

Precision Medical Care for Thoracic Aortic Disease: Findings from the Montalcino Aortic Consortium

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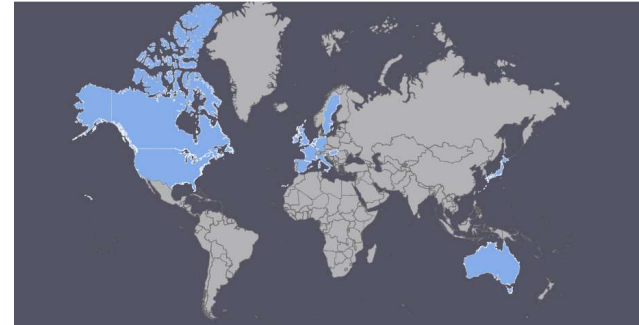
JOHN RITTER
RESEARCH PROGRAM
in AORTIC & VASCULAR DISEASE


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Heritable Thoracic Aortic Disease Genes “HTAD Genes”

ECM	<i>FBN1</i>	fibrillin-1	Marfan syndrome Vascular EDS
	<i>COL3A1</i>	collagen, type III, alpha-1	
	<i>LOX</i>	lysyl oxidase	
TGF-β Signaling	<i>TGFBR1</i>	TGF- β receptor type I	 Loeys-Dietz Syndrome
	<i>TGFBR2</i>	TGF- β receptor type II	
	<i>SMAD3</i>	SMAD family number 3	
	<i>TGFB2</i>	transforming growth factor β 2	
SMC Contraction	<i>ACTA2</i>	SM α -actin	
	<i>MYH11</i>	SM myosin heavy chain	
	<i>MYLK</i>	myosin light chain kinase	
	<i>PRKG1</i>	cGMP-dependent kinase I	

Montalcino Aortic Consortium: >30 Aortic Centers Worldwide



Inclusion criteria:

Any individual with a pathogenic or
VUS in an HTAD gene

Study Coordinator:

Ernesto Calderon Martinez, MD

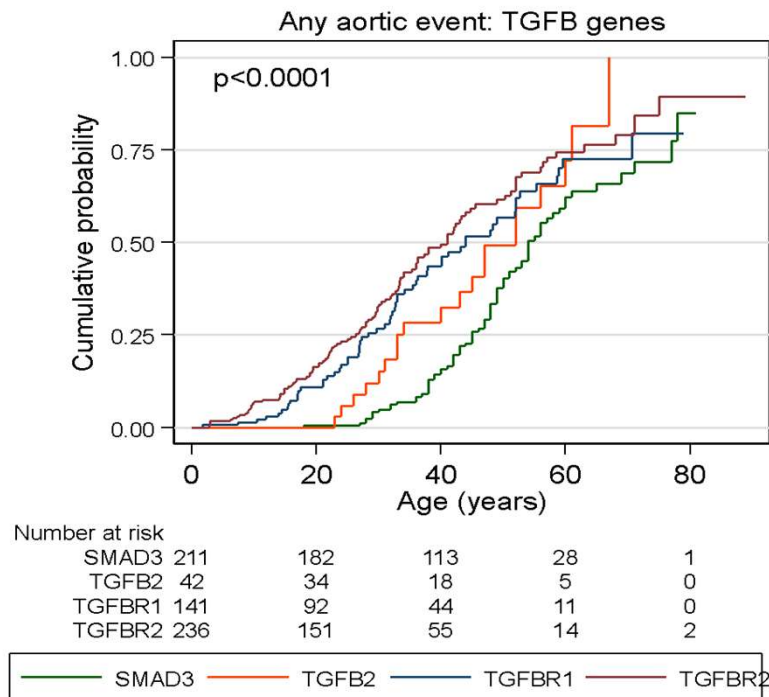
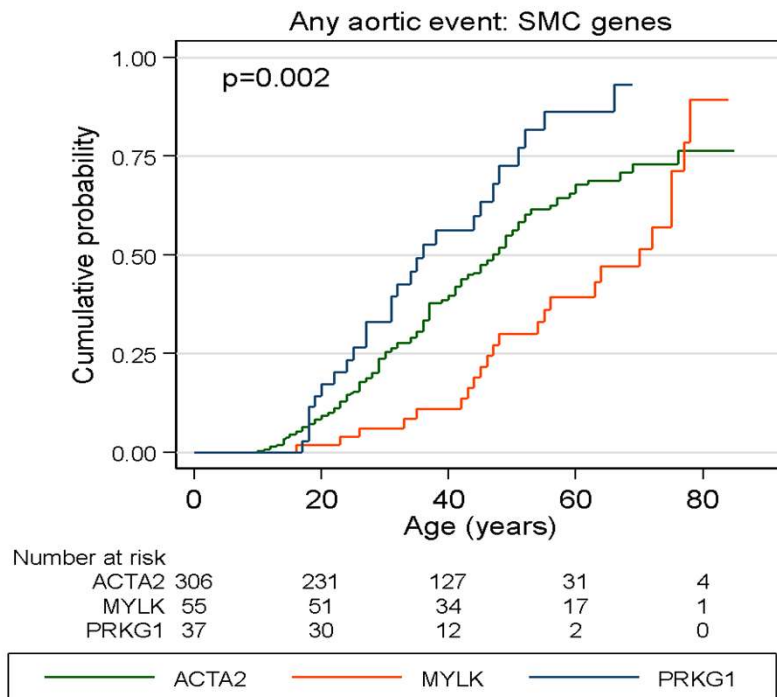
Email: MAC@uth.tmc.edu

