

02_haxby_fmri

February 25, 2019

1 Example 2: Visualizing the Haxby (fMRI) dataset with kmapper + dyneusr

1.1 1 import libraries

```
In [1]: # for auto-reloading external modules
        %load_ext autoreload
        %autoreload 2
```

```
In [2]: import os
```

```
In [3]: import numpy as np
        import pandas as pd
        import scipy as sp

        import networkx as nx
        from collections import Counter
```

```
In [4]: import matplotlib as mpl
        import matplotlib.pyplot as plt
        import seaborn as sns
        sns.set("paper", "white")

        %matplotlib inline
```

1.1.1 1.1 import kmapper

Here, we will use the KeplerMapper (kmapper) implementation of the Mapper algorithm.

We will also import sklearn implementations of PCA and TSNE to use as (linear and non-linear) filter functions for Mapper.

```
In [5]: import kmapper as km
        from sklearn.manifold import TSNE
        from sklearn.decomposition import PCA
```

1.1.2 1.2 import dyneusr

Note, dyneusr provides a wrapper around kmapper with support for caching each step of the Mapper algorithm. This will speed things up, especially when generating several shape graphs for different subjects. For a more detailed walk-through of how to generate a shape graph using kmapper, see 01_trefoil_knot/01_trefoil_knot.ipynb.

```
In [6]: import dyneusr as ds
```

```
# kmapper wrappers
from dyneusr.mapper.wrappers import KMapperWrapper
from dyneusr.mapper.utils import optimize_cover, optimize_dbscan
```

1.2 2 Load data

```
In [7]: from load_data import load_haxby, Bunch
```

Here, we actually use `nilearn.datasets.fetch_haxby` to fetch the files for the Haxby dataset. To make this note book more readable, we wrapped several steps into a single `load_haxby` function, stored in `load_data.py`.

These steps include: 1. fetching the Haxby dataset files; 2. performing some preprocessing; and 3. storing the data for each subject as an `sklearn.datasets.base.Bunch` object for easy access.

A list of these objects, one for each subject, is stored in a higher-level object, which also stores some other important meta-data shared across subjects. This higher-level object is returned by the function, `load_haxby`.

```
In [8]: # load data for all 6 subjects, but only sessions 4-5
haxby = load_haxby(subjects=-1, sessions=[4,5], targets=None)

# extract subjects, print some information about them
subjects = haxby.subjects
for i, subject in enumerate(subjects):
    print(i, subject.name, subject.data.shape, subject.target.shape)
```

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```

```

0 subj1_sess4_5 (242, 577) (242, 9)
1 subj2_sess4_5 (242, 464) (242, 9)
2 subj3_sess4_5 (242, 307) (242, 9)
3 subj4_sess4_5 (242, 675) (242, 9)
4 subj5_sess4_5 (242, 422) (242, 9)
5 subj6_sess4_5 (242, 348) (242, 9)

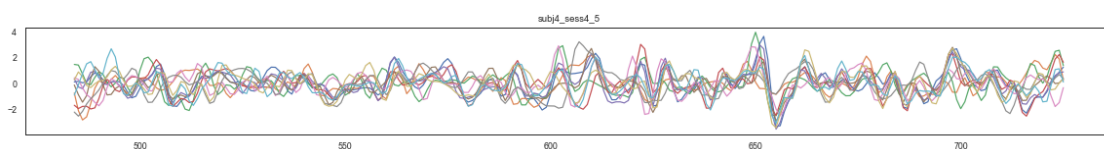
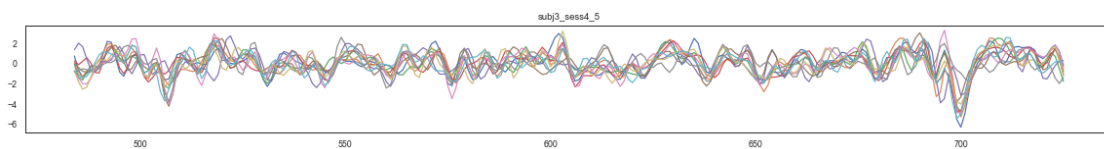
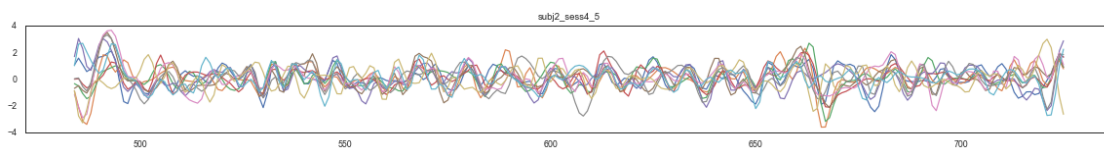
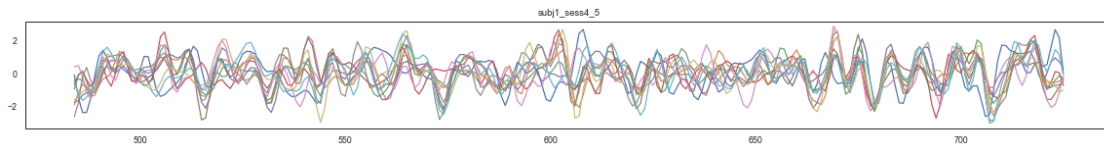
```

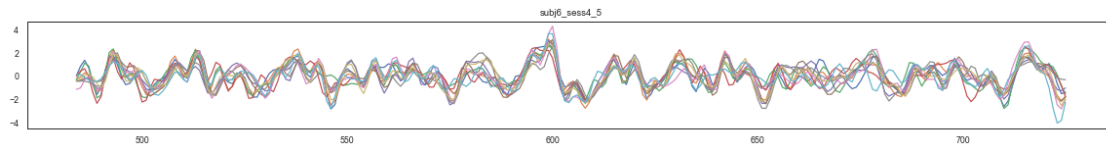
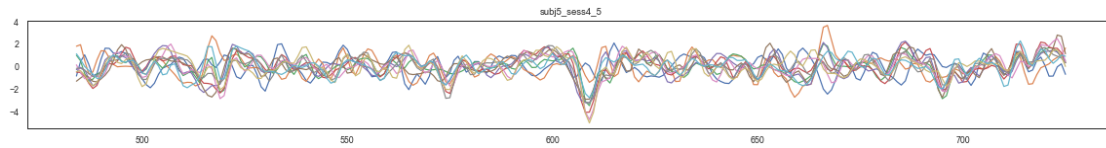
1.2.1 2.1 Visualize the data

```

In [9]: # inspect the first 10 voxels
for subject in haxby.subjects:
    _ = subject.data.iloc[:, :10].plot(
        figsize=(20, 2),
        legend=False,
        title=subject.name
    )

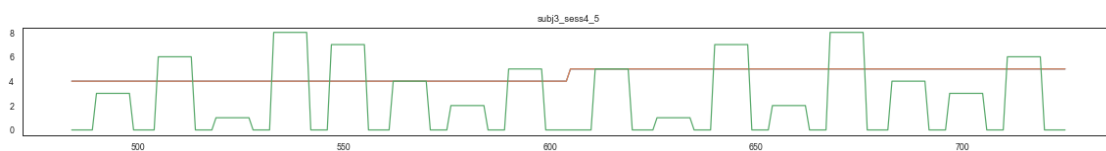
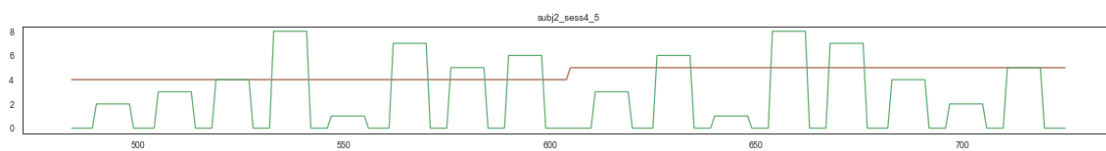
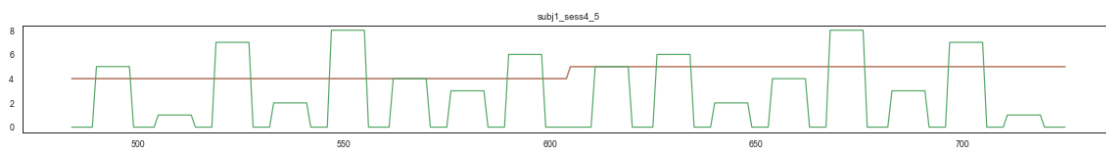
```

