A biased random-key genetic algorithm for a prize-collecting directed Steiner forest network design problem

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Joint work with

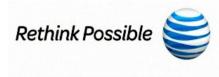
C.E. de Andrade & F.K. Miyazawa (UNICAMP, Brazil)

R.D. Doverspike, K. Reichmann, R.K. Sinha & W. Zhang (AT&T Labs Research, USA)



Summary

- Prize collecting directed k-hop Steiner forest (PCk-HSF) problem
- Wireless backhaul network planning as a PCk-HSF problem with additional constraints
- Biased random-key genetic algorithms (BRKGA)
- BRKGA for wireless backhaul network planning focusing on the decoder
- Application of the BRKGA to a "real" instance of wireless backhaul network planning
- Concluding remarks

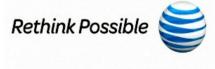


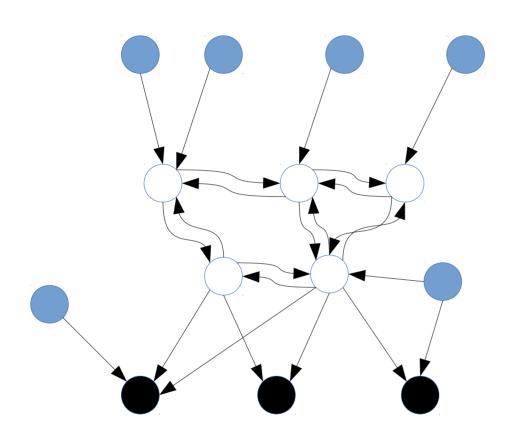
- Let G = (V,E) be a given directed graph
- Let V_r , V_s , and V_d partition the node set, i.e.

$$V_r \cup V_s \cup V_d = V$$
 and $V_r \cap V_s \cap V_d = \emptyset$

- V_r set of root nodes
- V_s set of Steiner nodes
- V_d set of demand nodes

- Let G = (V,E) be a given directed graph
- Arcs are directed
 - From demand nodes to Steiner nodes and root nodes
 - From Steiner nodes to root nodes
- For each pair of nodes v, $u \in V_s$
 - $-(u,v) \in E$
 - $-(v,u) \in E$





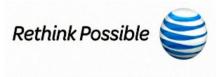


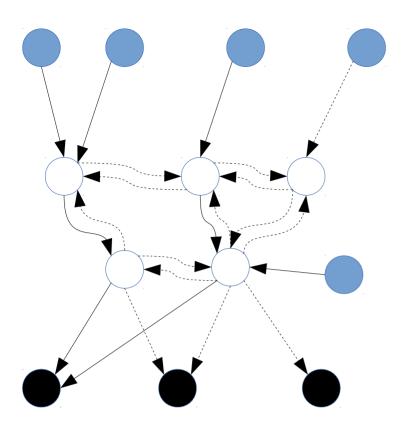
Steiner node

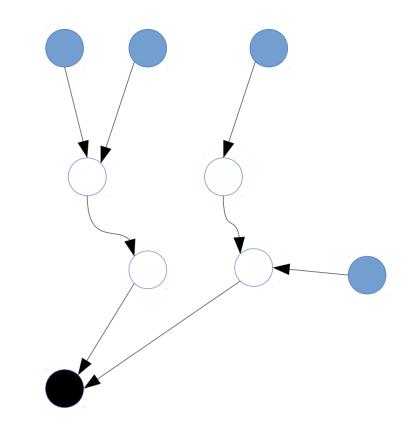
Root node

- A Directed Steiner Tree $T = (V[T] \subseteq V, E[T] \subseteq E)$ with root in $r \in V_r[T]$ is a loopless connected subgraph of G with a unique root r.
 - For each demand node $u \in V[T]$ there is a unique path in T from u to r.

- A k-Hop Directed Steiner Tree is a directed Steiner tree
 (connected subgraph of G with a unique root r) such that:
 - Any path from a demand node u to the root r has no more than k+2 nodes, including u and r.
- A k-Hop Directed Steiner Forest is a collection of disjoint k-Hop Directed Steiner Trees



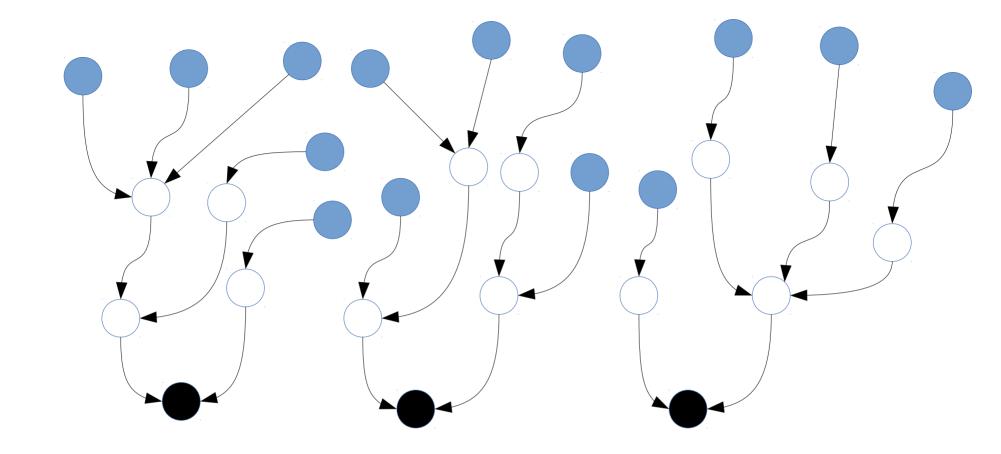




Acyclic directed graph

2-Hop Directed Steiner tree

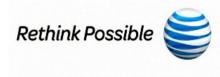




2-Hop Directed Steiner Forest

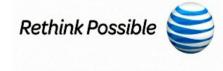


- A demand function d: V_d → R⁺ gives the prize to be collected from each demand node if it is a leaf on the tree.
- A cost function c: V_s → IR⁺ gives the cost of of each
 Steiner vertex.



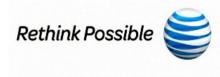
- Given a directed graph G = (V,E), vertex partition V_r , V_s , and V_d , hop parameter k, demand function d, and cost function c
- FIND: A k-hop directed Steiner forest F such that the profit

sum [$v \in V_d[F]$] d_v — sum [$v \in V_s[F]$] c_v is maximized.



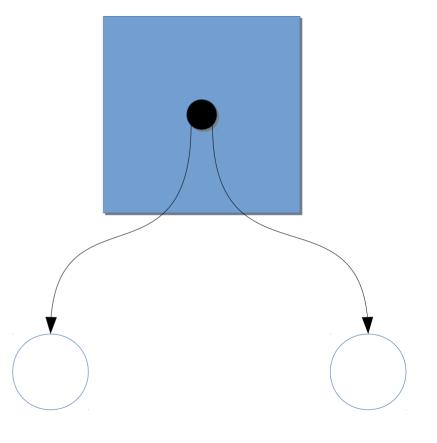
Given a geographical region where locations are represented as lat-long coordinates, where

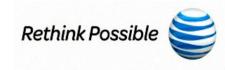
- V_d is the set of nodes representing traffic origination points (demand points) in the region
- V_s is the set of nodes representing locations where equipment for traffic collection and routing is located (e.g. utility poles)
- $-V_r$ is the set of nodes representing fibered access points (FAP), e.g. remote terminals (RT), macrocell (MC), or central offices.



