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# hw 2 Spam Classification

## Report

### • Program Overview

Different from the hw1, I modularized several parts of my program this time. All the functions are show as follows:

- 1. sigmoid
- 2. cross\_entropy
- 3. read data
- 4. logistic\_regression
- 5. gen\_model
- 6. read model
- 7. read test
- 8. gen\_ans

#### • sigmoid (Fig. 4)

It's a normal sigmoid function (support numpy array operation).

#### cross\_entropy (Fig. 5)

It will compute the cross-entropy for each weight. Since the predicted value may very close to either 0 or 1, which causing operation error to the numpy log function, I add a small value (1e-3) to the predicted value which is close to either 0 or 1 to avoid such runtime error. (Fig 2)

#### read\_data

It's a normal function to read the training data into the format of numpy array. (the data format is same with the hw1  $\rightarrow$  row: observation, col: variables)

#### logistic\_regression (Fig. 6)

It's similar with the linear regression version. The only differences are list below:

- 1. It replaces the results of inner product of weights and data with the same results but under the sigmoid function manipulation.
- 2. It changes the coefficients of computed gradient value of both weights and bias
- 3. It changes the way to calculate loss function (root-mean-square error to cross-entropy)
  Beyond these difference, I thought that the logistic one is almost the same with linear one.

#### gen\_model

I used this function to write back the weights and bias to the model file via json format.

#### · read model

It's a basic and trivial function which load the generated model.

#### read\_test

It's almost the same with the function "read\_data" which read the testing data.

#### • gen\_ans (Fig. 7)

Like previous functions, this function is similar with the part of the codes in hw1. It calculates the predicted value of testing data and fills the corresponding answer to the array according to their probability.

Finally, I ran the different functions according to the length of the system parameters and generated the "prediction.csv".

## Learning Rate

Different from the hw1, in this homework, only the adagrad and adamgrad can converge in specific epochs. The results of using adagrad is similar with using adamgrad.

#### Discussion

In this assignment, I have implemented 3 different methods:

- 1. Master → Logistic Regression (#iter: 8000, Adagrad, learning rate: 1 → Accuracy: **0.94333**)
- 2. Method2 → Multi-Logistic ("Neural Network", 4 layers, 4 neurons → Accuracy: **0.94667**) Fig. 8
- 3. Method3  $\rightarrow$  Probabilistic Generative Model (Accuracy: **0.94333**)  $\rightarrow$  Fig. 9
- 4. Method4 → Linear Regression (#iter: 8000, Adagrad, learning rate: 1 → Accuracy: **0.93667**)

I had tried some different combinations of the parameters such like #iteration, learning rate, feature extraction, etc. The best one's accuracy in the public leaderboard on the Kaggle is 0.95000, but I forgot the setting of the parameters... The other three methods I implemented are multi-logistic, probabilistic generative and linear regression. Take multi-logistic for further discussion, I setup 2 parameters "layer" and "neuron" to determine the structure of my neuron network. Then I pass the data to first layer, and take the output to be the next layer's input and so on. In the last layer, I output the final result to judge whether the mail is spam or not. The whole process is something like continuously doing the simple logistic regression, the only difference is to use the previous layer's output to be the training data for the next layer. Also, the data format should be check carefully; otherwise, the program will crash. For example, the format of the output model should be designed carefully such that it can be easily loaded later (I used nested dictionary). What's more, the keys of the loaded dictionary are all encoded in "unicode", so it need the preprocessing. Additionally, I did some experiments which are show in several line charts on the next page. After doing these experiments, I was attracted by the different feature extraction methods' effect; in other words, the difference between feature scale and combinations could result in different accuracy. Since this difference is bigger than I expected, I decided to try other possible combinations and plotted in the line chart (I only try 3 different methods at first). The reason why I came up with this idea is that I noticed that the last several features represent the rare characters which may not appear in the normal email content. (Note that all the experiments are under normal logistic version)

## Figures

#### **Different Epochs on Optimize Methods**

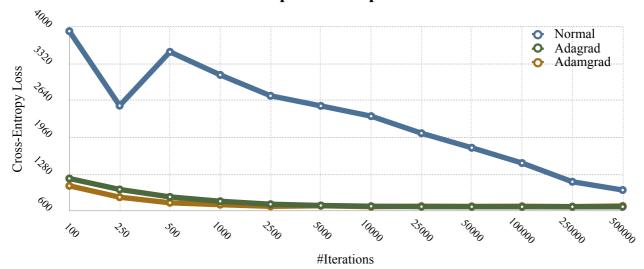


Figure 1.

