## For Management & Informatics



## **FINAL EXAM**

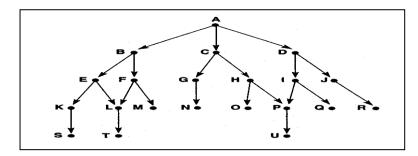
| Course Name: Artificial Intelligence | Term: First Academic Year: 2018-2019   |  |
|--------------------------------------|--|--|
| Course Code: CS6411                  | Date: 6/1/2019   |  |
| Intake & Major: Computer Science     | Time: 2 Hours  |  |
| 1                                    | The state of the s |  |

## Answer the following questions

## **Question One:**

Write and trace the algorithms of Backtrack, Depth-first and Breadth-First search by using the following graph. Begin from state A to goal R.

Keep track of the successive value of NSL, SL, CS and DE.



Where SL for state list, NSL for new state list, CS for current state and DE for dead ends.

## function backtrack;

```
begin
  SL := [Start]; NSL := [Start]; DE := []; CS := Start;
                                                                    % initialize:
                                             % while there are states to be tried
  while NSL ≠ [] do
    begin
      if CS = goal (or meets goal description)
        then return SL;
                                      % on success, return list of states in path.
      if CS has no children (excluding nodes already on DE, SL, and NSL)
        then begin
          while SL is not empty and CS = the first element of SL do
            begin
              add CS to DE;
                                                    % record state as dead end
              remove first element from SL;
                                                                    %backtrack
              remove first element from NSL;
              CS := first element of NSL;
            end
          add CS to SL:
        end
        else begin
          place children of CS (except nodes already on DE, SL, or NSL) on NSL;
          CS := first element of NSL;
          add CS to SL
        end
    end:
    return FAIL;
end.
```

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function breadth\_first\_search;

7.

8.

9.

```
begin
  open := [Start];
                                                                        % initialize
  closed := []:
                                                                  % states remain
  while open ≠ [] do
    begin
      remove leftmost state from open, call it X;
        if X is a goal then return SUCCESS
                                                                     % goal found
          else begin
            generate children of X;
            put X on closed;
            discard children of X if already on open or closed;
                                                                     % loop check
             put remaining children on right end of open
                                                                         % queue
          end
    end
                                                                   % no states left
  return FAIL
end.
1.
    open = [A]; closed = []
    open = [B,C,D]; closed = [A]
2.
    open = [C,D,E,F]; closed = [B,A]
3.
    open = [D,E,F,G,H]; closed = [C,B,A]
4.
    open = [E,F,G,H,I,J]; closed = [D,C,B,A]
5.
    open = [F,G,H,I,J,K,L]; closed = [E,D,C,B,A]
6.
```

open = [G,H,I,J,K,L,M] (as L is already on open); closed = [F,E,D,C,B,A]

open = [H,I,J,K,L,M,N]; closed = [G,F,E,D,C,B,A]

and so on until either U is found or **open** = []

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```
begin
  open := [Start];
                                                                             % initialize
  closed := [];
                                                                       % states remain
  while open ≠ [] do
    begin
       remove leftmost state from open, call it X;
       if X is a goal then return SUCCESS
                                                                          % goal found
         else begin
           generate children of X;
           put X on closed;
           discard children of X if already on open or closed:
                                                                          % loop check
           put remaining children on left end of open
                                                                                % stack
         end
    end:
  return FAIL
                                                                        % no states left
end.
```

