

# Biomechanics group

Status update, Zurich Jan 18<sup>th</sup> 2024

# Aims

- Develop a wrapper to export properties in **HDF5 files** towards a FE solver (ParOSol) (data robustness, accessibility)
- Develop the **automatic segmentation** pipeline for human femurs from clinical CT images (accessibility, multimodality)
- Widening the modelling strategies from **microCT images** of trabecular bone samples (data versatility)

# HDF5



- Why HDF5?
  - Hierarchical Data Format designed to store and organize large amounts of data
  - Open file format
  - Multi-datatype support
  - Architecture independent (data storage)
- Integrates with the ciclope pipeline
  - Originally using Calculix, now “migrating” towards ParOSol for increased scalability

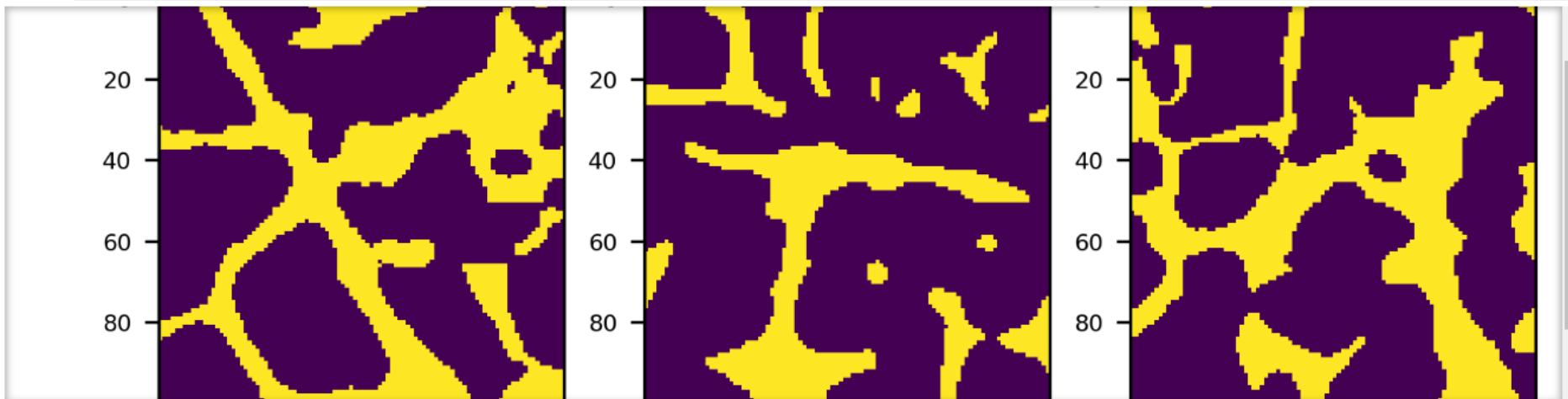
# HDF5

```
def vol2h5ParOSol(voldata, fileout, topDisplacement, voxelsize=1, poisson_ratio=0.3, young_modulus=18e3, topHorizontalFixedDisplacement=True, verbose=False):
    """Generate ParOSol HDF5 (.h5) input file from 3D volume data.
    Before to generate ParOSol HDF5 file, the Bounding BOX (bbox class) limits the input binary image.
    Info on HDF5 file type for ParOSol solver at: https://github.com/reox/parosol-tu-wien/blob/master/doc/file\_format.md

Parameters
-----
voldata : ndarray
    3D voxel data.
fileout : str
    Output .h5 file.
topDisplacement: float
    Vertical displacement imposed at the top.
voxelsize : float
    3D model voxelsize.
poisson_ratio: float
    Poisson ratio.
Young_modulus: float
    Young's modulus [MPa].
topHorizontalFixedDisplacement: bool
    if True X and Y displacements fixed at the top; if False no displacements fixed at the top.
verbose : bool
    Activate verbose output.
"""
```

