Statistical Relational Al & Robustness

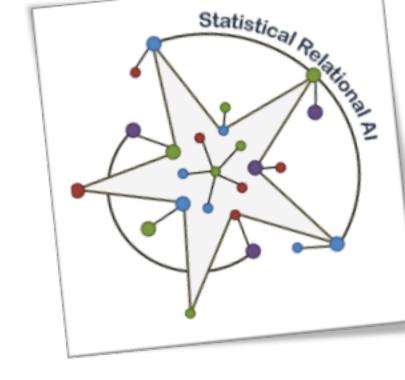
Workshop on Robust AI for High-Stakes Applications

@KI2022

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Agenda

- Probabilistic relational models (PRMs)
 - Application example
 - Semantics
 - Query answering / basic inference
- Lifting Algorithms for More Robust Inference
 - Cluster trees for efficient multi-query inference
 - Adaptive inference in cluster trees
 - Changing domains
 - Keeping inference going over time
- Summary



Application: Epidemics

- Atoms: Parameterised random variables = PRVs
 - With logical variables
 - E.g., X, M
 - Possible values (domain):

$$D(X) = \{alice, eve, bob\}$$

$$D(M) = \{injection, tablet\}$$

- Nat(D)Man(WEpid Travel(X)Treat(X, M

Sick(X)

- With range
 - E.g., Boolean
 - range(Travel(X)) r(Travel(X))

 $Nat(D) = natural\ disaster(D)$ $Man(W) = man - made \ disaster(W)$

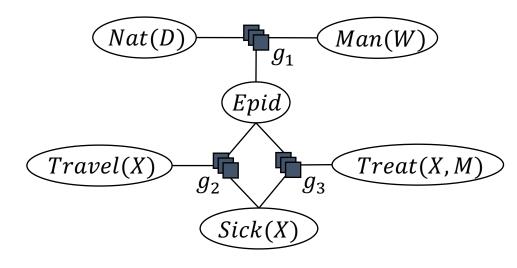


Encoding the Joint Distribution

- Factors with PRVs = parfactors
 - (Graphical) Model G
 - E.g., g_2

Travel(X)	Epid	Sick(X)	g_2
false	false	false	5
false	false	true	0
false	true	false	4
false	true	true	6
true	false	false	4
true	false	true	6
true	true	false	2
true	true	true	9

Sparse encoding of joint distribution







Factors

Grounding

• E.g., $gr(g_2) = \{f_2^1, f_2^2, f_2^3\}$

		Travel(eve)	Epid	Sick(eve)	f_{2}^{1}				
Travel(X) E	nid	false	false	false	5	(bob)	Epid	Sick(bob)	f_{2}^{3}
Travel(alice)	Еріс	false	false	true	0	e	false	false	5
false	fals	false	true	false	4	e	false	true	0
false	fals	false	true	true	6	e	true	false	4
false	true	true	false	false	4	e	true	true	6
false	true	true	false	true	6	e	false	false	4
true	fals	true	true	false	2	e	false	true	6
true	fals	true	true	true	9	e	true	false	2
true	true	e false	2		tru	ie	true	true	9
true	true	e true	9						

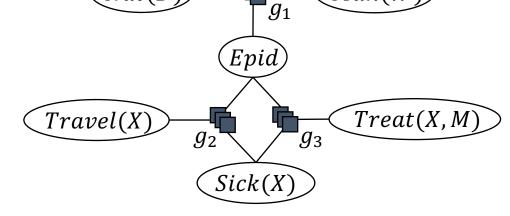


Semantics of a PRM

• Joint probability distribution P_G by grounding, multiplying all

grounded factors, and normalising the result

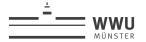
$$P_G = \frac{1}{Z} \prod_{f \in gr(G)} f$$



