02 Linear Regression with Regularization

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Information: Apply the concepts of generalization and regularization to Linear Regression

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1 Generalization for Polynomial Regression

Here is a simple example showing the concepts of overfitting and underfitting. Here directly use the **Scikit-Learn** package for convenience.

1.1 Generate Dataset

Try to fit a linear regression model to the data generated from

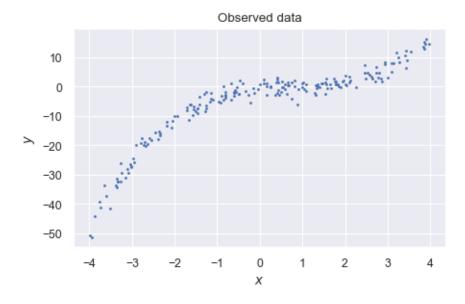
$$y_i = -0.5 + 1.5x_i - 2\frac{x_i^2}{2!} + 2.5\frac{x_i^3}{3!} + 2\epsilon_i$$

where $\epsilon_i \sim \mathcal{N}(0,1)$ and we sample $x_i \sim U([-4,4])$

First generate the dataset and visualize it

```
[1]: import seaborn as sns
     import numpy as np
     import matplotlib.pyplot as plt
     # Ensure reproducibility
     np.random.seed(123)
     # Plot setting
     sns.set()
     sns.set_context('paper')
     # Size of training set and validation set
     num_train, num_test = 160, 40
     # Number of total observations
     num_obs = num_train + num_test
     \# Sample x
     x = (8 * np.random.rand(num_obs) - 4.0).reshape(-1, 1)
     # True parameters
     theta_true = np.array([-0.5, 1.5, -2, 2.5]).reshape(-1, 1)
     # Get the polynomial featurtes
     y = -0.5 + 1.5 * x - 2 * x ** 2 / 2 + 2.5 * x ** 3 / 6
```

```
+ 2 * np.random.randn(num_obs).reshape(-1,1)
# Visualize the dataset
fig, ax = plt.subplots(figsize=(5, 3), dpi=100)
ax.plot(x, y, '.', markersize=2)
ax.set_xlabel('$x$')
ax.set_ylabel('$y$')
ax.set_title('Observed data')
plt.show()
```



1.2 Split training set and validation set

The naive way is to simply divide the observations into two parts

```
[2]: # Get x and y zipped
data = np.concatenate((x, y), axis=1)
# Randomly permute rows
data_permuted = np.random.permutation(data)
# Split in a training set by picking the first num_train rows
data_train = data_permuted[:num_train]
# Split in a validation set
data_val = data_permuted[num_train:]
# Get x_train and y_train
x_train = data_train[:, 0].reshape(-1, 1)
y_train = data_train[:, 1].reshape(-1, 1)
# Get x_val and y_val
x_val = data_val[:, 0].reshape(-1, 1)
y_val = data_val[:, 1].reshape(-1, 1)
# Sanity check
```

```
print('Shape of train set:\t\t', x_train.shape)
print('Shape of validation set:\t', x_val.shape)
fig, ax = plt.subplots(figsize=(5, 3), dpi=100)
ax.plot(x_train, y_train, 'x', label='Training Data')
ax.plot(x_val, y_val, 'o', label='Validation Data')
ax.set_xlabel('$x$')
ax.set_ylabel('$y$')
plt.legend(loc='best')
plt.show()
```

Shape of train set: (160, 1) Shape of validation set: (40, 1)



We can also use the **train_test_split** utility in **Scikit-Learn** to separate the training set and validation set easily

```
[3]: from sklearn.model_selection import train_test_split

x_train, x_val, y_train, y_val = \
train_test_split(x,y,test_size=0.2,random_state=123)

# Sanity check
print('Shape of train set:\t\t', x_train.shape)
print('Shape of validation set:\t', x_val.shape)
fig, ax = plt.subplots(figsize=(5, 3), dpi=100)
ax.plot(x_train, y_train, 'x', label='Training Data')
ax.plot(x_val, y_val, 'o', label='Validation Data')
ax.set_xlabel('$x$')
ax.set_ylabel('$y$')
plt.legend(loc='best')
```

```
plt.show()
```

