

# Biased Random-Key Genetic Algorithms

Algoritmos Genéticos de Chaves Aleatórias Viciadas

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Este material foi desenvolvido baseando-se em  
<http://mauricio.resende.info/talks/2012-09-CLAI02012-brkga-tutorial-both-days.pdf>

# Summary

- 1 Genetic algorithms (GAs)
- 2 Random-key genetic algorithms (RKGAs)
- 3 Biased random-key genetic algorithm (BRKGAs)

# Genetic algorithms

## Introduction

- ▶ Genetic algorithms (GAs) are metaheuristics inspired by the process of natural selection
- ▶ GAs evolve population of individuals (solutions) applying Darwin's principle of survival of the fittest
- ▶ A GA maintains a population of candidate individuals (solutions) for the problem at hand, and makes it evolve by iteratively applying a set of **stochastic operators** (selection, recombination and mutation)
  - selection: replicates the most successful individuals found in a population at a rate proportional to their relative quality
  - recombination (crossover): decomposes two or more distinct individuals and then randomly mixes their parts to form novel solutions
  - mutation: randomly perturbs a candidate individual

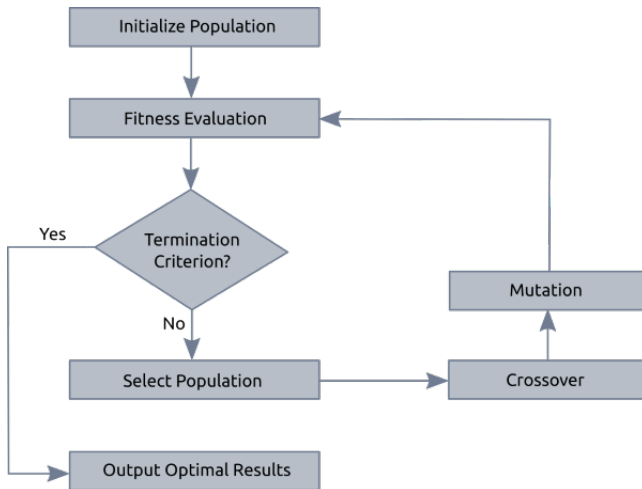
# Genetic algorithms

## Components

- ▶ Encoding principles (gene, chromosome)
- ▶ Initialization procedure (creation)
- ▶ Selection of parents (reproduction)
- ▶ Genetic operators (mutation, recombination)
- ▶ Evaluation function (environment)
- ▶ Termination condition

# Genetic algorithms

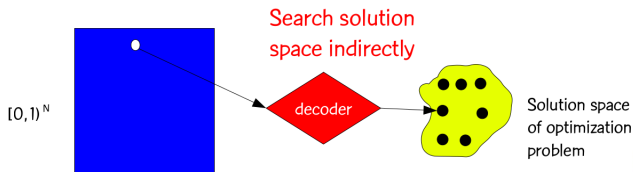
## Evolutionary cycle



# Random-key genetic algorithms

## Introduction

- ▶ Introduced by Bean (1994)<sup>1</sup> for sequencing problems
- ▶ A random-key is a real random number in the continuous interval  $[0,1]$
- ▶ Individuals (solutions) of optimization problems can be encoded by random-keys
- ▶ Individuals are strings of real-valued numbers (random-keys)
- ▶ A decoder is a deterministic algorithm that takes a vector of random-keys as input and outputs a solution of the optimization problem



<sup>1</sup>Bean, J. C. (1994). Random-key genetic algorithms for sequencing and optimization. ORSA journal on computing, 6(2), 154-160.

