

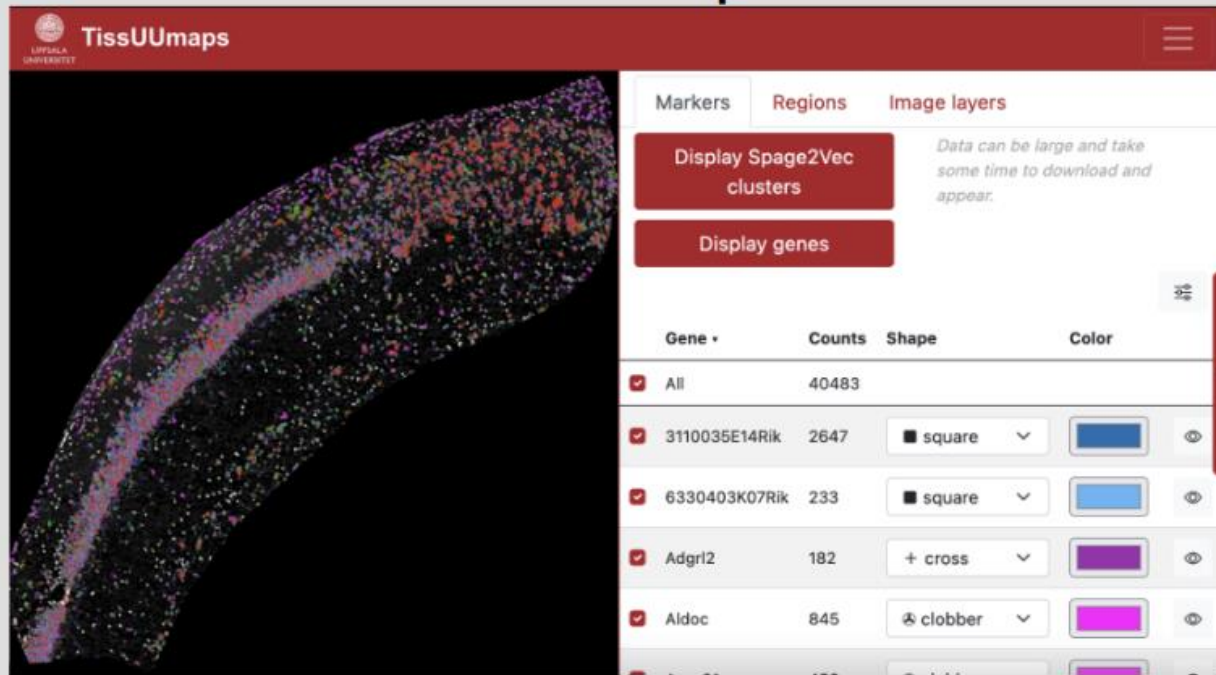
TissUUmaps 3: Interactive visualization and quality assessment of large-scale spatial omics

Christophe Avenel

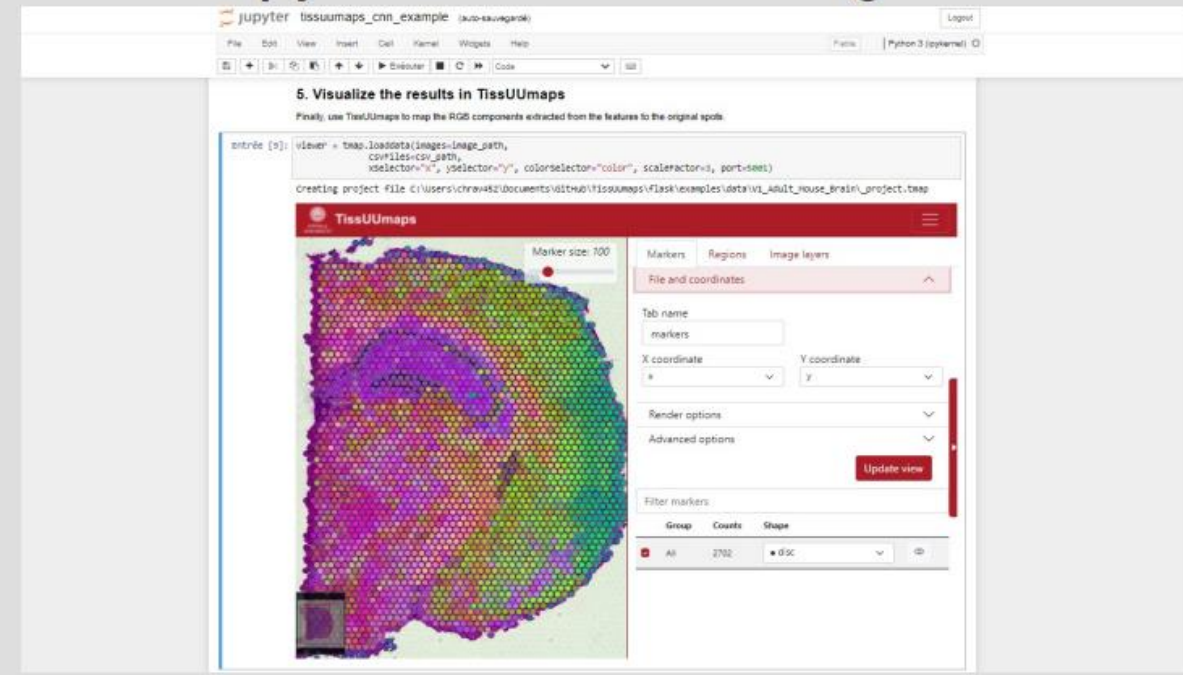
*Dept. of Information Technology, Uppsala University, and
NBIS / BioImage Informatics Facility*

