

# A Pilot Study Characterizing Flow Patterns in the Thoracic Aorta of Patients With Connective Tissue Disease: Comparison to Age- and Gender-Matched Controls *via* Fluid Structure Interaction

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Prior computational and imaging studies described changes in flow patterns for patients with Marfan syndrome, but studies are lacking for related populations. This pilot study addresses this void by characterizing wall shear stress (WSS) indices for patients with Loeys-Dietz and undifferentiated connective tissue diseases. Using aortic valve-based velocity profiles from magnetic resonance imaging as input to patient-specific fluid structure interaction (FSI) models, we determined local flow patterns throughout the aorta for four patients with various connective tissue diseases (Loeys-Dietz with the native aorta, connective tissue disease of unclear etiology with native aorta in female and male patients, and an untreated patient with Marfan syndrome, as well as twin patients with Marfan syndrome who underwent valve-sparing root replacement). FSI simulations used physiological boundary conditions and material properties to replicate available measurements. Time-averaged WSS (TAWSS) and oscillatory shear index (OSI) results are presented with localized comparison to age- and gender-matched control participants. Ascending aortic dimensions were greater in almost all patients with connective tissue diseases relative to their respective control. Differences in TAWSS and OSI were driven by local morphological differences and cardiac output. For example, the model for one twin had a more pronounced proximal descending aorta in the vicinity of the ductus ligamentum that impacted WSS indices relative to the other. We are optimistic that the results of this study can serve as a foundation for larger future studies on the connective tissue disorders presented in this article.

Keywords: Marfan syndrome, computational modeling, patient-specific modeling, wall shear stress, Loeys-Dietz syndrome

# INTRODUCTION

Blood flow patterns in the aortic arch and descending thoracic aorta (dAo) are unique when compared with other portions of the arterial vasculature. For example, helical flow is present within the aortic arch under normal conditions and is thought to influence flow patterns at the origins of the carotid and subclavian arteries (1). Characterizing any deviations from these normal flow patterns may be important for optimal operative interventions involving the thoracic aorta with the goal of decreasing the likelihood of later, long-term aortic pathology. For instance, transient periods of turbulence during systole (due to modest differences in local vessel geometry from thoracic aortic diseases or surgery) could cause downstream flow disturbances. Such disturbances have been associated with local dilation (2, 3)and can subsequently impact indices of wall shear stress (WSS; defined as the tangential force per unit area exerted on a vessel wall as a result of flowing blood). Abnormal WSS has been related to pathology in the thoracic aorta. In a study of 10 middleaged adults with preexisting plaques, areas of low time-averaged WSS (TAWSS) were found in a rotating pattern progressing down the dAo and correlated with areas of atherosclerosis (4). Excessively high WSS can also be deleterious by initiating platelet aggregation (5).

Marfan syndrome is one type of connective tissue disorder that impacts several tissues/organs, including the cardiovascular system, where it can lead to thoracic aortic aneurysms (6). Prior research indicates that thoracic aortic aneurysms are the leading cause of death for patients with Marfan syndrome (7). Despite the uniqueness and importance of flow patterns in the thoracic aorta, there are few studies characterizing local blood flow patterns for connective tissue diseases beyond Marfan syndrome (8). To date, most computational studies of patients with Marfan syndrome have employed rigid computational fluid dynamics (CFD) models, which have the potential to reveal detailed spatiotemporal quantification of hemodynamic indices, including WSS, based on magnetic resonance imaging (MRI) and blood pressure (BP) data (9-19). A recent report quantified several indices by CFD, including WSS, for patients with Marfan syndrome before and after surgery to implement personalized external aortic root support (20). Some local differences were noted after surgery, but values were largely similar. This investigation builds from this prior work by conducting fluid structure interaction (FSI) modeling that includes local wall deformation of the aorta for patients with Loeys-Dietz, connective tissue disease of unclear etiology, and native (i.e., untreated) Marfan syndrome, as well as fraternal (i.e., dichorionic) twin patients with Marfan syndrome who underwent valve sparing root repair. All results are interpreted relative to those from age- and gender-matched control patients. The approaches employed for this study assigned local tissue properties as well as physiological inflow profiles and outlet boundary conditions to replicate clinical measurements. With these advancements, we aim to more accurately replicate the physiological conditions for patients with these connective tissue anomalies, and therefore help to further reveal differences in WSS and related indices vs. control participants. Thus, this pilot investigation may aid future long-term studies of morbidity related to aortic vascular disease.

