



UiO : **Department of Informatics**
University of Oslo

Biologically inspired computing

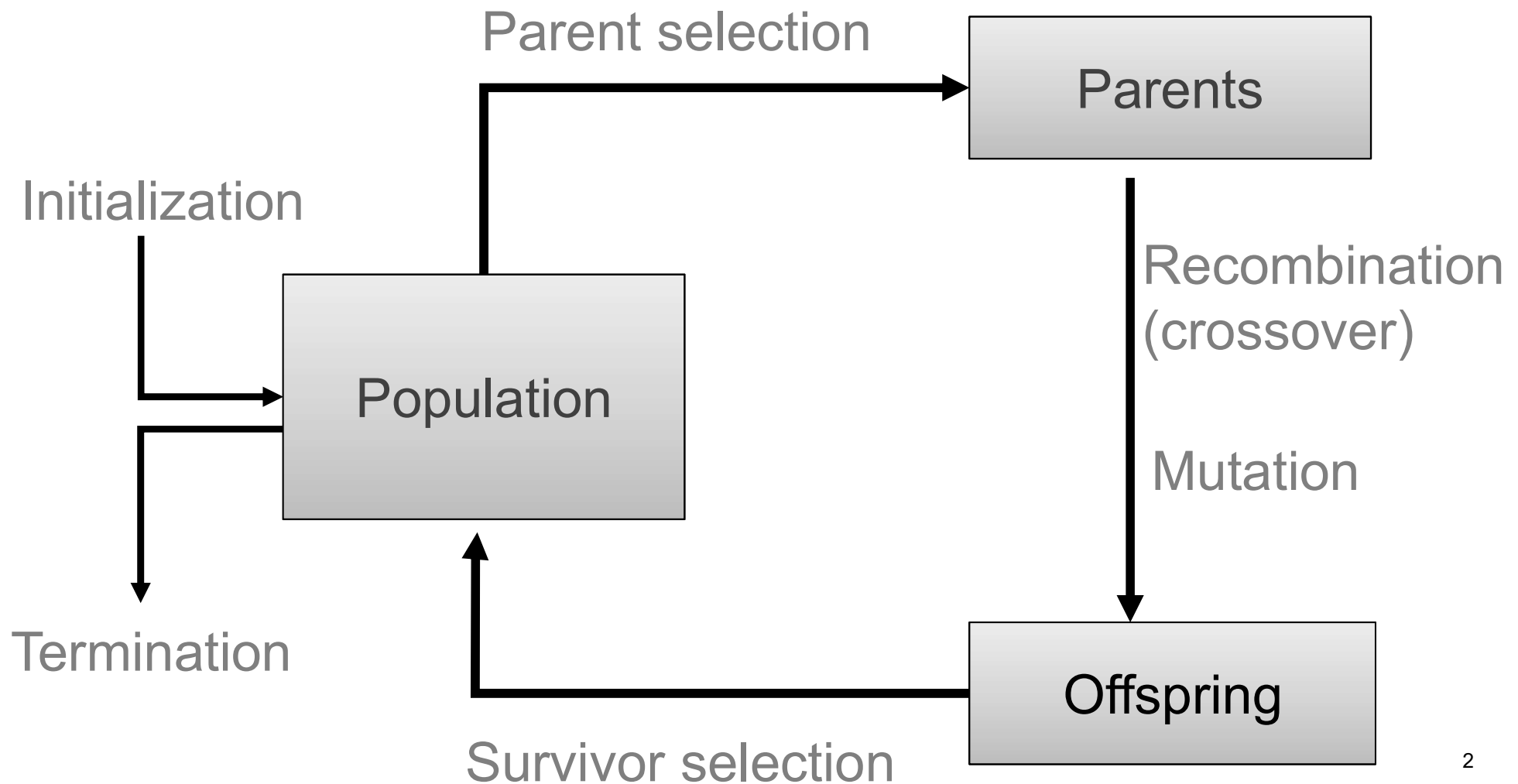
Lecture 3: Eiben and Smith, chapter 5-6

Evolutionary Algorithms - Population management and popular algorithms

Kai Olav Ellefsen



Repetition: General scheme of EAs



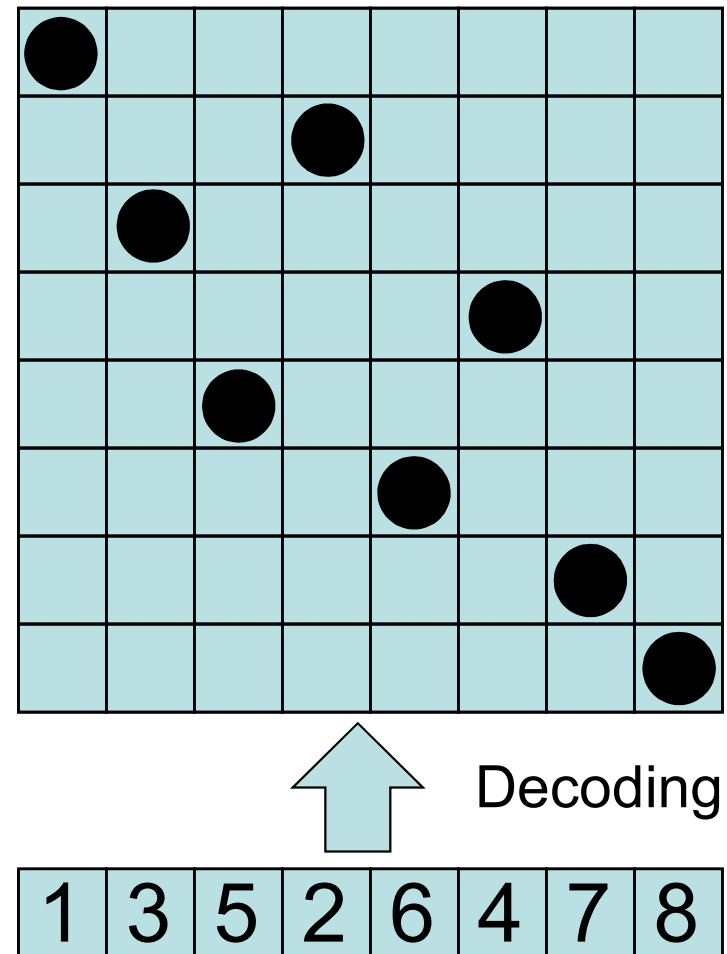
Repetition: Genotype & Phenotype

Phenotype:

A solution representation
we can **evaluate**

Genotype:

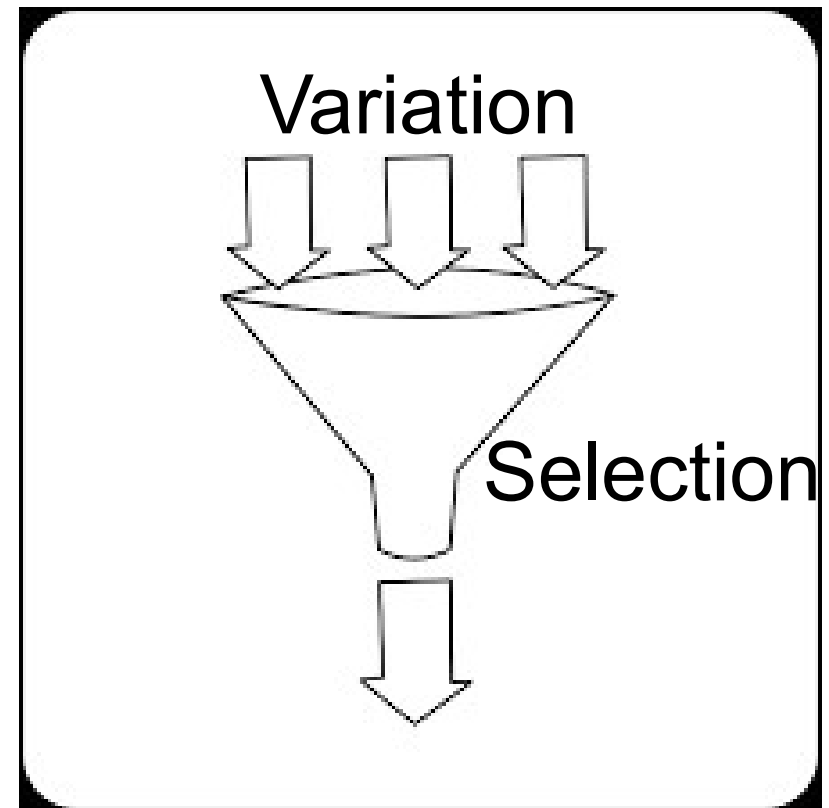
A solution representation
applicable to **variation**



Chapter 5:

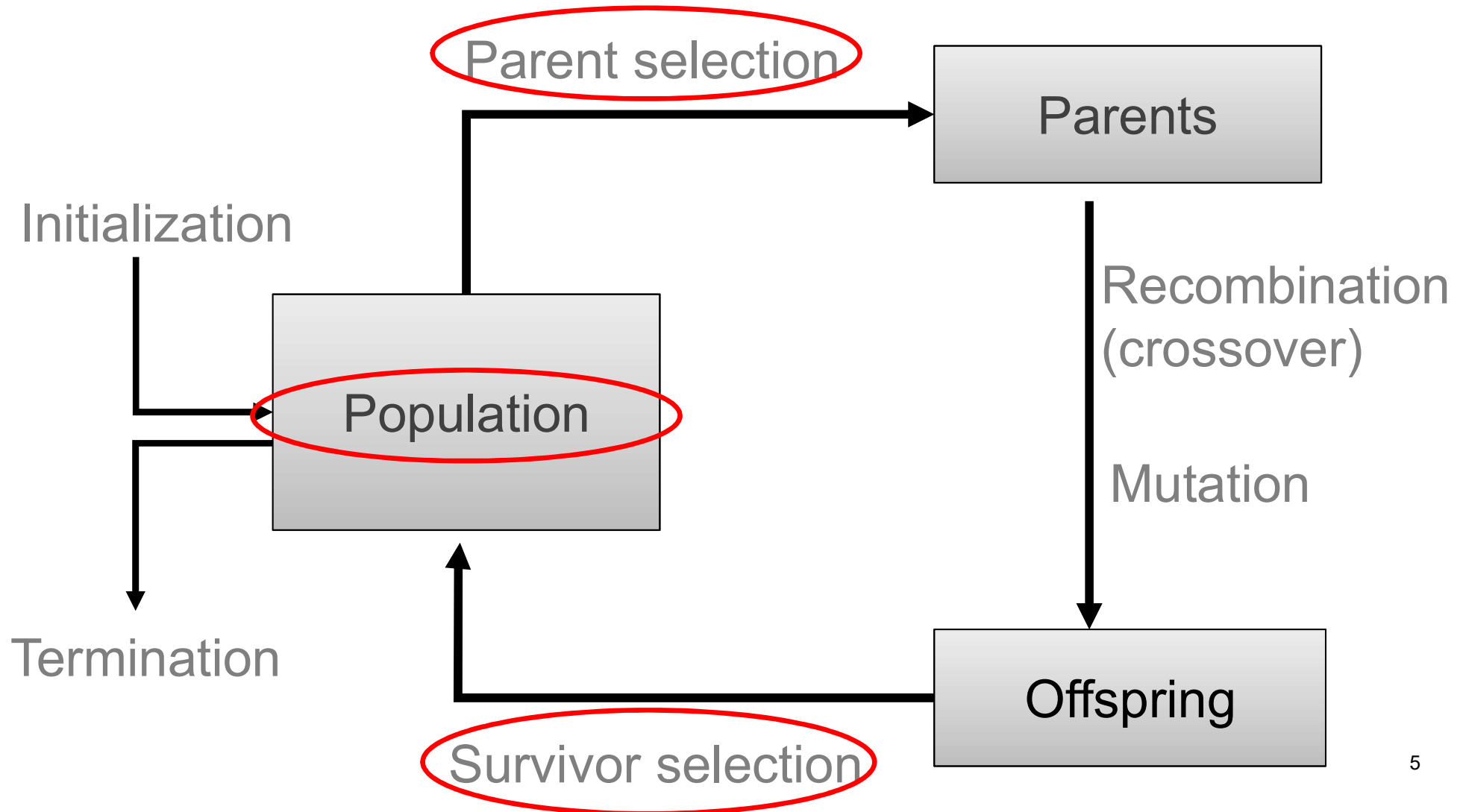
Fitness, Selection and Population Management

