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HW<sub>4</sub>

1a) 
$$J(w,a) = (Xw + \alpha 1 - y)^{T}(Xw + \alpha 1 - y) + \lambda w^{T}w$$

$$= \left[\sum_{i=1}^{n} (X_{i1}w_{1} + X_{i2}w_{2} + \dots + X_{id}w_{d} + \alpha - y_{i})^{2}\right] + \lambda(w_{1}^{2} + w_{2}^{2} + \dots + w_{d}^{2})$$

Set  $dJ/d\alpha = 0$  and solve for  $\alpha$ :

$$\begin{split} \frac{dJ}{d\alpha} &= 2 \left[ \sum_{i=1}^{n} \left( X_{i1} w_{1} + X_{i2} w_{2} + \dots + X_{id} w_{d} + \alpha - y_{i} \right) \right] \\ 0 &= 2 \left[ \sum_{i=1}^{n} \left( X_{i1} w_{1} + X_{i2} w_{2} + \dots + X_{id} w_{d} + \alpha - y_{i} \right) \right] \\ \hat{\alpha} &= \frac{1}{n} \left[ \sum_{i=1}^{n} \left( y_{i} \right) \right] - \frac{1}{n} \left[ \sum_{i=1}^{n} \left( X_{i1} w_{1} + X_{i2} w_{2} + \dots + X_{id} w_{d} \right) \right] = \bar{y} - \bar{x} w = \bar{y} \end{split}$$

Set dJ/dw = 0 and solve for w:

$$\frac{dJ}{dw_i} = \left[ \sum_{j=1}^n 2X_{ji} (X_{i1}w_1 + X_{i2}w_2 + \dots + X_{id}w_d + \alpha - y_i) \right] + 2\lambda w_i$$

$$0 = \left[ \sum_{j=1}^n X_{ji} (X_{j1}w_1 + X_{j2}w_2 + \dots + X_{jd}w_d) + X_{ji}\alpha - X_{ji}y_j \right] + \lambda w_i$$

$$\sum_{j=1}^n X_{ji}y_j = n\bar{x}\alpha + \lambda w_i + \sum_{j=1}^n X_{ji} (X_{j1}w_1 + X_{j2}w_2 + \dots + X_{jd}w_d)$$

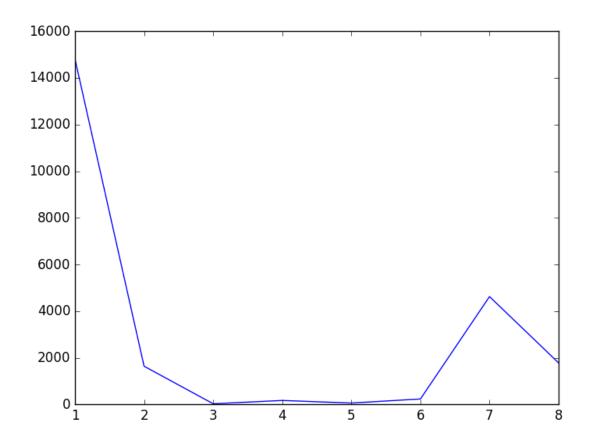
$$X_i^T y = \lambda w_i + X_i^T X_i w_i$$

$$X^T y = \lambda I \widehat{w} + X^T X \widehat{w} = (\lambda I + X^T X) \widehat{w}$$

$$\widehat{w} = (X^T X + \lambda I)^{-1} X^T y$$

b) The RSS on the validation set is 1.39e13. This is about twice what it was in HW3.

# c) Regression Coefficient plot:



The regression coefficients have a similar trend to those of HW3 (1, 7, and 8 are the most significant) but differ in that the magnitude of coefficients are smaller since the ridge regression model penalizes large weights.

```
2)
```

a) R(w0): 1.98837241413

b) mu0: [ 0.95257413 0.73105858 0.73105858 0.26894142]

c) w1: [-2. 0.94910188 -0.68363271]

d) R(w1): 1.72061709562

e) mu1: [ 0.89693957 0.54082713 0.56598026 0.15000896]

