

## Background

Precision functional mapping (PFM) refers to a suite of approaches for delineating functional brain areas and networks at the individual level.

## Example PFM dataset

An example PFM dataset (“ME01”, study author CJL) is available online at OpenNeuro.org (<https://openneuro.org/datasets/ds005118/>) as example data for this tutorial. Please cite Lynch et al., 2024 (citation pending) when using this code, and <sup>1</sup> when publishing any work that includes the ME01 dataset.

## Usage

The goal of this tutorial is to describe step-by-step how PFM is performed in our lab. This tutorial focuses on the ME01 dataset, but in principle the code can be adapted to accommodate any fMRI dataset that has been mapped to `fs_LR_32k` surface space (CIFTI format, “.dtseries.nii” file with 64k vertices, 32k vertices per hemisphere when including medial wall). For more information regarding the CIFTI format, see Glasser et al. 2013 Neuroimage (<https://pubmed.ncbi.nlm.nih.gov/23668970/>).

If you have issues, please email me (Chuck Lynch; [cjl2007@med.cornell.edu](mailto:cjl2007@med.cornell.edu)), and I will try my best to help you.

## Before you begin.

Check that all of the necessary dependencies are available.

Start Matlab, open the `pfm_tutorial.m` script in the Matlab editor window, and run lines 5-6 to add all of the necessary dependencies to the Matlab search path.

```
3 % Before you begin.
4
5 % add dependencies to Matlab search path
6 - addpath(genpath([pwd '/PFM-Tutorial/Utilities']));
7
8 % define path to some software packages that will be needed
9 - InfoMapBinary = '/home/charleslynch/miniconda3/bin/infomap'; % path to infomap binary; code tested on version 2.0.0
10 - WorkbenchBinary = '/usr/local/workbench/bin_linux64/wb_command'; % path to workbench binary; code tested on version 1
11
12 % number of
13 % workers
14 - nWorkers = 5;
```

You will need to install Infomap (<https://www.mapequation.org/infomap/#Install>) and Connectome Workbench (<https://www.humanconnectome.org/software/connectome-workbench>). Once installed, run lines 8-10 to define two (string) variables called

`InfomapBinary` and `WorkbenchBinary`, representing the paths to the binary executables for each of these programs.

```
3 % Before you begin.
4
5 % add dependencies to Matlab search path
6 - addpath(genpath([pwd '/PFM-Tutorial/Utilities']));
7
8 % define path to some software packages that will be needed
9 - InfoMapBinary = '/home/charleslynch/miniconda3/bin/infomap'; % path to infomap binary; code tested on version 2.0.0
10 - WorkbenchBinary = '/usr/local/workbench/bin_linux64/wb_command'; % path to workbench binary; code tested on version 1
11
12 % number of
13 % workers
14 - nWorkers = 5;
```

Next, run lines 12-14 to define the number of workers for parallelization of certain procedures. The appropriate number will depend on your particular computing environment.

```
3 % Before you begin.
4
5 % add dependencies to Matlab search path
6 - addpath(genpath([pwd '/PFM-Tutorial/Utilities']));
7
8 % define path to some software packages that will be needed
9 - InfoMapBinary = '/home/charleslynch/miniconda3/bin/infomap'; % path to infomap binary; code tested on version 2.0.0
10 - WorkbenchBinary = '/usr/local/workbench/bin_linux64/wb_command'; % path to workbench binary; code tested on version 1
11
12 % number of
13 % workers
14 - nWorkers = 5;
```

Finally, the example dataset can be obtained from OpenNeuro.org (<https://openneuro.org/datasets/ds005118/versions/1.0.0/download>). There are multiple ways to download the data — if using your browser to download the data is unsuccessful, try an alternative method like the AWS CLI. Please note that this is a large download (~100 GB).

## Step 1: Temporal Concatenation of fMRI data from all sessions.

The first step is to temporally concatenate all the individual denoised and fs\_LR\_32k surface-registered CIFTI (“.dtseries.nii”) files to obtain a single CIFTI file for the subsequent analysis.

For context, the validity and test-retest reliability of functional connectivity measurements and their derivatives (functional network parcellations) at the single-subject level increases rapidly with additional data. This is why PFM is often performed in individuals that have been scanned repeatedly over an extended period of time, for total scan durations of several hours or more. For example, ME01 was scanned 44 times over 10 imaging sessions (> 10 hours of functional MRI data total). In principle, however, PFM may be performed with far less data when specialized acquisitions are used to improve signal-to-noise.

In the `pfm_tutorial.m` script, run lines 18-20 to define the path to the subject’s folder and identifier.

```

16 %% Step 1: Temporal Concatenation of fMRI data from all sessions.
17
18 % define subject directory and name;
19 Subdir = [pwd '/WCM-ME/derivatives/sub-ME01/'];
20 Subject = 'ME01';
21
22 % define & create
23 % the pfm directory;
24 PfmDir = [Subdir '/pfm/'];
25 mkdir(PfmDir);

```

Next, run lines 22-25 to define and create the folder (`PfmDir`) where all the pfm related outputs will be stored. Note that for convenience, the dataset is distributed online with this folder and all outputs pre-generated. You can delete `PfmDir` before proceeding to start from scratch.

```

16 %% Step 1: Temporal Concatenation of fMRI data from all sessions.
17
18 % define subject directory and name;
19 Subdir = [pwd '/WCM-ME/derivatives/sub-ME01/'];
20 Subject = 'ME01';
21
22 % define & create
23 % the pfm directory;
24 PfmDir = [Subdir '/pfm/'];
25 mkdir(PfmDir);

```

Next, run lines 27-53 to iteratively load each of the denoised and fs\_LR\_32k surface-registered CIFTI (“.dtseries.nii”) files. The resting-state time courses will be extracted from the `.data` field of each CIFTI file, and demeaned and motion-censored (time points with framewise displacement > 0.3 mm are discarded) before being temporally concatenated into the `ConcatenatedData` variable.

```

27 % count the number of imaging sessions;
28 nSessions = length(dir([Subdir '/processed_restingstate_timecourses/ses-func*']));
29
30 % preallocate;
31 ConcatenatedData = [];
32
33 % sweep through
34 % the sessions;
35 for i = 1:nSessions
36
37 % count the number of runs in this session
38 nRuns = length(dir([Subdir '/processed_restingstate_timecourses/ses-func' sprintf('%02d',i) '/run-*'.dtseries.nii']));
39
40 % sweep
41 % through
42 % the runs;
43 for ii = 1:nRuns
44
45 % load the denoised & fs_lr_32k surface-registered CIFTI file for run "ii" from session "i"...
46 Cifti = ft_read_cifti_mod([Subdir '/processed_restingstate_timecourses/ses-func' sprintf('%02d',i) '/sub-' Subject '_ses-' sprintf('%02d',ii) '.dtseries.nii']);
47 Cifti.data = Cifti.data - mean(Cifti.data,2); % demean
48 Tmask = load([Subdir '/processed_restingstate_timecourses/ses-func' sprintf('%02d',i) '/sub-' Subject '_ses-' sprintf('%02d',ii) '_Tmask.mat']); % 1 (Low motion timepoints) == FD < 0.3mm, 0 (High motion)
49 ConcatenatedData = [ConcatenatedData Cifti.data(:,Tmask==1)]; % 1 (Low motion timepoints) == FD < 0.3mm, 0 (High motion)
50
51 end
52
53 end
54
55 % make a single CIFTI containing
56 % time-series from all scans;
57 ConcatenatedCifti = Cifti;
58 ConcatenatedCifti.data = ConcatenatedData;
59

```

Finally, run lines 55-58 to create a single CIFTI file (`ConcatenatedCifti`) that contains all available resting-state data for this subject.

```

27 % count the number of imaging sessions;
28 nSessions = length(dir([Subdir '/processed_restingstate_timecourses/ses-func*']));
29
30 % preallocate;
31 ConcatenatedData = [];
32
33 % sweep through
34 % the sessions;
35 for i = 1:nSessions
36
37 % count the number of runs in this session
38 nRuns = length(dir([Subdir '/processed_restingstate_timecourses/ses-func' sprintf('%02d',i) '/run-*_.dtseries.nii']));
39
40 % sweep
41 % through
42 % the runs;
43 for ii = 1:nRuns
44
45 % load the denoised & fs_lr_32k surface-registered CIFTI file for run "ii" from session "i"...
46 Cifti = ft_read_cifti_mod([Subdir '/processed_restingstate_timecourses/ses-func' sprintf('%02d',i) '/sub-' Subject '_ses-' sprintf('%02d',i) '_run-' sprintf('%02d',ii) '_dtseries.nii']);
47 Cifti.data = Cifti.data - mean(Cifti.data,2); % demean
48 Tmask = load([Subdir '/processed_restingstate_timecourses/ses-func' sprintf('%02d',i) '/sub-' Subject '_ses-' sprintf('%02d',i) '_run-' sprintf('%02d',ii) '_Tmask.npz']);
49 ConcatenatedData = [ConcatenatedData Cifti.data(:,Tmask==1)]; % 1 (Low motion timepoints) == FD < 0.3mm, 0 (high motion)
50
51 end
52
53 end
54
55 % make a single CIFTI containing
56 % time-series from all scans;
57 ConcatenatedCifti = Cifti;
58 ConcatenatedCifti.data = ConcatenatedData;
59

```

## Step 2: Create distance matrix & regress nearby cortical signals from subcortical structures.

The next step is to create a matrix summarizing the distance between all points in the brain. Geodesic and Euclidean space is used for cortico-cortical (vertex-to-vertex) and subcortical-cortical distance (voxel-to-vertex), respectively.

