

# Aberrant Regulation of EndMT in Turner Syndrome: Implications for the Pathogenesis of Congenital Cardiovascular Disease

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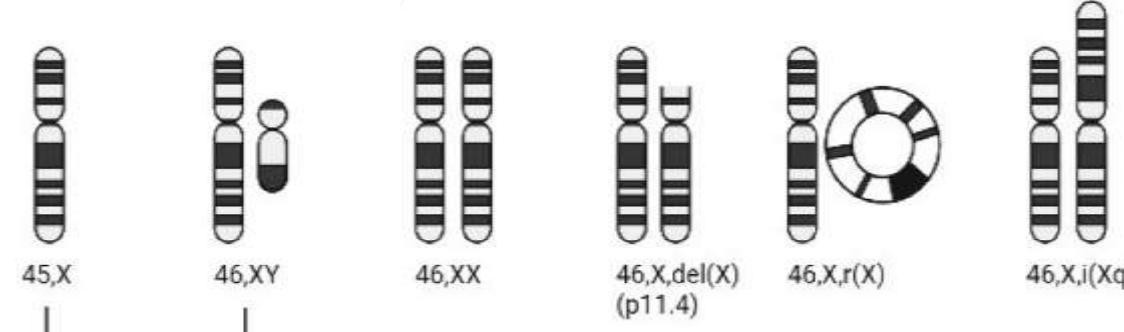
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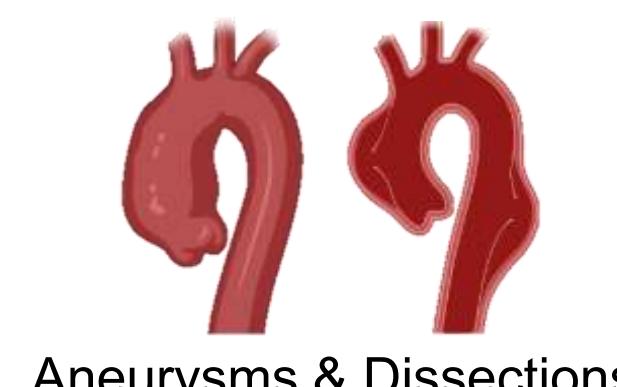
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## Introduction

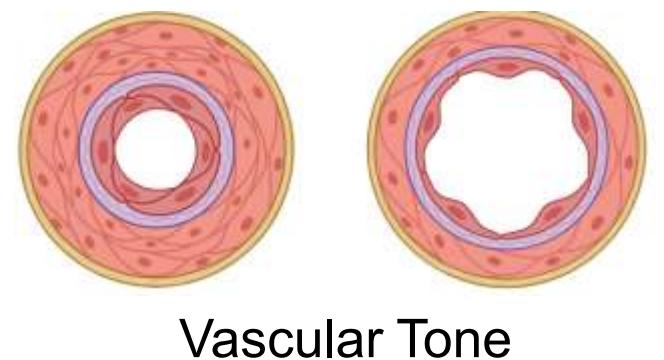
Turner Syndrome (TS) - Condition in which biological women have a half/partial dose of X-chromosome genes



Cardiac Complication



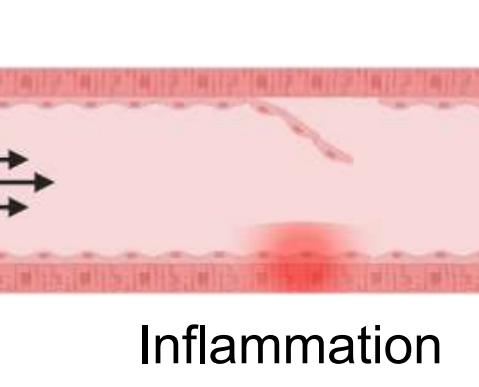
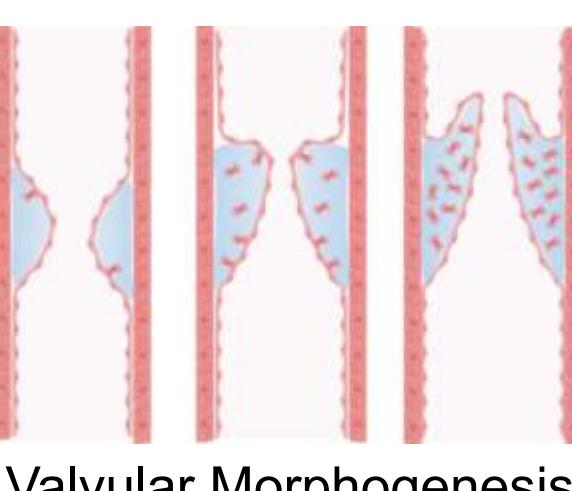
Endothelial Cell Dysfunction



