# **Classification Performance**

#### Nate Wells

Math 243: Stat Learning

October 2nd, 2020

Nate Wells (Math 243: Stat Learning)

# Outline

In today's class, we will...

- Implement KNN in R
- Analyze the performance of classification models
- Work in groups on a classification problem

# Section 1

 $\mathsf{KNN}\xspace$  in  $\mathsf{R}\xspace$ 

Nate Wells (Math 243: Stat Learning)

KNN in R	Additional Practice
000000	

Recall: The KNN model estimates the conditional probability  $P(Y = A_j | X)$  as

$$P(Y = A_j \mid X = x_0) \approx \frac{1}{K} \sum_{i \in N_0} I(y_i = A_j)$$

Recall: The KNN model estimates the conditional probability  $P(Y = A_j | X)$  as

$$P(Y = A_j \mid X = x_0) \approx \frac{1}{K} \sum_{i \in N_0} I(y_i = A_j)$$

In R, we use the knn function in the class library.

Recall: The KNN model estimates the conditional probability  $P(Y = A_j | X)$  as

$$P(Y = A_j \mid X = x_0) \approx \frac{1}{K} \sum_{i \in N_0} I(y_i = A_j)$$

In R, we use the knn function in the class library.

The knn function fits a model and makes predictions all in one command.

Recall: The KNN model estimates the conditional probability  $P(Y = A_j | X)$  as

$$P(Y = A_j \mid X = x_0) \approx \frac{1}{K} \sum_{i \in N_0} I(y_i = A_j)$$

In R, we use the knn function in the class library.

The knn function fits a model and makes predictions all in one command.

• Unlike the lm and glm, which first fit a model and then make predictions using the predict function

KNN in R 00●0000	Model Performance 0000000	Additional Practice OO

# Simulated Data

Suppose  $X|Y = 1 \sim N(0, 1)$  and  $X|Y = 0 \sim N(1, 1)$ , and that each class is of the same size.

KNN in R	Model Performance	Additional Practice
00●0000	0000000	00

## Simulated Data

Suppose  $X|Y = 1 \sim N(0, 1)$  and  $X|Y = 0 \sim N(1, 1)$ , and that each class is of the same size.

• We'll also subset our data into test and training sets.

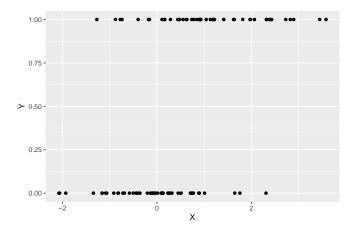
# Simulated Data

Suppose  $X|Y = 1 \sim N(0,1)$  and  $X|Y = 0 \sim N(1,1)$ , and that each class is of the same size.

• We'll also subset our data into test and training sets.

```
set.seed(100)
n<-100
Y<-rep(c(0,1), c(n/2, n/2))
X<-c(rnorm(n/2, 0, 1), rnorm(n/2, 1, 1) )
d<-data.frame(X,Y)
library(dplyr)
train_d<-d %>% sample_frac(.75)
test_d<-anti_join(d, train_d)</pre>
```

# Scatterplot



# The KNN Model

The knn function takes 4 arguments.

- A data frame containing the predictors associated to the training data
- 2 A data frame containing the predictors associated to the test data
- 8 A vector containing the response associated to the training data
- **4** A value for K, the number of nearest neighbors.

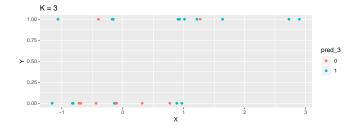
# The KNN Model

The knn function takes 4 arguments.

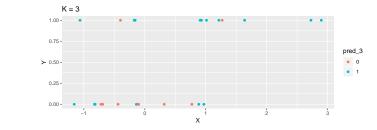
- ① A data frame containing the predictors associated to the training data
- A data frame containing the predictors associated to the test data
- 8 A vector containing the response associated to the training data
- **4** A value for K, the number of nearest neighbors.

```
library(class)
set.seed(200)
pred_3<-knn(train_d %>% select(X),
    test_d %>% select(X),
    train_d$Y,
    3)
pred_3
## [1] 1 0 1 0 0 1 0 1 1 1 0 0 1 0 1 0 1 1 1 1 1 0 1 1 1 1
## Levels: 0 1
```

# Results K = 3

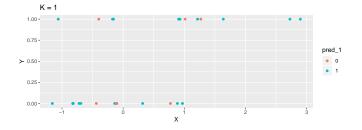


#### Results K = 3

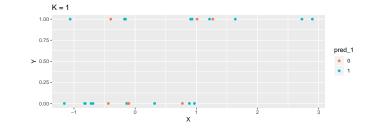




# Results K = 1



#### Results K = 1



##
## 0 1
## 0 4 9
## 1 3 9
## [1] 0.52

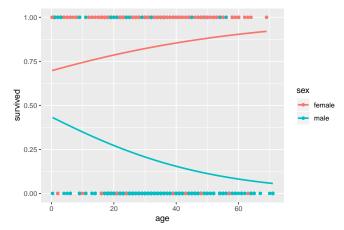
# Section 2

# Model Performance

Nate Wells (Math 243: Stat Learning)

