

# SPATIAL SHAPE ANALYSIS OF CELL NUCLEI\*

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**Abstract.** In a histopathology image, to assess if the shapes of tumor cell nuclei influence tumor infiltration in surrounding tissues it is important to first quantify and visualize variability in the shape of cell nuclei whilst accounting for their spatial correlations. To avoid loss of shape information when using summaries, we represent the nucleus boundary as a planar curve, and develop a geometric framework to compute local spatially-informed summaries of shapes of planar curves. The framework is based on the definition of a novel shape trace-variogram that captures dependence between shapes of curves. The variogram is used to compute a local spatially-weighted shape average and covariance, using which, shape variation is quantified via principal component analysis applied to the weighted covariance operator. Primary directions of spatial shape variation are subsequently visualized as vector fields (and their pointwise magnitudes) on the spatially-weighted average shape. Our approach serves as a practical exploratory tool for clinicians who routinely assess morphological and spatial heterogeneity of cell nuclei, and also represents the first step toward probabilistic modeling of spatially varying cell shapes. We demonstrate utility of the framework on synthetic data, and on data pertaining to cell nuclei from whole slide images of breast cancer tissues.

**Relevance to Life Sciences.** Pathologists routinely examine tumor regions in histopathology images to assess severity and progression of cancer. Nuclear shape heterogeneity plays a key role in this process and provides valuable information that guides clinical treatment decisions. To aid in such assessments, there is need for tools that enable statistical quantification, summarization and visualization of cell nucleus shape variation. While such tools exist for independent shape data, they are not applicable in the current context due to the inherent spatial correlation among cell nuclei. Thus, we define local spatially-informed shape summaries which, when computed based on cell nuclei within histopathology image regions, reveal their shape heterogeneity. The proposed framework is applied to whole slide images of breast cancer tissues.

**Mathematical Content.** Cell nucleus boundaries are represented using planar closed curves. Their shape is a property that is unaffected by translation, rescaling, rotation and reparameterization. To ensure that our framework is invariant to all shape-preserving transformations, we build on the Riemannian geometric elastic shape analysis framework. We first define a shape trace-variogram, which models spatial dependence among the shapes of planar curves. The trace-variogram is then used to define a spatially-weighted shape average and covariance, which enable exploration of shape variability in a local spatial region. Finally, we provide algorithms for estimation of the proposed spatially-informed summaries.

**Key words.** Shape Trace-variogram; Weighted Shape Average; Weighted Covariance and Principal Component Analysis

**MSC codes.** 62H11, 62R30, 62R10

**1. Introduction.** Statistical shape analysis tools have become indispensable in many scientific domains, including medical imaging [3, 6, 28] and biology (morphometrics) [19, 21]. In these fields, researchers are often interested in understanding how the geometry or shape of structures observed in images vary across space, usually defined as (a subset of) the image domain, under different conditions. For example, in medical applications, the shape of organs or cells along with their spatial configurations can provide crucial information about physiological conditions, disease progression and treatment outcomes [12, 22, 23]. Pathologists have long relied on nuclear shape abnormalities to distinguish benign from malignant cells [7], demon-

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\*Submitted to the editors February 10, 2026.

**Funding:** This work was supported in part by NIH R37-CA214955 (SK and KB), NSF DMS-2015374 (KB), and NSF DMS-2413747 (SK).

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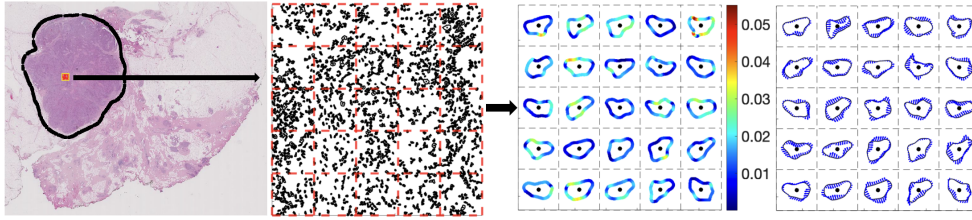


FIG. 1. *Local spatially-informed summarization of cell nucleus shapes in a histopathology image.*

46 strating the critical role of shape in characterizing cell states. In a tumor, these cells  
 47 interact with stromal (supportive) cells and other components to form the tumor  
 48 microenvironment (TME) [8, 13]. The TME exhibits a distinct structure featuring  
 49 spatial patterns that can guide cancer grading and predict treatment outcomes. For  
 50 instance, [9] showed that the location of immune cells in colorectal cancer is a key  
 51 prognostic factor, while [25] discovered a correlation between the local spatial struc-  
 52 ture of tumor-infiltrating lymphocytes (TILs) and overall survival. These findings  
 53 suggest that both, shape variation among cells and the spatial arrangement of TME  
 54 components, hold valuable information for cancer research. Thus, there is a clear need  
 55 for specialized methods capable of capturing spatially correlated shape data.

56 Many studies integrate cell-level morphological features with spatial information  
 57 to characterize tumors. Some are based on nuclear shape/texture descriptors, e.g.,  
 58 size, irregularity or fractal features, combined with spatial measures like cell-cell dis-  
 59 tances to predict recurrence or treatment response [2, 15, 31]. [14] combined single-cell  
 60 morphological features, e.g., area and eccentricity, with spatial interaction networks  
 61 to identify multicellular communities predictive of breast cancer recurrence and sur-  
 62 vival. [33] classified individual nuclei into cancer, stromal and immune cell types based  
 63 on morphology, and quantified their spatial distribution using Ripley’s K-function, re-  
 64 vealing prognostic patterns of stromal organization. However, all of these methods  
 65 represent shapes using scalar summaries, which do not capture their full geometric  
 66 structure.

67 When boundaries of objects observed in two-dimensional images are represented  
 68 as parameterized curves, i.e., functions of the form  $\beta : \mathcal{D} \rightarrow \mathbb{R}^2$  ( $\mathcal{D}$  is the param-  
 69 70 eterization domain of  $\beta$ ), their shape is a physical property that remains unchanged under  
 71 translation, rescaling, rotation and reparameterization. Statistical shape analysis is  
 72 then concerned with various tasks including (i) quantification of shape differences  
 73 via a distance, (ii) statistical summarization, i.e., computation of a representative  
 74 (sample average) shape and exploration of shape variability via principal component  
 75 analysis (PCA), and (iv) shape modeling and inference. While shape analysis ap-  
 76 proaches for independent shape data are well-established, see e.g., [18, 26], methods  
 77 which are appropriate for shapes that exhibit spatial dependence are scarce. Thus, we  
 78 define a statistical framework for local summarization of spatially dependent shape  
 79 data. The framework is accompanied by visualization tools, which enable easy inter-  
 80 pretability. To model spatial dependence among shapes, we extend the concept of the  
 81 trace-variogram, which has been effectively used for kriging and hierarchical cluster-  
 82 ing of spatial functional data [10, 11]. In essence, the proposed shape trace-variogram  
 83 encodes the dependence of shape variation on spatial distance. It allows efficient esti-  
 mation of weights for each observed shape that are used to define spatially-informed

84 shape summaries.

85 We provide an illustration of our approach in Figure 1. We start with a high-  
86 resolution histopathology image and focus on a region of interest (ROI, red square)  
87 inside a tumor area (black outline). We extract cell nucleus outlines and divide the  
88 ROI into equally sized, disjoint subregions. We then summarize shape variation of  
89 cell nuclei, while incorporating their spatial correlation, within each subregion. We  
90 compute a spatially-weighted shape average, based on a linear estimator that incor-  
91 porates weights estimated from the shape trace-variogram. We further quantify local  
92 shape variation by performing PCA, using a spatially-weighted covariance matrix,  
93 on the tangent space at this average shape. Summarization results are displayed as  
94 PC vector fields, and their pointwise magnitudes, on the spatially-weighted mean  
95 shape. Our techniques build on elastic shape representations and metrics, wherein  
96 parameterization variability is removed via registration [20, 26, 27, 32].

97 We demonstrate the effectiveness of spatially-informed summarization using syn-  
98 thetic and real data examples. In the synthetic study, we generate spatially correlated  
99 planar shapes, and show that the spatially-weighted average shape and associated  
100 principal modes of variation are effective at summarizing the data. In the real data  
101 application, we analyze cell nucleus outlines extracted from histopathology images of  
102 breast cancer [1, 30], which are known to exhibit complex morphological and spatial  
103 variability [12]. These nuclear shapes, along with their spatial distribution, can offer  
104 insights into how cells interact under various disease conditions. Our results indicate  
105 that local spatially-informed summaries can uncover morphological differences across  
106 tissue regions, which may be overlooked by global summaries that do not consider  
107 spatial dependence. As such, we demonstrate the potential of the developed tools for  
108 broader biomedical investigations.

109 Our contributions are as follows.

- 110 1. We define a shape trace-variogram to model spatial dependence between the *shapes*  
111 of curves, a property that is invariant to translation, rescaling, rotation and repara-  
112 meterization. The proposed variogram thus represents a novel extension of the  
113 functional trace-variograms proposed in [10, 11].
- 114 2. We use the shape trace-variogram to define a spatially-weighted shape average and  
115 covariance that can be used to explore shape variation in a local spatial region.  
116 The weights ensure that shapes close to a given spatial location have a stronger  
117 impact on the computed summaries.
- 118 3. We introduce concise, yet effective, visualizations of the spatially-informed sum-  
119 maries that aid in assessment of shape heterogeneity.
- 120 4. We apply our approach to histopathology images of breast cancer, resulting in  
121 meaningful summaries that effectively capture spatial shape heterogeneity of cell  
122 nuclei.

123 The rest of this article is organized as follows. Section 2 reviews summarization  
124 tools from elastic shape analysis under the assumption of independence. Section 3  
125 introduces the shape trace-variogram and describes its use in the definition and com-  
126 putation of (i) a spatially-weighted shape average, and (ii) spatially-weighted covari-  
127 ance. It additionally outlines how the proposed summaries are visualized. Section 4  
128 presents simulation results, while Section 5 describes an application to histopathology  
129 images of breast cancer. We close with a brief discussion in Section 6. The supple-  
130 ment includes (i) illustrations of the elastic shape analysis framework (Section 1), (ii)  
131 a proof of Proposition 3.1, which is related to the estimation of spatial weights via the  
132 shape trace-variogram (Section 2), (iii) empirical convergence analysis for Algorithm  
133 3.1 (Section 3), (iv) additional simulation results and sensitivity analysis (Section 4),

134 and (v) additional real data results (Section 5).

135 **2. Elastic Shape Analysis.** We provide an overview of the elastic shape analy-  
 136 sis framework; more details can be found in [26, 27]. Our particular focus is on the  
 137 case of planar closed curves, i.e., curves without a natural start/end point, as this is  
 138 the most pertinent case when analyzing cell nucleus shapes in histopathology images.  
 139 Illustrations of the framework are given in Section 1 in the supplement. Throughout  
 140 this section, we use  $|\cdot|$  and  $\langle \cdot, \cdot \rangle$  to denote the Euclidean norm and inner product in  
 141  $\mathbb{R}^2$ , and  $\|\cdot\|$  and  $\langle \cdot, \cdot \rangle$  to denote the norm and inner product on  $\mathbb{L}^2(\mathbb{S}^1, \mathbb{R}^2)$ , hereafter  
 142 denoted as  $\mathbb{L}^2$ , where  $\mathbb{S}^1$  is the unit circle on the plane.

