

# Computational Intelligence

## Master in Artificial Intelligence

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### Introduction to Evolution Strategies

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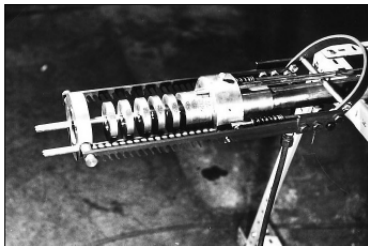
School of Professional & Executive Development

# The nozzle experiment (I)



device for clamping nozzle parts

collection of conical nozzle parts



# The nozzle experiment (II)



Hans-Paul Schwefel  
while changing nozzle parts



# The nozzle experiment (III)



the nozzle in operation ...

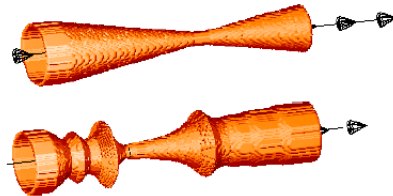
... while measuring degree of efficiency

# The nozzle experiment (IV)

– Initial:



– Evolution:



32% of increase in efficiency!

J. Klockgether and H.-P. Schwefel, "*Two-phase nozzle and hollow core jet experiments*". Proceedings of the 11th Symposium on Engineering Aspects of Magneto-Hydrodynamics, Caltech, Pasadena, California, USA, 1970.

# The Gaussian Distribution

A continuous  $d$ -variate random vector  $\mathbf{X} = (X_1, \dots, X_d)^T$  is **normally distributed**, written  $\mathbf{X} \sim \mathcal{N}(\boldsymbol{\mu}, \Sigma)$ , when its joint *pdf* is:

$$p(\mathbf{x}) = \frac{1}{(2\pi)^{\frac{d}{2}} |\Sigma|^{\frac{1}{2}}} \exp \left\{ -\frac{1}{2} (\mathbf{x} - \boldsymbol{\mu})^T \Sigma^{-1} (\mathbf{x} - \boldsymbol{\mu}) \right\}$$

where  $\boldsymbol{\mu}$  is the mean vector and  $\Sigma_{d \times d} = (\sigma_{ij}^2)$  is the (real symmetric and p.d.) covariance matrix.

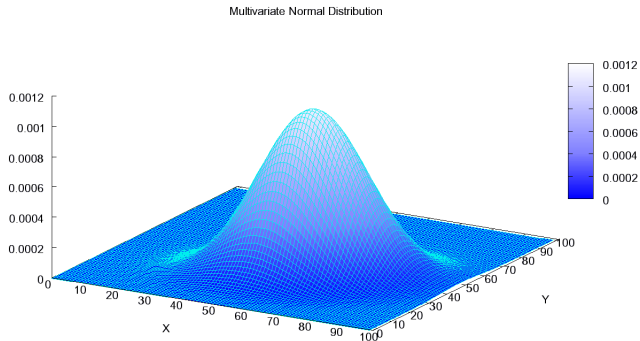
- $\mathbb{E}[\mathbf{X}] = \boldsymbol{\mu}$  and  $\mathbb{E}[(\mathbf{X} - \boldsymbol{\mu})(\mathbf{X} - \boldsymbol{\mu})^T] = \Sigma$ .
- $\text{CoVar}[X_i, X_j] = \sigma_{ij}^2$  and  $\text{Var}[X_i] = \sigma_{ii}^2$

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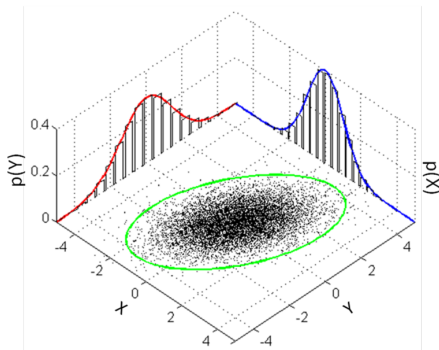
if  $\mathbf{X} \sim \mathcal{N}(\boldsymbol{\mu}, \Sigma)$ , then  $X_i, X_j$  are independent  $\iff \text{CoVar}[X_i, X_j] = 0$

(in general, only the left-to-right implication holds)

