

# Unit 4

# Model Order Selection

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EE-UY 4563/EL-GY 9143: INTRODUCTION TO MACHINE LEARNING  
PROF. SUNDEEP RANGAN (WITH MODIFICATION BY YAO WANG)

# Learning Objectives

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- ❑ Compute the model order for a given model class
- ❑ Visually identify overfitting and underfitting of a model in a scatterplot
- ❑ Determine if there is under-modeling for a given true function and model class
- ❑ Compute the bias and variance for linear models (advanced)
- ❑ Perform cross-validation for selecting an optimal order selection

# Outline

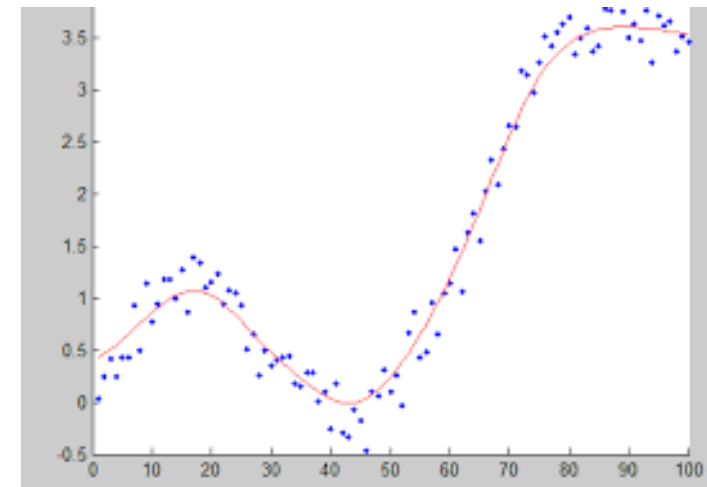
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 Motivating Example: What polynomial degree should a model use?

- ☐ Bias and variance
- ☐ Bias and variance for linear models (Advanced)
- ☐ Cross-validation


# Polynomial Fitting



- Last lecture: polynomial regression
- Given data  $(x_i, y_i), i = 1, \dots, N$
- Learn a polynomial relationship:
$$y = \beta_0 + \beta_1 x + \dots + \beta_d x^d + \epsilon$$
  - $d$  = degree of polynomial. Called **model order**
  - $\boldsymbol{\beta} = (\beta_0, \dots, \beta_d)$  = coefficient vector
- Given  $d$ , can find  $\boldsymbol{\beta}$  via least squares
- How do we select  $d$  from data?
- This problem is called **model order selection**.



# Demo on Github

□ Demo on github: [https://github.com/sdrangan/introml/blob/master/unit04\\_model\\_sel/demo1\\_polyfit.ipynb](https://github.com/sdrangan/introml/blob/master/unit04_model_sel/demo1_polyfit.ipynb)

□  GitHub, Inc. [US] | [https://github.com/sdrangan/introml/blob/master/model\\_sel/polyfit.ipynb](https://github.com/sdrangan/introml/blob/master/model_sel/polyfit.ipynb)

Suggested Sites  Web Slice Gallery  Import to Mendeley

## Demo: Polynomial Model Order Selection

In this demo, we will illustrate the process of cross-validation for model order selection. We derive data for a polynomial fit. The lab will demonstrate how to:

- Characterize the model order for a simple polynomial model
- Measure training and test error for a given model order
- Select a suitable model order using cross-validation
- Plot the results for the model order selection process

We first load the packages as usual.

```
In [2]: import numpy as np
import matplotlib
import matplotlib.pyplot as plt
from sklearn import datasets, linear_model, preprocessing
%matplotlib inline
```

## Polynomial Data

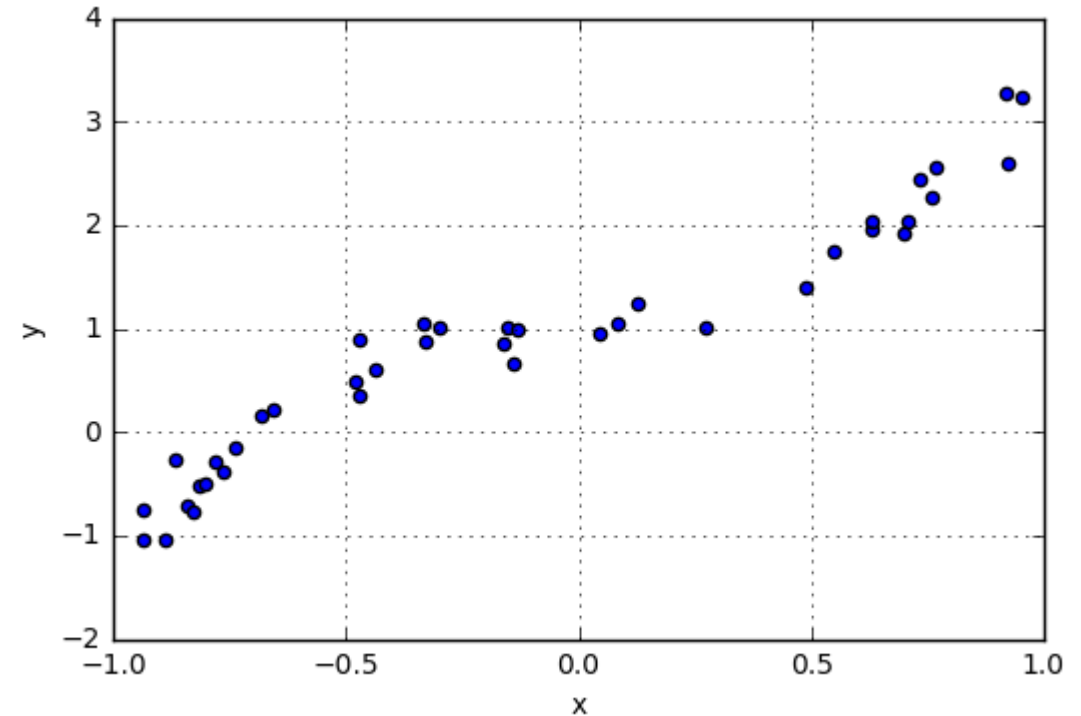
To illustrate the concepts, we consider a simple polynomial model:

$$y = \beta_0 + \beta_1 x + \dots + \beta_d x^d + \epsilon,$$

where  $d$  is the polynomial degree. We first generate synthetic data for this model.

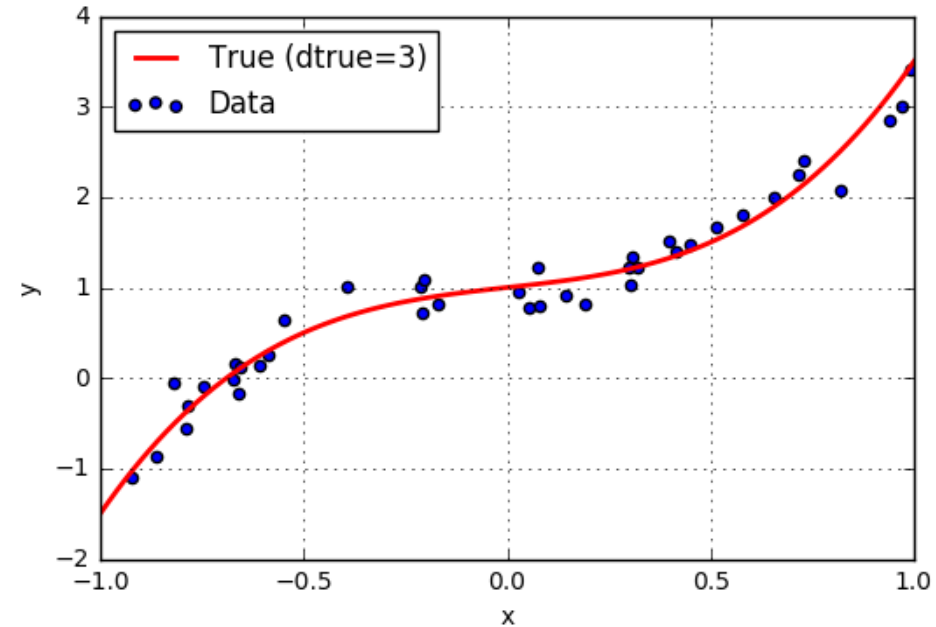
# Example Question

- ❑ You are given some data.
- ❑ Want to fit a model:  $y \approx f(x)$
- ❑ Decide to use a polynomial:  
$$f(x) = \beta_0 + \beta_1 x + \cdots + \beta_d x^d$$
- ❑ What model order  $d$  should we use?
- ❑ Thoughts?



# Synthetic Data

- Previous example is synthetic data
- $x_i$ : 40 samples uniform in  $[-1,1]$
- $y = f(x) + \epsilon$ ,
  - $f(x) = \beta_0 + \beta_1 x + \dots + \beta_d x^d = \text{"true relation"}$
  - $d = 3$ ,  $\epsilon \sim N(0, \sigma^2)$
- Synthetic data useful for analysis
  - Know "ground truth"
  - Can measure performance of various estimators



```
# Import useful polynomial library
import numpy.polynomial.polynomial as poly

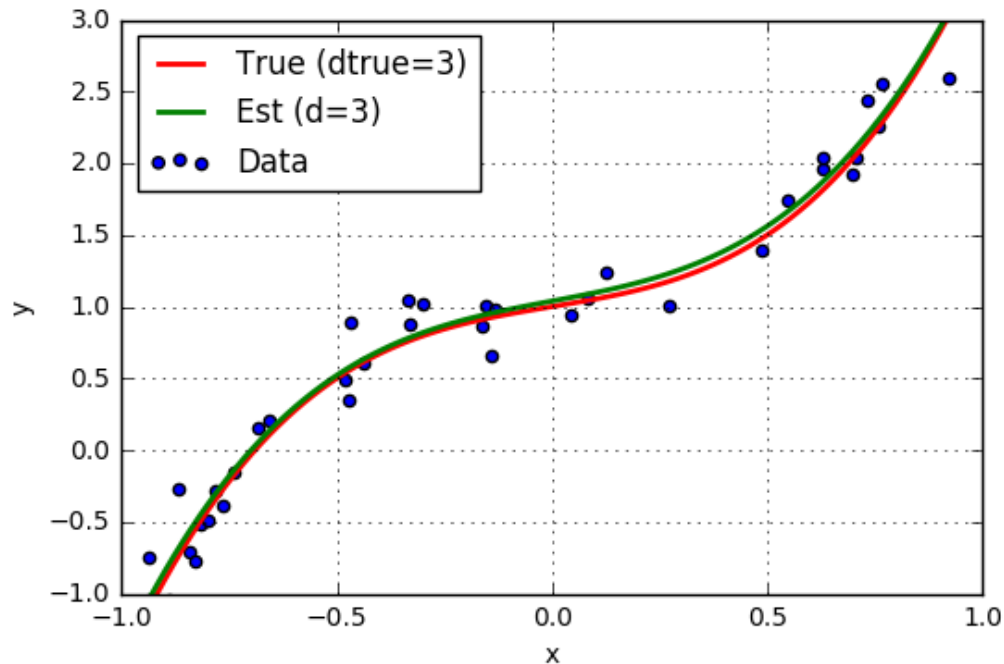
# True model parameters
beta = np.array([1,0.5,0,2]) # coefficients
wstd = 0.2 # noise
dtrue = len(beta)-1 # true poly degree

# Independent data
nsamp = 40
xdat = np.random.uniform(-1,1,nsamp)

# Polynomial
y0 = poly.polyval(xdat,beta)
ydat = y0 + np.random.normal(0,wstd,nsamp)
```

