

Generating Random Logic Programs Using Constraint Programming

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AI/ML Seminar



THE UNIVERSITY OF EDINBURGH

informatics



EDINBURGH CENTRE FOR
ROBOTICS



Engineering and
Physical Sciences
Research Council

Probabilistic Logic Programs (PROBLOG)

“Smokers” (Domingos et al. 2008; Fierens et al. 2015)

```
0.2::stress(P):-person(P).
0.3::influences(P1,P2):-friend(P1,P2).
0.1::cancer_spont(P):-person(P).
0.3::cancer_smoke(P):-person(P).
    smokes(X):-stress(X).
    smokes(X):-smokes(Y),influences(Y,X).
    cancer(P):-cancer_spont(P).
    cancer(P):-smokes(P),cancer_smoke(P).
    person(michelle).
    person(timothy).
    friend(timothy,michelle).
```

Inference

Let $a \oplus b = a + b - ab$.

$$\begin{aligned}\Pr[\text{cancer}(\textit{michelle})] &= \Pr[\text{cancer_spont}(\textit{michelle})] \\ &\quad \oplus \Pr[\text{smokes}(\textit{michelle})] \\ &\quad \times \Pr[\text{cancer_smoke}(\textit{michelle})]\end{aligned}$$

`cancer(P) :- cancer_spont(P).`

`cancer(P) :- smokes(P), cancer_smoke(P).`

Inference

Let $a \oplus b = a + b - ab$.

$$\Pr[\text{cancer}(\text{michelle})] = 0.1 \oplus 0.3 \times \Pr[\text{smokes}(\text{michelle})]$$

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0.1::cancer_spont(P):-person(P).  
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Inference

Let $a \oplus b = a + b - ab$.

$\Pr[\text{cancer}(\text{michelle})] = 0.1 \oplus 0.3 \times \Pr[\text{smokes}(\text{michelle})]$

$\Pr[\text{smokes}(\text{michelle})] = \Pr[\text{stress}(\text{michelle})]$
 $\oplus \Pr[\text{smokes}(\text{timothy})]$
 $\times \Pr[\text{influences}(\text{timothy}, \text{michelle})]$

```
smokes(X) :- stress(X).
```

```
smokes(X) :- smokes(Y), influences(Y, X).
```

Inference

Let $a \oplus b = a + b - ab$.

$\Pr[\text{cancer}(\text{michelle})] = 0.1 \oplus 0.3 \times \Pr[\text{smokes}(\text{michelle})]$

$\Pr[\text{smokes}(\text{michelle})] = 0.2 \oplus 0.3 \times \Pr[\text{smokes}(\text{timothy})]$

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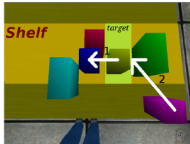
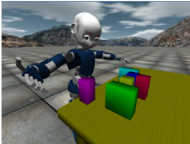
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$\Pr[\text{smokes}(\text{michelle})] = 0.2 \oplus 0.3 \times \Pr[\text{smokes}(\text{timothy})]$

$\Pr[\text{smokes}(\text{timothy})] = \Pr[\text{stress}(\text{timothy})] = 0.2$

```
0.2::stress(P):-person(P).  
      smokes(X):-stress(X).
```

Applications



Moldovan et al. 2012

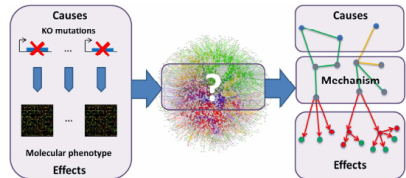
Q1: In a group of 10 people, 60 percent have brown eyes. Two people are to be selected at random from the group. What is the probability that neither person selected will have brown eyes?

Q2: Mike has a bag of marbles with 4 white, 8 blue, and 6 red marbles. He pulls out one marble from the bag and it is red. What is the probability that the second marble he pulls out of the bag is white?