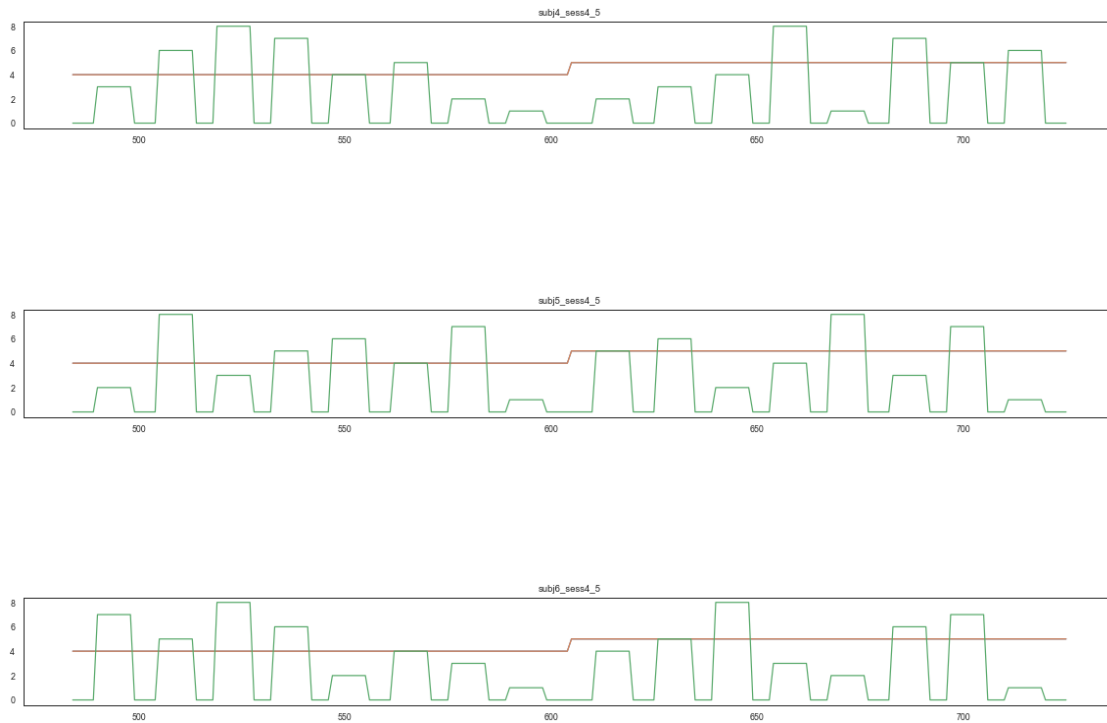




1.2.2 2.2 Visualize the meta data

```
In [10]: # inspect the first 10 voxels
for subject in haxby.subjects:
    _ = subject.meta.iloc[:, :10].plot(
        figsize=(20,2),
        legend=False,
        title=subject.name
    )
```





1.3 3 Generate a shape graph for each subject with knapper

```
In [11]: # loop over sessions, run for each
         for i, subject in enumerate(subjects):

             ### Setup: extract X, y for each subject
             X = subject.data.values.copy()
             y = subject.target.copy()
             y = y.reindex(columns=haxby.target_names)
             print('[subject {}]' .format(subject.name))
             print(' X has shape: {}'.format(X.shape))
             print(' y has shape: {}'.format(y.shape))

             ### Run knapper
             print("\nGenerating shape graph...")
             projection = TSNE(
                 perplexity=50, random_state=0,
                 init='pca', method='exact'
             )
             clusterer = optimize_dbscan(X)
             cover = optimize_cover(
                 X, r=15, g=0.67,
                 scale_r=not True,
```

```

        scale_limits=True
    )
    print('\n projection = {}'.format(projection))
    print('\n clusterer = {}'.format(clusterer))
    print('\n cover = {}'.format(cover.__dict__))

    # run kmapper
    mapper = KMapperWrapper(
        memory='dyneusr_cache/{}'.format(subject.name)
    )
    mapper.fit_lens(X, projection=projection)
    mapper.fit_graph(data=X, clusterer=clusterer, cover=cover)

    # extract results
    lens = mapper.lens_.copy()
    graph = dict(mapper.graph_)

    ### Fit DyNeuGraph
    print("\nProcessing shape graph...")
    dG = ds.DyNeuGraph(
        G=graph, y=y,
        labels=haxby.target_names,
        colors=haxby.target_colors,
        cmap=haxby.cmap
    )

    # store data
    subject.mapped = Bunch(
        data=X,
        lens=lens,
        graph=graph,
        dG=dG, X=X, y=y
    )
    subjects[i] = subject
    print()
    print()

```

```

[subject subj1_sess4_5]
X has shape: (242, 577)
y has shape: (242, 9)

```

Generating shape graph...

```

projection = TSNE(angle=0.5, early_exaggeration=12.0, init='pca', learning_rate=200.0,
    method='exact', metric='euclidean', min_grad_norm=1e-07, n_components=2,
    n_iter=1000, n_iter_without_progress=300, perplexity=50, random_state=0,
    verbose=0)

```

```
clusterer = DBSCAN(algorithm='auto', eps=26.3271381223901, leaf_size=15,
    metric='minkowski', metric_params=None, min_samples=2, n_jobs=1, p=2)

cover = {'n_cubes': 17, 'perc_overlap': 0.67, 'limits': array([[ -0.04466667,  1.04466667],
    [ -0.04466667,  1.04466667]])}
```

Processing shape graph...

<IPython.core.display.HTML object>

```
[subject subj2_sess4_5]
X has shape: (242, 464)
y has shape: (242, 9)
```

Generating shape graph...

```
projection = TSNE(angle=0.5, early_exaggeration=12.0, init='pca', learning_rate=200.0,
    method='exact', metric='euclidean', min_grad_norm=1e-07, n_components=2,
    n_iter=1000, n_iter_without_progress=300, perplexity=50, random_state=0,
    verbose=0)
```

```
clusterer = DBSCAN(algorithm='auto', eps=27.605001592992302, leaf_size=15,
    metric='minkowski', metric_params=None, min_samples=2, n_jobs=1, p=2)
```

```
cover = {'n_cubes': 17, 'perc_overlap': 0.67, 'limits': array([[ -0.04466667,  1.04466667],
    [ -0.04466667,  1.04466667]])}
```

Processing shape graph...

<IPython.core.display.HTML object>

```
[subject subj3_sess4_5]
X has shape: (242, 307)
y has shape: (242, 9)
```

Generating shape graph...

```
projection = TSNE(angle=0.5, early_exaggeration=12.0, init='pca', learning_rate=200.0,
    method='exact', metric='euclidean', min_grad_norm=1e-07, n_components=2,
    n_iter=1000, n_iter_without_progress=300, perplexity=50, random_state=0,
```

```
verbose=0)
```

```
clusterer = DBSCAN(algorithm='auto', eps=24.668255970446275, leaf_size=15,  
metric='minkowski', metric_params=None, min_samples=2, n_jobs=1, p=2)
```

```
cover = {'n_cubes': 17, 'perc_overlap': 0.67, 'limits': array([[ -0.04466667,  1.04466667],  
[ -0.04466667,  1.04466667]])}
```

Processing shape graph...

<IPython.core.display.HTML object>

```
[subject subj4_sess4_5]  
X has shape: (242, 675)  
y has shape: (242, 9)
```

Generating shape graph...

```
projection = TSNE(angle=0.5, early_exaggeration=12.0, init='pca', learning_rate=200.0,  
method='exact', metric='euclidean', min_grad_norm=1e-07, n_components=2,  
n_iter=1000, n_iter_without_progress=300, perplexity=50, random_state=0,  
verbose=0)
```

```
clusterer = DBSCAN(algorithm='auto', eps=28.004985102867675, leaf_size=15,  
metric='minkowski', metric_params=None, min_samples=2, n_jobs=1, p=2)
```

```
cover = {'n_cubes': 17, 'perc_overlap': 0.67, 'limits': array([[ -0.04466667,  1.04466667],  
[ -0.04466667,  1.04466667]])}
```

Processing shape graph...

<IPython.core.display.HTML object>

```
[subject subj5_sess4_5]  
X has shape: (242, 422)  
y has shape: (242, 9)
```

Generating shape graph...

```
projection = TSNE(angle=0.5, early_exaggeration=12.0, init='pca', learning_rate=200.0,  
method='exact', metric='euclidean', min_grad_norm=1e-07, n_components=2,
```



```
n_iter=1000, n_iter_without_progress=300, perplexity=50, random_state=0,
verbose=0)
```

```
clusterer = DBSCAN(algorithm='auto', eps=23.616781132201332, leaf_size=15,
metric='minkowski', metric_params=None, min_samples=2, n_jobs=1, p=2)
```

```
cover = {'n_cubes': 17, 'perc_overlap': 0.67, 'limits': array([[ -0.04466667,  1.04466667],
[ -0.04466667,  1.04466667]])}
```

Processing shape graph...

