

Introduction to Inductive Logic Programming

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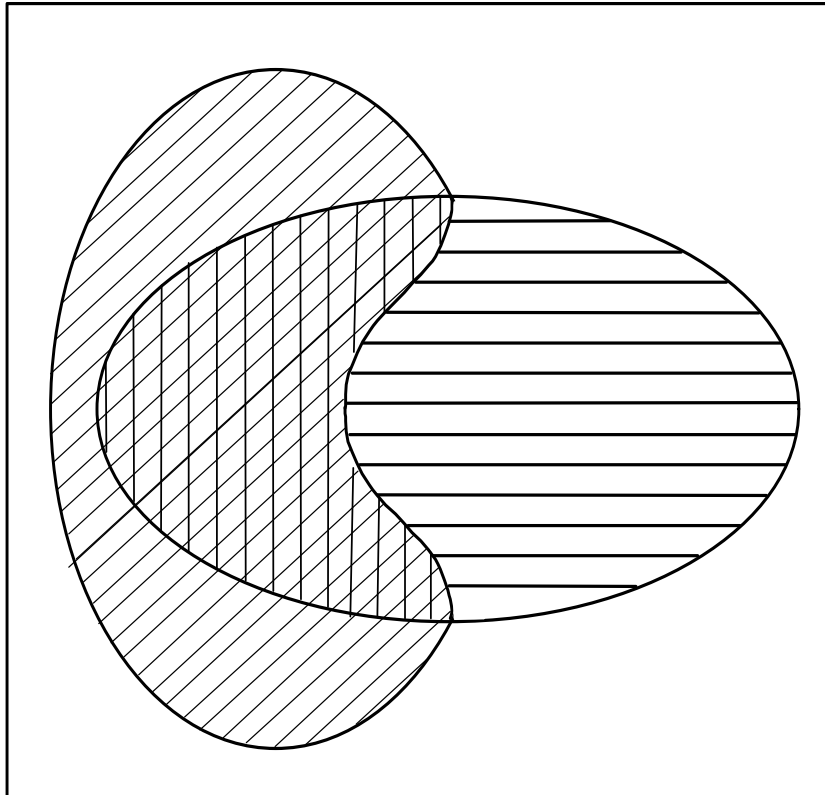
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In collaboration with Olga Štěpánková

Goal of inductive machine learning

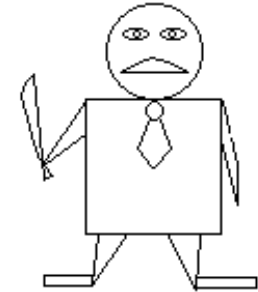
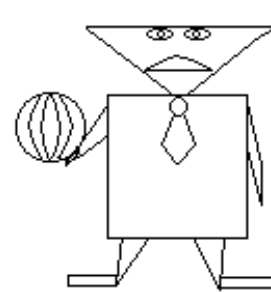
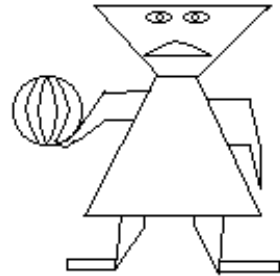
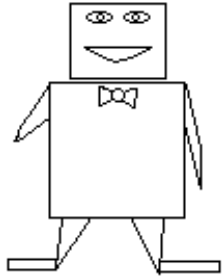


Given a finite subset of positive and negative examples E^+ and E^- , find a general description H of the whole set (a concept) so that

- H fits E^+ best, and
- discriminates also other positive and negative examples

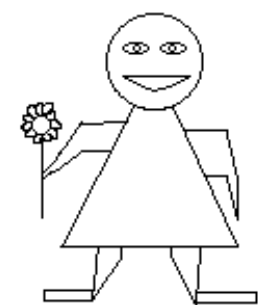
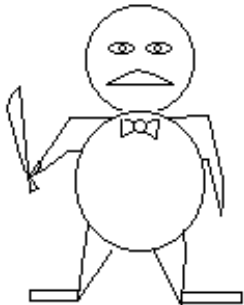
Example:

Can we recognize robots after short experience?



friendly

unfriendly



Example:

Robots and an attribute-value description

<i>head</i>	<i>smile</i>	<i>neck</i>	<i>body</i>	<i>In hand</i>	<i>friendly</i>
Circle	ne	Tie	Rectangle	Sword	no
Rectangle	ano	Butterfly	Rectangle	Nothing	yes
Circle	ne	Butterfly	Circle	Sword	yes
Triangle	ne	Tie	Rectangle	Ball	no
Circle	ano	Nothing	Triangle	Flower	no
Triangle	ne	Nothing	Triangle	Ball	yes
Triangle	ano	Tie	Circle	Nothing	no
Circle	ano	Tie	Circle	Nothing	yes

Example: hypothesis and testing

In the form of a decision tree

if neck = butterfly then yes
= nothing then
 if head = triangle then yes
 else no
= tie then
 if body = rectangle then no
 else
 if head = circle then yes
 else no

head	smile	neck	body	in hand	friendly
circle	no	tie	circle	sword	yes
triangle	yes	nothing	rectangle	nothing	yes

Example: hypothesis and testing (cont.)

Using a relation of equality

if neck = body then yes
else no

head	smile	neck	body	in hand	friendly
circle	no	tie	circle	sword	yes
triangle	yes	nothing	rectangle	nothing	no

Both trees classify the learning examples in the same way but they differ on testing set.

How many hypotheses can we have for a given learning set E ?

- **Fact:** a number of concepts is much more than a number of possible hypotheses.
- **Consequence:** mostly we have to accept a hypothesis that is only “almost correct” (semi-optimal).
- Even in that case, we usually have more than one hypothesis for a given E .

Hypothesis selection and **Ockham razer**



William of Ockham
(c. 1285- c.1349)
Franciscan Monk
Philosopher

Williamu of Ockham :

*„Entia non sunt
multiplicanda praeter
necessitatem“,*

- *„ A number of entities
should not be more
than is necessary.“*
- **Einstein:** *„... but also
not less.“*

Example 2: a need for domain knowledge

Pø.	139	319	854	468	349	561	756	789	987	256	189	354
Kl.	+	-	-	+	+	-	-	+	-	+	+	-

What hypothesis language to use?