Interactive viewer

Web & Desktop viewer



Jupyter Notebook integration

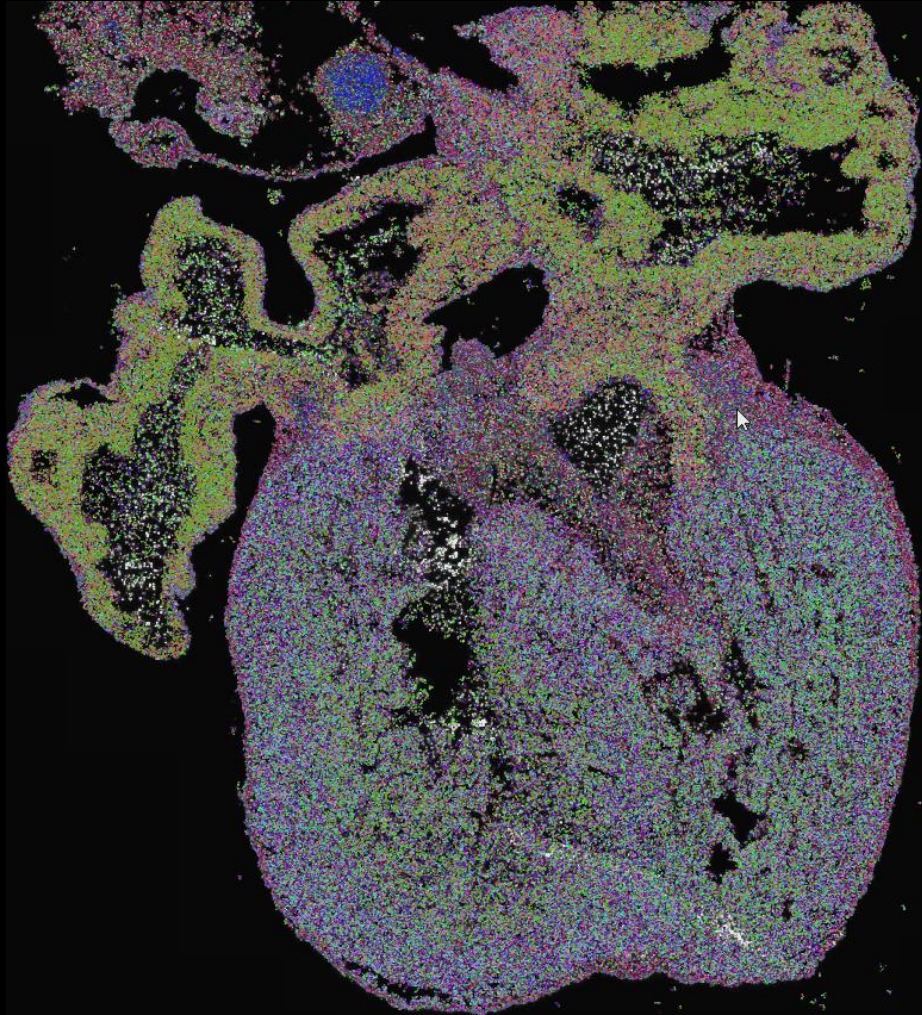


- Install locally with *pip* / *Windows installer* / *Docker*

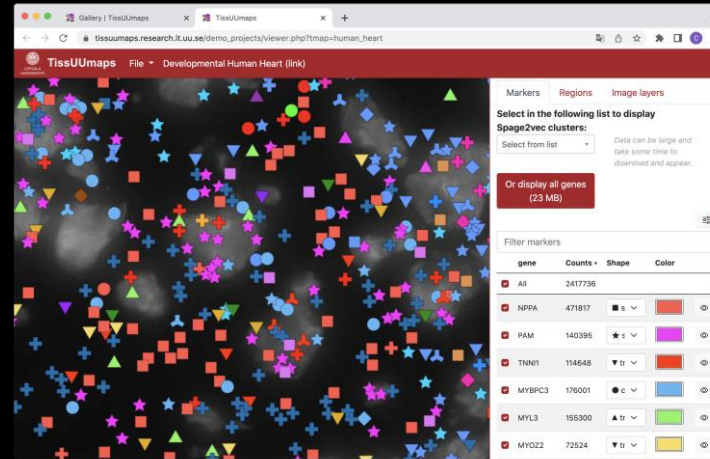
Challenge:

Viewing and sharing large spatial omics datasets without having to download large files or install software.

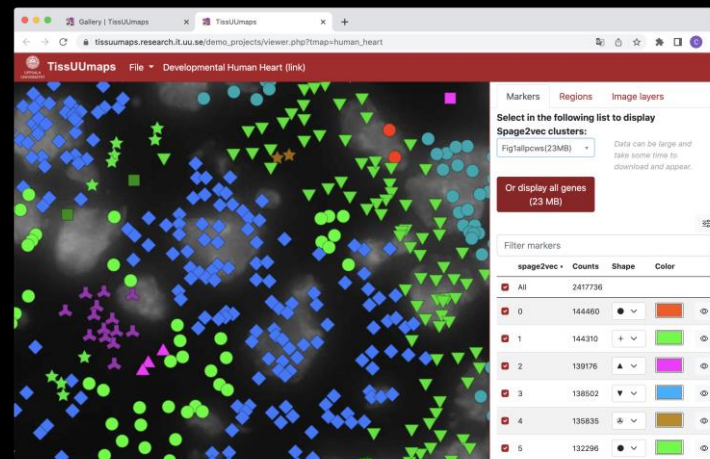
...and interact with the data, do spatial statistics, define ROIs, play with clustering alternatives, and check decoding quality...



Organ resolution (human heart at 6 pcws)



Sub-cellular resolution in situ sequencing data

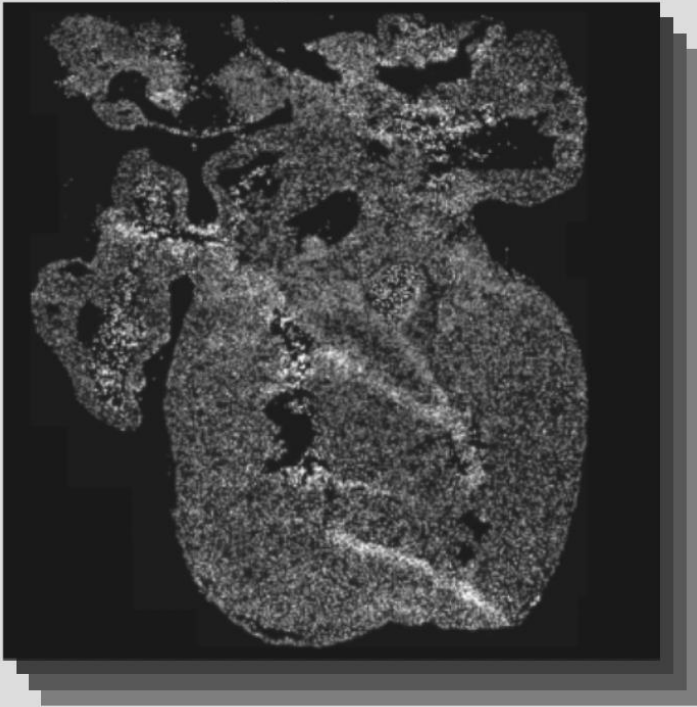


Cell-typing by spatial statistics

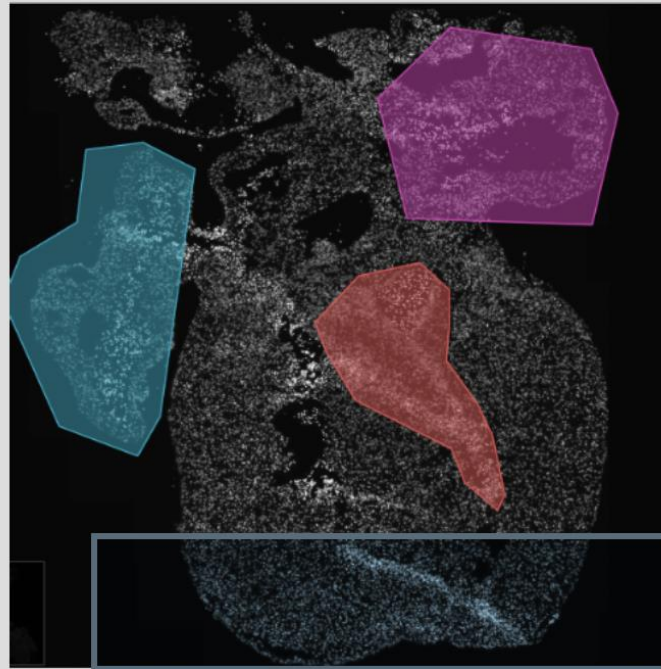
Types of input data

- 2D images, 8-bit, (of tissue), any modality, multiple layers
- Regions, drawn manually or defined using tools such as QuPath, Cellpose etc (GeoJSON)
- Markers, with x- and y coordinates, type and value, or even advanced info such as pie-charts (input as .csv file, could be generated from AnnData)
- Coming soon: *AnnData* objects directly