Dries et al. 2017

```
is_malignant(Case):-
    biopsyProcedure(Case,usCore),
    changes_Sizeinc(Case,missing),
    feature_shape(Case).
is_malignant(Case):-
    assoFinding(Case,asymmetry),
    breastDensity(Case,scatteredFDensities),
    vacuumAssisted(Case,yes).
is_malignant(Case):-
    needleGauge(Case,9),
    offset(Case,14),
    vacuumAssisted(Case,yes).
```

Côte-Real, Dutra, and Rocha 2017



De Maeyer et al. 2013

Inference Algorithms and Knowledge Compilation Maps

- NNF negation normal form
- d-DNNF deterministic decomposable negation normal form
- BDD binary decision diagrams
- SDD sentential decision diagrams
- k*-Best only use the *k* most probable proofs

Example Diagrams for $C \wedge (A \vee \neg B)$

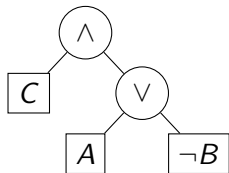


Figure: NNF

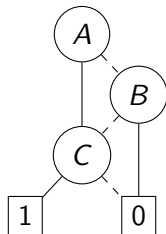


Figure: BDD

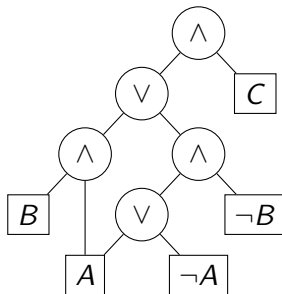


Figure: d-DNNF

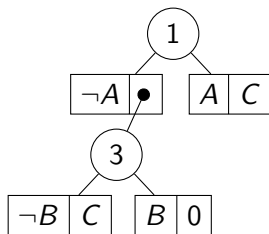


Figure: SDD

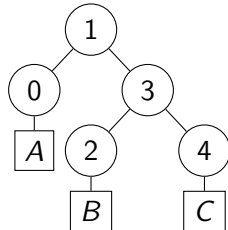


Figure: vtree

How Many Programs Are Used to Test Algorithms?

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Anytime Inference in Probabilistic Logic Programs with T_p -Compilation

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Inference and learning in probabilistic logic programs using weighted Boolean formulas

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k-Optimal: a novel approximate inference algorithm for ProbLog

Joris Renkens · Guy Van den Broeck · Siegfried Nijssen

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ProbLog Technology for Inference in a Probabilistic First Order Logic

Maurice Bruynooghe and Theodoros Mantas and Angelika Kimmig and Bernd Gutmann
and Joost Vennekens and Gerda Janssens and Luc De Raedt¹

What Characterizes a (Probabilistic) Logic Program?

