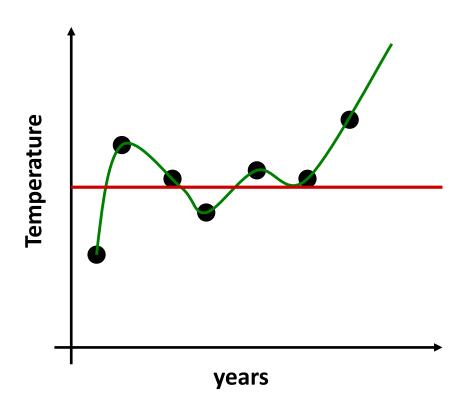
Cross validation, training and evaluation of data driven prediction methods

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Data driven method training

- A prediction method contains a very large set of parameters
 - A matrix for predicting binding for 9meric peptides has 9x20=180 weights
- Over fitting is a problem



Evaluation of predictive performance

- Train PSSM on raw data
 - No pseudo counts, No sequence weighting
 - Fit 9*20 (=180) parameters to 9 (*10 = 90) data points
- Evaluate on training data

$$-PCC = 0.97$$

- -AUC = 1.0
- Close to a perfect prediction method

ALAKAAAAM AT.AKAAAAN ALAKAAAAR **ALAKAAAAT Binders ALAKAAAAV GMNERPILT GILGFVFTM** TLNAWVKVV KLNEPVLLL **AVVPFIVSV MRSGRVHAV** Binders VRFNIDETP ANYIGODGL **AELCGDPGD QTRAVADGK GRPVPAAHP MTAQWWLDA FARGVVHVI** LORELTRLO

Evaluation of predictive performance

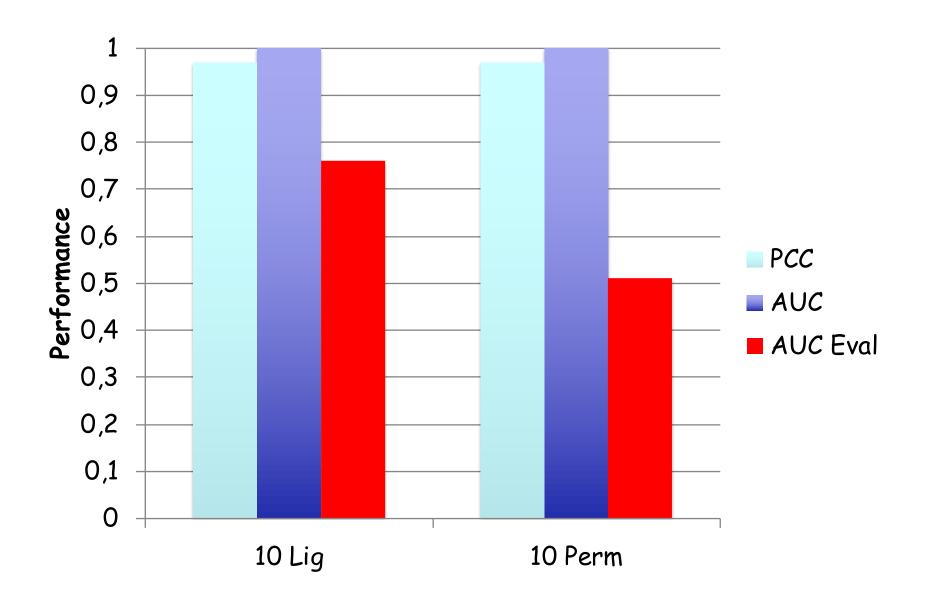
- Train PSSM on <u>Permuted</u> (random) data
 - No pseudo counts, No sequence weighting
 - Fit 9*20 parameters to 9*10 data points
- Evaluate on training data
 - -PCC = 0.97
 - -AUC = 1.0
- Close to a perfect prediction method AND
- Same performance as on the original data

3inders

Jone Binders

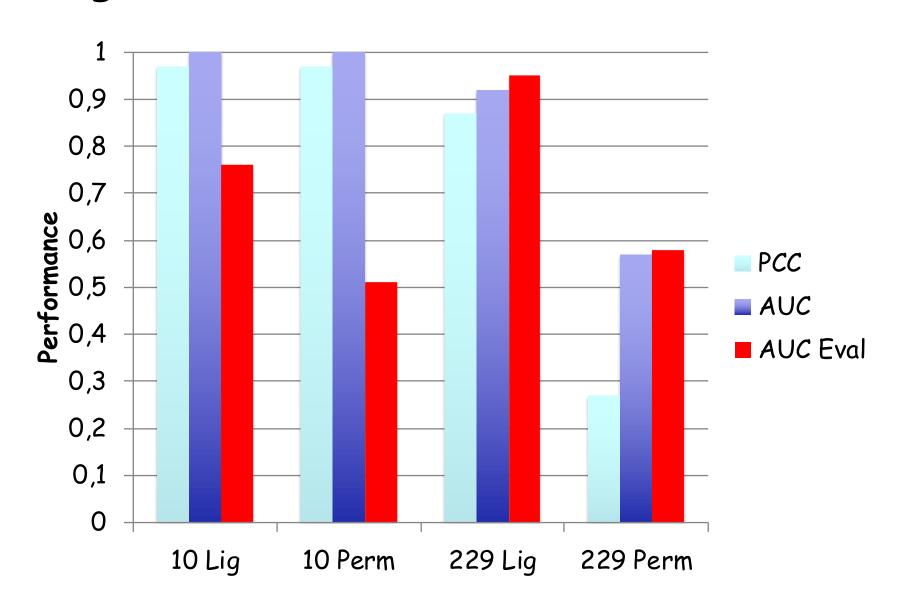
AAAMAAKLA
AAKNLAAAA
AKALAAAAR
AAAAKLATA
ALAKAVAAA
IPELMRTNG
FIMGVFTGL
NVTKVVAWL
LEPLNLVLK
VAVIVSVPF
MRSGRVHAV
VRFNIDETP

