

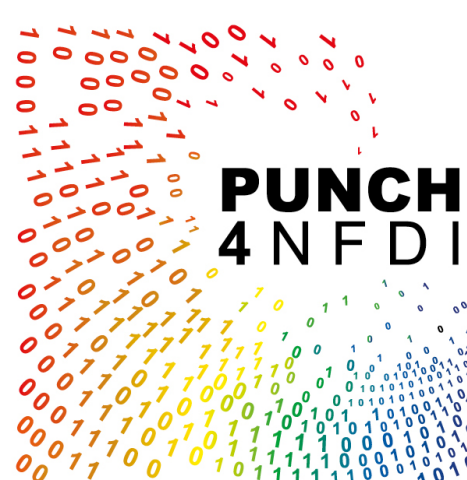
SciTrace

A Digital Research Product manager

Nicola Malavasi (LMU Munich)

In collaboration with: Yori Fournier, Kirill Makan, Anastasia Galkin, Olaf Michaelis, Harry Enke

PUNCH Young Academy Tutorial - 11/10/2023



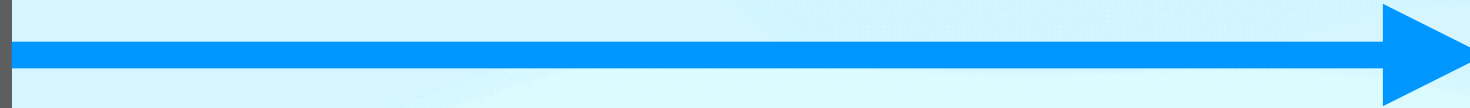
From abstract DRP to real world implementation

Several pieces of software need to be brought together to go from an abstract DRP idea to a prototype. Each piece addresses a specific problem.

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Code installation



Docker container +
installation
metadata

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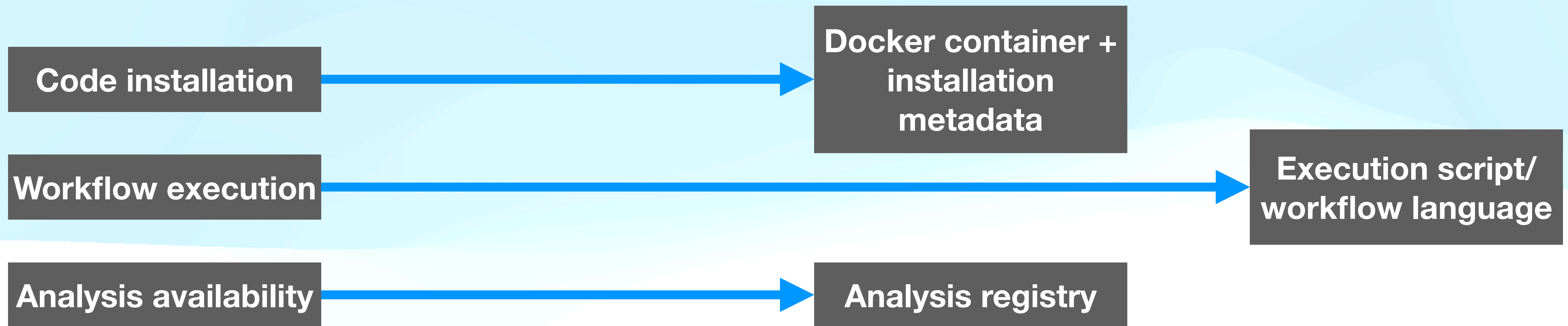
Workflow execution

Execution script/
workflow language

```
graph LR; A[Code installation] --> B[Docker container + installation metadata]; C[Workflow execution] --> D[Execution script/ workflow language];
```


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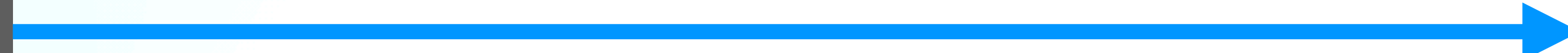
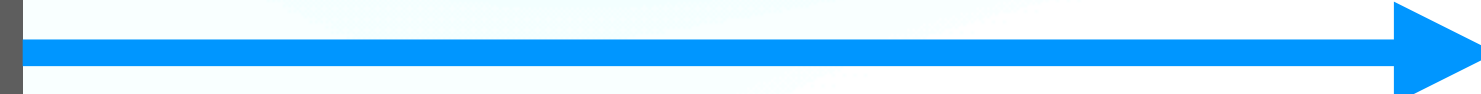
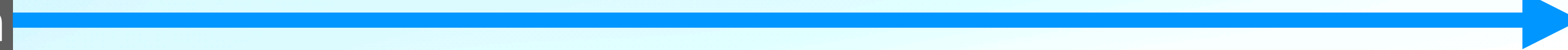
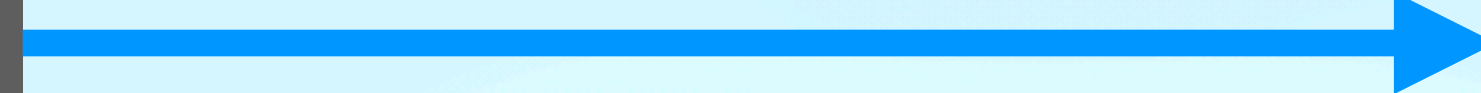
Execution script/
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Analysis availability

Analysis registry

Reproducibility/
traceability

Use of hashes/
integration with
GitLab



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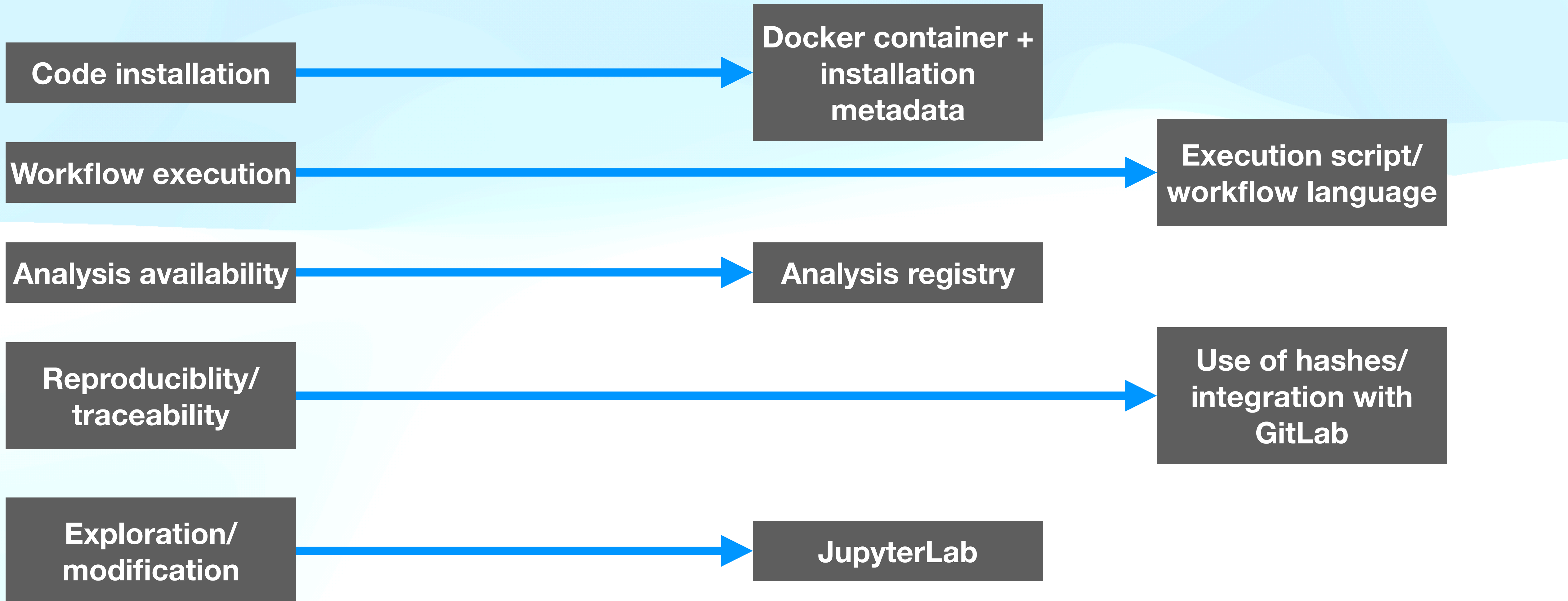
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Use of hashes/
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Exploration/
modification

JupyterLab



SciTrace

Having established the abstract concept of a DRP and a check list of software needed to build a prototype, we can move forward.

Prototype software created at AIP Potsdam by Yori Fournier, Kirill Makan, Anastasia Galkin, Olaf Michaelis: **SciTrace**

The DRP in the SciTrace formalism

Necessary elements

In the SciTrace formalism, a DRP is a package formed by a structure hosted in an existing [GitLab repository](#).

This repository can be ingested by the [SciTrace program](#), that can use its parts to create the DRP.

Name	Last commit	Last update
📁 env_disperse	Added readmes, renamed setup-env.sh ...	3 months ago
📁 env_python3.9	Added readmes, renamed setup-env.sh ...	3 months ago
📁 step0_get_data	Added readmes, renamed setup-env.sh ...	3 months ago
📁 step1_run_delaunay_3d	Added readmes, renamed setup-env.sh ...	3 months ago
📁 step2_run_mse	fixed bug in step2 parameters with nam...	3 months ago
📁 step3_analysis_and_plots	Updated parameters and fixed bug in ru...	3 months ago
📄 README.md	Update step1_run_delaunay_3d/paramet...	1 year ago
⚙️ setup-env-disperse.toml	Created setup-step-3.toml file	1 year ago
⚙️ setup-env-python3.9.toml	adapted the packages to the new Packa...	1 year ago
⚙️ setup-step-0.toml	adapted the packages to the new Packa...	1 year ago
⚙️ setup-step-1.toml	adapted the packages to the new Packa...	1 year ago
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The DRP in the SciTrace formalism

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Package repository

Necessary requirements











- **Data folder:** contains the input
- **Parameters folder:** contains the parameters for the analysis (toml file)
- **Products folder:** will contain the output
- Execution script: **drun.sh**
- **Analysis scripts**
- Installation scripts: **install.sh**, **install-deps.sh**, and **install-user-deps.sh**

Package folder

Necessary requirements

Data folder Parameters folder Products folder Execution script: `drun.sh` Analysis scripts

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Installation

install-deps.sh

This script installs the dependencies necessary for the code.

Treated as sequence of bash commands, it is transformed in an ad-hoc Dockerfile to generate a container.

Example of install-deps.sh for DisPerSE

```
install-deps.sh 594 B
Edit Replace Delete

1 # make sure apt do not prompt interactively
2 export DEBIAN_FRONTEND=noninteractive
3
4 # apt need access to /tmp
5 chmod 777 /tmp
6
7 # apt update is not necessary (done in base image)
8 apt-get update
9
10 # install some deps
11 apt-get install -y \
12     cmake \
13     wget
14
15 # disperse libs
16 apt-get install -y \
17     libboost-all-dev \
18     libgsl-dev \
19     libcgall-dev \
20     libcfitsio-dev
21
22 # Add symbolic link to solve xlocale.h to locale.h problem
23 ln -s /usr/include/locale.h /usr/include/xlocale.h
24
25 # Add a symbolic link for Delaunay3D (somehow it links to -lBoost instead of -lboost)
26 ln -s /usr/include/boost /usr/include/Boost
```

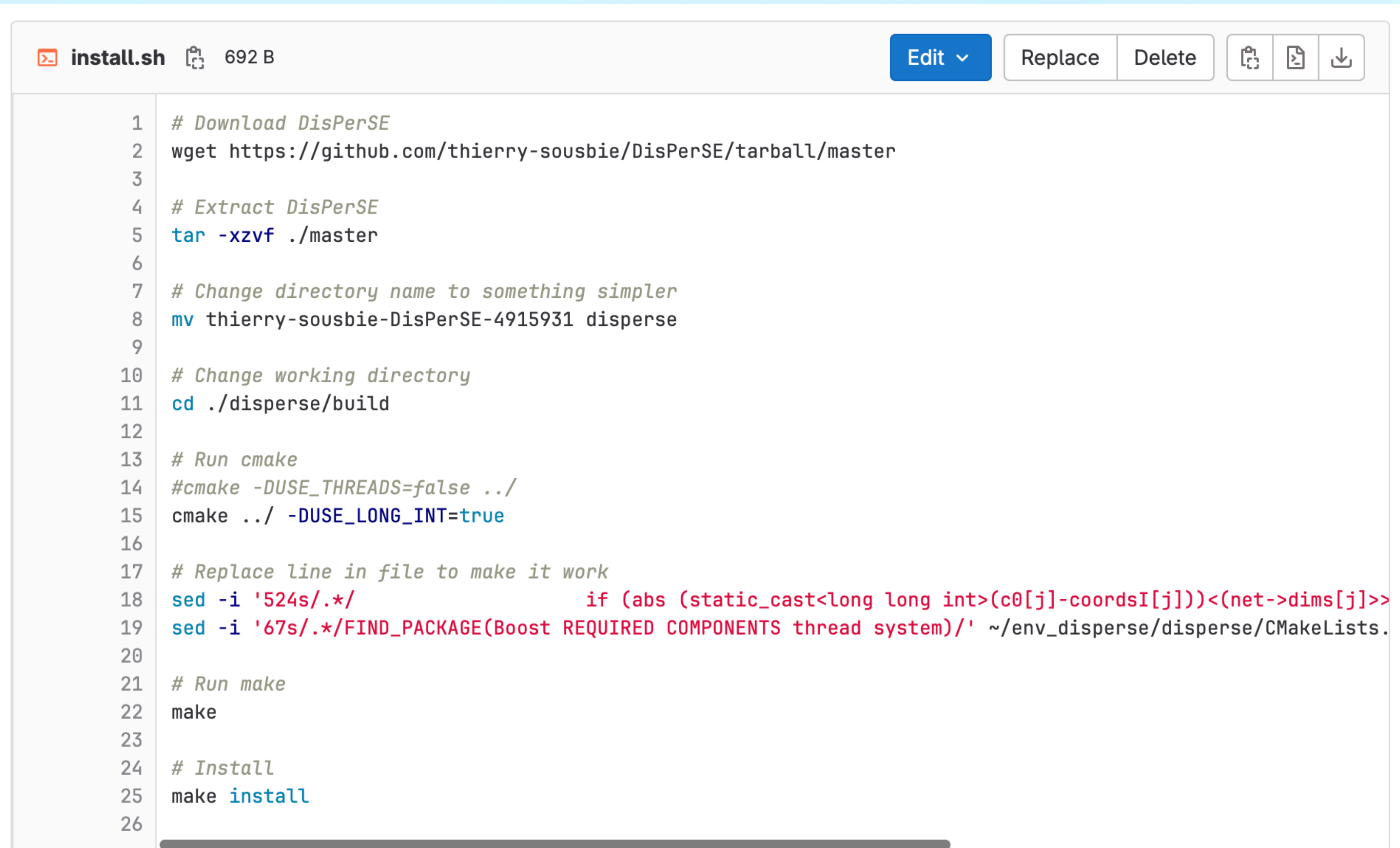
Installation

install.sh

This script installs the necessary code.

