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Overview

1. Estimated time required to complete

Between 1 and 2 hours.

2. Responses to submit on Canvas

Your article's cognitive skill

Item 1:

Is the brain region that you named in your answer to Part 1, question 2.B.iii.b part of the Neurosynth brain map?

[Jump to Item 1 instructions](#)

Item 2:

Screenshot of the brain region from Item 1

[Jump to Item 2 instructions](#)

Item 3:

Paste the table that lists the first 10 Neurosynth "Associations" for the brain region from Item 1

[Jump to Item 3 instructions](#)

Item 4:

Based on the names of the associated Neurosynth meta-analyses, what one other cognitive skill do you think your brain region might be involved in?

[Jump to Item 4 instructions](#)

Comparing your article's cognitive skill with another skill

Item 5:

Screenshot of the 3D surface model, with blobs from both Neurosynth maps

revealed by the "cut planes".

[Jump to Item 5 instructions](#)

Item 6:

Screenshot of one ROI cluster from the intersection of both Neurosynth maps

[Jump to Item 6 instructions](#)

Item 7:

Where in the brain is this cluster, and why did you choose it?

[Jump to Item 7 instructions](#)

Item 8:

Paste the table that lists the first 10 Neurosynth "Associations" for this cluster's coordinates

[Jump to Item 8 instructions](#)

Item 9:

What do you think this cluster's brain region does?

[Jump to Item 9 instructions](#)

Questions about this activity

Item 10:

What was confusing about this assignment, or what questions did it bring to mind that you would like to understand about neuroimaging or meta-analysis?

(This question does not appear elsewhere in the instructions, only right here.)

Item 11:

What would be another interesting thing to do with Mango and/or Neurosynth for this assignment? Is there anything you would like to do, even if you're not sure whether it's possible to do with these tools?

(This question does not appear elsewhere in the instructions, only right here.)

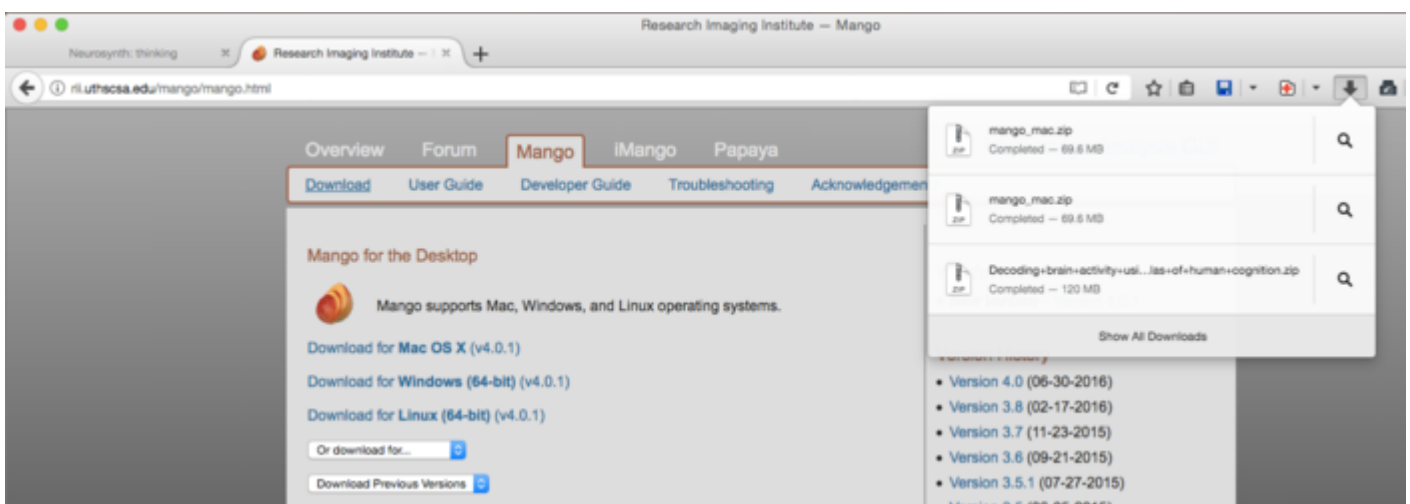
Get ready to neuroimage!

1. Install Mango software

You will need to install the Mango viewer software to do this part of the assignment. In general, I prefer not to ask students to download software, but currently there are no browser-based MRI image viewers available online that provide all the features we need. I have chosen Mango to download because it is non-commercial (so that there are no ads or additional software that come with it), free¹⁾, and designed to work with all major operating systems.

<http://ric.uthscsa.edu/mango/mango.html>

- Click on the link for your operating system, and follow the prompts. The website includes some instructions too.



Mango download page

2. Download a file to use with Mango

As you know, functional MRI (fMRI) measures differences in blood oxygenation levels. For technical reasons, fMRI images usually need to be relatively low-resolution (they look blurry). Also, the goal of most fMRI studies is to identify small, specific regions of the brain that are associated with a cognitive task. The final results of most fMRI studies (which are usually called “activation maps”) don't look like a picture of a whole brain, or even of a recognizable part of a brain. They just depict little three-dimensional blobs that mark the location of the brain regions most highly associated with the cognitive task from the study. So, the blobs that are visible in an fMRI activation map are like the Xs that “mark the spot,” but without the underlying treasure map! Looking at activation maps on their own is like looking at a blank piece of paper with an X on it – it tells you that there is, in fact, a treasure/important brain region, but not where to find it.

To understand which locations the activation blobs (the “Xs” in the treasure map analogy) mark in the brain, we need to “put the map under the Xs.” In our case, the “map” is an image of a whole brain (in 3D of course). The image that we will use is a composite image based on averaging together the structural MRI images of 152 individual brains. (Structural MRI images are high-resolution images that look more like a photograph than functional MRI images, and they don't measure blood oxygenation.) This average image is a little too smooth around the edges to look like a real, single person's brain, but that's the point: we want to understand where an activation blob would be located on average, in anyone's brain.

So, please download a file available in this Canvas folder:

https://canvas.vt.edu/courses/57758/files/folder/writing_assignment/brain

File name: `mni_icbm152_t1_tal_nlin_asym_09a_acMasked.nii.gz`

(NOTE: If your computer asks you to unzip this (or do anything else) with this file, respond “No.” The only reason this file is zipped (compressed to reduce the size of the file) is to make it easier to download and save. Mango will read this file without you having to unzip it first.)

Make a note of where the downloaded file gets saved, so that you can open it later. You can ask your web browser to show you where the file was saved in a number of ways, including:

- In Firefox: Under the “Tools” menu, select “Downloads.”
- In Chrome: Under the “Window” menu, select “Downloads.”

This file is a version of the standard 152-person average brain created by the Montreal Neurological Institute (MNI) in Canada. You will see this abbreviation a few times during this activity, because the MNI originated a number of widespread neuroimaging techniques. The only difference between this version of the file and the standard version available online from the MNI is that your instructor has removed the face, skull and other non-brain portions of the image, so that it will look more like a brain when we eventually view a 3D rendering of it.

3. View the structural average brain in Mango

Now that we have ...

1. The Mango viewing software
2. A copy of the structural average brain file

... we can look at the average brain.

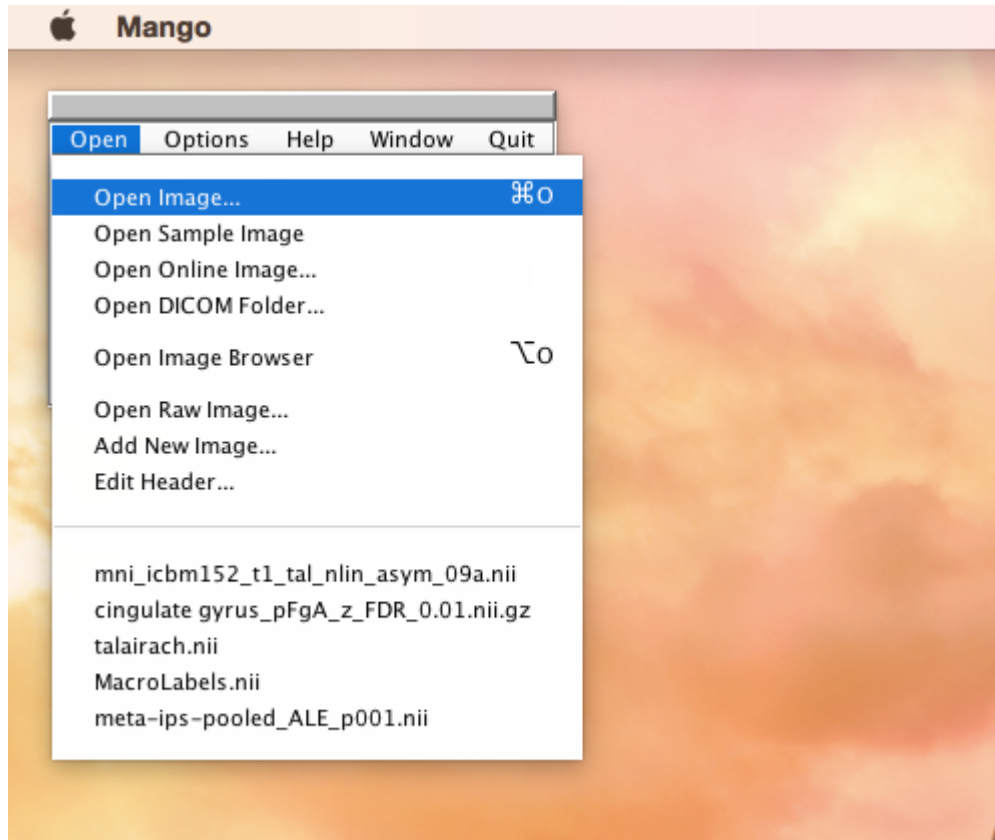
3.1 Start the Mango app

I feel funny writing instructions to you youngsters about how to open apps, because you know, you're more familiar with "apps" than me. However, problems can occur, and please don't hesitate to ask if they do.

3.2 Open the structural average brain file: the "image"

Somewhat confusingly, Mango uses different words when it refers to 1) the structural brain image, and 2) the functional activation map image. To me, these both count as "brain images." However, Mango calls the structural brain file an "**image**," and it refers to functional activation maps as "**overlays**." This is only important because there are different menu commands for the two kinds of files.

To load the average structural brain image, use the mouse menus to select "Open image," as seen here:



Load the average structural brain image using the “Open image” command

Select the file that you downloaded from Canvas.

Congratulations! You made a brain appear! It should look like this:



This is what you should see in Mango after loading the average structural brain image.