Constraints: Demand splitting

- Estimate for each block total demand is placed in center of block
- Block can be served by one or more antennae so demand can be split among them
- Because of this undirected cycles can be introduced resulting in a directed acyclic graph (DAG)





Constraints: Access equipment action radii & capacities

Action radii

- Wi-Fi: 100 m

- LTE: 400 m

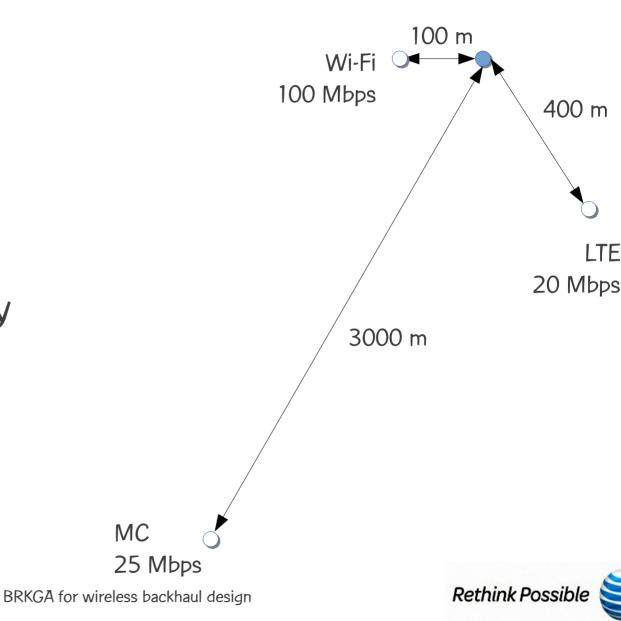
macrocell: 3 km

Processing capacity

- Wi-Fi: 100 Mbps

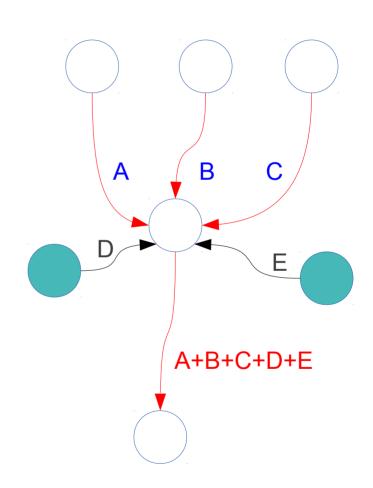
LTE: 20 Mbps

macrocell: 25 Mbps

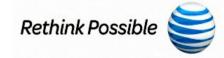


Constraints: Retransmitter equipment capacity

- Action radii: 1 km
- Processing capacity
 - Flow that equipment receives from other wireless retransmitters plus flow it sends to other retransmitters is limited to at most 100 Mbps



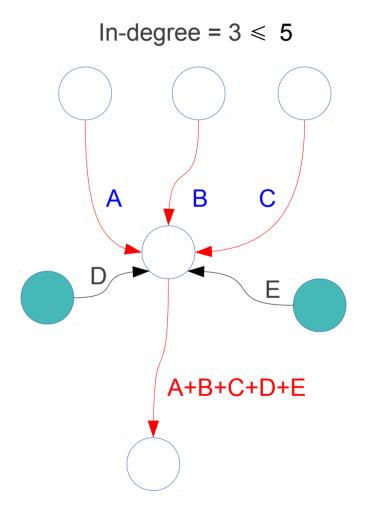
A+B+C+A+B+C+D+E ≤100



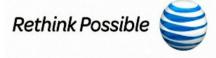
Constraints: Retransmitter equipment capacity

Fan-in constraint

 A limited number (5) of neighboring transmission equipment can flow traffic into equipment



A+B+C+A+B+C+D+E ≤100

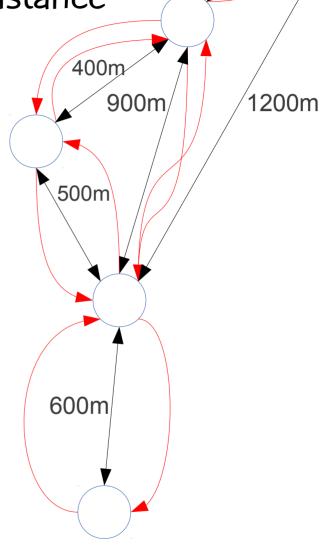


Constraints: Line of sight and minimum distance

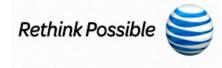
PCSF Problem assumed, for each pair $u, v \in V_s$ that $(u,v) \in E$ and $(v,u) \in E$

Not so in wireless backhaul planning: $(u,v) \in E$ and $(v,u) \in E$ only if

 Poles u and v are within 1000 m of each other



400m

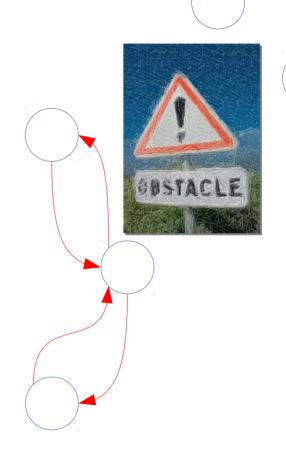


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PCSF Problem assumed, for each pair $u, v \in V_s$ that $(u,v) \in E$ and $(v,u) \in E$

Not so in wireless backhaul planning: $(u,v) \in E$ and $(v,u) \in E$ only if

- Poles u and v are within 1000 m of each other
- Poles u and v are in each other's line of sight

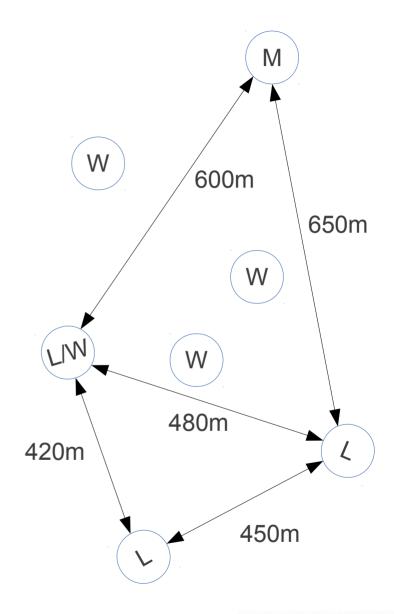


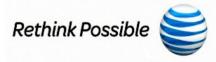


Constraints: Interference

LTE and MC use licensed spectrum and can interfere with each other.

- Pairs of LTE antennae must be separated by at least 400m
- LTE and macrocells by at least 500m
- No constraint on Wi-Fi exists since it uses non-licensed spectrum





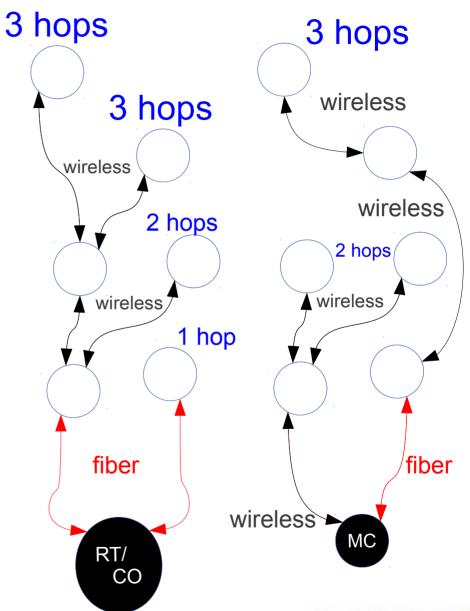
Constraints: k-Hops

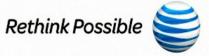
First hop is from FAP (root) to pole

- If root is Central Office (CO) or RT link must use fiber
- If root is macrocell link can be fiber or wireless

All other links are wireless

Number of hops is limited to k = 2 or 3 (in case first link is fiber)

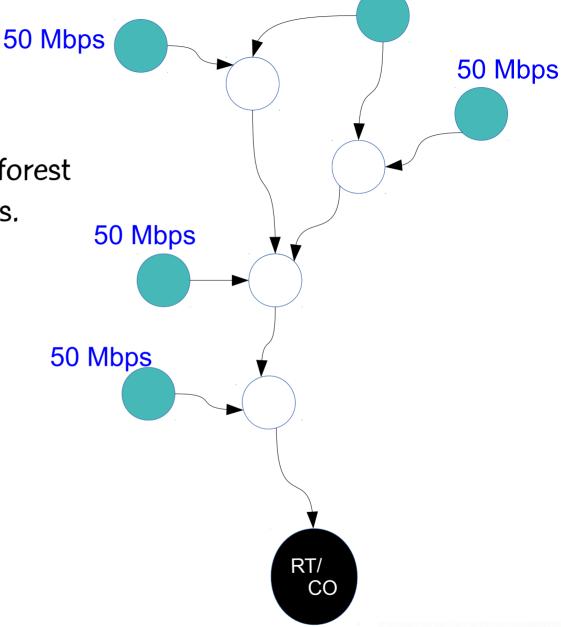


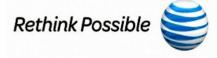


Constraints: Traffic flow

Traffic that is backhauled from demand points to root nodes of forest is limited by equipment capacities.

