

Genetic & Evolutionary Algorithms: Further Investigation on Selection

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**Further Detailed
Investigation on
Selection(2)**





New Population (1)



❖ Now, isn't there any problem at all in Original GA?

- **Original GA** uses a **generational replacement** strategy
 - ✓ Selection, crossover, mutation are applied to **N individuals** (i.e., population)
 - ✓ This set becomes a **new population**
- This process seems somewhat weird in view of **optimization**
 - ✓ We have spent big effort, **getting a good solution**, to run the **risk of throwing it away**

Fitness Order:
Red > Blue > Green > Black

Degenerative Evolution!

I_1	110 ... 101
I_2	101 ... 100
I_3	011 ... 100
I_4	010 ... 111
I_5	101 ... 011
I_6	011 ... 001

i th Population

I_1	110 ... 101
I_2	101 ... 100
I_1	110 ... 101
I_3	011 ... 100
I_2	101 ... 100
I_6	011 ... 001

Selection

110 ... 100
100 ... 101
110 ... 100
011 ... 101
101 ... 001
011 ... 100

Crossover+Mutation

$I_1 > 0_1$	110 ... 100
$I_1 > 0_2$	100 ... 101
$I_1 > 0_3$	110 ... 100
$I_1 > 0_4$	011 ... 101
$I_1 > 0_5$	101 ... 001
$I_1 > 0_6$	011 ... 110

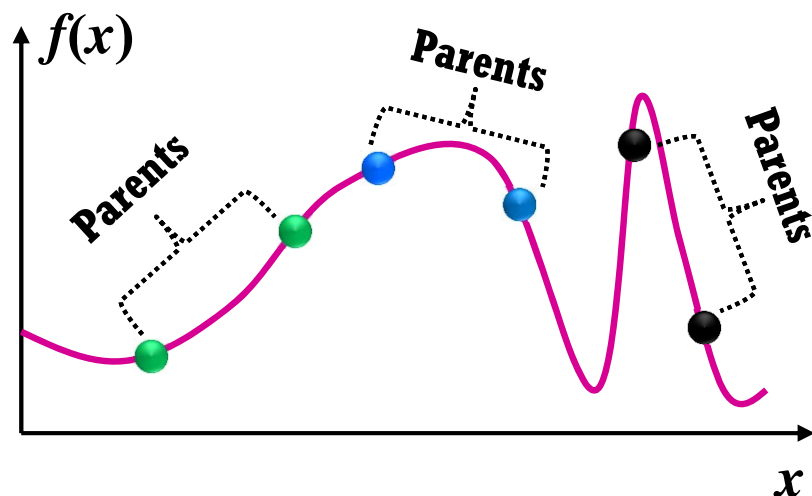
$(i+1)$ th Population



New Population (2)

What's Degenerative Evolution?

- ✓ We have **spent big effort**,
getting a good solution,
to run the **risk of throwing it away**
→ **Very good solutions found so far can be lost!**

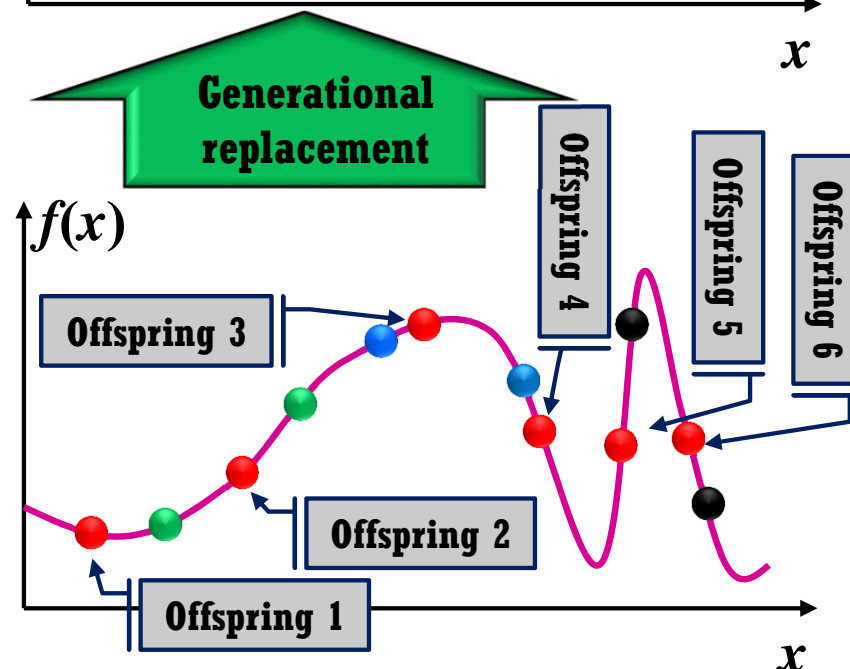
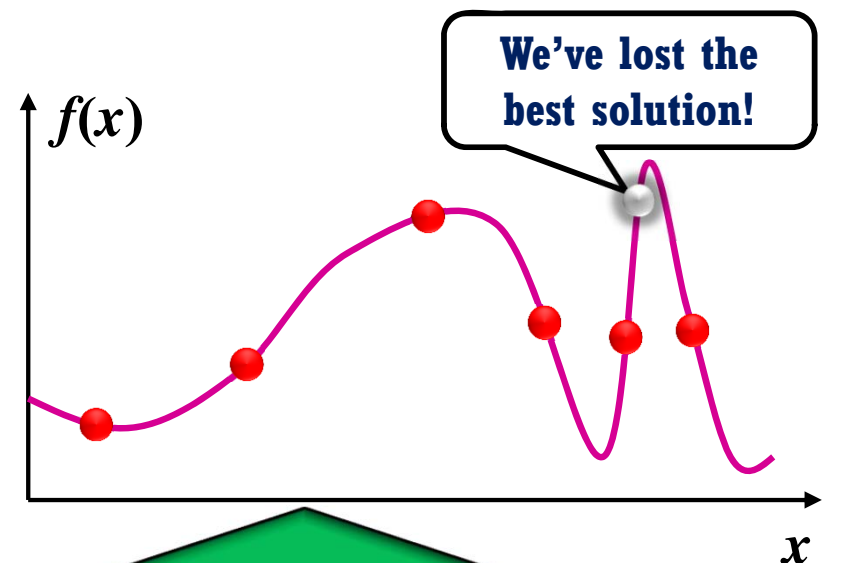


