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## ***Myh11*<sup>R247C/R247C</sup> Mutations Increase Thoracic Aorta Vulnerability to Intramural Damage Despite a General Biomechanical Adaptivity**

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### **Abstract**

Genetic studies in patients reveal that mutations to genes that encode contractile proteins in medial smooth muscle cells can cause thoracic aortic aneurysms and dissections. Mouse models of such mutations, including *Acta2*<sup>-/-</sup> and *Myh11*<sup>R247C/R247C</sup>, surprisingly do not present with any severe vascular phenotype under normal conditions. This observation raises the question whether these mutations nevertheless render the thoracic aorta increasingly vulnerable to aneurysms or dissections in the presence of additional, epigenetic, factors such as hypertension, a known risk factor for thoracic aortic disease. Accordingly, we compared the structure and biaxial mechanical properties of the ascending and descending thoracic aorta from male wild-type and *Myh11*<sup>R247C/R247C</sup> mice under normotension and induced hypertension. On average, the mutant aortas exhibited near normal biomechanics under normotensive hemodynamics and near normal adaptations to hypertensive hemodynamics, yet the latter led to intramural delaminations or premature deaths in over 20 percent of these mice. Moreover, the delaminated vessels exhibited localized pools of mucoid material, similar to the common histopathologic characteristic observed in aortas from humans affected by thoracic aortic aneurysms and dissections. The present findings suggest, therefore, that mutations to smooth muscle cell contractile proteins may place the thoracic aorta at increased risk to epigenetic factors and that there is a need to focus on focal, not global, changes in aortic structure and properties, including the pooling of glycosaminoglycans / proteoglycans that may lead to thoracic aortic dissection.

### **Keywords**

familial thoracic aortic aneurysms and dissections; smooth muscle myosin heavy chain; actomyosin functionality; wall stress and stiffness

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Conflict of interest statement  
None to declare.

## 1. Introduction

Advances in medical genetics have revealed diverse mutations that predispose individuals to thoracic aortic aneurysms and dissections (Lindsay and Dietz, 2011; Milewicz et al., 2008). One sub-class of these mutations directly affects actomyosin contractile activity within the medial smooth muscle cells (Guo et al., 2007, 2013; Wang et al., 2010; Zhu et al., 2006). That is, causative mutations include those that encode either the contractile proteins (e.g., *ACTA2*, which encodes smooth muscle specific  $\alpha$ -actin, and *MYH11*, which encodes smooth muscle myosin heavy chain) or kinases involved in the actomyosin-mediated contraction and relaxation (e.g., *MLCK*, which encodes myosin light chain kinase, and *PKG-1*, which encodes a type 1 cGMP-dependent protein kinase). Notwithstanding accumulating clinical and histopathological data from patients harboring these mutations, the continuing lack of sufficient healthy and diseased tissue limits our ability to elucidate the underlying molecular and biomechanical mechanisms that lead to aneurysms or dissections. There is strong motivation, therefore, to exploit mouse models that can help increase our understanding of genetically induced aneurysms and dissections of the thoracic aorta.

Kuang et al. (2012) recently reported a knock-in mouse model (*Myh11*<sup>R247C/R247U</sup>) of a recurrent, rare variant in *Myh11* that results in a substitution of a cysteine for an arginine in the motor head region of the smooth muscle myosin heavy chain, which affects both ATP and actin-binding. The mutant aorta displays a decreased contractile response as expected, but there is no overall vascular phenotype under normal conditions. That is, the mice exhibit normal growth, aortic structure, systemic blood pressure, and life expectancy. Yet, smooth muscle cells explanted from the mutant aortas exhibit increased proliferation compared with wild type cells both *in vitro* and *in vivo* in a carotid artery ligation model (i.e., increased neointimal proliferation under no-flow conditions). The latter observation suggests that, despite an apparently near normal vascular development and maintenance under physiologic conditions, the elastic arteries in these mutants may nevertheless be more vulnerable to maladaptive responses to alterations in hemodynamic loading or vascular injury. Noting that hypertension is a risk factor for thoracic aortic aneurysms and dissections (Chan and Rabkin, 2014; Elefteriades, 2008), the goal of this work was to biomechanically phenotype the ascending and proximal descending thoracic aorta in *Myh11*<sup>R247C/R247C</sup> mice under both normotensive and induced hypertensive conditions and to assess possible increased structural vulnerability in hypertension.

## 2. Methods

### Mouse Models

All animal procedures conformed with NIH guidelines and were approved by the Institutional Animal Care and Use Committee of the University of Texas Health Science Center at Houston, and all transfer of tissue between Houston and New Haven was in compliance with a Material Transfer Agreement established between the two institutions. As described previously (Kuang et al., 2012), a germline *Myh11*<sup>R247C/R247C</sup> mutation was achieved in mice having a mixed C57BL/6J  $\times$  129SvEv background. Male mice, both wild-type (*Myh11*<sup>+/+</sup>, denoted herein as WT) and homozygous mutant (*Myh11*<sup>R247C/R247C</sup>, denoted herein as R247C) were challenged, beginning at 8 to 9 weeks of age, with a 18-

week long combined treatment with NG-nitro-L-arginine methyl ester hydrochloride (3 g/L of drinking water), or L-NAME, and a high salt diet (83 NaCl chow, Harlan TD.92012). L-NAME increases blood pressure by blocking endothelial-derived nitric oxide synthase and salt loading is a common risk factor for hypertension. For completeness, data were collected for both normotensive (i.e., untreated, denoted uWT or uR247C) and hypertensive (i.e., treated, denoted tWT or tR247C) mice, the former of which allowed the first detailed assessments of the effects of genotype alone on the biomechanical properties. Blood pressures were measured in all four groups of mice using a standard tail-cuff method.

