For written notes on this lecture, please read chapter 3 of *The Practical Bioinformatician*,

# CS2220: Introduction to Computational Biology Unit 1a: Essence of Knowledge Discovery

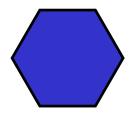
Wong Limsoon



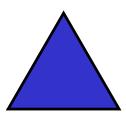
# What is knowledge discovery?

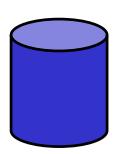


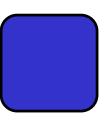
Jonathan's blocks



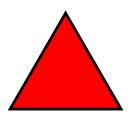


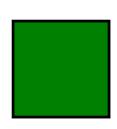


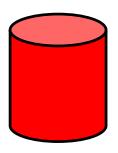


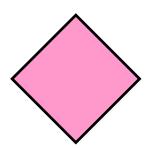


Jessica's blocks









Whose block is this?

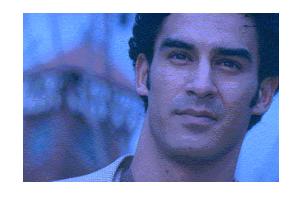
Jonathan's rules Jessica's rules : Blue or Circle

: All the rest

# What is knowledge discovery?











Question: Can you explain how?

## Key steps



- Training data gathering
- Feature generation
  - k-grams, colour, texture, domain know-how, ...
- Feature selection
  - Entropy, χ2, CFS, t-test, domain know-how...
- Feature integration
  - SVM, ANN, PCL, CART, C4.5, kNN, ...

Some classifiers / machine learning methods

## What is accuracy?





#### What is accuracy?

	predicted	predicted
	as positive	as negative
positive	TP	FN
negative	FP	TN

Accuracy = 
$$\frac{\text{No. of correct predictions}}{\text{No. of predictions}}$$
$$= \frac{\text{TP + TN}}{\text{TP + TN + FP + FN}}$$

## Examples (Balanced population)



classifier	TP	TN	FP	FN	Accuracy
Α	25	25	25	25	50%
В	50	25	25	0	75%
С	25	50	0	25	75%
D	37	37	13	13	74%

- Clearly, B, C, D are all better than A
- Is B better than C, D?
- Is C better than B, D?
- Is D better than B, C?

Accuracy may not tell the whole story

# Examples (Unbalanced population National University of Singapore

classifier	TP	TN	FP	FN	Accuracy
Α	25	75	75	25	50%
В	0	150	0	50	75%
С	50	0	150	0	25%
D	30	100	50	20	65%

- Clearly, D is better than A
- Is B better than A, C, D?

Exercise: What is B's Prediction strategy?

High accuracy is meaningless if population is unbalanced



# What is sensitivity (aka recall)?

	predicted	predicted
	as positive	as negative
positive	TP	FN
negative	FP	TN

Sensitivity = 
$$\frac{\text{No. of correct positive predictions}}{\text{No. of positives}}$$

$$= \frac{\text{TP}}{\text{TP} + \text{FN}}$$

Sometimes sensitivity wrt negatives is termed specificity

Exercise: Write down the formula for specificity



#### What is precision?

	predicted	predicted		
	as positive	as negative		
positive	TP	FN		
negative	FP	TN		

Precision = wrt positives

No. of correct positive predictions

No. of positives predictions

$$\frac{TP}{TP + FP}$$

# Unbalanced population revisited National University of Singapore

classifier	TP	TN	FP	FN	Accuracy	Sensitivity	Precision
A	25	75	75	25	50%	50%	25%
В	0	150	0	50	75%		
С	50	0	150	0	25%		
D	30	100	50	20	65%	60%	38%

- What are the sensitivity and precision of B and C?
- Is B better than A, C, D?



### Abstract model of a classifier



- Given a test sample S
- Compute scores p(S), n(S)
- Predict S as negative if p(S) / n(S) < t</li>
- Predict S as positive if  $p(S)/n(S) \ge t$

t is the decision threshold of the classifier

changing *t* affects the recall and precision, and hence accuracy, of the classifier

## Example



ន	P(S)	N(S)	Actual	Actual Predicted	
			Class	Class	Class
				0 t = 3	0 t = 2
2	0.961252	0.038748	P	Р	P
3	0.435302	0.564698	N	N	N
6	0.691596	0.308404	P	N	P
7	0.180885	0.819115	N	N	N
8	0.814909	0.185091	P	P	P
10	0.887220	0.112780	P	P	P
			accuracy	5 / 6	6/6
			recall	3 / 4	4/4
			precision	3 / 3	4/4

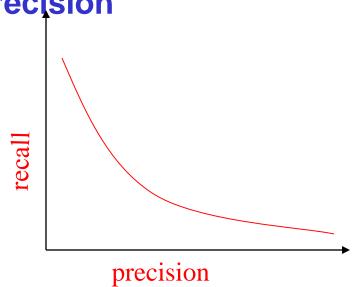
#### Recall that ...

