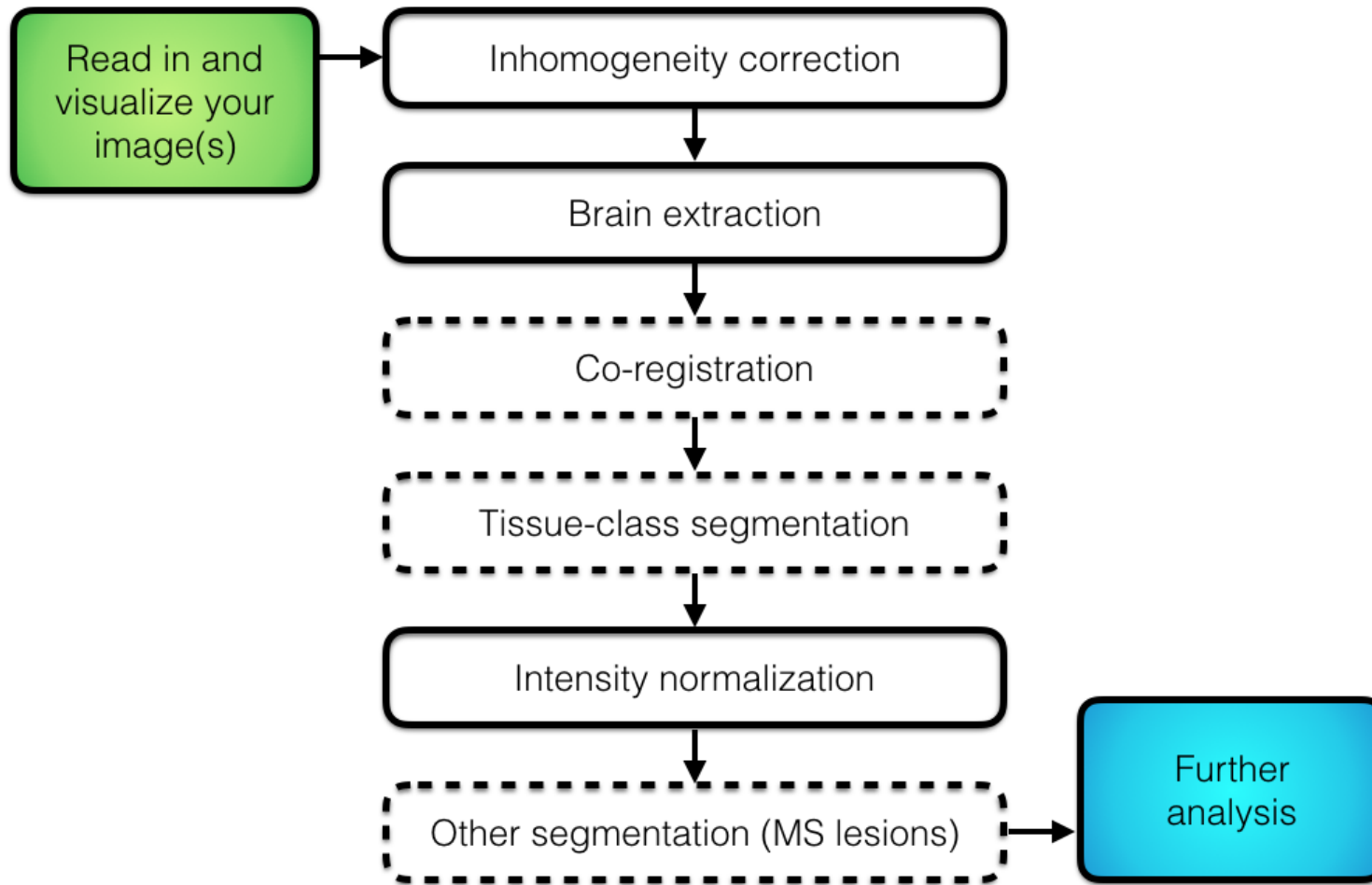


MS Lesion Segmentation

Overall Pipeline



Background

- Obtaining manual lesion segmentations is often resource intensive.
 - "Gold standard": Inter- and Intra-rater variability.
- Accurate and efficient methods for automatic segmentation are necessary for scalability and research progress.
- In this tutorial, we will learn how to train and apply OASIS (Sweeney et al. 2013), an automatic lesion segmentation model, to obtain predicted lesion probability maps.
 - Relies on intensity-normalized data.

