

# Lab session 9:

## Between-subjects multi-voxel pattern classification analysis (MVPA)

Andrew Bauer

03/23/16

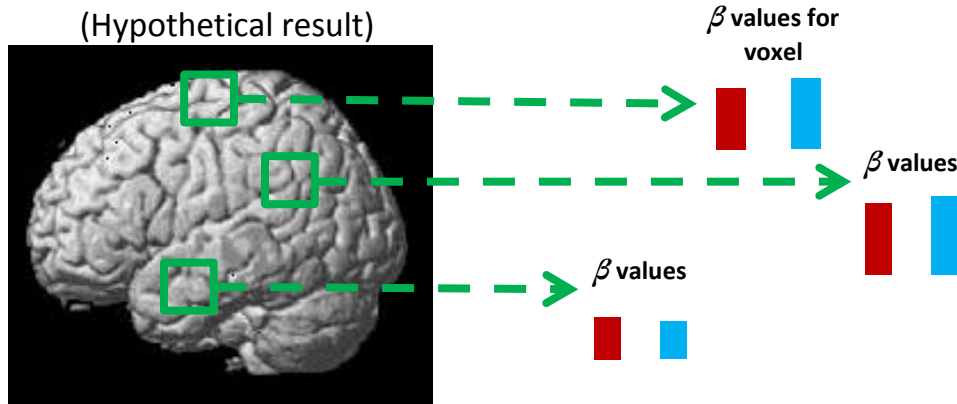
Session no.	Date (all Wednesday)	Topic/activity	Topic of quiz that day	Topic of lab write-up (assignment) due that day
1	13-Jan	Lab overview		
2	20-Jan	Brain anatomy		
3	27-Jan	Data preprocessing	Brain anatomy (no. 1)	
4	3-Feb	Set up GLM model	Functional brain anatomy (no. 2)	
5	10-Feb	Single-subject SPM contrasts	Data preprocessing and GLM model (no. 3)	Brain anatomy (no. 1)
6	17-Feb	Within-subject MVPA		Single-subject SPM contrasts (no. 2)
7	24-Feb	SIBR tour and review for mid-term exam		Within-subject MVPA (no. 3)
No lab	2-Mar	No lab (mid-term exam)		
No lab	9-Mar	No lab (spring break)		
8	16-Mar	Group-level SPM contrasts		
9	23-Mar	Between-subjects MVPA		Group-level SPM contrasts (no. 4)
10	30-Mar	Voxel-wise modeling		Between-subjects MVPA (no. 5)
11	6-Apr	Functional connectivity analysis (no assignment)		
12	13-Apr	Review for final exam		Voxel-wise modeling (no. 6)
No lab	20-Apr	No lab		
No lab	27-Apr	No lab (final exam)		

# Are there differences in activation between object concepts?

## GLM and MVPA approach this question differently

**Tools** vs. **other** object categories

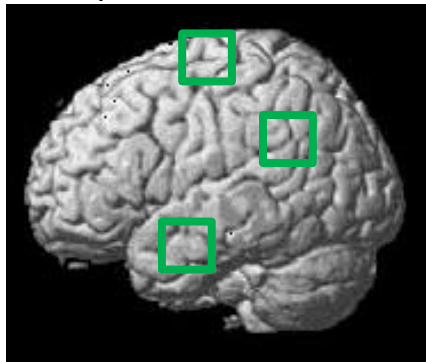
GLM  
contrast  
in each  
voxel



- GLM considers each single voxel *in isolation*
- Within a voxel, GLM compares conditions' activation *levels* ( $\beta$ )
- On the left: It appears **Tools** does not activate the brain differently, compared to **other** categories, in each voxel (see the three voxels)

Get response signal from each selected voxel (not to scale) for each object concept...

MVPA  
using  
pattern  
of 3  
voxels



Object concept	Voxel activation pattern (3 voxels)
hammer:	
saw:	
church:	
butterfly:	
.	.
.	.
.	.

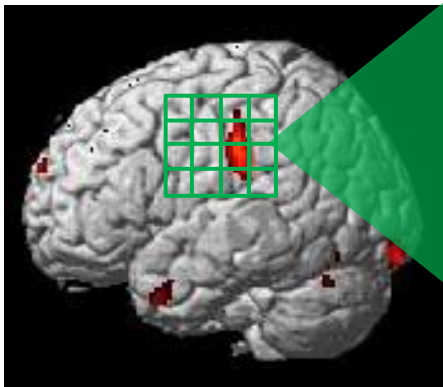
- MVPA considers *interactions* of voxels, and compares *patterns*; it uses the data to a fuller extent, and is thus more sensitive, than GLM
- On the left: Although GLM (above) shows no significant difference in activation *levels* in any single voxel for **Tools** vs. **others**, the voxel activation *patterns* between **Tools** and **other** concepts *are* different (i.e. are reliably distinguishable)

# Neural representation/storage of concept meaning

- MVPA asks: Are individual tool concepts represented in some brain areas?
- I.e. are there brain areas that encode rich information about different tool concepts?
  - GLM is not a sensitive enough method

**GLM contrast** (our actual result):  
Tools vs. other object categories

Say we select these 16 voxels  
(not to scale) for MVPA...



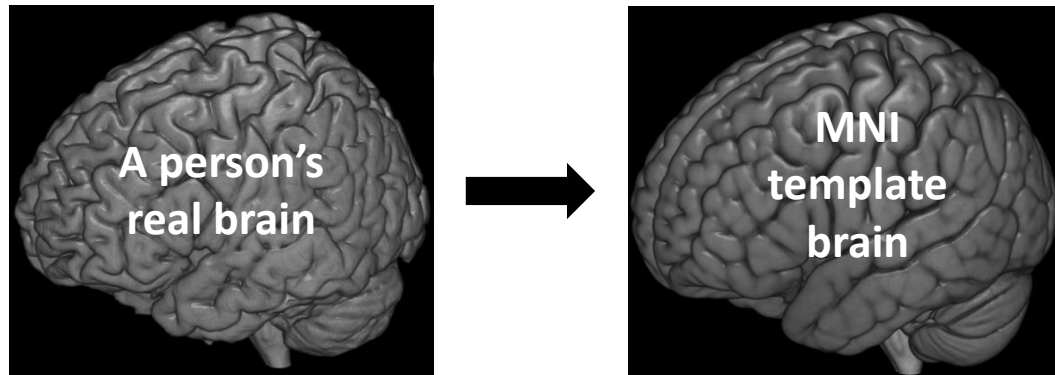
**Two possible MVPA results:**  
**Representation of tool concepts?**

	<i>No</i>		<i>Yes</i>
	Voxel pattern		Voxel pattern
hammer		OR	
drill			
.			
.			
.			
<tool <sub>i</sub> >			<tool <sub>i</sub> >

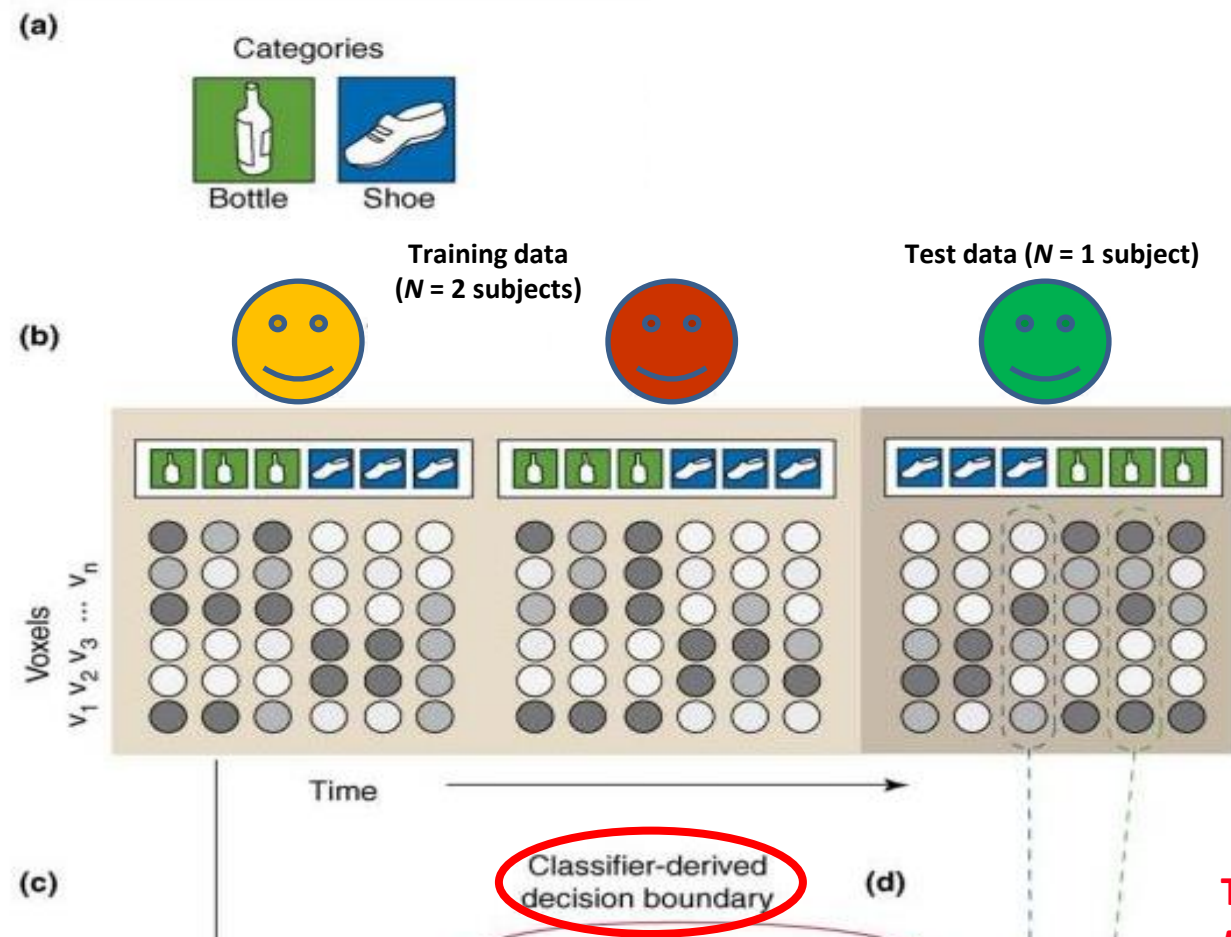
We can also use MVPA to ask:

Are neural representations similar between different people?