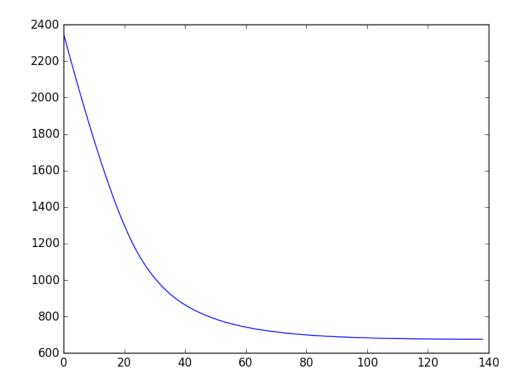
f) w2: [-1.69083609 1.91981257 -0.83738862]

g) R(w2): 1.85469978479

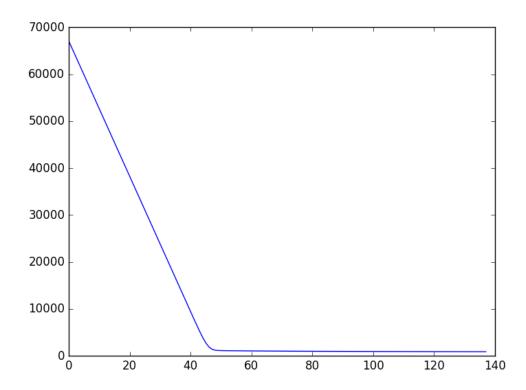
### Code used to generate values:

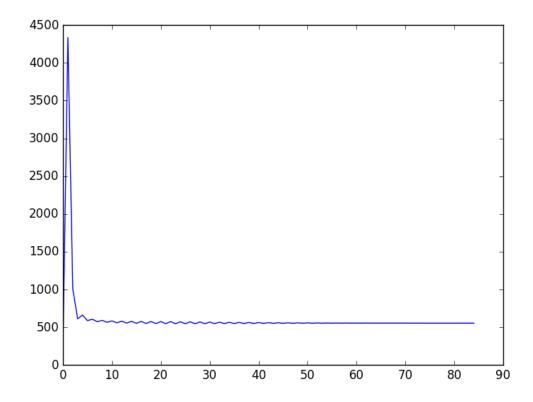
$$X = np.array([[0,3,1],[1,3,1],[0,1,1],[1,1,1]])$$
  
 $y = np.array([1,1,0,0])$   
 $w0 = np.array([-2,1,0])$   
 $e = 1$ 

```
def s(gamma):
  return 1/(1+np.exp(-gamma))
def R(w):
  total = 0
  for i in range(X.shape[0]):
     wXi = np.dot(w,X[i])
    total -= (y[i]*math.log(s(wXi)) + (1-y[i])*math.log(1 - s(wXi)))
  return total
print('R(w0): ',R(w0))
print('mu0: ', s(np.dot(w0,X.T)))
# iteration 1
total = 0
for i in range(X.shape[0]):
  total += (y[i] - s(np.dot(w0,X[i])))*X[i]
w1 = w0 + e*total
print('w1: ',w1)
print('R(w1): ',R(w1))
print('mu1: ',s(np.dot(w1,X.T)))
# iteration 2
total = 0
for i in range(X.shape[0]):
  total += (y[i] - s(np.dot(w1,X[i])))*X[i]
w2 = w1 + e*total
print('w2: ',w2)
print('R(w2): ',R(w2))
```

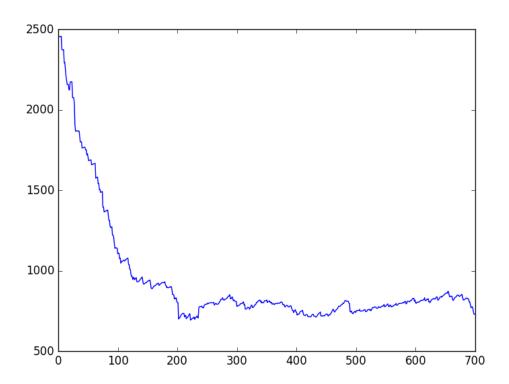


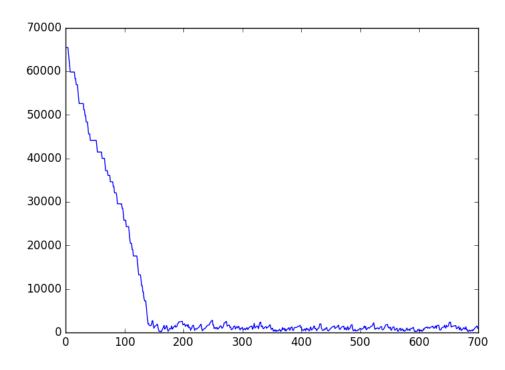
ii)