# Fitting with True Model Order

- Suppose true polynomial order,  $d=3$ , is known
- Use linear regression
  - `numpy.polynomial` package
- Get very good fit



```
d = 3
beta_hat = poly.polyfit(xdat,ydat,d)

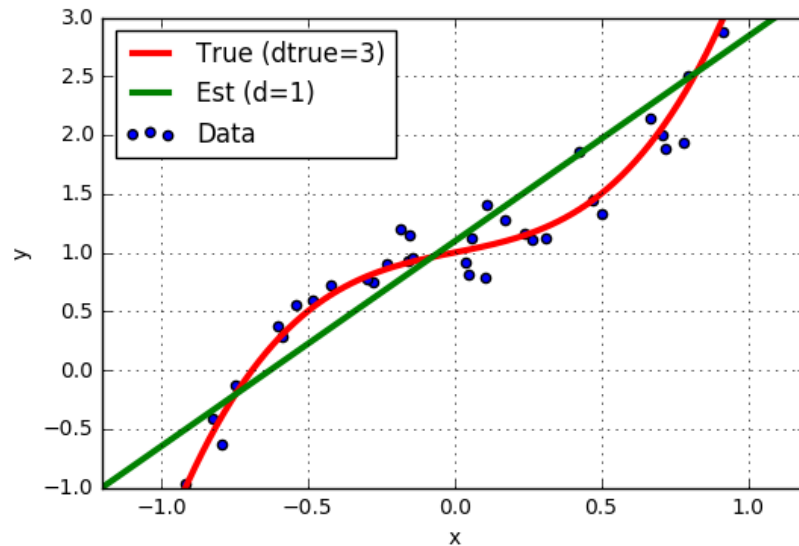
# Plot true and estimated function
xp = np.linspace(-1,1,100)
yp = poly.polyval(xp,beta)
yp_hat = poly.polyval(xp,beta_hat)
plt.xlim(-1,1)
plt.ylim(-1,3)
plt.plot(xp,yp,'r-',linewidth=2)
plt.plot(xp,yp_hat,'g-',linewidth=2)

# Plot data
plt.scatter(xdat,ydat)
plt.legend(['True (dtrue=3)', 'Est (d=3)', 'Data'], loc='upper left')
plt.grid()
plt.xlabel('x')
plt.ylabel('y')
```

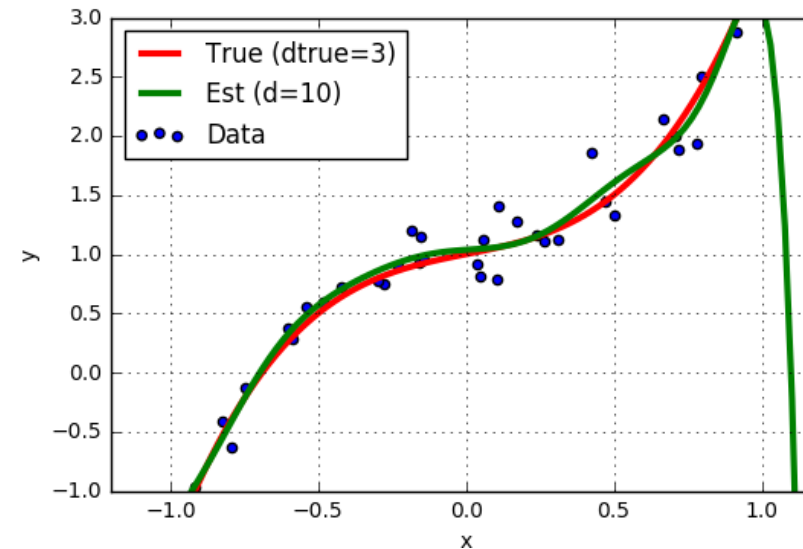


# But, True Model Order not Known

□ Suppose we guess the wrong model order?

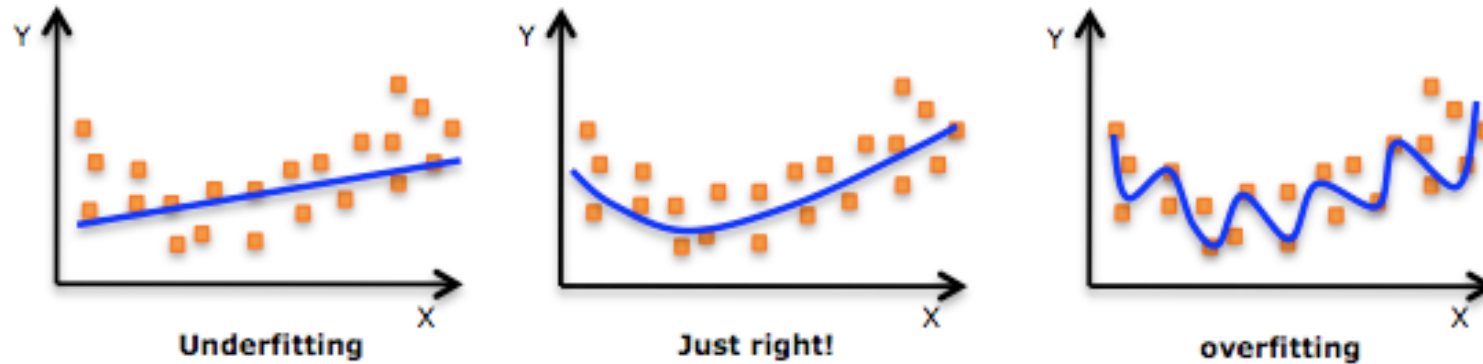


$d=1$  “Underfitting”



$d=10$  “Overfitting”

# How Can You Tell from Data?



- ❑ Is there a way to tell what is the correct model order to use?
- ❑ Must use the data. Do not have access to the true  $d$ ?
- ❑ What happens if we guess:
  - $d$  too big?
  - $d$  too small?

# Using RSS on Training Data?

## ❑ Simple (but bad) idea:

- For each model order,  $d$ , find estimate  $\hat{\beta}$
- Compute predicted values on training data

$$\hat{y}_i = \hat{\beta}^T x_i$$

- Compute RSS

$$RSS(d) = \sum_i (y_i - \hat{y}_i)^2$$

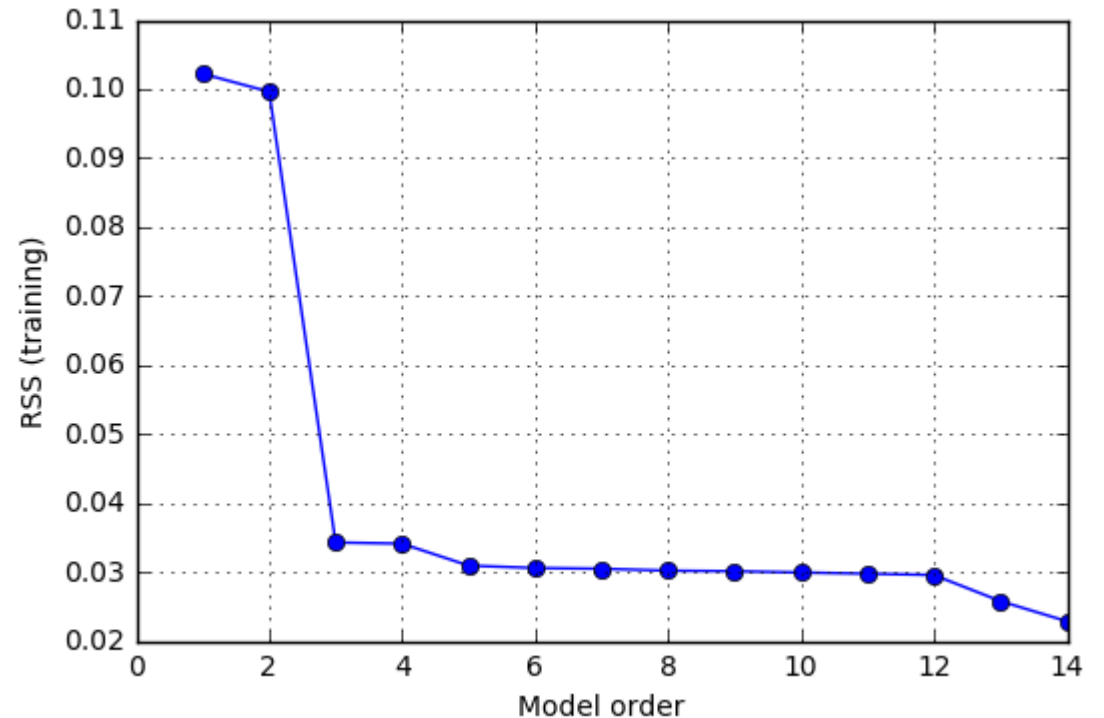
- Find  $d$  with lowest  $RSS$

## ❑ This doesn't work

- $RSS(d)$  is always decreasing (Question: Why?)
- Minimizing  $RSS(d)$  will pick  $d$  as large as possible
- Leads to overfitting

## ❑ What went wrong?


## ❑ How do we do better?



# Outline

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- ☐ Motivating Example: What polynomial degree should a model use?

