

# Covariance Matrix Adaptation Evolution Strategy (CMA-ES)

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Inria

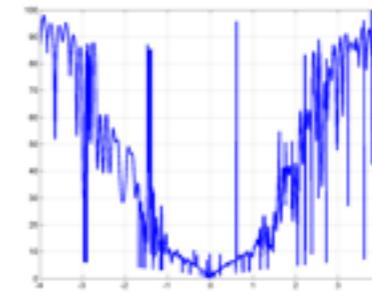
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# Problem Statement: Black-Box Optimization

Given an objective function

$$f : \mathcal{X} \subset \mathbb{R}^n \rightarrow \mathbb{R}$$

Minimize  $f$  in a *black-box scenario* (direct search, no gradients)



Problem domain specific knowledge is only used *within* the black-box

## *Objective*

- convergence to a global essential infimum of  $f$  as fast as possible  
linear convergence,  $\mathcal{O}(n \log 1/\epsilon)$  black-box evaluations
- find  $x \in \mathcal{X}$  with small  $f(x)$  value using as few black-box calls as possible

The black box can

- be non-convex, multi-modal/rugged, discontinuous, noisy, dynamic
- take from milli-seconds to hours to evaluate

# Stochastic Search

A black box search template to minimize  $f : \mathbb{R}^n \rightarrow \mathbb{R}$

Initialize distribution parameters  $\theta$ , set population size  $\lambda \in \mathbb{N}$   
While not terminate

- ① Sample distribution  $P(x|\theta) \rightarrow x_1, \dots, x_\lambda \in \mathbb{R}^n$
- ② Evaluate  $x_1, \dots, x_\lambda$  on  $f$
- ③ Update parameters  $\theta \leftarrow F_\theta(\theta, x_1, \dots, x_\lambda, f(x_1), \dots, f(x_\lambda))$

Everything depends on the definition of  $P$  and  $F_\theta$

deterministic algorithms are covered as well

In many Evolutionary Algorithms the distribution  $P$  is implicitly defined via **operators on a population**, in particular, selection, recombination and mutation

Natural template for (incremental) *Estimation of Distribution Algorithms*

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# Landscape of Continuous Search Methods

## *Gradient-based (Taylor, local)*

- Conjugate gradient methods [Fletcher & Reeves 1964]
- Quasi-Newton methods (BFGS) [Broyden et al 1970]

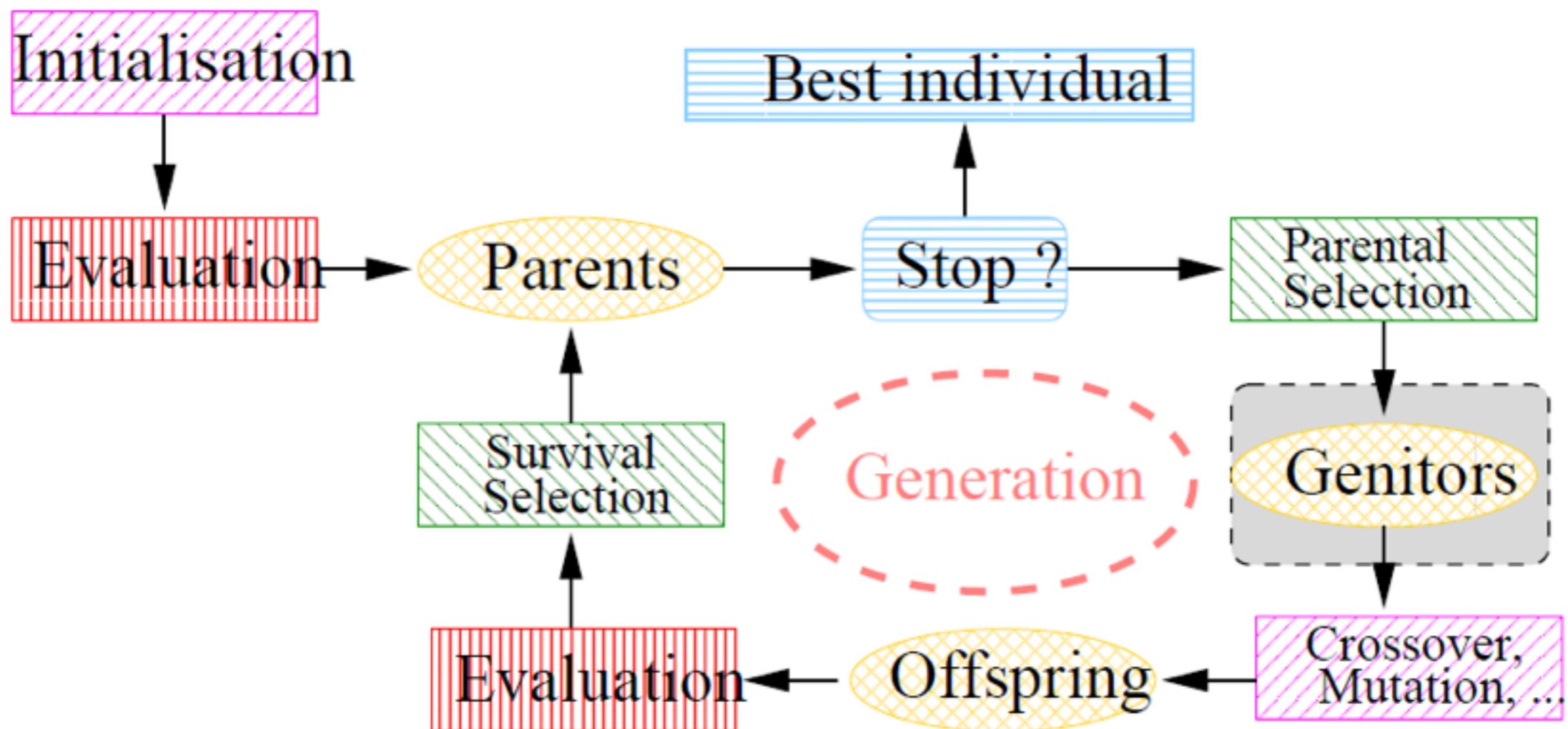
## *Derivative-free optimization (DFO)*

- Trust-region methods (NEWUOA, BOBYQA) [Powell 2006, 2009]
- Simplex downhill [Nelder & Mead 1965]
- Pattern search [Hooke & Jeeves 1961, Audet & Dennis 2006]

## *Stochastic (randomized) search methods*

- Evolutionary algorithms (broader sense, continuous domain)
  - Differential Evolution [Storn & Price 1997]
  - Particle Swarm Optimization [Kennedy & Eberhart 1995]
  - **Evolution Strategies** [Rechenberg 1965, Hansen & Ostermeier 2001]
- Simulated annealing [Kirkpatrick et al 1983]
- Simultaneous perturbation stochastic approximation (SPSA) [Spall 2000]

# A Different Viewpoint: Evolutionary Algorithm Scheme



- Stochastic operators Representation dependent
- Darwinian Evolution Engine (can be stochastic or deterministic)
- Main CPU cost
- Checkpointing: stopping criterion, statistics, updates, ...

# Metaphors

## Evolutionary Computation

## Optimization/Nonlinear Programming

individual, offspring, parent       $\longleftrightarrow$

candidate solution  
decision variables  
design variables  
object variables

population       $\longleftrightarrow$   
fitness function       $\longleftrightarrow$

set of candidate solutions  
objective function  
loss function  
cost function  
error function  
iteration

generation       $\longleftrightarrow$

... methods: ESs

# The CMA-ES

**Input:**  $\mathbf{m} \in \mathbb{R}^n$ ,  $\sigma \in \mathbb{R}_+$ ,  $\lambda$

**Initialize:**  $\mathbf{C} = \mathbf{I}$ , and  $\mathbf{p}_c = \mathbf{0}$ ,  $\mathbf{p}_\sigma = \mathbf{0}$ ,

**Set:**  $c_c \approx 4/n$ ,  $c_\sigma \approx 4/n$ ,  $c_1 \approx 2/n^2$ ,  $c_\mu \approx \mu_w/n^2$ ,  $c_1 + c_\mu \leq 1$ ,  $d_\sigma \approx 1 + \sqrt{\frac{\mu_w}{n}}$ ,  
and  $w_{i=1\dots\lambda}$  such that  $\mu_w = \frac{1}{\sum_{i=1}^\mu w_i^2} \approx 0.3 \lambda$

**While not terminate**

$$\mathbf{x}_i = \mathbf{m} + \sigma \mathbf{y}_i, \quad \mathbf{y}_i \sim \mathcal{N}_i(\mathbf{0}, \mathbf{C}), \quad \text{for } i = 1, \dots, \lambda \quad \text{sampling}$$

$$\mathbf{m} \leftarrow \sum_{i=1}^\mu w_i \mathbf{x}_{i:\lambda} = \mathbf{m} + \sigma \mathbf{y}_w \quad \text{where } \mathbf{y}_w = \sum_{i=1}^\mu w_i \mathbf{y}_{i:\lambda} \quad \text{update mean}$$

$$\mathbf{p}_c \leftarrow (1 - c_c) \mathbf{p}_c + \mathbf{1}_{\{\|\mathbf{p}_\sigma\| < 1.5\sqrt{n}\}} \sqrt{1 - (1 - c_c)^2} \sqrt{\mu_w} \mathbf{y}_w \quad \text{cumulation for } \mathbf{C}$$

$$\mathbf{p}_\sigma \leftarrow (1 - c_\sigma) \mathbf{p}_\sigma + \sqrt{1 - (1 - c_\sigma)^2} \sqrt{\mu_w} \mathbf{C}^{-\frac{1}{2}} \mathbf{y}_w \quad \text{cumulation for } \sigma$$

$$\mathbf{C} \leftarrow (1 - c_1 - c_\mu) \mathbf{C} + c_1 \mathbf{p}_c \mathbf{p}_c^T + c_\mu \sum_{i=1}^\mu w_i \mathbf{y}_{i:\lambda} \mathbf{y}_{i:\lambda}^T \quad \text{update } \mathbf{C}$$

$$\sigma \leftarrow \sigma \times \exp \left( \frac{c_\sigma}{d_\sigma} \left( \frac{\|\mathbf{p}_\sigma\|}{\mathbb{E}[\|\mathcal{N}(\mathbf{0}, \mathbf{I})\|]} - 1 \right) \right) \quad \text{update of } \sigma$$

**Not covered** on this slide: termination, restarts, useful output, boundaries and encoding

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Natural template for (incremental) *Estimation of Distribution Algorithms*

# Evolution Strategies

New search points are sampled normally distributed

$$x_i \sim \mathbf{m} + \sigma \mathcal{N}_i(\mathbf{0}, \mathbf{C}) \quad \text{for } i = 1, \dots, \lambda$$

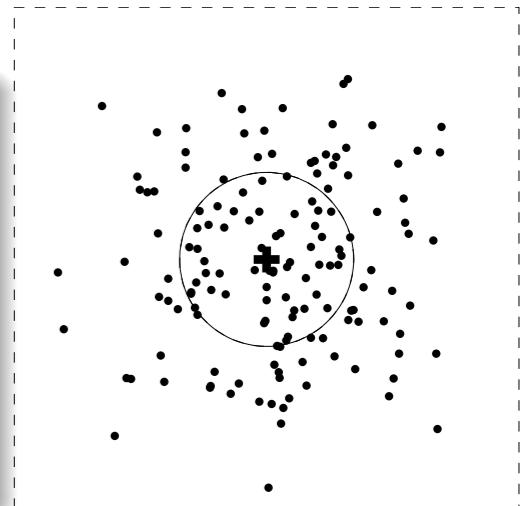
as perturbations of  $\mathbf{m}$ , where  $x_i, \mathbf{m} \in \mathbb{R}^n$ ,  $\sigma \in \mathbb{R}_+$ ,  $\mathbf{C} \in \mathbb{R}^{n \times n}$

where

- the **mean** vector  $\mathbf{m} \in \mathbb{R}^n$  represents the favorite solution
- the so-called **step-size**  $\sigma \in \mathbb{R}_+$  controls the *step length*
- the **covariance matrix**  $\mathbf{C} \in \mathbb{R}^{n \times n}$  determines the **shape** of the distribution ellipsoid

here, all new points are sampled with the same parameters

The question remains how to update  $\mathbf{m}$ ,  $\mathbf{C}$ , and  $\sigma$ .



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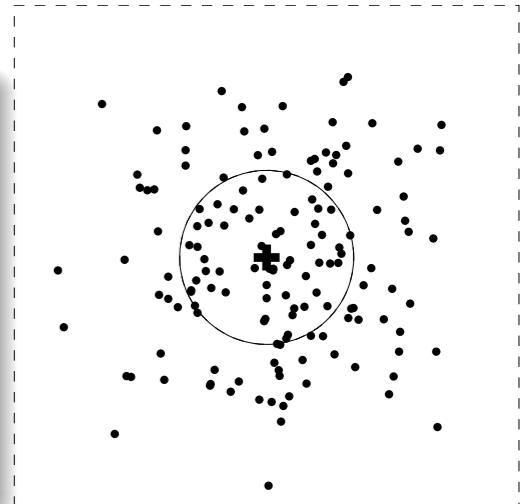
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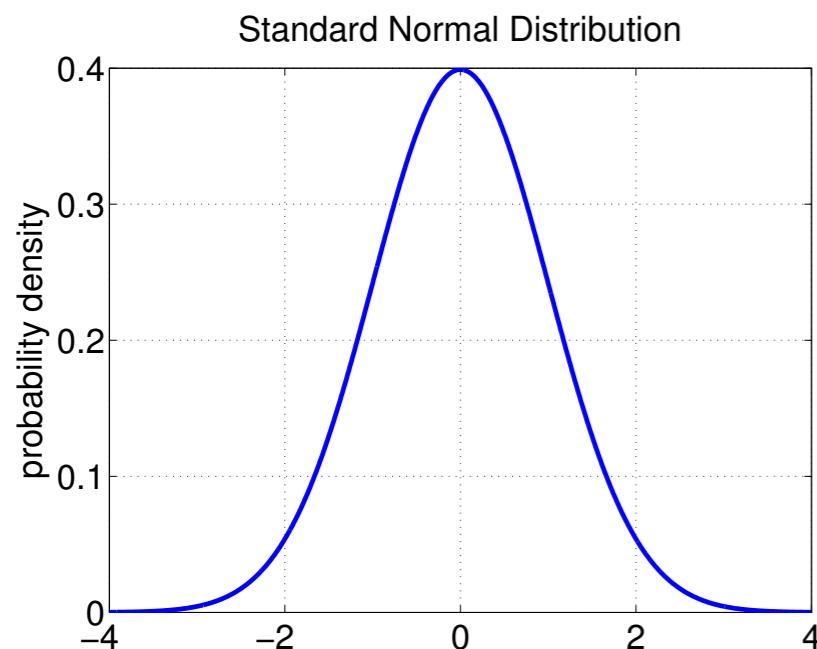


The (multivariate) normal distribution (Gaussian distribution)

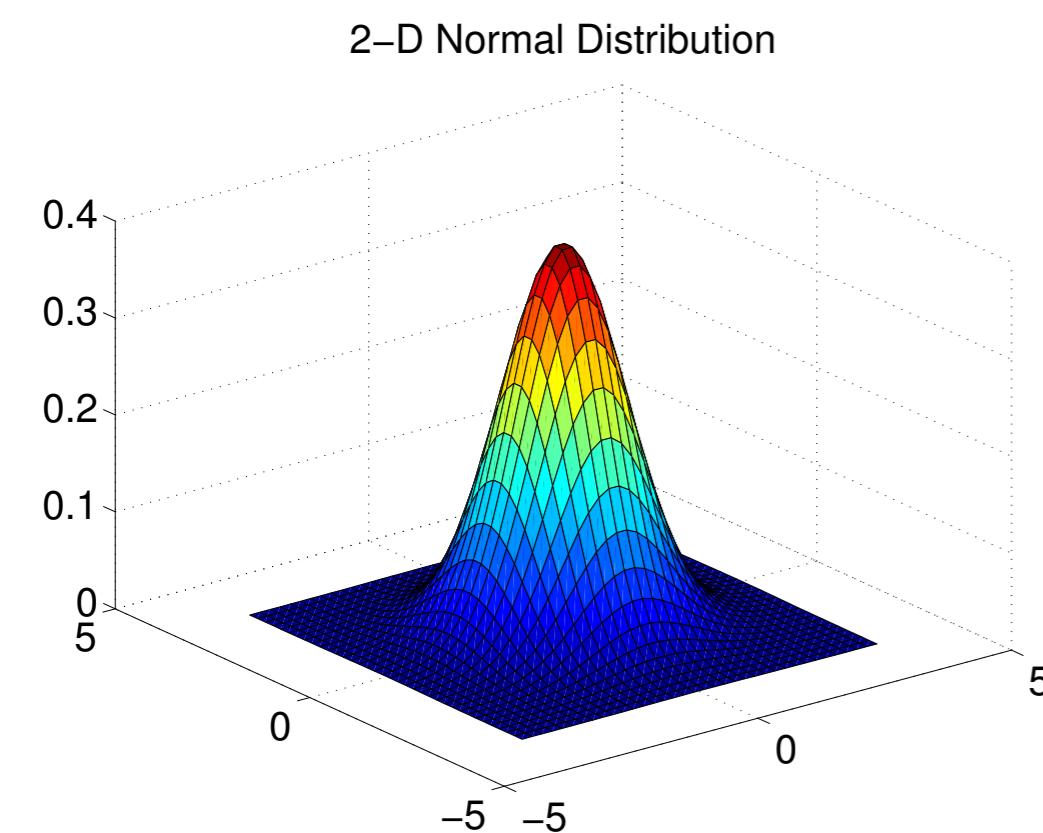
# Why Normal Distributions?

- ① widely observed in nature, for example as phenotypic traits
- ② only stable distribution with finite variance
  - stable means that the sum of normal variates is again normal:
$$\mathcal{N}(\mathbf{x}, \mathbf{A}) + \mathcal{N}(\mathbf{y}, \mathbf{B}) \sim \mathcal{N}(\mathbf{x} + \mathbf{y}, \mathbf{A} + \mathbf{B})$$
  - helpful in design and analysis of algorithms related to the *central limit theorem*
- ③ most convenient way to generate isotropic search points
  - the isotropic distribution does not favor any direction, rotational invariant
- ④ maximum entropy distribution with finite variance
  - the least possible assumptions on  $f$  in the distribution shape

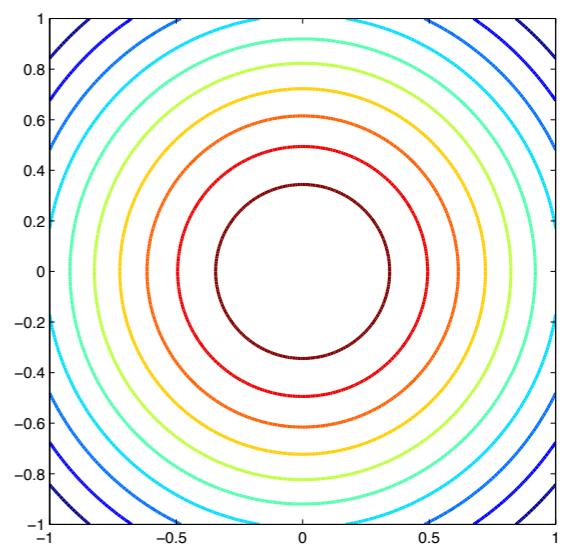
# Normal Distribution



probability density of the 1-D standard normal distribution



probability density of a 2-D normal distribution

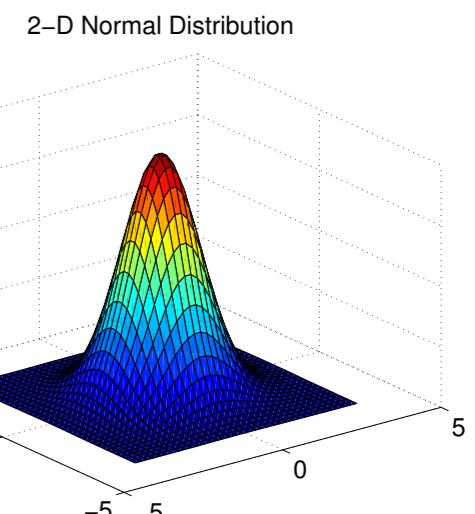


# The Multi-Variate ( $n$ -Dimensional) Normal Distribution

Any multi-variate normal distribution  $\mathcal{N}(\mathbf{m}, \mathbf{C})$  is uniquely determined by its mean value  $\mathbf{m} \in \mathbb{R}^n$  and its symmetric positive definite  $n \times n$  covariance matrix  $\mathbf{C}$ .

The **mean** value  $\mathbf{m}$

- determines the displacement (translation)
- value with the largest density (modal value)
- the distribution is symmetric about the distribution mean

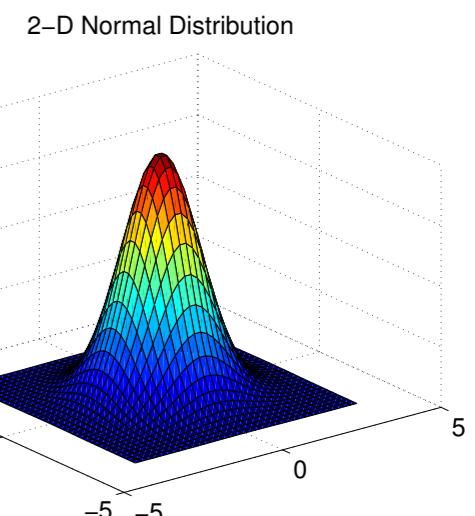


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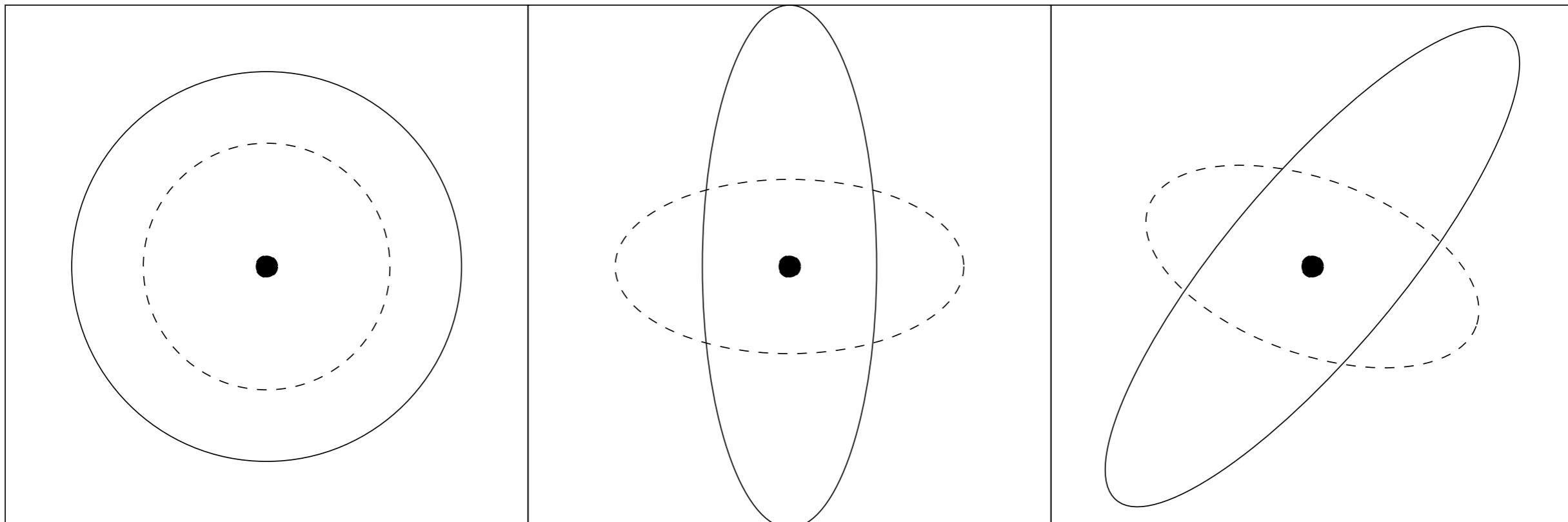


The **covariance matrix**  $\mathbf{C}$

- determines the shape
- **geometrical interpretation:** any covariance matrix can be uniquely identified with the iso-density ellipsoid  $\{\mathbf{x} \in \mathbb{R}^n \mid (\mathbf{x} - \mathbf{m})^T \mathbf{C}^{-1} (\mathbf{x} - \mathbf{m}) = 1\}$

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### Lines of Equal Density



$$\mathcal{N}(\mathbf{m}, \sigma^2 \mathbf{I}) \sim \mathbf{m} + \sigma \mathcal{N}(\mathbf{0}, \mathbf{I})$$

**one degree of freedom**  $\sigma$

components are independent standard normally distributed

$$\mathcal{N}(\mathbf{m}, \mathbf{D}^2) \sim \mathbf{m} + \mathbf{D} \mathcal{N}(\mathbf{0}, \mathbf{I})$$

**$n$  degrees of freedom**

components are independent, scaled

$$\mathcal{N}(\mathbf{m}, \mathbf{C}) \sim \mathbf{m} + \mathbf{C}^{\frac{1}{2}} \mathcal{N}(\mathbf{0}, \mathbf{I})$$

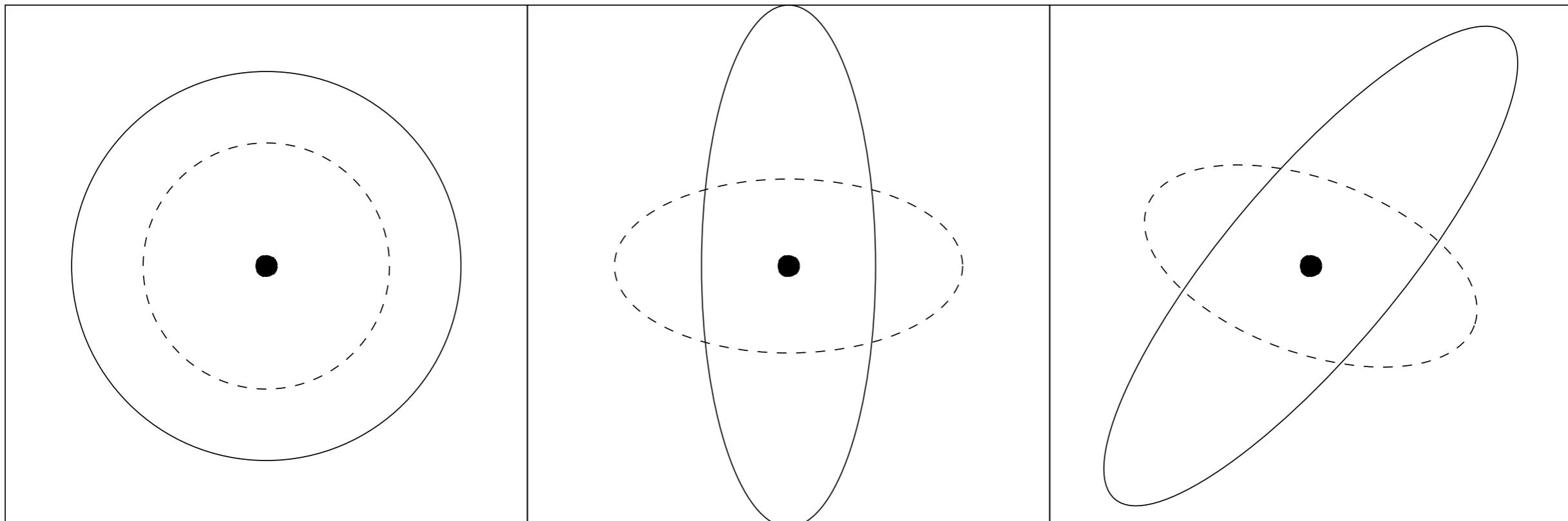
**$(n^2 + n)/2$  degrees of freedom**

components are correlated

where  $\mathbf{I}$  is the identity matrix (isotropic case) and  $\mathbf{D}$  is a diagonal matrix (reasonable for separable problems) and  $\mathbf{A} \times \mathcal{N}(\mathbf{0}, \mathbf{I}) \sim \mathcal{N}(\mathbf{0}, \mathbf{AA}^T)$  holds for all  $\mathbf{A}$ .