ACC/AHA Treatment Guidelines 2022: Surgical Thresholds for Loeys Dietz Syndrome

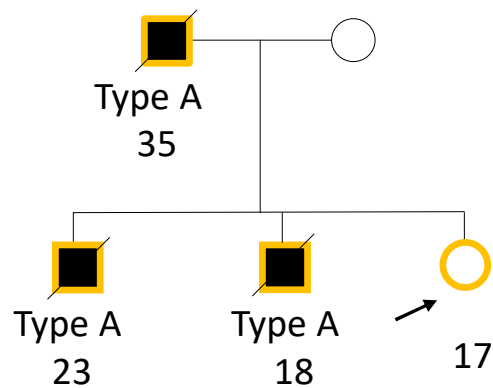
TABLE 11		Surgical Thresholds for Prophylactic Aortic Root and Ascending Aortic Replacement in Loeys-Dietz Syndrome Based on Genetic Variant		
COR	LOE (references)	Genetic Variant	Presence of High-Risk Features*	Aortic Diameter (cm)
1	C-LD ²	TGFBR1	No	≥4.5
1	C-LD ²	TGFBR2	No	≥4.5
2b	C-EO ²	TGFBR1	Yes	≥4.0
2a	C-LD ^{1,2}	TGFBR2	Yes	≥4.0
2a	C-EO ^{13,16}	SMAD3	-	≥4.5†
2b	C-EO ⁵⁻⁷	TGFB2‡	-	≥4.5†
2b	C-EO ^{9,23}	TGFB3	-	≥5.0†

