

A methodology for detection of causal relationships between discrete time series on systems

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Theoretical Foundations - Entropy

Definition

The amount of information produced by an information source.

$$H = \sum_i^n p_i(i) \log_2 \frac{1}{p(i)} \quad (1)$$

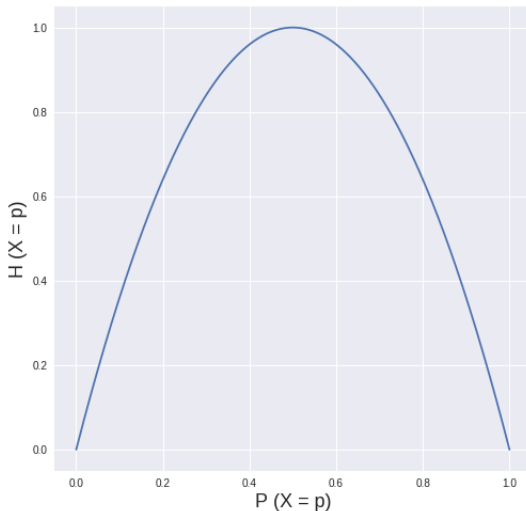
Theoretical Foundations - Entropy

It is related to the frequency of the appearance of each value or the amount of surprise obtained given the appearance of a state or value.

Properties:

- It is continuous in the domain of p_i , which is the probability mass function.
- It is monotonically increasing, in the n domain, when all events are likely equally.
- It is weighted additive when a choice is broken down into successive choices.

Theoretical Foundations - Entropy



Theoretical Foundations - Kullback-Leibler Divergence

Definition

- Given a variable I , with probability distribution p . It measures the error or divergence when it is assumed that the probability distribution of I is q , instead of p .

$$KL_I = \sum_i^n p(i) \log \frac{p(i)}{q(i)} \quad (2)$$

Theoretical Foundations - Kullback-Leibler Divergence

$$K_{I,J} = \sum_{i,j}^n p(i,j) \log \frac{p(i,j)}{q(i,j)} \quad (3)$$

$$K_{I|J} = \sum_{i,j}^n p(i,j) \log \frac{p(i|j)}{q(i|j)} \quad (4)$$

Theoretical Foundations - Mutual Information

Definition

It is a measure derived from the Kullback-Leibler divergence. It is calculated between two processes I and J and quantifies the amount of information obtained about one process when observing the other.

It can be seen as the “information produced by erroneously assuming that the two processes are independent”.⁴

$$MI_{I,J} = \sum_{i,j}^n p(i,j) \log \frac{p(i,j)}{p(i) \cdot p(j)} \quad (5)$$

⁴(SCHREIBER, 2000)

Theoretical Foundations - Transfer Entropy

Definition

It is a theoretical measure “that shares some of the desired properties of mutual information but takes the dynamics of information transport into account”.⁵

$$TE_{J \rightarrow I} = \sum_{i, i_{t+h}, j} p(i_{t+h}, i_t^k, j_t^l) \log \frac{p(i_{t+h} | i_t^k, j_t^l)}{p(i_{t+h} | i_t^k)} \quad (6)$$

⁵(SCHREIBER, 2000)

Theoretical Foundations - Transfer Entropy

$$i^k = [i_t, \dots, i_{t-k+1}] \quad (7)$$

$$j^l = [j_t, \dots, j_{t-l+1}] \quad (8)$$

- It infers the veracity of the equation:

$$p(i_{t+h} \mid i_t^k, j_t^l) = p(i_{t+h} \mid i_t^k) \quad (9)$$

Theoretical Foundations - Transfer Entropy

Understanding of the parameters (Example)

- Series size = 6 samples
- $k = 3$ samples
- $l = 2$ samples
- $h = 1$ sample

I series

0	0	0	1	1	0	0
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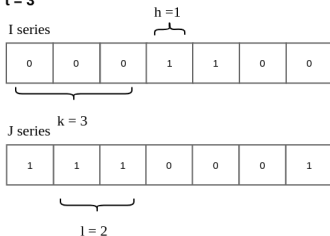
J series

1	1	1	0	0	0	1
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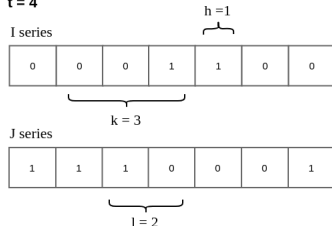
Figure 2: I and J series.