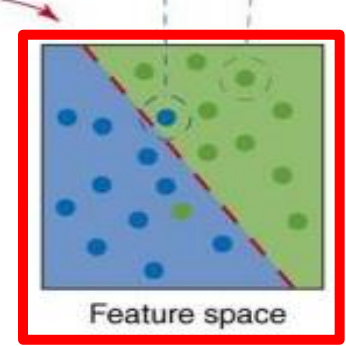
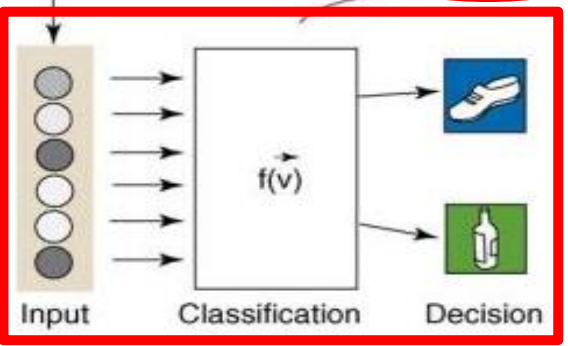
... where each person's brain is spatially normalized:



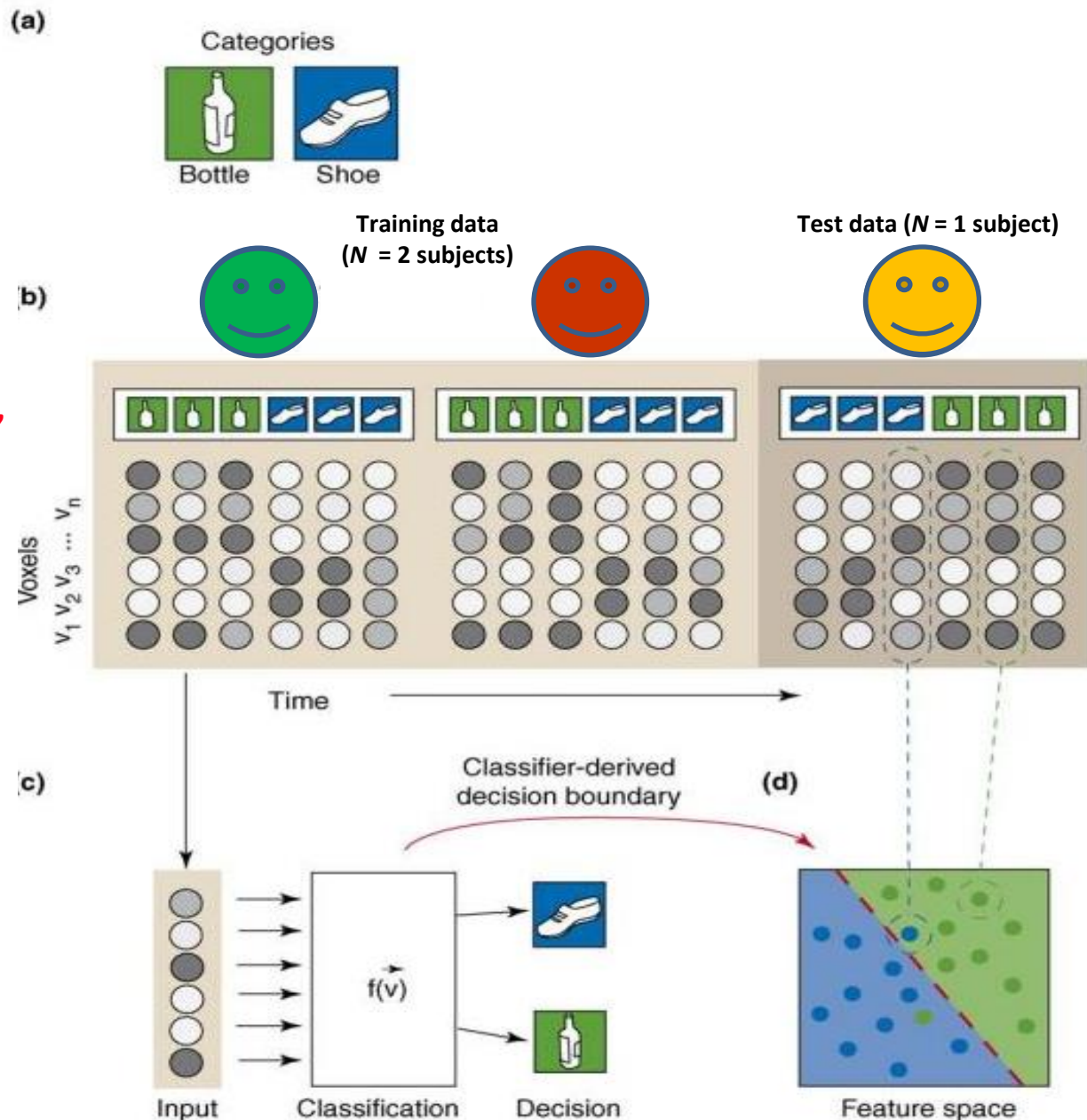
# How between-subjects MVPA works (assume $N = 3$ subjects)



The classifier is constructed from TRAINING data (from  $N = 2$  subjects); its *decision boundary* in feature (voxel) space is used to classify the TEST data



The TEST data (from  $N = 1$  subject) are compared to the *decision boundary* (feature space is simplified to 2D here; # dimensions = # voxels that make up the activation patterns)

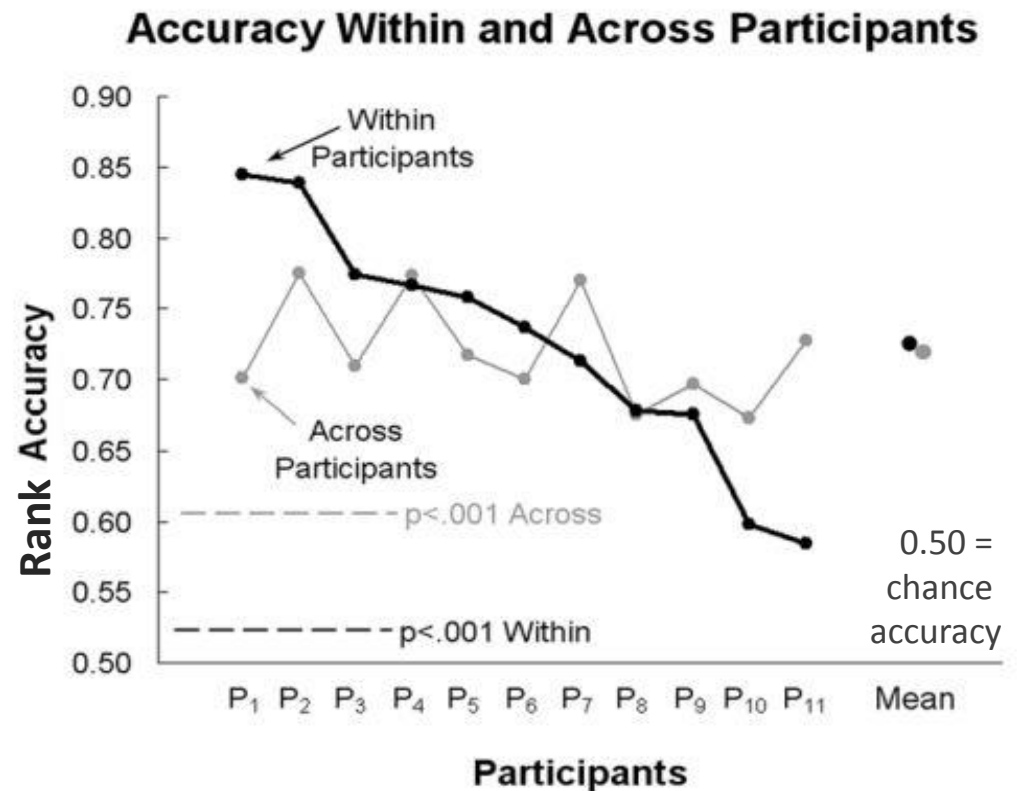
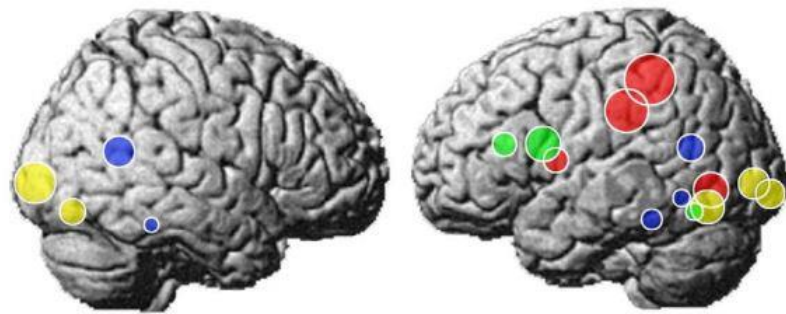


The process goes through many iterations, or “folds”, partitioning the data into TRAINING and TEST sets differently every time (each subject is used as the TEST data once)

The final output is the classification accuracy averaged over each subject (i.e. how well the subjects’ data were labeled)



We can classify each subject's data using a classifier trained on all the *other* subjects (gray line to the right)  
(Just et al., 2010's study on object concepts)



Category	Exemplar 1	Exemplar 2	Exemplar 3	Exemplar 4	Exemplar 5
body parts	leg	arm	eye	foot	hand
furniture	chair	table	bed	desk	dresser
vehicles	car	airplane	train	truck	bicycle
animals	horse	dog	bear	cow	cat
kitchen utensils	glass	knife	bottle	cup	spoon
tools	chisel	hammer	screwdriver	pliers	saw
buildings	apartment	barn	house	church	igloo
building parts	window	door	chimney	closet	arch
clothing	coat	dress	shirt	skirt	pants
insects	fly	ant	bee	butterfly	beetle
vegetables	lettuce	tomato	carrot	corn	celery
man-made objects	refrigerator	key	telephone	watch	bell

