# PCA analysis

April 16, 2020

Navigate to directory containing .mat data.

```
[1]: cd ../_data/matlabData/
```

/home/gustav/Documents/DD142X/code/\_data/matlabData

Function definitions for data extraction and feature extraction. Basic imports.

```
[2]: import numpy as np
     import matplotlib.pyplot as plt
     from h5py import File
     def getMatlabValues(fileName):
         with File(fileName, "r") as data:
             return {
                 key: np.array(data[key]["values"]).flatten() for key in data.keys()
             }
     # Fourier Feature Vector
     def ffv(xs, Fs = 16000., epoch_size = 2 ** 11, fft_n = 2 ** 14):
         # Pad with zeroes for more frequency outputs
         # Compare np.fft.fftfreq(n, 1/16000) for n = 2**11, 2**14
         fft_in = np.zeros((xs.shape[0], fft_n))
         fft_in[ : , 0:epoch_size] = xs
         frqs = np.fft.fftfreq(fft_n, 1./Fs)
         lo = np.where(frqs > 12)[0][0]
         hi = np.where(frqs > 30)[0][0]
         fftxs = np.abs(np.fft.fft(fft_in)[:,lo:hi])
         return fftxs, frqs[lo:hi]
```

I/O

```
[3]: mlDict = getMatlabValues("NPR-075.b11.mat")
vals_str = np.array([v for k, v in mlDict.items() if "str_lfp" in k])
```

```
vals_gp = np.array([v for k, v in mlDict.items() if "gp_lfp" in k])
     vals_all = np.concatenate((vals_str, vals_gp), axis = 0)
     print(vals_str.shape)
     print(vals_gp.shape)
     print(vals_all.shape)
    (11, 1587696)
    (15, 1587696)
    (26, 1587696)
    First, consider PCA of "raw" data. Trim excess datapoints in order to work with 2 n length
    epochs.
[4]: print( vals_str.shape[1] / 2 ** 14 )
     print( 2 ** 14 / 16000 )
    96.9052734375
    1.024
    2 ** 14 seems like a reasonable maximum epoch size.
[5]: vals_str = vals_str[ : , 0 : 2 ** 14 * 96]
     vals_gp = vals_gp [:, 0:2**14*96]
     vals_all = vals_all[ : , 0 : 2 ** 14 * 96]
     print(vals_str.shape)
     print(vals_gp.shape)
     print(vals_all.shape)
    (11, 1572864)
    (15, 1572864)
    (26, 1572864)
    Consider amount of required principal components to explain variance to some degree.
[6]: from sklearn.decomposition import PCA
[7]: print("### PCA explained variance ratio per component and sum, n_components = ___
     →10")
     for epoch_size in [
         2 ** 14, 2 ** 13, # ~1s, 0.5s
         2 ** 12, 2 ** 11, # ~0.25s, 0.125s
         2 ** 10, 2 ** 9  # Very short, may cause memory issues - hardly optimized
     ]:
         # Investigate for large n_components
```

