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# MUSE Software Manual

*Release v3.0 (r352)*

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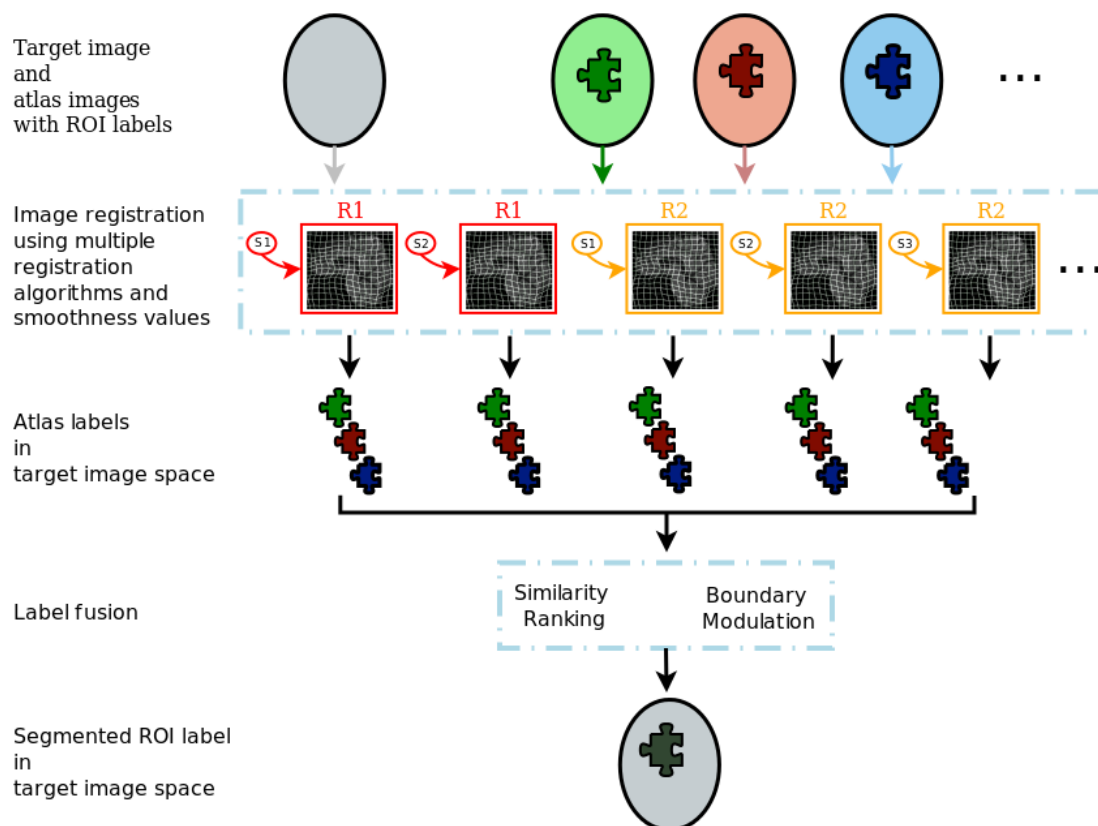
## Contents

<b>1</b>	<b>About the Algorithm</b>	<b>2</b>
<b>2</b>	<b>Download</b>	<b>2</b>
2.1	Software License . . . . .	2
2.2	Documentation . . . . .	2
2.3	System Requirements . . . . .	3
2.4	Register for Download . . . . .	3
<b>3</b>	<b>Installation</b>	<b>3</b>
3.1	Prerequisites . . . . .	3
3.2	Job Scheduler . . . . .	3
3.3	Configure . . . . .	3
3.4	Build . . . . .	4
3.5	Test (Optional) . . . . .	4
3.6	Install . . . . .	4
<b>4</b>	<b>Manual</b>	<b>5</b>
4.1	MUSE Default Command . . . . .	5
4.2	MUSE Options . . . . .	5
4.3	Calculating ROI Volumes generated from MUSE . . . . .	6
<b>5</b>	<b>Publications</b>	<b>6</b>
<b>6</b>	<b>People</b>	<b>6</b>
6.1	Advisors . . . . .	6
6.2	Software Development . . . . .	6
6.3	Contributors . . . . .	6
6.4	Testers . . . . .	7
	<b>Bibliography</b>	<b>7</b>

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# 1 About the Algorithm

MUSE generates a large ensemble of candidate labels in the target image space using multiple atlases, registration algorithms and smoothness values for these algorithms. The ensemble is then fused into a final segmentation. An illustration of the MUSE algorithm is given in the following figure.



