

Advanced Algorithms

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Lecture 10 – Evolutionary algorithms (cont.)

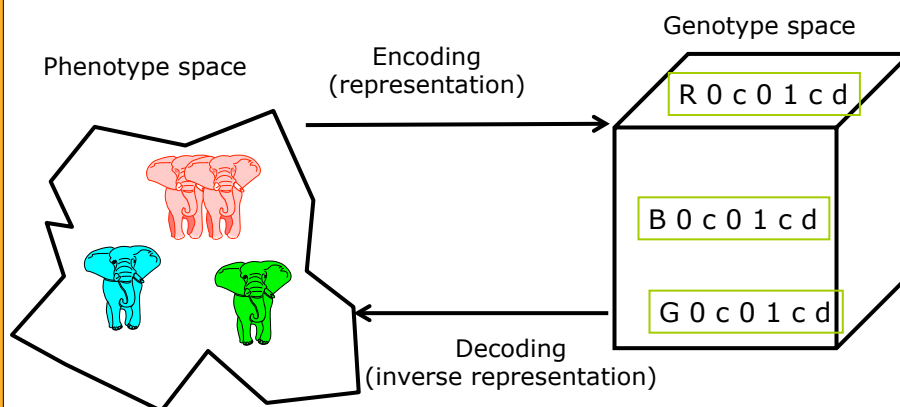
These slides are mainly taken from **A.E. Eiben** and **J.E. Smith**,
Introduction to Evolutionary Computing

Evolutionary algorithms

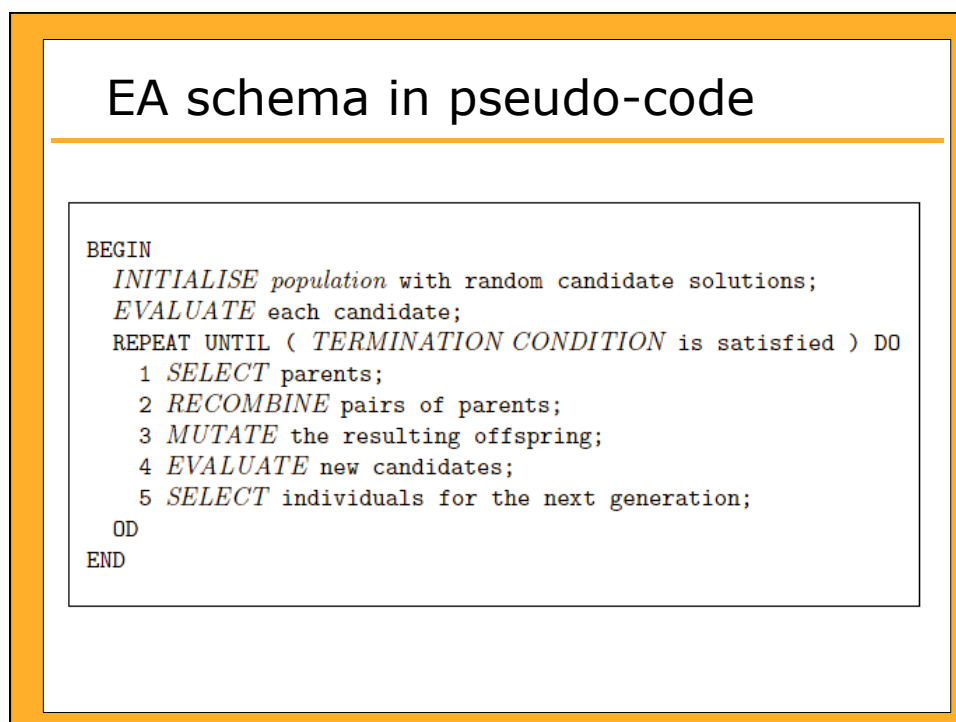
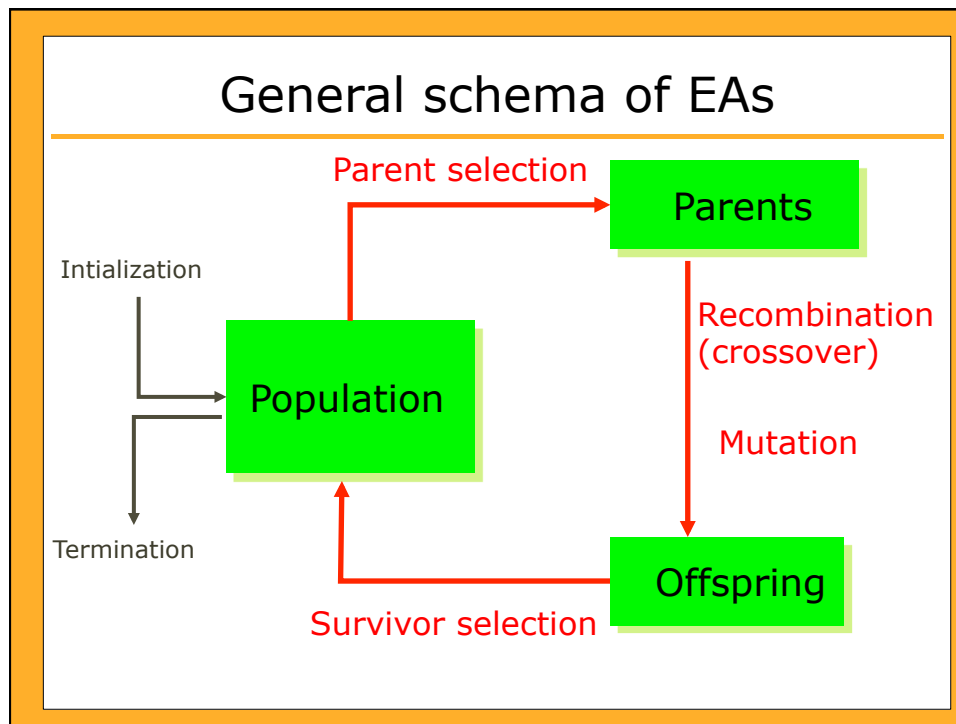
EAs are **randomized Monte Carlo** algorithms that can be used to find (approximate) solutions to a **wide range of computational problems** such as:

- **Optimization**
 - E.g., university timetabling
- **Modeling**
 - E.g., finding a classification model
- **Simulation**
 - E.g., evolving economical models

Representation

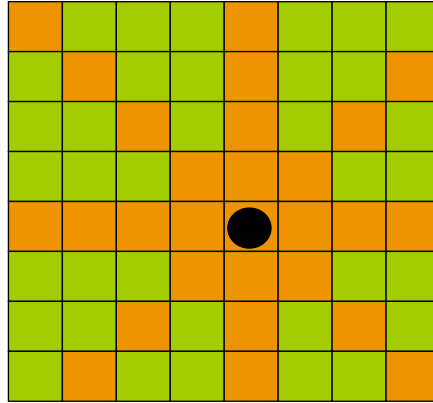


In order to find the global optimum, every feasible solution must be represented in genotype space



Example: the 8-queens problem

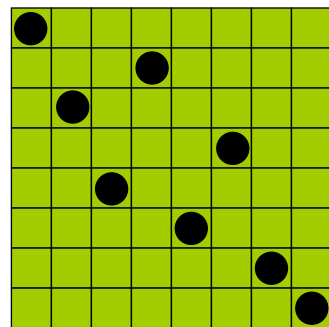
Place 8 queens on an 8x8 chessboard in such a way that they cannot attack each other



The 8-queens problem: representation

Phenotype:
a board configuration

Genotype:
a permutation of
the numbers 1 - 8



1 3 5 2 6 4 7 8

↑↓ Obvious mapping

The 8-queens problem: fitness evaluation

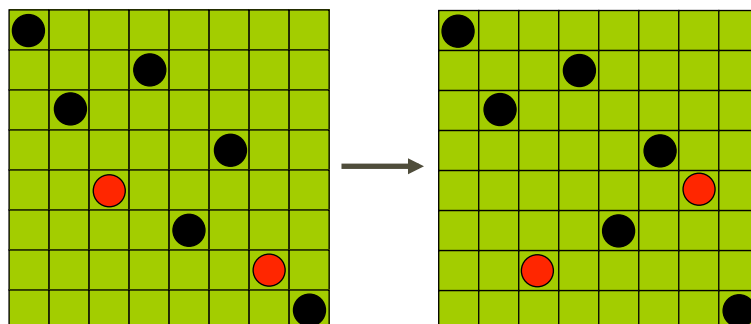
- **Penalty** of one **queen**: the number of queens she can attack
- **Penalty** of a **configuration**: the sum of penalties of all queens
- **Note**: penalty is to be minimized
- **Fitness** of a configuration: inverse penalty to be maximized

