

new read result

March 18, 2021

```
[1]: import os;
import sys;
import random;
import math
import json
import imutils;

import numpy as np;
import cv2;
import matplotlib.pyplot as plt;

from skimage.feature import greycomatrix, greycoprops, local_binary_pattern

import tensorflow as tf
from tensorflow import keras;

from tensorflow.keras.callbacks import ModelCheckpoint

from sklearn.metrics import roc_curve, auc

from keras import backend as K

import mahotas as mt
import sklearn

from skimage.filters import gabor

from sklearn.manifold import TSNE

from mpl_toolkits.mplot3d import Axes3D

import umap
```

```
import phate

from pywt import dwt2

import csv
```

/usr/lib/python3/dist-packages/h5py/___init___py:36: FutureWarning: Conversion of the second argument of issubdtype from `float` to `np.floating` is deprecated. In future, it will be treated as `np.float64 == np.dtype(float).type`.

from ._conv import register_converters as _register_converters
Using TensorFlow backend.

/home/ubuntu/dlennv3/lib/python3.6/site-packages/numba/errors.py:137:
UserWarning: Insufficiently recent colorama version found. Numba requires
colorama >= 0.3.9
warnings.warn(msg)

[1]:

[2]: mapfile="/home/ubuntu/project/pythologyImageDL/scatterplot/data/
→new_gland_tsne32.csv"

[3]: data=[]
with open(mapfile) as f:
f2 =csv.reader(f)
data=list(f2)

data=np.asarray(data)

[5]: colors=[];
for i in data:
if i[1] =="normal":
colors.append("green")
elif i[1] =="treatment":
colors.append("blue")
elif i[1] =="dss":
colors.append("red")

else:
print(i)

[6]: x=[float(i) for i in data[:, 2]];
y=[float(i) for i in data[:,3]];

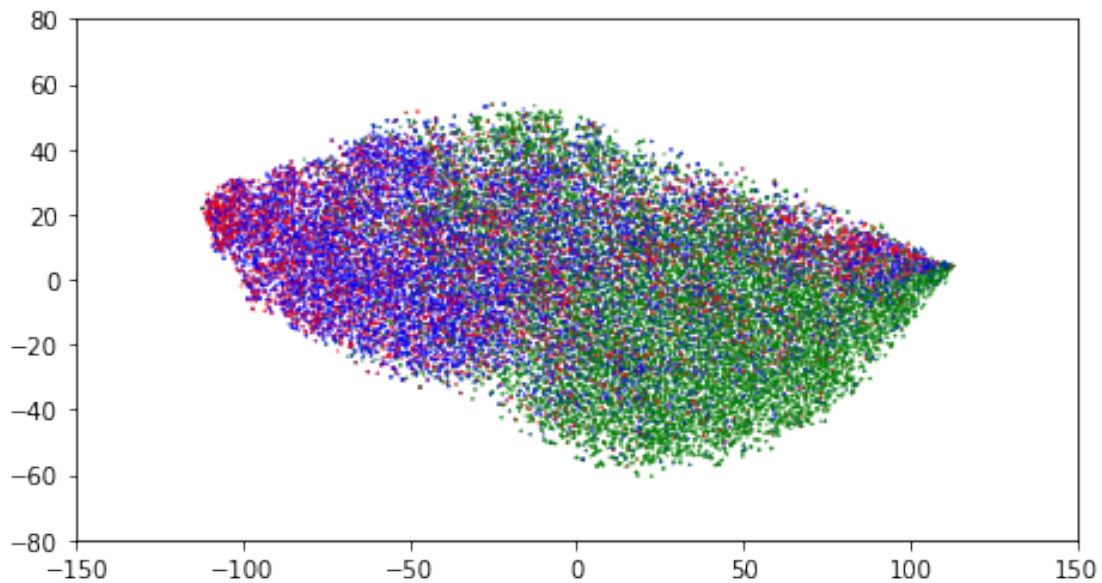
[274]: fig= plt.figure(figsize=(7.5,4))
plt.xlim([-150,150])
plt.ylim([-80,80])

ax = fig.add_subplot(111)
ax.scatter(x, y,c=colors,s=0.1)

```
/home/ubuntu/dlennv3/lib/python3.6/site-packages/ipykernel_launcher.py:5:
MatplotlibDeprecationWarning: Adding an axes using the same arguments as a
previous axes currently reuses the earlier instance.  In a future version, a new
instance will always be created and returned.  Meanwhile, this warning can be
suppressed, and the future behavior ensured, by passing a unique label to each
axes instance.
```

```
"""
```

[274]: <matplotlib.collections.PathCollection at 0x7ff39b3099e8>



```
[10]: import seaborn as sns
```

```
[74]: dssx = []
dssy = []
dssid=[];

normalx = []
normaly = []
normalid=[];

treatx = []
treaty = []
treatid=[];

for i in data:
```

```

if i[1] == "normal":
    normalx.append(float(i[2]))
    normaly.append(float(i[3]))
    normalid.append(i[0])

elif i[1] == "treatment":
    treatx.append(float(i[2]))
    treaty.append(float(i[3]))
    treatid.append(i[0])
elif i[1] == "dss":
    dssx.append(float(i[2]))
    dssy.append(float(i[3]))
    dssid.append(i[0])

else:
    print(i)

```

```

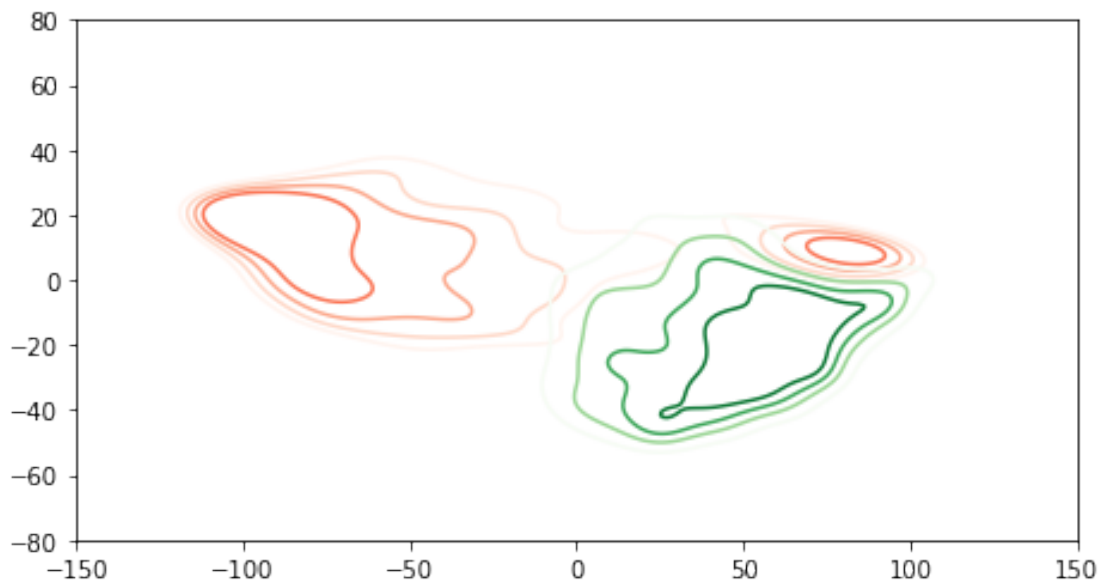
[508]: fig = plt.figure(figsize=(7.5, 4))

sns.kdeplot(x=dssx, y=dssy, levels=5, cmap="Reds", thresh=0.4)
sns.kdeplot(x=normalx, y=normaly, levels=5, cmap="Greens", thresh=0.4)

plt.xlim([-150,150])
plt.ylim([-80,80])

plt.show()

