

# Biased random-key genetic algorithms

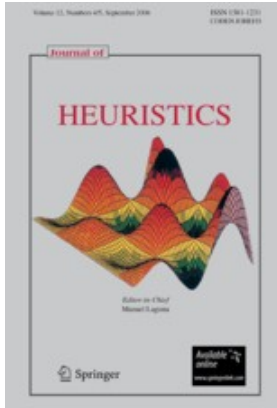
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Lecture given at LION 7  
Catania, Italy ♣ January 8, 2013



# Reference



J.F. Gonçalves and M.G.C.R., “**Biased random-key genetic algorithms for combinatorial optimization,**” J. of Heuristics, vol.17, pp. 487-525, 2011.

Tech report version:

<http://www.research.att.com/~mgcr/doc/srkga.pdf>

# Summary

- Metaheuristics and basic concepts of genetic algorithms
- Random-key genetic algorithm of Bean (1994)
- Biased random-key genetic algorithms (BRKGA)
  - Encoding / Decoding
  - Initial population
  - Evolutionary mechanisms
  - Problem independent / problem dependent components
  - Multi-start strategy
  - Specifying a BRKGA
  - Application programming interface (API) for BRKGA
- Example of a BRKGA for 2-dim packing
- Brief overview of literature

# Metaheuristics

Metaheuristics are heuristics to devise heuristics.

# Metaheuristics

**Metaheuristics** are high level procedures that coordinate simple heuristics, such as **local search**, to find solutions that are of better quality than those found by the simple heuristics alone.

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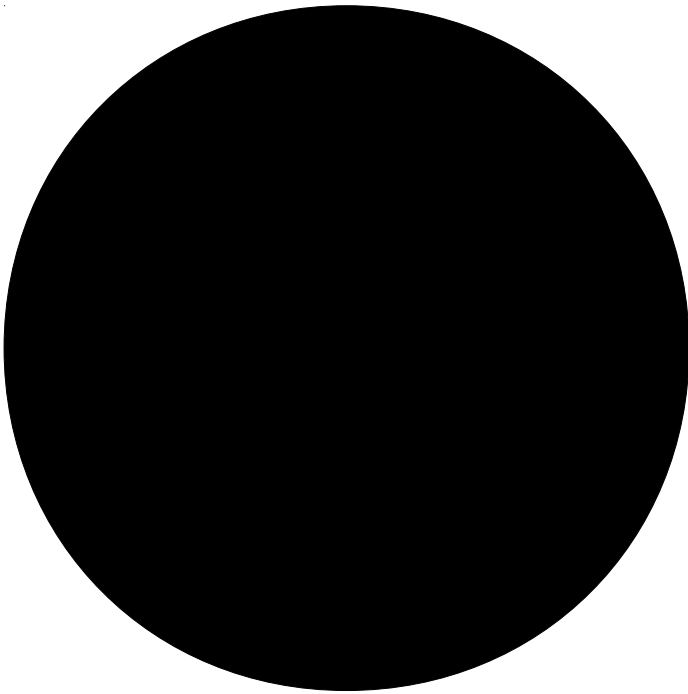
**Examples:** GRASP and C-GRASP, simulated annealing, genetic algorithms, tabu search, scatter search, ant colony optimization, variable neighborhood search, and **biased random-key genetic algorithms (BRKGA)**.

# Genetic algorithms

# Genetic algorithms

Holland (1975)

Adaptive methods that are used to solve search and optimization problems.

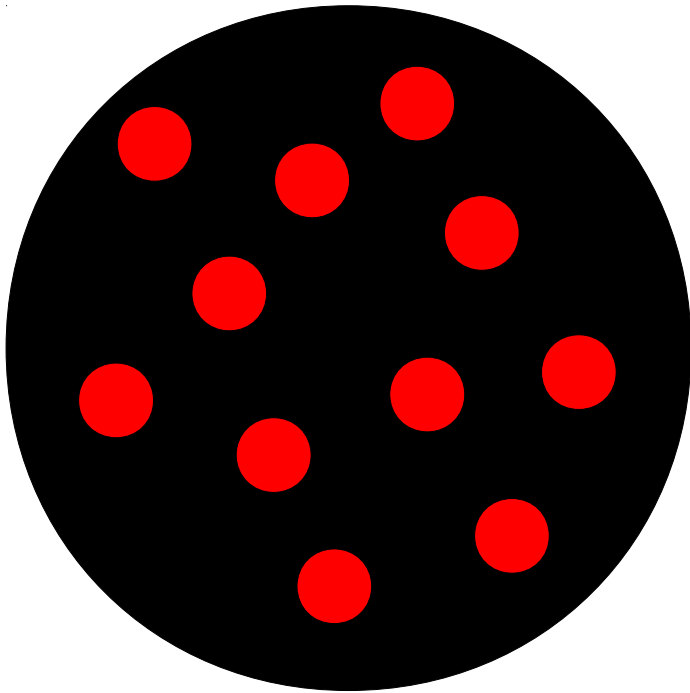


Individual: solution





# Genetic algorithms

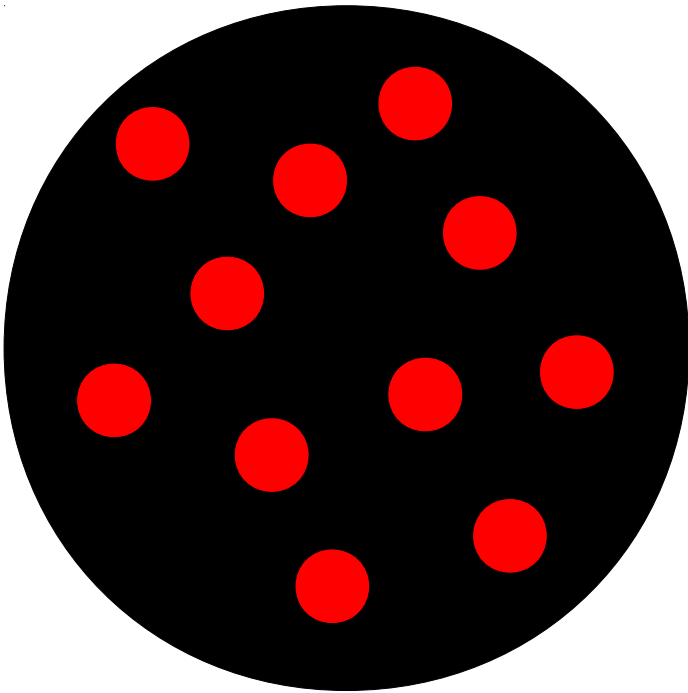


Individual: solution (chromosome = string of genes)

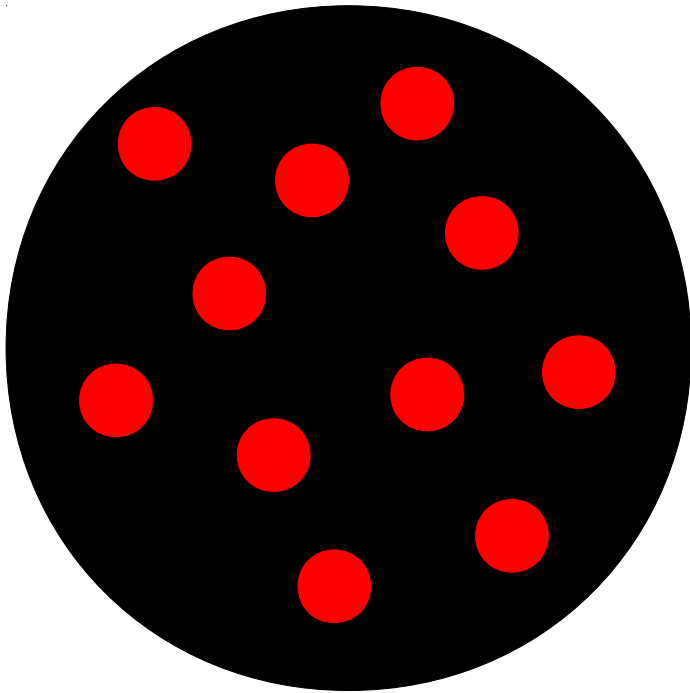
Population: set of fixed number of individuals

# Genetic algorithms

Genetic algorithms evolve population applying Darwin's principle of survival of the fittest.



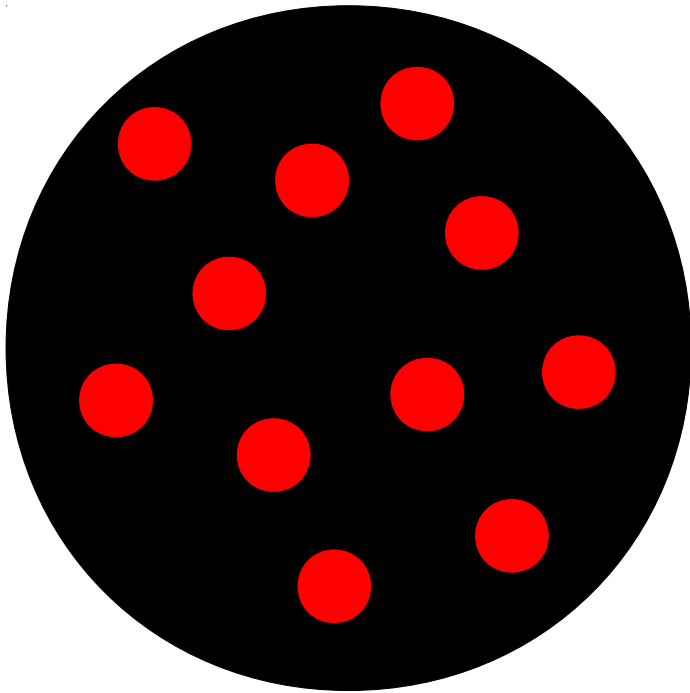
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A series of generations are produced by the algorithm. The most fit individual of the last generation is the solution.

# Genetic algorithms

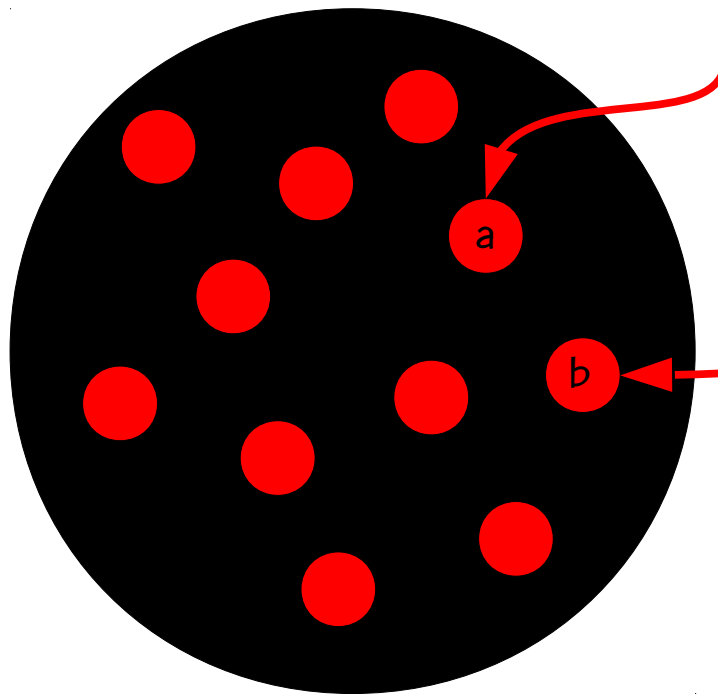


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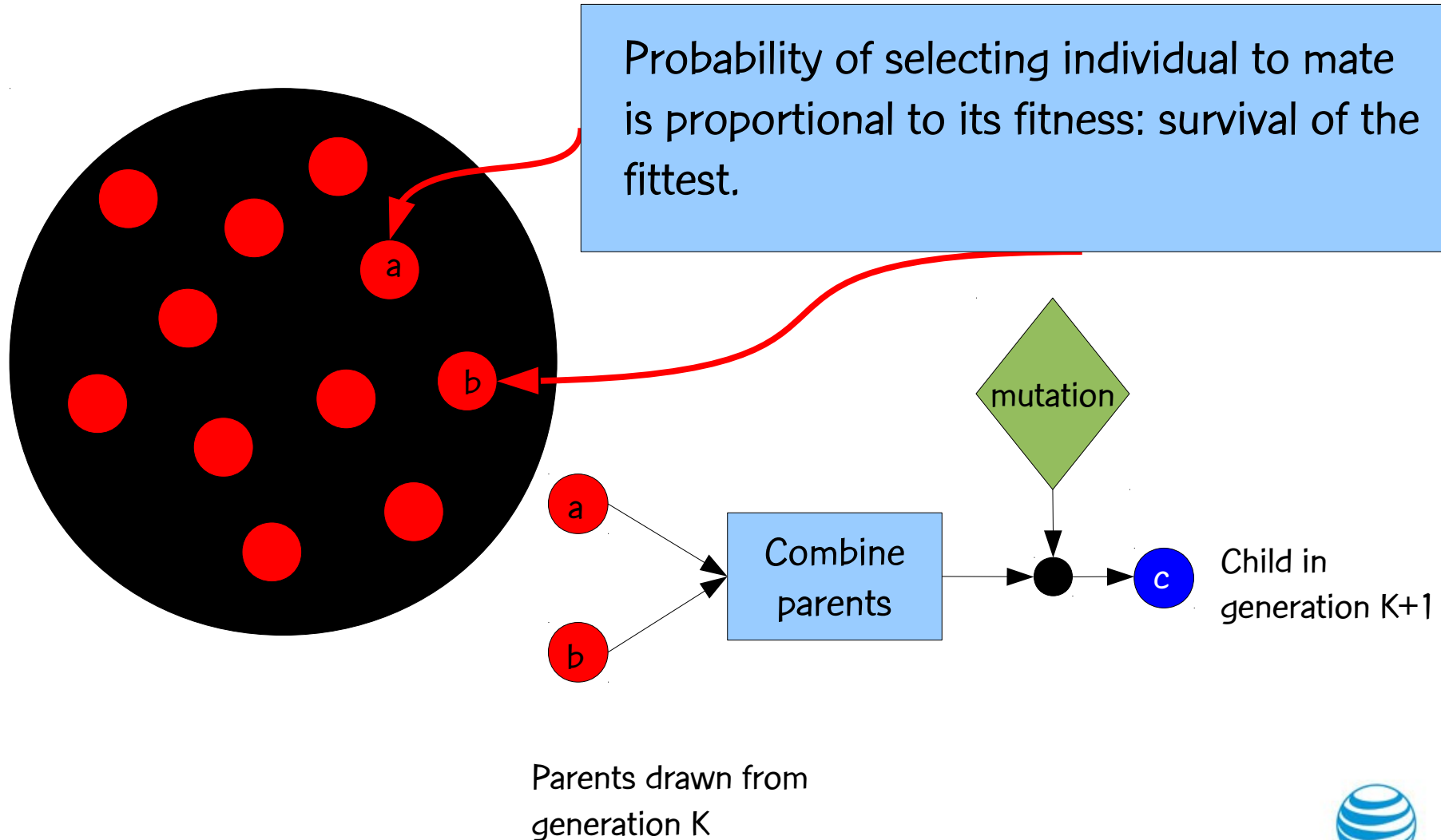
Individuals from one generation are combined to produce offspring that make up next generation.

# Genetic algorithms

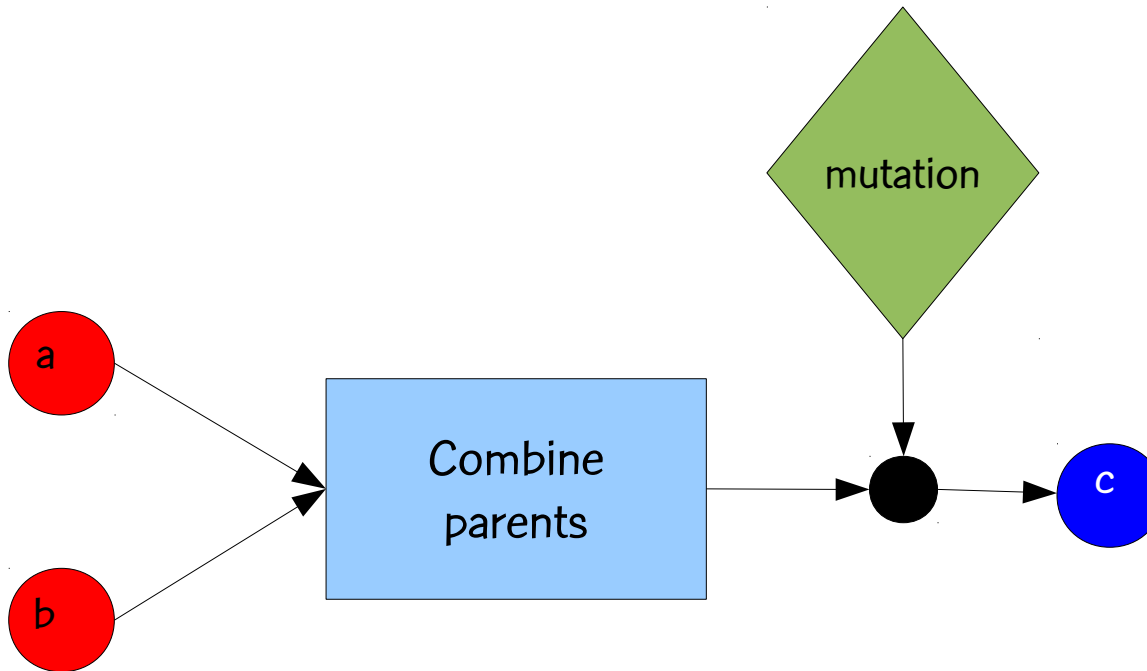


Probability of selecting individual to mate is proportional to its fitness: survival of the fittest.

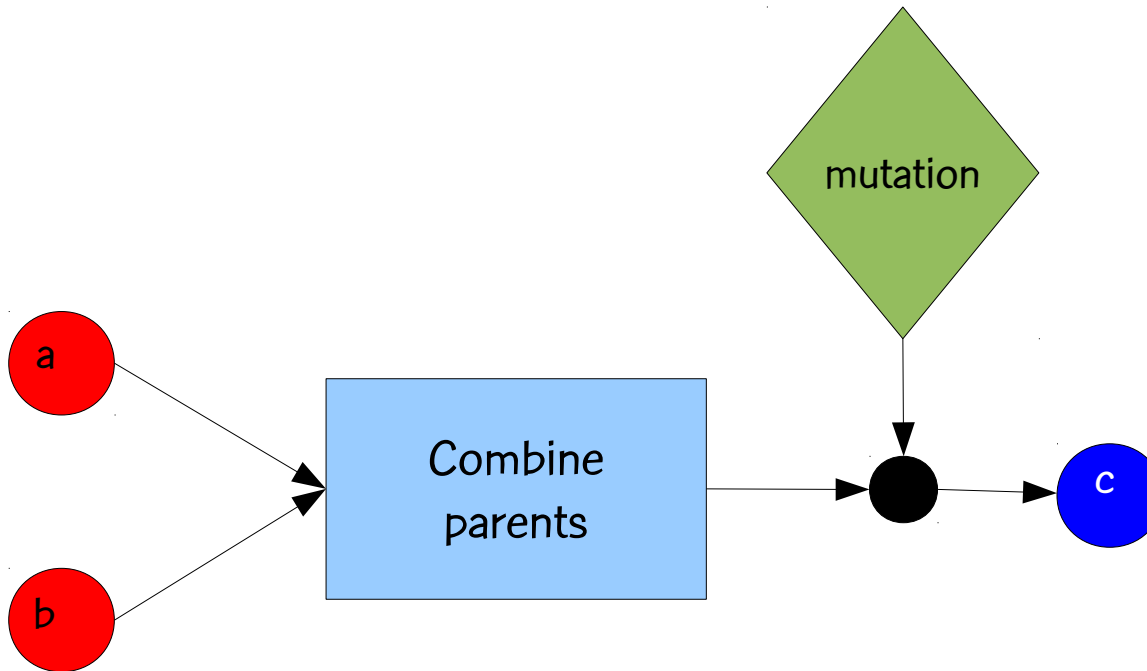
# Genetic algorithms



# Crossover and mutation



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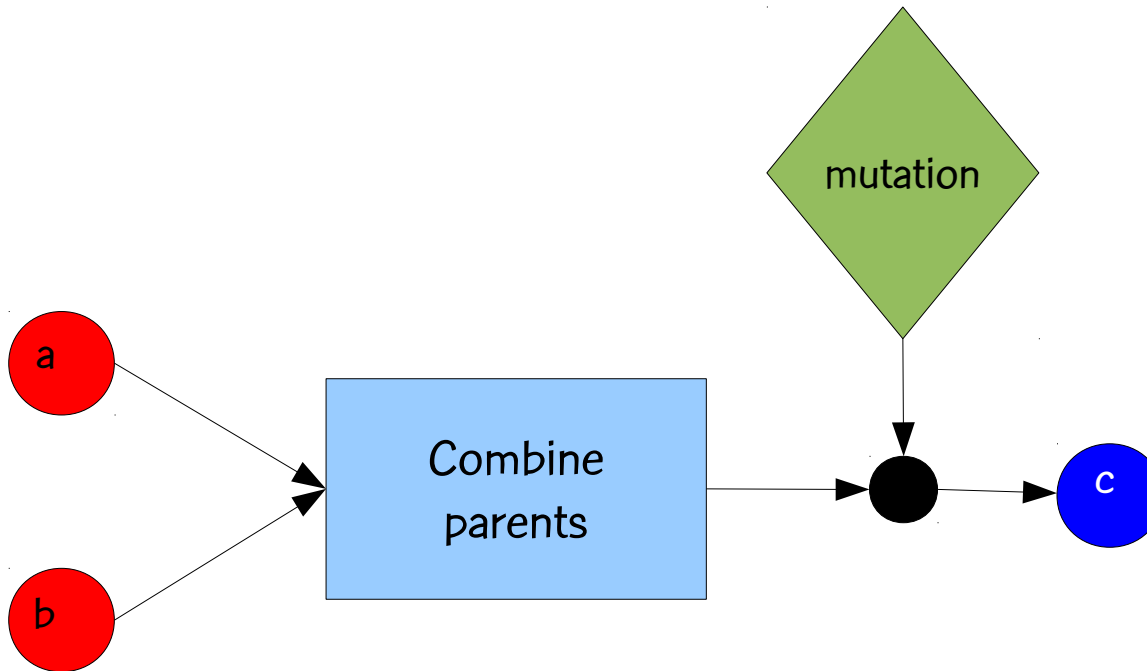


Crossover: Combines parents ... passing along to offspring characteristics of each parent ...

Intensification of search



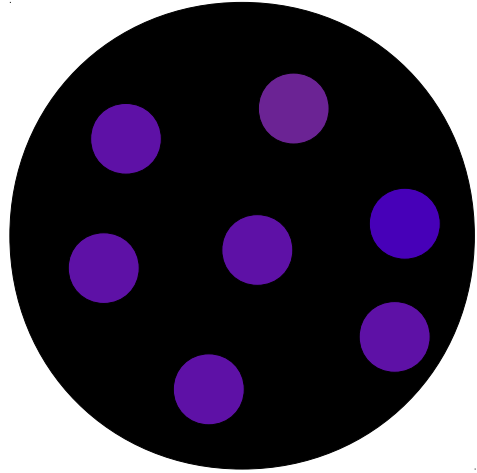
# Crossover and mutation



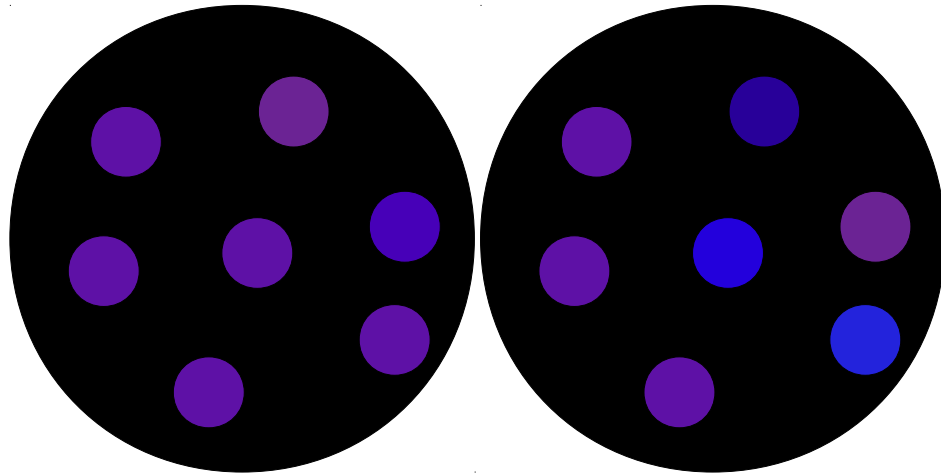
Mutation: Randomly changes chromosome of offspring ...  
Driver of evolutionary process ...

Diversification of search

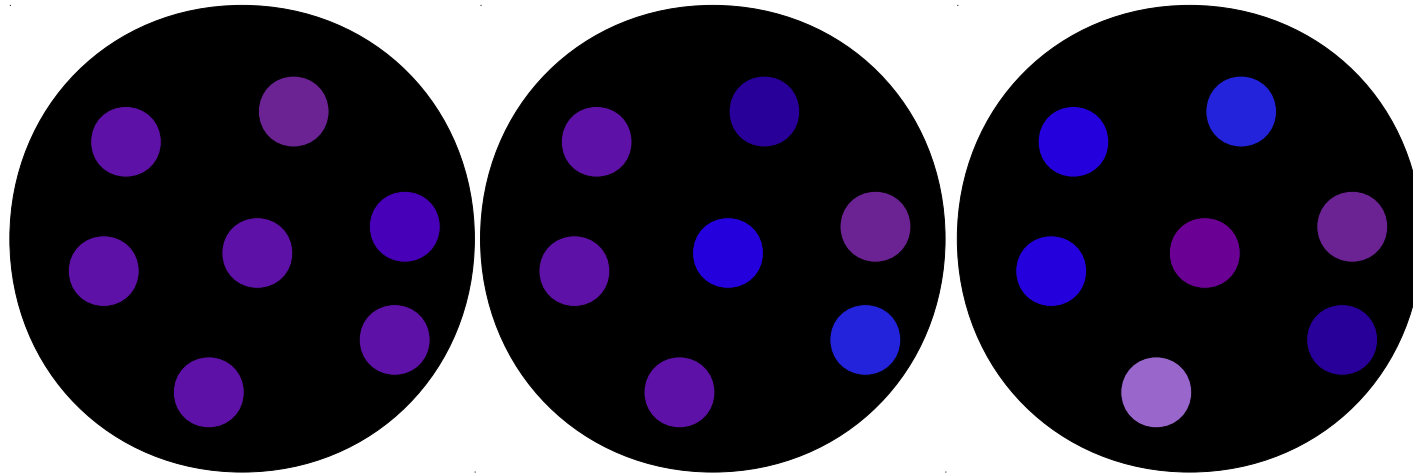
# Evolution of solutions



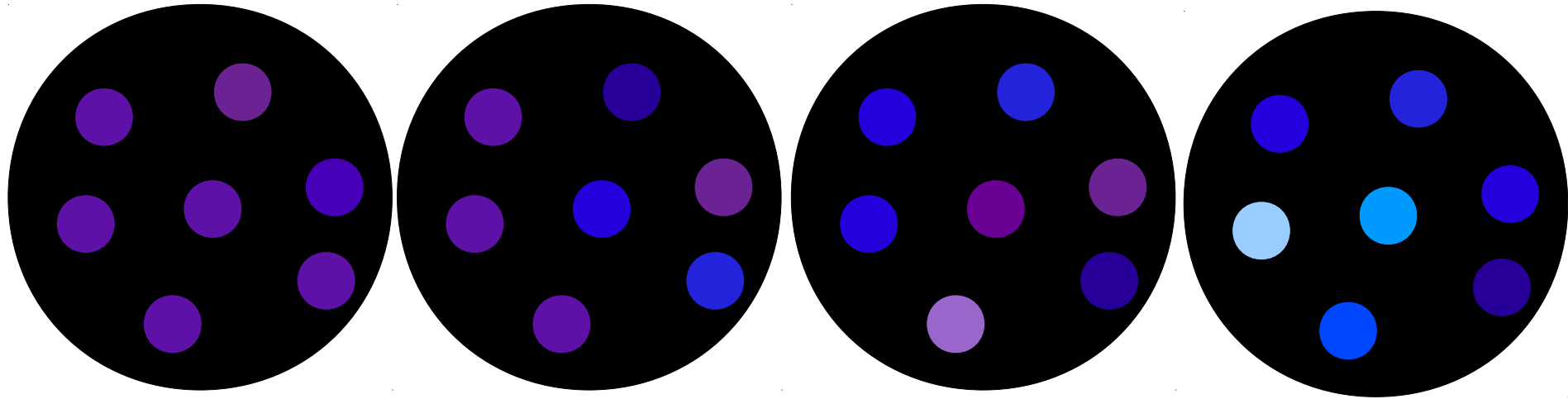
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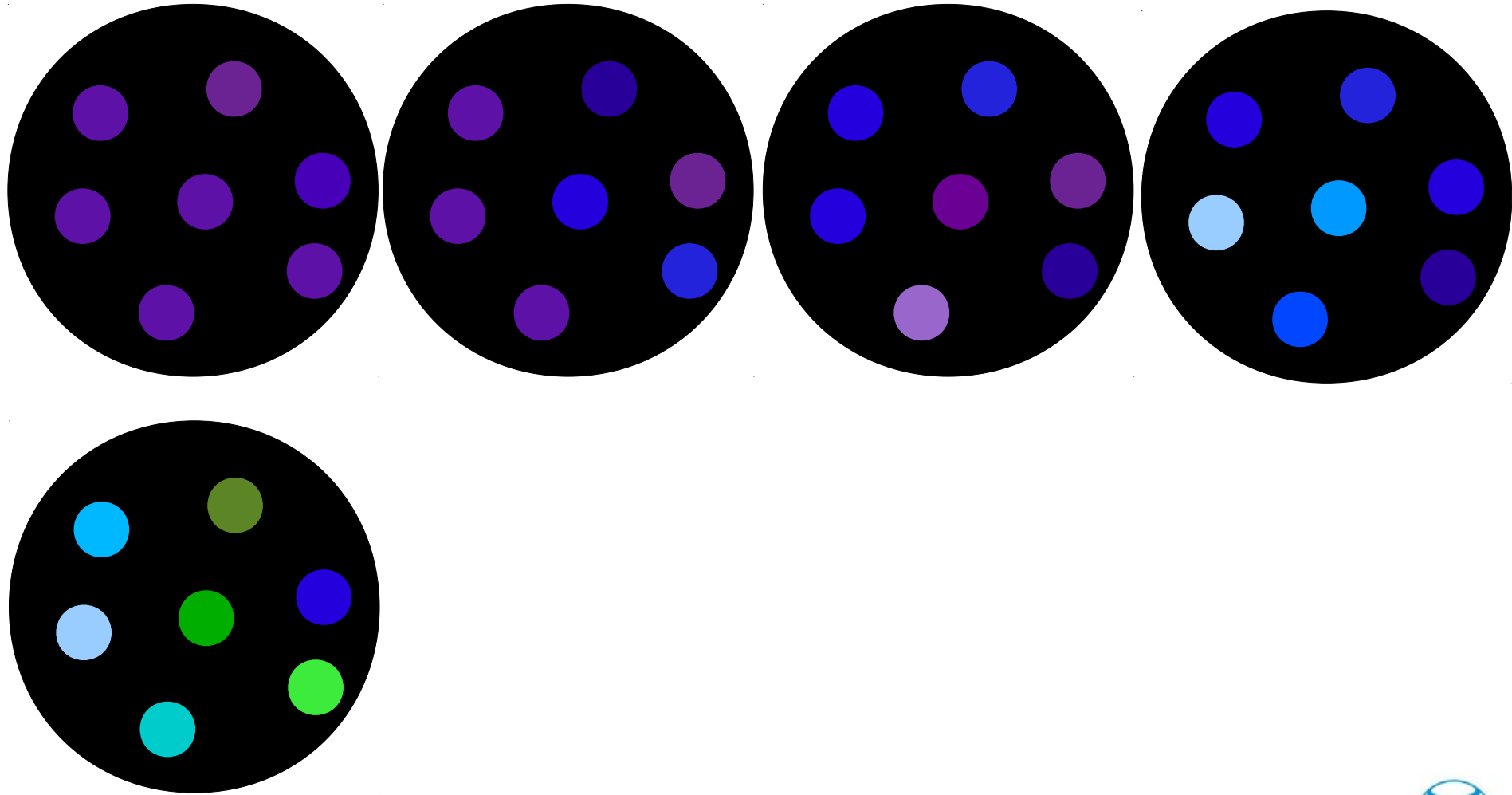
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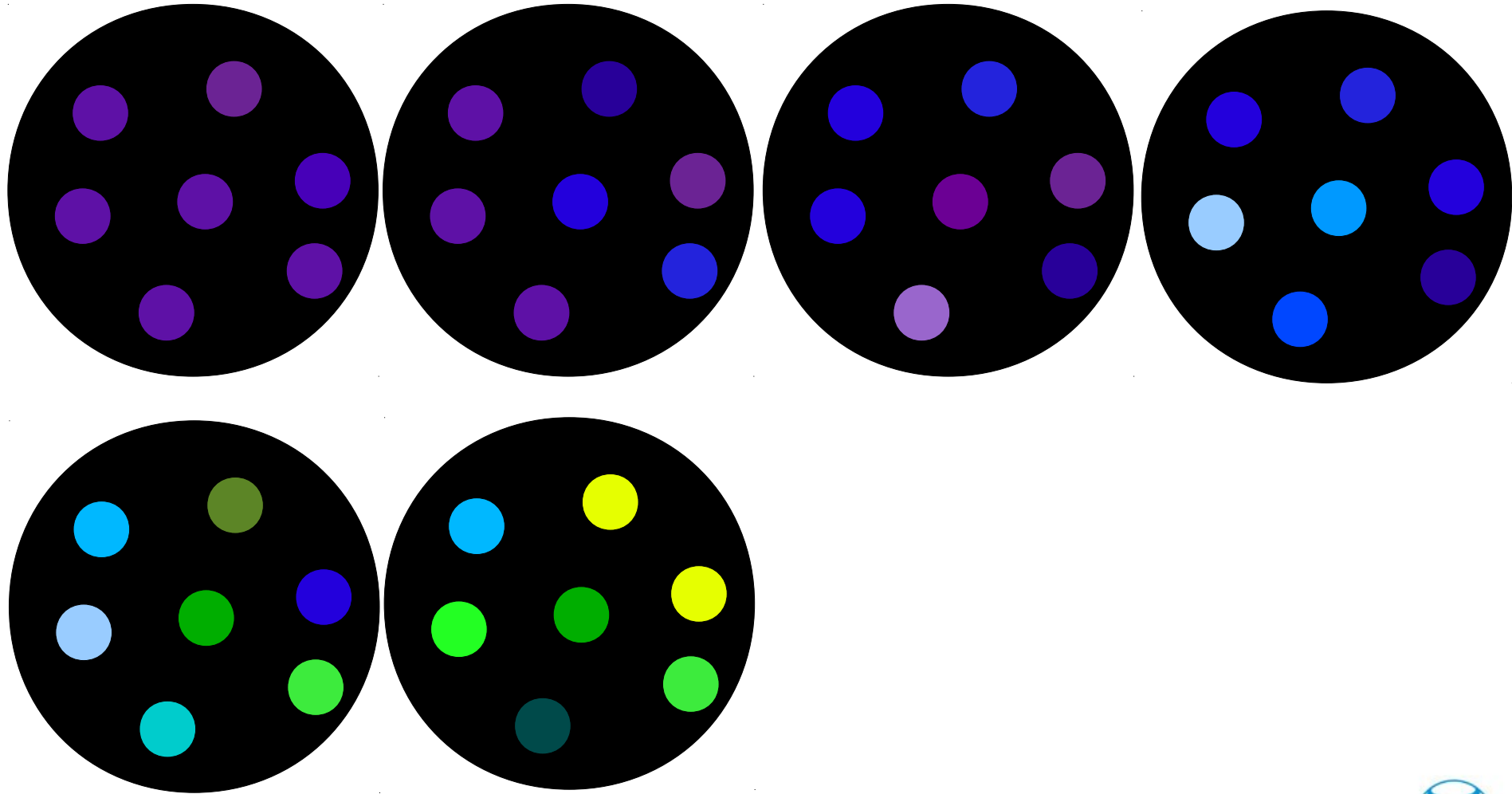
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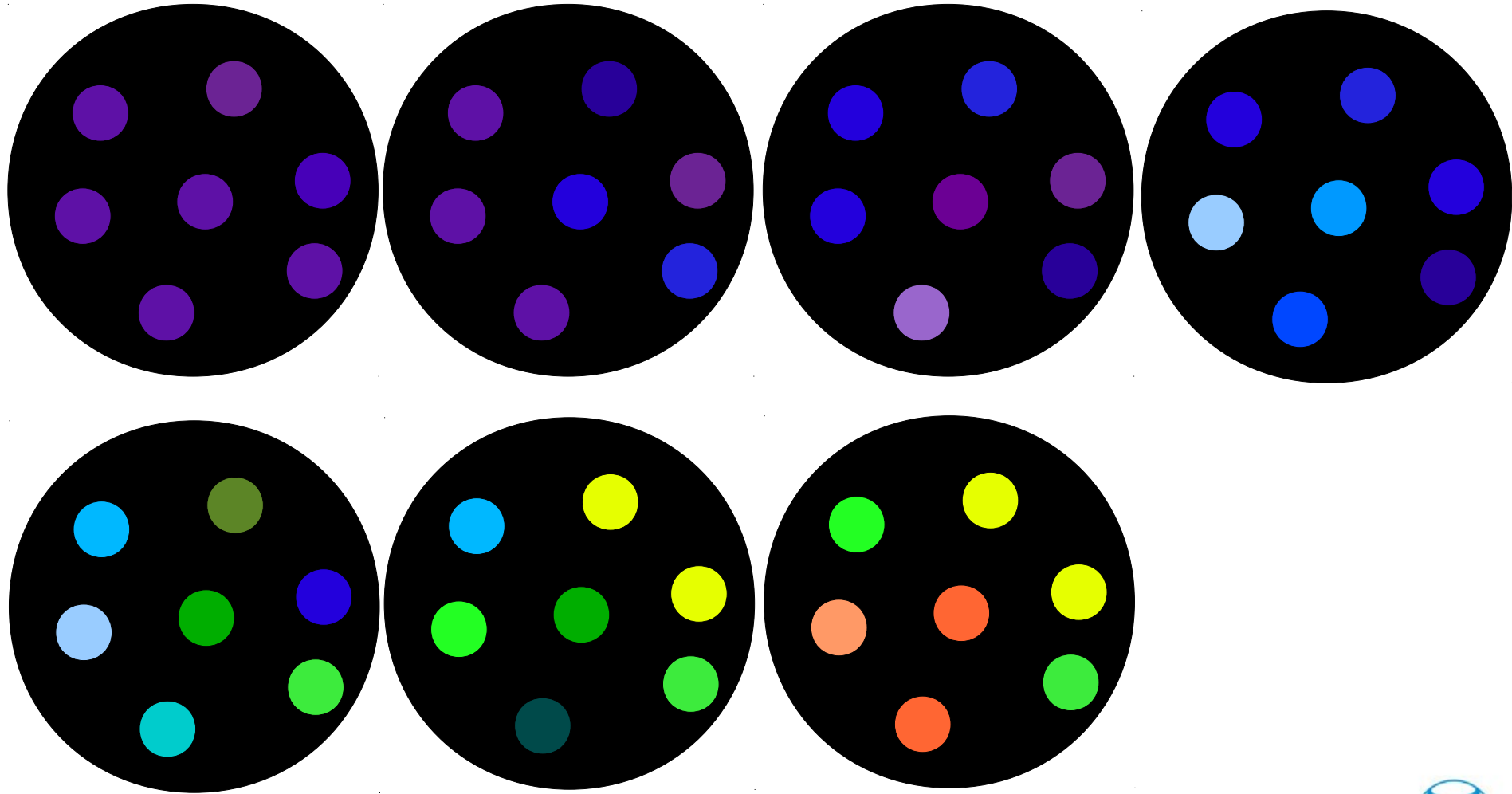
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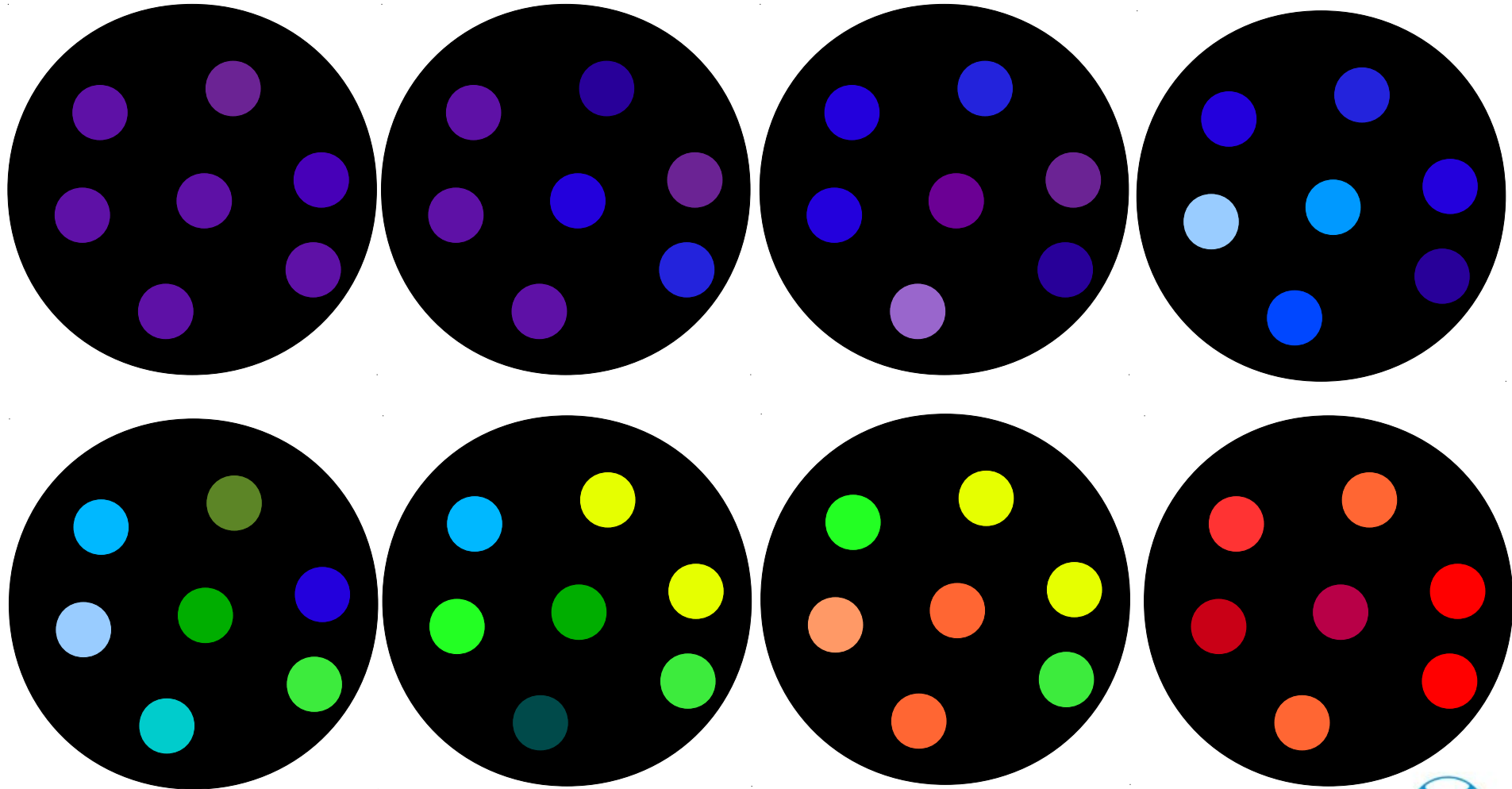


