```
Assignment2 (Score: 10.0 / 12.0)

1. Test cell (Score: 2.0 / 2.0)

2. Test cell (Score: 2.0 / 2.0)

3. Test cell (Score: 1.0 / 1.0)

4. Test cell (Score: 2.0 / 2.0)

5. Test cell (Score: 2.0 / 2.0)

6. Test cell (Score: 0.0 / 2.0)

7. Test cell (Score: 1.0 / 1.0)
```

You are currently looking at **version 0.1** of this notebook. To download notebooks and datafiles, as well as get help on Jupyter notebooks in the Coursera platform, visit the Jupyter Notebook FAQ course resource.

# Assignment 2¶

In this assignment you'll explore the relationship between model complexity and generalization performance, by adjust key parameters of various supervised learning models. Part 1 of this assignment will look at regression and Part 2 will I at classification.

## Part 1 - Regression¶

In [1]:

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
np.random.seed(0)
n = 15
x = np.linspace(0,10,n) + np.random.randn(n)/5
y = np.sin(x)+x/6 + np.random.randn(n)/10
X_train, X_test, y_train, y_test = train_test_split(x, y, random_state=0)
def intro():
    %matplotlib notebook
    plt.figure()
    plt.scatter(X_train, y_train, label='training data')
    plt.scatter(X_test, y_test, label='test data')
    plt.legend(loc=4);
intro()
```

	?		

Question 1¶

```
def answer one():
    from sklearn.linear model import LinearRegression
    from sklearn.preprocessing import PolynomialFeatures
    # first step starts with creating an array with values bretween (1,10) using linspace w
    input array = np.linspace(0, 10, 100)
    # We will create a new training data (X train1) , with required shape (-1,1) which must
    # and also because, we will have the need of X train for the next questions
    X_{train} = X_{train.reshape}(-1, 1)
    # I just created a 'predicted array' with a shape (4,100)
    # here, it is created with dtype ='f8' which is float type because, if it is created in
    # we add float values to it, it will append them as int(that float number)
    # thats why, it is initially created as a float type array
    predicted_array = np.arange(400, dtype="f8").reshape(4, 100)
    # list of degrees that can be used for iteration
    degree = [1, 3, 6, 9]
    for x in range(4):
        # Now for each loop, we will create a new training data 'Xtrain_2' using 'X_train1'
        Xtrain 2 = PolynomialFeatures(degree=degree[x]).fit transform(X train1)
        # Now, we create a regressor that trains over 'Xtrain 2' and 'y train'
        linreg = LinearRegression().fit(Xtrain 2, y train)
        # now, we will predict the result of each regression using '.predict' method of the
        # be the polynonial featured data set for 'input array'
        # here each row will be assigned with the predicted values of each degree
        predicted array[x] = linreg.predict(
            PolynomialFeatures(degree=degree[x]).fit transform(
                input array.reshape(-1, 1)
        )
    return predicted array
answer_one()
```

#### Out[2]:

```
array([[ 2.53040195e-01, 2.69201547e-01,
                                         2.85362899e-01,
        3.01524251e-01, 3.17685603e-01,
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                                         5.76267232e-01,
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        6.89396695e-01, 7.05558047e-01, 7.21719399e-01,
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```

```
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```

```
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```

In [3]:

```
Grade cell: cell-4b3a4b2c2971710c

# feel free to use the function plot_one() to replicate the figure
# from the prompt once you have completed question one
def plot_one(degree_predictions):
    plt.figure(figsize=(10,5))
    plt.plot(X_train, y_train, 'o', label='training data', markersize=10)
    plt.plot(X_test, y_test, 'o', label='test data', markersize=10)
    for i,degree in enumerate([1,3,6,9]):
        plt.plot(np.linspace(0,10,100), degree_predictions[i], alpha=0.8, lw=2, label='degr
    plt.ylim(-1,2.5)
    plt.legend(loc=4)

plot_one(answer_one())
```

## Question 2¶

Write a function that fits a polynomial LinearRegression model on the training data  $X_{train}$  for degrees 0 through 9. F each model compute the  $\mathbb{R}^2$  (coefficient of determination) regression score on the training data as well as the the test  $\mathfrak{c}$  and return both of these arrays in a tuple.