### Mechanical testing

The ascending thoracic aorta (ATA) and proximal descending thoracic aorta (DTA) were obtained via a euthanize and harvest procedure (i.e., 350 mg/kg tribromoethanol IP and exsanguination upon removal of the aorta) and shipped overnight from Houston to New Haven in an iced Hank's balanced salt solution (HBSS, GIBCO by Life Science Technologies Corporation, Grand Island, NY). Consistent with prior reports (cf. Humphrey (2002)), preliminary experiments revealed that overnight storage does not affect the passive mechanical properties. Upon receipt, the vessels were cleaned of excessive perivascular tissue and mounted on glass cannulae for mechanical testing. In particular, the ATA was cannulated between the aortic root and brachiocephalic artery, while the left subclavian artery, left common carotid artery, and aortic arch (just distal to the left carotid) were ligated to allow pressurization. Similarly, the proximal DTA was cannulated approximately between the left subclavian artery and the third pair of intercostal arteries, with each included intercostal branch ligated to enable pressurization. The two extremities of the arteries were coupled securely to the glass cannulae using sutures. The mounted specimens were placed within a custom, computer-controlled biaxial device (Gleason et al., 2004), and tested in HBSS at  $\sim 37^{\circ} C$  using established protocols (Ferruzzi et al., 2013). Briefly, following mechanical reacclimation (i.e., pulsatile loading from 90 to 110 mmHg at the *in vivo* axial length for 15 minutes) and preconditioning (i.e., 4 cycles of pressurization from 10 to 140 mmHg near the *in vivo* axial length), the luminal pressure, outer diameter, axial force, and axial extension were measured on-line using standard transducers, a video microscope, and custom software during two classes of tests: cyclic pressure-diameter (*P-d*) tests at three different levels of axial extension (*in vivo* and  $\pm 53$  of *in vivo*) and cyclic axial force-length (*f-l*) tests at four different levels of luminal pressure (10, 60, 100, and 140 mmHg). To minimize possible testing-induced mechanical damage during these seven protocols, imposed transmural pressures did not exceed 140 mmHg and imposed axial loads did not exceed 5.5 grams, each consistent with prior studies of normotensive vessels (Ferruzzi et al., 2013). In this way, the associated data can also be compared directly, if desired, with those collected in similar studies of other mouse models (Ferruzzi et al., 2015; Wan et al., 2010). Immediately following biaxial testing, a  $\sim 2$  mm long ring was cut from the proximal region of the specimen and imaged with a dissecting microscope at  $20\times$  magnification to measure the unloaded (i.e., traction-free) thickness *H*.

### Quantification of Mechanical Properties

Pressure-diameter and axial force-length data provide insights into the *structural stiffness* of an artery, including via the clinical metric of distensibility

$$D = \frac{d_{i,sys} - d_{i,dias}}{d_{i,dias} (P_{sys} - P_{dias})} \quad (1)$$

where  $d_i$  is the current inner diameter and subscripts *sys* and *dias* refer to configurations corresponding to systolic and diastolic pressures, respectively. Nevertheless, a detailed understanding of arterial mechanics requires an analysis of the associated stress-stretch behaviors and multiple metrics that can be inferred from them (Humphrey, 2002). Mean values of the circumferential and axial stretches were calculated via

$$\lambda_{\vartheta} = \frac{1+h/2}{A+H/2}, \quad \lambda_z = \frac{l}{L} \quad (2)$$

where  $a$ ,  $h$ , and  $l$  are values of the inner radius, wall thickness, and axial length in any loaded configuration and  $A$ ,  $H$ , and  $L$  are similar values in an unloaded reference configuration. Note that the inner radii were calculated from the measured outer diameters,  $d_0$  and  $D_0$ , via  $a = d_0/2 - h$  and  $A = D_0/2 - H$  and the assumption of volume preserving (isochoric) motions during testing. Mean values of the biaxial (i.e., in-plane) components of the Cauchy stress were calculated via

$$\sigma_{\vartheta} = \frac{P^{exp} a}{h}, \quad \sigma_z = \frac{f^{exp}}{\pi h (2a+h)} \quad (3)$$

where,  $p_{exp}$  is the luminal pressure measured by the transducers and  $f^{exp} = f_T + \pi a^2 P^{exp}$  is the axial force acting on the closed-ended vessel in the device, with  $f_T$  being the axial force measured by the force transducer. Following established methods (Ferruzzi et al., 2013), we quantified the mechanical behavior using a microstructurally motivated constitutive equation that relates the biaxial stresses and stretches via a stored energy function. When evaluated using unloading data, this function represents the energy stored elastically in the artery upon pressurization that would be available to do work on the distending fluid when depressurized. Our constitutive equation contains 8 parameters that can be determined via nonlinear regression by minimizing differences between theoretically predicted and experimentally measured  $P$ - $d$  and  $f$ - $l$  data, the latter of which can also be used to estimate energy dissipation during loading. These best-fit model parameters, in turn, were used to compute the *material stiffness* and related metrics. Details can be found elsewhere (Ferruzzi et al., 2013), but the basic equations are summarized in the Appendix. Finally, note that it is often best to evaluate the stress, stiffness, and energy at *in vivo* values of pressure and axial stretch, the latter of which is estimated easily *in vitro* as the value of axial stretch,  $A_z$ , at which the transducer measured axial force does not change with changes in pressure. This value of stretch appears to be energetically favorable and has proven to be a good indicator of arterial adaptations to altered hemodynamic loads as well as genetic mutations (Humphrey et al., 2009).

### Statistical Analyses and Data Presentation

Values of select geometric, structural, and material metrics were organized into quartiles for each of the eight experimental groups (i.e., uWT, tWT, uR247C, and tR247C for both aortic

segments, ATA and DTA) and graphically represented in the form of boxplots to highlight the median (2<sup>nd</sup> quartile), spread (interquartile range between the 1<sup>st</sup> and the 3<sup>rd</sup> quartiles), and outliers (values below the 1<sup>st</sup> quartile or above the 3<sup>rd</sup> quartile by at least 1.5 the interquartile range). Non-parametric Kruskal-Wallis analyses were performed to test differences amongst groups for each metric; post-hoc pairwise comparisons were used to assess statistical significance ( $p < 0.05$ , with adjustment for multiple comparisons). Finally, data collected during the three *P-d* and four *f-l* protocols on each specimen were averaged using methods in Ferruzzi et al. (2013) to create ‘average datasets’ for each of the four treatment/genotype pairs (uWT, tWT, uR247C, tR247C) at each of the two anatomical locations (ATA, DTA). Specimens identified as outliers by the quartile analysis were not included in these averaged data sets.

## Histology

Arterial segments were fixed in formalin following mechanical testing and stored in 703 EtOH until all specimens could be embedded in paraffin, sectioned, and stained together. Standard Hematoxylin & Eosin (H&E), Verhoeff-Van Gieson (VVG), MassonaAZs Trichrome (MTC), and Movat pentachrome (Mov) stains were used to assess cell morphology and the organization of extracellular constituents such as the elastic laminae, collagen fibers, smooth muscle cells, and glycosaminoglycans (GAGs). An Olympus BX/51 microscope (Olympus Inc., Center Valley, PA), equipped with an Olympus DP70 digital camera, was used to acquire images of the stained aortic cross-sections using consistent setting. Fdisentisbility

## 3. Results

Table 1 lists physiological data (mean  $\pm$  SD) for all surviving mice for the four basic experimental groups: untreated wild-type (uWT), L-NAME and salt treated wild-type (tWT), untreated *Myh11*<sup>R247C/R247C</sup> (uR247C), and L-NAME and salt treated *Myh11*<sup>R247C/R247C</sup> (tR247C). The blood pressure of untreated R247C mice was not significantly different from that of the WT mice. Treatment with L-NAME and salt significantly increased the diastolic and systolic pressures in the WT (by 40 and 41%, respectively) and especially in the R247C (by 80 and 58%) mice. Percent increase in pulse pressure due to treatment was higher in WT (41%, statistically significant) than in R247C (26%) mice. Neither genotype nor hypertension affected body mass.

