

Statistical  
modelling and  
Learning vs  
Machine  
Learning

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Introduction  
Fundamental  
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Causal  
Modelling

Predictive  
Modelling

# Statistical modelling and Learning vs Machine Learning

‘Causal modelling vs Predictive modelling’  
‘Health Data Science Short Course’  
‘University of Kwazulu-Natal’

Innocent Maposa (PhD)  
Stellenbosch University  
Faculty of Medicine and Health Sciences  
Department of Global Health  
Division of Epidemiology & Biostatist



25 August 2024

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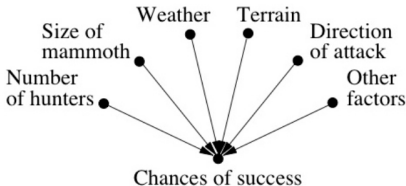
## Fundamental concepts

# The goal

- The purpose of statistics is to summarize data and quantify uncertainty around the *statistics*.
  - Descriptive
  - Inference including → Hypothesis testing, p-values, confidence intervals → Generalization
    - Bivariate
    - Regression (inference on the parameters)
- Predictive modelling mainly aims to find a *function* which can predict unseen outcomes based on new *feature* inputs with high **accuracy**.
- First we lay the foundational thoughts and philosophy in the *learning goals and processes*

# The Causal framework

- The *Hunter* → the mental model → the chances of success



*Figure 1: Why do we observe a success? : credit: Judea Pearl*

- The human mental models are always seeking to address this question: WHY?<sup>[ref-Pearl]</sup>
  - Causal modelling and inference is all about taking this question seriously
  - Understanding the mechanism of occurrence led to human progress over centuries!

# Causal Framework

## ■ The ladder of causation

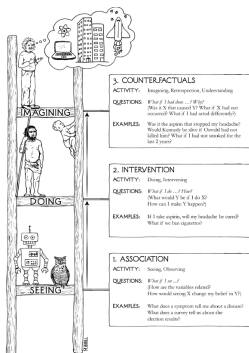


Figure 2: Three levels of causation? : credit: Judea Pearl

# Association level

## 1. ASSOCIATION

ACTIVITY: Seeing, Observing

QUESTIONS: *What if I see ...?*  
(How are the variables related?)  
How would seeing X change my belief in Y?)

EXAMPLES: What does a symptom tell me about a disease?  
What does a survey tell us about the election results?

Figure 3: Three levels of causation? : credit: Judea Pearl

- *How would seeing X change my belief in Y?*
- Can rephrase to: *How would seeing X influence my understanding of Y?*
  - By observing X, can I say something about unobserved Y?
    - Under what circumstances (assumptions)?
- Challenges with this level of evidence includes *BIAS* - confounding, selection, mediation, moderation,

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Correlation vs. causation

This keeps happening. How heavy  
are cats?



Figure 4: Correlation challenges

- Causation as a limit for correlation (*causation always implies correlation*) - Pearson



# Intervention Level

## 2. INTERVENTION

**ACTIVITY:** Doing, Intervening

**QUESTIONS:** *What if I do ...? How?*  
(What would Y be if I do X?)  
How can I make Y happen?)

**EXAMPLES:** If I take aspirin, will my headache be cured?  
What if we ban cigarettes?

Figure 5: Three levels of causation? : credit: Judea Pearl

- *How can I make Y happen?*
  - In other words, can I do something to influence the outcome of interest?
- Study design elements are optimized to *minimize (eliminate) bias*

# Counterfactual

## 3. COUNTERFACTUALS

**ACTIVITY:** Imagining, Retrospection, Understanding

**QUESTIONS:** *What if I had done ...? Why?*  
(Was it X that caused Y? What if X had not occurred? What if I had acted differently?)

**EXAMPLES:** Was it the aspirin that stopped my headache?  
Would Kennedy be alive if Oswald had not killed him? What if I had not smoked for the last 2 years?