# Evolution of solutions





# Evolution of solutions



# Encoding solutions with random keys

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- A vector  $X$  of random keys, or simply random keys, is an array of  $n$  random keys.
- Solutions of optimization problems can be encoded by random keys.
- A decoder is a deterministic algorithm that takes a vector of random keys as input and outputs a feasible solution of the optimization problem.

# Encoding with random keys: Sequencing

## Encoding

[ 1, 2, 3, 4, 5 ]

$X = [ 0.099, 0.216, 0.802, 0.368, 0.658 ]$

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## Decode by sorting vector of random keys

[ 1, 2, 4, 5, 3 ]

$X = [ 0.099, 0.216, 0.368, 0.658, 0.802 ]$



# Encoding with random keys: Sequencing

Therefore, the vector of random keys:

$$X = [ 0.099, 0.216, 0.802, 0.368, 0.658 ]$$

encodes the sequence: 1 – 2 – 4 – 5 – 3

# Encoding with random keys: Subset selection (select 3 of 5 elements)

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Therefore, the vector of random keys:

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encodes the subset:  $\{1, 2, 4\}$

# Encoding with random keys: Assigning integer weights $\in [0,10]$ to a subset of 3 of 5 elements

## Encoding

[ 1, 2, 3, 4, 5 | 1, 2, 3, 4, 5 ]

$X = [ 0.099, 0.216, 0.802, 0.368, 0.658 \mid 0.4634, 0.5611, 0.2752, 0.4874, 0.0348 ]$

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Decode by sorting the first 5 keys and assign as the weight the value

$W_i = \mathbf{floor} [ 10 X_{5+i} ] + 1$  to the 3 elements with smallest keys  $X_i$ , for  $i = 1, \dots, 5$ .

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Therefore, the vector of random keys:

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encodes the weight vector  $W = (5, 6, -, 5, -)$

# Genetic algorithms and random keys



# GAs and random keys

- Introduced by Bean (1994) for sequencing problems.

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- Individuals are strings of real-valued numbers (random keys) in the interval  $[0,1)$ .

$$S = ( 0.25, 0.19, 0.67, 0.05, 0.89 )$$

$s(1) \quad s(2) \quad s(3) \quad s(4) \quad s(5)$

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- Individuals are strings of real-valued numbers (random keys) in the interval  $[0,1)$ .
- Sorting random keys results in a sequencing order.

$$S = ( \begin{matrix} 0.25, & 0.19, & 0.67, & 0.05, & 0.89 \end{matrix} ) \\ \begin{matrix} s(1) & s(2) & s(3) & s(4) & s(5) \end{matrix}$$

$$S' = ( \begin{matrix} 0.05, & 0.19, & 0.25, & 0.67, & 0.89 \end{matrix} ) \\ \begin{matrix} s(4) & s(2) & s(1) & s(3) & s(5) \end{matrix}$$

Sequence: 4 – 2 – 1 – 3 – 5

# GAs and random keys

- Mating is done using parametrized uniform crossover (Spears & DeJong , 1990)

$a = ( 0.25, 0.19, 0.67, 0.05, 0.89 )$   
 $b = ( 0.63, 0.90, 0.76, 0.93, 0.08 )$

# GAs and random keys

- Mating is done using parametrized uniform crossover (Spears & DeJong , 1990)
- 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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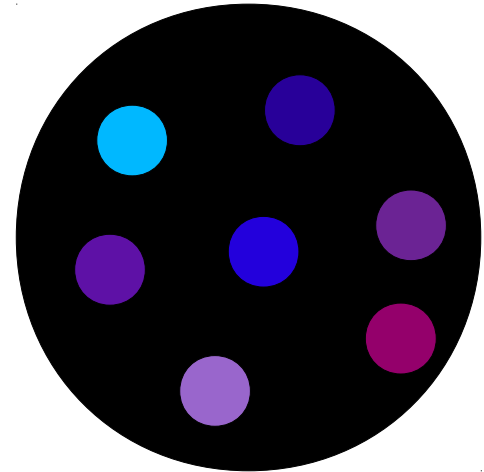
$b = ( 0.63, 0.90, 0.76, 0.93, 0.08 )$

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

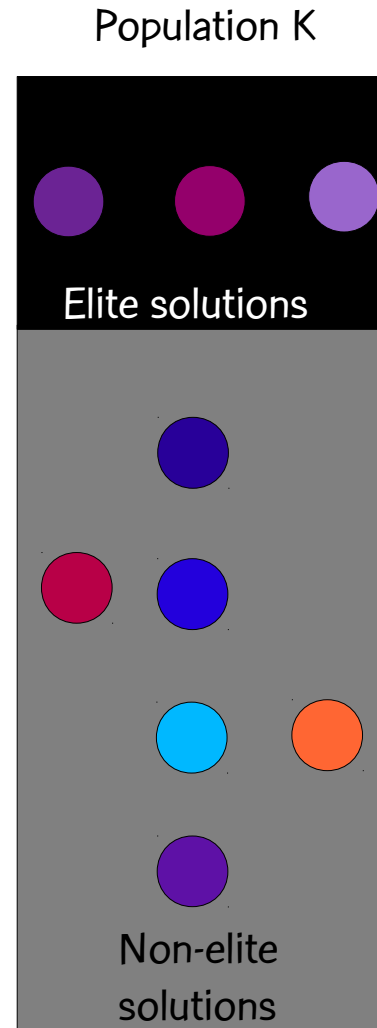
# GAs and random keys

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  $[0,1)$ .



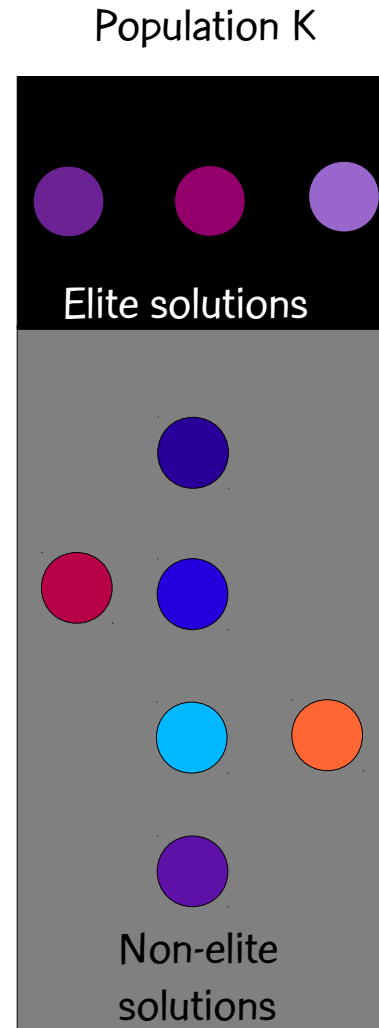
# GAs and random keys

At the K-th generation,  
compute the cost of each  
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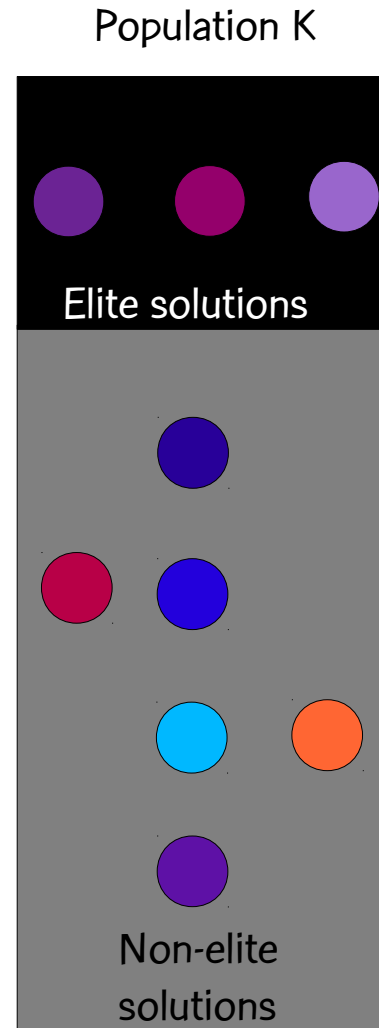
# GAs and random keys

At the K-th generation,  
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# GAs and random keys

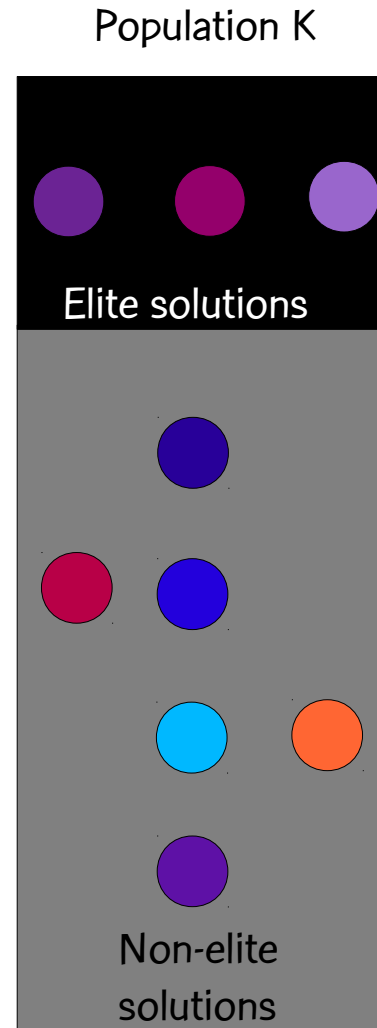
At the K-th generation,  
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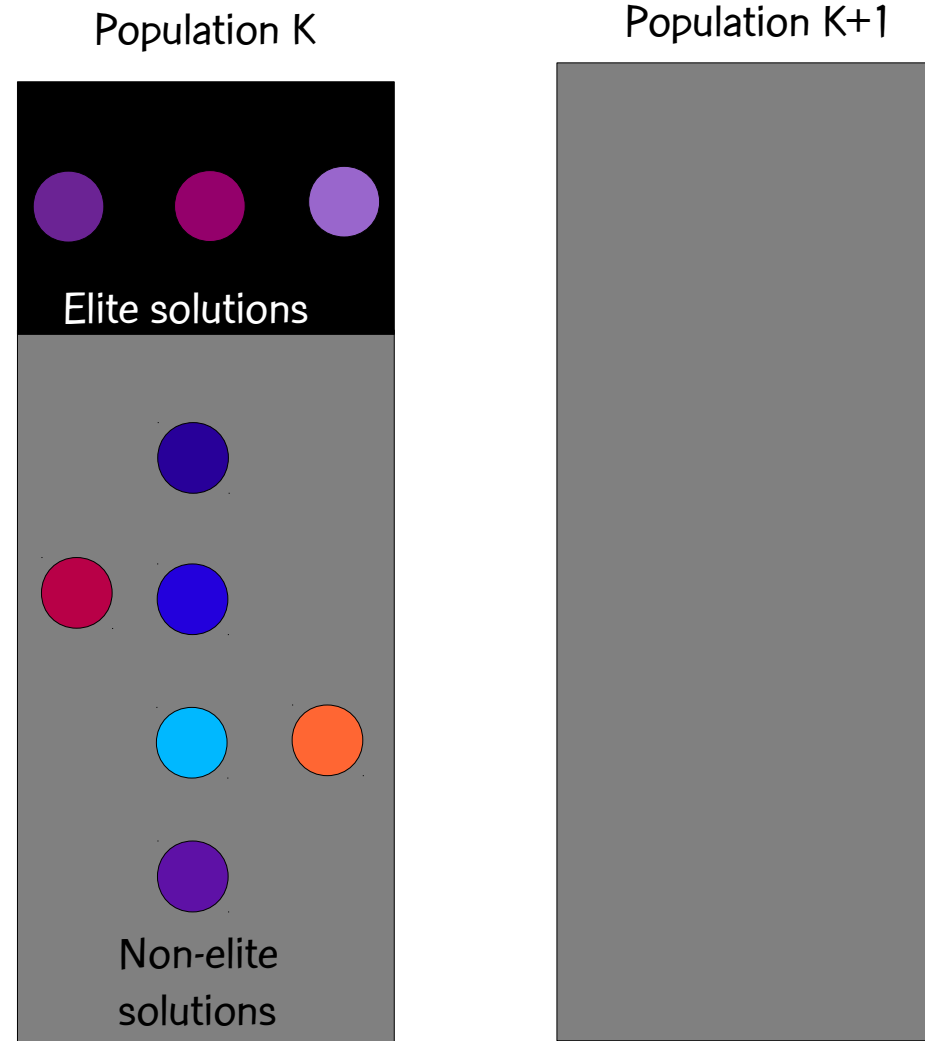
# GAs and random keys

At the K-th generation, compute the cost of each solution and partition the solutions into two sets: elite solutions and non-elite solutions. Elite set should be smaller of the two sets and contain best solutions.



# GAs and random keys

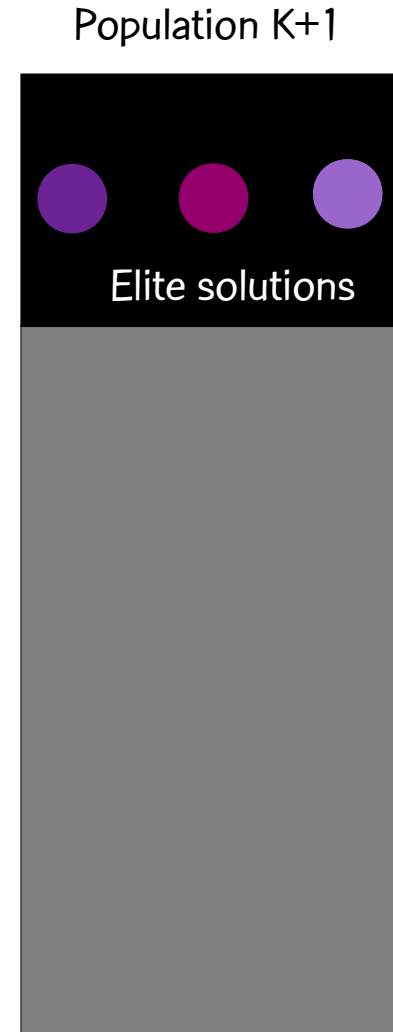
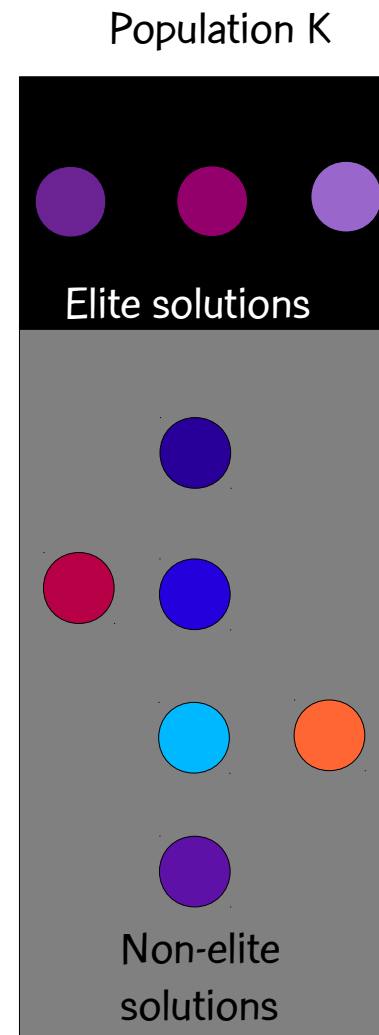
## Evolutionary dynamics



# GAs and random keys

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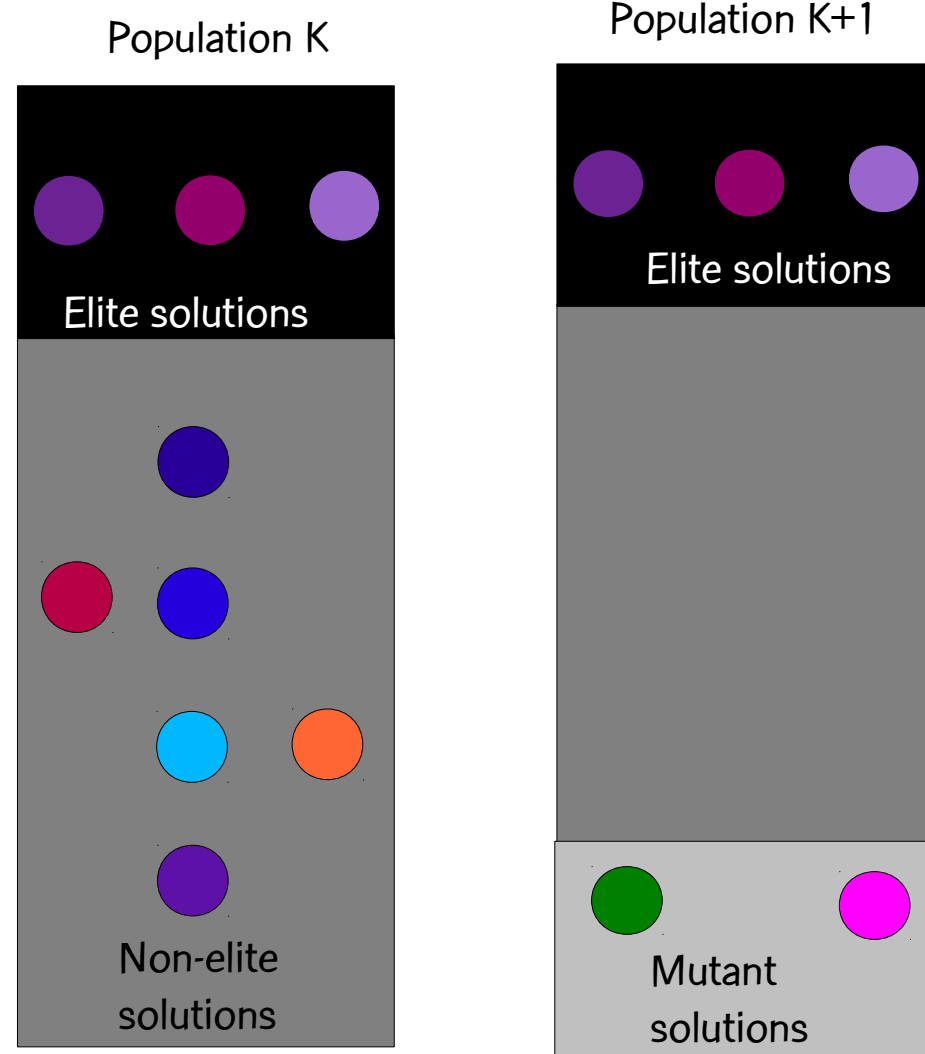
- Copy elite solutions from population K to population K+1



# GAs and random keys

## Evolutionary dynamics

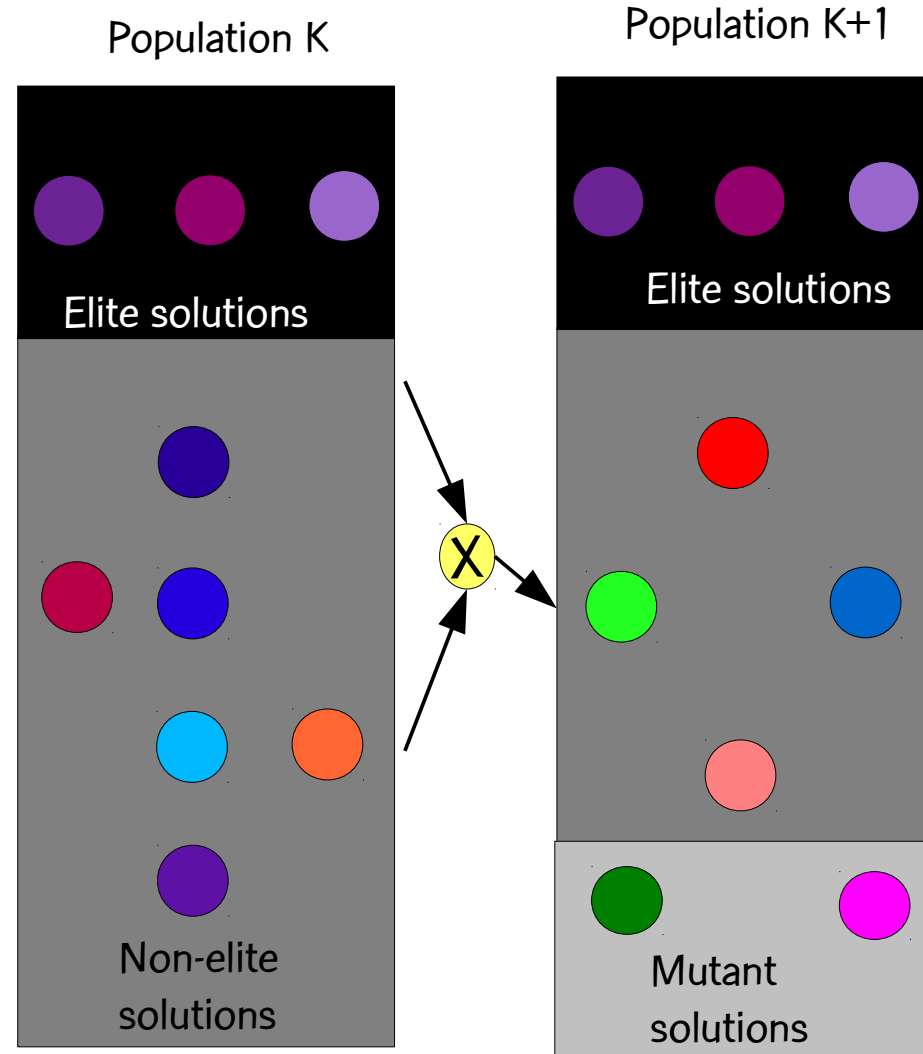
- Copy elite solutions from population K to population K+1
- Add R random solutions (mutants) to population K+1



# GAs and random keys

## Evolutionary dynamics

- Copy elite solutions from population K to population K+1
- Add R random solutions (mutants) to population K+1
- While K+1-th population  $< P$ 
  - **RANDOM-KEY GA:** Use any two solutions in population K to produce child in population K+1. Mates are chosen at random.



BRKGA

# Biased random key genetic algorithm

- A biased random key genetic algorithm (BRKGA) is a random key genetic algorithm (RKGA).

# Biased random key genetic algorithm

- 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 parametrized uniform crossover is applied.

# How RKGA & BRKGA differ

## RKGA

both parents chosen at  
random from entire  
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## BRKGA



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both parents chosen at random but one parent chosen from population of elite solutions

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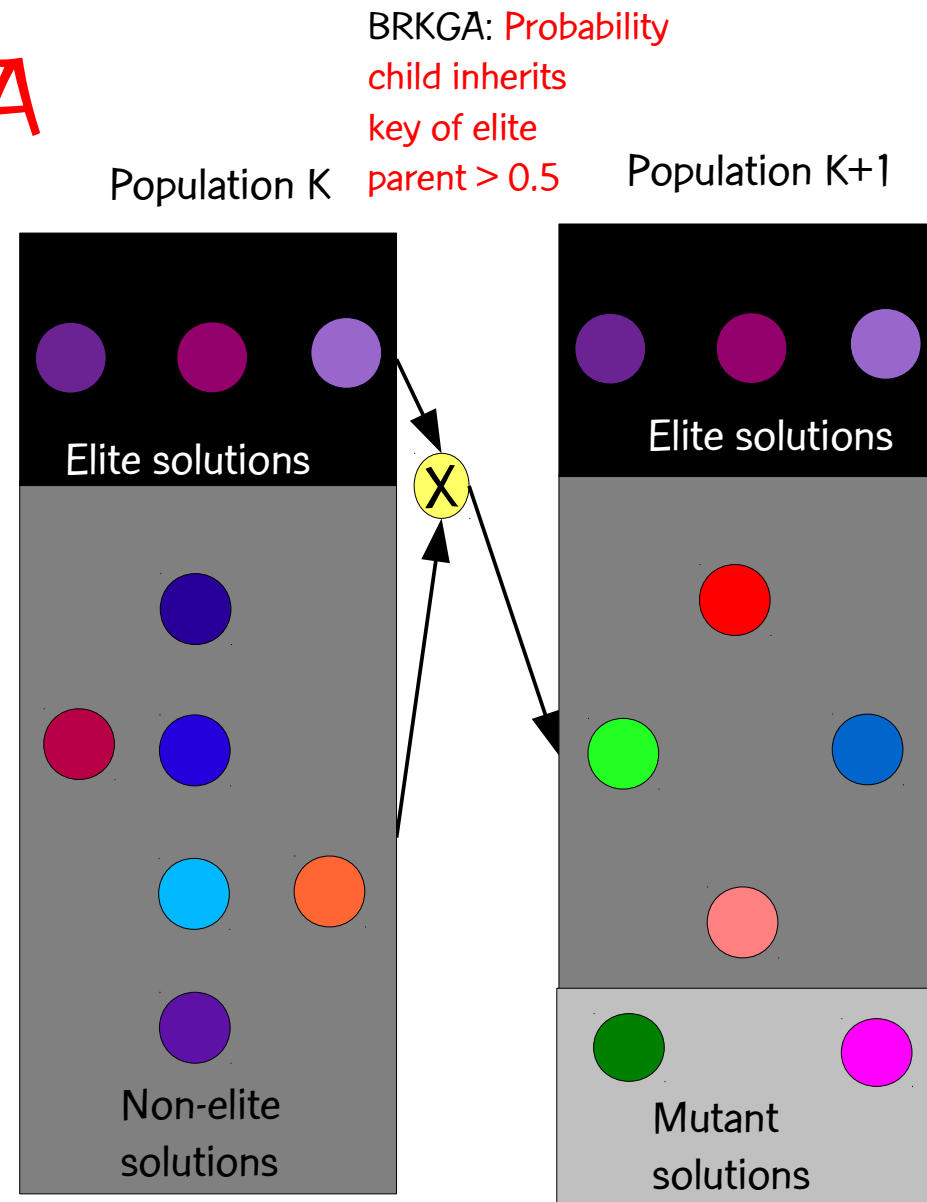
both parents chosen at random but one parent chosen from population of elite solutions

best fit parent is parent A in parametrized uniform crossover

# Biased random key GA

## Evolutionary dynamics

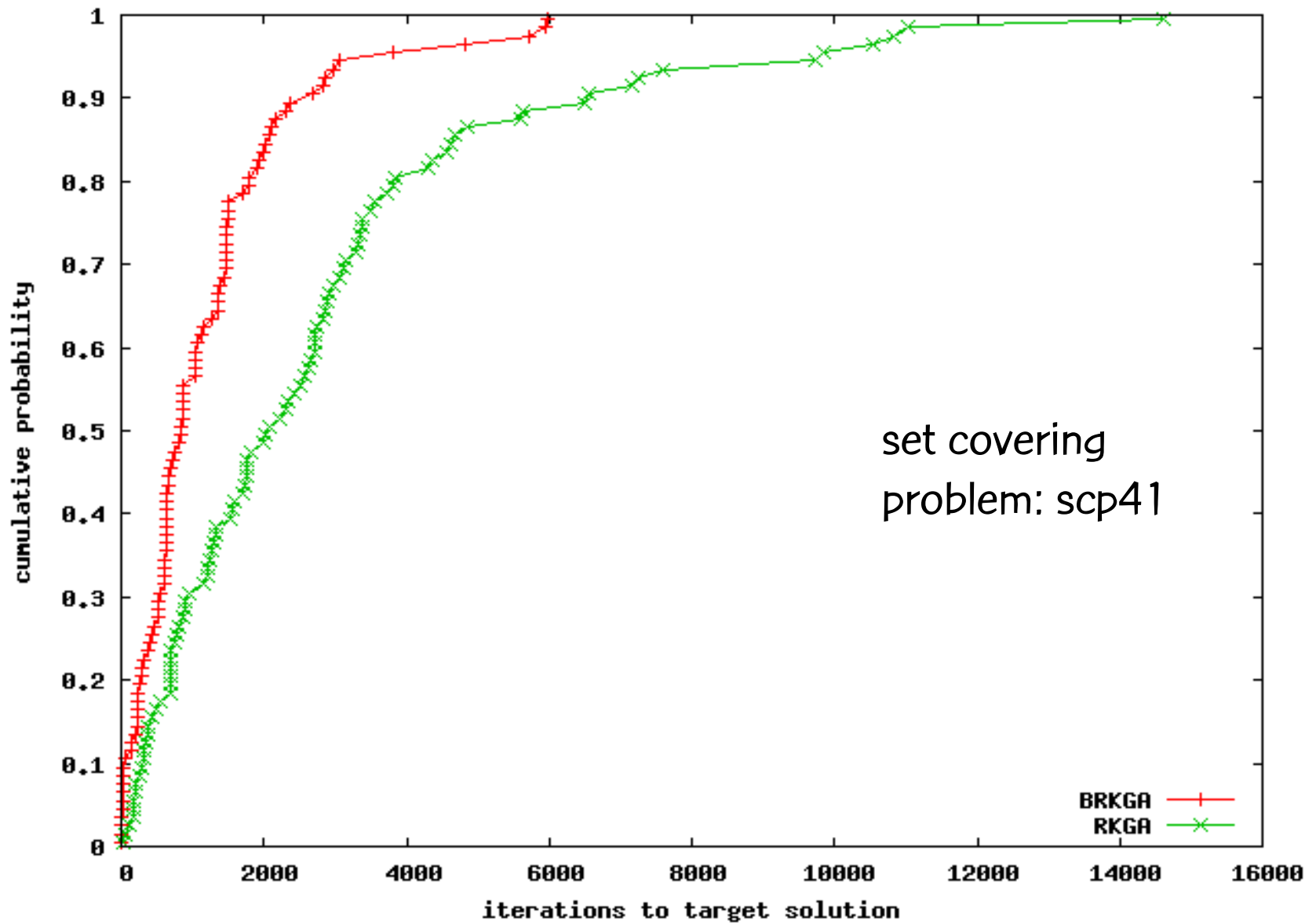
- Copy elite solutions from population K to population K+1
- Add R random solutions (mutants) to population K+1
- While K+1-th population  $< P$ 
  - **RANDOM-KEY GA:** Use any two solutions in population K to produce child in population K+1. Mates are chosen at random.
  - **BIASED RANDOM-KEY GA:** Mate elite solution with other solution of population K to produce child in population K+1. Mates are chosen at random.