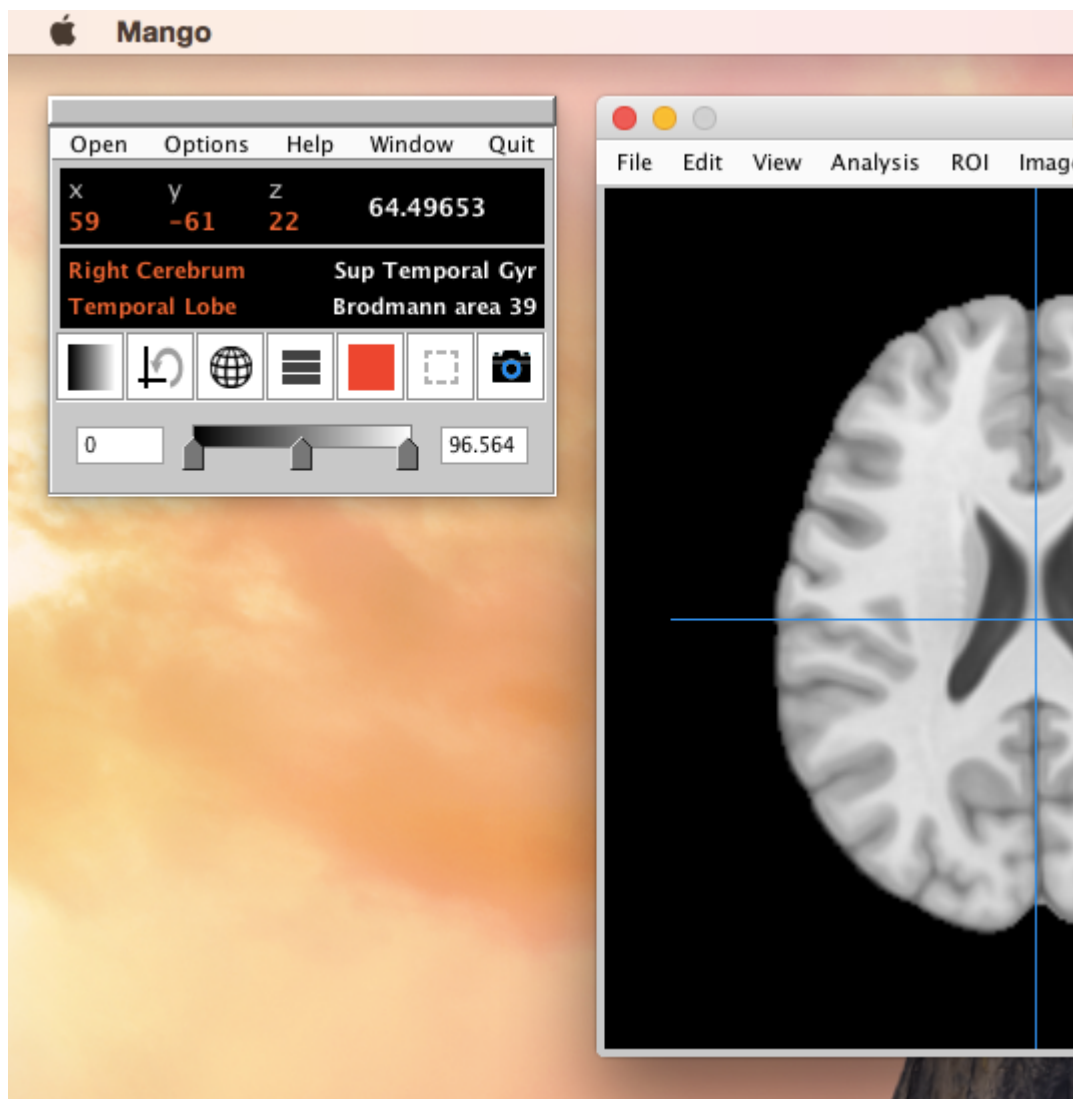
3.3 Enable the atlas feature

Mango can show us the name of the brain region that is under the mouse, but we need to enable this feature first. Mango looks up the most likely name for each brain region using a data set called an “atlas.” Click on the button in the main Mango window (which is actually the smaller window) that looks like a sphere, and then select “Atlas → MNI (Nearest Gray Matter)”.



Enable the atlas feature

Technically, the names of the brain regions (the “atlas labels”) only describe regions of gray matter. However, because our structural brain image is an **average** brain, it is very possible that some individual people's brains will have gray matter in locations that correspond to white matter in the average brain. A consequence of this is that functional activation blobs can sometimes appear to be located in white matter on the average brain. In those cases, we still want to know what to call that part of the brain.



Look in the main Mango window to see the name of the brain region located under the mouse pointer

Mango will display four different names with different levels of specificity. The one we want is the one at the upper right of the atlas window, e.g. "Sup Temporal Gyr" for superior temporal gyrus.

HINT: if the atlas abbreviations are unclear, try Googling them.

Get brain maps from the Neurosynth website

1. Go to the Neurosynth homepage in your browser

<http://neurosynth.org/>

2. Get a meta-analysis for your article's topic

Neurosynth provides a very interesting service: it allows us to view something like the most probable functional activation map that we could expect from an fMRI study if all we knew about the study were its keywords.

Because of this, the terms or keywords that we enter in Neurosynth can make a big difference in the activation maps that we get back. In particular, the level of specificity of the terms can make a big difference. For example, as we know from class, the English word “memory” refers to many different cognitive processes, including many that probably rely on distinctly different brain regions. If we enter the query “memory,” Neurosynth will return a map showing the likely locations of functional activation across all published studies that featured the word “memory.” This will lump together “apples and oranges” - distinctly different patterns of activity produced by qualitatively different cognitive processes that happen to share the term “memory” in their names, like “episodic memory” and “working memory.”

2.1 Find the Neurosynth term to use for your article

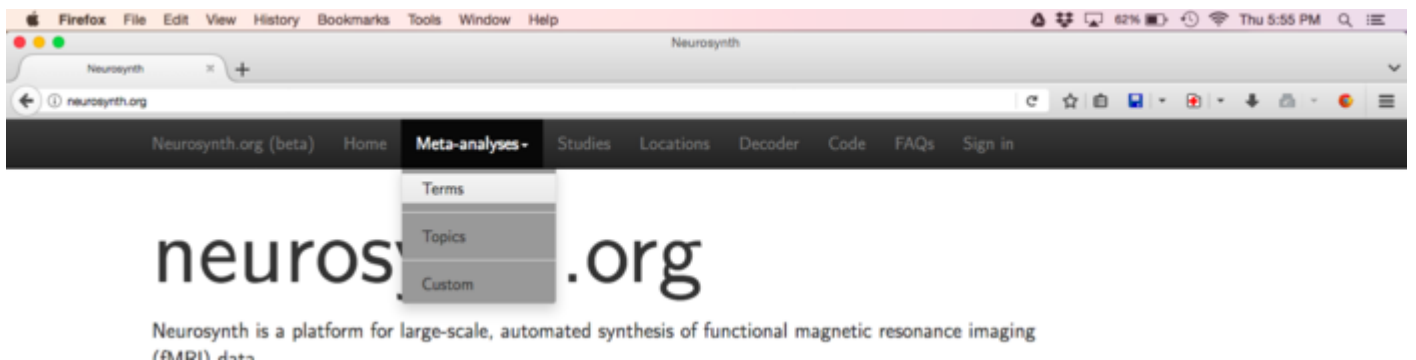
Please use the term that corresponds to the article you reviewed for Part 1 of this assignment:

Neurosynth terms for each article from Part 1

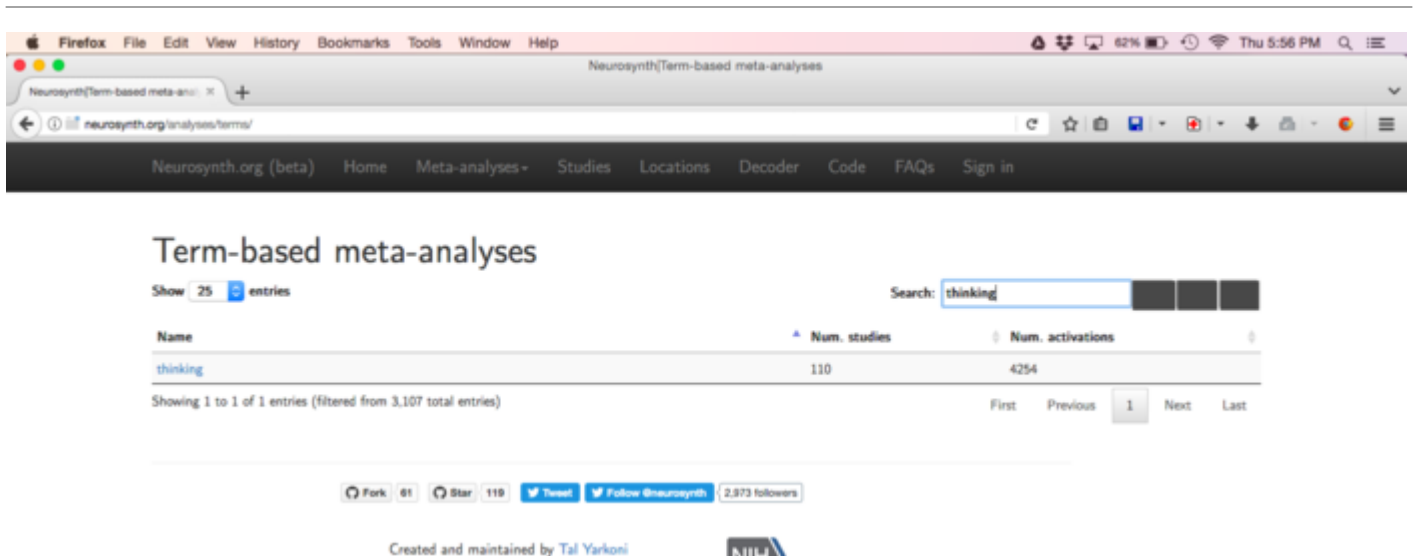
General Topic	Article Title	Neurosynth term to use
Perception	Untangling invariant object recognition.	Object recognition
	Towards a neural basis of music perception.	Music
Attention	What is specific to music processing? Insights from congenital amusia.	Musicians
	Why visual attention and awareness are different.	Awareness
	Moving towards solutions to some enduring controversies in visual search	Visual attention
Memory	The neurobiology of semantic memory.	Semantic memory
	Working memory capacity and its relation to general intelligence.	Working memory
	False memories and confabulation.	Memory retrieval
	The contribution of sleep to hippocampus-dependent memory consolidation.	Consolidation
	Observing the transformation of experience into memory.	Memory encoding

General Topic	Article Title	Neurosynth term to use
Language	Towards a functional neuroanatomy of speech perception.	Speech perception
	Reflections on mirror neurons and speech perception.	Mirror neuron
Decisions/Problems	Neuroeconomics: cross-currents in research on decision-making.	Decision making
	Cognitive and AI models of reasoning.	Reasoning
Intelligence	Genetics and general cognitive ability (g).	Cognitive control
	Evolution of the brain and intelligence.	Intelligence
Consciousness	Consciousness cannot be separated from function.	Consciousness
Neurosynth terms validated 2017-10-29 by adc		

2.2 View the meta-analysis for your term



Select the Meta-analyses → Terms menu



Type in your term and click the link that appears below

The screenshot shows the Neurosynth website interface for a meta-analysis on the term "thinking". The browser is Firefox, and the URL is <https://neurosynth.org/analyses/terms/thinking/>. The page title is "thinking" and it is described as "An automated meta-analysis of 156 studies".

The interface includes a navigation menu with "Neurosynth.org", "Home", "Meta-analyses", "Studies", "Locations", "Genes", "Decoder", "Code", and "FAQs". A search bar for another term is located at the top right.

The main content area has three tabs: "Maps", "Studies", and "FAQs". The "Maps" tab is active, displaying three brain slices: a coronal slice (labeled D and V), a sagittal slice (labeled A and P), and an axial slice (labeled L and R). The axial slice is labeled with "z = 0". Red clusters are visible on all three slices, indicating areas of significant activation.

Below the slices, there are controls for the z-score (set to 0), a "What's here?" button, and coordinate input fields for X, Y, and Z (all set to 0). To the right, there is a "Layers" panel with three layers: "thinking: association test", "thinking: uniformity test", and "anatomical". The "Color palette" is set to "red", and "Crosshairs" and "Labels" are checked. The "Positive/Negative" is set to "positive". The "Thresholds" and "Opacity" sliders are also visible.

Viola! You see before you the results of a neuroimaging meta-analysis.

2.3 Download the brain map image for your topic

The screenshot shows the Neurosynth website interface for the term "thinking". At the top, there is a navigation bar with links for Home, Meta-analyses, Studies, Locations, Genes, Decoder, Code, and FAQs. Below the navigation bar, the word "thinking" is displayed in a large font, followed by the text "An automated meta-analysis of 156 studies". To the right of the title is a search box labeled "Search for another term:". Below the title, there are three tabs: "Maps", "Studies", and "FAQs", with "Maps" being the active tab. The main content area displays three brain slices: a coronal slice (top left) labeled "D" and "V" with "y = 0", a sagittal slice (top right) labeled "A" and "P" with "x = 0", and an axial slice (bottom left) labeled "L" and "R" with "z = 0". Each slice shows red clusters of activation. To the right of the brain slices is a control panel. The "Layers" section contains three items: "thinking: association test", "thinking: uniformity test", and "anatomical". Each item has a trash icon and a download icon. The "Color palette" section has a dropdown menu set to "red" and a checked checkbox for "Crosshairs". The "Positive/Negative" section has a dropdown menu set to "positive" and a checked checkbox for "Labels". The "Thresholds" section has a slider set to 0. The "Opacity" section has a slider set to 1. Below the brain slices, there is a "z-score: 0" label and a "What's here?" button. Below that, there are input fields for X: 0, Y: 0, and Z: 0.

