

# Part 2

# MDL in Action



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# Explicit Coding

Ad hoc sounds bad, but is it really?

- Bayesian learning for instance, is **inherently subjective**, plus
- biasing search is a time-honoured tradition in data analysis

Using an **explicit encoding** allows us  
to **steer towards** the  
type of structure **we want to discover**

We so also mitigate one of the practical weak spots of AIT

- all data is a string, but wouldn't it be nice if the structure you found would not depend on the order of the data?

# Matrix Factorization

The rank of a matrix  $A$  is

- number of rank-1 matrices that when summed form  $A$  (**Schein rank**)

The diagram shows a large green square labeled  $A$  on the left. To its right is an equals sign, followed by three rank-1 matrices added together, and an ellipsis. Each rank-1 matrix is represented by a vertical green bar on the left and a horizontal green bar on the right. The first rank-1 matrix has a vertical bar labeled  $b_1$  and a horizontal bar labeled  $c_1$ , with the expression  $b_1 \circ c_1$  below it. The second rank-1 matrix has a vertical bar labeled  $b_2$  and a horizontal bar labeled  $c_2$ , with the expression  $b_2 \circ c_2$  below it. The third rank-1 matrix has a vertical bar labeled  $b_3$  and a horizontal bar labeled  $c_3$ , with the expression  $b_3 \circ c_3$  below it. Plus signs are placed between the rank-1 matrices, and an ellipsis  $\dots$  is placed to the right of the third one.

$$A = b_1 \circ c_1 + b_2 \circ c_2 + b_3 \circ c_3 + \dots$$

# Boolean Matrix Factorization

The rank of a Boolean matrix  $A$  is

- number of rank-1 matrices that when summed form  $A$  (**Schein rank**)

$$A = b_1 \circ c_1 + b_2 \circ c_2 + b_3 \circ c_3 + \dots$$

# Boolean Matrix Factorization

The rank of a Boolean matrix  $A$  is

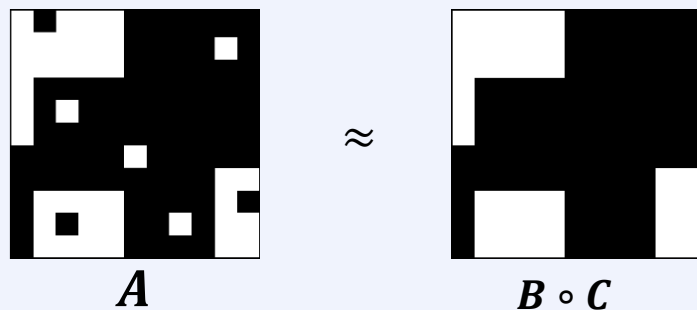
- number of rank-1 matrices that when summed form  $A$  (**Schein rank**)
- noise quickly inflate the 'true' latent rank to  $\min(n, m)$

The diagram illustrates the Boolean Matrix Factorization process. On the left, a matrix  $A$  is shown as a 5x5 grid of black and white squares. This matrix is equal to the sum of three rank-1 matrices, each represented as a 5x5 grid of black and white squares. The first rank-1 matrix, labeled  $b_1 \circ c_1$ , has a black square in the top-right corner. The second rank-1 matrix, labeled  $b_2 \circ c_2$ , has a black square in the center. The third rank-1 matrix, labeled  $b_3 \circ c_3$ , has a black square in the bottom-right corner. The sum of these three matrices is equal to the matrix  $A$ . The diagram is followed by an ellipsis, indicating that there can be more than three rank-1 matrices in the sum.

# Boolean Matrix Factorization

Noise quickly inflates the rank to  $\min(n, m)$

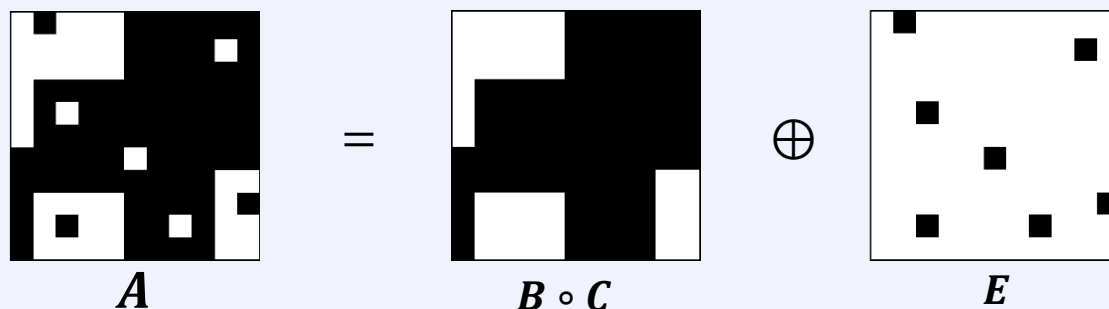
- how can we determine the 'true' latent rank?



# Boolean Matrix Factorization

## Separating structure and noise

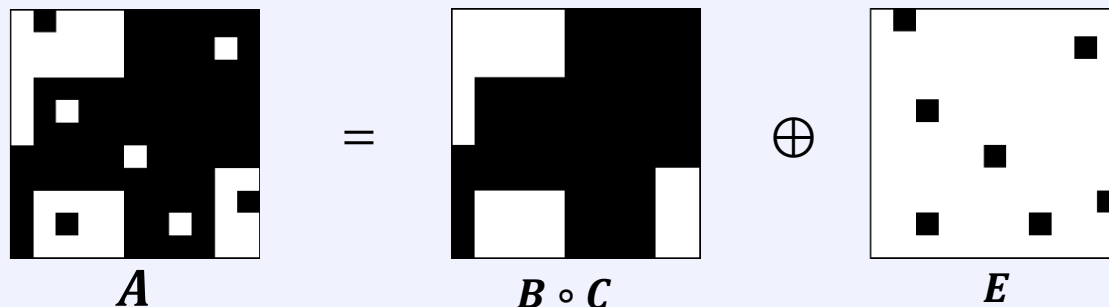
- matrices  $B$  and  $C$  contain structure, matrix  $E$  contains noise



# Boolean Matrix Factorization

Encoding the structure

$$L(\mathbf{B}) = \log n + \sum_{b \in \mathbf{B}} \left[ \log n + \log \binom{n}{|b|} \right]$$