E.g. attributes **c1**, **c2** and **c3** describe the 1st, 2nd and 3rd digit

Example 2: a need for domain knowledge

Př.	139	319	854	468	349	561	756	789	987	256	189	354
Kl.	+	-	-	+	+	-	-	+	-	+	+	-

Not enough

because classification depends on an order of digits

a relation of ordering is needed
= (apriori) domain knowledge

Result: **if $c1 < c2$ & $c2 < c3$ then '+'.**

Example 3: parity

Pø.	C1	C2	C3	C4	C5	C6	C7	C8	Kl.
1	1	0	1	1	0	0	0	0	-
2	1	0	1	1	0	0	0	1	+
3	1	1	1	1	0	0	0	1	-
4	0	1	1	1	0	0	0	1	+

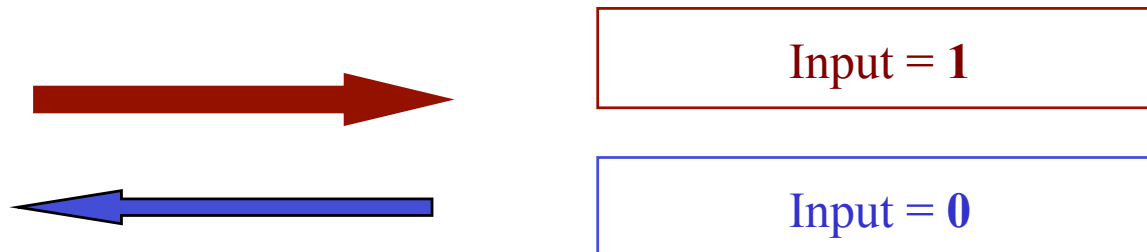
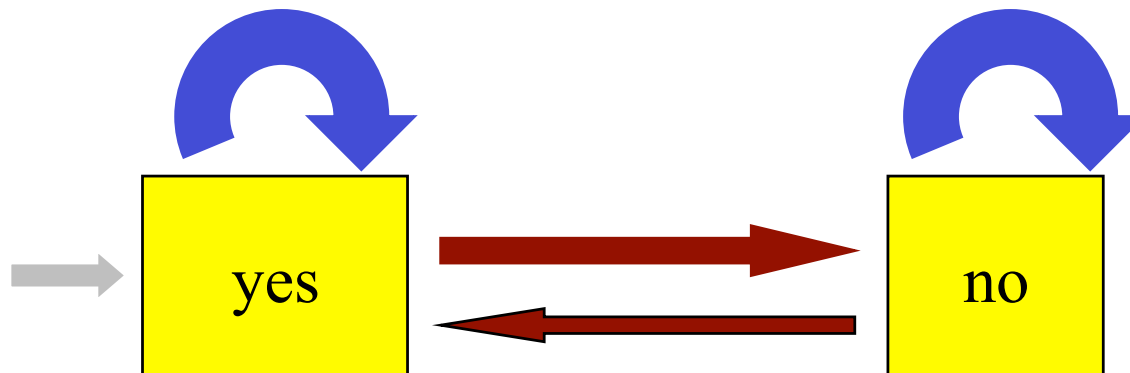
No attribute can be removed – all are important

A decision tree would be too complex –

Why not an automaton?

Recursion is useful

Automaton that decides
„odd number of 1“



Selection of a good hypothesis

depends on a right

- language for representation of examples
- hypothesis language
- domain knowledge

When an attribute-value representation is insufficient?

- Examples do not have a uniform description (e.g. are of a different length)
- A structure of examples is important
- Domain knowledge is (multi-)relational

Inductive logic programming: Basic task

(Muggleton94)

A set of positive E^+ and negative E^- examples

Domain knowledge B (a logic program)

goal: to find a logic program P that together with B covers
(almost all) positive examples and
not cover (almost no) negative example

+: much more flexible

data of any structure can be processed

-: some effort needed

more time consuming(even though \ll NeuroN)

Inductive logic programming: Example

Example: find a path in an oriented graph

path(X,Y) :- edge(X,Y).

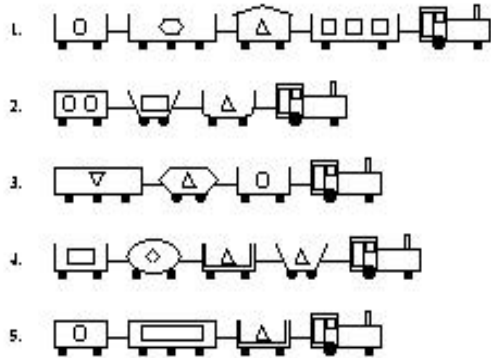
path(X,Y) :- path(X,U),edge(U,Y).

edge(1,2). edge(1,3). edge(2,3). edge(2,4). ...

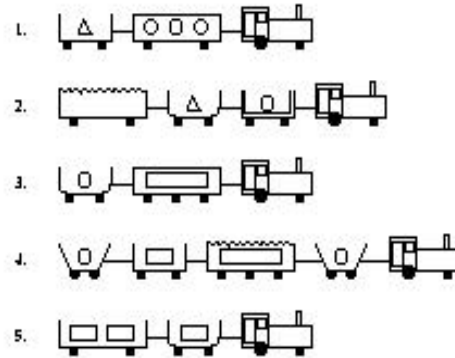
= domain knowledge

East-West Trains (1)