- Predict S as negative if p(S) / n(S) < t
- Predict S as positive if  $p(S) / n(S) \ge t$

#### Precision-recall trade-off



- A predicts better than
   B if A has better recall
   and precision than B
- There is a trade-off between recall and precision



- In some apps, once you reach satisfactory precision, you optimize for recall
- In some apps, once you reach satisfactory recall, you optimize for precision

# Comparing prediction performance National University of Singapore

- Accuracy is the obvious measure
  - But it conveys the right intuition only when the positive and negative populations are roughly equal in size
- Recall and precision together form a better measure
  - But what do you do when A has better recall than B and B has better precision than A?

# F-measure (Used in info extraction NUS National University of Singapore

Take the harmonic mean of recall and precision

$$F = \frac{2 * recall * precision}{recall + precision}$$
 (wrt positives)

classifier	TP	TN	FP	FN	Accuracy	F-measure
Α	25	75	75	25	50%	33%
В	0	150	0	50	75%	undefined
С	50	0	150	0	25%	40%
D	30	100	50	20	65%	46%

Does not accord with intuition:

C predicts everything as +ve, but still rated better than A

#### Adjusted accuracy



#### Weigh by the importance of the classes

Adjusted accuracy = 
$$\alpha$$
 \* Sensitivity +  $\beta$  \* Specificity where  $\alpha + \beta = 1$  typically,  $\alpha = \beta = 0.5$ 

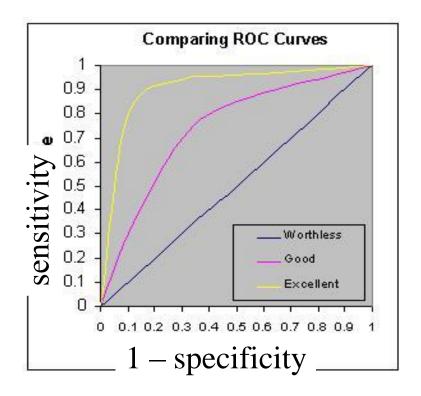
classifier	TP	TN	FP	FN	Accuracy	Adj Accuracy
Α	25	75	75	25	50%	50%
В	0	150	0	50	75%	50%
С	50	0	150	0	25%	50%
D	30	100	50	20	65%	63%

#### **ROC** curves



- By changing t, we get a range of sensitivities and specificities of a classifier
- A predicts better than B if A has better sensitivities than B at most specificities
- Leads to ROC curve that plots sensitivity vs. (1 – specificity)

 Then the larger the area under the ROC curve, the better



## Food for thought



- You have a classifier. On a test set having 20%
   +ve and 80% -ve cases, the classifier's recall and precision are both 80%
- Suppose you test it on a new test set having 80% +ve and 20% -ve cases. What do you expect its accuracy to be?
- You may assume that the +ve (resp. –ve) cases in both test sets are equally sufficiently representative of the +ve (resp. –ve) real-world population
- What lesson have you learned?

