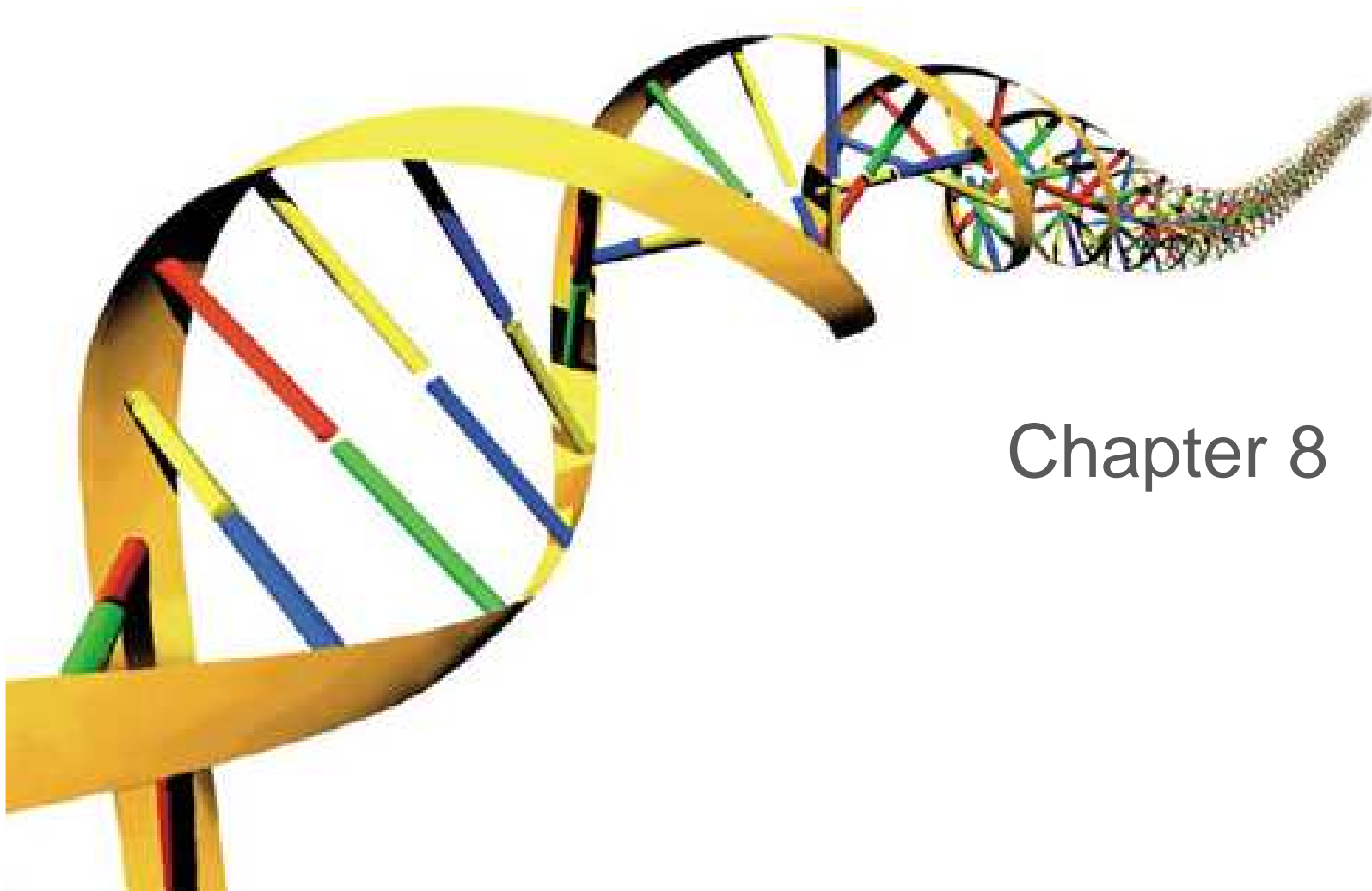


# Evolutionary Computing



## Chapter 8

# Chapter 8: Parameter Control

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- Motivation
- Parameter setting
  - Tuning
  - Control
- Examples
- Where to apply parameter control
- How to apply parameter control

# Motivation (1/2)

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An EA has many strategy parameters, e.g.

- mutation operator and mutation rate
- crossover operator and crossover rate
- selection mechanism and selective pressure (e.g. tournament size)
- population size

Good parameter values facilitate good performance

Q1 How to find good parameter values ?

## Motivation (2/2)

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EA parameters are rigid (constant during a run)

BUT

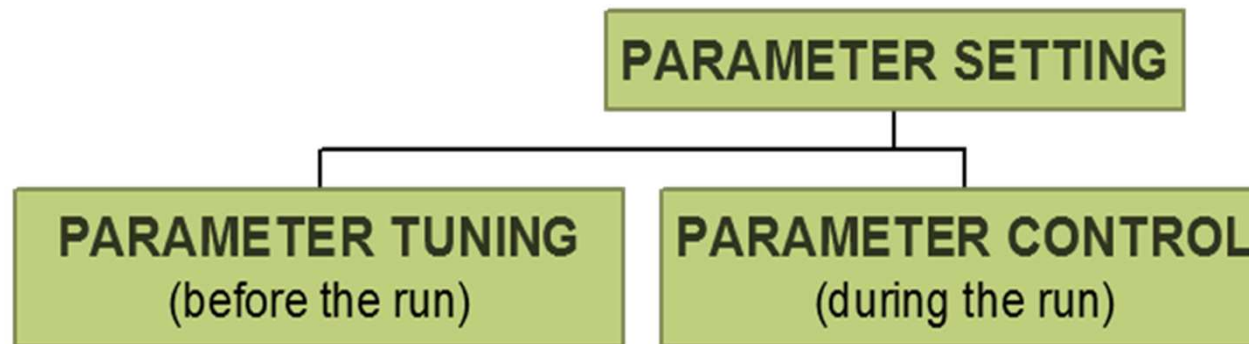
an EA is a dynamic, adaptive process

THUS

optimal parameter values may vary during a run

Q2: How to vary parameter values?

# Parameter Setting



# Parameter Settings: Tuning

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**Parameter tuning:** the traditional way of testing and comparing different values **before the “real” run**

## Problems:

- users mistakes in settings can be sources of errors or sub-optimal performance
- parameters interact: exhaustive search is not practicable
- costs much time even with “smart” tuning
- good values may become bad during the run

# Parameter Settings: Control

**Parameter control:** setting values on-line, during the actual run, e.g.

- predetermined time-varying schedule  $p = p(t)$
- using feedback from the search process
- encoding parameters in chromosomes and rely on selection

Problems:

- finding optimal  $p$  is hard, finding optimal  $p(t)$  is harder
- still user-defined feedback mechanism, how to ``optimize"?
- when would natural selection work for strategy parameters?

# Examples:

## Varying mutation step size

### Task to solve:

- $\min f(x_1, \dots, x_n)$
- $L_i \leq x_i \leq U_i$  for  $i = 1, \dots, n$  bounds
- $g_i(x) \leq 0$  for  $i = 1, \dots, q$  inequality constraints
- $h_i(x) = 0$  for  $i = q+1, \dots, m$  equality constraints

### Algorithm:

- EA with real-valued representation  $(x_1, \dots, x_n)$
- arithmetic averaging crossover
- Gaussian mutation:  $x'_i = x_i + N(0, \sigma)$   
standard deviation  $\sigma$  is called mutation step size



# Examples:

## Varying mutation step size, option 1

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Replace the constant  $\sigma$  by a function  $\sigma(t)$

$$\sigma(t) = 1 - 0.9 \times \frac{t}{T}$$

$0 \leq t \leq T$  is the current generation number

- Features:
  - changes in  $\sigma$  are independent from the search progress
  - strong user control of  $\sigma$  by the above formula
  - $\sigma$  is fully predictable
  - a given  $\sigma$  acts on all individuals of the population

# Examples:

## Varying mutation step size, option 2

Replace the constant  $\sigma$  by a function  $\sigma(t)$  updated after every  $n$  steps by the 1/5 success rule:

$$\sigma(t) = \begin{cases} \sigma(t-n)/c & \text{if } p_s > 0.2 \\ \sigma(t-n) \cdot c & \text{if } p_s < 0.2 \\ \sigma(t-n) & \text{otherwise} \end{cases}$$

- Features:
  - changes in  $\sigma$  are based on feedback from the search progress
  - some user control of  $\sigma$  by the above formula
  - $\sigma$  is not predictable
  - a given  $\sigma$  acts on all individuals of the population

