# CMSC/LING/STAT 208: Machine Learning

Abhishek Chakraborty [Much of the content in these slides have been adapted from *ISLR2* by James et al. and *HOMLR* by Boehmke & Greenwell]

## Remedies for Class Imbalance

For binary classification problems, in some contexts, the class of interest (Positive class) might have a very low frequency. We will see how to approach the ML process when such cases arise.

We will look at three possible remedies.

- Model tuning
- · Alternative cutoffs
- Subsampling

We will work with the attrition.rds dataset.

## Linear Model Selection and Regularization

The standard linear model,

$$Y = \beta_0 + \beta_1 X_1 + \dots + \beta_p X_p + \epsilon$$

- · Benefits: Simple, interpretable, often shows good predictive performance.
- · Limitations:
  - **Prediction Accuracy**: If n is not big (also when p>n), variance can be high resulting in overfitting and poor predictive performance.
  - Model Interpretability: Irrelevant features lead to unnecessary model complexity.

## Alternatives to Least Squares

Extensions/Modifications/Improvements to Least Squares:

- Subset Selection: Identify a subset of p predictors and then fit a linear model using least squares.
- Shrinkage/Regularization: Fit a model using p predictors, but shrink some of the estimated coefficients towards zero. Reduces variance and can also perform variable selection.
- **Dimension Reduction**: Project p predictors onto a M-dimensional subspace, where M < p. Achieved by computing M different linear combinations or projections of the p predictors. Fit a linear model using these M predictors by least squares.

# Shrinkage/Regularization Methods

Fit a model containing all p predictors using a technique that **shrinks** the coefficient estimates towards zero.

- · Ridge Regression
- · Lasso

Shrinking the coefficient estimates significantly reduces their variance.

## The Lasso

Acronym for Least Absolute Shrinkage and Selection Operator.

· Standard Linear Model

Given a training dataset, for  $i=1,\ldots,n$ 

$$\hat{y}_i = b_0 + b_1 x_{i1} + \ldots + b_p x_{ip}$$

$$ext{SSE} = \sum_{i=1}^n \left( y_i - \hat{y}_i 
ight)^2 = \sum_{i=1}^n \left( y_i - (b_0 + b_1 x_{i1} + \ldots + b_p x_{ip}) 
ight)^2$$

· Lasso

$$\mathrm{SSE} + \lambda \sum_{j=1}^p |b_j|$$

- .  $\lambda \sum_{j=1}^p |b_j|$ : Shrinkage Penalty
- ·  $\lambda \geq 0$ : Tuning/Regularization Parameter

## The Lasso

- This method not only shrinks the coefficient estimates towards zero, but also makes some of the coefficient estimates exactly equal to zero (when the tuning parameter  $\lambda$  is sufficiently large).
- · Hence, the lasso performs variable selection.
- · We say that the lasso yields **sparse models**, that is, models that involve only a subset of the variables.

# The Lasso: Scaling of Predictors

- Standard least squares (regression) coefficient estimates are **scale equivariant**: multiplying  $X_j$  by a constant c simply leads to a scaling of the least squares coefficient estimates by a factor of 1/c. In other words, regardless of how the  $j^{th}$  predictor is scaled,  $X_j \hat{\beta}_j$  will remain the same.
- · In contrast, the lasso coefficient estimates can change substantially when multiplying a given predictor by a constant. Apply lasso after **standardizing** the predictors.