# Random-key genetic algorithms

## Encoding/decoding principles

- ▶ Bean (1994)<sup>1</sup> proposed decoders based on sorting the random-key vector to produce a sequence

- ▶ Encoding:

$$s = \langle \underset{1}{0.25}, \underset{2}{0.19}, \underset{3}{0.67}, \underset{4}{0.05}, \underset{5}{0.89} \rangle$$

- ▶ Decode by sorting vector of random-keys:

$$s' = \langle \underset{4}{0.05}, \underset{2}{0.19}, \underset{1}{0.25}, \underset{3}{0.67}, \underset{5}{0.89} \rangle$$

- ▶ Therefore, the vector of random-keys:

$$s = \langle 0.25, 0.19, 0.67, 0.05, 0.89 \rangle$$

encodes the sequence:  $\langle 4, 2, 1, 3, 5 \rangle$

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# Random-key genetic algorithms

## Encoding/decoding principles

- ▶ Other decodings:
  - subset selection (select 3 of 5 elements)



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- subset selection (select 3 of 5 elements)

encoding:

$$s = \langle \underset{1}{0.099}, \underset{2}{0.216}, \underset{3}{0.802}, \underset{4}{0.368}, \underset{5}{0.658} \rangle$$

decode by sorting vector of random-keys:

$$s' = \langle \underset{1}{0.099}, \underset{2}{0.216}, \underset{4}{0.368}, \underset{5}{0.658}, \underset{3}{0.802} \rangle$$

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- subset selection (select  $k$  of 5 elements, where  $0 \leq k \leq 5$ )

encoding:

$$s = \langle \underset{1}{0.82}, \underset{2}{0.12}, \underset{3}{0.54}, \underset{4}{0.89}, \underset{5}{0.26} \rangle$$

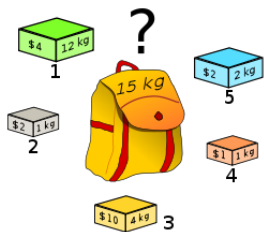
decoding: if  $s_i \geq 0.5$  then select  $i$

encodes the subset:  $\{1, 3, 4\}$

# Random-key genetic algorithms

## Encoding/decoding principles

- ▶ For some cases of complex decoders is necessary to adjust the chromosome in order to produce a feasible solution
- ▶ Knapsack problem:



encoding:

$$s = \langle 0.82, 0.12, 0.54, 0.89, 0.26 \rangle$$

1            2            3            4            5

decoding: if  $s_i \geq 0.5$  then get item  $i$

encodes the subset items:  $\{1, 3, 4\}$

$$\sum_{i=1,3,4} w_i \geq W \quad (12 + 4 + 1 \geq 15)$$

so, a **deterministic** strategy must be applied to make the solution viable

# Random-key genetic algorithms

## Initialize population

- Initial population is made up of  $P$  random-key vectors, each with  $N$  keys, each having a value generated uniformly at random in the interval  $[0,1)$ .

$$s_1 = \langle \text{KEY}_1^1, \text{KEY}_2^1, \dots, \text{KEY}_N^1 \rangle$$

$$s_2 = \langle \text{KEY}_1^2, \text{KEY}_2^2, \dots, \text{KEY}_N^2 \rangle$$

...

$$s_P = \langle \text{KEY}_1^P, \text{KEY}_2^P, \dots, \text{KEY}_N^P \rangle$$

$$\text{KEY}_j^i \leftarrow \text{rand}[0, 1)$$

$$\forall i = 1..P, \forall j = 1..N$$

# Random-key genetic algorithms

## Selection of parents and recombination

- ▶ Two parents  $a$  and  $b$  are randomly selected from the entire  $P$  population
- ▶ Mating is done using parameterized uniform crossover (Spears & DeJong, 1990)<sup>2</sup> for sequencing problems.

$$a = \langle 0.25, 0.19, 0.67, 0.05, 0.89 \rangle$$

$$b = \langle 0.63, 0.90, 0.76, 0.93, 0.08 \rangle$$

- ▶ For each gene, flip a biased coin to choose which parent passes the allele (key, or value of gene) to the child.

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If every random-key array corresponds to a feasible solution: Mating always produces feasible offspring.

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# Random-key genetic algorithms

## ~~Mutation~~ Mutants

- ▶ **No mutation:** mutants are used instead (they play same role as mutation in GAs ... help escape local optima)
- ▶ A mutant is a new individual generated uniformly at random in the interval  $[0,1)$

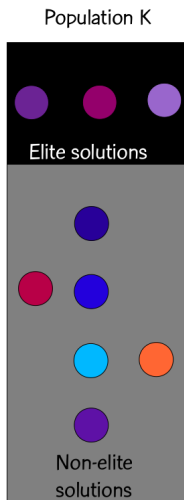
$$m = \langle \text{KEY}_1^m, \text{KEY}_2^m, \dots, \text{KEY}_N^m \rangle$$

$$\text{KEY}_j^m \leftarrow \text{rand}[0, 1) \quad \forall j = 1..N$$

# Random-key genetic algorithms

## Next generation

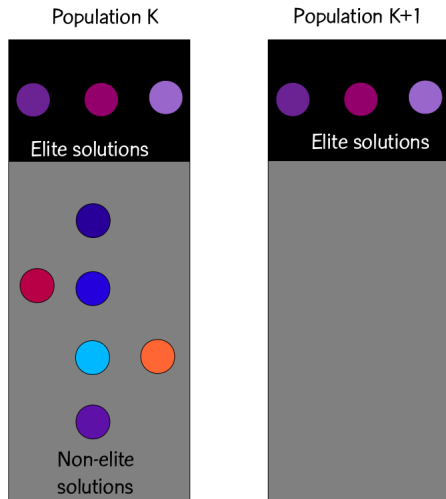
- ▶ At the  $k$ -th generation, compute the cost of each solution and partition the individuals into two sets:
  - elite individuals ( $P_e$ )
  - non-elite individuals ( $\bar{P}_e = P \setminus P_e$ )
- ▶ Elite set should be smaller of the two sets and contain best individuals, i.e.,  $|P_e| < |P|/2$



# Random-key genetic algorithms

Next generation

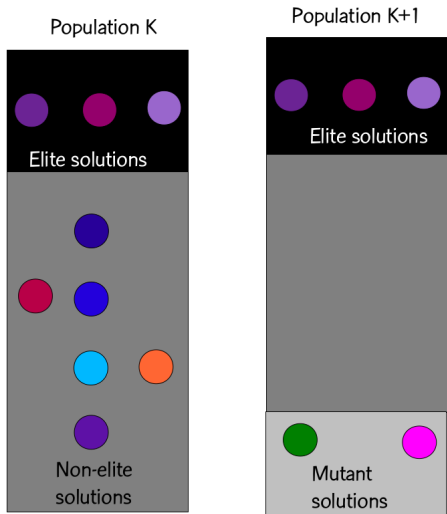
- Copy elite individuals  $P_e$  from population  $k$  to population  $k + 1$



# Random-key genetic algorithms

Next generation

- ▶ Copy elite individuals  $P_e$  from population  $k$  to population  $k + 1$
- ▶ Add  $|P_m|$  random individuals (mutants) to population  $k + 1$

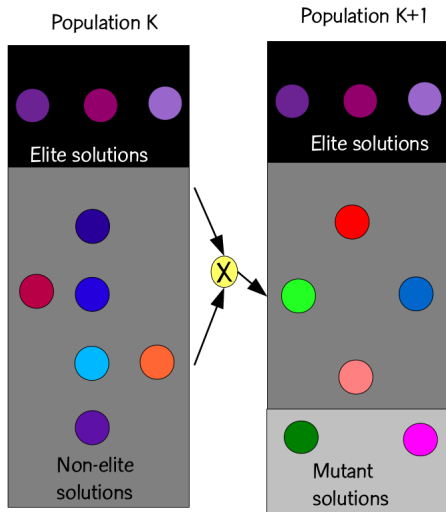




# Random-key genetic algorithms

Next generation

- ▶ Copy elite individuals  $P_e$  from population  $k$  to population  $k + 1$
- ▶ Add  $|P_m|$  random individuals (mutants) to population  $k + 1$
- ▶ While  $k+1$ -th population  $< P$  do
  - Use any two individuals in population  $k$  to produce child in population  $k+1$ . Mates are chosen at random.



# Biased random-key genetic algorithm

## Introduction

- ▶ A biased random-key genetic algorithm (BRKGA) is a random-key genetic algorithm (RKGA).
- ▶ BRKGA and RKGA differ in how mates are chosen for crossover and how parameterized uniform crossover is applied.

# Biased random-key genetic algorithm

## Selection of parents and recombination

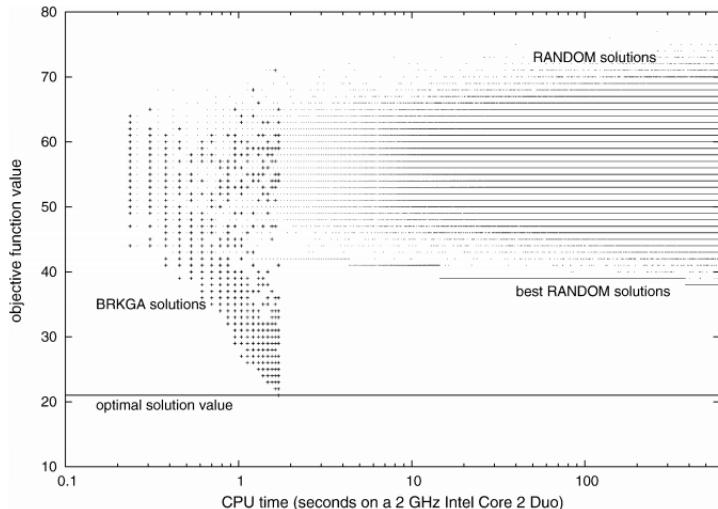
- ▶ A parent  $a$  is randomly selected from the elite population  $P_e$
- ▶ A parent  $b$  is randomly selected from the non-elite population  $\bar{P}_e$  **or** is randomly selected from entire population  $P$
- ▶ For  $i = 1, \dots, n$ , the  $i$ -th allele  $c_i$  of the offspring  $c$  takes on the value of the  $i$ -th allele  $a_i$  of the elite parent  $a$  with probability  $\rho_e$  and the value of the  $i$ -th allele  $b_i$  of the non-elite parent  $b$  with probability  $1 - \rho_e$

Parent $a$	0.32	0.77	0.53	0.85
Parent $b$	0.26	0.15	0.91	0.44
Random number	0.58	0.89	0.68	0.25
$\rho_e = 0.7$	<	>	<	<
Offspring $c$	0.32	0.15	0.53	0.85

- ▶ In this way, the offspring is more likely to inherit characteristics of the elite parent than those of the non-elite parent.

# Biased random-key genetic algorithm

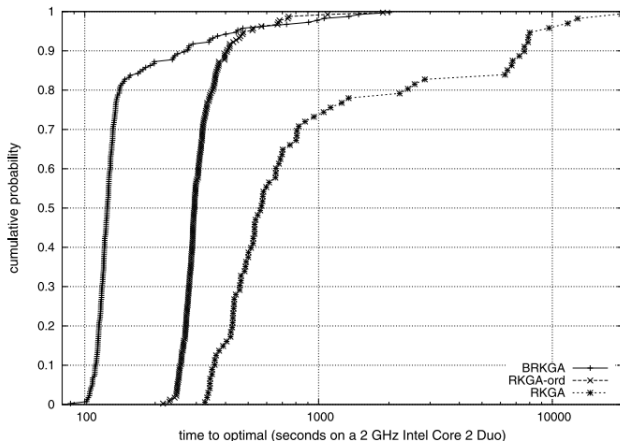
## Random solutions vs BRKGA solutions



Comparing a BRKGA with a random multistart heuristic on an instance of a covering by pairs problem