```
1.
    open = [A]; closed = []
    open = [B,C,D]; closed = [A]
2.
    open = [C,D,E,F]; closed = [B,A]
3.
    open = [D,E,F,G,H]; closed = [C,B,A]
4.
    open = [E,F,G,H,I,J]; closed = [D,C,B,A]
5.
    open = [F,G,H,I,J,K,L]; closed = [E,D,C,B,A]
6.
    open = [G,H,I,J,K,L,M] (as L is already on open); closed = [F,E,D,C,B,A]
7.
    open = [H,I,J,K,L,M,N]; closed = [G,F,E,D,C,B,A]
8.
    and so on until either U is found or open = []
9.
```

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# **Question Two:**

Write and trace of the execution of best\_first\_search by using evaluation function f(n) for following state space generated in heuristic search of the 8-puzzle graph.

Consider the evaluation function f(n)=g(n)+h(n),

where n is any state encountered in the search.

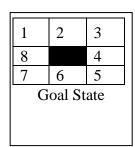
g(n) is the cost of n from the start state.

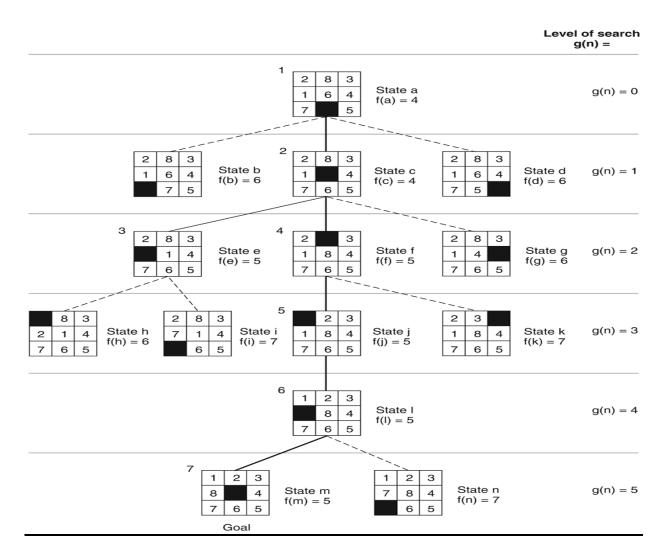
h(n) is the heuristic estimate of the cost of going from n to the goal.

What is the role of the g(n), answer by another trace with f(n)=h(n).

Write your comments.







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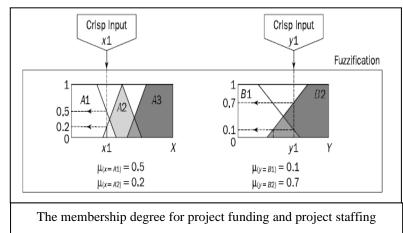
## **Question Three:**

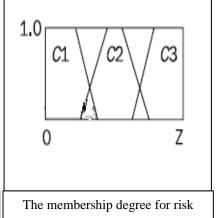
Draw the basic structure (Mamdani-style) that simulate the Fuzzy inference(Fuzzification, Rule evaluation, Aggregation of rule consequents, Defuzzification) for the following rules

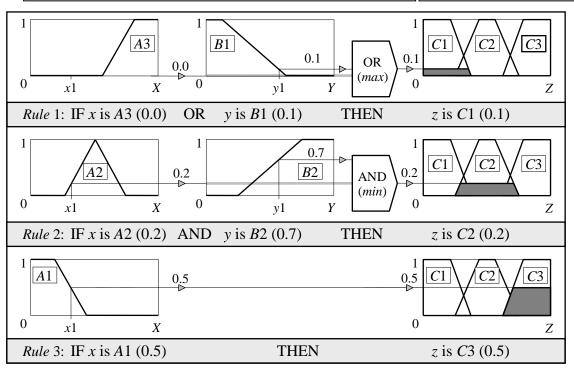
- 1. IF project\_funding is adequate OR project\_staffing is small THEN risk is low
- 2. IF project\_funding is marginal AND project\_staffing is large THEN risk is normal
- 3. IF project\_funding is inadequate THEN risk is high

Suppose the ranges of project funding and project staffing between 1 to 100 per cent. And the crisp input x1=0.35 and y1=0.6

The membership degree for project funding and project staffing and risk as follow:





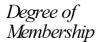


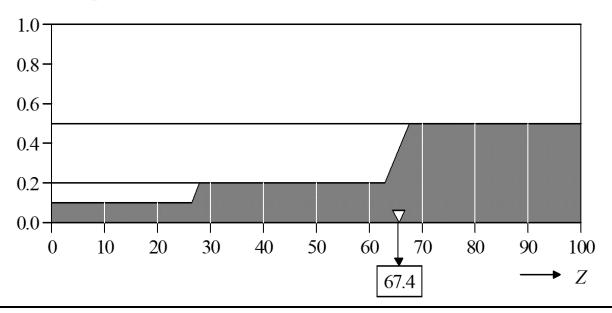
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$$COG = \frac{(0+10+20)\times0.1 + (30+40+50+60)\times0.2 + (70+80+90+100)\times0.5}{0.1+0.1+0.1+0.2+0.2+0.2+0.2+0.5+0.5+0.5+0.5+0.5} = 67.4$$





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## **Question Four:**

- **a.** Define the ontology.
- **b.** How to develop an ontology.
- c. Consider the following tabular data (bibliography) and answer the following questions.

| Books |                                     |                 |           |                 |                       |  |
|-------|-------------------------------------|-----------------|-----------|-----------------|-----------------------|--|
| Id    | Title                               | Author          | Publisher | Category        | ISBN                  |  |
| 1     | Introduction to computers           | Jim Hendler     | springer  | Semantic Web    | 978-0-12-385965-<br>5 |  |
| 2     | Essential bioinformatics            | JIN XIONG       | springer  | Bioinformatics  | 978-0-470-02001-<br>2 |  |
| 3     | Pattern discovery in bioinformatics | David L. Olson  | springer  | Data Mining     | 978-3-540-76916-<br>3 |  |
| 4     | Advanced databases                  | Borko Furht     | springer  | Cloud Computing | 978-1-4419-6523-<br>3 |  |
| 5     | Algorithms of bioinformatics        | Frédéric Dardel | springer  | Bioinformatics  | 978-0-470-12321-<br>2 |  |

- **1.** Convert the tabular data into XML formats where the "id", "title", "author", "publisher", "category" and "ISBN" are attributes for the element book and the root element is library.
- 2. Convert the tabular data into the RDF representation using the following:
  - i. The global URI for the rdf namespace is http://www.w3.org/1999/02/22-rdf-syntax-ns#.
  - ii. "bib" stands for <a href="http://www.amazon.com/books-used-books-textbooks">http://www.amazon.com/books-used-books-textbooks</a>. (Convert 2 records only).
    - **a.** Define the ontology.

An **ontology** is a formal explicit description of concepts in a domain of discourse (**classes** (sometimes called **concepts**)), properties of each concept describing various features and attributes of the concept (**slots** (sometimes called **roles** or **properties**)), and restrictions on slots (**facets** (sometimes called **role restrictions**)).

An ontology together with a set of individual **instances** of classes constitutes a **knowledge base**. In reality, there is a fine line where the ontology ends and the knowledge base begins.

**b.** How to develop an ontology defining classes in the ontology arranging the classes in a taxonomic (subclass–superclass) hierarchy, defining slots and describing allowed values for these slots, filling in the values for slots for instances.

