

# Derivative Free Optimization

## Topic 3: Genetic Algorithms

Warren Hare

# Genetic Algorithms

# Survival of the fittest

- In 1864 Charles Darwin published “On the origin of species”
- In 1869, Darwin published edition 5 and include the phrase “survival of the fittest”
- The phrase is originally attributed to Herbert Spencer, an economist

## GENETIC ALGORITHM philosophy

Use *survival of the fittest* to remove poor solutions and create new better solutions

# Genetic Algorithms, history

- Late 1960s, Rechenberg writes “Evolutionary Strategies for Optimization” (in German) and uses them to optimize aerofoils
- 1971, Rechenberg’s PhD thesis
- 1975, John Holland writes “Adaptation in Natural and Artificial Systems”
- 2020, Rechenberg’s thesis has 125 citations,  
Holland’s book has 67,000+ citations

# Genetic Algorithms, Overview

- Points in constraint set = potential members in a population
- Collection of points = population
- Function values = quality of individual (fitness)
- Algorithms Iterations = reproduction & survival

# Algorithm: Genetic Algorithms (GA)

Given  $f : \mathbb{R}^n \mapsto \mathbb{R}$  and an initial population  $P^0 = \{x^1, x^2, \dots, x^{\bar{p}}\}$

0. Initialize:

$\gamma \in (0, 1)$	mutation probability
$k \leftarrow 0$	iteration counter

1. Fitness:

Use  $f(x)$  to assign a fitness to each individual  $x \in P^k$

2. Reproduce:

- 2a) Selection: select 2 parents from  $P^k$  and proceed to 2b)  
or select 1 survivor from  $P^k$  and proceed to 2d)
- 2b) Crossover: use the 2 parents to create an offspring
- 2c) Mutation: with probability  $\gamma$  mutate the offspring  
if the offspring is infeasible  
declare the offspring deceased and return to 2a)  
otherwise declare the offspring a survivor and proceed to 2d)
- 2d) Update next generation: place survivor into  $P^{k+1}$   
if  $|P^{k+1}| \geq \bar{p}$  declare population complete and proceed to 3  
otherwise declare population incomplete and go to 2a)

3. Update:

increment  $k \leftarrow k + 1$ , stop or go to 1

# Notes

- GA is more of a framework than an algorithm
- The great flexibility has resulted in a lot of custom-made GA

# Fitness & Selection



# Genetic Algorithm

To maintain the spirit of genetics,

- **GA** begins by assigning each point a **fitness**
- Fitness is used to select **survivors** and **parents**
- Parents create **offspring**
- Offspring are **mutated** to create more variation

# Fitness

Fitness converts function value into a new number

Fitness should follow two rules

- When *minimizing*, if  $f(x) < f(y)$  then  $x$  is more fit than  $y$   
(When *maximizing*, if  $f(x) > f(y)$  then  $x$  is more fit than  $y$ )
- Fitness should be strictly positive

# Rank Fitness (minimization)

**Definition:** Given a population  $P^k$  with function values  $\{f(x^i)\}$   
 The **rank fitness** of  $x^i$  is the number of points  $x^j$  where  $f(x^i) \leq f(x^j)$

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**Example:** Suppose  $P = \{x^1, x^2, \dots, x^5\}$  and

$$f(x^1) = -3, \quad f(x^2) = 4, \quad f(x^3) = 1, \quad f(x^4) = 1, \quad f(x^5) = 2$$

Find the rank fitness

# Function Value Fitness (minimization)

**Definition:** Given a population  $P^k$  with function values  $\{f(x^i)\}$   
 The **function valued fitness** is obtained by multiplying the function value by  $-1$  and then shifting the results until the lowest value is equal to 1

$$\text{Fit}(x^i) = -f(x^i) + \bar{f} + 1 \quad \text{where } \bar{f} = \max_{i=1,2,\dots,\bar{p}} \{f(x^i)\}$$

**Example:** Suppose  $P = \{x^1, x^2, \dots, x^5\}$  and

$$f(x^1) = -3, \quad f(x^2) = 4, \quad f(x^3) = 1, \quad f(x^4) = 1, \quad f(x^5) = 2$$

