# solution05

November 10, 2020

## Exercise Sheet 5 Validation & Regularization

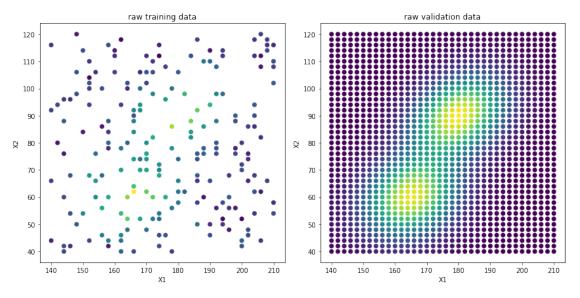
```
[1]: import numpy as np
   from mpl_toolkits.mplot3d import Axes3D
   import matplotlib.pyplot as plt
   from numpy import linalg as LA
   import scipy.cluster.vq as vq
   mm = np.matmul
```

```
[2]: # abbreviations
             training input
    # Xt
    # Xv
             validation input
             training output
    # Yt
    # Yυ
             validation output
             centered training input
    # Xtc
             decorrelated training input
    # Xtd
    # Xtw
             whitened training input
    # Xtwm
             monomials of training input after whitening
    # Wtwmct weight resulted from training set after cross validation training
     \rightarrow with optimum LambdaT
    # Wvwmcq weight resulted from validation set after cross validation training
     →with optimum LambdaG
    # Yp
               predicted output
```

#### 5.1: Cross-validation

#### (a) Preprocessing

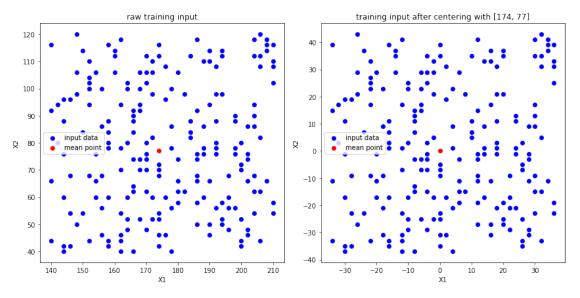
```
plt.scatter(Xt[0,:],Xt[1,:],c=Yt)
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('raw training data')
plt.subplot(1,2,2)
plt.scatter(Xv[0,:],Xv[1,:],c=Yv)
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('raw validation data')
plt.tight_layout()
plt.show()
```



```
[]:
[]:
[4]: def centering(x):
    return np.median(x, axis=1)
    xc = centering(Xt)
    Xtc = Xt - xc.reshape(2,1) # centered inputs

# plotting (no computation value)
plt.figure(figsize=(12,6))
plt.subplot(1,2,1)
plt.scatter(Xt[0,:],Xt[1,:],color='b',label='input data')
plt.scatter(xc[0],xc[1],color='r',label='mean point')
plt.xlabel('X1')
plt.ylabel('X2')
```

```
plt.title('raw training input')
plt.legend()
plt.subplot(1,2,2)
plt.scatter(Xtc[0,:],Xtc[1,:],color='b',label='input data')
plt.scatter(centering(Xtc)[0],centering(Xtc)[1],color='r',label='mean point')
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('training input after centering with [%d, %d]'%(xc[0],xc[1]))
plt.tight_layout()
plt.legend()
plt.show()
```



```
[5]: def covariance(x):
    return mm(x,x.T)/np.shape(x)[1]

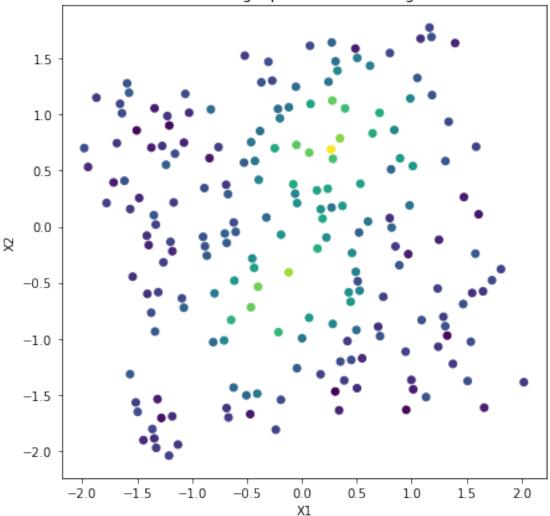
# Covarience, eigenvalues and eigenvectors based on training data
C = covariance(Xtc)
A, V = LA.eig(C) # A: Eigenvalues V: Eigenvectors

# whitening of the training data
Xtd = mm(V.T,Xtc) # decorrelation
Xtw = (np.sqrt(1/A).reshape(2,1)*Xtd) # whitening

# plotting (no computation value)
plt.figure(figsize=(7,7))
plt.scatter(Xtw[0,:],Xtw[1,:],c=Yt)
plt.xlabel('X1')
plt.ylabel('X2')
```

```
plt.title('training input after whitening')
plt.show()
```