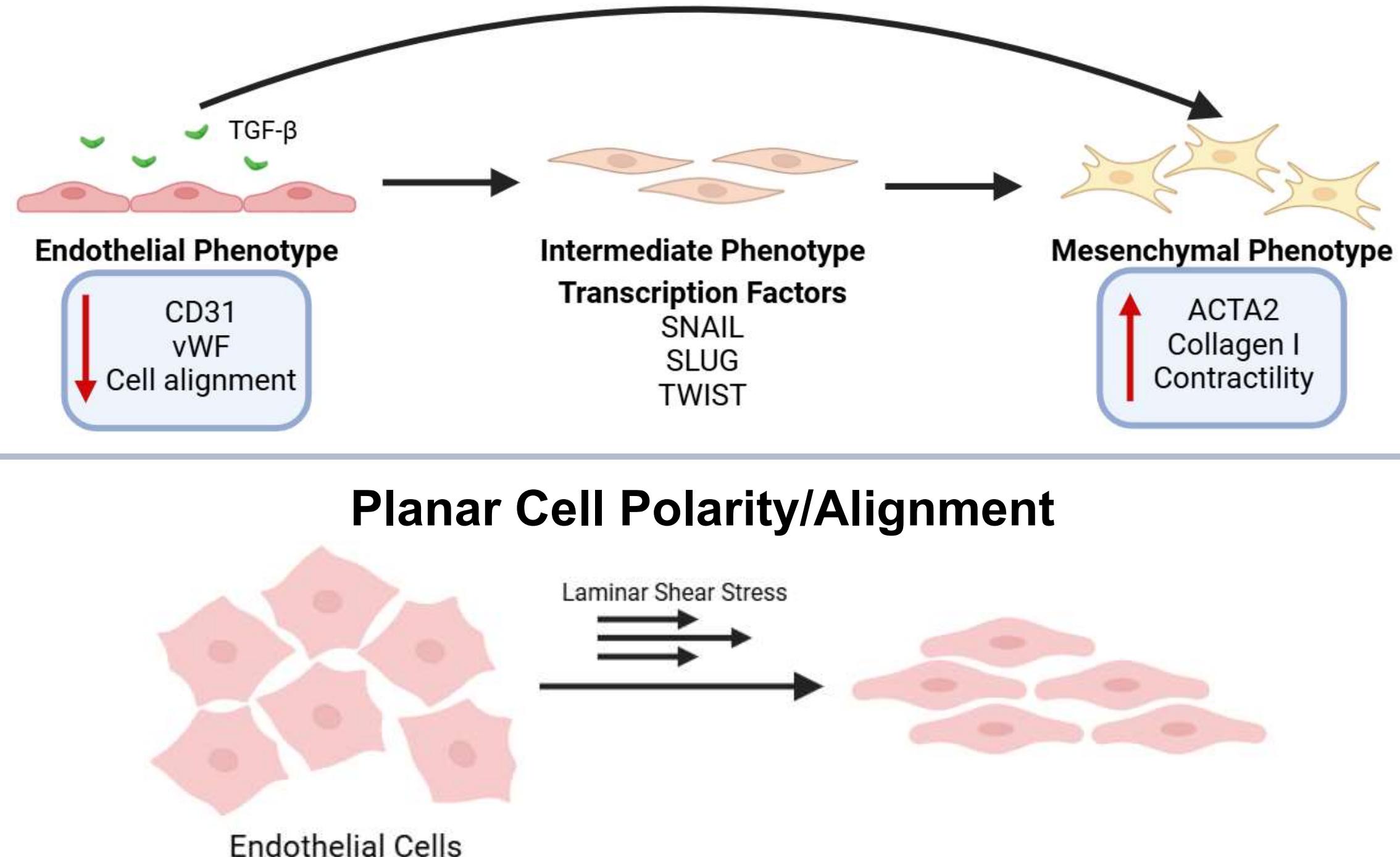
Planar Cell Polarity/Alignment  
Endothelial-Mesenchymal Transition  
Shear Stress  
Extracellular Matrix