-  ☐ Bias and variance

- ☐ Bias and variance for linear models (Advanced)

- ☐ Cross-validation

# Generalization

## Machine learning:

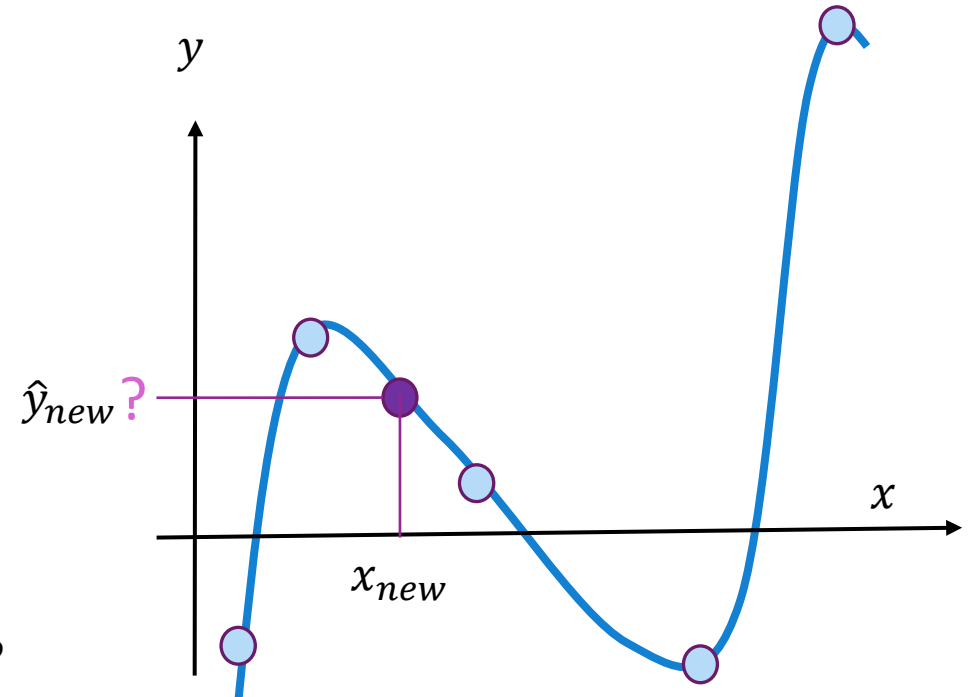
- Get **data** points  $(x_i, y_i), i = 1, \dots, n$
- **Learn** some function  $\hat{y} = f(x)$

## Implicitly, we are

- **Inferring** the value of  $y$  at **new** values of  $x$
- Called **generalization**

## Basic question for all ML:

- *How well do models we train generalize to new samples?*



# A Model To Understand Generalization

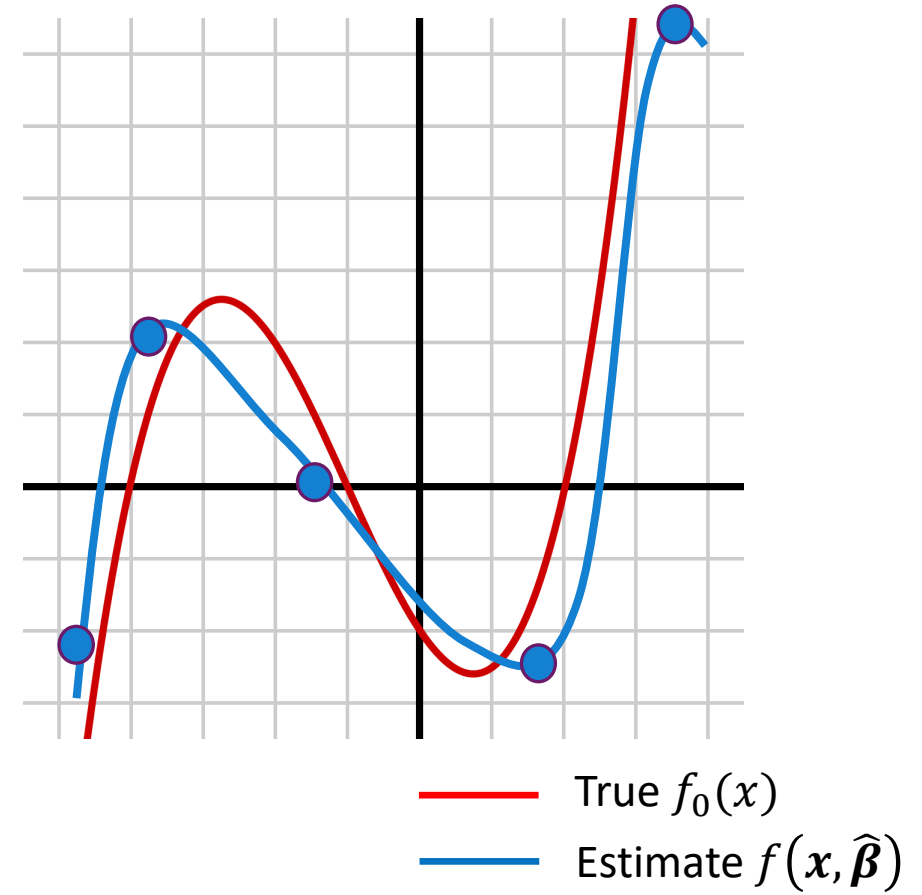
□ Assume a **true** relation:  $y = f_0(x) + \epsilon$ ,  $\epsilon \sim N(0, \sigma_\epsilon^2)$

□ Get **data** points  $(x_i, y_i), i = 1, \dots, n$   
 $y_i = f_0(x_i) + \epsilon_i$

□ Assume a **model**  $\hat{y} = f(x, \beta)$   
◦ Parameters  $\beta$

□ **Fit** a parameter  $\hat{\beta}$  from training data  
◦ Results in estimated function  $f(x, \hat{\beta})$

□ **Question**: How “good” is the estimated function?



# Output Mean Squared Error

□ To evaluate generalization, suppose we are given:

- A **test point**  $\mathbf{x}_{test}$
- New point, generally different from training samples.

□ Actual value:  $y = f_0(\mathbf{x}_{test}) + \epsilon$

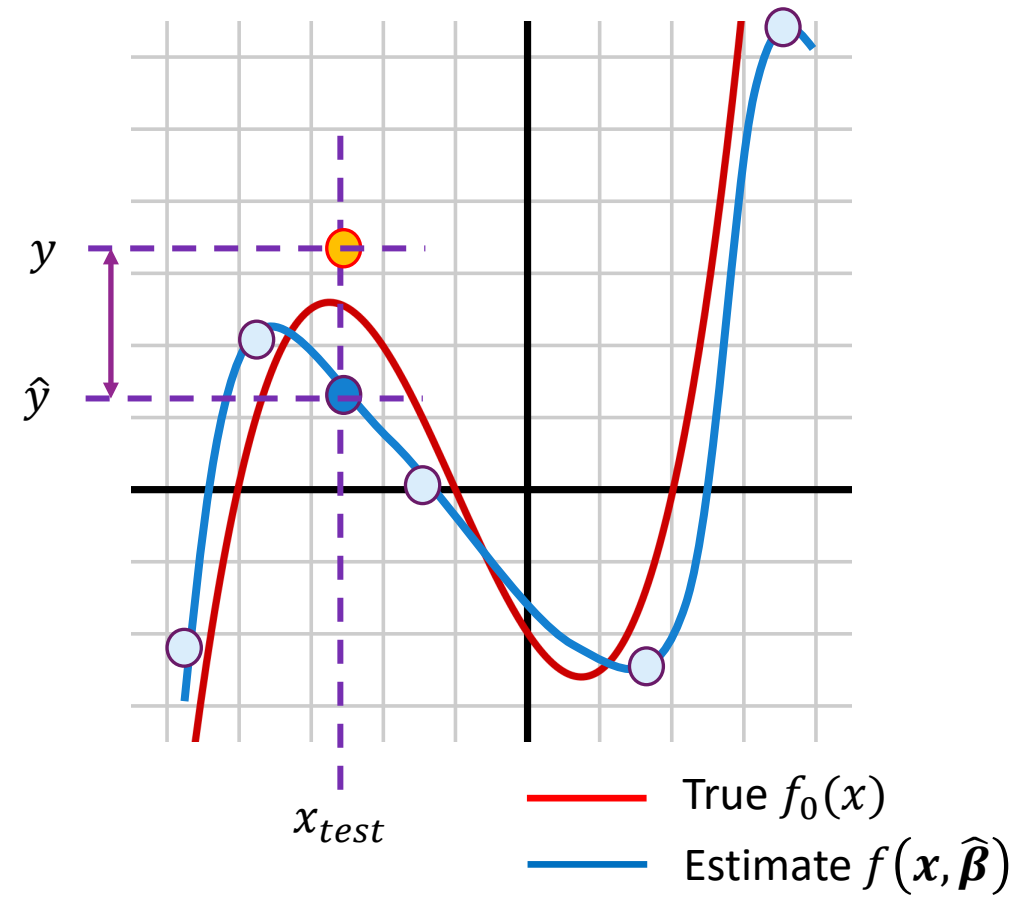
□ Predicted value:  $\hat{y} = f(\mathbf{x}_{test}, \hat{\beta})$

- Note that  $\hat{\beta}$  is random due to noise in training data

□ Define **output mean squared error** :

$$MSE_y(\mathbf{x}_{test}) := E[y - \hat{y}]^2$$

- Expectation is over noise  $\epsilon$  on the training and test data.



# Function MSE and Irreducible Error

□ **Output MSE decomposition:** Output MSE,  $MSE_y(\mathbf{x}_{test}) := E[y - \hat{y}]^2$ , satisfies:

$$MSE_y(\mathbf{x}_{test}) = MSE_f(\mathbf{x}_{test}) + \sigma_\epsilon^2$$

□ **Function MSE:**  $MSE_f(\mathbf{x}_{test}) = E[f_0(\mathbf{x}_{test}) - f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}})]^2$

- Represents difference between estimated and true function

□ **Irreducible error:**  $\sigma_\epsilon^2 = E(\epsilon^2)$  in output  $y = f_0(x) + \epsilon$

- Occurs since  $y$  is influenced by other factors than  $x$
- Fundamental limit on ability to predict  $y$
- Lower bound on  $MSE_y(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}}) \geq \sigma_\epsilon^2$



# Proof of the MSE Decomposition

□ **Output MSE decomposition:** Output MSE  $MSE_y(\mathbf{x}_{test}) := E[y - \hat{y}]^2$  is:

$$MSE_y(\mathbf{x}_{test}) = MSE_f(\mathbf{x}_{test}) + \sigma_\epsilon^2$$

□ **Proof:**

- $MSE_y(\mathbf{x}_{test}) := E[y - \hat{y}]^2 = E[f_0(\mathbf{x}_{test}) + \epsilon - f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}})]^2 = M_1 + M_2 + 2M_3$
- $M_1 = E[f_0(\mathbf{x}_{test}) - f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}})]^2 = MSE_f(\mathbf{x}_{test})$
- $M_2 = E[\epsilon^2] = \sigma^2$
- Noise on test sample is independent of  $\hat{\boldsymbol{\beta}}$  and  $\mathbf{x}_{test}$  and  $E(\epsilon) = 0$
- Therefore  $M_3 = E[\epsilon(f_0(\mathbf{x}_{test}) - f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}}))] = E(\epsilon)E[f_0(\mathbf{x}_{test}) - f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}})] = 0$

# Model Class and Under-Modeling

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❑ **Model class:** The set of possible functions,  $\hat{y} = f(x, \beta)$

- Set is parametrized by  $\beta$

❑ **Definition:** A true function  $f_0(x)$  is in the model class  $\hat{y} = f(x, \beta)$  if:

$$f_0(x) = f(x, \beta_0) \text{ for all } x$$

for some parameter  $\beta_0$ .