Table 2 lists morphological and mechanical data for all 45 aortic specimens that were tested, namely, ascending thoracic aorta (ATA) and proximal descending thoracic aorta (DTA) from the four experimental groups (uWT, tWT, uR247C, tR247C). On average, aortas from the untreated mutants had smaller outer diameters than did the untreated wildtype mice despite wall thickness being slightly greater in the mutants. Treatment with L-NAME and salt resulted in similar increases in outer diameter (8 to 16%) independent of genotype. As expected, treatment-induced hypertension also tended to increase wall thickness and decrease the in vivo (i.e., preferred) axial stretch. Interestingly, however, the treatment-induced increase in wall thickness was significant only for the WT mice, in which pulse pressure increased the most (cf. Table 1). Note that the computed mean wall stresses at mean

arterial pressure (MAP) were not significantly different as a function of genotype or blood pressure, with the only exception of the circumferential stress in the treated R247C mice. However, the large standard deviations in the treated mutants suggested a need to consider the possible existence of outliers.

Arrangement of the data into quartiles indeed identified one mouse as an outlier in the treated wild-type group (i.e., 1 DTA and 0 ATA, or 1 out of 10 vessels) and three mice as outliers in the treated mutant group (i.e., 2 DTA and, albeit less marked, 1 ATA, or 3 out of 15 vessels), each consistent with visual inspections of the specimens during testing and examination of the raw mechanical data (Table 2). All specimens from these four outlier mice were thus excluded from the averaging procedure that was used to generate 'representative' datasets for the eight experimental groups (ATA and DTA for uWT, tWT, uR247C, and tR247C). Figure 1 (left panels) shows the averaged pressure-diameter responses for each group. Note the lack of a significant difference due to the mutation and only modest increases in structural stiffness with treatment in both the WT and R247C vessels (i.e., leftward shift in the curves), consistent with the aforementioned trend towards increased wall thickness due to hypertension. Table 3 lists the associated best-fit constitutive parameters for these eight groups based on the averaged datasets, and Figure 1 (right panels) shows the associated averaged circumferential stress-stretch responses. These results reveal similar material behaviors, again independent of genotype and treatment. Both Table 2 (mean  $\pm$  SD for all data) and model predictions based on the averaged datasets (i.e., excluding outliers) suggested that the *Myh11*<sup>R247C/R247C</sup> mutation modestly lowered both the values of circumferential and axial stress at mean arterial pressure and the individual preferred axial stretch in the untreated mice. Interestingly, the results suggested further that, when excluding outliers, L-NAME and salt treatment of the mutant mice seemed to have a partial compensatory effect in the axial direction whereby stress (279 kPa in the ATA and 254 kPa in the DTA) was restored toward control levels. Conversely, the circumferential stress (335 kPa in ATA and 290 kPa in DTA) increased above control values in the treated mutant mice, which suggested some remodeling interplay between the two principal directions. Overall, however, these findings collectively suggested that the mutant ascending and proximal descending aorta managed, on average, to compensate well biomechanically, even when challenged with induced hypertension, and to achieve a near normal gross mechanical behavior, albeit at a different in vivo working point.

Although stress-stretch plots (Figure 1, right panels) and associated modeling (Table 3) provide considerable insight into the material behavior, the primary function of an elastic artery is to store energy when distended during systole and to use that energy to work on the blood during diastole to augment blood flow and smooth the pulse pressure. Plots of average energy storage as a function of biaxial stretch (i.e., circumferential distension and axial extension) thus provide useful functional information. For example, Figure 2 reveals that the ATA was more compliant and isotropic in its biaxial behavior than the DTA, independent of genotype and treatment, and that the average energy stored at the group specific mean arterial pressure and in vivo axial stretch (solid circle) was greater in the ATA than in the DTA in the untreated vessels, independent of genotype. These two observations are consistent with in vivo differences in loading of the ATA and DTA, with the ATA alone

experiencing both cyclic distension and extension. Figure 2 reveals further, however, that treatment-induced hypertension reduced energy storage in all cases, due in part to the decreased in vivo axial stretch (cf. Table 2). Finally, stored energy was less for untreated mutants compared with untreated wild-type for both the ATA and DTA, yet the hypertension, on average, restored the mutant stored energy contours closer to control. Overall, therefore, consistent with the initial report of no vascular phenotype due to the *Myh11* <sup>R247C/R247C</sup> mutation (Kuang et al., 2012), differences in biaxial mechanical behaviors between average wild-type and average mutant aortas were subtle, not dramatic, in both normotension and induced hypertension. In other words, with the exception of the few outliers, the thoracic aorta appeared both to develop and to adapt to increased blood pressures reasonably well despite the mutation.

Figure 3 supports this conclusion, but shows additional measures of stiffness and energy dissipation for all specimens in terms of boxplots to highlight the overall median and spread of these metrics as well as the associated outliers (circled cross). As it can be seen, most of the metrics followed similar patterns in the ATA and DTA. Circumferential (material) stiffness increased with hypertension, but only significantly so in the mutant, whereas axial stiffness was not different statistically. Consistent with this finding, plus that of increased wall thickness due to treatment (cf. Table 2), distensibility D (Equation 1) decreased dramatically with hypertension in all cases. Due to similarities between the wild-type and the ‘adapted mutants’ in both normotension and hypertension, however, this common clinical metric would likely not be revealing of disease or dissection potential.

Consistent with observed phenotypic diversity in patients harboring various mutations, we hypothesized that the mutant outliers identified by statistical analysis of the gross mechanical data may have actually manifested thoracic disease, likely driven in part by the epigenetic factor of induced hypertension. Figure 4 thus contrasts averaged pressure-diameter (left panels) and stress-stretch (right panels) results versus those for the 3 mutant outliers identified by analysis of stiffness. These outliers clearly represent maladaptations that did not compensate for the combined effects of mutation and treatment. In particular, outlier 3 exhibited a much stiffer behavior compared with the average, mainly in the DTA, whereas the other two outliers were globally more compliant, mostly in the circumferential direction. Most strikingly, however, Figure 5 shows gross images, histological sections, and energy storage from four representative R247C descending thoracic aortas: one untreated ‘average compensated’ DTA (first column), one treated ‘average adapted’ DTA (second column), and two treated outlier DTAs (third and fourth columns). Similar to prior tests on thoracic aorta from wild-types (Ferruzzi et al., 2013), the untreated mutant vessels tended to have a smooth, nearly translucent appearance during in vitro testing. Induced hypertension reduced this translucency, consistent with a thickening of the wall that appeared to include changes in both the media (e.g., increased inter-lamellar spacing) and adventitia. Whereas cross-sections from the average mutant appeared contiguous whether treated or not, note the marked intramural delaminations within the two outliers (white arrows). Note, too, the dramatic thickening of the wall in the two outliers, perhaps due to an attempted reparative response (i.e., these delaminations appeared to have occurred in vivo, not during testing, which would likely have elicited a healing response). Although there was some increase in

GAGs (blue in Movat stained sections) due to the induced hypertension (compare sections in the first and second columns), there was a much greater accumulation in the two outliers (black arrows), including a marked pooling of GAGs in outlier 2 in the outer media that appeared to correlate with the location of the delamination. Unfortunately, it was not possible to correlate the precise regions of delaminations and mucoid staining because these segments were used for different measurements. Finally, note the marked differences in energy storage in the vessels exhibiting a delamination, which revealed decreased ranges of distension and extension as well as energy storage, all indicators of a compromised mechanical functionality.

