

# Artificial Intelligence

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# Evolution

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*"... no limit to this power of slowly and beautifully adapting each form to the most complex relations of life ... "*

———— *Charles Darwin*

# Example

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**Maximum**  $f(x) = x^2, x \in [1, 31]$

- **Representation**

$$x \in \{0,1\}^5$$

- **Initialization**

1st generation      01101, 11000, 01000, 10011

Interpretation      13,      24,      8,      19

Fitness      169,      576,      64,      361

# Example

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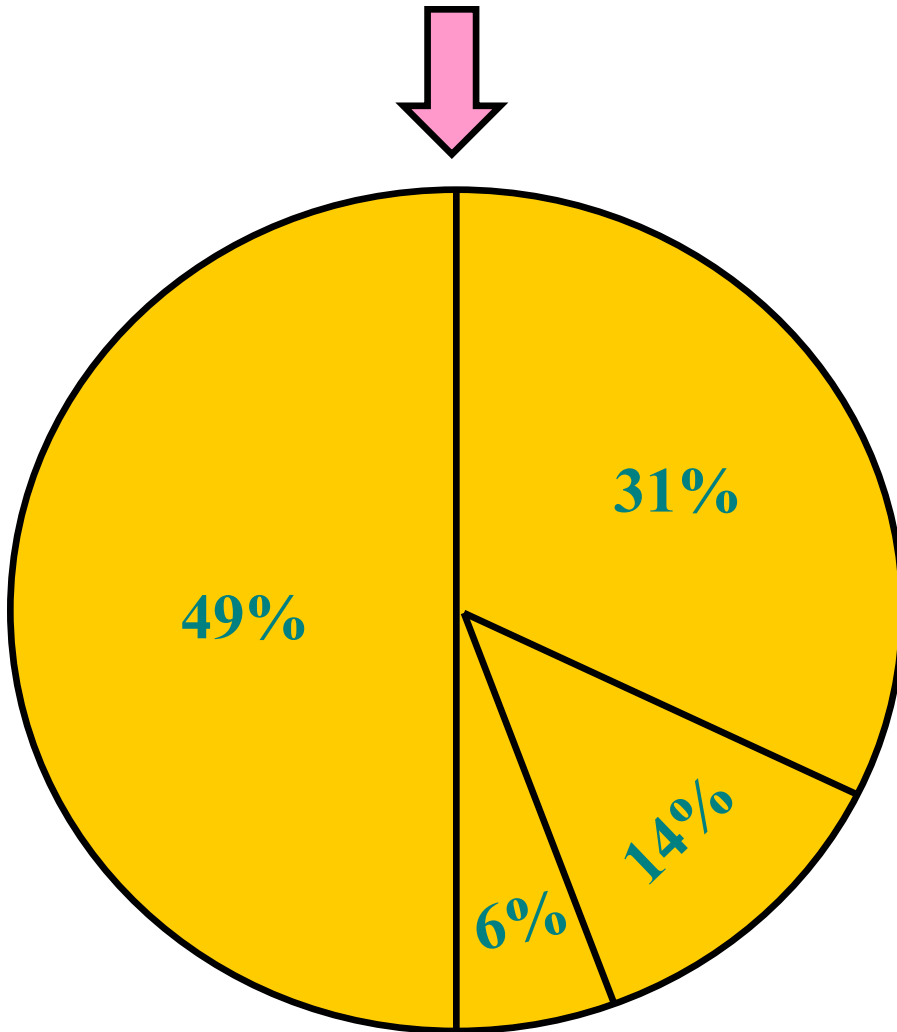
**Maximum**  $f(x) = x^2, x \in [1, 31]$

- **Selection**

Individual	01101, 11000, 01000, 10011
Fitness	169, 576, 64, 361 = 1170
Probability	0.14, 0.49, 0.06, 0.31 = 1.0
Result	01101, 11000, 11000, 10011

