

# Validation and Optimization of Machine Learning Models Regression



2021-06-17

ESCAPE Summer School

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# Topics

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- 1 Introduction
- 2 Figures of Merit
- 3 Verification Schemes
- 4 Resampling Techniques
- 5 Model Stability
- 6 Sample Size Planning
- 7 Validation
- 8 Data-driven Model Optimization and Hyperparameter Tuning
- 9 Regression

## Validation & Optimization

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**Verification:** Making sure/measuring/showing that the model meets the specifications.

**Validation:** Making sure that the model meets the application needs.

- Chemometric model validation  
↳ typically verification rather than validation is done.
- Characterize model by measuring its predictive performance

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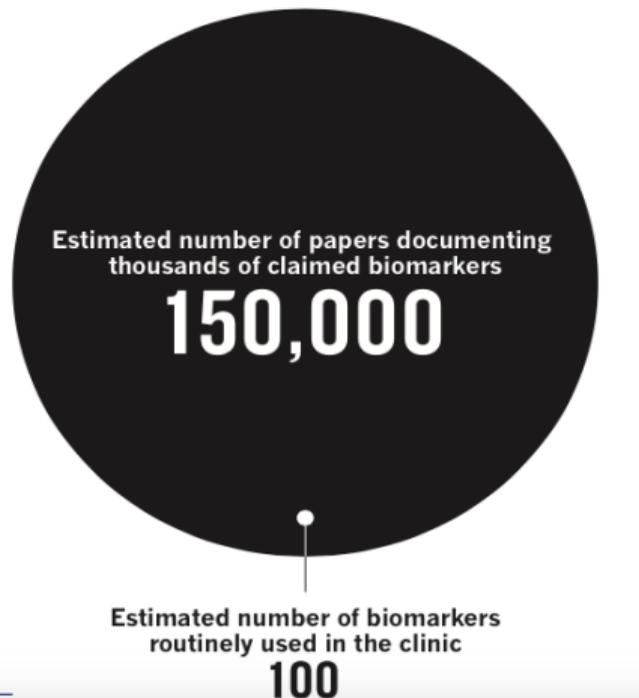
Regression

# Reproducibility!?

## A DROP IN THE OCEAN

Few of the numerous biomarkers so far discovered have made it to the clinic.

*Nature* **469**, 156-157



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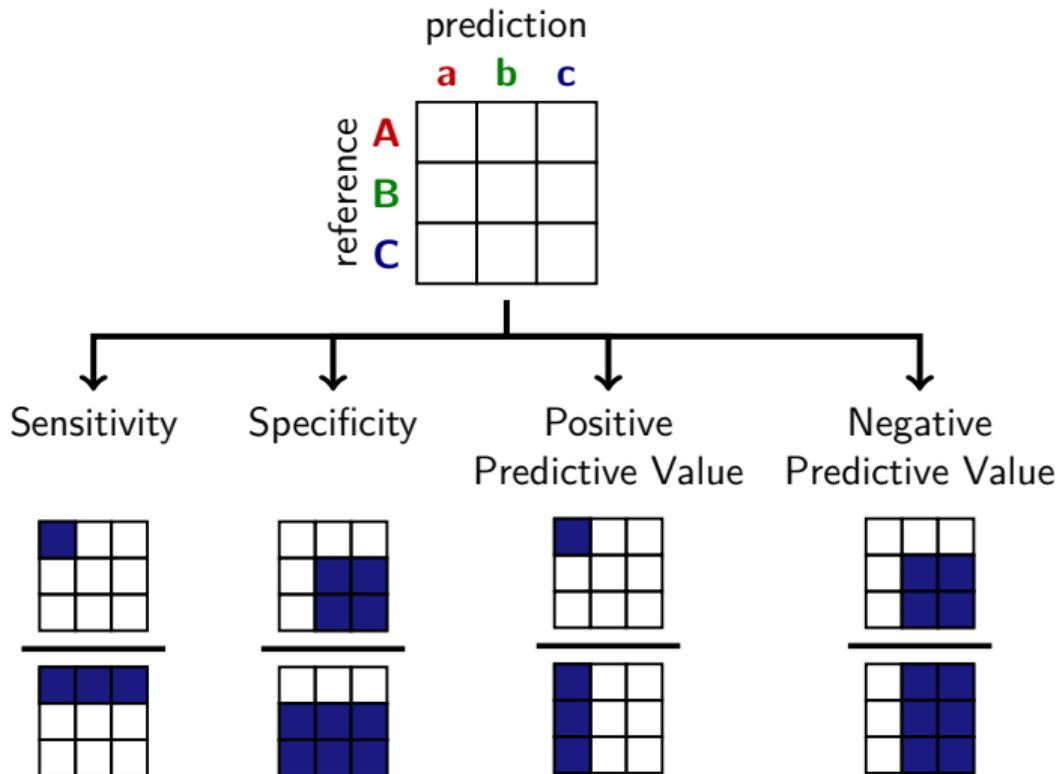
# Recipe: Verification/Validation/Testing

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## Ingredients

- Ready-to-use model  
treated as black box: *case*  $\mapsto$  *prediction*
- Figures of merit (performance measure)  
Overall Accuracy, Sensitivity, Specificity, Predictive Values,  
MSE, RMSE,  $R^2$ , ...
- Validation *scheme*: *How to get test cases?*  
Autoprediction, Resampling, Test Set, Validation study

# Figures of Merit: Proportions



# Proportion Questions

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**Sensitivity = Recall:** of all truly class A cases, which fraction is correctly recognized as class A?

**Specificity:** of all cases truly not belonging to class A, which fraction is correctly recognized as not belonging to class A?

**Positive Predictive Value = Precision:** of all cases predicted to belong to class A, which fraction does truly belong to class A?

**Negative Predictive Value:** of all cases predicted not to belong to class A, which fraction does truly not belong to class A?

**accuracy:** correct proportion among all predicted cases

**error rate:** misclassified proportion among all predicted cases

**K:** chance-corrected accuracy, inter-observer agreement

# Proportions: Characteristics

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- ✓ well-known, widely used
- ✗ often misunderstood:
  - sensitivity & specificity
    - ✓ easy to measure: test  $n$  cases of each class, record results
    - ✗ low relevance for application
  - predictive values (positive/negative)
    - ✓ high relevance for application
    - ✗ difficult to measure: need to know relative class frequencies under application conditionsweight rows of confusion matrix accordingly
  - figures of merit spanning rows of confusion matrix
    - ✗ correct for relative class frequencies under application conditions

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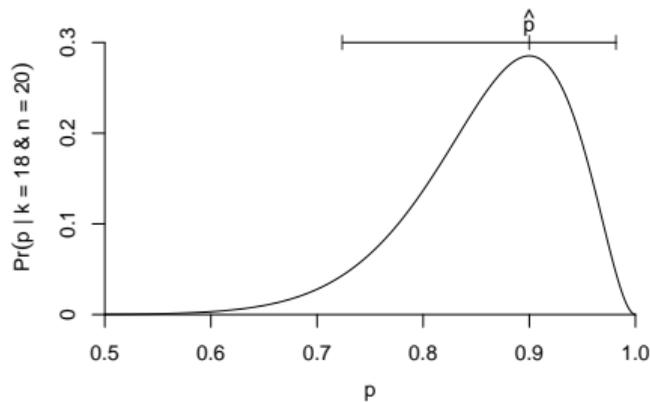
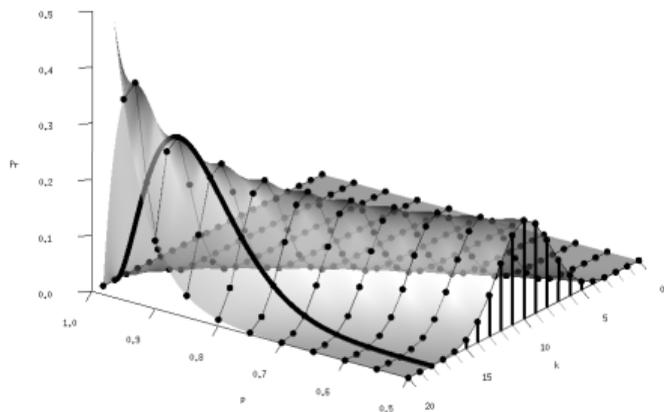
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# More figures of merit

- chance-corrected:  $\kappa$ 
  - ✓ rescaling possible for other figures of merit
  - ✓ alternative: report chance agreement (or naive model performance) together with figure of merit
  
- Information gain
  - **positive likelihood ratio:**  $LR_A^+ = \frac{Sens_A}{1-Spez_A}$   
How much do the odds to belong to class A increase when a case is predicted to belong to class A?
  - **negative likelihood ratio:**  $LR_A^- = \frac{Spez_A}{1-Sens_A}$   
How much do the odds to belong to class A decrease when a case is predicted not to belong to class A?
  - ✓ independent of relative class frequencies under application conditions

# Confidence Intervals for Sensitivity



- Statistical description: Bernoulli trial

✓  $\rightsquigarrow$  use binomial distribution

- Uncertainty on proportion:  $var(\hat{p}) = \frac{p(1-p)}{n_{test}}$

✗ normal approximation appropriate only with  $np \geq 5$  and  $n(1-p) \geq 5$

$\rightsquigarrow$  Estimate necessary  $n_{test}$

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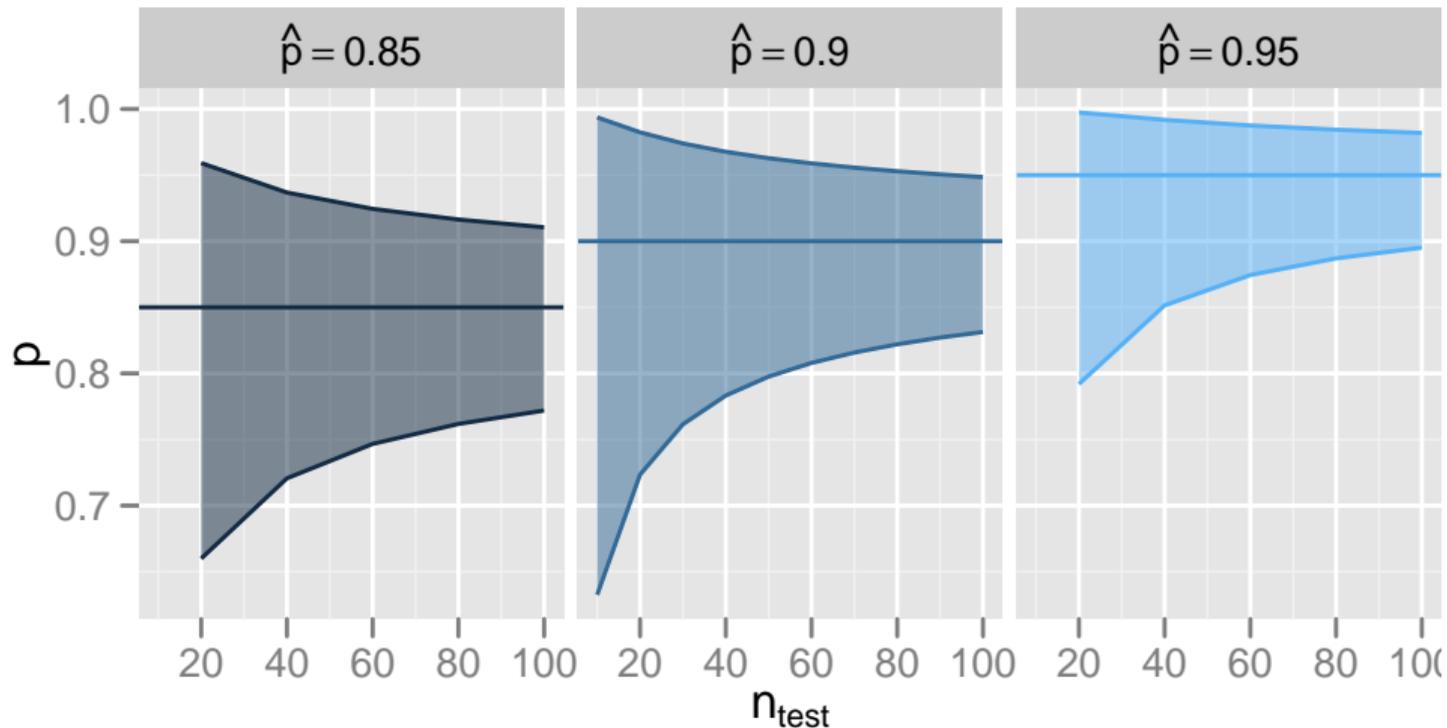
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# Sample size from Confidence Interval