Figure 6 shows an unusual feature in the DTA of the second mutant outlier. On gross imaging, there was a significant perivascular accumulation of collagen near one of the intercostal branches. Movat and VVG-stained sections suggested that there was an underlying focal intramural delamination near the inlet to the intercostal artery as well as frank interlamellar delaminations at sites adjacent to this branch, again corresponding to an apparent pooling of GAGs in the outer medial lamellae (Figure 5, Movat-stained cut in the third column). It thus appeared that this marked increase in perivascular tissue may have been stimulated as a healing response by an increased density of cells (H&E staining), likely fibroblasts, given the nearby delamination.

#### 4. Discussion

Genetic mutations affecting either smooth muscle  $\alpha$ -actin (encoded by the gene *ACTA2*) or smooth muscle myosin heavy chain (encoded by the gene *MYH11*) are causative for human thoracic aortic aneurysms and dissections (Guo et al., 2007; Zhu et al., 2006). Histopathological characteristics in the media include disruption or loss of elastic fibers, disarray or loss of smooth muscle cells, and significant accumulations of mucoid material (i.e., glycosaminoglycans / proteoglycans). Many of these patients present with aortic dissection, with or without aneurysm, at ages ranging from teenagers to older adults. Zhu et al. (2006) reported further that the *MYH11* mutation correlated with a ~ 3-fold increased aortic structural stiffness and an associated ~ 1.7-fold increase in pulse wave velocity, not unlike that seen in Marfan syndrome patients (de Wit et al., 2013) who are similarly susceptible to thoracic aneurysms and dissections (El-Hamamsy and Yacoub, 2009; Lindsay and Dietz, 2011). It is now generally accepted, therefore, that the smooth muscle contractile apparatus is important in establishing, maintaining, or restoring the structural integrity of the thoracic aorta (Humphrey et al., 2014; Milewicz et al., 2008; Prakash et al., 2010; Renard et al., 2013).

It has been somewhat surprising, therefore, that mouse models of altered smooth muscle  $\alpha$ -actin (*Acta2*<sup>-/-</sup>, see Schildmeyer et al. 2000) and smooth muscle myosin heavy chain (*Myh11*<sup>R247C/R247C</sup>, see Kuang et al. 2012) do not exhibit a severe vascular phenotype under normal conditions. Yet, a full understanding of the biomechanical phenotype requires a detailed assessment of the biaxial mechanical properties under normal and hemodynamically stressed conditions, which has not been attempted heretofore for these two mouse models. Consistent with prior histological findings by Kuang et al. (2012), including lack of marked elastic fiber disruption, we found that, on average, the biaxial



mechanical properties of the ascending and proximal descending thoracic aorta from mutants exhibited modest, not dramatic, differences from those of age-matched wild-type controls. The mutant aortas tended to be slightly less stiff materially (Figure 3), but somewhat stiffer structurally as revealed by pressure-diameter data (Figure 1) and calculated values of distensibility (Figure 3, right panels). This increased structural stiffness was likely due in part to the slight, though not statistically significant, increase in wall thickness (Table 2), but it was not nearly as dramatic as that reported for humans (Zhu et al., 2006). The mutant vessels also tended to store less elastic energy upon pressurization and to dissipate more energy than the controls (Figures 2 and 3), but the latter was marked only in the ascending aorta. Diminished energy storage decreases overall aortic function, but does not necessarily imply a compromised structural integrity (Ferruzzi et al., 2015). We conclude, therefore, that the thoracic aorta in *Myh11*<sup>R247C/R247C</sup> mice compensated during development so as to exhibit sufficient mechanical functionality and structural integrity and not develop aneurysms or dissections under normotensive conditions. Indeed, it appears that many viable mutant mouse arteries similarly compensate under normal conditions to achieve sufficient biomechanical properties and functionality (Louis et al., 2007; Bersi et al., 2014; Ferruzzi et al., 2015; Sather et al., 2012).

A similar comparison of the biomechanical properties of wild-type and mutant aortas subjected to an 18-week period of induced hypertension revealed that, on average, the aortas from the surviving mutants were also able to adapt well. In particular, similar to wild-types, the two primary changes with hypertension were a further increase in wall thickness and decrease in the in vivo value of axial stretch, both of which helped to decrease the pressure-induced increases in intramural stress toward normal values. A similar adaptation to induced hypertension has been reported for an elastin haploinsufficient mouse model (Wagenseil et al., 2007). Yet, there was an increased vulnerability to intramural delaminations by the R247C aortas in hypertension (Figure 5): only 1 of 10 wild-type specimens (103), but 3 of 15 mutant specimens (203) showed signs of intramural delamination, that is, separation of layers within the wall without frank dissection. In each of these cases, however, the adventitia appeared robust, which likely prevented any progression to rupture (Li et al., 2014). To these numbers, we add that the overall mortality of the R247C mice with hypertension was also greater than WT (263 vs. 73 after 8 weeks of treatment), with deaths in the R247C ascribed to acute aortic dissections based on necropsy (unpublished data). It should be noted, therefore, that complementary studies in wild-type mice have not identified thoracic aortic failures in either the two-kidney, 1-clip (2K1C) model or the aortic arch banding model of hypertension (Chen et al., 2011; Kuang et al., 2013; Wagenseil et al., 2007), though chronic infusion of a high concentration of angiotensin-II can lead to dissection (Ju et al., 2013).

It is well known that GAGs accumulate within the aorta in both aging and hypertension (Sawabe, 2010; Wight, 2008), yet these accumulations tend to be diffuse, not focal, and not to result in aneurysms or dissections. In contrast, one of the most distinguishing histopathological features of thoracic aortic aneurysms and dissections is a focal accumulation, often pooling, of GAGs (Humphrey, 2013; Jain et al., 2011). We observed herein that the hypertensive aortas appearing mechanically as outliers (i.e., not adaptive)

tended both to have focal accumulations of GAGs within the media and evidence of intralamellar delaminations (cf. Figure 5), though not frank dissections. Although we cannot establish causality based on the present results (e.g., GAGs could accumulate in response to prior injury), a recent computational model supports the hypothesis that pooled GAGs can precede spontaneous intramural delaminations since they represent sites of geometric and material discontinuity as well as possibly increased osmotic (Donnan) swelling pressures (Roccabianca et al., 2014). This hypothesis thus merits further testing. Finally, although it is not clear what leads to a local production and pooling of GAGs in the case of diminished actomyosin activity, we suggest that compromised mechancosensing may play an important role (Humphrey et al., 2014). This hypothesis also merits testing.

