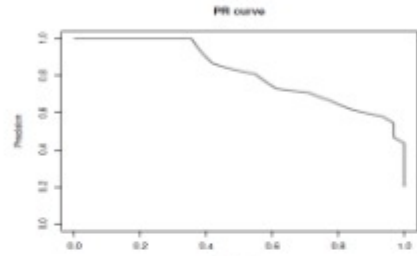


## Precision-recall curves are often used for imbalanced applications

- Precision ( $\frac{TP}{P}$ ) against the recall ( $\frac{TP}{TP+FN}$ ).
- To plot a precision-recall curve we can use the package PRRROC.

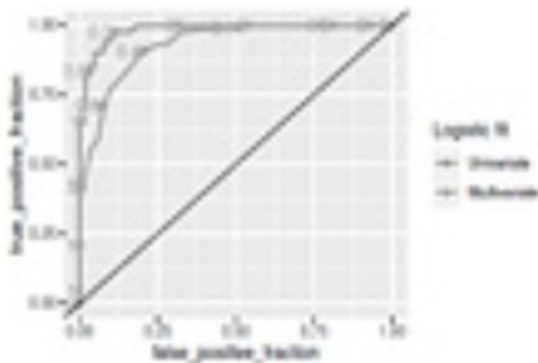
```
library(PRRROC)
```

```
PRROC_obj <- pr.curve(scores.class0 = imbalanced_heights$mu_hat,
                      weights.class0=imbalanced_heights$y,
                      curve=TRUE)
plot(PRRROC_obj, auc.main=FALSE, color=2)
```



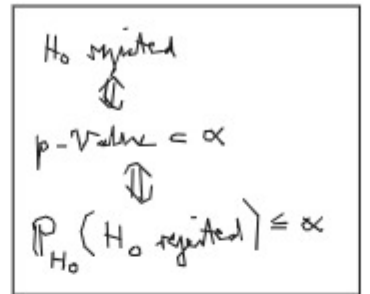
### Area under the ROC curve

- The <sup>AUC</sup> area under the ROC curve (AUC) captures the overall performance.
- AUC = 0.5 : random
- AUC = 1 : perfect



### Problem 13

We analyze 5000 independent experiments. In each experiment we sample the variable  $H_0$  (1 = 0, 2 = 1) with equal probability. We use a hypothesis test with a significance level of  $\alpha = 0.05$ . We reject  $H_0$  if the test statistic  $T$  is greater than the critical value  $c = 1.645$ . The test statistic  $T$  is the sum of the test statistics for each experiment. We reject  $H_0$  if the test statistic  $T$  is greater than the critical value  $c = 1.645$ . We reject  $H_0$  if the test statistic  $T$  is greater than the critical value  $c = 1.645$ .

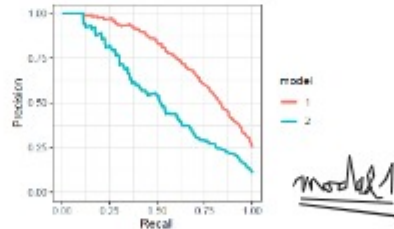


a)  $E[\text{false positive}] = 5000 \cdot 0.05 = 250 \checkmark$   
 number of cases where  $H_0$  is true      probability that  $H_0$  is rejected although  $H_0$  true

b)  $E[\text{false positive}] = 250 \checkmark$

### Problem 14 (1 credit)

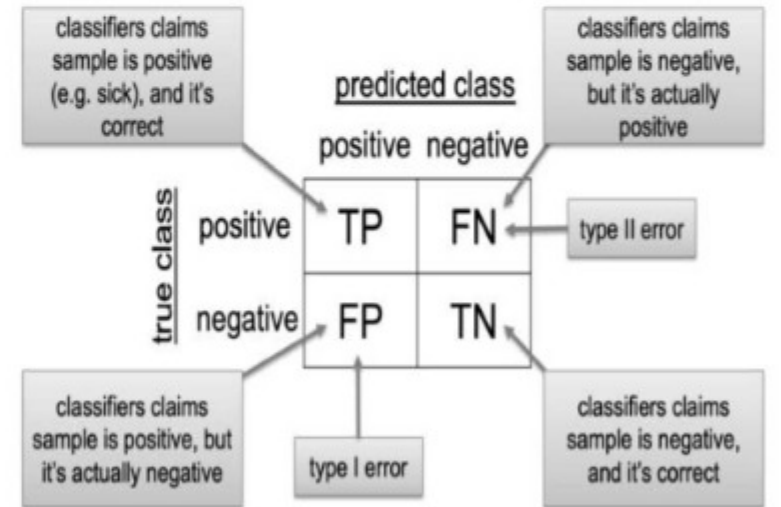
Consider the precision-recall curves provided below, which compare the performance of two models.



