

Computational Intelligence

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Introduction Evolutionary Algorithms Initialization and Selection Variation **EA Parameters** Typical EAs EA Design

Important Parameters of EAs (1)

- **dimension n of search space**
 - no parameter of EA, but given by the problem
 - measures the size of the search space: $\{0,1\}^n, \mathbb{R}^n, S_n$
 - plays the same role as input length in classical runtime analysis
 - other parameters are often chosen dependent on n (e.g. mutation probability $p_m = 1/n$)
- **population size μ**
 - obviously $\mu = n^{O(1)}$
 - often $\mu = \Theta(n)$ or $\mu = \Theta(\sqrt{n})$
 - $\mu = O(1)$ or even $\mu = 1$ are not unusual
- **number of offspring λ**
 - obviously $\lambda = n^{O(1)}$
 - often $\lambda = 1$
 - $\lambda = \mu$ or $\lambda \gg \mu$ not unusual
 - selection method influences reasonable choice of λ

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Important Parameters of EAs (2)

- **crossover probability p_c**
 - in general $p_c \in [0;1]$ arbitrary
 - often $p_c \in [1/2; 4/5]$ constant
- **probability of applying mutation**
 - don't confuse with mutation probability!
 - we will always use 1
 - Remark
 - $p_m = 1/n \Rightarrow \text{Prob (no mutation)} = (1 - 1/n)^n \approx 1/e$

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Methods for parameter control

- **static parameter control**
 - parameter values constant during the whole run
 - often used
 - + simple
 - maybe it's better to vary the parameter value during the run?!
- **dynamic parameter control**
 - parameter values change during the run according to some time-dependent scheme
 - + more flexible than static approach
 - cannot deal with non-time-dependent changes
 - unusual for EAs
- **adaptive parameter control**
 - parameter values can change dependently on every individual and any random experiment
 - + very flexible
 - hard to analyze
 - computationally expensive
 - often used for EAs

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Self-adaptation

Idea good parameter values evolve together with good individuals

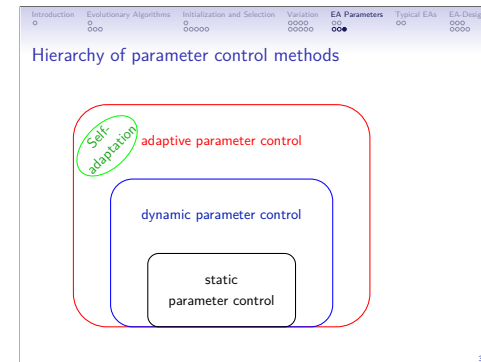
implementation code parameter values together with individual

formally $S \times Q$ instead of S
unchanged $f: S \rightarrow R$

e. g. for mutation probability

- every individual has its own mutation probability
- first vary the mutation probability
- then mutate with varied mutation probability
- afterwards normal selection
- important **don't swap steps**

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Evolutionary Algorithms: Historical Notes

Lecture 11

Idea emerged independently several times: about late 1950s / early 1960s.
Three branches / “schools“ still active today.

• Evolutionary Programming (EP):

Pioneers: Lawrence Fogel, Alvin Owen, Michael Walsh (New York, USA).

Original goal: Generate intelligent behavior through simulated evolution.

Approach: Evolution of finite state machines predicting symbols.

Later (~1990s) specialized to optimization in \mathbb{R}^n by David B. Fogel.

• Genetic Algorithms (GA):

Pioneer: John Holland (Ann Arbor, MI, USA).

Original goal: Analysis of adaptive behavior.

Approach: Viewing evolution as adaptation. Simulated evolution of bit strings.

Applied to optimization tasks by PhD students (Kenneth de Jong, 1975; et al.).

• Evolution Strategies (ES):

Pioneers: Ingo Rechenberg, Hans-Paul Schwefel, Peter Bienert (Berlin, Germany).

Original goal: Optimization of complex systems.

Approach: Viewing variation/selection as improvement strategy. First in \mathbb{Z}^n , then \mathbb{R}^n .

Evolutionary Algorithms: Historical Notes

Lecture 11

“Offspring“ from GA branch:

• Genetic Programming (GP):

Pioneers: Michael Lynn Cramer 1985, then: John Koza (Stanford, USA).

Original goal: Evolve programs (parse trees) that must accomplish certain task.

Approach: GA mechanism transferred to parse trees.

Later: Programs as successive statements → Linear GP (e.g. Wolfgang Banzhaf)

Already beginning early 1990s:

Borders between EP, GA, ES, GP begin to blurr ...

⇒ common term **Evolutionary Algorithm** embracing all kind of approaches

⇒ broadly accepted name for the field: **Evolutionary Computation**

scientific journals: *Evolutionary Computation* (MIT Press) since 1993,

IEEE Transactions on Evolutionary Computation since 1997,

several more specialized journals started since then.

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Design of EAs

Idea Methodology to apply standard EAs

Goal standard EAs do not have to be changed

Requirement problem is given as $g: A \rightarrow B$
 g has to be maximized (or minimized)
 A arbitrary set, B partially ordered

EA operates on search space S
 'maximizes' fitness $f: S \rightarrow \mathbb{R}$

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Definition of mappings

Fitness $f := h_2 \circ g \circ h_1$
 h_1 is genotype-phenotype-mapping.

