# Understanding Experimental Data, cont.

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## Remember our goal

- Want to find a model that fits experimental data well
- •Model will then allow us to explain phenomena, and to make predictions about behavior in new settings
- •Know that data is unlikely to be perfect, so have to account for uncertainty in measurements or observations
- Sometimes have theoretical knowledge of structure of model, but not always
  - In latter case, want to try to find best model from class of options

# Solving for Least Squares (Recap)

$$\sum_{i=0}^{len(observed)-1} (observed[i]-predicted[i])^{2}$$

- Given observed data, and model prediction of expected values, can measure goodness of fit of model to observation using sum-of-squared-differences (or meansquared-error)
- Want to find best model for predicting values
- Predicted values often come from mathematical expression, with set of parameters that can vary – typically a polynomial expression
- Use linear regression to find best model that minimizes difference – for polynomial model, this include coefficients, and may include order of polynomial

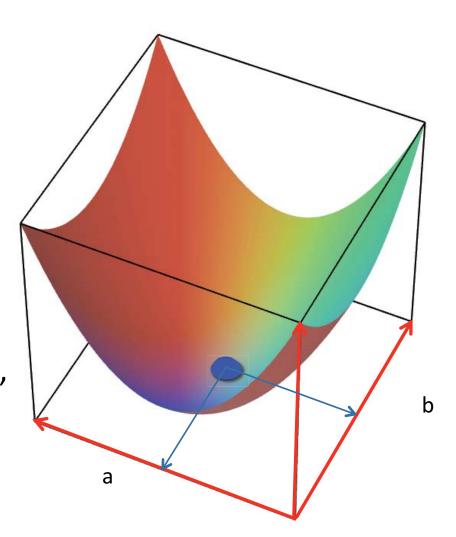
# Solving for Least Squares (Recap)

$$\sum_{i=0}^{len(observed)-1} (observed[i]-predicted[i])^{2}$$

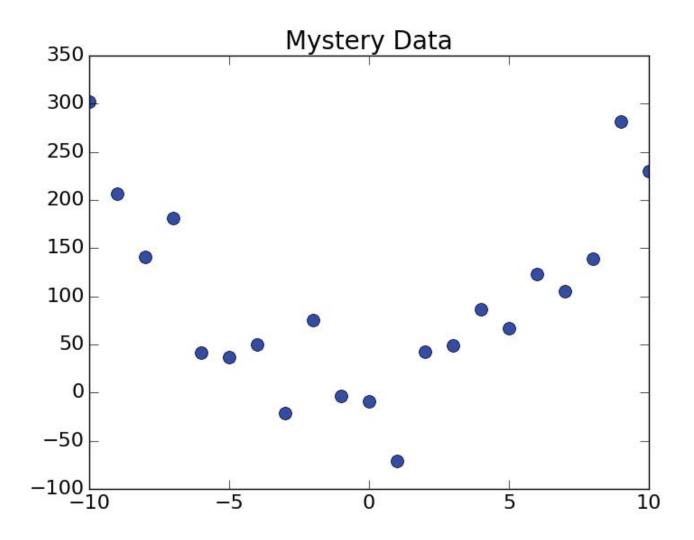
- Simple example:
  - Use a degree-one polynomial, y = ax+b, as model of our data (we want best fitting line)
- •Find values of a and b such that when we use the polynomial to predict y values for all of the x values in our experiment, the squared difference of these values and the corresponding observed values is minimized
- A linear regression problem

# Finding the best curve (simplest case)

- The set of all possible lines can be represented by a point in a-b space
- Imagine a surface in this space, where height of the surface is the value of the objective function
- Starting at any point on the surface, walk "downhill", until you reach the "bottom"
- Corresponding point is best line to fit to data
- Can generalize to higher order models



