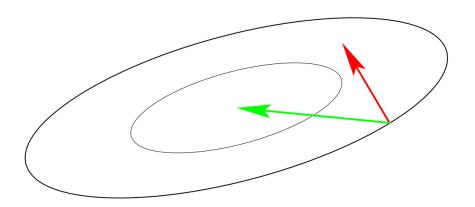
consider the curvature of the level sets of a function

ill-conditioned means "squeezed" lines of equal function value (high curvatures)



gradient direction  $-f'(\mathbf{x})^{\mathrm{T}}$ Newton direction  $-\mathbf{H}^{-1}f'(\mathbf{x})^{\mathrm{T}}$ 

Condition number equals nine here. Condition numbers up to  $10^{10}$  are not unusual in real world problems.

# Part II: Algorithms

#### **Deterministic Algorithms**

Quasi-Newton with estimation of gradient (BFGS) [Broyden et al. 1970] Simplex downhill [Nelder and Mead 1965] Pattern search, Direct Search [Hooke and Jeeves 1961] Trust-region/Model Based methods (NEWUOA, BOBYQA) [Powell, 06,09]

#### Stochastic (randomized) search methods

Evolutionary Algorithms (continuous domain)
Differential Evolution [Storn, Price 1997]
Particle Swarm Optimization [Kennedy and Eberhart 1995]
Evolution Strategies, CMA-ES [Rechenberg 1965, Hansen, Ostermeier 2001]
Estimation of Distribution Algorithms (EDAs) [Larrañaga, Lozano, 2002]
Cross Entropy Method (same as EDAs) [Rubinstein, Kroese, 2004]
Genetic Algorithms [Holland 1975, Goldberg 1989]

Simulated Annealing [Kirkpatrick et al. 1983]

Define  $\{P_{\theta} : \theta \in \Theta\}$ , a family of probability distributions on  $\mathbb{R}^{n}$ 

Generic template to optimize  $f : \mathbb{R}^n \to \mathbb{R}$ 

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

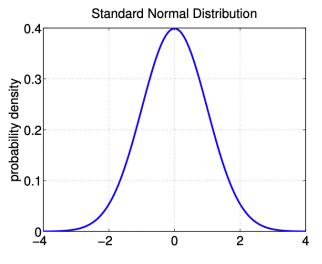
- 1. Sample  $x_1, \ldots, x_{\lambda}$  according to  $P_{\theta}$
- 2. Evaluate  $x_1, \ldots, x_{\lambda}$  on f
- 3. Update parameters  $\theta \leftarrow F(\theta, x_1, ..., x_{\lambda}, f(x_1), ..., f(x_{\lambda}))$

the update of  $\theta$  should drive  $P_{\theta}$  to concentrate on the optima of f

To obtain an optimization algorithm we need: **1** to define  $\{P_{\theta}, \theta \in \Theta\}$ **2** to define F the update function of  $\theta$ 

# Which probability distribution to sample candidate solutions?

### Normal distribution - 1D case



probability density of the 1-D standard normal distribution  $\mathcal{N}(0,1)$ 

(expected (mean) value, variance) = (0,1)

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

General case

▶ Normal distribution  $\mathcal{N}(\boldsymbol{m}, \sigma^2)$ 

(expected value, variance) = 
$$(\boldsymbol{m}, \sigma^2)$$
  
density:  $p_{\boldsymbol{m},\sigma}(x) = \frac{1}{\sqrt{2\pi\sigma}} \exp\left(-\frac{(x-\boldsymbol{m})^2}{2\sigma^2}\right)$ 

- A normal distribution is entirely determined by its mean value and variance
- The family of normal distributions is closed under linear transformations: if X is normally distributed then a linear transformation aX + b is also normally distributed

• Exercice: Show that 
$$m + \sigma \mathcal{N}(0, 1) = \mathcal{N}(m, \sigma^2)$$

