2 Probabilistic Model of Linear Regression

Both ordinary least squares and ridge regression have interpretations from a probabilistic standpoint. In particular, assuming a generative model for our data and a particular noise distribution, we will derive least squares and ridge regression as the maximum likelihood and maximum a-posteriori parameter estimates, respectively. This problem will walk you through a few steps to do that. (Along with some side digressions to make sure you get a better intuition for ML and MAP estimation.)

(a) Assume that \( X \) and \( Y \) are both one-dimensional random variables, i.e. \( X, Y \in \mathbb{R} \). Assume an affine model between \( X \) and \( Y \): \( Y = Xw_1 + w_0 + Z \), where \( w_1, w_0 \in \mathbb{R} \), and \( Z \sim N(0, 1) \) is a standard normal (Gaussian) random variable. Assume \( w_1, w_0 \) are fixed parameters (i.e., they are not random). What is the conditional distribution of \( Y \) given \( X \)?

**Solution:** When we condition on the event \( X = x \), the expression \( Xw_1 + w_0 \) becomes \( xw_1 + w_0 \), so

\[
Y | (X = x) \sim xw_1 + w_0 + Z.
\]

Now \( xw_1 + w_0 + Z \) is a constant plus a standard normal, so

\[
xw_1 + w_0 + Z \sim N(xw_1 + w_0, 1).
\]

The conditional density becomes:

\[
p(Y | X = x) = \frac{1}{\sqrt{2\pi}} \exp\left\{-\frac{1}{2}(Y - (xw_1 + w_0))^2\right\}.
\]

(b) Given \( n \) points of training data \( \{(X_1, Y_1), (X_2, Y_2), \ldots, (X_n, Y_n)\} \) generated in an iid fashion by the probabilistic setting in the previous part, derive the maximum likelihood estimator for \( w_1, w_0 \) from this training data.
Solution: The log likelihood function is given by
\[
\sum_{i=1}^{n} \log p(Y_i | X = X_i) = \sum_{i=1}^{n} \log \left( \frac{1}{\sqrt{2\pi}} \exp \left( -\frac{1}{2} (Y_i - (X_i w_1 + w_0))^2 \right) \right) = -\frac{1}{2} \sum_{i=1}^{n} (Y_i - (X_i w_1 + w_0))^2 + \text{Constant} \tag{2}
\]

Differentiating the log likelihood with respect to \(w_1\) and \(w_0\), we obtain:
\[
\frac{\partial}{\partial w_1} \sum_{i=1}^{n} \log p(Y_i | X = X_i) = -\sum_{i=1}^{n} X_i (X_i w_1 + w_0 - Y_i)
\]
\[
\frac{\partial}{\partial w_0} \sum_{i=1}^{n} \log p(Y_i | X = X_i) = -\sum_{i=1}^{n} (X_i w_1 + w_0 - Y_i)
\]

Setting both of the partial derivatives to zero, we immediately get two equations with two unknowns. The terms that only have \(Y_i\) and \(Y_i X_i\) are pulled to the other side of the equality:
\[
w_1 \sum_{i=1}^{n} X_i^2 + w_0 \sum_{i=1}^{n} X_i = \sum_{i=1}^{n} X_i Y_i
\]
\[
w_1 \sum_{i=1}^{n} X_i + w_0 n = \sum_{i=1}^{n} Y_i
\]

Then, these can be solved directly:
\[
w_1 = \frac{n \sum_{i=1}^{n} X_i Y_i - \sum_{i=1}^{n} X_i \sum_{i=1}^{n} Y_i}{n \sum_{i=1}^{n} X_i^2 - (\sum_{i=1}^{n} X_i)^2}
\]
\[
w_0 = \frac{\sum_{i=1}^{n} Y_i \sum_{i=1}^{n} X_i^2 - \sum_{i=1}^{n} X_i \sum_{i=1}^{n} X_i Y_i}{n \sum_{i=1}^{n} X_i^2 - (\sum_{i=1}^{n} X_i)^2}.
\]

To observe the significance of these terms, we now introduce a few notations. Define the sample mean, sample variance (unadjusted) and sample cross-covariance as follows:
\[
\bar{X} = \frac{1}{n} \sum_{i=1}^{n} X_i, \quad \bar{Y} = \frac{1}{n} \sum_{i=1}^{n} Y_i,
\]
\[
\hat{s}_X^2 = \frac{1}{n} \sum_{i=1}^{n} (X_i - \bar{X})^2 = \frac{1}{n} \sum_{i=1}^{n} X_i^2 - \bar{X}^2
\]
\[
\hat{s}_{XY} = \frac{1}{n} \sum_{i=1}^{n} (X_i - \bar{X})(Y_i - \bar{Y}) = \frac{1}{n} \sum_{i=1}^{n} X_i Y_i - \bar{X} \bar{Y},
\]

Then, we have
\[
w_0 = \bar{Y} - \bar{X} w_1, \quad \text{and}
\]
\[
w_1 = \frac{\hat{s}_{XY}}{\hat{s}_X^2}.
\]
(c) Now, consider a different generative model. Let \( Y = Xw + Z \), where \( Z \sim U[-0.5, 0.5] \) is a continuous random variable uniformly distributed between \(-0.5\) and \(0.5\). Again assume that \( w \) is a fixed parameter. **What is the conditional distribution of \( Y \) given \( X \)?**

**Solution:** As before, \( w \) is fixed, so we have

\[
P(Y = y | X = x) = P(z = y - Xw | X = x) = P(z = y - xw) = \begin{cases} 1 & \text{if } -0.5 < y - xw < 0.5 \\ 0 & \text{otherwise} \end{cases}
\]

(3)

(d) Given \( n \) points of training data \( \{(X_1, Y_1), (X_2, Y_2), \cdots, (X_n, Y_n)\} \) generated in an i.i.d. fashion in the setting of the part (c) derive a maximum likelihood estimator of \( w \). Assume that \( X_i > 0 \) for all \( i = 1, \ldots, n \). (Note that MLE for this case need not be unique; but you are required to report only one particular estimate.)

**Solution:** Noting that \( X_i > 0 \), we find that the likelihood function is given by

\[
\Pi_i p(Y_i | X_i) = \Pi_{i=1}^n 1 \left\{ -0.5 + X_i w < Y_i < 0.5 + X_i w \right\} = \Pi_{i=1}^n 1 \left\{ \frac{Y_i - 0.5}{X_i} < w < \frac{Y_i + 0.5}{X_i} \right\}.
\]

For this case, maximizing likelihood is equivalent to ensuring that all the indicators have value 1, which in turn requires that all those regions intersect:

\[
\max_i \left\{ \frac{Y_i - 0.5}{X_i} \right\} < w < \min_i \left\{ \frac{Y_i + 0.5}{X_i} \right\}.
\]

So any value of \( w \) satisfying these constraints is a valid MLE.

(e) Take the model \( Y = Xw + Z \), where \( Z \sim U[-0.5, 0.5] \). Use a computer to simulate \( n \) training samples \( \{(X_1, Y_1), (X_2, Y_2), \cdots, (X_n, Y_n)\} \) and illustrate what the likelihood of the data looks like as a function of \( w \) after \( n = 5, 25, 125, 625 \) training samples. Qualitatively describe what is happening as \( n \) gets large.