Click on the "download arrow" icon at the end of the "reverse inference" line.



SAVE the file to a location that you can find later.

3. Get a meta-analysis for an "other" topic

Repeat the instructions for one of the “other topics” that you chose for your answer to Part 1, question 2.B.ii (“What is known about the relationship between this cognitive skill and two of the following general cognitive skills?”).

Please use the following Neurosynth terms:

Neurosynth terms for each “other skill” from Part 1, question 2.B.ii

General cognitive skill	Neurosynth term to use
Perceptual skills	Perception
Selective attention	Selective attention
Memory	Memory
Mental imagery	Imagery
Language	Language comprehension
Intelligence or reasoning skills	Intelligence
Neurosynth terms validated 2017-10-31 by adc	

Explore brain map images using the Mango viewer

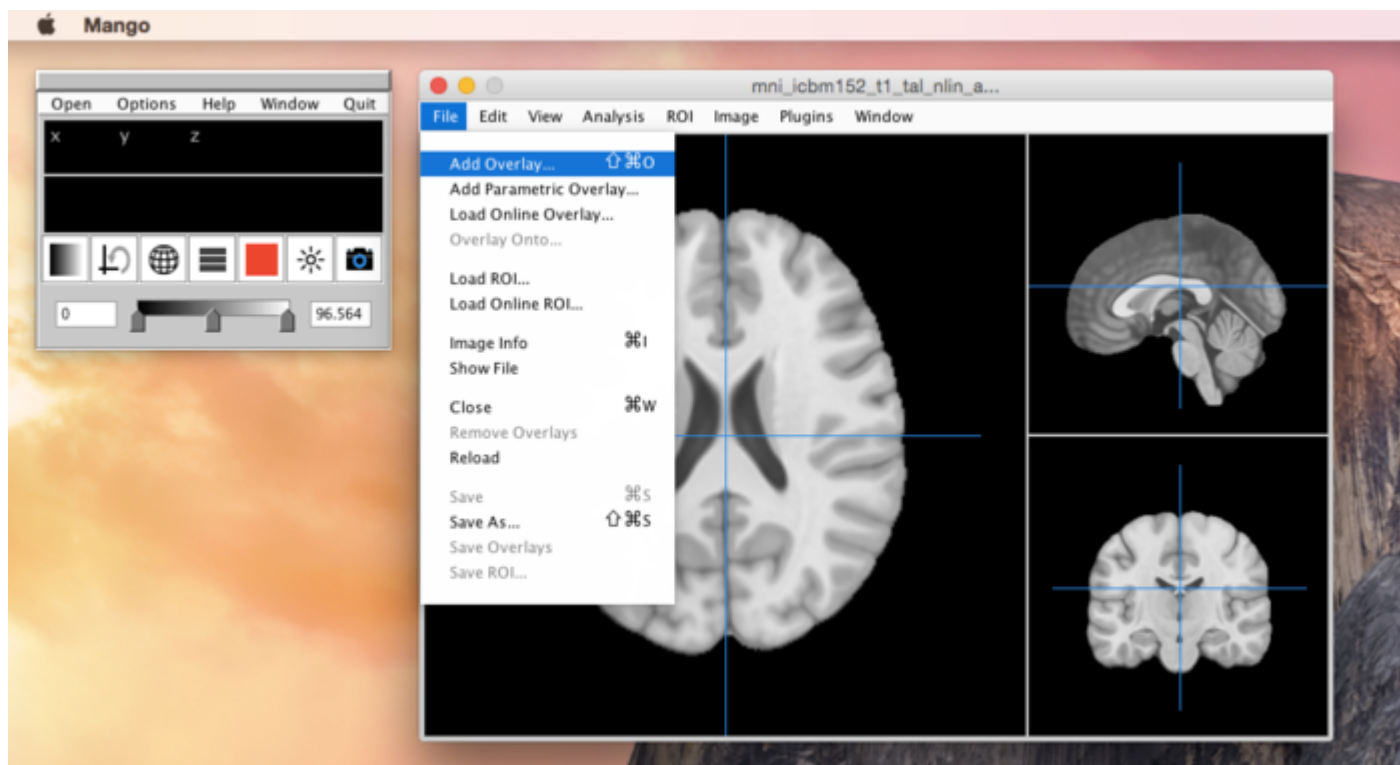
Follow the instructions above to open the average structural brain image and enable the atlas feature.

Again, the average structural brain image forms the background over which your functional activation maps will be overlaid.

1. Examine the Neurosynth brain map for your article's cognitive skill

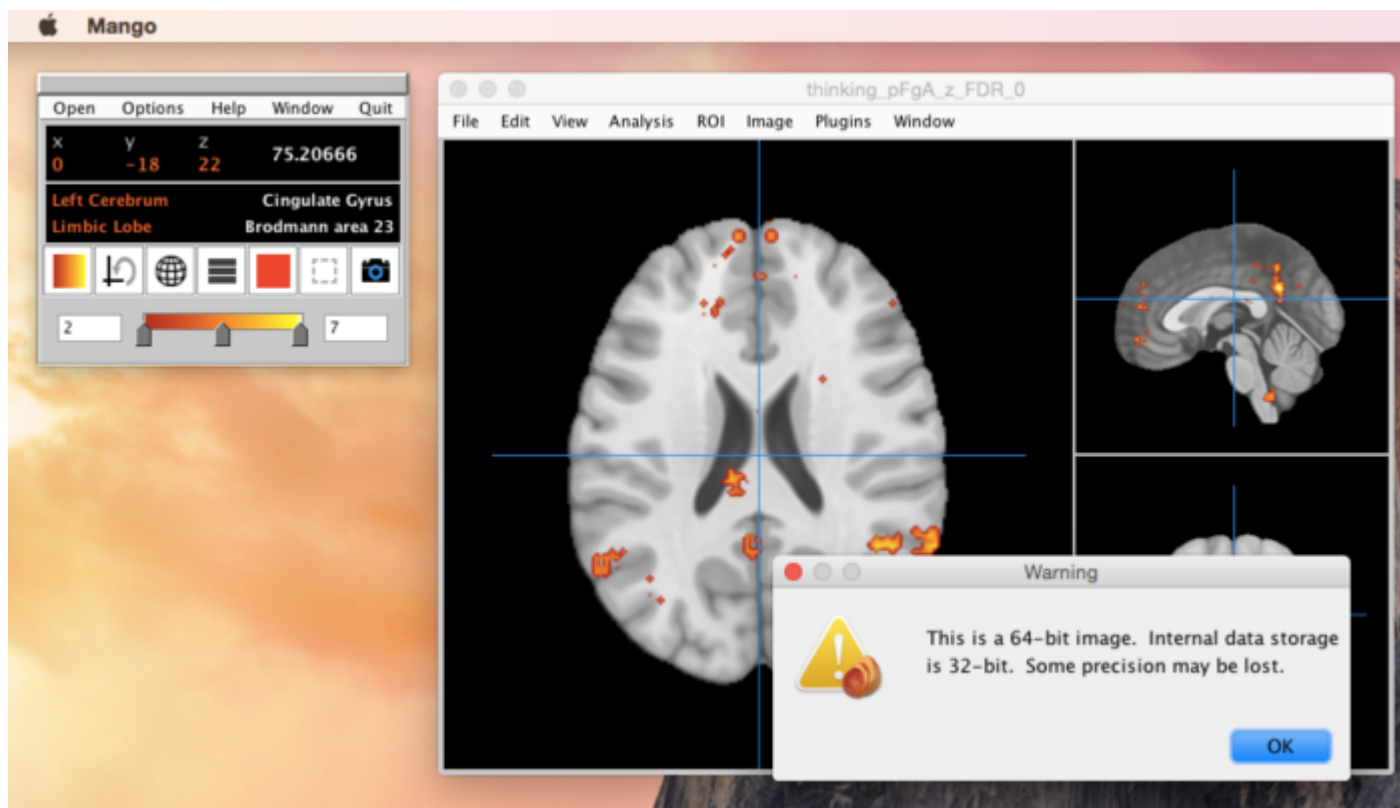
Mango refers to functional brain maps as “overlays,” in contrast to the structural brain map, which is the “image.” Mango can only open one “image” at a time, but it can have multiple overlays open simultaneously.

Now we can open the first file that you downloaded from Neurosynth. **DON'T** use the first Mango window's “Open” menu, even though that sounds like a good choice. Instead, use the “File” menu on the viewing window that is displaying the structural brain image.



Use the File menu in the viewing window to open a functional activation map as an overlay.

If you get the warning shown in the figure below, just click "OK."



You can ignore this warning and click OK.

With any luck, you will now see bright orange/red blobs superimposed atop the structural brain image.

NOTE: Mango (and most other neuroimaging software) cannot display the entire brain at once. Mango's viewing window shows three different 2D views centered on the same crosshairs, but these three views still can't reveal the entire 3D volume of the brain. This means that there can always be functional activation blobs "hiding" somewhere just out of view.

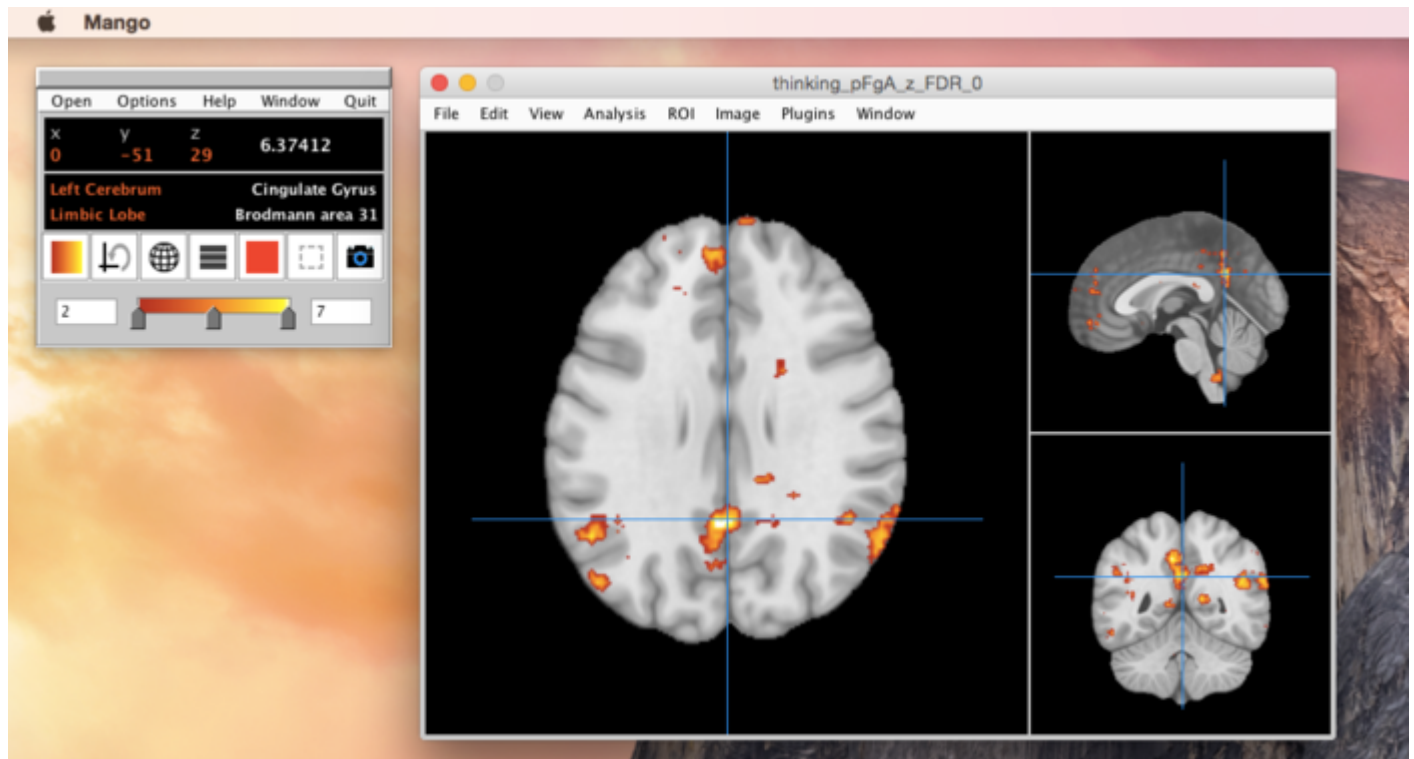
If you don't see any orange/red activation blobs after loading your first overlay, try clicking and dragging the mouse all around the brain, moving the blue crosshairs, until you've looked all over the entire brain. There is almost certainly a blob there somewhere.