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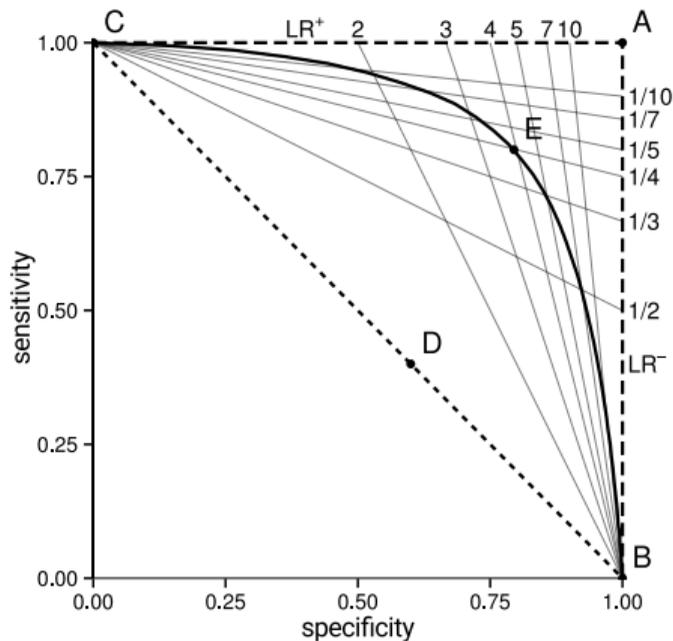
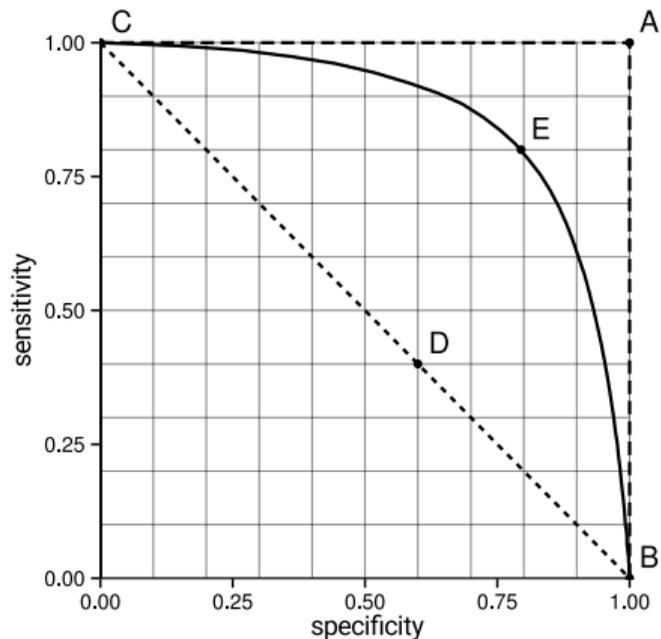
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# Receiver Operating Characteristic/Specificity-Sensitivity-Diagram



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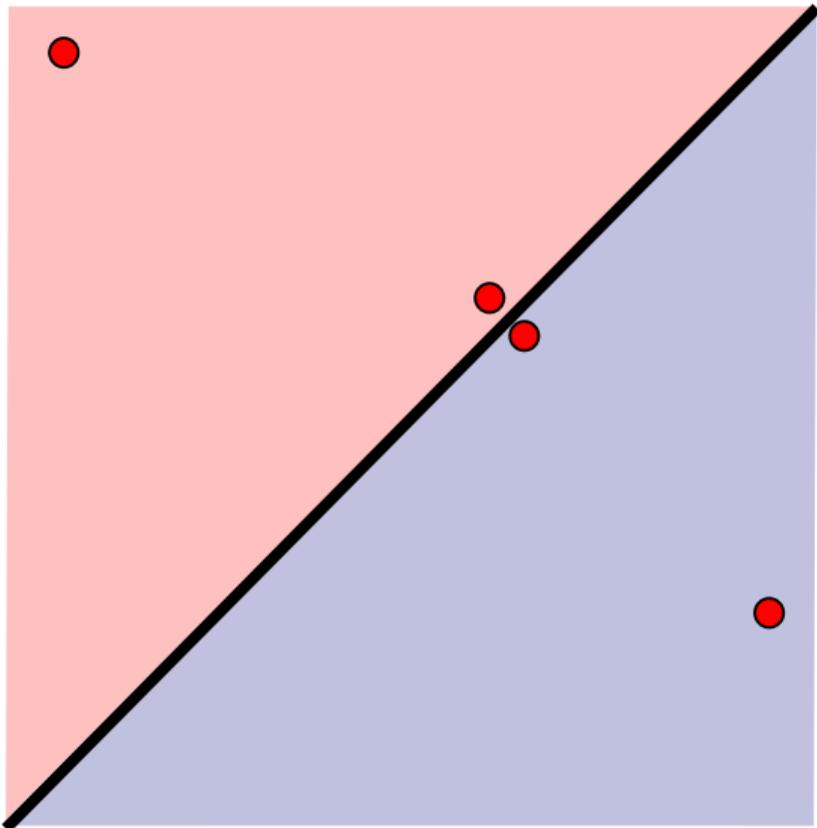
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# Proportions: Behaviour in Hyperparameter Optimization



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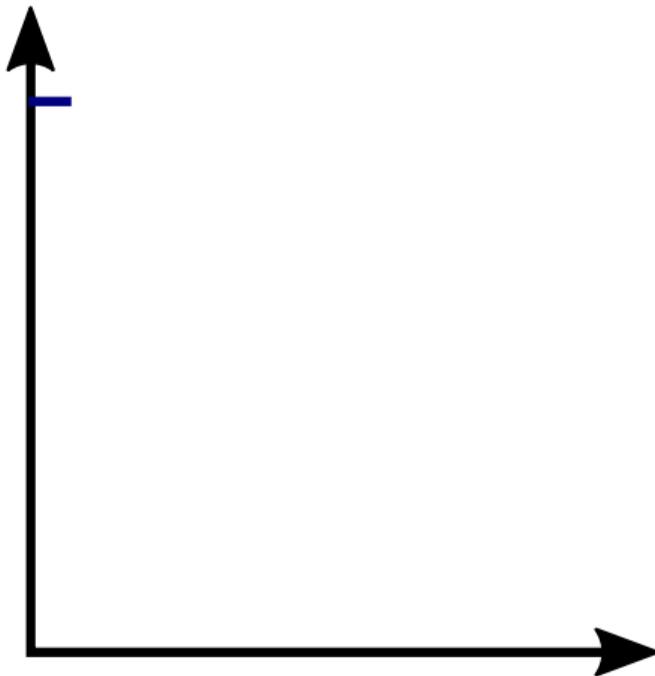
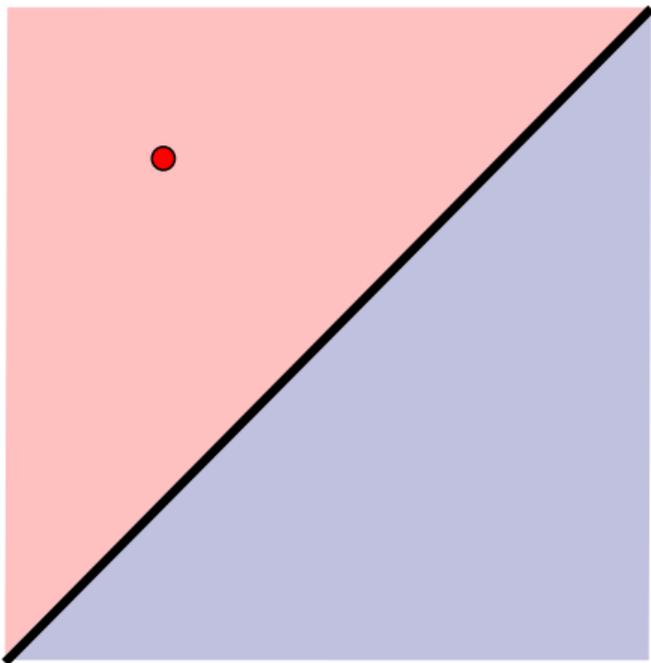
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# Proportions: Behaviour



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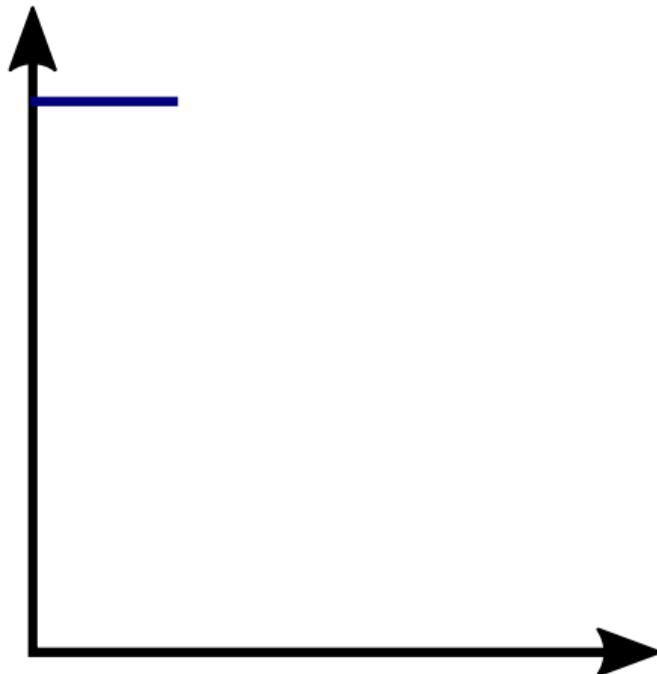
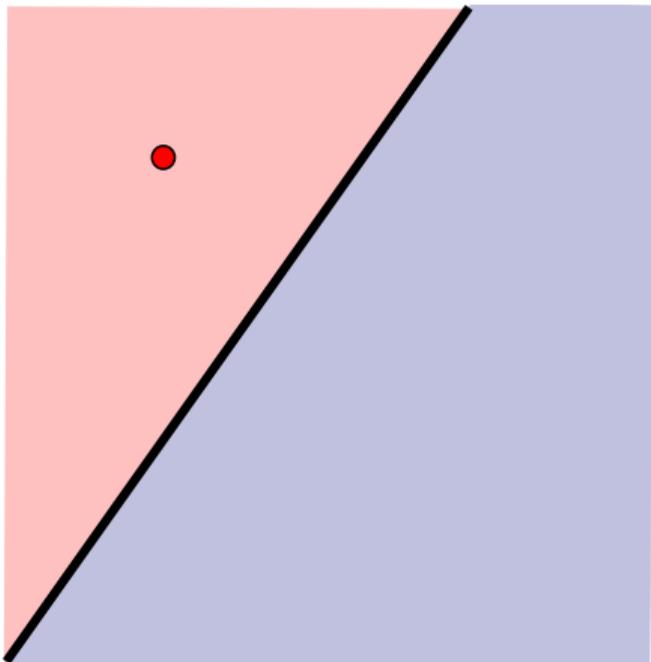
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# Proportions: Behaviour



