

Supplemental file 1: Detailed Patient Demographics

For MFS TAA tissue, sex was distributed: 63.64% male and 36.36% female. One individual identified as Hispanic while all others were non-Hispanic. The average age was 31 ± 2 years. Non-TAA control aortic tissues were harvested from the ascending aorta of heart transplant donors or recipients; mean ages are 58 ± 6 years; aortic diameters are 3.8 ± 0.2 cm with no prior diagnosis of MFS. For MFS TAA plasma, sex distribution was 54.79% male and 45.21% female. The average age was 38.48 ± 16.98 years. Non-TAA control plasma was 50.82% male 49.18% female. The average age was 42.08 ± 16.63 years. Further details are listed below:

Control Plasma Demographics Summary

- Sex:
 - Male (n = 62)
 - Age: 42.11 ± 15.72
 - Female (n = 60)
 - Age: 42.05 ± 17.66
- Ethnicity:
 - Unknown (n = 19)
 - Age: 43.37 ± 16.55
 - American Indian/Alaska Native (n = 6)
 - Age: 41.83 ± 10.87
 - Asian (n = 2)
 - Age: 19.5 ± 2.12
 - Black/African American (n = 3)
 - Age: 34.33 ± 14.57
 - Hispanic/Latino (n = 12)
 - Age: 33.92 ± 19.84
 - Other (n = 1)
 - Age: 61
 - White/Caucasian (n = 79)
 - Age: 43.66 ± 16.25

Marfan TAA Plasma Demographics Summary

- Sex:
 - Male (n = 40)
 - Age: 39.52 ± 17.72
 - Female (n = 33)
 - Age: 37.22 ± 16.21
- Ethnicity:
 - Asian (n = 1)
 - Age: 66.86
 - Black/African American (n = 2)
 - Age: 46.73 ± 12.93
 - Native Hawaiian/Pacific Islander (n = 2)
 - Age: 28.18 ± 6.00
 - Not Specified (n = 32)
 - Age: 35.18 ± 16.31
 - White (n = 36)
 - Age: 40.74 ± 17.42

SUPPLEMENT - DETAILED METHODOLOGY

Multiplex Suspension Array (MSA): MSAs were used to measure MMPs and TIMPs in human plasma and aortic tissue (FCSTM07 and LKTM003, R&D Systems) following manufacturer's instructions. In short, aortic tissues were homogenized in 4°C buffer (volume 1:6 weight/volume) containing 10 mM cacodylic acid pH 5.0, 0.15 M NaCl, 10 mM ZnCl₂, 1.5 mM NaN₃, and 0.01% Triton X-100. Homogenates were centrifuged (800 x g, 10 min, 4°C) and protein concentration was determined by BCA assay and 20 µg of total protein was analyzed. Plasma specimens were thawed on ice, centrifuged (16,000 x g, 4 min, 4°C) and diluted according to the manufacturer instructions. For MMP-2 and -9, samples were diluted 1:100; for all other MMPs, samples were diluted 1:10; for TIMPs, samples were diluted 1:20. The above assays were quantitated using a Bio-Rad Bioplex-200 analyzer. Results were compared to a standard curve from the same plate. Using analyte concentrations, fold changes from control values were calculated.

Immunoblotting analysis: Thawed tissues were transferred to 4°C homogenization buffer (buffer volume 1:6 weight/volume) containing 137 mM NaCl, 2.7 mM KCl, 8 mM Na₂HPO₄, and 2 mM KH₂PO₄, and disrupted in a 1mL glass tissue grinder. Homogenates were centrifuged (10,000 rpm, 2 min, 4°C) and the supernatants were used for immunoblotting. Ten µg, determined by BCA assay, of each aortic homogenate were loaded on a 4–15% gradient, bis-tris gel and fractionated by electrophoresis. Proteins were transferred to nitrocellulose membranes (0.45 µm, Bio-Rad, Hercules, CA) and incubated in antisera specific for MT1-MMP (1:2000; ab38971, Abcam) which was diluted in 5% bovine serum albumin (BSA)/PBS. A secondary peroxidase-conjugated antibody (primary antibody species-specific) was applied (1:5000, 5% BSA/PBS) and signals were detected with chemiluminescent substrate (SuperSignal West Pico PLUS Chemiluminescent Substrate, Thermo Fisher Scientific, Waltham, MA) and recorded using the Bio-Rad ChemiDoc XRS. Band intensity was quantified using ImageJ. Gels were normalized using two methods (further described in Results: *MT1-MMP protein abundance was increased in*