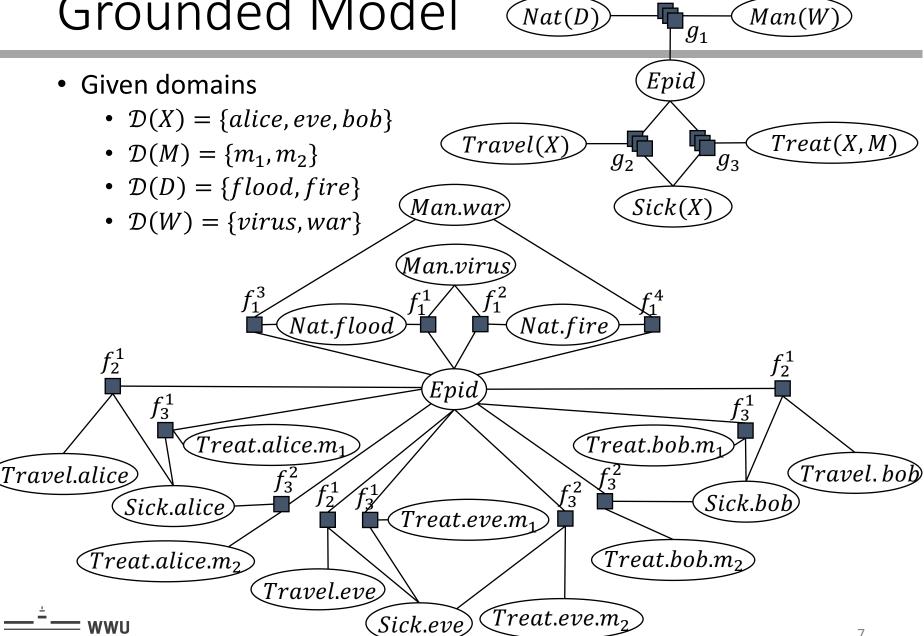
Man(W)

Nat(D)

$$Z = \sum_{v \in r(rv(gr(G)))} \prod_{f \in gr(G)} f_i(\pi_{rv(f_i)}(v))$$



Grounded Model



PRMs and Variants

- Probabilistic Relational Models (raw PRMs)
 [Poole 03, Taghipour et al. 13, B & Möller 16-19, Gehrke, B & Möller 18-19]
- Markov Logic Networks (MLNs) [Richardson & Domingos 06]
 - Use logical formulas to specify potential functions
- Probabilistic Soft Logic (PSL) [Bach et al. 17]
 - Use density functions to specify potential functions
- Based on grounding semantics [Sato 95, Fuhr 95]



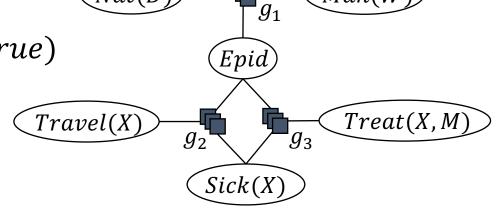
Queries

Marginal distribution

Avoid groundings!

Nat(D

- *P*(*Sick*(*eve*))
- $P(Travel(eve,) Treat(eve, m_1))$
- Conditional distribution
 - P(Sick(eve)|Epid)
 - P(Epid|Sick(eve) = true)
- Assignment queries
 - MPE
 - MAP



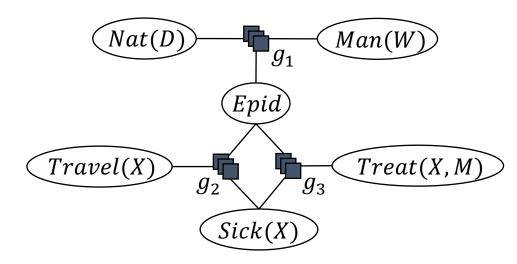


Man(W)

QA: Lifted Variable Elimination (LVE)

 Eliminate all variables not appearing in query [Poole 03, de Salvo Braz et al. 05, 06, Milch et al. 08, Taghipour et al. 13, 13a]

- Lifted summing out
 - Sum out
 representative
 instance as in
 propositional
 variable elimination
 - 2. Exponentiate result for isomorphic instances

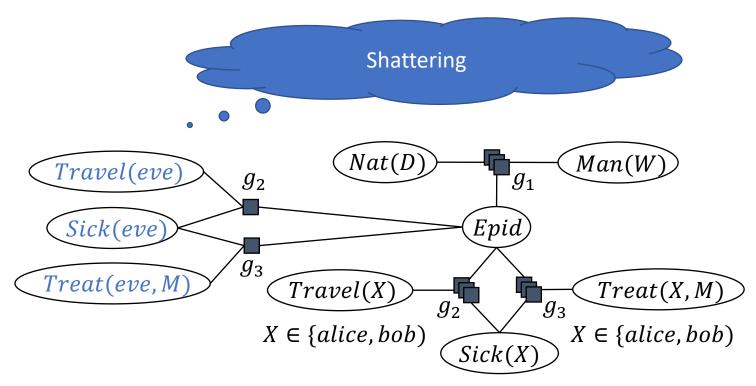


Avoid groundings!