(You may use the starter code. Note that you have considerable design freedom in this problem part. You get to choose how you draw the \( X_i \) as well as what true value \( w \) you want to illustrate. You have total freedom in using additional python libraries for this problem part. No restrictions.)

**Solution:** We generated samples assuming \( Y = 3X + Z \). We assumed that the model parameter is between 0 and 4 (you can assume any range that includes the true parameters.) and we calculated the likelihood by using the formula we obtained above for different values of \( w \) in
this interval. To do so, for a specific $w$ we compute the likelihood $\prod_{i=1}^{n} Pr(Y_i|X_i; w)$ which is either zero or 1 because the probability distribution is uniform. In the figure below, likelihood of the data can be seen as a function of $w$ for different sizes of the data. Notice the $y$-axis values in the different plots. We can observe from the plots that increasing the number of samples leads the posterior to concentrate around the true value of $w = 3$ in this case.

The code here uses a simple approach of just giving each training sample a veto into whether or not a particular $w$ is feasible or not.

```python
import numpy as np
import matplotlib.pyplot as plt

sample_size = [5, 25, 125, 625]
plt.figure(figsize=(12, 10))
for k in range(4):
    n = sample_size[k]
    X = 2+2*np.random.random(n)  # generating n X with U[2, 4]
    Z = np.random.uniform(-0.5, 0.5, n)  # generating i.i.d random uniform noise
    Y = 3*X+Z  # computing y from x and z assuming w=3
    N = 1001
    W = np.linspace(0,4,N)
    likelihood = np.ones(N)  # likelihood as a function of w
    for i1 in range(N):
        w = W[i1]
        for i in range(n):
            # Yi has uniform distribution in [wXi-0.5,wXi+0.5]
            if abs(Y[i]-w*X[i]) > 0.5:
                likelihood[i1]=0
    likelihood /= sum(likelihood)
    plt.subplot(2, 2, k+1)
    plt.plot(W, likelihood)
    plt.xlabel('w', fontsize=10)
    plt.legend(['n=' + str(n)], fontsize=14)
    # plt.savefig('uni_noise_post.pdf')  # command to save figure
plt.show()
```

Figure 1: Plot of the likelihood of data as a function of $w$ for different sample size
Now, let us return to the case of Gaussian noise. Given \( n \) points of training data \( \{(X_1, Y_1), (X_2, Y_2), \ldots, (X_n, Y_n)\} \) generated according to \( Y_i = X_i W + Z_i \), where \( Z_i \sim N(0, 1) \) are iid standard normal random variables. Assume \( W \sim N(0, \sigma^2) \) is also a normal random variable and is independent of both the \( Z_i \)'s and the \( X_i \)'s. Use Bayes' Theorem to derive the posterior distribution of \( W \) given the training data. What is the mean of the posterior distribution of \( W \) given the data?

**Hint:** Compute the posterior up-to proportionality and try to identify the distribution by completing the square.

**Solution:** By Bayes' Theorem, we have

\[
P(w|X_1, Y_1, \ldots, X_n, Y_n) \propto \prod_{i=1}^{n} P(Y_i|X_i, w) \cdot P(w)
\]

\[
\propto \prod_{i=1}^{n} \exp \left\{ -\frac{1}{2} (Y_i - w X_i)^2 \right\} \exp \left\{ -0.5 \frac{w^2}{\sigma^2} \right\}
\]

\[
\propto \prod_{i=1}^{n} \exp \left\{ -\frac{1}{2} (Y_i^2 - 2Y_i w X_i + w^2 X_i^2) \right\} \exp \left\{ -0.5 \frac{w^2}{\sigma^2} \right\}
\]

\[
\propto \exp \left\{ -\frac{1}{2} \left( \sum_{i=1}^{n} Y_i^2 - 2 \sum_{i=1}^{n} Y_i w X_i + w^2 \sum_{i=1}^{n} X_i^2 \right) \right\} \exp \left\{ -0.5 \frac{w^2}{\sigma^2} \right\}
\]

\[
\propto \exp \left\{ -\frac{1}{2} \left( \sum_{i=1}^{n} X_i^2 + \frac{1}{\sigma^2} \right) w^2 - \left( \sum_{i=1}^{n} X_i Y_i \right) w \right\} + \text{Constant}
\]

\[
\propto \exp \left\{ -\frac{1}{2} \left( \sum_{i=1}^{n} X_i^2 + \frac{1}{\sigma^2} \right) \right\} w^2 - \left( \sum_{i=1}^{n} X_i Y_i \right) w \right\} + \text{Constant}
\]

\[
\propto \exp \left\{ -\frac{1}{2} \left( \sum_{i=1}^{n} X_i^2 + \frac{1}{\sigma^2} \right) \right\} \left( w - \frac{\sum_{i=1}^{n} X_i Y_i}{\sum_{i=1}^{n} X_i^2 + \frac{1}{\sigma^2}} \right)^2 + \text{Constant}
\]

To get to the last line, we used a trick of completing the square. Consider the term we would need to add to

\[
\left( w^2 - \frac{2 \sum_{i=1}^{n} X_i Y_i}{\sum_{i=1}^{n} X_i^2 + \frac{1}{\sigma^2}} \right)
\]

to get

\[
\left( w - \frac{\sum_{i=1}^{n} X_i Y_i}{\sum_{i=1}^{n} X_i^2 + \frac{1}{\sigma^2}} \right)^2.
\]

The term we would add does not depend on \( w \), only the sample points. From the perspective of the distribution on \( w \), this is just a constant, so we folded it into the constant term.
The final step is to notice that this posterior of $w$ given $X_1, Y_1, \cdots, X_n, Y_n$ is a normal distribution. To see the connection, ignore the constant term in the posterior

$$P(w|X_1, Y_1, \cdots, X_n, Y_n) \propto \exp \left\{ -0.5 \left( \sum_{i=1}^{n} X_i^2 + \frac{1}{\sigma^2} \right) \left( w - \frac{\sum_{i=1}^{n} X_i Y_i}{\sum_{i=1}^{n} X_i^2 + \frac{1}{\sigma^2}} \right)^2 \right\}.$$ 

Recall the density for the normal distribution $N(\mu, \sigma^2)$ is given by

$$p(x) = \frac{1}{\sqrt{2\pi \sigma^2}} \exp \left\{ -\frac{1}{2\sigma^2} (x - \mu)^2 \right\} \propto \exp \left\{ -\frac{1}{2\sigma^2} (x - \mu)^2 \right\}.$$ 

Comparing the two we conclude that

$$W|\{X_i, Y_i, i = 1, \ldots, n\} \sim N \left( \frac{\sum_{i=1}^{n} X_i Y_i}{\sum_{i=1}^{n} X_i^2 + \frac{1}{\sigma^2}}, \frac{1}{\sum_{i=1}^{n} X_i^2 + \frac{1}{\sigma^2}} \right).$$ 

Note that the mean of the posterior is

$$\frac{\sum_{i=1}^{n} X_i Y_i}{\sum_{i=1}^{n} X_i^2 + \frac{1}{\sigma^2}}. \quad (4)$$ 

Since the posterior is Gaussian, we also conclude that the mode of the distribution is the mean and hence the mean is also the maximum a posteriori estimate (MAP) of $W$.