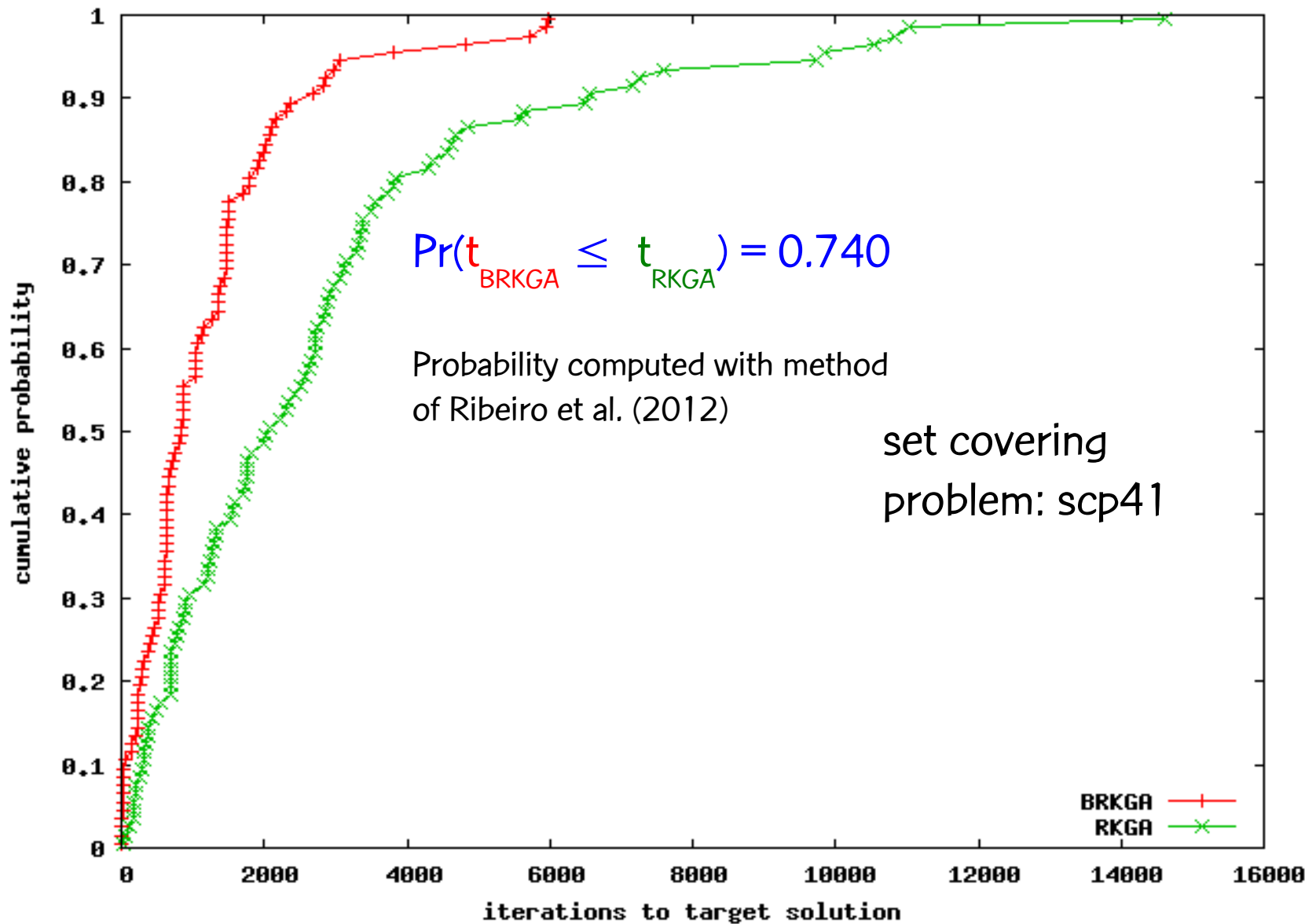


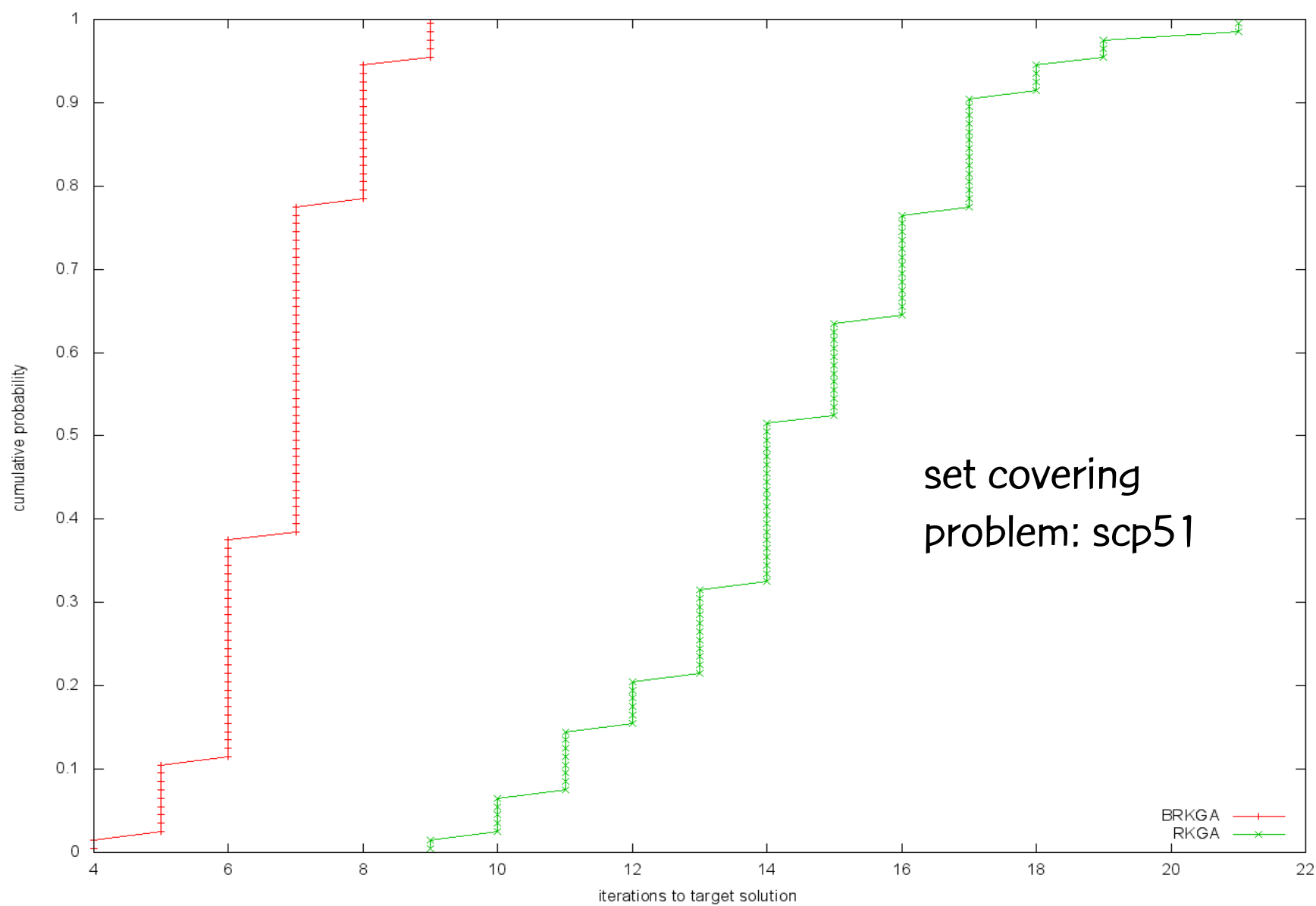
BRKGA

# Paper comparing BRKGA and Bean's Method

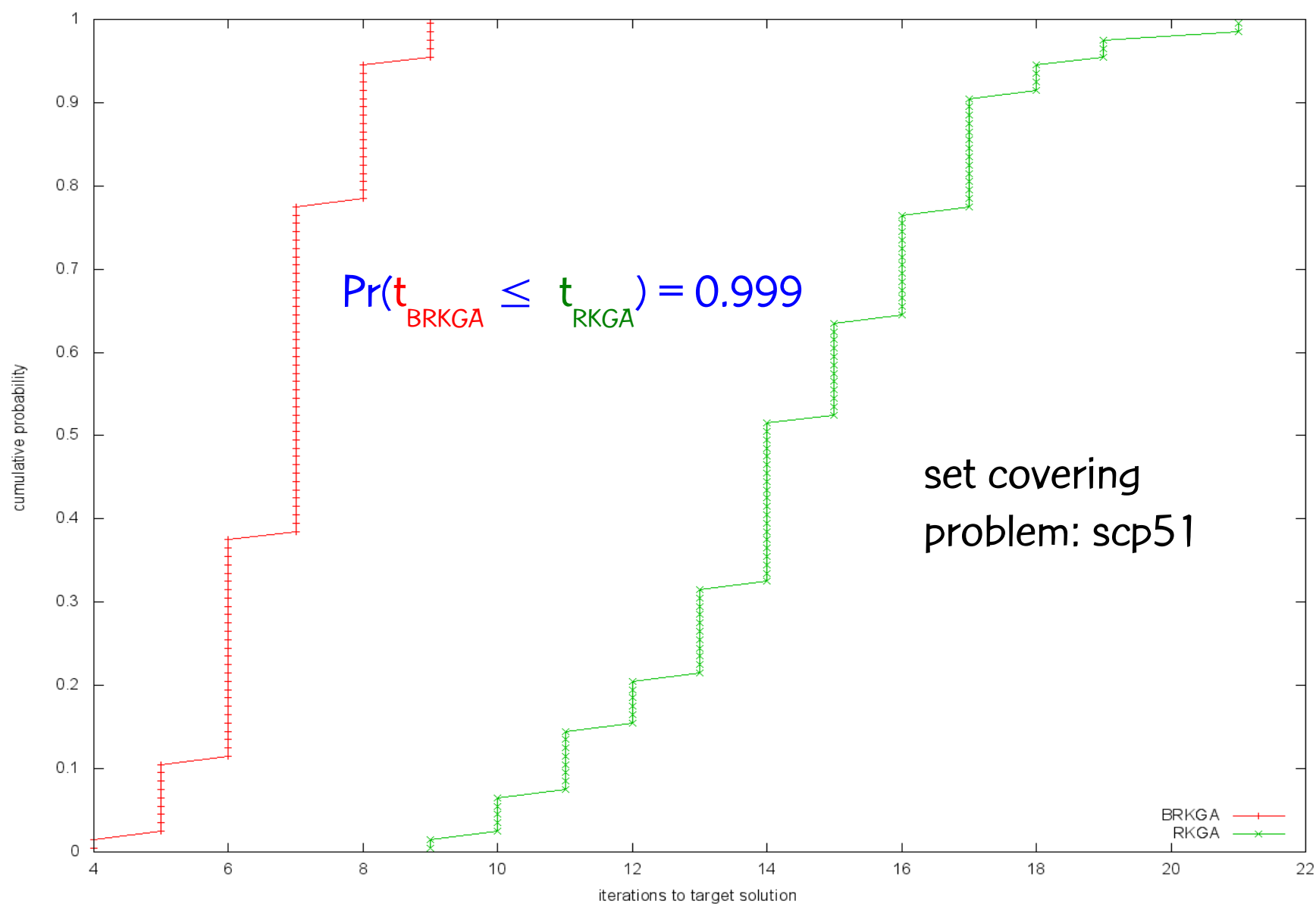
Gonçalves, R., and Toso, “Biased and unbiased random-key genetic algorithms: An experimental analysis”, AT&T Labs Research Technical Report, Florham Park, December 2012.