QA: LVE in Detail

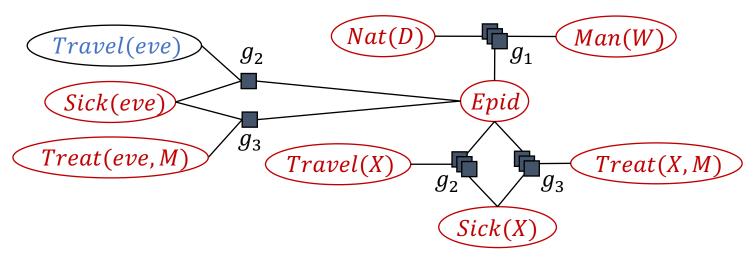
- E.g., marginal
 - P(Travel(eve))
 - Split atoms P(...,X,...) w.r.t. eve if eve in $\mathcal{D}(X)$





QA: LVE in Detail

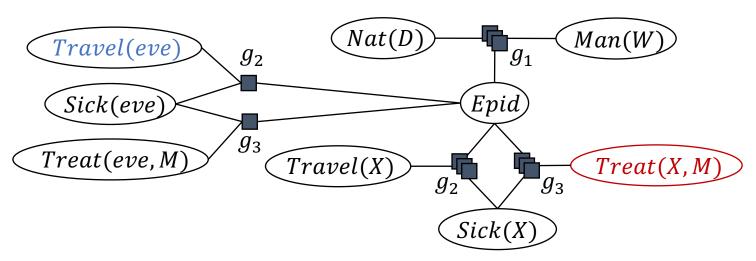
- E.g., marginal
 - P(Travel(eve))
 - Split atoms P(...,X,...) w.r.t. eve if eve in $\mathcal{D}(X)$
 - Eliminate all non-query variables





QA: LVE in Detail

- Eliminate Treat(X, M)
 - Appears in only one $g: g_3$
 - Contains all logical variables of g_3 : X, M
 - For each X constant: the same number of M constants
 - ✓ Preconditions of lifted summing out fulfilled, lifted summing out possible



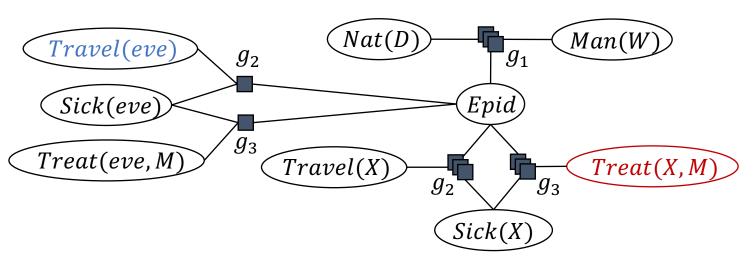


LVE in Detail: Lifted Summing Out

- Eliminate Treat(X, M) by lifted summing out
 - 1. Sum out representative
 - 2. Exponentiate for indistinguishable objects

Only here, domain size comes into play → no change in structure / equation if domain size changes

$$\left(\sum_{t \in r(Treat(X,M))} g_3(Epid = e, Sick(X) = s, Treat(X,M) = t)\right)^{|M|}$$





Tractability

- Given a model that allows for lifted calculations
 - I.e., no groundings during solving an instance of the problem
- Solving an instance of the problem is possible in time polynomial in domain sizes
 - → The query answering algorithm is domain-lifted
- An query answering problem is tractable
 - when it is solved by an efficient algorithm, running in time polynomial in the number of random variables
- Assume that the number of random variables is characterised by domain sizes
 - Then, solving a query answering problem is tractable under domain-liftability
 - Runtime might still be exponential in other terms
 - More general results by Niepert and Van den Broeck (2014)



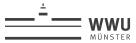
Agenda

- Probabilistic relational models (PRMs)
 - Application example
 - Semantics, static vs. dynamic behavior
 - Query answering / basic inference
- Algorithms for More Robust Inference
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Cluster Trees for Efficient Multi-query Inference