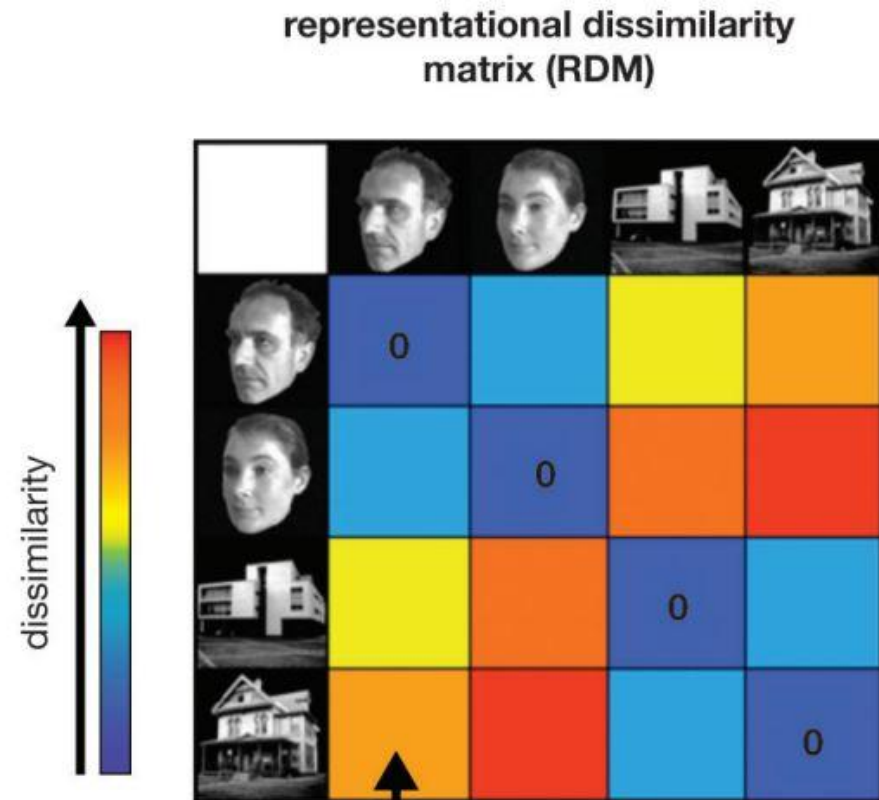


# Representational similarity analysis (RSA):

Between-subjects MVPA *without needing* spatial normalization

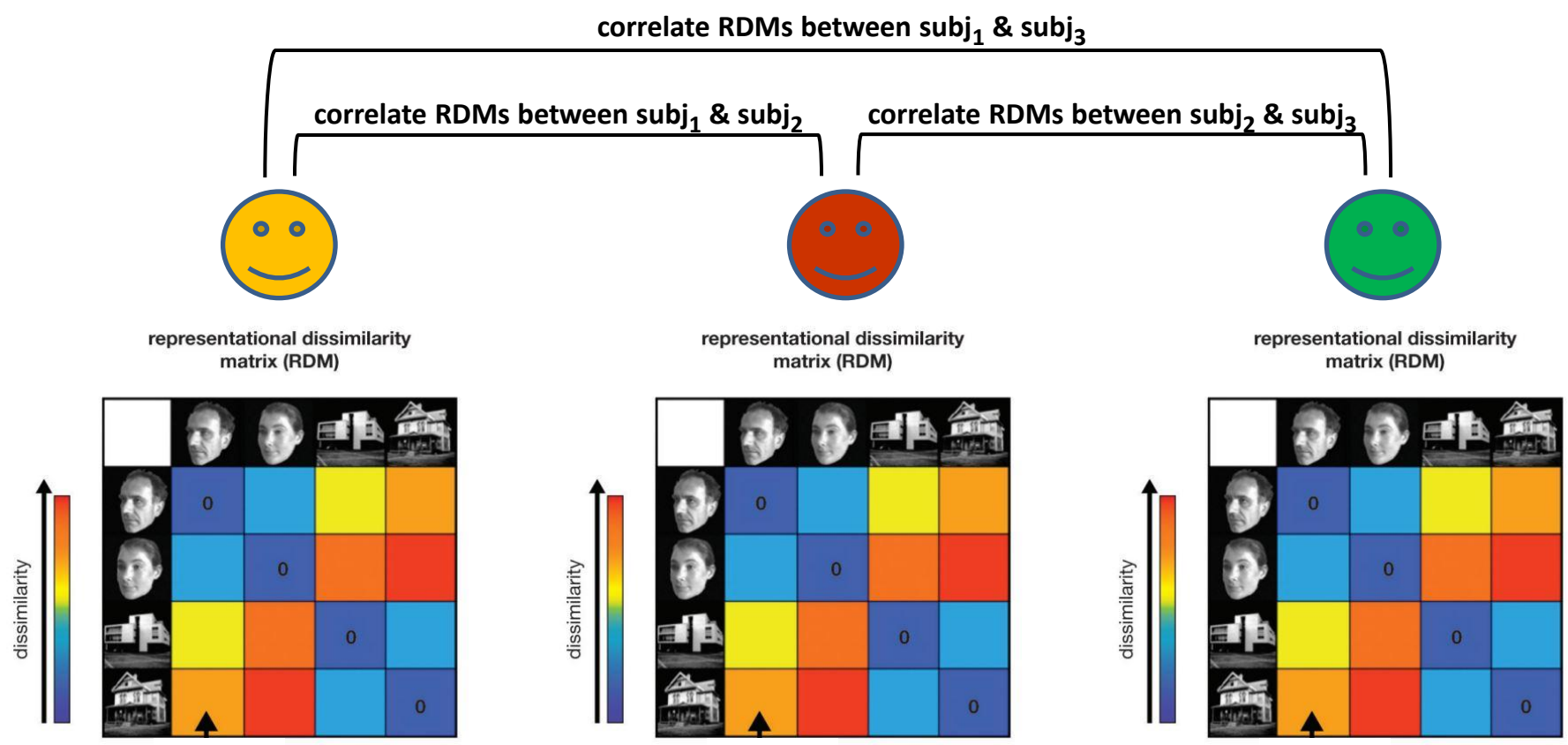
**Step 1, for each subject's data:** After selecting a set of voxels, calculate the correlation distance (i.e. dissimilarity) between the activation patterns of each pair of concepts

- The dissimilarities among the concepts reveal what kinds of information are represented in their activation patterns (e.g. visual information on the right)
- Each element in the RDM =  $1 - \text{correlation between activation pattern of concept}_1 \text{ and activation pattern of concept}_2$
- Dissimilarity between two faces or two buildings is low; dissimilarity between a face and building is high
- The dissimilarity between a stimulus and itself is 0



# Step 2: Calculate how well each subject's RDM correlates with the other subjects' RDMs

- Then, assess whether the average correlation between different subjects is high (and statistically significant)
  - (To see this analysis in use, see Raizada & Connolley, 2012)



# Take-home points about representational similarity analysis

- A form of between-subjects MVPA that assesses the similarity between different people's neural representations *without needing* spatial normalization
- This analysis compares different subjects' RDMs; it does NOT compare the same activation patterns between subjects
- Thus, this analysis abstracts away from the unique brain of each subject; the voxels selected from each subject can be different

# Representational similarity analysis can determine how similar the neural representations are between humans and monkeys!

(impossible to do using MVPA that requires spatial normalization)

Image from  
Kriegeskorte  
(2009)

## Legend:

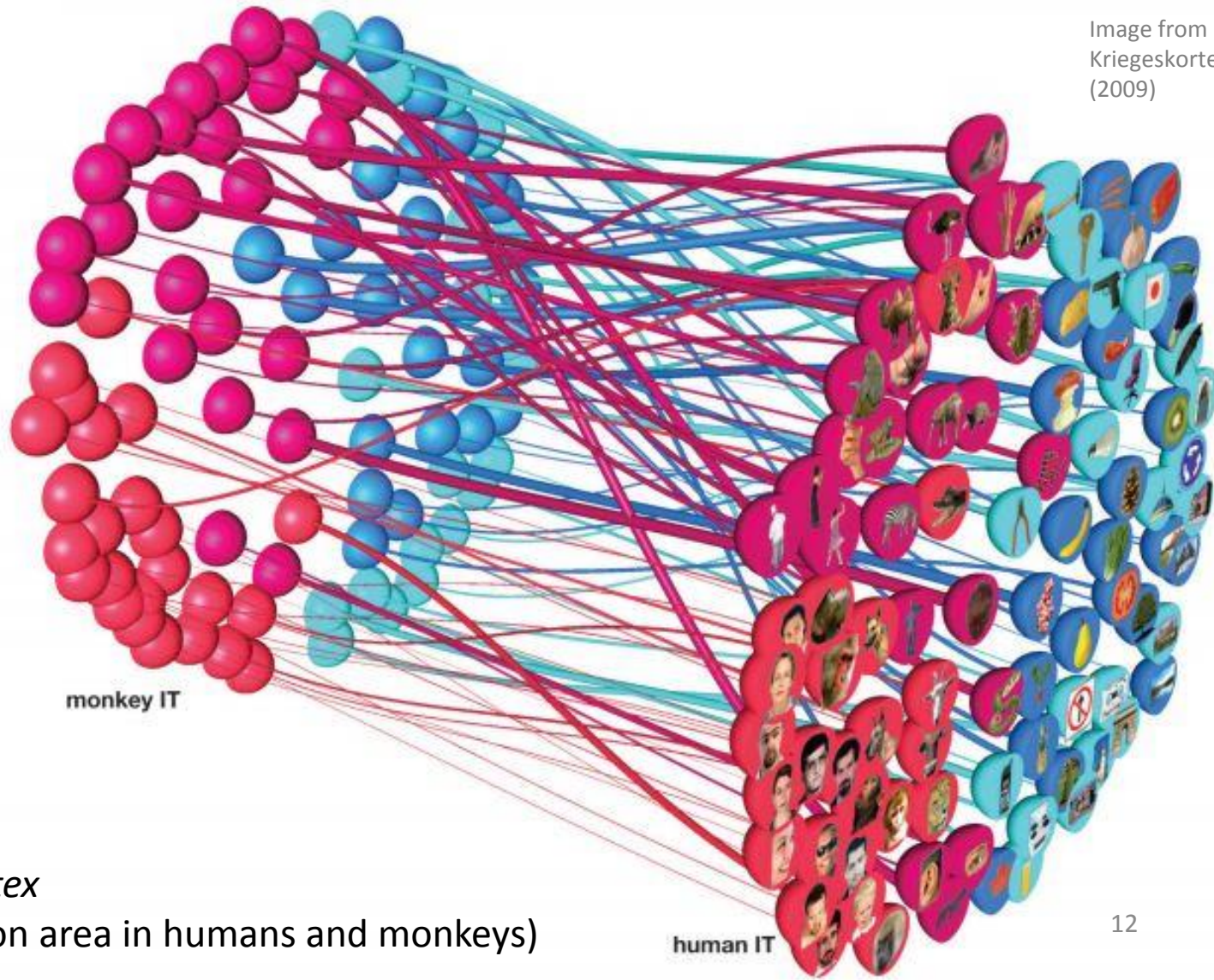
Face

Body

Natural object

Unnatural object

NOTE: Thick connecting lines mean “less alike” between monkey and human



*IT = inferotemporal cortex*  
(visual object recognition area in humans and monkeys)