None of these pathways & regulators have been explored with respect to X-chromosome dosage

Signaling Pathways & Mechanical Regulators



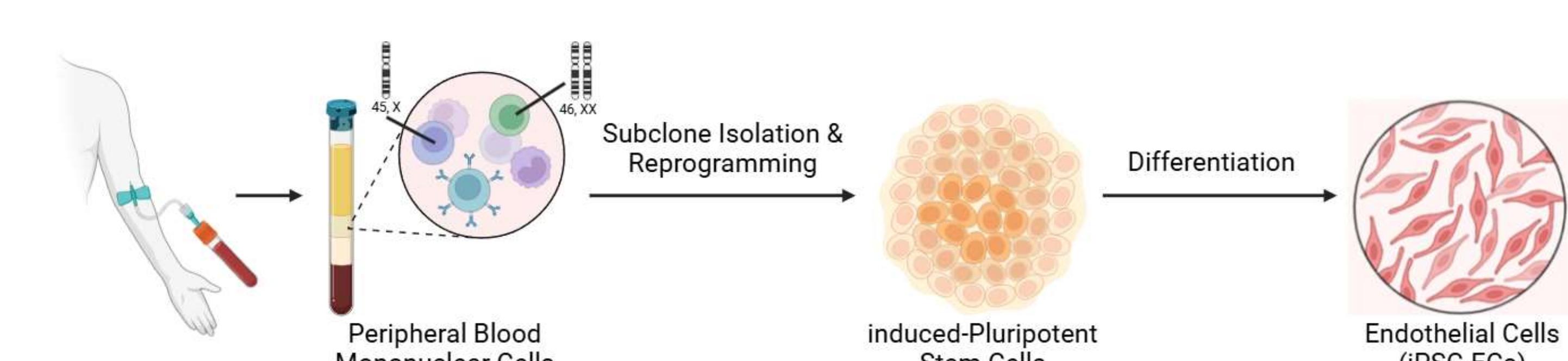
## Endothelial-Mesenchymal Transition (EndMT)



## Hypothesis

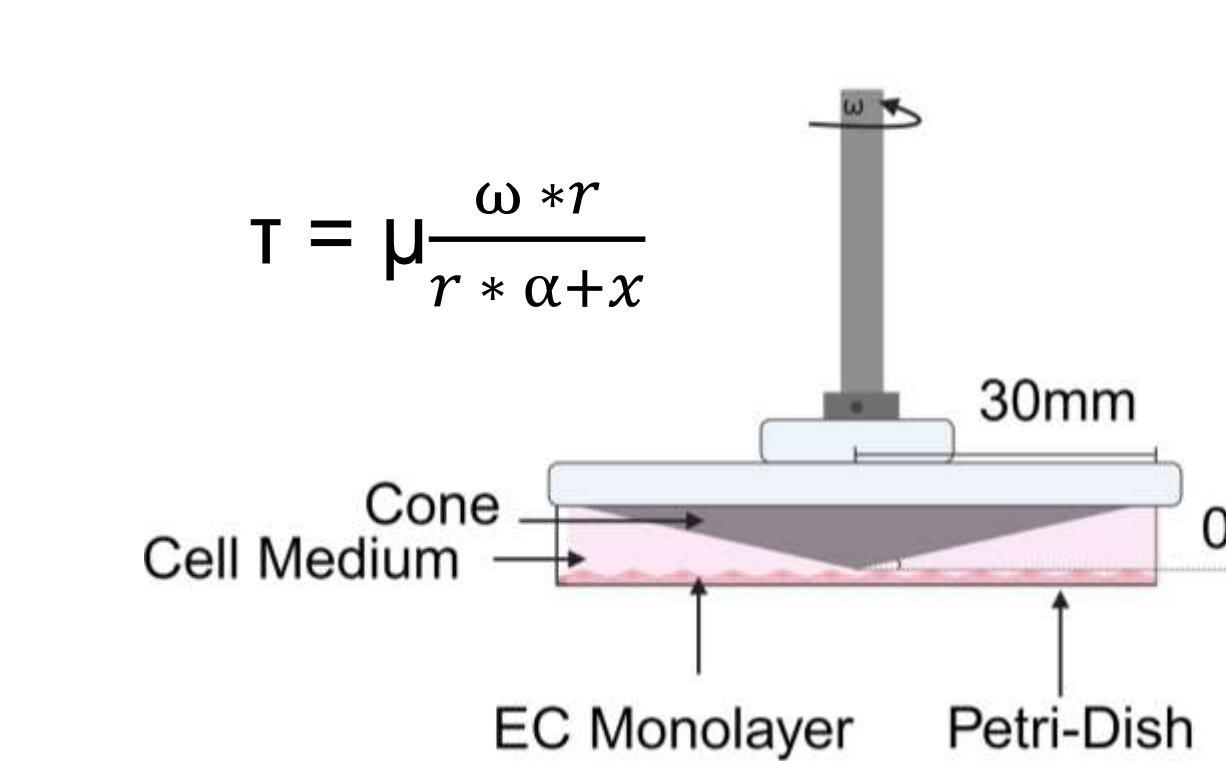
We hypothesize that disregulation of EndMT during cardiac development, at least partially due to abnormal sensing of shear stress by ECs, predisposes to congenital heart and vascular diseases in TS.