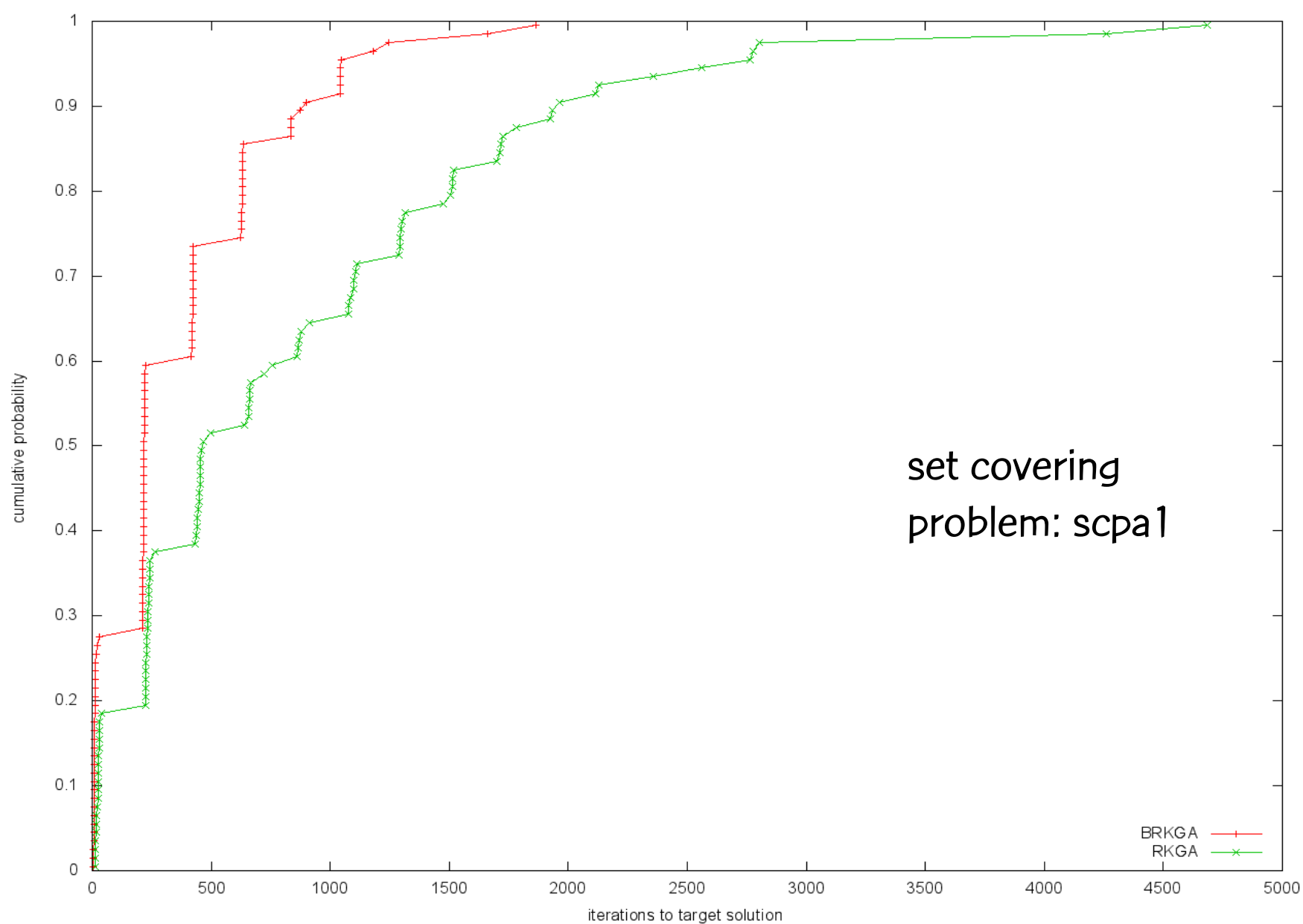


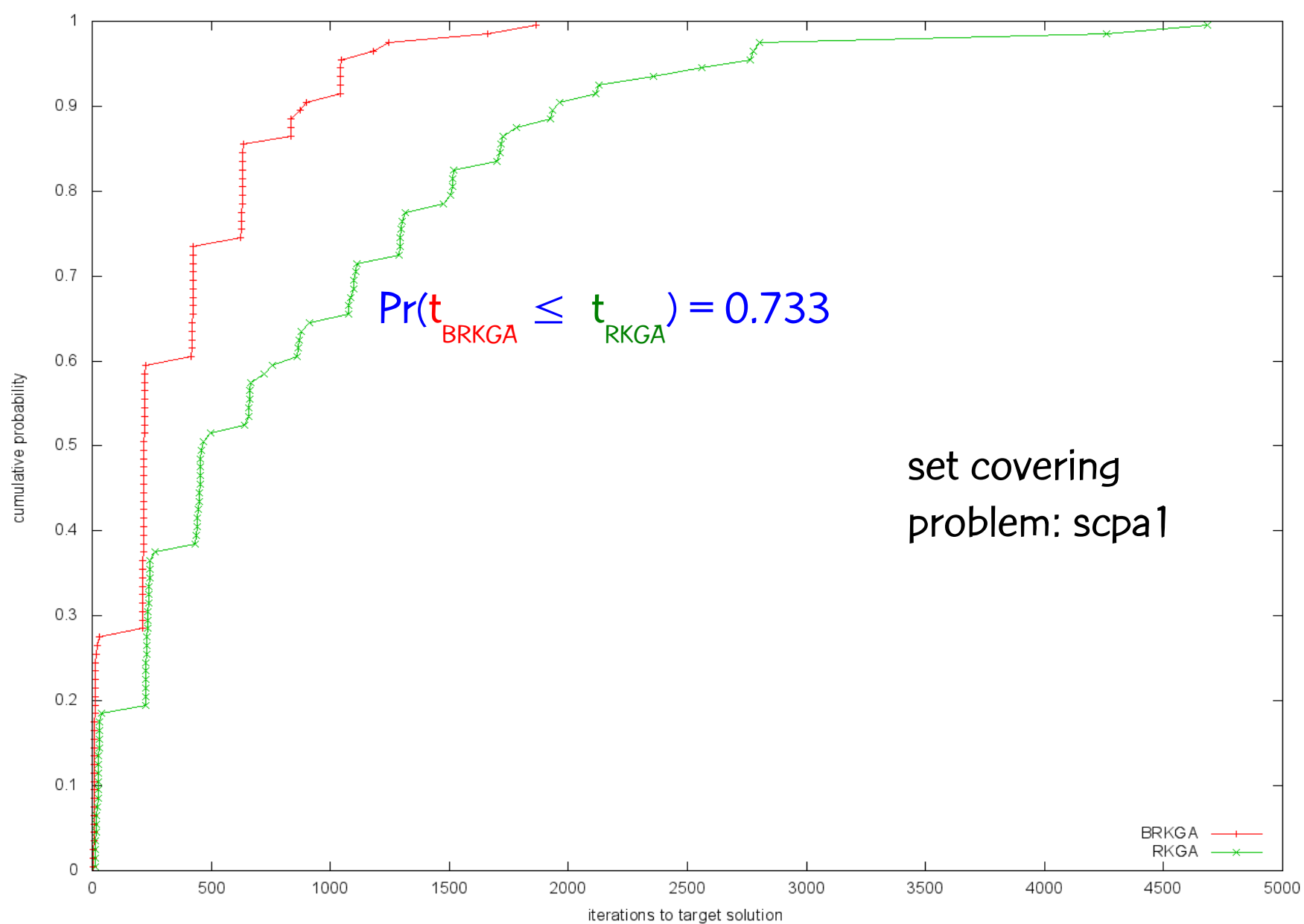


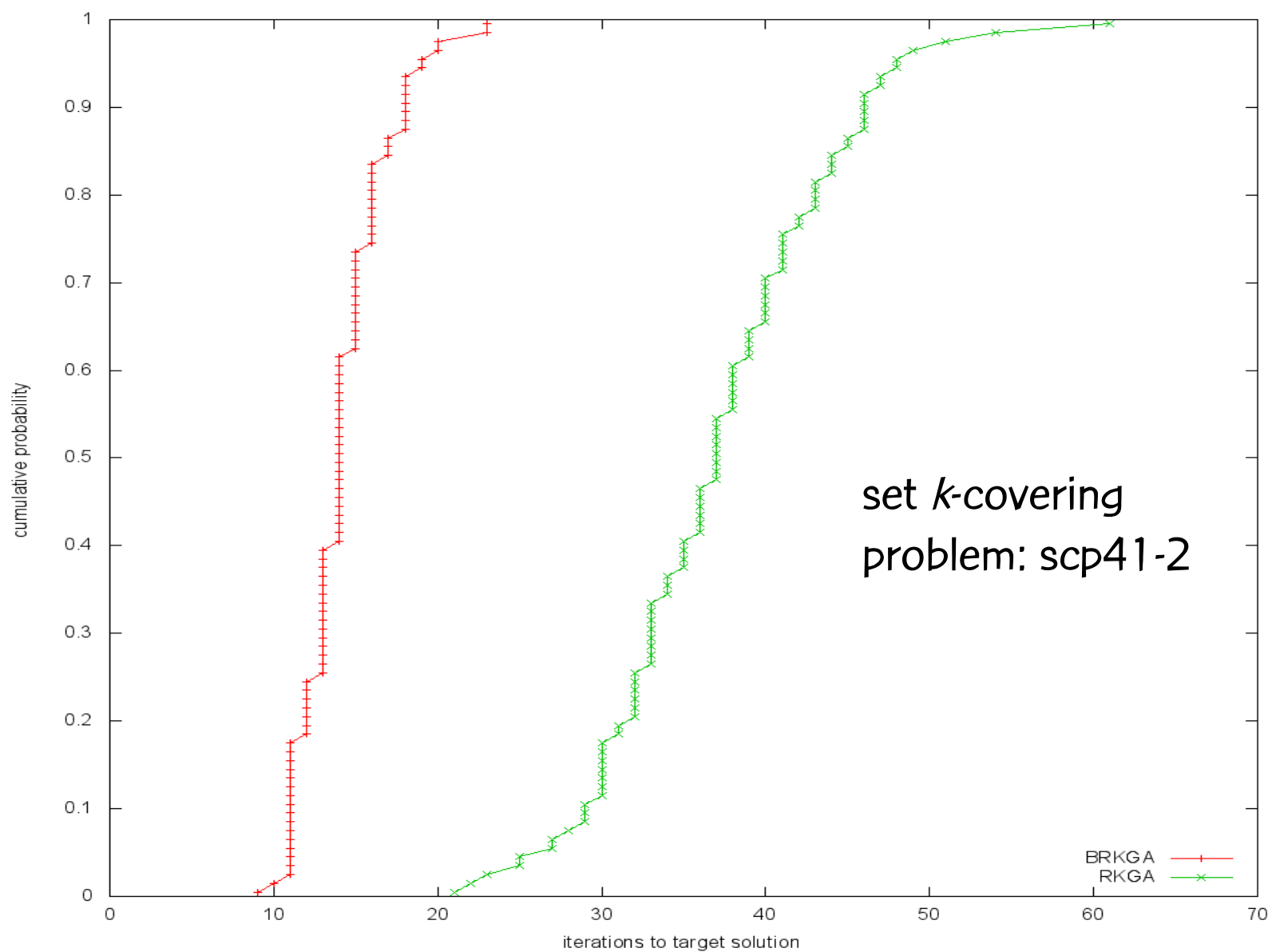


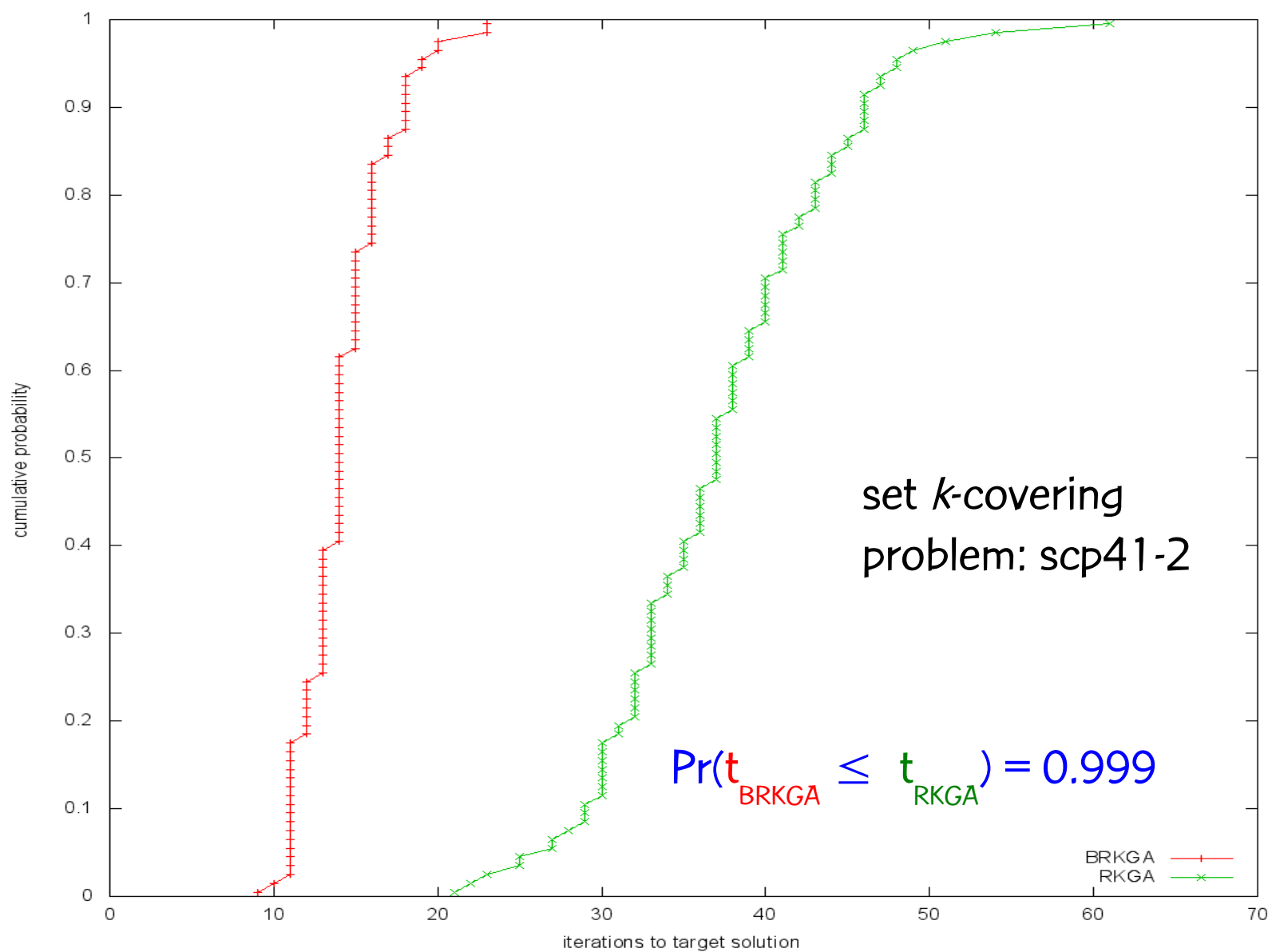


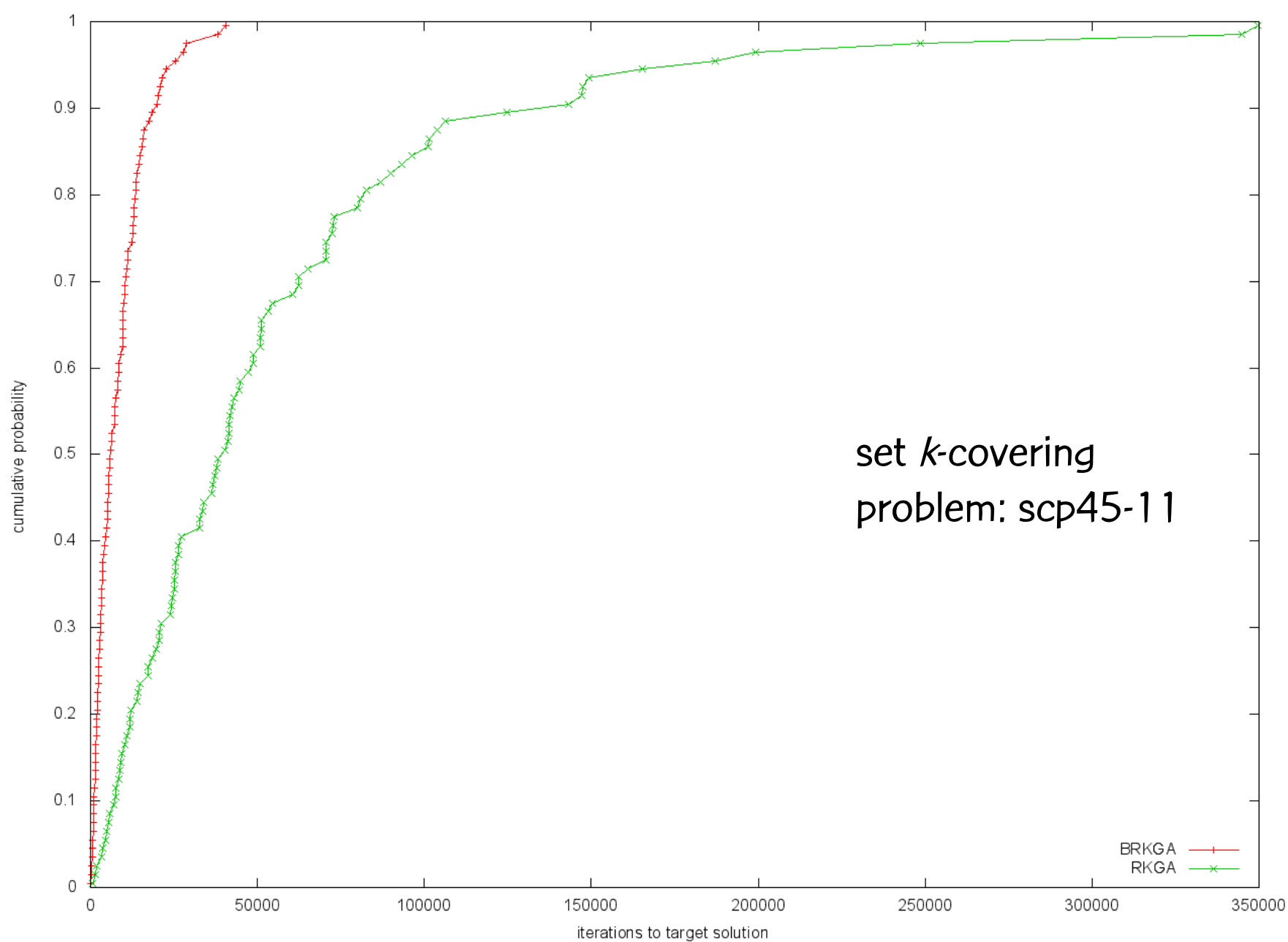


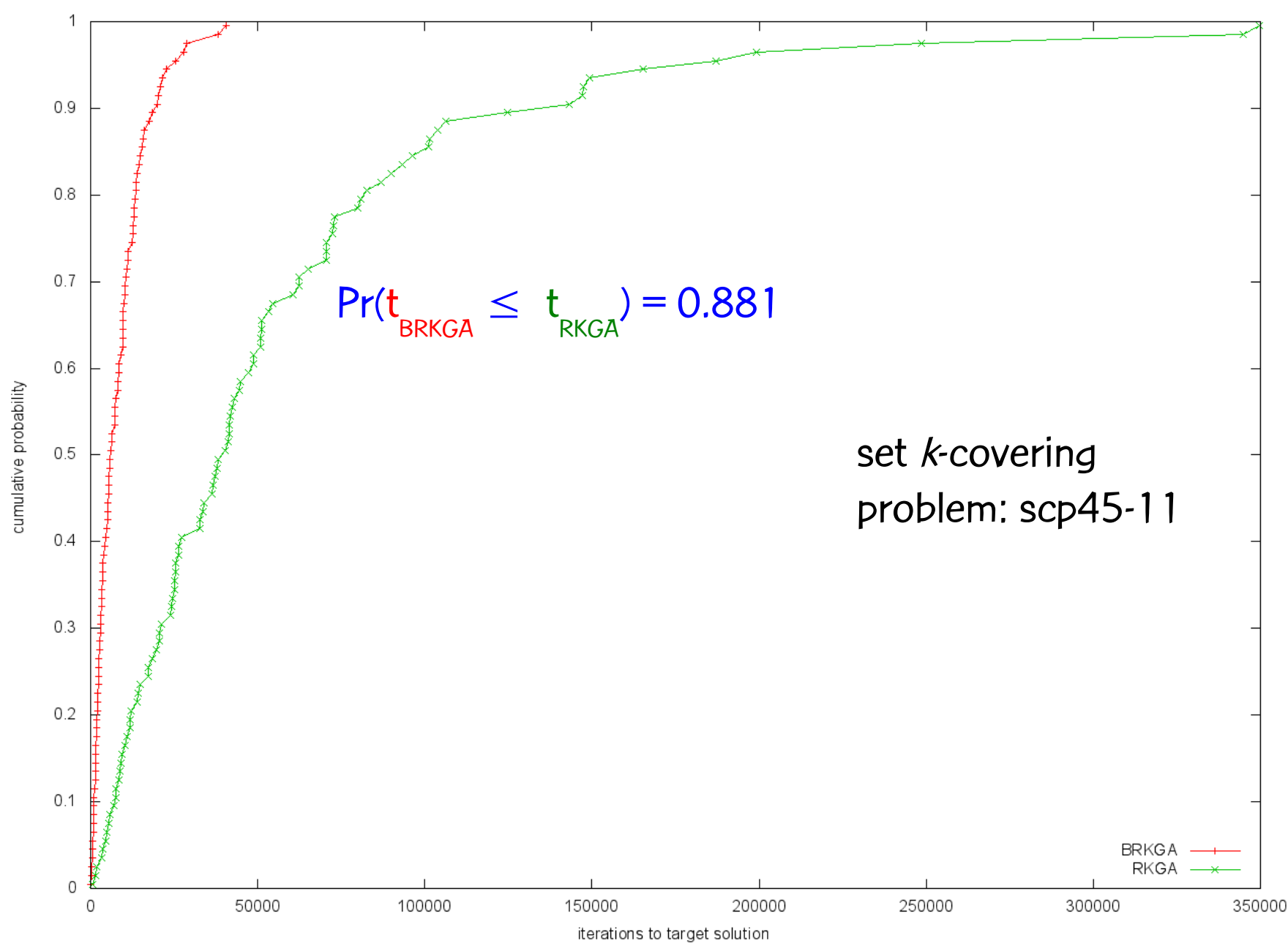


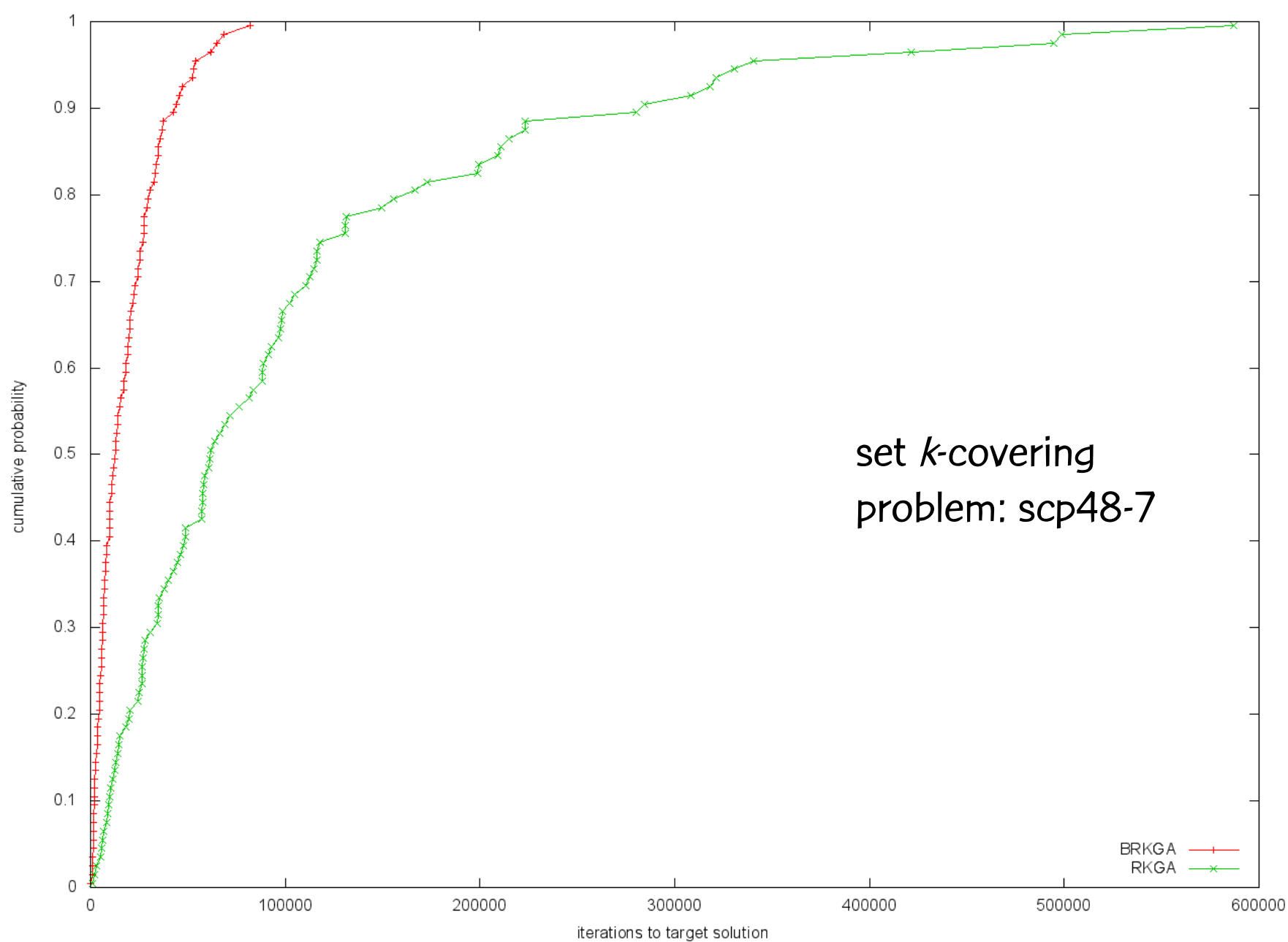




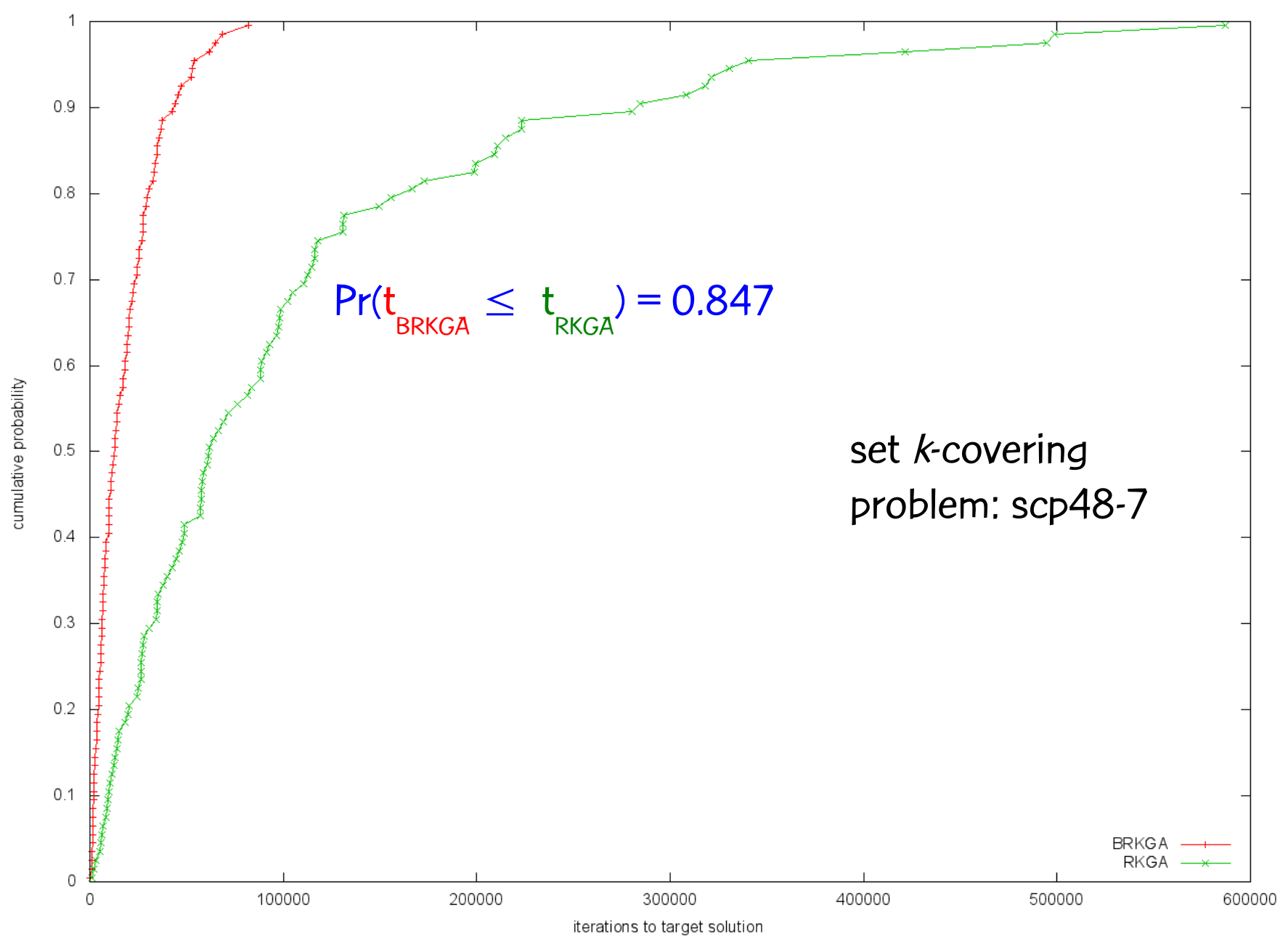












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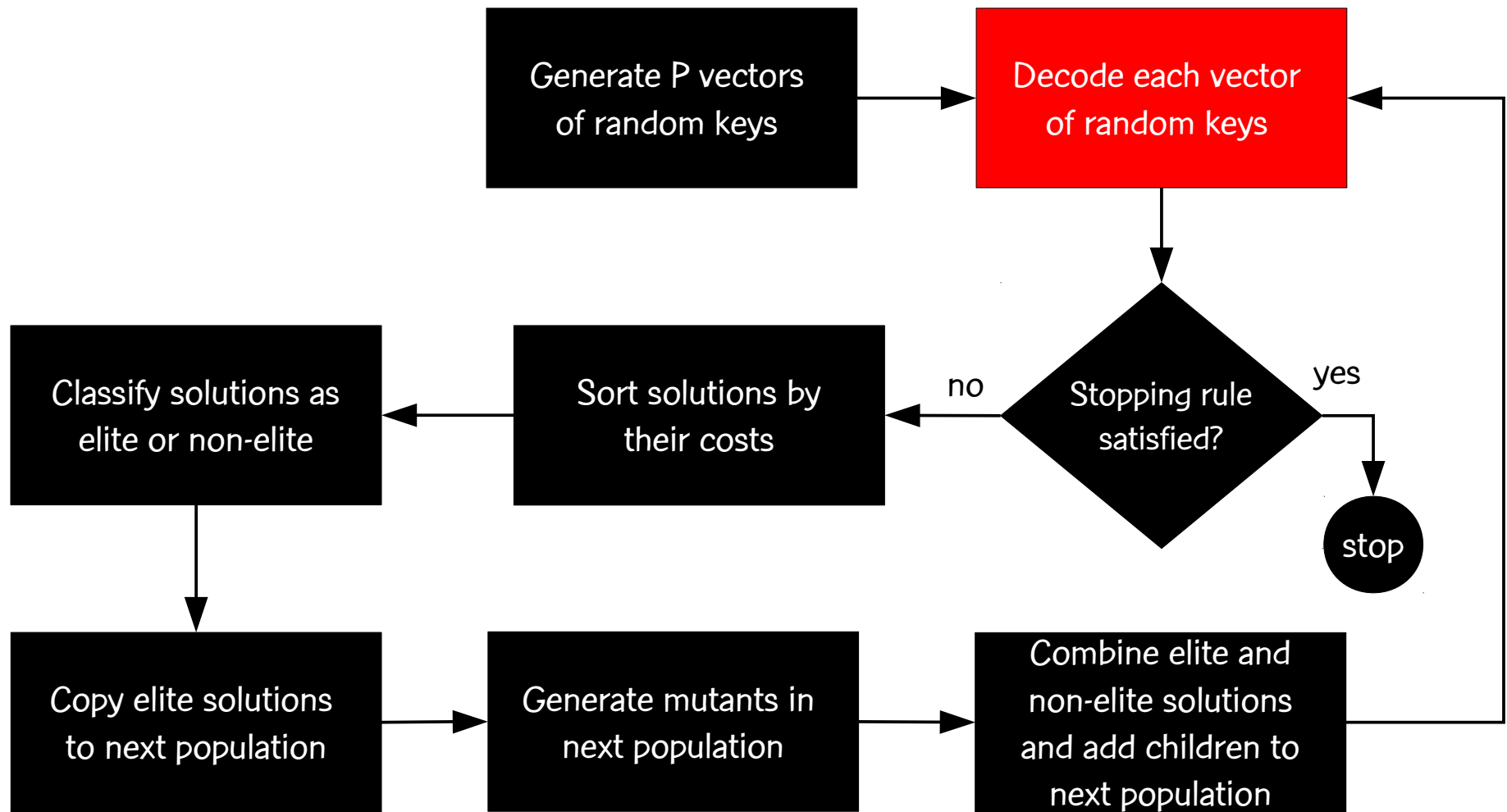
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- Random method: keys are randomly generated so solutions are always vectors of random keys
- Elitist strategy: best solutions are passed without change from one generation to the next (incumbent is kept)
- Child inherits more characteristics of elite parent: one parent is always selected (with replacement) from the small elite set and probability that child inherits key of elite parent  $> 0.5$  Not so in the RKGA of Bean.

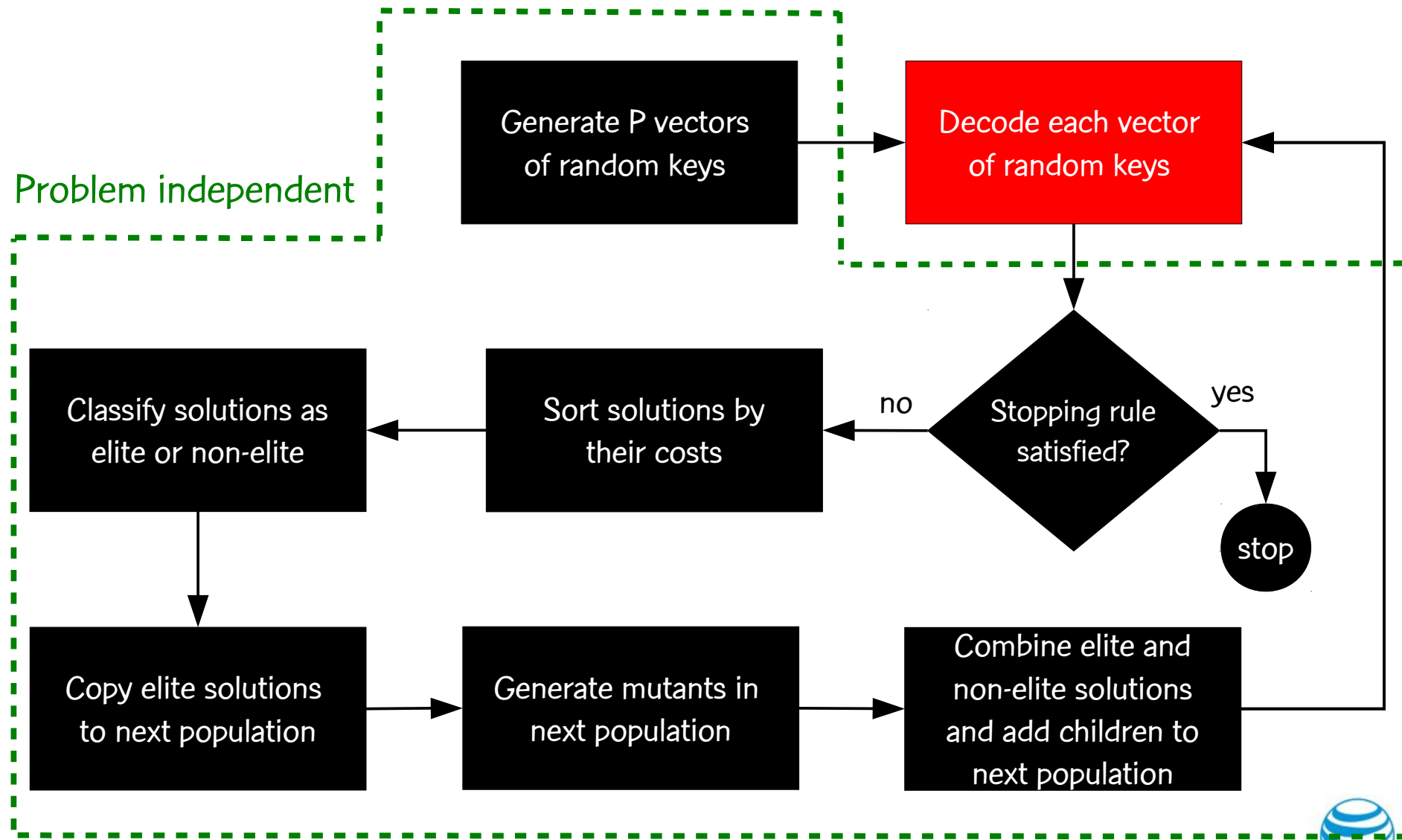
# Observations

- Random method: keys are randomly generated so solutions are always vectors of random keys
- Elitist strategy: best solutions are passed without change from one generation to the next (incumbent is kept)
- Child inherits more characteristics of elite parent: one parent is always selected (with replacement) from the small elite set and probability that child inherits key of elite parent  $> 0.5$  Not so in the RKGA of Bean.
- No mutation in crossover: mutants are used instead (they play same role as mutation in GAs ... help escape local optima)

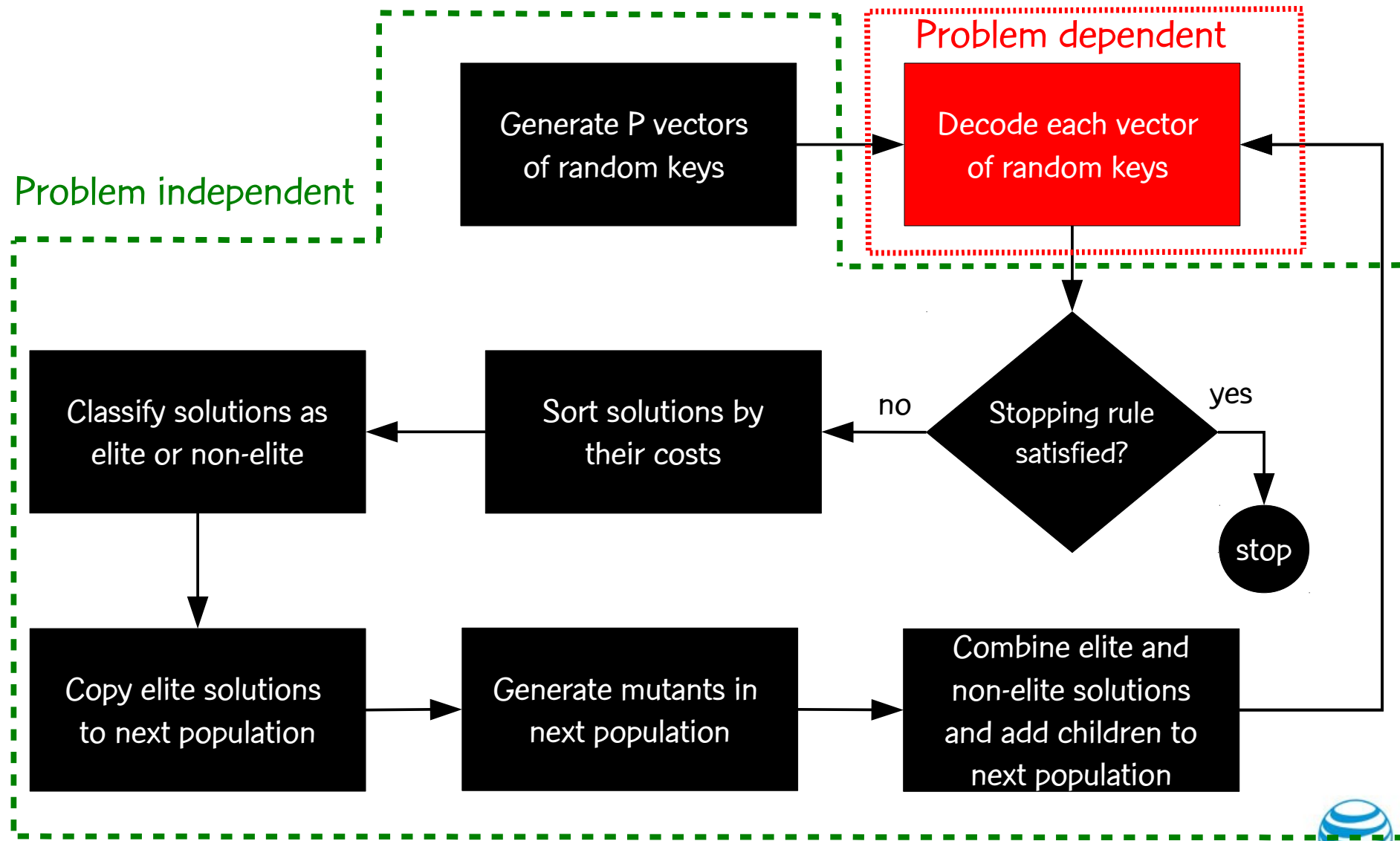
# Framework for biased random-key genetic algorithms



# Framework for biased random-key genetic algorithms

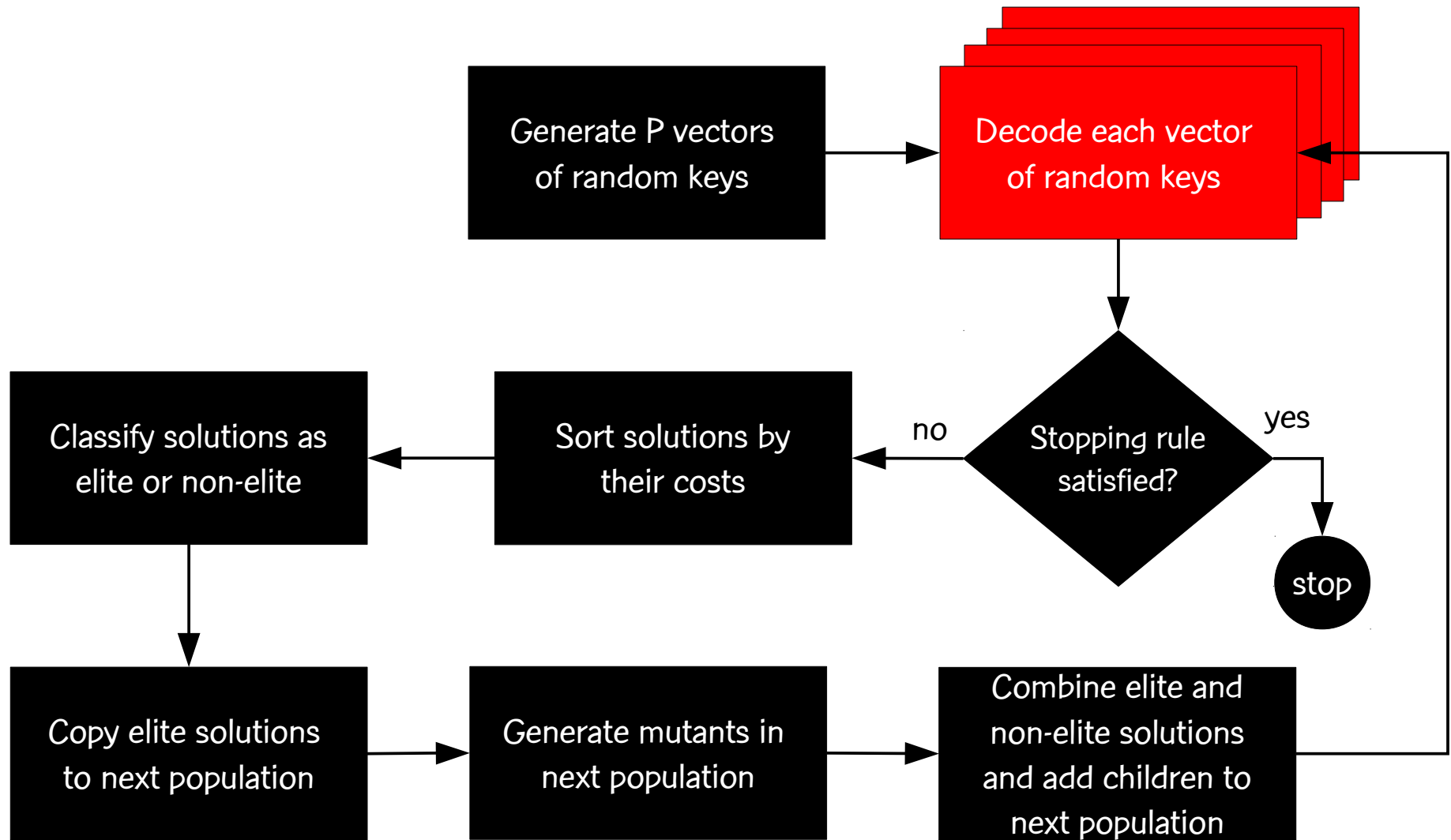


# Framework for biased random-key genetic algorithms





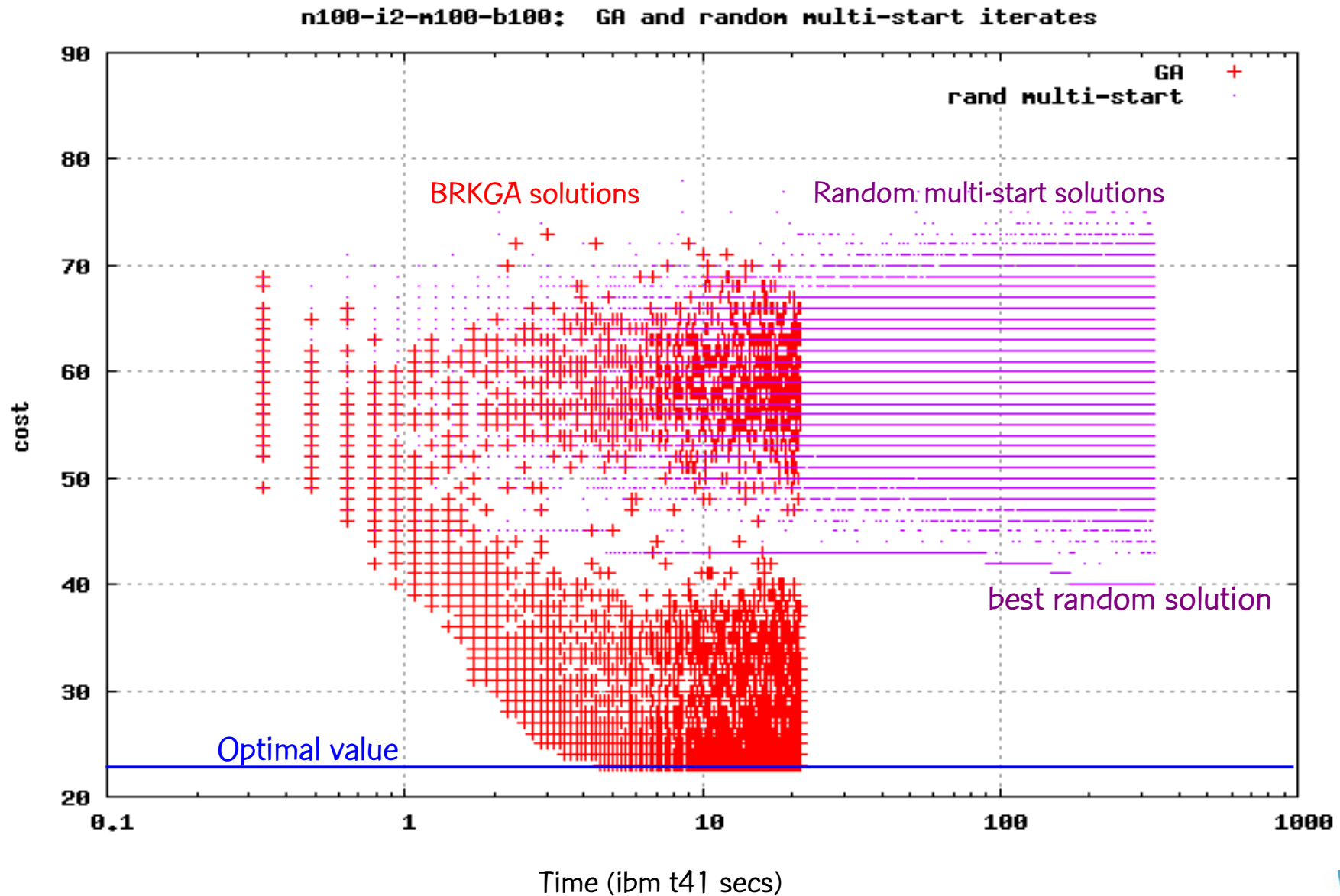
# Decoding of random key vectors can be done in parallel



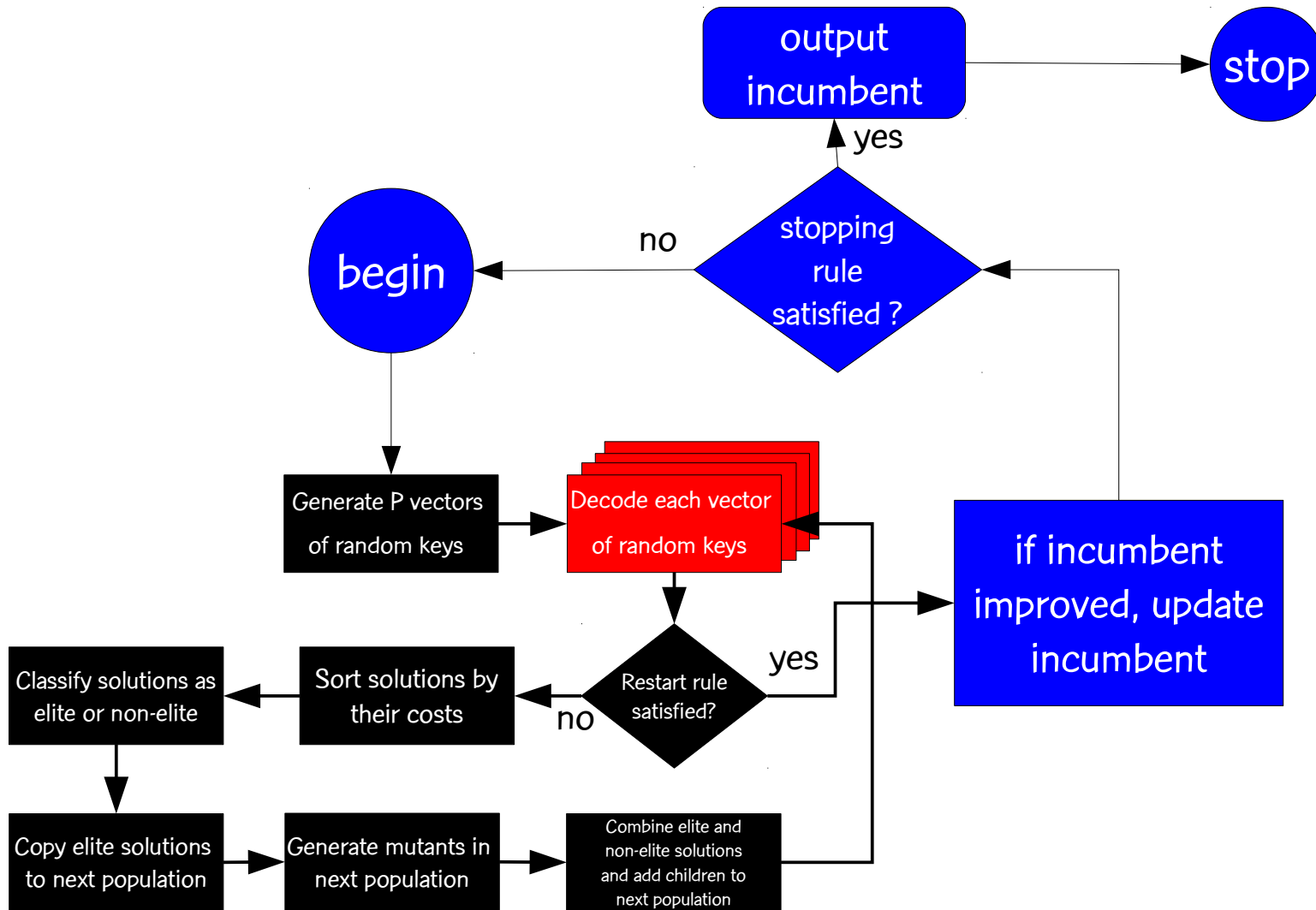
# Is a BRKGA any different from applying the decoder to random keys?

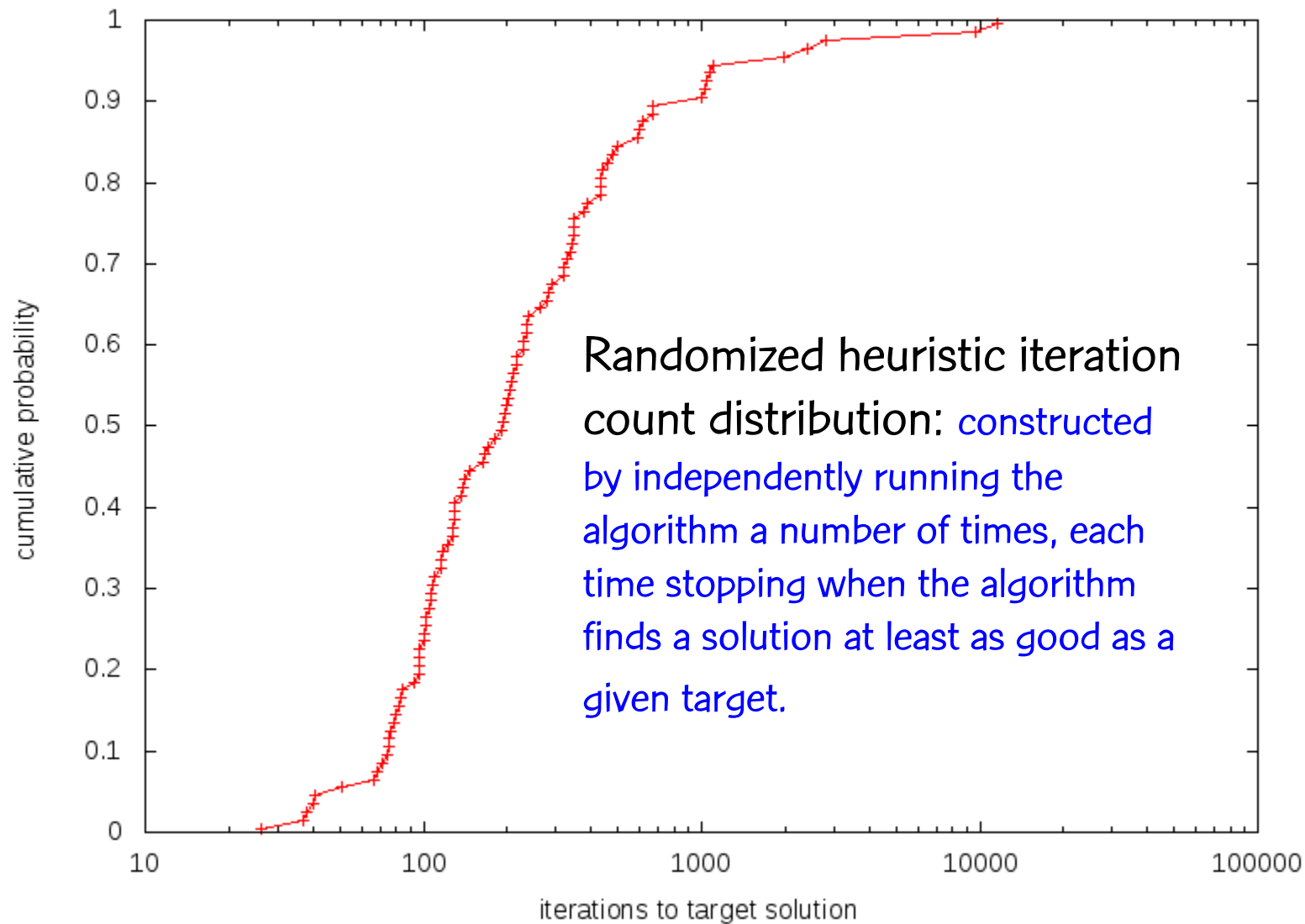
- Simulate a random multi-start decoding method with a BRKGA by setting size of elite partition to 1 and number of mutants to  $P-1$
- Each iteration, best solution is maintained in elite set and  $P-1$  random key vectors are generated as mutants ... no mating is done since population already has  $P$  individuals

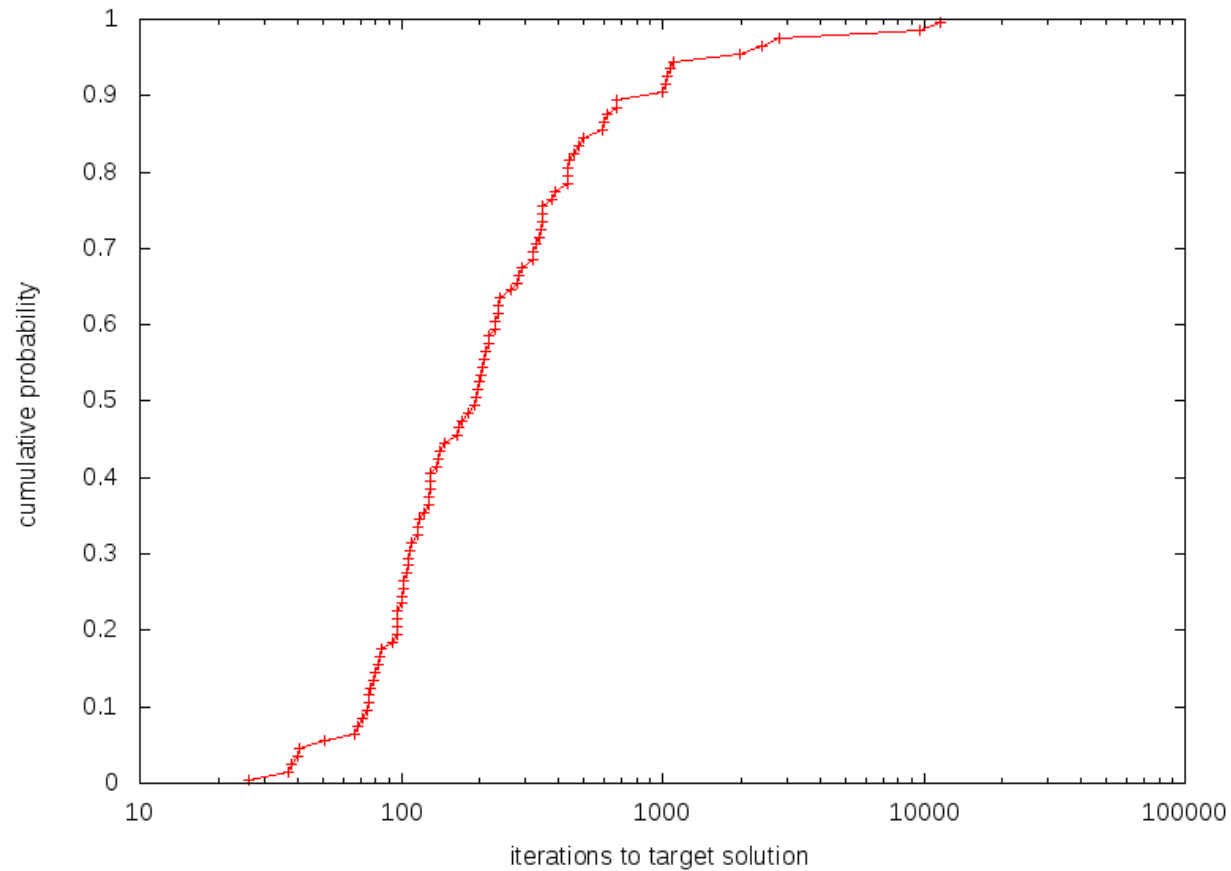
solution



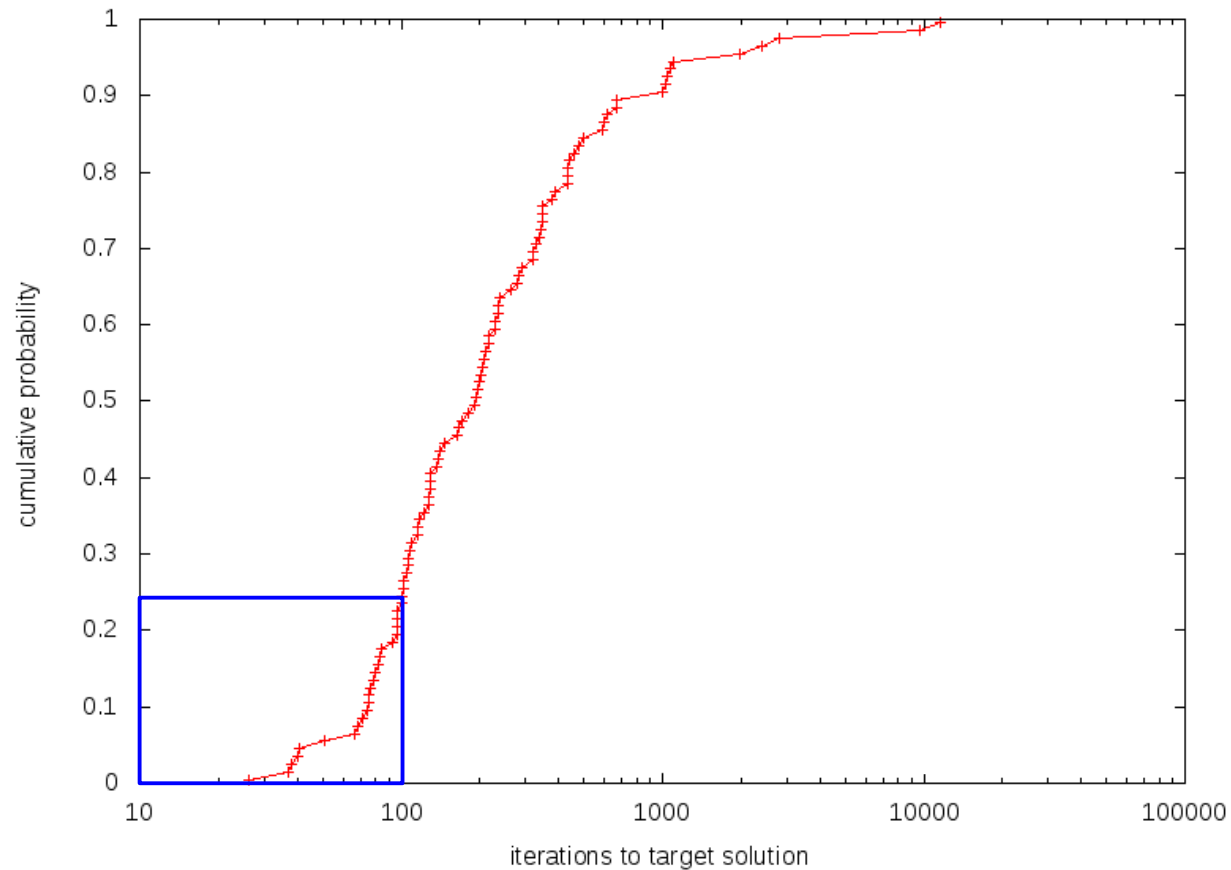
# BRKGA in multi-start strategy



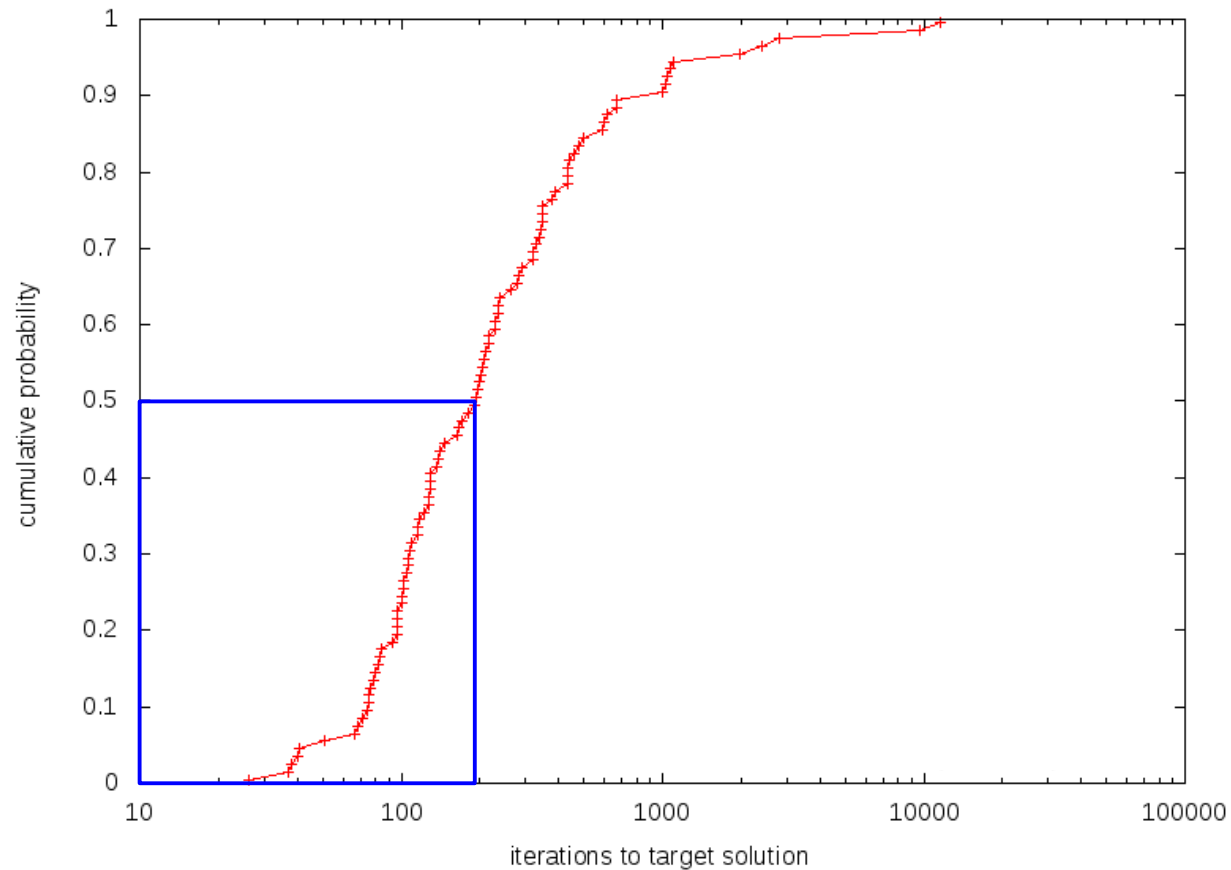




In most of the independent runs, the algorithm finds the target solution in relatively few iterations:

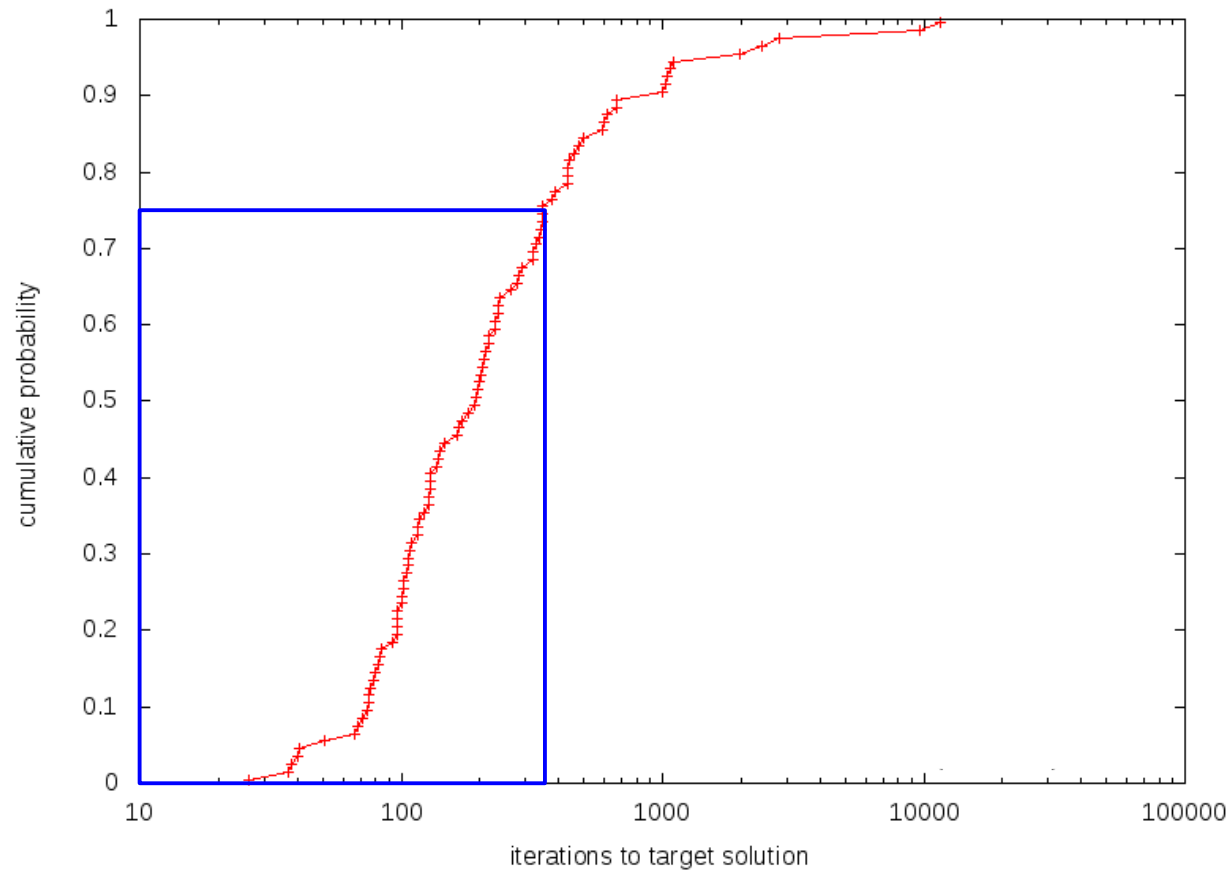


In most of the independent runs, the algorithm finds the target solution in relatively few iterations: 25% of the runs take fewer than 101 iterations

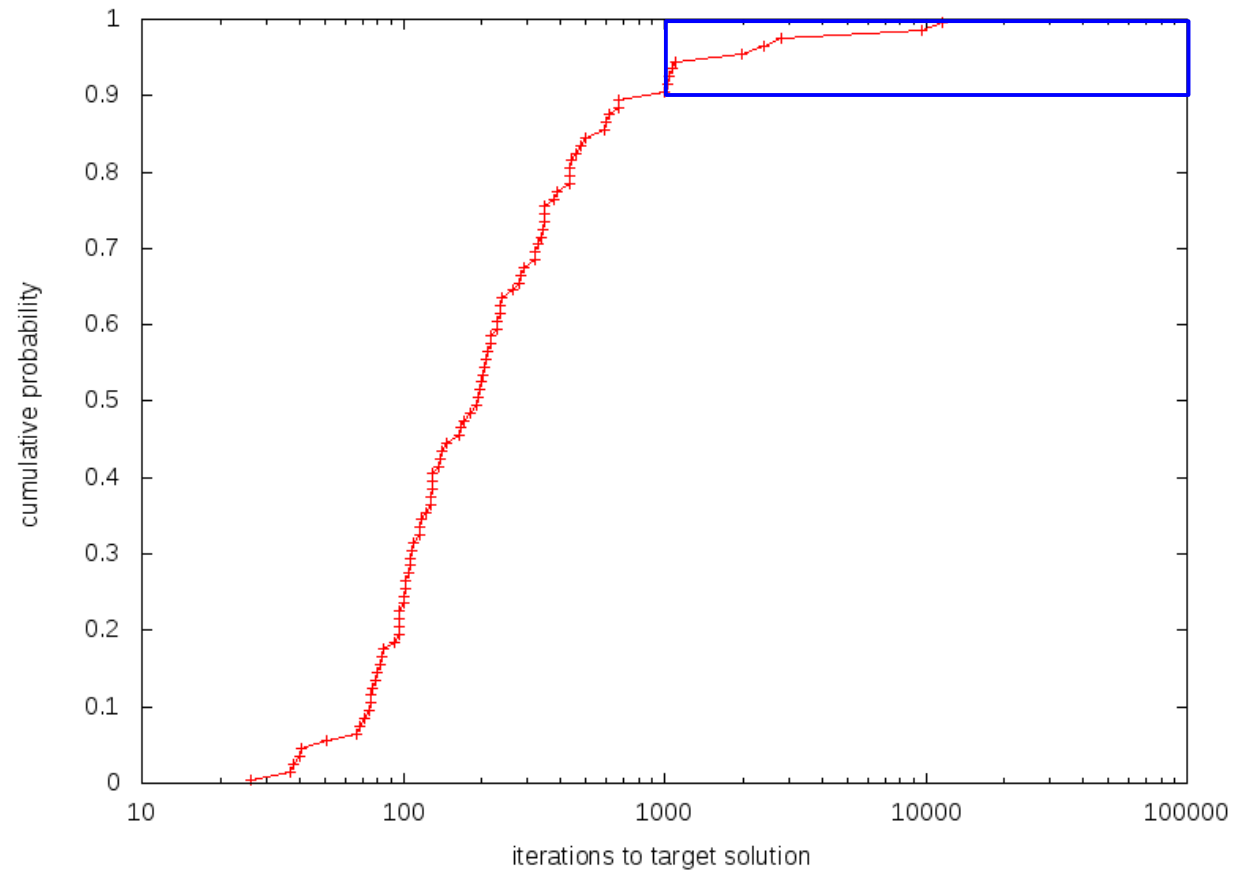


In most of the independent runs, the algorithm finds the target solution in relatively few iterations: 50% of the runs take fewer than 192 iterations

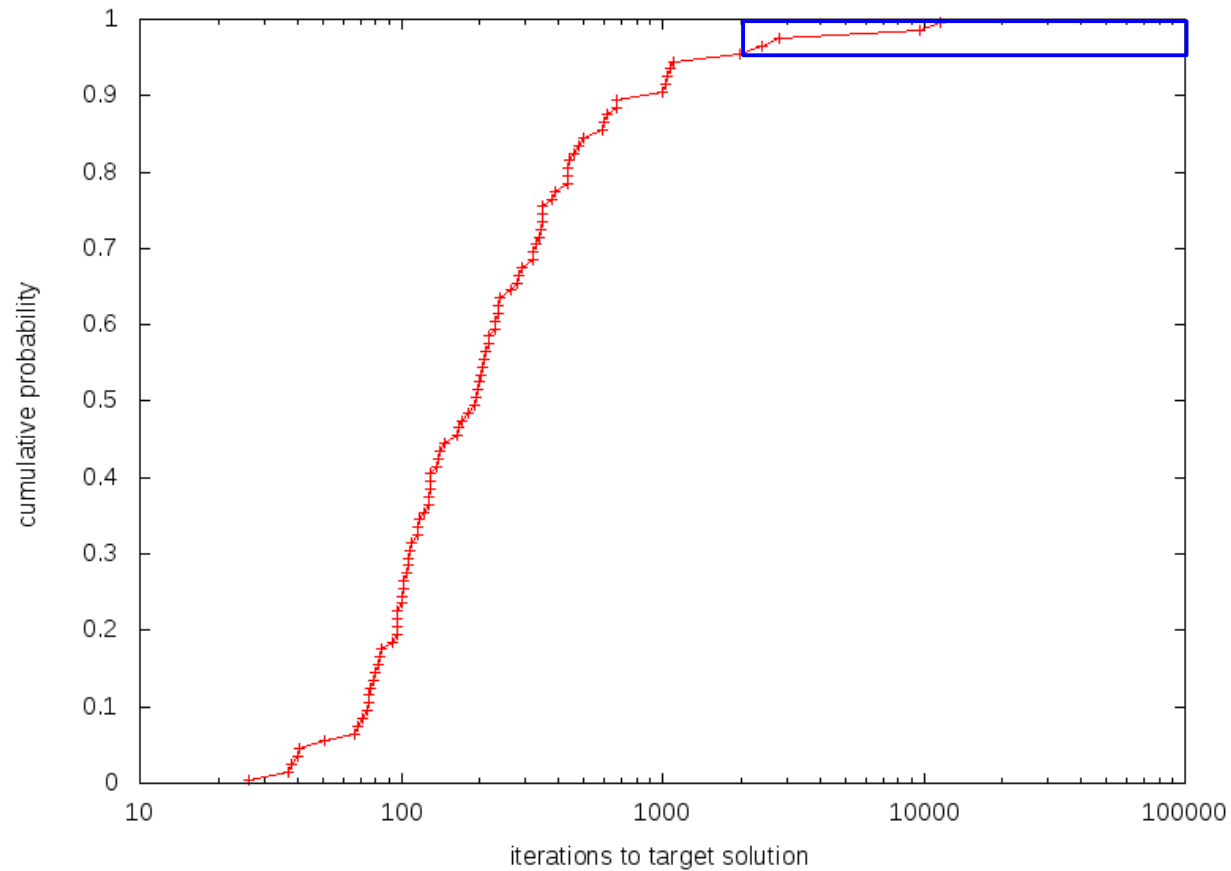




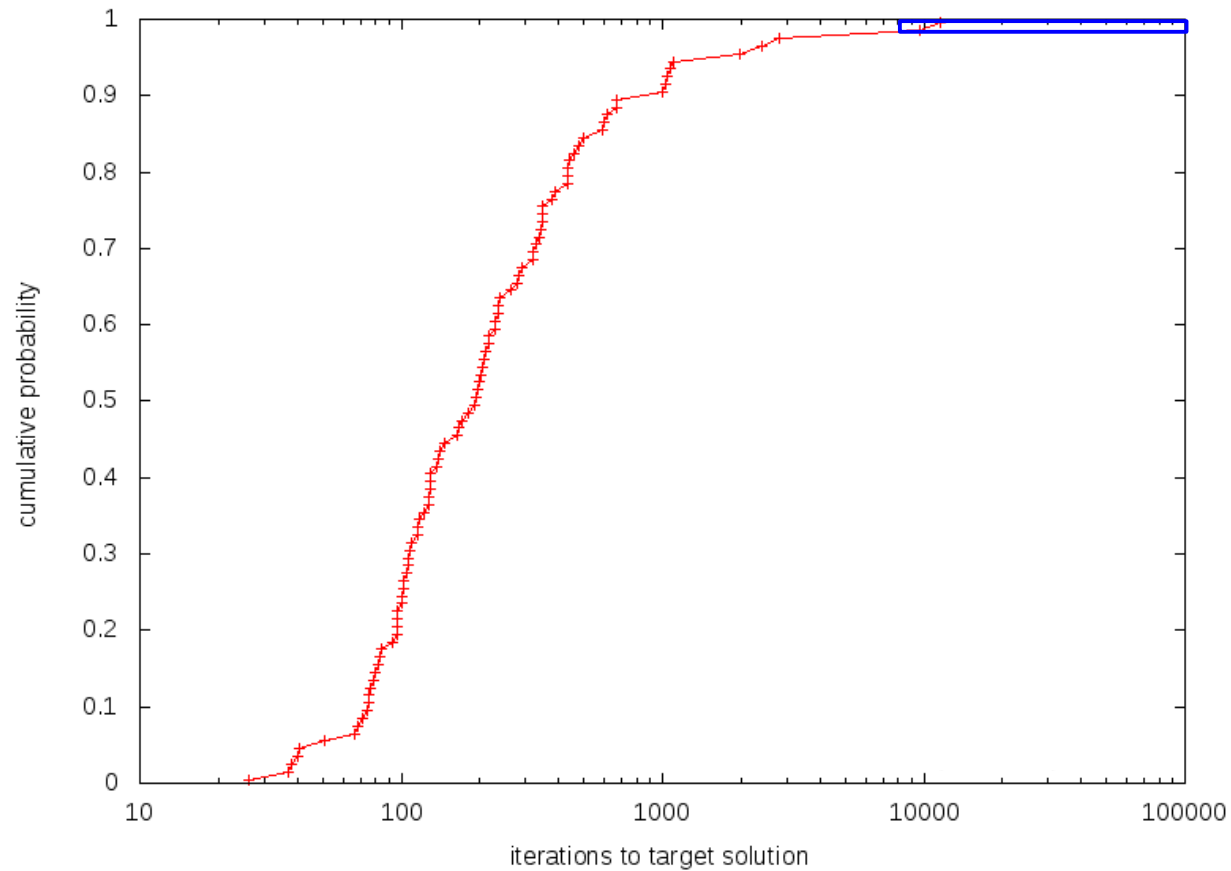
In most of the independent runs, the algorithm finds the target solution in relatively few iterations: 75% of the runs take fewer than 345 iterations



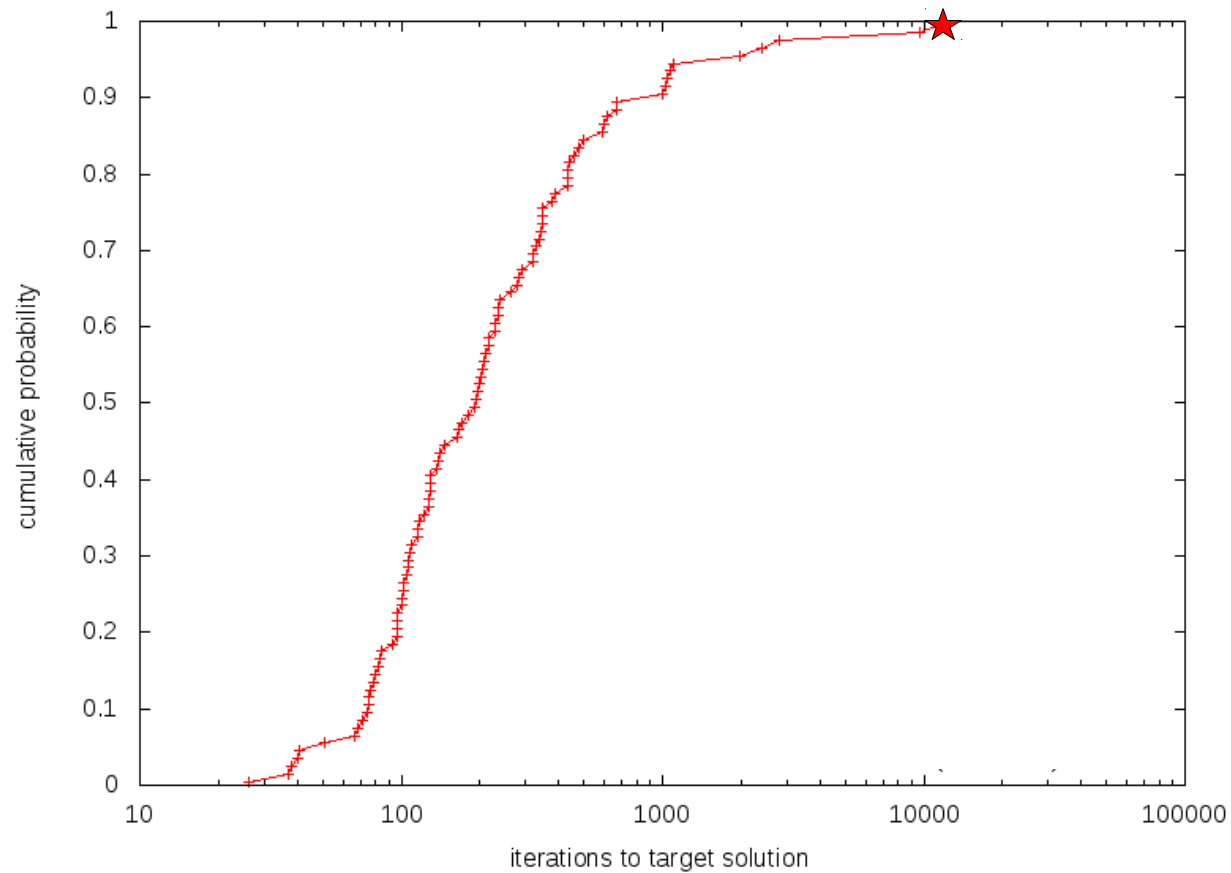
However, some runs take much longer: 10% of the runs take over 1000 iterations



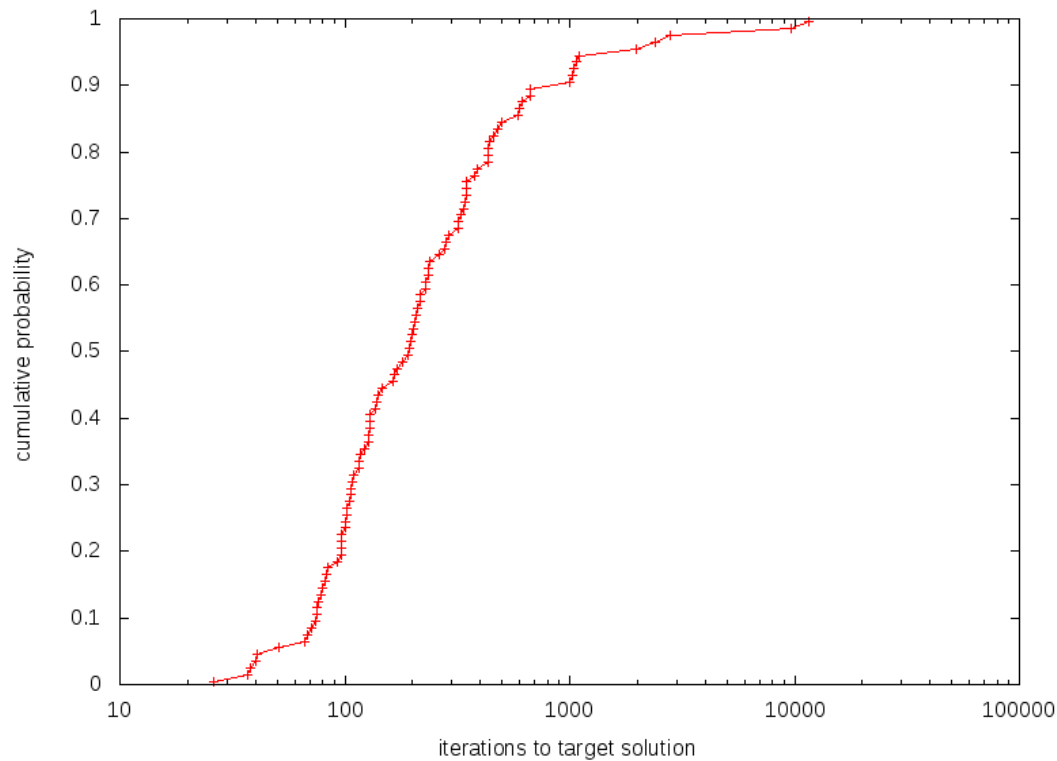
However, some runs take much longer: 5% of the runs take over 2000 iterations



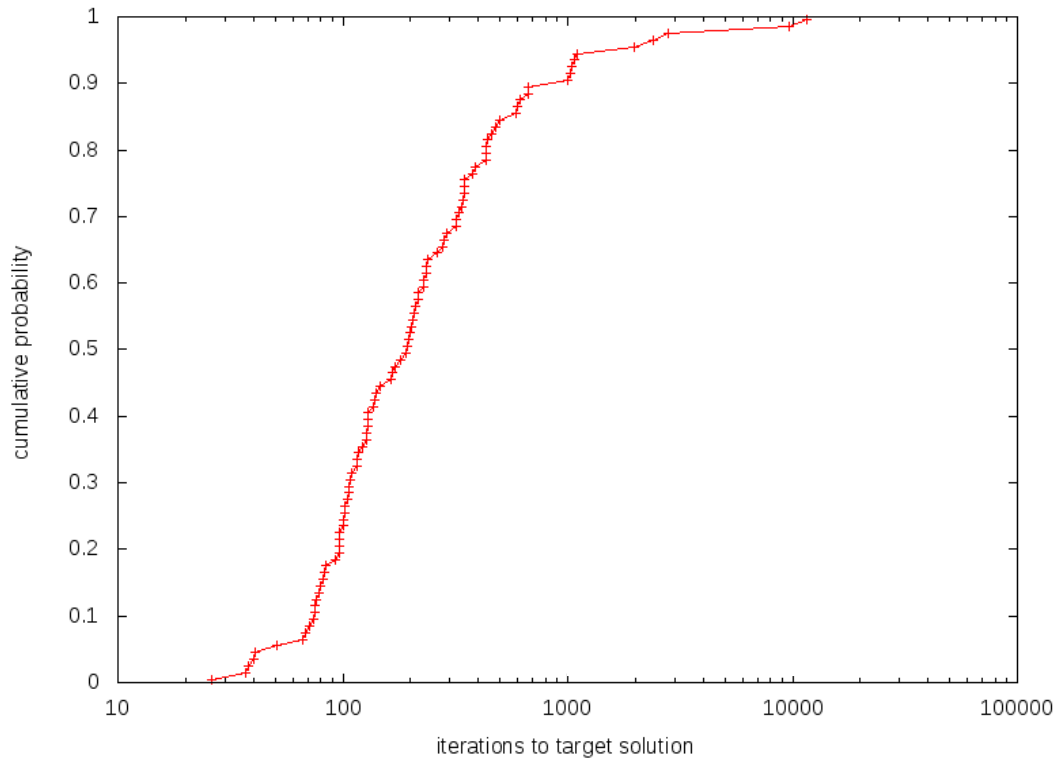
However, some runs take much longer: 2% of the runs take over 9715 iterations



However, some runs take much longer: the longest run took 11607 iterations



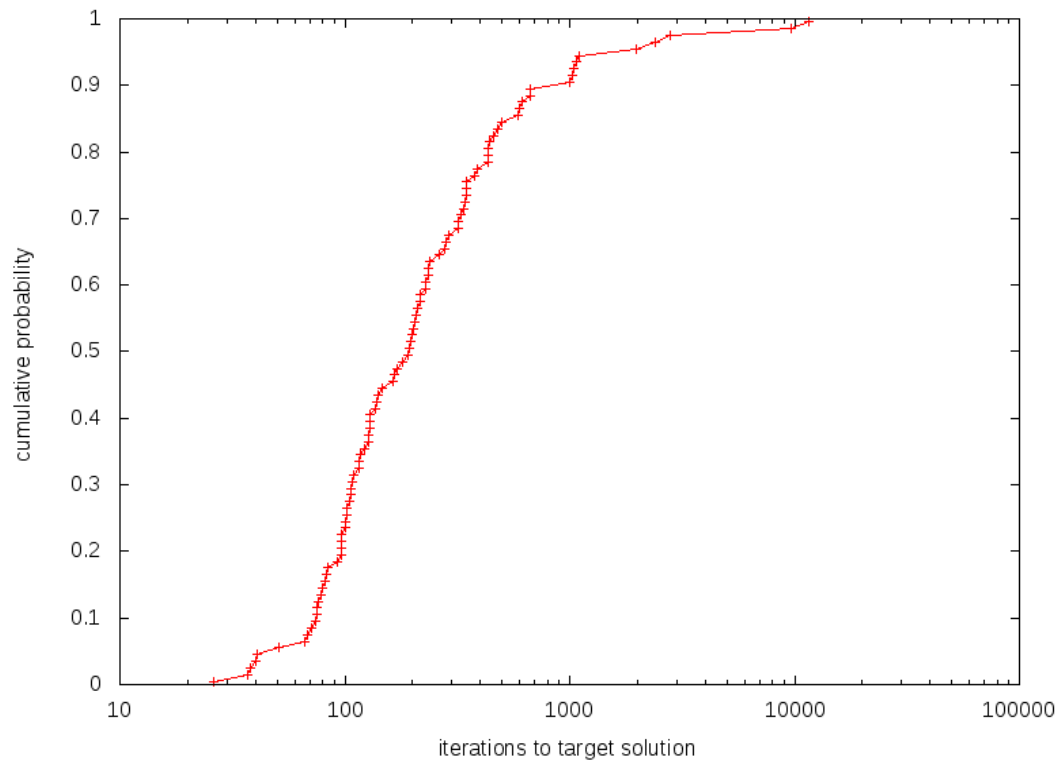
Probability that algorithm will take  
over 345 iterations:  $25\% = 1/4$



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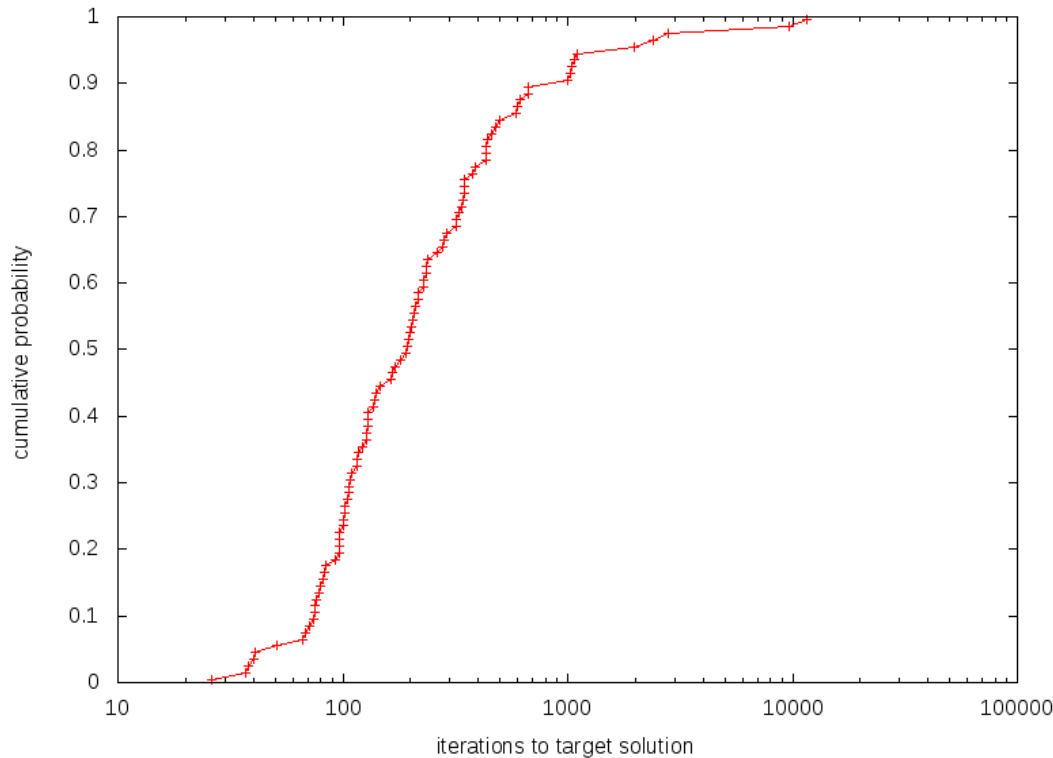
By restarting algorithm after 345 iterations, probability that new run will take over 690 iterations:  $25\% = 1/4$

Probability that algorithm with restart will take over 690 iterations: probability of taking over 345  $\times$  probability of taking over 690 iterations given it took over 345 =  $1/4 \times 1/4 = 1/4^2$



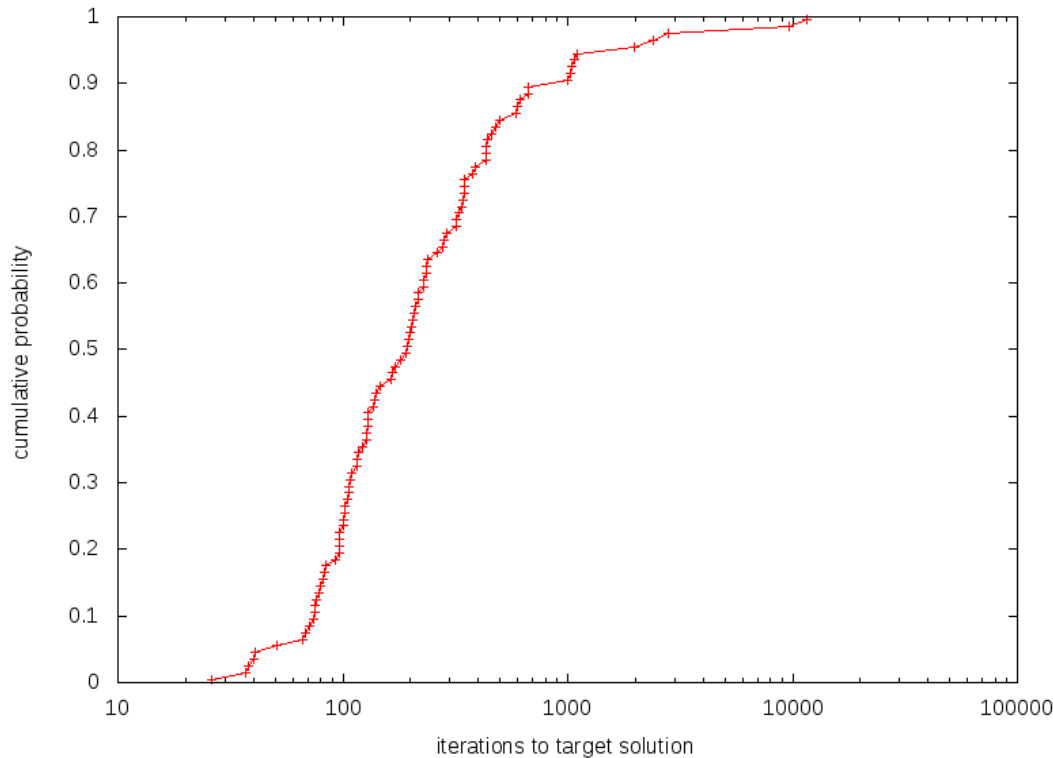
Probability that algorithm will still be  
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Probability that algorithm will still be running after  $K$  periods of 345 iterations:  $1/4^K$

For example, probability that algorithm with restart will still be running after 1725 iterations (5 periods of 345 iterations):  $1/4^5 \cong 0.0977\%$



Probability that algorithm will still be running after  $K$  periods of 345 iterations:  $1/4^K$

For example, probability that algorithm with restart will still be running after 1725 iterations (5 periods of 345 iterations):  $1/4^5 \cong 0.0977\%$

This is much less than the 5% probability that the algorithm without restart will take over 2000 iterations.

# Restart strategies

- First proposed by Luby et al. (1993)
- They define a restart strategy as a finite sequence of time intervals  $S = \{\tau_1, \tau_2, \tau_3, \dots\}$  which define epochs  $\tau_1, \tau_1 + \tau_2, \tau_1 + \tau_2 + \tau_3, \dots$  when the algorithm is restarted from scratch.
- Luby et al. (1993) prove that the optimal restart strategy uses  $\tau_1 = \tau_2 = \tau_3 = \dots = \tau^*$ , where  $\tau^*$  is a constant.