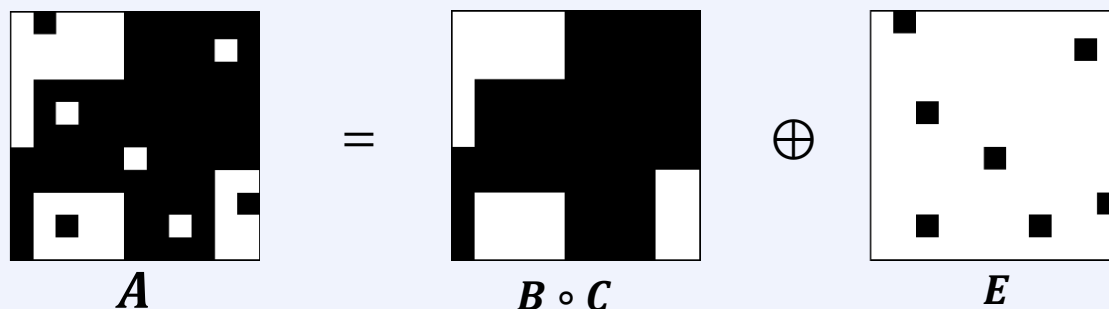




# Boolean Matrix Factorization

Encoding the structure

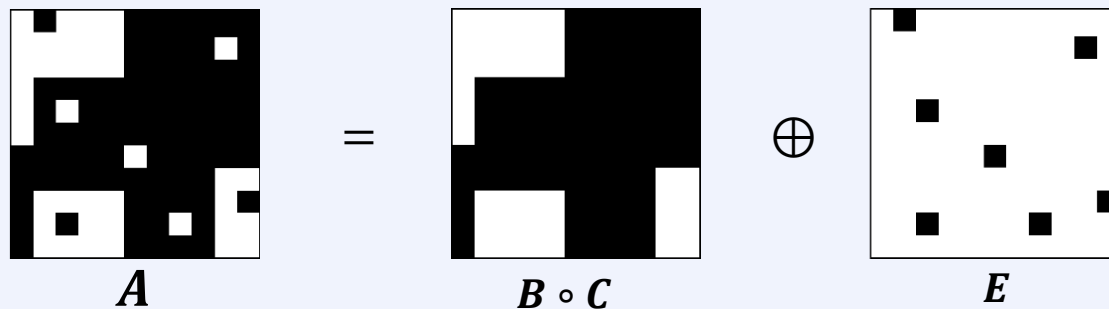
$$L(\mathcal{C}) = \log m + \sum_{c \in \mathcal{C}} \left[ \log m + \log \binom{m}{|c|} \right]$$



# Boolean Matrix Factorization

Encoding the noise

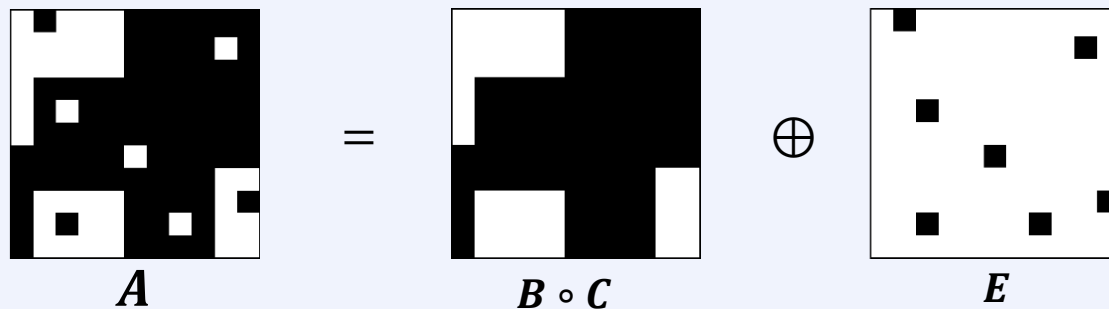
$$L(\mathbf{E}) = \log nm + \log \binom{nm}{|\mathbf{E}|}$$



# Boolean Matrix Factorization

MDL for BMF

$$L(D, H) = L(\mathbf{B}) + L(\mathbf{C}) + L(\mathbf{E})$$



# Pattern Mining

The **ideal** outcome of pattern mining

- patterns that show the structure of the data
- preferably a small set, without redundancy or noise

Frequent pattern mining does **not** achieve this

- pattern explosion → overly many, overly redundant results

MDL allows us to effectively pursue the ideal

- we want a group of patterns that summarise the data well
- we take a **pattern set** mining approach

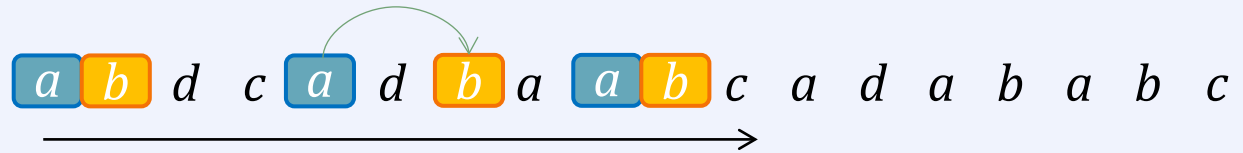


# Event sequences

Alphabet  $\Omega$

$\{ a, b, c, d, \dots \}$

Data  $D$

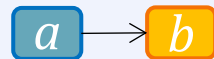


one, or  
multiple  
sequences

$\{ a b d c a d b a a b c,$   
 $a b d c a d b,$   
 $a b d c a d b a a, \dots \}$

Patterns

serial  
episodes



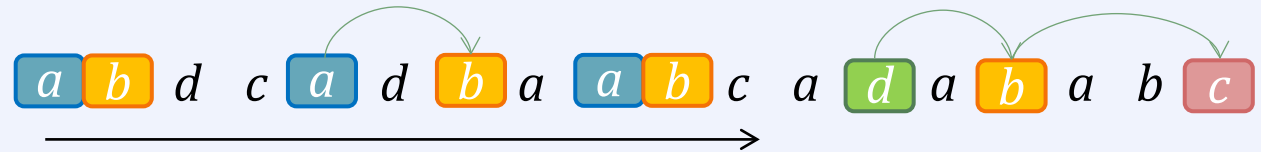
'subsequences  
allowing gaps'

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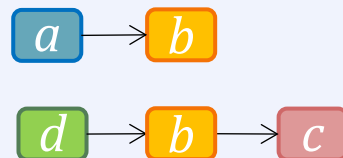


one, or  
multiple  
sequences

$\{ a b d c a d b a a b c, \\ a b d c a d b, \\ a b d c a d b a a, \dots \}$

Patterns

serial  
episodes



'subsequences  
allowing gaps'

# Models

pattern	code	gap	non-gap
<i>abc</i>	<i>p</i>	<i>?</i>	<i>!</i>
<i>da</i>	<i>q</i>	<i>?</i>	<i>!</i>
<i>a</i>	<i>a</i>	-	-
<i>b</i>	<i>b</i>	-	-
<i>c</i>	<i>c</i>	-	-
<i>d</i>	<i>d</i>	-	-

As models we use **code tables**

- dictionary of patterns & codes
- always contains all singletons

We use optimal prefix codes

- easy to compute,
- behave predictably,
- good results,
- more details follow




# Encoding Event Sequences

Data  $D$ :  $a \ b \ d \ c \ a \ d \ b \ a \ a \ b \ c$

Encoding 1: using only singletons

$C_p$  

$CT_1$ : 

The length of the code  $\boxed{X}$  for pattern  $X$

$$L(\boxed{X}) = -\log(p(\boxed{X})) = -\log\left(\frac{usg(X)}{\sum usg(Y)}\right)$$

The length of the code stream

$$L(C_p) = \sum_{X \in CT} usg(X) L(\boxed{X})$$

# Encoding Event Sequences

Data  $D$ :       
*a b d c a d b a a b c* →

Encoding 2: using patterns

$C_p$  p d a q b p  
 $C_g$  ! ? ! ? ! ! !