Based on the provided curves, is it possible to determine which of the two models has superior performance?

height 1,80 m (true positive)  
 height 1,9 m (true positive)  
 height 1,7 m (false negative)  
 You say, person is male if height > 1,75.

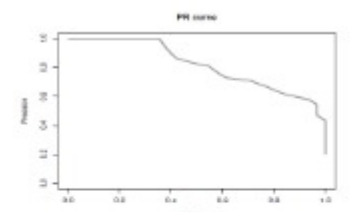
### Confusion Matrix



Precision-recall curves are often used for imbalanced applications

- Precision ( $\frac{TP}{P}$ ) against the recall ( $\frac{TP}{TP+FP}$ ).
- To plot a precision-recall curve we can use the package PRROC.

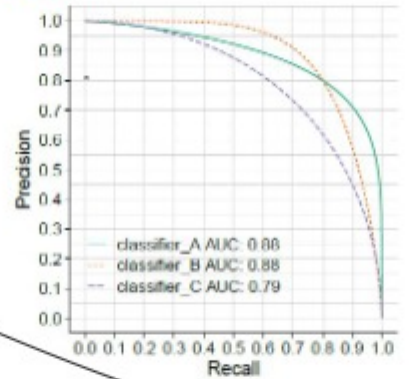
```
library(PRROC)
PRROC_obj <- pr.curve(scores.class0 = imbalanced_heights$mu_hat,
                      weights.class0=imbalanced_heights$y,
                      curve=TRUE)
plot(PRROC_obj, auc.main=FALSE, color=2)
```



positive =  
Kriminelle

Problem 13 (1 credit)

A police department uses the help of a machine learning classifier to find criminals. They want to ensure that less than 10% of individuals classified as criminal are actually innocent. Which one of the three classifiers A, B or C shown in the Precision-Recall curves do you recommend to use? Justify your choice.



gute Kurve = beste Wahl

da die grüne Kurve ab Precision von 0.8 den höchsten Recall hat  
Hoher Recall -> niedrigeres FP -> wenige unschuldige werden als Kriminelle verurteilt

Classification performance metrics

- The *sensitivity* refers to the fraction of actual positives that is predicted to be positive:  
Sensitivity =  $\frac{TP}{P} = \frac{TP}{TP+FN}$   
The sensitivity is also referred to as "recall", "true positive rate", or "power".
- The *specificity* refers to the fraction of actual negatives that is predicted to be negative:  
Specificity =  $\frac{TN}{N} = \frac{TN}{TN+FP}$   
The specificity is also known as "true negative rate" or "sensitivity of the negative class"
- The *precision* refers to the fraction of predicted positives that are indeed positives:

Recall = Precision =  $\frac{TP}{TP+FP}$

## Problem 3 (10 points)

Two particles, each of mass  $m$ , are moving with velocity  $v$  towards each other.



After the collision, the particles move with velocities  $v_1$  and  $v_2$ . The collision is elastic.

- (a) Find  $v_1$  and  $v_2$  in terms of  $v$ .
- (b) Find the change in kinetic energy.
- (c) Find the change in momentum.

*abhängig von  $v$ !*

The collision is elastic, so the total kinetic energy is conserved.



---

## Problem 7 (2 credits)

You have a dataset named `dt`, which includes information on cars from the `mtcars` dataset in R. The dataset contains information on miles per gallon (`mpg`) and transmission type (automatic or manual) of the cars.

<code>mpg</code>	<code>transmission</code>
21	Manual
21	Manual
23	Manual
21	Automatic
19	Automatic
18	Automatic
...	...

A colleague intends to investigate whether there is a significant difference in the median miles per gallon between cars with automatic and manual transmissions using a permutation test. The colleague visualizes a histogram of the test statistics across the permuted datasets alongside the value of the statistic `T` in the original unpermuted dataset (`T` reference). Examine the plot and the code used.

```
1 # Permutation function
2 perm_test <- function(data, n_permutations = 1000) {
3   # get reference statistics
4   T_ref <- data[transmission == "Manual", median(mpg)] -
5     data[transmission == "Automatic", median(mpg)]
6
7   # get permutation statistics
8   T_star <- sapply(1:n_permutations, function(x){
9     # shuffle groups
10    data[, transmission := sample(transmission), by='transmission']
11    # compute statistics
12    data[transmission == "Manual", median(mpg)] -
13      data[transmission == "Automatic", median(mpg)] })
14
15  g <- ggplot(data = data.table(T_star = T_star), aes(T_star)) +
16    geom_histogram() +
17    geom_vline(aes(xintercept=T_ref, color="T_ref")) + xlim(0, 10)
18  print(g)
19  return(list(T_ref=T_ref, T_star=T_star))
20 }
```

**Problem 7 (2 credits)**

You have a dataset named `dt`, which includes information on cars from the `mtcars` dataset in R. The dataset contains information on miles per gallon (`mpg`) and transmission type (automatic or manual) of the cars.

