Exploring biological shape analysis through topology, geometry and statistics

Ph. D. summer school: Biomedical image analysis, 2024/08/13

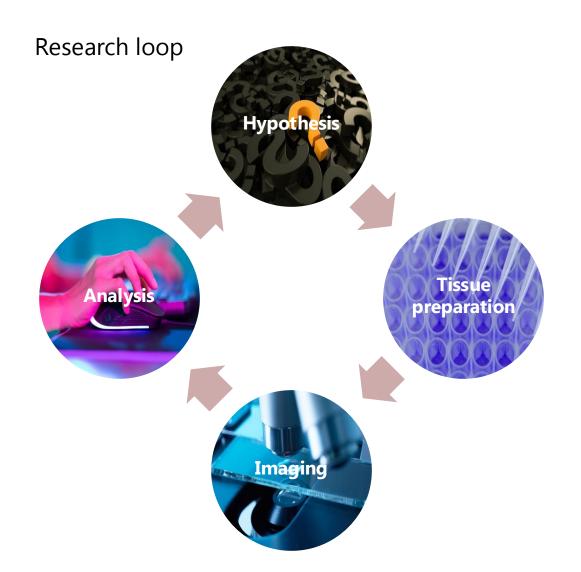
Jon Sporring,
Department of Computer Science

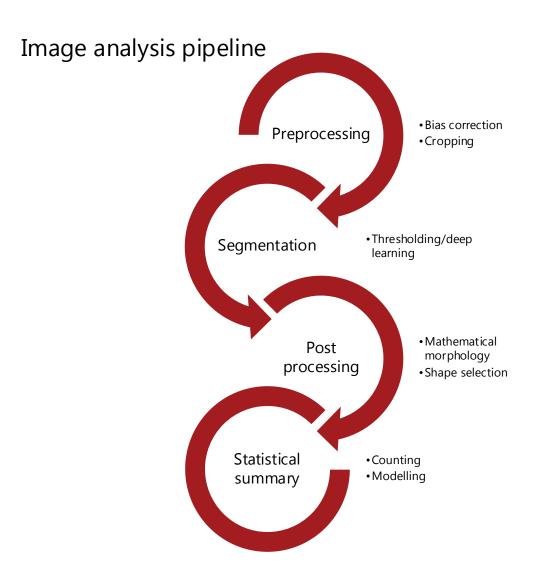


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Imaging research and pipeline





AI is pretty good at segmenting stuff, what's next?

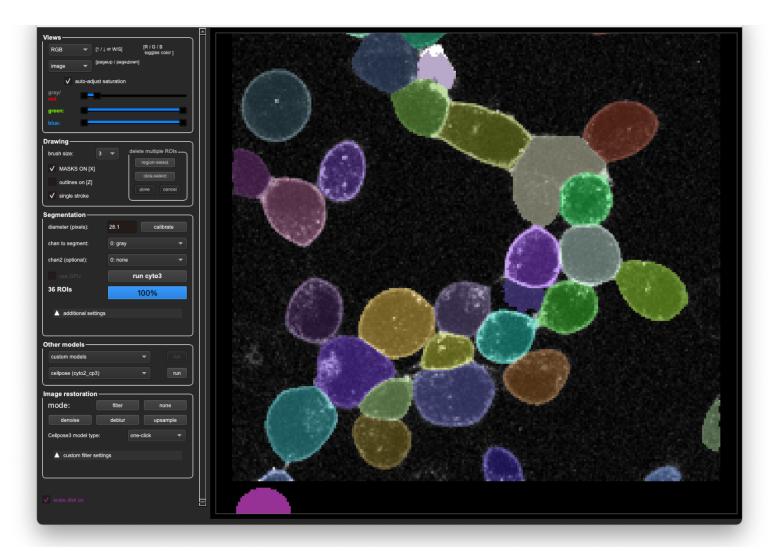
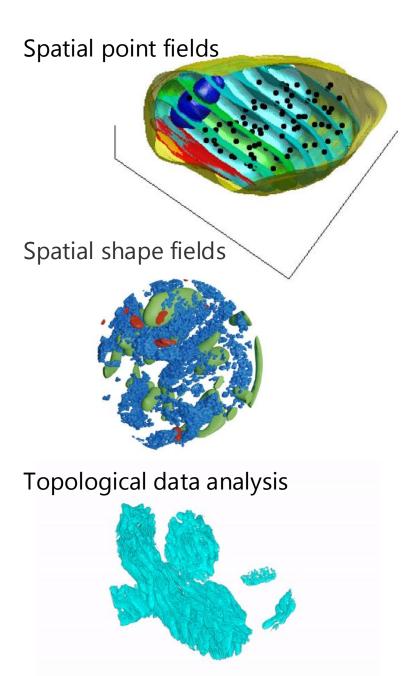


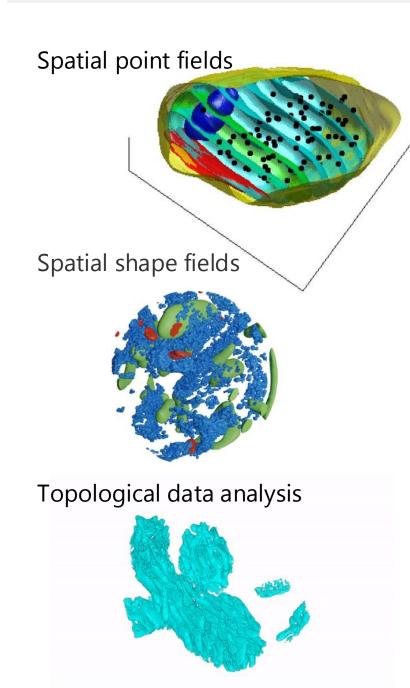
Image courtesy: Karen Martinez & Gabriella von Scheel von Rosing; AI: http://www.cellpose.org/