```



```
[510]: # I score each level of contour. disease is -4,-3,-2,-1,0 control is
→0,1,2,3,4
#we observe all the node location from treatment. and get the score of
→node(each gland).
```

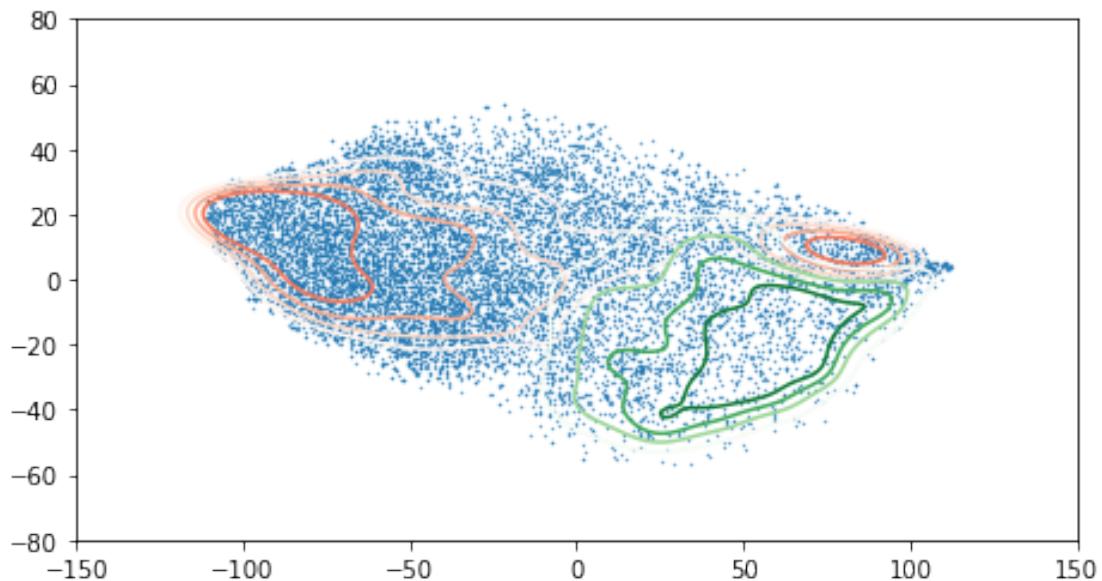
```
[509]: fig = plt.figure(figsize=(7.5, 4))

sns.kdeplot(x=dssx, y=dssy, levels=5, cmap="Reds", thresh=0.4)
sns.kdeplot(x=normalx, y=normaly, levels=5, cmap="Greens", thresh=0.4)

plt.xlim([-150,150])
plt.ylim([-80,80])

plt.scatter(treatx, treaty,s=0.1)

plt.show()
```



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[ ]:
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```
[511]: fig= plt.figure(figsize=(7.5/1.8,4/1.8))
sns.kdeplot(x=dssx, y=dssy, levels=5 ,thresh=0.4,fill=True,colors=["#COCOC0",
→"#808080", "#404040", "#000000"])

plt.xlim([-150,150])
plt.ylim([-80,80])

plt.axis('off')
```

```
plt.savefig("dss_contour.png",pad_inches=0)
```



```
[513]: fig= plt.figure(figsize=(7.5/1.8,4/1.8))

sns.kdeplot(x=normalx, y=normaly, levels=5 ,thresh=0.
→4,fill=True,colors=["#C0C0C0", "#808080", "#404040", "#000000"])

plt.xlim([-150,150])
plt.ylim([-80,80])

plt.axis('off')

plt.savefig("normal_contour.png",pad_inches=0)
```



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[ ]:
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```
[514]: nc = cv2.imread("normal_contour.png",cv2.IMREAD_GRAYSCALE)
```

```
[515]: nc.shape
```

[515]: (160, 300)

```
[516]: dc = cv2.imread("dss_contour.png",cv2.IMREAD_GRAYSCALE)
```

```
[517]: dc.shape
```

[517]: (160, 300)

```
[ ]:
```

```
[518]: dss_score=[];
control_score=[];
eval_score=[];
for i in range(len(treatid)):
    dsscolor=dc[round(treaty[i]) +80 ][round(treatx[i]) +150 ]
    dss_score.append(4-round(dsscolor/64))

    controlcolor=nc[round(treaty[i]) +80 ][round(treatx[i]) +150 ]
    control_score.append(4-round(controlcolor/64))
    eval_score.append(control_score[i]-dss_score[i])
```

```
[519]: score2=[x*-1 for x in eval_score]
```

```
[520]: score3=[ ]
for x in score2:
    if x <=0:
        score3.append(0)
    else:
        score3.append(x)
```

```
[521]: result=dict()
for i in treatid:
    result[i.split("___")[0]]={"dss_score":[],"combine_score":[]};

for i in range(len(treatid)):
    result[treatid[i].split("___")[0]]["dss_score"].append(dss_score[i]);
    result[treatid[i].split("___")[0]]["combine_score"].append(score3[i]);
```

```
[522]: for i in result:
    result[i]['dss_score']=np.mean(result[i]['dss_score'])
    result[i]['combine_score']=np.mean(result[i]['combine_score'])
```

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[ ]:
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```
[524]: #score method 2
```

```
[540]: slidedata=dict();

ids32 = data[:,0]
for i in range(len(ids32)):
    name = ids32[i].split("___")[0];
    slidedata[name]={"color":colors[i],"umap":[],"phate":[],"tsne":[]};
```