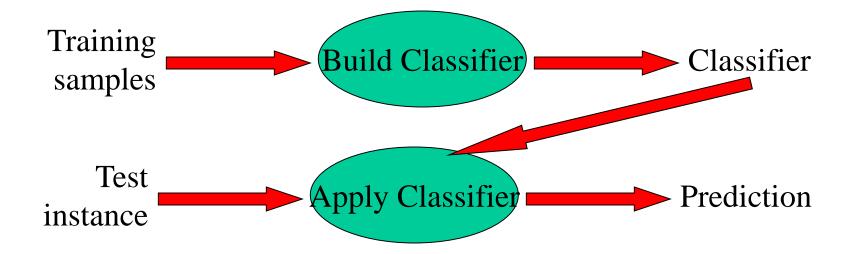


#### What is cross validation?



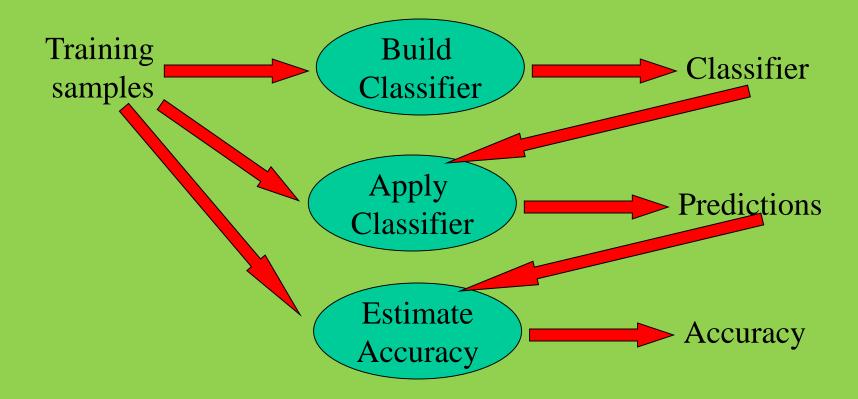
#### Construction of a classifier





## Estimate accuracy: Wrong way





Why is this way of estimating accuracy wrong?

Exercise #3

#### Recall ...



#### ...the abstract model of a classifier

- Given a test sample S
- Compute scores p(S), n(S)
- Predict S as negative if p(S) / n(S) < t</li>
- Predict S as positive if p(S) / n(S) ≥ t

t is the decision threshold of the classifier

# K-nearest neighbour classifier (k-New National University of Singapore

- Given a sample S, find the k observations S<sub>i</sub> in the known data that are "closest" to it, and take majority vote of their responses
- Assume S is well approximated by its neighbours

$$p(S) = \sum_{S_i \in N_k(S) \cap D^P} 1 \qquad n(S) = \sum_{S_i \in N_k(S) \cap D^N} 1$$

where  $N_k(S)$  is the neighbourhood of S defined by the k nearest samples to it.

Assume distance between samples is Euclidean distance for now

## Illustration of kNN (k=8)



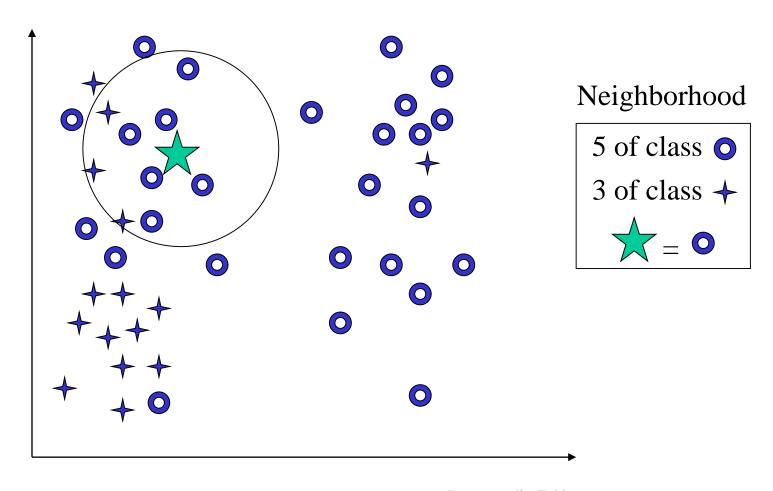
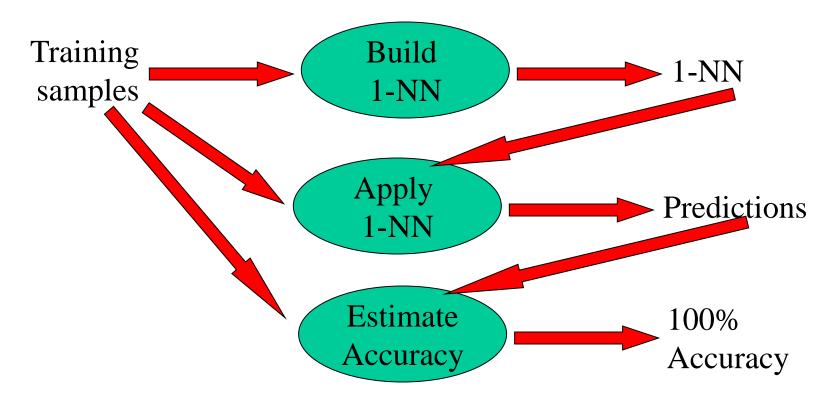