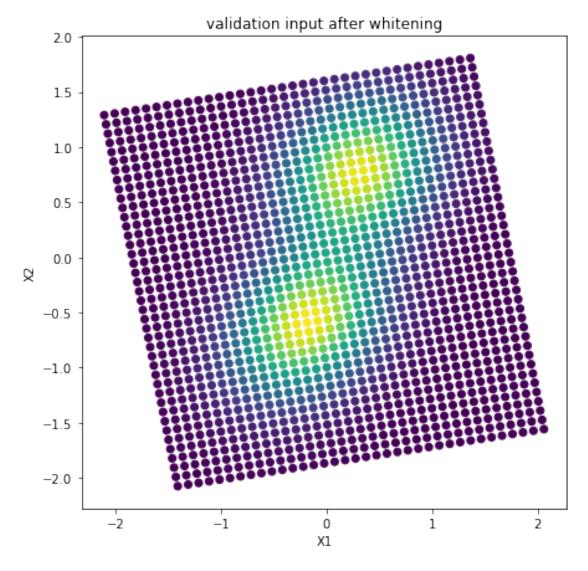


```
[6]: # test of successful whitening:
print(np.round(covariance(Xtw),6))
# should result in Identity matrix
```

[[1. 0.] [0. 1.]]

```
[7]: # whitening of the validation data
Xvc = Xv - xc.reshape(2,1) # centered the validation input
Xvd = mm(V.T,Xvc) # decorrelation
Xvw = (np.sqrt(1/A).reshape(2,1)*Xvd) # whitening
```

```
# plotting (no computation value)
plt.figure(figsize=(7,7))
plt.scatter(Xvw[0,:],Xvw[1,:],c=Yv)
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('validation input after whitening')
plt.show()
```



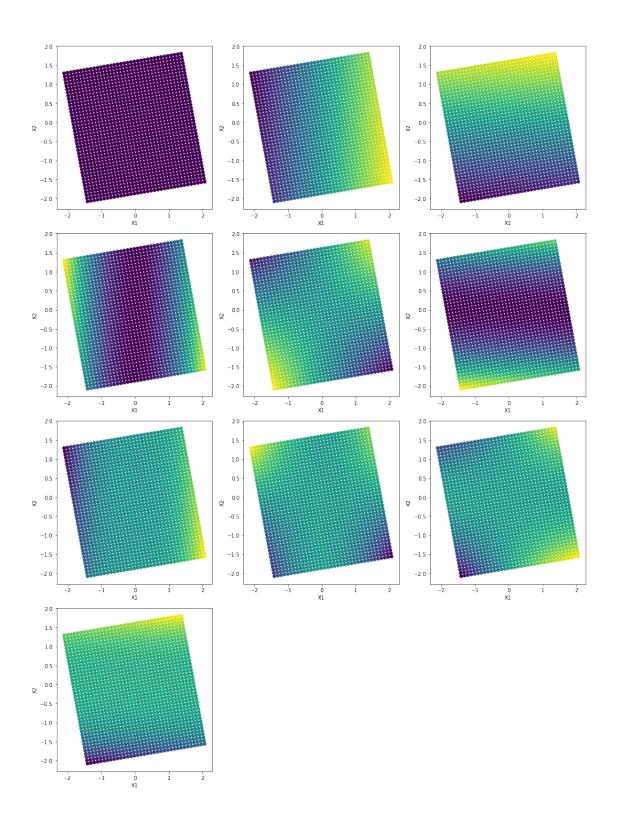
```
[8]: # Xtw is the training input after whitening
# Xvw is the validation input after whitening
# A is the eigenvalues matrix
# V is the eigenvectors matrix
```