# HDF5

```
[25]: plot_midplanes(L)
plt.show()
```



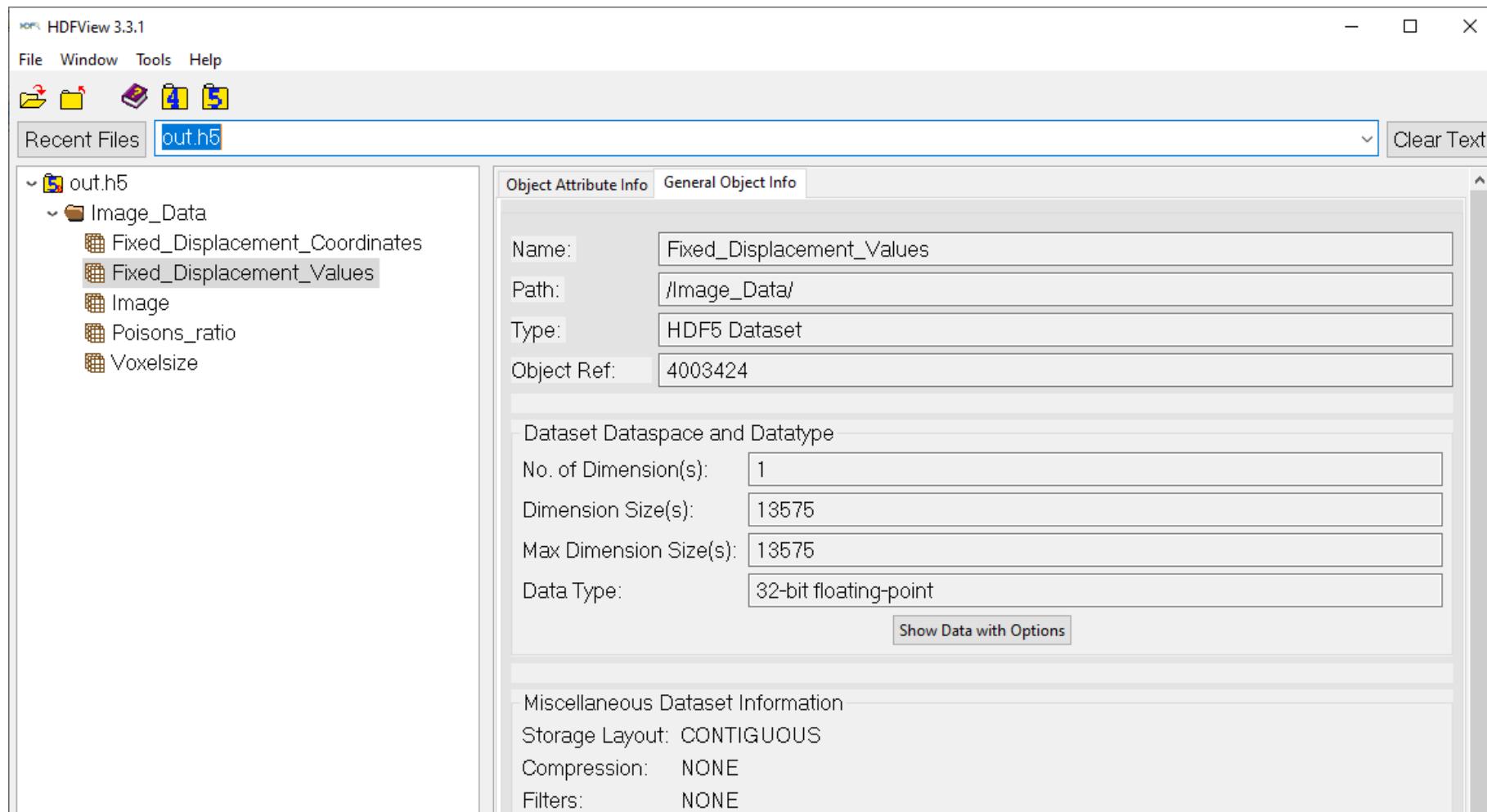
Generate ParOSol hdf5 from volume data

```
[29]: ciclope.core.voxelFE.vol2h5ParOSol(L, 'paroSOLout.h5', -0.4, 1, 0.3, 18e3, True, True )
```



```
INFO:root:Opening Output file
INFO:root:creating h5 Image_Data Group
INFO:root:creating Image dataset...
INFO:root:Setting voxel size
INFO:root:Setting Poisson ratio
INFO:root:Creating Fixed_Displacement_Coordinates
INFO:root:Creating Fixed_Displacement_Values
INFO:root:hdf5 export done!
```

# HDF5



# HDF5

- Next steps:
  - Solve the FE problem with ParOSol
  - Read and visualize the hdf5 output from ParOSol

# Automatic segmentation



- Not much code
- No organization
- No logo

# Automatic segmentation



A photograph of a human femur bone that has been artistically painted with a vibrant, psychedelic pattern. The design includes a large peace sign on the upper right side, various flowers, and intricate line work. The bone is set against a plain white background.

Hippyfemur

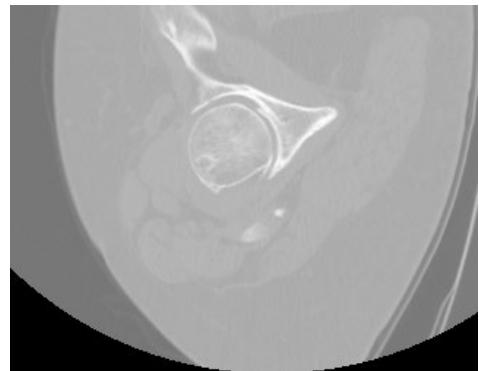
Type  to search | >

Overview    Repositories 3    Projects    Packages    Teams    People 4    Settings

Hippyfemur

The image shows a screenshot of a GitHub-like interface for a repository named "Hippyfemur". The repository page features the same colorful, abstractly painted femur bone as the background image. The GitHub navigation bar is visible at the top, with "Overview" being the active tab. Below the header, there's a search bar and several navigation links: "Repositories" (3), "Projects", "Packages", "Teams", "People" (4), and "Settings". The main content area displays the repository name "Hippyfemur" next to its icon.

