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Biologically inspired computing - Lecture 3
Representations
(Genetic algorithms \& Genetic programming)


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Optimization problems

- Continuous optimization
- 0-1 knapsack problem
- Other knapsack problems
- Travelling salesman problem
- Task solving problems

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## This lecture

- Representations
- Recombination
- Mutation

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## Real-valued representations

- As shown in the previous lecture
- Represents continuous solution spaces
- The solution parameters are often accompanied by strategy parameters for adaptive normal distribution-based mutation

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## Binary representation

- The representation used in the simple genetic algorithm (SGA)
- Directly inspired by low-level encoding in DNA
- Uses a binary $(0,1)$ coding instead of the quaternary ( $\mathrm{G}, \mathrm{T}, \mathrm{A}, \mathrm{C}$ ) coding used in nature

\section*{| 0 | 1 | 1 | 0 | 1 | 0 | 0 | 0 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |}

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Permutation representation

- Used to solve problems like the travelling salesman
- Known set of actions (go to town X)
- Want to optimize their sequence

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## Integer representation

- Each element is directly coded as an integer - Usually restricted to some pre-defined ranges



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## Tree representation

- Tree representations of programs or arithmetic expressions
- Mainly used in genetic programming


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## Representations

def evol ve() :
P. $x=i$ nitialize_populationd
P.fitness $=$ eval uate(P. x )
while not_done():
$Q \cdot x=$ mutate ( $Q \times \times$ )
$x=$ mut ate $Q \times \times$
= survival ( $\mathrm{P}, \mathrm{Q}$ )
return best(P). . $x$

- The central concepts in evolutionary algorithms are independent of representation
- Mutation and recombination must be tailored to the representation used

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## Indirect representations

- Most problems will have a fixed solution representation associated with it
- However, sometimes it is beneficial to evolve solutions using a different representation and then transform them to do the evaluation

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Expanding the analogy

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Binary representation operators

| Optimization | Biology |
| :--- | :--- |
| Candidate solution | Individual |
| Representation used in the EA | Genotype, chromosome |
| Problem-defined representation | Phenotype |
| Position/element of the genotype | Locus, gene |
| Old solution | Parent |
| New solution | Offspring |
| Solution quality | Fitness |
| Random displacements added to offspring | Mutation |
| Search strategy | Mutation rate, gene robustness |
| A set of solutions | Population |


$\square$ | 0 | 1 | 1 | 0 | 1 | 0 | 0 | 0 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |

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## Bit flip mutation

- Each bit is inverted with a probability $p_{m}$


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## N-point crossover

- $N$ random points in the genotype is chosen
- At each point the source parent changes
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## Uniform crossover

## Binary coding of integers

- Encoding integers as blocks of a binary string has been quite common each position
- Identical to discrete recombination
- Keeps the analogy to DNA clean
- Problematic because mutations are not local
- Small changes to the solution are not more probable
- The result of flipping a single bit varies enormously with bit position and the value of all bits that encode the same integer

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## Integer representation operators

- Can use the same crossover operators as the binary representation


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## Random reset mutation

- Each element is reset with probability $p_{m}$ to a random number in the range


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## Integer coding of symbols

- Sometimes a vector of symbols with no clear order is the most reasonable representation choice
- In such cases, the symbols are usually enumerated and treated as integers, but without using the creep mutation


| 0 | 5 | 8 | 3 | 1 | 3 | 7 | 5 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |


| 0 | 6 | 8 | 2 | 1 | 3 | 8 | 4 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |

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Real-valued representation operators


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## Uniform mutation

- Each element has a probability $p_{m}$ of being replaced with a number from some range

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## Arithmetic recombination

## Single arithmetic recombination

- Arithmetic recombination is applied to only one $k$
- Picks one or more random positions $k$ and replaces those elements with the interpolation $\alpha x_{k}+$ $(1-\alpha) y_{k}$, where $\alpha$ is either a fixed number or a random variable
- Intermediate recombination: $\alpha$ is 0.5 for all $k$


