

## Multi-modal Pipeline for Comprehensive Validation of Mitral Valve Geometry and Functional Computational Models



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#### Motivation

#### Mitral Valve (MV) Regurgitation

Most common form of valvular heart disease
Significant mortality & morbidity rates

#### **Data Acquisition**

#### **Mechanical Simulator**

In-vitro closed-loop left heart simulator<sup>[2]</sup>



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• Traditional treatment: MV *replacement* 

#### MV Repair<sup>[1]</sup>

- Better preservation of heart function & long-term survival
- Requires experienced surgeon & pre-operative planning
   Æfficient training/planning tools to optimize intervention outcome
  - Development of computational MV models
  - ➔ Models remain simplifications
- Assess clinical applicability of models by validation against comprehensive ex-vivo data





#### **Model Extraction**

#### **In-Vivo TEE**

#### High-Fidelity Model

**Ex-Vivo MicroCT** 

- Identify annulus and papillary muscle locations
  - Seed points for Random Walker segmentation
- Convert mask to mesh
  - ➔ Marching Cubes
- Segment chordae tendineae

#### State-of-the-Art Geometrical Model<sup>[3]</sup>

- Physiological point distribution model
- Based on 9 anatomical landmarks
- Marginal Space Learning framework

#### **Biomechanical Modeling**<sup>[4]</sup>



#### Novel path tracing approach



- Transverse isotropic linear elasticity
- Co-rotational finite elements method





### **Model Validation**

#### Geometric Comparison of Anatomical Models

- TEE model vs. microCT model
- MV excised from ovine heart
- Measured clinically-related parameters

#### MV Closure Computation (Biomechanics)

- Based on end-diastolic TEE model
- Pressure profile: 0 *mmHg* to 120 *mmHg*



# Anterior Leaflet Length

#### oaptation Length

#### Conclusion

#### Contributions

- Novel complete model validation pipeline
- Bridge between ex-vivo and clinical modalities
- Integration of geometric and functional models
- Controlled setup to acquire images from invasive and non-invasive modalities at almost identical conditions
- Robust algorithms to extract **reproducible** models
- First experiments on real data
  - → Utilized TEE model can accurately represent important biomarkers

#### Future Work

- Experiments on more specimen
- Evaluation of prediction power of current and future in-vivo computational frameworks

#### References

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