## METHODS

#### Magnetic Resonance Imaging

Following Children's Wisconsin Institutional Review Board approval, MRI was performed for patients with connective tissue anomalies. MRI data from control participants with ages, genders, and Reynolds numbers aiming to closely match these patients were also obtained. Prior to protocol enrollment, verbal and written information was provided, and informed consent was obtained from participants. Details of the patients and control participants are shown in **Table 1** (10 men and two women aged 18–55 years). Patients had various connective tissue diseases (Loeys-Dietz with native aorta; connective tissue disease of unclear etiology with native aorta in female and male patients; and an untreated patient with Marfan syndrome). All imaging was conducted as part of clinically ordered sessions or ongoing research.

Gadolinium-enhanced (0.4 ml/kg; gadodiamide, Omniscan®, GE Healthcare, Waukesha, WI, USA) MR angiography (MRA) was performed with a breath-held 3D fast gradient echo sequence using a 1.5T Symphony® scanner (Siemens Healthcare, Erlangen, Germany). Slice thickness was 2.0 mm, with 56–60 sagittal slices per volume. A  $384 \times 192$ acquisition matrix (reconstructed to  $384 \times 256$ ) was used with a field of view (FoV) of  $25 \times 42$  cm<sup>2</sup> (spatial resolution of 0.65 × 1.64 mm). Other parameters included a repetition time (TR) of 4.3 ms, echo time (TE) of 1.4 ms, and a flip angle of  $25^{\circ}$ .

Time-resolved, velocity-encoded two-dimensional anatomic and through-plane phase-contrast MRI (PC-MRI) was performed orthogonally in the ascending aorta (AscAo) near the main pulmonary artery, in the dAo at the level of the diaphragm, and orthogonal to the arch origins of the head and neck vessels. An additional PC-MRI scan was obtained through the aortic valve for all patients, and when possible for control participants. The data were used to create spatiotemporally varying computational model inlets delineated by the patient's aortic valve as previously described in detail (21, 22) and briefly summarized in the boundary conditions section below. Heart rates ranged from 82 to 92 bpm (with R-R ranging from 652 to 732 ms); 25 images were reconstructed for the average R-R interval. Imaging parameters included TR, TE, and flip angle of 46 ms, 3.8 ms, and 30°, respectively. The FoV was 30  $\times$ 22.5 cm<sup>2</sup> with an acquisition matrix of 256  $\times$  192 and a slice thickness of 7 mm, resulting in a voxel size of 1.17 mm  $\times$  $1.17 \,\mathrm{mm} \times 7 \,\mathrm{mm}$ . Subjects breathed freely during PC-MRI acquisition, and data were retrospectively gated to the cardiac cycle. After scanning, supine, bilateral upper and lower extremity BP assessment was performed using a Dinamap BP system (GE Healthcare, Waukesha, WI, USA). Cardiac indexes, aortic dimensions, and mean Reynolds numbers are provided in Table 1. Dimensions are taken from the computational models created using MRA data according to the details below, which we understand to generally be a diastolic representation of vessel morphology.

| <b>TABLE 1</b> Hemodynamic indices, diagnosis, and aortic dimensions. |
|-----------------------------------------------------------------------|
|-----------------------------------------------------------------------|