WRSGRVHAV
VRFNIDETP
ANYIGQDGL
AELCGDPGD
QTRAVADGK
GRPVPAAHP
MTAQWWLDA
FARGVVHVI
LQRELTRLQ
AVAEEMTKS



Repeat on large training data (229 ligands)





FLAFFSNGV

FLAFFSNGV

WLGNHGFEV

TLNAWVKVV

LLATSIFKL

LLSKNTFYL

KVGNCDETV

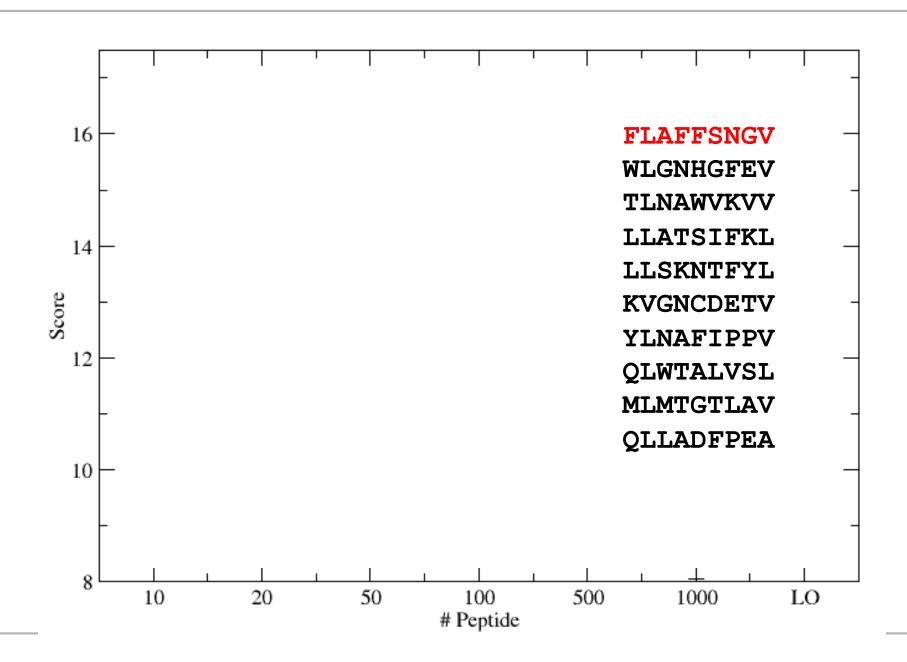
YLNAFIPPV

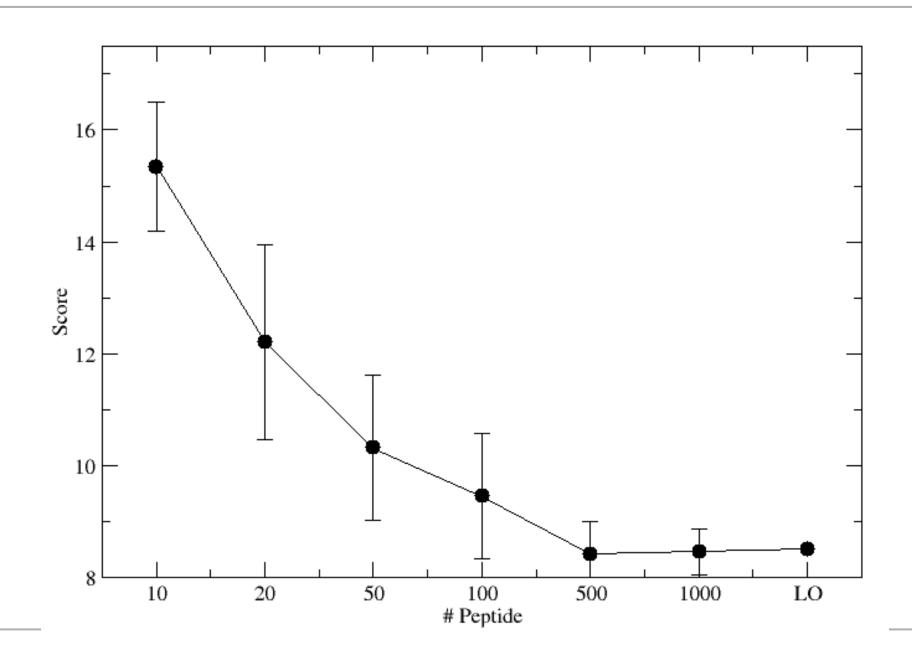
OLWTALVSL

MLMTGTLAV

QLLADFPEA

FLAFFSNGV VLMEAQQGI ILLLDQVLV KMYEYVFKG HLMRDPALL WLVHKOWFL ALAPSTMKI MLLTFLTSL FLIVSLCPT ITWQVPFSV RMPAVTDLV ALYSYASAK YFLRRLALV FLLDYEGTL FLITGVFDI LLVLCVTQV MTSELAALI MLLHVGIPL GLIIISIFL IVYGRSNAI GLYEAIEEC SLSHYFTLV GLYYLTTEV AQSDFMSWV KLFFAKCLV VLWEGGHDL YLLNYAGRI RLEELLPAV VLQAGFFLL AIDDFCLFA KVVSLVILA LLVFACSAV TLKDAMLQL GLFQEAYPL YQLGDYFFV GMVIACLLV MSDIFHALV MVVKVNAAL FMTALVLSL WLSTYAVRI GMRDVSFEL FLGFLATAG ILAKFLHWL IVLGNPVFL **QLPLESDAV SLYPPCLFK MTPSPFYTV LLVAPMPTA** KVGNCDETV RIFPATHYV IIDQVPFSV YLNKIQNSL ILYQVPFSV YLMKDKLNI AIMEKNIML LLNNSLGSV GLKISLCGI ALGLGIVSL MMCPFLFLM FMFNELLAL WLETELVFV ALYWALMES GLDPTGVAV GMLPVCPLI WQDGGWQSV LLIEGIFFI SILNTLRFL