Only traffic that reaches roots is counted as revenue.



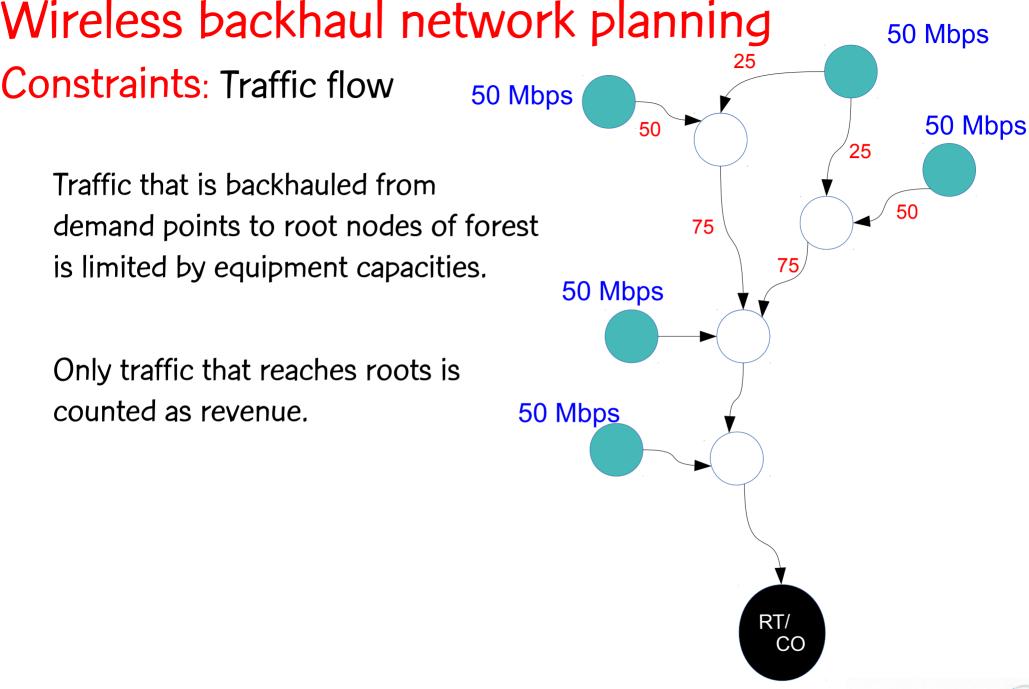


50 Mbps

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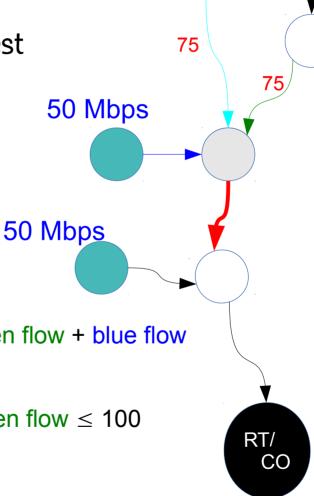
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Flow conservation: red flow = cyan flow + green flow + blue flow

Capacity constraint: red flow + cyan flow + green flow ≤ 100



50

50 Mbps

25

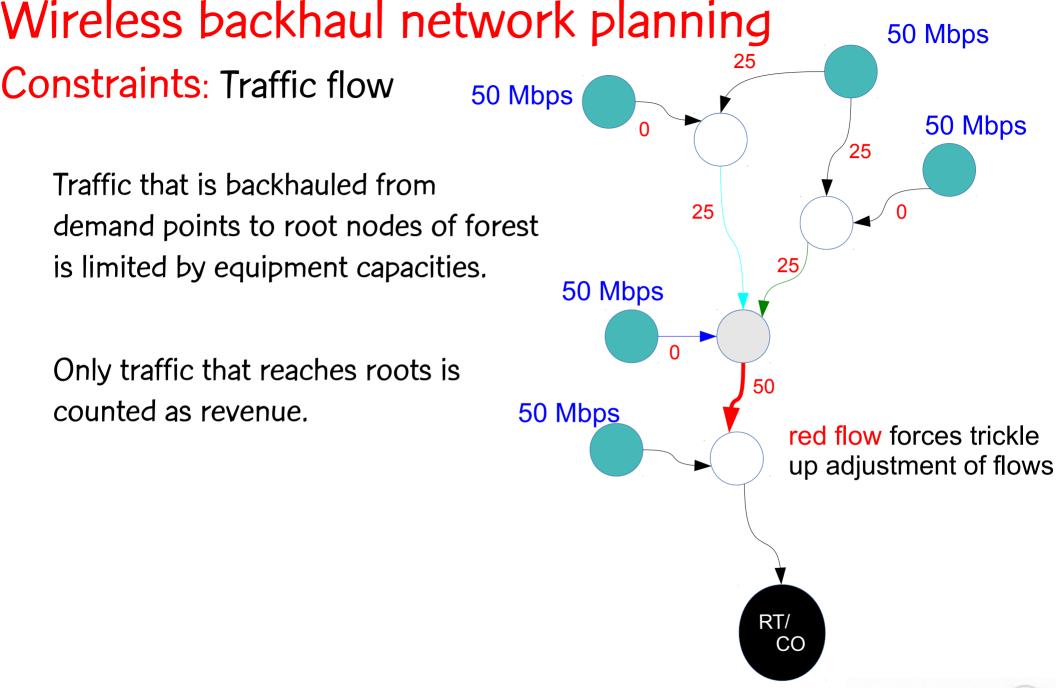
50 Mbps

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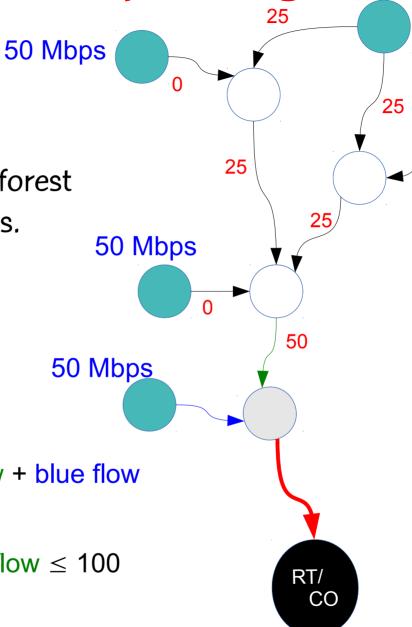
Constraints: Traffic flow

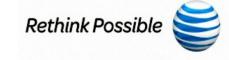
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50 Mbps

50 Mbps

Constraints: Traffic flow

Traffic that is backhauled from demand points to root nodes of forest is limited by equipment capacities.

Only traffic that reaches roots is counted as revenue.

50 Mbps

red flow forces trickle up adjustment of flows

75

RT/ CO

Flow conservation: red flow = green flow + blue flow = 75

Capacity constraint: red flow + green flow = 100 ≤ 100



50 Mbps

12.5

50 Mbps

50 Mbps

0

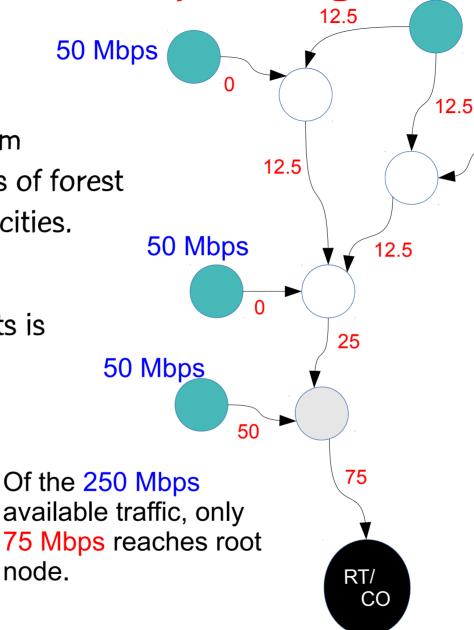
12.5

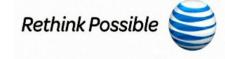
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Wireless backhaul network planning

Only traffic that reaches roots is counted as revenue.