- **Selection** is second fundamental force for evolutionary systems
- Components exist of:
 - Population management models
 - Selection operators
 - Preserving diversity



Scheme of an EA:

General scheme of EAs



Population Management Models: Introduction

- Two different population management models exist:
 - **Generational model**
 - each individual survives for exactly one generation
 - λ offspring are generated
 - the entire set of μ parents is replaced by μ offspring
 - **Steady-state model**
 - $\lambda (< \mu)$ parents are replaced by λ offspring
- Generation Gap
 - The proportion of the population replaced
 - Parameter = 1.0 for G-GA, $=\lambda/\text{pop_size}$ for SS-GA

Population Management Models: Fitness based competition

- Selection can occur in two places:
 - **Parent selection** (selects mating pairs)
 - **Survivor selection** (replaces population)
- Selection works on the population
 - > selection operators are representation-independent !
- **Selection pressure:** As selection pressure increases, fitter solutions are more likely to survive, or be chosen as parents

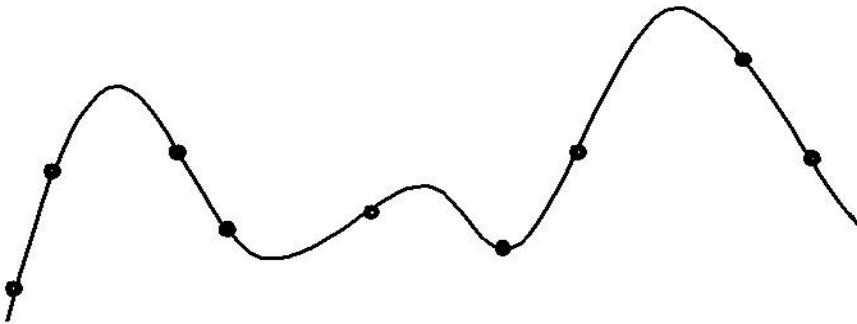
Effect of Selection Pressure

- Low Pressure
- High Pressure

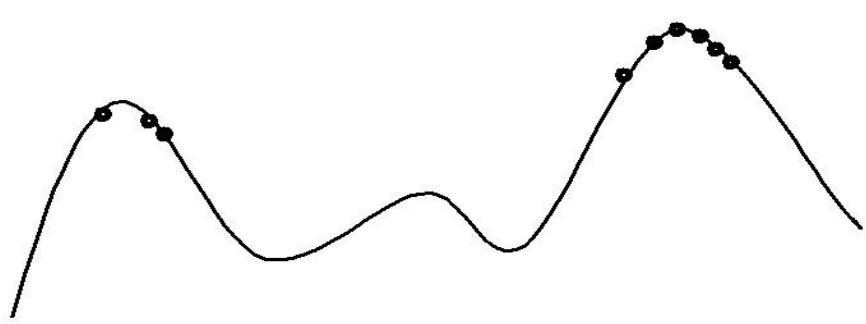


Why Not Always High Selection Pressure?

Exploration

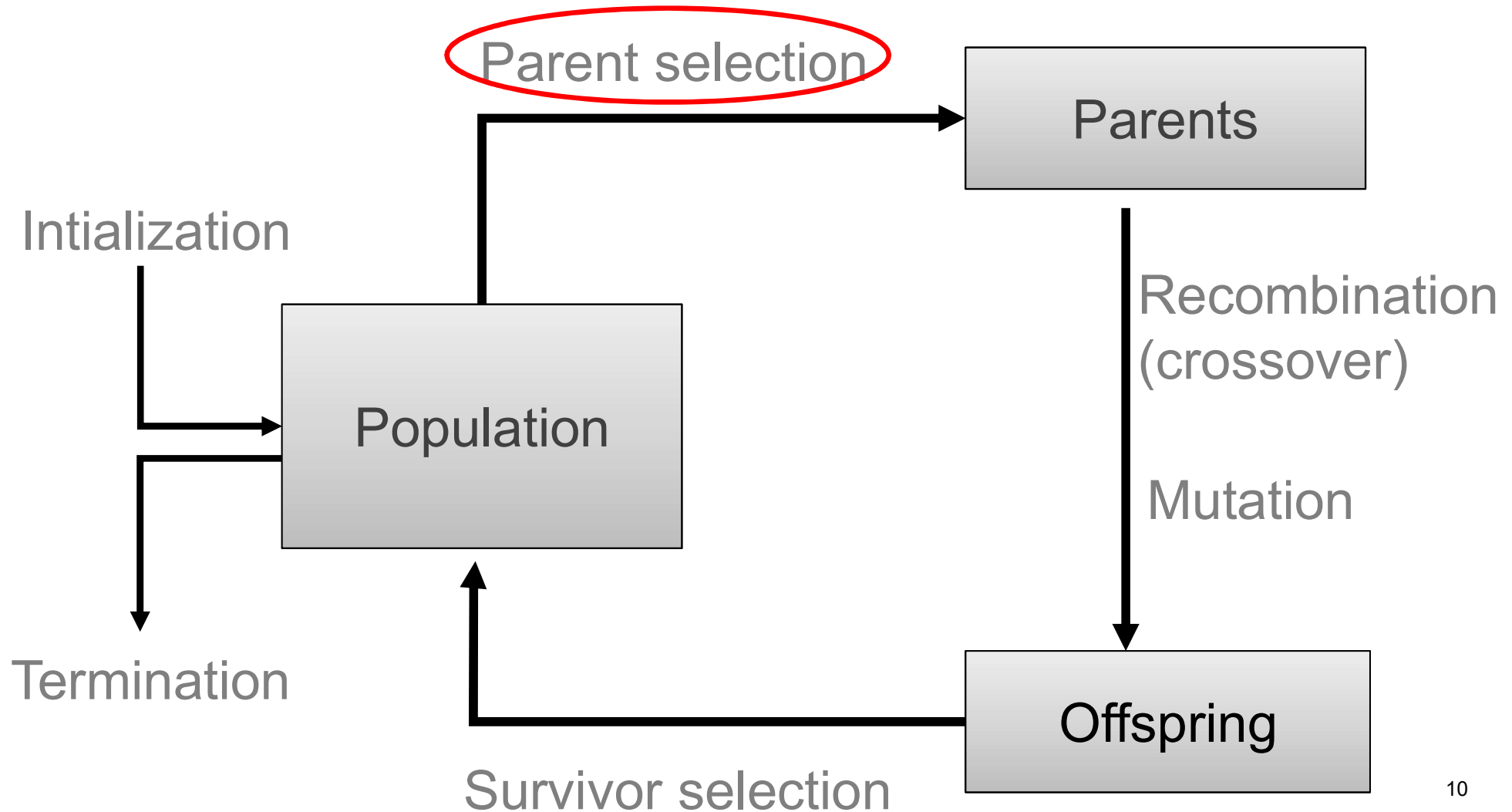


Exploitation



Scheme of an EA:

General scheme of EAs



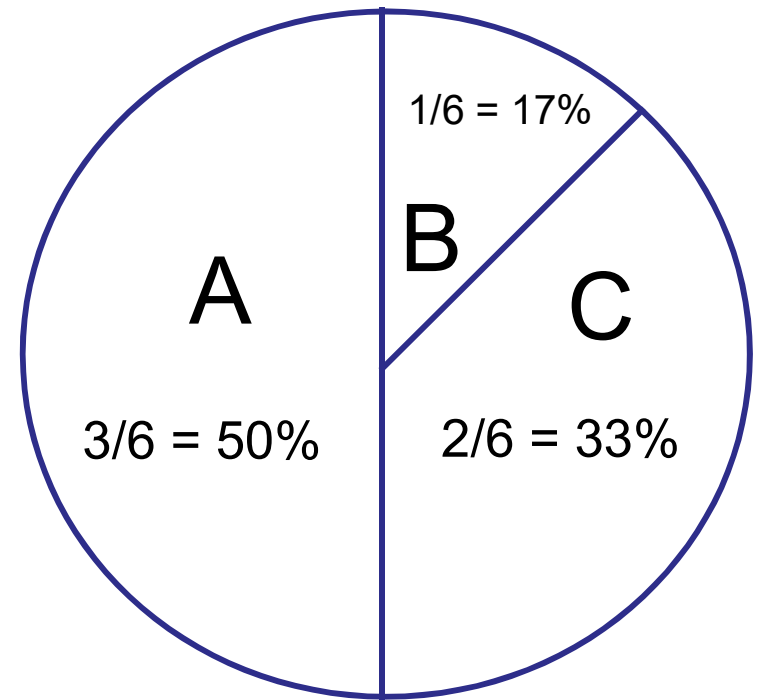
Parent Selection: Fitness-Proportionate Selection

Example: roulette wheel selection

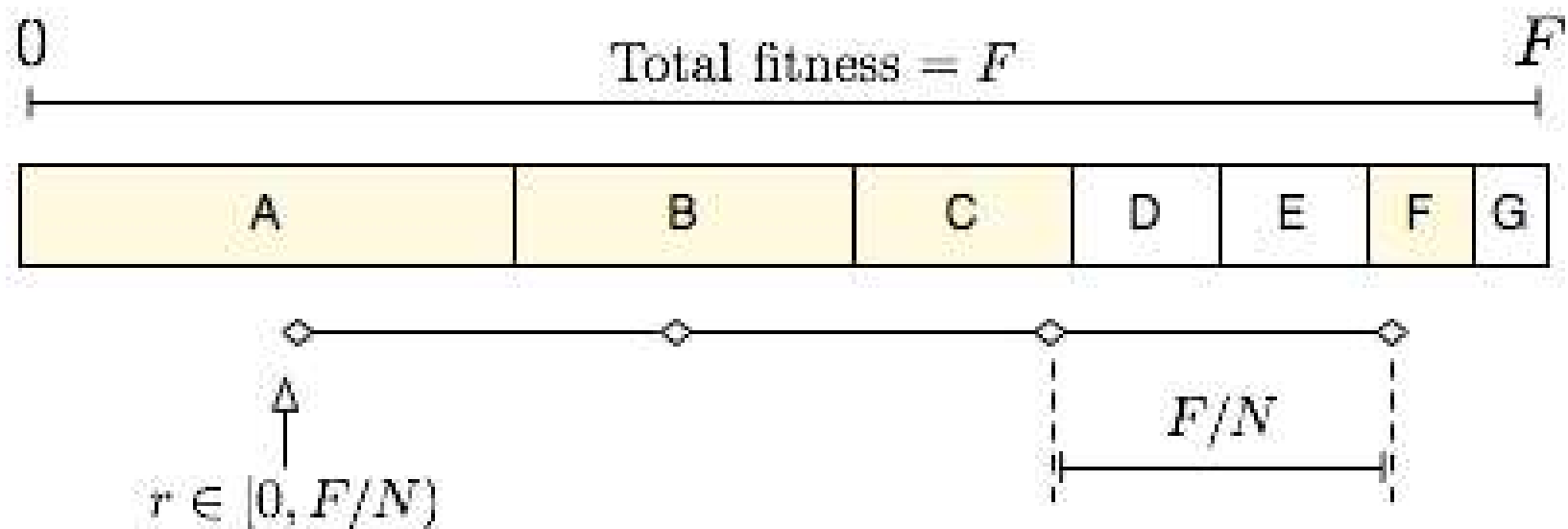
$\text{fitness}(A) = 3$

$\text{fitness}(B) = 1$

$\text{fitness}(C) = 2$



Stochastic Universal Sampling



Stochastic universal sampling (SUS)

Select multiple individuals by making **one** spin of the wheel with **a number of equally spaced arms**

Parent Selection:

Fitness-Proportionate Selection (FPS)