(g) Consider $n$ training data points $\{(x_1, Y_1), (x_2, Y_2), \cdots, (x_n, Y_n)\}$ generated according to $Y_i = \mathbf{w}^\top x_i + Z_i$ where $Y_i \in \mathbb{R}$, $\mathbf{w}, x_i \in \mathbb{R}^d$ with $\mathbf{w}$ fixed, and $Z_i \sim N(0, 1)$ iid standard normal random variables. Argue why the maximum likelihood estimator for $\mathbf{w}$ is the solution to a least squares problem.

**Solution:** Since the logarithm is a monotonically increasing function, we can just look at the log likelihood function.

$$\sum_{i} \log p(Y_i|x_i) = -\frac{1}{2} \sum_{i=1}^{n} (Y_i - (x_i^\top \mathbf{w}))^2 + \text{Constant}$$

$$= -\frac{1}{2} \|\mathbf{Xw} - \mathbf{Y}\|^2 + \text{Constant},$$

with $\mathbf{X} \in \mathbb{R}^{n \times d}$ whose $i^{th}$ row is $x_i^\top$ and $\mathbf{Y} \in \mathbb{R}^n$ whose $i^{th}$ entry is $Y_i$.

At this point, we are done. We have shown that maximizing the maximum-likelihood objective is the same as maximizing a negative squared error objective, which is the same as minimizing the squared error objective, the same as ordinary least squares.

However, if we wanted to, we could just keep solving this.

Differentiate the log likelihood with respect to $\mathbf{w}$ and setting the gradient to zero, we get

$$\nabla_{\mathbf{w}} \sum_{i} \log p(Y_i|x_i) = \mathbf{X}^\top (\mathbf{Xw} - \mathbf{Y}).$$
Setting the gradient to zero, we get

\[ w = (X^TX)^{-1}X^TY. \]

which is the same solution as the least squares problem.

(h) (Multi-dimensional ridge regression) Consider the setup of the previous part: \( Y_i = W^T x_i + Z_i \), where \( Y_i \in \mathbb{R}, W, x_i \in \mathbb{R}^d \), and \( Z_i \sim N(0, 1) \) iid standard normal random variables. Now we treat \( W \) as a random vector and assume a prior knowledge about its distribution. In particular, we use the prior information that the random variables \( W_j \) are i.i.d. \( \sim N(0, \sigma^2) \) for \( j = 1, 2, \ldots, d \). **Derive the posterior distribution of \( W \) given all the \( x_i, Y_i \) pairs. What is the mean of the posterior distribution of the random vector \( W \)?**

Hint: Use hints from part (f) and the following identities: For \( X = \begin{bmatrix} x_1^T \\ \vdots \\ x_n^T \end{bmatrix} \) and \( Y = \begin{bmatrix} Y_1 \\ \vdots \\ Y_n \end{bmatrix} \) we have \( X^TX = \sum_{i=1}^n x_i x_i^T \) and \( X^TY = \sum_{i=1}^n x_i Y_i \).

**Solution:** Note that, we have the following prior distribution on \( W \):

\[
P(w) = \Pi_{j=1}^d P(w_j)
= \Pi_{j=1}^d \frac{1}{\sqrt{2\pi \sigma^2}} \exp \left( -\frac{w_j^2}{2\sigma^2} \right)
= \frac{1}{\sqrt{(2\pi \sigma^2)^d}} \exp \left( -\frac{w^2}{2\sigma^2} \right)
\propto \exp \left\{ -\frac{\|w\|_2^2}{2\sigma^2} \right\}.
\]

Furthermore, to compute the likelihood of the data, we observe that \( Y|X, w \sim N(X^Tw, 1) \). Concretely, we have

\[
p(Y|X, w) \propto \exp \left\{ -\frac{1}{2}\|Y - (X^Tw)\|^2 \right\}.
\]

(5)
Now by Bayes’ Theorem, we have

\[ P(w|x_1, Y_1, \ldots, x_n, Y_n) \propto \prod_{i=1}^n P(Y_i|x_i, w) \prod_{j=1}^d P(w_j) \]

\[ \propto \prod_{i=1}^n \exp \left\{ -0.5 \left( Y_i - x_i^\top w \right)^2 \right\} \exp \left\{ -\frac{\|w\|_2^2}{2\sigma^2} \right\} \]

\[ \propto \prod_{i=1}^n \exp \left\{ -0.5 \left( Y_i^2 - 2Y_i x_i^\top w + (x_i^\top w)^2 \right) \right\} \exp \left\{ -\frac{\|w\|_2^2}{2\sigma^2} \right\} \]

\[ \propto \exp \left\{ -0.5 \left( \sum_{i=1}^n Y_i^2 - 2 \sum_{i=1}^n Y_i x_i^\top w + \sum_{i=1}^n (x_i^\top w)^2 \right) \right\} \exp \left\{ -\frac{\|w\|_2^2}{2\sigma^2} \right\} \]

\[ \propto \exp \left\{ -0.5 \left( \sum_{i=1}^n Y_i^2 - 2 \sum_{i=1}^n Y_i x_i^\top w + \sum_{i=1}^n w^\top x_i x_i^\top w \right) \right\} \exp \left\{ -\frac{\|w\|_2^2}{2\sigma^2} \right\} \]

\[ \propto \exp \left\{ -0.5 \left( \sum_{i=1}^n Y_i^2 - 2 \sum_{i=1}^n Y_i x_i^\top w + w^\top \sum_{i=1}^n x_i x_i^\top w \right) \right\} \exp \left\{ -\frac{\|w\|_2^2}{2\sigma^2} \right\} \]

\[ \propto \exp \left\{ -0.5 \left( \sum_{i=1}^n (x_i Y_i)^\top w + w^\top \sum_{i=1}^n x_i x_i^\top w \right) \right\} \exp \left\{ -\frac{\|w\|_2^2}{2\sigma^2} \right\} \]

\[ \propto \exp \left\{ -\frac{1}{2} \left[ w^\top (X^\top X + \frac{1}{\sigma^2} I_d) w - 2 \left( X^\top Y \right)^\top w \right] \right\}, \]  

(6)

with \( X \in \mathbb{R}^{n \times d} \) whose \( i \th \) row is \( x_i^\top \) and \( Y \in \mathbb{R}^n \) whose \( i \th \) entry is \( Y_i \). The last line uses the fact that

\[ X^\top X = \sum_{i=1}^n x_i x_i^\top \]

and

\[ X^\top Y = \sum_{i=1}^n x_i Y_i. \]

Now, as we did earlier, we complete the square, except in the matrix case. To make the notation easier, let’s define the matrix \( M \) as

\[ M = (X^\top X + \frac{1}{\sigma^2} I_d). \]

Then, noticing that \( M \) is symmetric, we see that:

\[ P(w|x_1, Y_1, \ldots, x_n, Y_n) \propto \exp \{-\frac{1}{2} [w^\top M w - 2(X^\top Y)^\top w] \} \]

\[ \propto \exp \{-\frac{1}{2} [(w - M^{-1}(X^\top Y))^\top M(w - M^{-1}(X^\top Y))] + \text{Constant} \} \]  

(7)
How did we know this was how we should complete the square? The \( w \) part on the sides is clear as is the central \( M \) since we knew we wanted the \( w^\top Mw \) term. Now, because we knew that we wanted to get an \( (X^\top Y)^\top w \) term as the linear term in the square, we had to cancel the \( M \) in the middle with an \( M^{-1} \). Of course, as argued earlier, the constant here absorbs the compensation for the term that we added while completing the square.