It is possible that your particular Neurosynth map might not have any activation blobs that happen to be visible at the current view of the brain.



The display should look like this after adding the first functional activation map overlay.

Even if you are rewarded by seeing a brainful of orange/red activation blobs immediately after loading your overlay, you should still click and drag the mouse around to explore the entire brain. You can do this within any of the three different panes of the viewing window. Mango will keep all three panes in sync, i.e. focused on the same point marked by the crosshairs.



Click on activation blobs to focus all 3 viewing panes.

1.1 Find the key brain region from your article

Refer to your answer to Part 1, question 2.B.iii.b (“Pick what you think is the most important of these brain regions, and briefly state why you think it's the most important.”).

Does any of the Neurosynth brain map activation fall in this region?

For this assignment, we can answer the question using what we can call “the click/scroll/hover method.”

1. Find the part of the brain in question by clicking/scrolling around on the average structural brain map, and then getting the proper atlas label to appear by hovering the mouse over the correct part of the structural brain map. This will put you in the correct general area of the brain.
 - Get a good approximate sense of the boundaries of the region that corresponds to the atlas label, by clicking/scrolling/hovering the mouse around until the atlas label changes.
 - Now you can judge whether any of the Neurosynth brain map activation regions (i.e. the colorful blobs) fall in that region.

This is not a precise method, but it is more than good enough for our purposes. The small amount of extra precision that we would get by using other methods would not be worth our time here. It is actually quite tricky to name the brain regions covered by an activation blob. Just for starters, the boundaries of brain regions are not always clear-cut, and even the names for various brain regions are not agreed upon by all scientists. Then there is the whole issue of using an average structural brain to guide us, when individual brains can be so different from each other.

Potential problems with finding a brain region

Here are two problems that we **do** need to address before going ahead with the click/scroll/hover method.

Different names for the same brain region

“What if the atlas label names are different from the names that our review article used for brain regions?”

Answer: You can trust your judgment and decide when you think the Mango atlas label is just another name for the region mentioned in your article. More to the point, I (the instructor) will trust your judgment, too. Just be sure to write down **both** names in your answer, for example:

“My article named the 'middle cingulate gyrus,' but Mango doesn't have an atlas label with that name. Part of the Neurosynth activation map did overlap with the middle of the region that the atlas labeled 'Cingulate gyrus,' though. Therefore, my answer is yes, the middle cingulate gyrus does appear on the Neurosynth activation map.”

I'm not a brain expert (yet)

“How do I find a brain region in Mango without clicking/scrolling/hovering over the whole dang brain? You (the instructor) said I didn't need to know how to read MRIs.”

Answer: Try Googling the name of the brain region you're looking for. Try searching for images, if necessary. Then use the results as a guide of where to look in Mango.

Is the key brain region from your article part of the Neurosynth brain map?

Item 1:

Is the brain region that you named in your answer to Part 1, question 2.B.iii.b part of the Neurosynth brain map?

(This is a one-word answer: yes or no.)

Take a screenshot of the region

Item 2:

Screenshot of the brain region from Item 1

Regardless of whether your answer is yes or no, take a screenshot of Mango when you have placed the **crosshairs** on the region in question.

NOTE that the crosshairs are not the same thing as the mouse pointer; to place the crosshairs (the two thin blue lines) you need to click the mouse.

PLEASE USE THE MANGO SCREENSHOT TOOL and not some other method for taking screenshots. Clicking on the “camera” icon in the main (smaller) Mango window will save an image file to the same folder that holds the **average structural brain image** file (i.e. `mni_icbm152_t1_tal_nlin_asym_09a_acMasked.nii`).

Rename the file “[name of the key brain region].png” and upload it to Canvas. Examples of a good file name: “SupTempGyr.png” or “Superior Temporal Gyrus.png”.

NOTE If your answer for Item 1 was “no,” then the crosshairs will mark an “unactivated” region of the structural brain image. That is OK. You still need to upload the screenshot.

1.2 What other cognitive skills are associated with this region?

Copy down the MNI coordinates for the region

Copy down the stereotactic coordinates of the point in the brain marked by the crosshairs.

While the crosshairs are still marking the region, hover the mouse right on top of them. The reason for moving the mouse is that by default, the main Mango window (the small one) displays the coordinates for the location the mouse, and not for the crosshairs.

The 3D (or “stereotactic”) coordinates are given by 3 numbers (X,Y,Z), which appear in red font. The fourth number that appears in white font to the right of them is not important for us here.

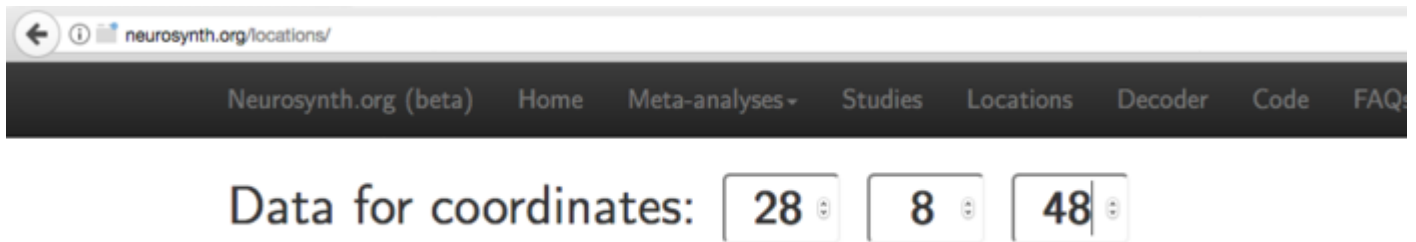
There are two main coordinate systems that scientists have developed for referring to locations in the average brain: Talairach coordinates, and MNI coordinates. The two systems are similar, but not identical. The brain images that we are using (both the average structural brain and the Neurosynth functional maps) are in MNI coordinates.

Enter the coordinates in a Neurosynth “locations” search

Click the “Locations” link at the top of any Neurosynth website page, or navigate your browser to this url:

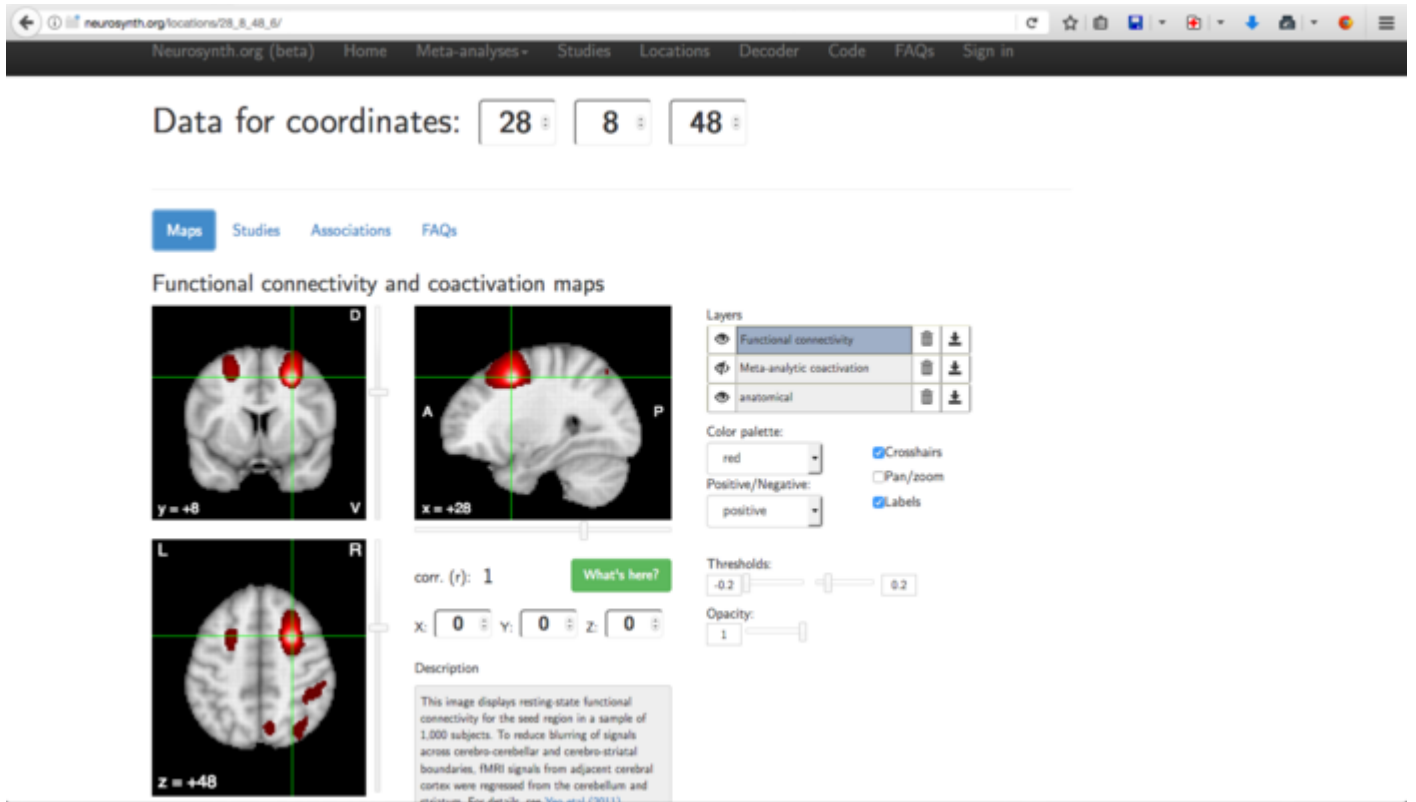
<http://neurosynth.org/locations/>

Type the three numbers (including any minus signs) into the “Data for coordinates” fields, and press the Enter key on your keyboard when you're done.



Type the X,Y,Z numbers into these three fields.

The web page will reload, showing a brain map with crosshairs and bright colors centered on the coordinates that you entered.



We will not use the “Functional connectivity and coactivation maps” that appear, but here is a brief explanation about them for those who might be interested. This color map represents something different from the meta-analysis activation maps that you produced earlier. This time, the color map indicates the probability that the location at these coordinates is “connected” with other locations in the brain. I put “connected” in quotes because the color map is not based on measures of **anatomical** connectivity, even though those exist (e.g. diffusion tensor imaging maps of white matter pathways).

Neurosynths' "Functional connectivity and coactivation maps" display something like the probability that other regions will be active in a functional MRI, given that the region under the crosshairs is active.

We will click on the "Association" link that appears above the brain map. The resulting web page gives a table listing the names of other Neurosynth meta-analysis maps (that already exist, somewhere on the Neurosynth web server) that have especially high values at the coordinates we entered.

