

# A survey of methods for model validation with applications in R

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- ▶ Get training data
- ▶ Build your model
- ▶ (validate your model??)
- ▶ Get score on a public data set
- ▶ Be ranked **solely** on unknown data

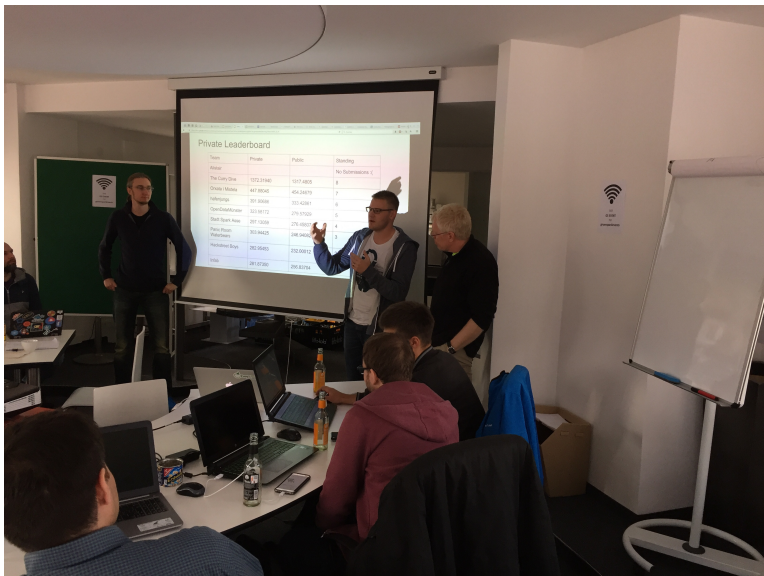


Figure 1: After the Data Dive

## Private Leaderboard

Team	Private	Public
Alistair		
The Curry Dive	1372.31940	1317.4805
Orxata i Mistela	447.88045	454.24679
hafenjungs	391.90686	333.42861
OpenDataMünster	323.58172	279.57929
Stadt Spark Asse	297.13059	276.45807
Panic Room Waterbears	303.94425	246.94082
Hackstreet Boys	282.95453	232.00012
Infab	281.87350	256.83704

Figure 2: After the Data Dive (zoomed in)



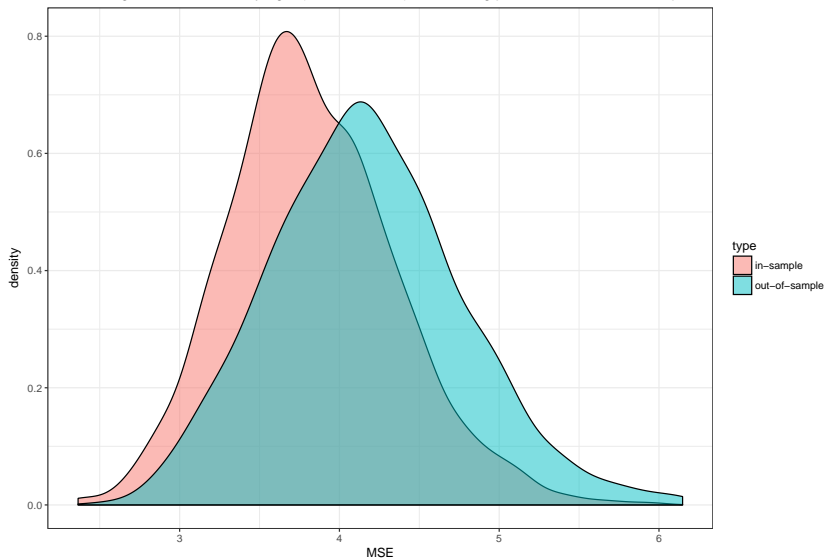
# Goals for today

- ▶ See how different validation methods stack up
- ▶ Some examples in R (`caret` package)

## Is it worthwhile to consider the performance on known data?

```
# Adapted from: http://eranraviv.com
n_sim <- 1000
MSE <- oosMSE <- rep(NA, n_sim)
# define the noise level
sig <- 2
# 3 explanatories
pp <- 3
# define the signal level
signall <- 10
for (i in 1:n_sim) {
  beta = seq(from = 1, to = signall, length.out = 4)
  X = cbind(rep(1, 100), matrix(rnorm(100 * pp), nrow = 100))
  Y = X %*% beta + sig * rnorm(100)
  Ynew = X %*% beta + sig * rnorm(100)
  mod = lm(Y ~ X - 1)
  Yhat = predict(mod)
  MSE[i] = mean((Y - Yhat)^2)
  oosMSE[i] = mean((Ynew - Yhat)^2)
}
```