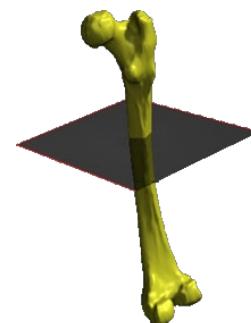
# Automatic segmentation



CT images  
Metadata  
Patient information

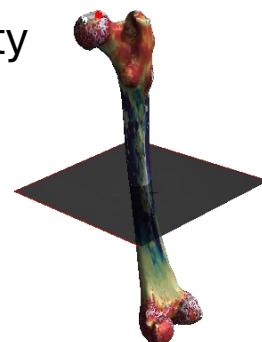
Automatic  
segmentation DL

Binarized volume  
Triangulated surface



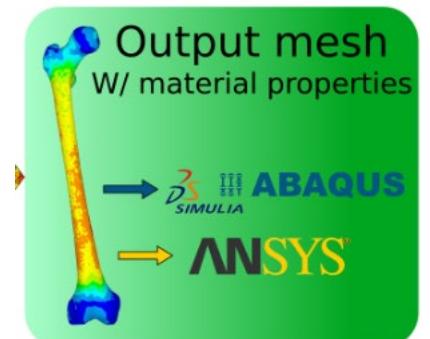
Refined  
segmentation with  
sub-pixel accuracy

Binarized volume  
Triangulated surface  
Cortical thickness  
and density



Automatic meshing

FE mesh  
Material properties  
Boundary conditions



# Automatic segmentation

**AutomaticFemurSegmentation** Private

Repository for the python code of automatic femur segmentation, inspired by Väänänen & Grassi et al., 2019  
<https://doi.org/10.1016/j.medengphy.2019.06.015>

● Python 0 ⚙ 0 ⭐ 0 ⏺ 0 Updated 32 minutes ago

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**AutomaticFemurSegmentationDL** Public

● Python 0 ⚙ 0 ⭐ 0 ⏺ 0 Updated 32 minutes ago

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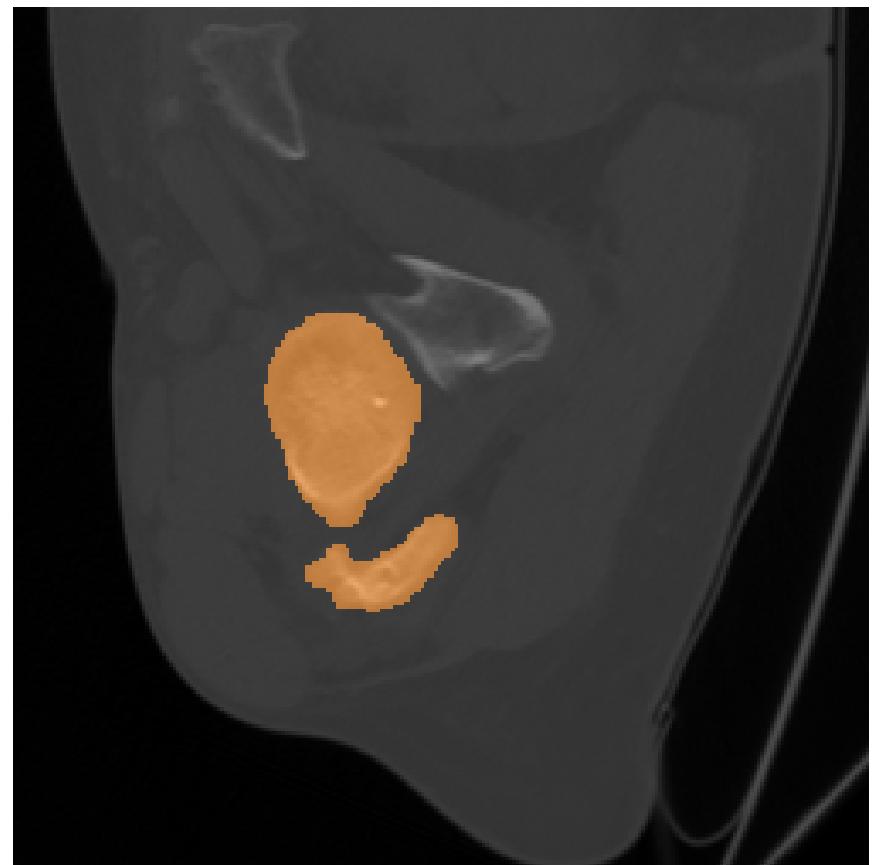
**AutomaticFemurSegmentationMATLAB** Private

MATLAB code from the automatic femur segmentation described in Väänänen&Grassi et al., MEP 2019

