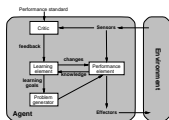


Clinical Decision Support Systems, 23/24

Inês Dutra and Pedro Rodrigues
DCC-FCUP & MEDCIDS-FMUP

ines@dcc.fc.up.pt, pprodrigues@med.up.pt

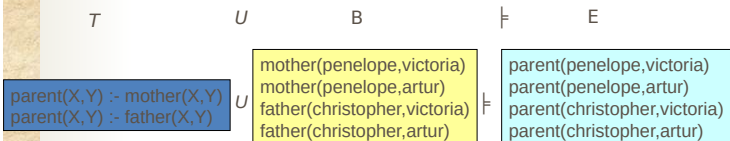
March 1st, 2024



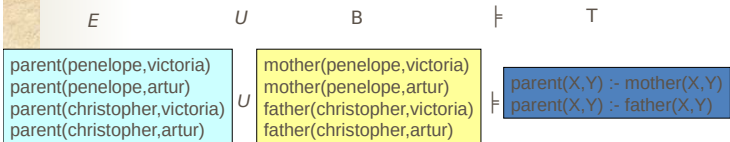
(Multi) Relational Data Mining

- Inductive Logic Programming (ILP)
- Probabilistic Reasoning – PR
- $ILP + PR = SRL$
- Statistical Relational Learning

Deductive Reasoning


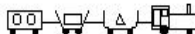
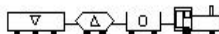




Inductive Reasoning


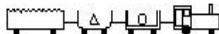





Inductive Logic Programming

TRAINS GOING EAST

1. 
2. 
3. 
4. 
5. 

TRAINS GOING WEST

1. 
2. 
3. 
4. 
5. 

Inductive Logic Programming: example



```
short(car_12).  
closed(car_12).
```

```
long(car_11).
```

```
long(car_13).
```

```
short(car_14).
```

```
open(car_11).
```

```
open_car(car_13).
```

```
open_car(car_14).
```

```
shape(car_11,rectangle).
```

```
shape(car_12,rectangle).
```

```
shape(car_13,rectangle).
```

```
shape(car_14,rectangle).
```

```
load(car_11,rectangle,3).
```

```
load(car_12,triangle,1).
```

```
load(car_13,hexagon,1).
```

```
load(car_14,circle,1).
```

```
wheels(car_11,2).
```

```
wheels(car_12,2).
```

```
wheels(car_13,3).
```

```
wheels(car_14,2).
```

```
has_car(east1,car_11).
```

```
has_car(east1,car_12).
```

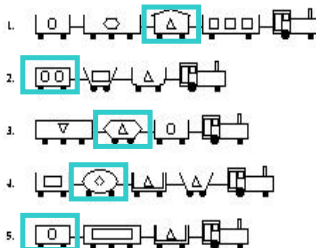
```
has_car(east1,car_13).
```

```
has_car(east1,car_14).
```

Inductive Logic Programming: example

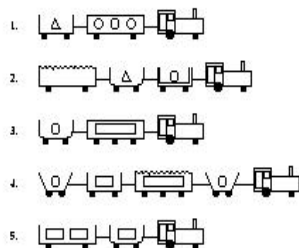
TRAINS GOING EAST

1. TRAINS GOING EAST



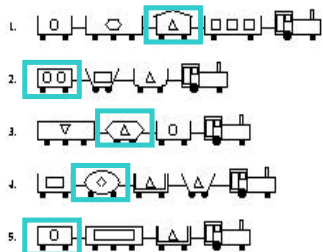
TRAINS GOING WEST

2. TRAINS GOING WEST

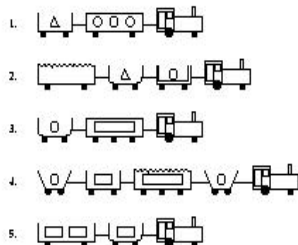


Inductive Logic Programming: example

TRAINS GOING EAST



TRAINS GOING WEST





eastbound(T) IF has_car(T,C) AND short(C) AND closed(C)

