

Evolutionary Programming (EP)

anfangs: endliche Automaten zur Symbolketten-Vorhersage

heute: numerische Optimierung (ähnlich ES)

wichtige Eigenschaften:

- evolvierende Einheit ist die Art (Spezies)
- keine Rekombination; kein Geburtenüberschuss ($\lambda = \mu$)
- normalverteilte Mutationen (ähnlich ES)
- Turnierselektion zwischen alten und neuen Spezies (Majoritätsauswahl)
- Selbstanpassung der Mutations-Varianzen (ähnlich ES)

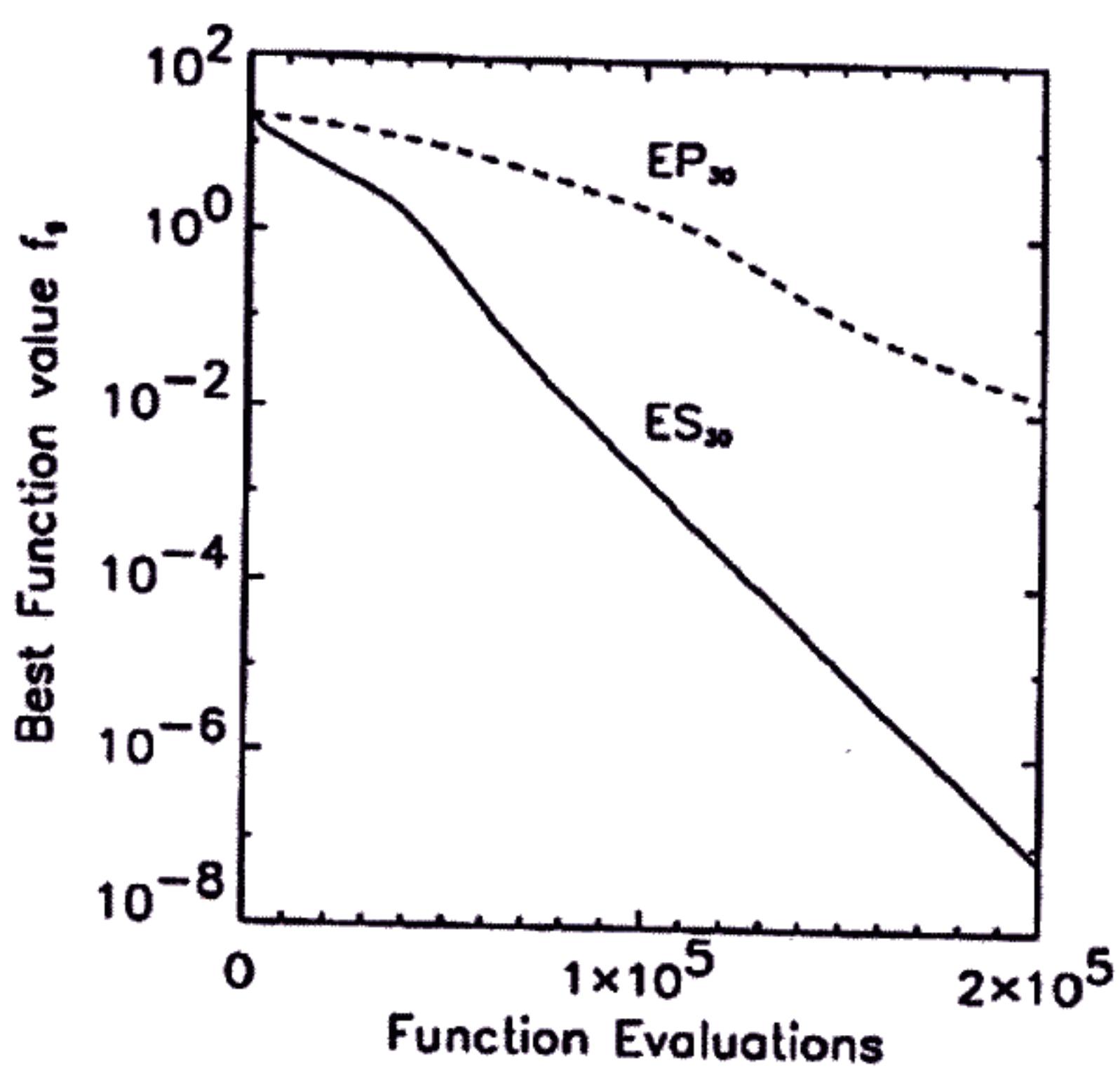
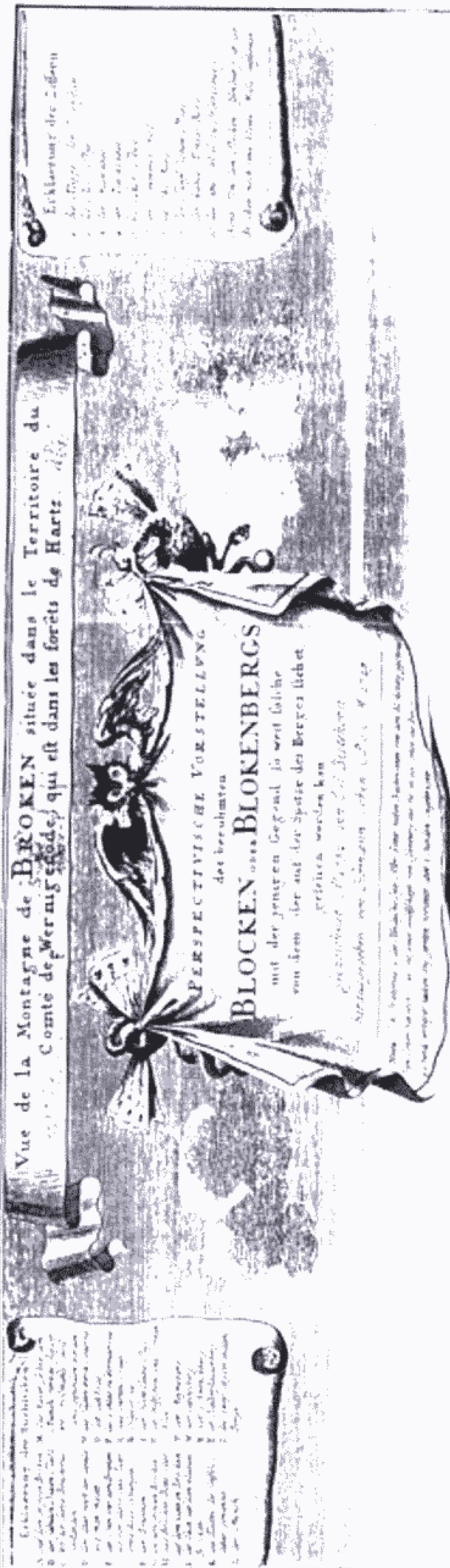


Figure 3: Experimental runs of an evolution strategy and evolutionary programming on f_9 .

↑

Ackley - Funktion



Evolutionsstrategien (ES)

anfangs: experimentelles Optimierverfahren

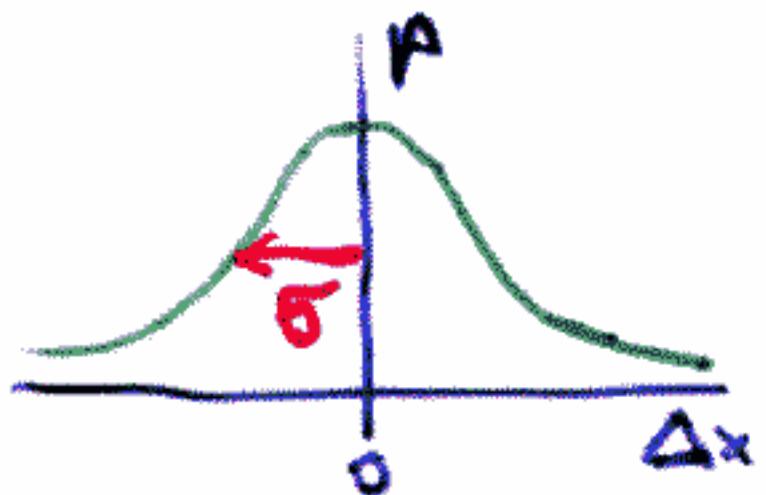
später: numerische Parameteroptimierung in Simulationsmodellen

wichtige Eigenschaften:

- evolvierende Einheit ist das Individuum (Phänotyp)
- Reproduktion mit Geburtenüberschuss ($\lambda > \mu$); rein zufällige Paarung
- Rekombination und Gaußsche Mutation als Quellen der Variation / Innovation
- Umweltselektion: Elimination der $\lambda - \mu$ Schlechtesten
- Selbstadaptation der (internen) Strategieparameter (Varianzen, Kovarianzen)
- gemischtes Diffusions-/Migrationsmodell für skalierbare Parallelität

ES Evolutionstrategien

Mutationen: Änderungen normalverteilt
(für reellwertige Variable)



$$p(\Delta x_i) = \frac{1}{\sqrt{2\pi}\sigma_i} e^{-\frac{\Delta x_i^2}{2\sigma_i^2}}$$

Rekombination:

a) intermediär

Mittelwert \bar{x} aus Elternpas.

b) diskret

Werte komponentenweise von verschied. Eltern

Reproduktion:

$\lambda > \mu$ Geburtenüberschuss
(à la Darwin/Malthus)

Selektion

: deterministisch

μ beste aus λ

(μ, λ) ES

bzw. $\lambda + \mu$ $(\mu + \lambda)$ ES

Selbstadaptation der σ_i :



Werden ebenfalls mutiert!

Individuum:

$x_1, x_2, \dots, x_n, \sigma_1, \sigma_2, \dots, \sigma_n, \dots$
--

Evolutionssstrategie (kanonisch)

Individuen : Vektor aus Objektvariablen und Strategievariablen (internes Modell)

Mutation : zufällige Änderung aller Variablen

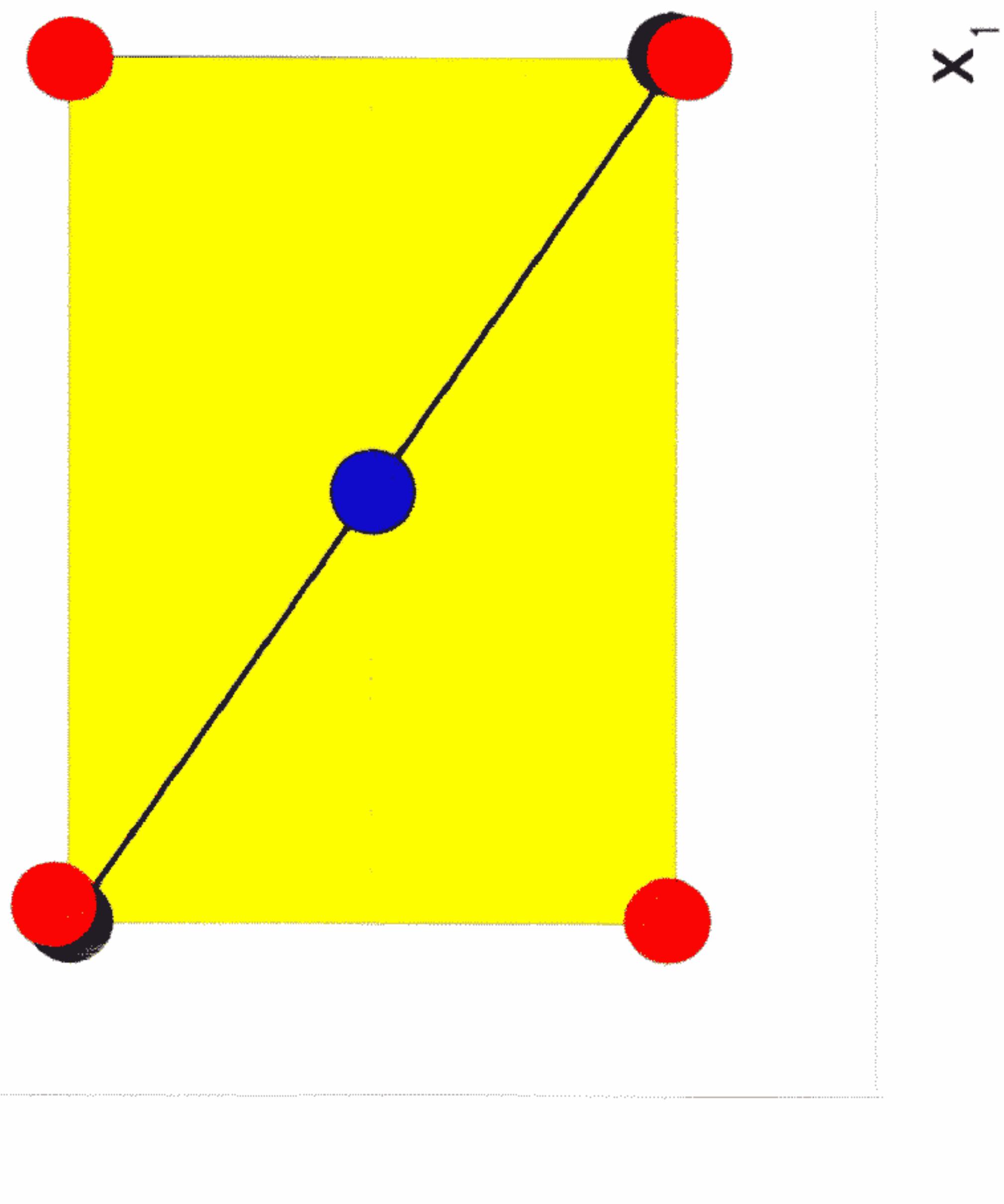
Rekombination: mittelnd oder komponentenweise

Reproduktion : $\lambda > \mu$, Geburtenüberschuss

Selektion : Die μ Besten von $\lambda(+\mu)$ werden Eltern mit gleicher Wahrscheinlichkeit (environmental selection)

Selbstanpassung der Strategievariablen!