In the pfm\_tutorial.m script in the Matlab editor window, and run lines 62-67 to define the path the the subject's fs\_LR\_32k midthickness (the midpoint between the white and pial surfaces) and run the `pfm_make_dmat` function.

```

60 %% Step 2: Make a distance matrix.
61
62 % define fs_lr_32k midthickness surfaces;
63 MidthickSurfs{1} = [Subdir '/fs_LR/fsaverage_LR32k/' Subject '.L.midthickness.32k_fs_LR.surf.gii'];
64 MidthickSurfs{2} = [Subdir '/fs_LR/fsaverage_LR32k/' Subject '.R.midthickness.32k_fs_LR.surf.gii'];
65
66 % make the distance matrix;
67 pfm_make_dmat(ConcatenatedCifti, MidthickSurfs, PfmDir, nWorkers, WorkbenchBinary); %
68
69 % optional: regress adjacent cortical signal from subcortex to reduce artifactual coupling
70 % (for example, between cerebellum and visual cortex, or between putamen and insular cortex)
71 [ConcatenatedCifti] = pfm_regress_adjacent_cortex(ConcatenatedCifti, [PfmDir '/DistanceMatrix.mat'], 20);
72
73 % write out the CIFTI file;
74 ft_write_cifti_mod([Subdir '/pfm/sub-ME01_task-rest_concatenated_32k_fsLR.dtseries.nii'], ConcatenatedCifti);

```

For context, this distance matrix is used in multiple different ways during PFM. As one example, spurious coupling between subcortical voxels and adjacent cortical tissue (e.g., inflated FC between between occipital cortex and the cerebellum) can be mitigated by regressing the time-series of cortical tissue within a specified distance from any subcortical voxel<sup>2,3</sup>.

In the pfm\_tutorial.m script in the Matlab editor window, run lines 69-74 to run the `pfm_regress_adjacent_cortex` function and save the resultant CIFTI file.

```

60 %% Step 2: Make a distance matrix.
61
62 % define fs_lr_32k midthickness surfaces;
63 - MidthickSurfs{1} = [Subdir '/fs_LR/fsaverage_LR32k/' Subject '.L.midthickness.32k_fs_LR.surf.gii'];
64 - MidthickSurfs{2} = [Subdir '/fs_LR/fsaverage_LR32k/' Subject '.R.midthickness.32k_fs_LR.surf.gii'];
65
66 % make the distance matrix;
67 - pfm_make_dmat(ConcatenatedCifti, MidthickSurfs, PfmDir, nWorkers, WorkbenchBinary); %
68
69 % optional: regress adjacent cortical signal from subcortex to reduce artifactual coupling
70 % (for example, between cerebellum and visual cortex, or between putamen and insular cortex)
71 - [ConcatenatedCifti] = pfm_regress_adjacent_cortex(ConcatenatedCifti, [PfmDir '/DistanceMatrix.mat'], 20);
72
73 % write out the CIFTI file;
74 - ft_write_cifti_mod([Subdir '/pfm/sub-ME01_task-rest_concatenated_32k_fsLR.dtseries.nii'], ConcatenatedCifti);

```

### Step 3: Apply the desired amount of spatial smoothing.

In the Matlab editor window, specify a range of kernel sizes (in sigma) at line 80. The example range of sigma values specified below of 0.85, 1.7, and 2.55 correspond to a FWHM of 2mm, 4mm, and 6mm, respectively (FWHM  $\approx 2.355 * \text{sigma}$ ).

```

76 %% Step 3: Apply spatial smoothing.
77
78 % define a range of gaussian
79 % smoothing kernels (in sigma)
80 - KernelSizes = [0.85 1.7 2.55];
81
82 % sweep a range of
83 % smoothing kernels;
84 - for k = KernelSizes
85
86 % smooth with geodesic (for surface data) and Euclidean (for volumetric data) Gaussian kernels;
87 - system([WorkbenchBinary ' -cifti-smoothing ' PfmDir '/sub-ME01_task-rest_concatenated_32k_fsLR.dtseries.nii '...
88 num2str(k) ' ' num2str(k) ' COLUMN ' PfmDir '/sub-ME01_task-rest_concatenated_smoothed' num2str(k) '_32k_fsLR.dt...
89 end

```

Next, run lines 82-90 to apply the specified levels of spatial smoothing to the concatenated CIFTI file with geodesic (for cortical vertices) and Euclidean (for subcortical voxels) Gaussian kernels using Connectome Workbench command line utilities. Note that this step can be considered optional, but is recommended for most datasets.

```

76 %% Step 3: Apply spatial smoothing.
77
78 % define a range of gaussian
79 % smoothing kernels (in sigma)
80 - KernelSizes = [0.85 1.7 2.55];
81
82 % sweep a range of
83 % smoothing kernels;
84 - for k = KernelSizes
85
86 % smooth with geodesic (for surface data) and Euclidean (for volumetric data) Gaussian kernels;
87 - system([WorkbenchBinary ' -cifti-smoothing ' PfmDir '/sub-ME01_task-rest_concatenated_32k_fsLR.dtseries.nii '...
88 num2str(k) ' ' num2str(k) ' COLUMN ' PfmDir '/sub-ME01_task-rest_concatenated_smoothed' num2str(k) '_32k_fsLR.dt...
89 end

```

### Step 4: Run infomap.

The Infomap community detection algorithm (<https://www.mapequation.org/infomap/>) is one of the most widely used approaches for delineating functional brain networks and their boundaries in individuals. The `pfm_infomap` function is a wrapper that encompasses multiple steps — including creating and thresholding the functional connectivity (FC) matrix, calling the Infomap algorithm, and saving the resultant Infomap communities to a CIFTI file.

In the Matlab editor window, run line 95 to load the CIFTI file that you want to run Infomap on. In the example below, we selected the CIFTI file with 2.55 sigma spatial smoothing.

```

92 %% Step 4: Run infomap.
93
94 % load your concatenated resting-state dataset, pick whatever level of spatial smoothing you want
95 ConcatenatedCifti = ft_read_cifti_mod([PfmDir '/sub-ME01_task-rest_concatenated_smoothed2.55_32k_fsLR.dtseries.nii']);
96
97 % define inputs;
98 DistanceMatrix = [Subdir '/pfm/DistanceMatrix.mat']; % can be path to file
99 DistanceCutoff = 10; % in mm; usually between 10 to 30 mm works well.
100 GraphDensities = flip([0.0001 0.0002 0.0005 0.001 0.002 0.005 0.01 0.02 0.05]); %
101 NumberReps = 50; % number of times infomap is run;
102 BadVertices = []; % optional, but you could include regions to ignore, if you know there is bad signal there.
103 Structures = {'CORTEX_LEFT','CEREBELLUM_LEFT','ACCUMBENS_LEFT','CAUDATE_LEFT','PALLIDUM_LEFT','PUTAMEN_LEFT','THALAMI';
104
105 % run infomap
106 pfm_infomap(ConcatenatedCifti,DistanceMatrix,PfmDir,GraphDensities,NumberReps,DistanceCutoff,BadVertices,Structures);
107
108 % remove some intermediate files (optional)
109 system(['rm ' Subdir '/pfm/*.net']);
110 system(['rm ' Subdir '/pfm/*.clu']);
111 system(['rm ' Subdir '/pfm/*Log*']);
112
113 % define inputs;
114 Input = [PfmDir '/Bipartite_PhysicalCommunities.dtseries.nii'];
115 Output = 'Bipartite_PhysicalCommunities+SpatialFiltering.dtseries.nii';
116 MinSize = 50; % in mm^2
117
118 % perform spatial filtering
119 pfm_spatial_filtering(Input,PfmDir,Output,MidthickSurfs,MinSize,WorkbenchBinary);
120

```

Next, in the Matlab editor window, the following inputs must be defined by the user at lines 97-103 in the pfm\_tutorial.m script.

```

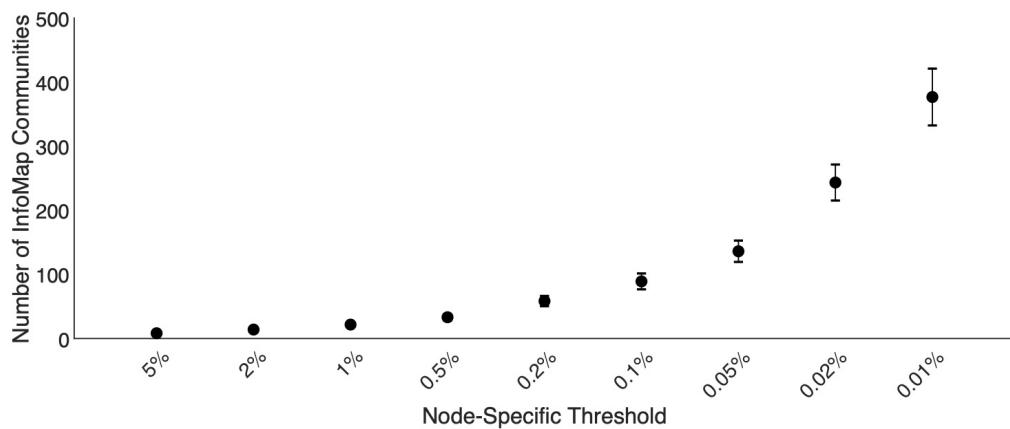
92 %% Step 4: Run infomap.
93
94 % load your concatenated resting-state dataset, pick whatever level of spatial smoothing you want
95 ConcatenatedCifti = ft_read_cifti_mod([PfmDir '/sub-ME01_task-rest_concatenated_smoothed2.55_32k_fsLR.dtseries.nii']);
96
97 % define inputs;
98 DistanceMatrix = [Subdir '/pfm/DistanceMatrix.mat']; % can be path to file
99 DistanceCutoff = 10; % in mm; usually between 10 to 30 mm works well.
100 GraphDensities = flip([0.0001 0.0002 0.0005 0.001 0.002 0.005 0.01 0.02 0.05]); %
101 NumberReps = 50; % number of times infomap is run;
102 BadVertices = []; % optional, but you could include regions to ignore, if you know there is bad signal there.
103 Structures = {'CORTEX LEFT','CEREBELLUM LEFT','ACCUMBENS LEFT','CAUDATE LEFT','PALLIDUM LEFT','PUTAMEN LEFT','THALAMUS LEFT'};
104
105 % run infomap
106 pfm_infomap(ConcatenatedCifti,DistanceMatrix,PfmDir,GraphDensities,NumberReps,DistanceCutoff,BadVertices,Structures);
107
108 % remove some intermediate files (optional)
109 system(['rm ' Subdir '/pfm/*.net']);
110 system(['rm ' Subdir '/pfm/*.clu']);
111 system(['rm ' Subdir '/pfm/*Log*']);
112
113 % define inputs;
114 Input = [PfmDir '/Bipartite_PhysicalCommunities.dtseries.nii'];
115 Output = 'Bipartite_PhysicalCommunities+SpatialFiltering.dtseries.nii';
116 MinSize = 50; % in mm^2
117
118 % perform spatial filtering
119 pfm_spatial_filtering(Input,PfmDir,Output,MidthickSurfs,MinSize,WorkbenchBinary);
120

```

`DistanceMatrix` is the path (string) to the distance matrix created earlier during Step 2. `DistanceCutoff` is the threshold (numeric, in mm) for removing short-distance correlations in the FC matrix. Correlations between nodes < `DistanceCutoff` from each other will be set to zero. This is done to mitigate the effects of spatial autocorrelation on the network structures identified.

