Rodent Cortical Thickness Analysis

Measure cortical thickness of rodent brain



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Principle

- "RodentThickness" is an automatic cortical thickness measurement tool for rat brains. The differents parts of the pipeline are :
- Preprocessing to create binary mask and label map
- Run measureThicknessFilter which produces laplacian field and thickness map in order.
- Run particle correspondence.
- Thickness sampling using particle correspondence result.
- Statistical analysis resulting in mean thickness color map and t-test result.

Different uses

- You can use Rodent Cortical Thickness Analysis :
 - With GUI
 - in Slicer (check the box noGUI)
 - in Command Line (-- nogui)

GUI

• Run the interface : ./RodentThickness



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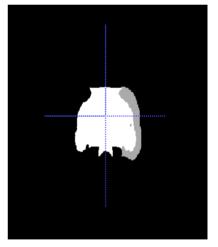
Input and Output

- The input file is the dataset CSV containig : SubjId and Input Image are required and a third input Group is needed if you compute statistical part.
- The input image is a label map of the rodent brain.
- Subjld is the name of the subject.
- The group is the name for each subjects. You must have exactly two groups if you want to compute the statistics. If you don't want to, you don't need to fill this column
- All the outputs are summarized in a csv file named outputdataset.csv

Parameters

- The label values of the binary segmentation image and the input parameters must match.
 - Cortex level (default 1).
 - Idl : Id number for the low Dirichlet boundary condition : it's the rest of the brain (default 4).
 - Idh : Id number for the high Dirichlet boundary condition: it's the background (default 0).

Label Map



- Grey : Cortex
- White : Brain
- Black : Outside

Load CSV File if available

• You can load a CSV file containing paths to a Dataset you saved before or you wrote yourself with this style:

subjld,labelMapInput,group 1,image1.nrrd,A 2,image2.nrrd,A 3,image3.nrrd,B 4,image4.nrrd,B



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Else configuring images to process

- Clicking on the table will allow you to select images, or enter the subject and the group.
- You have to fill the column group only if you check the box "Compute Statistics"

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Save CSV File

File Help

 You can save your Dataset into a CSV file so you can load it and use it again later.

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Add and delete row

• If you select a row , you can add and delete a row at this place , if no row are selected it is on the last of the table.



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Give an output folder

• You need to give an output folder: folders will be created in your output folder, and all the files generated by the program will be put in it.

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Statistical Analysis

- If you check the box Statistical , you have to fill the column Group.
- Statistic part needs two different groups.
- If you don't compute the statistics , you don't have to fill the column Group.



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Software configuration

- If you have your own version of the programs or if you need to use a particular version of it, you can write the path manually or click the button to search it.
- If you don't compute the statistical part, you don't have to fill the path for Rscript (2), concatToColumns and vtkPointAttributes.



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ParaToSPHARMMeshCLP		R
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ShapeWorksGroom		R
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Automatic Configuration

 By clicking the "Default" button, the program will automatically search all the programs in the PATH, and tell you if some of them are missing.

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Automatic Configuration

 If you want to reset the path for one or more programs, just push the "R" button and it will search the corresponding program in the PATH.

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Directory containing Batchmake Script	
Output Folder	
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Load and Save

- Save your configuration: In the « File » Menu, you can Save or Load a configuration file generated by the program.
- When opening the program, it will automatically search and load any file called "RodentThicknessconfigfile.bms" in the directory where the executable is and in the current work directory.



Software Configuration

- Python version : Python 2.6 or Python 2.7 with vtk installed.
- If you want to compute Statistics , you need to put niralvtk.py in the same folder as vtkPointAttributes.py.



Compute

 When you fill the table, the configuration and gave an output folder, you can compute
 Rodent Thickness by clicking the "Compute"
 button. It will test the existence and type of the files and the folders and tell you if some files are not images.

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Output	Folder					
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Command line

« RodentThickness --help » :

•Load :

--dataset <std ::string> : CSV file containing the dataset

- --configfile <std::string> : Software configuration file
- --PathBms <std::string> : Path where Batchmake scripts are

•Options:

--nogui : If you do not need the GUI (default: 0)

--CortexLevel : Label cortex level

--idl : Label Brain level

--ids : Label Outside level

•Output :

--WorkDir <std::string> Output Directory



No Gui Mode

- In « No GUI » mode, you need to set the dataset, the configile and the path for BatchMake Script by giving these in command line : DTIAtlasBuilder --nogui -dataset -- configile -- PathBms
- The program will not display the GUI and will run automatically with the given parameters, as if you had pushed the « Compute » button.