Algorithms for More Robust Inference



Statistical Rela

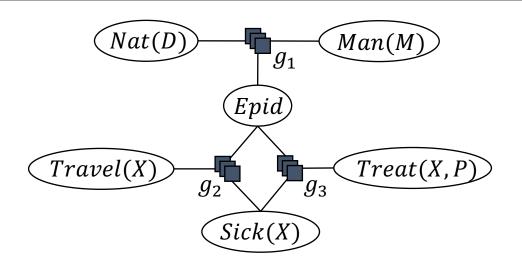
Many Queries: LJT

Set of queries

- P(Travel(eve))
- *P*(*Sick*(*bob*))
- $P(Treat(eve, m_1))$
- P(Epid)
- *P*(*Nat*(*flood*))
- P(Man(virus))
- Combinations of variables

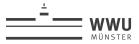
Under evidence

- Sick(X') = true
- $X' \in \{alice, eve\}$



Challenges:

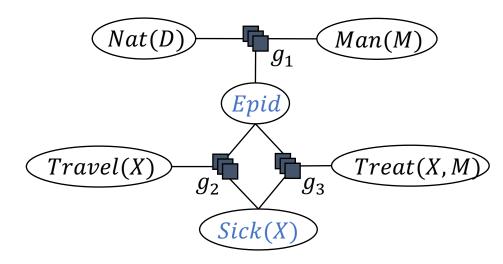
- Do not start from scratch for every query
- Support QA on subset of atoms
- Avoid groundings
- Cluster tree based on (conditional) independences



Solution: Submodels

- Identify submodel sufficient for query
 - Find PRVs that make submodel independent from remaining model
 - Separator
 - "Query" over separator collects all influences of remaining model on PRVs in submodel
 - PRVs of submodel = parcluster

Lauritzen and Spiegelhalter (1988), Shafer and Shenoy (1989), Jensen et al. (1990), Braun and Möller (2016)

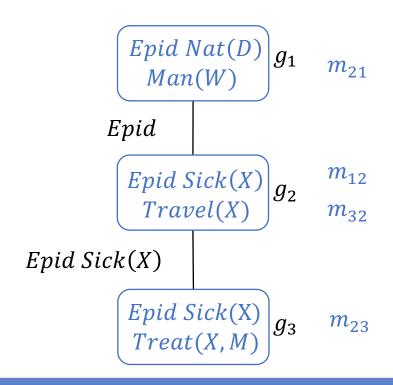




Solution: Submodels

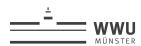
- Network of submodels with separators
 - Recursive "queries" to make submodels independent from each other
 - (First-order) Junction tree
 - Acyclic, running intersection property
- Recursive queries from each node
 - Arrange queries using dynamic programming
 - Also known as message passing

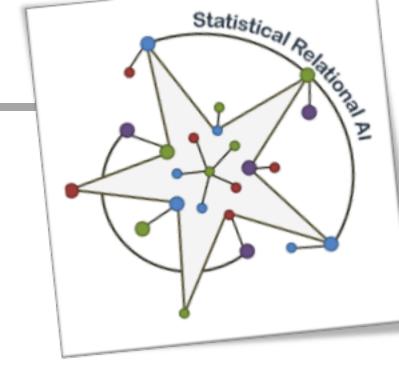
Lauritzen and Spiegelhalter (1988), Shafer and Shenoy (1989), Jensen et al. (1990), Braun and Möller (2016)



Answer queries on subtree over the query terms

P Use middle cluster for Pig(Sick(eve)ig)





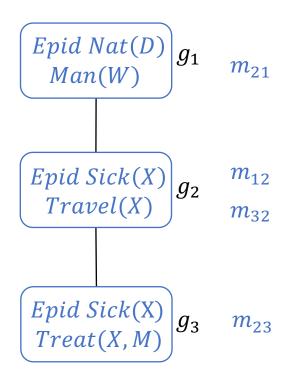
Adaptive Inference in Cluster Trees

Algorithms for More Robust Inference



Adaptive Inference

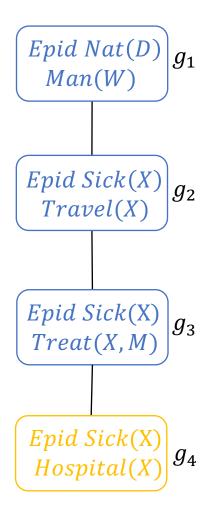
- After changes in queries, evidence, model: Avoid starting from scratch to fast reach the point of answering queries again
 - → adaptive inference
- Small, local model changes may preserve much of tree
 - If only local changes, up to half of messages still valid
 - Only resend messages if local model or incoming information changed