Therefore, the posterior of \( w \) given \( x_1, Y_1, \cdots, x_n, Y_n \) is a multivariate Gaussian. The mean of this multivariate Gaussian is

\[
(X^\top X + \frac{1}{\sigma^2} I_d)^{-1} X^\top Y.
\]

(i) Consider \( d = 2 \) and the setting of the previous part. Use a computer to simulate and illustrate what the a-posteriori probability looks like for the \( W \) model parameter space after \( n = 5, 25, 125 \) training samples for different values of \( \sigma^2 \).

(Again, you may use the starter code. And like problem (e), there are no restrictions for using additional python libraries for this part as well.)

**Solution:** First, we assumed a variance for the random variable \( W \) and we generate \( d \) numbers from \( N(0, \sigma^2) \). Then, we generated the samples using these parameters and computed a value proportional to the probability using Equation 6 from above. We could have normalized to get the exact posterior probability; our results would have been equivalent as far the plots would go.

In the figures below, the a-posteriori probability of the model parameters given data can be seen in the model space \( W = (W_1, W_2) \). As we can see in the plots, increasing the number of samples results in shrinking the zone of likely parameters.

![Heatplots of the a-posteriori probability of the model given the training data, plotted in the model space \( (W_1, W_2) \) for different sample sizes with \( \sigma = 0.5 \)](image1)

![Heatplots of the a-posteriori probability of the model given the training data, plotted in the model space \( (W_1, W_2) \) for different sample sizes with \( \sigma = 1.5 \)](image2)
import numpy as np
import matplotlib.pyplot as plt

Samples = [5, 25, 125]

Sigma = [0.5**2, 1.5**2]  # plotting for two different variance
w1_s = [0.4, 1.3]  # true W1 drawn from the gaussian with s.d 0.5 and 1.5 respectively
w2_s = [0.5, -1.7]  # true W2 drawn from the gaussian with s.d 0.5 and 1.5 respectively

for s in range(2):
    sigma = Sigma[s]
    A = w1_s[s]
    B = w2_s[s]
    plt.figure()
    for count in range(3):
        n = Samples[count]
        X = (X1, X2) Y = AX1 + AX2 + Z
        X1 = np.random.normal(0, 1, n)  # generating random samples for X1 from normal dist
        X2 = np.random.normal(0, 1, n)
        Z = np.random.normal(0, 1, n)
        Y = A * X1 + B * X2 + Z
        N = 201
        W = np.linspace(-5, 5, N)
        prob = np.ones([N, N])
        for i1 in range(N):
            w1 = W[i1]  # taking different values for w1 from -5 to 5 to compute something proportional to the posterior probability
            for i2 in range(N):
                w2 = W[i2]
                L = 1
                for i in range(n):
                    L = L * np.exp(-0.5 * (Y[i] - X1[i] * w1 - X2[i] * w2) ** 2)
                L = L * np.exp(-0.5 * (w1**2 + w2**2) / sigma)
                prob[i1][i2] = L
        plt.subplot(1, 3, count + 1)
        plt.imshow(np.flipud(prob), cmap='hot', aspect='auto', extent=[-5, 5, -5, 5])
        plt.show()

3 Simple Bias-Variance Tradeoff

Consider a random variable $X$, which has unknown mean $\mu$ and unknown variance $\sigma^2$. Given $n$ iid realizations of training samples $X_1 = x_1, X_2 = x_2, \ldots, X_n = x_n$ from the random variable, we wish to estimate the mean of $X$. We will call our estimate of $X$ the random variable $\hat{X}$, which has mean $\hat{\mu}$. There are a few ways we can estimate $\mu$ given the realizations of the $n$ samples:

1. Average the $n$ samples: $\frac{x_1 + x_2 + \ldots + x_n}{n}$.
2. Average the $n$ samples and one sample of 0: $\frac{x_1 + x_2 + \ldots + x_n}{n + 1}$.
3. Average the $n$ samples and $n_0$ samples of 0: $\frac{x_1 + x_2 + \ldots + x_n}{n + n_0}$.
4. Ignore the samples: just return 0.

In the parts of this question, we will measure the bias and variance of each of our estimators. The bias is defined as

$$E[\hat{X} - \mu]$$

and the variance is defined as

$$\text{Var}[\hat{X}]$$.
(a) **What is the bias of each of the four estimators above?**

**Solution:** Using the linearity of expectation, we write $E[\hat{X} - X]$ as $E[\hat{X}] - E[X] = E[\hat{X}] - \mu$, so we have the following biases:

(a) $E[\hat{X}] = E[\frac{X_1 + X_2 + \ldots + X_n}{n}] = \frac{n\mu}{n} \implies \text{bias} = 0$

(b) $E[\hat{X}] = E[\frac{X_1 + X_2 + \ldots + X_{n+1}}{n+1}] = \frac{n\mu}{n+1} \implies \text{bias} = -\frac{1}{n+1}\mu$

(c) $E[\hat{X}] = E[\frac{X_1 + X_2 + \ldots + X_{n+n_0}}{n+n_0}] = \frac{n\mu}{n+n_0} \implies \text{bias} = -\frac{n_0}{n+n_0}\mu$

(d) $E[\hat{X}] = 0 \implies \text{bias} = -\mu$

(b) **What is the variance of each of the four estimators above?**

**Solution:** The two key identities to remember are $\text{Var}[A + B] = \text{Var}[A] + \text{Var}[B]$ (when $A$ and $B$ are independent) and $\text{Var}[kA] = k^2\text{Var}[A]$, where $A$ and $B$ are random variables and $k$ is a constant.

(a) $\text{Var}[\hat{X}] = \text{Var}[\frac{X_1 + X_2 + \ldots + X_n}{n}] = \frac{1}{n^2}\text{Var}[X_1 + X_2 + \ldots + X_n] = \frac{1}{n^2}(n\sigma^2) = \frac{\sigma^2}{n}$

(b) $\text{Var}[\hat{X}] = \text{Var}[\frac{X_1 + X_2 + \ldots + X_{n+1}}{n+1}] = \frac{1}{(n+1)^2}\text{Var}[X_1 + X_2 + \ldots + X_n] = \frac{1}{(n+1)^2}(n\sigma^2) = \frac{n\sigma^2}{(n+1)^2}$

(c) $\text{Var}[\hat{X}] = \text{Var}[\frac{X_1 + X_2 + \ldots + X_{n+n_0}}{n+n_0}] = \frac{1}{(n+n_0)^2}\text{Var}[X_1 + X_2 + \ldots + X_n] = \frac{1}{(n+n_0)^2}(n\sigma^2) = \frac{n\sigma^2}{(n+n_0)^2}$

(d) $\text{Var}[\hat{X}] = 0$

(c) Suppose we have constructed an estimator $\hat{X}$ from some samples of $X$. We now want to know how well $\hat{X}$ estimates a fresh (new) sample of $X$. Denote this fresh sample by $X'$. Note that $X'$ is an i.i.d. copy of the random variable $X$. **Derive a general expression for the expected squared error** $E[(\hat{X} - X')^2]$ **in terms of** $\sigma^2$ **and the bias and variance of the estimator** $\hat{X}$. Similarly, derive an expression for the expected squared error $E[(\hat{X} - \mu)^2]$. Compare the two expressions and comment on the differences between them, if any.

**Solution:** Since $\hat{X}$ is a function of $X$, we conclude that the random variables $\hat{X}$ and $X'$ are independent of each other. Now we provide two ways to solve the first problem.