Two iPSC-EC karyotypes isolated from one mosaic patient



## Methods

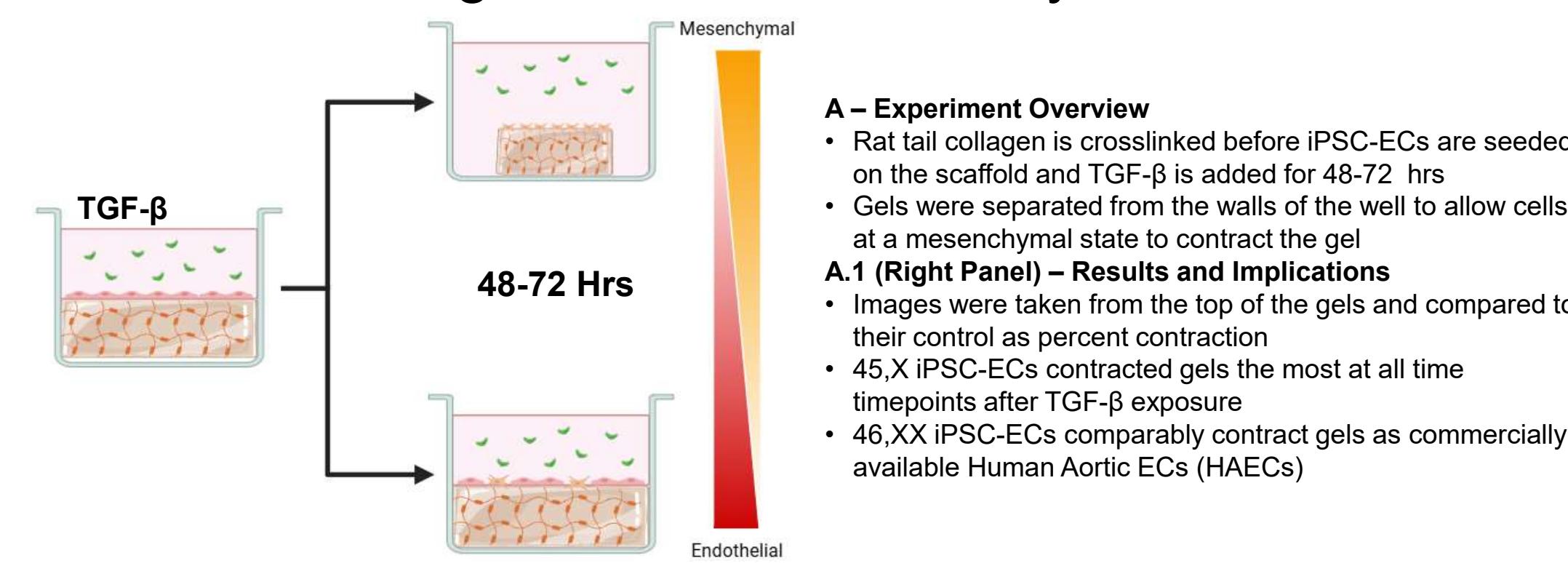
Uniform shear stress applied with a cone-plate bioreactor



Symbol	Description
$\tau$	Wall shear stress
$\mu$	Dynamic viscosity
$r$	Radius of cone
$\omega$	Angular velocity
$\alpha$	Angle between cone and plate