The 8-queens problem: mutation

Small variation in one permutation, e.g.:

- swapping values of two randomly chosen positions

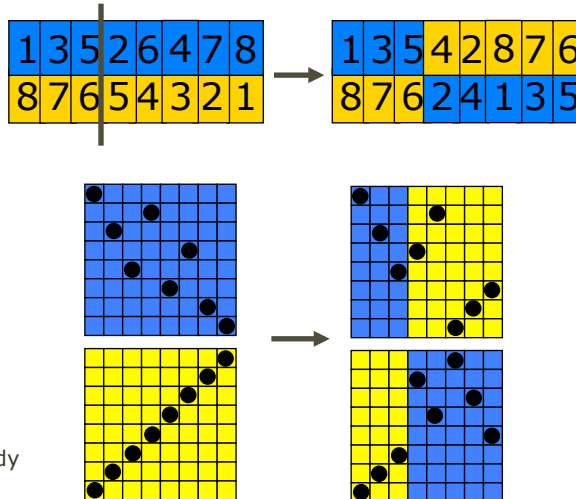
1 3 5 2 6 4 7 8 → 1 3 7 2 6 4 5 8



The 8-queens problem: recombination

Combining two permutations into two new permutations:

- choose random crossover point
- copy first parts into children
- create second part by inserting values from other parent:
 - in the order they appear there
 - beginning after crossover point
 - skipping values already in child



The 8-queens problem: selection

- **Parent selection:**
 - Pick 5 parents at random and take best two to undergo crossover
- **Survivor selection (replacement)**
 - When inserting a new child into the population, choose an existing member to replace by:
 - sorting the whole population by decreasing fitness
 - enumerating this list from high to low
 - replacing the first with a fitness lower than the given child

8 Queens Problem: summary


Representation	Permutations
Recombination	“Cut-and-crossfill” crossover
Recombination probability	100%
Mutation	Swap
Mutation probability	80%
Parent selection	Best 2 out of random 5
Survival selection	Replace worst
Population size	100
Number of Offspring	2
Initialisation	Random
Termination condition	Solution or 10,000 fitness evaluation

Note that is **only one possible** set of choices of operators and parameters

Typical behavior of an EA

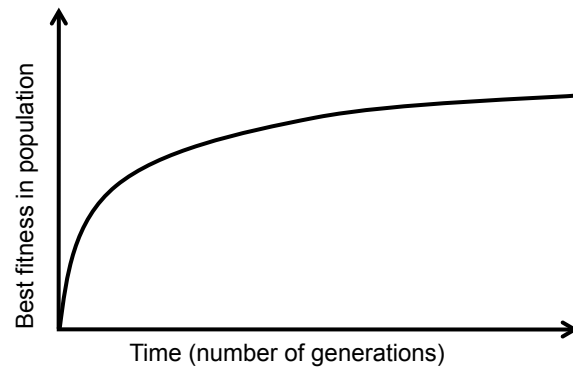
Stages in **optimizing** on a **1-dimensional** fitness landscape

 **Early stage:**
quasi-random population distribution

 **Mid-stage:**
population arranged **around/on hills**

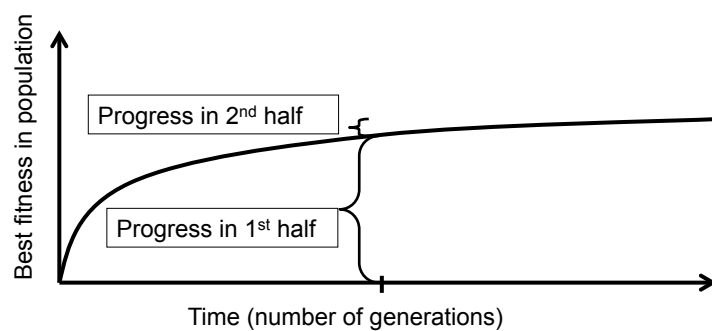
 **Late stage:**
population concentrated on **high hills**