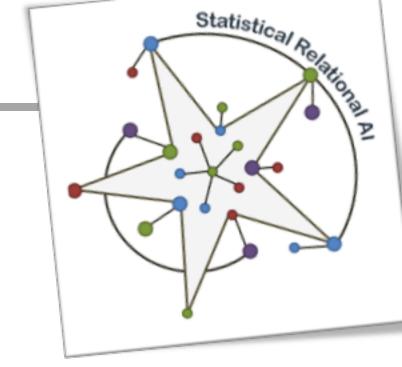




Adaptive Inference: Changes

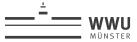
- Queries: no change
- Evidence: changes local models
 - New observations incoming
- Model
 - Potentials: changes local models
 - Domain sizes: changes local models
 - Nice property of relational models:
 No effect on model structure!
 - E.g., more people in dom(X)
 - Propositional models: number of variables changes, which changes the tree structure
 - Parfactors (addition, deletion): changes tree structure





Changing Domains

Algorithms for More Robust Inference



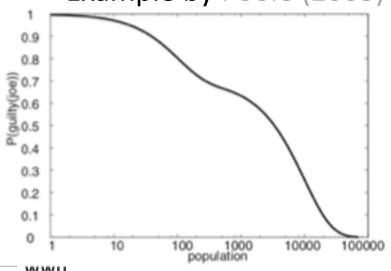
Changing Domains

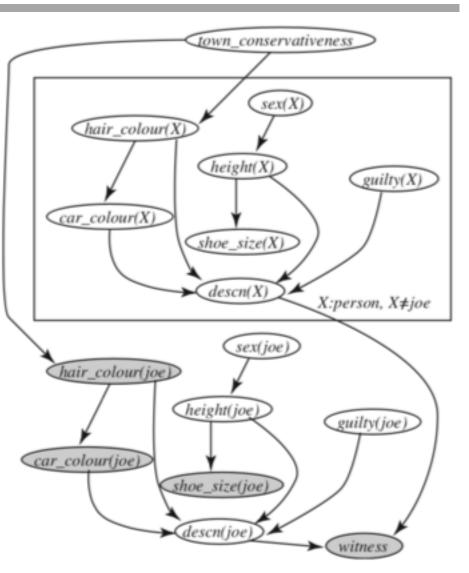
Keep semantics as before

Assume that parfactors accurately describe world

 Posterior probabilities change depending on domain sizes

Example by Poole (2003)

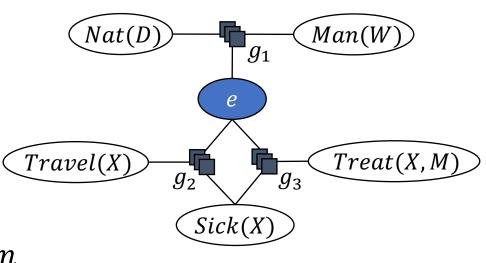




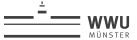
... Without Effects

(Conditional) Independence
 PRVs, containing logical variables X, that are (conditionally) independent from query terms → domains of X have no influence on query results

- E.g., given Epid = e,
 - $\mathcal{D}(D)$ and $\mathcal{D}(W)$ do not matter for queries regarding Travel, Sick, and Treat
 - $\mathcal{D}(X)$ and $\mathcal{D}(M)$ do not matter for queries regarding Nat and Man

