Chem277B: Machine Learning Algorithms

Homework assignment #5: Regression

```
In [3]: 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:

$$P[M|+] = \frac{P[+|M] * P[M]}{P[+|M] * P[M] + P[+|notM] * P[notM]}$$

$$=\frac{0.95\times0.01}{(0.95\times0.01)+(0.05\times0.99)}=0.161$$

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:

$$P[M|+]' = rac{P[+|M]' * P[M]'}{P[+|M]' * P[M]' + P[+|notM]' * P[notM]'}$$

$$=rac{0.95 imes0.1}{(0.95 imes0.1)+(0.05 imes0.9)}=0.679$$

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 [4]: 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 {
m 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:
        quess=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 quess
```

```
In [5]: # Import wines.csv
wines = pd.read_csv('wines.csv')
wines.head()
```

Out[5]:		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 [6]: # 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 [7]: # 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[7]:		Alcohol %	Malic Acid	Ash	Alkalinity	Mg	Phenols	Flavanoids	Phenols.1	Proantho- cyanins	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	1.013009
	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	0.949319
	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	1.69767
	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	1.82505{

```
In [8]: # 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()}')
```

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 [9]: # 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)
```

```
tensor([[-4.5661e+01, -1.3691e+01, -1.7287e+02],
        [-3.8358e+01, -4.7560e+00, -1.1983e+02],
        [-3.7913e+01, -1.7992e+01, -1.6826e+02],
        [-4.0346e+01, -2.5418e+01, -2.0616e+02],
        [-3.9303e+01, -2.7642e+01, -2.1222e+02],
        [-4.6351e+01, -2.1539e+01, -2.0694e+02],
        [-4.1492e+01, -5.2367e+00, -1.2725e+02],
        [-3.6758e+01, -1.7042e+01, -1.6418e+02],
        [-3.4566e+01, -1.1880e+01, -1.3688e+02],
        [-4.0340e+01, -2.5281e+01, -2.0730e+02],
        [-3.7320e+01, -1.7710e+01, -1.6708e+02],
        [-4.9081e+01, -1.7736e+01, -2.0063e+02],
        [-3.8823e+01, -1.0421e+01, -1.4267e+02],
        [-3.8718e+01, -1.9284e+01, -1.7850e+02],
        [-3.9538e+01, -1.8746e+01, -1.7665e+02],
        [-3.8050e+01, -1.1400e+01, -1.4363e+02],
        [-3.8947e+01, -1.7775e+01, -1.7098e+02],
        [-3.9717e+01, -2.5400e+01, -2.0357e+02],
        [-4.3327e+01, -2.0096e+01, -1.9217e+02],
        [-4.3692e+01, -2.3754e+01, -2.0740e+02],
        [-3.3863e+01, -5.3006e+00, -1.1086e+02],
        [-2.6572e+01, -6.6408e-01, -6.8693e+01],
        [-2.5513e+01, -5.0173e+00, -8.1033e+01],
        [-3.6483e+01, -1.1233e+01, -1.4137e+02],
        [-3.3833e+01, 2.1536e+00, -8.2306e+01],
        [-2.6807e+01, -8.2227e+00, -1.0236e+02],
        [-2.9793e+01, -2.4009e+00, -8.5069e+01],
        [-3.0256e+01, 8.2731e-01, -7.4071e+01],
        [-2.8945e+01, -8.5113e+00, -1.1031e+02],
        [-2.4809e+01, -9.2888e+00, -1.0138e+02],
        [-5.3220e+01, -3.3267e+00, -1.5456e+02],
        [-4.1035e+01, 1.7825e+00, -1.0353e+02],
        [-2.6666e+01, 6.0689e-01, -6.5912e+01],
        [-3.3164e+01, -7.2037e+00, -1.1552e+02],
        [-2.6325e+01, 7.8301e-02, -6.8174e+01],
        [-3.3421e+01, -9.2280e-01, -9.2022e+01],
        [-2.6889e+01, -6.3643e-01, -7.1323e+01],
        [-2.5043e+01, 1.6478e-01, -6.3227e+01],
        [-2.5847e+01, -2.9022e-02, -6.6891e+01],
        [-3.1516e+01, -4.8608e+00, -1.0194e+02],
        [-2.4972e+01, 1.3769e+00, -6.1246e+01],
        [-2.5443e+01, 1.9359e+00, -5.6399e+01],
        [-2.6197e+01, -5.6942e+00, -9.3593e+01],
        [-2.9933e+01, -3.2600e+00, -9.1049e+01],
        [-3.2797e+01, -2.4039e+00, -9.7599e+01],
        [-2.7745e+01, -4.8797e+00, -9.3400e+01],
        [-3.1232e+01, -2.9310e+00, -9.6214e+01],
        [-2.9239e+01, -1.5035e+00, -8.4516e+01],
        [-2.3179e+01, -9.4961e-01, -6.6196e+01],
        [-3.2858e+01, -7.6090e-01, -8.9076e+01],
```