| Connective<br>tissue<br>disorder | Age | Gender | Diagnosis                                                                            | Operation                               | Cardiac<br>index<br>(L/min/m <sup>2</sup> ) | AscAo<br>diameter<br>(mm) | dAo<br>diameter<br>(mm) | AscAo/dAo<br>diameter<br>ratio | Difference in<br>AscAo/dAo<br>diameter vs.<br>control (%) | Reynolds<br>number<br>(dimension-<br>less) |
|----------------------------------|-----|--------|--------------------------------------------------------------------------------------|-----------------------------------------|---------------------------------------------|---------------------------|-------------------------|--------------------------------|-----------------------------------------------------------|--------------------------------------------|
| Loeys Dietz                      | 55  | Μ      | Genetic-TGFBR 2<br>mutation                                                          | n/a                                     | 3.0                                         | 35.0                      | 21.4                    | 1.63                           | 29.3                                                      | 923                                        |
| Control                          | 57  | Μ      | -                                                                                    | -                                       | 2.5                                         | 32.2                      | 25.5                    | 1.26                           | -                                                         | 910                                        |
| Unknown<br>etiology-male         | 38  | М      | Phenotypic diagnosis;<br>Genetic-negative for<br>FBN-1, TGFBR1,<br>TGFBR 2           | n/a                                     | 3.1                                         | 28.9                      | 19.0                    | 1.52                           | 17.4                                                      | 1,380                                      |
| Control                          | 32  | М      | -                                                                                    | -                                       | 4.5                                         | 26.9                      | 20.7                    | 1.30                           | -                                                         | 1,520                                      |
| Unknown<br>etiology-<br>female   | 24  | F      | Genetic-negative for<br>FBN1, FBN1 del,<br>TGFBR1, TGFBR2,<br>MYLK, MYH11, ACTA<br>2 | n/a                                     | 3.2                                         | 32.9                      | 18.1                    | 1.82                           | 20.7                                                      | 1,130                                      |
| Control                          | 23  | F      | -                                                                                    | -                                       | 2.0                                         | 22.0                      | 14.6                    | 1.51                           | -                                                         | 823                                        |
| Marfan<br>syndrome–<br>twin<br>A | 22  | Μ      | Genetic-FBN1 exon 30 mutation                                                        | Valve<br>sparing<br>root<br>replacement | 2.9                                         | 28.0                      | 17.9                    | 1.56                           | 24.0                                                      | 1,190                                      |
| Control                          | 26  | М      | -                                                                                    | -                                       | 2.5                                         | 26.3                      | 20.9                    | 1.26                           | -                                                         | 1,190                                      |
| Marfan<br>syndrome–<br>twin<br>B | 22  | Μ      | Genetic-FBN1 exon 30 mutation                                                        | Valve<br>sparing<br>root<br>replacement | 3.3                                         | 26.8                      | 16.2                    | 1.65                           | 13.5                                                      | 1,350                                      |
| Control                          | 24  | Μ      | -                                                                                    | -                                       | 3.5                                         | 28.8                      | 19.8                    | 1.46                           | -                                                         | 1,550                                      |
| Marfan<br>syndrome-<br>native    | 18  | М      | Phenotypic diagnosis                                                                 | n/a                                     | 6.0                                         | 22.8                      | 16.2                    | 1.41                           | 10.6                                                      | 2,710                                      |
| Control                          | 18  | М      | -                                                                                    | -                                       | 3.2                                         | 25.1                      | 19.7                    | 1.28                           | -                                                         | 1,520                                      |

Reynolds number calculations assume a blood density of 1.06 g/cm<sup>3</sup> and dynamic viscosity of 4 cP.

### **Computational Model Construction**

Computational versions of the aorta and arteries of the head and neck were created from MRA imaging data using Simvascular (Simtk.org) software as discussed previously (15). Models were discretized using a commercially available, automatic mesh generation program (MeshSim, Simmetrix, Clifton Park, NY, USA). Meshes contained  $\sim$ 4 million tetrahedral elements, and localized refinement was performed until results were independent of the mesh as discussed elsewhere (12) using an adaptive technique (23, 24) to deposit more elements near the luminal surface and in anatomical regions prone to flow disruption (14).

#### **Inlet Boundary Conditions**

PC-MRI data were used to calculate time-resolved volumetric blood flow as previously described (16, 25). A time-varying plug flow inlet based on the measured AscAo flow was created, but with a restricted cross section determined from time-varying PC-MRI magnitude data at the level of the valve (21). A normal trileaflet valve was assumed for control participants and confirmed using the imaging data mentioned above, when possible.