# Biased random-key genetic algorithm

## RKGA solutions vs BRKGA solutions

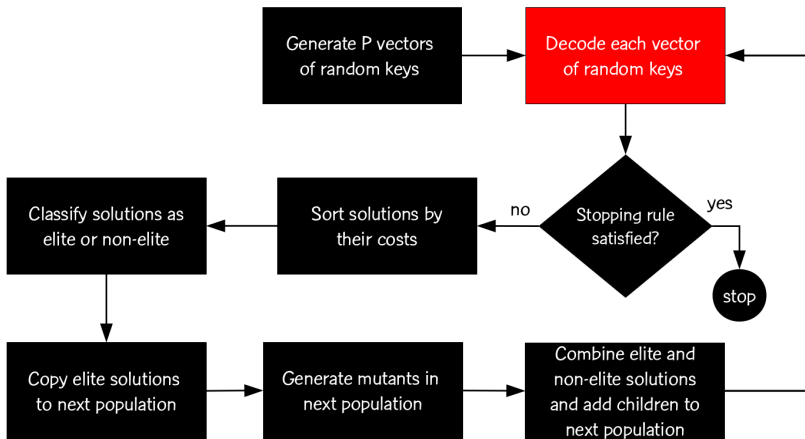


Time to target plots compare running times needed to find the optimal solution of a 220 element covering by pairs problem with a BRKGA and two variants of a RKGA

RKGA-ord is similar to a RKGA except that the offspring inherit the allele of the better fit of the two parents with probability  $\rho_e$

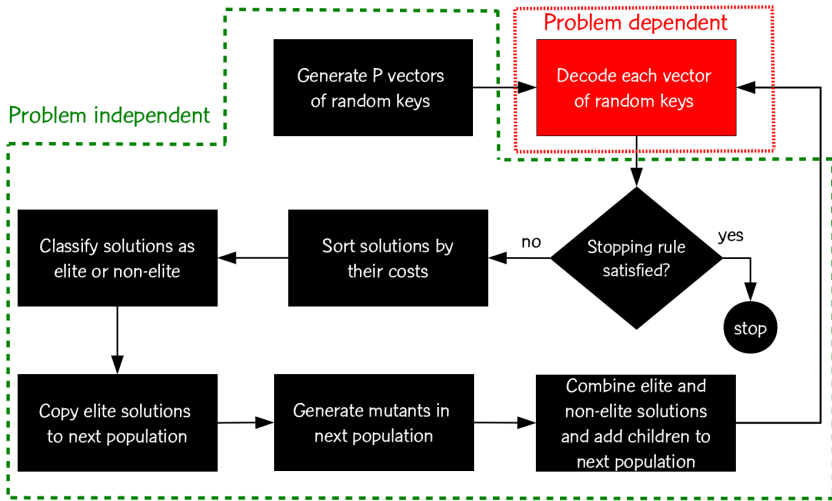
# Biased random-key genetic algorithm

## Framework



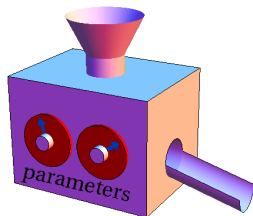
# Biased random-key genetic algorithm

## Framework



# Biased random-key genetic algorithm

## Parameters





# Biased random-key genetic algorithm

## Parameters

Parameter	Description	Recommended value
$P$	size of population	$P = aN$ , where $1 \leq a \in \mathbb{R}$ is a constant and $N$ is the length of the chromosome
$P_e$	size of elite population	$0.10P \leq P_e \leq 0.25P$
$P_m$	size of mutant population	$0.10P \leq P_m \leq 0.30P$
$\rho_e$	elite allele inheritance probability	$0.5 < \rho_e \leq 0.8$
<i>STOP</i>	stopping criterion	e.g. time, # generations, solution quality, # generations without improvement

# Biased random-key genetic algorithm

brkgaAPI: A C++ API for BRKGA

- ▶ Efficient and easy-to-use object oriented application programming interface (API) for the algorithmic framework of BRKGA
- ▶ Download: <http://mauricio.resende.info/src/brkgaAPI/>

- ▶ Resende, M. G. (2012). Biased random-key genetic algorithms: A tutorial  
<http://mauricio.resende.info/talks/2012-09-CLAI02012-brkga-tutorial-both-days.pdf>
- ▶ Gonçalves, J. F., & Resende, M. G. (2011). Biased random-key genetic algorithms for combinatorial optimization. *Journal of Heuristics*, 17(5), pp. 487-525.