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The Multiple Mechanistic Faces of a Pure Volume Overload: Implications for Therapy

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Abstract

Mitral regurgitation, and other conditions marked by a pure isolated volume overload (VO) of the heart, result in a progressive form of eccentric left ventricular (LV) remodeling and dysfunction. As opposed to the more extensively studied pressure overload, there are no approved medical therapies because an understanding of the underlying pathological mechanisms at work in VO is lacking. Over the past twenty years, our laboratory has identified multiple key biological functions involved in the pathological remodeling in VO. Specifically, we have noted perturbed matrix homeostasis, detrimental adrenergic signaling, increased intracellular reactive oxygen species, and an intense inflammatory response that implicates mast cells and their product chymase, which appears to cause extensive remodeling both inside and outside the cardiomyocyte. How these multiple pathways intersect over the course of VO and their response to various single and combined interventions are now the subject of intense investigation.

Keywords

Mitral Regurgitation; Volume Overload; Extracellular Matrix; Myocardial Inflammation; Adrenergic Drive; Oxidative Stress; LV dysfunction; LV remodeling

Introduction

Isolated mitral regurgitation (MR) is defined as MR due to intrinsic valvular disease. It is different from ischemic MR or functional MR, in which the LV dilatation occurs due to ischemic heart disease or cardiomyopathy that can cause the MR despite normal mitral valve leaflets. Mitral regurgitation creates a unique hemodynamic stress by inducing a low pressure form of volume overload (VO) caused by ejection into the low pressure left atrium. Mechanisms of LV dysfunction in isolated MR are not well understood, nor are there any approved medical therapies for this condition. Vasodilator therapy in other forms of LV dysfunction reduces LV wall stress and improves LV function; however, studies in isolated

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Work in this laboratory was the first to show that, as opposed to other forms of heart failure in which renin-angiotensin system (RAS) blockade has been highly successful, chronic VO is associated with loss of interstitial collagen surrounding cardiomyocytes¹ and a primary upregulation of the kallikrein kinin system.² This upregulation of the kallikrein kinin system promotes collagen loss, as well as inflammatory cell infiltration, and neither are improved by ACE inhibitor therapy.^{1,3} Studies of prolonged inflammatory cytokine or inflammatory cell blockade in human heart failure with pure volume overload have not met success. Mechanistic studies of a pure VO are further fueled by the difficulty in determining optimal timing for surgical intervention in valvular regurgitation because of the well defined decrease in LV ejection fraction, and the not uncommon incidence of heart failure postvalve repair/replacement. These problems may be improved by an appropriate medical therapy, which has yet to be determined, for the extended period of observation of chronic MR prior to surgery.

Extracellular Matrix in LV Remodeling in VO

To better understand why RAS blockade fails to improve remodeling and function in both the dog with experimentally induced MR, 1,4 and the rat with aorto-caval fistula (ACF) 3,5 , we performed gene arrays in both animal models. $^{6-8}$ In the dog with four months of MR, there is a significant downregulation of multiple noncollagen microfibillar and glycoprotein genes essential to collagen assembly and total extracellular matrix (ECM) structure,⁶ which matches the loss of collagen weave that connects individual cardiomyocytes. We also found marked upregulation of pro-fibrotic growth factors. In particular, there is a down-regulation of transforming growth factor-beta (TGF- β) mRNA and protein activity, but upregulation of matrix metalloproteinase (MMP) genes and the antifibrotic kallikrein kinin system (Figure 1). In the rat with 15 weeks of ACF, we find patchy areas of LV endocardial perivascular fibrosis, despite the loss of collagen surrounding cardiomyocytes (Figure 2).⁷ As in the dog, there is no increase in hydroxyproline at any time point, despite the increase in multiple mRNAs for various collagens at two weeks, thereby reinforcing the failure to increase ECM production despite an increase in LV angiotensin II (Ang II) levels in both animal models at early and late time points.^{1,2,3,9} Taken together, these findings suggest a compartmentalization of collagen homeostasis in the volume overloaded heart, with collagen loss surrounding cardiomyocytes persisting throughout the course of volume overload, and a perivascular fibrosis in later stages largely in the LV endocardium, which is subjected to higher wall stress and lower myocardial blood flow due to the elevation in LV diastolic filling pressures.

