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INF3490 - Biologically inspired computing

Lecture 3: Eiben and Smith, chapter 5-6

Evolutionary Algorithms - Population management and popular algorithms

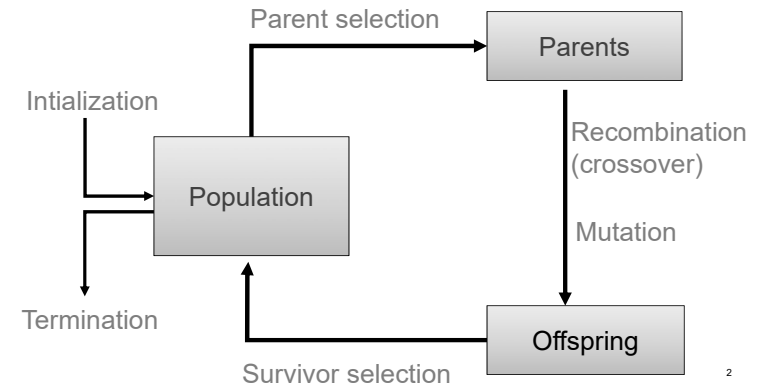


Kai Olav Ellefsen



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Repetition: General scheme of EAs



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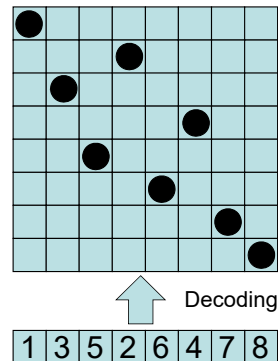
Repetition: Genotype & Phenotype

Phenotype:

A solution representation
we can **evaluate**

Genotype:

