

# Probabilistic Answer Set Programming

## A Research Draft

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May 23, 2022

In short. . .

. . . a word wall. I'm sorry.

- **Machine Learning** has important limitations:
  - The *one table, conditionally independent rows* assumption.
  - *Background knowledge* is hard to include.
  - *Training* requires “large” amounts of data.
  - *Models* are hard to interpret.
- **Inductive Logic Programming** is based on first order logic — solves all the problems above but is sensitive to *noise*.
- **Distribution Semantics** defines the probability of a proposition from probabilities of the (marginally independent) facts.
- **Answer Set Programs** resets the common syntax and semantics of logic programs; A “program” defines *stable models*, not a computation neither a variable substitution.

# ~~Goals~~ Wish list

## Extend distribution semantics to answer sets

- Within a theoretical framework.
- Computationally applicable to “real world” scenarios.
- Easy to include background knowledge.
- Perform common tasks such as *margin*, *mle*, *map*, *etc.*
- Learn program “parameters” and “structure” from *noisy samples* — possibly using *templates*.
- Related to Bayesian Networks, HMMs, *etc.*

**1 Development**

**2 Conclusions**

# The seed on an idea

We want to define the **joint distribution** of the stable models.

- 1 A **boolean random variable** can be described by a disjunction  $a; \neg a$ .
- 2 This ASP program has two stable models:  $a$  and  $\neg a$ .
- 3 A program with  $n$  such facts  $a_i; \neg a_i$  has  $2^n$  stable models, the distinct combinations of those choices.
- 4 **If each  $a_i$  has probability  $p_i$  then the probability of a stable model  $W$  would be**

$$P(W) = \prod_{a_i \in W} p_i \prod_{\neg a_i \in W} (1 - p_i).$$

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**But this is wrong.**

Even assuming that those facts are marginally independent — which we will do.

# Problem 1: Disjunctive Clauses

The ASP program with probabilistic facts

$$b \vee \neg b$$

$$h_1 \vee h_2 \leftarrow b$$

has **three** stable models:  $\{\neg b\}$ ,  $\{b, h_1\}$  and  $\{b, h_2\}$ .

**How to assign a probability to each model?**

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**How to assign a probability to each model?**

Possible approaches:

- 1 Pre-assign a **conditional distribution of the head**:

$$P(h_1, h_2 | b).$$

- 2 Bayesian learn from **observations**:

$$P(h_1, h_2 | b, z) \propto P(b, z | h_1, h_2) P(h_1, h_2).$$

- 3 Start with the former as **prior** and **update** with the latter.

# Questions to address

- How to **match** an observation  $z$  with a clause case  $h, b$ ?
- How do observations **update** the probabilities?
- Why match observations with clauses and **not with stable models**?
- Is this just **bayesian networking**?
- How to frame this in a **sound theoretic setting**?
- Is this enough to compute the **joint distribution of the atoms**?

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## Counters

Instead of setting and updating probabilities, we associate **counters** to disjunctive clauses and their cases.

# Bayesian updates: Matching observations

- An **observation** is a subset of the literals from a program<sup>1</sup>.
- A **consistent** observation has no subset  $\{p, \neg p\}$ .
- A **consistent** observation  $z$  is **relevant** for the clause  $h \leftarrow b$  if  $b \subseteq z$ .
- A disjunctive clause

$$h_1 \vee \dots \vee h_n \leftarrow b_1 \wedge \dots \wedge b_m$$

has  $n$  **cases**:  $\{h_i, b_1, \dots, b_m\}$ ,  $i = 1 : n$ .

- The **consistent** observation  $z$  **matches** the case  $\{h, b_*\}$  if  $\{h, b_*\} \subseteq z$ .

The above definitions also apply to **facts** *i.e.* clauses with an empty body and **constraints** *i.e.* clauses with no head.

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<sup>1</sup>The set of atoms,  $a$ , of the program and their classic negations,  $\neg a$ .

# Bayesian updates: Clauses Update

A consistent observation **relevant** for a clause

$h_1 \vee \dots \vee h_n \leftarrow b$  should:

- Increase the *probability of any matched case*.
- Decrease the *probability of any unmatched case*.

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## Update algorithm

- ① Associate three **counters**,  $r, u, n$ , to each clause  $h \leftarrow b$ .
- ② Associate a **counter**,  $m_i$ , to each case  $h_i, b$  of each clause.
- ③ **Initial** values result from *prior* knowledge.
- ④ Each *consistent* observation **increments**:
  - The  $r$  counters of **r**elevant clauses.
  - The  $u$  counters of **u**nmatched relevant clauses.
  - The  $n$  counters of **n**ot relevant clauses.
  - The  $m_i$  counters of **m**atched cases  $h_i, b$ .
  - Clause counters must verify  $r \leq u + \sum_i m_i$ .