Data for coordinates:

Neurosynth.org (beta) Home Meta-analyses - Studies Locations Decoder Code FAQs Sign in

Maps Studies **Associations** FAQs

Associations with meta-analysis maps

Show entries Search:

Name	Individual voxel		Seed-based network	
	z-score	Posterior prob.	Func. conn. (r)	Meta-analytic coact. (r)
working	6.73	0.74	0.43	0.54
working memory	6.7	0.74	0.44	0.55
load	5.55	0.77	0.32	0.44
memory wm	5.21	0.78	0.31	0.37
wm task	4.92	0.83	0.15	0.16
parietal	4.76	0.65	0.47	0.53
memory	4.75	0.66	0.28	0.29
demands	4.67	0.72	0.2	0.31
prefrontal	4.47	0.64	0.12	0.18
early stages	4.45	0.84	0	0

Showing 1 to 10 of 3,107 entries

First Previous **1** 2 3 4 5 ... 311 Next Last

Click the "Associations" link to see this list of terms associated with your coordinates.

We can use the names of these meta-analyses as a set of rough guesses about what other cognitive skills might be associated with this brain location. Again, the activation maps from Neurosynth meta-analyses are interesting because they represent pretty good answers to questions like, "If all I know about a neuroimaging study is that the term X appeared in the article, which brain regions were likely to be active?" Right now we are considering something like the reverse question: "If all I know about a neuroimaging study is that coordinate X,Y,Z was active, what terms were likely to appear in the article?" It requires an additional leap of faith to guess that just because an article included the term "language comprehension" (for example), that our brain region is actively involved in producing that cognitive skill, but it's a great start.

Copy the table from the "Associations" page and paste it at the bottom of your answer.

Item 3:

Paste the table that lists the first 10 Neurosynth "Associations" for the brain region from Item 1

Again, this table should have only 10 rows.

What else might this region do?

Item 4:

Based on the names of the associated Neurosynth meta-analyses, what one other cognitive skill do you think your brain region might be involved in?

- 1. Give the name of the skill
- 2. Explain why you concluded that.

By default, Neurosynth will display only 10 entries in this table, sorted in descending order by z-score. That is enough. There is no need to examine or submit more than that.

Data for coordinates: 28 8 48

Neurosynth.org (beta) Home Meta-analyses Studies Locations Decoder Code FAQs Sign in

Maps Studies **Associations** FAQs

Associations with meta-analysis maps

Show 10 entries Search: [] [] []

Name	Individual voxel		Seed-based network	
	z-score	Posterior prob.	Func. conn. (r)	Meta-analytic coact. (r)
working	6.73	0.74	0.43	0.54
working memory	6.7	0.74	0.44	0.55
load	5.55	0.77	0.32	0.44
memory w/m	5.21	0.78	0.31	0.37
w/m task	4.92	0.83	0.15	0.16
parietal	4.76	0.65	0.47	0.53
memory	4.75	0.66	0.28	0.29
demands	4.67	0.72	0.2	0.31
prefrontal	4.47	0.64	0.12	0.18
early stages	4.45	0.84	0	0

Showing 1 to 10 of 3,107 entries

First Previous 1 2 3 4 5 ... 311 Next Last

Select the rows of the "Associations" table and copy the text.

In a few cases this will be a very straightforward answer to give. Many of the existing Neurosynth meta-analyses are based on terms that describe methodological details or non-cognitive aspects of neuroimaging studies, e.g. “dti” (for diffusion tensor imaging), “multivariate,” or “able” (too generic to be a cognitive skill). It's possible that your coordinates will be associated with a list that includes only one cognitive skill. In that case, you can just name it, and the only explanation you would need to give would be “this was the only cognitive skill on the list.”

In most cases you will have to use your judgement to decide on the answer. For example, if not one but several different cognitive skills appear in the list, you will have to pick one. If several different skills all have something in common (e.g. “language comprehension,” “speech production,” “syntax), the right approach would be to give the name of that common topic as your answer (“language” seems good). For your explanation, just state what you did to arrive at that decision.

A final possibility is that there won't be **any** cognitive skills on your list. In that case, the appropriate answer is “none.” (You could examine more than the first 10 associations, but I just said that we don't need to do that.) For your explanation, just state that none of the meta-analyses on the list corresponded to a cognitive skill. (You will still need to copy and paste the table).

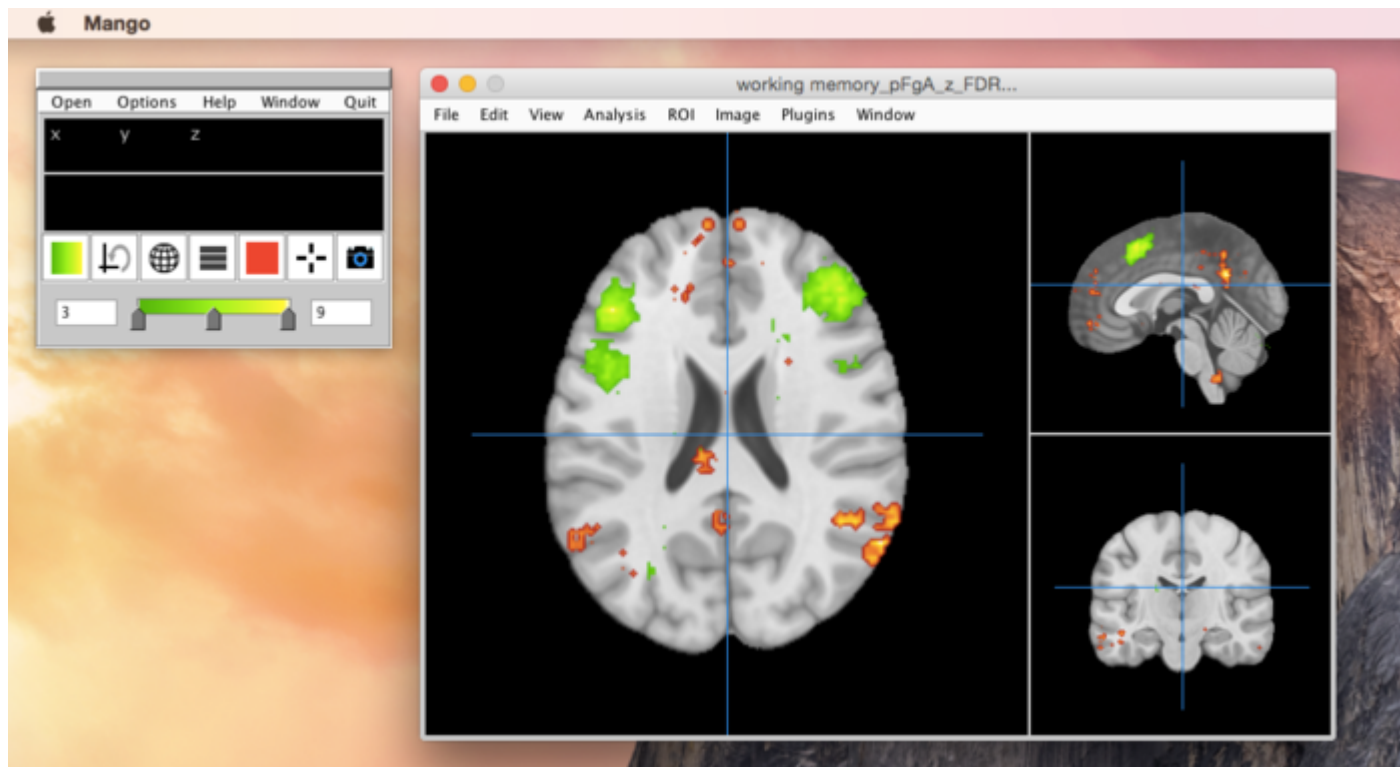
2. Compare the Neurosynth brain map for your "other" skill

2.1 Add your "other" brain image as a second overlay

Just to check in, this part of the instructions assumes that

1. Mango is running
 - The atlas feature is enabled
 - The Neurosynth functional map overlay for your article's cognitive skill is still open

Good. Now, open your second Neurosynth map (the one corresponding to your “other” cognitive skill). Mango allows you to have multiple overlays open at once, and it will even automatically assign each one a different color. The second one will be greenish.



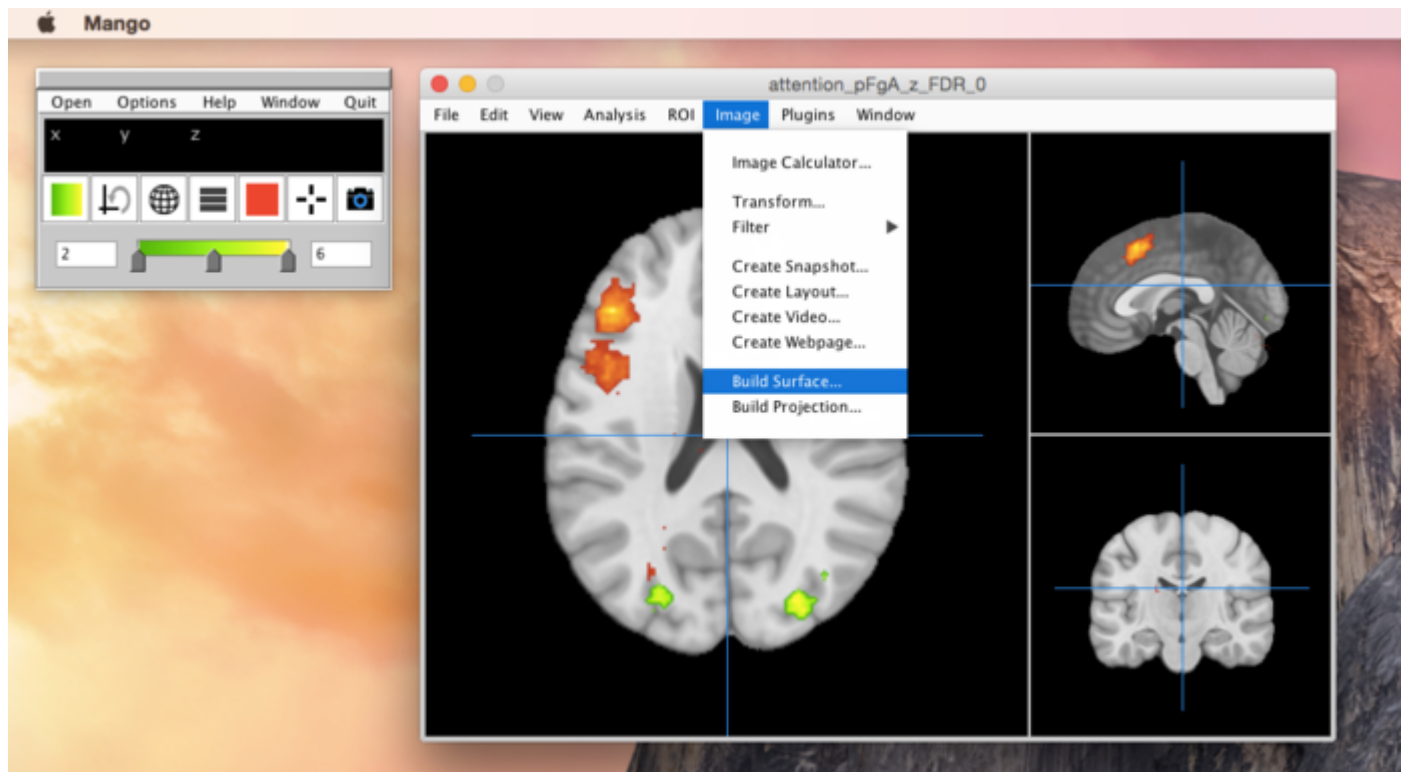
Open your second Neurosynth map as an overlay.

If you hate green, fear not. You can change any overlay's color (and tweak other things as well) by clicking on the overlay icon in the main (smaller) Mango window. The icon at the left of the row of icons below the coordinates display, and looks like a little square with a color gradient. Clicking on the icon once will bring up a drop-down list with icons for both of your current overlays and an icon for the structural "image" (in gray). Hovering the mouse over one of these icons will bring up a clickable menu.



The overlay menu expanded to show icons for the two overlays and the one structural “image.”