Figure 6: Three levels of causation? : credit: Judea Pearl

- *Was it X that caused Y? What if X had not happened? What if I had acted differently?*
  - These are high level questions that cannot be answered by just seeing and observing.
  - Most statistical paradigms that rely on learning from data are limited at this level

# Law of Regression

- The average regression of the offspring to a constant fraction of their respective mid-parental deviations, which was first observed in the diameters of seeds, and then confirmed by observations on human stature, is now shown to be a perfectly reasonable law which might have been deductively foreseen<sup>[ref-Galton]</sup>
  - The introduction of *regression* as a principle that can help us understand relationship
    - be they causal or *associational*
- There are two goals in analysing the data:
  - Explanation (Information): To extract some information about how nature is associating (relating) the response variables to the input variables
  - Prediction: To be able to predict what the responses are going to be to future input values<sup>[ref-Breiman]</sup>

# What is regression?

- The goal of regression is to model the relationship between the response (outcome or target) variable  $Y$  and predictor(s) variable(s)  $X$  using the form

$$Y = f(X) + \epsilon$$

- where the function  $f$  describes the functional form of the relationship between variables and  $\epsilon$  accounts for error. This relationship can qualitatively be thought of in different ways:
  - response = deterministic + random
  - response = signal + noise
  - response = model + unexplained
  - response = prediction + error
- Linear and generalized linear models make strong assumptions about the data generating process ie the structure of this model and restricts  $f(X)$  to linear functions of  $X$  ie  $Y = X\beta + \epsilon$ .

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# Causal Modelling

# Classical Statistical Modelling

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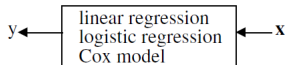
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- Classical statistical modelling refers to practices aiming to conduct model validation, and thus, statistical inference on one or several quantities of interest eg distributions, model parameters, errors etc.
- With inference, the goal is to estimate  $\hat{\beta}$  that estimates the true quantity  $\beta$ 
  - This true quantity is assumed to exist independently of the statistical model<sup>[ref-Daoud2023Statistical]</sup>
- These models are aimed at explaining relationships between variables as main focus, prediction is of little interest
  - *Fundamental to scientific enquiry*

# Classical Statistical Modelling

- Scientific methods consist of cycles of deductively formulating a hypothesis from substantive theory, testing this hypothesis in a model and against the data, and then revising the theory based on empirical results.
- The requirement of testing substantive theories through an interpretable statistical model is one of the appeals for classical statistical modelling<sup>[ref-Daoud2023Statistical]</sup>.



*Figure 7: Statistical Causal Modelling : credit: L.Breiman*

- Model validation: generally uses some form of goodness-of-fit tests and residual examination.

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# Predictive Modelling



# Predictive modelling

- Sometimes referred to as *algorithmic modelling*, entails practices defining a procedure  $f$ , that generates accurate predictions,  $\hat{\mathbf{Y}}$ , about an event (outcome),  $\mathbf{Y}$ .
  - by accurate, we mean, predictions that are as similar as possible to the true event that  $f$  has not yet encountered.
- A procedure is an *algorithm*, or a *function*, that takes some input  $\mathbf{X} = x$ , operates on this input  $f(\phi(x))$ , and then produces  $f(x) = \hat{y}$  where  $\phi(x)$  are features derived from  $\mathbf{X}$  and may include polynomials, interactions, etc.
  - Kernels
- The main goal is prediction and optimizing prediction function is key!
  - modern machine learning methods heavily rely on expanding the feature space in order to improve predictive accuracy

# Predictive modelling

- Under predictive modelling framework and related assumptions, the relationship between  $X$  and  $Y$  may or may not be causal.
- The overarching goal is to develop a model  $f$  that operates on data inputs, producing the best possible predictions  $\hat{Y}$  of  $Y$  that  $f$  has not observed yet.
- Absence of causal reasoning is a major limitation - however, according to Pearson, *causation is "sorely the conceptual limit to correlation or association"*.