# Updates and counters: An example

Given the following ASP program with **annotated counters**,

$b \vee \neg b$           counters: 7, 2; 12, 3, 0

$h_1 \vee h_2 \leftarrow b$     counters: 4, 3; 6, 2, 5

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## Counters of $b \vee \neg b$

0 observations where not relevant  
(because the body is  $\top$ );

There where 12 relevant  
observations;

Of those,  $b$  was matched by 7,  
 $\neg b$  by 2 and 3 observations  
matched neither ( $\models \sim b, \sim \neg b$ ).

## Counters of $h_1 \vee h_2 \leftarrow b$

There where  $11 = 6 + 5$   
observations, 6 relevant to this  
clause;

From these, 4 matched  $h_1$ , 3  
matched  $h_2$  and 2 matched no  
case.

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## What can be computed?

- $P(\neg b) = \frac{2}{12}$  because  $\neg b$  matched 2 of 12 relevant observations.
- $P(h_1|b) = \frac{4}{6}$  because  $h_1$  matched 4 of 6 relevant observations.
- $P(b)$  **can't be computed** without further information. *E.g.* supposing that **observations are independent** then

$$P(b) = \frac{7 + 6}{12 + 0 + 6 + 5}.$$

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## Note...

Counters are local to clauses and, for distinct clauses, may result from distinct sources. *E.g. the relevant counter of  $h_1 \vee h_2 \leftarrow b$  and the match counter of  $b$  in  $b \vee \neg b$ .*

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**Note...**

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## Note...

Some observations may have neither  $b$  nor  $\neg b$ :

$$P(b) + P(\neg b) < 1.$$

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Given the following ASP program with **annotated counters**,

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$h_1 \vee h_2 \leftarrow b$     counters: 4, 3; 6, 2, 5

## Note...

Since  $h_1$  and  $h_2$  are not independent,

$$\sum_m P(m) \approx 1.02 > 1.$$

# Updates and counters: An example

Given the following ASP program with **annotated counters**,

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## Note...

What is missing to compute the **joint distribution** of the program's atoms

$$P(H_1, H_2, B)?$$

## Shortcomming 2: Default Negation

- How to deal with rules with  $\sim a$  parts?
- Should missing elements on observations be replaced with  $\sim a$  atoms?

**1 Development**

**2 Conclusions**

# Background Material

# Machine Learning

Models are numeric functions:  $y \approx f_{\theta}(x)$ ,  $\theta_i, x_j, y \in \mathbf{R}$ .

- Amazing achievements.
- Noise tolerant.
- (as of today) Huge enterprise funding .

but

- (essentially) Academically solved.
- Models trained from “large” amounts of samples.
- Hard to add background knowledge.
- Models are hard to interpret.
- Single table, independent rows assumption.

# Inductive Logic Programming

Models are logic program:  $p_{\theta}(x, y)$ ,  $\theta_i, x_j, y \in \mathcal{A}$ .

- Amazing achievements, at scale.
- Models trained from “small” amounts of samples.
- Compact, readable models.
- Background knowledge is easy to incorporate and edit.

but

- as of today, Little enterprise commitment.
- as of today, Mostly academic interest.
- Noise sensitive.

# Distribution Semantics

Assigns probability to (marginally independent) facts and derives probability of ground propositions.

Let  $F$  be set of facts,  $S \subseteq F$ ,  $R$  a set of definite clauses and  $p$  a proposition:

$$P_F(S) = \prod_{f \in S} P(f) \prod_{f \notin S} (1 - P(f))$$

$$P(W) = \sum_{S \subseteq F: W = M(S \cup R)} P_F(S)$$

$$P(p) = \sum_{S: S \cup R \vdash p} P_F(S) = \sum_{W: p \in W} P(W)$$

- Amazing achievements, at scale.
- Lots of tools and research.
- The best of both “worlds”?

# Answer Set Programming

A “program” defines stable models *i.e.* minimal sets of derived ground atoms<sup>2</sup>.

- Pure declarative language, unlike Prolog.
- Uses *generate & test* methods instead of proofs .
- Uses both default  $\sim p$  and classical negation  $\neg p$ <sup>3</sup>
- Clauses can be disjunctive  $a; b \leftarrow c, d$ .

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<sup>2</sup>Alternative ~~fact~~ definition:  $X$  is a stable model of  $P$  if  $X = \text{Cn}(P^X)$ .

<sup>3</sup>Classic negation  $\neg a$  in ASP results from replacing the occurrences of  $\neg a$  by a new atom  $a_{\neg}$  and adding the restriction  $\leftarrow a_{\neg}, a$ .

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