Confusion Matrix

Machine Learning - Section 2.1.2

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We previously defined a prediction rule that predicts male any height above 64 inches. However, given the fact that female average height is about 65 inches, there appears to be a conflict with our rule: if the average female height is 65 inches, why does our rule tell us to predict male for those who are 65 inches tall?

Overall accuracy can be a deceptive measure. -Dr. Rafael Irizarry

We can see this by constructing a Confusion Matrix: A tabulation of prediction and actual value.

```
table(predicted = y_hat, actual = test_set$sex)
```

```
## actual
## predicted Female Male
## Female 50 27
## Male 69 379
```

The table above is read like so:

- Subjects that were actually female and predicted to be female: 50
- Subjects that were actually male but predicted to be female: 27
- Subjects that were actually female but predicted to be male: 69
- Subjects that were actually male and predicted to be male: 379

Looking at the above values closely, a problem emerges: male's are over represented in our data set. Computing the accuracy separately for each sex reflects as much:

```
test_set %>%
  mutate(y_hat = y_hat) %>%
  group_by(sex) %>%
  summarize(accuracy = mean(y_hat == sex))
```

sex	accuracy	
Female	0.420	
Male	0.933	

Our algorithm has very high accuracy when predicting male: 93.3%, but extremely low accuracy when predicting female: 42.0%. Too many female's are predicted to be male.

Our high overall accuracy despite our low female accuracy is due to **Prevalence**: There are more male's in our data set than female's.

```
prev = mean(y == "Male")
prev
```

[1] 0.773

What this means is that the *incorrect* predictions of female's is **outweighed** by correctly predicting more male's. This is a significant problem in Machine Learning: If our training data is biased, our algorithm will be biased. Therefore, we need to look at metrics other than overall accuracy, that are robust to prevalence, when evaluating a Machine Learning algorithm.

Derived Metrics from the Confusion Matrix: Sensitivity and Specificity

A general improvement to only studying overall accuracy is to study **Sensitivity** and **Specificity** separately. These two metrics can be defined with a binary outcome. When the outcomes are **categorical**, we can define these metrics for a *specific* category. For example, when predicting digits, we can calculate the Sensitivity and Specificity of correctly predicting "2" as opposed to some other digit. By selecting "2" we've specified a category of interest.

When Y = 1, we will define these outcomes as **positive outcomes**. When Y = 0, we will define these outcomes as **negative outcomes**.

Sensitivity is defined as the ability of an algorithm to **predict a positive outcome**, $\hat{Y} = 1$, when the **actual outcome is positive**, Y = 1. However, because an algorithm that predicts all Y's to be positive (i.e. an algorithm has $\hat{Y} = 1$, no matter what) has **perfect sensitivity**, Sensitivity alone is not adequate when evaluating algorithms.

Specificity is defined as the ability of an algorithm to **not predict a positive outcome**, $\hat{Y} = 0$, when the **actual outcome is not positive**, Y = 0. We can summarize like so:

High Sensitivity:
$$Y = 1 \Rightarrow \hat{Y} = 1$$

High Specificity :
$$Y = 0 \Rightarrow_{implies} \hat{Y} = 0$$

There is a second way to define **Specificity**: The proportion of **positive predictions** that are **actually positive**(i.e. (correct positive predictions) / (all positive predictions)). In this case high Specificity is defined as:

High Specificity:
$$\hat{Y} = 1 \underset{implies}{\Rightarrow} Y = 1$$

To differentiate between these metrics, each of the four entries in the confusion matrix has a unique name.

	Actually.Positive	Actually.Negative
Predicted + Predicted -	True Positives(TP) False Negatives(FN)	False Positives(FP) True Negatives(TN)

- When an **outcome** is **positive** and was **predicted** to be **positive**: True Positives (TP)
- When an outcome is negative but was predicted to be positive: False Positives (FP)
- When an outcome is positive but was predicted to be negative: False Negatives (FN)
- When an outcome is negative and was predicted to be negative: True Negatives (TN)

With these definitions, we can accurately quantify Sensitivity and Specificity:

$${\rm Sensitivity}_1 = \frac{TP}{(TP+FN)}$$

$${\rm Specificity}_2 = \frac{TN}{(TN+FP)}$$

There is another way of quantifying Specificity:

$${\rm Specificity}_3 = \frac{TP}{(TP+FP)}$$

To help in understanding the above, keep in mind that:

$$(TP + FN) = All positive outcomes$$

$$(TN + FP) = All negative outcomes$$

$$(TP + FP)$$
 = All positive predictions

Note that unlike the True Positive Rate and the True Negative Rate, **Precision** is affected by **Prevalence**. The higher the Prevalence, the higher the Precision.

 $^{^{1}}$ This formulation of Sensitivity is also known as the **True Positive Rate** (**TPR**) or **Recall**.

 $^{^2}$ This formulation of Specificity is also known as the $\bf True\ Negative\ Rate\ (TNR).$

 $^{^3}$ This formulation of Specificity is also known as the **Positive Predictive Value** (**PPV**) or **Precision**.

Summary Table

Measure	Name_1	Name_2	Calculation	As_Probability
Sensitivity Specificity Specificity	TNR	Recall 1 - FPR Precision	TN/(TN+FP)	$Pr(Y_hat = 1 Y = 1)$ $Pr(Y_hat = 0 Y = 0)$ $Pr(Y = 1 Y_hat = 1)$

```
TPR = True Positive Rate
TNR = True Negative Rate
PPV = Positive Predictive Value
FPR = False Positive Rate (Not covered in this lecture)
```

confusionMatrix()

The confusionMatrix function in the caret package computes all of the aforementioned metrics once we define what our algorithm should consider to be positive.

The function expects factors as inputs. The first factor level is considered to represent our "positive" outcome Y = 1. In our example, female is the first level.

The usage of confusionMatrix is as follows:

```
confusionMatrix(data = y_hat, reference = test_set$sex)
```

```
## Confusion Matrix and Statistics
##
##
             Reference
## Prediction Female Male
##
       Female
                  50
                       27
##
       Male
                  69 379
##
##
                  Accuracy: 0.817
##
                    95% CI: (0.781, 0.849)
##
       No Information Rate: 0.773
##
       P-Value [Acc > NIR] : 0.00835
##
##
                     Kappa: 0.404
##
##
   Mcnemar's Test P-Value : 2.86e-05
##
##
               Sensitivity: 0.4202
##
               Specificity: 0.9335
##
            Pos Pred Value: 0.6494
            Neg Pred Value: 0.8460
##
##
                Prevalence: 0.2267
            Detection Rate: 0.0952
##
##
      Detection Prevalence: 0.1467
##
         Balanced Accuracy: 0.6768
##
##
          'Positive' Class : Female
##
```