

# DSC5103 Statistics

Session 5. Validation

# Review of last session

- Logistic Regression
  - $Y \sim \text{Bernoulli}(p)$
  - Logistic function as a nonlinear mapping from  $\eta$  to  $p$
- Classification in general
  - From  $p_{\text{hat}}$  to classes
- Other Generalized Linear Models:  $E[Y] \sim X$ 
  - Poisson Regression
  - Survival Analysis

# Plan for today

- Model selection in (generalized) linear models
  - The model selection workflow
  - The traditional vs. modern performance measures
- Validation methods: a tool for numerically estimating out-of-sample error
  - Validation set
  - Leave-One-Out Cross-Validation
  - K-fold Cross-Validation

# Linear Model Selection

- To choose the optimal subset of predictors to be included in the model
- Workflow
  - Best subset: the best out of all possible combinations ( $2^p$ )
  - Forward selection: start from none, iteratively add the variable that improve the performance measure the most
  - Backward selection: start from all, iteratively remove the least significant variable
- Performance measures
  - In-sample measures, such as RSS and  $R^2$ , are not enough (over-fitting!)
  - ~~traditional~~ Adj.  $R^2$ , AIC, BIC, and Mallows'  $C_p$  take model complexity ( $p$ ) into account
  - ~~modern~~ Validation-based methods for approximating out-of-sample errors

Group # of variables	$X_1$	$X_2$	$X_3$	...	$X_p$	FWD Selection	* BWD Selection	Best Subset	Best Model in Group
0						P		1	$M_0$
1		✓				P-1		$\binom{P}{1} = P$	$M_1$
2		✓	✓			P-2		$\binom{P}{2} = \frac{P \cdot (P-1)}{2}$	$M_2$
						⋮		$\binom{P}{3}$	⋮
		✓	✓		✓	⋮	1	⋮	⋮
			⋮			⋮		⋮	⋮
P-1	✓	✓	✓	✓	✓	1	1	$\binom{P}{P-1} = P$	$M_{P-1}$
P	✓	✓	✓	✓	✓		1	$\binom{P}{P} = 1$	$M_P$

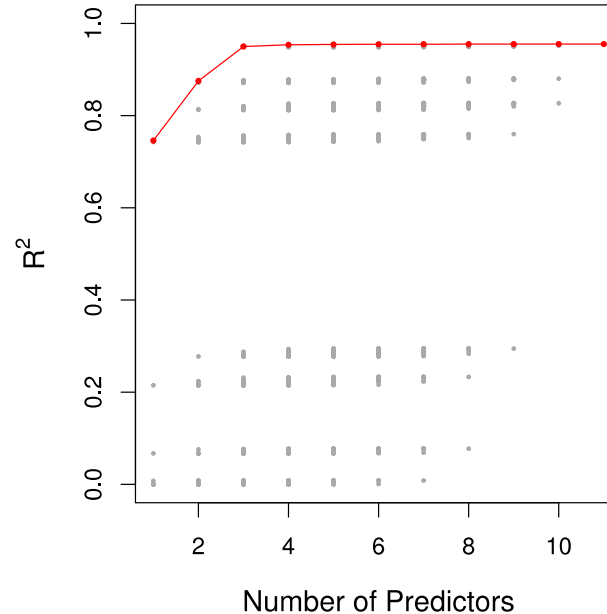
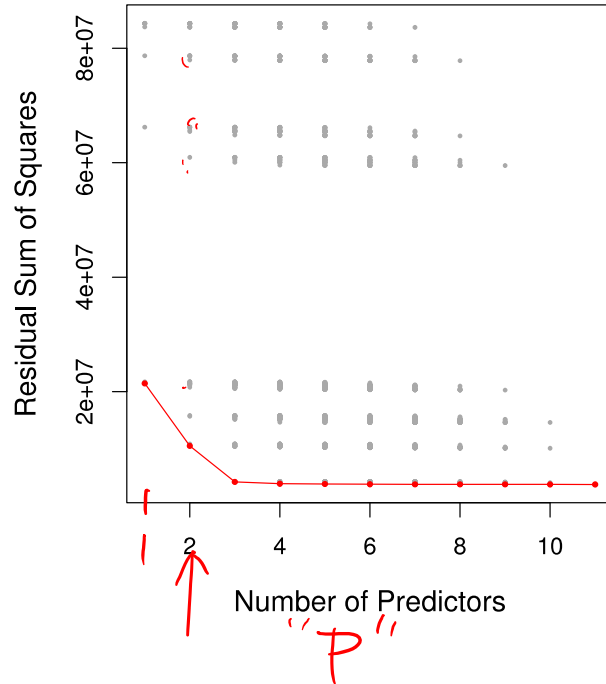
Within Group  $\Rightarrow$  same "P"  $\Rightarrow R^2$  / RSS / deviance (In-Sample)

Across Group  $\Rightarrow$  Adj.  $R^2$  / AIC /  $C_p$  / BIC

$\hookrightarrow$  Validation Error

# Credit Data: $R^2$ vs. Subset Size

- The RSS/ $R^2$  will always decline/increase as the number of variables increase so they are not very useful
- The red line tracks the best model for a given number of predictors, according to RSS and  $R^2$



# Other Measures for Model Comparison

- *Indirectly* estimate test error by making an adjustment to the training error to account for the bias due to over-fitting

- Adjusted  $R^2$

$$\text{adjusted } R^2 = 1 - \frac{\text{RSS}/(n - \textcircled{p} - 1)}{\text{TSS}/(n - 1)}$$

- AIC (Akaike information criterion)

(GLM)  $\text{AIC} = -2\text{Log-Likelihood} + \textcircled{2p}$   $\text{AIC} = (\text{RSS} + 2p\hat{\sigma}^2)/(n\hat{\sigma}^2)$  (Special: Linear / Gaussian)

