



# Shape analysis via skeletal models

## User Tutorial

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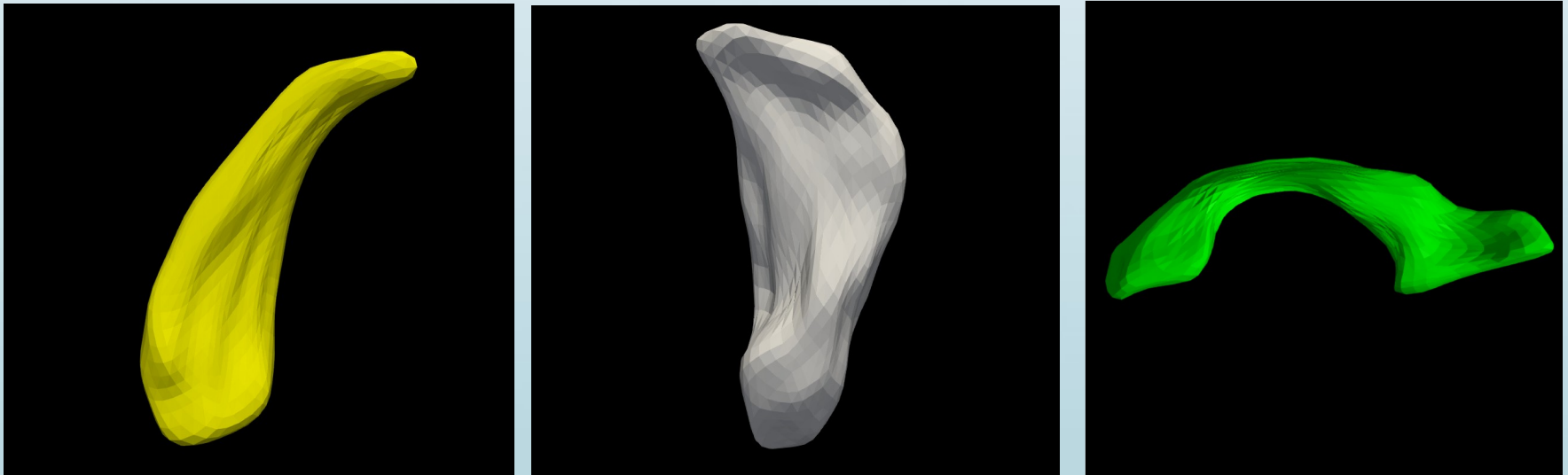
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# Skeletal representations

Shape Analysis allows studying morphology in populations of anatomic structures. A **skeletal representation (s-rep)** is used to model the structure of a non-branching object, providing a rich geometric representation with correspondence across cases.

The S-rep Extension (short for **Skeletal Representation Extension**) in SlicerSALT provides utilities to visualize, establish and refine s-reps of 3-dimensional objects.



# The Slabular S-rep

A 3D object whose length is notably larger than its width which is notably larger than its thickness is suitable for modeling by a slabular s-rep

A slabular s-rep (from now on just referred to as an s-rep) consists first of a folded, two-sided quasi-medial sheet inside the object. This sheet is called the “skeletal sheet”. It is sampled into a network of skeletal positions

Special skeletal samples are at positions where the sheet folds back onto itself

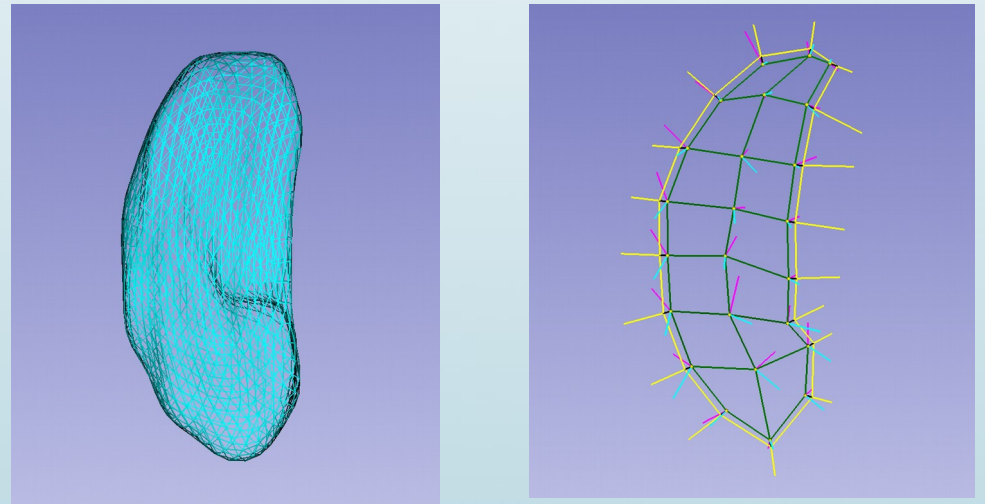


Figure: A skeletal model for a hippocampus. Left: the surface mesh of hippocampus. Right: the skeletal sheet of the s-rep of this hippocampus

# The Slabular S-rep (2)

From each sample point on the skeleton, a vector to and approximately orthogonal to the boundary is provided as part of the s-rep. These vectors are called “spokes”.

The spokes emanating from the fold curve of the sheet meet the object surface at crest points.

The spokes can be interpolated into a finer mesh, and the spoke endpoints can be interpolated into a implied boundary for the object.

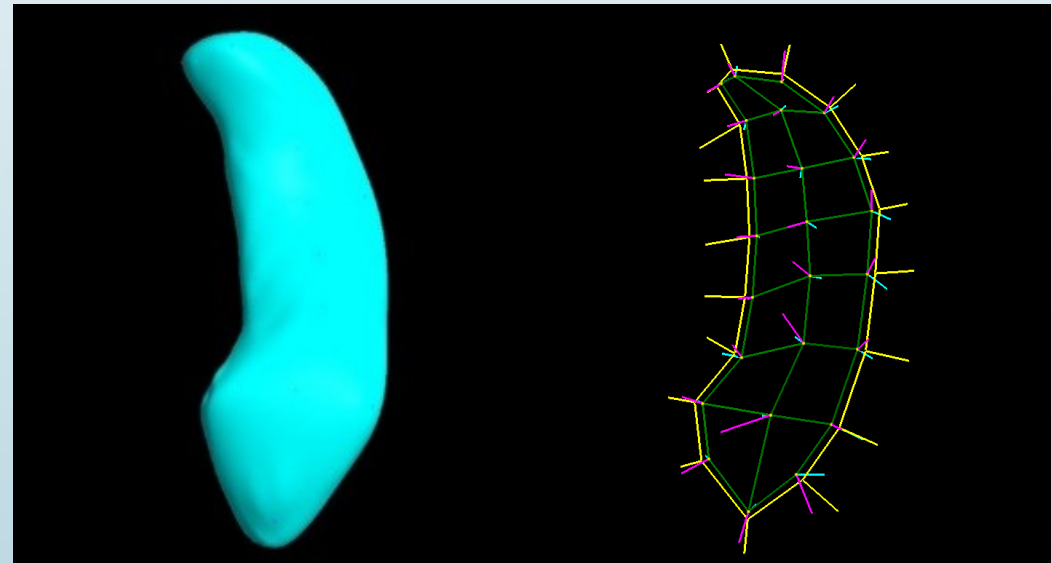


Figure: A skeletal model for a hippocampus.  
Left: implied boundary of hippocampus.  
Right: the s-rep of hippocampus

# Modules in S-rep Extension

Module 1: Establish an s-rep by mean curvature flow of an object boundary

Module 2: Refine an s-rep to fit an object boundary

Module 3: Visualize an s-rep

# Description of S-rep Extension

This extension consists of the following detailed steps:

*Input: Surface Mesh*

Step 1 (a) : Flow surface mesh into quasi-ellipsoidal surface

*Outputs: quasi-ellipsoidal surface & the best fitting ellipsoid*



*Input: Surface Mesh*

Step 1 (b) : Reverse the flow to derive s-rep of the original object

*Output: the s-rep of input object*



*Inputs: Surface Mesh + initial s-rep*

Step 2 : Refine the input s-rep to be better fit

*Output: Optimized s-rep*



*Input: an s-rep*

Step 3 : Visualize the details of the s-rep

**This tutorial focuses on this part of the extension.**

# Inputs of the S-rep Extension

The input describes an object as either a label image, a tile mesh of the object surface, or a signed distance image of distances to the object boundary, where negative distances are interior to the object, positive distances are exterior to the object, and the object boundary is the zero level set.

