

Learning Hierarchical Probabilistic Logic Programs

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Probabilistic Logic Programming (PLP) and Hierarchical PLP

- Probabilistic logic programming is a powerful tool for reasoning with **uncertain** relational models.
- Learning probabilistic logic programs is expensive due to the high cost of inference.
- We consider a restriction of the language of Logic Programs with Annotated Disjunctions called hierarchical PLP in which clauses and predicates are hierarchically organized.
- Generic clause C :

$$C = p(\vec{X}) : \pi :- \phi(\vec{X}, \vec{Y}), b_1(\vec{X}, \vec{Y}), \dots, b_m(\vec{X}, \vec{Y})$$

where $\phi(\vec{X}, \vec{Y})$ is a conjunction of literals for the input predicates.

Generic Hierarchical PLP: <https://rdcu.be/cmCIA>

$$\begin{aligned} C_1 &= r(\vec{X}) : \pi_1 & :- \phi_1, b_{1,1}, \dots, b_{1,m_1} \\ && \dots \\ C_n &= r(\vec{X}) : \pi_n & :- \phi_n, b_{n,1}, \dots, b_{n,m_n} \\ C_{1,1,1} &= r_{1,1}(\vec{X}) : \pi_{1,1,1} & :- \phi_{1,1,1}, b_{1,1,1,1}, \dots, b_{1,1,1,m_{1,1}} \\ && \dots \\ C_{1,1,n_{11}} &= r_{1,1}(\vec{X}) : \pi_{1,1,n_{11}} & :- \phi_{1,1,n_{11}}, b_{1,1,n_{11},1}, \dots, b_{1,1,n_{11},m_{1,n_{11}}} \\ && \dots \\ C_{n,1,1} &= r_{n,1}(\vec{X}) : \pi_{n,1,1} & :- \phi_{n,1,1}, b_{n,1,1,1}, \dots, b_{n,1,1,m_{n,1}} \\ && \dots \\ C_{n,1,n_{n1}} &= r_{n,1}(\vec{X}) : \pi_{n,1,n_{n1}} & :- \phi_{n,1,n_{n1}}, b_{n,1,n_{n1},1}, \dots, b_{n,1,n_{n1},m_{n,n_{n1}}} \\ && \dots \end{aligned}$$

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Example

- UW-CSE domain, the objective is to predict the “advised_by” relation
 $C_1 = \text{advised_by}(A, B) : 0.3 :- \text{student}(A), \text{professor}(B), \text{project}(C, A), \text{project}(C, B), r_{11}(A, B, C).$
- $C_2 = \text{advised_by}(A, B) : 0.6 :- \text{student}(A), \text{professor}(B), \text{ta}(C, A), \text{taught_by}(C, B).$
- $C_{111} = r_{11}(A, B, C) : 0.2 :- \text{publication}(D, A, C), \text{publication}(D, B, C).$
- $r = \text{advised_by}(\text{harry}, \text{ben})$ where *harry* is a student, *ben* is a professor, they have two joint courses, two joint projects and two joint publications from each jointed project.

Parameter learning for Hierarchical probabilistic Logic programs

- Given a HPLP T with parameters Π , and a training set $E = \{e_1, \dots, e_M, \text{not } e_{M+1}, \dots, \text{not } e_N\}$ **PHIL** finds the values of Π that maximize the log likelihood:

$$\arg \max_{\Pi} \sum_{i=1}^M \log P(e_i) + \sum_{i=M+1}^N \log(1 - P(e_i)) \quad (1)$$

$P(e_i)$ is the probability of example e_i .

- Or equivalently minimizing the sum of *cross entropy errors* for all the examples $\text{err}_i = -y_i \log(p_i) - (1 - y_i) \log(1 - p_i)$
- Deep PHIL (DPHIL) and Expectation Maximization PHIL (EMPHIL) learn the parameters using Gradient Descent and Expectation Maximization.