# Examples:

## Varying mutation step size, option 3

- Assign a personal  $\sigma$  to each individual
- Incorporate this  $\sigma$  into the chromosome:  $(x_1, \dots, x_n, \sigma)$
- Apply variation operators to  $x_i$ 's and  $\sigma$

$$\sigma' = \sigma \times e^{N(0, \sigma)}$$

$$x_i' = x_i + N(0, \sigma')$$

- Features:
  - changes in  $\sigma$  are results of natural selection
  - (almost) no user control of  $\sigma$
  - $\sigma$  is not predictable
  - a given  $\sigma$  acts on one individual

# Examples:

## Varying mutation step size, option 4

Assign a personal  $\sigma$  to each variable in each individual

Incorporate  $\sigma$ 's into the chromosomes:  $(x_1, \dots, x_n, \sigma_1, \dots, \sigma_n)$

Apply variation operators to  $x_i$ 's and  $\sigma_i$ 's

$$\sigma_i' = \sigma_i \times e^{N(0, \tau)}$$

$$x_i' = x_i + N(0, \sigma_i')$$

- Features:
  - changes in  $\sigma_i$  are results of natural selection
  - (almost) no user control of  $\sigma_i$
  - $\sigma_i$  is not predictable
  - a given  $\sigma_i$  acts on one gene of one individual

# Examples: Varying penalties

## Constraints

- $g_i(x) \leq 0$  for  $i = 1, \dots, q$  inequality constraints
- $h_i(x) = 0$  for  $i = q+1, \dots, m$  equality constraints

are handled by penalties:

$$eval(x) = f(x) + W \times penalty(x)$$

where

$$penalty(x) = \sum_{j=1}^m \begin{cases} 1 & \text{for violated constraint} \\ 0 & \text{for satisfied constraint} \end{cases}$$

# Examples:

## Varying penalties, option 1

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Replace the constant  $W$  by a function  $W(t)$

$$W(t) = (C \times t)^\alpha$$

$0 \leq t \leq T$  is the current generation number

- Features:
  - changes in  $W$  independent from the search progress
  - strong user control of  $W$  by the above formula
  - $W$  is fully predictable
  - a given  $W$  acts on all individuals of the population

# Examples:

## Varying penalties, option 2

Replace the constant  $W$  by  $W(t)$  updated in each generation

$$W(t+1) = \begin{cases} \beta \times W(t) & \text{if last } k \text{ champions all feasible} \\ \gamma \times W(t) & \text{if last } k \text{ champions all infeasible} \\ W(t) & \text{otherwise} \end{cases}$$

$\beta < 1, \gamma > 1, \beta \times \gamma \neq 1$  champion: best of its generation

- Features:
  - changes in  $W$  are based on feedback from the search progress
  - some user control of  $W$  by the above formula
  - $W$  is not predictable
  - a given  $W$  acts on all individuals of the population

# Examples:

## Varying penalties, option 3

Assign a personal  $W$  to each individual

Incorporate this  $W$  into the chromosome:  $(x_1, \dots, x_n, W)$

Apply variation operators to  $x_i$ 's and  $W$

Alert:

$$eval((x, W)) = f(x) + W \times penalty(x)$$

while for mutation step sizes we had

$$eval((x, \sigma)) = f(x)$$

this option is thus sensitive “cheating”  $\Rightarrow$  makes no sense



# Examples:

## Lessons learned (1/2)

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Various forms of parameter control can be distinguished by:

- primary features:
  - **what** component of the EA is changed
  - **how** the change is made
- secondary features:
  - **evidence/data** backing up changes
  - **level/scope** of change

# Examples: Lessons learned (2/2)

Various forms of parameter control can be distinguished by:

	$\sigma(t) = 1 - 0.9^{t/T}$	$\sigma' = \sigma/c$ , if $r > 1/5$ ...	$(x_1, \dots, x_n, \sigma)$	$(x_1, \dots, x_n, \sigma_1, \dots, \sigma_n)$	$W(t) = (C^*t)^{\alpha}$	$W' = \beta * W$ , if $b_i \in F$	$(x_1, \dots, x_n, W)$
<b>What</b>	Step size	Step size	Step size	Step size	Penalty weight	Penalty weight	Penalty weight
<b>How</b>	Deterministic	Adaptive	Self-adaptive	Self-adaptive	Deterministic	Adaptive	Self-adaptive
<b>Evidence</b>	Time	Successful mutations rate	(Fitness)	(Fitness)	Time	Constraint satisfaction history	(Fitness)
<b>Scope</b>	Population	Population	Individual	Gene	Population	Population	Individual