Start Matlab 2012b (on desktop, or type "matlab" in Start menu to find it)

**NOTE: You MUST select Matlab 2012b, do NOT select 2014b**

MATLAB 7.10.0 (R2010a)

File Edit Debug Parallel Desktop Window Help

Current Folder: C:\Users\andrewba\Documents\MATLAB

Shortcuts How to Add What's New

Command Window

New to MATLAB? Watch this [Video](#), see [Demos](#), or read [Getting Started](#).

MATLAB desktop keyboard shortcuts, such as Ctrl+S, are now customizable.  
In addition, many keyboard shortcuts have changed for improved consistency across the desktop.

To customize keyboard shortcuts, use [Preferences](#). From there, you can also restore previous default settings by selecting "R2009a Windows Default Set" from the active settings drop-down list. For more information, see [Help](#).

[Click here](#) if you do not want to see this message again.

>>

Workspace

Stack: Select data to plot

Name	Value	Min
------	-------	-----

Command History

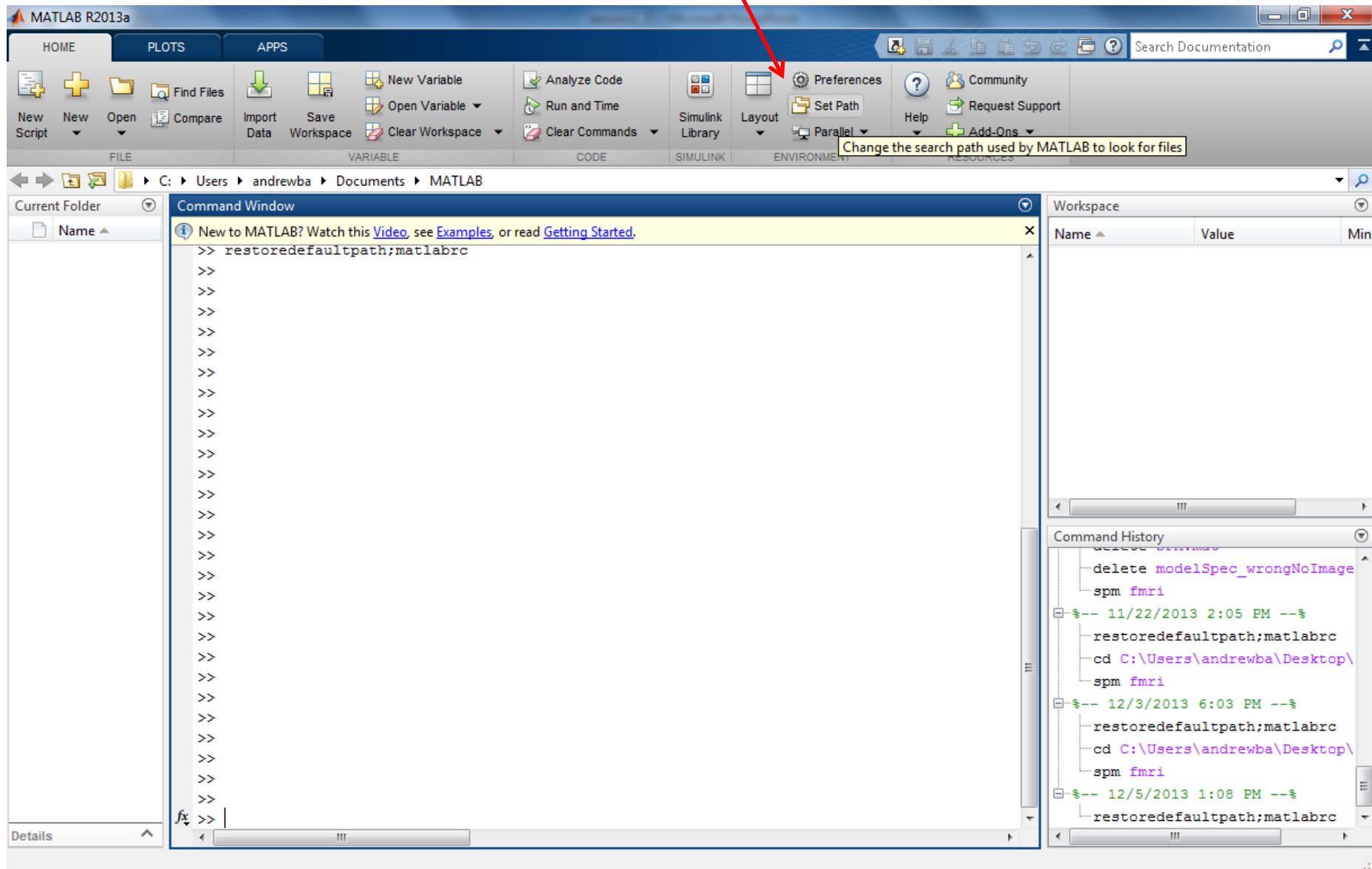
```
%-- 2/21/13 7:08 PM --%
spm fmri
%-- 2/22/13 2:30 PM --%
1-tcdf(2.75,30)
1-tcdf(3,30)
1-tcdf(3,40)
1-tcdf(2.75,40)
%-- 2/25/13 10:30 AM --%
spm fmri
clc
pwd
ll
ls
clc
%-- 2/27/13 3:25 PM --%
```