**Method 1:** In this method, we use the trick of adding and subtracting a term to derive the desired expression:

$$
E[(\hat{X} - X')^2] = E[(\hat{X} - \mu + \mu - X')^2]
$$

$$
= E[(\hat{X} - \mu)^2 + E[(\mu - X')]^2]
$$

$$
= E[(\hat{X} - \mu)^2] + \sigma^2
$$

$$
= E[(\hat{X} - E[\hat{X}]) + E[\hat{X}] - \mu)^2] + \sigma^2
$$

$$
= E[(\hat{X} - E[\hat{X}])^2] + (E[\hat{X}] - \mu)^2 + 2E[(\hat{X} - E[\hat{X}]) \cdot (E[\hat{X}] - \mu)] + \sigma^2
$$

$$
= \text{Var}(\hat{X}) + \text{bias}^2 + 2(\text{bias})^2 + \sigma^2
$$

$$
= \text{Var}(\hat{X}) + 2(\text{bias})^2 + \sigma^2
$$

The two key identities to remember are $\text{Var}[A + B] = \text{Var}[A] + \text{Var}[B]$ (when $A$ and $B$ are independent) and $\text{Var}[kA] = k^2\text{Var}[A]$, where $A$ and $B$ are random variables and $k$ is a constant.
Method 2: In this method, we make use of the definition of variance. We have

\[
E[(\hat{X} - X')^2] = E[\hat{X}^2] + E[X'^2] - 2E[\hat{X}X']
\]

\[
= (\text{Var}(\hat{X}) + (E[\hat{X}])^2) + (\text{Var}(X') + (E[X'])^2) - 2E[\hat{X}X']
\]

\[
= ((E[\hat{X}])^2 - 2E[\hat{X}X'] + (E[X'])^2) + \text{Var}(\hat{X}) + \text{Var}(X')
\]

\[
= (E[\hat{X}] - E[X'])^2 + \text{Var}(\hat{X}) + \text{Var}(X)
\]

\[
= (E[\hat{X}] - \mu)^2 + \text{Var}(\hat{X}) + \sigma^2
\]

The first term is equivalent to the bias of our estimator squared, the second term is the variance of the estimator, and the last term is the irreducible error.

Now let’s do \(E[(\hat{X} - \mu)^2]\).

\[
E[(\hat{X} - \mu)^2] = E[\hat{X}^2] + E[\mu^2] - 2E[\hat{X}\mu]
\]

\[
= (\text{Var}(\hat{X}) + (E[\hat{X}])^2) + \text{Var}(\mu) + (E[\mu])^2 - 2E[\hat{X}\mu]
\]

\[
= (E[\hat{X}]^2 - 2E[\hat{X}\mu] + (E[\mu])^2) + \text{Var}(\hat{X}) + \text{Var}(\mu)
\]

\[
= (E[\hat{X}] - E[\mu])^2 + \text{Var}(\hat{X}) + \text{Var}(\mu)
\]

\[
= (E[\hat{X}] - \mu)^2 + \text{Var}(\hat{X}).
\]

Notice that these two expected squared errors resulted in the same expressions except for the \(\sigma^2\) in \(E[(\hat{X} - X')^2]\). The error \(\sigma^2\) is considered “irreducible error” because it is associated with the noise that comes from sampling from the distribution of \(X\). This term is not present in the second derivation because \(\mu\) is a fixed value that we are trying to estimate.

(d) For the following parts, we will refer to expected total error as \(E[(\hat{X} - \mu)^2]\). It is a common mistake to assume that an unbiased estimator is always “best.” Let’s explore this a bit further. Compute the expected squared error for each of the estimators above. Solution: Adding the previous two answers:

(a) \(\frac{\sigma^2}{n}\)

(b) \(\frac{1}{(n+1)^2} (\mu^2 + n\sigma^2)\)

(c) \(\frac{1}{(n+n_0)^2} (\mu_0^2 + n\sigma^2)\)

(d) \(\mu^2\)

(e) Demonstrate that the four estimators are each just special cases of the third estimator, but with different instantiations of the hyperparameter \(n_0\).

Solution: The derivation for the third estimator works for any value of \(n_0\). The first estimator is just the third estimator with \(n_0\) set to 0:

\[
\frac{x_1 + x_2 + \ldots + x_n}{n + n_0} = \frac{x_1 + x_2 + \ldots + x_n}{n + 0} + \frac{x_1 + x_2 + \ldots + x_n}{n}.
\]
The second estimator is just the third estimator with \( n_0 \) set to 1:

\[
\frac{x_1 + x_2 + \ldots + x_n}{n + n_0} = \frac{x_1 + x_2 + \ldots + x_n}{n + 1}.
\]

The last estimator is the limiting behavior as \( n_0 \) goes to \( \infty \). In other words, we can get arbitrarily close to the fourth estimator by setting \( n_0 \) very large:

\[
\lim_{n_0 \to \infty} \frac{x_1 + x_2 + \ldots + x_n}{n + n_0} = 0.
\]

(f) **What happens to bias as \( n_0 \) increases? What happens to variance as \( n_0 \) increases?**

**Solution:**

One reason for increasing the samples of \( n_0 \) is if you have reason to believe that \( X \) is centered around 0. In increasing the number of zeros we are injecting more confidence in our belief that the distribution is centered around zero. Consequently, in increasing the number of ”fake” data, the variance decreases because your distribution becomes more peaked. Examining the expressions for bias and variance for the third estimator, we can see that larger values of \( n_0 \) result in decreasing variance \( (\frac{n}{(n+n_0)^2}\sigma^2) \) but potentially increasing bias \( (\frac{n_0\mu}{n+n_0}) \). Hopefully you can see that there is a trade-off between bias and variance. Using an unbiased estimator is not always optimal nor is using an estimator with small variance always optimal. One has to carefully trade-off the two terms in order to obtain minimum squared error.

(g) Say that \( n_0 = \alpha n \). **Find the setting for \( \alpha \) that would minimize the expected total error, assuming you secretly knew \( \mu \) and \( \sigma \).** Your answer will depend on \( \sigma \), \( \mu \), and \( n \).

**Solution:** First, we write our expression for the total error in terms of \( \alpha \):

\[
\frac{1}{(n+n_0)^2}(n_0^2\mu^2 + n\sigma^2)
\]

\[
\frac{1}{(n+\alpha n)^2}((\alpha n)^2\mu^2 + n\sigma^2)
\]

\[
\frac{1}{(1+\alpha)^2}\frac{1}{n^2}(\alpha^2n^2\mu^2 + n\sigma^2)
\]

\[
\frac{1}{(1+\alpha)^2}(\alpha^2\mu^2 + \frac{\sigma^2}{n})
\]

\[
\frac{\alpha^2}{(1+\alpha)^2}\mu^2 + \frac{1}{(1+\alpha)^2} \frac{\sigma^2}{n}
\]

Now take the derivative with respect to \( \alpha \) and set it equal to 0:

\[
\frac{2\alpha}{(1+\alpha)^3}\mu^2 - \frac{2}{(1+\alpha)^3} \frac{\sigma^2}{n} = 0.
\]
\[
\frac{2\alpha}{(1 + \alpha)^3} \mu^2 = \frac{2}{(1 + \alpha)^3} \frac{\sigma^2}{n}
\]
\[
2\alpha \mu^2 = 2 \frac{\sigma^2}{n}
\]
\[
\alpha = \frac{\sigma^2}{n \mu^2}.
\]