ES Rekombination



Eltern

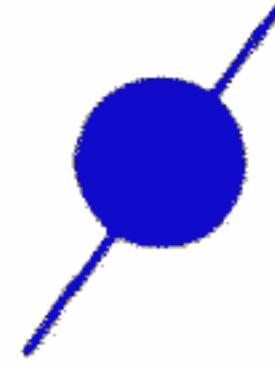


Nachkommen:

diskrete
Rekombination



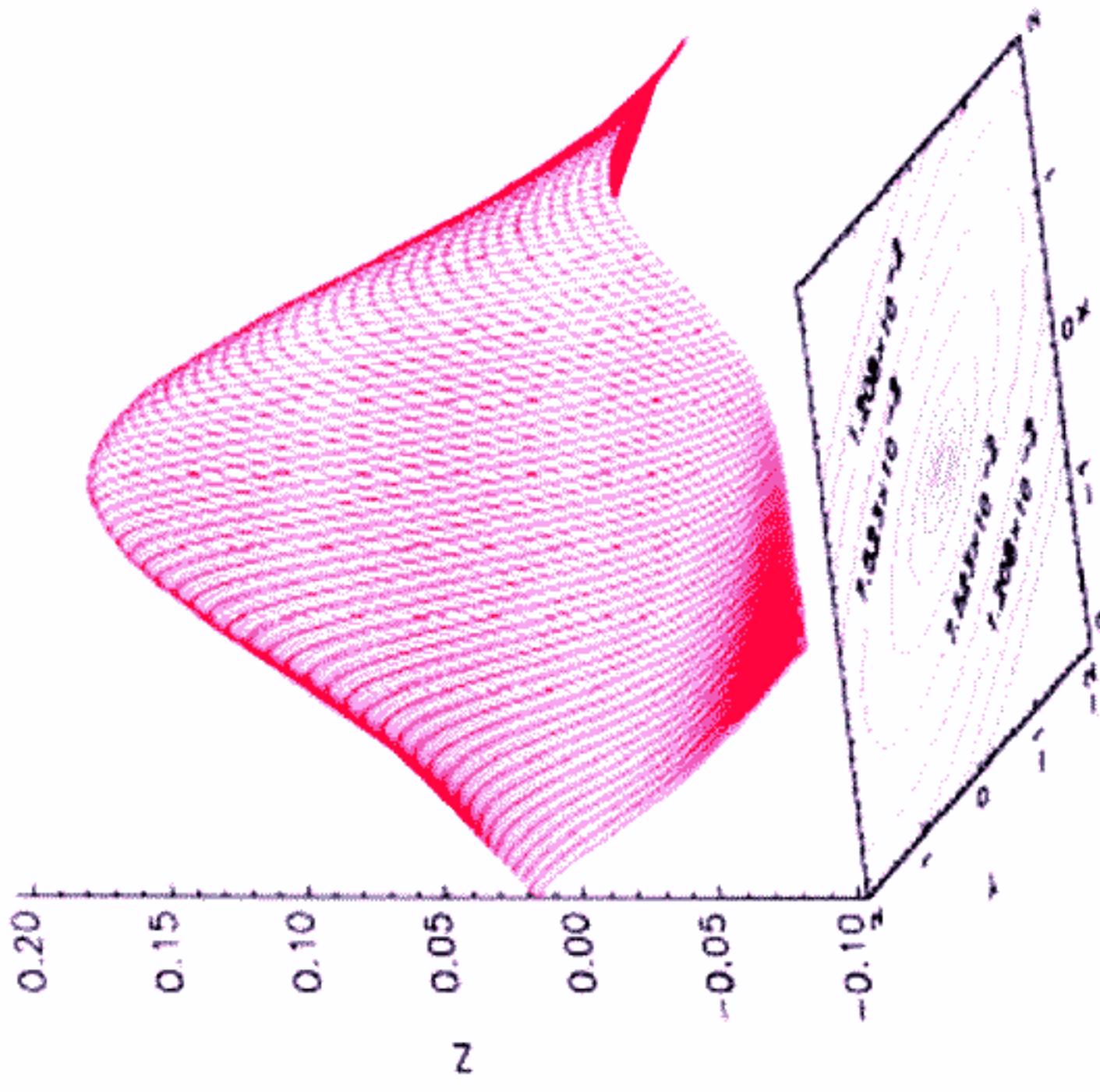
intermediäre
Rekombination



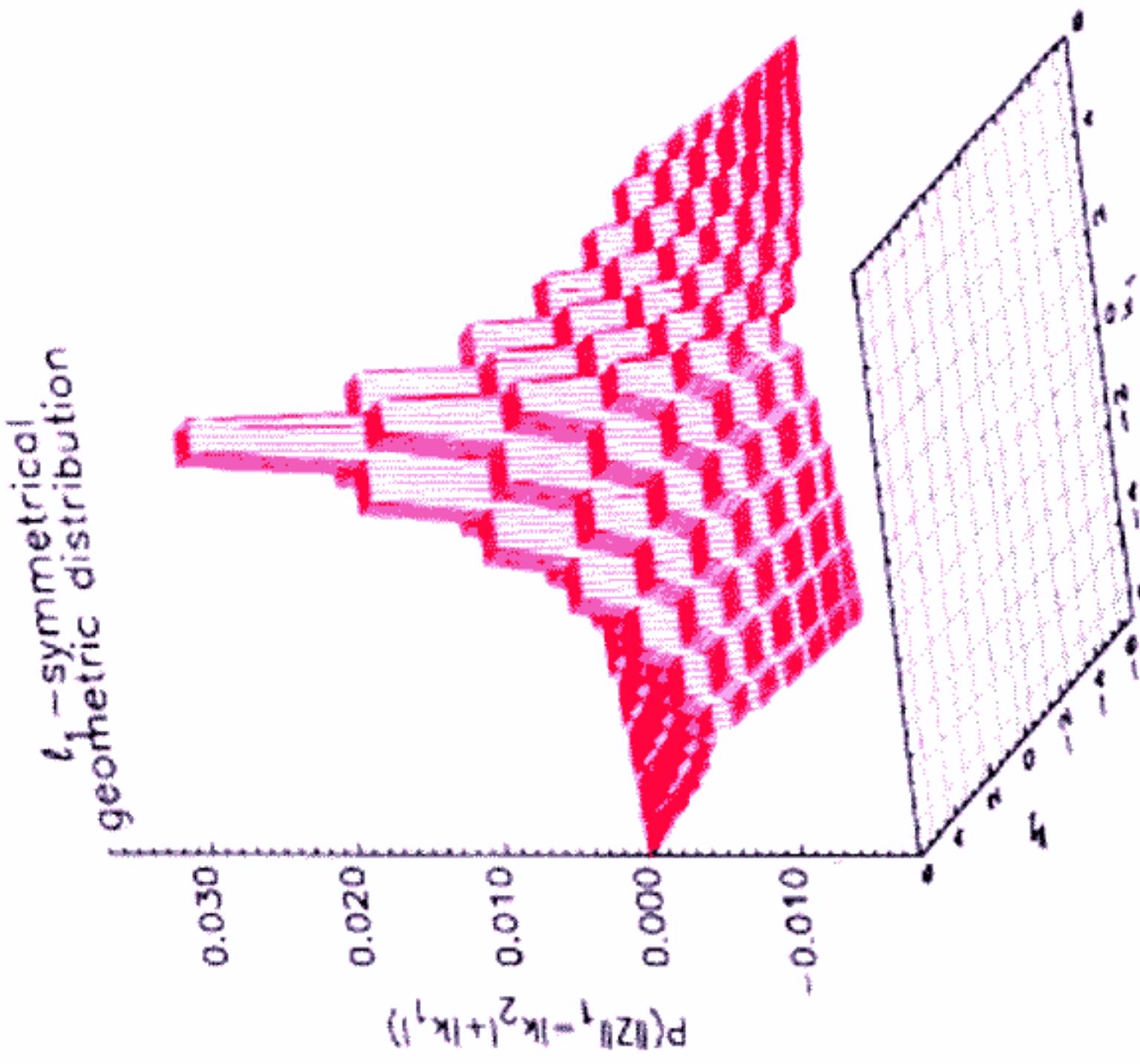
allgemeine
Rekombination



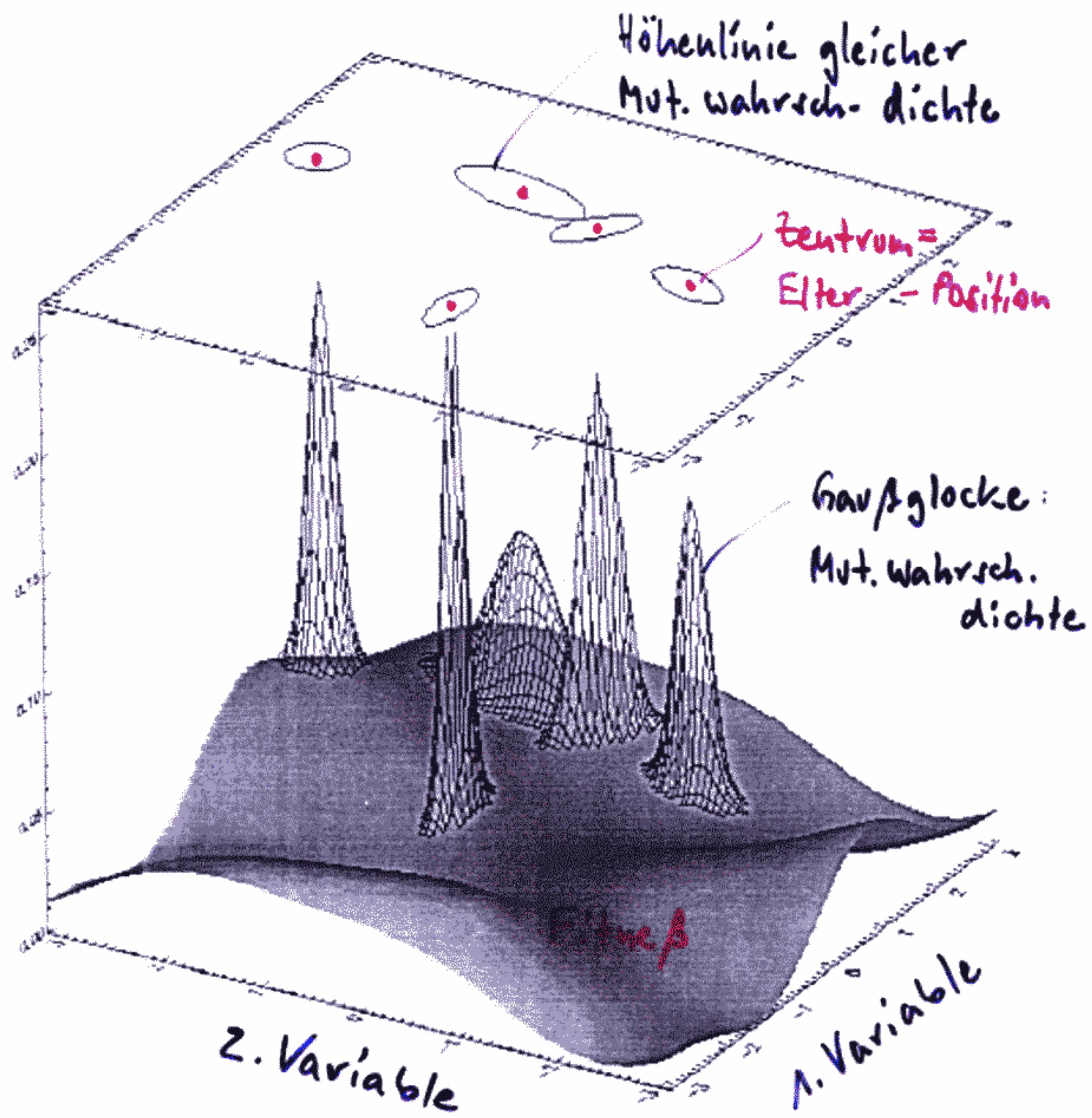
Mutationsdichtheitverteilung



Normalverteilung für kontinuierliche Variable



geometrische Verteilung für diskrete Variable



normalverteilte Mutationen kontinuierlich

Erwartungswert für Abweichung: δ

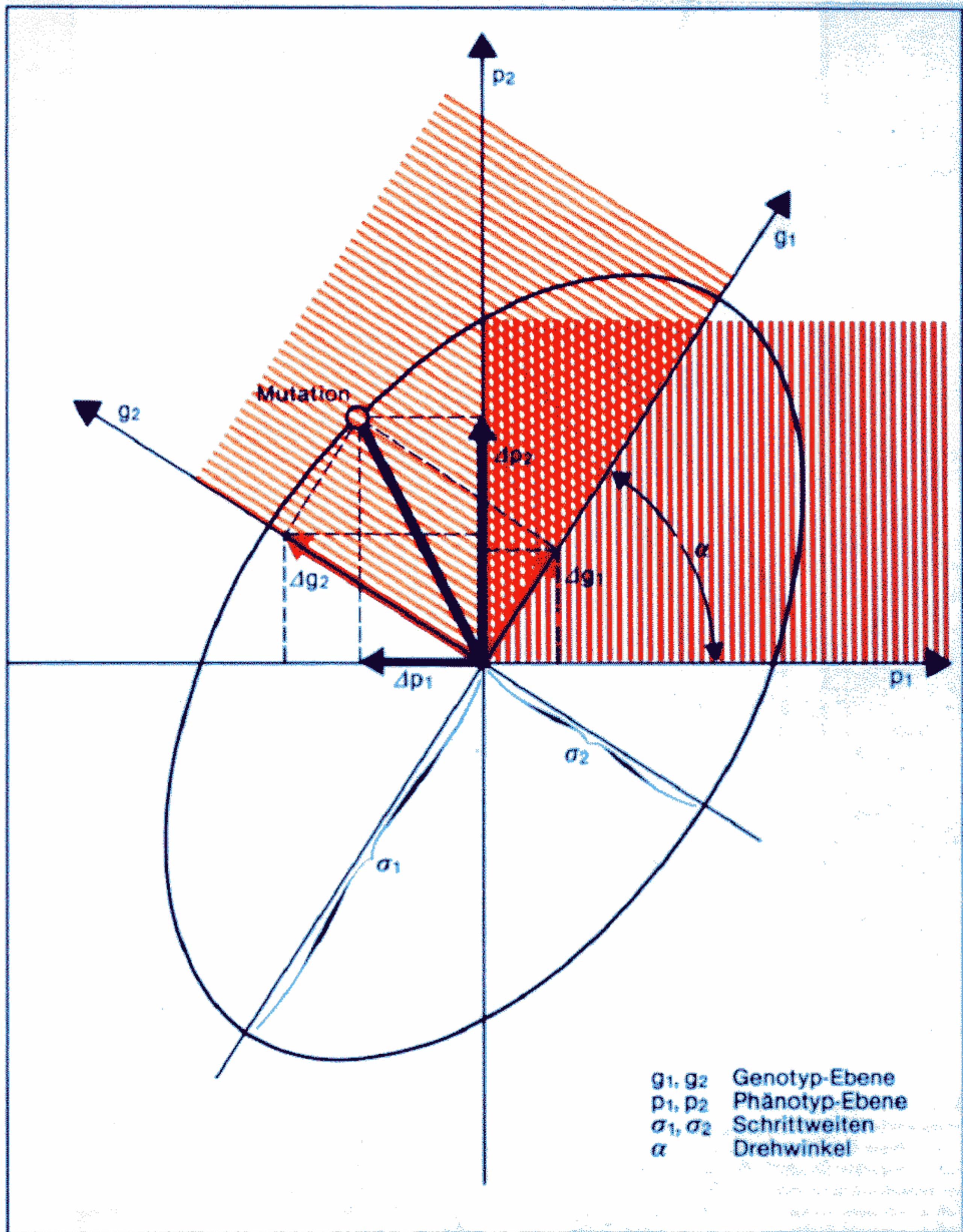
Freiheitsgrade: Varianzen σ_i^2

Kovarianzen c_{ij}

einfachster Fall: $c_{ij} = 0 \quad \forall i \neq j$

$$\sigma_i^2 = \sigma \quad \forall i = 1, 2, \dots, n$$

Polygenie, Pleiotropie, korrelierte phänotypische Veränderungen



The evolution of Evolution Strategies

Cons

two individuals

just creeping random search

$(1+1)$ ES

Pros

June 12, 1964

worked in experiments

(as opposed to other methods)

μ ancestors

$(\mu+1)$ ES

ridiculous waste of storage capacity

speedup through recombination

λ descendants

$(1+\lambda)$ ES

ridiculous not to use new info. immediately

if parallel: speedup $\sim \log \lambda$

non-elitist

$(1,\lambda)$ ES

ridiculous to forget good intermed. solution

self-adaptation of mutability works (one common σ)