print("### Epoch size " + str(epoch\_size))

```
pca_str = PCA(n_components = 10).fit(vals_str.copy().reshape((-1, __
 →epoch_size)))
    pca_gp = PCA(n_components = 10).fit(vals_gp.copy().reshape((-1,__
 →epoch size)))
    pca_all = PCA(n_components = 10).fit(vals_all.copy().reshape((-1,__
 →epoch_size)))
    print("\tStriatum")
    print(pca_str.explained_variance_ratio_)
    print(pca_str.explained_variance_ratio_.sum())
    print("\tGlobus pallidus")
    print(pca_gp.explained_variance_ratio_)
    print(pca_gp.explained_variance_ratio_.sum())
    print("\tAll")
    print(pca_all.explained_variance_ratio_)
    print(pca_all.explained_variance_ratio_.sum())
### PCA explained variance ratio per component and sum, n_components = 10
### Epoch size 16384
        Striatum
[0.13201564 0.09549128 0.08267939 0.0709507 0.06386331 0.05941619
0.6310981415229998
       Globus pallidus
[0.13970142\ 0.09407339\ 0.08341252\ 0.05959512\ 0.05431434\ 0.04960502
0.04085061 0.03476214 0.02840412 0.02579094]
0.6105096281327804
       All
[0.13595805 0.09411479 0.08162108 0.06524052 0.05891072 0.05358219
          0.03069342 0.02821657 0.02719611]
0.6176594579235939
### Epoch size 8192
        Striatum
[0.15187149 0.12482922 0.1079342 0.09913861 0.08795753 0.05288573
0.03481608 0.0186003 0.01585444 0.01430079]
0.7081883828881758
       Globus pallidus
[0.15902287 \ 0.11923743 \ 0.10791143 \ 0.08742827 \ 0.07933465 \ 0.0506975
0.03192745 0.02011312 0.01737053 0.01659388]
0.6896371362816823
       A11
          0.11412229 0.1134704 0.09218164 0.0860637 0.05165942
Γ0.154862
0.03317962 0.01921961 0.0163357 0.01554296]
0.6966373323654936
### Epoch size 4096
```

Striatum

- [0.24321588 0.1905 0.16359195 0.06412213 0.02667111 0.0245737
- 0.02319225 0.02064487 0.01704692 0.01646161]
- 0.7900204237254138

Globus pallidus

- $\begin{bmatrix} 0.26572307 & 0.17554352 & 0.13646579 & 0.05858759 & 0.0294942 & 0.02782842 \end{bmatrix}$
- 0.02592714 0.02423092 0.02231361 0.02044605]
- 0.7865603148082775

All

- [0.25507889 0.18175209 0.14923832 0.06116898 0.02756925 0.02579797
- 0.02503227 0.02305841 0.01966268 0.01857207]
- 0.7869309174324854

#### ### Epoch size 2048

Striatum

- $[0.40333083\ 0.23222896\ 0.06722281\ 0.03842049\ 0.0361776\ 0.03197513$
- 0.02217892 0.01402925 0.01211575 0.01169958]
- 0.8693793177856861

Globus pallidus

- $[0.41422713 \ 0.20006564 \ 0.0651903 \ \ 0.04534713 \ 0.04263925 \ 0.03980649$
- 0.02984995 0.0143692 0.00859884 0.00816665]
- 0.8682605732519141

A11

- $\hbox{\tt [0.40943561 \ 0.21405838 \ 0.06612751 \ 0.04061722 \ 0.04050694 \ 0.03719184 }$
- 0.02663702 0.0142366 0.00992589 0.00965878]
- 0.868395782439071

### ### Epoch size 1024

Striatum

- $\hbox{\tt [0.59032811\ 0.14154979\ 0.06555062\ 0.04238349\ 0.01959164\ 0.01372395] }$
- 0.0126338 0.01261415 0.01141206 0.00879597]
- 0.9185835912056062

Globus pallidus

- $[0.57932412\ 0.1326198\ 0.08038562\ 0.05201119\ 0.01938242\ 0.01218263$
- 0.00965168 0.00853804 0.00824053 0.00717727]
- 0.9095133056389795

All

- $[0.58428032 \ 0.13632042 \ 0.0738323 \ \ 0.04791357 \ 0.0194909 \ \ 0.01283283$
- 0.01079399 0.01031238 0.00977479 0.00792491]
- 0.9134764181012286

## ### Epoch size 512

Striatum

- 0.0115724 0.00671003 0.0045563 0.00453135]
- 0.944875222722342

Globus pallidus

[0.69615742 0.12981686 0.04567163 0.0176672 0.01094913 0.00946829

```
0.00841649 0.00533999 0.00359789 0.00348624]
0.9305711499013999
All
[0.70249968 0.12381821 0.0441189 0.0191505 0.01190735 0.0114102 0.0100411 0.00594364 0.00401489 0.00392155]
0.9368260280254591
```

At n >= 2048 points per epoch, lots of variance can be explained with very few components. Even at lower greater n, much variance can be explained with somewhat few components. Could be an interesting alternative to fourier transform. High dimensionality (large number of principal components) not a big problem - very large dataset.