Cumulative Risk for an Aortic Event Based on HTAD Gene

Aortic Event = Surgical repair of an aneurysm, type A, or B dissection



Gene-Based Management: Repair of a Normal Aorta in a 17 year old



PRKG1

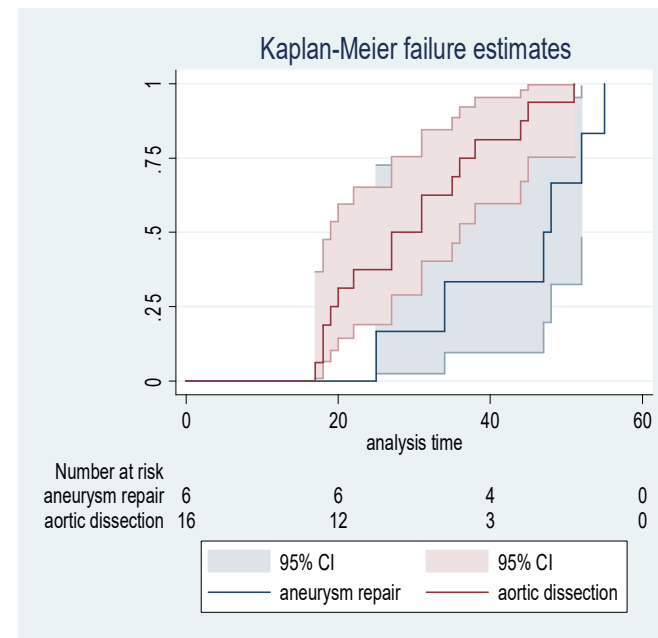





Photo used with permission

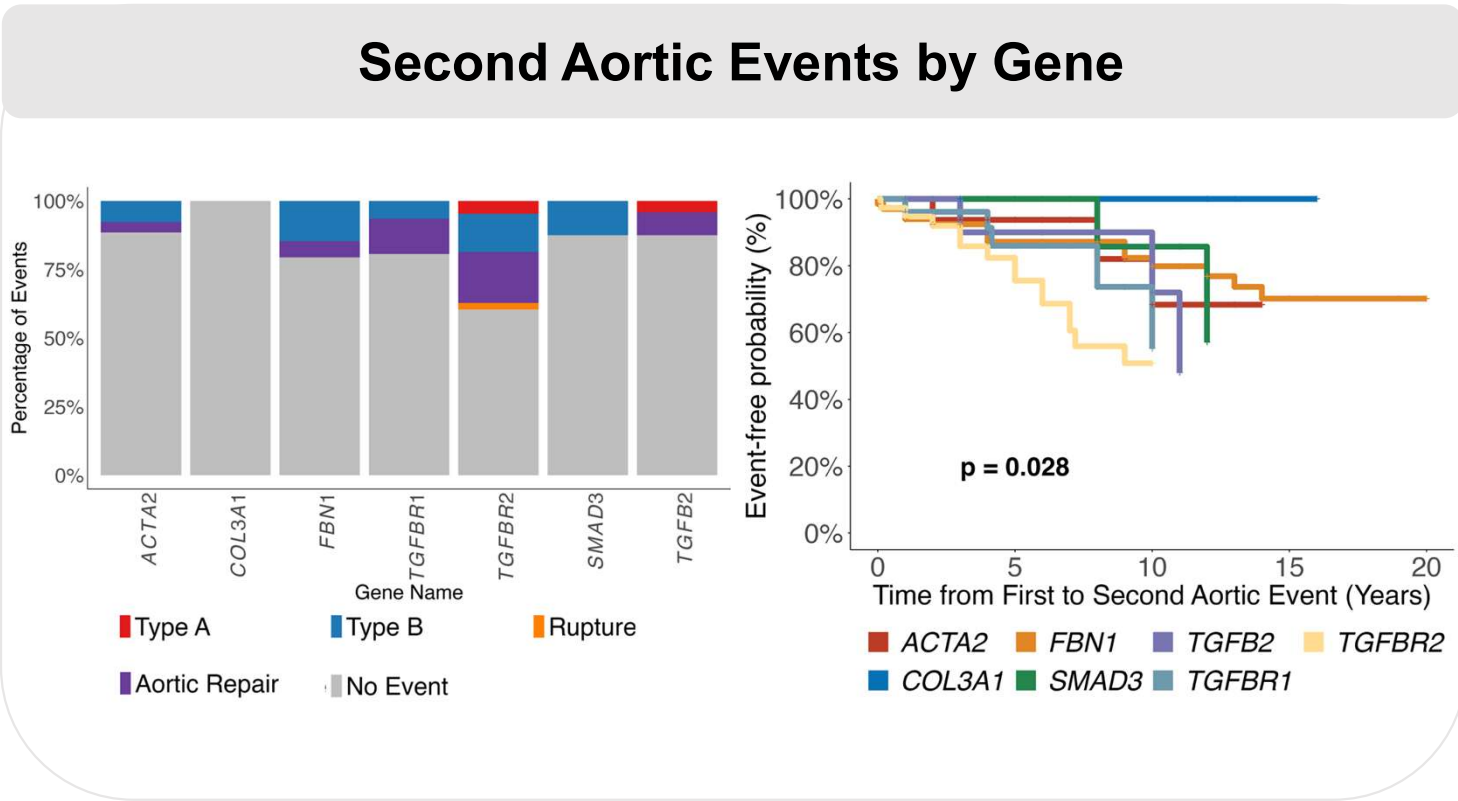
Asari A et al. *J Vasc Cases Innov Tech* 2025 12(2): 102088