In addition to the lack of global extracellular matrix production in the pure VO, McDermot and coworkers report that chronic isolated MR in the dog does not trigger substantial increases in the rate of protein synthesis, but instead a decrease in the rate of protein degradation.¹⁰ This matches the modest (at most) increase in LV mass seen in this model and is consistent with studies from both the Carabello laboratory¹¹ and our laboratory^{12,13}, which demonstrate extensive myofibrillar degeneration/loss with the characteristic long and

thin cardiomyocytes. In preliminary studies by the late Dr. George Cooper years ago, he found a greater than two-fold decrease in total tubulin in the MR dog heart (unpublished data). Careful inspection of Western blots in Figure 3 demonstrates that, despite equal protein loading in the 15 week ACF rats, there is a decrease in tubulin expression. Thus, in the pure VO, the loss of extracellular interstitial collagen is matched by an intracellular loss of cytoskeletal and myofibrillar proteins. Current experiments in our laboratory are studying the connection of increased cardiomyocyte oxidative stress and MMP activation as the cause of myofibrillar and cytoskeletal breakdown. However, the lack of adequate replacement of collagen and myofibrils is a question that requires further investigation.

In part related to this "failure of hypertrophy" is the loss of cellular surface proteins that connect the collagen weave to the cell surface, resulting in the dephosphorylation of focal adhesion kinase (FAK) in both the dog with isolated MR¹⁴ and rat with ACF.¹⁵ An intact focal adhesion complex is essential for cell growth, survival, and myofibril assembly. As opposed to the volume overloaded MR heart, increased FAK phosphorylation has been documented in animal models of pressure overload.¹⁶ Interestingly, mice with selective inactivation of cardiomyocyte FAK demonstrate increased myofibrillar degeneration, elongation and thinning of cardiomyocytes, and eccentric LV remodeling and heart failure after six months.¹⁷ The disruption of FAK and cellular integrins are also associated with disruption, disorganization, and decreased respiratory capacity of subsarcolemmal mitochondria early in the course of volume overload in the rat with ACF, and this is prevented by MMP inhibition in the acute 24 hour ACF.¹⁸

Inflammatory Cell Infiltration in Volume Overload

The pure stretch of VO also demonstrates a molecular and cellular inflammatory response that contributes to MMP activation and ECM degradation. This inflammatory feature is consistent with an early and persistent increase in mast cells and chymase activity in the MR dog and ACF rat.^{1,2,3,5,9} As opposed to the constant increase in mast cells, there is a cyclical pattern of polymorphonuclear and macrophage infiltration only seen in the very early (acute) and late (decompensated) stages of ACF in the rat (Figure 3).⁷ Mast cells contain a collection of cytokines and proteolytic enzymes, including tryptase and chymase, which are released upon stress and activate MMPs. Mast cell stabilizing drugs have been reported to improve LV remodeling in the rat.¹⁹ However, in the dog with chronic MR, long term treatment with a mast cell stabilizer had no effect on LV remodeling and worsened cardiomyocyte shortening and calcium transients;²⁰ this was most likely due to the stabilizers' calcium entry blocking effect, which is essential in preventing mast cell degranulation.

As with studies in human heart failure, prolonged inflammatory cytokine or inflammatory cell blockade in pure volume overload have not met with success. Neutrophil infiltration blockade attenuates the MMP activation, ECM degradation, and myocyte apoptosis induced by ACF at 24 hours, as well as the development of eccentric hypertrophy induced by ACF at 2 and 3 weeks. In spite of this acute protective effect, however, sustained neutrophil depletion for more than 4 weeks results in adverse cardiac remodeling with further increase in cardiac dilatation and macrophage infiltration.²¹ In a similar fashion, we have found that