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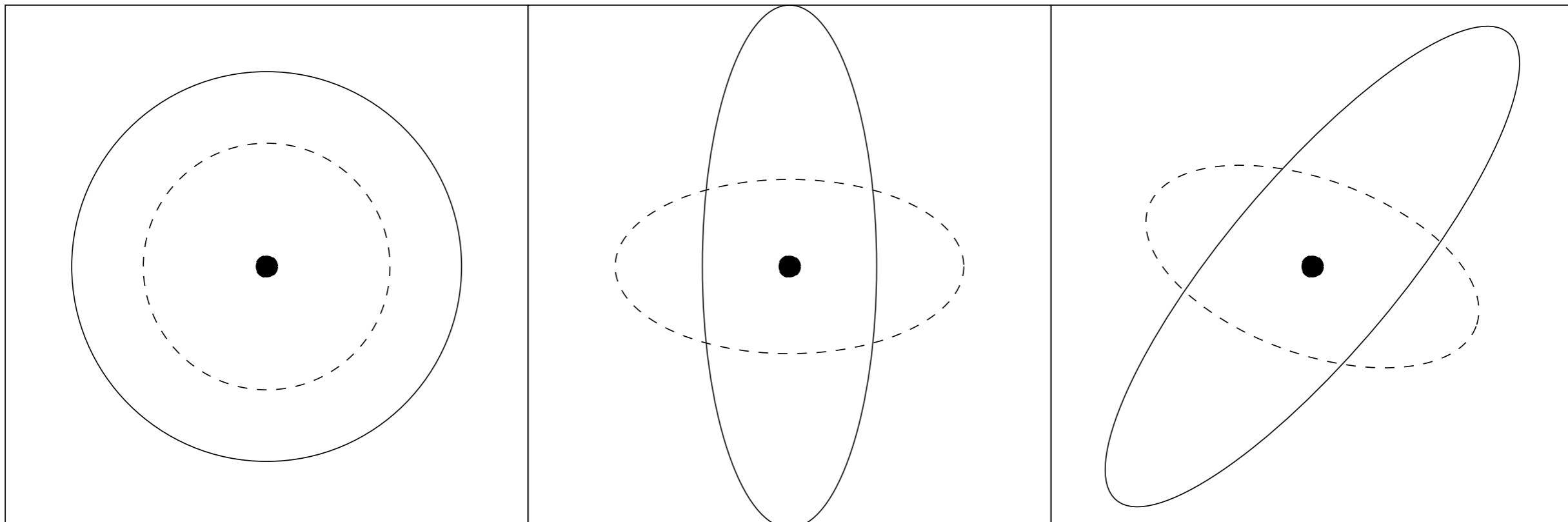
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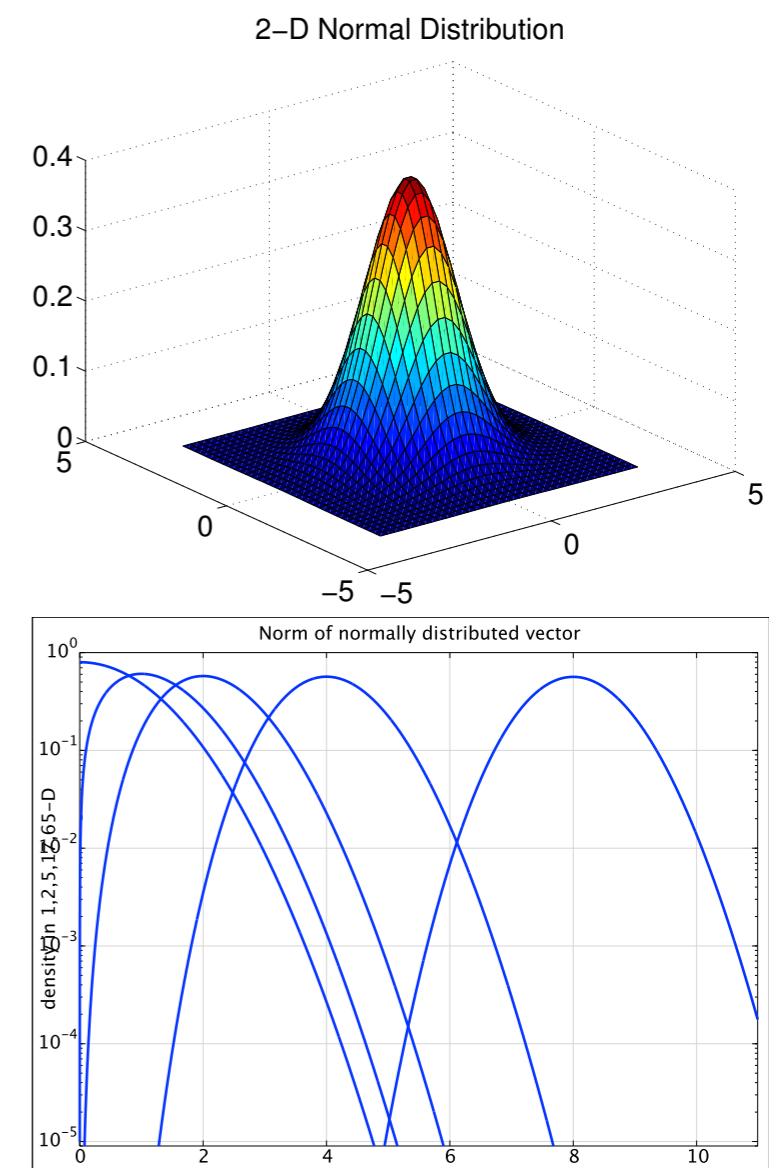
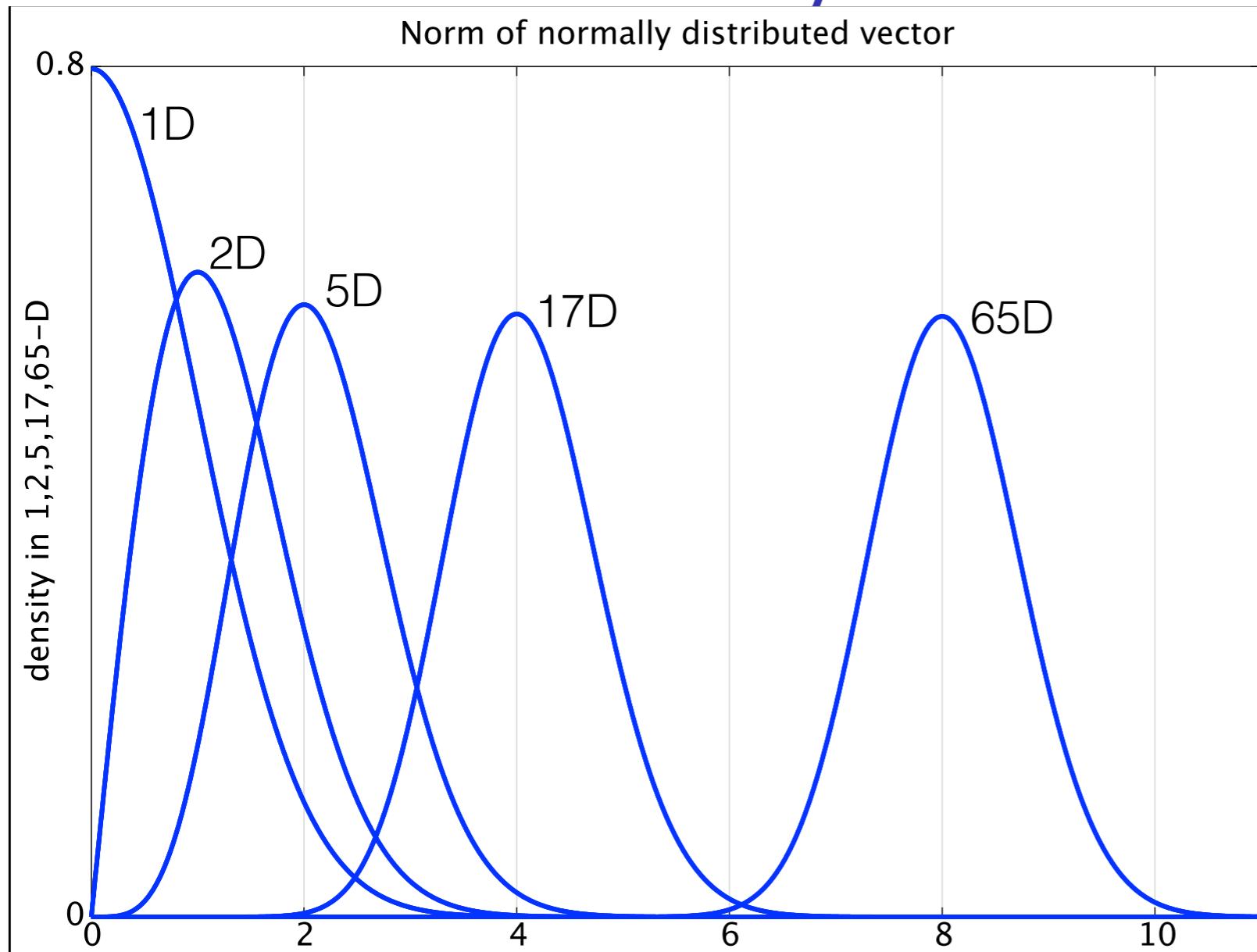
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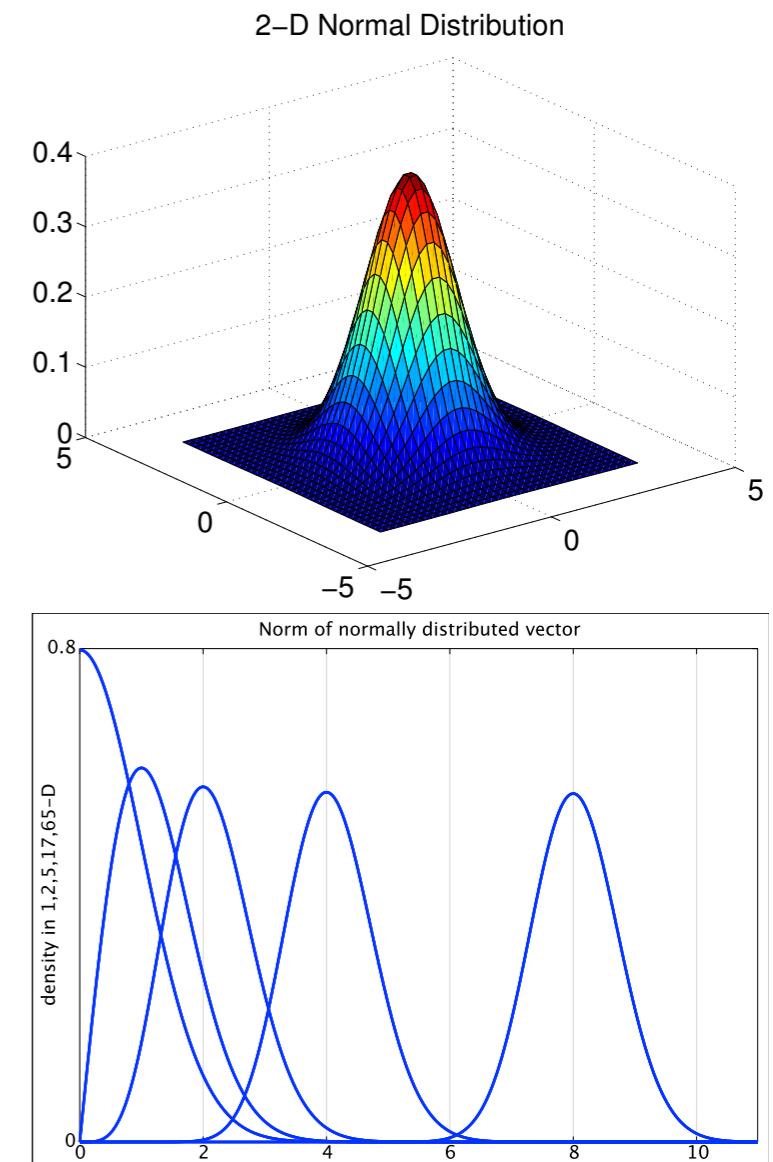
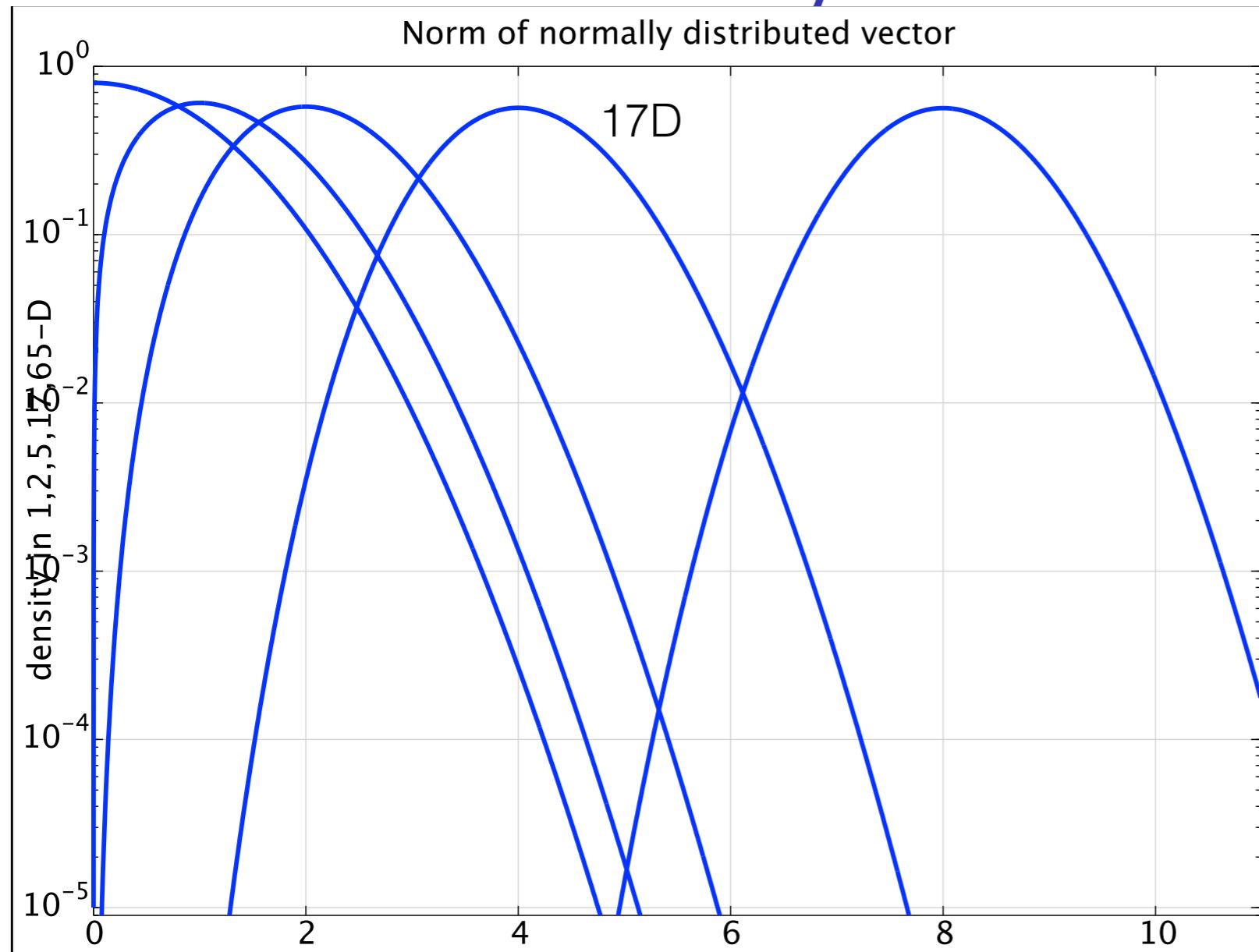
$$\|\mathcal{N}(\mathbf{0}, \mathbf{I})\| \rightarrow \mathcal{N}\left(\sqrt{n - 1/2}, 1/2\right) \text{ with modal value } \sqrt{n - 1}$$

yet: maximum entropy distribution

also consider a difference between two vectors:

$$\|\mathcal{N}(\mathbf{0}, \mathbf{I}) - \mathcal{N}(\mathbf{0}, \mathbf{I})\| \sim \|\mathcal{N}(\mathbf{0}, \mathbf{I}) + \mathcal{N}(\mathbf{0}, \mathbf{I})\| \sim \sqrt{2}\|\mathcal{N}(\mathbf{0}, \mathbf{I})\|$$

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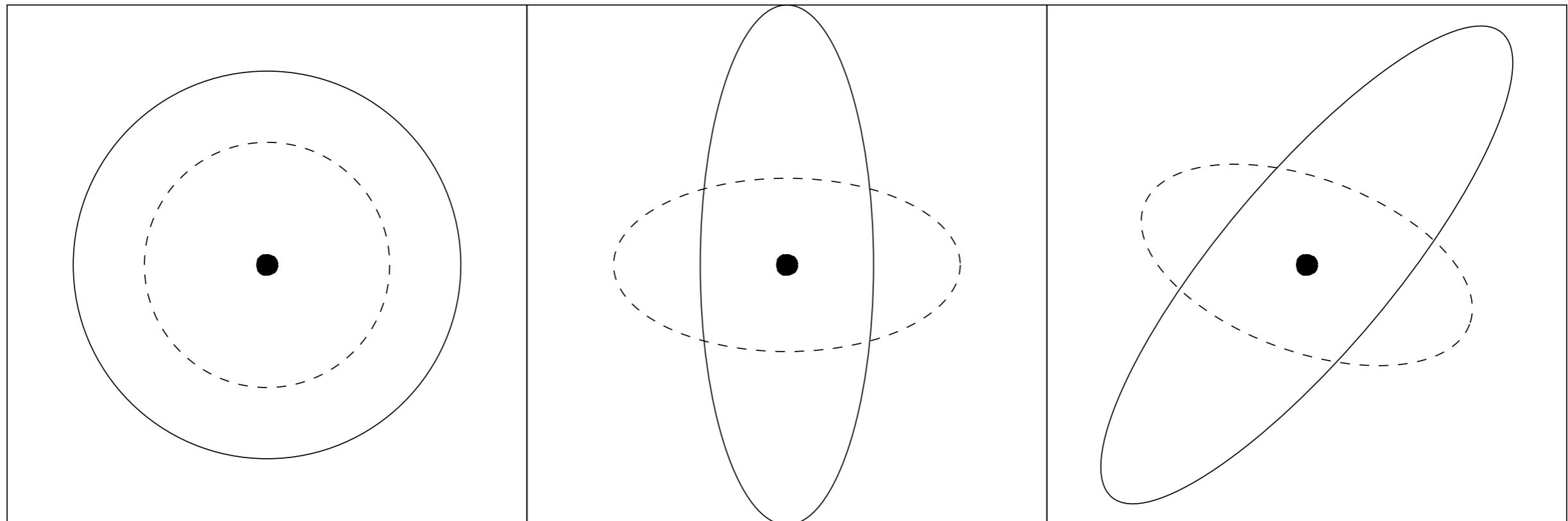
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What is the implication for the distribution in this picture (considering large dimension)?

...ESs

Update of the distribution mean

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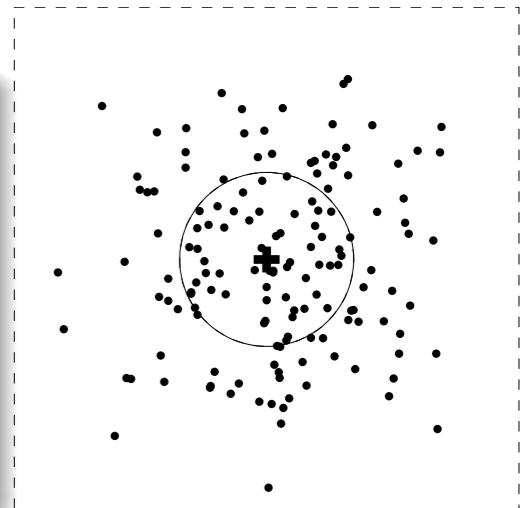
as perturbations of  $\mathbf{m}$ , where  $x_i, \mathbf{m} \in \mathbb{R}^n$ ,  $\sigma \in \mathbb{R}_+$ ,  $\mathbf{C} \in \mathbb{R}^{n \times n}$

where

- the **mean** vector  $\mathbf{m} \in \mathbb{R}^n$  represents the favorite solution
- the so-called **step-size**  $\sigma \in \mathbb{R}_+$  controls the *step length*
- the **covariance matrix**  $\mathbf{C} \in \mathbb{R}^{n \times n}$  determines the **shape** of the distribution ellipsoid

here, all new points are sampled with the same parameters

The question remains how to update  $\mathbf{m}$ ,  $\mathbf{C}$ , and  $\sigma$ .



# The $(\mu/\mu, \lambda)$ -ES

Non-elitist selection and intermediate (weighted) recombination

Given the  $i$ -th solution point  $\mathbf{x}_i = \mathbf{m} + \sigma \underbrace{\mathcal{N}_i(\mathbf{0}, \mathbf{C})}_{=: \mathbf{y}_i} = \mathbf{m} + \sigma \mathbf{y}_i$

Let  $\mathbf{x}_{i:\lambda}$  the  $i$ -th ranked solution point, such that  $f(\mathbf{x}_{1:\lambda}) \leq \dots \leq f(\mathbf{x}_{\lambda:\lambda})$ .

The new mean reads

$$\mathbf{m} \leftarrow \sum_{i=1}^{\mu} w_i \mathbf{x}_{i:\lambda}$$

where

$$w_1 \geq \dots \geq w_\mu > 0, \quad \sum_{i=1}^{\mu} w_i = 1, \quad \frac{1}{\sum_{i=1}^{\mu} w_i^2} =: \mu_w \approx \frac{\lambda}{4}$$

The best  $\mu$  points are selected from the new solutions (non-elitistic) and weighted intermediate recombination is applied.

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The best  $\mu$  points are selected from the new solutions (non-elitistic) and weighted intermediate recombination is applied.

# Evolution Strategies

## Terminology

$\mu$ : # of parents,  $\lambda$ : # of offspring

## Plus (elitist) and comma (non-elitist) selection

$(\mu + \lambda)$ -ES: selection in  $\{\text{parents}\} \cup \{\text{offspring}\}$

$(\mu, \lambda)$ -ES: selection in  $\{\text{offspring}\}$

## $(1 + 1)$ -ES

Sample one offspring from parent  $\mathbf{m}$

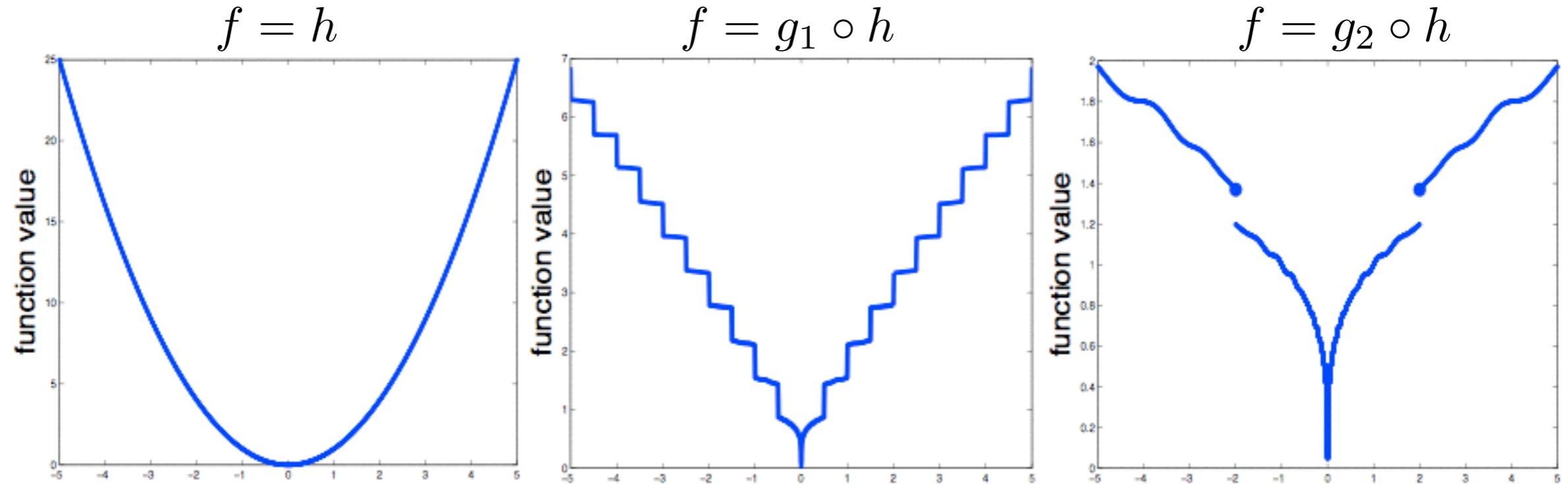
$$\mathbf{x} = \mathbf{m} + \sigma \mathcal{N}(\mathbf{0}, \mathbf{C})$$

If  $x$  better than  $\mathbf{m}$  select

$$\mathbf{m} \leftarrow \mathbf{x}$$

# Invariance

# Invariance: Function-Value Free Property



Three functions belonging to the same equivalence class

A *function-value free search algorithm* is invariant under the transformation with any *order preserving* (strictly increasing)  $g$ .

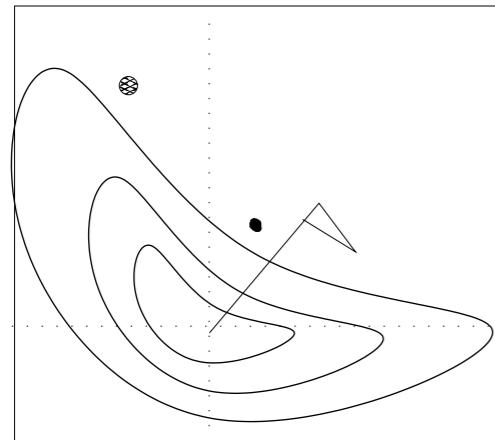
Invariances make

- observations meaningful as a rigorous notion of generalization
- algorithms predictable and/or "robust"

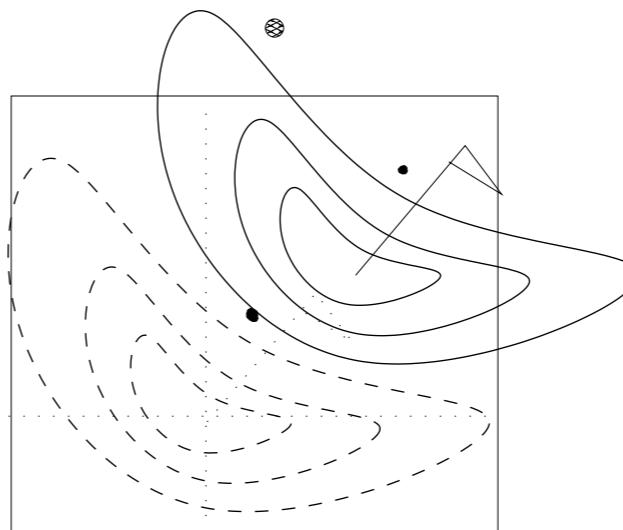
# Basic Invariance in Search Space

- translation invariance

is true for most optimization algorithms



$$f(\mathbf{x}) \leftrightarrow f(\mathbf{x} - \mathbf{a})$$



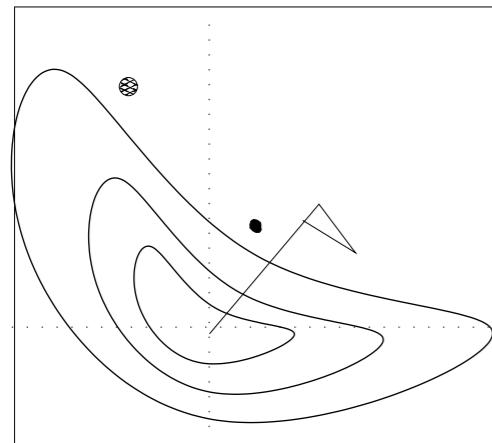
Identical behavior on  $f$  and  $f_a$

$$\begin{aligned} f : \quad \mathbf{x} &\mapsto f(\mathbf{x}), & \mathbf{x}^{(t=0)} &= \mathbf{x}_0 \\ f_a : \quad \mathbf{x} &\mapsto f(\mathbf{x} - \mathbf{a}), & \mathbf{x}^{(t=0)} &= \mathbf{x}_0 + \mathbf{a} \end{aligned}$$

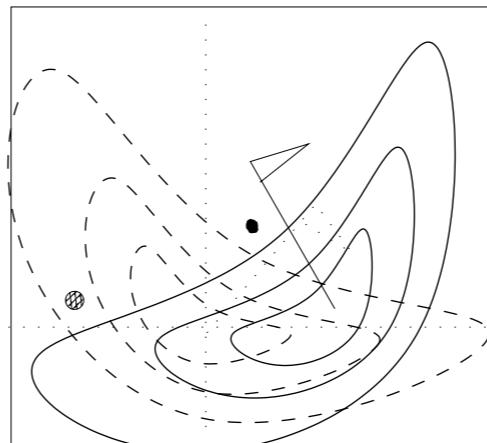
No difference can be observed w.r.t. the argument of  $f$

# Rotational Invariance in Search Space

- invariance to orthogonal (rigid) transformations  $\mathbf{R}$ , where  $\mathbf{RR}^T = \mathbf{I}$   
e.g. true for simple evolution strategies  
recombination operators might jeopardize rotational invariance



$$f(\mathbf{x}) \leftrightarrow f(\mathbf{Rx})$$



Identical behavior on  $f$  and  $f_{\mathbf{R}}$

$$\begin{aligned} f : \quad \mathbf{x} &\mapsto f(\mathbf{x}), & \mathbf{x}^{(t=0)} &= \mathbf{x}_0 \\ f_{\mathbf{R}} : \quad \mathbf{x} &\mapsto f(\mathbf{Rx}), & \mathbf{x}^{(t=0)} &= \mathbf{R}^{-1}(\mathbf{x}_0) \end{aligned}$$

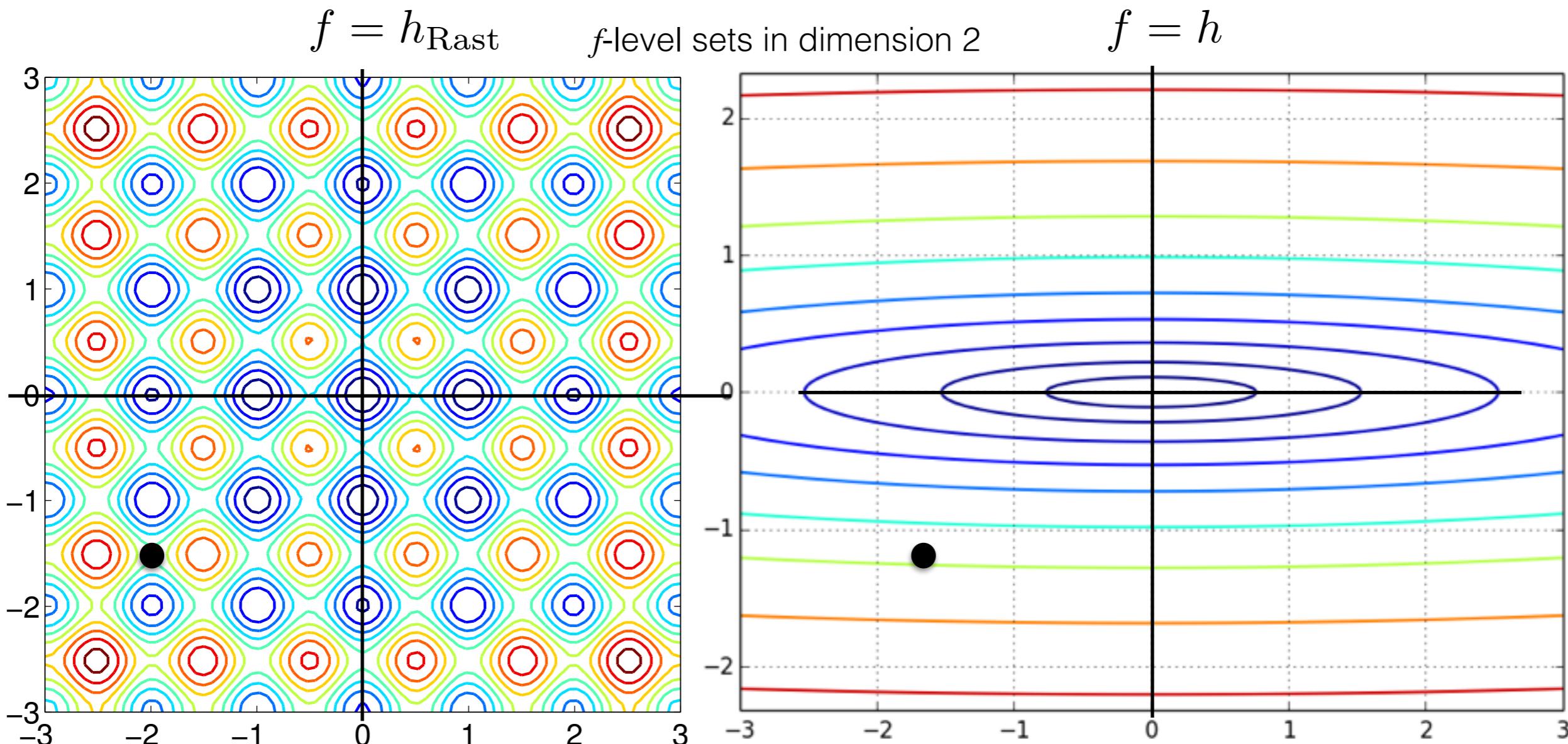
45

No difference can be observed w.r.t. the argument of  $f$

<sup>4</sup> Salomon 1996. "Reevaluating Genetic Algorithm Performance under Coordinate Rotation of Benchmark Functions; A survey of some theoretical and practical aspects of genetic algorithms." BioSystems, 39(3):263-278

<sup>5</sup> Hansen 2000. Invariance, Self-Adaptation and Correlated Mutations in Evolution Strategies. *Parallel Problem Solving from Nature PPSN VI*

# Invariance Under Rigid Search Space Transformations

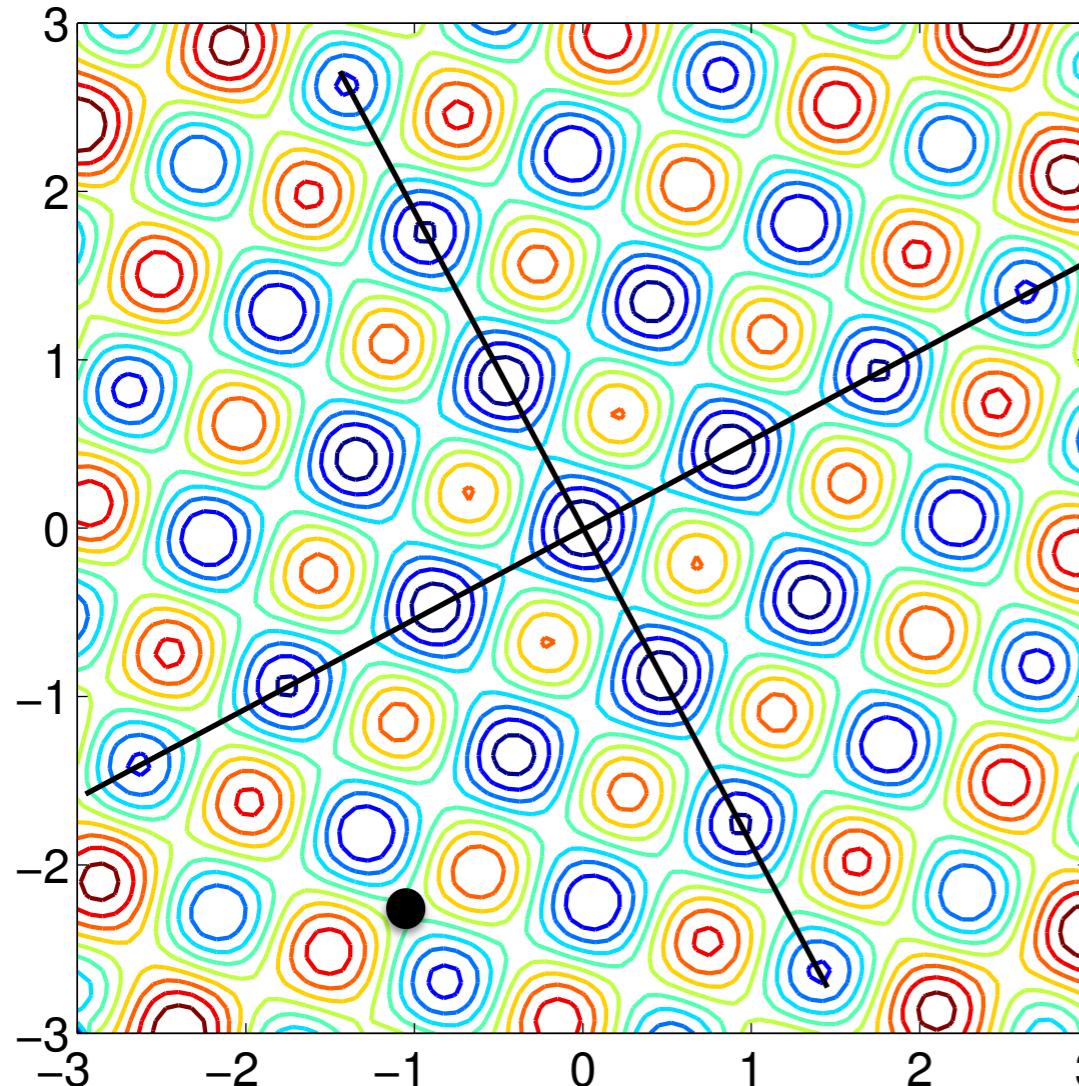


for example, invariance under search space rotation  
(**separable**  $\Leftrightarrow$  non-separable)

