

# Computational Intelligence

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

### ad 2) design guidelines for variation operators **in practice**

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

- a) reachability
- b) unbiasedness
- c) control

- every recombination results in some  $z \in \mathbb{Z}^n$   
- mutation of  $z$  may then lead to any  $z^* \in \mathbb{Z}^n$  with positive probability in one step

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

ad 2) design guidelines for variation operators **in practice**

$$X = Z^n$$

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

$\Rightarrow$  need analytic solution of an  $\infty$ -dimensional, nonlinear optimization problem with constraints!

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

$$\text{s.t.} \quad p_k = p_{-k} \quad \forall k \in Z, \quad (\text{symmetry w.r.t. } 0)$$

$$\sum_{k=-\infty}^{\infty} p_k = 1, \quad (\text{normalization})$$

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

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

**result:**

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

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

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

**generation of pseudo random numbers:**

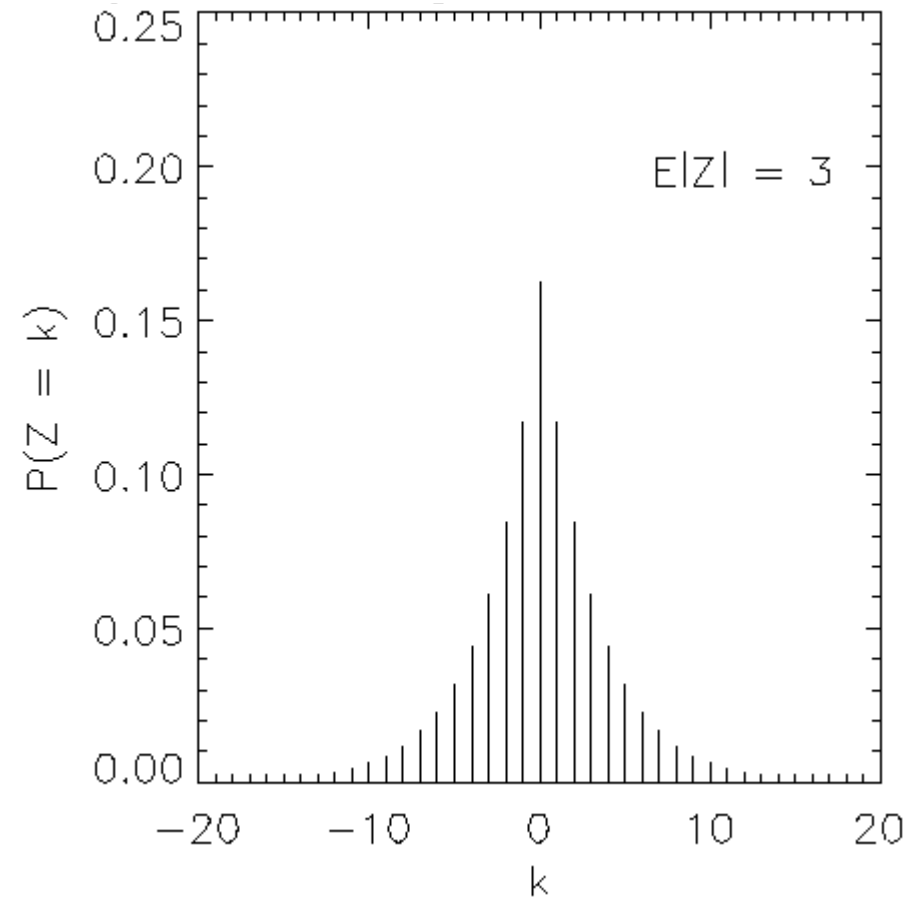
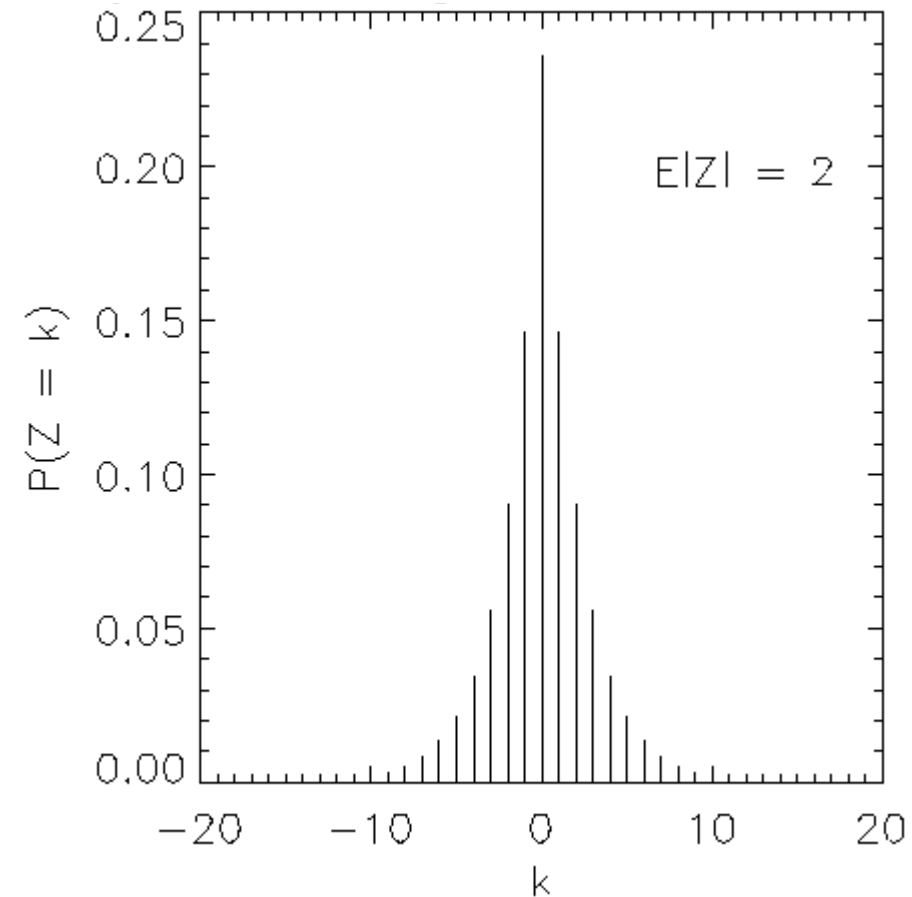
$$Z = G_1 - G_2$$