In summary, it is widely recognized that the primary function of the thoracic aorta is mechanical and consequently that there is a pressing need for more analyses of the mechanics (de Wit et al., 2013; Elefteriades, 2008; Wu et al., 2013). The present results represent the first assessment of the biaxial mechanical properties of the thoracic aorta in any mouse model of altered smooth muscle functionality. Somewhat surprisingly, we found that, on average, the thoracic aorta *inMyh11*<sup>R247C/R247C</sup> mice appears both to develop under physiologic loading and to adapt under induced hypertensive loading so as to maintain sufficient mechanical functionality and structural integrity. Nevertheless, however, phenotypic diversity manifested as a 263 early mortality of the hypertensive mutants and evidence of intramural delaminations in 203 percent of the surviving hypertensive mutants. The latter associated with focal, not diffuse, accumulations of GAGs. Because of the apparent highly focal nature of these intramural defects, one must be careful not to over interpret any results that are necessarily averaged, including those from Western blots, PCR, or measures of structural stiffness such as distensibility (D). Rather, there is a need for increased spatially-specific immunohistochemical data and computational models to test further the hypothesis that highly focal defects may increase overall vulnerability to intramural delaminations, which in turn may lead to lethal dissections or ruptures.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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## Appendix 1

The structurally motivated ‘four-fiber family’ model for the stored energy  $W$  can be written as (Ferruzzi et al., 2013)

$$W(\mathbf{C}, \mathbf{M}^i) = \frac{c}{2} (I_C - 3) + \sum_{i=1}^4 \frac{c_1^i}{4c_2^i} \left\{ \exp \left[ c_2^i (IV_C^i - 1)^2 \right] - 1 \right\}, \quad (4)$$

where  $c$  and  $c_1^i$  ( $i=1, 2, 3, 4$ ) are material parameters having the dimension of stress,  $c_2^i$  ( $i=1, 2, 3, 4$ ) are dimensionless material parameters,  $C = \mathbf{F}^T \mathbf{F}$  is the left Cauchy-Green deformation tensor, with  $\mathbf{F} = \text{diag}(\lambda_r, \lambda_v, \lambda_z)$  the deformation gradient tensor, and  $\mathbf{M}^i = (0, \sin \alpha_0^i, \cos \alpha_0^i)$  is a unit vector defining the orientation of the  $i$ th family of fibers ( $i = 1, 2, 3, 4$ ) in the reference configuration, with  $\alpha_0^i$  the reference angle between the  $i$ th family of fibers and the axial direction. Note, too, that  $I_C = \text{tr}C$  and  $IV_C^i = \mathbf{M}^i \cdot \mathbf{C} \mathbf{M}^i$  ( $i=1, 2, 3, 4$ ) are coordinate-invariant measures of deformation that can be expressed in terms of the principal stretches as

$$I_C = \lambda_r^2 + \lambda_\theta^2 + \lambda_z^2, \quad IV_C^i = (\lambda_\theta \sin \alpha_0^i)^2 + (\lambda_z \cos \alpha_0^i)^2, \quad (5)$$

where  $IV_C^i$  has the physical interpretation of the square of the stretch in the direction of the  $i$ th family of fibers. Assuming incompressibility,  $\det \mathbf{F} = 1$ , reduces the number of independent principal stretches and allows calculation of the radial stretch as  $\lambda_r = 1/(\lambda_v \lambda_z)$ , where  $\lambda_r = h/H$ .

It can be shown (Humphrey, 2002) that the Cauchy stress tensor is determined by

$$\mathbf{t} = -p \mathbf{i} + \bar{\mathbf{t}} = -p \mathbf{i} + 2\mathbf{F} \frac{\partial W}{\partial \mathbf{C}} \mathbf{F}^T, \quad (6)$$

where  $p$  is a Lagrange multiplier that enforces incompressibility and  $\mathbf{i} = \delta_{ij} \hat{e}_i \otimes \hat{e}_j$  is the spatial second order identity tensor, with  $\delta_{ij}$  the Kronecker delta. Hence, the two theoretically determined components of Cauchy stress are

$$t_{\theta\theta} = c \left( \lambda_\theta^2 - \frac{1}{\lambda_\theta^2 \lambda_z^2} \right) + \sum_{i=1}^4 c_1^i \exp \left[ c_2^i (IV_C^i - 1)^2 \right] (\lambda_\theta \sin \alpha_0^i)^2, \quad (7)$$

$$t_{zz} = c \left( \lambda_z^2 - \frac{1}{\lambda_\theta^2 \lambda_z^2} \right) + \sum_{i=1}^4 c_1^i \exp \left[ c_2^i (IV_C^i - 1)^2 \right] (\lambda_z \cos \alpha_0^i)^2, \quad (8)$$

where the Lagrange multiplier  $p = c/(\lambda_v^2 \lambda_z^2)$  results from the assumption of plane stress, namely  $t_{rr} < t_{vv}, t_{zz}$ .

These equations for stress, in turn, were used to determine theoretical values of luminal pressure ( $P^{\text{th}}$ ) and axial force ( $f^{\text{th}}$ ) as a function of the 8 unknown model parameters. A non-linear least-square regression, based on the Levenberg-Marquardt algorithm, was then used to estimate the unknown parameters by minimization of the objective function  $e$  proposed by Wan et al. 2010

$$e = \sum_{i=1}^N \left[ \left( \frac{P^{exp} - P^{th}}{\bar{P}^{exp}} \right)_i^2 + \left( \frac{f^{exp} - f^{th}}{\bar{f}^{exp}} \right)_i^2 \right], \quad (9)$$

where  $N$  is the total number of data points, the superscripts *exp* and *th* denote experimentally measured and theoretically predicted, respectively, with the overbars denoting to average values.

Finally, circumferential and axial components of the linearized stiffness,  $C_{vvvv}$  and  $C_{zzzz}$ , were determined from the stored energy function, viz.

$$C_{ijkl} = \delta_{ik} \bar{t}_{lj} + \delta_{il} \bar{t}_{jk} + 4F_{iP} F_{jQ} F_{kR} F_{lS} \frac{\partial^2 W}{\partial C_{PQ} \partial C_{RS}}, \quad (10)$$

