

Introduction

Pathology image analysis suffers from **limited annotations!**

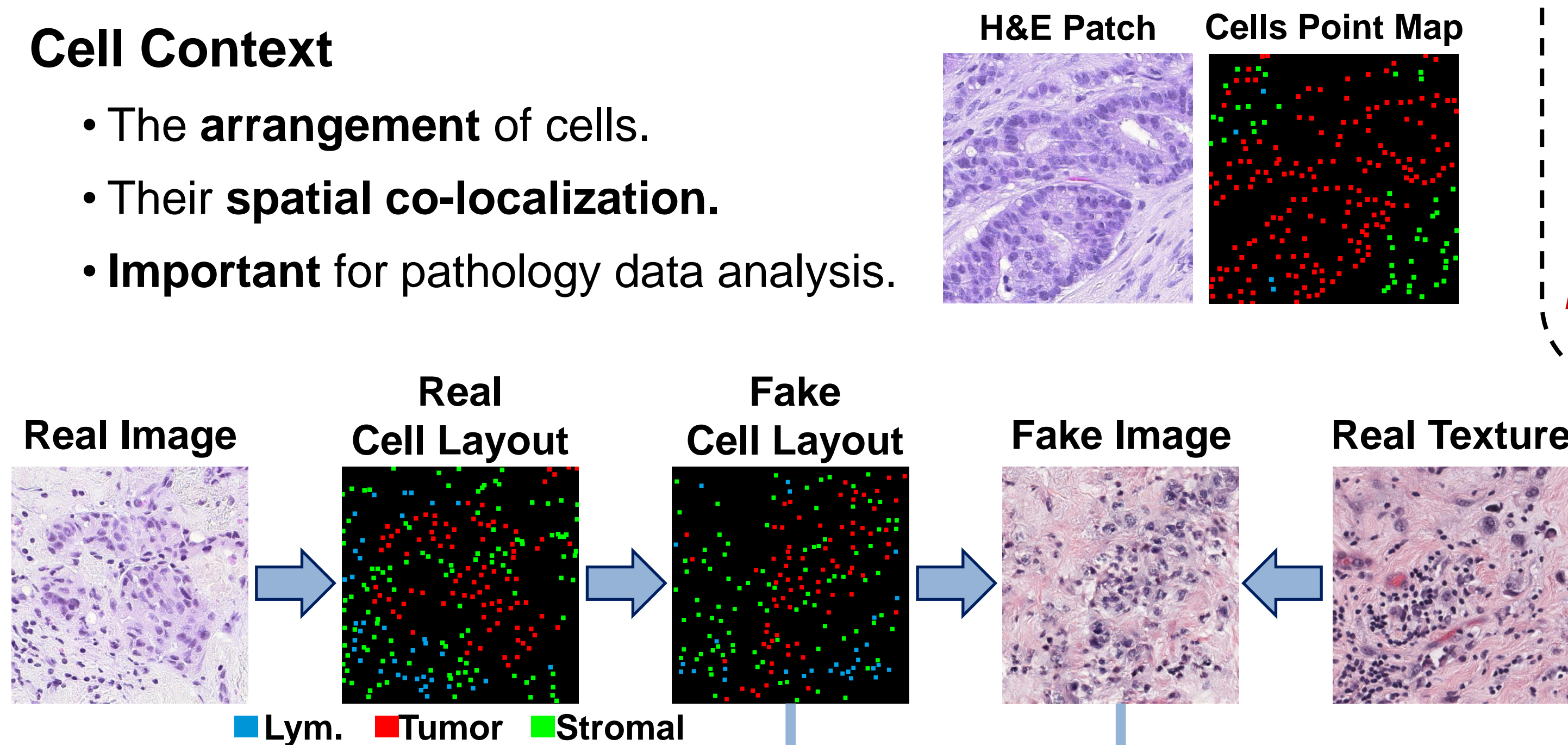
➔ **Solution: Augment training with generated labeled data.**

Generating pathology images usually involves **two steps**:

1. Generating spatial layout of cells. **Our Focus**
2. Filling in stains and textures.

Cell Context

- The **arrangement** of cells.
- Their **spatial co-localization**.
- **Important** for pathology data analysis.



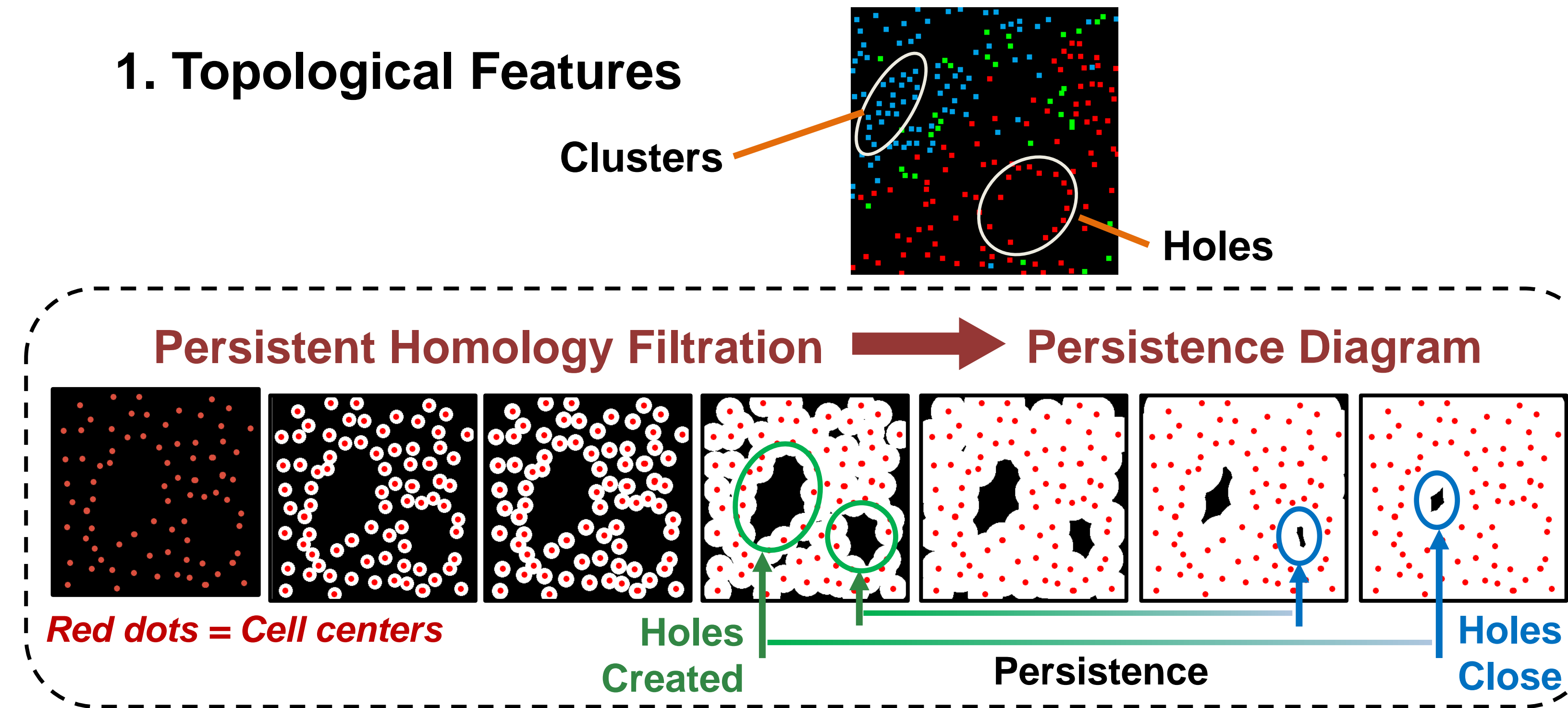
Challenges:

- **Complexity** of the cell layout.
- **How to model** and learn the underlying distribution.

We introduce mathematical descriptors to model and learn the *spatial distribution* of multi-class cells and their *structural patterns*.