# The Gaussian Distribution ( $d = 2$ )



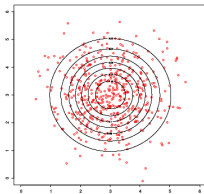
# The Gaussian Distribution ( $d = 2$ )



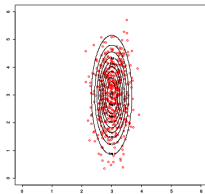
Observations from a bivariate normal distribution, a contour ellipsoid, the two marginal distributions, and their histograms (all images from the Wikipedia)



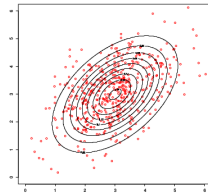
# Linear algebra point of view ( $d = 2$ )



$$\mu = \begin{bmatrix} 3 \\ 3 \end{bmatrix} \quad \Sigma = \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix}$$



$$\Sigma = \begin{bmatrix} 0.1 & 0 \\ 0 & 1 \end{bmatrix}$$



$$\Sigma = \begin{bmatrix} 1 & 0.5 \\ 0.5 & 1 \end{bmatrix}$$

- The principal directions (a.k.a. PCs) of the hyperellipsoids are given by the eigenvectors  $\mathbf{u}_i$  of  $\Sigma$ , which satisfy  $\Sigma \mathbf{u}_i = \lambda_i \mathbf{u}_i$ .
- The lengths of the hyperellipsoids along these axes are proportional to  $\sqrt{\lambda_i}$  (note  $\lambda_i > 0$ ), where  $\lambda_i$  are the eigenvalues associated with  $\mathbf{u}_i$ .

- What is behind the choice of a **multivariate Gaussian**?

Examples from a class are noisy versions of an ideal class member (a prototype):

- Prototype: modeled by the mean vector
  - Noise: modeled by the covariance matrix
- The quantity

$$d(\mathbf{x}) := \sqrt{(\mathbf{x} - \boldsymbol{\mu})^T \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu})}$$

is called the **Mahalanobis distance** for  $\mathbf{x}$

- Very important! the number of parameters is  $\frac{d(d+1)}{2} + d$

## Positive definiteness

For a Gaussian distribution to be well-defined,  $\Sigma$  has to be real symmetric and positive definite (p.d.):

- for all non-null vectors  $\mathbf{x} \in \mathbb{R}^d$ ,  $\mathbf{x}^T \Sigma \mathbf{x} > 0$  must hold true.
- alt., all eigenvalues must be positive (note they are real)

Examples: are these matrices p.d.?

$$a. \begin{pmatrix} 1 & 0 \\ 0 & 1 \end{pmatrix} \quad b. \begin{pmatrix} 1 & \frac{1}{2} \\ \frac{1}{2} & 1 \end{pmatrix}$$

$$c. \begin{pmatrix} 3 & -1 \\ -1 & 2 \end{pmatrix} \quad d. \begin{pmatrix} 1 & 4 \\ \frac{1}{2} & 1 \end{pmatrix}$$

- a. YES;      b. YES  
c. YES;      d. NO

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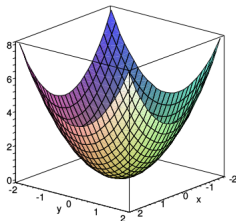
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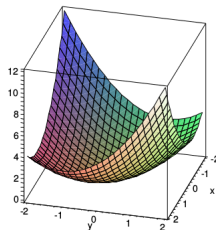
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- |         |        |
|---------|--------|
| a. YES; | b. YES |
| c. YES; | d. NO  |

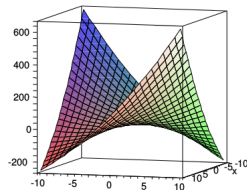
# Mathematical view



a.  $z_1^2 + z_2^2;$



b.  $z_1^2 + z_1 z_2 + z_2^2;$



d.  $z_1^2 + \frac{9}{2} z_1 z_2 + z_2^2$

# Evolution Strategies: main characteristics

- Continuous search space  $\mathbb{R}^n$  ( $n$  **objective** parameters)
- Various *ad hoc* recombination operators
- Deterministic  $(\mu, \lambda)$ -replacement
- Generation of an offspring *surplus*:  $\lambda \gg \mu$
- Emphasis on mutation:  $n$ -dimensional Gaussian
- Self-adaptation of mutation parameters  
(first self-adaptive EA!)