Even if we get the model exactly right (which we don't), the training performance tends to be optimistic



# Resampling methods

5-fold Cross Validation



5 times 60/40 split



5 times LOO-Bootstrap



## Training / Test split

- ▶ **Idea:** Split dataset in two sets
- ▶ **Variants:**
  - ▶ Repetitions: Can be repeated by randomly sampling new splits  $n$  times and averaging the results

## Cross validation

- ▶ **Idea:** Split into  $k$  distinct sets of size  $\frac{n}{k}$ . For 1 to  $k$ :
  - ▶ Fit the model using the remaining  $n - \frac{n}{k}$  observations
  - ▶ Average the  $k$  test set results
- ▶ **Variants:**
  - ▶ Number of folds ( $k$  usually between 2 and 10)
  - ▶ Repetitions: Average the results of multiple CV runs

## Bootstrapping

- ▶ **Idea:** Draw with replacement a random bootstrap-sample  $x_b$  with size equal to the original sample  $x$
- ▶ **Variants:**
  - ▶ Conventional Bootstrap: Train on  $x_b$ , test on  $x$
  - ▶ Leave-one-out bootstrap: Train on  $x_b$ , test on  $x \setminus x_b$

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- ▶ These methods suffer from bias and variance just like an “ordinary” Machine Learning model!
- ▶ Bias: Resampling can result in a systematically higher (or lower) estimate of the true external performance
- ▶ Variance: Even if unbiased, the estimate may vary considerably around the true external performance

# Simulation

## Data

- ▶ 795 observations
- ▶ Binary outcome: Migraine yes (56%) / no (44%)
- ▶ 9 independent variables (frequency of pain, auxiliary symptoms, age, etc.)

## Classification and regression tree (CART, via rpart)

- ▶ Variables: All 9 variables
- ▶ Maximum tree depth: 4
- ▶ Minimum observations in a node to attempt a split: 6

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## Nested simulation procedure

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4. Compare the performance of the resampling methods in terms of bias and variance