Another example: extracting knowledge from mammogram annotations

```
is_malignant(A) if
  'BIRADS_category'(A,b5), 'MassPAO'(A,present),
  'Age'(A,age6570),
  previous_finding(A,B), 'MassesShape'(B,none),
  'Calc_Punctate'(B,notPresent),
  previous_finding(A,C), 'BIRADS_category'(C,b3).
```

The learned rule above says:

A is classified as **BI-RADS 5** AND had a mass present 
in a patient who:
was between the ages of 65 and 70
had two prior mammograms (**B**, **C**)
AND prior mammogram (**B**):
had no mass shape described
had no punctate calcifications
AND prior mammogram (**C**) was classified as **BI-RADS 3** 

BI-RADS: Breast Imaging Reporting And Data System

Inductive Logic Programming

- More formally:
- Given:
 - A set of examples \mathbf{e} (observations, cases, instances) labelled as positive or negative (class \mathbf{c})
 - A language
 - Possibly, a set of constraints
- Find:
 - A hypothesis \mathbf{h} , such that $\mathbf{h}(\mathbf{e}_i) = \mathbf{c}_i$
 - For most examples

Inductive Logic Programming

- Advantages:
 - Utilization of a language that is easy to interpret
 - More concise classifiers
 - More powerful representation: relations
- Disadvantages:
 - Very large search space
 - Non-probabilistic classification

Properties

- Prior satisfiability

$$B \wedge E^- \not\models \square$$

(H is not a consequence of B and E-)

- Posterior sufficiency

$$B \wedge H \models E^+$$

(H allows to explain E+ relative to B)

- Posterior satisfiability

$$B \wedge H \wedge E^- \not\models \square$$

(B and H are consistent with E-)

- Prior necessity

$$B \not\models E^+$$

(some e+ must be false relative to the model found for B)

ILP: A Common Approach

- Use a greedy covering algorithm.
 - Repeat while some positive examples remain uncovered (not entailed):
 - Find a *good clause* (one that covers as many positive examples as possible but no/few negatives).
 - Add that clause to the current theory, and remove the positive examples that it covers.
- ILP algorithms use this approach but vary in their method for finding a *good clause*.

Some ILP Systems

- PROGOL, ALEPH (top-down): saturates first uncovered positive example, and then performs top-down admissible search of the lattice above this saturated example.
- GOLEM (bottom-up), FOIL (top-down), LINUS/DINUS.
- Tilde, Claudien, IndLog, ...

ILP Saturation

- Consists of building a *bottom clause* (seed)
- Incorporates background knowledge to an atomic formula
- Example: (gene that codes for a protein responsible for metabolism)

```
metabolism(A) :-  
  essential(A,'Non-Essential'), motif(A,'PS00510'), chromosome(A,'14'),  
  interaction(A,B,C,E),  
  essential(B,'Non-Essential'), motif(B,'PS00188'), chromosome(B,'2'),  
  interaction(A,F,D,G),  
  intertype(C,'Genetic'), intertype(D,?),  
  interaction(B,A,C,E),  
  interaction(B,H,C,I),  
  interaction(F,A,D,G),  
  interaction(H,B,C,I), interaction(H,_,_).
```

ILP: Aleph

- Procedure to extract theories from examples
- Complete (branch-and-bound) search for best clause in the *whole* space
- Search subject to several user control settings
 - Max clause length
 - Max chaining length
 - Minacc
 - Max nodes
 - Search strategy, etc.

ILP: Aleph

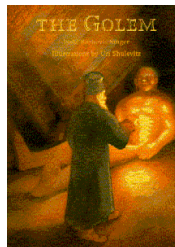
- Aleph
 - Desenvolvido na Universidade de Oxford por Ashwin Srinivasan

<http://www.comlab.ox.ac.uk/oucl/research/areas/machinelearn/Aleph/>

ILP: Aleph

Then the Rabbi said,
“Golem, you have not been completely formed, but I am about to finish you now...You will do as I will tell you.”

