



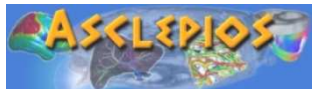
# **Interfaces development for medical imaging Applications in the framework of the Multiple Sclerosis disease**

**Erik Pernod<sup>1</sup>**

**Supervisor: Jean-Christophe Souplet<sup>2</sup>**

<sup>1</sup>Student engineer in “Calcul Scientifique”, ISITV, Toulon

<sup>2</sup>Asclépios Project team, INRIA, Sophia Antipolis



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Thursday the 4<sup>th</sup> of September, 2008



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## **INRIA**

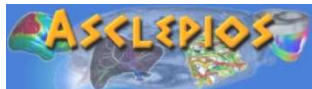
<http://www.inria.fr>

- Institut National de Recherche en Informatique et en automatique.
- Goals:
  - Dedicated to fundamental and applied research.
  - Play a major role in technology transfer.

## **Asclepios**

<http://www-sop.inria.fr/asclepios/>

- In medical image processing domain.
- Goals:
  - Analysis of medical and biomedical images with advanced geometrical, statistical, physical and functional models.
  - Provides optimized tools to clinicians



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## **NeuroLOG** <http://neurolog.polytech.unice.fr>

- A three years ANR scientific project (2007-2009)
- Goal: Federate medical data and algorithms, and sharing computing resources on grid infrastructure.
- On three different pathologies:
  - Multiple Sclerosis
  - Brain Stroke
  - Tumours
- Partners from different disciplines:
  - Software technologies
  - Databases and knowledge
  - Medical imaging



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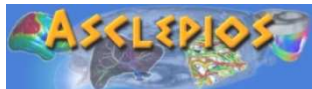


## My work

- To understand an application of brain MRI segmentation by implementing tools in the software SepINRIA.
- To deploy this application on the EGEE computational grid

## Plan

- Brain MRI segmentation pipeline
- SepINRIA software
- Workflow deployment on the EGEE grid
- Time performance and study of a method's parameter influence
- Conclusion



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# The pipeline

1- Introduction

2- The pipeline

3- SepINRIA

4- Workflow creation

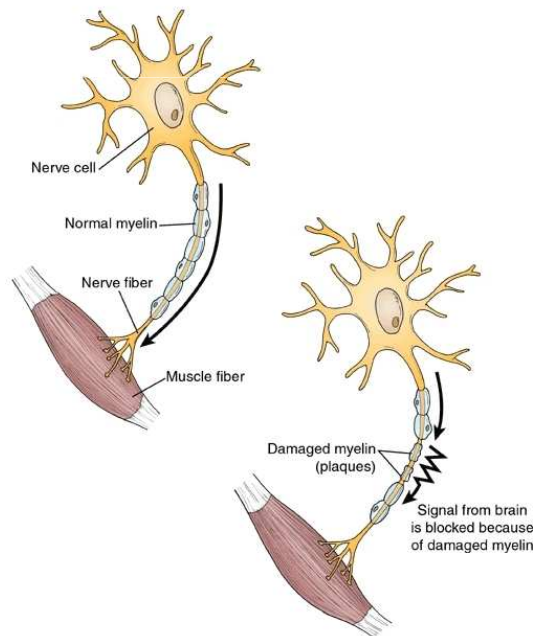
5- Performances

6- Conclusion

## Multiple Sclerosis (MS) brain MRI segmentation

- Segmentation of lesions on brain MRI is required for diagnosis or follow-up purpose in MS.

Damaged Myelin in Multiple Sclerosis



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*Improved EM-based tissue segmentation and partial volume effect quantification in multi-sequence brain MRI.*  
G. Dugas-Phocion, M. Angel G. Ballester, G. Malandain, C. Lebrun, and N. Ayache. MICCAI'04.

# The pipeline

1- Introduction

2- The pipeline

3- SepINRIA

4- Workflow creation

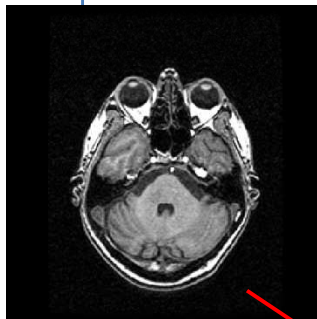
5- Performances

6- Conclusion

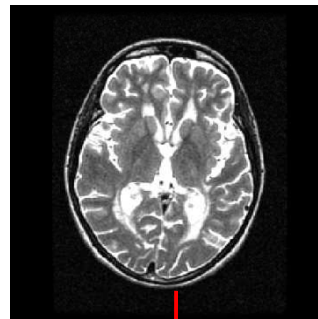
## Brain MRI segmentation pipeline: Five main steps

A rigid registration of the T1 on the T2 sequence is performed.

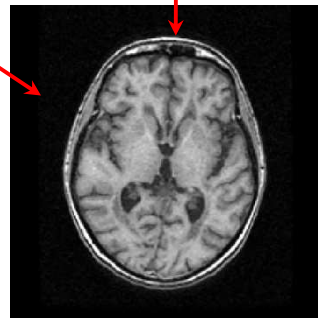
Spatial normalization



T1 sequence



T2 sequence



T1 sequence registered

*Block matching: A general framework to improve robustness of rigid registration of medical images.*

*S. Ourselin, A. Roche, S. Prima, and N. Ayache. MICCAI'00*

# The pipeline

1- Introduction

2- The pipeline

3- SepINRIA

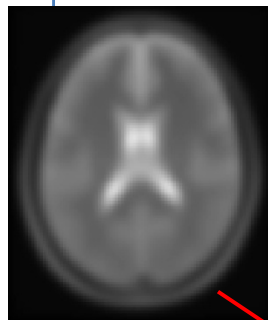
4- Workflow creation

5- Performances

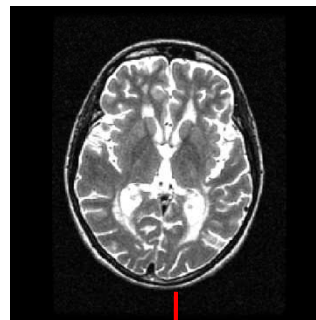
6- Conclusion

## Brain MRI segmentation pipeline: Five main steps

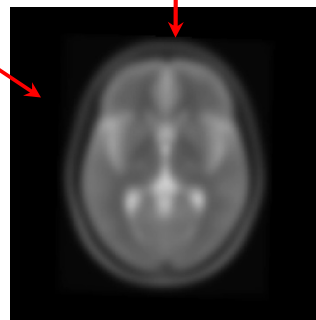
A rigid registration of the atlas images on the T2 sequence is performed.