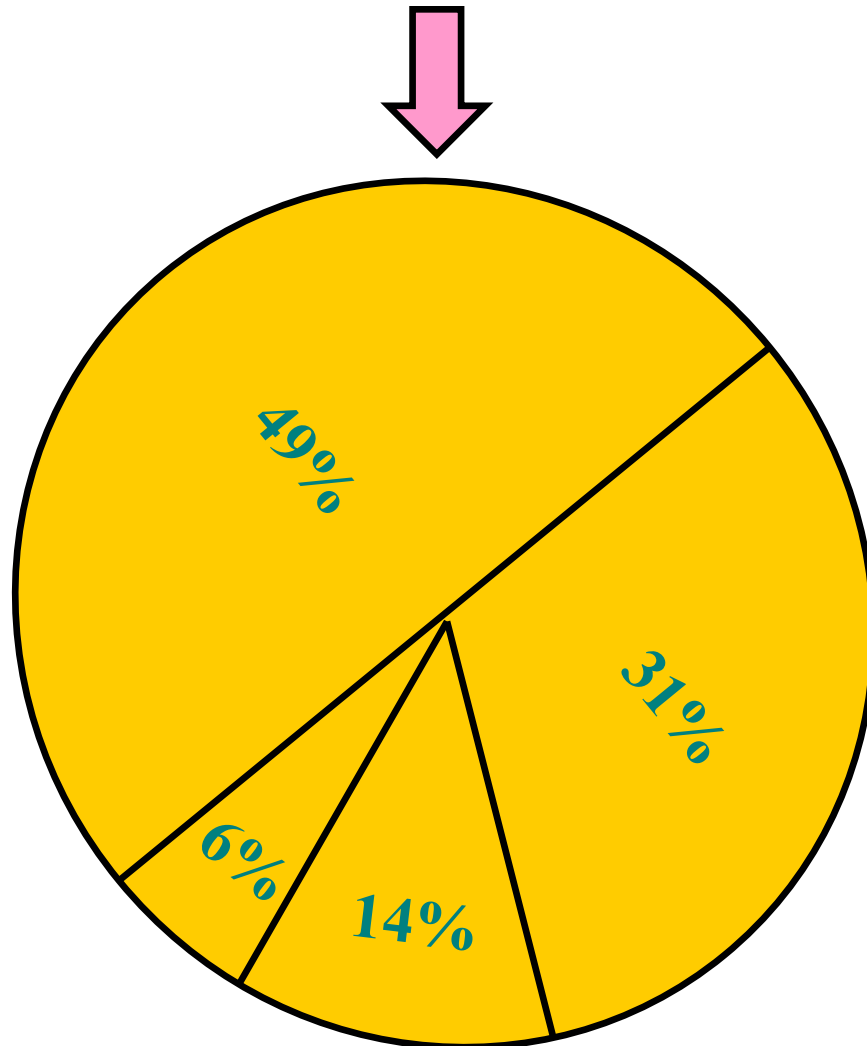
# Russian roulette

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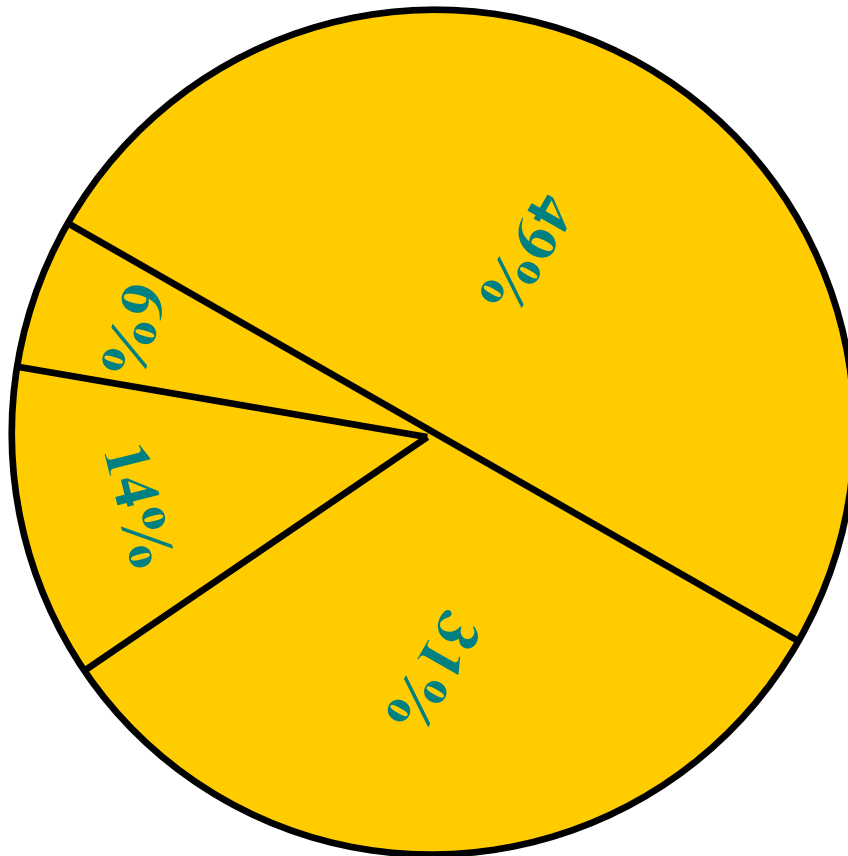
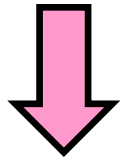
# Russian roulette

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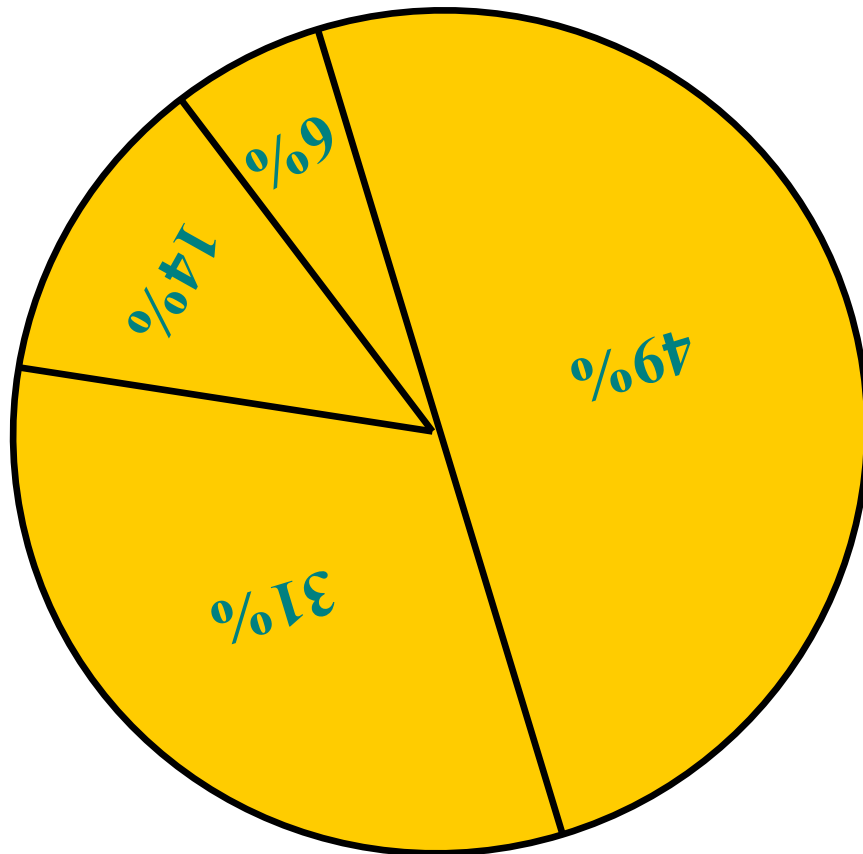
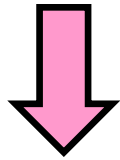
# Russian roulette