<IPython.core.display.HTML object>

```
[subject subj6_sess4_5]
X has shape: (242, 348)
y has shape: (242, 9)
```

Generating shape graph...

```
projection = TSNE(angle=0.5, early_exaggeration=12.0, init='pca', learning_rate=200.0,
method='exact', metric='euclidean', min_grad_norm=1e-07, n_components=2,
n_iter=1000, n_iter_without_progress=300, perplexity=50, random_state=0,
verbose=0)
```

```
clusterer = DBSCAN(algorithm='auto', eps=20.11123226958869, leaf_size=15,
metric='minkowski', metric_params=None, min_samples=2, n_jobs=1, p=2)
```

```
cover = {'n_cubes': 17, 'perc_overlap': 0.67, 'limits': array([[ -0.04466667,  1.04466667],
[ -0.04466667,  1.04466667]])}
```

Processing shape graph...

<IPython.core.display.HTML object>

1.4 4 Visualize and annotate the shape graph with dyneusr

```
In [12]: # select subject to inspect below
subject = subjects[0]
```

```

# copy inputs to Mapper
data = subject.data.copy()
meta = subject.meta.copy()

# copy outputs from Mapper
X = subject.mapped.X.copy()
y = subject.mapped.y.copy()
lens = subject.mapped.lens.copy()
dG = subject.mapped.dG

```

1.4.1 4.1 Visualize the lens

```

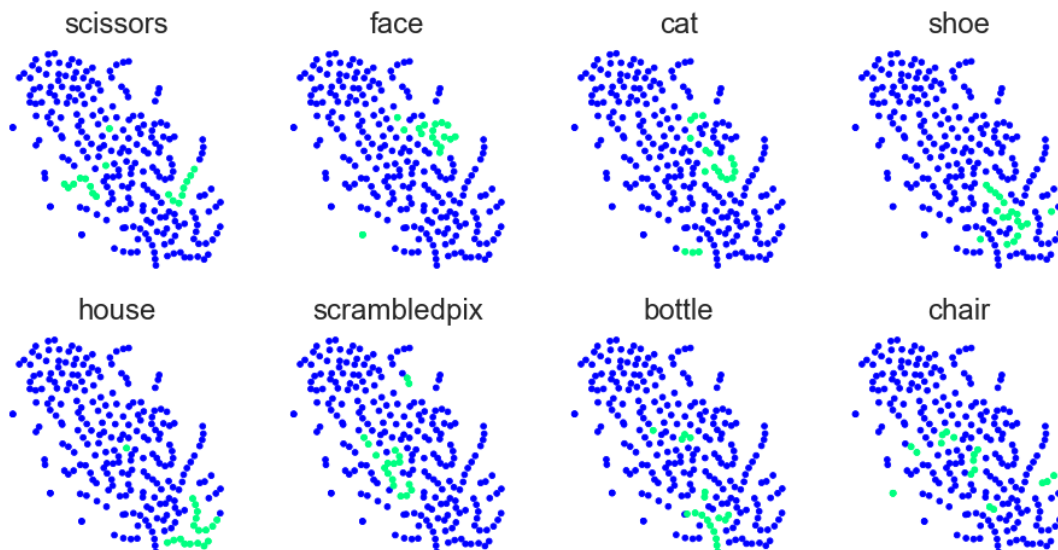
In [13]: ### View some quick results (e.g., the lens)
fig, axes = plt.subplots(2, 4, figsize=(16, 8))

# loop over axis for each category of visual stimuli
c_cols = [_ for _ in y.columns if _ not in ['rest']]
for ax, stim in zip(np.ravel(axes), c_cols):

    # mask the stimuli for this category
    y_mask = (y[stim] == 1)

    # plot the lens, coloring only the category == stim
    ax.scatter(*lens.T,
               c=y[stim], cmap='winter')
    ax.scatter(*lens[y_mask].T,
               c=y[y_mask][stim], cmap='winter_r')
    ax.set_title(stim, fontsize=24)
    ax.axis('off')

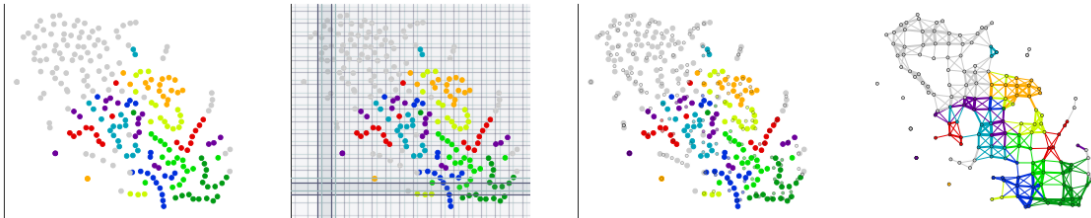
```



1.4.2 4.2 Visualize the stages of Mapper

```
In [14]: # we need to set y to a coloring
subject.y = subject.meta.target.values.copy()

# draw Mapper stages (and intermediates)
_ = ds.tools.networkx_utils.visualize_mapper_stages(
    data=subject, lens=lens, cover=cover, graph=graph, dG=dG,
    node_size=10,
)
```



1.4.3 4.3 Visualize the shape graph with dyneusr

```
In [15]: dG.visualize(
    'dyneusr_output_haxby_{}.html'.format(subject.name),
    show=True, port=8802
)
```

Already serving localhost:8802

[Force Graph] http://localhost:8802/dyneusr_output_haxby_subj1_sess4_5.html

<IPython.core.display.HTML object>

<IPython.lib.display.IFrame at 0x11e2c57f0>

Out[15]: DyNeuGraph()

<Figure size 432x288 with 0 Axes>

1.4.4 4.4 Qualitative analysis

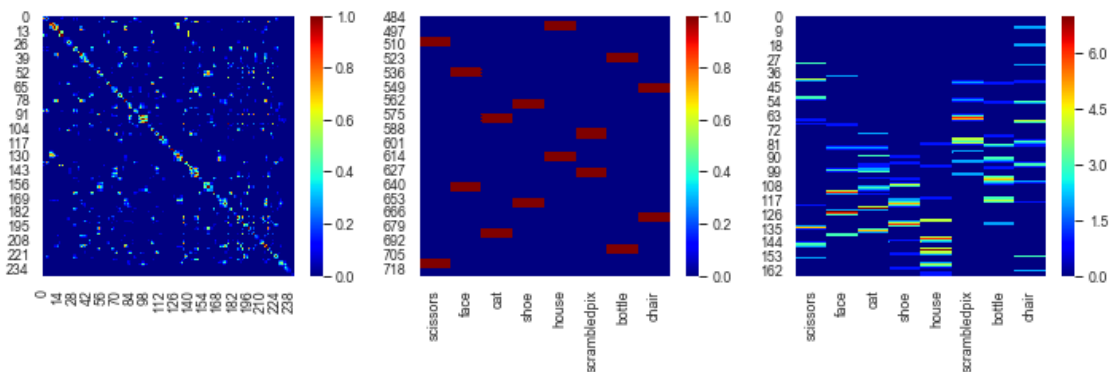
```
In [16]: # define TR x Node, TR x TR, and TR x Stimuli matrices
TxN = dG.M.copy()
TxT = dG.TCM.copy()
TxS = y.copy().iloc[:, 1:]

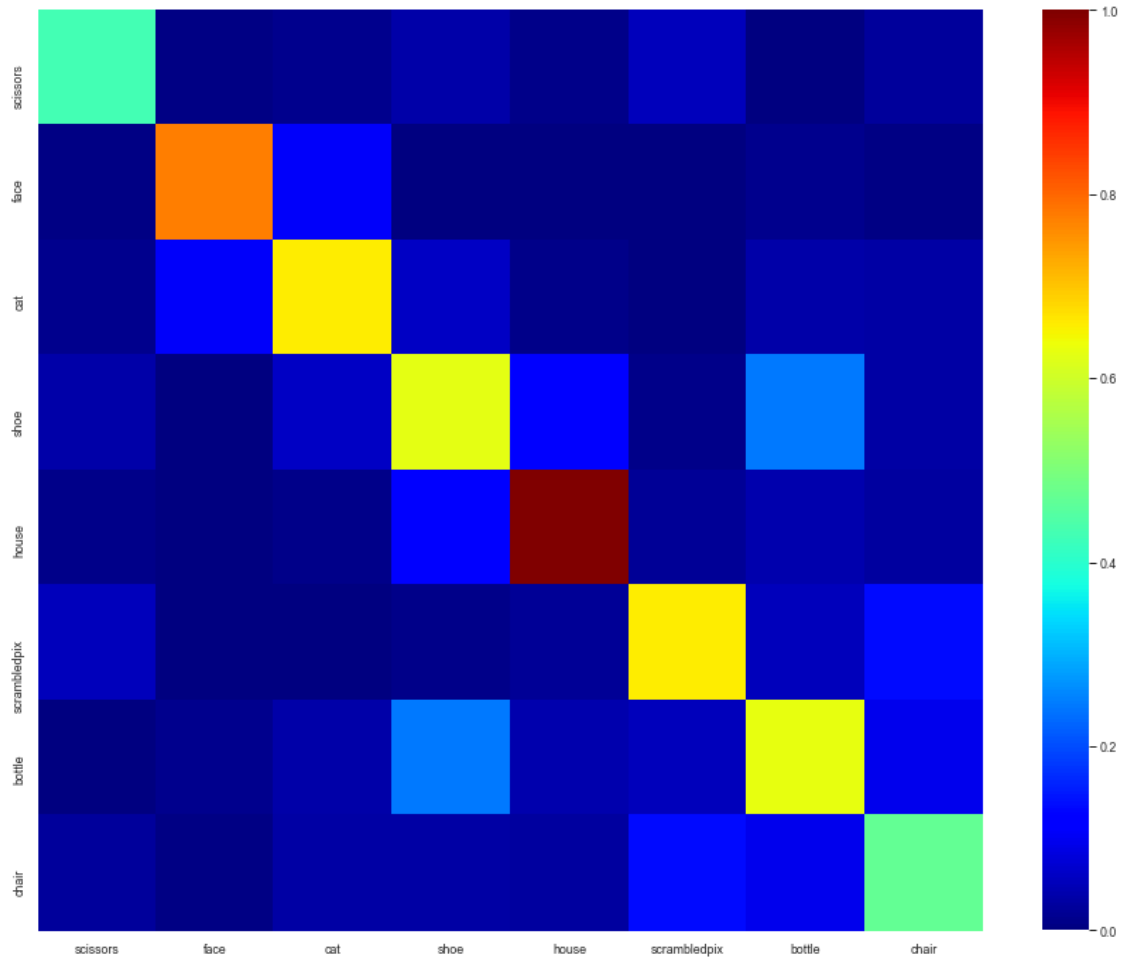
# Compute Node x Stimuli, Stimuli x Stimuli
NxS = TxN.T @ TxS
SxS = NxS.T @ NxS
SxS /= SxS.max()

# plot the heatmaps
fig, axes = plt.subplots(1, 3, figsize=(12, 3))
sns.heatmap(TxT, ax=axes[0], cmap='jet')
sns.heatmap(TxS, ax=axes[1], cmap='jet', xticklabels=TxS.columns)
sns.heatmap(NxS, ax=axes[2], cmap='jet', xticklabels=TxS.columns)

# plot the heatmap of category similarity
fig, axes = plt.subplots(1, 1, figsize=(15, 12))
axes = np.ravel(axes)
sns.heatmap(SxS, ax=axes[0], cmap='jet',
            xticklabels=TxS.columns, yticklabels=TxS.columns)
```

