

Parameter control

Chapter 8

A.E. Eiben and J.E. Smith, Introduction to Evolutionary Computing
Parameter Control in EAs

Motivation 1

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 ?

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Motivation 2

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?

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Parameter tuning

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
- costs much time
- parameters interact: exhaustive search is not practicable
- good values may become bad during the run

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Parameter 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 natural 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?

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Example

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 σ is called mutation step size

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Varying mutation step size: option1

Replace the constant σ 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 σ are independent from the search progress
- strong user control of σ by the above formula
- σ is fully predictable
- a given σ acts on all individuals of the population

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Varying mutation step size: option2

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

$$\sigma(t) = \begin{cases} \sigma(t-n)/c & \text{if } p_s > 1/5 \\ \sigma(t-n) \cdot c & \text{if } p_s < 1/5 \\ \sigma(t-n) & \text{otherwise} \end{cases}$$

Features:

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

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Varying mutation step size: option3

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

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

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

Features:

changes in σ are results of natural selection
(almost) no user control of σ
 σ is not predictable
a given σ acts on one individual

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Varying mutation step size: option4

Assign a personal σ to each variable in each individual
Incorporate σ 's into the chromosomes: $(x_1, \dots, x_n, \sigma_1, \dots, \sigma_n)$
Apply variation operators to x_i 's and σ_i 's

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

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

Features:

changes in σ_i are results of natural selection
(almost) no user control of σ_i
 σ_i is not predictable
a given σ_i acts on 1 gene of one individual

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Example cont'd

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

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

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Varying penalty: option 1

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 are 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

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Varying penalty: 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

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Varying penalty: 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:

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

while for mutation step sizes we had

$$\text{eval}((x, \sigma)) = f(x)$$

this option is thus sensitive "cheating" \Rightarrow makes no sense

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Lessons learned from examples

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

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What

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)

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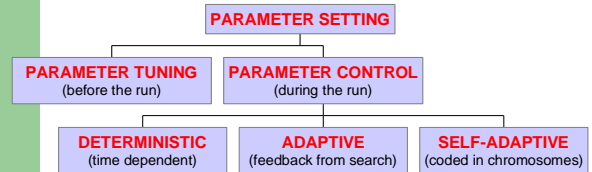
How

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

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Global taxonomy



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Evidence informing the change

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)

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Evidence informing the change

- **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 mut. Rates
- Direction and magnitude of change is not fixed

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Scope/level

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

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Refined taxonomy

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

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

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Evaluation / Summary

- 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)**
- Robustness, insensitivity of EA for variations assumed
 - If no. of parameters is increased by using (self)adaptation
 - For the "meta-parameters" introduced in methods

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