chronic TNF-a blockade failed to prevent eccentric LV remodeling or improve LV function after 2 or 4 weeks of ACF. It is of interest that the potent antioxidant protein, HMOX-1, was upregulated 5-fold at 2 weeks ACF and was significantly decreased by TNF- α blockade (Figure 4). Further, the increase in HMOX-1 is directly linked to TNF- α and not the hemodynamic stress of VO, because TNF-a neutralization decreased HMOX-1 transcript and protein expression but did not affect LV end-diastolic dimension (LVEDD) or pressure (LVEDP). Thus, TNF- α blockade actually abrogates the potentially beneficial effects of HMOX-1 and other cell-protective molecules in the course of volume overload. Further, it is now established that TNF receptor 1 exacerbates, and TNF receptor 2 ameliorates, LV remodeling, hypertrophy, NF-kB, inflammation, and apoptosis in heart failure, suggesting that future drug strategies should specifically target TNF receptor 1 effects.²² Finally, our gene array results from the volume overload of ACF in the rat demonstrate that the 24 hour,⁸ 2 week, and 15 week time points⁷ are marked by global inflammatory gene expression, while the 5 week interval was relatively quiescent for inflammatory gene expression. Thus, the expression of inflammatory cytokines in the MR heart may be dependent upon the clinical stage of LV remodeling and/or heart failure symptoms.

At this time it is not clear how the extracellular matrix loss in the pure volume overload can be therapeutically targeted, especially in view of the results in other models which demonstrated that prolonged MMP inhibition results in excessive collagen deposition (fibrosis), and that blockade of the bradykinin type 2 (BK₂) receptor results in a marked increase in blood pressure.²³ It is of interest that kallikrein, interstitial fluid bradykinin, and ACE2 activity are increased in rats with the VO of ACF at 4 and 15 weeks.² These in vivo anti-fibrotic responses in the rat with ACF were also recapitulated in isolated adult rat fibroblasts and cardiomyocytes after cyclical cell stretch, demonstrating increases in Kallikrein mRNA and bradykinin and MMP-2 activation.² Interestingly, blockade of kallikrein with the nonspecific protease inhibitor aprotinin prevented the decrease in collagen with chronic ACF and improved systolic function while having no effect on LV dilatation.²

Chymase has offered an interesting target for ECM degradation because mast cell infiltration and chymase activity are increased at all stages of VO in the rat and dog.^{1–3,5–9,23} In addition to an Ang II-forming capacity from Ang I that is 20-fold higher than ACE,^{24–26} chymase directly activates MMPs,^{27–30} degrades fibronectin,³¹ and activates kallikrein.³² The addition of chymase to isolated smooth muscle cells³³ or neonatal cardiomyocytes³⁴ results in cell death due to chymase degradation of cell adhesion proteins and disruption of the focal adhesion complex. A newly described propeptide cleaved from angiotensinogen, Ang-(1–12), may be an alternate substrate for the formation of biologically active angiotensins.^{35,36} Although the enzymatic mechanism responsible for cardiac Ang-(1–12) formation from angiotensinogen remains an open question, a recent study suggested a chymase-mediated mechanism in the ischemia/reperfusion rat heart.³⁷ In collaboration with Ferrario and coworkers we have reported that this chymase-mediated Ang II formation from Ang-(1–12) represents the predominant mechanism of Ang II formation in the left atria of humans undergoing corrective surgery for chronic atrial fibrillation³⁸, and also in the human LV.³⁹

It is important to note that the many unique features of isolated MR in the dog—MMP activation, increased bradykinin, ECM loss, disruption of the focal adhesion complex, and cardiomyocyte myofibrillar loss—are all either directly or indirectly beneficially affected by chronic inhibition of chymase, which leads to an improvement in LV systolic function and cardiomyocyte shortening.¹³

Increased Adrenergic Drive in VO

Another key factor in the pure VO is the role of increased adrenergic drive, which is central to both early and late stage volume overload of MR in both animal models⁴⁰⁻⁴² and humans.⁴³ Prolonged excessive adrenergic stimulation has a cytotoxic effect on cardiomyocytes,⁴⁴ in particular breakdown and loss of myofibrillar proteins in the cardiomvocyte of the dog and human with isolated MR.¹¹⁻¹³ Using a cardiac microdialysis technique in open-chest, anesthetized dogs in the early two and four week phase of the volume overload of isolated MR, we have shown that maximal electrical stimulation of the stellate ganglion activates intrathoracic adrenergic neurons to release norepinephrine (NE) and epinephrine (EP) into the cardiac interstitial fluid (ISF), and that β_1 receptor blockade $(\beta_1$ -RB) mitigates the excessive sympathetic neuronal EP and NE release.^{45,46} Neural ganglia cells in the heart express plentiful β receptors, and therefore a paracrine/autocrine feed-forward loop can be created where EP acts on the neuronal β_2 receptors and stimulates further release of NE and EP.^{40,41} This positive feedback can lead to abrupt and substantial increases in catecholamine exposure of cardiomyocytes, fibroblasts, and other cardiac cells. After synaptic release, NE and EP undergo metabolism and can be cleared by neurons (uptake 1); NE and EP can also be taken up by non-neuronal cells (uptake 2).⁴⁷ The remaining NE and EP escape these processes and enter the general circulation, and this is known as "spillover". Importantly, our studies in the dog indicate that measuring transcardiac plasma NE and EP concentrations substantially underestimates sympathetic efferent neuronal NE and EP release into the cardiac ISF space.⁴⁸ Thus, in addition to the direct effects on cardiomyocyte toxicity, β_1 -RB provides the additional benefit of reducing further catecholamine release form neural ganglia cells into the ISF during volume overload.

The combined loss of interstitial collagen with cardiomyocyte myofibrillar proteins and breakdown of the focal adhesion complex are improved by β_1 -RB.14,15 In the dog with isolated MR, β_1 -RB also results in significant improvements in cardiomyocyte function, calcium homeostasis, and LV function, but no improvement in extracellular matrix loss, LV dilatation, or cardiomyocyte elongation.^{11,49} In patients with severe MR and normal LV ejection fraction (EF), β -RB is associated with a survival benefit in patients, regardless of the presence or absence of coronary artery disease.⁵⁰ We have completed a phase IIb clinical trial of 38 asymptomatic subjects with moderate to severe isolated MR, who were randomized to either Placebo or β_1 -RB (Toprol-XL; average dose of drug: 75 mg q day) for two years.⁵¹ Magnetic resonance imaging with 3-dimensional analysis was performed at baseline and at approximate 6-month intervals. Rate of progression analysis was performed for primary end-point variables. Significant treatment effects were found on LVEF (p=0.0060) and LV early diastolic filling rate (p=0.0011), such that in untreated patients these systolic and diastolic parameters decreased over time on an intention to treat analysis. However, β_1 -AR blockade, as in the dog with isolated MR, does not attenuate the LV

dilatation. Thus, it would appear that chronic β_1 -RB protects the cardiomyocyte from excessive adrenergic drive; but the question remains, is valve surgery the only treatment that can arrest the continued LV dilatation.

Increased Cardiomyocyte Oxidative Stress in VO

A major step forward in identifying a potentially important target in an isolated volume overload was recently accomplished by examination of LV biopsies taken from patients with isolated MR. These patients had decreased LV systolic function 6 months post-mitral valve (MV) repair, despite LV ejection fractions (EF) > 60% prior to surgery.¹² These LVs had significant cardiomyocyte myofibrillar degeneration along with increased protein nitration and lipofuscin accumulation, consistent with an increased formation of reactive oxygen and nitrogen species, and activity of the protein xanthine oxidase (XO). In addition to being a major source of reactive oxygen species, XO is linked to bioenergetic dysfunction since its substrates derive from ATP catabolism. Correspondingly, there was also evidence of aggregates of small mitochondria in cardiomyocytes, which is generally considered a response to a cellular bioenergetic deficit.

To determine whether XO activation is linked to the hemodynamic stress of VO, we induced a 24 hour ACF in adult rats and isolated cardiomyocytes from sham and ACF left ventricles. Isolated cardiomyocytes from ACF LVs demonstrated increased XO activity, hydrogen peroxide and superoxide formation, increased MMP activity, and disruption of the subsarcolemmal mitochondrial network (Figure 5).^{18,51} The global MMP-inhibitor PD166793 preserved interstitial collagen, integrin- α 5 and the SSM structural arrangement, and prevented the decrease in state 3 mitochondrial respiration. These studies established a link between cardiomyocyte oxidative stress, MMP activation, and a disruption of the cytoskeletal network, which is a new emerging field for non-collagen targets of MMPs in cardiomyocyte ischemia/reperfusion injury.⁵²