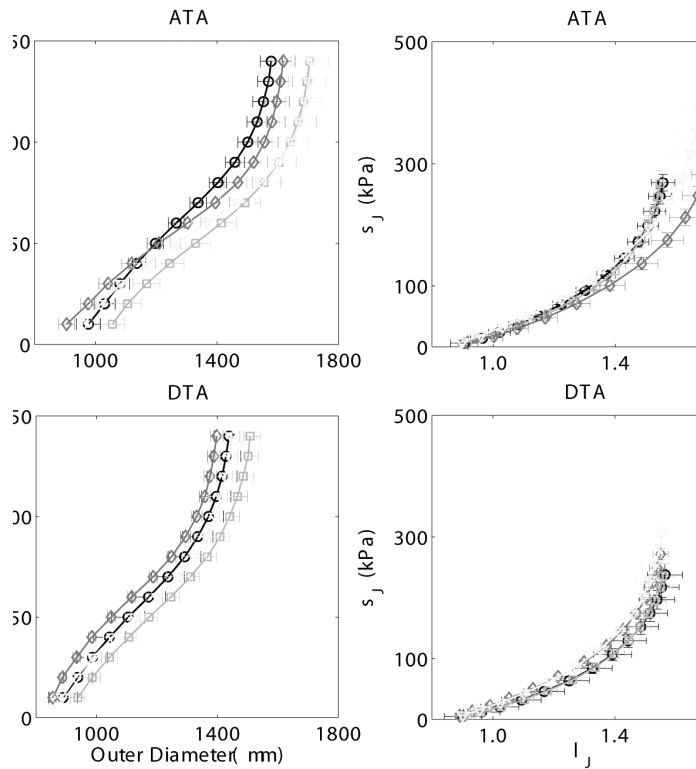
where lower case indices can be  $r, v,$  or  $z$  and upper case indices can be  $R, \Theta,$  or  $Z$ , and the components of the deformation gradient tensor ( $\mathbf{F}$ ) as well as of the deformation-dependent part of the Cauchy stress tensor ( $\bar{\mathbf{t}}$ ) are referred to the configuration about which the linearization is performed. Note that the values of linearized stiffness can be interpreted as the local slope of the corresponding stress-stretch curve about the configuration of interest provided that experiments are performed appropriately.

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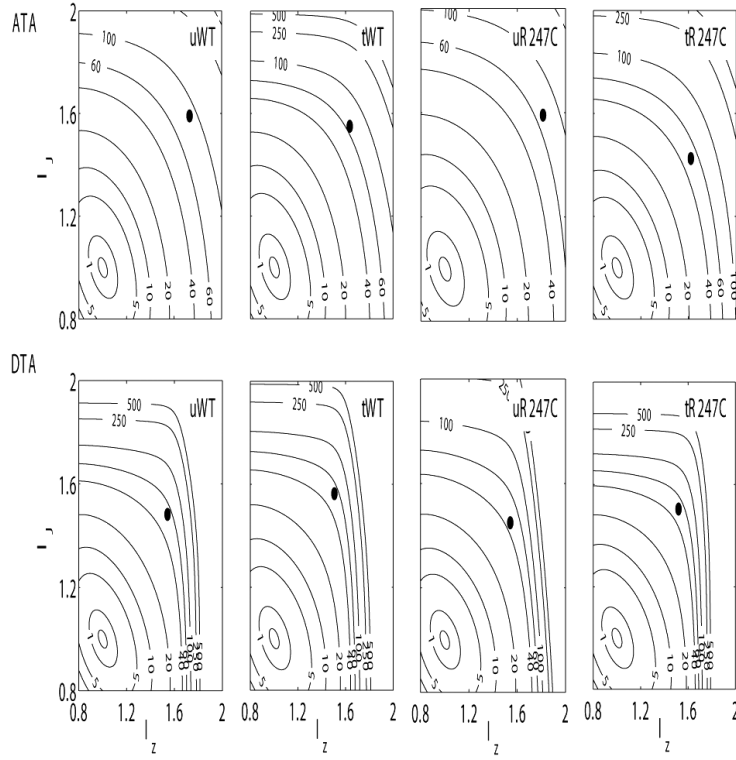
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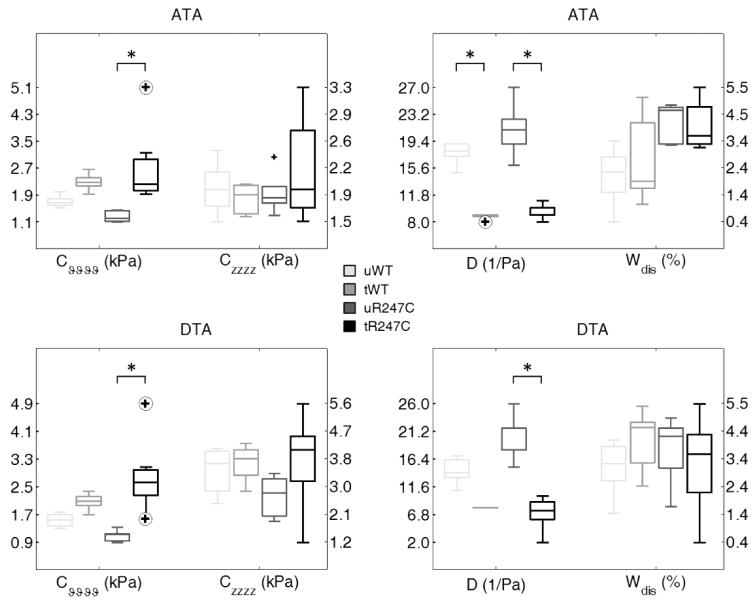
**Figure 1.**

Mean  $\pm$  SEM pressure - diameter responses (left panels) and circumferential Cauchy stress - stretch responses (right panels) for the ascending thoracic aorta (ATA, top panels) and the proximal descending thoracic aorta (DTA, bottom panels), excluding data from the 4 outliers. Datasets for the untreated (u) and treated (t, L-NAME + NaCl) wild-type (WT) as well as for untreated (u) and treated (t) *Myh11*<sup>R247C/R247C</sup> (R247C) mice reveal only modest differences due to either the mutation or the induced hypertension (i.e., L-NAME plus high salt treatment). Mean  $P$ - $d$  curves were obtained by averaging 5 ATAs/DTAs for the uWT, the uR247C, and the tR247C groups, and 4 ATAs/DTAs for the tWT group.

