

AQ1 **On the Three-Dimensional Correlation**
 AQ2 **Between Myofibroblast Shape and**
 AQ3 **Contraction**
 AQ4

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*Myofibroblasts are responsible for wound healing and tissue 58
 repair across all organ systems. In periods of growth and disease, 59
 myofibroblasts can undergo a phenotypic transition characterized 60
 by an increase in extracellular matrix (ECM) deposition rate, 61
 changes in various protein expression (e.g., alpha-smooth muscle 62
 actin (α SMA)), and elevated contractility. Cell shape is known to 63
 correlate closely with stress-fiber geometry and function and is 64
 thus a critical feature of cell biophysical state. However, the rela- 65
 tionship between myofibroblast shape and contraction is complex, 66
 even as well in regards to steady-state contractile level (basal 67
 tonus). At present, the relationship between myofibroblast shape 68
 and basal tonus in three-dimensional (3D) is poorly understood. 69
 Herein, we utilize the aortic valve interstitial cells (AVICs) as a 70
 representative myofibroblast to investigate the relationship 71
 between basal tonus and overall cell shape. AVICs were embed- 72
 ded within 3D poly(ethylene glycol) (PEG) hydrogels containing 73
 degradable peptide crosslinkers, adhesive peptide sequences, and 74
 submicron fluorescent microspheres to track the local displace- 75
 ment field. We then developed a methodology to evaluate the cor- 76
 relation between overall AVIC shape and basal tonus induced 77
 contraction. We computed a volume averaged stretch tensor (U) 78
 for the volume occupied by the AVIC, which had three distinct 79
 eigenvalues ($\lambda_{1,2,3} = 1.08, 0.99, \text{ and } 0.89$), suggesting that AVIC 80
 shape is a result of anisotropic contraction. Furthermore, the 81
 direction of maximum contraction correlated closely with the lon- 82
 gest axis of a bounding ellipsoid enclosing the AVIC. As gel- 83
 embedded AVICs are known to be in a stable state by 3 days of 84
 incubation used herein, this finding suggests that the overall qui- 85
 escent AVIC shape is driven by the underlying stress-fiber direc- 86
 tional structure and potentially contraction level. 87*

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1 Introduction 88

The resident cellular population of myofibroblasts within most 89
 soft tissues maintain normal extracellular matrix (ECM) turnover 90
 by excreting, degrading, and remodeling. Myofibroblast function 91
 can be modulated by local mechanical cues as well as cytokines 92
 such as transforming growth factor beta (TGF- β) [1]. During nor- 93
 mal function, myofibroblasts are known to exhibit a baseline level 94
 of contraction (i.e., basal tonus) which establishes tensional home- 95
 ostasis between the cell and surrounding ECM [2]. When this 96
 homeostasis is perturbed, as in the case of physical insult or nor- 97
 mal growth, myofibroblasts can increase their contraction level 98
 and concomitantly change their cell shape. These processes play a 99
 crucial role in ECM remodeling and repair [1]. As a result, myofi- 100
 broblasts are often characterized by the presence of alpha-smooth 101
 muscle actin (α SMA) as well as smooth muscle myosin heavy 102
 chains which ultimately regulate myofibroblast contractile 103
 strength [3]. 104

Myofibroblast contractile properties have been studied in native 105
 tissues [4,5] and on two-dimensional (2D) substrates [6]. In native 106
 tissue, cell contraction has been shown to significantly increase 107
 bulk tissue stiffness as assessed by mechanical evaluation of 108
 excised animal tissues in which contraction is induced in the 109
 underlying cells [4,5]. However, native tissue suffers from low 110
 throughput and limited visualization of cell mechanical function, 111
 which limits the assessment of the resulting kinematic effects. 112

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113 Other studies have reported the relationship between cell shape
 114 and contractility using 2D culture surfaces [7,8], which are advan-
 115 tageous over native tissues in terms of visualization and through-
 116 put. Previously, it has been shown with 2D in vitro studies that
 117 cellular shape is correlated with a variety of intracellular mecha-
 118 nisms that regulate cell mechanical function [9–12]. For example,
 119 the eccentricity of vascular smooth muscle cell shape was found to
 120 be positively correlated with cytoplasmic and nuclear concentra-
 121 tions of calcium, which subsequently leads to higher myosin
 122 light chain kinase activity and thus increased cellular contractility
 123 [9]. However, 2D in vitro studies are limited because they cannot
 124 accurately represent the three-dimensional (3D) nature of the
 125 native tissue micro-environment. As a result, the correlation
 126 between 3D cell shape and contractile function has yet to be fully
 127 elucidated. A better understanding of this relationship could pro-
 128 vide insight into the differences in myofibroblast activation
 129 between periods of normal physiological growth and disease. Fur-
 130 thermore, a better understanding of this relationship could shed
 131 light on why the activation process can become misregulated and
 132 prolonged, subsequently leading to pathological conditions such
 133 as excessive fibrosis [2] as found in the case of Dupuytren’s con-
 134 tracture [13].

135 More recently, 3D hydrogel matrices have been used to assess
 136 the biological and mechanical response of myofibroblasts
 137 [14–16]. Three-dimensional hydrogel matrices allow for direct
 138 cell visualization, high throughput, and tunable control of the
 139 micro-environment [14–18]. Notably, recent advances have
 140 allowed for independent control of the mechanical and biochemi-
 141 cal properties of 3D hydrogels through incorporation of various
 142 types of adhesive peptide sequences and crosslinking peptides
 143 [14–16,19–21]. In previous work, synthetic poly(ethylene glycol)
 144 (PEG) hydrogels have been used to elucidate myofibroblast bio-
 145 logical and mechanical response in 3D. Specifically, it has been
 146 shown that aortic valve interstitial cells (AVICs) cultured within
 147 3D hydrogels were more similar to that of freshly isolated AVICs

148 than AVICs cultured on tissue culture polystyrene in terms of the
 149 expression levels of genes related to the cytoskeleton and contrac-
 150 tility, TGF- β signaling, and matrix remodeling [21]. This finding
 151 highlights the importance of employing realistic 3D culture envi-
 152 ronments for the assessment of myofibroblast contractile
 153 behaviors.