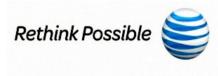




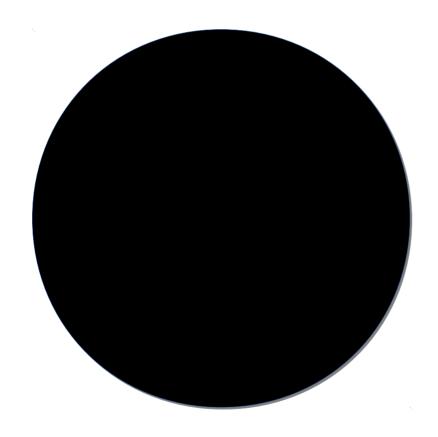
50 Mbps

50 Mbps

node.



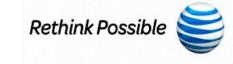
Holland (1975)

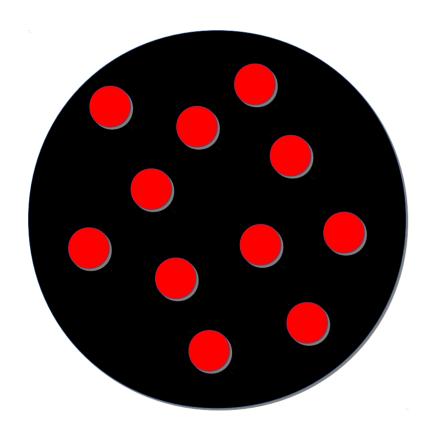


Adaptive methods that are used to solve search and optimization problems.

Individual: solution

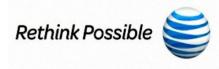


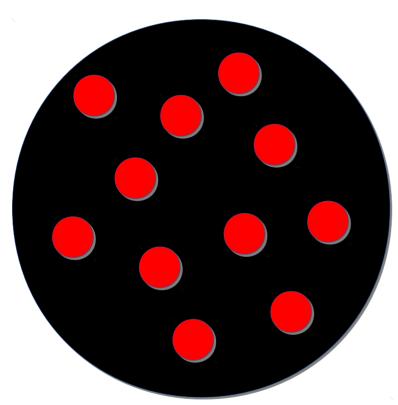




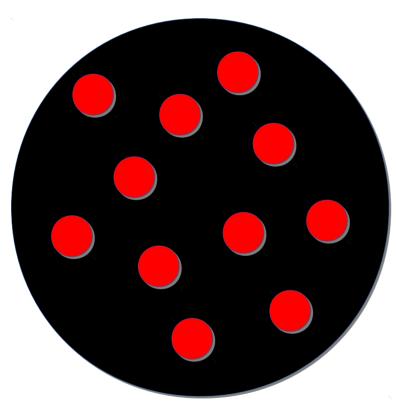
Individual: solution (chromosome = string of genes)
Population: set of fixed number of individuals

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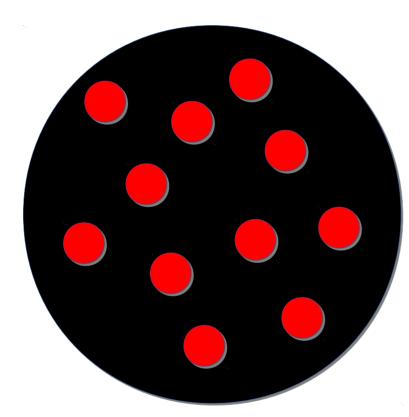


Genetic algorithms evolve population applying Darwin's principle of survival of the fittest.



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

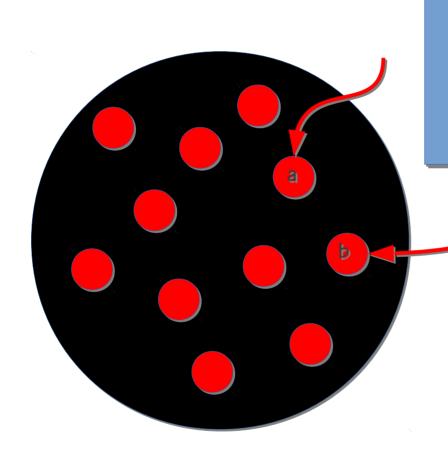
A series of generations are produced by the algorithm. The most fit individual of the last generation is the solution.



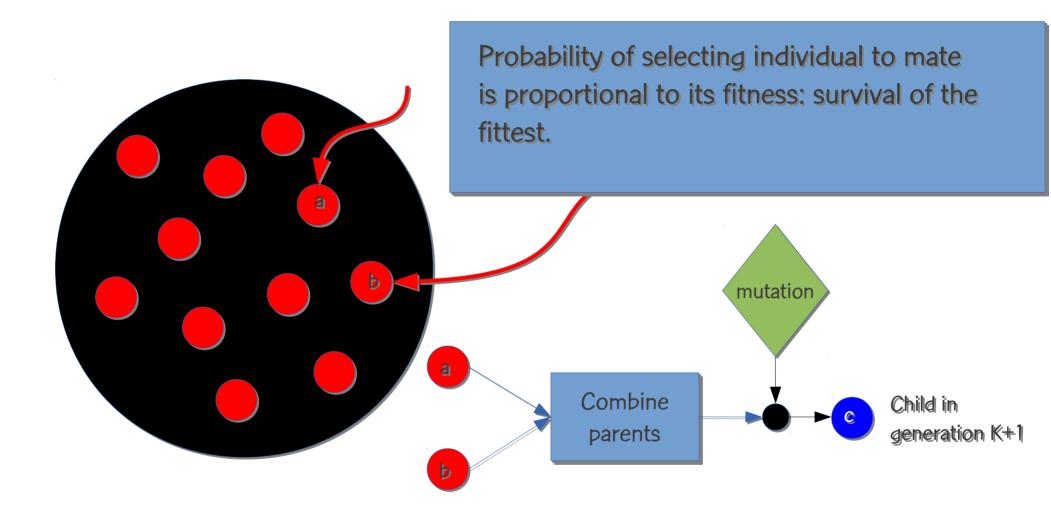
Genetic algorithms evolve population applying Darwin's principle of survival of the fittest.

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

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



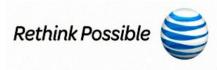
Probability of selecting individual to mate is proportional to its fitness; survival of the fittest.