#### Generalization to n Variables: Independent Case

Assume X1 ~ 
$$\mathcal{N}(\mu_1, \sigma_1^2)$$
 denote its density  $p(x_1) = \frac{1}{Z_1} \exp\left(-\frac{1}{2\sigma_1^2}(x_1 - \mu_1)^2\right)$   
Assume X2~  $\mathcal{N}(\mu_2, \sigma_2^2)$  denote its density  $p(x_2) = \frac{1}{Z_2} \exp\left(-\frac{1}{2\sigma_2^2}(x_2 - \mu_2)^2\right)$ 

Assume X1 and X2 are **independent**, then (X1,X2) is a Gaussian vector with

$$p(x_1, x_2) =$$

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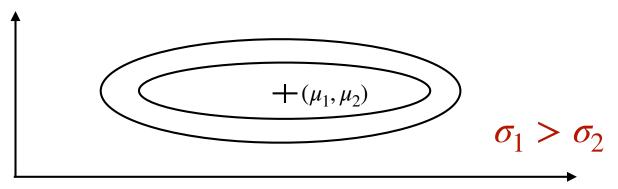
$$p(x_1, x_2) = p(x_1)p(x_2) = \frac{1}{Z_1 Z_2} \exp\left(-\frac{1}{2}(x-\mu)^T \Sigma^{-1}(x-\mu)\right)$$
  
with  $x = (x_1, x_2)^T$   $\mu = (\mu_1, \mu_2)^T$   $\Sigma = \begin{pmatrix} \sigma_1^2 & 0\\ 0 & \sigma_2^2 \end{pmatrix}$ 

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Assume X1 and X2 are **independent**, then (X1,X2) is a Gaussian vector with

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#### **Gaussian Vector - Multivariate Normal Distribution**

A random vector  $X = (X_1, ..., X_n) \in \mathbb{R}^n$  is a Gaussian vector (or multivariate normal) if and only if for all real numbers  $a_1, ..., a_n$ , the random variable  $a_1X_1 + ... + a_nX_n$  has a normal distribution.

#### Gaussian Vector - Multivariate Normal Distribution

A random variable following a 1-D normal distribution is determined by its mean value m and variance  $\sigma^2$ .

In the *n*-dimensional case it is determined by its mean vector and covariance matrix

#### **Covariance Matrix**

If the entries in a vector  $\mathbf{X} = (X_1, \dots, X_n)^T$  are random variables, each with finite variance, then the covariance matrix  $\Sigma$  is the matrix whose (i, j) entries are the covariance of  $(X_i, X_j)$ 

$$\Sigma_{ij} = \operatorname{cov}(X_i, X_j) = \operatorname{E}\left[(X_i - \mu_i)(X_j - \mu_j)\right]$$

where  $\mu_i = E(X_i)$ . Considering the expectation of a matrix as the expectation of each entry, we have

$$\boldsymbol{\Sigma} = \mathrm{E}[(\boldsymbol{X} - \boldsymbol{\mu})(\boldsymbol{X} - \boldsymbol{\mu})^{T}]$$

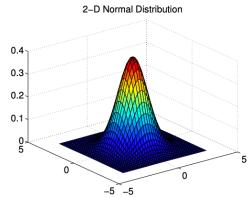
 $\Sigma$  is symmetric, positive definite

Density of a n-dimensional Gaussian vector  $\mathcal{N}(m, C)$ :

$$p_{\mathcal{N}(m,C)}(x) = \frac{1}{(2\pi)^{n/2} |C|^{1/2}} \exp\left(-\frac{1}{2}(x-m)^{\mathsf{T}} C^{-1}(x-m)\right)$$

#### The mean vector *m*:

determines the displacement is the value with the largest density



the distribution is symmetric around the mean

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

The covariance matrix:

determines the geometrical shape (see next slides)