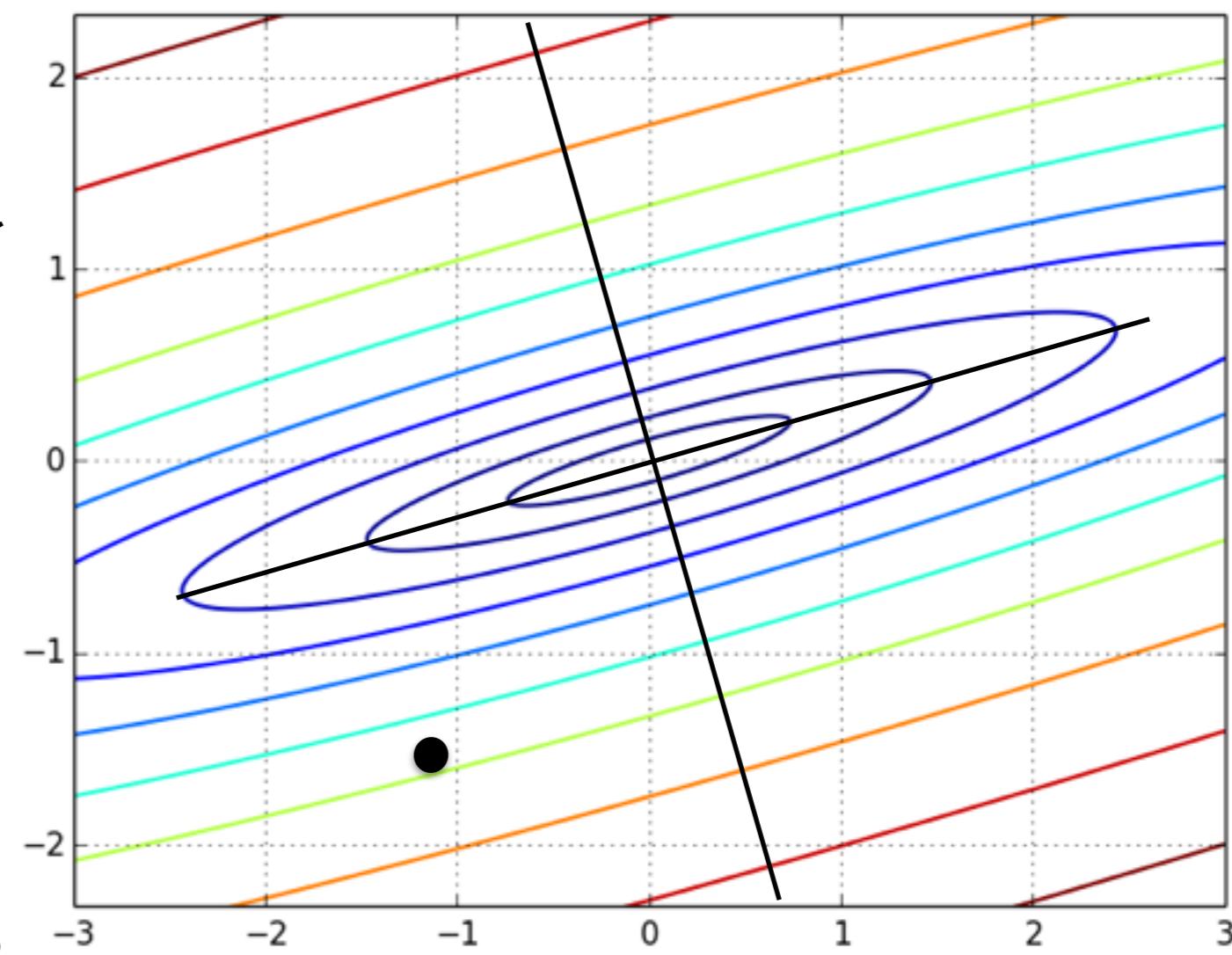
# Invariance Under Rigid Search Space Transformations

$$f = h_{\text{Rast}} \circ R$$

$f$ -level sets in dimension 2



$$f = h \circ R$$



for example, invariance under search space rotation  
(separable  $\Leftrightarrow$  non-separable)

# Landscape of Continuous Search Methods

## *Gradient-based (Taylor, local)*

- Conjugate gradient methods [Fletcher & Reeves 1964]
- Quasi-Newton methods (BFGS) [Broyden et al 1970]

## *Derivative-free optimization (DFO)*

- Trust-region methods (NEWUOA, BOBYQA) [Powell 2006, 2009]
- Simplex downhill [Nelder & Mead 1965]
- Pattern search [Hooke & Jeeves 1961, Audet & Dennis 2006]

## *Stochastic (randomized) search methods*

- Evolutionary algorithms (broader sense, continuous domain)
  - Differential Evolution [Storn & Price 1997]
  - Particle Swarm Optimization [Kennedy & Eberhart 1995]
  - **Evolution Strategies** [Rechenberg 1965, Hansen & Ostermeier 2001]
- Simulated annealing [Kirkpatrick et al 1983]
- Simultaneous perturbation stochastic approximation (SPSA) [Spall 2000]

# Invariance

*The grand aim of all science is to cover the greatest number of empirical facts by logical deduction from the smallest number of hypotheses or axioms.*

— Albert Einstein

- Empirical performance results

- ▶ from benchmark functions
  - ▶ from solved real world problems

are only useful if they do **generalize** to other problems

- **Invariance** is a strong **non-empirical** statement about generalization

generalizing (identical) performance from a single function to a whole class of functions

consequently, invariance is important for the evaluation of search algorithms

# Step-Size Control

# Evolution Strategies

Recalling

New search points are sampled normally distributed

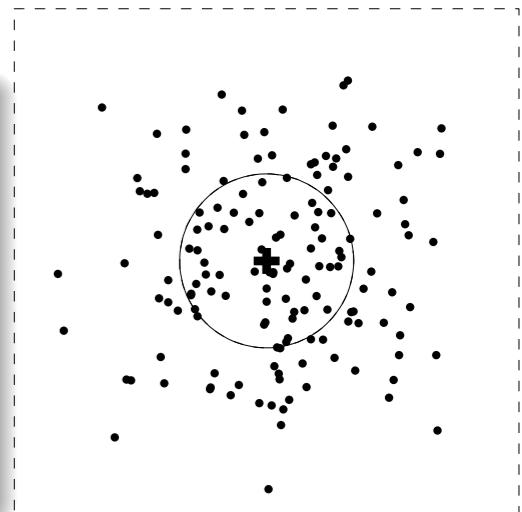
$$\mathbf{x}_i \sim \mathbf{m} + \sigma \mathcal{N}_i(\mathbf{0}, \mathbf{C}) \quad \text{for } i = 1, \dots, \lambda$$

as perturbations of  $\mathbf{m}$ , where  $\mathbf{x}_i, \mathbf{m} \in \mathbb{R}^n$ ,  $\sigma \in \mathbb{R}_+$ ,  $\mathbf{C} \in \mathbb{R}^{n \times n}$

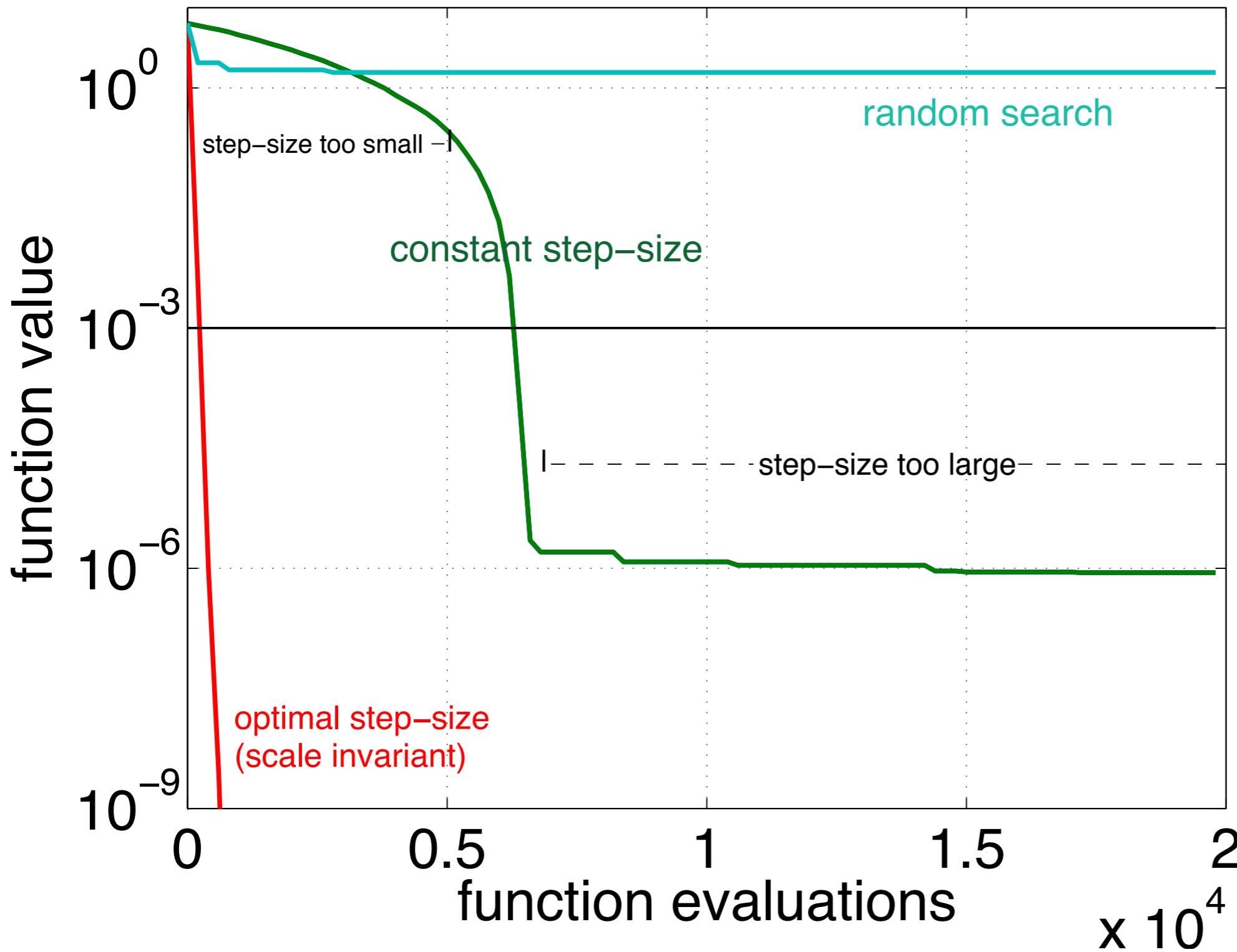
where

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- the so-called **step-size**  $\sigma \in \mathbb{R}_+$  controls the *step length*
- the **covariance matrix**  $\mathbf{C} \in \mathbb{R}^{n \times n}$  determines the **shape** of the distribution ellipsoid

The remaining question is how to update  $\sigma$  and  $\mathbf{C}$ .



# Why Step-Size Control?



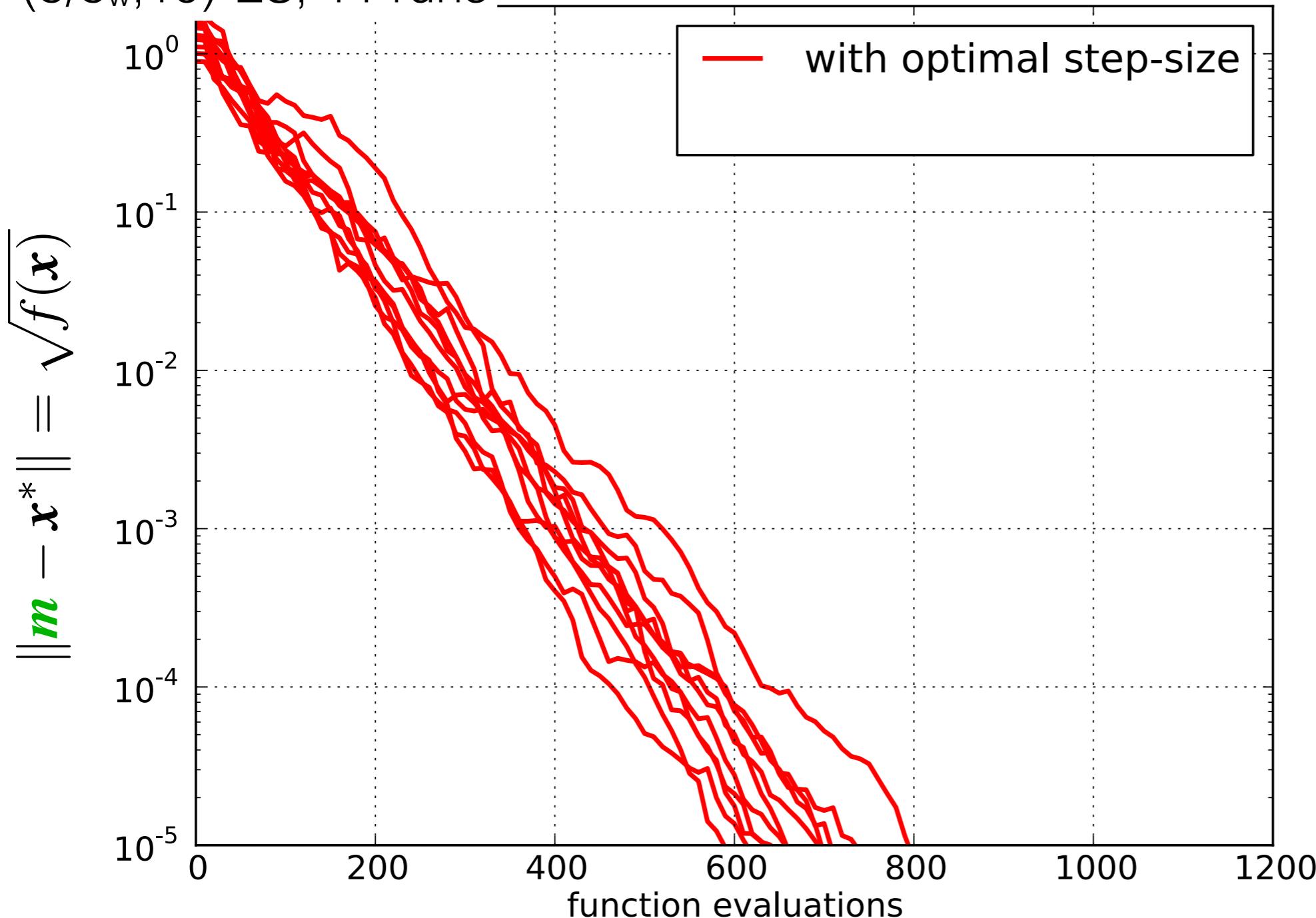
(1+1)-ES  
(red & green)

$$f(\mathbf{x}) = \sum_{i=1}^n x_i^2$$

in  $[-2.2, 0.8]^n$   
for  $n = 10$

# Why Step-Size Control?

(5/5<sub>w</sub>, 10)-ES, 11 runs



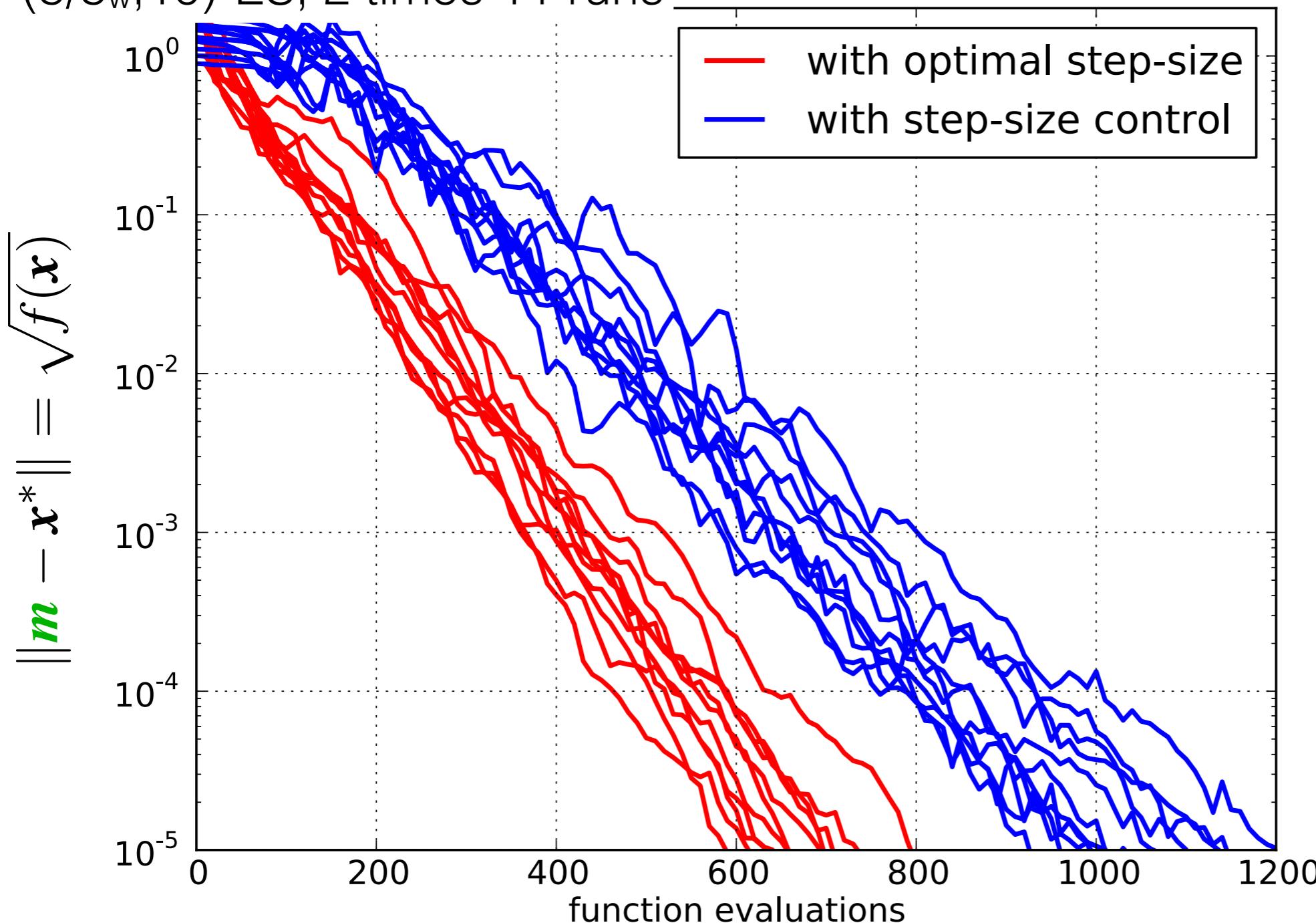
$$f(\mathbf{x}) = \sum_{i=1}^n x_i^2$$

for  $n = 10$  and  
 $\mathbf{x}^0 \in [-0.2, 0.8]^n$

with optimal step-size  $\sigma$

# Why Step-Size Control?

(5/5<sub>w</sub>, 10)-ES, 2 times 11 runs



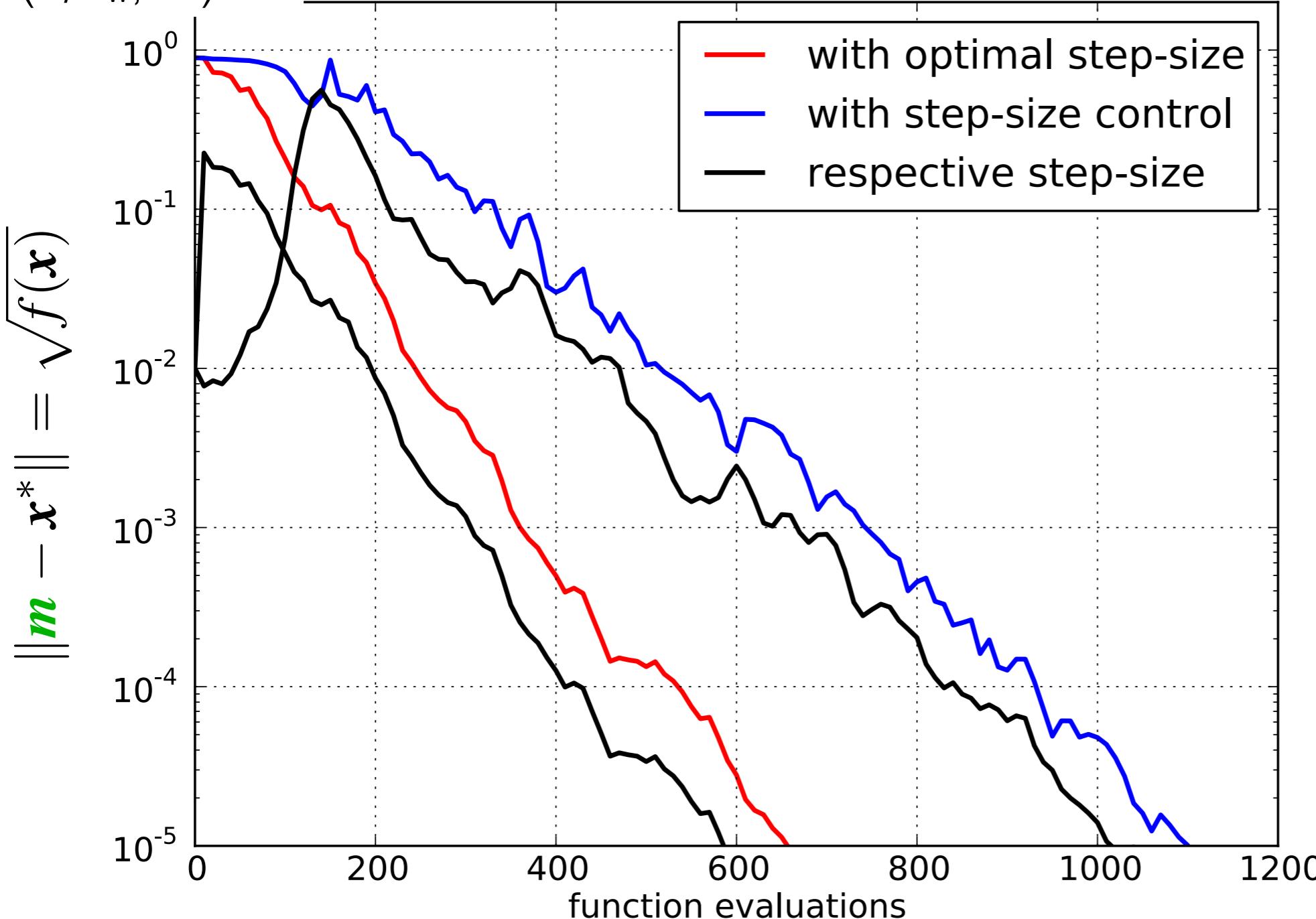
$$f(\mathbf{x}) = \sum_{i=1}^n x_i^2$$

for  $n = 10$  and  
 $\mathbf{x}^0 \in [-0.2, 0.8]^n$

with **optimal** versus adaptive step-size  $\sigma$  with too small initial  $\sigma$

# Why Step-Size Control?

(5/5<sub>w</sub>, 10)-ES



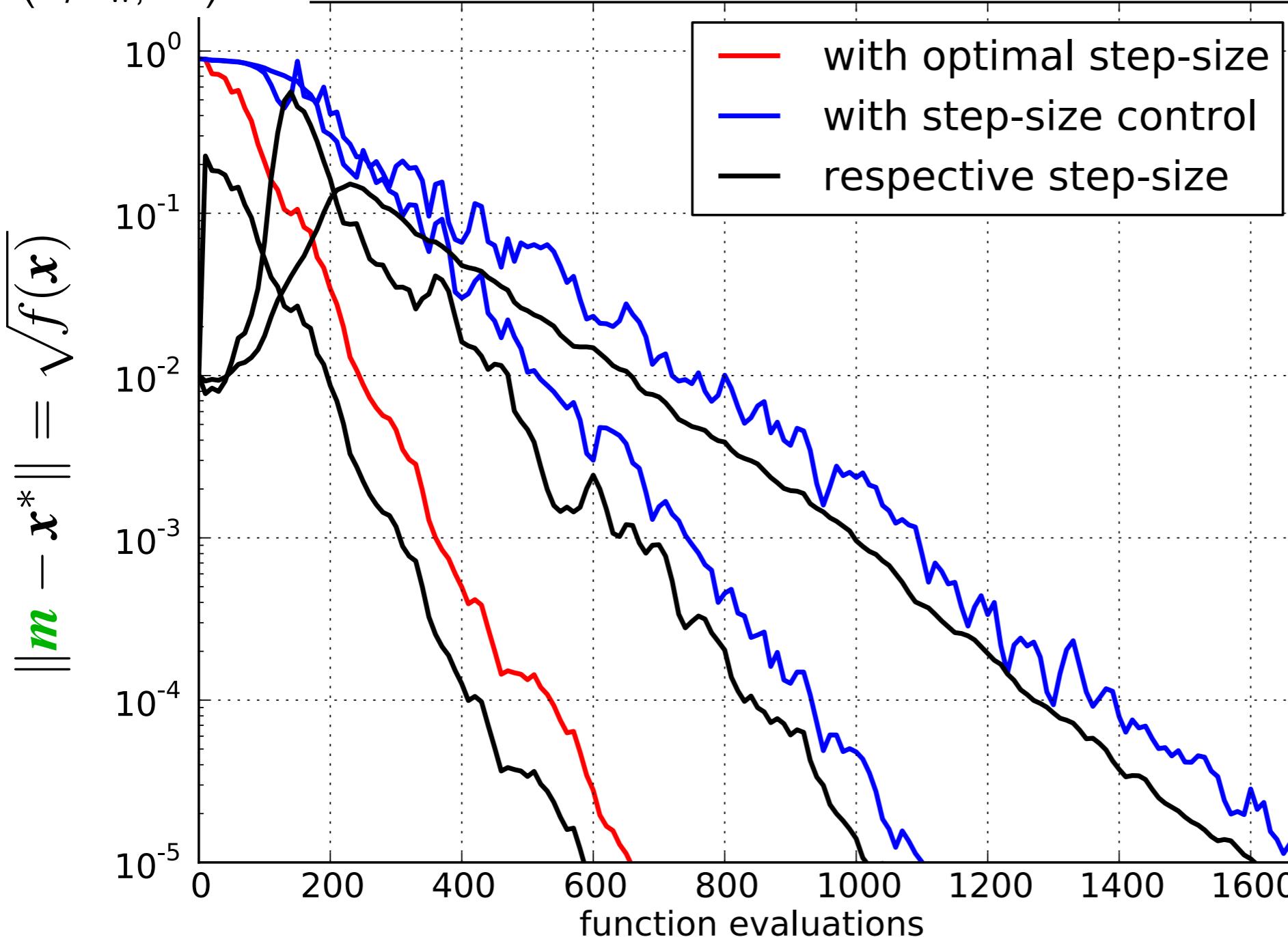
$$f(\mathbf{x}) = \sum_{i=1}^n x_i^2$$

for  $n = 10$  and  
 $\mathbf{x}^0 \in [-0.2, 0.8]^n$

comparing number of  $f$ -evals to reach  $\|\mathbf{m}\| = 10^{-5}$ :  $\frac{1100-100}{650} \approx 1.5$

# Why Step-Size Control?

(5/5<sub>w</sub>, 10)-ES



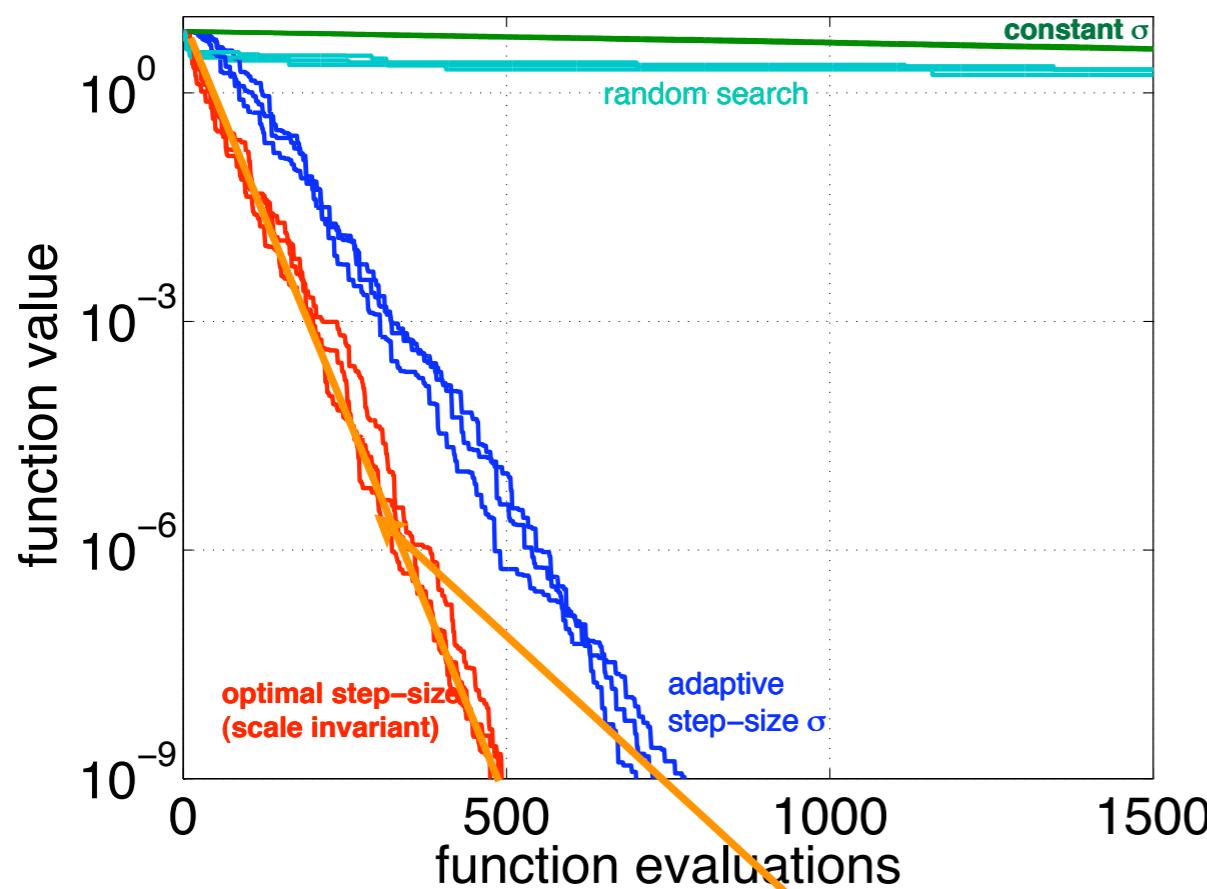
$$f(\mathbf{x}) = \sum_{i=1}^n x_i^2$$

in  $[-0.2, 0.8]^n$   
for  $n = 10$

comparing optimal versus default damping parameter  $d_\sigma$ :  $\frac{1700}{1100} \approx 1.5$