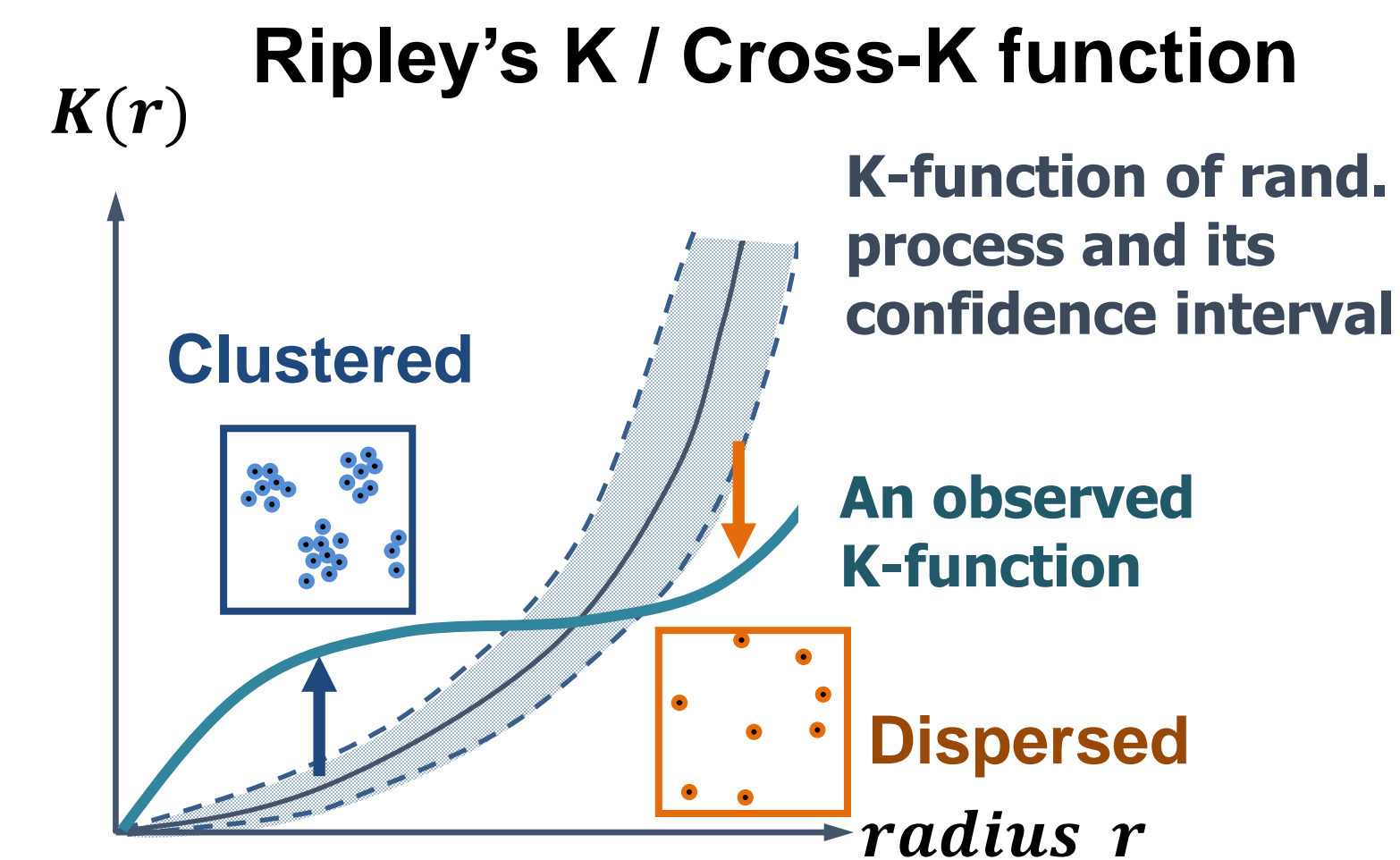
Cell Configuration Descriptors

1. Topological Features



2. Spatial Statistics Features

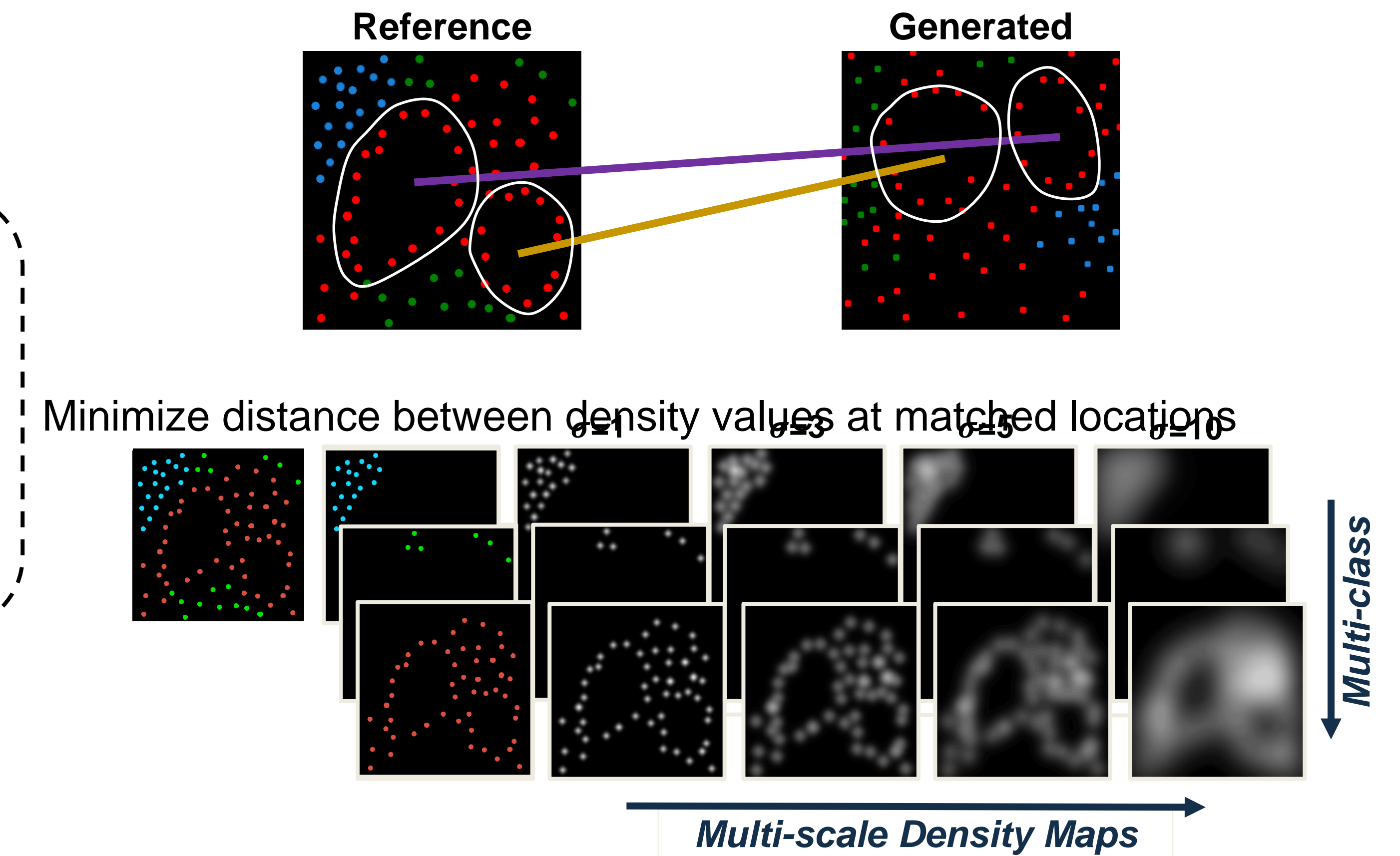
- Cell types co-localization
- Characterize neighborhoods of holes



Acknowledgements: Work funded by NSF grants CCF-2144901, IIS-2123920 and IIS-2212046, NIH and NCI grants UH3CA225021, U24CA215109, 5R01CA253368, and generous private fund from Bob Beals and Betsy Barton.

Cell Configuration Loss \mathcal{L}_{CC}

Match holes based on:
size (*persistence*), and spatial context (*cross K functions*)

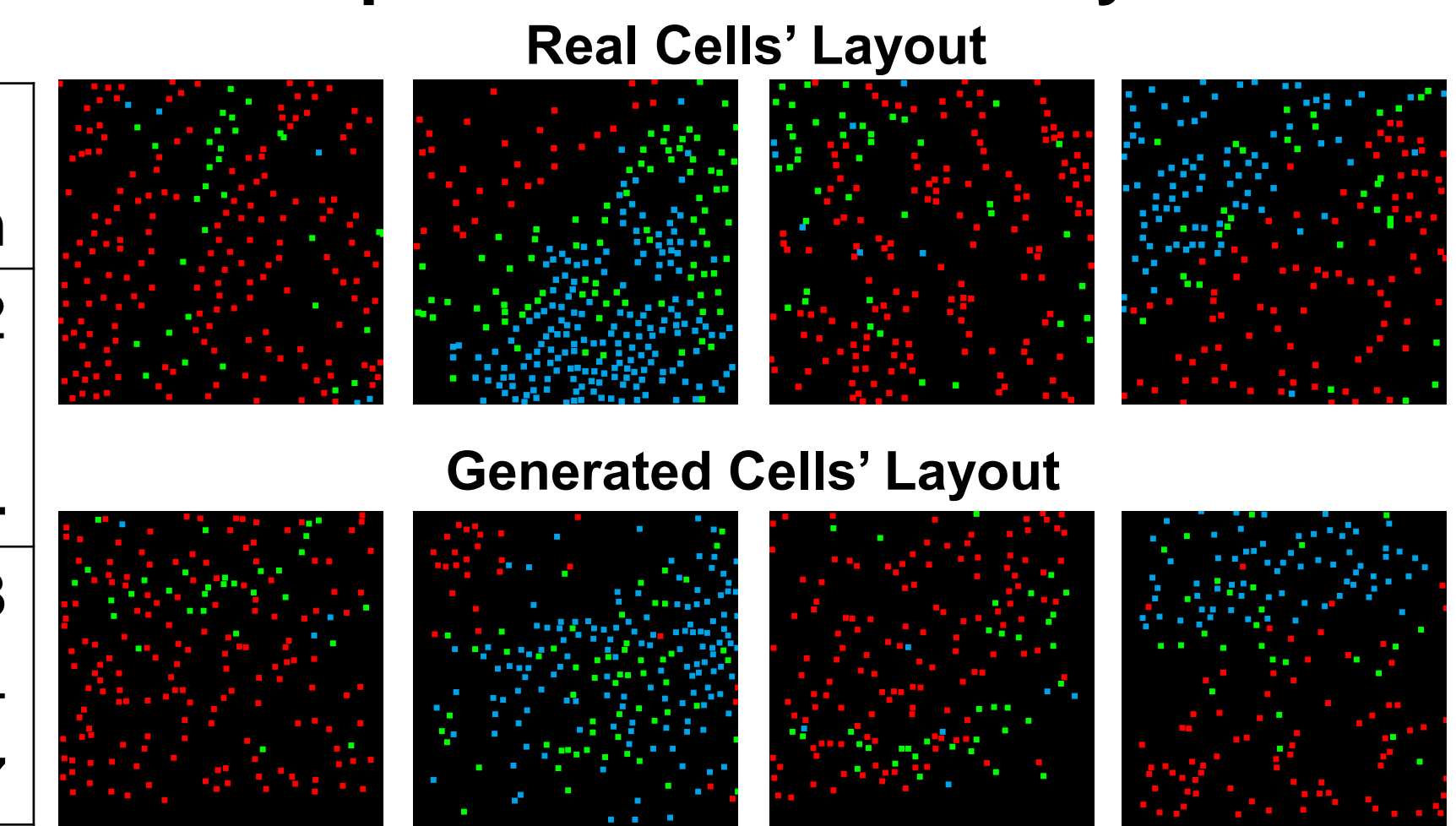


Results

Eval. on Cell Classification

| Method | F-Score | | | |
|--------------------------------|--------------|--------------|--------------|--------------|
| | Lym. | Tumor | Stro. | Mean |
| U-Net | 0.498 | 0.744 | 0.476 | 0.572 |
| U-Net + Aug. (Rand.) | 0.625 | 0.735 | 0.472 | 0.611 |
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Sample Generated Cell Layouts