MS Lesion Segmentation with OASIS

- **OASIS** is **A**utomated **S**tatistical **I**nference for **S**egmentation (Sweeney et al. 2013).
- OASIS takes FLAIR, T1, T2, and PD (optional) images.
 - Produces OASIS probability maps of MS lesion presence.
 - These can be thresholded into a binary lesion segmentation.
- The OASIS model is based on a logistic regression.
 - Regress binary manual segmentation labels on the images, smoothed versions of the images, and some interaction terms (e.g., supervised learning).
- Performs well compared to common machine learning models (Sweeney et al. 2014)

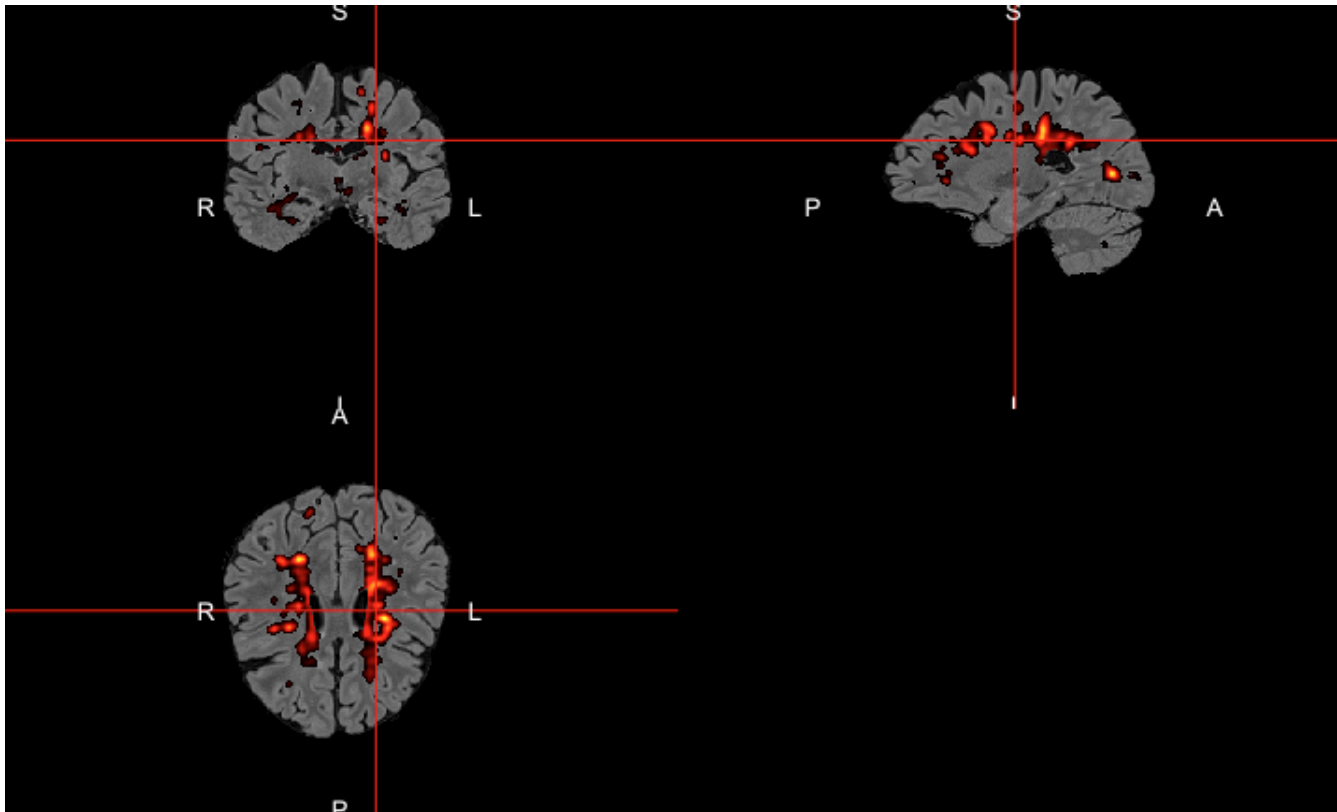
Default OASIS Model

- The OASIS library comes with default parameters that can be used to generate probability maps for new test subjects.