GLSLSLCTL VMLIGIEIL RLNKVISEL KVEKYLPEV YLVAYOATV SVMDPLIYA IMSSFEFQV FTLVATVSI ILLVAVSFV GMFGGCFAA RLLDDTPEV SLDSLVHLL LVLOAGFFL VLAGYGAGI VILWFSFGA VLNTLMFMV FLQGAKWYL



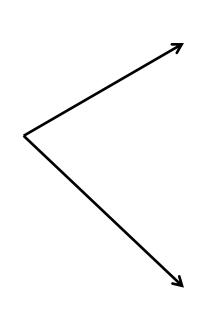


Gibbs clustering (multiple specificities)



Multiple motifs!

SLFIGLKGDIRESTV
DGEEEVQLIAAVPGK
VFRLKGGAPIKGVTF
SFSCIAIGIITLYLG
IDQVTIAGAKLRSLN
WIQKETLVTFKNPHAKKQDV
KMLLDNINTPEGIIP
ELLEFHYYLSSKLNK
LNKFISPKSVAGRFA
ESLHNPYPDYHWLRT
NKVKSLRILNTRRKL
MMGMFNMLSTVLGVS
AKSSPAYPSVLGQTI
RHLIFCHSKKKCDELAAK



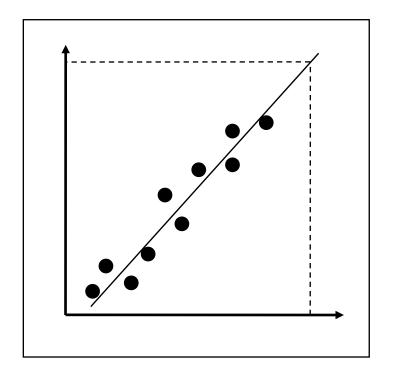
Cluster 1

```
----SLFIGLKGDIRESTV--
--DGEEEVQLIAAVPGK----
----VFRLKGGAPIKGVTF
---SFSCIAIGIITLYLG---
----IDQVTIAGAKLRSLN--
WIQKETLVTFKNPHAKKQDV-
-----KMLLDNINTPEGIIP
```

Cluster 2

```
--ELLEFHYYLSSKLNK----
-----LNKFISPKSVAGRFA
ESLHNPYPDYHWLRT-----
-NKVKSLRILNTRRKL-----
--MMGMFNMLSTVLGVS----
AKSSPAYPSVLGQTI-----
--RHLIFCHSKKKCDELAAK-
```

Always



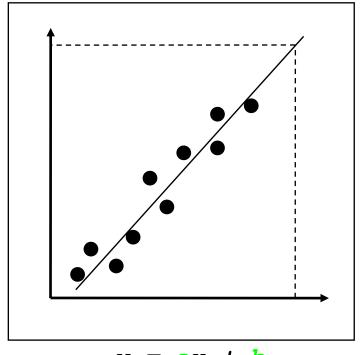
Observations (training data): a set of x values (input) and y values (output).