What to do next: Shape analysis

Focused ion-beam scanning electron microsopy (FIB-SEM) Voxel size: (5 nm)³ Mitochondria Vessicles Active zone



Graham Knott and Marco Cantoni. Electron microscopy dataset. https://cvlab.epfl.ch/data/data-em/



Literature

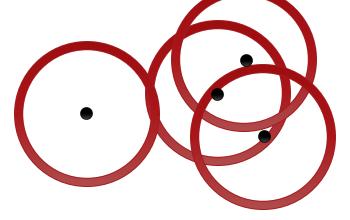
- Analysis of shape and spatial interaction of synaptic vesicles using data from focused ion beam scanning electron microscopy (FIB-SEM); M Khanmohammadi, RP Waagepetersen & J Sporring, Frontiers in Neuroanatomy, 2015
- Stoyan, D. (2006). Fundamentals of Point Process Statistics. In: Baddeley, A. et al. (eds) Case Studies in Spatial Point Process Modeling. Lecture Notes in Statistics, vol 185. Springer
- Mrkvička, Tomáš, et al. "A one-way ANOVA test for functional data with graphical interpretation." Kybernetika 56.3 (2020): 432-458.
- Stephensen, H.J.T., Svane, A.M., Villanueva, C.B. et al. Measuring Shape Relations Using r-Parallel Sets. J Math Imaging Vis, vol 63, 2021.

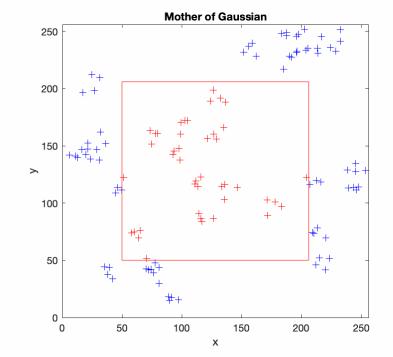
 Chazal F., Michel B., An Introduction to Topological Data Analysis: Fundamental and Practical Aspects for Data Scientists, In: Frontiers in Artificial Intelligence, vol 4, 2021

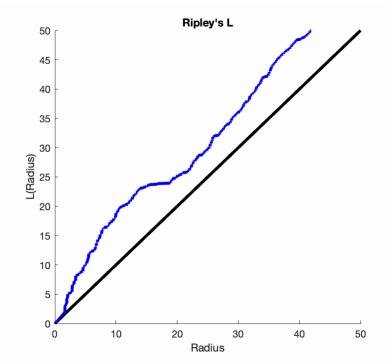
Analysis of shape and spatial interaction of synaptic vesicles using data from focused ion beam scanning electron microscopy (FIB-SEM); M Khanmohammadi, RP Waagepetersen & J Sporring, Frontiers in Neuroanatomy, 2015

Ripley's K- and L-functions: expected number of neighboring points by radius

$$K(r) = \frac{1}{\lambda} \mathbb{E}[I(d_{ij} < r)]$$
$$L = \sqrt{\frac{K}{\pi}}$$