Details

Start Ready

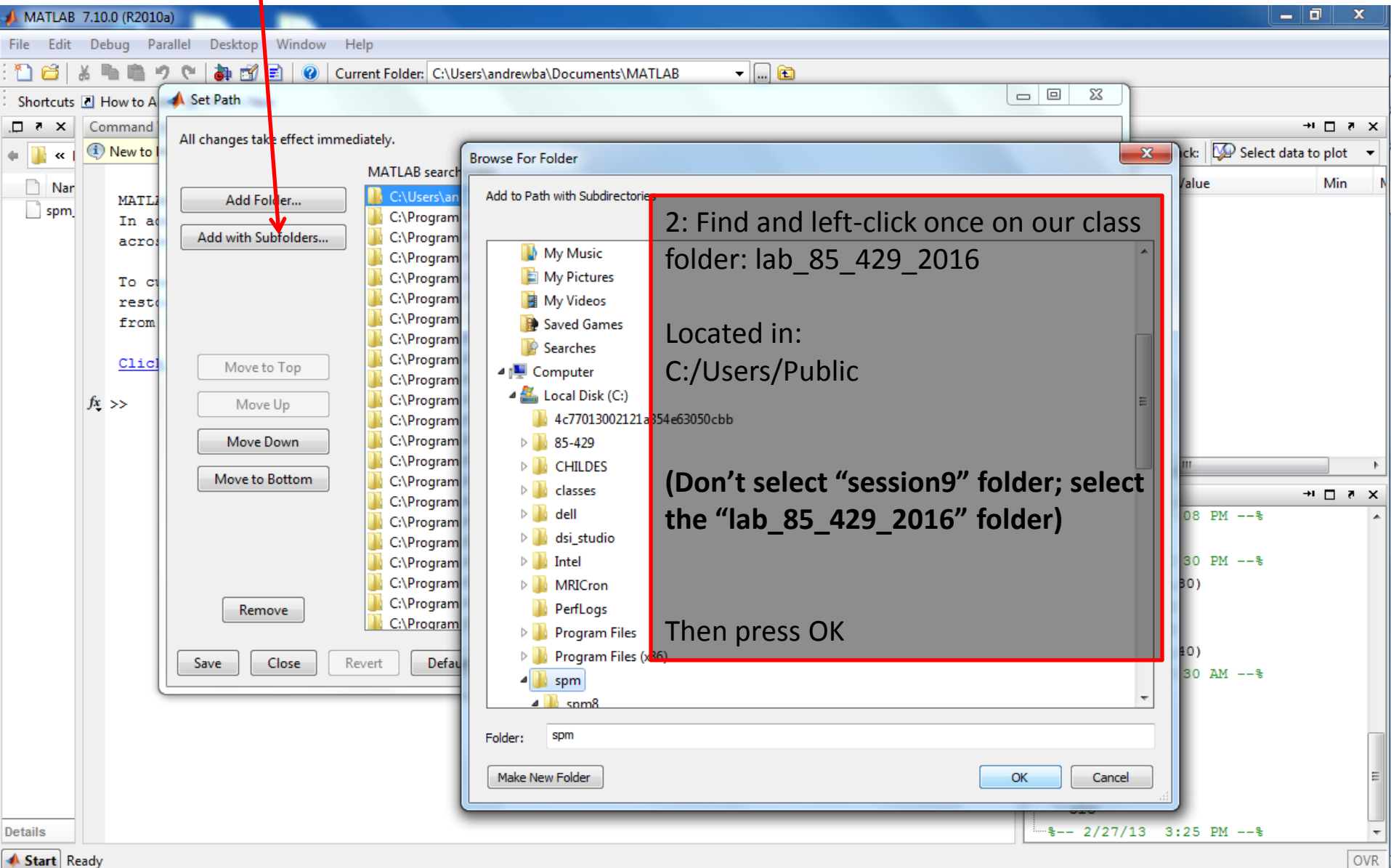
OVR

Select Set Path



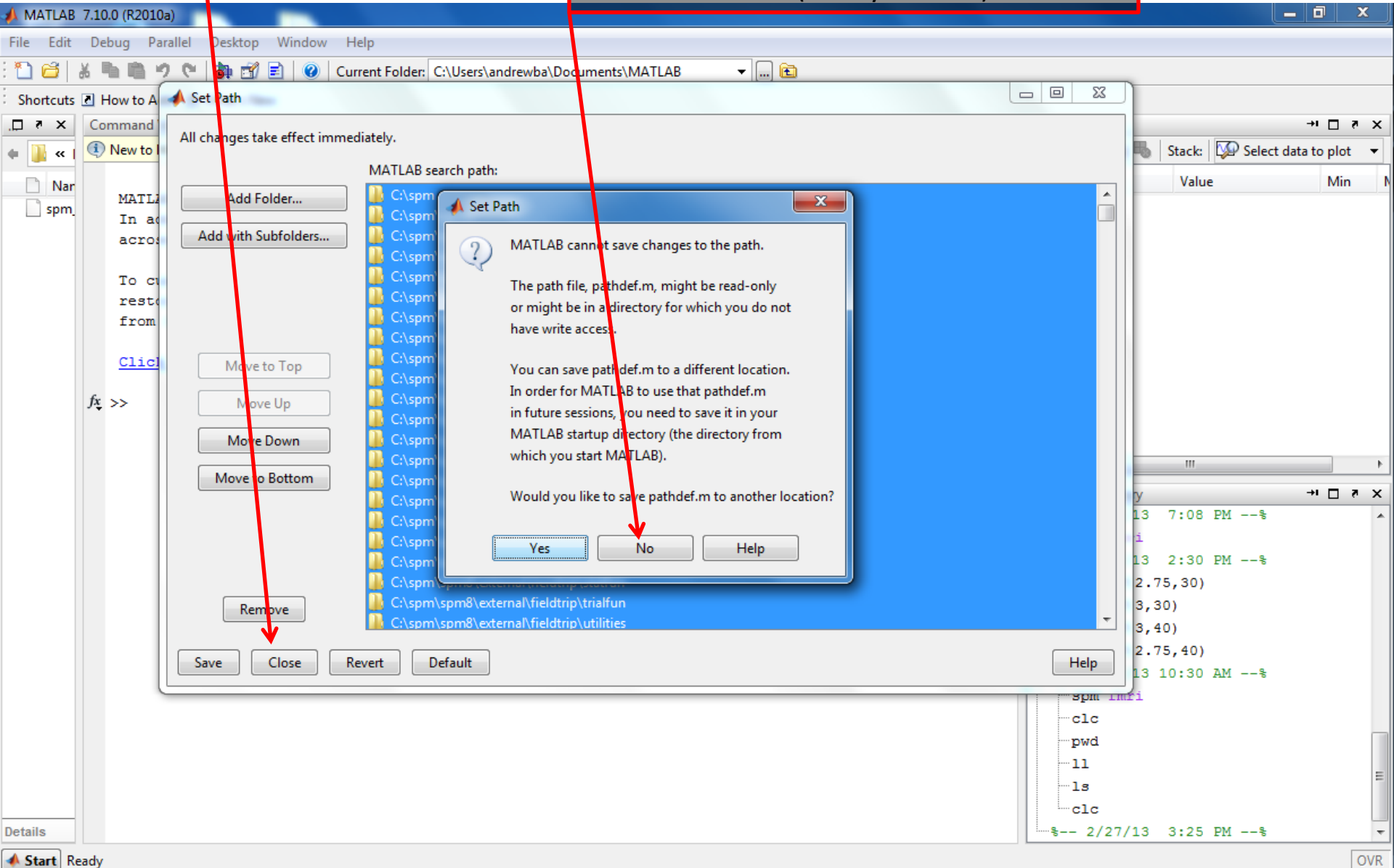


# 1: Select Add with Subfolders



1: Select Close

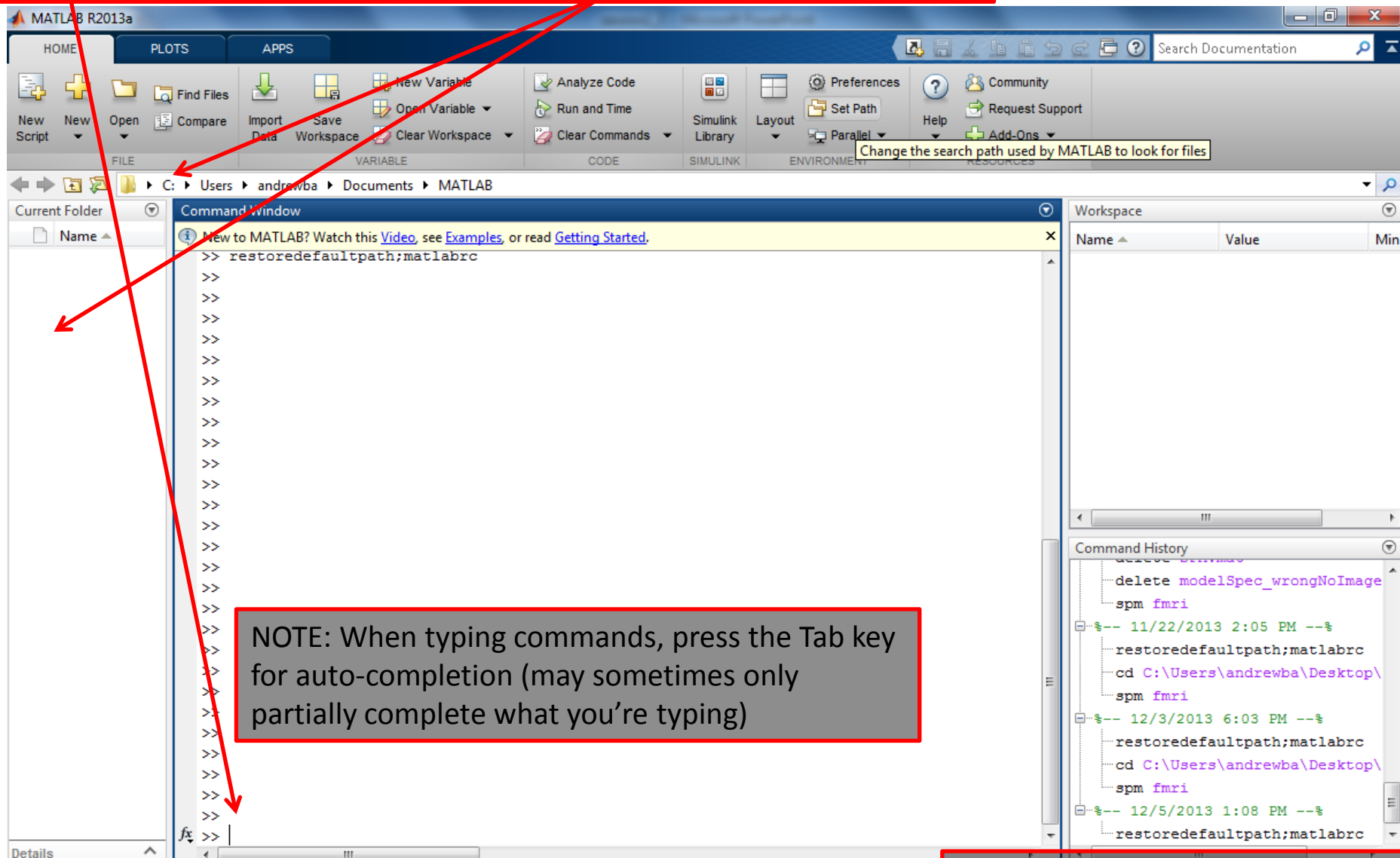
2: Select No if it asks to save the path file somewhere else (it may not ask)



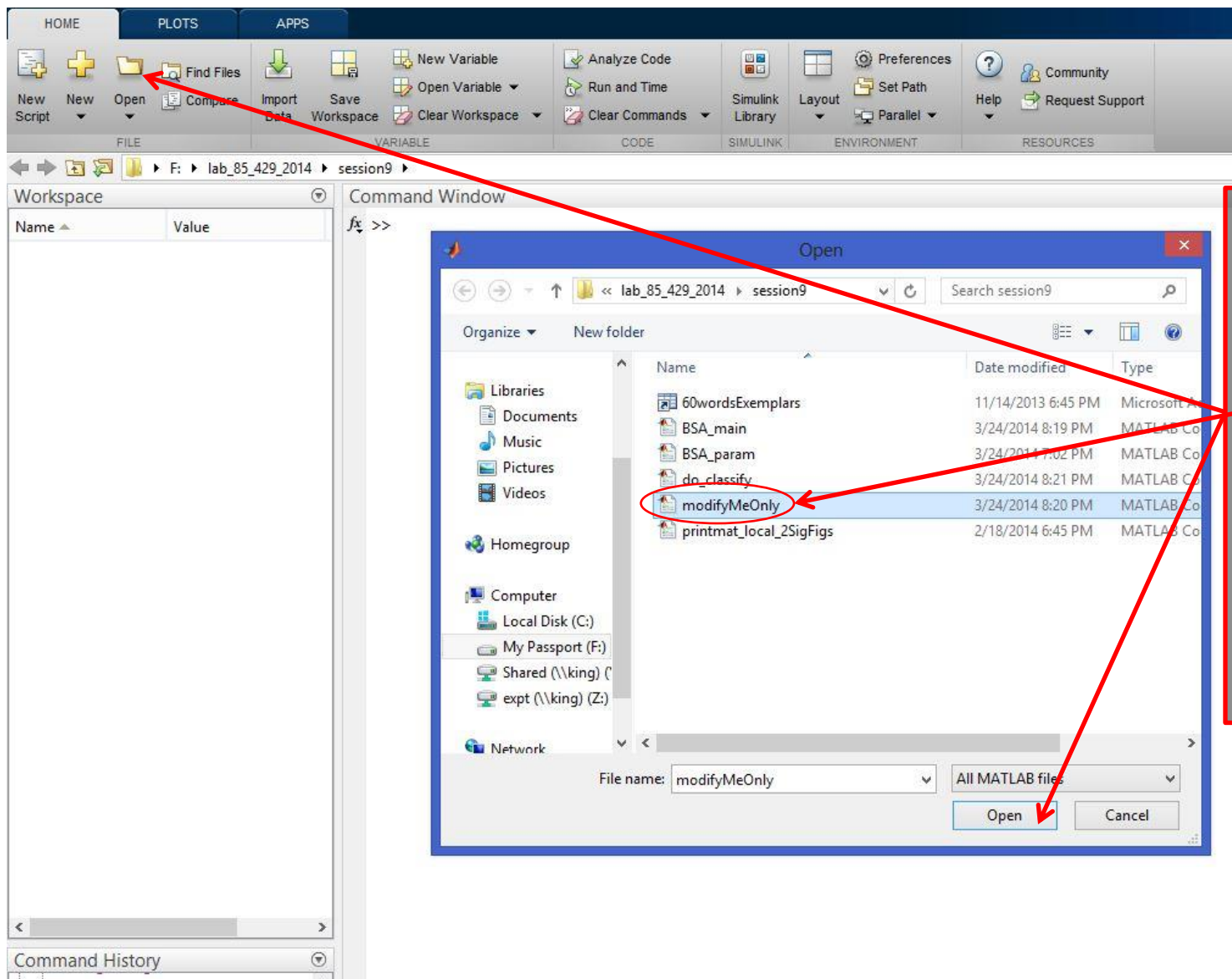
1: Go to the Matlab Command Window and type:

`cd C:/Users/Public/lab_85_429_2016/session9`

...(OR navigate there using the browser)



2: Next slide...