2.2 Make a 3D surface model



Select Image → Build surface to make a 3D surface model

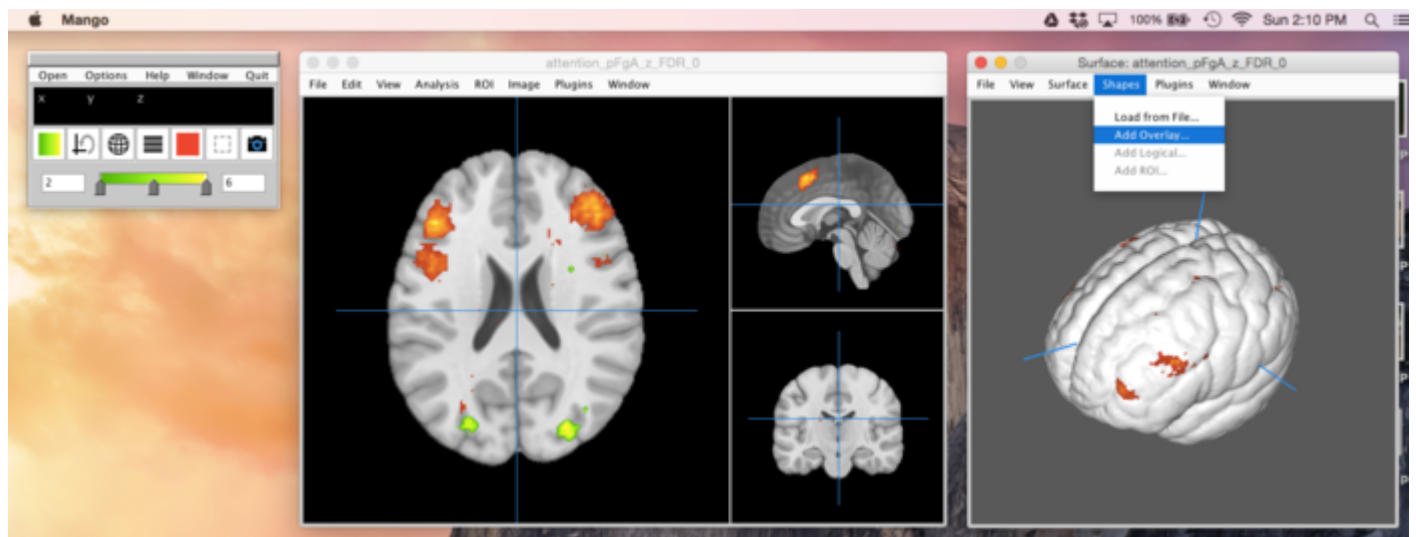


Click "OK" to accept the default options.



Behold! A brain you can spin around by clicking and dragging with the mouse.

By default, only the first overlay that you opened will be visible on the 3D surface model. Add the second overlay using the “Shapes” menu in the surface window.



Select “Shapes → Add Overlay” to add the second Neurosynth map overlay.

A dialog window will appear, asking you which overlay to add. Your second overlay (the green one) should be listed by default, so click “Add.”



Click "Add" to add your second overlay to the 3D surface model.

The only activation blobs that will be visible on the 3D surface model are those that actually extend all the way to the outside of the average structural brain image. However, we can see any blobs we want by enabling "cut planes" in the 3D surface window. A "cut plane" is a noun, short for "cutTING plane." It is like an invisible knife that cuts through the brain in one direction and removes all the brain stuff on the side facing us, the viewers seated at our computers.

The "All Cut Planes" option sets up a set of three "cut planes" that intersect at the point marked by the crosshairs.



Select "Surface" → "All Cut Planes"

This is how the 3D surface model should look after selecting the “All Cut Planes” menu item.



Congratulations! You now have a hole in your head.

The locations of the “cut planes” are linked to the crosshairs in the viewing window. To reveal different parts of the 3D surface model's interior, click and drag the mouse around in the *viewing window* (and not in the surface window).



The 3D surface model cut away to reveal the location marked by the crosshairs in the viewing window.

Take a screenshot of the 3D surface model

Item 5:

Screenshot of the 3D surface model, with blobs from both Neurosynth maps revealed by the "cut planes".

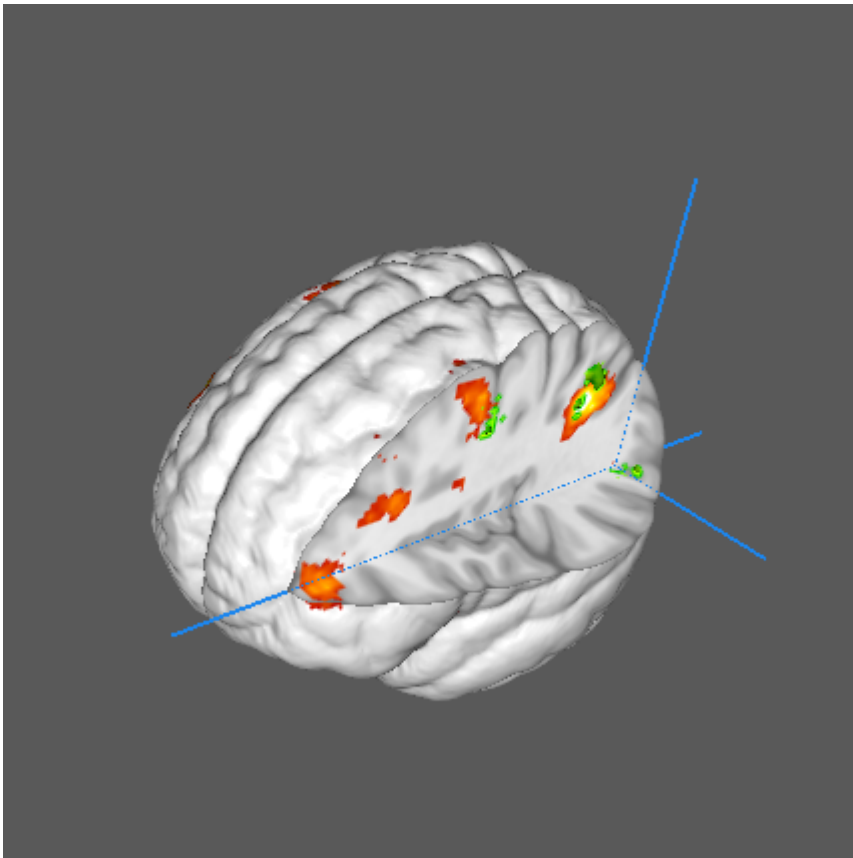
Move the "cut planes" around until you have revealed a nice view of the 3D surface model's interior, so that blobs from both overlays are showing. That is, both orange/red and green/yellow blobs should be visible.

Submit a screenshot of your awesome, holey brain by doing this:

- Take a screenshot of the surface window.
- Rename the file something similar to "[name of main cognitive skill] and [name of "other" skill] surface model.png"
- Upload it to the appropriate place on Canvas.

NOTE: You must have the Mango **surface window** "highlighted" or "in focus" (by clicking on it) before you use the main window's "camera" screenshot icon. Only the "highlighted" window's contents will be saved to the screenshot.

It should look similar to this:



An example screenshot of the surface window's contents

COMPLETELY NON-CREDIT OPTIONAL BONUS ACTIVITY:

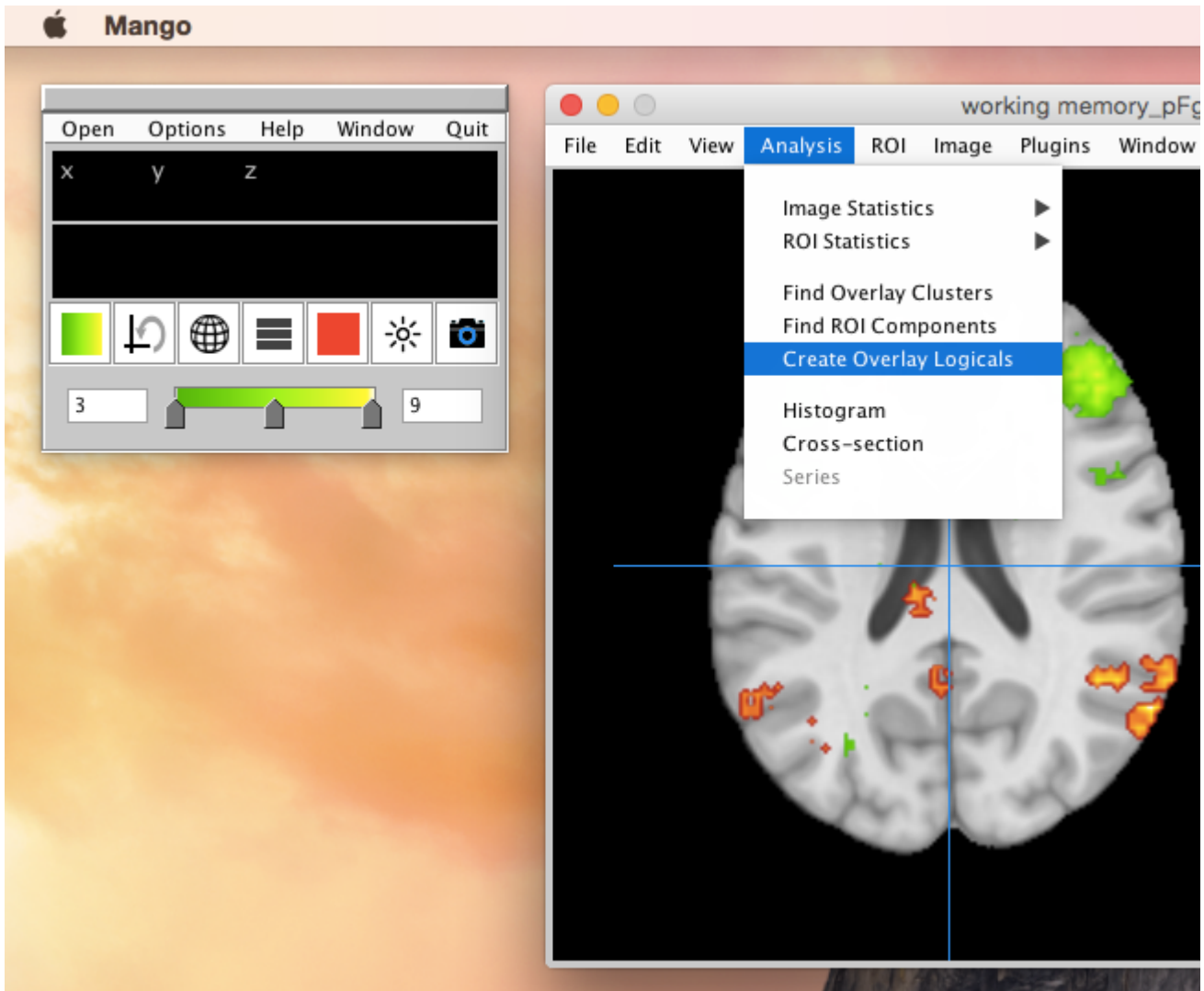
You can explore the “Surface” menu to create a video of a spinning brain. The world is your oyster now.

2.3 Find brain regions activated by both cognitive skills

Scroll/click/hover around the brain to get a sense of how the Neurosynth activation map for your “other” skill differs from the map for your primary cognitive skill. In most cases, the two maps will overlap in many places, but not always. Think about the similarities and differences between the two cognitive skills, and whether these are reflected by the degree to which their functional maps overlap.

Now, let's do a more precise investigation of the locations where the two maps overlap. We can use Mango to define a “region of interest” (ROI) that marks all of the overlapping locations.