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Genotype-Phenotype-Mapping $\mathbb{B}^n \rightarrow [L, R] \subset \mathbb{R}$

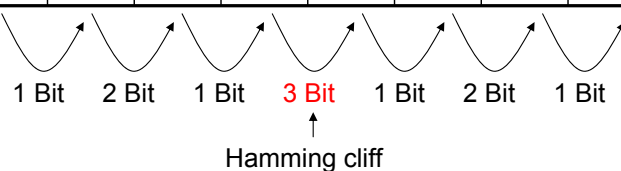
- Standard encoding for $b \in \mathbb{B}^n$

$$x = L + \frac{R - L}{2^n - 1} \sum_{i=0}^{n-1} b_{n-i} 2^i$$

→ Problem: *hamming cliffs*

000	001	010	011	100	101	110	111
0	1	2	3	4	5	6	7

$L = 0, R = 7$
 $n = 3$



Genotype-Phenotype-Mapping $\mathbb{B}^n \rightarrow [L, R] \subset \mathbb{R}$

- Gray encoding for $b \in \mathbb{B}^n$

Let $a \in \mathbb{B}^n$ standard encoded. Then $b_i = \begin{cases} a_i, & \text{if } i = 1 \\ a_{i-1} \oplus a_i, & \text{if } i > 1 \end{cases}$ $\oplus = \text{XOR}$

000	001	011	010	110	111	101	100
0	1	2	3	4	5	6	7

← genotype
 ← phenotype

OK, no hamming cliffs any longer ...

⇒ small changes in phenotype „lead to“ small changes in genotype

since we consider evolution in terms of Darwin (not Lamarck):

⇒ small changes in genotype lead to small changes in phenotype!

but: 1-Bit-change: $000 \rightarrow 100 \Rightarrow \ominus$

Genotype-Phenotype-Mapping $\mathbb{B}^n \rightarrow \mathbb{P}^n$ (example only)

- e.g. standard encoding for $b \in \mathbb{B}^n$

individual:

010	101	111	000	110	001	101	100	← genotype
0	1	2	3	4	5	6	7	← index

consider index and associated genotype entry as unit / record / struct;
 sort units with respect to genotype value, old indices yield permutation:

000	001	010	100	101	101	110	111	← genotype
3	5	0	7	1	6	4	2	← old index

= permutation

Requirements on h_1 and h_2

obvious requirements

- h_1 and h_2 can be computed efficiently
- h_2 suits g , i.e. good points in B are mapped to good points in \mathbb{R}
- h_1 maps on many (all) important points of A
- Optima of f correspond to optima of g

Caution requirements can be hard to achieve in practice
 for non-obvious requirements a metric is important

Definition

Mapping $d: M \times M \rightarrow \mathbb{R}_0^+$ is a metric on the set M \Leftrightarrow

- $\forall x, y \in M: x \neq y \Leftrightarrow d(x, y) > 0$ (positivity)
- $\forall x, y \in M: d(x, y) = d(y, x)$ (symmetry)
- $\forall x, y, z \in M: d(x, y) + d(y, z) \geq d(x, z)$ (triangle inequality)

Metric-based EAs

Assumption Metric d_A on A known
 (d_A reflects application knowledge)

Requirement metric d_S is known

if h_1 injective, $d_S(x, x') := d_A(h_1(x), h_1(x'))$ is metric

Requirement monotonicity

$$\forall x, x', x'' \in S: \begin{aligned} d_S(x, x') &\leq d_S(x, x'') \\ \Rightarrow d_A(h_1(x), h_1(x')) &\leq d_A(h_1(x), h_1(x'')) \end{aligned}$$

Variation as randomized mapping

now Design-rules for variation operators

hence Formalize variation operators as randomized mappings

$r: X \rightarrow Y$ randomized mapping
 $\Leftrightarrow r(x) \in Y$ depends on $x \in X$ and random experiment

formally probability space (Ω, p)

$$r: X \times \Omega \rightarrow Y$$

$$\text{Prob}(r(x) = y) = \sum_{\omega \in \Omega: r(x, \omega) = y} p(\omega)$$

Example 1-bit mutation

$$\Omega := \{1, 2, \dots, n\}, \forall i \in \Omega: p(i) = 1/n$$

1-bit mutation is randomized mapping $m: \{0, 1\}^n \rightarrow \{0, 1\}^n$
 where $m(x, i) := x \oplus 0^{i-1}10^{n-i}$

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Design-rules for mutation

favor small changes

$$\forall x, x', x'' \in S: \quad d_S(x, x') < d_S(x, x'') \\ \Rightarrow \text{Prob}(m(x) = x') > \text{Prob}(m(x) = x'')$$

no bias

$$\forall x, x', x'' \in S: \quad d_S(x, x') = d_S(x, x'') \\ \Rightarrow \text{Prob}(m(x) = x') = \text{Prob}(m(x) = x'')$$

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Design-rules for crossover

offspring similar to parents

$$\forall x, x', x'' \in S: \quad \text{Prob}(c(x, x') = x'') > 0 \\ \Rightarrow \max\{d_S(x, x''), d_S(x', x'')\} \leq d_S(x, x')$$

no bias

$$\forall x, x' \in S: \forall \alpha \in \mathbb{R}_0^+: \\ \text{Prob}(d_S(x, c(x, x')) = \alpha) = \text{Prob}(d_S(x', c(x, x')) = \alpha)$$

Any EA that fulfills these four design-rules is called a metric-based EA (MBEA).

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