Typical run: progression of fitness



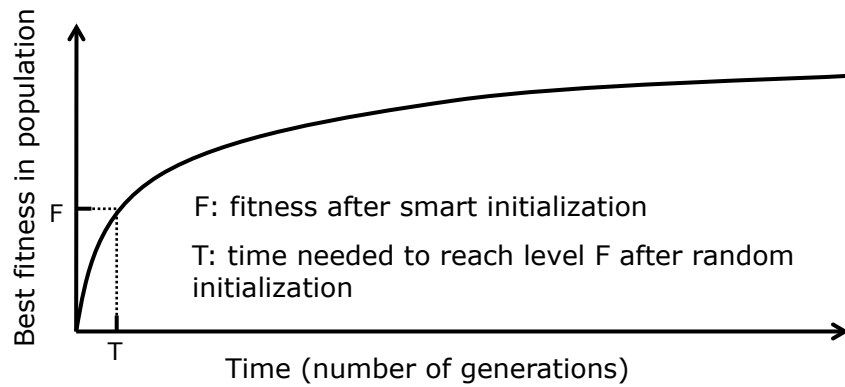
Typical run of an EA shows so-called “**anytime behavior**”

Typical run: progression of fitness



- Are long runs beneficial?
 - It depends on how much you want the last bit of progress
 - May be better to do more short runs

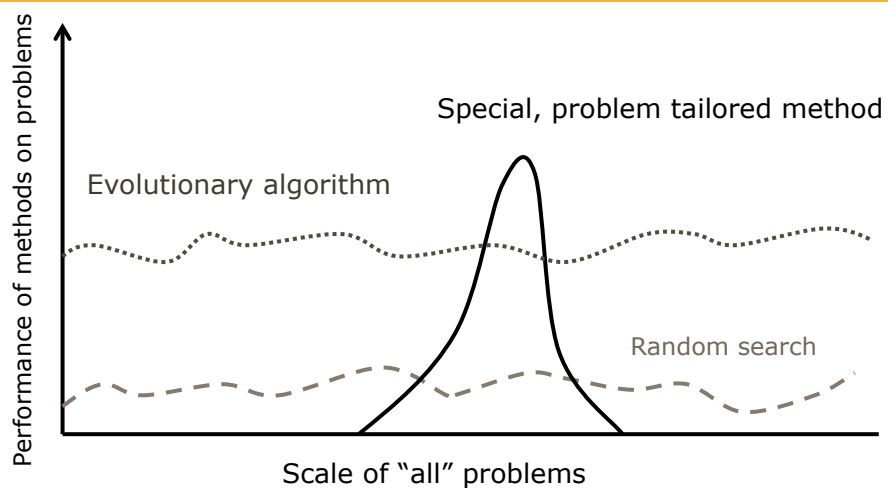
Is it worth expending effort on smart initialization?

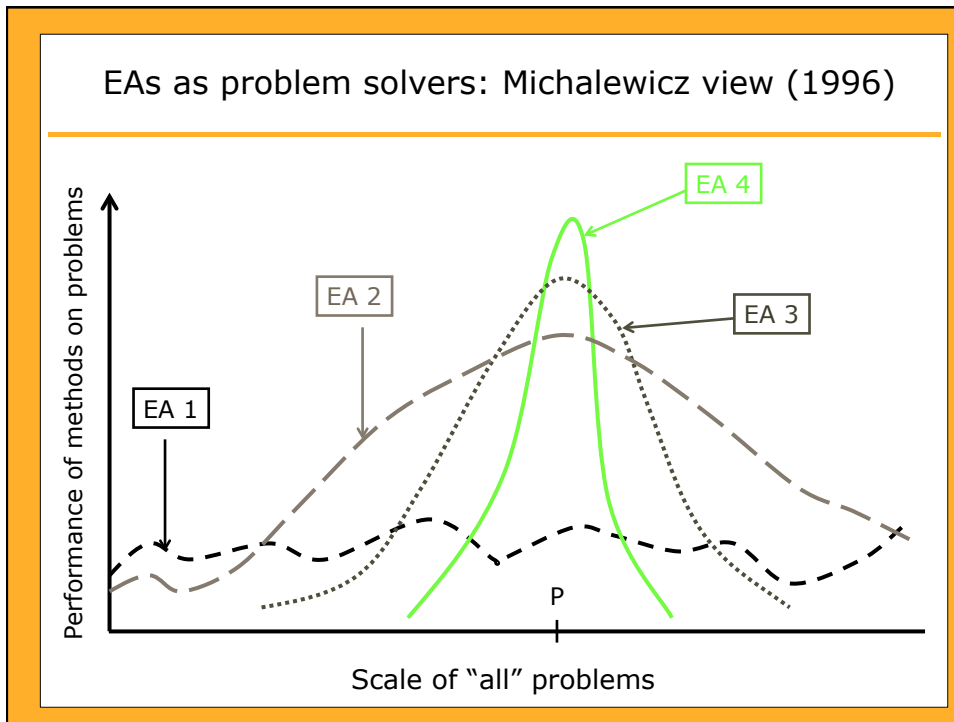


Answer: **it depends**

- Possibly good, if good solutions/methods exist
- Care is needed

EAs as problem solvers: Goldberg view (1989)