Alignment:       
*a b d c a d b a a b c* →  
p ! ? !     q ? ! p ! !

$CT_2$ : 

<i>a</i>	a		
<i>b</i>	b		
<i>c</i>	c		
<i>d</i>	d		
<i>abc</i>	p	?	!
<i>da</i>	q	?	!

gaps ↓

non-gaps ↓

# Encoding Event Sequences

Data  $D$ :       
*a b d c a d b a a b c* →

Encoding 2: using patterns

$C_p$  p d a q b p  
 $C_g$  ! ? ! ? ! ! !

$CT_2$ : 

<i>a</i>	<i>a</i>		
<i>b</i>	<i>b</i>		
<i>c</i>	<i>c</i>		
<i>d</i>	<i>d</i>		
<i>abc</i>	<i>p</i>	?	!
<i>da</i>	<i>q</i>	?	!

gaps

↓

non-gaps

↓

The length of a gap code ? for pattern  $X$

$$L(\text{?}) = -\log(p(\text{?} \mid p))$$

and analogue for non-gap codes !

# Encoding Event Sequences

By which, the encoded size of  $D$  given  $CT$  and  $C$  is

$$L(D | CT) = L(C_p | CT) + L(C_g | CT)$$

which leaves us to define  $L(CT | C)$

# Encoding a Code Table

$L(CT \mid C, D)$  consists of

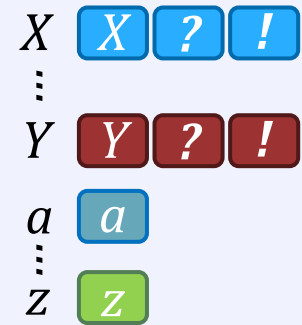
$X$	$X$	$?$	$!$
$\vdots$			
$Y$	$Y$	$?$	$!$
$a$	$a$		
$\vdots$			
$Z$	$Z$		

# Encoding a Code Table

$L(CT | C, D)$  consists of

- 1) base singleton counts in  $D$

$$L_{\mathbb{N}}(|\Omega|) + L_{\mathbb{N}}(|D|) + \log \binom{|D| - 1}{|\Omega| - 1}$$

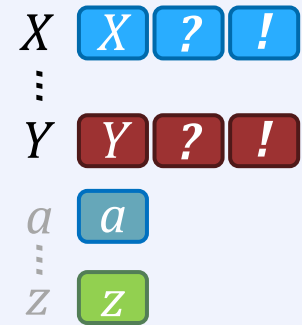


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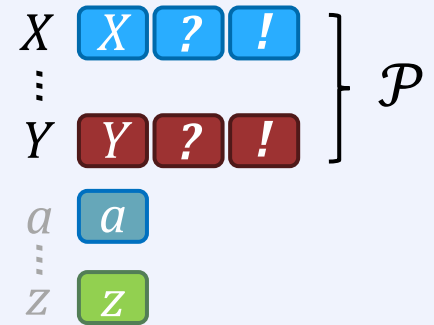
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$$L_{\mathbb{N}}(|\Omega|) + L_{\mathbb{N}}(|D|) + \log \binom{|D| - 1}{|\Omega| - 1}$$

- 2) number of patterns, total, and per pattern usage

$$L_{\mathbb{N}}(|\mathcal{P}| + 1) + L_{\mathbb{N}}(usg(\mathcal{P}) + 1) + \log \binom{usg(\mathcal{P}) - 1}{|\mathcal{P}| - 1}$$





# Encoding a Code Table

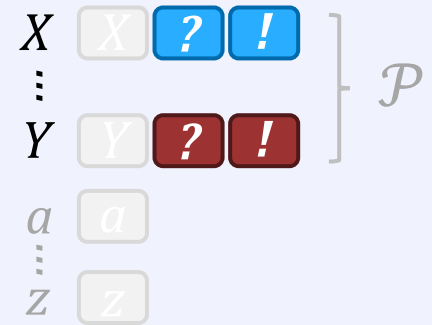
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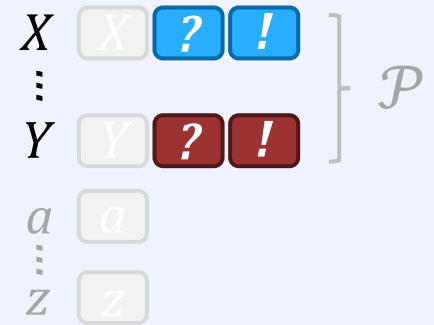
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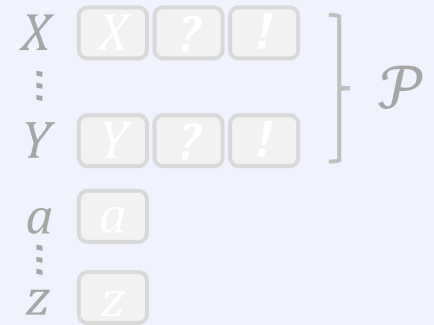
$$L_{\mathbb{N}}(|\mathcal{P}| + 1) + L_{\mathbb{N}}(usg(\mathcal{P}) + 1) + \log \binom{usg(\mathcal{P}) - 1}{|\mathcal{P}| - 1}$$

- 3) per pattern  $X$  : its length, elements, and number of gaps

$$L_{\mathbb{N}}(|X|) - \left[ \sum_{x \in X} \log p(x \mid D) \right] + L_{\mathbb{N}}(gaps(X) + 1)$$

# Encoding a Code Table

$L(CT \mid C, D)$  consists of



- 1) base singleton counts in  $D$

$$L_{\mathbb{N}}(|\Omega|) + L_{\mathbb{N}}(|D|) + \log \binom{|D| - 1}{|\Omega| - 1}$$

- 2) number of patterns, total, and per pattern usage

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$$L_{\mathbb{N}}(|X|) - \left[ \sum_{x \in X} \log p(x \mid D) \right] + L_{\mathbb{N}}(gaps(X) + 1)$$

# Encoding Event Sequences

By which we have a lossless encoding.  
In other words, an objective function.

By MDL, our goal is now to minimise

$$L(CT, D) = L(CT | C) + L(D | CT)$$

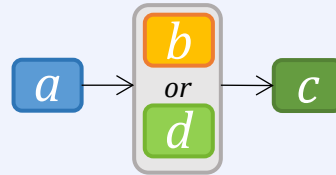
for how to do so, please see the papers



# Selected Results



Serial Episodes



Choice-episode



Ontological Episodes

## PRES. ADDRESSES

unit[ed] state[s]  
take oath  
army navy  
under circumst.  
econ. public expenditur  
exec. branch. governm.