Selected Population

Crossover
+ Mutation



New Population



Crossoverd/Mutated Population

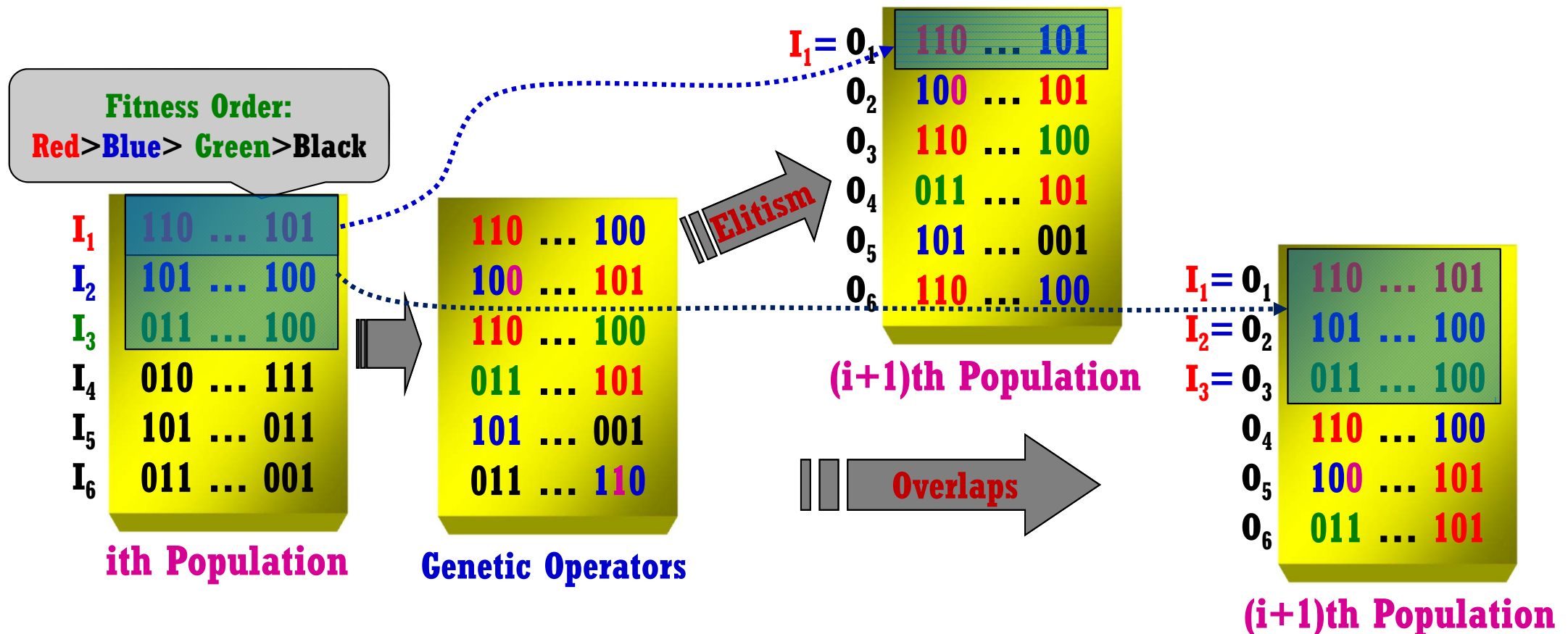


New Population (3)

❖ Any Idea for Resolving Degenerative Evolution?

➤ Use the concept of *Elitism* and *Population Overlaps*

- ✓ **Elitism** ensures the **survival** of the **best** individual
- ✓ **Population overlaps** replace only a **fraction G** (generation gap) of the population



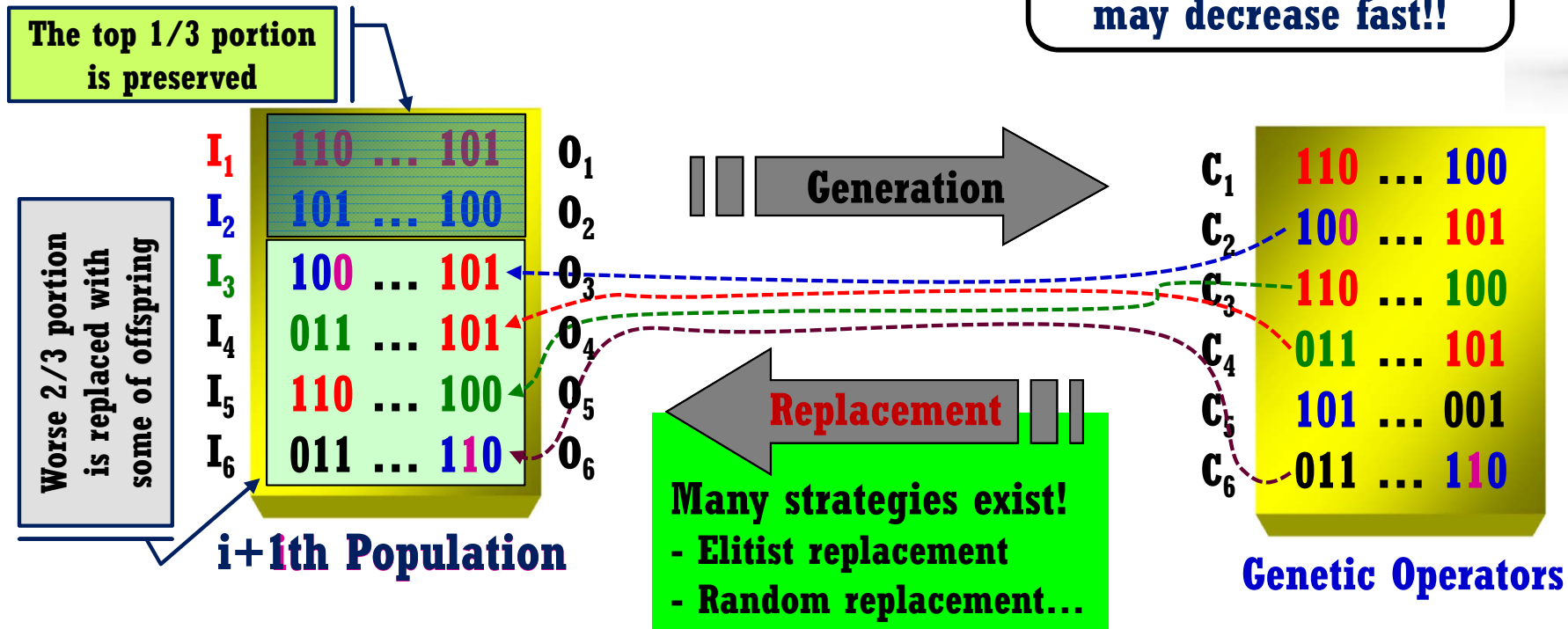


New Population (4)

❖ More General Elitism/Overlap-type Selection?

- The **top τ portion** of the parents are **preserved**
- **Worse $(1 - \tau)$ portion** of the population are **replaced** with **some of offspring**
- ➔ This set becomes a **new (next) population**.

Due to the elitism, the population diversity may decrease fast!!





New Population (5)

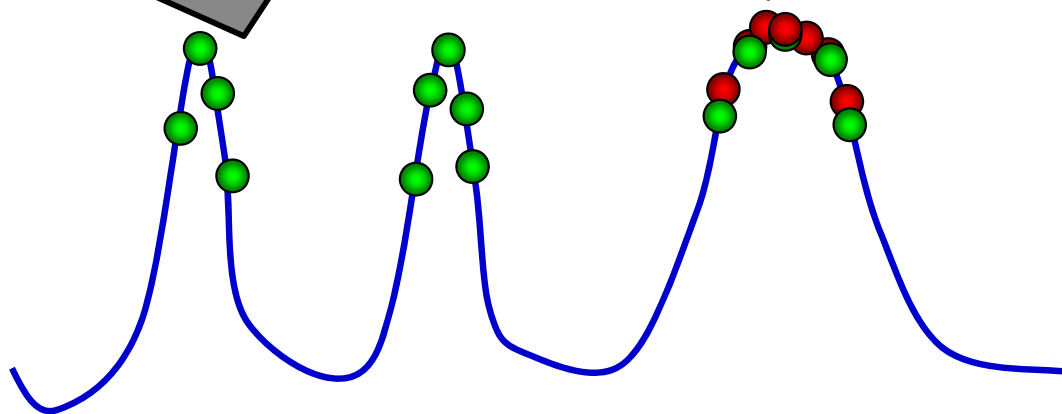
❖ Why Population Diversity Important?

1. Necessary to **cope with the premature convergence**
2. Essential to **discover all solutions in multimodal problems**

Remind Premature Convergence!

