Homework 3

School & Student ID: NYCU / 513 559 004 Course Title: Machine Learning

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Suppose the union $H_1 \cup H_2$ can leverage hypotheses from both sets to shatter more points than either set could alone.

$$d_{vc}(H_1 \cup H_2) > d_{vc}(H_1) + d_{vc}(H_2)$$

However, for any labeling on a set of $d_{vc}(H_1) + d_{vc}(H_2)$ points, at least one of the hypothesis sets H_1 or H_2 must fully accommodate that labeling.

Thus, the inequality above would require one of the sets to shatter more points than its individual VC dimension, which is impossible.

Therefore,

$$d_{vc}(H_1 \cup H_2) \le d_{vc}(H_1) + d_{vc}(H_2)$$

Given that,

$$f_{0/1}(x) = sign(P(y = +1 \mid x) - \frac{1}{2})$$

It implies that we classify y = +1 only if $P(y = +1 \mid x) > 0.5$.

Then, we adjust the classification rule with asymmetric error costs:

$$f_{MKT}(x) = sign(P(y = + 1 | x) - \alpha)$$

Since false negatives are 10 times more costly, it is necessary to have a threshold much more likely to classify as +1 to avoid the penalty:

$$\frac{P(y=+1 \mid x)}{P(y=-1 \mid x)} \ge 10$$

$$\Rightarrow \frac{P(y=+1 \mid x)}{1-P(y=+1 \mid x)} \ge 10$$

$$\Rightarrow P(y=+1 \mid x) \ge \frac{10}{11}$$

Hence, we can put $P(y = +1 \mid x)$ into the function $f_{MKT}(x)$, and we'll find:

$$\alpha = \frac{10}{11}$$

Given that,

$$E_{out}^{(2)}(h) = E_{x \sim P(x), y \sim (y|x)}[h(x) \neq y]$$

$$\Rightarrow E_{out}^{(2)}(h) = E_{x \sim P(x)}[P(h(x) \neq y \mid x)]$$

Decomposing it with the law of total probability:

$$P(h(x) \neq y \mid x) = P(h(x) \neq f(x) \text{ and } f(x) = y \mid x) + P(f(x) \neq y \mid x)$$

By linearity of expectation:

$$E_{out}^{(2)}(h) = E_x[P(h(x) \neq f(x) \text{ and } f(x) = y \mid x)] + E_x[P(f(x) \neq y \mid x)]$$

The first term relates to $E_{out}^{(1)}(h)$, as:

$$P(h(x) \neq f(x) \text{ and } f(x) = y \mid x) \leq P(h(x) \neq f(x) \mid x)$$

 $\Rightarrow E_x[P(h(x) \neq f(x) \text{ and } f(x) = y \mid x)] \leq E_{out}^{(1)}(h)$

Combining components, therefore:

$$E_{out}^{(2)}(h) \le E_{out}^{(1)}(h) + E_{out}^{(2)}(f)$$

Knowing that,

$$W_{LIN} = \left(X^T X\right)^{-1} X^T y$$

The matrix *X* can be written as:

$$X = egin{pmatrix} 1 & x_{11} & x_{12} & \dots & x_{1d} \ 1 & x_{21} & x_{22} & \dots & x_{2d} \ dots & dots & dots & dots \ 1 & x_{N1} & x_{N2} & \dots & x_{Nd} \end{pmatrix}$$

Using the diagonal matrix X' = XD to scale the intercept term by 1126.

$$\mathbf{D} = egin{pmatrix} 1126 & 0 & 0 & \dots & 0 \ 0 & 1 & 0 & \dots & 0 \ 0 & 0 & 1 & \dots & 0 \ dots & dots & dots & \ddots & dots \ 0 & 0 & 0 & \dots & 1 \end{pmatrix}$$

$$X' = egin{pmatrix} 1126 & x_{11} & x_{12} & \dots & x_{1d} \ 1126 & x_{21} & x_{22} & \dots & x_{2d} \ dots & dots & dots & \ddots & dots \ 1126 & x_{N1} & x_{N2} & \dots & x_{Nd} \end{pmatrix}$$