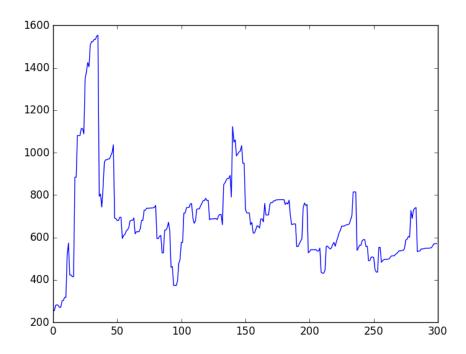


3.2) i)

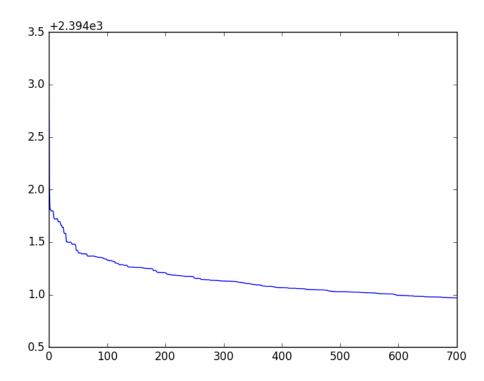




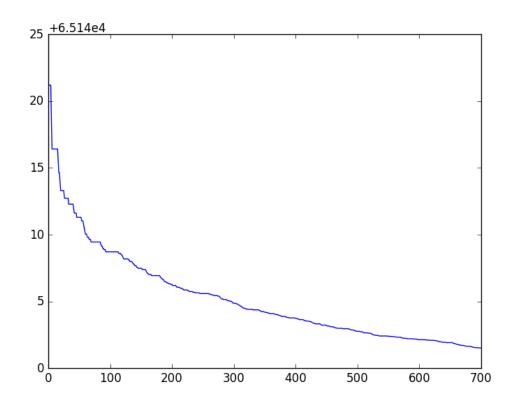
iii)

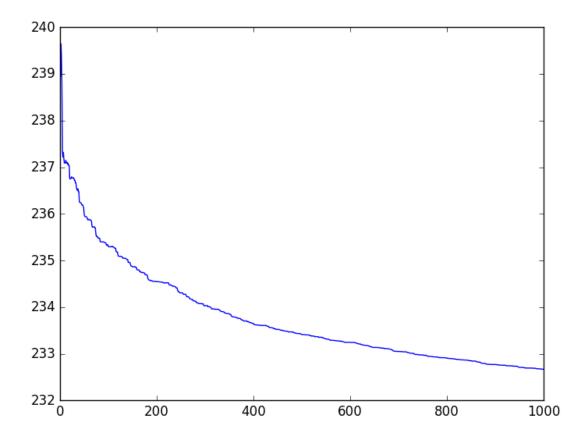


The plots differ from those of part (1) in that they do not strictly decrease, even though there is an overall decreasing trend. The sample chosen for each iteration does not necessarily decrease the risk.



ii)

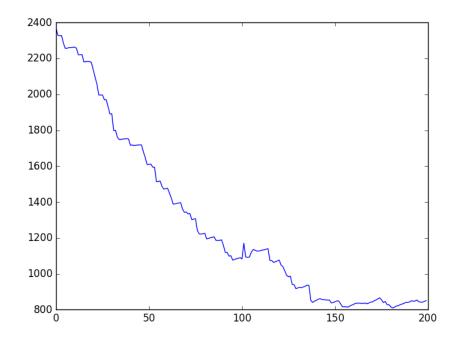




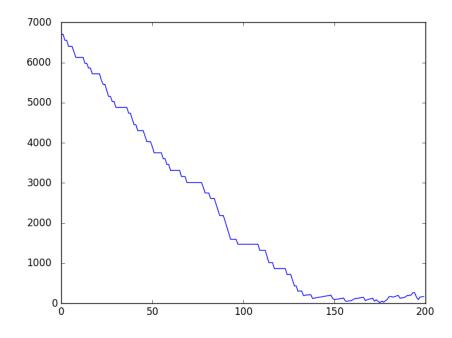
This strategy does not help improve accuracy in this case because it causes the model to converge faster, even when the risk can be further decreased with additional iterations. It does, however, help reduce volatility in the risk updates.