A solution representation
applicable to **variation**

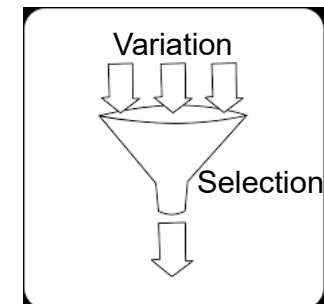


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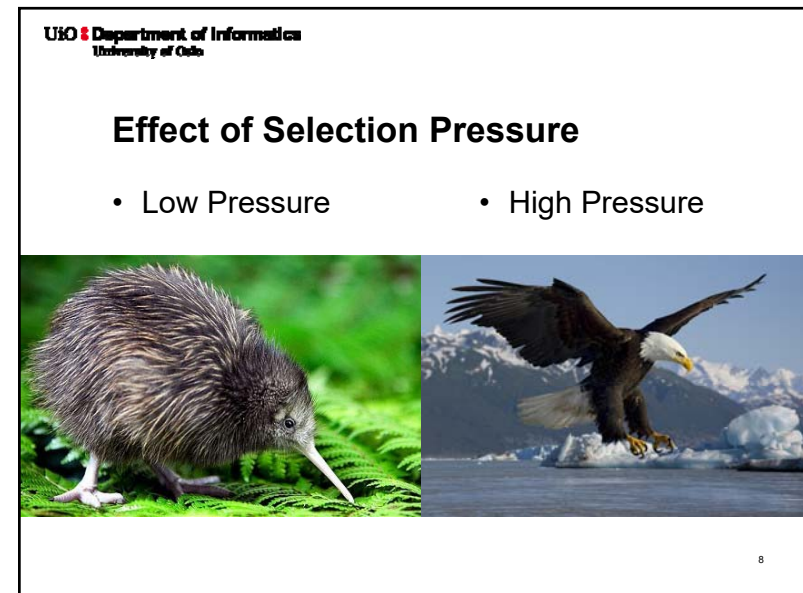
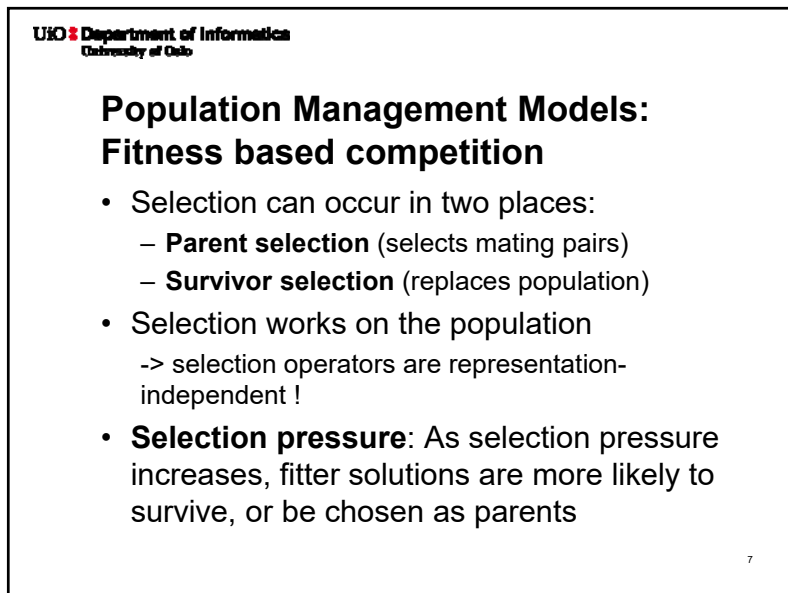
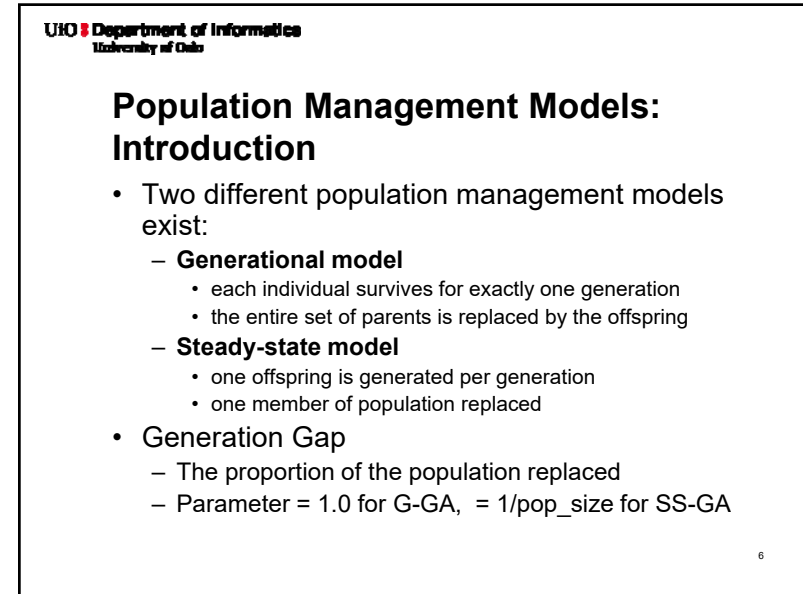
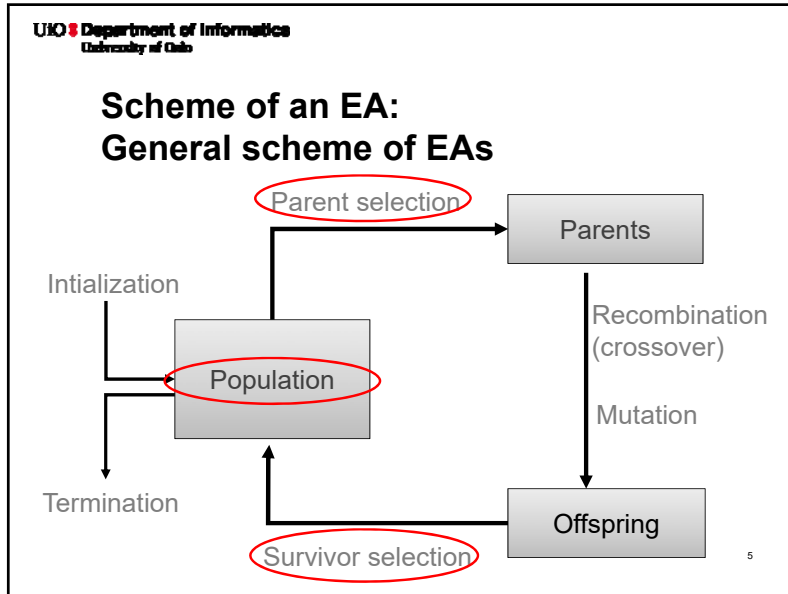
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Chapter 5: Fitness, Selection and Population Management

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



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Why Not Always High Selection Pressure?