Atlas T2 sequence



T2 sequence



Atlas T2 sequence registered

Spatial normalization

MNI atlas registration

# The pipeline

1- Introduction

2- The pipeline

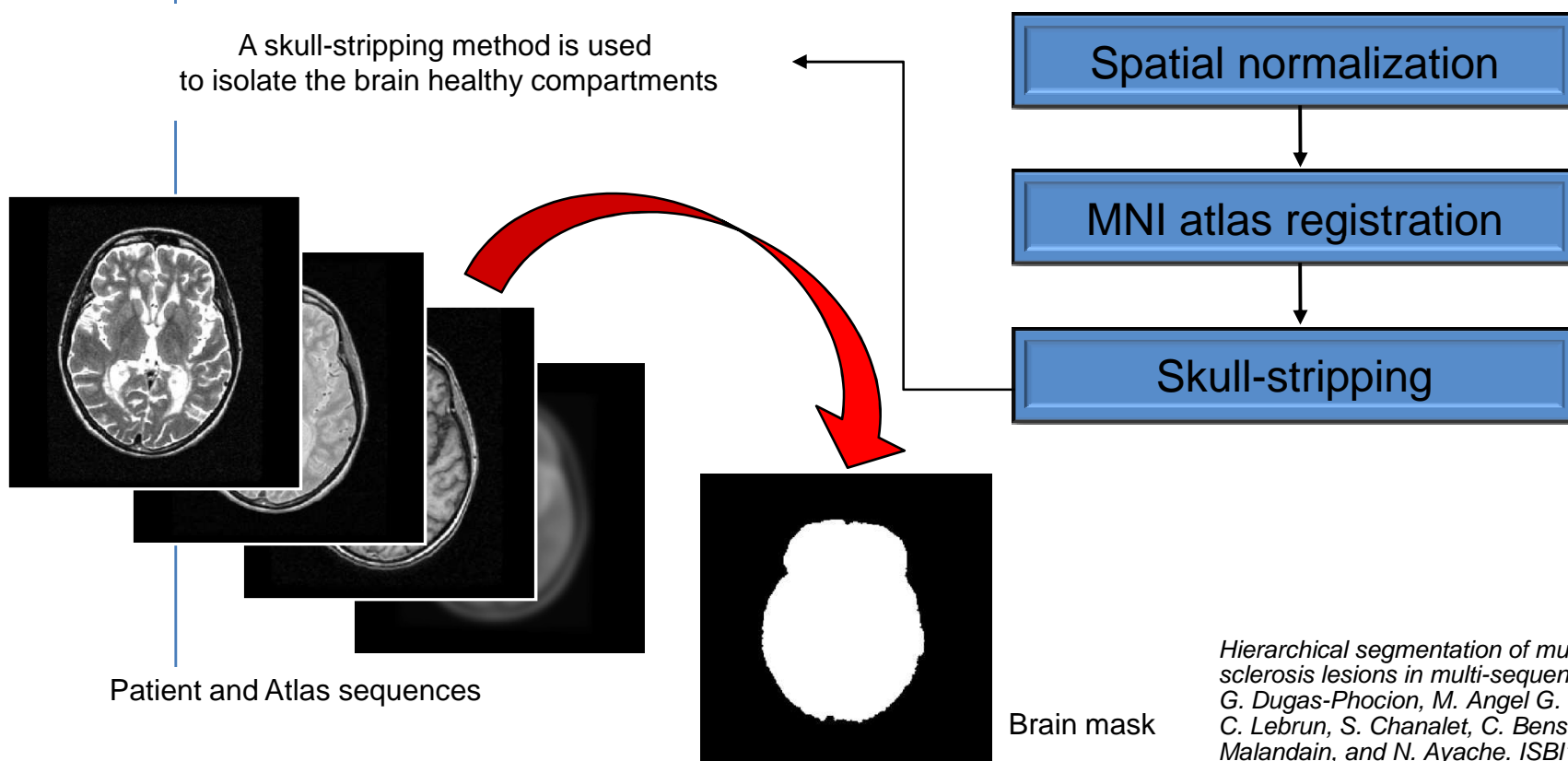
3- SepINRIA

4- Workflow creation

5- Performances

6- Conclusion

## Brain MRI segmentation pipeline: Five main steps



*Hierarchical segmentation of multiple sclerosis lesions in multi-sequence MRI. G. Dugas-Phocion, M. Angel G. Ballester, C. Lebrun, S. Chanalet, C. Bensa, G. Malandain, and N. Ayache. ISBI'04.*

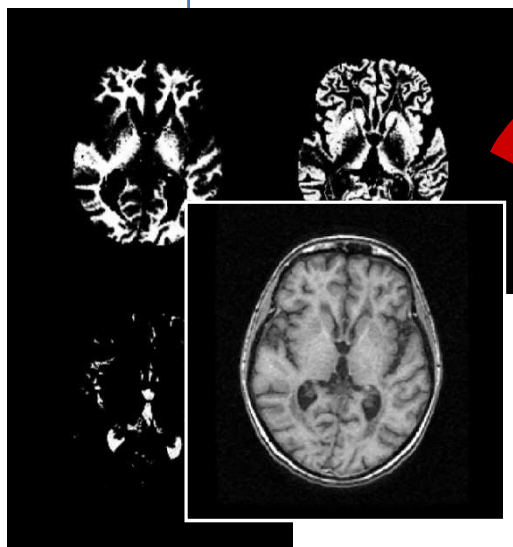
# The pipeline

1- Introduction 2- The pipeline 3- SepINRIA 4- Workflow creation 5- Performances 6- Conclusion

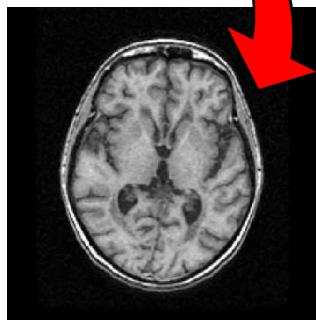
## Brain MRI segmentation pipeline: Five main steps

A first classification of the brain into WM, GM and CSF classes is realized.

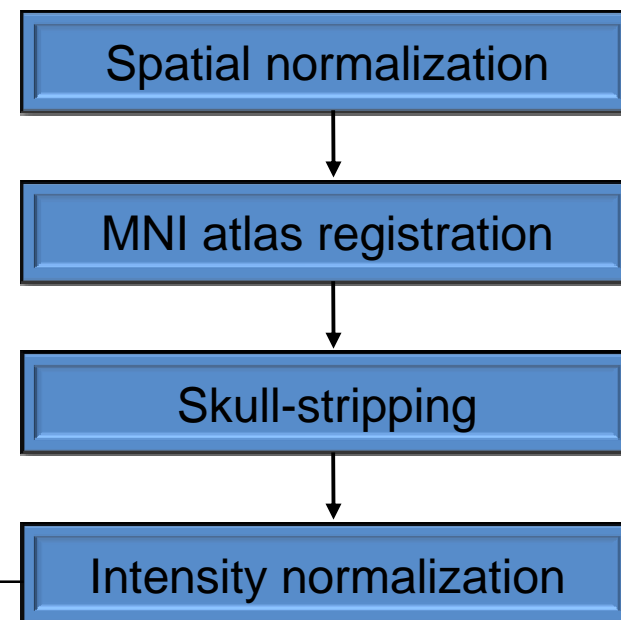
Parameters of bias field are then computed from these segmentations.