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0.1::cancer_spont(P):-person(P).
0.3::cancer_smoke(P):-person(P).
    smokes(X):-stress(X).
    smokes(X):-smokes(Y),influences(Y,X).
    cancer(P):-cancer_spont(P).
    cancer(P):-smokes(P),cancer_smoke(P).
    person(michelle).
    person(timothy).
    friend(timothy,michelle).
```

What Characterizes a (Probabilistic) Logic Program?

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0.3::`cancer_smoke`(P):-`person`(P).

`smokes`(X):-`stress`(X).

`smokes`(X):-`smokes`(Y),`influences`(Y, X).

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`person`(*michelle*).

`person`(*timothy*).

`friend`(*timothy*, *michelle*).

- predicates, arities

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- variables

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- predicates, arities
- variables
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- length

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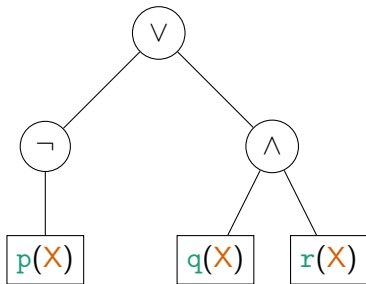
- predicates, arities
- variables
- constants
- probabilities
- length
- complexity

Clauses As Trees

$$\neg p(x) \vee (q(x) \wedge r(x))$$

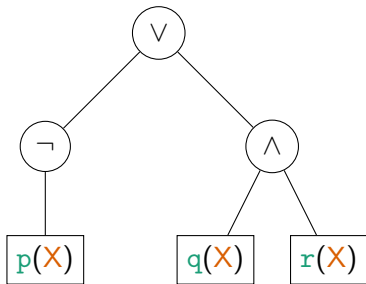
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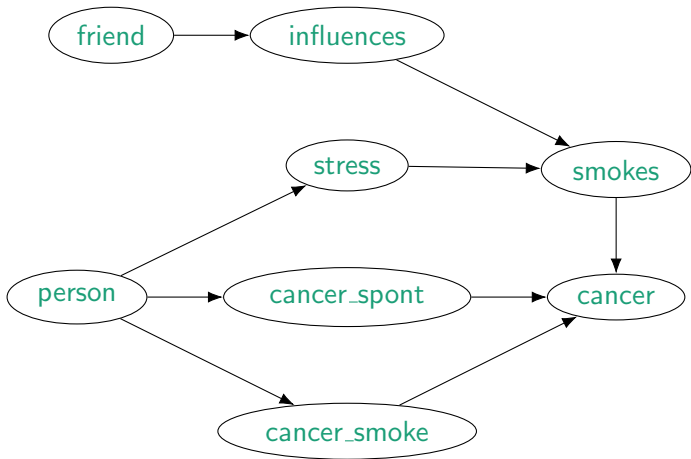
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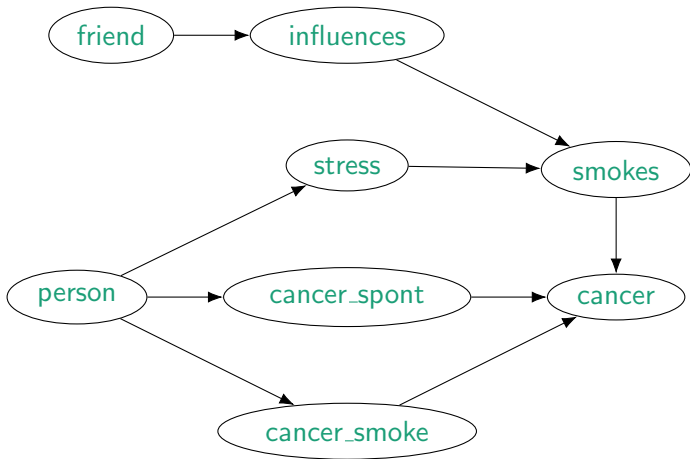


0	0	0	1	2	2	6
∨	¬	∧	p(X)	q(X)	r(X)	⊤

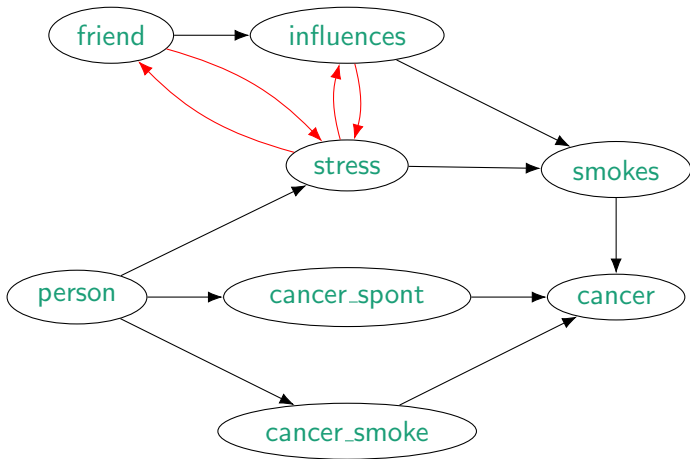
Predicate Dependency Graph



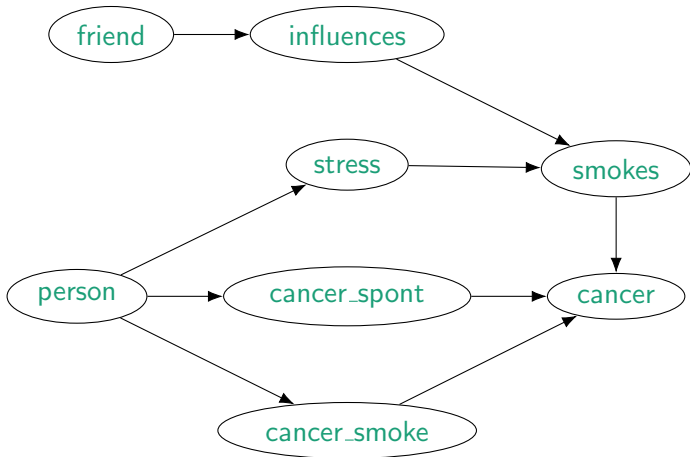
Independence: influences \perp stress



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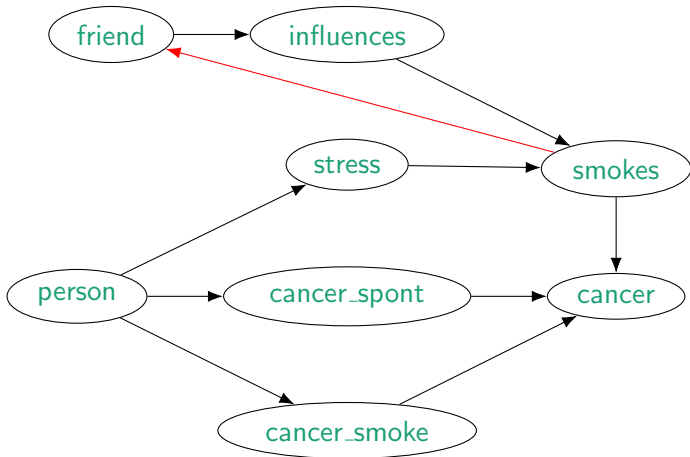


Stratification and Negative Cycles



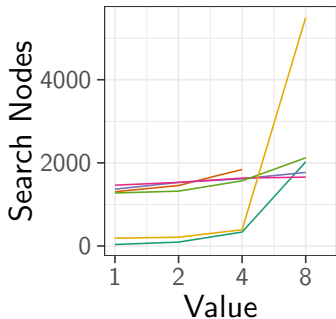
```
0.1::friend(X,Y):- \+ smokes(Y).
```