```
[-3.2474e+01, 3.4966e+00, -7.0152e+01],
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[-3.3932e+01, -5.7373e+00, -1.1218e+02],
[-2.8227e+01, -9.0485e+00, -1.0985e+02],
[-2.8914e+01, -8.0324e+00, -1.0596e+02],
[-3.0905e+01, -8.5678e+00, -1.1858e+02],
[-3.3078e+01, -8.1377e+00, -1.2107e+02],
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[-3.7636e+01, -1.8309e+01, -1.6853e+02],
[-3.8030e+01, -2.2234e+01, -1.8445e+02],
[-4.3009e+01, -1.7953e+01, -1.8281e+02],
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[-3.8831e+01, -1.4875e+01, -1.6010e+02],
[-4.1581e+01, -1.7072e+01, -1.7730e+02],
[-3.7959e+01, -1.6594e+01, -1.6493e+02],
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[-3.2016e+01, -7.1346e+00, -1.1076e+02],
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[-2.7294e+01, -2.6138e+00, -8.1114e+01],
[-2.3699e+01, 5.1893e-01, -5.9989e+01],
```

```
[-2.8457e+01, -3.9164e+00, -9.1628e+01],
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[-2.7501e+01, -1.2250e+00, -7.6370e+01],
[-3.1365e+01, -8.9508e-01, -8.6172e+01],
[-2.9951e+01, -8.2420e+00, -1.1598e+02],
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[-2.6079e+01, -3.6662e+00, -8.2860e+01],
[-2.8006e+01, -2.0993e+00, -8.0897e+01],
[-2.6025e+01, -1.6696e+00, -7.3420e+01],
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[-2.6492e+01, -7.9709e-01, -6.9775e+01],
[-3.0302e+01, -7.8865e+00, -1.0711e+02],
```

```
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                 [-3.4280e+01, 2.4063e+00, -7.6883e+01],
                 [-2.8654e+01, 4.2772e+00, -6.0093e+01],
                 [-2.4173e+01, 2.7400e-01, -6.1546e+01],
                 [-2.5307e+01, 6.6712e-01, -6.1938e+01],
                 [-3.8186e+01, -8.0562e-01, -1.0344e+02],
                 [-2.8584e+01, -7.7505e+00, -1.0517e+02],
                 [-3.0867e+01, -7.7628e+00, -1.1226e+02],
                 [-2.8262e+01, -1.1577e+00, -8.3253e+01],
                 [-3.7245e+01, -9.5428e+00, -1.3832e+02],
                 [-2.7080e+01, -6.8124e+00, -1.0027e+02],
                 [-3.4871e+01, 2.2189e+00, -8.1707e+01],
                 [-3.1473e+01, 6.0737e-01, -7.8050e+01],
                 [-3.0747e+01, -7.2281e+00, -1.0843e+02],
                 [-3.2382e+01, -5.1479e+00, -1.0411e+02],
                 [-3.0099e+01, -8.7073e+00, -1.1630e+02],
                 [-2.9919e+01, -8.1043e+00, -1.0625e+02],
                 [-2.7449e+01, -7.8829e+00, -9.9681e+01],
                 [-2.7839e+01, -6.8794e+00, -9.8818e+01],
                 [-3.3616e+01, -8.4037e+00, -1.2130e+02],
                 [-3.3476e+01, -2.9543e+00, -1.0206e+02],
                 [-3.8444e+01, -8.0253e+00, -1.3495e+02],
                 [-3.8981e+01, -8.3581e+00, -1.3623e+02],
                 [-2.8324e+01, -3.8579e+00, -8.9997e+01],
                 [-2.5720e+01, -4.4445e+00, -8.2650e+01]], grad fn=<AddmmBackward0>)
In [10]: # 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)
```

[-2.5326e+01, 3.3629e+00, -5.1772e+01], [-2.6120e+01, -3.2871e+00, -7.9219e+01],

```
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```

```
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```

```
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[1.3066e-33, 0.0000e+00, 1.0000e+00],
[7.8519e-31, 0.0000e+00, 1.0000e+00]], grad fn=<SoftmaxBackward0>)
```

(b) The finished codes are listed below. I unfortunately always encounter an error of dead kernel when trying to evaluate the train_and_val function in Jupyter Notebook. Hence I ran all the codes in Google Colab and got some interesting results.

```
In [11]: def train_and_val(model, train_X, train_y, epochs, draw_curve=True):
    """
    Train and validate a PyTorch model using cross-entropy loss.