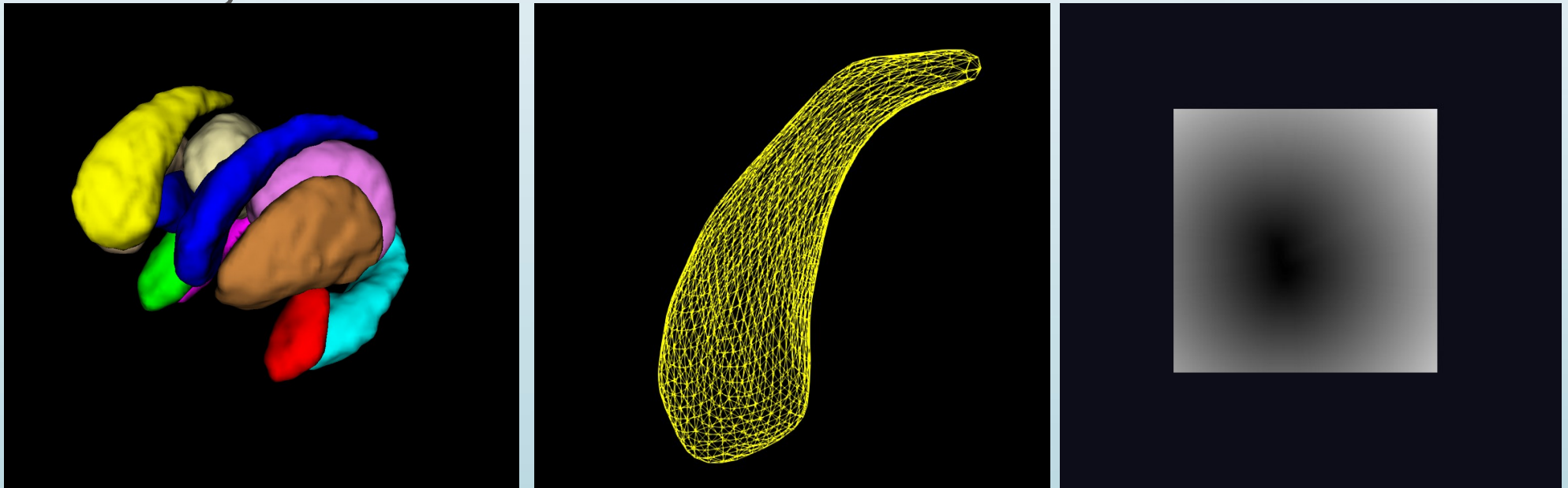


Figure: Different input image types:

Left: 3D model of a label image for multiple objects.

Middle: Surface mesh of a caudate nucleus

Right: one slice of signed distance image



# Output of the S-rep Extension

The output is an s-rep object model that captures both boundary shape and object width.

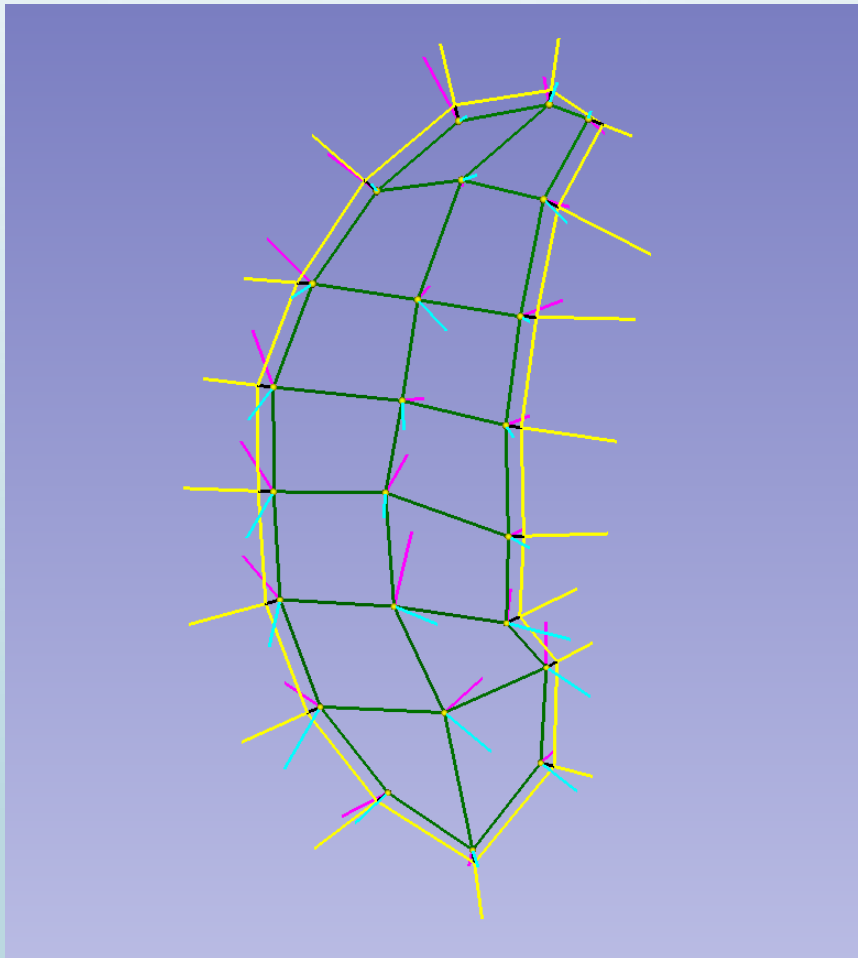


Figure: an example of output



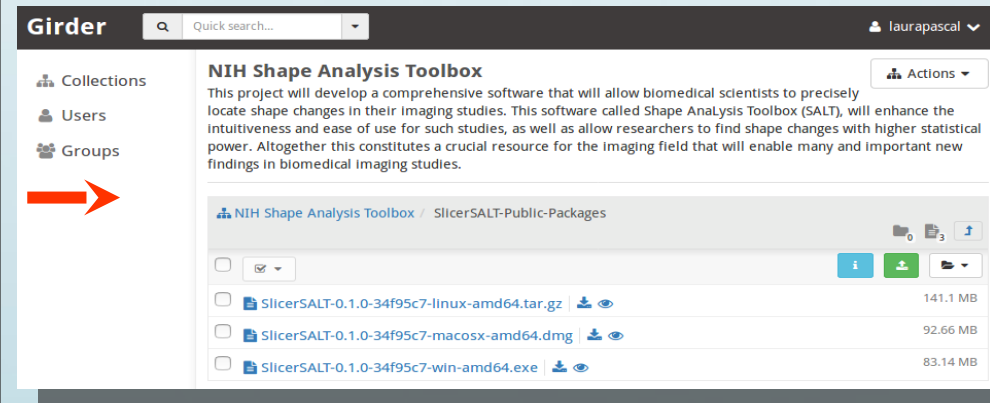
# Installation of the S-rep Extension

The S-rep Extension can be used with two open-source software platforms:

- **SlicerSALT**: which is the dissemination vehicle of powerful shape analysis methodology. This software is a light-weight, customized version of 3D Slicer. It contains the S-rep extension as modules
- **3D Slicer**: which is an open-source and free software platform for medical image informatics, image processing, and three-dimensional visualization. “SkeletalRepresentation” can be downloaded as an extension

# S-rep Extension Installation through SlicerSALT

Download the SlicerSALT packages for your respective operating system from the SlicerSALT website and install it. *SkeletalRepresentation* will be ready to use when



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# S-rep Extension Installation on 3D Slicer

1. Download 3D Slicer packages for your respective operating system on the [3D Slicer website](https://www.slicer.org/) and install it



 **3DSlicer** A multi-platform, **free and open source** software package for **visualization** and **medical image computing**

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[Download Slicer](#)

[Slicer Wiki](#)

**About Slicer**

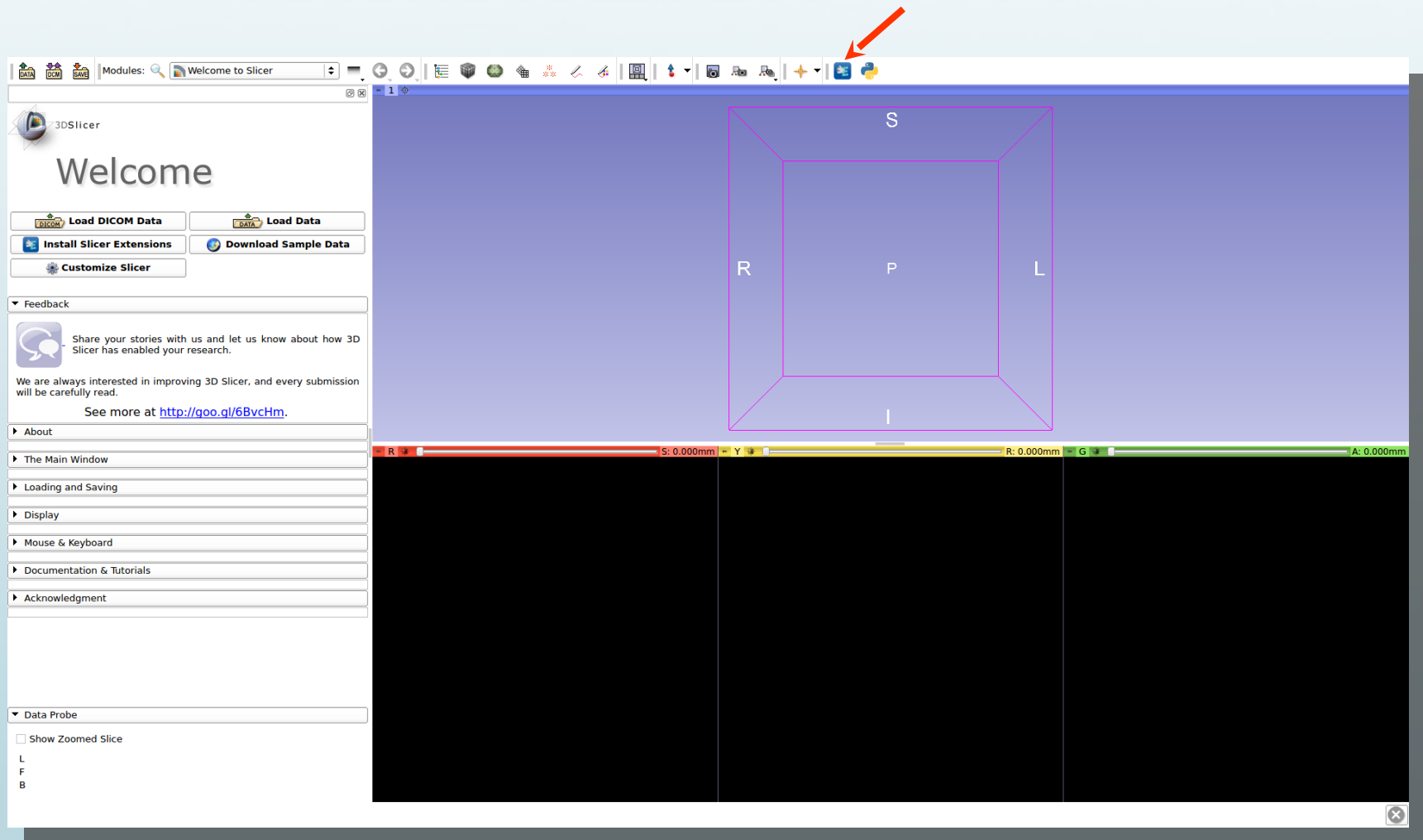
- [Introduction](#)
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**Slicer 4.6 released**

3D Slicer is an open source software platform for medical image informatics, image processing, and three-dimensional visualization. Built over two decades through support from the National Institutes of Health and a worldwide developer community, Slicer brings free, powerful cross-platform processing tools to physicians, researchers, and the general public.

