

Sub-quadratic Markov tree mixture models for probability density estimation

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A simple idea

Proposition

*Develop density estimation techniques that could scale to very high-dimensional spaces, by exploiting the **Perturb & Combine** idea with probabilistic graphical models.*

Outline

- Background
- Our proposal
- Some results
- Conclusions and Further works

P&C principle in supervised learning

■ Principle :

(Bagging, Random forests, Extremely randomized trees)

How can we apply this idea to density estimation with Bayesian networks (BN)?

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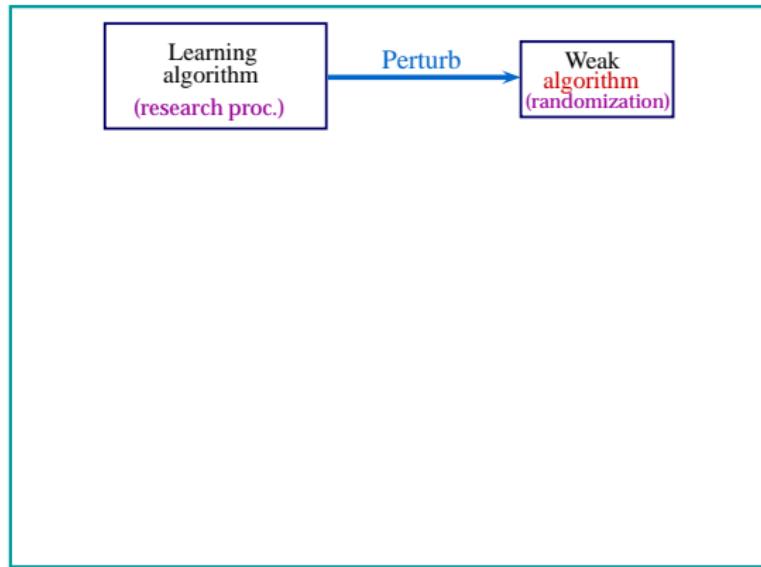
Learning
algorithm
(research proc.)

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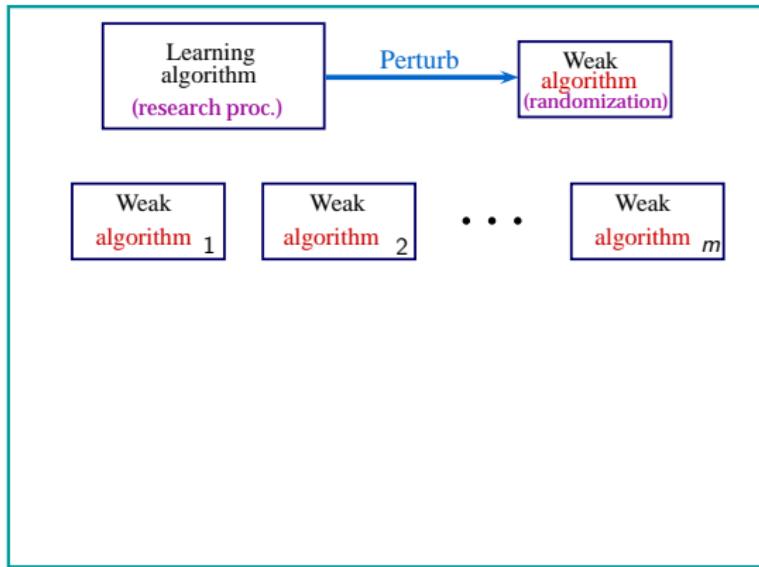


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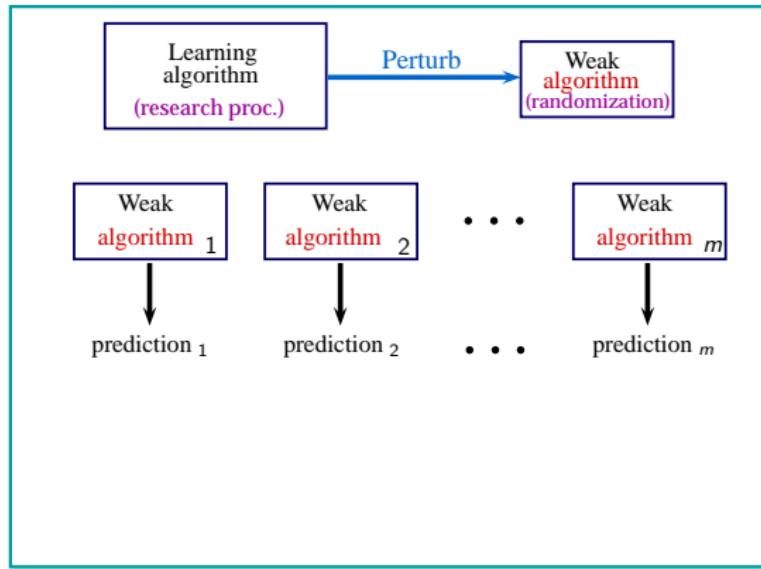


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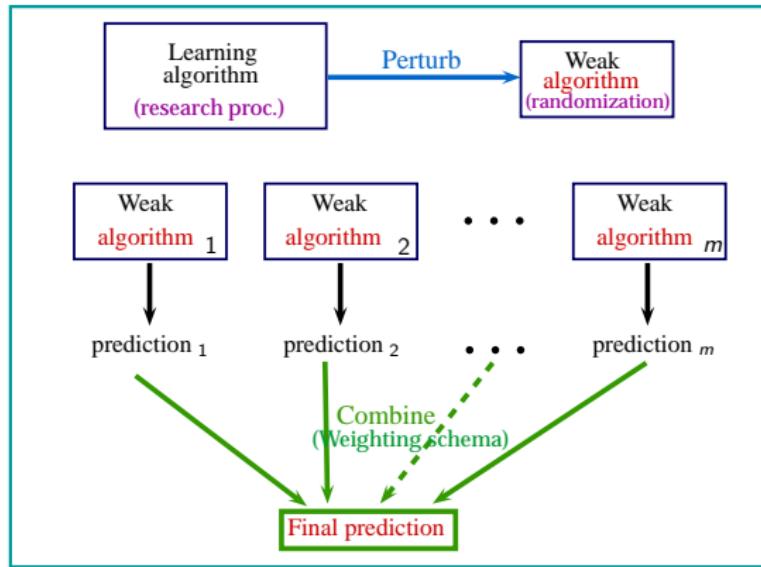


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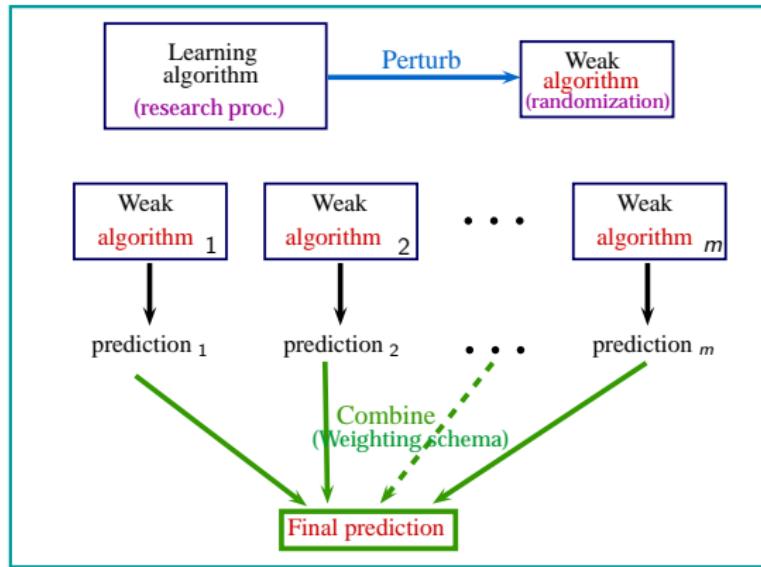


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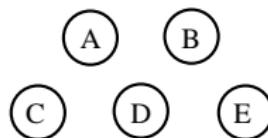
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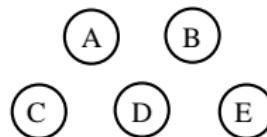
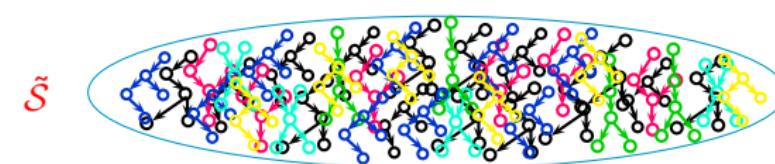


