

Evolutionary Algorithms

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Evolutionary Algorithms (EA)

group of search and optimization algorithms
based on

- Mendelian genetics
- Darwinian theory of evolution

which are

- heuristic
- stochastic
- population based

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Evolutionary Algorithms

- genetic algorithms (Holland 1975)
- genetic programming (Koza 1989)
- evolutionary strategies (Rechenberg 1973)

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Evolutionary Algorithms

- genetic algorithms
- genetic programming
- evolutionary strategies
- evolutionary programming
- differential evolution
- grammatical evolution
- memetic algorithm

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EA Outline

Algorithm EA:

```
INITIALIZE population randomly  
CALCULATE_FITNESS of each individual  
while not STOP_CRITERIA do  
    SELECT parents  
    RECOMBINE pairs of parents  
    MUTATE offspring  
    CALCULATE_FITNESS of offspring  
    REPLACE (some) parents by offspring  
end_do
```

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EA Terminology

- gene
- chromosome
- fitness
- individual (contains a chromosome and has a fitness) = candidate solution
- parents, offspring
- population
- generations = iterations

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Components of an EA

- representation for candidate solutions
- a population of candidate solutions
- method for creating an initial population of candidate solutions
- evaluation function to rate candidate solutions
- parent selection mechanism
- genetic operators to alter composition of population of candidate solutions
- replacement mechanism
- various parameters to control a run

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Parameters of an EA

- number of generations
 - or some other stopping criteria
- population size, i.e. no. of individuals in a population
- probability of applying variation operators
- chromosome length (depends on problem instance)
- other parameters

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The Simple Genetic Algorithm (SGA)

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Simple GA -SGA

- a.k.a., Canonical GA
- Operators of a SGA
 - selection
 - crossover
 - mutation

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SGA

```
generate initial population  
repeat  
    evaluate individuals  
    perform reproduction  
    select pairs  
    recombine pairs  
    apply mutation  
until end_of_generation
```

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Representation & Encoding

- population size constant
- individual has one chromosome
- chromosome length constant
- individual has a fitness value
- binary genes (0/1)
- generational

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Initial Population

random initial population



each gene value for each individual determined
randomly to be either *0* or *1*
with equal probability

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Fitness Evaluation

- fitness function
 - objective function(s)
 - constraints
- shows fitness of individual
 - degree to which the candidate solution meets the objective
- apply fitness function to individual

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Example Problem: One-Max

Objective: maximize the number of **1**s in a string of length 5, composed only of **1**s and **0**s

➡ *population size = 4*

chromosome length = 5

*fitness function = no. of genes that are **1***

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Example Population

individual 1:

chromosome = 11001

fitness = 3

individual 3:

chromosome = 11111

fitness = 5

individual 2:

chromosome = 00001

fitness = 1

individual 4:

chromosome = 01110

fitness = 3

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Reproduction

- consists of
 - selection
 - mating pool (size same as population)
 - possibly more than one copy of some individuals
 - crossover
 - mutation

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Selection: Roulette-Wheel Selection

- fitness proportionate selection
 - expected no. of representatives of each individual is proportional to its fitness

$$prob_i = \frac{fitness_i}{\sum_j fitness_j} , j = 1, \dots, pop.size$$

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Selection: Roulette-Wheel Selection

```
begin
  set current_member=1;
  while (current_member ≤ m)do
    pick uniform r.v. r from [0,1];
    set i=1;
    while (ai < r) do
      set i=i+1;
    od
    set mating_pool[current_member]=parents[i];
    set current_member=current_member+1;
  od
end
```

m: population size, $a_i = \sum_1^i P_{sel}(i)$, $i = 1, 2, \dots, m$

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Selection: Tournament Selection

- ordinal based
- roulette wheel selection uses info on whole population
 - info may not be available
 - population too large
 - population distributed on a parallel system
 - maybe no universal fitness definition (e.g., game playing, evolutionary art, evolutionary design)

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Selection: Tournament Selection

- relies on an ordering relation to rank any n
- individuals
- most widely used approach
- tournament size k
 - if k large, more of the fitter individuals get a chance

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Selection: Tournament Selection

```
begin
  set current_member=1;
  while (current_member  $\leq$  m) do
    pick k individuals randomly;
    select best from k individuals;
    denote this individual i;
    set mating_pool[current_member]=i;
    set current_member=current_member+1;
  od
end
```

m: population size , k: tournament size

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Example: Roulette Wheel Selection

