. Together, these findings suggest that losartan treatment is effective in normalization of reduced collagen protein deposition in volume overloaded hearts. In our experiments dealing with MMP activation, we observed that 1 week losartan treatment was associated with significant attenuation of LV MMP-9 activity (zymography) while 1 week RV MMP-9 activity was not affected. Thus short-term losartan treatment was associated with a ventricle specific differential response regarding MMP-9 activity. On the other hand, 8 week treatment with losartan was associated with significant attenuation of MMP-1 activity in both ventricles. The precise mechanism for the normalized MMP-1 activity subsequent to AT₁ blockade in the experimental model of volume overload is unclear. The lack of a uniform response of activities of different MMP species to losartan suggests that a direct regulatory role by the AT₁ receptor is not solely responsible for these changes. In this regard, a number of cytokines and growth factors have been shown to induce or stimulate the synthesis of MMPs including IL-1, PDGF and TNF- α . Thus it is possible that the effect of AT₁ blockade on MMP activity is mediated via the secondary activation of one or more of these cytokines and/or growth factors after AT₁ receptor activation. Further studies to identify and assess the role of various cytokines and growth factors that may contribute to modulation of MMP activity seen in AV shunt model of volume overload needs to be carried out.

In conclusion, the result of the current study indicates that early activation of cardiac collagenase (and gelatinase) in experimental volume overload may be responsible for loss

of cardiac matrix collagen content during ventricular remodeling associated with volume overload. Losartan treatment was associated with normalization of MMP activity and collagen content in volume overloaded hearts.

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