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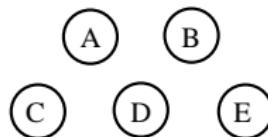
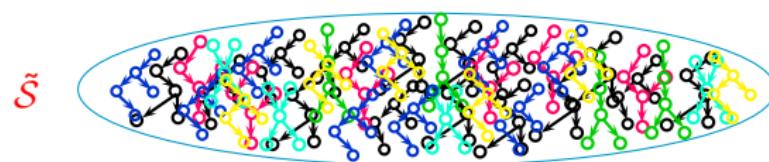
Density estimation with BN



Density estimation with BN

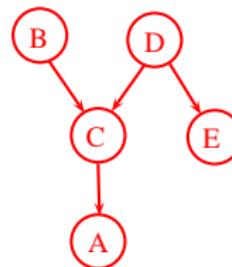
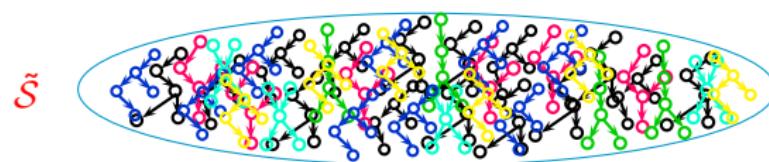


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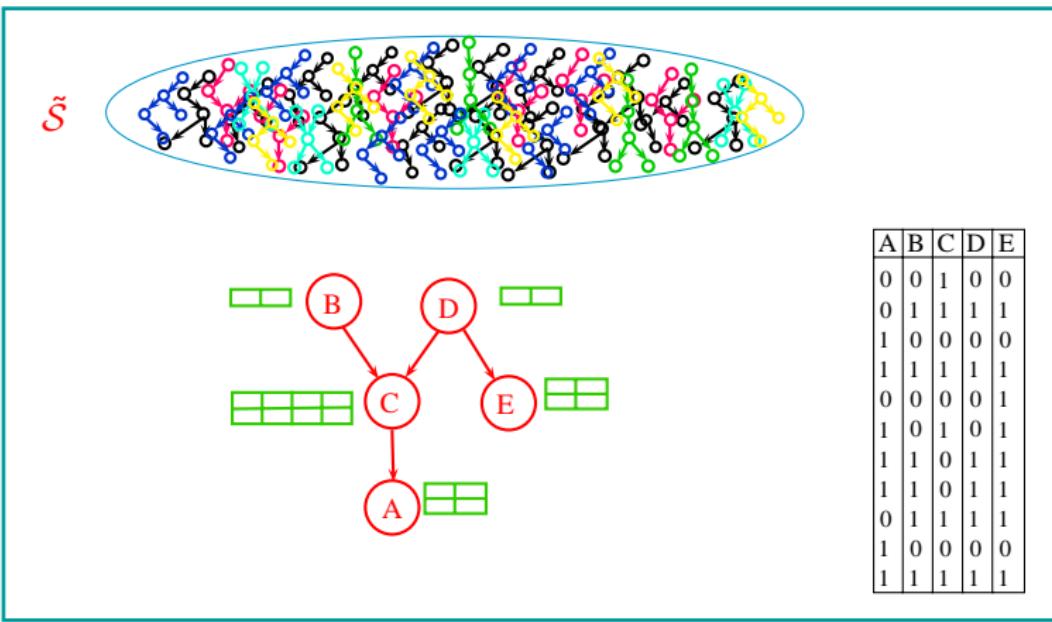
A	B	C	D	E
0	0	1	0	0
0	1	1	1	1
1	0	0	0	0
1	1	1	1	1
0	0	0	0	1
1	0	1	0	1
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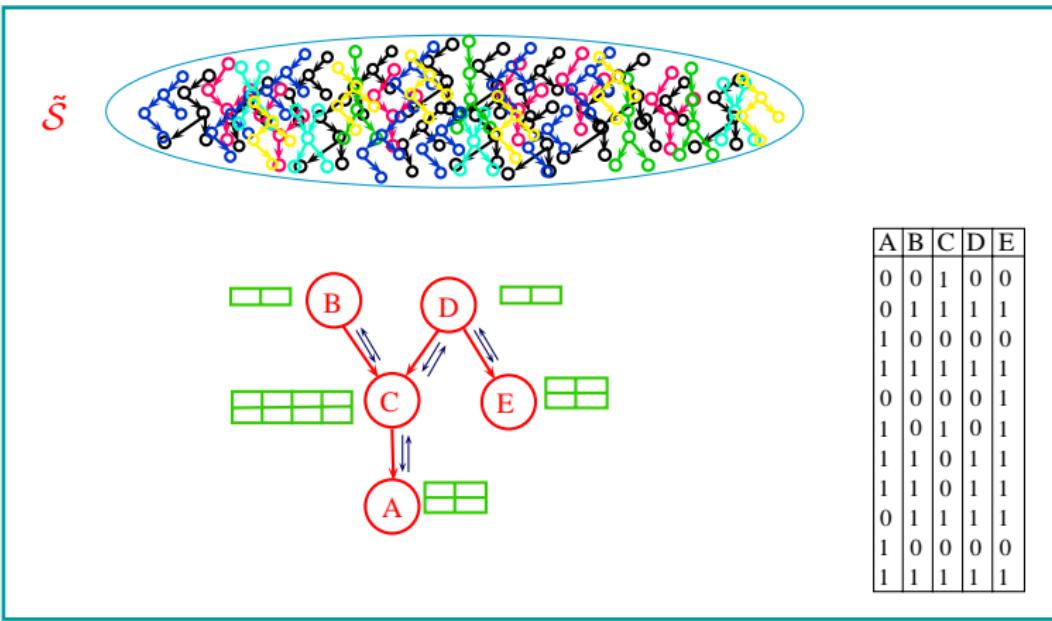


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1	0	0	0	0
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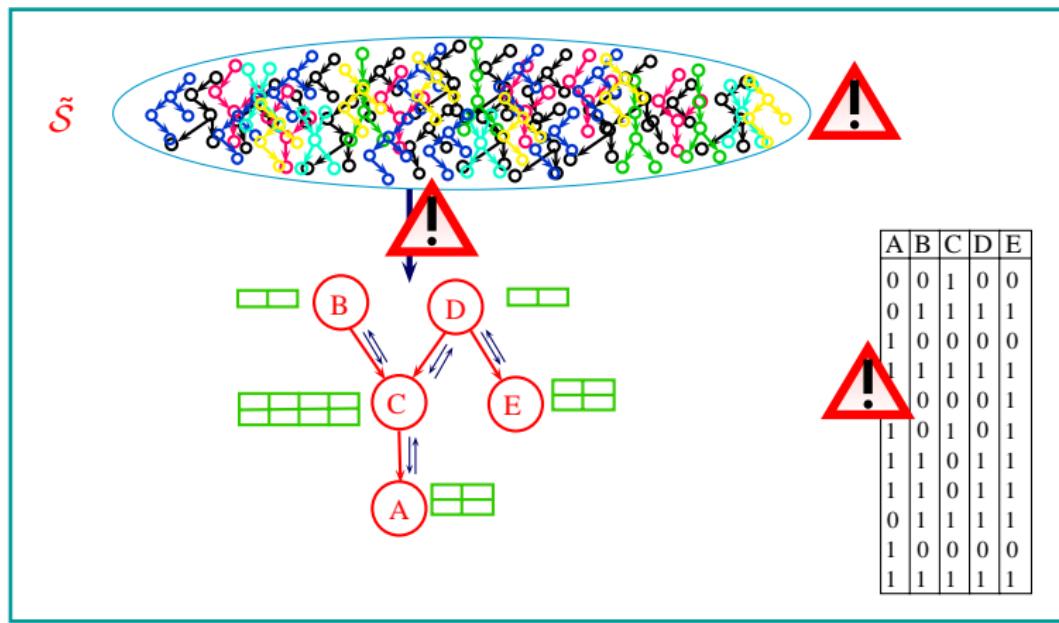
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Density estimation with BN