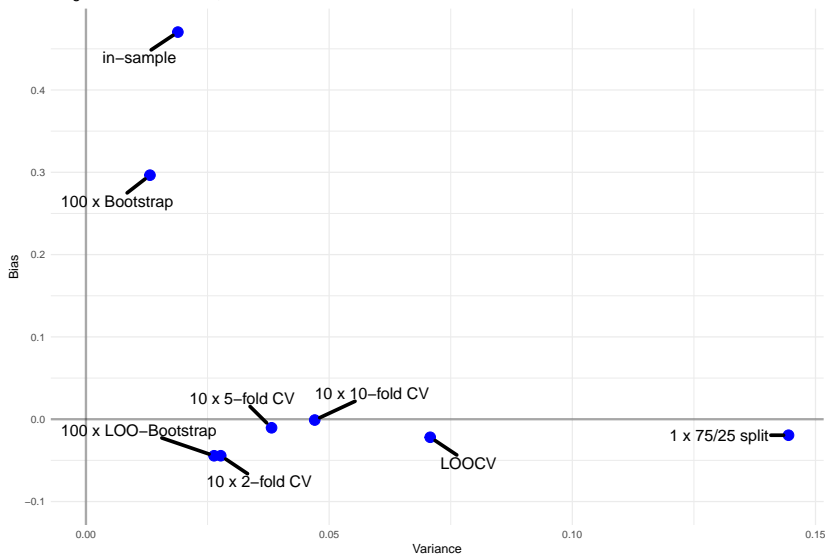


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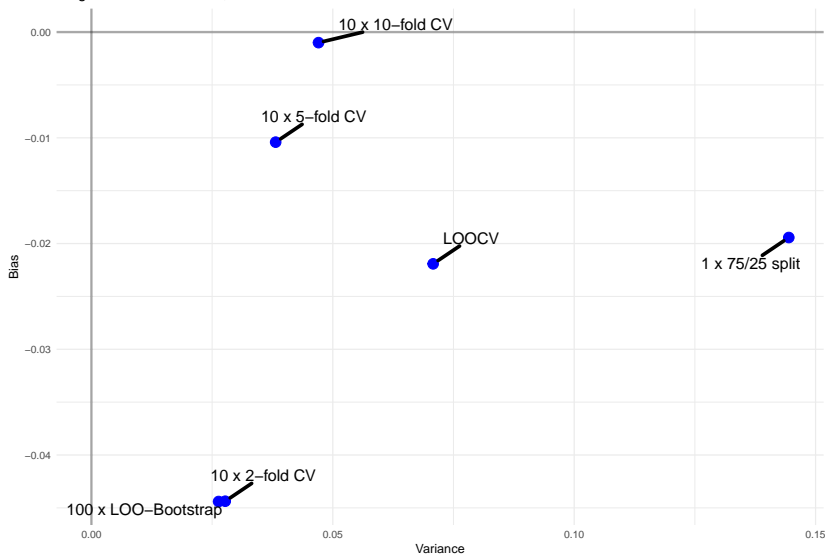
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3. Check pseudo out-of-sample performance on the rest of the data set and compare with the estimate from the internal validation
4. Compare the performance of the resampling methods in terms of bias and variance
5. Repeat simulation steps at least 6000 times (13000 in the case of training / test split)

Single Classification Tree,  $n = 30$

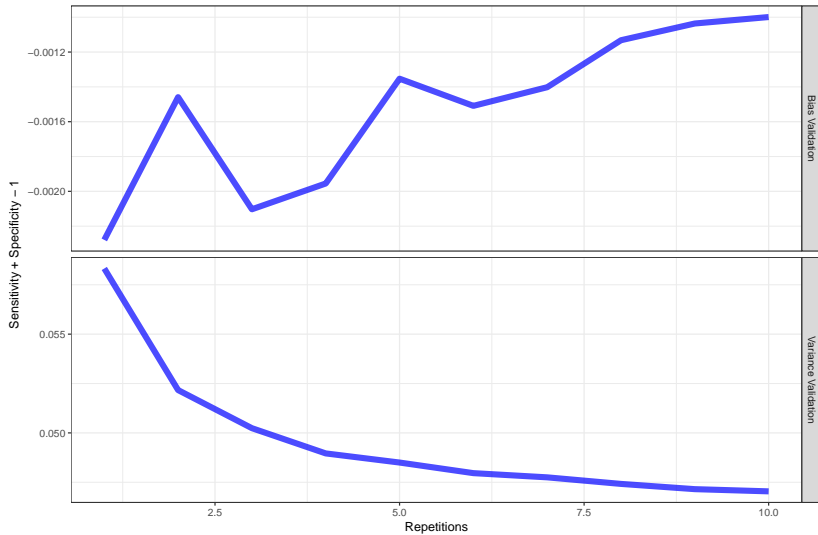


Single Classification Tree,  $n = 30$



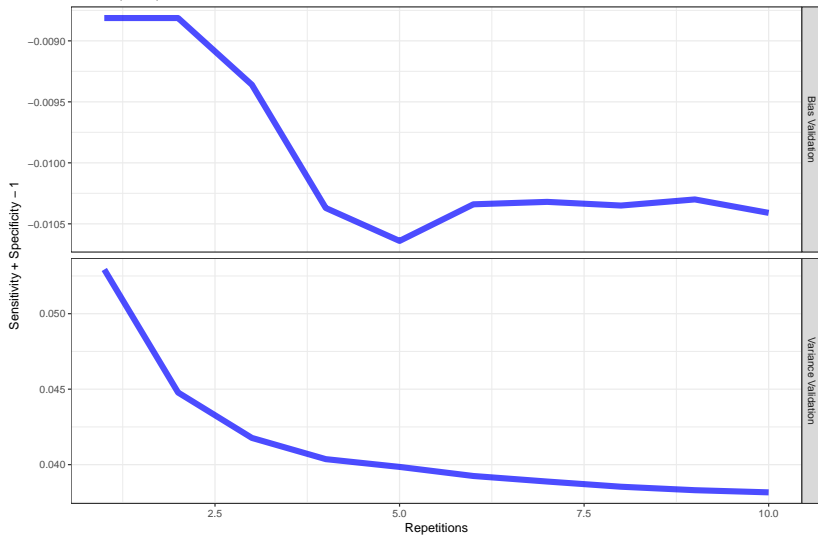
## Effect of more repetitions

10-fold CV, CART,  $n = 30$



## Effect of more repetitions

5-fold CV, CART,  $n = 30$



## Preliminary wrap-up

We've seen that all methods are somewhat biased and variable.  
By doing more repetitions we can lower the variance.