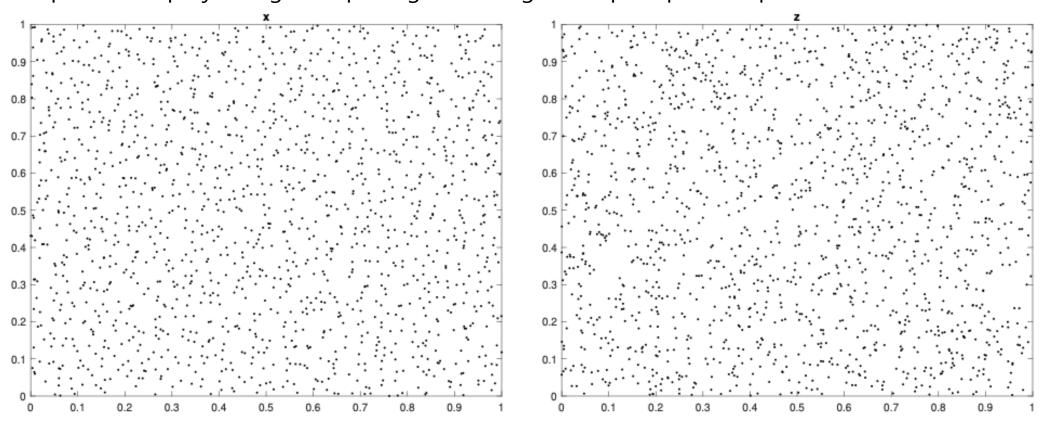


R and rpy2 demo: https://sporring.github.io/bia2024/ demoRpy2.py

https://cran.r-project.org/

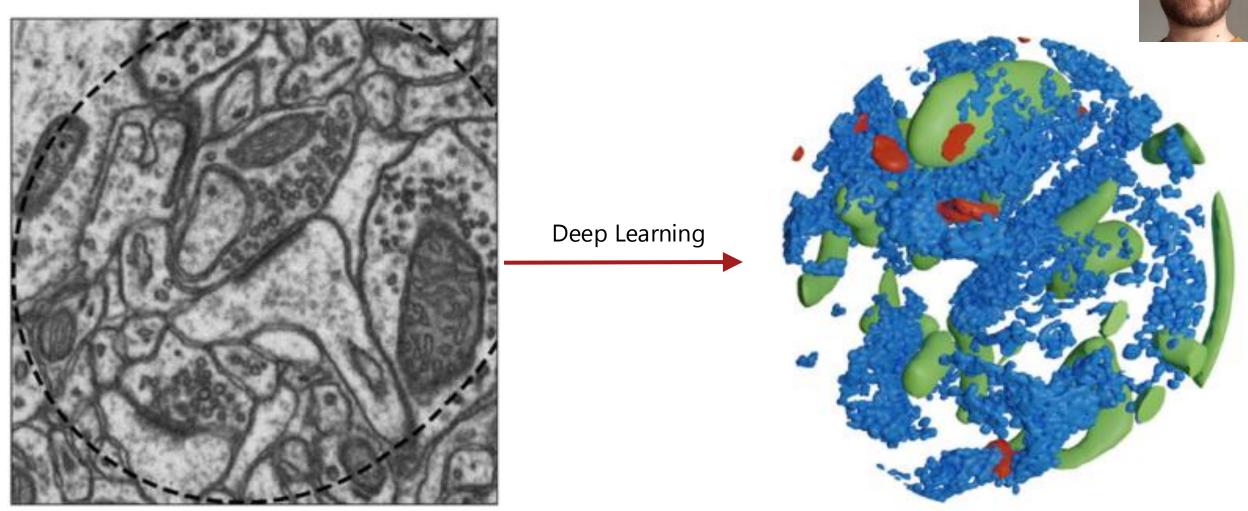
https://spatstat.org/

https://cran.r-project.org/web/packages/GET/vignettes/pointpatterns.pdf



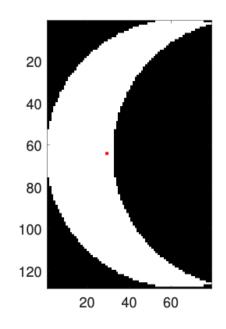
Exercise: Use GET's global_envelope_test to test whether x and/or z are likely to be random

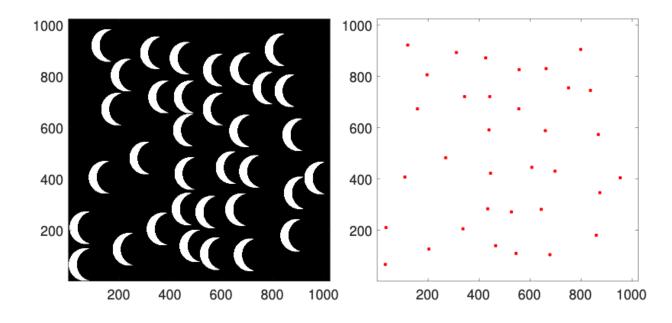
2. Spatial shape fields: Real structures are not points, small structures are difficult to separate

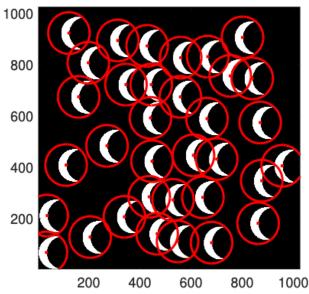


Measuring Shape Relations Using r-Parallel Sets; HJT Stephensen, AM Svane, CB Villanueva, SA Goldman, & J Sporring; Journal of mathematical imaging and vision, 2021

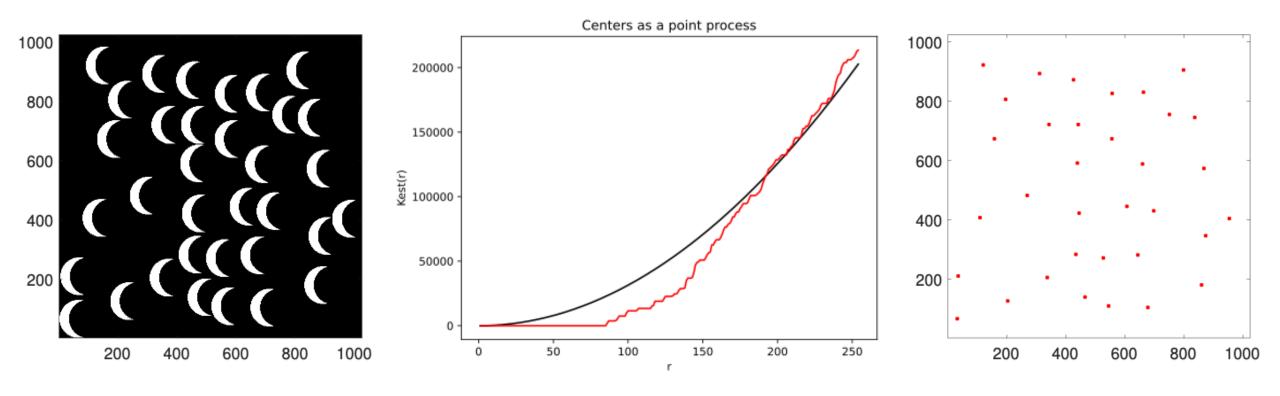
Not all shapes are well summarized as a reference point



