

# Lecture 17: Model Selection Algorithms

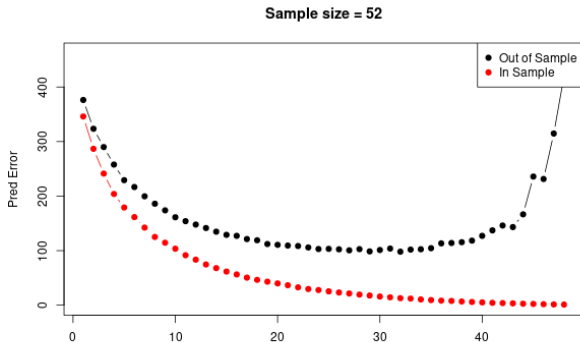
Module 5: part 2

Spring 2025

# Model Selection Recap

# Recap

- When selecting a model which will perform well on new data, we can't just consider how well it performs on the data it is fit to
- Choosing the level of complexity involves a bias vs variance trade-off
- Including more “truly relevant” covariates can improve prediction (decrease bias)
- Including more covariates means  $\hat{b}$  are generally not estimated as well (increase variance)



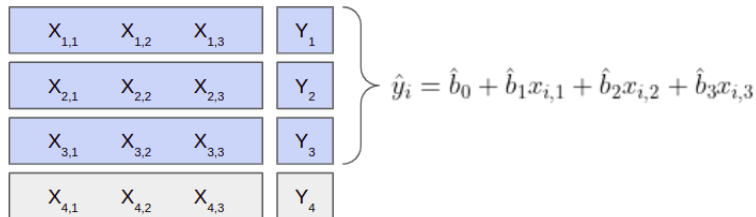
# Leave one out cross validation

Cross validation gives us a way to estimate how a model will perform on “new data”

|           |           |           |       |
|-----------|-----------|-----------|-------|
| $X_{1,1}$ | $X_{1,2}$ | $X_{1,3}$ | $Y_1$ |
| $X_{2,1}$ | $X_{2,2}$ | $X_{2,3}$ | $Y_2$ |
| $X_{3,1}$ | $X_{3,2}$ | $X_{3,3}$ | $Y_3$ |
| $X_{4,1}$ | $X_{4,2}$ | $X_{4,3}$ | $Y_4$ |

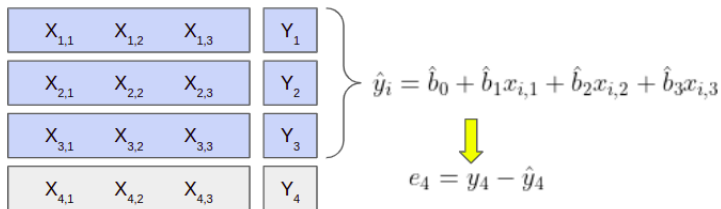
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## Leave one out

- This is called **leave one out cross-validation**
- The cross validation score is:

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

- Not quite RSS since the model we use for  $\hat{y}_i$  is different each time
- The model that produces  $\hat{y}_i$  was fit without using the  $i$ th observation
- This is an unbiased estimator of generalization error; i.e., how well your model will generalize to new data
- Depending on how computationally intensive your model is, and how many data points you have, this may be computationally infeasible