This function should return a tuple of numpy arrays (r2\_train, r2\_test). Both arrays should have shape (10,) In [4]:

Student's answer

```
def answer two():
    from sklearn.linear_model import LinearRegression
    from sklearn.metrics import r2 score
    from sklearn.preprocessing import PolynomialFeatures
   # create empty lists for r2 train and r2 test
    r2 train = []
    r2_{test} = []
   # then, create new data sets based on X train and X test with the required shape of onl
   X_{train1} = X_{train.reshape(-1, 1)}
   X \text{ test1} = X \text{ test.reshape}(-1, 1)
    for x in range(10):
        # Now, create polynomial featured datasets from X train1, X test1 which is used to
        X train2 = PolynomialFeatures(degree=x).fit transform(X train1)
        X_test2 = PolynomialFeatures(degree=x).fit_transform(X_test1)
        # Now, train the regressor with the above obtained polynomial featured training dat
        linreg = LinearRegression().fit(X_train2, y_train)
        y predicted train = linreq.predict(X train2)
        y_predicted_test = linreg.predict(X_test2)
        # Now, now append the scores of training and testing data sets into the respective
        r2_train.append(r2_score(y_train, y_predicted_train))
        r2_test.append(r2_score(y_test, y_predicted_test))
   # Now, convert the lists into arrays of required shape(10,1)
   # Note - in this question, it is asked to make the shape of array as (10,1) but auto gr
   # reshape it as (1,10)
    r2 train = np.array(r2 train).reshape(10, 1)
    r2 test = np.array(r2 test).reshape(10, 1)
   # Your code here
    ans = tuple([r2 train, r2 test])
    return ans
answer two()
```

## Out[4]:

```
(array([[0.
        [0.42924578],
        [0.4510998],
        [0.58719954],
        [0.91941945],
        [0.97578641],
        [0.99018233],
        [0.99352509],
        [0.99637545],
        [0.99803706]]),
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        [-0.45237104],
        [-0.06856984],
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        [ 0.73004943],
        [ 0.87708301],
        [ 0.9214094 ],
        [0.92021504],
        [0.63247942],
        [-0.64525285]]))
```

## In [5]:

Grade cell: cell-84a9de5d00b87d2f Score: 2.0 / 2.0

## Question 3¶

Based on the  $R^2$  scores from question 2 (degree levels 0 through 9), what degree level corresponds to a model that is underfitting? What degree level corresponds to a model that is overfitting? What choice of degree level would provide a model with good generalization performance on this dataset?

(Hint: Try plotting the  $R^2$  scores from question 2 to visualize the relationship)

This function should return a tuple with the degree values in this order: (Underfitting, Overfitting, Good Generalization)

In [6]:

Student's answer

```
def answer_three():
    # Your code here
    (r2_train, r2_test) = answer_two()
      print(r2_test)
#
      print(r2_train)
#
      import matplotlib.pyplot as plt
#
      %matplotlib notebook
#
      plt.figure()
#
      plt.plot(r2_train, label='data')
      plt.plot(r2_test, label='data')
    order = (2,9,7)
    return (order) # Return
answer_three()
```

Out[6]:

```
(2, 9, 7)
```

In [7]:

Grade cell: cell-877e9f32963e0d5a

Score: 1.0 / 1.0

## Question 4¶

Training models on high degree polynomial features can result in overfitting. Train two models: a non-regularized LinearRegression model and a Lasso Regression model (with parameters alpha=0.01,  $max_iter=10000$ , tol=0.1) c polynomial features of degree 12. Return the  $R^2$  score for LinearRegression and Lasso model's test sets.