3.4) a) Used preprocessing method 1 since that seemed to give the highest accuracy. Optimal value of  $\rho$  was found to be 0.5.

#### Validation risk vs iterations:



## Training risk vs iterations:



b) Training accuracy: 0.701857134 Validation accuracy: 0.712959535

Here we don't see overfitting as the validation error is slightly lower than the training error. It is possible for the quadratic kernel to overfit the data, and if it did overfit the data then we should

increase  $\lambda$  to increase performance. This increases the penalty on large weight coefficients, reducing overfitting. For the linear kernel,  $\lambda$  could be decreased for increased performance by preventing underfitting.

3.5) My Kaggle score is 0.79482. I used batch gradient descent with 0 mean and unit variance for preprocessing. My learning rate was 0.0001.

4) a) 
$$\tanh(z) = 2s(2z) - 1$$

$$s(z) = \frac{1}{(1 + exp(-z))}$$

$$g(z) = \frac{(tanh(z) + 1)}{2} = \frac{(2s(2z) - 1 + 1)}{2} = s(2z) = \frac{1}{1 + exp(-2z)}$$

Multiply top and bottom by exp(z),:

$$g(z) = \frac{exp(z)}{\exp(z) + \exp(-z)} = \frac{exp(z)}{\exp(z) + \exp(-z)} - \frac{1}{2} + \frac{1}{2}$$

$$= \frac{2exp(z)}{2(\exp(z) + \exp(-z))} - \frac{\exp(z) + \exp(-z)}{2(\exp(z) + \exp(-z))} + \frac{1}{2}$$

$$= \frac{\exp(z) - \exp(-z)}{2(\exp(z) + \exp(-z))} + \frac{1}{2}$$

$$g(z) = \frac{\tanh(z) + 1}{2} = \frac{(2s(2z) - 1 + 1)}{2} = s(2z)$$

$$d(s(z))/dz = s(z)(1 - s(z))$$

$$g'(z) = \frac{d(s(2z))}{dz} = 2s(2z)(1 - s(2z)) = 2\left(\frac{\tanh(z)}{2} + \frac{1}{2}\right)\left(1 - \frac{\tanh(z)}{2} - \frac{1}{2}\right)$$

$$= (\tanh(z) + 1)\left(\frac{1}{2} - \frac{\tanh(z)}{2}\right) = \frac{1}{2}\tanh(z) - \frac{1}{2}\tanh(z) - \frac{1}{2}(\tanh(z))^2 + \frac{1}{2}$$

$$= \frac{1}{2} - \frac{1}{2}(\tanh(z))^2$$

c)

$$\nabla_{w}J(w) = \sum_{i=1}^{n} \frac{y_{i}X_{i}(\frac{1}{2}(1-\tanh^{2}(X_{i}w)))}{(\tanh(X_{i}w)+1)/2} + \frac{(1-y_{i})X_{i}(-\frac{1}{2}(1-\tanh^{2}(X_{i}w)))}{1-(\tanh(X_{i}w)+1)/2}$$

$$= \sum_{i=1}^{n} [y_{i}(1-\tanh(X_{i}w)) - (1-y_{i})(1+\tanh(X_{i}w))]X_{i}$$

$$= \sum_{i=1}^{n} [2y_{i}-1-\tanh(X_{i}w)]X_{i}$$

$$w \leftarrow w - \lambda\nabla_{w}J(w) = w + \lambda \sum_{i=1}^{n} [2y_{i}-1-\tanh(X_{i}w)]X_{i}$$

5) The linear SVM currently does not utilize the new feature well because the spike of spam messages occurs both before and after midnight. The new feature is stored as the number of seconds since the previous midnight, which means the additional spike in spam messages will consist of spam samples that have very high values (right before midnight) or very low values (right after midnight) for the new feature. This makes it impossible to linearly separate the low values from the high values. One way to improve the results would be to use a quadratic kernel and shift the feature vector by 12 hours so that the messages right after midnight have large negative values and the messages just before midnight have large positive values. Squaring this new feature with the quadratic kernel will result in a parabola in which the spam messages near midnight are all at the top, making it possible for the hyperplane to separate the spam messages from the ham messages at the bottom of the parabola.