143 **2.1. Representation, Pre-shape Space and Distance.** Let  $\mathcal{B} = \{\beta : \mathbb{S}^1 \rightarrow$   
 144  $\mathbb{R}^2 \mid \beta \text{ is absolutely continuous}\}$  denote the space of planar closed curves. As briefly  
 145 discussed in Section 1, the shape of  $\beta$  is unchanged under translation, rescaling,  
 146 rotation and reparameterization. Thus, we seek a metric structure on  $\mathcal{B}$  that is  
 147 invariant to these shape-preserving transformations. The reparameterization group  
 148  $\Gamma = \{\gamma : \mathbb{S}^1 \rightarrow \mathbb{S}^1 \mid \gamma \text{ is orientation-preserving diffeomorphism}\}$  acts on  $\mathcal{B}$  via compo-  
 149 sition:  $(\beta, \gamma) = \beta \circ \gamma$ . It is easy to see that the usual  $\mathbb{L}^2$  distance between two curves  
 150  $\beta_1, \beta_2 \in \mathcal{B}$  is not invariant to the reparameterization action:  $\|\beta_1 \circ \gamma - \beta_2 \circ \gamma\| \neq$   
 151  $\|\beta_1 - \beta_2\|$  for all  $\gamma \in \Gamma$  except  $\gamma_{id}(t) = t$ . This, in turn, means that the  $\mathbb{L}^2$  distance on  
 152  $\mathcal{B}$  is not suitable for parameterization-invariant comparison and modeling of shape.

153 To overcome the aforementioned challenge, we equip  $\mathcal{B}$  with an elastic Riemann-  
 154 ian metric [20]. Let  $\dot{\beta}$  be the coordinatewise derivative of  $\beta$ , which exists almost  
 155 everywhere on  $\mathbb{S}^1$ ; set  $\dot{\beta}(t) = 0$  on the set of all  $t \in \mathbb{S}^1$  where it is not defined point-  
 156 wise. Outside of this set assume  $|\dot{\beta}| \neq 0$ . Then,  $p = |\dot{\beta}|$  and  $\theta = \dot{\beta}|\dot{\beta}|^{-1}$  denote  
 157 the speed and angle functions, which fully characterize  $\beta$  since  $\dot{\beta} = p\theta$ . Thus, this  
 158 defines a bijective map (up to a translation) from  $\mathcal{B}$  to the product space  $\mathcal{P} \times \Theta$ ,  
 159 where  $\mathcal{P} = \{p : \mathbb{S}^1 \rightarrow \mathbb{R}_+\}$  and  $\Theta = \{\theta : \mathbb{S}^1 \rightarrow \mathbb{S}^1\}$ . Letting  $(\delta p_1, \delta \theta_1)$  and  $(\delta p_2, \delta \theta_2)$   
 160 denote two vectors in the tangent space at  $(p, \theta)$ ,  $T_{(p, \theta)}(\mathcal{P} \times \Theta)$ , the family of elastic  
 161 Riemannian metrics is defined as

(2.1)

$$162 \quad \langle \langle (\delta p_1, \delta \theta_1), (\delta p_2, \delta \theta_2) \rangle \rangle_{(p, \theta)} := a \int_{\mathbb{S}^1} \delta p_1(t) \delta p_2(t) \frac{1}{p(t)} dt + b \int_{\mathbb{S}^1} \langle \delta \theta_1(t), \delta \theta_2(t) \rangle p(t) dt,$$

163 where  $a, b > 0$ . Note that the first term measures variations in the speed function  
 164 while the second term measures variations in the angle function. The constants  $a$  and  
 165  $b$  provide relative weights for the two terms. An important property of this metric is  
 166 that it is preserved when two curves are reparameterized in the same way; this is true  
 167 for any choice of  $a$  and  $b$ . However, the resulting Riemannian distance does not have  
 168 a closed-form expression and is difficult to compute.

169 To simplify the metric structure and retain the desirable property of invariance  
 170 to reparameterization, we employ the square-root velocity function (SRVF) [27]. The  
 171 SRVF of a curve  $\beta \in \mathcal{B}$  is defined via the mapping  $Q : \mathcal{B} \rightarrow \mathbb{L}^2$  as

$$172 \quad (2.2) \quad q \equiv Q(\beta) := \begin{cases} \frac{\dot{\beta}}{\sqrt{|\dot{\beta}|}}, & |\dot{\beta}| \neq 0 \\ 0, & |\dot{\beta}| = 0 \text{ or } \dot{\beta} \text{ is not defined.} \end{cases}$$

173 A key consequence of this transformation is that the elastic Riemannian metric on  
 174  $\mathcal{B}$ , with  $a = 1/4$  and  $b = 1$ , simplifies to the  $\mathbb{L}^2$  metric on the space of SRVFs [27].  
 175 The SRVF of a closed curve  $\beta$  satisfies the following constraint:  $\int_{\mathbb{S}^1} q(t) |q(t)| dt = 0$ .  
 176 Finally, the inverse mapping,  $Q^{-1}$ , can be computed (up to a translation) using

177  $\int_0^t q(u)|q(u)|du$ , where the domain  $\mathbb{S}^1$  has been unwrapped to the interval  $[0, 2\pi]$ ; note  
 178 that this inverse mapping subsumes projection of the SRVF  $q$  into the space of closed  
 179 curves using Algorithm 18 in [26, Ch. 6] to ensure that the aforementioned closure  
 180 condition is satisfied.

181 We now return to the problem of removing nuisance variation from the representa-  
 182 tion space. Translation is automatically removed under the SRVF since its definition  
 183 only involves the derivative of  $\beta$ . To remove scale, we normalize all curves to have unit  
 184 length, resulting in SRVFs with unit  $\mathbb{L}^2$  norm:  $L(\beta) = \int_{\mathbb{S}^1} |\dot{\beta}(t)|dt = \int_{\mathbb{S}^1} |q(t)|^2 dt =$   
 185  $\|q\|^2 = 1$ . This leads to the pre-shape space for planar closed curves, defined as  
 186  $\mathcal{C} = \{q \in \mathbb{L}^2 \mid \|q\| = 1, \int_{\mathbb{S}^1} q(t)|q(t)|dt = 0\}$ , which is a subset of the Hilbert sphere.  
 187 The space  $\mathcal{C}$  is endowed with the ambient  $\mathbb{L}^2$  metric, resulting in the extrinsic pre-  
 188 shape distance  $d_{\mathcal{C}}(q_1, q_2) = \|q_1 - q_2\|$  between  $q_1, q_2 \in \mathcal{C}$ . Removal of rotation and  
 189 parameterization variability cannot be achieved via normalization. Instead, we use  
 190 the notion of equivalence classes as a formal definition of shape.

191 **2.2. Registration, Shape Space and Distance.** Let  $O \in SO(2) = \{R \in$   
 192  $\mathbb{R}^{2 \times 2} \mid RR^T = R^T R = I_2, \det(R) = 1\}$  represent a rotation matrix. The rotation of  
 193 a curve  $\beta \in \mathcal{B}$  by  $O \in SO(2)$  is given by  $O\beta$ . The corresponding transformation of  
 194 the SRVF of  $\beta$ ,  $Q(\beta) = q$ , is the same, i.e.,  $Oq$ . The reparameterization of  $\beta$  via  $\gamma$ ,  
 195  $(\beta, \gamma) = \beta \circ \gamma$ , results in the following transformation of its SRVF:  $(q, \gamma) = (q \circ \gamma)\sqrt{\dot{\gamma}}$ .  
 196 It is now easy to verify that the distance  $d_{\mathcal{C}}$  is preserved if two curves  $\beta_1, \beta_2 \in \mathcal{B}$   
 197 are reparameterized using the same  $\gamma \in \Gamma$  and rotated using the same  $O \in SO(2)$   
 198 ( $q_1 = Q(\beta_1), q_2 = Q(\beta_2)$ ):

$$199 \quad (2.3) \quad \|O(q_1, \gamma) - O(q_2, \gamma)\| = \|q_1 - q_2\|.$$

200 To account for rotation and parameterization variation, we define equivalence classes  
 201 of the form  $[q] := \{O(q, \gamma) \mid O \in SO(2), \gamma \in \Gamma\}$ , which serve as the definition of shape.  
 202 In other words, all possible rotations and reparameterizations of a curve are unified  
 203 via an equivalence class, which uniquely represents the shape of that curve. This leads  
 204 to the definition of the shape space as the quotient space  $\mathcal{S} = \mathcal{C}/(SO(2) \times \Gamma) = \{[q]\}$ .  
 205 The extrinsic shape distance on  $\mathcal{S}$  is defined via the distance  $d_{\mathcal{C}}$  as (for  $[q_1], [q_2] \in \mathcal{S}$ )  
 206

$$207 \quad (2.4) \quad d_{\mathcal{S}}([q_1], [q_2]) = \inf_{(O, \gamma) \in SO(2) \times \Gamma} \|q_1 - O(q_2, \gamma)\|.$$

208 To compute this distance, we have to determine an optimal rotation  $\hat{O}$  and an optimal  
 209 reparameterization  $\hat{\gamma}$  of  $q_2$  with respect to  $q_1$ . Optimization over  $SO(2)$  is performed  
 210 via Procrustes analysis [16]. Optimization over  $\Gamma$  is split into (i) an exhaustive search  
 211 over starting points on  $\mathbb{S}^1$ , and (ii) Dynamic Programming given each starting point  
 212 [24]. The joint optimization over  $SO(2) \times \Gamma$  is referred to as elastic registration, with  
 213  $\hat{O}(\beta_2 \circ \hat{\gamma})$  being optimally registered to  $\beta_1$ , and ensures that shape comparisons are  
 214 natural, i.e., they preserve geometric features of the shapes being compared.

215 **2.3. Shape Summaries Under Independence.** Let  $\beta_1, \dots, \beta_n \in \mathcal{B}$  denote a  
 216 sample of *independent* planar closed curves with SRVFs  $q_1, \dots, q_n \in \mathcal{C}$ . Their extrinsic

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<sup>2</sup>Under the SRVF, a specific instance of the elastic Riemannian metric simplifies to the  $\mathbb{L}^2$  metric. When restricted to the pre-shape space of closed curves,  $\mathcal{C}$ , the resulting *intrinsic* distance does not have a closed-form expression [27]. However, throughout this manuscript, we employ the *extrinsic* distance  $d_{\mathcal{C}}$ , which does have a closed-form expression and results in simplified computation.