Treated as sequence of bash commands, it is transformed in an ad-hoc Dockerfile to generate a container.

Example of install.sh for DisPerSE



```
install.sh 692 B
Edit Replace Delete

1 # Download DisPerSE
2 wget https://github.com/thierry-sousbie/DisPerSE/tarball/master
3
4 # Extract DisPerSE
5 tar -xzvf ./master
6
7 # Change directory name to something simpler
8 mv thierry-sousbie-DisPerSE-4915931 disperse
9
10 # Change working directory
11 cd ./disperse/build
12
13 # Run cmake
14 #cmake -DUSE_THREADS=false ../
15 cmake ../ -DUSE_LONG_INT=true
16
17 # Replace line in file to make it work
18 sed -i '524s/./ if (abs (static_cast<long long int>(c0[j]-coordsI[j]))<(net->dims[j]>>
19 sed -i '67s/./FIND_PACKAGE(Boost REQUIRED COMPONENTS thread system)/' ~/env_disperse/disperse/CMakeLists.
20
21 # Run make
22 make
23
24 # Install
25 make install
26
```


Installation

install-user-deps.sh

This script activates the python environment for exploration and installs the necessary code.

Treated as sequence of bash commands, it is transformed in an ad-hoc Dockerfile to generate a container.

 **install-user-deps.sh**  141 B

Edit ▾

Replace

Delete



```
1  # activate the virtual env
2  . ~/env/bin/activate
3
4  # install the requirements
5  pip install -r ~/step3_analysis_and_plots/setup/requirements.txt
```

Example of install-user-deps.sh for a python environment

Run

Analysis scripts

Name
..
parameters
products
.gitkeep
README.md
drun.sh
format_from_survey.py
prepare_for_disperse.py

These are scripts created by the user that perform the analysis.

They are executed as they are within the container created by the installation procedure.

Run drun.sh

The analysis scripts are called by the drun.sh script which is executed inside the container.

The diagram illustrates the workflow for running analysis scripts. On the left, a file explorer shows a directory structure with files like `parameters`, `products`, `.gitkeep`, `README.md`, and `drun.sh`. A red arrow points from `drun.sh` to a code editor on the right. The code editor displays the contents of `drun.sh`, which is a shell script that activates a Python environment and runs two analysis scripts: `format_from_survey.py` and `prepare_for_disperse.py`. These two scripts are highlighted with a red box. Another red arrow points from this box back to the file explorer, where these two scripts are also listed and highlighted with a red box.

```
drun.sh 168 B
1  #!/bin/bash
2
3  # activate the python environment
4  . ~/env/bin/activate
5
6  # run the pre formatting script
7  python -U format_from_survey.py
8  python -U prepare_for_disperse.py
```

format_from_survey.py

prepare_for_disperse.py

Example of analysis scripts for the get_data step

Parameters

Toml file

```
22
23 # Cosmological parameters: if false, Planck Collaboration et al. 2015, Paper XIII cosmology is used. If set
24 # H0
25 H0 = false
26
27 # Omega matter
28 Om0 = false
29
30 # Omega Lambda
31 Olambda0 = false
32
33
34 # Center of the field. If false defaults to (0,0).
35 # RA center
36 ra_center = 186.183
37
38 # Dec center
39 dec_center = 26.845
40
41
42 # Names of the columns with the quantities. These should come from the catalogue information (e.g. paper, r
43 # RA column
44 rc = "ra"
45
46 # Dec column
47 dc = "dec"
48
49 # Redshift column
50 zc = "zfinal"
```

These are parameters that can be used in an analysis by the analysis routines. They are looked for in the parameters folder.

SciTraceWeb

DRP creation and (re-)use in a user friendly way

What is SciTraceWeb?

SciTraceWeb is an instance of SciTrace running at AIP, accessible through a web page.

It allows efficient and user-friendly DRP creation, as well as possibilities for DRP manipulation such as:

- **Exploration**: a DRP can be accessed and its content inspected but not modified.
- **Run**: a DRP is executed.
- **Modification**: a DRP can be accessed, its content modified, and a new DRP is created, the difference between the two is recorded by the system.

Creation

- DRP creation starts with the creation of a GitLab repository.
- The structure of the repository is fixed, with the necessary files found at the correct position.
- A given DRP can be based on a previous one. In that case it will have access to the previous's code and environment.
- Input data can be mounted in the data folder. They can be the products of a previous step.
- The scripts `install.sh`, `intall-deps.sh`, and `install-user-deps.sh` are run automatically.

DRP Run

- Right after creation a DRP has no products. To create the products it must be run.
- Running happens within the container.
- The script `drun.sh` is executed automatically.
- It then runs the analysis scripts.
- Running the container uses resources.

Exploration

- Package exploration is possible thanks to a JupyterLab instance installed within the container.
- Data are accessible, so are products, and the analysis code.
- Exploration is performed in read-only mode.

Modification

- Modification is similar to exploration but it allows also writing.
- Several operations are possible: the code can be modified and run, data accessed, parameters modified.
- GitLab integration means that once the modifications are done they can be saved, pushed to a cloned version of the repository and a new package created and run.
- Parameters can also be downloaded, modified, and re-uploaded to generate a new DRP.

Tracing

- All the operations described before are traced via the hash of the container image.
- Package creation generates an hash.
- A new hash is generated for package run that indicates that products have been created.
- Package modification also generates a new hash, different from the starting one to indicate that the package has been modified and is different from the original.
- There is a DRP registry where generated DRPs are saved and can be explored/used as starting point by the community.

Practical example

The analysis of the cosmic web around the Coma cluster as detected by DisPerSE implemented in SciTraceWeb

Scientific analysis based on:

Malavasi et al. 2020a, A&A, 634, A30

Malavasi et al. 2020b, A&A, 642, A19

Malavasi et al. 2023, A&A, 675, A76

Implementation in PUNCH based on:

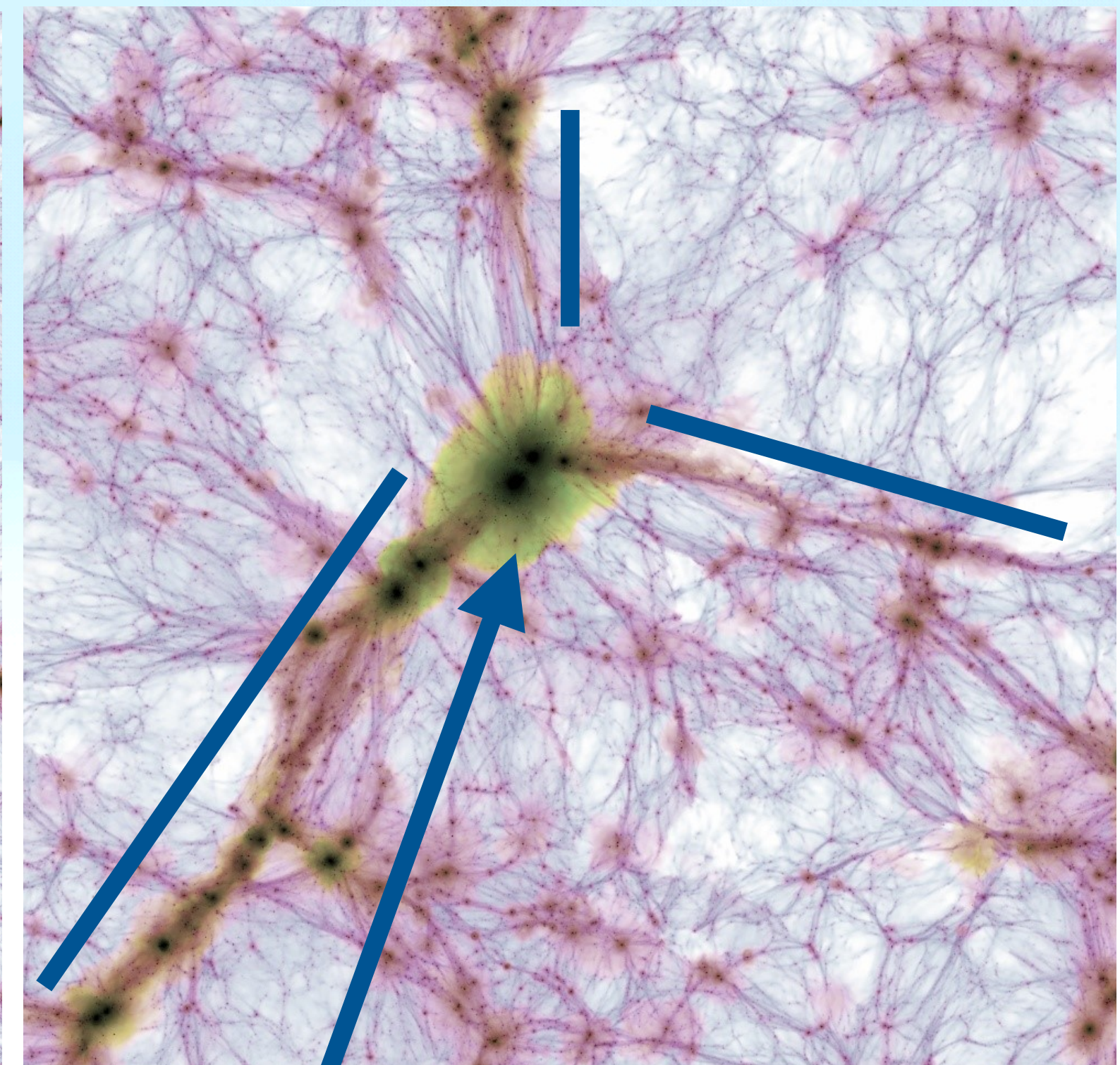
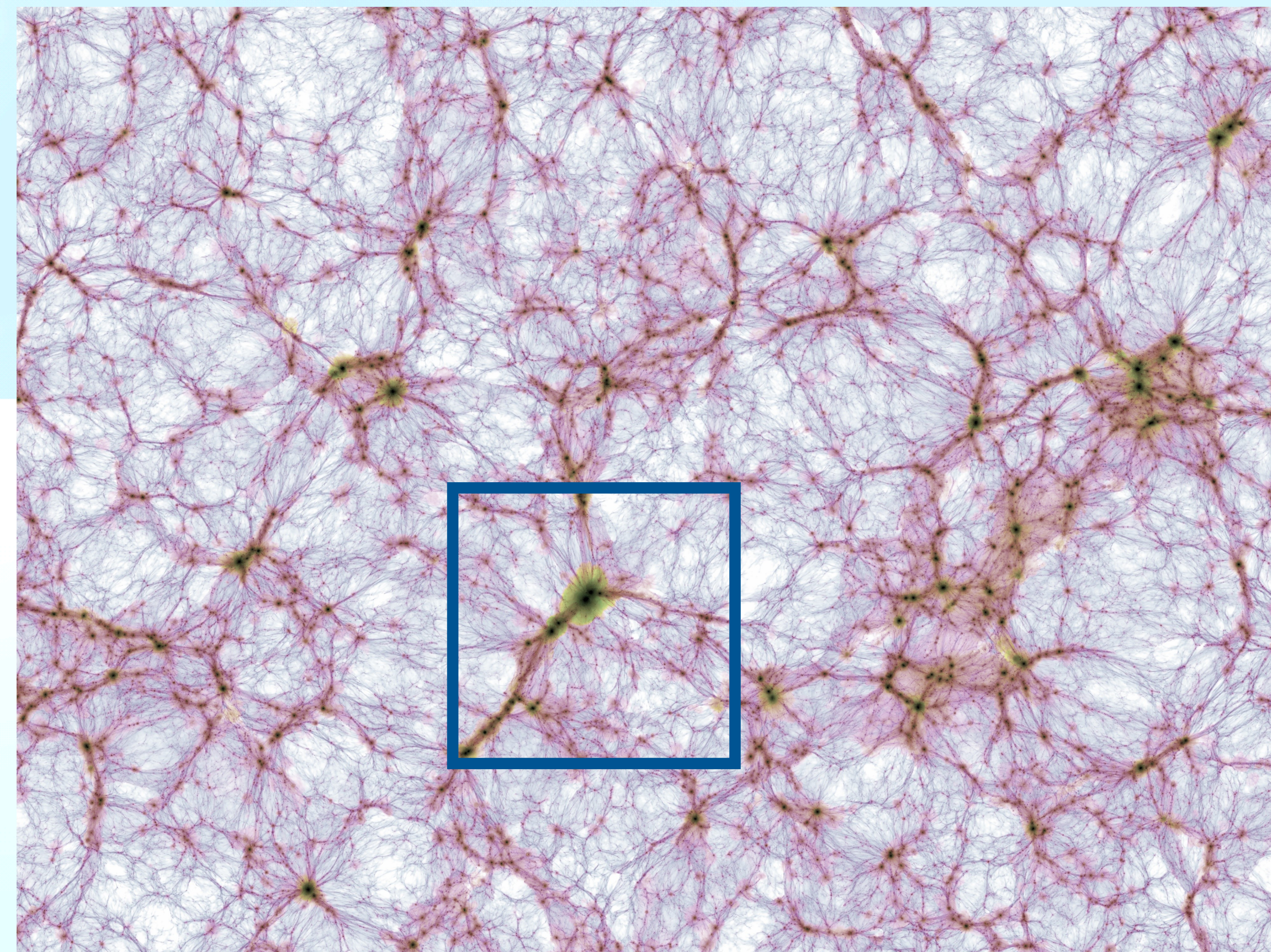
Fournier et al., in prep.