$$\tilde{x}_{ij} = \frac{x_{ij}}{\sqrt{\frac{1}{n} \sum_{i=1}^{n} (x_{ij} - \overline{x}_j)^2}}$$

```
# create recipe and blueprint, prepare and apply blueprint
set.seed(050724) # set seed
ames recipe <- recipe(Sale Price ~ ., data = ames train) # set up recipe
blueprint <- ames recipe %>%
 step_nzv(Street, Utilities, Pool_Area, Screen_Porch, Misc_Val) %>% # filter out zv/nzv predictors
 step impute mean(Gr Liv Area) %>%
                                                                       # impute missing entries
                                                                       # numeric conversion of levels of the predictors
 step integer(Overall Qual) %>%
 step center(all numeric(), -all outcomes()) %>%
                                                                       # center (subtract mean) all numeric predictors
 step scale(all numeric(), -all outcomes()) %>%
                                                                       # scale (divide by standard deviation) all numeric predictors
                                                                       # Lumping required predictors
 step other(Neighborhood, threshold = 0.01, other = "other") %>%
 step dummy(all nominal(), one hot = FALSE)
                                                                       # one-hot/dummy encode nominal categorical predictors
prepare <- prep(blueprint, data = ames_train) # estimate feature engineering parameters based on training data</pre>
baked train <- bake(prepare, new data = ames train) # apply blueprint to training data
baked test <- bake(prepare, new data = ames test)</pre>
                                                     # apply blueprint to test data
```

#### **Ames Housing Dataset**

## [1] 42593.99

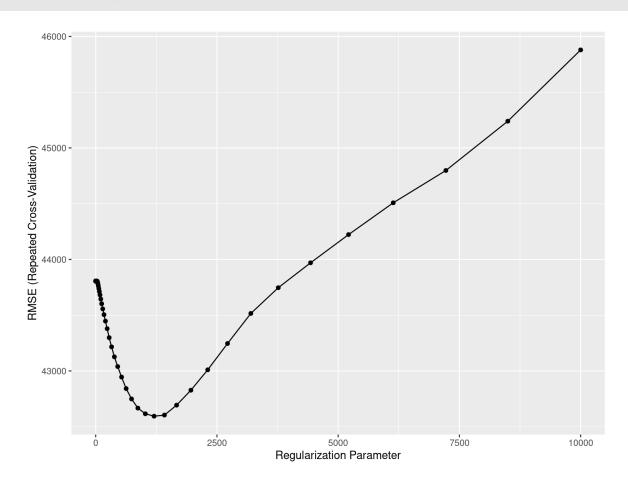
Implement CV to tune the hyperparameter  $\lambda$ .

```
set.seed(050724) # set seed
cv_specs <- trainControl(method = "repeatedcv", number = 5, repeats = 5) # CV specifications</pre>
lambda grid <- 10^seq(-3, 4, length = 100) # grid of lambda values to search over
library(glmnet) # for LASSO
lasso_cv <- train(blueprint,</pre>
                  data = ames train,
                  method = "glmnet", # for lasso
                  trControl = cv_specs,
                  tuneGrid = expand.grid(alpha = 1, lambda = lambda grid), # alpha = 1 implements lasso
                  metric = "RMSE")
# results from the CV procedure
lasso cv$bestTune$lambda # optimal Lambda
## [1] 1204.504
min(lasso cv$results$RMSE) # RMSE for optimal Lambda
```

## **Ames Housing Dataset**

Results from the CV procedure.

ggplot(lasso\_cv) # Lambda vs. RMSE plot



#### **Ames Housing Dataset**

We will now build the optimal lasso model on the modified training data using the optimal  $\lambda$ .

```
# create datasets required for 'glmnet' function
X_train <- model.matrix(Sale_Price ~ ., data = baked_train)[, -1] # training features without intercept</pre>
Y_train <- baked_train$Sale_Price # training response
X test <- model.matrix(Sale Price ~ ., data = baked test)[, -1] # test features without intercept
# build optimal lasso model
final model <- glmnet(x = X train,</pre>
                      y = Y train,
                      alpha = 1,
                                                           # alpha = 1 builds lasso model
                      lambda = lasso cv$bestTune$lambda, # using optimal lambda from CV
                      standardize = FALSE)
                                                           # already standardized during data preprocessing
# obtain predictions and test set RMSE
final model preds <- predict(final model, newx = X test)</pre>
                                                            # obtain predictions
sqrt(mean((final model preds - baked test$Sale Price)^2))
                                                            # calculate test set RMSE
## [1] 29903.92
```

