## EC-350 AI and Decision Support Systems

# Week 6 Local Search Algorithms

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Acknowledgement: Lecture slides material from Stuart Russell

## Local Search Algorithms

- In many problems the path to the goal is irrelevant
  - 8 queens problem, integrated circuit design, automatic prog
  - Factory floor layout, Telecom network optimization
- Algorithms to solve such problems are called local search algorithm
  - Start from the current state and then gradually only move to the neighbor of the state
  - Don't have to remember all the previous states

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#### Local Search Algorithms

- They use very little memory usually constant amount
- They can often find reasonable solution in a large or infinite space for which systematic algorithms are unsuitable
- Useful for solving pure optimization problems in which the aim is to find best state according to an objective function

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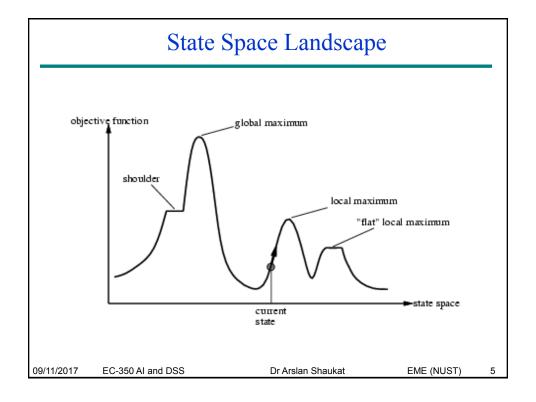
#### Local Search Algorithms

- Consider the state space as a land scape
- Locations and elevation defined by state and value of the objective function respectively
- If elevation is a cost then aim is to find the lowest valley
   a global minimum
- If elevation is objective function then aim is to find the highest peak – a global maximum
- A complete local search algorithm always finds a goal if one exist, an optimal always find a global min/max

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## Genetic Algorithms

- Organisms (animals or plants) produce a number of offsprings which are almost, but not entirely, like themselves
  - Variation may be due to mutation (random changes)
  - Variation may be due to the fact that an offspring has some characteristics from each parent)
- Some of these offspring may survive to produce offspring of their own—some won't
  - The "better adapted" offspring are more likely to survive
  - Over time, later generations become better and better adapted
- Genetic algorithms use this same process to "evolve" better programs.

#### Genes and Chromosomes

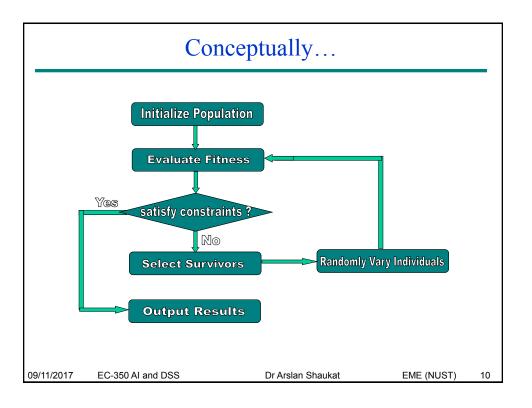
- Genes are the basic "instructions" for building an organism
- A chromosome is a sequence of genes
- Biologists distinguish between an organism's genotype (the genes and chromosomes) and its phenotype (what the organism actually is like)
  - Example: You might have genes to be tall, but never grow to be tall for other reasons (such as poor diet)
- Similarly, "genes" may *describe* a possible solution to a problem, without actually *being* the solution.

#### **Quick Overview**

- Developed: USA in the 1970's
- Early names: J. Holland, K. DeJong, D. Goldberg
- Typically applied to:
  - discrete optimization
- Attributed features:
  - Not too fast

#### The Basic GA

- Start with a large "population" of randomly generated "attempted solutions" to a problem
- Repeatedly do the following:
  - Evaluate each of the attempted solutions through a fitness function
  - Keep a subset of these solutions (ones with the "best" fitness)
  - Use these solutions to generate a new population
- Quit when you have a satisfactory solution (or you run out of time)



#### A Really Simple Example

- Suppose your "organisms" are 32-bit computer words
- You want a string in which all the bits are ones
- Here's how you can do it:
  - Create 100 randomly generated computer words
  - Repeatedly do the following:
    - Count the 1 bits in each word
    - Exit if any of the words have all 32 bits set to 1
    - Keep the ten words that have the most 1s (discard the rest)
    - From each word, generate 9 new words as follows:
      - Pick a random bit in the word and toggle (change) it
- Note that this procedure does not guarantee that the next "generation" will have more 1 bits, but it's likely.