1. TRAINS GOING EAST



2. TRAINS GOING WEST



East-West Trains (2)



```
eastbound(east1).           % eastbound train 1
eastbound(east2).           short(car_12). closed(car_12).
eastbound(east3).           long(car_11). open_car(car_11).
eastbound(east4).           ...
eastbound(east5).           shape(car_11,rectangle). shape(car_12,rectangle).
                             ...
eastbound(west6).           load(car_11,rectangle,3). load(car_12,triangle,1).
eastbound(west7).           ...
eastbound(west8).           wheels(car_11,2). wheels(car_12,2).
eastbound(west9).           ...
eastbound(west10).         has_car(east1,car_11). has_car(east1,car_12).
                             has_car(east1,car_13). has_car(east1,car_14).
```

East-West Trains (3)

`:- modeh(1, eastbound(+train)).`

`:- modeb(*, has_car(+train, -car)).`

`:- modeb(1, short(+car)).`

`:- modeb(1, load(+car, #shape, #int)).`

...

`:- determination(eastbound/1, has_car/2).`

`:- determination(eastbound/1, short/1).`

`:- determination(eastbound/1, load/3).`

...

`:- set(...).`

`?- [aleph].`

`?- read_all(train).`

`?- induce.`

...

[Rule 1] [Pos cover = 5 Neg cover = 0]

`eastbound(A) :-`

`has_car(A,B), short(B), closed(B).`

Actual

	+	-		Accuracy = 1.0
--	---	---	--	----------------

	+ 5	0	5	
--	-----	---	---	--

Pred -	0	5	5	[time taken] [0.07]
--------	---	---	---	---------------------

	5	5	10	[total clauses constructed] [100]
--	---	---	----	-----------------------------------

East-West Trains (4)



[bottom clause][literals] [25][saturation time] [0.01]

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).

[reduce]

eastbound(A).	[5/5]	...
eastbound(A) :- has_car(A,B).	[5/5]	
eastbound(A) :-		eastbound(A) :- has_car(A,B), closed(B), shape(B,rectangle).
has_car(A,B), short(B).	[5/5]	eastbound(A) :- has_car(A,B), closed(B), wheels(B,2).
...		eastbound(A) :-has_car(A,B), closed(B), load(B,triangle,1). [2/0]
eastbound(A) :-		
has_car(A,B),wheels(B,3).	[3/1]	
eastbound(A) :-		...
has_car(A,B), closed(B).	[5/2]	
eastbound(A) :- has_car(A,B),		eastbound(A) :- has_car(A,B), short(B), closed(B). [5/0]
load(B,triangle,1).	[5/2]	
...		

Specialization and generalization

A formula F is a **specialization** of a formula G iff

F is a logical consequence of G

$G \models F$ (any model of G is also a model of F).

Specialization operator (refinement operator)

assigns to a clause a set of all its specializations

Most of ILP systems use two basic operations of specialization

binding two variables

$\text{spec}(\text{path}(X, Y)) = \text{path}(X, X)$

adding a goal into a clause body

$\text{spec}(\text{path}(X, Y)) = (\text{path}(X, Y) :- \text{edge}(U, V))$

and also

substitution a variable with a constant

$\text{spec}(\text{number}(X)) = \text{number}(0)$

substitution a variable with a most general term

$\text{spec}(\text{number}(X)) = \text{number}(s(Y))$.