(h) For this part, let’s assume that we had some reason to believe that \(\mu \) should be small (close to 0) and \(\sigma \) should be large. In this case, what happens to the expression in the previous part? **Solution:** The value of \(\alpha \) can be quite large, since the solution has a small value of \(\mu \) in the denominator. In mathematical terms, we could write the limit as \(\mu \) goes to 0:
\[
\lim_{\mu \to 0} \frac{\sigma^2}{n \mu^2} = \infty.
\]

(i) In the previous part, we assumed there was reason to believe that \(\mu \) should be small. Now let’s assume that we have reason to believe that \(\mu \) is not necessarily small, but should be close to some fixed value \(\mu_0\). In terms of \(X \) and \(\mu_0\), how can we define a new random variable \(X' \) such that \(X' \) is expected to have a small mean? Compute the mean and variance of this new random variable. **Solution:** Shift the random variable \(X \) by the constant guess \(\mu_0 \) to get the random variable \(X' = X - \mu_0 \). Let’s calculate the mean and variance of this new random variable:
\[
E[X'] = E[X - \mu_0] = E[X] - \mu_0 = \mu - \mu_0 \approx 0.
\]
The last line \((\mu - \mu_0 \approx 0)\) comes from the assumption that \(\mu \) is close to \(\mu_0 \).

We can also calculate the variance of \(X' \):
\[
\text{Var}[X'] = \text{Var}[X - \mu_0] = \text{Var}[X] - \text{Var}[\mu_0] = \text{Var}[X] - 0 = \text{Var}[X].
\]

This is a useful step to understand the relation between \(X \) and \(X' \), but not necessary for a full solution to the question asked.

(j) Draw a connection between \(\alpha \) in this problem and the regularization parameter \(\lambda \) in the ridge-regression version of least-squares. **What does this problem suggest about choosing a regularization coefficient and handling our data-sets so that regularization is most effective?** This is an open-ended question, so do not get too hung up on it. **Solution:** The key lesson is another reminder that regularization reduces variance, at the cost of increasing bias, by forcing solutions towards zero. But the bias-variance trade-off is not always the same. If we first center our data around some prior, so that the model is supposed to be close to zero anyways, then we can use larger values of \(\alpha \) or \(\lambda \) and reduce variance considerably for a small cost in bias. It may also be instructive to realize that regularization can be thought of as adding “fake” training data which is uniformly zero.
Robotic Learning of Controls from Demonstrations and Images

Huey, a home robot, is learning to retrieve objects from a cupboard, as shown in Fig. 4. The goal is to push obstacle objects out of the way to expose a goal object. Huey’s robot trainer, Anne, provides demonstrations via tele-operation. When tele-operating the robot, Anne can look at the images captured by the robot and provide controls to Huey remotely.

During a demonstration, Huey records the RGB images of the scene for each timestep, \(x_0, x_1, \ldots, x_n\), where \(x_i \in \mathbb{R}^{30 \times 30 \times 3}\) and the controls for his body, \(u_0, u_1, \ldots, u_n\), where \(u_i \in \mathbb{R}^{3}\). The controls correspond to making small changes in the 3D pose (i.e. translation and rotation) of his body. Examples of the data are shown in the figure.

Under an assumption (sometimes called the Markovian assumption) that all that matters for the current control is the current image, Huey can try to learn a linear policy \(\pi\) (where \(\pi \in \mathbb{R}^{2700 \times 3}\)) which linearly maps image states to controls (i.e. \(\pi^T x = u\)). We will now explore how Huey can recover this policy using linear regression. Note please use numpy and numpy.linalg to complete this assignment.

Figure 4: A) Huey trying to retrieve a mustard bottle. An example RGB image of the workspace taken from his head mounted camera is shown in the orange box. The angle of the view gives Huey and eye-in-hand perspective of the cupboard he is reaching into. B) A scatter plot of the 3D control vectors, or \(u\) labels. Notice that each coordinate of the label lies within the range of \([-1, 1]\) for the change in position. Example images, or states \(x\), are shown for some of the corresponding control points. The correspondence is indicated by the blue lines.

(a) To get familiar with the structure of the data, please visualize the 0th, 10th and 20th images in the training dataset. Also find out what’s their corresponding control vectors.

Solution: The corresponding controls are \([0, -1, 0]\), \([-1, -0.45111084, -1]\), \([0, 0, 0.37368774]\)

(b) Load the \(n\) training examples from \(x\_train.p\) and compose the matrix \(X\), where \(X \in \mathbb{R}^{n \times 2700}\). Note, you will need to flatten the images to reduce them to a single vector. The flattened image
Figure 5: The 0th, 10th and 20th images in the training set.

vector will be denoted by $\bar{x}$ (where $\bar{x} \in \mathbb{R}^{2700 \times 1}$). Next, load the $n$ examples from $y_{\text{train}.p}$ and compose the matrix $U$, where $U \in \mathbb{R}^{n \times 3}$. Try to perform ordinary least squares to solve:

$$\min_{\pi} \|X\pi - U\|_F$$

to learn the policy $\pi \in \mathbb{R}^{2700 \times 3}$. Report what happens as you attempt to do this and explain why.

**Solution:** The matrix is singular and not invertible. The reason is there isn’t enough data to cover the high dimensional image space, so many solutions exist to solve the problem. Specifically, we only train the policy on 91 images, however a single RGB image is 30x30x3 in dimensionality. So, with that many parameters, we can basically fit any arbitrary set of controls. (This is called the temptation to memorize in machine learning.) We can use ridge regression though to help condition the optimization to favor solutions with a small $\ell_2$ norm.

(c) Now try to perform ridge regression:

$$\min_{\pi} \|X\pi - U\|_2^2 + \lambda \|\pi\|_2^2$$

on the dataset for regularization values $\lambda = \{0.1, 1.0, 10, 100, 1000\}$. Measure the average squared Euclidean distance for the accuracy of the policy on the training data:

$$\frac{1}{n} \sum_{i=0}^{n-1} \|\bar{x}_i^T\pi - u_i\|_2^2$$

Report the training error results for each value of $\lambda$.

**Solution:**

The learned policy should match the training data with very low error. For all values of $\lambda$, in the specified range, you should see the error is below $10^{-8}$, which is a very good fit for the dataset. It should be noted that the ability to fit training data perfectly does not necessarily
mean the robot will perform well on unseen data. Even with the ridge penalties set to these values, the policy has apparently just memorized the controls.

Why was it able to do this? Remember, all the controls are small numbers between $-1$ and $1$ while the training image data has large numbers. Consequently, small numbers for the parameters will allow these controls to be recovered. Furthermore, there are so many parameters that the work of memorizing the controls can be distributed across them. This further shrinks the parameters required to do this memorization. The ridge penalties are simply incapable of discouraging this memorization.

(d) Next, we are going to try standardizing the states. For each pixel value in each data point, $x$, perform the following operation:

$$x \mapsto \frac{x}{255} \times 2 - 1.$$  

Since we know the maximum pixel value is 255, this rescales the data to be between $[-1, 1]$.

Repeat the previous part and report the average squared training error for each value of $\lambda$.

**Solution:**

The answers for the fitting error for $\lambda = \{0.1, 1.0, 10, 100, 1000\}$ are $\{0.000, 0.000, 0.0016, 0.035, 0.25\}$.