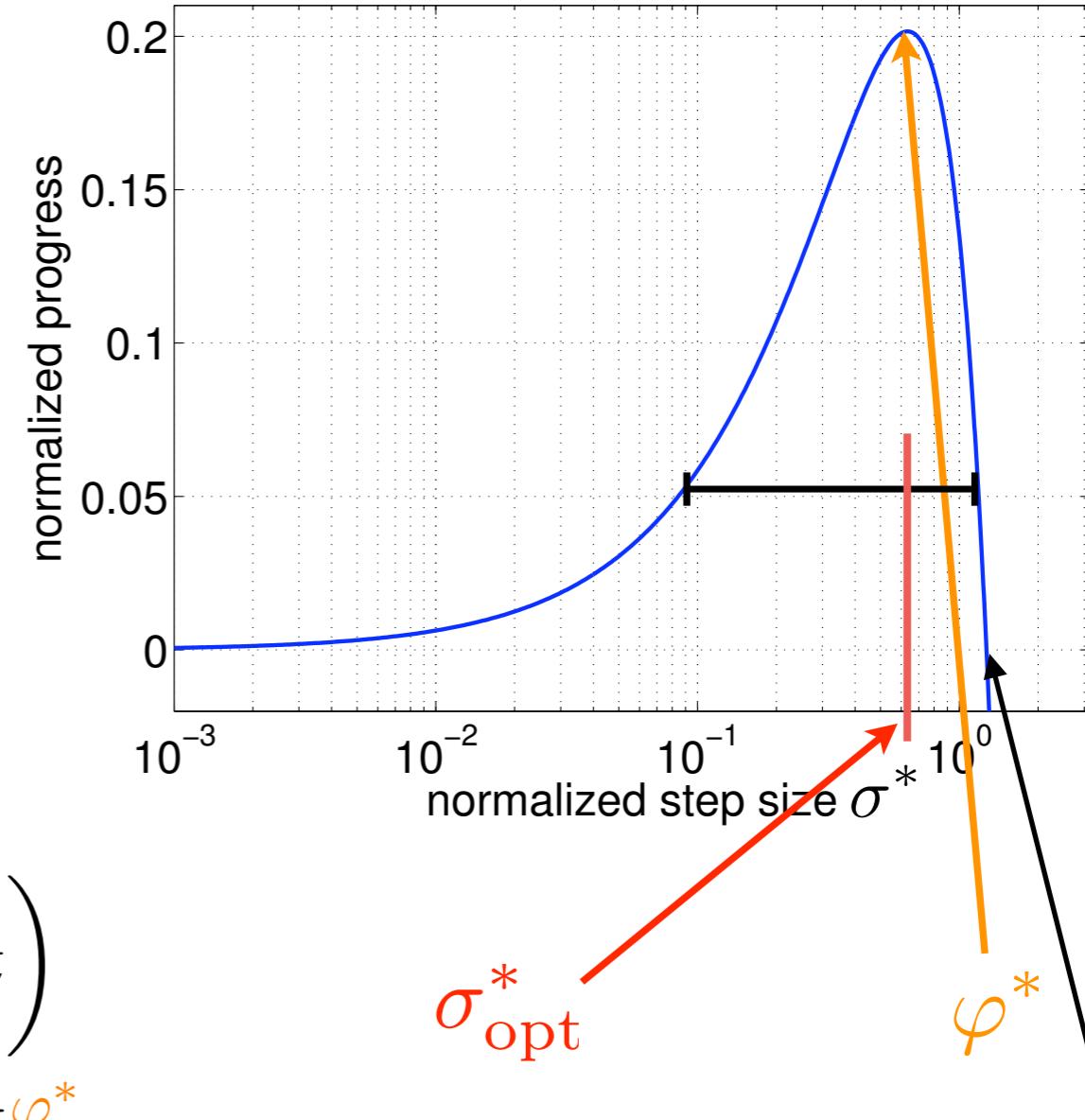
# Why Step-Size Control?

$$\sigma_{\text{opt}} \approx \sigma_{\text{opt}}^* \mu_w \frac{\|m\|}{n}$$



$$f(\mathbf{m}^{(t)}) \approx f(\mathbf{m}^{(0)}) \times \exp \left( 2 \frac{-\varphi^*}{n} \lambda t \right)$$

$$\frac{\lg(f(\mathbf{m}^{(t+1)})) - \lg(f(\mathbf{m}^{(t)}))}{\lambda} \approx \lg(e) \times 2 \frac{-\varphi^*}{n}$$



*evolution window* refers to the step-size interval ( $\text{---}$ ) where reasonable performance is observed

# Methods for Step-Size Control

- **1/5-th success rule<sup>ab</sup>**, often applied with “+”-selection  
increase step-size if more than 20% of the new solutions are successful,  
decrease otherwise
- **$\sigma$ -self-adaptation<sup>c</sup>**, applied with “,”-selection  
mutation is applied to the step-size and the better, according to the  
objective function value, is selected  
simplified “global” self-adaptation
- **path length control<sup>d</sup>** (Cumulative Step-size Adaptation, CSA)<sup>e</sup>  
self-adaptation derandomized and non-localized

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<sup>a</sup> Rechenberg 1973, *Evolutionsstrategie, Optimierung technischer Systeme nach Prinzipien der biologischen Evolution*, Frommann-Holzboog

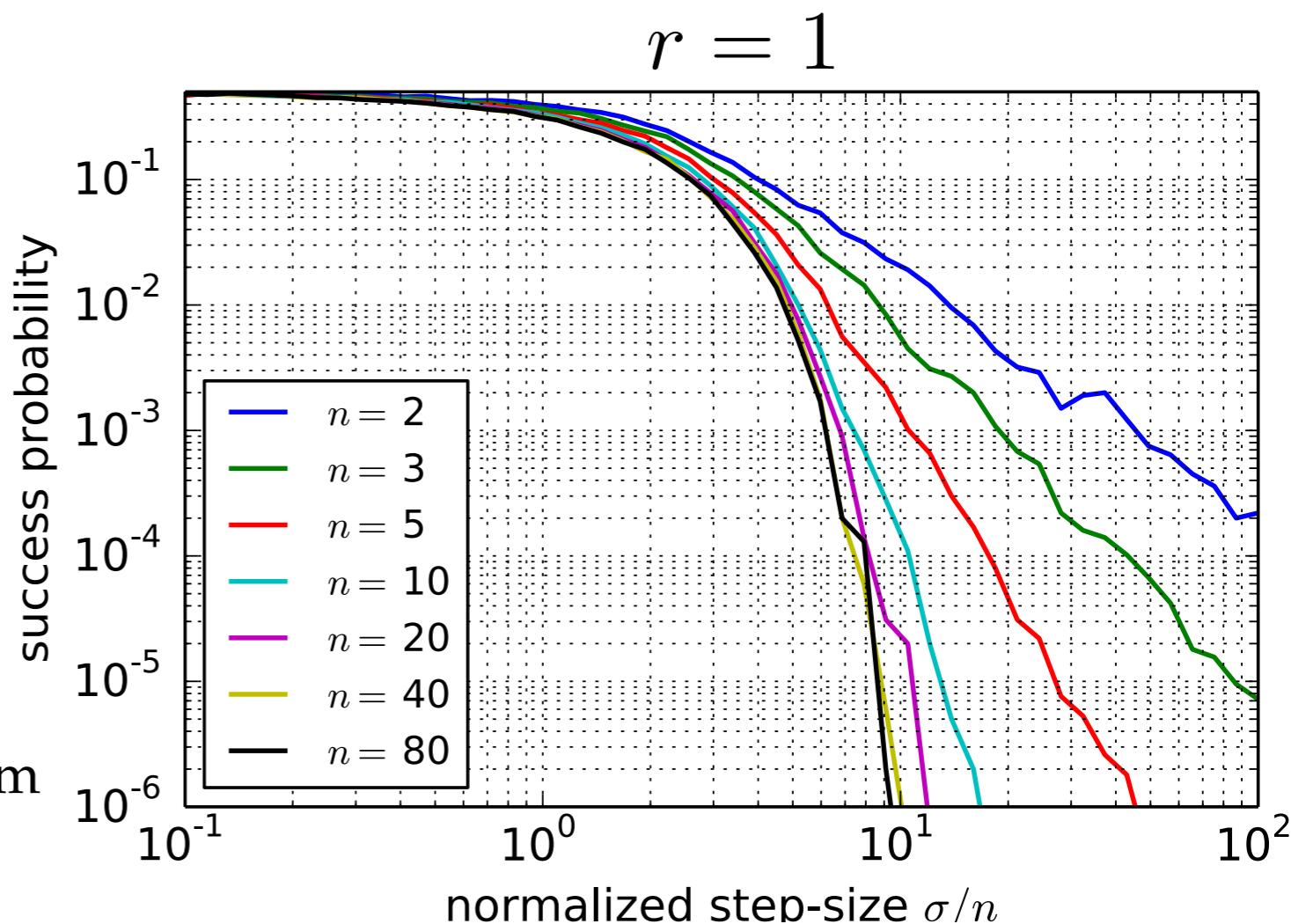
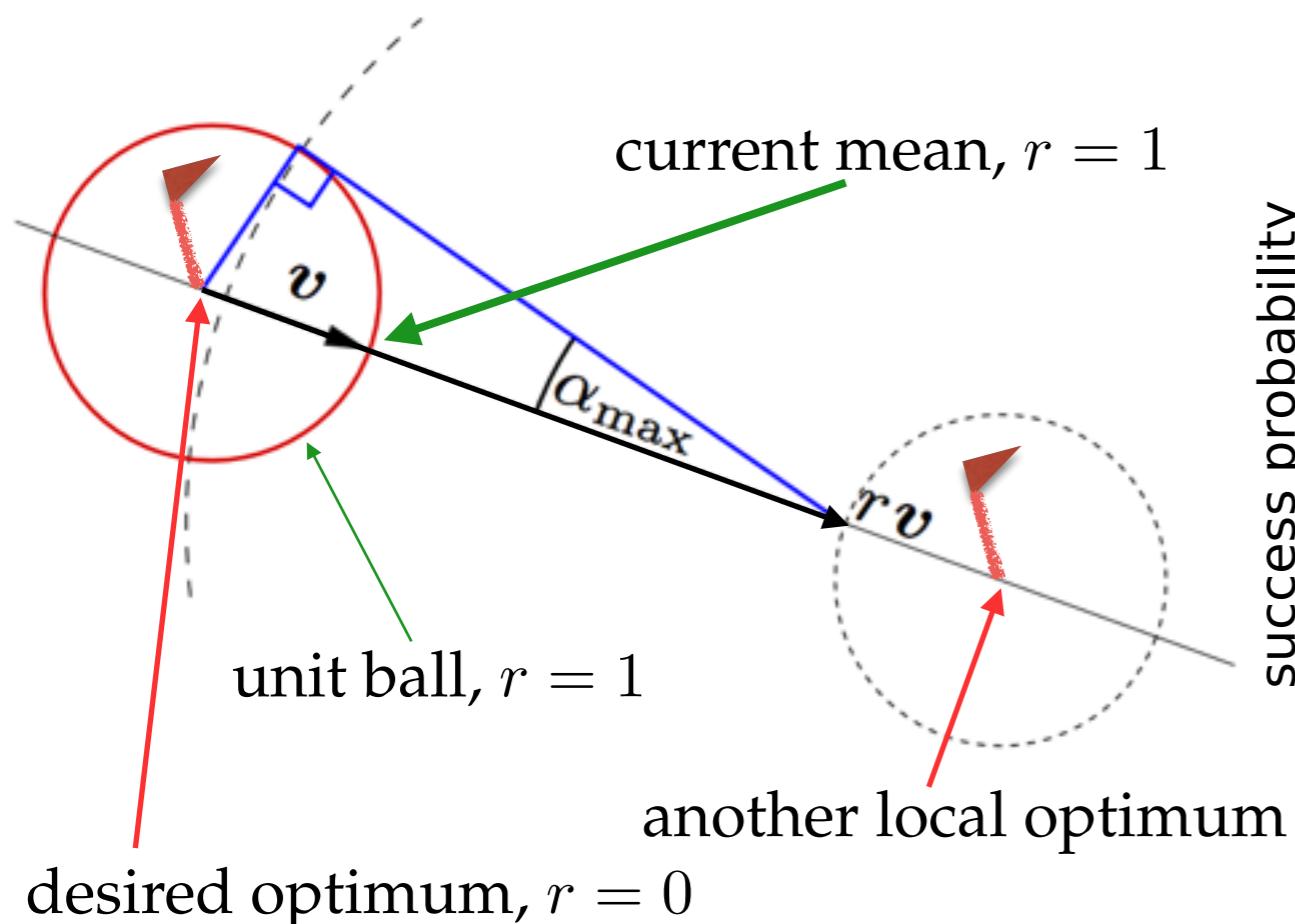
<sup>b</sup> Schumer and Steiglitz 1968. Adaptive step size random search. *IEEE TAC*

<sup>c</sup> Schwefel 1981, *Numerical Optimization of Computer Models*, Wiley

<sup>d</sup> Hansen & Ostermeier 2001, Completely Derandomized Self-Adaptation in Evolution Strategies, *Evol. Comput.*  
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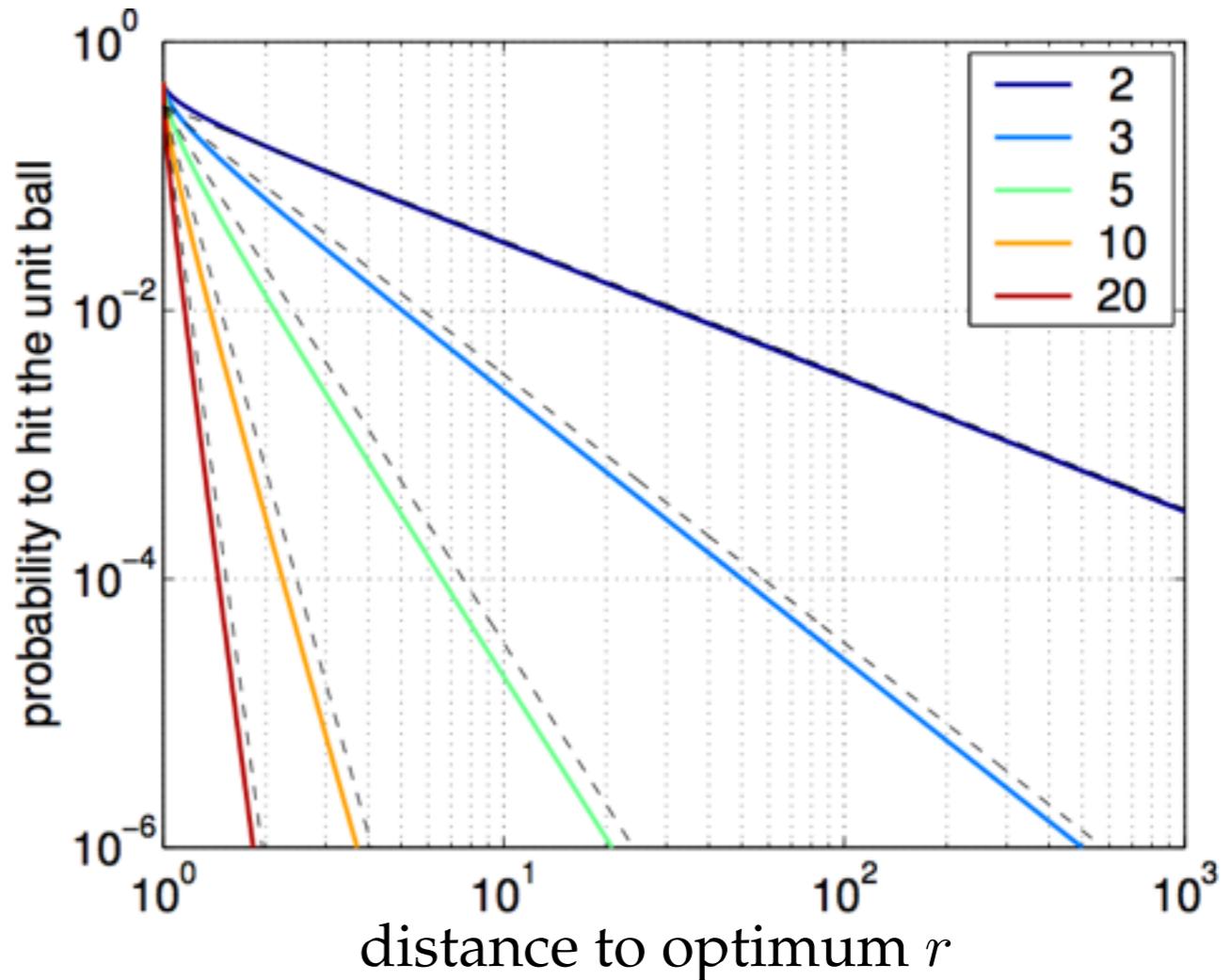
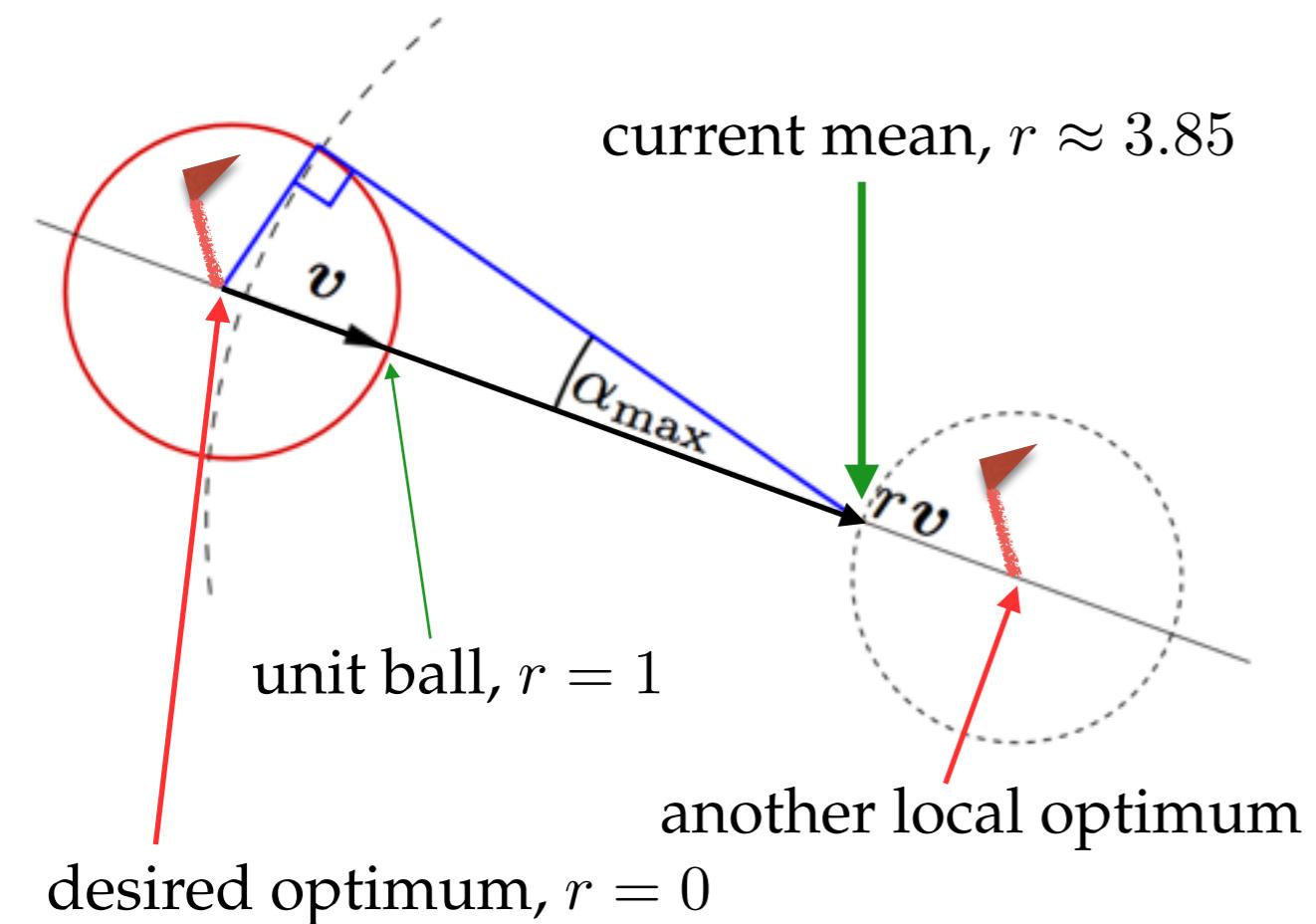
<sup>e</sup> Ostermeier *et al* 1994, Step-size adaptation based on non-local use of selection information, *PPSN IV*

# Success vs step-size



probability to hit the unit ball as a function of step-size

# Success vs distance



**Fig. 1.** Probability to hit the unit hyperball (solid) sampling from  $rv$  as mean with an optimal isotropic distribution, where  $v \in \mathbb{R}^n$  and  $\|v\| = 1$ . The plots on the right show results for  $n = 2, 3, 5, 10, 20$ , from above to below. Dashed lines depict the approximation  $\frac{1}{3r^{n-1}}$ .

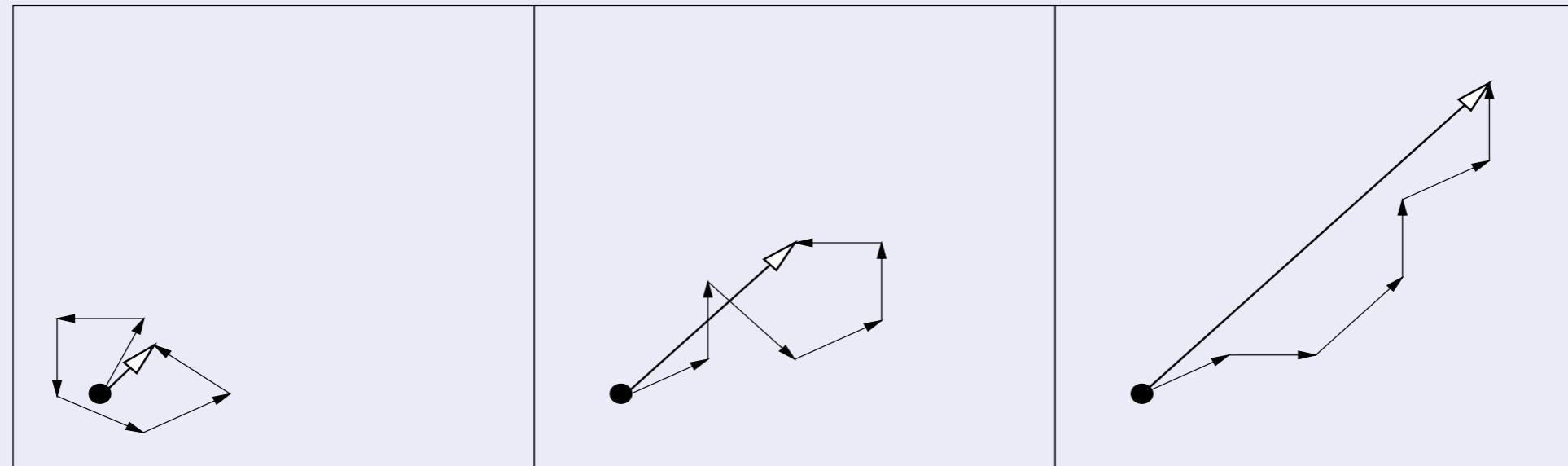
# Path Length Control (CSA)

The Concept of Cumulative Step-Size Adaptation

$$\begin{aligned} \mathbf{x}_i &= \mathbf{m} + \sigma \mathbf{y}_i \\ \mathbf{m} &\leftarrow \mathbf{m} + \sigma \mathbf{y}_w \end{aligned}$$

Measure the length of the *evolution path*

the pathway of the mean vector  $\mathbf{m}$  in the generation sequence



loosely speaking steps are

- perpendicular under random selection (in expectation)
- perpendicular in the desired situation (to be most efficient)

# Path Length Control (CSA)

## The Equations

Initialize  $\mathbf{m} \in \mathbb{R}^n$ ,  $\sigma \in \mathbb{R}_+$ , evolution path  $\mathbf{p}_\sigma = \mathbf{0}$ ,  
set  $c_\sigma \approx 4/n$ ,  $d_\sigma \approx 1$ .

$$\begin{aligned} \mathbf{m} &\leftarrow \mathbf{m} + \sigma \mathbf{y}_w \quad \text{where } \mathbf{y}_w = \sum_{i=1}^{\mu} w_i \mathbf{y}_{i:\lambda} && \text{update mean} \\ \mathbf{p}_\sigma &\leftarrow (1 - c_\sigma) \mathbf{p}_\sigma + \underbrace{\sqrt{1 - (1 - c_\sigma)^2}}_{\text{accounts for } 1 - c_\sigma} \underbrace{\sqrt{\mu_w}}_{\text{accounts for } w_i} \mathbf{y}_w \\ \sigma &\leftarrow \sigma \times \underbrace{\exp \left( \frac{c_\sigma}{d_\sigma} \left( \frac{\|\mathbf{p}_\sigma\|}{\mathbb{E}\|\mathcal{N}(\mathbf{0}, \mathbf{I})\|} - 1 \right) \right)}_{>1 \iff \|\mathbf{p}_\sigma\| \text{ is greater than its expectation}} && \text{update step-size} \end{aligned}$$

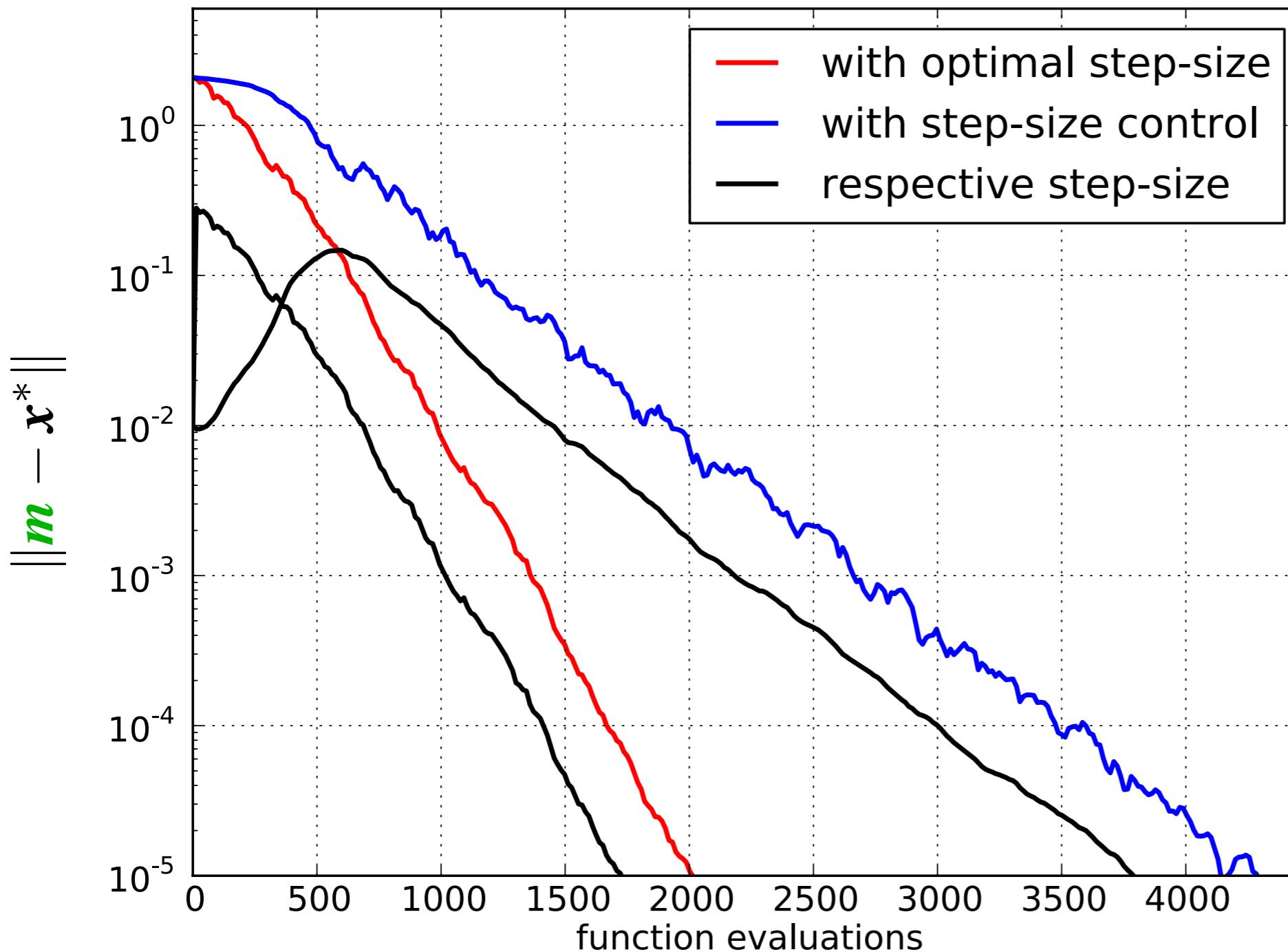
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## (5/5, 10)-CSA-ES, default parameters



$$f(\mathbf{x}) = \sum_{i=1}^n x_i^2$$

in  $[-0.2, 0.8]^n$   
for  $n = 30$

# Covariance Matrix Adaptation

# Evolution Strategies

Recalling

New search points are sampled normally distributed

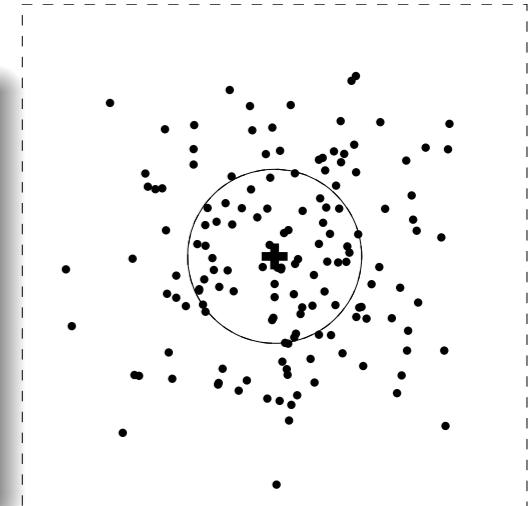
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as perturbations of  $\mathbf{m}$ , where  $\mathbf{x}_i, \mathbf{m} \in \mathbb{R}^n$ ,  $\sigma \in \mathbb{R}_+$ ,  $\mathbf{C} \in \mathbb{R}^{n \times n}$

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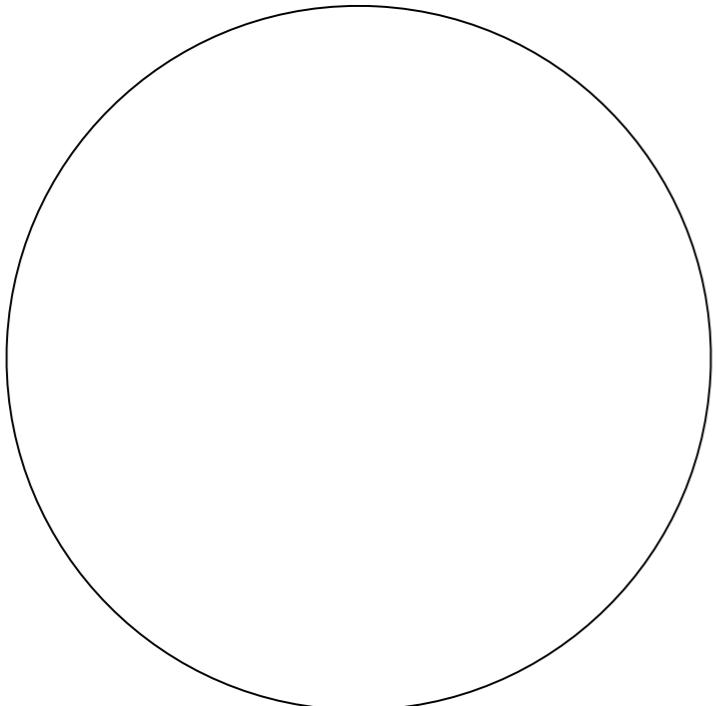
The remaining question is how to update  $\mathbf{C}$ .