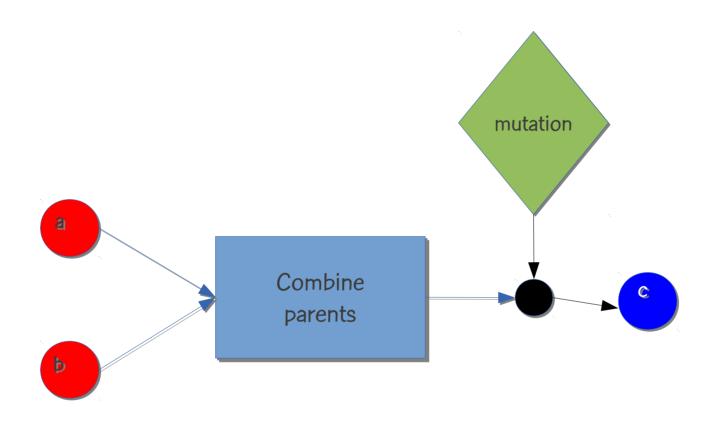
Parents drawn from generation K

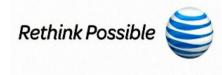
BRKGA for wireless backhaul design

C

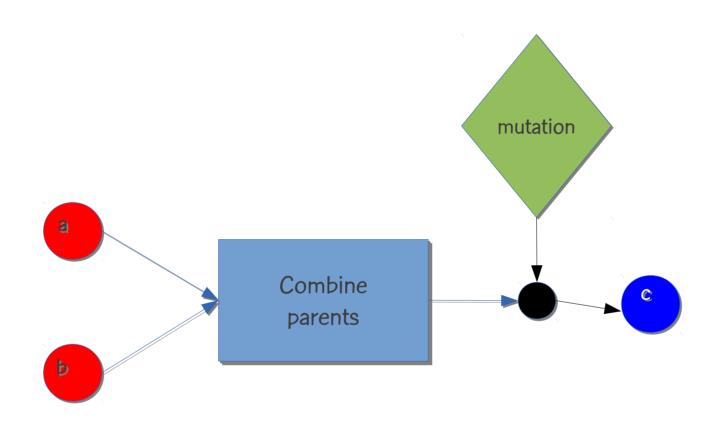


Crossover and mutation



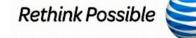


Crossover and mutation

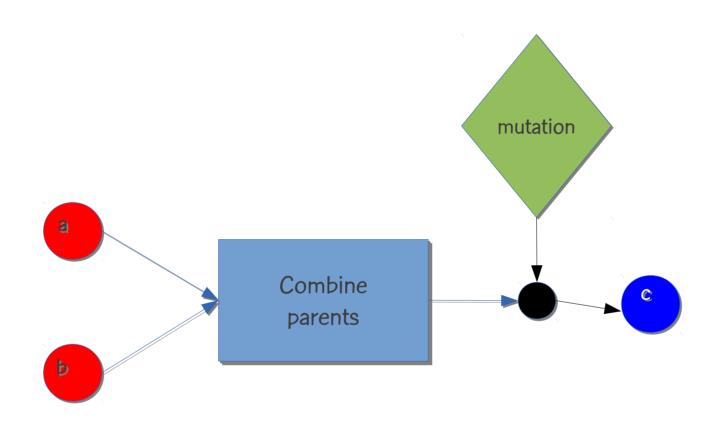


Crossover: Combines parents ... passing along to offspring characteristics of each parent ...

Intensification of search



Crossover and mutation

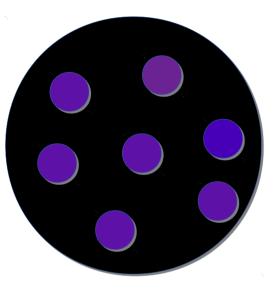


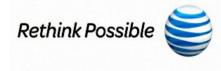
Mutation: Randomly changes chromosome of offspring ...

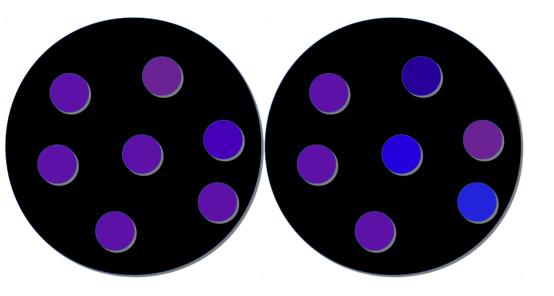
Driver of evolutionary process ...

