

# Chem277B: Machine Learning Algorithms

## Homework assignment #5: Regression

```
In [4]: import numpy as np
import pandas as pd
import math
import matplotlib.pyplot as plt
import torch
import torch.nn as nn
import torch.optim as optim
from sklearn.model_selection import train_test_split, KFold
```

### 1. Baye's Theorem.

(a) From the given data, the categories of testing results and their probabilities within proportion have been summarized in the table below:

Category	Probability within Proportion	Kidney Disease Positive	Marker	Proportion
P[+ M]	0.95	+	+	0.95 * 0.01
P[- M]	(1-0.95)	-	+	(1-0.95) * 0.01
P[+ not M]	(1-0.95)	+	-	(1-0.95) * 0.99
P[- not M]	0.95	-	-	0.95 * 0.99

Hence the quantities for the questions are:

(a1)  $P[-|M] = (1-0.95) = 0.05$  within its proportion, the absolute probability is  $0.05 * 0.01 = 0.0005$

(a2)  $P[+|not M] = (1-0.95) = 0.05$  within its proportion, the absolute probability is  $0.05 * 0.99 = 0.0495$

(a3)  $P[not M] = (1-0.95) * 0.99 + 0.95 * 0.99 = 0.99$ , or  $1 - 0.01 = 0.99$

(b) Using the Baye's Theorem, we try to differentiate between positive marker + positive test and positive marker + negative test. Hence the calculation is defined as:

$$P[M|+] = \frac{P[+|M] * P[M]}{P[+|M] * P[M] + P[+|not M] * P[not M]}$$

$$= \frac{0.95 \times 0.01}{(0.95 \times 0.01) + (0.05 \times 0.99)} = 0.161$$

Hence the chance of testing positive and actually have the marker is 16.1%. It warrants additional testing to confirm.

(c) When  $P[M] = 0.10$ , the categories and probabilities become the following:

Category	Probability within Proportion	Kidney Disease Positive	Marker	Proportion
$P[+ M]$	0.95	+	+	$0.95 * 0.10$
$P[- M]$	$(1-0.95)$	-	+	$(1-0.95) * 0.10$
$P[+ not M]$	$(1-0.95)$	+	-	$(1-0.95) * 0.90$
$P[- not M]$	0.95	-	-	$0.95 * 0.90$

Hence with the new frequency, the individual who test positive actually has the marker is:

$$P[M|+] = \frac{P[+|M]' * P[M]'}{P[+|M]' * P[M]' + P[+|not M]' * P[not M]'}$$

$$= \frac{0.95 \times 0.1}{(0.95 \times 0.1) + (0.05 \times 0.9)} = 0.679$$

## 2. Gaussian Naive Bayes.

(a) The finished codes are shown below.

I chose Gaussian distribution because it's a widely used normal probability distributions in statistics, data analysis and visualization.

The Gaussian distribution has a bell curve with mean and standard deviation, hence suitable for modeling many real-world phenomena.

With the finished function, I calculated that a wine from cultivar 1 has a 23.24% probability of containinh 13% Alcohol.

```
In [5]: class NaiveBayesClassifier():
        def __init__(self):
            self.type_indices={}      # store the indices of wines that belong to each cultivar as a boolean array of length
            self.type_stats={}        # store the mean and std of each cultivar
            self.ndata = 0
```

```

        self.trained=False

    @staticmethod
    def gaussian(x,mean,std):
        exponent = -(x - mean)**2 / (2 * std**2)

        return (np.exp(exponent) / (np.sqrt(2 * np.pi) * std))

    @staticmethod
    def calculate_statistics(x_values):
        # Returns a list with length of input features. Each element is a tuple, with the input feature's average and s
        n_feats=x_values.shape[1]
        return [(np.average(x_values[:,n]),np.std(x_values[:,n])) for n in range(n_feats)]

    @staticmethod
    def calculate_prob(x_input,stats):
        """Calculate the probability that the input features belong to a specific class(P(X|C)), defined by the statist
        x_input: np.array shape(nfeatures)
        stats: list of tuple [(mean1,std1),(means2,std2),...]
        """

        init_prob = 1
        for i in range(len(x_input)):
            mean, std = stats[i]
            init_prob *= NaiveBayesClassifier.gaussian(x_input[i], mean, std)
        return init_prob

    def fit(self,xs,ys):
        # Train the classifier by calculating the statistics of different features in each class
        self.ndata = len(ys)
        for y in set(ys):
            type_filter= (ys==y)
            self.type_indices[y]=type_filter
            self.type_stats[y]=self.calculate_statistics(xs[type_filter])
        self.trained=True

    def predict(self,xs):
        # Do the prediction by outputing the class that has highest probability
        if len(xs.shape)>1:
            print("Only accepts one sample at a time!")
        if self.trained:
            guess=None
            max_prob=0
            # P(C|X) = P(X|C)*P(C) / sum_i(P(X|C_i)*P(C_i)) (deniminator for normalization only, can be ignored)
            for y_type in self.type_stats:
                pre = sum(self.type_indices[y_type]) / self.ndata
                prob= self.calculate_prob(xs, self.type_stats[y_type]) * pre
                if prob>max_prob:
                    max_prob=prob
                    guess=y_type
            return guess