Exploration Exploitation

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Scheme of an EA: General scheme of EAs

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Parent Selection: Fitness-Proportionate Selection

Example: roulette wheel selection

fitness(A) = 3
fitness(B) = 1
fitness(C) = 2

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Stochastic Universal Sampling

Stochastic universal sampling (SUS)
Select multiple individuals by making **one** spin of the wheel with a **number of equally spaced arms**

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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

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Parent Selection: Rank-based Selection

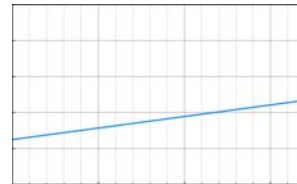
- Attempt to remove problems of FPS by basing selection probabilities on **relative** rather than **absolute** fitness
- Rank population** according to fitness and then base selection probabilities on rank (fittest has rank $\mu-1$ and worst rank 0)
- This imposes a sorting overhead on the algorithm



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Rank-based Selection: Linear Ranking

$$P_{lin-rank}(i) = \frac{(2-s)}{\mu} + \frac{2i(s-1)}{\mu(\mu-1)}$$

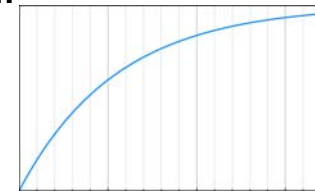


- Parameterised by factor s : $1 < s \leq 2$
 - Tunes selection pressure
- Simple 3 member example

Individual	Fitness	Rank	P_{selFP}	$P_{selLR} (s=2)$	$P_{selLR} (s=1.5)$
A	1	0	0.1	0	0.167
B	4	1	0.4	0.33	0.33
C	5	2	0.5	0.67	0.5
Sum	10		1.0	1.0	1.0

Rank-based selection: Exponential Ranking

$$P_{exp-rank}(i) = \frac{1 - e^{-i}}{c}$$



- Linear Ranking is limited in selection pressure
- Exponential Ranking can allocate more than 2 copies to fittest individual
- Normalise constant factor c according to population size

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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

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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



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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

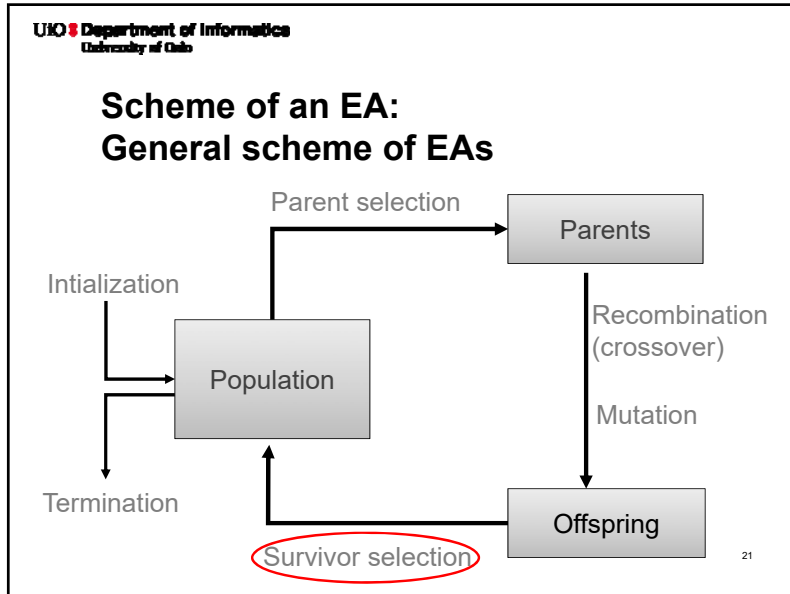
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Parent Selection: Uniform

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

- 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

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Survivor Selection (Replacement)

- From a set of μ parents 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
 - In SS-GA can implement as “delete-random” (not recommended) or as first-in-first-out (a.k.a. delete-oldest)
 - Fitness-Based Replacement**

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Fitness-based replacement (1/2)

- Elitism
 - Always **keep** at least one copy of **the fittest solution** so far
 - Widely used in both population models (GGA, SSGA)
- Delete Worst
 - The worst λ individuals are replaced
- Round-robin tournament (from EP)
 - 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
 - Typically $q = 10$

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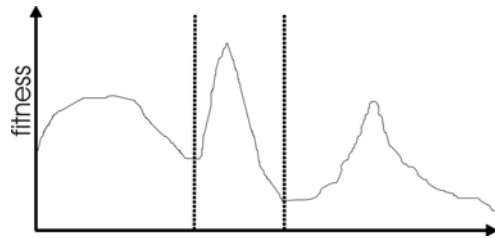
Fitness-based replacement (2/2) (from ES)

- (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

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Multimodality

Most interesting problems have more than one locally optimal solution.