**Algorithm 2.1** Extrinsic Karcher Mean Shape

- 
- 1: Input:  $q_1, \dots, q_n$ .
  - 2: Output:  $\hat{\mu}_q; \hat{O}_1(q_1, \hat{\gamma}_1), \dots, \hat{O}_n(q_n, \hat{\gamma}_n)$ .
  - 3: Initialize  $k = 1$ ,  $\epsilon > 0$  (small),  $\delta > 0$ ,  $\hat{\mu}_q^{(1)}$ ,  $\|\bar{z}^{(0)}\| > 0$  (large).
  - 4: **while**  $\|\bar{z}^{(k-1)}\| > \epsilon$  **do**
  - 5: Register each  $q_i$  to  $\hat{\mu}_q^{(k)}$  using (2.4) to obtain  $\hat{O}_i^{(k)}(q_i, \hat{\gamma}_i^{(k)})$ .
  - 6: Compute deviation vector from  $\hat{\mu}_q^{(k)}$  for each  $\hat{O}_i^{(k)}(q_i, \hat{\gamma}_i^{(k)})$ :  $z_i = \hat{O}_i^{(k)}(q_i, \hat{\gamma}_i^{(k)}) - \hat{\mu}_q^{(k)}$ .
  - 7: Compute average direction:  $\bar{z}^{(k)} = \frac{1}{n} \sum_{i=1}^n z_i$ .
  - 8: Update:  $\hat{\mu}_q^{(k+1)} \leftarrow \hat{\mu}_q^{(k)} + \delta \bar{z}^{(k)}$ .
  - 9: Normalize:  $\hat{\mu}_q^{(k+1)} \leftarrow \frac{\hat{\mu}_q^{(k+1)}}{\|\hat{\mu}_q^{(k+1)}\|}$ .
  - 10: Project  $\hat{\mu}_q^{(k+1)}$  using Algorithm 18 in [26, Ch. 6] to ensure that closure condition is satisfied.
  - 11: Set  $k \leftarrow k + 1$ .
  - 12: **end while**
  - 13: Register each  $q_i$  to  $\hat{\mu}_q$  using (2.4) to obtain  $\hat{O}_i(q_i, \hat{\gamma}_i)$ .
- 

217 sample Karcher mean shape,  $[\hat{\mu}_q]$ , is defined as a minimizer of

$$218 \quad (2.5) \quad \mathcal{S} \ni [q] \mapsto \sum_{i=1}^n d_{\mathcal{S}}([q], [q_i])^2.$$

219 In practice, we select a single element  $\hat{\mu}_q \in [\hat{\mu}_q]$  as a representative for further analy-  
 220 ses. The mean can be visualized after inversion,  $\hat{\mu}_\beta = Q^{-1}(\hat{\mu}_q)$ . The procedure for  
 221 computing the extrinsic Karcher mean shape is presented as Algorithm 2.1.

222 To explore variation in a sample of shapes, we compute the covariance matrix  
 223 in the tangent space  $T_{\hat{\mu}_q}(\mathcal{S})$  as follows. With respect to the extrinsic spherical  
 224 metric on  $\mathcal{C}$ , we first project the registered curves (shape representatives)  $q_1^* =$   
 225  $\hat{O}_1(q_1, \hat{\gamma}_1), \dots, q_n^* = \hat{O}_n(q_n, \hat{\gamma}_n)$  to  $T_{\hat{\mu}_q}(\mathcal{S})$  using the inverse-exponential map [26,  
 226 Ch. 3]:  $v_i := \exp_{\hat{\mu}_q}^{-1}(q_i^*) = \theta \sin(\theta)^{-1} (q_i^* - \cos(\theta)\hat{\mu}_q)$ , where  $\theta = \cos^{-1}(\langle \hat{\mu}_q, q_i^* \rangle)$ .  
 227 While theoretically shape is an infinite-dimensional object, at the implementation  
 228 stage, it is approximated using a finite number of points  $T$ . Thus, each  $v_i$  is a matrix  
 229 of size  $2 \times T$ . We further reshape  $v_i$  to a vector of size  $2T$  resulting in  $v_i^* \in \mathbb{R}^{2T}$ . The  
 230 sample shape covariance matrix is then given by

$$231 \quad (2.6) \quad \hat{K} := \frac{1}{n-1} \sum_{i=1}^n v_i^* v_i^{*\top} \in \mathbb{R}^{2T \times 2T}.$$

232 To perform PCA, we apply singular value decomposition (SVD) to the sample covari-  
 233 ance matrix to obtain  $\hat{K} = U \Sigma U^\top$ . The columns of  $U$  contain the principal directions  
 234 of shape variation; the diagonal elements of  $\Sigma$  are the variances along each direction  
 235 (ordered from largest to smallest). The total sample shape variance can be computed  
 236 using the trace of  $\Sigma$ .

237 To visualize shape variation along the  $k$ th principal direction, we compute a dis-  
 238 placement, in units of standard deviation, from the mean shape on the tangent space  
 239 and map it back to the shape space via the exponential map,  $q_k^l = \exp_{\hat{\mu}_q} \left( l \sqrt{\Sigma_{kk}} U_k \right)$ ,

240 where  $l$  controls the size of the displacement,  $\Sigma_{kk}$  is the  $k$ th diagonal element of  $\Sigma$ ,  
 241 and  $U_k$  is the  $k$ th column of  $U$  reshaped to a matrix of size  $2 \times T$ . For  $v \in T_{\hat{\mu}_q}(\mathcal{S})$ ,  
 242 the exponential map is given by  $\exp_{\hat{\mu}_q}(v) = \cos(\|v\|)\hat{\mu}_q + \sin(\|v\|)v\|v\|^{-1}$  [26, Ch. 3].  
 243 As before, the principal directions of shape variation can be visualized after inversion,  
 244  $\beta_k^l = Q^{-1}(q_k^l)$ .

245 **3. Spatially-informed Shape Summaries.** While the elastic shape analysis  
 246 framework described in Section 2 is effective in summarizing independent shape data,  
 247 as illustrated in Section 1 in the supplement, it does not account for potential de-  
 248 pendence among observed shapes. Depending on the specific application of interest,  
 249 shape dependence may arise in many different ways. However, in the current work,  
 250 we focus on spatial dependence among cell nucleus shapes in histopathology images.  
 251 Thus, we introduce a new framework, with elastic shape analysis serving as a building  
 252 block, that leverages local spatial dependence for shape summarization.

253 Let  $X \subset \mathbb{R}^2$  denote the spatial domain; our focus here is on two-dimensional  
 254 histopathology images, but the concepts presented in this section can be extended  
 255 to higher-dimensional (and irregular) spatial domains. In spatial statistics, the vari-  
 256 ogram [5] is a core tool for summarizing dependence among spatially-indexed objects.  
 257 In particular, [10] proposed the trace-variogram for functional data. Under the as-  
 258 sumption that the functional random field  $\{f_s, s \in X\}$  with values in  $\mathbb{L}^2([a, b], \mathbb{R})$  is  
 259 second-order stationary and isotropic<sup>3</sup>, the trace-variogram is defined as

$$260 \quad (3.1) \quad V_f(h) := \frac{1}{2}E\left[\|f_s - f_{s'}\|^2\right],$$

261 where  $h = |s - s'|$  denotes the Euclidean distance between spatial locations  $s$  and  $s'$ .  
 262 In this definition, each spatially indexed function  $f_s$  is univariate with domain given  
 263 by the interval  $[a, b]$ . The trace-variogram  $V_f$  is a function of the spatial distance  $h$ ,  
 264 and captures variation among the functional random field at different distance lags  
 265 via the  $\mathbb{L}^2$  distance. Then, given a realization of the random field  $\{f_{s_i}, s_i \in X\}$   
 266 ( $i = 1, \dots, n$ ) at  $n$  locations, one can compute the empirical trace-variogram using

$$267 \quad (3.2) \quad \hat{V}_f(h) = \frac{1}{2|N(h)|} \sum_{i,j \in N(h)} \|f_{s_i} - f_{s_j}\|^2,$$

268 where  $N(h) = \{(s_i, s_j) \mid |s_i - s_j| = h\}$  is a spatial neighborhood. For irregularly spaced  
 269 functional data,  $N(h)$  is often modified to  $N(h) = \{(s_i, s_j) \mid |s_i - s_j| \in (h - \epsilon, h + \epsilon)\}$ .

270 If there exists spatial dependence among the functions  $\{f_{s_i}\}$ , the values of the  
 271 empirical trace-variogram will be smaller at shorter spatial distance lags. This is due  
 272 to the fact that functions that are located close to each other in the spatial domain  
 273  $X$  will exhibit less variation than those located far away from each other. The trace-  
 274 variogram allows estimation of spatial weights for the sample  $\{f_{s_i}\}$ , which can be  
 275 subsequently used for prediction at a particular location and hierarchical clustering  
 276 of spatial functional data. The definition of the trace-variogram was extended to the  
 277 case of spatial functional data with phase variation in [11]. In this work, we define  
 278 a *shape trace-variogram* to construct local shape summaries: (i) a spatially-weighted  
 279 shape average, and (ii) a spatially-weighted covariance matrix for PCA.

<sup>3</sup>Recall that a second-order stationary and isotropic random field has a constant mean, finite variance, and covariance that only depends on the spatial distance between observations.

280 **3.1. Shape Trace-variogram and its Estimation.** We begin with a spatial  
 281 random field of planar closed curves,  $\{\beta_s, s \in X\}$  with values in  $\mathcal{B}$ , which is trans-  
 282 formed to a spatial random field of SRVFs,  $\{q_s, s \in X\}$  with values in  $\mathcal{C} \subset \mathbb{L}^2$ . In  
 283 most applications, for example those involving cell nucleus shapes in histopathology  
 284 images, assuming that the field  $\{q_s, s \in X\}$  is second-order stationary with constant  
 285 mean shape is inappropriate. As such, the spatially-weighted average shape we wish  
 286 to estimate is the shape of the value  $q_{s_0} \equiv q_0$  of the SRVF random field at a *fixed*,  
 287 *known location*  $s_0 \in X$ ; for all results presented in Sections 4 and 5 we choose  $s_0$  to  
 288 be the center of the spatial domain  $X$ . The estimate will be a local spatially-weighted  
 289 average of the shapes of observed values of  $\{q_s, s \in X\}$  in a neighborhood of  $s_0$ , and  
 290 is referred to within the spatial statistics literature as a kriging predictor at  $s_0$ .

291 The shape of the random field  $\{q_s, s \in X\}$  is encoded within the random field  
 292  $\{[q_s], s \in X\}$  of equivalence classes. This object is difficult to work with in practice:  
 293 probability distributions on the shape space  $\mathcal{S}$  may behave in non-standard ways  
 294 due to the infinite-dimensionality of the equivalence class  $[q_s]$  for each  $s \in X$  and  
 295 that of the non-compact reparameterization group  $\Gamma$ . To avoid such difficulties, we  
 296 instead work with the transformed field  $\{O_s(q_s, \gamma_s) \in [q_s], s \in X\}$  that represents,  
 297 for each  $s \in X$ , the registration of the field  $\{q_s, s \in X\}$  to a preferred element  
 298  $O_{s_0}(q_{s_0}, \gamma_{s_0}) \equiv O_0(q_0, \gamma_0)$  of the shape  $[q_0]$ , where  $\{O_s, s \in X\}$  and  $\{\gamma_s, s \in X\}$  are,  
 299 respectively, random fields with values in  $SO(2)$  and  $\Gamma$  satisfying

$$300 \quad (O_s, \gamma_s) = \underset{(O, \gamma) \in SO(2) \times \Gamma}{\operatorname{arg\,inf}} \|O(q_s, \gamma) - O_0(q_0, \gamma_0)\|^2, \quad s \in X.$$

301 Abusing terminology, we refer to  $\{O_s(q_s, \gamma_s) \in [q_s], s \in X\}$ , consisting of shape rep-  
 302 resentatives for  $\{q_s, s \in X\}$ , as the *shape random field*<sup>4</sup>, which implicitly depends  
 303 on the representative  $O_0(q_0, \gamma_0)$  of the shape  $[q_0]$  at location  $s_0$ . At a different loca-  
 304 tion,  $\tilde{s}_0$ , we obtain a different shape SRVF random field defined for the purpose of  
 305 estimating a different spatially-weighted average shape  $O_{\tilde{s}_0}(q_{\tilde{s}_0}, \gamma_{\tilde{s}_0})$ .