### (b) Plot monomials

```
[9]: # monomial
     \operatorname{def\ arisum}(n)\colon # arithmetic sum of combination sequences only for X with dim. 2
         s = 0
         for i in range(n+2):
             s += i
         return s
     def monomial(X,k): # calculating the monomial series of order k
         Xm = np.zeros((arisum(k),np.shape(X)[1]))
         for j in range(np.shape(X)[1]):
             i = 0
             for n in range(k+1):
                  1 = n
                 m = 0
                  while 1+m == n and 1>= 0:
                      Xm[i,j] = ((X[0,j]**1)*(X[1,j]**m))
                      1 -= 1
                      m += 1
                      i += 1
                  if i == arisum(k):
                      break
         return Xm
```

```
[10]: Xvwm3 = monomial(Xvw,3) # monomials using the white validation set

# plotting (no computation value)
# the monomial as a function of x1 and x1
plt.figure(figsize=(15,20))
for i in range(10):
    plt.subplot(4,3,i+1)
    plt.scatter(Xvw[0,:],Xvw[1,:],c=Xvwm3[i,:])
    plt.xlabel('X1')
    plt.ylabel('X2')
plt.tight_layout()
plt.show()
```

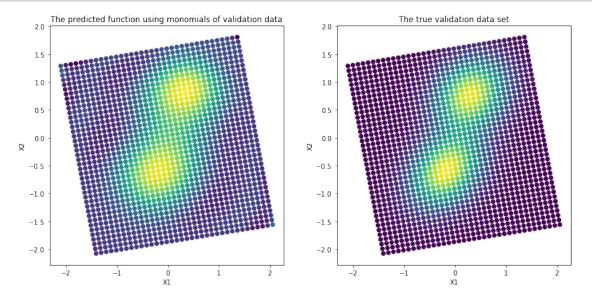


```
[11]: Xtwm = monomial(Xtw,9) # monomials of training set
Xvwm = monomial(Xvw,9) # monomials of validation set
```

```
Wtwm = mm(LA.inv(mm(Xtwm,Xtwm.T)),mm(Xtwm,Yt.T)) # weights as validation set
Wvwm = mm(LA.inv(mm(Xvwm,Xvwm.T)),mm(Xvwm,Yv.T)) # weights as training set

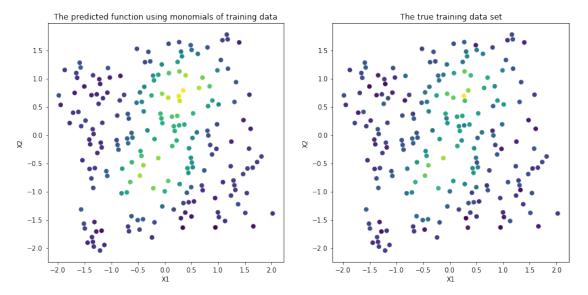
Yts = mm(Xtwm.T,Wtwm) # simulated output based on training data
Yvs = mm(Xvwm.T,Wvwm) # simulated output based on validation data
```

```
[12]: # prediction vs validation
plt.figure(figsize=(12,6))
plt.subplot(1,2,1)
plt.scatter(Xvw[0,:],Xvw[1,:],c=Yvs)
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('The predicted function using monomials of validation data')
plt.subplot(1,2,2)
plt.scatter(Xvw[0,:],Xvw[1,:],c=Yv)
plt.xlabel('X1')
plt.ylabel('X1')
plt.ylabel('X2')
plt.title('The true validation data set')
plt.tight_layout()
plt.show()
```



```
[13]: # prediction vs Training
   plt.figure(figsize=(12,6))
   plt.subplot(1,2,1)
   plt.scatter(Xtw[0,:],Xtw[1,:],c=Yts)
   plt.xlabel('X1')
   plt.ylabel('X2')
```

```
plt.title('The predicted function using monomials of training data')
plt.subplot(1,2,2)
plt.scatter(Xtw[0,:],Xtw[1,:],c=Yt)
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('The true training data set')
plt.tight_layout()
plt.show()
```

