

Computational Intelligence

Winter Term 2021/22

Prof. Dr. Günter Rudolph

Lehrstuhl für Algorithm Engineering (LS 11)

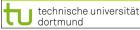
Fakultät für Informatik

TU Dortmund

Plan for Today

Lecture 07

- Design of Evolutionary Algorithms
 - Case Study: Integer Search Space
 - **Towards CMA-ES**



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Design of Evolutionary Algorithms

Lecture 07

- every recombination results

- mutation of z may then lead

to any $z^* \in \mathbb{Z}^n$ with positive probability in one step

in some $z \in \mathbb{Z}^n$

ad 2) design guidelines for variation operators in practice

integer search space $X = \mathbb{Z}^n$

- a) reachability
- b) unbiasedness
- c) control
- ad a) support of mutation should be \mathbb{Z}^n
- ad b) need maximum entropy distribution over support \mathbb{Z}^n
- ad c) control variability by parameter
 - → formulate as constraint of maximum entropy distribution

Design of Evolutionary Algorithms

Lecture 07

ad 2) design guidelines for variation operators in practice

 $X = \mathbb{Z}^n$

task: find (symmetric) maximum entropy distribution over \mathbb{Z} with $\mathbb{E}[|Z|] = \theta > 0$

⇒ need *analytic* solution of an ∞-dimensional, nonlinear optimization problem with constraints!

$$H(p) = -\sum_{k=-\infty}^{\infty} p_k \log p_k \quad \longrightarrow \quad \max!$$

$$p_k =$$

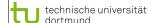
$$= p_{-k} \quad \forall k \in \mathbb{Z}$$
 , (symme

$$\sum_{k=0}^{\infty} p_k = 1$$
 , (normalization

$$p_k = p_{-k} \quad \forall k \in \mathbb{Z}$$
 , (symmetry w.r.t. 0)
$$\sum_{k=-\infty}^{\infty} p_k = 1 \; , \qquad \qquad \text{(normalization)}$$

$$\sum_{k=-\infty}^{\infty} |k| \, p_k = \theta \qquad \qquad \text{(control "spread")}$$

$$p_k \geq 0 \quad \forall k \in \mathbb{Z}$$
 . (nonnegativity)



Design of Evolutionary Algorithms

Lecture 07

result:

a random variable Z with support $\ensuremath{\mathbb{Z}}$ and probability distribution

$$p_k := P\{Z = k\} = \frac{q}{2-q} (1-q)^{|k|}, k \in \mathbb{Z}, q \in (0,1)$$

symmetric w.r.t. 0, unimodal, spread manageable by q and has max. entropy

generation of pseudo random numbers:

$$Z = G_1 - G_2$$

where

$$U_i \sim U(0,1) \quad \Rightarrow \quad G_i = \left| \frac{\log(1 - U_i)}{\log(1 - q)} \right| \quad , \quad i = 1, 2.$$

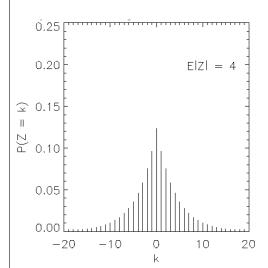
stochastic independent!

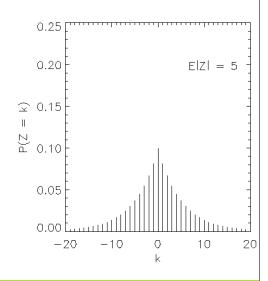


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Design of Evolutionary Algorithms Lecture 07

probability distributions for different mean step sizes $E|Z| = \theta$





Design of Evolutionary Algorithms Lecture 07 probability distributions for different mean step sizes $E|Z| = \theta$ 0.20 0.20 F[7] = 2E|Z| = 3♀ 0.15 0.15 Z) 0.10 0.05 0.05 -20-1010 -20

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Design of Evolutionary Algorithms

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How to control the spread?

We must be able to adapt $q \in (0,1)$ for generating Z with variable $E|Z| = \theta$! self-adaptation of q in open interval (0,1)?

 \longrightarrow make mean step size E[|Z|] adjustable!

$$E[|Z|] = \sum_{k=-\infty}^{\infty} |k| p_k = \theta = \frac{2(1-q)}{q(2-q)} \Leftrightarrow q = 1 - \frac{\theta}{(1+\theta^2)^{1/2} + 1}$$

$$\in \mathbb{R}_+ \qquad \qquad \in (0,1)$$

 $\rightarrow \theta$ adjustable by mutative self adaptation

 $\rightarrow \text{get q from } \theta$

like mutative step size size control of σ in EA with search space \mathbb{R}^n !

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Lecture 07

Mutative Step Size Control

Individual $(\mathbf{x}, \boldsymbol{\theta}) \in \mathbb{Z}^n \times \mathbb{R}_+$

First, mutate step size

 $\theta_{t+1} = \theta_t \cdot L$

Second, mutate parent $Y = x + \theta_{t+1} \cdot Z$

Often: assure minimal step size ≥ 1

 $\theta_{t+1} = \max\{1, \theta_t \cdot L\}$

→ invented: Schwefel (1977) for real variables

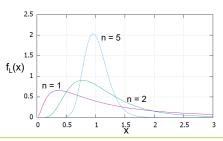
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→ transferred: Rudolph (1994) for integer variables

where L = $\exp(N)$ with N ~ N(0, 1/n)

log-normal distributed

 $P\{L > c\} = P\{L < 1/c\} \text{ for } c \ge 1$



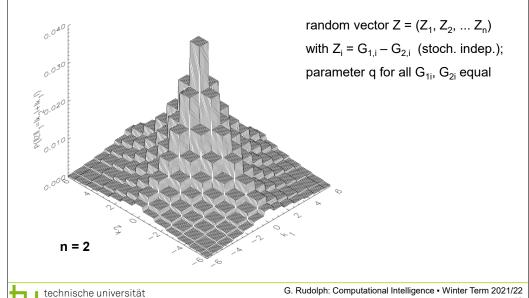
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n - dimensional generalization



Lecture 07

n - dimensional generalization

$$P\{Z_i = k\} = \frac{q}{2-q} (1-q)^{|k|}$$

$$P\{Z_1 = k_1, Z_2 = k_2, \dots, Z_n = k_n\} = \prod_{i=1}^n P\{Z_i = k_i\} =$$

$$\left(\frac{q}{2-q}\right)^n \prod_{i=1}^n (1-q)^{|k_i|} = \left(\frac{q}{2-q}\right)^n (1-q)^{\sum_{i=1}^n |k_i|}$$

$$= \left(\frac{q}{2-q}\right)^n (1-q)^{\|k\|_1}.$$

- \Rightarrow n-dimensional distribution is symmetric w.r.t. ℓ_1 norm!
- ⇒ all random vectors with same step length have same probability!

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How to control E[$||Z||_1$]?

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$$E[\|Z\|_1] = E\left[\sum_{i=1}^n |Z_i|\right] = \sum_{i=1}^n E[|Z_i|] = n \cdot E[|Z_1|]$$
 by def. linearity of E[·] identical distributions for Z_i

$$n \cdot E[|Z_1|] = n \cdot \frac{2(1-q)}{q(2-q)} \quad \Leftrightarrow \quad q = 1 - \frac{\theta/n}{(1+(\theta/n)^2)^{1/2}+1}$$

$$= \theta \qquad \qquad \text{self-adaptation}$$

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Algorithm:

individual $(x,\theta) \in \mathbb{Z}^n \times \mathbb{R}_+$

: $\theta^{(t+1)} = \theta^{(t)} \cdot \exp(N)$. $N \sim N(0, 1/n)$. mutation

if $\theta^{(t+1)} < 1$ then $\theta_{t+1} = 1$

calculate new q for G_i from θ_{t+1}

 $\forall j = 1, \dots, n : X_j^{(t+1)} = X_j^{(t)} + (G_{1,j} - G_{2,j})$

recombination: discrete (uniform crossover)

: (μ, λ) -selection selection

(Rudolph, PPSN 1994)



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Excursion: Maximum Entropy Distributions

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ad 2) design guidelines for variation operators in practice

continuous search space $X = \mathbb{R}^n$

a) reachability → mutation distribution with unbounded support

b) unbiasedness → mutation distribution with maximum entropy

→ mutation distribution with parameters c) control

⇒ leads to CMA-ES!

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Covariance Matrix **A**daptation

Design of Evolutionary Algorithms

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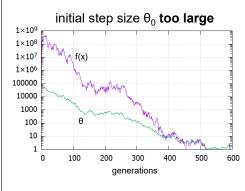
Example: $(1, \lambda)$ -EA with $\lambda = 10$; $f(x) = x'x \rightarrow min!$; n = 10

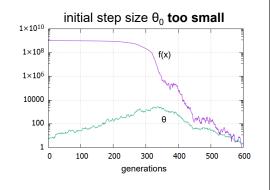
$$\mathbf{X}^{(0)} \in [100, 101]^n \cap \mathbb{Z}^n$$

$$\mathbf{X}^{(0)} \in [10000, 10100]^n \cap \mathbb{Z}^n$$

$$\theta_0 = 50~000$$

$$\theta_0 = 5$$





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Towards CMA-ES

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mutation: Y = X + Z

Z ~ N(0, C) multinormal distribution

maximum entropy distribution for support Rn, given expectation vector and covariance matrix

how should we choose covariance matrix C?

unless we have not learned something about the problem during search

⇒ don't prefer any direction!

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 \Rightarrow covariance matrix C = I_n (unit matrix)

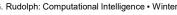






 $C = diag(s_1,...,s_n)$

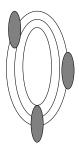
C orthogonal



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claim: mutations should be aligned to isolines of problem (Schwefel 1981)



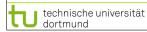
if true then covariance matrix should be inverse of Hessian matrix!

$$\Rightarrow$$
 assume f(x) $\approx \frac{1}{2}$ x'Ax + b'x + c \Rightarrow H = A

$$Z \sim N(0, C)$$
 with density
$$f_Z(x) = \frac{1}{(2\pi)^{n/2} |C|^{1/2}} \exp\left(-\frac{1}{2}x'C^{-1}x\right)$$

since then many proposals how to adapt the covariance matrix

 \Rightarrow extreme case: use n+1 pairs (x, f(x)), apply multiple linear regression to obtain estimators for A, b, c invert estimated matrix A! OK, **but**: O(n⁶)! (Rudolph 1992)



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 $Z = rQu, A = B'B, B = Q^{-1}$

$$f(x + rQu) = \frac{1}{2}(x + rQu)'A(x + rQu) + b'(x + rQu) + c$$

$$= \frac{1}{2}(x'Ax + 2rx'AQu + r^2u'Q'AQu) + b'x + rb'Qu + c$$

$$= f(x) + rx'AQu + rb'Qu + \frac{1}{2}r^2u'Q'AQu$$

$$= f(x) + r(Ax + b + \frac{r}{2}AQu)'Qu$$

$$= f(x) + r(\nabla f(x) + \frac{r}{2}AQu)'Qu$$

$$= f(x) + r\nabla f(x)'Qu + \frac{r^2}{2}u'Q'AQu$$

$$= f(x) + r\nabla f(x)'Qu + \frac{r^2}{2} \rightarrow \min!$$

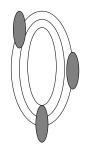
if Qu were deterministic ...

 \Rightarrow set Qu = $-\nabla f(x)$ (direction of steepest descent)

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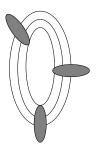
doubts: are equi-aligned isolines really optimal?



principal axis

should point into negative gradient direction!

(proof next slide)



most (effective) algorithms behave like this:

run roughly into negative gradient direction, sooner or later we approach longest main principal axis of Hessian,

now negative gradient direction coincidences with direction to optimum, which is parallel to longest main principal axis of Hessian, which is parallel to the longest main principal axis of the inverse covariance matrix

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(Schwefel OK in this situation)

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Apart from (inefficient) regression, how can we get matrix elements of Q?

