

C0656 / COMPUTATIONAL INTELLIGENCE IN BUSINESS, ECONOMICS AND FINANCE

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housekeeping

- sample solutions to class exercises will be available on Moodle a week after the class
 - try the tasks first!
- quizzes are available on Moodle for you to check your understanding
 - there are no marks associated with them
 - automatic feedback

outline:

1. a little bit more on GAs

2. another GA example

why GA works?

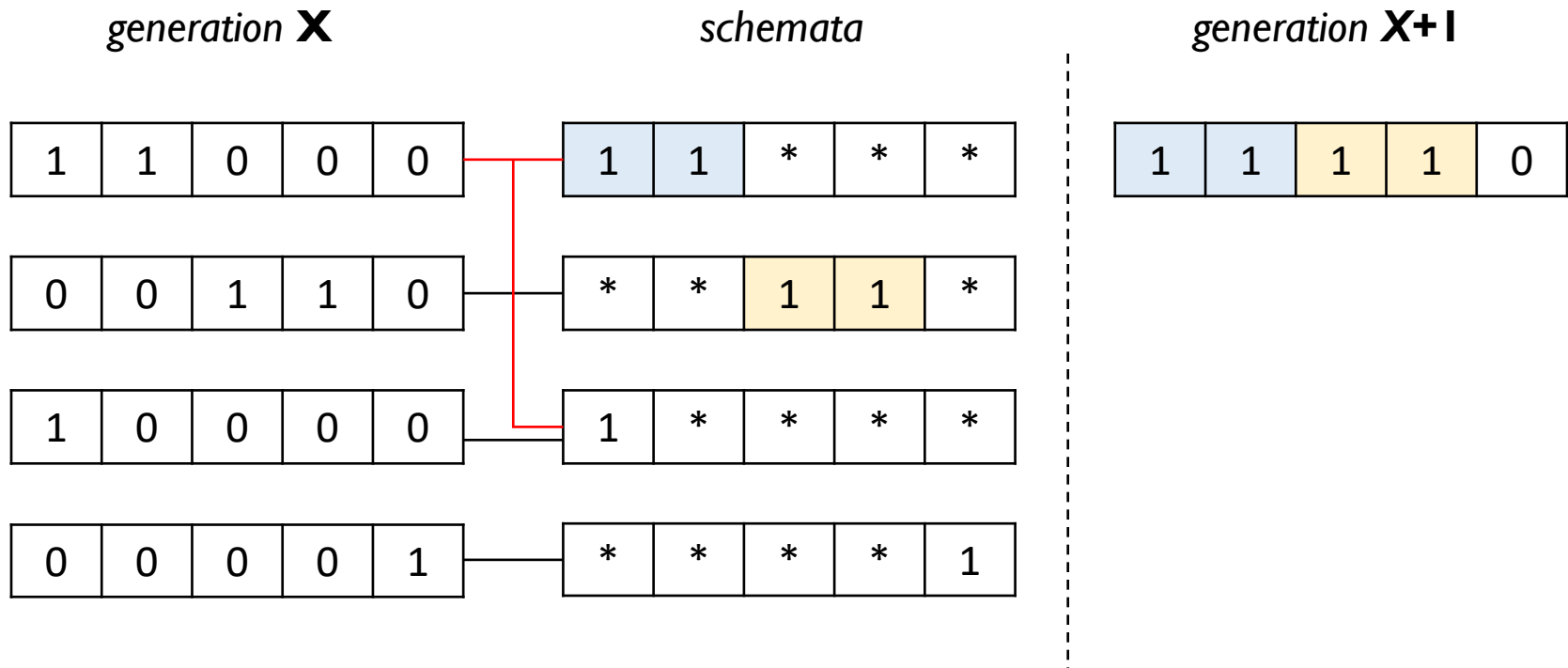
- many random choices
- ...but still we see improvements (evolution)
- **stochastic** but different than random search
 - selection is based on fitness
 - ... therefore search is biased towards good regions of the search space
- need for multiple runs to estimate the “true” performance of a GA

why GA works?