`GraphDensities` is a numeric vector of graph densities — the percentage of the top connections retained by each node after thresholding. So for example, in a hypothetical 1000 x 1000 FC matrix, a graph density of 0.05 (5%) means that each node will retain its top 50 strongest connections. These connections will maintain their weights (they will not be binarized), and all other connections will be set to zero.

By default the total number of communities identified by Infomap is data-driven, but can be controlled in part by how many connections are retained in the functional connectivity matrix after thresholding. For example, fewer communities are identified at the 5% threshold (on average,  $8.28 \pm 1.21$ ) than at the 0.1% threshold (on average,  $89.13 \pm 8.04$ ). We recommend running Infomap over a range of graph densities (e.g., 5% to 0.01%, as done in <sup>4</sup>).



`NumberReps` is a numeric value representing the number of times the Infomap algorithm is run before selecting the best solution.

`BadVertices` is an optional index of all points in the brain the user would like to omit from the analysis. For example, if the user knows that data quality is especially poor (low tSNR or test-retest reliability) in a particular set of brain regions. Otherwise, set to `[]` to include all vertices and voxels.

`BrainStructures` is a cell array of brain structures of interest from the `.brainstructurelabel` field of the `ConcatenatedCifti` file. The FC matrix will omit nodes from brain structures not included in this variable.

Run lines 105-106 in the `pfm_tutorial.m` script to run the `pfm_infomap` function.

```

92 %% Step 4: Run infomap.
93
94 % load your concatenated resting-state dataset, pick whatever level of spatial smoothing you want
95 ConcatenatedCifti = ft_read_cifti_mod([PfmDir '/sub-ME01_task-rest_concatenated_smoothed2.55_32k_fsLR.dtseries.nii'])
96
97 % define inputs;
98 DistanceMatrix = [Subdir '/pfm/DistanceMatrix.mat']; % can be path to file
99 DistanceCutoff = 10; % in mm; usually between 10 to 30 mm works well.
100 GraphDensities = flip([0.0001 0.0002 0.0005 0.001 0.002 0.005 0.01 0.02 0.05]); %
101 NumberReps = 50; % number of times infomap is run;
102 BadVertices = []; % optional, but you could include regions to ignore, if you know there is bad signal there.
103 Structures = {'CORTEX_LEFT','CEREBELLUM_LEFT','ACCUMBENS_LEFT','CAUDATE_LEFT','PALLIDIUM_LEFT','PUTAMEN_LEFT','THALAMUS_LEFT'};
104
105 % run infomap
106 pfm_infomap(ConcatenatedCifti,DistanceMatrix,PfmDir,GraphDensities,NumberReps,DistanceCutoff,BadVertices,Structures,1);
107
108 % remove some intermediate files (optional)
109 system(['rm ' Subdir '/pfm/*.net']);
110 system(['rm ' Subdir '/pfm/*.clu']);
111 system(['rm ' Subdir '/pfm/*Log*']);
112
113 % define inputs;
114 Input = [PfmDir '/Bipartite_PhysicalCommunities.dtseries.nii'];
115 Output = 'Bipartite_PhysicalCommunities+SpatialFiltering.dtseries.nii';
116 MinSize = 50; % in mm^2
117
118 % perform spatial filtering
119 pfm_spatial_filtering(Input,PfmDir,Output,MidthickSurfs,MinSize,WorkbenchBinary);
120

```

Optionally, users can discard implausibly small (i.e., smaller than the effective resolution of underlying data) patches of vertices or voxels of a community by specifying a minimum cluster size (in mm<sup>2</sup>). This procedure acts as a spatial filter — removing small objects without imposing additional spatial smoothing on the underlying data. The neighboring network identities are dilated one vertex at a time until the region is filled, as done in <sup>5</sup>.

Run lines 113-119 in the pfmTutorial.m script to define inputs and run the `pfm_spatial_filtering` function.

```

92 %% Step 4: Run infomap.
93
94 % load your concatenated resting-state dataset, pick whatever level of spatial smoothing you want
95 ConcatenatedCifti = ft_read_cifti_mod([PfmDir '/sub-ME01_task-rest_concatenated_smoothed2.55_32k_fsLR.dtseries.nii'])
96
97 % define inputs;
98 DistanceMatrix = [Subdir '/pfm/DistanceMatrix.mat']; % can be path to file
99 DistanceCutoff = 10; % in mm; usually between 10 to 30 mm works well.
100 GraphDensities = flip([0.0001 0.0002 0.0005 0.001 0.002 0.005 0.01 0.02 0.05]); %
101 NumberReps = 50; % number of times infomap is run;
102 BadVertices = []; % optional, but you could include regions to ignore, if you know there is bad signal there.
103 Structures = {'CORTEX_LEFT','CEREBELLUM_LEFT','ACCUMBENS_LEFT','CAUDATE_LEFT','PALLIDIUM_LEFT','PUTAMEN_LEFT','THALAMUS_LEFT'};
104
105 % run infomap
106 pfm_infomap(ConcatenatedCifti,DistanceMatrix,PfmDir,GraphDensities,NumberReps,DistanceCutoff,BadVertices,Structures,1);
107
108 % remove some intermediate files (optional)
109 system(['rm ' Subdir '/pfm/*.net']);
110 system(['rm ' Subdir '/pfm/*.clu']);
111 system(['rm ' Subdir '/pfm/*Log*']);
112
113 % define inputs;
114 Input = [PfmDir '/Bipartite_PhysicalCommunities.dtseries.nii'];
115 Output = 'Bipartite_PhysicalCommunities+SpatialFiltering.dtseries.nii';
116 MinSize = 50; % in mm^2
117
118 % perform spatial filtering
119 pfm_spatial_filtering(Input,PfmDir,Output,MidthickSurfs,MinSize,WorkbenchBinary);
120

```

## Step 5: Algorithmic assignment of network identities to infomap communities.

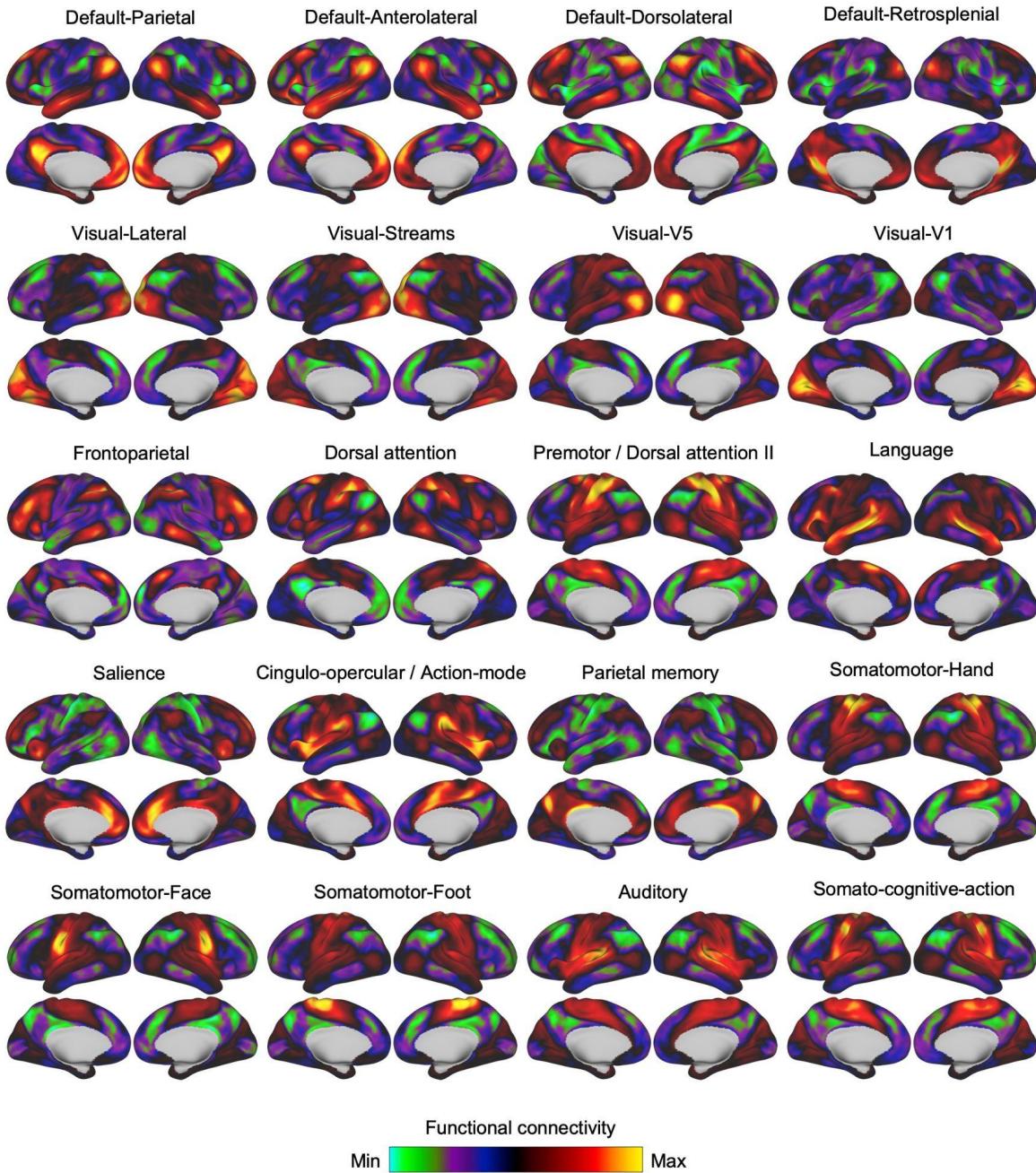
The main output of `pfm_infomap` is a CIFTI file called `"Bipartite_PhysicalCommunities.dtseries.nii"`, which contains the Infomap communities obtained at each graph density (each column represents a different graph density). These community labels are arbitrary — in other words, community number 1 will not represent the same functional network in different individuals, or be interpretable as any particular functional network, such as the salience or default mode network.