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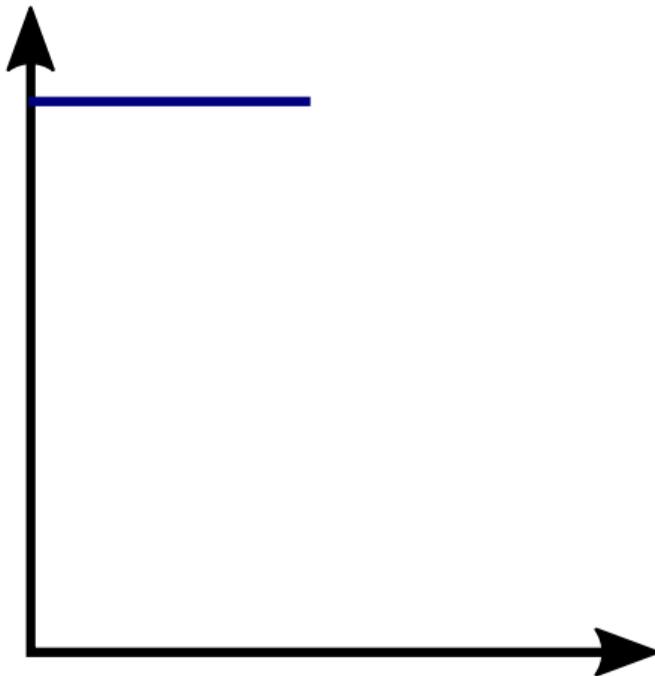
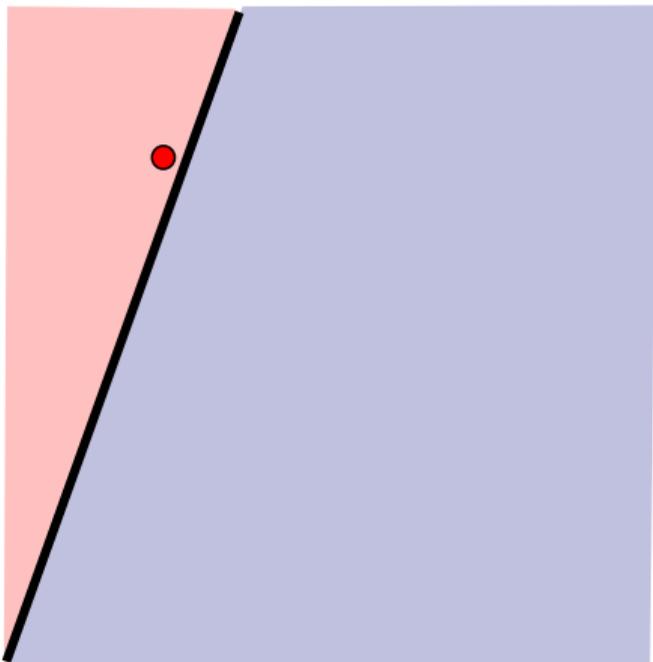
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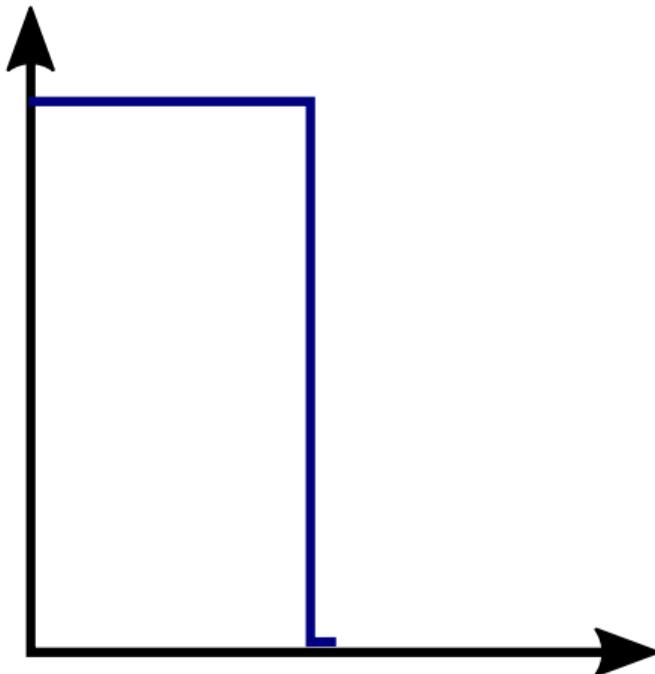
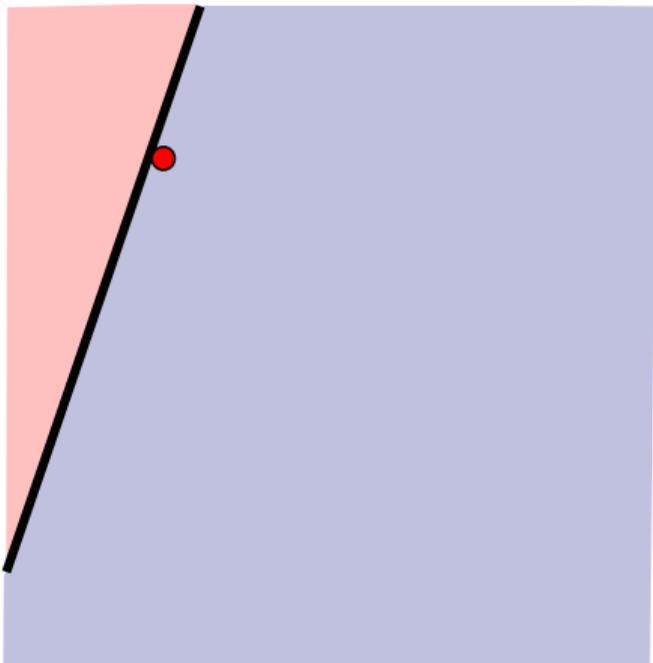
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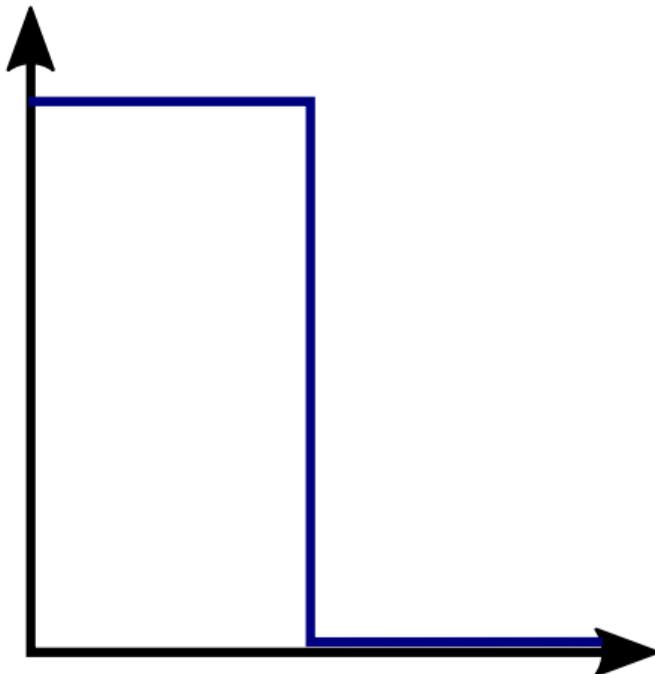
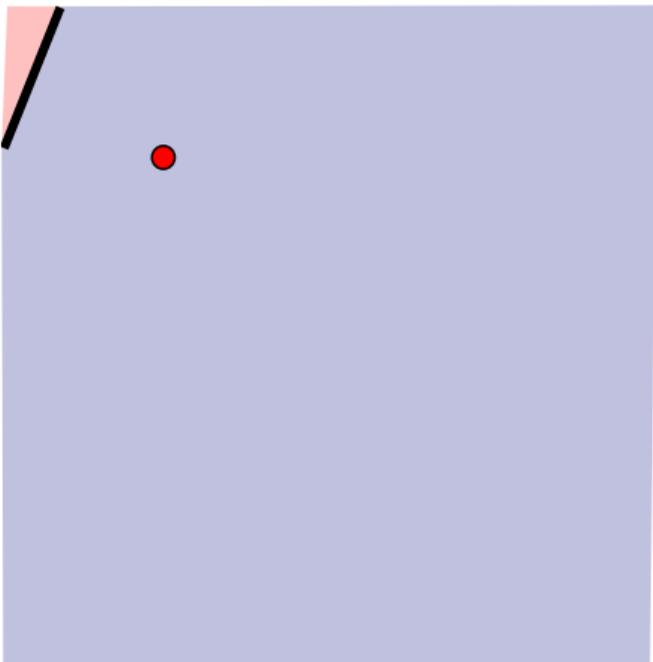
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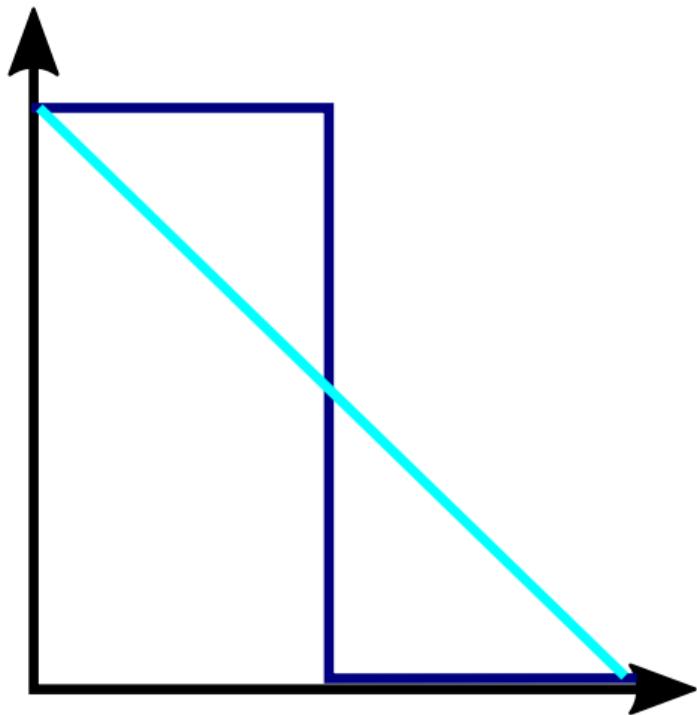
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# (Strictly) Proper Scoring Rules



Wanted: Figure of merit that ...

- ... continuously penalizes closeness to class boundary
- ... continuously reacts to changes in the model
- ... slight deterioration  $\rightsquigarrow$  slight drop in measured performance
- ... has exactly one optimum
- at the best classifier.