# Restart strategies

- Luby et al. (1993)
- Kautz et al. (2002)
- Palubeckis (2004)
- Sergienko et al. (2004)
- Nowicki & Smutnicki (2005)
- D'Apuzzo et al. (2006)
- Shylo et al. (2011a)
- Shylo et al. (2011b)
- Resende & Ribeiro (2011)

# Restart strategy for BRKGA

- Recall the restart strategy of Luby et al. where equal time intervals  $\tau_1 = \tau_2 = \tau_3 = \dots = \tau^*$  pass between restarts.
- Strategy requires  $\tau^*$  as input.
- Since we have no prior information as to the runtime distribution of the heuristic, we run the risk of:
  - choosing  $\tau^*$  too small: restart variant may take long to converge
  - choosing  $\tau^*$  too big: restart variant may become like no-restart variant

# Restart strategy for BRKGA

- We conjecture that number of iterations between improvement of the incumbent (best so far) solution varies less w.r.t. heuristic/ instance/ target than run times.
- We propose the following restart strategy: Keep track of the last generation when the incumbent improved and restart BRKGA if  $K$  generations have gone by without improvement.
- We call this strategy  $\text{restart}(K)$

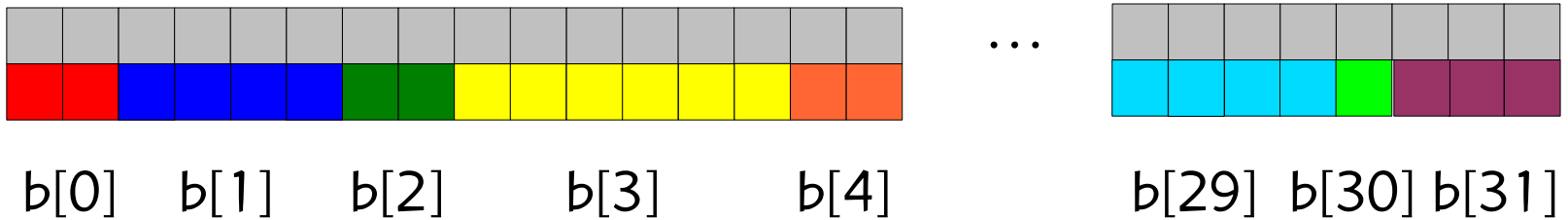
# Example of restart strategy for BRKGA: Load balancing

Given an unordered sequence of 1024 integers  $p[0], p[1], \dots, p[1023]$



# Example of restart strategy for BRKGA: Load balancing

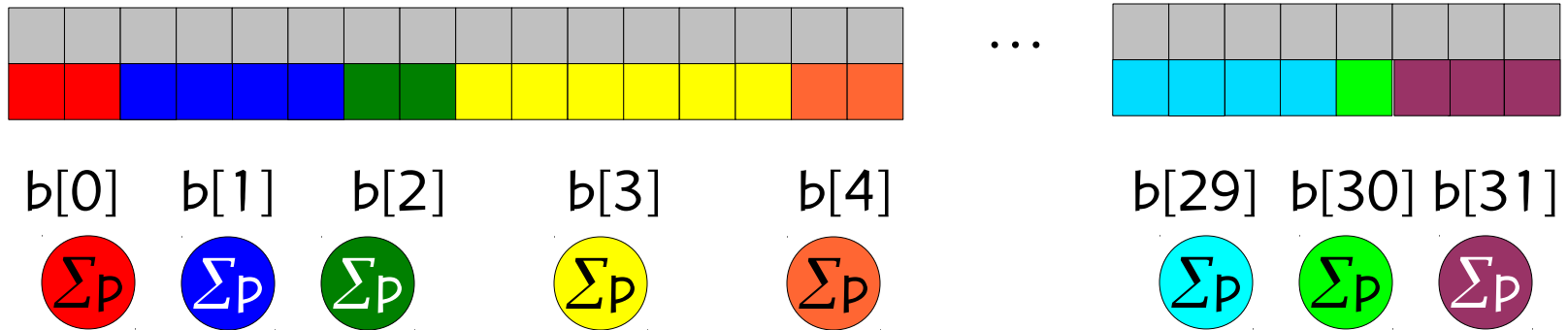
Place consecutive numbers in 32 buckets  $b[0]$ ,  $b[1]$ , ...,  $b[31]$





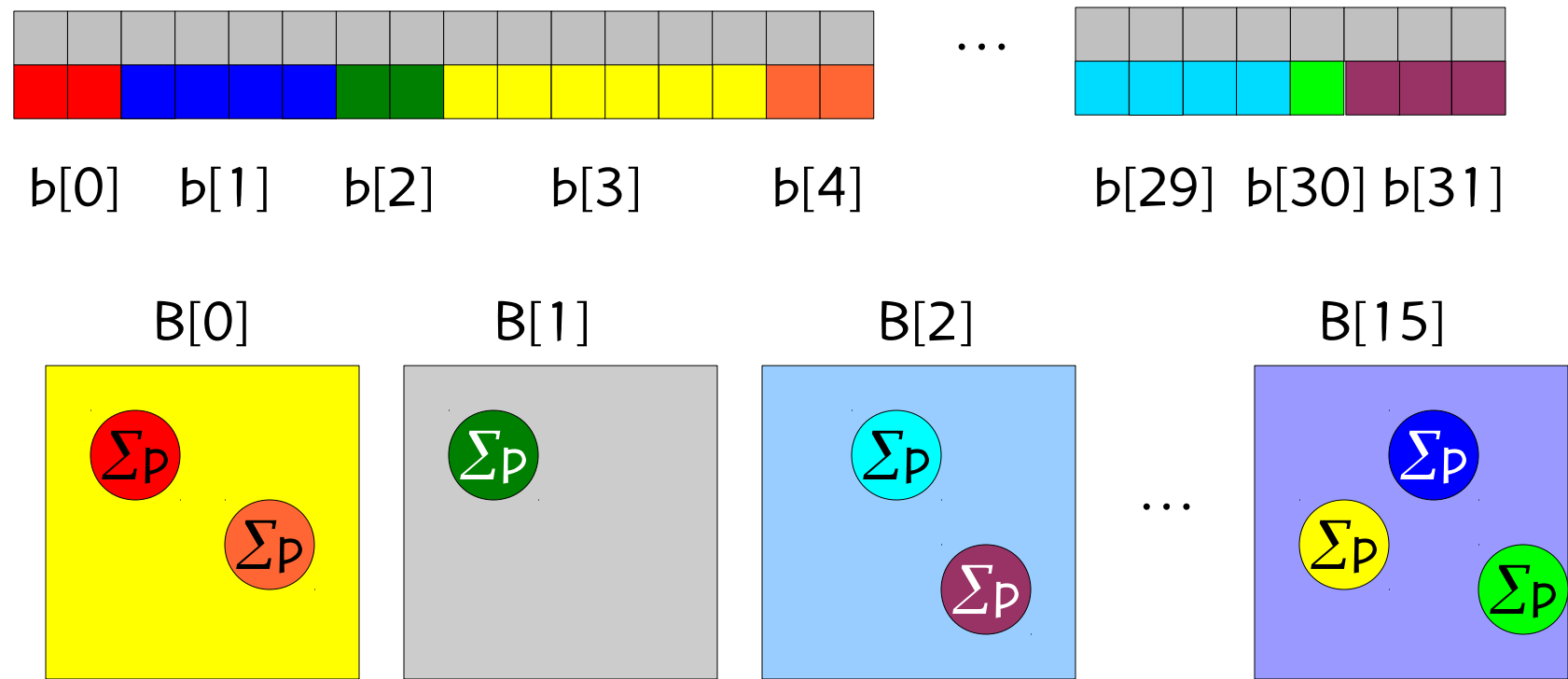
# Example of restart strategy for BRKGA: Load balancing

Add the numbers in each bucket  $b[0]$ ,  $b[1]$ , ...,  $b[31]$



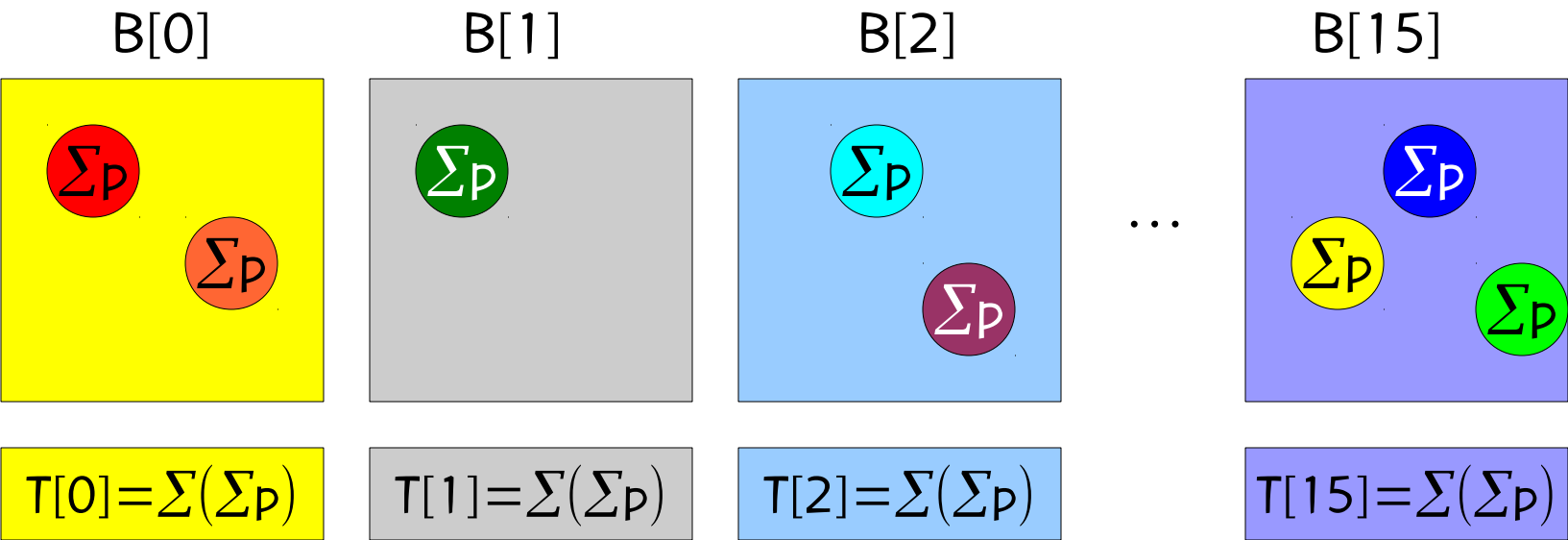
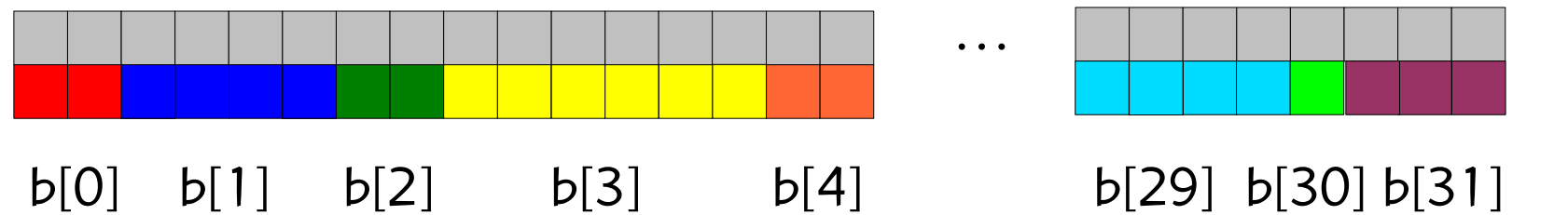
# Example of restart strategy for BRKGA: Load balancing

Place the buckets in 16 bins  $B[0], B[1], \dots, B[15]$



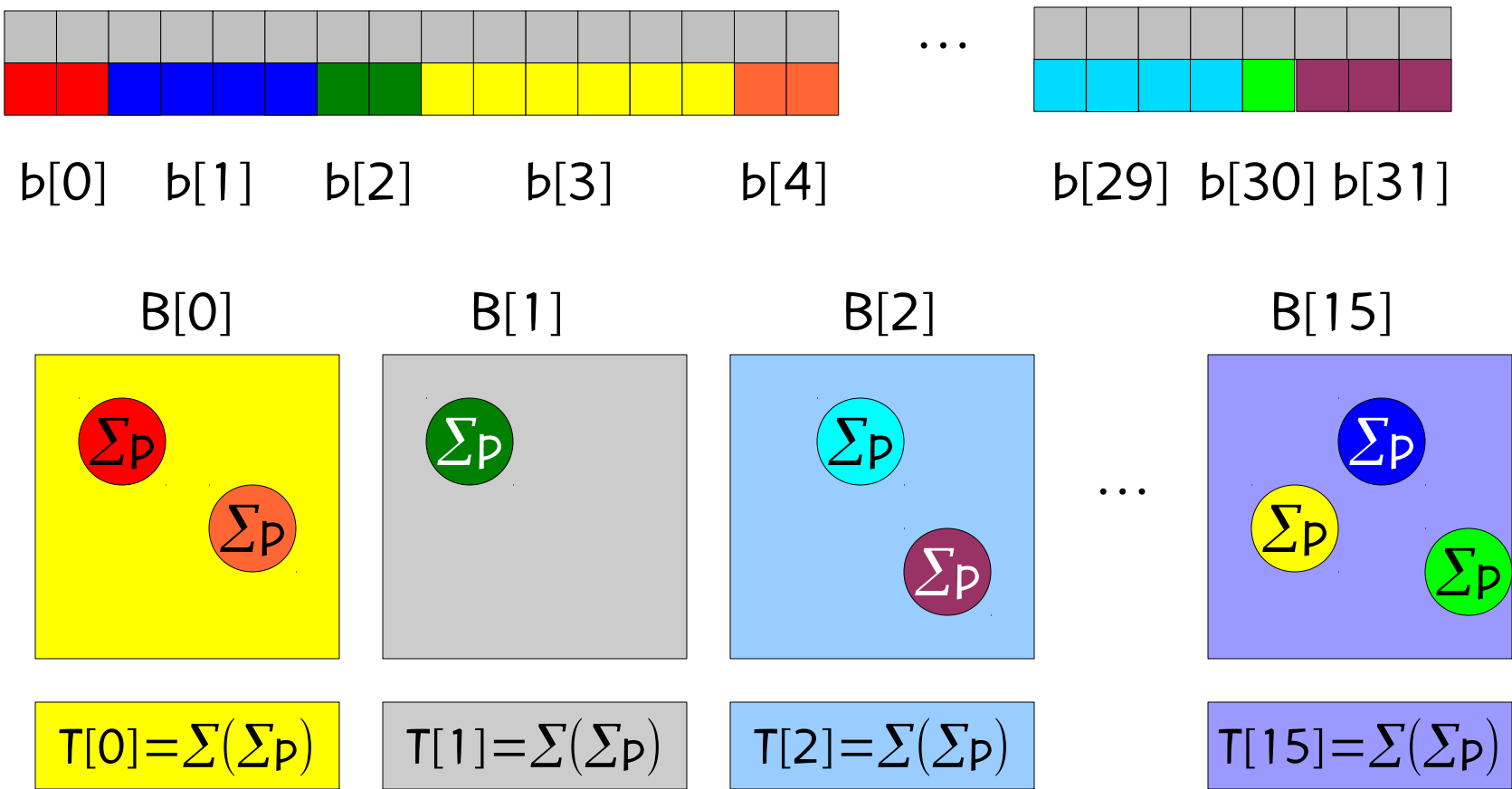
# Example of restart strategy for BRKGA: Load balancing

Add up the numbers in each bin  $B[0], B[1], \dots, B[15]$



# Example of restart strategy for BRKGA: Load balancing

OBJECTIVE: Minimize { Maximum ( $T[0], T[1], \dots, T[15]$ ) }



# Example of restart strategy for BRKGA: Load balancing

## Encoding

$$X = [ x[1], x[2], \dots, x[32] \quad | \quad x[32+1], x[32+2], \dots, x[32+16] ]$$

## Decoding

$x[1], x[2], \dots, x[32]$  are used to define break points for buckets

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## Decoding

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Size of bucket  $i = \text{floor}(1024 \times x[i] / (x[1] + x[2] + \dots + x[32]))$ ,  $i=1, \dots, 15$

Size of bucket 16 =  $1024 - \text{sum of sizes of first 15 buckets}$

# Example of restart strategy for BRKGA: Load balancing

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Bin that bucket  $i$  is assigned to =  $\text{floor}(16 \times x[32+i]) + 1$

# Example of restart strategy for BRKGA: Load balancing

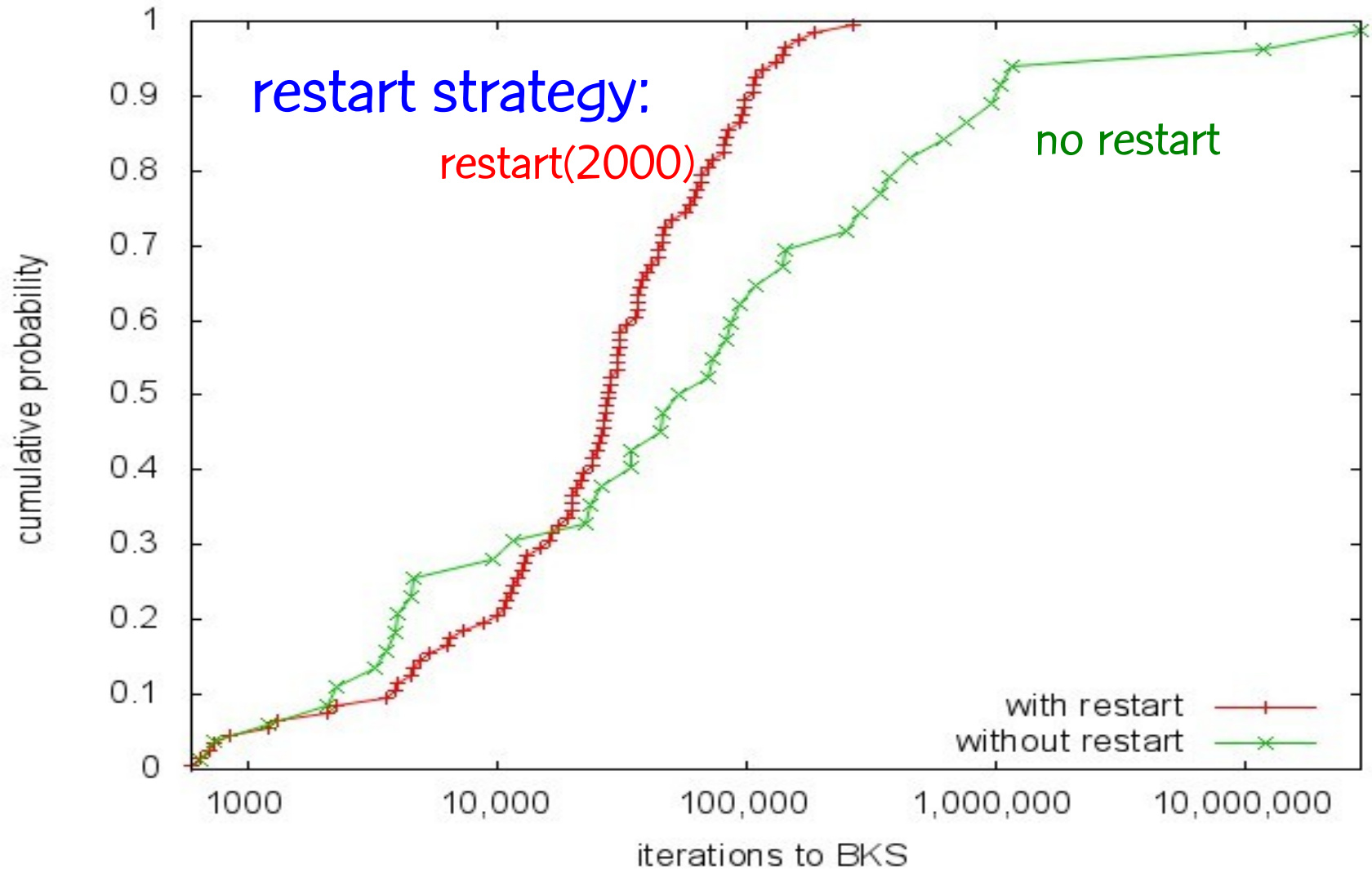
## Decoding (Local search phase)

- **while** (there exists a bucket in the most loaded bin that can be moved to another bin and not increase the maximum load) **then**
  - move that bucket to that bin
- **end while**

Make necessary chromosome adjustments to last 16 random keys of vector of random keys to reflect changes made in local search phase: Add or subtract an integer value from chromosome of bucket that moved to new bin.



# Example of restart strategy for BRKGA: Load balancing



# Specifying a BRKGA

# Specifying a biased random-key GA

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# Specifying a biased random-key GA

## Parameters:

- Size of population
- Parallel population parameters
- Size of elite partition
- Size of mutant set
- Child inheritance probability
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- Stopping criterion

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- Restart strategy parameter: a function of  $N$ , say  $2N$  or  $10N$
- Stopping criterion: e.g. time, # generations, solution quality, # generations without improvement

# brkgaAPI: A C++ API for BRKGA

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- Implemented in C++ and may benefit from shared-memory parallelism if available.
- User only needs to implement problem-dependent decoder.

# brkgaAPI: A C++ API for BRKGA

Paper: Rodrigo F. Toso and M.G.C.R., "A C++  
Application Programming Interface for  
Biased Random-Key Genetic Algorithms,"  
AT&T Labs Technical Report, Florham Park, August 2011.

Software: <http://www.research.att.com/~mgcr/src/brkgaAPI>

# An example BRKGA: Packing weighted rectangles

# Reference



J.F. Gonçalves and M.G.C.R., "A parallel multi-population genetic algorithm for a constrained two-dimensional orthogonal packing problem," Journal of Combinatorial Optimization, vol. 22, pp. 180-201, 2011.

Tech report:

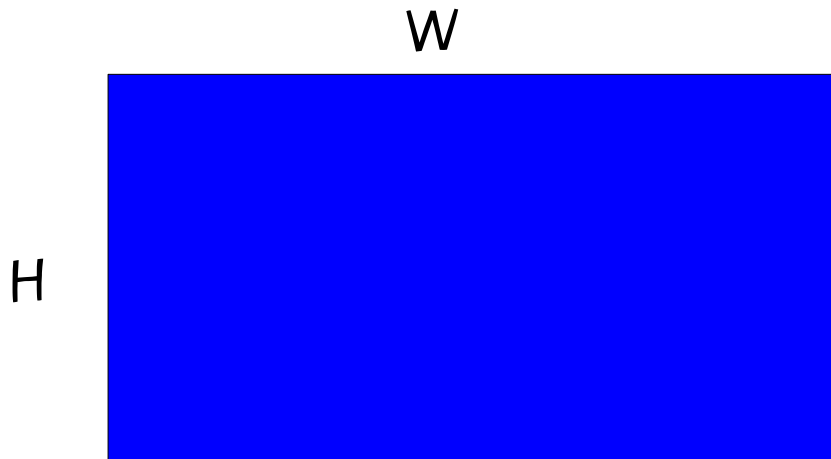
<http://www.research.att.com/~mgcr/doc/pack2d.pdf>

# Constrained orthogonal packing

- Given a large planar stock rectangle ( $W$ ,  $H$ ) of width  $W$  and height  $H$ ;

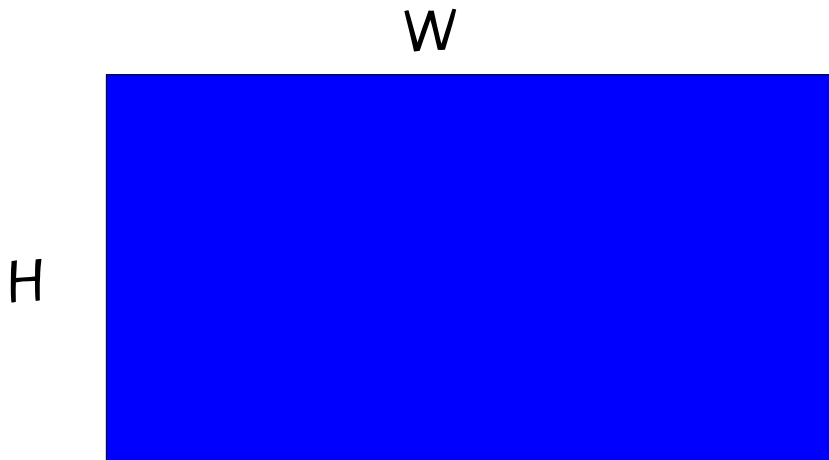
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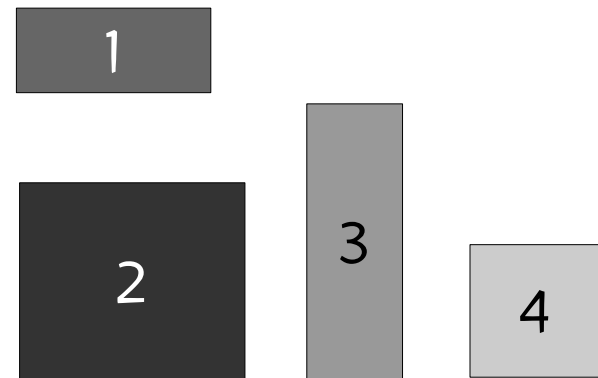
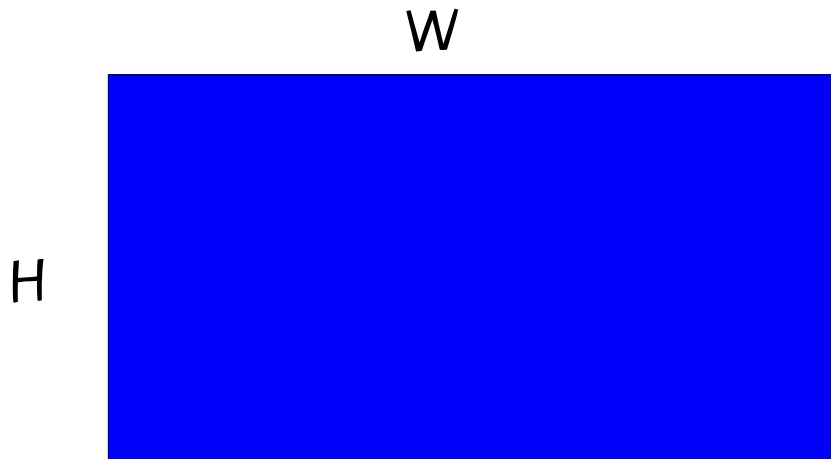
# Constrained orthogonal packing

- Given a large planar stock rectangle ( $W$ ,  $H$ ) of width  $W$  and height  $H$ ;
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# Constrained orthogonal packing

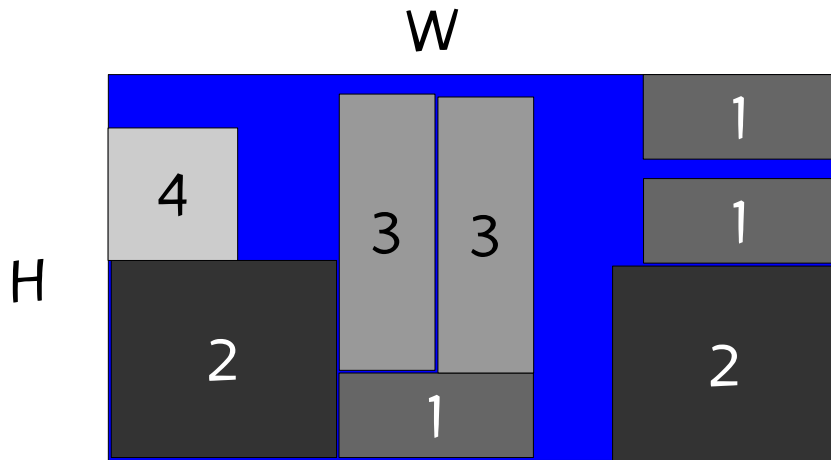
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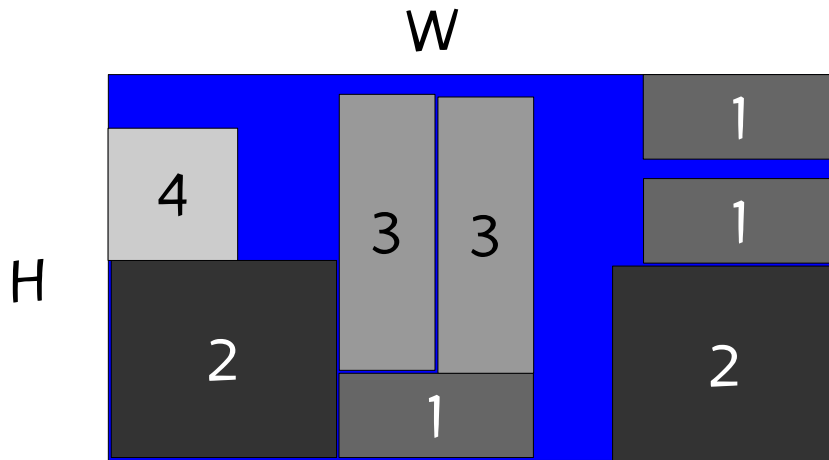
- $r[i]$  rectangles of type  $i = 1, \dots, N$  are to be packed in the large rectangle without overlap and such that their edges are parallel to the edges of the large rectangle;



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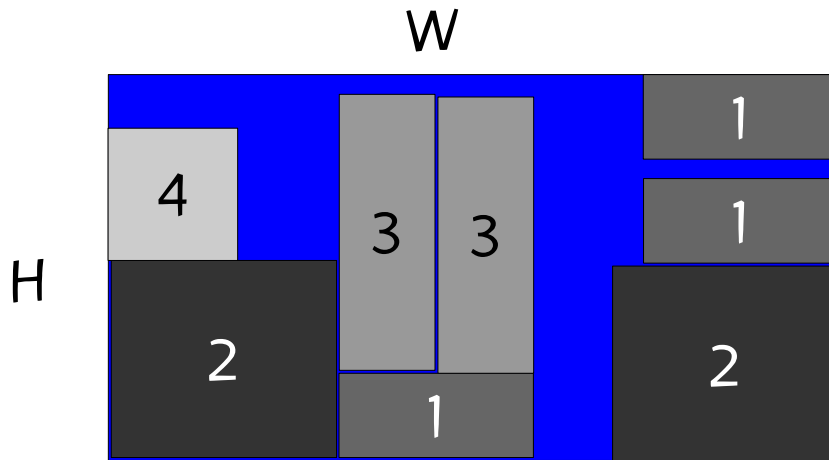
$$0 \leq P[i] \leq r[i] \leq Q[i]$$



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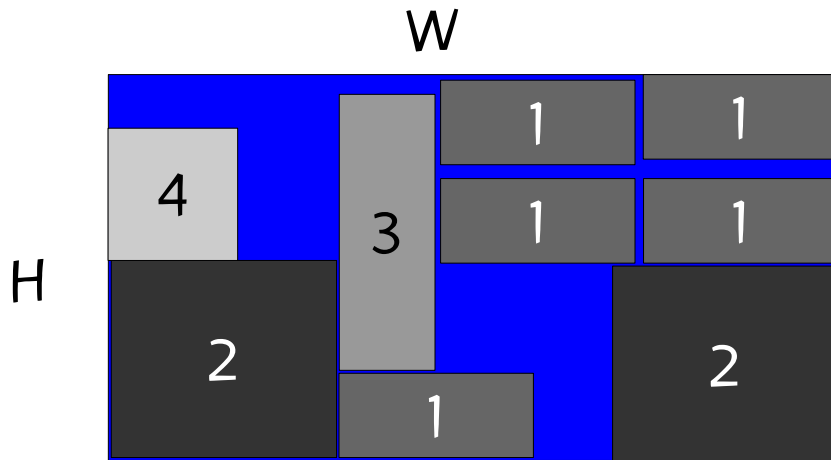


Suppose  $5 \leq r[1] \leq 12$

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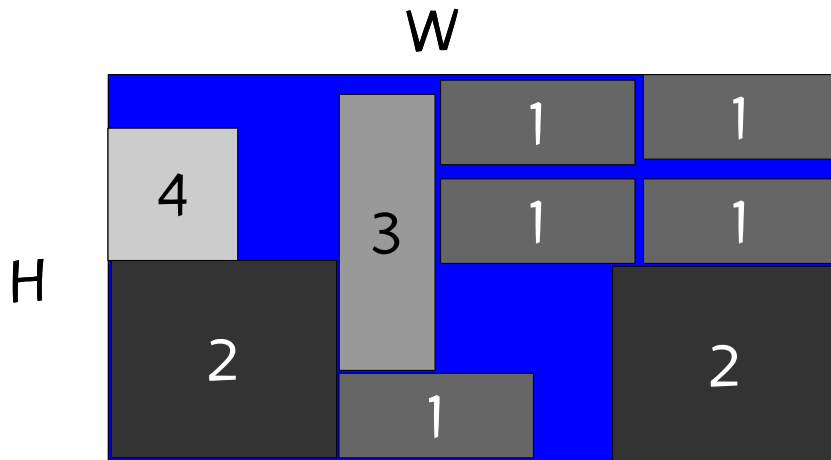


Suppose  $5 \leq r[1] \leq 12$

# Objective

Among the many feasible packings, we want to find one that maximizes total value of packed rectangles:

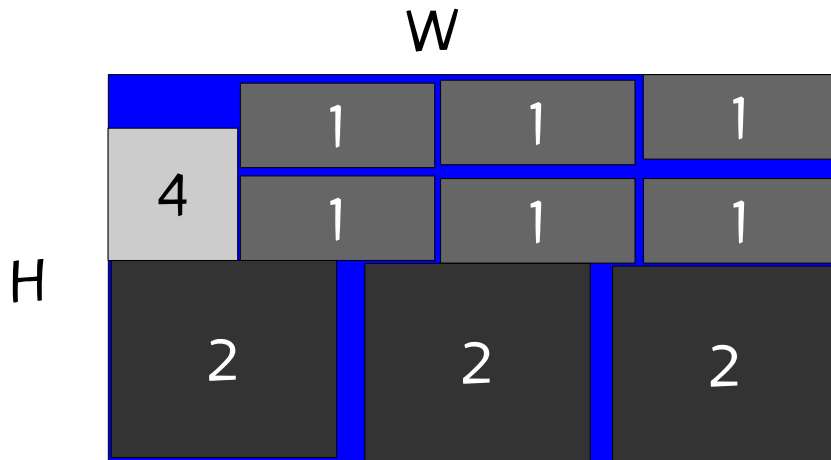
$$v[1] r[1] + v[2] r[2] + \dots + v[N] r[N]$$



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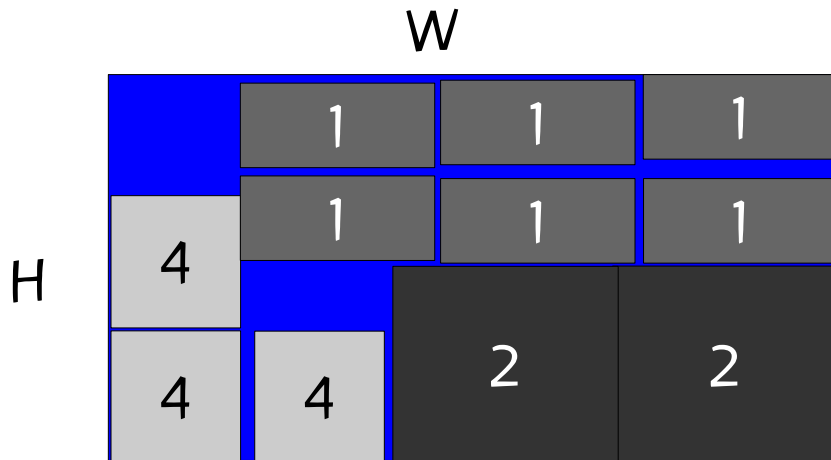
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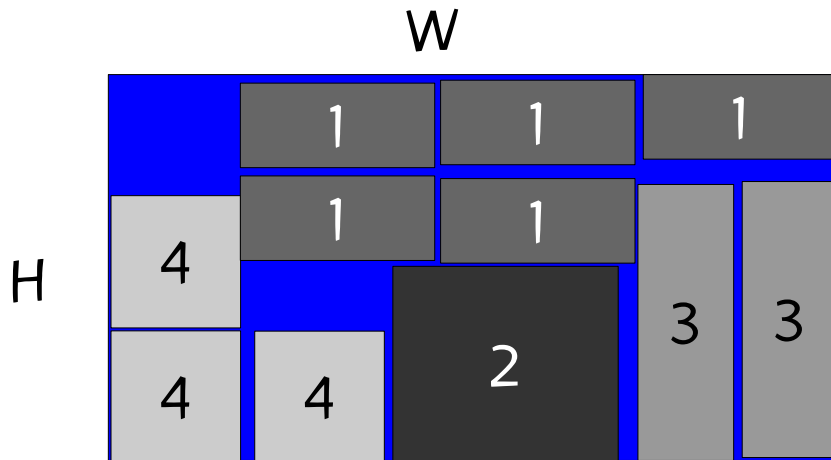
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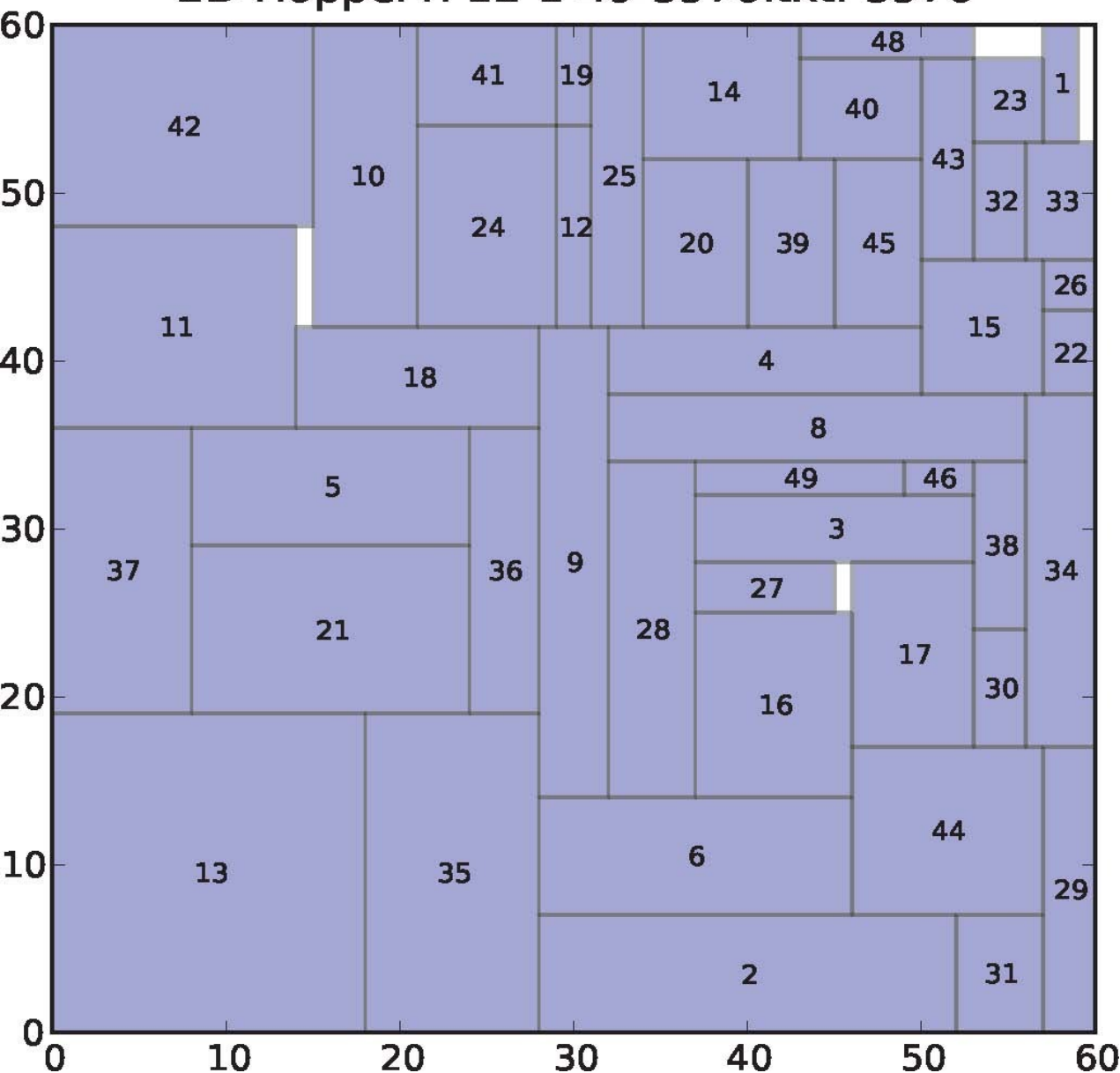
# Applications

Problem arises in several production processes, e.g.