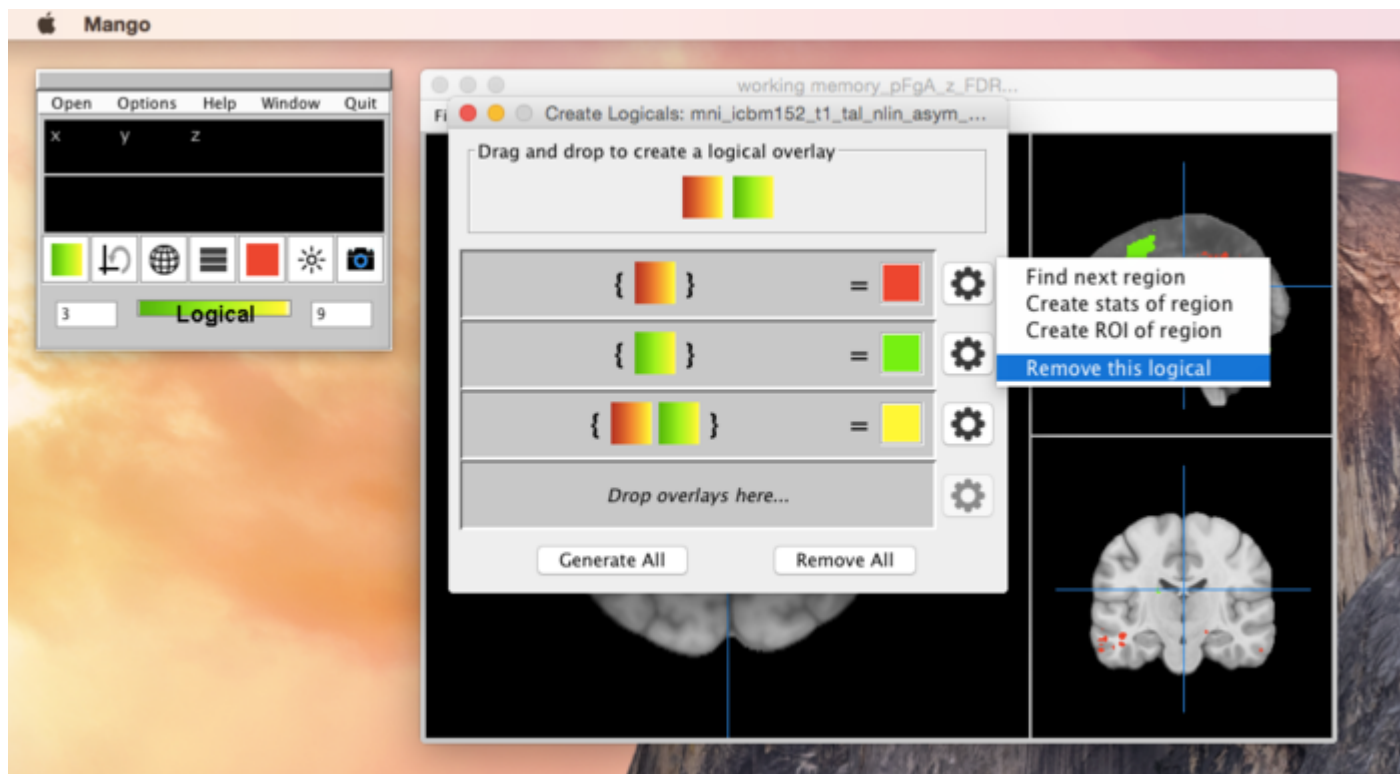
In the Mango viewing window, select the “Analysis → Create Overlay Logicals” menu item.



Select the “Create Overlay Logicals” menu item to make a third overlay.

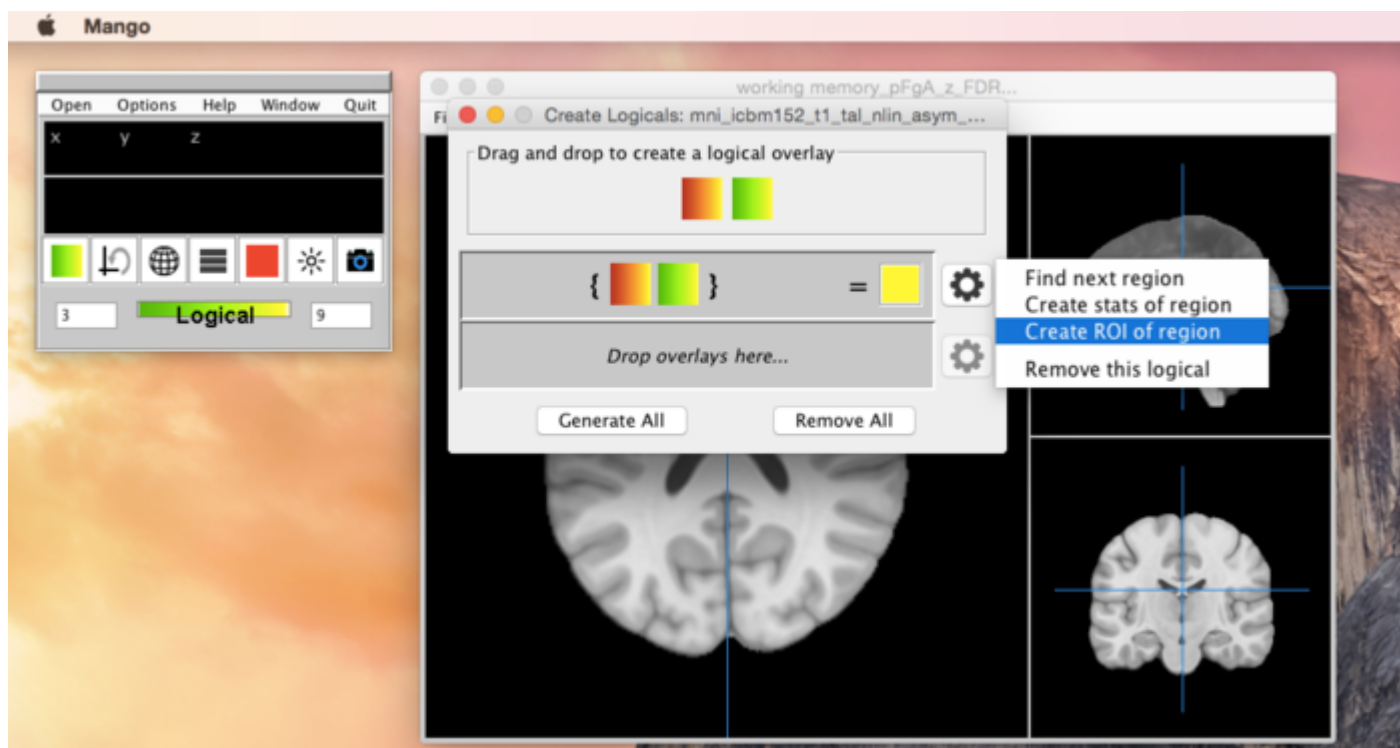
This will open a “Create Logicals” dialog window. When Mango currently has two overlays open (as we do now), it will list three possible ways for us to define a set of locations based on the locations covered in the overlays. The first two ways are trivial: they just consist of selecting all the locations covered by one of the overlays. The third way, even though it's not clearly explained in the dialog window, is to select all the locations covered by both of our overlays; this is the one we want.

To make this work correctly, it is best to remove the first two rows in this list. Click on the “gear” icon to the right of these rows and select “Remove this logical.”



Remove the first two rows from the “Create Logicals” dialog window.

Now, create a new overlay by clicking the “gear” icon to the right of the yellow square, and select “Create ROI of region.”



Select “Create ROI of region”

When this is done, you can close the “Create Logicals” window.

ROI stands for “Region of Interest” in neuroimaging. ROI is a vague term. It is mainly used when referring to some set of locations that are smaller than the entire volume of the brain.

The new ROI will be drawn on top of the structural brain image, although it can be hard to see. Right now it will appear as a set of thin red lines that circumscribe the ROI's locations.

Instead of scrolling/clicking/hovering around to examine the hard-to-see ROI lines, we will ask Mango to make us a nice table listing all the locations included in the ROI. Mango will helpfully group all of these locations into several discrete “components” that consist of contiguous points. Mango uses the word “component” to describe what most neuroimagers call “clusters.”

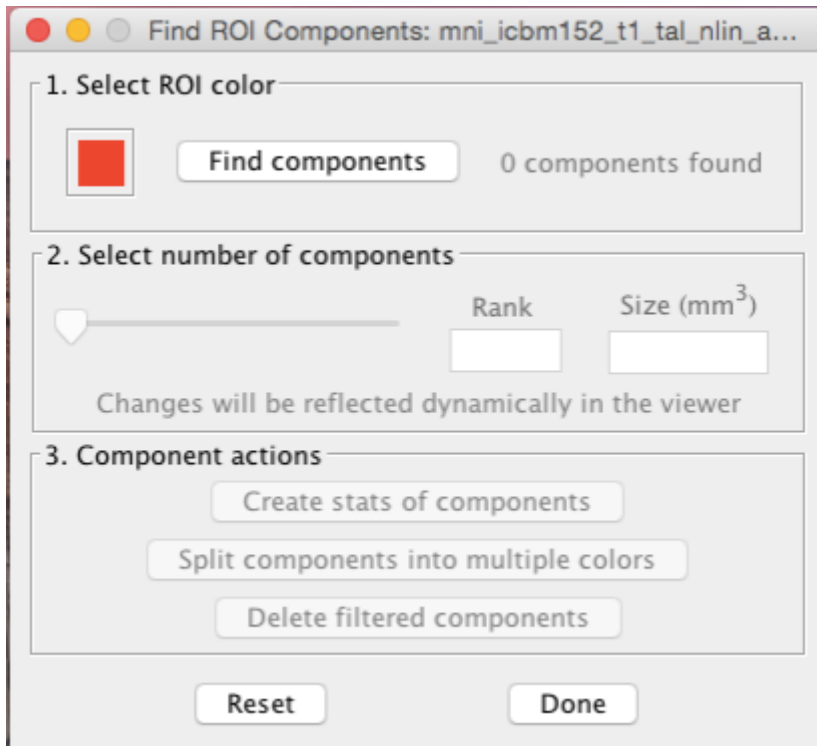
I prefer the word “clusters.” It is more tangible, and it makes me think of candy. When I use the word “cluster” below, I am referring to the same thing as a Mango “component.”

In the Mango viewing window, select the “Analysis → Find ROI Components” menu item.



Select "Find ROI Components" to generate a table of the ROI's locations.

This will open up the "Find ROI Components" dialog window:



The “Find ROI Components” dialog window.

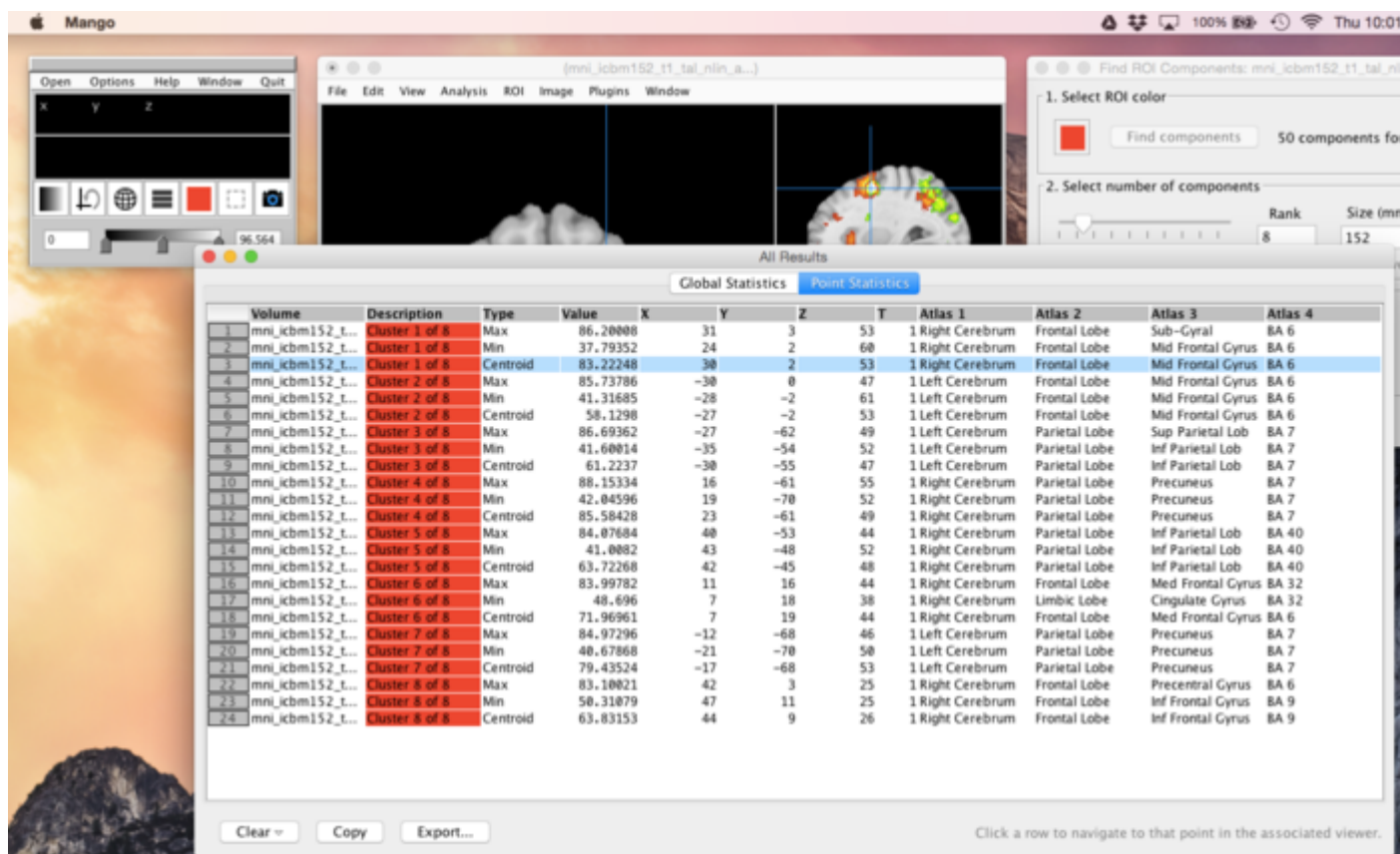
Click the “Find components” button. After message to the right of this button changes from “0 components found” to “[some number greater than zero] components found,” click on the “Create stats of components” button in the “3. Component actions” section of the dialog window. You do not need to adjust the number of components slider in section 2 of the dialog window.