Gene-Specific Risks for Aortic Events After Proximal Aneurysm Repair

Montalcino Aortic Consortium

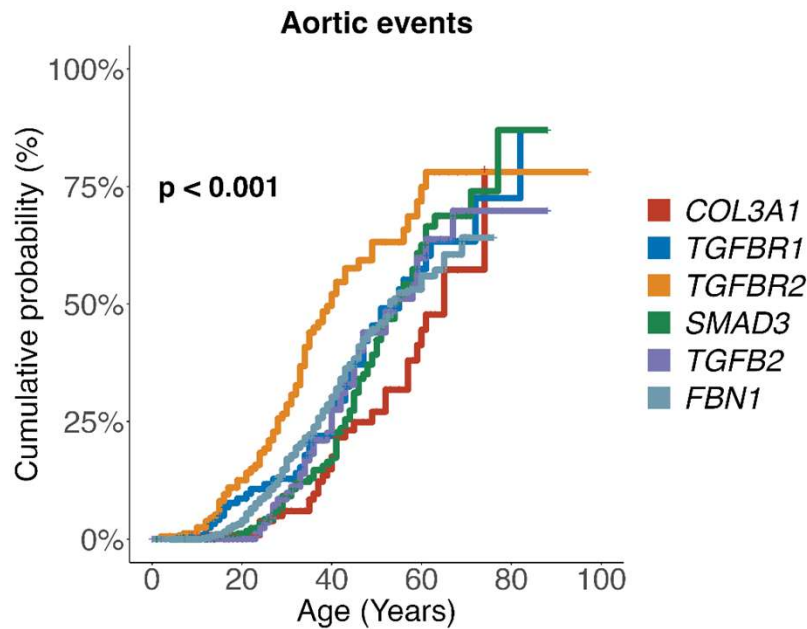
**Elective proximal aneurysm repair
n=229**



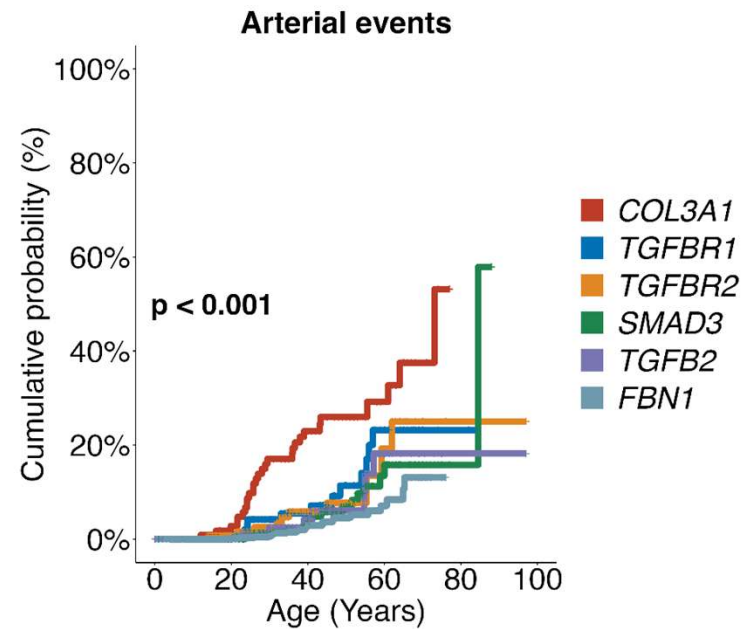
Second aortic events most prevalent in *TGFB2* cases (40%).

Calderon-Martinez et al., 2026, under revision.