## The Unsinkable Example

The Titanic data set contains information on passengers of the Titanic



#### A better confusion matrix

The confusionMatrix function in the caret package provides a confusion matrix along withe relevant statistics:

```
library(caret)
confusionMatrix(data = factor(preds) , reference = factor(Titanic1$survived) )
## Confusion Matrix and Statistics
##
##
             Reference
## Prediction 0 1
##
            0 308 82
            1 44 199
##
##
                  Accuracy : 0.8009
##
##
                    95% CI : (0.7677, 0.8314)
##
       No Information Rate : 0.5561
       P-Value [Acc > NTR] : < 2.2e-16
##
##
##
                     Kappa : 0.5912
##
##
    Moneman's Test P-Value : 0.0009799
##
##
               Sensitivity : 0.8750
##
               Specificity : 0.7082
##
            Pos Pred Value : 0.7897
##
            Neg Pred Value : 0.8189
##
                Prevalence : 0.5561
##
            Detection Rate : 0.4866
##
      Detection Prevalence : 0.6161
##
         Balanced Accuracy : 0.7916
##
##
          'Positive' Class · 0
##
```

Sensitivity: Rate of correct positive identification

• Type II Error rate: 1 – Sensitivity

Specificity: Rate of correct negative identification

• Type I Error rate: 1 - Specificity

Sensitivity: Rate of correct positive identification

• Type II Error rate: 1 – Sensitivity

Specificity: Rate of correct negative identification

• Type I Error rate: 1 - Specificity

By changing our classification cutoff, we can increase sensitivity to the detriment of specificity (or vice versa)

• But the tradeoff is non-linear

Sensitivity: Rate of correct positive identification

• Type II Error rate: 1 – Sensitivity

Specificity: Rate of correct negative identification

• Type I Error rate: 1 – Specificity

By changing our classification cutoff, we can increase sensitivity to the detriment of specificity (or vice versa)

- But the tradeoff is non-linear
  - Increasing specificity by .1 may decrease sensitivity by .15 when specificity is .8
  - But the same increase in specificity may decrease sensitivity by .25 when specificity is .9.

Sensitivity: Rate of correct positive identification

• Type II Error rate: 1 – Sensitivity

Specificity: Rate of correct negative identification

• Type I Error rate: 1 - Specificity

By changing our classification cutoff, we can increase sensitivity to the detriment of specificity (or vice versa)

- But the tradeoff is non-linear
  - Increasing specificity by .1 may decrease sensitivity by .15 when specificity is .8
  - But the same increase in specificity may decrease sensitivity by .25 when specificity is .9.

When might we want high specificity? High sensitivity?

Sensitivity: Rate of correct positive identification

• Type II Error rate: 1 – Sensitivity

Specificity: Rate of correct negative identification

• Type I Error rate: 1 - Specificity

By changing our classification cutoff, we can increase sensitivity to the detriment of specificity (or vice versa)

- But the tradeoff is non-linear
  - Increasing specificity by .1 may decrease sensitivity by .15 when specificity is .8
  - But the same increase in specificity may decrease sensitivity by .25 when specificity is .9.

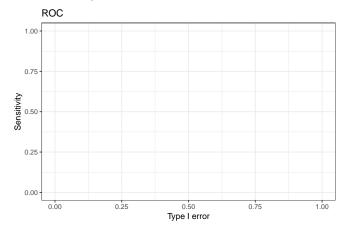
When might we want high specificity? High sensitivity?

What are the ramifications of changing the classification cutoff?

	Model Performance	Additional
000000	0000000	00

### **ROC Curves**

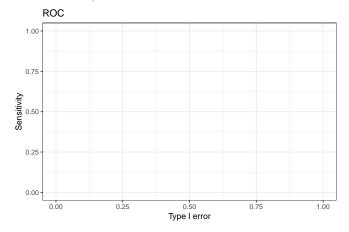
A Receiver Operating Characteristic (ROC) curve is a plot of sensitivity vs. type I error rate, based on classification probabilities.



	Model Performance	Additional Pra
000000	0000000	00

## **ROC Curves**

A Receiver Operating Characteristic (ROC) curve is a plot of sensitivity vs. type I error rate, based on classification probabilities.

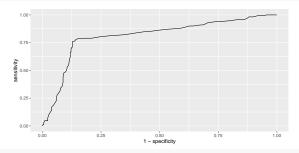


Poll: For a perfectly accurate model, what is the expected area under the ROC curve?

### ROC Curves in R

The roc function in the pROC package can create ROC curves.

```
library(pROC)
roc_curve <- roc(response = Titanic1$survived, predictor = probs)
ggroc(roc_curve, legacy.axes=TRUE)</pre>
```

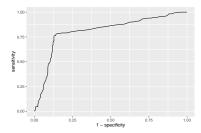


auc(roc\_curve)

**##** Area under the curve: 0.8095

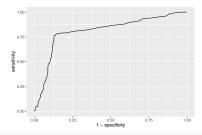
### ROC Curves in R

What threshold corresponds to the "kink" in the ROC curve?



### ROC Curves in R

What threshold corresponds to the "kink" in the ROC curve?



```
coords(roc_curve, "best", ret = "threshold")
```

```
## threshold
```

```
## 1 0.2533806
```

coords(roc\_curve, .253)

## threshold specificity sensitivity
## 1 0.253 0.8522727 0.7829181

# Section 3

# Additional Practice

Nate Wells (Math 243: Stat Learning)

### Mushroom Hunting

The mushrooms data set on the schedule page of the course website contains information on several species of mushrooms, including edibility.

## Mushroom Hunting

The mushrooms data set on the schedule page of the course website contains information on several species of mushrooms, including edibility.

Can we predict whether a mushroom is edible?

### Mushroom Hunting

The mushrooms data set on the schedule page of the course website contains information on several species of mushrooms, including edibility.

Can we predict whether a mushroom is edible?

- Create a Logistic Regression model using your choice of a small subset of predictors
  - You will need to recode your response class to take values 0 or 1.
- Then create an ROC curve and select a threshold that seems appropriate for this situation.
- Time permitting, create a KNN model for various values of K and compare to the logistic regression model.