Image credit: Zaki

## Estimate accuracy: Wrong way

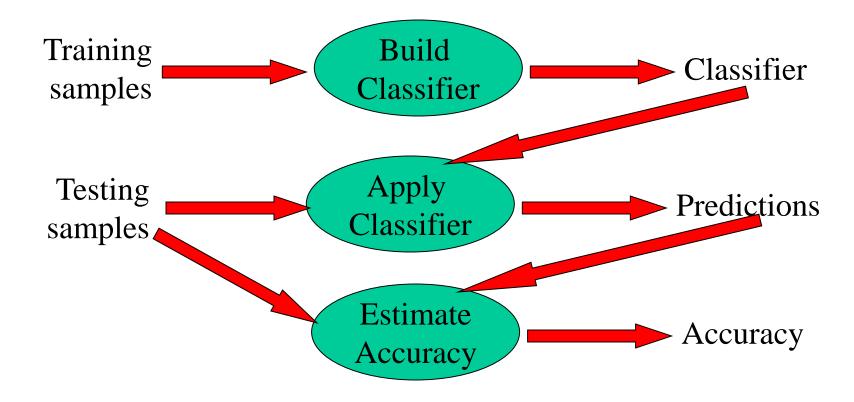




For sure k-NN (k = 1) has 100% accuracy (Why?) in the "accuracy estimation" procedure above. Does this accuracy generalize to new test instances?

## Estimate accuracy: Right way





Testing samples are NOT to be used during "Build Classifier"

## How many training and testing sample

- No fixed ratio between training and testing samples; but typically 2:1 ratio
- Proportion of instances of different classes in testing samples should be similar to proportion in the real world, and preferably also to proportion in the training samples
- What if there are insufficient samples to reserve 1/3 for testing?

#### **Cross validation**

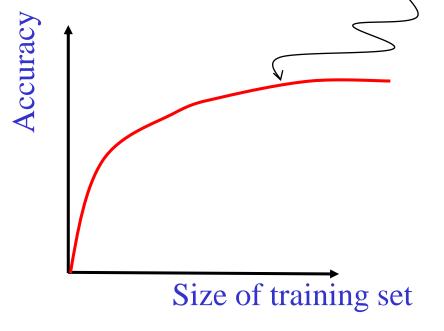


- 2.Train 3.Train 4.Train 5.Train 2.Test 3.Train 4.Train 5.Train 1.Train 1.Train 2.Train 3. Test 4. Train 5. Train 1.Train 2.Train 3.Train 4.Test 5.Train 1.Train 2.Train 3.Train 4.Train
- Divide samples into k roughly equal parts
- Each part has similar proportion of samples from different classes
- Use each part to test other parts
- Total up accuracy

#### How many fold?



 If samples are divided into k parts, we call this k-fold cross validation



#### Choose k so that

- the k-fold cross
   validation accuracy
   does not change
   much from k-1 fold
- each part within the kfold cross validation has similar accuracy
- k = 5 or 10 are popular choices for k

#### Bias and variance



Suppose a butcher weighs a steak with his thumb on the scale. That causes an error in the measurement, but little has been left to chance. Take another example. Suppose a drapery store uses a cloth tape measure which has stretched from 36 inches to 37 inches in length. Every "yard" of cloth they sell to a customer has an extra inch tacked onto it. This isn't a chance error, because it always works for the customer. The butcher's thumb and the stretched tape are two examples of bias, or systematic error.

Bias affects all measurements the same way, pushing them in the same direction. Chance errors change from measurement to measurement, sometimes up and sometimes down.

The basic equation has to be modified when each measurement is thrown off by bias as well as chance error:

individual measurement = exact value + bias + chance error.

If there is no bias in a measurement procedure, the long-run average of repeated measurements should give the exact value of the thing being measured: the

Source: Freedman et al., Statistics, Norton, 1998

#### Bias-variance decomposition



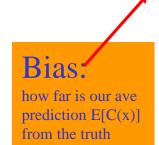
- Suppose classifiers C<sub>j</sub> and C<sub>k</sub> were trained on different sets S<sub>j</sub> and S<sub>k</sub> of 1000 samples each
- Then C<sub>j</sub> and C<sub>k</sub> might have different accuracy
- What is the expected accuracy of a classifier C trained this way?