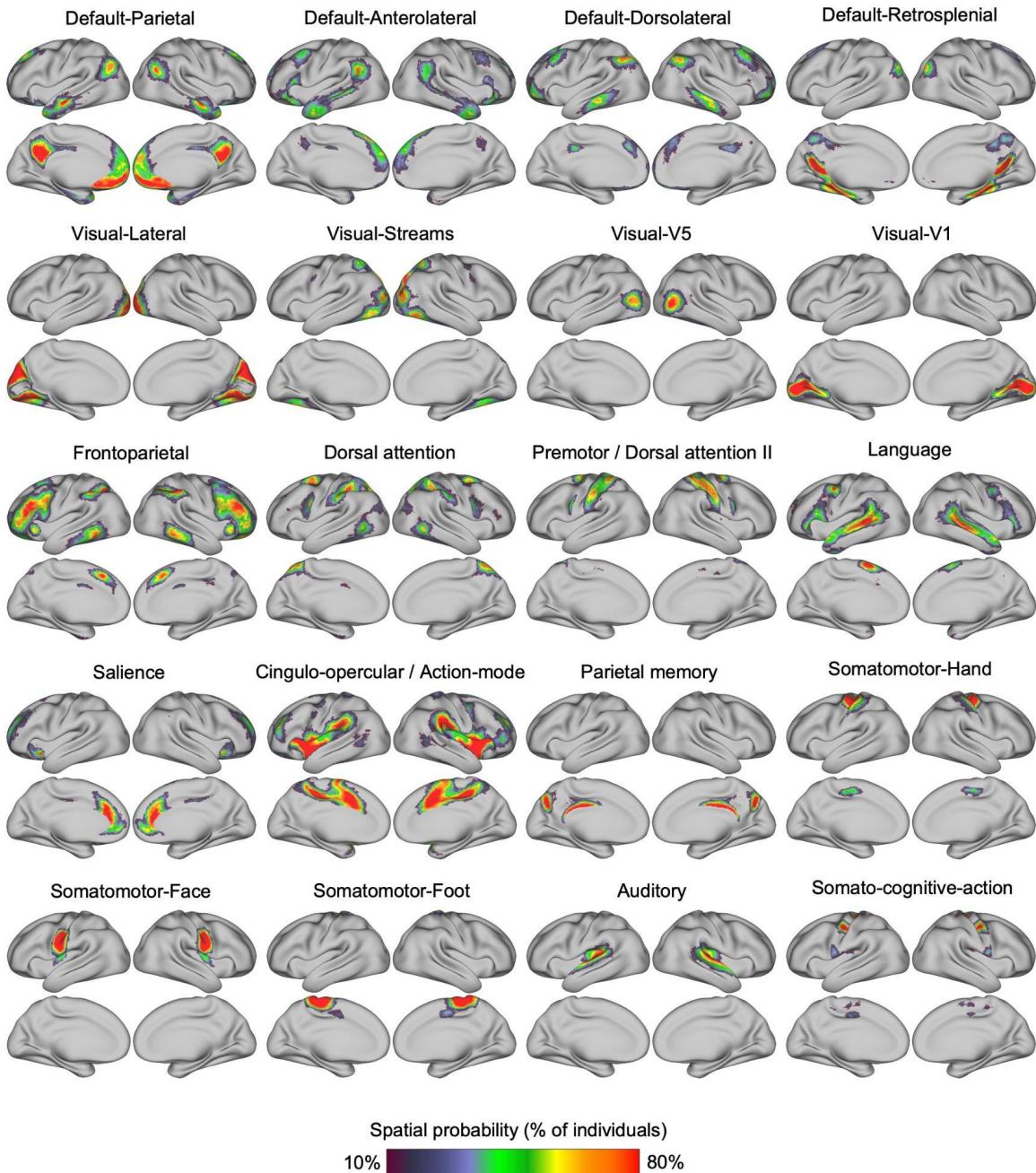
We assign each Infomap community to one of 20 known functional network identities based on their spatial locations and functional connectivity. In principle, this could be accomplished manually by an expert familiar with functional network topography in individuals, but this can be time consuming and difficult to scale. To help accelerate and standardize the network identification process, we have created a semi-automated procedure for quantifying the likelihood of an Infomap community belonging to a particular functional brain network using priors, and specifying the best match as the initial assignment. The entire procedure is implemented using the `pfm_identify_networks` function.

First, in the Matlab editor window, run lines 123-124 in the `pfm_tutorial.m` script to load the default priors for the network identification algorithm.

```
121 %% Step 5: Algorithmic assignment of network identities to infomap communities.
122
123 % load the priors;
124 - load('priors.mat');
125
126 % define inputs;
127 - Ic = ft_read_cifti_mod([PfmDir '/Bipartite_PhysicalCommunities+SpatialFiltering.dtseries.nii']);
128 - Output = 'Bipartite_PhysicalCommunities+AlgorithmicLabeling';
129 - Column = 6; % column 6, representing graph density 0.01% in this example.
130
131 % run the network identification algorithm;
132 - pfm_identify_networks(ConcatenatedCifti,Ic,MidthickSurfs,Column,Priors,Output,PfmDir,WorkbenchBinary);
133
```

`Priors` is a structure array containing the average FC (`Priors.FC`) and spatial locations (`Priors.Spatial`) of 20 functional brain networks — (Default-Parietal, Default-Anterolateral, Default-Dorsolateral, Default-Retrosplenial, Visual-Lateral, Visual-Dorsal/Ventral Stream, Visual-V1, Visual-V5, Frontoparietal, Dorsal Attention, Premotor / Dorsal Attention II, Language, Salience, Cingulo-opercular / Action-mode, Parietal memory, Auditory, Somatomotor-Hand, Somatomotor-Face, Somatomotor-Foot, Auditory, or Somato-Cognitive-Action). The code will accept other priors if they are organized in the same way.





Next, in the Matlab editor window, run lines 124-127 in the pfmTutorial.m script to define the other inputs for the `pfmIdentifyNetworks` function.

```

121 %% Step 5: Algorithmic assignment of network identities to infomap communities.
122
123 % load the priors;
124 load('priors.mat');
125
126 % define inputs;
127 Ic = ft_read_cifti_mod([PfmDir '/Bipartite_PhysicalCommunities+SpatialFiltering.dtseries.nii']);
128 Output = 'Bipartite_PhysicalCommunities+AlgorithmicLabeling';
129 Column = 6; % column 6, representing graph density 0.01% in this example.
130
131 % run the network identification algorithm;
132 pfm_identify_networks(ConcatenatedCifti,Ic,MidthickSurfs,Column,Priors,Output,PfmDir,WorkbenchBinary);
133

```

`InfomapCommunities` is the CIFTI file created by `pfm_infomap`. It contains the Infomap communities obtained at each graph density (each column represents a different graph density). `Column` is the column of `InfomapCommunities` that the user wants to assign network identities. `Output` is the name for the output file.

Next, in the Matlab editor window, run lines 131-132 in the `pfm_tutorial.m` script to run the `pfm_identify_networks` function.

```

121 %% Step 5: Algorithmic assignment of network identities to infomap communities.
122
123 % load the priors;
124 load('priors.mat');
125
126 % define inputs;
127 Ic = ft_read_cifti_mod([PfmDir '/Bipartite_PhysicalCommunities+SpatialFiltering.dtseries.nii']);
128 Output = 'Bipartite_PhysicalCommunities+AlgorithmicLabeling';
129 Column = 6; % column 6, representing graph density 0.01% in this example.
130
131 % run the network identification algorithm;
132 pfm_identify_networks(ConcatenatedCifti,Ic,MidthickSurfs,Column,Priors,Output,PfmDir,WorkbenchBinary);
133

```

## Step 6: Review algorithmic network assignments, adjust if needed.

There are multiple outputs generated by `pfm_identify_networks` that we recommend reviewing carefully before proceeding.

First, there is an .XLS sheet containing the winning and runner-up network assignments for each Infomap community, as well as a brief summary of the information that guided the algorithmic assignments. Second, there are multiple important CIFTI files that are highlighted and described below.

Name	Size	Type
Bipartite_PhysicalCommunities.dtseries.nii	3.6 MB	Binary
Bipartite_PhysicalCommunities+AlgorithmicLabeling.dlabel.nii	919.6 kB	Binary
Bipartite_PhysicalCommunities+AlgorithmicLabeling.L.border	582.2 kB	Marku
Bipartite_PhysicalCommunities+AlgorithmicLabeling.R.border	591.8 kB	Marku
Bipartite_PhysicalCommunities+AlgorithmicLabeling_FC_btwn_InfoMapCommunities.dtseries.nii	26.8 MB	Binary
Bipartite_PhysicalCommunities+AlgorithmicLabeling_FC_btwn_InfoMapCommunities.pdf	143.3 kB	Docum
Bipartite_PhysicalCommunities+AlgorithmicLabeling_FC_WholeBrain.dtseries.nii	26.8 MB	Binary
Bipartite_PhysicalCommunities+AlgorithmicLabeling_InfoMapCommunities.dlabel.nii	27.0 MB	Binary

`Bipartite_PhysicalCommunities+AlgorithmicLabeling.dlabel.nii` is a CIFTI file containing the initial algorithmic network assignments for all Infomap communities (set in Tab 1 in images below). The colors used to represent different functional brain networks are set by the RGB values in `Priors.NetworkColors`.