Theoretical Foundations - Transfer Entropy

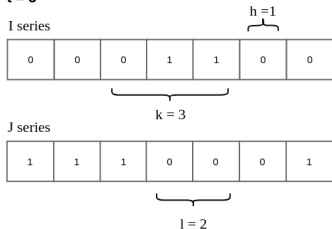
t = 3



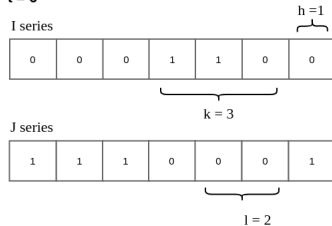
t = 4



t = 5



t = 6



Theoretical Foundations - Bayesian Networks

- Probabilistic models based on direct acyclic graphs.
- The nodes represent variables of interest, while the connections represent relationships of informational or causal dependencies.⁶
- Each variable is independent from its nondescendants, given its parents.

⁶(PEARL, 2011)

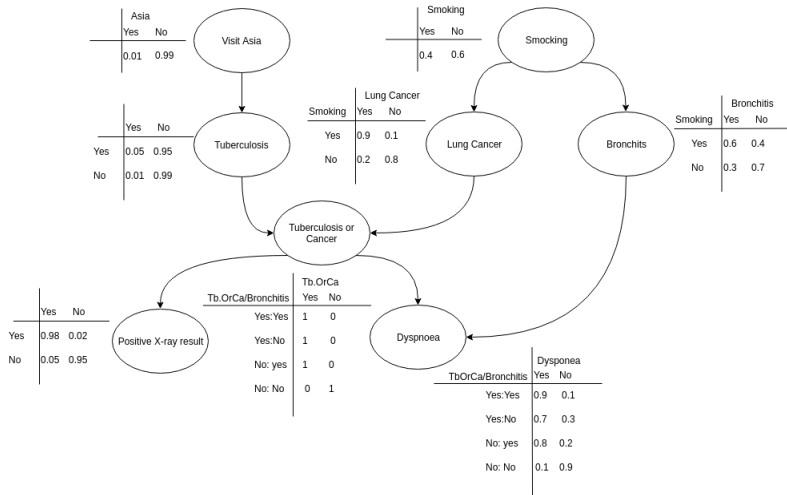
- It factors the joint the distribution of the variable into local joint distributions.

$$P(A_n, \cap \dots \cap A_1) = \prod_i^n P(A_i \mid \cap_{j=1}^{i-1} A_j) \quad (10)$$

$$P(A_n, \cap \dots \cap A_1) = \prod_{i=1}^n P(A_i \mid pa_i) \quad (11)$$

$$P(TbOrCa \mid Cancer, \textit{Bronch.}, \textit{Asia}) = P(TbOrCa \mid Cancer) \quad (12)$$

Theoretical Foundations - Bayesian Networks



Theoretical Foundations - K2 Algorithm

K2 - Algorithm

- Bayesian method for estimating a probabilistic network from data.
- The algorithm searches for the network that has the highest posterior probability given a database of records.
- Necessity of a prior ordering on nodes.
- Scores each local network in order to find the most probable structure.