Find the function value fitness

# Selection

Given population  $P^k$

In selection we create two subsets

- 1 Survivors  $S^k \subseteq P^k$
- 2 Parents  $T^k \subseteq \{P^k, P^k\}$

To maintain the spirit of genetics, typically some randomness is used

# Elitism Selection

**Definition:** Given a population  $P^k$  with fitness values  $\{f^i = \text{Fit}(x^i)\}$  **Elitism selection** of size  $s$  selects the  $s$  individuals with the highest fitness level with ties broken by random selection with uniform probability

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**Example:** Suppose  $P = \{x^1, x^2, \dots, x^5\}$  and

$$\text{Fit}(x^1) = 3, \quad \text{Fit}(x^2) = 2, \quad \text{Fit}(x^3) = 1, \quad \text{Fit}(x^4) = 5, \quad \text{Fit}(x^5) = 3$$

Describe the result if elitism selection is used to select 2 individuals

# Roulette Wheel Selection

**Definition:** Given a population  $P^k$  with fitness values  $\{f^i = \text{Fit}(x^i)\}$  **Roulette wheel selection** randomly selects individuals with probability

$$\text{prob}\{x^i\} = f^i / \sum_{j=1}^{\bar{p}} f^j$$

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**Example:** Suppose  $P = \{x^1, x^2, \dots, x^5\}$  and

$$\text{Fit}(x^1) = 3, \quad \text{Fit}(x^2) = 2, \quad \text{Fit}(x^3) = 1, \quad \text{Fit}(x^4) = 5, \quad \text{Fit}(x^5) = 3$$

Describe the result if roulette wheel selection is used to select 1 individual

# Tournament Selection

**Definition:** Given a population  $P^k$  with fitness values  $\{f^i = \text{Fit}(x^i)\}$

Given a **tournament size**  $1 \leq N \leq \bar{p}$

**Tournament selection** first randomly chooses  $N$  *distinct* individuals with equal probability, then selects from this subset using either elitism or roulette selection

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**Example:** Suppose  $P = \{x^1, x^2, \dots, x^5\}$  and

$$\text{Fit}(x^1) = 3, \quad \text{Fit}(x^2) = 2, \quad \text{Fit}(x^3) = 1, \quad \text{Fit}(x^4) = 5, \quad \text{Fit}(x^5) = 3$$

Describe the result if tournament selection is used to select 2 individuals when

- ① the tournament size is 2 and elitism is used
- ② the tournament size is 2 and roulette wheel is used
- ③ the tournament size is 3 and roulette wheel is used



# Tournament Selection, size 2

Tournaments of size 2 for  $P = \{x^1, x^2, \dots, x^5\}$

Tournament	Probability
$\{x^1, x^2\}$	1/10
$\{x^1, x^3\}$	1/10
$\{x^1, x^4\}$	1/10
$\{x^1, x^5\}$	1/10
$\{x^2, x^3\}$	1/10
$\{x^2, x^4\}$	1/10
$\{x^2, x^5\}$	1/10
$\{x^3, x^4\}$	1/10
$\{x^3, x^5\}$	1/10
$\{x^4, x^5\}$	1/10

# Tournament Selection, Elitism size 2

Elitism winners when

$$\text{Fit}(x^1) = 3, \text{Fit}(x^2) = 2, \text{Fit}(x^3) = 1, \text{Fit}(x^4) = 5, \text{Fit}(x^5) = 3$$

Tournament	Probability	Winner
$\{x^1, x^2\}$	1/10	$x^1$
$\{x^1, x^3\}$	1/10	$x^1$
$\{x^1, x^4\}$	1/10	$x^4$
$\{x^1, x^5\}$	1/10	$x^1$ and $x^5$ tie
$\{x^2, x^3\}$	1/10	$x^2$
$\{x^2, x^4\}$	1/10	$x^4$
$\{x^2, x^5\}$	1/10	$x^5$
$\{x^3, x^4\}$	1/10	$x^4$
$\{x^3, x^5\}$	1/10	$x^5$
$\{x^4, x^5\}$	1/10	$x^4$