154 The overall objective of this work is to establish a methodology
 155 to assess the correlation between overall 3D myofibroblast shape
 156 and contractile behavior. We utilized the AVIC as an example
 157 myofibroblast and investigate their responses within a 3D PEG gel
 158 environment after 3 days of culture. By suspending fluorescent
 159 microspheres within the PEG gel, we were able to track the local
 160 gel displacements that result from AVIC basal contraction. This
 161 approach follows from previous work in the literature on 3D trac-
 162 tion force microscopy (TFM) [22,23]. The AVICs were first
 163 assessed in a homeostatic state and then within a relaxed state eli-
 164 cited by exposure to cytochalasin-D (CytoD). Through compari-
 165 son of both states, we gained insight into how AVIC basal tonus
 166 affects homeostatic AVIC shape.

2 Methods 167

2.1 Cell Isolation and Culture. 168 Porcine hearts were obtained
 169 from a local abattoir (Harvest House Farms, Johnsonville, TX) on
 170 the day of slaughter, and the aortic valves (AVs) were dissected
 171 immediately upon arrival to the lab. AVICs were extracted from
 172 the AV using previously published methods [24]. In brief, the AV
 173 leaflets were washed in Earle’s balanced salt solution (Thermo
 174 Fisher Scientific) before being subjected to a digestion solution
 175 (0.75 mg/ml of collagenase in Earle’s balanced salt solution) for
 176 half an hour to remove the valve endothelial cells lining the out-
 177 side of the leaflets. A second round of digestion was implemented,
 178 and the leaflets were incubated within a fresh collagenase solution
 179 for 1 h before vortexing the solution to dislodge the AVICs. After-
 180 ward, the solution was filtered using a 100 μ m cell strainer, and

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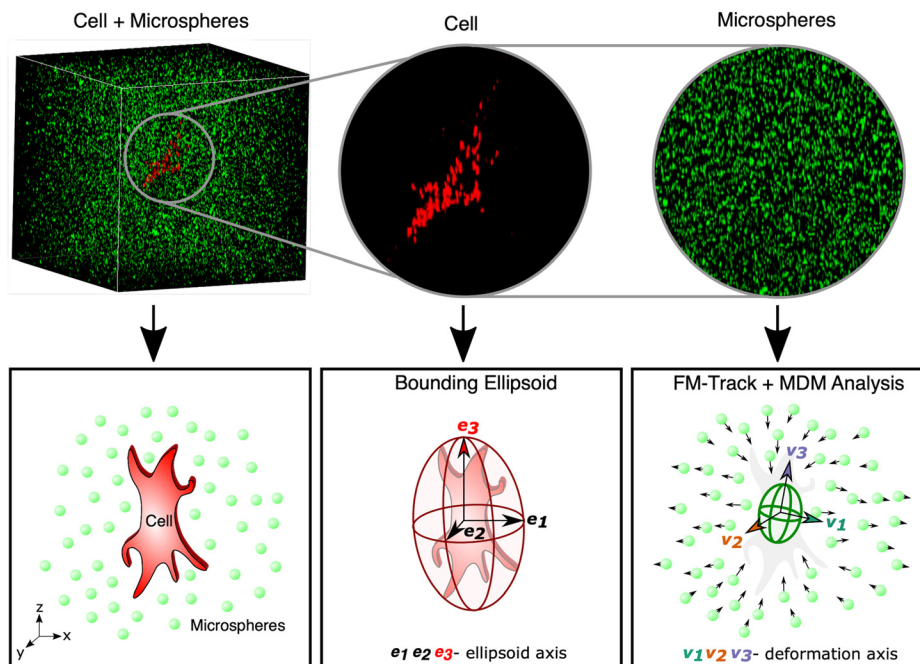


Fig. 1 Description of the work flow for each of the 20 cells analyzed. Top row: two 3D images of a single cell surrounded by 0.5 μ m fluorescent microspheres were obtained via confocal microscopy before and after treatment with CytoD (only one image is shown here for brevity). Bottom left: the 3D cell geometry and the position of each fluorescent microsphere were segmented from 3D images. Bottom center: a bounding ellipsoid is computed for each cell geometry. Bottom right: the displacements of the fluorescent microspheres between the two image sets are computed using our recently developed software “FM-TRACK” [25].

181 the filtered solution was centrifuged. The cell pellet was resus-
 182 pended and plated in standard growth media (Dulbecco's modified
 183 eagle medium, 10% fetal bovine serum, 2% pen-strep, 0.4% fun-
 184 gizone, Thermo Fisher Scientific). Only AVICs with passage
 185 numbers 2–4 were used for experimentation.

186 2.2 Three-Dimensional Traction Force Microscopy

187 *Valve Interstitial Cell Imbedded Gel Specimen Preparation.*
 188 Porcine AVICs were stained with CellBrite™ Red (Biotum) to
 189 fluorescently label the cell membranes. The 3D TFM samples
 190 were then fabricated by seeding stained AVICs at a density of
 191 500,000 cells/ml within a synthetic PEG hydrogel containing
 192 eight-arm 40kDa norbornene functionalized PEG molecules,
 193 matrix metalloproteinase degradable cross linking peptides,
 194 cysteine–arginine–glycine–aspartate–serine adhesive peptide
 195 sequences, and 0.5 μm yellow–green fluorescent microspheres
 196 (Polysciences) suspended at a density of 3×10^9 beads/ml. During
 197 the mixing process, the fluorescent microspheres were added first
 198 followed by the hydrogel precursor solutions and then the stained
 199 AVICs. Adding the components in this order ensured that the fluo-
 200 rescent microspheres did not clump together during the fabrication
 201 process. The 3D TFM samples were then polymerized within well
 202 dishes with 12 mm glass well inserts to hold the samples in place
 203 during the imaging process (Fisher Scientific). The samples
 204 were incubated for 72 h in standard growth media before
 205 experimentation.