The weight of each PC in a resulting PC-based feature vector could be scaled by explained variance ratio, or some other technique along those lines.

**PCA**, however, ignores spectral domain. This could be problematic. PCA gives information only on "entire" LFP, not beta-range, which we are targeting(?). PCA can be applied to Fourier-transform-based feature vector "ffv".

```
[]: # Number of features extracted remains (in this example) constant at 18
    print("### PCA explained variance ratio per component and sum, for DFT-based ∪
     for epoch size in [
        2 ** 14, 2 ** 13,
        2 ** 12, 2 ** 11,
        2 ** 10, #2 ** 9 # Greater risk of memory issues, workaround not
     \rightarrow implemented
    ]:
        print("### Epoch size " + str(epoch_size))
        ffv_str, _ = ffv(vals_str.reshape((-1, epoch_size)), epoch_size=epoch_size)
        ffv_gp, _ = ffv(vals_gp.reshape((-1, epoch_size)), epoch_size=epoch_size)
        ffv_all, _ = ffv(vals_all.reshape((-1, epoch_size)), epoch_size=epoch_size)
        pca_str = PCA(n_components = 10).fit(ffv_str)
        pca_gp = PCA(n_components = 10).fit(ffv_gp)
        pca_all = PCA(n_components = 10).fit(ffv_all)
        print("\tStriatum")
        print(pca_str.explained_variance_ratio_)
        print(pca_str.explained_variance_ratio_.sum())
        print("\tGlobus pallidus")
        print(pca_gp.explained_variance_ratio_)
        print(pca_gp.explained_variance_ratio_.sum())
        print("\tAll")
        print(pca_all.explained_variance_ratio_)
```

```
print()
### PCA explained variance ratio per component and sum, for DFT-based feature
vector, n_components = 10
### Epoch size 16384
        Striatum
[0.1282068 \quad 0.10920921 \quad 0.09645611 \quad 0.08493102 \quad 0.07250854 \quad 0.06865115
0.06047625 0.05241997 0.04776931 0.0458104 ]
0.7664387619805519
        Globus pallidus
[0.12801003 0.11145797 0.1017971 0.08601633 0.08069299 0.06828279
0.06331946 0.05063642 0.04935059 0.0448874 ]
0.7844510793717202
        All
[0.12412244\ 0.10926272\ 0.10029431\ 0.08297152\ 0.07811252\ 0.06657244
0.0614023  0.05190005  0.05048729  0.04812489]
0.7732504782016221
### Epoch size 8192
        Striatum
[0.1748784 0.12133393 0.10323539 0.08420862 0.08053167 0.06975029
0.05514768 0.05259114 0.04747176 0.04044812]
0.8295969892428826
        Globus pallidus
[0.60077462\ 0.0662128\ 0.05387123\ 0.03661241\ 0.03646467\ 0.03411644
0.03013468 0.02531864 0.02213112 0.01950109]
0.9251377003498068
        All
[0.1725584 0.12378346 0.11180106 0.08366169 0.07423475 0.06750929
0.06530015 0.05014109 0.04681424 0.04113517]
0.8369392858187161
### Epoch size 4096
        Striatum
[0.273996
            0.17234378 0.13692129 0.11220554 0.09711271 0.06560395
 0.05480532 0.03874422 0.02356104 0.01272468]
0.9880185432577007
        Globus pallidus
[0.26164328 0.18845243 0.14646079 0.11317954 0.09679361 0.0653134
0.04899552 0.03669224 0.02074158 0.01018742]
0.9884598037893412
        All
[0.26624254 0.18184926 0.14380045 0.1129786 0.09698456 0.06495389
0.05114059 0.03749518 0.02175697 0.01113666]
```

print(pca\_all.explained\_variance\_ratio\_.sum())

#### 0.988338712738876

Good potential for feature reduction for sufficiently short epoch sizes. 10 components do a very good job at n=4096. 6 components good enough for  $\sim 85\%$  variance. 6 components do a very good job at n=2048. 4 components good enough for >90% variance. At n=1024, 2 components  $\sim 85\%$  variance. 3 components >90%. Memory issues at n=512. Workaround pending.