# Regression: (Root) Mean Squared Error

- Loss behind e.g. Gaussian Least Squares
- penalizes large deviations
- $MSE = \frac{1}{N} \sum_{i=1}^N (\hat{y}_i - y_i)^2$   
 $N$  ... number of cases  
 $i$  ... case in question
- $MSE = bias^2 + variance$
- $RMSE = \sqrt{\frac{1}{N} \sum_{i=1}^N (\hat{y}_i - y_i)^2}$   
same scale as  $y$

# Brier's Score: Mean Squared Error for Classification

- Classifier that predicts class membership probability rather than labels
- Idea: of all cases where classifier predicts  $x\%$  class membership,  $x\%$  should belong to class in question  
a.k.a. *well calibrated prediction*
- Brier's score:  $BS = \frac{1}{N} \sum_{i=1}^N (\hat{p}_i - p_i)^2$  or  
 $BS = \frac{1}{N} \sum_{j=1}^R \sum_{i=1}^N (\hat{p}_{ij} - p_{ij})^2$  (multiclass version) with  
 $N$  ... number of cases  
 $i$  ... case in question  
 $R$  ... number of classes  
 $j$  ... class in question  
 $p$  ... class membership, usually  $\in \{0, 1\}$   
 $\hat{p}$  ... predicted class membership  $\in [0, 1]$

# Summary Figures of Merit

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- Many exist  $\rightsquigarrow$  choose relevant ones
  - Usually, several figures of merit are needed for characterization
  - Figures of merit are *measured*  $\rightsquigarrow$  subject to bias and variance like any other measurement
  - Regression: figures of merit “well-behaved”, but no back-of-the-envelope variance guesstimates
  - Classification: proportions easy to understand & widespread but have bad variance properties & discontinuous behaviour
- $\rightsquigarrow$  use (strictly) proper scoring rules for optimization

# Cases and Statistical Independence

0 1 2 3 4 5 6 7 8 9  
0 1 2 3 4 5 6 7 8 9  
0 1 2 3 4 5 6 7 8 9

0 1 2 3 4 5 6 7 8 9  
0 1 2 3 4 5 6 7 8 9  
0 1 2 3 4 5 6 7 8 9

- structure in data: clusters, spatial and/or temporal
  - e.g. repeated measurements: repetitions are more similar to each other
  - special case: time series
- ✓ Think hard about factors affecting independence

# Model Testing: Measure the Model's Performance

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## Different kinds of test samples

↪ different performance measures

**Goodness of fit:** *training samples*

↪ residuals

**Generalization error:** statistically independent samples

resampling,

test set measured at same time as training set

**Future performance:** samples measured *after* training samples

dedicated test set for detection of drift

# Resampling for Model Validation

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- ✘ We don't have enough samples
- Training:
  - Model quality depends on ratio  $n_{train} : d.f.$
  - Linear model: 5 samples/(variate · class)
  - ✓ We want to use all samples for training

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✘ We don't have enough samples

- Training:

- Model quality depends on ratio  $n_{train} : d.f.$

- Linear model: 5 samples/(variate · class)

- ✓ We want to use all samples for training

- Testing:

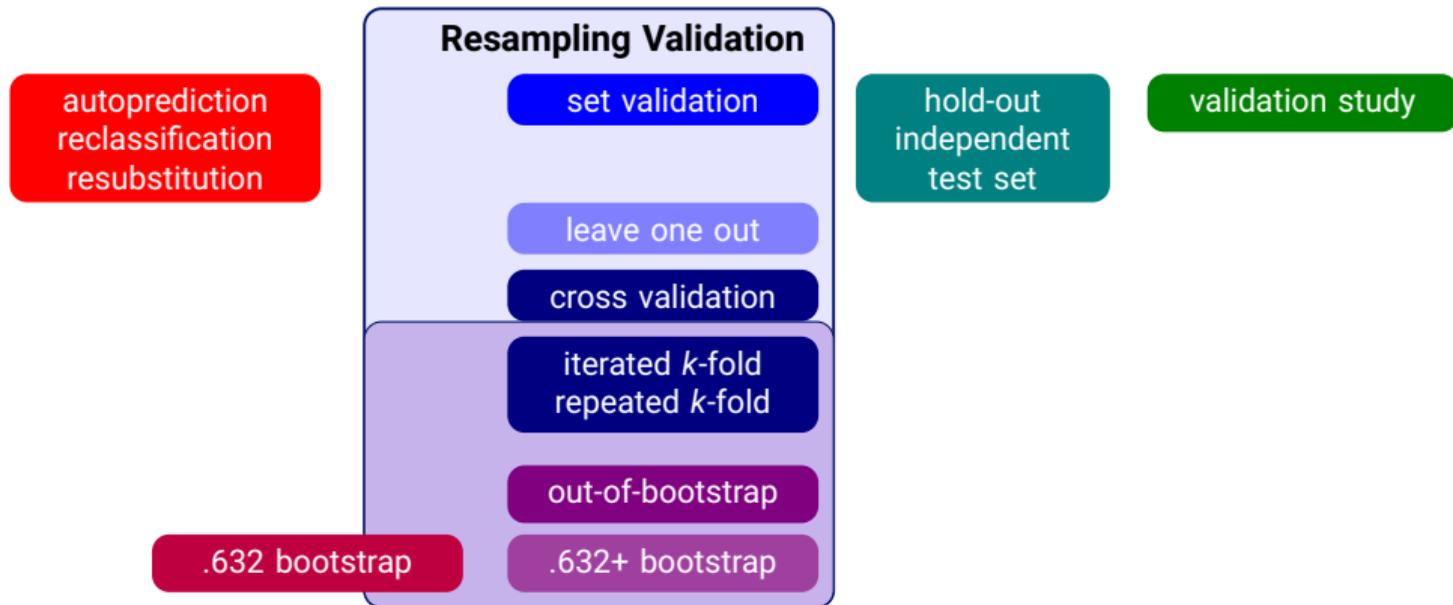
- ✓ We want to know whether the model is stable

- Quality of the performance measure depends on  $n_{test}$

- Width of 95 % confidence interval  $\overset{!}{\leq} 10\%$  for  $p = 90\%$ :  $n_{test} \geq 140$

- ✓ We want to use all samples for testing

# Validation Schemes: Overview



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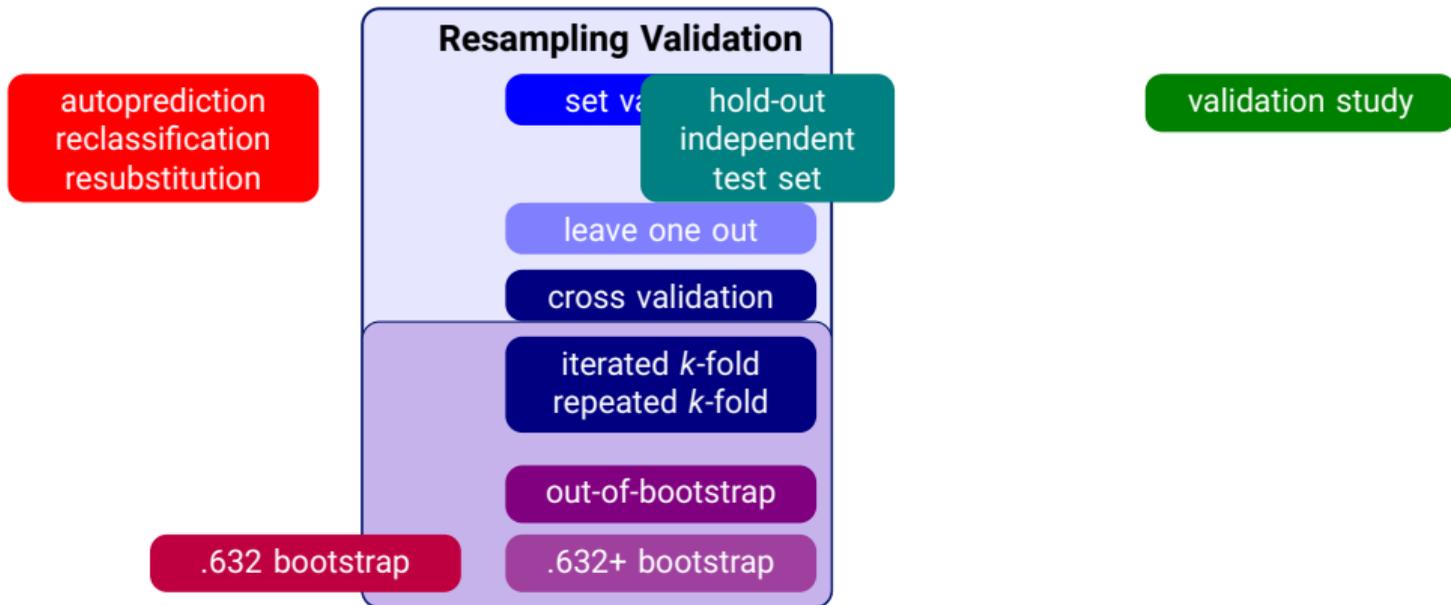
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R Kohavi: A Study of Cross-Validation and Bootstrap for Accuracy Estimation and Model Selection. In: Proc. 14<sup>th</sup> International Joint Conference on Artificial Intelligence, 1995, Morgan Kaufmann, USA, 1995, 1137–1145.

Beleites, C. et al.: Variance reduction in estimating classification error using sparse datasets, Chemom Intell Lab Syst, 2005, 79, 91 - 100

# Validation Schemes: Overview



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# Validation Schemes: Overview