Recall the notation:

$$(\mu/\rho, \lambda) - \text{ES}$$

The three parts of an individual

- 1 object variables  $\mathbf{x} \in \mathbb{R}^n$  to compute fitness  $F(\mathbf{x})$
- 2 standard deviations  $\boldsymbol{\sigma} \in \mathbb{R}_+^{n_\sigma}$  to express variances
- 3 rotation angles  $\boldsymbol{\alpha} \in (-\pi, \pi]^{n_\alpha}$  to express covariances  
(all Gaussians are zero mean)

# Evolution Strategies: Mutation (I)

## Simple self-adaptive Mutation

$$n_\sigma = 1, n_\alpha = 0$$

(one mutation parameter per individual)

$$\sigma := \sigma \cdot \exp(\mathcal{N}(0, \tau_0))$$

**For**  $i \in \{1, 2, \dots, n\}$

①  $x_i := x_i + \mathcal{N}_i(0, \sigma^2)$

where

$$\tau_0 \propto \frac{1}{n}$$

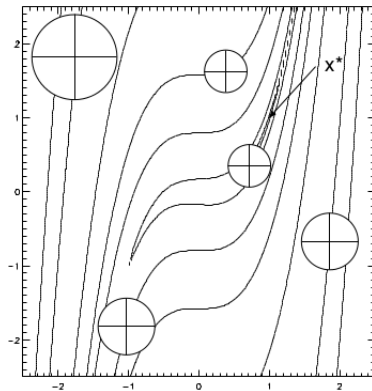


# Evolution Strategies: Mutation (I)

## Simple self-adaptive Mutation ( $n = 2$ )



equal probability to place an offspring



# Evolution Strategies: Mutation (II)

## Diagonal self-adaptive Mutation

$$n_\sigma = n, n_\alpha = 0$$

(one mutation parameter per individual and variable)

**For**  $i \in \{1, 2, \dots, n_\sigma\}$

①  $\sigma_i := \sigma_i \cdot \exp(\mathcal{N}(0, \tau') + \mathcal{N}_i(0, \tau))$

**For**  $i \in \{1, 2, \dots, n\}$

①  $x_i := x_i + \mathcal{N}_i(0, \sigma_i^2)$

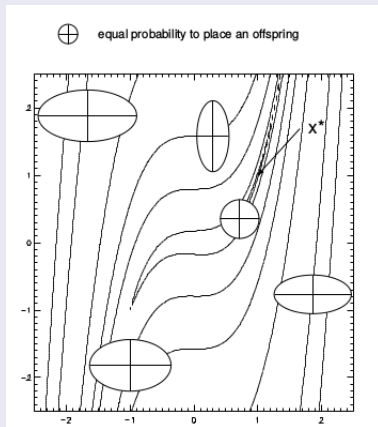
where

$$\tau \propto \frac{1}{2\sqrt{n}}$$

$$\tau' \propto \frac{1}{2n}$$

# Evolution Strategies: Mutation (II)

## Diagonal Self-Adaptive Mutation ( $n = 2$ )



# Evolution Strategies: Mutation (III)

## Correlated self-adaptive Mutation

$$n_\sigma = n, n_\alpha = \left(n - \frac{n_\sigma}{2}\right) (n_\sigma - 1)$$

(one covariance matrix per individual, represented by a collection of  $n_\alpha$  rotation angles)

**For**  $i \in \{1, 2, \dots, n_\sigma\}$

$$\textcircled{1} \quad \sigma_i := \sigma_i \cdot \exp(\mathcal{N}(0, \tau') + \mathcal{N}_i(0, \tau)), \quad \tau \propto \frac{1}{2\sqrt{n}}, \tau' \propto \frac{1}{2n}$$

**For**  $i \in \{1, 2, \dots, n_\alpha\}$

$$\textcircled{1} \quad \alpha_i := \alpha_i + \mathcal{N}_i(0, \beta^2), \quad \beta \propto 5^\circ = \pi/36 \text{ radians}$$

Build  $\Sigma$  using the  $\sigma$  and  $\alpha$  for individual  $\mathbf{x}$  and then