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#### A More Realistic Example (I)

- Suppose you have a large number of (x, y) data points
  - For example, (1.0, 4.1), (3.1, 9.5), (-5.2, 8.6), ...
- You would like to fit a polynomial (of up to degree 5) through these data points
  - That is, you want a formula  $y = ax^5 + bx^4 + cx^3 + dx^2 + ex + f$  that gives you a reasonably good fit to the actual data
  - Here's the usual way to compute goodness of fit: Compute the sum of (actual y - predicted y)<sup>2</sup> for all the data points: The lowest sum of differences represents the best fit
- There are some standard curve fitting techniques, but let's assume you don't know about them
- You can use a genetic algorithm to find a "pretty <sub>1</sub>good" solution.

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#### A More Realistic Example (II)

- Your formula is  $y = ax^5 + bx^4 + cx^3 + dx^2 + ex + f$
- Your "genes" are a, b, c, d, e, and f
- Your "chromosome" is the array [a, b, c, d, e, f]
- Your evaluation function for *one* array is:
  - For every actual data point (x, y), (red means "actual data")
    - Compute  $\dot{y} = ax^5 + bx^4 + cx^3 + dx^2 + ex + f$
    - Find the sum of  $(y \dot{y})^2$  over all x
    - The sum is your measure of "badness" (larger numbers are worse)
  - Example: For [0, 0, 0, 2, 3, 5] and the data points (1, 12) and (2, 22):
    - $\dot{y} = 0x^5 + 0x^4 + 0x^3 + 2x^2 + 3x + 5$  is 2 + 3 + 5 = 10 when x is 1
    - $\dot{y} = 0x^5 + 0x^4 + 0x^3 + 2x^2 + 3x + 5$  is 8 + 6 + 5 = 19 when x is 2
    - $(12 10)^2 + (22 19)^2 = 2^2 + 3^2 = 13$
    - If these are the only two data points, the "badness" of [0, 0, 0, 2, 3, 5] is 13

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### A More Realistic Example (III)

- Your algorithm might be as follows:
  - Create 100 six-element arrays of random numbers
  - Repeat 500 times (or any other number):
    - For each of the 100 arrays, compute its badness (using all data points)
    - Keep the ten best arrays (discard the other 90)
    - From each array you keep, generate nine new arrays as follows:
      - Pick a random element of the six
      - Pick a random floating-point number between 0.0 and 2.0
      - Multiply the random element of the array by the random floating-point number
  - After all 500 trials, pick the best array as your final answer.

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#### Generating a Population

- Through one parent:
  - In the previous example, each solution had only one parent
  - The only way to introduce variation was through mutation (random changes)
- Through two parents:
  - Each solution has two parents
  - New solutions are produced by combining parts of the chromosomes of each parent – more commonly known as crossover.

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## The Really Simple Example Again

- Suppose your "organisms" are 32-bit computer words, and you want a string in which all the bits are ones
- Here's how you can do it:
  - Create 100 randomly generated computer words
  - Repeatedly do the following:
    - Count the 1 bits in each word
    - Exit if any of the words have all 32 bits set to 1
    - Keep the ten words that have the most 1s (discard the rest)
    - From each word, generate 9 new words as follows:
      - Choose one of the other words
      - Take the first half of this word and combine it with the second half of the other word

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#### The Example Continued

- Half from one, half from the other:
   0110 1001 0100 1110 1010 1101 1011 0101
   1101 0100 0101 1010 1011 0100 1010 0101
   0110 1001 0100 1110 1011 0100 1010 0101
- Or we might choose "genes" (bits) randomly:
   0110 1001 0100 1110 1010 1101 1011 0101
   1101 0100 0101 1010 1011 0100 1010 0101
   0100 0101 0100 1010 1010 1100 1011 0101
- Or we might consider a "gene" to be a larger unit:
   0110 1001 0100 1110 1010 1101 1011 0101
   1101 0100 0101 1010 1011 0100 1010 0101
   1101 1001 0101 1010 1010 1101 1010 0101

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#### Comparison of Simple Examples

- In the simple example (trying to get all 1s):
  - The two-parent-no mutation approach, if it succeeds, is likely to succeed much faster
    - Because up to half of the bits change each time, not just one bit
  - However, with no mutation, it may not succeed at all
    - By pure bad luck, maybe *none* of the first (randomly generated) words have (say) bit 17 set to 1
      - Then there is no way a 1 could ever occur in this position
    - Another problem is lack of genetic diversity
      - Maybe some of the first generation did have bit 17 set to 1, but none of them were selected for the second generation
- The best technique *in general* turns out to be a twoparent approach with a *small* probability of mutation

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#### **Un-Directed Evolution**

- In the previous examples, child organisms were formed randomly
  - We didn't choose the "best" gene from each parent
  - That's how biological evolution works: it's not necessary that a child will inherit only the best characteristics of both the mother and father.