- Mallows's  $C_p$  (equivalent to AIC for linear regression)

$$C_p = (\text{RSS} + \textcircled{2p}\hat{\sigma}^2)/n$$

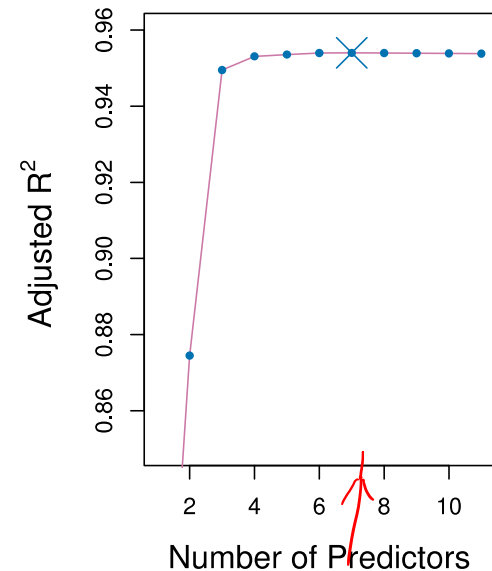
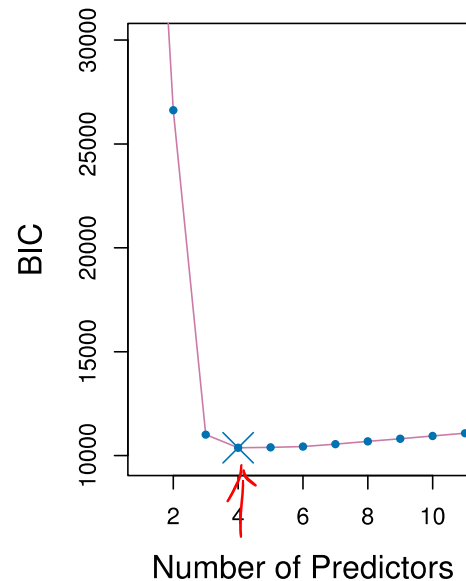
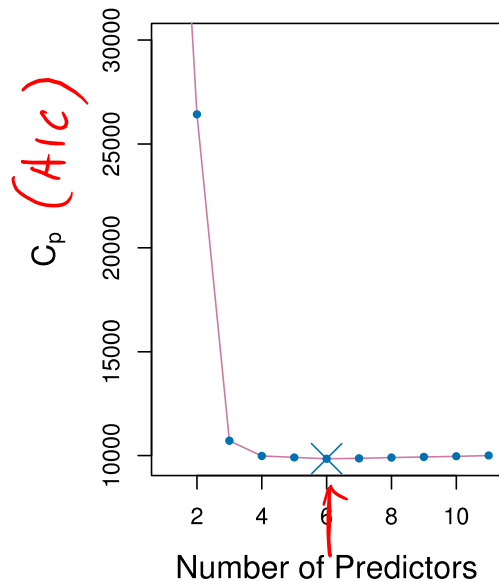
- BIC (Bayesian information criterion)

$$\text{BIC} = (\text{RSS} + \textcircled{\log(n)p}\hat{\sigma}^2)/n$$

- These methods add penalty to RSS for the number of variables (i.e. complexity) in the model, but none are perfect (e.g., how to estimate  $\sigma^2$ ?)

# Credit Data: $C_p$ , BIC, and Adjusted $R^2$

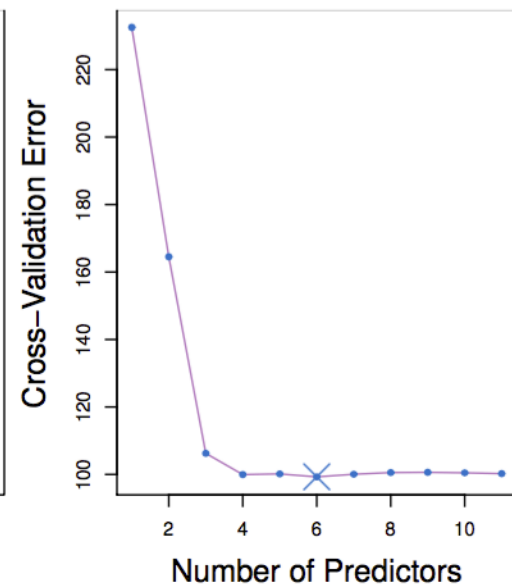
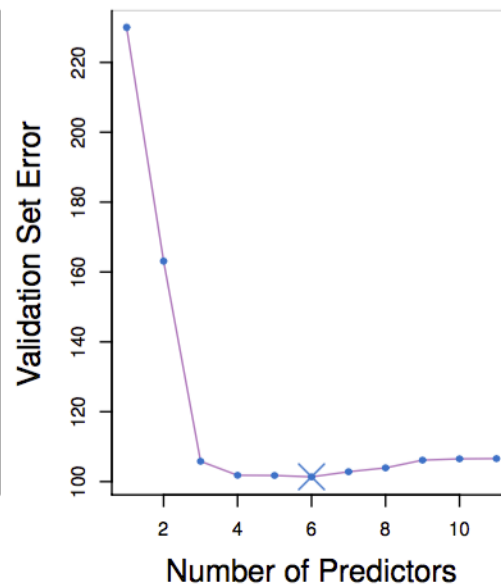
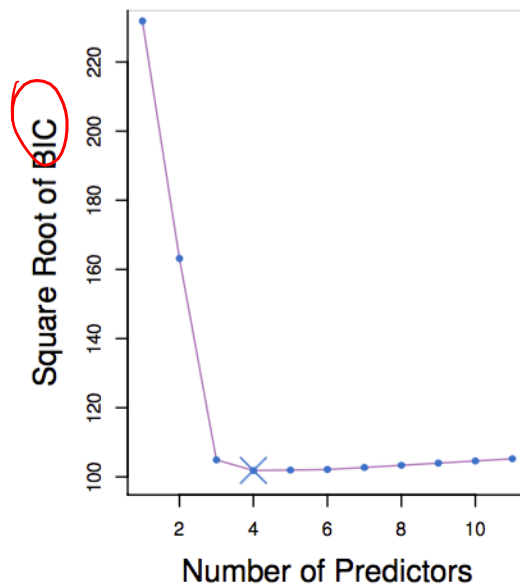
- A small value of  $C_p$  and BIC indicates a low error, and thus a better model
- A large value for the Adjusted  $R^2$  indicates a better model





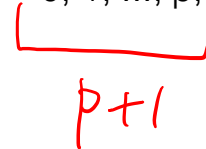
# Model Comparison by Cross-Validation

- *Directly* estimate the out-of-sample error using validation/cross-validation



# Model Selection

- Model selection with out-of-sample error in mind
  - CV is computationally intensive, it is not practical to do it for all possible models
  - A hybrid approach:
    - For each fixed model size  $k = 0, 1, \dots, p$ , select the best  $k$  predictors by RSS or  $R^2$ . We obtain the best model if we choose to have  $k$  predictors. Let's call it  $\mathbf{M}_k$ .
    - Use Cp/AIC/BIC or cross-validation to compare  $\mathbf{M}_k$  for  $k = 0, 1, \dots, p$ , and choose the best  $k$ .