```

```

    else:
        print("Please train the classifier first!")

def calculate_accuracy(model,xs,ys):
    y_pred=np.zeros_like(ys)
    for idx,x in enumerate(xs):
        y_pred[idx]=model.predict(x)
    return np.sum(ys==y_pred)/len(ys)

```

```

In [6]: # Import wines.csv
wines = pd.read_csv('wines.csv')
wines.head()

```

```

Out[6]:

```

	Alcohol %	Malic Acid	Ash	Alkalinity	Mg	Phenols	Flavanoids	Phenols.1	Proantho- cyanins	Color intensity	Hue	OD280 315	Proline	Start assignment	ranking
0	14.23	1.71	2.43	15.6	127	2.8	3.06	0.28	2.29	5.64	1.04	3.92	1065	1	1
1	13.24	2.59	2.87	21.0	118	2.8	2.69	0.39	1.82	4.32	1.04	2.93	735	1	1
2	14.83	1.64	2.17	14.0	97	2.8	2.98	0.29	1.98	5.20	1.08	2.85	1045	1	1
3	14.12	1.48	2.32	16.8	95	2.2	2.43	0.26	1.57	5.00	1.17	2.82	1280	1	1
4	13.75	1.73	2.41	16.0	89	2.6	2.76	0.29	1.81	5.60	1.15	2.90	1320	1	1

```

In [7]: # Define the instance from the Naive Bayes Classifier
nbc = NaiveBayesClassifier()

# Fit the Naive Bayes Classifier
nbc.fit(wines.loc[:, 'Alcohol %':'Proline'].values, wines.loc[:, 'ranking'].values)

# First get the stats for cultivar 1
type_stats = nbc.type_stats[1]

# Then calculate the probability of alcohol% = 13
probability = nbc.gaussian(13, type_stats[0][0], type_stats[0][1])

print(f"A wine from cultivar 1 has a {round(probability*100, 2)}% probability of containinh 13% Alcohol")

```

A wine from cultivar 1 has a 23.24% probability of containinh 13% Alcohol

(b) After 3-fold training, I can achieve close to 100% accuracy in very short term. The Naive Baye's method performs much better and much faster than the simulated annealing method.

```

In [8]: # First normalize the wines dataframe
wines_norm = wines.loc[:, 'Alcohol %':'Proline']
wines_norm = (wines_norm - np.mean(wines_norm, axis=0)) / np.std(wines_norm, axis=0)

```

```
wines_norm = wines_norm.merge(wines[['Start assignment', 'ranking']], left_index=True, right_index=True)
wines_norm.head()
```

Out[8]:

	Alcohol %	Malic Acid	Ash	Alkalinity	Mg	Phenols	Flavanoids	Phenols.1	Proanthocyanins	Color intensity	Hue	OD280 315	Proline
0	1.518613	-0.562250	0.232053	-1.169593	1.913905	0.808997	1.034819	-0.659563	1.224884	0.251717	0.362177	1.847920	1.013005
1	0.295700	0.227694	1.840403	0.451946	1.281985	0.808997	0.663351	0.226796	0.401404	-0.319276	0.362177	0.449601	-0.037874
2	2.259772	-0.625086	-0.718336	-1.650049	-0.192495	0.808997	0.954502	-0.578985	0.681738	0.061386	0.537671	0.336606	0.949315
3	1.382733	-0.768712	-0.170035	-0.809251	-0.332922	-0.152402	0.402320	-0.820719	-0.036617	-0.025128	0.932531	0.294232	1.697675
4	0.925685	-0.544297	0.158946	-1.049479	-0.754202	0.488531	0.733629	-0.578985	0.383884	0.234414	0.844785	0.407228	1.825055