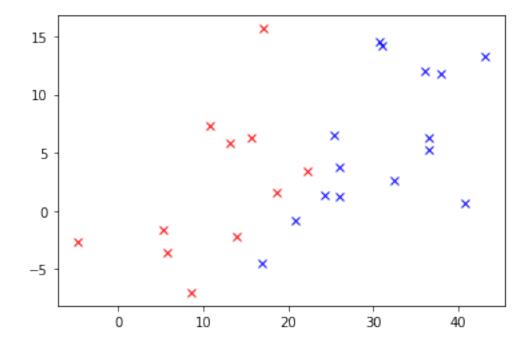
If used as feature vectors, should take explained variance ratio into account (as previously discussed).

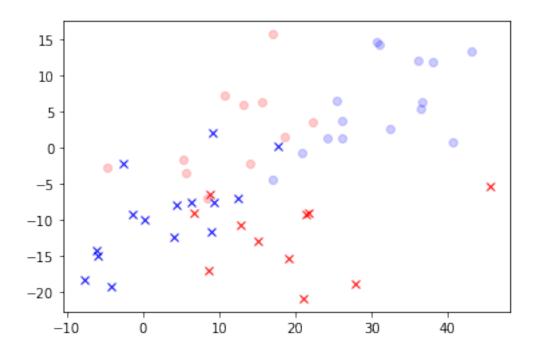
```
[9]: EPOCH_MAX = 20

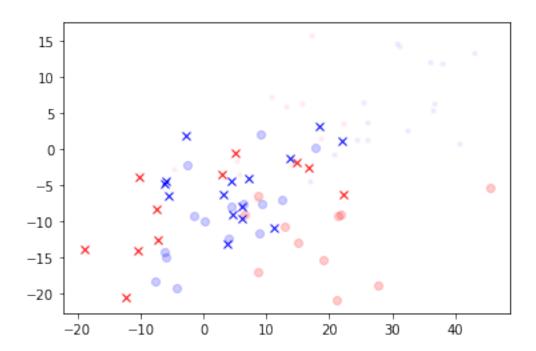
plt.clf()
for hi in range(0, EPOCH_MAX):
    for ep in range(0, hi):
```

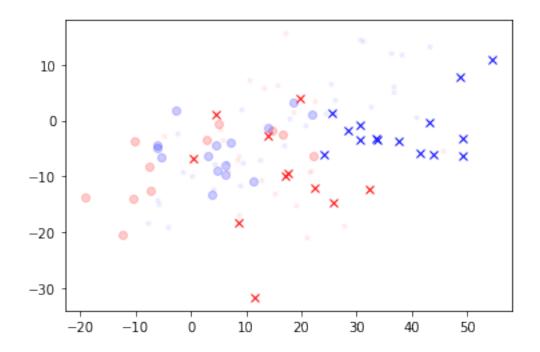
```
alpha = 0.05
    shape = '.'
    if ep == hi - 1:
        alpha = 1
        shape = 'x'
    elif ep == hi - 2:
        shape = 'o'
        alpha = 0.2
    plt.plot(
        [x for x,_ in str_tmp[ : , ep]],
        [y for _,y in str_tmp[ : , ep]],
        'r' + shape,
        alpha=alpha
    )
    plt.plot(
        [x for x,_ in gp_tmp[ : , ep]],
        [y for _,y in gp_tmp[ : , ep]],
        'b' + shape,
        alpha=alpha
    )
plt.show()
```