# Another Experiment (Recap)

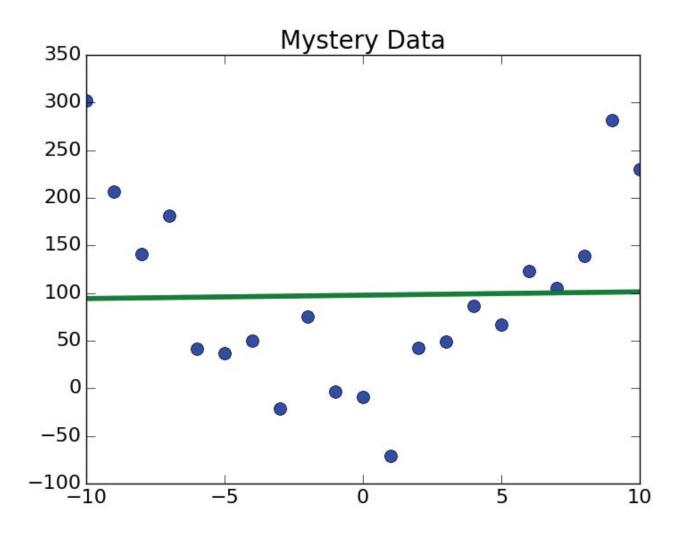


#### Fit a Line

- Remember that pylab.polyfit will find parameters of best fitting polynomial of described order
  - In this case (with argument n = 1), find the values of a and b, such that y = ax + b best matches the observed yVals
- •Remember that pylab.polyval will generate predicted yVals given parameters of model

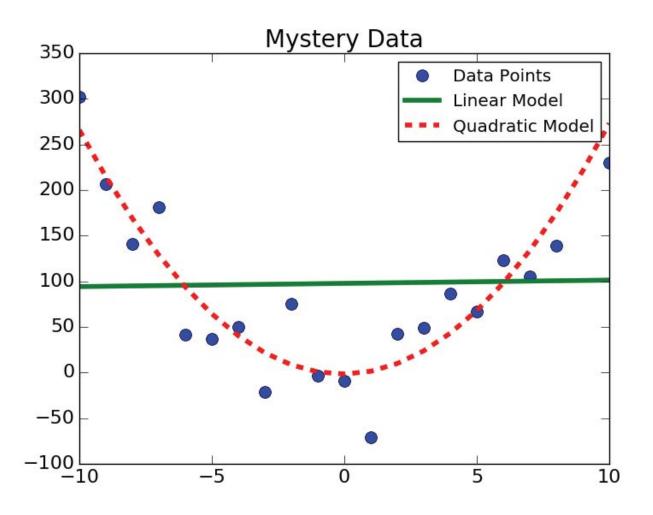
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## Fit a Line



## Let's Try a Higher-degree Model

# Quadratic Appears to be a Better Fit



## Can We Get a Tighter Fit?

- •What if we try fitting higher order polynomials to the data?
  - Does this give us a better fit?
- •How would we measure that?
  - In absence of other information (e.g., theoretical insights into order of model), R<sup>2</sup> (coefficient of determination) gives us decent measure of the tightness of the model fit
  - In principle, a model with a higher R<sup>2</sup> value is a "better" fit

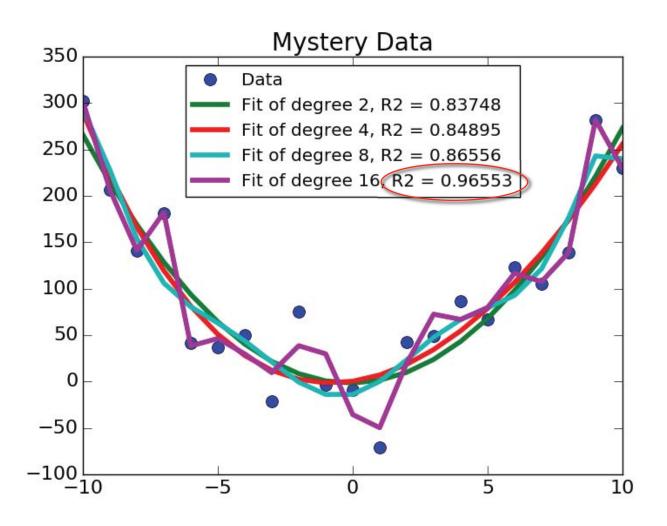
$$R^2 = 1 - \frac{\sum_i (y_i - p_i)^2}{\sum_i (y_i - \mu)^2}$$
 Error in estimates