```

for i in range(len(ids32)):
    name = ids32[i].split("___")[0];

    coor = [float(data[i][2]),float(data[i][ 3])];
    slidedata[name]["tsne"].append(coor);

for i in slidedata:

    coors = slidedata[i]["tsne"];
    xs=[];
    ys=[];
    for c in coors:
        xs.append(c[0])
        ys.append(c[1])

    mean = np.mean(xs);
    std = np.std(xs)
    xs = [x for x in xs if (x > mean - 2 * std)]
    xs = [x for x in xs if (x < mean + 2 * std)]

    mean = np.mean(ys);
    std = np.std(ys)
    ys = [x for x in ys if (x > mean - 2 * std)]
    ys = [x for x in ys if (x < mean + 2 * std)]

    xs = sum(xs)/len(xs)
    ys = sum(ys)/len(ys)

    slidedata[i]["tsne"]=(xs,ys);

```

```

[542]: slidecolors=[];

slidetsne=[];
slideid=[]
for i in slidedata:
    slidecolors.append(slidedata[i]["color"]);

    slidetsne.append(list(slidedata[i]["tsne"]));
    slideid.append(i)

slidetsne=np.asarray(slidetsne)

```

```

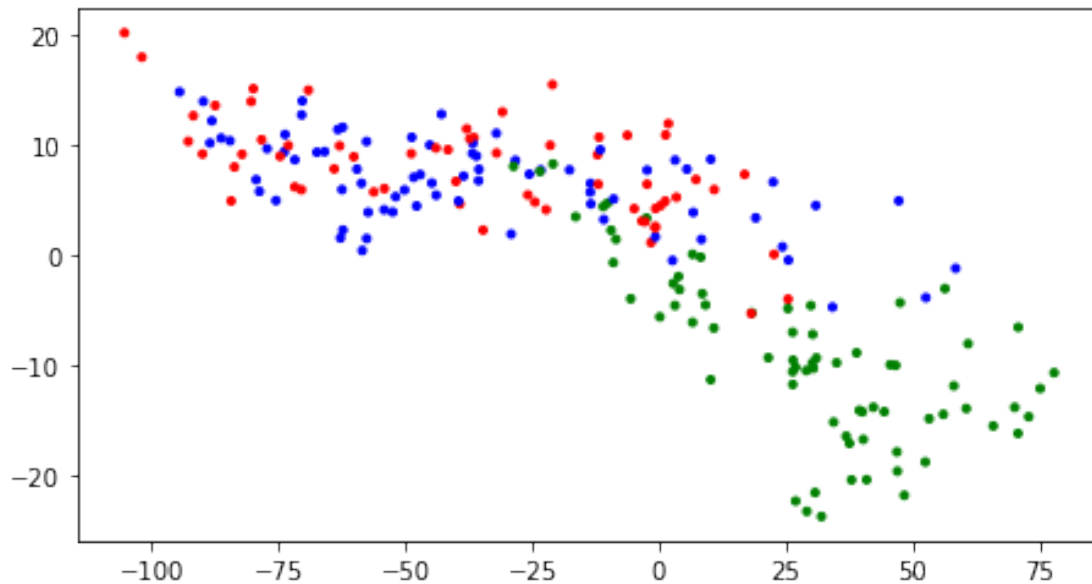
[543]: fig = plt.figure(figsize=(7.5, 4))

ax = fig.add_subplot(111)

ax.scatter(slidetsne[:, 0], slidetsne[:, 1],c=slidecolors,s=9)

```


[543]: <matplotlib.collections.PathCollection at 0x7ff34424f908>



```
[614]: scoredata=[]
with open("dss32_slidelevel_v4.csv") as f:
    f2 =csv.reader(f)
    scoredata=list(f2)

scoredata=np.asarray(scoredata)
```

```
[639]: keyvalue=dict();
for i in scoredata[1:]:
    keyvalue[i[0]]=int( float(i[-3]))

scorecolor=["green","steelblue","blue","orange","red"]
```

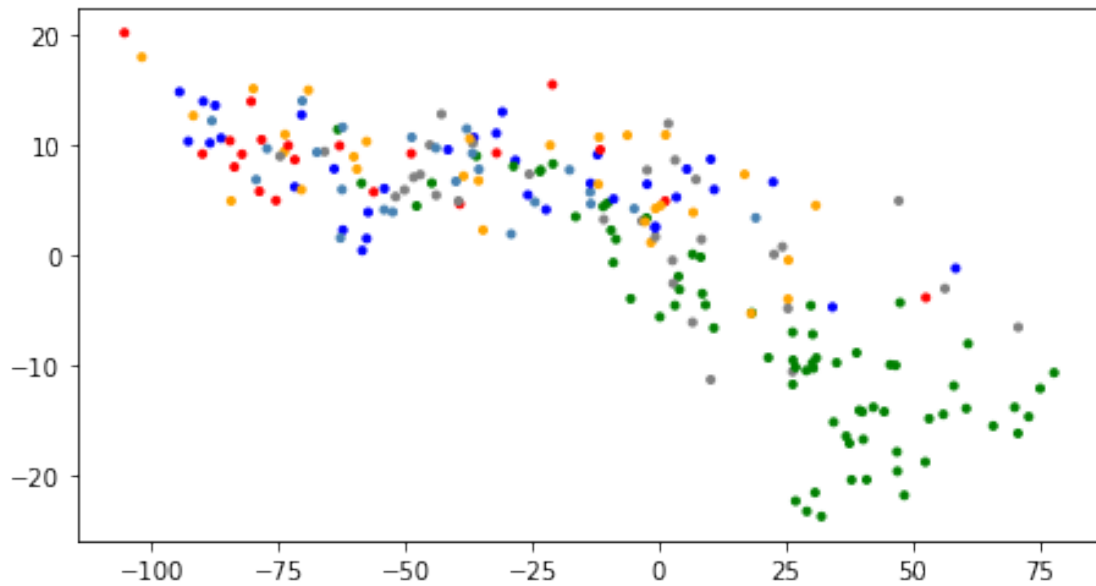
```
[640]: scorecolors=[];
for i in slideid:
    if i in keyvalue:
        val=keyvalue[ i]
        scorecolors.append(scorecolor[val])
    else:
        scorecolors.append("gray")
```