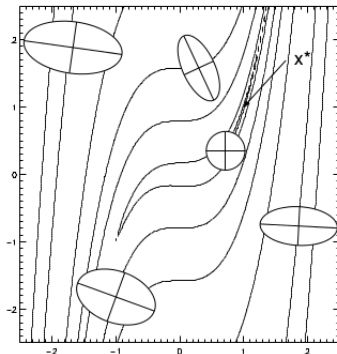
$$\mathbf{x} := \mathbf{x} + \mathcal{N}(0, \Sigma)$$

# Evolution Strategies: Mutation (III)

## Correlated Self-Adaptive Mutation ( $n = 2$ )

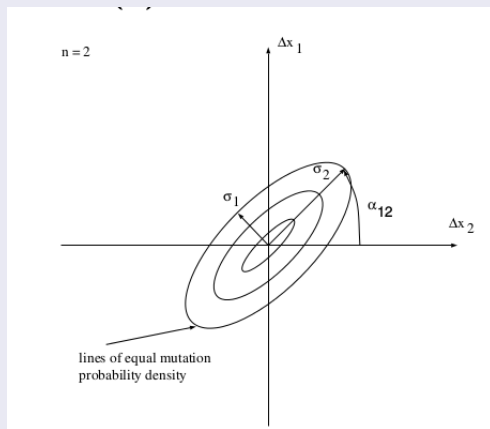


equal probability to place an offspring



# Evolution Strategies: Mutation (III)

## Illustration of the mutation ellipsoid for the case ( $n = 2$ )



# Evolution Strategies: Mutation (III)

## Theorem (Rudolph 1992)

A real symmetric matrix  $\Sigma_{n \times n}$  is p.d. iff it can be decomposed as  $\Sigma = (ST)^T(ST)$ , with  $T$  orthogonal and  $S$  diagonal with  $s_{ii} > 0$  and:

$$T := \prod_{i=1}^{n-1} \prod_{j=i+1}^n T_{ij}(\alpha_{f(i,j)})$$

- $T$  is the product of  $\frac{n(n-1)}{2}$  elementary rotation matrices  $T_{ij}$ .
- $\alpha_{f(i,j)}$  are the rotation angles (between axes  $i$  and  $j$ ), represented in the chromosomic vector in position  $f(i,j)$ .
- $T_{ij}(\alpha_{f(i,j)})$  is built as the identity matrix and modified as:

$$\begin{aligned} r_{ii} &= r_{jj} := \cos(\alpha_{f(i,j)}) \\ r_{ij} &= -r_{ji} := -\sin(\alpha_{f(i,j)}), \quad i \neq j \end{aligned}$$

# Evolution Strategies: Mutation (III)

- The *dummy* function  $f(i, j)$  is used to index the vector of self-adaptive parameters (angles)  $\alpha$ , using a single index.
- As a consequence, a total of  $\frac{n(n+1)}{2}$  angles and scaling parameters are sufficient to generate arbitrary correlated Normal random vectors with 0 mean and covariance matrix  $\Sigma = (ST)^\top(ST)$  via:

$$\mathbf{x} := \mathbf{x} + T\mathbf{z}$$

with  $\mathbf{z} \sim \mathcal{N}(0, S)$  and  $S = \text{diag}(\sigma_1^2, \dots, \sigma_{n_\sigma}^2)$ .



## Log-normal distribution

It is a continuous probability distribution whose logarithm is normally distributed. A random variable which is log-normally distributed takes only positive real values.

$$\sigma_i := \sigma_i \cdot \exp(\mathcal{N}(0, \tau'))$$

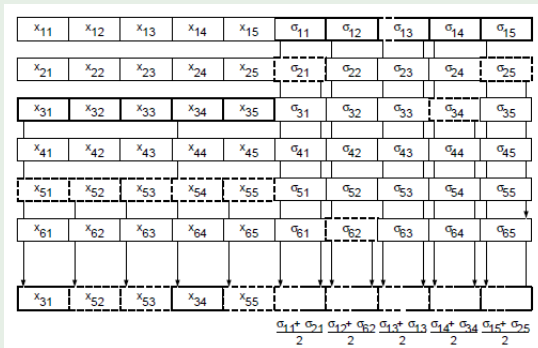
- 1 Multiplication by positive values preserves positivity
- 2  $Pr\{X = x\} = Pr\{X = \frac{1}{x}\}, x > 0$
- 3 Small modifications are more probable than larger ones