# Outlet Boundary Conditions and Wall Deformation

Flow from the innominate, left common carotid artery, left subclavian artery, and dAo were used together with BP data to prescribe physiological outflow boundary conditions using threeelement Windkessel approximations (26). The three-element Windkessel accounts for vessels distal to computational model branches using three parameters with physiological meaning, namely, characteristic resistance ( $R_c$ ), capacitance (C), and distal resistance ( $R_d$ ). The total arterial capacitance (TAC) for each patient was determined from inflow and BP data, assuming a characteristic to total resistance ratio of 6% (27). The TAC was then distributed among outlets according to their blood flow distributions (28). The terminal resistance ( $R_t = R_c + R_d$ ) was then calculated from mean BP and PC-MRI flow measurements and distributed between the remaining resistance parameters by adjusting  $R_c$  to  $R_t$  ratios (6–10%) for each outlet using the pulse pressure method, (29, 30) thereby replicating measured BP. An augmented-Lagrangian formulation (31) for constraining velocity profiles at model outlets was used to mitigate instabilities often occurring during flow deceleration and diastole. Vessel deformability (32) was included in FSI simulations as discussed elsewhere (15). Briefly, a wall thickness of 0.15 cm (33) was implemented from literature for all models and the Young's modulus was then adjusted iteratively until the AscAo mean luminal displacement was within 5% of the values obtained from PC-MRI magnitude measurements.

#### **Computational Simulations**

Simulations were performed using a novel stabilized finite element method to solve the conservation of mass (continuity), balance of fluid momentum (Navier-Stokes), and vessel wall elastodynamics equations (32). Simulations were run for 4–6 cardiac cycles until the flow rate and BP fields yielded periodic solutions.

Blood flow velocity, BP, and wall displacement were visualized using ParaView (Kitware, Clifton Park, NY, USA). TAWSS (34) and the oscillatory shear index (OSI) (25) was then calculated as previously described. Low TAWSS is generally thought to promote atherogenesis, as is elevated OSI, an index of directional changes in WSS. Low OSI indicates WSS is unidirectional, while a value of 0.5 is indicative of bidirectional WSS with a timeaverage value of zero. These indices were quantified in several ways. Values for TAWSS and OSI were extracted longitudinally along the inner and outer curvatures of the thoracic aorta, as well as along its anatomic right, and left sides as described previously (14, 35). This was done because prior imaging studies found that local values for these indices were statistically different from circumferential averages (36), thereby motivating the need to report detailed local WSS maps in computational studies. To visualize these indices locally, the surface of each vessel was unwrapped and mapped into a  $\theta$ , *l* rectangular domain, where  $\theta$  and l are the circumferential and longitudinal coordinates of each point on the vessel wall (37). Data presented represent an average of nearest-neighbor values in 2% increments along the length of the aorta. This distance was made consistent between patients and control participants using dimensional information from imaging data, and then normalized from 0 to 1. TAWSS values from 0 to 50 dyn/cm<sup>2</sup> are particularly interesting from the perspective of the vascular response to hemodynamics and are also highlighted in histograms (2 dyn/cm<sup>2</sup> bins) for values within this range. Histograms are also provided for the full range of OSI, 0-0.5 (0.2 unit bins) (22, 38).

### RESULTS

Fluid structure interaction simulations for all patients, and their respective controls, yielded the TAWSS and OSI results shown in **Figures 1**, **3**, respectively. The size of the models shown is relative to each other using the descending aortic outlet dimension as reference. A summary of dimensions following model construction is also provided in **Table 1**. In nearly all cases, the AscAo dimension was greater in patients than the respective control participant. Conversely, with the exception of the female

patient having connective tissue disease of unknown etiology, the dAo dimension was of smaller caliber compared to the age- and gender-matched control participant. The ratio of AscAo to dAo for patients was consequently 10.6–29.3% larger than AscAo:dAo ratio for control participants.