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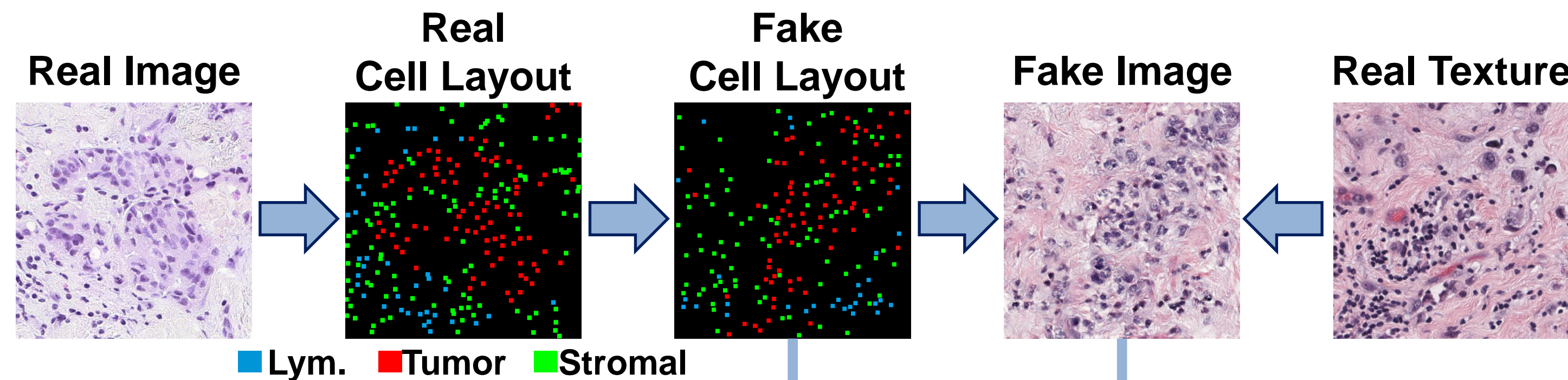
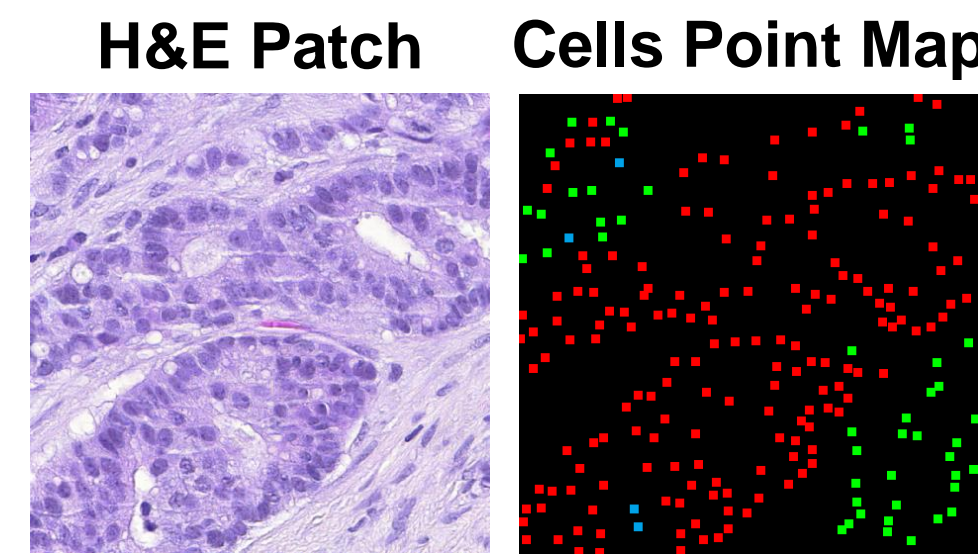
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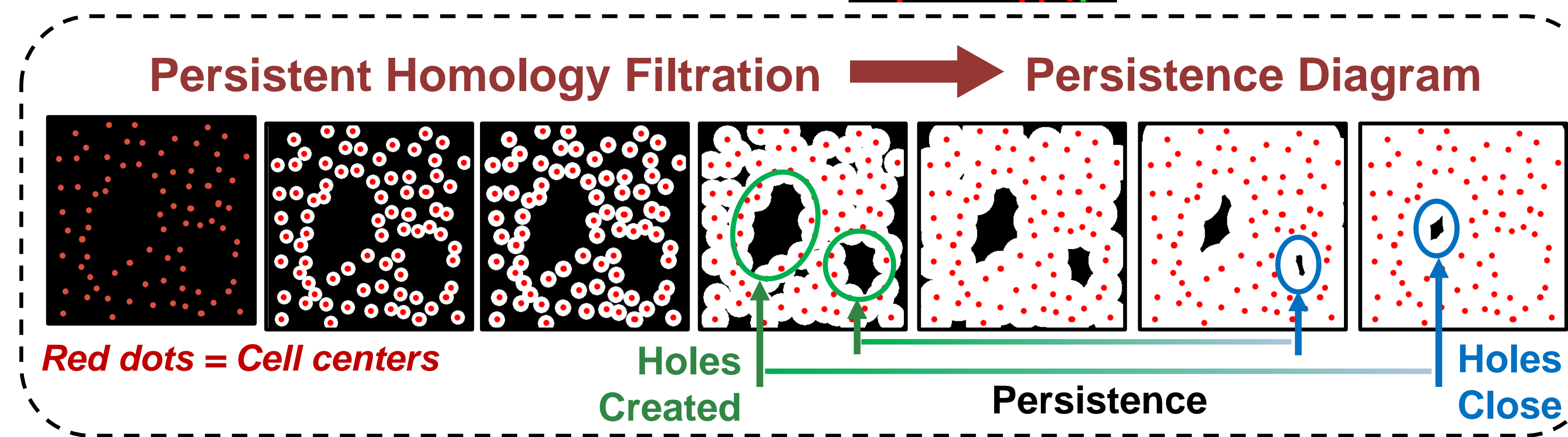
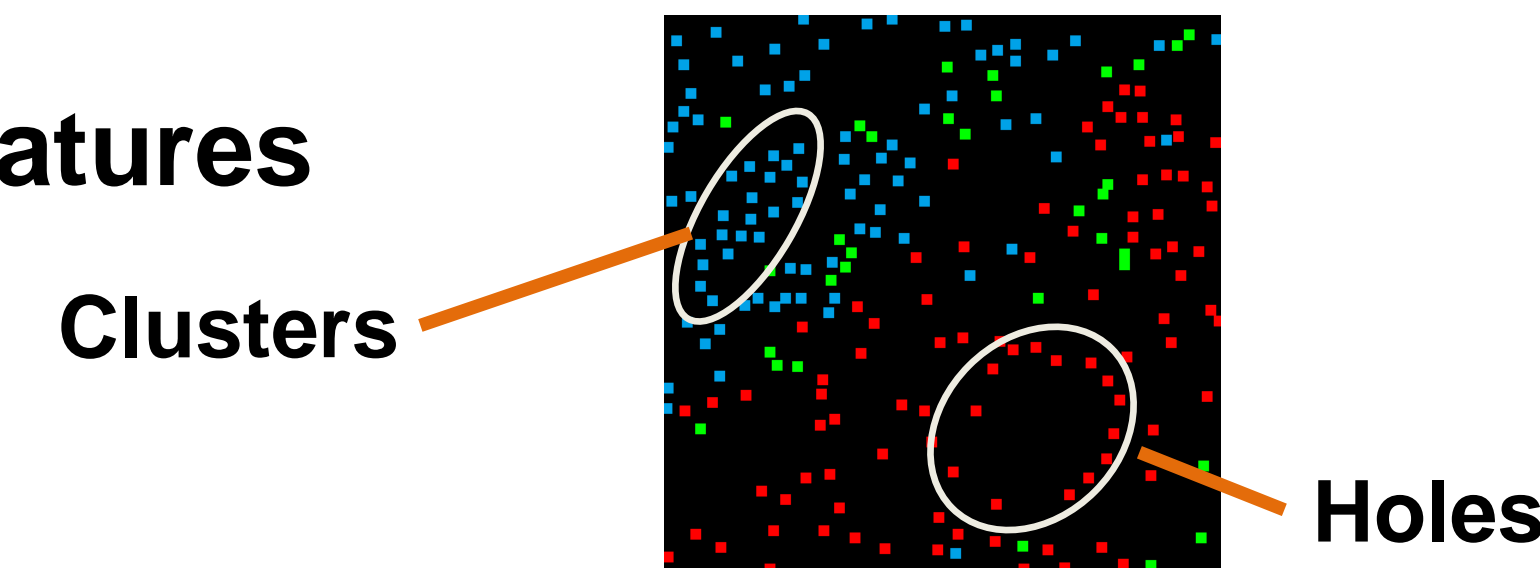
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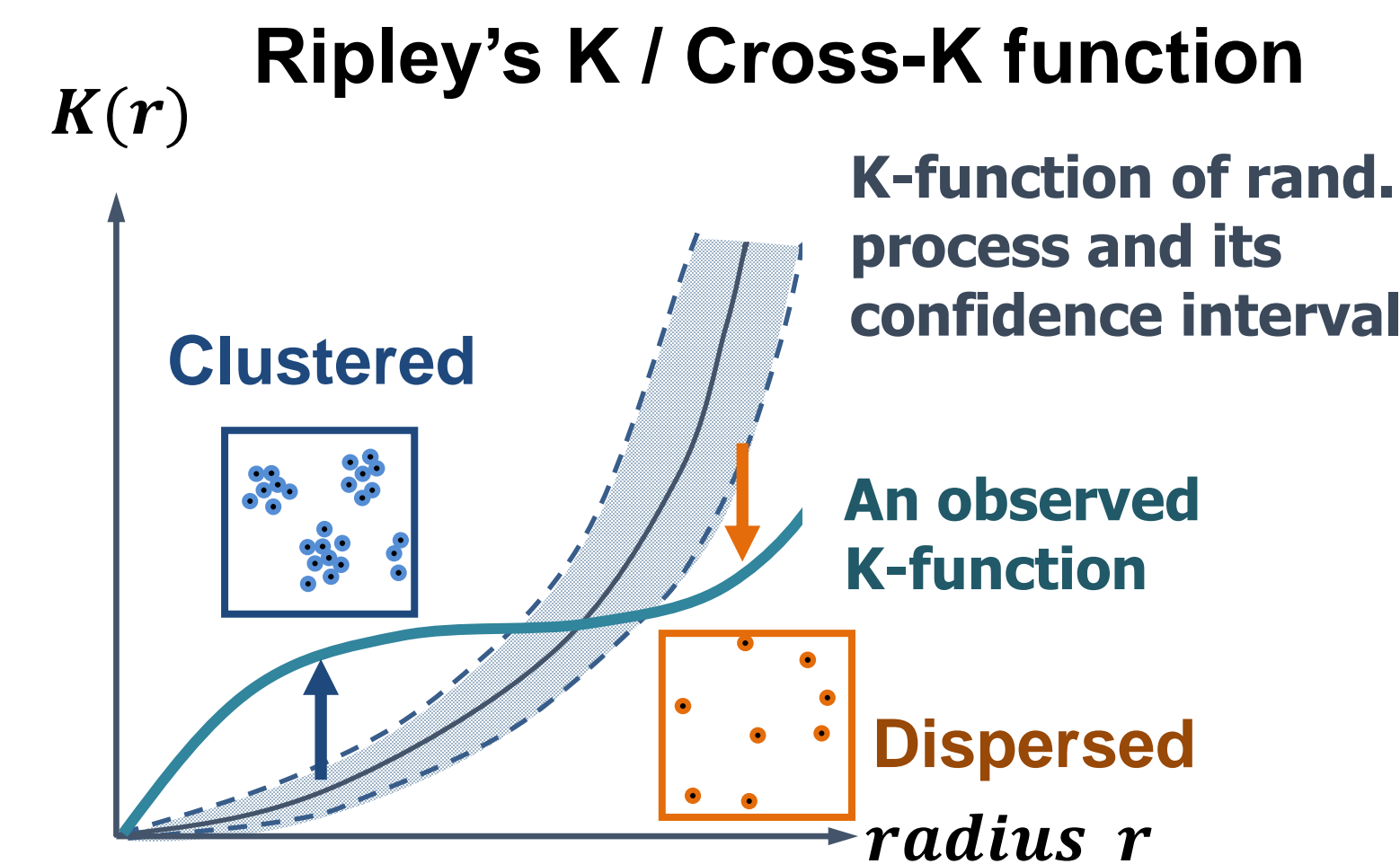
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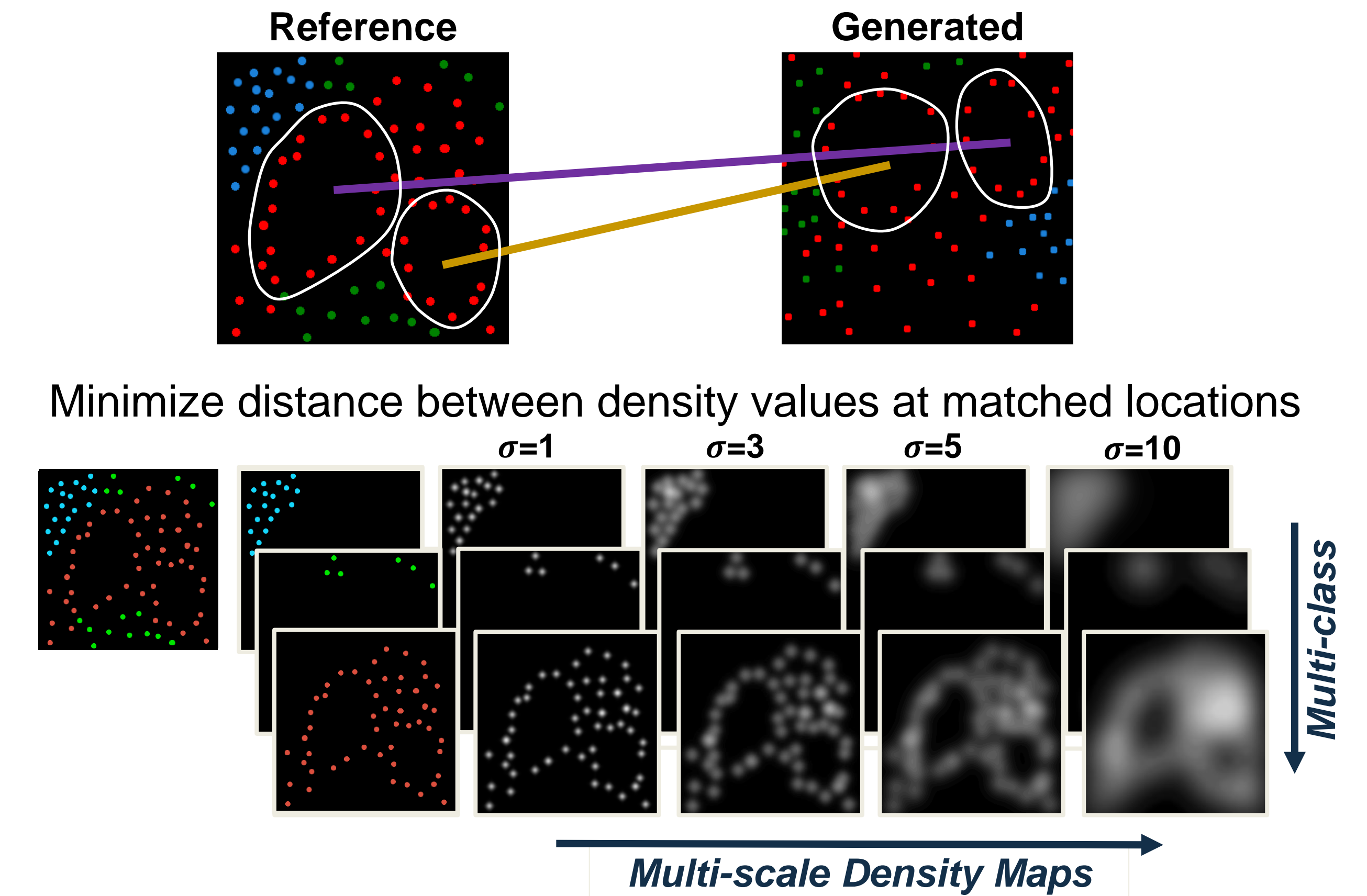
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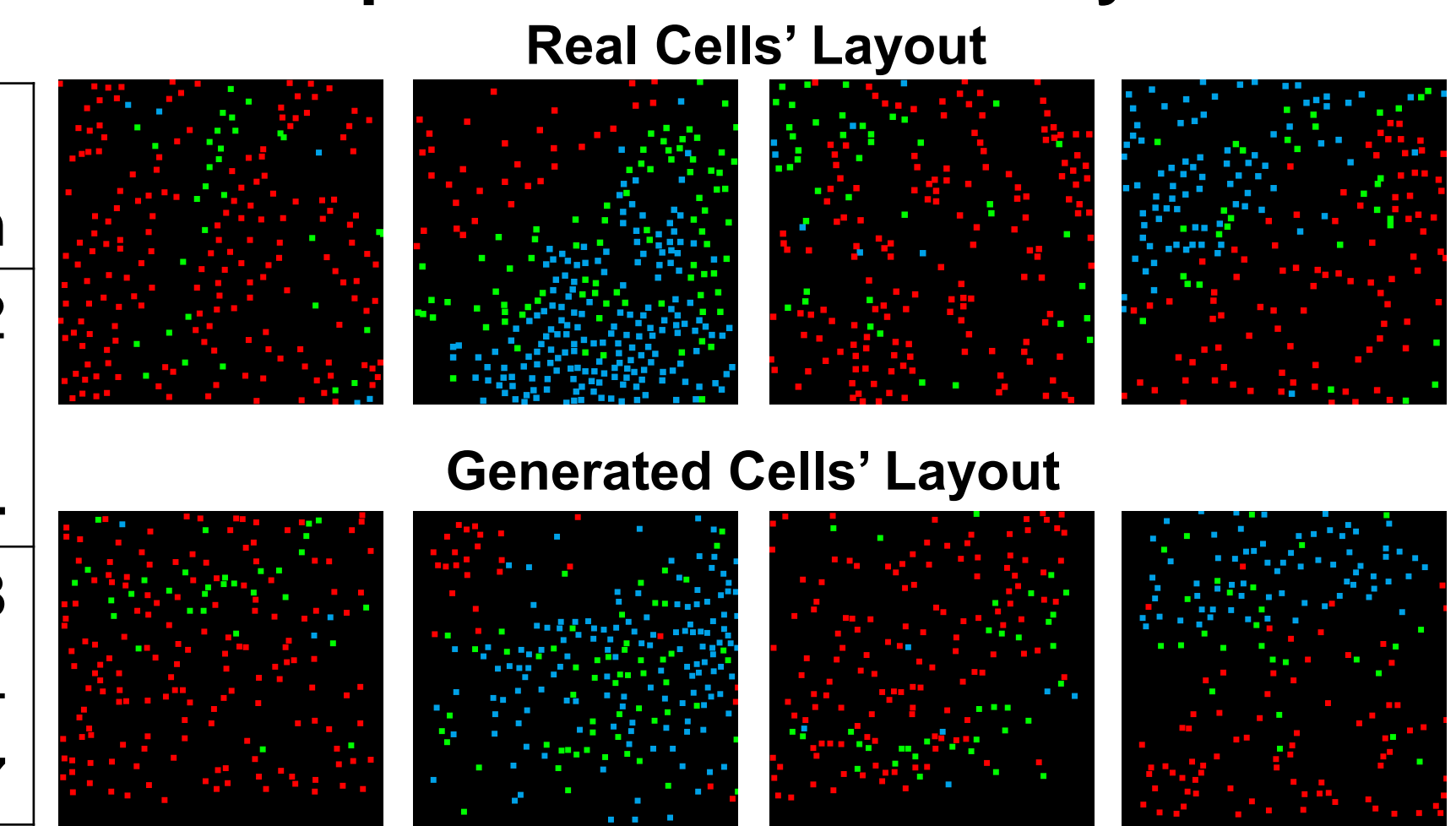