# S-rep Extension Installation on 3D Slicer

2. In 3D Slicer, open the Extension Manager



# S-rep Extension Installation on 3D Slicer

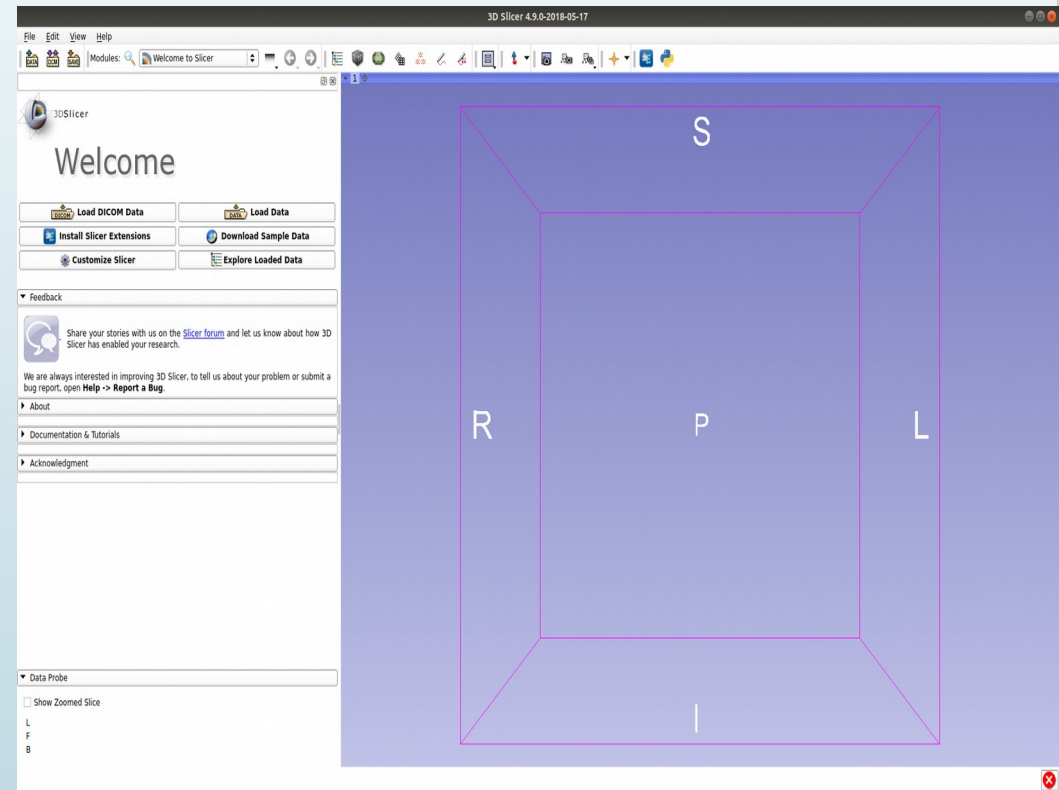
3. In the *Install Extension* tab, select **Shape Analysis** under *Categories*
4. Under **SkeletalRepresentation**, select the *Install* button and restart Slicer when prompted

# Feature 1: Visualization of s-rep

This tutorial is about features that

- Visualize an s-rep as a whole
- Toggle different parts of an s-rep

Start 3DSlicer



**Figure:** Welcome module of Slicer

# How to visualize an s-rep

## Steps:

1. Set the module path in Slicer:

*Edit->Application Settings ->Modules*

Add the position of module scripts. Then restart Slicer.