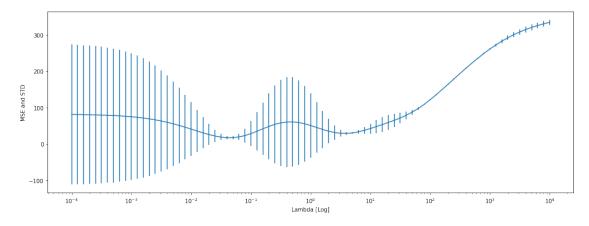


#### (c) apply regularization using a weight-decay term

```
[14]: def _MSE_(W,X,Y):
          Ys = mm(W.T,X)
          sqe = ((Y-Ys)**2)
          return np.sum(sqe)/np.shape(X)[1]
      setLambda = 10**(np.linspace(-4,4,81,endpoint=True))
      ebd = [[],[],[]] # error bar data: [Lambda, MSE, STD]
      for Lambda in setLambda:
          ident = np.identity(55)*Lambda
          Err = []
          for fold in range(10):
              vind = list(range(20*fold,20*(fold+1)))
              tind = list(range(0,20*fold))+list(range(20*(fold+1),200))
              Xcv = Xtwm[:,vind] # validation X for cross validation
              Xct = Xtwm[:,tind] # Training X for cross validation
              Ycv = Yt[vind] # validation Y for cross validation
              Yct = Yt[tind] # training Y for cross validation
              Wtwmc = mm(LA.inv(mm(Xct,Xct.T)+ident),mm(Xct,Yct.T)).reshape(55,1)
```

```
Err.append(_MSE_(Wtwmc,Xtwm,Yt)) # calculated the whole training set
Err = np.array(Err)
mse = np.sum(Err)/10
std = np.std(Err)
ebd[0].append(Lambda)
ebd[1].append(mse)
ebd[2].append(std)

plt.figure(figsize=(17,6))
ax = plt.subplot()
plt.errorbar(ebd[0], ebd[1], yerr=ebd[2])
ax.set_xscale("log")
plt.xlabel('Lambda [Log]')
plt.ylabel('MSE and STD')
plt.show()
```