  
 $p+1$

# Model Selection Algorithm

- Best Subset Selection

## *Best Subset Selection*

1. Let  $\mathcal{M}_0$  denote the *null model*, which contains no predictors. This model simply predicts the sample mean for each observation.
2. For  $k = 1, 2, \dots, p$ :
  - (a) Fit all  $\binom{p}{k}$  models that contain exactly  $k$  predictors.
  - (b) Pick the best among these  $\binom{p}{k}$  models, and call it  $\mathcal{M}_k$ . Here *best* is defined as having the smallest RSS, or equivalently largest  $R^2$ .
3. Select a single best model from among  $\mathcal{M}_0, \dots, \mathcal{M}_p$  using cross-validated prediction error,  $C_p$  (AIC), BIC, or adjusted  $R^2$ .

# Model Selection Algorithm

- Forward Stepwise Selection

## *Forward Stepwise Selection*

1. Let  $\mathcal{M}_0$  denote the *null* model, which contains no predictors.
2. For  $k = 0, \dots, p - 1$ :
  - 2.1 Consider all  $p - k$  models that augment the predictors in  $\mathcal{M}_k$  with one additional predictor.
  - 2.2 Choose the *best* among these  $p - k$  models, and call it  $\mathcal{M}_{k+1}$ . Here *best* is defined as having smallest RSS or highest  $R^2$ .
3. Select a single best model from among  $\mathcal{M}_0, \dots, \mathcal{M}_p$  using cross-validated prediction error,  $C_p$  (AIC), BIC, or adjusted  $R^2$ .

# Model Selection Algorithm

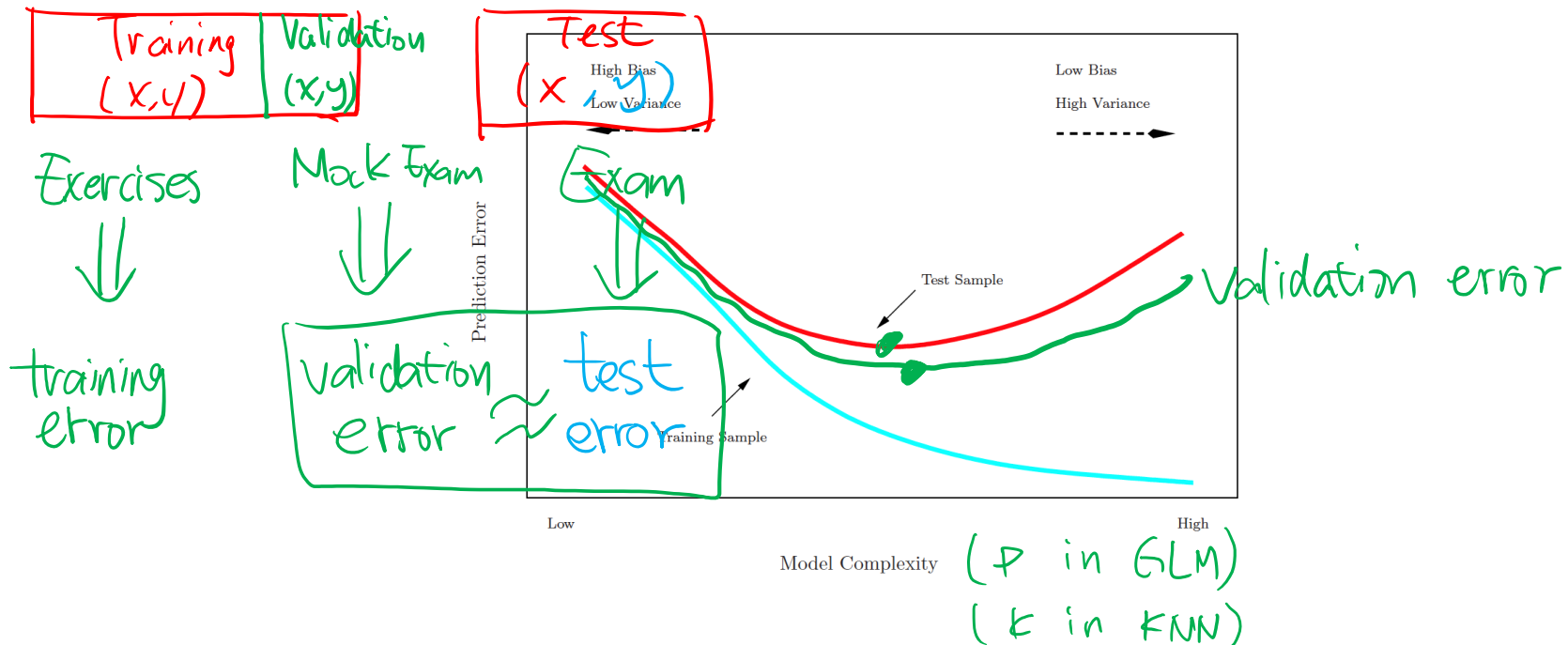
- Backward Stepwise Selection

## *Backward Stepwise Selection*

1. Let  $\mathcal{M}_p$  denote the *full* model, which contains all  $p$  predictors.
2. For  $k = p, p - 1, \dots, 1$ :
  - 2.1 Consider all  $k$  models that contain all but one of the predictors in  $\mathcal{M}_k$ , for a total of  $k - 1$  predictors.
  - 2.2 Choose the *best* among these  $k$  models, and call it  $\mathcal{M}_{k-1}$ . Here *best* is defined as having smallest RSS or highest  $R^2$ .
3. Select a single best model from among  $\mathcal{M}_0, \dots, \mathcal{M}_p$  using cross-validated prediction error,  $C_p$  (AIC), BIC, or adjusted  $R^2$ .

# Evaluating Out-of-Sample Error

- Motivation
  - In-sample vs. Out-of-sample Error (Bias-Variance Trade-off)
  - How to estimate out-of-sample error (and then do model selection) **without** a test dataset?!
    - Theoretical adjustment (Adjusted  $R^2$ , AIC, BIC,  $C_p$ ): penalize model complexity (P)
    - “Estimate” the out-of-sample error using validation data