Model: y = ax + b (2 parameters, which are estimated from the training data)

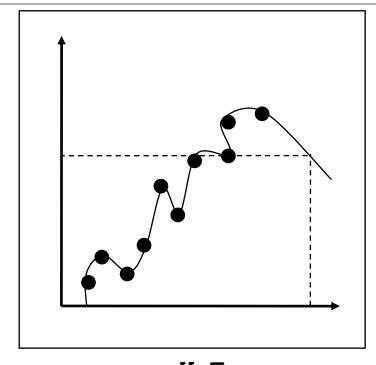
Prediction: Use the model to calculate a y value for a new x value

Note: the model does not fit the observations exactly. Can we do better than this?

Overfitting



y = ax + b2 parameter model Good description, poor fit



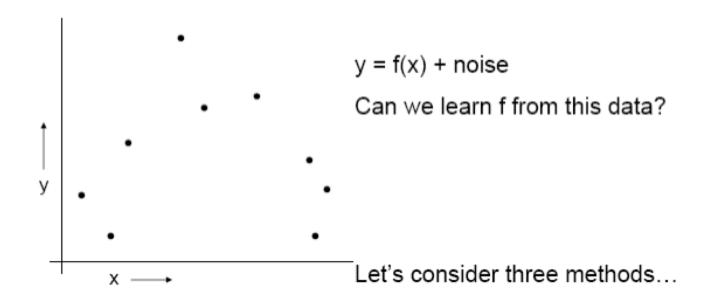
 $y = ax^6 + bx^5 + cx^4 + dx^3 + ex^2 + fx + g$ 7 parameter model
Poor description, good fit

Note: It is not interesting that a model can fit its observations (training data) exactly.

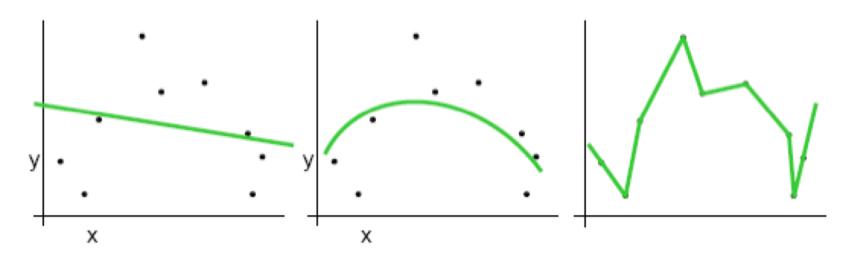
To function as a prediction method, a model must be able to generalize, i.e. produce sensible output on new data.

How to estimate parameters for prediction?

A Regression Problem



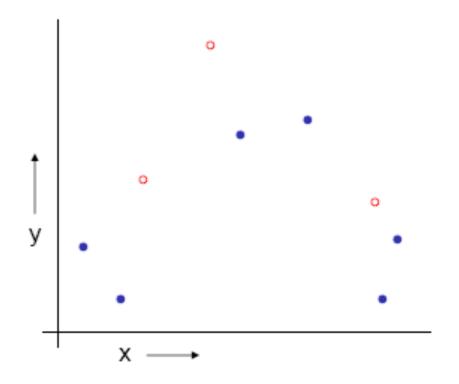
Which is best?



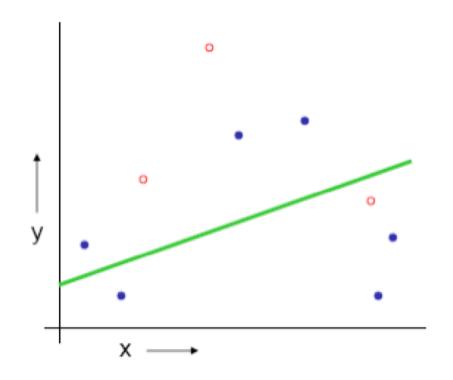
Linear Regression

Quadratic Regression

Join-the-dots

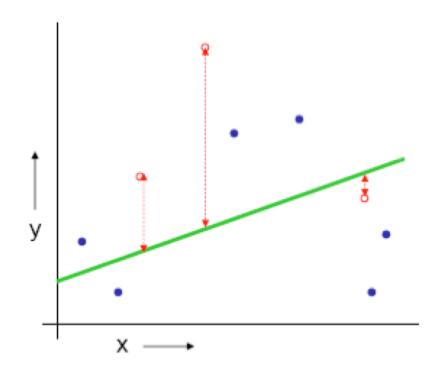


- 1. Randomly choose 30% of the data to be in a test set
- The remainder is a training set



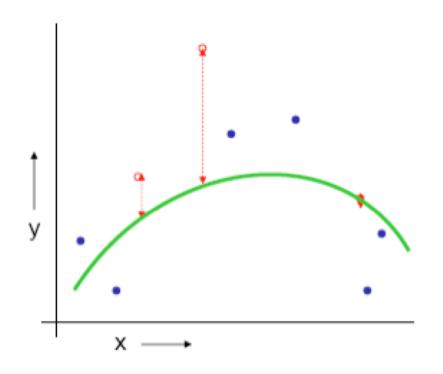
(Linear regression example)

- Randomly choose
 of the data to be in a test set
- 2. The remainder is a training set
- Perform your regression on the training set