## JMLR

empirical, structural risk minimization  
indep, principial component analysis  
Mahalanobis, edit, Euclidean, pairwise distance

## LOTR

he Verb Conj he  
[he said that he]  
Conf \_ the Noun of  
[and even the end of]  
the Adj Noun and  
[the young Hobbits and]

# Clustering

The best clustering is the one that costs the least bits

- similar structure (patterns) within clusters
- different structure (patterns) between clusters

Partition your data such that

$$L(C) + \sum_{(D_i, H_i) \in C} L(D_i, H_i)$$

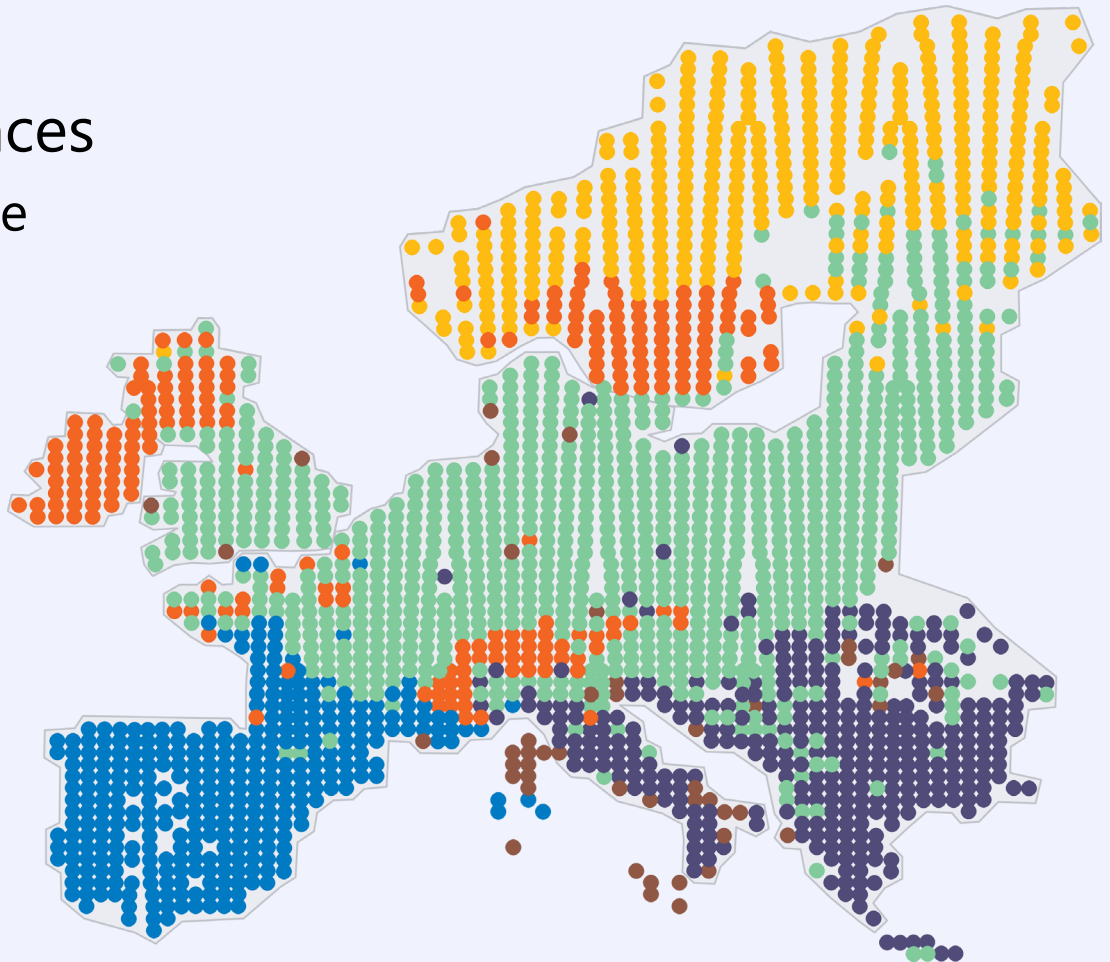
is minimal

(similar to mixture modelling, but descriptive instead of predictive)

# Clustering

## Mammals occurrences

- 2221 areas in Europe
- 50x50km each
- 123 mammals
- no location info



$k=6$ , MDL 'optimal'