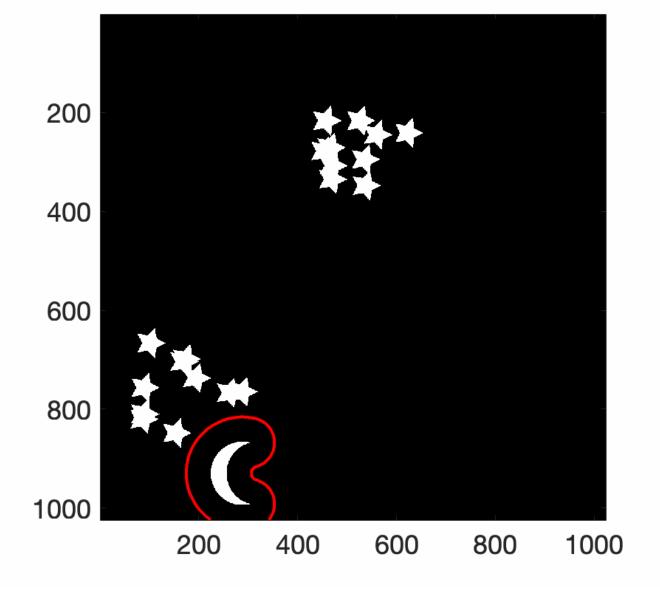




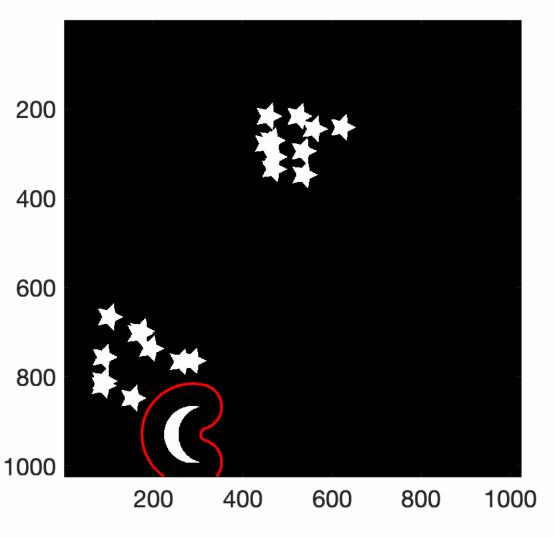
Ripley's K-function indicates strong repulsion

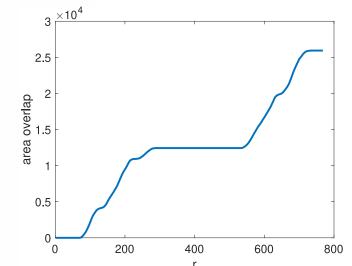


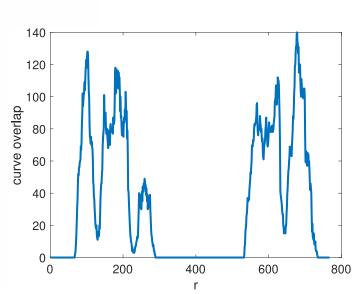
Shape relation measures = Interaction with distance fields



Shape relation measures: Some characteristic functions



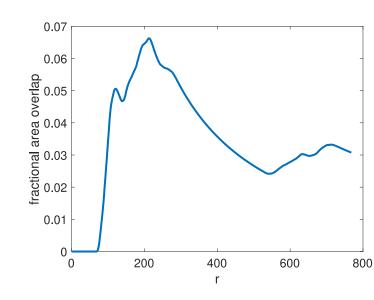




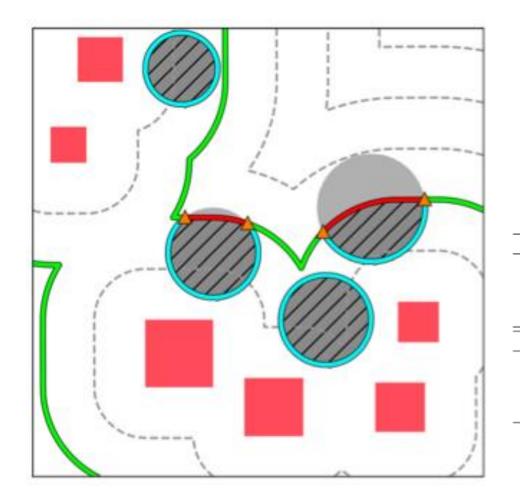
$$\mu_{00}(r) = \mathcal{H}(X \cap Y^r)$$

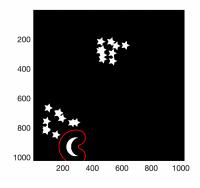
$$g_{00}(r) = \frac{d\mu_{00}(r)}{dr} = \mu_{01}(r)$$

$$f_{00}(r) = \frac{\mu_{00}(r)}{\mathcal{H}(Y^r)}$$



There can be only 4





$$Y^r = \{ \alpha \in \mathbb{R}^d \mid \inf_{y \in Y} d(\alpha, y) \le r \}$$
 (1)

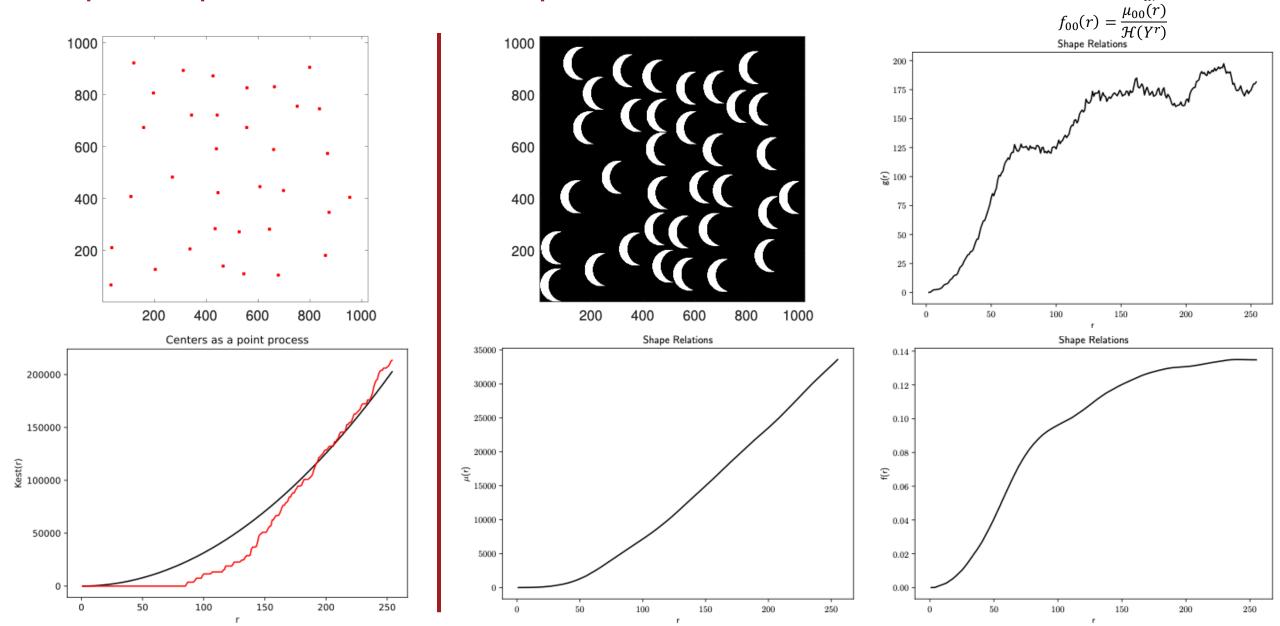
$$\mu_{\varepsilon,\varepsilon'}(X,Y^r) = \mathcal{H}^{d-\varepsilon-\varepsilon'}(\partial^{\varepsilon}X \cap \partial^{\varepsilon'}Y^r) . \tag{2}$$