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# Russian roulette

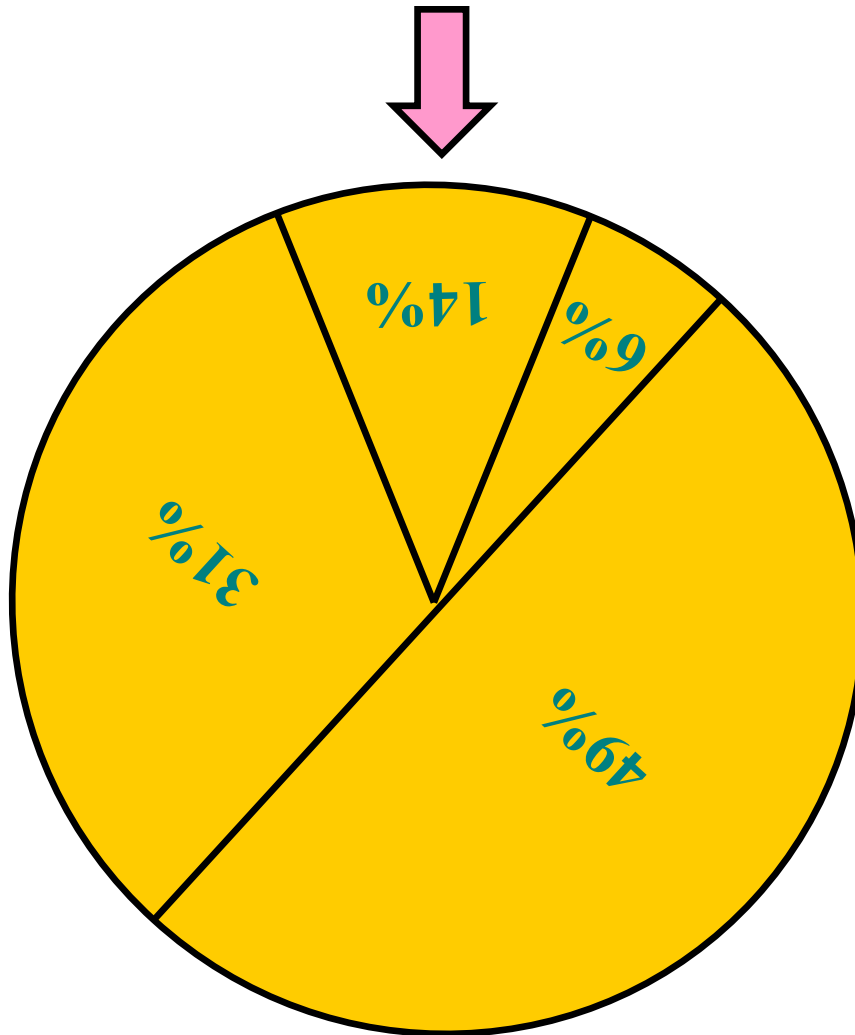
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# Russian roulette

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# Example

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**Maximum**  $f(x) = x^2, x \in [1, 31]$

- **Selection**

Individual	01101, 11000, 01000, 10011
Fitness	169, 576, 64, 361 = 1170
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- **Crossover**

0110 1	⇒	01100	11 000	⇒	11011
1100 0		11001	10 011		10000

- **Mutation**

01100 ⇒ 11100

# Genetic algorithm

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**begin**

set time  $t = 0$

initialize the *population*  $P(t)$

**while** the termination condition is not met **do**

**begin**

evaluate fitness of each member of the population  $P(t)$ ;

select members from population  $P(t)$  based on *fitness*;

produce the *offspring* of these pairs using *genetic operators*;

replace candidates of  $P(t)$ , with these offspring;