● MATLAB 0 ⚙ 0 ⭐ 0 ⏺ 0 Updated 3 hours ago

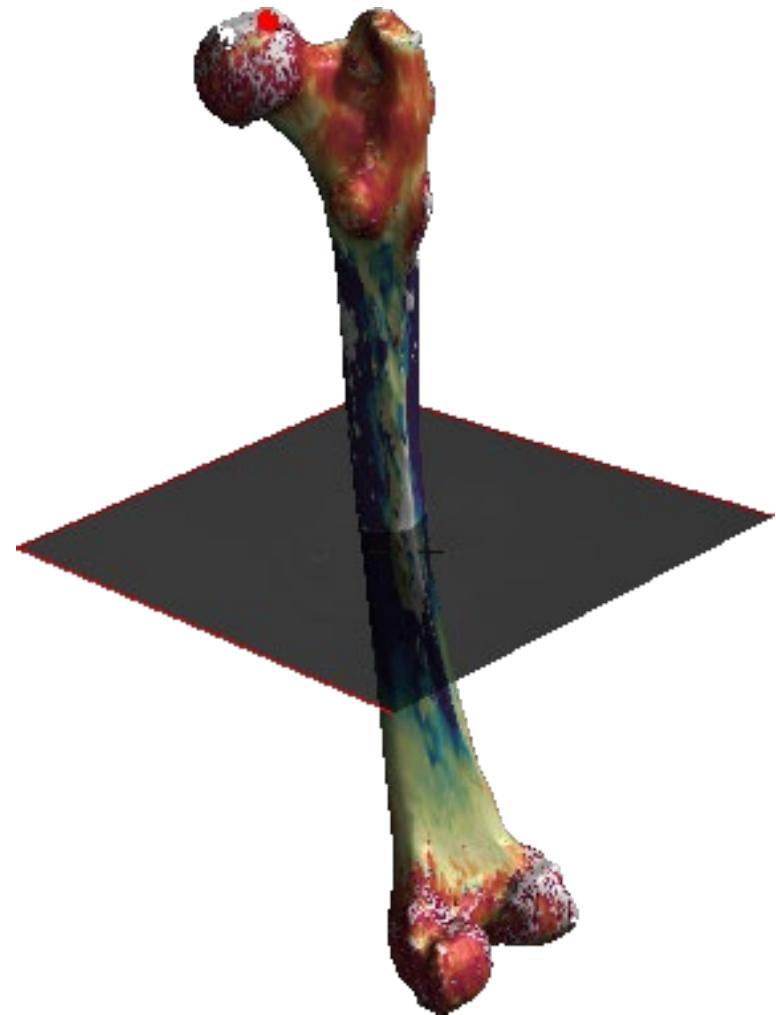
# Automatic segmentation DL

- Trained on openly available repository
  - Zenodo link
- Works on each individual 2D slice
- Intersection over union = 0.95
- Public repository
  - Needs increased testing
  - Documentation



# Automatic segmentation sub-pixel resolution

- Main idea:
  - run optimization using Stradview (freeware) using 3 sets of parameters
  - Combine output from the 3 runs
- Struggling with changes in Stradview compared to 2017
  - The 3 optimizations run, need to combine the outputs



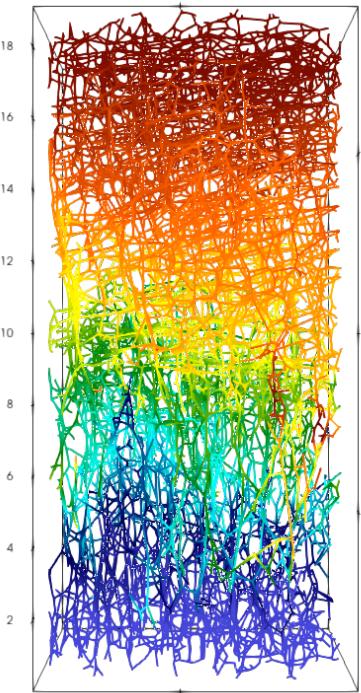
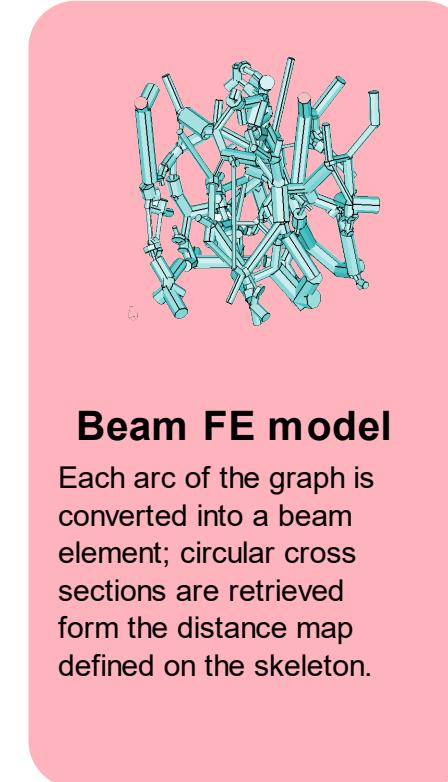
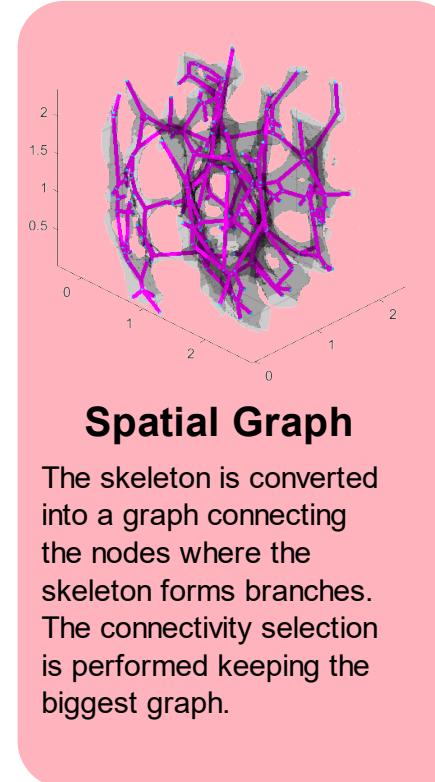
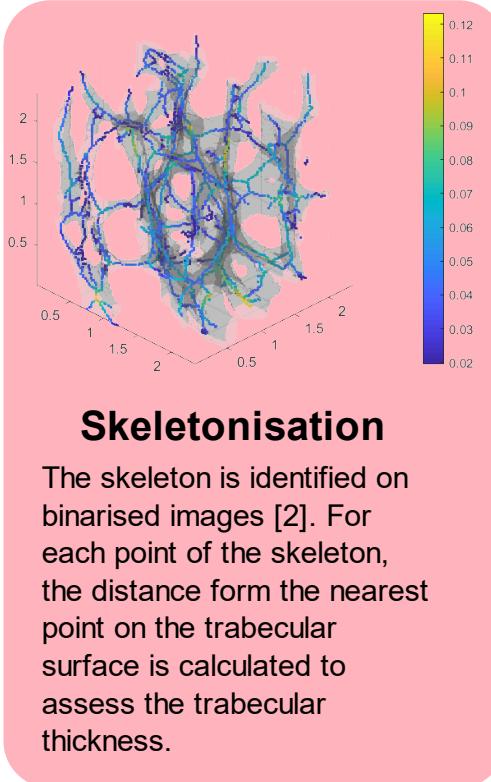
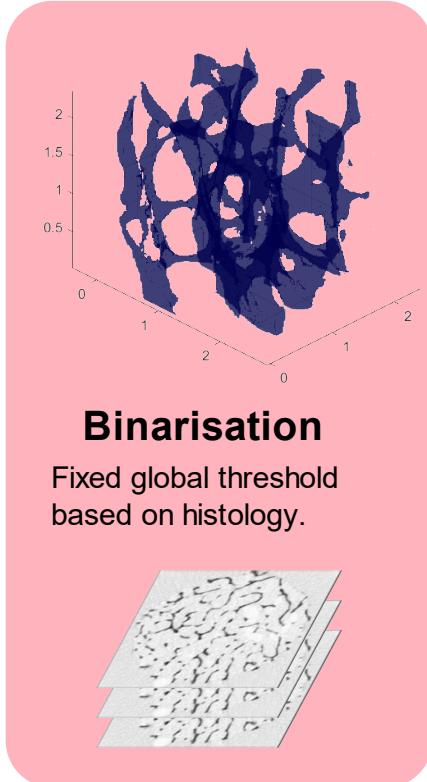
# Beam FE modelling