Saying these words, Rabbi Leib finished engraving the letter **Aleph**. Immediately the golem began to rise.



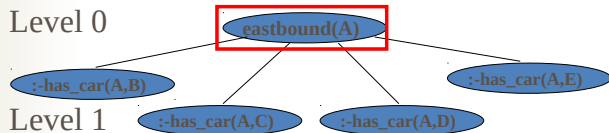
Aleph: algorithm

- **Initial State:**
 - Examples or observations
 - Descriptions: background knowledge (BK)
- **Final State:** hypothesis or theory or model
- **Transitions:** intermediate hypotheses

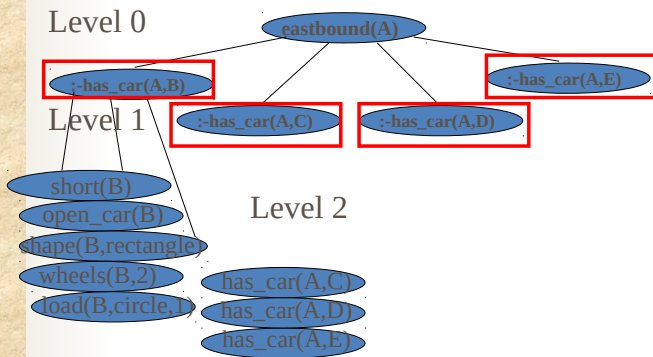
Aleph: algorithm

- Select example
- Build most-specific-clause (**bottom clause**)
- Search. Find a clause more general than the bottom clause
- Remove redundant. The clause with the best score is added to the current theory, and all examples made redundant are removed. This step is sometimes called the "**cover removal**" step. Note here that the best clause may make clauses other than the examples redundant
- Return to first step

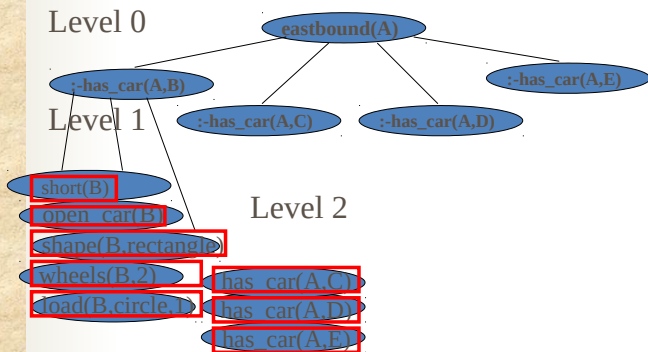
Aleph: Search



Aleph: Search



Aleph: Search



Aleph: Knowledge Representation

Input Files: Prolog Syntax

dtp.b: BK

dtp.f: pos examples

dtp.n: neg examples

Representation: BK

```
chromosome('G234064','1').  
chromosome('G234065','1').  
chromosome('G234070','1').  
chromosome('G234073','1').  
chromosome('G234074','1').  
chromosome('G234076','1').  
chromosome('G234084','2').  
chromosome('G234085','2').  
chromosome('G234089','2').
```


Representation: BK

interaction('G234062','G235011','Physical',?).

interaction('G234064','G234126','Genetic-Physical','0.9141').

interaction('G234064','G235065','Genetic-Physical','0.7515').

interaction('G234064','G235571','Physical','0.9691').

interaction('G234065','G234073','Physical','0.7492').

interaction('G234065','G235042','Physical','-0.4659').

Representation: Examples

metabolism('G239098').
metabolism('G234980').
metabolism('G235245').
metabolism('G234108').
metabolism('G238387').
metabolism('G240504').
metabolism('G236733').