# K-fold cross validation

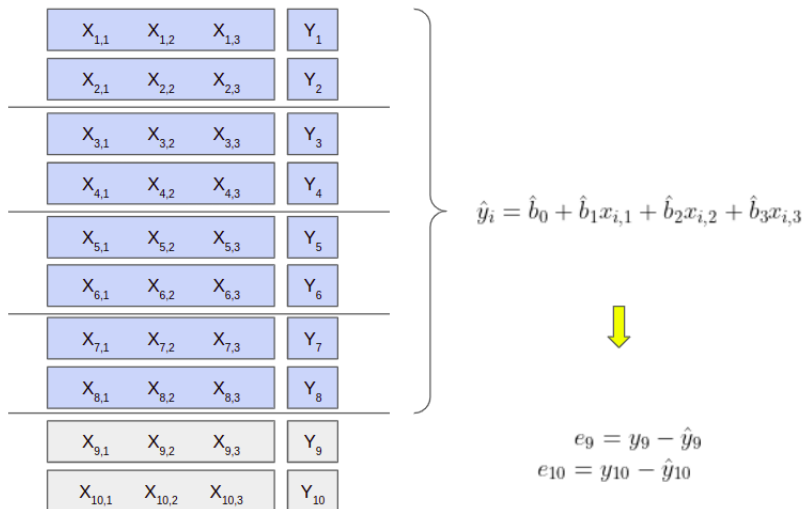
|            |            |            |          |
|------------|------------|------------|----------|
| $X_{1,1}$  | $X_{1,2}$  | $X_{1,3}$  | $Y_1$    |
| $X_{2,1}$  | $X_{2,2}$  | $X_{2,3}$  | $Y_2$    |
| <hr/>      |            |            |          |
| $X_{3,1}$  | $X_{3,2}$  | $X_{3,3}$  | $Y_3$    |
| $X_{4,1}$  | $X_{4,2}$  | $X_{4,3}$  | $Y_4$    |
| <hr/>      |            |            |          |
| $X_{5,1}$  | $X_{5,2}$  | $X_{5,3}$  | $Y_5$    |
| $X_{6,1}$  | $X_{6,2}$  | $X_{6,3}$  | $Y_6$    |
| <hr/>      |            |            |          |
| $X_{7,1}$  | $X_{7,2}$  | $X_{7,3}$  | $Y_7$    |
| $X_{8,1}$  | $X_{8,2}$  | $X_{8,3}$  | $Y_8$    |
| <hr/>      |            |            |          |
| $X_{9,1}$  | $X_{9,2}$  | $X_{9,3}$  | $Y_9$    |
| $X_{10,1}$ | $X_{10,2}$ | $X_{10,3}$ | $Y_{10}$ |

# K-fold cross validation

|            |            |            |          |
|------------|------------|------------|----------|
| $X_{1,1}$  | $X_{1,2}$  | $X_{1,3}$  | $Y_1$    |
| $X_{2,1}$  | $X_{2,2}$  | $X_{2,3}$  | $Y_2$    |
| <hr/>      |            |            |          |
| $X_{3,1}$  | $X_{3,2}$  | $X_{3,3}$  | $Y_3$    |
| $X_{4,1}$  | $X_{4,2}$  | $X_{4,3}$  | $Y_4$    |
| <hr/>      |            |            |          |
| $X_{5,1}$  | $X_{5,2}$  | $X_{5,3}$  | $Y_5$    |
| $X_{6,1}$  | $X_{6,2}$  | $X_{6,3}$  | $Y_6$    |
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| $X_{10,1}$ | $X_{10,2}$ | $X_{10,3}$ | $Y_{10}$ |

$$\hat{y}_i = \hat{b}_0 + \hat{b}_1 x_{i,1} + \hat{b}_2 x_{i,2} + \hat{b}_3 x_{i,3}$$

# K-fold cross validation



# Recap

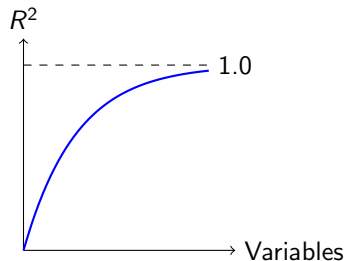
- Several ways to select a training set (data used to estimate parameters of model) and test set (data used to evaluate performance)
  - Leave One Out
  - K-fold CV
- Because the “best” model selection will also depend on the number of observations you have, Leave One Out will generally be best, but also has the highest computational cost
- Use approximations instead which only require fitting the model once
- Models with the lower cross validation scores are better

# The Model Selection Problem

- **Core challenge:** Finding the optimal balance between model complexity and predictive performance
- **Overfitting:** Complex models may fit training data perfectly but perform poorly on new data
- **Underfitting:** Simple models may miss important patterns in the data

# Penalized Scores for Model Selection

# $R^2$ and Its Limitations



$$R^2 = 1 - \frac{\text{RSS}}{\sum_i (y_i - \bar{y})^2}$$

- Measures how well model fits *training data*
- **Key issue:**  $R^2$  *always increases* with additional predictors
- This incentivizes unnecessarily complex models