206 *Imaging.* The growth media surrounding the 3D TFM sample
 207 was replaced with Tyrode's salt solution (TSS) and incubated for
 208 40 min before a z-stack image set containing a single AVIC and
 209 surrounding fluorescent microspheres in the field of view

(FOV) was obtained with dimensions $150 \times 150 \times 140 \mu\text{m}$ 210
 (512 \times 512 \times 176 pixels) at a z-step of 0.8 μm using a Zeiss LSM 211
 710 inverted confocal microscope with a 63 \times , 1.2 numerical aper- 212
 ture water-immersion objective lens (Fig. 1). Then, a stock solu- 213
 tion of CytoD was added to achieve a working concentration of 214
 4 μM to relax the basal tonus of the AVIC through actin depoly- 215
 merization. To allow for sufficient time for the actin depolymeriz- 216
 ing effects of CytoD to be observed while minimizing 217
 downstream effects on gene expression, the samples were allowed 218
 to incubate for 40 min before an additional z-stack image set of 219
 the same FOV was obtained. A total of 20 experimental trials 220
 were analyzed in this study. 221

222 *Tracking of Fiducial Markers.* For all experimental trials, 222
 microsphere displacements were tracked from the CytoD treated 223
 state to the TSS state (opposite of how the experiments were con- 224
 ducted) to assess the effect of AVIC basal tonus on hydrogel dis- 225
 placements. Each set of two 3D image stacks was analyzed with 226
 our recently developed open source software FM-TRACK [25]. The 227
 workflow of the software is as follows. First, it segments each 228
 image and determines the 3D position of the centroid of each fluo- 229
 rescent microsphere. Next, it constructs feature vectors extending 230
 from the centroid of each microsphere to the centroid of each of 231
 the microsphere's five closest neighbors. Then, markers in the first 232
 configuration are matched to markers in the second configuration 233
 by comparing feature vectors within a spatially defined neighbor- 234
 hood of 15 candidate matches. The difference between the feature 235
 vectors of the original marker and each of the 15 candidate 236
 markers is computed and recorded as a "test score." Test scores 237
 are computed for all permutations of the feature vectors, and the 238
 particle with the overall lowest test score is assigned as a match. 239
 If a matching conflict occurs (e.g., two markers are paired with 240

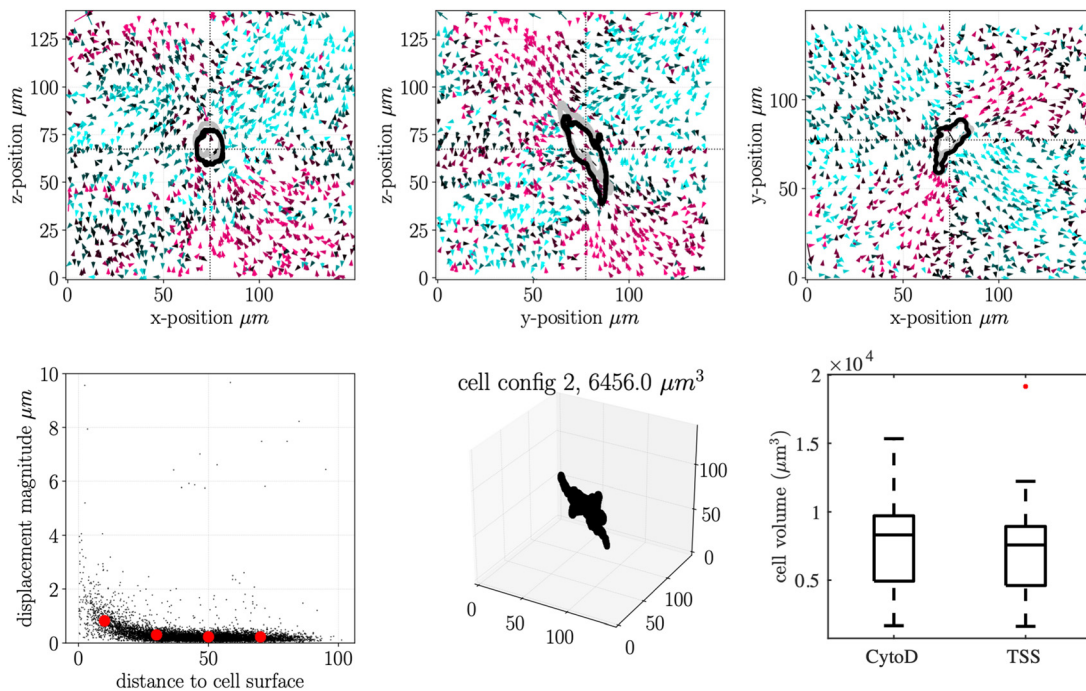


Fig. 2 Representative data sheet from analyzing the experimental data with FM-TRACK [25]. Top row: plane slices depicting microsphere displacements in the y -, x -, and z -planes with magenta denoting microsphere displacement vectors oriented toward the AVIC surface and cyan denoting those away from the AVIC surface. The gray and black cross-sectional outlines show the AVIC in its first configuration and second configuration, respectively. Bottom left: microsphere displacement magnitudes decrease with respect to increasing distance from the AVIC surface. Both the microsphere displacement magnitude and distance to cell surface are computed in 3D. Bottom center: segmented AVIC geometry and volume in the second configuration (TSS). Bottom right: AVIC volume within the CytoD and TSS treated states. AVIC volume is not statistically different between each state as determined by a paired t -test (p -value = 0.23). The middle line of the box plot denotes the median. Red markers denote outliers that are greater than 1.5 times the interquartile range above the third quartile. A total of $n = 20$ cells were tested.