Example of clause learned

metabolism(A) :-

chromosome(A,'15'),

interaction(A,B,_,_),

*complex(B,'Transcription
complexes/Transcriptosome').*

A and **B** are variables that represent **genes**

Aleph: algorithm

- Example: Michalski's trains

Aleph: algorithm

- Saturation (saturated / bottom clause):

eastbound(A) :-

has_car(A,B), has_car(A,C), has_car(A,D), has_car(A,E),
short(B), short(D), closed(D), long(C),
long(E), open_car(B), open_car(C), open_car(E),
shape(B,rectangle), shape(C,rectangle), shape(D,rectangle),
shape(E,rectangle),
wheels(B,2), wheels(C,3), wheels(D,2), wheels(E,2),
load(B,circle,1), load(C,hexagon,1), load(D,triangle,1),
load(E,rectangle,3).

Aleph: algorithm

- Search: most general clause

eastbound(A) :-

```
has_car(A,B), has_car(A,C), has_car(A,D), has_car(A,E),  
short(B), short(D), closed(D), long(C),  
long(E), open_car(B), open_car(C), open_car(E),  
shape(B,rectangle), shape(C,rectangle), shape(D,rectangle),  
shape(E,rectangle),  
wheels(B,2), wheels(C,3), wheels(D,2), wheels(E,2),  
load(B,circle,1), load(C,hexagon,1), load(D,triangle,1),  
load(E,rectangle,3).
```

Aleph: algorithm

- Search: add possible descendants (candidate literals of level 1)

eastbound(A) :-

```
has_car(A,B), has_car(A,C), has_car(A,D), has_car(A,E),  
short(B), short(D), closed(D), long(C),  
long(E), open_car(B), open_car(C), open_car(E),  
shape(B,rectangle), shape(C,rectangle), shape(D,rectangle),  
shape(E,rectangle),  
wheels(B,2), wheels(C,3), wheels(D,2), wheels(E,2),  
load(B,circle,1), load(C,hexagon,1), load(D,triangle,1),  
load(E,rectangle,3).
```

Aleph: algorithm

- Search: add possible descendants of level 2

```
eastbound(A) :-  
  has_car(A,B), has_car(A,C), has_car(A,D), has_car(A,E),  
  short(B), short(D), closed(D), long(C),  
  long(E), open_car(B), open_car(C), open_car(E),  
  shape(B,rectangle), shape(C,rectangle), shape(D,rectangle),  
  shape(E,rectangle),  
  wheels(B,2), wheels(C,3), wheels(D,2), wheels(E,2),  
  load(B,circle,1), load(C,hexagon,1), load(D,triangle,1),  
  load(E,rectangle,3).
```


Aleph: algorithm

- Search: second descendant of level 1

eastbound(A) :-

```
has_car(A,B), has_car(A,C), has_car(A,D), has_car(A,E),
short(B), short(D), closed(D), long(C),
long(E), open_car(B), open_car(C), open_car(E),
shape(B,rectangle), shape(C,rectangle), shape(D,rectangle),
shape(E,rectangle),
wheels(B,2), wheels(C,3), wheels(D,2), wheels(E,2),
load(B,circle,1), load(C,hexagon,1), load(D,triangle,1),
load(E,rectangle,3).
```

Aleph: algorithm

- Search: descendants of second descendant...

eastbound(A) :-

```
has_car(A,B), has_car(A,C), has_car(A,D), has_car(A,E),  
short(B), short(D), closed(D), long(C),  
long(E), open_car(B), open_car(C), open_car(E),  
shape(B,rectangle), shape(C,rectangle), shape(D,rectangle),  
shape(E,rectangle),  
wheels(B,2), wheels(C,3), wheels(D,2), wheels(E,2),  
load(B,circle,1), load(C,hexagon,1), load(D,triangle,1),  
load(E,rectangle,3).
```

Aleph: example of run

 aleph_trains

Aleph: how to run?

- You need to have a Prolog system
 - Yap: <http://yap.sourceforge.net> OU
 - SWI: <http://www.swi-prolog.org>
- Aleph:
<http://www.comlab.ox.ac.uk/oucl/research/areas/machlearn/Aleph/>
- Files: .b, .f, .n
- To make things easier: everything in the same directory!