In [9]:

```
# Divide the normalized wines data into 3-fold training and testing groups
# and use 2/3 training and 1/3 testing for the three divisions
kf = KFold(n_splits=3, shuffle=True)
xs = wines.loc[:, 'Alcohol %': 'Proline'].values
ys = wines.loc[:, 'ranking'].values
nbc = NaiveBayesClassifier()
accuracy = []

for train_index, test_index in kf.split(xs):
    x_train, x_test = xs[train_index], xs[test_index]
    y_train, y_test = ys[train_index], ys[test_index]

    # train the classifier
    nbc.fit(x_train, y_train)
    accuracy.append(calculate_accuracy(nbc, x_test, y_test))
    print(f'Accuracy: {calculate_accuracy(nbc, x_test, y_test)}')
print(f'Average accuracy after 3-fold training is {np.array(accuracy).mean()}')
```

```
Accuracy: 0.95
Accuracy: 0.9661016949152542
Accuracy: 1.0
Average accuracy after 3-fold training is 0.9720338983050847
```

### 3. Softmax and Cross Entropy Loss.

(a) I did one PyTorch model without softmax and one PyTorch model with softmax. The output without softmax is a cluster of large positive or negative values. The output with softmax is more like probabilities that sum up to 1.

In [10]:

```
# First convert the features and labels to PyTorch tensors
pytorch_features = torch.tensor(wines.loc[:, 'Alcohol %': 'Proline'].values, dtype=torch.float32)
pytorch_labels = torch.tensor(wines.loc[:, 'ranking'].values, dtype=torch.int64)
```

```
# Then define a pytorch model without softmax
model_no_softmax = nn.Sequential(
    nn.Linear(pytorch_features.shape[1], len(np.unique(pytorch_labels)))
)

# Then pass the data through the model once without backpropagation
outputs_no_softmax = model_no_softmax(pytorch_features)

# Finally print out the outputs_no_softmax
print(outputs_no_softmax)
```

```
tensor([[ -95.7595, -80.1433, -271.1181],
        [ -70.3680, -55.5473, -186.5312],
        [ -89.5526, -77.3873, -266.7516],
        [-105.8480, -94.7881, -328.4174],
        [-107.6909, -97.2809, -338.8018],
        [-109.6877, -95.2311, -327.6151],
        [ -75.2029, -59.1946, -197.1850],
        [ -86.9012, -75.8067, -259.8968],
        [ -74.8449, -63.2743, -215.8418],
        [-105.9978, -95.4717, -329.8702],
        [ -88.4649, -76.9907, -264.4617],
        [-108.5639, -93.0019, -315.7316],
        [ -79.9209, -66.1011, -224.1377],
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        [ -92.5969, -72.4996, -237.2635],
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        [ -65.3198, -53.5460, -180.6982],
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        [ -56.0758, -42.3140, -141.3992],
        [ -44.3542, -33.0702, -109.6971],
        [ -39.9226, -29.4330, -97.2258],
        [ -41.9280, -30.9954, -102.9244],
        [ -58.9969, -47.1536, -158.7496],
        [ -39.0347, -27.9414, -95.3846],
        [ -37.5140, -26.1886, -85.9839],
        [ -52.5705, -42.9040, -147.9189],
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        [ -58.1301, -44.9303, -152.1356],
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        [ -50.9161, -39.1205, -131.8629],
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        [ -55.1265, -40.5552, -138.4230],
```

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[ -48.6012, -37.7607, -130.6048],  
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[ -43.6103, -32.3360, -107.6418],  
[ -60.5434, -49.1194, -167.6660],

```

[ -35.6747, -24.3274, -78.7909],
[ -47.1800, -36.5285, -123.9038],
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[ -38.8315, -27.9552, -95.4119],
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[ -63.7631, -48.4041, -159.2106],
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[ -55.9494, -45.4670, -158.8678],
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[ -56.0662, -44.7320, -155.9557],
[ -68.2930, -55.3038, -190.7732],
[ -60.3732, -46.4871, -159.5021],
[ -76.4289, -61.6658, -211.7994],
[ -77.3098, -62.5640, -213.1622],
[ -52.7916, -40.5824, -141.4767],
[ -48.2669, -38.1544, -129.4740]], grad_fn=<AddmmBackward0>)