# Classification

Split your data **per class**

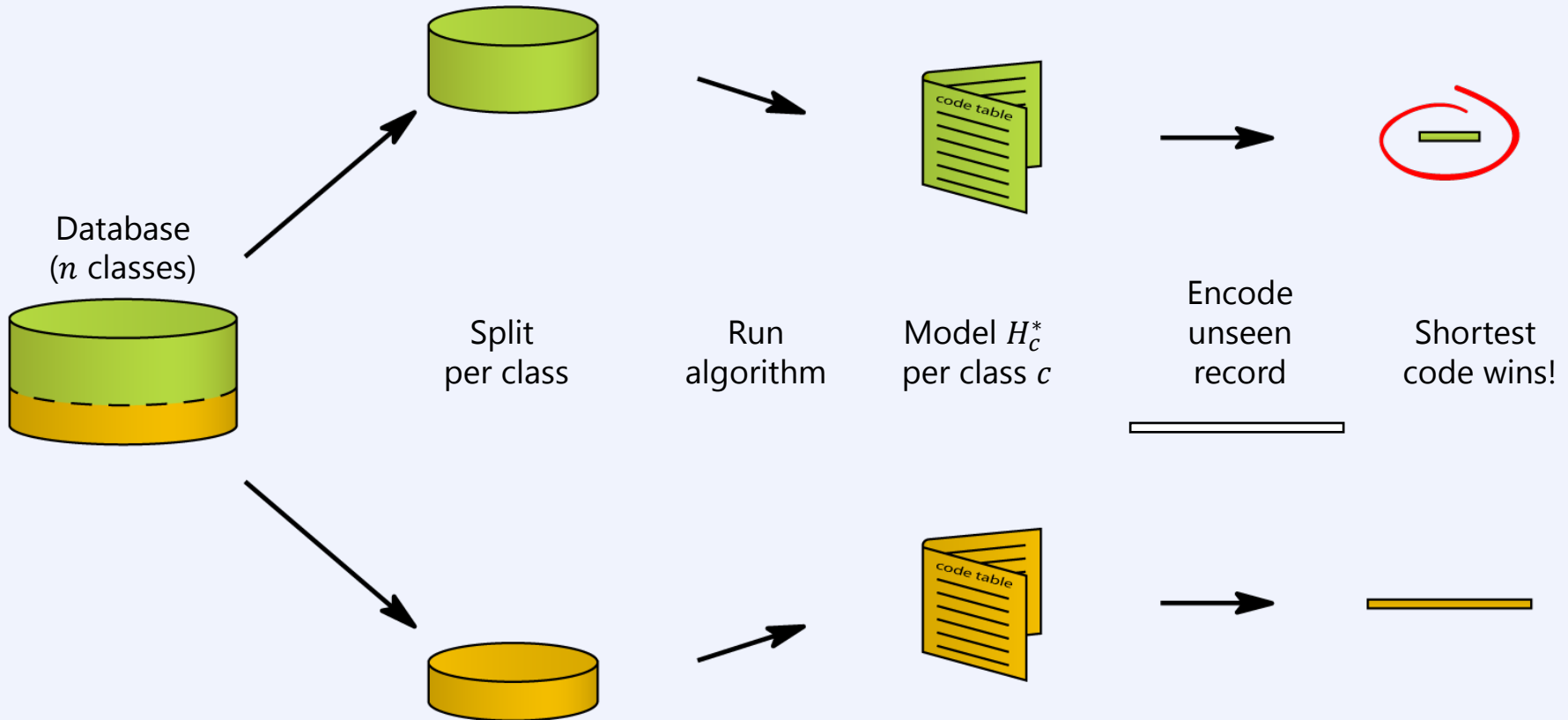
- induce model per class

Then, for **unseen** instances

- **assign** class label of model that encodes it **shortest**

$$L(x | H_1) < L(x | H_2) \rightarrow P(x | H_1) > P(x | H_2)$$

# Classification by MDL



$$L(x | H_1) < L(x | H_2) \rightarrow P(x | H_1) > P(x | H_2)$$

# Outlier Detection

## One-Class Classification (aka anomaly detection)

- lots of data for **normal** situation – insufficient data for **target**

## Compression models the **norm**

- anomalies will have **high** description length  $L(t \mid H_{norm}^*)$

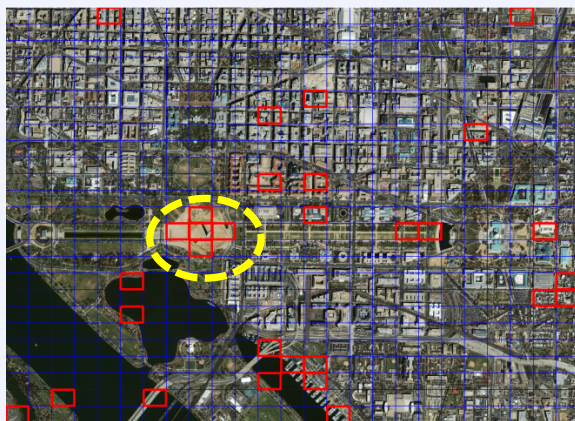
## Very nice properties

- **performance**            high accuracy
- **versatile**                no distance measure needed
- **characterisation**        *'this part of t is incompressible'*

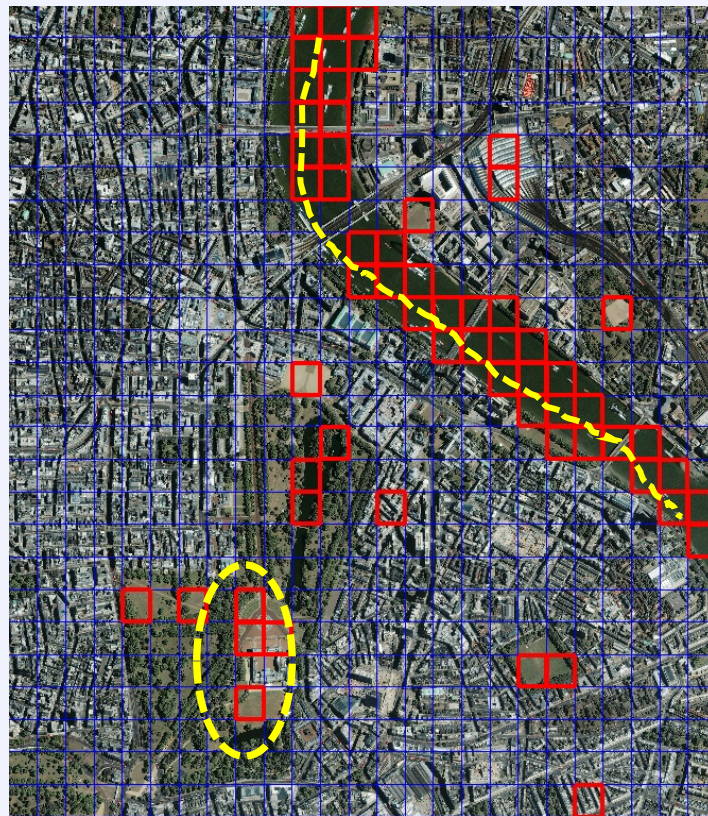
# CompreX on Images



Catholic church, Vatican



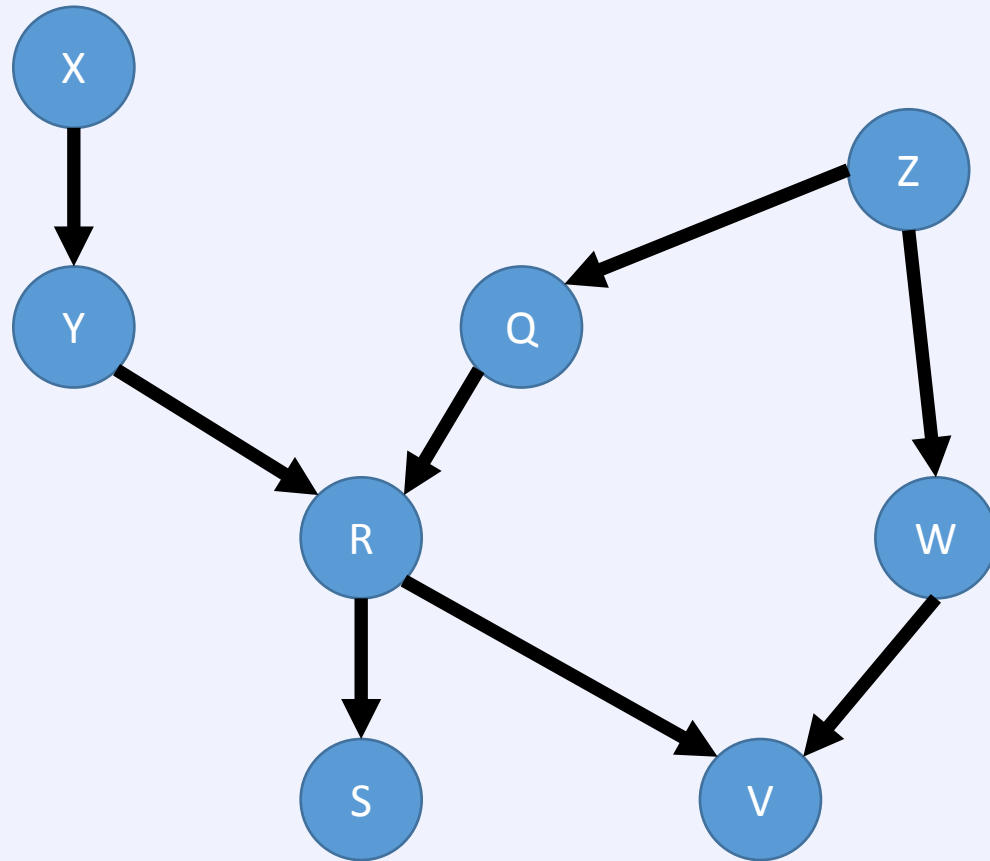
Washington Memorial, D.C.