We can discover all optimal solutions by maintaining diversity!

A simple GA will find one solution!



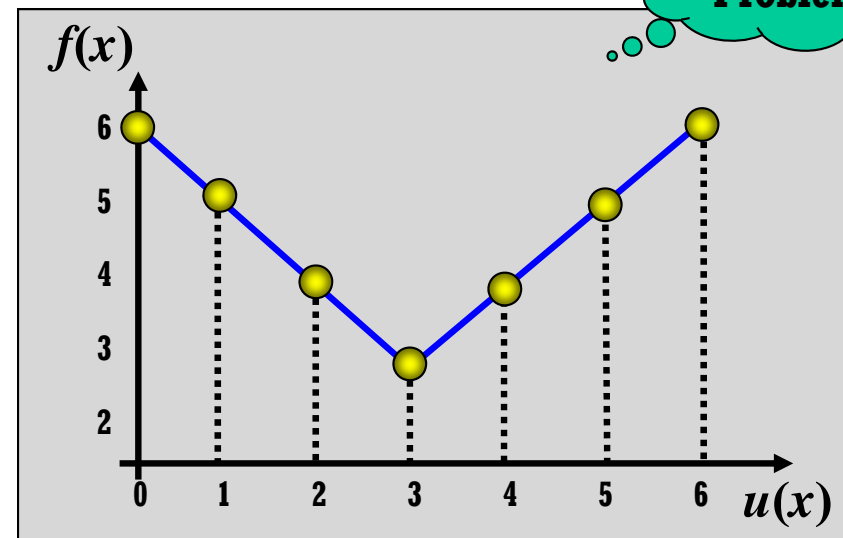
Multimodal problem:
- It has **three optimal** solutions

$$u(x) = \sum x_i, \quad \bar{u}(x) = \sum (1 - x_i), \quad \text{where } x_i \in \{0,1\}$$

Annotations: $\text{count } x_i=1$ points to x_i ; $\text{count } x_i=0$ points to $(1-x_i)$.

$$f(x) = \max \{u(x), \bar{u}(x)\}$$

TwoMax Problem



A multimodal problem $f(x_1, \dots, x_6)$



New Population (6)



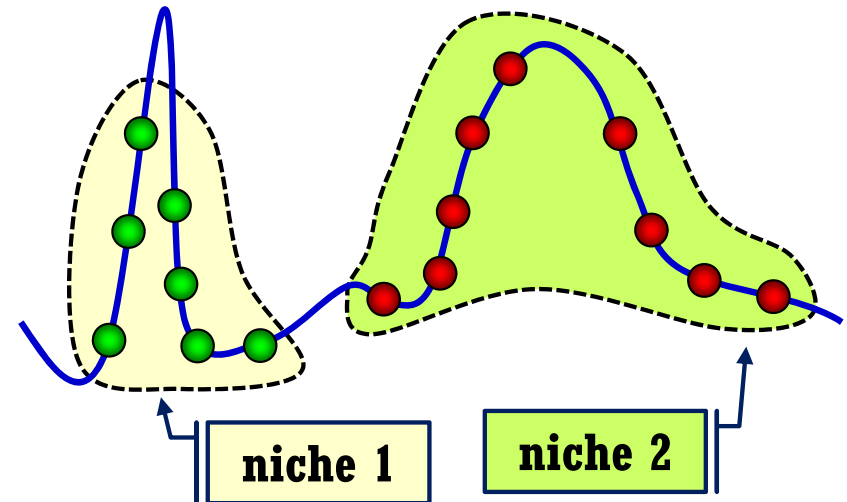
❖ Any Idea for Maintaining the Population Diversity?

Remind Premature Convergence!

- Another aspect of **generating a new population** is the **diversity maintenance**
 - ✓ The concepts of **crowding** and **niching** can be employed:
A **niche** in the nature is a set of conditions to which a species is well adapted.
- In the GA, a **niche** can be treated as a set of **similar individuals**
 - ✓ IDEA: A **newly** generated **individual** replaces one in **its own niche**!



A Real World Example: Tulips



A GA Example



New Population (7)