Malavasi et al., in prep.

The cosmic web

Network of connected structures made of galaxies, dark matter, and gas.

The cosmic web impacts the formation of structures and the formation of galaxies.

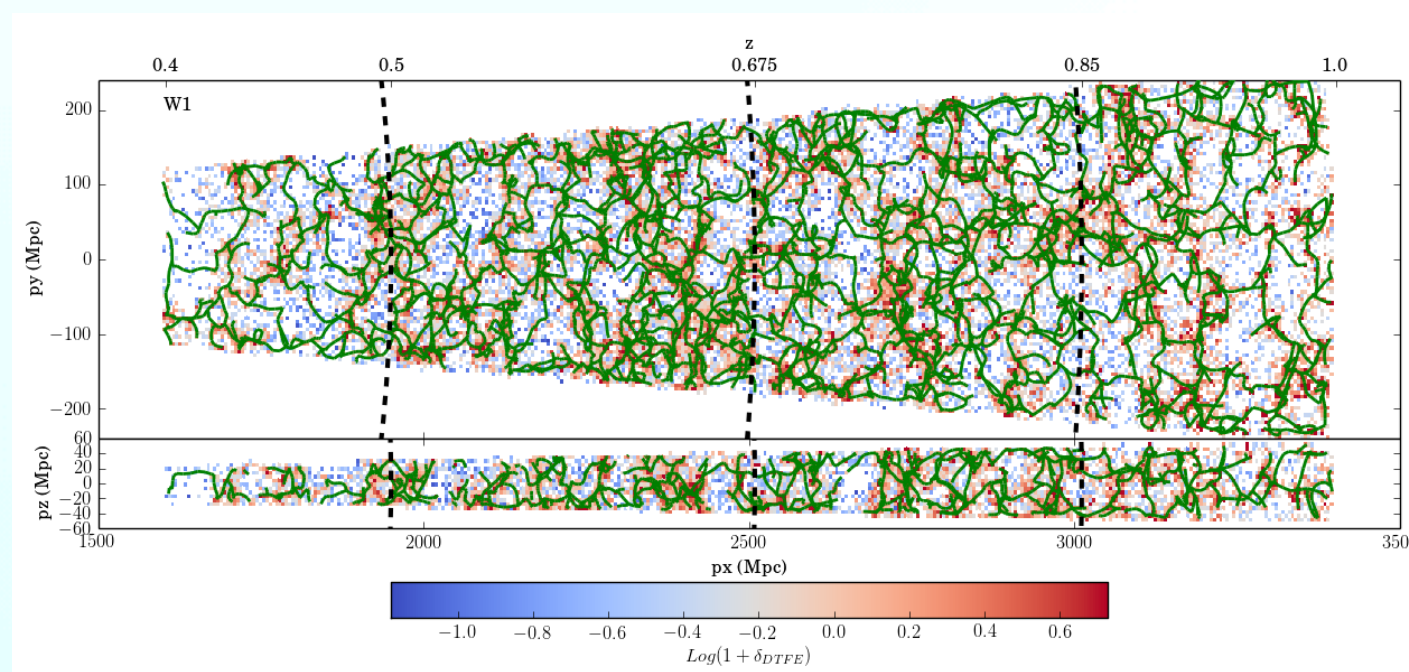


Clusters as nodes connected by filaments.

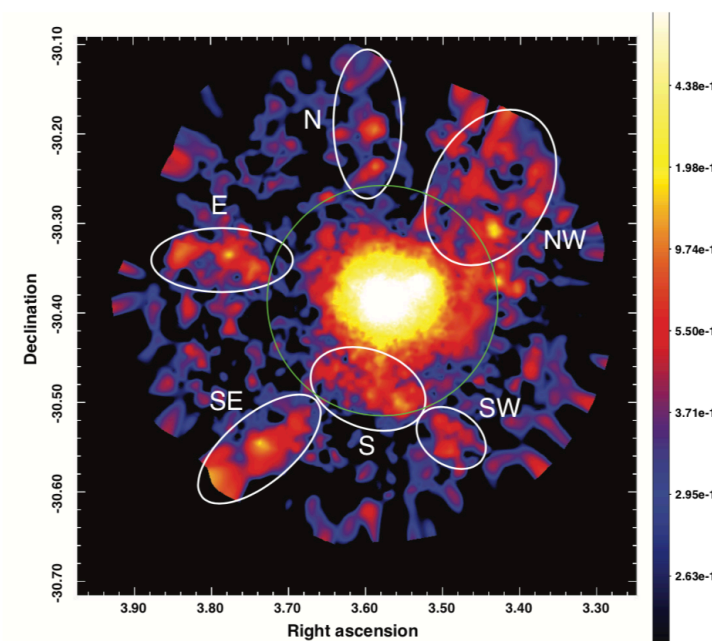
Our goal

- Study the connections and the connectivity of a massive, nearby, well known cluster.
- Study the LSS in a large region around a cluster.
- Perform a case study of a well known object.
- Study filament properties.

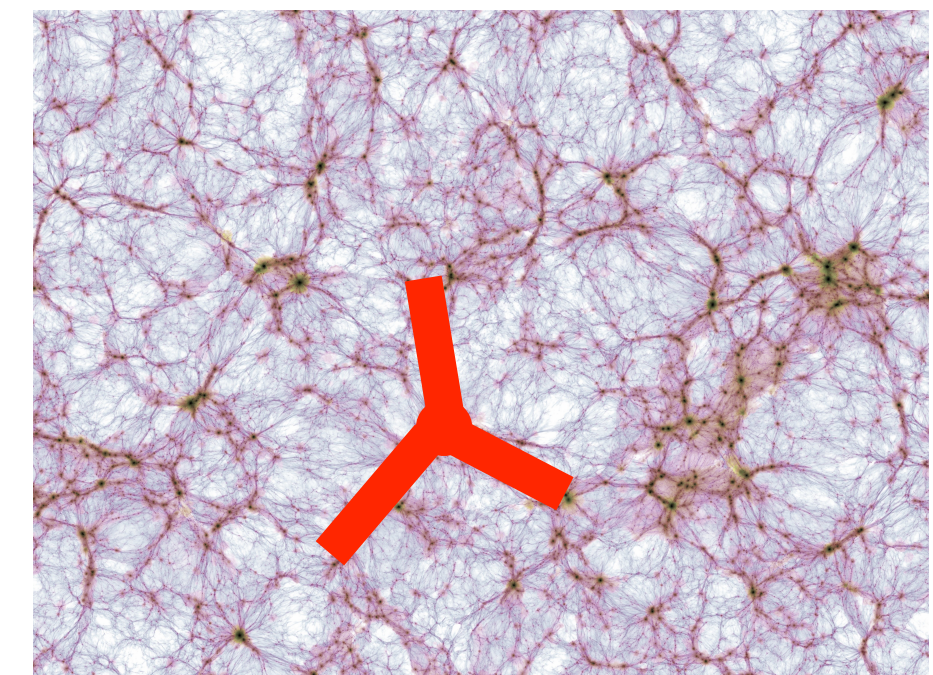
Apply a filament detection algorithm to a spectroscopic survey, then study the properties of the filaments detected around a cluster.



+



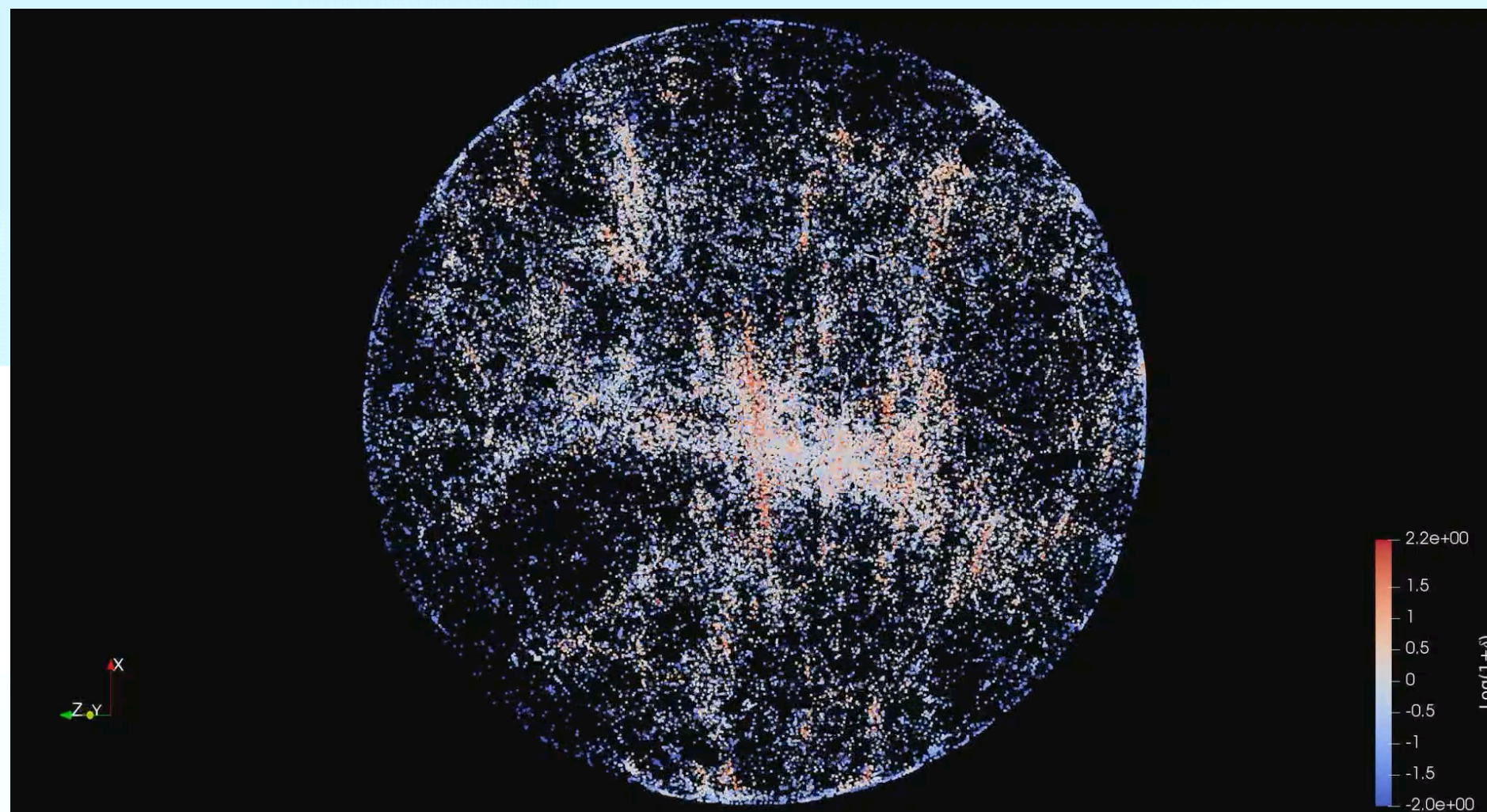
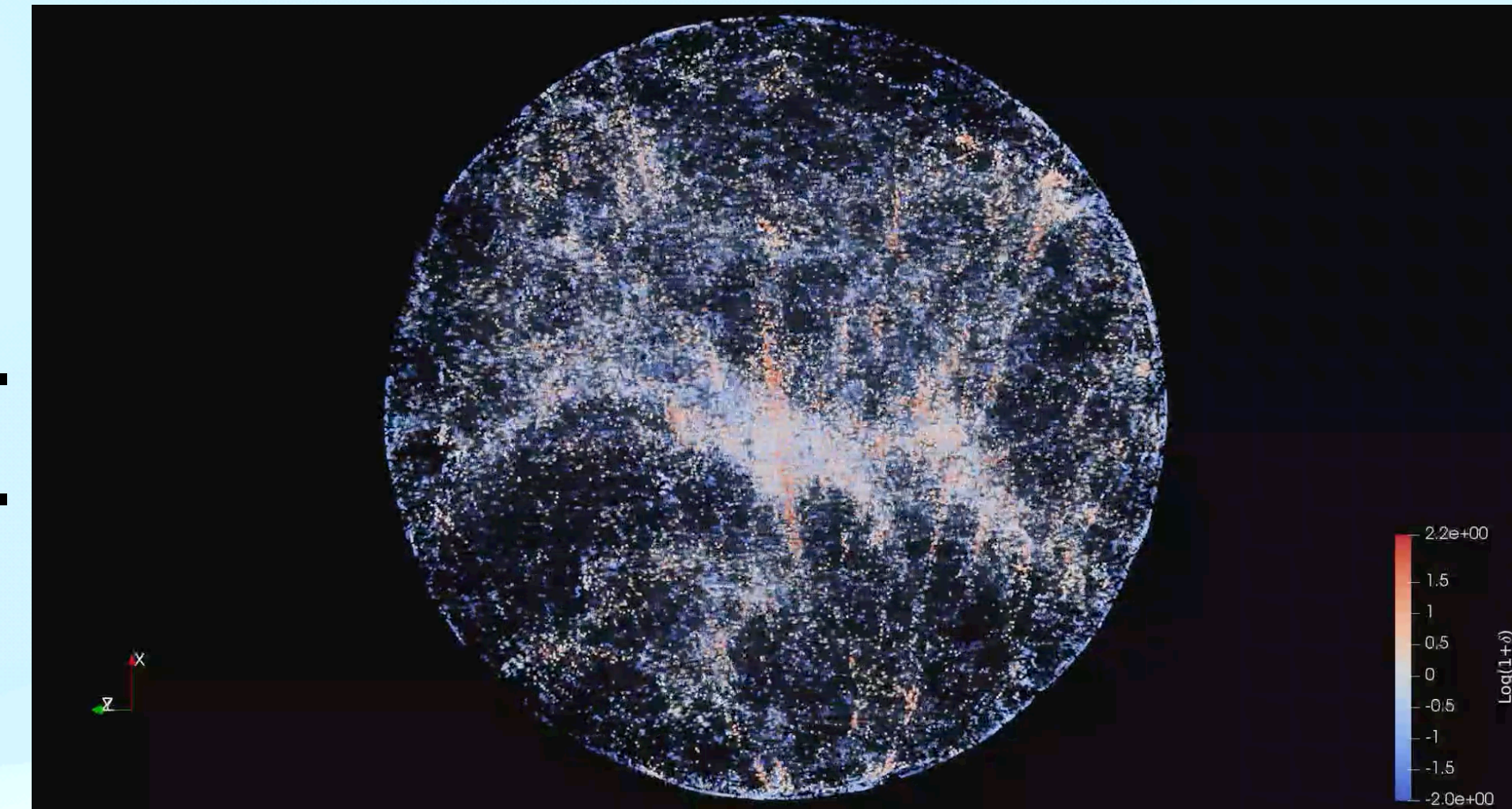
=