set time  $t = t + 1$

**end**

**end**

# Knapsack problem

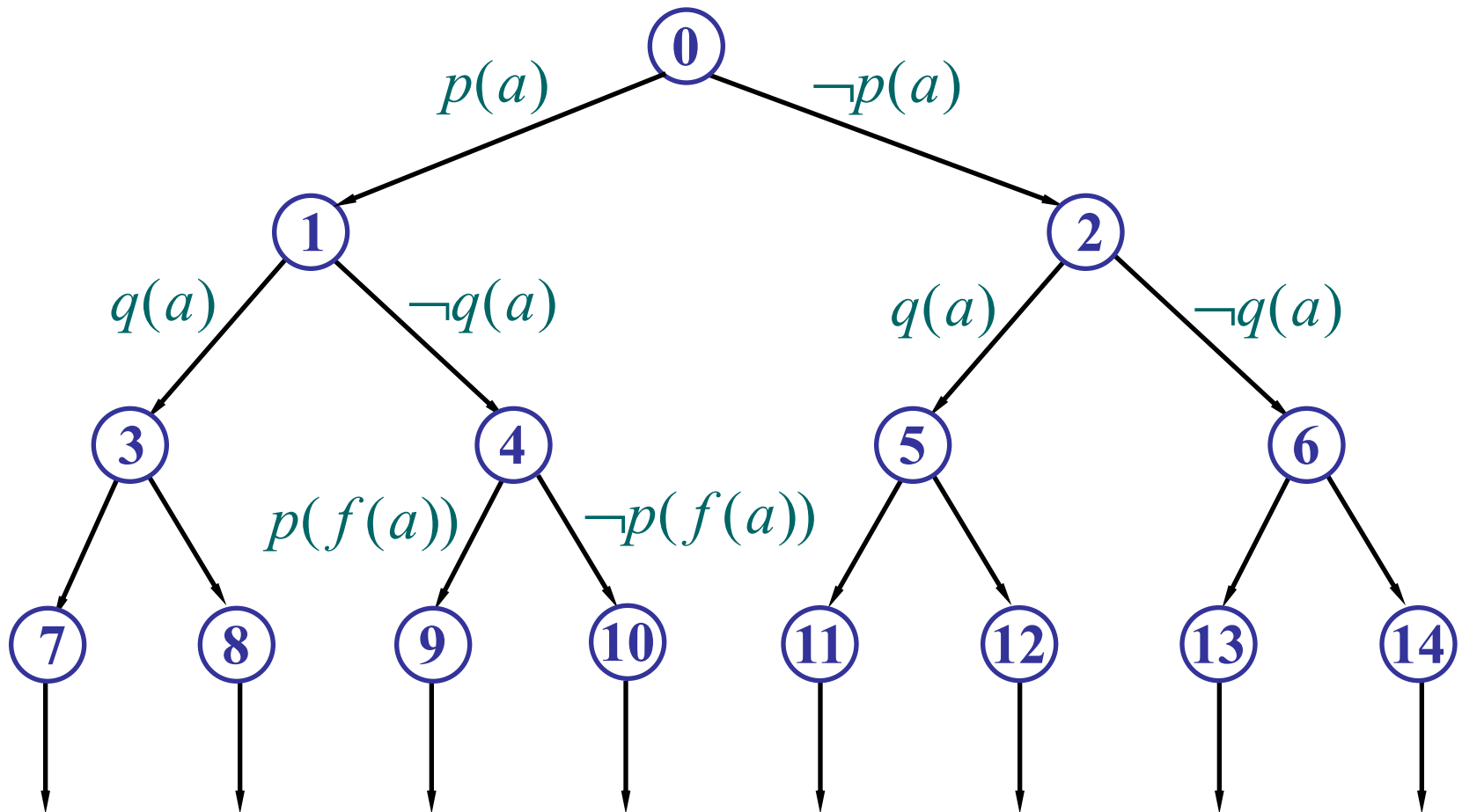
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A thief robbing a store finds  $n$  items; the  $i$ th item is worth  $v_i$  dollars and weights  $w_i$  pounds, where  $v_i$  and  $w_i$  are integers. He wants to take as valuable a load as possible, but he can carry at most  $W$  pounds in his knapsack for some integer  $W$ .

*Which items should he take?*

# Semantic tree

$\{p(x), \neg p(x) \vee q(x), \neg q(f(a))\}$



# CNF-satisfaction problem

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The *conjunctive normal form* (CNF) satisfiability problem is straightforward: an expression of propositions is in conjunctive normal form when it is a sequence of clauses joined by an **AND** relation. Each of these clauses is in the form of a disjunction, the **OR** of literals.

$$(\neg a \vee c) \wedge (\neg a \vee c \vee \neg e) \wedge (\neg b \vee c \vee d \vee \neg e) \wedge \\ (a \vee \neg b \vee c) \wedge (\neg e \vee v)$$

# Analysis of traveling salesman problem

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Given a finite number of "cities" along with the cost of travel between each pair of them, find the *cheapest way* of visiting each city exactly once and finishing at the city he starts from.

- 19 Cities
- Possible routes =  $18! = 6.40237 \times 10^{15}$
- life time =  $80 \times 365 \times 24 \times 60 \times 60$   
 $= 2.52288 \times 10^9$
- Computer speed = 10000 routes/second

253.77 *Generation!*

# Traveling salesman problem

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## Crossover

$$p_1 = ( 1 \ 9 \ 2 \ 4 \ 6 \ 5 \ 7 \ 8 \ 3 )$$
$$p_2 = ( 4 \ 5 \ 9 \ 1 \ 8 \ 7 \ 6 \ 2 \ 3 )$$

Start from the second cut point of one parent, the cities from the other parent are copied in the same order, omitting cities already present. When the end of the string is reached, continue on from the beginning. The sequence of cities from  $p_2$  is :

2 3 4 5 9 1 8 7 6

$$c_1 = ( 2 \ 3 \ 9 \ 4 \ 6 \ 5 \ 7 \ 1 \ 8 )$$
$$c_2 = ( 3 \ 9 \ 2 \ 1 \ 8 \ 7 \ 6 \ 4 \ 5 )$$