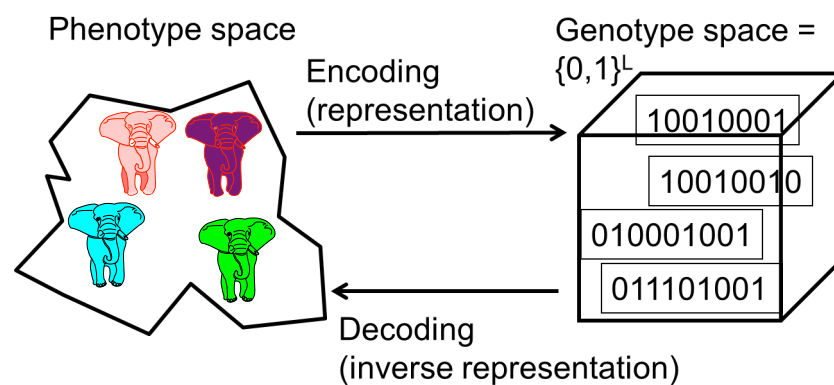


Genetic Algorithms

GA Quick Overview

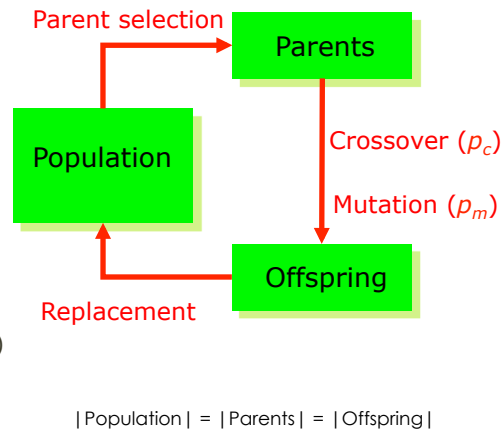
- Developed: USA in the 1970's
- Early names: J. Holland, K. DeJong, D. Goldberg
- Holland's original GA is now known as the simple genetic algorithm (SGA)
- Other GAs use different:
 - Representations
 - Mutations
 - Crossovers
 - Selection mechanisms

Representation



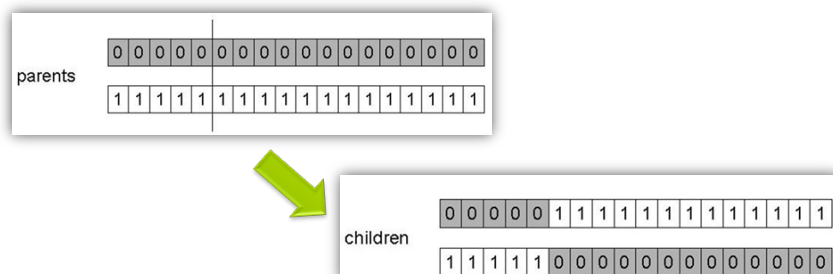
SGA reproduction cycle

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



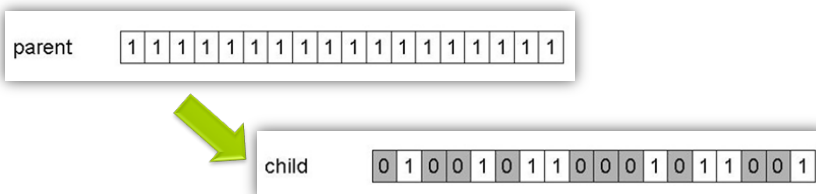
SGA operators: 1-point crossover

- Apply with probability p_c
 - p_c typically in range (0.6, 0.9)
- Choose a random point on the two parents
- Split parents at this crossover point
- Create children by exchanging tails



