```
1 !wget -c https://uni-bonn.sciebo.de/s/f81JgGZmEHXxMnB/download?path=%2F&files=steinmetz 2016-12-14 Cori.nc
   --2023-10-24 12:53:34-- https://uni-bonn.sciebo.de/s/f81JqGZmEHXxMnB/download?path=%2F
   Resolving uni-bonn.sciebo.de (uni-bonn.sciebo.de)... 128.176.1.2
   Connecting to uni-bonn.sciebo.de (uni-bonn.sciebo.de) | 128.176.1.2 | :443... connected.
   HTTP request sent, awaiting response... 200 OK
   Length: unspecified [application/zip]
   Saving to: 'download?path=%2F'
   download?path=%2F
                                  <=>
                                                1 1.16G 27.1MB/s
                                                                      in 43s
   2023-10-24 12:54:18 (27.6 MB/s) - 'download?path=%2F' saved [1241807443]
1 !unzip /content/download?path=%2F
   Archive: /content/download?path=%2F
      creating: steinmetz neuromatch dataset/
    extracting: steinmetz neuromatch dataset/steinmetz 2016-12-14 Cori.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2016-12-17 Cori.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2016-12-18 Cori.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-01-07 Muller.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-01-08 Muller.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-01-08 Radnitz.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-01-09 Muller.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-01-09 Radnitz.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-01-10 Radnitz.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-01-11 Radnitz.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-01-12 Radnitz.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-05-15 Moniz.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-05-16 Moniz.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-05-18 Moniz.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-06-15 Hench.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-06-16 Hench.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-06-17 Hench.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-06-18 Hench.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-10-11 Theiler.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-10-29 Richards.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-10-30 Richards.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-10-31 Richards.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-11-01 Forssmann.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-11-01 Richards.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-11-02 Forssmann.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-11-02 Richards.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-11-04 Forssmann.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-11-05 Forssmann.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-12-05 Lederberg.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-12-06 Lederberg.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-12-06 Tatum.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-12-07 Lederberg.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-12-07 Tatum.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-12-08 Lederberg.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-12-08 Tatum.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-12-09 Lederberg.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-12-09 Tatum.nc
    extracting: steinmetz_neuromatch_dataset/steinmetz_2017-12-10 Lederberg.nc
    extracting: steinmetz neuromatch dataset/steinmetz 2017-12-11 Lederberg.nc
```

```
1 dataset = xa.load dataset('/content/steinmetz neuromatch dataset/steinmetz 2016-12-14 Cori.nc')
```

## 1 dataset

```
xarray.Dataset
    ▶ Dimensions:
                         (trial: 364, time: 250, cell: 734, waveform_component: 3, sample: 82, probe: 384,
                         brain_area_lfp: 7, spike_id: 2446173)
    ▼ Coordinates:
       trial
                         (trial)
                                                 int32 1 2 3 4 5 6 ... 360 361 362 363 364
                                                                                                 time
                         (time)
                                               float64 0.01 0.02 0.03 ... 2.48 2.49 2.5
                                                                                                 cell
                         (cell)
                                                 int32 1 2 3 4 5 6 ... 730 731 732 733 734
                                                                                                 waveform co...
                         (waveform_component)
                                                 int32 123
                                                                                                 probe
                         (probe)
                                                 int32 1 2 3 4 5 6 ... 380 381 382 383 384
                                                                                                 brain_area_lfp
                         (brain_area_lfp)
                                                object 'ACA' 'LS' 'MOs' ... 'SUB' 'VISp'
                                                                                                 spike_id
                                                 int32 1 2 3 4 ... 2446171 2446172 2446173
                         (spike_id)
                                                                                                 ▶ Data variables: (31)
    ▶ Indexes: (7)
    ▼ Attributes:
       session date:
                         2016-12-14
                         Cori
       mouse:
       stim_onset:
                         0.5
       bin_size:
                         0.01
1 dataset.wheel.values
```

```
array([[-1, 0, 0, ..., 1, 0, 1],
      [0, -1, 0, \ldots, 1, 0, 0],
     [0, 0, -1, \ldots, -1, 0, 0],
      . . . ,
     [ 0, 0, 0, ..., 0, 0, 0],
     [ 0, 0, 0, ..., 0, 0, 0],
     [ 0, 0, 0, ..., 0, 0, 0]], dtype=int8)
```