# Traveling salesman problem

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## Mutation

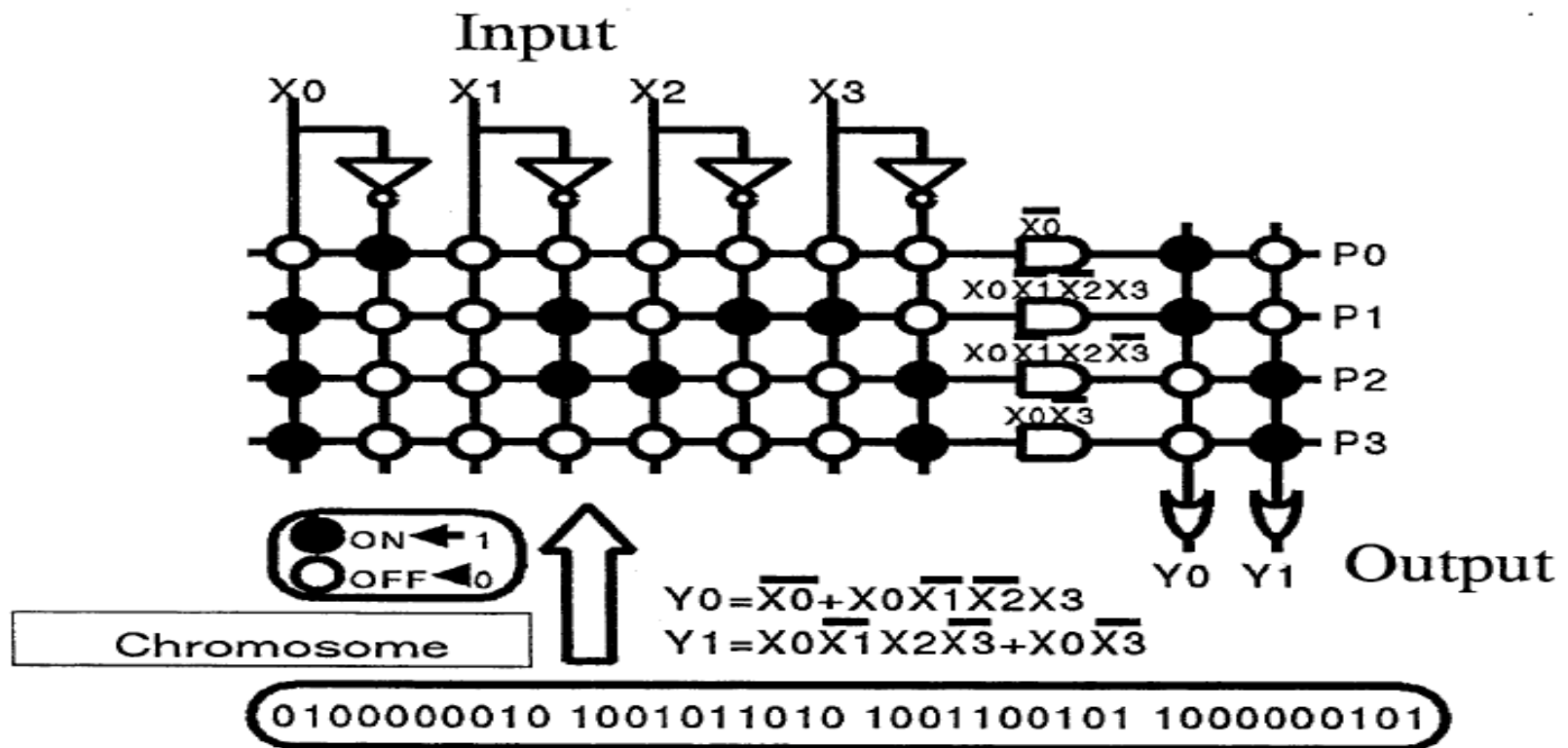
$$c_1 = (2 \ 3 \ 9 \ 4 \ 6 \ 5 \ 7 \ 1 \ 8)$$

$$c_1 = (2 \ 3 \ 9 \ 7 \ 5 \ 6 \ 4 \ 1 \ 8)$$

*Inversion*

# Evolutionary hardware

## PLA (Programmable Logic Array)



# Gray coding

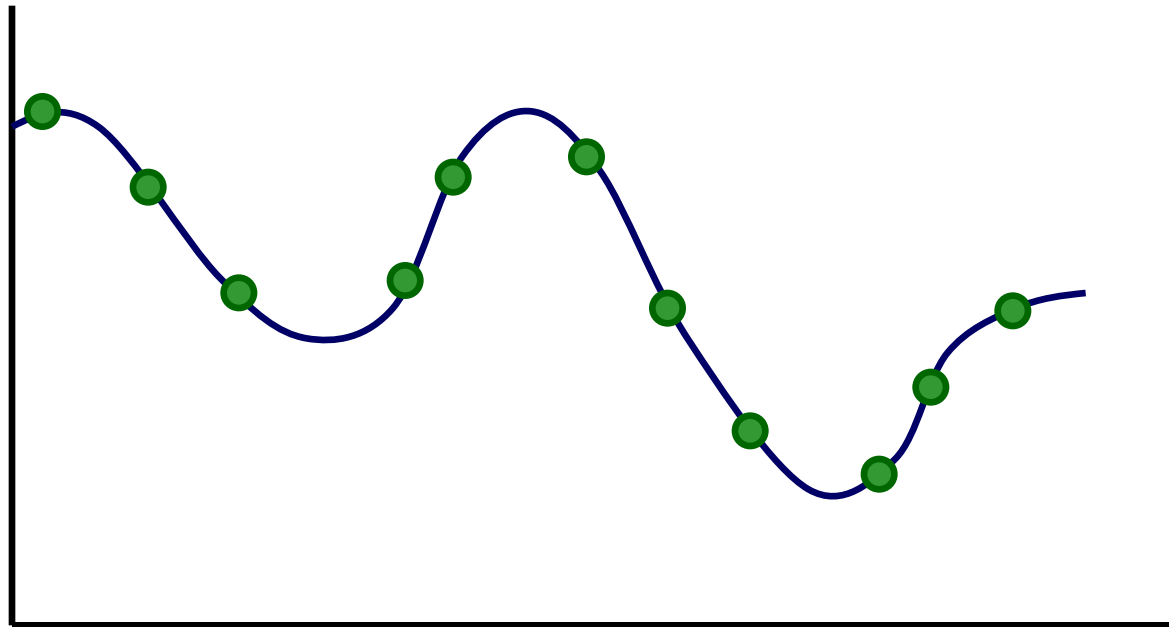
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Binary	Gray	Binary	Gray
0000	0000	1000	1100
0001	0001	1001	1101
0010	0011	1010	1111
0011	0010	1011	1110
0100	0110	1100	1010
0101	0111	1101	1011
0110	0101	1110	1001
0111	0100	1111	1000