SGA operators: mutation

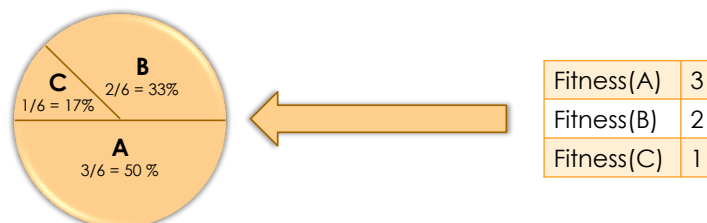
- Alter each gene independently with a probability p_m
- p_m is called the mutation rate
 - Typically between $1/\text{pop_size}$ and $1/\text{chromosome_length}$



SGA operators: Selection

Main idea: better individuals get higher chance

- Chances proportional to fitness
- Implementation: roulette wheel technique
 - Assign to each individual a part of the roulette wheel
 - Spin the wheel n times to select n individuals



An example after Goldberg '89 (1)

- 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 crossover, bitwise mutation
 - Roulette wheel selection
 - Random initialization

X^2 example: selection

We show one generational cycle done by hand

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

The simple GA

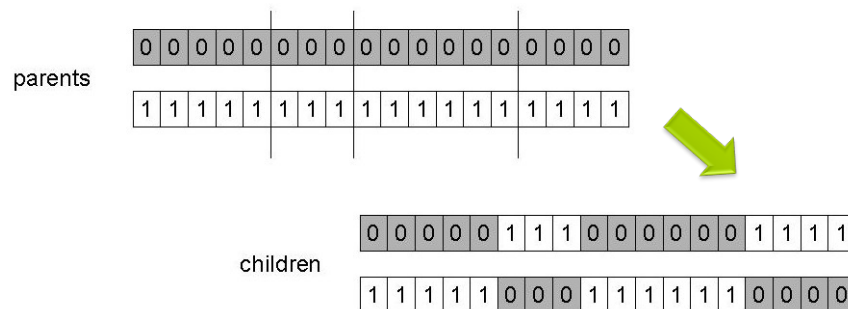
- Has been subject of many (early) studies
 - still often used as benchmark for novel GAs
- Shows many shortcomings, e.g.,
 - Representation is too restrictive
 - Mutation & crossover operators only applicable for bit-string
 - Selection mechanism sensitive for converging populations with close fitness values
 - Generational population model (step 5 in SGA repr. cycle) can be improved with explicit survivor selection

Alternative Crossover Operators

- Performance with 1 Point Crossover depends on the order that variables occur in the representation
 - more likely to keep together genes that are near each other
 - can never keep together genes from opposite ends of string
 - this is known as *Positional Bias*
 - can be exploited if we know about the structure of our problem, but this is not usually the case

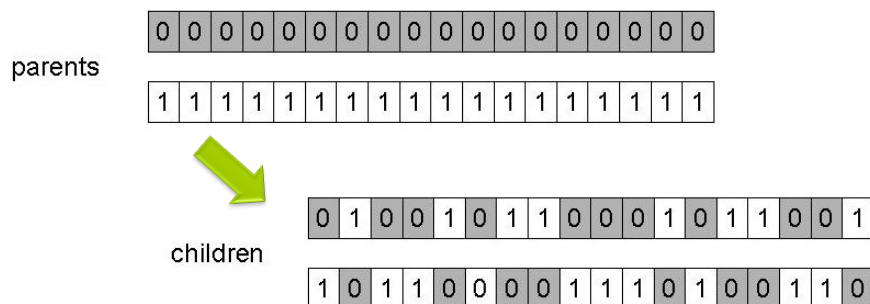
n-point crossover

- Choose n random crossover points
- Split along those points
- Glue parts, alternating between parents
- Generalisation of 1 point (still some positional bias)



Uniform crossover

- Flip a coin for each gene of the first child
 - Keep the value if 'heads', change the value if 'tails'
 - Make an inverse copy of the gene for the second child
- Inheritance is independent of position



Crossover OR mutation?

- Decade long debate:
 - which one is better?
 - Are both necessary?