To gain further insight into the role of oxidative stress in MR, we subsequently showed that cyclical stretch of adult rat cardiomyocytes caused increased oxidative stress, mitochondrial swelling, and cytoskeletal and myofibrillar disruption-all of which were prevented by XO inhibitor or the mitochondrial targeted antioxidant drug MitoQ (Figure 6).⁵¹ More importantly, MitoO prevented XO activation and myofibrillar degeneration, suggesting that the mitochondria themselves represent a more direct target in preventing oxidative stress, XO activation, and cellular remodeling in cardiomyocyte stretch. As mentioned previously, increased adrenergic drive is central to early and late stage MR. In the dog and human with isolated MR, β_1 -RB results in improvement in cardiomyocyte function, calcium homeostasis, and LV function, but no improvement in LV dilatation and cardiomyocyte elongation. However, β_1 -RB in the dog with isolated MR has no effect on a two-fold increase in LV XO activity (unpublished data), suggesting that additional protection against cardiomyocyte mitochondrial reactive oxygen species (ROS) production may be synergistic with β_1 -RB in the treatment of isolated MR. These studies have led us to hypothesize that the pure VO of isolated MR causes continual cellular damage emanating from pure stretchmediated mitochondrial ROS production. This, when combined with continual and mast cell infiltration and degranulation, sets up a self-perpetuating process of cytoskeletal and

myofibrillar breakdown without replacement, as described by McDermott and colleagues. Taken together, this suggests that a combined therapeutic approach of β_1 -RB and direct targeting of cardiomyocyte oxidative stress may be required to attenuate the adverse LV remodeling in VO.

Summary and Future Perspectives

The marked increase in oxidative stress in our in vitro studies of stretched cardiomyocytes that is prevented by MitoQ—has now directed our focus to prevention of ROS release form the numerous mitochondria in the cell (which occupy 40% of the cardiomyocyte interior volume). We are now testing the hypothesis that inhibition of mitochondrial ROS production will prevent: 1) MMP activation and digestion of cytoskeletal components, 2) the impressive loss of myofibrils, and 3) the disruption of interfibrillar mitochondrial structure and their connection to contractile units of the sarcomere. Figure 7 demonstrates the underlying myocardial events that are operating throughout the course of volume overload. Future directions will study the how the increased adrenergic drive, protease (chymase) and MMP activation, and mitochondrial-derived ROS production are interrelated and whether these are independent or interrelated processes in the pathophysiology of the unique hemodynamic stress of a pure stretch on the myocardium without a systolic pressure overload.

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Α	B Heat map of genes related to ECM structure			
	Gene name	Fold		
	Multimerin 1	-6.14		
	Vitronectin	-2.19		
	Decorin	-1.80		
	Fibrillin 1	-1.79		
	Fibulin 1	-1.78		
	Versican	-1.67		
	Integrin alpha V	-1.60		
	Lumican	-1.53		
	C Heat map of genes related to matrix degradation and ECM synthesis			
	Gene name		Fold	
	MMP1 MMP9		5.14	
			10.13	
	Plasminogen activator inhibitor type 1		-2.99	
	Thrombospondin 1		-2.41	
	Latent TGF-		-1.87	
and the second	TGF- β receptor 2		-1.68	
0.2 1.0 5.0	Connective tissue grow	vth factor	-1.56	
	TGF-receptor 3		-1.52	

B Heat map of genes related to ECM structure

Figure 1.

(A). Heatmap of the 659 genes altered greater than 1.5 fold (p<0.05) in the four week MR dog vs. baseline (1-4). Two-color gene array with dye swap was applied. Red = upregulation, black = no change, green = downregulation vs. baseline with scale of color corresponding to magnitude of fold-change. (B) Genes altered in MR related to ECM structure and (C) TGF- β pathway and ECM degradation. Previously reported in Zheng et al., Circulation 2009.