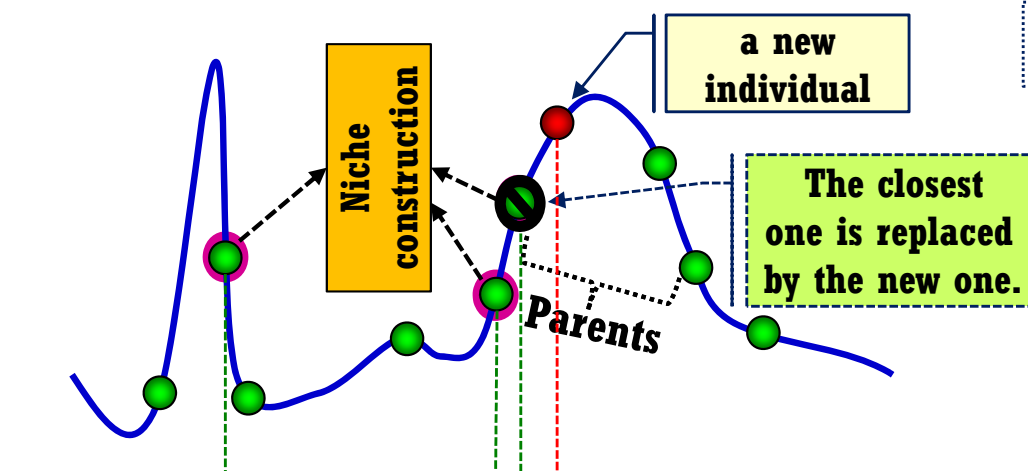
❖ Crowding Method

- A **subset** of individuals **randomly chosen** is constructed (as a **niche**)
- The **closest one** for the new individual is **replaced** with it

❖ Sharing Method

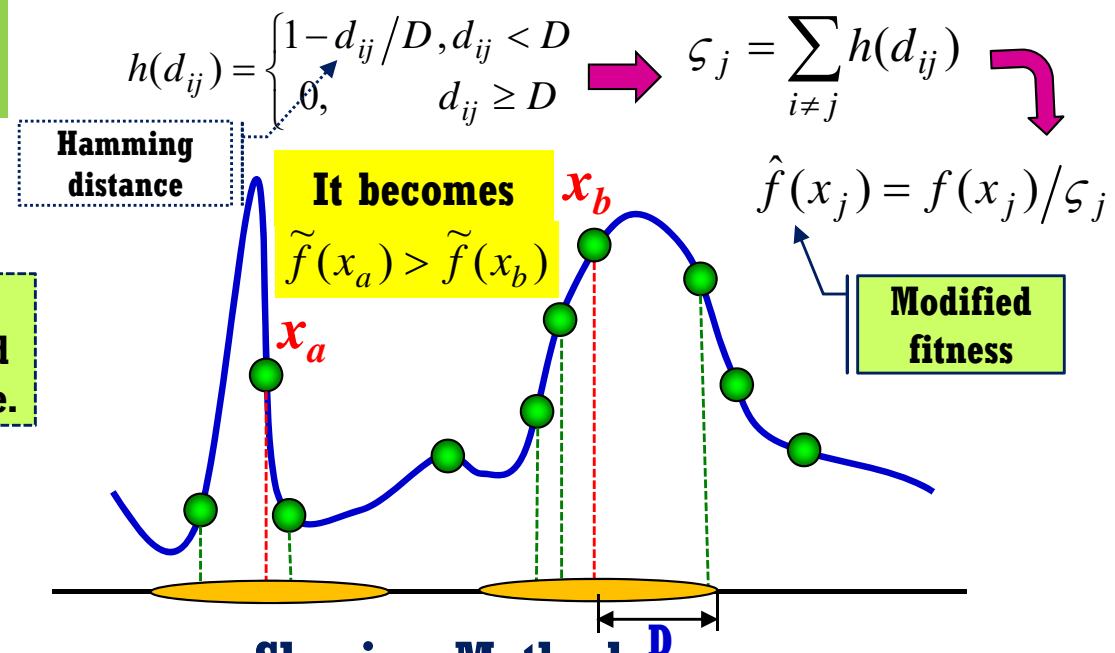
- A **sharing function**, which **adjusts** the raw fitness function, is used; thereby **degrading** individuals that occur in **clusters** comparing to those fairly isolated.

Replacement is done when the new fitness is better; called “Restricted Tournament Replacement”.



Crowding Method

- Apt for handling premature convergence



Sharing Method

- Good at solving multimodal problems

**Further
Studies on
Selection(3)**





Takeover Time (1)

❖ Takeover Time (Convergence Analysis by Applying Selection Alone)