```

```

In [11]: # Second is to define a pytorch model with softmax
model_softmax = nn.Sequential(
    nn.Linear(pytorch_features.shape[1], len(np.unique(pytorch_labels))),
    nn.Softmax(dim=1)
)

# Then pass the data through the model once without backpropagation
outputs_softmax = model_softmax(pytorch_features)

# Finally print out the outputs_softmax
print(outputs_softmax)

```

```
tensor([[ 1.8472e-04,  0.0000e+00,  9.9982e-01],
        [ 8.6887e-05,  0.0000e+00,  9.9991e-01],
        [ 3.2321e-02,  0.0000e+00,  9.6768e-01],
        [ 5.2833e-01,  0.0000e+00,  4.7167e-01],
        [ 8.3785e-01,  0.0000e+00,  1.6215e-01],
        [ 1.6370e-02,  0.0000e+00,  9.8363e-01],
        [ 8.5346e-06,  0.0000e+00,  9.9999e-01],
        [ 7.0744e-02,  0.0000e+00,  9.2926e-01],
        [ 1.5047e-02,  0.0000e+00,  9.8495e-01],
        [ 6.7305e-01,  0.0000e+00,  3.2695e-01],
        [ 5.4550e-02,  0.0000e+00,  9.4545e-01],
        [ 8.9474e-04,  0.0000e+00,  9.9911e-01],
        [ 4.5460e-04,  0.0000e+00,  9.9955e-01],
        [ 1.0055e-01,  0.0000e+00,  8.9945e-01],
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```

(b) I unfortunately always encounter an error of dead kernel when trying to evaluate the train\_and\_val function. I don't really understand the root cause of the issue.

```
In [14]: def train_and_val(model,train_X,train_y,epochs,draw_curve=True):
    """
    Parameters
    -----
    model: a PyTorch model
    train_X: np.array shape(ndata,nfeatures)
    train_y: np.array shape(ndata)
    epochs: int
    draw_curve: bool
    """

    ### Define your loss function, optimizer. Convert data to torch tensor ###
    loss = nn.CrossEntropyLoss()
    optimizer = optim.Adam(model.parameters(), lr = 0.001)

    train_y -= 1
    Xs = torch.tensor(train_X).float()
```

```

ys = torch.tensor(train_y).long()

# Define Kfolds
kf = KFold(n_splits = 3, shuffle = True)
for train_index, test_index in kf.split(Xs):
    train_X, test_X = Xs[train_index], Xs[test_index]
    train_y, test_y = ys[train_index], ys[test_index]

### Split training examples further into training and validation ###
train_X, val_X, train_y, val_y = train_test_split(train_X, train_y, test_size = 0.20)
val_array=[]
lowest_val_loss = np.inf

for i in range(epochs):
    ### Compute the loss and do backpropagation ###
    optimizer.zero_grad()
    train_out = model(train_X)
    train_loss = loss(train_out, train_y)
    train_loss.backward()
    optimizer.step()

    ### compute validation loss and keep track of the lowest val loss ###
    # compute validation loss
    val_out = model(val_X)
    val_loss = loss(val_out, val_y)

    # append val loss to val_array
    val_array.append(val_loss.item())

    # keep track of the lowest val loss
    if val_loss < lowest_val_loss:
        lowest_val_loss = val_loss
        torch.save(model.state_dict(), 'model.pt')

# The final number of epochs is when the minimum error in validation set occurs

final_epochs = np.argmin(val_array) + 1
print("Number of epochs with lowest validation:", final_epochs)
### Recover the model weight ###
model.load_state_dict(torch.load('model.pt'))
model.eval()

### Plot the validation loss curve ###

if draw_curve:
    plt.figure()
    plt.plot(np.arange(len(val_array))+1, val_array, label='Validation loss')
    plt.xlabel('Epochs')
    plt.ylabel('Loss')
    plt.legend()

```

```
In [ ]: train_and_val(model_softmax,pytorch_features,pytorch_labels,1000,draw_curve=True)
```

```
In [ ]:
```