d=2	$(\varepsilon, \varepsilon')$	$\mathcal{H}^{d-arepsilon-arepsilon'}$	$\partial^{\varepsilon} X \cap \partial^{\varepsilon'} Y^r$	Interpretation of $\mu_{\varepsilon,\varepsilon'}(X,Y^r)$
////	(0,0)	Area	$X \cap Y^r$	Area of intersection
	(0, 1)	Curve length	$X \cap \partial Y^r$	Boundary length of intersection inside interior of X
	(1, 0)	Curve length	$\partial X \cap Y^r$	Boundary length of intersection inside boundary of X
_	(1, 1)	Point counts	$\partial X \cap \partial Y^r$	Number of points in intersection of boundaries
d = 3	$(\varepsilon, \varepsilon')$	$\mathcal{H}^{d-arepsilon-arepsilon'}$	$\partial^{\varepsilon} X \cap \partial^{\varepsilon'} Y^r$	Interpretation of $\mu_{\varepsilon,\varepsilon'}(X,Y^r)$
	(0, 0)	Volume	$X \cap Y^r$	Volume of intersection
	$(0,0) \\ (0,1)$	Volume Surface area	$X \cap Y^r$ $X \cap \partial Y^r$	Volume of intersection Surface area of intersection inside interior of X
	` ' /			
	(0, 1)	Surface area	$X\cap \partial Y^r$	Surface area of intersection inside interior of X

 $\mu_{00}(r)=\mathcal{H}(X\cap Y^r)$

• Chirphori or Corphinadin

Spatial point fields vs. shape fields

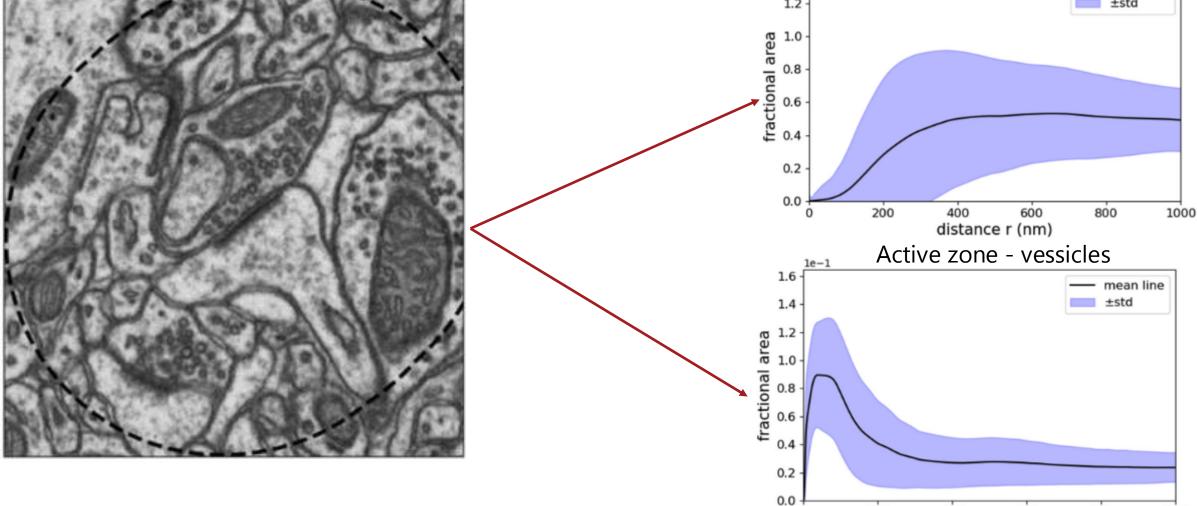


800

1000

Shape relations for statistical summary of families of shapes and their relations