Parameters
------
model: PyTorch model
    The model to train.
    train_X: numpy.ndarray
    The input training data of shape (n_samples, n_features).
    train_y: numpy.ndarray
    The target training data of shape (n_samples,).
    epochs: int
    The number of training epochs.
    draw_curve: bool, optional
    Whether to draw a validation loss curve (the default is True).
```

```
Returns
float
    The final validation loss.
# Convert data to torch tensor
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]
    # Subtract one from labels to make them 0-based
    train y -= 1
    test y -= 1
    # 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=0.2)
    # Define loss function and optimizer
    loss fn = nn.CrossEntropyLoss()
    optimizer = optim.Adam(model.parameters(), lr=0.001)
    # Keep track of the lowest validation loss
    lowest val loss = np.inf
    # Train the model
    val array = []
    for epoch in range(epochs):
        # Compute training loss and update model parameters
        optimizer.zero grad()
        train out = model(train X)
        train loss = loss fn(train out, train y)
        train loss.backward()
        optimizer.step()
        # Compute validation loss and keep track of the lowest validation loss
        val out = model(val X)
        val loss = loss fn(val out, val y)
        val array.append(val loss.item())
        if val loss < lowest val loss:</pre>
            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()
    # Compute test accuracy
    test out = model(test X)
    test loss = loss fn(test out, test y)
    test preds = torch.argmax(test out, dim=1).numpy()
    test acc = np.mean(test preds == test y.numpy())
    print("Test accuracy:", test acc)
    # 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()
return lowest val loss.item()
```

For evaluation of the function, I used the 2 models generated above, model_no_softmax and model_softmax. I also created two additional models, one with softmax as the last layer activation function and one with ReLu as the last layer activation function. Their performance in terms of convergence and accuracy of prediction as shown below. In general it seems model_no_softmax, the simpliest model gives the best prediction performance. My guess is the other models are over-fitting?

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

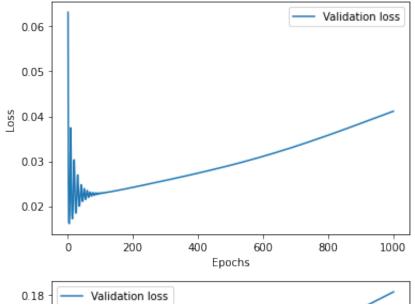
```
In [ ]: class Classifier softmax(nn.Module):
            def init (self, input dim, hidden dim, num classes):
                super(Classifier softmax, self). init ()
                self.fc1 = nn.Linear(input dim, hidden dim)
                self.fc2 = nn.Linear(hidden dim, num classes)
            def forward(self, x):
                out = self.fcl(x)
                out = nn.functional.relu(out)
                out = self.fc2(out)
                out = nn.functional.softmax(out, dim=1)
                return out
        wine classifier = Classifier softmax(pytorch features.shape[1], len(np.unique(pytorch labels)), 3)
        train and val(wine classifier, pytorch features, pytorch labels, 1000, draw curve=True)
In [ ]: class Classifier_Relu(nn.Module):
            def init (self, input dim, hidden dim, num classes):
                super(Classifier Relu, self). init ()
                self.fc1 = nn.Linear(input dim, hidden dim)
                self.fc2 = nn.Linear(hidden_dim, num_classes)
                self.relu = nn.ReLU()
            def forward(self, x):
                out = self.fcl(x)
                out = self.relu(out)
                out = self.fc2(out)
                return out
        wine classifier 2 = Classifier Relu(pytorch features.shape[1], len(np.unique(pytorch labels)), 3)
        train and val(wine classifier 2, pytorch features, pytorch labels, 1000, draw curve=True)
```

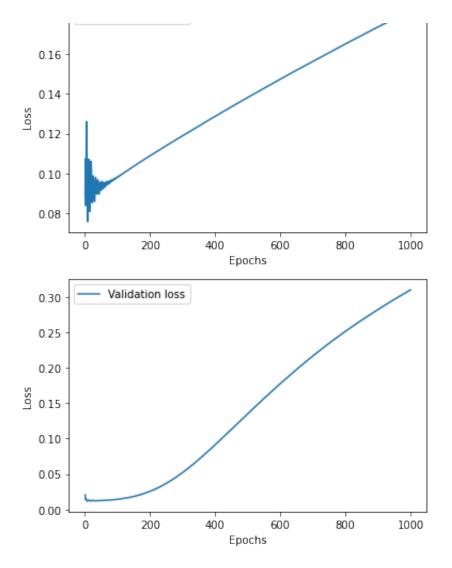
```
# Plot the validation loss curve
if draw_curve:
    plt.figure()
    plt.plot(np.arange(len(val_ar
        plt.xlabel('Epochs')
        plt.ylabel('Loss')
        plt.legend()
```

return lowest_val_loss.item()

train_and_val(model_no_softmax, pytorch_f

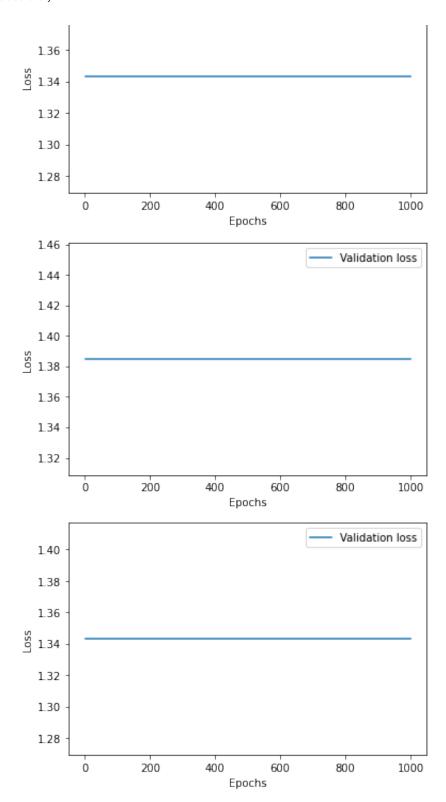
```
<ipython-input-97-f0eea764c926>:24: UserWarning: To copy construct from a tent
    Xs = torch.tensor(train_X).float()
<ipython-input-97-f0eea764c926>:25: UserWarning: To copy construct from a tent
    ys = torch.tensor(train_y).long()
Number of epochs with lowest validation: 5
Test accuracy: 0.966666666666667
Number of epochs with lowest validation: 9
Test accuracy: 0.9830508474576272
Number of epochs with lowest validation: 7
Test accuracy: 1.0
0.011702748946845531
```





train_and_val(model_softmax, pytorch_feat

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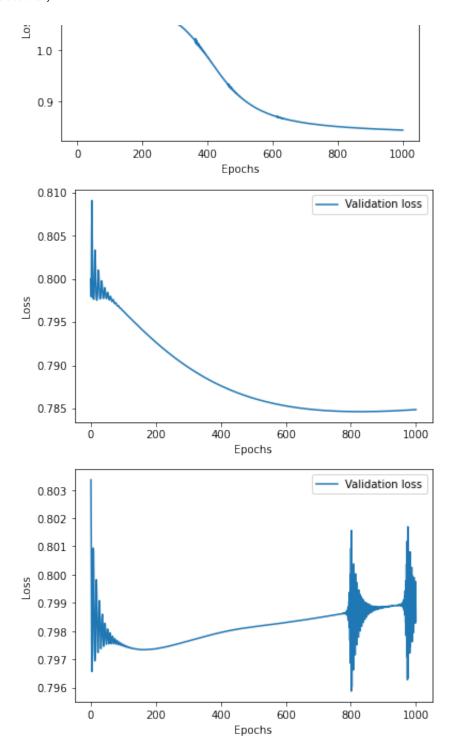