# Covariance Matrix Adaptation

## Rank-One Update

$$\textcolor{red}{m} \leftarrow \textcolor{red}{m} + \sigma \mathbf{y}_w, \quad \mathbf{y}_w = \sum_{i=1}^{\mu} \textcolor{blue}{w}_i \mathbf{y}_{i:\lambda}, \quad \mathbf{y}_i \sim \mathcal{N}_i(\mathbf{0}, \textcolor{red}{C})$$



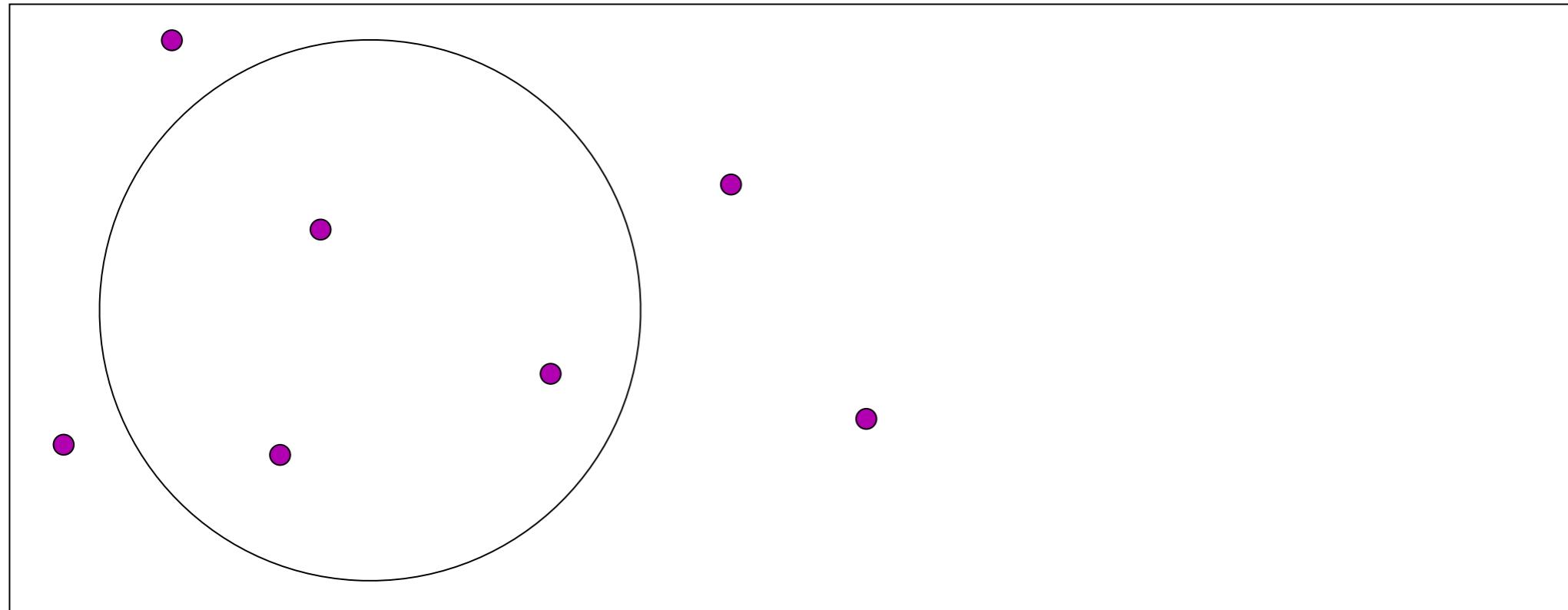
initial distribution,  $\textcolor{red}{C} = \mathbf{I}$

... equations

# Covariance Matrix Adaptation

## Rank-One Update

$$\mathbf{m} \leftarrow \mathbf{m} + \sigma \mathbf{y}_w, \quad \mathbf{y}_w = \sum_{i=1}^{\mu} \mathbf{w}_i \mathbf{y}_{i:\lambda}, \quad \mathbf{y}_i \sim \mathcal{N}_i(\mathbf{0}, \mathbf{C})$$



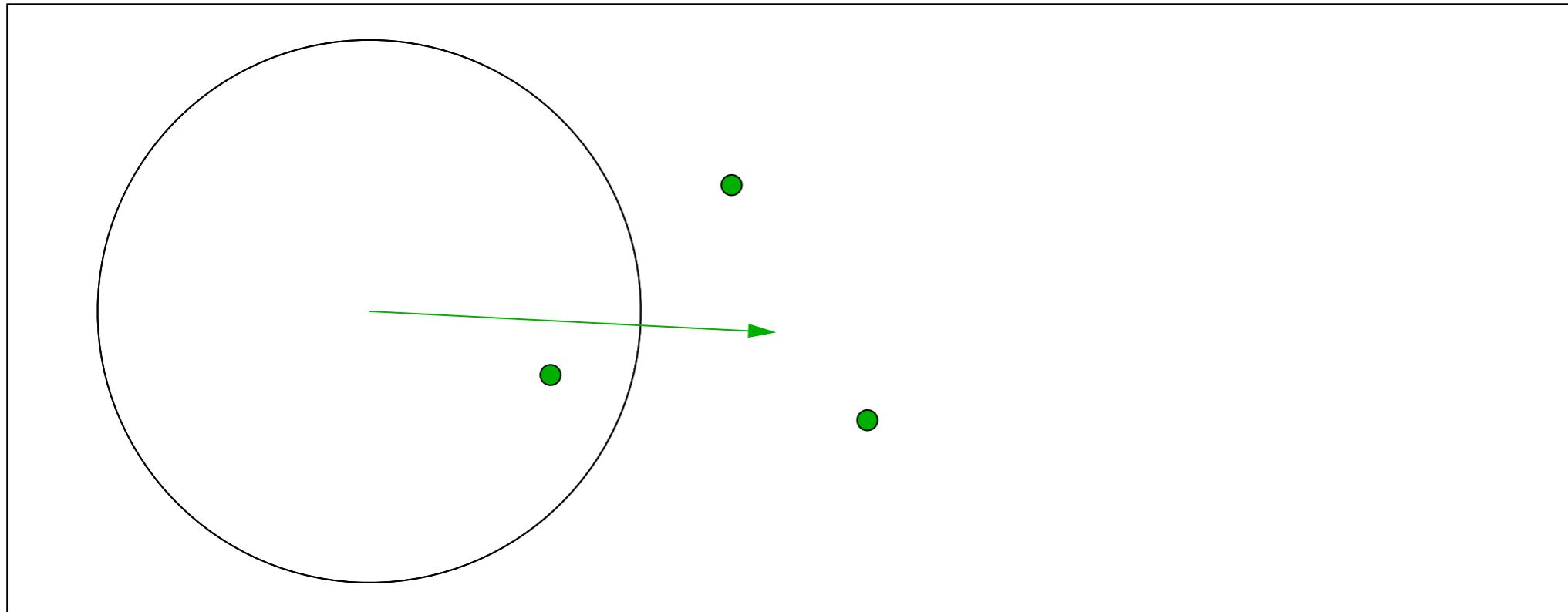
initial distribution,  $\mathbf{C} = \mathbf{I}$

... equations

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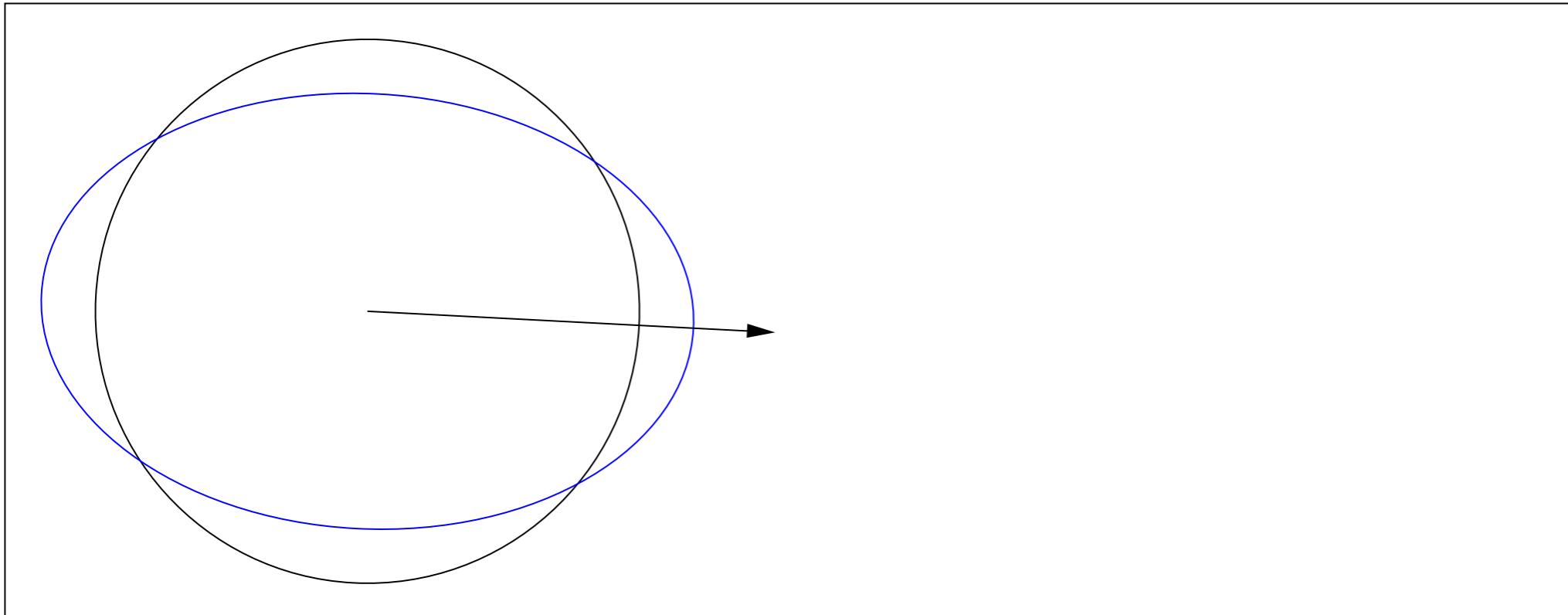
$\mathbf{y}_w$ , movement of the population mean  $\mathbf{m}$  (disregarding  $\sigma$ )

... equations

# Covariance Matrix Adaptation

## Rank-One Update

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mixture of distribution  $\mathbf{C}$  and step  $\mathbf{y}_w$ ,

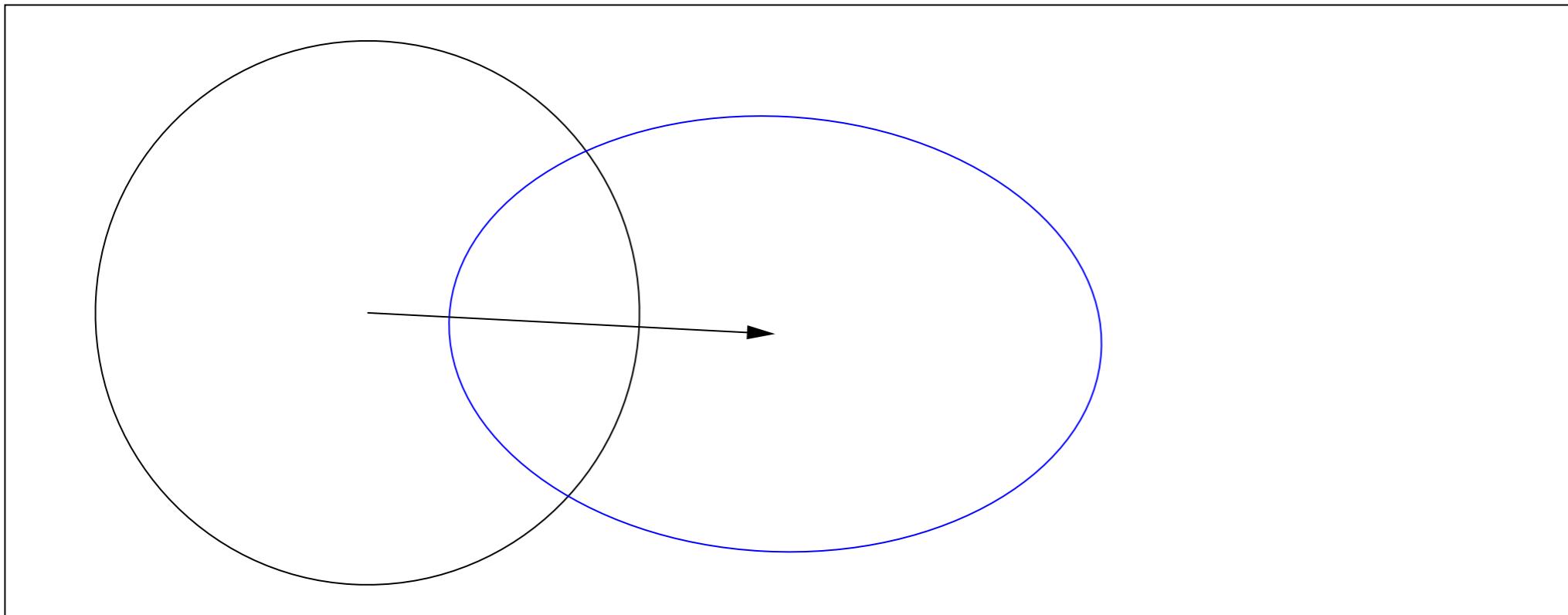
$$\mathbf{C} \leftarrow 0.8 \times \mathbf{C} + 0.2 \times \mathbf{y}_w \mathbf{y}_w^T$$

... equations

# Covariance Matrix Adaptation

## Rank-One Update

$$\mathbf{m} \leftarrow \mathbf{m} + \sigma \mathbf{y}_w, \quad \mathbf{y}_w = \sum_{i=1}^{\mu} \mathbf{w}_i \mathbf{y}_{i:\lambda}, \quad \mathbf{y}_i \sim \mathcal{N}_i(\mathbf{0}, \mathbf{C})$$



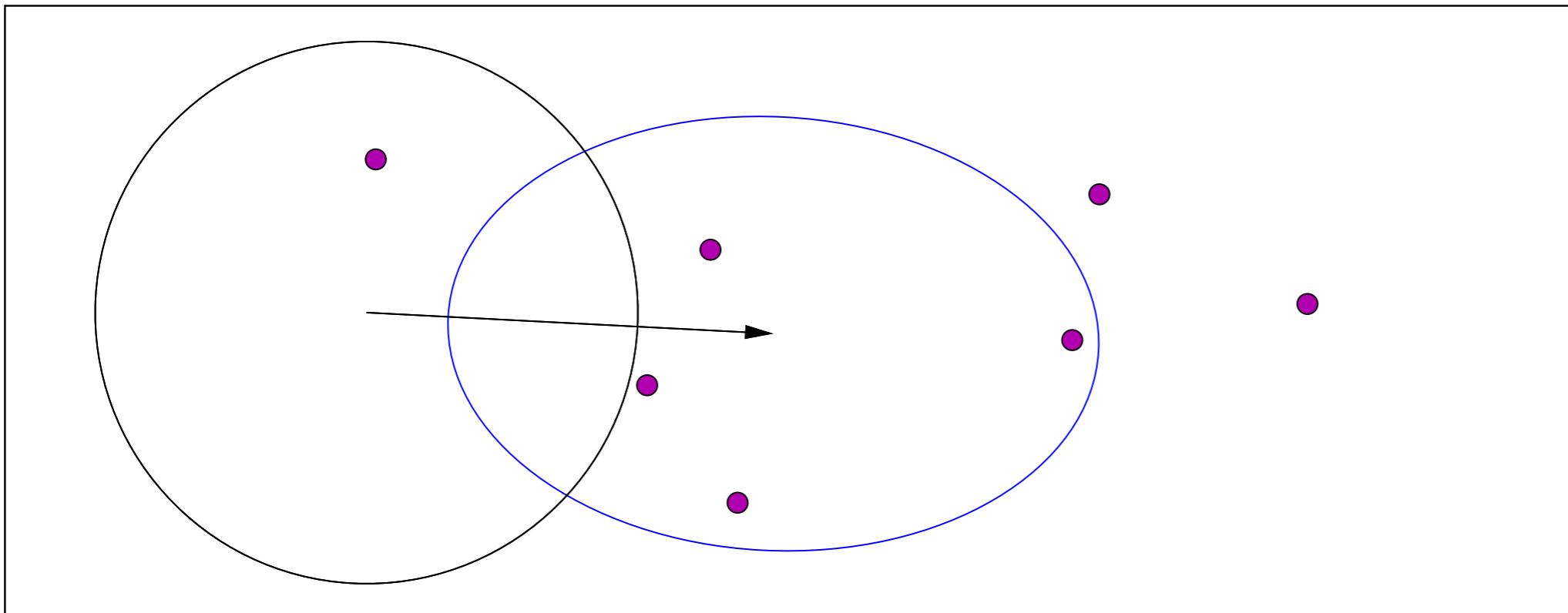
new distribution (disregarding  $\sigma$ )

... equations

# Covariance Matrix Adaptation

## Rank-One Update

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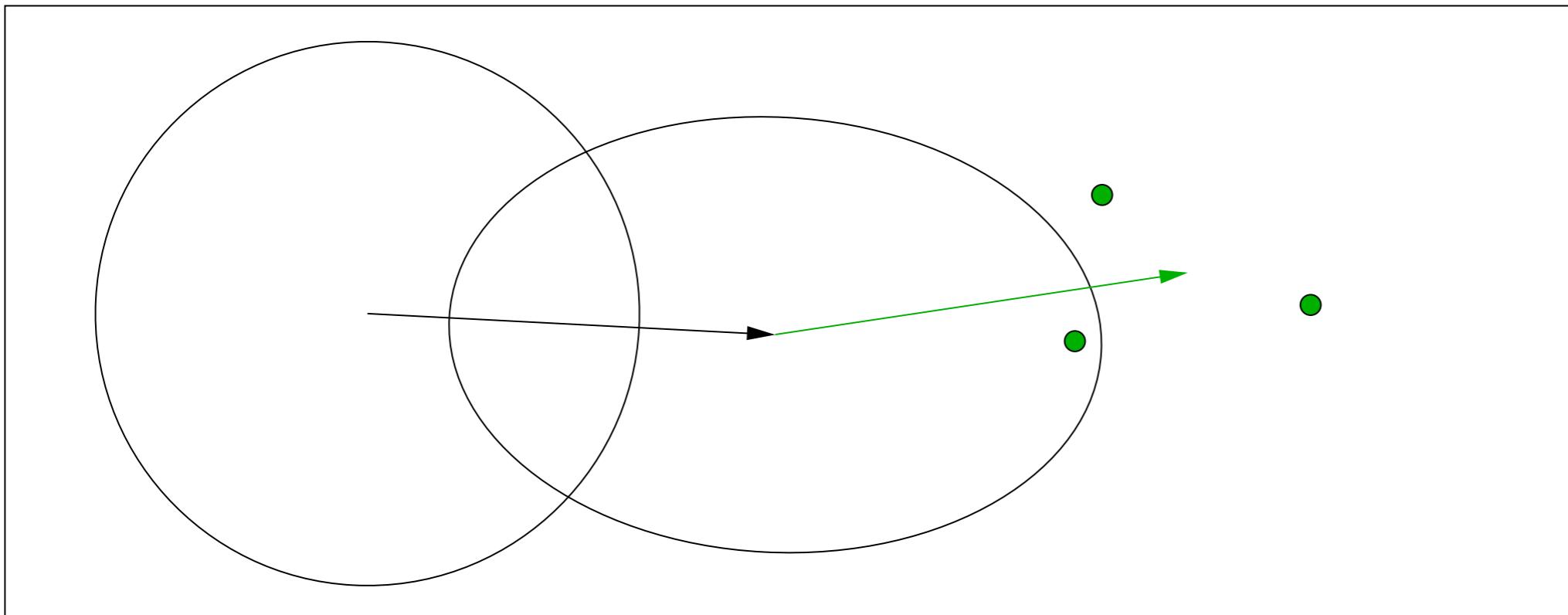
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... equations

# Covariance Matrix Adaptation

## Rank-One Update

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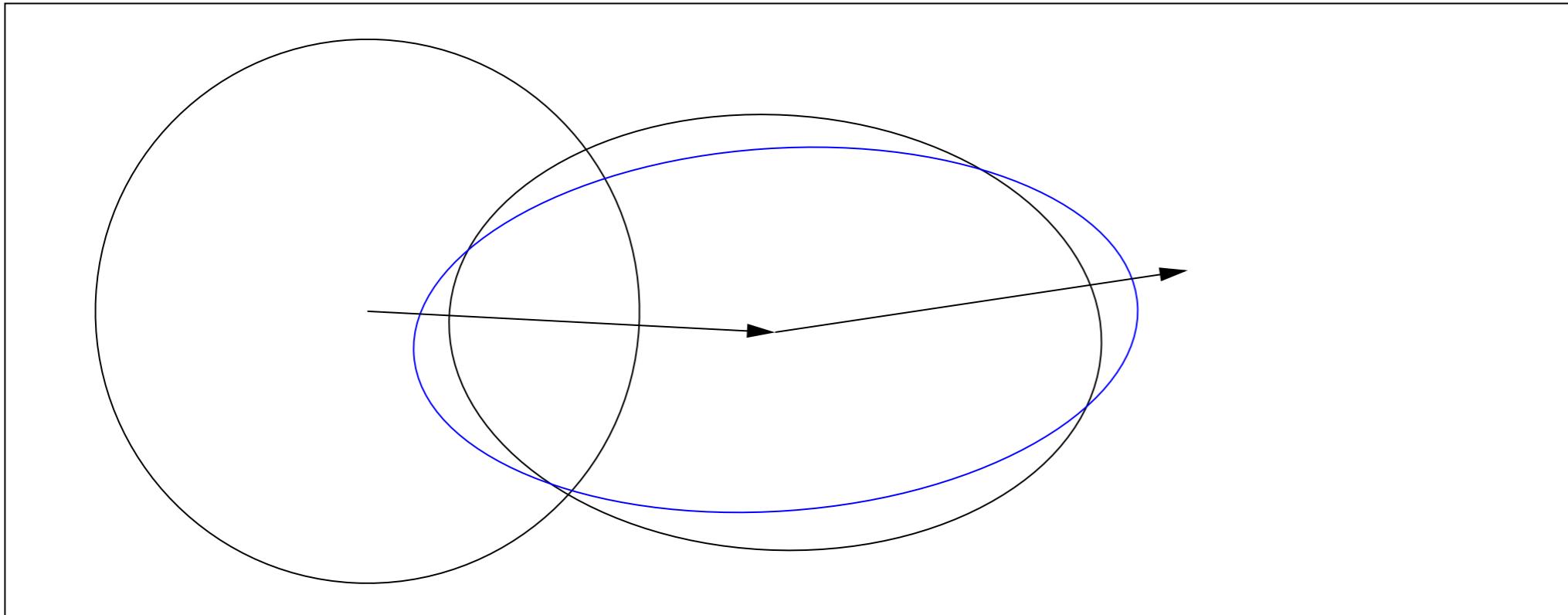
movement of the population mean  $\mathbf{m}$

... equations

# Covariance Matrix Adaptation

## Rank-One Update

$$\mathbf{m} \leftarrow \mathbf{m} + \sigma \mathbf{y}_w, \quad \mathbf{y}_w = \sum_{i=1}^{\mu} \mathbf{w}_i \mathbf{y}_{i:\lambda}, \quad \mathbf{y}_i \sim \mathcal{N}_i(\mathbf{0}, \mathbf{C})$$



mixture of distribution  $\mathbf{C}$  and step  $\mathbf{y}_w$ ,

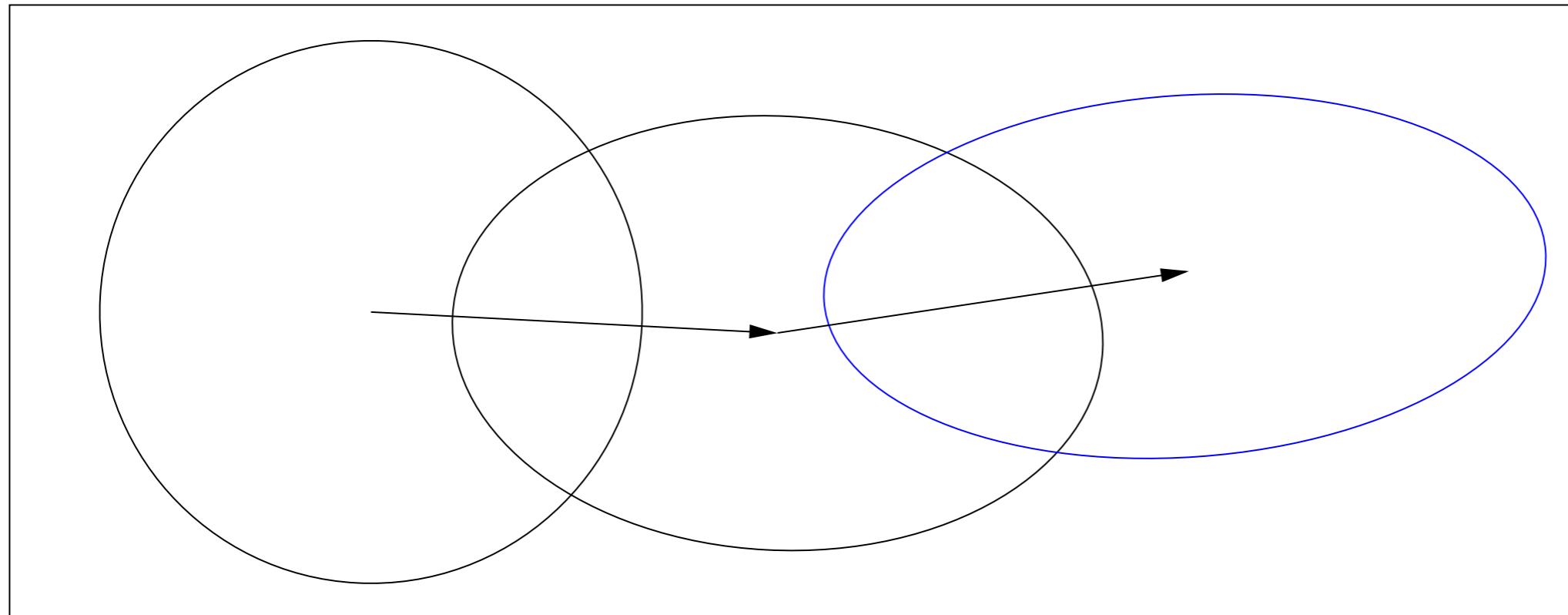
$$\mathbf{C} \leftarrow 0.8 \times \mathbf{C} + 0.2 \times \mathbf{y}_w \mathbf{y}_w^T$$

... equations

# Covariance Matrix Adaptation

## Rank-One Update

$$\textcolor{violet}{m} \leftarrow \textcolor{violet}{m} + \sigma y_w, \quad y_w = \sum_{i=1}^{\mu} \textcolor{blue}{w}_i y_{i:\lambda}, \quad y_i \sim \mathcal{N}_i(\mathbf{0}, \mathbf{C})$$



## new distribution,

$$\mathbf{C} \leftarrow 0.8 \times \mathbf{C} + 0.2 \times \mathbf{y}_w \mathbf{y}_w^T$$

the ruling principle: the adaptation **increases** the likelihood of successful steps,  $y_w$ , to appear again

another viewpoint: the adaptation follows a natural gradient approximation of the expected fitness

## ... equations

# Covariance Matrix Adaptation

## Rank-One Update

Initialize  $\mathbf{m} \in \mathbb{R}^n$ , and  $\mathbf{C} = \mathbf{I}$ , set  $\sigma = 1$ , learning rate  $c_{\text{cov}} \approx 2/n^2$

While not terminate

$$\mathbf{x}_i = \mathbf{m} + \sigma \mathbf{y}_i, \quad \mathbf{y}_i \sim \mathcal{N}_i(\mathbf{0}, \mathbf{C}),$$

$$\mathbf{m} \leftarrow \mathbf{m} + \sigma \mathbf{y}_w \quad \text{where } \mathbf{y}_w = \sum_{i=1}^{\mu} \mathbf{w}_i \mathbf{y}_{i:\lambda}$$

$$\mathbf{C} \leftarrow (1 - c_{\text{cov}}) \mathbf{C} + c_{\text{cov}} \mu_w \underbrace{\mathbf{y}_w \mathbf{y}_w^T}_{\text{rank-one}} \quad \text{where } \mu_w = \frac{1}{\sum_{i=1}^{\mu} \mathbf{w}_i^2} \geq 1$$

The rank-one update has been found independently in several domains<sup>6 7 8 9</sup>

<sup>6</sup> Kjellström&Taxén 1981. Stochastic Optimization in System Design, IEEE TCS

<sup>7</sup> Hansen&Ostermeier 1996. Adapting arbitrary normal mutation distributions in evolution strategies: The covariance matrix adaptation, ICEC

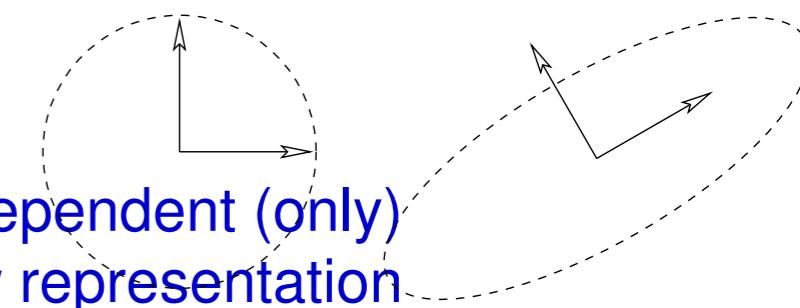
<sup>8</sup> Ljung 1999. System Identification: Theory for the User

<sup>9</sup> Haario et al 2001. An adaptive Metropolis algorithm, JSTOR

$$\mathbf{C} \leftarrow (1 - c_{\text{cov}}) \mathbf{C} + c_{\text{cov}} \mu_w \mathbf{y}_w \mathbf{y}_w^T$$

# covariance matrix adaptation

- learns all **pairwise dependencies** between variables  
off-diagonal entries in the covariance matrix reflect the dependencies
  - conducts a **principle component analysis** (PCA) of steps  $y_w$ , sequentially in time and space  
eigenvectors of the covariance matrix  $\mathbf{C}$  are the principle components / the principle axes of the mutation ellipsoid
  - learns a new **rotated problem representation**  
components are independent (only) in the new representation
  - learns a **new** (Mahalanobis) **metric**  
variable metric method
  - approximates the **inverse Hessian** on quadratic functions  
transformation into the sphere function
  - for  $\mu = 1$ : conducts a **natural gradient ascent** on the distribution  $\mathcal{N}$   
entirely independent of the given coordinate system



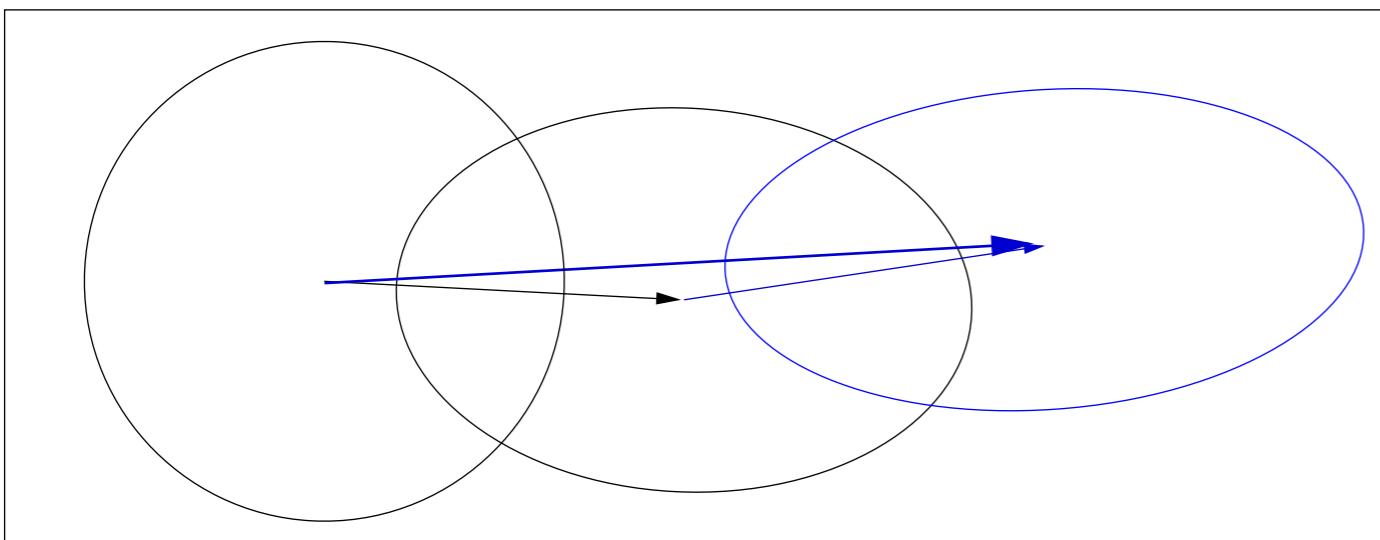
The Evolution Path  
or  
Cumulation

# Cumulation

## The Evolution Path

### Evolution Path

Conceptually, the evolution path is the **search path** the strategy takes **over a number of generation steps**. It can be expressed as a sum of consecutive *steps* of the mean  $\mathbf{m}$ .