- Probability for individual i to be selected for mating in a population size μ with FPS is

$$P_{FPS}(i) = f_i / \sum_{j=1}^{\mu} f_j$$

- Problems include
 - One highly fit member can rapidly take over if rest of population is much less fit: **Premature Convergence**
 - At end of runs when fitnesses are similar, loss of selection pressure
- **Scaling** can fix the last problem by:
 - **Windowing:** $f'(i) = f(i) - \beta^t$

where β is worst fitness in this (last n) generations

- **Sigma Scaling:** $f'(i) = \max(f(i) - (\bar{f} - c \cdot \sigma_f), 0)$

where c is a constant, usually 2.0

Parent Selection: Rank-based Selection

- Attempt to remove problems of FPS by basing selection probabilities on **relative** rather than **absolute** fitness
- **Rank population** according to fitness and then base selection probabilities on rank (fittest has rank $\mu-1$ and worst rank 0)
- This imposes a sorting overhead on the algorithm



Parent Selection: Tournament Selection (1/3)

- All methods above rely on global population statistics
 - Could be a bottleneck esp. on parallel machines, very large population
 - Relies on presence of external fitness function which might not exist: e.g. evolving game players

Parent Selection: Tournament Selection (2/3)

Idea for a procedure using only local fitness information:

- Pick k members at random then select the best of these
- Repeat to select more individuals



Parent Selection: Tournament Selection (3/3)

- Probability of selecting i will depend on:
 - Rank of i
 - Size of sample k
 - higher k increases selection pressure
 - Whether contestants are picked with replacement
 - Picking without replacement increases selection pressure
 - Whether fittest contestant always wins (deterministic) or this happens with probability p

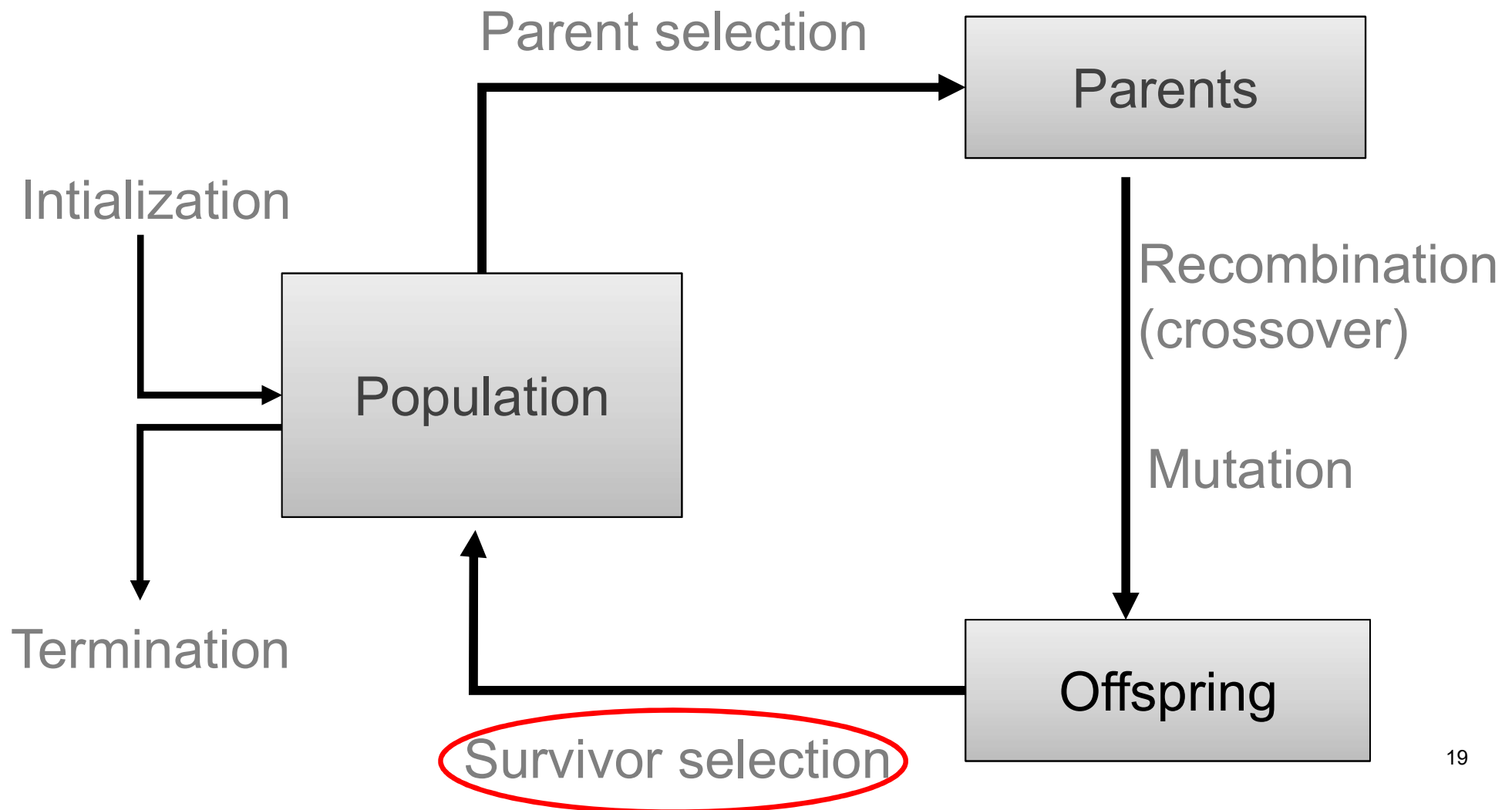
Parent Selection: Uniform

$$P_{uniform}(i) = \frac{1}{\mu}$$

- Parents are selected by uniform random distribution whenever an operator needs one/some
- Uniform parent selection is unbiased - every individual has the **same probability** to be selected

Scheme of an EA:

General scheme of EAs



Survivor Selection (Replacement)

- From a set of μ old solutions and λ offspring:
Select a set of μ individuals **forming the next generation**
- Survivor selection can be divided into two approaches:
 - **Age-Based Replacement**
 - Fitness is not taken into account
 - **Fitness-Based Replacement**
 - Usually with deterministic elements

Fitness-based replacement (1/2)

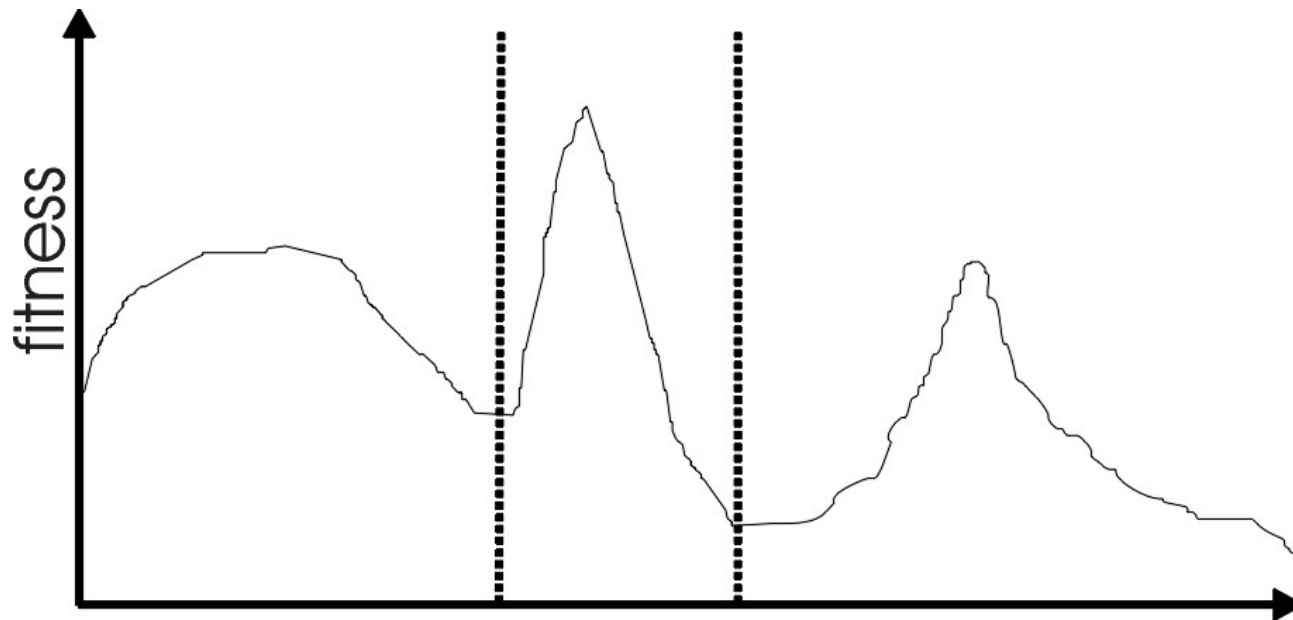
- Elitism
 - Always **keep** at least one copy of **the N fittest solution(s)** so far
 - Widely used in both population models (GGA, SSGA)
- Delete Worst
 - The worst λ individuals are replaced
- Round-robin tournament (from Evolutionary Programming)
 - Pairwise competitions in round-robin format:
 - Each individual x is **evaluated against q other** randomly chosen individuals in 1-on-1 tournaments
 - For each comparison, a "win" is assigned if x is better than its opponent
 - The μ solutions with the greatest number of wins are the winners of the tournament
 - Parameter q allows tuning selection pressure

Fitness-based replacement (2/2) (from Evolution Strategies)

- **(μ, λ) -selection** (best candidates can be lost)
 - based on the set of **children only** ($\lambda > \mu$)
 - choose the **best** μ offspring for next generation
- **$(\mu + \lambda)$ -selection** (elitist strategy)
 - based on the set of **parents and children**
 - choose the **best** μ offspring for next generation
- Often (μ, λ) -selection is preferred because it is better in leaving local optima

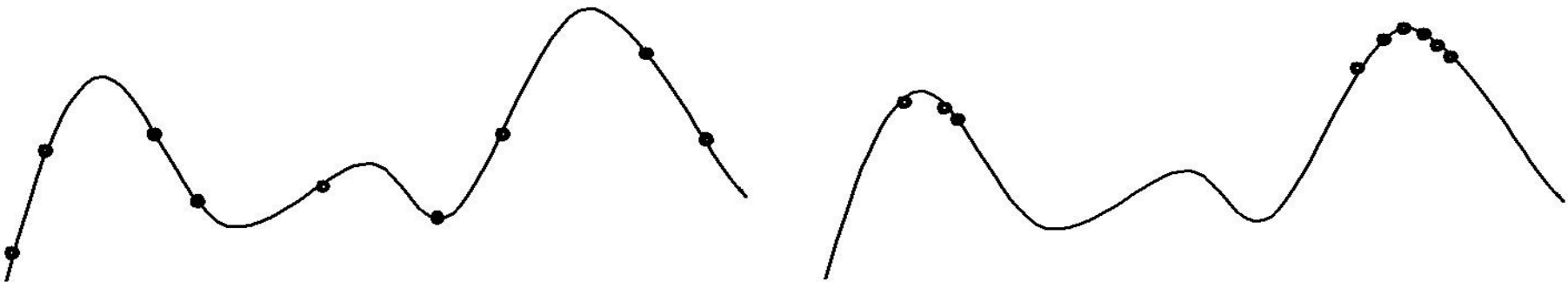
Multimodality

Most interesting problems have more than one locally optimal solution.