Out[16]: <matplotlib.axes._subplots.AxesSubplot at 0x120c86978>





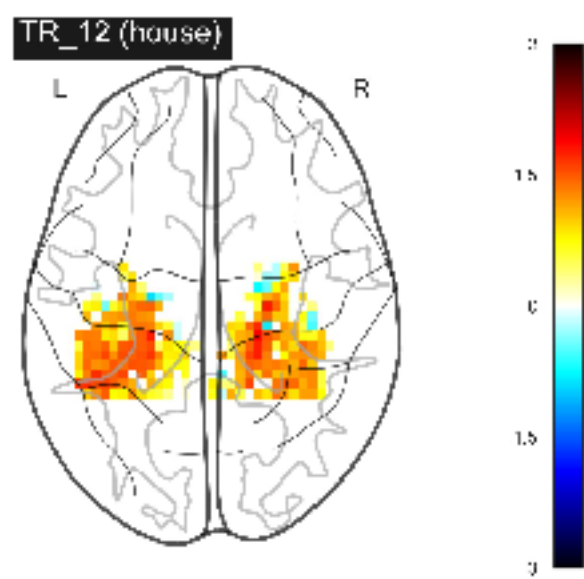
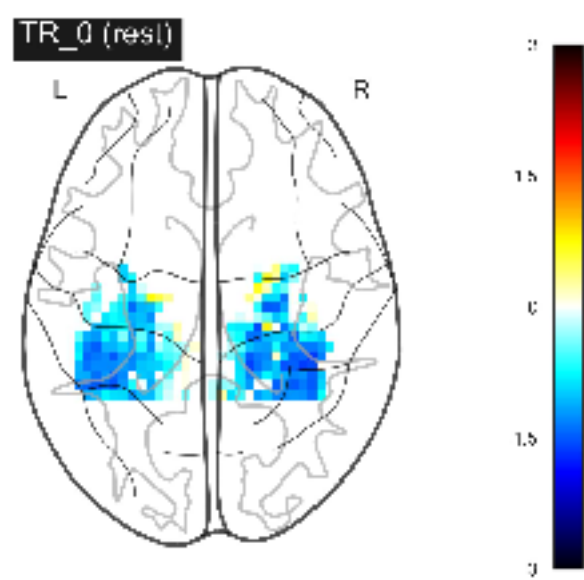
1.5 5 Anchor the shape graph to brain anatomy with dyneusr

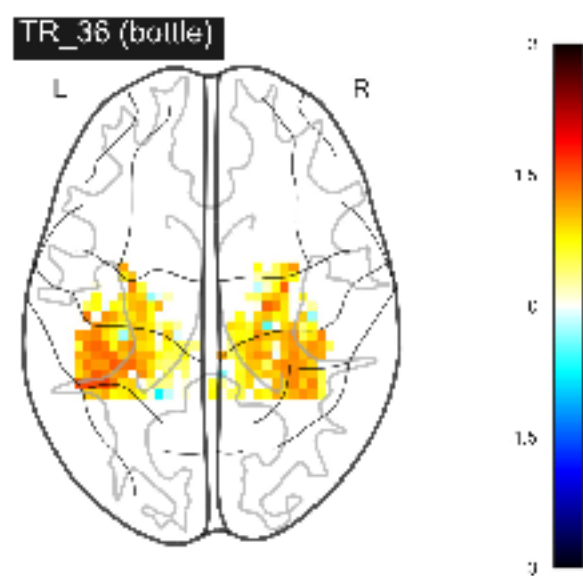
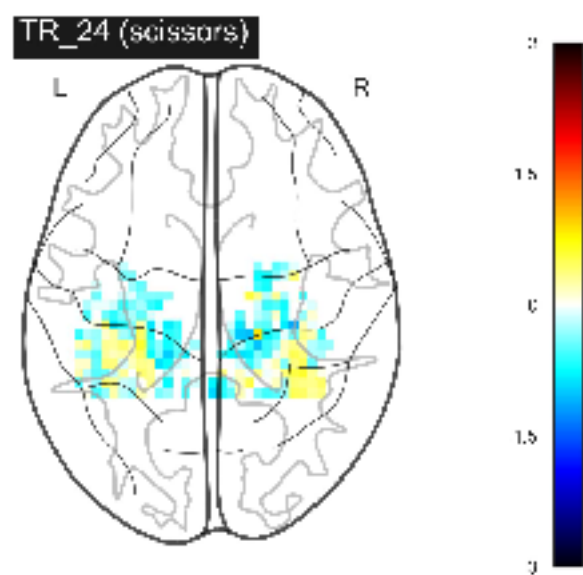
1.5.1 5.1 Estimate average brain images for each time frame