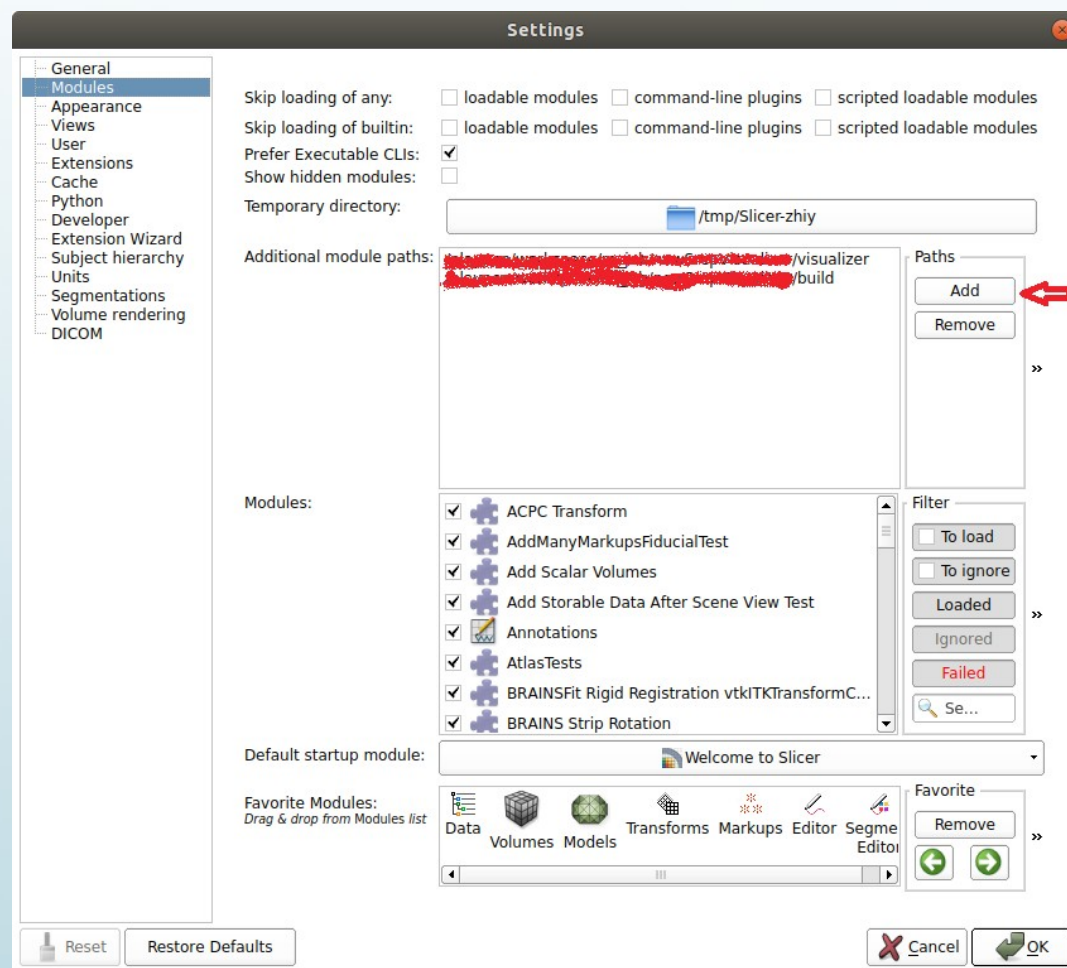


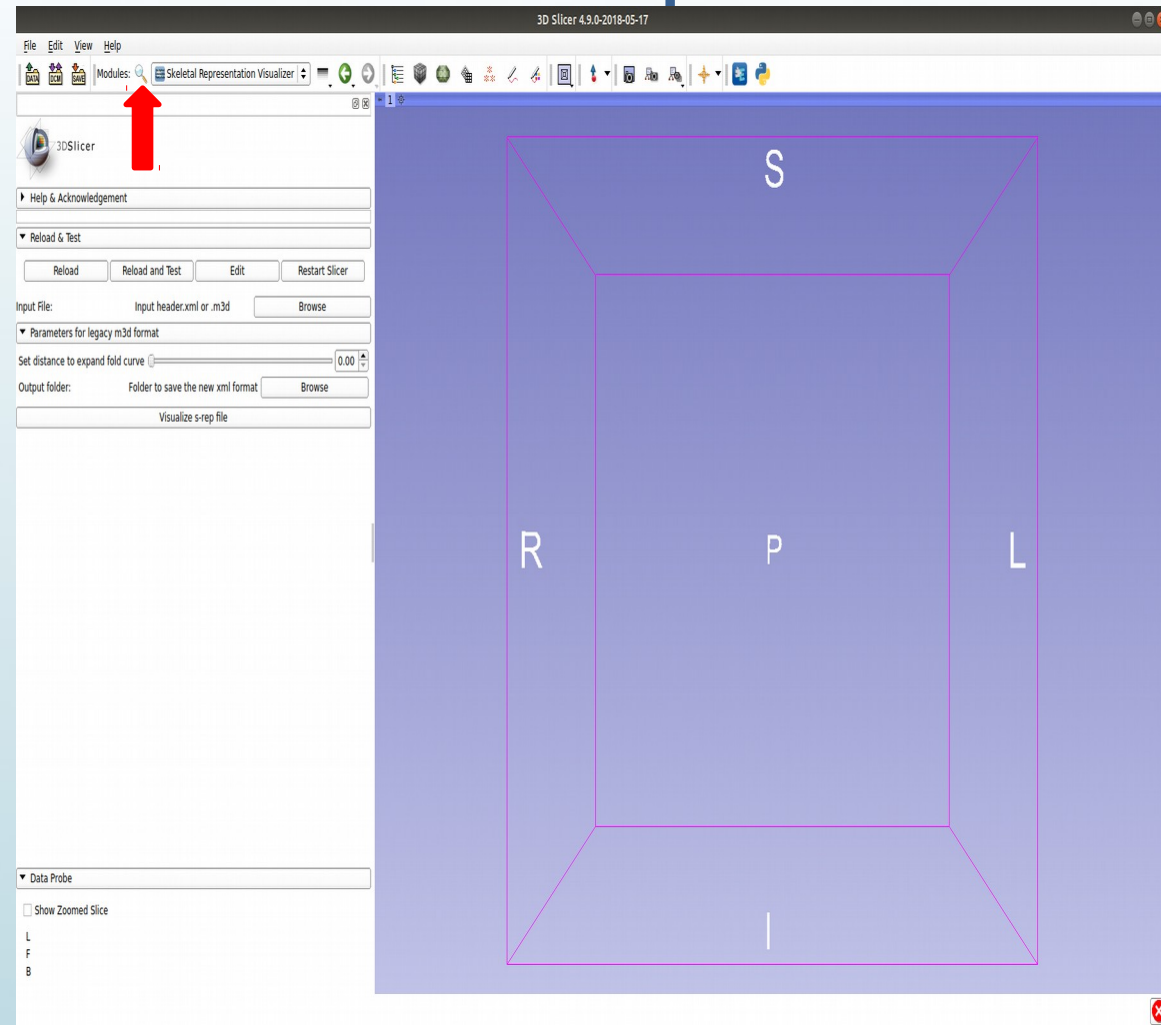
Figure: Set module path



# How to visualize an s-rep

Steps :

2. Load the *Skeletal Representation Visualizer* module in Slicer



**Figure:** Use the magnifier to search and load *visualizer* module

# How to visualize an s-rep

Steps :

3. Select an s-rep file.

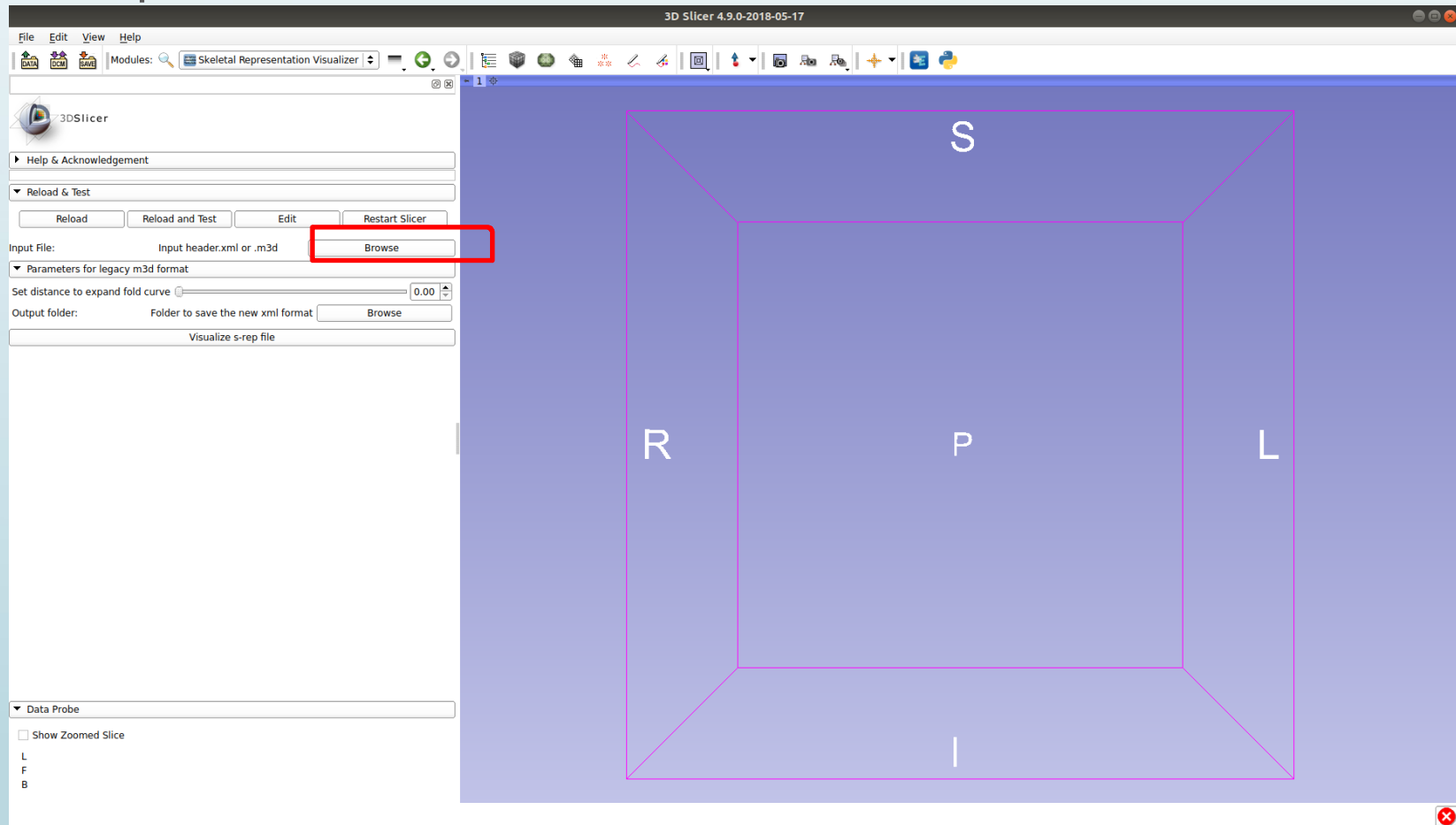


Figure: Click this button to select s-rep file

# How to visualize an s-rep

Steps :

4. Select a header file for an s-rep. In Slicer-SALT, the s-rep would be rendered in 3D window.

The header file is an xml file that stores general information of the s-rep, such as the dimension of the s-rep and the location of other related files. It looks like following:

```
<?xml version="1.0" ?>
<s-rep>
  <nRows>3</nRows>
  <nCols>8</nCols>
  <meshType>Quad</meshType>
  <color>
    <red>0</red>
    <green>0.5</green>
    <blue>0</blue>
  </color>
  <isMean>False</isMean>
  <meanStatPath/>
  <upSpoke>/home/up.vtp</upSpoke>
  <downSpoke>/home/down.vtp</downSpoke>
  <crestSpoke>/home/crest.vtp</crestSpoke>
</s-rep>
```