# Evolution Strategies: recombination (I)

- Usually introduced as the *first* operator (before mutation)
- Generates an intermediate population size of  $\lambda$  by generating *one individual at a time* out of  $\rho$  parents by looping  $\lambda \gg \mu$  times (generation of a **surplus**)
- Typically  $\rho = 2$  (**dual**) or  $\rho = \mu$  (**global** recombination):
  - dual**: the two parents are chosen at random, per individual
  - global**: one parent is held fixed and the other is chosen anew per each gene
- Applied to both objective and strategy parameters (and often differently)
- Two basic ways: choose randomly (**discrete**) and average (**intermediate**)

# Evolution Strategies: recombination (II)

## Recombination example



- $\mu = 6, n = 5, n_\sigma = n, n_\alpha = 0$  (one mutation parameter per individual and gene)
- dual discrete recombination on  $x_i$ ; global intermediate on  $\sigma_i$  (first parent held fixed, second chosen anew)

# Evolution Strategies: replacement

- Strictly deterministic, rank-based
- The  $\mu$  best are treated equally
- $(\mu, \lambda)$  selection:
  - offspring surplus  $\lambda \gg \mu$
  - important (necessary?) for self-adaptation
  - useful for moving optima, noisy  $F$ , ...

⇒ Very strong selective pressure

## The crucial claim (Schwefel '87 '92)

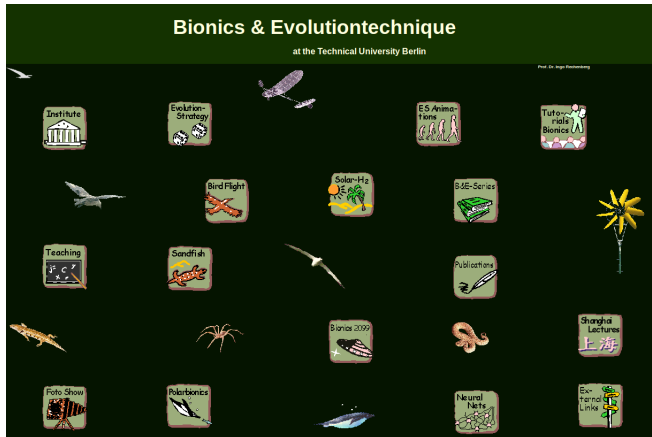
Self-adaptation of strategy parameters works!

- without exogenous or centralized control
- needs mutation of all parameters
- needs generation of a surplus and  $(\mu, \lambda)$  replacement
- needs recombination of all parameters

default (recommended) settings:

- $\lambda \propto 7\mu$ ; historically  $(\mu = 7, \lambda = 105)$
- dual discrete recombination on objective parameters
- global intermediate on strategy parameters

# Evolution Strategies: demos



Prof. Dr. Ingo Rechenberg

<https://web.archive.org/web/20180425010001/http://www.bionik.tu-berlin.de/institut/xstart.htm>

# Evolution Strategies: Modern developments

The CMA-ES (Covariance Matrix Adaptation Evolution Strategy, by N. Hansen) is the more recent development of ESs:

- Uses a more sophisticated method to update the covariance matrix, particularly useful if the fitness function is complex.
- Learns a second order model of the underlying function (similar to the approximation of the inverse Hessian matrix in quasi-Newton methods, used for example in neural networks).

## Resources:

- [www.cmap.polytechnique.fr/~nikolaus.hansen/cma-es.github.io/index.html](http://www.cmap.polytechnique.fr/~nikolaus.hansen/cma-es.github.io/index.html)  
[arxiv.org/pdf/1604.00772.pdf](https://arxiv.org/pdf/1604.00772.pdf)
- **C, C++, Java, Matlab, Octave, Python, Scilab**  
[cma-es.github.io/cmaes\\_sourcecode\\_page.html](http://cma-es.github.io/cmaes_sourcecode_page.html)
- **Julia** [github.com/bionik-berlin/PURE\\_ES](https://github.com/bionik-berlin/PURE_ES)
- **R** packages {rCMA, cmaes, adagio, parma}
- **Python** [github.com/CMA-ES/pycma](https://github.com/CMA-ES/pycma)
- **Tensorflow 2** [pypi.org/project/cma-es/](https://pypi.org/project/cma-es/)