- Textile
- Glass
- Wood
- Paper

where rectangular figures are cut from large rectangular sheets of materials.

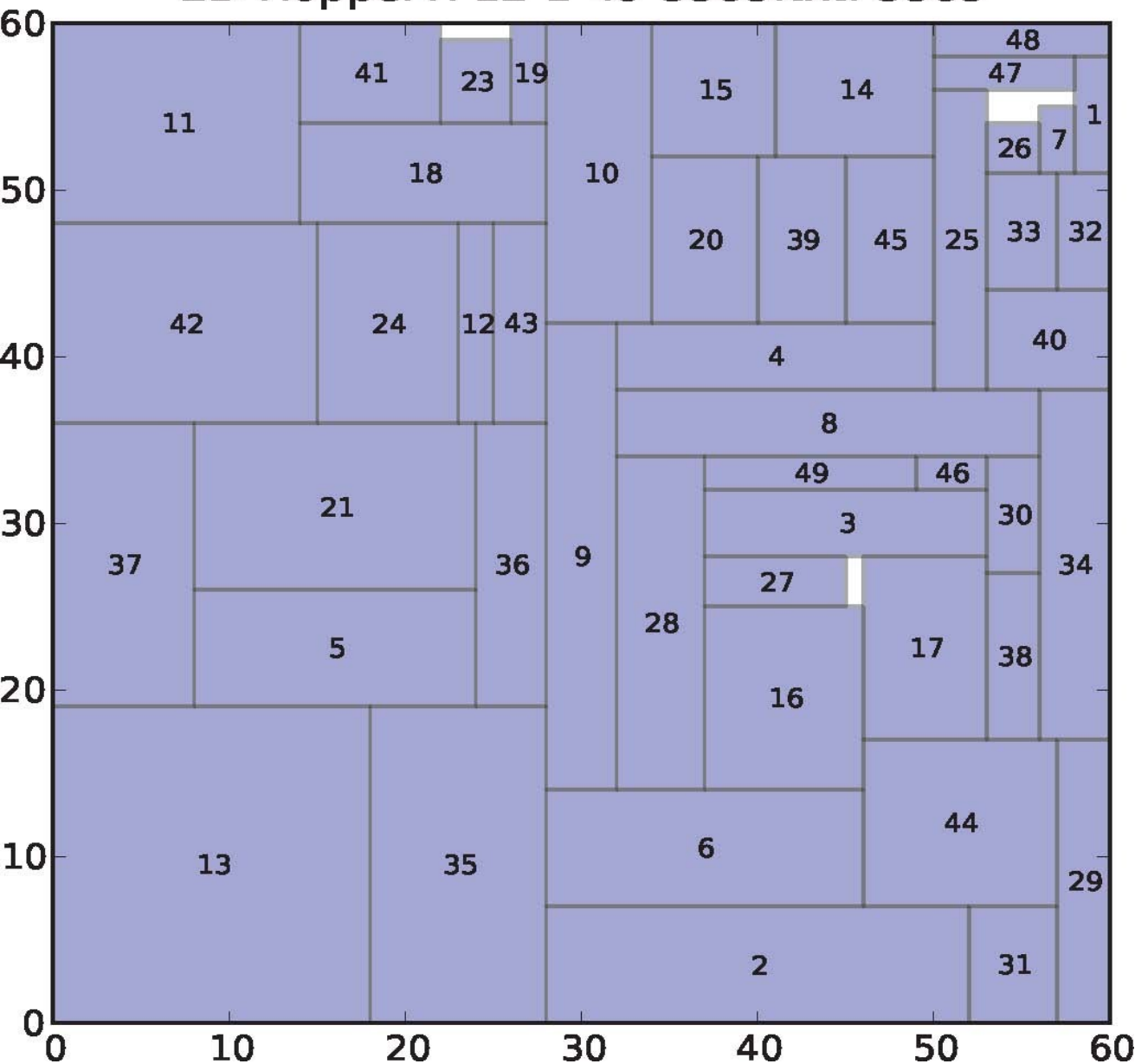
# 2D-HopperTP12-1-49-3576.txt: 3576



Hopper & Turton, 2001  
 Instance 4-1 60 x 60  
 Value: 3576

Previous best: 3580 by a  
 Tabu Search heuristic  
 (Alvarez-Valdes et al., 2007)

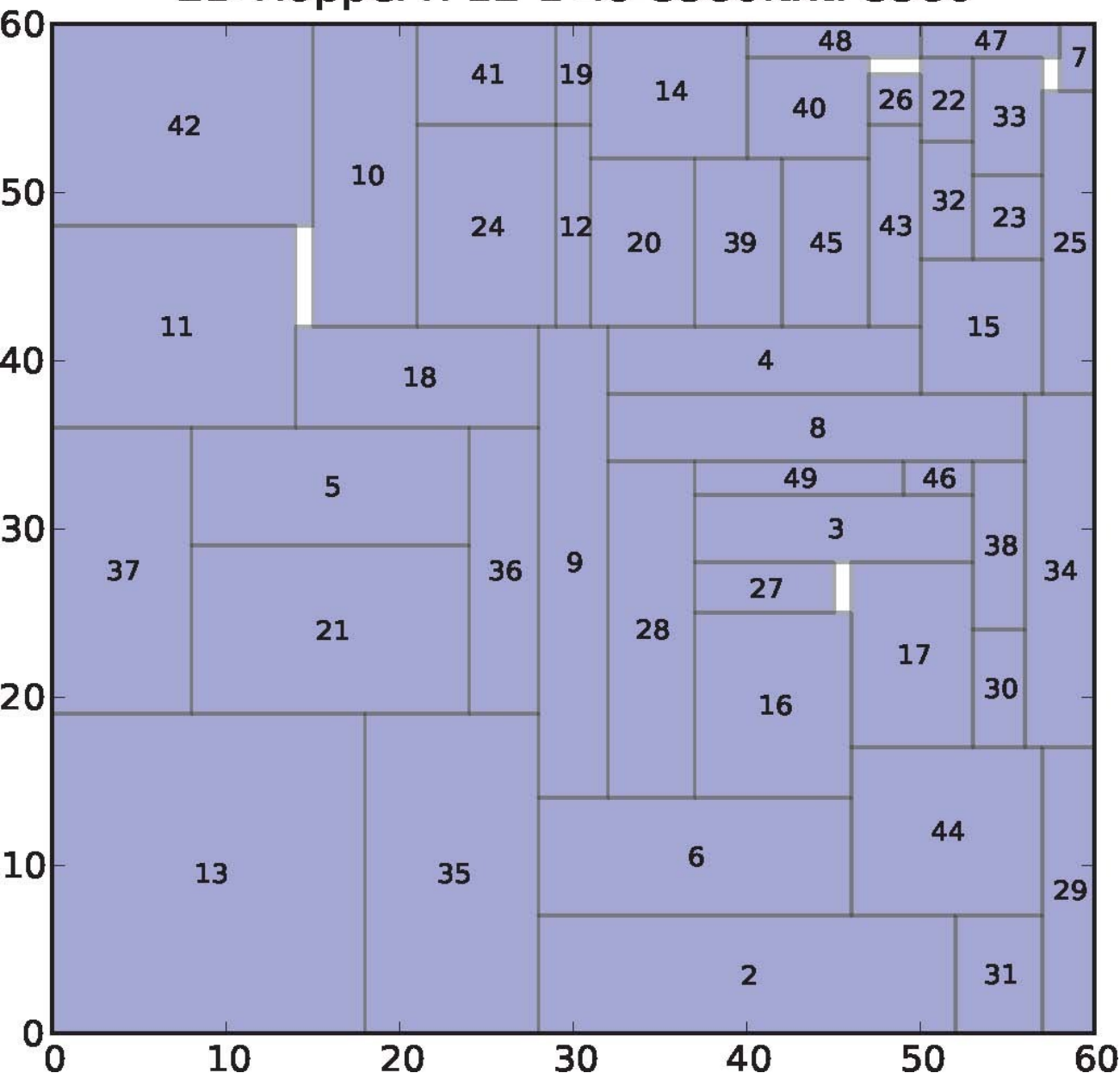
# 2D-HopperTP12-1-49-3585.txt: 3585



Hopper & Turton, 2001  
 Instance 4-2 60 x 60  
 Value: 3585

Previous best: 3580 by a  
 Tabu Search heuristic  
 (Alvarez-Valdes et al., 2007)

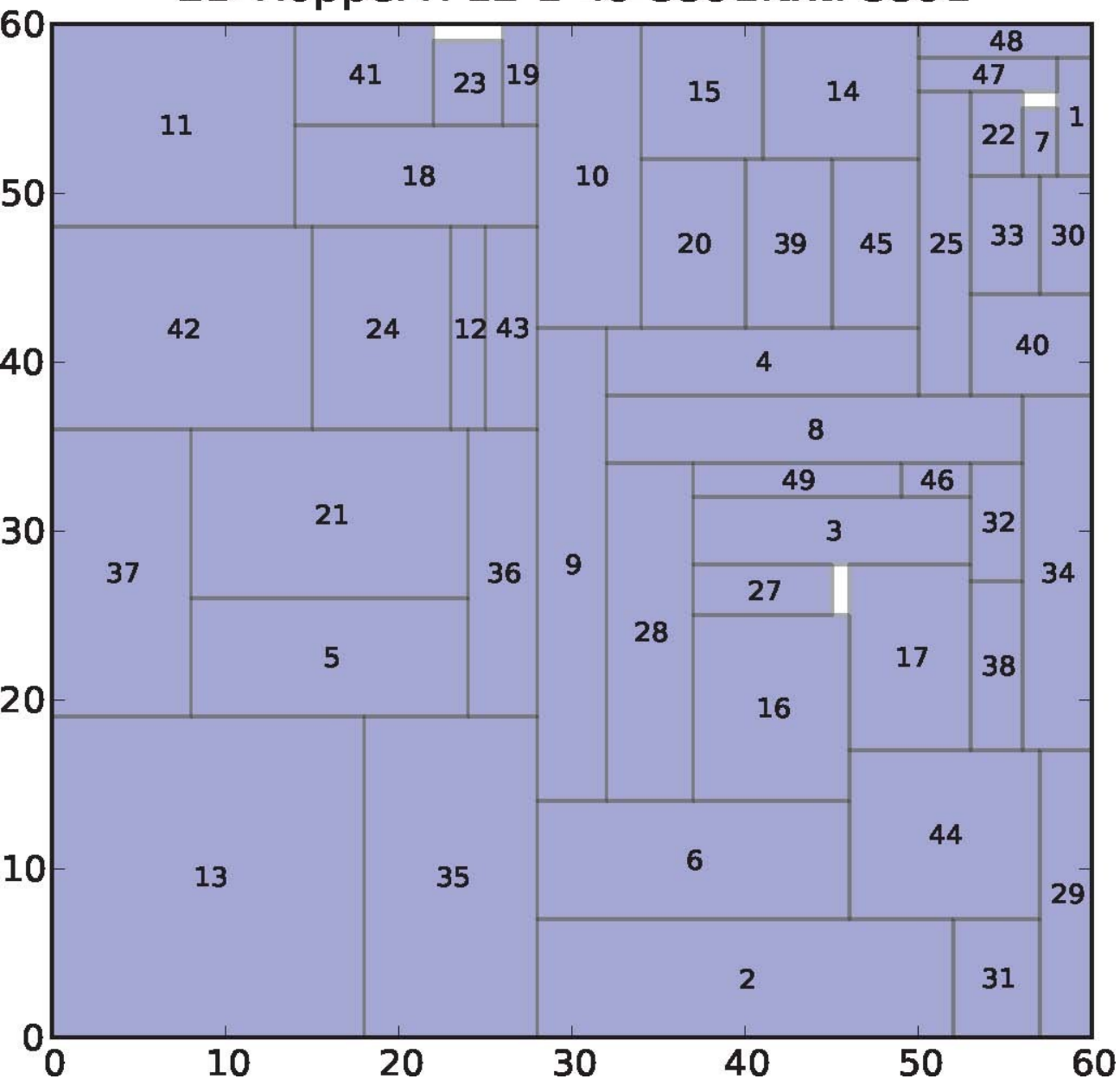
# 2D-HopperTP12-1-49-3586.txt: 3586



Hopper & Turton, 2001  
Instance 4-2 60 x 60  
Value: 3586

Previous best: 3580 by a  
Tabu Search heuristic  
(Alvarez-Valdes et al., 2007)

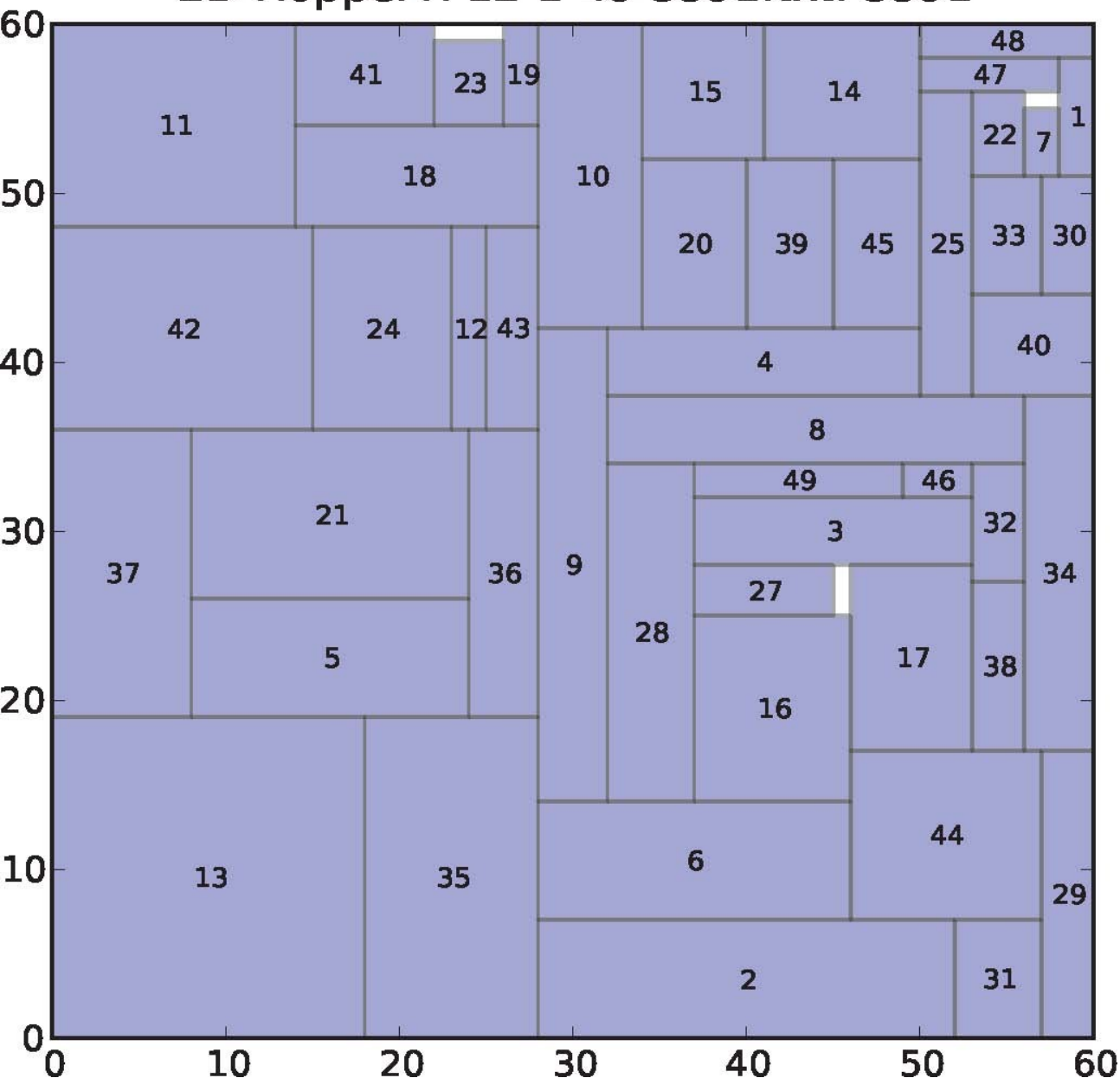
# 2D-HopperTP12-1-49-3591.txt: 3591



Hopper & Turton, 2001  
 Instance 4-2 60 x 60  
 Value: 3591

Previous best: 3580 by a  
 Tabu Search heuristic  
 (Alvarez-Valdes et al., 2007)

# 2D-HopperTP12-1-49-3591.txt: 3591



Hopper & Turton, 2001

Instance 4-2 60 x 60

Value: 3591

New best known solution!

Previous best: 3580 by a  
Tabu Search heuristic  
(Alvarez-Valdes et al., 2007)

# BRKGA for constrained 2-dim orthogonal packing

# Encoding

- Solutions are encoded as vectors  $K$  of  
$$2N' = 2 \{ Q[1] + Q[2] + \dots + Q[N] \}$$
random keys, where  $Q[i]$  is the maximum number of rectangles of type  $i$  (for  $i = 1, \dots, N$ ) that can be packed.
- $K = ( k[1], \dots, k[N'], \quad k[N'+1], \dots, k[2N'] )$



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# Decoding

- Simple heuristic to pack rectangles:
  - Make  $Q[i]$  copies of rectangle  $i$ , for  $i = 1, \dots, N$ .
  - Order the  $N' = Q[1] + Q[2] + \dots + Q[N]$  rectangles in some way.
  - Process the rectangles in the above order. Place the rectangle in the stock rectangle according to one of the following heuristics: **bottom-left (BL)** or **left-bottom (LB)**. If **rectangle cannot be positioned, discard it** and go on to the next rectangle in the order.

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# Decoding

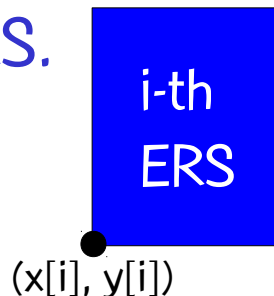
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  - Process the rectangles in the above order. Place the rectangle in the stock rectangle according to one of the following heuristics: **bottom-left (BL)** or **left-bottom (LB)**. If **rectangle cannot be positioned, discard it** and go on to the next rectangle in the order. **Use the last  $N'$  keys to determine which heuristic to use. If  $k[N'+i] > 0.5$  use LB, else use BL.**

# Decoding

- A maximal empty rectangular space (ERS) is an empty rectangular space not contained in any other ERS.
- ERSs are generated and updated using the Difference Process of Lai and Chan (1997).
- When placing a rectangle, we limit ourselves only to maximal ERSs. We order all the maximal ERSs and place the rectangle in the first maximal ERS in which it fits.
- Let  $(x[i], y[i])$  be the coordinates of the bottom left corner of the  $i$ -th ERS.

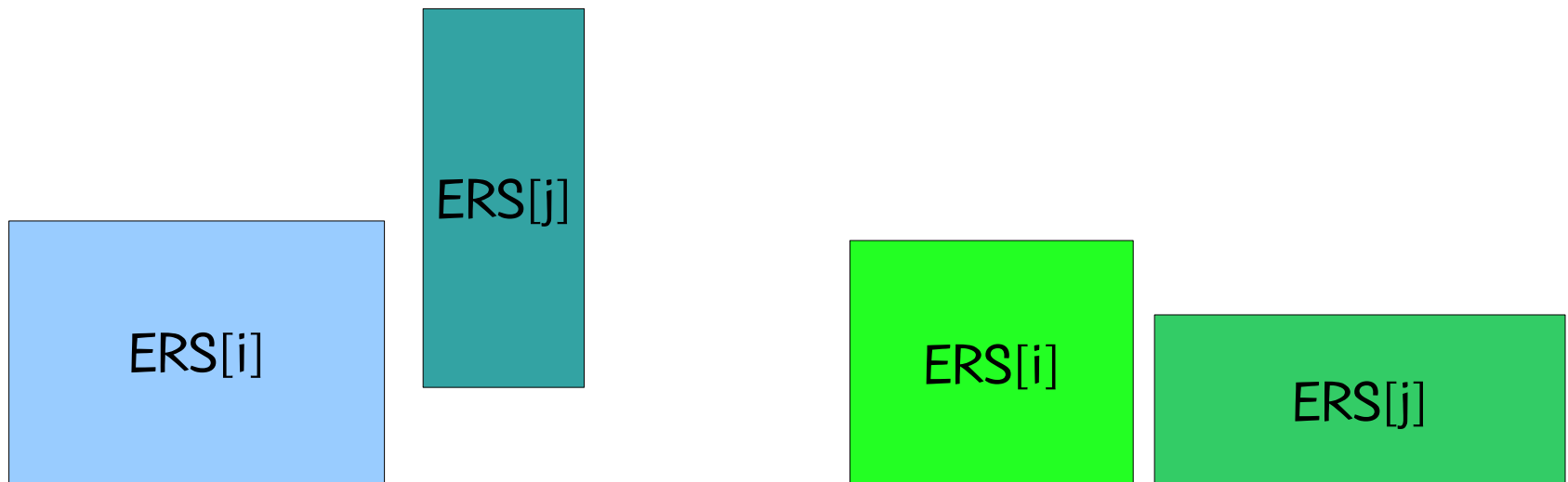
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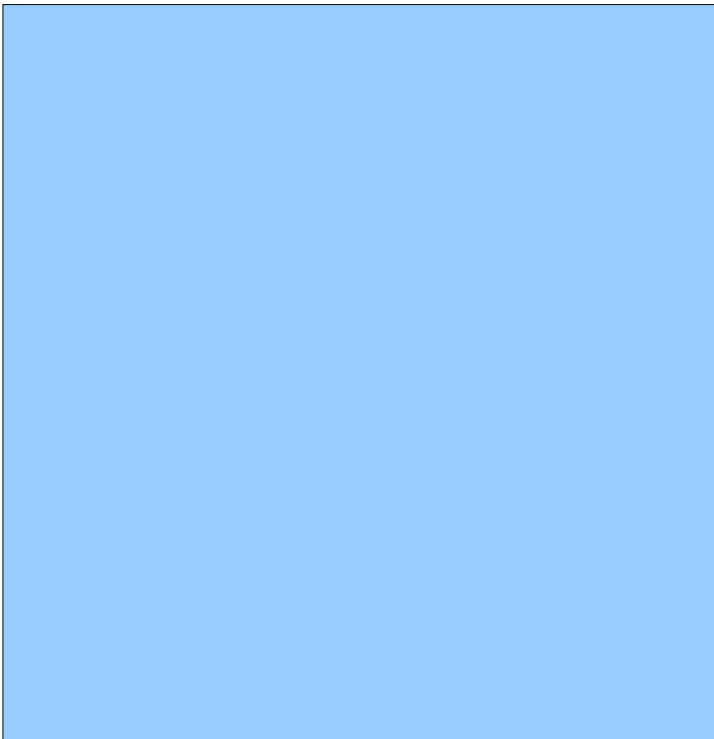
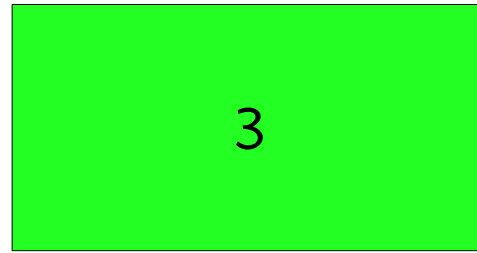
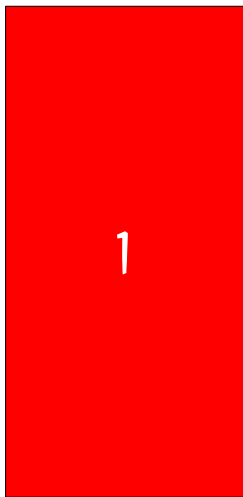
# Decoding

- If BL is used, ERSs are ordered such that  $ERS[i] < ERS[j]$  if  $y[i] < y[j]$  or  $y[i] = y[j]$  and  $x[i] < x[j]$ .



$ERS[i] < ERS[j]$

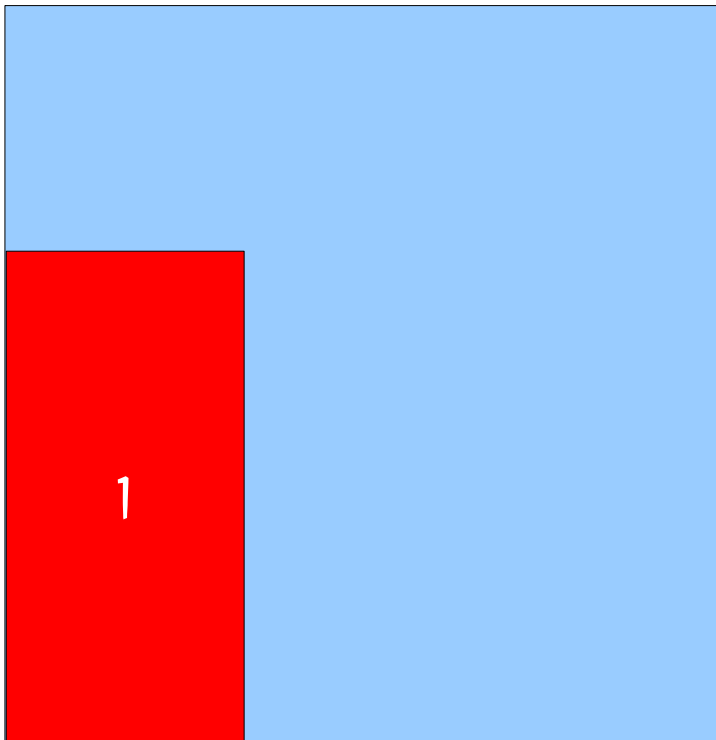
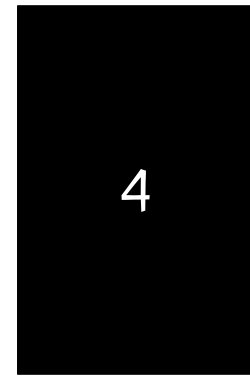
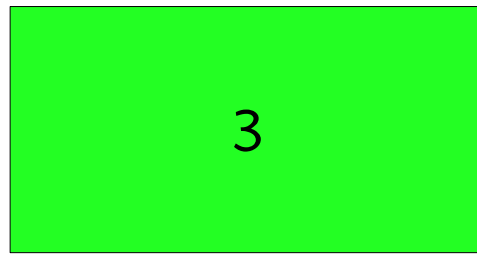




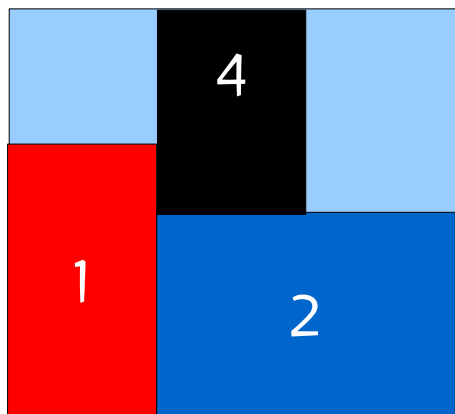
BL can run into problems even on small instances (Liu & Teng, 1999).

Consider this instance with 4 rectangles.

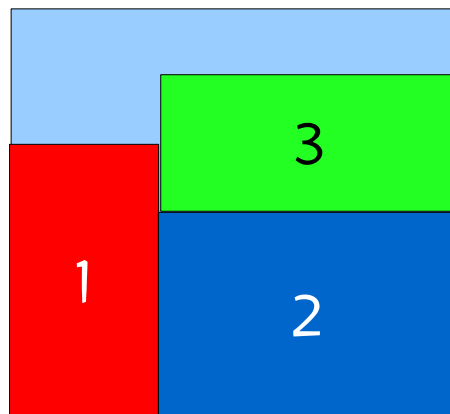
BL cannot find the optimal solution for any RTPS.



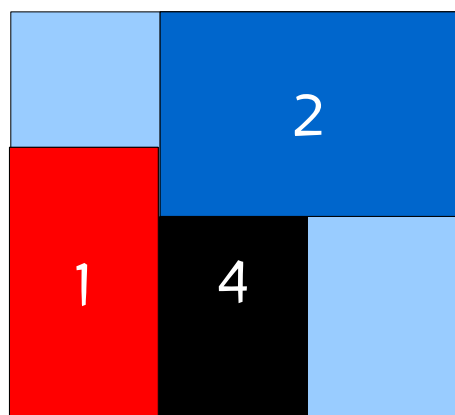
We show 6 rectangle type packing sequences (RTPS's) where we fix rectangle 1 in the first position.



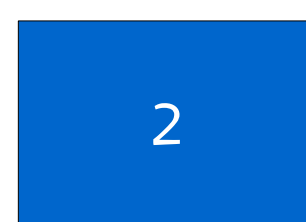
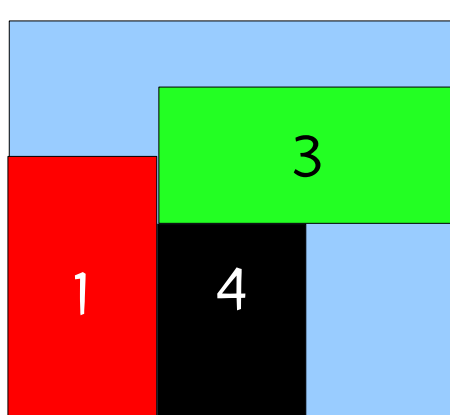
RTPS: 1-2-4-3



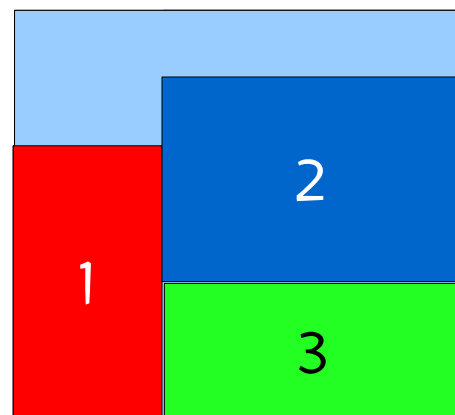
RTPS: 1-2-3-4



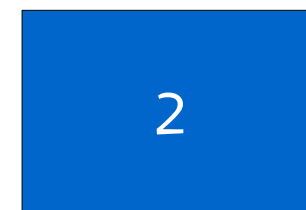
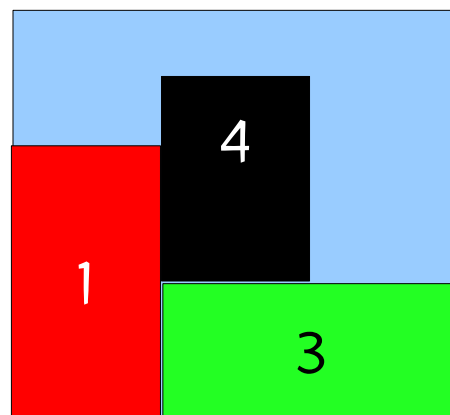
RTPS: 1-4-2-3



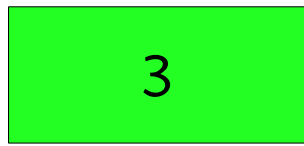
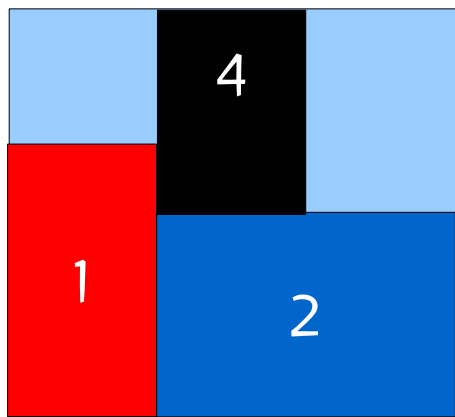
RTPS: 1-4-3-2



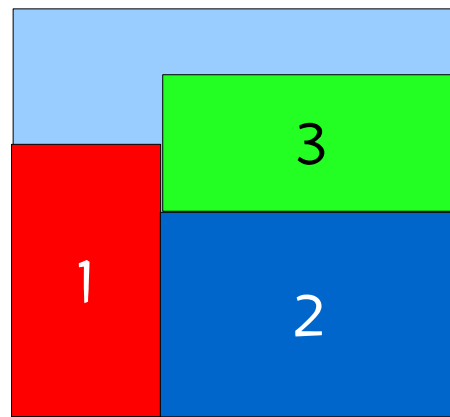
RTPS: 1-3-2-4



RTPS: 1-3-4-2

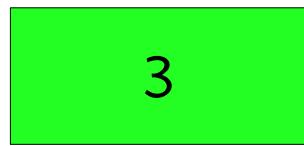
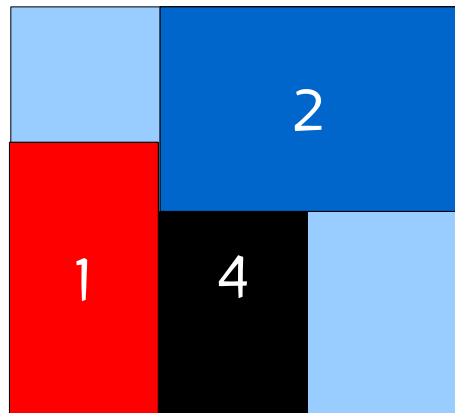


RTPS: 1-2-4-3

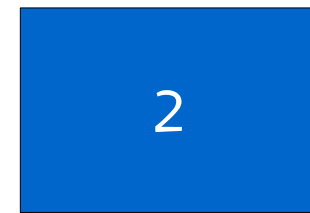
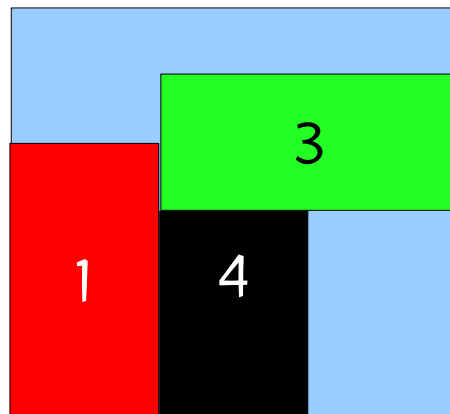


RTPS: 1-2-3-4

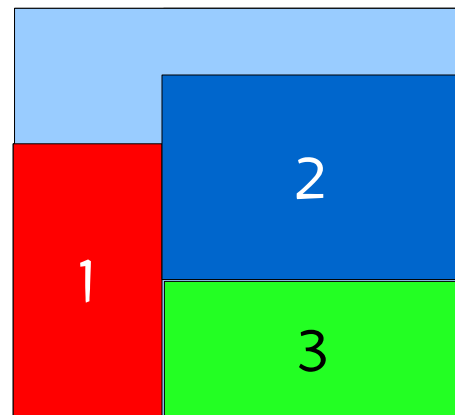
Similar infeasibilities are observed if 2, 3, or 4 is the first rectangle in the RTPS.



