Example of identifying relationships between neural activity and movement behavior in AJILE12

This script loads data directly from the DANDI remote repository using JupyterHub (https://hub.dandiarchive.org/ (https://hub.dandiarchive.org/)). It can be adjusted to work on downloaded AJILE12 data files stored locally by using the commented out line in the cell after 'Load the data file' and commenting out all preceding lines in the cell.

In [1]:

```
# Uncomment and run this cell if using JupyterHub
# %pip install natsort
# %pip install seaborn
# %pip install nilearn
# %pip install dandi
# %pip install nwbwidgets==0.7.0
# %pip install git+https://github.com/catalystneuro/brunton-lab-to-nwb.git
```

In [2]:

```
import seaborn as sns
import numpy as np
import pandas as pd
import statsmodels.api as sm
from scipy.signal import sosfiltfilt, butter, hilbert

from pynwb import NWBHDF5IO
from dandi.dandiapi import DandiAPIClient
from ndx_events import LabeledEvents, AnnotatedEventsTable, Events
```

User-defined parameters

These parameters can be adjusted to analyze other electrodes, frequency bands, behavior types, participants, sessions, etc.

We will select data from participant 1, session 3 only during times that the participant was eating (because this is an active behavior). ECoG data will be converted to spectral power in the gamma band (80-100 Hz) for electrode 7, which is located over the motor cortex. We will also look at the vertical velocity of the right wrist.

about:srcdoc Page 1 of 9

In [3]:

```
sbj, session = 1, 3
behavior_type = 'Eat'
neural_freq_range = [80, 100] # Hz
ecog_ch_num = 7
keypoint_of_interest = 'R_Wrist'
pose_direction = 'vertical' # 'vertical' or 'horizontal'
```

Load the data file

Data is loaded in directly from DANDI remote repository.

In [4]:

```
with DandiAPIClient() as client:
    asset = client.get_dandiset("000055").get_asset_by_path(
        "sub-{0:>02d}/sub-{0:>02d}_ses-{1:.0f}_behavior+ecephys.nwb".format(sbj,
session)
    )
    s3_path = asset.get_content_url(follow_redirects=1, strip_query=True)

io = NWBHDF5IO(s3_path, mode='r', load_namespaces=True, driver='ros3')
# io = NWBHDF5IO(local_file_path, mode='r', load_namespaces=False)
nwb = io.read()
```

/opt/conda/lib/python3.9/site-packages/hdmf/spec/namespace.py:532: U serWarning: Ignoring cached namespace 'hdmf-common' version 1.4.0-al pha because version 1.5.0 is already loaded.

warn("Ignoring cached namespace '%s' version %s because version %s
is already loaded."

/opt/conda/lib/python3.9/site-packages/hdmf/spec/namespace.py:532: U serWarning: Ignoring cached namespace 'core' version 2.2.5 because v ersion 2.4.0 is already loaded.

warn("Ignoring cached namespace '%s' version %s because version %s
is already loaded."

Print information about selected electrode

about:srcdoc Page 2 of 9

```
In [5]:

nwb.electrodes[ecog_ch_num]

Out[5]:

x y z imp location filtering group ç

id

7 -53.896469 -29.059873 62.709102 NaN unknown lowpass pynwb.ecephys.ElectrodeGroup at 0x1402148...
```

Data pre-processing

Identify the start and stop times when the behavioral label of interest occurs.

```
In [6]:
```

```
min_len = 100 # (sec) only keep times when the given label appears for longer t
han this amount of time at once

coarse_labels = nwb.intervals['epochs'].to_dataframe()
coarse_labels = coarse_labels[coarse_labels['labels'].str.contains(behavior_type
)]
coarse_labels['diff'] = coarse_labels['stop_time'] - coarse_labels['start_time']
coarse_labels = coarse_labels[coarse_labels['diff'] > min_len]
coarse_labels.reset_index(inplace=True, drop=True)
```

Load the corresponding ECoG data for each behavioral label chunk and convert to spectral power via the Hilbert transform.

about:srcdoc Page 3 of 9

In [7]:

```
filter order = 4 # order of butterworth filter used to bandpass filter the ECoG
neural data = nwb.acquisition['ElectricalSeries'].data
sampling rate = nwb.acquisition['ElectricalSeries'].rate # (Hz) ECoG sampling r
ate
neural power = []
for i in range(coarse labels.shape[0]):
    # Identify the start/end indices for each continuous chunk of the given beha
vioral label
    start t = int(coarse labels.loc[i, 'start time']*sampling rate)
    end t = int(coarse labels.loc[i, 'stop time']*sampling rate)
    # Load data snippet
   neur data curr = neural data[start t:end t, ecog ch num]
    # Bandpass filter
    sos = butter(filter order, neural freq range, btype='bandpass', output='sos'
, fs=sampling rate)
    neur data filtered = sosfiltfilt(sos, neur data curr)
    # Apply Hilbert transform and convert to decibels
    neur pow = np.abs(hilbert(neur data filtered))
    neur pow = 10*np.log(neur pow)
    # Take the difference between neighboring timepoints
    neural power.append(np.diff(neur pow))
```

Load the corresponding pose data for each behavioral label chunk and convert to vertical velocity.