Discrete Persistent Structure Extractor

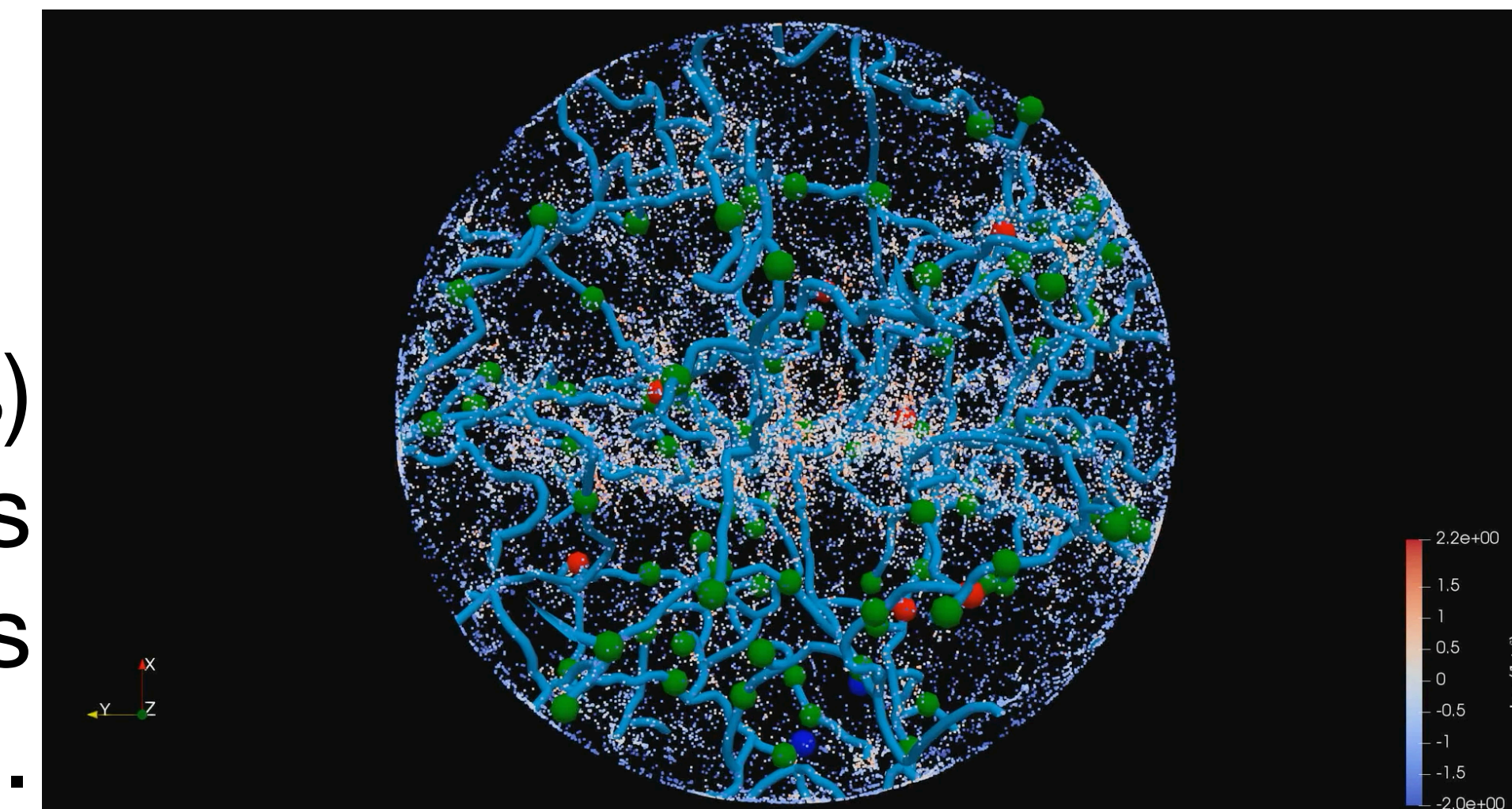
Sousbie 2011, Sousbie et al. 2011

Measure of the density field (DTFE).
Possibility of smoothing (although not necessary).



Computation of the discrete gradient.
Detection of the critical points (maxima, minima, and saddles).

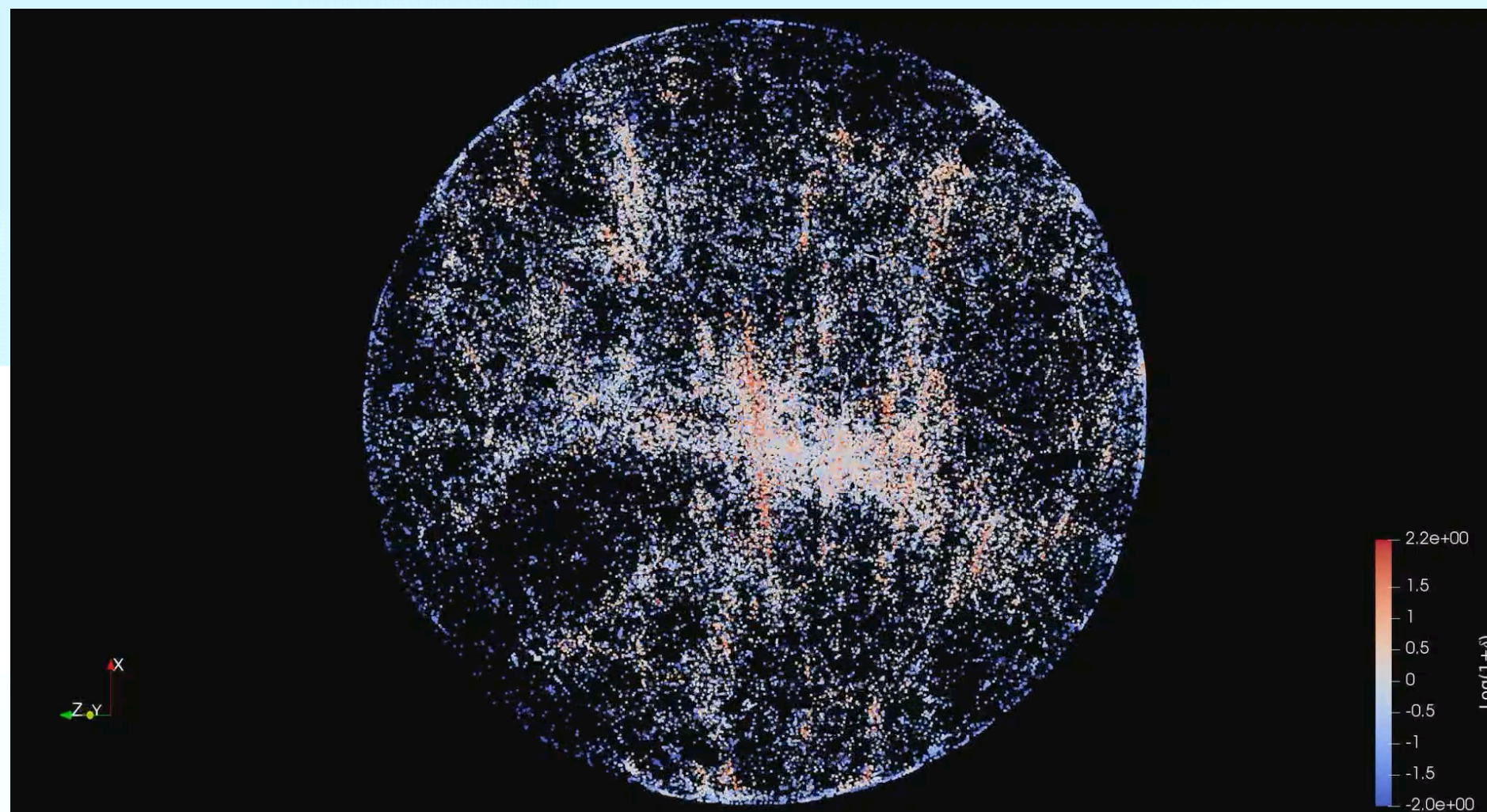
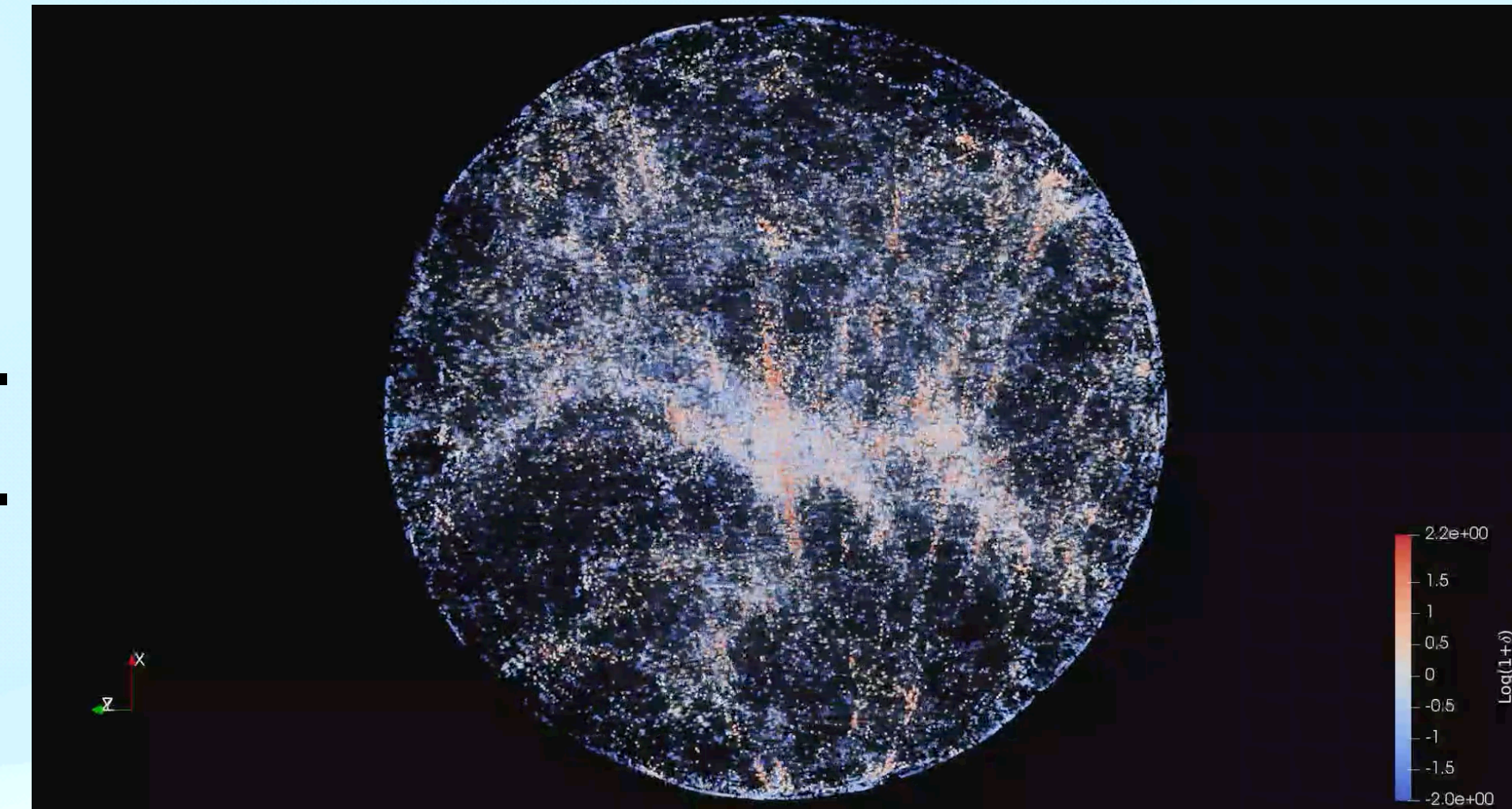
Connection of the critical points (maxima and saddles) with filaments. Persistence cut to eliminate spurious structures due to noise (expressed in terms of numbers of sigma, similar to S/N threshold).