#### CODE:

import scipy.io import numpy as np import random import matplotlib.pyplot as plt import math from sklearn import preprocessing import sklearn

```
# Problem 1
housing_data = scipy.io.loadmat("housing_dataset/housing_data.mat")
Xtrain= housing data['Xtrain']
Xvalidate= housing_data['Xvalidate']
Ytrain= housing data['Ytrain']
Yvalidate= housing_data['Yvalidate']
\# bias = np.ones((Xtrain.shape[0],1))
# Xtrain = np.hstack((Xtrain,bias))
\#Xtrain = Xtrain - np.tile(np.mean(Xtrain, axis = 0), (Xtrain.shape[0],1)) \# center x
Ytrain = np.ndarray.astype(Ytrain.ravel(), dtype = float)
Yvalidate = np.ndarray.astype(Yvalidate.ravel(), dtype = float)
lambdas = [pow(10,i)*1e-20 \text{ for } i \text{ in range}(40)]
# lambdas = [1e9]
best lambda = 0
best_error = float('inf')
indices = list(range(Xtrain.shape[0]))
random.shuffle(indices)
# find best lambda
errors = []
for 1 in lambdas:
  chosen x = []
  chosen_y = []
  for i in range(10):
     subset = indices[i*Xtrain.shape[0]//10:(i+1)*Xtrain.shape[0]//10]
     subsetX = []
     subsetY = []
    for j in subset:
       subsetX.append(Xtrain[j])
       subsetY.append(Ytrain[j])
     chosen_x.append(np.array(subsetX))
     chosen y.append(np.array(subsetY))
  loss = 0
  for i in range(10):
     x_validate = chosen_x[i]
     y_validate = chosen_y[i]
     x = np.vstack(chosen_x[:i] + chosen_x[i+1:])
     x = x - np.tile(np.mean(x, axis = 0), (x.shape[0],1))
```

```
y = np.hstack(chosen_y[:i] + chosen_y[i+1:])
     w = np.linalg.solve(np.dot(x.T,x)+l*np.identity(x.shape[1]),np.dot(x.T,y))
     \# w = np.dot(np.linalg.inv(np.dot(x.T,x) + l*np.identity(x.shape[1])),np.dot(x.T,y))
     a = np.mean(y)
     predictions = np.dot(x validate, w) + a*np.ones(x validate.shape[0])
     loss += np.dot((predictions - y_validate)).T,(predictions - y_validate)) + 1*np.dot(w.T,w)
  errors.append(loss/10)
  print(l, loss/10)
  if loss/10 < best_error:
     best error = loss/10
     best lambda = 1
plt.semilogx(lambdas,errors)
plt.show()
# part ii and part iii
1 = 1e6
Xtrain = Xtrain - np.tile(np.mean(Xtrain, axis = 0), (Xtrain.shape[0], 1)) # center x
x = Xtrain
y = Ytrain
w = np.linalg.solve(np.dot(x.T,x)+l*np.identity(x.shape[1]),np.dot(x.T,y))
a = np.mean(y)
predictions = np.dot(Xvalidate, w) + a*np.ones(Xvalidate.shape[0])
RSS = np.dot((predictions - Yvalidate).T,(predictions-Yvalidate))# + 1*np.dot(w.T,w)
plt.figure()
plt.plot(range(1,9), abs(w))
plt.show()
# problem 2
X = \text{np.array}([[0,3,1],[1,3,1],[0,1,1],[1,1,1]])
y = np.array([1,1,0,0])
w0 = np.array([-2,1,0])
e = 1
def s(gamma):
  return 1/(1+np.exp(-gamma))
def R(w):
  total = 0
  for i in range(X.shape[0]):
     wXi = np.dot(w,X[i])
     total = (y[i]*math.log(s(wXi))+(1-y[i])*math.log(1 - s(wXi)))
  return total
```

```
print('R(w0): ',R(w0))
print('mu0: ', s(np.dot(w0,X.T)))
# iteration 1
total = 0
for i in range(X.shape[0]):
  total += (y[i] - s(np.dot(w0,X[i])))*X[i]
w1 = w0 + e*total
print('w1: ',w1)
print('R(w1): ',R(w1))
print('mu1: ',s(np.dot(w1,X.T)))
# iteration 2
total = 0