autoprediction  
reclassification  
resubstitution

## Resampling Validation

set validation

leave one out

cross validation

iterated  $k$ -fold  
repeated  $k$ -fold

out-of-bootstrap

.632 bootstrap

.632+ bootstrap

hold-out  
independent  
test set

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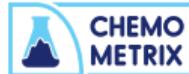
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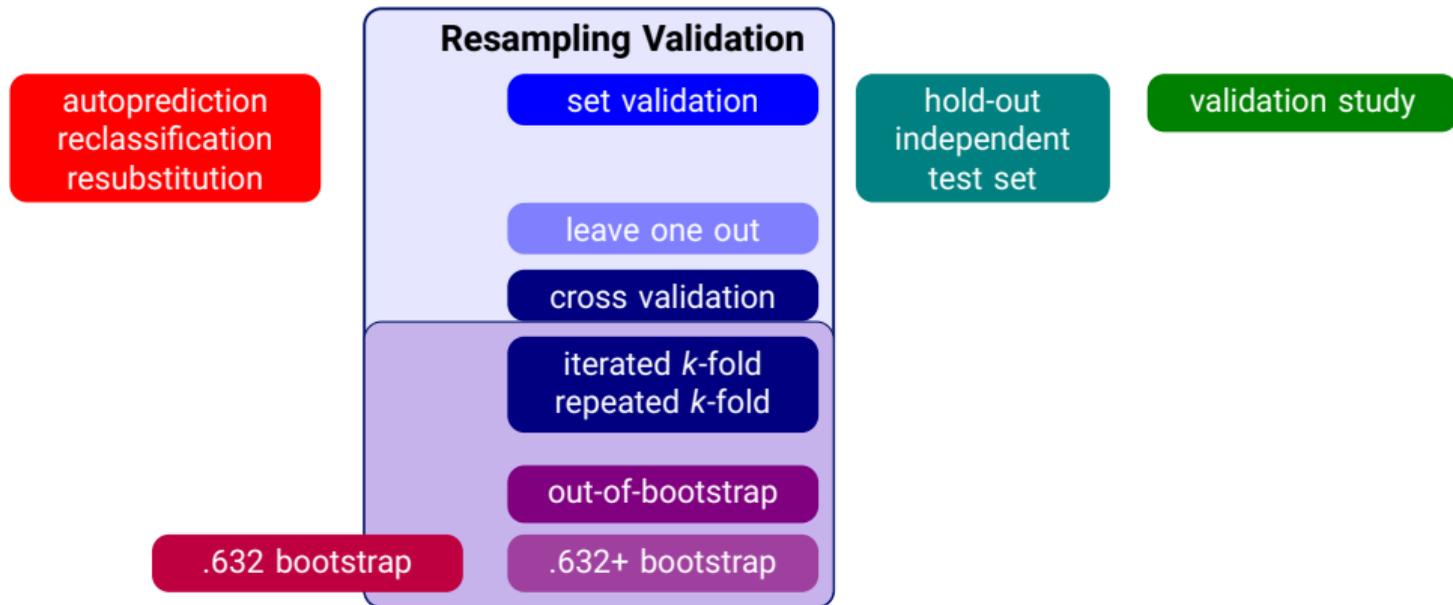
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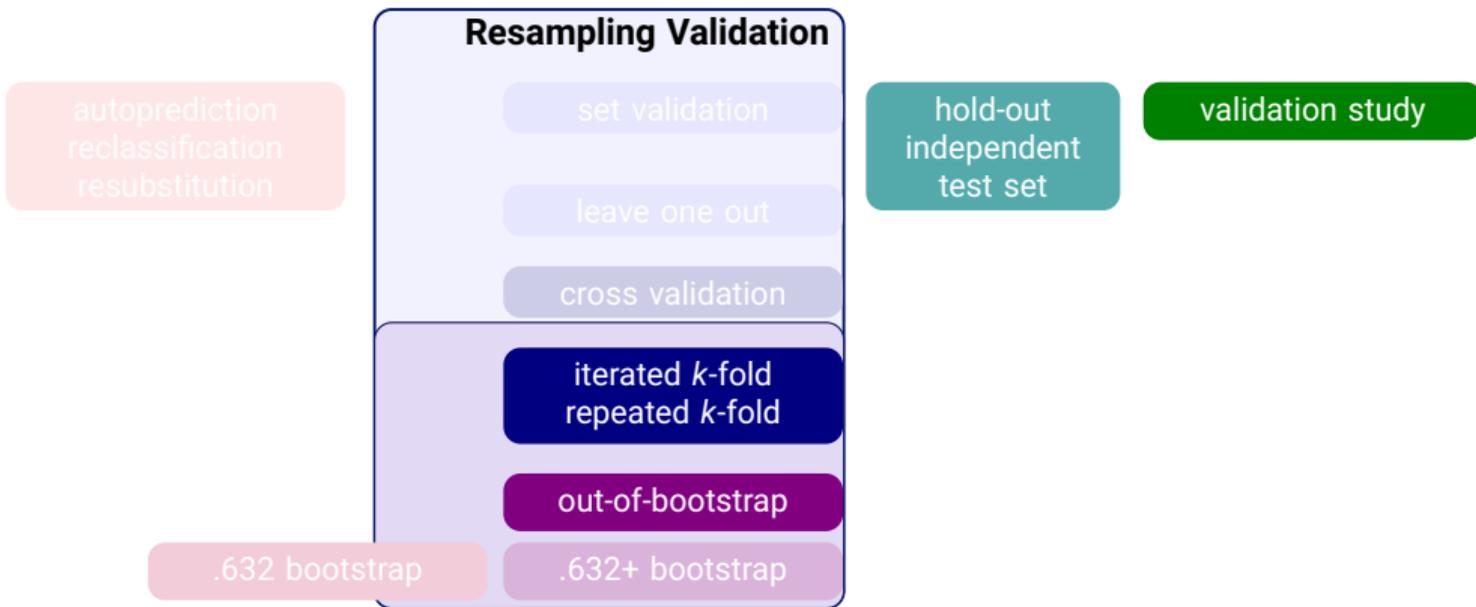
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# Validation Schemes: Recommendations



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# Resampling vs. Validation Study

## statistical properties

bias  
variance  
efficient use of cases  
measure model stability  
measure drift  
future case performance  
out-of-spec cases

## Resampling

✓ pessimistic (low)  
 $f(n)$   
✓  
✓ iterated  
✗  
✗  
✗

## Validation Study

✓ unbiased  
 $f(n_{test})$   
✓  
✓/✗  
✓ DoE  
✓ DoE  
✓ DoE

## practical properties

independence  
effort

⚠ splitting error prone  
✓ computational  
✗ experimental

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# Resampling vs. Hold Out

## statistical properties

bias  
variance  
efficient use of cases  
measure model stability  
measure drift  
future case performance  
out-of-spec cases

## Resampling

✓ pessimistic (low)  
✓  $f(n)$  lower  
✓  
✓ iterated  
✗  
✗  
✗

## Split off Hold Out Set

✓ unbiased  
✗  $f(n_{test})$  large  
✗  
✗  
✗  
✗  
✗(✓)

## practical properties

independence  
effort

⚠ splitting error prone  
✓ computational

⚠ same as resampling  
✓ low

## Validation & Optimization

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Beleites, C. & Salzer, R.: Assessing and improving the stability of chemometric models in small sample size situations Anal Bioanal Chem, 2008, 390, 1261-1271

Esbensen, K. H. & Geladi, P.: Principles of Proper Validation: use and abuse of re-sampling for validation J Chemom, 2010, 24, 168-187



# Validation Check List

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- Randomize order of measurements
- Split at highest level in sample hierarchy/data structure
- Split before 1<sup>st</sup> step that involves multiple cases
- Additional independent validation for data-driven optimization/tuning/model selection
- Test cases: reference labels must be independent of cases (measurements, spectra, ...)
- Make sure labelling procedure does not distort difficulty for test cases
- Ensure correctness of code

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# Validation Check List

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- Randomize order of measurements
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patients, strains, cell lines,
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# Validation Check List

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- Randomize order of measurements
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patients, strains, cell lines,  
day of measurement, before/after new calibration, ...
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- Split before 1<sup>st</sup> step that involves multiple cases  
centering, PCA preprocessing, ...
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patients, strains, cell lines,  
day of measurement, before/after new calibration, ...
- Split before 1<sup>st</sup> step that involves multiple cases  
centering, PCA preprocessing, ...
- Additional independent validation for data-driven optimization/tuning/model selection  
nested/double cross validation or train-validate-test ↔ necessary case numbers HUGE
- Test cases: reference labels must be independent of cases (measurements, spectra, ...)
  
- Make sure labelling procedure does not distort difficulty for test cases
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centering, PCA preprocessing, ...
- Additional independent validation for data-driven optimization/tuning/model selection  
nested/double cross validation or train-validate-test ↔ necessary case numbers HUGE
- Test cases: reference labels must be independent of cases (measurements, spectra, ...)  
cluster analysis to assign labels → OK for training cases
- Make sure labelling procedure does not distort difficulty for test cases
- Ensure correctness of code

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- Split before 1<sup>st</sup> step that involves multiple cases  
centering, PCA preprocessing, ...
- Additional independent validation for data-driven optimization/tuning/model selection  
nested/double cross validation or train-validate-test ↔ necessary case numbers HUGE
- Test cases: reference labels must be independent of cases (measurements, spectra, ...) cluster analysis to assign labels → OK for training cases semi-supervised learning → OK for training cases
- Make sure labelling procedure does not distort difficulty for test cases
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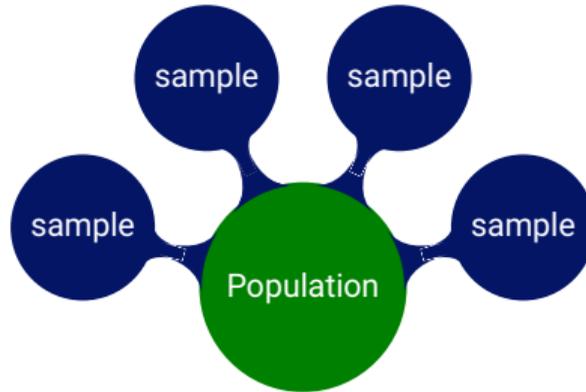
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# The Concept behind Resampling

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**Resampling**

Model Stability

Sample Size

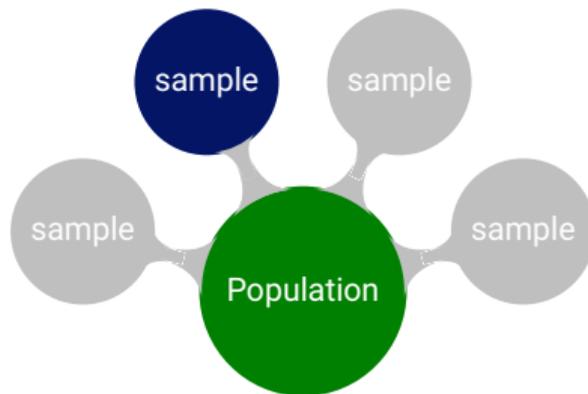
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# The Concept behind Resampling

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## Validation & Optimization

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Model Stability

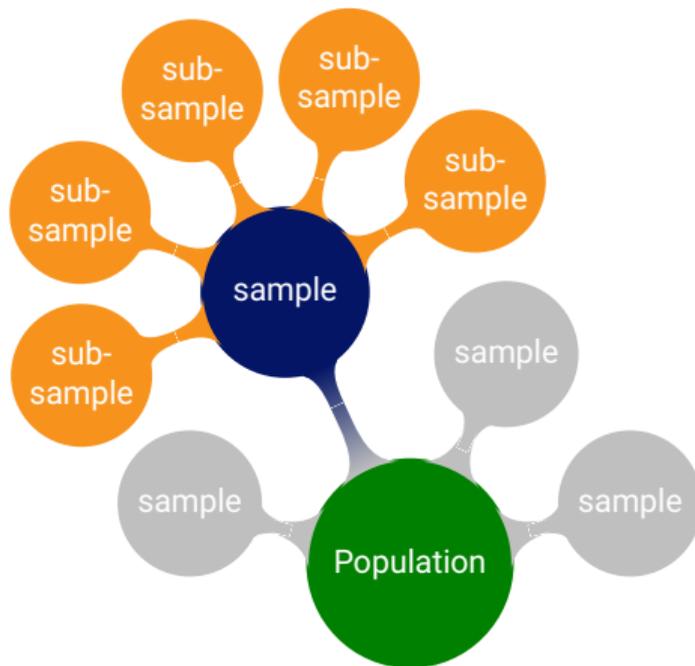
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# The Concept behind Resampling



- Subsamples are approximations of (more) real samples
- Subsample is perturbed version of the real sample

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# Cross Validation: Drawing without Replacement

1	2	3	4	5	6
1	2	3	4	5	6
1	2	3	4	5	6

5	4	2	6	1	3
5	4	2	6	1	3
5	4	2	6	1	3

✓ Each case is left out exactly once

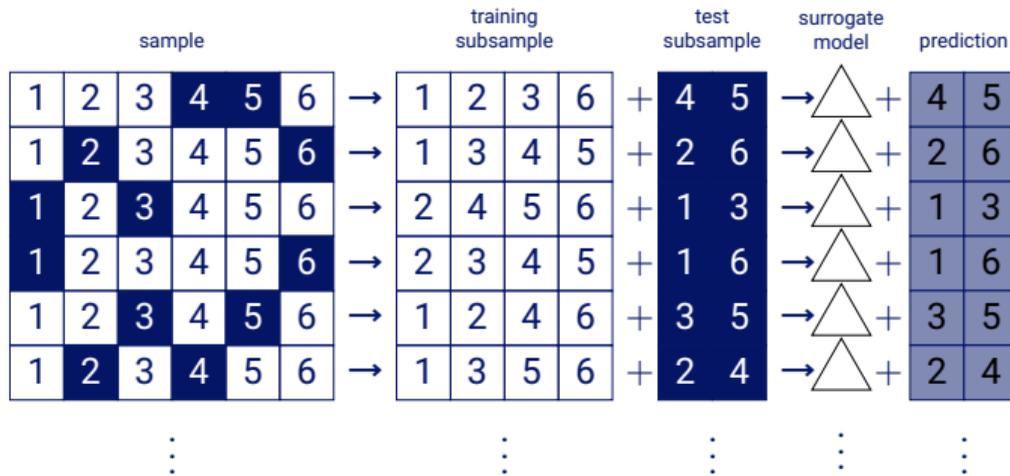
# Cross Validation: Drawing without Replacement