Case	Variable values for each case		
	x_1	x_2	x_3
1	<i>present</i>	<i>absent</i>	<i>absent</i>
2	<i>present</i>	<i>present</i>	<i>present</i>
3	<i>absent</i>	<i>absent</i>	<i>present</i>
4	<i>present</i>	<i>present</i>	<i>present</i>
5	<i>absent</i>	<i>absent</i>	<i>absent</i>
6	<i>absent</i>	<i>present</i>	<i>present</i>
7	<i>present</i>	<i>present</i>	<i>present</i>
8	<i>absent</i>	<i>absent</i>	<i>absent</i>
9	<i>present</i>	<i>present</i>	<i>present</i>
10	<i>absent</i>	<i>absent</i>	<i>absent</i>

Figure 5: Database of cases

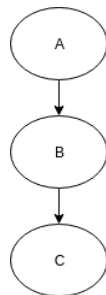


Figure 6: Possible structure.

K2-Algorithm

Algorithm 1: K2 Algorithm

Input: a set of n nodes - $(x_1 \dots x_n)$
 an ordering for the nodes ,
 a data set of n columns and m cases - database

Output: Bayesian Network Topology

```

1  $\pi_i \leftarrow$  parents of node  $i$ 
2
3 for  $i \leftarrow 1$  to  $n$  do
4    $\pi_i \leftarrow \emptyset$ 
5 end
6 for  $i \leftarrow 1$  to  $n$  do
7    $P_{old} \leftarrow g(i, \pi_i)$ 
8   while True do
9      $pred_{x_i} \leftarrow \text{Pred}(x_i)$  - set of nodes that precede  $x_i$ 
10    select the node  $x_j \in pred_{x_i} \setminus \pi_i$  that maximizes:  $g(i, \pi_i \cup \{x_j\})$ 
11     $P_{new} \leftarrow g(i, \pi_i \cup \{x_j\})$ 
12     $\text{sigma} \leftarrow P_{new} > P_{old}$ 
13    if  $\text{sigma} = \text{True}$  then
14       $P_{old} \leftarrow P_{new}$ 
15       $\pi_i \leftarrow \pi_i \cup x_j$ 
16    end
17    if  $\text{not}(pred_{x_i} = \emptyset)$  then
18       $pred_{x_i} \leftarrow pred_{x_i} \setminus x_j$ 
19    end
20    if  $(\text{not sigma or } pred_{x_i} = \emptyset)$  then
21      break
22    end
23  returns the parent set of each node
24 end

```

Figure 7: K2-Algorithm

Proposed Methodology

General Concept

By the use of the concepts of Information Theory and Bayesian Networks it is intended to unite the method of Transfer Entropy and the K2 algorithm in order to generate a single methodology for the detection of causal relationships.

Approach

To model the system as a graph, in which the nodes will be the entities related to each other, by a causality relationship. This detection will be made in five stages

Proposed Methodology

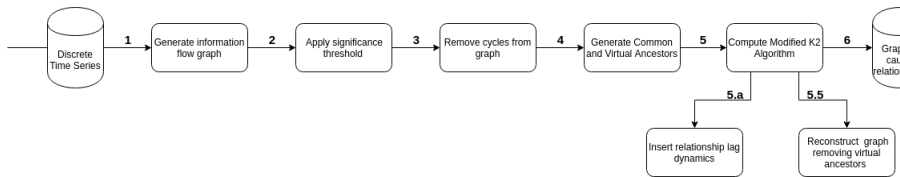


Figure 8: Proposed Methodology

Generation of graph of information flow

- Given a system with N variables, it computes the Transfer Entropy pairwise for the set of variables.
- For each pair of variables it computes the method h times.
- Chooses the greatest entropy value from the h computations.