Density estimation with BN



Bayesian averaging

- Instead of searching for an optimal model (structure + parameters):
 - Assume prior probabilities over the space of structures
 - Determine posterior probabilities of each model given a dataset
 - approach the target distribution by averaging the different models weighted by their posterior probabilities
- **Caveats :** Exact Bayesian averaging over large space of models is not 'scalable'
⇒ requires to strongly constrain the space of structures



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Strategy

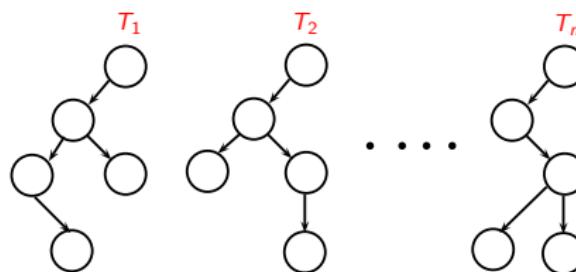
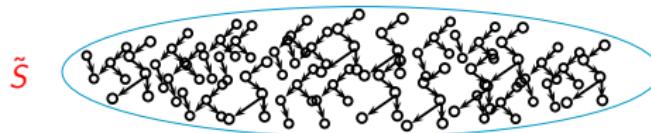
- Use simple spaces of graphical structures $\tilde{\mathcal{S}}$ (e.g. chains, trees, poly-trees etc.)
- Do not assume that target distribution is representable by one of these structures
- Rather, assume that target distribution may be approximated well by a mixture of a reasonable number of (S, θ^*) pairs,
 $S \in \tilde{\mathcal{S}}$

Generic algorithm principle



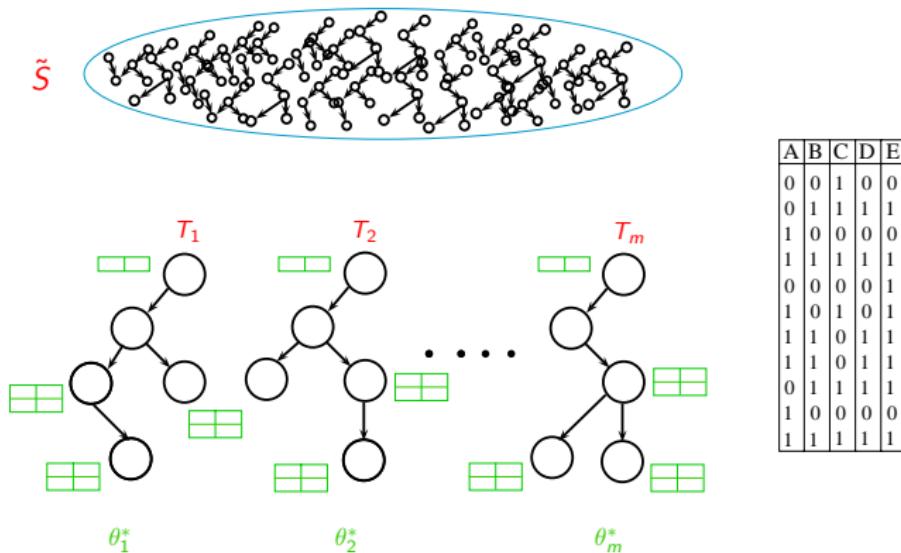
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0	1	1	1	1
1	0	0	0	0
1	1	1	1	1
0	0	0	0	1
1	0	1	0	1
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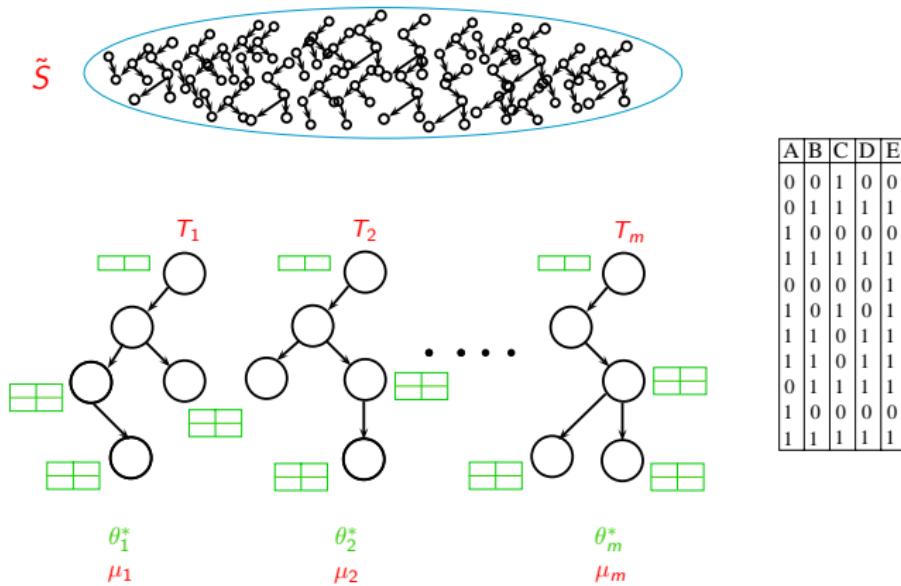


A	B	C	D	E
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0	1	1	1	1
1	0	0	0	0
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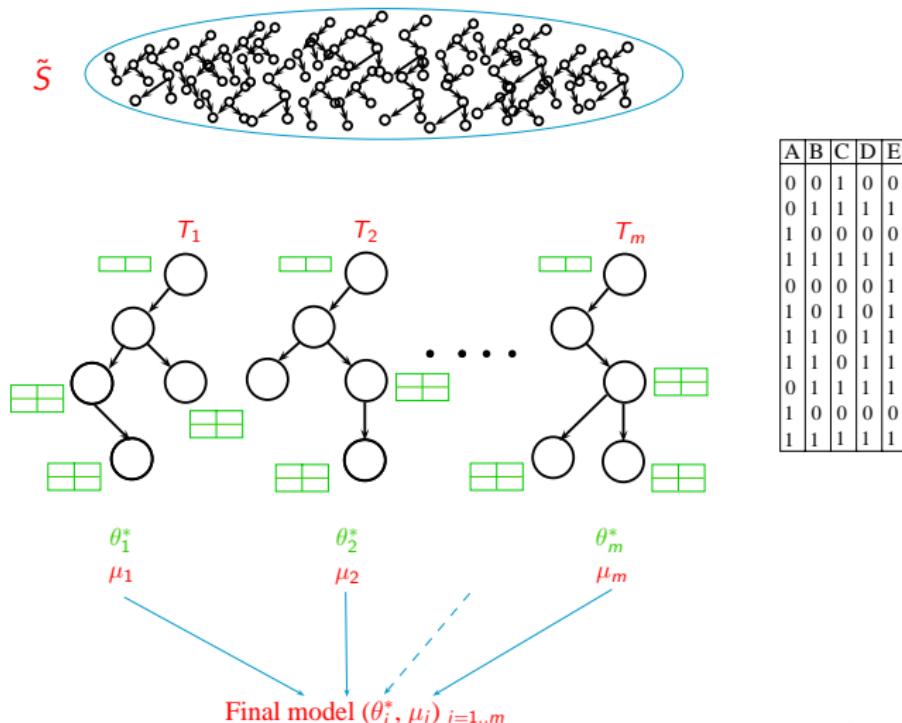
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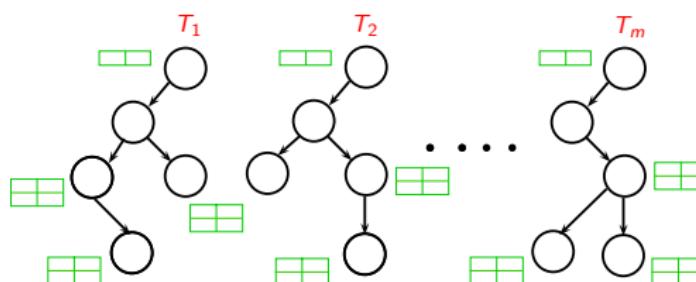
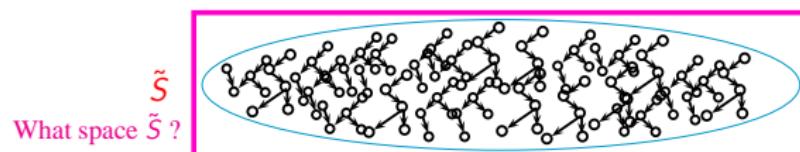
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Generic algorithm principle