Variability in measured data

Y<sub>i</sub> are measured values

P<sub>i</sub> are predicted values μ is mean of measured values

# Can We Get a Tighter Fit?

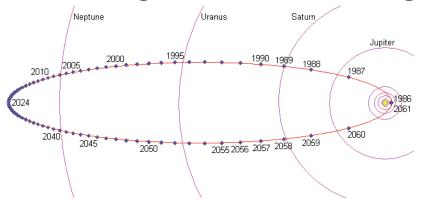


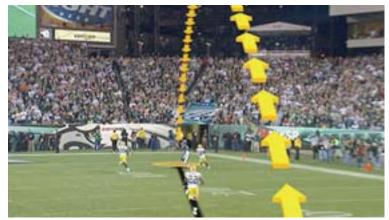
## Why We Build Models

- Looks like an order 16 fit is really good so should we just use this as our model?
  - To answer, need to ask why build models in first place?
- •Help us understand process that generated the data
  - E.g., the properties of a particular linear spring
- Help us make predictions about out-of-sample data
  - E.g., predict the displacement of a spring when a force is applied to it
  - E.g., predict the effect of treatment on a patient
  - E.g., predict the outcome of an election
- A good model helps us do both of these things

## Motivation for Mystery Data – Parabola

- Trajectory of a particle under the influence of a uniform gravitational field (e.g. Halley's Comet)
- Position of center of mass of a football pass
- Design of a load-bearing arch







Images of particle trajectory, load-bearing arch, football pass center of mass diagram © sources unknown. All rights reserved. This content is excluded from out Creative Commons license. For more information, see <a href="https://ocw.mit.edu/help/faq-fair-use/">https://ocw.mit.edu/help/faq-fair-use/</a>.

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## How Mystery Data Was Generated

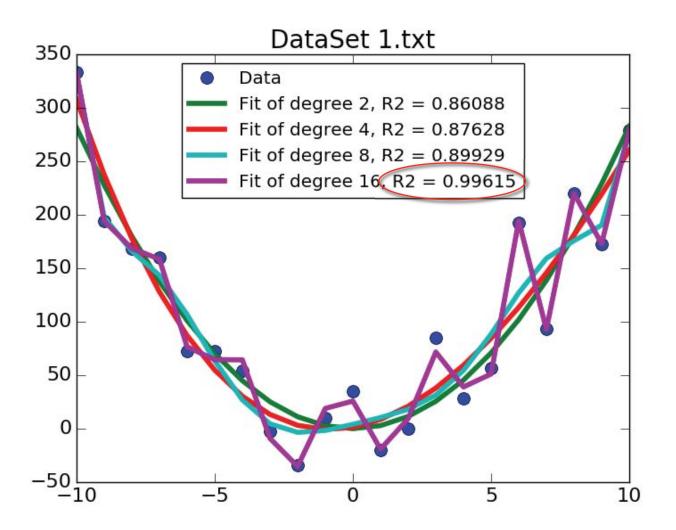
```
def genNoisyParabolicData(a, b, c, xVals, fName):
   yVals = []
    for x in xVals:
        theoreticalVal = a*x**2 + b*x + a
        yVals.append(theoreticalVal + random.gauss(0, 35))
    f = open(fName, 'w')
    f.write('x y\n')
    for i in range(len(yVals)):
        f.write(str(yVals[i]) + ' ' + str(xVals[i]) + '\n')
    f.close()
#parameters for generating data
xVals = range(-10, 11, 1)
a, b, c = 3, 0, 0
genNoisyParabolicData(a, b, c, xVals, 'Mystery Data.txt')
```

If data was generated by quadratic, why was 16<sup>th</sup> order polynomial the "best" fit?

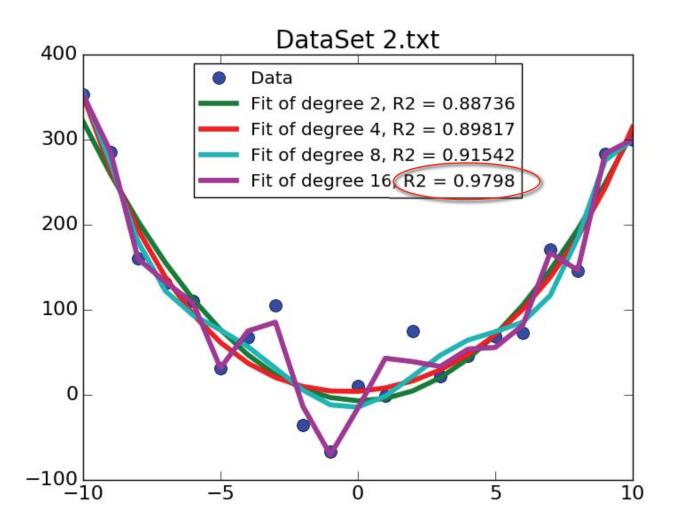
#### Let's Look at Two Data Sets

```
degrees = (2, 4, 8, 16)
random.seed(0)
xVals1, yVals1 = getData('Dataset 1.txt')
models1 = genFits(xVals1, yVals1, degrees)
testFits(models1, degrees, xVals1, yVals1,
        'DataSet 1.txt')
pylab.figure()
xVals2, yVals2 = getData('Dataset 2.txt')
models2 = genFits(xVals2, yVals2, degrees)
testFits(models2, degrees, xVals2, yVals2,
         'DataSet 2.txt')
```