306 As done for the functional random field in [10], we assume that the shape random  
 307 field is second-order stationary and isotropic, and accordingly define the *shape trace-*  
 308 *variogram*

$$309 \quad (3.3) \quad V_{sh}(h) := \frac{1}{2} E \left[ \|O_s(q_s, \gamma_s) - O_{s'}(q_{s'}, \gamma_{s'})\|^2 \right],$$

310 where  $h = |s - s'|$ . It is important to note that the definition of the shape trace-  
 311 variogram  $V_{sh}$  is novel and can be viewed as a modification of the functional trace-  
 312 variogram in (3.1) that incorporates registration: the presence of shape-preserving  
 313 nuisance variation in  $\{q_s, s \in X\}$  is handled via the random fields  $\{O_s, s \in X\}$   
 314 and  $\{\gamma_s, s \in X\}$ , which determine the shape random field via the registration of  
 315  $\{q_s, s \in X\}$  to the target  $O_0(q_0, \gamma_0)$ . Thus, importantly, the definition of the shape  
 316 trace-variogram implicitly depends on the chosen spatial location,  $s_0$ , through the esti-  
 317 mation target  $O_0(q_0, \gamma_0)$ . As such, one can define a family of shape trace-variograms  
 318 by varying the location  $s_0$  resulting in different shape random fields and spatially-  
 319 weighted average shapes.

320 Similarly to the functional trace-variogram, the shape trace-variogram captures  
 321 variation among the shape random field at different spatial distance lags. Evidently,

<sup>4</sup>Note that the original random field  $\{q_s\}$  and the shape random field  $\{O_s(q_s, \gamma_s)\}$  are both random fields of SRVFs. Their difference lies in the fact that the latter depends on the registration of  $\{q_s\}$  to  $O_0(q_0, \gamma_0)$ , and models the variation in  $\{q_s\}$  relative to the spatially-weighted average  $O_0(q_0, \gamma_0)$ .

322 the shape trace-variogram is invariant to simultaneous rotation and reparameteriza-  
 323 tion of  $\{q_s, s \in X\}$  using any  $(O, \gamma) \in SO(2) \times \Gamma$ , which is a direct consequence of  
 324 the isometry property verified in (2.3): it captures pure spatial shape variation.

325 We now define an empirical version of the shape trace-variogram, needed to  
 326 compute an estimator,  $\tilde{q}_0$ , of the spatially-weighted shape average  $O_0(q_0, \gamma_0)$ . Let  
 327  $\{q_{s_i}, s_i \in X\}$  denote an observed sample of  $n$  curves represented by their SRVFs.  
 328 The definition requires the estimator  $\tilde{q}_0$ , which will be defined in the sequel. To sim-  
 329 plify notation, we use the subscript  $i$  for the spatial location  $s_i$ . Then, the empirical  
 330 shape trace-variogram is defined as

$$331 \quad (3.4) \quad \hat{V}_{sh}(h) = \frac{1}{2|N(h)|} \sum_{i,j \in N(h)} \|\hat{O}_i(q_i, \hat{\gamma}_i) - \hat{O}_j(q_j, \hat{\gamma}_j)\|^2,$$

332 where  $N(h) = \{(s_i, s_j) \mid |s_i - s_j| \in (h - \epsilon, h + \epsilon)\}$  and  $(\hat{O}_i, \hat{\gamma}_i)$  minimize  $(O, \gamma) \mapsto$   
 333  $\|\tilde{q}_0 - O(q_i, \gamma)\|$  for  $i = 1, \dots, n$ . To ensure that the estimated variogram is condition-  
 334 ally negative definite [5], we additionally fit a Matérn model (with a predetermined  
 335 smoothness) to  $\hat{V}_{sh}$  using ordinary least squares. The resulting Matérn fit is used in  
 336 subsequent analyses, and with a slight abuse in notation, will henceforth be denoted  
 337 by  $\hat{V}_{sh}$ .

338 **3.2. Estimation of Spatially-weighted Average Shape.** In (3.4),  $\hat{O}_i(q_i, \hat{\gamma}_i)$   
 339 serves the role of a representative of the shape equivalence class  $[q_i]$ . Given  $(\hat{O}_i, \hat{\gamma}_i)$   
 340 ( $i = 1, \dots, n$ ), the empirical shape trace-variogram (and the Matérn fit) is computed  
 341 once the estimator  $\tilde{q}_0$  is prescribed. On the other hand, definition of the kriging  
 342 estimator  $\tilde{q}_0$  requires using  $\hat{V}_{sh}$ . We now detail the issue and present an algorithm  
 343 that enables computation of  $\hat{V}_{sh}$  while simultaneously allowing for the computation  
 344 of  $\tilde{q}_0$  and  $(\hat{O}_i, \hat{\gamma}_i)$  ( $i = 1, \dots, n$ ) via registration to  $\tilde{q}_0$ .

345 For a given vector  $\boldsymbol{\eta} = (\eta_1, \dots, \eta_n)^\top \in \Delta := \{\mathbf{x} \in \mathbb{R}^n \mid \sum_{i=1}^n x_i = 1\}$ , the  
 346 spatially-weighted average shape estimator,  $\tilde{q}_0 \in [\tilde{q}_0]$ , is defined as a minimizer of  
 347

$$348 \quad (3.5) \quad q \mapsto \sum_{i=1}^n \eta_i d_{\mathcal{S}}([q], [q_i])^2,$$

349 which is directly available as

$$350 \quad (3.6) \quad \tilde{q}_0 = \frac{\sum_{i=1}^n \eta_i \hat{O}_i(q_i, \hat{\gamma}_i)}{\|\sum_{i=1}^n \eta_i \hat{O}_i(q_i, \hat{\gamma}_i)\|}.$$

351 For clarity of exposition, for just this section, we make explicit the dependence of the  
 352 estimator  $\tilde{q}_0$  on  $\boldsymbol{\eta}$  and denote it by  $\tilde{q}_0(\boldsymbol{\eta})$ . Note that  $\tilde{q}_0(\boldsymbol{\eta})$  is not a weighted sample  
 353 Karcher mean in a strict sense since some of the elements of  $\boldsymbol{\eta}$  may be negative. On the  
 354 other hand, given the nuisance transformations  $(\hat{O}_i, \hat{\gamma}_i)$  ( $i = 1, \dots, n$ ), the coefficient  
 355 vector  $\boldsymbol{\eta}$  is obtained by minimizing the expected shape prediction error functional:

$$356 \quad (3.7) \quad \boldsymbol{\eta} = \operatorname{argmin}_{\mathbf{x} \in \Delta} E [\|\tilde{q}_0(\mathbf{x}) - O_0(q_0, \gamma_0)\|^2].$$

357 If  $(\hat{O}_i, \hat{\gamma}_i)$  ( $i = 1, \dots, n$ ) can be recovered exactly, the optimization problem in (3.7) is  
 358 equivalent to minimizing  $\mathbf{x} \mapsto \mathbf{x}^\top V_{sh} \mathbf{x}$ , where  $V_{sh} = V_{sh}(h_{i0}) + V_{sh}(h_{0j}) - V_{sh}(h_{ij})$ ,  
 359  $h_{ij} = |s_i - s_j|$  is the population shape trace-variogram in (3.3). Hence, under this  
 360 setting, the spatially-weighted average shape estimator depends only on the shape

**Algorithm 3.1** Spatially-weighted Average Shape

- 
- 1: Input:  $(s_1, q_1), \dots, (s_n, q_n)$ ;  $s_0 \in X$ .
  - 2: Output:  $\tilde{q}_0$ ;  $V_{sh}$ ;  $\boldsymbol{\eta}$ ;  $\hat{O}_1(q_1, \hat{\gamma}_1), \dots, \hat{O}_n(q_n, \hat{\gamma}_n)$ .
  - 3: Initialize  $k = 1$ ,  $\epsilon > 0$  (small),  $\delta > 0$ ,  $\tilde{q}_0^{(0)}$  as  $q_j$  where  $j = \underset{i \in \{1, \dots, n\}}{\operatorname{argmin}} |s_0 - s_i|$ ,  $\|\tilde{z}^{(0)}\| > 0$  (large).
  - 4: **while**  $\|\tilde{z}^{(k-1)}\| > \epsilon$  **do**
  - 5:   Register each  $q_i$  to  $\tilde{q}_0^{(k)}$  using (2.4) to obtain  $\hat{O}_i^{(k)}(q_i, \hat{\gamma}_i^{(k)})$ .
  - 6:   Compute  $\hat{V}_{sh}^{(k)}$ , using  $\{\hat{O}_i^{(k)}(q_i, \hat{\gamma}_i^{(k)})\}$  and (3.4), and corresponding Matérn fit.
  - 7:   Estimate weight vector  $\boldsymbol{\eta}^{(k)}$  using  $\hat{V}_{sh}^{(k)}$ .
  - 8:   Compute deviation vector from  $\tilde{q}_0^{(k)}$  for each  $\hat{O}_i^{(k)}(q_i, \hat{\gamma}_i^{(k)})$ :  $z_i = \hat{O}_i^{(k)}(q_i, \hat{\gamma}_i^{(k)}) - \tilde{q}_0^{(k)}$ .
  - 9:   Compute spatially-weighted average direction:  $\tilde{z}^{(k)} = \frac{1}{n} \sum_{i=1}^n \eta_i^{(k)} z_i$ .
  - 10:   Update:  $\tilde{q}_0^{(k+1)} \leftarrow \tilde{q}_0^{(k)} + \delta \tilde{z}^{(k)}$ .
  - 11:   Normalize:  $\tilde{q}_0^{(k+1)} \leftarrow \frac{\tilde{q}_0^{(k+1)}}{\|\tilde{q}_0^{(k+1)}\|}$ .
  - 12:   Project  $\tilde{q}_0^{(k+1)}$  using Algorithm 18 in [26, Ch. 6] to ensure that closure condition is satisfied.
  - 13:   Set  $k \leftarrow k + 1$ .
  - 14: **end while**
  - 15: Register each  $q_i$  to  $\tilde{q}_0$  using (2.4) to obtain  $\hat{O}_i(q_i, \hat{\gamma}_i)$ .
  - 16: Compute  $\hat{V}_{sh}$ , using  $\{\hat{O}_i(q_i, \hat{\gamma}_i)\}$  and (3.4), and corresponding Matérn fit.
  - 17: Estimate weight vector  $\boldsymbol{\eta}$  using  $\hat{V}_{sh}$ .
- 

361 trace-variogram  $V_{sh}$ . In practice, the minimization problem is solved using the empirical  
 362 trace-variogram  $\hat{V}_{sh}$ . This is also discussed in detail in [11] for functional data  
 363 with phase variation. A formal statement of this result is provided as Proposition 3.1;  
 364 the proof is included in Section 2 in the supplement.