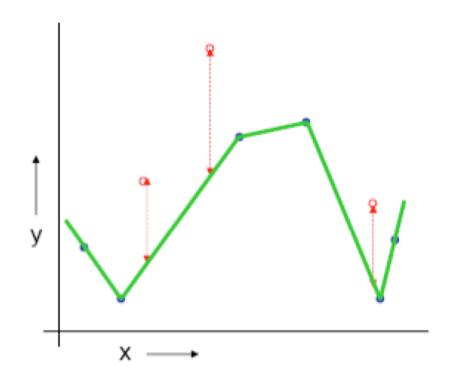
(Linear regression example)
Mean Squared Error = 2.4

- 1. Randomly choose 30% of the data to be in a test set
- The remainder is a training set
- 3. Perform your regression on the training set
- 4. Estimate your future performance with the test set



(Quadratic regression example) 4. Estimate your future Mean Squared Error = 0.9

- 1. Randomly choose 30% of the data to be in a test set
- 2. The remainder is a training set
- Perform your regression on the training set
- performance with the test set



(Join the dots example)
Mean Squared Error = 2.2

- 1. Randomly choose 30% of the data to be in a test set
- The remainder is a training set
- 3. Perform your regression on the training set
- 4. Estimate your future performance with the test set

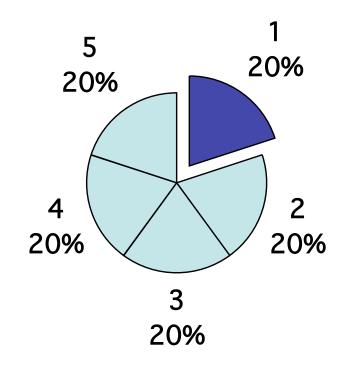
So quadratic function is best

How to deal with overfitting? Cross validation

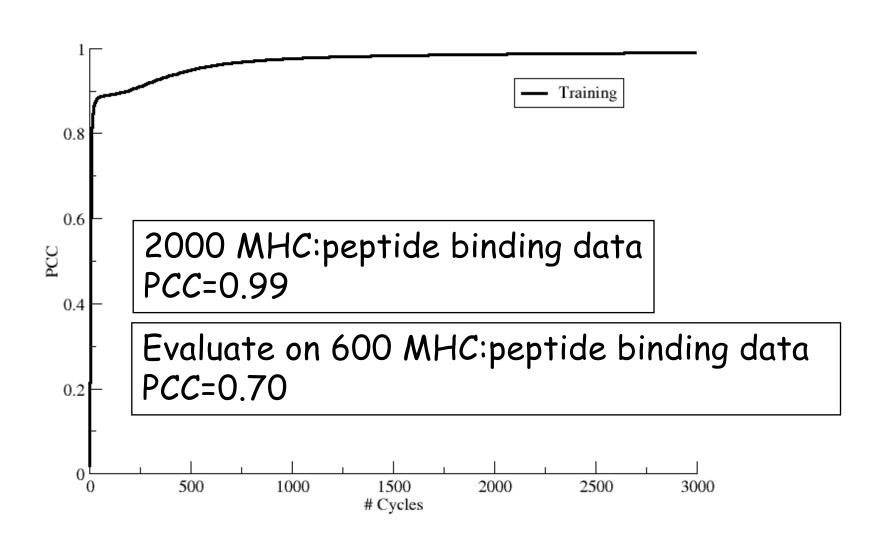
Cross validation

Train on 4/5 of data
Test/evaluate on 1/5
=>

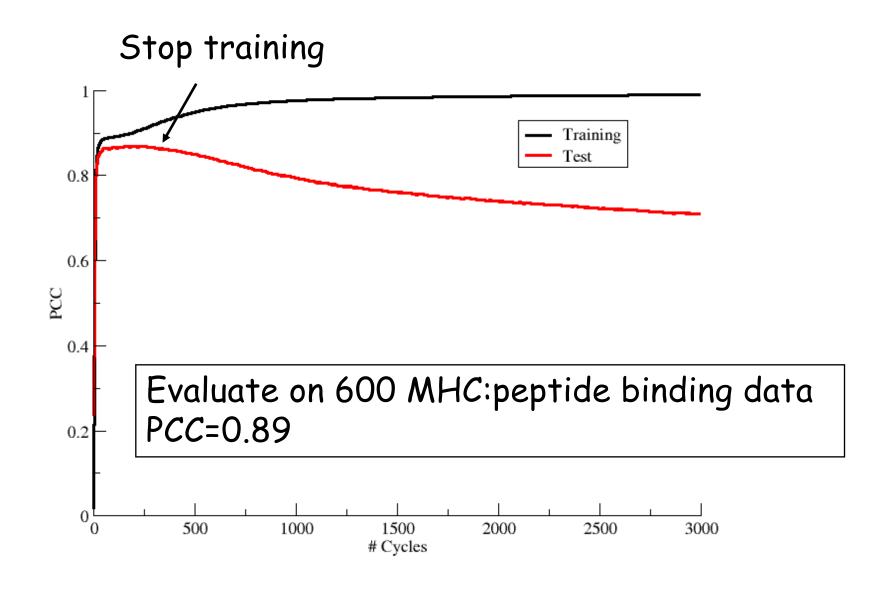
Produce 5 different methods each with a different prediction focus