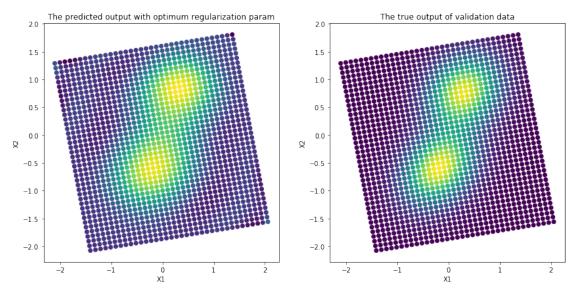


The Lambda at which the MSE is minimum over the whole training set: best regularization parameter is 0.0398

```
[16]: # using best regularization parameter
ident = np.identity(55)*LambdaT
Wvwmc = mm(LA.inv(mm(Xvwm,Xvwm.T)+ident),mm(Xvwm,Yv.T)).reshape(55,1)
Yp = mm(Wvwmc.T,Xvwm)

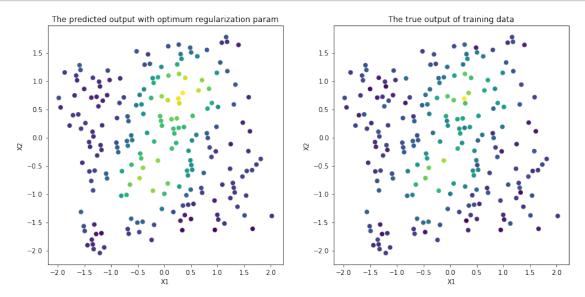
# plotting the raw data
plt.figure(figsize=(12,6))
plt.subplot(1,2,1)
plt.scatter(Xvw[0,:],Xvw[1,:],c=Yp[0,:])
```

```
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('The predicted output with optimum regularization param')
plt.subplot(1,2,2)
plt.scatter(Xvw[0,:],Xvw[1,:],c=Yv)
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('The true output of validation data')
plt.tight_layout()
plt.show()
```



```
[17]: # using best regularization parameter
      ident = np.identity(55)*LambdaT
      Wtwmc = mm(LA.inv(mm(Xtwm,Xtwm.T)+ident),mm(Xtwm,Yt.T)).reshape(55,1)
      Yp = mm(Wtwmc.T,Xtwm)
      # plotting the raw data
      plt.figure(figsize=(12,6))
      plt.subplot(1,2,1)
      plt.scatter(Xtw[0,:],Xtw[1,:],c=Yp[0,:])
      plt.xlabel('X1')
      plt.ylabel('X2')
      plt.title('The predicted output with optimum regularization param')
      plt.subplot(1,2,2)
      plt.scatter(Xtw[0,:],Xtw[1,:],c=Yt)
      plt.xlabel('X1')
      plt.ylabel('X2')
      plt.title('The true output of training data')
```

```
plt.tight_layout()
plt.show()
```

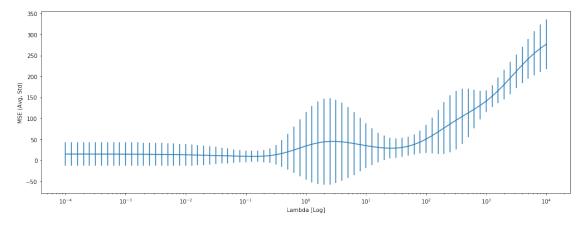


# (d) Compare empirical estimates of bias and variance with the true generalization error

```
[26]: def MSE (W,X,Y):
          Ys = mm(W.T,X)
          sqe = ((Y-Ys)**2)
          return np.sum(sqe)/np.shape(X)[1]
      setLambda = 10**(np.linspace(-4,4,81,endpoint=True))
      ebd = [[],[],[]] # error bar data: [Lambda, MSE, STD]
      for Lambda in setLambda:
          ident = np.identity(55)*Lambda
          Err = \prod
          for fold in range(9): # 9 folds of 164 samples
              vind = list(range(164*fold,164*(fold+1))) #validation indecis
              tind = list(range(0,164*fold))+list(range(164*(fold+1),1476)) #__
       \hookrightarrow training indecis
              Xcv = Xvwm[:,vind] # validation X for cross validation
              Xct = Xvwm[:,tind] # Training X for cross validation
              Ycv = Yv[vind] # validation Y for cross validation
              Yct = Yv[tind] # training Y for cross validation
              Wvwmc = mm(LA.inv(mm(Xct,Xct.T)+ident),mm(Xct,Yct.T)).reshape(55,1)
              Err.append(_MSE_(Wvwmc,Xvwm,Yv)) # calculated the whole training set
          Err = np.array(Err)
          mse = np.sum(Err)/9
          std = np.std(Err)
```

```
ebd[0].append(Lambda)
  ebd[1].append(mse)
  ebd[2].append(std)

plt.figure(figsize=(17,6))
ax = plt.subplot()
plt.errorbar(ebd[0], ebd[1], yerr=ebd[2])
ax.set_xscale("log")
plt.xlabel('Lambda [Log]')
plt.ylabel('MSE (Avg, Std)')
plt.show()
```