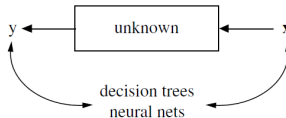


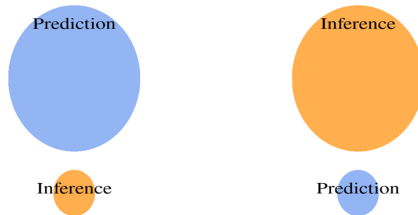
Figure 8: Predictive Modelling : credit: L.Breiman

- Model validation. Measured by predictive accuracy

# Comparisons of the frameworks

Machine Learning

Statistics



*Figure 9: Model framework goals*

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# Comparisons

	Data modeling culture (DMC)	Algorithmic modeling culture (AMC)
<b>Exemplifying question</b>	What is the causal relationship between food supply and famines?	How well can famines be predicted from available data?
<b>Goal</b>	Estimating unbiased parameters for causal estimation, to populate the magnitudes of the edges of a directed acyclic graph (DAG).	To develop and train an algorithm $f$ for accurate prediction.
<b>A key assumption</b>	Assuming a DAG, a stipulated and interpretable statistical model such as $y_i = c_0 + \beta w_i + e_i$ produces unbiased estimates of the true causal quantity $\beta$ .	The algorithm $f$ can produce accurate predictions of $Y$ from data source, $D$ .
<b>Limitation</b>	Although the parametric model is interpretable, its statistical structure may be a poor representation of the causal system.	Although $f$ produces accurate predictions, the model is a black-box restricting causal interpretations.
<b>Quantity of interest</b>	$\hat{\beta}$	$\hat{Y}$

*Figure 10: Central practices of two statistical cultures: credit: L. Breiman*

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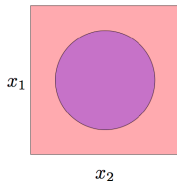
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# Optimization functions

- For statistical causal models
  - *Loss function + penalty*
  - $L + \lambda \sum_{j=1}^p \beta_j^2$  where  $L$  is the log loss function for generalized linear models and  $\lambda$  parameter controls how much emphasis is given to the penalty term. The higher the  $\lambda$  value, the more coefficients in the regression will be pushed towards zero.
- Generally, we optimize the function based on the observed variables
- For predictive models, we optimize *featurised or kernelized loss functions*
  - high dimensional
    - number of features
    - interactions etc

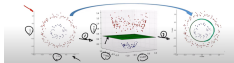
# Two class example



*Figure 11: Two class geometric problem: credit: D Rosenberg, NYU*

- With linear feature map  $\phi(X) = (X_1, X_2)$  and linear models, no hope to separate the classes
- With appropriate nonlinearity  $\phi(X) = (X_1, X_2, X_1^2 + X_2^2)$ , simple.
- Example Video

# Decision boundary in higher dimension



*Figure 12: Two class geometric problem:credit:D Rosenberg, NYU*

- The kernel trick optimizes expressiveness and hence prediction accuracy
- A kernel  $\phi(X_i, X_j)$  is a function that quantifies the similarities between observations by summarizing the relationship between every single pairs in the training set.

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# Examples



# Data

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```
## # A tibble: 5 x 8
```

```
##   L_SEP L_ethnicity cancer hba1c sample id BME deprived
##   <dbl>    <dbl>    <dbl> <dbl>    <dbl> <dbl> <dbl>    <dbl>
## 1  1.11    -0.218      0  9.31      1     1     0      5
## 2 -0.206     2.29      1 10.7      1     2     1      3
## 3  1.22    -0.0640     1 11.1      1     3     0      5
## 4  0.0993   -0.692      1  9.68      1     4     0      3
## 5  1.51     -1.38      0  9.30      1     5     0      5
```