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Structure LEarning of Hierarchical Probabilistic logic programming

Given a set of interpretations, each containing positive and negative examples for a target predicate and facts for input predicates,
 $I = \{e_1, \dots, e_M, \text{not } e_{M+1}, \dots, \text{not } e_N\}$, **SLEAHP** finds an HPLP with parameters Π that maximizes the (log) likelihood

$$LL = \arg \max_{\Pi} \sum_k^{NI} \left(\sum_{i=1}^M \log P(e_{ik}) + \sum_{i=M+1}^N \log(1 - P(e_{ik})) \right) \quad (2)$$

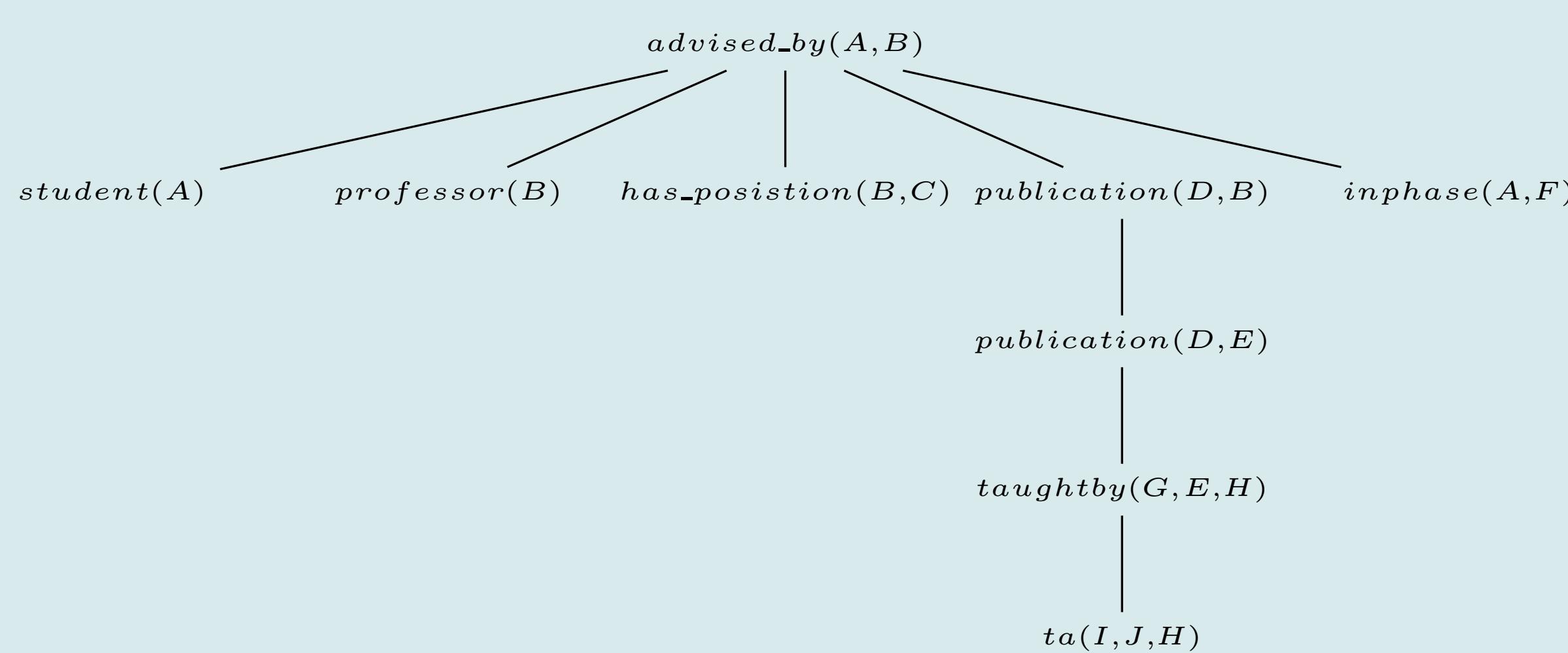
Structure Learning: SLEAHP algorithm

1. **Input**: a set of interpretations and a target predicate (examples).
2. Generate a set of bottom clauses from a language bias.
3. Generate a program tree from the bottom clauses.
4. **Randomly** generate an initially **large** HPLP from the tree.
5. Apply a regularized version of PHIL on the HPLP.
6. Discard Clauses with very small probability values.

