

# Evolutionary Algorithms

## Applications of evolutionary algorithms

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# Outline

## 1. Parallelization of evolutionary algorithms

Which steps can be parallelized?

Island Model and Migration

Cellular Evolutionary Algorithms

Approach of Mühlenbein

## 2. Random numbers

# Parallelization of evolutionary algorithms

EA: fairly **expensive optimization methods** since one often has to work with

- large population (a few thousand up to several tens of thousands of individuals)
- large number of generations (a couple of hundreds)

**Advantage:** slightly better solution quality compared to other approaches

**Disadvantage:** unpleasantly long execution time

one way to improve this: **parallelization**

distribution of necessary operations on several processors

Questions:

- **Which steps can be parallelized?**
- **What additional, specialized techniques are inspired by a parallel organization of the algorithm?**

# Which steps can be parallelized?

## Creating an initial population:

- often easy to parallelize, because usually the chromosomes of the initial population are created randomly and independently of each other
  - attempt to prevent duplicate individuals may pose obstacles to a parallel execution
- fairly little importance overall though, because the initial population is created just once

## Evaluation of chromosomes:

- easily parallelizable because usually an individual is evaluated independently of any other ones
- even in prisoner's dilemma: process pairings in parallel

## Computing the (relative) fitness values or a ranking of the individuals

- central agent that collects and processes evaluations

## Which steps can be parallelized?

**Selection:** whether the selection of the individuals is parallelizable, depends heavily on the chosen selection method

- **expected value model** and **elitism:**  
all require to consider the population as a whole and therefore are difficult to parallelize
- **Roulette-wheel** and **rank-based selection:** can be parallelized after the initial step of computing the relative fitness values
- **Tournament selection:** best suited for a parallel execution, especially for small tournament sizes, because all tournaments are independent and thus can be held in parallel

# Which steps can be parallelized?

## **Applying genetic operators**

can easily be applied in parallel, since they affect only one (mutation) or two chromosomes (crossover), and are independent of any other chromosomes (If combined with tournament selection, a steady-state evolutionary algorithm can thus be parallelized very well)

## **termination criterion:**

simple test whether a certain number of generations is reached  
does not cause any problems

Termination criteria like

- the best individual of the population exceeds a user-specified fitness threshold, or
- the best individual has not changed (a lot) over a certain number of generations

need a central agency that collects this information about the individuals

# Island Model

if we require a selection method that causes some troubles for parallelization:

achieving a parallel execution by simply processing several independent populations

each population can be seen as inhabiting an island, therefore **island model**.

pure island  $\equiv$  executing the same evolutionary algorithm multiple times

yields results that are somewhat worse than those of a single run with a larger population

# Migration

one may consider exchanging individuals between the island populations at certain fixed points in time (*not* in every generation)

## **Migration** (Wanderung)

usually no direct recombination of chromosomes from different islands

after migration: Recombination of genetic material of one island with another



# Control the migration between islands

## Random model:

pairs of islands are chosen randomly, which then exchange some of their inhabitants

any two island can, in principle, exchange individuals

## Network model:

Islands are arranged into a network or graph

Individuals can migrate only between islands that are connected by an edge in the graph

Along which of the edges individuals are exchanged is determined randomly.

# Contest model

evolutionary algorithms that are applied on the islands differ in approaches and/or parameters

population size of an island is increased or decreased according to the average fitness of its individuals

usually: a lower bound for the population size is set

# Cellular Evolutionary Algorithms

also called: “isolation by distance”

Processors are arranged in a rectangular grid usually in the shape of a torus in order to avoid boundary effects  
Selection and crossover are restricted to adjacent processors (to those connected by an edge), mutation to single processors  
Example: each processor is responsible for one chromosome

- **Selection:** processor chooses the best chromosome of the (four) processors adjacent to it (or one of these chromosomes randomly based on their fitness)
- **Crossover:** processor then performs crossover of the selected chromosome with its own and may also mutate the chromosome (The better child resulting from such a crossover replaces the chromosome of the processor)

construction of groups of adjacent processors that maintain similar chromosomes

mitigates the usually destructive effect of the crossover

# Approach of Mühlenbein

## Combination of EAs and hill climbing

- every individual is optimized with hill climbing, that is
    - random mutations are applied and kept if they are advantageous
    - otherwise they are retracted
  - hill climbing can be easily parallelized
  - individuals search for a crossover partner in their (local) neighborhood (this requires a distance measure for the individuals — relates to niche techniques, e.g. *power law sharing*)
  - offspring perform local hill climbing
  - individuals of the next generation are selected with a **local elite principle**
- ⇒ best two individuals among the four involved individuals (two parents and two children) replace the parents

# Outline

## 1. Parallelization of evolutionary algorithms

## 2. Random numbers

Generating uniform distributed random numbers

Linear-congruential method

Normal-distributed random numbers

Random numbers for EA

# Random numbers

EAs base in general on random changes of existing solution candidates and selection

all created „Random numbers“  $\neq$  really random

created by deterministic algorithm: **Pseudorandomness**

- attempts to use random behaviour of physical processes  
numbers with worse properties than deterministic methods
- on top: reducability of simulations with deterministic methods

# Historical

aus [Knuth, 1997]

before 1900: Scientist who needed random numbers drew balls from a „well shuffled urn“, rolled dices or dealt cards

1927, L. Tippett published a table with  $\geq 40000$  random numbers

Disadvantages of tables: expensive to create and store  
machines to generate random numbers

mechanic methods: error-prone and no reproducibility

arithmet. operations on computer: John von Neumann (1946)

- calculate square of the last random number and extract middle digits
- e.g. 10-place random numbers and last number be 5772156649
- square is 33317792380594909201
- therefore, next number is 7923805949

## Argument against arithmetic methods

Can we call a generated sequence "random" if every number is completely determined by its predecessor?