## 2 Download

### 2.1 Software License

The MUSE software is freely available under a BSD-style open source license that is compatible with the Open Source Definition by [The Open Source Initiative](#) and contains no restrictions on use of the software. The full [license](#) text is included with the distribution package and available online.

### 2.2 Documentation

[MUSE Manual](#): Online version of this manual.

[MUSE ChangeLog](#): Summary of changes, new features, and bug fixes.

## 2.3 System Requirements

**Operating System:** Linux

## 2.4 Register for Download

Please [register online](#) to receive an email with the download links of the software.

# 3 Installation

See the [BASIS guide on software installation](#) for a complete list of build tools and detailed installation instructions.

## 3.1 Prerequisites

Dependency	Version	Description
<a href="#">BASIS</a>	2.1.0	A meta-project developed at <a href="#">SBIA</a> to standardize the software development.
<a href="#">DRAMMS</a>	1.4.1	A registration algorithm developed at <a href="#">SBIA</a> to warp images.
<a href="#">ANTS</a>	1.9.x	A registration algorithm developed at <a href="#">PICSL</a> to warp images.
<a href="#">MICO</a>	1.0.0	A segmentation algorithm developed at <a href="#">SBIA</a> to segment images.
<a href="#">AFNI</a>		Using the version built on 2008_07_18_1710
<a href="#">FSL</a>	4.1.5	A comprehensive library of analysis tools for brain imaging data
<a href="#">NIBABEL</a>	1.2.0	A python package for read and write access to common medical file formats
<a href="#">NUMPY</a>	1.6.1	A python package for scientific computing

\* The versions listed are the minimum versions of the softwares for which the MUSE package was tested.

## 3.2 Job Scheduler

If you have access to a computing cluster which has a job scheduler/queuing software (SGE, PBS etc) installed, it can be used to significantly reduce the (wall-clock) time it will take for the MUSE software to produce the results. During the installation process, you can initialize the `SCHEDULER` variable with the particular version of your job scheduler. Currently, there are four options that are supported. You can select the one that best fits your system:

```
SGE - Sun Grid Engine
PBS - Portable Batch System
NONE - No queuing system (default)
MISC - User defined setting
```

If you have a different queuing software and you select the “MISC” option, you need to modify the `src/schedulerSettings/SettingsMISC.sh` file within the package with the appropriate options and arguments that are specific to your queuing system. You can refer to the corresponding files for SGE and PBS as examples.

## 3.3 Configure

1. Extract source files:

```
tar -xzf muse-3.0.0-source.tar.gz
```

2. Create build directory:

```
mkdir muse-3.0.0-build
```

3. Change to build directory:

```
cd muse-3.0.0-build
```

4. Run **CMake** to configure the build tree by using one of the following commands:

```
cmake -D CMAKE_INSTALL_PREFIX:STRING=/Full/path/to/install/muse/  
-D SCHEDULER:STRING=???  
../muse-3.0.0-source
```

## 3.4 Build

After the configuration of the build tree, the software can be build using **GNU Make**:

```
make
```

## 3.5 Test (Optional)

After the build of the software, optionally run the tests using the command:

```
make test
```

Allow 30-60 mins for the test to finish. In case of a test failure, re-run the test, but this time by executing **CTest** directly with the `-V` option to enable verbose output and redirect the output to a text file:

```
ctest -V >& muse-test.log
```

and attach the file `muse-test.log` to the issue report.

## 3.6 Install

The final installation copies the built files and additional data and documentation files to the installation directory specified using the `CMAKE_INSTALL_PREFIX` option during the configuration of the build tree:

```
make install
```

After the successful installation, the build directory can be removed again.