T1 sequence and brain compartments



T1 sequence unbiased



Maximum likelihood estimation of the bias field in MR brain images:  
Investigating different modelings of the imaging process.  
S. Prima, N. Ayache, T. Barrick, and N. Roberts. MICCAI'01

# The pipeline

1- Introduction

2- The pipeline

3- SepINRIA

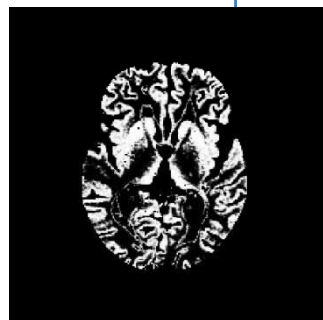
4- Workflow creation

5- Performances

6- Conclusion

## Brain MRI segmentation pipeline: Five main steps

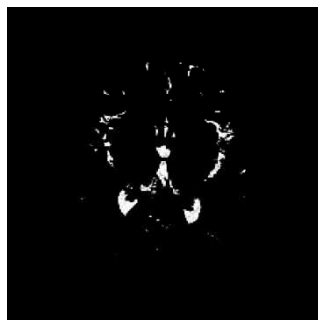
The Expectation Maximization method is used once again to classify brain MRI voxels from unbiased images.



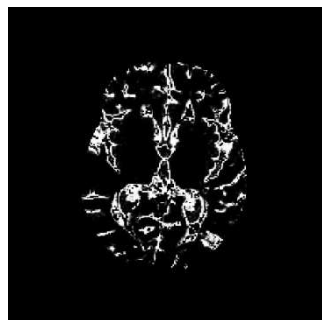
Grey matter



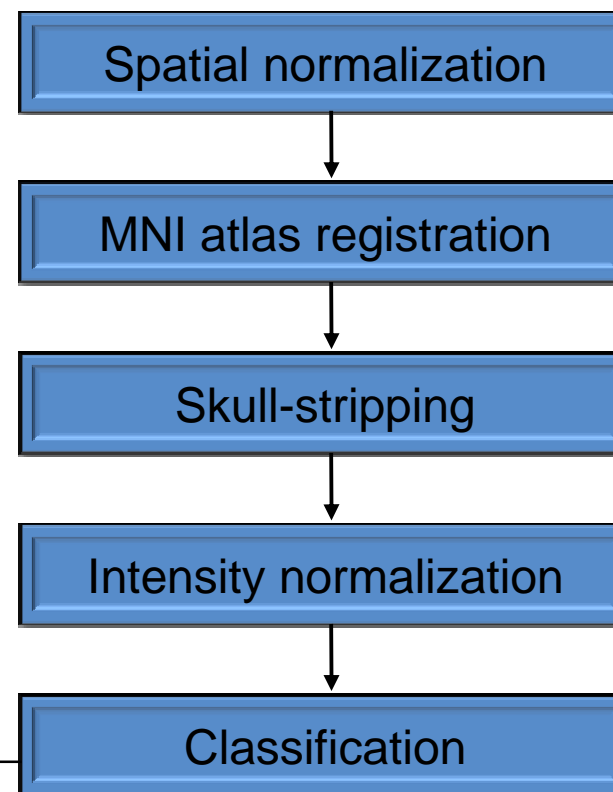
White matter



CSF



PVE



# SepINRIA

1- Introduction

2- The pipeline

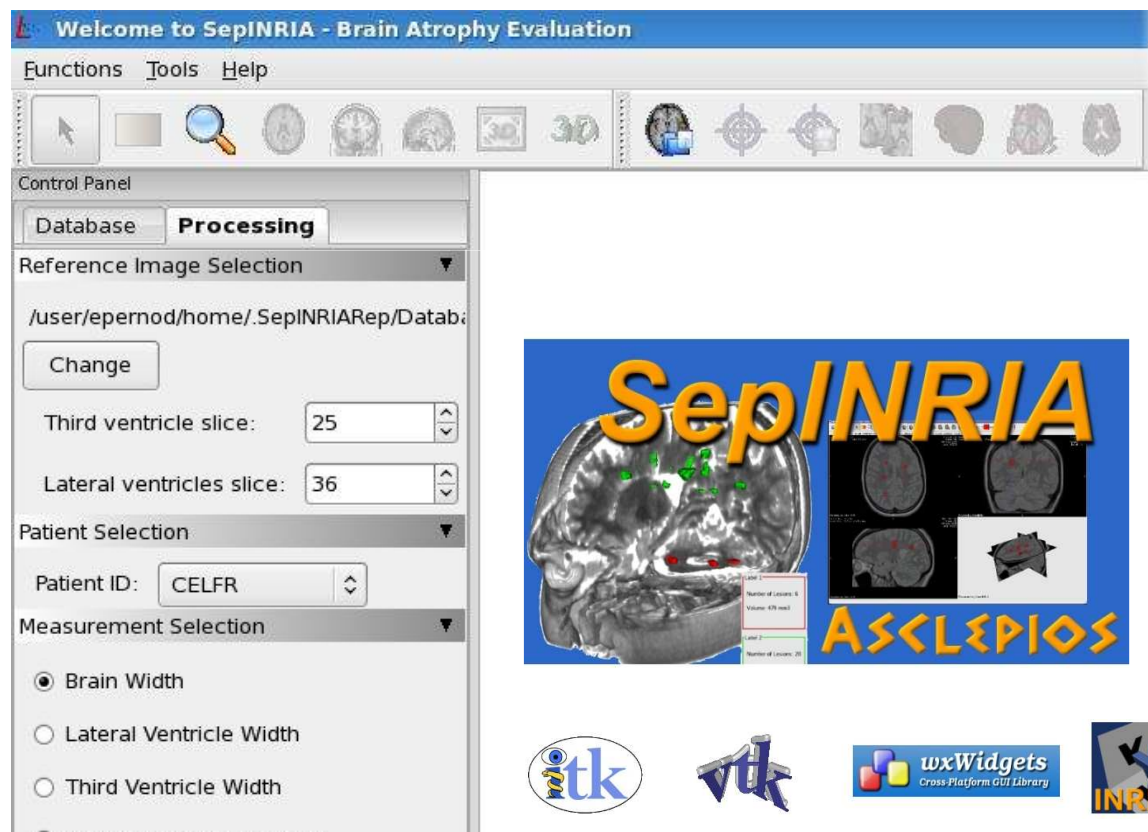
3- SepINRIA

4- Workflow creation

5- Performances

6- Conclusion