This function should return a tuple (LinearRegression R2 test score, Lasso R2 test score)

#### In [8]:

#### Student's answer

```
def answer four():
    from sklearn.preprocessing import PolynomialFeatures
    from sklearn.linear_model import Lasso, LinearRegression
    from sklearn.metrics import r2_score
    from sklearn.preprocessing import MinMaxScaler
    scaler = MinMaxScaler()
    poly = PolynomialFeatures(degree=12)
   X_poly = poly.fit_transform(X_train.reshape(11,1))
   X_test_poly = poly.fit_transform(X_test.reshape(4,1))
    # Create regressor
    linreg = LinearRegression().fit(X_poly, y_train)
   X_train_scaled = scaler.fit_transform(X_poly)
    X_test_scaled = scaler.transform(X_test_poly)
    linlasso = Lasso(alpha=0.01, max_iter = 10000, tol = 0.1).fit(X_poly, y_train)
    return r2_score(y_test, linreg.predict(X_test_poly)) , r2_score(y_test, linlasso.predic
answer_four()
```

#### Out[8]:

(-4.3119675863308435, 0.6051396919570036)

#### In [9]:

Grade cell: cell-f1ccfec3713e840c

Score: 2.0 / 2.0

## Part 2 - Classification¶

For this section of the assignment we will be working with the UCI Mushroom Data Set

(http://archive.ics.uci.edu/ml/datasets/Mushroom?ref=datanews.io) stored in mushrooms.csv. The data will be used to trian a model to predict whether or not a mushroom is poisonous. The following attributes are provided:

#### Attribute Information:

- 1. cap-shape: bell=b, conical=c, convex=x, flat=f, knobbed=k, sunken=s
- 2. cap-surface: fibrous=f, grooves=g, scaly=y, smooth=s
- 3. cap-color: brown=n, buff=b, cinnamon=c, gray=g, green=r, pink=p, purple=u, red=e, white=w, yellow=y
- 4. bruises?: bruises=t, no=f
- 5. odor: almond=a, anise=l, creosote=c, fishy=y, foul=f, musty=m, none=n, pungent=p, spicy=s
- 6. gill-attachment: attached=a, descending=d, free=f, notched=n
- 7. gill-spacing: close=c, crowded=w, distant=d
- 8. gill-size: broad=b, narrow=n
- 9. gill-color: black=k, brown=n, buff=b, chocolate=h, gray=g, green=r, orange=o, pink=p, purple=u, red=e, white=w, yellow=y
- 10. stalk-shape: enlarging=e, tapering=t
- 11. stalk-root: bulbous=b, club=c, cup=u, equal=e, rhizomorphs=z, rooted=r, missing=?
- 12. stalk-surface-above-ring: fibrous=f, scaly=y, silky=k, smooth=s
- 13. stalk-surface-below-ring: fibrous=f, scaly=y, silky=k, smooth=s
- 14. stalk-color-above-ring: brown=n, buff=b, cinnamon=c, gray=g, orange=o, pink=p, red=e, white=w, yellow=y
- 15. stalk-color-below-ring: brown=n, buff=b, cinnamon=c, gray=g, orange=o, pink=p, red=e, white=w, yellow=y
- 16. veil-type: partial=p, universal=u
- 17. veil-color: brown=n, orange=o, white=w, yellow=y
- 18. ring-number: none=n, one=o, two=t
- 19. ring-type: cobwebby=c, evanescent=e, flaring=f, large=l, none=n, pendant=p, sheathing=s, zone=z
- 20. spore-print-color: black=k, brown=n, buff=b, chocolate=h, green=r, orange=o, purple=u, white=w, yellow=y
- 21. population: abundant=a, clustered=c, numerous=n, scattered=s, several=v, solitary=y
- 22. habitat: grasses=g, leaves=l, meadows=m, paths=p, urban=u, waste=w, woods=d

The data in the mushrooms dataset is currently encoded with strings. These values will need to be encoded to numeric work with sklearn. We'll use pd.get\_dummies to convert the categorical variables into indicator variables.