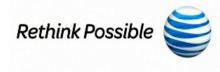
Diversification of search

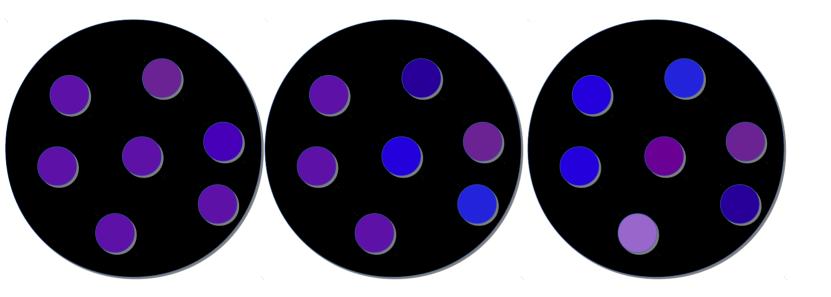


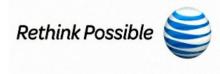


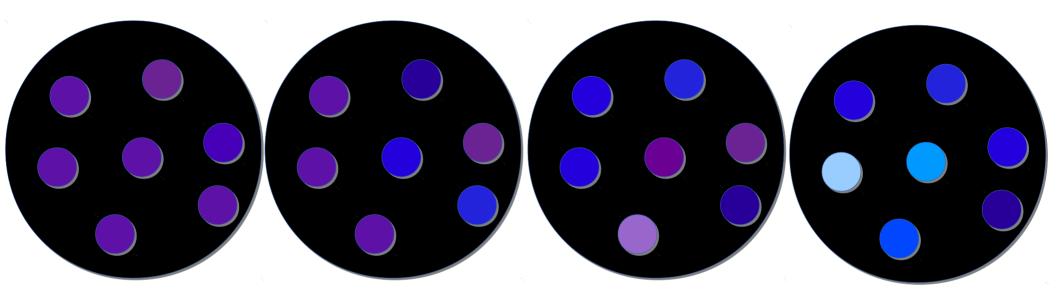


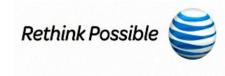


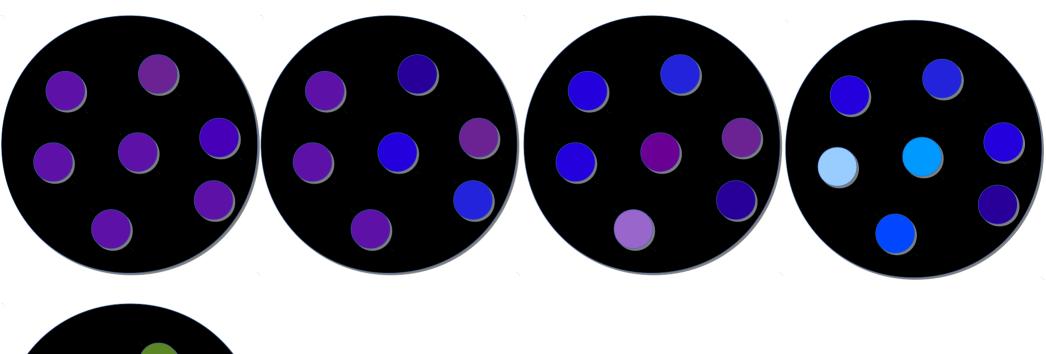


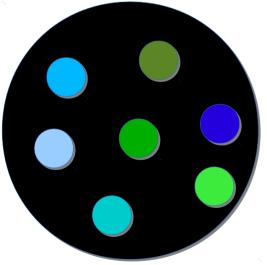




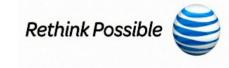


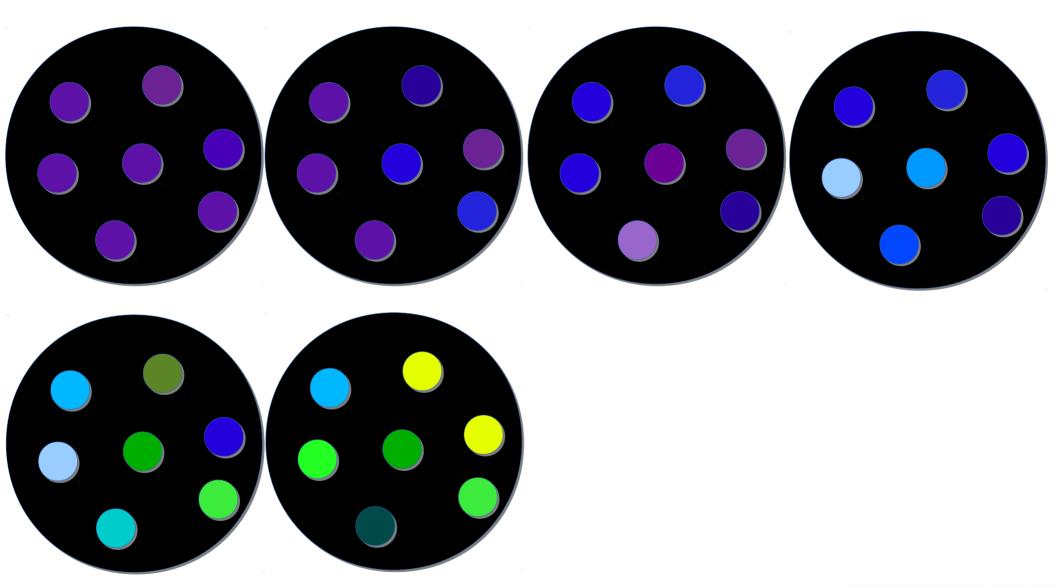




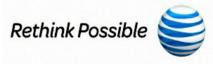


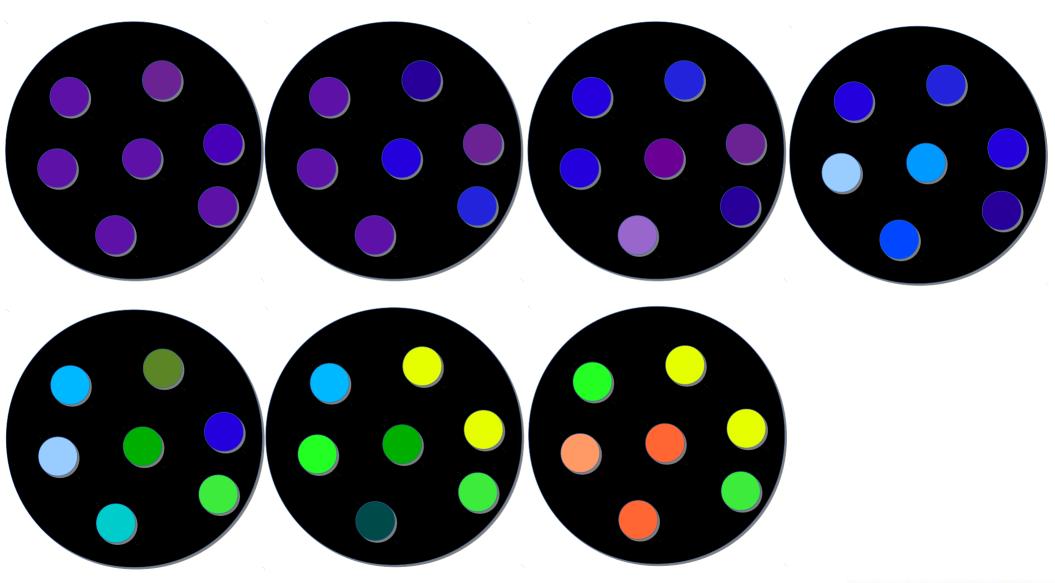
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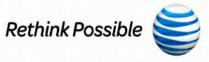


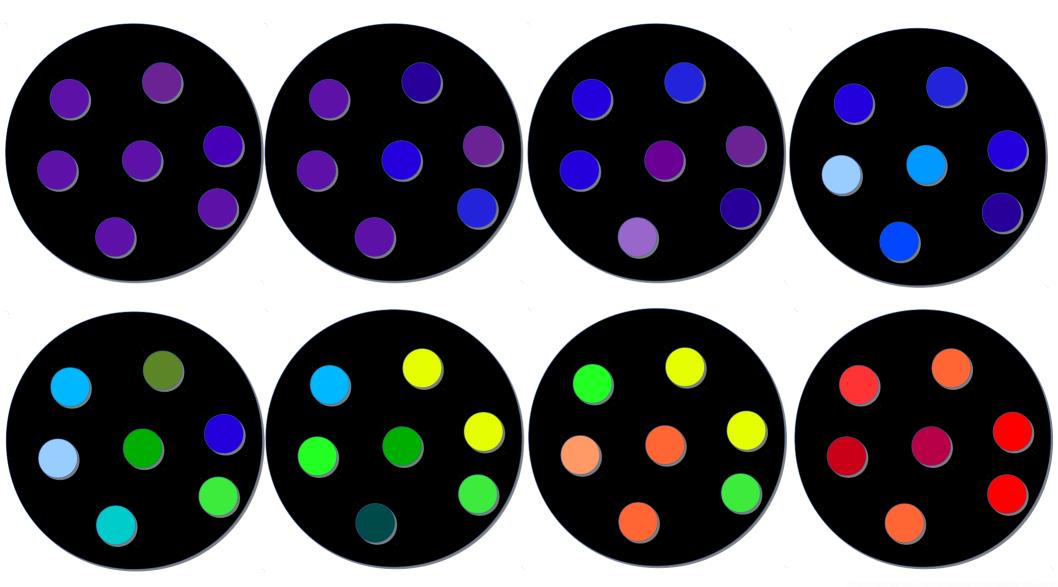
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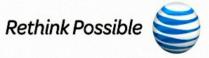


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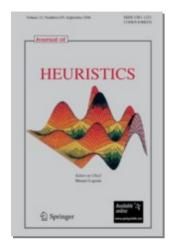




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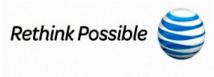
Reference



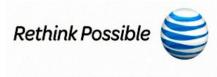
J.F. Gonçalves and M.G.C.R., "Biased random-key genetic algorithms for combinatorial optimization," J. of Heuristics, vol.17, pp. 487-525, 2011.

Tech report version:

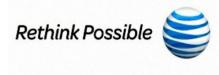
http://www.research.att.com/~mgcr/doc/srkga.pdf



Genetic algorithms and random keys



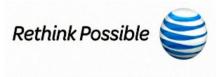
• Introduced by Bean (1994) for sequencing problems.



- Introduced by Bean (1994) for sequencing problems.
- Individuals are strings of real-valued numbers (random keys) in the interval [0,1).

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

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



- Introduced by Bean (1994) for sequencing problems.
- Individuals are strings of real-valued numbers (random keys) in the interval [0,1).
- Sorting random keys results in a sequencing order.

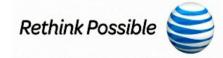
$$S = (0.25, 0.19, 0.67, 0.05, 0.89)$$

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

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

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

Sequence: 4 - 2 - 1 - 3 - 5



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

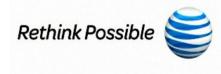
$$a = (0.25, 0.19, 0.67, 0.05, 0.89)$$

 $b = (0.63, 0.90, 0.76, 0.93, 0.08)$

- 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.

```
a = (0.25, 0.19, 0.67, 0.05, 0.89)

b = (0.63, 0.90, 0.76, 0.93, 0.08)
```



- Mating is done using parametrized uniform crossover (Spears & DeJong, 1990)
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```
a = (0.25, 0.19, 0.67, 0.05, 0.89)

b = (0.63, 0.90, 0.76, 0.93, 0.08)

c = (
```

- 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.

```
a = (0.25, 0.19, 0.67, 0.05, 0.89)

b = (0.63, 0.90, 0.76, 0.93, 0.08)

c = (0.25)
```

- 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.

```
a = (0.25, 0.19, 0.67, 0.05, 0.89)

b = (0.63, 0.90, 0.76, 0.93, 0.08)

c = (0.25, 0.90)
```

- 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.

```
a = (0.25, 0.19, 0.67, 0.05, 0.89)

b = (0.63, 0.90, 0.76, 0.93, 0.08)

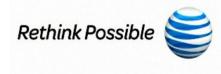
c = (0.25, 0.90, 0.76)
```

- 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.

```
a = (0.25, 0.19, 0.67, 0.05, 0.89)

b = (0.63, 0.90, 0.76, 0.93, 0.08)

c = (0.25, 0.90, 0.76, 0.05)
```



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 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.

