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# CS 178 Homework 2

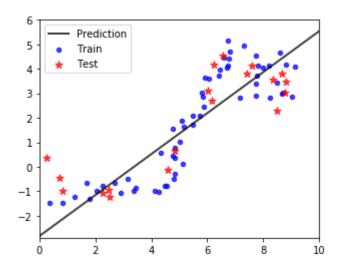
## **Problem 1**

1.

```
In [237]:
          import numpy as np
          import matplotlib.pyplot as plt
          import mltools as ml
          # Load data
          data = np.genfromtxt("data/curve80.txt", delimiter=None)
          #split data
          X = data[:,0]
          X = np.atleast_2d(X).T # code expects shape (M,N) so make sure it's 2-dimen
          sional
          Y = data[:,1]
          Xtr, Xte, Ytr, Yte = ml.splitData(X,Y,0.75) # split data set 75/25
          # print the shapes of these four objects
          print(Xtr.shape)
          print(Xte.shape)
          print(Ytr.shape)
          print(Yte.shape)
          (60, 1)
          (20, 1)
           (60,)
          (20,)
```

```
In [244]: print("a)")
          lr = ml.linear.linearRegress( Xtr, Ytr ) # create and train model
          xs = np.linspace(0,10,200) \# densely sample possible x-values
          xs = xs[:,np.newaxis] # force "xs" to be an Mx1 matrix (expected by our cod
          e)
          ys = lr.predict( xs ) # make predictions at xs
          f, ax = plt.subplots(1, 1, figsize=(5, 4))
          ax.scatter(Xtr, Ytr, s=20, color='blue', alpha=0.75, label='Train')
          ax.scatter(Xte, Yte, s=60, marker='*', color='red', alpha=0.75, label='Tes
          t')
          ax.plot(xs, ys, lw=2, color='black', alpha=0.75, label='Prediction')
          ax.set xlim(0, 10)
          ax.set ylim(-2.9, 6)
          ax.legend(fontsize=10, loc=0)
          plt.show()
          print("b)")
          print(lr.theta)
          print('''The linear regression coefficients do match the plot
          because the y-intercept does appear to be somewhere close to
          -3 and taking the points (2,-1) and (8,4) on the graph, which
          are close approximates that fall on the line of regression,
          we can calculate that the slope is 5/6 = 0.833, which is
          a value close to the coefficient 0.8361.''')
          print("\nc)")
          def MSE(lr, X, Y):
              Yhat = lr.predict(X)
              return np.mean((Y - Yhat.reshape(Y.shape))**2 , axis=0)
          print("Training data MSE:", MSE(lr, Xtr, Ytr))
          print("Test data MSE:", MSE(lr, Xte, Yte))
```

a)



b) [[-2.82765049 0.83606916]]

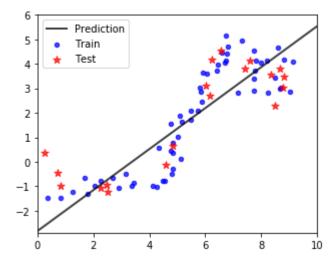
The linear regression coefficients do match the plot because the y-intercept does appear to be somewhere close to -3 and taking the points (2,-1) and (8,4) on the graph, which are close approximates that fall on the line of regression, we can calculate that the slope is 5/6 = 0.833, which is a value close to the coefficient 0.8361.

c)
Training data MSE: 1.127711955609391
Test data MSE: 2.2423492030101246

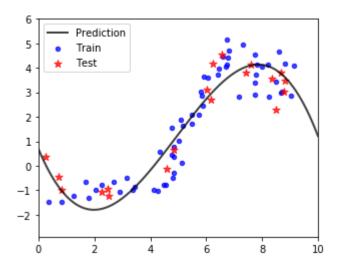
```
In [239]: print("a)\n")
          degrees = np.array([1,3,5,7,10,18])
          train errors = np.zeros(degrees.shape[0])
          test errors = np.zeros(degrees.shape[0])
          for i in range(6):
              degree = degrees[i]
              XtrP = ml.transforms.fpoly(Xtr, degree, False)
              XtrP,params = ml.transforms.rescale(XtrP)
              XteP,_ = ml.transforms.rescale(ml.transforms.fpoly(Xte, degree, False),
           params)
              lr = ml.linear.linearRegress(XtrP, Ytr)
              xs = np.linspace(0, 10, 200)
              xs = np.atleast 2d(xs).T
              # Transform the predicting xs
              xsP,_ = ml.transforms.rescale(ml.transforms.fpoly(xs, degree, False), p
          arams)
              ys = lr.predict(xsP)
              train errors[i] = MSE(lr, XtrP, Ytr)
              test errors[i] = MSE(lr, XteP, Yte)
              # Plot the data
              f, ax = plt.subplots(1, 1, figsize=(5, 4))
              ax.scatter(Xtr, Ytr, s=20, color='blue', alpha=0.75, label='Train')
              ax.scatter(Xte, Yte, s=60, marker='*', color='red', alpha=0.75, label=
           'Test')
              # Plot the regression line
              ax.plot(xs, ys, lw=2, color='black', alpha=0.75, label='Prediction')
              ax.set_xlim(0, 10)
              ax.set_ylim(-2.9, 6)
              # Control the size of the legend and the location.
              ax.legend(fontsize=10, loc=0)
              print("degree", degrees[i])
              plt.show()
```

a)