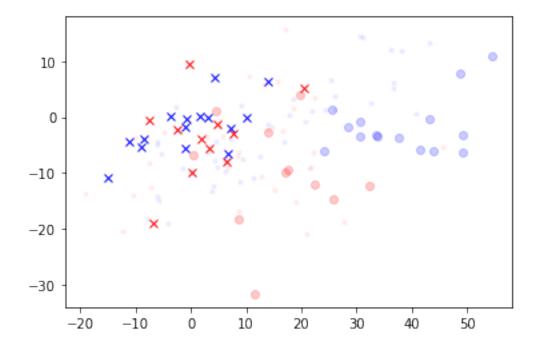
<Figure size 432x288 with 0 Axes>

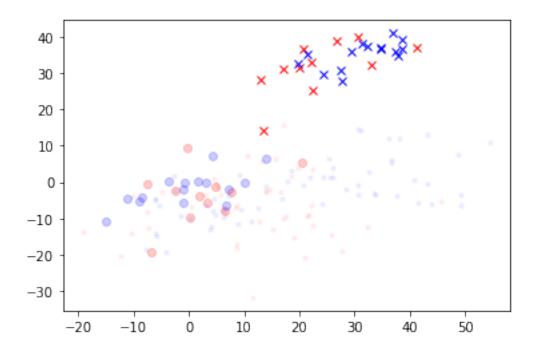


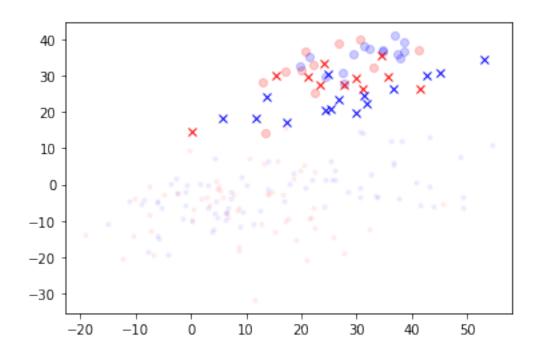


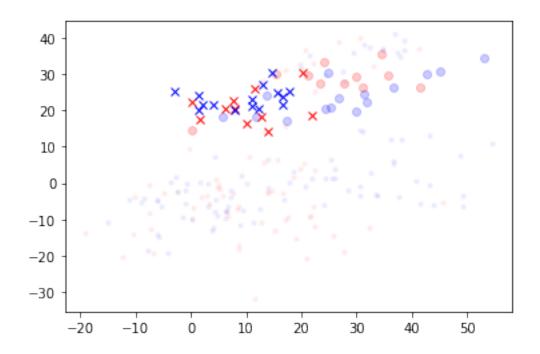


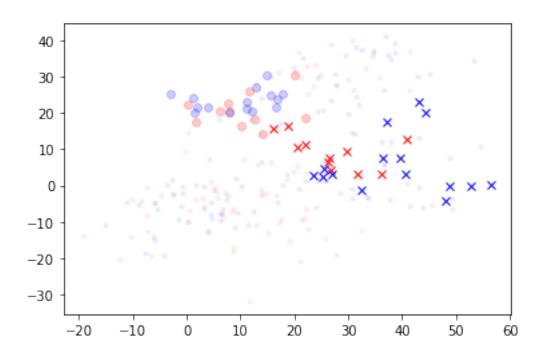


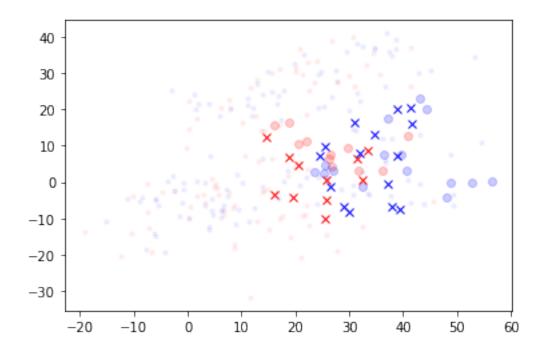


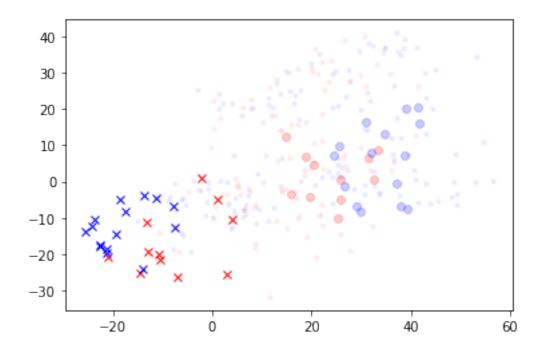


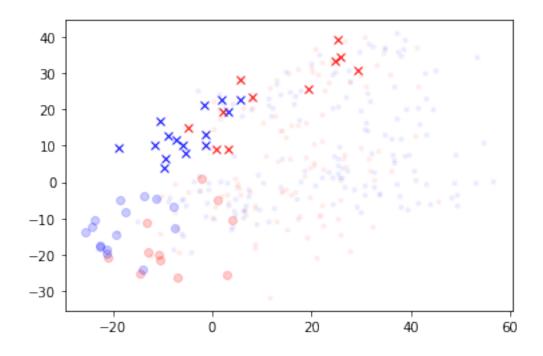


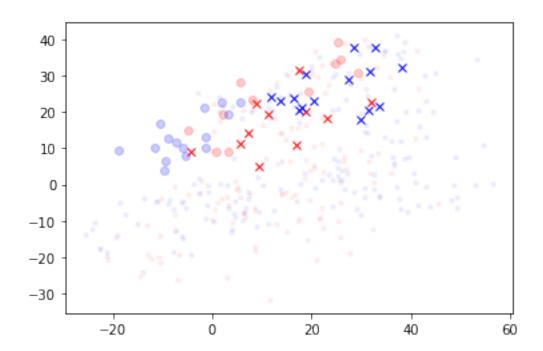


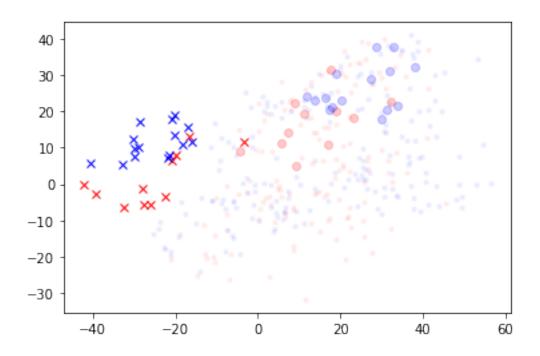


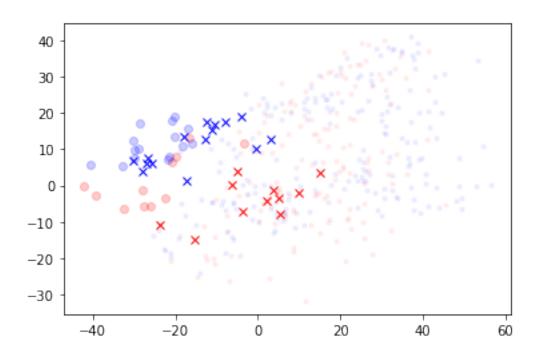


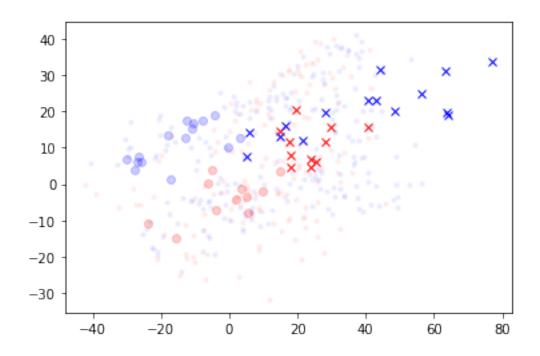


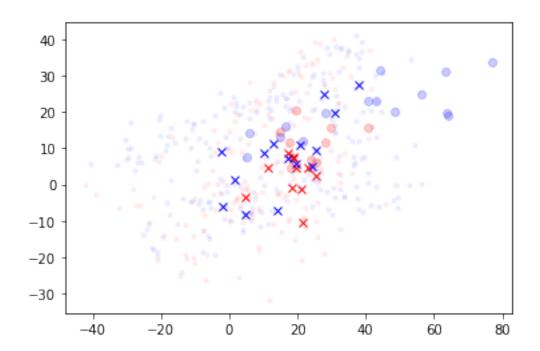


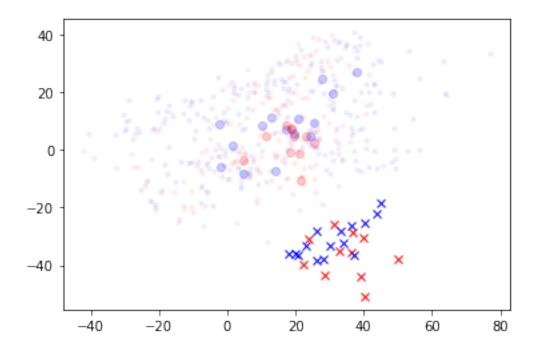


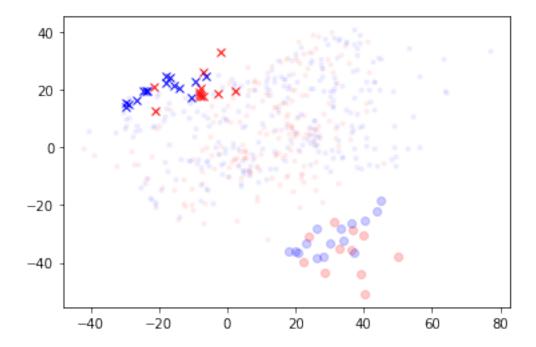












Data seems to move "together in a cloud", though consistently slightly separate perchannel-type. Similar results as use of TSNE.

## Attempt to rasterize PCA

```