Degrees of freedom

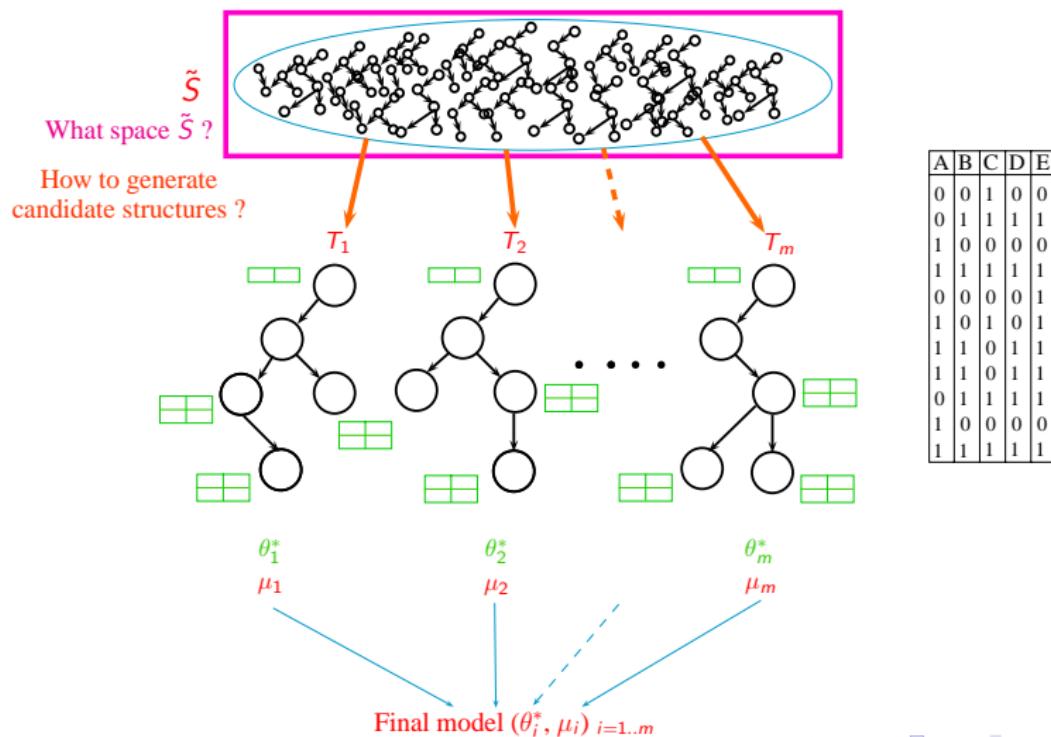


$$\begin{array}{l} \theta_1^* \\ \mu_1 \end{array} \quad \begin{array}{l} \theta_2^* \\ \mu_2 \end{array} \quad \dots \quad \begin{array}{l} \theta_m^* \\ \mu_m \end{array}$$

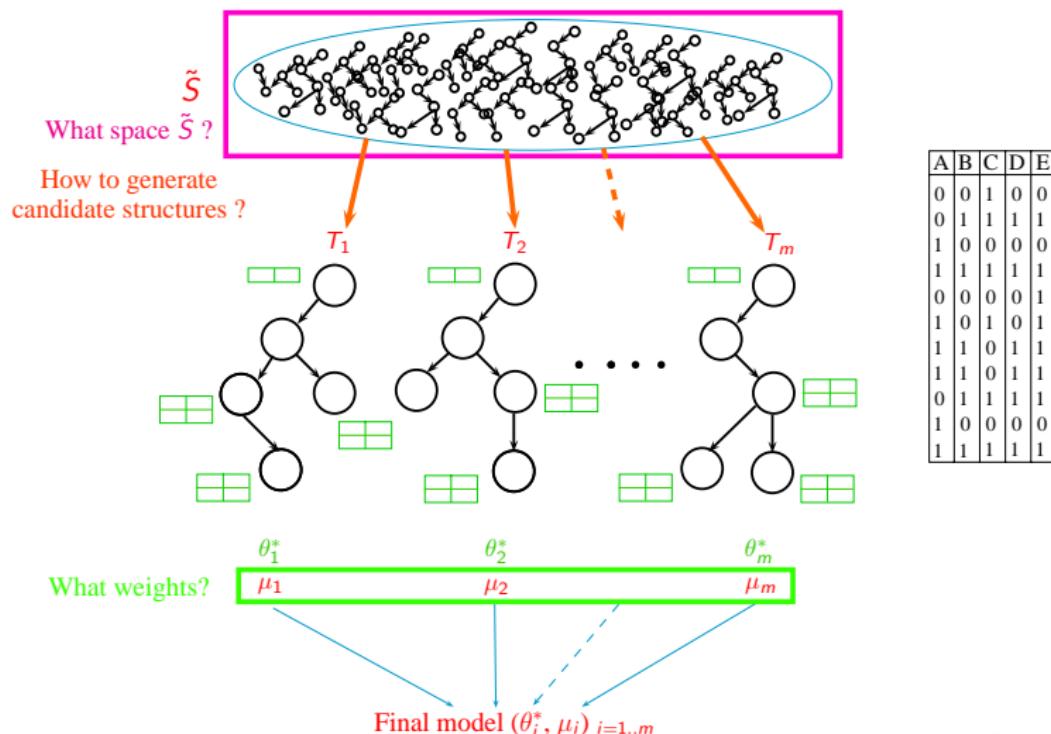
Final model $(\theta_i^*, \mu_i)_{i=1..m}$

A	B	C	D	E
0	0	1	0	0
0	1	1	1	1
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1	1	1	1	1
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Degrees of freedom



Degrees of freedom



Why Markov tree space ?

- Polytrees, although more expressive, do not yield more accurate ensemble models than undirected trees [Ammar et al.2008].
- Markov trees
 - (-) "poor" independency models

But

- (+) Inference and parameters learning are scalable (linear complexity)
- (+) Uniform sampling of trees (linear complexity)
- (+) Optimal tree structure learning is polynomial (MWST)

How to generate candidate structures ?

- Random uniform sampling [Ammar et al.2009] : linear complexity $O(n)$ and good results
- Build the optimal tree (MWST) over a bootstrap replica of original dataset D [Ammar et al.2009] : quadratic complexity $O(n^2 \log(n^2))$ and better results

Goal : How can we improve the complexity of our quadratic methods and keep the same accuracy ?

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Goal : *How can we improve the complexity of our quadratic methods and keep the same accuracy ?*

Principle of MWST

- Step 1 : fill a $(n \times n)$ mutual information matrix
complexity $O(n^2)$
- Step 2 : build optimal tree (Kruskal)
complexity $E \log(E)$; $E = n^2$
- Step 3 : learn parameters
linear complexity

⇒ **Solution** : reduce the first step complexity by applying the Perturb & Combine principle

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sub-quadratic research heuristics

■ Partial matrix M_I

	X			X		
		X				
			X			
M_I_K		X		X	X	
	X			X		
		X				
	X		X			
				X		

■ $K = n \log(n) \Rightarrow$

Step 1: $n \log(n)$

Etape 2: $n \log(n) \log(n \log(n))$

\Rightarrow Total complexity = $n \log(n) \log(n \log(n))$

↖↖ quadratic and \propto quasilinear

sub-quadratic research heuristics

■ Partial matrix MI

MI_K	X		X	
		X		
			X	
	X		X	X
		X		
	X			
			X	

$K = \text{number of considered terms in } MI$

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		X				
			X			
				X	X	
	X				X	
		X				
	X		X			