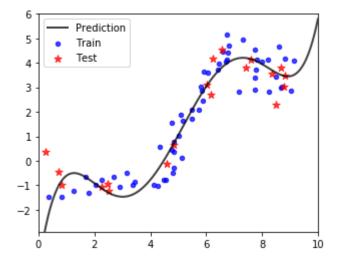
degree 1



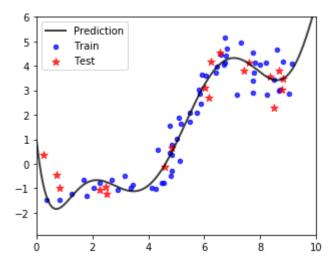
degree 3



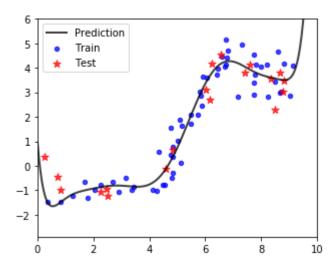
degree 5



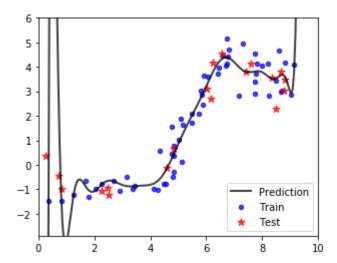
degree 7



degree 10



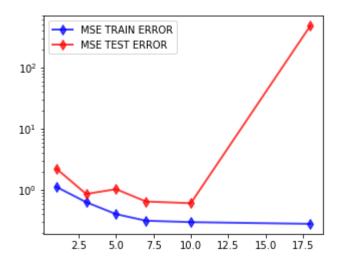
degree 18



```
In [240]: print("b)")
    f, ax = plt.subplots(1, 1, figsize=(5, 4))
    ax.semilogy(degrees, train_errors, lw=2, marker='d', color = "blue", marker
    size=7, alpha=0.75, label='MSE TRAIN ERROR')
    ax.semilogy(degrees, test_errors, lw=2, marker='d', color = "red", markersi
    ze=7, alpha=0.75, label='MSE TEST ERROR')
    ax.legend(fontsize=10, loc=0)
    plt.show()

    print("\nc) I would recommend polynomial degree 5.")
```

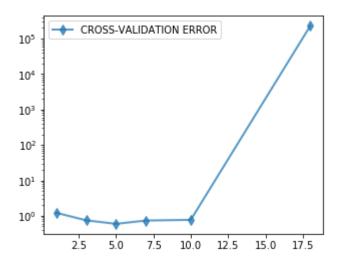
b)



c) I would recommend polynomial degree 5.

# **Problem 2**

```
In [241]:
          def find cross validation error(degree: int, folds: int):
               J = np.zeros(folds)
               for iFold in range(folds):
                   Xti, Xvi, Yti, Yvi = ml.crossValidate(Xtr, Ytr, folds, iFold) # use ith b
           lock as validation
                   XtiP = ml.transforms.fpoly(Xti, degree, False)
                   XtiP,params = ml.transforms.rescale(XtiP)
                   XviP,_ = ml.transforms.rescale(ml.transforms.fpoly(Xvi, degree, Fal
           se), params)
                   lr = ml.linear.linearRegress(XtiP, Yti)
                   J[iFold] = MSE(lr, XviP, Yvi)
               return np.mean(J)
           errors = np.zeros(degrees.shape)
          for i,degree in enumerate(degrees):
               errors[i] = find_cross_validation_error(degree, 5)
           f, ax = plt.subplots(1, 1, figsize=(5, 4))
           ax.semilogy(degrees, errors, lw=2, marker='d', markersize=7, alpha=0.75, la
          bel='CROSS-VALIDATION ERROR')
           ax.legend(fontsize=10, loc=0)
           plt.show()
```



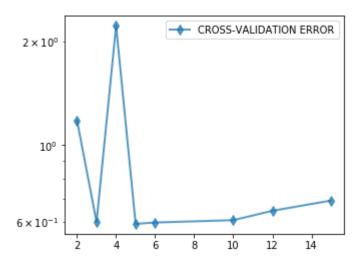
## 2.

The MSE estimates from five-fold cross-validation are pretty similar compared to the MSE's evaluated on Problem 1, though the higher degrees are notably worse and start overfitting sooner.

### 3.

I would recommend polynomial degree 5 based on five-fold cross-validation error.

```
In [242]: nFolds = np.array([2,3,4,5,6,10,12,15])
          errors = np.zeros(nFolds.shape)
          for i,fold in enumerate(nFolds):
              errors[i] = find cross validation error(5, fold)
          f, ax = plt.subplots(1, 1, figsize=(5, 4))
          ax.semilogy(nFolds, errors, lw=2, marker='d', markersize=7, alpha=0.75, lab
          el='CROSS-VALIDATION ERROR')
          ax.legend(fontsize=10, loc=0)
          plt.show()
          print('''There's a large amount of fluctuation of cross-validation
          errors at the beginning because we don't have enough test samples
          to average out our errors. However, past a certain number of folds,
          the errors start to increase a lot more steadily as the numbres of
          folds increases. This can be explained by the fact that we have
          more test results to average as we increaes the number of folds,
          but the validation errors will steadily increaseover time because
          we are reducing the amount of training data each time we create a
          new fold, which will make way for more errors.''')
```



There's a large amount of fluctuation of cross-validation errors at the beginning because we don't have enough test samples to average out our errors. However, past a certain number of folds, the errors start to increase a lot more steadily as the numbres of folds increases. This can be explained by the fact that we have more test results to average as we increases the number of folds, but the validation errors will steadily increaseover time because we are reducing the amount of training data each time we create a new fold, which will make way for more errors.

### **Problem 3: Statement of Collaboration**

I collaborated with Jordan Rayfield on Problem 1.2 to discuss how to define an MSE function.