241 the same marker), the marker with the lowest score is assigned,
 242 and the other marker(s) are paired to their next best match. The
 243 tracking software is programed to only accept bidirectional
 244 matches that hold from both the first configuration to the second
 245 configuration and from the second configuration to the first confi-
 246 guration. Marker pairs that did not produce the same result from
 247 both matching directions are not considered for further analysis.
 248 The tracking algorithm outputs the initial coordinates and dis-
 249 placement vectors for each successfully paired marker (Fig. 2).
 250 Furthermore, the software provides a displacement interpolation
 251 between each microsphere location via Gaussian process regres-
 252 sion [26].

253 *Generating Aortic Valve Interstitial Cell Surface Meshes.* A tri-
 254 angulated surface mesh for each AVIC was produced by the FM-
 255 TRACK software. This was accomplished by thresholding the 3D
 256 experimental image stack containing the AVIC, identifying the
 257 largest connected 3D volume of voxels which was assumed to be
 258 the AVIC geometry, and using the Lewiner marching cubes algo-
 259 rithm to construct a surface mesh [27]. The FM-TRACK software
 260 also computed the volume of the AVICs by multiplying the num-
 261 ber of voxels making up the AVIC geometry by the voxel spatial
 262 dimensions. This was done for images acquired after the TSS and
 263 CytoD treatment.

264 2.3 Aortic Valve Interstitial Cell Shape Characterization.

265 Imbedded myofibroblasts generally have very complex geometries
 266 within 3D environments, often containing multiple protrusions
 267 that vary in overall size and length [15]. For this study, we are
 268 interested in overall cell shape rather than the granular details. To
 269 this end, approximate descriptors of AVIC shape were obtained
 270 through computing the minimum volume enclosing ellipsoids
 271 (bounding ellipsoids) of each AVIC geometry. The resulting
 272 length of the axes and the orientation of the ellipsoid are used to
 273 characterize the AVIC shape. Minimum volume enclosing ellip-
 274 soids were computed using a modified version of a script [28]
 275 based on Khachiyan's algorithm (Algorithm 1) [29]. The algo-
 276 rithm finds the minimum enclosing ellipsoid of a data point cloud
 277 stored in matrix P by solving the following optimization problem:

$$\begin{aligned} \text{min. } & \Phi = \log(\det(A)) \\ \text{subject to } & (P_i - c)^T \times A \times (P_i - c) \leq 1, \forall i \in P \end{aligned} \quad (1)$$

279 where A is a 3×3 matrix of the ellipse equation in center form
 280 such that

$$(x - c)^T \times A \times (x - c) = 1 \quad (2)$$

282 and c is the center of the ellipsoid. The optimization problem was
 283 solved using Algorithm 1 to obtain A and c . Then, the radii and
 284 orientation of the ellipsoid are obtained by singular value decom-
 285 position of A into matrices U , S , and V such that

$$A = U \times S \times V^T \quad (3)$$

286 The radii are simply the inverse of the square root of the diagonal
 287 elements of matrix S , and the orientation of the bounding ellipsoid
 288 is obtained from the rotation matrix V . From this method, the
 289 length and orientation of three orthogonal axes that describe the
 290 bounding ellipsoid are obtained. The orientation of the major axis
 291 (e_3) is used to represent the approximate orientation of the AVIC
 292 shape.

294 **2.4 Describing Aortic Valve Interstitial Cell Average**
 295 **Deformation.** In order to determine the inter-relationship
 296 between AVIC shape and overall contraction pattern, we utilized
 297 our ability to interpolate TFM microsphere displacements using
 298 FM-TRACK [25]. Gaussian process regression [26] was used to inter-
 299 polate the local hydrogel displacement field to compute the

Algorithm 1 Bounding ellipsoid algorithm [28] based on Kha-
 chiyan's algorithm [29] as implemented in MATLAB.

Result: bounding ellipsoid
 $P = x, y, z$ point cloud of cell geometry;
 $n =$ number of points in P ;

$$Q = \begin{bmatrix} x_1 & x_2 & x_3 & \dots & x_n \\ y_1 & y_2 & y_3 & \dots & y_n \\ z_1 & z_2 & z_3 & \dots & z_n \\ 0 & 0 & 0 & \dots & 0 \end{bmatrix};$$

error = 1;
 tolerance = 0.05;
 $\mathbf{u} = (1/n) \times \text{ones}(n, 1)$;
 $\mathbb{R}^d, d = 3$;
while error > tolerance **do**
 $\mathbf{X} = Q \times \text{diag}(\mathbf{u}) \times Q^T$;
 $\mathbf{m} = \text{diag}(Q^T \times \text{inv}(\mathbf{X}) \times Q)$;
 $[\text{maximum}, \text{index}] = \text{max}(\mathbf{m})$;
 $\text{step_size} = (\text{maximum} - d - 1) / ((d + 1) \times (\text{maximum} - 1))$;
 $\text{new_u} = (1 - \text{step_size}) \times \mathbf{u}$;
 $\text{new_u}(\text{index}) = \text{new_u}(\text{index}) + \text{step_size}$;
 $\text{error} = \text{norm}(\text{new_u} - \mathbf{u})$;
 $\mathbf{u} = \text{new_u}$;
end
 $\mathbf{U} = \text{diag}(\mathbf{u})$;
 $\mathbf{A} = \frac{1}{d} \times (\mathbf{P} \times \mathbf{U} \times \mathbf{P}^T - (\mathbf{P} \times \mathbf{u}) \times (\mathbf{P} \times \mathbf{u})^T)^{-1}$;
 $\mathbf{c} = \mathbf{P} \times \mathbf{u}$