An exponentially weighted sum of steps  $\mathbf{y}_w$  is used

$$\mathbf{p}_c \propto \sum_{i=0}^g \underbrace{(1 - c_c)^{g-i}}_{\text{exponentially fading weights}} \mathbf{y}_w^{(i)}$$

The recursive construction of the evolution path (cumulation):

$$\mathbf{p}_c \leftarrow \underbrace{(1 - c_c)}_{\text{decay factor}} \mathbf{p}_c + \underbrace{\sqrt{1 - (1 - c_c)^2} \sqrt{\mu_w}}_{\text{normalization factor}} \underbrace{\mathbf{y}_w}_{\text{input} = \frac{\mathbf{m} - \mathbf{m}_{\text{old}}}{\sigma}}$$

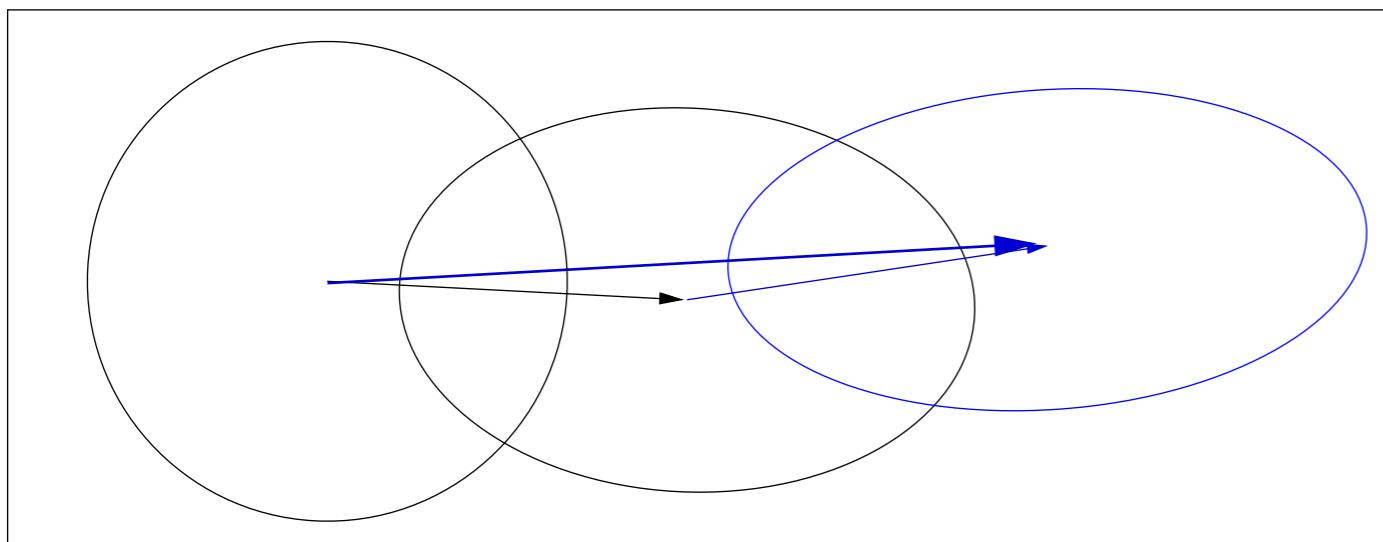
where  $\mu_w = \frac{1}{\sum w_i^2}$ ,  $c_c \ll 1$ . History information is accumulated in the evolution path.

# Cumulation

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“Cumulation” is a widely used technique and also know as

- *exponential smoothing* in time series, forecasting
- exponentially weighted *mooving average*
- *iterate averaging* in stochastic approximation
- *momentum* in the back-propagation algorithm for ANNs
- ...

“Cumulation” conducts a *low-pass filtering*, but there is more to it...

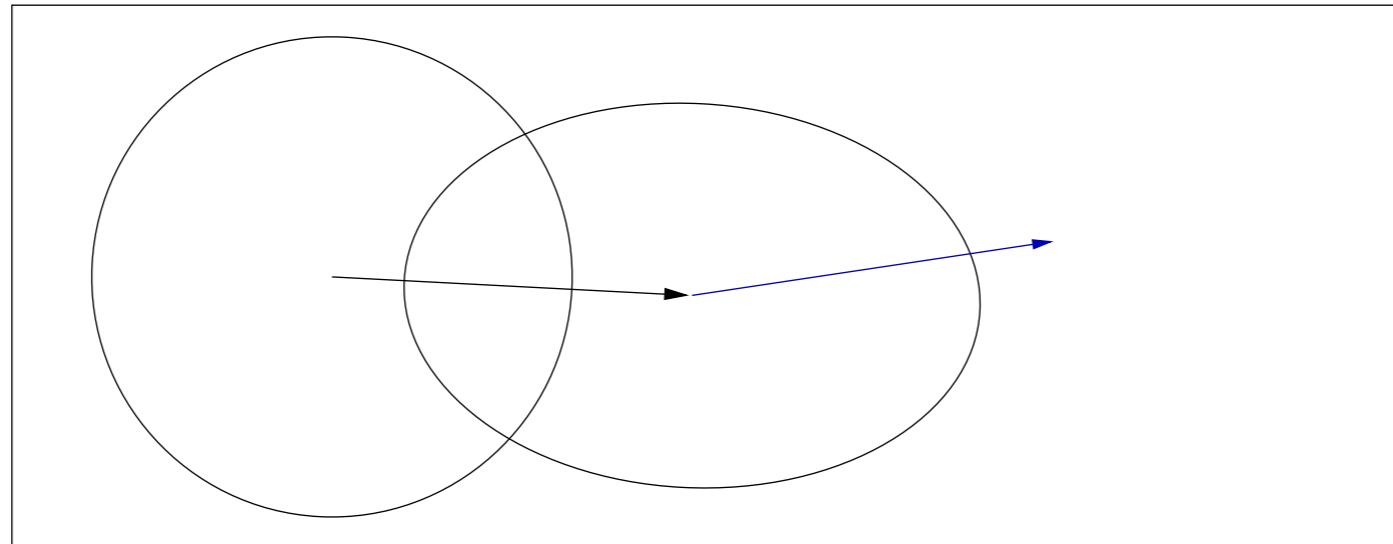
... why?

# Cumulation

$$\mathbf{C} \leftarrow (1 - c_{\text{cov}})\mathbf{C} + c_{\text{cov}}\mu_w \mathbf{y}_w \mathbf{y}_w^T$$

## Utilizing the Evolution Path

We used  $\mathbf{y}_w \mathbf{y}_w^T$  for updating  $\mathbf{C}$ . Because  $\mathbf{y}_w \mathbf{y}_w^T = -\mathbf{y}_w (-\mathbf{y}_w)^T$  the sign of  $\mathbf{y}_w$  is lost.



The **sign information** (signifying correlation *between steps*) is (re-)introduced by using the *evolution path*.

$$\begin{aligned} \mathbf{p}_c &\leftarrow \underbrace{(1 - c_c)}_{\text{decay factor}} \mathbf{p}_c + \underbrace{\sqrt{1 - (1 - c_c)^2} \sqrt{\mu_w}}_{\text{normalization factor}} \mathbf{y}_w \\ \mathbf{C} &\leftarrow (1 - c_{\text{cov}})\mathbf{C} + c_{\text{cov}} \underbrace{\mathbf{p}_c \mathbf{p}_c^T}_{\text{rank-one}} \end{aligned}$$

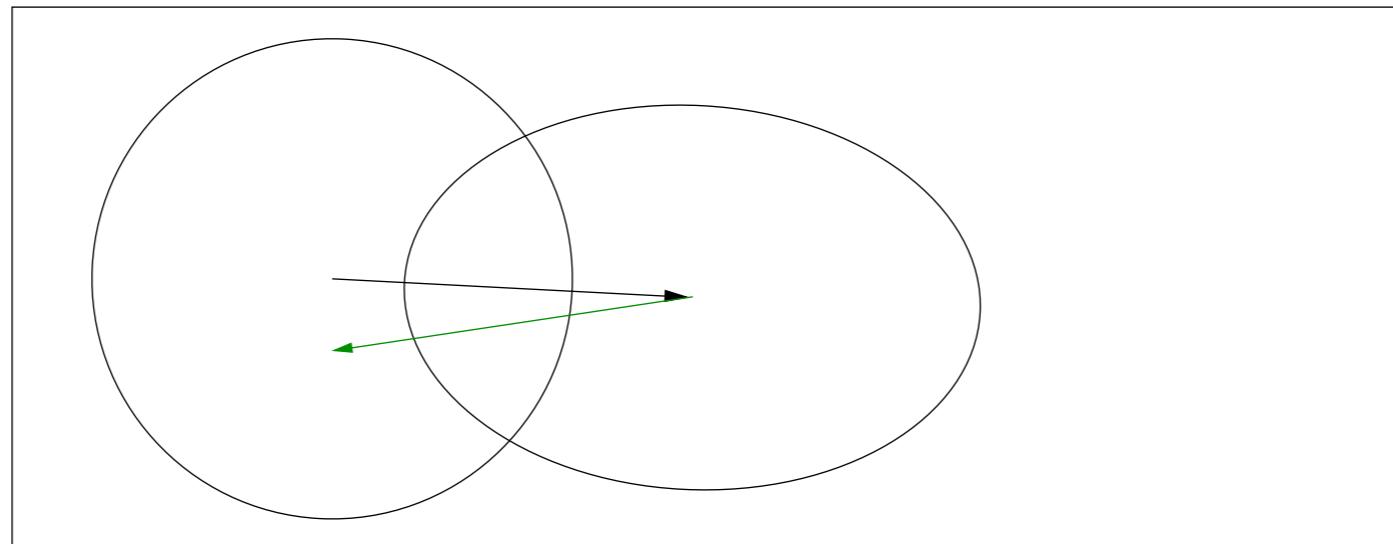
where  $\mu_w = \frac{1}{\sum w_i^2}$ ,  $c_{\text{cov}} \ll c_c \ll 1$  such that  $1/c_c$  is the “backward time horizon”.

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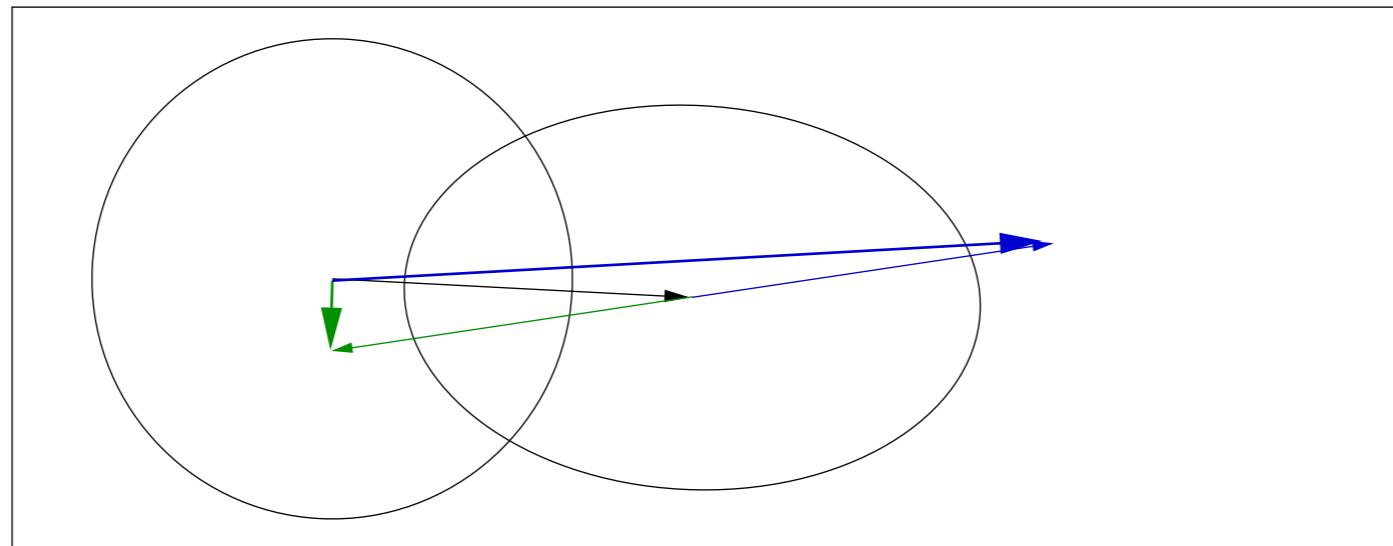
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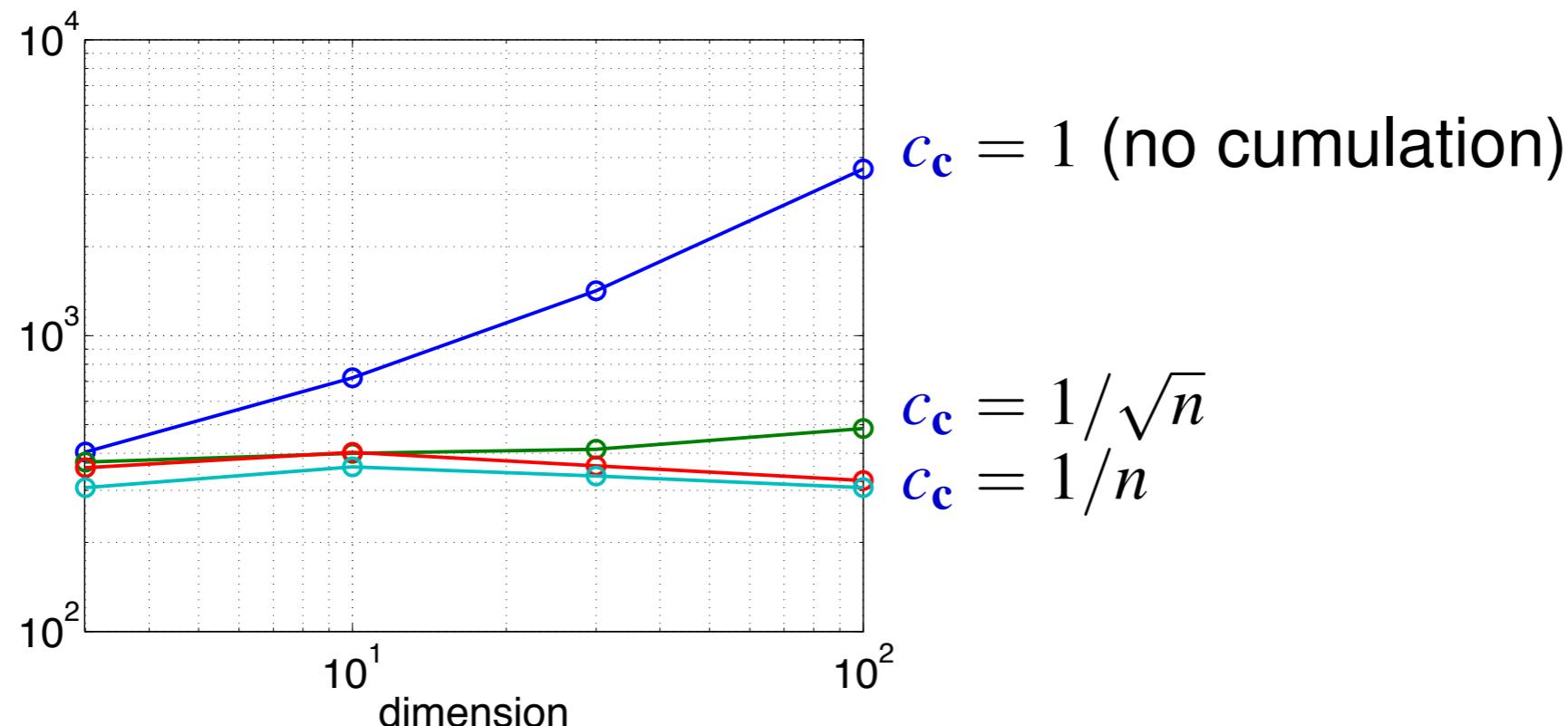
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Using an **evolution path** for the **rank-one update** of the covariance matrix reduces the number of function evaluations to adapt to a straight ridge from about  $\mathcal{O}(n^2)$  to  $\mathcal{O}(n)$ .<sup>(a)</sup>

<sup>a</sup>Hansen & Auger 2013. Principled design of continuous stochastic search: From theory to practice.

Number of  $f$ -evaluations divided by dimension on the cigar function  $f(\mathbf{x}) = x_1^2 + 10^6 \sum_{i=2}^n x_i^2$



The overall model complexity is  $n^2$  but important parts of the model can be learned in time of order  $n$

# Rank- $\mu$ Update

$$\begin{aligned} \mathbf{x}_i &= \mathbf{m} + \sigma \mathbf{y}_i, & \mathbf{y}_i &\sim \mathcal{N}_i(\mathbf{0}, \mathbf{C}), \\ \mathbf{m} &\leftarrow \mathbf{m} + \sigma \mathbf{y}_w & \mathbf{y}_w &= \sum_{i=1}^{\mu} w_i \mathbf{y}_{i:\lambda} \end{aligned}$$

The rank- $\mu$  update extends the update rule for **large population sizes**  $\lambda$  using  $\mu > 1$  vectors to update  $\mathbf{C}$  at each generation step.

The weighted empirical covariance matrix

$$\mathbf{C}_\mu = \sum_{i=1}^{\mu} w_i \mathbf{y}_{i:\lambda} \mathbf{y}_{i:\lambda}^T$$

computes a weighted mean of the outer products of the best  $\mu$  steps and has rank  $\min(\mu, n)$  with probability one.

with  $\mu = \lambda$  weights can be negative <sup>10</sup>

The rank- $\mu$  update then reads

$$\mathbf{C} \leftarrow (1 - c_{\text{cov}}) \mathbf{C} + c_{\text{cov}} \mathbf{C}_\mu$$

where  $c_{\text{cov}} \approx \mu_w/n^2$  and  $c_{\text{cov}} \leq 1$ .

<sup>10</sup> Jastrebski and Arnold (2006). Improving evolution strategies through active covariance matrix adaptation. *ES*.

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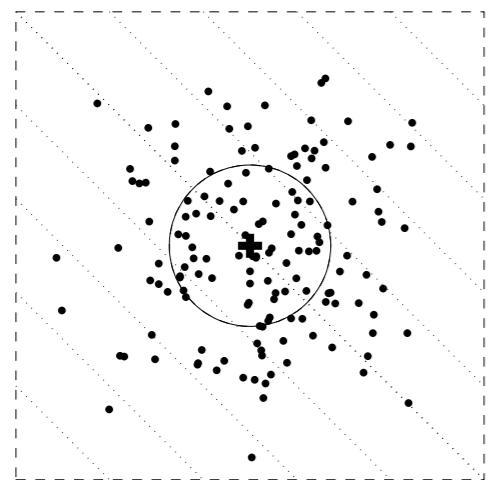
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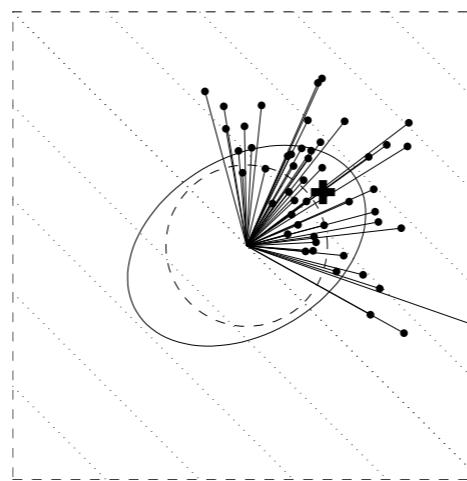
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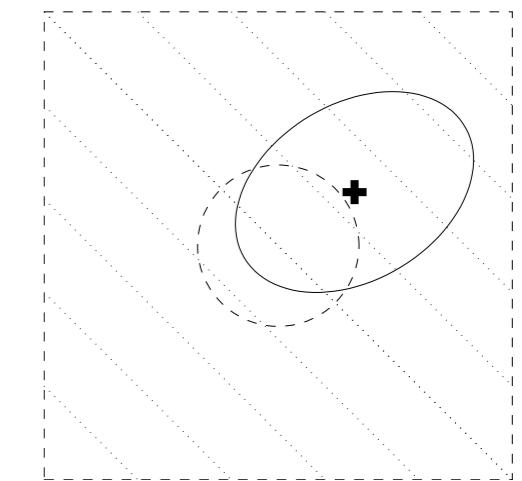
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$$\begin{aligned} \mathbf{C}_\mu &= \frac{1}{\mu} \sum \mathbf{y}_{i:\lambda} \mathbf{y}_{i:\lambda}^T \\ \mathbf{C} &\leftarrow (1 - 1) \times \mathbf{C} + 1 \times \mathbf{C}_\mu \end{aligned}$$



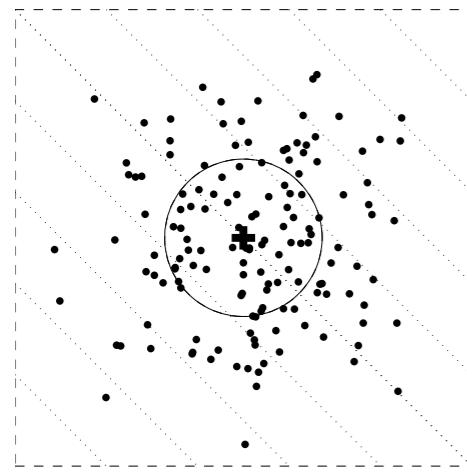
$$\mathbf{m}_{\text{new}} \leftarrow \mathbf{m} + \frac{1}{\mu} \sum \mathbf{y}_{i:\lambda}$$

sampling of  $\lambda = 150$   
solutions where  
 $\mathbf{C} = \mathbf{I}$  and  $\sigma = 1$

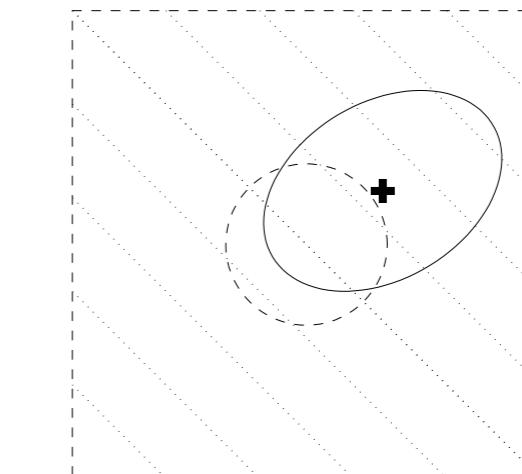
calculating  $\mathbf{C}$  where  
 $\mu = 50$ ,  
 $w_1 = \dots = w_\mu = \frac{1}{\mu}$ ,  
and  $c_{\text{cov}} = 1$

new distribution

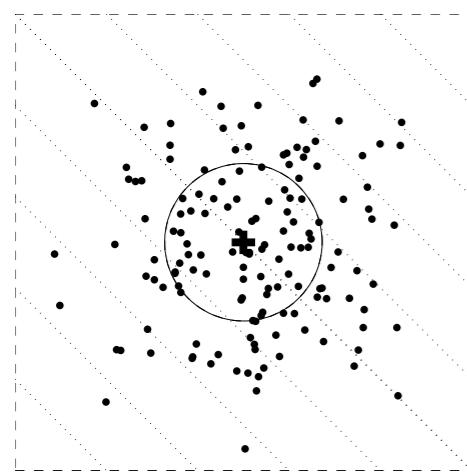
# Rank- $\mu$ CMA versus Estimation of Multivariate Normal Algorithm EMNA<sub>global</sub><sup>11</sup>



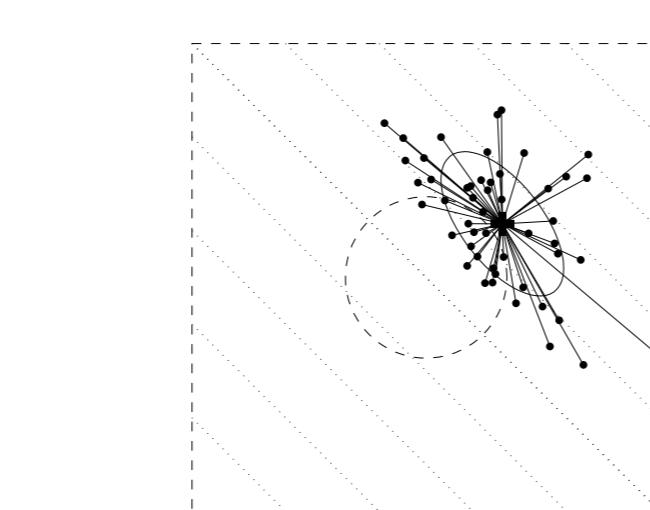
$$x_i = \mathbf{m}_{\text{old}} + \mathbf{y}_i, \quad \mathbf{y}_i \sim \mathcal{N}(\mathbf{0}, \mathbf{C}) \quad \mathbf{C} \leftarrow \frac{1}{\mu} \sum (x_{i:\lambda} - \mathbf{m}_{\text{old}})(x_{i:\lambda} - \mathbf{m}_{\text{old}})^T$$



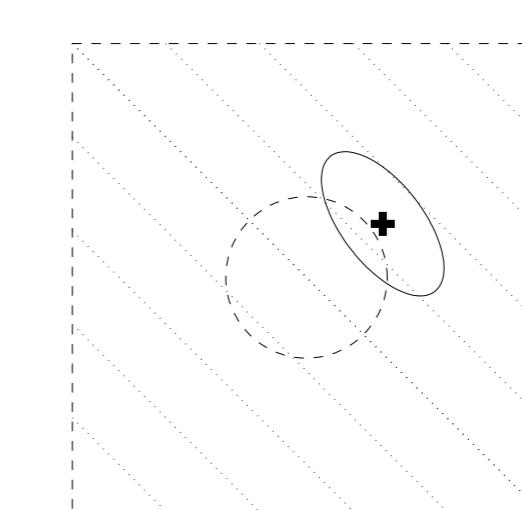
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$$\mathbf{C} \leftarrow \frac{1}{\mu} \sum (x_{i:\lambda} - \mathbf{m}_{\text{new}})(x_{i:\lambda} - \mathbf{m}_{\text{new}})^T$$



$$\mathbf{m}_{\text{new}} = \mathbf{m}_{\text{old}} + \frac{1}{\mu} \sum \mathbf{y}_{i:\lambda}$$

sampling of  $\lambda = 150$  solutions (dots)

calculating  $\mathbf{C}$  from  $\mu = 50$  solutions

$\mathbf{m}_{\text{new}}$  is the minimizer for the variances when calculating  $\mathbf{C}$

rank- $\mu$  CMA  
conducts a  
PCA of  
steps

EMNA<sub>global</sub>  
conducts a  
PCA of  
points

<sup>11</sup> Hansen, N. (2006). The CMA Evolution Strategy: A Comparing Review. In J.A. Lozano, P. Larranga, I. Inza and E. Bengoetxea (Eds.). Towards a new evolutionary computation. Advances in estimation of distribution algorithms. pp. 75-102

## The rank- $\mu$ update

- increases the possible learning rate in large populations  
roughly from  $2/n^2$  to  $\mu_w/n^2$
- can reduce the number of necessary **generations** roughly from  $\mathcal{O}(n^2)$  to  $\mathcal{O}(n)$ <sup>12</sup>  
given  $\mu_w \propto \lambda \propto n$

Therefore the rank- $\mu$  update is the primary mechanism whenever a large population size is used

say  $\lambda \geq 3n + 10$

## The rank-one update

- uses the evolution path and reduces the number of necessary **function evaluations** to learn straight ridges from  $\mathcal{O}(n^2)$  to  $\mathcal{O}(n)$ .

Rank-one update and rank- $\mu$  update can be combined

... all equations

---

<sup>12</sup> Hansen, Müller, and Koumoutsakos 2003. Reducing the Time Complexity of the Derandomized Evolution Strategy with Covariance Matrix Adaptation (CMA-ES). *Evolutionary Computation*, 11(1), pp. 1-18

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# Summary of Equations

The Covariance Matrix Adaptation Evolution Strategy

**Input:**  $\mathbf{m} \in \mathbb{R}^n$ ,  $\sigma \in \mathbb{R}_+$ ,  $\lambda$  (problem dependent)

**Initialize:**  $\mathbf{C} = \mathbf{I}$ , and  $\mathbf{p}_c = \mathbf{0}$ ,  $\mathbf{p}_\sigma = \mathbf{0}$ ,

**Set:**  $c_c \approx 4/n$ ,  $c_\sigma \approx 4/n$ ,  $c_1 \approx 2/n^2$ ,  $c_\mu \approx \mu_w/n^2$ ,  $c_1 + c_\mu \leq 1$ ,  $d_\sigma \approx 1 + \sqrt{\frac{\mu_w}{n}}$ ,  
and  $w_{i=1\dots\lambda}$  such that  $\mu_w = \frac{1}{\sum_{i=1}^\mu w_i^2} \approx 0.3 \lambda$

**While not terminate**

$$\mathbf{x}_i = \mathbf{m} + \sigma \mathbf{y}_i, \quad \mathbf{y}_i \sim \mathcal{N}_i(\mathbf{0}, \mathbf{C}), \quad \text{for } i = 1, \dots, \lambda \quad \text{sampling}$$

$$\mathbf{m} \leftarrow \sum_{i=1}^\mu w_i \mathbf{x}_{i:\lambda} = \mathbf{m} + \sigma \mathbf{y}_w \quad \text{where } \mathbf{y}_w = \sum_{i=1}^\mu w_i \mathbf{y}_{i:\lambda} \quad \text{update mean}$$

$$\mathbf{p}_c \leftarrow (1 - c_c) \mathbf{p}_c + \mathbf{1}_{\{\|\mathbf{p}_\sigma\| < 1.5\sqrt{n}\}} \sqrt{1 - (1 - c_c)^2} \sqrt{\mu_w} \mathbf{y}_w \quad \text{cumulation for } \mathbf{C}$$

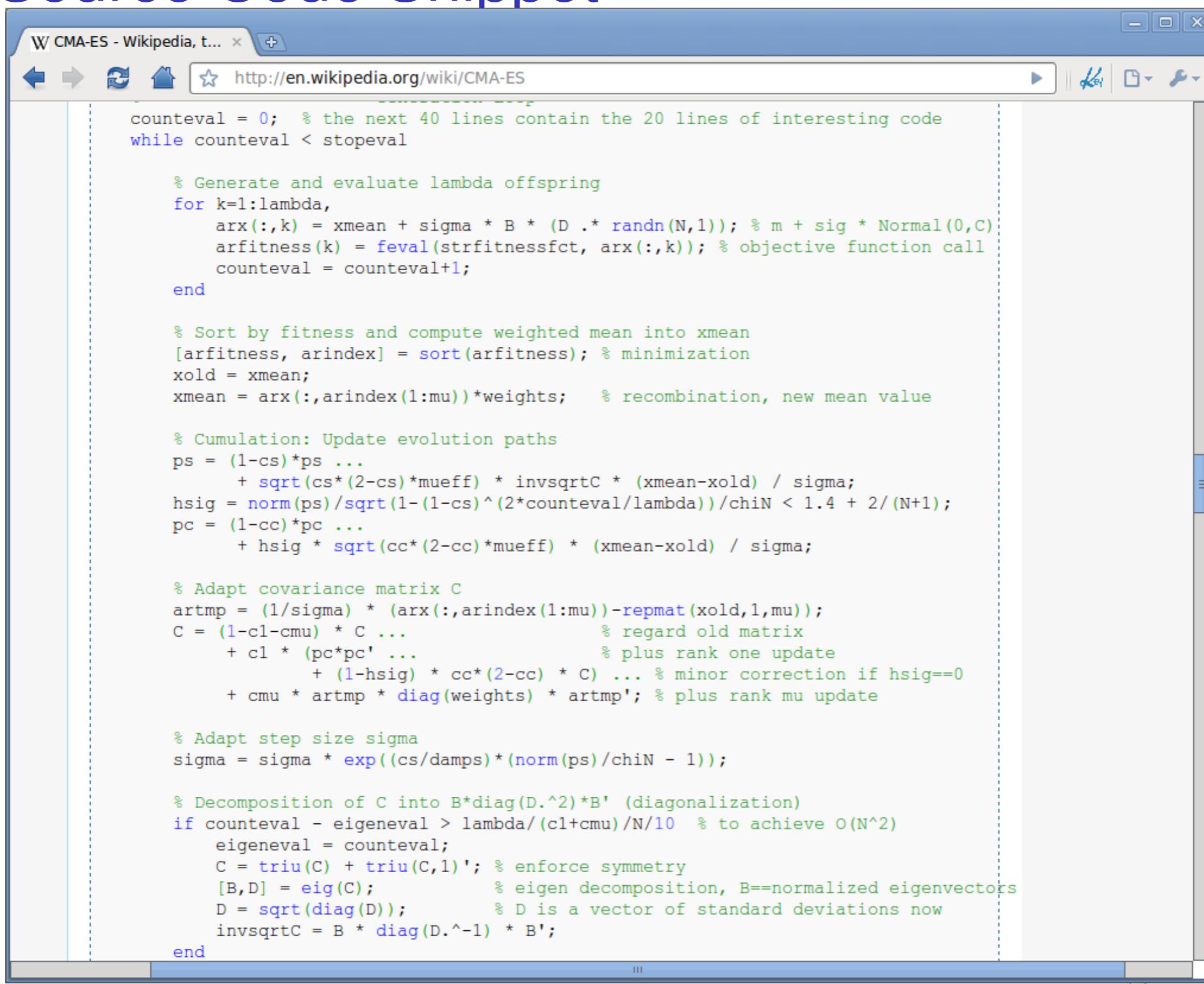
$$\mathbf{p}_\sigma \leftarrow (1 - c_\sigma) \mathbf{p}_\sigma + \sqrt{1 - (1 - c_\sigma)^2} \sqrt{\mu_w} \mathbf{C}^{-\frac{1}{2}} \mathbf{y}_w \quad \text{cumulation for } \sigma$$

$$\mathbf{C} \leftarrow (1 - c_1 - c_\mu) \mathbf{C} + c_1 \mathbf{p}_c \mathbf{p}_c^T + c_\mu \sum_{i=1}^\mu w_i \mathbf{y}_{i:\lambda} \mathbf{y}_{i:\lambda}^T \quad \text{update } \mathbf{C}$$

$$\sigma \leftarrow \sigma \times \exp \left( \frac{c_\sigma}{d_\sigma} \left( \frac{\|\mathbf{p}_\sigma\|}{\mathbb{E} \|\mathcal{N}(\mathbf{0}, \mathbf{I})\|} - 1 \right) \right) \quad \text{update of } \sigma$$