Multimodality

- Often might want to identify several possible peaks
- Different peaks may be different good ways to solve the problem.
- We therefore need methods to **preserve diversity** (instead of converging to one peak)



Approaches for Preserving Diversity: Introduction

- Explicit vs implicit
- Implicit approaches:
 - Impose an equivalent of geographical separation
 - Impose an equivalent of speciation
- Explicit approaches
 - Make similar individuals compete for resources (fitness)
 - Make similar individuals compete with each other for survival

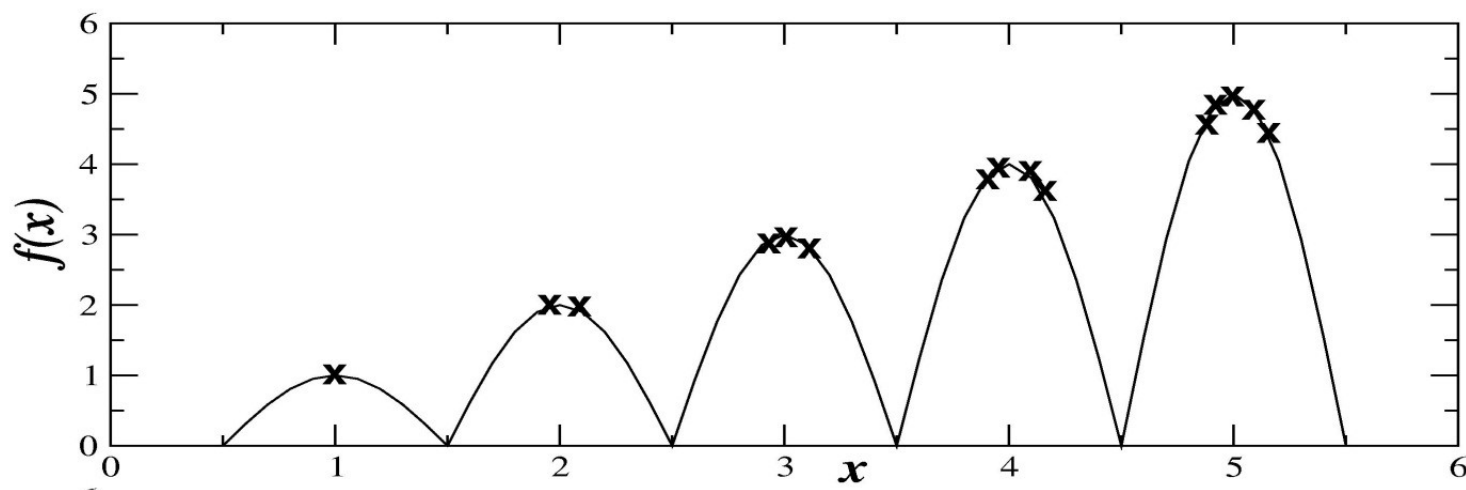
Explicit Approaches for Preserving Diversity: Fitness Sharing (1/2)

- Restricts the number of individuals within a given niche by “sharing” their fitness
- Need to set the size of the niche σ_{share} in either genotype or phenotype space
- run EA as normal but after each generation set

$$f'(i) = \frac{f(i)}{\sum_{j=1}^{\mu} sh(d(i, j))} \quad sh(d) = \begin{cases} 1 - d / \sigma & d \leq \sigma \\ 0 & otherwise \end{cases}$$

Explicit Approaches for Preserving Diversity: Fitness Sharing (2/2)

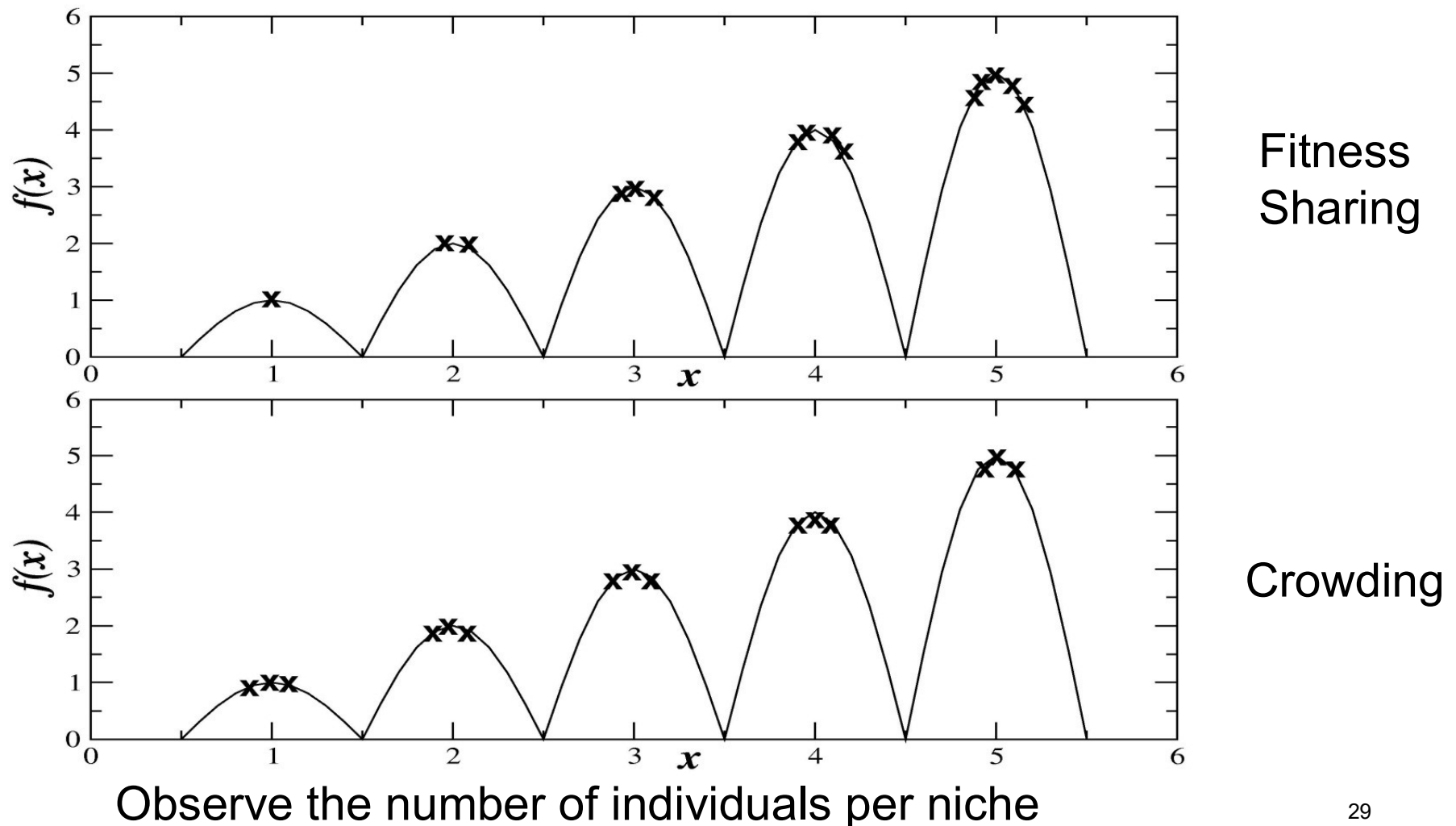
$$f'(i) = \frac{f(i)}{\sum_{j=1}^{\mu} sh(d(i, j))} \quad sh(d) = \begin{cases} 1 - d / \sigma & d \leq \sigma \\ 0 & otherwise \end{cases}$$



Explicit Approaches for Preserving Diversity: Crowding

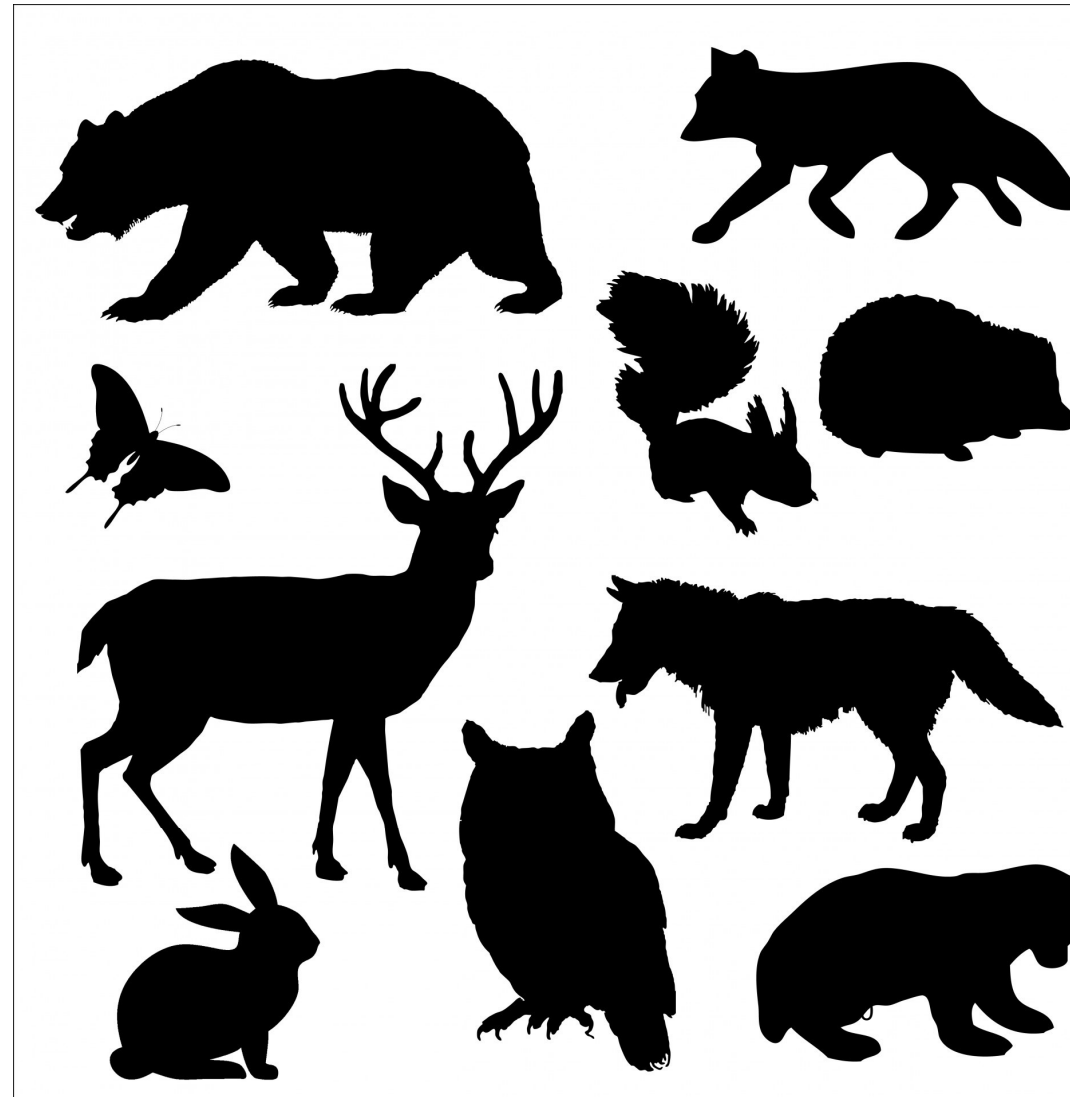
- Idea: New individuals replace *similar* individuals
- Randomly shuffle and pair parents, produce 2 offspring
- Each offspring competes with their **nearest** parent for survival (using a distance measure)
- Result: Even distribution among niches.

Explicit Approaches for Preserving Diversity: Crowding vs Fitness sharing



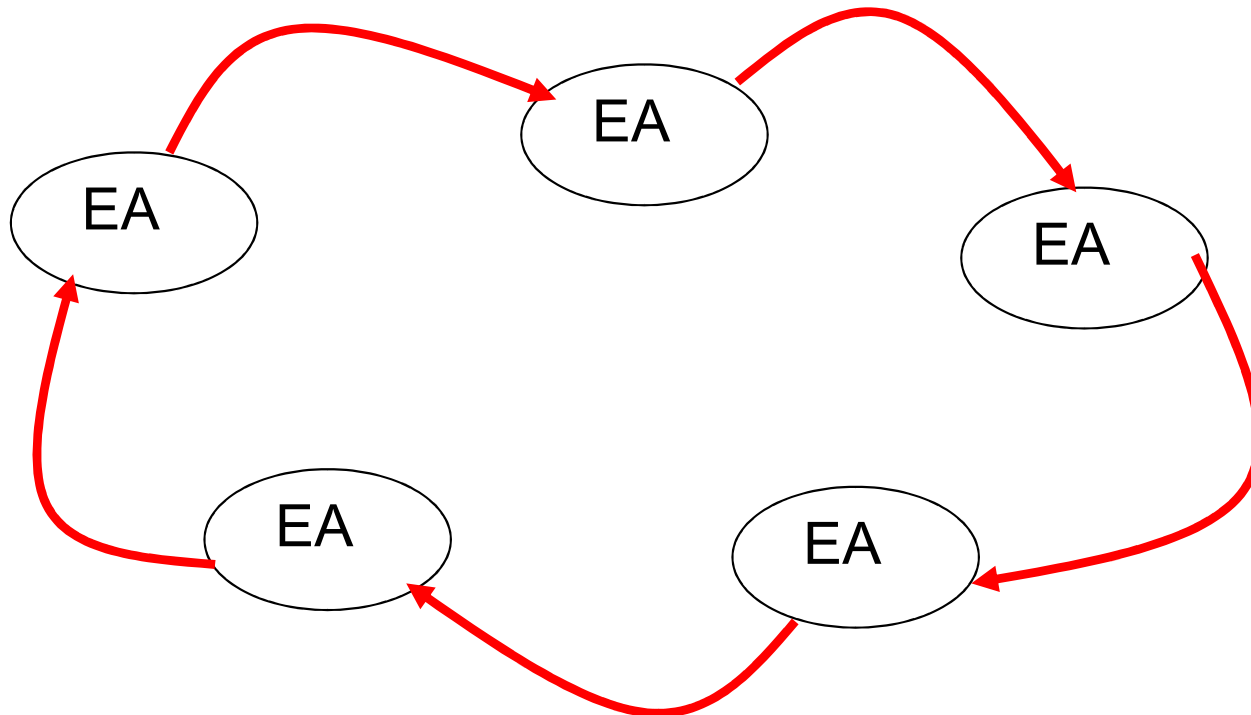
Implicit Approaches for Preserving Diversity: Automatic Speciation