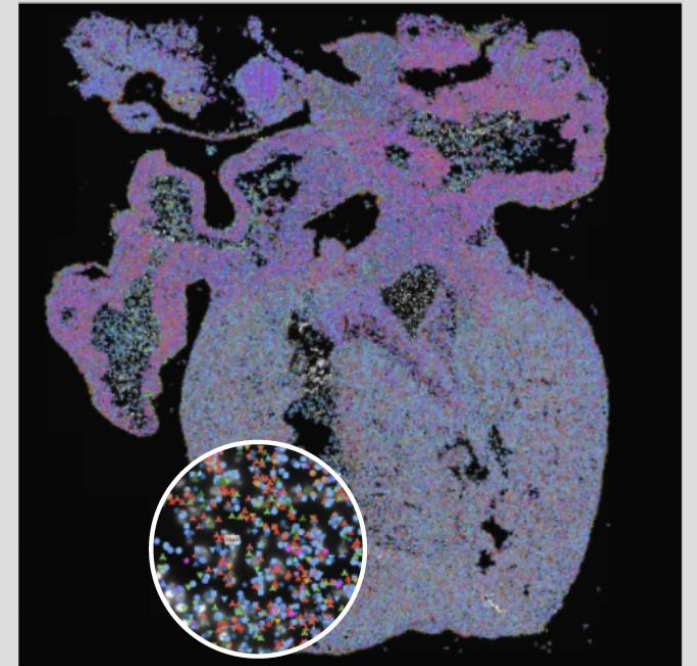
Image stack



Regions

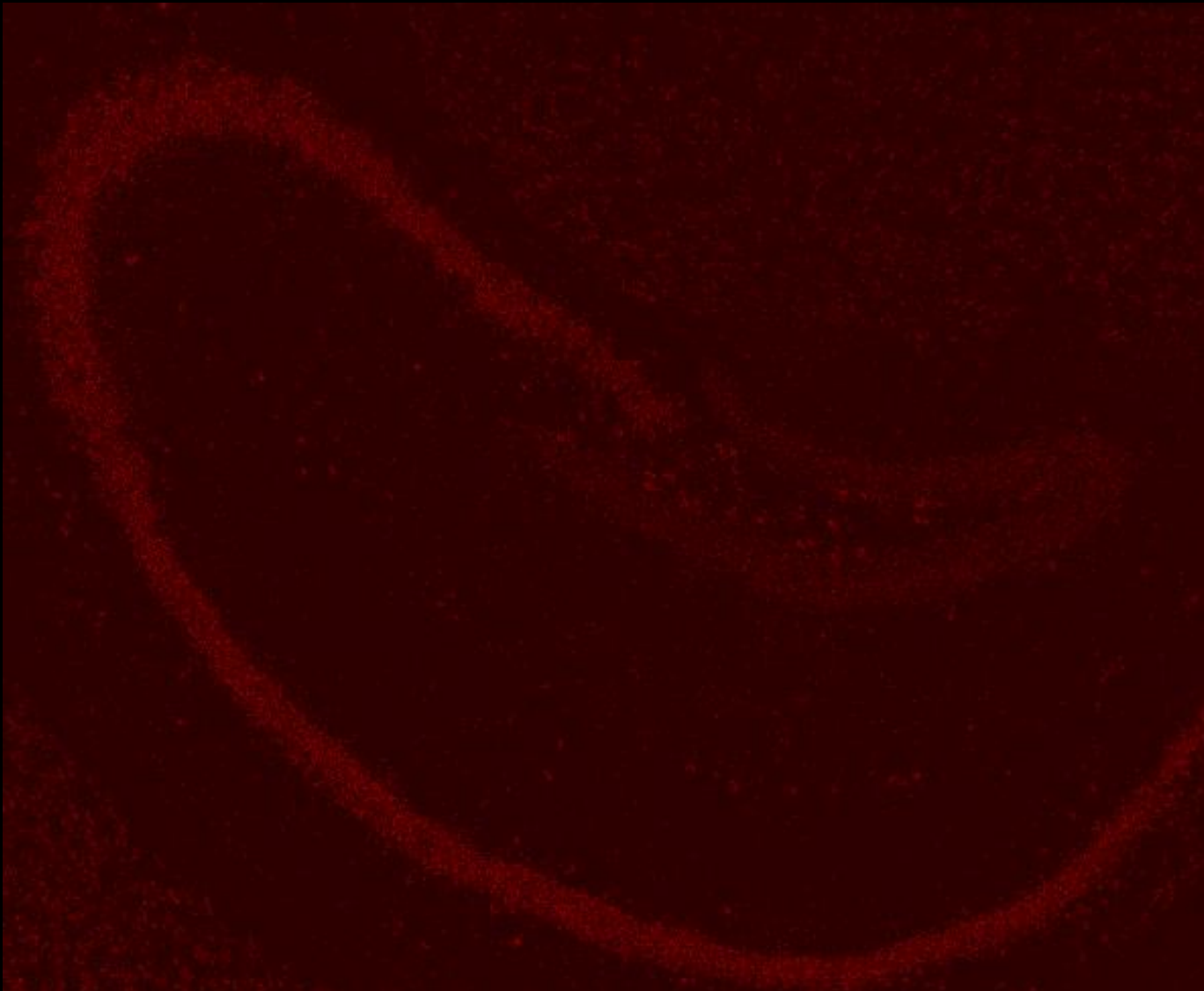


Markers



2D images

8-bit, (of tissue), any modality, multiple layers



The image shows a 2D medical scan, likely a histological section, with a prominent, bright, curved boundary. The image is displayed in a dark red color, which is the result of the '6.tif' layer being visible in the 'Image layers' panel.

The right side of the image shows a control panel with the following sections:

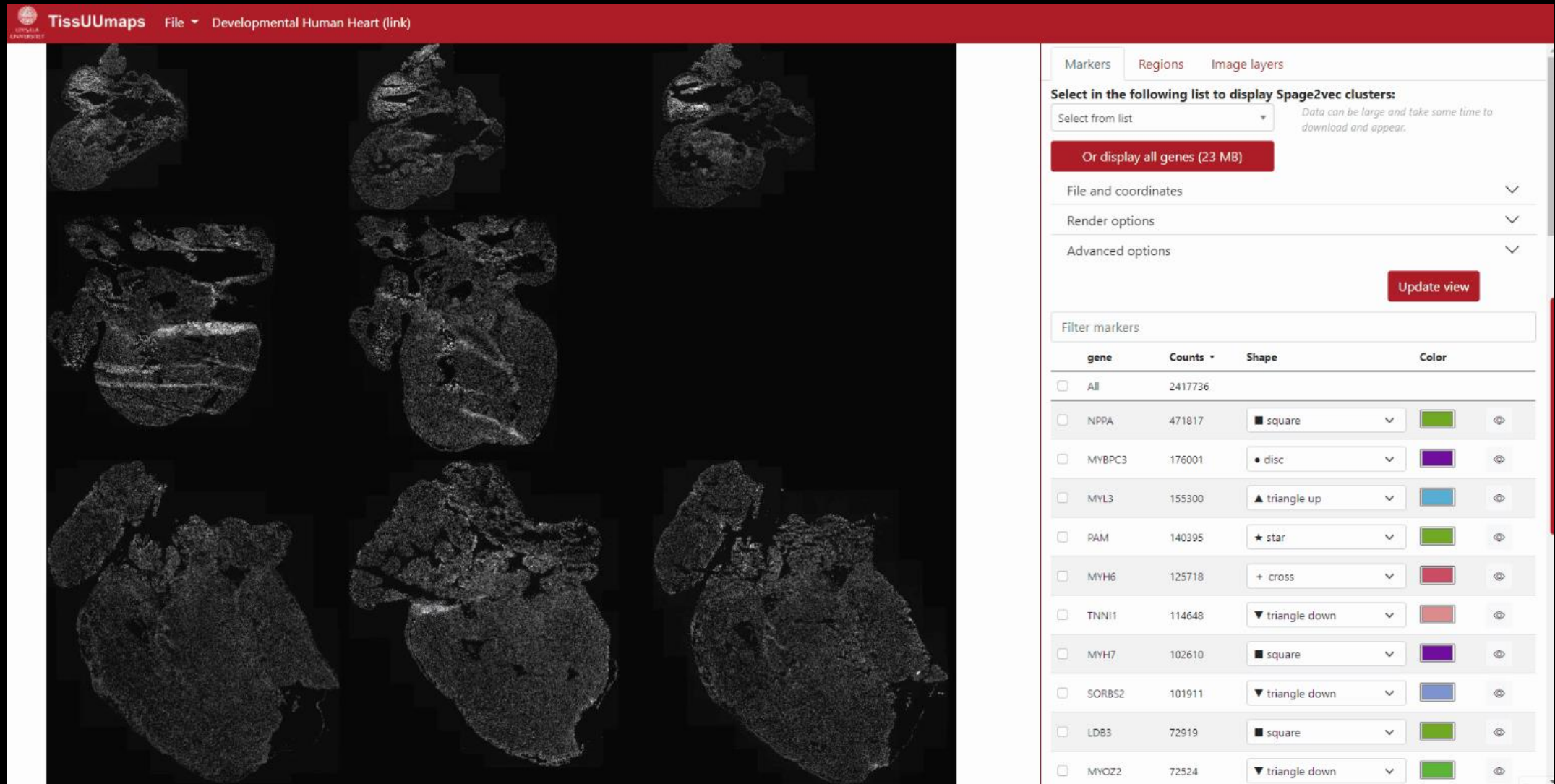
- Markers** (selected)
- Regions**
- Image layers**

Name	Visible	Opacity	Saturation	Brightness	Color
6.tif ⓘ	<input checked="" type="checkbox"/>	<input type="range"/>	<input type="range"/>	<input type="range"/>	Red
5.tif ⓘ	<input type="checkbox"/>	<input type="range"/>	<input type="range"/>	<input type="range"/>	Blue
3.tif ⓘ	<input type="checkbox"/>	<input type="range"/>	<input type="range"/>	<input type="range"/>	Green
2.tif ⓘ	<input type="checkbox"/>	<input type="range"/>	<input type="range"/>	<input type="range"/>	Gray

Below the table is a section for **Filter settings**, which is currently collapsed (indicated by a downward arrow).

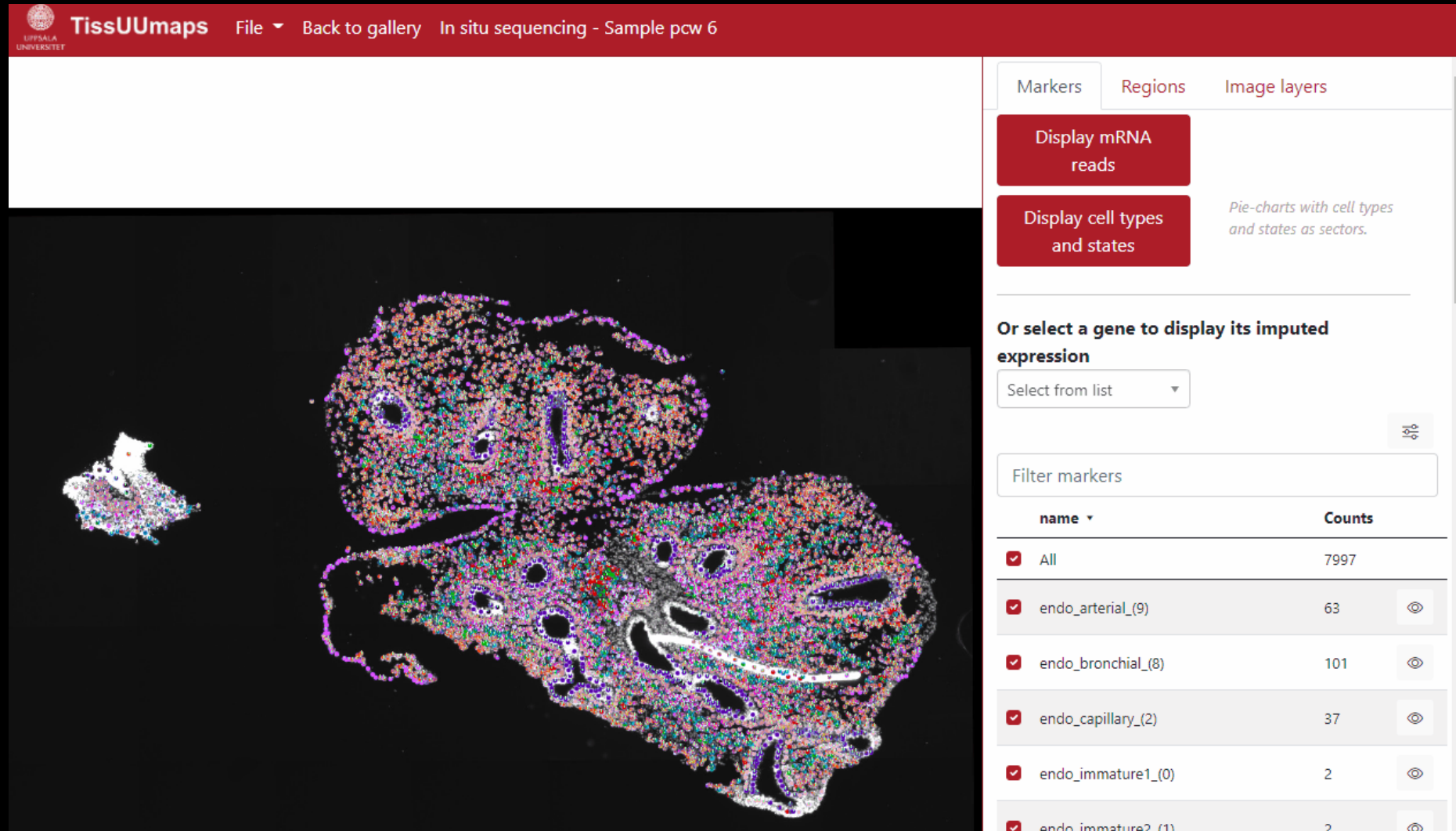
Markers

Markers, with x- and y coordinates, type and value, or even advanced info such as pie-charts (input as .csv file, could be generated from AnnData)