#### Fits for Dataset 1



#### Fits for Dataset 2



## Hence Degree 16 Is Tightest Fit

- "Best" fitting model is still order 16 polynomial for both data sets, but we know data was generated using an order 2 polynomial?
- What we are seeing comes from training error
  - How well the model performs on the data from which it was learned
  - Small training error a necessary condition for a great model,
     but not a sufficient one
- We want model to work well on other data generated by the same process
  - Measurements for other weights on the spring
  - Positions of comets under different forces
  - Voters other than those surveyed
- In other words, the model needs to generalize

#### **Cross Validate**

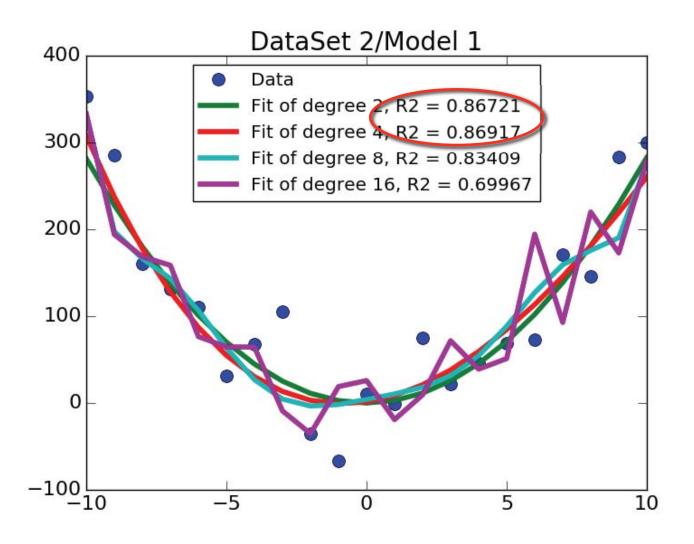
- Generate models using one dataset, and then test them on another dataset
  - Use models for Dataset 1 to predict points for Dataset 2
  - Use models for Dataset 2 to predict points for Dataset 1
- Expect testing error to be larger than training error
- A better indication of generalizability than training error

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#### Test Code

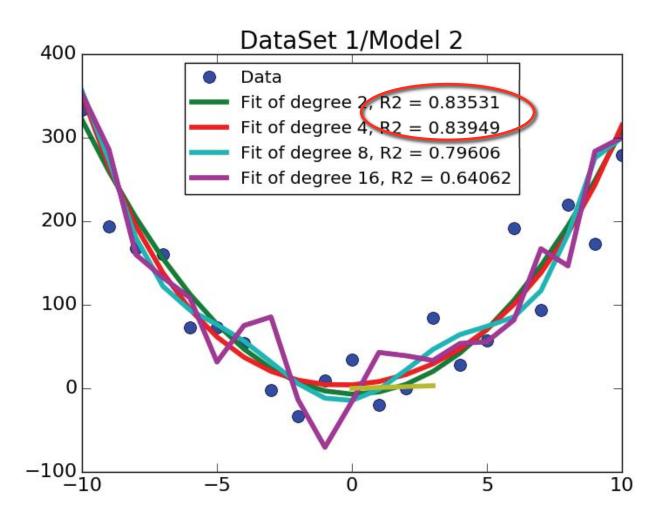
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## Train on Dataset 1, Test on Dataset 2