displacements \mathbf{u} at the center of each triangular facet of the AVIC 300
 surface mesh. Next, mean deformation metrics (MDMs) were 301
 computed to quantify the change that the AVIC shape undergoes 302
 due to AVIC basal tonus [30]. This was done by first computing a 303
 mean displacement gradient $\langle \nabla \mathbf{u} \rangle$ using 304

$$\langle \nabla \mathbf{u} \rangle := \frac{1}{\text{vol}(V_{\text{AVIC}})} \int_{\partial V_{\text{AVIC}}} \mathbf{u} \otimes \mathbf{n} dA \quad (4)$$

where $\text{vol}(V_{\text{AVIC}})$ represents the volume of the AVIC geometry in 306
 the initial configuration (V_{AVIC}), ∂V_{AVIC} is the boundary of V_{AVIC} , 307
 dA is an infinitesimal element on the boundary ∂V_{AVIC} , and \mathbf{n} is 308
 the unit surface normal of dA . This approach provides a simple 309
 and convenient method to compute a volume average displace- 310
 ment gradient from cell surface displacements only. From this, the 311
 mean deformation gradient tensor $\langle \mathbf{F} \rangle$ is simply computed as 312
 follows: 313

$$\langle \mathbf{F} \rangle = \mathbf{I} + \langle \nabla \mathbf{u} \rangle \quad (5)$$

From polar decomposition theorem, $\langle \mathbf{F} \rangle = \langle \mathbf{R} \rangle \langle \mathbf{U} \rangle$, the mean 314
 stretch $\langle \mathbf{U} \rangle$ and rotation $\langle \mathbf{R} \rangle$ tensors are obtained. The eigenvalues 316
 $(\langle \lambda_i \rangle, i = 1, 2, 3)$ and eigenvectors (v_1, v_2, v_3) of $\langle \mathbf{U} \rangle$ are computed 317
 and represent the magnitudes and directions of principal AVIC 318
 shape deformation. Because the analysis is conducted from the 319
 CytoD treated state to the TSS state, maximum contraction is cap- 320
 tured by (λ_3, v_3) , and maximum expansion is captured by (λ_1, v_1) . 321
 We noted that $\langle \mathbf{R} \rangle \cong \mathbf{I}$, with a mean rotation angle along any axis 322
 of less than 2 deg, so that here $\langle \mathbf{U} \rangle \cong \langle \mathbf{F} \rangle$. 323

2.5 Investigating the Correlation Between Aortic Valve 324
Interstitial Cell Eccentricity and Magnitude of Aortic Valve 325
Interstitial Cell Contraction. It has been previously demon- 326
 strated with 2D in vitro studies that increased cell eccentricity 327
 leads to increased cell contractility [7–9,31,32]. Here, this obser- 328
 vation is tested in 3D. The eccentricity of the AVIC shapes was 329
 quantified by computing the aspect ratios of the bounding ellip- 330
 soids (length of long axis/average of two minor axes), and the 331
 magnitude of AVIC contraction was represented by λ_3 . The 332

333 correlation of these two variables is assessed using Pearson's correlation coefficient.
 334

335 **2.6 Statistical Methods**

336 *Data Exclusion Criteria.* Experimental trials were excluded
 337 from analysis based on two criteria. First, if the AVIC surface was
 338 less than $10\ \mu\text{m}$ from any boundary of the FOV within either the
 339 TSS or CytoD treated states, the trial was excluded from further
 340 analysis under the assumption that the experiment failed to capture
 341 the full displacement field produced by the AVIC. Then, the
 342 effect size of the trial was assessed. The mean of the displacement
 343 magnitudes within $10\ \mu\text{m}$ of the cell surface was used as a metric
 344 to represent the effect size of the trial. This value was compared
 345 to synthetic data obtained by randomly shuffling the displacement
 346 magnitude versus distance to cell surface data (e.g., Fig. 2, bottom
 347 left). This was done by assigning each displacement magnitude
 348 value to a random distance from cell surface value. Next, the average
 349 and standard deviation of the synthetic displacement magnitudes
 350 within $10\ \mu\text{m}$ of the cell surface were computed. If the
 351 experimental average was within three standard deviations from
 352 the synthetic mean, the effect size was considered insignificant,
 353 and the experimental trial was excluded from analysis. A total of
 354 20 experimental trials were deemed acceptable and were analyzed
 355 in this study.

356 *Testing the Relationship Between Aortic Valve Interstitial Cell*
 357 *Orientation and Direction of Maximum Contraction.* The dot
 358 product between the unit vectors describing the longest axis of the
 359 bounding ellipsoid (e_3) and the direction of maximum contraction
 360 (v_3) was computed to assess if a correlation existed between
 361 AVIC orientation and the direction of maximum contraction. A
 362 dot product of $e_3 \cdot v_3 = 1$ indicates parallel unit vectors and complete
 363 agreement between the direction of AVIC contraction and
 364 AVIC shape orientation, whereas $e_3 \cdot v_3 = 0$ indicates orthogonal
 365 unit vectors.