## SepINRIA improvement: Automatic atrophy measurement functionality (Created by Jean-Christophe Souplet)



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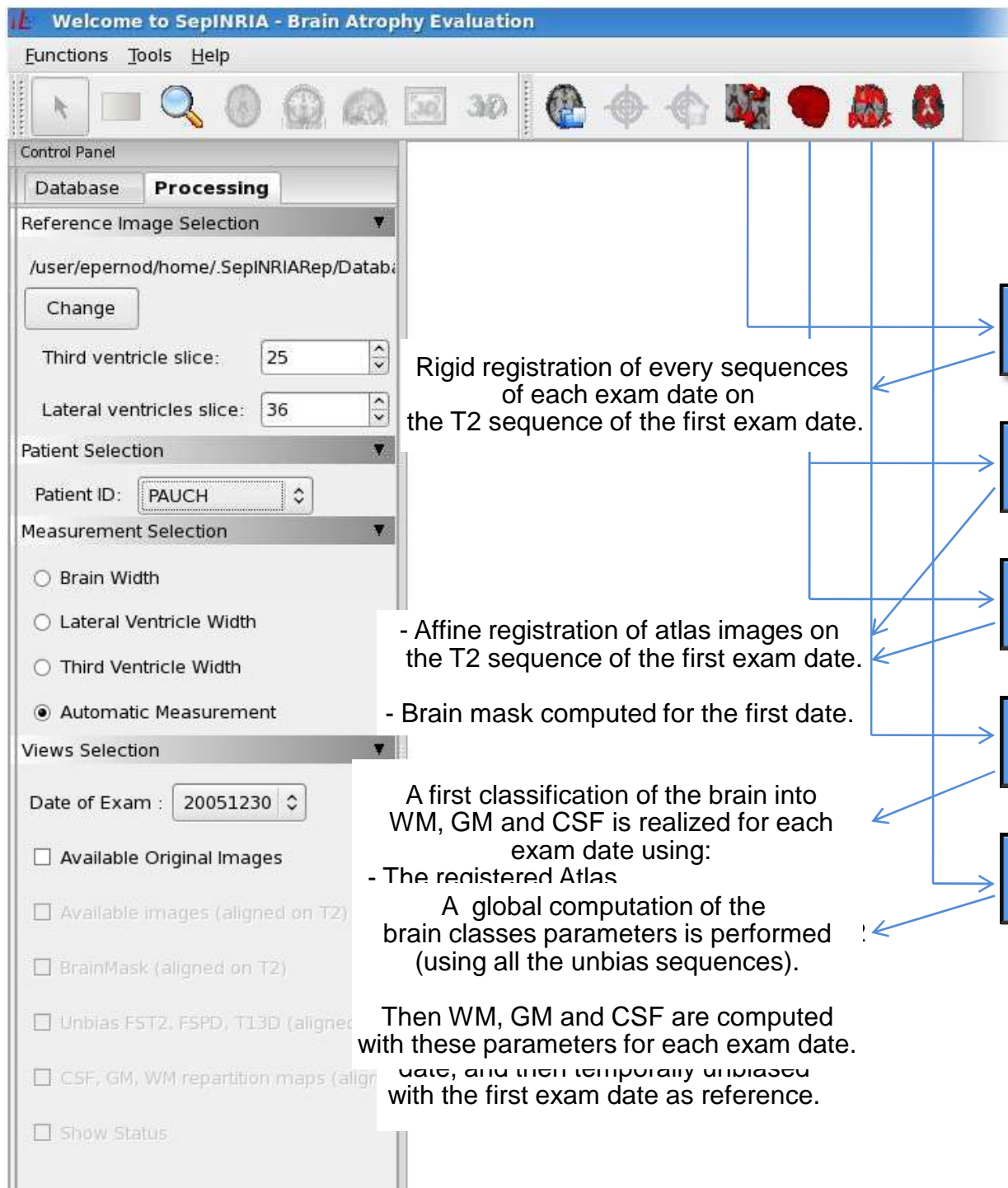


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6- Conclusion

Rigid registration of every sequences of each exam date on the T2 sequence of the first exam date.

- Affine registration of atlas images on the T2 sequence of the first exam date.

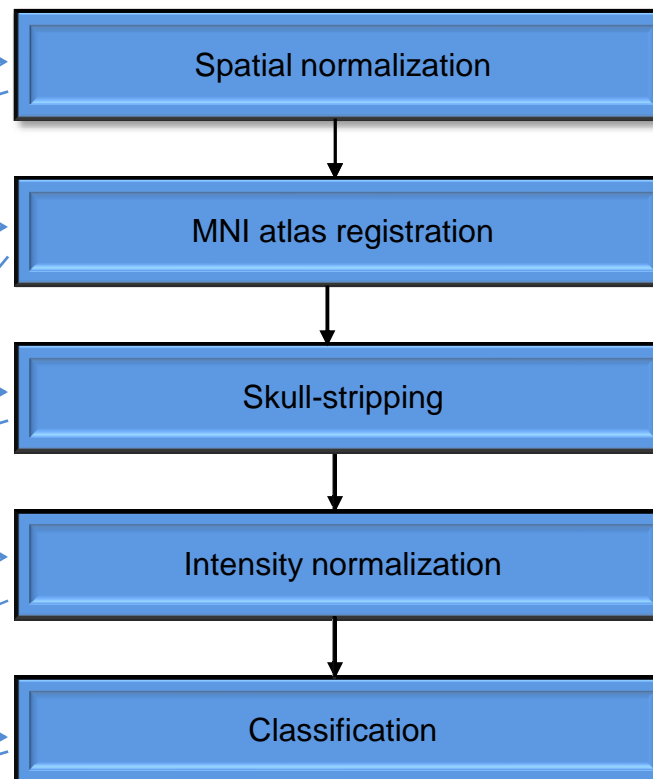
- Brain mask computed for the first date.

A first classification of the brain into WM, GM and CSF is realized for each exam date using:

- The registered Atlas

A global computation of the brain classes parameters is performed (using all the unbiased sequences).

Then WM, GM and CSF are computed with these parameters for each exam date. date, and then temporarily unbiased with the first exam date as reference.