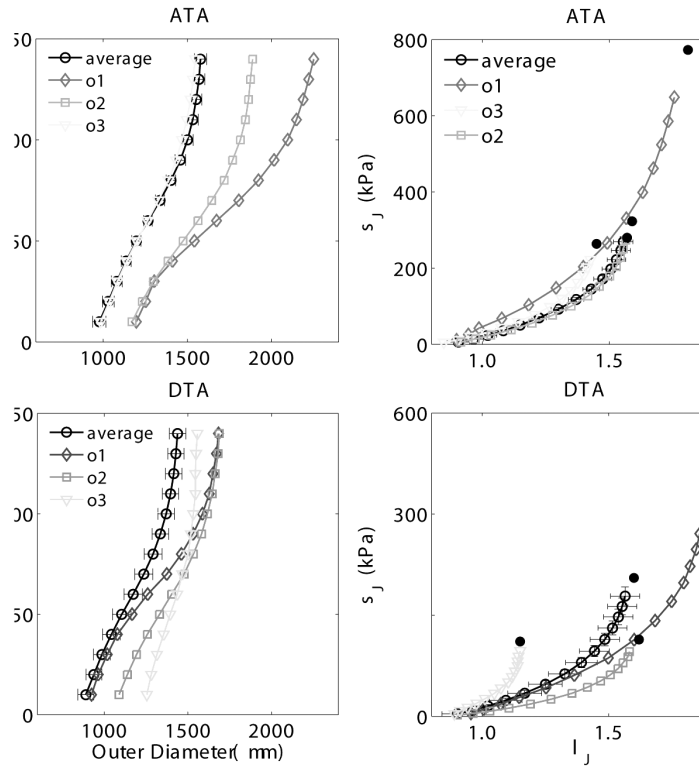


**Figure 2.** Contour plots of constant strain energy ( $W$ ) as a function of biaxial stretch  $\lambda_\theta - \lambda_z$  for the untreated wild-type (first column), treated wild-type (second column), untreated *Myh11*<sup>R247C/R247C</sup> (third column) and treated *Myh11*<sup>R247C/R247C</sup> (fourth column) mice. The strain energy functions were estimated from average datasets for the ascending thoracic aorta (ATA, top panels) and proximal descending thoracic aorta (DTA, bottom panels). The solid black circle in each plot indicates the in vivo conditions (mean arterial pressure and preferred axial stretch). The superimposed numerical values on the contours denote the values of strain energy in kPa. The mean strain energy functions ( $W$ ) representative for each combination of genotype, treatment, and location were obtained by averaging experimental data from 5 ATAs/DTAs for the uWT, the uR247C, and the tR247C groups, and 4 ATAs/DTAs for the tWT group.

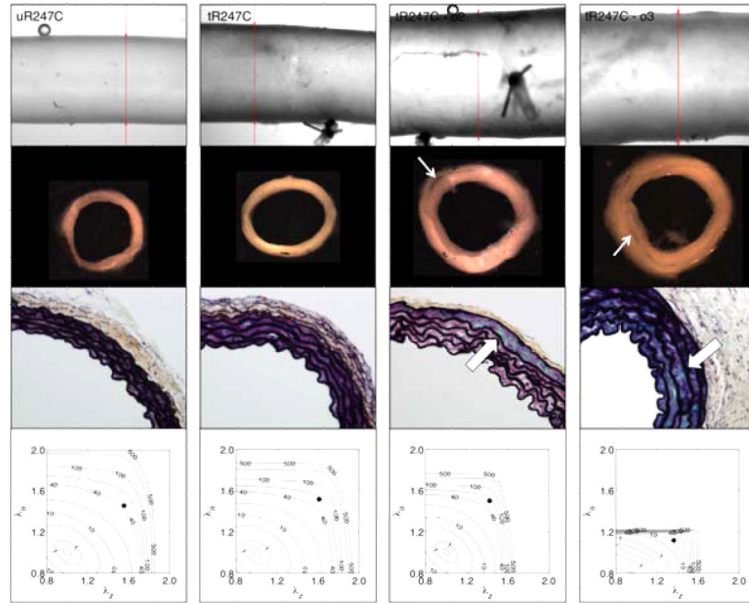




**Figure 3.** Mechanical metrics of material stiffness (left panels), and structural stiffness and energy dissipation (right panels) for the ascending thoracic aorta (ATA, top panels) and proximal descending thoracic aorta (DTA, bottom panels) of untreated (u) and treated (L-Name + NaCl, t) wild-type (WT) as well as for untreated (u) and treated (t) *Myh11<sup>R247C/R247C</sup>* (R247C) mice, showed at in vivo conditions (mean arterial pressure and preferred axial stretch).  $C_{\theta\theta\theta\theta}$  and  $C_{zzzz}$  are values of linearized stiffness in circumferential and axial direction, respectively.  $D$  is the distensibility computed at in vivo conditions (mean arterial pressure and preferred axial stretch).  $W_{dis}$  is the percentage of energy dissipated during the mechanical test, relative to the energy stored after loading. Note: circled cross symbols indicate outliers. Data from all collected vessels were included in the statistical analysis, i.e. 5 ATAs/DTAs for the uWT, tWT, and the uR247C groups, 7 ATAs and 8 DTAs for the tR247C group.

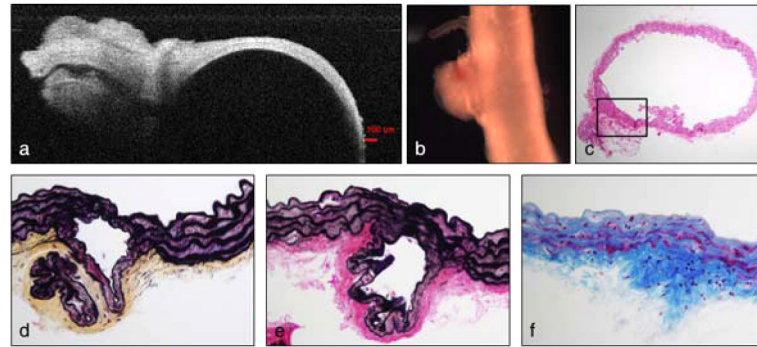


**Figure 4.** Pressure - diameter (left panels) and circumferential stress - stretch (right panels) data for the ascending thoracic aorta (ATA, top panels) and the proximal descending thoracic aorta (DTA, bottom panels). The datasets corresponding to the 3 treated *Myh11*<sup>R247C/R247C</sup> mutant outliers (o1, o2, o3) are compared with the average dataset for the treated mutant group in each plot, for both ATA and DTAs. The standard error of the mean is shown for selected points of the average dataset. Mean *P-d* curves for the tR247C group were obtained by averaging the data from the 5 ATAs and the 5 DTAs (5 treated mutant mice) that were not classified as outliers.



**Figure 5.**

Gross appearance, microstructure and mechanical behavior of descending thoracic aortas from one representative ‘average’ untreated *Myh11*<sup>R247C/R247C</sup> mouse (first column, uR247C), one representative ‘average’ treated *Myh11*<sup>R247C/R247C</sup> mouse (second column, tR247C), and two of the three treated *Myh11*<sup>R247C/R247C</sup> ‘outlier’ mice (third and fourth columns, tR247C-o2 and tR247C-o3 ). The panels in the first row show the gross appearance during biaxial testing, the panels in the second row show cross-sections of rings cut from the vessel after mechanical testing, the panels in the third row show Movat-stained histological sections (60x magnification), and the panels in the fourth row show the contour plots of constant strain energy ( $W$ ). The thin white arrows highlight intramural delaminations that were evident by eye. The thick white arrows highlight accumulated mucoid material, which tended to increase diffusely with induced hypertension but to accumulate more focally in the outliers having evidence of intramural delaminations. Note that the histological sections were necessarily taken from different regions than the isolated rings. Similar to Figure 3, the solid black circle in each contour plot indicates the specimen-specific *in vivo* condition (mean arterial pressure and preferred axial stretch). The superimposed numerical values on the contours denote the values of strain energy in kPa.