Active zone - mitochondria



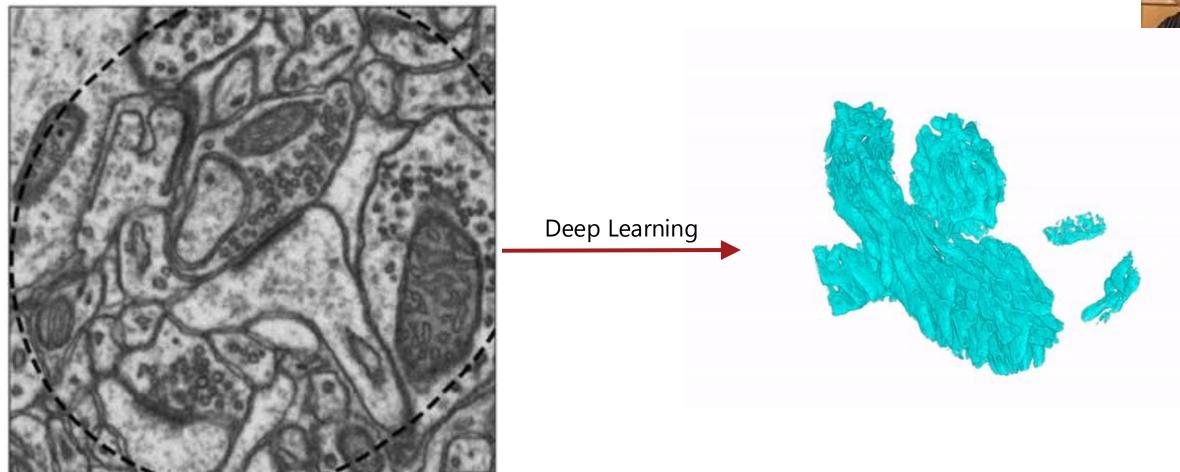
200

400

distance r (nm)

3. With a little help from topology: Cristae membranes in Mitochondria



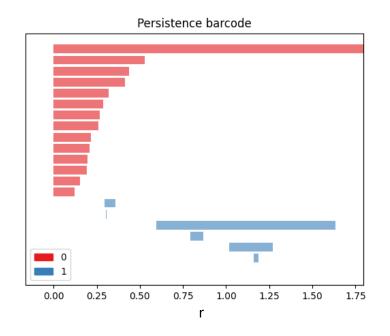


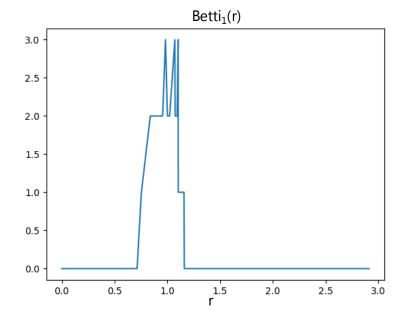
Extracting Mitochondrial Cristae Characteristics from 3D Focused Ion Beam Scanning Electron Microscopy Data, C Wang, L Østergaard, S Hasselholt, & J Sporring, to appear in Communications Biology, 2024

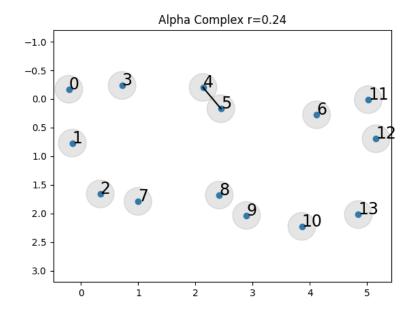
Persistent homology and friends

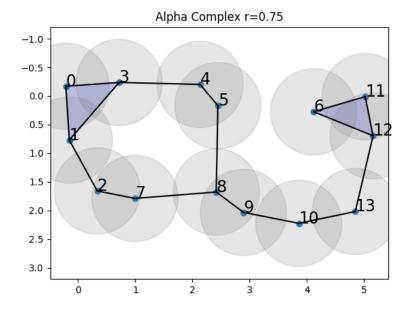
Simplex $\sigma = [x_0, x_1, ... x_k]$ is in the alpha complex if

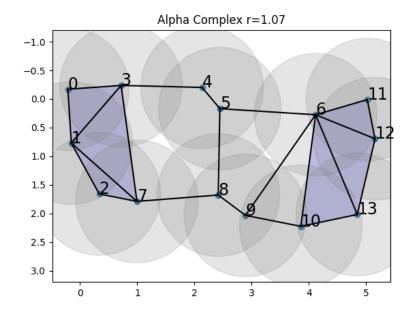
$$\bigcap_{x_i \in \sigma} B(x_i, r) \neq \emptyset$$