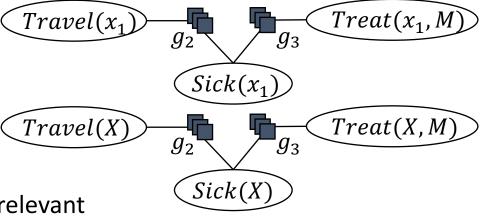


→ Partly invariant under increasing domain sizes



... Without Effects

- A simple case of so-called projectivity
 After shattering, query terms are independent of model parts containing logical variables X → domains of X have no influence on query results
 - Depends on model structure
 - More by Jaeger and Schulte (2018)
- E.g., $P(Sick(x_1))$
 - $\mathcal{D}(X) = \{x_1, ..., x_n\}$
 - After shattering:
 - $\mathcal{D}(X) = \{x_2, \dots, x_n\}$
 - Upper part independent from lower part; $\mathcal{D}(X)$ irrelevant



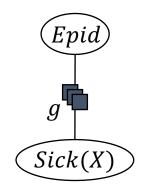
→ Partly invariant under increasing domain sizes



Growing Domain Sizes

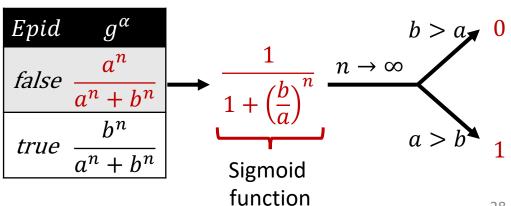
Poole et al. (2014)

- Let domain size n grow
 - With grounding semantics, posteriors change
 - Can lead to extreme behaviour in the posteriors
- Example: Epid gets more and more neighbours with n rising



$$P(Epid) \propto \left(\sum_{s \in r(Sick(X))} g(Epid, Sick(x) = s)\right)^{n}$$
$$= \left(g'(Epid)\right)^{n} = g''(Epid) = g^{\alpha}(Epid)$$

Epid	g'
false	a
true	b



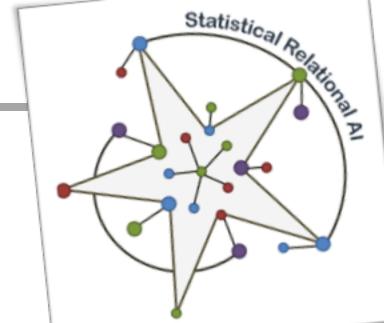


Growing Domain Sizes

Mittal et al. (2019)

- How to avoid extreme behaviour?
- → Adapt values in model dependent on domain size
 - Approach for MLNs: Domain-size aware MLNs
 - Assume predicates P_1, \dots, P_m occur in a first-order formula F
 - Count number of connections c_j for each predicate P_j given new domains
 - Build a connection vector $[c_1, ..., c_m]$
 - Choose $\max_{c_1}[c_1, ..., c_m]$ as scaling-down factor
 - Instead of max, other functions possible
 - Works best if the values in $[c_1, \dots, c_m]$ do not vary that much
 - Given an MLN with a set of formulas F_i with weights w_i
 - Rescale each w_i with scaling-down factor s_i computed for F_i as $\frac{w_i}{s_i}$
 - Analogous approach possible for parfactors





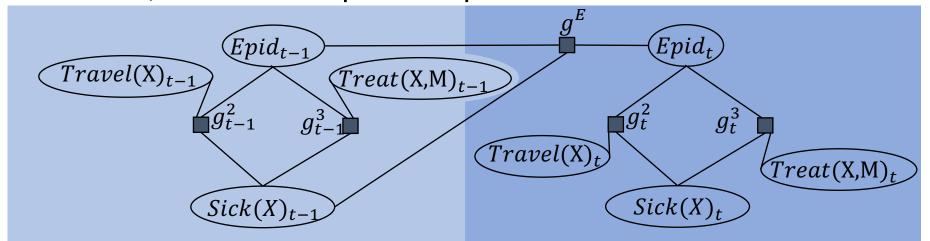
Keeping Inference Going over Time

Algorithms for More Robust Inference



Dynamic PRMs

- Marginal distribution query: $P(A_{\pi}^{i} \mid E_{0:t})$ w.r.t. the model:
 - Hindsight: $\pi < t$ (Was there an epidemic $t \pi$ days ago?)
 - Filtering: $\pi = t$ (Is there currently an epidemic?)
 - Prediction: $\pi > t$ (Will there be an epidemic in πt days?)
- MPE, MAP on temporal sequence



- Define the interface for relational case (avoid groundings)
- Taming reasoning w.r.t. evidence over time (avoid creeping groundings)