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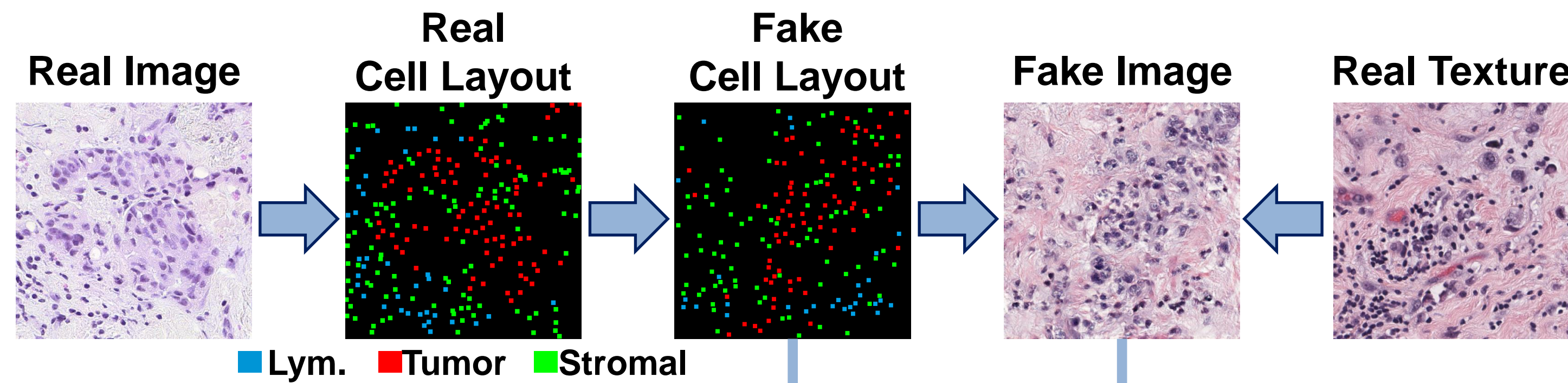
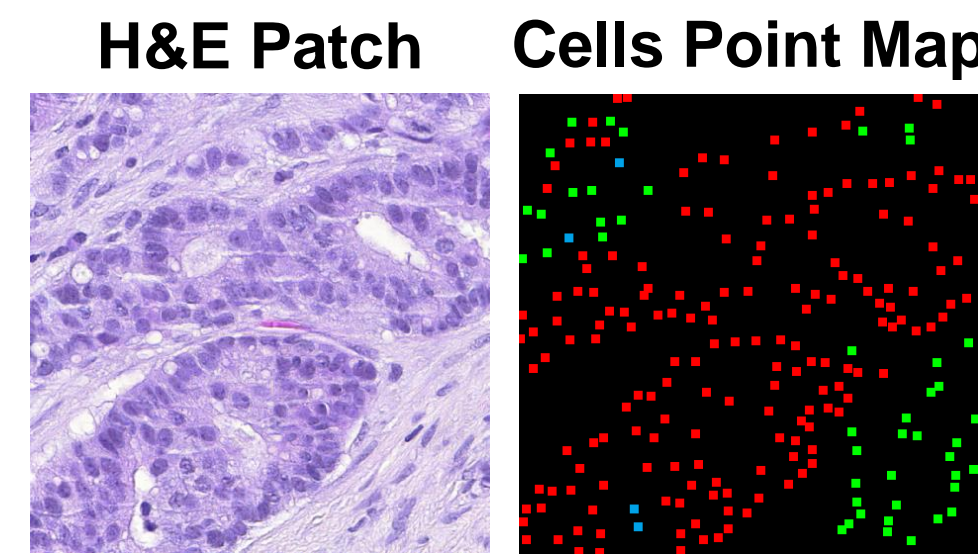
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Train Downstream Tasks