## Patient With Loeys-Dietz Disease

The patient with Loeys-Dietz disease and corresponding control had the lowest overall distributions of TAWSS due primarily to the large AscAo dimensions in the setting of normal range cardiac output values (e.g., Loeys-Dietz =  $3.0 \text{ L/min/m}^2$ ; control =  $2.5 \text{ L/min/m}^2$ ). The histogram of TAWSS values shown in **Figure 2** reveals the similarity in distributions between the Loeys-Dietz and control models. The smaller caliber of the dAo relative to the AscAo in the patient with Loeys-Dietz disease relative to its control (mentioned above) also results in slightly higher distributions of TAWSS along the last half of the thoracic aorta quantified (distances from ~0.5 to 1.0).

There were modest differences in OSI for the patient with Loeys-Dietz disease relative to the control participant (**Figure 3**). The histogram (**Figure 4**) shows similar amounts of elevated OSI values >0.4 for both models. The histogram from the patient with Loeys-Dietz disease showed that less of the luminal surface was exposed to elevated OSI (0.2–0.4) shown to be atherogenic in prior reports. The difference in OSI values within this region corresponded to the transverse arch of the control patient (**Figure 3**), which also seems to have slight out-of-plane morphology in this region compared to the more traditional arch of the patient with Loeys-Dietz disease. These differences can also be seen in the longitudinal plots (e.g., right, outer, and left surfaces at distances of ~0.4–0.6).

# Patients With Connective Tissue Disease of Unknown Etiology

The TAWSS histogram for the male patient with connective tissue disease of unknown etiology reveals a greater portion of the aorta and its branches were exposed to lower TAWSS, particularly in the ascending aorta, relative to control (**Figure 2**). This is clearly shown in longitudinal TAWSS plots along the outer and left luminal surfaces. Vascular dimensions and morphology were similar for both models, suggesting that these differences may primarily be due to a higher cardiac index for the control participant (e.g., 3.1 L/min/m<sup>2</sup> for the patient with connective tissue disease vs. 4.5 L/min/m<sup>2</sup> vs. for the control).

The ascending aorta of the male patient with connective tissue disease of unknown etiology was exposed to a greater area of elevated OSI values in the range of  $\sim$ 0.1–0.4 (**Figure 3** and histogram of **Figure 4**), which occurred along nearly the full length of the left luminal surface and lasted 50–70% of the distance for the right and outer luminal surfaces (**Figure 4**).

The female patient with connective tissue disease of unknown etiology experienced higher TAWSS along the transverse arch and dAo (**Figure 1**). This is reflected in all longitudinal plots (**Figure 2**). This finding appears to correspond to a larger AscAo/dAo ratio for the patient compared to the control.

The larger caliber ascending aorta and its branches for the female patient with connective tissue disease of unknown



etiology resulted in a greater area of potentially deleterious OSI values in this region (**Figure 3**) and more area exposed to the full range of OSI compared to the corresponding control participant. Elevated OSI values were most pronounced along the left luminal surface (**Figure 4**).

### Patients With Marfan Syndrome

TAWSS for the native (i.e., uncorrected) patient with Marfan syndrome was extremely high relative to the corresponding control participant primarily due to a cardiac index exceeding normal conditions (6.0 L/min/m<sup>2</sup> vs. 3.2 L/min/m<sup>2</sup> for the control). The distributions of OSI values were somewhat similar for values > 0.2. By qualitative assessment, the AscAo of the native patient with Marfan syndrome had a greater area exposed to lower OSI (**Figure 3**), primarily along the inner and left luminal surfaces (**Figure 4**). Of note, the native patient with Marfan syndrome had the highest Reynolds number of all patients or participants studied (**Table 1**), and the value suggests flow was not laminar in the aorta of this patient, likely leading to spatial differences in OSI within the AscAo.

The twin patients with Marfan syndrome, who previously underwent aortic root replacement, had similar TAWSS distributions. The computational model labeled as Twin B for this study had a more dilated proximal descending aorta in the vicinity of the ductus ligamentum, which impacted indices of WSS. Differences between the computational models of each twin and their respective control model were mainly a function of cardiac index. TAWSS was generally lower in the dAo of both twins after surgery compared to their respective controls (**Figure 1**). This was most pronounced along the left luminal surface of the AscAo (**Figure 2**).