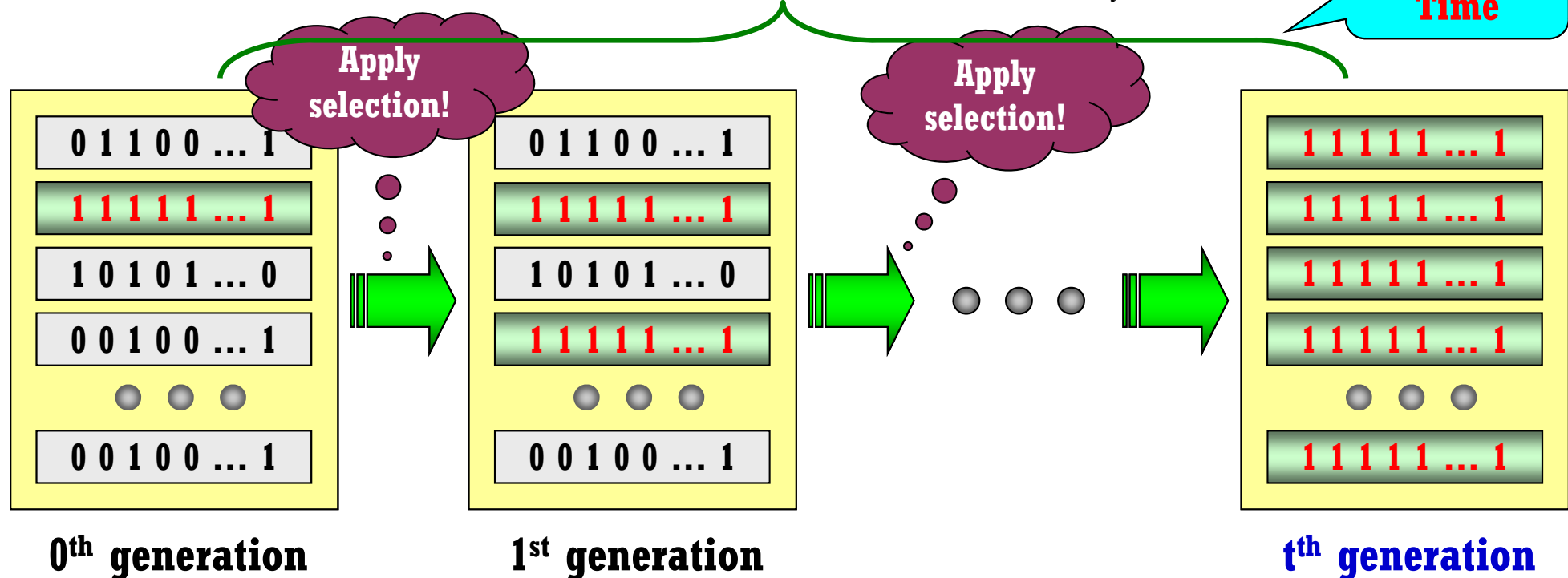
➤ Time from (an) initial best individuals until the population is converged

➤ Assumption:

- ✓ Two alternatives: 0 and 1
- ✓ Initial proportion $P_0=1/N$, Final proportional $P_f=(N-1)/N$
- ✓ The best fitness: f_1 , The average fitness: $\overline{f_t}$

Worst-case Scenario: There is only one best individual!

Takeover Time





Takeover Time (2)

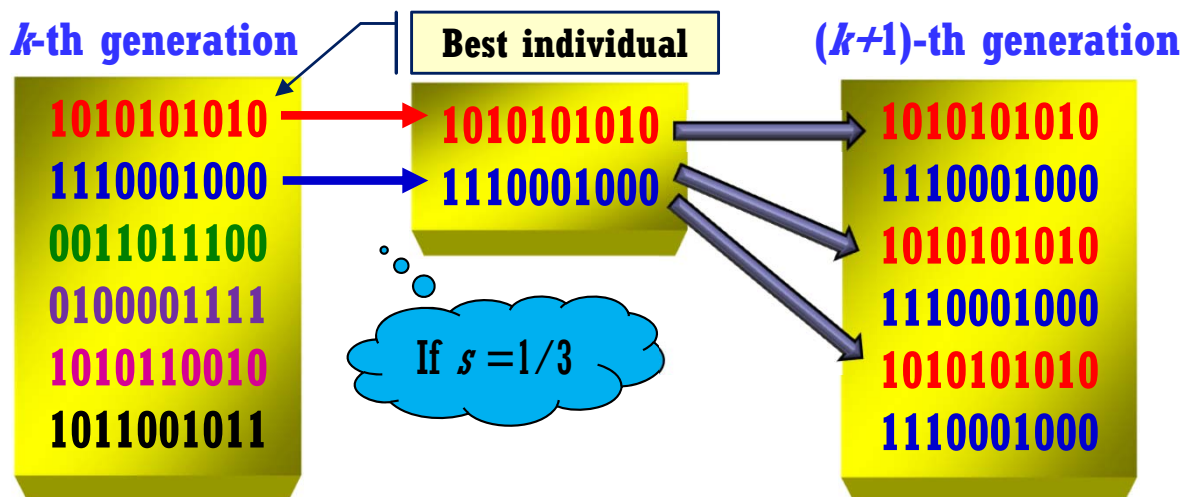


Truncation Selection

- Truncation selection is easy to analyze:
the **top $1/s$** individuals are given **s copies each!**
- If the **proportion P_t** of **best individuals** in a population is less than $1/s$, the growth is geometric:

$$P_{t+1} = sP_t$$

- Once the proportion reaches or exceeds $1/s$ ($P_t \geq s^{-1}$), it saturates and the final proportion is one: **$P_{t+1}=1$**



- Thus, the **proportion of best individual** under truncation selection alone can be written as:

$$P_t = sP_{t-1} = s^2P_{t-2} = \dots = s^t P_0$$

- Assuming $P_0=1/N$, $P_f=1$, and calculating the number of generations to takeover:

$$1 = s^{t^*} \frac{1}{N} \rightarrow s^{t^*} = N \rightarrow t^* \ln s = \ln N$$

- Thus, we have

$$t^* = \frac{\ln N}{\ln s} = O(\ln N)$$

A population is taken to converge as little as $O(\log N)$ generations!



Takeover Time (3)



Proportional Selection

You don't have to follow up all the derivations in detail.

