# sheet06

May 31, 2022

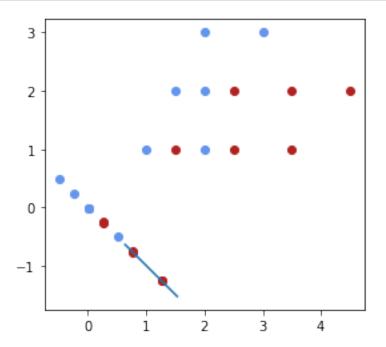
#### 0.0.1 Exercise 13

## 0.1 c)

```
[1]: import numpy as np
     import matplotlib.pyplot as plt
     import matplotlib.axes as ax
     import sympy as smp
     #define populations per axis
     pop0x = [1, 2, 1.5, 2, 2, 3]
     pop0y = [1, 1, 2, 2, 3, 3]
     pop1x = [1.5, 2.5, 3.5, 2.5, 3.5, 4.5]
     pop1y = [1, 1, 1, 2, 2, 2]
     #linspace for the plot below. the vector lambda doesn't need limitaions, but
      →limitations are used here to make a better plot.
     # the limitation -2.5 is the lambda_cut
     x = np.linspace(-6, -2.5)
     #plot of the data and lambda with a cutoff at lambda_cut
     fig = plt.figure()
     axx = fig.add_subplot(111, aspect='equal')
     axx.scatter(pop0x, pop0y, c = "cornflowerblue")
     axx.scatter(pop1x, pop1y, c = "firebrick")
     axx.plot(-(370/1447)*x,(367/1447)*x)
     # d.)
     #the following to functions create the projection onto lambda
     def H(x,y,param1,param2):
        return param1*x+param2*y
     def projection(x1,y1,x2,y2,param1,param2):
         a = H(x1,y1,param1,param2)
         b = H(x2,y2,param1,param2)
         s, t = smp.symbols('s, t', real = True)
         s_1 = smp.solve(param1*(param1*s)+param2*(param2*s)-a,s)
         s_2 = smp.solve(param1*(param1*t)+param2*(param2*t)-b,t)
```

```
return [s_1,s_2]

#projection of each datapoint onto lambda
for i in range(len(pop0x)):
    p = projection(pop0x[i],pop0y[i],pop1x[i],pop1y[i],-370,367)
    axx.scatter(-370*p[0][0],367*p[0][0], c = "cornflowerblue")
    axx.scatter(-370*p[1][0],367*p[1][0], c = "firebrick")
```



### 0.1.1 e)

 $\lambda_{\rm cut} = -2.5$  is choosen, because with this  $\lambda_{\rm cut}$  we have no false true datapoints. The rusulting vector, with  $\forall \lambda < -2.5$ , is shown in the above output.

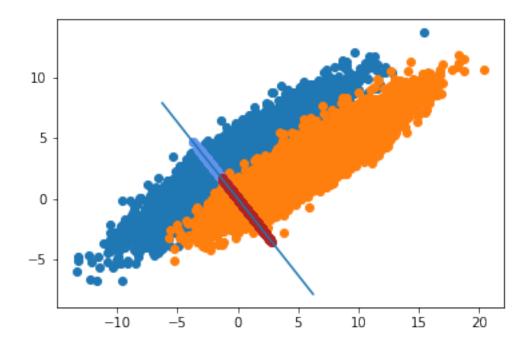
### 0.2 Exercise 13

```
[2]: import pandas as pd
  import numpy as np
  import numpy.linalg as alg
  import matplotlib.pyplot as plt
  import matplotlib.axes as ax
  #reading the data from the .h5-file
  p0 = pd.read_hdf('two_populations.h5', key='P_0_10000')
  p1 = pd.read_hdf('two_populations.h5', key='P_1')
  p0_1000 = pd.read_hdf('two_populations.h5', key='P_0_1000')
  # a)
```

```
#converting pd.dataframe into numpy arrays
pOn = p0.to_numpy()
p1n = p1.to_numpy()
#calculation of the mean vectors
mu1 = [np.mean(p0n[:,0]), np.mean(p0n[:,1])]
mu2 = [np.mean(p1n[:,0]), np.mean(p1n[:,1])]
#print(p0)
# b)
#calculation of the covariance matrices
VP1 = np.cov([p0n[:,0], p0n[:,1]])
#print(VP1)
VP2 = np.cov([p1n[:,0], p1n[:,1]])
#print(VP2)
VP12 = VP1 + VP2
#print(VP12)
# c)
#calculation and plot of lambda with the data
e_lam = np.matmul(alg.inv(VP12), [mu1[0]-mu2[0], mu1[1]-mu2[1]])
#print(e lam)
x = np.linspace(-5,5,10000)
fig = plt.figure()
axd = fig.add_subplot(111, aspect='equal')
axd.scatter(p0n[:,0], p0n[:,1])
axd.scatter(p1n[:,0], p1n[:,1])
axd.plot(e_lam[0]*x,e_lam[1]*x)
# d)
x1 = np.zeros(len(p0n[:,0]))
x2 = np.zeros(len(p0n[:,0]))
x3 = np.zeros(len(p0n[:,0]))
x4 = np.zeros(len(p0n[:,0]))
for i in range(len(p0n[:,0])):
    p = projection(p0n[:,0][i],p0n[:,1][i],p1n[:,0][i],p1n[:
 \rightarrow,1][i],e_lam[0],e_lam[1])
    #print(p[0][0])
    x1[i] = p[0][0]
    \#x2[i] = e_lam[1]*p[0][0]
    x3[i] = p[1][0]
    #x4[i] = e_lam[1]*p[1][0]
    #print("p00",i,p[0][0])
```

```
#print("p10",p[1][0])
#print("x1",x1)
#print("x3",x3)
#print(e_lam[0])
#print(e_lam[1])
axd.scatter(e_lam[0]*x1,e_lam[1]*x1, c = "cornflowerblue")
axd.scatter(e_lam[0]*x3,e_lam[1]*x3, c = "firebrick")
```

[2]: <matplotlib.collections.PathCollection at 0x7f9a533a5550>



[]: