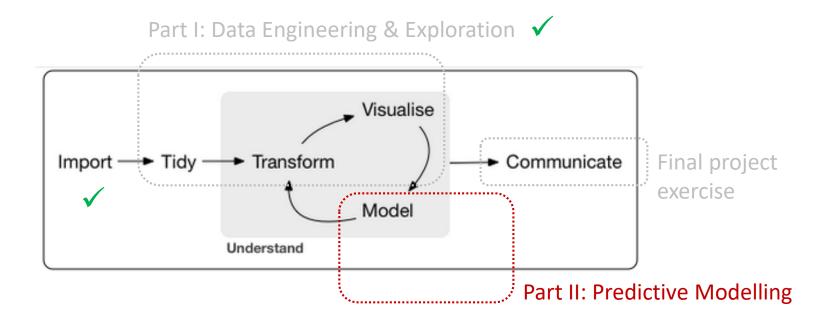


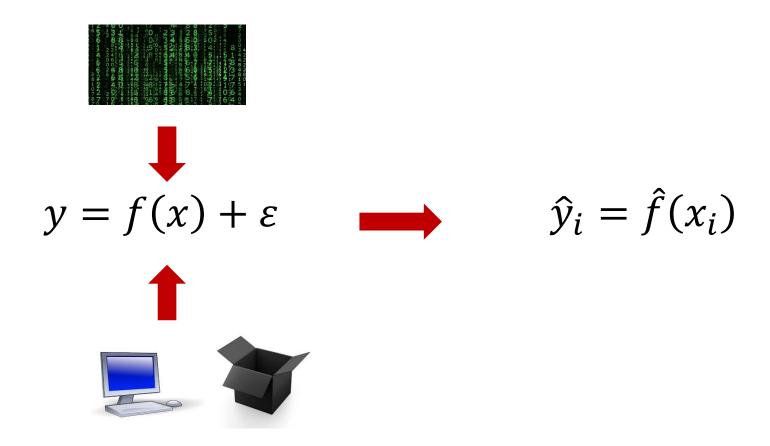
# Data Science Predictive Modelling

Gero Szepannek

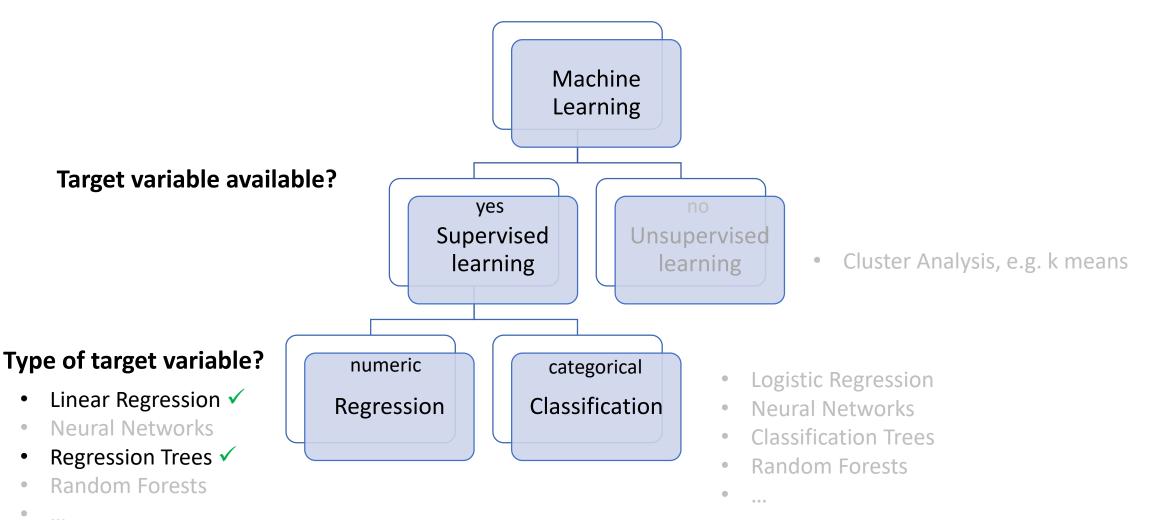
#### Data Science Process



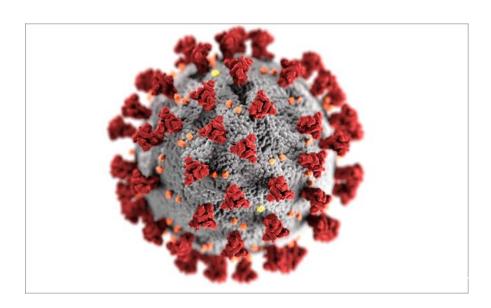
#### **Prediction Model**



### Regression vs. Classification

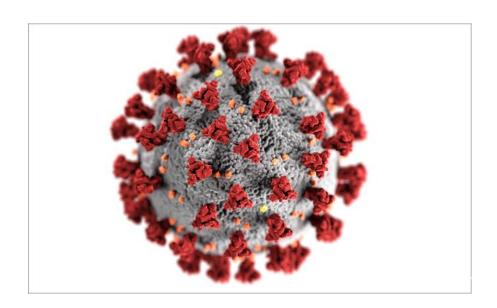
















### A Point-and-click Framework developed @

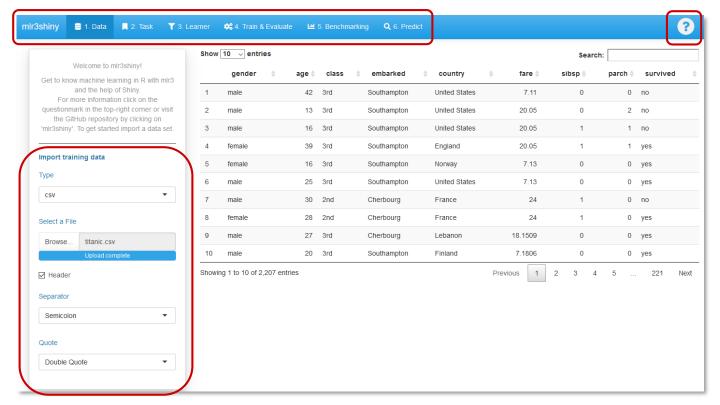


# install

remotes::install\_github("https://github.com/LamaTe/mlr3shiny.git")

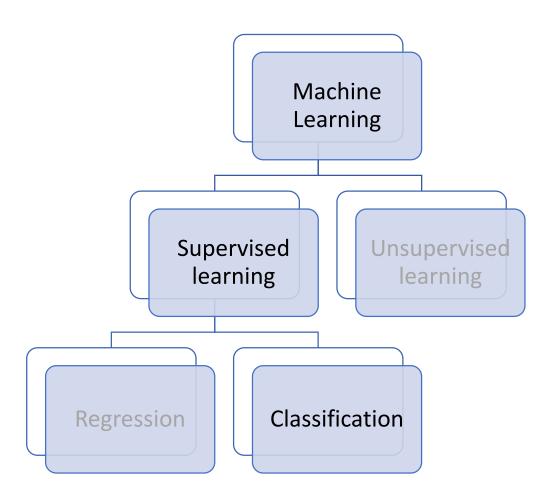
# start
library(mlr3shiny)
launchMlr3Shiny()





### Classification

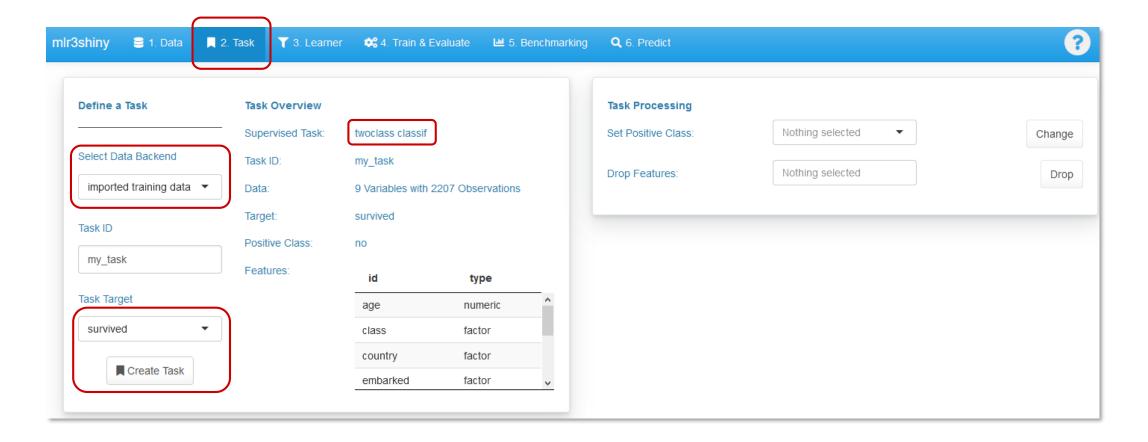
 $\hat{y}_i$ 



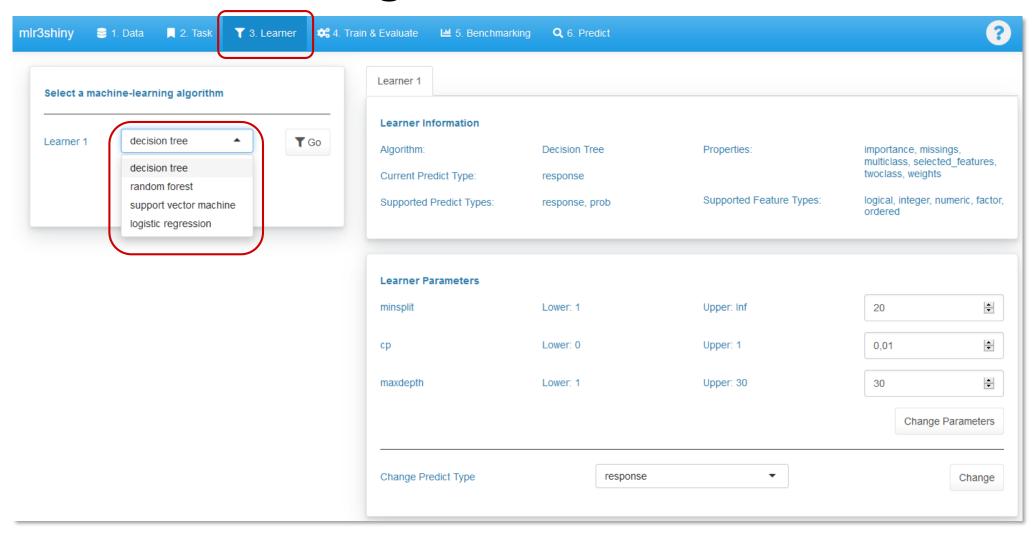
$$\hat{y}_i = \hat{f}(x_i)$$

- 1. In classification  $y_i$  is categorical.
- 2. ...most relevant practical case: binary classification  $\rightarrow$   $y_i \in \{0,1\}$

### Classification



## Classification Algorithms



### Flavours of Classification Models

 $\hat{y}_i$ 

$$\hat{y}_i = \hat{f}(x_i)$$

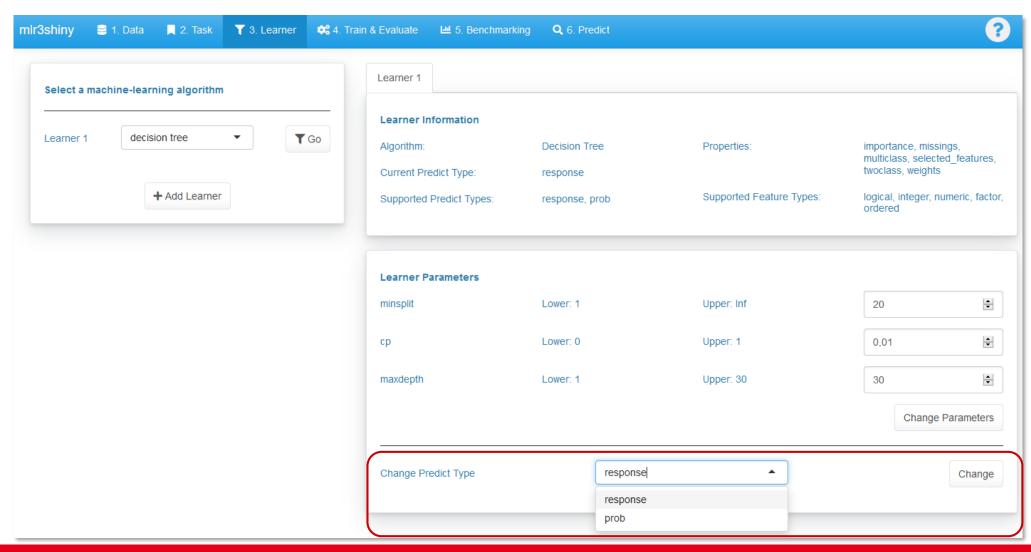
- 1. In classification  $y_i$  is categorical.
- 2. ...most relevant practical case: binary classification  $\rightarrow y_i \in \{0,1\}$
- 3. Predicting class labels:  $\hat{y}_i \in \{0,1\}$
- 4. Predicting posterior probabilities:  $\hat{P}(Y_i = 1) \in [0,1]$



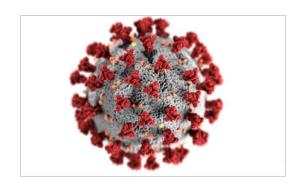
What is a natural rule for prediction of class labels in order to minimize the probability of making an error?

$$\hat{y}_i = \begin{cases} 1 & \dots if \ \hat{P}(Y_i = 1) \ge cut \ off \\ 0 & \dots if \ \hat{P}(Y_i = 1) < cut \ off \end{cases}$$

## Classification Algorithms



#### Performance Measures for Classification



**Epidemiologisches Bulletin** 

8 2021

25. Februar 2021 (online vorab)

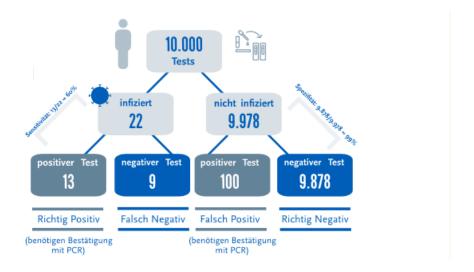
Was ist bei Antigentests zur Eigenanwendung (Selbsttests) zum Nachweis von SARS-CoV-2 zu beachten?

- Accuracy
- Sensitivity
- Specifity
- Precision

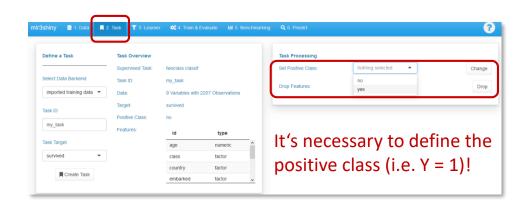
Test detects infection.

$$P(\hat{Y} = 1 | Y = 1)$$

$$P(\hat{Y}=0|Y=0)$$



https://edoc.rki.de/bitstream/handle/176904/7826/EB-8-21-Heimtest-Beitrag\_22-02-21-zur%20Ver%C3%B6ffentlichung.pdf?sequence=1&isAllowed=y



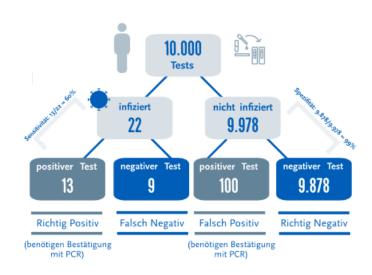
- 1. Accuracy
- 2. Sensitivity
- 3. Specifity
- 4. Precision

Test detects infection.

$$P(\hat{Y}=1|Y=1)$$

$$P(\hat{Y}=0|Y=0)$$

	$\hat{Y} = 1$	$\hat{Y} = 0$	
Y = 1			
Y = 0			



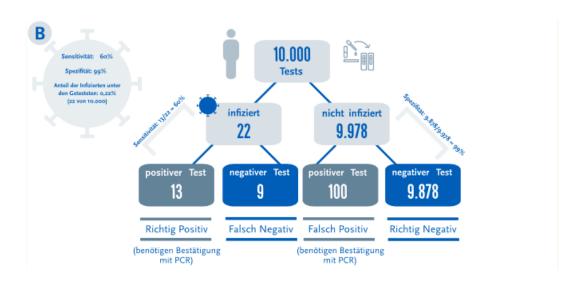
	$\hat{Y} = 1$	$\hat{Y} = 0$	
Y = 1			
Y = 0			

- 1. Accuracy
- 2. Sensitivity
- 3. Specifity
- 4. Precision

Test detects infection.

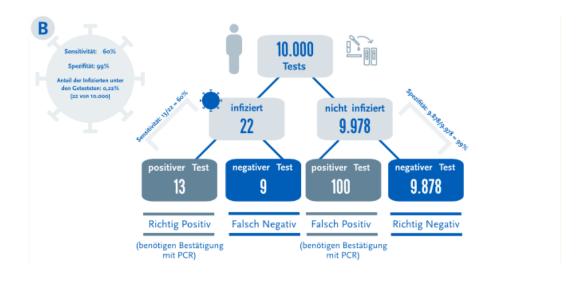
$$P(\hat{Y}=1|Y=1)$$

$$P(\widehat{Y}=0|Y=0)$$



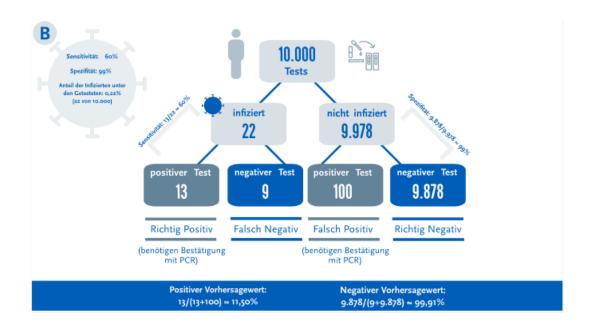
	$\hat{Y} = 1$	$\hat{Y} = 0$	
Y = 1			
Y = 0			

- 1. Accuracy
- 2. Sensitivity
- 3. Specifity
- 4. Precision



	$\hat{Y} = 1$	$\hat{Y} = 0$	
Y = 1			
Y = 0			

- 1. Accuracy
- 2. Sensitivity
- 3. Specifity
- 4. Precision / PPV

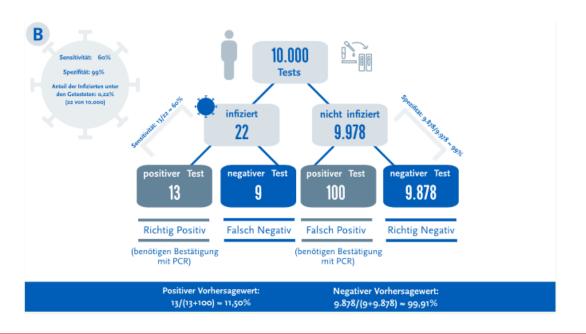


	$\hat{Y} = 1$	$\hat{Y} = 0$	
Y = 1			
Y = 0			



- 1. Imagine: How can we change the cut off in order to increase the sensitivity of the test?
- 2. ...What will typically happen with the precision (aka PPV) at the same time?

$$\hat{y}_i = \begin{cases} 1 & \dots if \ \hat{P}(Y_i = 1) \ge cut \ off \\ 0 & \dots if \ \hat{P}(Y_i = 1) < cut \ off \end{cases}$$



## Takeaway...

#### Bei Veranstaltungen nicht auf Schnelltests verlassen!

Deshalb sei es gefährlich, sich bei Einlasskontrollen auf das Ergebnis eines Schnelltests zu verlassen - etwa beim Theater- oder Konzert-Besuch, an der Eingangstür eines Restaurants. Wenn ein Schnelltest eine Infektion "übersieht", wird diese Person herumlaufen in der Annahme, dass sie nicht ansteckend ist - und kann so mitunter andere infizieren. "Es ist nicht so simpel, wie es in der Politik dargestellt wird - nach dem Motto: Jetzt kann alles öffnen, weil wir ja die Schnelltests haben." Zwischen 40 Prozent und 60 Prozent der Infektionen werden bei Schnelltests übersehen, so Drosten.

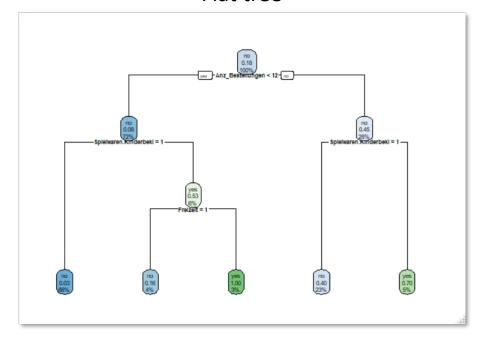
#### An Schulen machen Antigen-Schnelltests Sinn

Drosten stellt aber auch klar: In Schulen sei der Einsatz von Antigen-Schnelltests trotzdem gerechtfertigt - wenn die Schülerinnen und Schüler mindestens zweimal in der Woche getestet werden. "Selbst wenn bei einer Testung nicht alle Infektionen entdeckt werden, bei der nächsten Testung nach zwei oder drei Tagen werden die Infektionen dann nachgewiesen. In Clustern ist solch ein geringer zeitverzögerter Effekt kein Problem", meint der Virologe. Wichtig sei nur, Infektionen in einem Cluster aufzuspüren, um so die Kontrolle in den Schulen während der Pandemie zu behalten und dann entsprechend schnell mit Cluster-Quarantäne zu reagieren.

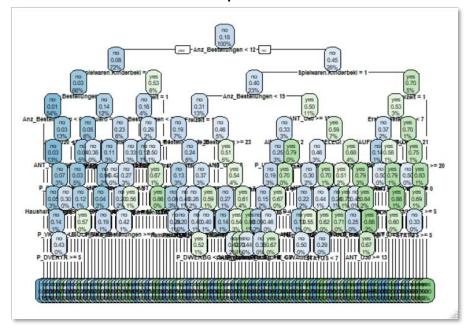
## Optimizing a Decision Tree



"Flat tree"



"Deep tree"



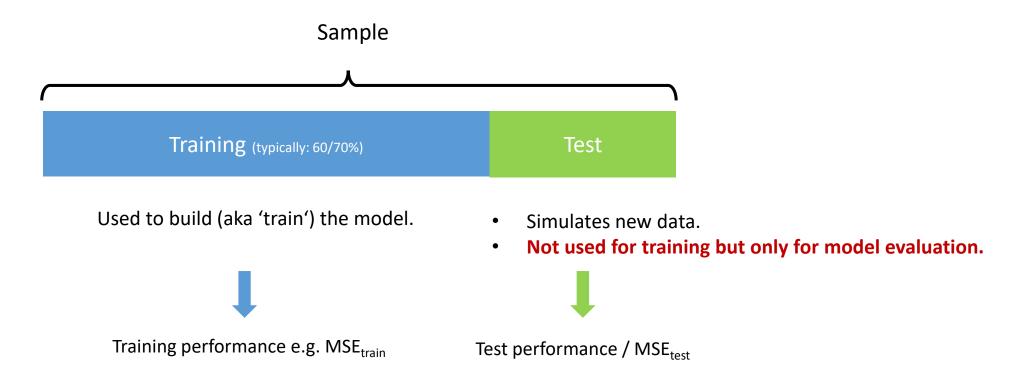


1. What might be the problem with a flat tree (w.r.t. accuracy)?

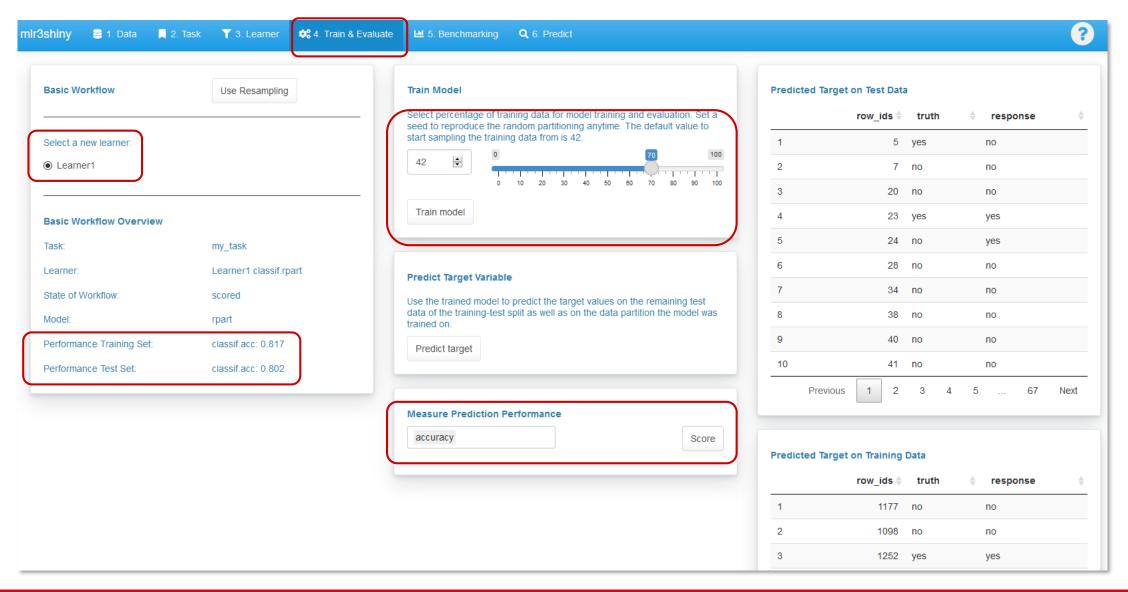
2. ...What might be the problem with a deep tree?

#### Test Data

- A model should have strong predictive power also for new data!
- In practice: no new data is available to test the model ⊗.
- Therfore use a trick: (Randomly) split the data in two parts:



### Performance Evaluation



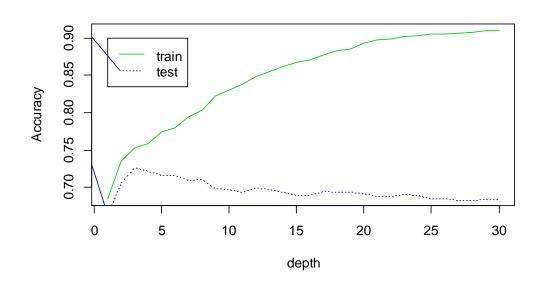


#### On your own:

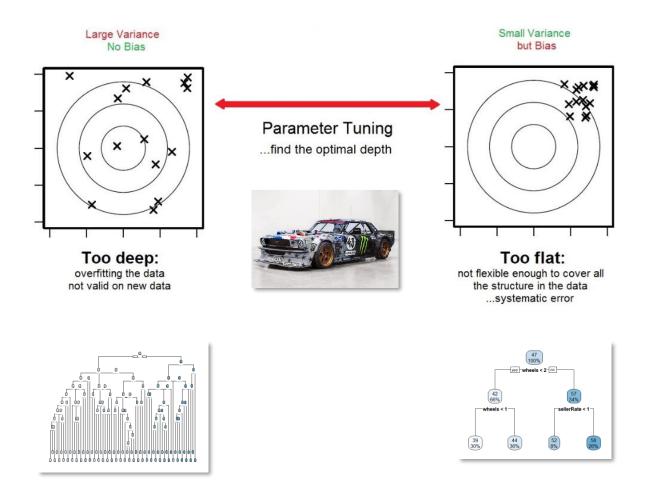
- 1. Modify the hyperparameters of the tree...
- 2. ...Try to find the best model, i.e. the model that maximizes accuracy!
- 3. What do you observe w.r.t. the accuracy on both:
  - 1. Training data...
  - 2. ...and Test data?



# Overfitting and Underfitting

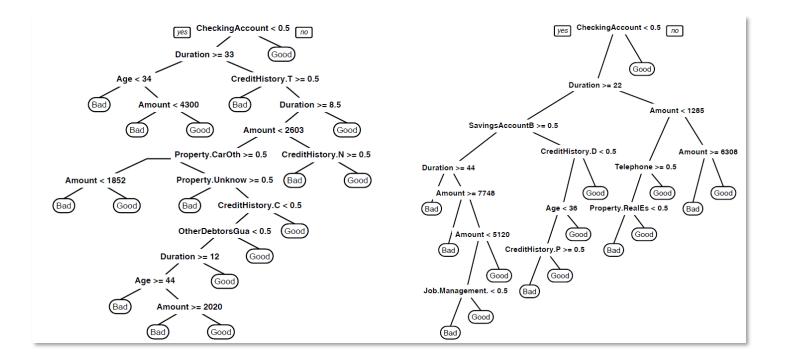


...e.g. set the parameters minsplit = minbucket = 1 and cp = 0 such that and then increase maxdepth.



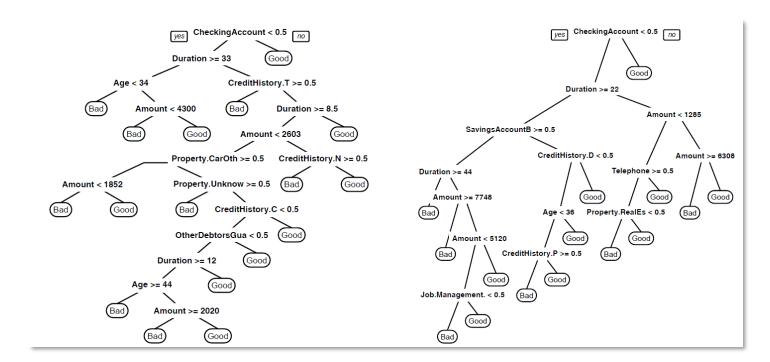
#### Variance of Models

- Deep trees will look pretty different on different samples...
- ...except for the first splits.
- Try to explain, why?



#### Variance of Models

- Deep trees will look pretty different on different samples...
- ...except for the first splits.
- Try to explain, why?



...from Statistics, we know:

$$Var(\bar{X}) = \frac{Var(X)}{n}$$

- 1. What denotes  $\bar{X}$  in Statistics?
- 2. ...How can we translate this to the variance of a single model?



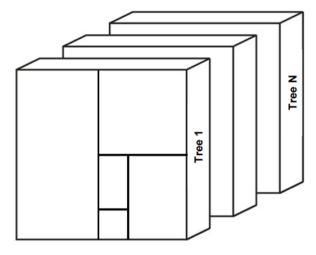
## Random Forests and Bootstrapping



- A large number of random samples of size n with replacement is drawn.
- For each sample a (deep) tree is built.
- The final forest model counts the number of predictions of each class for an observation.
- The class with the most predcitions is assigned.
- This process of aggregating predictions is called "voting".

Stichprobe	1	L 2	2 3	3 4	1 5	6
Bootstrap-Sample 1	6	5 (	5 2	2 5	5 4	4
Bootstrap-Sample 2	5	5 1	L 4	4 5	5 3	5
Bootstrap-Sample 3	6	5 2	2 3	3 6	5 6	1
Bootstrap-Sample 4	3	3 4	1 (	5 1	. 6	6

#### **Ensemble of trees**





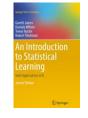
# ...From Bootstrapping to Random Forests<sup>1</sup>

- In addition to Bootstrapping random forests make use of a 2<sup>nd</sup> trick:
- For each tree at each split only a subset of variables is randomly offered for split search.

#### • Idea:

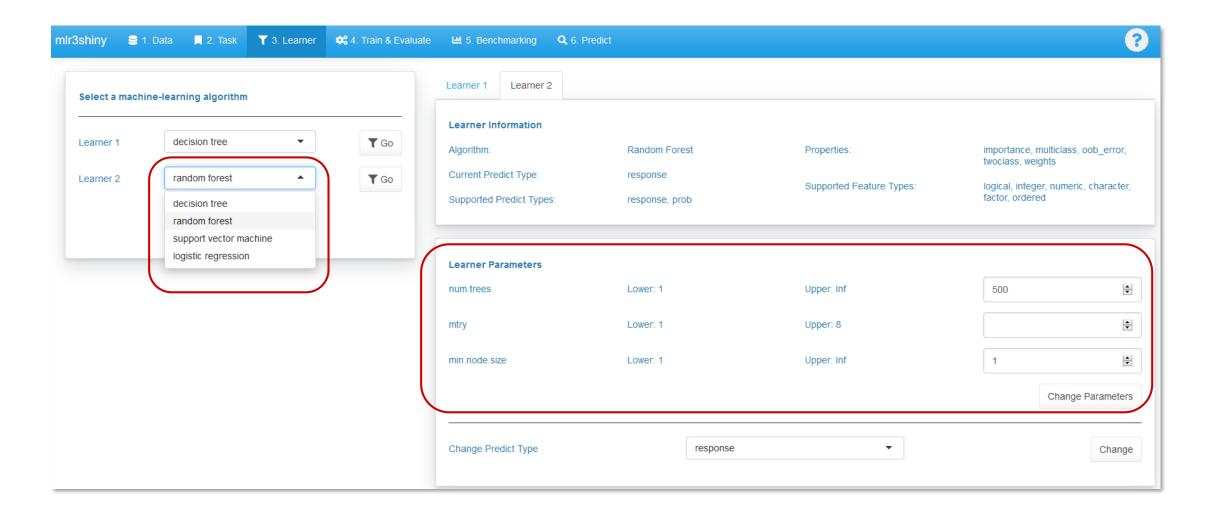
- Variance reduction formula is for independent observations.
- ...If all variables are available to all splits of all trees the trees will look pretty much the same ('be correlated').





Random Forests: chp. 8.2

#### Random Forests in Practice



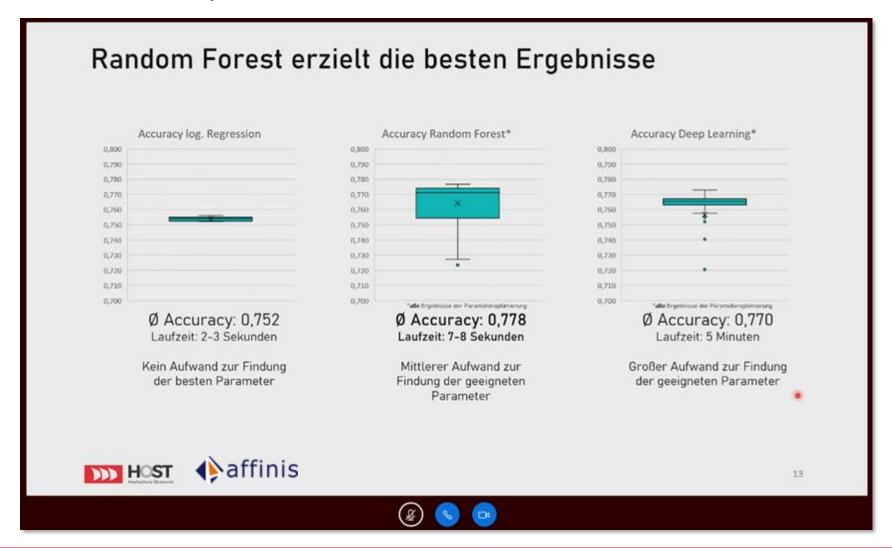
## Random Forest Tuning Parameters

Parameter	Description	Name in	Default
# Trees	Number of trees to grow: - Generally: the larger the better, but increases computation timethumb rule: not smaller than 100.	ntree	500
Min. nodesize	Tree parameter: Minimal number of observations in each terminal node: - Idea of forests: deep trees (overfitting will be averaged out) Strong impact on computation time => can be increased for large data.	nodesize	5
# Variables   Split	Number of randomly sampled variables @ each split: - Important parameter, optimal choice depends on data:too large: similar treestoo small: trees/splits with only unpredictive values may occur.	mtry	√#Variables in data

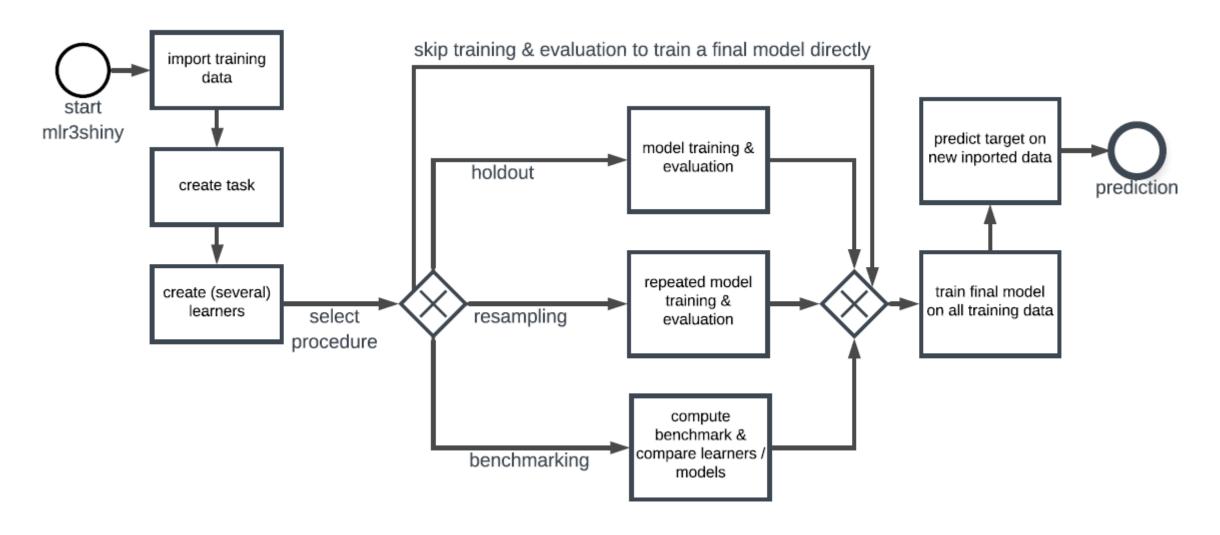
#### For further information, see:

Szepannek, G. (2017): On the Practical Relevance of Modern Machine Learning Algorithms for Credit Scoring Applications, In: Mucha, H.: *Big Data Clustering: Data Preprocessing, Variable Selection and Dimension Reduction*, WIAS Report 29, S. 88-96, DOI: 10.20347/WIAS.REPORT.29.

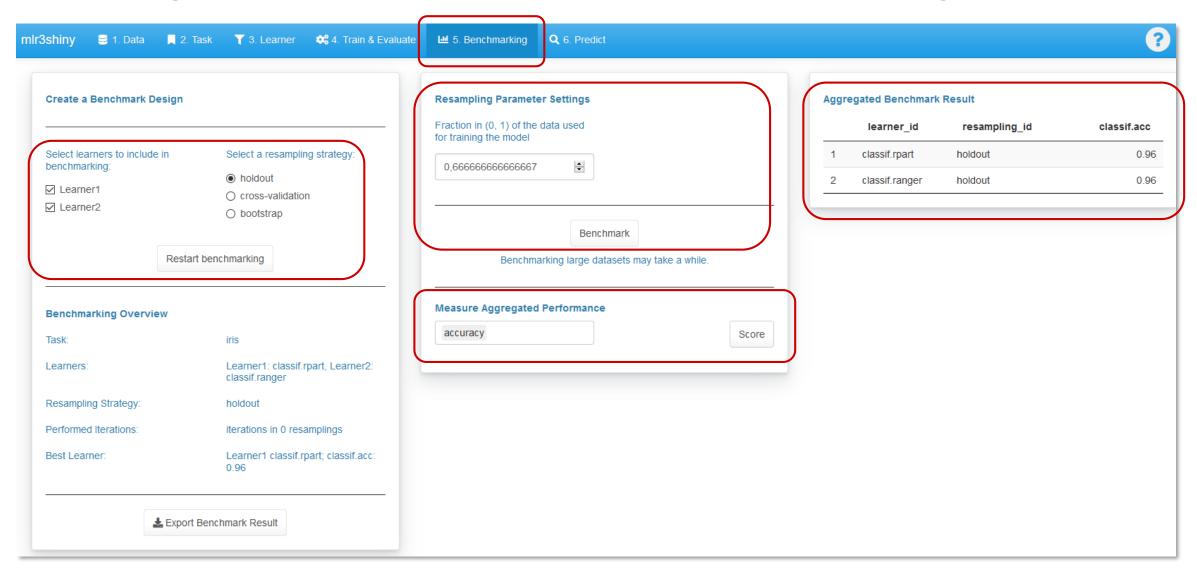
## Recent Example



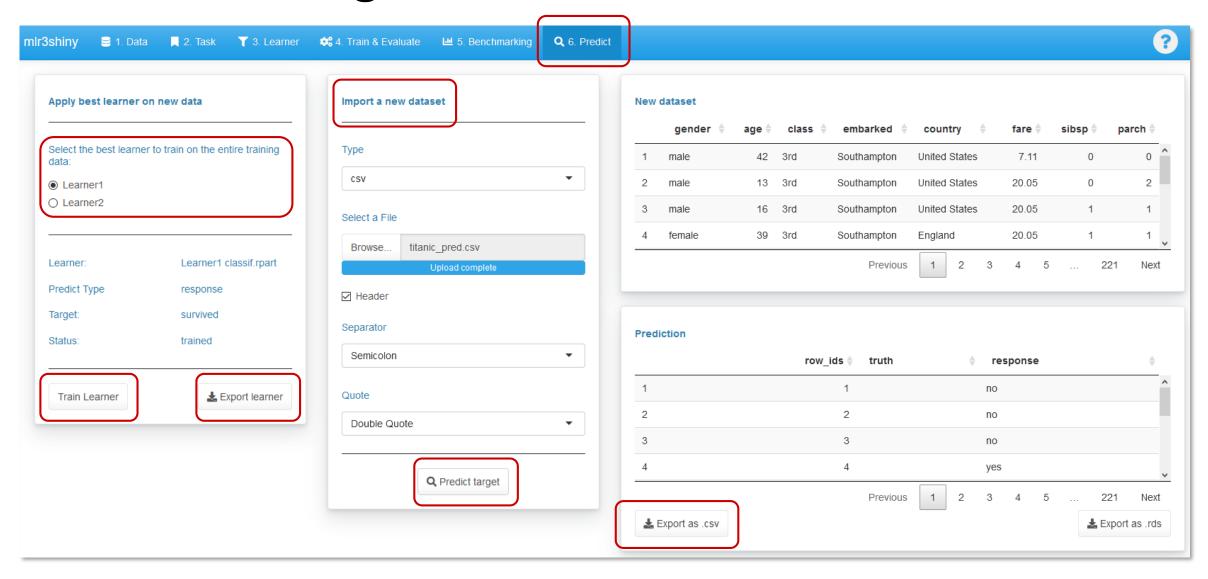
# The General Machine Learning Workflow



# Finding the Best Learner: Benchmarking



### Final Retraining on All Data



# Further Reading