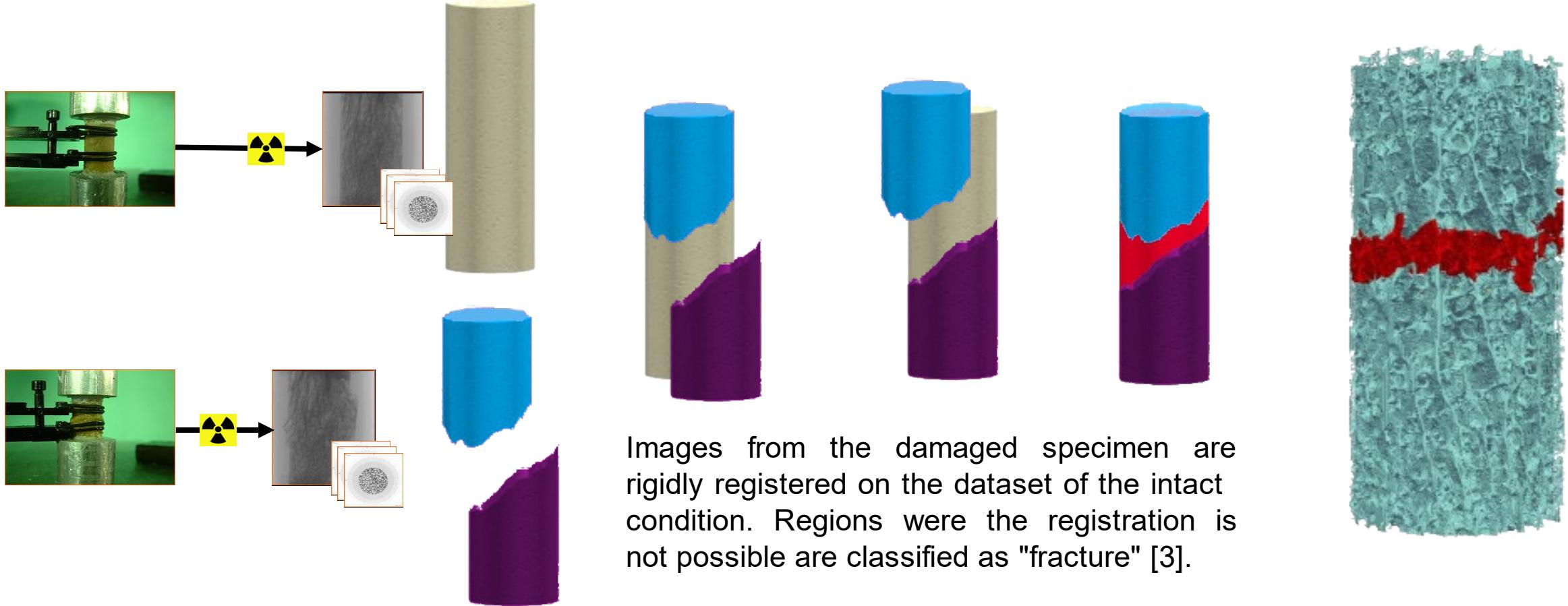
- [https://github.com/gianthk/ciclope/blob/master/examples/jupyter/ciclope\\_ex05.1\\_beamFE-skel.ipynb](https://github.com/gianthk/ciclope/blob/master/examples/jupyter/ciclope_ex05.1_beamFE-skel.ipynb)

# Beam FE model of the trabecular structure

- Each trabecula can be modelled as a beam element (circular cross = thickness of each trabecula).
- Isotropic and homogeneous material with linear elastic constitutive law ( $E = 19 \text{ GPa}$ ,  $\nu = 0.3$ ).
- Kinematic Boundary Conditions to replicate the uni-axial compression test (nodes at the bottom layer fully constrained, nodes at the top layer imposed a vertical displacement and null transversal displacement).