Distributions (Figure 3) and histograms (Figure 4) of OSI values for the twin patients with Marfan syndrome, who previously underwent aortic root replacement, were also similar with morphological differences in the vicinity of the ductus ligamentum mostly impacting the luminal surface at a normalized distance of  $\sim 0.4-0.6$ . This area was not replaced in either subject and reflects their native aortic properties.

## DISCUSSION

Blood flow patterns in the aortic arch and dAo are unique compared with other portions of the arterial vasculature. This is evident when considering the Reynolds number, a dimensionless index used to characterize fluid flow (2). Specifically, Reynolds number describes the ratio of convective inertial forces to viscous forces. In general, Reynolds number values <2,200 constitute laminar flow where adjacent layers of fluid move in layers without mixing, while those >2,200 may be characterized as transitional or turbulent depending on specific details of the local flow domain. Under normal conditions, the thoracic aorta experiences Reynolds numbers on the order of 1,500 (mean) as a result of its large caliber and high blood flow rates. These Reynolds number values indicate blood flow is generally laminar throughout the cardiac cycle, but there are undoubtedly portions



rigure 2 [Local quantification of TAWSS results. The top row shows histograms (2 dyn/cm<sup>2</sup> bins) of the area (cm<sup>2</sup>) from each model surface exposed to TAWSS values from 0 to 50 dyn/cm<sup>2</sup> (connective tissue disease = solid bars; age- and gender-matched controls = hollow bars). Longitudinal TAWSS distributions along the outer, anatomic right, anatomic left, and inner curvatures of the aorta are also shown in subsequent rows for the patients with various connective tissue disease (dashed lines) vs. their associated age- and gender-matched controls (solid lines). Low TAWSS is generally thought to promote atherogenesis, so skewing of histogram results toward lower values could be interpreted as less ideal, as could overall lower values of TAWSS along the aortic surfaces.

of the cardiac cycle during which blood flow transiently becomes transitional and/or turbulent. Characterizing any deviations from these normal flow patterns may be important for optimal operative interventions for the thoracic aorta with the aim being to decrease the likelihood of later, long-term aortic pathology. For example, transient periods of turbulence during systole (due to modest differences in local vessel geometry from thoracic aortic diseases or surgery) could cause downstream disturbances.

This investigation builds upon prior work by conducting FSI, in contrast to rigid CFD, with modeling that includes local wall deformation of the aorta for patients with Loeys-Dietz, connective tissue disease of unclear etiology, and native (i.e., untreated) Marfan syndrome, as well as twin patients with Marfan syndrome who underwent valve sparing aortic root repair. We matched changes in dimension by PC-MRI measurements by assigning local tissue properties and iterating Young's modulus until the deformation matched that observed *in vivo*. Physiological inflow profiles are uniquely implemented by imposing a restricted cross-sectional flow determined from time-varying PC-MRI magnitude data at the level of the valve (21)

and outlet boundary conditions for PC-MRI data are added to reflect clinical measurements (14, 15). Our goal with these model improvements was to more accurately replicate the physiological conditions for patients with these connective tissue anomalies, and therefore reveal differences in WSS and OSI vs. age- and gender-matched control participants in a pilot study.

In nearly all cases, the AscAo dimension was greater in patients with connective tissues diseases relative to their respective control. With the exception of the female patient with connective tissue disease of unknown etiology, the dAo dimension was of smaller caliber when compared to the corresponding control. Differences in TAWSS and OSI were driven by these local morphological differences and cardiac output. Unfortunately, it is difficult to speculate on the cause of smaller dAo dimensions in most patients relative to controls given the small sample size and heterogeneity. This finding should be validated in larger studies with groups of patients having similar genetic anomalies.