## 4 Manual

### 4.1 MUSE Default Command

The main command of MUSE which labels an input image into a set of desired regions of interest is named `muse`. The simplest use is:

```
muse -i /path/to/source/sourceimage.hdr
```

This command will internally submit 22 registrations ( 11 DRAMMS + 11 ANTS ) and 1 labelFusion job. Once finished, it'll generate the output file in the `/Path/To/Source/Directory/` directory:

*Input\_n3\_str\_muse.nii.gz* - The final labeled image

**Supported File Formats:** [NIfTI-1](#) (recommended)

**Supported Datatypes:** byte (unsigned char, uint8), int8, short, int16, uint16, float, float32, int32.

### 4.2 MUSE Options

To parcellate the input brain image using the default options, but without using the computing cluster:

```
muse
-i /Path/To/Source/Directory/Input_n3_str.nii.gz
-Q;
```

To parcellate the input brain image using the default options, but without the cerebellum and storing the results at a user-specified destination :

```
muse
-i /Path/To/Source/Directory/Input_n3_str.nii.gz
-D /Path/To/Destination/Directory/
-C;
```

To parcellate the input brain image, but use the WMLS mask to exclude those regions from intensity weighting/correction:

```
muse
--in /Path/To/Source/Directory/Input_n3_str.nii.gz
--dest /Path/To/Destination/Directory/
--WML /Path/To/WMLS/Directory/Input_n3_str_WMLS_mask.nii.gz;
```

To parcellate the input brain image into ROIs using the default options. Additionally, use 6 CPU cores during processing of individual ROIs to speed up the process:

```
muse
-i /Path/To/Source/Directory/Input_n3_str.nii.gz
-D /Path/To/Destination/Directory/
-P 6;
```

To parcellate the input brain image into ROIs using Majority Voting to combine the results of different registrations:

```
muse
--in /Path/To/Source/Directory/Input_n3_str.nii.gz
--dest /Path/To/Destination/Directory/
--noIC
--noSim
--noFuzzy;
```

To parcellate the input brain image into ROIs and keep some of the important intermediate results:

```
muse
-i /Path/To/Source/Directory/Input_n3_str.nii.gz
-D /Path/To/Destination/Directory/
-k 1;
```

## 4.3 Calculating ROI Volumes generated from MUSE

To calculate the ROI volumes for each unique ROI within the labeled ROI image:

```
muse-calculateVolumes -i /Path/To/Destination/Directory/Input_n3_str_muse.nii.gz;
```

To calculate the ROI volumes for each unique ROI within the labeled ROI image and output the results in a csv file:

```
muse-calculateVolumes
-i /Path/To/Destination/Directory/Input_n3_str_muse.nii.gz
-s Input
-o /Path/To/Destination/Directory/Input_n3_str_muse.csv;
```

To calculate the derived ROI volumes (based on a hierarchical parcellation) from the labeled ROI image:

```
muse-calculateVolumes
-i /Path/To/Destination/Directory/Input_n3_str_muse.nii.gz
-s Input
-o /Path/To/Destination/Directory/Input_n3_str_muse.csv
-d
-v /Path/To/Source/Directory/Input_n3_str.nii.gz;
```

## 5 Publications

[TEMP2015]

## 6 People

### 6.1 Advisors

- Christos Davatzikos ([Christos.Davatzikos@uphs.upenn.edu](mailto:Christos.Davatzikos@uphs.upenn.edu))

### 6.2 Software Development

- Jimit Doshi ([Jimit.Doshi@uphs.upenn.edu](mailto:Jimit.Doshi@uphs.upenn.edu))

### 6.3 Contributors

- Guray Erus
- Yangming Ou
- Meng-Kang Hsieh

## 6.4 Testers

- Harsha Battapady
- Meng-Kang Hsieh
- Xiao Da
- Martin Rozycki
- Irem Aselcioglu

## References

[TEMP2015] Jimit Doshi, Guray Erus, Yangming Ou, Susan M. Resnick, Ruben C. Gur, Raquel E. Gur, Theodore D. Satterthwaite, Susan Furth, Christos Davatzikos, for the Alzheimer's Neuroimaging Initiative. MUSE: MUlti-atlas region Segmentation utilizing Ensembles of registration algorithms and parameters, and locally optimal atlas selection (Under Revision)