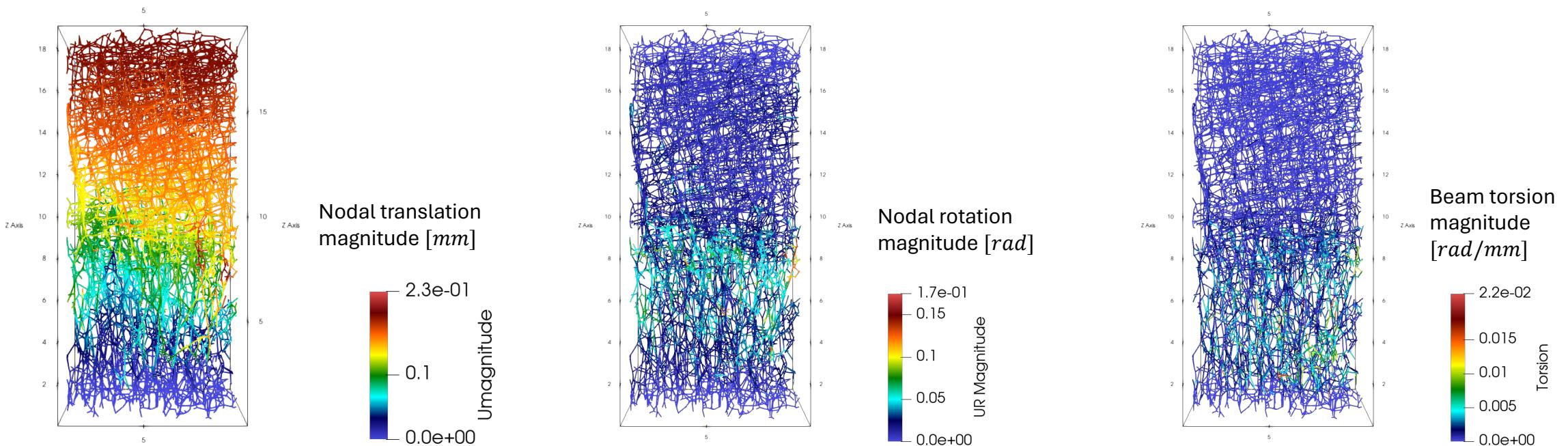


# Fracture identification (as per the previous study)

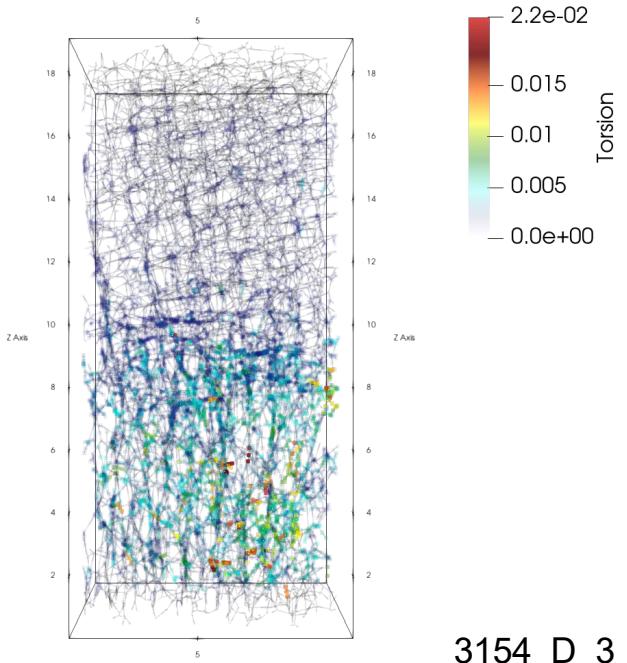


# Output/Comparison

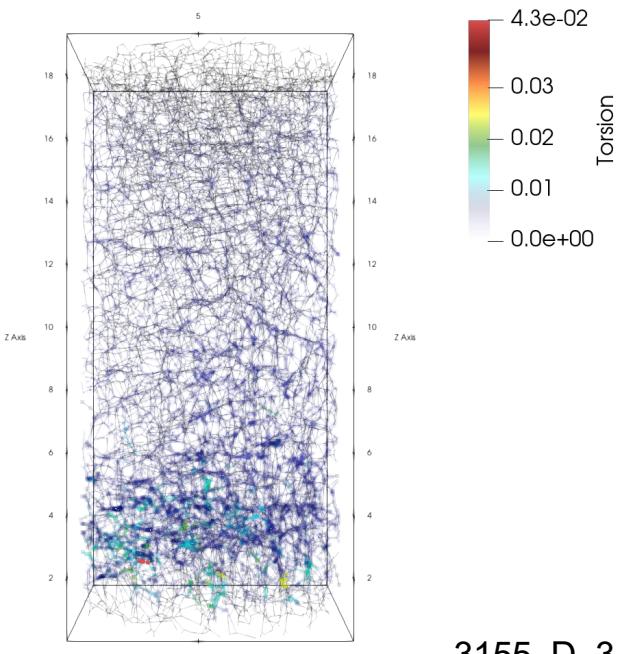
- For arch beam element, the differential axial rotation of its nodes was divided by the length of the element, defining the specific torsion [rad/mm].
- The map of the element specific torsion presented peaks matching the map of the fractured volumes.
- The difference in the axial positioning between torsion peaks and fractured volumes ranged between 0.67% and 2.46% of the total length of the specimen (20 mm).



