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Code for problem 4.
import numpy as np
import os
os.chdir('/home/tuomas/Documents/DATA.STAT.770/koe')
#%%
iris_data = np.loadtxt('iris-commaseparated.txt', delimiter=',')
iris_data = np.concatenate((np.array([[5.1,3.5,1.4,0.2,0]]), iris_data))
labels = iris_data[:,-1]
iris data = iris data[:,0:4]
#%% a)
from sklearn.decomposition import PCA
pca = PCA(n_components=2)
pca.fit(iris_data)
iris reduced = pca.transform(iris data)
#%% Plot
import matplotlib.pyplot as plt
iris1 = iris reduced[(labels==0),:]
iris2 = iris_reduced[(labels==1),:]
iris3 = iris_reduced[(labels==2),:]
plt.plot(iris1[:,0], iris1[:,1], 'bo')
plt.plot(iris2[:,0], iris2[:,1], 'ro')
plt.plot(iris3[:,0], iris3[:,1], 'go')
plt.title('Blue=setosa\n Red=versicolor\n Green=virginica')
#%% b)
from numpy.linalg import norm
prop_of_var = np.sum(pca.explained_variance_ratio_)
print('Proportion of variance explained = {}'.format(prop_of_var))
back proj = pca.inverse transform(iris reduced)
loss = norm((iris data-back proj), None)
print('Reconstruction error = {}'.format(loss))
#%% c)
from sklearn.manifold import TSNE
semeion_data = np.loadtxt('semeion-commaseparated.txt', delimiter=',')
labels = semeion_data[:,-1]
semeion_data = semeion_data[:,0:256]
#%%
tsne = TSNE(n components=2)
semeion_proj = tsne.fit_transform(semeion_data)
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