```
[641]: fig = plt.figure(figsize=(7.5, 4))

ax = fig.add_subplot(111)

ax.scatter(slidetsne[:, 0], slidetsne[:, 1],c=scorecolors,s=9)
```

[641]: <matplotlib.collections.PathCollection at 0x7ff3442e4c18>



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[ ]:
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```
[544]: dssdict={"treatment":set(),"dss":set(),"normal":set()};
for i in range(len(slideid)):
    if slidecolors[i] == "blue":
        dssdict["treatment"].add(slideid[i].split("_")[0])

    if slidecolors[i] == "red":
        dssdict["dss"].add(slideid[i].split("_")[0])

    if slidecolors[i] == "green":
        dssdict["normal"].add(slideid[i].split("_")[0])
```

```
[546]: for i in dssdict:
        dssdict[i]=list(dssdict[i])
        dssdict[i].sort()
```

```
[547]: pltindex=1;

for i in dssdict["treatment"]:
    sizelist=[];
    newcolors=[];

    for index in range(len(slideid)):

        if slideid[index].startswith(i+"_"):
            sizelist.append(50)
```

```

        location = slideid[index].split("_")[1];
        location=location[0];
        if location == "a":
            newcolors.append("indigo")
        elif location == "b":
            newcolors.append("darkorange")
        else:
            sizelist.append(1)
            newcolors.append(slidecolors[index])
    if pltindex==1:
        fig = plt.figure(figsize=(16, 5))
        ax = fig.add_subplot(120+pltindex)

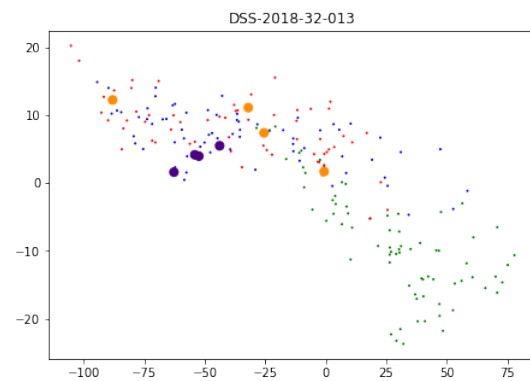
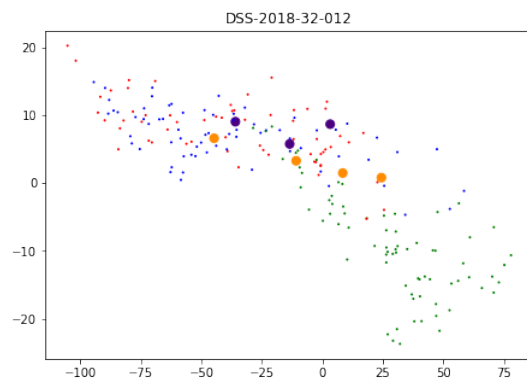
        ax.set_title(i)

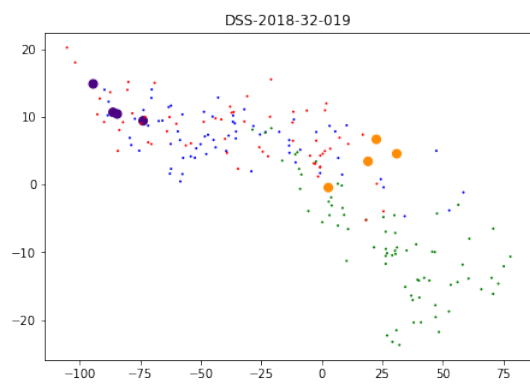
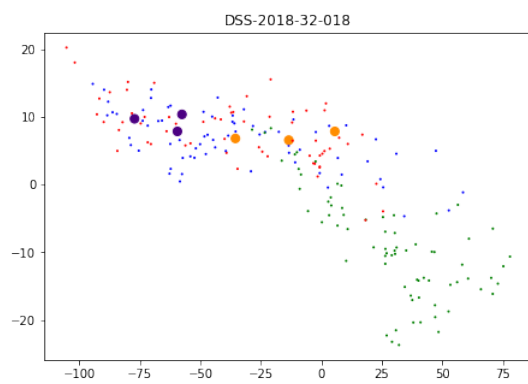
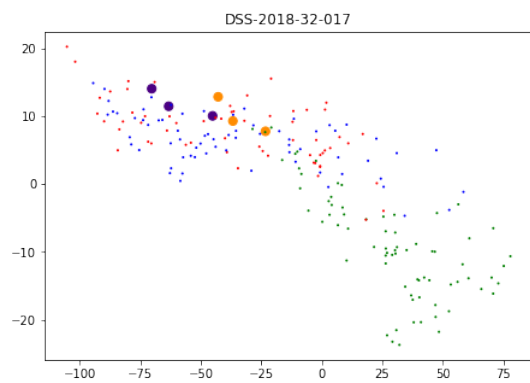
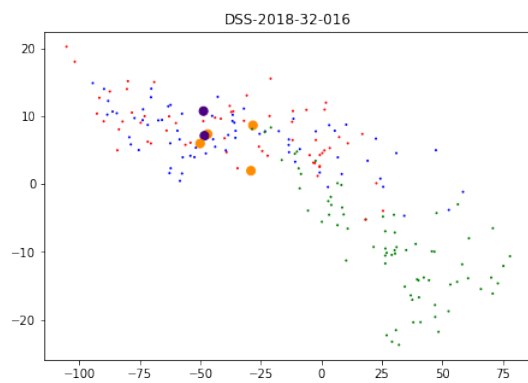
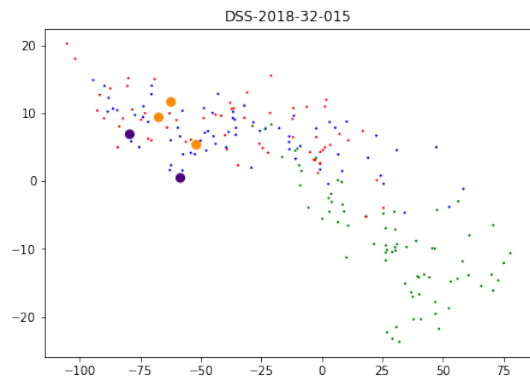
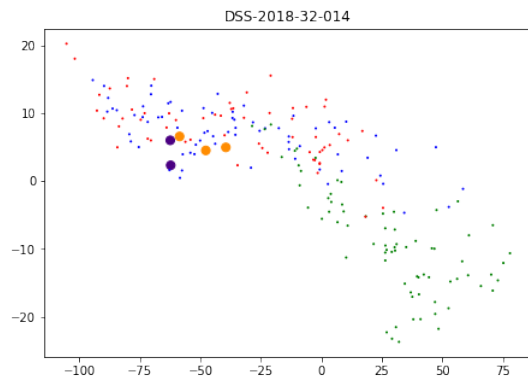
        ax.scatter(slidetsne[:, 0], slidetsne[:, 1],c=newcolors,s=sizelist)

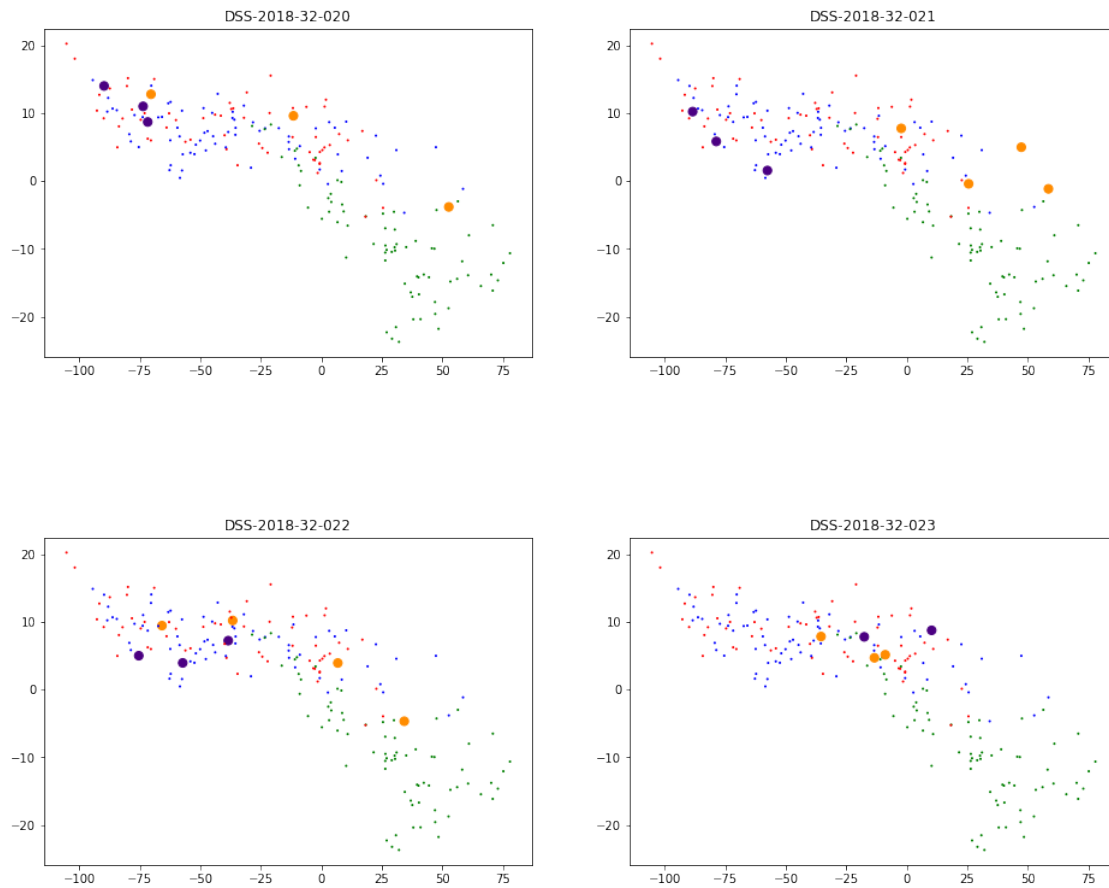
    pltindex+=1;

    if pltindex>2:
        pltindex=1;