- Let Y = f(X) be what C is trying to predict
- The expected squared error at a test instance x, averaging over all such training samples, is

$$E[C(x) - f(x)]^{2}$$

$$= E[C(x) - E[C(x)]]^{2}$$

$$+ [E[C(x)] - f(x)]^{2}$$



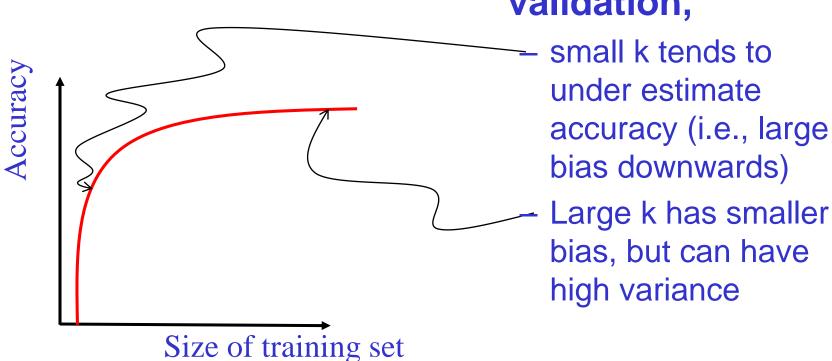
# Proof of bias-variance decomposition Nation of Sir

 $E[C(x) - f(x)]^2$  $= E[C(x) - E[C(x)] + E[C(x)] - f(x)]^{2}$  $= E[(C(x) - E[C(x)])^2 + (E[C(x)] - f(x))^2 - 2(C(x) - E[C(x)])(E[C(x)] - f(x))]$  $= E[C(x) - E[C(x)]]^{2} + E[E[C(x)] - f(x)]^{2} - 2 E(C(x) - E[C(x)]) (E[C(x)] - f(x))$  $= E[C(x) - E[C(x)]]^{2} + (E[C(x)] - f(x))^{2} - 2(E[C(x)] - E[C(x)])(E[C(x)] - f(x))$  $= E[C(x) - E[C(x)]]^2 + (E[C(x)] - f(x))^2$ Bias: Variance: how much our how far is our ave estimate C(x) will prediction E[C(x)]vary across the from the truth different training sets

#### Bias-variance trade-off



 In k-fold cross validation,



### Food for thought



- What is the logical basis of cross validation?
- Hint: Central limit theorem
- What / whose accuracy does it really estimate?



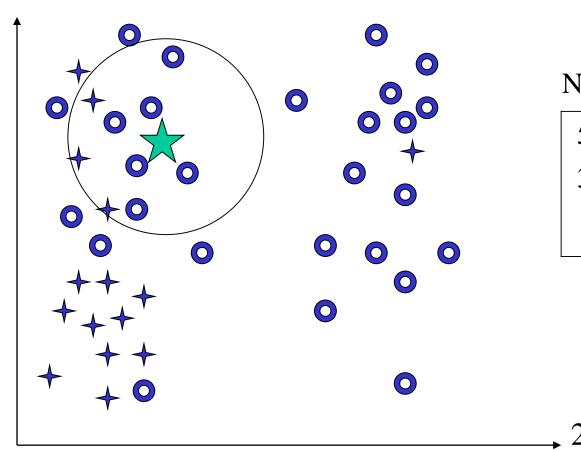
## Curse of dimensionality



#### Recall kNN ...



1st dimension



#### Neighborhood

5 of class O 3 of class →

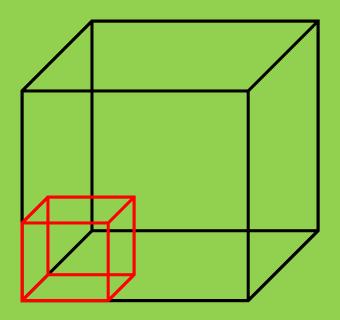
2<sup>nd</sup> dimension

Image credit: Zaki

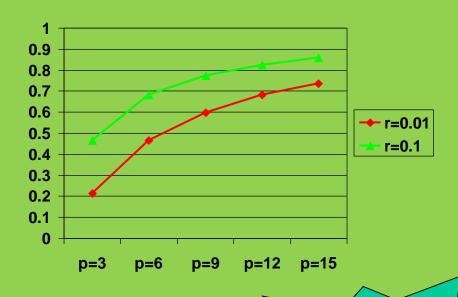
#### Curse of dimensionality



 How much of each dimension is needed to cover a proportion r of total sample space?