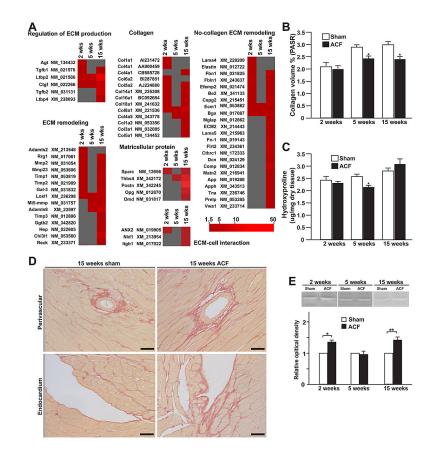


Figure 2. Expression of extracellular matrix protein in LV of sham and ACF rats

LV interstitial collagen measured by (**A**) PASR staining and (**B**) hydroxyproline content in 2, 5 and 15 week age-matched sham and ACF rats. (**C**) Representative images of perivascular and endocardial collagen staining by PASR. Excessive collagen accumulation is observed in perivascular LV areas in ACF rats. (**D**) Representative gel-zymography image demonstrating MMP-2 activity in 2, 5 and 15 weeks sham and ACF rats. Results are expressed as fold-changes to corresponding control. (**E**) Representative western blot with tubulin loading controls and (**F**) immunohistological staining of periostin in age-matched sham and ACF rats. Endo: endocardium; V: vessel. Bars: 20um. **P*<0.05, ***P*< 0.01 vs. sham. Previously reported in Chen et al., Am J Physiol, 2011.

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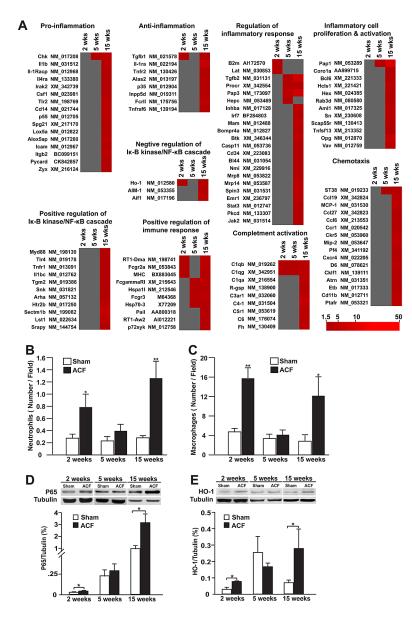
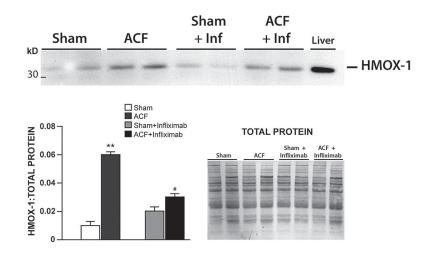


Figure 3. Heatmap demonstrating upregulated inflammation-related genes identified by DAVID analysis and inflammatory cell infiltration and protein expression of inflammation related genes (A) Upregulated genes related to inflammation and immune response at 2, 5 and 15 week. (B) Number of neutrophils (MPO+) and (C) macrophages (CD68+) in ACF and agematched shams at each time point. Representative western blots demonstrating expression of (D) NF- κ B p65, (E) HMOX-1 (heme oxygenase 1) with densitometric analysis after normalization to tubulin. **P* < 0.05, ***P* < 0.01 vs. sham. Previously reported in Chen et al., Am J Physiol, 2011.





HMOX-1 in sham, ACF, sham + infliximab (Inf) and ACF + Inf with total protein demonstrated by commassie stain. **P < 0.01 vs. sham, #P < 0.01 vs. ACF. Unpublished data.