# Tournament Selection

Tournaments of size 3 for  $P = \{x^1, x^2, \dots, x^5\}$

Tournament	Probability
$\{x^1, x^2, x^3\}$	1/10
$\{x^1, x^2, x^4\}$	1/10
$\{x^1, x^2, x^5\}$	1/10
$\{x^1, x^3, x^4\}$	1/10
$\{x^1, x^3, x^5\}$	1/10
$\{x^1, x^4, x^5\}$	1/10
$\{x^2, x^3, x^4\}$	1/10
$\{x^2, x^3, x^5\}$	1/10
$\{x^2, x^4, x^5\}$	1/10
$\{x^3, x^4, x^5\}$	1/10

# Encoding

# What is encoding

To maintain the spirit of genetics, most **GA** start by encoding the points as *chromosomes*

$$x \in \mathbb{R}^n \mapsto c \in \{0, 1\}^d$$

This is called **encoding**

## Example:

Suppose  $\Omega = [0, 1]$

Encode  $x \in \Omega$  by

$$\begin{aligned} x < \frac{1}{2} &\mapsto c = 0 \\ x \geq \frac{1}{2} &\mapsto c = 1 \end{aligned}$$

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## Example:

Suppose  $\Omega = [0, 1]$

Encode  $x \in \Omega$  by

$$\begin{aligned} x < \frac{1}{2} &\mapsto c = 0 \\ x \geq \frac{1}{2} &\mapsto c = 1 \end{aligned}$$

Alternately

$$\begin{aligned} 0 \leq x < \frac{1}{4} &\mapsto c = [0, 0] \\ \frac{1}{4} \leq x < \frac{1}{2} &\mapsto c = [0, 1] \\ \frac{1}{2} \leq x < \frac{3}{4} &\mapsto c = [1, 0] \\ \frac{3}{4} \leq x \leq 1 &\mapsto c = [1, 1] \end{aligned}$$

# Natural encodings

Some problems have a *natural structure* that makes encoding easy

**Example:**

$$\min \left\{ \text{expected overtime} : \begin{array}{l} \text{all shifts filled} \\ \text{staff are 'happy'} \end{array} \right\}$$

uses the variables

$$x_{i,j} \in \{0, 1\}$$

with  $x_{i,j} = 1$  if staff member  $i$  works shift  $j$

(See MATH 441 for more info on scheduling problems)

# Natural encoding for bounded variables

**Definition:** If  $x \in \mathcal{Z}^n$  with  $\ell \leq x \leq u$ , then the **natural encoding for bounded integer variables** replaces each  $x_i$  with the binary string  $c^i \in \{0, 1\}^{d_i}$  where  $d_i$  is the number of bits required

The final chromosome is defined by  $c = [c^1, c^2, \dots, c^n]^\top$

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**Example:** Let  $S = \{x \in \mathbb{R}^2 : 0 \leq x_1 \leq 8, -2 \leq x_2 \leq 1, x_i \in \mathcal{Z}\}$

Develop the natural encoding for  $S$



# Discretization encoding

**Definition:** Suppose  $x \in \mathbb{R}^1$  with  $\ell \leq x \leq u$   
 Select  $d$  and set  $\Delta = (u - \ell)/(2^d)$   
 Separate  $[l, u]$  into  $2^d$  equal sized subintervals

$$[\ell, \ell + \Delta) \quad [\ell + \Delta, \ell + 2\Delta) \quad \dots \quad [\ell + (2^d - 1)\Delta, \ell + (2^d)\Delta]$$

The **discretization encoding** is created by enumerating the interval list and encoding the integer enumeration

If  $x \in \mathbb{R}^n$ , the encode each  $x_i \mapsto c^i$ , then set  $c = [c^1, c^2, \dots, c^n]^\top$

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**Example:** Let  $S = \{x \in \mathbb{R}^2 : 0 \leq x_1 \leq 8, -2 \leq x_2 \leq 1\}$   
 Develop a discretization encoding using  $2^2 = 4$  equally spaced subintervals in each coordinate

# Notes

- Using  $2^d$  subintervals in a discretization encoding avoids waste
- Another option is to not encode (leave  $x \in \mathbb{R}^n$ )

# Reproduction

# Reproduction

*Survivors* of iteration  $k$  are placed into  $P^{k+1}$

*Parents* are combined to create *offspring*

$$\text{parent1} \oplus \text{parent2} \rightarrow \text{child}$$

To maintain the spirit of genetics...