Thames river, Buckingham palace, plain fields, London

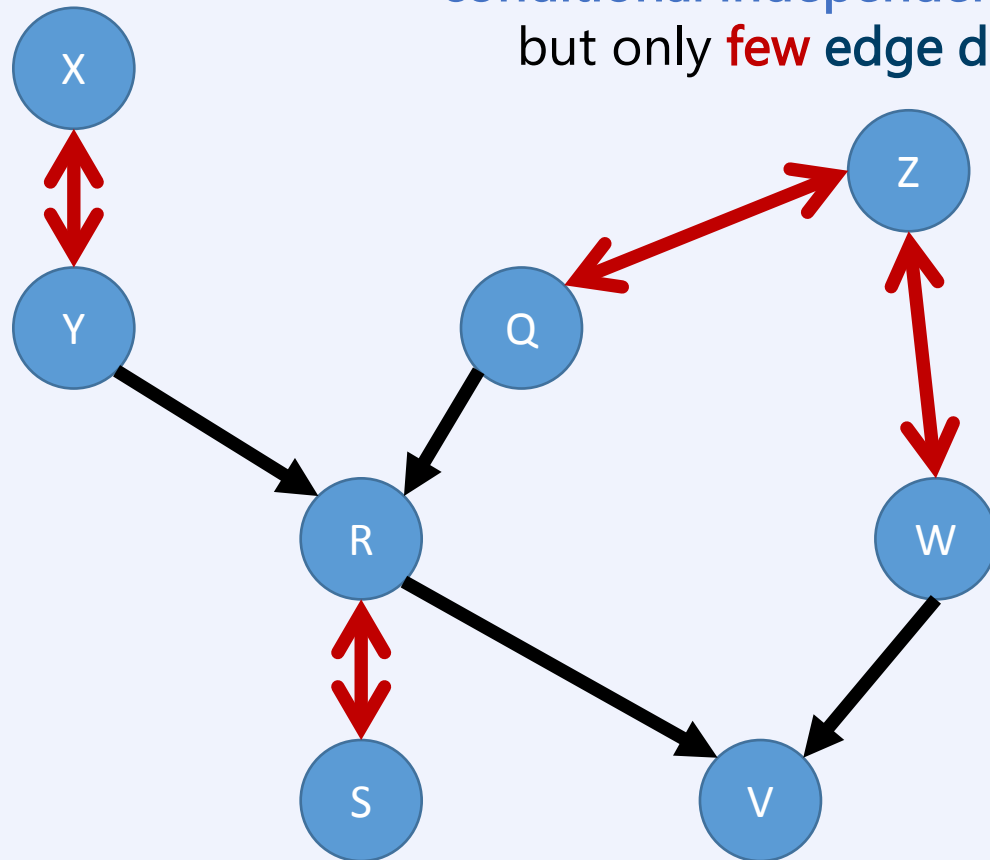


# Causal Discovery



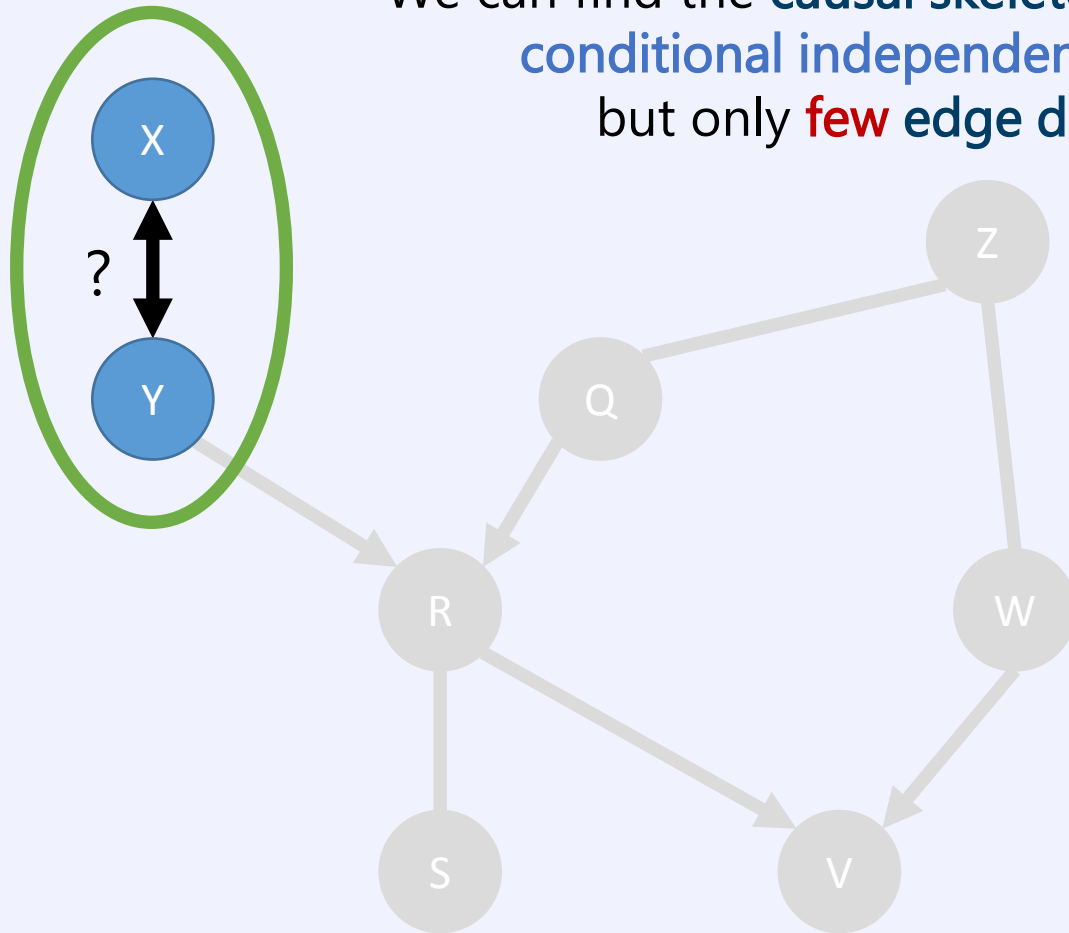
# Causal Discovery

We can find the **causal skeleton** using **conditional independence** tests, but only **few** edge directions



# Causal Inference

We can find the **causal skeleton** using **conditional independence** tests, but only **few edge directions**