**Figure 6.**

Full vessel and stained cross-sectional views of the proximal descending thoracic aorta from the second outlier in the *Myh11*<sup>R247C/R247C</sup> treated group. A dense mesh of fibrotic tissue originated from the adventitial layer of the aorta (b, naked eye), in the proximity of one of the intercostal branches (a, Optical Coherence Tomographic image during mechanical testing). Cross-sectional cuts stained with H&E (c, 10x magnification), Movat (d, 40x magnification), VVG (e, 40x magnification), and MTC (f, 40x magnification) selectively show cells and/or ECM components and reveal a separation between the elastic laminae within the fibrotic region (magnified).

**Table 1**

Mean  $\pm$  SD physiologic data for the four groups: untreated wild-type (uWT), treated wild-type (tWT), untreated *Myh11<sup>R247C/R247C</sup>* (uR247C), and treated *Myh11<sup>R247C/R247C</sup>* (tR247C), where treatment combined L-NAME and high salt for 18 weeks beginning at 8 to 9 weeks of age.

	uWT	tWT	uR247C	tR247C
Number of Animals	5	5	5	8
Age (weeks)	27.3 $\pm$ 1.0	26.7 $\pm$ 1.0	26.0 $\pm$ 2.0	26.9 $\pm$ 1.4
Body Mass (g)	27.4 $\pm$ 4.3	27.2 $\pm$ 4.0	29.0 $\pm$ 2.6	23.6 $\pm$ 2.4
Blood Pressure (mmHg)				
Diastolic	91 $\pm$ 6	127 $\pm$ 8*	82 $\pm$ 2	141 $\pm$ 18*
Systolic	124 $\pm$ 7	175 $\pm$ 12*	117 $\pm$ 3	185 $\pm$ 16*

\* p < 0.05 between treated and untreated mice with the same genotype.

**Table 2**

Mean  $\pm$  SD morphologic and mechanical data for the ascending thoracic aorta (ATA) and the proximal descending thoracic aorta (DTA) for the four groups: untreated wild-type (uWT), treated wild-type (tWT), untreated *Myh11*<sup>R247C/R247C</sup> (uR247C), and treated *Myh11*<sup>R247C/R247C</sup> (tR247C). Note that one ATA was too short to test reliably. MAP denotes the group-specific mean arterial pressure.

<b>Ascending Thoracic Aorta (ATA)</b>				
	uWT	tWT	uR247C	tR247C
Number of Specimens ( <i>n</i> )	5	5	5	7
Unloaded Dimensions				
Outer Diameter ( $\mu\text{m}$ )	1099 $\pm$ 106	1254 $\pm$ 128	1025 $\pm$ 58	1154 $\pm$ 121
Wall Thickness ( $\mu\text{m}$ )	104 $\pm$ 11	142 $\pm$ 13*	117 $\pm$ 9	124 $\pm$ 19
In-vitro Axial Length (mm)	2.50 $\pm$ 0.29	2.96 $\pm$ 0.44	2.99 $\pm$ 0.57	2.56 $\pm$ 0.44
Dimensions at MAP				
Outer Diameter ( $\mu\text{m}$ )	1619 $\pm$ 132	1783 $\pm$ 190	1486 $\pm$ 110	1712 $\pm$ 275
Wall Thickness ( $\mu\text{m}$ )	38 $\pm$ 5	56 $\pm$ 6*	41 $\pm$ 5	48 $\pm$ 11
In-vitro Axial Stretch (–)	1.73 $\pm$ 0.05	1.63 $\pm$ 0.09	1.81 $\pm$ 0.15	1.64 $\pm$ 0.09
Cauchy Stresses (kPa) at MAP				
Circumferential	276 $\pm$ 38	278 $\pm$ 26	212 $\pm$ 21	387 $\pm$ 182*
Axial	313 $\pm$ 40	243 $\pm$ 31	272 $\pm$ 31	312 $\pm$ 112
<b>Descending Thoracic Aorta (DTA)</b>				
	uWT	tWT	uR247C	tR247C
Number of Specimens ( <i>n</i> )	5	5	5	8
Unloaded Dimensions				
Outer Diameter ( $\mu\text{m}$ )	1001 $\pm$ 62	1089 $\pm$ 76	978 $\pm$ 19	1088 $\pm$ 182
Wall Thickness ( $\mu\text{m}$ )	98 $\pm$ 8	142 $\pm$ 31*	106 $\pm$ 9	141 $\pm$ 42
In-vitro Axial Length (mm)	5.31 $\pm$ 0.78	7.49 $\pm$ 0.64	5.01 $\pm$ 1.38	5.64 $\pm$ 1.19
Dimensions at MAP				
Outer Diameter ( $\mu\text{m}$ )	1382 $\pm$ 93	1561 $\pm$ 125	1336 $\pm$ 85	1547 $\pm$ 141*
Wall Thickness ( $\mu\text{m}$ )	43 $\pm$ 5	60 $\pm$ 13*	46 $\pm$ 3	62 $\pm$ 25
In-vitro Axial Stretch (–)	1.54 $\pm$ 0.09	1.49 $\pm$ 0.05	1.54 $\pm$ 0.07	1.50 $\pm$ 0.09
Cauchy Stresses (kPa) at MAP				
Circumferential	205 $\pm$ 16	229 $\pm$ 30	164 $\pm$ 12	274 $\pm$ 101
Axial	225 $\pm$ 36	214 $\pm$ 30	190 $\pm$ 23	223 $\pm$ 75

\*  $p < 0.05$  between treated and untreated mice with the same genotype.

**Table 3**

Material parameters estimated by fitting the microstructurally motivated constitutive relation (Appendix 1) to average datasets (excluding outliers) for the ascending thoracic aorta (ATA) and proximal descending thoracic aorta (DTA) from untreated wild-type (uWT), treated wild-type (tWT), untreated R247C mutant (uR247C), and treated R247C mutant (tR247C) mice. RMSE denotes the root mean sum-of-the-squares of the error for the regression. Although individual parameters can be compared across groups, better interpretations come from resulting values of the linearized stiffness and stored energy (cf. Figures 3 and 5).

	$c$ (kPa)	$c_1^1$ (kPa)	$c_2^1$ (-)	$c_1^2$ (kPa)	$c_2^2$ (-)	$c_1^{3,4}$ (kPa)	$c_2^{3,4}$ (-)	$\alpha_0$ (deg)	RSME (-)
<b>ATA</b>									
uWT	32.38	12.67	0.04	15.00	0.24	4.57	0.50	42.08	0.060
tWT	29.06	9.54	0.10	14.40	0.47	3.50	0.82	41.76	0.053
uR247C	20.01	6.62	0.12	16.02	0.16	3.70	0.40	41.31	0.064
tR247C	26.10	6.89	0.19	20.94	0.26	5.69	0.63	41.25	0.058
<b>DTA</b>									
uWT	45.17	0.65	1.60	13.14	0.64	0.20	2.50	30.75	0.061
tWT	33.73	2.96	1.26	14.04	0.48	0.27	2.53	28.57	0.054
uR247C	36.19	4.66	0.34	23.21	0.25	0.27	2.26	26.85	0.087
tR247C	28.25	2.11	0.99	18.18	0.36	1.12	1.46	25.52	0.078