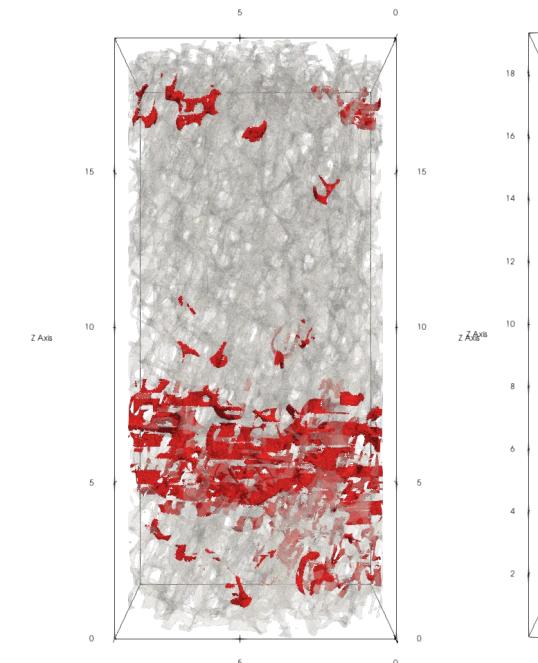
# Results



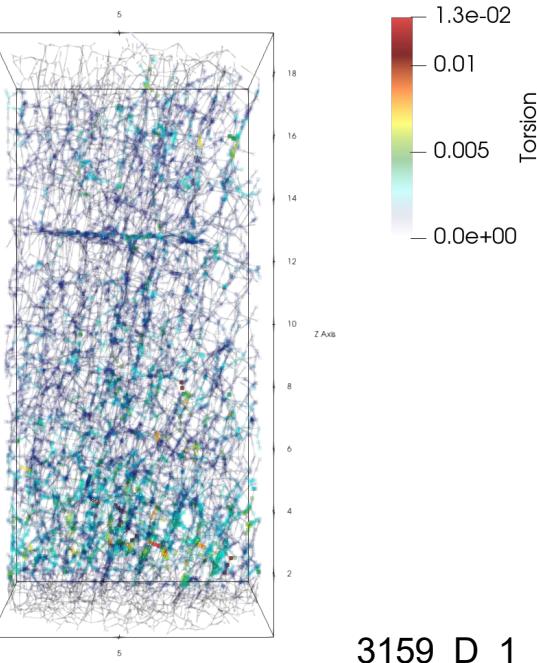
3154\_D\_3



3155\_D\_3

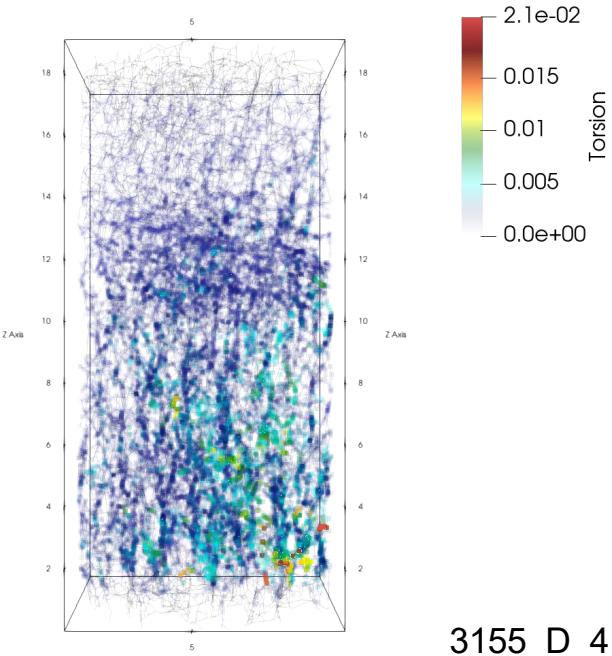


Tp\_2923\_P\_1