# SepINRIA

1- Introduction

2- The pipeline

3- SepINRIA

4- Workflow creation

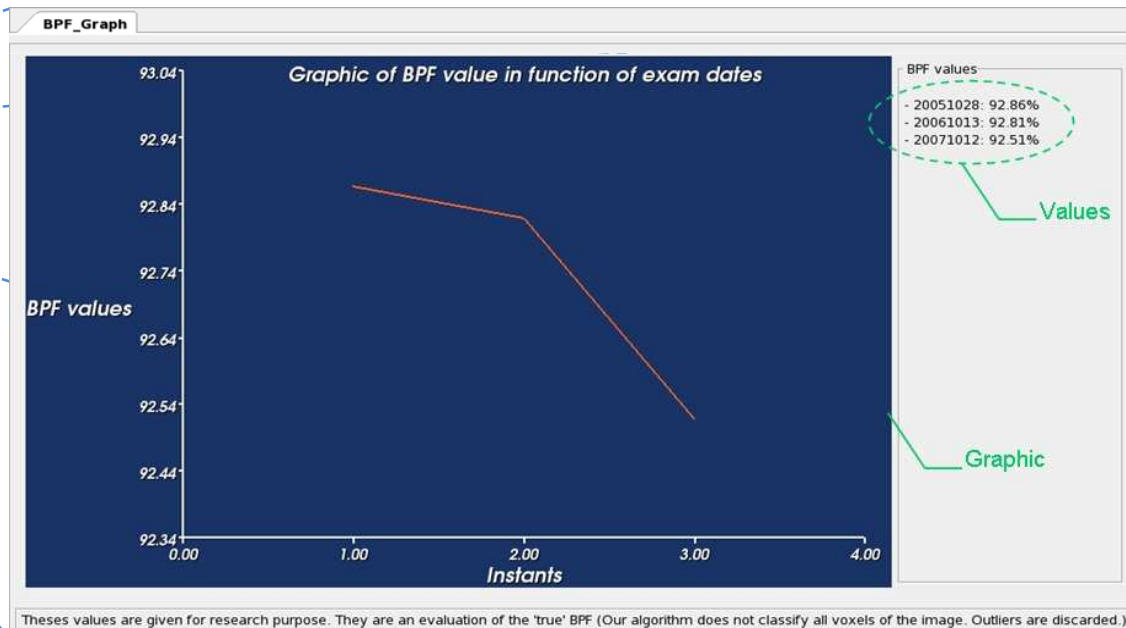
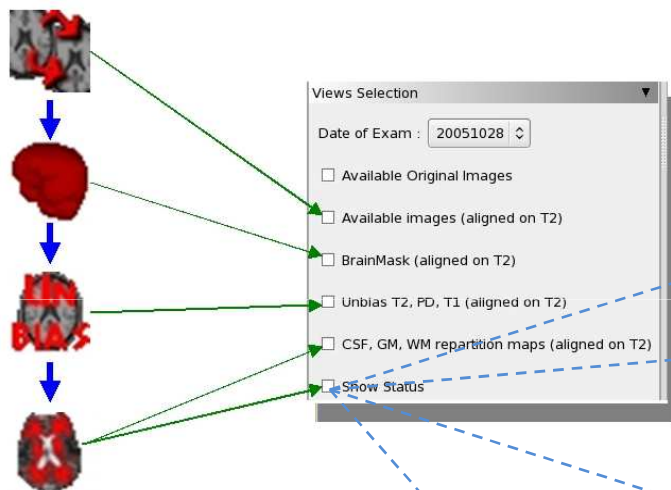
5- Performances

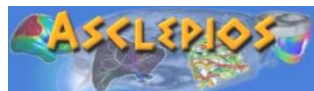
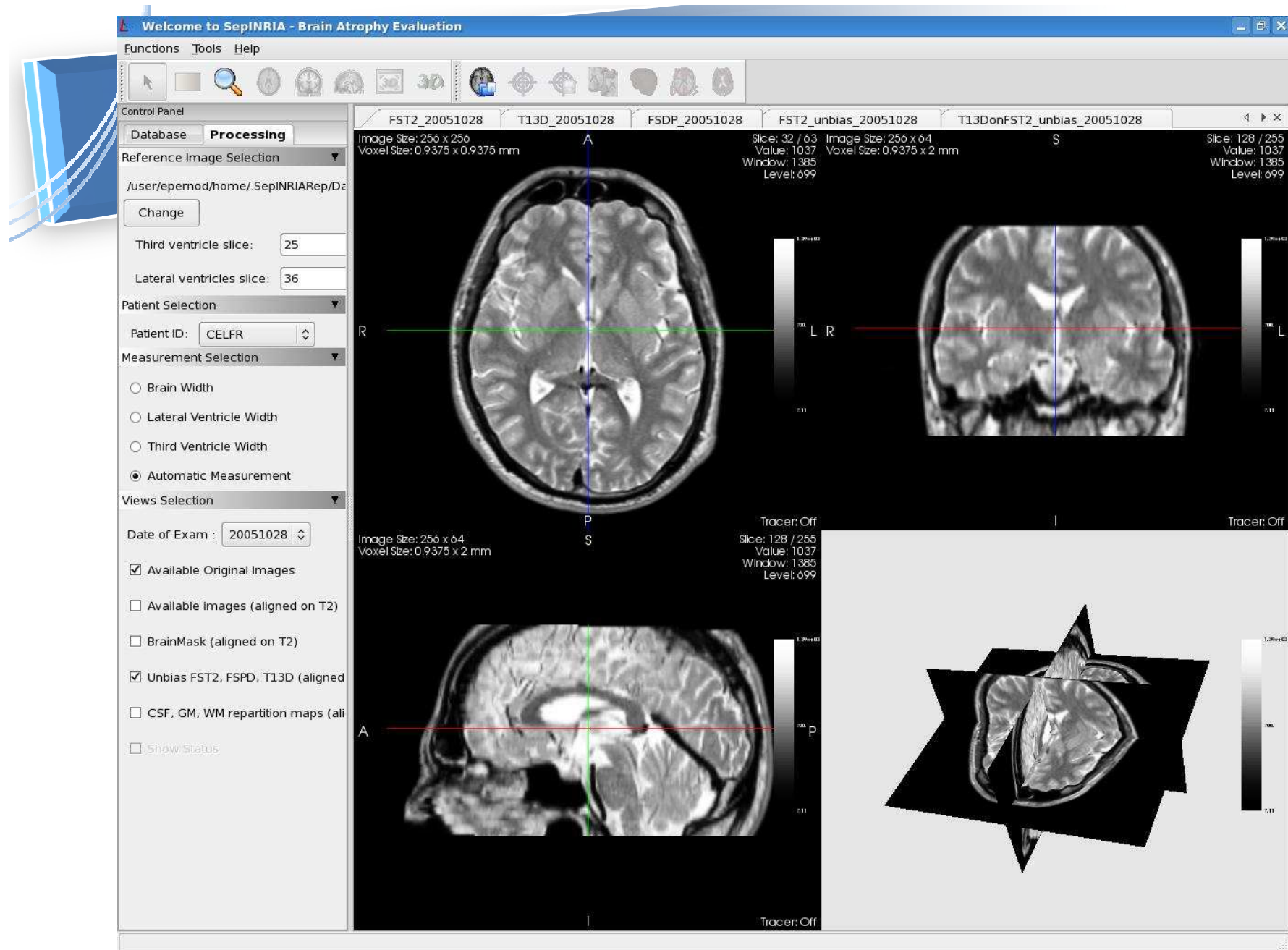
6- Conclusion

## Final step:

- WM, GM and CSF segmentations are used to compute the brain parenchymal fraction (BPF).
- The graph showing BPF value in function of the exam date reflects the evolution of atrophy

$$BPF = \frac{\text{Volume(GM)} + \text{Volume(WM)}}{\text{Volume(GM)} + \text{Volume(WM)} + \text{Volume(CSF)}}$$





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## **Problematic:** How to deploy and parallelize an algorithm on a grid?

- Needed transformations of the pipeline ?
- How to create and execute a workflow ?
- Performances ?
- First, what is a grid?
  - Network of shared computing resources.
  - Different from a cluster.