Aleph: Basic Commands

- `read_all`
- `reduce`
- `induce`

Aleph: Parameters

```
:- set(clauselength,5).
:- set(depth, 200).
:- set(i,3).
:- set(noise,0).
:- set(minacc,0.7).
:- set(nodes,1000000).
:- set(m,20).
:- set(evalfn,mestimate).
:- set(test_pos,'/u/dutra/Protein/prot_test_set.f').
:- set(test_neg,'/u/dutra/Protein/prot_test_set.n').
:- set(optimise_clauses,true).

:- set(record,true).
:- set(recordfile,'prot_train_set.out').
:- set(samplesize,0).
```

$Strength\ estimate = (support + m * prior) / (coverage + m)$

$M \rightarrow 0, strength \rightarrow precision$

Support = True positives

Coverage = True positives + false negatives

Aleph: Modes and Types

```
:- modeh(1,eastbound(+train)).
:- modeb(1,short(+car)).
:- modeb(1,closed(+car)).
:- modeb(1,long(+car)).
:- modeb(1,open_car(+car)).
:- modeb(1,double(+car)).
:- modeb(1,jagged(+car)).
:- modeb(1,shape(+car,#shape)).
:- modeb(1,load(+car,#shape,#int)).
:- modeb(1,wheels(+car,#int)).
:- modeb(*,has_car(+train,-car)).

:- determination(eastbound/1,short/1).
:- determination(eastbound/1,closed/1).
:- determination(eastbound/1,long/1).
:- determination(eastbound/1,open_car/1).
:- determination(eastbound/1,double/1).
:- determination(eastbound/1,jagged/1).
:- determination(eastbound/1,shape/2).
:- determination(eastbound/1,wheels/2).
:- determination(eastbound/1,has_car/2).
:- determination(eastbound/1,load/3).
```

Aleph: Modes and Types

```
:- modeh(1,metabolism(+gene)).  
:- modeb(1,essential(+gene,#essential)).  
:- modeb(1,class(+gene,#class)).  
:- modeb(1,complex(+gene,#complex)).  
:- modeb(1,phenotype(+gene,#phenotype)).  
:- modeb(1,motif(+gene,#motif)).  
:- modeb(1,chromosome(+gene,#chromosome)).  
:- modeb(*,gte(+number,#number)).  
:- modeb(*,interaction(+gene,-gene,-intertype,-number)).  
:- modeb(1,intertype(+intertype,#intertype)).
```


Example: drug discovery using Aleph refinement operators

- Given:
 - Molecules active and inactive for dtp
 - Their description in terms of coordinates and bonds
- Find small structures that model active molecules

Examples: drug discovery

- Examples of dtp groups:

hydrophobic(m752,
hyphob([a2, a3, a5, a8, a7, a4, a2],
2.16452, -0.833917, 3.6379)).

hacc(m9706,
hacc(a10, -6.2969, -1.3684, -0.4631)).

Example: drug discovery

- Utilisation of **refinement operator**

`refine(false,Clause):-`

```
member(Point1, [hydrophobic(M,P1), hdonor(M,P1),halogen(M,P1),hacc(M,P1)]),  
member(Point2,[hydrophobic(M,P2),hdonor(M,P2),halogen(M,P2),hacc(M,P2)]),  
Clause = (active(M) :- Point1, Point2, dist(M,P1,P2,D1,E)).
```

`refine(Clause1,Clause2):-`

```
Clause1 = (active(M) :- Point1,Point2, dist(M,P1,P2,D1,E)), member(Point3,  
[hydrophobic(M,P3),hdonor(M,P3),halogen(M,P3),hacc(M,P3)]),  
Clause2 = (active(M) :- Point1, Point2, dist(M,P1,P2,D1,E),  
Point3, dist(M,P1,P3,D2,E), dist(M,P2,P3,D3,E)).
```

- Reduce search space!!!