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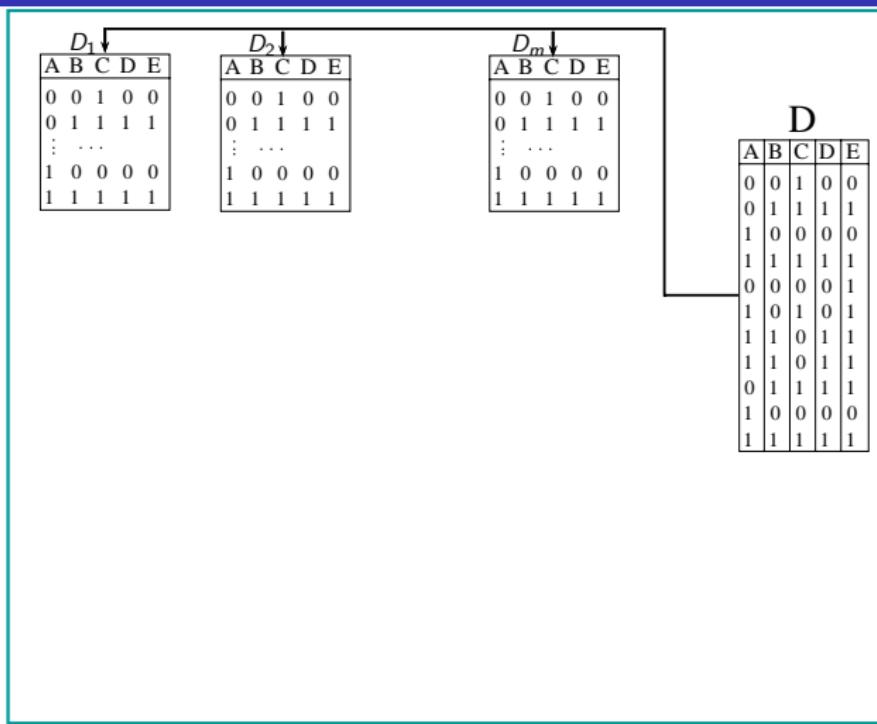
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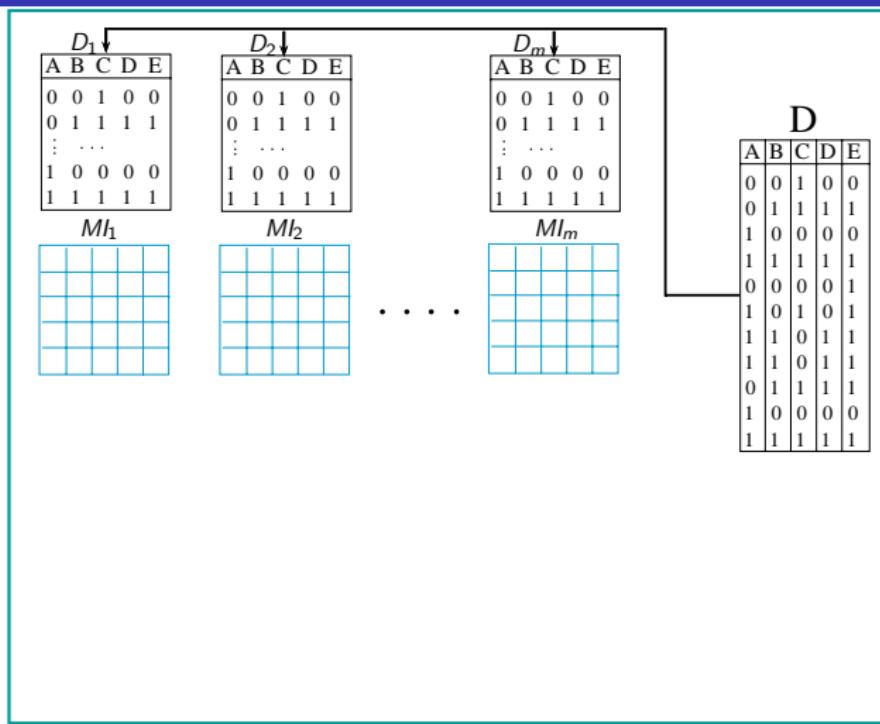
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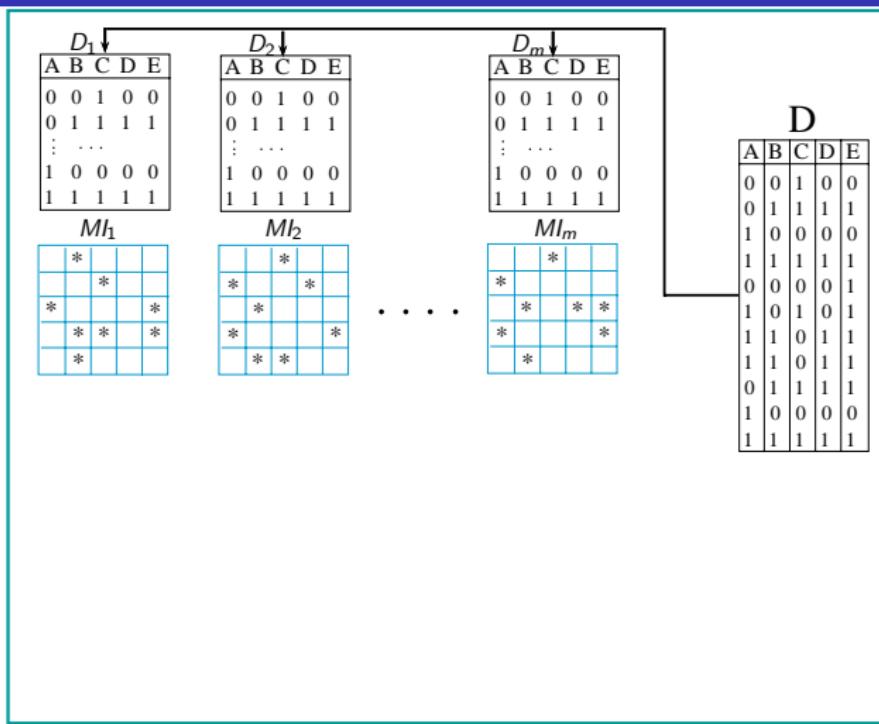
General algorithm : Naive Heuristic