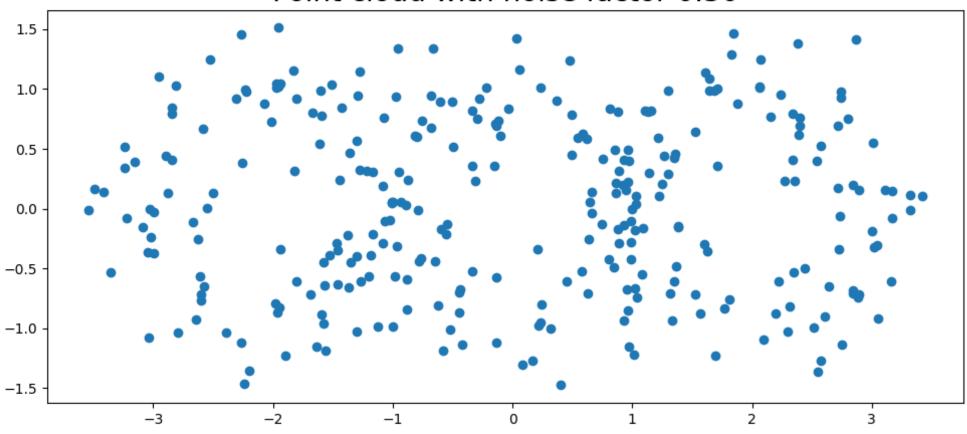




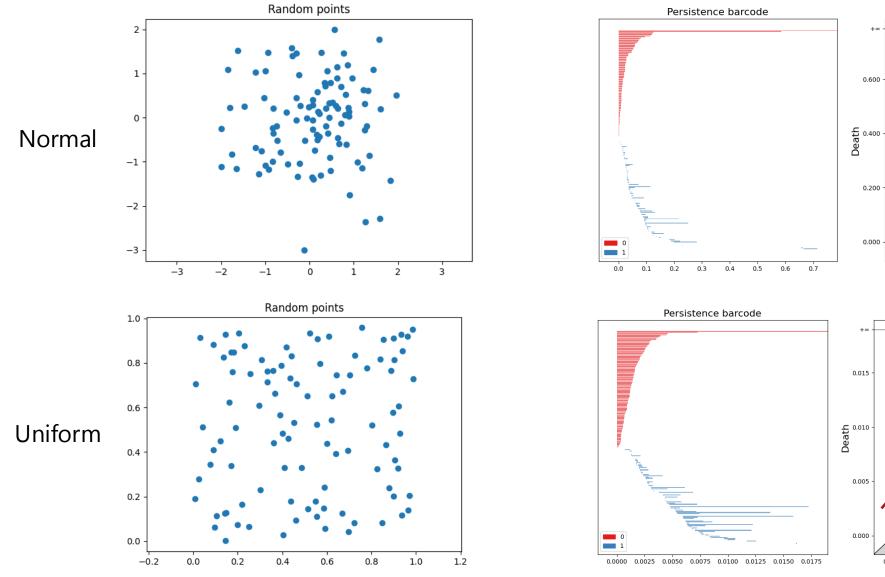


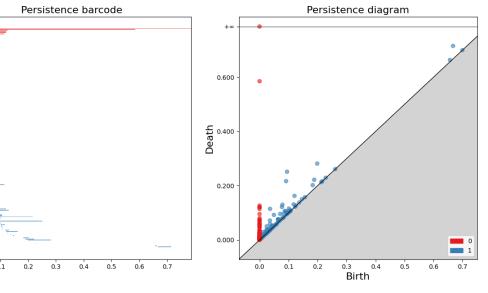
Persistence under noise: $dgm = dgm^{Signal} \cup dgm^{Noise}$

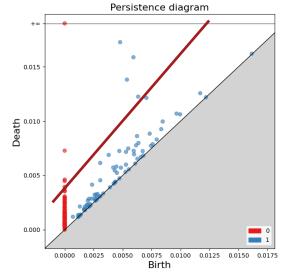




Pure noise







Bobrowski & Skraba, "A universal null-distribution for topological data analysis", Nature/Scientific Reports, 2023

Random points:

$$x \in S(d)$$
, $x \sim f$, $p = (r_{birth}, r_{death})$

Left-skewed Gumbel distribution:

$$F(x) = 1 - e^{-e^x}$$
, $f(x) = e^{x - e^x}$, $\mu = -\gamma = -0.57721$, $\sigma^2 = \frac{\pi^2}{6}$

Transformation:

$$\rho = \ln \ln \frac{r_{\text{death}}}{r_{\text{birth}}}$$

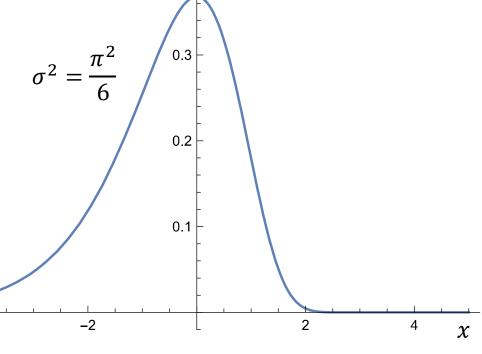
$$x = \frac{(\rho - \bar{\rho})}{\beta} - \gamma, \qquad \bar{\rho} = \frac{1}{|\text{dgm}_k|} \sum_{n \in \text{dgm}_k} \rho, \qquad \beta = \begin{cases} 1, \text{Rips} \\ 2, \text{Čech} \end{cases}$$

Bonferroni testing (family-wise error rate $< \alpha$):

$$P(x \ge x_0 | x \text{ is noise}) = 1 - F(x) = e^{-e^{x_0}}$$

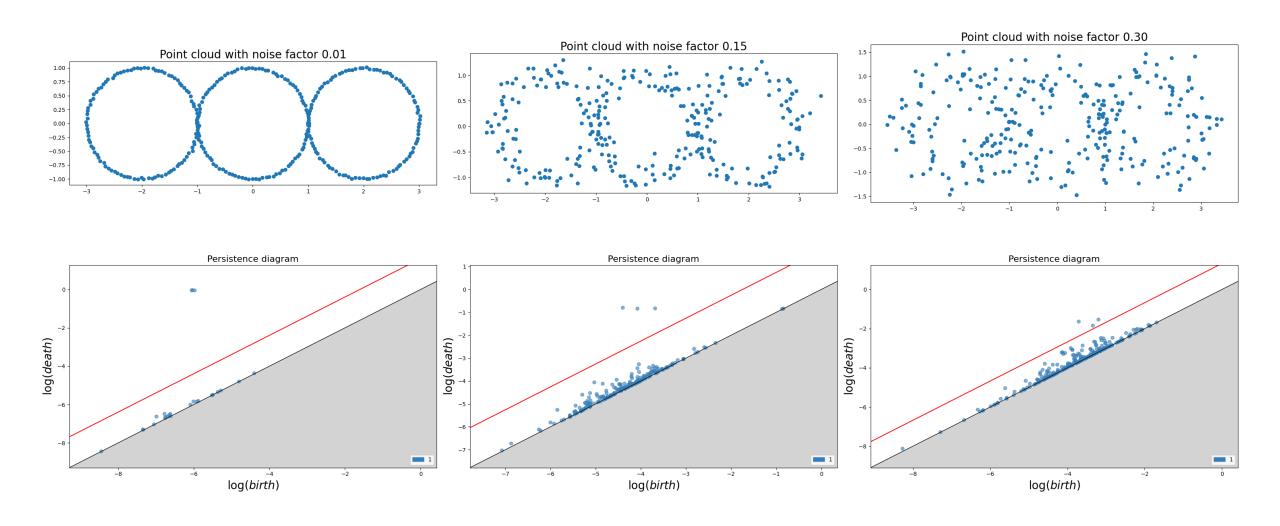
$$\operatorname{dgm}_k^{\text{Signal}}(\alpha) = \left\{ p \in \operatorname{dgm}_k : e^{-e^x} < \frac{\alpha}{|\operatorname{dgm}_k|} \right\}$$

$$e^{\rho} = \ln \frac{r_{\text{death}}}{r_{\text{birth}}} = (-1)^{\beta} e^{\beta \gamma + \overline{\rho}} \left(\ln \frac{\alpha}{|\operatorname{dgm}_k|} \right)^{\beta}$$