3. Open the following file found in the session9 folder (you should already be there):

modifyMeOnly.m

NOTE: Never modify any other file but this one

```
%% Andrew Bauer
% brainImagingLabSpring2014

% modify ***ONLY*** the contents of this .m file
% modifying ***any other files*** could result in your code not working
```

```
%% (1)Where are the voxels being selected from? Type in the ID number
% directly after "voxelsID = "
%
% But first, determine which ID number goes with which set of voxels:
%
% only frontal lobe ID = 1
% only temporal lobe ID = 2
% only parietal lobe ID = 3
% only occipital lobe ID = 4
% only subcortical areas ID = 5
% anywhere EXCEPT occipital lobe ID = 6
%
% NOTE: every lobe refers to both right and left lobes combined
```

```
voxelsID = 5
```

Change to "1" if  
it's not already

```
%% (2)How many voxels? Type the number in directly after "noVoxels = "
```

```
noVoxels = 80
```

```
%% (3)SAVE THIS FILE AFTER YOU CHOOSE (1) & (2) ABOVE!
%
% Now you are ready to run the classification analysis. Type (without
% the quotes) "do_classify" into the MATLAB command prompt and then press
% the enter key. Wait a little for it to run (it will display fold
% accuracies/etc. on the screen); then, when it is done, your screen will
% show the mean classification accuracies for all the object categories
% individually and combined, as well as the parameters that you chose
% (steps (1) & (2) above)
```

NOTE: There are three steps  
every time you will run MVPA:

- (1) Select where to draw voxels from;
- (2) Set how many voxels to select;
- (3) Type "do\_classify" (without quotes) into the MATLAB command prompt to start the classification

1: To acquaint yourself with how to do the classification, you will run classification on the frontal lobe only. Type "1" (without quotes) directly after "voxelsID = "; this is the numerical ID associated with the frontal lobe

(See the typed instructions in this file too)

NOTE: For the assignment, you will have to change the voxelsID value... do *not* choose "1" every time

2: Next slide...



```
## Andrew Bauer
## brainImagingLabSpring2014

## modify ***ONLY*** the contents of this .m file
## modifying ***any other files*** could result in your code not working
```

```
## (1)Where are the voxels being selected from? Type in the ID number
## directly after "voxelsID = "
##
## But first, determine which ID number goes with which set of voxels:
##
## only frontal lobe ID = 1
## only temporal lobe ID = 2
## only parietal lobe ID = 3
## only occipital lobe ID = 4
## only subcortical areas ID = 5
## anywhere EXCEPT occipital lobe ID = 6
##
## NOTE: every lobe refers to both right and left lobes combined
```

```
voxelsID = 1
```

```
## (2)How many voxels? Type the number in directly after "noVoxels = "
```

```
noVoxels = 50
```

Change to "50" if it's not already

```
## (3)SAVE THIS FILE AFTER YOU CHOOSE (1) & (2) ABOVE!
##
## Now you are ready to run the classification analysis. Type (without
## the quotes) "do_classify" into the MATLAB command prompt and then press
## the enter key. Wait a little for it to run (it will display fold
## accuracies/etc. on the screen); then, when it is done, your screen will
## show the mean classification accuracies for all the object categories
## individually and combined, as well as the parameters that you chose
## (steps (1) & (2) above)
```

3: Now we will set the number of voxels. In the PLoS One paper (the study that published these data), 80 voxels were drawn from the whole brain; so we should use fewer than 80 if we are focusing on just one lobe of the brain

**NOTE: For the assignment, you will have to change the noVoxels value... do *not* choose "50" every time**

4: Next slide...



```
1 %% Andrew Bauer
2 % brainImagingLabSpring2014
3
4 % modify ***ONLY*** the contents of this .m file
5 % modifying ***any other files*** could result in your code not working
6
7
8
9
10 %% (1)Where are the voxels being selected from? Type in the ID number
11 % directly after "voxelsID = "
12 %
13 % But first, determine which ID number goes with which set of voxels:
14 %
15 % only frontal lobe ID = 1
16 % only temporal lobe ID = 2
17 % only parietal lobe ID = 3
18 % only occipital lobe ID = 4
19 % only subcortical areas ID = 5
20 % anywhere EXCEPT occipital lobe ID = 6
21 %
22 % NOTE: every lobe refers to both right and left lobes combined
23 %
24 voxelsID = 1
25
26
27
28
29 %% (2)How many voxels? Type the number in directly after "noVoxels = "
30
31 noVoxels = 50
32
33
34
35
36 %% (3)SAVE THIS FILE AFTER YOU CHOOSE (1) & (2) ABOVE!
37 % Now you are ready to run the classification analysis. Type (without
38 % the quotes) "do_classify" into the MATLAB command prompt and then press
39 % the enter key. Wait a little for it to run (it will display fold
40 % accuracies/etc. on the screen); then, when it is done, your screen will
41 % show the mean classification accuracies for all the object categories
42 % individually and combined, as well as the parameters that you chose
43 % (steps (1) & (2) above)
```

5: Next, you *must* save the “modifyMeOnly.m” file before running classification

PLOTS APPS

Find Files Compare Import Data Save Workspace New Variable Open Variable Clear Workspace

VARIABLE

F:\lab\_85\_429\_2014\session6

Command Window

```
>>
>>
fx >> do_classify
```

Value

'F:\lab_85_429_2014\session6'
120
1

111111  
111101

efScor

8)/(fa

1:13

lustCe

s

entral

6: After saving, go to the MATLAB command prompt and type “do\_classify” (without the quotes), then press enter

7: Next slide...

MVPA report: Object category BY subject accuracy (%) =

	04383B	04408B	04550B	04647B	04564B
MEAN_category	61.69	59.94	56.44	55.82	55.65
animals	47.80	55.59	62.37	67.46	68.81
bodyparts	63.73	57.29	33.90	67.12	34.58
buildings	51.86	24.75	63.05	61.02	46.78
buildprts	61.02	46.78	63.73	55.93	42.71
clothing	63.39	47.12	51.53	60.00	53.22
frniture	67.80	81.02	70.85	22.71	68.47
insects	49.83	73.56	58.98	57.29	67.46
kitchen	69.49	61.02	62.71	55.25	47.12
manmade	62.71	83.05	77.63	64.07	50.17
tools	65.76	51.86	42.71	42.03	56.61
vegetbles	82.37	61.02	35.59	68.81	71.19
vehicles	54.58	76.27	54.24	48.14	60.68

	04619B	04480B	04639B	04597B	04605B
MEAN_category	54.32	53.33	53.08	52.40	51.72
animals	50.17	28.81	48.47	66.44	68.81
bodyparts	64.07	60.00	57.63	56.95	35.59
buildings	44.07	60.00	72.54	43.05	46.44
buildprts	48.47	61.69	60.68	52.20	38.64
clothing	46.78	50.17	51.86	25.08	44.75
frniture	52.20	27.12	73.22	66.10	50.85
insects	62.03	80.34	65.76	53.90	51.86
kitchen	54.24	48.14	46.78	57.63	49.15
manmade	48.14	43.73	46.78	32.54	82.03
tools	66.44	73.22	50.17	62.71	34.24
vegetbles	47.80	43.73	22.03	57.97	73.90
vehicles	67.46	63.05	41.02	54.24	44.41

	04617B	MEAN_subj
MEAN_category	49.49	54.90
animals	39.32	54.92
bodyparts	49.15	52.73
buildings	51.53	51.37
buildprts	63.73	54.14
clothing	55.93	49.98
frniture	54.58	57.72
insects	27.46	58.95
kitchen	47.12	54.42
manmade	57.63	58.95
tools	49.83	54.14
vegetbles	33.56	54.36
vehicles	64.07	57.10

8: After the classification is finished, you will see an “MVPA report” that lists the classification accuracy for each of the twelve object categories (i.e. mean over the five items per category); and the mean accuracy over ALL categories at the top (called “MEAN\_category”)

You will also see a column of accuracies for each subject. The last column is the mean subject (mean over all subjects)