Experiment results of different epochs on different optimize methods

#### **Different Feature Extractions (Adamgrad, learning rate = 0.001)**

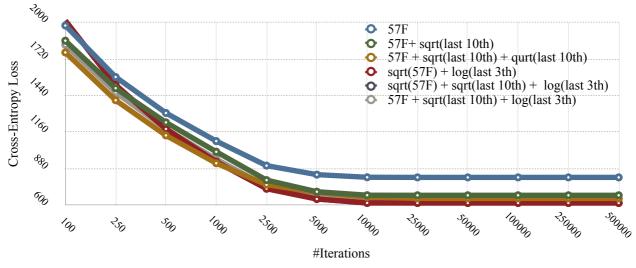


Figure 2.

Experiment results of different feature extraction method (F = feature, Adamgrad, learning rate = 0.001)

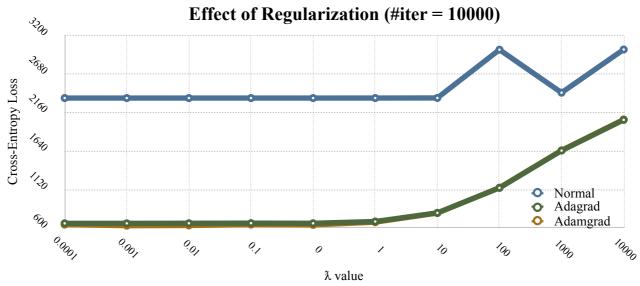


Figure 3.

Experiment results of the effect of regularization (#iter = 10000)

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#### Codes

```
def sigmoid(z):
return 1 / (1 + np.exp(-z))
```

Figure. 4

Sigmoid function implemented by numpy

Figure. 5

Compute the cross-entropy for each weight (they will be summed as the value of loss function)

```
logistic_regression(X_TRAIN, '
            W, b = np.zeros(X_TRAIN.shape[1]), 0
             SUM_SQDW, SUM_SQDB = np.zeros(X_TRAIN.shape[1])+1, 0
             norm, adag, adam = 0.00000001, 0.1, 0.01 # adam-default = 0.001
             beta1, beta2 = 0.9, 0.999
            Wmt, Wvt = 0, 0
Bmt, Bvt = 0, 0
            epoch, Lambda, t, eps = 10000, 0, 0, 1e-8
for i in range(epoch):
                  fwb = sigmoid(np.dot(X_TRAIN, W) + b)
ERR = Y_TRAIN - fwb
DW = -1 * np.dot(X_TRAIN.T, ERR)
DB = -1 * np.sum(ERR)
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                  DW += Lambda * 2 * W
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                  # SUM_SQDB += np.square(DB)
                  # W -= adag / np.sqrt(SUM_SQDW) * DW # / X_TRAIN.shape[0]
# b -= adag / np.sqrt(SUM_SQDB) * DB # / X_TRAIN.shape[0]
                  Wmt = beta1 * Wmt + (1-beta1) * DW
64
                  Wvt = beta2 * Wvt + (1-beta2) * np.square(DW)
                  Wmthat = Wmt / (1-np.power(beta1, t))
Wvthat = Wvt / (1-np.power(beta2, t))
Bmt = beta1 * Bmt + (1-beta1) * DB
                  Bvt = beta2 * Bvt + (1-beta2) * np.square(DB)
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                  Bmthat = Bmt / (1-np.power(beta1, t))
Bvthat = Bvt / (1-np.power(beta2, t))
                         (adam*Wmthat) / (np.sqrt(Wvthat) + eps)
                  b
                         (adam*Bmthat) / (np.sqrt(Bvthat) +
             return W, b
```