 $C^{(k+1)}$ = update($C^{(k)}$, Population(k)) ⇒ iteratively:

 $C^{(k)}$ must be positive definite (p.d.) and symmetric for all $k \ge 0$, basic constraint:

otherwise Cholesky decomposition impossible: C = Q'Q

Lemma

Let A and B be quadratic matrices and α , $\beta > 0$.

- a) A, B symmetric $\Rightarrow \alpha A + \beta B$ symmetric.
- b) A positive definite and B positive semidefinite $\Rightarrow \alpha A + \beta B$ positive definite

ad a) C = α A + β B symmetric, since $c_{ii} = \alpha a_{ii} + \beta b_{ii} = \alpha a_{ii} + \beta b_{ii} = c_{ii}$

ad b)
$$\forall x \in \mathbb{R}^n \setminus \{0\}$$
: $x'(\alpha A + \beta B) x = \alpha x'Ax + \beta x'Bx > 0$

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Theorem

A quadratic matrix $C^{(k)}$ is symmetric and positive definite for all $k \ge 0$, if it is built via the iterative formula $C^{(k+1)} = \alpha_k C^{(k)} + \beta_k v_k v_k^t$ where $C^{(0)} = I_{n_k} v_k \ne 0$, $\alpha_k > 0$ and liminf $\beta_k > 0$.

Proof:

If $v \neq 0$, then matrix V = vv' is symmetric and positive semidefinite, since

- as per definition of the dyadic product $v_{ij} = v_i \cdot v_j = v_i \cdot v_j = v_{ij}$ for all i, j and
- for all $x \in \mathbb{R}^n$: $x'(vv') x = (x'v) \cdot (v'x) = (x'v)^2 \ge 0$.

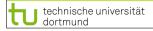
Thus, the sequence of matrices $v_k v'_k$ is symmetric and p.s.d. for $k \ge 0$.

Owing to the previous lemma matrix C(k+1) is symmetric and p.d., if

 $C^{(k)}$ is symmetric as well as p.d. and matrix $v_k v_k'$ is symmetric and p.s.d.

Since $C^{(0)} = I_n$ symmetric and p.d. it follows that $C^{(1)}$ is symmetric and p.d.

Repetition of these arguments leads to the statement of the theorem.



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CMA-ES

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State-of-the-art: CMA-EA (currently many variants)

→ many successful applications in practice

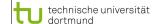
C, C++, Java Fortran, Python, Matlab, R, Scilab

available in WWW:

- http://cma.gforge.inria.fr/cmaes_sourcecode_page.html
- http://image.diku.dk/shark/ (EAlib, C++)
- ...

advice:

before designing your own new method or grabbing another method with some fancy name ... try CMA-ES – it is available in most software libraries and often does the job!



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CMA-ES Lecture 07

Idea: Don't estimate matrix C in each iteration! Instead, approximate <u>iteratively!</u>

(Hansen, Ostermeier et al. 1996ff.)

→ Covariance Matrix Adaptation Evolutionary Algorithm (CMA-EA)

Set initial covariance matrix to $C^{(0)} = I_n$

$$C^{(t+1)} = (1-\eta) C^{(t)} + \eta \sum_{i=1}^{\mu} w_i (x_{i:\lambda} - m^{(t)}) (x_{i:\lambda} - m^{(t)})$$

$$\eta$$
: "learning rate" \in (0,1)

$$\textbf{w}_{\text{i}}$$
 : weights; mostly 1/ $\!\mu$

$$m = \frac{1}{\mu} \sum_{i=1}^{\mu} x_{i:\lambda}$$
 mean of all selected parents

complexity: $O(\mu n^2 + n^3)$

sorting:
$$f(x_{1:\lambda}) \le f(x_{2:\lambda}) \le ... \le f(x_{\lambda:\lambda})$$

Caution: must use mean m(t) of "old" selected parents; not "new" mean m(t+1)!

⇒ Seeking covariance matrix of fictitious distribution pointing in gradient direction!



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