```
import numpy as np
from sklearn import svm
from google.colab import files
from sklearn.manifold import TSNE
import matplotlib.pyplot as plt
import pickle
uploaded = files.upload()
for fn in uploaded.keys():
 print('User uploaded file "{name}" with length {length} bytes'.format(
      name=fn, length=len(uploaded[fn])))
С⇒
     Choose Files 4 files
       elbow_table.txt(text/plain) - 31134 bytes, last modified: 5/4/2019 - 100% done
      arm_straight.txt(text/plain) - 29533 bytes, last modified: 5/4/2019 - 100% done

    arm_down.txt(text/plain) - 33172 bytes, last modified: 5/4/2019 - 100% done

    arm_across.txt(text/plain) - 28697 bytes, last modified: 5/4/2019 - 100% done

    Saving elbow_table.txt to elbow_table (1).txt
    Saving arm straight.txt to arm straight (1).txt
    Saving arm down.txt to arm down (1).txt
    Saving arm across.txt to arm across (1).txt
    User uploaded file "elbow table.txt" with length 31134 bytes
    User uploaded file "arm_across.txt" with length 28697 bytes
    User uploaded file "arm down.txt" with length 33172 bytes
    User uploaded file "arm straight.txt" with length 29533 bytes
arr list = []
data = np.zeros([1, 6])
labels = np.zeros([1, 1])
for k, v in uploaded.items():
 print(k)
 text_file = open(k, "w")
 text_file.write(uploaded[k].decode("utf-8") )
 text file.close()
 # Thanks to Aura Barrera for this short cut!
 arr = np.loadtxt(fname=k, encoding="ISO-8859-1")
 print(arr.shape)
 arr list.append(arr.shape)
 data = np.vstack((data, arr))
 num data = arr.shape[0]
 labels = np.vstack((labels, np.full((num_data, 1), k)))
data = np.delete(data, (0), axis=0)
labels = np.delete(labels, (0), axis=0)
    elbow table.txt
     (357, 6)
    arm across.txt
     (333, 6)
    arm down.txt
     (388, 6)
    arm straight.txt
     (334, 6)
svm model = svm.SVC(gamma='scale')
svm model.fit(data, labels )
svm model.get params()
```

```
C→
      {'C': 1.0,
        'cache_size': 200,
        'class weight': None,
        'coef0': 0.0,
        'decision function shape': 'ovr',
        'degree': 3,
        'gamma': 'scale',
        'kernel': 'rbf',
        'max iter': -1,
        'probability': False,
        'random state': None,
        'shrinking': True,
        'tol': 0.001,
        'verbose': False}
pickle fname = "pickle elnaz model.sav"
pickle.dump(svm model, open(pickle fname, "wb"))
from google.colab import files
files.download(pickle fname)
# random testing
point1 = np.asarray([[7.28, 0.1915, -7.867, -5.02, 1.67, -8.27]]) # should be arm strepoint2 = np.asarray([[2.79, 9.82, -1.939, -1.79, 9.32, 1.4]]) # should be arm down point3 = np.asarray([[8.56, 2.87, -6.00, -6.7, 6.45, -2.2]]) # should be arm across point4 = np.asarray([[5, 5, 5, 5, 5]]) # should be random
svm model.predict(point3)
      array([u'elbow table.txt'], dtype='<U32')</pre>
reduced = TSNE(n components=2).fit transform(data)
index = 0
for sh in arr list:
  plt.scatter \overline{(reduced[index:index + sh[0], 0], reduced[index: index + sh[0], 1])}
   index += sh[0]
plt.show()
 \Box
         60
         40
         20
          0
       -20
       -40
       -60
                             -<u>2</u>0
                                                    20
                   -40
                                                               40
```

# import sklearn.discriminant\_analysis as QDA
from sklearn.discriminant analysis import QuadraticDiscriminantAnalysis as QDA