```







```
[548]: pltindex=1;

for i in dssdict["dss"]:
    sizelist=[];
    newcolors=[];

    for index in range(len(slideid)):

        if slideid[index].startswith(i+"_"):
            sizelist.append(50)

            location = slideid[index].split("_")[1];
            location=location[0];
            if location == "a":
                newcolors.append("indigo")
            elif location == "b":
                newcolors.append("darkorange")
        else:
            sizelist.append(1)
```

```

        newcolors.append(slidecolors[index])
    if pltindex==1:
        fig = plt.figure(figsize=(16, 5))
        ax = fig.add_subplot(120+pltindex)

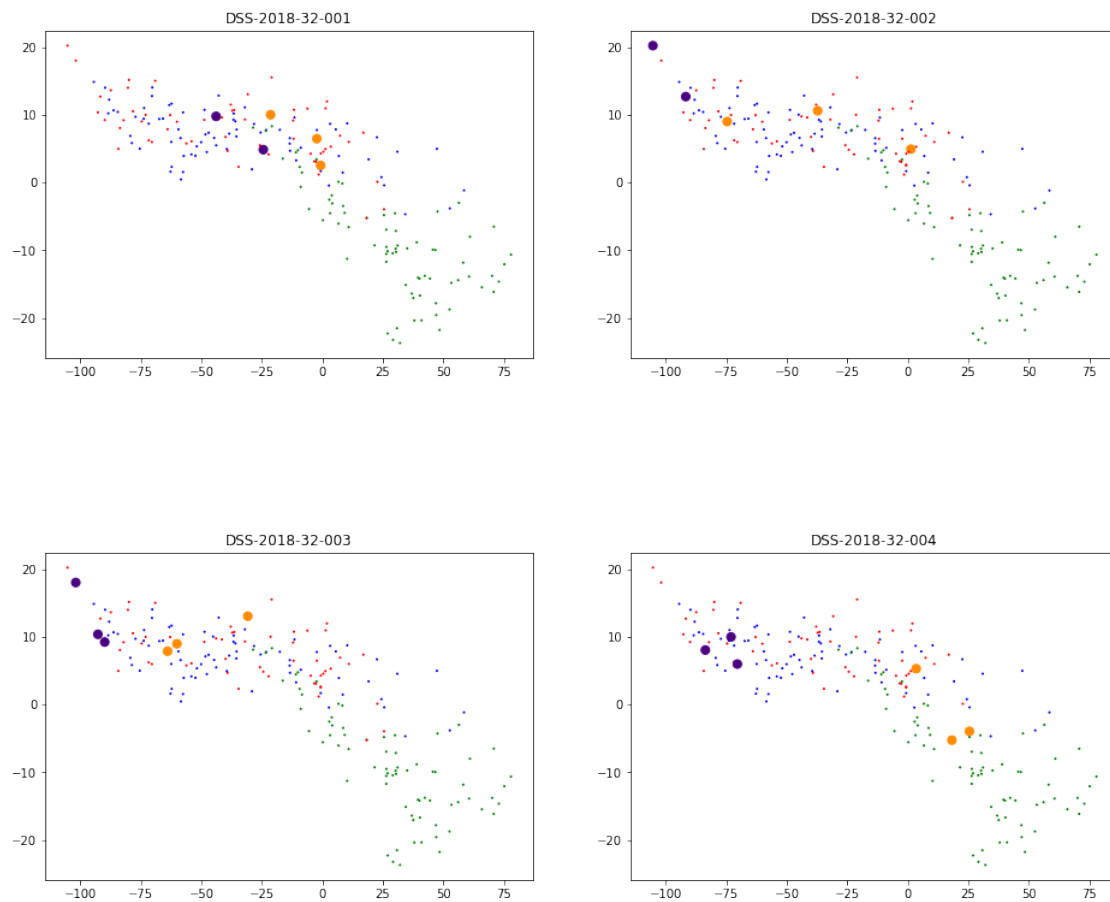
        ax.set_title(i)

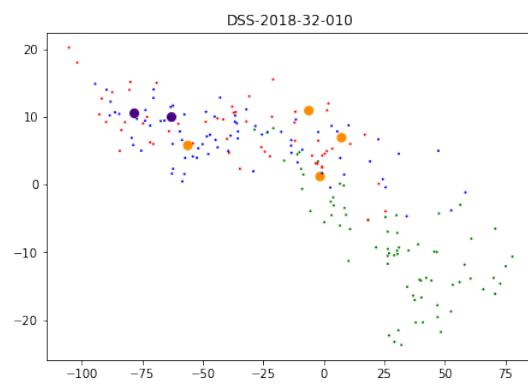
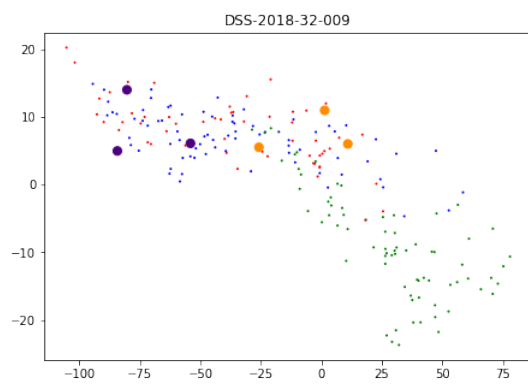
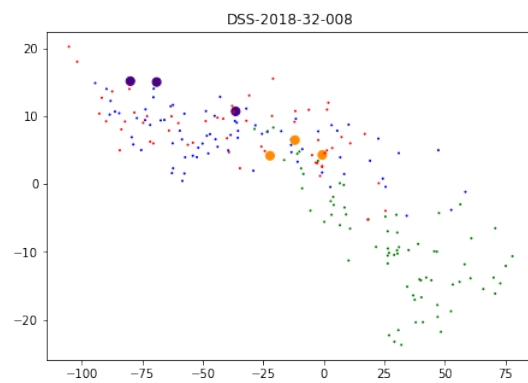
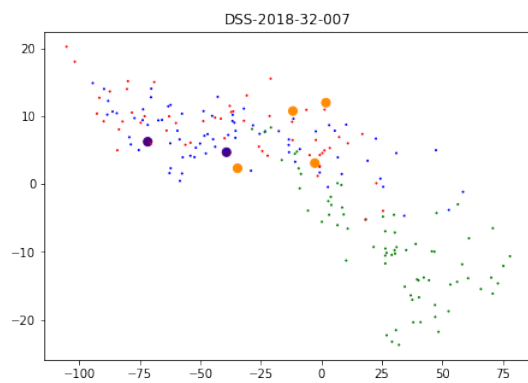
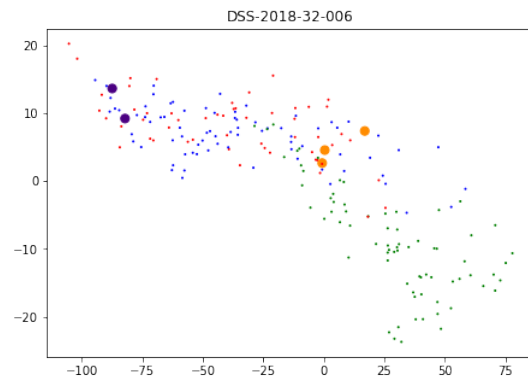
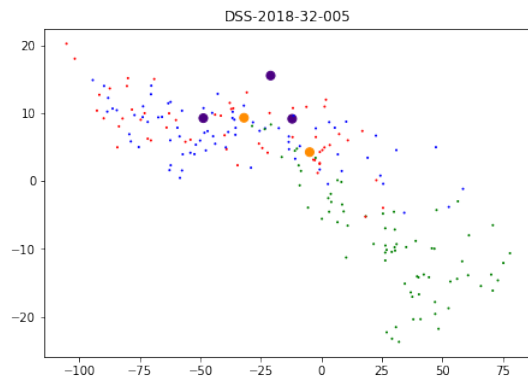
        ax.scatter(slidetsne[:, 0], slidetsne[:, 1],c=newcolors,s=sizelist)

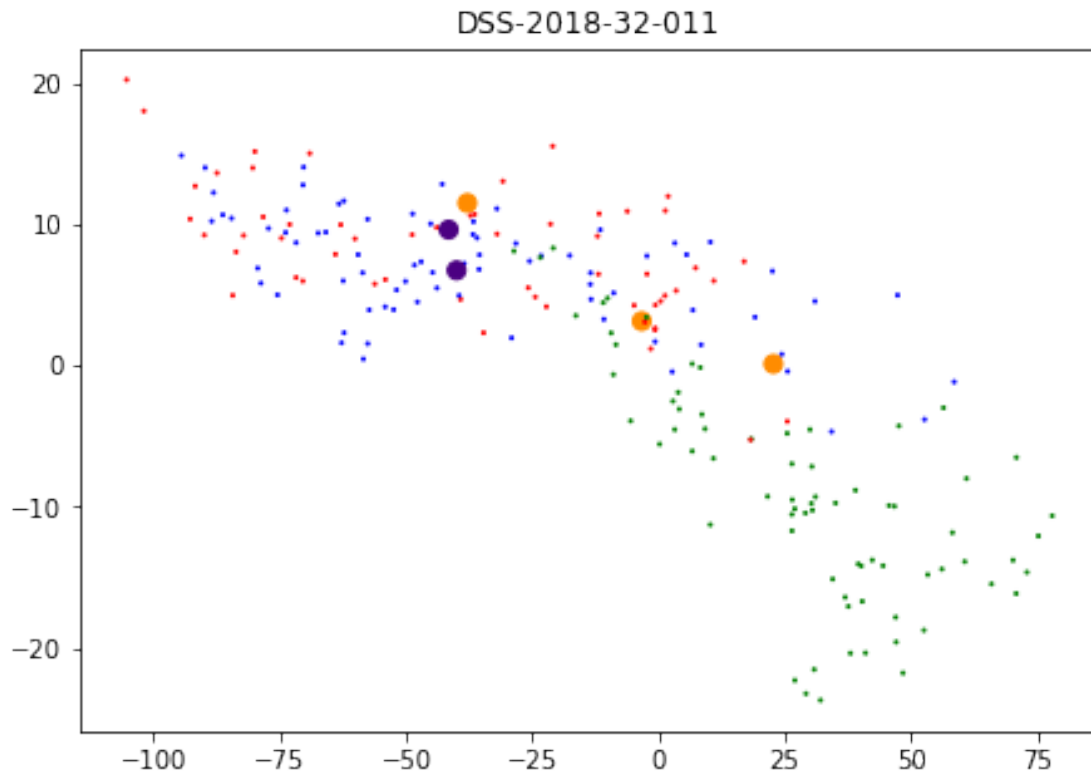
    pltindex+=1;

    if pltindex>2:
        pltindex=1;