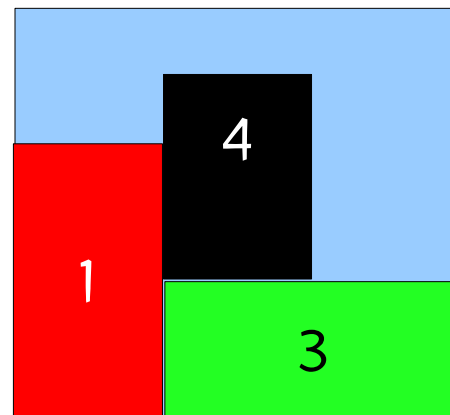
RTPS: 1-4-2-3



RTPS: 1-4-3-2



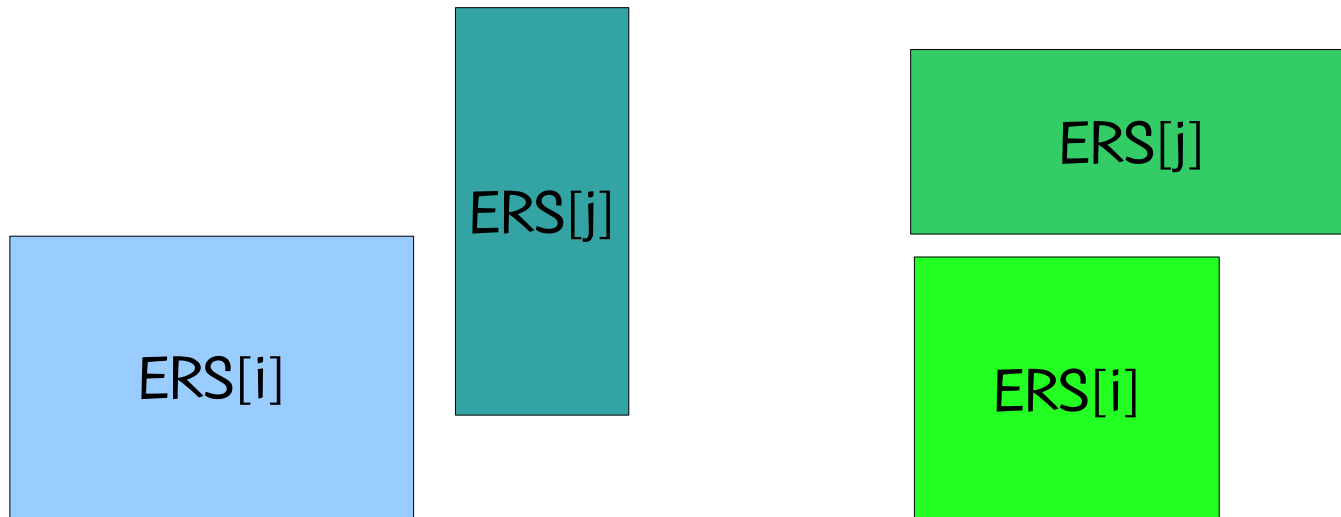
RTPS: 1-3-2-4



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# Decoding

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$ERS[i] < ERS[j]$

1  
BL

2  
BL

3  
LB

4  
BL

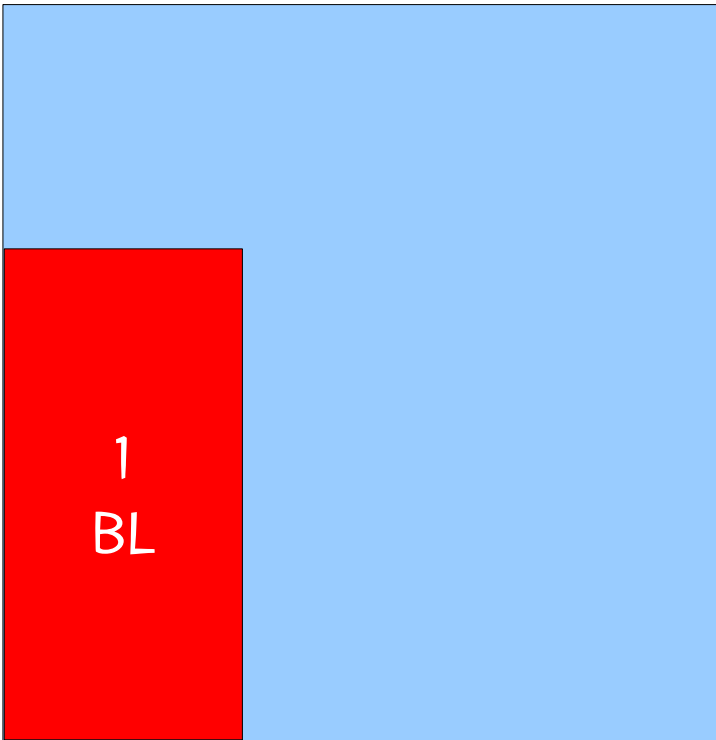
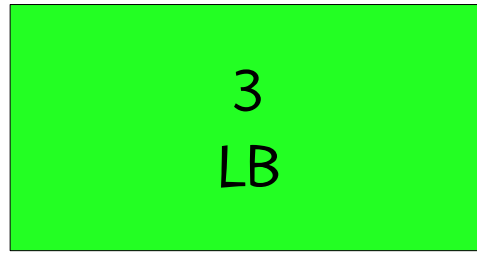
1  
BL

2  
BL

3  
LB

4  
BL

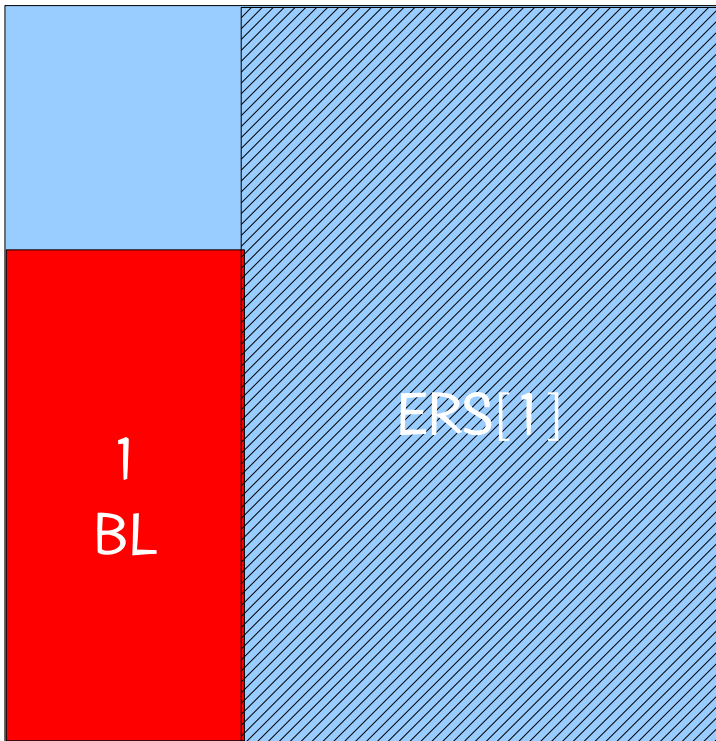
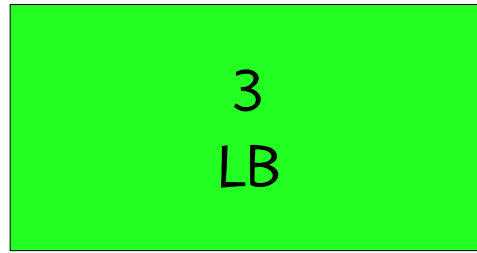
ERS[1]

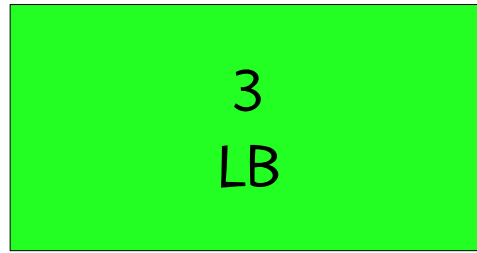


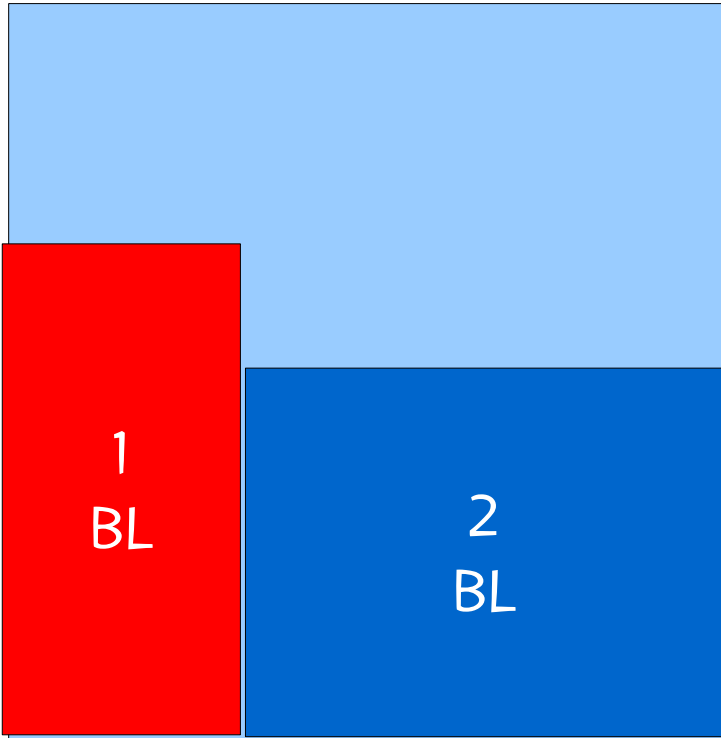
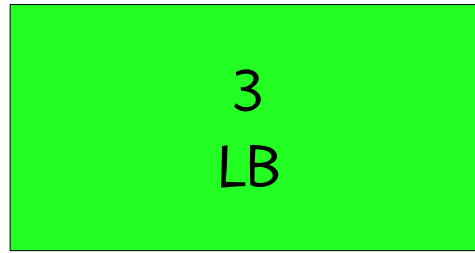
LION 7 ♣ January 2013

BRKGA







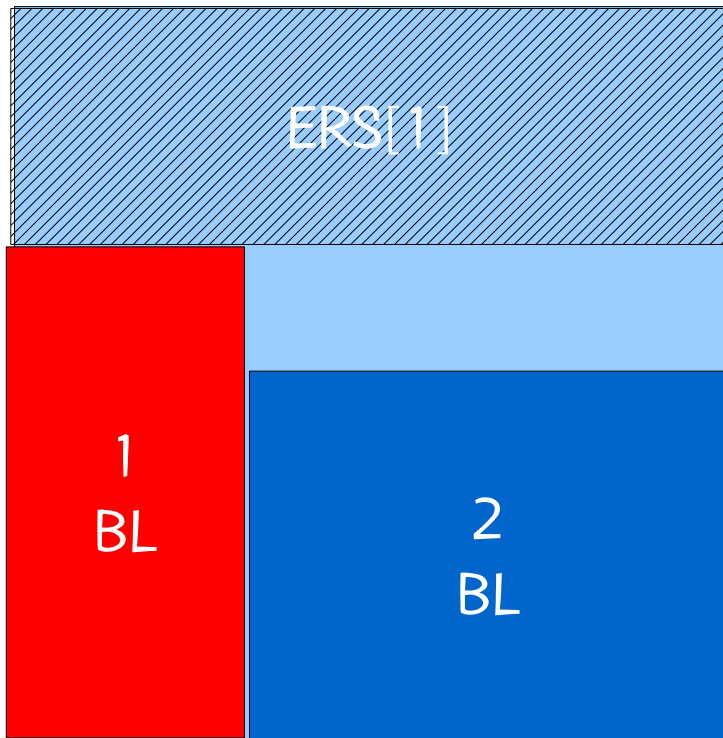


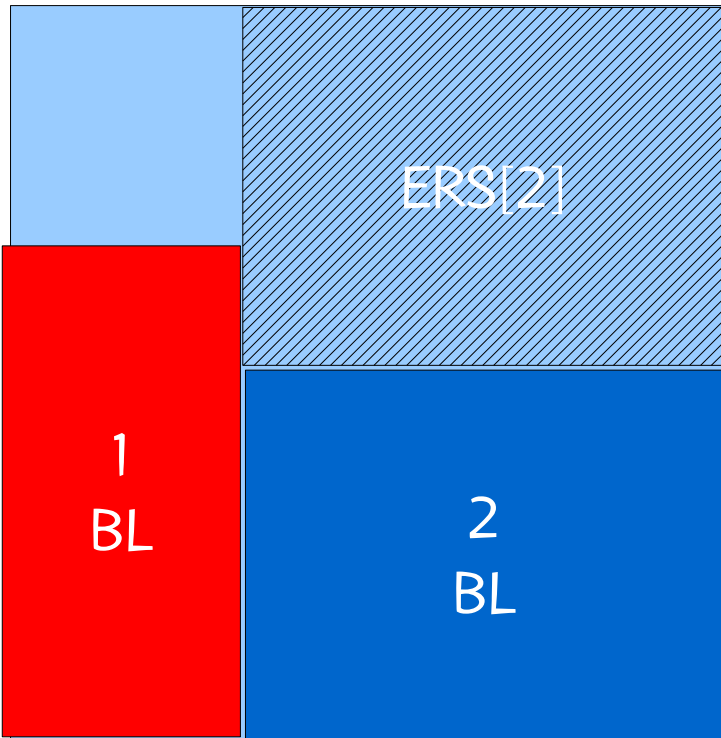
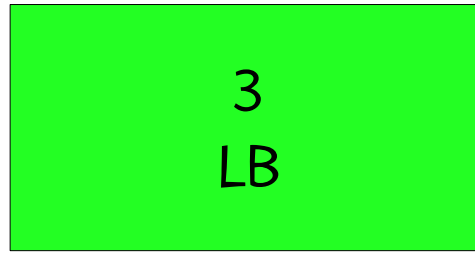
LION 7 ♣ January 2013

BRKGA

3  
LB

4  
BL





4  
BL

3  
LB

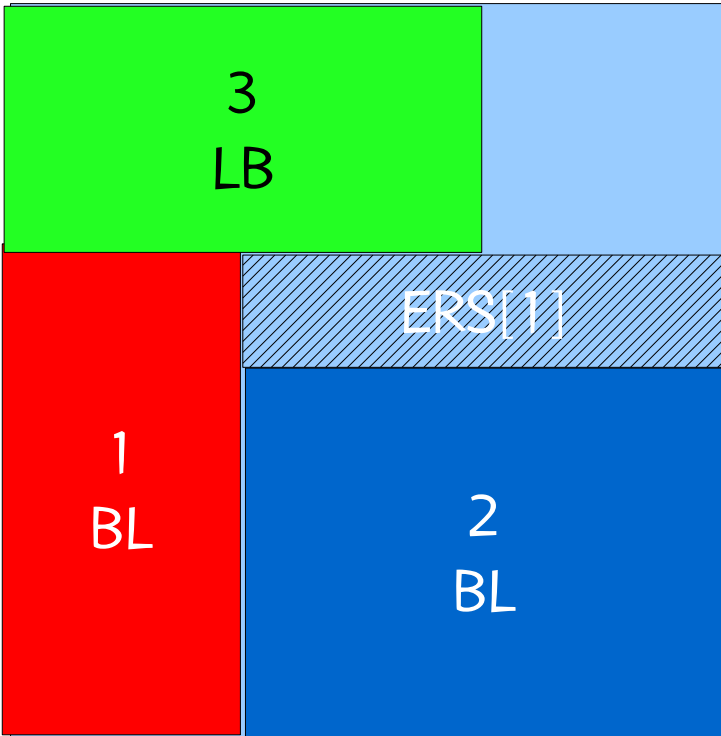
1  
BL

2  
BL

LION 7 ♣ January 2013

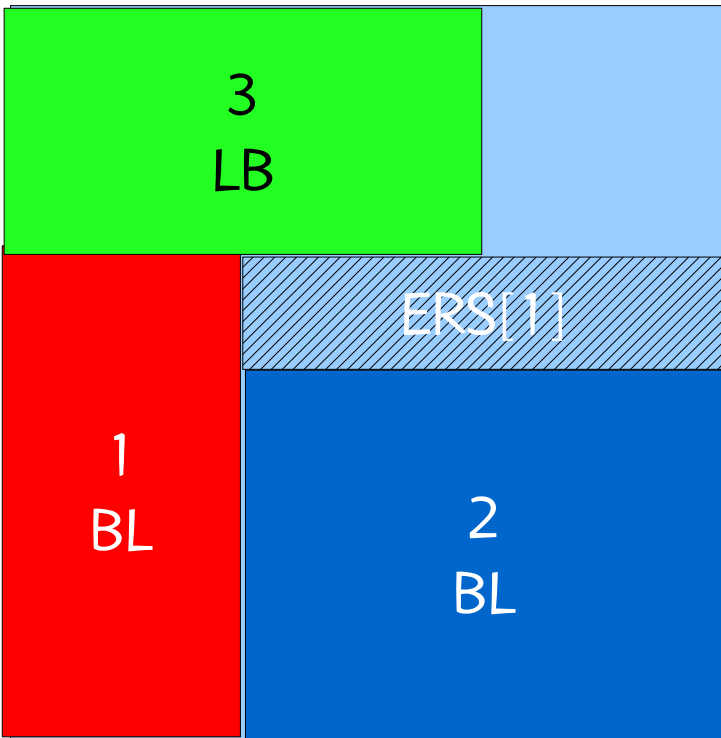
BRKGA

4  
BL



4  
BL

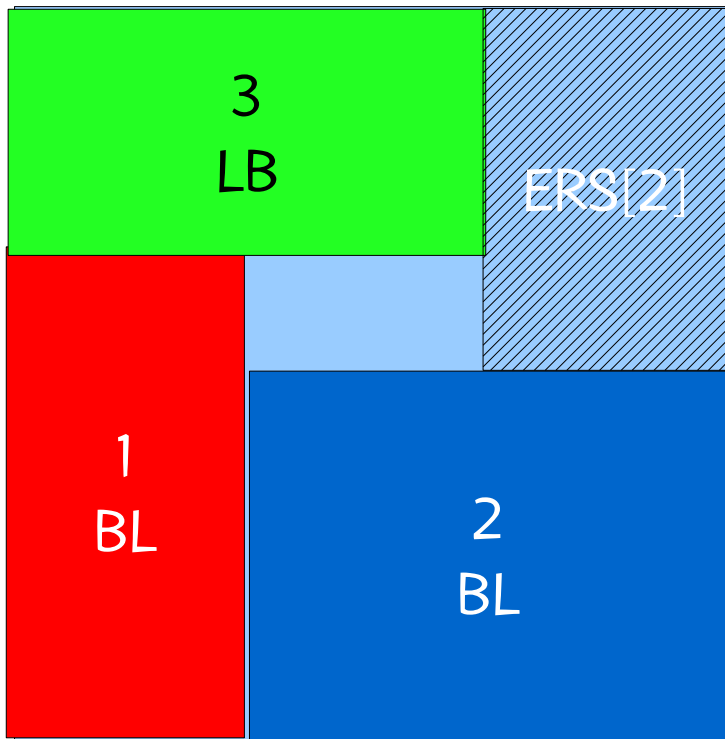
4 does not fit  
in ERS[1].

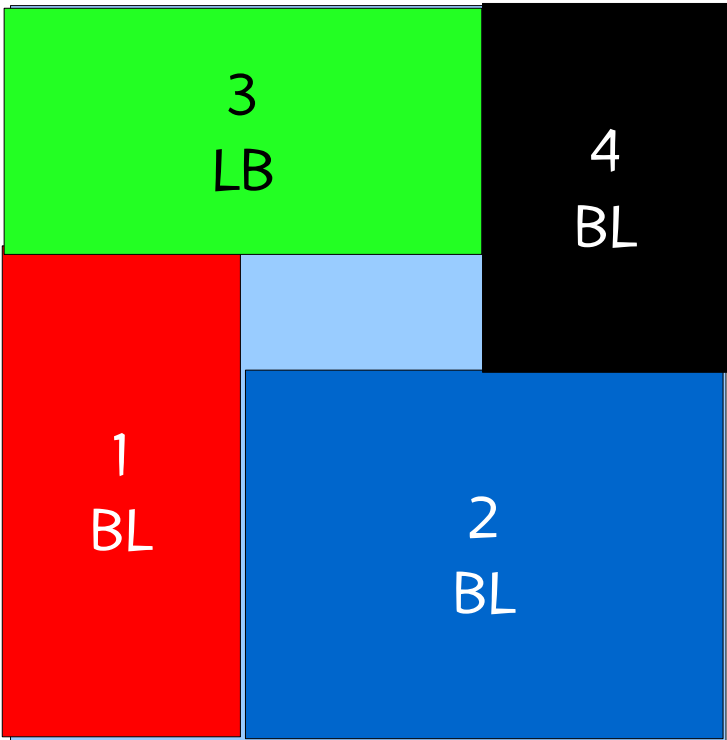






4 does fit  
in ERS[2].





Optimal solution!

# Experimental results

# Design

- We compare solution values obtained by the parallel multi-population BRKGA with solutions obtained by the heuristics that produced the best computational results to date:

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  - **TABU**: tabu search of Alvarez-Valdes et al. (2007)



# Number of best solutions / total instances

Problem	PH	GA	GRASP	TABU	BRKGA BL-LB-L-4NR
From literature (optimal)	13/21	<b>21/21</b>	18/21	<b>21/21</b>	<b>21/21</b>
Large random*	0/21	0/21	5/21	8/21	<b>20/21</b>
Zero-waste			5/31	17/31	<b>30/31</b>
Doubly constrained	11/21		12/21	17/21	<b>19/21</b>

\* For large random: number of best average solutions / total instance classes

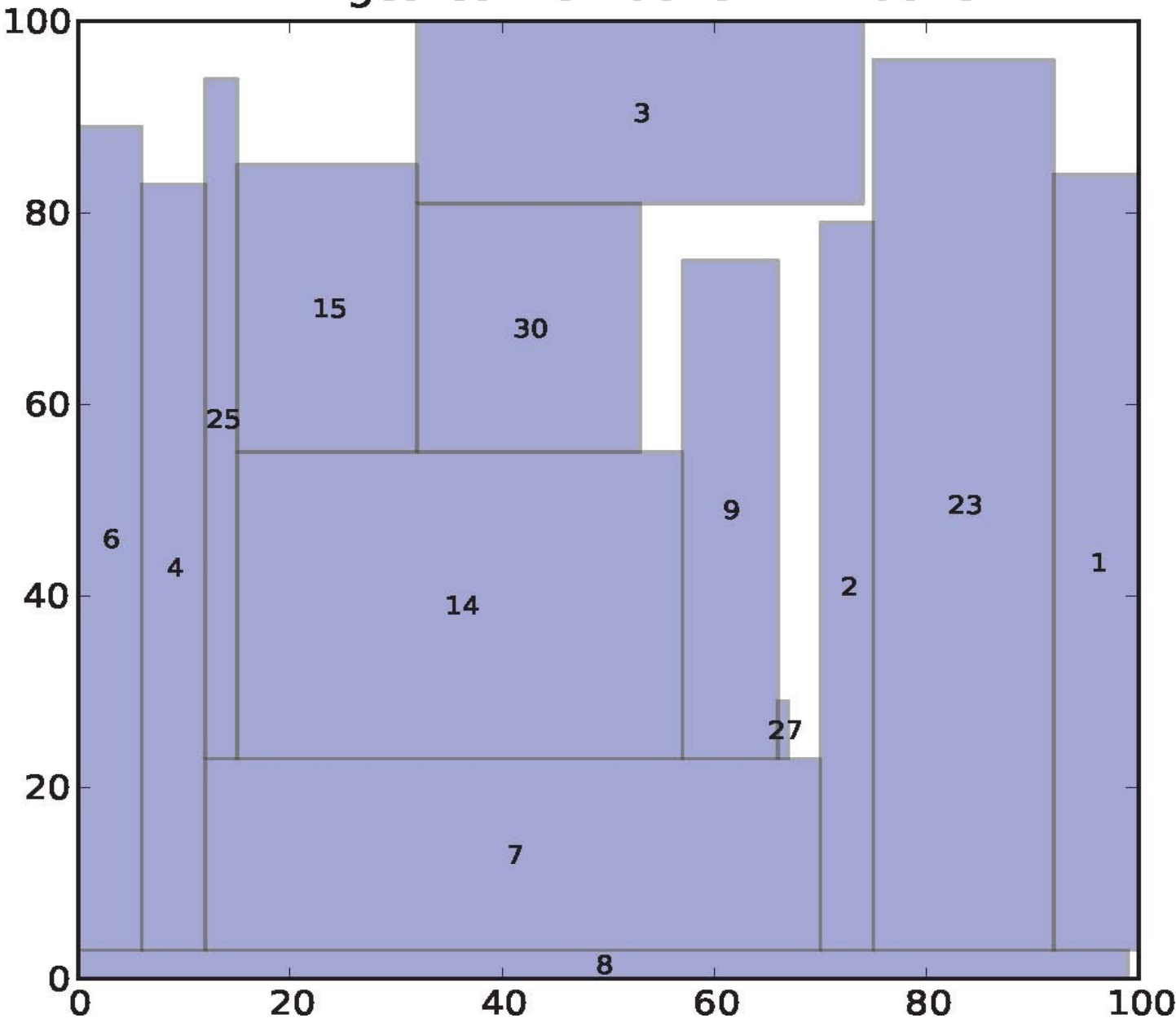
# Minimum, average, and maximum solution times (secs) for BRKGA (BL-LB-L-4NR)

Problem	Min solution time (secs)	Avg solution time (secs)	Max solution time (secs)
From literature (optimal)	0.00	0.05	0.55
Large random	1.78	23.85	72.70
Zero-waste	0.01	82.21	808.03
Doubly constrained	0.00	1.16	16.87

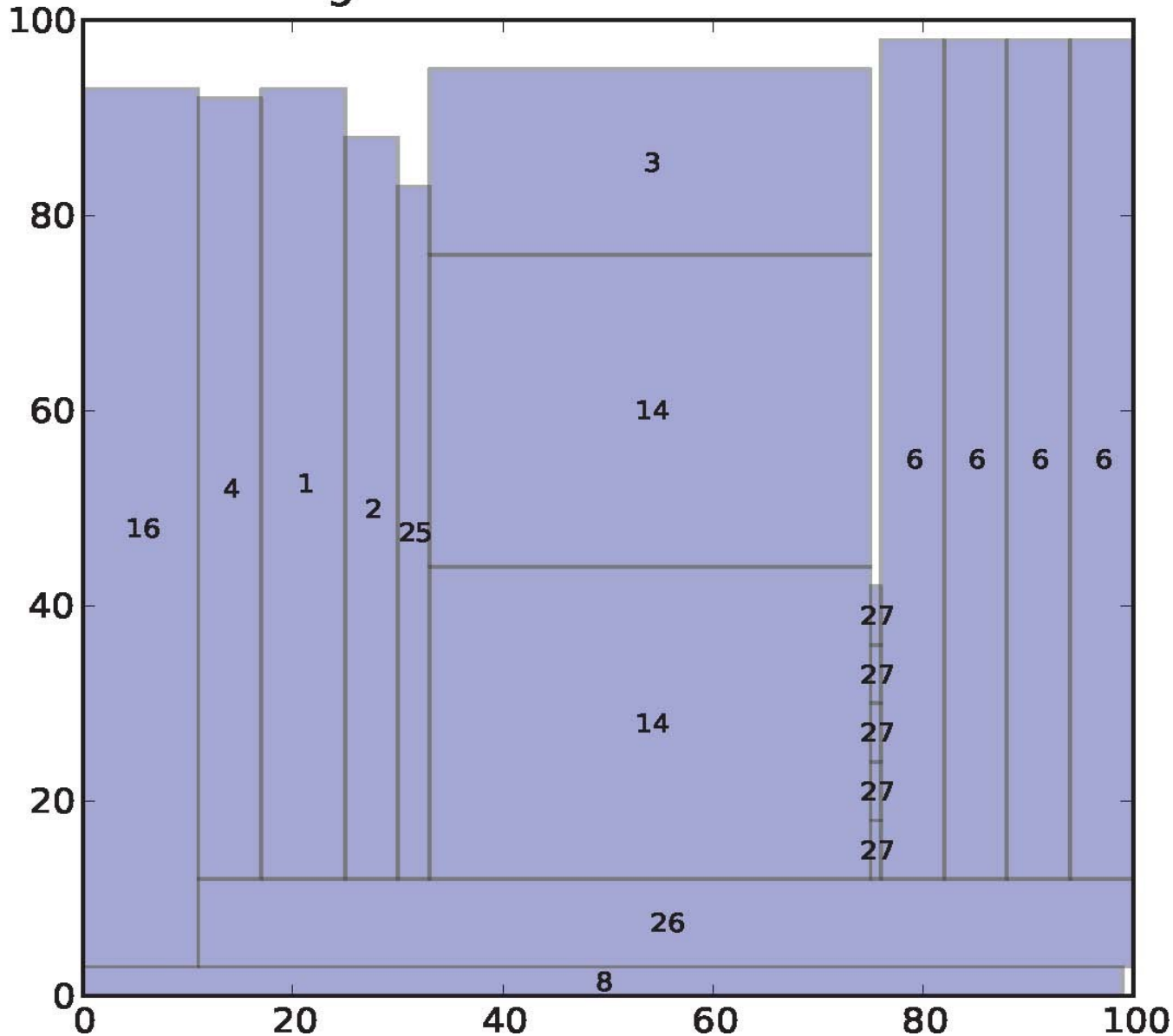
# 2D-ngcutcon18-20678.txt: 20678

New BKS  
for a 100 x100  
doubly  
constrained  
instance of  
Fekete &  
Schepers (1997)  
of value **20678**.  
Previous best  
was **19657** by  
tabu search of  
Alvarez-Valdes et  
al., (2007).

30 types  
30 rectangles



2D-ngcutcon21-22140-1.txt: 22140



New BKS for a 100 x 100 doubly constrained instance Fekete & Schepers (1997) of value **22140**.

Previous BKS was **22011** by tabu search of Alvarez-Valdes et al. (2007).

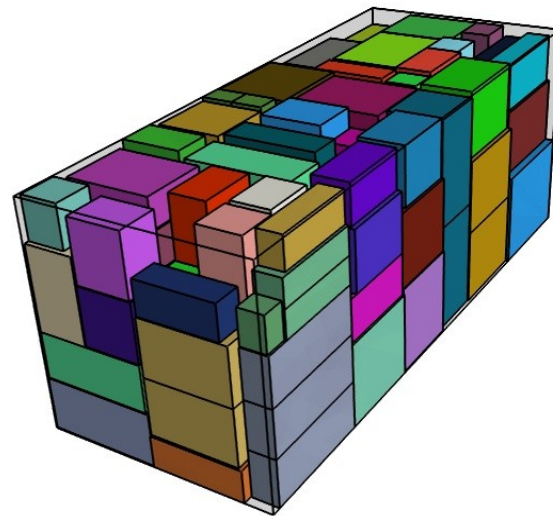
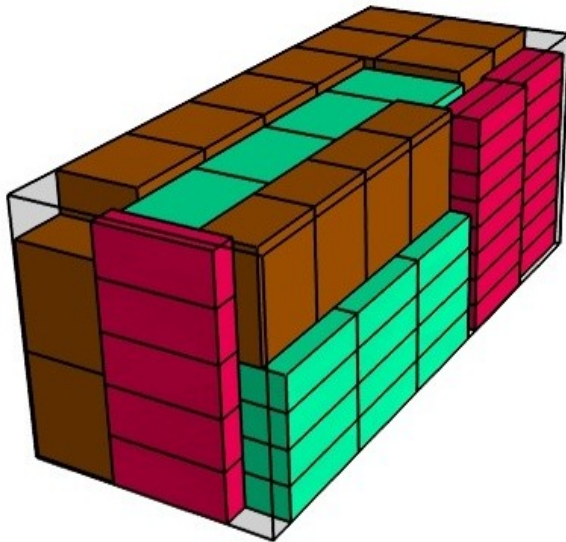
29 types  
97 rectangles

# Some remarks

We have extended this to 3D packing:

J.F. Gonçalves and M.G.C.R., "A parallel multi-population biased random-key genetic algorithm for a container loading problem," Computers & Operations Research, vol. 29, pp. 179-190, 2012.

Tech report: <http://www.research.att.com/~mgcr/doc/brkga-pack3d.pdf>



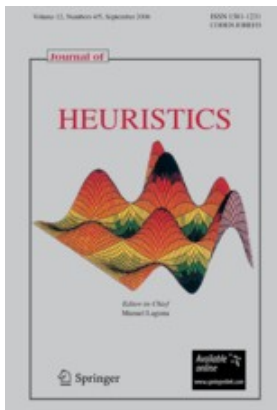
# Literature survey

# Literature

- BRKGAs have been applied in a wide range of areas.
- The following is a sampling of some papers that appeared in the literature applying BRKGAs.

# Survey

- Survey: Gonçalves and R. (2011)



J.F. Gonçalves and M.G.C.R., “**Biased random-key genetic algorithms for combinatorial optimization,**” J. of Heuristics, vol.17, pp. 487-525, 2011.



# Telecommunications

- Routing: Ericsson, R., Pardalos (2002), Buriol et al. (2002, 2005), Reis et al. (2011), Noronha, R., Ribeiro (2007, 2008, 2011), Heckeler et al. (2011)
- Design: Andrade et al. (2006), Buriol, R., Thorup (2007)
- Network monitoring: Breslau et al. (2011)
- Regenerator location: Duarte et al. (2011)
- Fiber installation in optical networks: Goulart et al. (2011)
- Path-based recovery in flexgrid optical networks: Castro et al. (2012)

# Telecommunications (cont'd)

- Handover minimization: [Morán-Mirabal et al. \(2012\)](#)
- Survivable IP/MPLS-over-WSN multi-layer network: [Ruiz et al. \(2011\)](#), [Pedrola et al. \(2011\)](#)
- Survey: [R. \(2012\)](#)

# Scheduling

- Job-shop scheduling: Gonçalves, Mendes, R. (2005), Gonçalves and R. (2012)
- Single machine scheduling: Valente et al. (2006), Valente and Gonçalves (2008)
- Resource constrained project scheduling: Gonçalves, Mendes, R. (2008, 2009), Gonçalves, R., Mendes (2011)
- Selection and scheduling of observations on Earth observing satellites: Tangpattanakul, Josefowicz, Lopez (2012)

# Production planning

- Assembly line balancing: Gonçalves and Almeida (2002)
- Manufacturing cell formation: Gonçalves and R. (2004)
- Single machine scheduling: Valente et al. (2006), Valente and Gonçalves (2008)
- Assembly line worker assignment and balancing: Moreira et al. (2010)
- Lot sizing and scheduling with capacity constraints and backorders: Gonçalves and Sousa (2011)

# Network optimization

- Concave minimum cost flow: [Fontes and Gonçalves \(2007\)](#)
- Robust shortest path: [Coco, Noronha, Santos \(2012\)](#)
- Tree of hubs location: [Pessoa, Santos, R. \(2012\)](#)
- Hop-constrained trees in nonlinear cost flow networks: [Fontes and Gonçalves \(2012\)](#)
- Capacitated arc routing: [Martinez, Loiseau, R. \(2011\)](#)

# Power systems

- Unit commitment: Roque, Fontes, Fontes (2010, 2011)
- Multi-objective unit commitment: Roque, Fontes, Fontes (2012)

# Packing

- 2D orthogonal packing: [Gonçalves and R. \(2011\)](#)
- 3D container loading: [Gonçalves and R. \(2012a\)](#)
- 2D/3D bin packing: [Gonçalves and R. \(2012b\)](#)

# Covering

- Steiner triple systems: [R. et al. \(2012\)](#)
- Covering by pairs: [Breslau et al. \(2011\)](#)



# Transportation

- Tollbooth assignment: [Buriol. et al. \(2009, 2010\)](#)

# Auctions

- Combinatorial auctions: [Andrade et al. \(2012\)](#)

# Automatic parameter tuning

- GRASP with path-relinking: Festa et al. (2010)
- GRASP with evolutionary path-relinking: Morán-Mirabal, González-Velarde, R. (2012)

# Continuous global optimization

- Bound-constrained optimization: Silva, Pardalos, R. (2012)

# Software

- C++ API: Toso and R. (2012)

# Thanks!

These slides and all of the papers cited in this lecture can be downloaded from my homepage:

<http://www.research.att.com/~mgcr>