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Structure Learning: program tree from a bottom clause

$\text{advised_by}(A, B) :- \text{student}(A), \text{professor}(B), \text{has_position}(B, C), \text{publication}(D, B), \text{publication}(D, E), \text{inphase}(A, F), \text{taughtby}(G, E, H), \text{ta}(I, J, H).$



Structure Learning: initial HPLP generated from the tree

$\text{advised_by}(A, B) : 0.5 :- \text{student}(A).$
 $\text{advised_by}(A, B) : 0.5 :- \text{professor}(B).$
 $\text{advised_by}(A, B) : 0.5 :- \text{has_position}(B, C).$
 $\text{advised_by}(A, B) : 0.5 :- \text{publication}(D, B), \text{hidden_1}(A, B, D).$
 $\text{advised_by}(A, B) : 0.5 :- \text{inphase}(A, E).$
 $\text{hidden_1}(A, B, D) : 0.5 :- \text{publication}(D, F), \text{hidden_11}(A, B, D, F).$
 $\text{hidden_11}(A, B, D, F) : 0.5 :- \text{taughtby}(G, F, H), \text{hidden_111}(A, B, D, F, G, H).$
 $\text{hidden_111}(A, B, D, F, G, H) : 0.5 :- \text{ta}(I, J, H).$

- Learning HPLPs on web browser: **phil on SWISH**
- Link: https://cplint.eu/e/phil/phil_examples.swinb
- Manual: <https://arnaudfadja.github.io/phil/>
- Paper: <https://rdcu.be/cmCIA>

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Experiments: Average time for PHIL

| Time | Mutagenesis | Carcinogenesis | Mondial | UWCSE | PAKDD15 |
|----------------------|---------------|----------------|-----------------|---------------|---------|
| DPHIL | 2.8573 | 178.2680 | 265.4160 | 0.0884 | 34.8170 |
| DPHIL ₁ | 5.2059 | 177.4270 | 311.3280 | 0.2910 | 20.5704 |
| DPHIL ₂ | 5.4450 | 88.5100 | 301.1392 | 0.2214 | 4.9517 |
| EMPHIL | 4.4430 | 106.5540 | 270.6688 | 0.2890 | 6.6106 |
| EMPHIL ₁ | 4.8940 | 181.0890 | 317.4202 | 1.0000 | 7.0691 |
| EMPHIL ₂ | 5.0046 | 146.8440 | 245.3830 | 0.8372 | 6.6334 |
| EMPHIL _{B1} | 2.6478 | 85.3210 | 248.1978 | 0.1572 | 6.0990 |
| EMPHIL _{B2} | 1.5820 | 80.4270 | 261.3612 | 0.1266 | 8.8722 |
| EMPHIL _{B3} | 1.4937 | 94.6850 | 274.9598 | 0.1190 | 6.0985 |
| EMBLEM | 125.62 | - | - | 0.9666 | 154.51 |
| ProbLog2 | 722.00 | 38685.00 | 1607.60 | 161.40 | 596.90 |
| Tuffy | - | - | - | - | - |

Experiments: Average time for SLEAHP

| Time | Mutagenesis | Carcinogenesis | Mondial | UWCSE | PAKDD15 |
|----------------------|----------------|----------------|----------------|-----------------|----------------|
| SLEAHP _{G1} | 41.8250 | 48.7600 | 59.5054 | 219.6410 | 192.4396 |
| SLEAHP _{G2} | 47.1344 | 10524.0900 | 14.0470 | 194.9706 | 162.9938 |
| SLEAHP _{E1} | 48.0152 | 303.0570 | 60.8316 | 387.6650 | 151.7217 |
| SLEAHP _{E2} | 45.9245 | 92.3820 | 61.0996 | 312.2604 | 68.2432 |
| SLEAHP _B | 13.1478 | 1399.0090 | 14.6698 | 295.6734 | 61.5347 |
| SLIPCOVER | 74610.7000 | 17419.4500 | 650.3630 | 141.3600 | 242.7077 |
| PROBFOIL+ | 1726.6000 | 15433.0000 | - | - | - |
| MLN-BC | 91.7700 | 59.73500 | 56.5007 | 376.2356 | * |
| MLN-BT | 360.8563 | 87.5020 | 53.5568 | 891.4226 | 915.5794 |
| RDN-B | 183.7140 | 61.5000 | 192.0487 | 501.1176 | 793.7661 |

- PHIL and SLEAHP Provide similar performance w.r.t SOTA in terms of area under the ROC and PR curves but appear to be scalable.

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