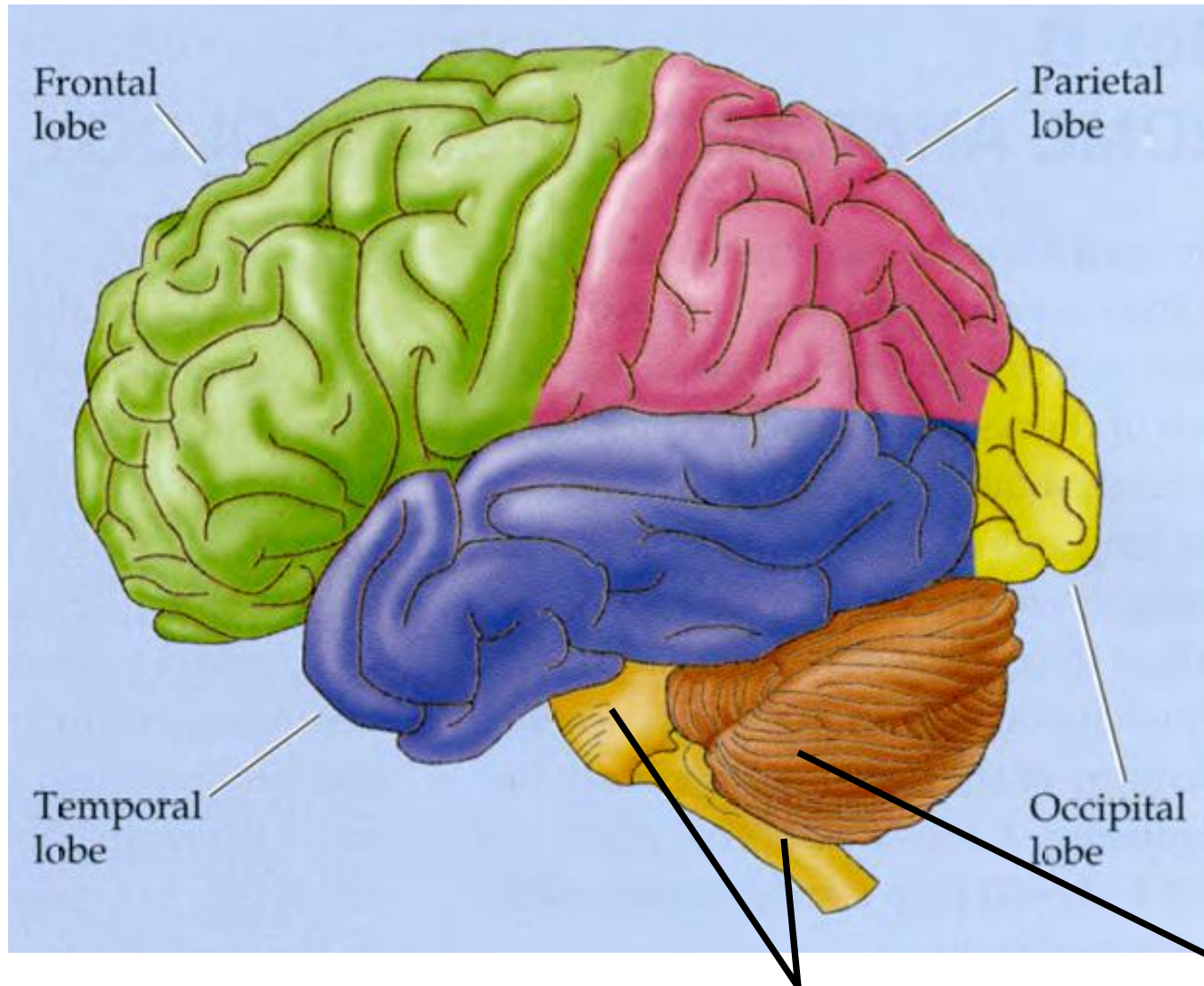
(You will also see the two parameters that you set directly before running the classification)

no. of voxels: 50  
voxels are from: only frontal

- This is all the guidance/information that you should need to do this session's assignment
- **You do not need to start up spm or xjview for this assignment**
- If you must restart MATLAB for whatever reason, after you start MATLAB **be sure to set the path again (just our lab folder) and cd back to the "session9" directory (see beginning slides)**
- See the slides below for any help with orientation terms and planes, brain anatomy, and general functional neuroanatomy

# Brain partitioned into four lobes

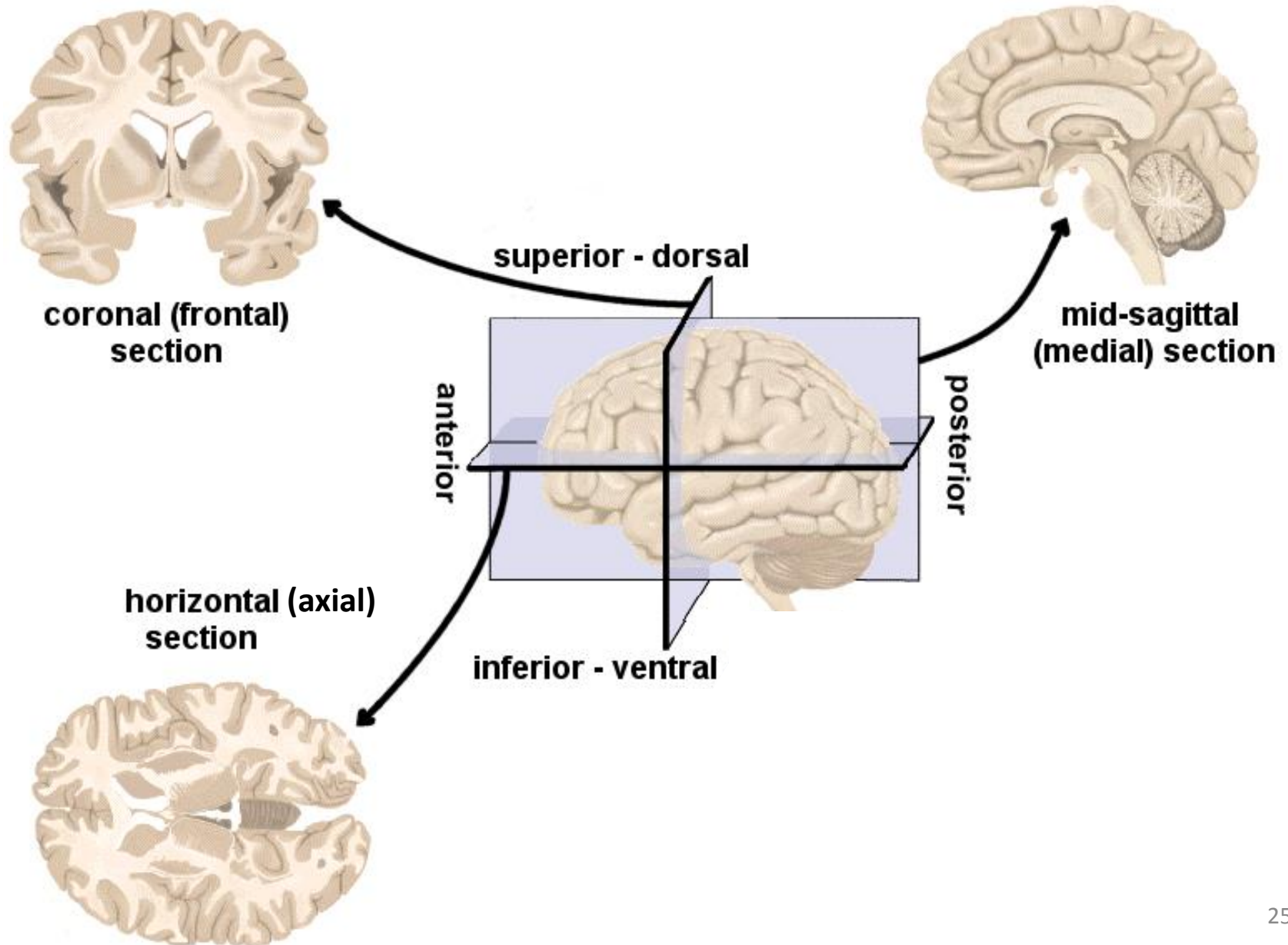
(... plus some extra regions)



NOT "lobes": Some subcortical brain regions & cerebellum



# Orientation terms and planes

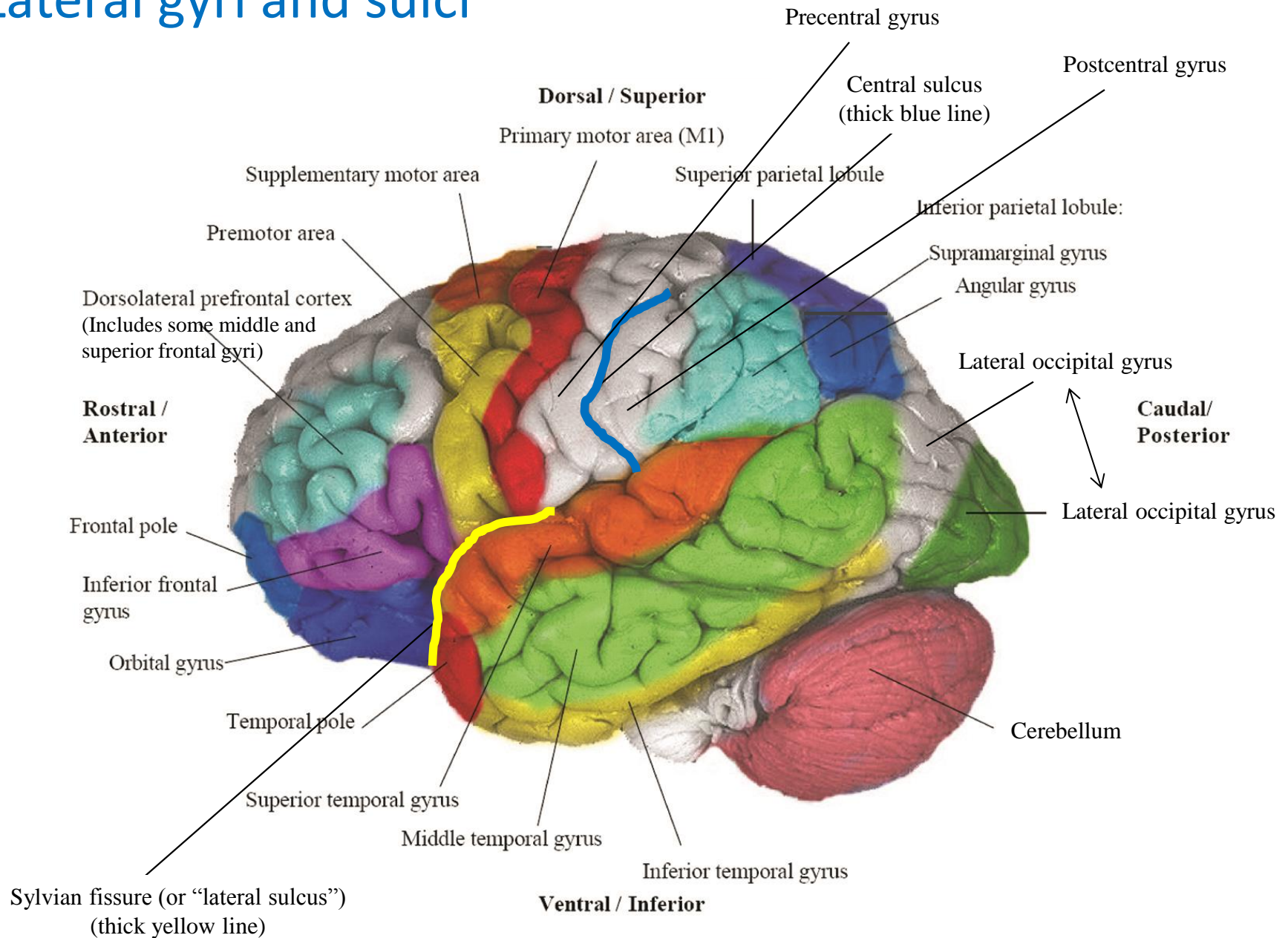


We are using the **Automated Anatomical Labeling (AAL)** atlas in this lab (very similar to the atlases of your two quizzes)