```
c. 1.<?xml version="1.0" encoding="UTF-8"?>
   tibrary>
   <book>
   <ID> 1 </ID>
   <title> Introduction to computers</title>
   <author>Jim Hendler</author>
   <publisher>springer</publisher>
   <category>Semantic Web</category>
   <ISBN>978-0-12-385965-5</ISBN>
   </book>
   <book>
   <ID> 2 </ID>
   <title>Essential Bioinformatics</title>
   <author>JIN XIONG</author>
   <publisher>springer</publisher>
   <category> Bioinformatics</category>
   <ISBN>978-0-470-02001-2</ISBN>
   </book>
   </library>
```

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```
2.<rdf:RDF
xmlns:bib="http://www.amazon.com/textbooks#"
xmlns:rdf="http://www.w3.org/1999/02/22-rdf-syntaxns#"
<bib:Book rdf:about="http://www.amazon.com/textbooks#Book1">
<bib:id>1</bib:id>
<bib:title> Introduction to computers </bib:title>
<bib:author> Jim Hendler</bib:author>
<bib:publisher> springer</bib:publisher>
<bib:category> Semantic Web</bib:category>
<bib:ISBN>978-0-12-385965-5</bib:ISBN>
</bib:Book>
<bib:Book rdf:about="http://www.amazon.com/textbooks#Book2">
<br/><bib:id>2</bib:id>
<bib:title>Essential Bioinformatics
<br/><bib:author> JIN XIONG</bib:author>
<bib:publisher> springer</bib:publisher>
<bib:category> Bioinformatics </bib:category>
<br/><bib:ISBN>978-0-470-02001-2</bib:ISBN>
</bib:Book>
</rdf:RDF>
```

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## **Question Five:**

By using the tabular data in the question 3 answer the following:

- **a.** By using the graph representation, represent the relation between
  - i. The publisher and the title of the book (published)
  - ii. The title and the author of the book (wroteby)
  - iii. Merge the two graphs in one graph.
- **b.** Write SPARQL query to determine the following
  - i. The books which published by the springer.
  - ii. The publisher of the book "Pattern Recognition"
  - iii. Write the answer of the following query:

**SELECT** ?who

**WHERE** {:springer :published ?what . ?what :wroteby ?who .}

. What ! Wrote y ! Who ! j

a.By using the graph representation, represent the relation between



B. The  $\underline{\text{title}}$  and the  $\underline{\text{author}}$  of the book ( $\underline{\text{wroteby}}$ )



C. Merge the two graphs in one graph.



#### b.

i. The books which published by the springer.

Representation: SELECT?what

WHERE {:Springer:Published?what.}

ii. The publisher of the book "Pattern Recognition"

SELECT?who

WHERE {: Pattern Recognition: Published By ?who.}

iii. Write the answer of the following query:

SELECT?who

WHERE {:springer:published?what

?what:WroteBy?who.}

**RESULT** 

| Author          | Title                                  |
|-----------------|--|
| Jim Hendler     | Bioinformatics Genomics& post genomics |
| JIN XIONG       | Introduction of bioinformatics         |
| David L. Olson  | Introduction to Semantic Web           |
| Borko Furht     | Advanced Data Mining Techniques        |
| Frédéric Dardel | Handbook of Cloud Computing            |