- Either only mate with genotypically / phenotypically similar members or
- Add species-tags to genotype
 - initially randomly set
 - when selecting partner for recombination, only pick members with a good match



Implicit Approaches for Preserving Diversity: Geographical Separation

- “Island” Model Parallel EA
- Periodic migration of individual solutions between populations



Implicit Approaches for Preserving Diversity: “Island” Model Parallel EAs

- Run multiple populations in parallel
- After a (usually fixed) number of generations (an ***Epoch***), exchange individuals with neighbours
- Repeat until ending criteria met
- Partially inspired by parallel/clustered systems

Island Model: Parameters

- How often to exchange individuals ?
 - too quick and all sub-populations converge to same solution
 - too slow and waste time
 - can do it adaptively (stop each pop when no improvement for (say) 25 generations)
- Operators can differ between the sub-populations

Chapter 6: Popular Evolutionary Algorithm Variants

Historical EA variants:

- Genetic Algorithms
- Evolution Strategies
- Evolutionary Programming
- Genetic Programming

Algorithm	Chromosome Representation	Crossover	Mutation
Genetic Algorithm (GA)	Array	X	X
Genetic Programming (GP)	Tree	X	X
Evolution Strategies (ES)	Array	(X)	X
Evolutionary Programming (EP)	No constraints	-	X

Genetic Algorithms:

Overview Simple GA (1/2)

- Developed: USA in the 1960's
- Early names: Holland, DeJong, Goldberg
- Typically applied to:
 - discrete function optimization
 - benchmark for comparison with other algorithms
 - straightforward problems with binary representation
- Features:
 - not too fast
 - missing new variants (elitism, sus)
 - often modelled by theorists

Genetic Algorithms:

Overview Simple GA (2/2)

- Holland's original GA is now known as the simple genetic algorithm (SGA)
- Other GAs use different:
 - Representations
 - Mutations
 - Crossovers
 - Selection mechanisms

Genetic Algorithms:

SGA reproduction cycle

- **Select parents** for the mating pool
(size of mating pool = population size)
- Shuffle the mating pool
- **Apply crossover** for each consecutive pair with probability p_c , otherwise copy parents
- **Apply mutation** for each offspring (bit-flip with probability p_m independently for each bit)
- **Replace the whole population** with the resulting offspring

Genetic Algorithms:

An example after Goldberg '89

- Simple problem: $\max x^2$ over $\{0,1,\dots,31\}$
- GA approach:
 - Representation: binary code, e.g., $01101 \leftrightarrow 13$
 - Population size: 4
 - 1-point x-over, bitwise mutation
 - Roulette wheel selection
 - Random initialisation
- We show one generational cycle done by hand

X² example: Selection

String no.	Initial population	x Value	Fitness $f(x) = x^2$	$Prob_i$	Expected count	Actual count
1	0 1 1 0 1	13	169	0.14	0.58	1
2	1 1 0 0 0	24	576	0.49	1.97	2
3	0 1 0 0 0	8	64	0.06	0.22	0
4	1 0 0 1 1	19	361	0.31	1.23	1
Sum			1170	1.00	4.00	4
Average			293	0.25	1.00	1
Max			576	0.49	1.97	2

X² example: Crossover

String no.	Mating pool	Crossover point	Offspring after xover	x Value	Fitness $f(x) = x^2$
1	0 1 1 0 1	4	0 1 1 0 0	12	144
2	1 1 0 0 0	4	1 1 0 0 1	25	625
2	1 1 0 0 0	2	1 1 0 1 1	27	729
4	1 0 0 1 1	2	1 0 0 0 0	16	256
Sum					1754
Average					439
Max					729

X² example: Mutation

String no.	Offspring after xover	Offspring after mutation	x Value	Fitness $f(x) = x^2$
1	0 1 1 0 0	1 1 1 0 0	26	676
2	1 1 0 0 1	1 1 0 0 1	25	625
2	1 1 0 1 1	1 1 0 1 1	27	729
4	1 0 0 0 0	1 0 1 0 0	18	324
Sum				2354
Average				588.5
Max				729

Genetic Algorithms:

The simple GA

- Has been subject of many (early) studies
 - still often used as benchmark for novel GAs
- Shows many shortcomings, e.g.,
 - Representation is too restrictive
 - Mutation & crossover operators only applicable for bit-string & integer representations
 - Selection mechanism sensitive for converging populations with close fitness values
 - Generational population model can be improved with explicit survivor selection

Genetic Algorithms:

Simple GA (SGA) summary

Representation	Bit-strings
Recombination	1-Point crossover
Mutation	Bit flip
Parent selection	Fitness proportional – implemented by Roulette Wheel
Survivor selection	Generational

Evolution Strategies:

Quick overview

- Developed: Germany in the 1960's by Rechenberg and Schwefel
- Typically applied to numerical optimisation
- Attributed features:
 - fast
 - good optimizer for real-valued optimisation
 - relatively much theory
- Special:
 - self-adaptation of (mutation) parameters standard

Evolution Strategies: Example (1+1) ES

- Task: minimise $f : \mathbb{R}^n \rightarrow \mathbb{R}$
- Algorithm: “two-membered ES” using
 - Vectors from \mathbb{R}^n directly as chromosomes
 - Population size 1
 - Only mutation creating one child
 - Greedy selection

Evolution Strategies: Representation

- Chromosomes consist of two parts:
 - Object variables: x_1, \dots, x_n
 - Strategy parameters (mutation rate, etc):
 p_1, \dots, p_m
- Full size: $\langle x_1, \dots, x_n, p_1, \dots, p_n \rangle$

Evolution Strategies:

Parent selection

- Parents are selected by **uniform random distribution** whenever an operator needs one/some
- Thus: ES parent selection is unbiased - every individual has the **same probability** to be selected

$$P_{uniform}(i) = \frac{1}{\mu}$$

Evolution Strategies: Recombination

- Two parents create one child
- Acts per variable / position by either
 - Intermediary crossover, or
 - Discrete crossover
- From two or more parents by either:
 - Local recombination: Two parents make a child
 - Global recombination: Selecting two parents randomly for each gene

Evolution Strategies:

Names of recombinations

	Two fixed parents	Two parents selected for each i
$z_i = (x_i + y_i)/2$	Local intermediary	Global intermediary
z_i is x_i or y_i chosen randomly	Local discrete	Global discrete

Evolution Strategies:

ES summary

Representation	Real-valued vectors
Recombination	Discrete or intermediary
Mutation	Gaussian perturbation
Parent selection	Uniform random
Survivor selection	(μ, λ) or $(\mu + \lambda)$

Evolutionary Programming:

Quick overview

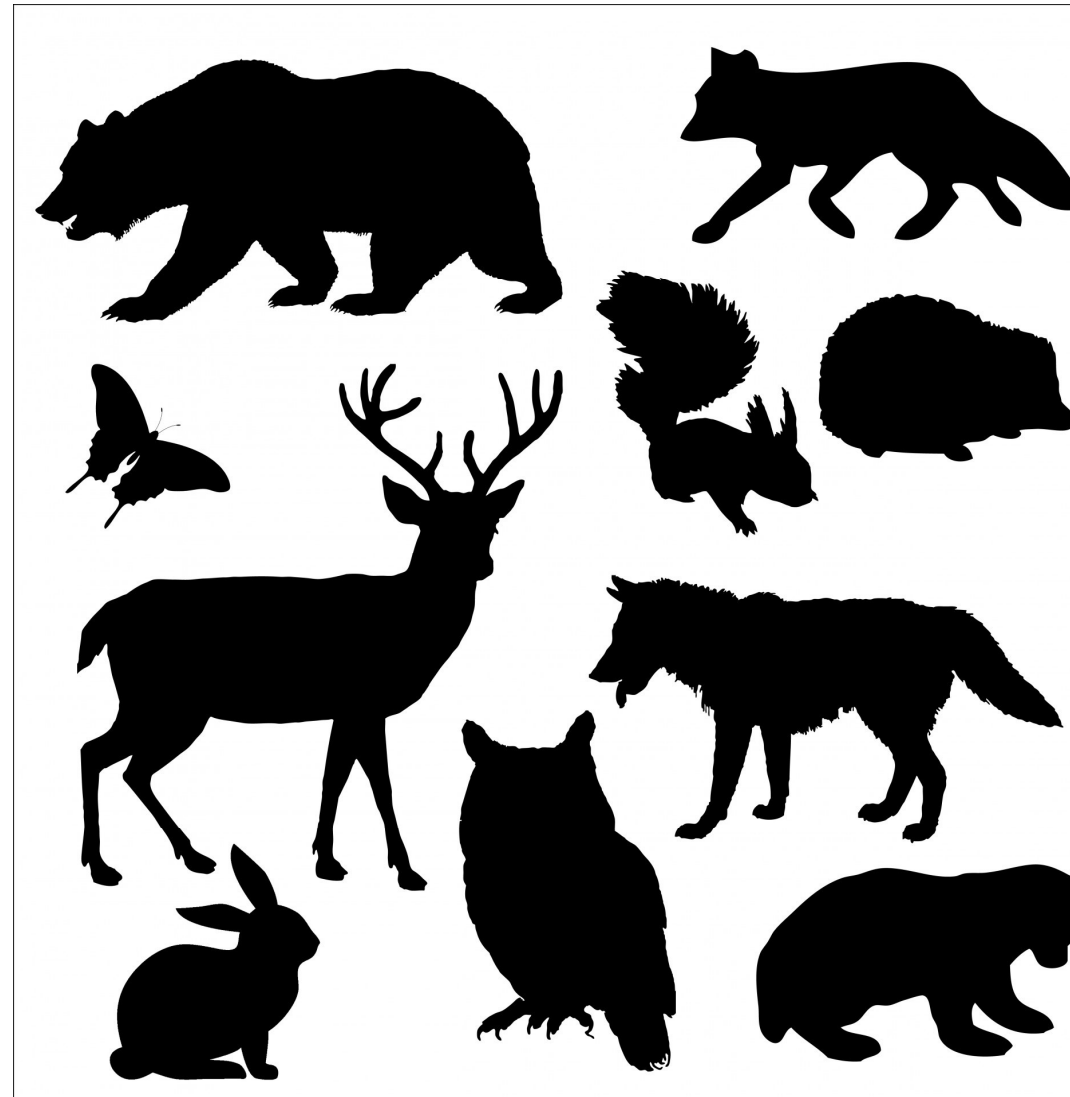
- Developed: USA in the 1960's by Fogel et al.
- Typically applied to:
 - traditional EP: prediction by finite state machines
 - contemporary EP: (numerical) optimization
- Attributed features:
 - very open framework: any representation and mutation op's OK
 - Contemporary EP has almost merged with ES
- Special:
 - **no recombination**
 - self-adaptation of parameters standard (contemporary EP)

Evolutionary Programming: Representation

- For continuous parameter optimisation
- Chromosomes consist of two parts:
 - Object variables: x_1, \dots, x_n
 - Mutation step sizes: $\sigma_1, \dots, \sigma_n$
- Full size: $\langle x_1, \dots, x_n, \sigma_1, \dots, \sigma_n \rangle$

Evolutionary Programming: Recombination

- None
- Rationale: one point in the search space stands for a species, not for an individual and there can be no crossover between species



Evolutionary Programming: Selection

- Each individual creates one child by mutation
 - Deterministic
 - Not biased by fitness
- Parents and offspring compete for survival in round-robin tournaments.