With standardization applied, we see as the regularization term is decreased the training loss is lowered. This can be interpreted as the variance of the model is increased as the possible function class is expanded.

We can also see that indeed having the smaller values in the images is forcing the system to use bigger values for the parameters if it wants to memorize the controls. The regularization penalty is now able to begin to discourage this.

(e) Evaluate both policies (i.e. with and without standardization on the new validation data $x_{test.p}$ and $y_{test.p}$) for the different values of $\lambda$. Report the average squared Euclidean loss and qualitatively explain how changing the values of $\lambda$ affects the performance in terms of bias and variance.

**Solution:**

The answer is for $\lambda = \{0.1, 1.0, 10, 100, 1000\}$ is $\{0.87, 0.86, 0.83, 0.72, 0.73\}$ for with standardization and $\{0.77, 0.77, 0.77, 0.77, 0.77\}$ for with out.

The results with standardization illustrate that because the state space is so high dimensional the policy has trouble generalizing, thus adding bias (i.e. increasing $\lambda$ can help generalization). However, increasing it too much can lead to worst performance.

We empirically see that without standardization the test error is higher, in the next section we will examine how we can characterize this.

It should be noted that the ability of Huey to generalize is still quite inadequate for a real robot policy. Later, in the course we will explore how to get significantly lower error on a larger dataset using convolutional neural networks.
(f) To better understand how standardizing improved the loss function, we are going to evaluate the condition number $\kappa$ of the optimization, which is defined as

$$\kappa = \frac{\sigma_{\max}(X^T X + \lambda I)}{\sigma_{\min}(X^T X + \lambda I)}$$

or the ratio of the maximum singular value to the minimum singular value of the relevant matrix. Roughly speaking, the condition number of the optimization process measures how stable the solution will be when some error exists in the observations. More precisely, given a linear system $Ax = b$, the condition number of the matrix $A$ is the maximum ratio of the relative error in the solution $x$ to the relative error of $b$.

For the regularization value of $\lambda = 100$, report the condition number with the standardization technique applied and without.

**Solution:** The condition number without standardization is $\kappa = 52711697.6679$ and with standardization is $\kappa = 444.725931711$. By standardizing our data, we are able to significantly reduce the ratio of the eigenvalues, which makes our optimization less sensitive to noise in the data when performing matrix inversion.

5 Jaina and her giant peaches

Make sure to submit the code you write in this problem to “HW2 Code” on Gradescope.

In another alternative universe, Jaina is a mage testing how long she can fly a collection of giant peaches. She has $n$ training peaches – with masses given by $x_1, x_2, \ldots, x_n$ – and flies these peaches once to collect training data. The experimental flight time of peach $i$ is given by $y_i$. She believes that the flight time is well approximated by a polynomial function of the mass

$$y_i \approx w_0 + w_1 x_i + w_2 x_i^2 + \cdots + w_D x_i^D$$

where her goal is to fit a polynomial of degree $D$ to this data. Include all text responses and plots in your write-up.

(a) **Show how Jaina’s problem can be formulated as a linear regression problem.**

**Solution:** The problem is to find the coefficients $w_d$ such that the squared error is minimized:

$$\min_{w_0, \ldots, w_D} \sum_i (w_0 + w_1 x_i + w_2 x_i^2 + \cdots + w_D x_i^D - y_i)^2$$

Assume that we construct the following matrix:

$$X = \begin{bmatrix} 1 & x_1 & x_1^2 & \cdots & x_1^D \\ 1 & x_2 & x_2^2 & \cdots & x_2^D \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 1 & x_n & x_n^2 & \cdots & x_n^D \end{bmatrix}$$
Then the problem can be written as:

\[
\min_w \| Xw - y \|^2
\]

(b) You are given data of the masses \( \{x_i\}_{i=1}^n \) and flying times \( \{y_i\}_{i=1}^n \) in the “x_train” and “y_train” keys of the file 1D_poly.mat with the masses centered and normalized to lie in the range \([-1, 1]\). Write a script to do a least-squares fit (taking care to include a constant term) of a polynomial function of degree \( D \) to the data. Letting \( f_D \) denote the fitted polynomial, plot the average training error \( R(D) = \frac{1}{n} \sum_{i=1}^n (y_i - f_D(x_i))^2 \) against \( D \) in the range \( D \in \{0, 1, 2, 3, \ldots, n-1\} \). You may not use any library other than numpy and numpy.linalg for computation.

**Solution:**

See Figure for the plot.

```python
#!/usr/bin/env python3
import matplotlib.pyplot as plt
import numpy as np
import scipy.io as spio

# There is numpy.linalg.lstsq, which you should use outside of this class
def lstsq(A, b):
    return np.linalg.solve(A.T @ A, A.T @ b)

def main():
    data = spio.loadmat('1D_poly.mat', squeeze_me=True)
    x_train = np.array(data['x_train'])
    y_train = np.array(data['y_train']).T
    
    n = 20  # max degree
    err = np.zeros(n - 1)
    
    # fill in err
    for d in range(n - 1):
        D = d + 1
        for i in range(D + 1):
            if i == 0:
                Xf = np.array([1] * x_train.size)
            else:
                Xf = np.vstack([np.power(x_train, i), Xf]).T
            
            w = lstsq(Xf, y_train)
            y_predicted = Xf @ w
            err[d] = (np.linalg.norm(y_train - y_predicted)**2) / n

    plt.plot(err)
    plt.xlabel('Degree of Polynomial')
    plt.ylabel('Training Error')
    plt.show()

if __name__ == '__main__':
    main()
```

1A early version of this homework uses \( D \in [1, n-3] \).
(c) **How does the average training error behave as a function of \(D\), and why?** What happens if you try to fit a polynomial of degree \(n\) with a standard matrix inversion method?

**Solution:** The training error decreases since we have more degrees of freedom to fit the dataset. If we try to fit a polynomial of degree \(n - 1\), we will have enough parameters to fit the data exactly, so the training error will be zero. Using a polynomial of degree \(n\) results in a non-invertible data matrix, and so we cannot use a standard matrix inversion method to find our solution.

(d) Jaina has taken Mystical Learning 189, and so decides that she needs to run another experiment before deciding that her prediction is true. She runs another fresh experiment of flight times using the same peaches, to obtain the data with key “y_fresh” in 1D.poly.mat. Denoting the fresh flight time of peach \(i\) by \(\tilde{y}_i\), **plot the average error** \(\tilde{R}(D) = \frac{1}{n} \sum_{i=1}^{n} (\tilde{y}_i - f_D(x_i))^2\) **for the same values of \(D\) as in part (b) using the polynomial approximations \(f_D\) also from the previous part.** How does this plot differ from the plot in (b) and why?