where

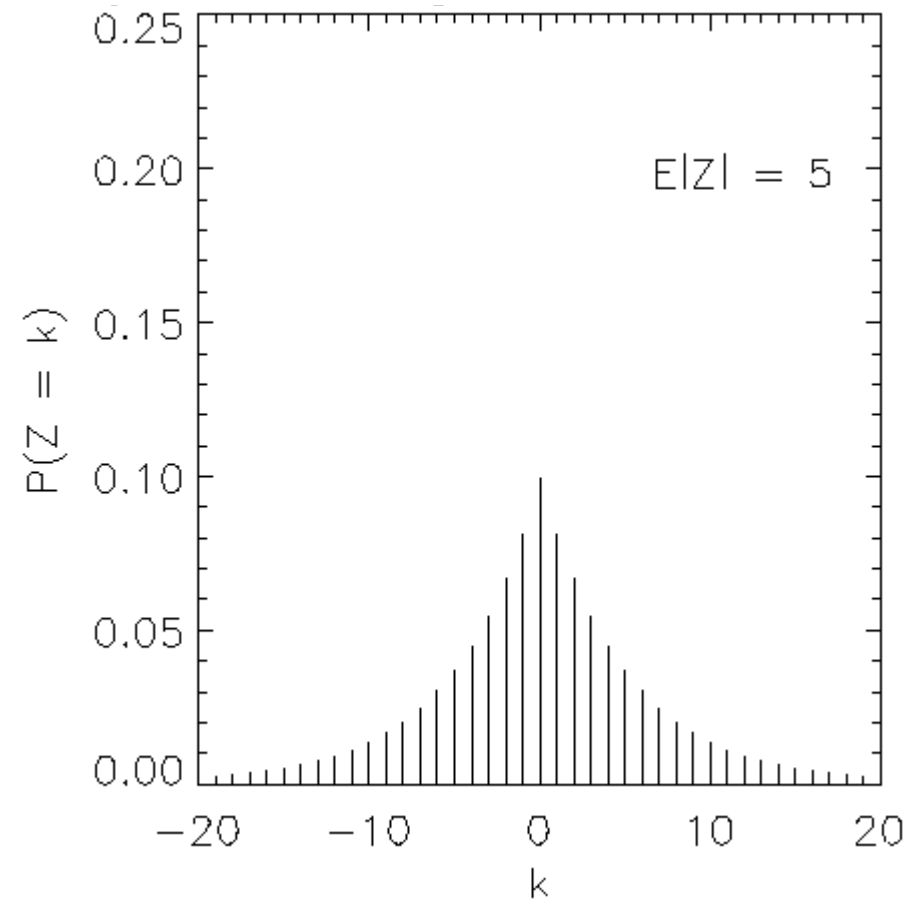
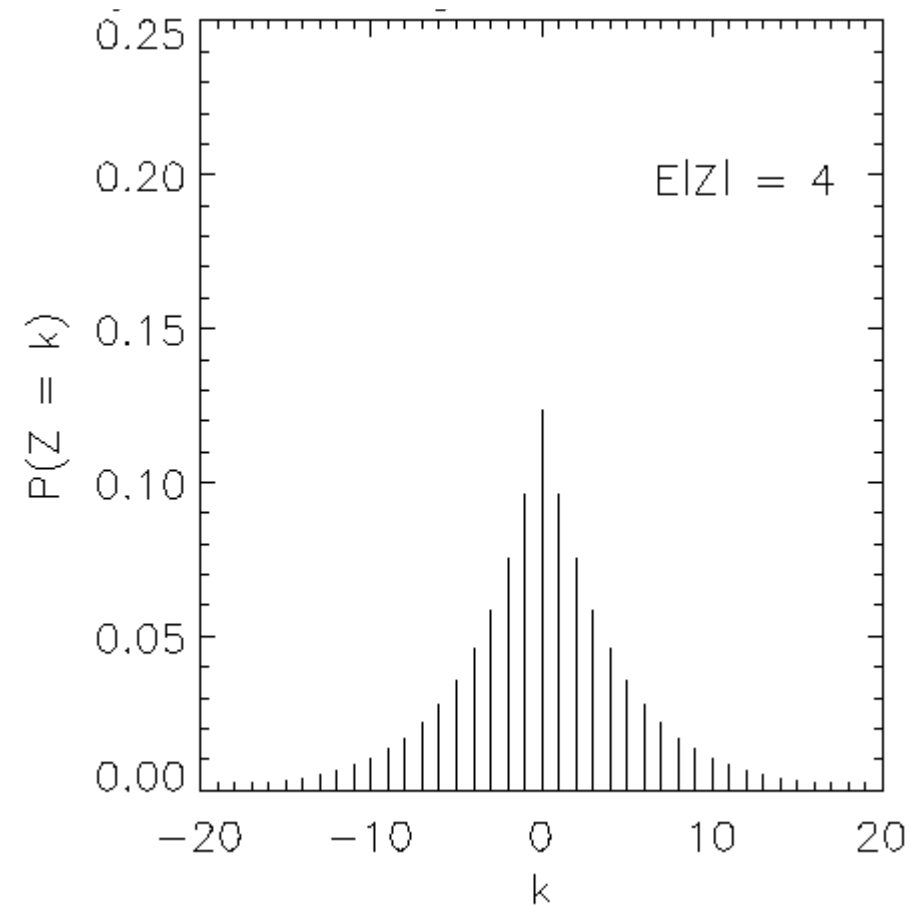
$$U_i \sim U(0, 1) \Rightarrow G_i = \left\lfloor \frac{\log(1 - U_i)}{\log(1 - q)} \right\rfloor, \quad i = 1, 2.$$

stochastic  
independent!

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



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



### 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  $(0,1)$  ?

→ make mean step size  $E[|Z|]$  adjustable!

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

$\downarrow$   
 $\in \mathbb{R}_+$

$\downarrow$   
 $\in (0,1)$

→  $\theta$  adjustable by mutative self adaptation

→ get  $q$  from  $\theta$

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



### Mutative Step Size Control

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

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  $\geq 1$

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

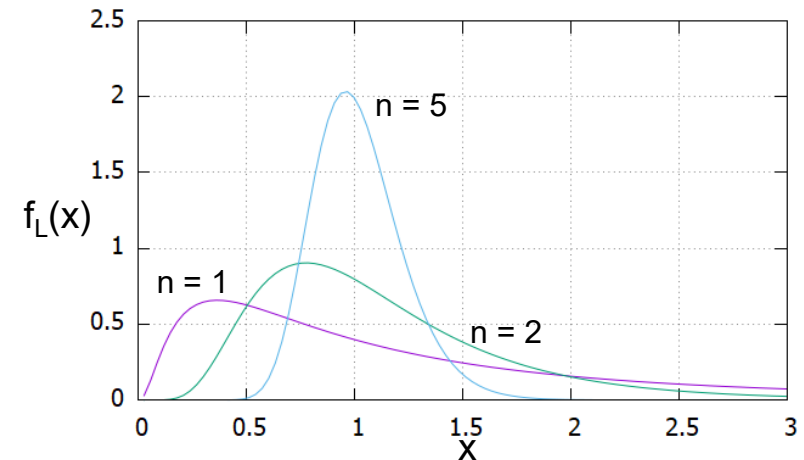
- invented: Schwefel (1977) for real variables
- transferred: Rudolph (1994) for integer variables

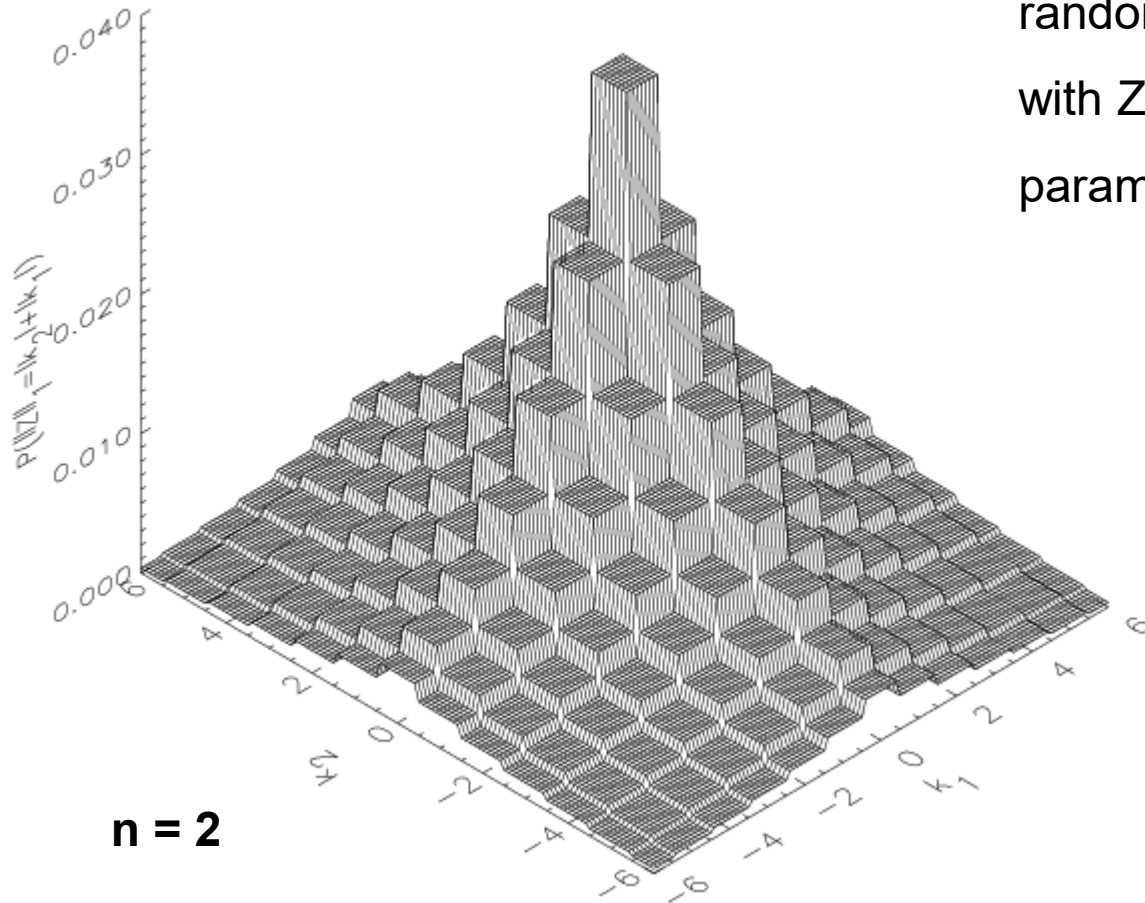
where  $L = \exp(N)$  with  $N \sim N(0, 1/n)$



log-normal distributed

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



**n - dimensional generalization****n = 2**

random vector  $Z = (Z_1, Z_2, \dots, Z_n)$   
with  $Z_i = G_{1,i} - G_{2,i}$  (stoch. indep.);  
parameter  $q$  for all  $G_{1i}, G_{2i}$  equal

n - dimensional generalization

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

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

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

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

⇒ n-dimensional distribution is symmetric w.r.t.  $\ell_1$  norm!

⇒ all random vectors with same step length have same probability!

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

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

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

self-adaptation
calculate from  $\theta$

**Algorithm:**

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

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

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

calculate new  $q$  for  $G_i$  from  $\theta_{t+1}$

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

recombination : discrete (uniform crossover)

selection :  $(\mu, \lambda)$ -selection

(Rudolph, PPSN 1994)

**Example:**  $(1, \lambda)$ -EA with  $\lambda = 10$ ;  $f(x) = x^2 \rightarrow \min!$ ;  $n = 10$

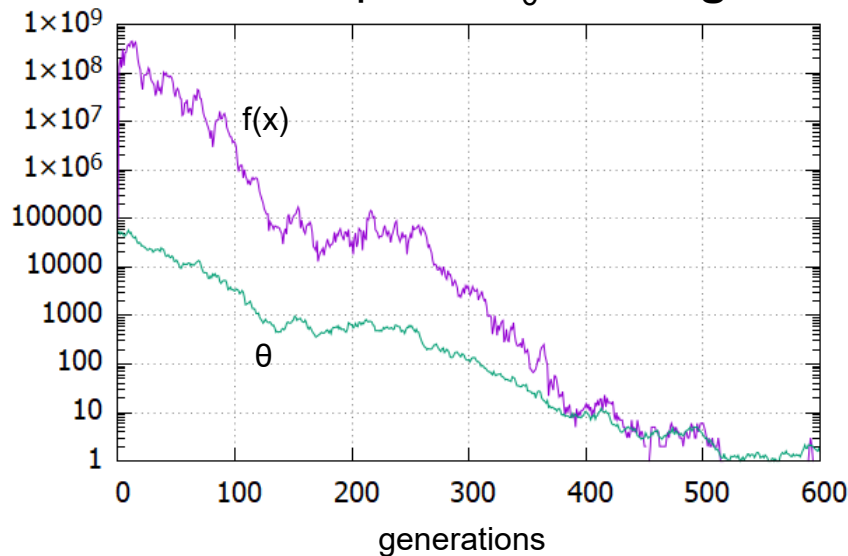
$$X^{(0)} \in [100, 101]^n \cap \mathbb{Z}^n$$

$$\theta_0 = 50\,000$$

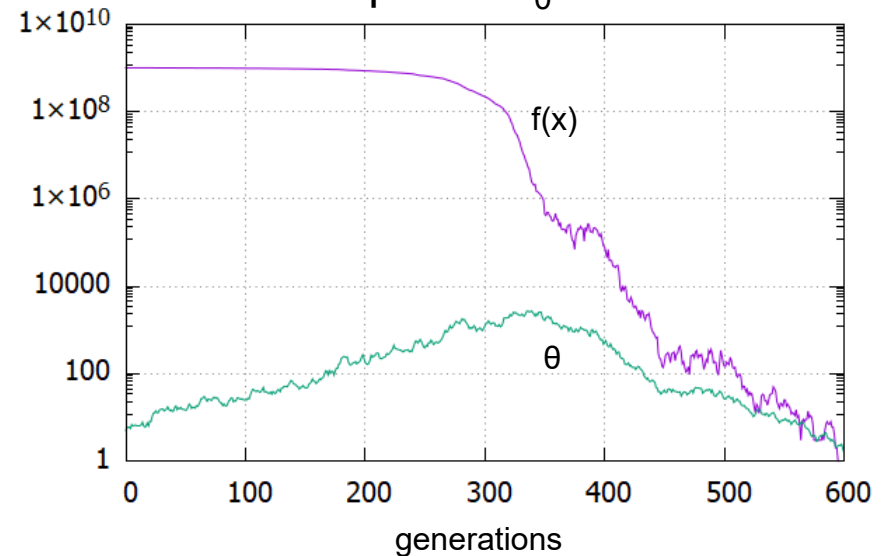
$$X^{(0)} \in [10000, 10100]^n \cap \mathbb{Z}^n$$