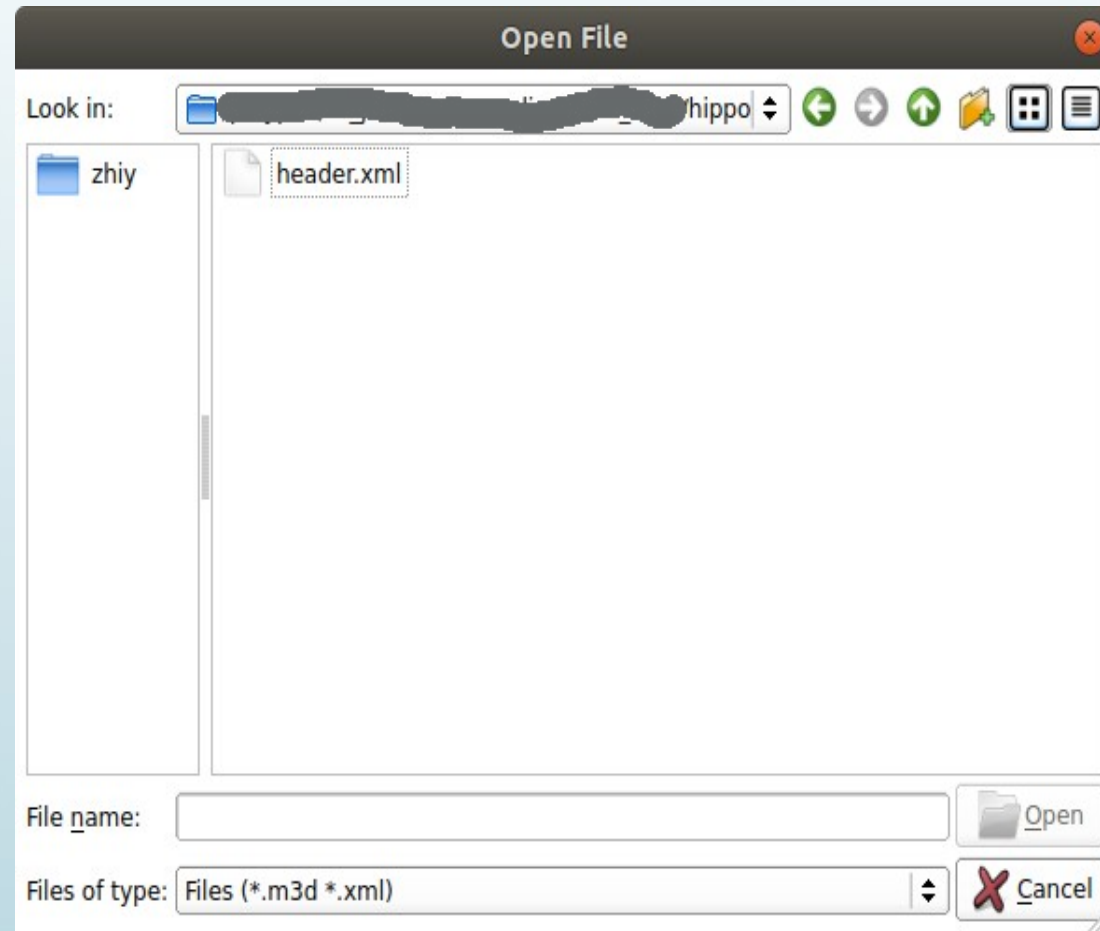
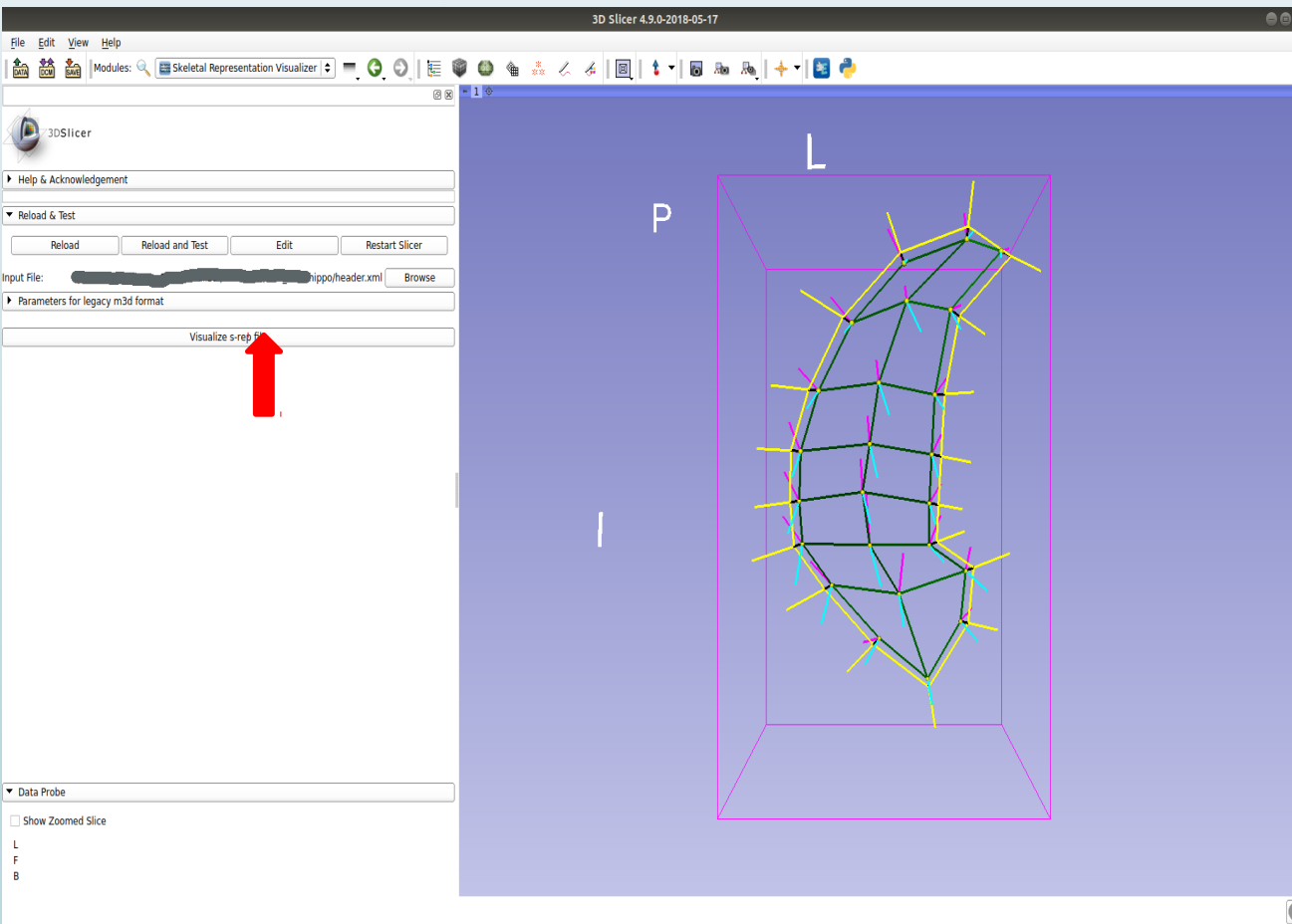


Figure: Select a new s-rep

# How to visualize an s-rep

Steps :

5. Click *Visualize s-rep file*. Center and zoom in to view s-rep in 3D window.

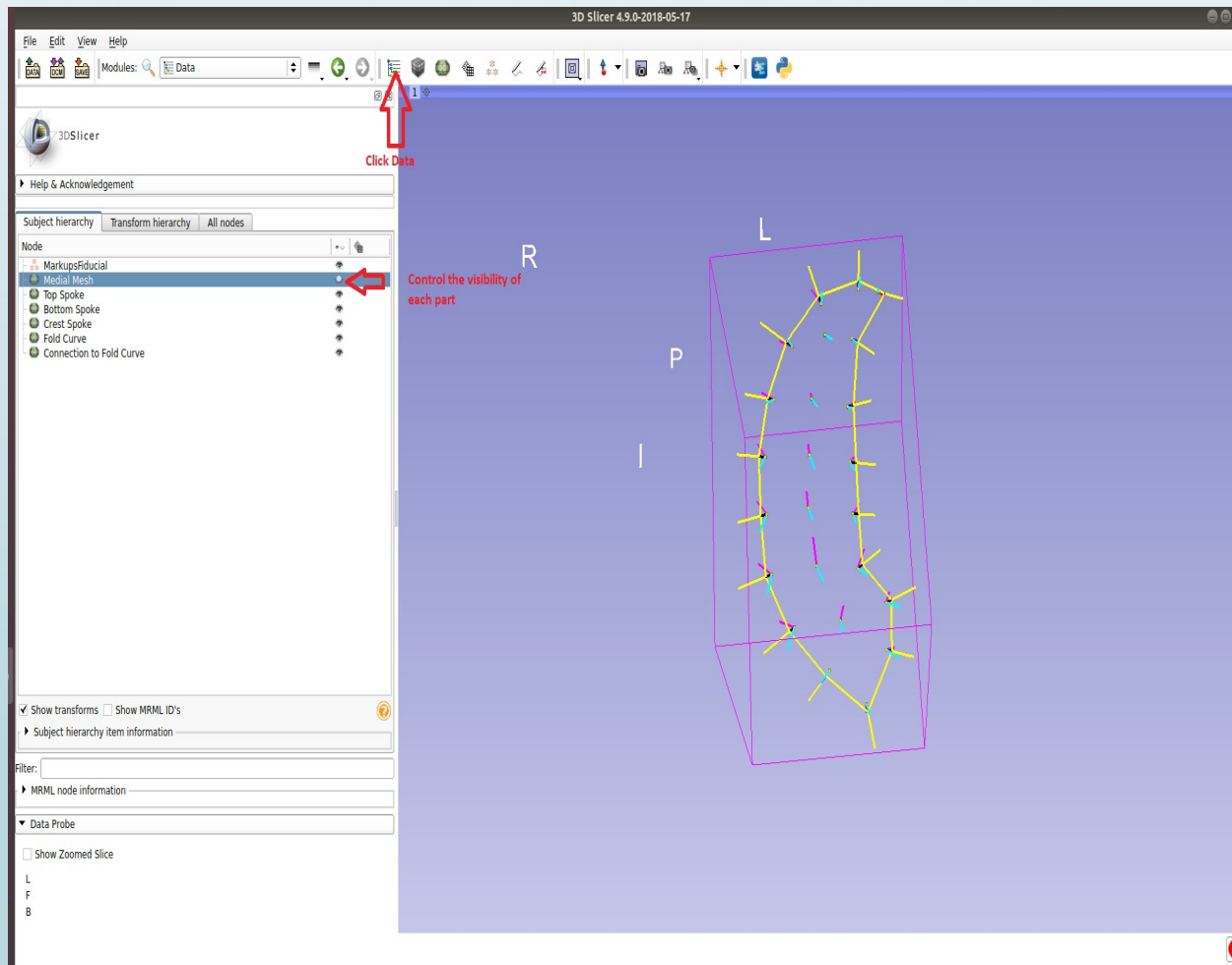


**Figure:** View the s-rep

# How to visualize an s-rep

Steps :

