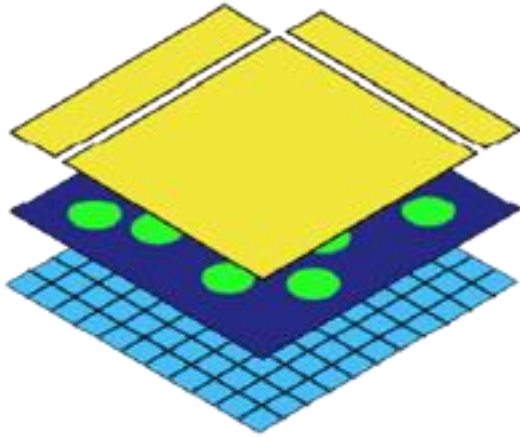


Spatialdata 101

Technical Intro to the
scVerse framework

Teach + Learn Session MannLabs
2025

Why spatialdata?



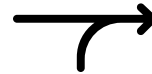
Integrated modalities



Unified data storage



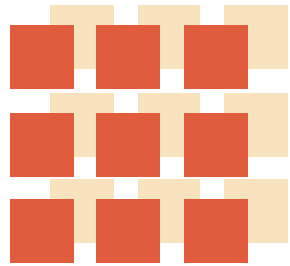
Scalable



Compatible with many packages

Spatial modalities

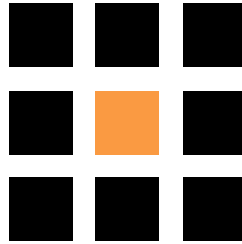
Images



$$c \times y \times x$$



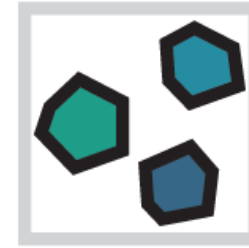
Rasterized annotation



$$y \times x$$



Vectorized annotation



+ Shapely

Tables



AnnData

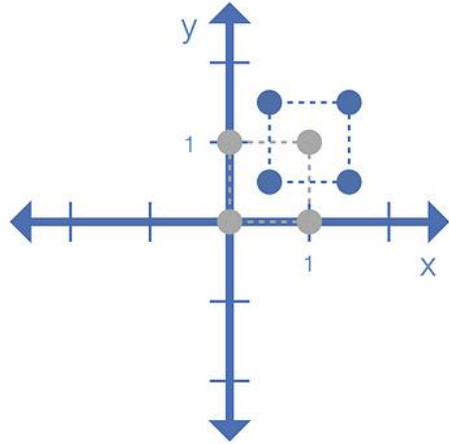
Knowing the fundamental packages is often helpful for debugging/troubleshooting

Transformations

SpatialData encodes spatial transformations as matrix operations

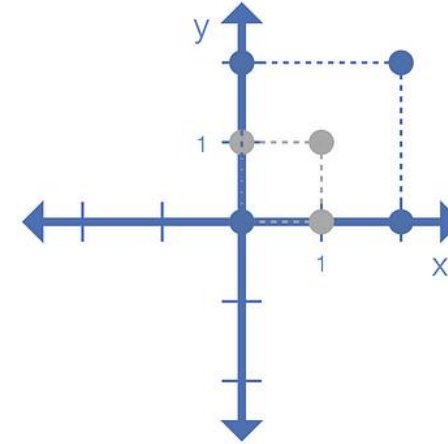
Translate

$$\begin{bmatrix} 1 & 0 & 1 \\ 0 & 1 & 1 \\ 0 & 0 & 1 \end{bmatrix}$$



Scale

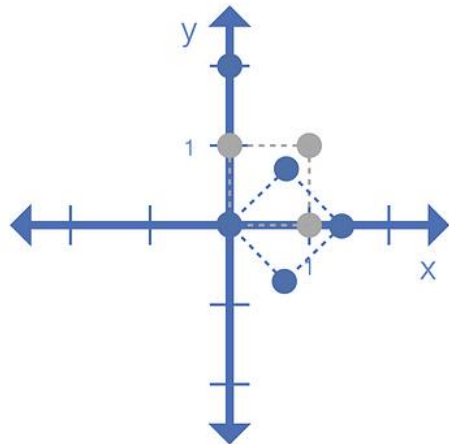
$$\begin{bmatrix} 2 & 0 & 0 \\ 0 & 2 & 0 \\ 0 & 0 & 1 \end{bmatrix}$$



Rotate

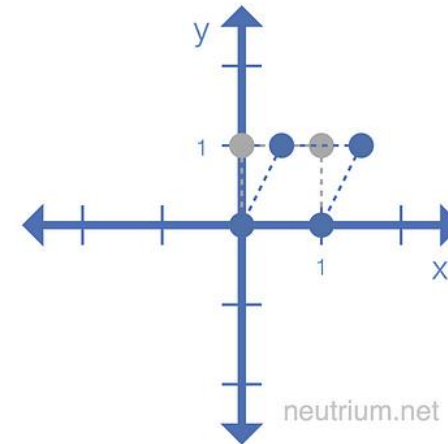
$$\begin{bmatrix} c & s & 0 \\ -s & c & 0 \\ 0 & 0 & 1 \end{bmatrix}$$

$$c = s = \sin(45^\circ)$$



Shear

$$\begin{bmatrix} 1 & 0.5 & 0 \\ 0 & 1 & 1 \\ 0 & 0 & 1 \end{bmatrix}$$



neutrium.net

The spatial ecosystem

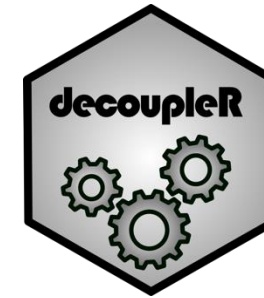
scRNAseq analysis



Spatial analysis



TF activity



Cell-cell-communication



Image analysis

(Cell segmentation + featurization)



scPortrait

HE Stains



Trajectory Inference



Optimal transport



Tutorial

See GitHub

- <https://github.com/lucas-diedrich/spatialdata-learning.git>

References

- Marconato, L. *et al.* SpatialData: an open and universal data framework for spatial omics. *Nat Methods* 1–5 (2024) doi:[10.1038/s41592-024-02212-x](https://doi.org/10.1038/s41592-024-02212-x).
- Virshup, I. *et al.* The scverse project provides a computational ecosystem for single-cell omics data analysis. *Nat Biotechnol* **41**, 604–606 (2023).