$$\theta_0 = 5$$

initial step size  $\theta_0$  too large



initial step size  $\theta_0$  too small



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
- c) control → mutation distribution with parameters

⇒ leads to CMA-ES !



**Covariance  
Matrix  
Adaptation**

mutation:  $Y = X + Z$

$Z \sim N(0, C)$  multinormal distribution



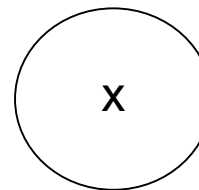
maximum entropy distribution for support  $\mathbb{R}^n$ , given expectation vector and covariance matrix

how should we choose covariance matrix  $C$ ?

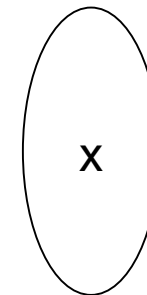
unless we have not learned something about the problem during search

⇒ don't prefer any direction!

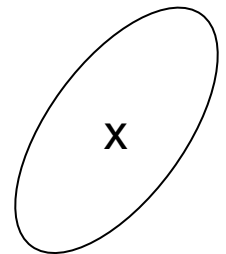
⇒ covariance matrix  $C = I_n$  (unit matrix)



$C = I_n$



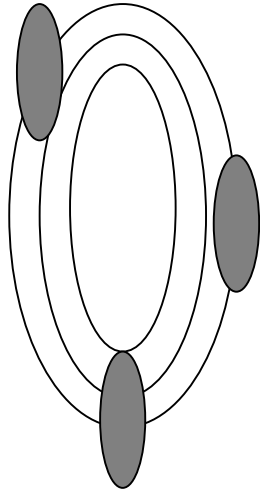
$C = \text{diag}(s_1, \dots, s_n)$



$C$  orthogonal



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



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

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

$Z \sim N(0, C)$  with density

$$f_Z(x) = \frac{1}{(2\pi)^{n/2} |C|^{1/2}} \exp\left(-\frac{1}{2} x' C^{-1} x\right)$$

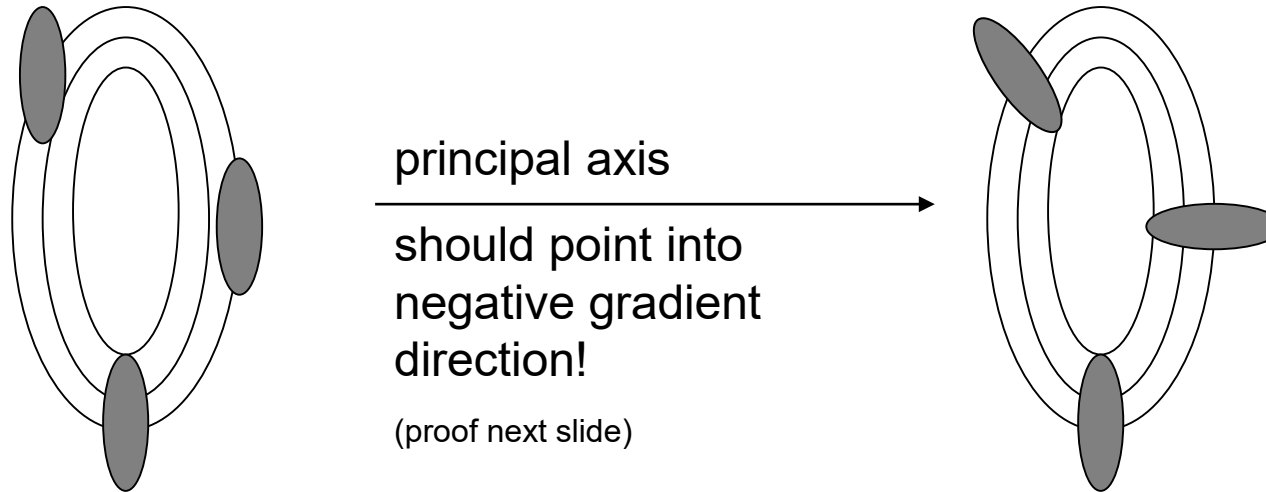
since then many proposals how to adapt the covariance matrix