365 **PROPOSITION 3.1.** *Under the assumption that  $(O_i, \gamma_i)$  can be estimated exactly*  
 366 *such that  $\hat{O}_i(q_i, \hat{\gamma}_i) = O_i(q_i, \gamma_i)$ , the vector of coefficients  $\boldsymbol{\eta} \in \Delta$  that minimizes*  
 367  *$\boldsymbol{x} \mapsto E [\|\tilde{q}_0(\boldsymbol{x}) - O_0(q_0, \gamma_0)\|^2]$  also minimizes  $\boldsymbol{x}^\top V_{sh} \boldsymbol{x}$ , where the matrix  $V_{sh} \in \mathbb{R}^{n \times n}$*   
 368 *is composed of elements  $V_{sh} = V_{sh}(h_{i0}) + V_{sh}(h_{0j}) - V_{sh}(h_{ij})$ ,  $h_{ij} = |s_i - s_j|$  ( $i, j =$   
 369  $1, \dots, n$ ).*

370 **Remark.** Evidently, the spatially-weighted average shape estimation problem  
 371 cannot be defined using a single cost function. Estimation of  $\tilde{q}_0(\boldsymbol{\eta})$  and  $(\hat{O}_i, \hat{\gamma}_i)$   
 372 ( $i = 1, \dots, n$ ) is only possible if the coefficient vector  $\boldsymbol{\eta}$  is known. Similarly, estimation  
 373 of  $\boldsymbol{\eta}$  requires the empirical shape trace-variogram, which in turn depends on  $\tilde{q}_0(\boldsymbol{\eta})$   
 374 and  $(\hat{O}_i, \hat{\gamma}_i)$  ( $i = 1, \dots, n$ ). Thus, as discussed next, we utilize an iterative algorithm  
 375 that alternates between the two optimization problems.

376 Unfortunately, exact recovery of rotation or reparameterization is difficult when  
 377 the observed curves exhibit complex noise; see [4, 17] for more details in the case  
 378 of functional data with phase variation, which is similar to the shape setting we  
 379 consider. In addition, accurate estimation of these nuisance transformations depends  
 380 on accurate recovery of the underlying spatially-weighted average shape. We propose  
 381 an iterative procedure, detailed in Algorithm 3.1, which simultaneously estimates the  
 382 (i) spatially-weighted average shape, and (ii) the shape trace-variogram and associated

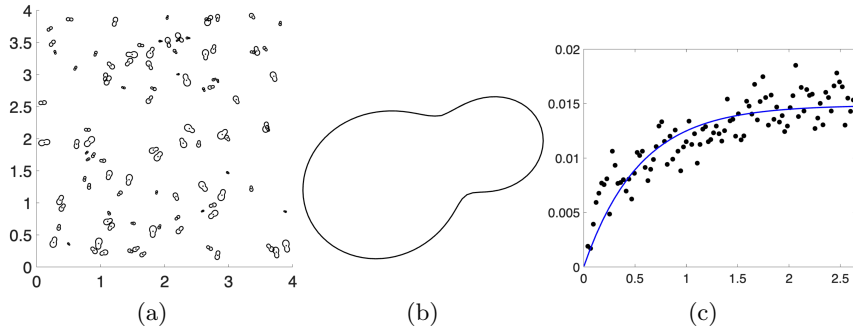


FIG. 2. (a) Simulated curves with spatially varying shapes. (b) Spatially-weighted average shape. (c) Empirical shape trace-variogram (black) and fitted Matérn model (blue).

383 vector of spatial weights  $\boldsymbol{\eta}$ . First, we initialize the shape average  $\tilde{q}_0(\boldsymbol{\eta})$  as the SRVF  
 384 observed at a spatial location that is closest to  $s_0$ . Empirically, we have found that the  
 385 final estimate of the spatially-weighted average shape is not sensitive to initialization  
 386 of Algorithm 3.1. We then iterate between the following steps until convergence: (i)  
 387 registration of the sample SRVFs  $\{q_i\}$  to the current shape average using (2.4), (ii)  
 388 estimation of the shape trace-variogram using (3.4) and corresponding Matérn fit,  
 389 (iii) computation of the coefficient vector  $\boldsymbol{\eta}$ , and (iv) incremental update of the shape  
 390 average using weighted deviations from the current estimate.

391 Recall that, under independence, computation of the Karcher mean shape of the  
 392 sample SRVFs  $q_1, \dots, q_n$  (Section 2.3; Algorithm 2.1) uses equal weights for each  
 393 sample. Under spatial dependence, we estimate a spatially-weighted average, where  
 394 the weight (coefficient) for each sample is determined by the shape trace-variogram  
 395  $\hat{V}_{sh}$ . In particular, spatial dependence among the shapes of  $\{q_i\}$  is incorporated  
 396 via the trace-variogram estimate at each iteration  $k$ ,  $\hat{V}_{sh}^{(k)}$ , to determine the weight  
 397 vector  $\boldsymbol{\eta}^{(k)}$ . Thus, the algorithm performs spatially local registration of curves, with  
 398 respect to a spatially-weighted average, that results in their shape representatives.  
 399 Empirically, we have not observed any issues with convergence of Algorithm 3.1; we  
 400 leave the formal convergence analysis as future work. Section 3 in the supplement  
 401 presents convergence plots for a few simulated examples.

402 Figure 2 shows an example of estimation of a spatially-weighted average shape,  
 403 and the associated shape trace-variogram, using Algorithm 3.1. Simulated spatially-  
 404 indexed planar closed curves are shown in (a), where  $X = [0, 4] \times [0, 4]$ ; here, the  
 405 sample size is  $n = 100$ . Panel (b) displays the estimated spatially-weighted average  
 406 shape  $\tilde{q}_0$  at spatial location  $s_0 = (2, 2)$ . In (c), we show the final empirical shape  
 407 trace-variogram (black points) and the Matérn fit with smoothness equal to 0.5.

408 **3.3. Spatially-weighted Covariance and PCA.** In addition to estimating a  
 409 spatially-weighted average shape, we want to characterize and visualize shape varia-  
 410 tion in the sample. The vector  $\boldsymbol{\eta}$  estimated by Algorithm 3.1 provides spatial weights  
 411 associated with shapes  $\{q_i^* = \hat{O}_i(q_i, \hat{\gamma}_i)\}$ , which can be used to incorporate the con-  
 412 tribution of each shape to the local variation around the spatially-weighted average  
 413 via a weighted covariance matrix.

414 As in Section 2.3, the weighted covariance matrix is computed on the tangent  
 415 space at the spatially-weighted average shape,  $T_{\tilde{q}_0}(\mathcal{S})$ . We project  $\{q_i^*\}$  to  $T_{\tilde{q}_0}(\mathcal{S})$  via  
 416 the inverse-exponential map, resulting in  $\{v_i\}$ . At the implementation stage, each  
 417  $v_i$  is a  $2 \times T$  matrix, which is reshaped to a vector of size  $2T$ , resulting in  $v_i^*$ . The

418 weighted covariance matrix is defined as

$$419 \quad (3.8) \quad \hat{K}_\eta = \frac{n}{n-1} \sum_{i=1}^n |\eta_i| v_i^* v_i^{*\top} \in \mathbb{R}^{2T \times 2T}.$$

420 In this definition, shape  $q_i^*$  is assigned a weight of  $n|\eta_i|$ . The reasoning behind this  
 421 weighting scheme is as follows. First, to ensure that the weighted covariance matrix  
 422 is non-negative definite we require each weight to be positive, but in practice, some  
 423 shapes may be assigned a small negative weight. Hence, we use the absolute value of  
 424 the weights. Second, scaling each  $\eta_i$  by  $n$  ensures that the weights sum to  $n$ , which  
 425 matches the unweighted scenario described in Section 2.3. One can alter the form  
 426 of the weighted covariance matrix to accommodate other weighting schemes, e.g.,  
 427 weights proportional to  $\eta_i^2$ . As in the unweighted case, PCA is performed via SVD of  
 428  $\hat{K}_\eta$ , resulting in spatially-informed principal directions of variation and the variances  
 429 along each direction. Additionally, the overall spatially-weighted shape variance can  
 430 be computed using the sum of the singular values of  $\hat{K}_\eta$ .

431 **3.4. Visualization of Spatial Shape Variation.** In Section 5 that considers  
 432 real data, we first select an ROI within a histopathology image. Motivated by the  
 433 assumption of second-order stationarity and isotropy of the underlying shape random  
 434 field, which is more reasonable on a smaller spatial domain, we further partition  
 435 this region into disjoint, equally sized subregions. The size/number of subregions  
 436 is user and application specific. Spatially-informed summarization of cell nucleus  
 437 shapes is then performed independently within each subregion. We first compute  
 438 a spatially-weighted average shape for each subregion. Because these averages are  
 439 computed independently across subregions, they may exhibit nuisance rotation and  
 440 reparameterization variation. Thus, we additionally optimally register them with  
 441 respect to their shape Karcher mean (Algorithm 2.1); while this step is not necessary  
 442 it allows for easier interpretation of the local shape summaries. We then estimate  
 443 the weighted covariance matrix and perform PCA. Instead of visualizing deformation  
 444 paths along each principal direction, we instead display the deformation vector field  
 445 associated with each direction, and its pointwise magnitude, on the spatially-weighted  
 446 average. This provides concise displays of spatially-informed shape variation in each  
 447 subregion.

448 Let  $U_k$  (reshaped to a matrix of size  $2 \times T$ ) and  $\Sigma_{kk}$  denote the  $k$ th spatially-  
 449 weighted principal direction of shape variation and the variance along this direction,  
 450 respectively, resulting from SVD of  $\hat{K}_\eta$ . We compute  $q_k^1 = \exp_{\tilde{q}_0}(\sqrt{\Sigma_{kk}}U_k)$ , i.e., the  
 451 SRVF that is one standard deviation away from the spatially-weighted shape average  
 452  $\tilde{q}_0$  along the  $k$ th direction. Next, we invert both  $\tilde{q}_0$  and  $q_k^1$  resulting in  $\tilde{\beta}_0 = Q^{-1}(\tilde{q}_0)$   
 453 and  $\beta_k^1 = Q^{-1}(q_k^1)$ , respectively. Since the inverse mapping is only determined up to a  
 454 translation, we additionally center them such that their centroids are at the origin in  
 455  $\mathbb{R}^2$ . Then, the vector field is defined as  $\beta_k^1 - \tilde{\beta}_0$ , i.e., the difference between the centered  
 456 shapes; for display purposes and easy interpretation, the vector field is normalized. In  
 457 addition, we compute the magnitude of the resulting vector field at each point along  
 458 the domain:  $|\beta_k^1(t) - \tilde{\beta}_0(t)| \forall t \in \mathbb{S}^1$ . The vector field and its pointwise magnitude are  
 459 finally displayed on the estimated spatially-weighted average shape  $\tilde{\beta}_0$ .

460 **4. Simulation Studies.** We evaluate the proposed spatially-informed approach  
 461 for summarization of shape variation using two simulation studies. In the first, we  
 462 show that if the underlying shapes are indeed correlated in space, then the spatially-  
 463 weighted shape average is a more accurate estimate of the underlying ground truth

464 shape average than the Karcher mean shape. In the second, we provide qualita-  
 465 tive and quantitative comparisons of spatially-weighted shape variation and variation  
 466 computed under the assumption of independence.