Reasoning over Time: Interfaces

- Main idea: Use temporal conditional independences to perform inference on smaller model
 - Normally only a subset of random variables influence next time step → interface variables
 - State description of interface variables from time slice t-1 suffice to perform inference on time slice t
- → Makes present independent from past / future
- Procedure
 - Build a helper structure of clusters (junction tree)
 - Proceed forward one time step at a time (forward message), using the same structure (vanilla junction tree)
 - Algorithms:
 - Propositional: Interface Algorithm (Murphy, 2002)
 - Lifted: Lifted Dynamic Junction Tree Algorithm (Gehrke et al, 2018)



Taming Reasoning

- Evidence can ground a model over time
- Non-symmetric evidence
 - Observe evidence for some instances in one time step
 - Observe evidence for a subset of these instances in another time step
 - Split the logical variable slowly over time
- Vanilla junction trees for each time step
- Forward message carries over splits, leading to slowly grounding a model over time



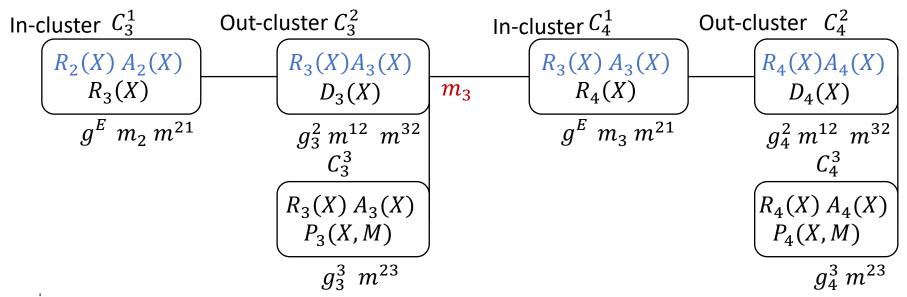
Undoing Splits

- Need to undo splits to
 - keep reasoning polynomial w.r.t. domain sizes
- Where can splits be undone efficiently?
- How to undo splits?
- Is it reasonable to undo splits?
 - Effect of slight differences in evidence?
 - Impact of evidence vs. temporal behaviour of model?



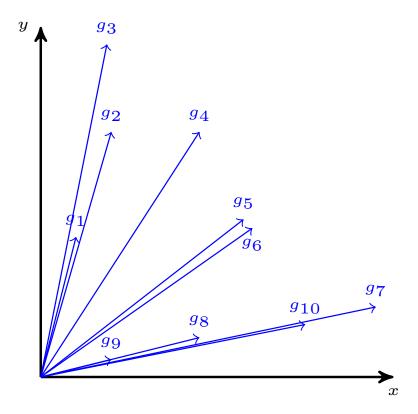
Where Can Splits Be Undone Efficiently?

- Evidence causes splits in a logical variable in the same way in all factors in a model
- LDJT always instantiates a vanilla junction tree
- Forward message carries over splits





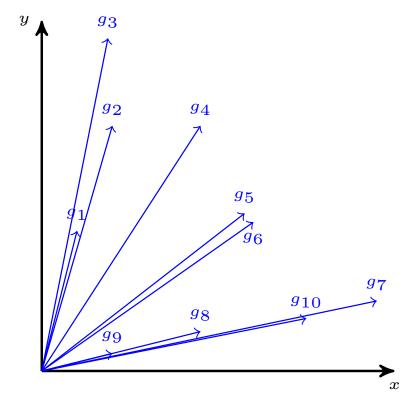
Identifying Similar Groups



- Groups are equal if they have the same full joint distribution
- Full joint distribution computationally hard to get
- → Use parfactors as vector
- → If vectors of two groups point in same direction, they have the same full joint distribution



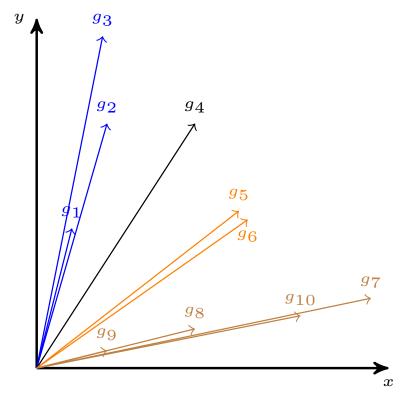
Cluster Groups



- Density-based clustering as unknown number of clusters
- Cosine similarity as distance function