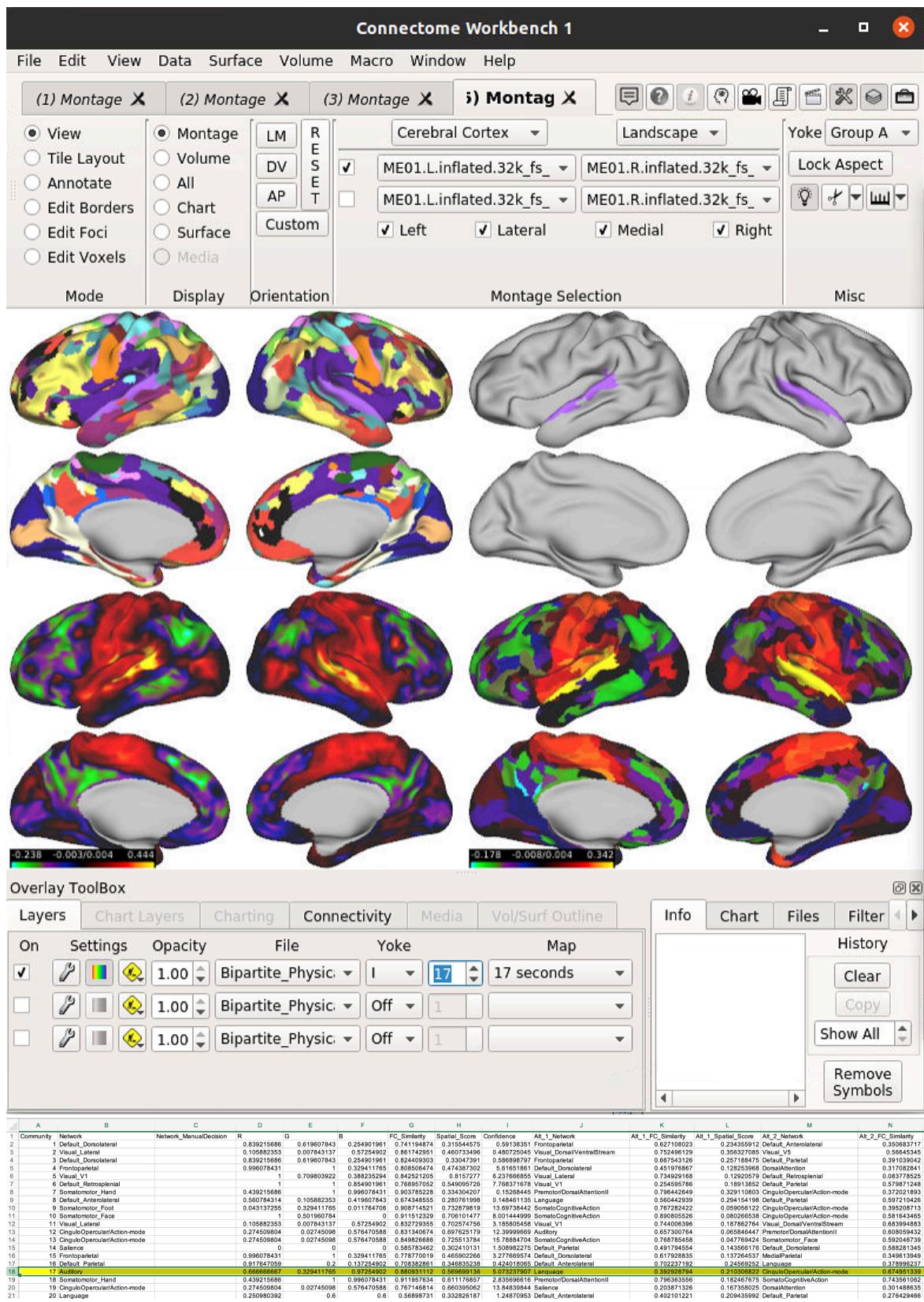
`Bipartite_PhysicalCommunities+AlgorithmicLabeling_InfoMapCommunities.dlabel.nii` is a CIFTI file containing the initial algorithmic network assignment for each Infomap community, stored separately in each column (set in Tab 2 in images below).

`Bipartite_PhysicalCommunities+AlgorithmicLabeling_FC_WholeBrain.dtseries.nii` is a CIFTI file containing the whole-brain functional connectivity of each Infomap community, stored separately in each column (set in Tab 3 in images below).

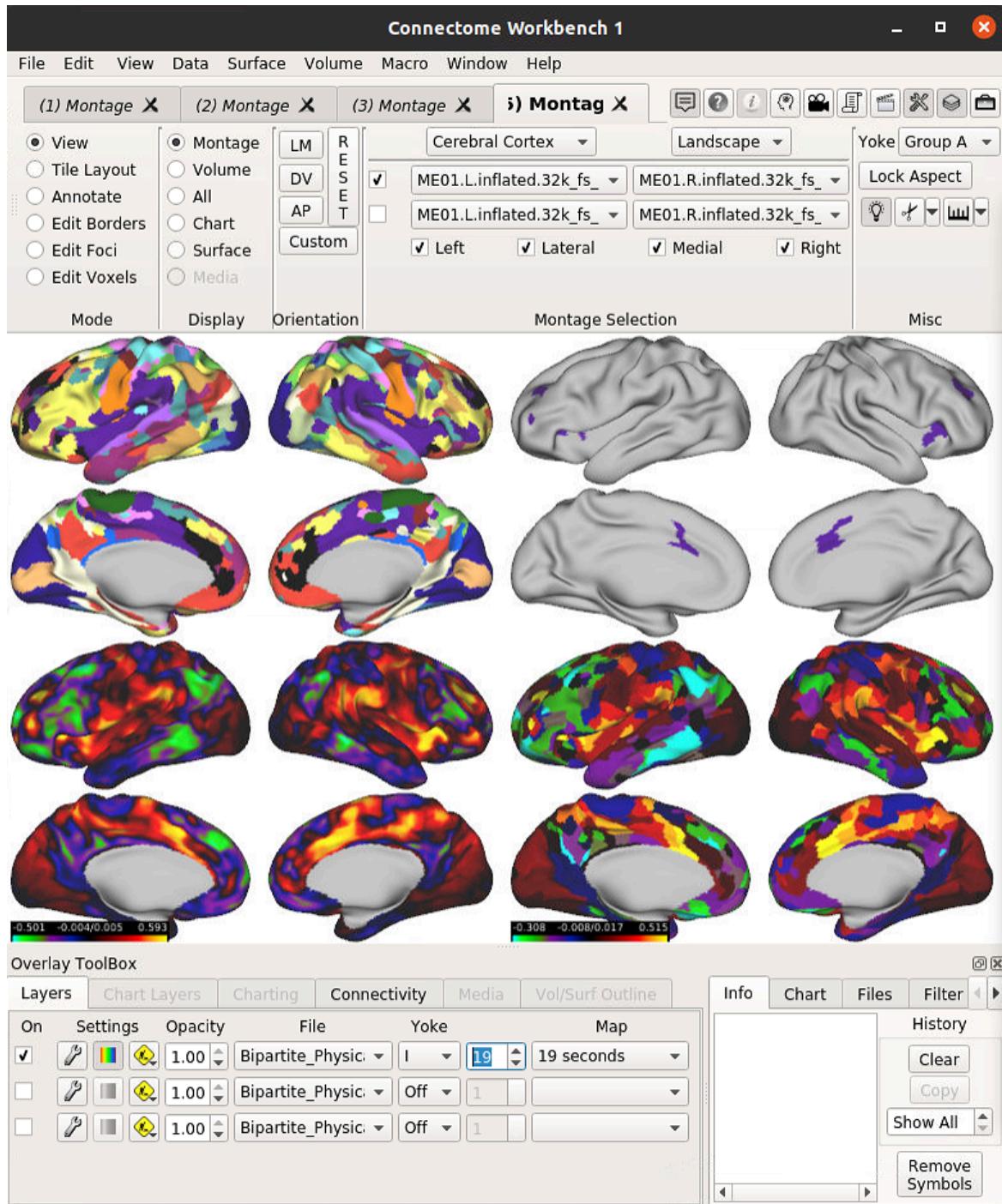
`Bipartite_PhysicalCommunities+AlgorithmicLabeling_FC_btwn_InfoMapCommunities.dtseries.nii` is a CIFTI file containing the functional connectivity of each Infomap community to all other Infomap communities, stored separately in each column (set in Tab 4 in images below).

To review the algorithmic network assignments, we recommend loading the subject's anatomical data and the four CIFTI files above into Connectome Workbench. Create a 2 x 2 tile tab configuration to simultaneously view all of the relevant information. Tabs 2-4 can be set to "Yoke I" in the *Overlay Toolbox*, so that when you flip through the columns of one file, the others will also update. This allows you to flip through individual communities and view their algorithmic assignment and spatial locations (Tab 2), as well as their functional connectivity with the rest of the brain (Tab 3) and with the other communities (Tab 4).

For example, community 17 (highlighted below), was labeled *Auditory* (Column B in the XLS). The spatial correlation of community 1 FC with the *Auditory* template FC was  $r = 0.88$  (Column G in the XLS). The average *Auditory* spatial probability value in community 1 was 0.57 (Column H in the XLS). The total score for this winning assignment is the product of the FC\_Similarity and Spatial\_Score ( $0.88 * 0.57 = 0.50$ ). The total score for the first runner-up assignment (*Language*) is also the product of the FC\_Similarity and Spatial\_Score ( $0.39 * 0.21 = 0.08$ ). The "Confidence" of the winning assignment (0.59) is quantified in Column I, and is defined as the relative difference in total score associated with the winning and first runner-up network assignments ( $[0.502 - 0.083] / 0.083 = 5.07$ ).



Another example is shown below. In this case, community 19 (highlighted below), was labeled *Cingulo-opercular / Action-mode*.