 - The default model was trained on approximately 100 MS subjects and 30 healthy subjects with manual segmentations.
- Here we apply `oasis_predict` with the default model to obtain OASIS probability maps for the test subjects.

```
library(oasis)
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 of probability map

- Here's the probability map for test subject 01.

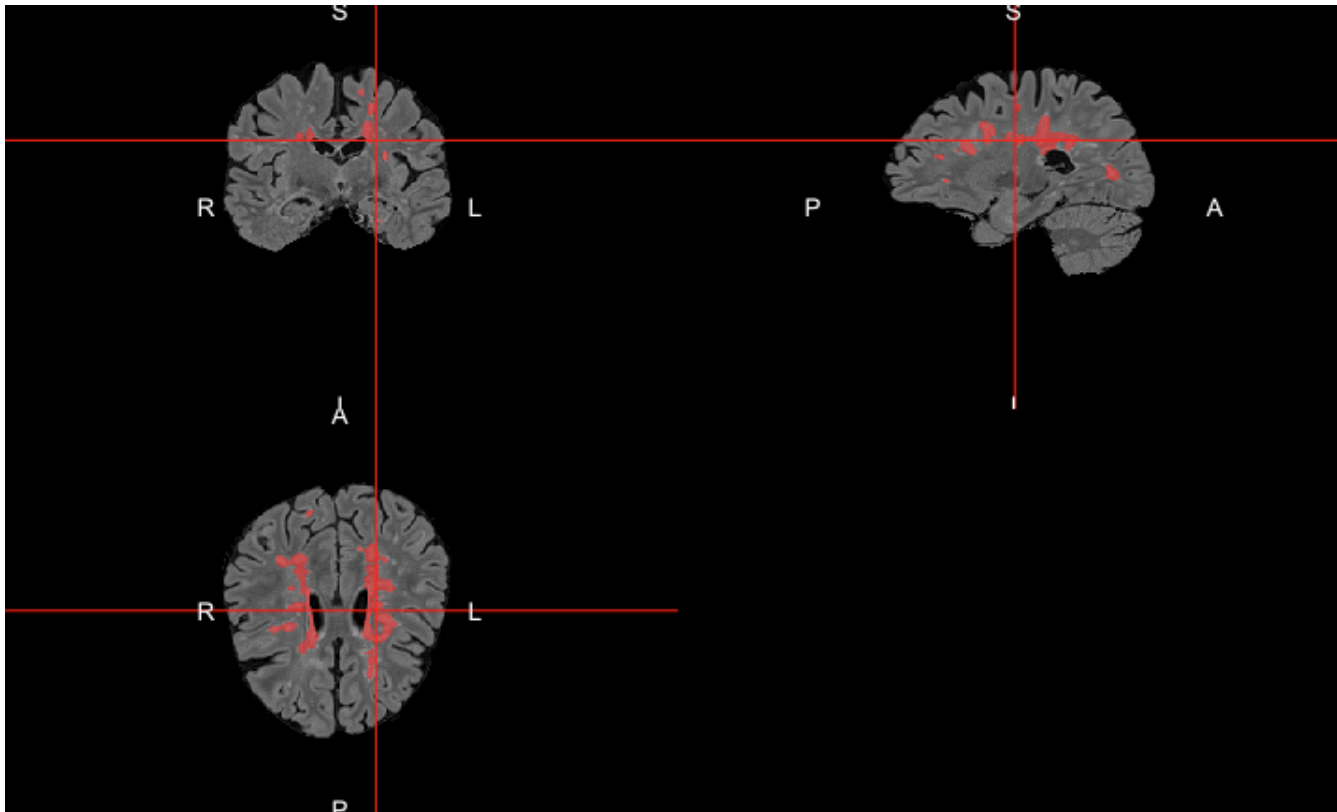


Thresholding: Getting a binary map

- 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).
 - 0.16 was obtained via a validation set allowing for a 0.5% false positive rate.
- In practice, we might want to use a grid search over thresholds and cross validation to choose the cutoff.

Vizualization of binary map