**Not covered** on this slide: termination, restarts, useful output, boundaries and encoding

# Source Code Snippet



The screenshot shows a web browser window with the title "CMA-ES - Wikipedia, t..." and the URL "http://en.wikipedia.org/wiki/CMA-ES". The main content area displays the MATLAB-like pseudocode for the CMA-ES algorithm. The code is color-coded for readability, with comments in green and code in blue.

```

counteval = 0; % the next 40 lines contain the 20 lines of interesting code
while counteval < stopeval

    % Generate and evaluate lambda offspring
    for k=1:lambda,
        arx(:,k) = xmean + sigma * (D .* randn(N,1)); % m + sig * Normal(0,C)
        arfitness(k) = feval(strfitnessfct, arx(:,k)); % objective function call
        counteval = counteval+1;
    end

    % Sort by fitness and compute weighted mean into xmean
    [arfitness, arindex] = sort(arfitness); % minimization
    xold = xmean;
    xmean = arx(:,arindex(1:mu))*weights; % recombination, new mean value

    % Cumulation: Update evolution paths
    ps = (1-cs)*ps ...
        + sqrt(cs*(2-cs)*mueff) * invsqrtC * (xmean-xold) / sigma;
    hsig = norm(ps)/sqrt(1-(1-cs)^(2*counteval/lambda))/chiN < 1.4 + 2/(N+1);
    pc = (1-cc)*pc ...
        + hsig * sqrt(cc*(2-cc)*mueff) * (xmean-xold) / sigma;

    % Adapt covariance matrix C
    artmp = (1/sigma) * (arx(:,arindex(1:mu))-repmat(xold,1,mu));
    C = (1-cl-cmu) * C ... % regard old matrix
        + cl * (pc*pc') ... % plus rank one update
        + (1-hsig) * cc*(2-cc) * C ... % minor correction if hsig==0
        + cmu * artmp * diag(weights) * artmp'; % plus rank mu update

    % Adapt step size sigma
    sigma = sigma * exp((cs/damps)*(norm(ps)/chiN - 1));

    % Decomposition of C into B*diag(D.^2)*B' (diagonalization)
    if counteval - eigeneval > lambda/(cl+cmu)/N/10 % to achieve O(N^2)
        eigeneval = counteval;
        C = triu(C) + triu(C,1)';
        [B,D] = eig(C); % eigen decomposition, B==normalized eigenvectors
        D = sqrt(diag(D)); % D is a vector of standard deviations now
        invsqrtC = B * diag(D.^-1) * B';
    end

```

# Strategy Internal Parameters

- related to selection and recombination
  - ▶  $\lambda$ , offspring number, new solutions sampled, population size
  - ▶  $\mu$ , parent number, solutions involved in updates of  $m$ ,  $C$ , and  $\sigma$
  - ▶  $w_{i=1,\dots,\mu}$ , recombination weights
- related to  $C$ -update
  - ▶  $c_c$ , decay rate for the evolution path
  - ▶  $c_1$ , learning rate for rank-one update of  $C$
  - ▶  $c_\mu$ , learning rate for rank- $\mu$  update of  $C$
- related to  $\sigma$ -update
  - ▶  $c_\sigma$ , decay rate of the evolution path
  - ▶  $d_\sigma$ , damping for  $\sigma$ -change

Parameters were identified in carefully chosen experimental set ups. **Parameters do not in the first place depend on the objective function** and are not meant to be in the users choice.

Only(?) the population size  $\lambda$  (and the initial  $\sigma$ ) might be reasonably varied in a wide range,  
*depending on the objective function*

Useful: restarts with increasing population size (IPOP)

# Experimentum Crucis (0)

What did we want to achieve?

- reduce any convex-quadratic function

$$f(\mathbf{x}) = \mathbf{x}^T \mathbf{H} \mathbf{x}$$

to the sphere model

$$f(\mathbf{x}) = \mathbf{x}^T \mathbf{x}$$

e.g.  $f(\mathbf{x}) = \sum_{i=1}^n 10^{6\frac{i-1}{n-1}} x_i^2$

without use of derivatives

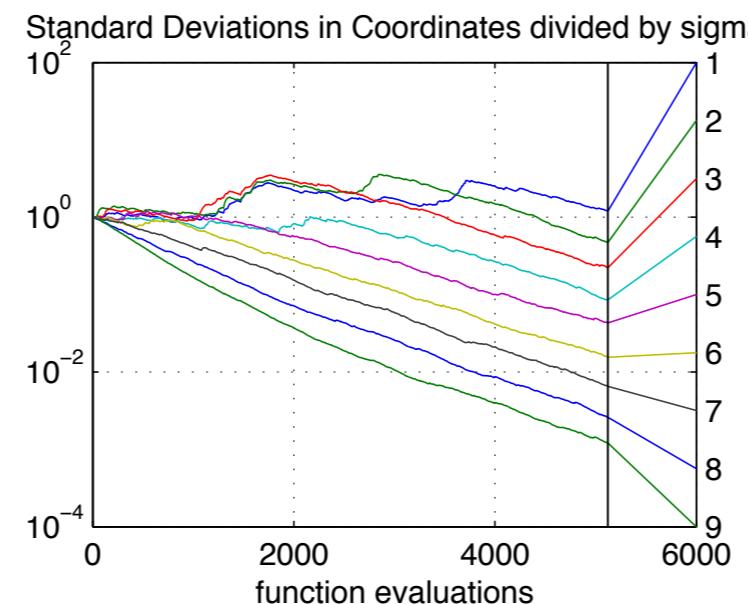
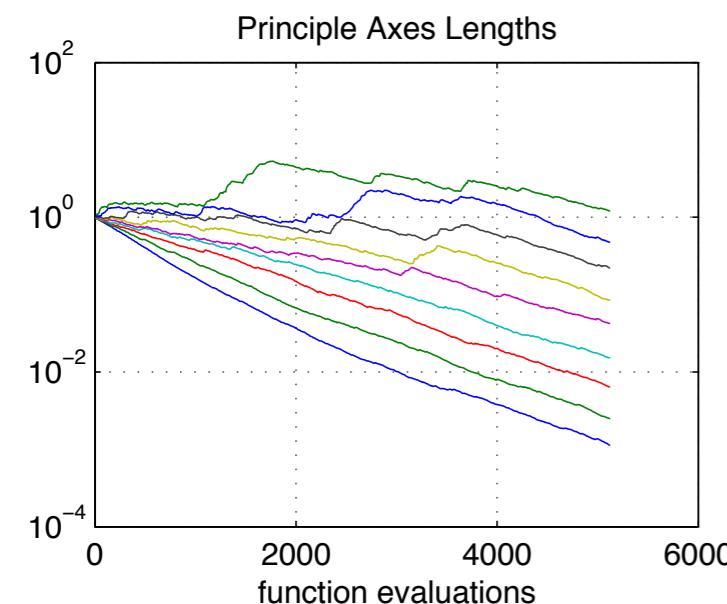
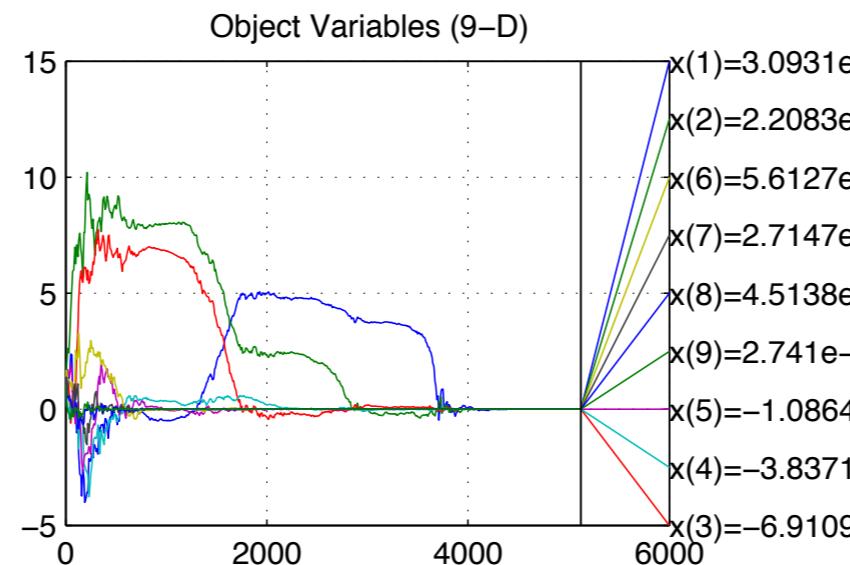
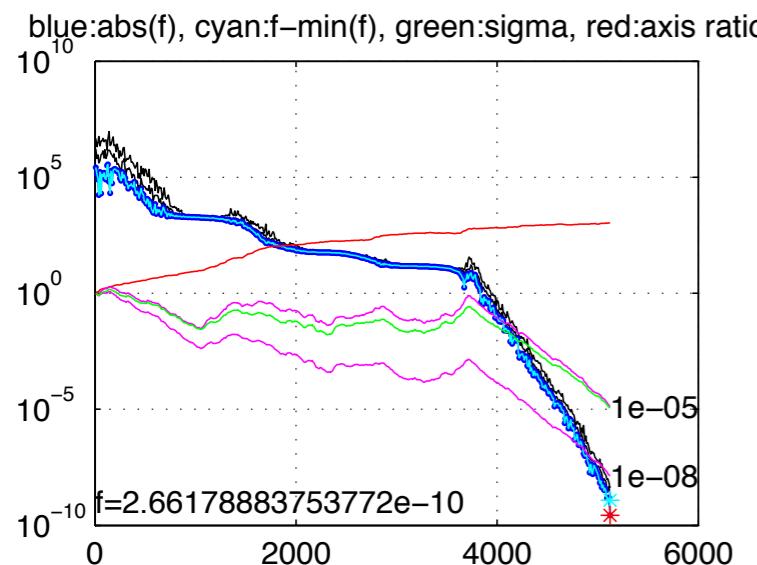
- lines of equal density align with lines of equal fitness

$$\mathbf{C} \propto \mathbf{H}^{-1}$$

in a stochastic sense

# Experimentum Crucis (1)

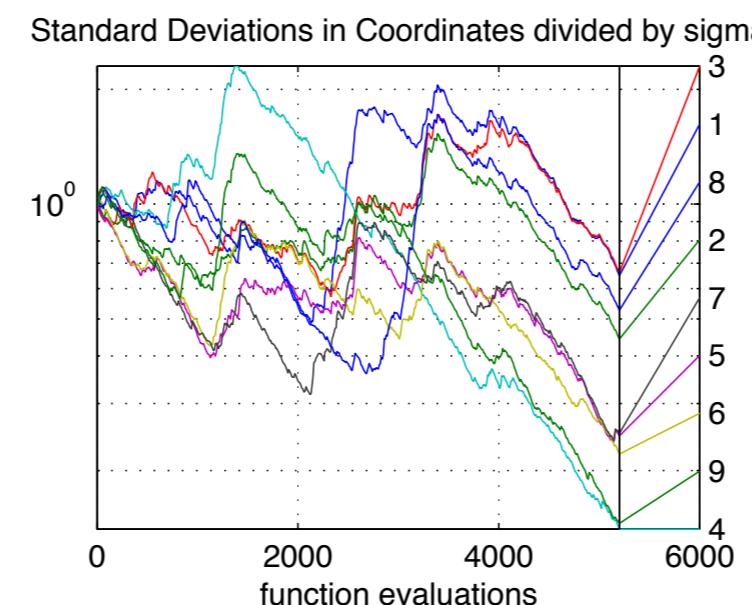
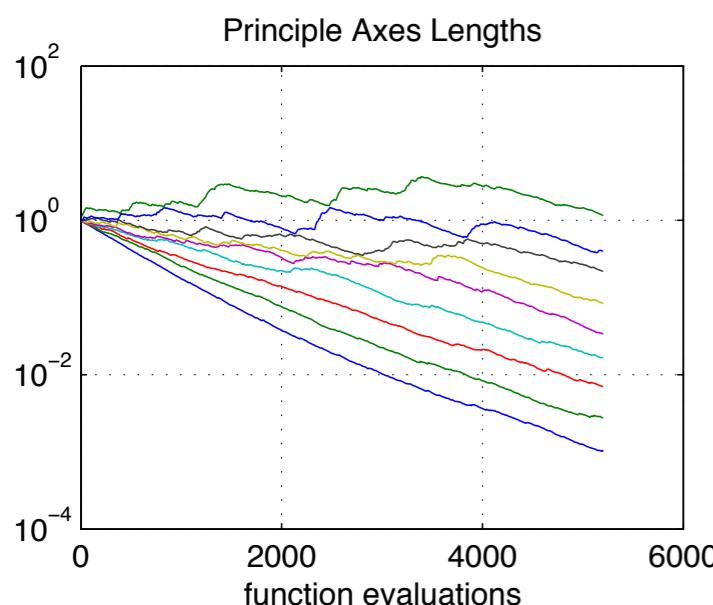
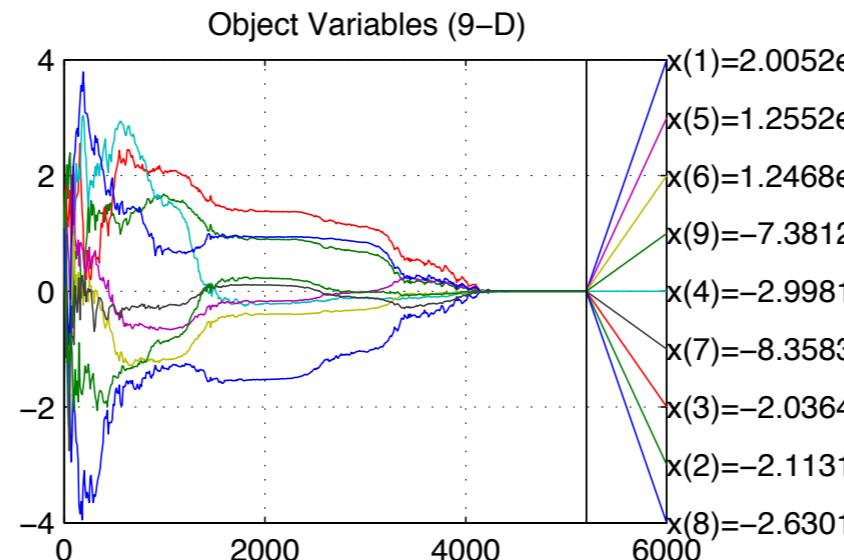
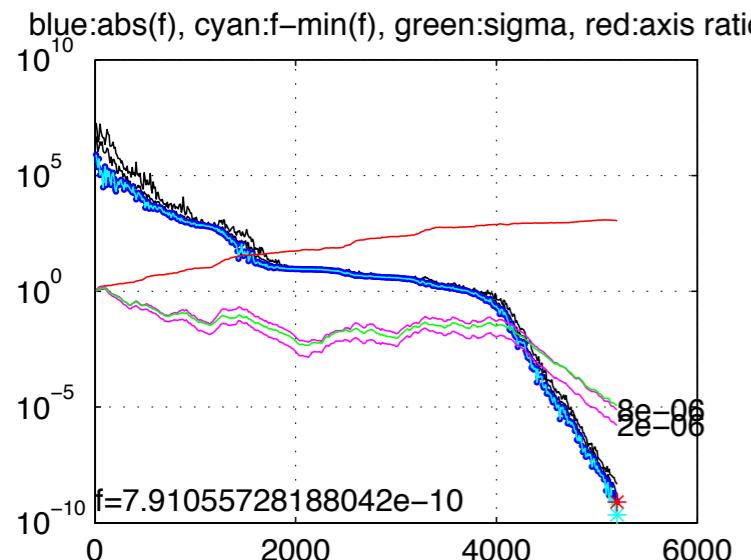
$f$  convex quadratic, separable



$$f(\mathbf{x}) = \sum_{i=1}^n 10^{\alpha \frac{i-1}{n-1}} x_i^2, \alpha = 6$$

# Experimentum Crucis (2)

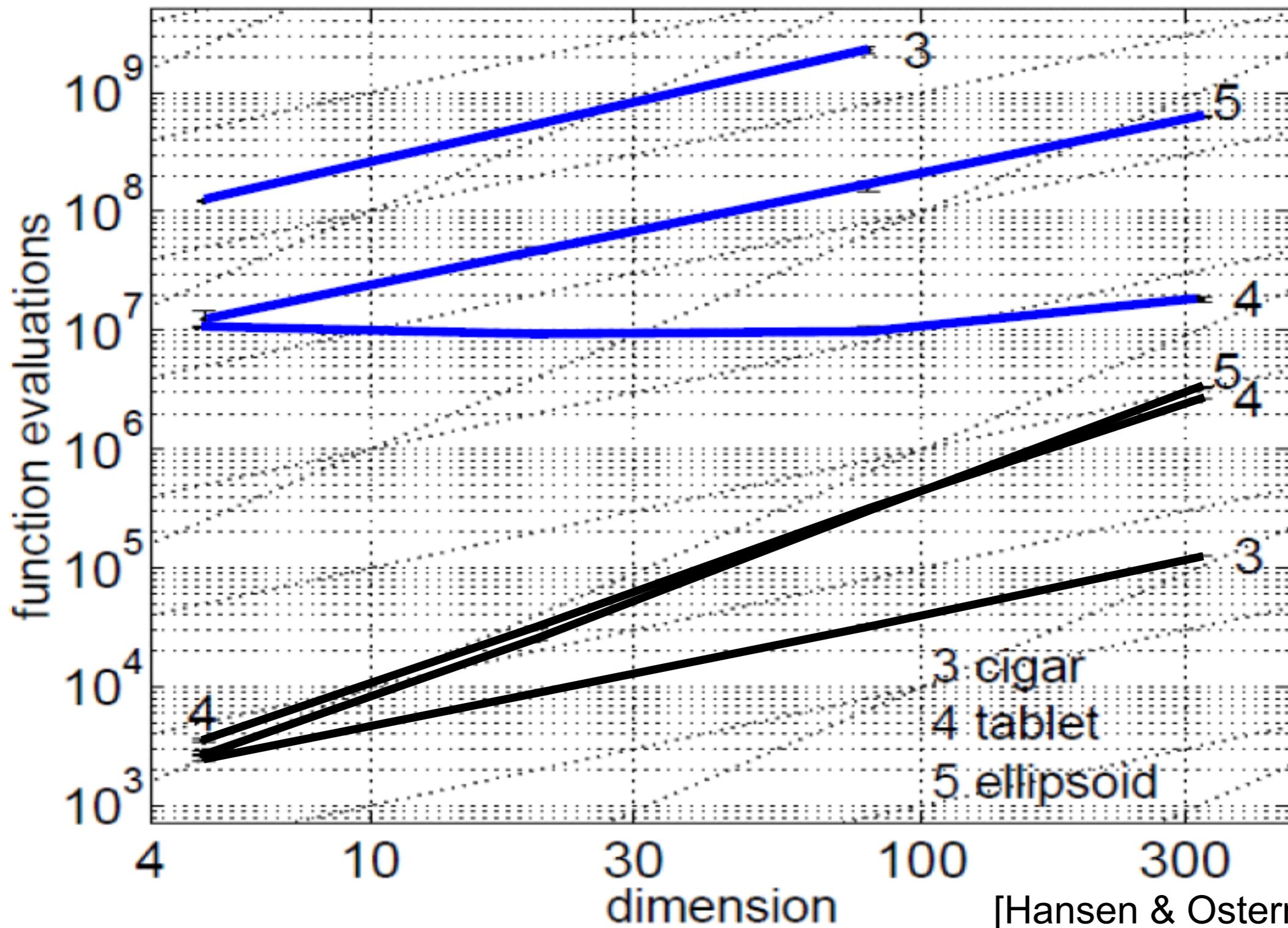
$f$  convex quadratic, as before but non-separable (rotated)



$$\mathbf{C} \propto \mathbf{H}^{-1} \text{ for all } g, \mathbf{H}$$

$$f(\mathbf{x}) = g(\mathbf{x}^T \mathbf{H} \mathbf{x}), g : \mathbb{R} \rightarrow \mathbb{R} \text{ strictly increasing}$$

# Quantifying the enhancement



[Hansen & Ostermeier 2001]

black: CMA-ES ( $c_1 \approx 2/n^2$ ), blue: CSA-ES ( $c_1 = 0$ )

# Theoretical Considerations

# CMA-ES = Natural Evolution Strategy + Cumulation

Natural gradient descend using the MC approximation and the normal distribution

- Rewriting the update of the distribution mean

$$\mathbf{m}_{\text{new}} \leftarrow \sum_{i=1}^{\mu} w_i \mathbf{x}_{i:\lambda} = \mathbf{m} + \underbrace{\sum_{i=1}^{\mu} w_i (\mathbf{x}_{i:\lambda} - \mathbf{m})}_{\text{natural gradient for mean } \frac{\partial}{\partial \mathbf{m}} \widehat{\mathbb{E}}(w \circ P_f(f(\mathbf{x}))) | \mathbf{m}, \mathbf{C}}$$

- Rewriting the update of the covariance matrix<sup>13</sup>

$$\begin{aligned} \mathbf{C}_{\text{new}} &\leftarrow \mathbf{C} + c_1 \overbrace{(\mathbf{p}_c \mathbf{p}_c^T - \mathbf{C})}^{\text{rank one}} \\ &+ \frac{c_\mu}{\sigma^2} \underbrace{\sum_{i=1}^{\mu} w_i \left( \overbrace{(\mathbf{x}_{i:\lambda} - \mathbf{m})(\mathbf{x}_{i:\lambda} - \mathbf{m})^T}^{\text{rank-}\mu} - \sigma^2 \mathbf{C} \right)}_{\text{natural gradient for covariance matrix } \frac{\partial}{\partial \mathbf{C}} \widehat{\mathbb{E}}(w \circ P_f(f(\mathbf{x}))) | \mathbf{m}, \mathbf{C}} \end{aligned}$$

<sup>13</sup> Akimoto et.al. (2010): Bidirectional Relation between CMA Evolution Strategies and Natural Evolution Strategies, PPSN XR Q

# Summary of Equations

The Covariance Matrix Adaptation Evolution Strategy

**Input:**  $\mathbf{m} \in \mathbb{R}^n$ ,  $\sigma \in \mathbb{R}_+$ ,  $\lambda$  (problem dependent)

**Initialize:**  $\mathbf{C} = \mathbf{I}$ , and  $\mathbf{p}_c = \mathbf{0}$ ,  $\mathbf{p}_\sigma = \mathbf{0}$ ,

**Set:**  $c_c \approx 4/n$ ,  $c_\sigma \approx 4/n$ ,  $c_1 \approx 2/n^2$ ,  $c_\mu \approx \mu_w/n^2$ ,  $c_1 + c_\mu \leq 1$ ,  $d_\sigma \approx 1 + \sqrt{\frac{\mu_w}{n}}$ ,  
and  $w_{i=1\dots\lambda}$  such that  $\mu_w = \frac{1}{\sum_{i=1}^\mu w_i^2} \approx 0.3 \lambda$

**While not terminate**

$$\mathbf{x}_i = \mathbf{m} + \sigma \mathbf{y}_i, \quad \mathbf{y}_i \sim \mathcal{N}_i(\mathbf{0}, \mathbf{C}), \quad \text{for } i = 1, \dots, \lambda \quad \text{sampling}$$

$$\mathbf{m} \leftarrow \sum_{i=1}^\mu w_i \mathbf{x}_{i:\lambda} = \mathbf{m} + \sigma \mathbf{y}_w \quad \text{where } \mathbf{y}_w = \sum_{i=1}^\mu w_i \mathbf{y}_{i:\lambda} \quad \text{update mean}$$

$$\mathbf{p}_c \leftarrow (1 - c_c) \mathbf{p}_c + \mathbf{1}_{\{\|\mathbf{p}_\sigma\| < 1.5\sqrt{n}\}} \sqrt{1 - (1 - c_c)^2} \sqrt{\mu_w} \mathbf{y}_w \quad \text{cumulation for } \mathbf{C}$$

$$\mathbf{p}_\sigma \leftarrow (1 - c_\sigma) \mathbf{p}_\sigma + \sqrt{1 - (1 - c_\sigma)^2} \sqrt{\mu_w} \mathbf{C}^{-\frac{1}{2}} \mathbf{y}_w \quad \text{cumulation for } \sigma$$

$$\mathbf{C} \leftarrow (1 - c_1 - c_\mu) \mathbf{C} + c_1 \mathbf{p}_c \mathbf{p}_c^T + c_\mu \sum_{i=1}^\mu w_i \mathbf{y}_{i:\lambda} \mathbf{y}_{i:\lambda}^T \quad \text{update } \mathbf{C}$$

$$\sigma \leftarrow \sigma \times \exp \left( \frac{c_\sigma}{d_\sigma} \left( \frac{\|\mathbf{p}_\sigma\|}{\mathbb{E} \|\mathcal{N}(\mathbf{0}, \mathbf{I})\|} - 1 \right) \right) \quad \text{update of } \sigma$$

**Not covered** on this slide: termination, restarts, useful output, boundaries and encoding

# Maximum Likelihood Update

The new distribution mean  $\mathbf{m}$  maximizes the log-likelihood

$$\mathbf{m}_{\text{new}} = \arg \max_{\mathbf{m}} \sum_{i=1}^{\mu} w_i \log p_{\mathcal{N}}(\mathbf{x}_{i:\lambda} | \mathbf{m})$$

independently of the given covariance matrix

$$\log p_{\mathcal{N}}(\mathbf{x} | \mathbf{m}, \mathbf{C}) = -\frac{1}{2} \log \det(2\pi \mathbf{C}) - \frac{1}{2} (\mathbf{x} - \mathbf{m})^T \mathbf{C}^{-1} (\mathbf{x} - \mathbf{m})$$

$p_{\mathcal{N}}$  is the density of the multi-variate normal distribution

# Maximum Likelihood Update

The new distribution mean  $\mathbf{m}$  maximizes the log-likelihood

$$\mathbf{m}_{\text{new}} = \arg \max_{\mathbf{m}} \sum_{i=1}^{\mu} w_i \log p_{\mathcal{N}}(\mathbf{x}_{i:\lambda} | \mathbf{m})$$

independently of the given covariance matrix

The rank- $\mu$  update matrix  $\mathbf{C}_\mu$  maximizes the log-likelihood

$$\mathbf{C}_\mu = \arg \max_{\mathbf{C}} \sum_{i=1}^{\mu} w_i \log p_{\mathcal{N}} \left( \frac{\mathbf{x}_{i:\lambda} - \mathbf{m}_{\text{old}}}{\sigma} \mid \mathbf{m}_{\text{old}}, \mathbf{C} \right)$$

$$\log p_{\mathcal{N}}(\mathbf{x} | \mathbf{m}, \mathbf{C}) = -\frac{1}{2} \log \det(2\pi\mathbf{C}) - \frac{1}{2} (\mathbf{x} - \mathbf{m})^T \mathbf{C}^{-1} (\mathbf{x} - \mathbf{m})$$

$p_{\mathcal{N}}$  is the density of the multi-variate normal distribution

# Variable Metric

On the function class

$$f(\mathbf{x}) = g \left( \frac{1}{2} (\mathbf{x} - \mathbf{x}^*) \mathbf{H} (\mathbf{x} - \mathbf{x}^*)^T \right)$$

the covariance matrix approximates the inverse Hessian up to a constant factor, that is:

$$\mathbf{C} \propto \mathbf{H}^{-1} \quad (\text{approximately})$$

In effect, ellipsoidal level-sets are transformed into spherical level-sets.