1	2	3	4	5	6
1	2	3	4	5	6
1	2	3	4	5	6
1	2	3	4	5	6
1	2	3	4	5	6
1	2	3	4	5	6
⋮					

5	4	2	6	1	3
5	4	2	6	1	3
5	4	2	6	1	3
1	6	5	3	2	4
1	6	5	3	2	4
1	6	5	3	2	4
⋮					

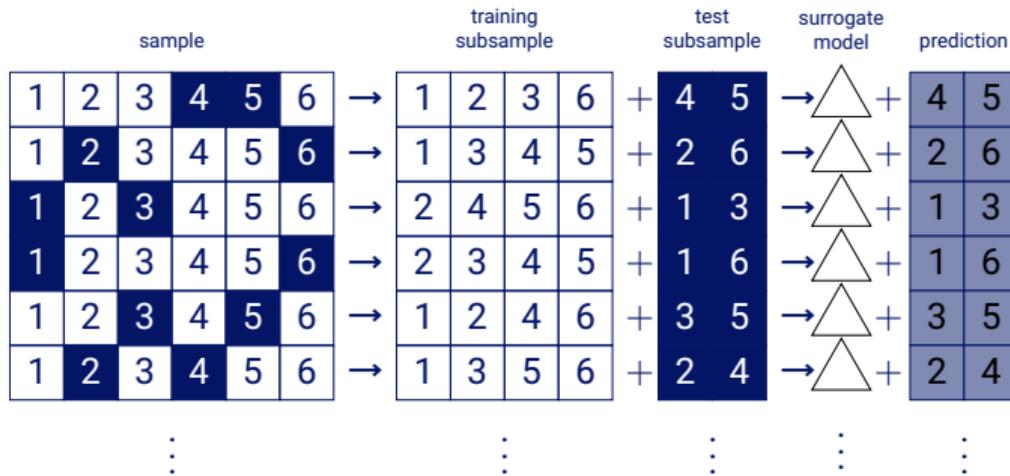
- ✓ Each case is left out exactly once per iteration
- Repetitions aka iterations possible with  $k$ -fold or leave- $n$ -out cross validation
- ✗ Leave-one-out cannot be iterated

# Resampling for Model Validation: Assumptions



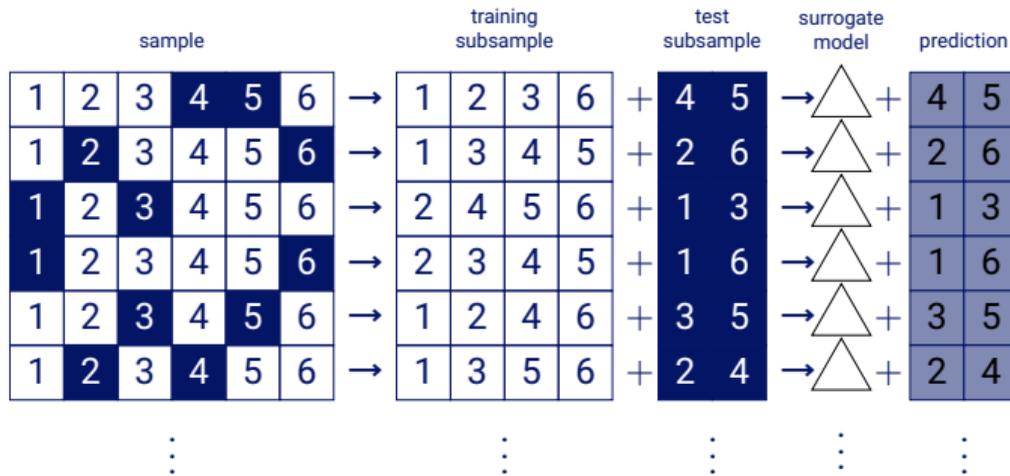
- Surrogate model equals model of whole sample
- Surrogate models equal to each other
- All cases come from the same distribution

# Resampling for Model Validation: Assumptions



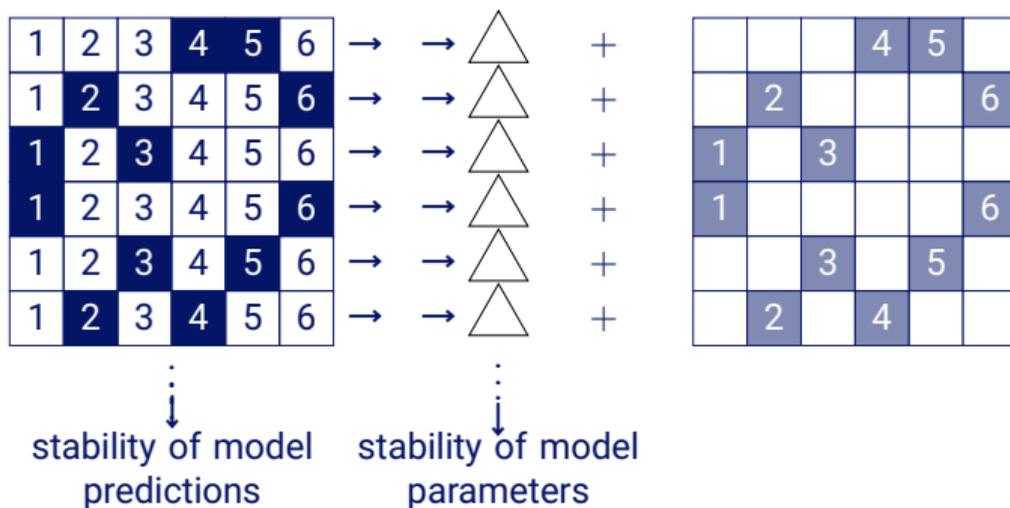
- Surrogate model equals model of whole sample
- ✗ Violation  $\rightsquigarrow$  pessimistic bias
- Surrogate models equal to each other
- All cases come from the same distribution

# Resampling for Model Validation: Assumptions



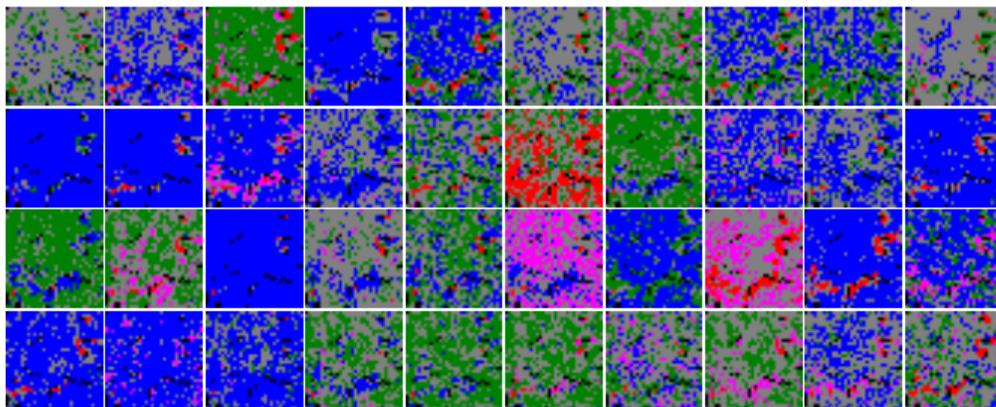
- Surrogate model equals model of whole sample
- ✗ Violation  $\rightsquigarrow$  pessimistic bias
- Surrogate models equal to each other
- ✗ Violation (instability)  $\rightsquigarrow$  higher variance
- All cases come from the same distribution

# Model Stability



- Subsamples are perturbed versions of real sample
- ✓ Measure stability of model
  - Stability of model parameters
  - Stability of predictions
- Repetitions reduce variance due to instability of surrogate models.

# Model Stability: 40× 8-fold cross validation



- FTIR images of tumour sections (normal, °II, °III, °IV)
- total: 150 images of 58 patients: 133 000 spectra  
smallest class: °II, 4 800 spectra (3 patients, 5 images)
- LDA after automatic selection of 8 spectral regions
- reject spectra with posterior probability <math>< 0.85</math>

# How many cases do we need?

## ... to train a good classifier?

- rules of thumb  
linear model:  $\frac{n}{p} \geq 3 - 5$  in each class

⇒ learning curve

## ... to measure the model's performance?

- ↔ confidence intervals for test results
- Rules of thumb  
100 test cases to estimate a proportion
- Regression ↔ needs preliminary experiment

# Validation: Questions

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- Did you ask the right question?
- Or did you use a surrogate?
  - Is that surrogate appropriate?
  - What are the limits?
- Is your model set up correctly?
  - Is it really a classification problem?
  - one-class vs. discriminative?
  - open-world vs. closed-world?
  - correct scale of  $y$ ? Other transformation better?
- Do you use the correct controls/base class or correct 0-point (center, origin) of regression?
- What happens with out-of-spec cases (unknown class? bad measurements?)

## Validation & Optimization

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## Validation: Questions

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- Bias introduced by data acquisition procedure?
  - Labeling procedure with self-fulfilling prophecies (e.g. cluster analysis as basis for labeling, semi-supervised label generation)?
- What about borderline cases?
  - Do your *labeled* cases correctly represent them?
  - No exclusion of “difficult” cases in the reference labeling step?
- What other confounders could exist?
- What are the limits of your method?
- reading suggestions on reproducibility issues in medical research:
  - Buchen, L.: Cancer: Missing the mark. *Nature*, 2011, 471, 428 – 432
  - Begley, C. G. & Ellis, L. M.: Drug development: Raise standards for preclinical cancer research. *Nature*, 2012, 483, 531 – 533
  - Ioannidis, J. P. A.: Why Most Published Research Findings Are False, *PLoS Med* 2(8): e124
  - ...

- How robust are the predictions?
- Which factors (confounders) have most influence?
- Perturb Data
  - Repeated cross validation:  
How do predictions vary if a *few* training cases are exchanged?  
↔ stability of predictions
  - Simulate instrument related distortions:  
Measure respective drop in performance

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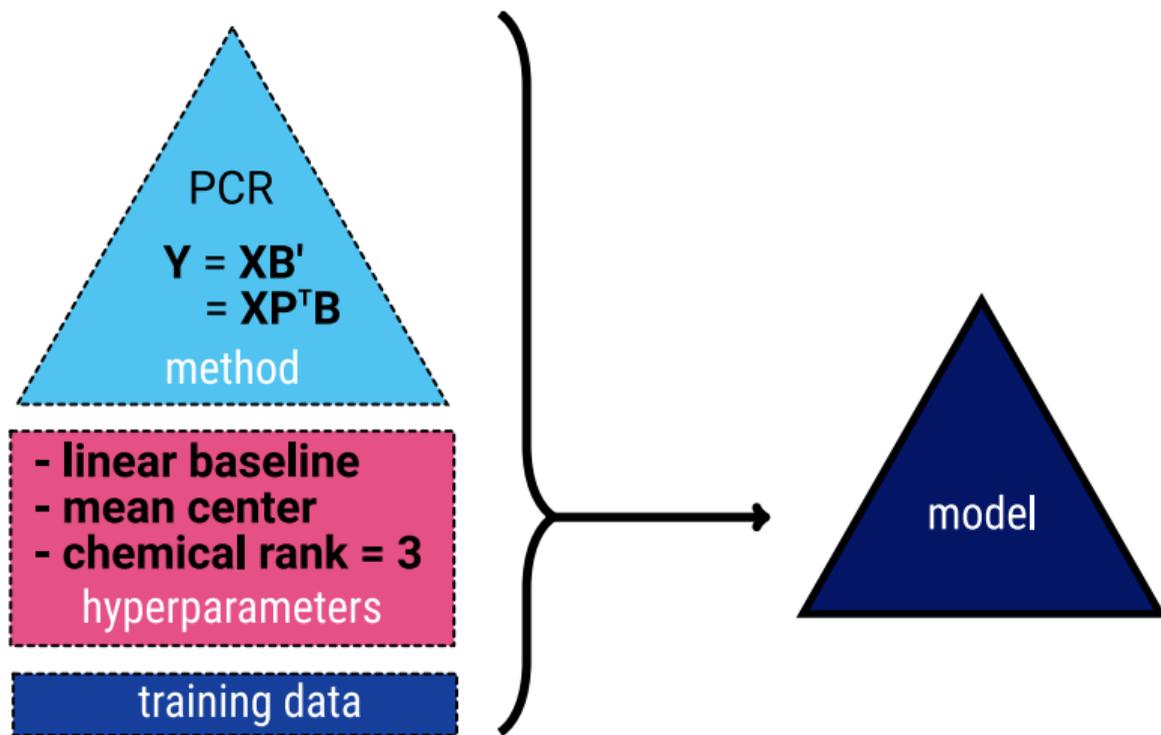
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Beleites, C. & Salzer, R.: Assessing and improving the stability of chemometric models in small sample size situations Anal Bioanal Chem, 2008, 390, 1261-1271