- Here's the binary mask for test subject 01, using the default 0.16 threshold:



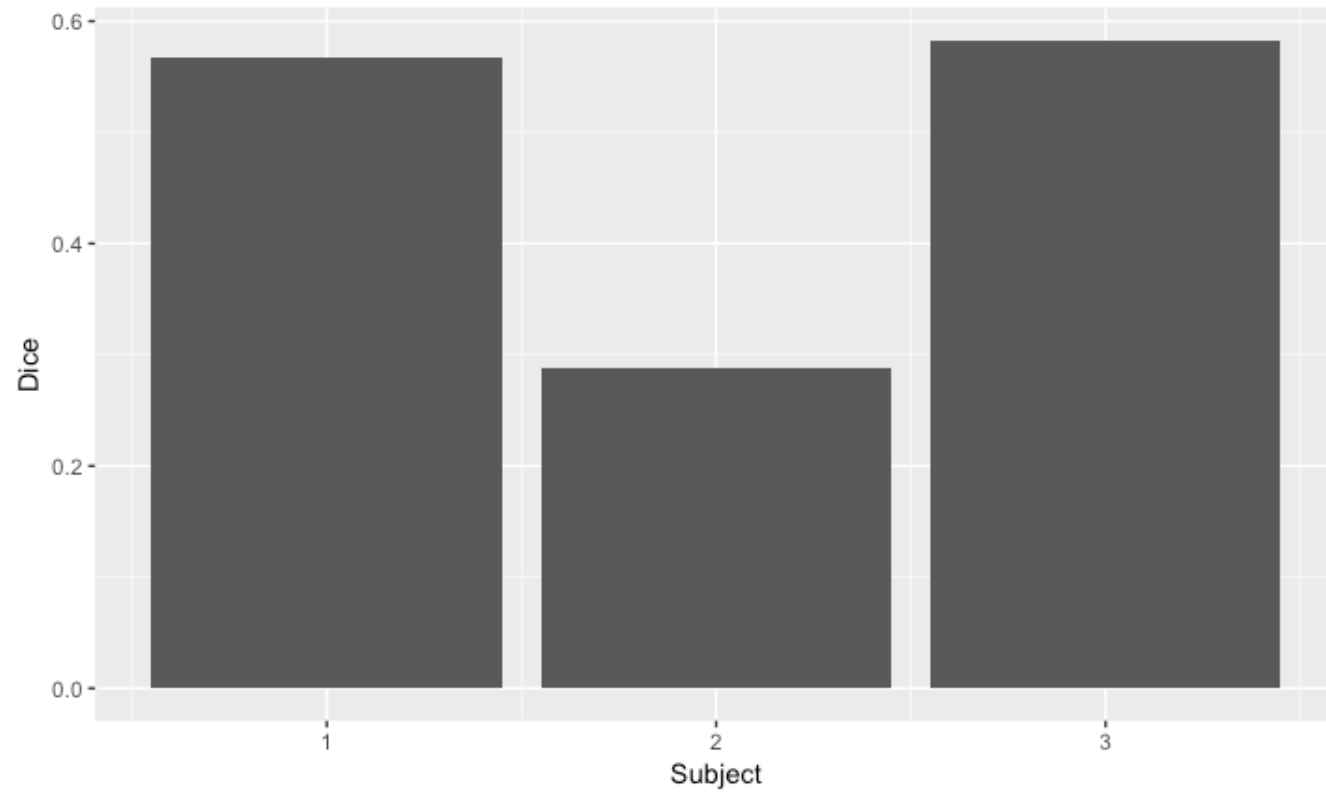
Default OASIS Model

- To evaluate how the default model with default threshold performs, we'll compare the predictions to our manual segmentations.
- Sorensen–Dice coefficient:
 - Similarity measure between two samples.
 - Ranges from 0 to 1.
 - (TP) - true positive, (FP) - false positive, (FN) - false negative.

$$D = \frac{2TP}{2TP + FP + FN}$$

Default OASIS Model Results

Dice coefficients for the test subjects



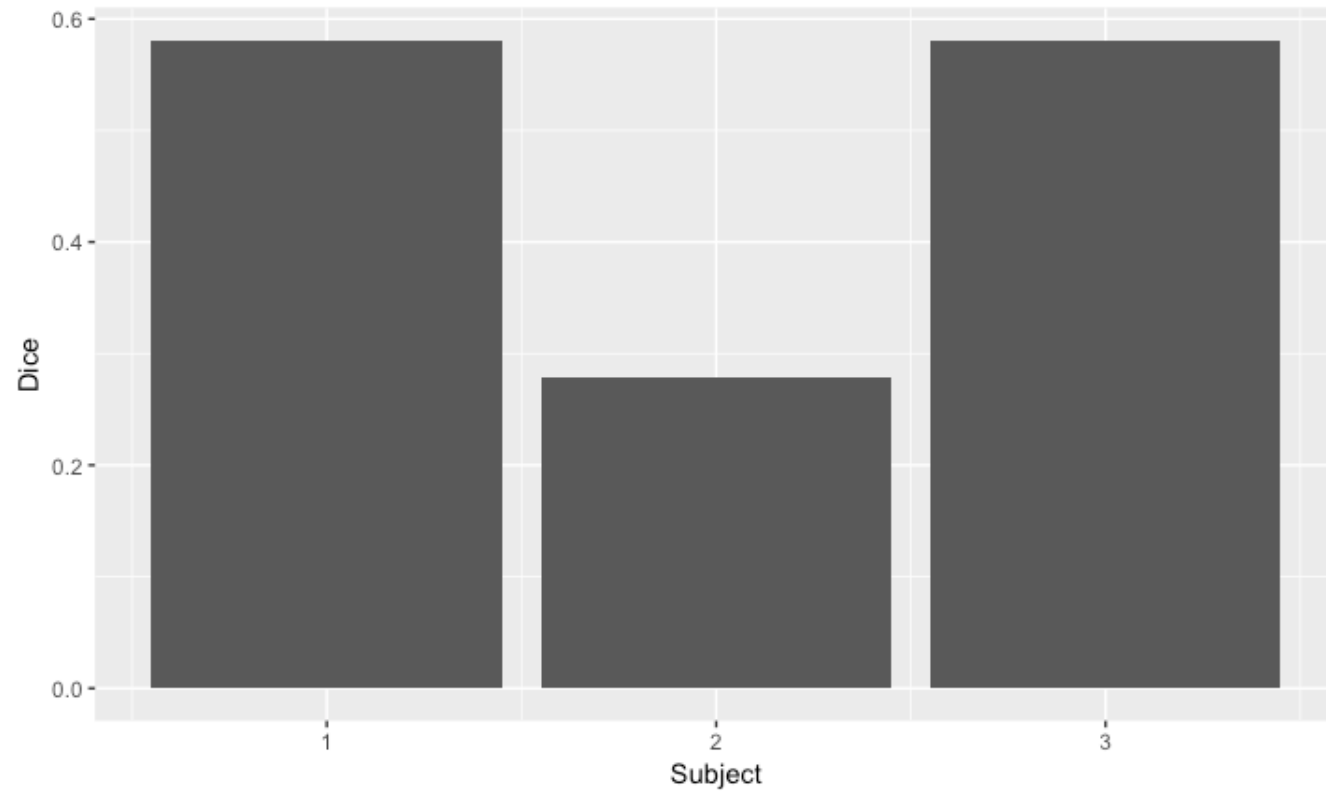
Improving Results

- We might be able to improve the results by adjusting the threshold.
- Let's optimize the threshold on the training data using a grid search (in practice, we might do cross-validation).

Threshold	0.050	0.100	0.150	0.200	0.250	0.300
Average Dice	0.242	0.272	0.273	0.261	0.231	0.194

Improving Results

- Turns out a coarse grid search chose a threshold of 0.15, so the results are nearly identical.



Improving Results

- We might be able to further improve the results by re-training the OASIS model using our five training subjects.
- To re-train 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, which we create using the function `oasis_train_dataframe`.
- Includes an option to preprocess your data (`preproc`), which does (1) inhomogeneity correction using `fsl_biascorrect` and (2) rigid coregistration using `flirt` to the T1 space.
- Includes an option to whole-brain intensity normalize (`normalize`).
- `make_df()` below is a helper function.