Slicer

- You can use RodentThickness in Slicer .
- You can download Slicer on : http://download.slicer.org/
- Open Slicer.
- On "Application Settings", click on "Modules" and add the path of the executable on "Additional Module Path".
- Close Slicer and reopen it .

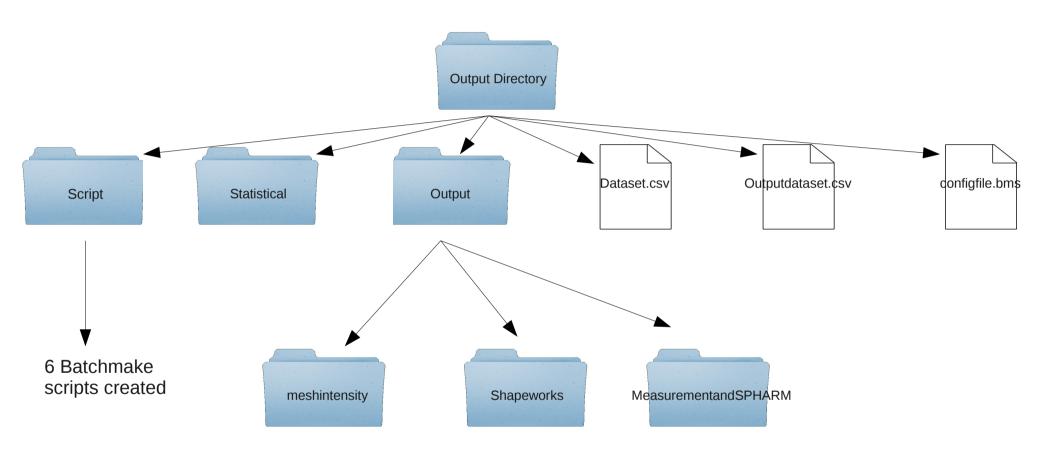
Slicer

- After added the path , click on Modules and in "All Modules " choose "Rodent Cortical Thickness Analysis"
- You must complete four parameters:
 - dataset : CSV file containing the input
 - configfile : the configuration file for executable path
 - WorkDir : The path of the Output Directory
 - -PathBms : the directory containing Batchmake script

Frequent Problem

- You will have a message box with "Item Empty" if one row are empty. Please delete the row or fill all the item of the row (except group if Statistic box is not check).
- You will have a message box with "Corrupt File" if the config file is not good, please save one for see the form and the order of the config file.
- You will have a message "Could not open csv file" if we can't read it or because the first line is not : subjld labelMapInput group

Data organisation





Output Directory

3 Folders in the directory :

MeasurementandSPHARM
\${subject}_measurementoutput.nrrd : Computed thickness volume
\${subject}_para.vtk : Mesh for the spherical parametrization
\${subject}_surf.vtk : Mesh for the surface
\${subject}.ip.SPHARM.vtk : output is a series of SPHARM coefficients and SPHARM-PDM meshes
\${subject}.subj.SPHARM.vtk

shapeworks\${subject}.correspondence.vtk\${subject}.warped.vtk

meshintensity \${subject}.txt
\${subject}.sampling.vtk : Mesh output with actual sampling points

Statistical Directory

4 files have created :

- data.{group}.txt : Table contains Value of thickness measurement for each point and each subjects of the group
- listThicknessSamplingResult_{group}.txt : File contains the path for \${subject}.txt
- stats.txt : Average of thickness measurement for each point for all the subject in the group

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- stats.vtk
 In the file stats.txt , you have the results :
 - ctrmean : mean of thickness measurement for the first group
 - exprmean : mean of thickness measurement for the second group
 - t.pvalues : pvalue if the number of subject is more than 20
 - w.pvalues : pvalue if the number of subject is less than 20

Rodent Thickness on line

• GitHub :

https://github.com/mjacquem/RodentThickness NITRC :

https://www.nitrc.org/projects/rodentthickness/