```
##   Variable      N   Mean Std. Dev.   Min Pctl. 25 Pctl. 75   Max
## 1   L_SEP    2500 -0.012      1   -3.7   -0.7    0.7   3.5
## 2 L_ethnicity 2500 -0.024      1   -3.4   -0.7    0.7   3.1
## 3   cancer    2500   0.25    0.43    0      0      0    1
## 4   hba1c    2500    9      1.5    3.7    7.9    9.9   15
## 5   sample    2500   0.75    0.43    0      1      1    1
## 6    id      2500 1250    722    1    626   1875  2500
## 7    BME     2500   0.25    0.43    0      0      1    1
## 8  deprived  2500    3      1.4    1      2      4    5
```

# Describe data

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Characteristic	N = 2,500 <sup>1</sup>
L_SEP	-0.01 (-0.69, 0.70)
L_ethnicity	-0.01 (-0.74, 0.68)
cancer	613 (25%)
hba1c	8.95 (7.95, 9.93)
BME	629 (25%)
deprived	
1	509 (20%)
2	512 (20%)
3	479 (19%)
4	488 (20%)
5	512 (20%)

<sup>1</sup>Median (Q1, Q3); n (%)

# Describe data visualization

Ethnicity vs hba1c



- Seems the separation problem here may be difficult

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# The DAG

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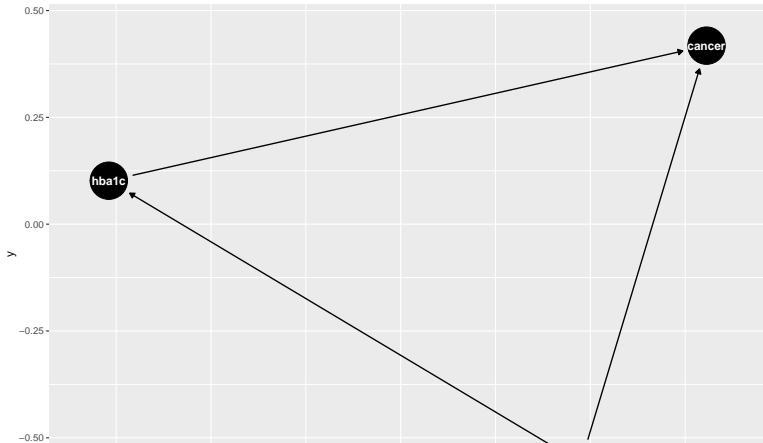
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```
##  
## Attaching package: 'ggdag'  
  
## The following object is masked from 'package:stats':  
##  
##   filter
```



# Causal statistical model

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- Question: What is the effect of hba1c on cancer?
- Unadjusted effect

Characteristic	OR <sup>1</sup>	95% CI <sup>1</sup>	p-value
hba1c	1.34	1.26, 1.43	<0.001

<sup>1</sup>OR = Odds Ratio, CI = Confidence Interval

# Logistic regression (adjusted effect)

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Characteristic	OR <sup>1</sup>	95% CI <sup>1</sup>	p-value
hba1c	1.06	0.98, 1.14	0.13
as.factor(BME)			
0	—	—	
1	1.42	1.03, 1.97	0.032
deprived	1.63	1.50, 1.77	<0.001
L_ethnicity	1.37	1.17, 1.60	<0.001

<sup>1</sup>OR = Odds Ratio, CI = Confidence Interval

# Within sample predict

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##	Variable	N	Mean	Std. Dev.	Min	Pctl. 25	Pctl. 75	Max
## 1	L_SEP	2500	-0.012	1	-3.7	-0.7	0.7	3.5
## 2	L_ethnicity	2500	-0.024	1	-3.4	-0.7	0.7	3.1
## 3	cancer	2500	0.25	0.43	0	0	0	1
## 4	hba1c	2500	9	1.5	3.7	7.9	9.9	15
## 5	sample	2500	0.75	0.43	0	1	1	1
## 6	id	2500	1250	722	1	626	1875	2500
## 7	BME	2500	0.25	0.43	0	0	1	1
## 8	deprived	2500	3	1.4	1	2	4	5
## 9	cancer_prob	2500	0.25	0.16	0	0.1	0.3	0.8
## 10	c_pred	2500	0.079	0.27	0	0	0	1