$\kappa = 1$

(μ,λ) ES

ridiculous to conserve inferior solutions

recombination helps auto-scaling, variable metric, and speeds up

$\kappa = \infty$

$(\mu+\lambda)$ ES

hampers self-adaptation, does not work for moving goals

elitist version - no loss of good results

(μ,κ,λ,g) ES

contemporary standard

$1 \leq \kappa \leq \infty$ max. life span (reproduction cycles)

$2 \leq g \leq \mu$ recombination uses g ancestors

Evolution der ES

- 12.6.1964 Geburt der (1+1)ES: experimentelle Optimierung
wichtig: alle Variablen zugleich etwas (zufällig) verändern
- 1965 meine Diplomarbeit zeigt: Stagnationsgefahr bei Binomialverlgl.
Düsensexperimente für die AEG
- 1968
- 1971 Diss. Rechenberg: erste Theorie der $(1+1)$ ES für $x \in \mathbb{R}^n$
und erste ES mit Rekombination: $(\mu+1)$ ES
- 1972-1975 DFG-Projekte: numerische ES mit autoadaptiven (Ko-)Varianzen
 $(1+\lambda) / (1,\lambda) / (\mu+\lambda) / (\mu,\lambda)$ / $(\mu,1)$ ES-Simulationen, z.B. (10,100)ES
- 1975 Diss. Schwefel: erste Theorie der $(1+\lambda) / (1,\lambda)$ ES
- Nomenklatur:**
- μ : Zahl der Eltern, λ : Zahl der Nachkommen einer Generation
Plus (+): max. Lebensdauer unendlich, Komma (,): max. 1 Generation
heute wesentlich verallgemeinerte ES-Versionen

$$(\mu, \kappa, \lambda, \rho)ES := (P^{(0)}, \mu, \kappa, \lambda, \text{rec}, p_r, \rho, \gamma, \omega, \text{mut}, p_m, \tau, \tau_0, \delta, \beta, \text{sel}, \zeta, t, \varepsilon)$$

with

$P^{(0)} := (\mathbf{a}_1, \dots, \mathbf{a}_\mu)^{(0)} \in I^\mu$	$I := \mathbb{N}_0 \times \mathbb{R}^n \times \mathbb{R}_+^{n_\sigma} \times [-\pi, \pi]^{n_\sigma}$	start population
$\mu \in \mathbb{N}$	$\mu \geq 1$	number of parents
$\kappa \in \mathbb{N}$	$\kappa \geq 1$	upper limit for life span
$\lambda \in \mathbb{N}$	$\lambda > \mu$ if $\kappa = 1$	number of offspring
$\text{rec} : I^\mu \rightarrow I$		recombination operator
$p_r \in \mathbb{R}_+^3$	$0 \leq p_r \leq 1$	recombination probability
$\rho \in \mathbb{N}^3$	$1 \leq \rho \leq \mu$	number of ancestors for each descendant
$\gamma \in \mathbb{N}^3$	$1 \leq \gamma \leq n_x - 1$ $\gamma \geq \rho - 1$	number of crossover sites in a string of n_x elements
$\omega \in \{0, 1, 2, 3, \dots\}^3$		type of recombination
$\text{mut} : I \rightarrow I$		mutation operator
$p_m \in \mathbb{R}_+^3$	$0 < p_m \leq 1$	mutation probability
$\tau, \tau_0, \delta \in \mathbb{R}_+$	$0 \leq \delta \leq 1$	step length variabilities
$\beta \in \mathbb{R}_+$	$0 \leq \beta \leq \frac{\pi}{4}$	correlation variability
$\text{sel} : I^{\mu+\lambda} \rightarrow I^\mu$		selection operator
$\zeta \in \mathbb{N}$	$2 \leq \zeta \leq \mu + \lambda$	tournament participators
$t : I^{2\mu} \rightarrow \{0, 1\}$		termination criterion
$\varepsilon \in \mathbb{R}_+^4$		accuracies required.

Weitere ES-Varianten

- mit Diploidie, Dominanz + Rezessivität
für Vektoroptimierung (mehrfache Zielfunktion)
- mit Annihilation / Migration
für globale Optimierung (großkörnig)
- mit lokaler Interaktion (Diffusion)
für globale Optimierung (feinkörnig)
- geschachtelte ES
für (sequentielle) globale Optimierung

$[\mu/s', \lambda'(\mu/s, \lambda), v]$ ES

λ' Subpopulationen agieren v
Generationen lang unabh. voneinander

Auslese unter Subpopulationen

Notation

- Individual: $\vec{a} \in I$.
- Individual space: $I = IR^n \times S$.
- Strategy parameters: S .
- Population: $P^{(t)} = \{\vec{a}_1, \dots, \vec{a}_n\} \in I^k$, $k \in \{\mu, \lambda\}$.
- Recombination: $\text{rec} : I^\mu \rightarrow I$.
- Mutation: $\text{mut} : I \rightarrow I$.
- Selection: $\text{sel}_\mu^k : I^k \rightarrow I^\mu$, $k \in \{\lambda, \mu + \lambda\}$.
- Generation transition: $opt_{ES}(P^{(t)}) : I^\mu \rightarrow I^\mu$

$$opt_{ES}(P^{(t)}) = \text{sel}_\mu^k(\sqcup_{i=1}^\lambda \{\text{mut}(\text{rec}(P^{(t)}))\} \sqcup Q)$$

$$(k = \lambda + |Q|, Q \in \{P^{(t)}, \emptyset\}).$$

- Realization of a normally distributed random variable:
 $z \sim N(\zeta, \sigma^2)$.

Random number generation:

$$z_1 = \sqrt{-2 \ln u_1} \sin(2\pi u_2) \quad , \quad z_2 = \sqrt{-2 \ln u_1} \cos(2\pi u_2)$$

$$u_1, u_2 \sim U((0, 1]), z_1, z_2 \sim N(0, 1).$$

The (1 + 1)-ES

- $\mu = \lambda = 1$, no recombination.
- Individual structure: $\vec{a} = (\vec{x}, \sigma) \in I\!\!R^n \times I\!\!R_+$.
- Mutation: 1/5-success rule

$$\tilde{\sigma} := \text{mu}_\sigma(\sigma) = \begin{cases} \sigma / \sqrt[n]{c} & , \text{ if } p > 1/5 \\ \sigma \cdot \sqrt[n]{c} & , \text{ if } p < 1/5 \\ \sigma & , \text{ if } p = 1/5 \end{cases}$$

$$\tilde{\vec{x}} := \text{mu}_{\vec{x}}(\vec{x}) = (x_1 + z_1, \dots, x_n + z_n)$$

- p denotes the measured relative frequency of successful mutations.
- $c = 0.817$ proposed by Schwefel.
- $z_i \sim N_i(0, \tilde{\sigma}^2)$.

- Selection:

$$\text{sel}_1^2(\{\vec{a}, \tilde{\vec{a}}\}) = \begin{cases} \{\tilde{\vec{a}}\} & , \text{ if } f(\tilde{\vec{x}}) \leq f(\vec{x}) \\ \{\vec{a}\} & , \text{ otherwise} \end{cases}$$

- Generation transition:

$opt_{(1+1)-ES}(\{\vec{a}\}) = \text{sel}_1^2(\{\text{mut}(\vec{a})\} \sqcup \{\vec{a}\})$

Pseudocode for a $(1 + 1)$ -ES

ALGORITHM 1 $((1 + 1)\text{-ES})$

```

 $t := 0;$ 
initialize  $P^{(t)} = \{(\vec{x}, \sigma)\};$ 
evaluate  $f(\vec{x});$ 
while ( $T(P^{(t)}) = 0$ ) do
     $(\tilde{\vec{x}}, \tilde{\sigma}) := \text{mut}((\vec{x}, \sigma));$ 
    evaluate  $f(\tilde{\vec{x}});$ 
    if ( $f(\tilde{\vec{x}}) \leq f(\vec{x})$ )
        then  $P^{(t+1)} := \{(\tilde{\vec{x}}, \tilde{\sigma})\};$ 
        else  $P^{(t+1)} := P^{(t)};$ 
     $t := t + 1;$ 
od

```

σ -modification according to Schwefel:

After every n mutations, check how many successes have occurred over the preceding $10 \cdot n$ mutations. If this number is less than $2 \cdot n$, multiply the step length by the factor $c = 0.85$; divide it by 0.85 if more than $2 \cdot n$ successes occurred.

The (μ, λ) -ES

One generation transition function ((μ, λ) -selection):

$$opt_{(\mu, \lambda)-ES}(P^{(t)}) = \text{sel}_{\mu}^{\lambda}(\sqcup_{i=1}^{\lambda} \{\text{mut}(\text{rec}(P^{(t)}))\})$$

ALGORITHM 2 $((\mu, \lambda)$ -ES)

```

 $t := 0;$ 
initialize  $P^{(0)} = \{\vec{a}_1, \dots, \vec{a}_{\mu}\} \in I^{\mu}$ ;
evaluate  $f(\vec{x}_1), \dots, f(\vec{x}_{\mu})$ ;
while  $(T(P^{(t)}) = 0)$  do
     $\tilde{P} := \emptyset$ ;
    for  $i := 1$  to  $\lambda$  do
         $(\tilde{\vec{x}}, \tilde{\vec{\sigma}}, \tilde{\vec{\alpha}}) := \text{mut}(\text{rec}(P^{(t)}))$ ;
        evaluate  $f(\tilde{\vec{x}})$ ;
         $\tilde{P} := \tilde{P} \sqcup \{(\tilde{\vec{x}}, \tilde{\vec{\sigma}}, \tilde{\vec{\alpha}})\}$ ;
    od
     $P^{(t+1)} := \text{sel}_{\mu}^{\lambda}(\tilde{P})$ ;
     $t := t + 1$ ;
od

```

For $(\mu + \lambda)$ -selection:

$$opt_{(\mu+\lambda)-ES}(P^{(t)}) = \text{sel}_{\mu}^{\mu+\lambda}(\sqcup_{i=1}^{\lambda} \{\text{mut}(\text{rec}(P^{(t)}))\} \sqcup P^{(t)})$$

Mutations

Normally distributed variations:

$$p(z) = \frac{1}{\sigma\sqrt{2\pi}} \exp\left(-\frac{(z - \xi)^2}{2\sigma^2}\right)$$

- Expectation $\xi = 0$ is assumed.
- Standard deviation σ must be adapted.

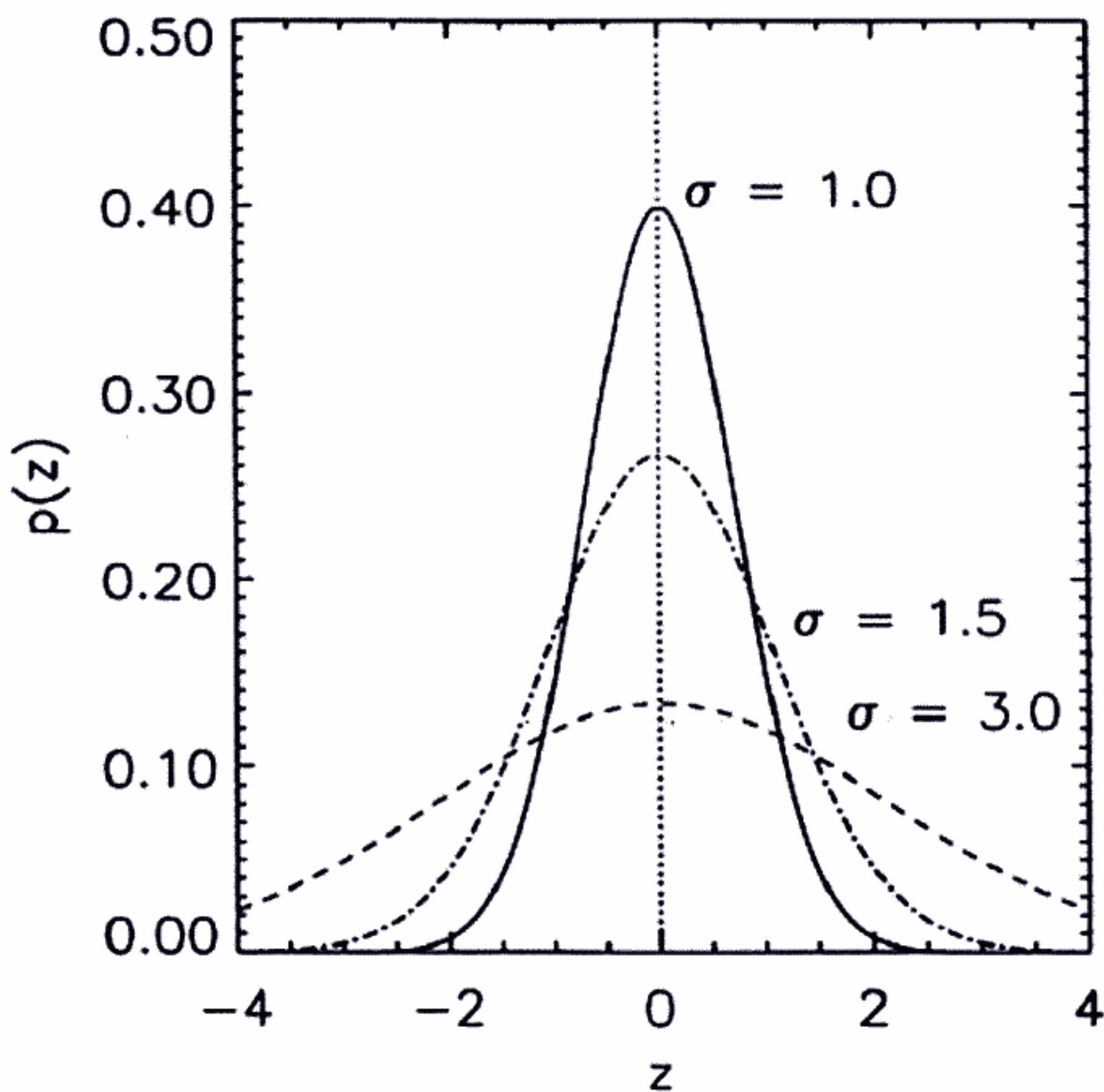


Figure 6: Probability density function $p(z)$ for a normal distribution with $\xi = 0$ and standard deviations $\sigma \in \{1, 1.5, 3\}$.