```
class Classifier_softmax(nn.Module):
    def __init__(self, input_dim, hidden_
        super(Classifier_softmax, self)._
        self.fc1 = nn.Linear(input_dim, h
        self.fc2 = nn.Linear(hidden_dim,

    def forward(self, x):
        out = self.fc1(x)
        out = nn.functional.relu(out)
        out = self.fc2(out)
        out = nn.functional.softmax(out, return out)
```

```
wine_classifier = Classifier_softmax(pyto
```

train_and_val(wine_classifier, pytorch_fe



```
class Classifier_Relu(nn.Module):
    def __init__(self, input_dim, hidden_
        super(Classifier_Relu, self).__in
        self.fc1 = nn.Linear(input_dim, h
        self.fc2 = nn.Linear(hidden_dim,
        self.relu = nn.ReLU()

    def forward(self, x):
        out = self.fc1(x)
        out = self.relu(out)
        out = self.fc2(out)
        return out
```

```
wine_classifier_2 = Classifier_Relu(pytor
train_and_val(wine_classifier_2, pytorch_
```

```
<ipython-input-71-f0eea764c926>:24: UserWarning: To copy construct from a ten
   Xs = torch.tensor(train_X).float()
<ipython-input-71-f0eea764c926>:25: UserWarning: To copy construct from a ten
   ys = torch.tensor(train_y).long()
Number of epochs with lowest validation: 1000
Test accuracy: 0.6166666666666667
Number of epochs with lowest validation: 992
Test accuracy: 0.6949152542372882
Number of epochs with lowest validation: 836
Test accuracy: 0.711864406779661
0.4345228970050812
Walidation loss
```