Current Population:

i1: 11001, 3

i2: 00001, 1

i3: 11111, 5

i4: 01110, 3

Expected copies of each individual in pool:

i1: $(3/12 * 4)$ 1

i2: $(1/12 * 4)$ 0

i3: $(5/12 * 4)$ 2

i4: $(3/12 * 4)$ 1

Probability of each individual being selected:

$\text{prob}(i1) = 3/12 = 0.25$

$\text{prob}(i2) = 1/12 = 0.08$

$\text{prob}(i3) = 5/12 = 0.42$

$\text{prob}(i4) = 3/12 = 0.25$

Assume:

wheel is turned 4 times

1 copy of i1

2 copies of i3

1 copy of i4

is copied into mating pool

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Recombination

- process for creating new individuals (offspring)
 - from two or more parents
- term used interchangeably with crossover
 - mostly refers to 2 parents
- crossover rate **pc**
 - typically in range [0.5,1.0]
 - acts on parent pair

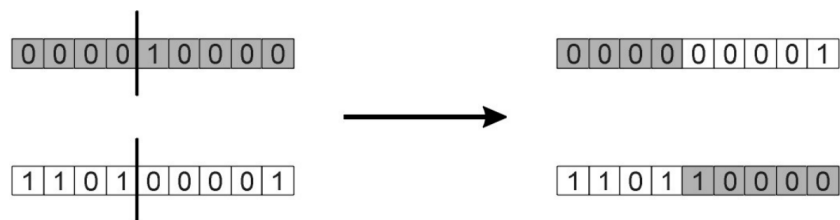
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Recombination

- two parents selected randomly
- a r.v. drawn from $[0,1)$
- if value $< p_c$, then two offspring created through recombination
- else, two offspring created asexually
 - i.e., copy of parent

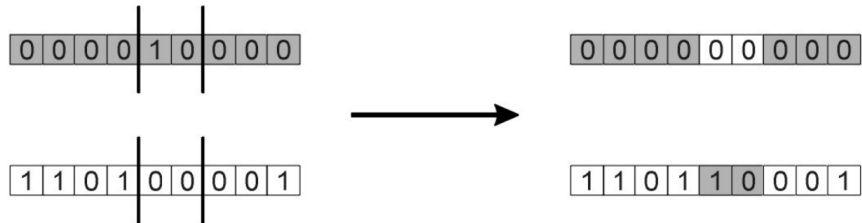
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One-Point Crossover



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Two-Point Crossover



Can be N point crossover. In the above example: N=2

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Example Pairing

Current mating pool:

mate 1: 11001 (i1)

mate 2: 11111 (i2)

mate 3: 11111 (i3)

mate 4: 01110 (i4)

Assume:

As a result of random drawing
(mate 1, mate 3)
and
(mate 2, mate 4)
are paired off for reproduction.

Pairs:

Pair 1:

11001

11111

Pair 2:

11111

01110

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Example One-Point Crossover

Assume: $p_c = 1.0$

for pair 1:

crossover site: 3

110 | 01 \rightarrow 11011

111 | 11 \rightarrow 11101

for pair 2:

crossover site: 1

1 | 1111 \rightarrow 11110

0 | 1110 \rightarrow 01111

the new individuals:

i1: 11011

i2: 11101

i3: 11110

i4: 01111

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Mutation

- bitwise mutation
- probability of mutation: p_m
 - a.k.a., mutation rate
 - equal probability for each gene
 - chromosome of length L ; expected no. of changes: $L \cdot p_m$
 - typically chosen to be small
 - depends on nature of problem

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Bitwise Mutation



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Example Mutation

Assume:

as a result of random draws,
1st gene of i1
and
4th gene of i3
are found to undergo mutation

i1: 11011 → 01011

i3: 11110 → 11100

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Generational SGA

- non-overlapping populations
- offspring replace parents
- elitism may be used
 - no. of elite individuals e
 - elite individuals replace worst individuals in the new population (if they are better)

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Example New Population (no elitism)

individual 1:

chromosome = 01011

fitness = 3

individual 2:

chromosome = 11101

fitness = 4

individual 3:

chromosome = 11100

fitness = 3

individual 4:

chromosome = 01111

fitness = 4

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Stopping Criteria

- main loop repeated until stopping criteria met
 - for a predetermined no. of generations ✓
 - until a goal is reached
 - until population converges

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Example Stopping Criteria

- *number of generations*= 250
 - loop repeated 250 times
 - best individual at each generation (*lbest*) found
 - overall best individual (*gbest*) becomes solution

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