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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)



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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

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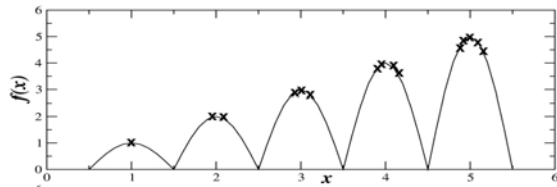
Explicit Approaches for Preserving Diversity: Fitness Sharing (1/2)

- 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

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

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

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

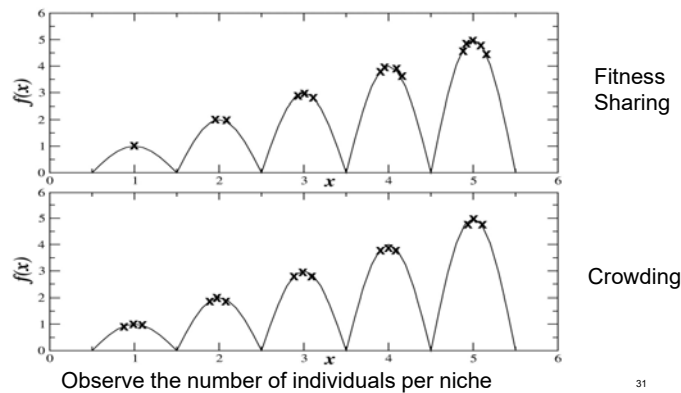


Explicit Approaches for Preserving Diversity: Crowding

- 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.

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Explicit Approaches for Preserving Diversity: Crowding vs Fitness sharing



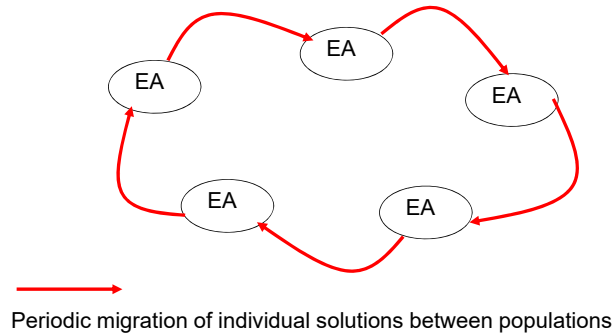
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Implicit Approaches for Preserving Diversity: Automatic Speciation

- 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: “Island” Model Parallel EAs



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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

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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

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Genetic Algorithms: Overview Simple GA

- 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

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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

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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

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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

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X^2 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

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X² example: Crossover

String no.	Mating pool	Crossover point	Offspring after crossover	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

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X² example: Mutation

String no.	Offspring after crossover	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

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**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

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**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

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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

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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

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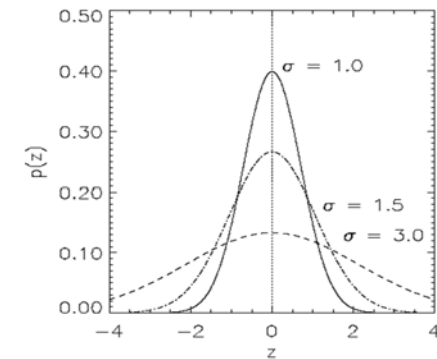
Evolution Strategies: Representation

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

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Evolution Strategies: Adaptive Mutation

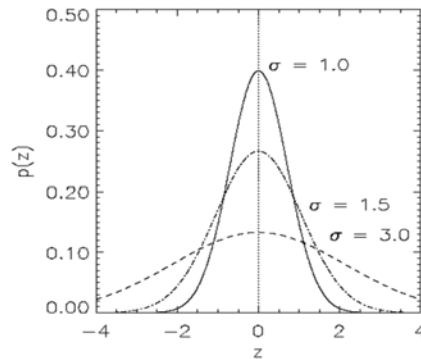
- z values drawn from normal distribution $N(\xi, \sigma)$
 - mean ξ is set to 0
 - variation σ is called **mutation step size**
- σ is varied on the fly by the “1/5 success rule”