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Regions

Regions, drawn manually or defined using tools such as QuPath, Cellpose etc or defined using a TissUUmaps plugin (GeoJSON)

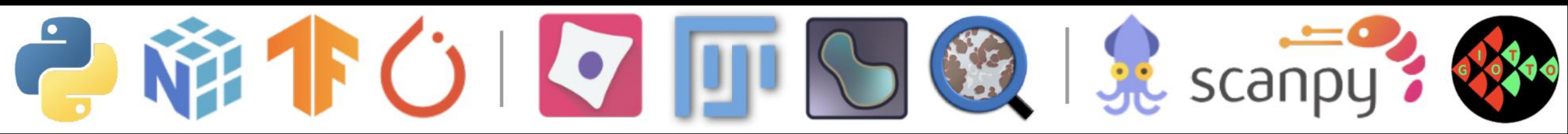


Regions

Regions, drawn manually or defined using tools such as QuPath, Cellpose etc or defined using a TissUUmaps plugin (GeoJSON)



Compatibilities



We aim for compatibility with other open source initiatives rather than building all functionality into TissUMaps.

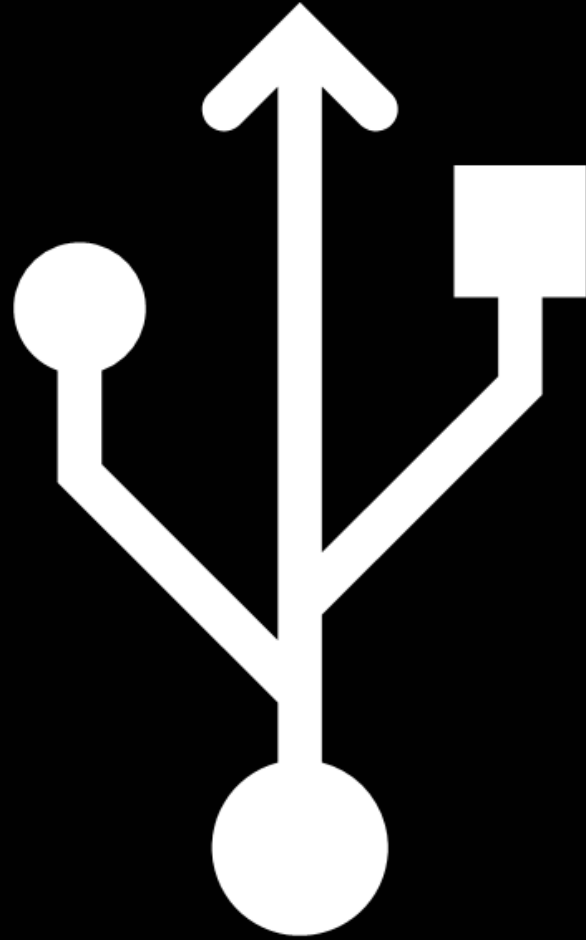
TissUMaps can be incorporated into Jupyter Notebooks to create modular data flows – see tutorials at tissuumaps.github.io