Stratification and Negative Cycles



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```

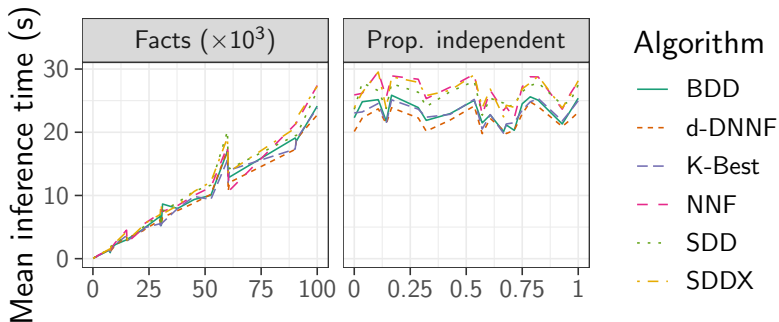

Scalability



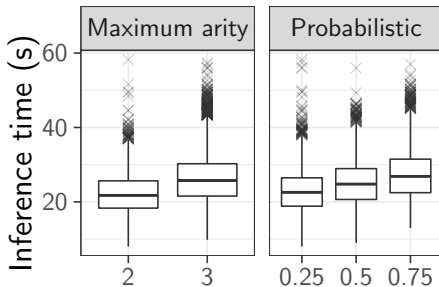
Variable

- The number of predicates
- Maximum arity
- The number of variables
- The number of constants
- The number of additional clauses
- The maximum number of nodes

Properties of Programs vs. Inference Algorithms



Properties of Programs vs. Inference Algorithms



Summary

- foo
- bar
- baz

The implementation of the model is available at

<https://github.com/dilkas/random-logic-programs>

Clause constraints

- `treeStructure` represents `numTrees` trees.
- `treeStructure[0] = 0`
- `numTrees + numNodes = maxNumNodes + 1`
- `treeStructure` is sorted
- For $i = 0, \dots, \text{maxNumNodes} - 1$,
 - If $\text{numNodes} \leq i$,
 - then `treeStructure[i] = i` and `treeValues[i] = \top` ,
 - else `treeStructure[i] < numNodes`.
 - has 0 children $\iff \text{treeValues}[i]$ is a predicate
 - has 1 child $\iff \text{treeValues}[i] = \neg$
 - has > 1 child $\iff \text{treeValues}[i] \in \{\wedge, \vee\}$
 - `treeStructure[i] \neq i \implies treeValues[i] \neq \top`
- If the clause should be disabled, `numNodes = 1` and `treeValues[0] = \top` .

Adjacency matrix representation

$A[i][j] = 0 \iff \nexists k : \text{clauseAssignments}[k] = j \text{ and } i \in \text{clauses}[k].\text{treeValues}$

New constraints

- No (negative) cycles
 - No clever propagation, just entailment checking.
- Independence. Propagation:
 - Two types of dependencies: determined and one-undetermined-edge-away-from-being-determined.
 - Look up the dependencies of both predicates. For each pair of matching dependencies:
 - If both are determined, fail.
 - If one is determined, the selected edge of the other must not exist.