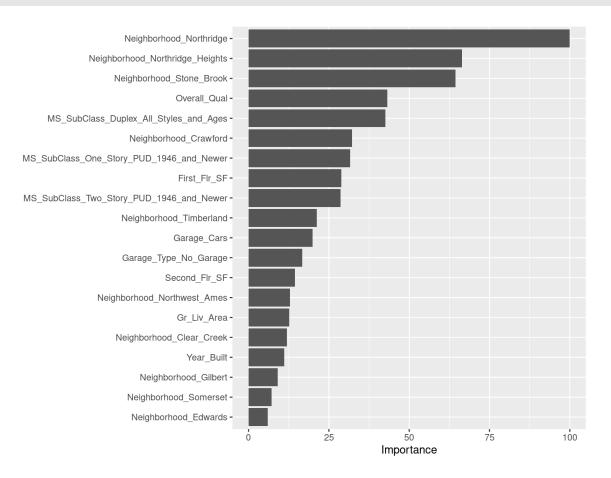
#### Ames Housing Dataset

The coefficients for the optimal lasso model can be obtained from

coef(final\_model) # estimated coefficients from final Lasso model

##	54 x 1 sparse Matrix of class "dgCMatrix"	
##		s0
##	(Intercept)	181313.5819
##	Gr_Liv_Area	6917.8414
##	Garage_Cars	10466.0529
##	Garage_Area	217.2498
##	Overall_Qual	35210.8237
##	Lot_Area	2714.6661
##	Lot_Frontage	612.9751
##	Open_Porch_SF	316.2876
##	TotRms_AbvGrd	-338.8492
##	First_Flr_SF	23960.4079
##	Second_Flr_SF	12380.3553
##	Year_Built	7099.8829
##	Garage_Type_Basment	
##	Garage_Type_BuiltIn	
##	Garage_Type_CarPort	
##	Garage_Type_Detchd	
##	Garage_Type_More_Than_Two_Types	
##	Garage_Type_No_Garage	
##	Neighborhood_College_Creek	
##	Neighborhood_Old_Town	
##	Neighborhood_Edwards	
##	Neighborhood_Somerset	
##	Neighborhood_Northridge_Heights	15388.8984
##	Neighborhood_Gilbert	
##	Neighborhood_Sawyer	•
##	Neighborhood_Northwest_Ames	•
##	Neighborhood_Sawyer_West	
##	Neighborhood_Mitchell	
##	Neighborhood_Brookside	
##	Neighborhood_Crawford	
##	Neighborhood_Iowa_DOT_and_Rail_Road	•
##	Neighborhood_Timberland	
##	Neighborhood_Northridge	5196.8736
##	Neighborhood_Stone_Brook	•
##	${\tt Neighborhood\_South\_and\_West\_of\_Iowa\_State\_University}$	•

```
# variable importance
vip(object = lasso_cv, num_features = 20, method = "model")
```



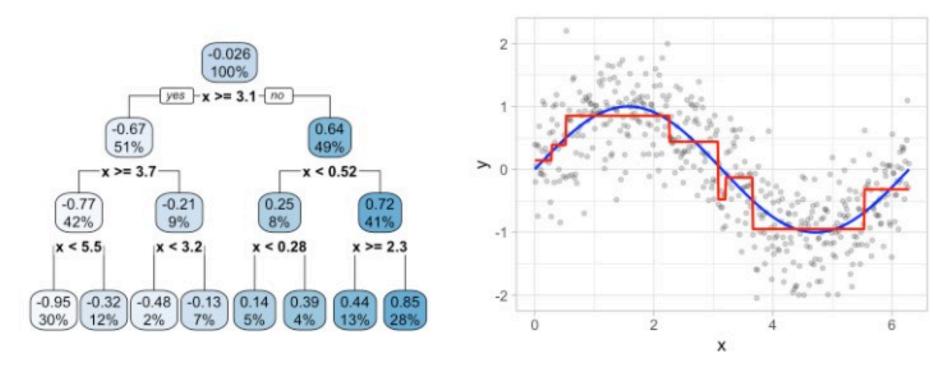
## **Tree-Based Methods**

- Involves **stratifying** or **segmenting** the predictor space into a number of simple regions.
- The set of splitting rules used to segment the predictor space can be summarized in a tree, thus, the name **decision tree** methods.
- · Can be used for both classification and regression.
- Tree-based methods are simple and useful for interpretation, however, not the best in terms of prediction accuracy.
- Methods such as bagging, random forests, and boosting grow multiple trees and then combine their results.

# **Terminology for Trees**

- · Every split is considered to be a **node**.
- We refer to the first node at the top of the tree as the **root node** (this node contains all of the training data).
- The final nodes at the bottom of the tree are called the **terminal nodes** or **leaves**.
- Decision trees are typically drawn **upside down**, in the sense that the leaves are at the bottom of the tree.
- The points along the tree where the predictor space is split are referred to as **internal nodes**, that is, every node in between the **root node** and **terminal nodes** is referred to as an **internal node**.
- The segments of the trees that connect the nodes are known as **branches**.

# **Terminology for Trees**



Adapted from HMLR, Boehmke & Greenwell

## **Building a Tree**

• First select the predictor  $X_j$  and the cutpoint s such that splitting the predictor space into the regions  $\{X|X_j < s\}$  and  $\{X|X_j \geq s\}$  leads to the greatest possible reduction in SSE.

For any j and s, define

$$R_1 = \{X | X_j < s\} \text{ and } R_2 = \{X | X_j > s\}$$

Find j and s that minimize

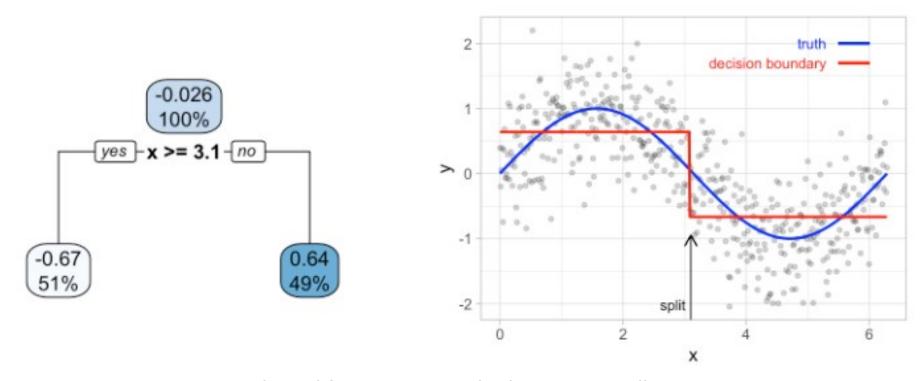
$$SSE = \sum_{i \in R_1} \left( y_i - \hat{y}_{R_1} 
ight)^2 + \sum_{i \in R_2} \left( y_i - \hat{y}_{R_2} 
ight)^2$$

- Next, repeat the process, look for the best predictor and best cutpoint in order to split the data further. However, this time, instead of splitting the entire predictor space, split one of the two previously identified regions.
- The process continues until a stopping criterion is reached; say, we may continue until no region contains more than five observations.

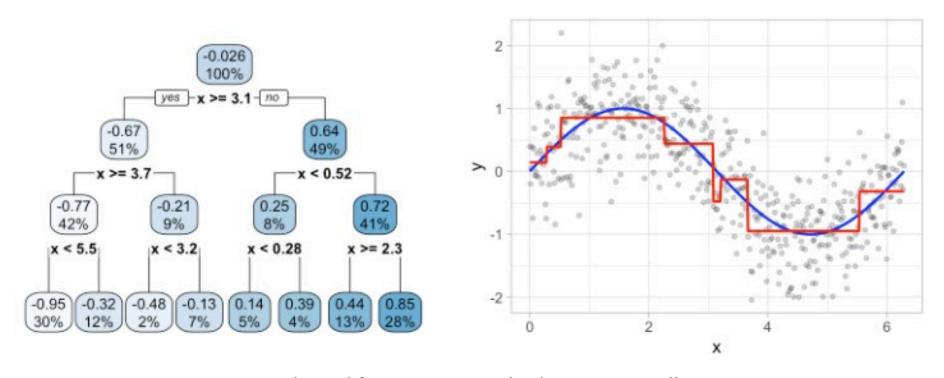
## **Prediction**

For every observation that falls into the region  $R_j$ , make the same prediction, which is

- · the mean response of the training set observations in  $R_j$  (for regression problems),
- · majority vote response of the training set observations in  $R_j$  (for classification problems).

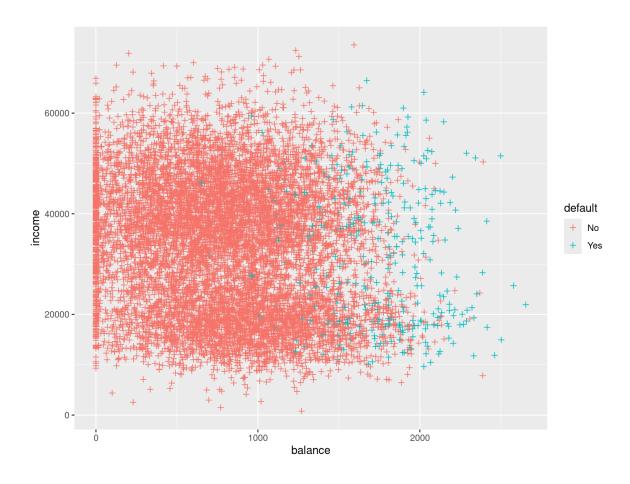


Adapted from HMLR, Boehmke & Greenwell

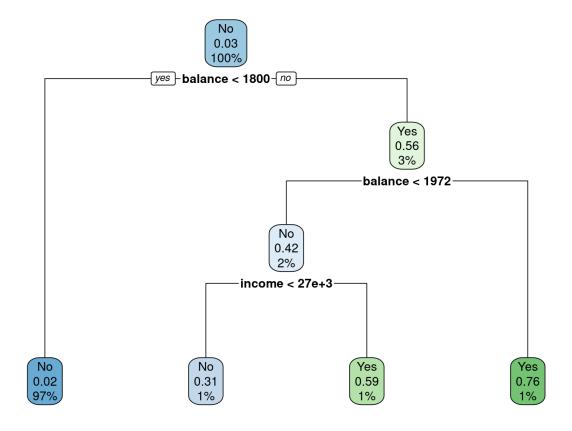


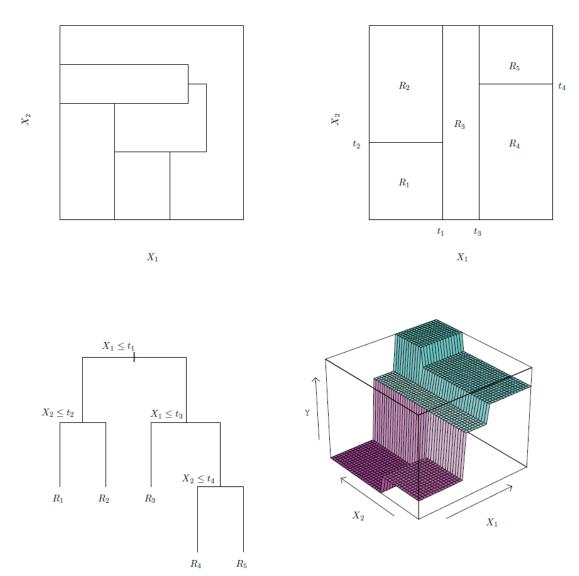
Adapted from HMLR, Boehmke & Greenwell

## **Default Dataset**



### **Default Dataset**





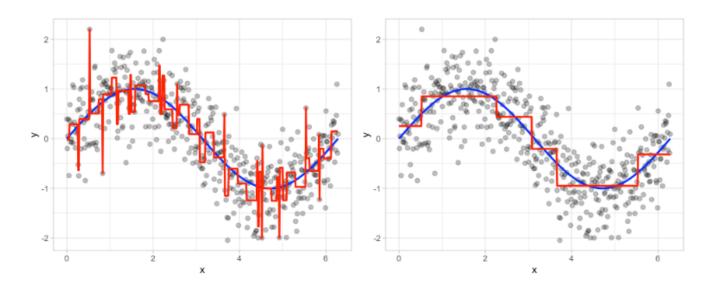
Adapted from ISLR, James et al.

# **Building a Tree**

- · It is computationally infeasible to consider every possible partition of the feature space into J boxes.
- For this reason, we take a **top-down**, **greedy** approach known as **recursive binary splitting**.
  - **top-down** because it begins at the top of the tree and then successively splits the predictor space.
  - **greedy** because at each step of the tree-building process, the best split is made at that particular step, rather than looking ahead and picking a split that will lead to a better tree in some future step.

# **Tree Pruning**

The process described above may **overfit** the data.



# **Tree Pruning**

- Grow a very large tree, and then **prune** it back to obtain a **subtree**.
- The technique uses is known as **cost complexity pruning** (also known as **weakest link pruning**).
- · Consider a sequence of trees indexed by  $\alpha$ . For each  $\alpha$ , consider the tree that minimizes

$$SSE + \alpha |T|$$

· Choose optimal  $\alpha$  by CV.

```
# create recipe and blueprint, prepare and apply blueprint
set.seed(050924) # set seed
ames recipe <- recipe(Sale Price ~ ., data = ames train) # set up recipe
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```

#### **Ames Housing Dataset**

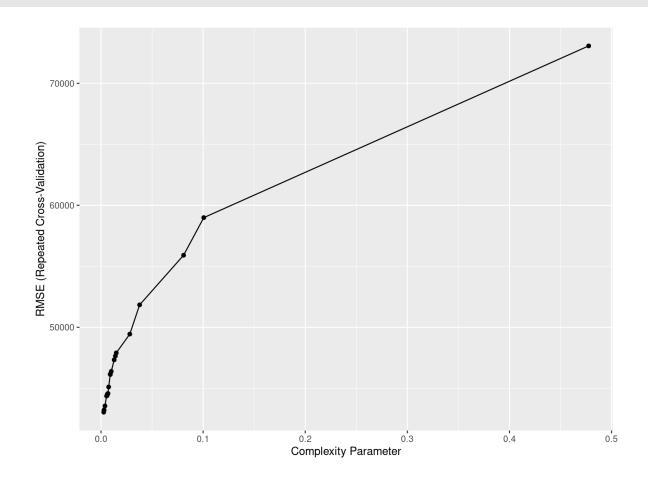
Implement CV to tune the hyperparameter.

```
set.seed(050924) # set seed
cv specs <- trainControl(method = "repeatedcv", number = 5, repeats = 5) # CV specifications</pre>
library(rpart) # for trees
tree cv <- train(blueprint,</pre>
                 data = ames_train,
                 method = "rpart",
                 trControl = cv_specs,
                 tuneLength = 20,
                                                    # considers a grid of 20 possible tuning parameter values
                 metric = "RMSE")
# results from the CV procedure
tree cv$bestTune
                   # optimal hyperparameter
              ср
## 1 0.002631828
min(tree_cv$results$RMSE) # optimal CV RMSE
## [1] 43039.21
```

#### **Ames Housing Dataset**

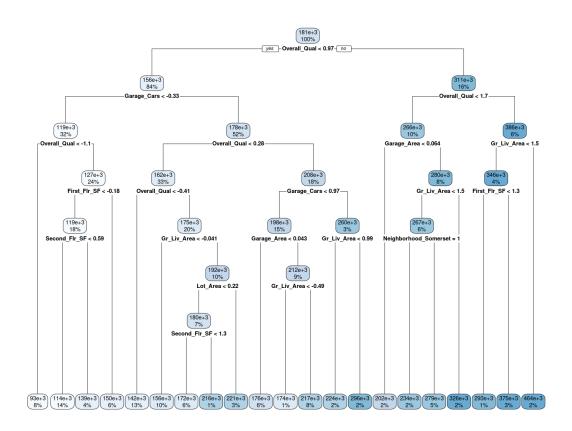
Results from the CV procedure.

ggplot(tree\_cv)



## **Ames Housing Dataset**

library(rpart.plot)
rpart.plot(final\_model)



```
# variable importance
vip(object = tree_cv, num_features = 20, method = "model")
```