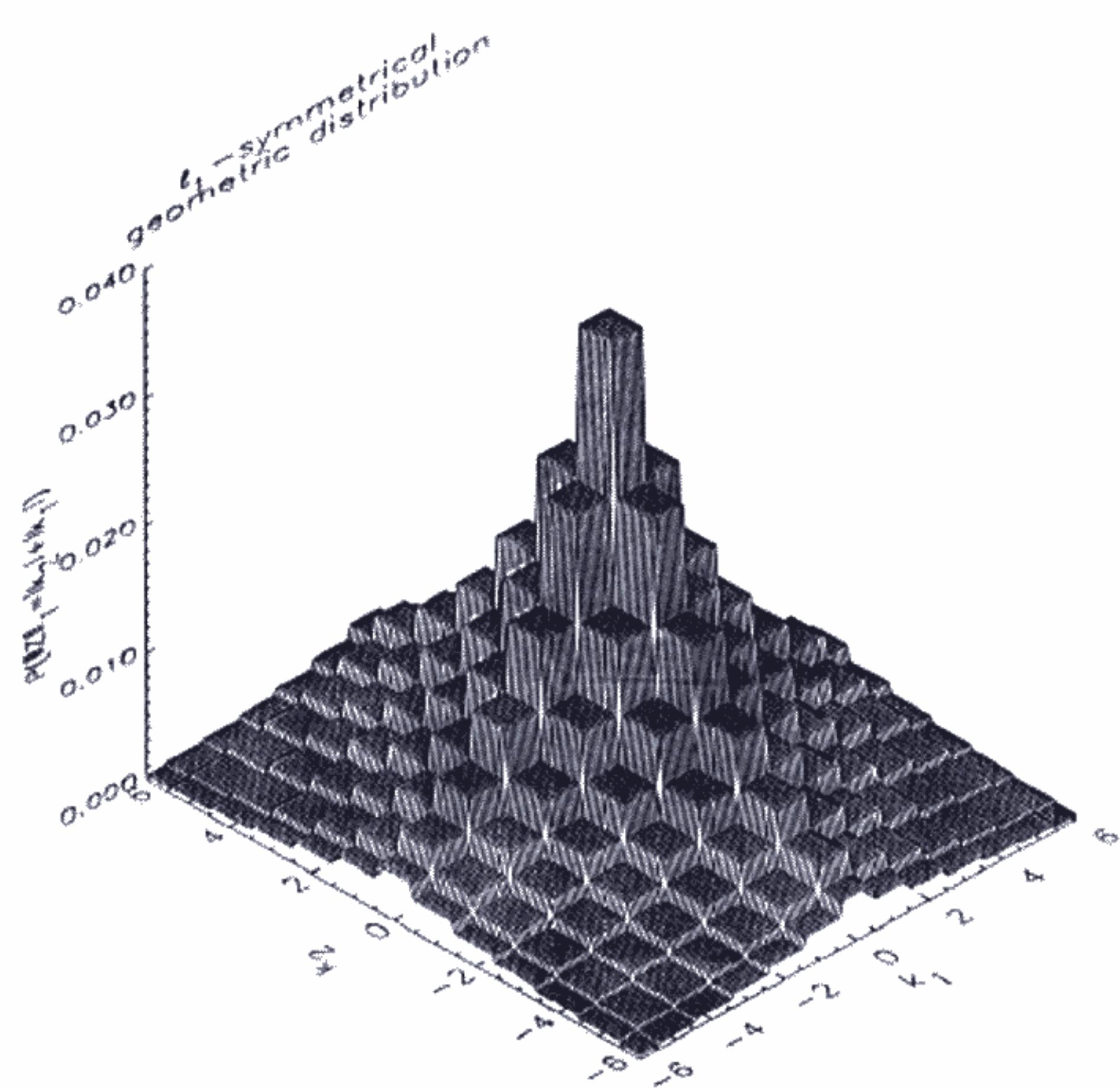


Figure 1: Maximum-entropy probability distribution in \mathbb{Z}^2 for $E[\|Z\|_1] = 5$.

Simple Mutations: $n_\sigma = 1$



equal probability to place an offspring

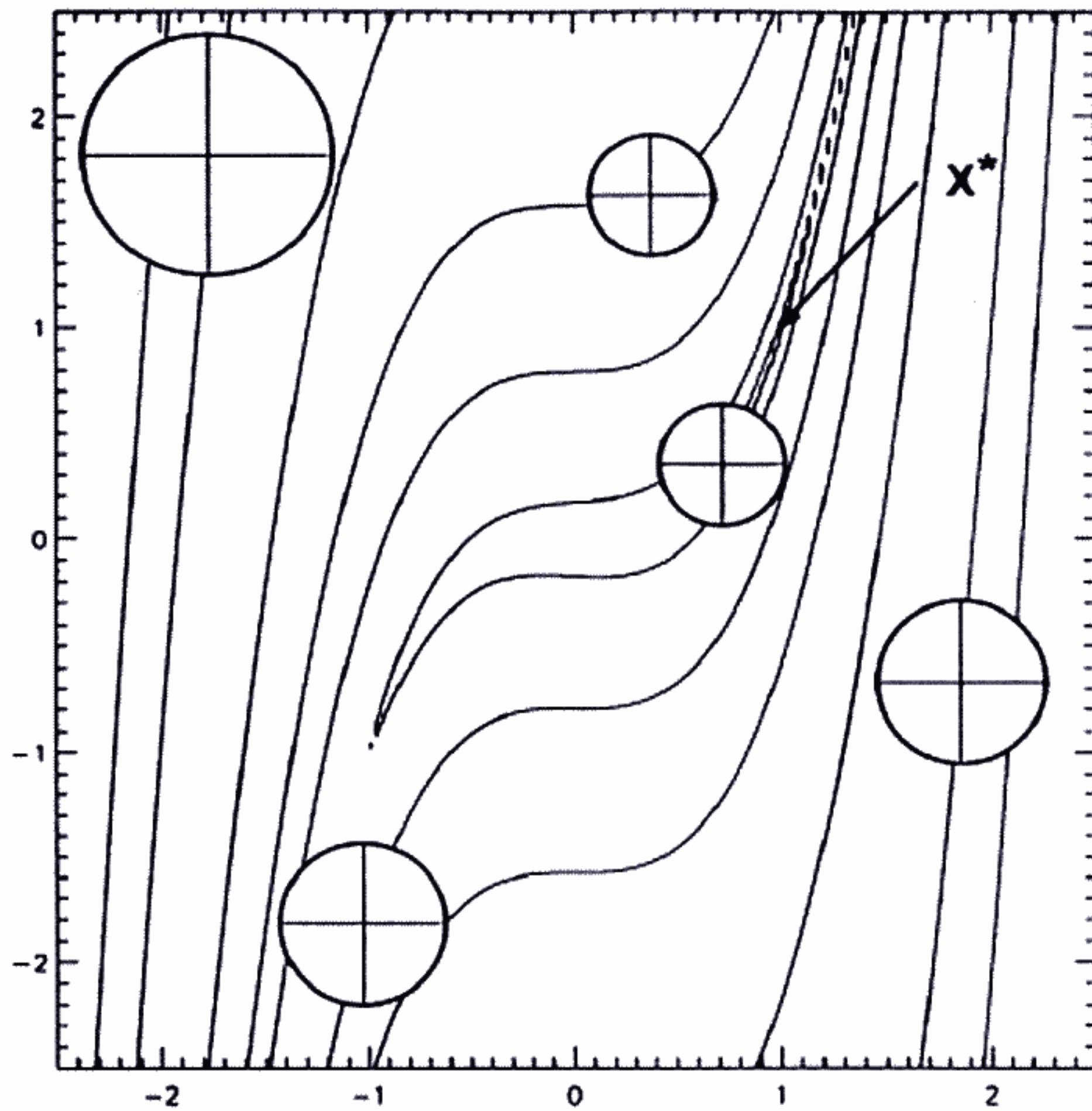
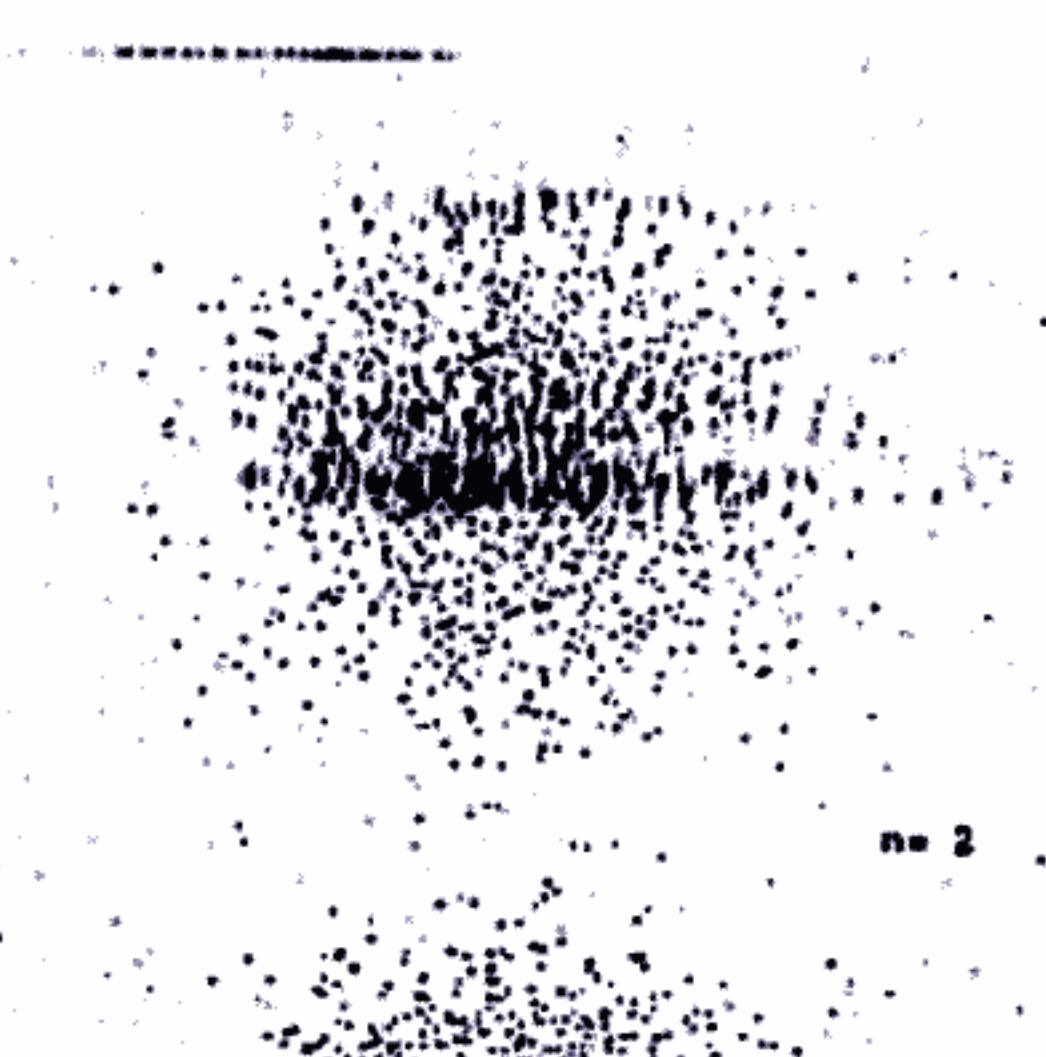


Figure 7: Simple mutations, $n = 2$, $n_\sigma = 1$, ($\Rightarrow n_\alpha = 0$).

$$\begin{aligned} I &= \text{IR}^n \times \text{IR}_+ \\ m'_{\{\tau_0\}}(\vec{x}, \sigma) &= (\vec{x}', \sigma') \\ \tau_0 &\sim 1/\sqrt{n} \end{aligned}$$

$$\begin{aligned} \sigma' &= \sigma \cdot \exp(\tau_0 \cdot N(0, 1)) \\ x'_i &= x_i + \sigma' \cdot N_i(0, 1) \end{aligned}$$



n= 2



n= 10



n= 3



n= 100



n= 5

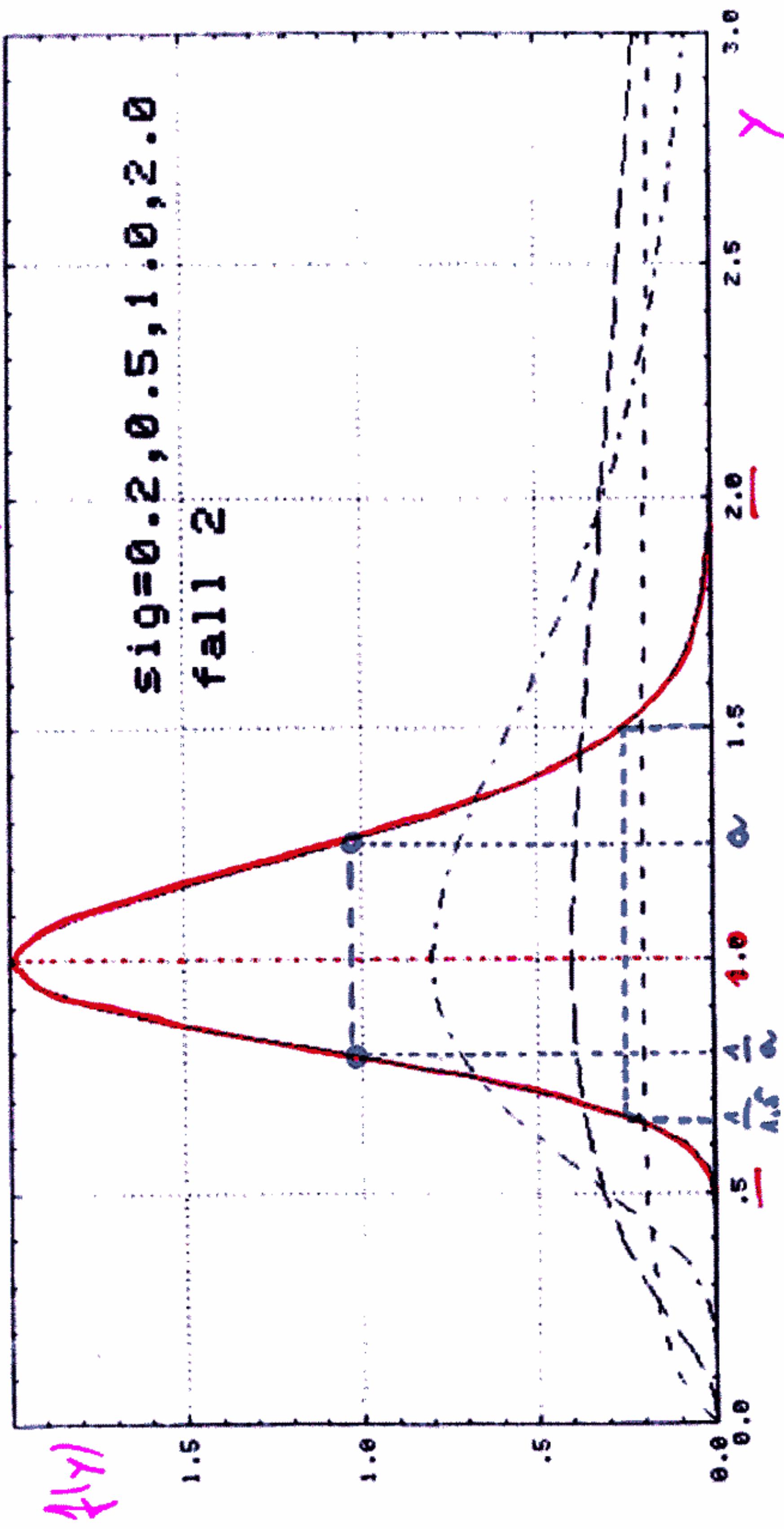


n= 1000



Radialsymmetrische Darstellung einer n=2,3,5,
10,100,1000,10 000 dimensionalen Normalverteilung

metrischer / fallweise Multiplikator



$\log \cdot \text{normal}$
 $y \sim N(\mu, \sigma^2)$
 dann ist

$y = e^z$ logistisch normalverteilt
 $\text{wahrscheinlichster Wert } e^z = 1$

An alternative method for $n_\sigma = 1$:
 (proposed by Rechenberg; see e.g. (Ostermeier 1992)).

$$\begin{aligned} I &= \mathbb{R}^n \times \mathbb{R}_+ \\ m'_{\{\alpha\}}(\vec{x}, \sigma) &= (\vec{x}', \sigma') \end{aligned}$$

$$\begin{aligned} \sigma' &= \sigma \cdot u \\ x'_i &= x_i + \frac{\sigma'}{\sqrt{n}} \cdot N_i(0, 1) \end{aligned}$$

- $u \in \{1, \alpha, 1/\alpha\}$ with equal probability for each of the three outcomes \Rightarrow
 - One-third of the offspring tries larger step-sizes,
 - one-third of the offspring tries smaller step-sizes, and
 - one-third of the offspring keeps step-sizes constant.
- Normally, $\alpha = 1.5$ is used (Ostermeier, 1992).
- Terminology: *Mutational stepsize control* (Rechenberg).
- Extension proposed by (Ostermeier 1992): *Momentum adaptation*.
 (Allow for an adaptation of expected values *different from zero* for the normally distributed random numbers).