DEMO

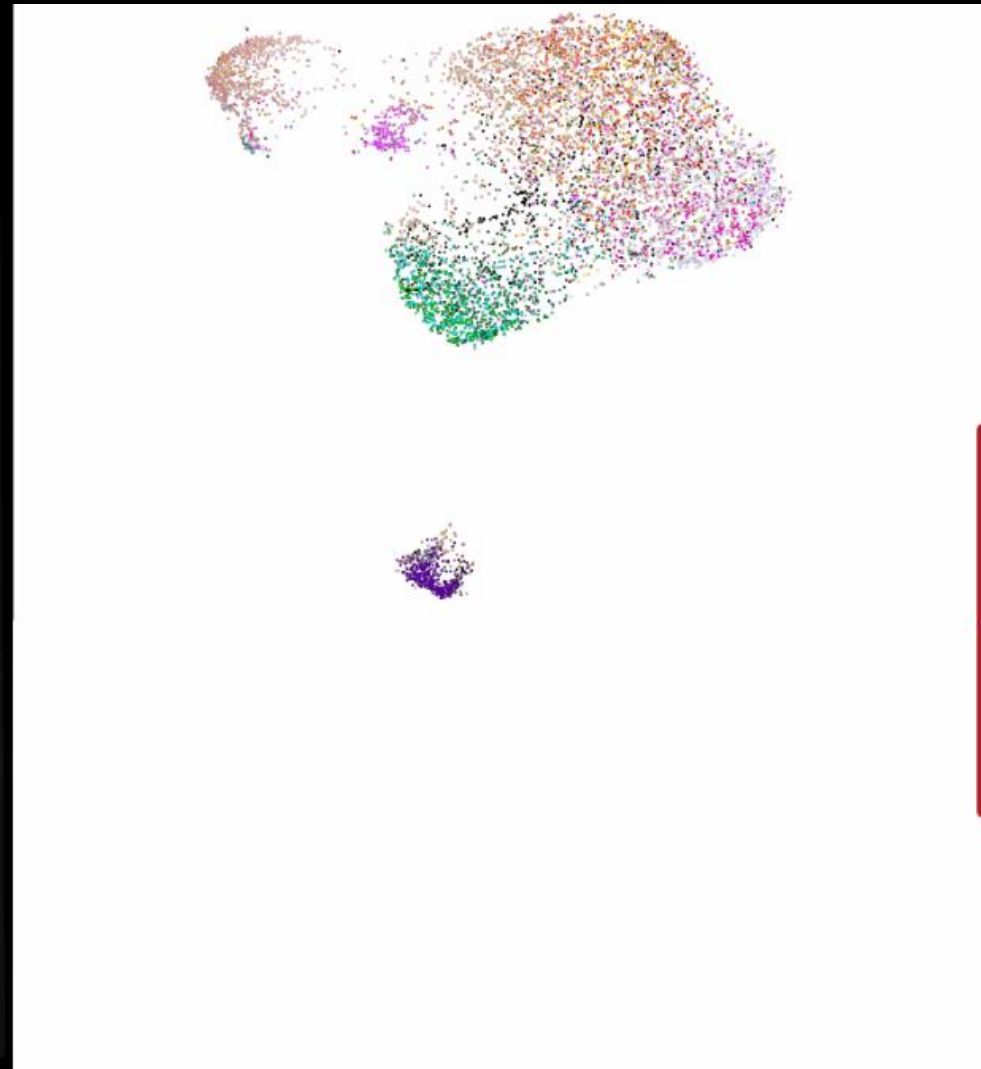
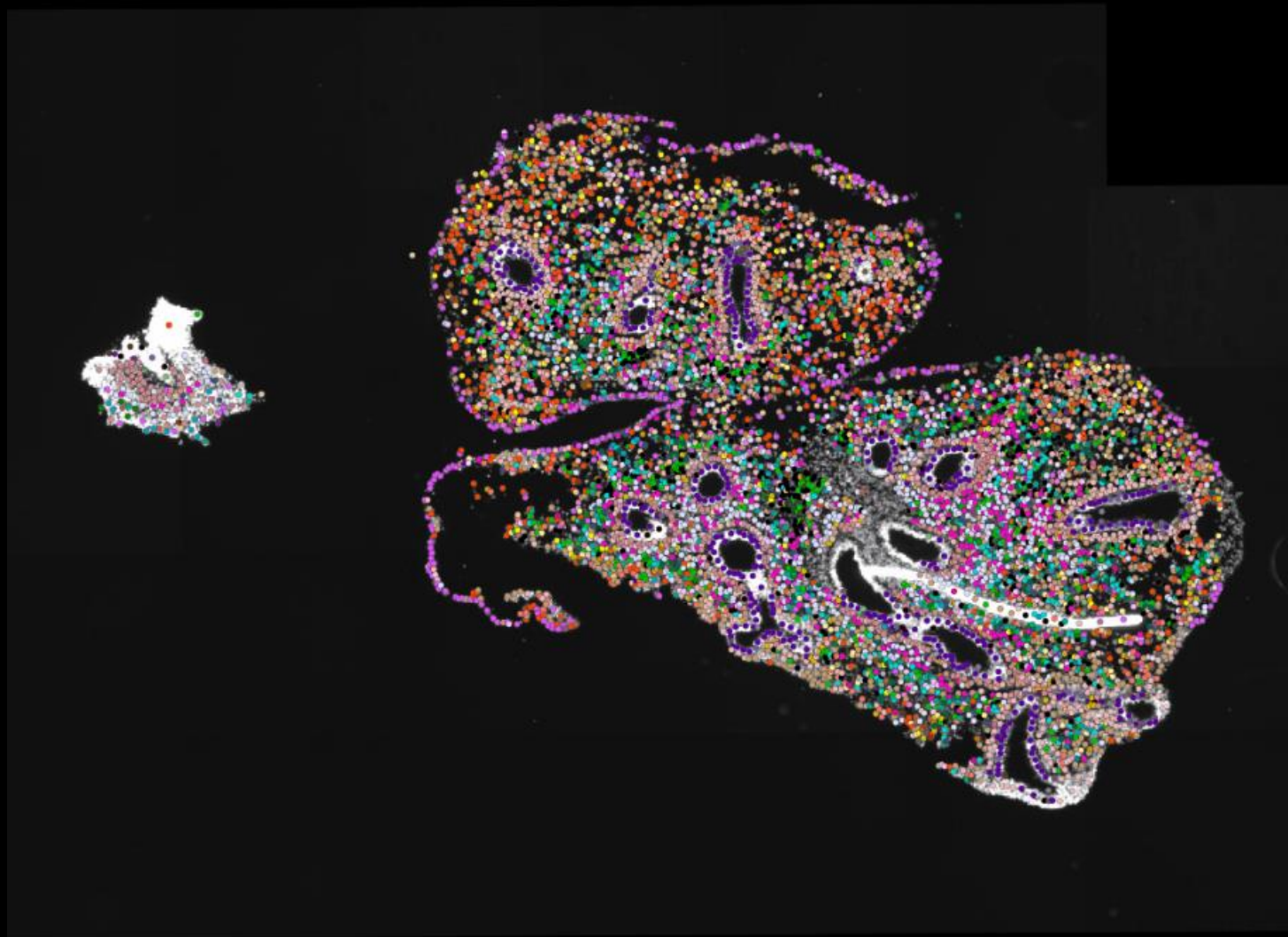
- <https://tissuumaps.github.io>