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### Genetic Algorithms

- A state is represented as a string over a finite alphabet (often a string of 0s and 1s)
- Start with *k* randomly generated states (*population*)
- Evaluation function (*fitness function*). Higher values for better states.
- A successor state is generated by combining two parent states
- Produce the next generation of states by
  - selection,
  - crossover, and
  - mutation

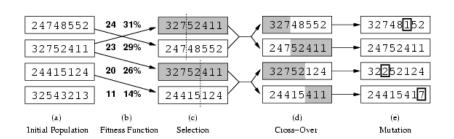
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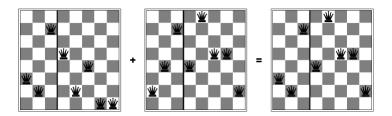
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## 8 Queens Example



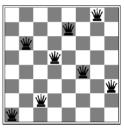
- Fitness function: number of non-attacking pairs of queens (min = 0, max = 28)
- 24/(24+23+20+11) = 31%
- 23/(24+23+20+11) = 29% etc

## Example



#### **Fitness Function**

- h: use the number of attacking pairs of queens.
- There are 28 pairs of different queens, so solutions have fitness 28. (Basically, **fitness function** is 28 h)
- for example, fitness of the state below is 27 (queens in columns 4 and 7 attack each other)



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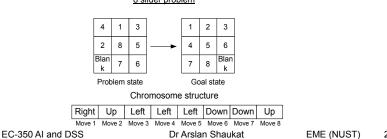
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### 8 Slider Example

- In a block slider problem, player bring blocks to their goal state, using 8 moves of the chromosome.
- Each Move in chromosome represents movement of the blank space. E.g.in the chromosome shown
  - if Move 1 = Right, the blank space will come to right, and block 7 will move to left
  - Move 2=Up, so blank space will move up and block 8 will come down.
  - Move 3=Left, so blank space will move left and block 2 will move right



#### Example

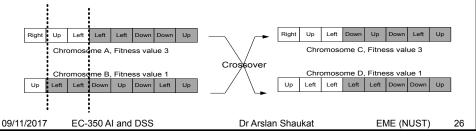
- In generation1: Find fitness, and select pairs for crossover.
- In generation 2: Crossover, Find fitness, and worst chromosome will mutate, rest will remain as it is.
- In generation 3: Find fitness and select pairs for crossover.
- In generation 4: Crossover and find fitness
- Fitness:
  - The number of moves in which BLANK space can move is fitness. E.g. If Move 1 = Up and Move 2 = Left. Then blank space will move just one space, up, and cannot go left after that. So fitness value is 1.

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#### Example

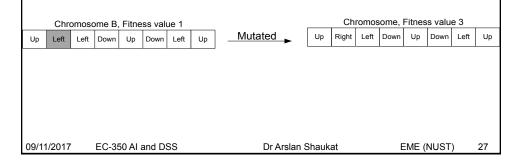
#### Crossover:

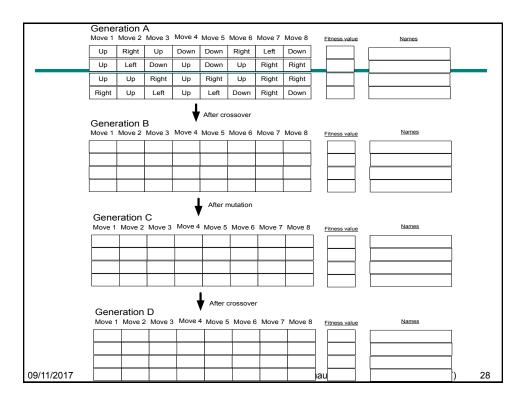
- Arrange chromosome in order of fitness.
- Pair 1: 1st and 4th
- Pair2: 2<sup>nd</sup> and 3<sup>rd</sup>
- Crossover point is chosen at the maximum of both fitness values, i.e. out of fitness 1 and fitness 3, the crossover point was after 3

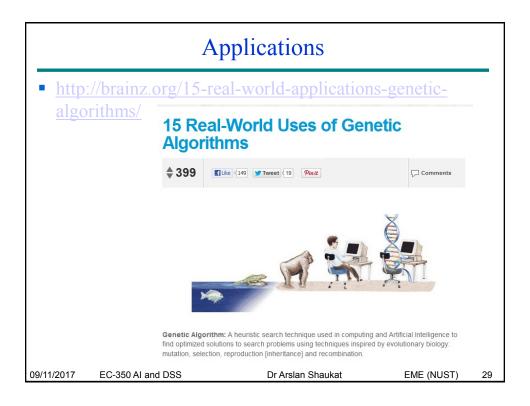


## Example

- Mutation:
- For mutation, the worst chromosome changes its (first problematic) direction. i.e.
- Up changes to down, and vice versa and left changes to right and vice versa







## Assignment# 2

- Finding solution with Genetic Algorithm
- Submission: After 2 weeks

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