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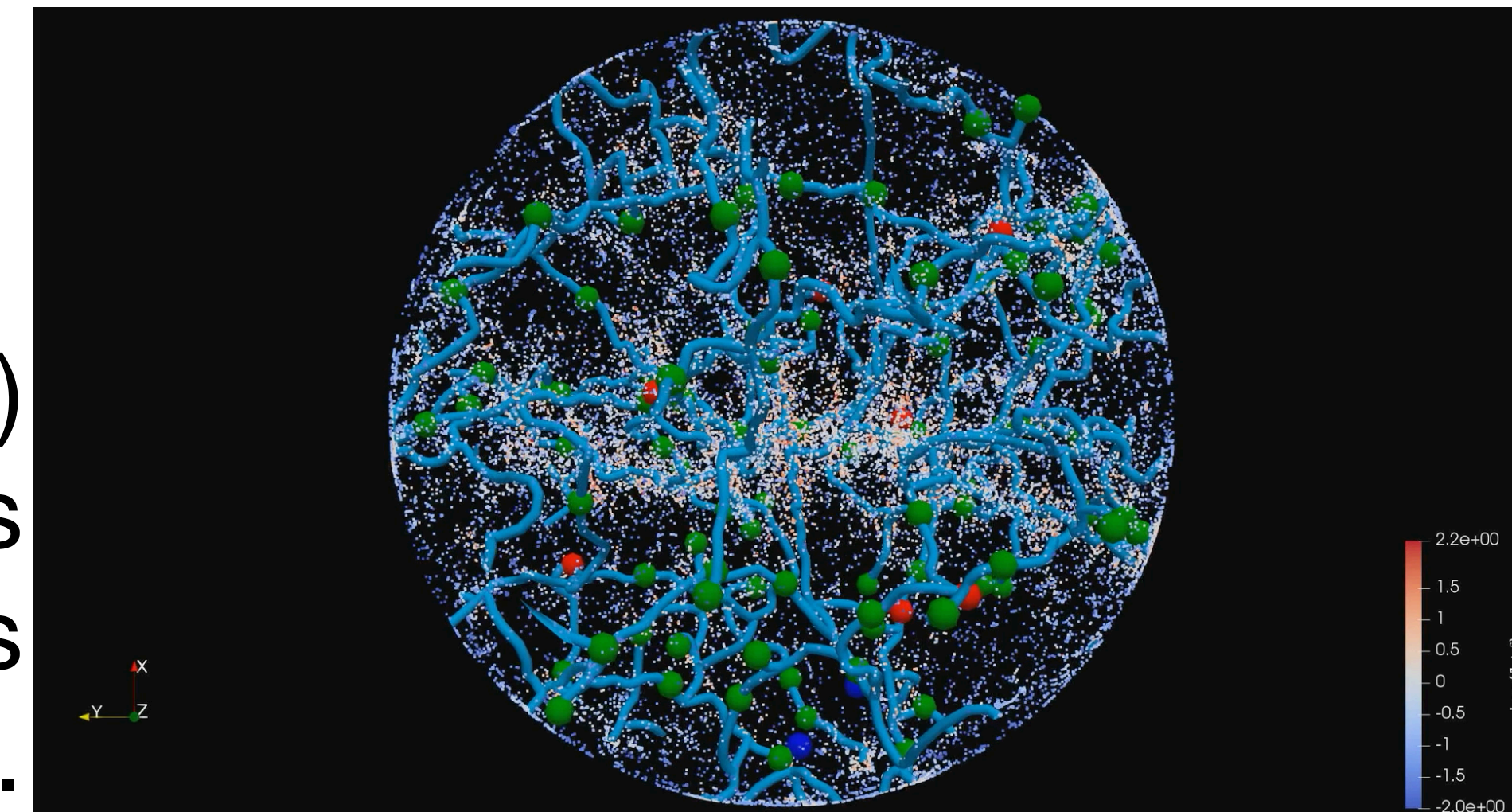
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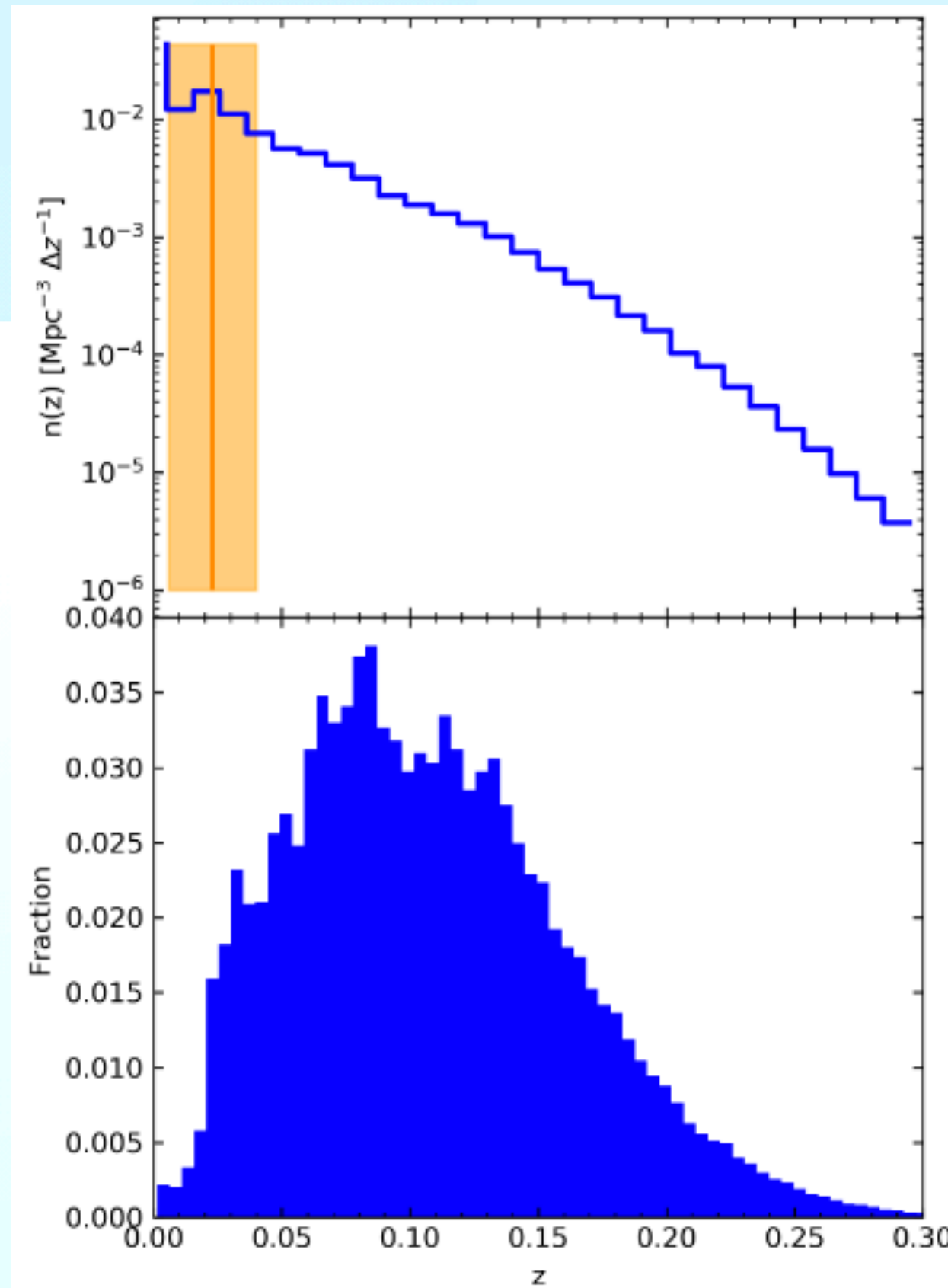
Connection of the critical points (maxima and saddles) with filaments. Persistence cut to eliminate spurious structures due to noise (expressed in terms of numbers of sigma, similar to S/N threshold).