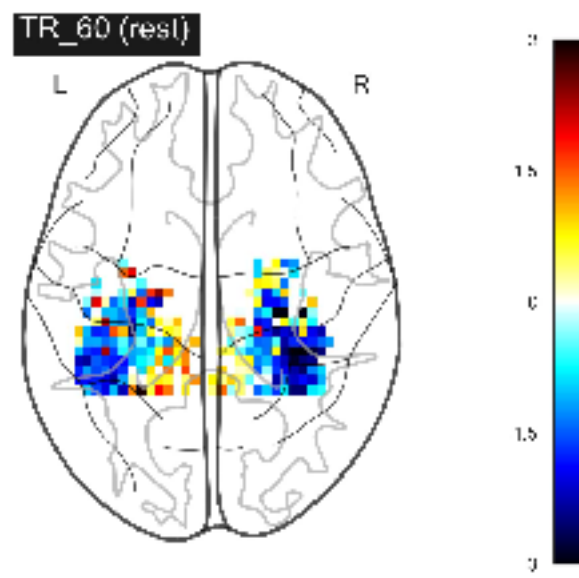
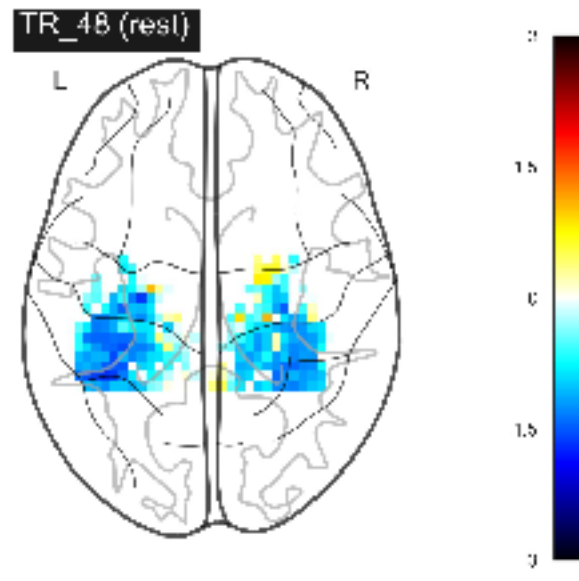
```
In [17]: # get mixtures (time frames to average over)
mixtures = dG.mixtures_

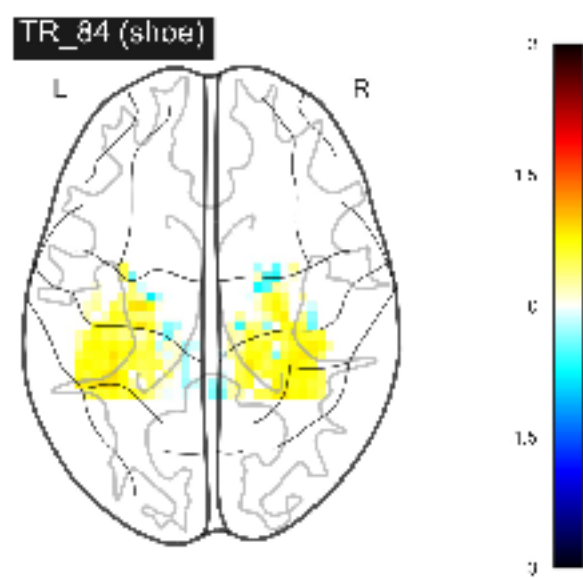
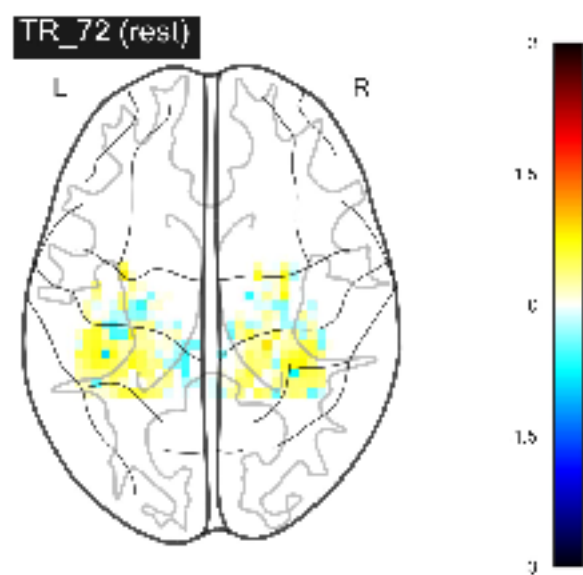
# estimate and plot average brain images
brain_images = ds.tools.mixture.simple_mixtures(
    subject, mixtures=mixtures,
    save_dir='tooltips_{}'.format(subject.name),
    prefix='TR_', targets=subject.meta.labels.tolist(),
    show_every_n=12, print_every_n=0,
    threshold=None, fwhm=None,
    kind='mean', mode='glass', figsize=(4, 4),
    plot_kws=dict(
        display_mode='z', cut_coords=1,
        vmin=-3, vmax=3, threshold=0, colorbar=True,
```

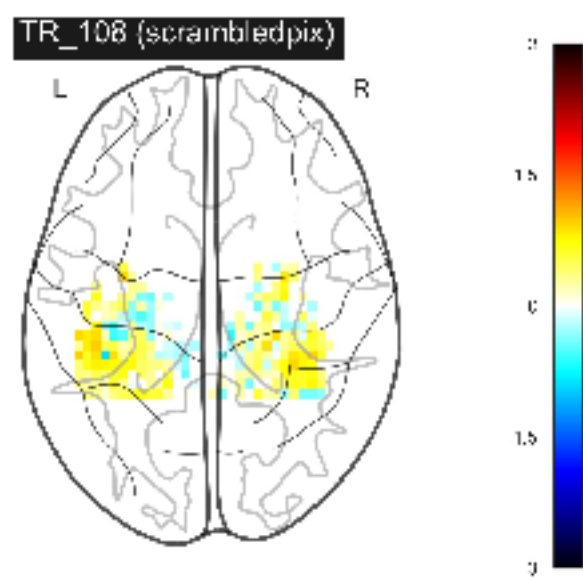
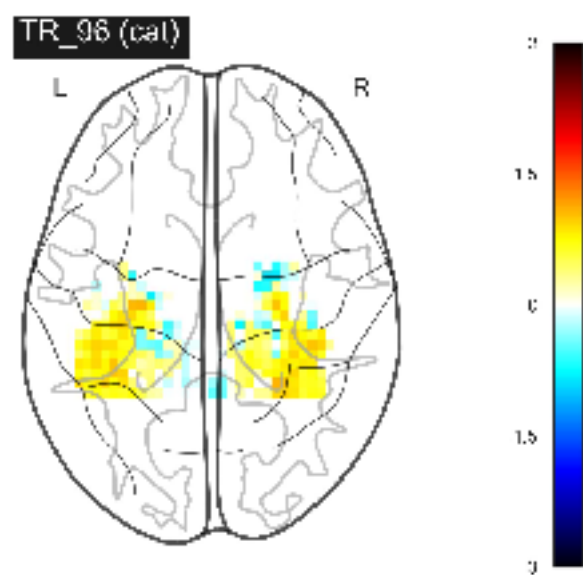
),

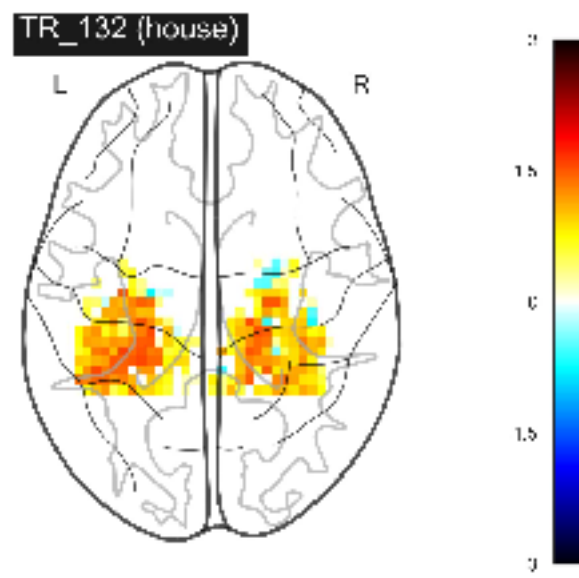
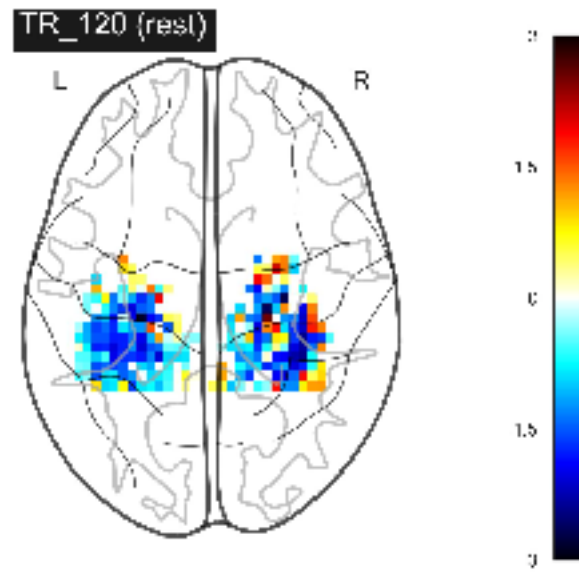