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The “1/5 success rule”

- Goal: Balance exploration and exploitation
- Resets σ after every k iterations by
 - $\sigma = \sigma / c$ if $p_s > 1/5$
 - $\sigma = \sigma \cdot c$ if $p_s < 1/5$
 - $\sigma = \sigma$ if $p_s = 1/5$
- where p_s is the % of successful mutations, $0.8 \leq c \leq 1$



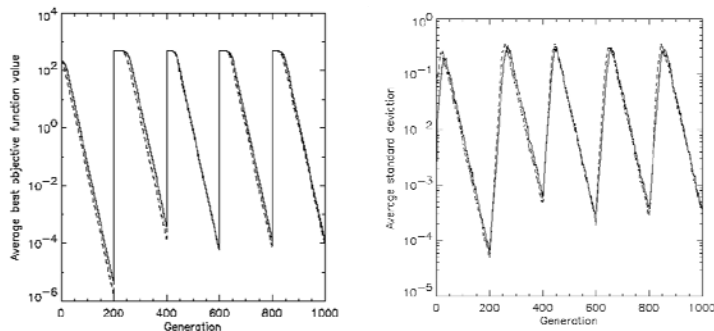
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Evolution Strategies: Self-adaptation illustrated (1/2)

- Given a dynamically changing fitness landscape (optimum location shifted every 200 generations)
- Self-adaptive ES is able to
 - follow the optimum and
 - adjust the mutation step size after every shift !

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Evolution Strategies: Self-adaptation illustrated cont'd (2/2)



Changes in the fitness values (left) and the mutation step sizes (right)

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

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Evolution Strategies: Recombination

- Two parents create one child
- Acts per variable / position by either
 - Averaging parental values, or
 - Selecting one of the parental values
- From two or more parents by either:
 - Local recombination: Two parents make a child
 - Global recombination: Selecting two parents randomly for each gene

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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

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Evolution Strategies: ES summary

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

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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)

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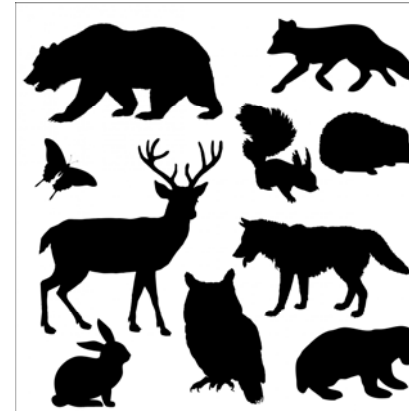
Evolutionary Programming: Representation

- 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: $\langle x_1, \dots, x_n, \sigma_1, \dots, \sigma_n \rangle$

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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



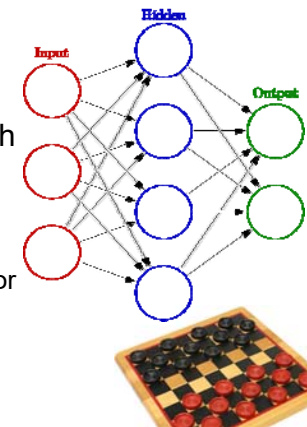
Evolutionary Programming: Selection

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

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Evolutionary Programming: Evolving checkers player (Fogel'02) (1/2)

- Neural nets for evaluating future values of moves are evolved
- NNs have fixed structure with 5046 weights, these are evolved
- Representation:
 - vector of 5046 real numbers for NN weights
- Population size 15



Evolutionary Programming: Evolving checkers player (Fogel'02) (2/2)

- Tournament size $q = 5$
- Programs (with NN inside) play against other programs, no human trainer
- After 840 generation (6 months!) best strategy was tested against humans
- Program earned “expert class” ranking outperforming 99.61% of all rated players

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Evolutionary Programming: Summary

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

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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

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Genetic Programming: Initialisation