Arterial Versus Aortic Events in VEDS, LDS and MFS: 2864 Cases

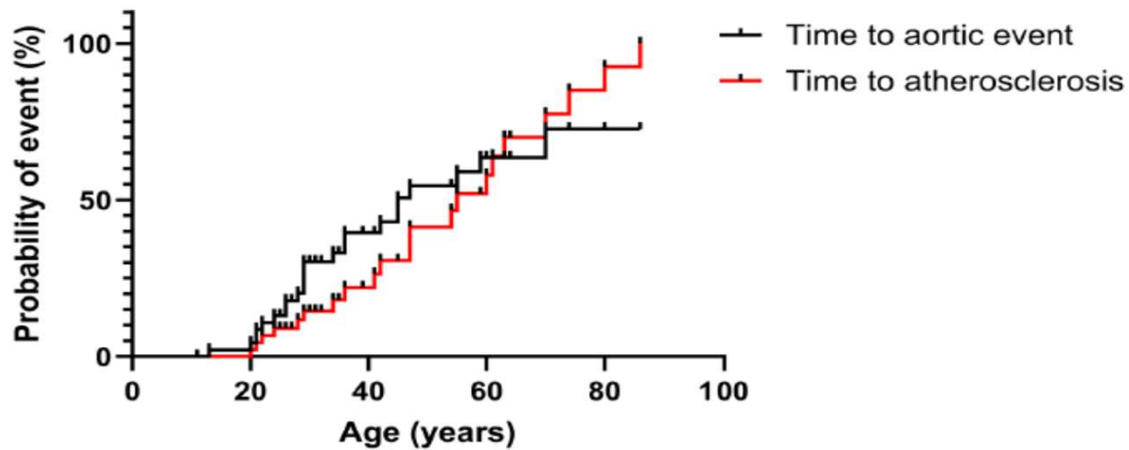
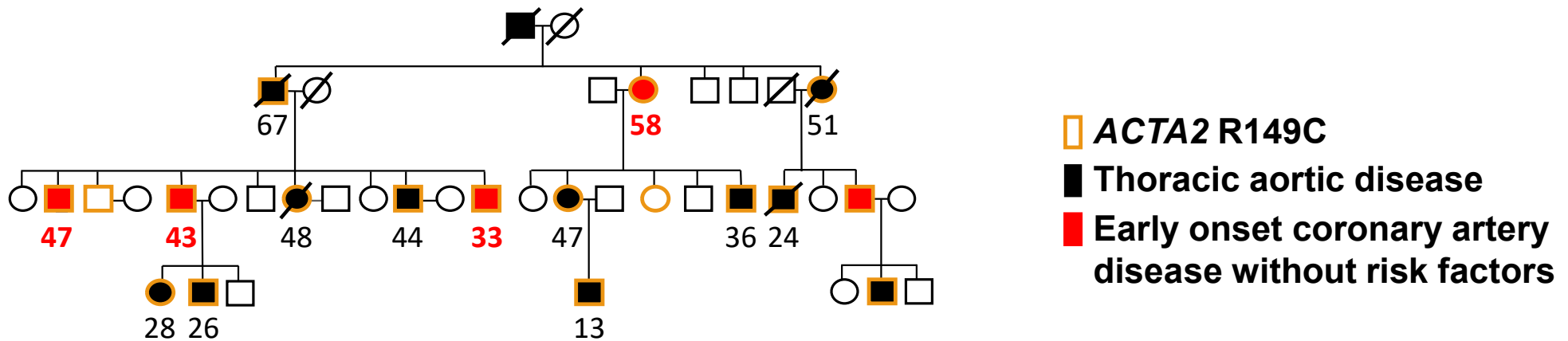


	0	20	40	60	80	100
COL3A1	125	101	63	18	0	0
TGFBR1	136	94	52	18	2	0
TGFBR2	167	112	40	9	1	0
SMAD3	196	167	95	22	1	0
TGFB2	125	90	47	11	1	0
FBN1	1009	678	262	50	0	0