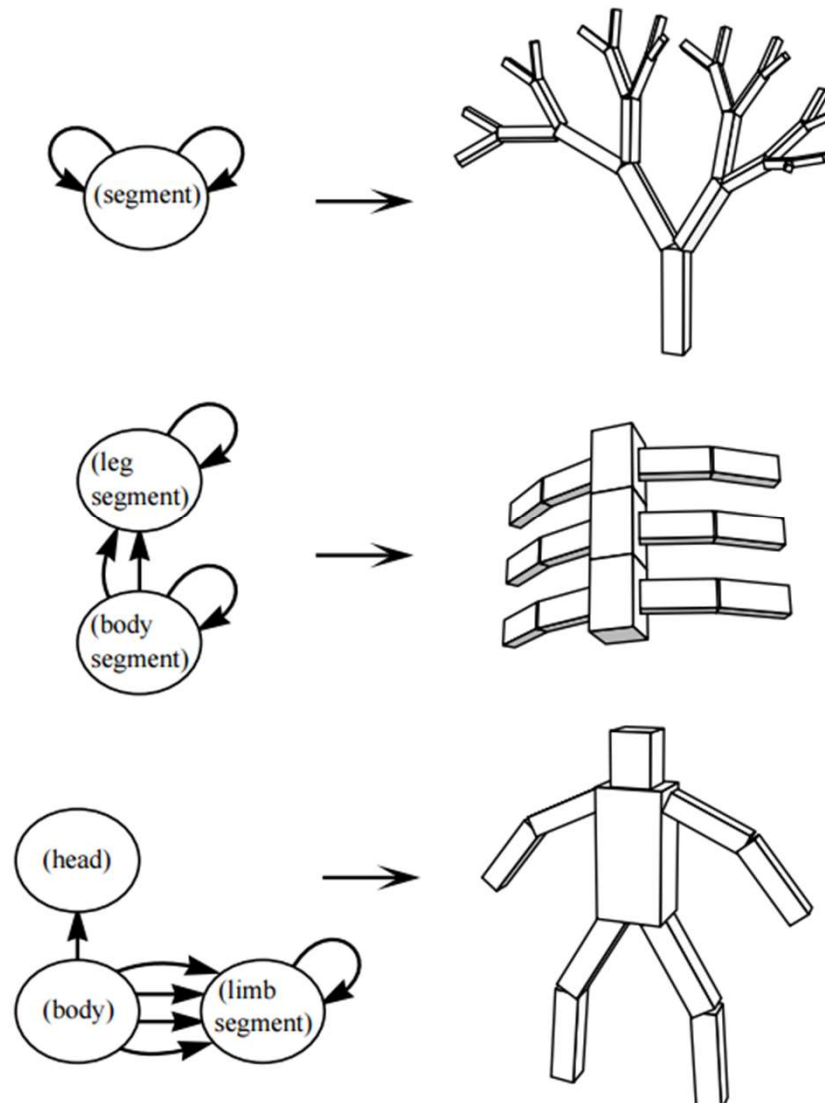
Evolutionary Programming: Summary

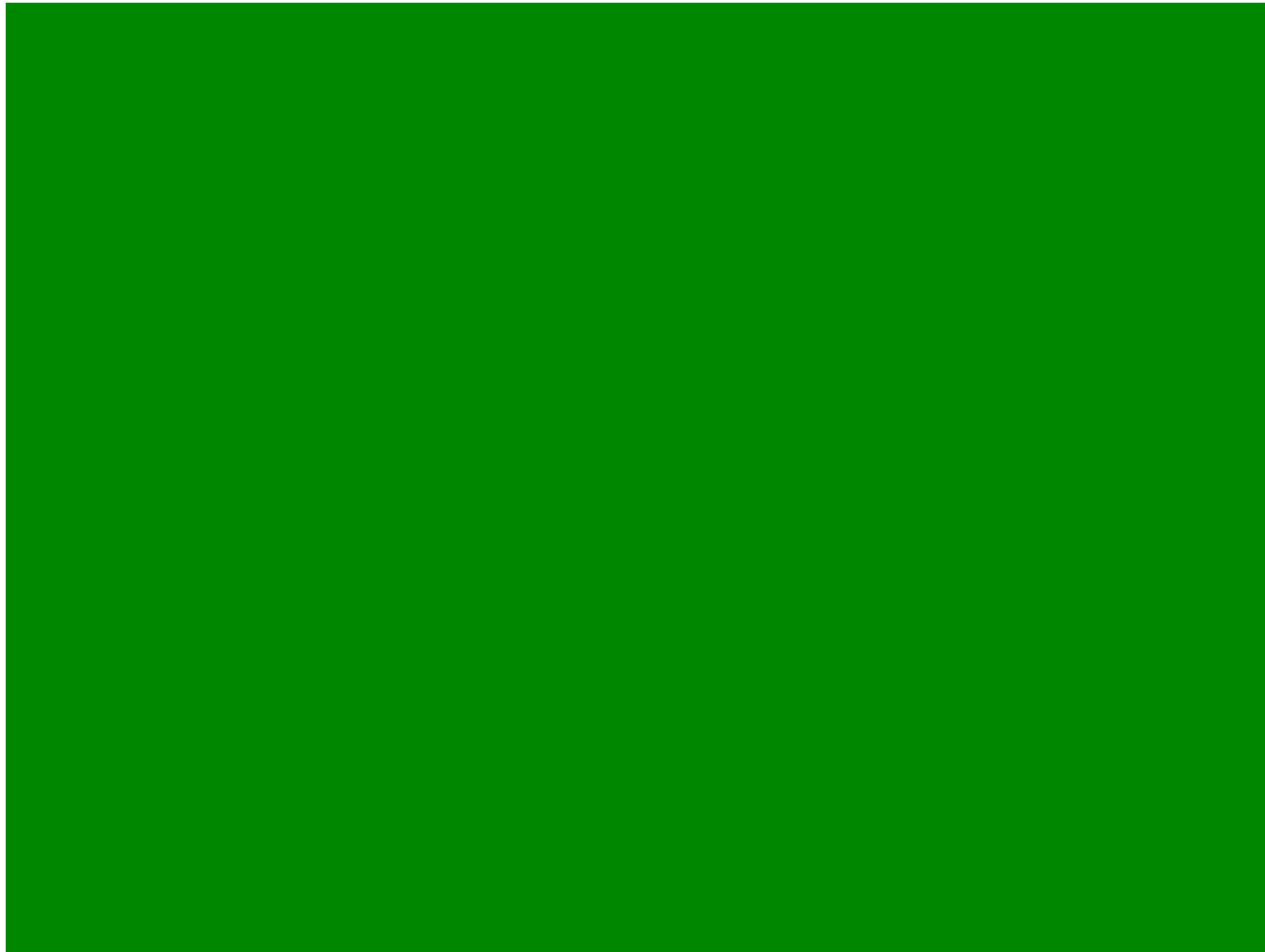
Representation	Real-valued vectors
Recombination	None
Mutation	Gaussian perturbation
Parent selection	Deterministic (each parent one offspring)
Survivor selection	Probabilistic ($\mu+\lambda$)

Virtual Creatures (Karl Sims, 1994)

Genotype: directed graph.

Phenotype: hierarchy of 3D parts.





Genetic Programming:

Quick overview

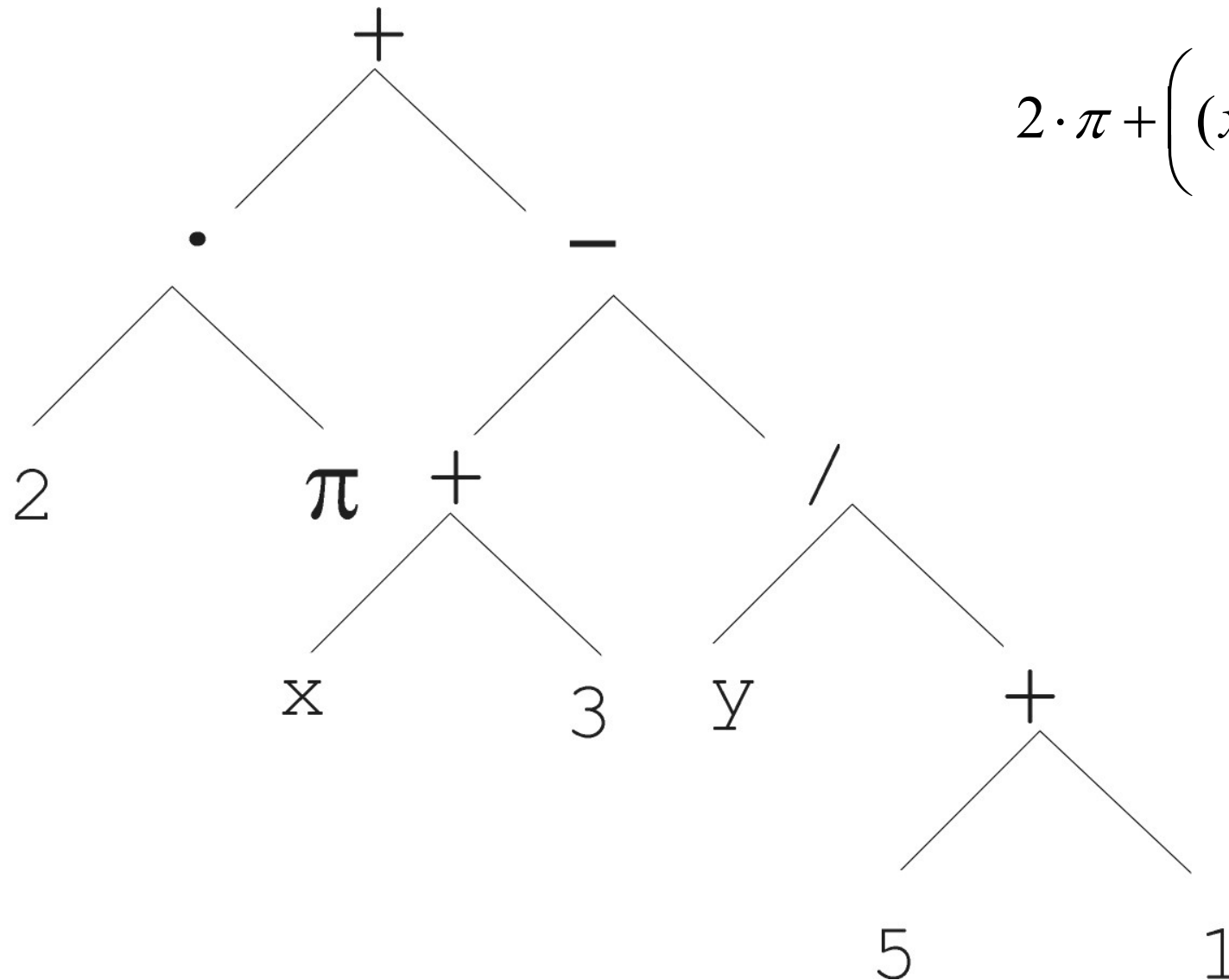
- Developed: USA in the 1990's by Koza
- Typically applied to:
 - machine learning tasks (prediction, classification...)
- Attributed features:
 - “automatic evolution of computer programs”
 - needs huge populations (thousands)
 - slow
- Special:
 - non-linear chromosomes: trees
 - mutation possible but not necessary

Tree Representation

- Trees are a universal form, e.g. consider
- Arithmetic formula: $2 \cdot \pi + \left((x + 3) - \frac{y}{5 + 1} \right)$
- Logical formula: $(x \wedge \text{true}) \rightarrow ((x \vee y) \vee (z \leftrightarrow (x \wedge y)))$
- Program:

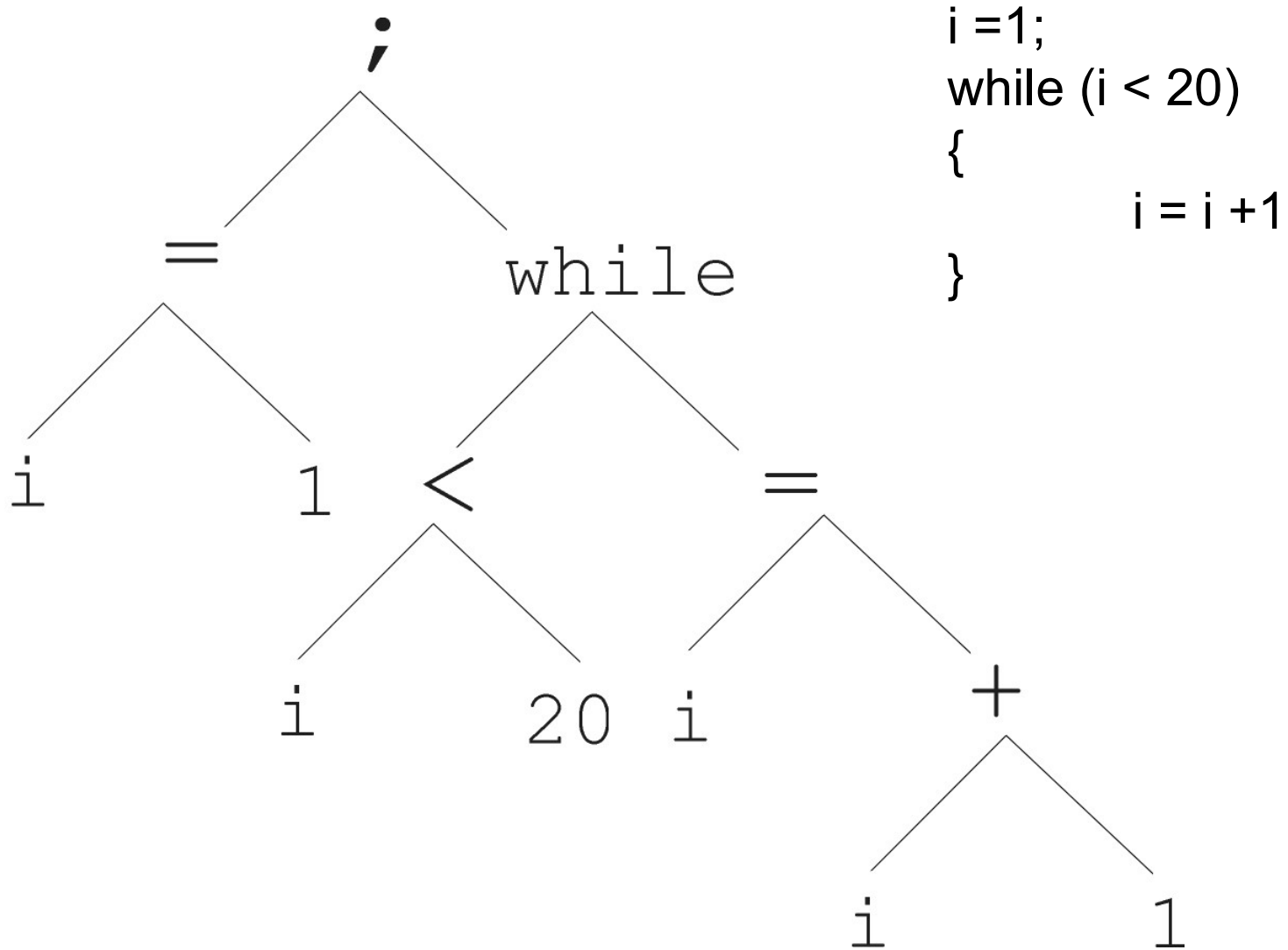
```
i = 1;
while (i < 20)
{
    i = i + 1
}
```

Tree Representation



$$2 \cdot \pi + \left((x + 3) - \frac{y}{5 + 1} \right)$$

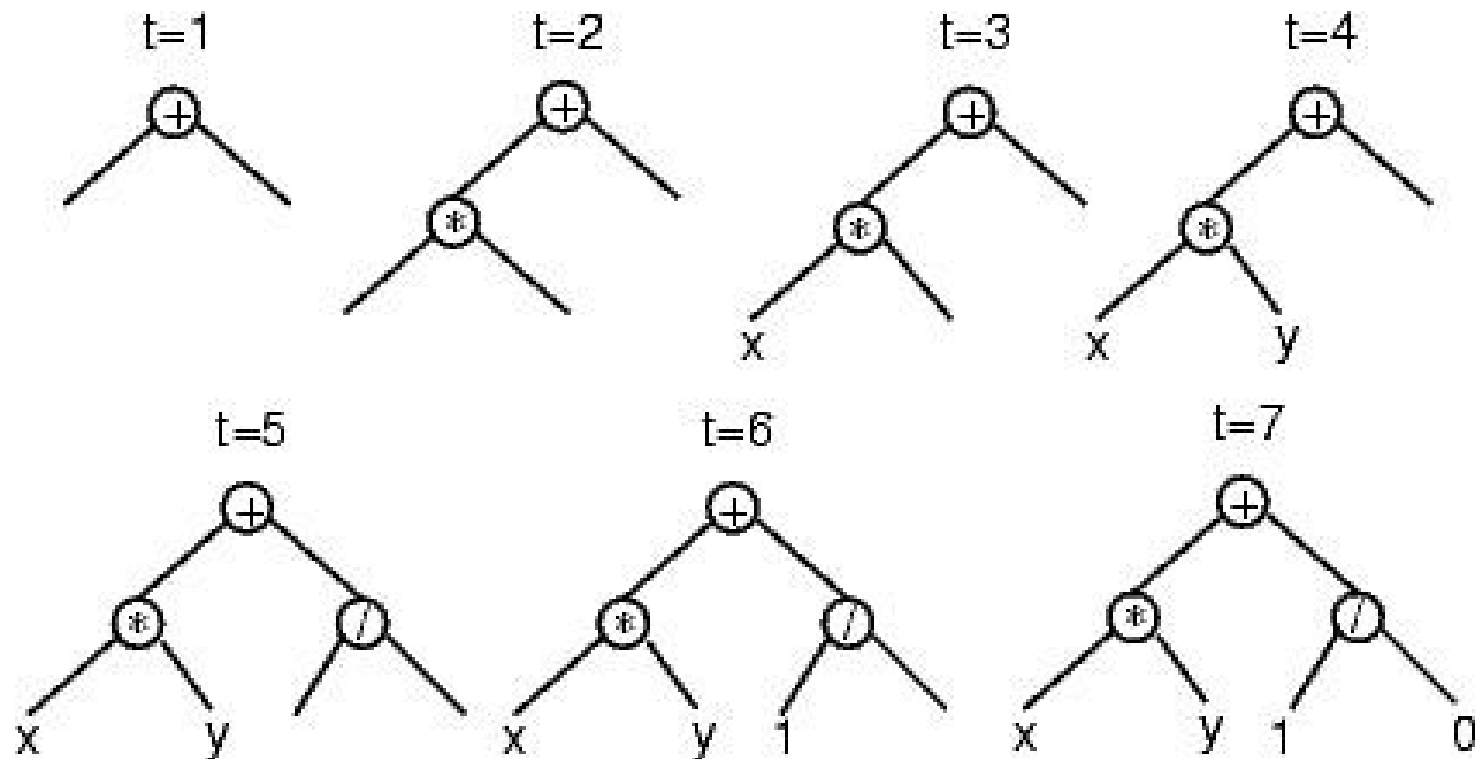
Tree Representation



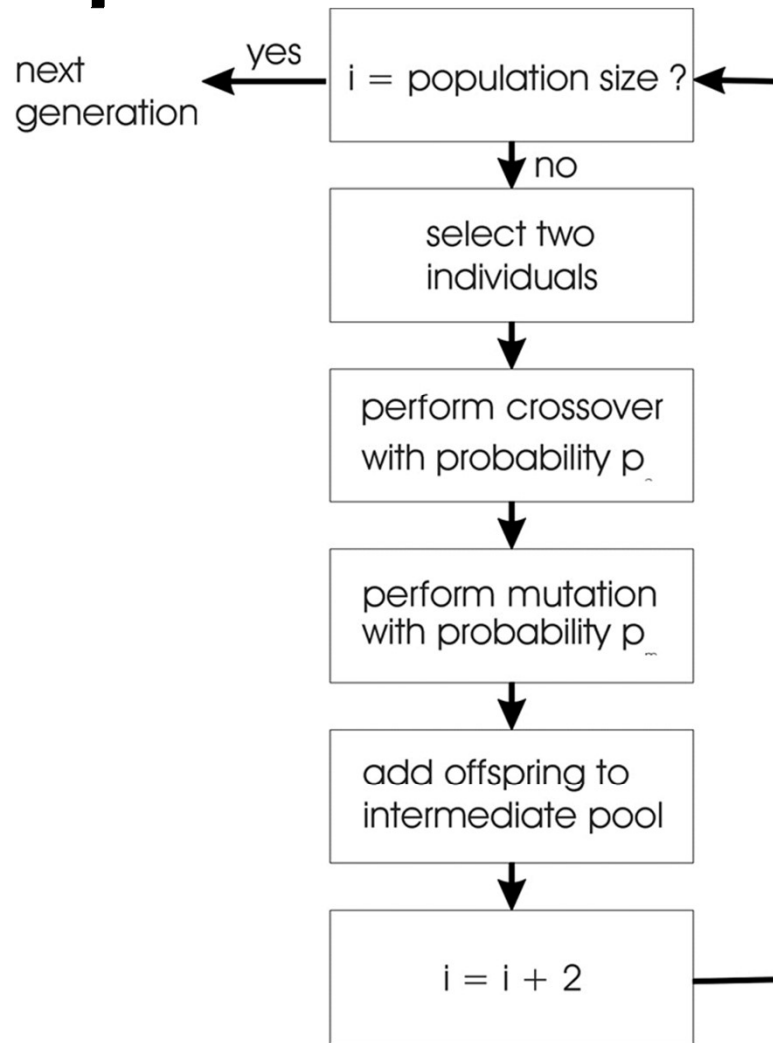
Tree Representation

- In GA, ES, EP chromosomes are linear structures (bit strings, integer string, real-valued vectors, permutations)
- Tree shaped chromosomes are non-linear structures
- In GA, ES, EP the size of the chromosomes is fixed
- Trees in GP (Genetic Programming) may vary in depth and width

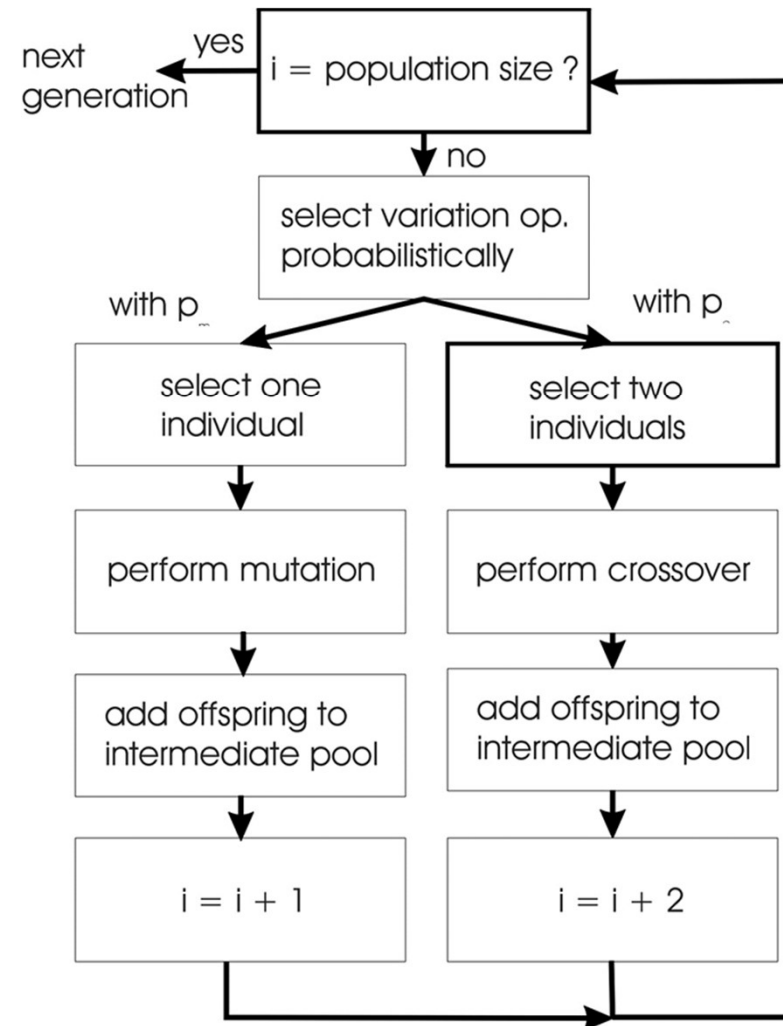
Example of how to initialize trees: Full initialisation to depth 2