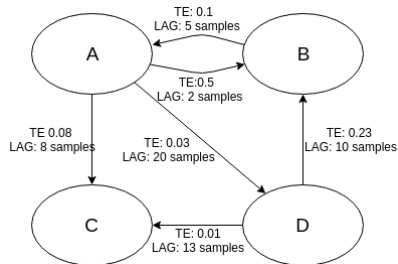
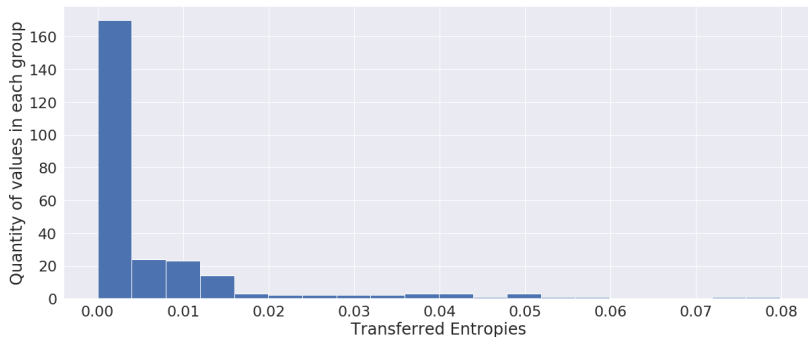


Figure 9: Example of output from stage 1

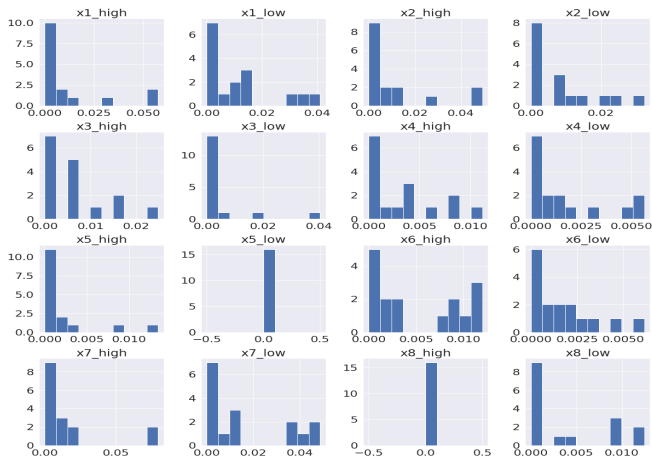
Application of Statistical Threshold

- Because the Transfer Entropy is calculated pairwise, it can produce a dense graph. Since due to the noise contained in the time series, the chance of a zero entropy becomes low.
- This stage aims, to the define a threshold for the entropy values based on the distribution of the computed values.

Application of Statistical Threshold



Application of Statistical Threshold



Removal of Cycles

- In order to prepare the graph to be used on K2, third stage does a removal of cycles.
- It consists of recursive search on the graph, in which the nodes “above” the node-cause, are marked as its ancestors. Therefore, the ancestors of a node-cause cannot be related to it as an effect.
- The removal is made based on the criterion of the higher information

Removal of Cycles

Removal of cycles

Generation of Common and Virtual Ancestors

What is a common and a virtual ancestor?

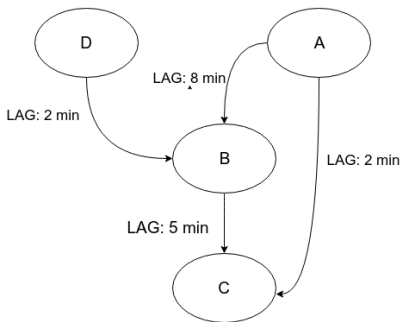


Figure 10: Ordinary Graph

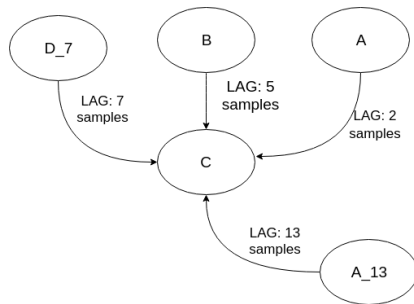


Figure 11: Modified Graph

Generation of Common and Virtual Ancestors

Prior Structure to K2





Modification and Computation of K2

Modification and Computation of K2

Case Study and Results

Resumo

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