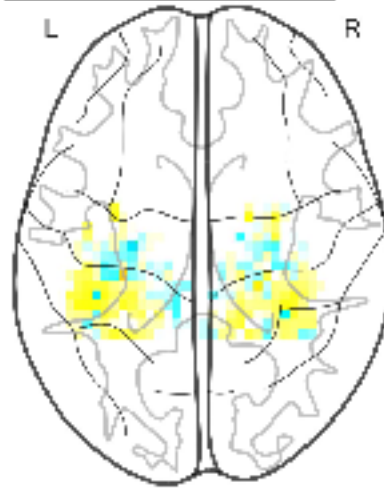




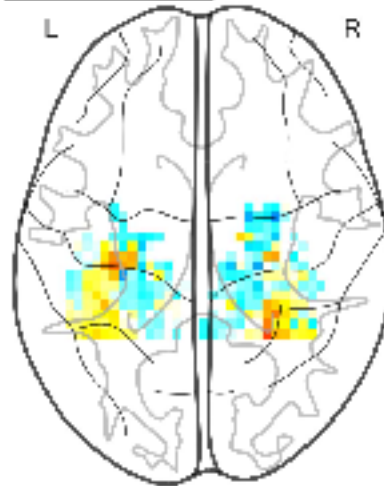


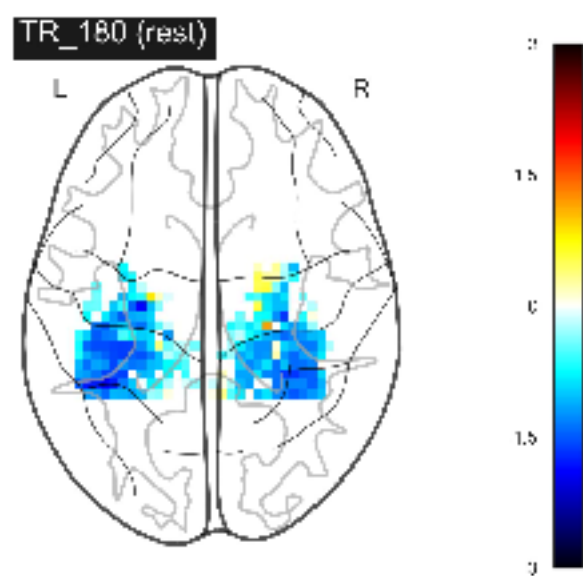
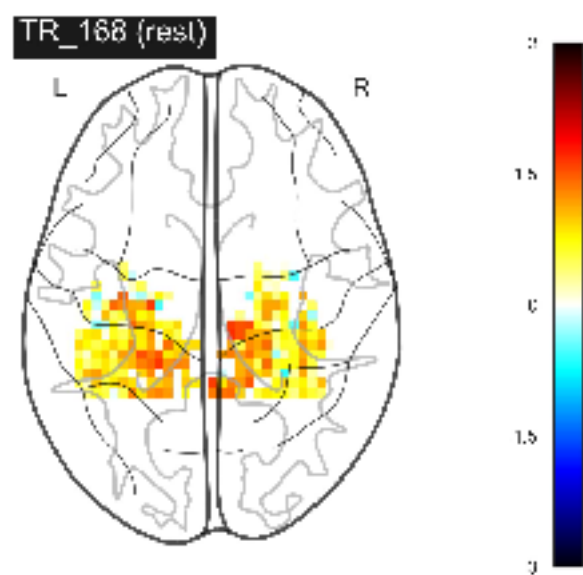


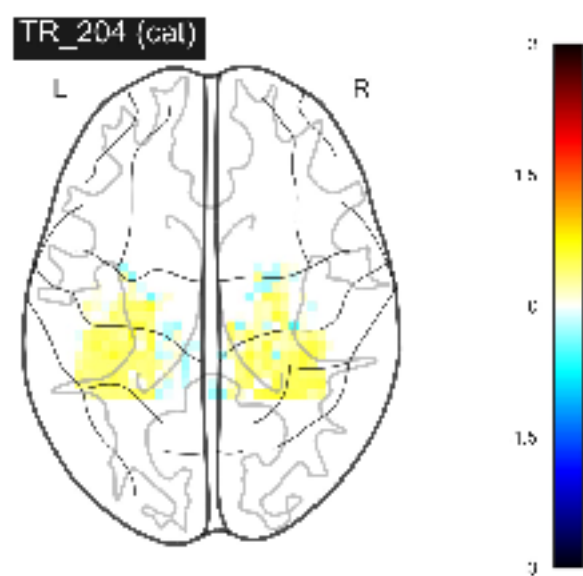
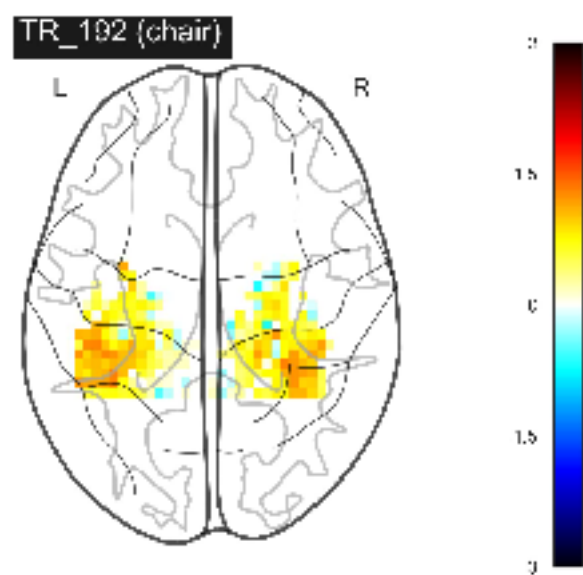
TR_144 (scrambledpix)

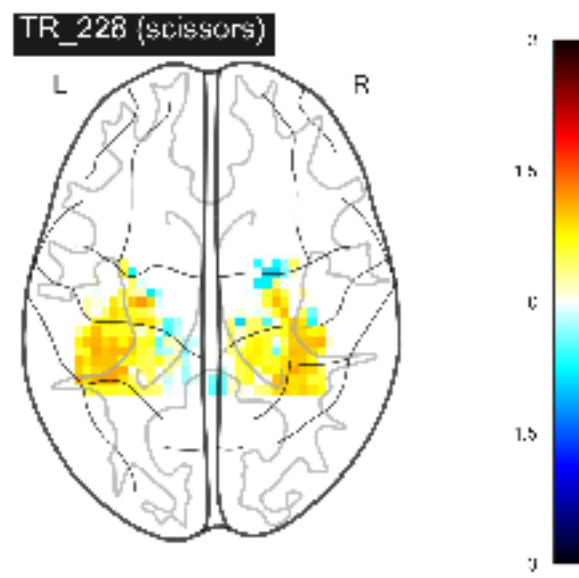
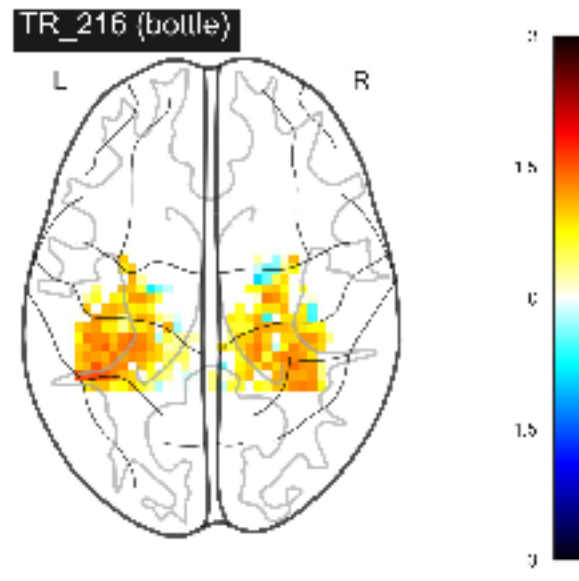


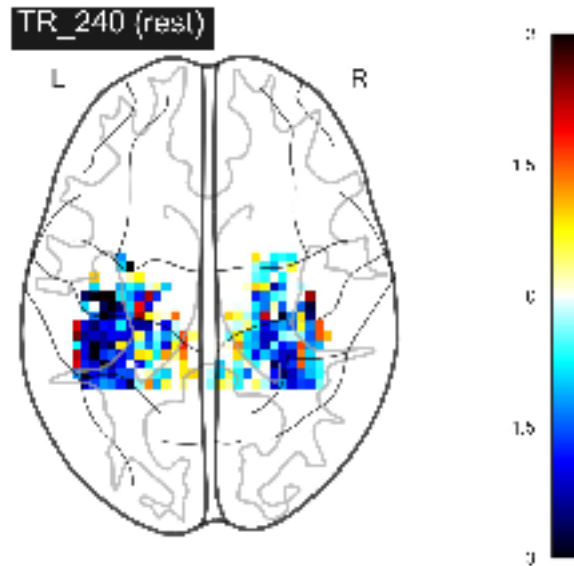
TR_156 (face)











[done]

1.5.2 5.2 Annotate nodes in the shape graph with the brain images

```
In [18]: _ = dG.annotate(image=brain_images)
```

1.5.3 5.3 Visualize the shape graph with dyneusr (using the d3-force layout)

```
In [19]: dG.visualize(
    'dyneusr_output_haxby_anat_{}.html'.format(subject.name),
    show=True, port=8802
)
```

Already serving localhost:8802

[Force Graph] http://localhost:8802/dyneusr_output_haxby_anat_subj1_sess4_5.html

<IPython.core.display.HTML object>

<IPython.lib.display.IFrame at 0x120a7c6d8>

Out[19]: DyNeuGraph()