[10]: # Hard to understand but efficient function.
      # Written in a "pythonic" (simpler) rather than "numpythonic" way, takes many_
      → many MANY times longer.
      def rasterize(samples, side = 40):
          # Find range of values; input is N x 2 numpy array
          maxx1 = np.max(samples[:, 0])
          minx1 = np.min(samples[:, 0])
          maxx2 = np.max(samples[:, 1])
          minx2 = np.min(samples[:, 1])
          # Decides the "pixels"
          x1incr = (maxx1 - minx1) / side
          x2incr = (maxx2 - minx2) / side
          raster = np.zeros((side, side))
          # Count amount of samples belonging to each pixel
          for i in range(0, side):
              for j in range(0, side):
                       Uses numpy boolean indexing
                  raster[i,j] += samples[
                      (samples[ : , 0] \geq minx1 + i * x1incr) & \
                      (samples[:, 0] < minx1 + (i + 1) * x1incr) & \setminus
                      (samples[:, 1] >= minx2 + j * x2incr) & \
                      [samples[:, 1] < minx2 + (j + 1) * x2incr)
                  ].flatten().shape[0]
          return raster
```

```
[11]: print(pca_plot.explained_variance_ratio_)
    raster_all = rasterize(embeddings.reshape((-1, 2)))
    plt.imshow(raster_all, cmap = 'gray', vmin = 0, vmax = np.max(raster_all))
    plt.colorbar()
    plt.show()

    raster_gp = rasterize(gp_tmp.reshape((-1, 2)))
    plt.imshow(raster_gp, cmap = 'gray', vmin = 0, vmax = np.max(raster_gp))
    plt.colorbar()
    plt.show()

    raster_str = rasterize(str_tmp.reshape((-1, 2)))
    plt.imshow(raster_str, cmap = 'gray', vmin = 0, vmax = np.max(raster_str))
    plt.colorbar()
    plt.show()
```

## [0.58872963 0.30003181]