- evolution preserves good building blocks
 - blocks of genes in a chromosome
- building blocks are propagated in the population by the fitness-based selection
 - crossover of 2 good individuals with different building block will likely result in an even better offspring
- there are theories that try to characterize the evolution of a GA
 - schema theorem and building block hypothesis
 - ...but there is not a universally accepted theory

building block hypothesis:

- better solutions are created from the best **partial** solutions of past generations



observations:

- you might have noticed that the best fitness of the population can **fluctuate** – e.g., be lower than the best fitness of the previous generation
- roulette wheel selection can be **affected** if you have extreme fitness values – e.g., one individual having a fitness value much higher than the rest of the population
- more options of genetic operators?

elitism:

- **best** individual of a population is guaranteed to be carried over to the next
- solution quality will **not** decrease from one generation to the next

tournament selection:

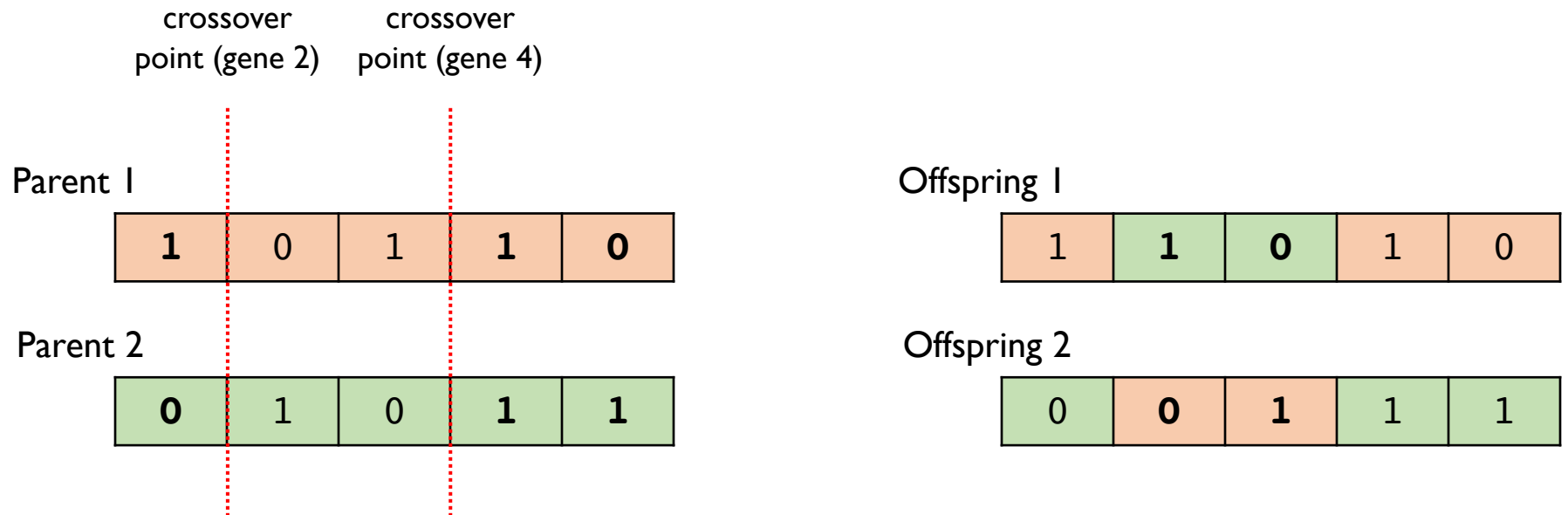
- small **subset** of k individuals is chosen at random
- best individual in this set is selected (tournament winner)
- k = tournament size (user-specified parameter)
- easy to control the selection pressure – the higher the value of k , the higher is the selective pressure

what happens if $k = 1$?

what happens if k = population size?

two-point crossover:

- same principle as the one-point variation, but uses 2 crossover points



uniform crossover:

- combines genes sampled uniformly from the 2 parents
 - each gene is subject to crossover subject to a probability
 - avoids positional bias

Parent 1

1	0	1	1	0
---	---	---	---	---

Parent 2

0	1	0	1	1
---	---	---	---	---

Offspring 1

0	0	0	1	0
---	---	---	---	---

Offspring 2

1	1	1	1	1
---	---	---	---	---

bit string mutation:

- each gene has a probability of $\frac{1}{length}$ of being mutated

Parent I

1	0	1	1	0
---	---	---	---	---

Offspring I

1	1	0	1	1
---	---	---	---	---



GA examples:

- **Goldberg's book (1989)**
 - **real-value encoding**

GA example:

- **toy** problem: finding the maximum value of the function x^2 in the interval $[0..31]$
- individual encoding: five bits representing x in $[0..31]$
- fitness function: x^2 (the larger the fitness, the better the individual)
 - decode 5-bits (genotype) and then compute x^2 (phenotype)

GA example:

- roulette wheel selection
- genetic operators:
 - one-point crossover
 - point mutation
- termination criteria:
 - optimal solution found
 - maximum number of generations

GA example:

- initial population (randomly generated):

1:	0	1	1	0	1
<hr/>					
2:	1	1	0	0	0
<hr/>					
3:	0	1	0	0	0
<hr/>					
4:	1	0	0	1	1

GA example:

- measuring the fitness of each individual in the population:

						x	x ² (fitness)	% of total fitness
1:	0	1	1	0	1	13	169	14.4
2:	1	1	0	0	0	24	576	49.2
3:	0	1	0	0	0	8	64	5.5
4:	1	0	0	1	1	19	361	30.9

decoding individual 1: $0 \times 2^4 + 1 \times 2^3 + 1 \times 2^2 + 0 \times 2^1 + 1 \times 2^0 = 13$

GA example:

- suppose the selected individuals are:
 - one copy of individual 1
 - one copy of individual 4
 - two copies of individual 2(individual 3 was not selected)
- selected individuals undergo crossover
 - user-defined probability: about 90% - 95%
- they can also undergo mutation
 - user-defined probability: about 1% - 5%(lower mutation rate since in nature most mutations are harmful)

GA example:

- one-point crossover

crossover of individuals 1 and 2

1:	0	1	1	0	1	0	1	1	0	0
2:	1	1	0	0	0	1	1	0	0	1

crossover of individuals 2 and 4

1:	1	1	0	0	0	1	1	0	1	1
2:	1	0	0	1	1	1	0	0	0	0

GA example:

population at
generation 0

		x	x ² (fitness)
1:	0 1 1 0 1	13	169
2:	1 1 0 0 0	24	576
3:	0 1 0 0 0	8	64
4:	1 0 0 1 1	19	361
Average:			293
Maximum:			576

population at
generation 1

		x	x ² (fitness)
1:	0 1 1 0 0	12	144
2:	1 1 0 0 1	25	625
3:	1 1 0 1 1	27	729
4:	1 0 0 0 0	16	256
Average:			439
Maximum:			729

note: generation 1 has better individuals than generation 0 – the population evolves

another example:

- **task:** find the weights (real numbers) of the polynomial

$$\mathbf{a}x^3 + \mathbf{b}x^2 + \mathbf{c}x + \mathbf{d}$$

- how would you encode your individual?



another example:

- each weight is represented by a gene (string of real-valued numbers)

<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>
0.5	0.7	0.2	0.3

- crossover works the same
- mutation generates a new real-valued number

another example:

- fitness calculation will be based on **fitness cases**
 - values of **x** for which the value of the polynomial is known
 - fitness is a notion of how far from the desired value the individual is

<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>
0.5	0.7	0.2	0.3

$$= 0.5 \cdot x^3 + 0.7 \cdot x^2 + 0.2 \cdot x + 0.3$$

x	individual	correct value	fitness (error)
2	7.5	8.8	1.3
3	20.7	27.3	6.6
total:			7.9

the lower the error, the better the fitness

advantages:

- perform a **global** search in the search space
 - work with a population of individuals, rather than a single individual (candidate solution)
 - broader exploration of the search space, less likely to get trapped in a local maxima
- candidate solution is represented in a declarative way, independent of the search method
- easy to implement

considerations:

- do not offer any guarantee of finding the optimal solution, nor any lower bound on the quality
- generally computationally expensive
 - although can be easily parallelised
- several parameters need to be set
 - population size
 - number of generations
 - mutation / crossover probabilities
 - tournament size
 - ... among others

finishing off:

Practical class **this week:**

- more on **GA**
- finish the implementation of the GA before moving on to this week's exercise



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