The Lambda at which the MSE is minimum over the whole validation set: best regularization parameter is 0.1259

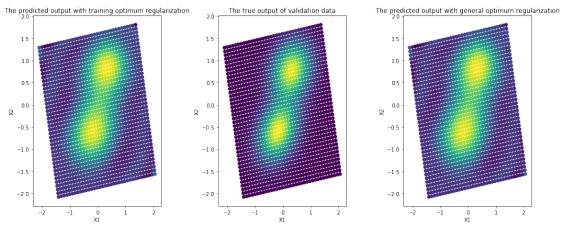
The  $\lambda G$  (= 0.1259) is different than  $\lambda T$  (= 0.0398) but the difference is small.

```
[20]: # using best regularization parameter
identg = np.identity(55)*LambdaG
Wvwmcg = mm(LA.inv(mm(Xvwm,Xvwm.T)+identg),mm(Xvwm,Yv.T)).reshape(55,1)
Ypg = mm(Wvwmcg.T,Xvwm)

identt = np.identity(55)*LambdaT
Wvwmct = mm(LA.inv(mm(Xvwm,Xvwm.T)+identt),mm(Xvwm,Yv.T)).reshape(55,1)
Ypt = mm(Wvwmct.T,Xvwm)

# plotting the raw data
plt.figure(figsize=(15,6))
```

```
plt.subplot(1,3,1)
plt.scatter(Xvw[0,:],Xvw[1,:],c=Ypt[0,:])
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('The predicted output with training optimum regularization')
plt.subplot(1,3,2)
plt.scatter(Xvw[0,:],Xvw[1,:],c=Yv)
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('The true output of validation data')
plt.subplot(1,3,3)
plt.scatter(Xvw[0,:],Xvw[1,:],c=Ypg[0,:])
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('The predicted output with general optimum regularization')
plt.tight_layout()
plt.show()
```

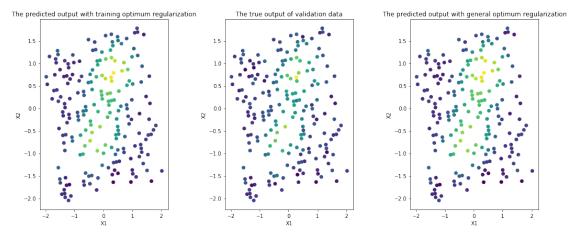


the mean square error for training lambda is 1.7 and for the generalized lambda is 1.8

## (e) Compare generalization and training optimal

```
[22]: # using best regularization parameter
identg = np.identity(55)*LambdaG
Wtwmcg = mm(LA.inv(mm(Xtwm,Xtwm.T)+identg),mm(Xtwm,Yt.T)).reshape(55,1)
Ypg = mm(Wtwmcg.T,Xtwm)
```

```
identt = np.identity(55)*LambdaT
Wtwmct = mm(LA.inv(mm(Xtwm,Xtwm.T)+identt),mm(Xtwm,Yt.T)).reshape(55,1)
Ypt = mm(Wtwmct.T,Xtwm)
# plotting the raw data
plt.figure(figsize=(15,6))
plt.subplot(1,3,1)
plt.scatter(Xtw[0,:],Xtw[1,:],c=Ypt[0,:])
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('The predicted output with training optimum regularization')
plt.subplot(1,3,2)
plt.scatter(Xtw[0,:],Xtw[1,:],c=Yt)
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('The true output of validation data')
plt.subplot(1,3,3)
plt.scatter(Xtw[0,:],Xtw[1,:],c=Ypg[0,:])
plt.xlabel('X1')
plt.ylabel('X2')
plt.title('The predicted output with general optimum regularization')
plt.tight_layout()
plt.show()
```



the mean square error for training lambda is 13.5 and for the generalized lambda is 14.8