Figure. 6

Logistic Regression (normal, adagrad, adamgrad)

```
gen_ans(anspath, X_TEST, W,
                fwb_TEST = sigmoid(np.dot(X_TEST, W) + b)
Y_TEST = []
for i in fwb_TEST:
                       Y_TEST.append(0) if np.less_equal(i, 0.5) else Y_TEST.append(1)
                with open(anspath, 'wb') as file:
    writer = csv.writer(file, delimiter=',')
    writer.writerow(('id', 'label'))
    for i in range(len(Y_TEST)): writer.writerow((i+1, Y_TEST[i]))
100
```

Figure, 7

Generate the predicted results according to the probability (logistic)

```
multi_logistic(neuron, layer, X, Y):
W_dic, b_dic = {}, {}
X_NEXT, WT, bT = backward_prop(X, Y, neuron)
W_dic, b_dic = update_neuron_info(W_dic, b_dic, WT, bT, neuron, 0)
                W_dic, b_dic = dpdate_intro(w_dic, b_dic, wi, br, hearon, e)
for i in range(1, layer):
    X_NEXT, WT, bT = backward_prop(X_NEXT, Y, neuron)
    W_dic, b_dic = update_neuron_info(W_dic, b_dic, WT, bT, neuron, i)
W_dic, b_dic = classification(X_NEXT, Y, W_dic, b_dic, neuron, layer)
return W_dic, b_dic
         def logistic_regression(X_TRAIN, Y_TRAIN):
         def backward_prop(X_TRAIN, Y_TRAIN, neuron):
    WFT, bFT, = [], []
    X_NEXT = [[] for i in range(neuron)]
    for i in range(neuron):
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                      tW, tb = logistic_regression(X_TRAIN, Y_TRAIN)
                      WFT.append(tW)
                       bFT.append(tb)
                      fwb = sigmoid(np.dot(X_TRAIN, tW) + tb)
for j in range(fwb.shape[0]):
    X_NEXT[i].append(fwb[j]) # shape=(neuron, 4001)
                X_NEXT = np.array(X_NEXT).T # shape=(4001, neuron)
return X_NEXT, WFT, bFT
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                | wdic[i], bdic[i] = W[i], b[i]
W_dic[layer], b_dic[layer] = wdic, bdic
return W_dic, b_dic
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         def classification(X_TRAIN, Y_TRAIN, W_dic, b_dic, neuron, layer):
    X_TRANS = [[] for i in range(neuron)]
                for i in range(neuron):
                W_dic[layer], b_dic[layer] = WC, bC
                return W_dic, b_dic
```

Figure. 8

Multi-Logistic Model (so called "Neural Network", normal, adagrad, adamgrad, #iter:7500)

```
f gen_mean(X, Y):
    X_0, X_1 = [], []
    for i in range(Y.shape[0]):
          X_0.append(X[i]) if Y[i] == 0 else X_1.append(X[i])
          X_0, X_1 = np.array(X_0).T, np.array(X_1).T
          u_0, u_1 = [], []
    col = X.shape[1]
    for i in range(col):
          u_0.append(col):
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                                                                                                      cot = X.snapel1)
for i in range(col):
    u_0.append(np.mean(X_0[i][:]))
    u_1.append(np.mean(X_1[i][:]))
return np.array(u_0), np.array(u_1), X_0.shape[1], X_1.shape[1]
                                                       def gen_cov(X):
                                                                                                                                              urn np.cov(X, rowvar=False)
                                                  def gen_model(modelpath, u_0, u_1, cov, N_0, N_1):
    inv_cov = np.linalg.inv(cov)
    W = np.dot((u_0 - u_1).T, inv_cov).T
    b = -0.5 * np.dot(np.dot(u_0.T, inv_cov), u_0) + \
        0.5 * np.dot(np.dot(u_1.T, inv_cov), u_1) + \
        np.log(N_0 / N_1)
    with open(modelpath, 'wb') as file:
        json.dump({'b': b, 'W': list(W)}, file)
```

Figure. 9

Probabilistic Generative Model