- $\beta_0$  called the true parameter

❑ **Under-modeling:** When  $f_0(x)$  is not in the model class

# Sample Questions 1 and 2

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□ For each pair, state if the true function is in the model class or not

- That is, is there under-modeling or not?
- If true function is in the model class, state the true parameter

□ Ex 1:

- True function:  $f_0(x) = 2 + 3x$  Model class:  $f(x, \beta) = \beta_0 + \beta_1x + \beta_2x^2$
- **No under-modeling**. True parameter:  $\beta = (2, 3, 0)$

□ Ex 2:

- True function:  $f_0(x) = 2 + 3x + 4x^2$  Model class:  $f(x, \beta) = \beta_0 + \beta_1x$
- **There is under-modeling**. Model class does not contain  $x^2$  term

# Sample Questions 3 and 4

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□ For each pair, state if the true function is in the model class or not

- That is, is there under-modeling or not?
- If true function is in the model class, state the true parameter

□ Ex 3:

- True function:  $f_0(x) = \sin(2\pi(5)x + 7)$  Model class:  $f(x, \beta) = \beta_0 \sin(2\pi(5)x) + \beta_1 \cos(2\pi(5)x)$
- **No under-modeling.**  $f_0(x) = \sin(2\pi(5)x + 7) = \sin(2\pi(5)x) \cos(7) + \cos(2\pi(5)x) \sin(7)$
- True parameter  $\beta = (\cos 7, \sin 7)$

□ Ex 4:

- True function:  $f_0(x) = \sin(2\pi(8)x + 7)$  Model class:  $f(x, \beta) = \beta_0 \sin(2\pi(5)x) + \beta_1 \cos(2\pi(5)x)$
- **There is under-modeling.** Model class does not contain  $\sin(2\pi 8)$  or  $\cos(2\pi 8)$  terms

# Under-Modeling and Irreducible Error

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□ Suppose that:

- There is no under-modeling:  $f_0(\mathbf{x}) = f(\mathbf{x}, \boldsymbol{\beta}_0)$  for some “true” parameter  $\boldsymbol{\beta}_0$ ; and
- Estimator selects the true parameter  $\hat{\boldsymbol{\beta}} = \boldsymbol{\beta}_0$  (somehow?)

□ Then, function MSE is zero:

$$MSE_f(\mathbf{x}_{test}) = E[f_0(\mathbf{x}_{test}) - f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}})]^2 = E[f_0(\mathbf{x}_{test}) - f(\mathbf{x}_{test}, \boldsymbol{\beta}_0)]^2 = 0$$

□ Output MSE = irreducible error

$$MSE_y(\mathbf{x}_{test}) := MSE_f(\mathbf{x}_{test}) + \sigma_\epsilon^2 = \sigma_\epsilon^2$$

# What We Have Learned So Far

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## □ If (A Big If!)

- There is no undermodeling (i.e. true function is in model class), and
- We can estimate the true parameter

## □ Then:

- Output error = irreducible error
- We can achieve the same error as if we knew the true function  $f_0(x)$

## □ This suggests: *Select the model class large!*

- Guarantees to approximately contains true function
- Ex: Take model class = set of polynomials with very high degree

## □ But, using large models has other problems...

# In-Class Exercise

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## Question

For each model and true function pair below: Determine if there is undermodeling (i.e. the true function is in the model class). If there is no undermodeling, find the true parameters

- (a) True function:  $f_0(x) = (1 + 2x)(3 + 4x)$ , Model:  $f(x, \beta) = \beta_0 + \beta_1 x + \beta_2 x^2$
- (b) True function:  $f_0(t) = 2(1 - e^{t-3})$  Model:  $f(t, \beta) = a + be^{ct}, \beta = (a, b, c)$
- (c) True function:  $f_0(t) = 2(1 - e^{t-3})$  Model:  $f(t, \beta) = a + be^{-t}, \beta = (a, b)$

# Bias and Variance

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❑ To understand potential problem of using a large model class introduce two key quantities:

❑ **Bias:**  $Bias(\mathbf{x}_{test}) := f_0(\mathbf{x}_{test}) - E[f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}})]$

- How much the average value of the estimate differs from the true function

❑ **Variance:**  $Var(\mathbf{x}_{test}) := E \left[ f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}}) - E[f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}})] \right]^2$

- How much the estimate varies around its average

❑ Bias and variance are (conceptually) measured as follows:

- Get many independent training data sets, each with same size  $N$  and input values  $\mathbf{x}_i$
- Each dataset has different output values  $y_i$  because of independent noise in the training data
- Obtain  $\hat{\boldsymbol{\beta}}$  for each training data set
- Bias and variances are computed over the different sets

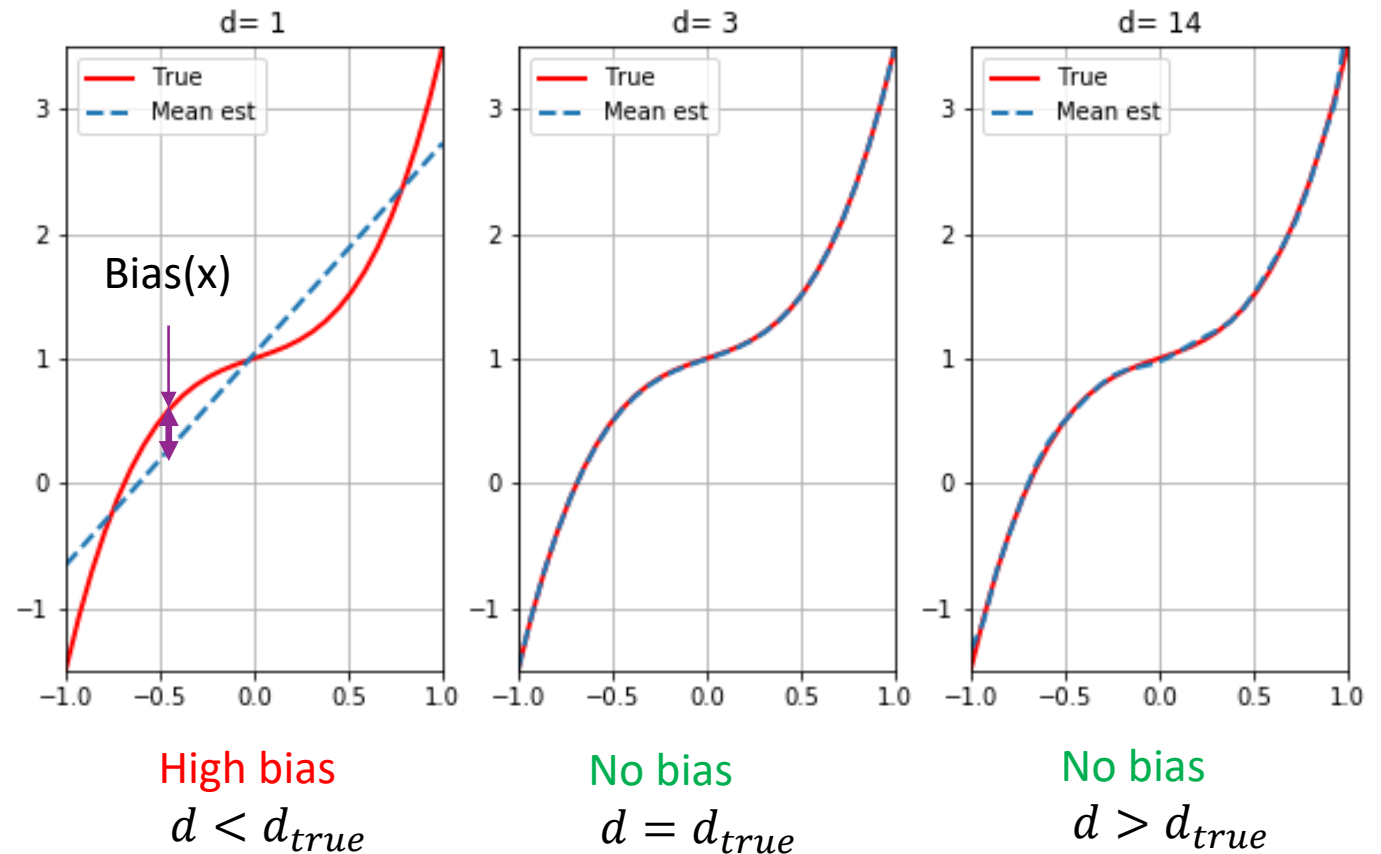
❑ Of course, in reality, we have only one training dataset

- Used to study theoretical averages over different experiments



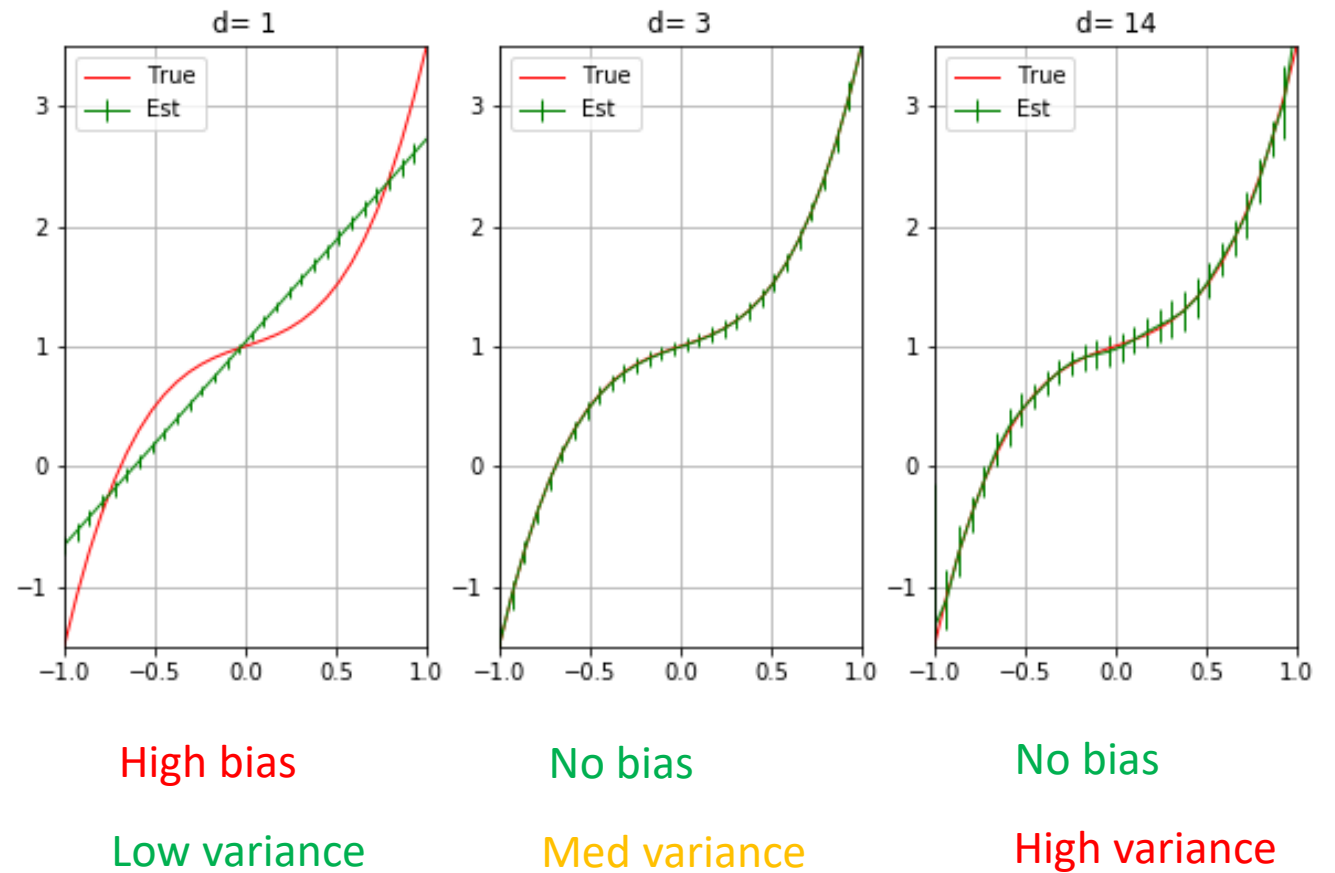
# Bias Illustrated

- Red: True function
- Repeat 100 trials
  - Each trial has independent data
  - Obtain estimate for each trial
- Dashed line: Mean estimate among all trials
- Bias = True – Mean estimate
- Conclusions:
  - Low model orders  $\Rightarrow$  bias high
  - High model orders  $\Rightarrow$  bias low