Q: "But what's the method I should use for my task at hand?"

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Figure 4: Well...

A2: Think of model selection vs. error estimation (is bias or variance more important?)



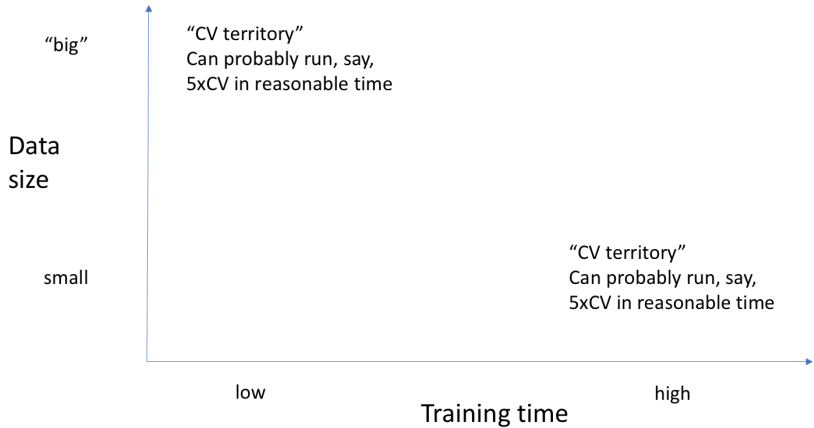


Figure 5:

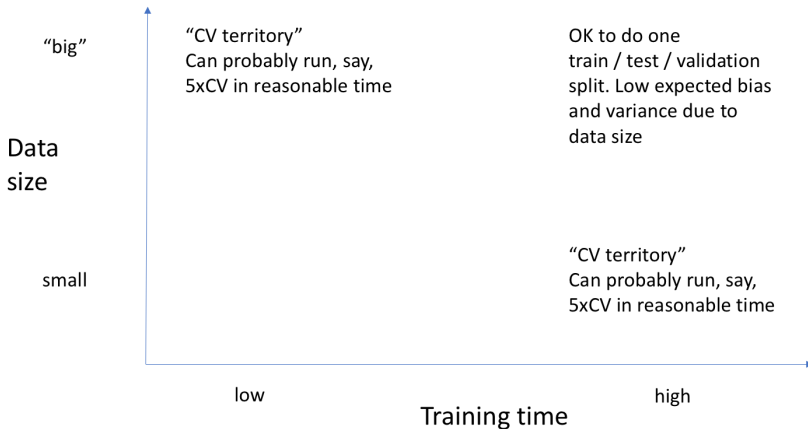


Figure 6:

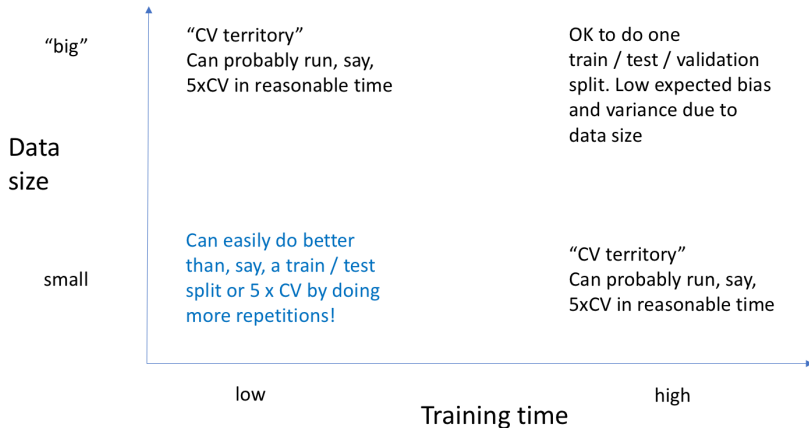


Figure 7:

## Aside: Some attempts to lower the bootstrap bias

- ▶ .632:  $0.632 * \text{conventional bootstrap} + (1 - 0.632) * \text{LOO-bootstrap}$
- ▶ .632+: weight not 0.632 but dependent on amount of in-sample overfitting
- ▶ optimism corrected: optimism = performance on out-of-bag data - conventional bootstrap. Subtract optimism from performance on full sample.