# Workflow creation

1- Introduction

2- The pipeline

3- SepINRIA

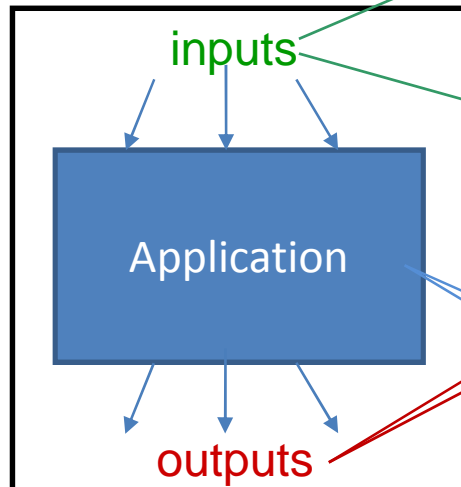
4- Workflow creation

5- Performances

6- Conclusion

## Services creation

- Splitting the pipeline in black boxes
- Description of each black box



- Use of GASW (Generic Application

```
<description>
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    <access type="LFN"/>
    <value value="baladin.sh"/>

    <input name="reference_image_inr" option="noa">
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    </input>

    <input name="float_image_inr" option="nob">
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    </input>

    <input name="baladin parameters" option="noc">
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    </input>

    <input name="Initial transformation_matrix" option="nod">
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    </input>

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    </output>

    <output name="trsf_matrix" option="nof">
      <template value="%s Reg.trsf"/>
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    </output>

    <sandbox name="baladin">
      <access type="LFN"/>
      <value value="baladin"/>
    </sandbox>

  </executable>
</description>
```

# Workflow creation

1- Introduction

2- The pipeline

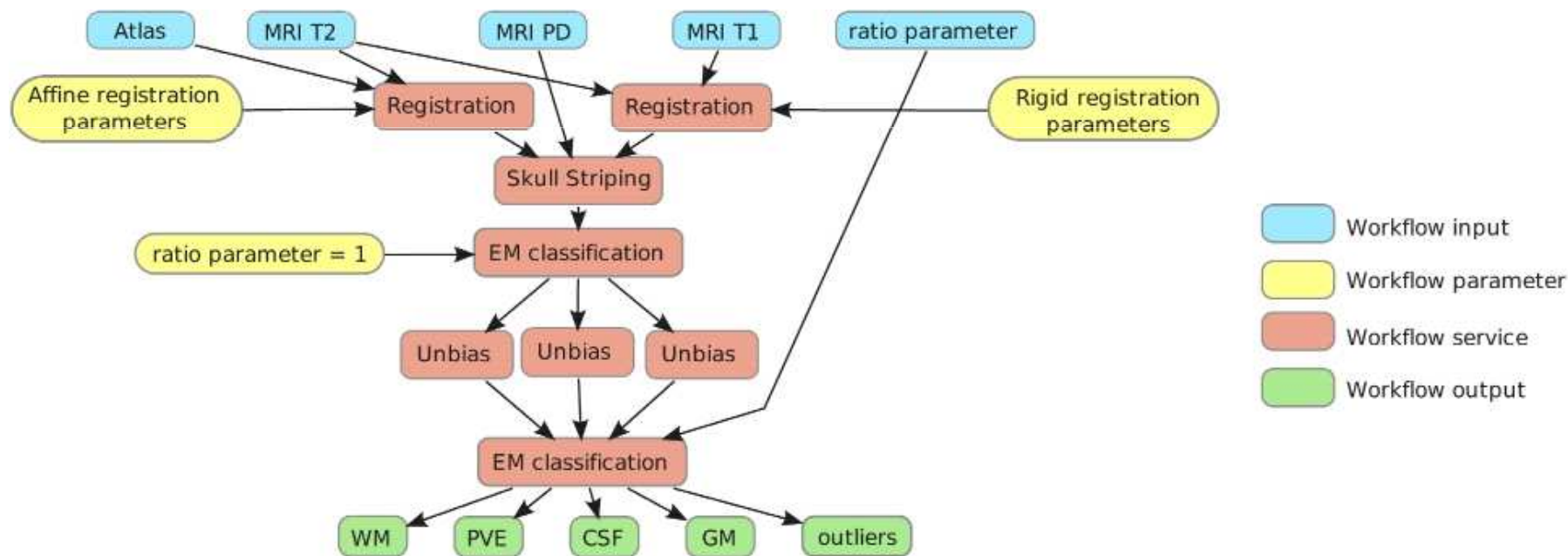
3- SepINRIA

4- Workflow creation

5- Performances

6- Conclusion

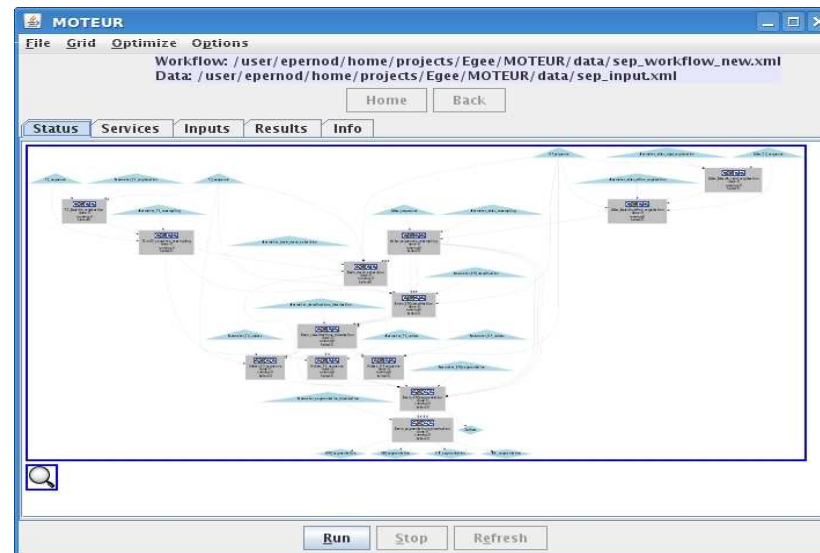
## Use of the software Taverna to build the structure of the workflow





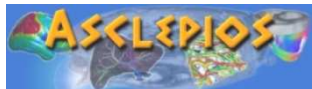
## Workflow execution

- Need a account on one computer of the grid
- Execution of the workflow (XML file) using MOTEUR



*Efficient services composition  
for grid-enabled data-intensive  
applications.  
T. Glatard, J. Montagnat, and X.  
Pennec. HPDC'06*