# Confusion Table

```
##
##
##      Cell Contents
## |-----|
## |              N |
## | Chi-square contribution |
## |      N / Row Total |
## |      N / Col Total |
## |      N / Table Total |
## |-----|
##
##
## Total Observations in Table:  2500
##
##
##                | popdatex$cancer
## popdatex$c_pred |          0 |          1 | Row Total |
## -----|-----|-----|-----|
##                0 |      1806 |      496 |      2302 |
##                |      2.697 |      8.301 |           |
##                |      0.785 |      0.215 |      0.921 |
##                |      0.957 |      0.809 |           |
##                |      0.722 |      0.198 |           |
## -----|-----|-----|-----|
##                1 |        81 |       117 |       198 |
##                |     31.351 |     96.509 |           |
##                |      0.409 |      0.591 |      0.079 |
##                |      0.043 |      0.191 |           |
##                |      0.032 |      0.047 |           |
## -----|-----|-----|-----|
##      Column Total |      1887 |       613 |      2500 |
##                |      0.755 |      0.245 |           |
```

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# Predictive modelling

# Predictive Model

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```
# Encoding the target feature as factor and splitting
suppressWarnings(suppressMessages(library(caTools)))
dcancer<-popdatex %>% select(c(cancer, hba1c, BME, de
dcancer$cancer = factor(dcancer$cancer, levels = c(0,
set.seed(123)
splitdat = sample.split(dcancer$cancer, SplitRatio =

train = subset(dcancer, splitdat == TRUE)
test = subset(dcancer, splitdat == FALSE)
```

# Logistic regression (try prediction out of sample)

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```
##
##
##      Cell Contents
## |-----|
## |                      N |
## | Chi-square contribution |
## |      N / Row Total |
## |      N / Col Total |
## |      N / Table Total |
## |-----|
##
##
## Total Observations in Table:  750
##
##
##      | test$cancer
## test$c_pred |      0 |      1 | Row Total |
## -----|-----|-----|-----|
##           0 |      547 |      155 |      702 |
##           |      0.560 |      1.723 |      |
##           |      0.779 |      0.221 |      0.936 |
##           |      0.966 |      0.842 |      |
##           |      0.729 |      0.207 |      |
## -----|-----|-----|-----|
##           1 |       19 |       29 |       48 |
##           |      8.190 |     25.192 |      |
##           |      0.396 |      0.604 |      0.064 |
##           |      0.034 |      0.158 |      |
##           |      0.025 |      0.039 |      |
## -----|-----|-----|-----|
## Column Total |      566 |      184 |      750 |
##           |      0.755 |      0.245 |      |
```

# Support vector machine

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```
##
```

```
## Call:
```

```
## svm(formula = cancer ~ ., data = train, type = "C-
```

```
##       kernel = "linear", gamma = 1)
```

```
##
```

```
##
```

```
## Parameters:
```

```
##       SVM-Type:  C-classification
```

```
##       SVM-Kernel: linear
```

```
##           cost:  1
```

```
##
```

```
## Number of Support Vectors:  884
```

# SVM linear

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```
##
##
##      Cell Contents
## |-----|
## |                               N |
## |      N / Table Total |
## |-----|
##
##
## Total Observations in Table:  750
##
##
##      test$cancer | y_pred
## |-----|-----|-----|
## |      0 |      566 |      566 |
## |      |      0.755 |      |
## |-----|-----|-----|
## |      1 |      184 |      184 |
## |      |      0.245 |      |
## |-----|-----|-----|
## Column Total |      750 |      750 |
## |-----|-----|-----|
##
##
```