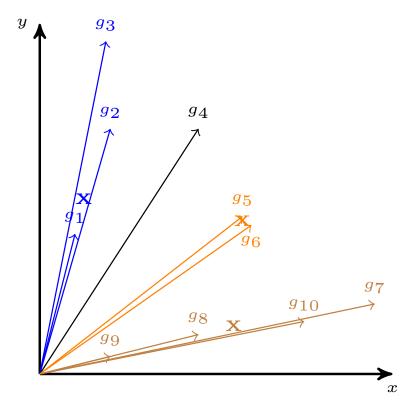
Cluster Groups



- Density-based clustering as unknown number of clusters
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Merge Clusters

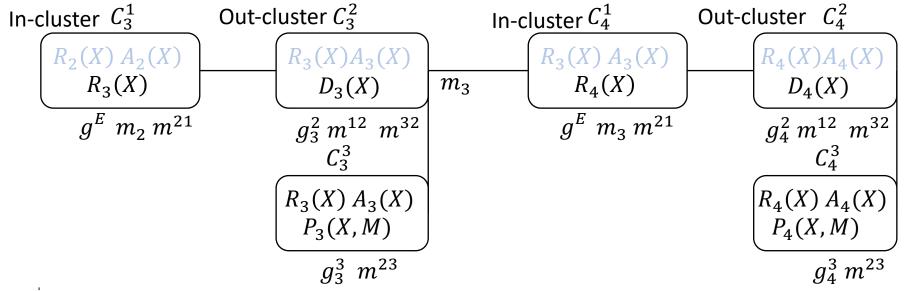


- Merge groups of cluster by calculating mean of cluster while accounting for groundings
- Replace old groups with merged group in temporal message



Is It Reasonable to Undo Splits?

- Approximate forward message
- For each time step the temporal behaviour is multiplied on the forward message
- Indefinitely bounded error due to temporal behaviour





Taming Reasoning

Gehrke et al. (2020)

Need to undo splits to

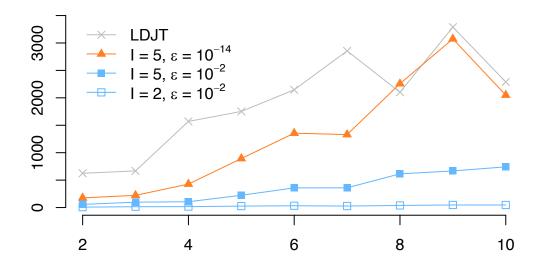
keep reasoning polynomial w.r.t. domain sizes

- Where can splits be undone efficiently?
 - Undo splits in a forward message
- How to undo splits?
 - Find approximate symmetries
 - Merge based on groundings
- Is it reasonable to undo splits
 - Yes, due to the temporal model behaviour (indefinitely bounded error)

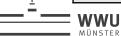


Results

- DBSCAN for Clustering
- ANOVA for checking fitness of clusters



π	Max	Min	Average
0	0.0001537746121	0.000000001720	0.0000191206488
2	0.0000000851654	0.0000000000001	0.0000000111949
4	0.0000000000478	0	0.0000000000068



Agenda

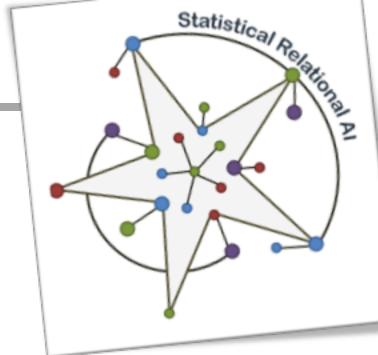
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 - Application example
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The Finish Line

- PRMs as a compact encoding of a full joint
 - Exploit symmetries
- Lifted inference
 - Use information about regular structures to speed up inference
- More robust infernce
 - Multi-query answering using junction tree as helper structure
 - Adaptive inference to get to the point of answering queries again fast
 - Changing domains with minimal effect
 - Keeping inference going over time





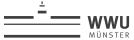
Bibliography & Further Papers

Ordered topic-wise and then alphabetically



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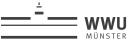
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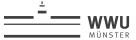
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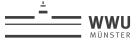
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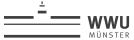
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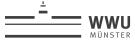
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