Sattlecker, M. et al.: Assessment of robustness and transferability of classification models built for cancer diagnostics using Raman spectroscopy J Raman Spectrosc, 2010, 897-903



# Hyperparameters



- available: PCR ( $X_{\text{train}}$ , no\_PCs, center)
- wanted: PCR\_tuned ( $X_{\text{train}}$ )

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# Data-driven Model Optimization

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training data

### Idea:

- 1 sensible range of hyperparameters
- 2 build covering this search space
- 3 measure performance
- 4 take the best

### ⇒ Optimize predictive performance

- ✓ Large variety of numerical optimizers available  
exhaustive grid search, genetic optimizers, simulated annealing, ...



# Data-driven Model Optimization

training data

validation data

### Idea:

- 1 sensible range of hyperparameters
- 2 build covering this search space
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⇒ **Optimize predictive performance**

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# Data-driven Model Optimization

training data

validation data

### Idea:

- 1 sensible range of hyperparameters
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### ⇒ Optimize predictive performance

- ✓ Large variety of numerical optimizers available  
exhaustive grid search, genetic optimizers, simulated annealing, ...
- ✗ Careful: validation data enters model building process  
⇒ need another independent set to validate the *final* model

# Data-driven Model Optimization

training data

validation data

test data

### Idea:

- 1 sensible range of hyperparameters
- 2 build covering this search space
- 3 measure performance
- 4 take the best

### ⇒ Optimize predictive performance

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# Data-driven Model Optimization

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training data

validation data

test data

- fit normal parameters (coefficients) with *training* set
- fit hyperparameters with validation set
- validate chosen model with test set

# Data-driven Model Optimization



- fit normal parameters (coefficients) with *training* set
- fit hyperparameters with validation set
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# Data-driven Model Optimization

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training data

optimization data

verification data

- fit normal parameters (coefficients) with *training* set
- fit hyperparameters with ~~validation set~~ optimization aka development set
- validate chosen model with ~~test set~~ final verification set

# Data-driven Model Optimization

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training data

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- fit normal parameters (coefficients) with *training* set
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- validate chosen model with ~~test set~~ final verification set

✓ resampling version: nested/double cross validation

# Data-driven Model Optimization

training data

optimization data

verification data

- fit normal parameters (coefficients) with *training* set
  - fit hyperparameters with ~~validation set~~ optimization aka development set
  - validate chosen model with ~~test set~~ final verification set
- ✓ resampling version: nested/double cross validation
- ✓ train ( $X$ , hyperparameters) vs. tuned\_train ( $X$ )  
– tuned training function: additional internal split for tuning

# Data-driven Model Optimization

training data

test data

- fit normal parameters (coefficients) with *training* set
  - fit hyperparameters with ~~validation set~~ optimization aka development set
  - validate chosen model with ~~test set~~ final verification set
- ✓ resampling version: nested/double cross validation
- ✓ `train (X, hyperparameters)` vs. `tuned_train (X)`
- tuned training function: additional internal split for tuning
- ✓ treat `tuned_train (X)` like any other training function

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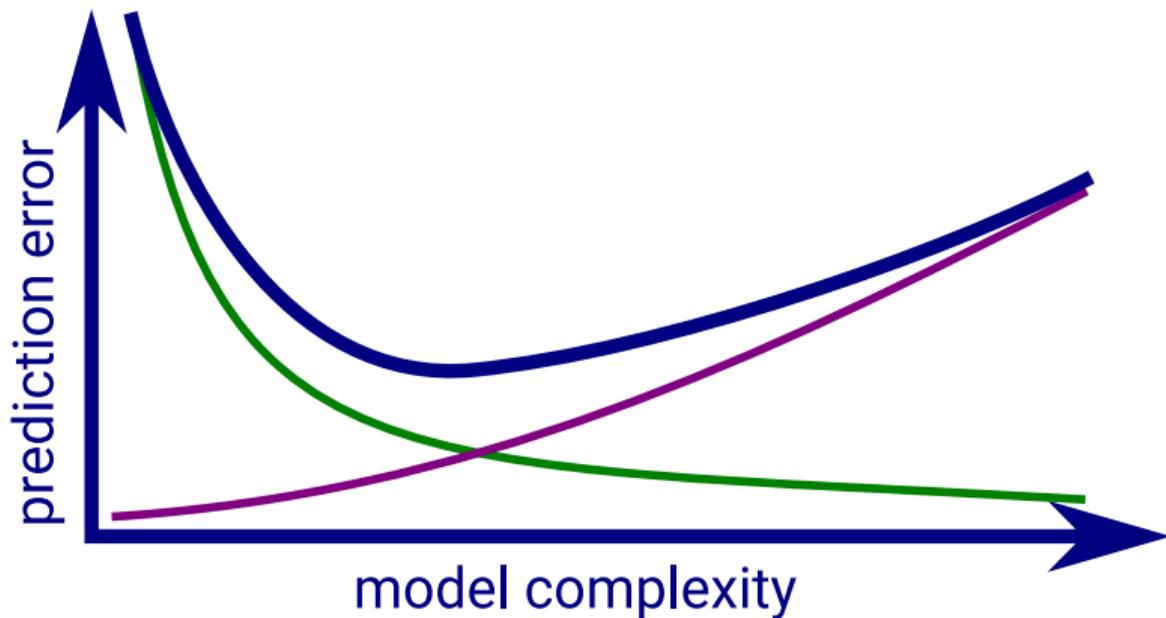
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# Grid search: Stability of Solution



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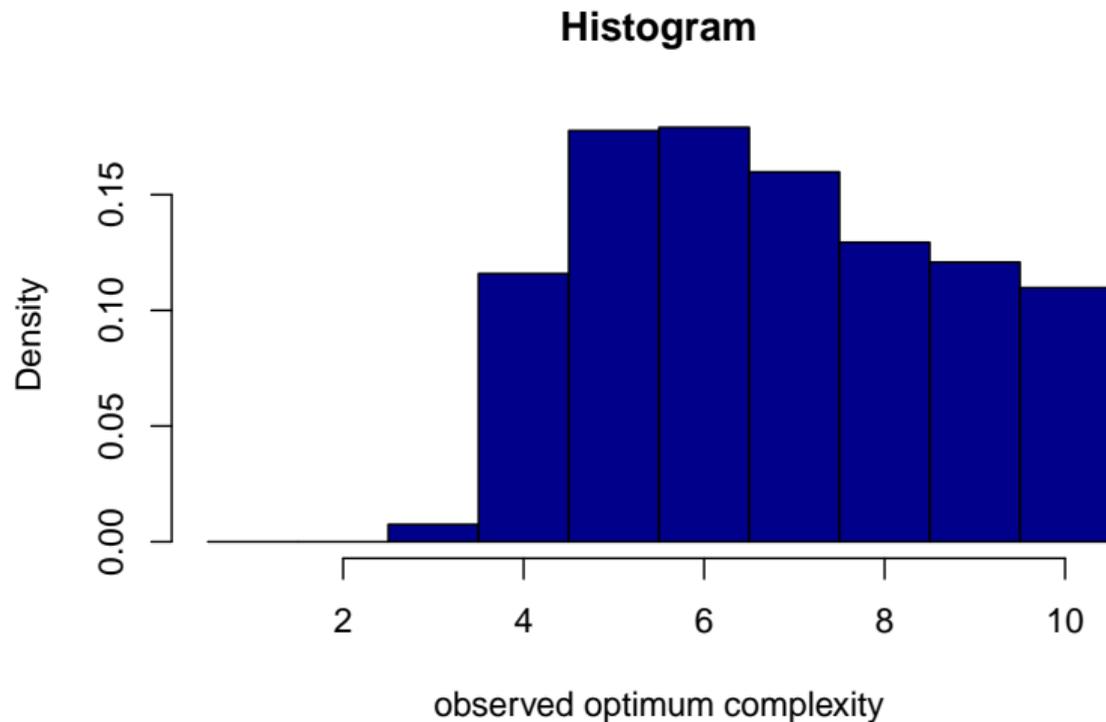
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# Grid search: Stability of Solution



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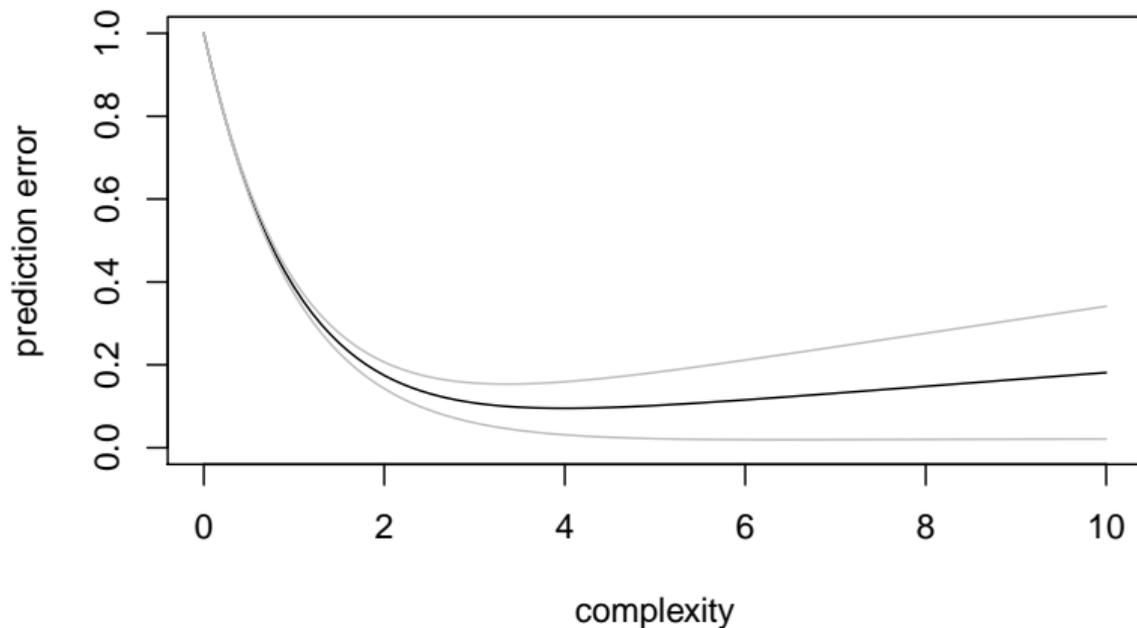
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# Grid search: Stability of Solution