# Evaluating Out-of-Sample Error

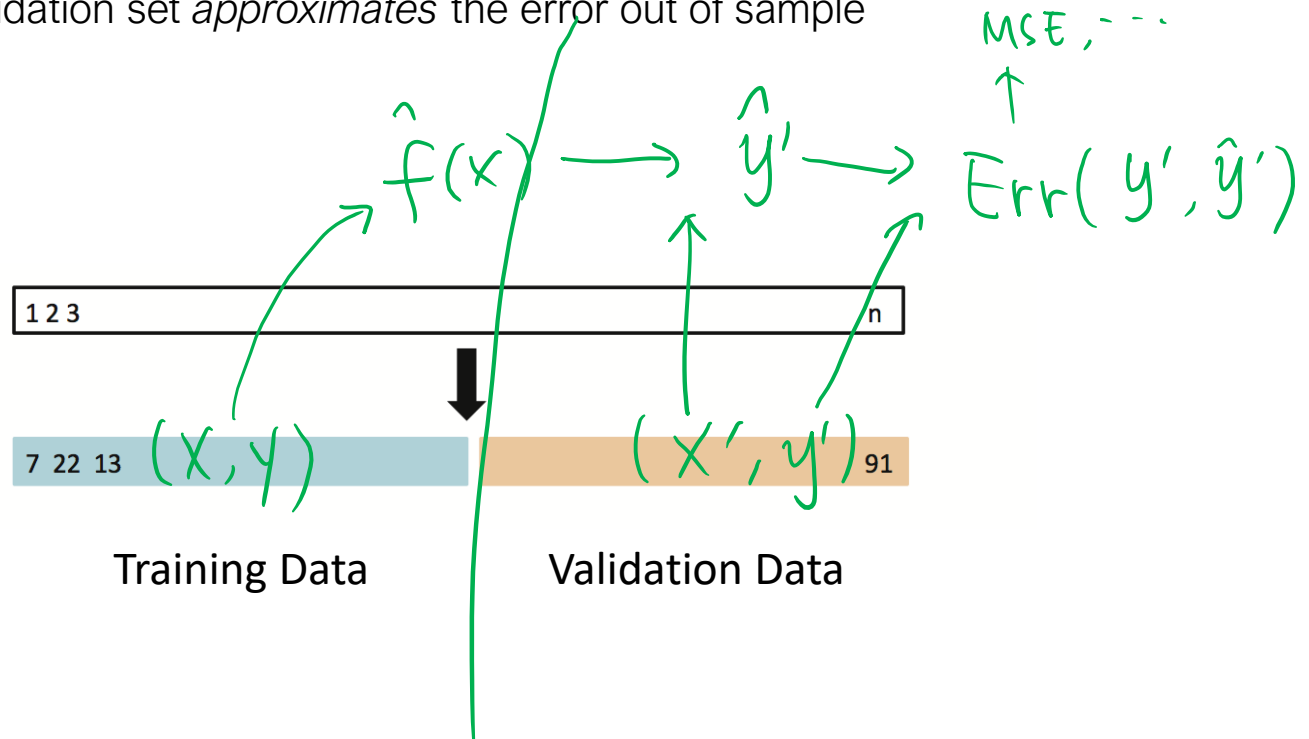
- Three common approaches
  - The Validation Set Approach
  - Leave-One-Out Cross Validation
  - K-fold Cross Validation

# The Validation Set Approach

Stratified sampling

- The validation-set approach

- Randomly divide the available data into **Training** and **Validation** set
- Fit the model using the Training set, evaluate the prediction on the Validation set
- Error in Validation set *approximates* the error out of sample

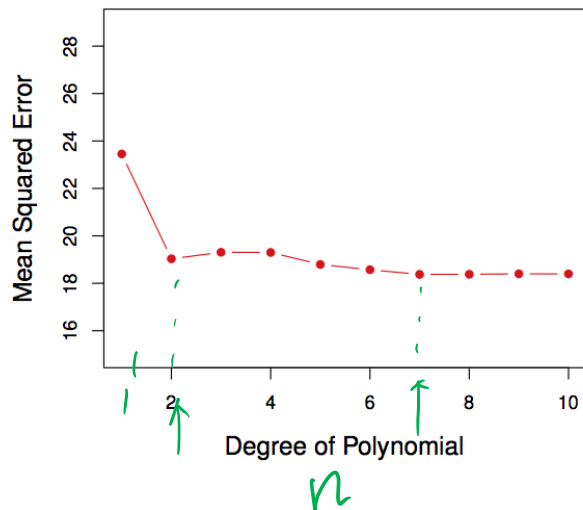




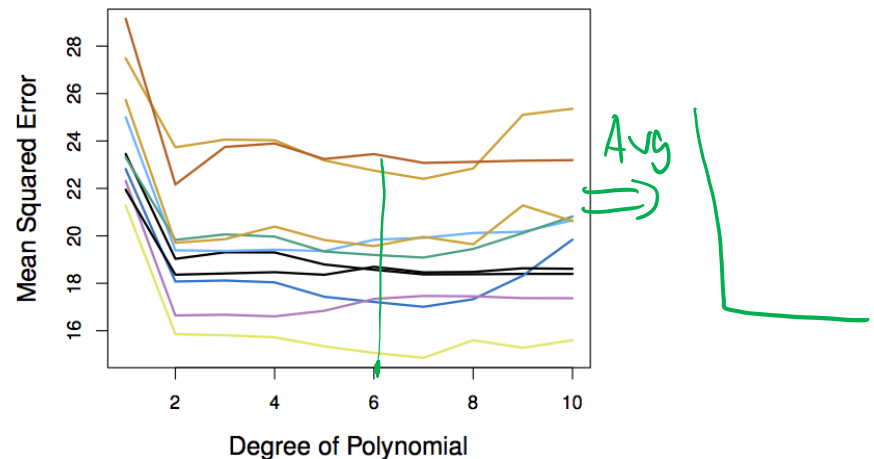
# Example: Auto Data

- Suppose that we want to predict  $y$   $\text{mpg}$  from  $x$   $\text{horsepower}$
- Compare models:
  - $\text{mpg} \sim \text{horsepower}^n, n=1, 2, \dots, 10$   $\leftarrow 10 \text{ model}, n^*$
- Which model gives a better fit?
  - Randomly split  $\text{Auto}$  data set into training (196 obs.) and validation data (196 obs.)