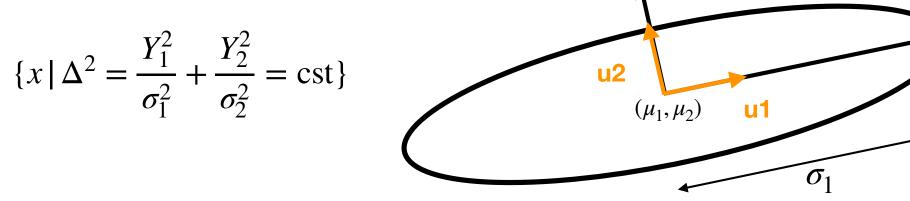
Consider a Gaussian vector  $\mathcal{N}(m, C)$ , remind that lines of equal densities are given by:

$$\{x \mid \Delta^2 = (x - m)^T C^{-1} (x - m) = \text{cst}\}\$$

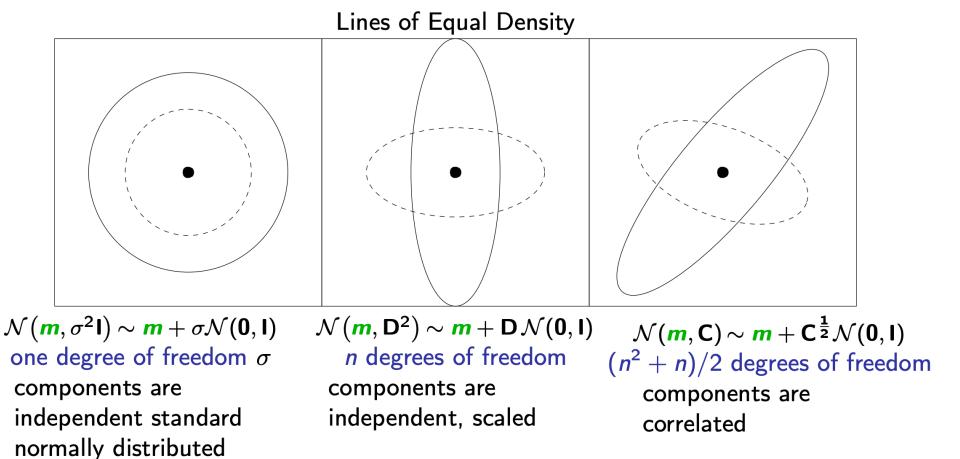
Decompose  $C = U \Lambda U^{\top}$  with U orthogonal, i.e.

$$C = \begin{pmatrix} u_1 & u_2 \\ | & | \end{pmatrix} \begin{pmatrix} \sigma_1^2 & 0 \\ 0 | & \sigma_2^2 \end{pmatrix} \begin{pmatrix} u_1 & - \\ u_2 & - \end{pmatrix}$$

Let  $Y = U^{\top}(x - m)$ , then in the coordinate system, (u1,u2), the lines of equal densities are given by



... any covariance matrix can be uniquely identified with the iso-density ellipsoid  $\{x \in \mathbb{R}^n | (x - m)^T C^{-1} (x - m) = 1\}$ 



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

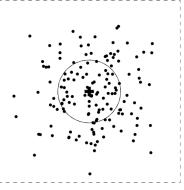
#### **Evolution Strategies**

# New search points are sampled normally distributed

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

as perturbations of *m*,

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



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

#### **Evolution Strategies**

# New search points are sampled normally distributed

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

- In fact, the covariance matrix of the sampling distribution is  $\sigma^2 \mathbb{C}$ but it is convenient to refer to  $\mathbb{C}$  as the covariance matrix (it is a covariance matrix but not of the sampling distribution)
  - the mean vector  $\boldsymbol{m} \in \mathbb{R}^n$  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

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#### How to update the different parameters $m, \sigma, \mathbf{C}$ ?