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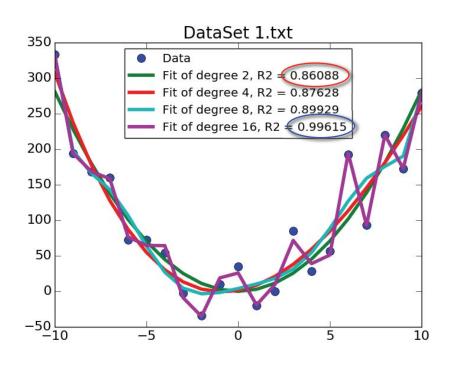
#### Train on Dataset 2, Test on Dataset 1

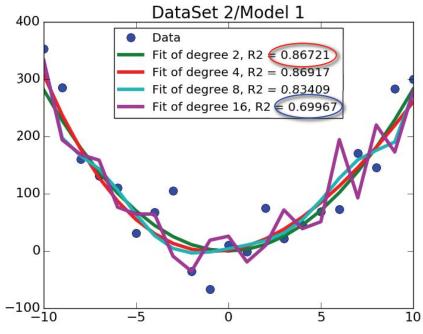


#### **Cross Validation**

- Now can see that based on R<sup>2</sup> numbers, best model is more likely to be 2<sup>nd</sup> order or 4<sup>th</sup> order polynomial (we know it is actually 2<sup>nd</sup> order, and difference in R<sup>2</sup> values is pretty small), but certainly not 16<sup>th</sup> order
- Example of over fitting to the data
- •Can see that if we only fit model to training data, we may not detect that model is too complex; but training on one data set, then testing on a second helps expose this problem

# **Training and Testing Errors**



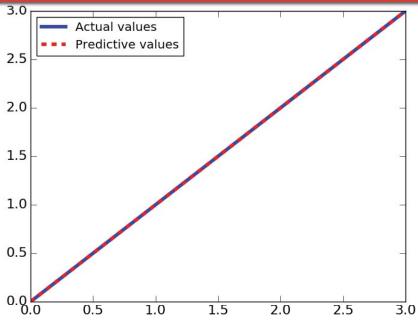


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## Increasing the Complexity

- •Why do we get a "better" fit on training data with higher order model, but then do less well on handling new data?
- •What happens when we increase order of polynomial during training?
  - Can we get a worse fit to training data?
- If extra term is useless, coefficient will merely be zero
- •But if data is noisy, can fit the noise rather than the underlying pattern in the data
  - May lead to a "better" R<sup>2</sup> value, but not really a "better" fit

## Fitting a Quadratic to a Perfect Line



$$y = ax^{2} + bx + c$$

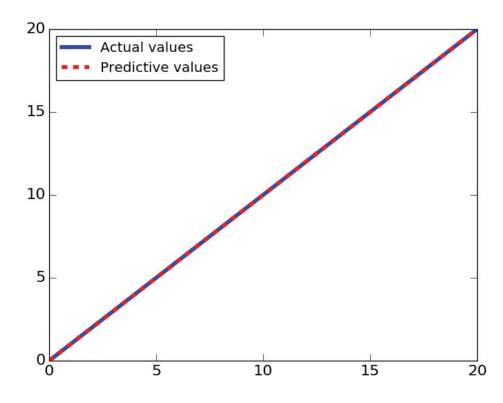
$$y = 0x^{2} + 1x + 0$$

$$y = x$$

R-squared = 1.0

## Predict Another Point Using Same Model

```
xVals = xVals + (20,)
yVals = xVals
pylab.plot(xVals, yVals, label = 'Actual values')
estYVals = pylab.polyval((a,b,c), xVals)
pylab.plot(xVals, estYVals, 'r--', label = 'Predictive values')
print('R-squared = ', rSquared(yVals, estYVals))
```