Monte Carlo Simulation. Standard statistical techniques often
 366 assume that the random variables being compared are from normal
 367 distributions. Here, we do not assume the distribution of the
 368 e_3 and v_3 vectors. Instead, a nonparametric statistical method was
 369 used to assess the statistical significance of the correlation
 370 between e_3 and v_3 . To this end, a Monte Carlo simulation of the
 371 random scenario was performed. Specifically, random unit vectors
 372 were generated for the direction of AVIC orientation (\mathbf{E}), and
 373 their dot products with the experimentally observed directions of
 374 maximum contraction (\mathbf{V}) were computed (Algorithm 2). This
 375 process was implemented for each of the 20 experimental trials,
 376 and the mean dot product was reported. For convergence, 10,000
 377 Monte Carlo simulations were ran, and the results are illustrated
 378 as the distribution of their means in a histogram (Fig. 3(b)).
 379 Outliers of the random distribution were defined as values greater
 380 than three standard deviations away from the mean of the
 381 random distribution. If the experimental observations were
 382

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Algorithm 2 Monte Carlo simulation algorithm used to obtain a distribution representing the hypothetical case where the relationship between AVIC orientation (e_3) and the direction of maximum contraction (v_3) is random.

```

Result: Monte Carlo simulation
n = 20;
V = [
    v_{3,x}^1  v_{3,y}^1  v_{3,z}^1
    v_{3,x}^2  v_{3,y}^2  v_{3,z}^2
    ...
    v_{3,x}^n  v_{3,y}^n  v_{3,z}^n
];
for i = 1:10,000 do
    E = rand(n, 3) (generates n random unit vectors);
    g = dot(E, V, 2) (row-wise dot product);
    avg(i) = mean(g)
end
    
```

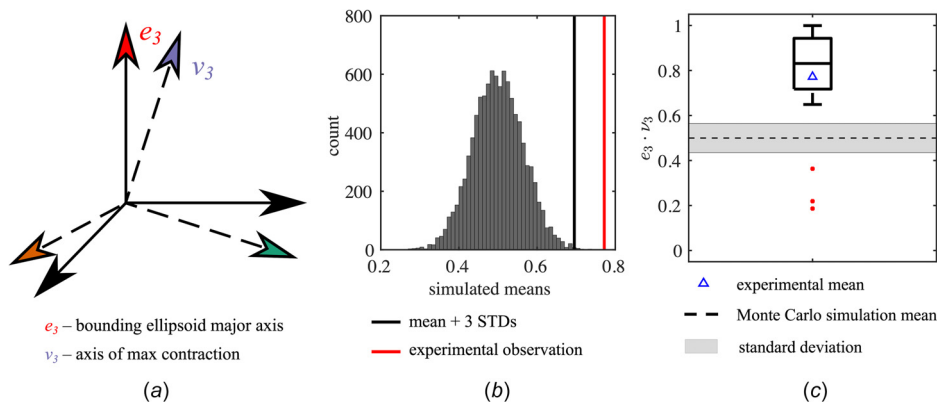


Fig. 3 Comparison between the direction of AVIC shape orientation from ellipsoidal approximation and the direction of maximum contraction computed by MDM. (a) Schematic illustrating the unit vector of the major axis of the best-fit ellipsoid (e_3) and the direction of maximum contraction (v_3). (b) The distribution of means for 10,000 Monte Carlo simulations which represent a random correlation between e_3 and v_3 . Each Monte Carlo simulation consisted of computing the dot products between 20 randomly oriented unit vectors and the experimentally observed values of v_3 . The mean of the experimental observation (red line) was more than three standard deviations above the mean of the Monte Carlo simulation (black line) which suggests that the correlation between e_3 and v_3 is not random. (c) Box-and-whiskers plot of the dot products between e_3 and v_3 from the experimental data is shown along with the experimental mean (0.77). The middle line of the box plot denotes the median, whereas the triangular marker denotes the mean. Red markers denote outliers that are less than 1.5 times the interquartile range below the first quartile. In addition, the mean and standard deviation of the Monte Carlo simulation results (dashed line and gray area, respectively) are superimposed. A total of $n = 20$ cells were tested.

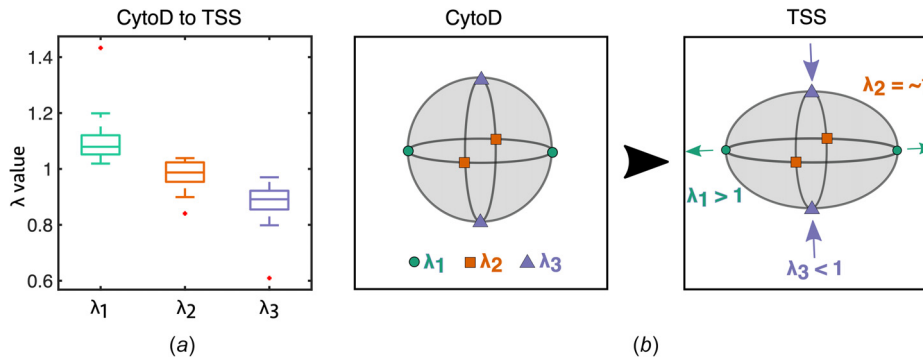


Fig. 4 Kinematic summary of changes in AVIC shape due to basal tonus. (a) Box-and-whiskers plot of the eigenvalues of the mean stretch tensor $\langle \mathbf{U} \rangle$. The middle line of the box plot denotes the median. Red markers denote outliers defined as values either greater than 1.5 times the interquartile range above the third quartile or less than 1.5 times the interquartile range below the first quartile. A total of $n = 20$ cells were tested. (b) A schematic demonstrating typical deformations that AVICs undergo from the CytoD state to the TSS state characterized by contraction in one direction ($\lambda_3 < 1$), expansion in an orthogonal direction ($\lambda_1 > 1$), and virtually no deformation in the region orthogonal to the previous two (λ_2).

383 substantially far from the mean of the random distribution, the
 384 null hypothesis that the relationship between the orientation of
 385 AVICs and the direction of maximum contraction is random can
 386 be rejected.