# Appendix: caret and nested CV

## How can I do all this training and testing in R?

Two main options: The caret and the mlr package. We're going to quickly cover the older caret.

```
library(caret)
data(iris) # Yawn...
head(iris)
```

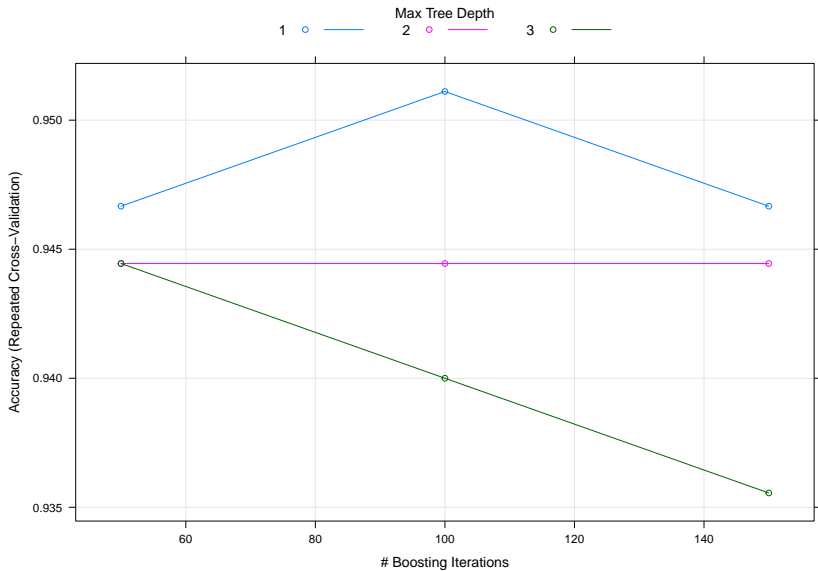
##	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
## 1	5.1	3.5	1.4	0.2	setosa
## 2	4.9	3.0	1.4	0.2	setosa
## 3	4.7	3.2	1.3	0.2	setosa
## 4	4.6	3.1	1.5	0.2	setosa
## 5	5.0	3.6	1.4	0.2	setosa
## 6	5.4	3.9	1.7	0.4	setosa

```
trcont <- trainControl(method = "repeatedcv",  
                        number = 5, repeats = 3,  
                        savePredictions = T,  
                        verboseIter = F,  
                        classProbs = T,  
                        preProcOptions = "nzv")  
mod <- train(Species ~ ., data = iris,  
             trControl = trcont, method = "gbm")
```

```
mod
```

```
## Stochastic Gradient Boosting
##
## 150 samples
## 4 predictor
## 3 classes: 'setosa', 'versicolor', 'virginica'
##
## No pre-processing
## Resampling: Cross-Validated (5 fold, repeated 3 times)
## Summary of sample sizes: 120, 120, 120, 120, 120, 120, ...
## Resampling results across tuning parameters:
##
## interaction.depth n.trees Accuracy Kappa
## 1 50 0.9466667 0.9200000
## 1 100 0.9511111 0.9266667
## 1 150 0.9466667 0.9200000
## 2 50 0.9444444 0.9166667
## 2 100 0.9444444 0.9166667
## 2 150 0.9444444 0.9166667
## 3 50 0.9444444 0.9166667
## 3 100 0.9400000 0.9100000
```

```
plot(mod)
```





```
str(mod$pred)
```

```
## 'data.frame':    4050 obs. of  11 variables:
## $ pred          : Factor w/ 3 levels "setosa","versicolor
## $ obs           : Factor w/ 3 levels "setosa","versicolor
## $ rowIndex      : int   12 16 17 19 22 23 24 30 34 50 ...
## $ setosa        : num   1 1 1 1 1 ...
## $ versicolor    : num   0.0000000573 0.0000002937 0.000000211
## $ virginica     : num   0.0000000000358 0.0000000000105 0.00
## $ shrinkage     : num   0.1 0.1 0.1 0.1 0.1 0.1 0.1 0.1 0.1 0.
## $ interaction.depth: int   1 1 1 1 1 1 1 1 1 1 ...
## $ n.minobsinnode : num   10 10 10 10 10 10 10 10 10 10 ...
## $ n.trees       : num   150 150 150 150 150 150 150 150 150 15
## $ Resample      : chr    "Fold1.Rep1" "Fold1.Rep1" "Fold1.R
```

```
# Select the "best" parameter set
resamples <- mod$pred %>%
  dplyr::filter(shrinkage == 0.1,
                n.minobsinnode == 10,
                n.trees == 50,
                interaction.depth == 1)
dim(iris)
dim(resamples) # 450 rows because 3 repeats of CV
confusionMatrix(reference = resamples$obs,
                 data = resamples$pred)
```