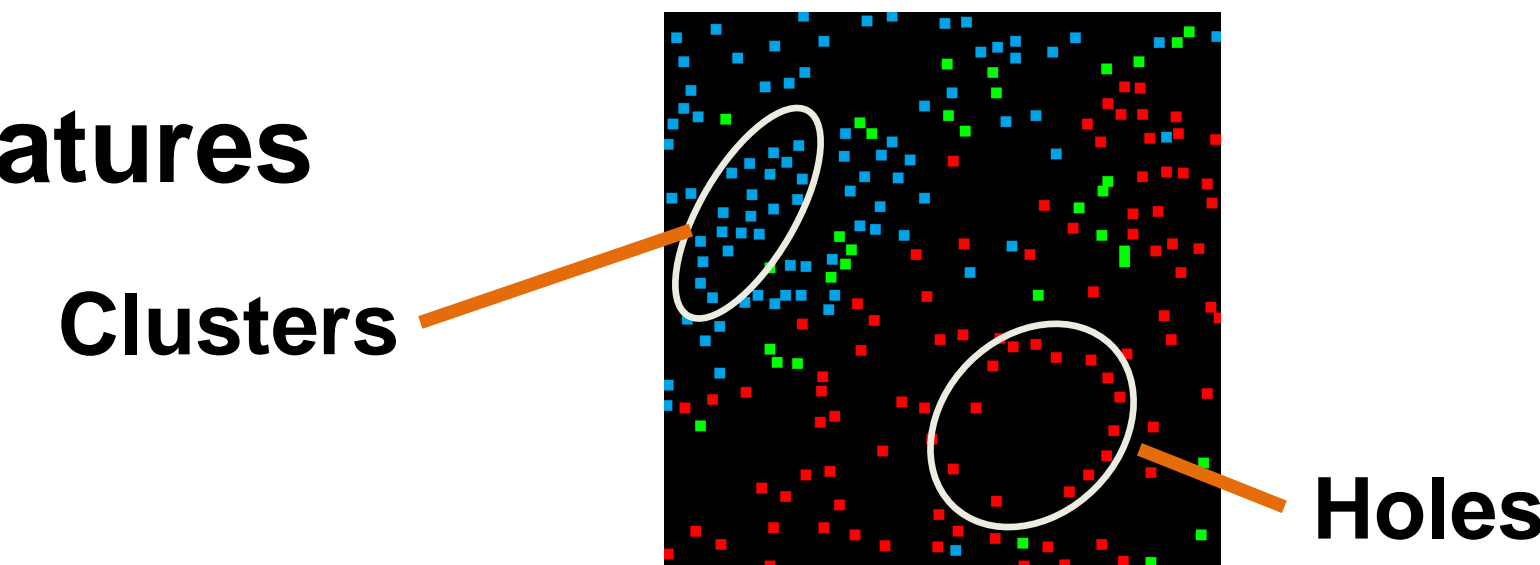
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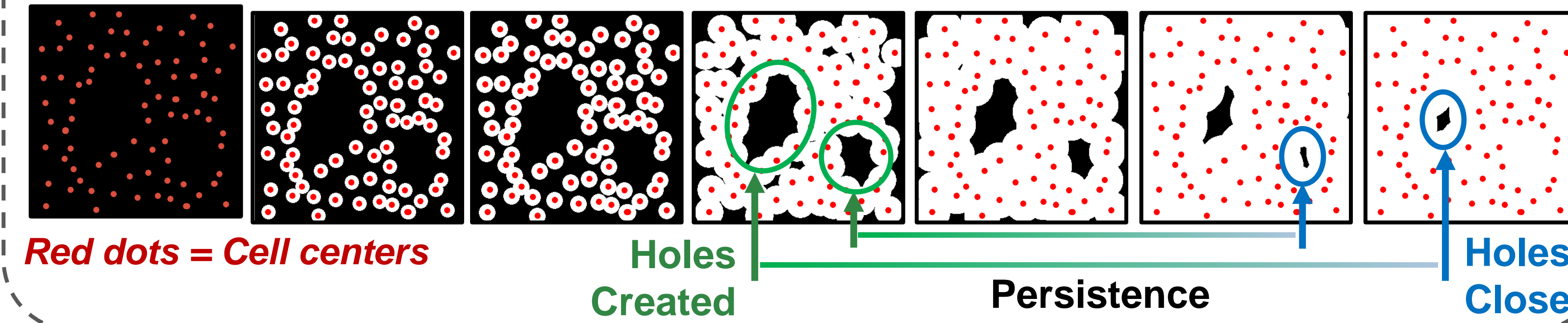
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Cell Configuration Descriptors