Validation error for a single split

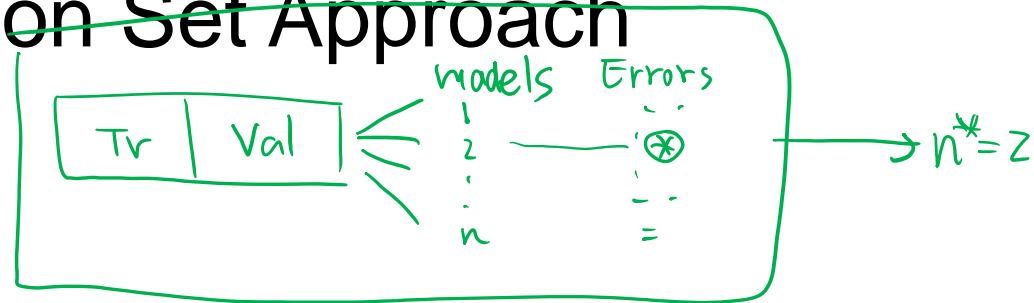


Validation error for 10 random splits



# The Validation Set Approach

Model Selection



- Advantages:

- Simple and easy to implement
- Computationally efficient: one run of fitting on part of the data

- Disadvantages:

- Less data: only a subset of observations are used to fit the model (training data)
  - an overestimation of the out-of-sample error

$$\text{Tr } (n=z) \rightarrow \hat{f}()$$

Random  
splitting

Higher variance: the validation MSE can be highly variable because of the randomness in constructing Training and Validation datasets

→ average over multiple runs

# Leave-One-Out Cross Validation (LOOCV)

- For a dataset of size  $n$ , repeat the following  $n$  times
  - In iteration  $i$ , use the  $i$ -th data point for validation, the rest for training ( $n-1$ )
  - Fit the model with training data, and obtain validation error on point  $i$
- The LOOCV error for the model is the average of the  $n$  validation errors:

$$CV_{(n)} = \frac{1}{n} \sum_{i=1}^n M SE_i.$$

$Err(y, \hat{y})$

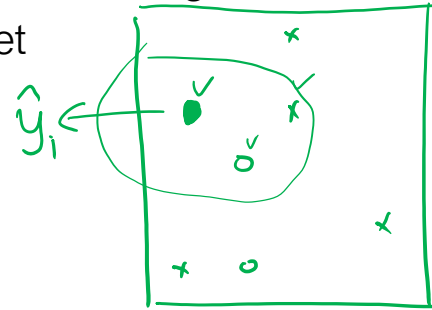
The diagram illustrates the LOOCV process. At the top, a horizontal bar represents the dataset of size  $n$ , labeled '123' and 'n'. A large black arrow points down to a vertical list of  $n$  rows, each representing an iteration. Each row shows the validation point (123) and the predicted value ( $\hat{y}$ ) for the training set (123). The first two rows show the validation point (123) and the predicted value ( $\hat{y}$ ) for the training set (123). The remaining rows show the validation point (123) and the predicted value ( $\hat{y}$ ) for the training set (123). A green bracket on the right indicates the total number of iterations is  $n$ .

# LOOCV vs. the Validation Set Approach

In-sample

LOOCV

- Designed to overcome the previous disadvantages  $k=3$ 
  - No randomness in sampling the dataset
  - Maximal utilization of data for training
- Disadvantages
  - LOOCV is computationally intensive (We fit the each model  $n$  times!)
    - Exception: least-squares linear or polynomial regression, KNN



Randomness  
in

the data sample

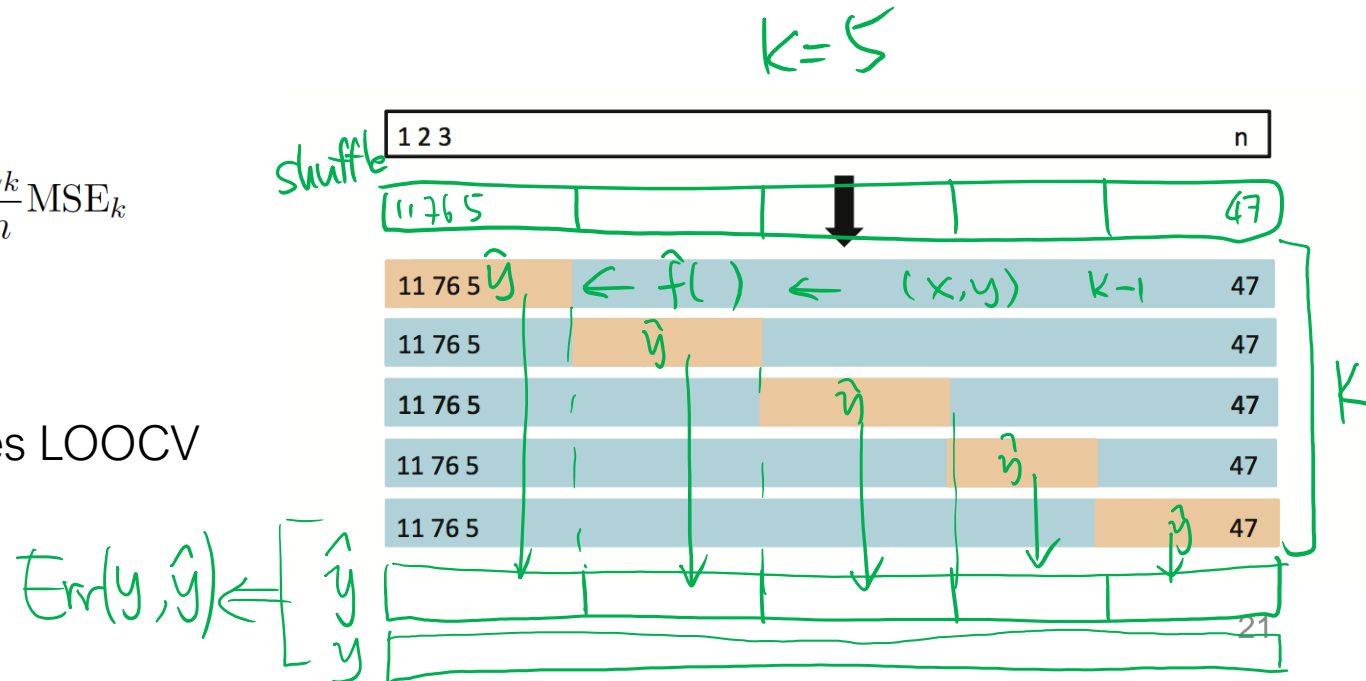
- High variance: each fold (iteration) is using almost the same data => high correlation

# K-fold Cross Validation

- A trade-off between the validation-set approach and LOOCV
- Randomly divide the data into  $K$  different parts, repeat the following  $K$  times
  - Use the  $i$ -th part for validation, the remaining  $K-1$  parts for training
  - Fit the model with training data, and obtain validation error
- The  $K$ -fold cross validation error (parts may be of different size  $n_k$ )