# Crossover

**Definition:** Given parents  $p = [p_1, p_2, \dots, p_d]^\top$  and  $q = [q_1, q_2, \dots, q_d]^\top$   
**Single-point crossover** selects a random number  $X \in \{1, 2, \dots, d-1\}$ ,  
 and creates the child

$$[p_1, p_2, \dots, p_X, q_{X+1}, q_{X+2}, \dots, q_d]^\top$$

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**Example:** Let  $p = [5, 2, -3, 5, 9, 1]^\top$  and  $q = [3, 0, 0, 5, 2, 1]^\top$   
 Describe the result of a single-point crossover

# Crossover

**Definition:** Given parents  $p = [p_1, p_2, \dots, p_d]^\top$  and  $q = [q_1, q_2, \dots, q_d]^\top$  **2-point crossover**, selects 2 random numbers  $X_1 \in \{1, 2, \dots, d-2\}$  and  $X_2 \in \{X_1 + 1, X_1 + 2, \dots, d-1\}$  and creates the child

$$[p_1, p_2, \dots, p_{X_1}, q_{X_1+1}, q_{X_1+2}, \dots, q_{X_2}, p_{X_2+1}, p_{X_2+2}, \dots, p_d]^\top$$

---

**Example:** Let  $p = [5, 2, -3, 5, 9, 1]^\top$  and  $q = [3, 0, 0, 5, 2, 1]^\top$   
Describe the result of a 2-point crossover

# Crossover

**Definition:** Given parents  $p = [p_1, p_2, \dots, p_d]^\top$  and  $q = [q_1, q_2, \dots, q_d]^\top$   **$n$ -point crossover**, selects  $n$  random numbers ...

# Probabilistic gene selection

**Definition:** Given parents  $p = [p_1, p_2, \dots, p_d]^\top$  and  $q = [q_1, q_2, \dots, q_d]^\top$   
Select a probability parameter  $\theta \in (0, 1)$

**Probabilistic gene selection** produces child  $c$  such that

$$\text{prob}(c_i = p_i) = \theta \quad \text{and} \quad \text{prob}(c_i = q_i) = (1 - \theta)$$

---

**Example:** Let  $p = [5, 2, -3, 5, 9, 1]^\top$  and  $q = [3, 0, 0, 5, 2, 1]^\top$   
Describe the result of a probabilistic gene selection for  $\theta = 0.25$



# Weighted Average

**Definition:** Given parents  $p \in \mathbb{R}^d$  and  $q \in \mathbb{R}^d$

Select a weight parameter  $\theta \in (0, 1)$

**Weighted average** produces the child

$$c = \theta p + (1 - \theta)q$$

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**Example:** Let  $p = [5, 2, -3, 5, 9, 1]^\top$  and  $q = [3, 0, 0, 5, 2, 1]^\top$

Suppose  $\text{Fit}(p) = 6$  and  $\text{Fit}(q) = 4$

Describe the result of a weighted average where  $\theta = \frac{\text{Fit}(p)}{\text{Fit}(p) + \text{Fit}(q)}$

# Mutation

# Mutation

Offspring in natural are never exact copies of their parents

To maintain the spirit of genetics, *mutation* is applied

# Bit Inversion

**Definition:** Given offspring  $c = [c_1, c_2, \dots, c_d]^T$  where  $c_i \in \{0, 1\}$

Fix a *mutation probability*  $\delta \in (0, 1)$

The **bit inversion** mutation creates *mutated offspring*  $m$  such that

$$\text{prob}(m_i = c_i) = (1 - \delta) \quad \text{and} \quad \text{prob}(m_i \neq c_i) = \delta.$$