```







```
[549]: pltindex=1;

for i in dssdict["normal"]:
    sizelist=[];
    newcolors=[];

    for index in range(len(slideid)):

        if slideid[index].startswith(i+"_"):
            sizelist.append(50)

            location = slideid[index].split("_")[1];
            location=location[0];
            if location == "a":
                newcolors.append("indigo")
            elif location == "b":
                newcolors.append("darkorange")
        else:
            sizelist.append(1)
            newcolors.append(slidecolors[index])
    if pltindex==1:
```



```

fig = plt.figure(figsize=(16, 5))
ax = fig.add_subplot(120+pltindex)

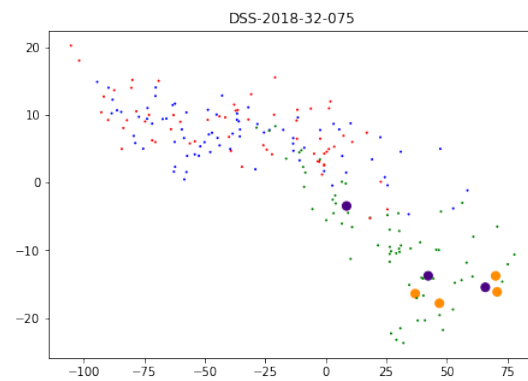
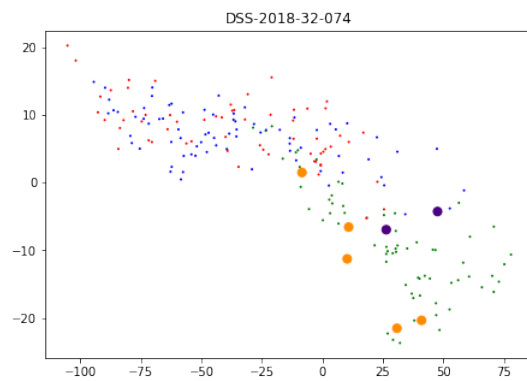
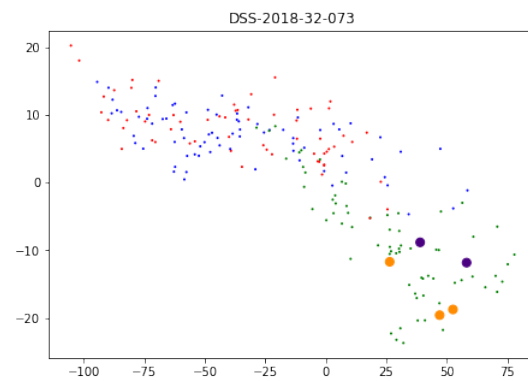
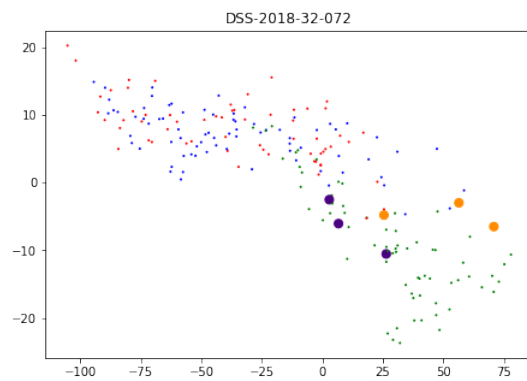
ax.set_title(i)

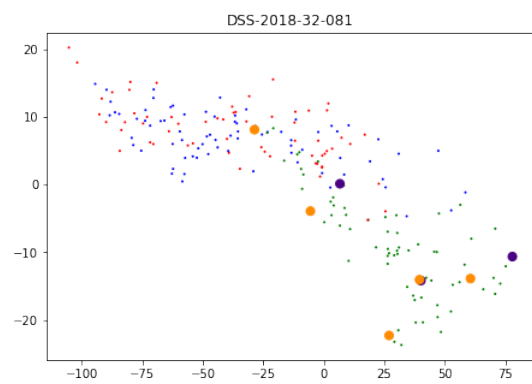
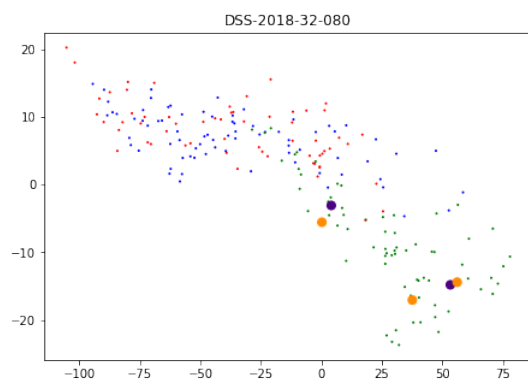
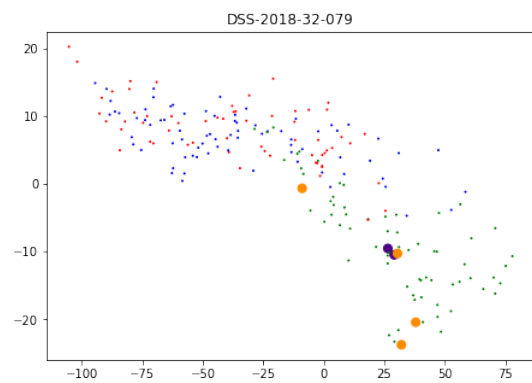
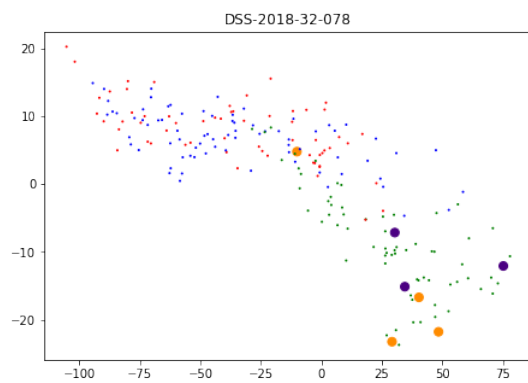
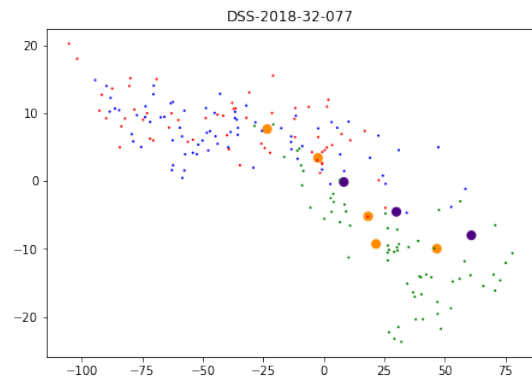
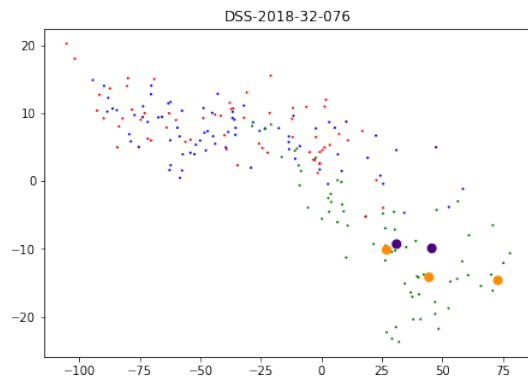
ax.scatter(slidetsne[:, 0], slidetsne[:, 1], c=newcolors, s=sizelist)

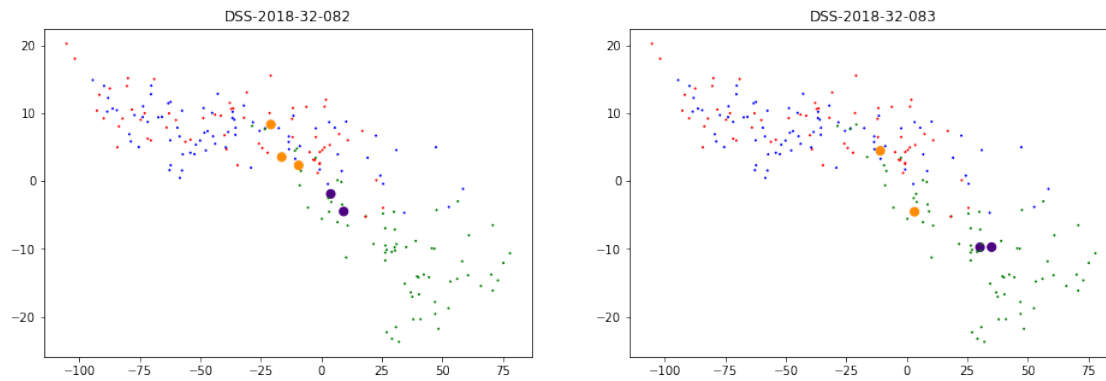
pltindex+=1;

if pltindex>2:
    pltindex=1;