387 3 Results

388 **3.1 Relaxing Aortic Valve Interstitial Cell Basal Tonus in**
 389 **a Three-Dimensional Synthetic Hydrogel Environment**
 390 **Produces a Local Gel Displacement Field.** To visualize the
 391 effects of cessation of stress-fiber contraction via CytoD, planar
 392 cross sections of the FOV taken through the center of each AVIC
 393 were performed. Results revealed that, qualitatively, the hydrogel
 394 displacement patterns were highly directional, with regions of
 395 contraction toward the AVIC surface and expansion away from
 396 the AVIC surface (Fig. 2, top row). In addition, we observed that
 397 the magnitude of bead displacement decreased nonlinearly with
 398 respect to increasing distance from the AVIC surface (Fig. 2, bot-
 399 tom left). Both the microsphere displacement magnitude and dis-
 400 tance to cell surface are computed in 3D. We also noted that the
 401 AVIC volumes did not change significantly between the two con-
 402 figurations (Fig. 2, bottom right).

403 **3.2 Basal Tonus Contribution to Aortic Valve Interstitial**
 404 **Cell Shape.** Following the methods described in Sec. 2.4, the
 405 average deformation of the volume occupied by the AVIC was
 406 determined using MDMs [30] to investigate the kinematic effects
 407 of AVIC basal tonus on the resulting AVIC shape. The MDMs for
 408 each of the 20 AVICs were computed and summarized in a box-
 409 and-whiskers plot (Fig. 4(a)). The median MDM eigenvalues
 410 were 1.08, 0.99, and 0.89 for λ_1 , λ_2 , and λ_3 , respectively. The
 411 results of this analysis revealed that AVIC basal tonus causes a
 412 consistent pattern of AVIC shape changes characterized by con-
 413 traction in one direction (λ_3), expansion in an orthogonal direction
 414 (λ_1), and virtually no deformation in the direction orthogonal to
 415 the previous two (λ_2) (Fig. 4(a)). A schematic is provided which
 416 demonstrates the effects of the deformation patterns on a refer-
 417 ence spherical geometry (Fig. 4(b)). The characteristic deforma-
 418 tion pattern will cause a sphere to morph to an ellipsoid.

419 **3.3 Aortic Valve Interstitial Cell Orientation and**
 420 **Direction of Maximum Contraction Are Correlated.** Following
 421 the methods outlined in Sec. 2.6, the correlation between AVIC
 422 orientation and the direction of maximum contraction was
 423 assessed by computing the dot product $e_3 \cdot v_3$, where e_3 is the
 424 direction of the longest axis of the AVIC bounding ellipsoid, and
 425 v_3 is the direction of maximum gel contraction (Fig. 3(a)). This

allows for a quantitative measure of the correlation between
 AVIC shape and AVIC basal tonus. The dot product $e_3 \cdot v_3$ was
 computed for all 20 AVICs and was found to be greater than three
 standard deviations above the mean of the Monte Carlo simula-
 tions which represent the random scenario (Fig. 3(b)). This indi-
 cates that the null hypothesis that the relationship between the
 orientation of AVICs and the direction of maximum contraction is
 random can be rejected. The distribution of the experimentally
 observed $e_3 \cdot v_3$ results is also summarized in a box-and-whiskers
 plot (Fig. 3(c)).

436 **3.4 Aortic Valve Interstitial Cell Eccentricity and**
 437 **Magnitude of Contraction Are Not Correlated in Three-**
 438 **Dimensional.** To assess the correlation between 3D AVIC eccen-
 439 tricity and contraction magnitude, we plotted the third eigenvalue
 440 (λ_3) of the mean stretch tensor $\langle \mathbf{U} \rangle$ against the aspect ratio of
 441 the bounding ellipsoid (length of long axis/average of two minor
 442 axes) for all 20 cells (Fig. 5) and computed Pearson's correlation
 443 coefficient. From this analysis, a Pearson's correlation coefficient
 444 of $r = -0.33$ was observed. However, the correlation was found
 445 to not be statistically significant (p -value = 0.16). Therefore, the
 446 eccentricity of AVIC shape does not influence the amount of con-
 447 traction brought about from the AVIC in 3D.

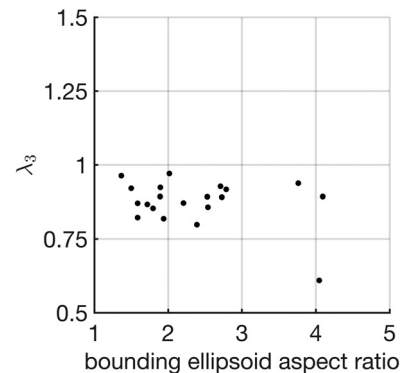


Fig. 5 Assessing the correlation between AVIC eccentricity and AVIC contraction magnitude. Plot of the third eigenvalue (λ_3) of the mean stretch tensor $\langle \mathbf{U} \rangle$ against the aspect ratio of the bounding ellipsoid showed no statistically significant correlation as assessed by Pearson's correlation coefficient ($r = -0.33$ and p -value = 0.16). This suggests that the magnitude of AVIC contraction is not meaningfully correlated with AVIC eccentricity in 3D.

448 **4 Discussion**

449 **4.1 Initial Analysis of the Relationship Between Aortic**
 450 **Valve Interstitial Cell Shape and Basal Tonus.** The average
 451 experimentally observed relationship between longest AVIC axis
 452 and maximum contraction (measured by the dot product $e_3 \cdot v_3$) is
 453 greater than what would be predicted if there was no correlation
 454 between AVIC orientation and direction of maximum contraction
 455 (Fig. 3(b)). This result suggests that the direction of AVIC orienta-
 456 tion and the direction of maximum contraction are correlated.
 457 This is consistent with observations in 2D studies that demon-
 458 strated that an aligned AVIC population produced substantial
 459 bending in a thin-film cantilever along the direction of the longest
 460 AVIC shape axis [8]. Broadly speaking, these results indicate that
 461 the direction of the longest AVIC shape axis is correlated with the
 462 direction of AVIC contraction in both 2D and 3D environments.
 463 This observation is possibly a result of the structure–function rela-
 464 tionship in AVICs. Specifically, cell shape in general is related to
 465 the orientation of subcellular stress-fibers [33,34]. In AVICs, fila-
 466 mentous actin (F-actin) stress-fiber orientation has been previ-
 467 ously shown to be correlated with AVIC shape [7].