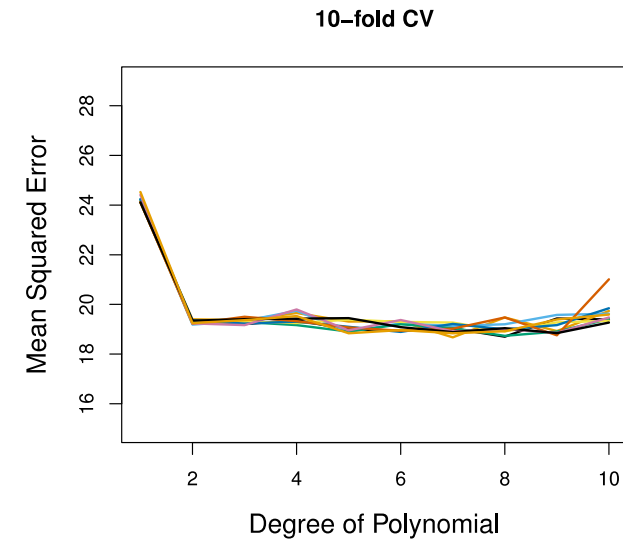
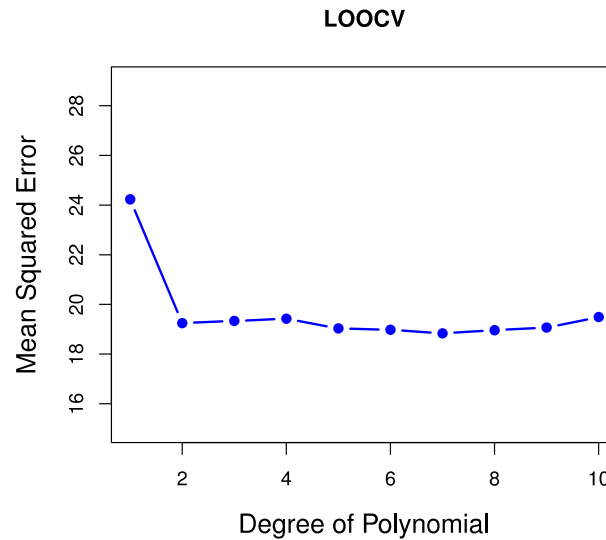
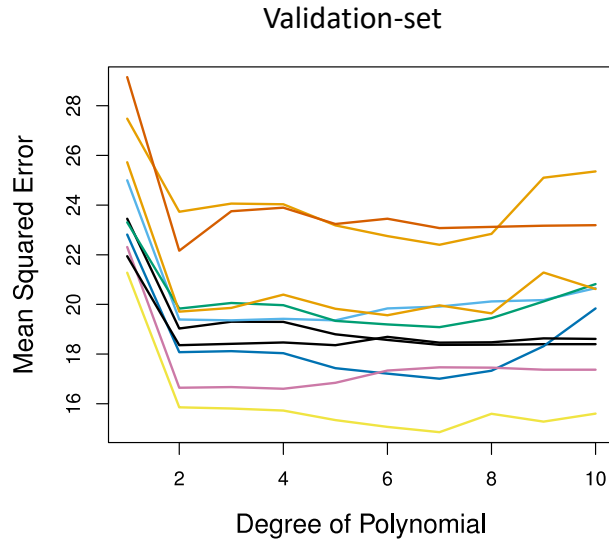
$$CV_{(K)} = \sum_{k=1}^K \frac{n_k}{n} MSE_k$$

- If  $K=n$ , it becomes LOOCV

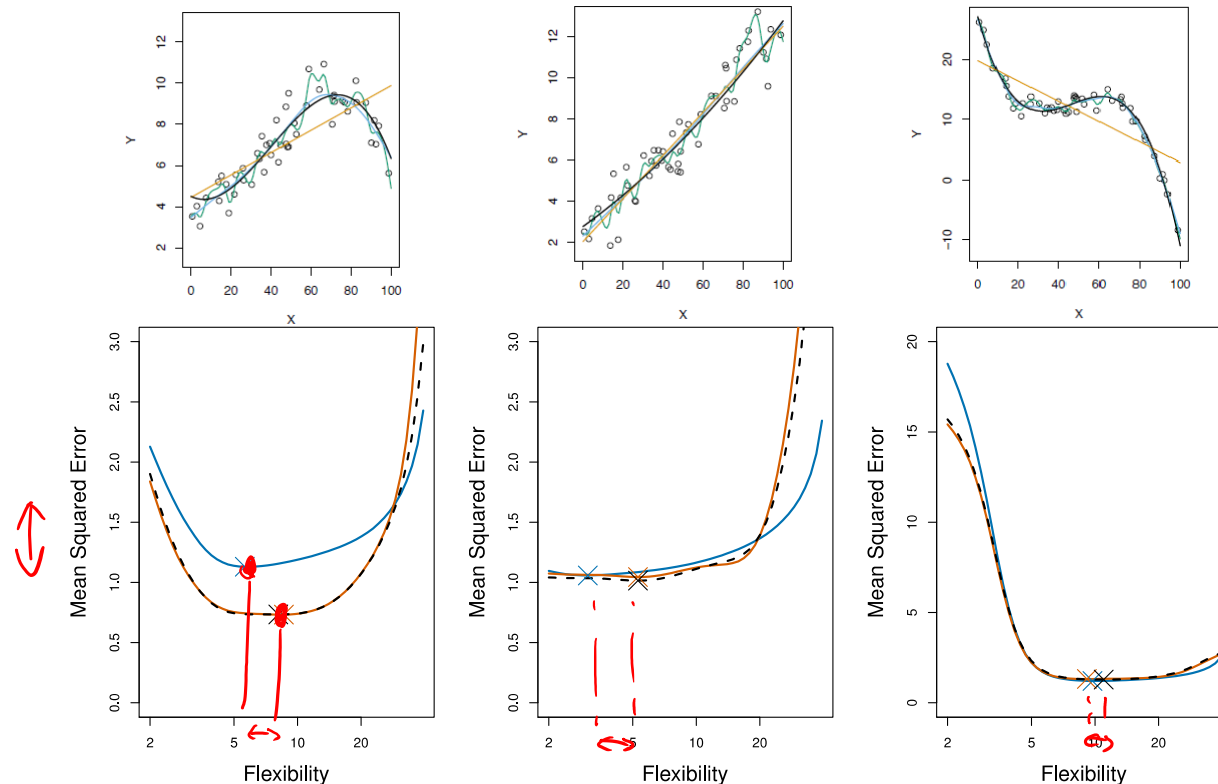


# Auto Data: LOOCV vs. K-fold CV

- Left: Validation-set, repeated many times
- Middle: LOOCV
- Right: 10-fold CV, repeated many times
  - K-fold CV is still random, but variability is small



# K-fold Cross Validation on Simulated Data



- Blue: Test MSE
- Black: LOOCV MSE
- Orange: 10-fold CV MSE
- Refer to chapter 2 for the top graphs, Fig 2.9, 2.10, and 2.11

# Bias-Variance Trade-off for K-fold CV

- How to choose K?