# Strength of the genetic algorithms

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*Solution  
Quality*

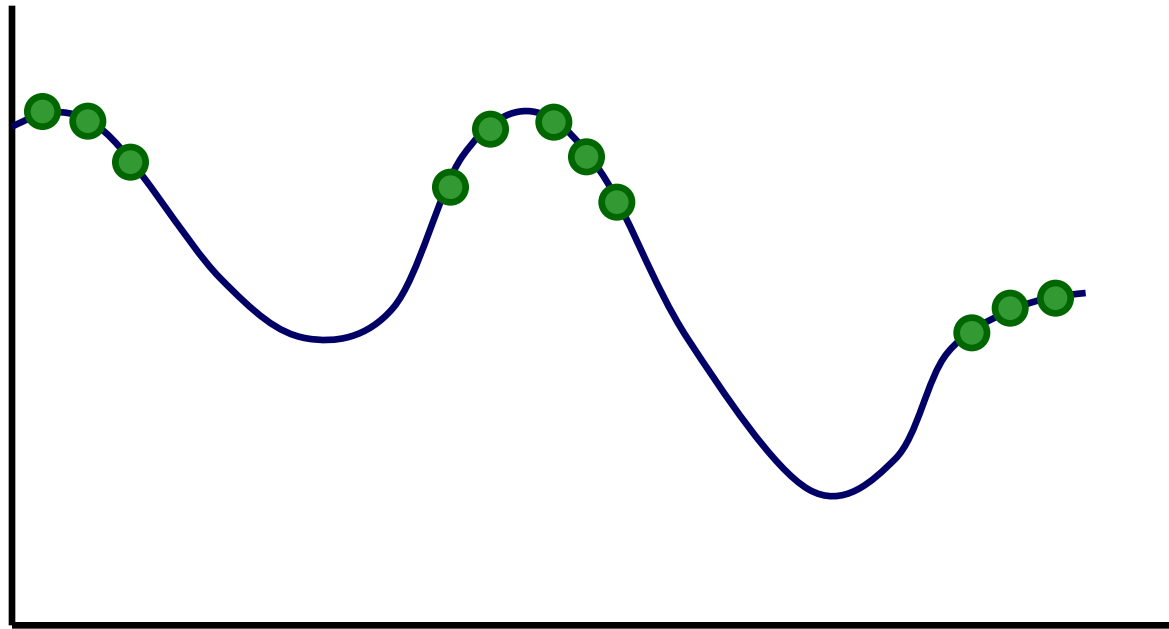


The beginning search space

# Strength of the genetic algorithms

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*Solution  
Quality*

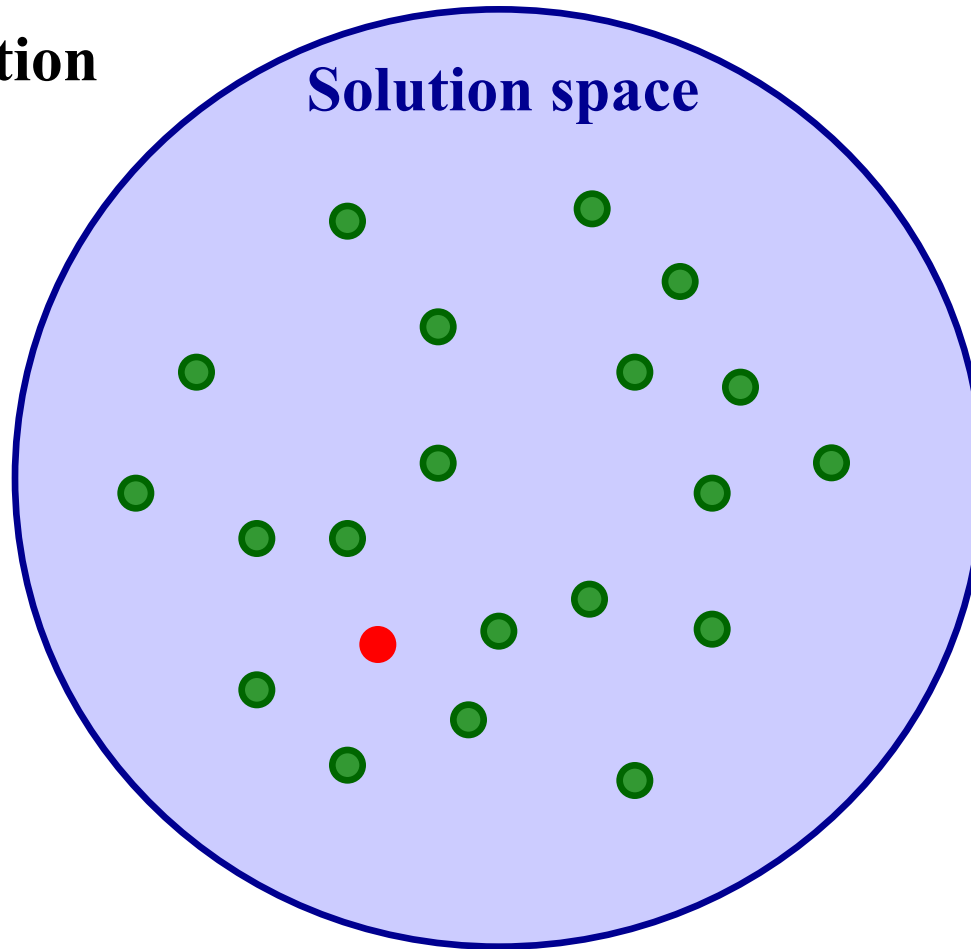