MFS TAA tissue). After performing MT1-MMP analysis, the nitrocellulose membranes were washed with a harsh stripping buffer containing 20 mL SDS 10%, 12.5 mL Tris HCl - pH 6.8 - 0.5 M, 67.5 mL distilled water, 0.8 mL β -mercaptoethanol. The membrane was then incubated in antisera specific for GAPDH (1:2500; Cat#31460, Thermo Fisher Scientific, Waltham, MA) diluted in 5% bovine serum albumin (BSA)/PBS. The second method of normalization was total protein densitometry from 20-120 kDa, using ultraviolet light activated trihalo. This normalization method provided the least amount of variation and was therefore used for data analysis.

Cell Culture: Primary aortic fibroblast cell lines (n=3) were established from human aortic biopsies of non-syndromic aneurysmal thoracic aortas using an established outgrowth technique as previously described.³ The isolated aortic fibroblasts were maintained in complete fibroblast-specific growth media (Fibroblast Growth Media 2; C-23020, PromoCell, Heidelberg, Germany) with added 10% fetal bovine serum (97068-085), and gentamicin (0.5 mg/mL; 15710-064, Gibco) at 37°C in 5% CO₂. Aortic fibroblasts from culture passages 2 to 10 were used in the following studies.

In Vitro Transfection: Human aortic fibroblasts were transfected for 18 hours at 50% confluence in 75 cm² flasks (430641U; Corning) using jetPRIME transfection reagent (114-15, Polypolus-transfection) according to the DNA transfection protocol with either 10 μ g MT1-MMP Over Expression vector (MT1-OE; OriGene, MC219253) or transfection reagent alone. Transfected fibroblasts were lifted with Tryp-LE (12604021, Thermo Fisher Scientific), and 200,000 cells per well were seeded in 6-well culture dishes. Serum-free culture media was exposed to the transfected cells and collected 48 hours later. Media was centrifuged at 2,000 x g for 10 minutes to remove debris and stored at -20°C.

Alteration of MT1-MMP Localization: MT1-MMP can be differentially located in aortic fibroblasts; protein kinase C-mediated phosphorylation of the C-terminal tail of MT1-MMP causes translocation of MT1-MMP from the membrane to endosomal vesicles in the cytoplasm.

Treatment with phorbol 12-myristate 13-acetate (PMA), which activates both classical and novel PKCs, results in enhanced internalization of MT1-MMP. Conversely, translocation of MT1-MMP can be inhibited with Röttlerin, a PKC- δ -specific inhibitor. PMA was used to induce internalization of MT1-MMP in isolated fibroblasts while Röttlerin was used to confine it to the cell surface. Primary human aortic fibroblasts were seeded in 6-well plates with 200,000 cells per well and cultured in complete growth media. Cells were cultured (37°C, 5% CO₂) until >75% confluence and treated for 24 hours: PMA (100 nM), Röttlerin (3 μ M), or vehicle control (DMSO, 1:1000). After transfection, the cells were exposed to serum-free culture media. The conditioned serum-free media was collected after 24 hours exposure, centrifuged at 2,000 x g for 10 minutes to remove debris, and stored at -20°C.

Soluble EMMPRIN ELISA: A human EMMPRIN ELISA (ab219631, Abcam) was performed on the conditioned media according to protocol. In brief, 50 μ L of standards or sample were added to the appropriate wells and exposed to 50 μ L of antibody cocktail. After 1 hour of incubation at room temperature, the liquid was aspirated, and the wells were washed three times with 350 μ L of the recommended wash buffer. Wells were incubated in 100 μ L of the recommended development solution for 10 minutes; development was stopped with an equal volume of the recommended stop solution. The plate was measured at an OD value of 450 nm.