Prediction	Reference		
	setosa	versicolor	virginica
setosa	150	0	0
versicolor	0	140	14
virginica	0	10	136

Overall Statistics

Accuracy : 0.9467  
 95% CI : (0.9217, 0.9655)  
 No Information Rate : 0.3333  
 P-Value [Acc > NIR] : < 2.2e-16

Kappa : 0.92  
 McNemar's Test P-Value : NA

Statistics by Class:

	Class: setosa	Class: versicolor	Class: virginica
Sensitivity	1.0000	0.9333	0.9067
Specificity	1.0000	0.9533	0.9667
Pos Pred Value	1.0000	0.9091	0.9315
Neg Pred Value	1.0000	0.9662	0.9539
Prevalence	0.3333	0.3333	0.3333
Detection Rate	0.3333	0.3111	0.3022
Detection Prevalence	0.3333	0.3422	0.3244
Balanced Accuracy	1.0000	0.9433	0.9367

Figure 8: Confusion matrix using the out-of-fold data

## Repeated CV is no nested CV!

Let's say we want to get an estimate of our model's performance after selecting the best parameter set or the best single model. We need an additional outer CV loop:

```

# Outer Loop
inTrain <- createFolds(iris$Species, k = 5,
                      returnTrain = T)

# I hear the cool kids are using purrr now...
nested_cv <- apply(inTrain, function(train_indices) {
  train_temp <- iris[train_indices, ]
  test_temp  <- iris[-train_indices, ]
  # If nothing is specified, caret trains a Random Forest
  # and evaluates it using Leave-one-out bootstrapping.
  # We're doing selection here, not error estimation,
  # after all. Then picks "best" parameter set.
  mod <- train(x = train_temp[, 1:4],
               y = train_temp$Species)
  preds_temp <- predict(mod, test_temp)
  # Let's calculate accuracy, could be any other metric
  sum(test_temp$Species == preds_temp) / nrow(test_temp)
})
summary(nested_cv) # Now only "normal" bias/variance left

```

```

##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
## 0.8667  0.9333  0.9667  0.9533  1.0000  1.0000

```

# What we did not cover

- ▶ calibration
- ▶ CV for time series
- ▶ t-tests for comparing validation results
- ▶ ...

End of appendix

...and of my presentation

**Thank you!**

# References

caret tutorial

- ▶ <http://topepo.github.io/caret/index.html>

Alternative: mlr

- ▶ <https://mlr-org.github.io/mlr-tutorial/>

Arlot, S., & Celisse, A. (2010). A survey of cross-validation procedures for model selection. *Statistics Surveys*, 4, 40–79.

- ▶ [http://www.di.ens.fr/sierra/pdfs/2010\\_Arlot\\_Celisse\\_SS.pdf](http://www.di.ens.fr/sierra/pdfs/2010_Arlot_Celisse_SS.pdf)

Hastie, Tibshirani, Friedman. The elements of statistical learning

- ▶ [http://statweb.stanford.edu/~tibs/ElemStatLearn/printings/ESLII\\_print10.pdf](http://statweb.stanford.edu/~tibs/ElemStatLearn/printings/ESLII_print10.pdf)

Yu, W., Ruibo, W., Huichen, J., & Jihong, L. (2014). Blocked 3x2 Cross-Validated t-Test for Comparing Supervised Classification Learning Algorithms. *Neural Computation*, 26(1), 208–235.

- ▶ [https://doi.org/10.1162/NECO\\_a\\_00532](https://doi.org/10.1162/NECO_a_00532)



Optimism of the training error

- ▶ <http://eranraviv.com/optimism-training-error-rate/>