This will open up a table in a window entitled “All Results.”

Image	Description	Mean	Sum	SD	Size	Count
1	working memory_pFgA_z_FDR_0 Cluster 1 of 8	6.35502	13993.7	1.68852	2202	2202
2	working memory_pFgA_z_FDR_0 Cluster 2 of 8	5.66482	7143.33	1.17689	1261	1261
3	working memory_pFgA_z_FDR_0 Cluster 3 of 8	6.04663	6538.36	1.5947	1080	1080
4	working memory_pFgA_z_FDR_0 Cluster 4 of 8	4.67784	4495.40	0.86953	961	961
5	working memory_pFgA_z_FDR_0 Cluster 5 of 8	6.08738	5521.25	1.43254	907	907
6	working memory_pFgA_z_FDR_0 Cluster 6 of 8	5.09552	2817.82	0.75816	553	553
7	working memory_pFgA_z_FDR_0 Cluster 7 of 8	4.56182	2262.66	0.86582	496	496
8	working memory_pFgA_z_FDR_0 Cluster 8 of 8	3.9834	605.476	0.5391	152	152

The ROI stats appear in the “All Results” window.

The “All Results” window displays two different tables. At first, the “Global Statistics” table will probably be visible. This is not the one we want. Click the “Point Statistics” button to switch to that view.



The screenshot shows the Mango software interface. The main window displays a brain map with colored clusters. The 'All Results' window is open, showing two tabs: 'Global Statistics' and 'Point Statistics'. The 'Point Statistics' tab is active, displaying a table with 24 rows of data. The table columns are: Volume, Description, Type, Value, X, Y, Z, T, Atlas 1, Atlas 2, Atlas 3, and Atlas 4. The data is organized into 8 clusters, with each cluster having three rows for Max, Min, and Centroid values.

Volume	Description	Type	Value	X	Y	Z	T	Atlas 1	Atlas 2	Atlas 3	Atlas 4
mni_icbm152_t...	Cluster 1 of 8	Max	86.20008	31	3	53	1	Right Cerebrum	Frontal Lobe	Sub-Gyrus	BA 6
mni_icbm152_t...	Cluster 1 of 8	Min	37.79352	24	2	60	1	Right Cerebrum	Frontal Lobe	Mid Frontal Gyrus	BA 6
mni_icbm152_t...	Cluster 1 of 8	Centroid	83.22248	30	2	53	1	Right Cerebrum	Frontal Lobe	Mid Frontal Gyrus	BA 6
mni_icbm152_t...	Cluster 2 of 8	Max	85.73786	-30	0	47	1	Left Cerebrum	Frontal Lobe	Mid Frontal Gyrus	BA 6
mni_icbm152_t...	Cluster 2 of 8	Min	41.31685	-28	-2	61	1	Left Cerebrum	Frontal Lobe	Mid Frontal Gyrus	BA 6
mni_icbm152_t...	Cluster 2 of 8	Centroid	58.1298	-27	-2	53	1	Left Cerebrum	Frontal Lobe	Mid Frontal Gyrus	BA 6
mni_icbm152_t...	Cluster 3 of 8	Max	86.69362	-27	-62	49	1	Left Cerebrum	Parietal Lobe	Sup Parietal Lob	BA 7
mni_icbm152_t...	Cluster 3 of 8	Min	41.60014	-35	-54	52	1	Left Cerebrum	Parietal Lobe	Inf Parietal Lob	BA 7
mni_icbm152_t...	Cluster 3 of 8	Centroid	61.2237	-30	-55	47	1	Left Cerebrum	Parietal Lobe	Inf Parietal Lob	BA 7
mni_icbm152_t...	Cluster 4 of 8	Max	88.15334	16	-61	55	1	Right Cerebrum	Parietal Lobe	Precuneus	BA 7
mni_icbm152_t...	Cluster 4 of 8	Min	42.04596	19	-70	52	1	Right Cerebrum	Parietal Lobe	Precuneus	BA 7
mni_icbm152_t...	Cluster 4 of 8	Centroid	85.58428	23	-61	49	1	Right Cerebrum	Parietal Lobe	Precuneus	BA 7
mni_icbm152_t...	Cluster 5 of 8	Max	84.07684	40	-53	44	1	Right Cerebrum	Parietal Lobe	Inf Parietal Lob	BA 40
mni_icbm152_t...	Cluster 5 of 8	Min	41.0082	43	-48	52	1	Right Cerebrum	Parietal Lobe	Inf Parietal Lob	BA 40
mni_icbm152_t...	Cluster 5 of 8	Centroid	63.72268	42	-45	48	1	Right Cerebrum	Parietal Lobe	Inf Parietal Lob	BA 40
mni_icbm152_t...	Cluster 6 of 8	Max	83.99782	11	16	44	1	Right Cerebrum	Frontal Lobe	Med Frontal Gyrus	BA 32
mni_icbm152_t...	Cluster 6 of 8	Min	48.696	7	18	38	1	Right Cerebrum	Limbic Lobe	Cingulate Gyrus	BA 32
mni_icbm152_t...	Cluster 6 of 8	Centroid	71.96961	7	19	44	1	Right Cerebrum	Frontal Lobe	Med Frontal Gyrus	BA 6
mni_icbm152_t...	Cluster 7 of 8	Max	84.97296	-12	-68	46	1	Left Cerebrum	Parietal Lobe	Precuneus	BA 7
mni_icbm152_t...	Cluster 7 of 8	Min	40.67868	-21	-70	50	1	Left Cerebrum	Parietal Lobe	Precuneus	BA 7
mni_icbm152_t...	Cluster 7 of 8	Centroid	79.43524	-17	-68	53	1	Left Cerebrum	Parietal Lobe	Precuneus	BA 7
mni_icbm152_t...	Cluster 8 of 8	Max	83.10021	42	3	25	1	Right Cerebrum	Frontal Lobe	Precentral Gyrus	BA 6
mni_icbm152_t...	Cluster 8 of 8	Min	50.31079	47	11	25	1	Right Cerebrum	Frontal Lobe	Inf Frontal Gyrus	BA 9
mni_icbm152_t...	Cluster 8 of 8	Centroid	63.83153	44	9	26	1	Right Cerebrum	Frontal Lobe	Inf Frontal Gyrus	BA 9

Switch to the “Point Statistics” table in the “All Results” window.

The “Point Statistic” table lets us do two helpful things:

First, it gives the name of the brain region where each cluster is located. That's right! No need to scroll/click/hover²! Simply read the names in the column titled “Atlas 3” to see a more or less complete list of the brain regions that the ROI clusters occupy. The “Atlas 2” column names the larger structure (usually a lobe of the cerebral cortex) that this region is part of. The “Atlas 1” column might look like it is too general to be useful (as in, “Thanks a lot, Mango, I know that the frontal lobe is part of the cerebrum! Sheesh. For real.”), but it also specifies which hemisphere (left or right) the cluster is in. You could figure that out by looking at the brain map, but sometimes it's easier just to read this from a table.

Second, the table is integrated with the Mango viewing window, so that when you click on a row Mango will recenter the views of the brain with the crosshairs located on that point. Mango lists three types of points for each cluster: the locations of the minimum and maximum activation values, and the centroid of the cluster. Sometimes all three of these can occupy different brain regions. The centroid is the most useful one for our purposes, because it corresponds to the center of mass or “middle” of the cluster. It is the most representative point in the cluster.

Click on rows for the different clusters, and look at the clusters in viewing window. By default, the clusters should be listed in order from biggest (occupying the most space) to smallest in the table. A large cluster does not necessarily mark the location of the most crucial brain activity, though.

Pay special attention to whether a given cluster occurs only in one hemisphere (left or right), or whether it is part of a pair of clusters (one in each hemisphere, in more or less the same brain region).

Choose a most important single cluster and take a screenshot

Item 6:

Screenshot of one ROI cluster from the intersection of both Neurosynth maps

After browsing around the brain maps and thinking for a while, decide which **single cluster** you think is a good guess for “the most important brain region that is active during both of these cognitive skills.”

- Take a screenshot of the Mango viewing window.
- Rename the file something like “most important cluster”
- Upload the file to the appropriate place on Canvas.

Explain why you chose this cluster

Item 7:

Where in the brain is this cluster, and why did you choose it?

- Use the “centroid” type when you look for the name of the brain region that a cluster occupies.
- Use the “Atlas 3” column to give the name of the brain region (e.g. “Precuneus” and not “Parietal lobe”).
- If the cluster is one of a pair of bilateral clusters (i.e. one in each hemisphere), then you can name the brain region for the cluster as “bilateral X.”
- You will need to explain how you made your choice. There is no clear right or wrong answer, and it is OK if the reasons for your choice don't seem especially scientific.

For our final exercise in this assignment, look up the coordinates for your cluster in Neurosynth's “Locations” search.

Copy the first 10 Neurosynth associations for your cluster's centroid coordinates

Item 8:

Paste the table that lists the first 10 Neurosynth "Associations" for this cluster's coordinates

- Copy down the 3D stereotactic coordinates (X,Y,Z) for your cluster.
- Repeat the instructions above ([jump to that section](#)) to view the list of the first 10 Neurosynth associations for those coordinates.

Copy and paste the list as your answer.

What does this brain region do?

Item 9:

What do you think this cluster's brain region does?

If (and only if) the first 10 associations returned by Neurosynth include **zero cognitive skills**, you may answer "Hard to say." In fact, it is *always* hard to say what a region of the brain "does." Don't let that stop you from making a good guess. Explain why you made your guess, even if your explanation sounds painfully straightforward (e.g. "I guessed 'language comprehension' because that was the only cognitive skill that appeared in the list of associations."). Your instructor appreciates the benefits of "painfully straightforward."

Footnotes

1)

Both "free as in speech" and "free as in beer." https://en.wikipedia.org/wiki/Gratis_versus_libre

2)

You might be asking yourself, "Wait, why did we have to 'scroll/click/hover' for the earlier part, then?" Answer: Because scroll/click/hovering is good for your soul. It is easier to receive a list of brain regions like the one in the "Point Statistics" table, but that won't help you to get an understanding of where these regions are in the brain, what they look like, and how close they are to each other. For that, you need to practice navigating the spatial layout of the brain.

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