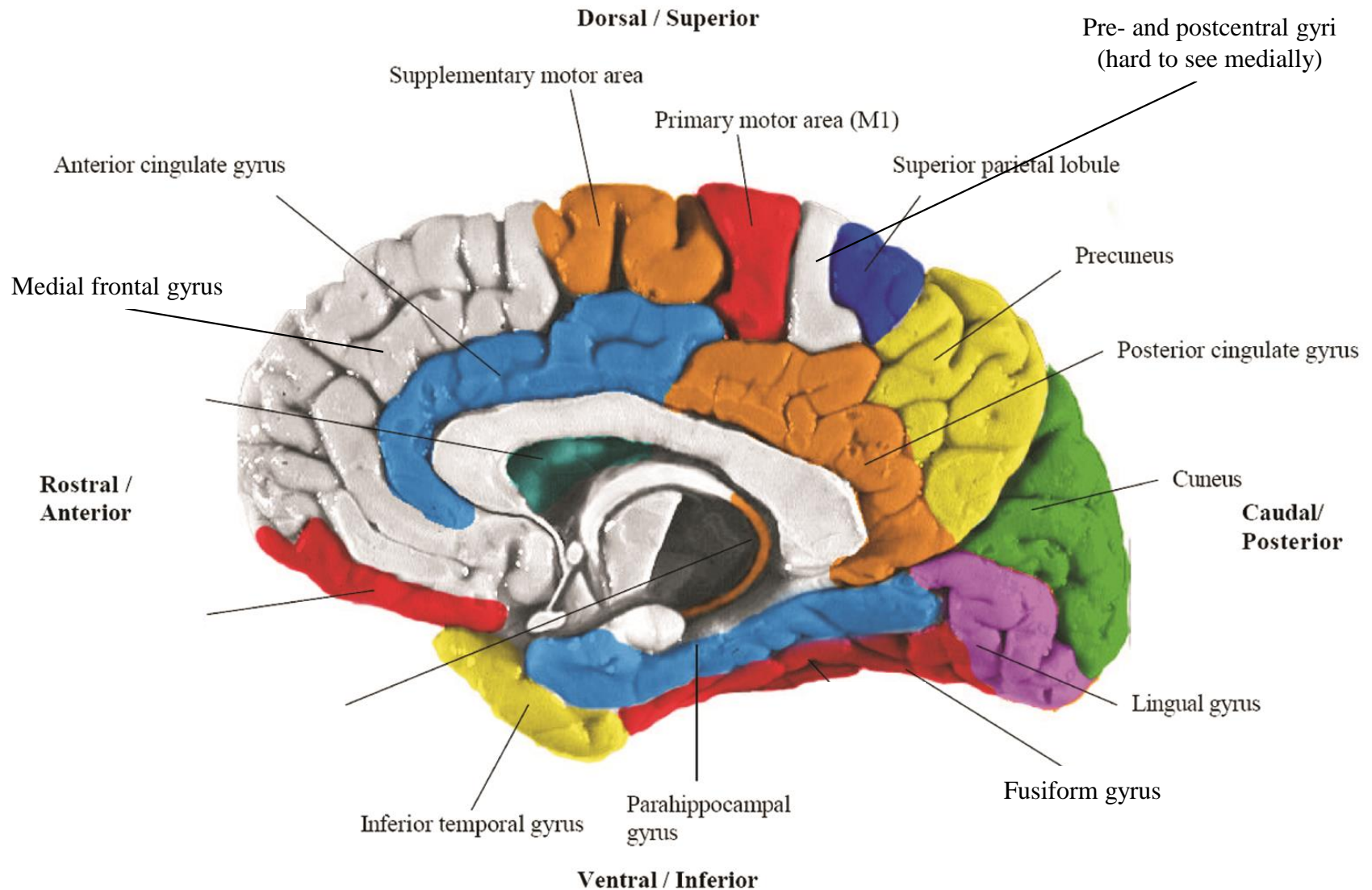




# Lateral gyri and sulci

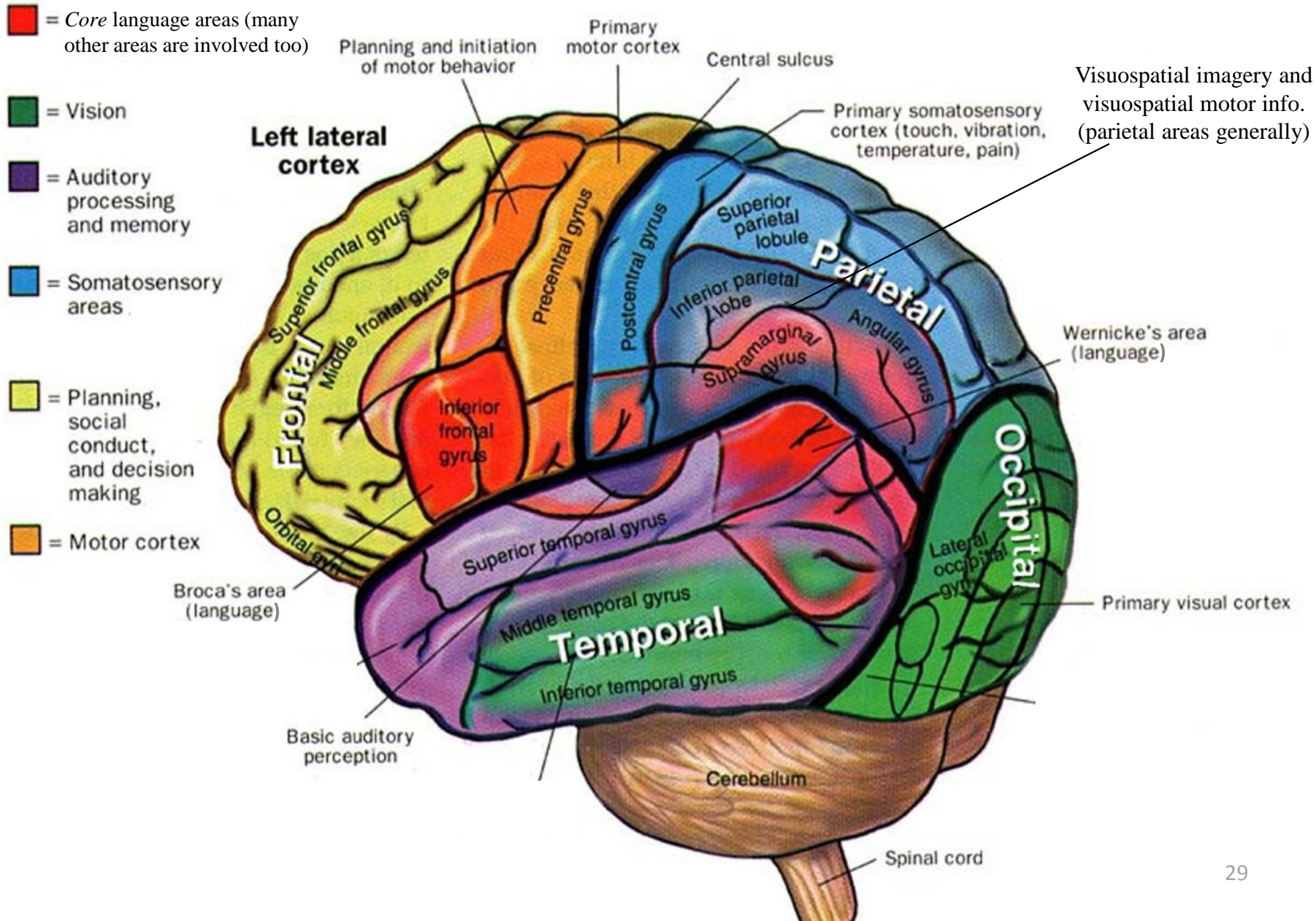


# Medial gyri (some redundancy w/previous slide)





# General functional neuroanatomy



# References

Just, M. A., Cherkassky, V. L., Aryal, S., & Mitchell, T. M. (2010). A neurosemantic theory of concrete noun representation based on the underlying brain codes. *PLoS One*, 5(1), e8622. doi:10.1371/journal.pone.0008622

Kriegeskorte, N., Mur, M., & Bandettini, P. (2008). Representational similarity analysis - connecting the branches of systems neuroscience. *Frontiers in Systems Neuroscience*, 2(November), 4. doi:10.3389/neuro.06.004.2008

Kriegeskorte, N. (2009). Relating Population-Code Representations between Man, Monkey, and Computational Models. *Frontiers in Neuroscience*, 3(3), 363–73. doi:10.3389/neuro.01.035.2009

Norman, K. A., Polyn, S. M., Detre, G. J., & Haxby, J. V. (2006). Beyond mind-reading: multi-voxel pattern analysis of fMRI data. *Trends in Cognitive Sciences*, 10(9), 424–30. doi:10.1016/j.tics.2006.07.005

Raizada, R. D. S., & Connolly, A. C. (2012). What Makes Different People's Representations Alike : Neural Similarity Space Solves the Problem of Across-subject fMRI Decoding. *Journal of Cognitive Neuroscience*, 24(4), 868–877.