**Solution:** The plots are shown in Figure 7. The plots are different from the training errors since the data was generated afresh. While increasing the polynomial degree served to fit the noise in the training data, we cannot hope to fit noise by using the same model for fresh data. Hence, our performance degrades as we increase polynomial degree, with a minimum seen at the true model order.
def main():
    data = spio.loadmat('1D_poly.mat', squeeze_me=True)
    x_train = np.array(data['x_train'])
    y_train = np.array(data['y_train']).T
    y_fresh = np.array(data['y_fresh']).T

    n = 20  # max degree
    err_train = np.zeros(n - 1)
    err_fresh = np.zeros(n - 1)

    # fill in err_fresh and err_train
    for d in range(n - 1):
        D = d + 1
        for i in range(D + 1):
            if i == 0:
                Xf = np.array([1] * n)
            else:
                Xf = np.vstack([np.power(x_train, i), Xf])
        Xf = Xf.T

        w = lstsq(Xf, y_train)
        err_train[d] = (np.linalg.norm(y_train - Xf @ w)**2) / n
        err_fresh[d] = (np.linalg.norm(y_fresh - Xf @ w)**2) / n

plt.figure()
plt.ylim([0, 6])
plt.plot(err_train, label='train')
plt.plot(err_fresh, label='fresh')
plt.legend()
plt.show()

if __name__ == '__main__':
    main()

Figure 7: Fresh Error yfresh: Result for 1D_poly.mat

(e) How do you propose using the two plots from parts (b) and (d) to “select” the right polynomial model for Jaina?
**Solution:** The right model is the one that minimizes the error in the fresh dataset. The minimizer is at $D = 4$.

(f) Jaina has a new hypothesis – the flying time is actually a function of the mass, smoothness, size, and sweetness of the peach, and some multivariate polynomial function of all of these parameters. A $D$-multivariate polynomial function looks like

$$f_D(x) = \sum_j \alpha_j \prod_i x_i^{p_{ji}},$$

where $\forall j : \sum_i p_{ji} \leq D$. Here $\alpha_j$ is the scale constant for $j$th term and $p_{ji}$ is the exponent of $x_i$ in $j$th term. The data in polynomial_regression_samples.mat ($100000 \times 5$) with columns corresponding to the 5 attributes of the peach. **Use 4-fold cross-validation to decide which of $D \in \{0, 1, 2, 3, 4, 5, 6\}$ is the best fit for the data provided.** For this part, compute the polynomial coefficients via ridge regression with penalty $\lambda = 0.1$, instead of ordinary least squares. You are not allowed to use any library other than numpy and numpy.linalg.

**Solution:** Please refer to the next part for the general implementation.

(g) Now **redo the previous part, but use 4-fold cross-validation on all combinations of $D \in \{1, 2, 3, 4, 5, 6\}$ and $\lambda \in \{0.05, 0.1, 0.15, 0.2\}$** - this is referred to as a grid search. **Find the best $D$ and $\lambda$ that best explains the data using ridge regression.** Print the average training/validation error per sample for all $D$ and $\lambda$.

**Solution:** Minimum of cross validation error happens at $D = 4$ and $\lambda = 0.15$.

```
Average train error:
[[0.05856762013 0.05856762066 0.05856762225 0.05856762491 0.05856762862]
 [0.05848920002 0.05848948229 0.0584902034 0.05849122 0.05849243261]
 [0.05846063295 0.05848702853 0.05848893458 0.05849036454 0.05849178744]
 [0.05839260718 0.05848700898 0.05848892445 0.05849035741 0.05849178171]
 [0.05831118703 0.05848700861 0.05848892426 0.05849035728 0.05849178161]
 [0.05815215946 0.0584870086 0.05848892426 0.05849035728 0.05849178161]]

Average valid error:
[[0.0585748661 0.05857468536 0.0585746856 0.0585748691 0.05857468927]
 [0.05852250667 0.05852112813 0.05852038752 0.05852010745 0.05852016181]
 [0.05854617135 0.0585210268 0.05852032221 0.05852006042 0.0585201252 ]
 [0.05860046897 0.05852102354 0.05852032035 0.05852005896 0.05852012386]
 [0.05869725681 0.05852102357 0.05852032036 0.05852005896 0.05852012386]
 [0.05887534948 0.05852102357 0.05852032036 0.05852005896 0.05852012386]]
```

In the following implementation, we run cross-validation to find errors for each $D$ separately, by varying the value of $\lambda$. This is purely for pedagogical purposes. You can combine these two steps by running through $D$ in an outer loop. Note that we can collect all of these errors, and then choose the model that minimizes the cross-validation error.

```python
import matplotlib.pyplot as plt
import numpy as np
import scipy.io as spio

data = spio.loadmat('polynomial_regression_samples.mat', squeeze_me=True)
data_x = data['x']
```
data_y = data['y']
Kc = 4  # 4-fold cross validation
KD = 6  # max D = 6
LAMBDA = [0, 0.05, 0.1, 0.15, 0.2]

feat_x = 0

def lstsq(A, b, lambda_=0):
    return np.linalg.solve(A.T @ A + lambda_ * np.eye(A.shape[1]), A.T @ b)

def assemble_feature(x, D):
    n_feature = x.shape[1]
    Q = [(np.ones(x.shape[0]), 0, 0)]
    i = 0
    while Q[i][1] < D:
        cx, degree, last_index = Q[i]
        for j in range(last_index, n_feature):
            Q.append((cx * x[:, j], degree + 1, j))
        i += 1
    return np.column_stack([q[0] for q in Q])

def fit(D, lambda_):
    Ns = int(data_x.shape[0] * (Kc - 1) / Kc)  # training
    Nv = int(Ns / (Kc - 1))  # validation
    Etrain = np.zeros(4)
    Evalid = np.zeros(4)
    for c in range(4):
        valid_x = feat_x[c * Nv:(c + 1) * Nv]
        valid_y = data_y[c * Nv:(c + 1) * Nv]
        train_x = np.delete(feat_x, list(range(c * Nv, (c + 1) * Nv)), axis=0)
        train_y = np.delete(data_y, list(range(c * Nv, (c + 1) * Nv)))
        w = lstsq(train_x, train_y, lambda_=lambda_)
        Etrain[c] = np.mean((train_y - train_x @ w)**2)
        Evalid[c] = np.mean((valid_y - valid_x @ w)**2)
    return np.mean(Etrain), np.mean(Evalid)

def main():
    np.set_printoptions(precision=11)
    Etrain = np.zeros((KD, len(LAMBDA)))
    Evalid = np.zeros((KD, len(LAMBDA)))
    for D in range(KD):
        global feat_x
        feat_x = assemble_feature(data_x, D + 1)
        for i in range(len(LAMBDA)):
            Etrain[D, i], Evalid[D, i] = fit(D + 1, LAMBDA[i])
        print('Average train error:', Etrain, sep='
')
        print('Average valid error:', Evalid, sep='
')
    D, i = np.unravel_index(Evalid.argmin(), Evalid.shape)
    print("D =", D + 1)
    print("lambda =", LAMBDA[i])

if __name__ == "__main__":
    main()
6 Your Own Question

Write your own question, and provide a thorough solution.

Writing your own problems is a very important way to really learn the material. The famous “Bloom’s Taxonomy” that lists the levels of learning is: Remember, Understand, Apply, Analyze, Evaluate, and Create. Using what you know to create is the top-level. We rarely ask you any HW questions about the lowest level of straight-up remembering, expecting you to be able to do that yourself. (e.g. make yourself flashcards) But we don’t want the same to be true about the highest level.

As a practical matter, having some practice at trying to create problems helps you study for exams much better than simply counting on solving existing practice problems. This is because thinking about how to create an interesting problem forces you to really look at the material from the perspective of those who are going to create the exams.

Besides, this is fun. If you want to make a boring problem, go ahead. That is your prerogative. But it is more fun to really engage with the material, discover something interesting, and then come up with a problem that walks others down a journey that lets them share your discovery. You don’t have to achieve this every week. But unless you try every week, it probably won’t happen ever.