# Adjusted $R^2$ : A First Approach

## Adjusted $R^2$ Formula

$$R_{\text{adj}}^2 = 1 - \frac{\frac{n}{n-p-1} \text{RSS}}{\sum_i (y_i - \bar{y})^2}$$

- **Improvement:** Penalizes for model complexity via  $\frac{n}{n-p-1}$  factor
- $p$  = number of predictors in the model
- **Theoretical limitation:** Penalty is too weak
- Often selects models that are still too complex
- Better than regular  $R^2$ , but inferior to AIC/BIC/CV for model selection

# Akaike Information Criterion (AIC)

## AIC for Linear Regression with Gaussian Errors

$$\text{AIC} = -\frac{n}{2} \log \left( \frac{\text{RSS}}{n} \right) - (p + 2)$$

- $p$  = number of coefficients (excluding intercept)
- **Interpretation:** Higher AIC values indicate better models
- **Note:** Different resources may multiply AIC by  $-2$  or  $2$ , changing interpretation
- **Theoretical foundation:** AIC is an unbiased estimator of expected out-of-sample prediction error
- **Computational advantage:** Fast approximation of cross-validation

# AIC and Cross-Validation: Theoretical Properties

- As sample size  $n \rightarrow \infty$ :
  - Both AIC and CV select models with near-optimal generalization error
  - Both tend to select slightly more complex models than the theoretical optimum
  - This happens because more complex models have higher variability in their error estimates

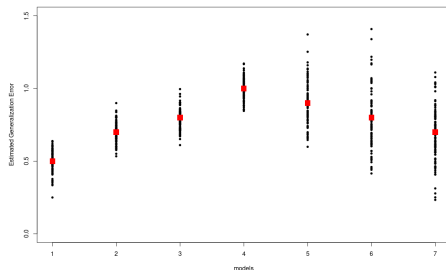


Figure: AIC vs. True Error

# Bayesian Information Criterion (BIC)

## BIC for Linear Regression with Gaussian Errors

$$\text{BIC} = -\frac{n}{2} \log \left( \frac{\text{RSS}}{n} \right) - \frac{\log(n)}{2} (p + 2)$$

- **Key difference from AIC:** Complexity penalty grows with sample size  $n$
- **Comparison:**
  - AIC penalty:  $(p + 2)$
  - BIC penalty:  $\frac{\log(n)}{2} (p + 2)$
- **Consequence:** BIC favors simpler models than AIC, especially with large datasets
- **Theoretical property:** BIC is consistent - will select the true model as  $n \rightarrow \infty$  (if true model exists in candidate set)

# Comparing Model Selection Criteria

| Criterion        | Complexity Penalty | Consistency | Prediction | Computation |
|------------------|--------------------|-------------|------------|-------------|
| $R^2$            | None               | No          | Poor       | Fast        |
| Adjusted $R^2$   | Weak               | No          | Moderate   | Fast        |
| AIC              | Moderate           | No          | Good       | Fast        |
| BIC              | Strong             | Yes         | Good*      | Fast        |
| Cross-Validation | Flexible           | No          | Excellent  | Slow        |

**Table:** \*BIC may underperform AIC for prediction with finite samples

- **No perfect criterion** - choice depends on goals:
  - **Pure prediction:** Cross-validation or AIC
  - **Finding true model:** BIC (if you believe a true model exists)

# Computational Strategies for Model Selection

# The Computational Challenge

## Number of Possible Models

With  $p$  potential predictors, we have  $2^p$  possible models

| Number of predictors | Possible models                                        | Feasibility |
|----------------------|--------------------------------------------------------|-------------|
| 5                    | $2^5 = 32$                                             | Trivial     |
| 10                   | $2^{10} = 1,024$                                       | Easy        |
| 20                   | $2^{20} = 1,048,576$                                   | Challenging |
| 30                   | $2^{30} \approx 1 \text{ billion}$                     | Impractical |
| 60                   | $2^{60} \approx \text{number of sand grains on Earth}$ | Impossible  |

**We need efficient search strategies!**

# Branch and Bound Algorithm

## Key Insight

We can eliminate entire groups of models without evaluating each one individually