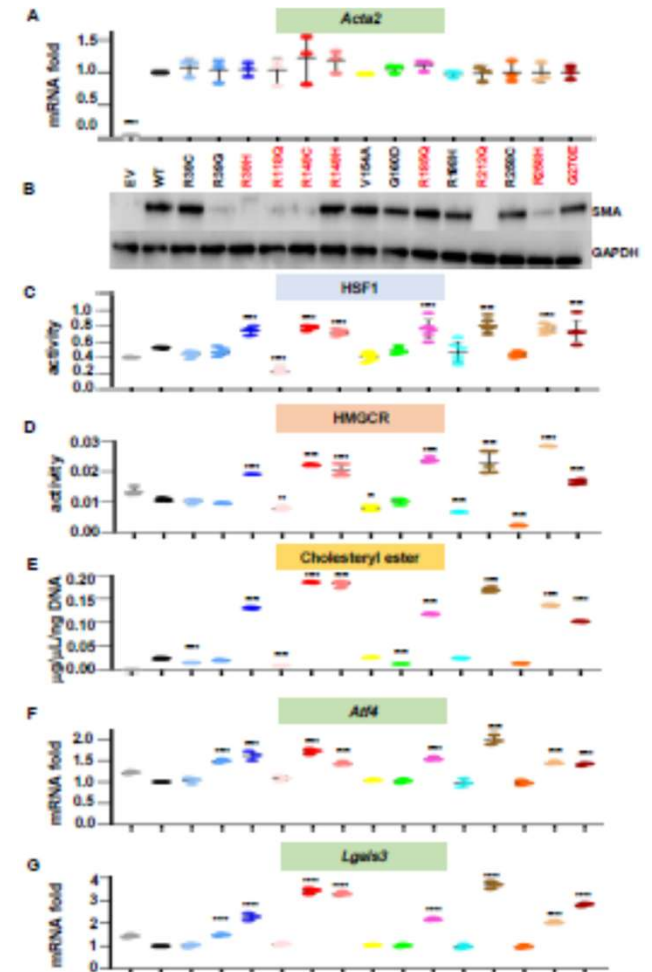
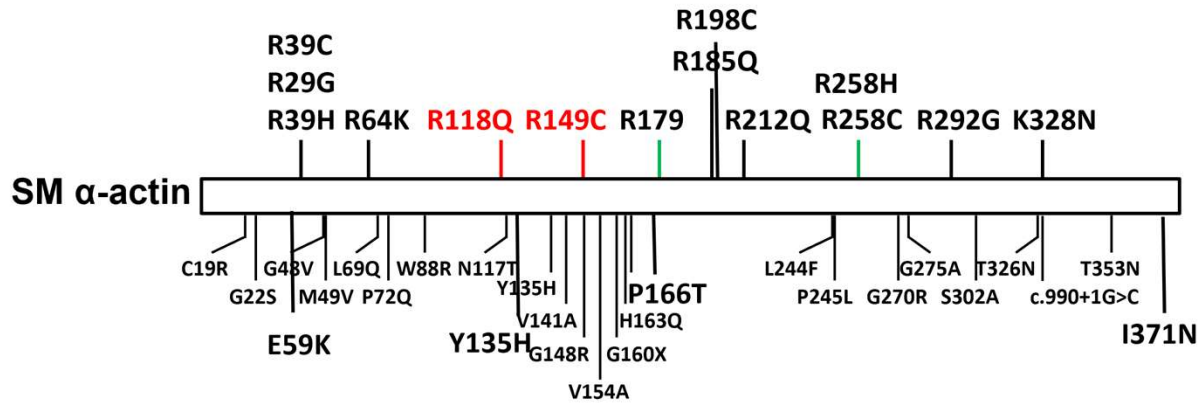


	0	20	40	60	80	100
COL3A1	125	99	63	20	0	0
TGFBR1	136	103	56	20	2	0
TGFBR2	167	121	68	14	1	0
SMAD3	196	169	105	37	2	0
TGFB2	125	90	54	13	2	0
FBN1	1009	696	324	75	0	0

ACTA2 R149C Variant and Early Onset Coronary Artery Disease



ACTA2 Pathogenic Variants Predisposing to Early Onset Atherosclerosis



Boerio et al, *Circulation Genome Prec Med*, 2026 Feb;19(1):e005169.

Putative Heritable Thoracic Aortic Disease Genes

ECM	<i>FBN2</i>	fibrillin 2
	<i>MFAP5</i>	microfibril associated protein 2
	<i>COL5A1</i>	collagen, type V, alpha-1
	<i>COL5A2</i>	collagen, type V, alpha-2
	<i>BGN</i>	biglycan
	<i>EFEMP2</i>	EGF-containing fibulin-like extracellular matrix protein 2
	<i>THSD4</i>	Adamts16
	<i>EMILIN1</i>	emilin
	<i>LTBP3</i>	latent TGF- β binding protein
TGF-β Signaling	<i>ELN</i>	elastin
	<i>SKI</i>	SKI protooncogene
	<i>SMAD2</i>	SMAD family number 2
	<i>SMAD4</i>	SMAD family number 4
	<i>TGFB3</i>	transforming growth factor, beta-3
SMC Contraction	<i>ARIH1</i>	ariadne RBR E3 ubiquitin protein ligase 1
	<i>FLNA</i>	filamin A
	<i>MYH9</i>	myosin, heavy chain 9
	<i>HCN4</i>	hyperpolarization-activated cyclic nucleotide-gated potassium channel 4
Other	<i>MAT2A</i>	methionine adenosyltransferases II α
	<i>FOXE3</i>	forkhead transcript factor E3
	<i>MTOR</i>	mechanistic target of rapamycin
	<i>CBS</i>	cystathionine beta-synthase
	<i>NOTCH1</i>	notch receptor 1
	<i>SLC2A10</i>	solute carrier protein 2, member 10
	<i>IPO8</i>	importin 8
	<i>PKD1</i>	polycystin 1
<i>PKD2</i>	polycystin 2	