

MS Lesion Segmentation

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Background

- ▶ Multiple sclerosis is a chronic disease of the central nervous system (brain, spinal chord, optic nerves)
- ▶ Symptoms can include blurred vision, loss of balance, poor coordination, slurred speech, tremors, numbness, extreme fatigue, problems with memory and concentration, paralysis, and blindness.
- ▶ Affects more than 2.3 million worldwide.
- ▶ Affects more women than men (approximately 2:1)

Background

- ▶ Multiple sclerosis lesions in the brain are areas of active inflammation, demyelination, or permanent tissue damage.
- ▶ MRI is well-suited for assessing lesion burden (volume and patterns) because lesions appear as hyperintensities on FLAIR, T2-w, and PD images and as hypointensities on T1-w images.
- ▶ Obtaining manual lesion segmentations is often resource intensive, so accurate and efficient methods for automatic segmentation are necessary for scalability and research progress.

Goals of this tutorial

- ▶ Apply OASIS (Sweeney et al. 2013), an automatic lesion segmentation model, to obtain predicted lesion probability maps.
- ▶ Compare the results using the default OASIS settings to those obtained after re-training the model using our data.

Loading Data

- Let's read in the images, brain masks, and manual lesion segmentations.

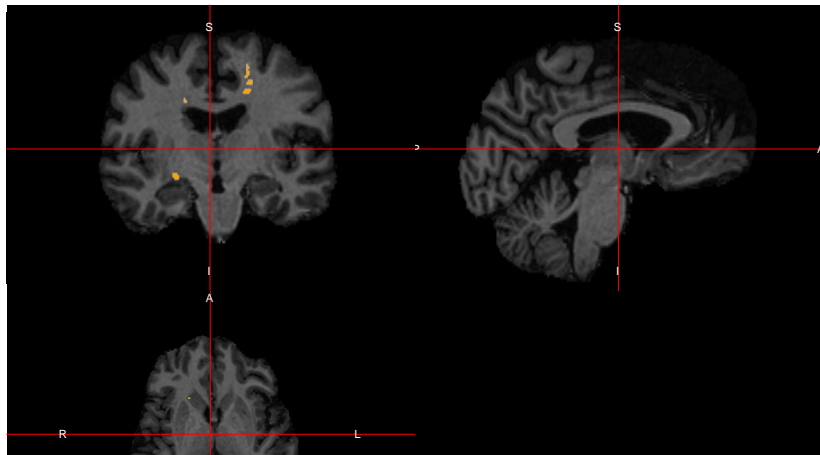
```
library(ms.lesion)
library(neurobase)
library(fslr)
library(scales)
library(oasis)

tr_files = get_image_filenames_list_by_subject(group = "tra
ts_files = get_image_filenames_list_by_subject(group = "tes
tr_t1s = lapply(tr_files, function(x) readnii(x["MPRAGE"]))
tr_t2s = lapply(tr_files, function(x) readnii(x["T2"]))
tr_flairs = lapply(tr_files, function(x) readnii(x["FLAIR"]))
tr_pds = lapply(tr_files, function(x) readnii(x["PD"]))
tr_masks = lapply(tr_files, function(x) readnii(x["Brain_Ma
tr_golds = lapply(tr_files, function(x) readnii(x["mask2"]))
ts_t1s = lapply(ts_files, function(x) readnii(x["MPRAGE"]))
ts_t2s = lapply(ts_files, function(x) readnii(x["T2"]))
```

Visualization

- Here's the T1 volume for training subject 05 with the 'gold standard' manual lesion segmentation overlaid.

```
les_mask = tr_golds$training05  
ortho2(tr_t1s$training05, les_mask, col.y = "orange")
```



MS Lesion Segmentation with OASIS

- ▶ OASIS is Automated Statistical Inference for Segmentation (Sweeney et al. 2013)
- ▶ The OASIS algorithm takes FLAIR, T1, T2, and PD images from patients with multiple sclerosis (MS) and produces OASIS probability maps of MS lesion presence, which can be thresholded into a binary lesion segmentation.
- ▶ OASIS uses logistic regression of the labels on the images, smoothed versions of the images, and some interaction terms

Default OASIS Model

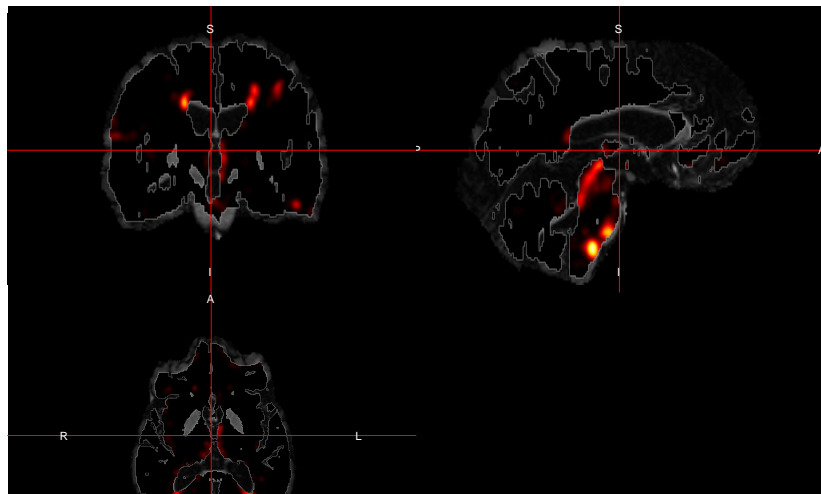
- ▶ The OASIS library comes with default parameters that can be used to generate probability maps for new test subjects
- ▶ Here we apply the default model to obtain OASIS probability maps for the test subjects.

```
default_predict_ts = function(x){  
  res = oasis_predict(  
    flair=ts_flairs[[x]], t1=ts_t1s[[x]],  
    t2=ts_t2s[[x]], pd=ts_pds[[x]],  
    brain_mask=ts_masks[[x]],  
    preproc=FALSE, normalize=TRUE,  
    model=oasis::oasis_model)  
  return(res)  
}  
default_probs_ts = lapply(1:3, default_predict_ts)
```