A unique presentation of the results of this study centers on the twin patients with Marfan syndrome who previously



underwent aortic root replacement. The computational model labeled as Twin B for this study had a more pronounced proximal descending aorta in the vicinity of the ductus ligamentum that impacted indices of WSS relative to Twin A. As with the other patients studied here, differences between the computational results of each twin and their respective control were mainly a function of cardiac index. Nonetheless, TAWSS was generally lower in the AscAo of both twins after surgery compared to their respective controls (Figure 1). Interpretation of these findings prompted review of literature that may be applicable. With twin gestation, there is a higher incidence of congenital heart disease. For monozygotic twins (65% with one chorion), the incidence of congenital heart malformations may be six times that for a singleton. For monozygotic twins, twin-twin transfusion syndrome may play a role in the development of CHD (39). The twins in our cohort were fraternal twins (i.e., dichorionic) inheriting the gene mutation for Marfan syndrome. Marfan syndrome is almost exclusively inherited in an autosomaldominant manner, although rare case reports have described recessive fibrillin 1 gene (FBN1) mutations (40). For these dichorionic twins, twin-twin transfusion does not explain the CHD, but rather genetic inheritance explains their clinical course. In other dichorionic twins with congenital heart diseases such a coarctation of the aorta, altered fetal-placental hemodynamics sometimes resulting from fetal growth restriction can contribute to the development of their congenital heart disease (41).

The results of this study build from and extend existing simulation results for these patient populations. For example, previous work has independently quantified stress and strain fields (42) for three patients with Marfan syndrome before and after surgery to implement personalized external aortic root support, as well as aortic flow patterns and resulting distributions of WSS using imposed flow and pressure waveforms allowing for qualitative comparison of velocity patterns to PC-MRI (20). Although the fluid flow version of the study used rigid wall CFD, the results are generally consistent with this study in that models created from data before personalized external aortic root support had larger AscAo diameters, leading to more flow disturbances (20). More recent work reported the ratio of circumferential to longitudinal WSS in idealized models informed by the data of patient with Marfan syndrome with stable or dilating aneurysms (43). To the best of our knowledge, there is a paucity of studies characterizing altered blood flow patterns for patients with connective tissue anomalies beyond Marfan syndrome. Despite the scarcity of such studies, there is a 2015 report that quantified aortic dimensions and indices of aortic stiffness in patients with connective tissue disorders using MRI (8). While the finding of elevated stiffness in patients with connective tissue disorders from this study suggests that rigid CFD models may be appropriate for this patient population,



FIGURE 4 | Local quantification of oscillatory shear index (OSI) results. The top row shows histograms (0.2 unit bins) of the area (cm<sup>2</sup>) from each model surface exposed to OSI values from 0 to 0.5 (connective tissue disease = solid bars; age- and gender-matched controls = hollow bars). Longitudinal OSI distributions along the outer, anatomic right, anatomic left, and inner curvatures of the aorta are also shown in subsequent rows for the patients with various connective tissue disease (dashed lines) vs. their associated age- and gender-matched controls (solid lines). Higher OSI values are generally thought to promote atherogenesis, so skewing of histogram results toward higher values could be interpreted as less ideal, as could overall higher values of OSI along the outer, anatomic right, anatomic left, and inner curvatures of the aorta.

it also points to the importance of accurately replicating deformation in patients with connective tissue disorders, who may not yet have experienced an increase in stiffness and associated adverse changes in WSS indices.

The results of this study should be considered relative to several potential limitations. Our investigation studied alterations in WSS indices locally in the proximal thoracic aorta and its branches given the hallmark capacitive function for this region of the arterial vasculature. There is evidence, however, that connective tissue disorders such as Marfan syndrome may also impact central aortic flow dynamics by virtue of altered distal resistance vessels (44). Although our outlet boundary conditions do account for downstream vascular resistance, the impact of changes in resistance vessel due to each connective tissue disease was not explicitly included in the outflow boundary conditions imposed for this study. To date, it is not known which specific, or combination of, WSS indices are directly linked to vascular pathologies for patients with connective tissue diseases. The influence of each index likely also depends on the patient population and its predominant pathology (e.g., stiffening, neointimal hyperplasia, and aneurysm rupture) (4, 45, 46). The small sample size for each connective tissue disease does undoubtedly present a limitation to extrapolating the results to the full population of patients with Loeys-Dietz, patients with connective tissue disease of unclear etiology, and those with Marfan syndrome (native and status post valve sparing root replacement). The cardiac index for the native (i.e., uncorrected) patient with Marfan syndrome was higher than expected based on elevated heart rate and hyperdynamic left ventricular ejection fraction. It is also possible that this patient had some anxiety during the MRI session. The goal of this study was to remain patient specific in terms of boundary conditions. However, when considering the results of this study, it may be interesting to conduct an idealized parameter-based study to quantify the impact of morphology and cardiac output independently. Such work is planned for the future. The Lagrangian multiplier