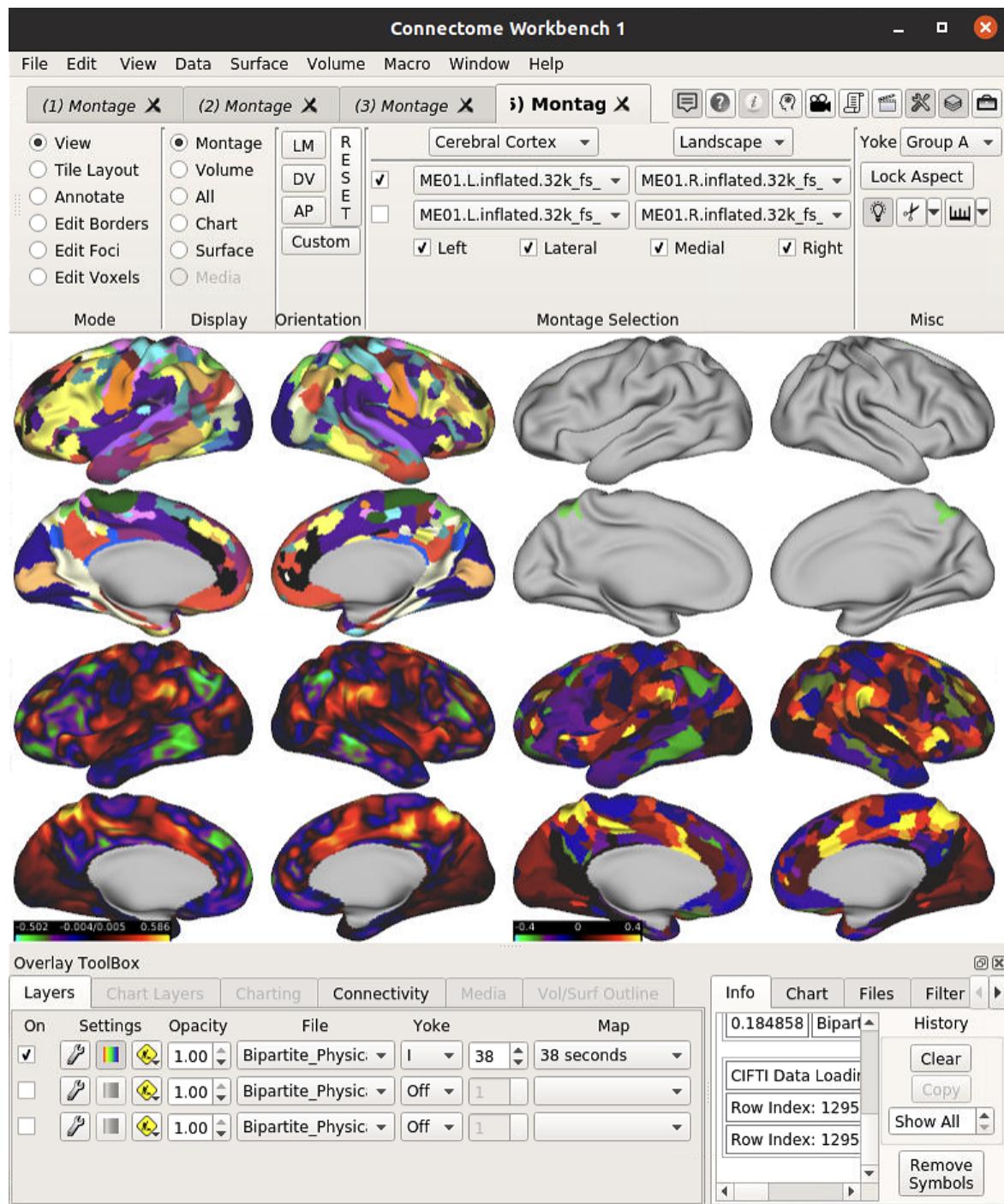


A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	
1	Community	Network	Network_ManualDecision	R	G	B	FC_Similarity	Spatial_Score	Confidence	Alt_1_Network	Alt_1_FC_Similarity	Alt_1_Spatial_Score	Alt_2_Network	Alt_2_FC_Similarity	Alt_2_Spatial_Score
2	1	Default_Dorsolateral		0.839015686	0.519607843	0.254001061	0.744194874	0.31544175	0.50130519	Frontoparietal	0.620168093	0.25435412	0x4a5b_Anteriorlateral	0.620168093	0.137070119
3	2	Visual_Lateral		0.105882353	0.078473137	0.57254029	0.861742951	0.460734075	0.480725045	Visual_V5	0.752406129	0.356327055	Visual_V5	0.56643345	0.04220592
4	3	Default_Dorsolateral		0.839215686	0.619607843	0.254001061	0.82409303	0.30047398	0.56869879	Frontoparietal	0.667564312	0.257188475	Default_Parietal	0.391309042	0.161668304
5	4	Premotor		0.996078431	0.105882353	0.57254029	0.845251205	0.460734075	0.480725045	Default_Dorsolateral	0.744929168	0.12625579	Default_Dorsolateral	0.317110037	0.105000138
6	5	Visual_V1		1	0.709803922	0.382352368	0.54095723	0.815727277	0.423766855	Visual_V1	0.25459578	0.1891385	Default_Parietal	0.57877248	0.04050129
7	6	Default_Retrospenial		1	0.854001061	0.76897052	0.54095723	7.768371678	0.117683716	Visual_V1	0.25459578	0.1891385	Default_Parietal	0.57877248	0.04050129
8	7	Somatotmotor_Hand		0.493615686	0.996078431	0.09087431	0.903785228	0.334304207	0.15268445	PremotorDorsalAttentionII	0.766442469	0.32911983	CinguloOpercularAction-mode	0.37202193	0.14529203
9	8	Default_Anterolateral		0.560784314	0.105882353	0.57254029	0.709803922	0.334304207	0.15268445	Default_Anterolateral	0.744929168	0.12625579	Default_Anterolateral	0.34910656	0.103700119
10	9	Somatotmotor_Foot		0.043137255	0.329417678	0.011768371	0.908745219	0.136973842	0.15268445	SomatoCognitiveAction	0.767328422	0.059056122	CinguloOpercularAction-mode	0.395209713	0.10564059
11	10	Visual_Lateral		1	0.205196078	0.050130519	0.911512329	0.760101477	0.800144499	SomatoCognitiveAction	0.89026655	0.256023226	CinguloOpercularAction-mode	0.581643465	0.07631635
12	11	Visual_Lateral		0.105882353	0.57254029	0.57254029	0.760101477	0.31364458	0.15268445	Visual_V1	0.25459578	0.1891385	Visual_V1	0.620168093	0.030000142
13	12	CinguloOpercularAction-mode		0.274508924	0.027450892	0.57254029	0.86762549	0.399600009	0.15268445	CinguloOpercularAction-mode	0.657307654	0.056564447	CinguloOpercularAction-mode	0.609503432	0.04519275
14	13	CinguloOpercularAction-mode		0.274508924	0.027450892	0.57254029	0.86762549	0.399600009	0.15268445	CinguloOpercularAction-mode	0.767884704	0.256023226	SomatoCognitiveAction	0.592047379	0.059131161
15	14	Salience		0	0	0.585783462	0.302410131	1.509882275	0.15268445	Default_Parietal	0.491749544	0.145568174	Default_Parietal	0.58821345	0.116094771
16	15	Anterolateral		0.996078431	1	0.329411768	0.86762549	0.15268445	Anterolateral	0.744929168	0.12625579	Anterolateral	0.34910656	0.103700119	
17	16	Default_Parietal		0.917647059	0.2	0.137254029	0.709803922	0.348853288	0.424010605	Default_Anterolateral	0.702231982	0.24529428	Language	0.378992337	0.08321582
18	17	Auditory		0.666666667	0.329411768	0.25245029	0.880931112	0.569699138	0.97237907	Language	0.392928794	0.210306822	CinguloOpercularAction-mode	0.674951339	0.11208222
19	18	Somatotmotor_Hand		0.439215686	0.996078431	0.09087431	0.919576324	0.611768578	0.253596122	SomatoCognitiveAction	0.763530556	0.182407675	SomatoCognitiveAction	0.743561052	0.04658793
20	19	DorsalAttention		0.043137255	0.205196078	0.050130519	0.911512329	0.136973842	0.15268445	DorsalAttention	0.767328422	0.059056122	CinguloOpercularAction-mode	0.767328422	0.131398697
21	20	Lanuseus		0.25090392	0.6	0.6	0.6	0.6	0.6	0.6	0.240945992	0.240945992	Default_Parietal	0.276429466	0.131398697

Note that this approach works well when functional networks are located more or less in their “typical” locations. However, communities may be “incorrectly” labeled if they have a low Spatial\_Score (i.e., minimal overlap with the typical network location).

These cases tend to be accompanied by low Confidence scores. Users can sort the XLS sheet by the Confidence values (as done below) and visually examine low confidence assignments to help guard against potential misclassifications. We have found that a good rule of thumb is to focus on Confidence values of 0.33 or less (highlighted below), but it is useful to carefully review all of the assignments, if time permits.