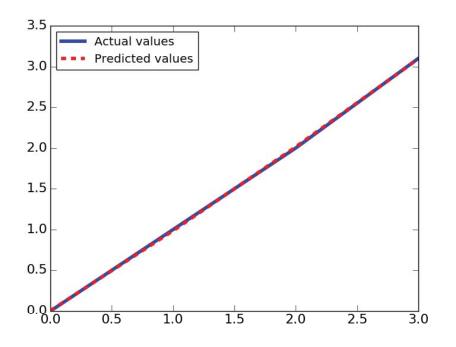


R-squared = 1.0

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#### Simulate a Small Measurement Error

```
xVals = (0,1,2,3)
yVals = (0,1,2(3.1))
pylab.plot(xVals, yVals, label = 'Actual values')
model = pylab.polyfit(xVals, yVals, 2)
print(model)
estYVals = pylab.polyval(model, xVals)
pylab.plot(xVals, estYVals, 'r--', label = 'Predicted values')
print('R-squared = ', rSquared(yVals, estYVals))
```



$$y = ax^2 + bx + c$$
  
 $y = .025x^2 + .955x + .005$ 

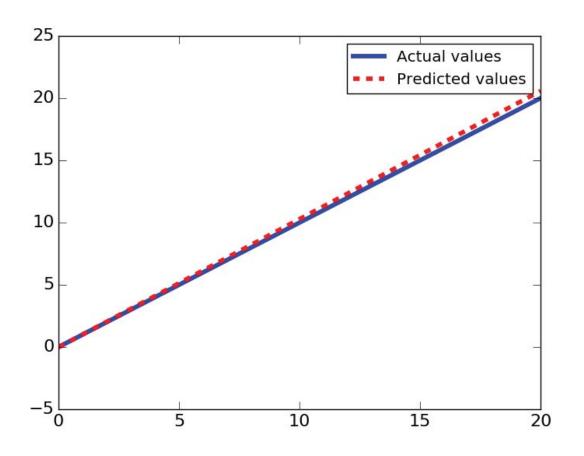
$$R$$
-squared =  $0.9994$ 

## Predict Another Point Using Same Model

```
xVals = xVals + ((20,))
yVals = xVals
estYVals = pylab.polyval(model, xVals)
print('R-squared = ', rSquared(yVals, estYVals))
pylab.figure()
pylab.plot(xVals, estYVals)
30
      Actual values
      Predicted values
25
20
                                      R-squared = 0.7026
15
10
 5
                   10
                           15
                                    20
```

## Suppose We Had Used a First-degree Fit

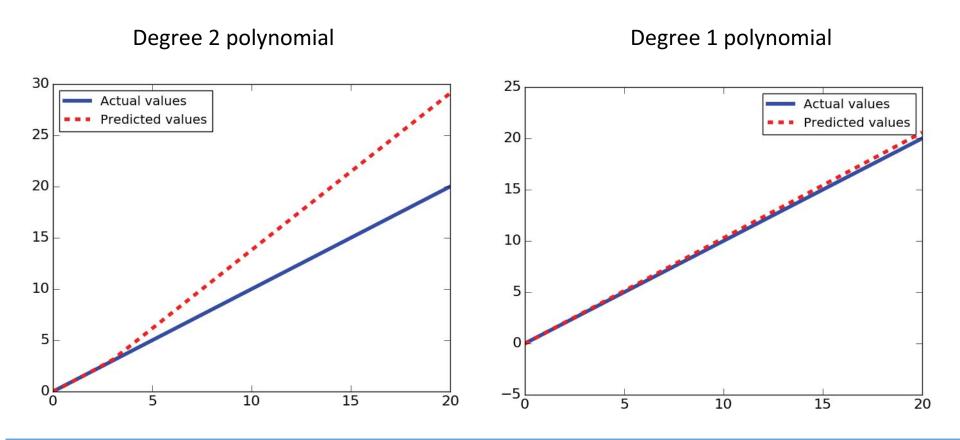
model = pylab.polyfit(xVals, yVals, 1)



R-squared = 0.9988

# Comparing first and second degree fits

 Predictive ability of first order fit much better than second order fit

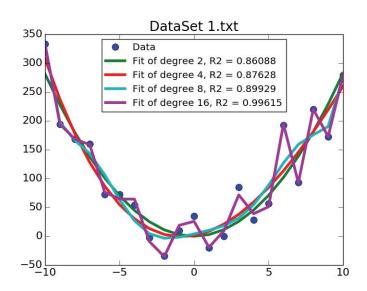


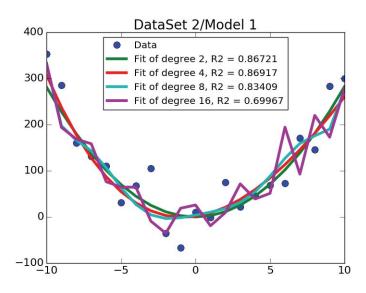
## The Take Home Message

- Choosing an overly-complex model leads to overfitting to the training data
- Increases the risk of a model that works poorly on data not included in the training set
- On the other hand choosing an insufficiently complex model has other problems
  - As we saw when we fit a line to data that was basically parabolic
  - "Everything should be made as simple as possible, but not simpler" – Albert Einstein