Tracers of the filaments: SDSS

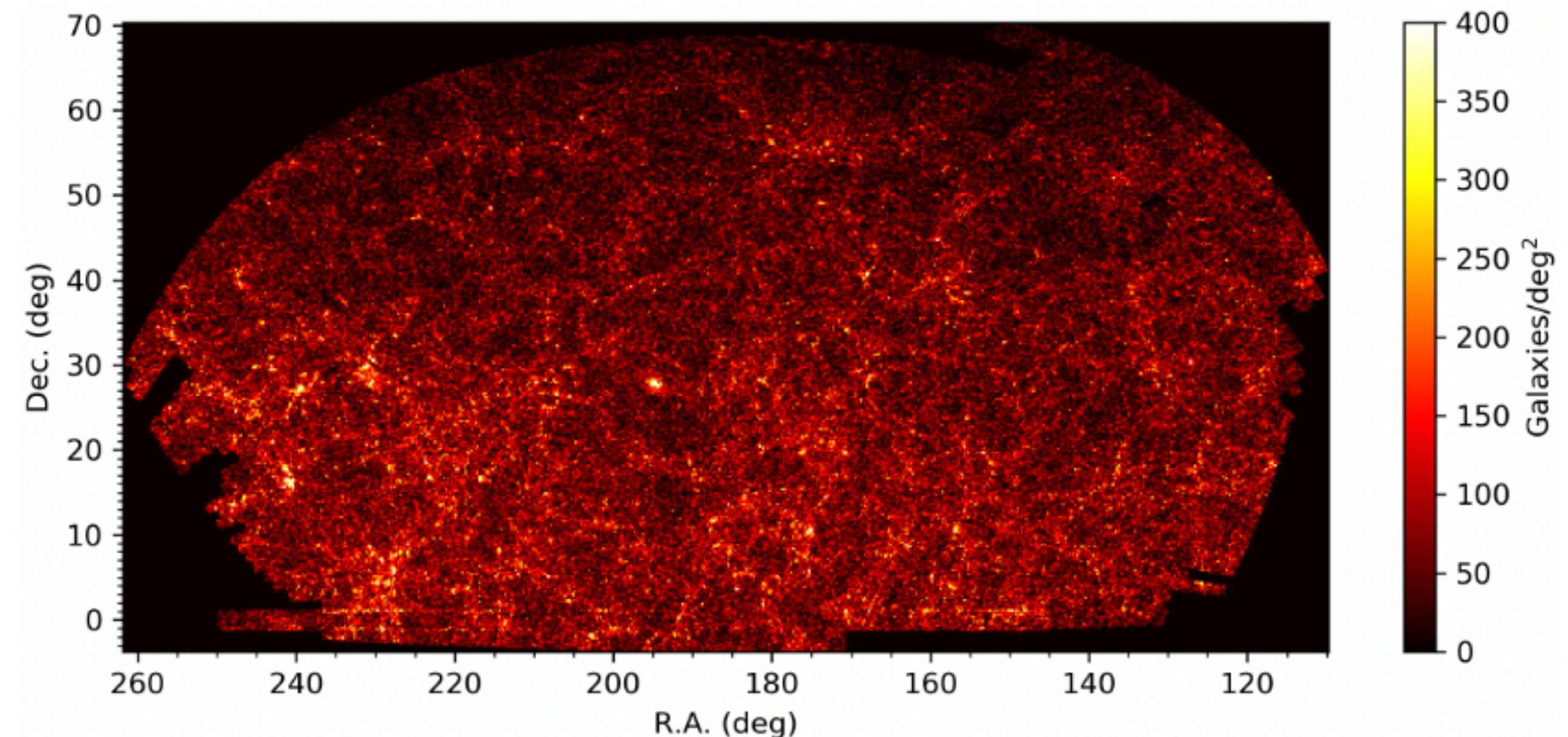
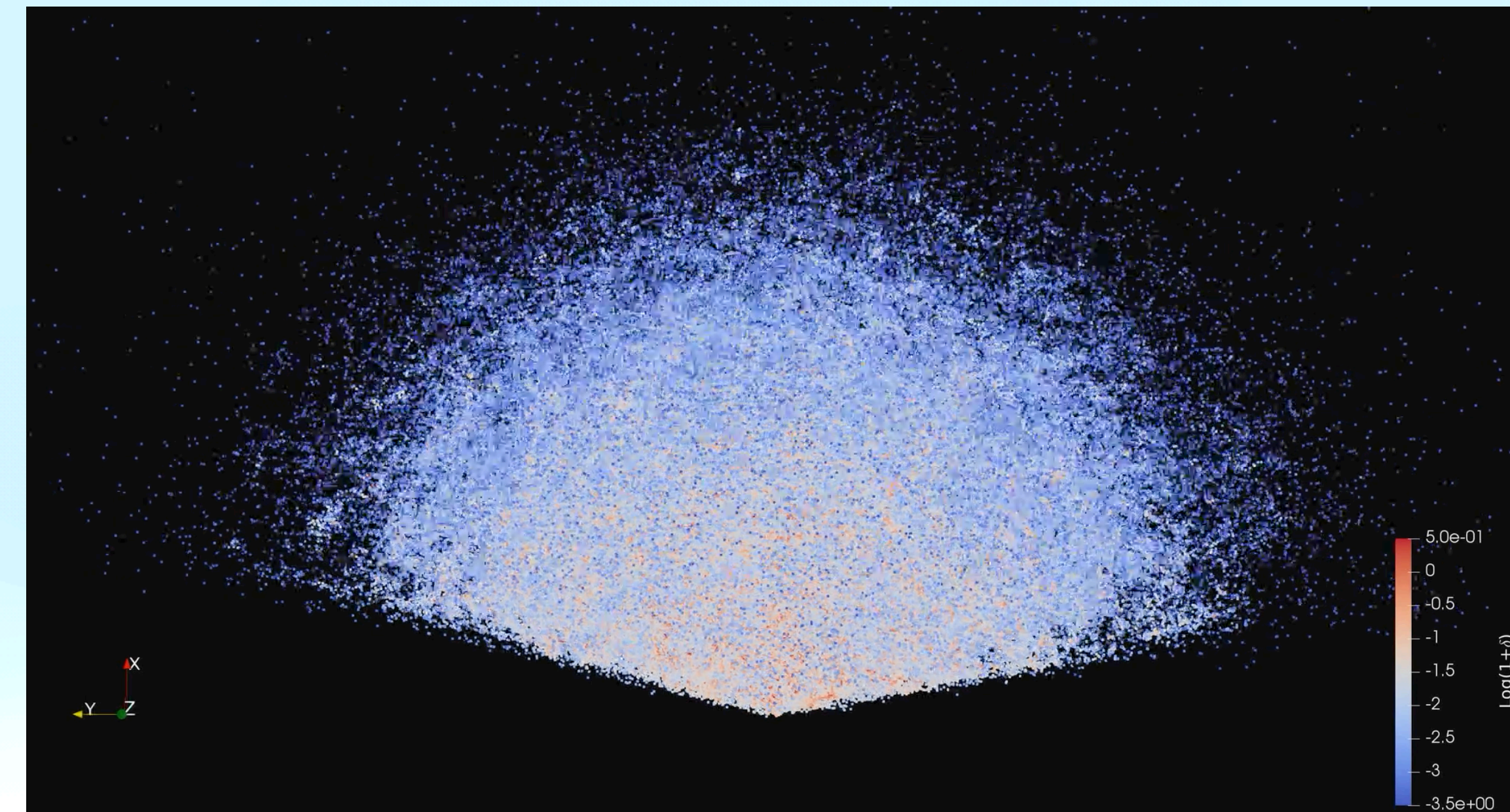
SDSS DR7 Legacy survey Main Galaxy Sample
(Strauss+02) ~600 000 galaxies at $z = 0-0.3$.

Uniform coverage in the plane of the sky and
smooth redshift distribution.



Volume density
distribution: $n(z)$

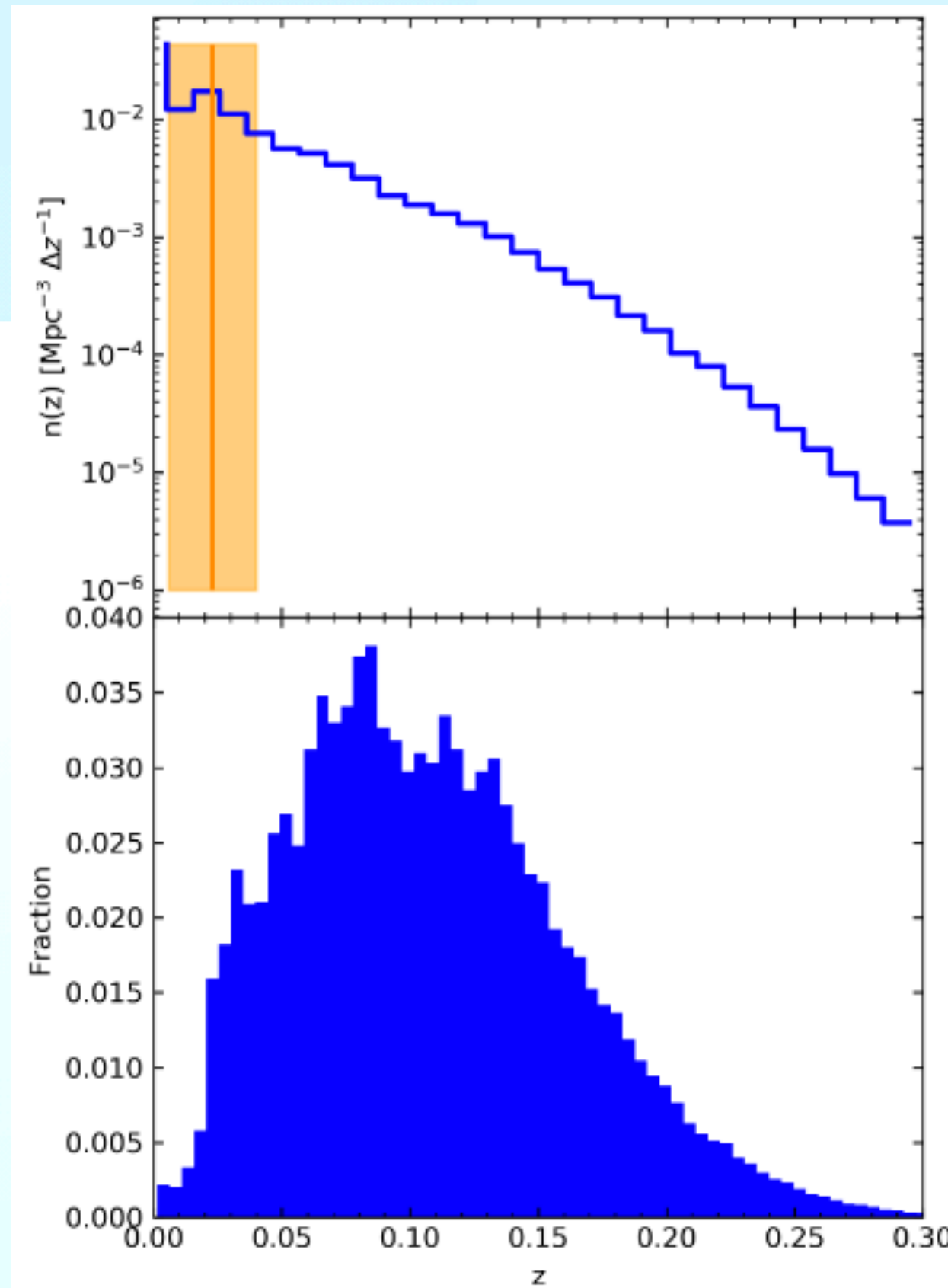
Redshift
distribution



Tracers of the filaments: SDSS

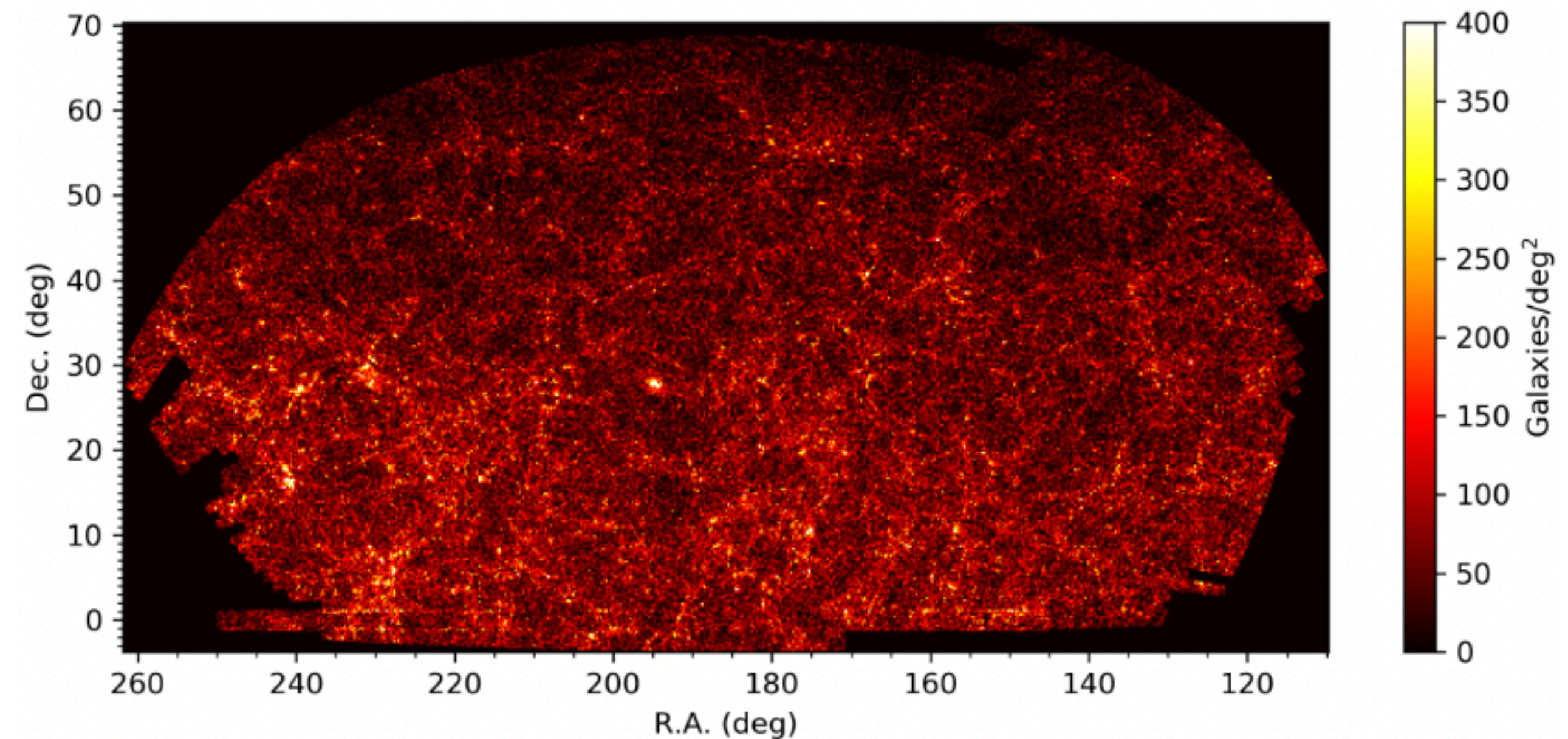
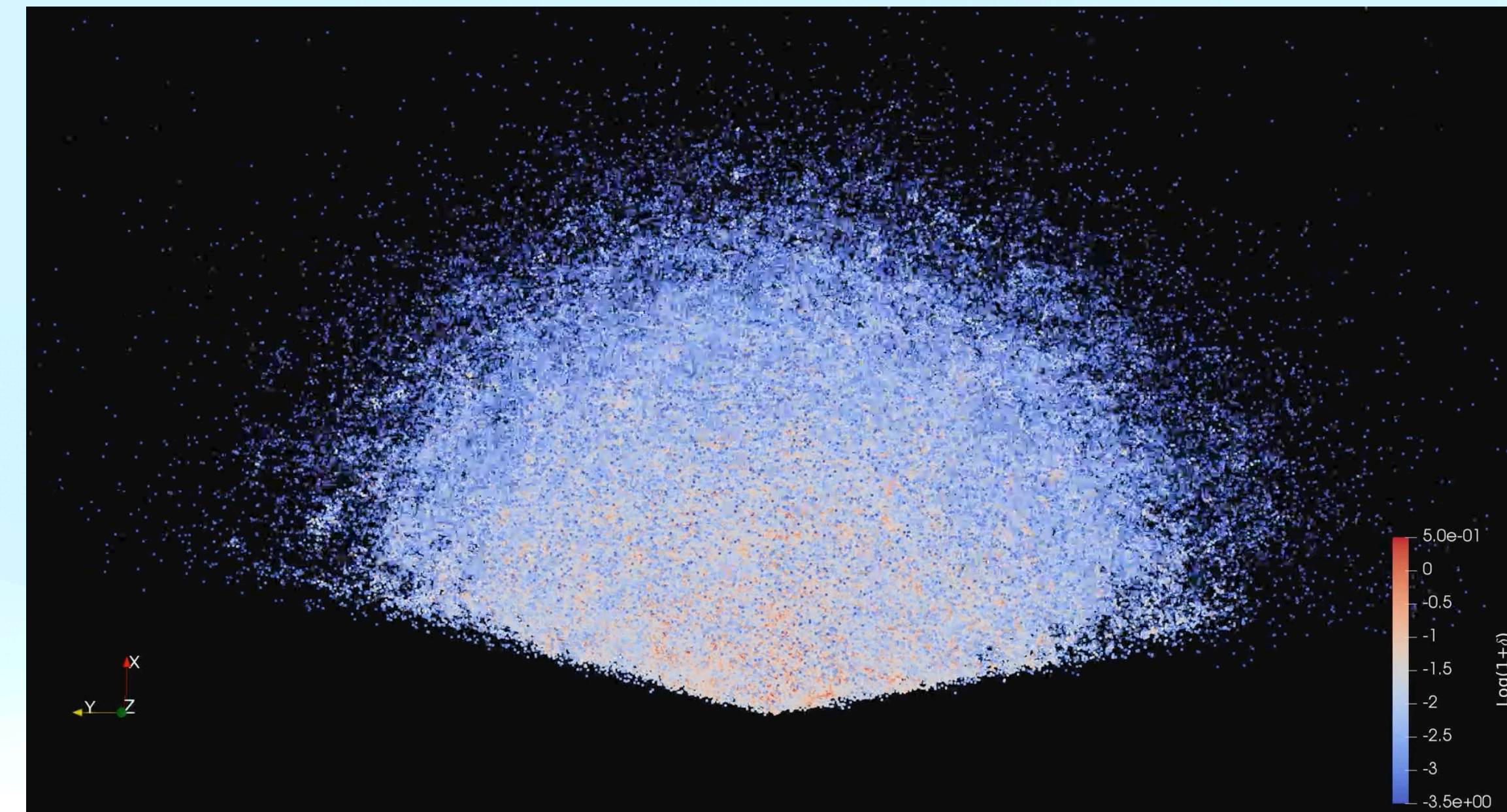
SDSS DR7 Legacy survey Main Galaxy Sample
(Strauss+02) ~600 000 galaxies at $z = 0-0.3$.