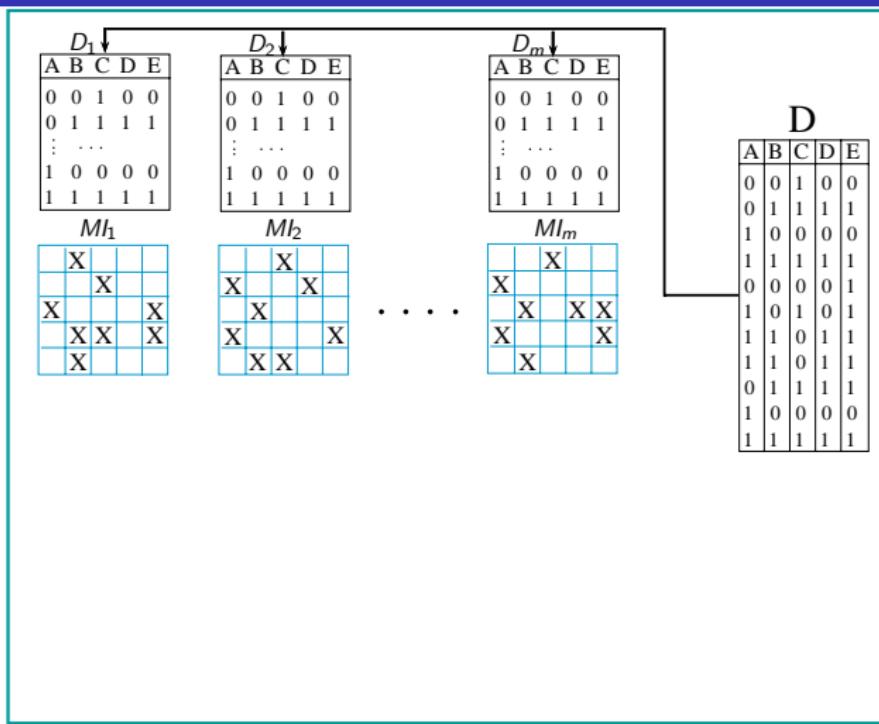
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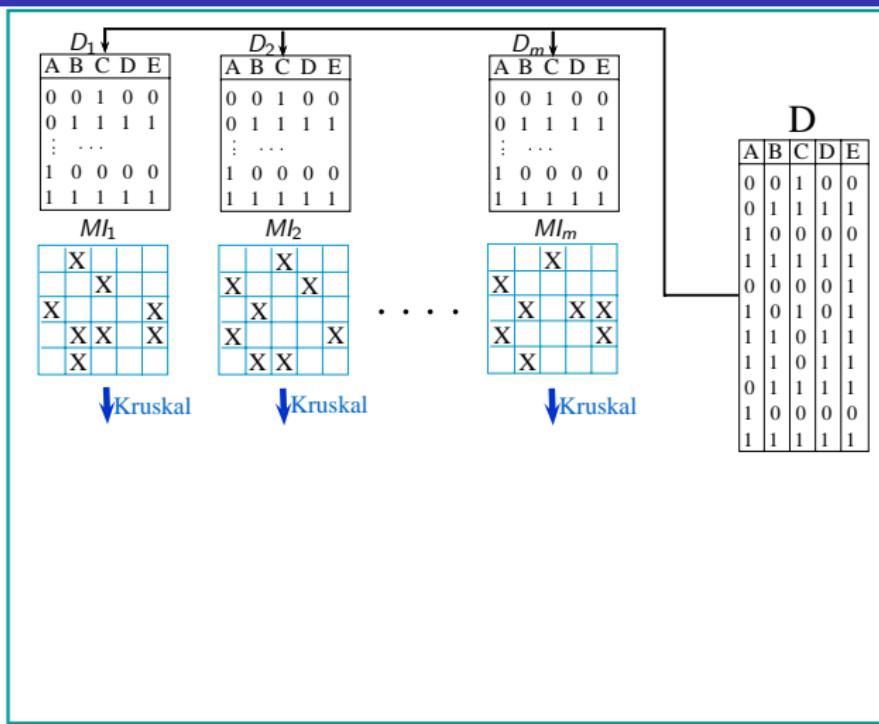
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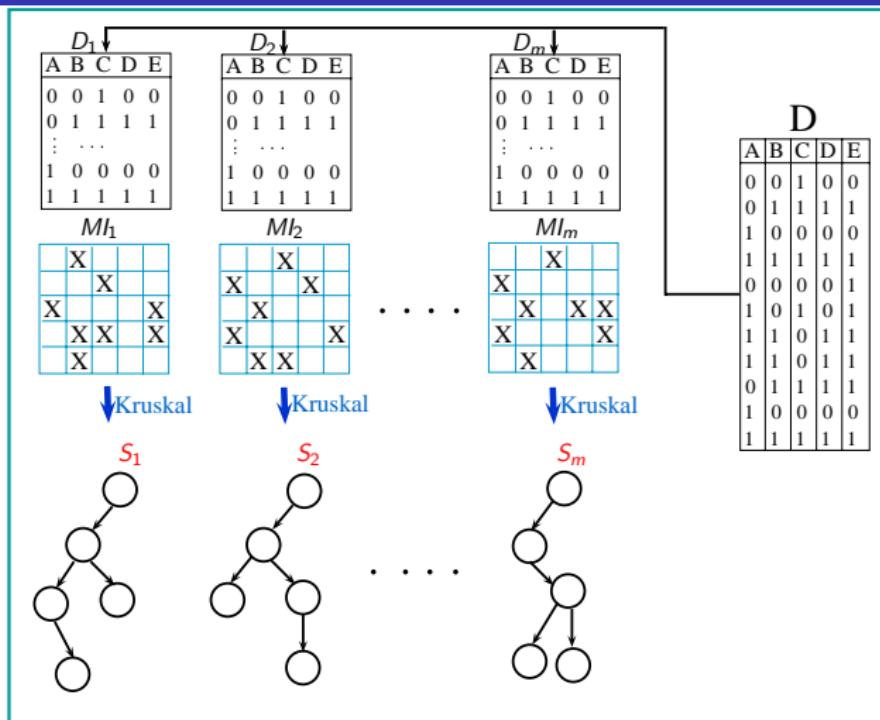
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General algorithm : Inertial Heuristic

$D_1 \downarrow$				
A	B	C	D	E
0	0	1	0	0
0	1	1	1	1
⋮	...			
1	0	0	0	0
1	1	1	1	1

D

A	B	C	D	E
0	0	1	0	0
0	1	1	1	1
1	0	0	0	0
1	1	1	1	1
0	0	0	0	1
1	0	1	0	1
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M_{I_1}				

D				
A	B	C	D	E
0	0	1	0	0
0	1	1	1	1
1	0	0	0	0
1	1	1	1	1
0	0	0	0	1
1	0	1	0	1
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0	0	1	0	0
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⋮	⋮	⋮	⋮	⋮
1	0	0	0	0
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M_{I_1}				
	*	*	*	
*	*			
			*	
*			*	
*	*	*	*	

D				
A	B	C	D	E
0	0	1	0	0
0	1	1	1	1
1	0	0	0	0
1	1	1	1	1
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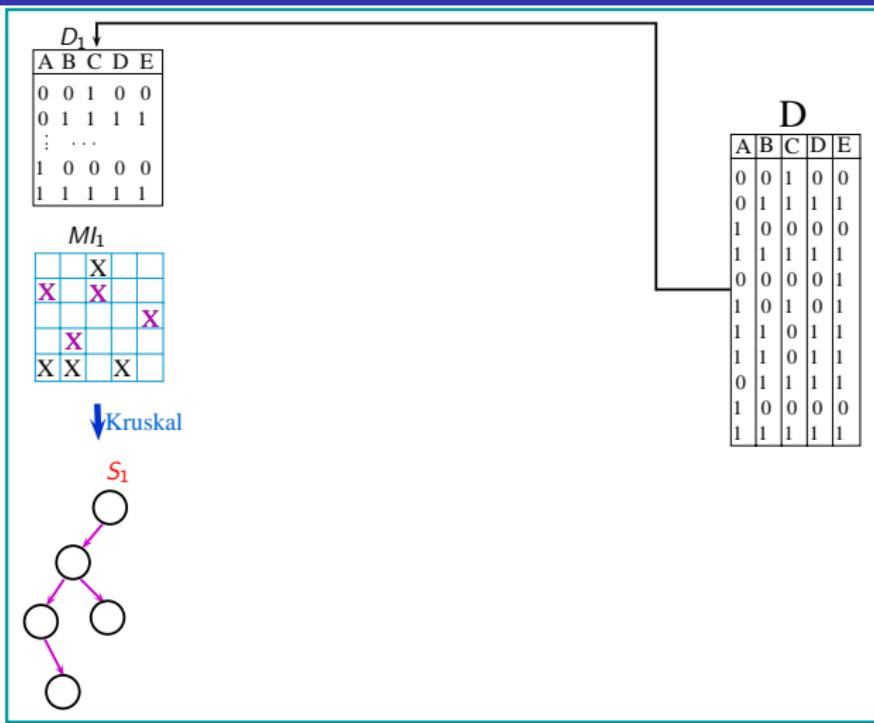
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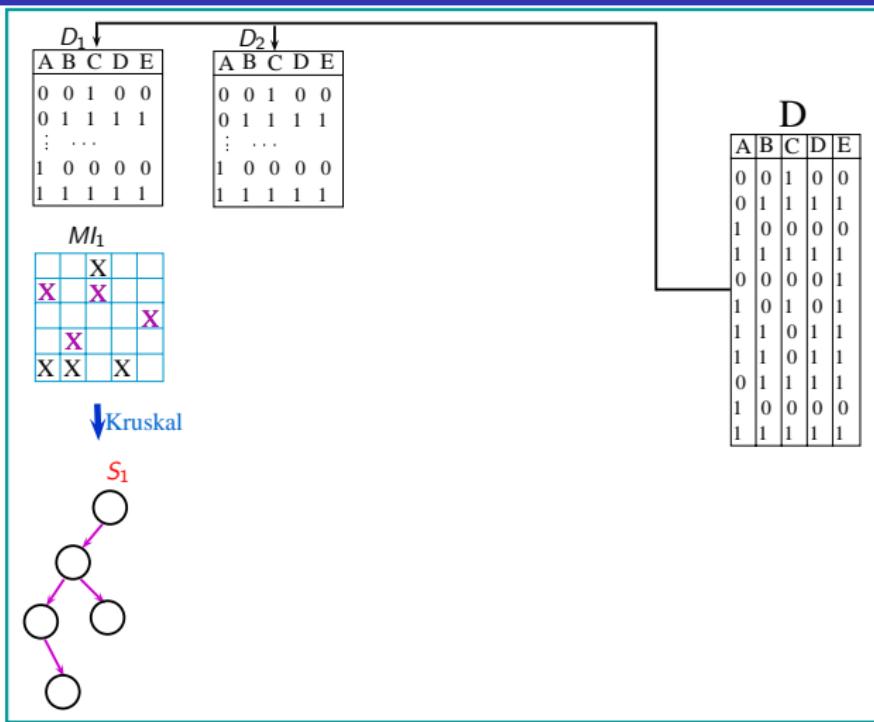
M_{I_1}				
	X			
X	X			
		X		
X	X	X		