## In [10]:

```
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split

mush_df = pd.read_csv('assets/mushrooms.csv')
mush_df2 = pd.get_dummies(mush_df)

X_mush = mush_df2.iloc[:,2:]
y_mush = mush_df2.iloc[:,1]

X_train2, X_test2, y_train2, y_test2 = train_test_split(X_mush, y_mush, random_state=0)
```

```
In [11]:
```

```
X_subset = X_test2
y_subset = y_test2
```

## Question 5¶

Using X\_train and y\_train from the preceding cell, train a DecisionTreeClassifier with default parameters and random state=0. What are the 5 most important features found by the decision tree?

This function should return a list of length 5 of the feature names in descending order of importance.

In [12]:

```
Student's answer
def answer five():
    from sklearn.tree import DecisionTreeClassifier
    # First, creat a classifer using the given data sets X_train and y_train2
    classifier = DecisionTreeClassifier(random state=0).fit(X train2, y train2)
    # then, using the feature importance attribute, create a list of those feature importan
    features list = list(classifier.feature_importances_)
    features_list.sort(reverse=True)
    top 5 features num list = features list[:5]
    # then, find those index positions, where the top 5 featueres are in the list
    index pos list = [
        list(classifier.feature importances).index(x) for x in top 5 features num list
    # then, find the name of the features from the .columns of the dataset X train2
    top 5 features list = [X train2.columns[x] for x in index pos list]
    # Your code here
    return top_5_features_list
answer_five()
```

#### Out[12]:

```
['odor_n', 'stalk-root_c', 'stalk-root_r', 'spore-print-color_r', 'odor_l']
```

#### In [13]:

Grade cell: cell-45bb9fef0723e714

Score: 2.0 / 2.0

## Question 6¶

For this question, use the validation\_curve function in sklearn.model\_selection to determine training and test scores for a Support Vector Classifier (SVC) with varying parameter values.

Create an SVC with default parameters (i.e. kernel='rbf', C=1) and  $random\_state=0$ . Recall that the kernel width c the RBF kernel is controlled using the gamma parameter. Explore the effect of gamma on classifier accuracy by using the validation\_curve function to find the training and test scores for 6 values of gamma from 0.0001 to 10 (i.e. np.logspace(-4,1,6)).

For each level of gamma, validation\_curve will use 3-fold cross validation (use cv=3, n\_jobs=2 as parameters for validation\_curve), returning two 6x3 (6 levels of gamma x 3 fits per level) arrays of the scores for the training and te sets in each fold.

Find the mean score across the five models for each level of gamma for both arrays, creating two arrays of length 6, an return a tuple with the two arrays.

e.g.

if one of your array of scores is

#### it should then become

```
array([ 0.5, 0.73333333, 0.83333333, 0.76666667, 0.63333333, 0.5])
```

This function should return a tuple of numpy arrays (training\_scores, test\_scores) where each array in the tuple shape (6,).

#### In [14]:

Student's answer

#### Out[14]:

#### In [15]:

Grade cell: cell-ae2a630e8862c3a3

Score: 0.0 / 2.0

You have failed this test due to an error. The traceback has been removed because it may cor AssertionError: Q6: The training\_scores has incorrect value at index 0

## Question 7¶

Based on the scores from question 6, what gamma value corresponds to a model that is underfitting? What gamma value corresponds to a model that is overfitting? What choice of gamma would provide a model with good generalization performance on this dataset?

(Hint: Try plotting the scores from question 6 to visualize the relationship)

This function should return a tuple with the degree values in this order: (Underfitting, Overfitting, Good\_Generalization)

# In [16]: Student's answer def answer seven(): #As, there are limited number of elements, we can conclude things, just by looking over tha #For overfitting, i think gamma = $10^1$ is the correct as, it has the best training score an #For underfitting, i think gamma = 9 gamma = $10^{-4}$ is the correct one, as it has least traini #For good generalization, gamma = $10^{-1}$ is the best, as it has a training score and testing # Your code here ans = tuple([0.0001,10,0.1]) return ans answer\_seven() Out[16]: (0.0001, 10, 0.1) In [17]: Score: 1.0 / 1.0 Grade cell: cell-6372c6701a14c068 In []: In []:

This assignment was graded by mooc\_adswpy:e5e20d3b91dd, v1.45.052423