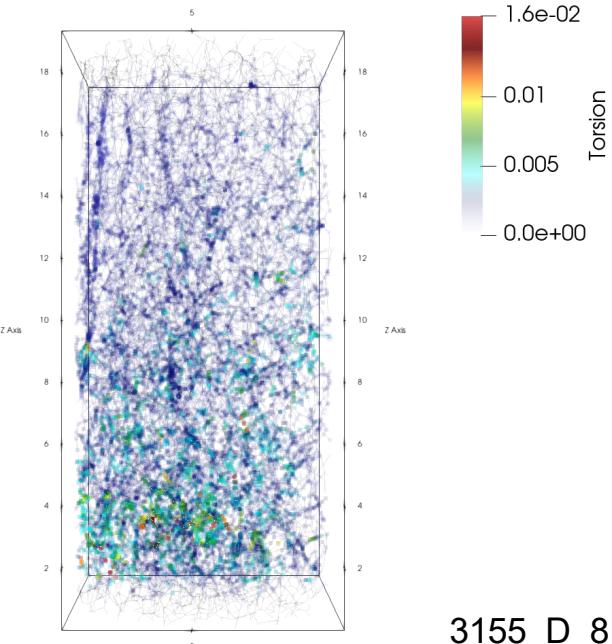


Torsion

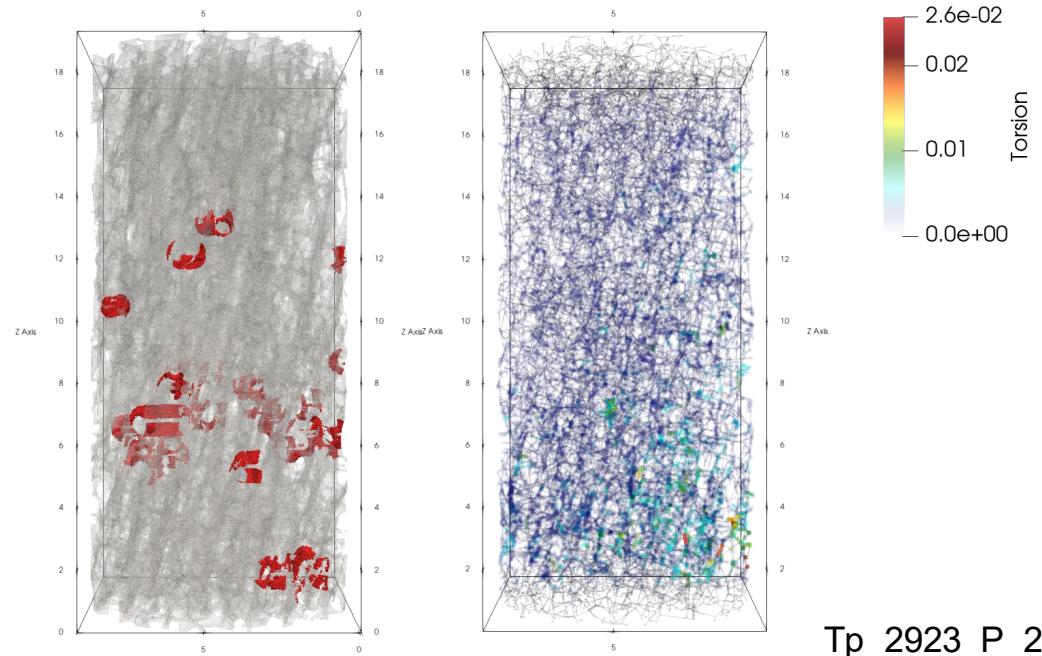
# Results



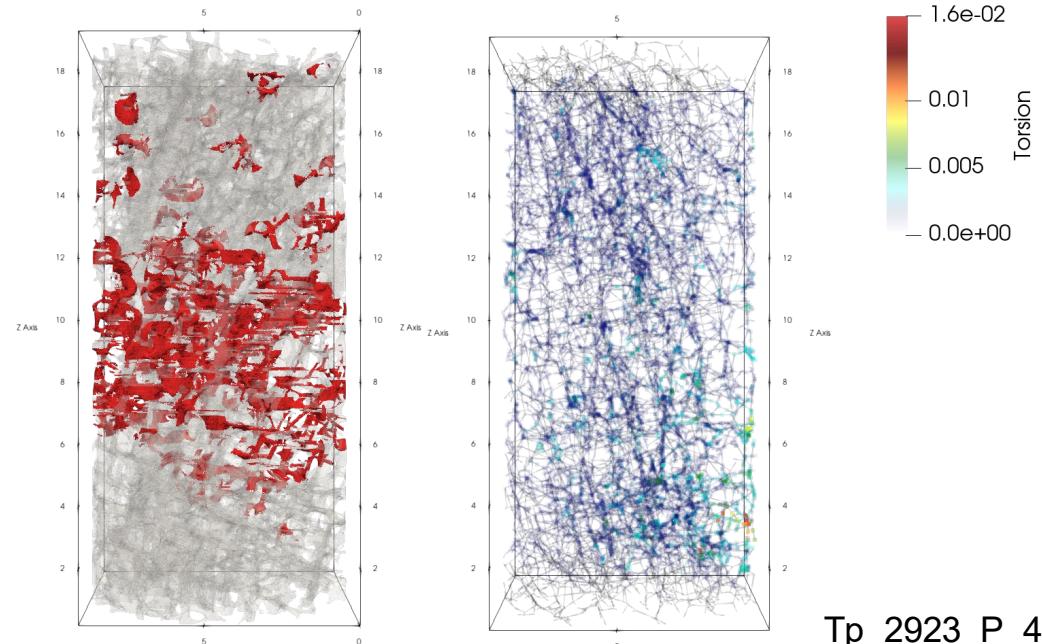
3155\_D\_4



3155\_D\_8



Tp\_2923\_P\_2



Tp\_2923\_P\_4