The search space after  $n$  generations

# Random research

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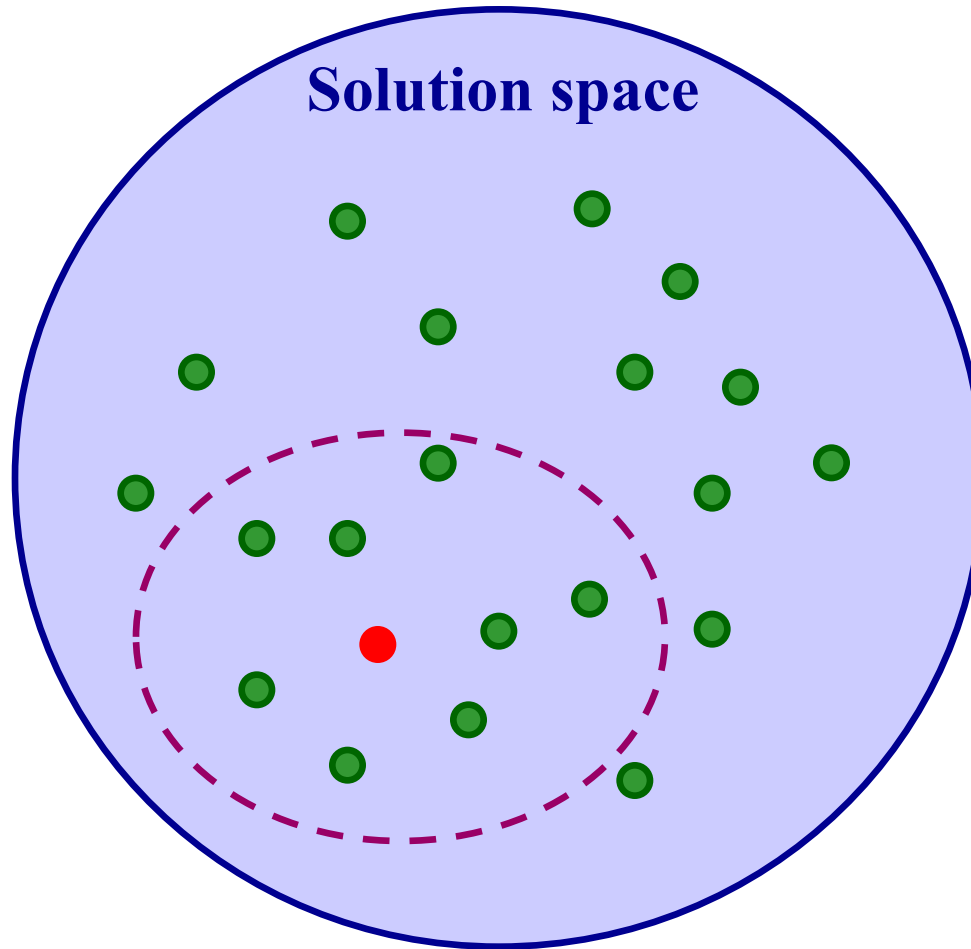
**Initialization**



# Random research

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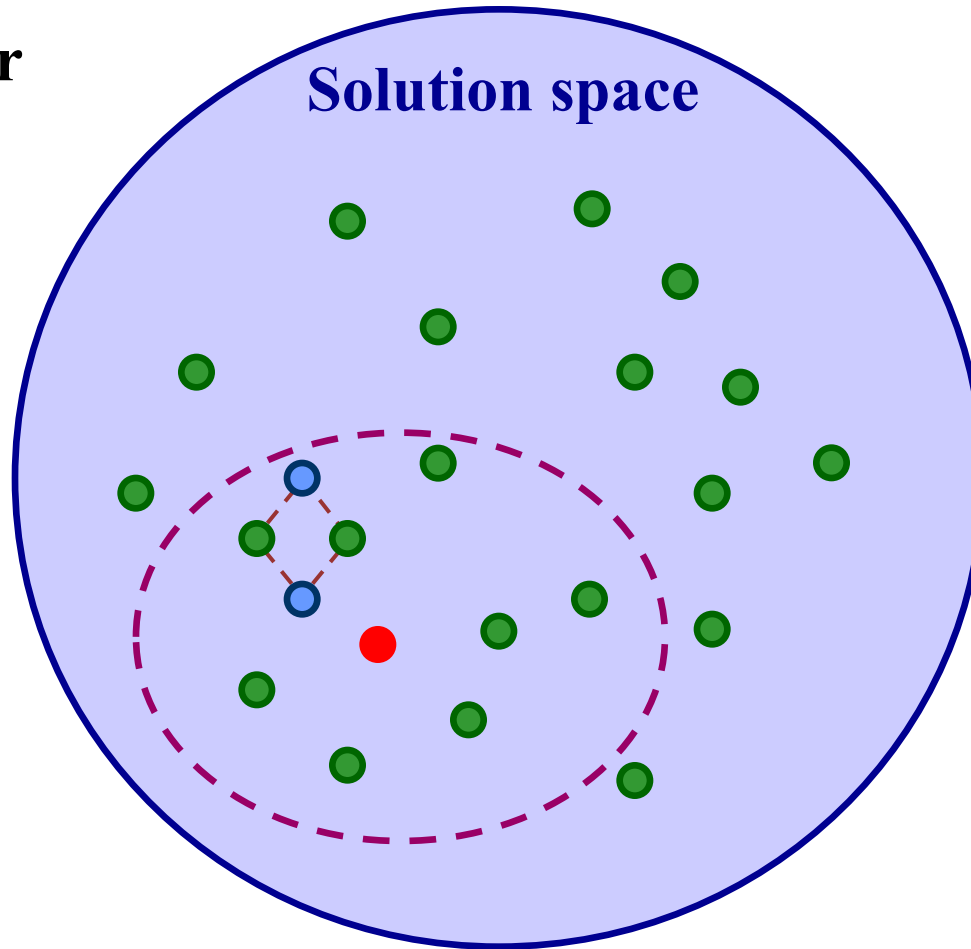
**Selection**