Standard Mutations: $n_\sigma = n$

\oplus equal probability to place an offspring

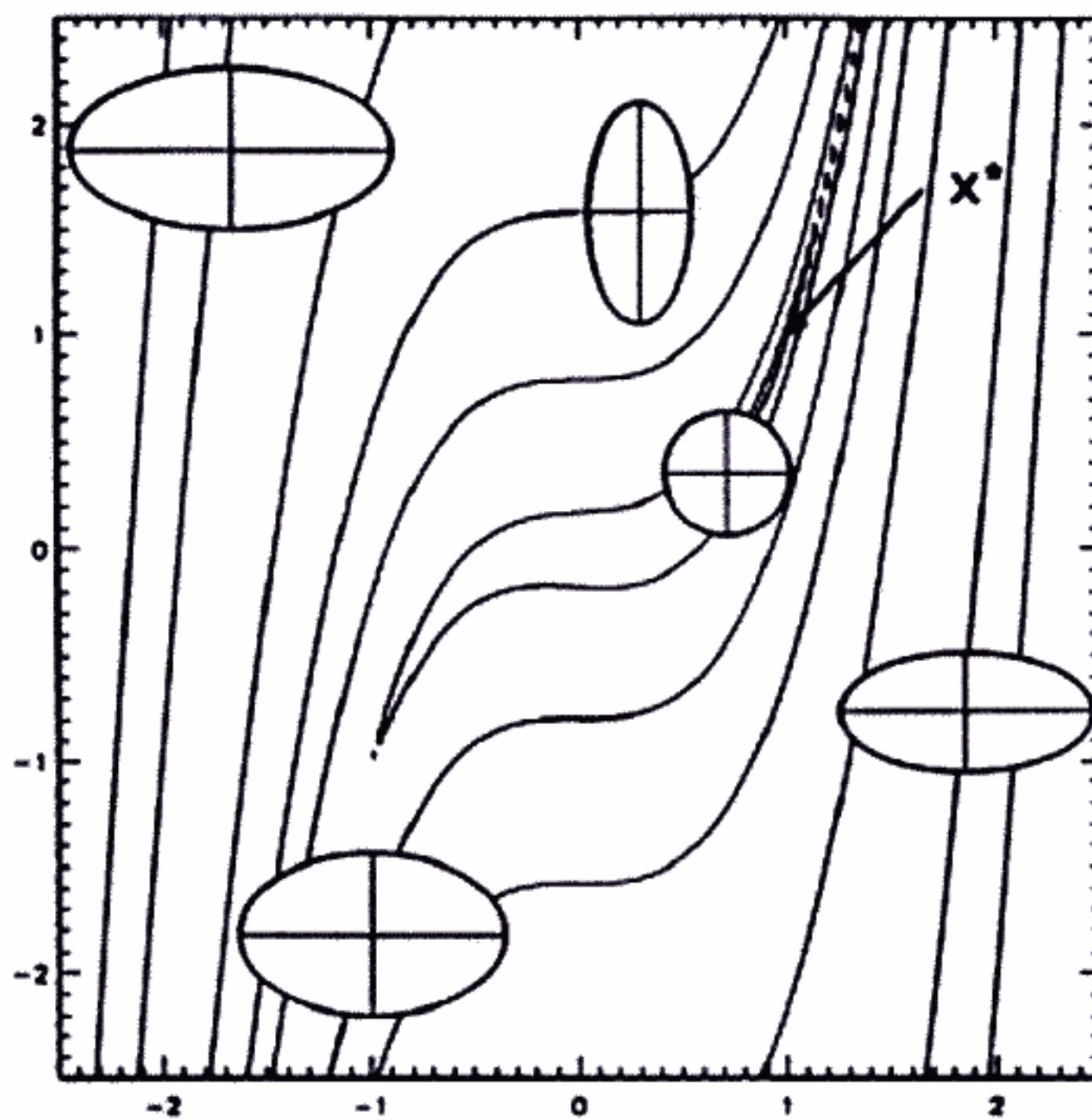


Figure 8: Simple mutations, $n = 2$, $n_\sigma = 2$, ($n_\sigma = 0$).

$$I = \mathbb{R}^n \times \mathbb{R}_+^n$$

$$m'_{\{\tau, \tau'\}}(\vec{x}, \vec{\sigma}) = (\vec{x}', \vec{\sigma}')$$

$$\tau \sim 1/\sqrt{2\sqrt{n}} \cdot c_{\mu, \lambda, \varsigma, \kappa}$$

$$\tau' \sim 1/\sqrt{2n} \cdot c_{\mu, \lambda, \varsigma, \kappa}$$

$$\sigma'_i = \sigma_i \cdot \exp(\tau' \cdot N(0, 1) + \tau \cdot N_i(0, 1))$$

$$x'_i = x_i + \sigma'_i \cdot N_i(0, 1)$$

Boundary rule for preserving standard deviations larger than zero:

$$\sigma'_i < \varepsilon_\sigma \Rightarrow \sigma'_i := \varepsilon_\sigma$$

Correlated Mutations

\oplus equal probability to place an offspring

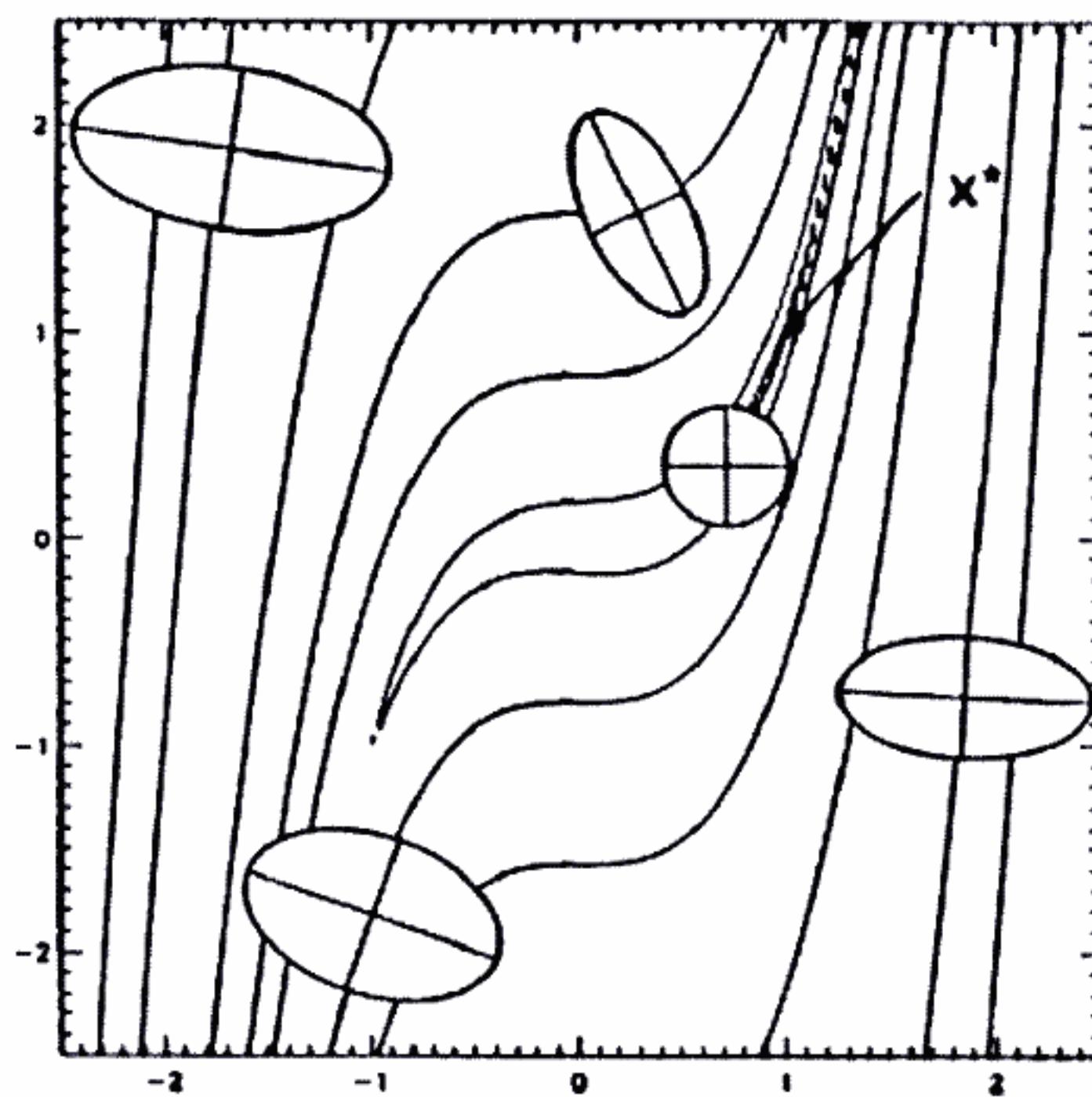


Figure 9: Correlated mutations, $n = 2$, $n_x = 2$, $n_\alpha = 1$.

$$\begin{aligned}
 I &= IR^n \times IR_+^n \times [-\pi, \pi]^{n \cdot (n-1)/2} \\
 m'_{\{\tau, \tau' \beta\}}(\vec{x}, \vec{\sigma}, \vec{\alpha}) &= (\vec{x}', \vec{\sigma}', \vec{\alpha}') \\
 \tau &\sim 1/\sqrt{2\sqrt{n}} \\
 \tau' &\sim 1/\sqrt{2n} \\
 \beta &\approx 5^\circ
 \end{aligned}$$

$$\begin{aligned}
 \sigma'_i &= \sigma_i \cdot \exp(\tau' \cdot N(0, 1) + \tau \cdot N_i(0, 1)) \\
 \alpha'_j &= \alpha_j + \beta \cdot N_j(0, 1) \\
 \vec{x}' &= \vec{x} + \vec{N}(\vec{0}, \mathbf{C}'')
 \end{aligned}$$

Boundary rule for keeping rotation angles feasible:

$$|\alpha'_j| > \pi \Rightarrow \alpha'_j := \alpha'_j - 2\pi \cdot \text{sign}(\alpha'_j)$$

⇒ A strategy parameter set

- is part of each individual,
- represents the p.d.f. for mutation of the individual:

$$p(\vec{z}) = \sqrt{\frac{\det C}{(2\pi)^n}} \exp\left(-\frac{1}{2}\vec{z}^T C \vec{z}\right) .$$

- C^{-1} : Covariance matrix:

$$\begin{aligned} c_{ii} &= \sigma_i^2 \\ c_{ij}, (i \neq j) &= \begin{cases} 0 & , \text{ no correlations} \\ \frac{1}{2}(\sigma_i^2 - \sigma_j^2) \tan(2\alpha_{ij}) & , \text{ correlations} \end{cases} \end{aligned}$$

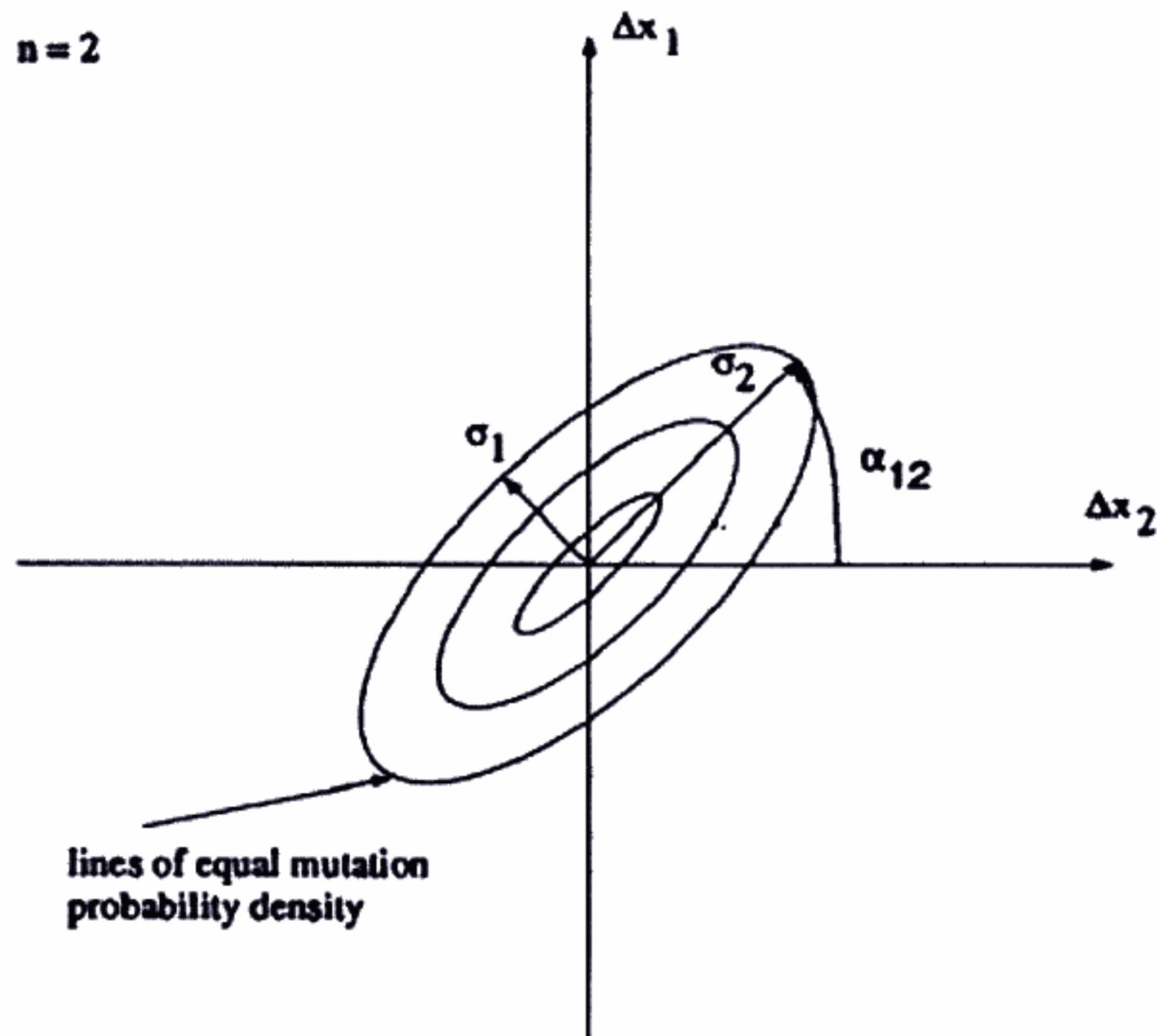


Figure 5: Illustration of the mutation ellipsoid for the case $n = 2$, $n_s = 2$, $n_a = 1$.