Uniform coverage in the plane of the sky and
smooth redshift distribution.



Volume density
distribution: $n(z)$

Redshift
distribution

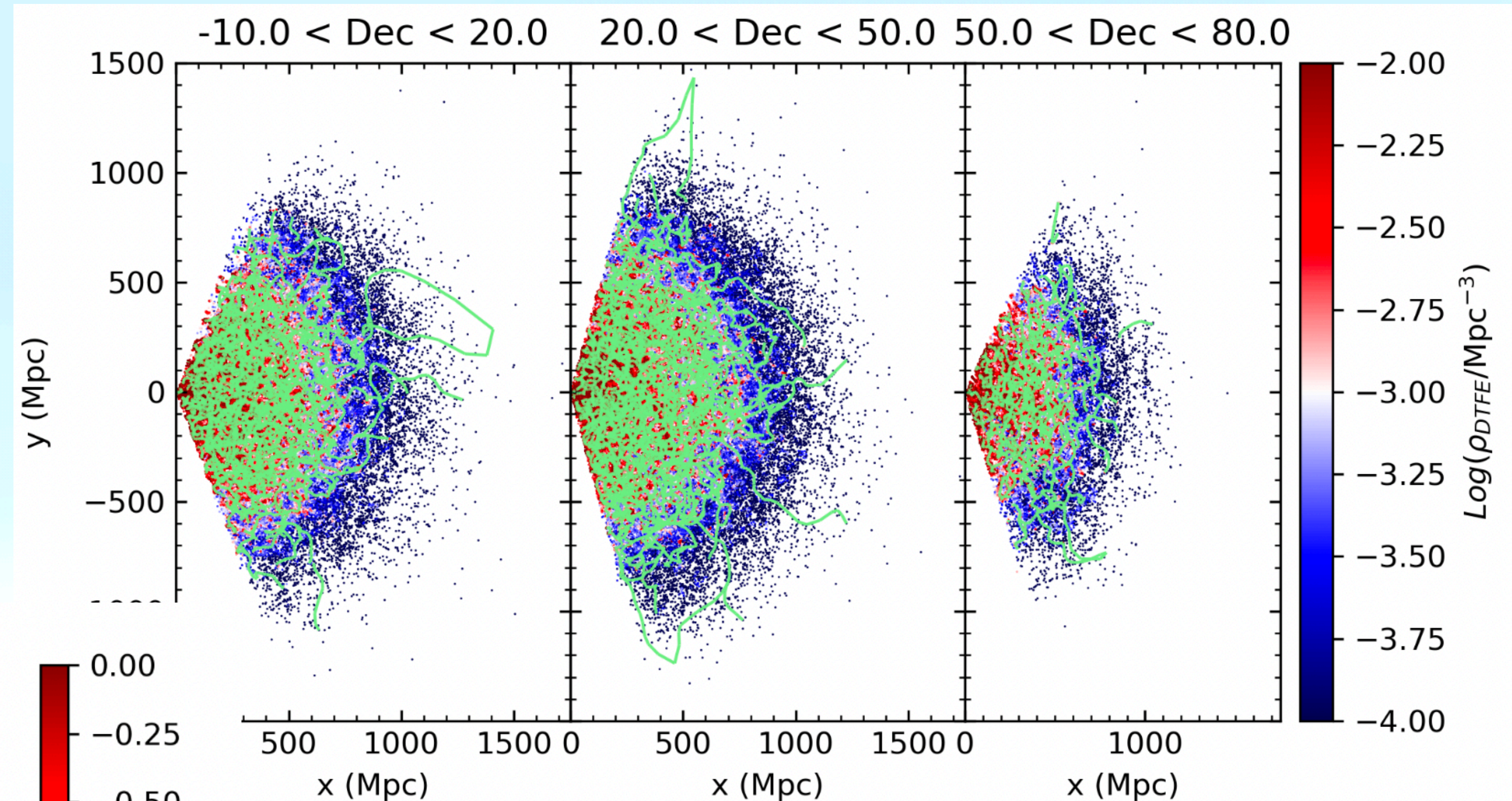
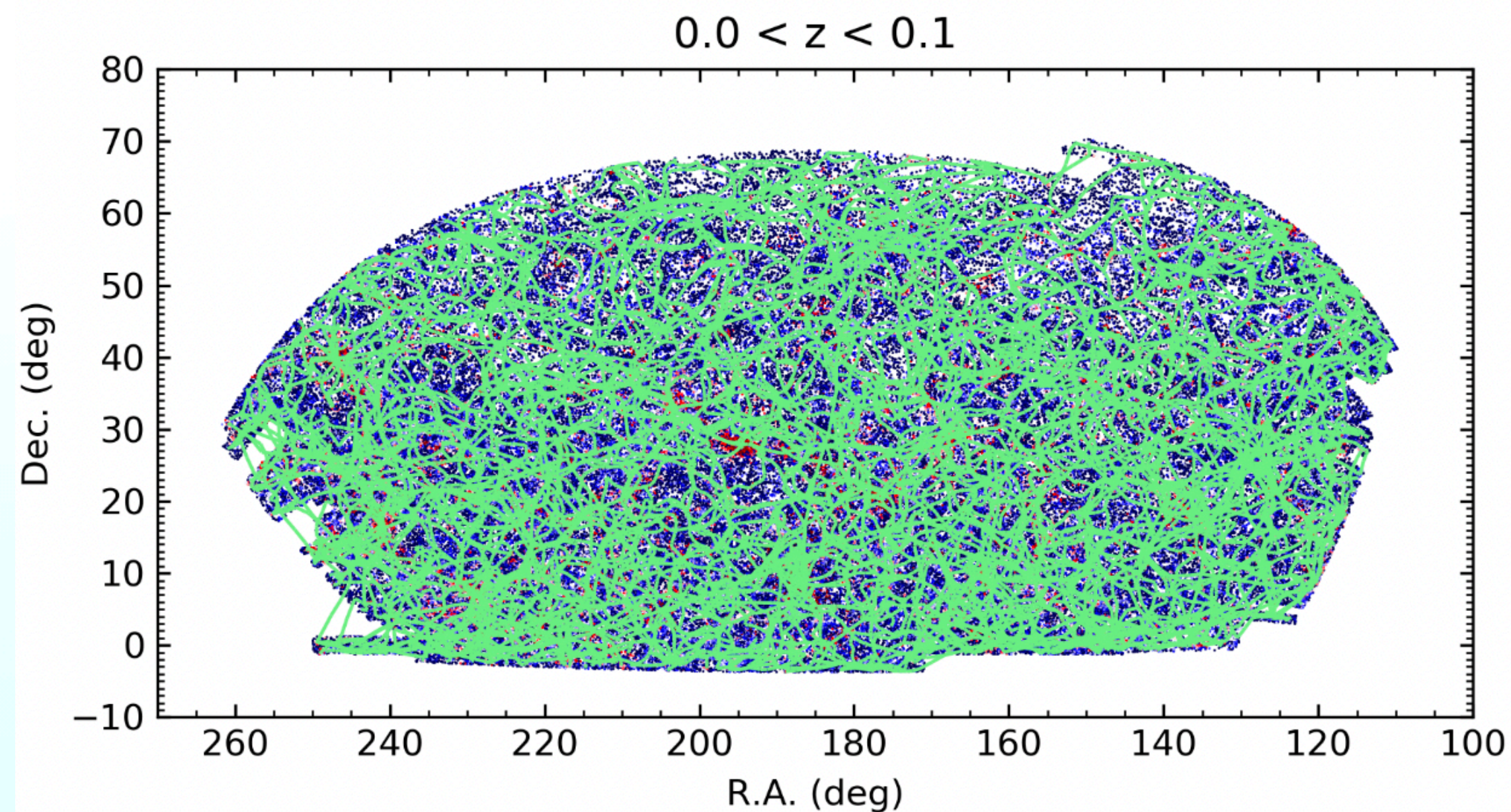


Catalogue of filaments in the SDSS

Malavasi et al. 2020b

We released the catalogue of filaments for use by the community.

Full characterization of properties and systematics.

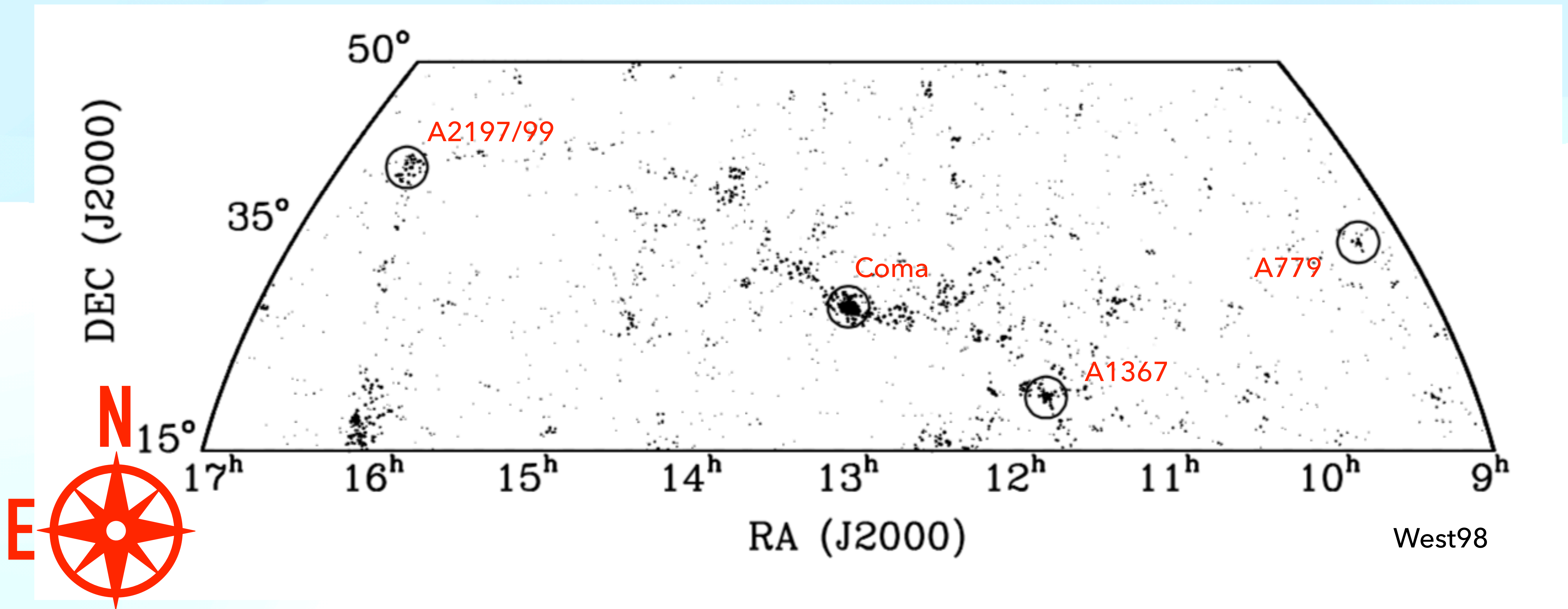


Available for DR7 and DR12.