- Answer (at least, rather wide agreement):
 - it depends on the problem
 - in general, **it is good to have both**
 - mutation-only-EA is possible, crossover-only-EA would not work

Crossover OR mutation? (cont'd)

Exploration: Discovering promising areas in the search space, i.e. gaining information on the problem

Exploitation: Optimising within a promising area, i.e. using information

There is co-operation AND competition between them

- **Crossover is explorative**
 - it makes a *big* jump to an area somewhere "in between" two (parent) areas
- **Mutation is exploitative**
 - it creates random *small* diversions, thereby staying near (in the area of) the parent

Crossover OR mutation? (cont'd)

- Only **crossover** can **combine information from two parents**
- Only **mutation** can **introduce new information** (alleles)
- To hit the **optimum** you often need a '**lucky**' **mutation**

Population Models

- SGA uses a **Generational model**:
 - each individual survives for exactly one generation
 - **the entire set of parents is replaced by the offspring**
- At the **other end** of the scale **are Steady-State** models:
 - one offspring is generated per generation
 - **one member of population replaced**
- **Generation Gap**
 - the **proportion** of the population **replaced**
 - makes a parameterized transition between generational and steady-state Gas
 - **gg = 1.0 for SGA, gg = 1/pop_size for SSGA**
- The name SSGA is often used for any GA with a generation gap < 1

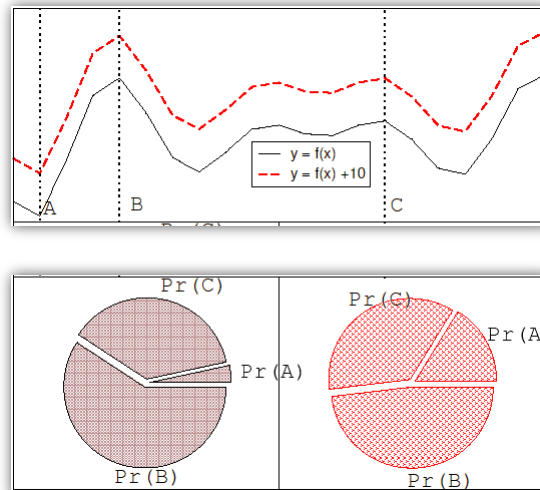
Fitness Based Competition

- **Selection** can occur in **two places**:
 - Selection from current generation to take part in mating (**parent selection**)
 - Selection from parents + offspring to go into next generation (**survivor selection**)
- **Selection operators** work on whole individual
 - i.e. they **are representation-independent** !

Fitness-Proportionate Selection

- 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**
 - Highly **susceptible to function transposition** (see next slide)

Function transposition for FPS



Fitness-Proportionate Selection

- 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**
 - Highly **susceptible** to **function transposition** (see next slide)
- **Scaling** can fix the last two problems
 - **Windowing** $f'(i) = f(i) - \beta$
 - where β is worst fitness in this generation
 - **Sigma Scaling** $f'(i) = \max(f(i) - (\bar{f} - c * \sigma_f), 0.0)$
 - where c is a constant, usually 2.0

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 μ and worst rank 1)
- This imposes a sorting overhead on the algorithm, but this is usually negligible compared to the fitness evaluation time

Linear Ranking

- Parameterised by factor s : $1.0 < s \leq 2.0$
 - measures **advantage of best individual**
 - In SGA this is the number of children allotted to it

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

- Simple 3 member example

	Fitness	Rank	P_{selFP}	$P_{selLR} (s = 2)$	$P_{selLR} (s = 1.5)$
A	1	1	0.1	0	0.167
B	5	3	0.5	0.67	0.5
C	4	2	0.4	0.33	0.33
Sum	10		1.0	1.0	1.0

Survivor Selection

- Most of **selection methods above** are used for **parent selection**
- **Survivor selection** can be divided into two approaches:
 - **Age-Based Selection**
 - In SGA the population is fully replaced ad each generation
 - In SSGA can implement as "delete-random" (not recommended) or as first-in-first-out (a.k.a. delete-oldest)
 - **Fitness-Based Selection**
 - Using one of the methods above

Two Special Cases of fitness-based survivor selection

- **Replace-worst**
 - The **worst** (in term of fitness) individuals are replaced and each generation by the offspring
- **Elitism**
 - Always keep at least one copy of the fittest solution so far
 - Widely used in both population models (SGA, SSGA)

SGA technical summary tableau

Representation	Binary Strings
Recombination	N-point or uniform crossover
Mutation	Bitwise bit-flipping with fixed probability
Parent selection	Fitness-Proportionate
Survivor selection	All children replace parents
Speciality	Emphasis on crossover