

Practical "Introduction to Artificial Intelligence"

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Block 1: Prolog

Sheet 8: Files

Hints:

- *In Block 1 (Prolog) you do not have to submit your solutions to me. Just solve the excercises and discuss your problems and solutions. The aim of Block 1 is that you become familiar with the prolog programming.*
- *If you do not succed with a task, just delay it and try it again later. Some constructs need time to settle in the brain and will become easier as you get more experienced.*

Preparation (at home):

Read Chapter 12 of LearnPrologNow!.

Excercise 6.1

Read the SWI-prolog documentation for the relevant predicates, at least

- `consult`, `ensure_loaded`
- `open`, `close`
- `write/2`, `format/3`
- `read/2`, `at_end_of_stream/1`, `read_string/3`

Excercise 6.2

Consider the following prolog program, save it as `family1.pl`.

```
father(anton, john) .
father(paul, mary) .
father(john, peter) .
father(john, elisabeth) .
father(peter, agneta) .
mother(mary, peter) .
mother(mary, elisabeth) .
mother(elisabeth, agneta) .
mother(elisabeth, sarah) .

parent(X, Y) :- father(X, Y) .
parent(X, Y) :- mother(X, Y) .
ancestor(X, Y) :- parent(X, Y) .
ancestor(X, Y) :- parent(X, Z), ancestor(Z, Y) .
```

a) Write a predicate that reads in the rules from `family1.pl` and print it on the screen.

b) Write a predicate that reads in the rules from `family1.pl` and writes it to a file `family2.pl`. Load the file into prolog and test if it runs correctly. (e.g. `ancestor(anton, agneta)` is true two 2 times).

c) Write a predicate that reads in the rules from `family1.pl` and writes it to a file `family3.pl` so that each rules is preceed by a comment line which contains

- a running number (e.g. the number of the clause/line/...),
- the name of the predicate defined as well as its arity and
- the information whether this is a fact or a proper rule (i.e. of form `head :- body`).

Load the file into prolog and test if it runs correctly.

d) Write a predicate that reads in the rules from `family1.pl` and writes it to a file `family4.pl` so that

- the head of each rule is enriched by an additional parameter which contains a running number (e.g. the number of the clause/line/...).
- e.g. `pred1(X, Y) :-` would be changed to `pred1(X, Y, 7) :-`
- the body of each clause is changed in such a way that all predicates are added an additional argument, which is an anonymous variable.

Example: The program

```
father(anton, john) .
ancestor(X, Y) :- parent(X, Z), ancestor(Z, Y) .
```

would be changed to

```
father(anton, john, 1) .
ancestor(X, Y, 2) :- parent(X, Z, _), ancestor(Z, Y, _)
```

Load the file into prolog and test if it runs correctly. Trace your queries. You now should see in the trace which clause is used.

e) Write a predicate that reads in the rules from `family1.pl` and writes it to a file `family5.pl` so that each rule in the input file is transformed into a fact rule(`Name`, `Head`, `Body`), where

- `Head` is the head of the rule and
- `Body` is the body of the rule in form of the lists of all predicates and
- `Name` is an atom `rulex`, where `x` is a running number as before.

Example:

```
father(anton, john) .
ancestor(X, Y) :- parent(X, Z), ancestor(Z, Y) .
```

would be transformed into

```
rule(rule1, father(anton, john), []).
rule(rule2, ancestor(X, Y), [parent(X, Z), ancestor(Z, Y)]) .
```

Put the following lines into a new prolog file and load it.

```
:-consult(family5).

interpret(Query) :-
    rule(_, Query, Body),
    interpret_body(Body).

interpret_body([]).
interpret_body([Query1 | RemainingQueries]):-
    interpret(Query1),
    interpret_body(RemainingQueries).

interpret2(Query, NameOfRule -> BodyDerivation) :-
    rule(NameOfRule, Query, Body),
    interpret_body2(Body, BodyDerivation).

interpret_body2([], []).
interpret_body2([Query1 | RemainingQueries], [Derivation1 |
DerivationRemainder] ):-
    interpret2(Query1, Derivation1),
    interpret_body2(RemainingQueries, DerivationRemainder).
```

Execute `interpret(ancestor(anton, agneta))` What happens? Trace the execution. Try with other queries.

Repeat this with `interpret2(ancestor(anton, agneta), Derivation)`. What happens? Trace the execution.

Experiment with the interpreter:

- Change it so that the derivation tree contains the particular predicate call (i.e. The instantiated head) instead of the rule number.
- Change the order of the execution, e.g. execute the body from right to left.
- Whenever a predicate is executed, print it on the screen.

Exercise 5.3

a) Write a predicate that does the following:

- Prints a prompt on the screen,
- reads in an atom *a*,
- prints out all atoms *b*, for which `ancestor(a, b)` is true (see 5.2),
- and loops, i.e. shows the prompt again until the user types *bye*.

b) Write a predicate that behaves as the one in a) but

- the input is now a query that is executed (use `call/1`) and the result is printed
 - e.g. when typing in `ancestor(anton, agneta)` the system should answer *yes*
- Can you extend your program so that it also outputs the variable substitutions?
- (advanced) Can you extend your program so that it is tolerant to errors, i.e. continues to prompt even if the input leads to a prolog error (illegal syntax, predicate not defined, ..)?

Exercise 5.4

Reconsider Exercise 6.2. Your program will not succeed for 4 disks due to heavily creation of duplicate states.

a) Log the execution of the algorithm to a file: after each iteration, print the fringe. Check for duplicate states.

b) Change your predicates so that you detect multiple states:

- Extend your predicate `search` by an additional argument `VisitedStates` that keeps the list of visited states.
- If you expand a node n , then
 - Add the state of n to the list of visited nodes.
 - Only insert successor nodes to the fringe that are not contained in `VisitedStates`.

c) Repeat your experiments.

- Check the log files that now no duplicate states occur.
- Run the algorithm with and without heuristics and compare the results.