Shape of train set: (160, 1) Shape of validation set: (40, 1)

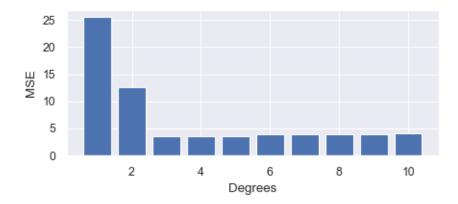


1.3 Train and validation

Let's find the best polynomial degree by training and validation

```
[4]: from sklearn.linear_model import LinearRegression
     from sklearn.pipeline import make_pipeline
     from sklearn.preprocessing import PolynomialFeatures
     # Try with different degree
     degrees = np.arange(1, 11)
     MSE = []
     Estimator = []
     for index, degree in enumerate(degrees):
         # Assign the regression model and preprocessing method
         estimator = make_pipeline(PolynomialFeatures(degree), LinearRegression())
         # Fit with the created model
         estimator.fit(x_train, y_train)
         # See the performance on evaluation points
         y_pred = estimator.predict(x_val).reshape(-1, 1)
         # Calculate the validation MSE
         loss = np.sum((y_pred - y_val)**2) / y_pred.shape[0]
         Estimator.append(estimator)
         MSE.append(loss)
     # Plot out the MSE for different degrees
```

```
fig, ax = plt.subplots(figsize=(5, 2), dpi=100)
ax.bar(range(1, len(MSE) + 1), MSE)
ax.set_xlabel('Degrees')
ax.set_ylabel('MSE')
plt.show()
```



It seems that the MSEs for degree 1 and 2 are so huge that we should not plot them together wit other degrees

```
[5]: # Get the minimum index

print("Degree %d gives the minimized validation mean square error."%(np.

→argmin(MSE)+1))

# Plot out the MSE for different degrees

fig, ax = plt.subplots(figsize=(5, 2), dpi=100)

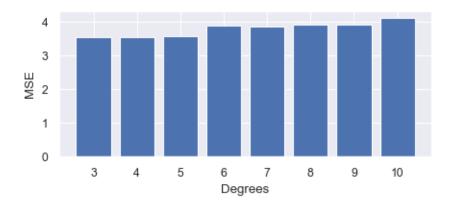
ax.bar(range(3, len(MSE) + 1), MSE[2:])

ax.set_xlabel('Degrees')

ax.set_ylabel('MSE')

plt.show()
```

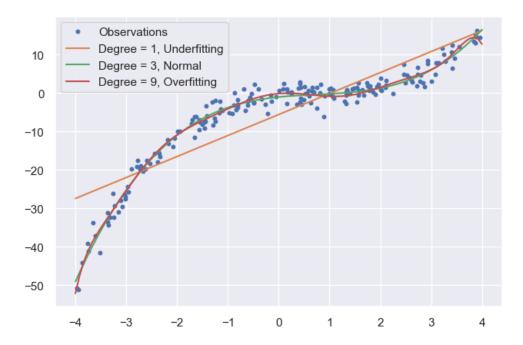
Degree 3 gives the minimized validation mean square error.



1.4 Visualization - Underfitting, Normal fitting and Overfitting

Here shows the visualization of three types of fitting

```
fig, ax = plt.subplots(figsize=(6,4),dpi=125)
xe = np.linspace(-4,4,100).reshape(-1,1)
ye_1 = Estimator[0].predict(xe)
ye_3 = Estimator[2].predict(xe)
ye_10 = Estimator[9].predict(xe)
ax.plot(x,y,'.',label='Observations')
ax.plot(xe,ye_1,label='Degree = 1, Underfitting')
ax.plot(xe,ye_3,label='Degree = 3, Normal')
ax.plot(xe,ye_10,label='Degree = 9, Overfitting')
plt.legend(loc='best')
plt.show()
```



As shown in the plot, the curve with one degree can not capture the features well and is said to be underfitting. The curve with 9 degrees shows strange predictions in the region (3.5,4) and is said to be overfitting. The situation would be far more severe when the size of the dataset is smaller.

2 Polynomial Regression with Regularization