D				
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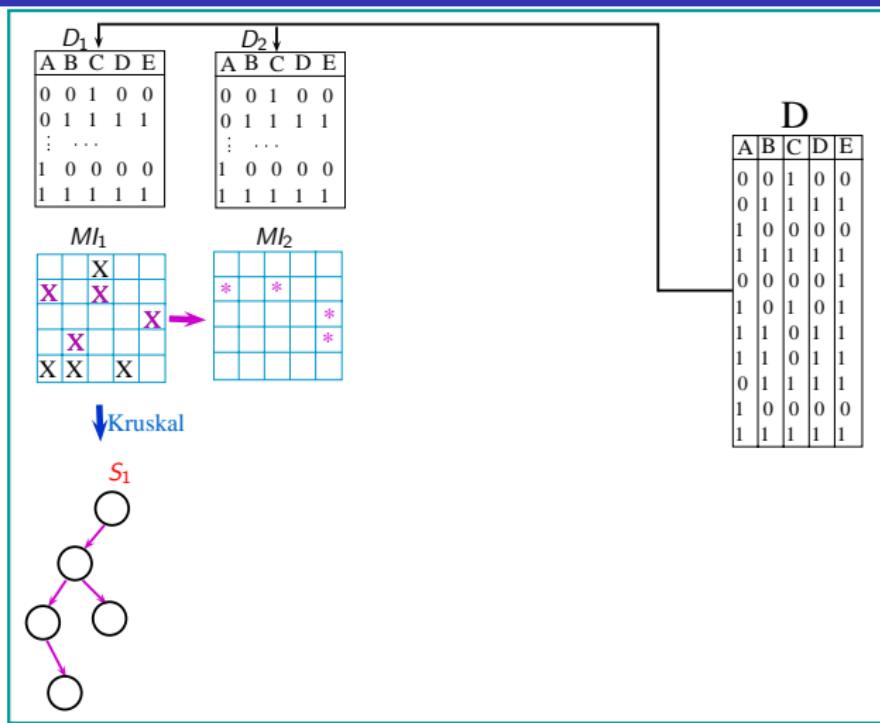
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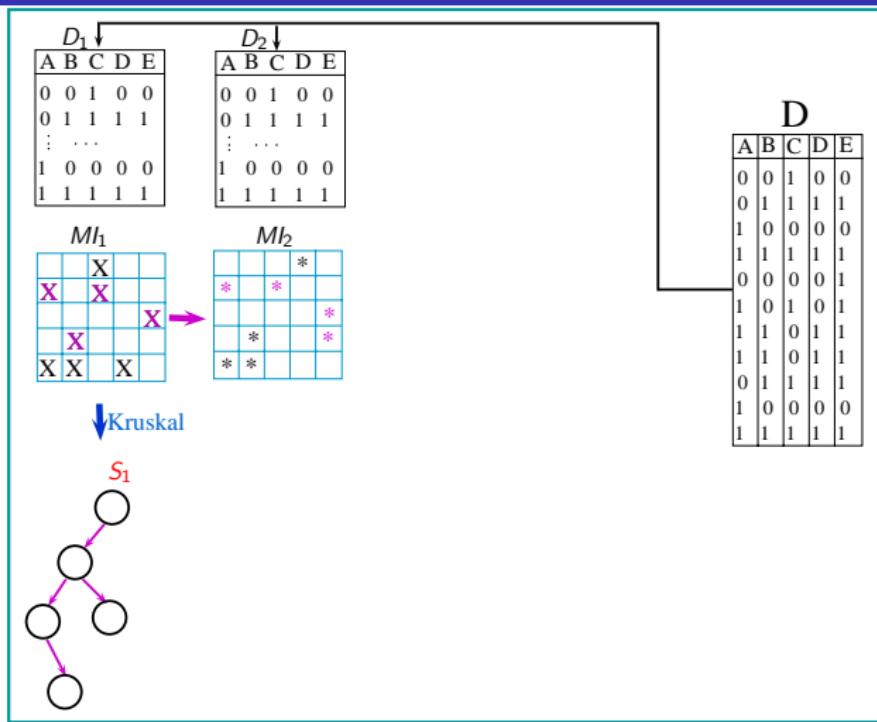
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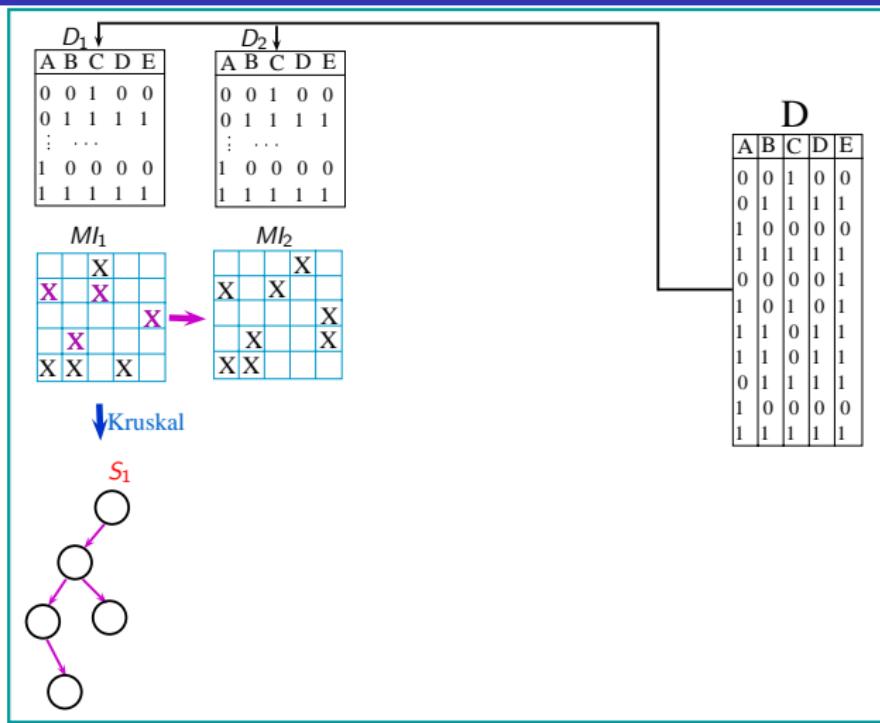
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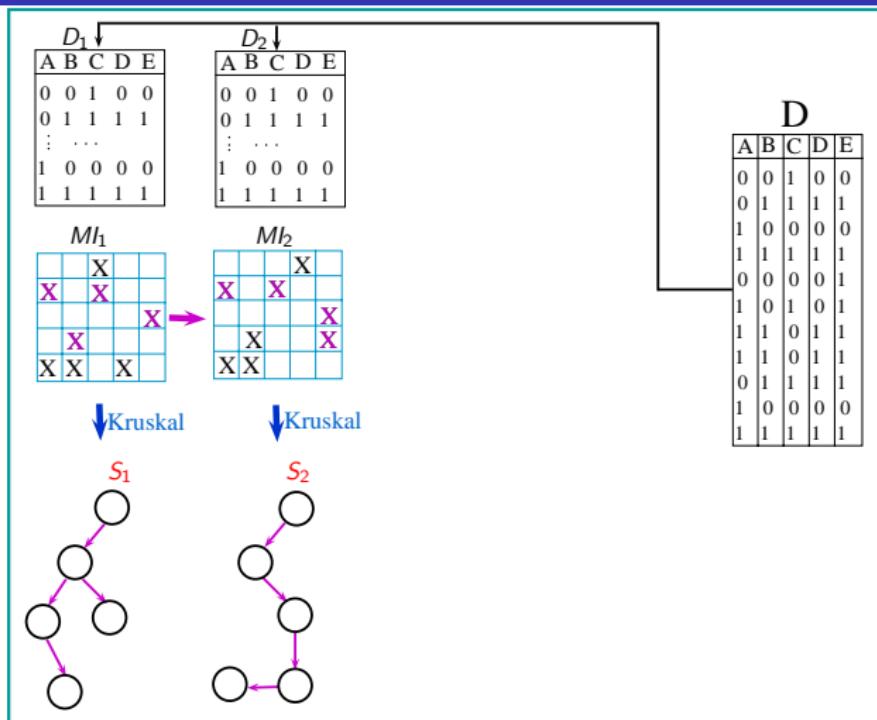
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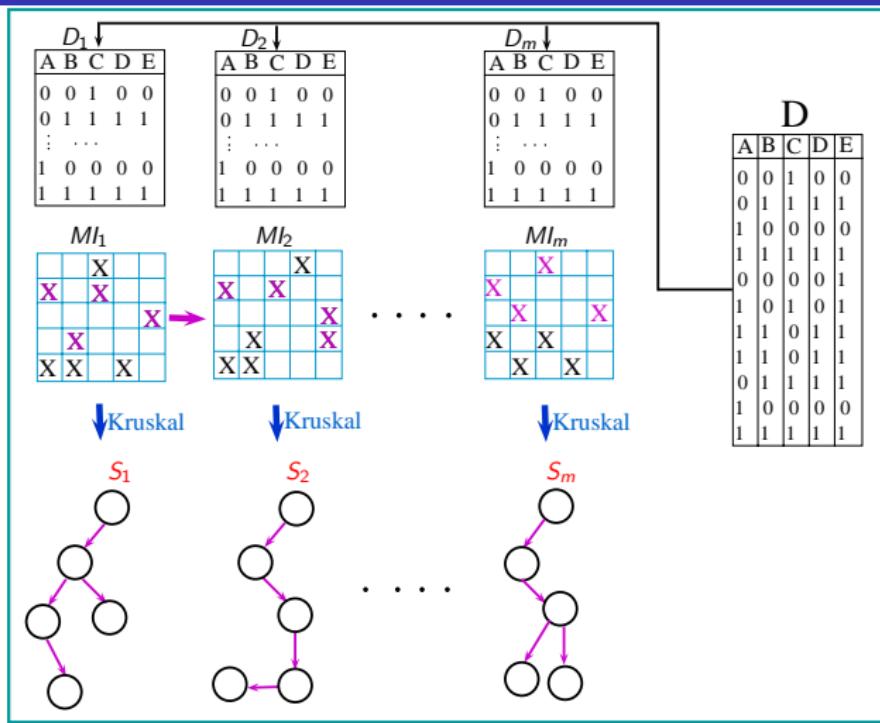
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Experimental protocol