$g : \mathbb{R} \rightarrow \mathbb{R}$  is strictly increasing

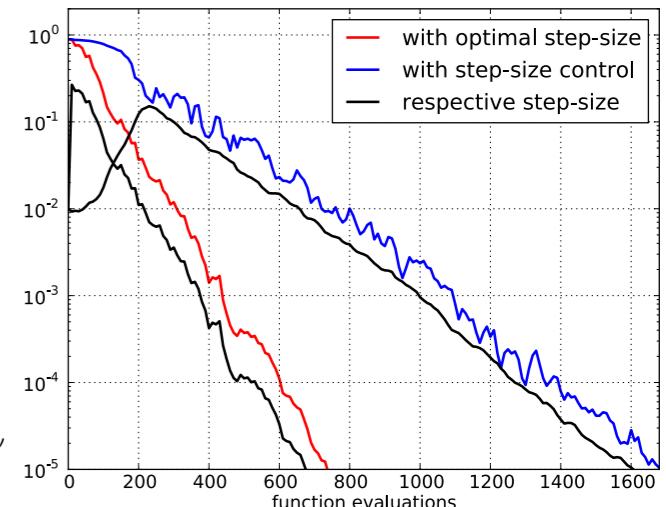
# On Convergence

Evolution Strategies converge with probability one on, e.g.,  $g\left(\frac{1}{2}\mathbf{x}^T \mathbf{H} \mathbf{x}\right)$  like

$$\|\mathbf{m}_k - \mathbf{x}^*\| \propto e^{-ck}, \quad c \leq \frac{0.25}{n}$$

where  $k$  is the number of  $f$ -evaluations. In practice:  $c \approx 0.1/n$

selection	$n \cdot c_{\max}$
(1+1)	0.202
$(\mu/\mu, \lambda)$	0.202
$(\mu/\mu_w, \lambda)$	0.25



Monte Carlo pure random search converges like

$$\|\mathbf{m}_k - \mathbf{x}^*\| \propto k^{-c} = e^{-c \log k}, \quad c = \frac{1}{n}$$

# On the Sphere Function (for $n$ large)

optimal population size  $\lambda$  (offspring)

$$\lambda \not\prec 5 \text{ and } \lambda \not\succ n \quad \text{for } \mu = 1 \text{ we have } \lambda^{\text{opt}} = 5$$

optimal recombination weights

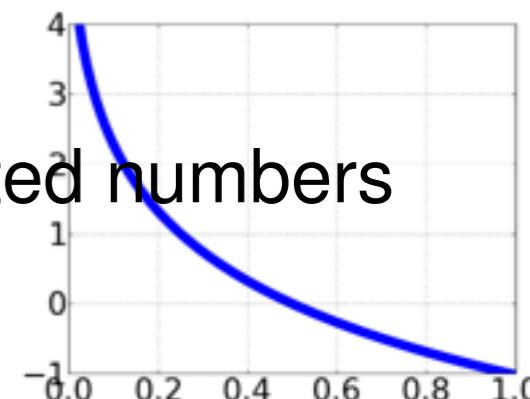
$$w_i^{\text{opt}} \propto -\mathcal{N}_{i:\lambda}(0, 1) \quad i\text{th order statistic of } \lambda \text{ normally distributed numbers}$$

optimal (effective) parent number

$$\mu_w := \frac{1}{\sum_i w_i^2} \leq \mu \quad \text{with } \sum_i |w_i| = 1$$

$$\mu_w^{\text{opt}} \approx 0.32\lambda \quad \text{if } w_i = \max(w_i^{\text{opt}}, 0)$$

$$\mu^{\text{opt}} \approx 0.27\lambda \quad \text{if } w_{i=1\dots\mu} = 1/\mu \text{ (truncation selection)}$$



# On the Sphere Function (for $n$ large)

optimal step-size if  $\mu_w < n$

$$\sigma^{\text{opt}} \propto c_w \frac{\mu_w}{n} \quad \text{where } c_w = -\sum_i w_i \mathcal{N}_{i:\lambda}(0, 1) \approx 1$$

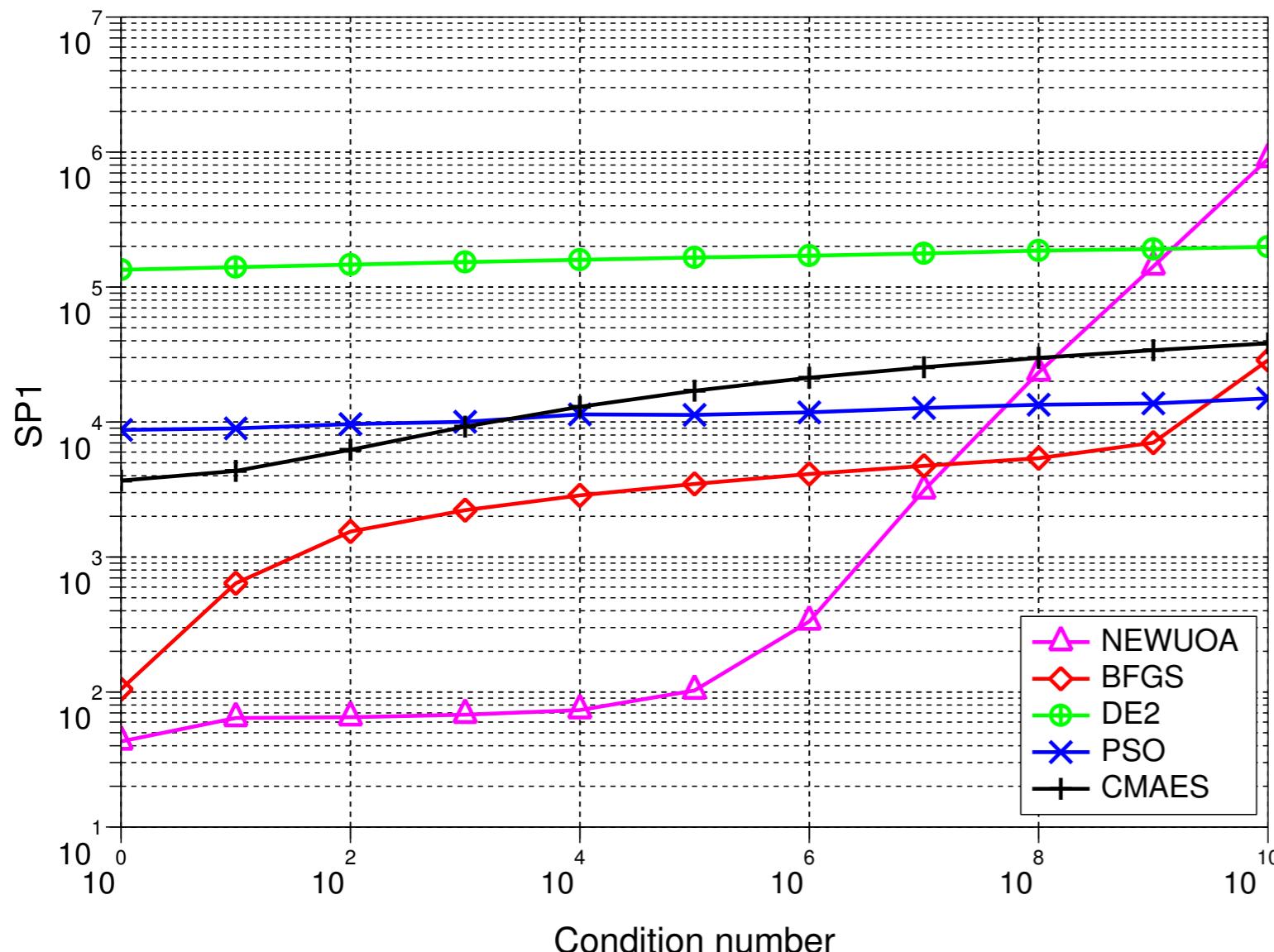
is proportional to  $\mu_w$  and therefore to the population size

# Comparing Experiments

# Comparison to BFGS, NEWUOA, PSO and DE

$f$  convex quadratic, separable with varying condition number  $\alpha$

Ellipsoid dimension 20, 21 trials, tolerance  $1e-09$ , eval max  $1e+07$



- BFGS** (Broyden et al 1970)
  - NEWUOA** (Powell 2004)
  - DE** (Storn & Price 1996)
  - PSO** (Kennedy & Eberhart 1995)
  - CMA-ES (Hansen & Ostermeier 2001)
- $f(\mathbf{x}) = g(\mathbf{x}^T \mathbf{H} \mathbf{x})$  with  
 $\mathbf{H}$  diagonal  
 $g$  identity (for **BFGS** and **NEWUOA**)  
 $g$  any order-preserving = strictly increasing function (for all other)

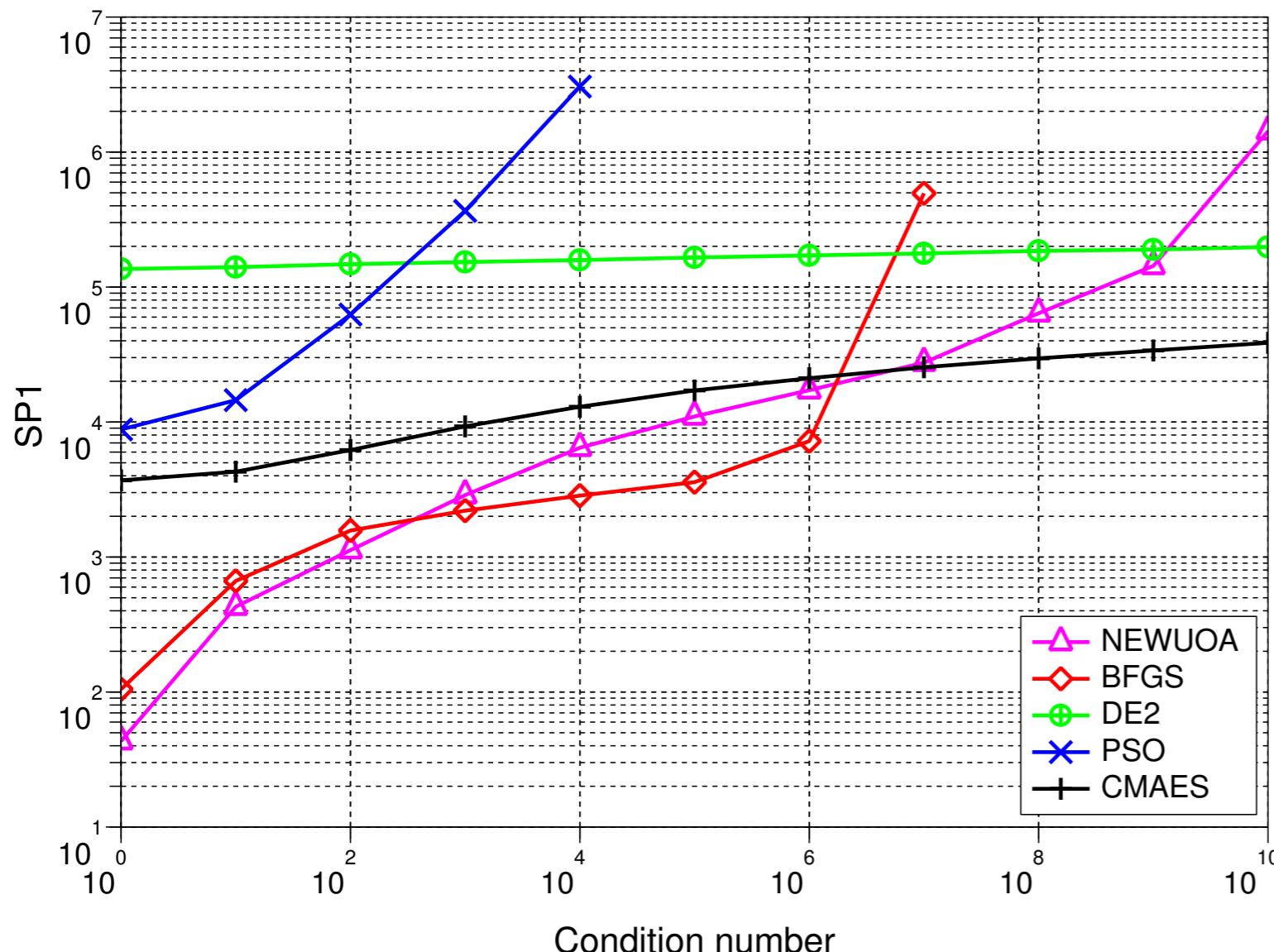
SP1 = average number of objective function evaluations<sup>14</sup> to reach the target function value of  $g^{-1}(10^{-9})$

<sup>14</sup> Auger et.al. (2009): Experimental comparisons of derivative free optimization algorithms, SEA

# Comparison to BFGS, NEWUOA, PSO and DE

$f$  convex quadratic, non-separable (rotated) with varying condition number  $\alpha$

Rotated Ellipsoid dimension 20, 21 trials, tolerance  $1e-09$ , eval max  $1e+07$



- BFGS** (Broyden et al 1970)
  - NEWUOA** (Powell 2004)
  - DE** (Storn & Price 1996)
  - PSO** (Kennedy & Eberhart 1995)
  - CMA-ES (Hansen & Ostermeier 2001)
- $f(\mathbf{x}) = g(\mathbf{x}^T \mathbf{H} \mathbf{x})$  with
- H** full
- $g$  identity (for **BFGS** and **NEWUOA**)
- $g$  any order-preserving = strictly increasing function (for all other)

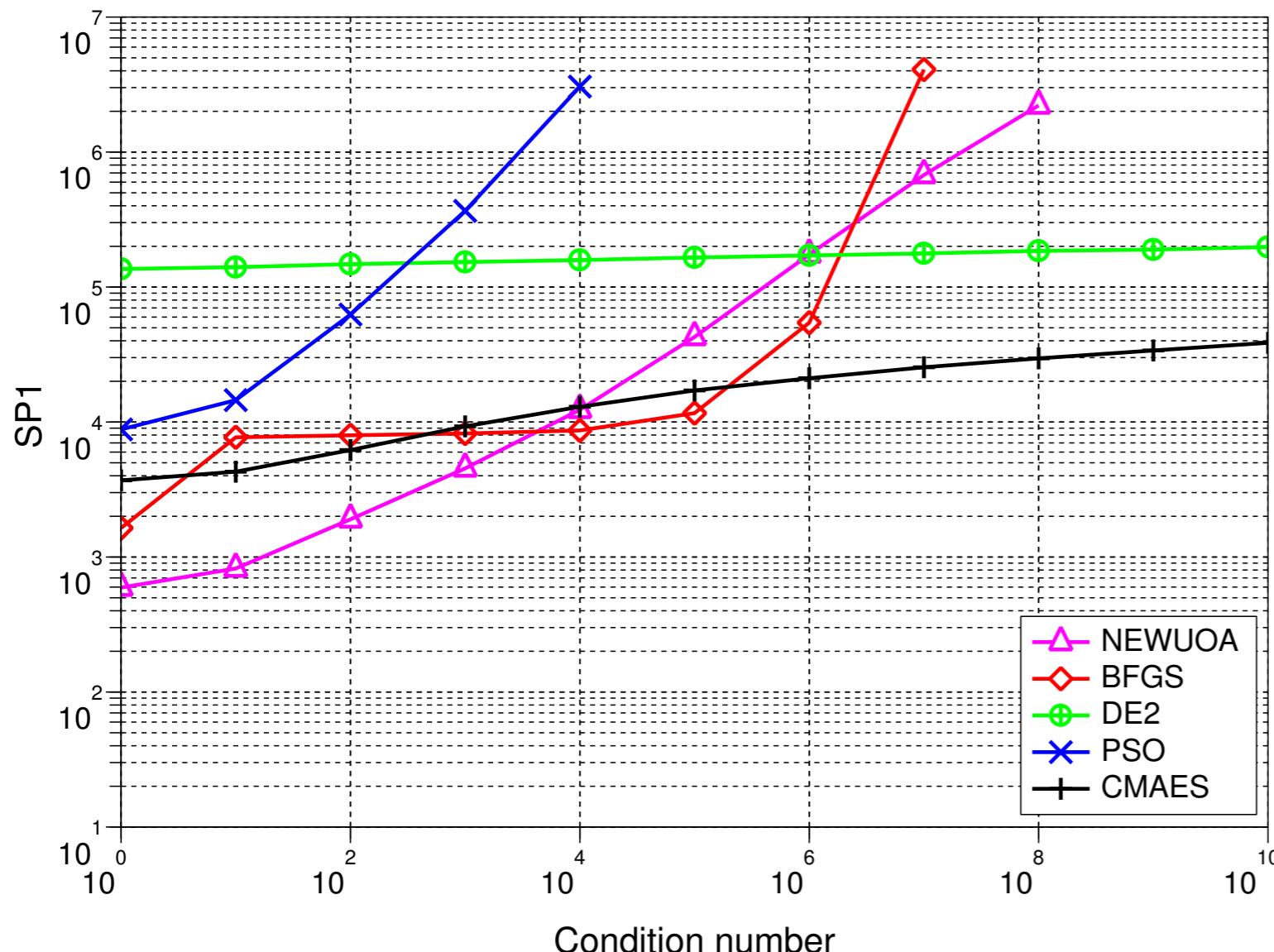
SP1 = average number of objective function evaluations<sup>15</sup> to reach the target function value of  $g^{-1}(10^{-9})$

<sup>15</sup> Auger et.al. (2009): Experimental comparisons of derivative free optimization algorithms, SEA

# Comparison to BFGS, NEWUOA, PSO and DE

$f$  non-convex, non-separable (rotated) with varying condition number  $\alpha$

Sqrt of sqrt of rotated ellipsoid dimension 20, 21 trials, tolerance  $1e-09$ , eval max  $1e+07$



- BFGS** (Broyden et al 1970)
  - NEWUOA** (Powell 2004)
  - DE** (Storn & Price 1996)
  - PSO** (Kennedy & Eberhart 1995)
  - CMA-ES (Hansen & Ostermeier 2001)
- $f(\mathbf{x}) = g(\mathbf{x}^T \mathbf{H} \mathbf{x})$  with  
 $\mathbf{H}$  full
- $g : x \mapsto x^{1/4}$  (for **BFGS** and **NEWUOA**)
- $g$  any order-preserving = strictly increasing function (for all other)

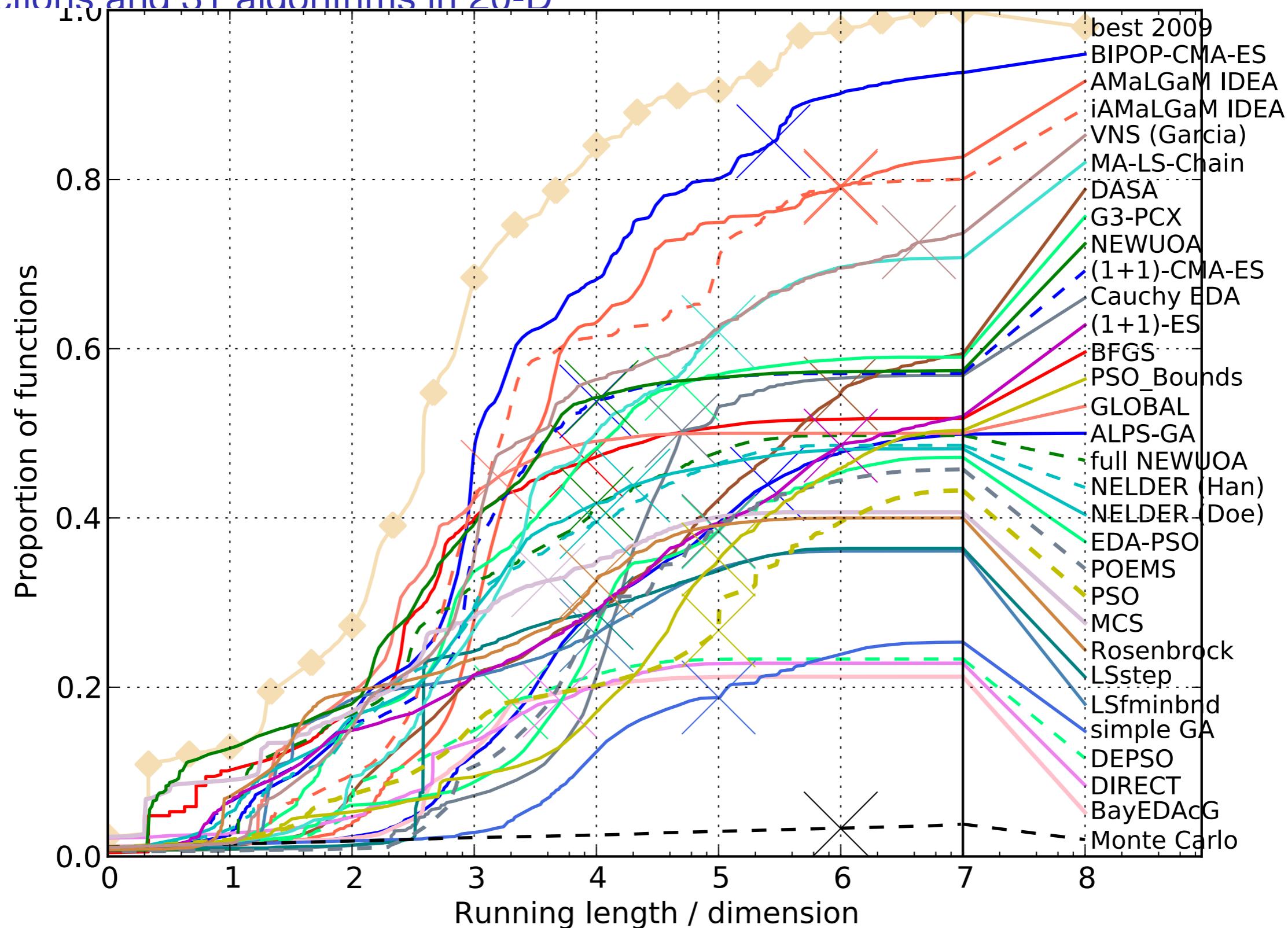
SP1 = average number of objective function evaluations<sup>16</sup> to reach the target function value of  $g^{-1}(10^{-9})$

<sup>16</sup>

Auger et.al. (2009): Experimental comparisons of derivative free optimization algorithms, SEA

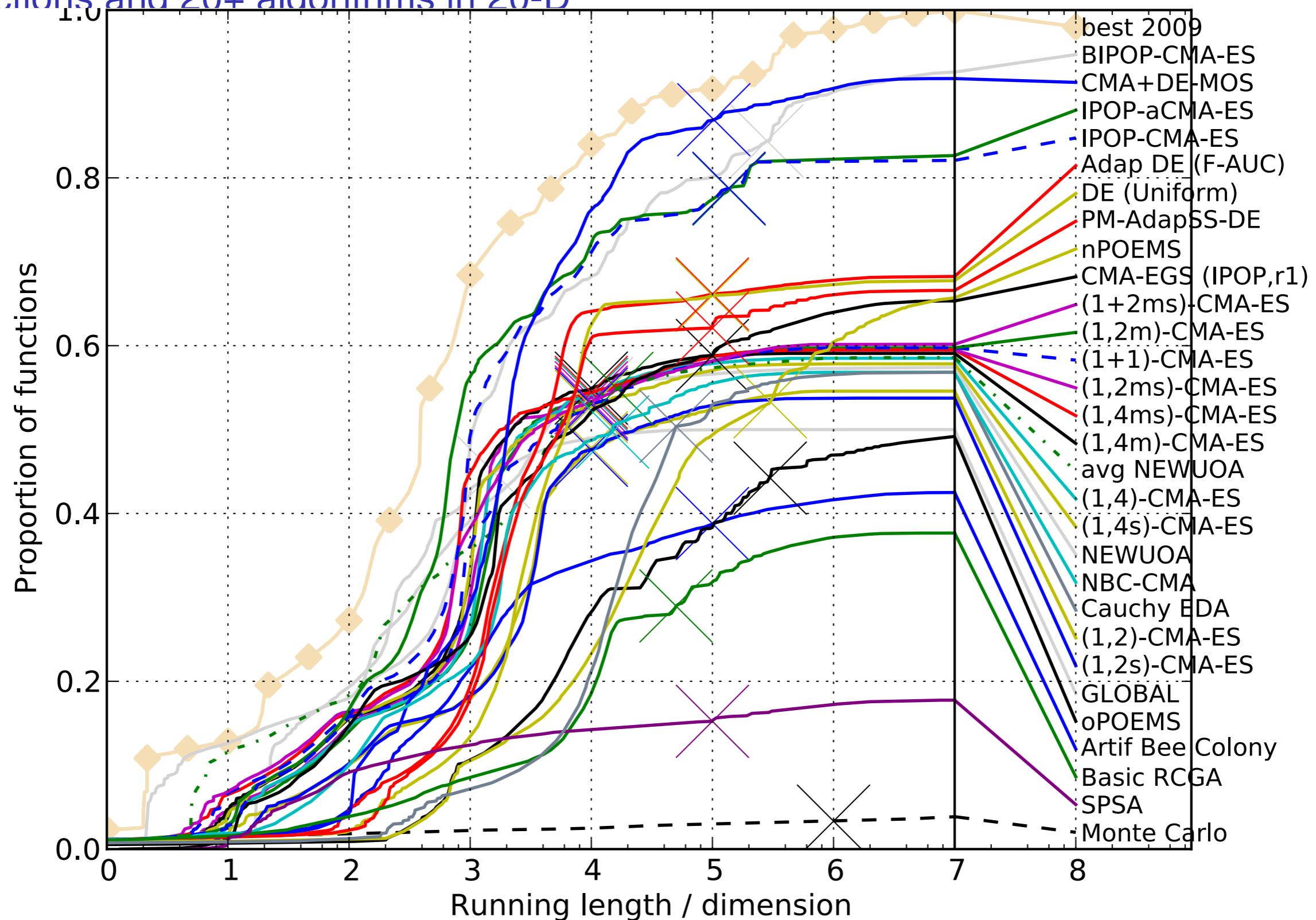
# Comparison during BBOB at GECCO 2009

24 functions and 31 algorithms in 20-D



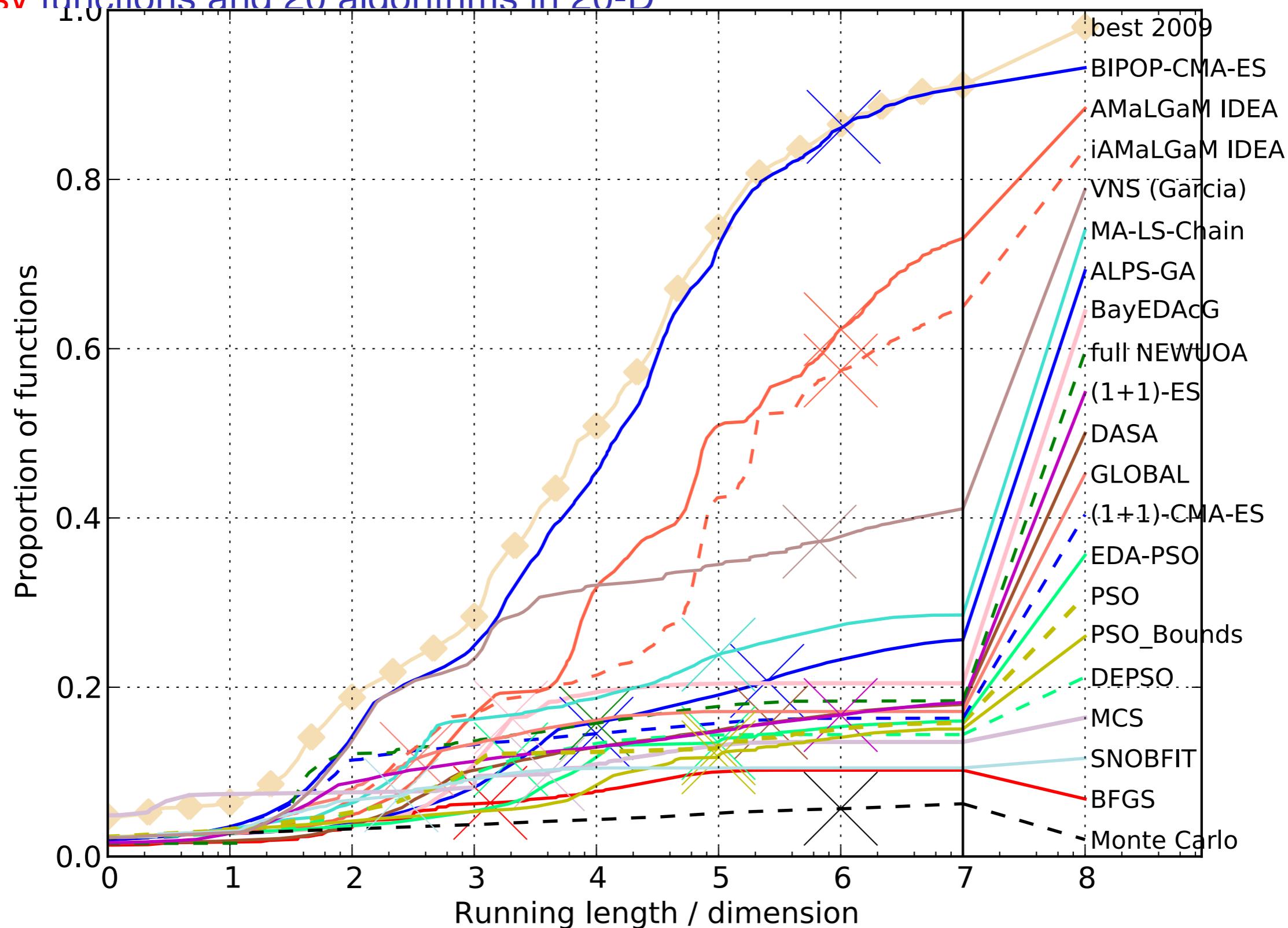
# Comparison during BBOB at GECCO 2010

24 functions and 20+ algorithms in 20-D



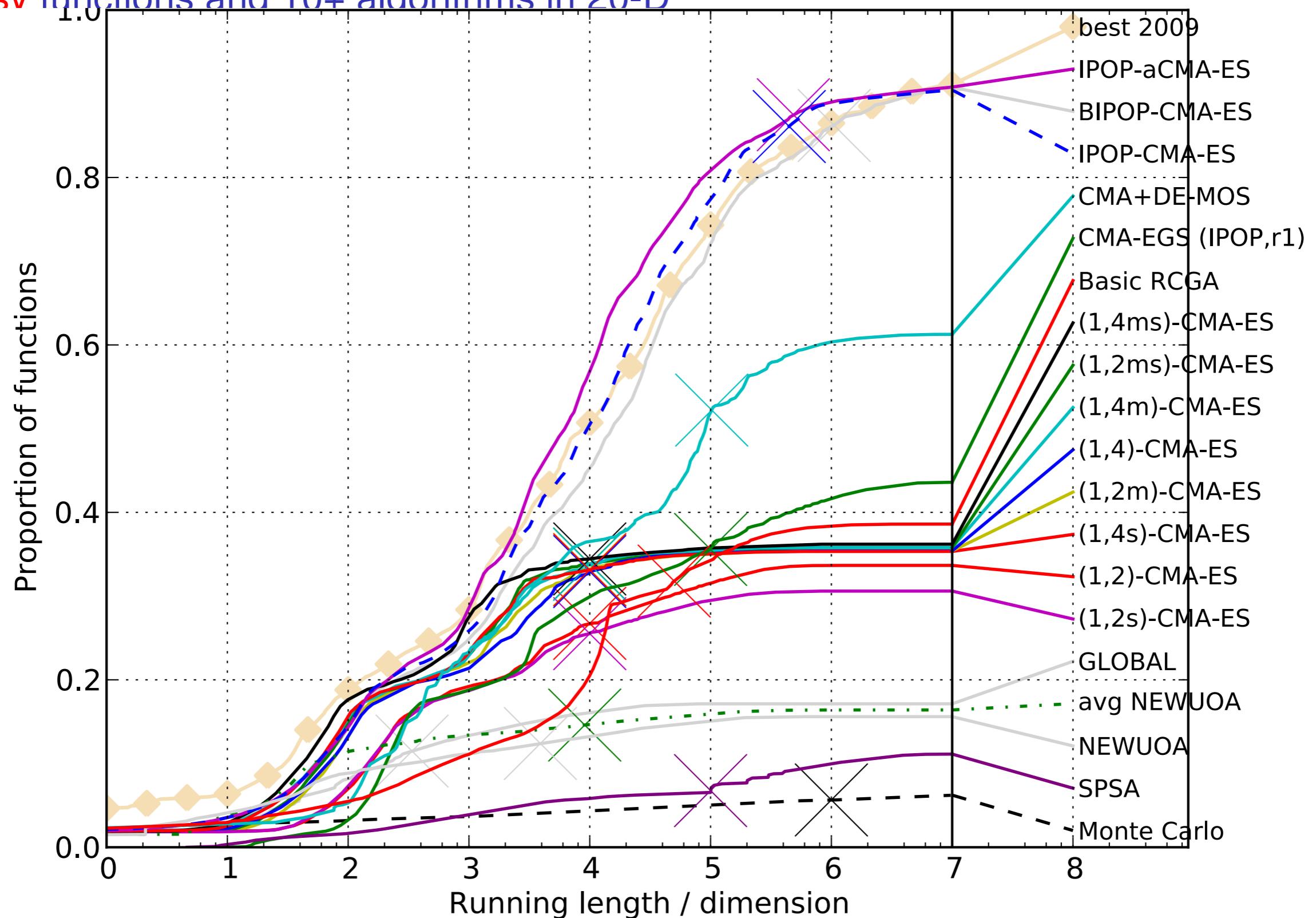
# Comparison during BBOB at GECCO 2009

30 noisv functions and 20 algorithms in 20-D



# Comparison during BBOB at GECCO 2010

30 noisv functions and 10+ algorithms in 20-D



## Final Remarks

# The Continuous Search Problem

**Difficulties** of a non-linear optimization problem are

- dimensionality and non-separability
  - demands to exploit problem structure, e.g. neighborhood  
cave: design of benchmark functions
- ill-conditioning
  - demands to acquire a second order model
- ruggedness
  - demands a non-local (stochastic? population based?) approach

# Main Characteristics of (CMA) Evolution Strategies

- ① Multivariate normal distribution to generate new search points  
follows the maximum entropy principle
- ② Rank-based selection  
implies invariance, same performance on  $g(f(\mathbf{x}))$  for any increasing  $g$   
more invariance properties are featured
- ③ Step-size control facilitates fast (log-linear) convergence and  
possibly linear scaling with the dimension  
in CMA-ES based on an **evolution path** (a non-local trajectory)
- ④ *Covariance matrix adaptation (CMA)* **increases the likelihood of previously successful steps** and can improve performance by orders of magnitude
  - the update follows the natural gradient
  - $\mathbf{C} \propto \mathbf{H}^{-1} \iff$  adapts a variable metric
  - $\iff$  new (rotated) problem representation
  - $\implies f : \mathbf{x} \mapsto g(\mathbf{x}^T \mathbf{H} \mathbf{x})$  reduces to  $\mathbf{x} \mapsto \mathbf{x}^T \mathbf{x}$

# Limitations of CMA Evolution Strategies

- internal CPU-time:  $10^{-8}n^2$  seconds per function evaluation on a 2GHz PC, tweaks are available  
1 000 000  $f$ -evaluations in 100-D take 100 seconds *internal CPU-time*
- better methods are presumably available in case of
  - ▶ partly separable problems
  - ▶ specific problems, for example with cheap gradients  
specific methods
  - ▶ small dimension ( $n \ll 10$ )  
for example Nelder-Mead
  - ▶ small running times (number of  $f$ -evaluations  $< 100n$ )  
model-based methods