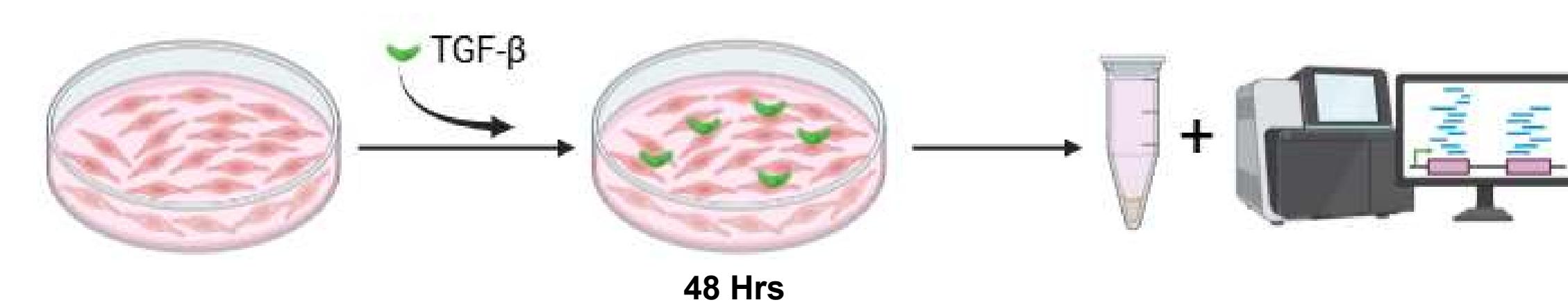
## Experiments

### Collagen Contraction Assay



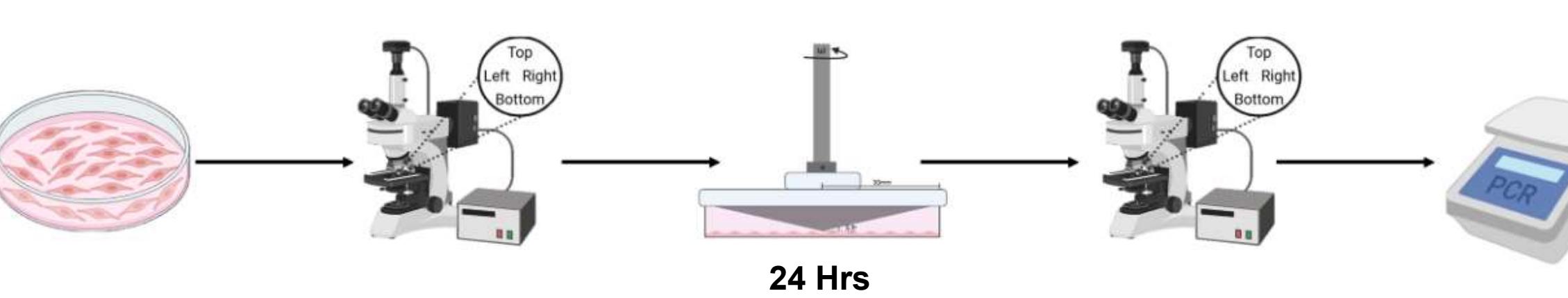
**A – Experiment Overview**  
• Rat tail collagen is crosslinked before iPSC-ECs are seeded on the scaffold and TGF-β is added for 48-72 hrs  
• Gels were separated from the walls of the well to allow cells at a mesenchymal state to contract the gel  
**A.1 (Right Panel) – Results and Implications**  
• Images were taken from the top of the gels and compared to their control as percent contraction  
• 45,X iPSC-ECs contracted gels the most at all time timepoints after TGF-β exposure  
• 46,XX iPSC-ECs comparably contract gels as commercially available Human Aortic ECs (HAECS)

### RNA-sequencing



**B – Experiment Overview**  
• iPSC-ECs (46,XX & 45,X subclones) were exposed to TGF-β for 48hrs before being sent for RNA-sequencing  
**B.1 (Right Panel) – Results and Implications**  
• Via gene ontology enrichment analysis, KDM5C, ZEB2, SNAIL and SLUG were upregulated in 45,X cells exposed to TGF-β compared to 46,XX