After changing x_0 to 1126, the weight vector W_{LUCKY} for X' is shown as:

$$W_{LUCKY} = \left((X')^T X' \right)^{-1} (X')^T y$$

Since X' = XD, we can substitute X' with XD as follows:

$$\Rightarrow W_{LUCKY} = \left(D^T X^T X D\right)^{-1} \left(D^T X^T y\right)$$

We know that $(D^T X^T X D)^{-1} = D^{-1} (X^T X)^{-1} D^{-1}$, we can replace it into the equation for W_{LUCKY} :

$$W_{LUCKY} = D^{-1} (X^{T} X)^{-1} X^{T} y$$

$$\Rightarrow W_{LIN} = D \cdot W_{LUCKY}$$

Given that,

$$\hat{h}(x) = \frac{1}{2} \left(\frac{w^T x}{\sqrt{1 + (w^T x)^2}} + 1 \right)$$

Minimize the negative log-likelihood for the data $\{(x_n, y_n)\}$:

$$\widehat{E}_{in}(w) = -\frac{1}{N} \left(y_n ln(\widehat{h}(x)) + (1 - y_n) ln(1 - \widehat{h}(x)) \right)$$

Let $u = w^T x$.

$$\frac{d\hat{h}}{du} = \frac{1}{2} \cdot \frac{(1+u^2)-u^2}{(1+u^2)^{\frac{3}{2}}} = \frac{1}{2} \cdot \frac{1}{(1+u^2)^{\frac{3}{2}}}$$

Then, the gradient $\nabla_{w} \hat{h}$ we find out would be:

$$\Rightarrow \nabla_{w} \hat{h} = \frac{d\hat{h}}{du} \cdot x$$

We can calculate the gradient $\nabla \widehat{E}_{in}(w)$:

$$\nabla \widehat{E}_{in}(w) = -\frac{1}{N} \sum_{n=1}^{N} \left(\frac{y_n}{\widehat{h}(x_n)} - \frac{1 - y_n}{1 - \widehat{h}(x_n)} \right) \nabla_w \widehat{h}(x_n)$$

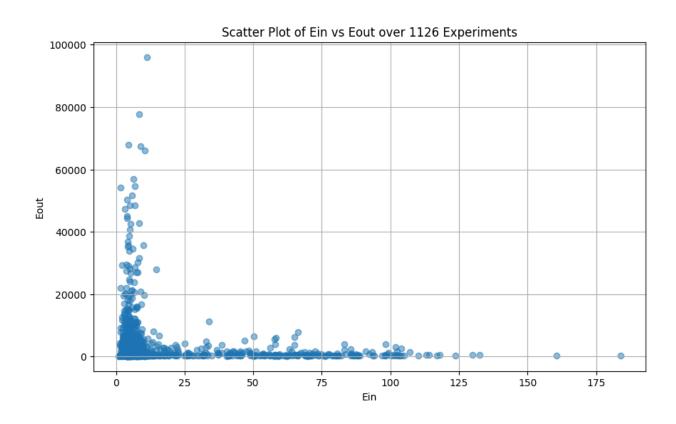
Substitute $\nabla_{w} \hat{h}(x)$:

$$\Rightarrow \nabla \widehat{E}_{in}(w) = -\frac{1}{N} \sum_{n=1}^{N} \left(\frac{y_n - \widehat{h}(x_n)}{\widehat{h}(x_n) \left(1 - \widehat{h}(x_n)\right)} \right) \cdot \frac{1}{2} \cdot \frac{x_n}{\left(1 + \left(w^T x_n\right)^2\right)^{\frac{3}{2}}}$$

The first page of the snapshot of my *code is on the next page*.

My findings:

- 1. The scatter plot shows the relationship between Ein and Eout for different experiments.
- 2. Generally, Ein and Eout are *positively correlated*, meaning that when the model performs well on training data, it also tends to perform well on unseen data.
- 3. However, some variability indicates that a low Ein does not always guarantee a low Eout. This suggests that *overfitting can still occur* in some cases, depending on the specific training samples.