# Algorithmic Markov Condition

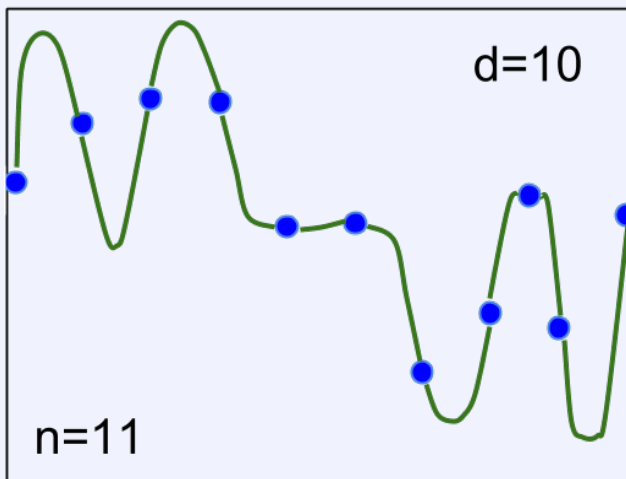
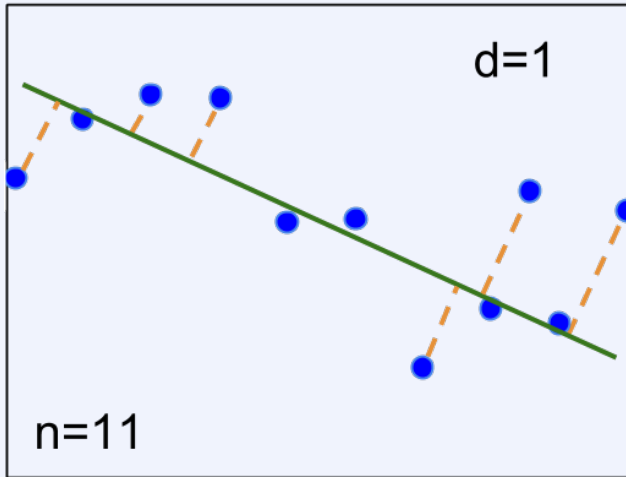
If  $X \rightarrow Y$ , we have,  
up to an additive constant,

$$K(P(X)) + K(P(Y|X)) \leq K(P(Y)) + K(P(X|Y))$$

That is, we can do **causal inference** by identifying the factorization of the joint with the **lowest Kolmogorov complexity**



# MDL and Regression



VS.

$$L(M) + L(D|M)$$

Arrows point from  $L(M)$  to  $a_1 x + a_0$  and from  $L(D|M)$  to **errors**.

$$a_{10} x^{10} + a_9 x^9 + \dots + a_0 \{ \}$$

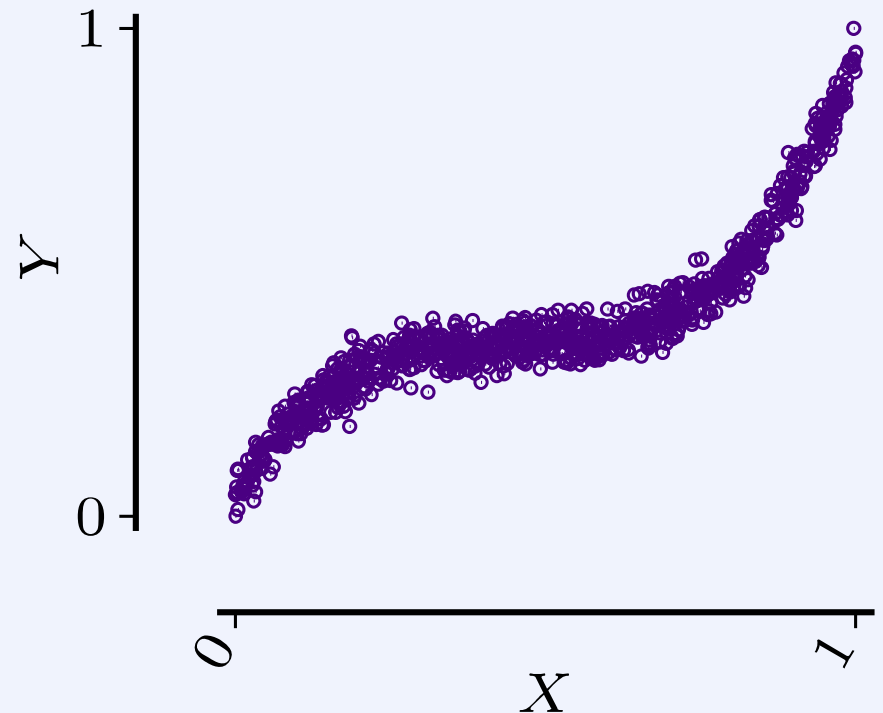
# Modelling the Data

We model  $Y$  as

$$Y = f(X) + \mathcal{N}$$

As  $f$  we consider **linear**, **quadratic**, **cubic**, **exponential**, and **reciprocal** functions, and model the noise using a 0-mean Gaussian. We choose the  $f$  that minimizes

$$L(Y | X) = L(f) + L(\mathcal{N})$$



# Confidence and Significance

How certain are we?

$$\mathbb{C} = \underbrace{|L(X) + L(Y | X)|}_{L(X \rightarrow Y)} - \underbrace{|L(Y) + L(X | Y)|}_{L(Y \rightarrow X)} \quad \blacksquare \text{ the higher the more certain}$$

# Confidence and Significance

How certain are we?

$$\mathbb{C} = \left| \frac{L(X) + L(Y | X)}{L(X) + L(Y)} - \frac{L(Y) + L(X | Y)}{L(X) + L(Y)} \right|$$

- the higher the more certain
- robust w.r.t. sample size

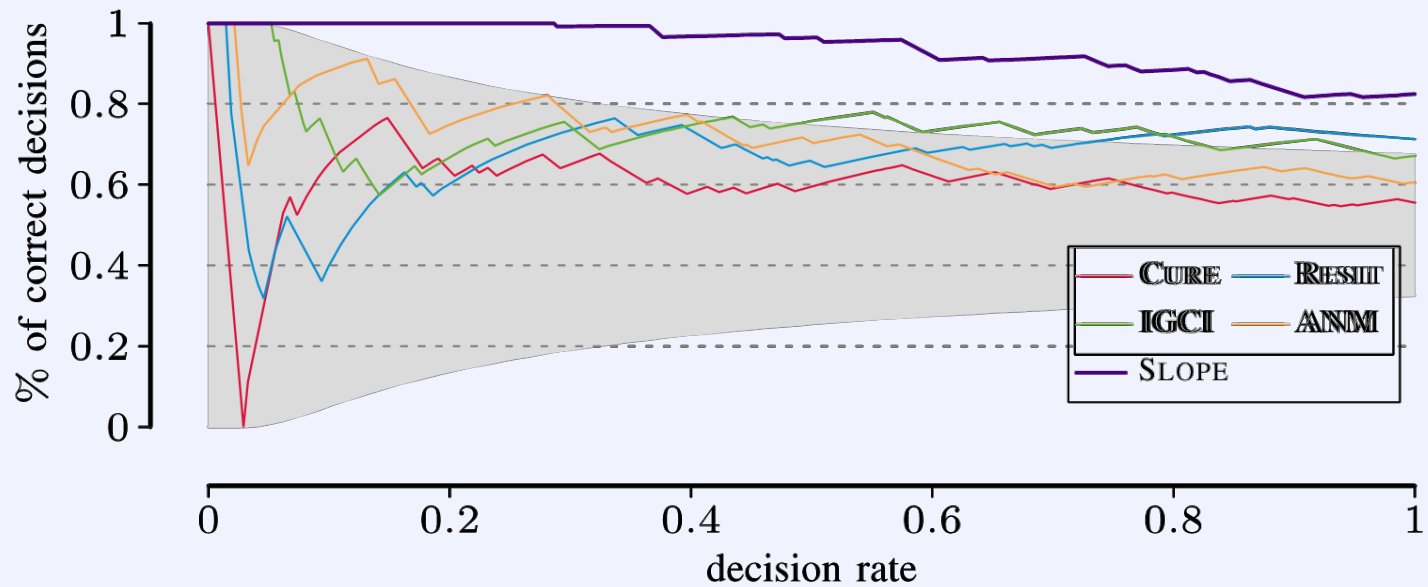
Is a given inference significant?