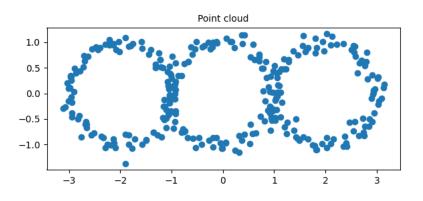


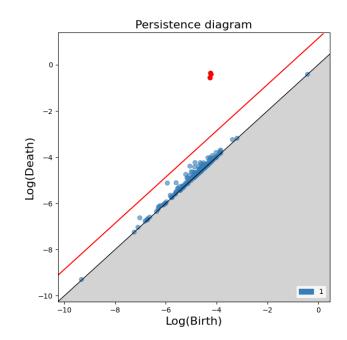
f(x)

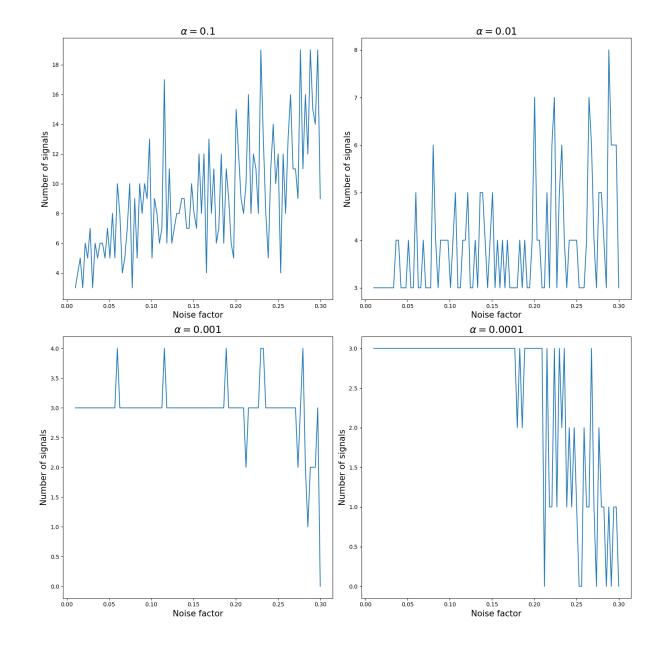
The universal distribution can separate very noisy cases



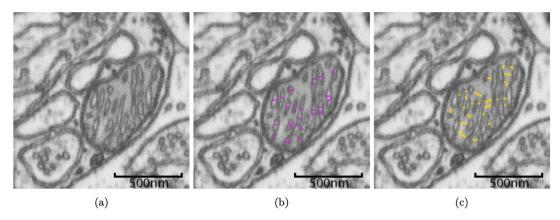
Family-wise error rate

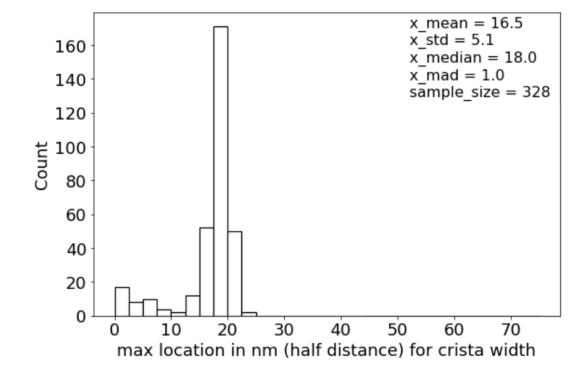


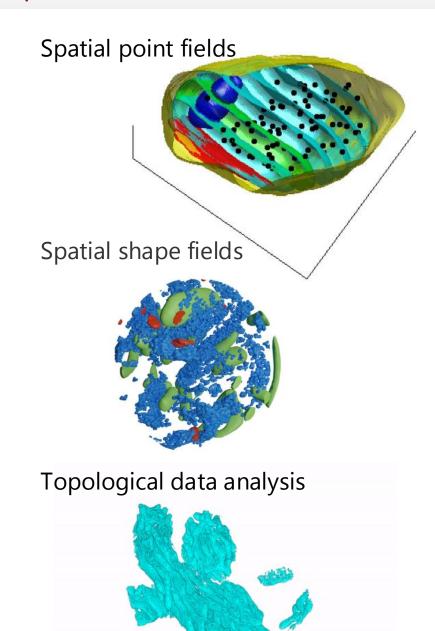




Persistent homology: Statistical measures on H_0







Statistical summary of object collections

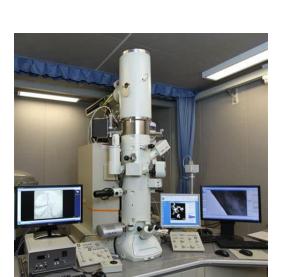
Pair correlation and Ripley's K functions summarizes 1st order point relations – e.g., do the vessicles cluster?

Hausdorf measures on overlaping sets extends notion of points to shapes – e.g., are mitochondria seen close to the synapse?

Filtrations brings topological concepts to measurements e.g., what is the average tubular radius of complicated objects

DANISH BIOIMAGING

NETWORK





Statistical analysis



Imaging