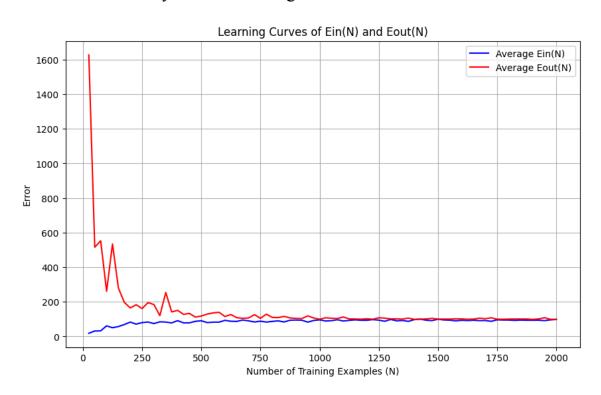


```
import numpy as np
       import matplotlib.pyplot as plt
       from sklearn.linear_model import LinearRegression
       from google.colab import drive
       # Load the dataset
       drive.mount('/d')
       file_path = '/d/MyDrive/ML_hw3_attachment-cpusmall_scale.txt'
       data = []
       with open(file_path, 'r') as f:
           for line in f:
               values = line.strip().split()
               y = float(values[0])
               x = [float(v.split(':')[1]) for v in values[1:]]
               data.append((x, y))
       X_full = np.array([x for x, y in data])
       y_full = np.array([y for x, y in data])
       # Parameters
       N = 32
       num_experiments = 1126
       Ein_list = []
       Eout_list = []
       # Run the experiments
       for _ in range(num_experiments):
           # Randomly sample N examples for training
           indices = np.random.choice(len(X_full), N, replace=False)
           X_train = X_full[indices]
           y_train = y_full[indices]
           # Add bias term (x_0 = 1)
           X_train = np.hstack((np.ones((X_train.shape[0], 1)), X_train))
           X_full_bias = np.hstack((np.ones((X_full.shape[0], 1)), X_full))
           # Fit linear regression model
           model = LinearRegression(fit_intercept=False)
           model.fit(X_train, y_train)
40
           w_lin = model.coef_
           # Calculate Ein (in-sample error)
           y_train_pred = X_train @ w_lin
           Ein = np.mean((y_train - y_train_pred) ** 2)
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           Ein_list.append(Ein)
```

The first page of the snapshot of my *code is on the next page*.

My findings:

- 1. As N increases,
 - a. both Ein and Eout tend to *decrease*, indicating improved model generalization with more training data.
 - b. The gap between Ein and Eout *narrows*, suggesting reduced overfitting with larger training sizes.
 - c. Both *converge* to similar values, indicating that the model is better at generalizing to new data, reducing bias and variance.
- 2. For smaller values of N, there is a significant difference between Ein and Eout, which indicates *overfitting*. *The model fits the training data well but struggles to generalize to unseen data*.
- 3. The learning curve shows that *adding more training data* helps the model improve its performance, especially for smaller training set sizes where the model initially suffers from high variance.