- our null hypothesis  $L_0$  is that  $X$  and  $Y$  are only **correlated**,  
we have  $L_0 = \frac{|L(X \rightarrow Y) - L(Y \rightarrow X)|}{2}$
- we can use the no-hypercompression inequality to test significance

$$P(L_0(D) - L(D) \geq k) \leq 2^{-k}$$

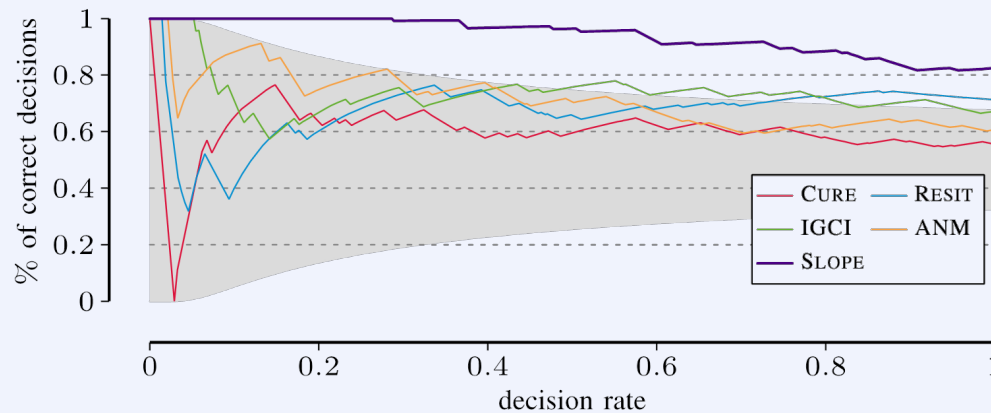
# Performance on Benchmark Data

(Tübingen 97 univariate numeric cause-effect pairs, weighted)

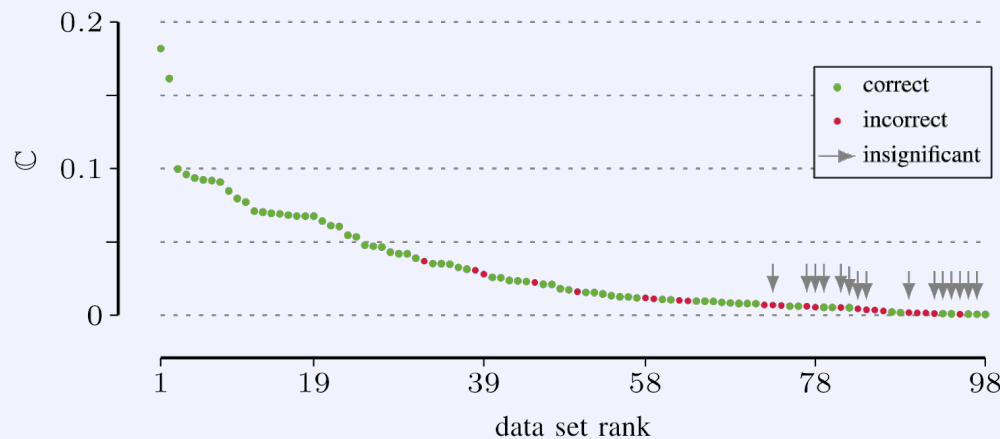


# Performance on Benchmark Data

(Tübingen 97 univariate numeric cause-effect pairs, weighted)



Inferences of state of the art algorithms **ordered** by **confidence** values.



SLOPE is 85% accurate with  $\alpha = 0.001$

# Deep Learning

Model selection in deep learning is hard

- way too many 'free' parameters for standard regularizers,
- no meaningful prior over networks, and
- uniform prior will lead to overfitting

How about an MDL approach?

- what is the description length of a neural network?

# MDL for Neural Networks

Suppose neural network  $H \in \mathcal{H}$  predicts target  $y$  given  $x$   
$$\hat{y} = H(x)$$

How do we encode data given the model?

- if  $H(x)$  is probabilistic, we have  $L(\mathbf{y} | H(\mathbf{x})) = -\sum_{y_i \in \mathbf{y}} \log p(y_i | x_i)$
- else we can simply encode the residual error,
  - e.g. if  $\mathbf{y}$  is binary, we have  $\mathbf{e} = \mathbf{y} \oplus \hat{\mathbf{y}}$ , and  $L(\mathbf{y} | H(\mathbf{x})) = \log n + \log \binom{n}{|\mathbf{e}|}$
  - e.g. if  $\mathbf{y}$  is continuous, we can encode using a zero-mean Gaussian



# MDL for Neural Networks

Suppose neural network  $H \in \mathcal{H}$  predicts target  $y$  given  $x$

$$\hat{y} = H(x)$$

How do we encode the model?

- we could encode all of the parameters, but that's highly ad hoc
- instead, we can use the notion of **prequential coding**

# Prequential Coding

Simple, elegant idea:

*“Update your model after every message”*

That is, we re-train our network after ‘every’ new label

- we initialize topology  $H \in \mathcal{H}$  with fixed weights
- we transmit the first  $k$  labels using  $H_0$
- we now train  $H$  on this first batch of  $k$  labelled points, we obtain  $H_1$
- we transmit the second  $k$  labels using  $H_1$
- we now train  $H$  on the first two batches, and obtain  $H_2$

# Prequential Coding

Simple, elegant idea:

*“Update your model after every message”*

$$L(D | \mathcal{H}) = \sum_{D_i} L(D_i | H_{i-1})$$

Best of all, this is not a crude, but a refined MDL code!

- depends fully on how  $H$  behaves on the data
- no arbitrary choices on how to encode  $H$
- within a constant of  $L(D|H^*)$ , and this constant only depends on  $\mathcal{H}$

# Schedule

8:00am Opening

8:10am Introduction to MDL

8:50am MDL in Action

9:30am  ~~—————*break*—————~~

10:00am Stochastic Complexity

11:00am MDL in Dynamic Settings