### Laminar Shear Stress & rt-qPCR



**C – Experiment Overview**  
• iPSC-ECs (46,XX & 45,X subclones) were exposed to uniform shear stress (0, 15 & 25 dynes/cm²) with a cone-plate bioreactor for 24Hrs  
• Brightfield images were taken before and after shear stress from 4 locations per biological replicate  
• RNA was isolated from the same samples for rt-qPCR  
**C.1 (Right Panel) – Results and Implications**  
• The average cell orientation distribution before (Blue) and after (Red) shear stress was depicted in histograms and their standard deviations as regions of their respective color  
• 46,XX iPSC-ECs increase in alignment after each shear stress condition while 45,X cells do not align with fluid flow  
• Rt-qPCR suggests EndMT activation in 45,X iPSC-ECs by shear stress while 46,XX increase transcription of endothelial CD31

## Conclusions

X-chromosome dosage on isogenic iPSC-ECs regulates:

- Epigenetic factors (KDM5C)

- TGF-β and shear stress induced EndMT (SNAIL, SLUG, ZEB2 & CD31)

- Shear stress induced cellular polarity & anti-inflammatory response

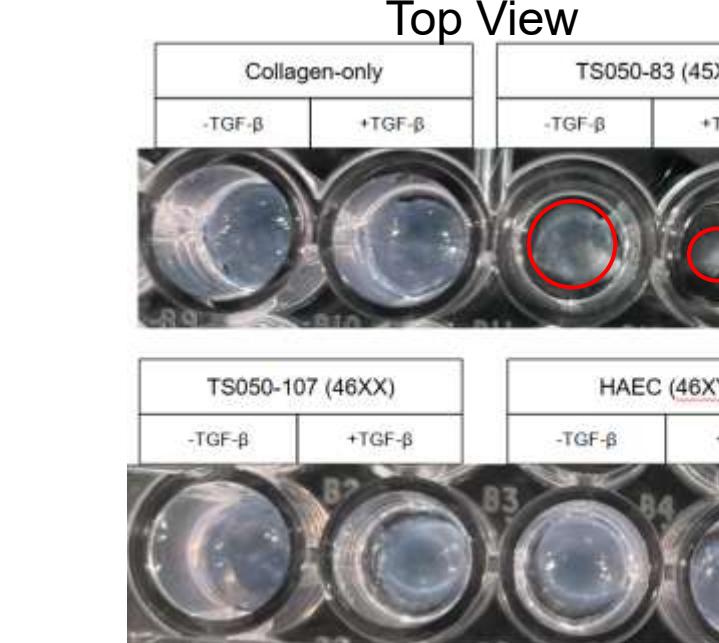
X-chromosome dosage on pathogenesis in Turner Syndrome

- Aberrant EndMT & epigenetic regulation by TGF-β in 45,X ECs may drive BAV formation

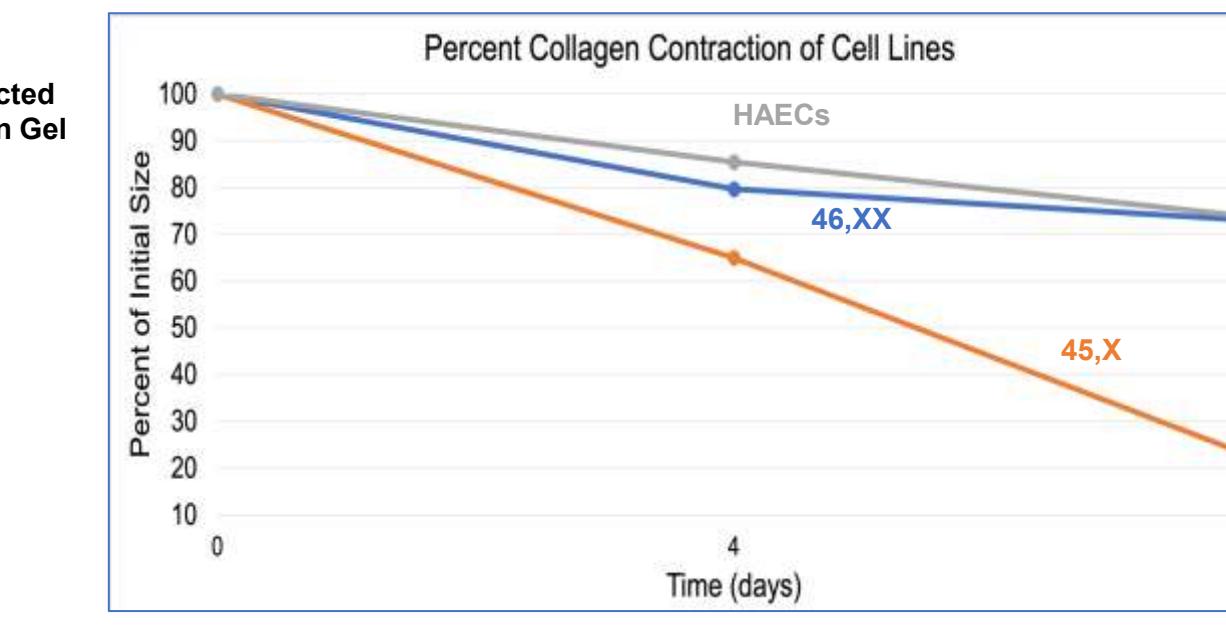
- Aberrant polarity & EndMT by shear stress in 45,X ECs may drive aneurysms/dissections