```
a = (0.25, 0.19, 0.67, 0.05, 0.89)

b = (0.63, 0.90, 0.76, 0.93, 0.08)

c = (0.25, 0.90, 0.76, 0.05, 0.89)
```

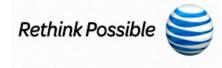
- Mating is done using parametrized uniform
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a = (0.25, 0.19, 0.67, 0.05, 0.89)

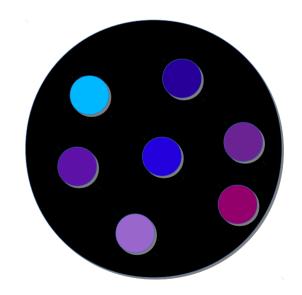
b = (0.63, 0.90, 0.76, 0.93, 0.08)

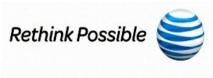
c = (0.25, 0.90, 0.76, 0.05, 0.89)
```

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

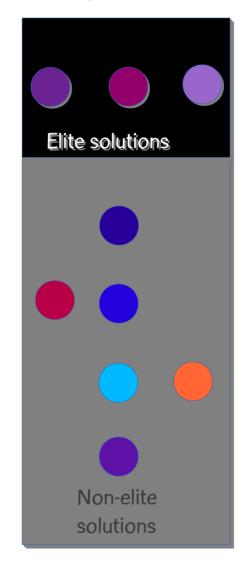


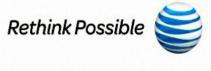
Initial population is made up of P random-key vectors, each with N keys, each having a value generated uniformly at random in the interval [0,1).



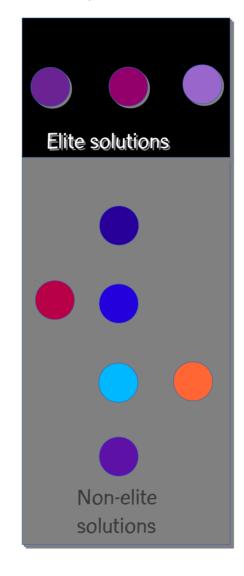


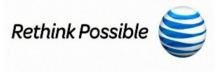
At the K-th generation, compute the cost of each solution ...



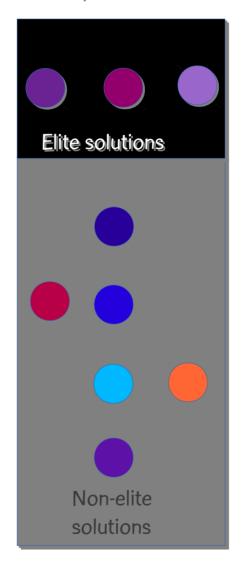


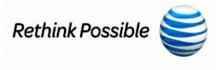
At the K-th generation, compute the cost of each solution and partition the solutions into two sets:



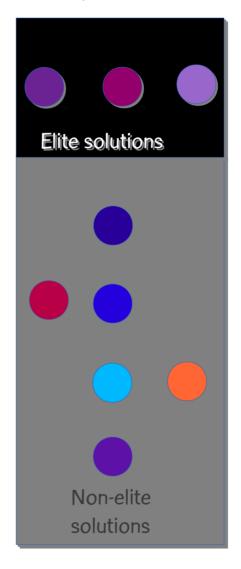


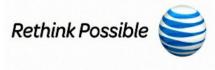
At the K-th generation, compute the cost of each solution and partition the solutions into two sets: elite solutions and non-elite solutions.





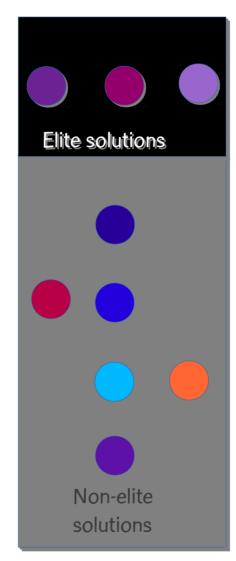
At the K-th generation, compute the cost of each solution and partition the solutions into two sets: elite solutions and non-elite solutions. Elite set should be smaller of the two sets and contain best solutions.





Evolutionary dynamics





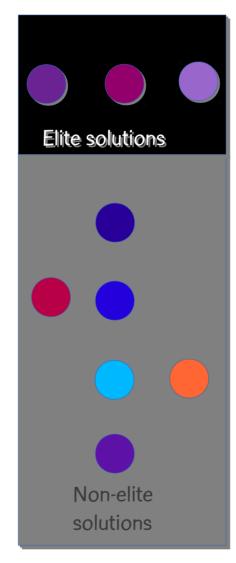




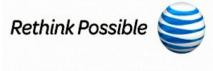
Evolutionary dynamics

 Copy elite solutions from population K to population K+1



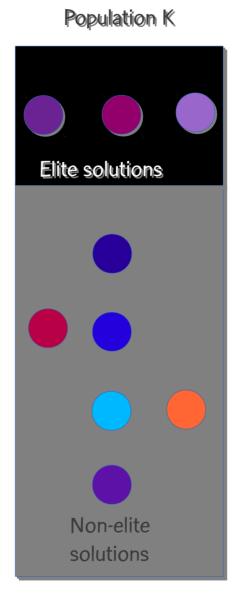






Evolutionary dynamics

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

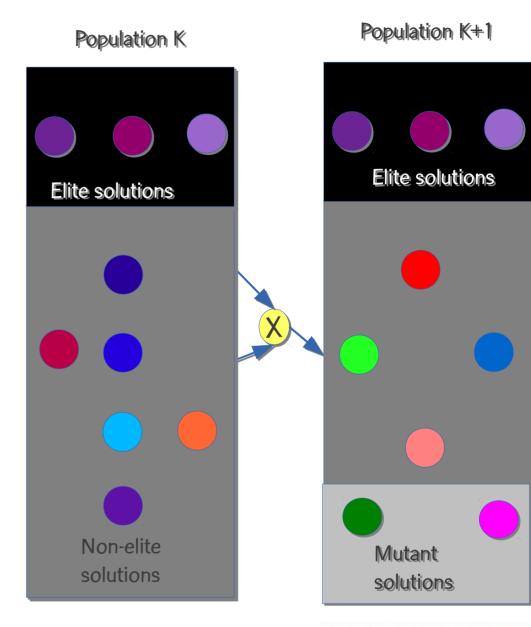


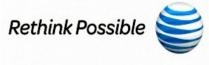




Evolutionary dynamics

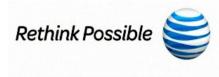
- Copy elite solutions from population K to population K+1
- Add R random solutions (mutants) to population K+1
- While K+1-th population < P
 - RANDOM-KEY GA: Use any two solutions in population K to produce child in population K+1. Mates are chosen at random.





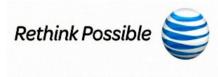
Biased random key genetic algorithm

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



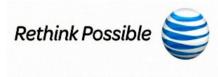
Biased random key genetic algorithm

- 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.



RKGA BRKGA

both parents chosen at random from entire population

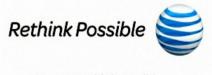


RKGA

both parents chosen at random from entire population

BRKGA

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



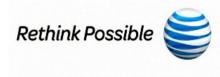
RKGA

both parents chosen at random from entire population

BRKGA

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

either parent can be parent A in parametrized uniform crossover



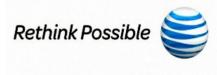
RKGA

both parents chosen at random from entire population

BRKGA

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

either parent can be parent A in parametrized uniform crossover best fit parent is parent A in parametrized uniform crossover



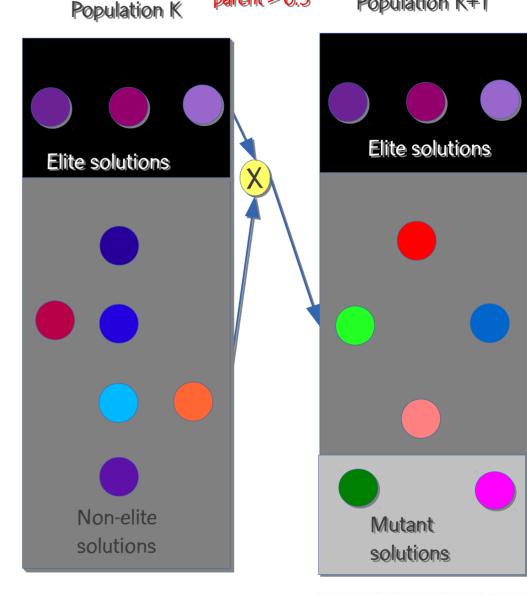
Biased random key GA

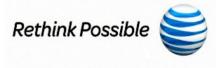
BRKGA: Probability child inherits key of elite parent ≥ 0.5

Population K+1

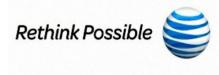
Evolutionary dynamics

- 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.