467 In both simulations, we generate dependent shapes as follows. We first define  
 468 the spatial domain  $X = [0, D] \times [0, D]$  and sample  $n$  spatial locations uniformly  
 469 over this domain. We generate spatially correlated shapes at the sampled locations  
 470 using the Fourier basis functions  $\{\cos(lt), \sin(lt), l = 0, 1, 2, t \in [-\pi, \pi]\}$ . Letting  
 471  $\{b_j, j = 1, \dots, 6\}$  denote the basis functions, the radial distance as a function of the  
 472 angle for shape  $i$  is defined as  $\Omega_i(t) = \sum_{j=1}^6 \nu_{ij} b_j(t)$ , where  $\{\nu_{ij}, i = 1, \dots, n, j =$   
 473  $1, \dots, 6\}$  are Fourier coefficients. Each shape is obtained by transforming the radial  
 474 distances to Cartesian coordinates ( $x_i(t) = \Omega'_i(t) \cos(t)$ ,  $y_i(t) = \Omega'_i(t) \sin(t)$ ), where  
 475  $\Omega'_i(t) = \Omega_i(t) + |\min_t \Omega_i(t)| + |\max_t \Omega_i(t)|$  is an adjusted radial function that ensures that  
 476 the resulting shapes are non-self-intersecting. Spatial dependence between the shapes  
 477 is imposed using the Fourier coefficients  $\nu_{ij}$ . Given a mean vector for the coefficients,  
 478  $\mathbf{m} = (m_1, \dots, m_6)^\top$ , we independently sample them for each  $j$  from the multivariate  
 479 normal distribution,  $\text{MVN}(m_j \mathbf{1}_n^\top, C_{mat})$ , where  $\mathbf{1}_n^\top$  is an  $n$ -dimensional vector of ones,  
 480 and  $C_{mat}$  is the Matérn covariance. The Matérn covariance has three parameters. In  
 481 all simulations, we fix the scale parameter to 1 and the smoothness parameter to 0.5.  
 482 The range parameter will depend on the simulation setting.

483 We further add nuisance variation to the simulated shapes. We generate random  
 484 rescalings  $c_i$  independently from the  $\text{Unif}(0.2, 0.6)$  distribution. Random rotations are  
 485 represented by angles  $\phi_i$ , which are sampled independently from the  $\text{Unif}(0, 2\pi)$  distri-  
 486 bution. The resulting spatially dependent shapes, with additional random rotations  
 487 and rescalings, are given by  $\beta_i(t) = (x_i^*(t) = c_i[x_i(t) \cos(\phi_i) - y_i(t) \sin(\phi_i)], y_i^*(t) =$   
 488  $c_i[x_i(t) \sin(\phi_i) + y_i(t) \cos(\phi_i)]$ .

489 **4.1. Comparison of Spatially-weighted Shape Average and Karcher**  
 490 **Mean Shape.** We set the spatial domain to  $X = [0, 2] \times [0, 2]$  and sample a mean  
 491 Fourier coefficient vector  $\mathbf{m} = (m_1, \dots, m_6)$  from  $\text{MVN}(\mathbf{0}_6^\top, 3I_6)$ , where  $\mathbf{0}_6^\top$  is a six-  
 492 dimensional vector of zeros and  $I_6$  is the  $6 \times 6$  identity matrix. We use  $\mathbf{m}$  to define  
 493 the radial distance function for the ground truth mean shape, defined at  $s_0 = (1, 1)$ ,  
 494 and transform to Cartesian coordinates resulting in  $\beta_{gt}$  and its SRVF  $q_{gt}$ . We set the  
 495 sample size to  $n = 50$  and generate spatially dependent shapes as described earlier.  
 496 We consider three settings for the range parameter in the Matérn covariance: 2, 1 and  
 497 0.5. We then use the generated data to estimate the Karcher mean shape  $\hat{\mu}_q$  (under  
 498 assumption of independence) using Algorithm 2.1, and the spatially-weighted shape  
 499 average  $\tilde{q}_0$  using Algorithm 3.1 with  $s_0 = (1, 1)$ . Accuracy of estimation is assessed  
 500 using the shape distance  $d_S([q_{gt}], [\hat{\mu}_q])$  (defined in (2.4)) for the Karcher mean and  
 501  $d_S([q_{gt}], [\tilde{q}_0])$  for the spatially-weighted average. We repeat this process 100 times.

502 Table 1 reports the means and standard deviations of the resulting estimation  
 503 errors. On average, for all values of the range parameter, the spatially-weighted  
 504 shape average is more similar to the ground truth than the Karcher mean shape; the  
 505 standard deviations of estimation errors are also smaller for the spatially-weighted  
 506 average shape. Further, the spatially-weighted shape average becomes more accurate  
 507 in estimating the underlying ground truth shape when the range parameter in the  
 508 data generating process decreases. We also repeated this simulation for  $n = 100$  and  
 509  $n = 200$  (see Section 4 in the supplement for results). In all cases, the spatially-  
 510 weighted shape average was more accurate in estimating the ground truth shape.  
 511 However, the trend of increasing accuracy as the range parameter decreased is not as  
 512 clear in those settings. This is likely due to the data generating process, which results

Range	Method	Mean	SD
0.5	Proposed	0.0947	0.0361
	Karcher Mean	0.1118	0.0488
1	Proposed	0.0982	0.0372
	Karcher Mean	0.1099	0.0521
2	Proposed	0.1035	0.0426
	Karcher Mean	0.1075	0.0447

TABLE 1

Means and standard deviations (SDs) of estimation errors for the spatially-weighted average shape and the Karcher mean shape across 100 replicates.

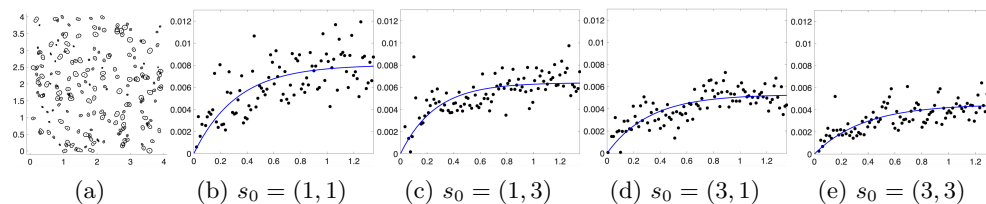


FIG. 3. (a) Simulated spatially dependent shapes. (b)-(e) Empirical shape trace-variograms (black points) and fitted Matérn models (blue curve) for subregions  $X_1$ - $X_4$ , respectively.

513 in more shape heterogeneity with increasing  $n$ . This simulation illustrates the benefit  
 514 of incorporating spatial shape dependence through the shape trace-variogram when  
 515 summarizing spatially correlated shape data.

516 **4.2. Comparison of Independent and Spatially-weighted Shape Varia-**  
 517 **tion.** In this simulation, we fix the spatial domain to  $X = [0, 4] \times [0, 4]$ . To generate  
 518  $n = 200$  spatially correlated shapes, we use  $\mathbf{m} = \mathbf{0}_6^\top$  and a range of 2 in the Matérn  
 519 covariance. To reflect the summarization procedure that will be used for histopathol-  
 520 ogy images, we further partition the spatial domain into four subregions, defined as  
 521  $X_1 = [0, 2] \times [0, 2]$ ,  $X_2 = [0, 2] \times [2, 4]$ ,  $X_3 = [2, 4] \times [0, 2]$  and  $X_4 = [2, 4] \times [2, 4]$ ,  
 522 and perform local summarization based on data within each subregion. For spatially-  
 523 informed summarization, the location  $s_0$  is chosen as  $(1, 1)$ ,  $(1, 3)$ ,  $(3, 1)$  and  $(3, 3)$   
 524 for each subregion, respectively.

525 The data is shown in Figure 3(a). In Figure 3(b)-(e), we show the estimated  
 526 shape trace-variograms within each subregion obtained via Algorithm 3.1. The em-  
 527 pirical variogram is shown as black points with the Matérn model fit, with smoothness  
 528 parameter of 0.5, in blue. Based on the structure of these trace-variograms, it is clear  
 529 that the shapes within each subregion are spatially dependent, i.e., there is less vari-  
 530 ation at smaller spatial distance lags; this is especially clear in regions  $X_1$  and  $X_2$   
 531 (panels (b) and (c)).

532 Figure 4 compares local spatially-informed shape summarization (panel (a)), as  
 533 described in Section 3, to local summarization under independence (panel (b)), as  
 534 described in Section 2.3. The columns show variation along the first two principal  
 535 directions of shape variation. First, there is a clear difference in the estimated average  
 536 shapes across the four subregions. The spatially-weighted shape averages appear to  
 537 have more geometric features than the Karcher mean shapes, especially in subregions  
 538  $X_1$  and  $X_2$ . Second, the estimated patterns of variation are markedly different across

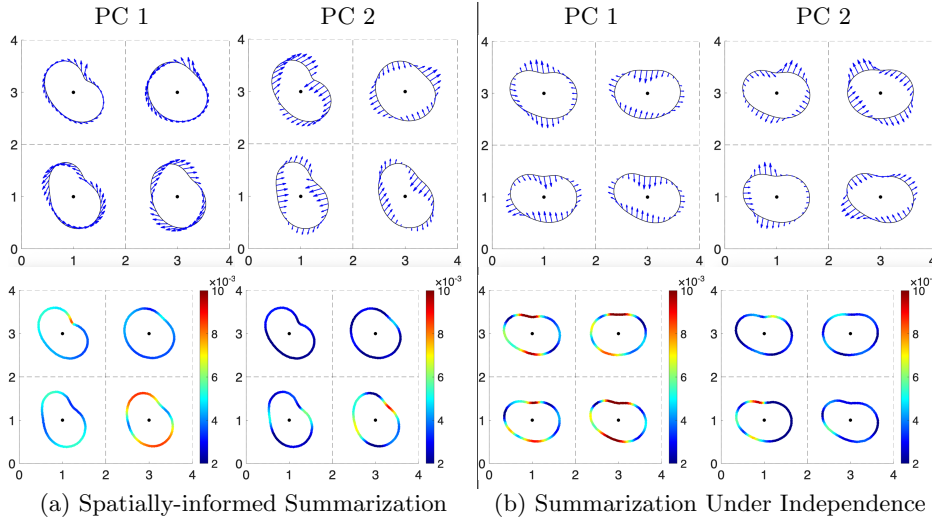


FIG. 4. (a) *Spatially-informed summarization of shape variation.* (b) *Summarization of shape variation under independence.* Columns correspond to the first and second principal directions of shape variation. We show variation as a vector field (top row), and its pointwise magnitude (bottom row), on the estimated average.

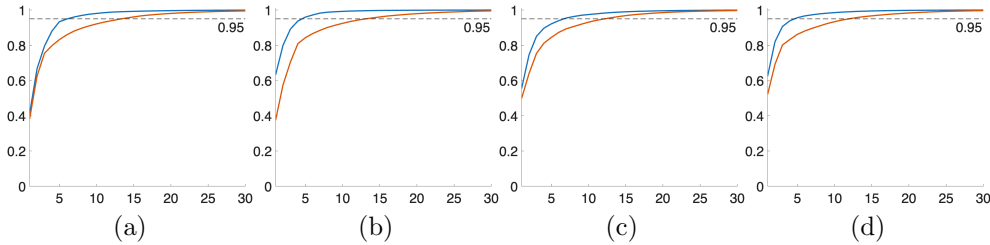


FIG. 5. *Cumulative explained variance as a function of the number of PCs based on spatially-weighted PCA (blue) and unweighted PCA (red) for subregions (a)  $X_1$ , (b)  $X_2$ , (c)  $X_3$ , and (d)  $X_4$ .*

539 the two summarization approaches as evident through the vector fields (top row)  
 540 and their pointwise magnitudes (bottom row). Finally, PCA carried out using the  
 541 spatially-weighted covariance captures primary variation more efficiently than PCA  
 542 carried out on the covariance matrix computed without spatial weighting. In partic-  
 543 ular, for regions  $X_1$ - $X_4$ , 7, 5, 7 and 5 spatially-weighted PCs explain at least 95% of  
 544 shape variation, respectively. For unweighted PCA, in the same four subregions, 14,  
 545 14, 13 and 12 shape PCs are needed to explain at least 95% of the variation. This  
 546 trend is consistent for other variation thresholds as seen in Figure 5. For all four  
 547 subregions, the blue cumulative explained variance curve, corresponding to spatially-  
 548 weighted PCA, is above the red one, which corresponds to unweighted PCA.