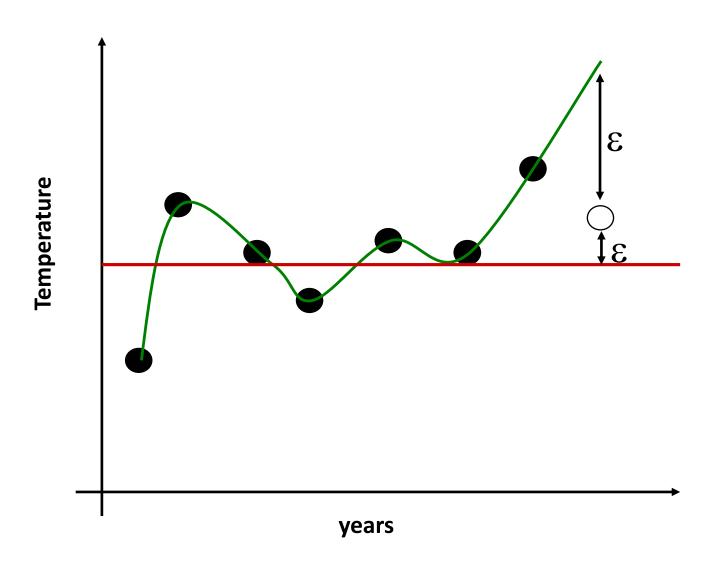
Model over-fitting



Model over-fitting (early stopping)

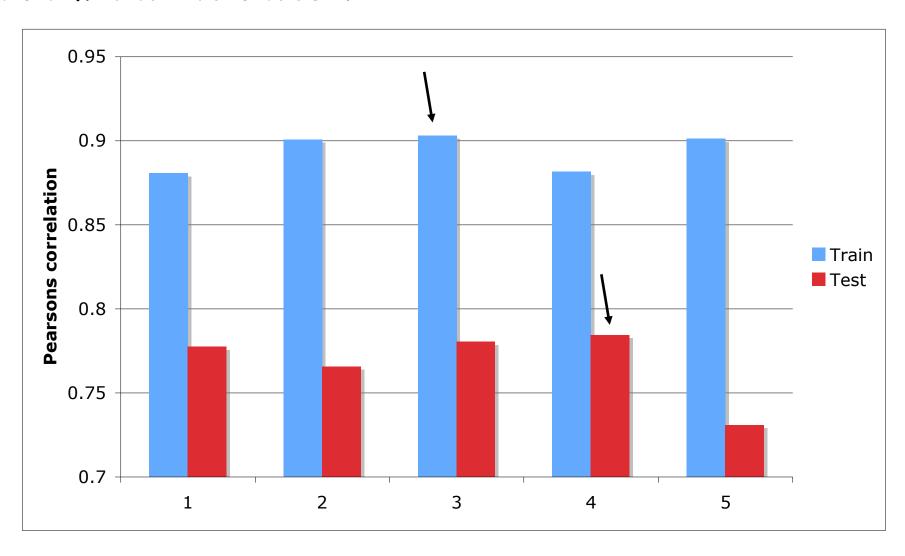


What is going on?

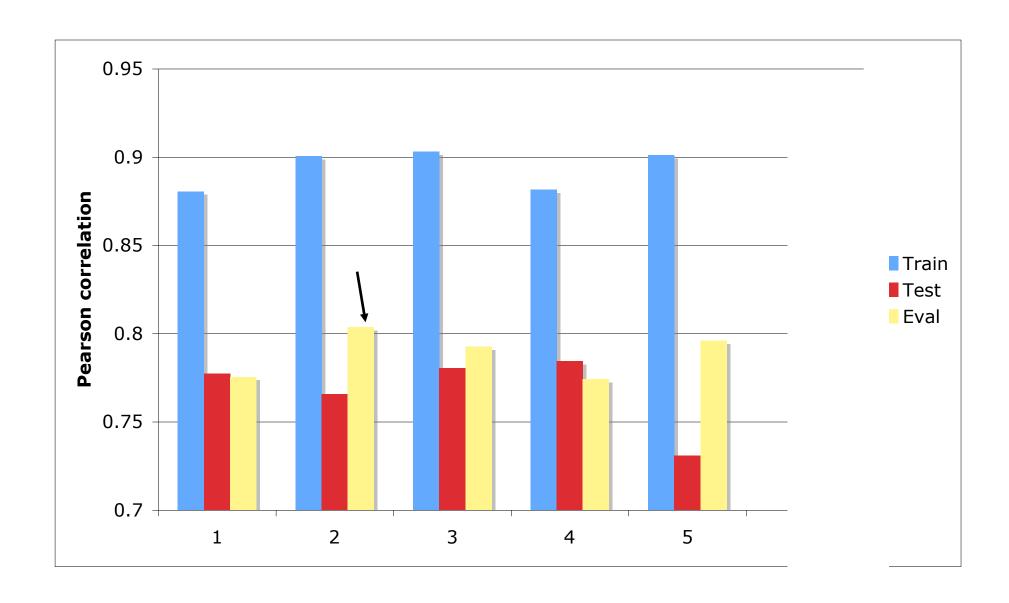


5 fold training

Which method to choose?