about:srcdoc Page 4 of 9

In [8]:

```
keypoints = list(nwb.processing['behavior'].data_interfaces['Position'].spatial_
series.keys())
assert keypoint of interest in keypoints
assert pose direction in ['vertical', 'horizontal']
keypoint series = nwb.processing['behavior'].data interfaces['Position'].spatial
series[keypoint of interest]
sampling rate keypoint = keypoint series.rate # Hz
keypoint velocity = []
for i in range(coarse labels.shape[0]):
    start_t = int(coarse_labels.loc[i, 'start_time']*sampling_rate_keypoint)
    end t = int(coarse labels.loc[i, 'stop time']*sampling rate keypoint)
    # Load pose data snippet
    pose data curr = keypoint series.data[start t:end t, :]
    pose mag curr = pose_data_curr[:, 1 if pose_direction == 'vertical' else 0]
    # Convert to velocity (delta X / delta t)
    velocity curr = np.diff(pose mag curr)/(1/sampling rate keypoint)
    keypoint velocity.append(velocity curr)
```

Align and combine neural and pose data into a pandas dataframe

about:srcdoc Page 5 of 9

In [9]:

```
assert len(neural power) == len(keypoint velocity)
measures all = []
for i in range(len(neural power)):
    # Neural power for the given chunk
    neur curr = neural power[i]
    l neur = len(neur curr)
    # Pose velocity for the given chunk
    accel curr = keypoint velocity[i]
    l accel = len(accel curr)
    # Downsample neural data to match pose data
    inds split = np.array split(np.arange(l neur), l accel)
    for j, inds in enumerate(inds split):
        measures all.append([neur curr[inds].mean(), accel curr[j]])
# Combine neural/pose data into a dataframe
df measures all = pd.DataFrame(np.asarray(measures all), columns=['Neural power
(dB)', 'Keypoint velocity (pixels/sec)'])
# Remove any NaN's
df measures all.dropna(inplace=True)
# Remove instances with velocity close to 0
df measures all = df measures all[(df measures all['Keypoint velocity (pixels/se
(c)'] > 100) | 
                                  (df measures all['Keypoint velocity (pixels/se
(c)'] < -100)]
```

Results

Print correlation between neural power and keypoint velocity

We find a small, positive correlation between neural power in the gamma band and right wrist vertical velocity.

about:srcdoc Page 6 of 9

In [10]:

```
df_measures_all.corr(method='pearson')
```

Out[10]:

Neural power (dB) Keypoint velocity (pixels/sec)

Neural power (dB)	1.000000	0.025379
Keypoint velocity (pixels/sec)	0.025379	1.000000

Perform robust linear regression to quantify any linear relationships

Regression identifies a small, but significant (p<0.05) positive relationship between neural power in the gamma band and right wrist vertical velocity. This result makes sense because moving one's wrist upward takes more effort (fighting against gravity) than moving one's arm downward and thus may require slightly more cortical control.

In [11]:

```
X = df_measures_all['Keypoint velocity (pixels/sec)']
Y = df_measures_all['Neural power (dB)']

X = sm.add_constant(X)
rlm_model = sm.RLM(Y, X, M=sm.robust.norms.HuberT())
rlm_results = rlm_model.fit()
rlm_results.summary()
```

about:srcdoc Page 7 of 9

Out[11]:

Robust linear Model Regression Results

Dep. Variable: Neural power (dB) No. Observations: 6447

Model: RLM **Df Residuals:** 6445

Method: IRLS Df Model: 1

Norm: HuberT

Scale Est.: mad

Cov Type: H1

Date: Thu, 03 Feb 2022

Time: 08:26:34

No. Iterations: 16

	coef	std err	Z	P> z	[0.025	0.975]
const	-0.0075	0.006	-1.251	0.211	-0.019	0.004
Keypoint velocity (pixels/sec)	1.635e-05	6.62e-06	2.470	0.014	3.37e-06	2.93e-05

If the model instance has been used for another fit with different fit parameters, then the fit options might not be the correct ones anymore .

Plot data with linear fit

Most of the data appears clustered near 0 velocity. The positive relationship between neural power and right wrist vertical velocity is only barely visible. Additional steps that may help better understand the relationship between neural spectral power and wrist velocity include: removing pose data with abnormally high standard deviation due to noisy tracking, subtracting spectral power in nearby periods with minimal movement from ECoG spectral power, and manually reviewing pose trajectories to remove noisy tracking periods.

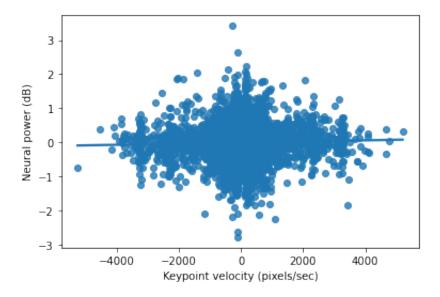
about:srcdoc Page 8 of 9

In [12]:

sns.regplot(data=df_measures_all, x='Keypoint velocity (pixels/sec)', y='Neural
power (dB)', robust=True, ci=None)

Out[12]:

<AxesSubplot:xlabel='Keypoint velocity (pixels/sec)', ylabel='Neural
power (dB)'>



In []:

about:srcdoc Page 9 of 9