6. Select the s-rep part(s) to display in the *Data* tree.

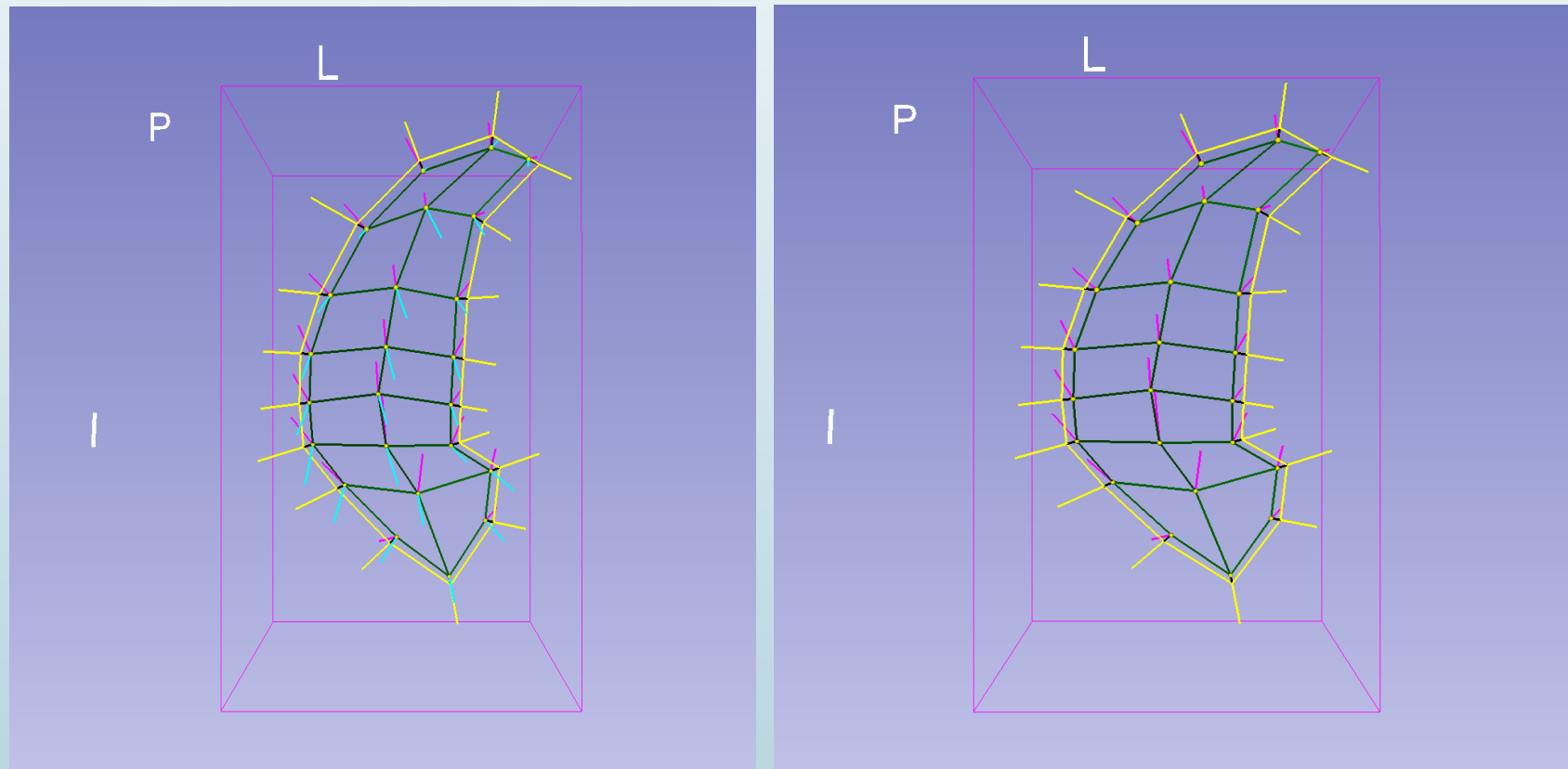


**Figure:** Illustration of how to toggle visibility of skeletal mesh in data tree

# How to visualize an s-rep

Steps :

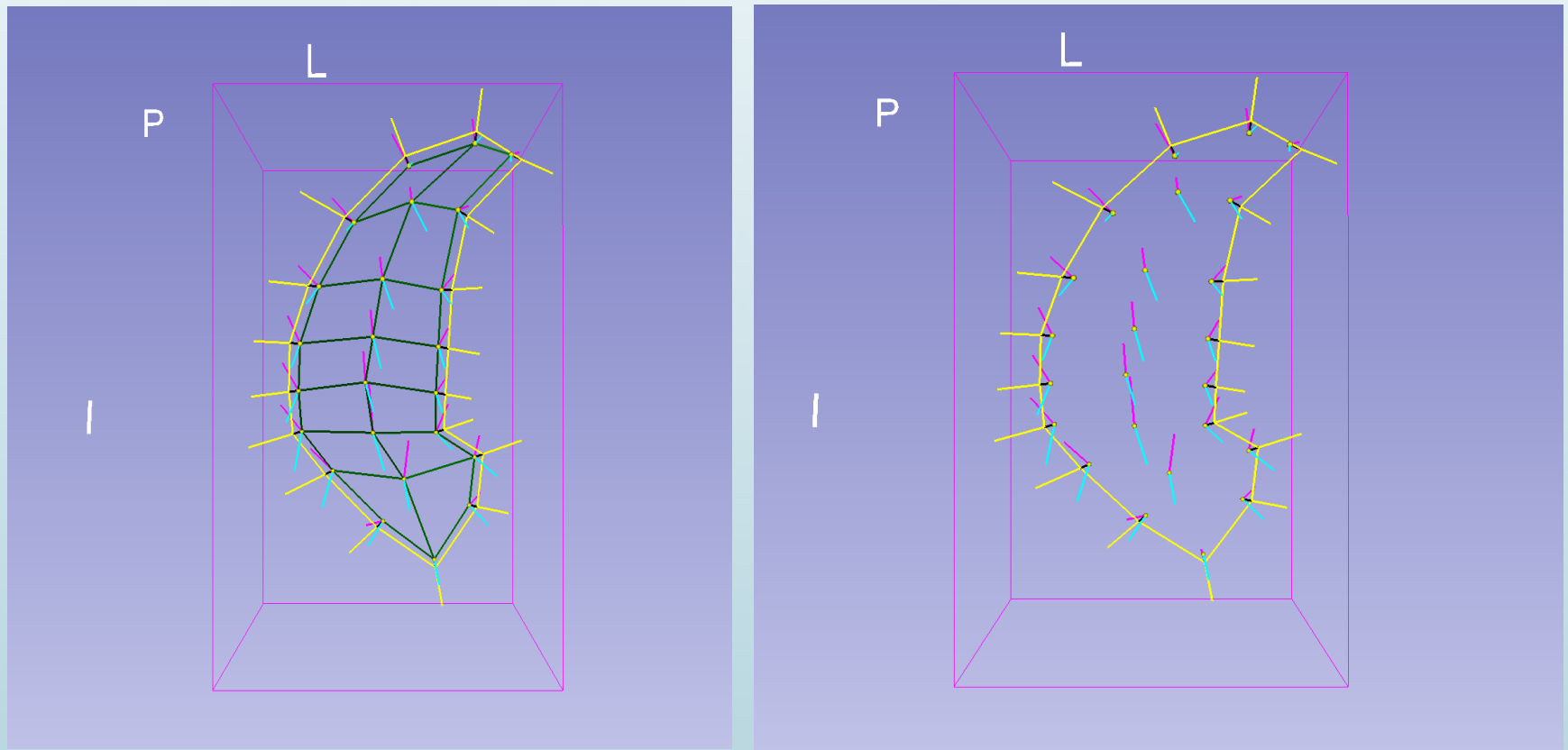
6. Select the s-rep part(s) to display in the *Data* tree.



# How to visualize an s-rep

Steps :

6. Select the s-rep part(s) to display in the *Data* tree.

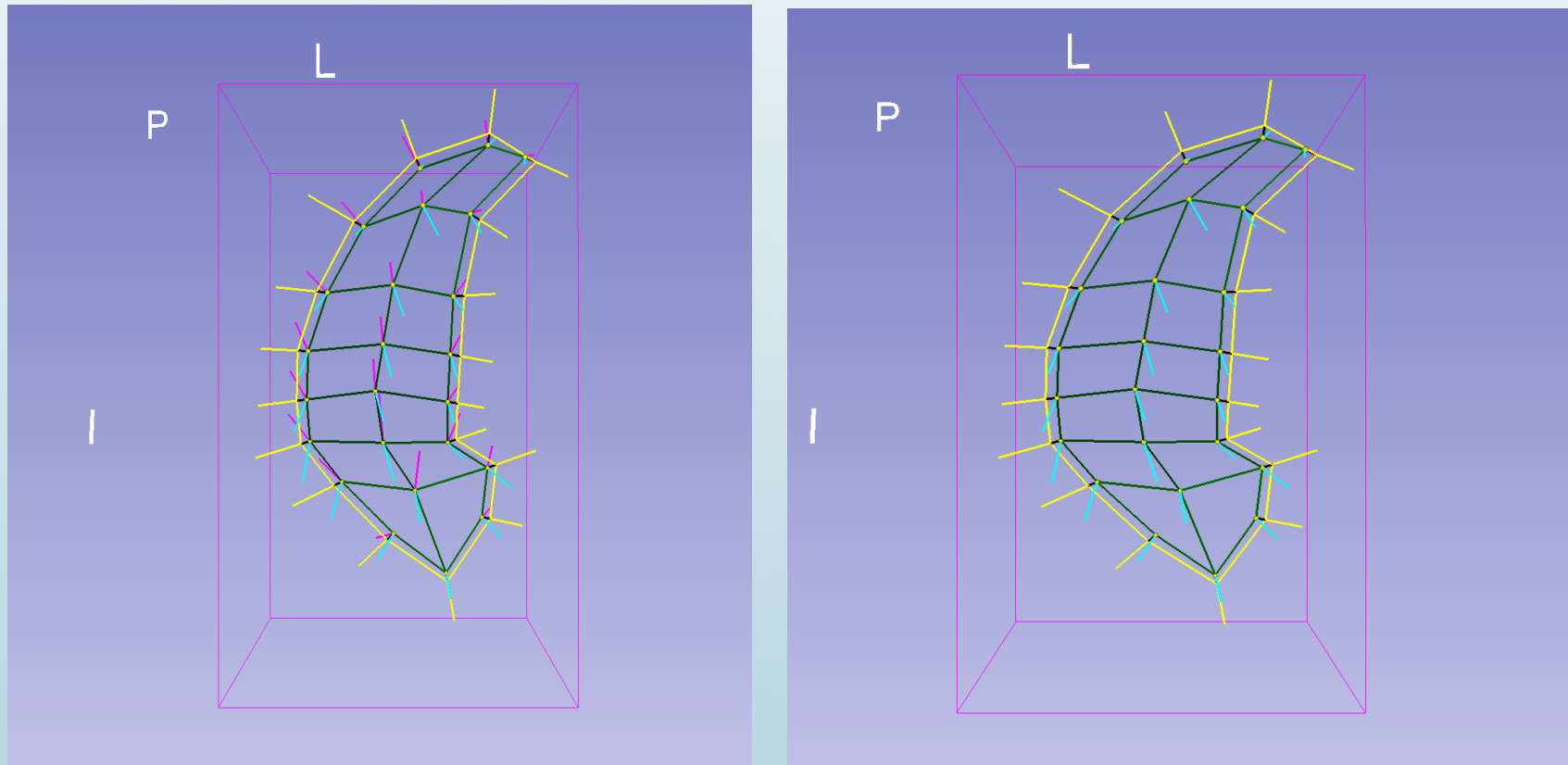




# How to visualize an s-rep

Steps :