# Variance Illustrated

- ❑ Red: True function
- ❑ Repeat 100 trials
  - Each trial has independent data
  - Obtain estimate for each trial
- ❑ Variance=STD around mean
- ❑ Conclusions:
  - Low model orders  $\Rightarrow$  low variance
  - High model orders  $\Rightarrow$  high variance



# Bias-Variance Formula

## □ Recall definitions:

- **Function MSE:**  $MSE_f(\mathbf{x}_{test}) := E[f_0(\mathbf{x}_{test}) - f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}})]:$
- **Bias:**  $Bias(\mathbf{x}_{test}) := f_0(\mathbf{x}_{test}) - E[f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}})]$
- **Variance:**  $Var(\mathbf{x}_{test}) := E[f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}}) - E[f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}})]]^2$

## □ Bias-Variance formula : $MSE_f(\mathbf{x}_{test}) = Bias(\mathbf{x}_{test})^2 + Var(\mathbf{x}_{test})$

- Will be proved below

## □ Bias-Variance tradeoff

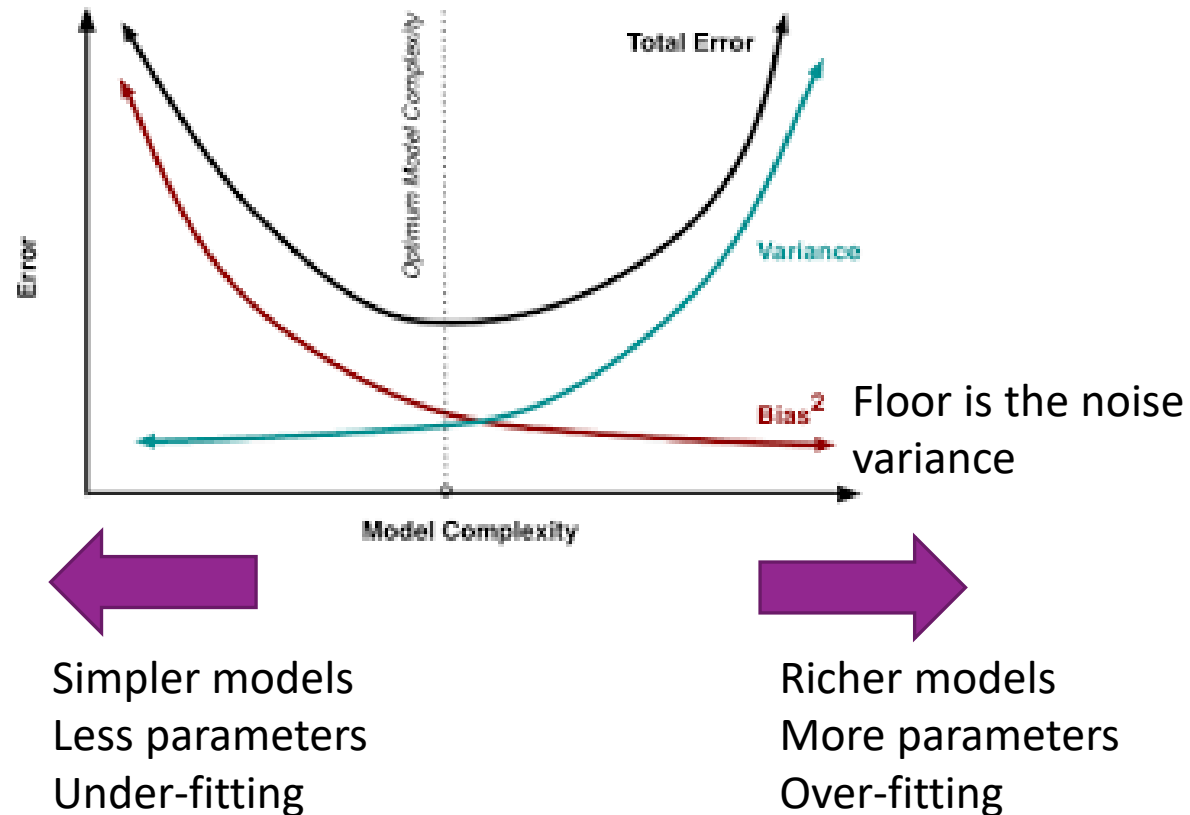
### □ Bias due to under-modeling

- Reduced with high model order

### □ Variance is due to noise in training data and number of parameters to estimate

- Increases with higher model order

# Bias-Variance Tradeoff



## □ Bias:

- Due to under-modeling
- Reduced with high model order

## □ Variance:

- Increases with noise in training data
- Increase with high model order

## □ Optimal model order depends on:

- Amount of samples available
- Underlying complexity of the relation

# Bias-Variance Formula Proof

- Define  $\bar{f}(\mathbf{x}_{test}) = E[f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}})]$  = average value of estimated function
- $MSE_f(\mathbf{x}_{test}) = E[f_0(\mathbf{x}_{test}) - f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}})]^2 = E[f_0(\mathbf{x}_{test}) - \bar{f}(\mathbf{x}_{test}) + \bar{f}(\mathbf{x}_{test}) - f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}})]^2$
- Three components:  $MSE_f(\mathbf{x}_{test}) = M_1 + M_2 - 2M_3$ 
  - $M_1 = E[f_0(\mathbf{x}_{test}) - \bar{f}(\mathbf{x}_{test})]^2 = [f_0(\mathbf{x}_{test}) - \bar{f}(\mathbf{x}_{test})]^2 = Bias(\mathbf{x}_{test})$
  - $M_2 = E[f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}}) - \bar{f}(\mathbf{x}_{test})]^2 = Var(\mathbf{x}_{test})$
  - $M_3 = E[(f_0(\mathbf{x}_{test}) - \bar{f}(\mathbf{x}_{test}))(f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}}) - \bar{f}(\mathbf{x}_{test}))]$   
 $= (f_0(\mathbf{x}_{test}) - \bar{f}(\mathbf{x}_{test}))E[f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}}) - \bar{f}(\mathbf{x}_{test})]$   
 $= (f_0(\mathbf{x}_{test}) - \bar{f}(\mathbf{x}_{test}))(\bar{f}(\mathbf{x}_{test}) - \bar{f}(\mathbf{x}_{test})) = 0$

# Summary of Results for Linear Models

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- ❑ Suppose model is linear with  $N$  = num samples,  $p$  = num parameters
- ❑ Can show the following results (need some math, see next section)
- ❑ Result 1: When  $N < p$ , linear estimate is not unique
  - Need at least as many samples as parameters
- ❑ Now assume that  $N \geq p$  and parameter estimate is unique
- ❑ Result 2: When there is no under-modeling, estimate is unbiased
$$E[f(x_{test}, \hat{\beta})] = f_0(x_{test}).$$
- ❑ Result 3: If test point drawn randomly from the training data:
$$Var = \frac{p}{N} \sigma_{\epsilon}^2$$
  - Variance increases linearly with number of parameters and inversely with number of samples

# In-Class Exercise

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## Question

A teacher tries to model the score of a child on a test as follows:


$$\text{Test score} \approx \beta_0 + \beta_1[\text{Hours studied}] + \beta_2[\text{Hours sleep before test}]$$

Below are three possible shortcomings of the model. For each shortcoming, indicate whether it would lead to (i) High irreducible error, (ii) High bias error or (iii) High variance error

- (a) The relation between test score and hours studied is not linear. After some amount of studying, the students test score will no longer improve.
- (b) The model should also include the age of the child.
- (c) The model was fit with data from one class with only 15 children.

# Outline

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- ❑ Motivating Example: What polynomial degree should a model use?
- ❑ Bias and variance
-  ❑ Bias and variance for linear models (Advanced)
- ❑ Cross-validation



# This Section is Advanced

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- ❑ This section requires more advanced probability and linear algebra
- ❑ Means and variances of random vectors
- ❑ Undergraduates: Skip to final slide for final conclusions
- ❑ Graduate students: We will cover this
  - You should review your multi-variable probability and linear algebra

# Linear Models

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- Consider linear model in general transformed feature space:

$$\hat{y} = f(x, \beta) = \phi(x)^T \beta = \beta_1 \phi_1(x) + \cdots + \beta_p \phi_p(x)$$

- See previous lecture

- Assume true data relation is:  $y = f_0(x) + \epsilon$ ,  $E(\epsilon) = 0$ ,  $E(\epsilon^2) = \sigma^2$

- When there is no under-modeling:  $f_0(x) = f(x, \beta^0) = \phi(x)^T \beta^0$

- $\beta^0 = (\beta_0^0, \dots, \beta_k^0)$  True parameter

- Get data  $(x_i, y_i)$ ,  $i = 1, \dots, N$

- Least squares fit  $\hat{\beta} = (A^T A)^{-1} A^T y$

$$A = \begin{bmatrix} \phi_1(x_1) & \cdots & \phi_p(x_1) \\ \vdots & \vdots & \vdots \\ \phi_1(x_N) & \cdots & \phi_p(x_N) \end{bmatrix}$$

# Minimum Number of Samples

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- ❑ LS estimate requires  $A^T A$  is invertible.
- ❑ Linear algebra fact: Since  $A \in R^{N \times p}$ , we need  $\text{Rank}(A) \geq p$ 
  - Otherwise solution is not unique
- ❑ Since  $\text{Rank}(A) \leq \min(N, p)$  we need  $N \geq p$ .
- ❑ Recall:
  - $N$  = number of data samples
  - $p$  = number of parameters
- ❑ **Conclusion:** Number of samples  $\geq$  number of parameters
- ❑ This places a basic limit on the model complexity that you can use

# Random Vectors Review

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- ❑ To analyze bias and variance in linear models, we need to review random vectors
- ❑ **Random vectors:**  $\mathbf{x} = (x_1, \dots, x_d)^T$ : Each component  $x_j$  is a random variable
- ❑ **Mean:** The vector of means of the components

$$\boldsymbol{\mu} = E\mathbf{x} = (Ex_1, \dots, Ex_d)^T = (\mu_1, \dots, \mu_d)^T$$