# Where to apply parameter control

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Practically any EA component can be parameterized and thus controlled on-the-fly:

- representation
- evaluation function
- variation operators
- selection operator (parent or mating selection)
- replacement operator (survival or environmental selection)
- population (size, topology)

# How to apply parameter control

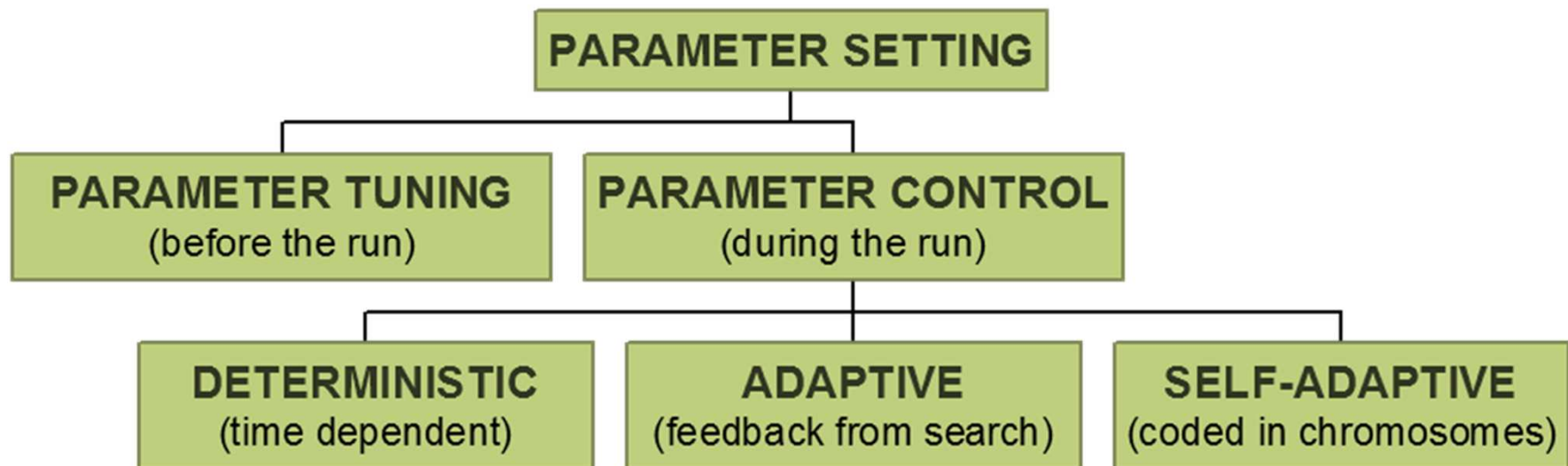
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Three major types of parameter control:

- **deterministic**: some rule modifies strategy parameter without feedback from the search (based on some counter)
- **adaptive**: feedback rule based on some measure monitoring search progress
- **self-adaptive**: parameter values evolve along with solutions; encoded onto chromosomes they undergo variation and selection

# How to apply parameter control

## Global taxonomy



# Evidence: Informing the change (1/2)

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The parameter changes may be based on:

- **time or nr. of evaluations** (deterministic control)
- **population statistics** (adaptive control)
  - progress made
  - population diversity
  - gene distribution, etc.
- **relative fitness** of individuals created with given values (adaptive or self-adaptive control)

# Evidence: Informing the change (2/2)

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- **Absolute evidence**: predefined event triggers change, e.g. increase  $p_m$  by 10% if population diversity falls under threshold  $x$
- Direction and magnitude of change is fixed
- **Relative evidence**: compare values through solutions created with them, e.g. increase  $p_m$  if top quality offspring came by high mutation rates
- Direction and magnitude of change is not fixed

# Evidence: Refined taxonomy

- Combinations of types and evidences
  - Possible: +
  - Impossible: -

	Deterministic	Adaptive	Self-adaptive
Absolute	+	+	-
Relative	-	+	+



# Scope/level

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The parameter may take effect on different levels:

- environment (fitness function)
- population
- individual
- sub-individual

Note: given component (parameter) determines possibilities

Thus: scope/level is a derived or secondary feature in the classification scheme

# Evaluation/Summary

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- Parameter control offers the possibility to use **appropriate values in various stages of the search**
- Adaptive and self-adaptive parameter control
  - offer users **“liberation” from parameter tuning**
  - delegate **parameter setting task to the evolutionary process**
  - the latter implies a double task for an EA: problem solving + **self-calibrating (overhead)**