approach implemented for constraining outlet velocity profiles can be replaced in future work by a backflow stabilization treatment that is thought to be less intrusive to the flow field, computationally inexpensive, and has been implemented in Simvascular (47). While this study did impose a plug velocity profile at model inlets, it was restricted by the time-varying cross section determined from PC-MRI using novel methods previously developed in our lab (21). Upon implementation of these methods, we quantified the impact of valve morphology on aortic hemodynamics and identified regions most influenced by the inlet, including the ascending aorta that has previously been a site of dilation for patients with Marfan syndrome as a result of local flow patterns. Besides this approach, computational studies of the thoracic aorta to date have typically introduced blood flow in one of two ways. In one approach, PC-MRI is used to temporally sample the velocity profile downstream of the valve and input this measured profile directly into the model. While not directly including the valve, its impact can be manifested in the data that is obtained, but this requires appropriate through- and in-plane velocity encoding to adequately resolve flow features being input into the CFD model. This approach may be difficult to implement within the constraints of a clinical setting as it can require specialized sequences not routinely implemented and obtains data that are more detailed than those commonly used in clinical diagnosis. One alternative approach has been to construct CFD models with their inlet beginning just distal to the aortic sinuses, impose the blood flow waveform measured downstream as an assumed velocity profile at the model inlet, and allow the curvature of the arch to influence resulting flow patterns (14). While this technique does not use the complete spatial velocity information, it does not require specialized sequences, minimizes the introduction of noise at the model inflow due to inadequate velocity encoding, and allows for improved temporal resolution compared to 3component PC-MRI (48). The methods of Wendell et al. used in this study (21) allow for more accurate representation of the impact of the aortic valve on computational studies of the thoracic aorta while still using data obtained as part of a routine clinical imaging session. The patients with connective tissue disorders of unknown etiology were suspected to have hereditary (or genetic) aortopathy, but negative testing for known genetic variants. The yield for current gene panels for thoracic aortic disease is only  $\sim$ 30%, even in patients with high clinical suspicion. Hence, unfortunately, it is a common situation in our clinic to have suspected genetic etiology but a negative gene panel. This may be interpreted as not yet identifying the applicable gene(s) in that individual/family. We often pursue whole-exome sequencing in such patients, but even then the results often do not identify a causative genetic variant. This is a limitation that we are working to mitigate in the future as having complete genetic data for future cohorts would greatly enhance our understanding of the results presented for larger populations of patients.

# CONCLUSION

The methods employed represent some of the most advanced vascular modeling tools available such as deformable walls, dynamically varying valvular area at the inlet of the model, physiological boundary conditions, and the use of age- and gender-matched controls. Despite some potential limitations outlined above in implementing these tools, the lack of computational modeling data for those patients with connective tissue diseases makes the current pilot data interesting and relevant. We are optimistic that the results of this study can serve as a foundation for larger future studies with the connective tissue disorders presented here.

# DATA AVAILABILITY STATEMENT

The datasets presented in this article are not readily available because raw data supporting the conclusions of this article can only be available by the authors to the extent possible by the institutional approvals governing the research presented. Requests to access the datasets should be directed to jladisa@mcw.edu.

# ETHICS STATEMENT

Ethical review and approval was obtained for the study using data from human participants in accordance with local legislation and institutional requirements. The patients/participants provided their written informed consent to participate in this study.

# **AUTHOR CONTRIBUTIONS**

JAC: patient recruitment clinical expertise for manuscript development/review. HW and RD: computational simulations, analysis/interpretation of computational results, and approval of the article. MS and JRC: patient recruitment, supervision of MRI scanning, and clinical expertise for manuscript development/review. JL: concept/design, methodological developments, analysis/interpretation of computational results, drafting article, and approval of article. All authors contributed to the article and approved the submitted version.

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