468 Previous work from our group investigated the roles of α SMA
 469 and F-actin stress-fibers in AVIC intrinsic stiffness and contractile
 470 force generation [35,36]. Our results indicated that α SMA stress-
 471 fibers were the dominant contributor toward stress-fiber force gen-
 472 eration, especially among activated AVICs that were treated with
 473 TGF- β 1 and potassium chloride to elicit a myofibroblast pheno-
 474 type. However, F-actin and α SMA stress-fibers were both found
 475 to be crucial in modulating AVIC contractile forces. Based on
 476 these observations, we hypothesize that F-actin and α SMA stress-
 477 fibers play large roles in modulating both the contractile behavior
 478 and the homeostatic cell shape of AVICs in 3D PEG hydrogel
 479 environments.

480 **4.2 Insight Into the Differences of Aortic Valve Interstitial**
 481 **Cell Biomechanical Behaviors Within Two-Dimensional and**
 482 **Three-Dimensional Environments.** In 2D environments, it has
 483 been demonstrated that AVIC shape is correlated with the direc-
 484 tion and magnitude of contraction for both single-cells [7] and
 485 cell populations [8]. Specifically, it has been shown that the basal
 486 tonus of elongated AVICs produced greater levels of stress than
 487 the basal tonus of less elongated AVICs [7]. This finding suggests
 488 that AVIC shape potentially influences internal stress-fiber func-
 489 tion and thus the resulting contractile behaviors. However, our
 490 investigations into the correlation between AVIC shape and con-
 491 tractile behaviors within a 3D isotropic PEG gel showed that
 492 AVIC eccentricity, as represented by the aspect ratio of the
 493 bounding ellipsoid, was not meaningfully correlated with the
 494 magnitude of contraction (λ_3) due to basal tonus (Fig. 5). This
 495 finding makes it clear that further investigation is required into the
 496 correlation between AVIC shape and basal tonus in a 3D environ-
 497 ment. Based on the previous work in the literature and our initial
 498 findings, it is likely that this phenomenon depends upon the
 499 dimension of the culture system (constrained 2D versus uncon-
 500 strained 2D versus 3D). Previous work by Mabry et al. demon-
 501 strated that AVIC gene expression varied substantially between
 502 2D and 3D cultures [21]. Most notably, AVICs seeded within 3D
 503 hydrogel matrices showed more similarities to freshly isolated
 504 AVICs than AVICs seeded on 2D substrates in terms of gene
 505 expression levels for cytoskeletal organization and focal adhe-
 506 sions. This finding suggests that 3D culture matrices elicit AVIC
 507 contractile and adhesion behaviors that are most similar to the
 508 native tissue environment. Furthermore, 2D versus 3D cell
 509 mechanical function may be influenced by the spatial distribution
 510 of adhesion sites [37]. For example, in 2D cultures, cell adhesion
 511 is constrained to a horizontal plane, whereas in 3D, cells can form
 512 adhesion sites in all three dimensions. Moreover, differences
 513 between cell behavior in 2D and 3D may arise because cell shape
 514 influences stress-fiber orientation and function differently in 3D
 515 gel environments. Clearly, further investigation is needed to

thoroughly elucidate the dimension-dependent mechanical
 response of myofibroblast cells. 516
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4.3 Limitations and Future Directions. In this study, we
 computed bounding ellipsoids of the AVIC membrane surfaces to
 obtain approximate AVIC shape descriptors. Although the bound-
 ing ellipsoids were useful, they cannot account for detailed cellu-
 lar shape/structures that may play a large role in AVIC contractile
 behaviors [33,34]. For example, bounding ellipsoids do not record
 the number, size, and curvature of pseudopodia [38]. Therefore,
 the bounding ellipsoid serves as an informative but ultimately first
 approximation. Regarding underlying relations to specific stress-
 fiber structures, direct observations of the relationship between
 3D cell shape and stress-fiber orientation are limited due to the
 feature size of individual stress-fibers (F-actin fibers are approxi-
 mately 7 nm in diameter [39]) falling well below the axial resolu-
 tion of light microscopy. We thus only report on the observed
 correlations between AVIC shape and the direction of greatest
 contraction. Ongoing work will focus on addressing the current
 limitations by employing more advanced methods to quantify cell
 shape and the kinematic effects of cell contraction. 518
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4.4 Summary. Our findings indicated that a clear correlation
 existed between AVIC overall shape and orientation and the direc-
 tion of maximum basal tonus-derived contraction. Specifically,
 AVIC basal tonus produced consistent changes in AVIC shape
 characterized by contraction in one primary direction, expansion
 in an orthogonal direction, and virtually no change in the third
 orthogonal direction. As gel-embedded AVICs are known to be in
 a relatively stable state by 3 days of incubation used herein
 [16,20], this finding suggested that the overall quiescent AVIC
 shape is driven by the underlying homeostatic stress-fiber direc-
 tional structure and contraction level. These findings may also
 hold for myofibroblasts as a whole. 536
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Nomenclature 560

AV = aortic valve	561
AVICs = aortic valve interstitial cells	562
CytoD = cytochalasin-D	563
ECM = extracellular matrix	564
F-actin = filamentous actin	565
FOV = field of view	566
MDMs = mean deformation metrics [30]	567
PEG = poly(ethylene glycol)	568
TFM = traction force microscopy	569
TGF- β = transforming growth factor beta	570
TSS = Tyrode's salt solution	571
α SMA = alpha-smooth muscle actin	572

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