Structure of Individuals

- Individual space

$$I = \mathbb{R}^n \times \mathcal{S}$$

- Strategy parameters

$$\mathcal{S} = \mathbb{R}_+^{n_\sigma} \times [-\pi, \pi]^{n_\alpha}$$

$$\vec{a} = (\underbrace{(x_1, \dots, x_n)}_{\vec{x}}, \underbrace{(\sigma_1, \dots, \sigma_{n_\sigma})}_{\vec{\sigma}}, \underbrace{(\alpha_1, \dots, \alpha_{n_\alpha})}_{\vec{\alpha}}) \in I$$

- | | | | |
|----------------|-----------------------|---------------|----------------------|
| \vec{x} | : Object variables | \Rightarrow | Fitness $f(\vec{x})$ |
| $\vec{\sigma}$ | : Standard deviations | \Rightarrow | Variances |
| $\vec{\alpha}$ | : Rotation angles | \Rightarrow | Covariances |

n_σ	n_α	Remark
1	0	standard mutations
n	0	standard mutations
n	$n \cdot (n - 1)/2$	correlated mutations
$1 \leq n_\sigma \leq n$	$(n - \frac{n_\sigma}{2})(n_\sigma - 1)$	general case (correlated mutations)

Table 1: Possible settings of n_σ and n_α .

- If $1 < n_\sigma < n$: All x_i ($i > n_\sigma$) are mutated according to σ_{n_σ} .
- E.g.: $n_\sigma = 2$, $n_\alpha = n - 1$ facilitates learning of one preference direction.

Some remarks:

- Biological model:
Repair enzymes, mutator genes.
- There exists *no deterministic control*.
Instead, strategy parameters evolve as object variables do.
- There is only an *indirect link* between fitness and useful strategy parameter settings.
- $\vec{\sigma}$, $\vec{\alpha}$ are conceivable as an *internal model* of the local topology.
- Standard strategy: $n_\sigma = n$, $n_\alpha = 0$.
- For correlated mutations:
 - $\vec{\sigma}_c \sim \tilde{N}(\vec{0}, \mathbf{C})$ is generated by a multiplication of the uncorrelated random vector $\vec{\sigma}_u$ by n_α rotation matrices (Schwefel 1981, Rudolph 1992).

$$\vec{\sigma}_c = \prod_{i=1}^{n-1} \prod_{j=i+1}^n \mathbf{R}(\alpha_{ij}) \cdot \vec{\sigma}_u .$$

- Exactly the feasible (positive definite) correlation matrices \mathbf{C} can be created this way (Rudolph 1992).

$\text{rec} : I^\mu \rightarrow I$, $\text{rec} = \text{re} \circ \text{co}$

- Creates one individual per application.
- Works according to:

– $\text{co} : I^\mu \rightarrow I^\varrho$

Chooses ϱ parents at random.

e.g. $\varrho = 2$

– $\text{re} : I^\varrho \rightarrow I$

Creates one offspring individual.

- Common cases:

– $\varrho = 2$ bisexual recombination.

– $\varrho = \mu$ global recombination ("gene pool recombination").

Recombination types:

- No recombination, i.e., rec performs just a random choice of an individual.
- Global intermediary recombination:

$$b'_i = \frac{1}{\varrho} \sum_{k=1}^{\varrho} b_{k,i}$$

(Averaging over all parents).

- Local intermediary recombination:

$$b'_i = u_i b_{k_1,i} + (1 - u_i) b_{k_2,i}$$

$u_i \sim U([0, 1])$ or $u_i = 1/2$

$k_1, k_2 \sim U(\{1, \dots, \varrho\})$ for each offspring.

- Discrete recombination:

$$b'_i = b_{k_i,i}$$

$k_i \sim U(\{1, \dots, \varrho\})$ at random for each i .

b stands for $\{x, \varsigma, \alpha\}$

Recombination example

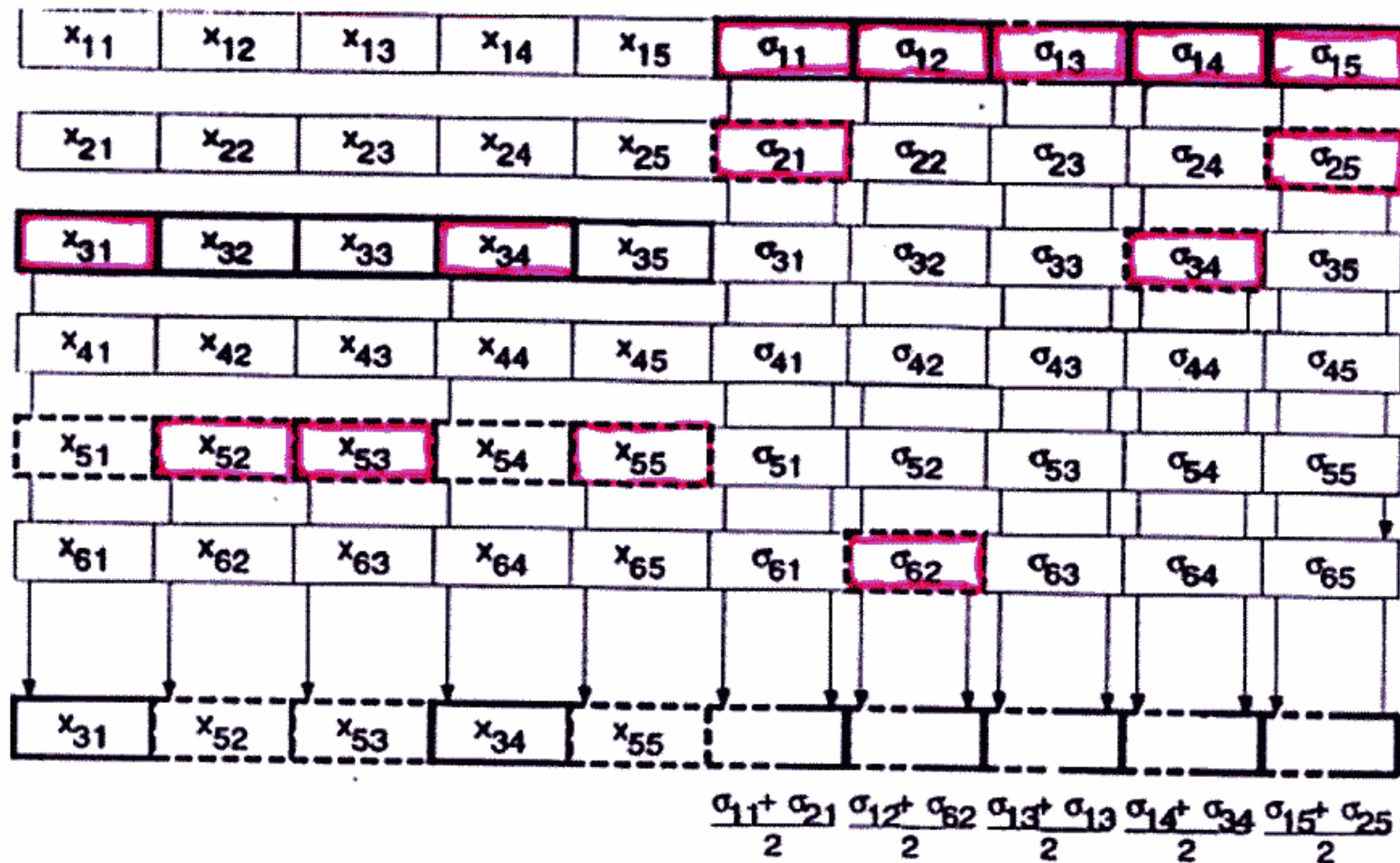


Figure 1: Recombination example for $n = 5$, $\mu = 6$, discrete recombination on x_i and global intermediate recombination on σ_i . The process is performed λ times for the creation of a complete offspring population.

Recombination type illustration

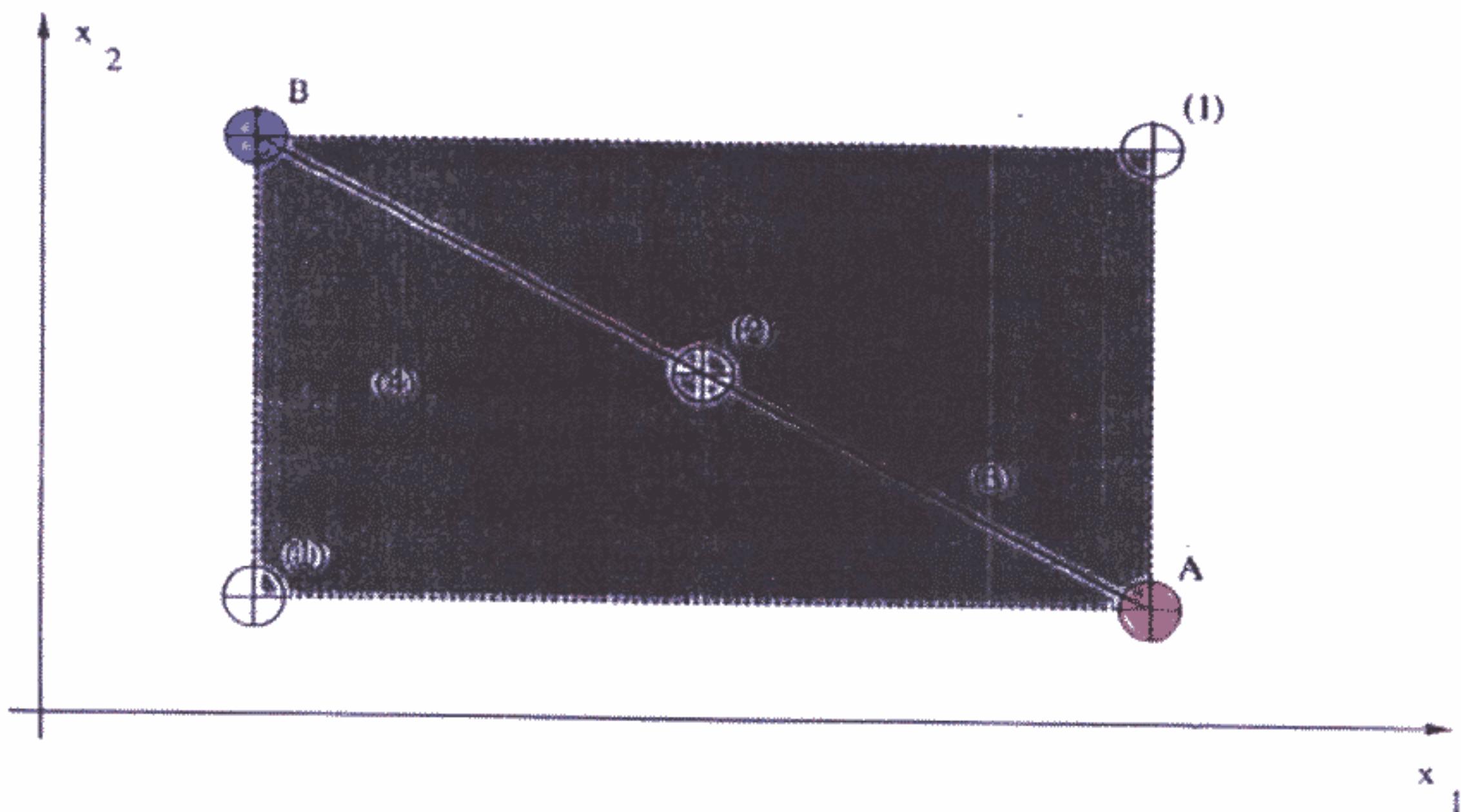


Figure 2: Illustration of the possible locations of recombination results according to the selected recombination scheme ($n = 2$). Notice that global random intermediate recombination is the most general mechanism.

Selection

- Strictly deterministic, rank-based.
- The μ best ranks are handled equally.
- Two forms:

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

The μ best of offspring and parents survive.

- (μ, λ) -selection:

The μ best offspring survives.

Formally: $\text{sel}_{\mu}^k(P) = \tilde{P}$, where $|\tilde{P}| = \mu$, $|P| = k \geq \mu$, and

$$\forall \vec{x} \in \tilde{P} : \forall \vec{a} \in P - \tilde{P} : f(\vec{x}) \leq f(\vec{a})$$

- Default: (μ, λ) -selection (e.g., (15,100)).

- Important for self-adaptation.

- Applicable also for noisy objective functions, moving optima.

- Selective pressure: Very high.

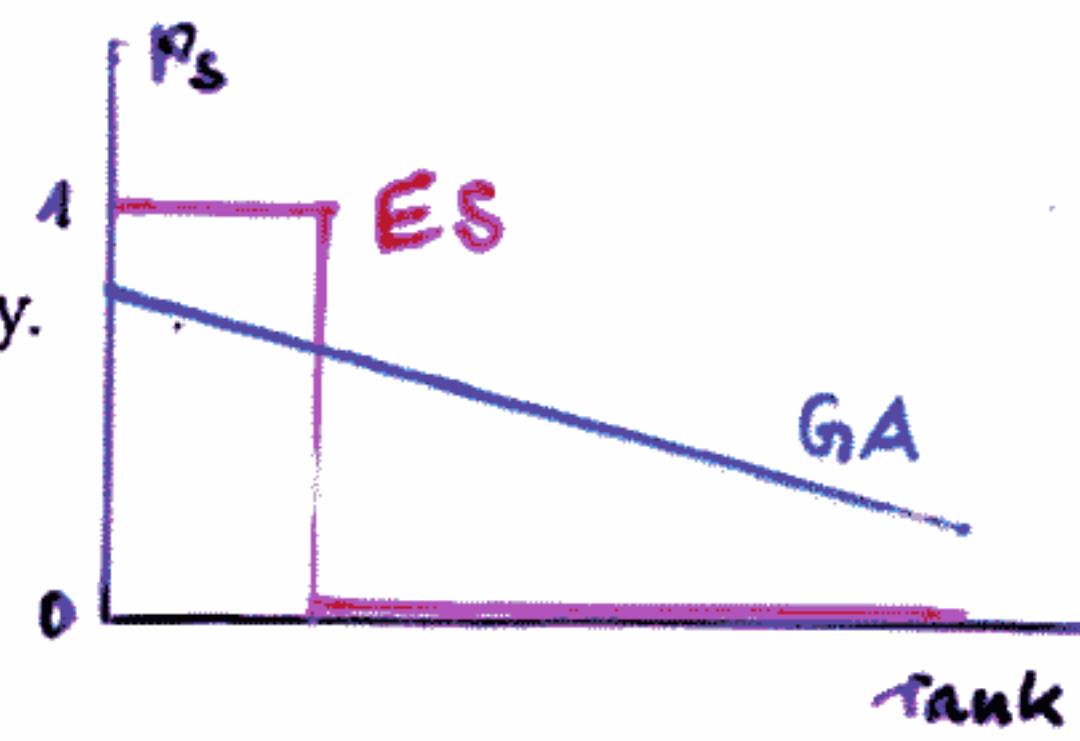
In terms of the *takeover time* τ^* (number of generations required until repeated application of selection completely fills the population with copies of the best individual; Goldberg & Deb 1991):

$$\tau^* = \frac{\ln \lambda}{\ln(\lambda/\mu)}$$

$\tau^* \approx 2$ generations for a (15,100)-ES

$\tau^* \approx 458$ for a (99,100)-ES

(cf. proportional selection in GAs: $\tau^* \approx \lambda \ln \lambda = 460$ generations!).



Self-Adaptation

1974/75

The critical claim (Schwefel 1987, 1992):

Self-adaptation of strategy parameters works

- without exogenous control,
- by recombining / mutating the strategy parameters,
- by exploiting the implicit link between fitness and useful internal model.