```







[]:

```
[568]: hc=sklearn.cluster.AgglomerativeClustering(n_clusters = 6,).
        fit_predict(slidetsne)
```

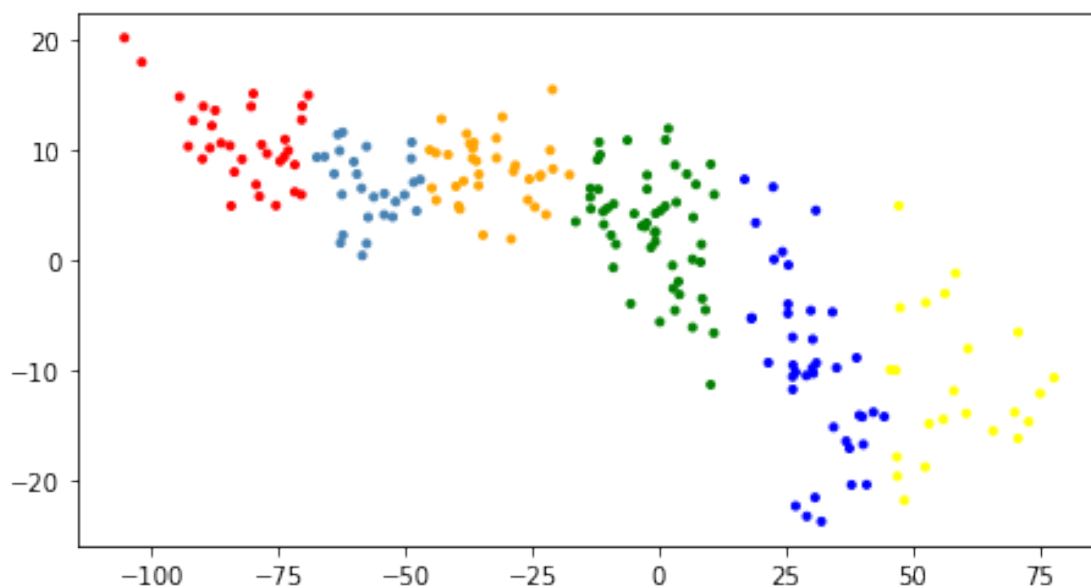
```
colordict=["blue","red","green","orange","yellow","steelblue","#aabbcc","#aabb00","#aa43cc","#
colorlabels=[]
for i in hc:
    colorlabels.append(colordict[i])
```

```
[569]: fig = plt.figure(figsize=(7.5, 4))

ax = fig.add_subplot(111)

ax.scatter(slidetsne[:, 0], slidetsne[:, 1],c=colorlabels,s=9)
```

```
[569]: <matplotlib.collections.PathCollection at 0x7ff33c17e550>
```



```
[570]: #score 2.1
```

```
[571]: score21=[]  
for i in range(len(slideid)):  
    hcolor = colorlabels[i]  
    if hcolor == "red":  
        score21.append(4)  
    elif hcolor == "steelblue":  
        score21.append(3)  
    elif hcolor == "orange":  
        score21.append(2)  
    elif hcolor == "green":  
        score21.append(1)  
    else:  
        score21.append(0)
```

```
[ ]: #score 2.2
```

```
[574]: threshold = 0.3
```

```
[573]: right=np.max(slidetsne[:, 0])  
left=np.min(slidetsne[:, 0])  
top=np.max(slidetsne[:, 1])  
bottom=np.min(slidetsne[:, 1])
```

```
[576]: xlength = right-left  
ylength = top - bottom
```

```
[586]: levels=4  
  
xlength2 =(1-threshold)*xlength  
ylength2 =(1-threshold)*ylength  
  
gapx=xlength2/4  
gapy=ylength2/4
```

```
[595]: score22=[]  
for i in range(len(slideid)):  
    x=slidetsne[i][0];  
    y=slidetsne[i][1];  
  
    xvalue = 0;  
    yvalue = 0;  
  
    if x - left > (1-threshold)*xlength :  
        xvalue = 0;  
    else:  
        temp =x-left;
```

```

        lv = temp/gapx
        xvalue = 4-(lv)

    if y- bottom < threshold*ylength:
        yvalue=0;

    else:
        temp = y-bottom-threshold*ylength
        lv = temp/gapy
        yvalue=(lv)

    score22.append(xvalue/2 + yvalue/2 )

```

```
[ ]:
```

```
[644]: scoredata_head = scoredata[0]
```

```
[647]: scoredata_head[23]
```

```
[647]: 'gland_divid_mucosa'
```

```
[654]: myscore=[]
```

```

for i in scoredata[1:]:
    gdm = float(i[23])*10;
    score = 4-gdm;
    if score <0:
        score =0;

    myscore.append(score)

```

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