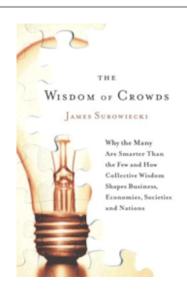
5 fold training



The Wisdom of the Crowds

• The Wisdom of Crowds. Why the Many are Smarter than the Few. James Surowiecki

One day in the fall of 1906, the British scientist Fracis
Galton left his home and headed for a country fair... He
believed that only a very few people had the
characteristics necessary to keep societies healthy. He
had devoted much of his career to measuring those
characteristics, in fact, in order to prove that the vast
majority of people did not have them. ... Galton came
across a weight-judging competition... Eight hundred people
tried their luck. They were a diverse lot, butchers,
farmers, clerks and many other no-experts... The crowd
had guessed ... 1.197 pounds, the ox weighted 1.198

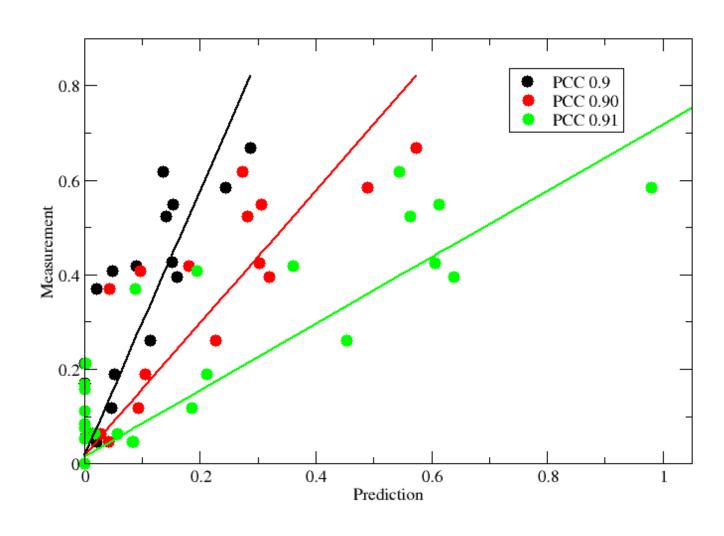


The wisdom of the crowd!

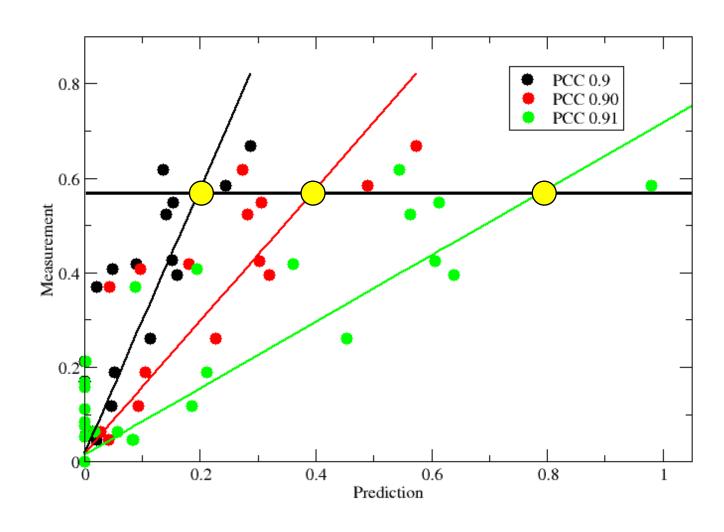
- The highest scoring hit will often be wrong
 - Not one single prediction method is consistently best
- Many prediction methods will have the correct fold among the top 10-20 hits
- If many different prediction methods all have a common fold among the top hits, this fold is probably correct

- Use cross validation
- Evaluate on concatenated data and <u>not</u> as an average over each cross-validated performance