## **Balancing Fit with Complexity**

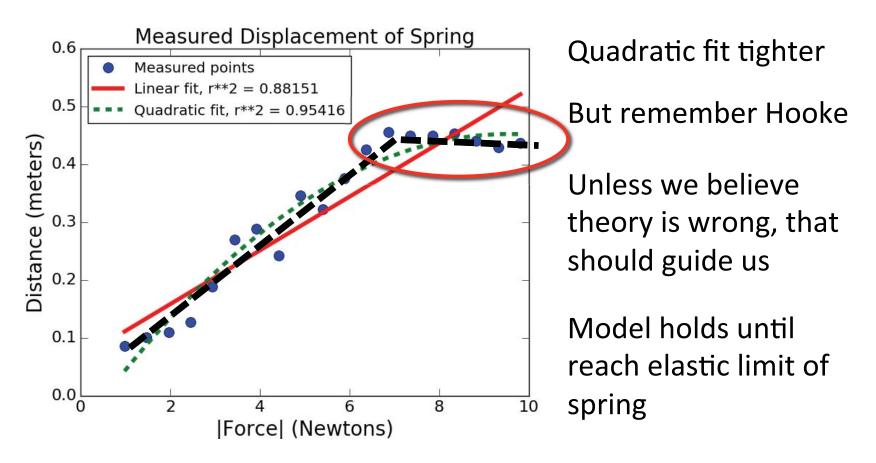
- In absence of theory predicting order of model, can engage in a search process
  - Fit a low order model to training data
  - Test on new data and record R<sup>2</sup> value
  - Increase order of model and repeat
  - Continue until fit on test data begins to decline





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## Returning to Where We Started



Should probably fit different models to different segments of data Can visualize as search process – find best place to break into two parts, such that both linear segments have high R<sup>2</sup> fits

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## Suppose We Don't Have a Solid Theory

- Use cross-validation results to guide the choice of model complexity
- •If dataset small, use leave-one-out cross validation
- If dataset large enough, use k-fold cross validation or repeated-random-sampling validation

### Leave-one-out Cross Validation

```
Let D be the original data set
testResults = []
for i in range(len(D)):
    training = D[:].pop(i)
    model = buildModel(training)
    testResults.append(test(model, D[i]))
Average testResults
   k-fold very similar
   Applies when we have large amount of data
   D partitioned into k equal size sets
   Model trained on k-1 sets, and tested on remaining set
```

# Repeated Random Sampling

```
Let D be the original data set
    n be the number of random samples
        usually n between 20% and 50%
    k be number of trials
testResults = []
for i in range(k)
    randomly select n elements for testSet,
       keep rest for training
    model = buildModel(training)
    testResults.append(test(model, testSet))
Average testResults
```

### An Example, Temperature By Year

- ■Task: Model how the mean daily high temperature in the U.S. varied from 1961 through 2015
- Get means for each year and plot them
- Randomly divide data in half n times
  - For each dimensionality to be tried
    - Train on one half of data
    - Test on other half
    - Record r-squared on test data
- Report mean r-squared for each dimensionality

## A Boring Class

```
class tempDatum(object):
    def __init__(self, s):
        info = s.split(',')
        self.high = float(info[1])
        self.year = int(info[2][0:4])
    def getHigh(self):
        return self.high
    def getYear(self):
        return self.year
```

### Read Data

```
def getTempData():
    inFile = open('temperatures.csv')
    data = []
    for l in inFile:
        data.append(tempDatum(l))
    return data
```