A	B	C	D	E	F	G	H	I	J	K	L	M	N	O		
1	Community	Network	Network_ManualDecision	R	G	B	FC_Similarity	Spatial_Score	Confidence	Alt_1_Network	Alt_1_FC_Similarity	Alt_1_Spatial_Score	Alt_2_Network	Alt_2_FC_Similarity	Alt_2_Spatial_Score	
2	1	CinguloOpercularAction-mode		0.974508924	0.027450892	0.576470586	0.760101477	0.399600009	0.15268445	Auditor	0.620168093	0.25435412	0x4a5b_Anteriorlateral	0.620168093	0.137070119	
3	2	Visual_DorsalVentralStream		0.184313725	0.458823529	0.709803922	0.71751487	0.105988227	0.15268445	Visual_DorsalVentralStream	0.572433468	0.257188475	CinguloOpercularAction-mode	0.814096292	0.107708979	
4	3	Visual_V5		0.725408924	0.811764705	0.564705882	0.72771487	0.1378614	0.04734886	Visual_DorsalVentralStream	0.48742482	0.203970981	DorsalAttention	0.232988137	0.142039846	
5	4	Selence		0	0	0	0	0	0	0	0	0	0	0		
6	5	Default_Anterolateral		0.560784314	0.105882353	0.419607843	0.891957632	0.334304207	0.15268445	CinguloOpercularAction-mode	0.170795867	0.303597172	Default_Anterolateral	0.340784685	0.122781775	
7	6	Somatotmotor_Hand		0.439215686	0.996078431	0.09087431	0.919576324	0.136973842	0.15268445	PremotorDorsalAttentionII	0.766442469	0.23911983	CinguloOpercularAction-mode	0.37202193	0.145292033	
8	7	Default_Dorsolateral		0.388235294	0.892156866	0.247058824	0.747833917	0.41598971	0.213530277	Visual_DorsalVentralStream	0.658829043	0.399030432	Frontoparietal	0.184934746	0.077568464	
9	8	PremotorDorsalAttentionII		1	0.57254029	0.050130519	0	0.760101477	0.334304207	0.15268445	CinguloOpercularAction-mode	0.215532077	0.275525258	Somatomotor_Foot	0.869176948	0.150847147
10	9	Visual_Face		0	0	0	0	0	0	0	0	0	0	0		
11	10	SematoCognitiveAction		0.892156866	0.247058824	0.414344646	0.40584569	0.321867499	0.213530277	CinguloOpercularAction-mode	0.636207924	0.276625258	Default_Retrospenial	0.205082247	0.050204169	
12	11	CinguloOpercularAction-mode		0.996078431	0.105882353	0.419607843	0.780199867	0.409666923	0.213530277	Default_Dorsolateral	0.463254157	0.151931147	Language	0.276124843	0.197250861	
13	12	Frontoparietal		0	0	0	0	0	0	0	0	0	0	0		
14	13	Anterolateral		0	0	0	0	0	0	0	0	0	0	0		
15	14	Default_Parietal		0.917647059	0.2	0.137254029	0.709803922	0.348853288	0.424010605	Default_Anterolateral	0.722337192	0.24599252	Language	0.378992337	0.08321582	
16	15	Default_Anterolateral		0.560784314	0.104882353	0.419607843	0.819573224	0.334450172	0.403820417	Default_Parietal	0.783900874	0.237514318	Default_Dorsolateral	0.532250342	0.136597938	
17	16	SematoCognitiveAction		0.709803922	0	0	0.078473137	0.09087431	0.15268445	CinguloOpercularAction-mode	0.490354045	0.24529428	SomatoCognitiveAction	0.56643345	0.04220592	
18	17	Visual_Lateral		0.105882353	0.078473137	0.57254029	0.861742951	0.460734045	0.213530277	DorsalAttention	0.732496129	0.356327055	Visual_V5	0.56643345	0.04220592	
19	18	Visual_DorsalVentralStream		0.184313725	0.458823529	0.709803922	0.735867793	0.369124877	0.483115741	DorsalAttention	0.424142233	0.434611603	PremotorDorsalAttentionII	0.740788217	0.082988184	
20	19	Default_Anterolateral		0.560784314	0.105882353	0.419607843	0.780199867	0.409666923	0.213530277	Default_Dorsolateral	0.693342509	0.399043939	Default_Parietal	0.694147953	0.104298459	
21	20	Default_Dorsolateral		0.839215686	0.619607843	0.254001061	0.760101477	0.334304207	0.15268445	Frontoparietal	0.620168093	0.25435412	Anterolateral	0.391309042	0.161668304	
22	21	PremotorDorsalAttentionII		1	0.560784314	0.247058824	0.334304207	0.300473981	0.586897797	Frontoparietal	0.675745126	0.257188475	Default_Parietal	0.391309042	0.161668304	
23	22	Default_Dorsolateral		0.839215686	0.619607843	0.254001061	0.760101477	0.334304207	0.15268445	CinguloOpercularAction-mode	0.591147449	0.24599252	Anterolateral	0.499847815	0.107800168	
24	23	PremotorDorsalAttentionII		0.839215686	0.619607843	0.254001061	0.760101477	0.334304207	0.15268445	Frontoparietal	0.757700383	0.253525258	Default_Dorsolateral	0.35088373	0.136700168	
25	24	Default_Parietal		0.917647059	0.2	0.137254029	0.709803922	0.345560531	0.420382398	Default_Dorsolateral	0.778910076	0.07243108	Default_Retrospenial	0.5351245	0.110944027	
26	25	Default_Retrospenial		1	0.329411768	0.25245029	0.743090474	0.708670047	0.308627501	DorsalAttention	0.395535319	0.14720738	MediaParietal	0.395595438	0.082174688	
27	26	Default_Anterolateral		0.709803922	0	0	0	0	0	0	0	0	0	0		
28	27	SematoCognitiveAction		0.470588235	0.078473137	0.093849813	0.451342876	0.853852876	0.253596122	Somatomotor_Hand	0.856528349	0.263285617	Somatomotor_Foot	0.877094206	0.032338193	
29	28	CinguloOpercularAction		0.470588235	0.078473137	0.093849813	0.451342876	0.853852876	0.253596122	PremotorDorsalAttentionII	0.826139053	0.201241215	PremotorDorsalAttention_Foot	0.797541585	0.153394687	
30	29	Language		0.329411768	0.25245029	0.743090474	0.708670047	0.308627501	0.15268445	Default_Parietal	0.621169697	0.232641166	Default_Parietal	0.261877622	0.04896782	
31	30	Anterolateral		0.388235294	0.892156866	0.247058824	0.676584551	0.341144603	0.209341166	Frontoparietal	0.621169697	0.232641166	Anterolateral	0.261877622	0.04896782	
32	31	Language		0.25090392	0.6	0	0	0	0	0	0	0	0	0		
33	32	PremotorDorsalAttentionII		0.25090392	0	0	0	0	0	0	0	0	0	0		
34	33	Anterolateral		0.917647059	1	0.329411768	0.25245029	0.743090474	0.420382398	1.318150177	DorsalAttention	0.596484824	0.230728178	Visual_DorsalVentralStream	0.437699331	0.087301587
35	34	Default_Parietal</td														



A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	
Community	Network	Network_ManualDecision	R	G	B	FC_Similarity	Spatial_Score	Confidence	Alt_1_Network	Alt_1_FC_Similarity	Alt_1_Spatial_Score	Alt_2_Network	Alt_2_FC_Similarity	Alt_2_Spatial_Score	
2	38	CinguOpercularAction-mode	0.274509804	0.027450984	0.676470588	0.805963345	0.412367367	0.013459784	Auditory	0.961796446	0.278202320	SomatoCognitiveAction	0.683511448	0.04787799	
3	50	Visual,DorsalVentralStream	0.184313725	0.498253592	0.709803923	0.737521487	0.196483353	0.02760549	DorsalAttention	0.572783436	0.267452301	CinguloOpercularAction-mode	0.614062292	0.107706697	
4	72	Visual_V5	0.725490198	0.817687407	0.564705882	0.756270312	0.137688114	0.047348585	Visual,DorsalVentralStream	0.48742482	0.203970861	DorsalAttention	0.232888137	0.142038946	
5	71	Visual_V5	0.725490198	0.817687407	0.564705882	0.756270312	0.137688114	0.047348585	Visual,DorsalVentralStream	0.48742482	0.203970861	AnteriorLateral	0.34320207	0.107706697	
6	6	Default_Anterolateral	0.560794314	0.105882353	0.419607843	0.674345555	0.260719196	0.146461153	Language	0.560442939	0.294154184	Default_Parietal	0.597210428	0.162536179	
7	7	Somatotor_Hand	0.439215686	1	0.996708431	0.903785228	0.334304207	0.15268445	Language	0.766426469	0.329110863	CinguloOpercularAction-mode	0.372021893	0.145292033	
8	62	DefaultAnterior	0.388253594	0.659215688	0.247058824	0.747833517	0.1556097	0.213532077	Visual,DorsalVentralStream	0.858282025	0.389030432	Frontoparietal	0.184934476	0.077955446	
9	40	PremotorDorsalAttention	0.105882353	0.078411785	0.260705198	0.460719196	0.1556097	0.213532077	Visual,DorsalVentralStream	0.276641144	0.321867499	Default_Foot	0.838253594	0.117706697	
10	56	Somatotor_Face	1	0.519697084	0	0.789737871	0.391477277	0.29308709	SomatotogaiveAction	0.803828255	0.276641144	CinguloOpercularAction-mode	0.785823011	0.190340903	
11	38	DorsalAttention	0.838253594	0.839215686	0.447058824	0.414434464	0.405845691	0.321867499	CinguloOpercularAction-mode	0.862079824	0.2	Language	0.205682241	0.05024169	
12	42	Default_Parietal	0.996708431	0.32911785	0.260705198	0.460719196	0.1556097	0.213532077	Visual,DorsalVentralStream	0.572783436	0.181805179	Language	0.572783436	0.04787799	
13	61	Default_Parietal	0.917647059	0.560803923	0.254019681	0.544413322	0.344413322	0.37468908	Default_Retroparietal	0.543117204	0.249615022	Default_DorsalAttention	0.769379712	0.110384146	
14	16	Default_Parietal	0.917647059	0.560803923	0.254019681	0.544413322	0.344413322	0.37468908	Default_Retroparietal	0.543117204	0.249615022	Default_DorsalAttention	0.769379712	0.110384146	
15	35	Default_Anterolateral	0.560794314	0.105882353	0.419607843	0.819573224	0.334401712	0.463820417	Default_Parietal	0.788300874	0.237514318	Default_DorsalAttention	0.532250342	0.136597938	
16	34	DefaultAnterior	0.725490198	0.078411785	0.260705198	0.460719196	0.1556097	0.213532077	Visual,DorsalVentralStream	0.276641144	0.321867499	Default_Parietal	0.733202239	0.107706697	
17	2	Visual_Lateral	0.105882353	0.078411785	0.260705198	0.460719196	0.1556097	0.213532077	Visual,DorsalVentralStream	0.724298129	0.306327085	Visual_V5	0.56643345	0.04220992	
18	68	Visual,DorsalVentralStream	0.184313725	0.498253592	0.709803922	0.735867933	0.369124877	0.483115704	DorsalAttention	0.421402232	0.434811633	PremotorDorsalAttention	0.740768217	0.082898184	
19	64	Default_Anterolateral	0.105882353	0.078411785	0.260705198	0.460719196	0.1556097	0.213532077	Visual,DorsalVentralStream	0.572783436	0.2	Default_Parietal	0.88911749	0.04787799	
20	41	DefaultAnterior	0.105882353	0.078411785	0.260705198	0.460719196	0.1556097	0.213532077	Visual,DorsalVentralStream	0.572783436	0.2	Default_Parietal	0.639677851	0.04525202	
21	3	Default_Dorsal	0.839215686	0.619607843	0.254019681	0.824403003	0.30304798	0.58689879	Frontoparietal	0.687543126	0.257118475	Default_Parietal	0.391030042	0.161668304	
22	33	PremotorDorsalAttention	0.105882353	0.078411785	0.260705198	0.460719196	0.1556097	0.213532077	Visual,DorsalVentralStream	0.687444667	0.202641168	Default_Parietal	0.235902323	0.107706697	
23	27	Default_Dorsal	0.839215686	0.619607843	0.254019681	0.824403003	0.30304798	0.58689879	Frontoparietal	0.687444667	0.202641168	Default_Parietal	0.235902323	0.107706697	
24	67	PremotorDorsalAttention	1	0.509803922	1	0.690088984	0.414481893	0.686611367	DorsalAttention	0.75720938	0.226966292	CinguloOpercularAction-mode	0.629767871	0.116664245	
25	77	Default_Parietal	0.917647059	0.2	0.137254902	0.520860531	0.245448953	0.687844647	Default_DorsalAttention	0.778910078	0.097243106	Default_Retroparietal	0.535132435	0.119940207	
26	37	Default_Retroparietal	0.917647059	0.2	0.137254902	0.520860531	0.245448953	0.687844647	Default_DorsalAttention	0.778910078	0.097243106	Default_Retroparietal	0.535132435	0.119940207	
27	73	PremotorDorsalAttention	1	0.509803922	1	0.902544974	0.427390163	0.736206187	Default_Frontal_Face	0.816977887	0.27200861	Default_Frontal_Hand	0.847987858	0.062533	
28	29	SomatotorAction	0.470582835	0.07841373	0.260705198	0.460719196	0.1556097	0.213532077	Somatotor_Hand	0.856628349	0.263265617	Somatotor_Foot	0.877042026	0.147511929	
29	21	CinguOpercularAction-mode	0.274509804	0.072450984	0.576470588	0.776588329	0.3408927	0.105944359	PremotorDorsalAttention	0.82613936	0.201242115	Somatotor_Foot	0.797541585	0.153394487	
30	60	DefaultAnterior	0.274509804	0.072450984	0.576470588	0.776588329	0.3408927	0.105944359	DefaultAnterior	0.82613936	0.201242115	DefaultAnterior	0.797541585	0.153394487	
31	29	DorsalAttention	0.388253594	0.839215686	0.247058824	0.876858681	0.332821867	1.209341168	Frontoparietal	0.621698722	0.202641168	MediaParietal	0.261877622	0.04898782	
32	30	Language	0.250980392	0.6	0.6	0.6	0.5898873	0.332821867	1.24870953	Default_Anterolateral	0.420210021	0.209435962	Default_Parietal	0.276424946	0.131396957
33	51	PremotorDorsalAttention	1	0.509803922	1	0.93141765	0.427390163	1.311501777	DorsalAttention	0.686236334	0.2	Visual,DorsalVentralStream	0.639647773	0.087301587	
34	58	Default_Parietal	0.917647059	0.2	0.137254902	0.74023657	0.369588263	1.346481893	Default_Parietal	0.686236334	0.180241327	Default_DorsalAttention	0.639647773	0.087301587	
35	34	14	Salience	0	0	0	0.585738462	0.30241013	1.508982275	Default_Parietal	0.491748454	0.143568178	Default_DorsalAttention	0.588281345	0.116094771
36	74	Default_Parietal	0.996708431	1	0.329411765	0.59823042	0.293867501	1.522512723	DorsalAttention	0.386844074	0.199979098	Default_Retroparietal	0.5717173183	0.022662266	
37	48	Default_Parietal	0.996708431	0.2	0.137254902	0.74023657	0.369588263	1.346481893	Default_Parietal	0.386844074	0.199979098	Default_Retroparietal	0.5717173183	0.022662266	
38	39	Frontoparietal	0.996708431	1	0.329411765	0.73421948	0.309435028	1.666751792	Salience	0.470625276	0.233804143	Default_Anterolateral	0.138642031	0.03955544	
39	59	CinguOpercularAction-mode	0.274509804	0.072450984	0.576470588	0.826305791	0.309435028	1.572725984	PremotorDorsalAttention	0.82207974	0.20090765	Somatotor_Face	0.669336388	0.10538602	
40	23	CinguOpercularAction-mode	0.274509804	0.072450984	0.576470588	0.826305791	0.309435028	1.572725984	PremotorDorsalAttention	0.649549276	0.148667868	Frontoparietal	0.217351764	0.1784071559	
41	49	Salence	0	0	0	0.645687453	0.249508903	1.004616562	Default_Parietal	0.308660523	0.142669297	Default_Parietal	0.362257685	0.116737073	