Necessary conditions:

- Generation of a surplus $\lambda > \mu$,
- (μ, λ) -selection (to guarantee extinction of misadapted individuals),
- a not too strong selective pressure (e.g., $(15, 100)$);
 $\lambda/\mu \approx 7$ turns out to be useful, but μ has to be clearly larger than 1),
- recombination also on strategy parameters
 (especially: intermediate recombination).

An illustration of self-adaptation:

Sphere model, (15,100)-ES with $n = 30$, $n_\sigma = 1$, and the optimum location is changed every 200 generations.

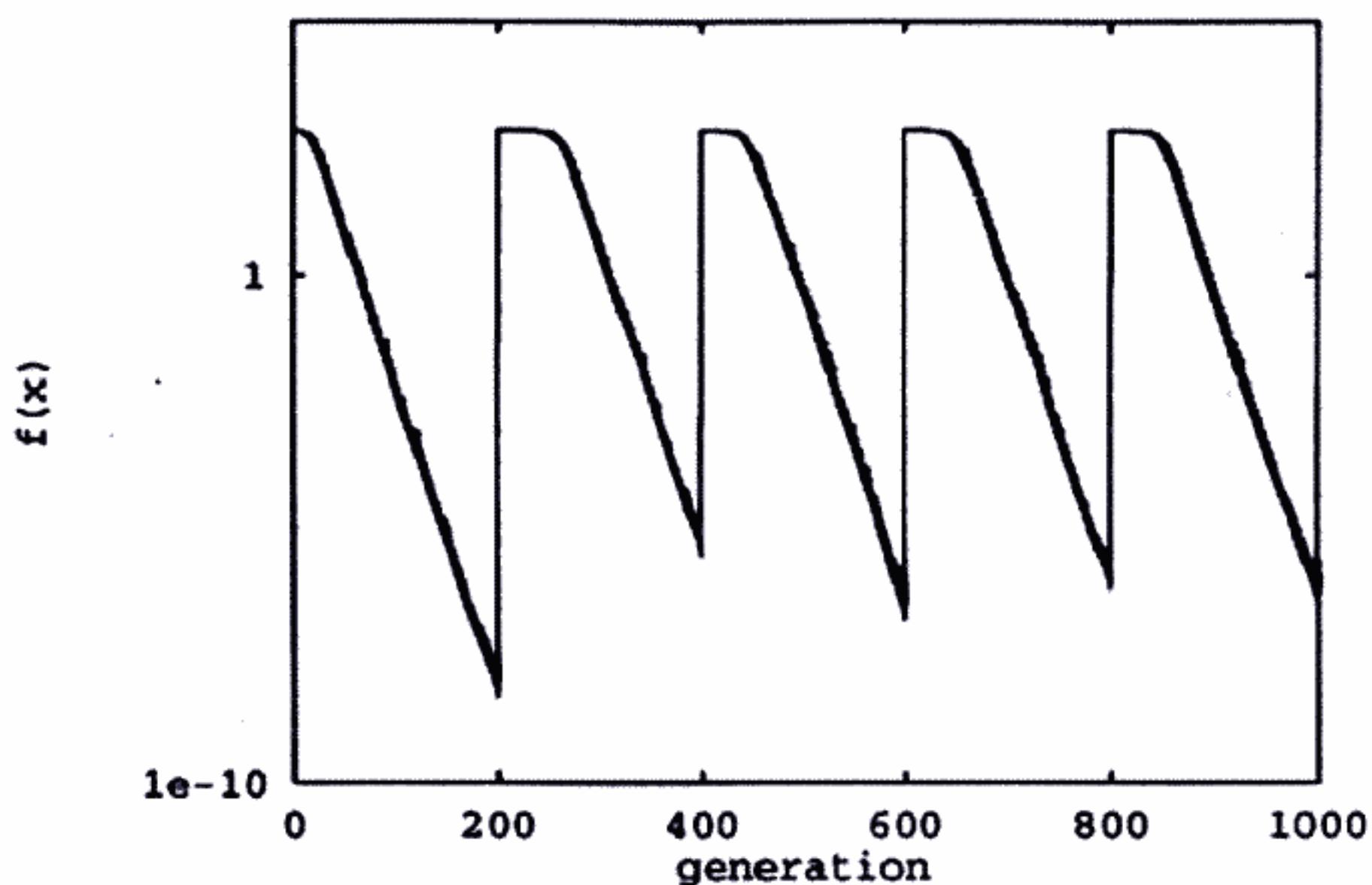


Figure 12: Behaviour of the objective function value.

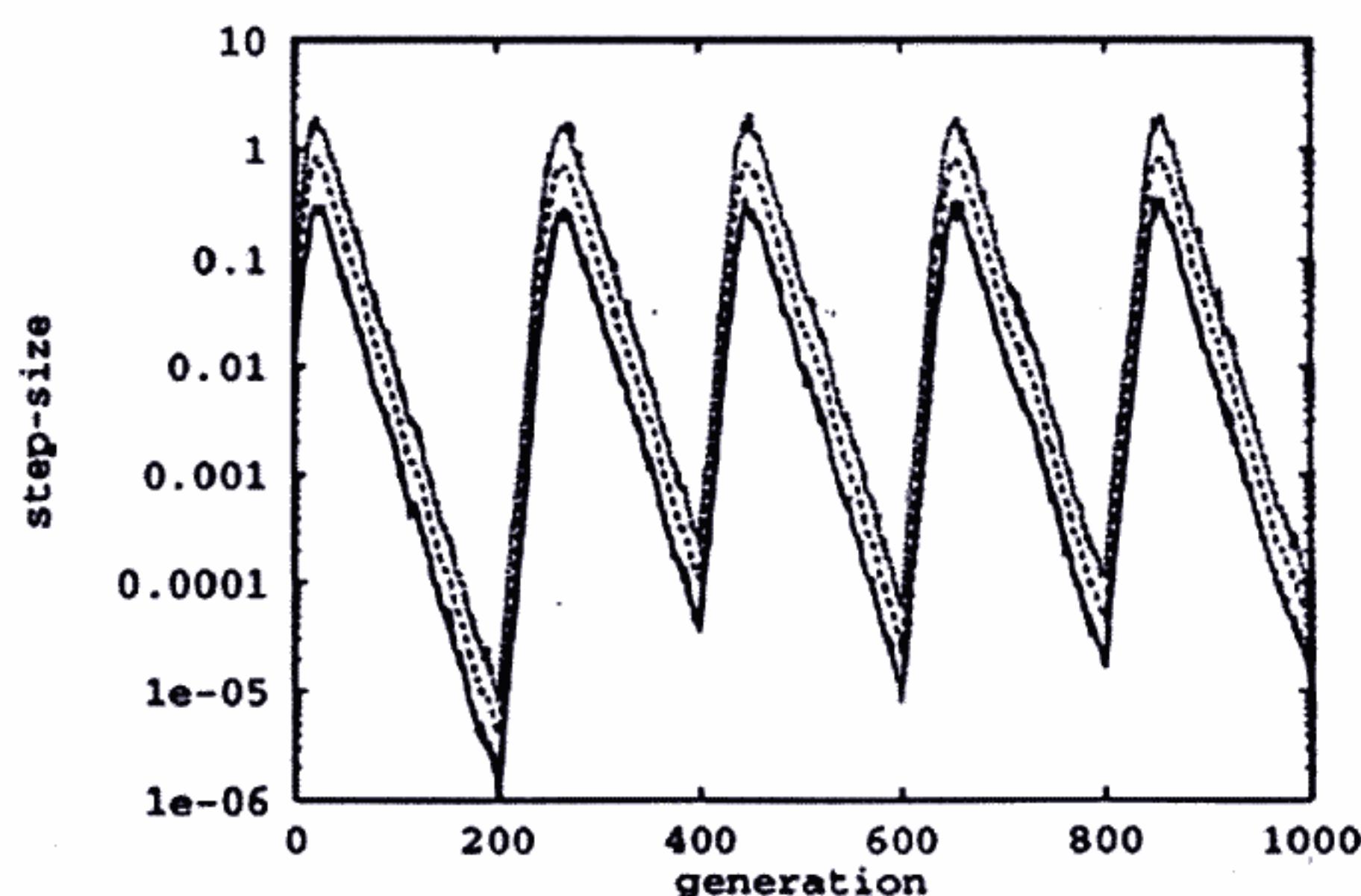


Figure 13: Behaviour of the standard deviations (minimum, average, and maximum within the population).

Collective learning ($n = 30$)

- of one common step size ($\Rightarrow 1$ step size):

$$f_1(\vec{x}) = \sum_{i=1}^n x_i^2$$

- of proper scalings ($\Rightarrow n$ step sizes):

$$f_2(\vec{x}) = \sum_{i=1}^n i \cdot x_i^2$$

- of a metric (\Rightarrow correlated mutations):

$$f_3(\vec{x}) = \sum_{i=1}^n \left(\sum_{j=1}^i x_j \right)^2$$

(Schwefel, 1987) compares the progress rate for f_2 and a $(\mu, 100)$ -ES for

- optimum prefixed scaling, i.e., *perfect information* ($\sigma_i = c/\sqrt{i}$) (A),
- prefixed arbitrary scaling ($\sigma_i = c'$) (B),
- adaptive scaling (C).

$$F(x) = \sum_{i=1}^{n=30} i \cdot x_i^2$$

$(\mu, 100) \in S$

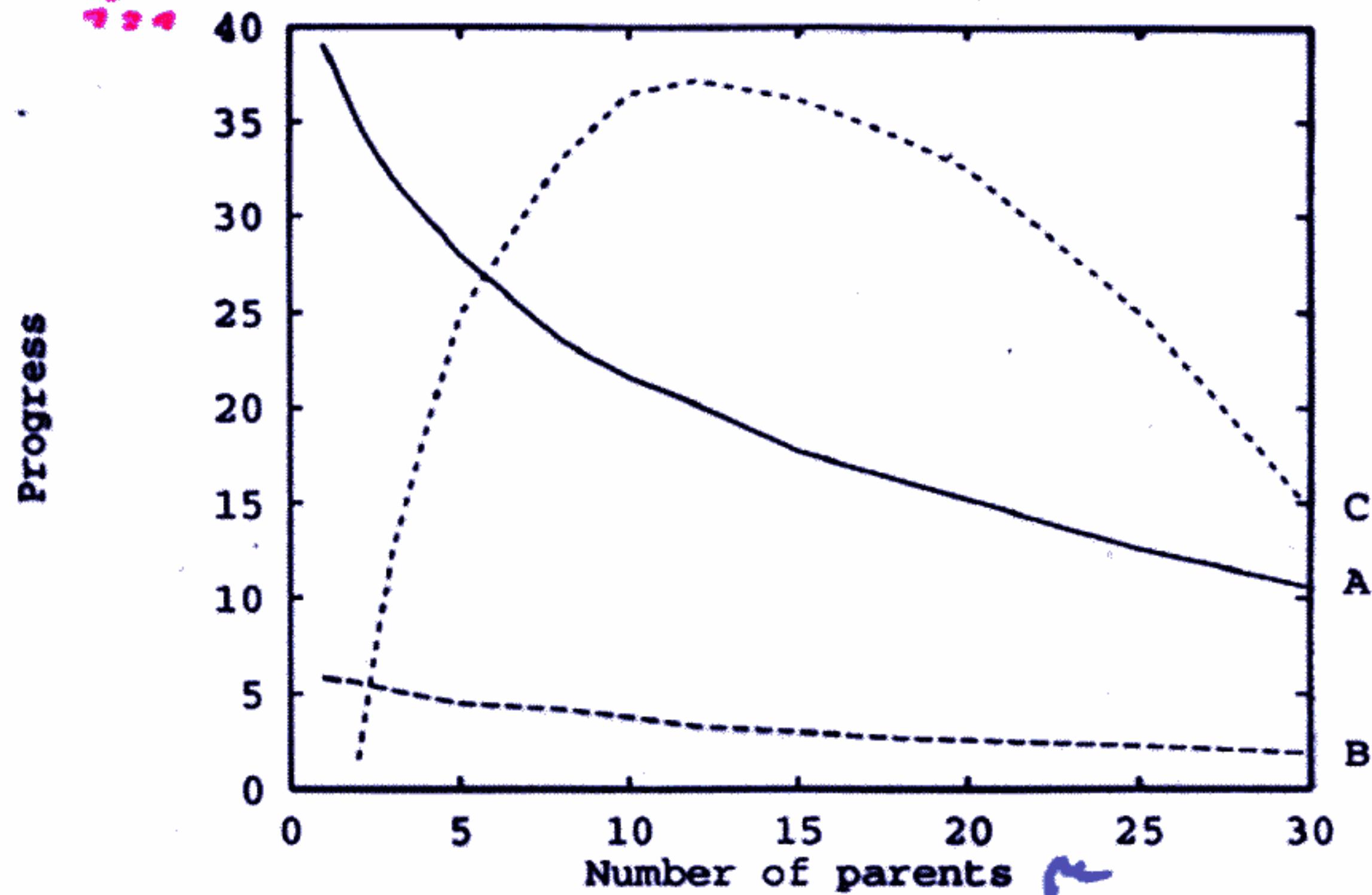


Figure 14: Comparison of progress rates. 15 imperfect, but different individuals (C) perform collectively better than the same number of cloned specialists (A) and almost as good as possible in case of perfect information (A, (1,100)).

Progress measure:

$$P = \ln \sqrt{\frac{f_0^{\min}}{f_{1000}^{\min}}}$$

Outlook: “Contemporary ESs”

Extensions of the $(\mu, \kappa, \lambda, \rho)$ -ES:

- Life span limited to $\kappa \geq 1$ generations.
 - $\kappa = 1$: (μ, λ) -selection.
 - $\kappa = \infty$: $(\mu + \lambda)$ -selection.
- Tournament selection as an alternative method.
- Further recombination types (e.g., crossover from GAs).
- Introduction of operator application probabilities p_m, p_c .

other ES variants

- with diploidy / dominance + recessivity
for vector optimization (MCDM)
- with niching / migration
for multimodal optimization (coarse grained)
- with local interaction (diffusion)
for multimodal optimization (fine grained)
- with predator-prey selection
for multimodal optimization (tournament-like)
- nested ESSs [Achenberg & Co.]