TissUUmaps plugins

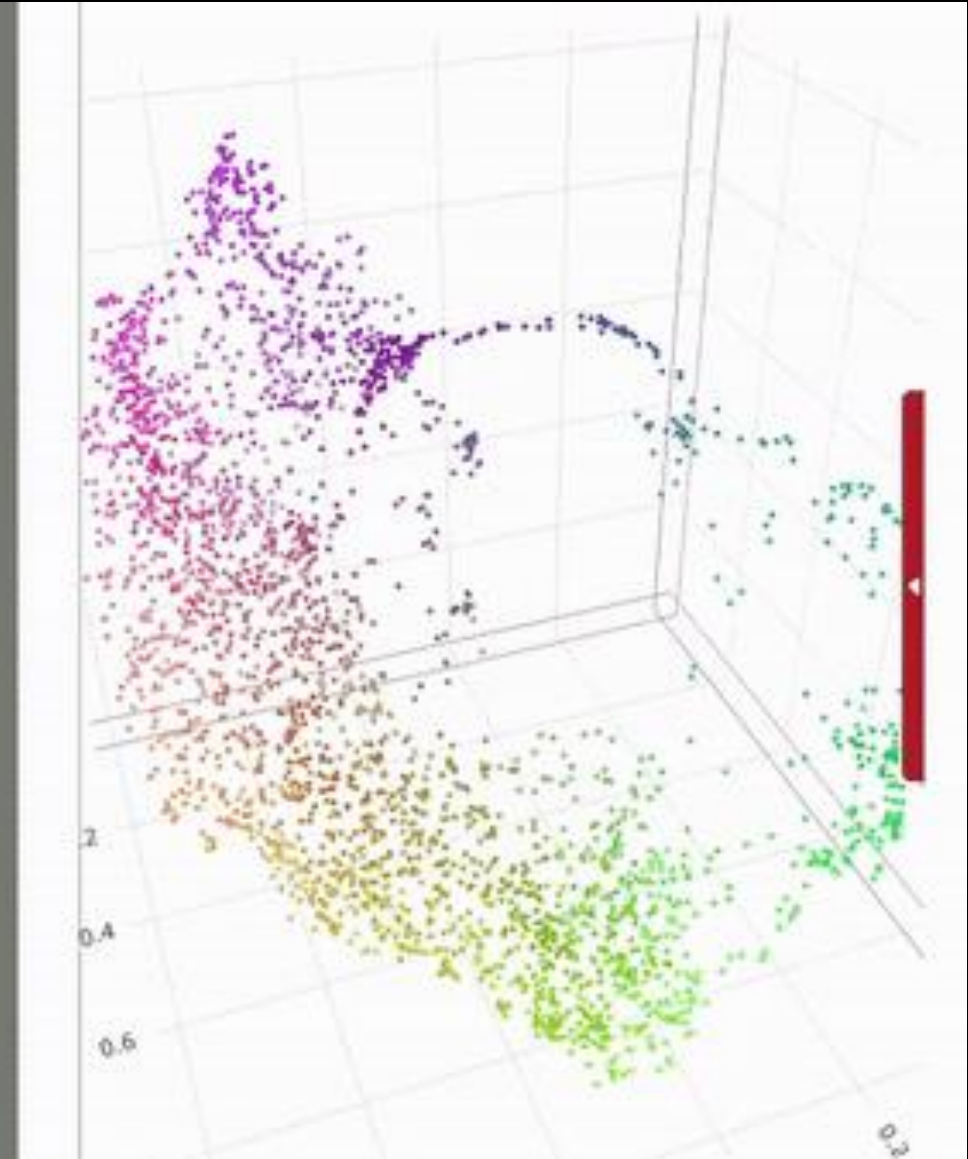
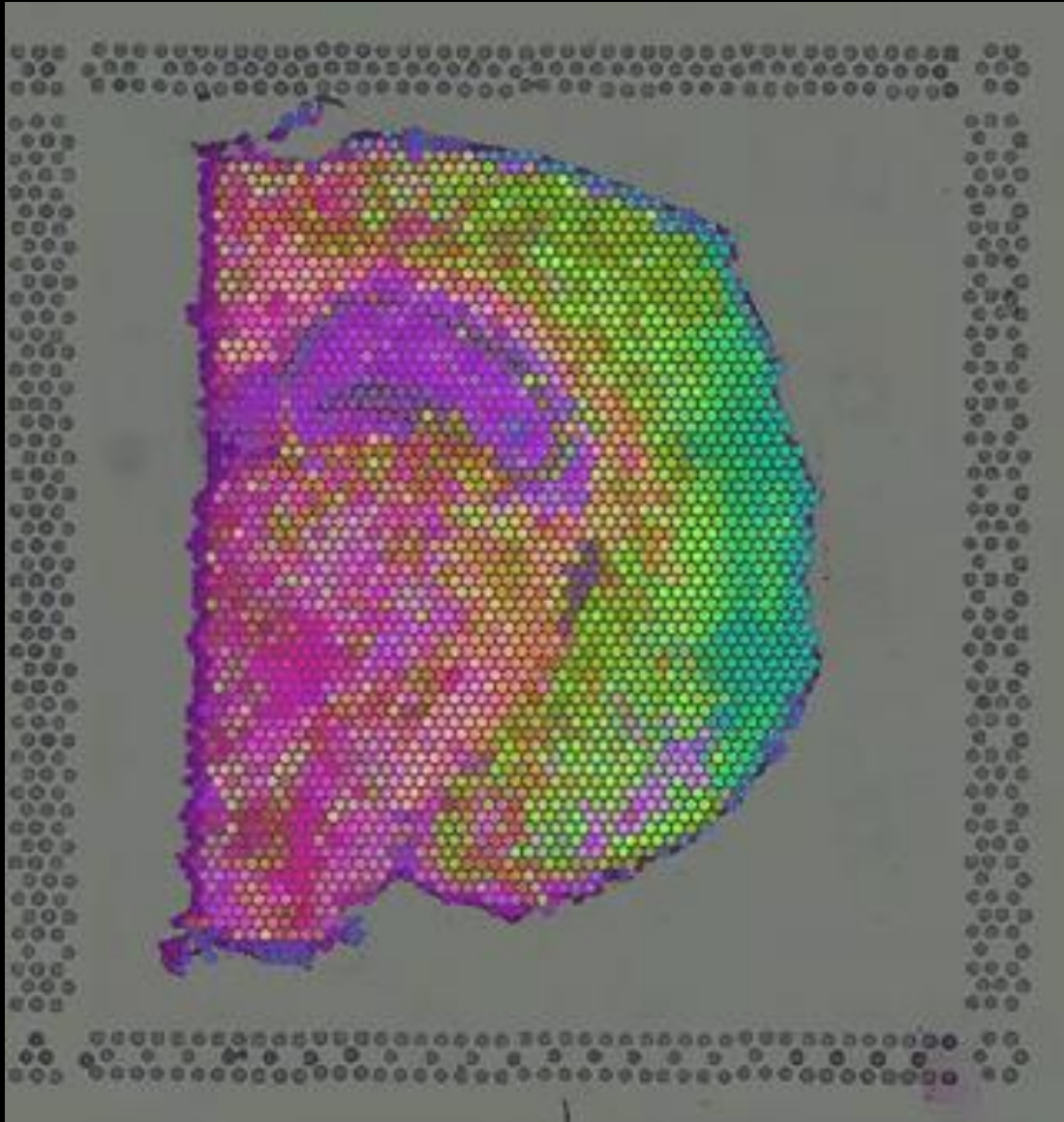
- Feature Space 2D
- Feature Space 3D
- Points2Regions
- Spot Inspector