549 **5. Application to Histopathology Images.** We apply our framework to cell  
 550 nucleus outlines extracted from whole slide images (WSIs) of breast cancer tissue. The  
 551 images were obtained from the TiGER challenge [30], which focused on HER2-positive  
 552 and triple-negative breast cancers; both types are linked to notably poor prognosis.  
 553 We first selected two images from this database; these are shown in Figure 6(a),  
 554 where the tumor tissue is outlined in black. For each image, we selected two ROIs

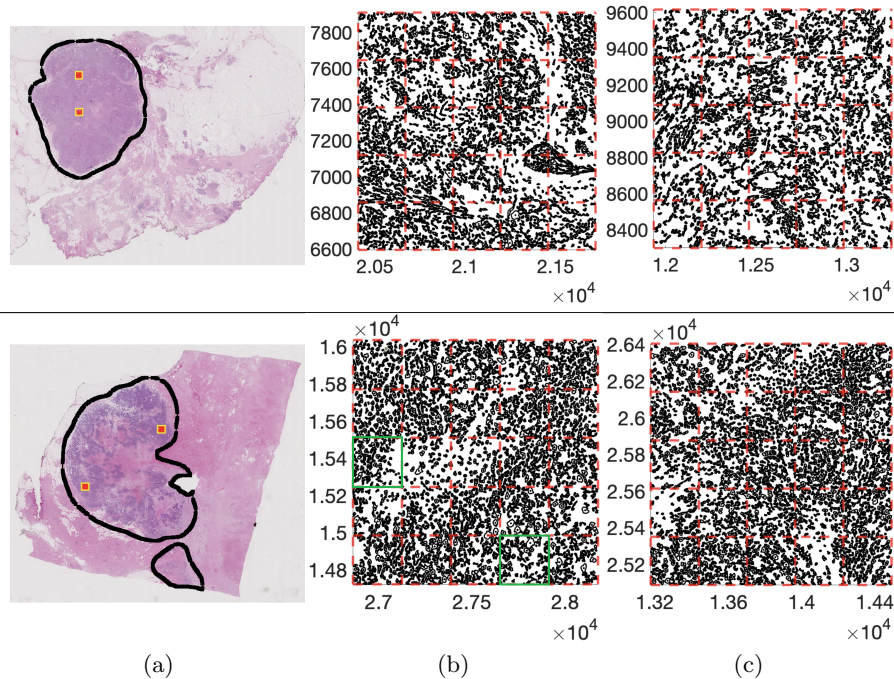


FIG. 6. (a) Breast cancer tissue images; tumor outlined in black. (b)&(c) Cell nucleus outlines in ROIs (red squares) from (a); subregions are denoted by red dashed lines. The two subregions highlighted in green in the bottom row in (b) are visualized in more detail in Figure 9(a).

555 within tumor tissue, shown as red squares in (a). Using QuPath [1], we segmented cell  
 556 nucleus outlines from these ROIs. The extracted outlines are displayed in (b) and (c)  
 557 for each of the two images and ROIs. Each ROI was subsequently partitioned into 25  
 558 equally sized subregions for local spatially-informed summarization and visualization  
 559 of cell nucleus shape variation. The two subregions highlighted in green in the bottom  
 560 row of Figure 6(b) are displayed in more detail in Figure 9(a).

561 The empirical shape trace-variograms within each subregion exhibited structure  
 562 that led us to fit a Matérn model with a smoothness parameter of 0.2. Depending  
 563 on the application and spatial dependence pattern, the smoothness parameter should  
 564 be tuned to obtain a satisfactory Matérn fit to the empirical shape trace-variogram.  
 565 However, as we show in Section 4 in the supplement, spatially-informed summarization  
 566 is not sensitive to the choice of the smoothness parameter, or the density of cell nucleus  
 567 shapes in a fixed spatial subregion. Figures 7 and 8 show the resulting local spatially-  
 568 informed summarizations of cell nucleus shape variation. We display the first principal  
 569 direction of shape variation in Figure 7; the second direction is shown in Figure 8.  
 570 Panels (a)-(d) consider the ROIs shown in the top row of Figure 6(b)&(c) and bottom  
 571 row of Figure 6(b)&(c), respectively. The results highlighted in green in the top row  
 572 of Figure 7(c) are shown in more detail in Figure 9(b). Clearly, the spatially-weighted  
 573 shape averages can be quite different across subregions within the same ROI, i.e.,  
 574 there appears to be quite a bit of spatial cell nucleus shape heterogeneity. This is also  
 575 true for the principal directions of shape variation. Certain subregions exhibit much  
 576 less variation than others. The proposed visualizations represent practical exploratory  
 577 tools for clinicians to assess cell nucleus shape variation in histopathology images. In

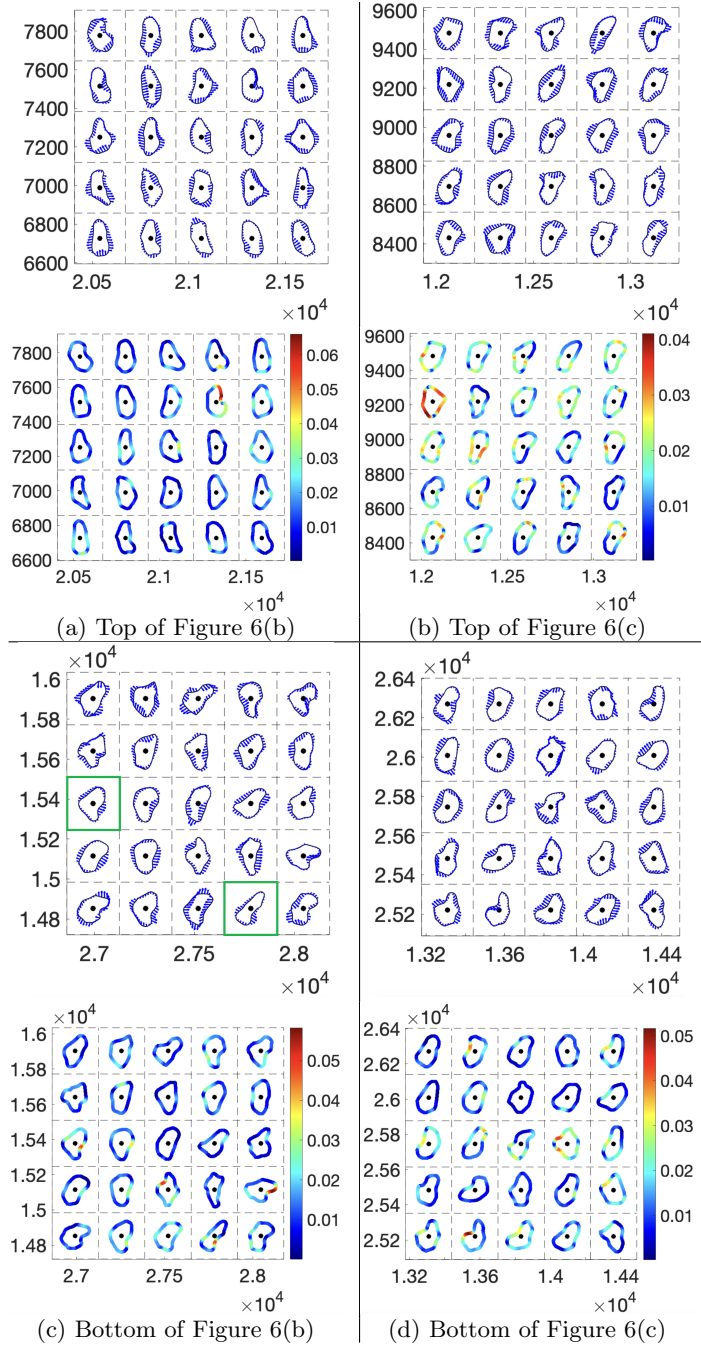


FIG. 7. Spatially-informed summarization of cell nucleus shape variation. Columns correspond to the first principal direction of shape variation for ROIs in top and bottom rows of Figure 6(b)&(c). We show variation as a vector field (top), and its pointwise magnitude (bottom), on the spatially-weighted average shape. The two results highlighted in green in the top row in (c) are visualized in more detail in Figure 9(b).

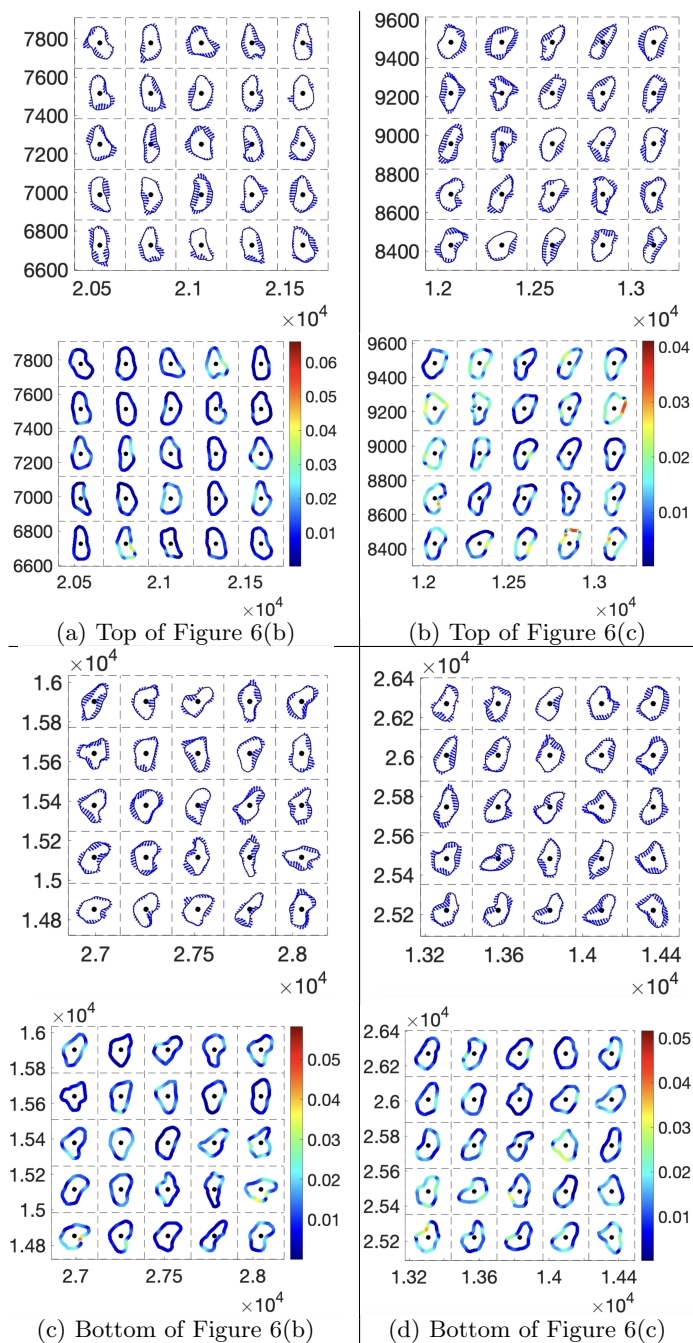


FIG. 8. Spatially-informed summarization of cell nucleus shape variation. Columns correspond to the second principal direction of shape variation for ROIs in top and bottom rows of Figure 6(b) & (c). We show variation as a vector field (top), and its pointwise magnitude (bottom), on the spatially-informed average shape.