<Figure size 432x288 with 0 Axes>

1.6 6 Capture temporal transitions in brain activity with dyneusr

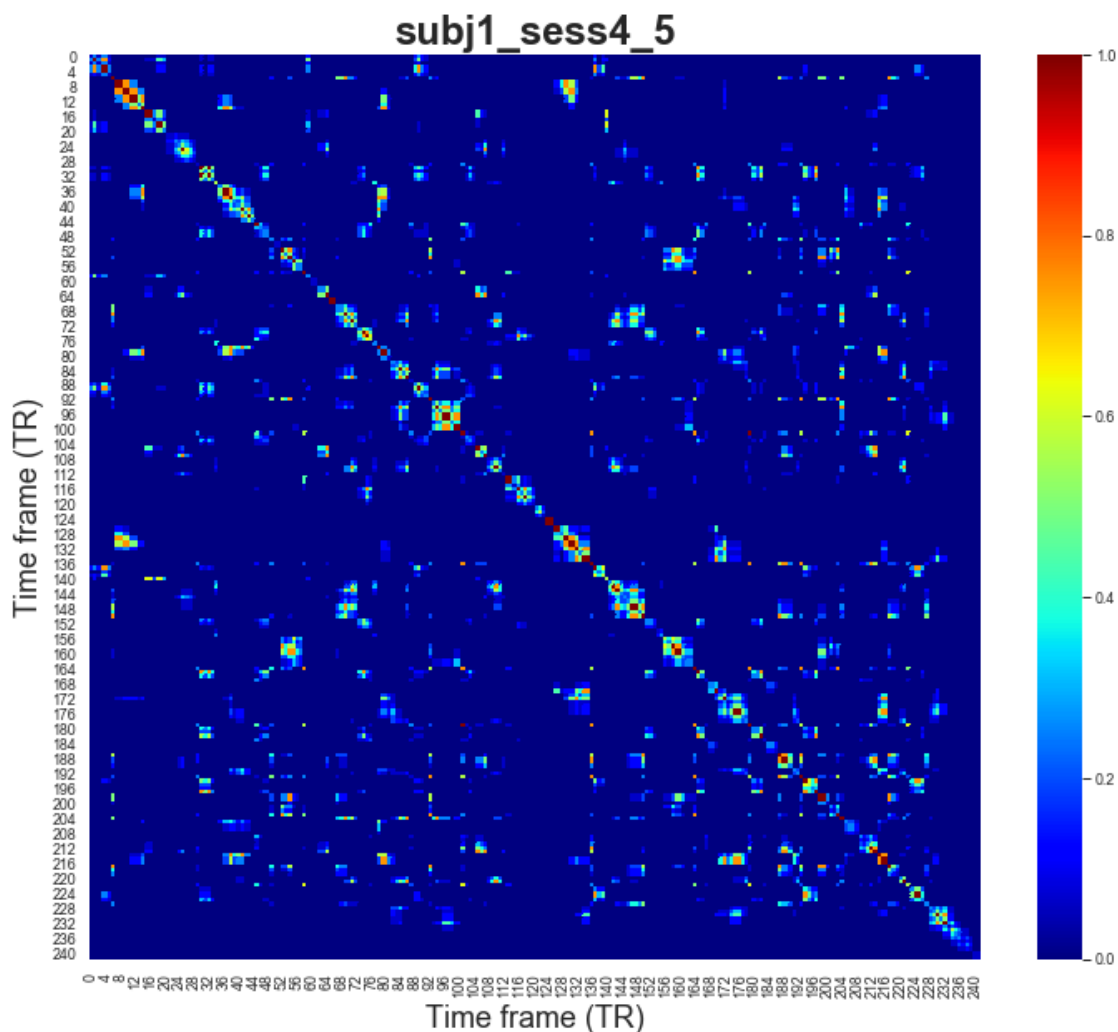
```
In [20]: # aggregate the TCMs into a single matrix
TCMs = np.array([_.mapped.dG.TCM.copy() for _ in subjects])
print("TCMs has shape:", TCMs.shape)
```

TCMs has shape: (6, 242, 242)

1.6.1 6.1 Visualize the temporal connectivity matrix (TCM)

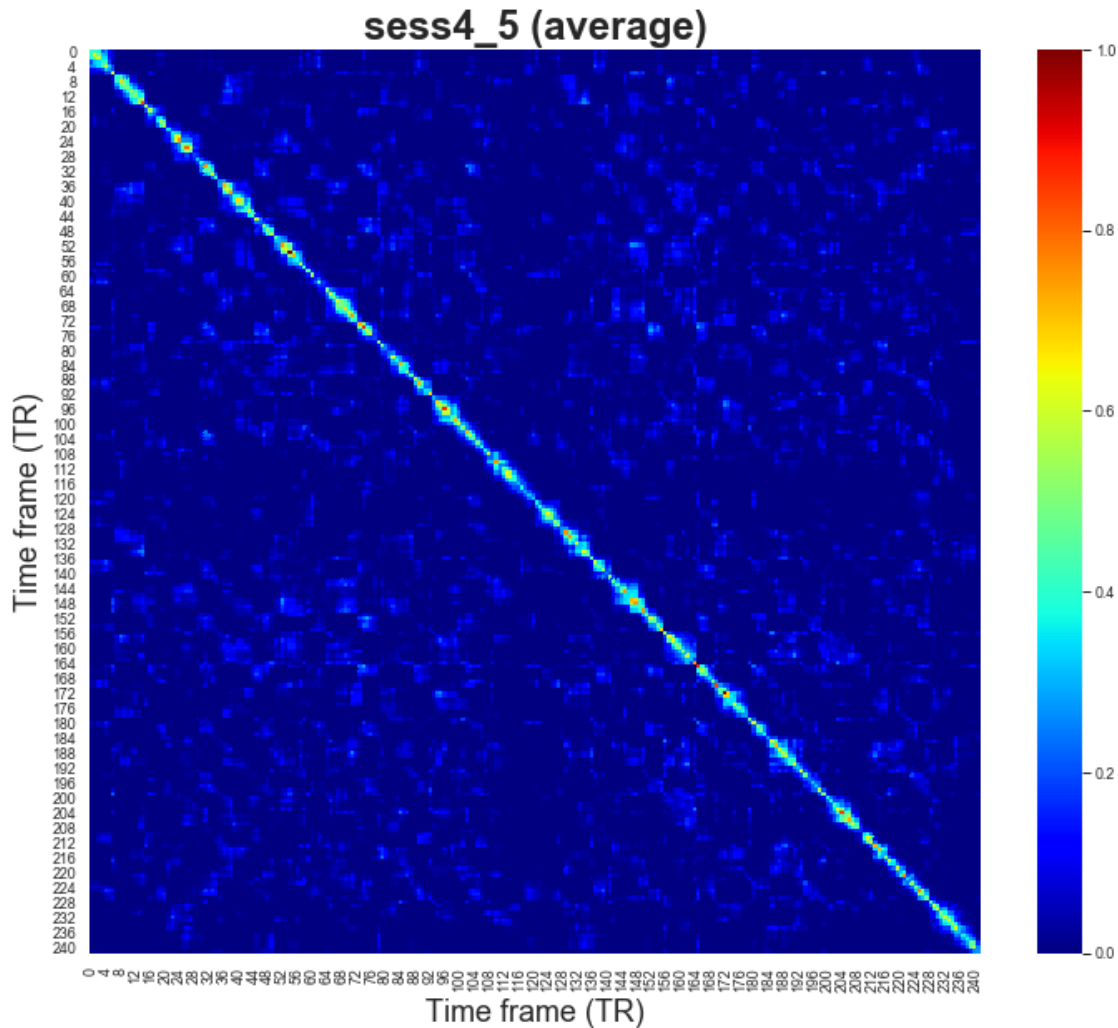
```
In [21]: sns.heatmap(dG.TCM, cbar=True, cmap='jet', figure=plt.figure(figsize=(12,10)))
plt.title('{}'.format(subject.name), fontsize=24, fontweight='bold')
plt.xlabel('Time frame (TR)', fontsize=18)
plt.ylabel('Time frame (TR)', fontsize=18)
```

Out[21]: Text(86.7,0.5,'Time frame (TR)')



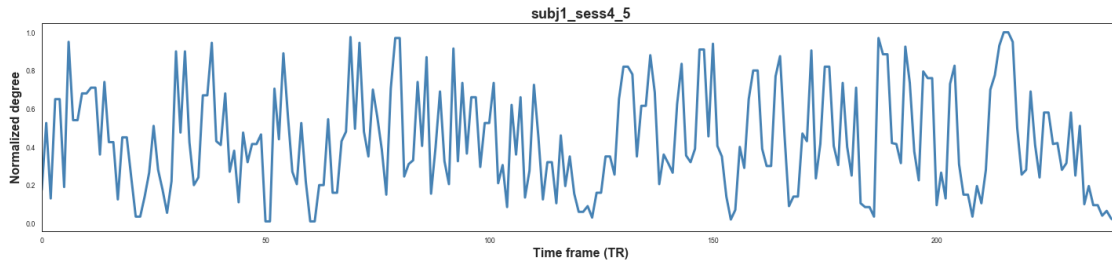
```
In [22]: sns.heatmap(TCMs.mean(axis=0), cbar=True, cmap='jet', figure=plt.figure(figsize=(12,12),
plt.title('{} (average)'.format(haxby.session_code), fontsize=24, fontweight='bold')
plt.xlabel('Time frame (TR)', fontsize=18)
plt.ylabel('Time frame (TR)', fontsize=18)
```

```
Out[22]: Text(86.7,0.5,'Time frame (TR)')
```