- Suppose we have some number of distinct individuals with **objective function values** $f_j, j \in J$
- The **proportion** of the i th individual at time t ($P_{i,t}$) is related to the **initial proportion** of the other individuals and their function values as follows:

$$P_{i,t} = \frac{f_i^t P_{i,0}}{\sum_{j \in J} f_j^t P_{j,0}}$$

- We restrict ourselves to the unit interval and track the **proportion** of the **best individuals**, $P_{Best,t}$, where $Best = \{x: 1 - 1/N \leq x \leq 1\}$

$$P_{Best,t} = \frac{\int_{1-N^{-1}}^1 f^t(x) p_0(x) dx}{\int_0^1 f^t(x) p_0(x) dx}$$

- With $p_0 = \text{constant}$, and $f(x) = x^c$, we get

$$P_{Best,t^*} = 1 - (1 - 1/N)^{ct^*+1}$$

- Assuming a **final proportion** of best individuals $P_f = (N-1)/N$, we have

$$t^* = \frac{1}{c} (N \ln N) = O(N \ln N)$$



Takeover Time (4)



Tournament Selection

i.e., Ordinal Selection

- τ -wise tournament selection is considered:
Draw τ individuals, and Select the best one!

- The case the best individual is copied into the next generation:
If the best individual is drawn at least one time among τ times,

$$P_{t+1} = 1 - (1 - P_t)^\tau$$

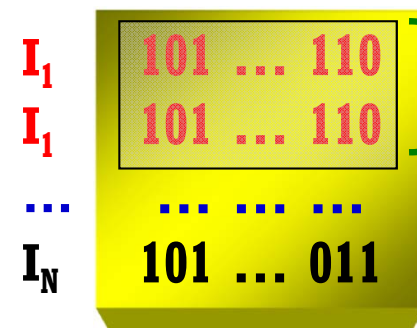
$$1 - P_{t+1} = (1 - P_t)^\tau$$

- We use the complementary proportion $Q_t = 1 - P_t$

$$Q_{t+1} = Q_t^\tau \Rightarrow Q_t = \underbrace{(((Q_0)^\tau)^\tau \cdots)^\tau}_{t \text{ times}} = Q_0^{\tau^t}$$

- By the complementary proportion, $Q_0 = (N-1)/N$, $Q_t = Q_{t^*} = 1/N$ and taking the natural log:

$$\ln(1/N) = \tau^{t^*} \ln((N-1)/N)$$



ith Population

$P_t \rightarrow P_{t+1}$ equals the probability that the best individual is picked and survives (by τ -wise tournament) under P_t

- Recognizing that $\ln(1-x) \cong -x$ for small x

$$-\ln N = \tau^{t^*} \ln(1 - 1/N)$$

$$\ln N = \tau^{t^*} (1/N)$$

- Taking the natural log again:

$$\ln \ln N = t^* \ln \tau - \ln N$$

$$t^* = \frac{\ln N + \ln \ln N}{\ln \tau} = O(\ln N)$$

Last But Not Least
Issues on
Selection(3)





Drift Time

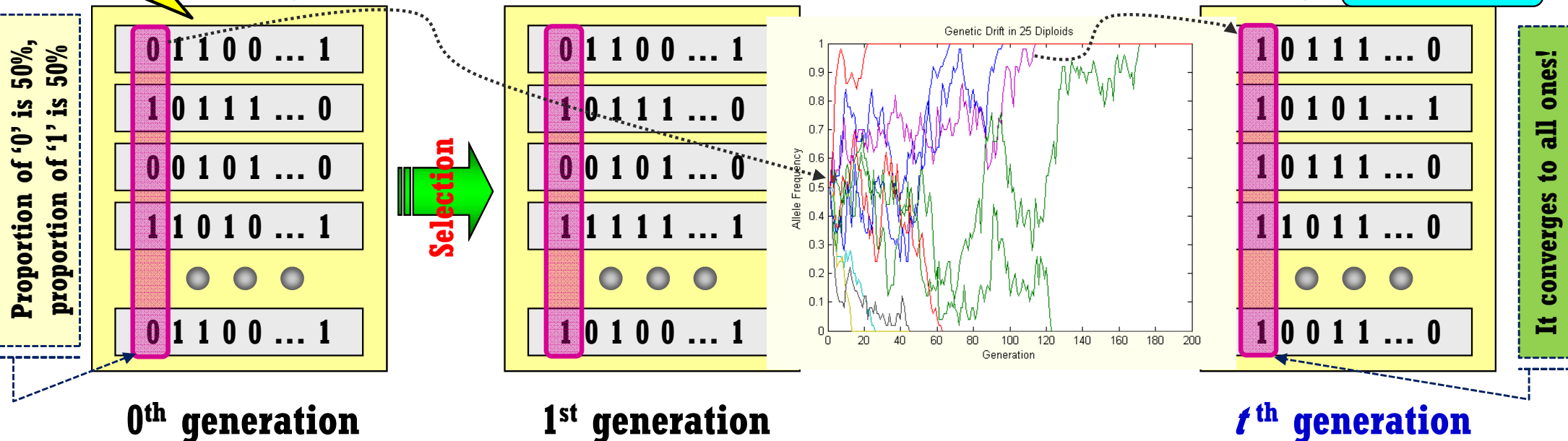


❖ Drift Time (Convergence when Selection chooses for No Reason)