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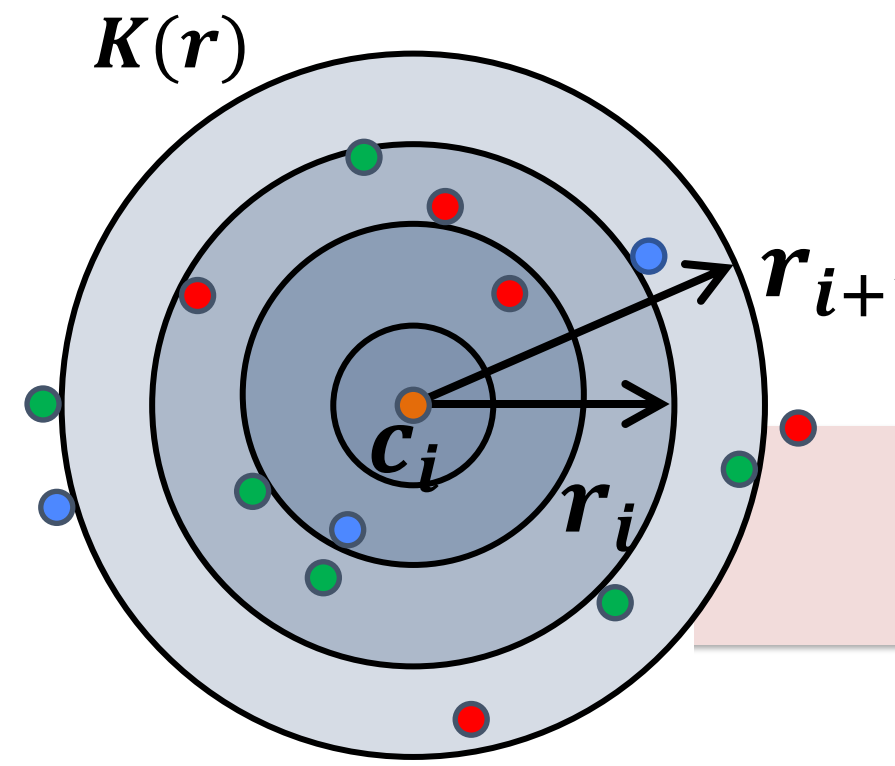
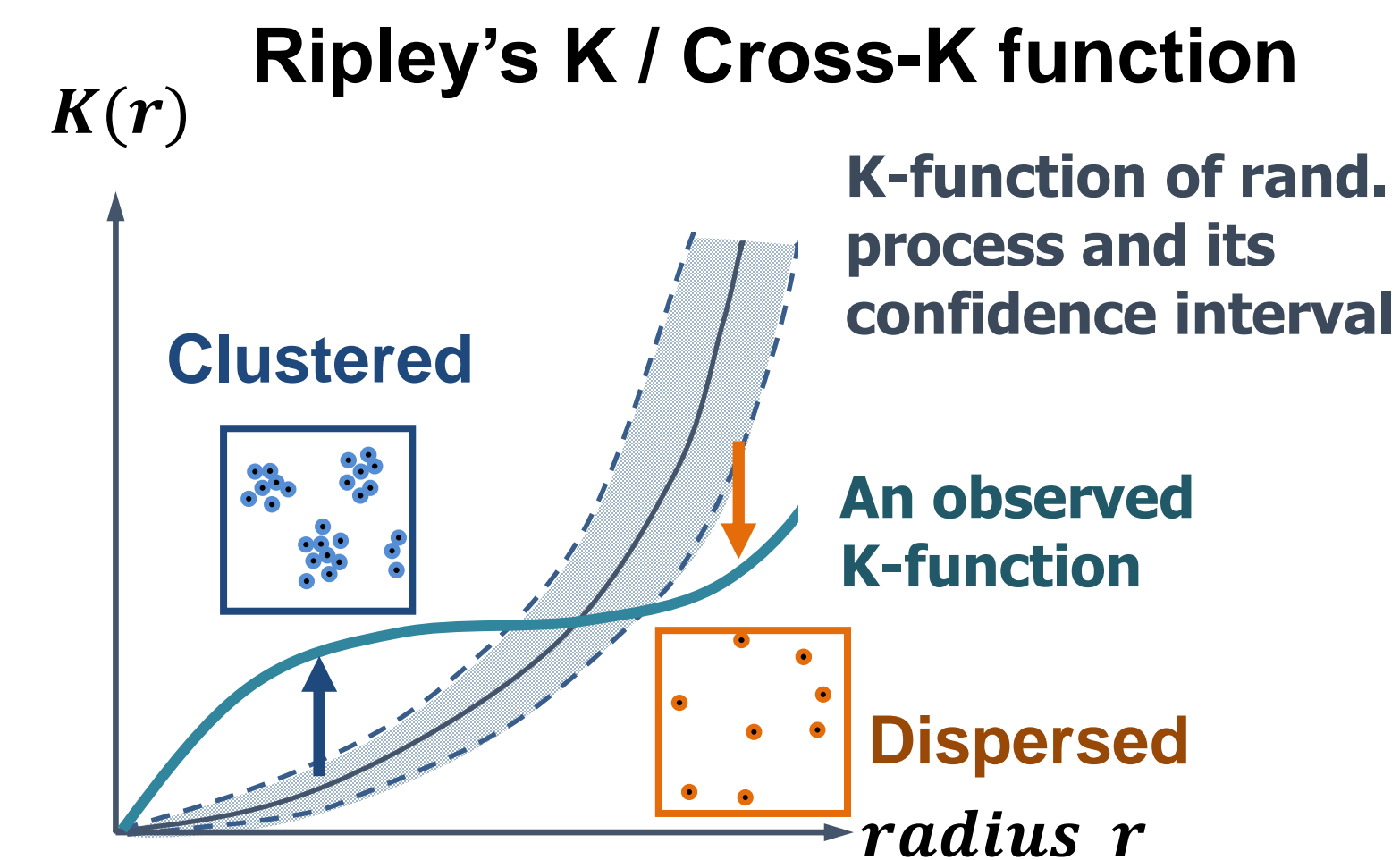


Persistent Homology Filtration ➔ Persistence Diagram



2. Spatial Statistics Features

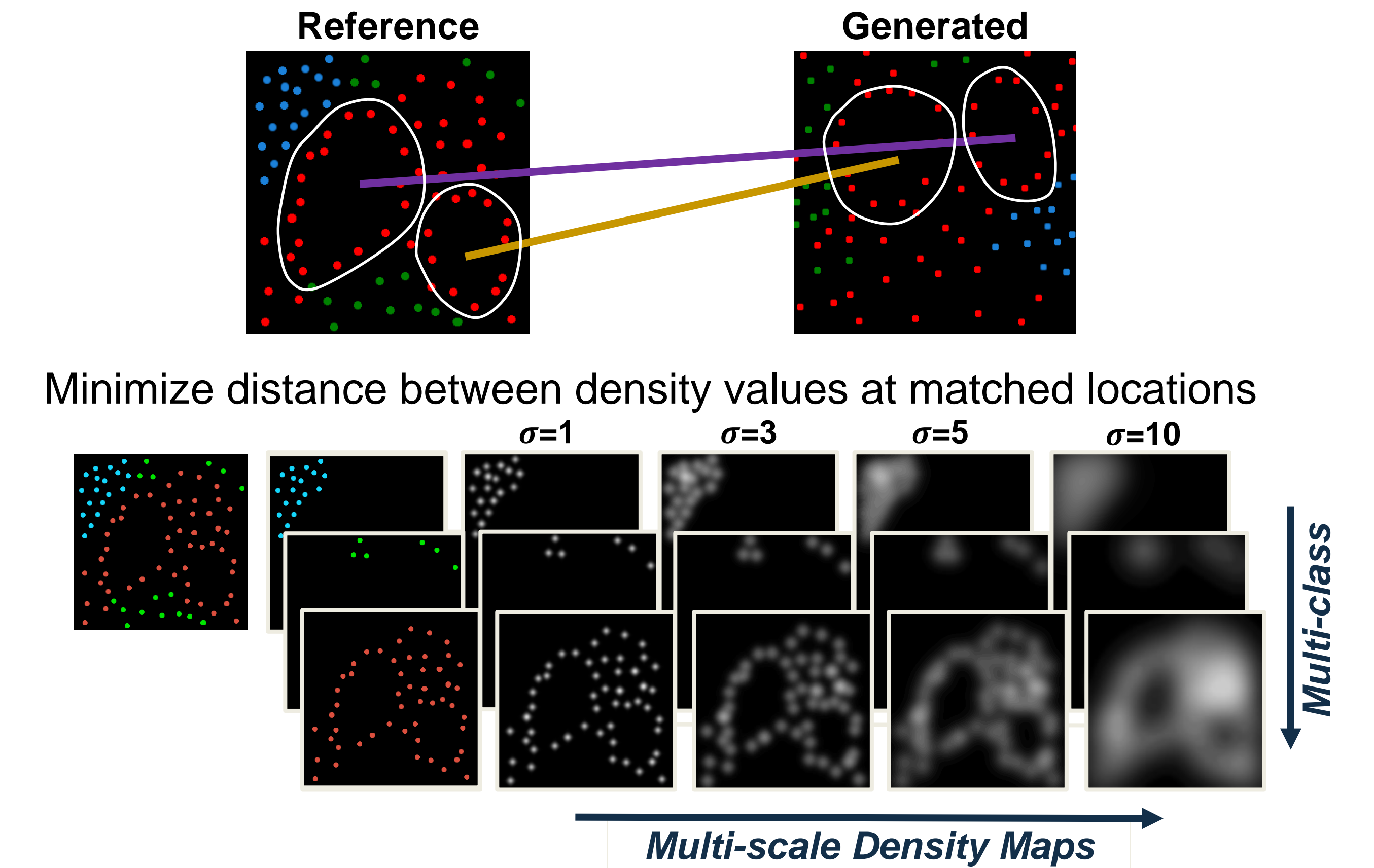
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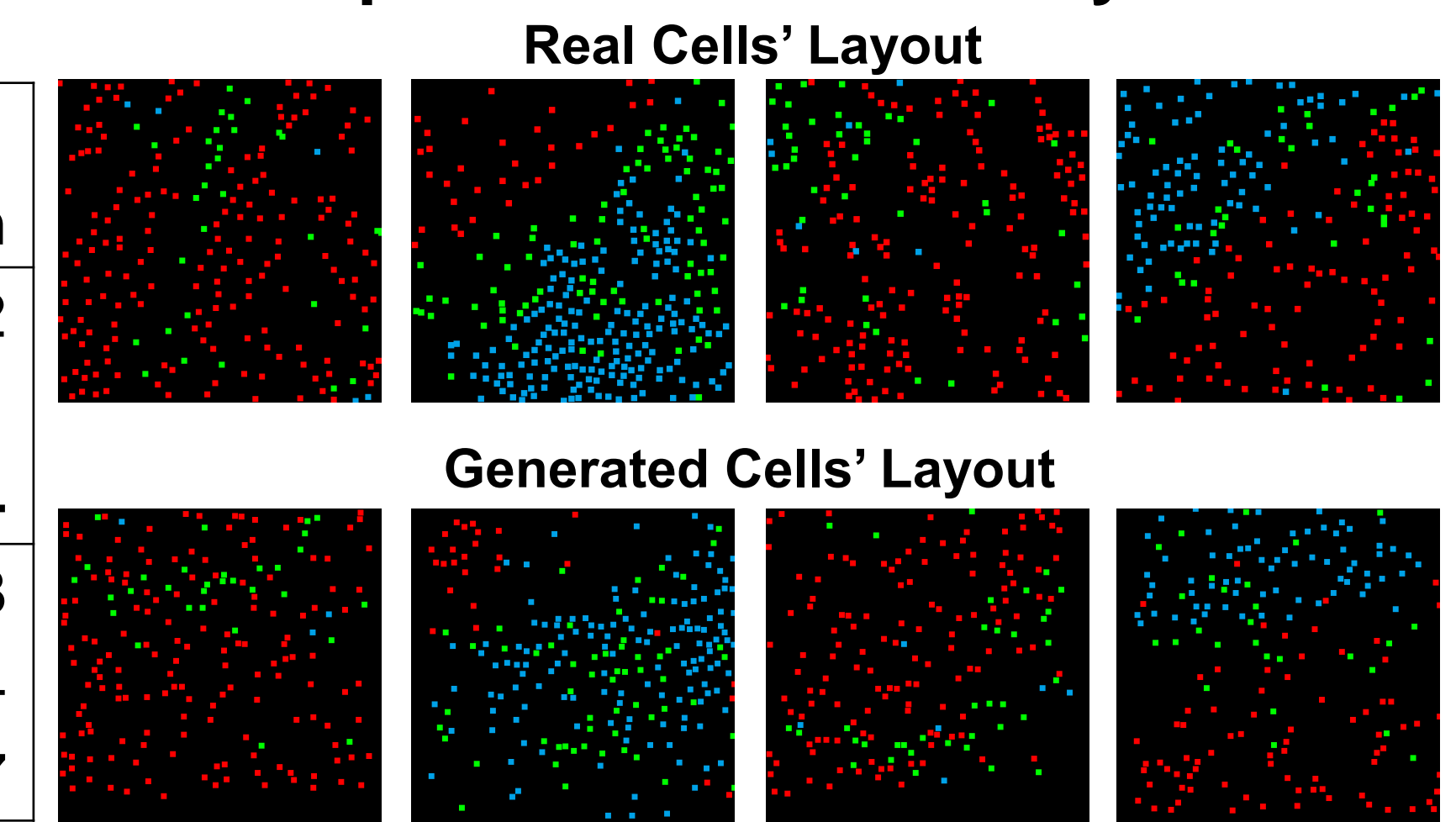