- Recommendation
  - We tend to use K-fold CV with  $K = 5$  or  $K = 10$
  - It has been empirically shown that they yield test error rate estimates that suffer neither from excessively high bias, nor from very high variance



# Cross Validation on Classification Problems

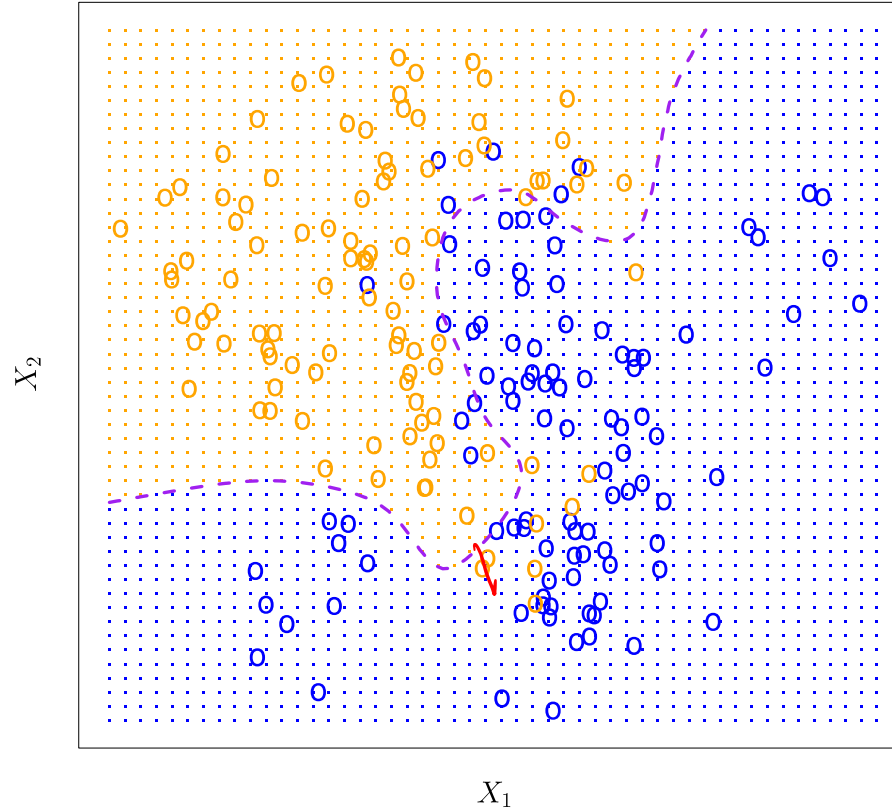
- Cross validation on classification problems in a similar manner
  - Replace MSE with error rate or other performance measures for classification
  - ROC / AUC on the validation dataset

$$\left. \begin{array}{c} \boxed{1 \quad \hat{y} \quad n} \\ \boxed{1 \quad y \quad n} \end{array} \right\} \Rightarrow \frac{\text{Err}(y, \hat{y})}{\hookrightarrow \text{mse}, \dots}$$

$$\left. \begin{array}{c} \boxed{1 \quad \hat{p} \quad n} \\ \boxed{1 \quad y \quad n} \end{array} \right\} \Rightarrow \begin{array}{l} \text{Err}(y, \hat{p}) \\ \hookrightarrow \text{misclassification} \\ \hookrightarrow \text{AUC} \\ \hookrightarrow \dots \end{array}$$

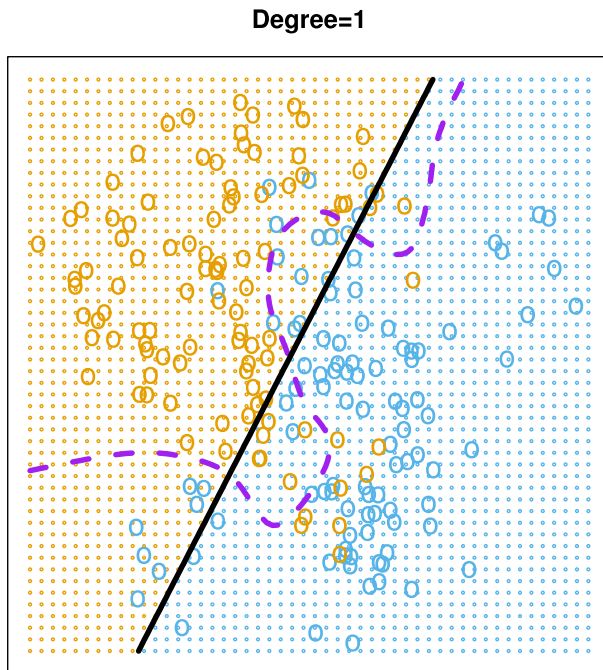
# CV to Choose Order of Polynomial

- The data set used is simulated (refer to Fig 2.13)
- The purple dashed line is the Bayes' boundary

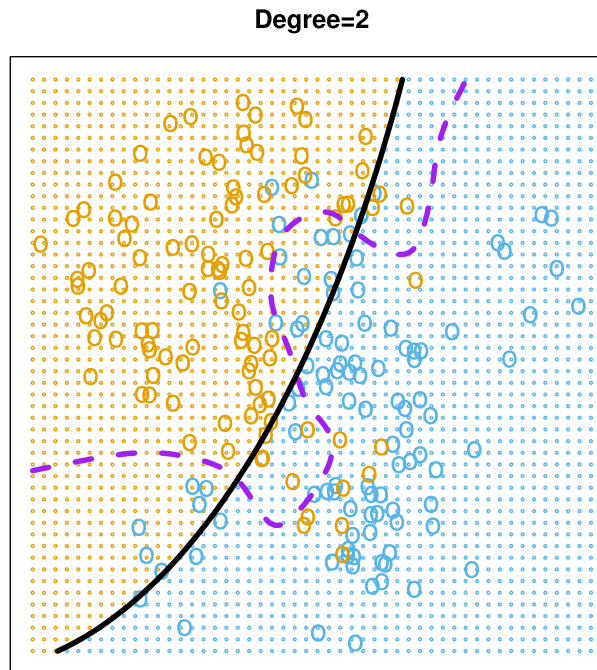


# CV to Choose Order of Polynomial

- Linear Logistic regression (Degree 1) is not able to fit the Bayes' decision boundary
- Quadratic Logistic regression does better than linear