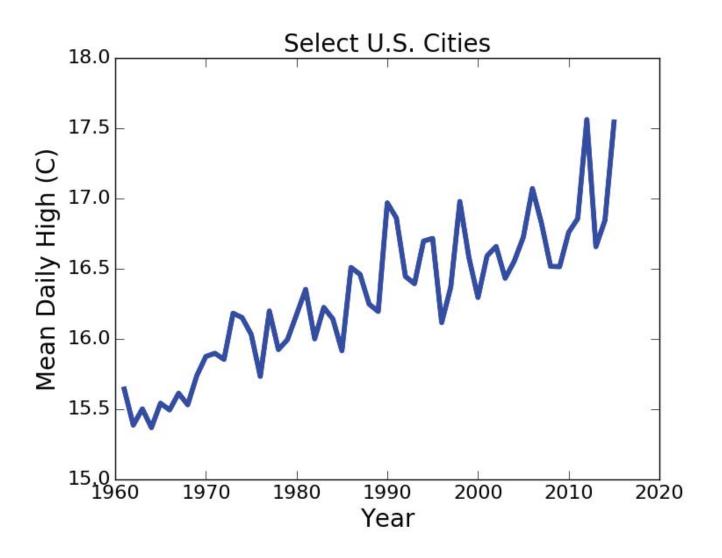
#### **Get Means**

```
def getYearlyMeans(data):
    years = {}
    for d in data:
        try:
        years[d.getYear()].append(d.getHigh())
        except:
        years[d.getYear()] = [d.getHigh()]
    for y in years:
        years[y] = sum(years[y])/len(years[y])
    return years
```

### Get and Plot Data

```
data = getTempData()
years = getYearlyMeans(data)
xVals, yVals = [], []
for e in years:
        xVals.append(e)
        yVals.append(years[e])
pylab.plot(xVals, yVals)
pylab.xlabel('Year')
pylab.ylabel('Mean Daily High (C)')
pylab.title('Select U.S. Cities')
```

### The Whole Data Set



## **Initialize Things**

```
numSubsets = 10
dimensions = (1, 2, 3, 4)
rSquares = {}
for d in dimensions:
    rSquares[d] = []
```

## Split Data

```
def splitData(xVals, yVals):
    toTrain = random.sample(range(len(xVals)),
                             len(xVals)//2)
    trainX, trainY, testX, testY = [],[],[],[]
    for i in range(len(xVals)):
        if i in toTrain:
            trainX.append(xVals[i])
            trainY.append(yVals[i])
        else:
            testX.append(xVals[i])
            testY.append(yVals[i])
    return trainX, trainY, testX, testY
```

## Train, Test, and Report

```
for f in range(numSubsets):
    trainX, trainY, testX, testY = splitData(xVals, yVals)
    for d in dimensions:
        model = pylab.polyfit(trainX, trainY, d)
        #estYVals = pylab.polyval(model, trainX)
        estYVals = pylab.polyval(model, testX)
        rSquares[d].append(rSquared(testY, estYVals))
print('Mean R-squares for test data')
for d in dimensions:
    mean = round(sum(rSquares[d])/len(rSquares[d]), 4)
    sd = round(numpy.std(rSquares[d]), 4)
    print('For dimensionality', d, 'mean =', mean,
          'Std =', sd)
```

### Results

```
Mean R-squares for test data For dimensionality 1 mean = 0.7535 Std = 0.0656 For dimensionality 2 mean = 0.7291 Std = 0.0744 For dimensionality 3 mean = 0.7039 Std = 0.0684 For dimensionality 4 mean = 0.7169 Std = 0.0777
```

- Line seems to be the winner
  - Highest average r-squared
  - Smallest deviation across trials
  - Simplest model

## Why we should run multiple sets

- Note that deviations are a decimal order of magnitude smaller than means
  - Suggests that while there is good agreement, deviations are large enough there could be a noticeable range of variation across trials
- Suppose we had just run one trial
  - Here are the R<sup>2</sup> values for each trial of linear fit
    - [0.7828002156420516, 0.80637964025052067, 0.79637132757274265, 0.78433885743211906, 0.76001112024853124, 0.57088936507035748, 0.72115408562589023, 0.74358276762149023, 0.79031455375148507, 0.77920238586399471]
  - If we had only run one split, and happened to get this result, we might have reached a different conclusion about validity of linear model

## Wrapping Up Curve Fitting

- •We can use linear regression to fit a curve to data
  - Mapping from independent values to dependent values
- •That curve is a model of the data that can be used to predict the value associated with independent values we haven't seen (out of sample data)
- R-squared used to evaluate model
  - Higher not always "better" because of risk of over fitting
- Choose complexity of model based on
  - Theory about structure of data
  - Cross validation
  - Simplicity

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