Vizualization

- Let's look at the probability map for test subject 01:

```
les_mask = default_ts[[1]]  
ortho2(ts_t1s$test01, les_mask)
```



Thresholding

- ▶ To get a final estimated segmentation, we must choose a cutoff to binarize the OASIS probability maps.
- ▶ The `binary` argument in the `oasis_predict` function is `FALSE` by default, resulting in the output being the probability map
- ▶ Setting `binary=TRUE` will return the thresholded version, using the input to the `threshold` argument (default = 0.16).
- ▶ In practice, we might want to use a grid search over thresholds and cross validation to choose the cutoff.

Default OASIS Model

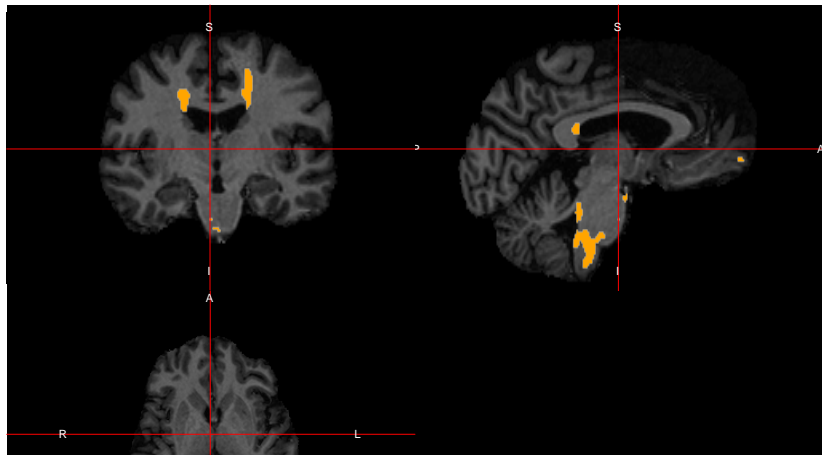
- ▶ To evaluate how the default model performs, we need to compare the predictions to a gold standard.
- ▶ Let's therefore obtain OASIS probability maps for our training subjects.
- ▶ We will use the default threshold to binarize.

```
default_predict_tr = function(x){  
  res = oasis_predict(  
    flair=tr_flairs[[x]], t1=tr_t1s[[x]],  
    t2=tr_t2s[[x]], pd=tr_pds[[x]],  
    brain_mask=tr_masks[[x]],  
    preproc=FALSE, normalize=TRUE,  
    model=oasis::oasis_model, binary=TRUE)  
  return(res)  
}  
default_probs_tr = lapply(1:5, default_predict_tr)
```

Default OASIS Model Results

- Here's the T1 volume for training subject 05 with the OASIS segmentation overlayed.

```
les_mask = default_tr[[5]]  
ortho2(tr_t1s$training05, les_mask, col.y = "orange")
```



Default OASIS Model Results

- ▶ Let's see how well OASIS corresponds with the gold standard manual segmentations for the training subjects.

```
tbls = lapply(1:5, function(x) table(c(tr_golds[[x]]), c(de  
lapply(tbls, function(x) (2*x[2,2])/(2*x[2,2] + x[1,2] + x
```

```
[[1]]
```

```
[1] 0.4938917
```

```
[[2]]
```

```
[1] 0.6860189
```

```
[[3]]
```

```
[1] 0.4316734
```

```
[[4]]
```

```
[1] 0.2891665
```

Improving Results

- ▶ We might improve the results by re-training the OASIS model using our five training subjects.
- ▶ To retrain the model using new data, binary masks of 'gold standard' lesion segmentations are needed and should be in T1 space.

Making OASIS data frames

- ▶ OASIS requires a particular data frame format
- ▶ OASIS has an option to preprocess your data for you (preproc)
- ▶ OASIS has an option to normalize the intensities of your data for you using a whole-brain normalization (normalize)
- ▶ `make_df()` below is a helper function

```
library(oasis)
make_df = function(x){
  res = oasis_train_dataframe(
    flair=tr_flairs[[x]], t1=tr_t1s[[x]],
    t2=tr_t2s[[x]], pd=tr_pds[[x]],
    gold_standard=tr_golds[[x]],
    brain_mask=tr_masks[[x]],
    preproc=FALSE, normalize=TRUE,
    return_preproc=FALSE)
  return(res$oasis_dataframe)
}
oasis_dfs = lapply(1:5, make_df)
```

Training OASIS

- ▶ The function `oasis_training` takes the data frames we made and fits a logistic regression, where the outcome vector consists of all subjects' voxel-level data (top 85% in intensity)
- ▶ The function `do.call` is a useful R function that applies the function named in the first argument to all elements of the list specified in the second argument.

```
model = do.call("oasis_training", oasis_dfs)
```


References

Sweeney, Elizabeth M, Russell T Shinohara, Navid Shiee, Farrah J Mateen, Avni A Chudgar, Jennifer L Cuzzocreo, Peter A Calabresi, Dzung L Pham, Daniel S Reich, and Ciprian M Crainiceanu. 2013. "OASIS Is Automated Statistical Inference for Segmentation, with Applications to Multiple Sclerosis Lesion Segmentation in MRI." *NeuroImage: Clinical* 2. Elsevier: 402–13.