- **Selection** can cause **convergence** in **finite population** with **little fitness difference**
 - ✓ Assume a population initialized with 50 ones and 50 zeros where $f_1 = f_0$.
 - ✓ In a **random trial**, the population (allele) wanders about from the initial condition until it finally comes to **converge to zeros**
 - ✓ In **another trial**, the population (allele) is equally likely **to converge to all ones**
- ➔ Due to the **stochastic errors** in **selection**, which causes the proportion to wander and eventually this wandering **accumulates** to the point where **an allele converges**

All fitness are equal!

Drift Time

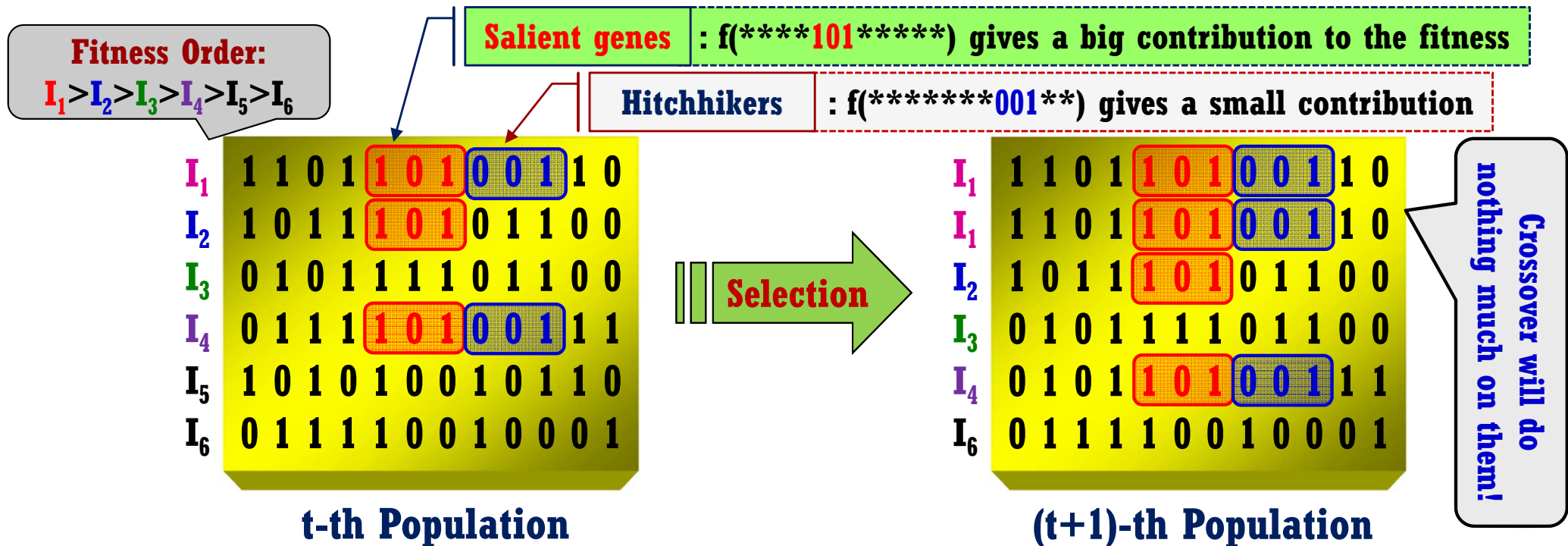




Hitchhiking

❖ Hitchhiking Problem

- In general, **GA** suffers from the **Hitchhiking problem**
 - ✓ Once a **high-fitness schema** (i.e., salient genes) is **discovered**, the **unfit genes**, especially those next to the fit parts, **spread along with the fit ones**
 - ✓ This **slows** the **discovery** of **good schemata** in those positions
 - ✓ **Early convergence** to **wrong** schemata **limits** the **effectiveness of crossover**
- ➔ This **degrades** the **performance of GA** w.r.t. optimality as well as convergence





Summary



- ❖ There are **Two Selection Categories!**
 - **Proportional Selection**; e.g., Roulette-Wheel selection, Scaling
 - **Ordinal Selection**; e.g., Ranking selection, Tournament selection, etc.
- ❖ Generally, **Proportional Selection** tends to have **Premature Convergence**
 - Thus, the **scaling method** has been employed.
 - But, it **needs** to do **re-scaling** at every generation.
- ❖ **Ordinal Selection** is quite **robust** in this regard.
 - It can adjust **selection pressure** at a **constant level** what we want.
 - But, the **ranking selection** is somewhat **restricted**.
 - **Tournament selection** does not have such constraints.
- ❖ In terms of **takeover time** (i.e., convergence with selection only)
 - **Proportional Selection** has $O(N \ln N)$, but **Ordinal Selection** has $O(\ln N)$.

Thus, we conclude that the **Tournament Selection is the most promising choice!**