<https://l3s.osups.universite-paris-saclay.fr/cosfil.html>

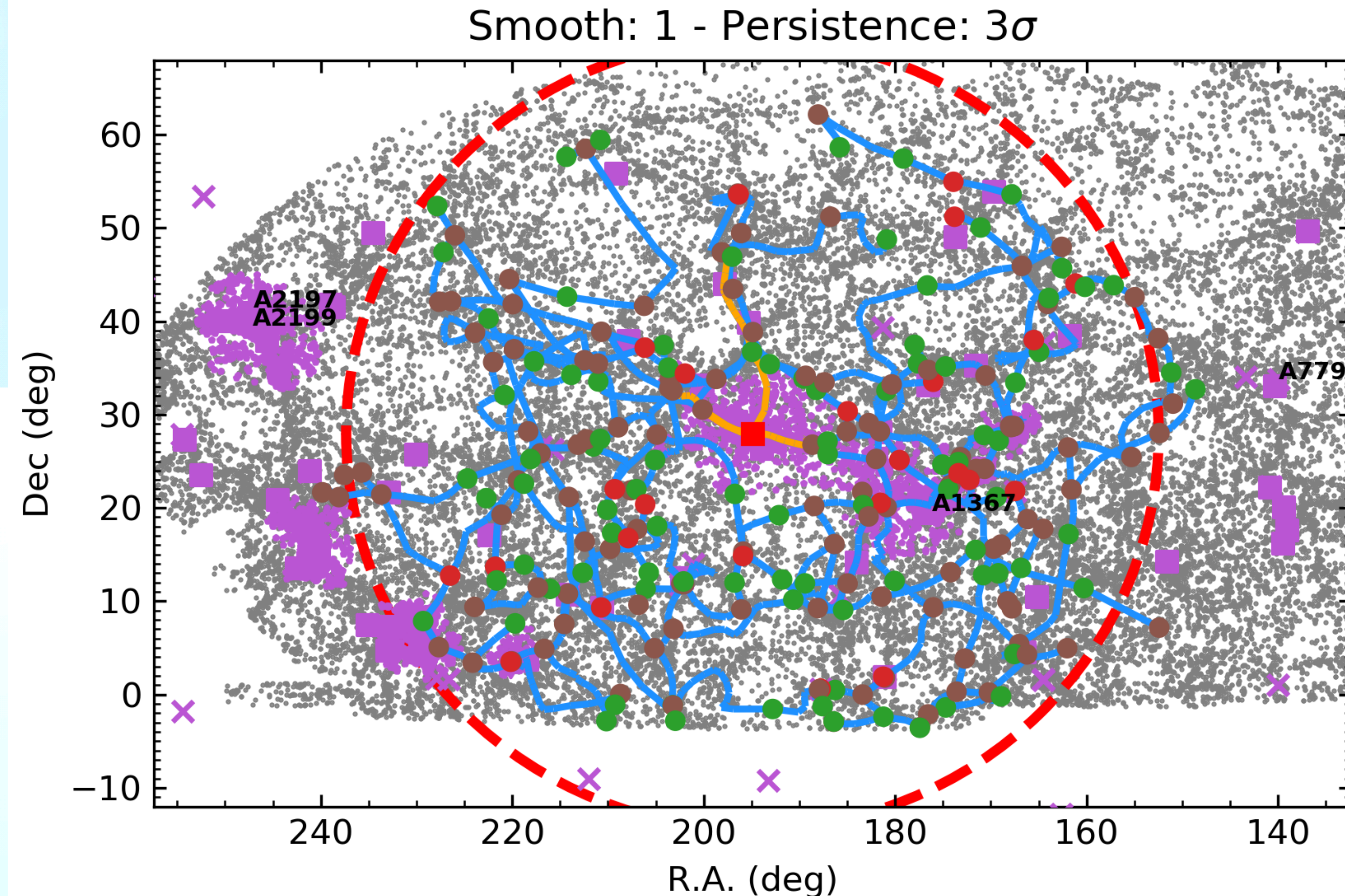
Coma and its LSS

- Physical parameters (Mass, radius, velocity dispersion, Łokas&Mamon03)
- Substructure (Subgroups, ICL) (Adami+05a, 05b, 09)
- Idea of the LSS in the region (other clusters in the region)



Filaments around the Coma cluster

DisPerSE: Sousbie11, Sousbie+11 - SDSS DR7: Abazajian+09, Strauss+02

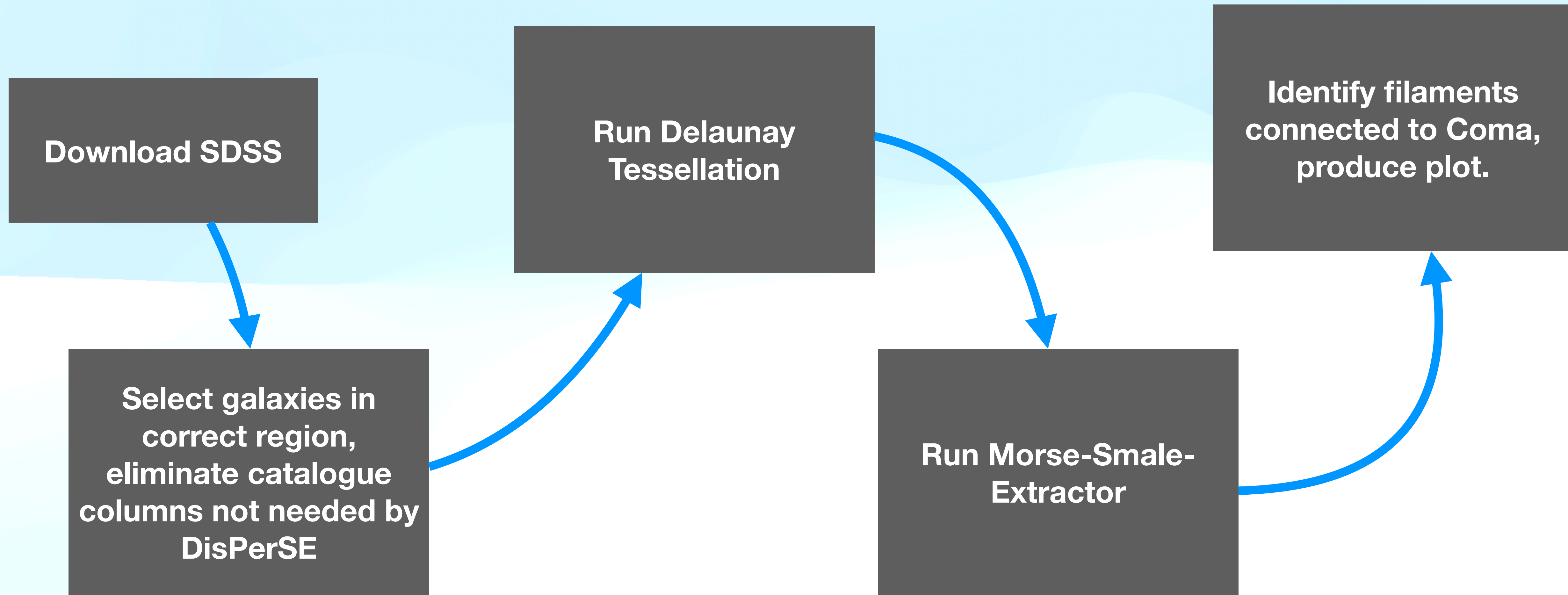


- Discrete Persistent Structure Extractor (DisPerSE) applied to Sloan Digital Sky Survey DR7.
- Focus on the Coma cluster: well known cluster with a lot of information available.
- Detected three filaments connected to the Coma cluster.

Implementing the analysis in SciTraceWeb

- Breakdown the analysis in sequential steps.
- Identify input and output.
- Identify parameters.
- Write script to install the code and its dependencies.
- Optimize and tidy up the code: your package will be explored by others.
- Create repository with the correct structure.
- Login to SciTraceWeb app and implement.

Identifying steps in the analysis



Identifying steps in the analysis

1

Download SDSS

Select galaxies in
correct region,
eliminate catalogue
columns not needed by
DisPerSE

2

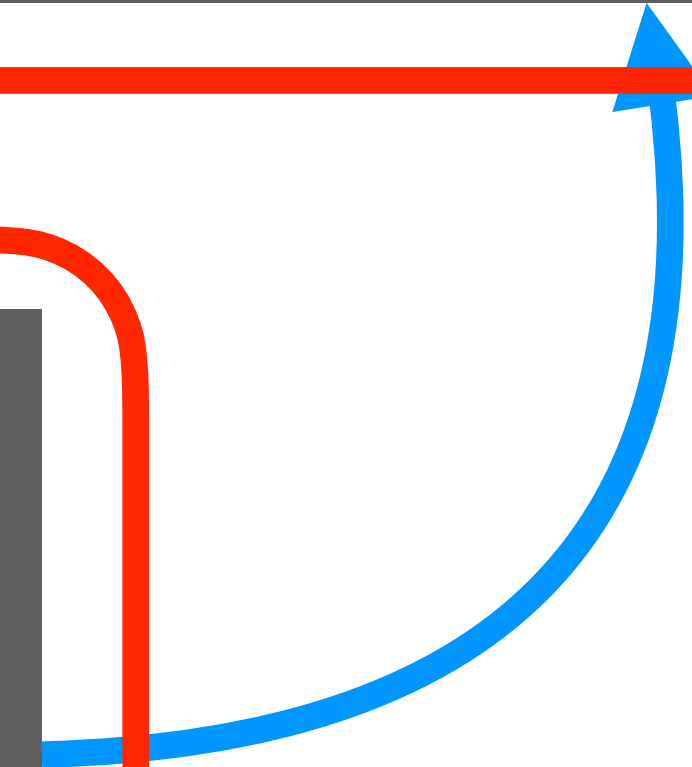
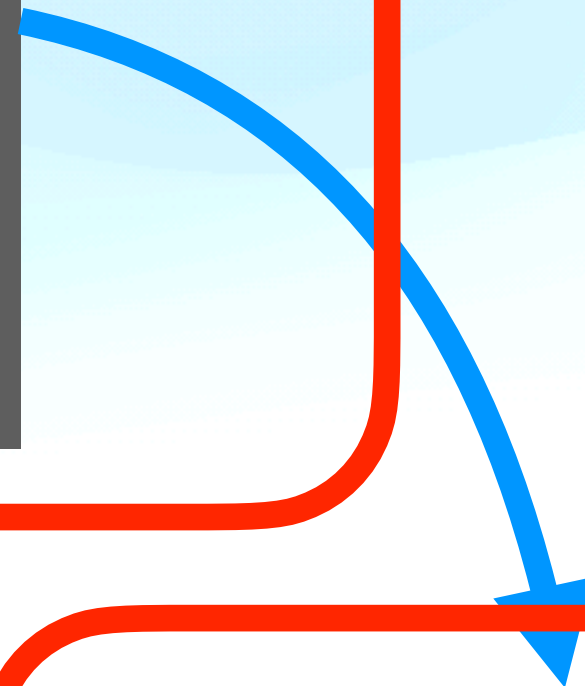
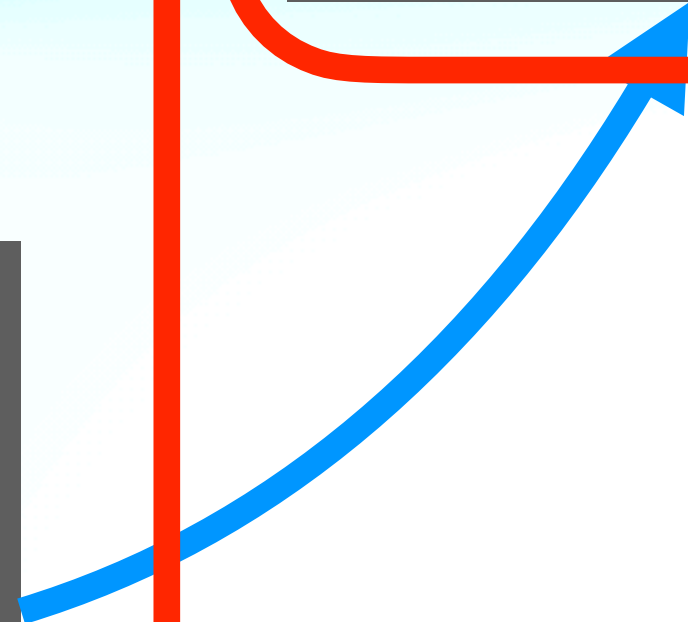
Run Delaunay
Tessellation

3

Run Morse-Smale-
Extractor

4

Identify filaments
connected to Coma,
produce plot.



Input & output

- Step 1: no input, output is a catalogue of galaxies ready for DisPerSE
- Step 2: input is catalogue output by step 1, output is tessellation
- Step 3: input is tessellation, output is skeleton
- Step 4: input is skeleton AND catalogue output by step 1, output is plot (PNG)

Parameters

- Step 1: file names, whether to output intermediate catalogues, coordinate centers and cosmology for coordinate conversion
- Step 2: DisPerSE parameter, number of smoothing cycles
- Step 3: DisPerSE parameter, persistence threshold
- Step4: radius up to which search for filaments connected to Coma