■ Test problems

- 1000 binary variables;
- 10 target distributions : DAG (1000 vars)
- 10 repeated experiments, for $|D| = 1000$

■ Algorithm variants

- Mixtures with growing sizes ($m = 1, 10, 20, \dots, 150$) of Markov trees, two variants of the generation of these trees :
 - Subquadratic Heuristics over the initial dataset D
 - Subquadratic Heuristics over a bootstrap replica of D
- parameter learning : MAP with uniform priors
- Different weighting schemas :
 - Uniform weighting (ie. $\mu_i = 1/m$)
 - Bayesian averaging (ie. $\mu_i \propto \text{score}_{BDeu}$)

■ Accuracy evaluation

- asymmetric approached Kullback-Leibler divergence
- Accuracy is better when \hat{KL} divergence is lower.

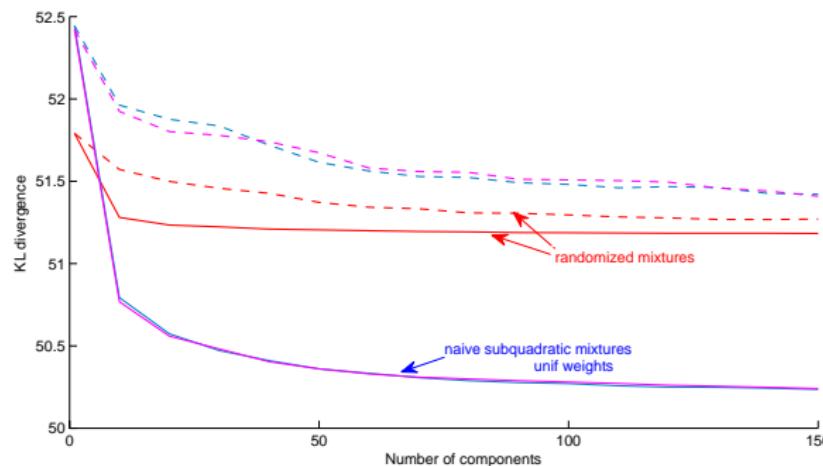
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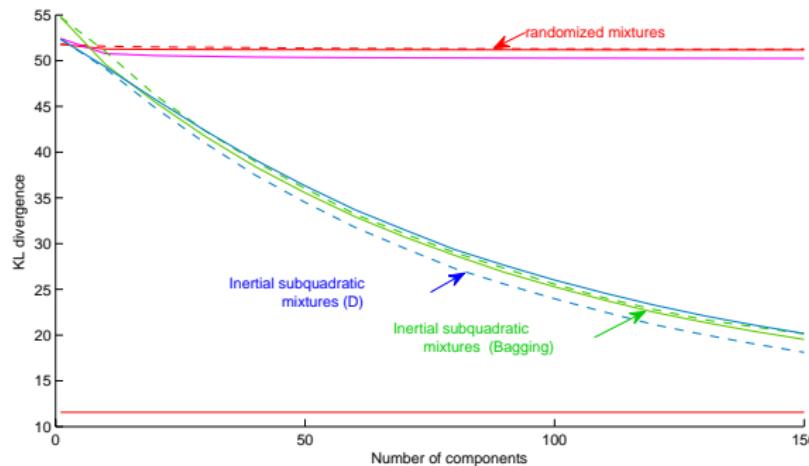
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 - Accuracy is better when \hat{KL} divergence is lower.

Naive Heuristic results



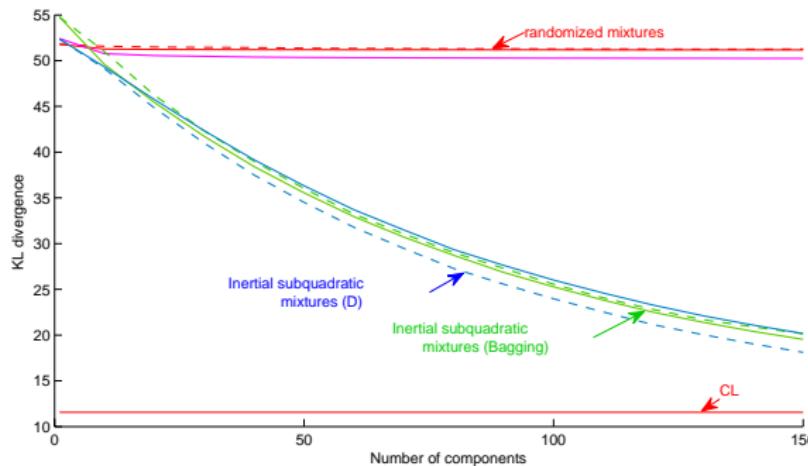
- Our naive subquadratic mixtures are clearly outperforming the randomized methods with uniform weighting

Inertial Heuristic results



- Our inertial methods outperform the randomized ones (Linear complexity)
- Our inertial methods approach the base method CL (Quadratic complexity)

Inertial Heuristic results



- Our inertial methods outperform the randomized ones (Linear complexity)
- Our inertial methods approach the base method CL (Quadratic complexity)

Conclusion

- Our subquadratic mixtures (quasilinear) outperform the randomized methods (linear)
- Our inertial methods approach the base method CL (quadratic)
- Bagging does not allow an improvement in the estimation quality

Further work

- Improve our methods : linear approximation of the optimal tree [Chazelle2000]
- Comparison with scalable optimal structure learning algorithms (sparse candidate [Friedman et al.1999], MMHC [Tsamardinos et al.2006] ...)
- Consider sequential methods of combination (Boosting, MCMC)

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