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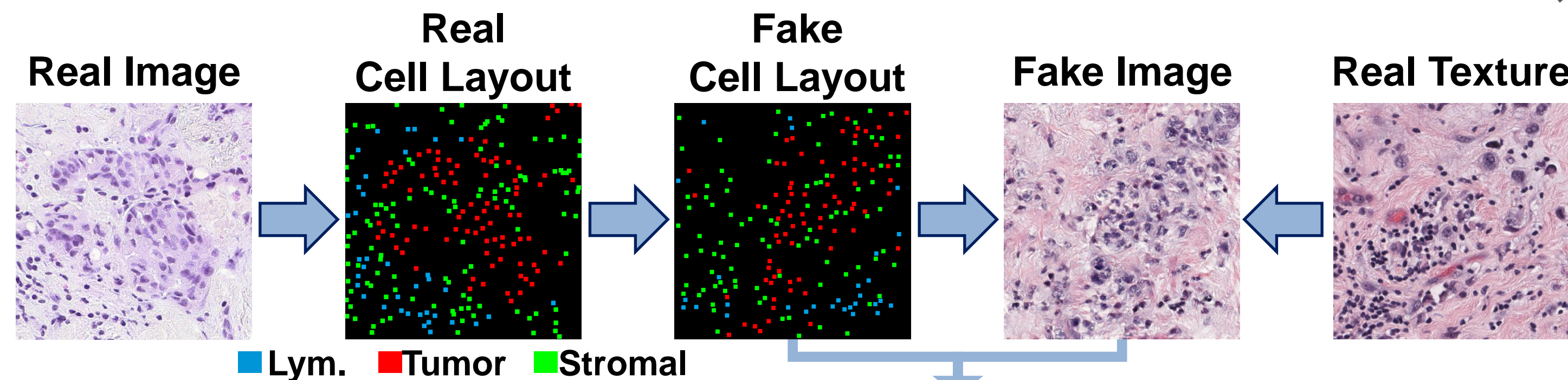
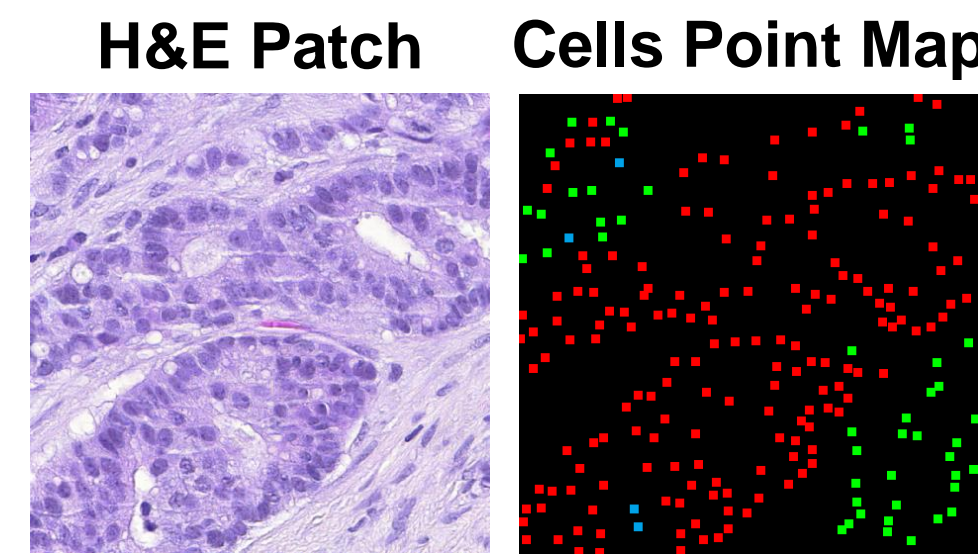
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Train Downstream Tasks

Challenges: • Complexity of the cell layout.

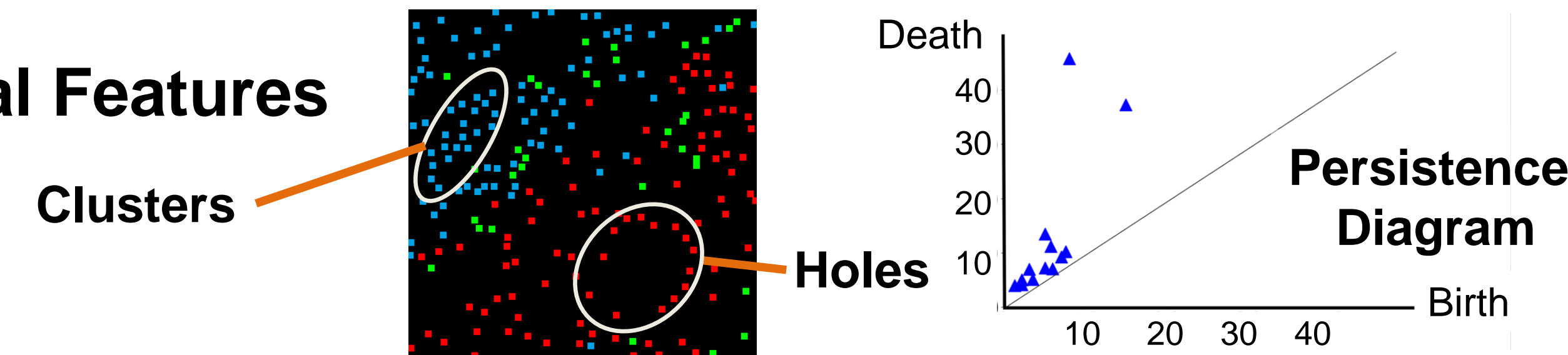
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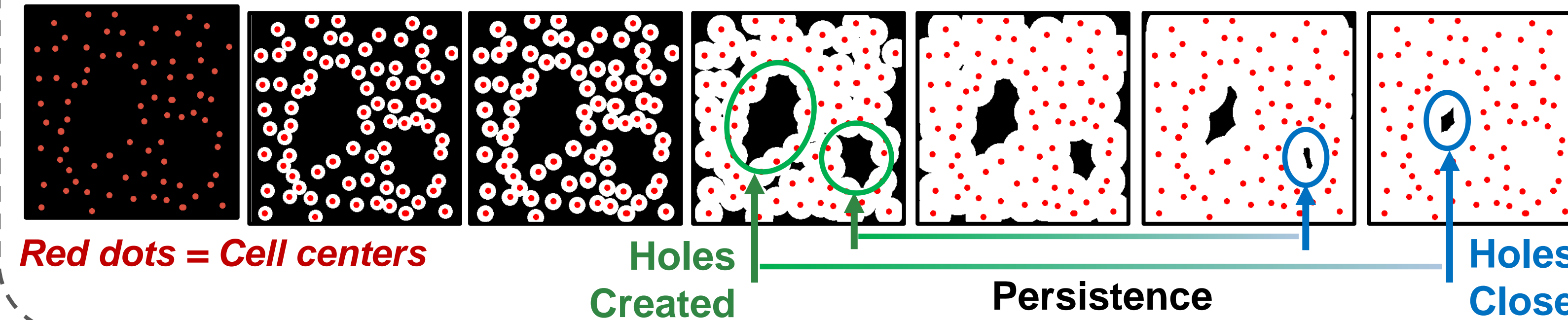
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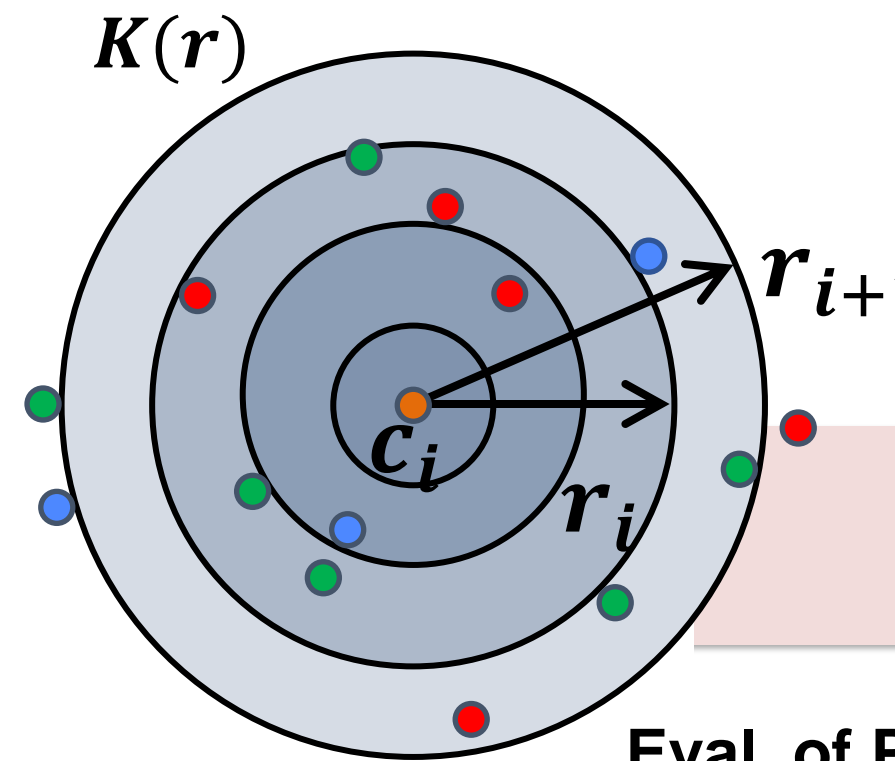


Persistent Homology Filtration → Persistence Diagram (PD)