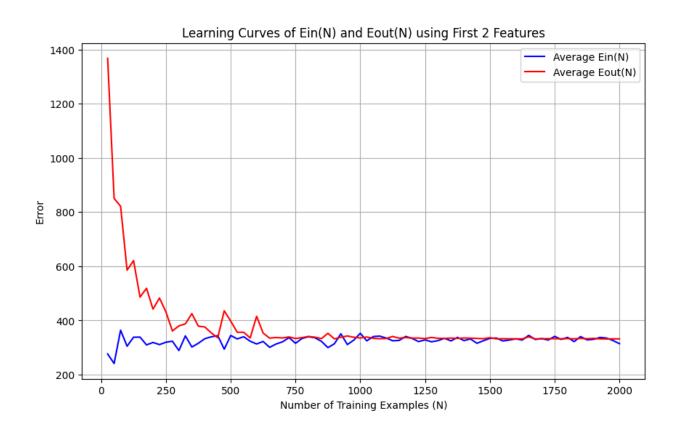


```
with open(file_path, 'r') as f:
10
           for line in f:
               values = line.strip().split()
               y = float(values[0])
               x = [float(v.split(':')[1]) for v in values[1:]]
               data.append((x, y))
       X_full = np.array([x for x, y in data])
       y_full = np.array([y for x, y in data])
       # Parameters
20
       N_values = np.arange(25, 2001, 25)
       num_experiments = 16
       average_Ein = []
       average_Eout = []
       # Run experiments for each value of N
       for N in N_values:
           Ein_total = 0
           Eout_total = 0
30
           for _ in range(num_experiments):
               # Randomly sample N examples for training
               indices = np.random.choice(len(X_full), N, replace=False)
               X_train = X_full[indices]
               y_train = y_full[indices]
               # Add bias term (x_0 = 1)
               X_train = np.hstack((np.ones((X_train.shape[0], 1)), X_train))
               X_full_bias = np.hstack((np.ones((X_full.shape[0], 1)), X_full))
               # Fit linear regression model
               model = LinearRegression(fit_intercept=False)
               model.fit(X_train, y_train)
44
               w_lin = model.coef_
               # Calculate Ein (in-sample error)
               y_train_pred = X_train @ w_lin
               Ein = np.mean((y_train - y_train_pred) ** 2)
               Ein_total += Ein
               # Calculate Eout (out-of-sample error)
               y_full_pred = X_full_bias @ w_lin
```

The first page of the snapshot of my *code is on the next page*.

My findings:

- 1. With only the first 2 features, both Ein and Eout are *generally higher compared to using all 12 features*, indicating a decrease in model performance.
- 2. The gap between Ein and Eout *remains more significant for smaller values of N*, suggesting that the reduced feature set leads to increased bias and potentially higher variance.
- 3. As N increases, the errors decrease, but they *do not reach as low values as when using all features*, demonstrating that the model benefits from having more features to learn from.



```
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       X_full = np.array([x[:2] for x, y in data]) # Use only the first 2 features
       y_full = np.array([y for x, y in data])
20
       # Parameters
       N_values = np.arange(25, 2001, 25)
       num experiments = 16
       average_Ein = []
       average_Eout = []
       # Run experiments for each value of N
       for N in N_values:
28
           Ein_total = 0
           Eout_total = 0
           for in range(num experiments):
               # Randomly sample N examples for training
               indices = np.random.choice(len(X_full), N, replace=False)
               X_train = X_full[indices]
               y_train = y_full[indices]
               # Add bias term (x_0 = 1)
               X_train = np.hstack((np.ones((X_train.shape[0], 1)), X_train))
               X_full_bias = np.hstack((np.ones((X_full.shape[0], 1)), X_full))
40
               # Fit linear regression model
               model = LinearRegression(fit_intercept=False)
               model.fit(X_train, y_train)
44
               w_lin = model.coef_
               # Calculate Ein (in-sample error)
               y_train_pred = X_train @ w_lin
               Ein = np.mean((y_train - y_train_pred) ** 2)
               Ein total += Ein
               # Calculate Eout (out-of-sample error)
               y_full_pred = X_full_bias @ w_lin
               Eout = np.mean((y_full - y_full_pred) ** 2)
               Eout total += Eout
           # Calculate average Ein and Eout
           average_Ein.append(Ein_total / num_experiments)
           average_Eout.append(Eout_total / num_experiments)
```

Knowing that B(N, k) covers every possible subset size from 0 up to k - 1.

$$B(N, k) \le \sum_{i=0}^{k-1} C_i^N$$

Precisely, B(N, k) should count all subsets that can be formed up to k elements. Such that, it appears B(N, k) is at least large enough to include all such subsets.

$$B(N, k) = \sum_{i=0}^{k} C_i^N \ge \sum_{i=0}^{k-1} C_i^N$$

$$\Rightarrow B(N, k) \ge \sum_{i=0}^{k-1} C_i^N$$

Since B(N, k) includes all subsets up to that size, it cannot be less than $\sum_{i=0}^{k-1} C_i^N$. In

addition, we have already known $B(N, k) \le \sum_{i=0}^{k-1} C_i^N$, we can eventually establish:

$$B(N, k) = \sum_{i=0}^{k-1} C_i^N$$