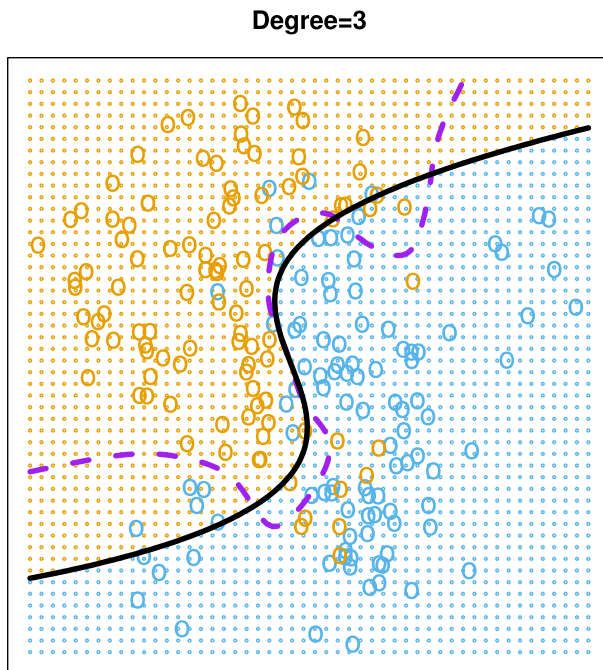
Error Rate: 0.201



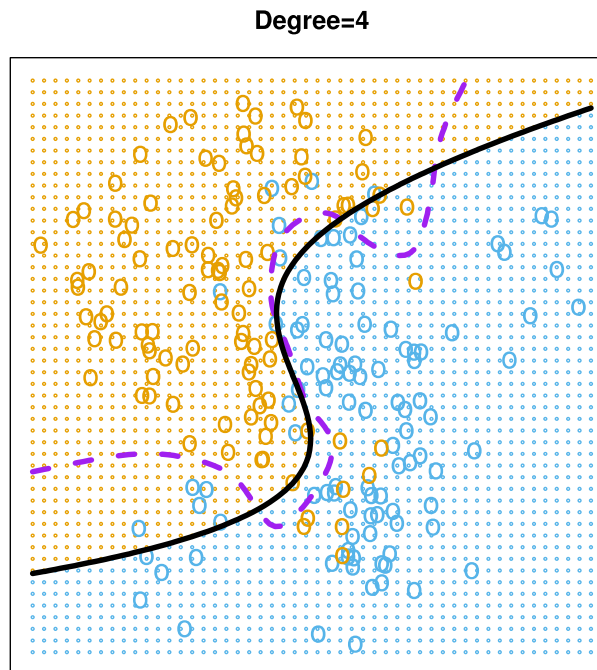
Error Rate: 0.197

# CV to Choose Order of Polynomial

- Using cubic and quartic predictors, the accuracy of the model improves



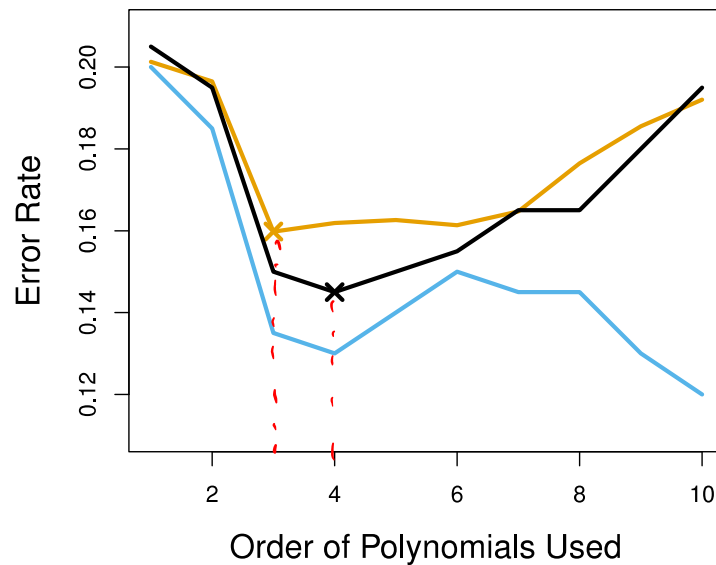
Error Rate: 0.160



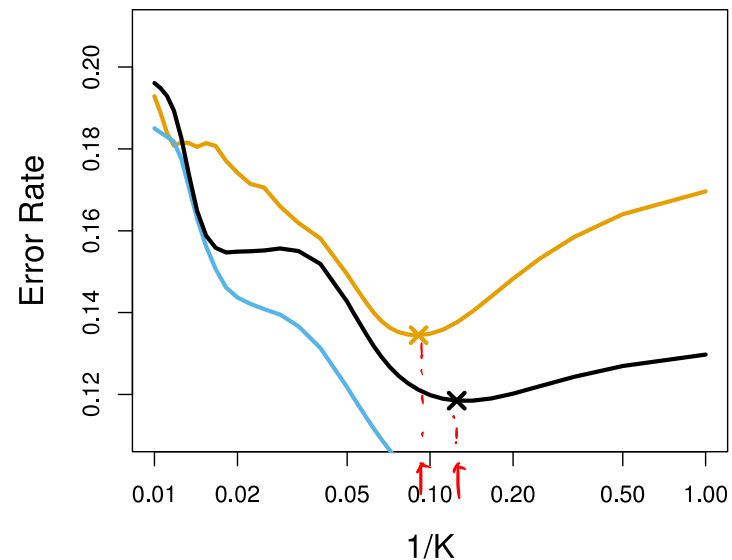
Error Rate: 0.162

# CV to Choose the Order

Logistic Regression



KNN



- Brown: Test Error
- Blue: Training Error
- Black: 10-fold CV Error

Even when the CV error and Test error are quite different, CV error serves as a good measure for model selection!

# Cross-Validation in R

- Demo of the three approaches on GLM (using `cv.glm()`)
    - `5-glm_validation.R`
  - Demo of LOOCV in kNN (using `knn.cv()`)
    - `5-Mixture_knn_cv.R`
- 
- Demo of k-fold CV in logistic regression (using `cv.glm()` and `cost`)
  - Demo of manual k-fold CV in logistic regression
    - `5-Mixture_LogisticRegression.R`

cv.glm()  
• set.seed()  
run = 1, ..., RUN

model  $i = 1, \dots, I$

cv.glm()

• Split data into  $K$  fold

$k = 1, \dots, K$

• fit model  
with  $k-1$

• predict on 1 fold

• Error measure

manual

• set.seed()

run = 1, ..., RUN

\* Split data into  $K$  folds

model family I  
 $i = 1, \dots, I$

$k = 1, \dots, K$

• fit  
• Predict  
• error

model family II  
 $j = 1, \dots, J$

$k = 1, \dots, K$

• fit  
• predict  
• error