for i in range(X.shape[0]):
  total += (y[i] - s(np.dot(w1,X[i])))*X[i]
w2 = w1 + e*total
print('w2: ',w2)
print('R(w2): ',R(w2))
# problem 3
spam_data = scipy.io.loadmat("spam_dataset/spam_data.mat")
X = \text{spam data}[\text{'training data'}]
y = spam_data['training_labels'].ravel()
test_data= spam_data['test_data']
indices = list(range(y.shape[0]))
random.shuffle(indices)
validation_x = []
validation_y = []
training_x = []
training_y = []
for i in indices[:y.shape[0]//3]:
  validation_x.append(X[i])
  validation_y.append(y[i])
for i in indices[y.shape[0]//3:]:
  training_x.append(X[i])
  training_y.append(y[i])
X = np.array(training_x)
y = np.array(training_y)
validation_x = np.array(validation_x)
validation_y = np.array(validation_y)
```

```
def s(gamma):
  return 1/(1+np.exp(-gamma))
def R(w,X):
  total = 0
  for i in range(X.shape[0]):
     sigmoid_wXi = s(np.dot(w,X[i]))
     total = (y[i]*math.log(sigmoid_wXi)+(1-y[i])*math.log(max((1 - sigmoid_wXi),1)))
  return total
# Problem 3.1
def batch(w, X, e = 0.0001, processing = 0, tolerance = 0.01, max iter = 200):
  if processing == 1:
     X = preprocessing.scale(X) # part (i)
  elif processing == 2:
     X = \text{np.vectorize}(\text{lambda Xij: math.log}(\text{Xij} + 0.1))(X) \# \text{part (ii)}
  elif processing == 3:
     X = sklearn.preprocessing.binarize(X,0.01) # part (iii)
  risk list = [R(w,X)]
  total = 0
  for i in range(X.shape[0]):
     total += (y[i] - s(np.dot(w,X[i])))*X[i]
  w = w + e*total
  risk list.append(R(w,X))
  while abs(risk list[-1] - risk list[-2]) > tolerance and len(risk list) < max iter:
     total = 0
     for i in range(X.shape[0]):
       total += (y[i] - s(np.dot(w,X[i])))*X[i]
     w = w + e*total
     risk_list.append(R(w,X))
  return w, risk_list
w0 = np.ones(X.shape[1])
w,risks = batch(w0, X, processing = 1)
plt.figure()
plt.plot(range(len(risks)),risks)
accuracy = np.count_nonzero(sklearn.preprocessing.binarize(s(np.dot(validation_x,w)),0.5) ==
validation y)/validation y.shape[0]
print(accuracy)
w, risks = batch(w0, X, e = 1e-5, tolerance = 1, processing = 2)
plt.figure()
```

```
plt.plot(range(len(risks)),risks)
accuracy = np.count_nonzero(sklearn.preprocessing.binarize(s(np.dot(validation_x,w)),0.5) ==
validation y)/validation y.shape[0]
print(accuracy)
w, risks = batch(w0, X, e = 1e-2, processing = 3)
plt.figure()
plt.plot(range(len(risks)),risks)
accuracy = np.count_nonzero(sklearn.preprocessing.binarize(s(np.dot(validation_x,w)),0.5) ==
validation_y)/validation_y.shape[0]
print(accuracy)
#Problem 3.2
def stochastic(w, X, e = 1e-1, processing = 0, max_iter = 1000):
  if processing == 1:
     X = preprocessing.scale(X) # part (i)
  elif processing == 2:
     X = \text{np.vectorize}(\text{lambda Xij: math.log}(\text{Xij} + 0.1))(X) \# \text{part (ii)}
  elif processing == 3:
     X = \text{sklearn.preprocessing.binarize}(X, 0.01) \# \text{ part (iii)}
  risk_list = [R(w,X)]
  idx = random.randint(0,(y.shape[0]-1))
  w = w + e*np.dot((y[idx] - s(np.dot(X[idx],w))),X[idx])