Once all of the assignments have been inspected and the XLS file containing the manual decisions has been saved, you can run `pfm_parse_manual_decisions` to incorporate the manual labels.

In the Matlab editor window, run lines 136-141 to define the inputs and run `pfm_parse_manual_decisions`.

```

134 %% Step 6: Review algorithmic network assignments, optionally adjust labels manually if needed.
135 %
136 % define inputs
137 - XLS = [PfmDir '/Bipartite_PhysicalCommunities+AlgorithmicLabeling_NetworkLabels.xls']; % note that the XLS
138 - Output = 'Bipartite_PhysicalCommunities+FinalLabeling';
139 -
140 % OPTIONAL: update network assignments according to manual decisions;
141 - pfm_parse_manual_decisions(Ic,Column,MidthickSurfs,Priors,XLS,Output,PfmDir,WorkbenchBinary);

```

## Step 7: Calculate size of each functional brain network.

The main output of `pfm_parse_manual_decisions` is a CIFTI file (called "Bipartite\_PhysicalCommunities+FinalLabeling.dlabel.nii"). The `pfm_calculate_network_size` function can be used to calculate the size of each functional brain network (the percentage of cortical surface area it occupies).

In the Matlab editor window, run lines 145-151 in the `pfm_tutorial.m` script to define the inputs and.

```

143 %% Step 7: Calculate size of each functional brain network
144 %
145 % define inputs
146 - FunctionalNetworks = ft_read_cifti_mod([PfmDir '/Bipartite_PhysicalCommunities+FinalLabeling.dlabel.nii']);
147 - VA = ft_read_cifti_mod(['Subdir '/fs_LR/fsaverage_LR32k/' Subject '.midthickness_va_32k_fs_LR.scalar.nii']);
148 - Structures = {'CORTEX_LEFT','CORTEX_RIGHT'}; % in this case, cortex only.
149 -
150 % calculate the size of each functional brain network
151 NetworkSize = pfm_calculate_network_size(FunctionalNetworks,VA,Structures);

```

`FunctionalNetworks` is a CIFTI file containing the “final” (after manual review) functional network maps.

`VA` is a CIFTI file containing the surface area (in mm^2) each vertex is responsible for.

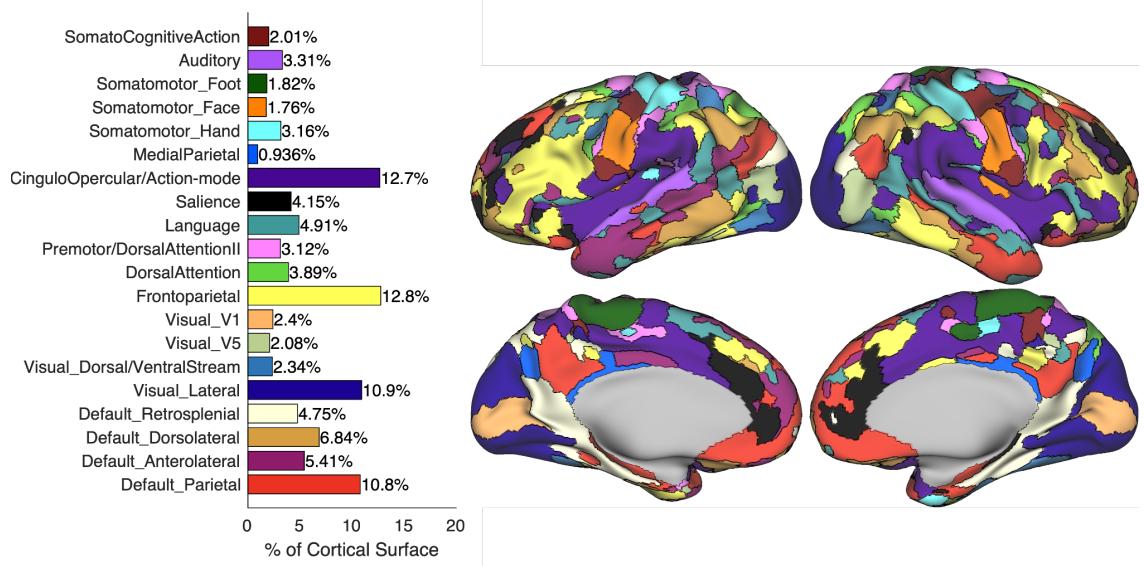
`Structures` is a cell array of brain structures. The network size calculation will be constrained to the structures in this variable.

In the Matlab editor window, run lines 153-175 to run the `pfm_calculate_network_size` function and visualize the results.

```

143 %% Step 7: Calculate size of each functional brain network
144
145 % define inputs
146 FunctionalNetworks = ft_read_cifti_mod([PfmDir '/Bipartite_PhysicalCommunities+FinalLabeling.dlabel.nii']);
147 VA = ft_read_cifti_mod([Subdir '/fs_LR/fsaverage_LR32k/' Subject '.midthickness_va.32k_fs_LR.dscalar.nii']);
148 Structures = {'CORTEX_LEFT','CORTEX_RIGHT'}; % in this case, cortex only.
149
150 % calculate the size of each functional brain network
151 NetworkSize = pfm_calculate_network_size(FunctionalNetworks,VA,Structures);
152
153 close all; % blank slate
154 H = figure; % preallocate parent figure
155 set(H,'position',[1 1 325 400]); hold;
156
157 % unique functional networks:
158 uCi = unique(nonzeros(FunctionalNetworks.data));
159
160 % sweep through
161 % the networks:
162 for i = 1:length(uCi)
163     Tmp = nan(1,length(Priors.NetworkLabels));
164     Tmp(i) = NetworkSize(i);
165     barh(Tmp,'FaceColor',Priors.NetworkColors(i,:));
166     text((NetworkSize(i)+0.1),i,[num2str(NetworkSize(i),3) '%']);
167 end
168
169 % make it pretty:
170 yticklabels(Priors.NetworkLabels);
171 yticks(1:length(uCi)); ylim([0 21]);
172 xlim([0 20]); xticks(0:5:20);
173 set(gcf,'fontname','arial','fontsize',10,'TickLength',[0 0],'TickLabelInterpreter','none');
174 xlabel('% of Cortical Surface');
175 print(gcf,[PfmDir '/FunctionalNetworkSizes'],'-dpdf');
176

```



## References

1. Lynch, C. J. *et al.* Rapid Precision Functional Mapping of Individuals Using Multi-Echo fMRI. *Cell Rep.* **33**, 108540 (2020).
2. Buckner, R. L., Krienen, F. M., Castellanos, A., Diaz, J. C. & Yeo, B. T. T. The organization of the human cerebellum estimated by intrinsic functional connectivity. *J. Neurophysiol.* **106**, 2322–2345 (2011).
3. Marek, S. *et al.* Spatial and Temporal Organization of the Individual Human Cerebellum. *Neuron* **100**, 977–993.e7 (2018).
4. Gordon, E. M. *et al.* Default-mode network streams for coupling to language and control systems. *Proc. Natl. Acad. Sci. U. S. A.* **117**, 17308–17319 (2020).
5. Gordon, E. M. *et al.* Precision Functional Mapping of Individual Human Brains. *Neuron* **95**, 791–807.e7 (2017).