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## Whole arithmetic recombination

- Arithmetic recombination is applied with the same $\alpha$ to all $k$


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## Permutation representation

- Special mutation/recombination operators
- Each item should appear once and only once
- Result should be "close" to the original solution(s)

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## Insert mutation

- Two random elements are picked
- The second is placed right after the first



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## Scramble \& invert mutation

- Two random points are selected
- The order of the elements in between is scrambled (scramble mutation) or reversed (invert mutation)


| 1 | 2 | 5 | 4 | 3 | 6 | 7 | 8 | 1 | 5 | 4 | 3 | 2 | 6 | 7 | 8 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |

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## Partially mapped crossover (PMX)

- Two random points are chosen
- All elements between the points in parent A are copied to the offspring

| 3 | 4 | 5 | 6 | 7 | 8 | 1 | 2 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |

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## Partially mapped crossover (PMX)

- For each element $x$ in parent $B$ between those points that is not in parent $A$
- Place it in the position in $B$ of the element with the same position in $A$ as $x$ has in $B$

| 3 | 4 | 5 | 6 | 7 | 8 | 1 | 2 |  |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
|  | 7 | 8 | 2 | 6 | 5 | 1 | 4 | 3 | 

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## Partially mapped crossover (PMX)

- For each element $x$ in parent $B$ between those points that is not in parent $A$
- Place it in the position in $B$ of the element with the same position in $A$ as $x$ has in $B$
- If that position is occupied, do one more redirection

| 3 | 4 | 5 | 6 | 7 | 8 | 1 | 2 | 7 | 8 | 2 | 6 | 5 | 1 | 4 | 3 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | | 2 | 4 |  |  |  |  |  |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| 2 | 4 | 5 | 6 | 7 |  | 8 |

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## Partially mapped crossover (PMX)

- Finally, the missing elements are copied from their places in parent $B$

| 3 | 4 | 5 | 6 | 7 | 8 | 1 | 2 | 7 | 8 | 2 | 6 | 5 | 1 | 4 | 3 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | | 2 | 4 | 5 | 6 | 7 | 1 | 8 | 3 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |

## Order crossover

- Two random points are chosen
- All elements between the points in parent A are copied to the offspring

| 3 | 4 | 5 | 6 | 7 | 8 | 1 | 2 | 7 | 8 | 2 | 6 | 5 | 1 | 4 | 3 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |



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## Edge crossover

- Heuristic to preserve as many edges as possible
def edge_xo(PA, PB, N):
e = construct_edge_table()
$\mathrm{k}=\mathrm{random} \mathrm{N}$ )
(fin inge( 1 , N)
e. remove( $k$ )
if e.empty $(k): k=\operatorname{reverse}(x)[-1]$
if e. empty $(k): k=\operatorname{dranc}($, e. remaining()
el se:
k = e. pick_common(k) or aram 1, e. pick_shortest(k))

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$$

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## Order crossover

- The rest of the elements are copied from parent $B$ in the order starting from the second random point



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Cycle crossover

- Identify next cycle
- Copy from parent $A$ and $B$ to offspring $B$ and $A$


| 3 | 4 | 5 | 6 | 7 | 8 | 1 | 2 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |$\quad$| 3 | 8 |
| :--- | :--- |

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Tree representation operators

## Cycle crossover

- Identify last cycle
- Copy from parent A and B to offspring A and B
$\square$
A

| 3 | 4 | 5 | 6 | 7 | 8 | 1 | 2 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |$\quad$| 3 | 8 |
| :--- | :--- |



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## Tree mutation

- Take a random node and replace it by a new randomly generated subtree


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## Tree crossover

- Take one random node from each parent and exchange them


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## Bloat in tree representations

- Larger trees will have greater fitness on average in most cases
- Without any active countermeasures the population will tend to grow indefinitely