  risk_list.append(R(w,X))
  while len(risk list) < max iter:
     idx = random.randint(0, y.shape[0]-1)
     w = w + e*np.dot((y[idx] - s(np.dot(X[idx],w))),X[idx])
     risk list.append(R(w,X))
  return w, risk_list
w0 = np.ones(X.shape[1])
w, risks = stochastic(w0, X, processing = 1, max_iter = 700)
plt.figure()
plt.plot(range(len(risks)),risks)
accuracy = np.count nonzero(sklearn.preprocessing.binarize(s(np.dot(validation x,w)),0.5) ==
validation y)/validation y.shape[0]
print(accuracy)
w,risks = stochastic(w0, X, e = 1e-2, processing = 2, max_iter = 700)
plt.figure()
plt.plot(range(len(risks)),risks)
accuracy = np.count nonzero(sklearn.preprocessing.binarize(s(np.dot(validation x,w)),0.5) ==
validation y)/validation y.shape[0]
print(accuracy)
```

```
w,risks = stochastic(w0, X, e = 1, processing = 3, max iter = 300)
plt.figure()
plt.plot(range(len(risks)),risks)
accuracy = np.count_nonzero(sklearn.preprocessing.binarize(s(np.dot(validation_x,w)),0.5) ==
validation_y)/validation_y.shape[0]
print(accuracy)
# Problem 3.3
def decreasing_e(w, X, e = 1e-1, processing = 0, max_iter = 1000):
  if processing == 1:
     X = preprocessing.scale(X) # part (i)
  elif processing == 2:
     X = \text{np.vectorize}(\text{lambda Xij: math.log}(\text{Xij} + 0.1))(X) \# \text{part (ii)}
  elif processing == 3:
     X = sklearn.preprocessing.binarize(X,0.01) # part (iii)
  risk list = [R(w,X)]
  idx = random.randint(0,(y.shape[0]-1))
  w = w + e*np.dot((y[idx] - s(np.dot(X[idx],w))),X[idx])
  risk list.append(R(w,X))
  while len(risk_list) < max_iter:
     idx = random.randint(0, y.shape[0]-1)
     w = w + e*np.dot((y[idx] - s(np.dot(X[idx],w))),X[idx])/len(risk_list)
     risk_list.append(R(w,X))
  return w, risk list
w,risks = decreasing e(w0, X, e = 1e-3, processing = 1, max iter = 700)
plt.figure()
plt.plot(range(len(risks)),risks)
accuracy = np.count_nonzero(sklearn.preprocessing.binarize(s(np.dot(validation_x,w)),0.5) ==
validation_y)/validation_y.shape[0]
print(accuracy)
w, risks = decreasing_e(w0, X, e = 1e-4, processing = 2, max_iter = 700)
plt.figure()
plt.plot(range(len(risks)),risks)
accuracy = np.count nonzero(sklearn.preprocessing.binarize(s(np.dot(validation x,w)),0.5) ==
validation_y)/validation_y.shape[0]
print(accuracy)
w,risks = decreasing_e(w0, X, e = 0.1, processing = 3)
plt.figure()
plt.plot(range(len(risks)),risks)
```

```
accuracy = np.count_nonzero(sklearn.preprocessing.binarize(s(np.dot(validation_x,w)),0.5) ==
validation_y)/validation_y.shape[0]
print(accuracy)
# Problem 3.4 (a)
def k(x,z,rho):
  return np.dot((np.dot(x.T,z)+rho).T, np.dot(x.T,z)+rho)
def kernal_R(a,x):
  gamma = np.dot(np.dot(a,x),x.T)
  r_i = y * np.log(s(gamma)) + (1-y) * np.log(1 - s(gamma))
  return -np.sum(r_i)
def stochastic_kernel(a, X, l=1e-3, e=0.1, rho = 0.5, tol = 1e-8, max_iter=200):
  K=np.zeros((X.shape[0],X.shape[0]))
  z1 = X[random.randint(0,y.shape[0]-1)]
  for i in range(K.shape[0]):
    for j in range(K.shape[0]):
       K[i][j] = k(X[i],z1,rho)
  r list = [kernal R(a, X)]