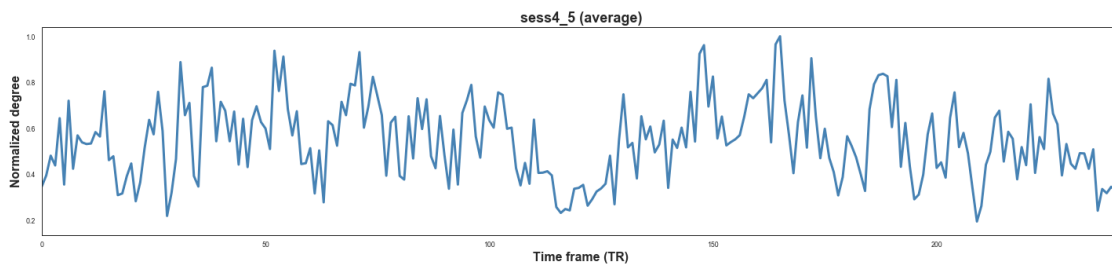


1.6.2 6.2 Visualize the normalized degree of nodes in the TCM over time

```
In [23]: ds.visuals.plot_temporal_degree(dG.TCM, title=subject.name)
```



```
In [24]: ds.visuals.plot_temporal_degree(TCMs.mean(axis=0), title='{} (average)'.format(haxby.))
```



1.6.3 6.3 Visualize the degree of TCM for a single stimulus category

```
In [25]: # define subject, category
subject = subjects[0]
target_name = 'face'

# mask target
target = subject.target.copy().reset_index(drop=True)
target = target.assign(data_id = target.index)
meta = subject.meta.copy().reset_index(drop=True)

# tcms
tcms = []
for session in [4, 5]:

    # define mask for each session
    target_mask = target[target_name].eq(1)
    session_mask = meta.session.eq(session)
    target_index = target[(target_mask & session_mask)].index.values
    target_bounds = np.min(target_index), np.max(target_index)

    # include 5 rest beforeblock
    rest_before = (target.loc[target['rest'].eq(1)]
                  .data_id.ge(target_bounds[0]-5))
```

```

rest_after = (target.loc[target['rest'].eq(1)]
               .data_id.le(target_bounds[1]+5))
rest_mask = (rest_before & rest_after)

# mask targets for session, and rest before, after block
target_mask = ((target_mask & session_mask) | (rest_mask & session_mask))
target_mask_ix = np.ix_(target_mask, target_mask)
trs = target_mask[target_mask].iloc[:, :].index.values

# print some info
print("Extracting TCM for category={}, session={}, onset={}, offset={}".format(
    target_name, session, trs[0]+5, trs[-1]-5
))

# extract tcm
tcm = subject.mapped.dG.TCM.copy()
tcm = tcm[target_mask_ix]

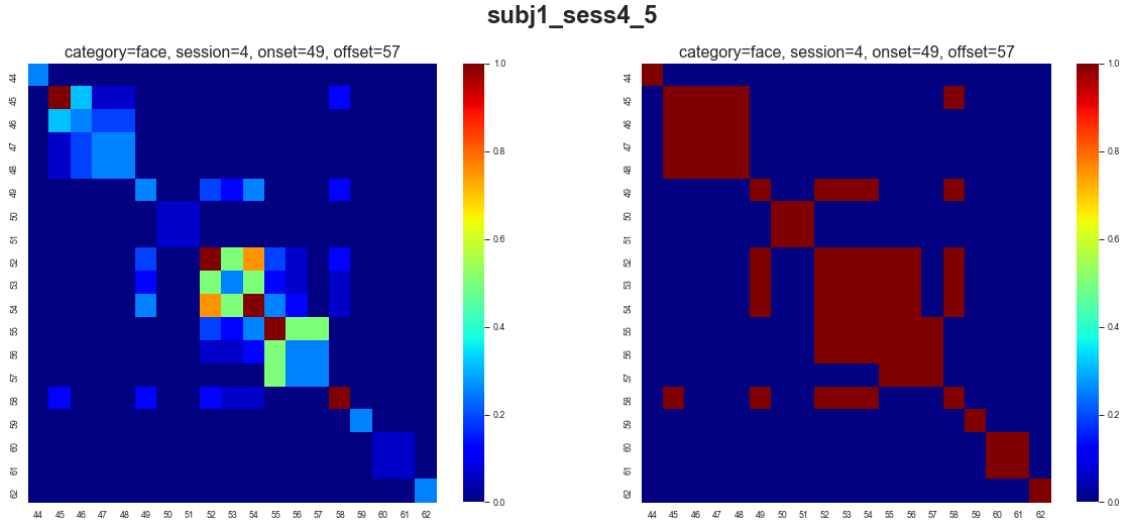
# convert to dataframe, save
df_tcm = pd.DataFrame(tcm, columns=trs, index=trs)
tcms.append(df_tcm)

# plot heatmap
figure, axes = plt.subplots(1, 2, figsize=(20,8))
ax = sns.heatmap(df_tcm, cbar=True, cmap='jet', ax=axes[0])
ax = sns.heatmap(df_tcm>0, cbar=True, cmap='jet', axes=axes[1])
for ax in axes:
    ax.set_title("category={}, session={}, onset={}, offset={}".format(
        target_name, session, trs[0]+5, trs[-1]-5
    ), fontsize=16)
plt.suptitle(subject.name, fontsize=24, fontweight='bold')
plt.show()

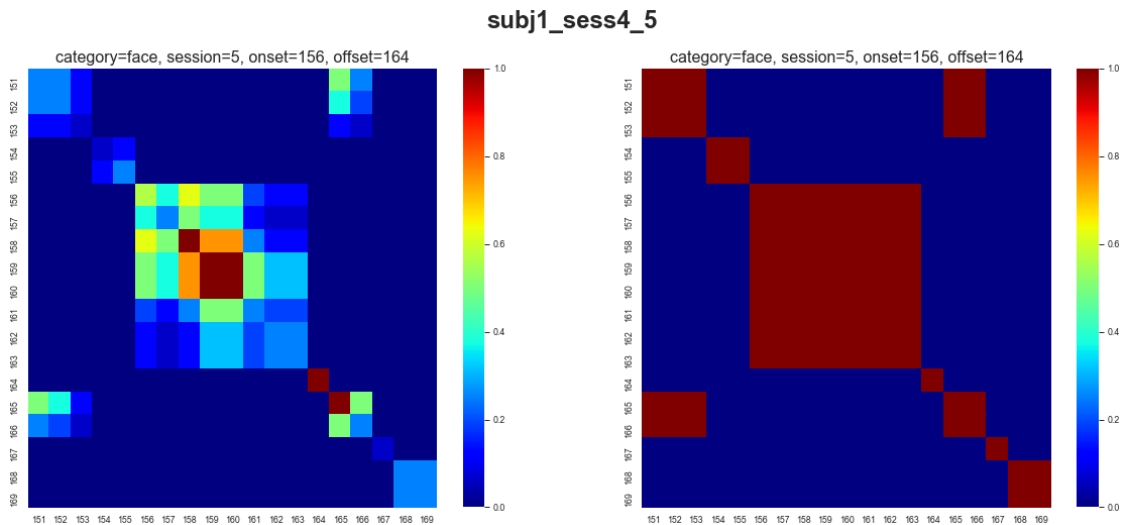
# convert to array
tcms = np.array([_.values for _ in tcms])

```

Extracting TCM for category=face, session=4, onset=49, offset=57



Extracting TCM for category=face, session=5, onset=156, offset=164



```
In [26]: # plot heatmap
figure, axes = plt.subplots(1, 2, figsize=(20,8))
ax = sns.heatmap(tcms.mean(0), cbar=True, cmap='jet', ax=axes[0])
ax = sns.heatmap(tcms.mean(0)>0, cbar=True, cmap='jet', axes=axes[1])
for ax in axes:
    ax.set_title("category={}, session={}, onset={}, offset={}".format(
        target_name, session, trs[0]+5, trs[-1]-5
    ), fontsize=16)
```

```
plt.suptitle(haxby.session_code, fontsize=24, fontweight='bold')
plt.show()
```

sess4_5