- Calculate by e<sub>p</sub>(r) = r<sup>1/p</sup>. Why?
- So, to cover 10% of a 15-D space, need 85% of each dimension!



#### Consequence of the curse



- Suppose the number of samples given to us in the total sample space is fixed
- Let the dimension increase
- Then the distance of the k nearest neighbours of any point increases
- Then the k nearest neighbours are less and less useful for prediction, and can confuse the k-NN classifier

#### What is feature selection?



#### Tackling the curse

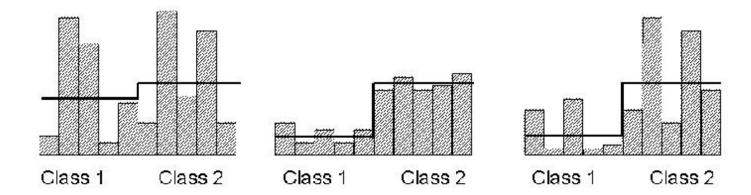


- Given a sample space of p dimensions
- It is possible that some dimensions are irrelevant
- Need to find ways to separate those dimensions (aka features) that are relevant (aka signals) from those that are irrelevant (aka noise)

## Signal selection (Basic idea)



- Choose a feature w/ low intra-class distance
- Choose a feature w/ high inter-class distance



# Signal selection (e.g., t-statistics National University of Singapore

The t-stats of a signal is defined as

$$t = \frac{|\mu_1 - \mu_2|}{\sqrt{(\sigma_1^2/n_1) + (\sigma_2^2/n_2)}}$$

where  $\sigma_i^2$  is the variance of that signal in class i,  $\mu_i$  is the mean of that signal in class i, and  $n_i$  is the size of class i.

## Food for thought



- How is the t-statistic typically used?
- What are the assumptions required for this way of using the t-statistic?



## Self-fulfilling oracle



- Construct artificial dataset with 100 samples, each with 100,000 randomly generated features and randomly assigned class labels

Select 20 features
 with the best t statistics (or other
 methods)

 The resulting accuracy can be ~90%

Evaluate accuracy by

the 20 selected

features

cross validation using

 But the true accuracy should be 50%, as the data were derived randomly

#### What went wrong?



- The 20 features were selected from whole dataset
- Information in the held-out testing samples has thus been "leaked" to the training process
- The correct way is to re-select the 20 features at each fold; better still, use a totally new set of samples for testing



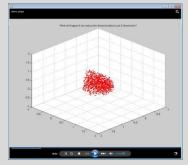
While **dimensionality reduction** is an important tool in machine learning/data mining, we must always be aware that it can distort the data in misleading ways.

Above is a two dimensional projection of an intrinsically three dimensional world....

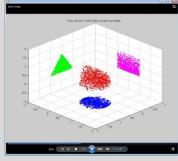


Original photographer unknown/
See also www.cs.gmu.edu/~jessica/DimReducDanger.htm

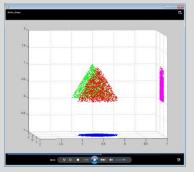
#### A cloud of points in 3D



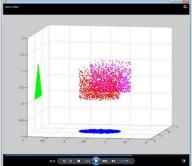
Can be projected into 2D XY or XZ or YZ



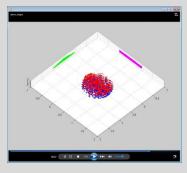
In 2D XZ we see a triangle



In 2D YZ we see a square



In 2D XY we see a circle



Screen dumps of a short video from www.cs.gmu.edu/~jessica/DimReducDanger.htm

# Concluding remarks



#### What have we learned?



- Methodology of data mining
  - Feature generation, feature selection, feature integration
- Evaluation of classifiers
  - Accuracy, sensitivity, precision
  - Cross validation
- Curse of dimensionality
  - Feature selection concept
  - Self-fulfilling oracle

# Any questions?



#### Acknowledgements



- The first two slides were shown to WLS ~20 years ago by Tan Ah Hwee
- The three slides on the dangers of dimensionality reduction were created by Eamonn Keogh

#### References



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