$[\mu'/s', \lambda' \underbrace{(\mu/s, \lambda)}^v]$ ES

λ' subpopulations (demes) act in parallel
for v generations independently

$\underbrace{\text{Intra-selection among } \lambda' \text{ subpopulations}}$

(μ, λ) ES good for dynamic Opt.
(changing environment)

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Strategien - Test

1. Konvergenzgeschwindigkeit

bzw. benötigte Rechenzeit (n) für geg. Approx.
zwei quadratische Probleme

a) $F = \sum_{i=1}^n x_i^2$

b) $F = \sum_{i=1}^n \left(\sum_{j=1}^i x_j \right)^2$

$$= x^T A x$$

↑ Kondition der Matrix $\sim n^2$

2. Zuverlässigkeit

erreichbare Approximation

50 Testprobleme mit $n \leq 6$

28 ohne Nebenbedingungen

22 mit Nebenbedingungen

8 davon multimodal

getestete Strategien

Koordinatenstrategie mit linearer Suche

Fibonacci:

FIBO

Gold. Schnitt

GOLD

Lagrange

LAGR

Hooke + Jeeves

H03E

Rosenbrock

ROSE

Davies, Swann, Lampsey

DSCG / DISC

Powell (konjug. R:)

POWE

variable Metrik

DFPS

Simplex

SIMP

Complex

COMP

Evolutionsstrategien

(1+1)

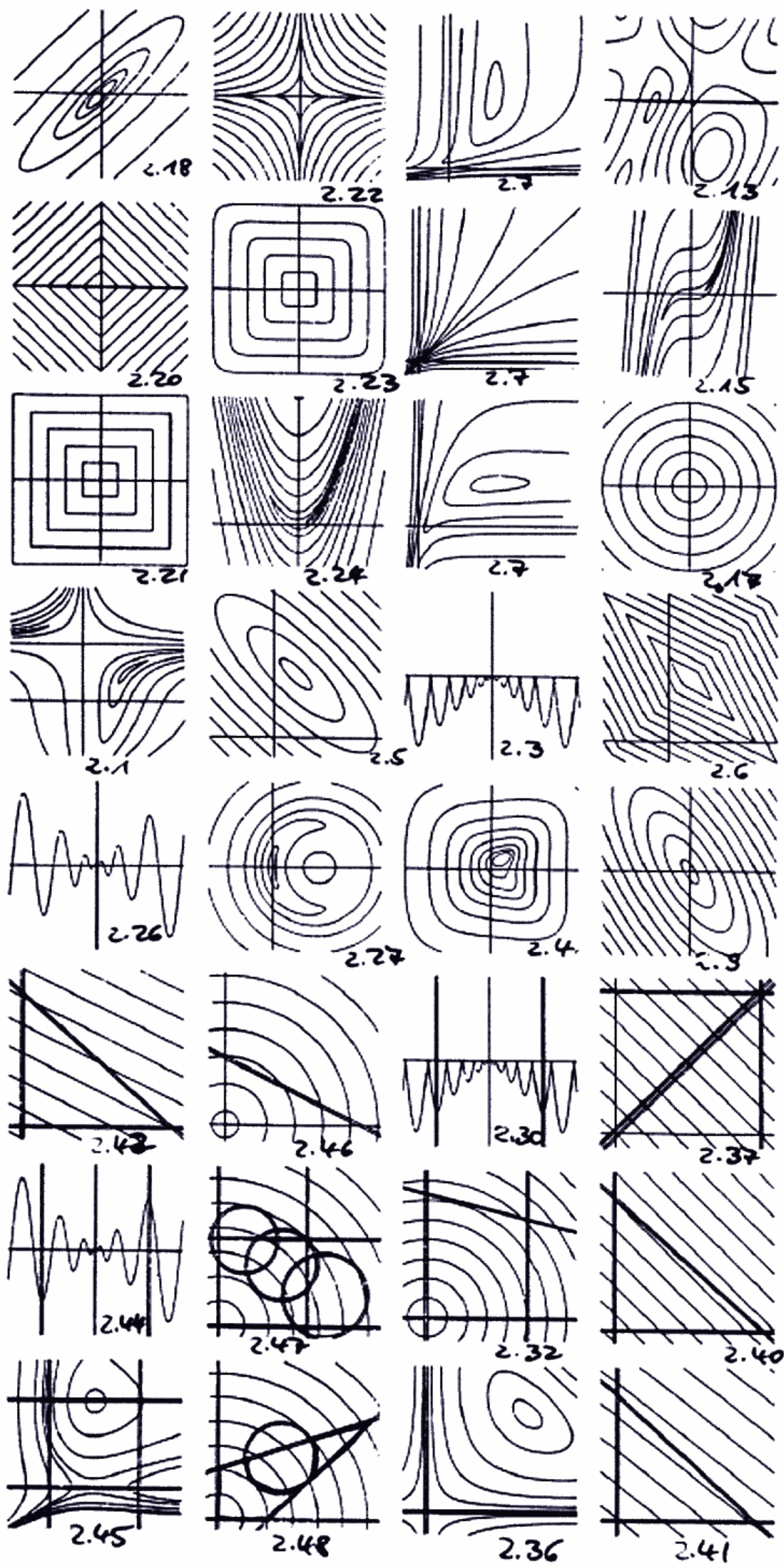
EVOL

(10,100)

GRUP

(10,100) mit Rekomb.

REKO



e: fataler Fehler

a: schaltet nicht ab

r: sehr langsam;
Zeitabstaltung

h: Neustart

1 Schwierigkeit
Bosser 10^{-38}

2 ~ 10^{-8}

3 ~ 10^{-4}

4 ~ 10^{-2}

5 abgelenkt 10^{-2}

Kennzeichen

Problem Nr.

	Strategie	ROSE	COMP	EVOL	GRUP	REKO
2.29	3	1	4	3	3	3
2.30	1	5	1	1	1	1
2.31	3v	3	1	1	1	1
2.32	3v	3	1	1	1	1
2.33	3	2	5	4	4	1
2.34	1	2	3	3	3	2
2.35	3v	1	4	4	4	4
2.36	1	1	1	1	1	1
2.37	3	1	1	1	1	1
2.38	3v	3	1	1	1	1
2.39	3	3	4	4	4	3
2.40	5	5	5	5	5	5
2.41	5	5	5	5	5	5
2.42	3	3	2	2	2	1
2.43	3v	3	2	2	2	1
2.44	1	5	1	1	1	1
2.45	3	2	4	2	2	1
2.46	3	2	3	3	3	1
2.47	3v	1	1	1	1	1
2.48	3v	3	1	1	1	1
2.49	3	2	4	3	3	1
2.50	3	1	1	1	1	1

Strategie

Anzahl der

	exakt	nicht gelösten Aufgaben	exakt	nicht
	ohne	mit	ohne	mit
Nebenbedingungen				
FIBO	3	9		
GOLD	4	9		
LAGR	2	7		
HOJE	6	2		
DSCG	11	2		
DSCP	12	2		
POWE	4	7		
DFPS	5	6		
SIMP	7	2		
ROSE	11	2	4	2
COMP	5	2	6	4
EVOL	17	0	10	3
GRUP	18	0	10	2
REKO	23	0	16	2
			Von 28	Von 22

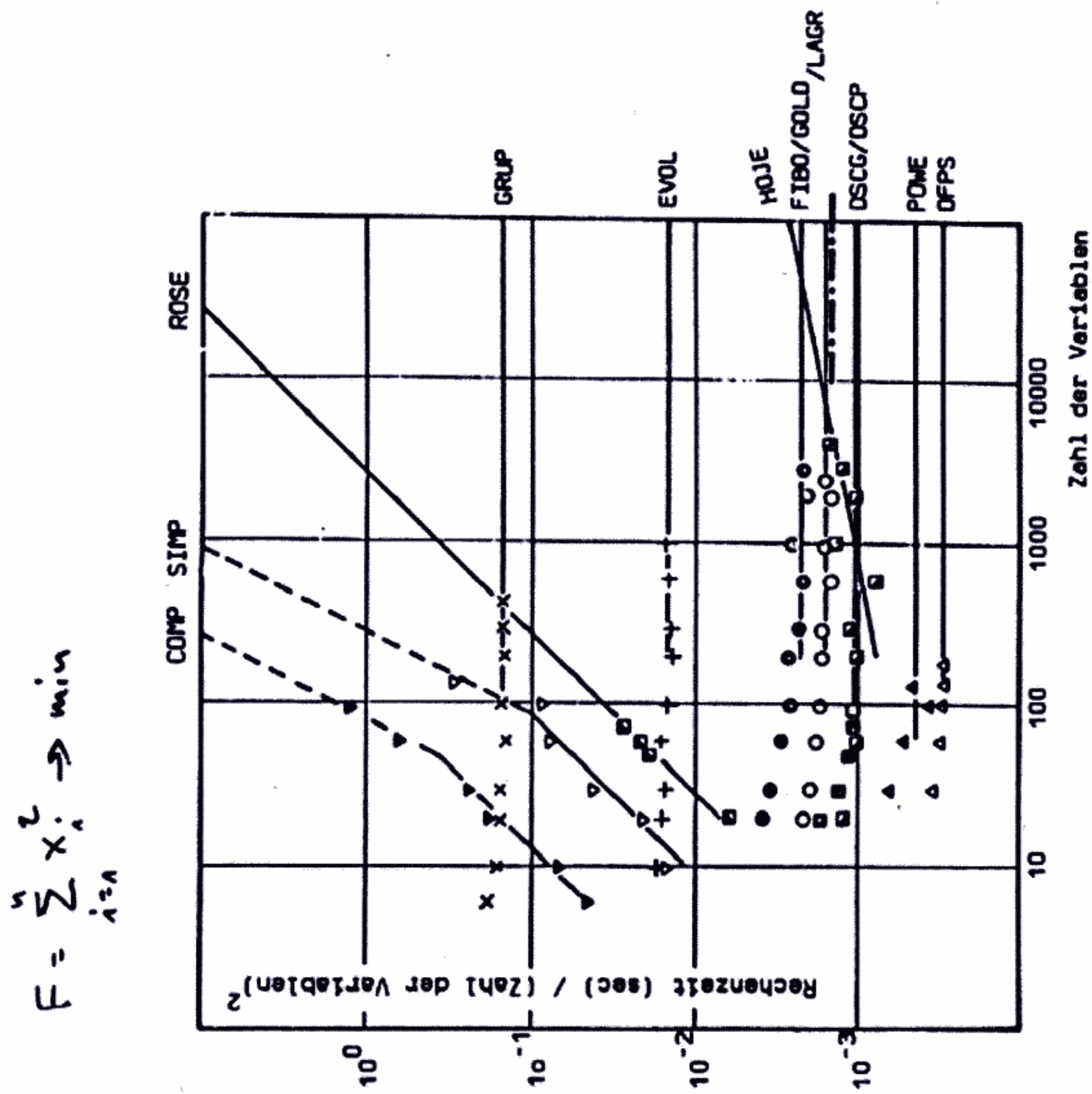
Zusammengefaßte Ergebnisse des Zuverlässigkeitstests im Hinblick auf lokale Konvergenz (28 Probleme ohne und 22 mit Nebenbedingungen; Strategie-Namen wie in Tabelle 1)

<u>Strategie</u>	Problem Nr.	FIBO	GOLD	LAGR	HOLE	SIMP	DFPS	POWE	DSCP	DSCG	ROSE	COMP	EVOL	GRUP	REKO
1		L1	L1	L3	L1	L7	L1	L3	L1	L6	L1	Lm	G1	G1	G1
2		L1	G1	G1	G1										
3									L4	L1	Lm				GL
4												G	G	G	
5											L1	L1	L1		
6											L	G	G		
7											L3	L1	L2		
8											L2	Lm	Lm		

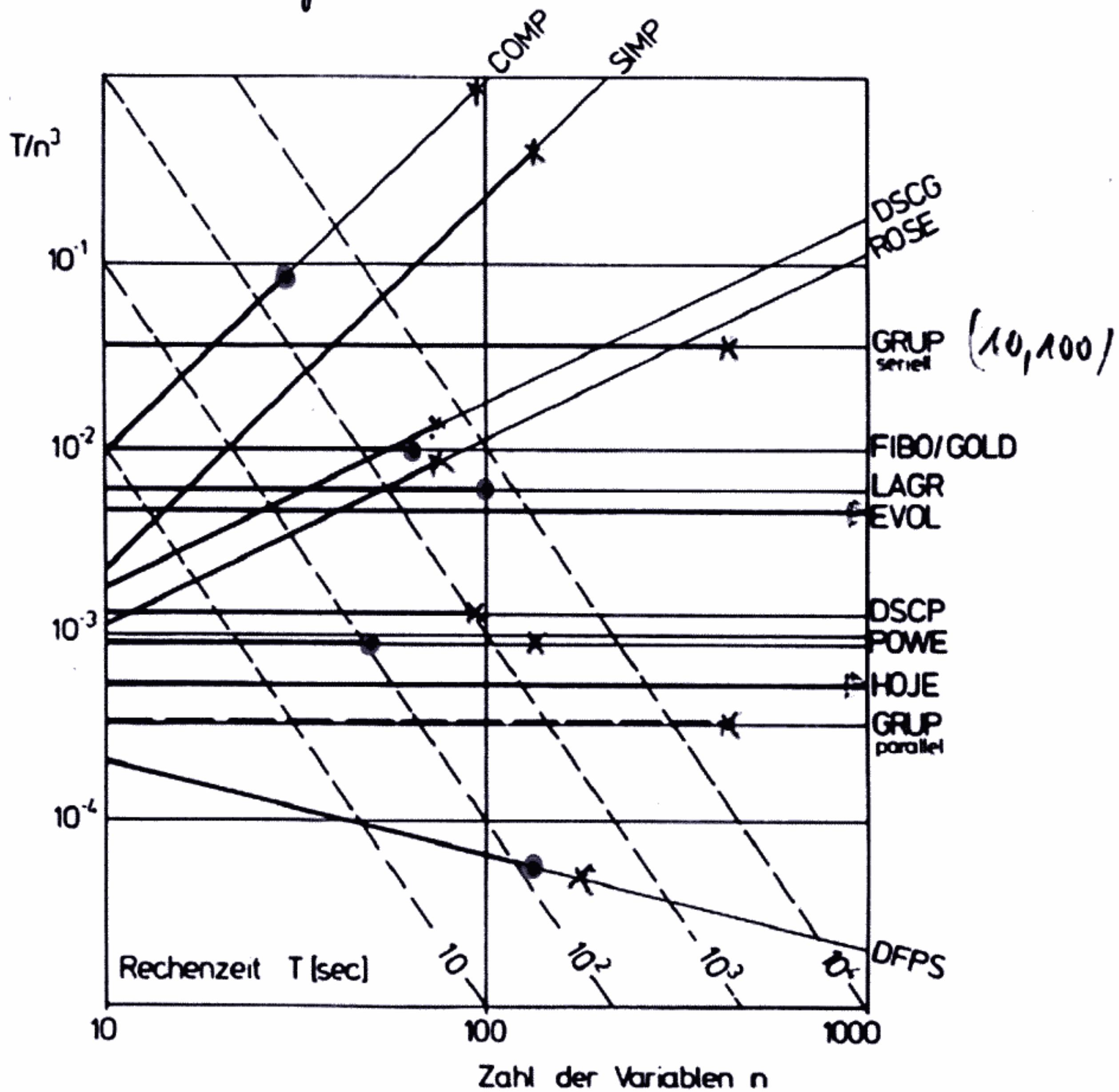
Ergebnisse des Tests auf globale Konvergenz
(Strategienamen wie in Tabelle 1)

Bedeutung der Symbole:

- L Suche konvergiert gegen lokales Minimum
- L3 Suche konvergiert gegen das 3. lokale Minimum
(Zählung der Reihenfolge nach absteigenden Zielfunktionswerten)
- Lm Suche konvergiert gegen verschiedene lokale Minima je nach Zufallszahlen
- G Suche konvergiert gegen globales Minimum
- GL Suche konvergiert gegen lokales oder globales Minimum je nach Zufallszahlen



$$F = \sum_{i=1}^m \left(\sum_{j=1}^n x_j \right)^2 \rightarrow \min$$



● fatale Fehler

✗ Speicherplatzbegrenzung

Benötigte Rechenzeit in Abhängigkeit von der Parameterzahl für die Erzielung einer vorgegebenen Approximation an die Lösung eines quadratischen Optimierproblems

Bedeutung der Symbole:

○ Strategie versagt bei höherer Parameterzahl

✗ Kernspeicher-Platzbedarf überschreitet 30 K Worte