## 1 dataset.wheel.to\_dataframe()

## wheel

trial	time	
1	0.01	-1

1 wheel\_data = dataset.wheel.to\_dataframe().reset\_index()

2 wheel data

	trial	time	wheel
0	1	0.01	-1
1	1	0.02	0
2	1	0.03	0
3	1	0.04	0
4	1	0.05	0
90995	364	2.46	0
90996	364	2.47	0
90997	364	2.48	0
90998	364	2.49	0
90999	364	2.50	0

91000 rows × 3 columns

<sup>1</sup> import seaborn as sns

<sup>1</sup> sns.lineplot(x=wheel\_data.time, y=wheel\_data.wheel)

<Axes: xlabel='time', ylabel='wheel'>
1.25 
Day 2

1 dataset

xarray.Dataset

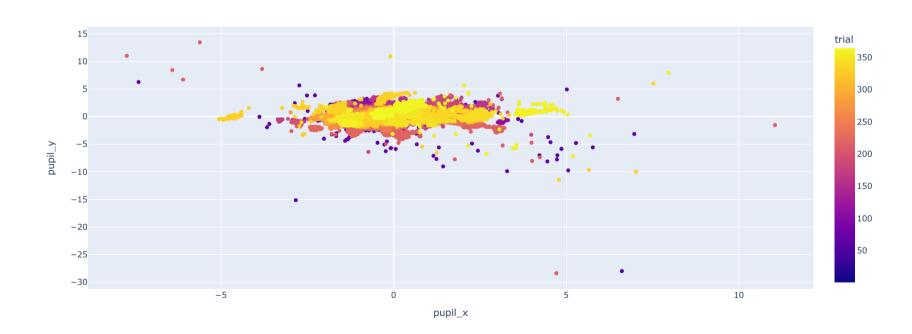
```
▶ Dimensions:
                    (trial: 364, time: 250, cell: 734, waveform_component: 3, sample: 82, probe: 384,
                    brain_area_lfp: 7, spike_id: 2446173)
    ▼ Coordinates:
      trial
                                                  int32 1 2 3 4 5 6 ... 360 361 362 363 ...
                    (trial)
      timo
                    /time)
                                                float64 0.01 0.00 0.00 0.40 0.40 0.5
1 x = dataset[['pupil_x','pupil_y','pupil_area']].to_dataframe()
2 r = dataset['reaction_time'].to_dataframe()
      probe
                    (probe)
                                                 int32 1 2 3 4 5 6 ... 380 381 382 383 ...
1 from matplotlib import pyplot as plt
                                                  -----
                   \-r-··-_
1 !pip install --upgrade plotly
                                                  ---
      ______
1 !pip install jupyter-dash
      stim onset
                    (trial)
                                                tloat64 0.5 0.5 0.5 0.5 ... 0.5 0.5 0.5 0.5
1 x
```

pupil\_x pupil\_y pupil\_area

trial	time					
1	0.01	0.812856	0.664247	0.165841		
	0.02		0.612752	0.158668		
	0.03	0.812352	0.632769	0.169667		
	0.04	0.760796	0.779100	0.172513		
	0.05	0.790401	0.680980	0.176774		
364	2.46	-0.446190	0.230169	0.117658		
	2.47	-0.472352	0.359197	0.120560		
	2.48	-0.453607	0.130451	0.113835		
	2.49	-0.486026	0.239184	0.109935		
	2.50	-0.491644	0.270002	0.119929		
91000 ro	ws × 3	columns				
пþ		(มเสเ	ıı_aıea_ııp, ıı	ıaı, шп <del>е</del> )	110at04 -2.001 -4.04 -4.180 10.04 0	
1 import p	lotly.	express a	s px			
=		er_3d(x,	x='pupil_x	x', y='pupil_y	', z='pupil_area')	



```
1 fig = px.scatter(x1, x='pupil_x', y='pupil_y',color='trial')
2 fig.show()
```