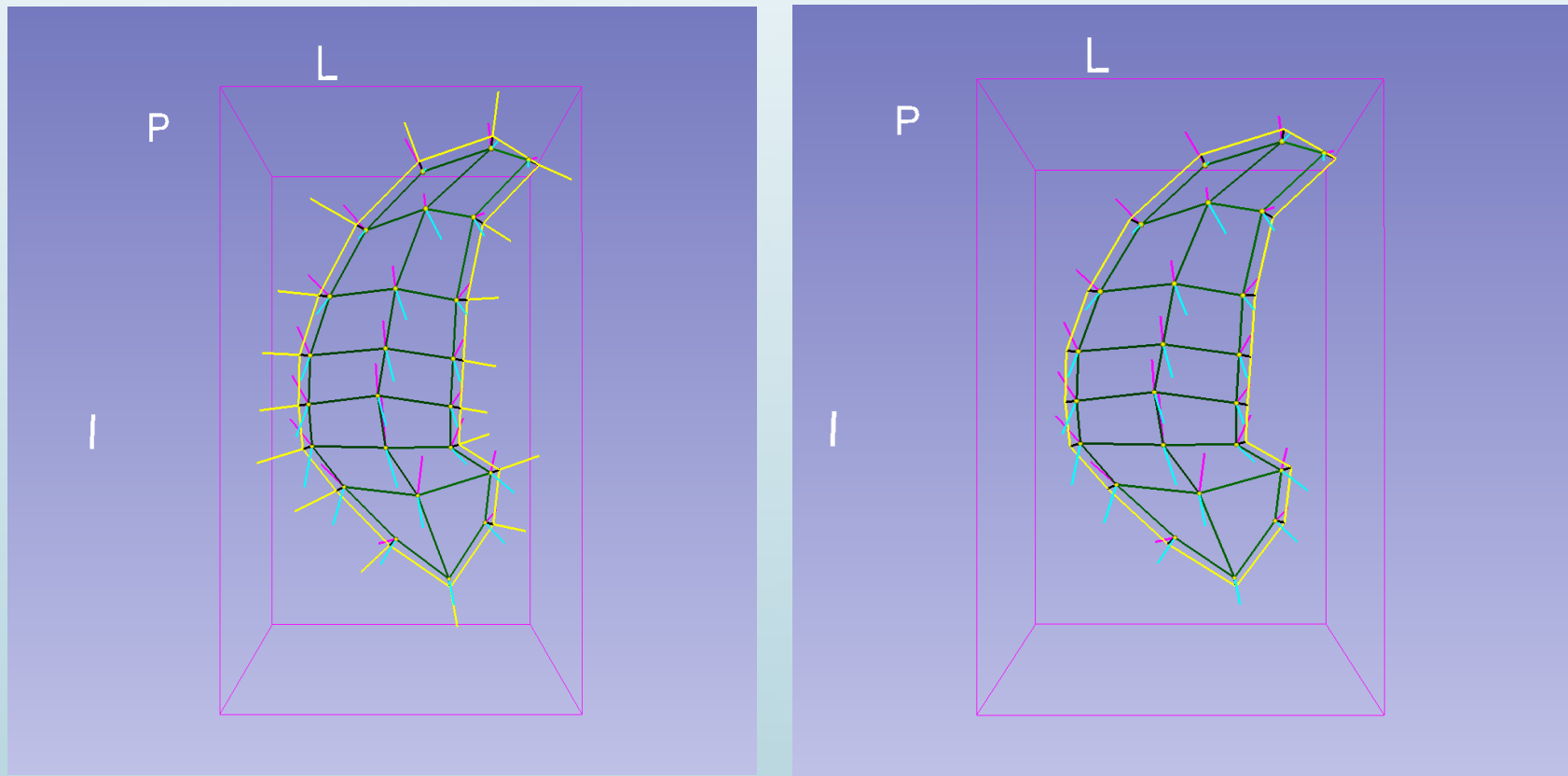
6. Select the s-rep part(s) to display in the *Data* tree.



# How to visualize an s-rep

Steps :

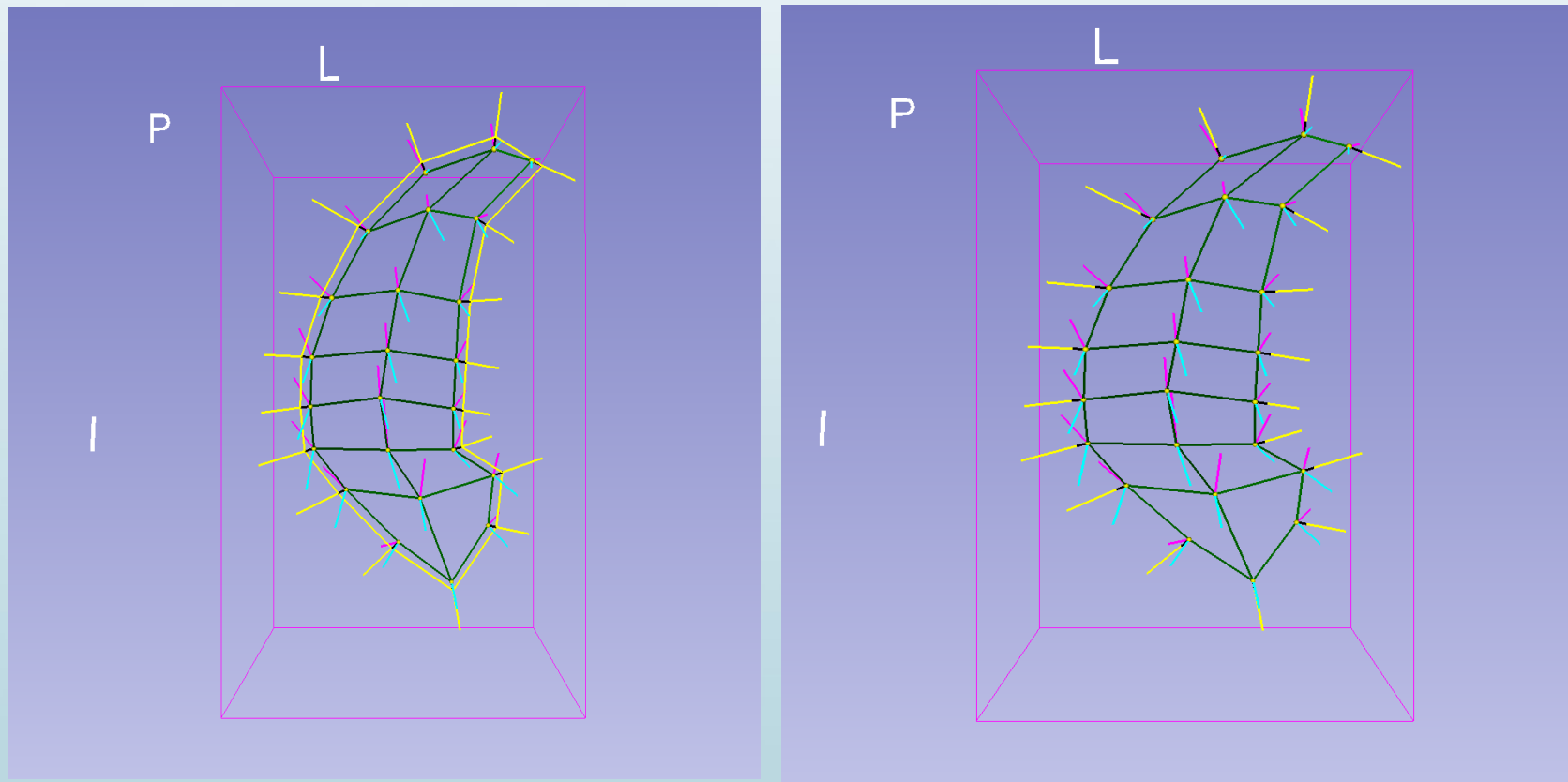
6. Select the s-rep part(s) to display in the *Data* tree.