# Random research

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## Crossover

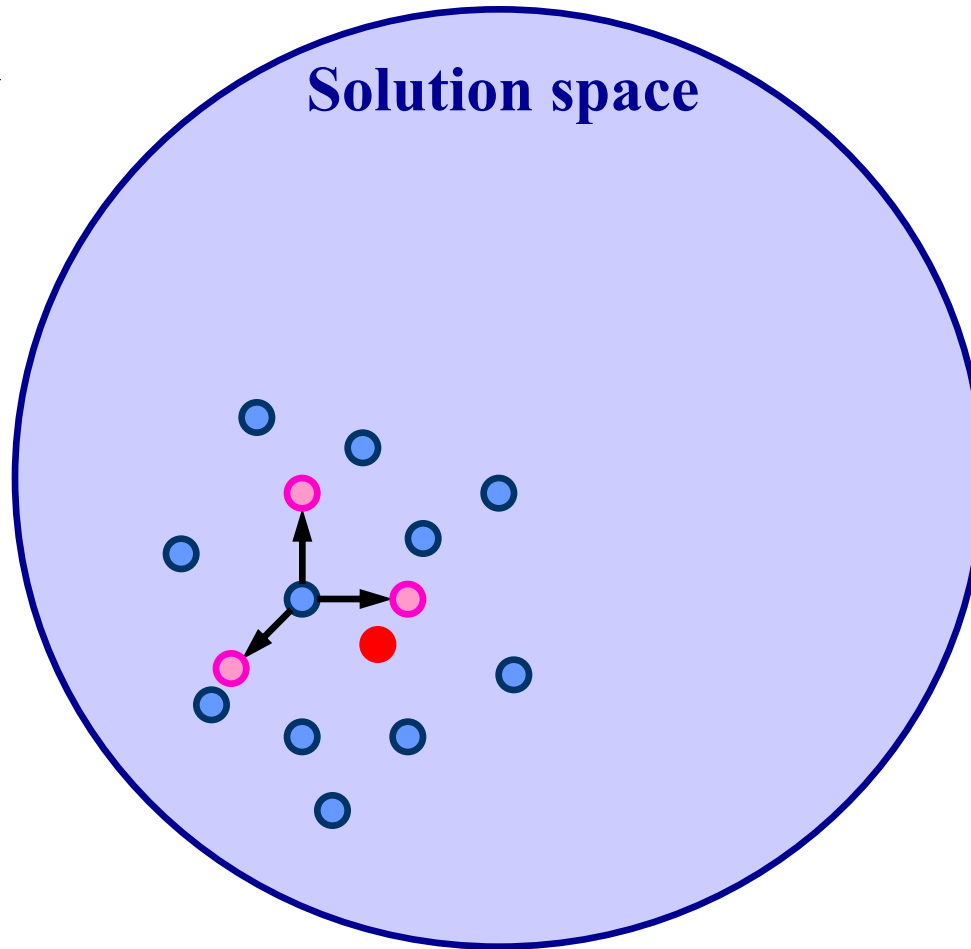




# Random research

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## Mutation



# Schema

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- The schema ( 1 \* \* 0 \* 1 ) describes the set of all strings of length 6 with 1's at positions 1 and 6 and a 0 at position 4. The \* is a *wildcard* symbol, which means that positions 2, 3 and 5 can have a value of either 1 or 0.
- The *order* of a schema,  $O(s)$ , is defined as the number of *fixed positions* in the template, while the defining *length*  $\delta(s)$  is the distance between the first and last specific positions. The order of ( 1 \* \* 0 \* 1 ) is 3 and its defining length is 5.
- The *fitness* of a schema is the *average* fitness of all strings matching the schema. The fitness of a string is a measure of the value of the encoded problem solution, as computed by a problem-specific evaluation function.

# Schema theorem

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The probability of the string  $i$  that is selected:

$$p_i = \frac{f(i)}{\sum_{j=1}^n f(j)}$$

$m(s,t)$ : the number of strings belonging to schema  $s$  at generation  $t$ .

$$E[m(s,t+1)] = m(s,t) \cdot n \cdot \frac{\overline{f(s)}}{\sum_{j=1}^n f(j)}$$

# Schema theorem

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$\bar{f}$  is the average fitness at generation  $t$ .

$$\bar{f} = \frac{\sum_{j=1}^n f(j)}{n}$$

Then,

$$E[m(s, t+1)] = m(s, t) \cdot \frac{\bar{f}(s)}{\bar{f}}$$

Let  $\bar{f}(s) = (1+c)\bar{f}$

$$\begin{aligned} E[m(s, t+1)] &= m(s, t) \cdot \frac{(1+c)\bar{f}}{\bar{f}} \\ &= m(s, t) \cdot (1+c) = m(s, 0) \cdot (1+c)^{t+1} \end{aligned}$$

# Schema theorem

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The probability of disruption  $p_s$  is the probability that crossover will not destroy the schema  $s$ .

$$p_s \geq 1 - \frac{\delta(s)}{l-1}$$

$p_c$  is the probability of crossover.

$$p_s \geq 1 - p_c \cdot \frac{\delta(s)}{l-1}$$

Then,

$$E[m(s, t+1)] = m(s, t) \cdot \frac{\bar{f}(s)}{\bar{f}} \left[ 1 - p_c \cdot \frac{\delta(s)}{l-1} \right]$$

# Schema theorem

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The probability of disruption  $p_s$  is the probability that mutation will not destroy the schema  $s$ , and  $p_m$  is the probability of mutation.

$$p_s = (1 - p_m)^{O(s)}$$

Assuming  $p_m \ll 1$

$$p_s \approx 1 - p_m \cdot O(s)$$

Then,

$$\begin{aligned} E[m(s, t+1)] &= m(s, t) \cdot \frac{\bar{f}(s)}{\bar{f}} \cdot \left[ 1 - p_c \cdot \frac{\delta(s)}{l-1} \right] \cdot [1 - O(s)p_m] \\ &\approx m(s, t) \cdot \frac{\bar{f}(s)}{\bar{f}} \cdot \left[ 1 - p_c \cdot \frac{\delta(s)}{l-1} - O(s) \cdot p_m \right] \end{aligned}$$

# Schema theorem

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- Holland's schema theorem is widely taken to be the foundation for explanations of the power of *genetic algorithms*.
- A *schema* is a template that identifies a *subset* of strings with similarities at certain string positions.
- The schema theorem states that *short, low-order, schemata* with *above-average fitness* increase exponentially in successive generations.

*Any question?*



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