- ❑ **Covariance components:**  $\text{Cov}(x_i, x_j) = E[(x_i - \mu_i)(x_j - \mu_j)]$

- ❑ **Variance matrix** ( $d \times d$ ):

$$\text{Var}(\mathbf{x}) := E[(\mathbf{x} - \boldsymbol{\mu})(\mathbf{x} - \boldsymbol{\mu})^T] = \begin{bmatrix} \text{Cov}(x_1, x_1) & \cdots & \text{Cov}(x_1, x_d) \\ \vdots & \vdots & \vdots \\ \text{Cov}(x_d, x_1) & \cdots & \text{Cov}(x_d, x_d) \end{bmatrix}$$



# Linear Transforms of Random Vectors

---

- A linear transform is a map:  $y = Ax + b$
- $A \in R^{M \times N}$  maps input  $x \in R^N$  to  $Ax \in R^M$
- Mean and variance matrix under linear map given by
  - Mean:  $E(y) = AE(x) + b$
  - Variance:  $Var(y) = AVar(x)A^T$

# Bias With No Under-Modeling

- Suppose that there is no undermodeling:  $f_0(x) = \phi(x)^T \beta^0$
- Then each training sample output is:  $y_i = \phi(x_i)^T \beta^0 + \epsilon_i$
- Hence: true data vector  $y = A\beta^0 + \epsilon$
- Parameter estimate is:  
$$\hat{\beta} = (A^T A)^{-1} A^T y = (A^T A)^{-1} A^T (A\beta^0 + \epsilon) = \beta^0 + (A^T A)^{-1} A^T \epsilon$$
- Since  $E\epsilon = 0$ ,  $E\hat{\beta} = \beta^0$ . Average of parameter estimate matches true parameter
- $Ef(x_{test}, \hat{\beta}) = \phi(x_{test})^T E\hat{\beta} = \phi(x_{test})^T \beta^0 = f_0(x_{test})$
- Therefore  $Bias(x_{test}) := f_0(x_{test}) - Ef(x_{test}, \hat{\beta}) = 0$
- **Conclusion:** When the model is linear and there is no under-modeling, there is no bias

# Variance of the Parameters in Linear Models

□ Since  $\epsilon_i$  are independent for different samples with  $E\epsilon_i = 0$ ,  $E\epsilon_i^2 = \sigma^2$

$$\text{Cov}(\epsilon_i, \epsilon_j) = \begin{cases} 0 & i \neq j \\ \sigma^2 & i = j \end{cases}$$

□ Therefore variance matrix is:  $\text{Var}(\epsilon) = \sigma^2 I$

□ From last slide:  $\hat{\beta} = \beta^0 + (A^T A)^{-1} A^T \epsilon$ .

□ Applying variance formula of a linear transformation of  $\epsilon$

$$\begin{aligned} E \left( (\hat{\beta} - \beta^0)(\hat{\beta} - \beta^0)^T \right) &= (A^T A)^{-1} A^T \text{Var}(\epsilon) A (A^T A)^{-1} \\ &= \sigma^2 (A^T A)^{-1} A^T A (A^T A)^{-1} = \sigma^2 (A^T A)^{-1} \end{aligned}$$



# Variance in Linear Estimate

- To compute variance use trick: Suppose  $\mathbf{a}$  and  $\mathbf{z}$  are vectors,  $\mathbf{a}$  is non-random,  $\mathbf{z}$  is random:

$$E|\mathbf{a}^T \mathbf{z}|^2 = E(\mathbf{a}^T \mathbf{z} \mathbf{z}^T \mathbf{a}) = \mathbf{a}^T E(\mathbf{z} \mathbf{z}^T) \mathbf{a}$$

- From earlier:  $E f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}}) = \phi(\mathbf{x}_{test})^T E \hat{\boldsymbol{\beta}} = \phi(\mathbf{x}_{test})^T \boldsymbol{\beta}^0$

- Therefore variance of linear model:

$$\begin{aligned} \text{Var}(\mathbf{x}_{test}) &= E[f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}}) - E f(\mathbf{x}_{test}, \hat{\boldsymbol{\beta}})]^2 = E[\phi(\mathbf{x}_{test})^T (\hat{\boldsymbol{\beta}} - \boldsymbol{\beta}^0)]^2 \\ &= \phi(\mathbf{x}_{test})^T E[(\hat{\boldsymbol{\beta}} - \boldsymbol{\beta}^0)(\hat{\boldsymbol{\beta}} - \boldsymbol{\beta}^0)^T] \phi(\mathbf{x}_{test}) \\ &= \sigma^2 \phi(\mathbf{x}_{test})^T (\mathbf{A}^T \mathbf{A})^{-1} \phi(\mathbf{x}_{test}) \end{aligned}$$

- Above calculation is for the case of no under-modeling
- But, similar calculation shows variance expression is the same when there is under-modeling



# Case with Equal Test & Training Distributions

□ Suppose that test point is distributed identically to training data

- Training data inputs  $\mathbf{x}_i, i = 1, \dots, N$
- $\mathbf{x}_{test} = \mathbf{x}_i$  with probability  $\frac{1}{N}$

□ Since rows of  $A$  are  $\phi(\mathbf{x}_i)^T$ :  $A^T A = \sum_{i=1}^N \phi(\mathbf{x}_i) \phi(\mathbf{x}_i)^T$

□ Now use trick: For random vectors  $\mathbf{u}, \mathbf{v}$ :  $E(\mathbf{u}^T \mathbf{v}) = \text{Tr } E(\mathbf{v} \mathbf{u}^T)$

- $\text{Tr}(A) = \sum_i A_{ii}$  = sum of diagonals

□ Therefore, variance averaged over  $\mathbf{x}_{test}$  is:

$$E \text{Var}(\mathbf{x}_{test}) = \sigma^2 E[\phi(\mathbf{x}_{test})^T (A^T A)^{-1} \phi(\mathbf{x}_{test})] = \sigma^2 \text{Tr}\{E[\phi(\mathbf{x}_{test}) \phi(\mathbf{x}_{test})^T] (A^T A)^{-1}\}$$

$$= \frac{\sigma^2}{N} \text{Tr}\left\{\sum_i \phi(\mathbf{x}_i) \phi(\mathbf{x}_i)^T (A^T A)^{-1}\right\} = \frac{\sigma^2}{N} \text{Tr}\{(A^T A)(A^T A)^{-1}\} = \frac{\sigma^2}{N} \text{Tr}\{I_p\} = \frac{\sigma^2 p}{N}$$

# Case with Equal Test & Training Distributions

---

□ Assumption on previous slide: Test point  $\mathbf{x}_{test}$  is randomly selected from training data

□ Then, average variance is given by

$$E \text{Var}(\mathbf{x}_{test}) = \frac{\sigma^2 p}{N}$$

□ Increases with number of parameters  $p$

- Shows that increasing model complexity increases variance error

□ Decreases with number of samples  $N$

□ What if test data point is distributed differently from training data?

- Then variance may be much larger  $\frac{\sigma^2 p}{N}$
- If test data is not like training data, we are extending model to regions not seen in training data
- Often leads to high error

# Summary of Results for Linear Models

---

□ Suppose model is linear with  $N$  = num samples,  $p$  = num parameters

□ Result 1: When  $N < p$ , linear estimate is not unique

- Need at least as many samples as parameters

□ Now assume that  $N \geq p$  and parameter estimate is unique

□ Result 2: When there is no under-modeling, estimate is unbiased

$$E[f(x_{test}, \hat{\beta})] = f_0(x_{test}).$$

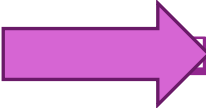
□ Result 3: If test point drawn from same distribution as training data:

$$Var = \frac{p}{n} \sigma_{\epsilon}^2$$

- Variance increases linearly with number of parameters and inversely with number of samples

# Outline

---

- ❑ Motivating Example: What polynomial degree should a model use?
- ❑ Bias and variance
- ❑ Bias and variance for linear models (Advanced)
-  ❑ Cross-validation

# Cross Validation

❑ **Key idea:** Evaluate on samples different from training

❑ Get data  $X, y$

❑ Split into training  $X_{tr}, y_{tr}$  and test  $X_{ts}, y_{ts}$

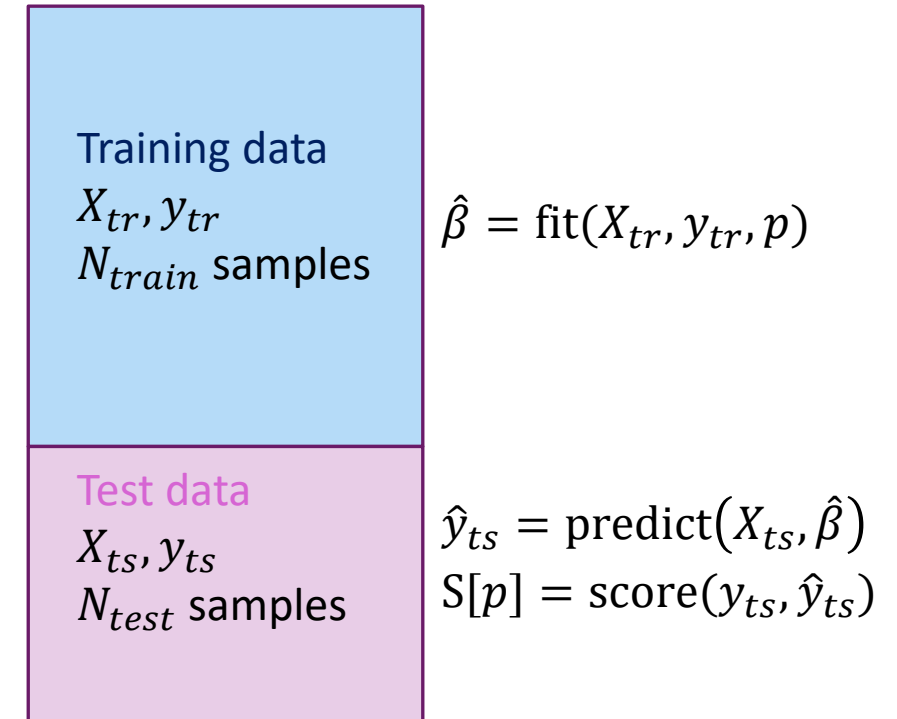
❑ For  $p = 1$  to  $p_{max}$  // Loop over model order

- **Fit** on training data with model order  $p$
- **Predict** values on test data
- **Score** fit on test data (e.g. measure RSS)

❑ Select model order with smallest score:

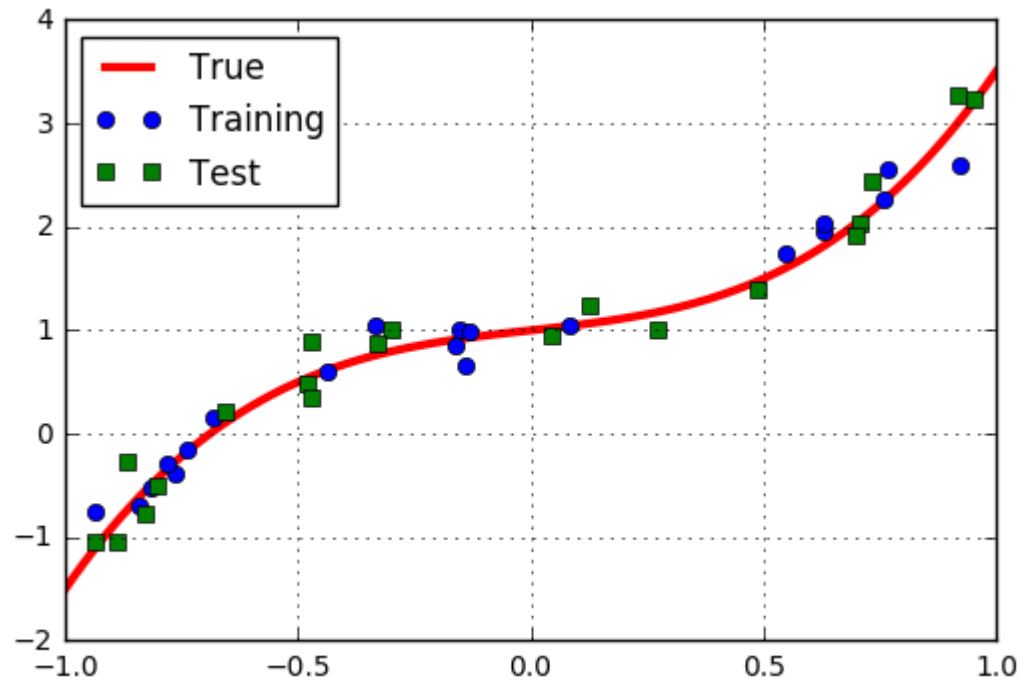
$$\hat{p} = \arg \min_p S[p]$$

- Maximize if higher score is better



# Polynomial Example: Training Test Split

□ Example: Split data into 20 samples for training, 20 for test



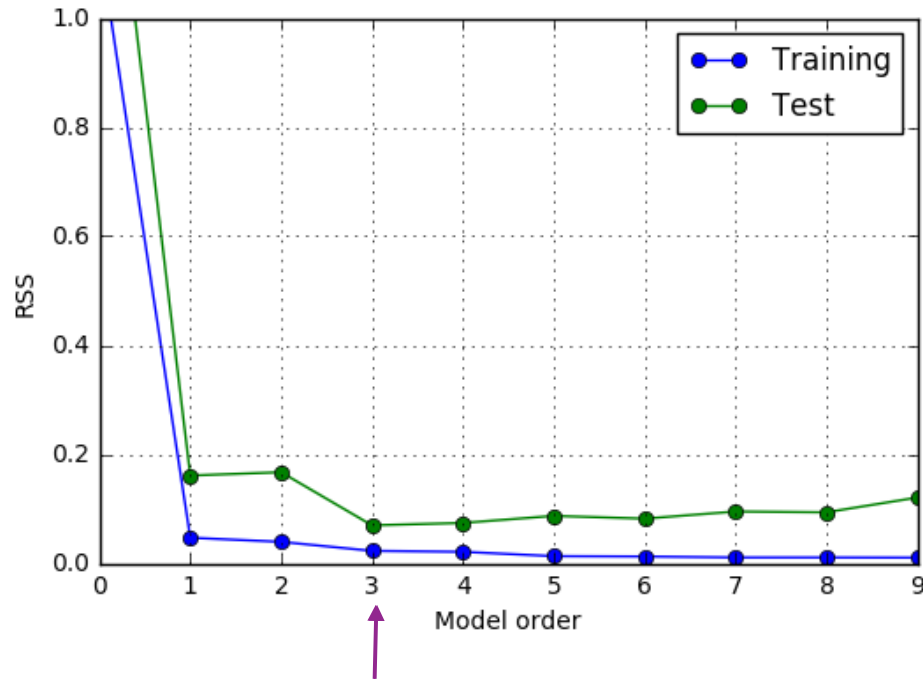
```
# Number of samples for training and test
ntr = nsamp // 2
nts = nsamp - ntr

# Training
xtr = xdat[:ntr]
ytr = ydat[:ntr]

# Test
xts = xdat[ntr:]
yts = ydat[ntr:]
```

# Finding the Model Order

Estimated optimal model order = 3



RSS test minimized at  $d = 3$   
RSS training always decreases

```
dtest = np.array(range(0,10))
RSStest = []
RSStr = []
for d in dtest:

    # Fit data
    beta_hat = poly.polyfit(xtr,ytr,d)

    # Measure RSS on training data
    # This is not necessary, but we do it just to show the training error
    yhat = poly.polyval(xtr,beta_hat)
    RSSd = np.mean((yhat-ytr)**2)
    RSStr.append(RSSd)

    # Measure RSS on test data
    yhat = poly.polyval(xts,beta_hat)
    RSSd = np.mean((yhat-yts)**2)
    RSStest.append(RSSd)

plt.plot(dtest,RSStr,'bo-')
plt.plot(dtest,RSStest,'go-')
plt.xlabel('Model order')
plt.ylabel('RSS')
plt.grid()
plt.ylim(0,1)
plt.legend(['Training','Test'],loc='upper right')
```

# Problems with Simple Train/Test Split

---

- ❑ Test error could vary significantly depending on samples selected
- ❑ Only use limited number of samples for training
- ❑ Problems particularly bad for data with limited number of samples



# K-Fold Cross Validation

## □ K-fold cross validation

- Divide data into  $K$  parts
- Use  $K - 1$  parts for training. Use remaining for test.
- Average over the  $K$  test choices
- More accurate, but requires  $K$  fits of parameters
- Typical choice:  $K=5$  or  $10$
- Average MSE over  $K$  folds estimates the total MSE
- $(= \text{Bias}^2 + \text{Variance} + \text{irreducible error})$



## □ Leave one out cross validation (LOOCV)

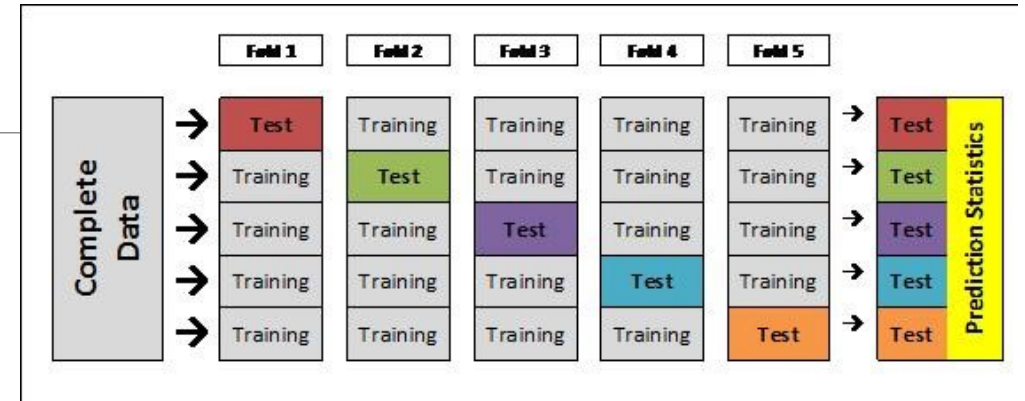
- Take  $K = N$  so one sample is left out.
- Most accurate, but requires  $N$  model fittings
- Necessary when  $N$  is small.

From

<http://blog.goldenhelix.com/goldenadmin/cross-validation-for-genomic-prediction-in-svs/>

# K-Fold Pseudo-Code

- Get data  $X, y$
- For  $i = 1$  to  $K$  // Loop over folds
  - Split into training  $X_{tr}, y_{tr}$  and test  $X_{ts}, y_{ts}$  for fold  $i$
  - For  $p = 1$  to  $p_{max}$  // Loop over model order
    - **Fit** on training data with model order  $p$ :  $\hat{\beta} = \text{fit}(X_{tr}, y_{tr}, p)$
    - **Predict** values on test data:  $\hat{y}_{ts} = \text{predict}(X_{ts}, \hat{\beta})$
    - **Score** the fit on test data:  $S[p, i] = \text{score}(y_{ts}, \hat{y}_{ts})$
- Find **average score** for each model order:  $\bar{S}[p] = \frac{1}{K} \sum_{i=1}^K S[p, i]$
- Select model order with lowest average score:  $\hat{p} = \arg \min_p \bar{S}[p]$



# Polynomial Example

❑ Use sklearn Kfold object

❑ Loop

- Outer loop: Over K folds
- Inner loop: Over D model orders
- Measure test error in each fold and order
- Averaging test errors from K folds for each model order
- Find the model order with the minimal average test errors
- Can be time-consuming

```
# Create a k-fold object
nfold = 20
kf = sklearn.model_selection.KFold(n_splits=nfold,shuffle=True)

# Model orders to be tested
dtest = np.arange(0,10)
nd = len(dtest)

# Loop over the folds
RSSs = np.zeros((nd,nfold))
for isplit, Ind in enumerate(kf.split(xdat)):

    # Get the training data in the split
    Itr, Its = Ind
    xtr = xdat[Itr]
    ytr = ydat[Itr]
    xts = xdat[Its]
    yts = ydat[Its]

    for it, d in enumerate(dtest):

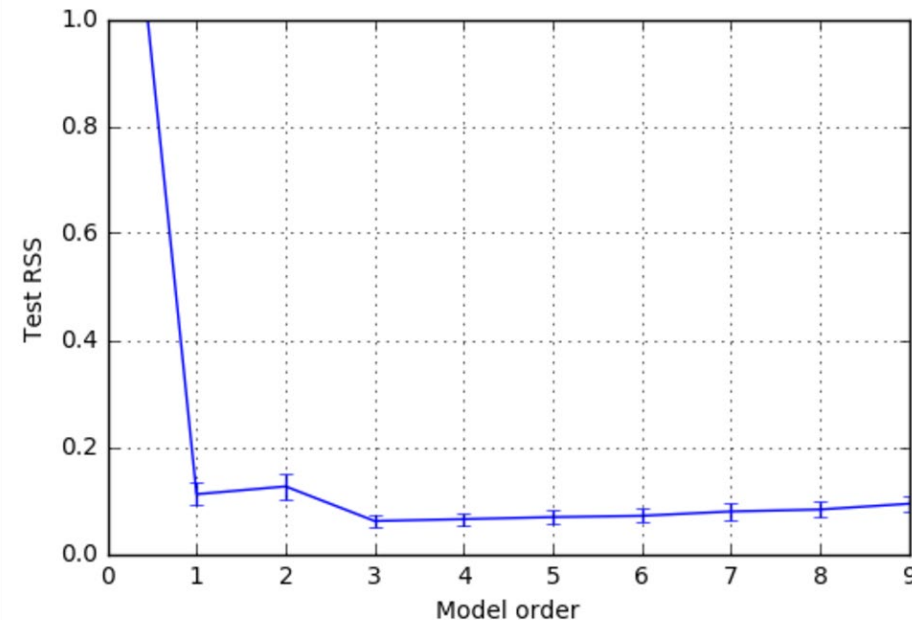
        # Fit data on training data
        beta_hat = poly.polyfit(xtr,ytr,d)

        # Measure RSS on test data
        yhat = poly.polyval(xts,beta_hat)
        RSSs[it,isplit] = np.mean((yhat-yts)**2)
```