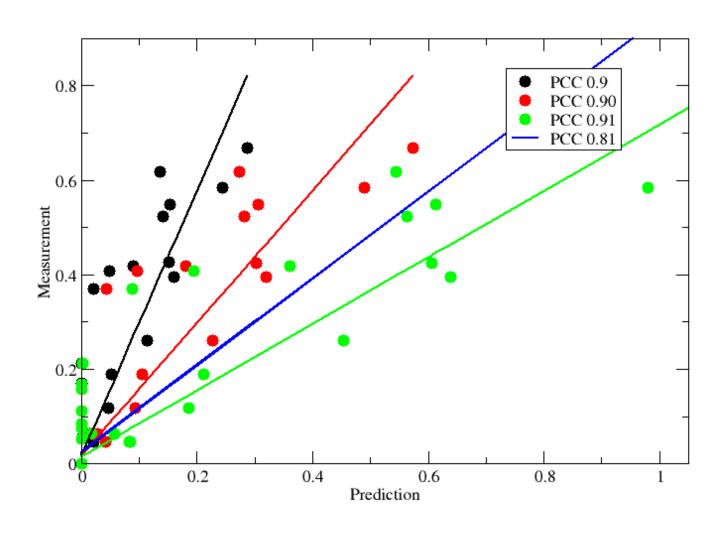
Method evaluation



Method evaluation



Method evaluation



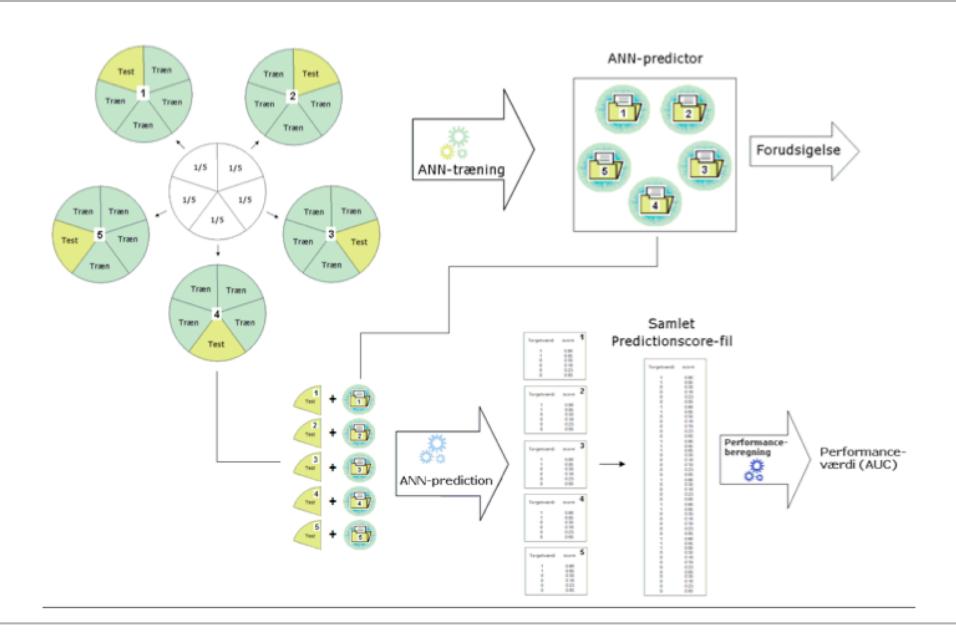
How many folds?

- Cross validation is always good!, but how many folds?
 - Few folds -> small training data sets
 - Many folds -> small test data sets
- 560 peptides for training
 - 50 fold (10 peptides per test set, few data to stop training)
 - 2 fold (280 peptides per test set, few data to train)
 - 5 fold (110 peptide per test set, 450 per training set)

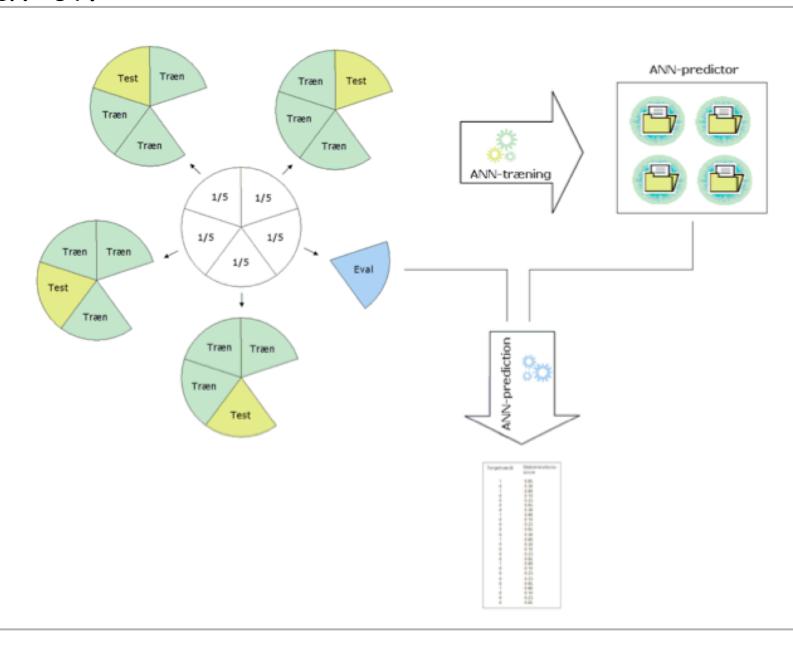
Problems with 5fold cross validation

- Use test set to stop training, and test set performance to evaluate training
 - Over-fitting?
- If test set is small, Yes
- If test set is large, No
- Confirm using "true" 5 fold cross validation
 - 1/5 for evaluation
 - 4/5 for 4 fold cross-validation

Conventional 5 fold cross validation



"Nested (or true)" 5 fold cross validation



When to be careful

- When data is scarce, the performance obtained used "conventional" versus "nested" cross validation can be very large
- When data is abundant the difference is in general small

Training/evaluation procedure

- Define method
- Select data
- Deal with data redundancy
 - In method (sequence weighting)
 - In data (Hobohm)
- Deal with over-fitting either
 - in method (SMM regulation term) or
 - in training (stop fitting on test set performance)
- Evaluate method using cross-validation