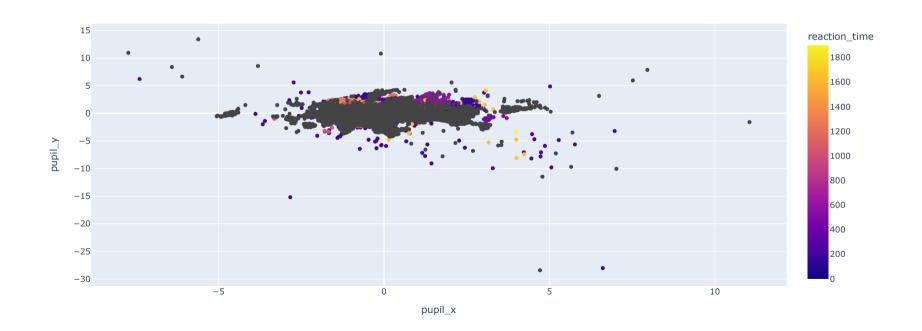


```
1 import pandas as pd
2 rt = pd.merge(x,r, on='trial')
3 rt
```

	pupil_x	pupil_y	pupil_area	reaction_time
trial				
1	0.812856	0.664247	0.165841	170.0
1	0.778162	0.612752	0.158668	170.0
1	0.812352	0.632769	0.169667	170.0
1	0.760796	0.779100	0.172513	170.0
1	0.790401	0.680980	0.176774	170.0
364	-0.446190	0.230169	0.117658	NaN
364	-0.472352	0.359197	0.120560	NaN
364	-0.453607	0.130451	0.113835	NaN
364	-0.486026	0.239184	0.109935	NaN
364	-0.491644	0.270002	0.119929	NaN

91000 rows × 4 columns

<sup>2</sup> fig.show()



<sup>1</sup> fig = px.scatter(rt, x='pupil\_x', y='pupil\_y',color='reaction\_time')



```
xarray.Dataset
```

1 spk = dataset[['spike\_trial','spike\_time','spike\_cell']].to\_dataframe()
2 spk

	spike_trial	spike_time	spike_cell	
spike_id				th
1	21	0.267635	1	
2	21	2.308335	1	
3	31	0.853476	1	
4	37	2 197501	1	



2.188634

2.399335

364

364

2446173 rows x 3 columns

2446172

2446173

1

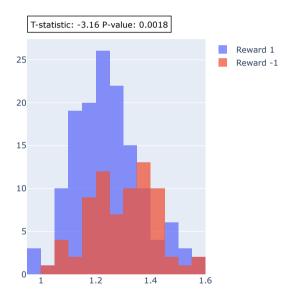
734

734

```
spike_trial spike_time spike_cell
                                                    th
     spike_id
      841385
                               0.050555
                                              364
                         1
      841386
                               0.108489
                                              364
      841387
                         1
                               0.172022
                                              364
      841388
                               0.376223
                                               364
 1 average_spike_time = all_cell.groupby(['spike_trial','spike_cell'])[['spike_time']].mean()
 2 average_by_cell = average_spike_time.groupby('spike_trial')['spike_time'].mean()
 3 average_by_cell
    spike_trial
           1.100715
           1.362811
           1.211432
    4
          1.339632
           1.139493
             . . .
    360
          1.218692
    361
          1.161034
    362
          1.214097
    363
          1.199479
    364
           1.118665
    Name: spike_time, Length: 364, dtype: float32
 1 ft = dataset['feedback_type'].to_dataframe()
 1 p_ft = ft[ft['feedback_type'] == 1].index.to_list()
 2 f1 = average_by_cell[p_ft]
 1
 1 p_ft = ft[ft['feedback_type'] == -1].index.to_list()
 1 f2 = average_by_cell[p_ft]
 1 from scipy.stats import ttest_ind
 2
 3 t_stat, p_val = ttest_ind(f1, f2)
 4
 1 import plotly.graph_objects as go
 2
 3 # Create histogram
 4 fig = go.Figure()
 6 # Add traces (data) for the histograms
 7 fig.add_trace(go.Histogram(x=f1, name='Reward 1', opacity=0.75,xbins=dict(size=0.05)))
 8 fig.add_trace(go.Histogram(x=f2, name='Reward -1', opacity=0.75,xbins=dict(size=0.05)))
10 # Overlay histograms
```

```
11 fig.update layout(barmode='overlay')
12
13 fig.add_annotation(
14
      x=0, y=1.1, # You might want to adjust these coordinates for better placement
       xref="paper", yref="paper", # Use "paper" coordinates for relative positioning
15
16
       text=f"T-statistic: {t_stat:.2f}\nP-value: {p_val:.4f}",
17
       showarrow=False,
18
       font=dict(size=12, color="black"),
19
       bordercolor="black",
20
       borderwidth=1,
21
       borderpad=4,
22
       bgcolor="white"
23)
24 # Display the figure
25 fig.show()
26
```





```
1 SUB = dataset['brain_area'].to_dataframe()
2 cell_label = SUB[SUB['brain_area']=='SUB'].index

1 rs_spk = dataset['spike_rate'].to_dataframe().reset_index()

1 sub_cell = rs_spk[rs_spk['cell'].isin(cell_label)]

1 average_spike_time = sub_cell.groupby(['trial','cell'])[['spike_rate']].mean()
2 average_spike_time
```

```
3 average_by_cell = average_spike_time.groupby('trial')['spike_rate'].mean()
4 average by cell
   trial
   1
          0.079307
   2
          0.067573
   3
          0.055947
          0.056853
   4
          0.063520
          . . .
   360
         0.090987
         0.073333
   361
   362
          0.083840
   363
         0.090827
   364
          0.085600
   Name: spike_rate, Length: 364, dtype: float64
1 rt_df = dataset['reaction_time'].to_dataframe()
1 st df = dataset[['spike time','spike cell','spike trial']].to dataframe()
1 st cell df = st df[st df['spike cell'].isin(cell label)]
2 average_spike_time = st_cell_df.groupby(['spike_trial','spike_cell'])[['spike_time']].mean()
3 average_spike_time = average_spike_time.groupby('spike_trial')['spike_time'].mean()
4 average spike time
   spike_trial
         1.134605
         1.268446
         1.088225
   3
   4
         1.205556
         1.177878
           . . .
   360
         1.251748
   361
         1.258168
   362
         1.285870
   363
         1.120888
   364
         1.388254
   Name: spike time, Length: 364, dtype: float32
1 import pandas as pd
2 import numpy as np
1 df = pd.DataFrame({
     'reaction time': rt df.values.flatten(),
      'spike_time': average_spike_time,
4
      'spike_rate': average_by_cell,
5 })
1 df = df.reset_index()
2 df
```