- MOTEUR takes in charge all the interactions with the grid



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# Performances

1- Introduction

2- The pipeline

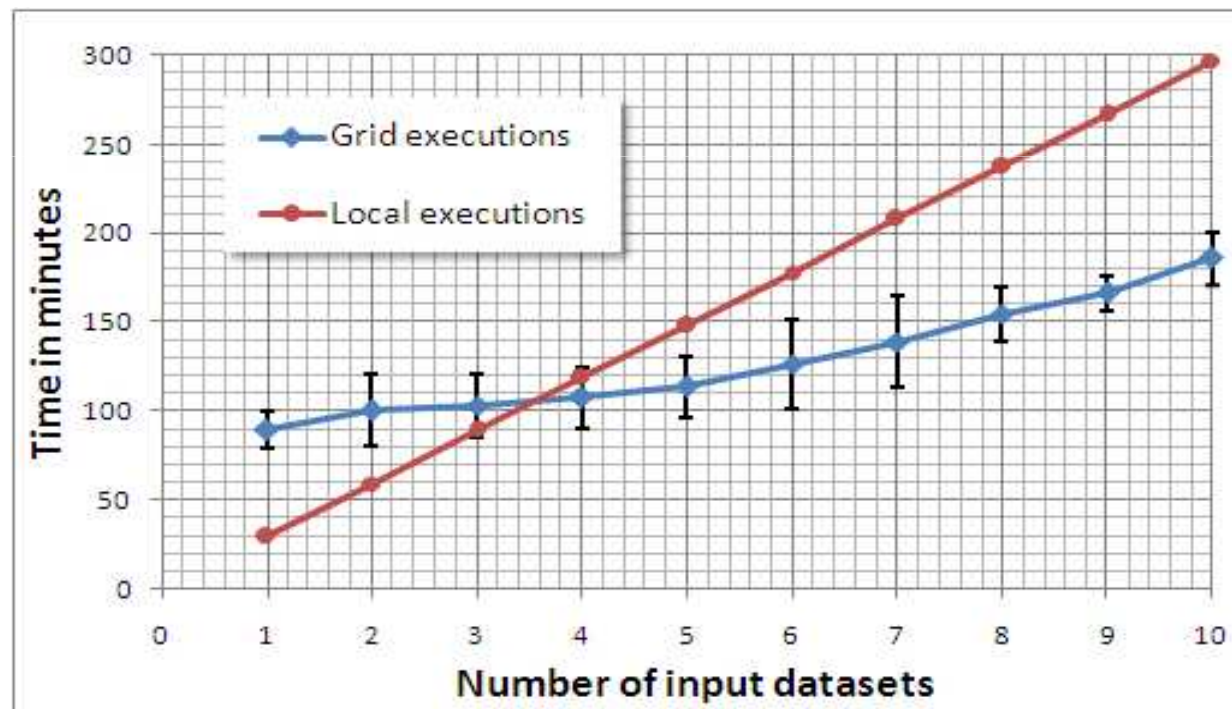
3- SepINRIA

4- Workflow creation

5- Performances

6- Conclusion

- Possibility of multiple concurrent executions
- Time performance:





## Potential issues

- The EGEE grid use the gLite middleware. <http://glite.web.cern.ch/glite/>
- In this framework, the Resource Broker is responsible for the matchmaking between job requests and resources.
  - Fastest responding resources are chosen. (after filtration)
  - Not necessarily the most powerful
  - Nor directly available.
- Workload management could becomes a bottleneck.



## Parameter-sweep test

- Validation of the deployment on the grid
  - Comparison with a sequential execution on one single computer: identical results
- The power of the grid allows to perform parameter sweeping in a reasonable amount of time
- Goal: To find a good compromise between accuracy and speed in the EM method. Study of the performance for different percentage of points used to estimate brain classes.

# Performances

1- Introduction 2- The pipeline 3- SepINRIA 4- Workflow creation

## Ratio Parameter of the EM

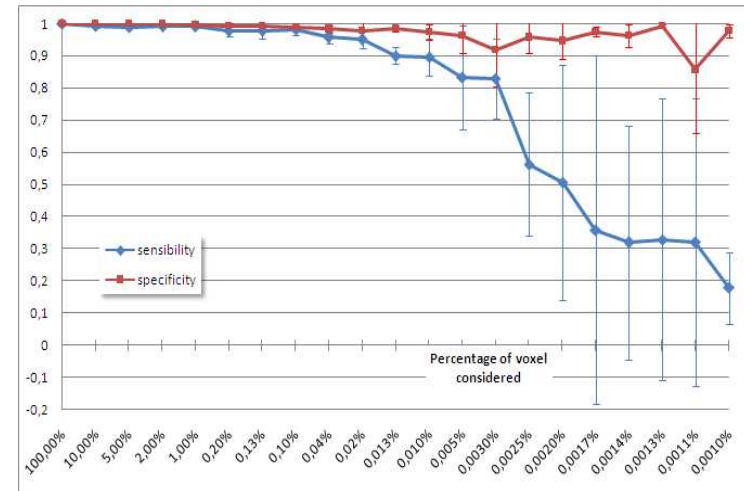
- Generation of WM segmentations obtained using different ratio value in the EM method.
- Comparaison of these segmentation to the segmentation of reference (ratio = 1)

$$\text{Percentage of voxel considered} = 100 \times \frac{1}{\text{ratio parameter}}$$

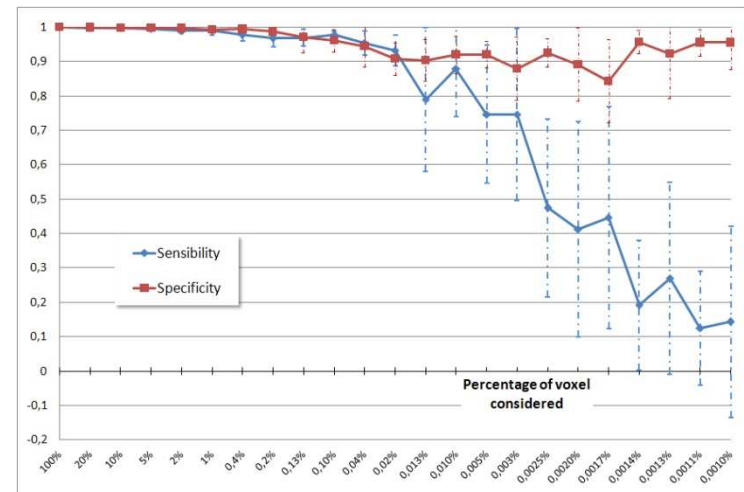
$$\text{Sensibility} = \frac{\text{true positives}}{\text{true positives} + \text{false negatives}}$$

$$\text{Specificity} = \frac{\text{true negatives}}{\text{true negatives} + \text{false positives}}$$