Genetic Programming: Variation Operators



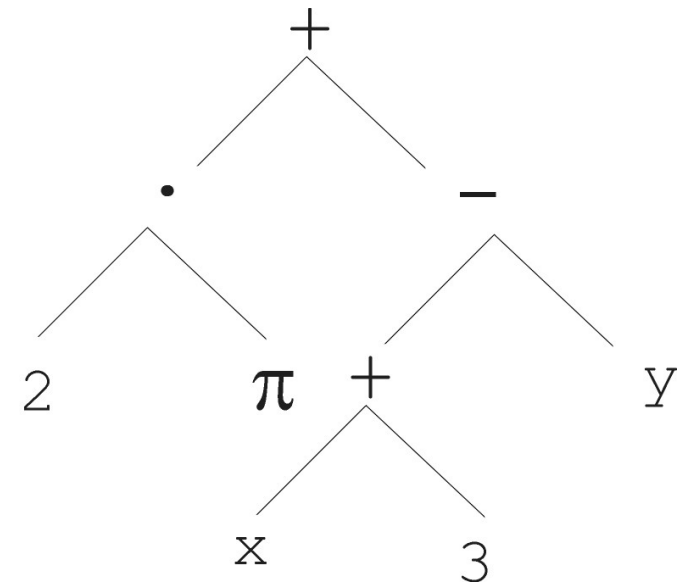
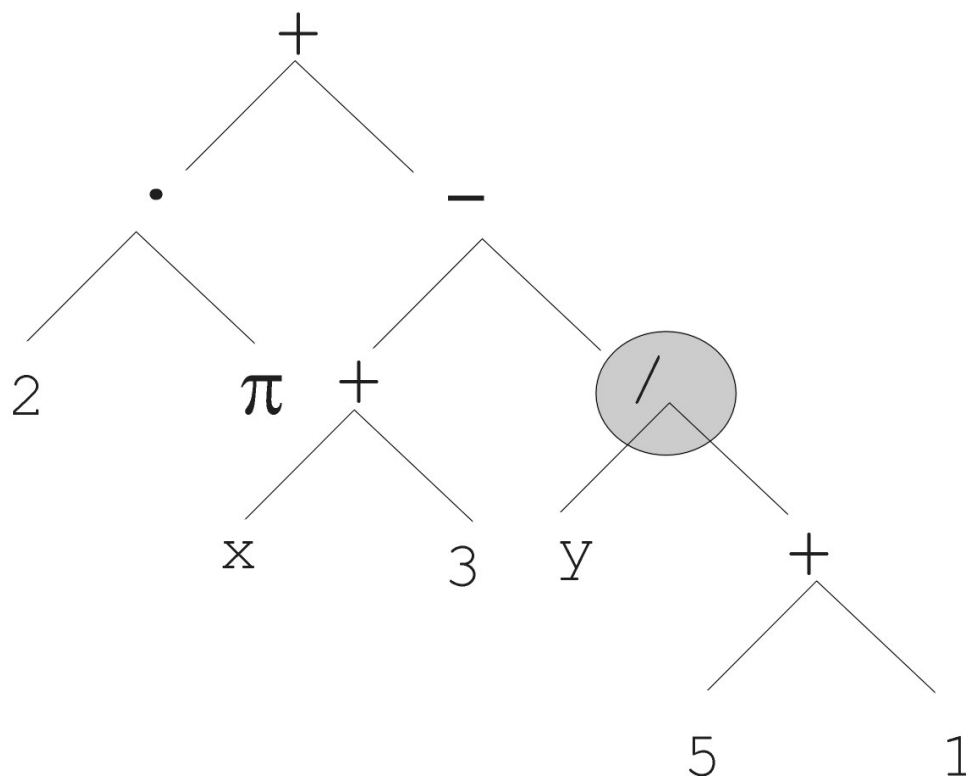
GA flowchart



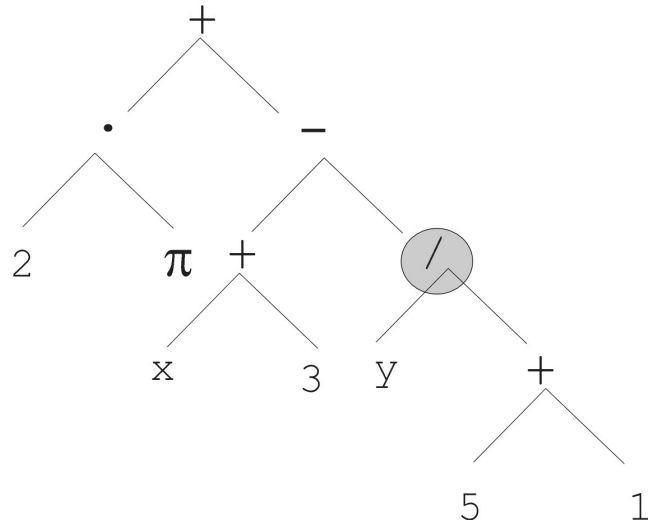
GP flowchart

Genetic Programming: Mutation

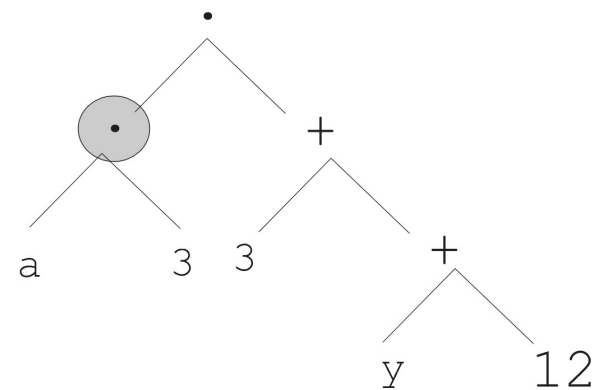
- Most common mutation: replace randomly chosen subtree by randomly generated tree



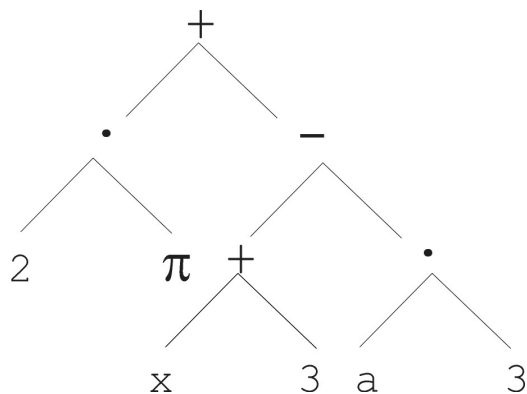
Genetic Programming: Recombination



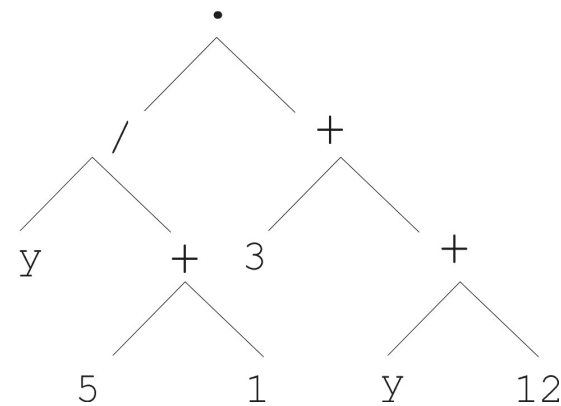
Parent 1



Parent 2



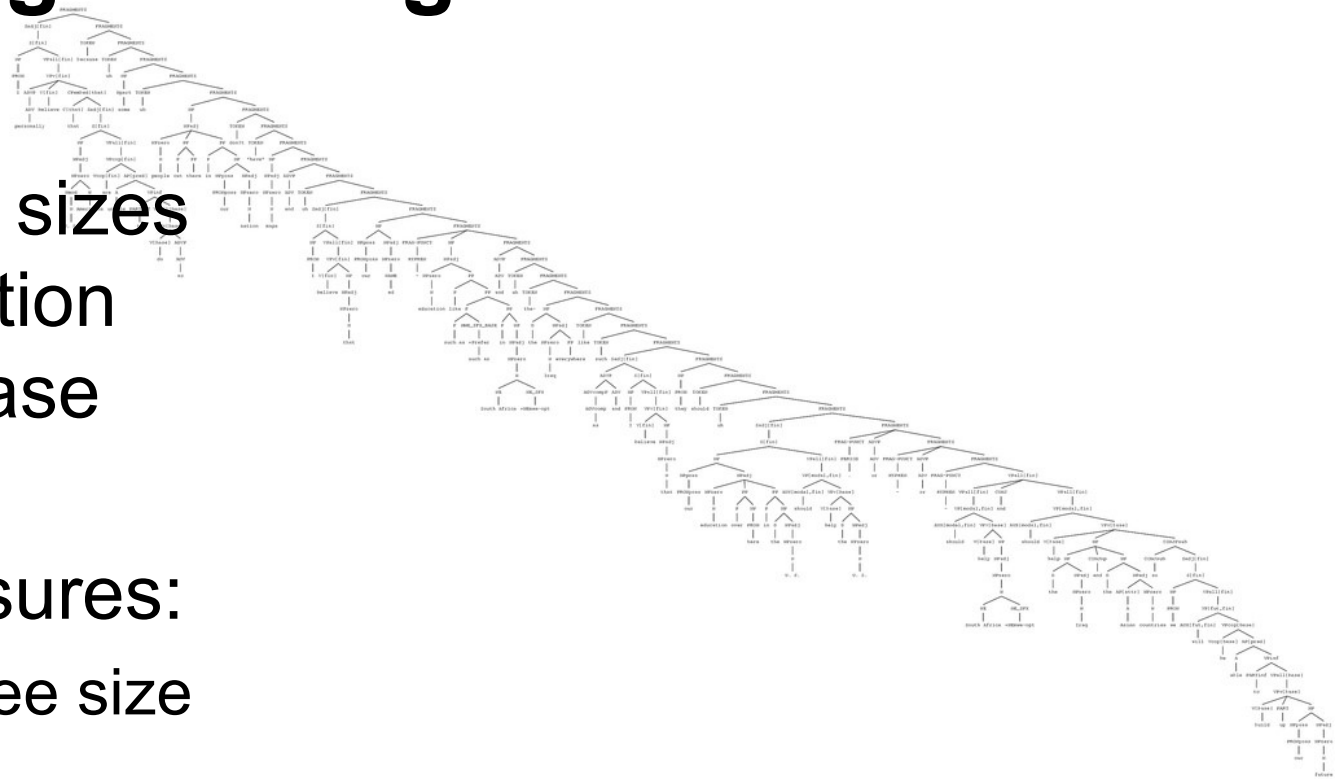
Child 1



Child 2

Genetic Programming: Bloat

- Average tree sizes in the population tend to increase over time
- Countermeasures:
 - Maximum tree size
 - Parsimony pressure: penalty for being oversized



Genetic Programming: Summary

Representation	Tree structures
Recombination	Exchange of subtrees
Mutation	Random change in trees
Parent selection	Fitness proportional
Survivor selection	Generational replacement

Summary: The standard EA variants

Name	Representation	Crossover	Mutation	Parent selection	Survivor selection	Specialty
Genetic Algorithm	Usually fixed-length vector	Any or none	Any	Any	Any	None
Evolution Strategies	Real-valued vector	Discrete or intermediate recombination	Gaussian	Random draw	Best N	Strategy parameters
Evolutionary Programming	Real-valued vector	None	Gaussian	One child each	Tournament	Strategy parameters
Genetic Programming	Tree	Swap sub-tree	Replace sub-tree	Usually fitness proportional	Generational replacement	None