```
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 using labels and features from a subset of voxels in each subject's brain mask (top 15% in FLAIR 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.

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

OASIS model object

```
print(ms.lesion::ms_model)
```

```
Call: glm(formula = form, family = binomial, data = df)
```

Coefficients:

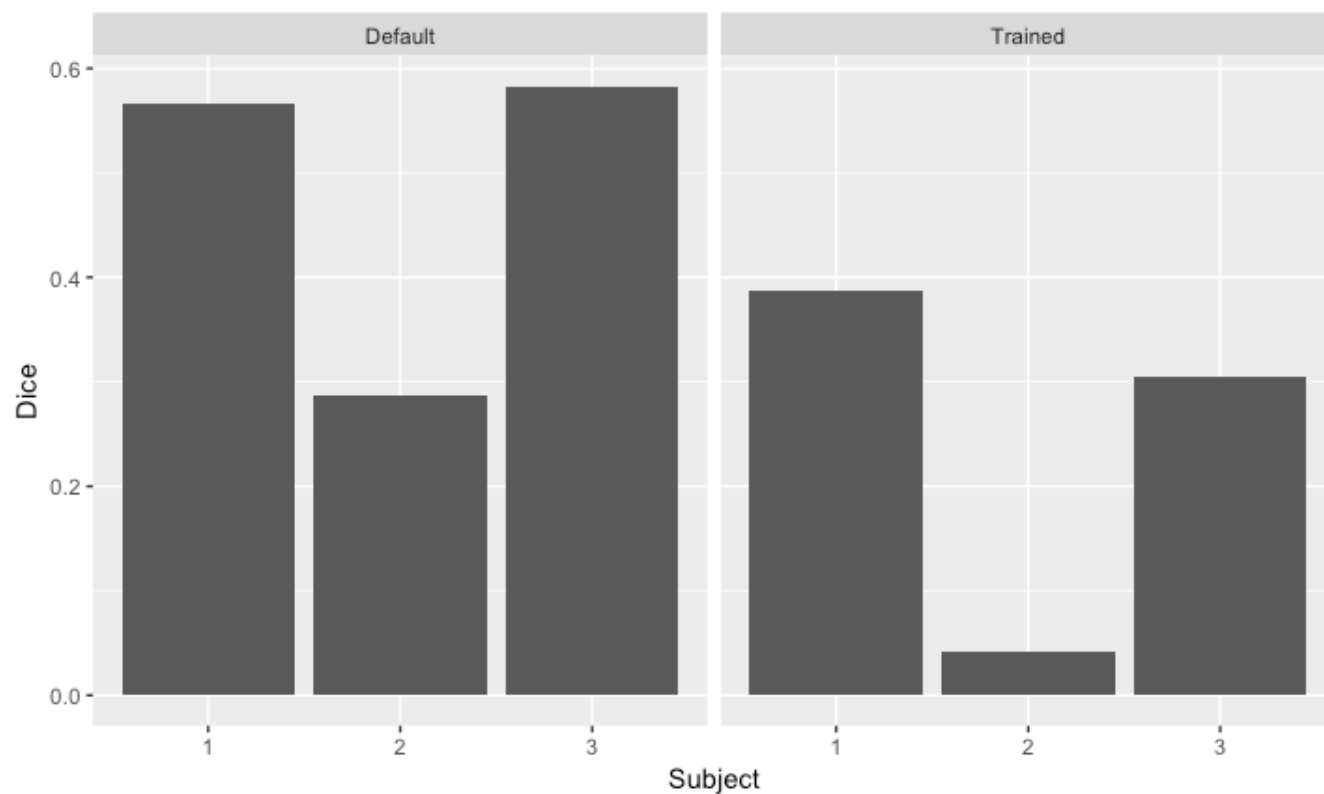
(Intercept)	FLAIR_10	FLAIR	FLAIR_20
-4.79369	13.10386	1.14120	-18.77010
T2_10	T2	T2_20	T1_10
4.85370	1.09444	-7.06750	13.63554
T1	T1_20	FLAIR_10:FLAIR	FLAIR:FLAIR_20
1.04771	-21.09848	-1.28891	1.03121
T2_10:T2	T2:T2_20	T1_10:T1	T1:T1_20
0.09151	3.18903	-1.04701	3.14265

Degrees of Freedom: 3930444 Total (i.e. Null); 3930429 Residual
Null Deviance: 2691000
Residual Deviance: 1842000 AIC: 1842000

Trained OASIS Model Results

```
Threshold    0.050 0.100 0.150 0.200 0.25 0.300
Average Dice 0.253 0.324 0.346 0.346 0.33 0.294
```

- Using a threshold of 0.15.
- Dice coefficients for default vs. re-trained OASIS model.



Improvement

- Percent improvement in Dice over the default model:

ID	Dice
01	-31.7
02	-85.7
03	-47.7

Website

http://johnmuschelli.com/imaging_in_r

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." 2. Elsevier: 402–13.

Sweeney, Elizabeth M, Joshua T Vogelstein, Jennifer L Cuzzocreo, Peter A Calabresi, Daniel S Reich, Ciprian M Crainiceanu, and Russell T Shinohara. 2014. "A Comparison of Supervised Machine Learning Algorithms and Feature Vectors for MS Lesion Segmentation Using Multimodal Structural MRI." 9 (4). Public Library of Science: e95753.