## Results

45,X iPSC-ECs readily transition to a mesenchymal phenotype with TGF-β



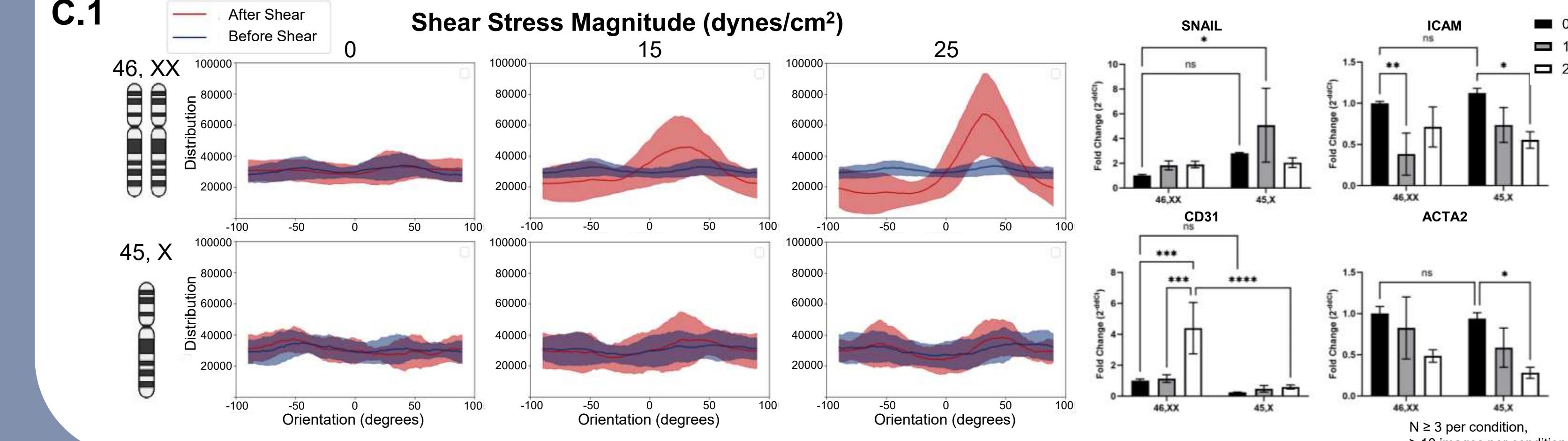
Percent Collagen Contraction of Cell Lines



TGF-β promotes phenotypic plasticity in 45,X iPSC-ECs by autosomal and X-chromosome transcriptional regulators

Expression Pattern	Gene(s)	Description	Role
Upregulation of EndMT in 45,X vs 46,XX	ZEB2, SNAIL, SLUG	Autosomal transcription factors driving mesenchymal phenotype	Vasculature Development, Cell Morphogenesis
Upregulation of epigenetic regulators in 45,X vs 46,XX	KDM5C	X-linked demethylase - increases transcriptional activity	Tube morphogenesis, Tissue Development

45,X iPSC-ECs undergo EndMT with shear stress and do not align with fluid flow



## Acknowledgements

- Funding Sources: American Heart Association Predoctoral Fellowship (24PRE1240096) and the National Science Foundation (2129122 & 2129088)

- We would like to thank all our summer undergraduate students and research assistants that helped us throughout this project: Erin Stout, Samantha Lydon, Navaneeth Pravin, Leopold Guo, Meera Krishnan, Jatin Khanna & Sama Alnassiry



References