# How to visualize an s-rep

Steps :

6. Select the s-rep part(s) to display in the *Data* tree.

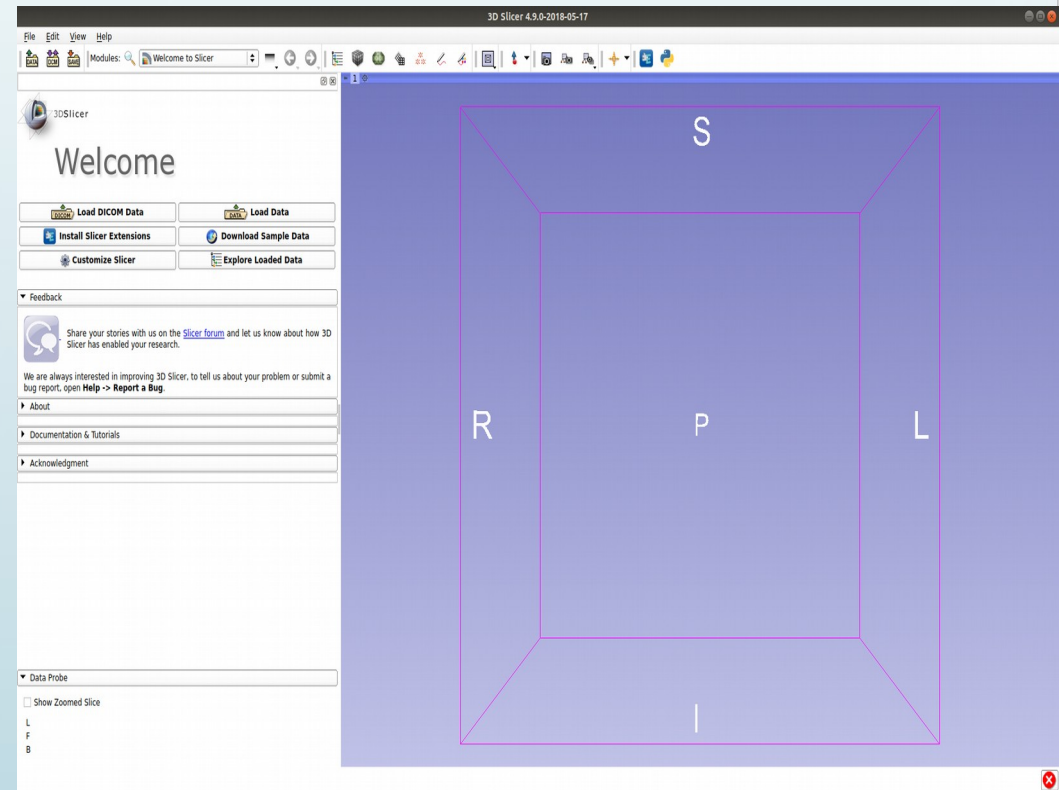


# Feature 2: S-rep filter

This function allows users to

- Transform a legacy s-rep to new s-rep, save it and visualize it as aforementioned

Start 3DSlicer



**Figure:** Welcome module of Slicer

# S-reps in the Legacy Representation

- If the visualizer is presented with an s-rep in the legacy representation (\*.m3d), auto filter and display

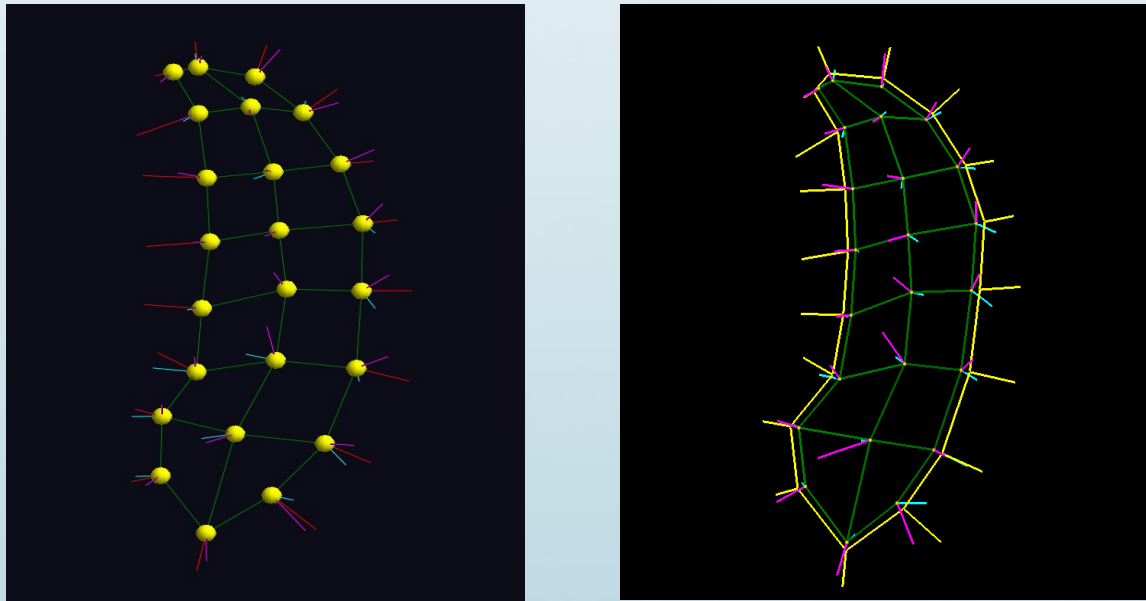


Figure: transformation from legacy s-rep to new s-rep

Left: legacy s-rep of hippocampus

Right: new s-rep of hippocampus

# How to visualize a legacy s-rep

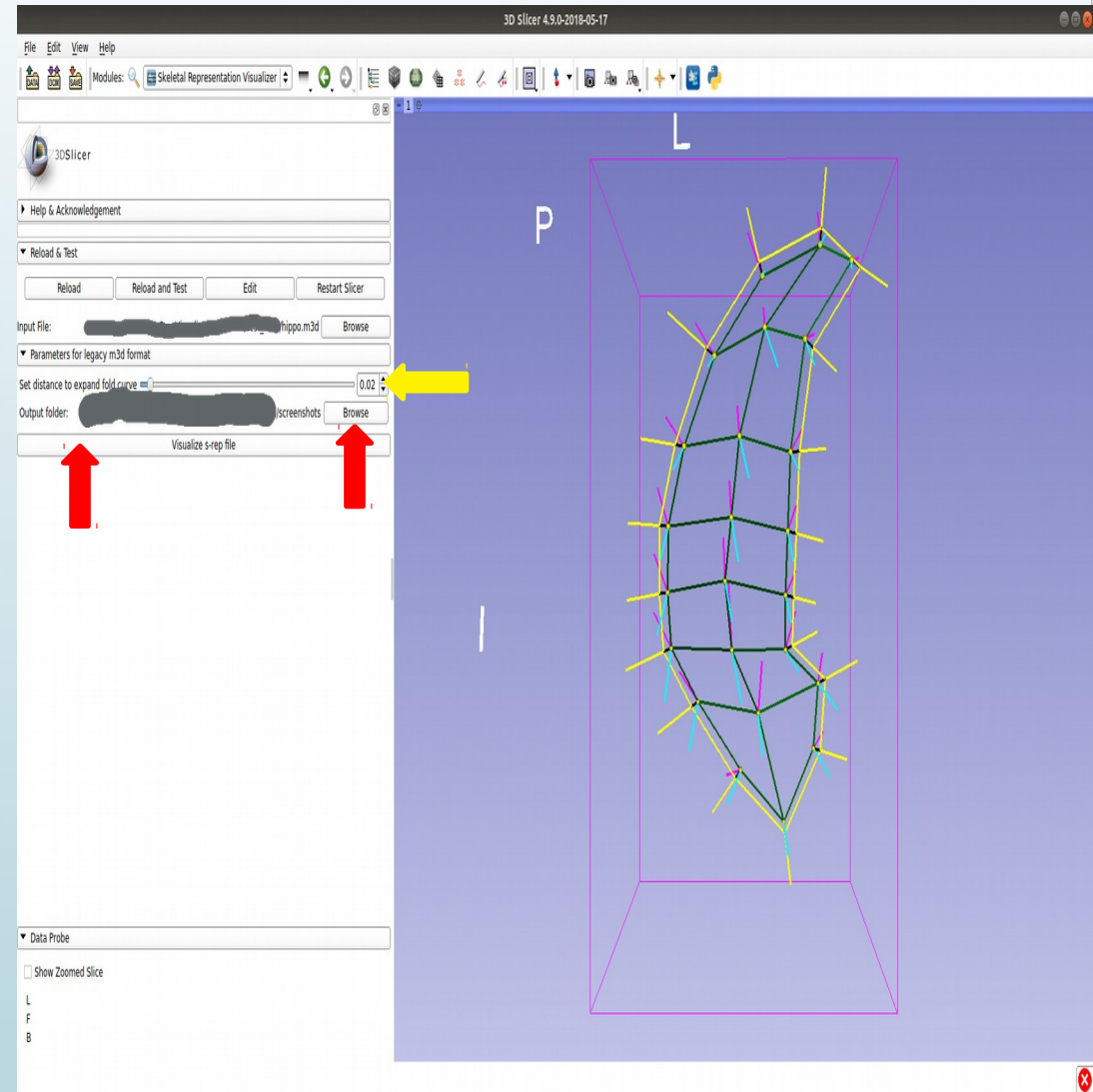
## Steps :

Select an s-rep file. If you select a legacy s-rep file (\*.m3d), the module would transform it to new s-rep and save it to the output folder. **A non-zero distance (see the yellow arrow) is recommended to expand the outer skeletal points in the legacy s-rep into the fold curve .**

*If the transformation succeeds, the s-rep will be rendered in the 3D window and related files would be saved in the output folder.*

*For hippo.m3d in [test data](#), we recommend distance as 0.02.*

*For more details of legacy s-reps, please refer section 2.1 of [this document](https://www.nitrc.org/docman/view.php/718/2065/Pablo_user_manual.pdf): [https://www.nitrc.org/docman/view.php/718/2065/Pablo\\_user\\_manual.pdf](https://www.nitrc.org/docman/view.php/718/2065/Pablo_user_manual.pdf).*



**Figure:** Transform, save and visualize a legacy s-rep

# Acknowledgements - Resources - Questions

- The S-rep module developers gratefully acknowledge funding for this project provided by NIH NIBIB R01EB021391 (Shape Analysis Toolbox for Medical Image Computing Projects), as well as the Slicer community.
- Github repository:
  - ♦ [SkeletalRepresentation](#)
  - ♦ [SlicerSALT](#)
  - ♦ [3D Slicer](#)
- Forums:
  - ♦ [SlicerSALT](#)
  - ♦ [3D Slicer](#)
- Papers & manuals:
  - [Object Shape Representation via Skeletal Models \(s-reps\) and Statistical Analysis](#)
  - [Skeletal shape correspondence through entropy](#)
  - [S-rep fitting and stats manual](#)



• For other remarks or questions, please email:

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