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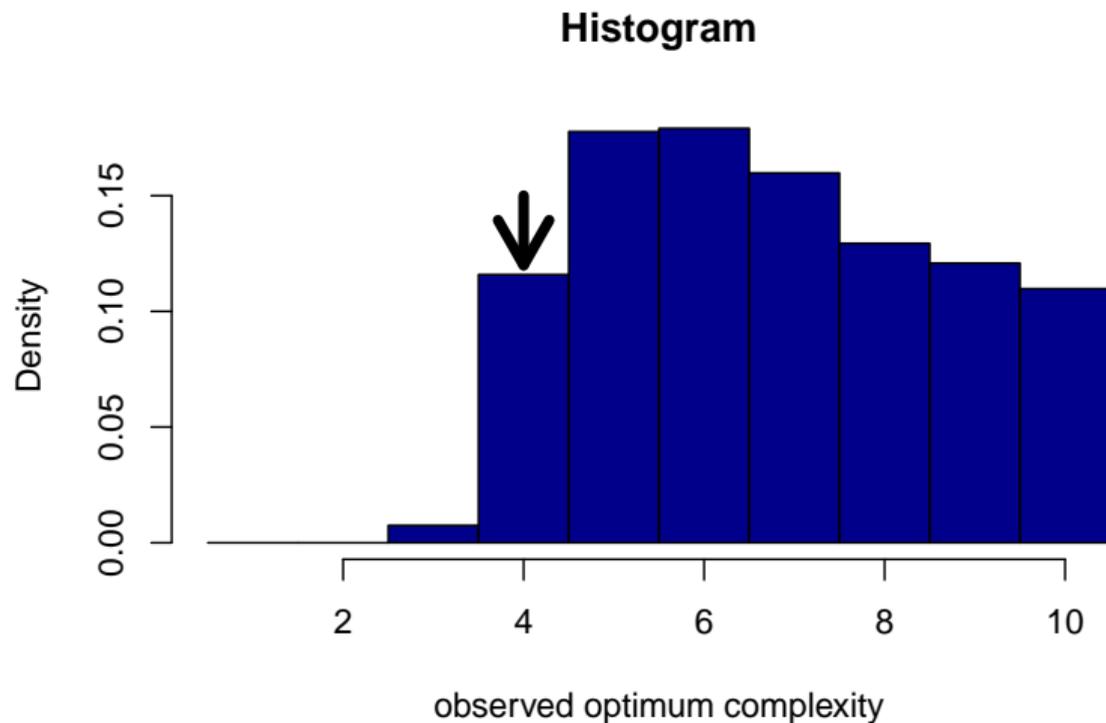
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# Grid search: Stability of Solution



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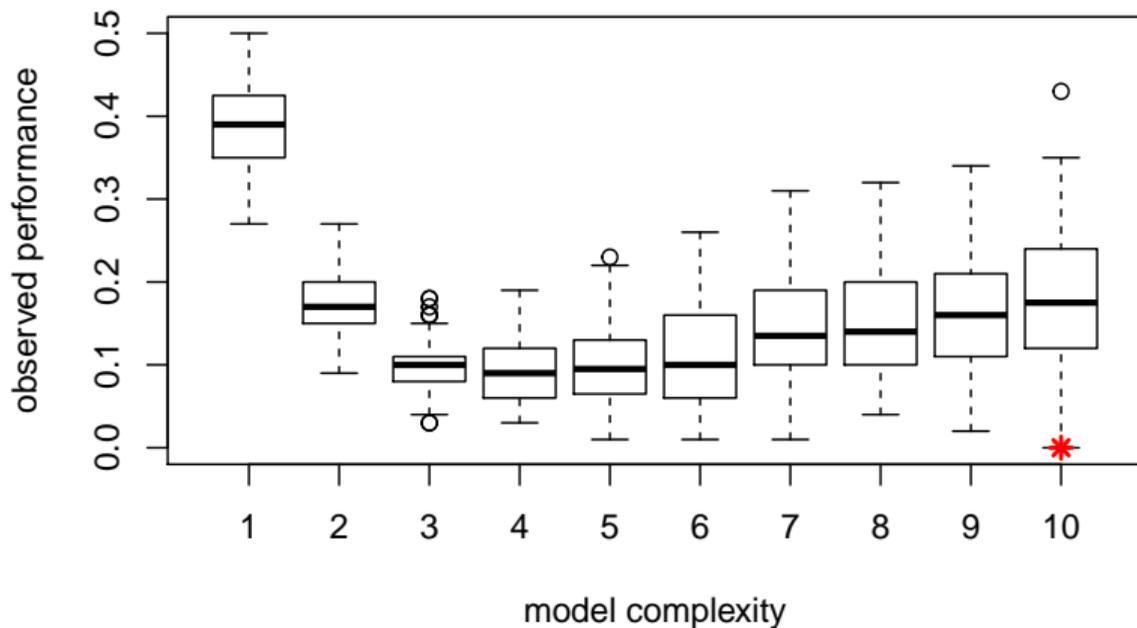
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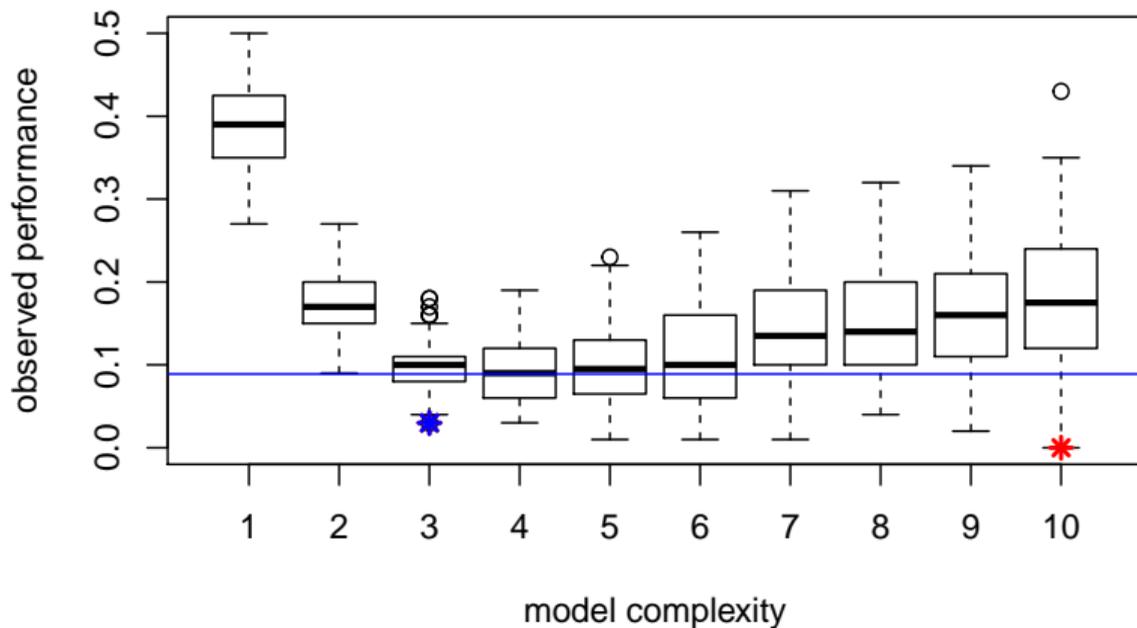
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# Grid search: Stability of Solution



# Grid search: Stability of Solution



## Summary: Validation

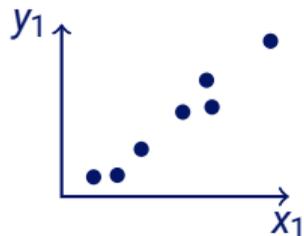
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- ✓ Think hard about your data, model, and application!
- ✓ Sample size planning: calculate from required precision of validation results possibly from preliminary experiment
  - At some point, validation studies are needed.  
Before that, use repeated cross validation or out-of-bootstrap.
- ✓ Determine independent splitting
- ✓ Check stability of predictions and – if possible – model parameters
- ✗ Resampling cannot detect drift
- ✗ Hold-out is inefficient and prone to the same errors as resampling!

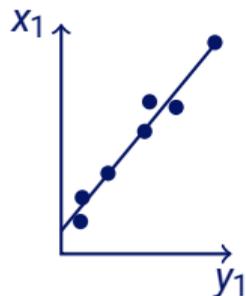
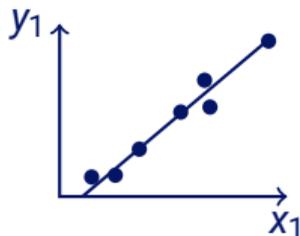
## Summary: Data-driven model Optimization

- ✓ Needs internal performance estimate plus outer independent validation
- ✗  $\rightsquigarrow$  large sample size required
- ✓ wrap optimization in `tuned_model` function
- ✓ validate output of `tuned_model` like any other model training function
- ✓ Check stability of optimization
- ✓ Use 1-sd-rule to guard against overfitting
- ✓ Class membership probability predicted: MSE (Brier's Score) has low variance and is proper scoring rule  
 $\rightsquigarrow$  suitable for optimization

# Regression



Regression



## Ordinary Regression

$$\mathbf{Y}^{(n \times m)} = \mathbf{X}^{(n \times p)} \mathbf{B}^{(p \times m)}$$

- assume error on  $y$  ( $l$ )
- ✓ causality:  $l = f(c)$
- ✓ efficient estimation of calibration line parameters

## Inverse Regression

- assume error on  $x$  ( $c$ )
- ✓ prediction:  $c = f(l)$
- ✓ efficient estimation of  $y$
- ✗ needs  $p \leq m$

# Univariate Linear Regression

---

$$\mathbf{Y}^{(n \times m)} = \mathbf{X}^{(n \times p)} \mathbf{B}^{(p \times m)}$$

<b>y</b>	<b>x</b>
1	2
2	3
3	4
4	5
5	6

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# Multivariate Linear Regression

$$\mathbf{Y}^{(n \times m)} = \mathbf{X}^{(n \times p)} \mathbf{B}^{(p \times m)}$$

<b>y</b>	<b>x<sub>1</sub></b>	<b>x<sub>2</sub></b>	<b>x<sub>3</sub></b>
1	2	7	3
2	3	5	5
3	4	3	-2
4	5	1	7
5	6	-1	0

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# Linear Models: Polynomial Features

$$\mathbf{Y}^{(n \times m)} = \mathbf{X}^{(n \times p)} \mathbf{B}^{(p \times m)}$$

y	x
1	2
2	3
3	4
4	5
5	6

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# Linear Models: Polynomial Features

$$\mathbf{Y}^{(n \times m)} = \mathbf{X}^{(n \times p)} \mathbf{B}^{(p \times m)}$$

y	$x^0$	$x^1$	$x^2$
1	1	2	4
2	1	3	9
3	1	4	16
4	1	5	25
5	1	6	36

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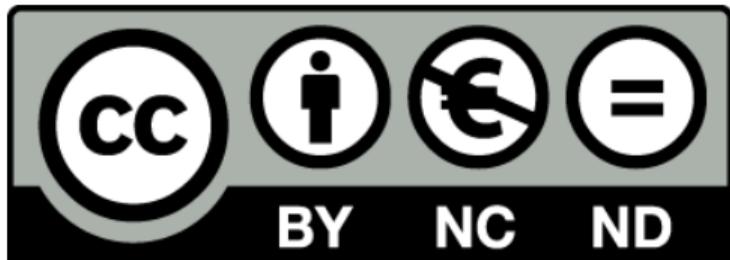


## Questions?

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Please contact me ([Claudia.Beleites@chemometrix.gmbh](mailto:Claudia.Beleites@chemometrix.gmbh)) if you

- have questions, or
- want to reuse these slides.



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