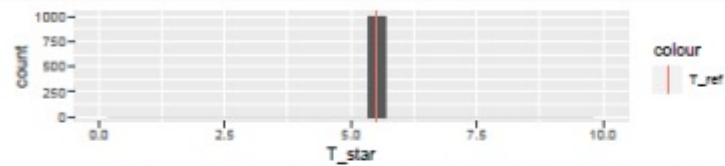
mpg	transmission
21	Manual
21	Manual
23	Manual
21	Automatic
19	Automatic
18	Automatic
...	...

Testgröße  $T_{obs}$  =  
 Durchschnittsverbrauch der  
 Gangschaltautos  
 • Durchschnittsverbrauch der  
 Automatikschaltautos  
 $H_0$ : Gangschaltautos verbrauchen weniger

A colleague intends to investigate whether there is a significant difference in the median miles per gallon between cars with automatic and manual transmissions using a permutation test. The colleague visualizes a histogram of the test statistics across the permuted datasets alongside the value of the statistic  $T$  in the original unpermuted dataset ( $T$  reference). Examine the plot and the code used.

```

1 # Permutation function
2 perm_test <- function(data, n_permutations = 1000) {
3   # get reference statistics
4   T_ref <- data[transmission == "Manual", median(mpg)] -
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17    geom_vline(aes(xintercept=T_ref, color="T_ref")) + xlim(0, 10)
18  print(g)
19  return(list(T_ref=T_ref, T_star=T_star))
20 }
    
```



- The code has a mistake. Clearly state the line of code that is wrong. Explain why it leads to the obtained histogram. Correct the line of code.
- Assume the code is now correct. Give the formula to compute the P-value estimate using  $T_{ref}$  and  $T_{star}$  in this particular case. No R code is required.

b)  $T_{ref} = T_{obs}$   
 $r = \#\{i \mid T_{star}[i] \geq T_{obs}\}$   
 $\hat{p} = \frac{r+1}{m+1}$   
 length(T\_star) // n-permutations

Falls  $T_{obs}$  falls  
 wird  $H_0$   
 abgelehnt  
 Falls  $T_{obs}$  groß, so  
 $r$  klein!  
 $\Downarrow \hat{p}$  klein  
 $\Downarrow H_0$  wird abgelehnt

**Permutation testing: P-value**

- Let
  - $m$  be the number of random (Monte Carlo) permutations
  - $r = \#\{T^* \geq T_{obs}\}$  be the number of these random permutations that produce a test statistic greater than or equal to that calculated for the actual data.
- The estimated one-sided P-value,  $\hat{P}$  is:

$$\hat{P} = \frac{r+1}{m+1}$$

- Permutation P-values should never be zero.

**Permutation testing: P-value**

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- Permutation P-values should never be zero.



Problem 3

Two screenshots of a data table. The first shows columns for 'Friend 1' and 'Friend 2' with rows of taste preferences. The second shows a similar table with a highlighted row.

$H_0$ : Friend 1 and 2 have independent music tastes

$T_{obs}$  = Anzahl der Beobachtungen, wo Freund 1 und 2 gleiche Tastes haben

$T_{obs} = 4$

```

# get permutation statistics
# stat = apply(X, MARGIN=2, FUN=function(x) {
#   shuffle_group
#   stat[, TRANSFORMED = apply(TRANSFORMED, MARGIN=2, FUN=function(x) {
#     compute_statistics
#     TRANSFORMED = TRANSFORMED[, ORDERING]
#   })
# })
    
```

$dt[, friend2 := sample(friend1)]$   
 $t\_stat[i] = dt[friend == friend2, .N]$

$T^*[i]$  = Wert der Teststatistik für die i-te Permutation

$r = 0;$   
 for  $i$  in  $1:N$ -perm {  
 if ( $T\_stat[i] \geq T_{ref}$ ) {  
 } }  
 $r = r + 1;$

$p = (r + 1) / (N\_perm + 1)$

Falls  $T_{obs}$  groß, dann wird  $H_0$  abgelehnt

$r = \# \{ T^* \geq T_{obs} \}$

it  $i$

$\hat{p} = \frac{r + 1}{m + 1}$  it  $i$

$H_0$  wird abgelehnt