# SVM radial

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```
##
##
##      Cell Contents
## |-----|
## |                                N |
## | Chi-square contribution |
## |           N / Row Total |
## |           N / Col Total |
## |           N / Table Total |
## |-----|
##
##
## Total Observations in Table:  750
##
##
##                                | y_pred2
```

# SVM polynomial

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```
##
##
##      Cell Contents
## |-----|
## |              N |
## | Chi-square contribution |
## |      N / Row Total |
## |      N / Col Total |
## |      N / Table Total |
## |-----|
##
##
## Total Observations in Table:  750
##
##
##              | y_pred3
## test$cancer | 0 |          1 | Row Total |
## -----|-----|-----|-----|
##              0 |    548 |         18 |         566 |
##              | 0.391 |        6.435 |         |
##              | 0.968 |         0.032 |        0.755 |
##              | 0.775 |         0.419 |         |
##              | 0.731 |         0.024 |         |
## -----|-----|-----|-----|
##              1 |    159 |         25 |         184 |
##              | 1.204 |        19.795 |         |
##              | 0.864 |         0.136 |        0.245 |
##              | 0.225 |         0.581 |         |
##              | 0.212 |         0.033 |         |
## -----|-----|-----|-----|
## Column Total |    707 |         43 |         750 |
##              | 0.942 |        0.057 |         |
```

# Random forest

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```
## randomForest 4.7-1.1

## Type rfNews() to see new features/changes/bug fixes.

##
## Attaching package: 'randomForest'

## The following object is masked from 'package:ggplot2':
##
##     margin

##
## Call:
## randomForest(formula = cancer ~ ., data = train, proximity = TRUE)
##           Type of random forest: classification
##           Number of trees: 500
## No. of variables tried at each split: 2
##
##           OOB estimate of  error rate: 23.43%
## Confusion matrix:
##           0  1 class.error
## 0 1244 77  0.05828917
## 1  333 96  0.77622378
```



# Random forest train performance

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```
## Confusion Matrix and Statistics
##
##           Reference
## Prediction    0    1
##           0 1303  147
##           1   18  282
##
##           Accuracy : 0.9057
##           95% CI : (0.891, 0.919)
##           No Information Rate : 0.7549
##           P-Value [Acc > NIR] : < 2.2e-16
##
##           Kappa : 0.7165
##
##  Mcnemar's Test P-Value : < 2.2e-16
##
##           Sensitivity : 0.9864
##           Specificity : 0.6573
##           Pos Pred Value : 0.8986
##           Neg Pred Value : 0.9400
##           Prevalence : 0.7549
##           Detection Rate : 0.7446
##           Detection Prevalence : 0.8286
##           Balanced Accuracy : 0.8219
##
##           'Positive' Class : 0
##
```

# Random forest test performance

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```
## Confusion Matrix and Statistics
##
##           Reference
## Prediction  0    1
##           0 525 144
##           1  41  40
##
##           Accuracy : 0.7533
##           95% CI : (0.7209, 0.7838)
##           No Information Rate : 0.7547
##           P-Value [Acc > NIR] : 0.5534
##
##           Kappa : 0.1787
##
## Mcnemar's Test P-Value : 6.421e-14
##
##           Sensitivity : 0.9276
##           Specificity : 0.2174
##           Pos Pred Value : 0.7848
##           Neg Pred Value : 0.4938
##           Prevalence : 0.7547
##           Detection Rate : 0.7000
##           Detection Prevalence : 0.8920
##           Balanced Accuracy : 0.5725
##
##           'Positive' Class : 0
##
```

# References I

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- 3 Breiman L. Statistical Modeling: The Two Cultures (with comments and a rejoinder by the author). *Statistical Science* 2001; **16**: 199–231.
- 4 Daoud A, Dubhashi D. Statistical Modeling: The Three Cultures. *Harvard Data Science Review* 2023; **5**.

# References II

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