5- Performances 6- Conclusion

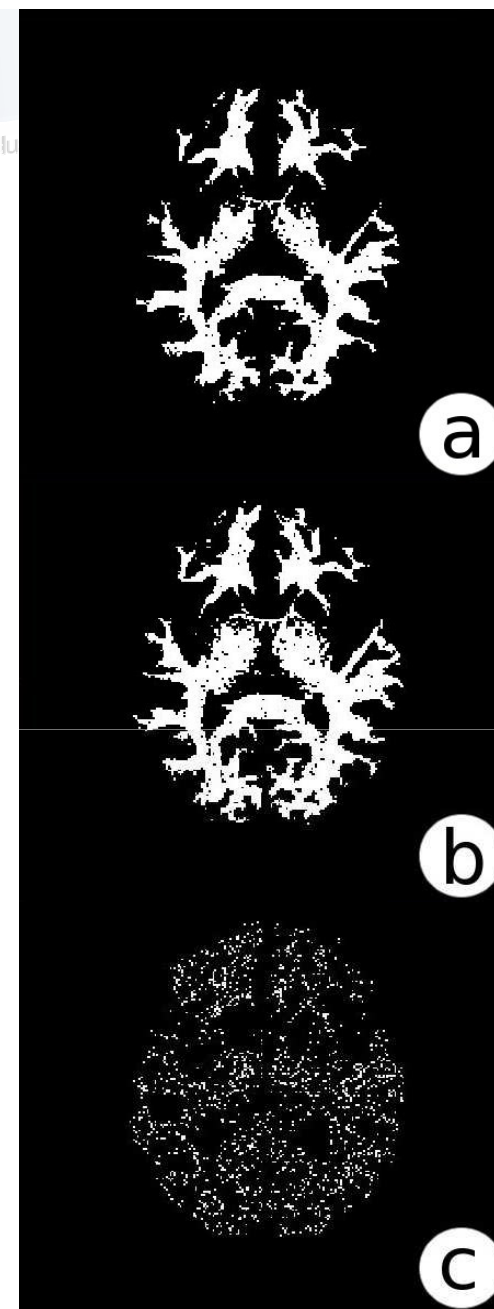
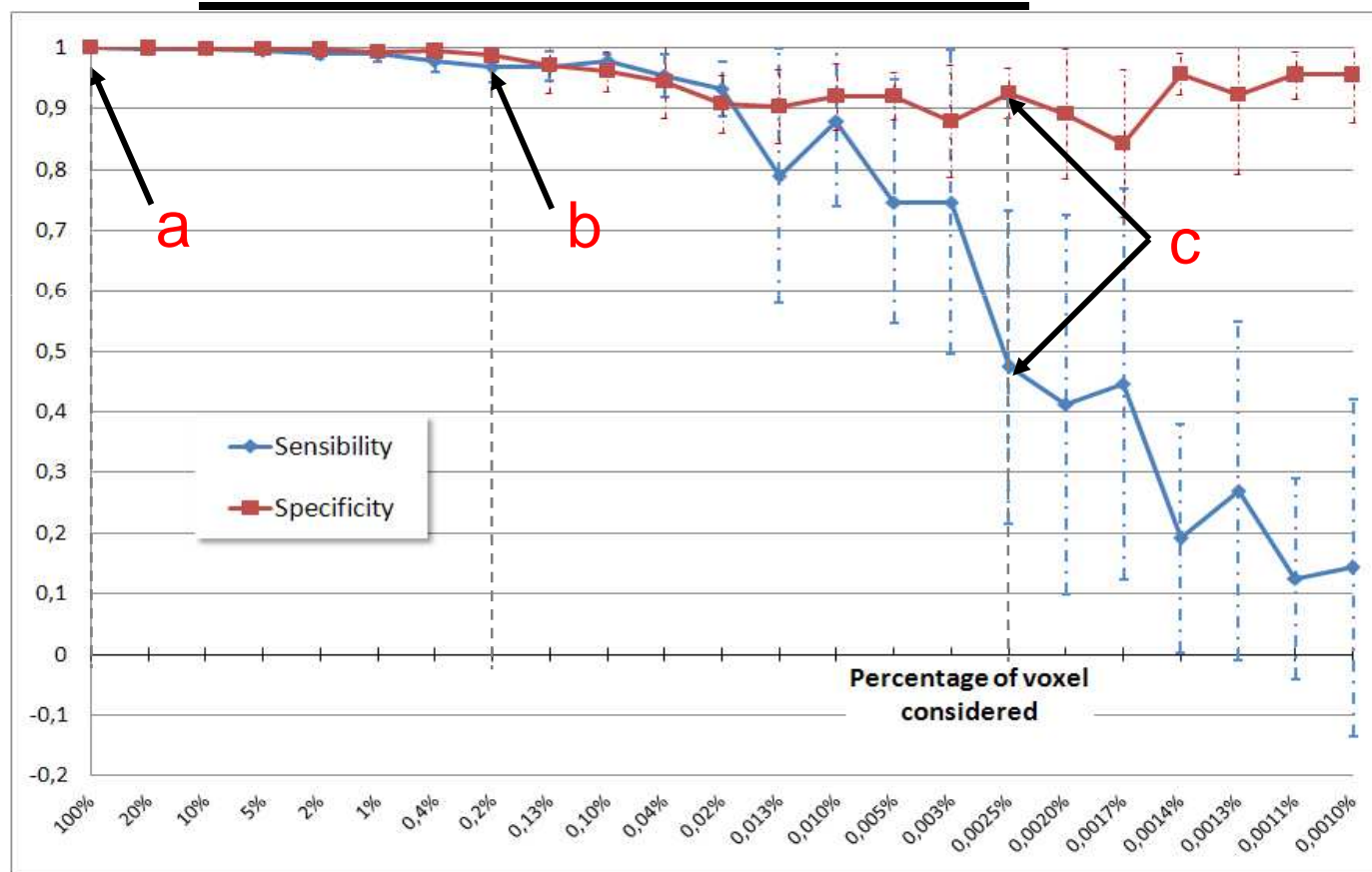


MR image of a control subject



MR image of a MS patient

## Ratio Parameter of the EM





## Ratio Parameter of the EM

- Using only 1% of the image voxels in the EM method:
  - Divides the execution time of the method by ~3
  - Still provides segmentation of sufficient quality
- Taking less than 1% of the voxels may leads to poor results
- ~ 210 workflow executions (10 per ratio value) have been computed (per image set).
  - Local execution time (sequential): ~ 100 hours (estimated)
  - Grid execution time: ~ 40 hours (4 hours per bunch of 21 workflow executions)

# Conclusion

1- Introduction 2- The pipeline 3- SepINRIA 4- Workflow creation 5- Performances 6- Conclusion

## Conclusion:

- Implemented tools in the software SepINRIA have been validated and recognized useful for clinician research and application (article accepted) :

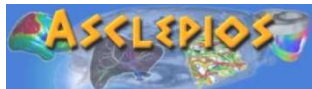
*Erik Pernod, Jean-Christophe Souplet, Mikael Cohen, Nicolas Toussaint, Christine Lebrun et Grégoire Malandain.  
SepINRIA v1.7.2: Multiple Sclerosis Brain MRI: visualisation, comparison and analysis Software. In World Congress  
for Treatment and Research in Multiple Sclerosis (WCTRIMS), Montreals, Canada, September 2008.*

- Deployment demonstration of a “real” medical image processing solution on the grid:
  - The power of the grid allows multiple concurrent executions and a sizeable gain of time.
  - As a consequence, it allows computation costly tests, e.g. parameter sweeping.
  - This implementation on the grid leads to the article:

*Erik Pernod, Jean-Christophe Souplet, Javier Rojas Balderrama, Diane Lingrand et Xavier Pennec.  
Multiple Sclerosis Brain MRI Segmentation Workow deployment on the EGEE Grid.  
In MICCAI-Grid Workshop (MICCAI-Grid), New York, NY, USA, September 2008*

## Future work:

- Generalization of the services to support more image formats
  - Does not require to modify the workflow nor the web service descriptions
  - Can be done directly at the application level
  - Allows the workflow diffusion to other research groups
- Add new services to the workflow to get the lesions segmentation.



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