  delta = float('inf')
  while(len(r_list) < max_iter):</pre>
     idx = random.randint(0, y.shape[0]-1)
     a = a-e^*l^*a[idx]+e^*(y[idx]^*s(np.dot(K,a)[idx]))
    r = kernal R(a, X)
     delta = r list[-1] - r
     r_list.append(r)
  return a, r list, K
a = np.zeros(y.shape[0])
X = preprocessing.scale(X)
a_{\text{final,r_list,K}} = \text{stochastic_kernel}(a,X)
accuracy = np.count_nonzero(sklearn.preprocessing.binarize(s(np.dot(K,a_final)),0.5) ==
y)/y.shape[0]
# Find best rho
# for rho in [i/10 for i in range(1,10)]:
    a final,r list,K = stochastic kernel(a,X, rho = rho)
    accuracy = np.count_nonzero(sklearn.preprocessing.binarize(s(np.dot(K,a_final)),0.5) ==
y)/y.shape[0]
# Problem 3.4 (b)
```

```
def k(x,z,rho):
  return np.dot(x.T,z)+rho
def kernal_R(a,x):
  gamma = np.dot(np.dot(a,x),x.T)
  r_i = y * np.log(s(gamma)) + (1-y) * np.log(1 - s(gamma))
  return -np.sum(r_i)
def stochastic_kernel(a, X, l=1e-3, e=0.1, rho = 0.5, tol = 1e-8, max_iter=200):
  K=np.zeros((X.shape[0],X.shape[0]))
  z1 = X[random.randint(0,y.shape[0]-1)]
  for i in range(K.shape[0]):
     for j in range(K.shape[0]):
       K[i][j] = k(X[i],z1,rho)
  r_list = [kernal_R(a, X)]
  delta = float('inf')
  while(len(r list) < max iter):
     idx = random.randint(0, y.shape[0]-1)
     a = a-e^*l^*a[idx]+e^*(y[idx]^*s(np.dot(K,a)[idx]))
     r = kernal_R(a, X)
     delta = r_list[-1] - r
     r list.append(r)
  return a, r list, K
a = np.zeros(y.shape[0])
X = preprocessing.scale(X)
a final, r list, K = stochastic kernel(a, X)
accuracy = np.count nonzero(sklearn.preprocessing.binarize(s(np.dot(K,a final)),0.5) ==
y)/y.shape[0]
# Find best rho
# for rho in [i/10 for i in range(1,10)]:
    a_final,r_list,K = stochastic_kernel(a,X, rho = rho)
    accuracy = np.count_nonzero(sklearn.preprocessing.binarize(s(np.dot(K,a_final)),0.5) ==
y)/y.shape[0]
# Problem 3.5
def batch(w, X, e = 0.0001, processing = 0, tolerance = 0.01, max iter = 200):
  if processing == 1:
     X = preprocessing.scale(X) # part (i)
  elif processing == 2:
     X = \text{np.vectorize}(\text{lambda Xij: math.log}(\text{Xij} + 0.1))(X) \# \text{part (ii)}
  elif processing == 3:
     X = \text{sklearn.preprocessing.binarize}(X, 0.01) \# \text{ part (iii)}
```

```
risk_list = [R(w,X)]
  total = 0
  for i in range(X.shape[0]):
     total += (y[i] - s(np.dot(w,X[i])))*X[i]
  w = w + e*total
  risk_list.append(R(w,X))
  while abs(risk_list[-1] - risk_list[-2]) > tolerance and len(risk_list) < max_iter:
     total = 0
     for i in range(X.shape[0]):
       total += (y[i] - s(np.dot(w,X[i])))*X[i]
     w = w + e*total
     risk_list.append(R(w,X))
  return w, risk_list
w0 = np.ones(X.shape[1])
w,risks = batch(w0, X, processing = 1)
plt.figure()
plt.plot(range(len(risks)),risks)
predictions = sklearn.preprocessing.binarize(s(np.dot(test_data,w)),0.5)
numbers = np.arange(predictions.ravel().shape[0]) + 1
test_predict = np.vstack((numbers,predictions))
np.savetxt("spam.csv", test_predict.transpose(), delimiter=",",fmt = '%u')
```