---

**Example:** Suppose  $c = [1, 0, 0]^T$  and let  $\delta = 1/3$

Describe the mutated offspring of a bit inversion mutation

# Coordinate perturbation

Given offspring  $c = [c_1, c_2, \dots, c_d]^T$  where  $c_i \in \mathbb{R}$

Fix a *mutation probability*  $\delta \in (0, 1)$

Fix a *mutation probability distribution* function  $F$

**Coordinate perturbation** creates *mutated offspring*  $m$  such that

$$\text{prob}(m_i = c_i) = (1 - \delta) \quad \text{and} \quad \text{prob}(m_i \neq c_i) = \delta$$

and

if  $m_i \neq c_i$ , then  $m_i = c_i + \xi$  where  $\xi$  follows the distribution  $F$

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**Example:** Suppose  $c = [0.5, 0, 0.3]^T$  and let  $\delta = 1/3$

Let  $F$  be uniform in  $[-0.1, 0.1]$

Describe the mutated offspring of a coordinate perturbation

# Convergence

# Stochastic

GA are stochastic

- i.e., the algorithm contains random subroutines, so running it twice on the same problem can get different results

So, we aim for *probabilistic convergence*

- i.e., we try to prove  $\text{prob}(\lim_{k \rightarrow \infty} f_{\text{best}}^k = f^*) = 1$

# Required assumptions

## Definition: (Monotonicity)

The **monotonicity** assumption for **GA** is that the best function value for population  $k + 1$  is at least as good as the best function value for population  $k$

$$\min\{f(x^i) : x^i \in P^{k+1}\} \leq \min\{f(x^i) : x^i \in P^k\}$$

## Definition: (Positive probabilities - binary encoded)

Suppose points are binary encoded

The **positive probabilities** assumption states that the set of encoded points is finite and there exists  $\varepsilon > 0$  such that

$$\text{prob}(c \in P^k) \geq \varepsilon \quad \text{for any encoded point } c \text{ and iteration } k$$

\* Positive probabilities can be rephrased for unencoded or other encodings



# Monotonicity

**Theorem:** Suppose a GA is used such that

- each generation has at least 1 survivor picked by elitism

Then the *monotonicity assumption holds*

# Positive probabilities

**Theorem:** Suppose a GA with binary encoding is used such that

- at least one offspring is generated per iteration
- bit inversion mutation is used

Then the *positive probabilities assumption holds*

\* Similar results hold for unencoded or other encodings

# Convergence

**Theorem:** Suppose a GA is used such that

- the points are binary encoded
- the monotonicity and positive probability assumptions hold

Let

$$f^* = \min_x \{f(x) : x \in \Omega\}, \quad X^* = \operatorname{argmin}_x \{f(x) : x \in \Omega\},$$

$$f_{\text{best}}^k = \min \{f(x^i) : x^i \in P^k\} \text{ and } x_{\text{best}}^k \in \operatorname{argmin} \{f(x^i) : x^i \in P^k\}.$$

Then

$$\operatorname{prob} \left( \lim_{k \rightarrow \infty} f_{\text{best}}^k = f^* \right) = 1 \quad \text{and} \quad \lim_{k \rightarrow \infty} \operatorname{prob}(P^k \cap X^* \neq \emptyset) = 1.$$

\* Similar results hold for unencoded or other encodings

# Final Remarks

# Genetic Algorithms

## Pros:

- Very popular
- Very flexible
- Theoretical proof that GA probably converges

## Cons:

- Requires infinite time
- No way to know when you can stop
  - Popular stopping methods are fixed budget and improvement stagnation

## Conclusion

- Not really a DFO method
- Can be useful for BBO

# Assignment 3

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## MATH 462

- Textbook # 4.1, 4.4, 4.5, 4.8, 4.9

## COSC 419K

- Textbook # 4.2, 4.3, 4.4, 4.5, 4.8

## MATH 562

- All MATH 462 and COSC 419K questions
- Textbook # 4.6b, 4.11