*Anyone who considers arithmetic methods of producing random digits is, of course, in a state of sin.* — John von Neumann (1951)

- Answer: sequence is **not** random, but it seems to be!
- deterministic algorithms generate **pseudorandom** sequences
- von Neumann's method has some issues:
  - sequence tends to short periods of repeating numbers
- z.B. bei 4-stelligen Zahlen: 6100, 2100, 4100, 8100, 6100, ...
- following: methods which are advantageous in comparison to the von Neumann method



# Generating uniform distributed random numbers

- Creating a sequence of real numbers (uniform-distributed from 0 to 1)
- fixed (floating point) precision actually leads to generating integers  $X_n$  between 0 and  $m$
- fraction  $U_n = X_n/m$  lies between 0 and 1
- in general:  $m$  is word size of the computer  $w$

# The linear-congruential method

- most popular random number generators are special implementations of [Lehmer, 1951]

$m$ , Modules;  $0 < m$ ,

$a$ , Factor;  $0 \leq a < m$ ,

$c$ , Increment;  $0 \leq c < m$ ,

$X_0$ , Initial value;  $0 \leq X_0 < m$

- favored sequence of random numbers  $\langle X_n \rangle$  by

$$X_{n+1} = (a \cdot X_n + c) \bmod m, \quad n \geq 0$$

- Remainder mod  $m$ : localisation of a ball on a rotating roulette wheel

# The linear-congruential method

- for e.g.  $m = 10$  and  $X_0 = a = c = 7$  yields to

$$\langle X_n \rangle = 7, 6, 9, 0, 7, 6, 9, 0,$$

- so, sequence is not always random for all values of  $m, a, c, X_0$
- such loops: on all sequences with  $X_{n+1} = f(X_n)$
- **periodic** behaviour
- Objective: useful sequences with a fairly long period

## Favourable choice of parameters

- arbitrary initial value  $X_0$
- Module  $m \geq 2^{30}$  or greatest prime number smaller  $w = 2^e$   
whereby  $e = \#$  representable bits

⇒ *sequence* with maximum length

- if  $m$  Power of 2: factor  $a = 2^k + 1$  whereas  $2 \leq k \leq w$

⇒ *period* with maximum length

- Increment  $c$  is subsidiary if  $a$  is well-chosen: but  $c$  may not have a common divisor with  $m$  (e.g.  $c = 1$  or  $c = a$ )

⇒ avoid multiplication by shift and add operations:

$$X_{n+1} = ((2^k + 1) \cdot X_n + 1) \bmod 2^e$$

- generate at most  $m/1000$  numbers

## Further methods

- square method of R. R. Coveyou: let

$$X_0 \bmod 4 = 2, \quad X_{n+1} = X_n(X_n + 1) \bmod 2^e, \quad n \geq 0$$

be

- computable with similar efficiency like linear-congruential method
- Mitchell and Moore (1958) proposed the approach

$$X_n = (X_{n-24} + X_{n-55}) \bmod m, \quad n \geq 55$$

whereas  $m$  is even and  $X_0, \dots, X_{54}$  is chosen arbitrarily

- very efficient to implement using a cyclic list
- period of  $2^{55} - 1 \Rightarrow$  probably best algorithm

# Generating normal-distributed random numbers

Polar method [Box and Muller, 1958]:

let  $U_1, U_2$  be independent and uniform distributed from  $[0, 1]$   
the following random numbers are generated from the same  
distribution  $N(0, 1)$

$$X_1 = \sqrt{-2 \ln U_1} \cos 2\pi U_2, \quad X_2 = \sqrt{-2 \ln U_1} \sin 2\pi U_2$$

- Proof: inverse relations are

$$U_1 = e^{-\frac{(X_1^2 + X_2^2)}{2}}, \quad U_2 = -\frac{1}{2\pi} \arctan \frac{X_2}{X_1}$$

⇒ multivariate density of  $X_1, X_2$  is

$$\begin{aligned} f(X_1, X_2) &= \frac{1}{2\pi} e^{-\frac{(X_1^2 + X_2^2)}{2}} \\ &= \frac{1}{\sqrt{2\pi}} e^{-\frac{X_1^2}{2}} \cdot \frac{1}{\sqrt{2\pi}} e^{-\frac{X_2^2}{2}} = f(X_1) \cdot f(X_2) \end{aligned}$$

# Generating normal-distributed random numbers

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## Algorithm Polar-method

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**Output:** two independent, normal-distributed random numbers  $X_1, X_2$

- 1: **do** {
  - 2:    $U_1, U_2 \leftarrow$  generate two independent random numbers from  $U([0, 1])$
  - 3:    $V_1 \leftarrow 2U_1 - 1$
  - 4:    $V_2 \leftarrow 2U_2 - 1$
  - 5:    $S \leftarrow V_1^2 + V_2^2$
  - 6: } **while**  $S < 1.0$  **and**  $S \neq 0$
  - 7:  $X_1 \leftarrow V_1 \sqrt{\frac{-2 \ln S}{S}}$
  - 8:  $X_2 \leftarrow V_2 \sqrt{\frac{-2 \ln S}{S}}$
  - 9: **return**  $X_1, X_2$
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# Random numbers for EA




- reasonable for an object-oriented approach:  
one random number generator per individual
- disadvantage: consequences due to randomness cannot be estimated

⇒ pro optimization method: only one random number generator

- a clearly defined initial value is advisable ⇒ experiments are reproducible
- system time or last created number as seed should be avoided



## Additional Literature I

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