- Maximum initial depth of trees D_{\max} is set
- Full method (each branch has depth = D_{\max}):
 - nodes at depth $d < D_{\max}$ randomly chosen from **function set F (IF, AND, =, >, *, etc.)**
 - nodes at depth $d = D_{\max}$ randomly chosen from **terminal set T (x,y,5000,NOC, etc.)**
- Grow method (each branch has depth $\leq D_{\max}$):
 - nodes at depth $d < D_{\max}$ randomly chosen from $F \cup T$
 - nodes at depth $d = D_{\max}$ randomly chosen from T
- Common GP initialisation: ramped half-and-half, where grow & full method each deliver half of initial population

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Genetic Programming: Full Initialisation to depth 2

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Source: www.gp-field-guide.org.uk

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Genetic Programming: Grow Initialisation to depth 2

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Source: www.gp-field-guide.org.uk

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Genetic Programming: Variation Operators

GA flowchart

GP flowchart

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Genetic Programming: Mutation

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

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Genetic Programming: Recombination

Parent 1

Parent 2

Child 1

Child 2

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Genetic Programming: Bloat

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

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Genetic Programming: Summary

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

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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

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Particle Swarm Optimisation: Quick overview

- Developed: in 1995 by Kennedy and Eberhart
- Inspired by social behavior of bird flocking/fish schooling



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Particle Swarm Optimisation: Representation (1/2)

- Every population member can be considered as a pair $\langle \bar{x}, \bar{p} \rangle$ where the first vector is candidate solution and the second one a perturbation vector in \mathbb{R}^n
- The perturbation vector determines how the solution vector is changed to produce a new one: $\bar{x}' = \bar{x} + \bar{p}'$, where \bar{p}' is calculated from \bar{p} and some additional information

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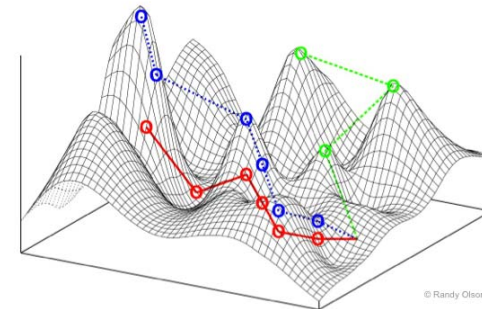
Particle Swarm Optimisation: Representation (2/2)

- A member is a point in space with a position and a velocity
- The perturbation is defined as the weighted sum of three components:
 - Current perturbation vector
 - Vector difference current position to best position of member so far
 - Vector difference from current position to best position of population so far

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Image source: <http://wirelesstechthoughts.blogspot.no/>

A Reminder about Search Landscapes

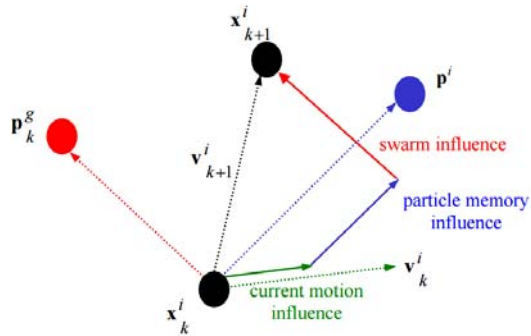


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PSO: Velocity and Position Update

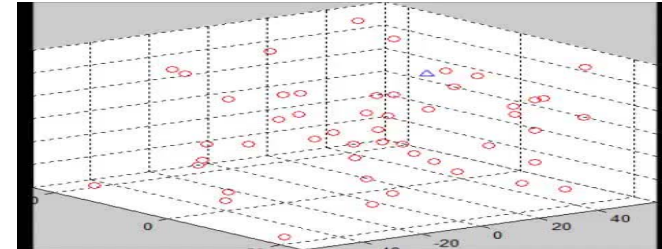


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Image source: Hassan et al. 2004: A COPMARISON OF PARTICLE SWARM OPTIMIZATION AND THE GENETIC ALGORITHM

Particle Swarm Optimisation: Example moving target

- Optimum moves randomly
- Particles do not know position of optimum but do know which particle is closest and are attracted to that one



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Particle Swarm Optimisation: Summary

Representation	Real-valued vectors
Recombination	None
Mutation	Adding velocity vector
Parent selection	Deterministic (each parent creates one offspring via mutation)
Survivor selection	Generational (offspring replaces parents)

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