578 particular, the estimated averages and principal directions of variation can be used to  
 579 identify heterogeneous tissue properties within different ROIs. Importantly, in future

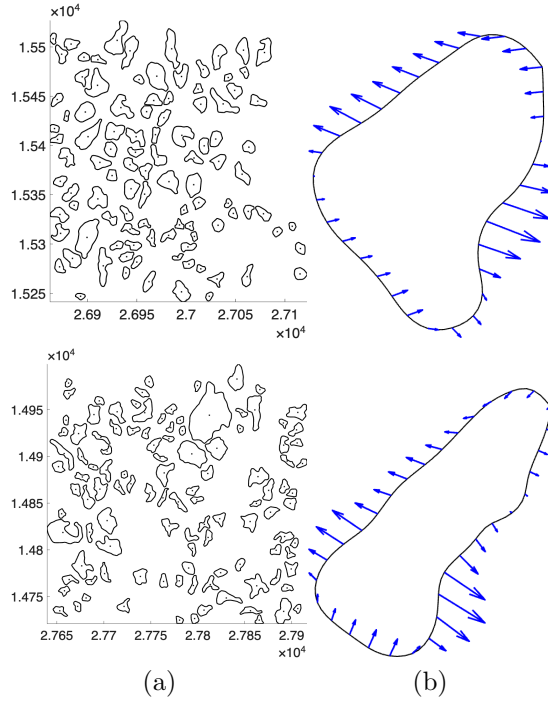


FIG. 9. (a) Zoomed-in visualizations of the subregions highlighted in green in the bottom row of Figure 6(b). (b) Detailed visualizations of the corresponding spatially-weighted shape averages with the first principal directions of shape variation shown as vector fields.

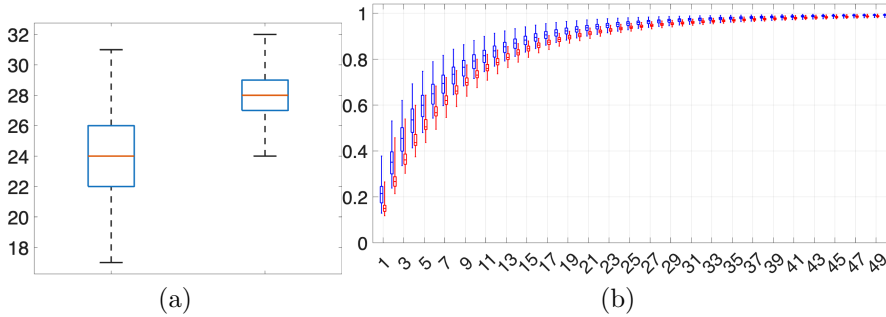


FIG. 10. (a) Boxplots of the number of PCs required to explain at least 95% of the total shape variance across 100 tumor tissue subregions. Left: Spatially-weighted PCA. Right: Unweighted PCA. (b) Cumulative explained variance as a function of the number of PCs, aggregated over the same 100 tumor tissue subregions: spatially-weighted in blue and unweighted in red.

580 studies, such summaries along with further visual aides can be used to characterize  
 581 subtypes and severity of tumors. Additional results on real data for different ROIs  
 582 are provided in Section 5 in the supplement.

583 **Comparison of spatially-weighted PCA and unweighted PCA.** We next  
 584 compare efficiency of spatially-weighted PCA and unweighted PCA, in terms of cap-  
 585 turing cell nucleus shape variation, based on all 100 subregions (four ROIs shown  
 586 in Figure 6, each split into 25 subregions). Within each subregion, we carried out  
 587 spatially-weighted PCA and unweighted PCA. Figure 10(a) presents boxplots of the

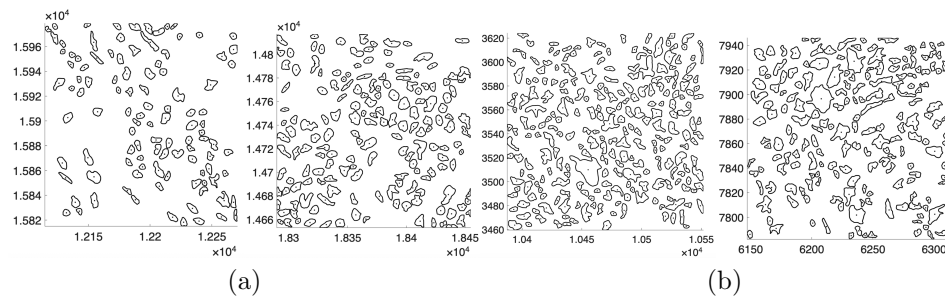


FIG. 11. Illustration of two randomly sampled regions within (a) normal and (b) tumor tissues.

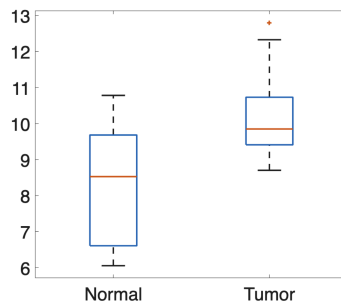


FIG. 12. Boxplots of spatially-weighted variance of cell nucleus shapes within ten randomly sampled normal and tumor tissue regions of the same size.

588 number of PCs required by spatially-weighted PCA (left) and unweighted PCA (right)  
 589 to capture at least 95% of total shape variation. The proposed method consistently  
 590 requires fewer PCs to reach the 95% threshold, indicating a more compact representa-  
 591 tion of shape variability. For 89 of the subregions, spatially-weighted PCA resulted in  
 592 fewer PCs required to capture at least 95% of the variation; for six of the subregions,  
 593 both PCA approaches resulted in an equal number of PCs required to capture at least  
 594 95% of the variation.

595 Figure 10(b) further displays cumulative explained variance curves (as boxplots)  
 596 aggregated across the 100 subregions for both methods (spatially-weighted in blue  
 597 and unweighted in red). The boxplots corresponding to the proposed approach rise  
 598 more rapidly as a function of the number of PCs compared to unweighted PCA. This  
 599 demonstrates that spatial weighting results in a larger proportion of shape variability  
 600 being captured by the leading PCs. Together, these results show that the spatially-  
 601 weighted covariance is more efficient in capturing shape variation than the unweighted  
 602 one, and thus has the potential to provide clearer insight into regional cell nucleus  
 603 shape heterogeneity.

604 **Comparison of cell nucleus shape heterogeneity within tumor and nor-**  
 605 **mal tissue regions.** Clinically, it is well known that tumor cells are more variable  
 606 than healthy cells due to their rapid proliferation and mutation. To confirm this,  
 607 we use the proposed spatially-informed summarization framework to compare overall  
 608 shape variation across different regions sampled within tumor and normal tissues.  
 609 In particular, we randomly sampled ten small regions within tumor tissue and ten  
 610 size matched regions within normal tissue; see Figure 11 for a visualization of some  
 611 of the sampled regions. We then computed the total spatially-weighted shape vari-

612 ance in each of the regions. Figure 12 displays boxplots of the variances. It is clear  
613 that spatially-weighted cell nucleus shape variation is larger in tumor tissue than in  
614 normal tissue. The average variance within tumor regions is 10.25, whereas the average  
615 variance within normal regions is 8.36. This shows that, as expected, cell nuclei  
616 exhibit substantially greater morphological heterogeneity within the tumor as compared  
617 to normal tissue, which provides quantitative support for increased structural  
618 heterogeneity in tumor microenvironments.

619 **6. Discussion and Future Work.** We introduced a novel framework for local  
620 spatially-informed summarization and visualization of planar curve shapes. By leveraging  
621 tools from elastic shape analysis and incorporating spatial dependence via a  
622 shape trace-variogram, the method estimates a spatially-weighted shape average and  
623 covariance matrix, which can be used to explore spatially-informed shape variation via  
624 PCA. Our approach addresses limitations of traditional global summaries by accounting  
625 for spatial correlation among shapes, which is especially pertinent in assessing  
626 heterogeneity of cell nucleus shapes in histopathology images in the context of cancer.  
627 Simulation studies demonstrate that the spatially-weighted shape average is a more  
628 accurate estimator of a ground truth shape average than the conventional Karcher  
629 mean shape when observed shapes exhibit spatial dependence. Further, spatially-  
630 weighted PCA is more efficient in capturing primary modes of shape variation.

631 We have identified a few directions for future work. First, the natural next step  
632 is to generalize the proposed framework to the case of 3D shapes. In many medical  
633 imaging applications, the outlines of structures of interest are surfaces rather than  
634 curves. Thus, we will define an appropriate analogue of the shape trace-variogram for  
635 the case of surfaces; a key aspect of this is to determine how to account for all relevant  
636 nuisance transformations. Additionally, we will develop novel visualization techniques  
637 for summaries of shape variation in this case. This advance can significantly broaden  
638 the framework’s applicability, particularly in fields where 3D morphological analysis  
639 is critical.

640 Second, the current approach relies on the assumption of isotropy, i.e., that spatial  
641 dependence is uniform in every direction. However, in many practical scenarios,  
642 strength of spatial correlation may not be isotropic. We will explore new models  
643 to account for directional dependence, potentially leading to better estimates of the  
644 spatially-weighted shape average and associated shape variation.

645 Third, the proposed framework provides point estimates for the spatially-weighted  
646 shape average and PCA. In many contexts, uncertainty associated with these estimates  
647 is also of interest. While a bootstrap procedure can be utilized to quantify  
648 uncertainty in the independent case, it does not apply in the current setting. Thus,  
649 we will explore alternative approaches to assess uncertainty when shape data are  
650 spatially dependent.

651 Finally, to translate this framework into clinical practice, we will study the correlation  
652 between the spatially-informed shape summaries and clinical outcomes, e.g.,  
653 tumor grade or survival, to evaluate how well these shape summaries perform as  
654 biomarkers in the clinical setting, potentially improving personalized treatment planning  
655 and disease monitoring. In particular, breast cancer grading is based on the  
656 Nottingham Histopathology Grading system, which includes nuclear pleomorphism  
657 as a key component. Nuclear pleomorphism refers to variation in the size, shape  
658 and appearance of cell nuclei. As such, the presented methodology can be used to  
659 objectively quantify nuclear pleomorphism to aid in tumor grading. Currently, most  
660 studies use a potentially small number of shape features to quantify morphological

661 irregularity of cell nuclei; see, e.g., [29]. However, this can result in significant loss  
 662 of information, and thus provide misleading results. In contrast, the proposed frame-  
 663 work offers spatially-weighted measures of variation estimated from the full geometric  
 664 shape of cell nuclei.

665 **Authorship and Contributorship Statement.** All authors have made substantial  
 666 intellectual contributions to the study conception, execution, and design of the work.  
 667 All authors have read and approved the final manuscript. In addition, the following  
 668 contributions occurred: Conceptualization: Ye Jin Choi, Sebastian Kurtek, Karthik  
 669 Bharath; Methodology: Ye Jin Choi, Sebastian Kurtek, Karthik Bharath; Formal  
 670 analysis and investigation: Ye Jin Choi; Writing - original draft preparation: Ye Jin  
 671 Choi, Sebastian Kurtek, Karthik Bharath; Writing - review and editing: Ye Jin Choi,  
 672 Sebastian Kurtek, Karthik Bharath; Funding acquisition: Sebastian Kurtek, Karthik  
 673 Bharath; Supervision: Sebastian Kurtek, Karthik Bharath.

674 **Conflicts of Interest.** The authors declare that there are no conflicts of interest.

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