#### **1. Adapting the mean** *m*

- 2. Adapting the step-size  $\sigma$
- **3.** Adapting the covariance matrix C

Update the Mean: a Simple Algorithm the (1+1)-ES

#### Notation and Terminology:

one solution kept from one iteration to the next

one new solution (offspring) sampled at each iteration

The + means that we keep the best between current solution and new solution, we talk about *elitist* selection

(1+1)-ES

(1+1)-ES algorithm (update of the mean)

sample one candidate solution from the mean  ${\boldsymbol{m}}$ 

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

if **x** is better than **m** (i.e. if  $f(\mathbf{x}) \leq f(\mathbf{m})$ ), select **m** 

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

The (1+1)-ES algorithm is a simple algorithm, yet:
the elitist selection is not robust to outliers
we cannot loose solutions accepted by "chance", for instance that look good because the noise gave it a low function value
there is no population (just a single solution is sampled) which makes it less robust

In practice, one should rather use a:

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

The  $\mu$  best solutions are selected and recombined (to form the new mean)

 $\lambda$  solutions are sampled at each iteration

#### The $(\mu/\mu, \lambda)$ -ES - Update of the Mean Vector

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

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

Notation: we denote  $y_{i:\lambda}$  the vector such that  $x_{i:\lambda} = m + \sigma y_{i:\lambda}$ Exercice: realize that  $y_{i:\lambda}$  is generally not distributed as  $\mathcal{N}(\mathbf{0}, \mathbf{C})$ The new mean reads

$$m{m} \leftarrow \sum_{i=1}^{\mu} m{w}_i \, m{x}_{i:\lambda}$$

where

$$w_1 \geq \cdots \geq w_\mu > 0, \quad \sum_{i=1}^\mu w_i = 1, \quad rac{1}{\sum_{i=1}^\mu w_i^2} =: \mu_w pprox rac{\lambda}{4}$$

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

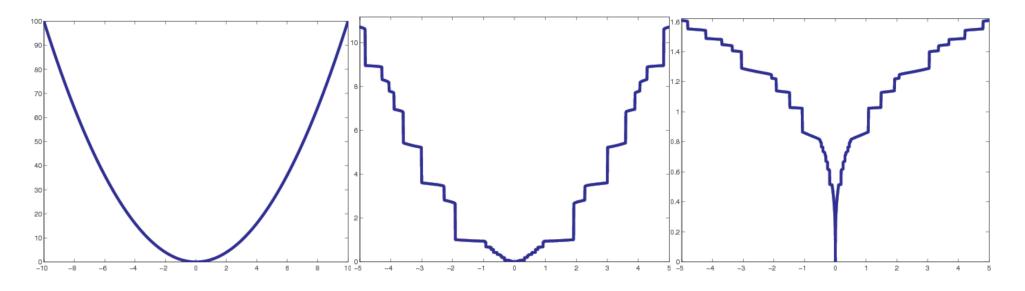
What changes in the previous slide if instead of optimizing f, we optimize  $g \circ f$  where  $g : \text{Im}(f) \to \mathbb{R}$ is strictly increasing?

#### Invariance Under Monotonically Increasing Functions

Comparison-based/ranking-based algorithms:

Update of all parameters uses only the ranking:

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



 $g(f(x_{1:\lambda})) \le g(f(x_{2:\lambda})) \le \dots \le g(f(x_{\lambda:\lambda}))$ for all  $g : \operatorname{Im}(f) \to \mathbb{R}$  strictly increasing

### A Template for Comparison-based Stochastic Search

Define  $\{P_{\theta} : \theta \in \Theta\}$ , a family of probability distributions on  $\mathbb{R}^{n}$ 

Generic template to optimize  $f : \mathbb{R}^n \to \mathbb{R}$ 

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

- 1. Sample  $x_1, \ldots, x_{\lambda}$  according to  $P_{\theta}$
- 2. Evaluate  $x_1, \ldots, x_{\lambda}$  on f
- 3. Rank the solutions and find  $\pi$  the permutation such  $f(x_{\pi(1)}) \leq f(x_{\pi(2)}) \leq \ldots \leq f(x_{\pi(\lambda)})$
- 4. Update parameters  $\theta \leftarrow F(\theta, x_1, ..., x_{\lambda}, \pi)$