	level_0	index	reaction_time	spike_time	spike_rate
0	0	1	170.0	1.134605	0.079307
1	1	2	230.0	1.268446	0.067573
2	2	3	200.0	1.088225	0.055947
3	3	4	860.0	1.205556	0.056853
4	4	5	140.0	1.177878	0.063520
				•••	•••
359	359	360	NaN	1.251748	0.090987
360	360	361	NaN	1.258168	0.073333
361	361	362	NaN	1.285870	0.083840
362	362	363	NaN	1.120888	0.090827

1 ft = dataset['feedback\_type'].to\_dataframe()
364 rows x 5 columns

1 p\_ft = ft[ft['feedback\_type'] == 1].index.to\_list()
2
3 f1 = df.iloc[p\_ft,:]
4 f1

	level_0	index	reaction_time	spike_time	spike_rate	$\blacksquare$
1	1	2	230.0	1.268446	0.067573	ıl
2	2	3	200.0	1.088225	0.055947	
3	3	4	860.0	1.205556	0.056853	
4	4	5	140.0	1.177878	0.063520	
9	9	10	200.0	1.082780	0.070240	
191	191	192	inf	1.193350	0.073280	
194	194	195	inf	1.251339	0.060587	
198	198	199	inf	1.114188	0.064107	
200	200	201	1370.0	1.427714	0.066933	
201	201	202	inf	1.275791	0.066133	

141 rows x 5 columns

<sup>1</sup> p\_ft = ft[ft['feedback\_type'] == -1].index.to\_list()
2
3 f2 = df.iloc[p\_ft,:]
4 f2

	level_0	index	reaction_time	spike_time	spike_rate		
5	5	6	1340.0	1.265247	0.058560	th.	
6	6	7	740.0	1.170172	0.062667		
7	7	8	990.0	1.086859	0.053440		
8	8	9	1180.0	1.454938	0.045440		
18	18	19	920.0	1.245060	0.058613		
210	210	211	1050.0	1.317677	0.070080		
211	211	212	inf	1.281139	0.067413		
212	212	213	inf	1.327681	0.072853		
213	213	214	1350.0	1.294707	0.073440		
<pre>1 # Create line plot 2 fig = go.Figure(data=go.Scatter(x=f1.index, y=f1.reaction_time, mode='lines+markers')) 3 4 # Add title and labels 5 fig.update_layout( 6</pre>							

yaxis\_title='reaction\_time')

9 # Display the figure

10 fig.show()