Generic algorithm

$QH := initialize(B; E^-, E^-) ;$

while not(*stopping_kriterium*(QH)) **do**

remove H from QH ;

choose_rules r_1, \dots, r_k from R ;

 applying r_1, \dots, r_k to H create a set H_1 ;

$QH := (QH-H) \cup H_1 ;$

remove_some_elements from QH ;

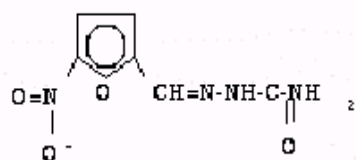
choose_hypothesis P from QH

Successful ILP applications

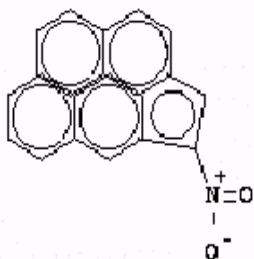
Bioinformatics

- **Structure Activity Relationships (SAR):** given
 - Chemical structure and
 - Empirical data about toxicity/ mutagenicity/ effect in therapy
- **Goal: to find a reason for the effect**

Positive

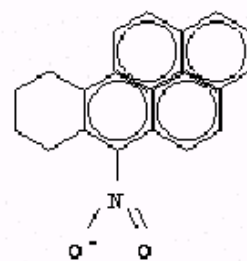


nitrofurazone

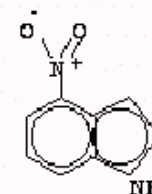


4-nitropenta[cd]pyrene

Negative



6-nitro-7,8,9,10-tetrahydrobenzo[a]pyrene



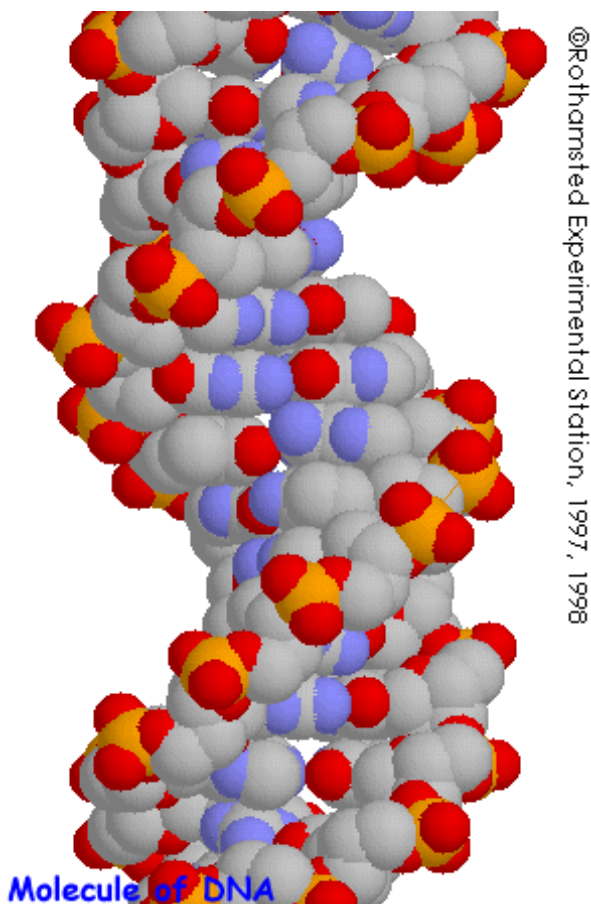
4-nitroindole

Result:

structural
indicator



Bioinformatics - prostor. uspořádání bílkovin



Bílkoviny = řetězce aminokyselin tvořících složité prostor. útvary.

- Posloupnost aminokyselin = **primární struktura**.
- *Lze předpovědět prostorovou strukturu molekuly na základě info. o její primární struktuře?*
- **Interpretace NMR spektra** - rozdělení do 23 strukturních typů. Klasické metody - 80% úspěšnost, **ILP 90%** - odpovídá výkonu zkušeného odborníka

Bioinformatika - karcinogenicita

- 230 aromatických a heteroaromatických dusíkatých sloučenin

188 sloučenin (lze je dobře klasifikovat regresí v rámci atributové reprezentace)

+ 42 RU sloučenin (regression-unfriendly skupina).

- Na RU skupině se prokázaly výhody relační reprezentace:

Hypotéza navržená PROGOLem dosahovala přesnosti 88%

zatímco klasické metody asi o 20 % méně.