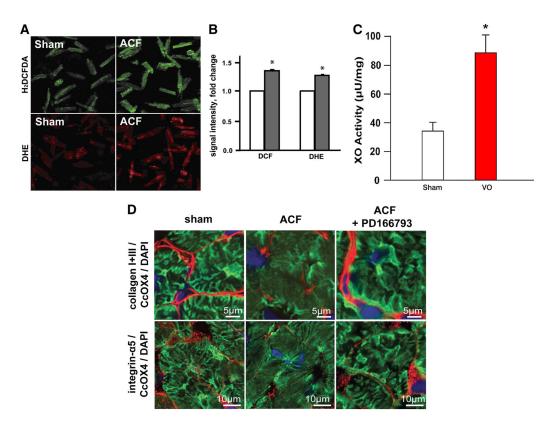


Figure 5. Oxidative stress in cardiomyocytes 24hr ACF

(A) Fluorescence microscopy of cardiomyocytes from sham and 24hr ACF rats probed with H₂DCFDA and DHE dyes for H₂O₂ and O₂⁻⁻ formation (respectively). (**B**) Quantitation of fluorescence intensity in cardiomyocytes from sham (white) and ACF (grey) rats. *p<0.05 vs. sham n=3 rats/group. (**C**) XO activity in sham and 24 hr ACF rat cardiomyocytes. (**D**) Overlay of collagen I and III, and Integrin α -5 with cytochrome oxidase 4 CcOX4 a marker of mitochondrial density and DAPI at higher power demonstrates a prominent loss of CcOX4 and Integrin α -5 in the subsarcolemmal area of cardiomyocytes in the ACF LV, which is restored by global MMP inhibitor PD 166793. Previously reported in Ulasova et al., J Mol Cell Cardiol, 2011.

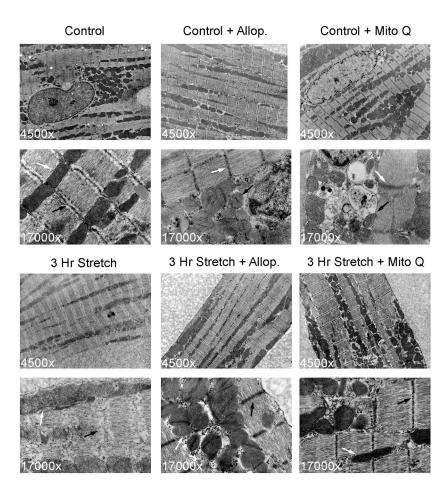


Figure 6. Cyclical stretch causes myofibrillar structural damage and mitochondrial swelling in rat cardiomyocytes

<u>Upper panels</u>: Transmission electron microscopic TEM) images of unstretched cardiomyocyte control, control + allopurinol (allo, 250 μ M), control + Mito-Q (50 nM); 4,500x and 17,000x. <u>Lower panels</u>: TEM images of cardiomyocytes after 3 hour stretch treated with 250 μ M allo or 50 nM Mito-Q. Stretch causes breakdown of myofilaments and Z-line (black arrows) and mitochondrial swelling (white arrows) in stretched cells vs. controls. Z-line structural integrity and mitochondrial morphology are preserved by MitoQ. Previously reported in Gladden et al., Free Rad Biol Med, 2011.

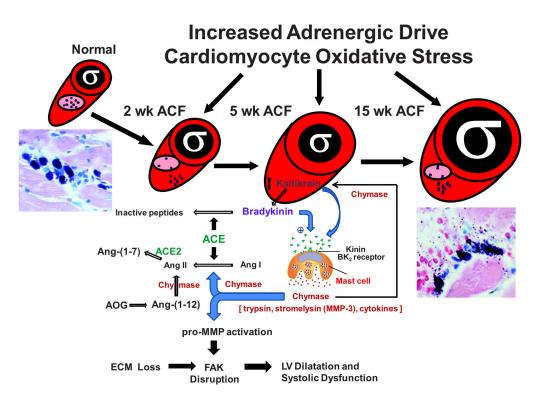


Figure 7. Mechanisms in the LV remodeling in a pure volume overload

VO has a number of simultaneous events with acute induction including increased adrenergic drive and cardiomyocyte oxidative stress. In addition, there is an acute influx of macrophages and neutrophils that abate while mast cell infiltration and degranulation continue throughout the course of VO. The release of chymase into the cardiac interstitium is associated with an increase in interstitial Ang II but also activation of MMPs and degradation of fibronectin, which combined with increased oxidative stress, leads to cardiomyocyte elongation and adverse LV eccentric remodeling. The failure to produce collagen despite an increase in Ang II may be counterbalanced by an increase in Kallikrein/ bradykinin and ACE2, which also result in the antifibrotic effects of Ang-(1–7). For all of these reasons, ACE inhibition increases BK₂–dependent chymase release from mast cells (60), which may further counteract a direct effect of ACE inhibition on Ang II formation by activating the chymase pathway of Ang II formation via Ang I and Ang-(1–12).