- **Consider** a set of models  $\{M_1, M_2, \dots, M_S\}$
- Let  $M_{\text{sup}}$  be a model containing all covariates that appear in at least one model in our set
- Let  $p_{\text{min}}$  be the number of covariates in the smallest model in our set
- For all models  $M_s$  in our set:  $\text{RSS}(M_s) \geq \text{RSS}(M_{\text{sup}})$

## Upper Bound for AIC

$$\text{AIC}(M_s) \leq -\frac{n}{2} \log \left( \frac{\text{RSS}(M_{\text{sup}})}{n} \right) - (p_{\text{min}} + 2)$$

# Branch and Bound: Practical Application

- **Key advantage:** If we find any model with AIC exceeding our upper bound, we can eliminate all models in our set without evaluating them
- **Effectiveness varies:**
  - Best case: Drastically reduces computation
  - Worst case: Similar to exhaustive search
- **Practical limit:** Up to 30 predictors
- **Software implementation:** leaps package in R
- **Applicability:** Works with both AIC and BIC
- **Limitation:** Specialized for linear regression (uses computational tricks specific to linear models)

# Stepwise Selection Methods

## When Exhaustive Search Is Infeasible

Stepwise methods provide heuristic approaches that:

- Are computationally efficient
- Work for various model types (not just linear regression)
- Usually find good (though not guaranteed optimal) solutions

## Two Main Approaches

- **Forward selection:** Start simple, add variables
- **Backward selection:** Start complex, remove variables

# Forward Selection Algorithm

## How Forward Selection Works

A greedy algorithm that builds a model iteratively:

- Start with an empty model (no predictors)
- At each step, add the most significant variable
- Continue until no remaining variables improve the model

## Algorithm Steps

- 1 Start with an intercept-only model
- 2 For each candidate variable not in the model:
  - Fit model with the variable added
  - Compute selection criterion (e.g., p-value, AIC, BIC)
- 3 Add variable that most improves the criterion (if significant)
- 4 Repeat steps 2-3 until no further improvement

# Forward Selection Algorithm

## Advantages & Limitations

### Advantages

- Computationally efficient
- Easy to implement
- Interpretable process

### Limitations

- May miss optimal subset
- Ignores multicollinearity
- Cannot remove variables once added

# Backward Selection Algorithm

## How Backward Selection Works

A pruning algorithm that simplifies a model iteratively:

- Start with the full model (all predictors)
- At each step, remove the least significant variable
- Continue until all remaining variables are significant

## Algorithm Steps

- 1 Start with all variables in the model
- 2 For each variable currently in the model:
  - Compute significance measure (e.g., p-value, t-statistic)
  - Identify least significant variable
- 3 Remove least significant variable if below threshold
- 4 Repeat steps 2-3 until all variables meet significance criterion

# Backward Selection Algorithm

## Advantages & Limitations

### Advantages

- Considers interactions between all variables initially
- May detect effects hidden by confounders
- Works well with many predictor variables

### Limitations

- Requires fitting full model first
- Computationally intensive with many variables
- Cannot reconsider variables once removed

# Computational Complexity of Stepwise Methods

- **Forward selection:**

- At step  $i$ : evaluate  $(p - i)$  models
- Maximum steps:  $p$
- Total evaluations:  $\mathcal{O}(p^2)$

- **Backward selection:**

- At step  $i$ : evaluate  $i$  models
- Maximum steps:  $p$
- Total evaluations:  $\mathcal{O}(p^2)$

- **Compared to exhaustive:**  $\mathcal{O}(p^2)$  vs.  $\mathcal{O}(2^p)$

- **Important note:** Forward and backward selection may yield different models
- **Hybrid approaches:** Can allow both addition and removal at each step

# Summary: Model Selection Best Practices

## Selection Criteria

- **Avoid:**  $R^2$  (no penalty)
- **Adequate:** Adjusted  $R^2$  (weak penalty)
- **Better:** AIC/BIC (theoretically grounded)
- **Best:** Cross-validation (directly measures generalization)

## Search Strategies

- **Small  $p$  ( $< 30$ ):** Branch and bound
- **Large  $p$ :** Stepwise methods
- **Modern alternatives:** LASSO, Elastic Net (next lecture)

## Final Recommendations

- Always validate your final model on holdout data
- Consider your goals: prediction vs. explanation
- Incorporate domain knowledge when possible
- Remember model selection introduces additional uncertainty