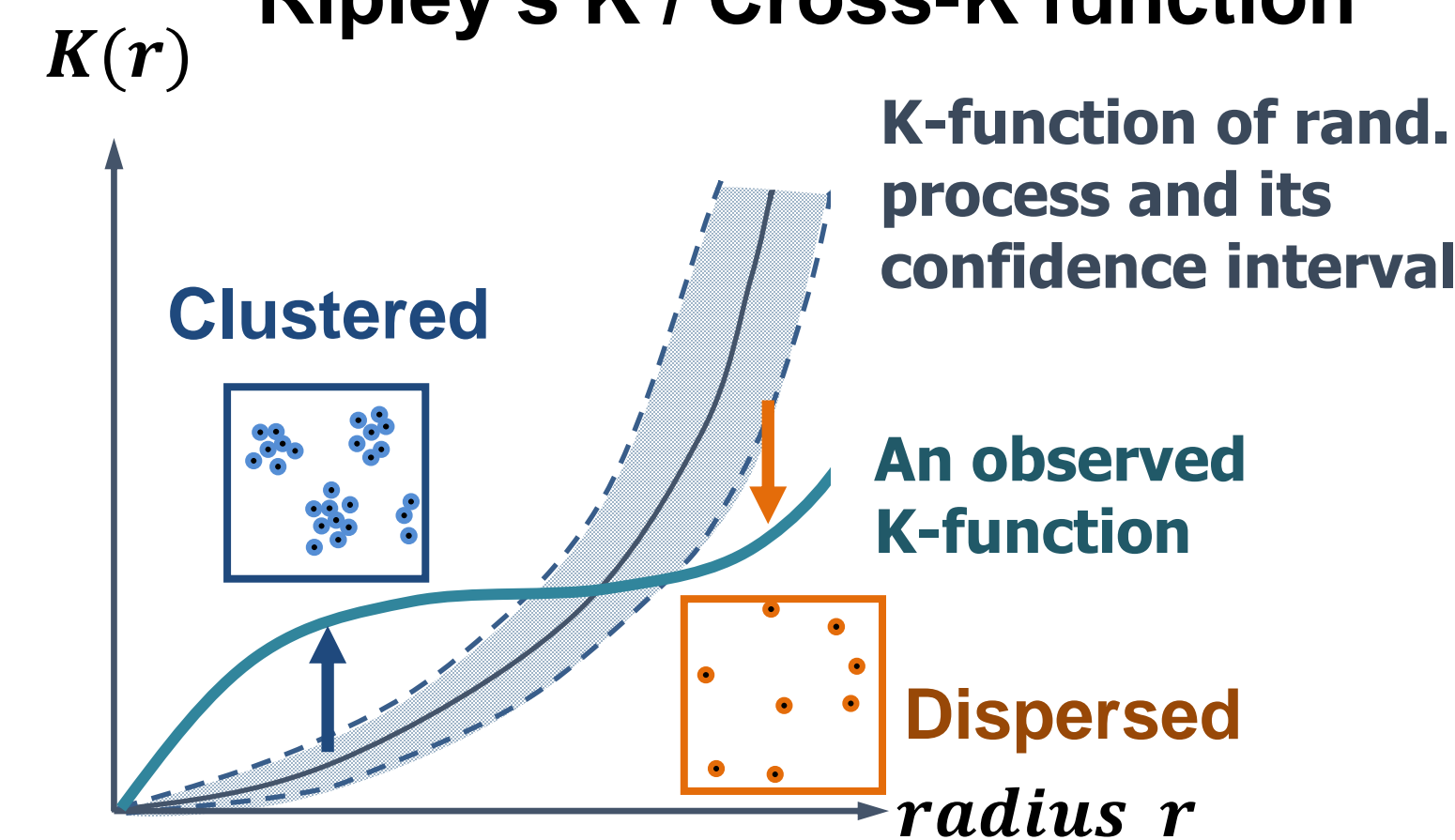


2. Spatial Statistics Features

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Ripley's K / Cross-K function

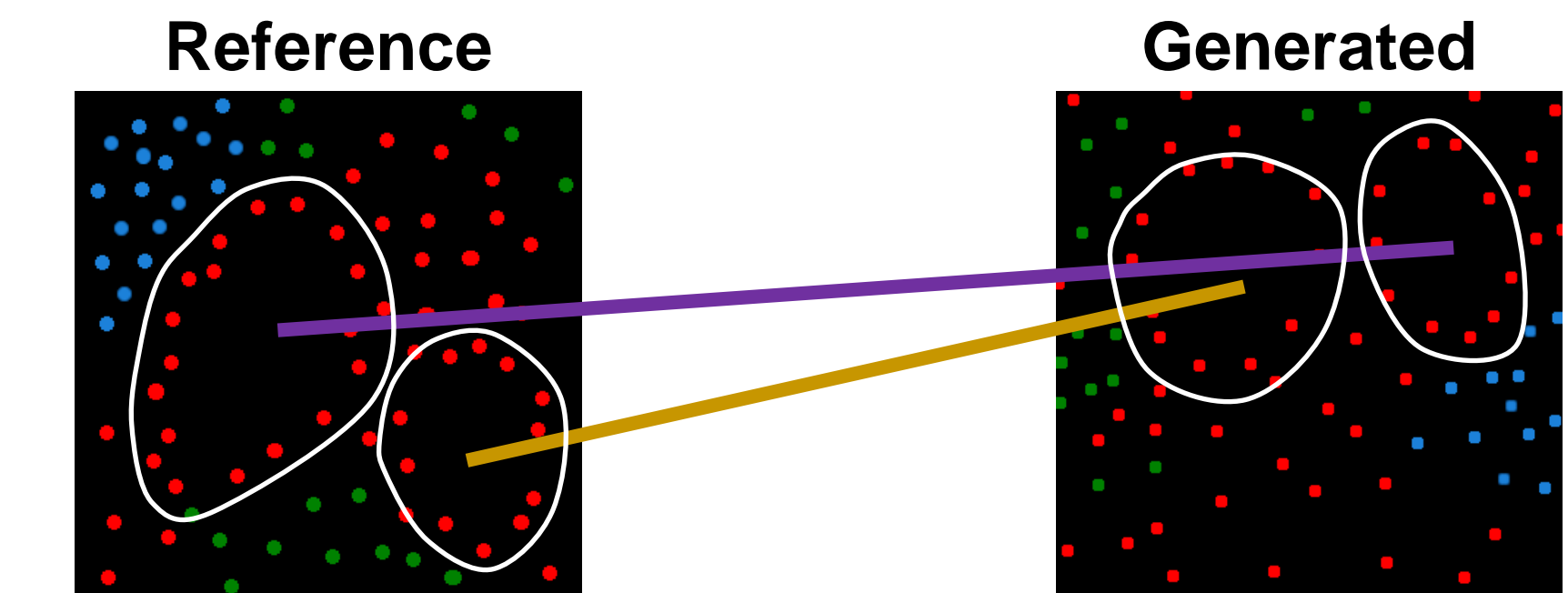


Model Backbone: A modified version of SPGAN*

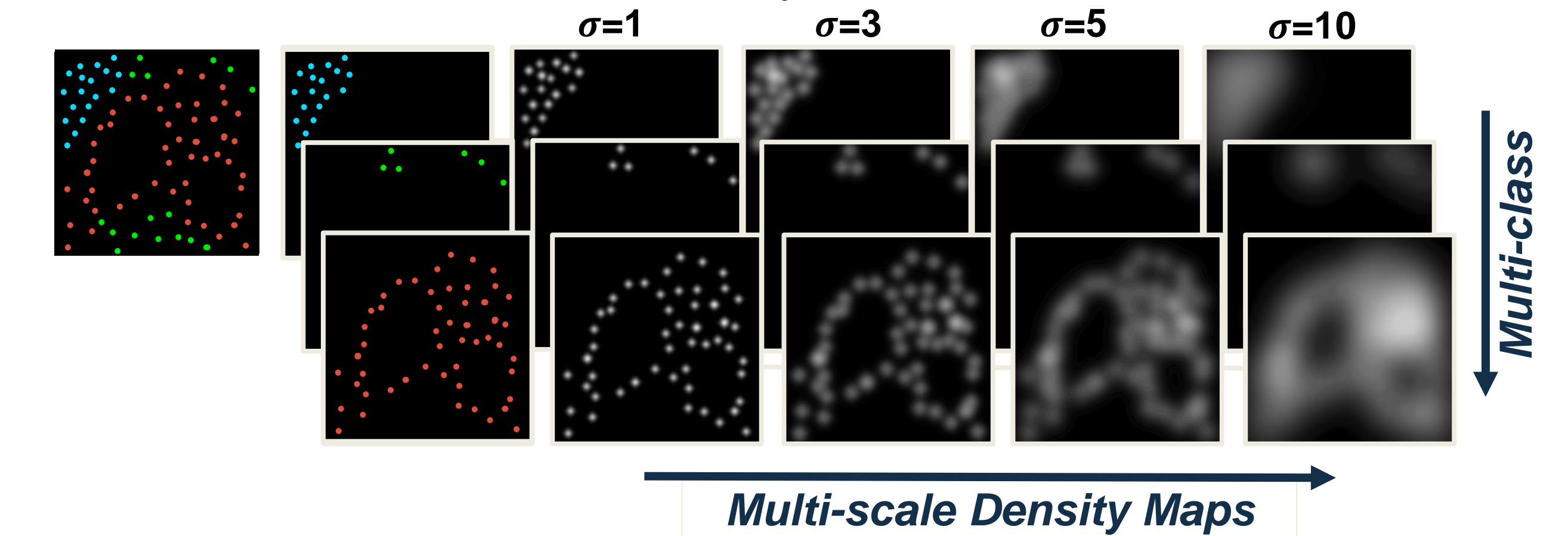
* Li et al. SP-GAN: sphere-guided 3d shape generation and manipulation. SIGGRAPH, 40(4), 2021.

Cell Configuration Loss \mathcal{L}_{CC}

Match holes based on:
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Minimize distance between density values at matched locations



Results

Eval. of Persistence Diagrams (Cell Config. Matching) ↓

| Method | Lym. | Tumor | Stro. | Mean |
|--|------|-------|-------|-------------|
| w/o Spatial Descriptors + w/o \mathcal{L}_{CC} | 0.8 | 1.74 | 1.66 | 1.4 |
| w/o \mathcal{L}_{CC} | 0.9 | 1.69 | 1.79 | 1.46 |
| w/o Cross K-function Descriptor | 0.75 | 1.74 | 1.77 | 1.42 |
| Ours | 0.74 | 1.64 | 1.71 | 1.36 |

Eval. on Cell Classification (F-Score) ↑

| Method | Lym. | Tumor | Stro. | Mean |
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