Submit

Please build a folder with your netID in **lowercase**, for example, *abc*123456.

Put all outputfiles (including the ones from reducedim1.py and reducedim2.py) in the folder and turn the whole folder to TA. There should be 30 + 12 files in the folder.

The output files should be the exactly same as given.

Part I

There are 3 test datasets: iris, wine and seed.

The .data and .labels files are the inputs. the .txt files are the outputs.

Run the following commands:

python3 pca1.py iris.data iris.labels pca1_vec_iris.txt pca1_reduced_iris.txt python3 pca1.py wine.data wine.labels pca1_vec_wine.txt pca1_reduced_wine.txt python3 pca1.py seed.data seed.labels pca1_vec_seed.txt pca1_reduced_seed.txt

python3 pca2.py iris.data iris.labels pca2_vec_iris.txt pca2_reduced_iris.txt python3 pca2.py wine.data wine.labels pca2_vec_wine.txt pca2_reduced_wine.txt python3 pca2.py seed.data seed.labels pca2_vec_seed.txt pca2_reduced_seed.txt

python3 scatter1.py iris.data iris.labels scatter1_vec_iris.txt scatter1_reduced_iris.txt python3 scatter1.py wine.data wine.labels scatter1_vec_wine.txt scatter1_reduced_wine.txt python3 scatter1.py seed.data seed.labels scatter1_vec_seed.txt scatter1_reduced_seed.txt

python3 scatter2.py iris.data iris.labels scatter2_vec_iris.txt scatter2_reduced_iris.txt python3 scatter2.py wine.data wine.labels scatter2_vec_wine.txt scatter2_reduced_wine.txt python3 scatter2.py seed.data seed.labels scatter2_vec_seed.txt scatter2_reduced_seed.txt

python3 scatter3.py iris.data iris.labels scatter3_vec_iris.txt scatter3_reduced_iris.txt python3 scatter3.py wine.data wine.labels scatter3_vec_wine.txt scatter3_reduced_wine.txt python3 scatter3.py seed.data seed.labels scatter3_vec_seed.txt scatter3_reduced_seed.txt

Part II

In Part II, there are 2 test datasets: iris, wine.

Run the followin commands:

Case 1:

python3 reducedim1.py iris.data iris.labels reducedim1_vec_iris.txt reducedim1_reduced_iris.txt python3 nearest.py reducedim1_reduced_iris.txt reducedim1_vec_iris.txt iris.labels reducedim1_iris_query.data reducedim1_iris_query.labels reducedim1_nn_iris.txt

 $python 3\ reducedim 1. py\ wine. data\ wine. labels\ reducedim 1_ vec_wine. txt\ reducedim 1_ reduced_wine. txt\ python 3\ nearest. py\ reducedim 1_ reduced_wine. txt\ reducedim 1_ vec_wine. txt\ wine. labels\ reducedim 1_ wine_ query. data\ reducedim 1_ wine_ query. labels\ reducedim 1_ nn_ wine. txt$

Case 2:

python3 reducedim2.py iris.data iris.labels reducedim2_vec_iris.txt reducedim2_reduced_iris.txt python3 nearest.py reducedim2_reduced_iris.txt reducedim2_vec_iris.txt iris.labels reducedim2_iris_query.data reducedim2_iris_query.labels reducedim2_nn_iris.txt

python3 reducedim2.py wine.data wine.labels reducedim2_vec_wine.txt reducedim2_reduced_wine.txt python3 nearest.py reducedim2_reduced_wine.txt reducedim2_vec_wine.txt wine.labels reducedim2_wine_query.data reducedim2_wine_query.labels reducedim2_nn_wine.txt