$\Rightarrow$  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:**  $O(n^6)!$  (Rudolph 1992)

**doubts:** are equi-aligned isolines really optimal?



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

(Schwefel OK in this situation)

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

$$\begin{aligned}
 f(x + rQu) &= \frac{1}{2} (x + rQu)' A (x + rQu) + b'(x + rQu) + c \\
 &= \frac{1}{2} (x'Ax + 2rx' AQu + r^2u'Q' AQu) + b'x + rb'Qu + c \\
 &= f(x) + rx' AQu + rb'Qu + \frac{1}{2} r^2u'Q' AQu \\
 &= f(x) + r(Ax + b + \frac{r}{2}AQu)'Qu \\
 &= f(x) + r(\nabla f(x) + \frac{r}{2}AQu)'Qu \\
 &= f(x) + r \nabla f(x)'Qu + \frac{r^2}{2} u'Q' AQu \\
 &= f(x) + r \nabla f(x)'Qu + \frac{r^2}{2} \quad \rightarrow \text{min!}
 \end{aligned}$$

if  $Qu$  were deterministic ...

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

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

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

basic constraint:  $C^{(k)}$  must be positive definite (p.d.) and symmetric for all  $k \geq 0$ ,  
otherwise Cholesky decomposition impossible:  $C = Q'Q$

### Lemma

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

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

### Proof:

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

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

■

**Theorem**

A quadratic matrix  $C^{(k)}$  is symmetric and positive definite for all  $k \geq 0$ ,

if it is built via the iterative formula  $C^{(k+1)} = \alpha_k C^{(k)} + \beta_k v_k v_k'$

where  $C^{(0)} = I_n$ ,  $v_k \neq 0$ ,  $\alpha_k > 0$  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  $v_{ij} = v_i \cdot v_j = v_j \cdot v_i = v_{ji}$  for all  $i, j$  and
- for all  $x \in \mathbb{R}^n$  :  $x' (vv') x = (x'v) \cdot (v'x) = (x'v)^2 \geq 0$ .

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

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. ■

**Idea:** Don't estimate matrix  $\mathbf{C}$  in each iteration! Instead, approximate iteratively!

(Hansen, Ostermeier et al. 1996ff.)

→ **C**ovariance **M**atrix **A**daptation **E**volutionary **A**lgorithm (CMA-EA)

Set initial covariance matrix to  $\mathbf{C}^{(0)} = \mathbf{I}_n$

$$\mathbf{C}^{(t+1)} = (1-\eta) \mathbf{C}^{(t)} + \eta \sum_{i=1}^{\mu} w_i (\mathbf{x}_{i:\lambda} - \mathbf{m}^{(t)}) (\mathbf{x}_{i:\lambda} - \mathbf{m}^{(t)})'$$

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

$w_i$  : weights; mostly  $1/\mu$

$$\mathbf{m} = \frac{1}{\mu} \sum_{i=1}^{\mu} \mathbf{x}_{i:\lambda} \quad \text{mean of all selected parents}$$

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

sorting:  $f(\mathbf{x}_{1:\lambda}) \leq f(\mathbf{x}_{2:\lambda}) \leq \dots \leq f(\mathbf{x}_{\lambda:\lambda})$

**Caution:** must use mean  $\mathbf{m}^{(t)}$  of “old“ selected parents; not „new“ mean  $\mathbf{m}^{(t+1)}$ !

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

**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://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!