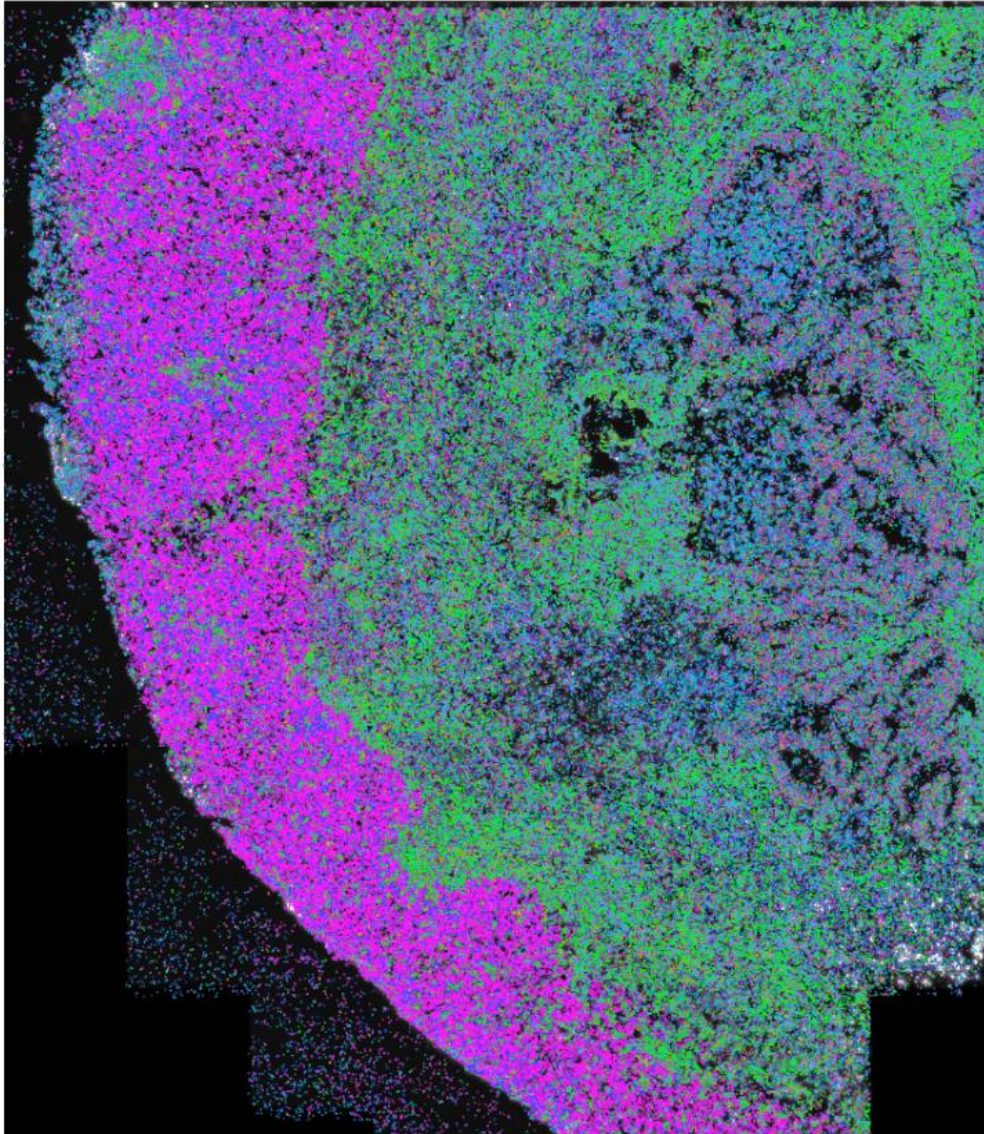
Feature Space 2D



Feature Space 3D



Points2Regions



Markers

Regions



























Image layers

Plugins

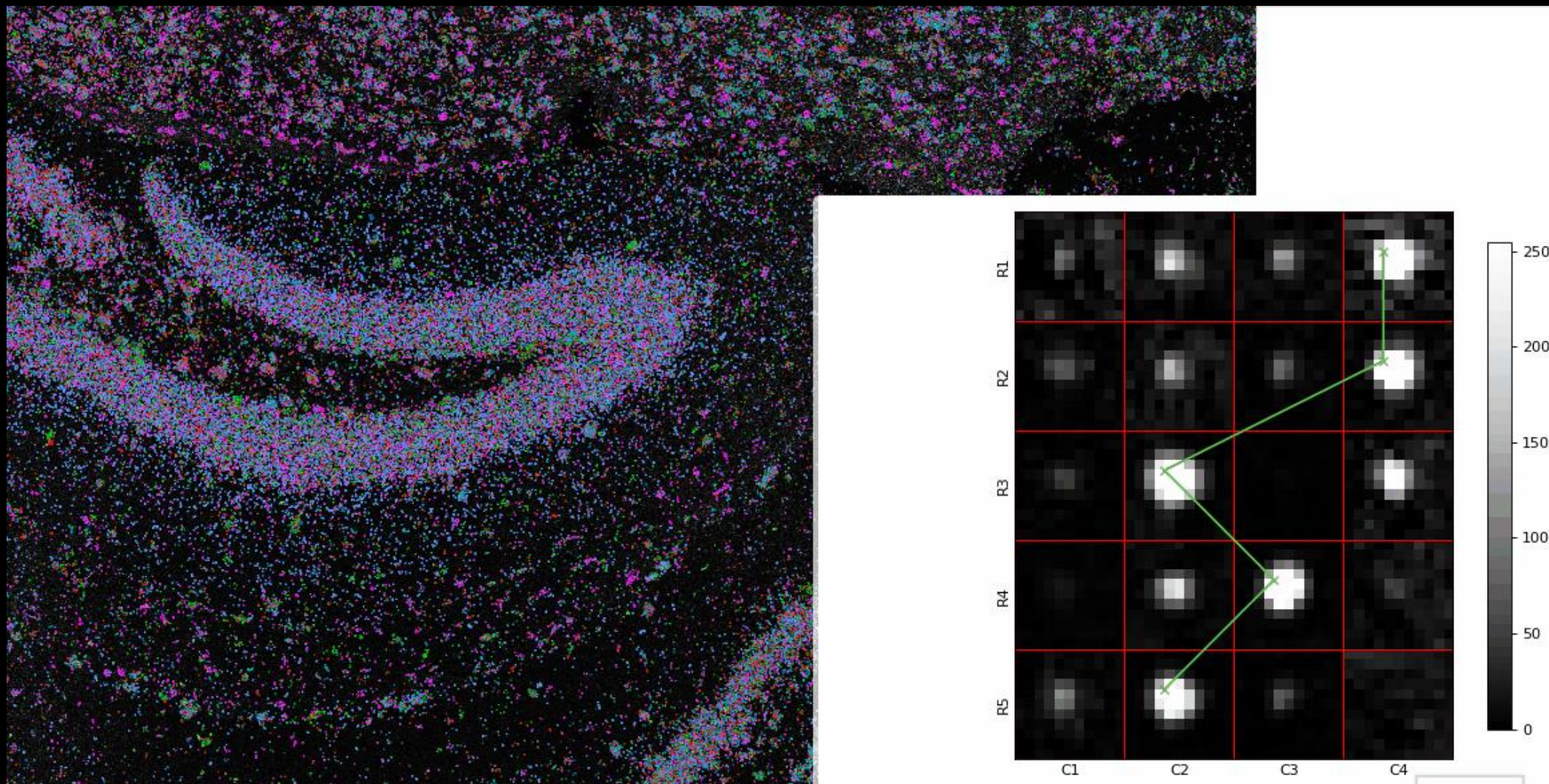
Download data

New markers × +

Filter markers

name ▾	Counts	Shape	Color	
<input checked="" type="checkbox"/> All	561485			
<input checked="" type="checkbox"/> ADAMDEC1	477	• disc ▾		
<input checked="" type="checkbox"/> ALBUMIN	178367	+ cross ▾		
<input checked="" type="checkbox"/> APC	804	■ square ▾		
<input checked="" type="checkbox"/> ARPC2	60	◆ diamond ▾		
<input checked="" type="checkbox"/> ARPC3	3184	▲ triangle up ▾		
<input checked="" type="checkbox"/> AXIN1	716	★ star ▾		
<input checked="" type="checkbox"/> AXIN2	1143	▼ triangle down ▾		
<input checked="" type="checkbox"/> BMP3B	134	⊗ clobber ▾		
<input checked="" type="checkbox"/> CALD1	49	• disc ▾		
<input checked="" type="checkbox"/> CALM3	1058	+ cross ▾		
<input checked="" type="checkbox"/> CAMK2A	1034	■ square ▾		
<input checked="" type="checkbox"/> CAMK2G	1852	◆ diamond ▾		
<input checked="" type="checkbox"/> CAV1	126	▲ triangle up ▾		

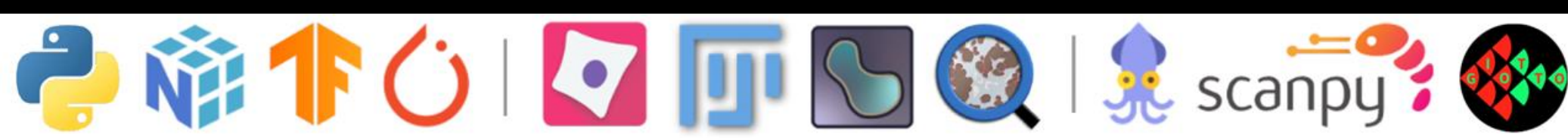
Spot Inspector



Conclusion

- TissUMaps is primarily a web-based interactive viewer for large-scale 2D images, regions, and markers.
- We collaborate with other open source initiatives for cell and tissue segmentation and spatial statistics, and aim for compatibility rather than building all functionality into TissUMaps
- Please reach out to us if you have ideas for how your tools can be combined with TissUMaps.

Examples and video tutorials on compatibilities with...



... can be found at website <https://tissuumaps.github.io>