# Polynomial Example CV Results

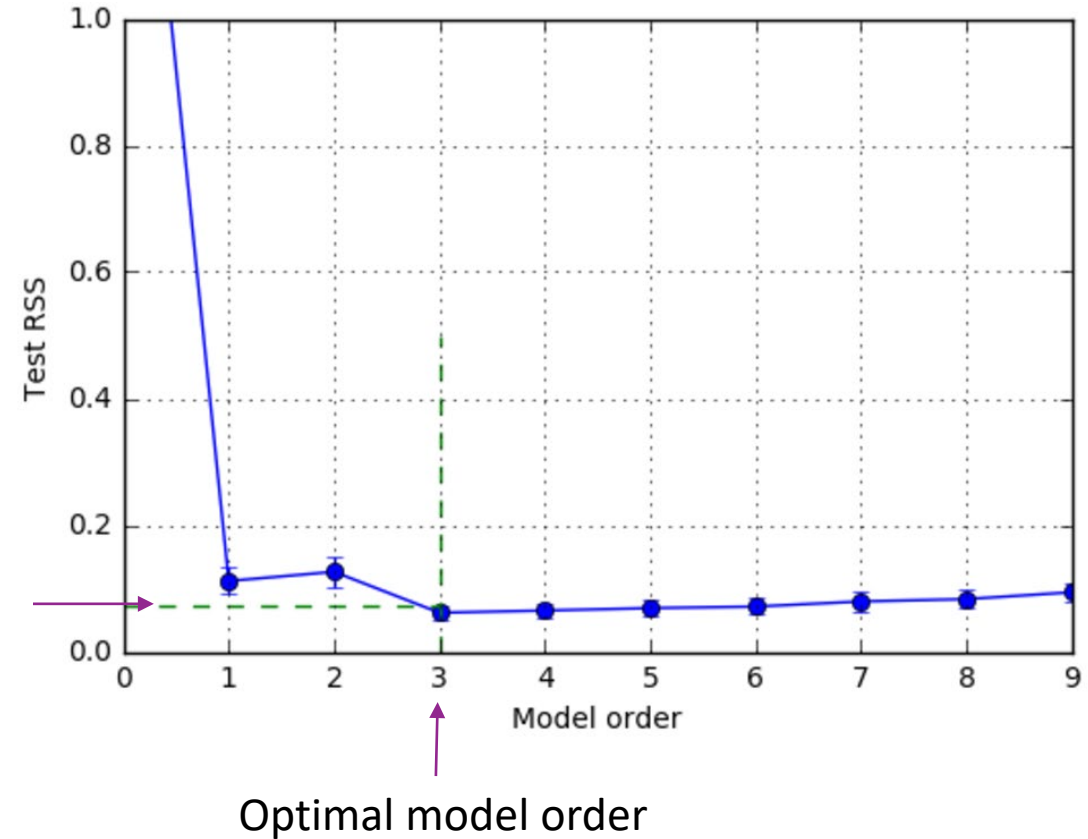
- ❑ For each model order  $d$ 
  - Compute mean test RSS over  $K$  folds
  - Compute standard error (SE) of test RSS
  - $SE = \text{STD of mean RSS} = \text{RSS std} / \sqrt{K - 1}$
  - (expectation over different realizations of data in each fold)
- ❑ Simple model selection
  - Select  $d$  with lowest mean test RSS
- ❑ For this example
  - Estimate model order = 3

```
RSS_mean = np.mean(RSSsts,axis=1)
RSS_std = np.std(RSSsts,axis=1) / np.sqrt(nfold-1)
plt.errorbar(dtest, RSS_mean, yerr=RSS_std, fmt='-.')
plt.ylim(0,1)
plt.xlabel('Model order')
plt.ylabel('Test RSS')
plt.grid()
```



# One Standard Error Rule

- ❑ Previous slide: Select  $d$  to minimize  $RSS\_mean[d]$ 
  - Average RSS across the folds and select lowest average
- ❑ Problem: Often over-predicts model order
- ❑ One standard deviation rule
  - Use simplest model “within one SE of minimum”



# One SE Rule Pseudo-Code

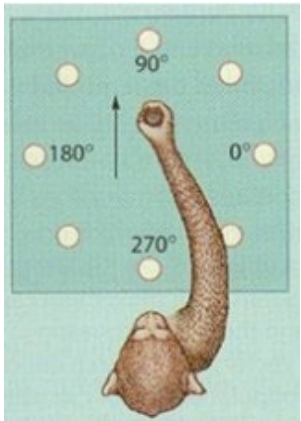
- ❑ Get data  $X, y$
- ❑ Compute score as before:  $S[p, i] = \text{score for model order } p \text{ on fold } i$
- ❑ Compute average, std deviation and standard error of the scores:
  - $\bar{S}[p] = \frac{1}{K} \sum_{i=1}^K S[p, i]$ ,  $\sigma^2[p] = \frac{1}{K} \sum_{i=1}^K (S[p, i] - \bar{S}[p])^2$ ,  $SE[p] = \frac{\sigma[p]}{\sqrt{K-1}}$
- ❑ Find model order via **normal rule**:  $\hat{p}_0 = \arg \min_p \bar{S}[p]$  (lowest average score)
- ❑ Compute target score:  $S_{tgt} = \bar{S}[p_0] + SE[p_0]$
- ❑ **One SE rule**: Find simplest model with score lower than target:

$$\hat{p} = \min\{p \mid \bar{S}[p] \leq S_{tgt}\}$$

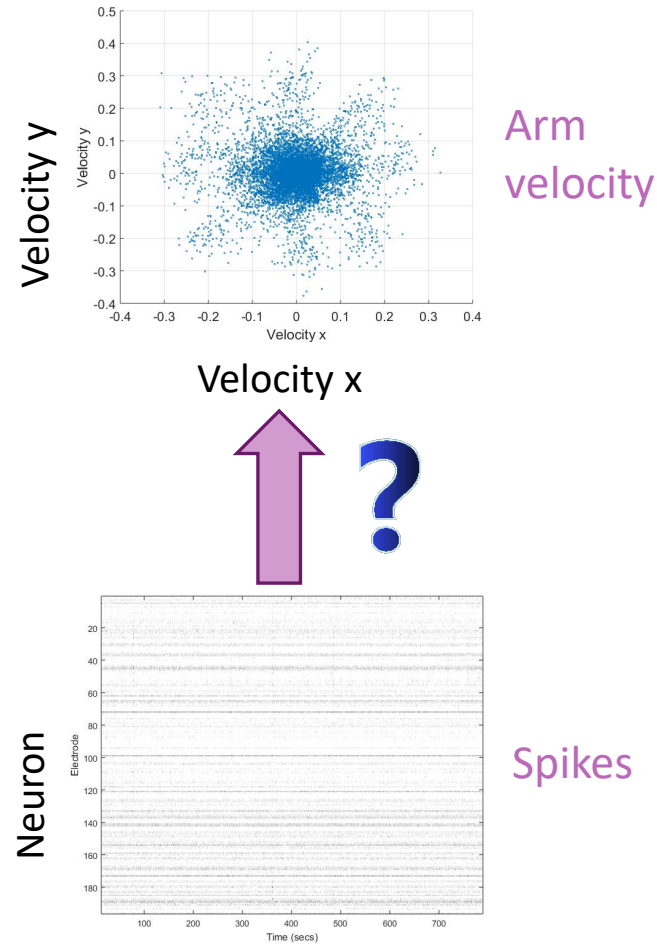
- ❑ Note that one SE rule always produce a model order  $\leq$  normal rule ( $\hat{p} \leq \hat{p}_0$ )

# Lab: Neural ECoG Data

Record arm motion

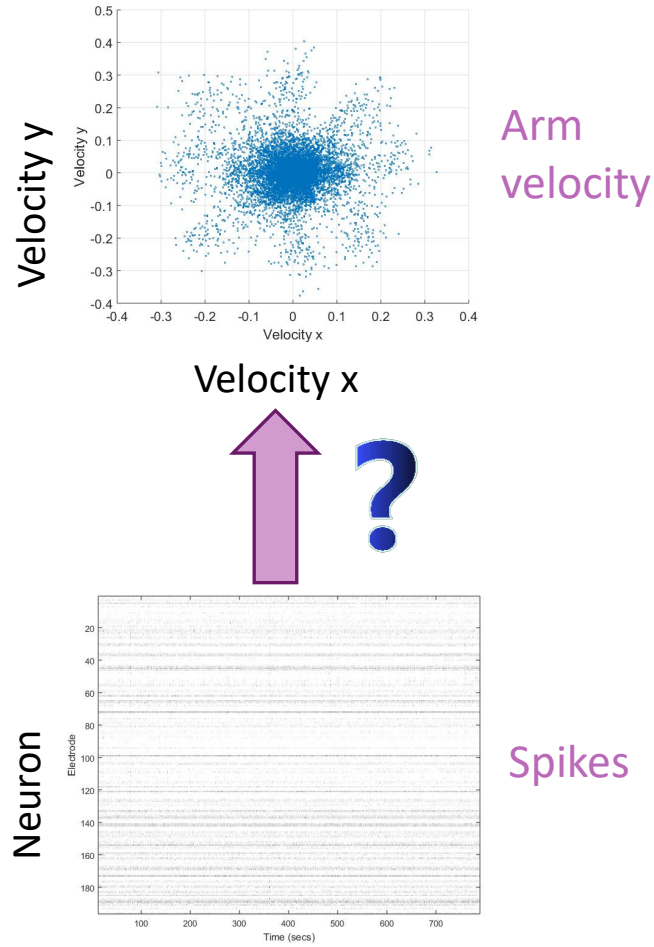


Record spikes  
(ECoG in M1)



- ☐ Read a monkey's brain!
- ☐ Predict hand motion from electrode measurements (number of spikes in each neuron)
- ☐ Data from <https://crcns.org/>
  - Open source website on neural data
  - Great for projects

# Lab: Neural ECoG Data



□ Linear filter model:

$$y[t, k] = \sum_{\ell=0}^d \sum_{j=0}^{p-1} X[t - \ell, j] W[\ell, j, k] + b[k]$$

- $X[t, j]$  = spikes from neuron  $j$  at time  $t$
- $y[t, k]$  = Output  $k$  at time  $t$  (two outputs for x and y motion)
- $W[\ell, j, k]$  = weight from neuron  $j$  to output  $k$  at time delay  $\ell$

□ Linear fitting: Find  $W$  and  $b$  from  $X$  and  $y$

□ Model order selection:

- Find optimal maximum delay  $d$
- Higher  $d$  allows better fit, but uses more parameters

Time