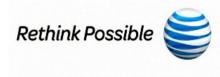




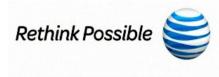
 Random method: keys are randomly generated so solutions are always vectors of random keys



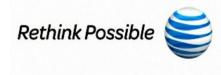
- Random method: keys are randomly generated so solutions are always vectors of random keys
- Elitist strategy: best solutions are passed without change from one generation to the next (incumbent is kept)



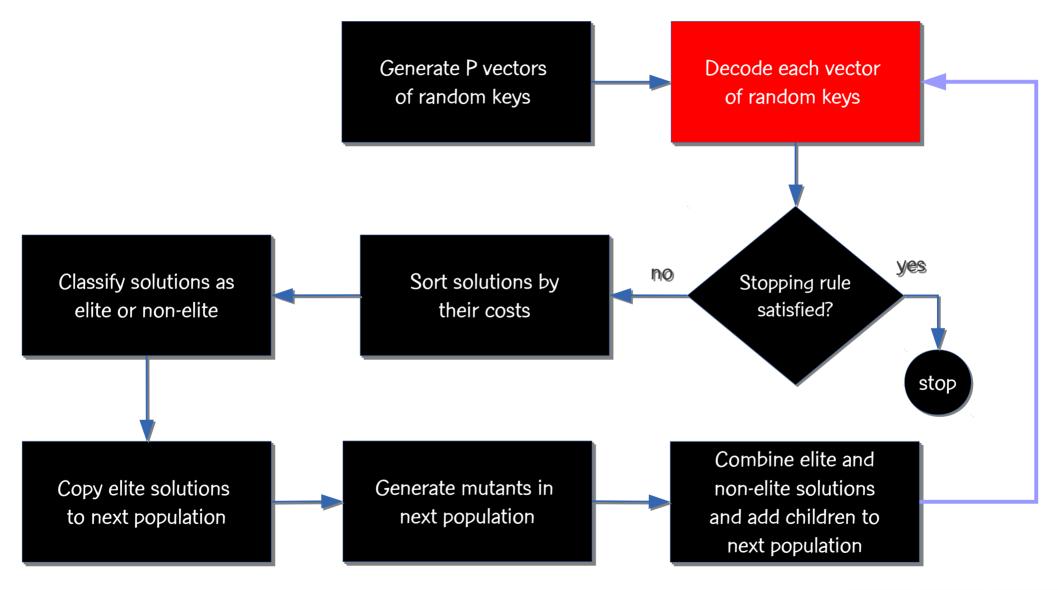
- Random method: keys are randomly generated so solutions are always vectors of random keys
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- Child inherits more characteristics of elite parent: one parent is always selected (with replacement) from the small elite set and probability that child inherits key of elite parent > 0.5 Not so in the RKGA of Bean.

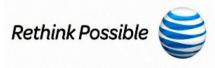


- Random method: keys are randomly generated so solutions are always vectors of random keys
- Elitist strategy: best solutions are passed without change from one generation to the next (incumbent is kept)
- Child inherits more characteristics of elite parent: one parent is always selected (with replacement) 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)

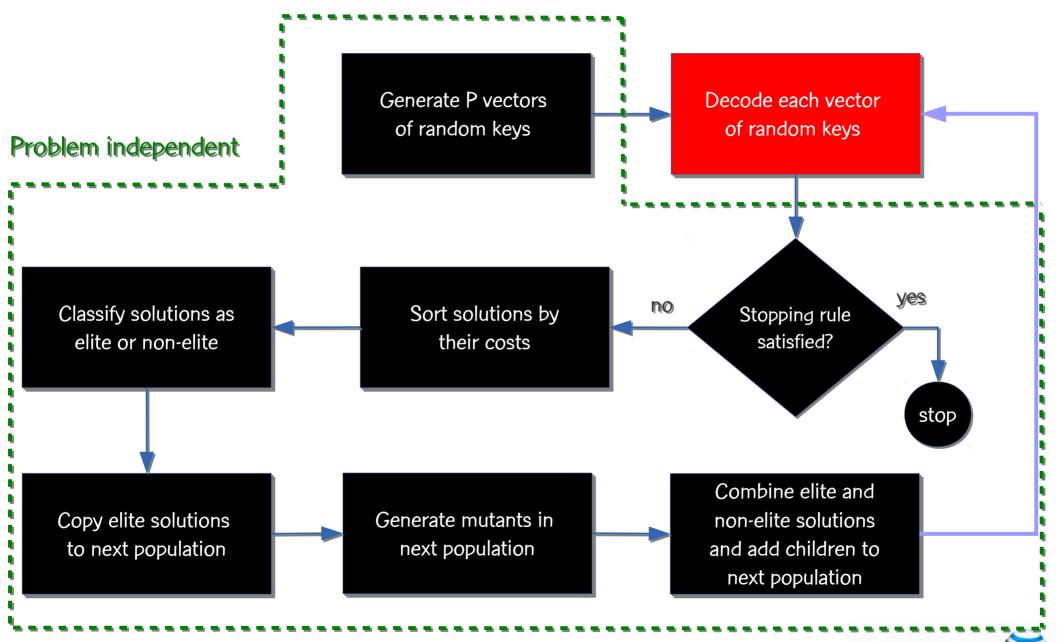


Framework for biased random-key genetic algorithms

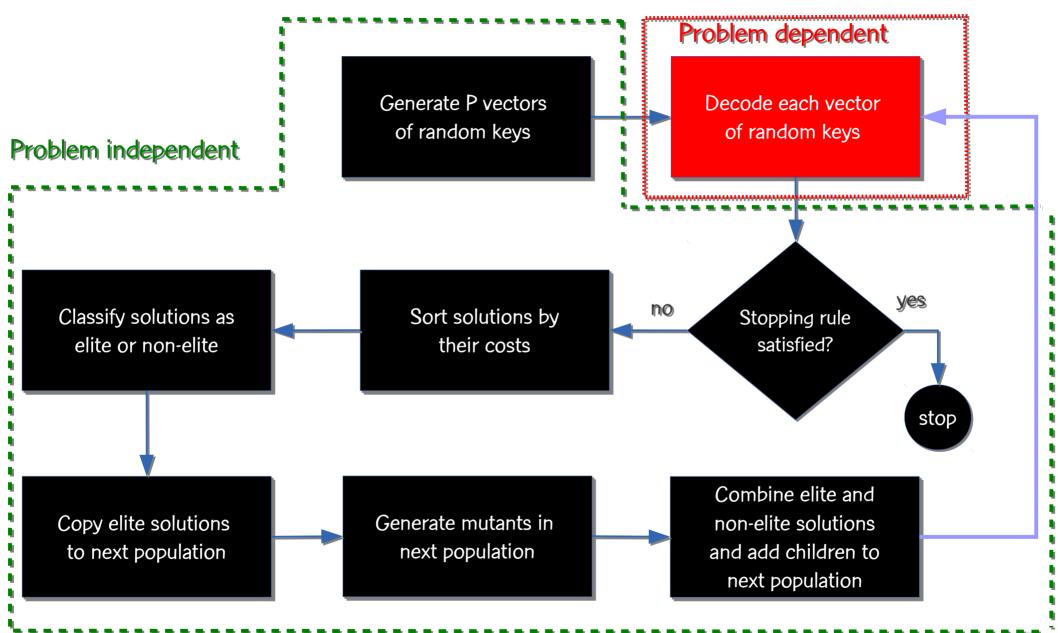




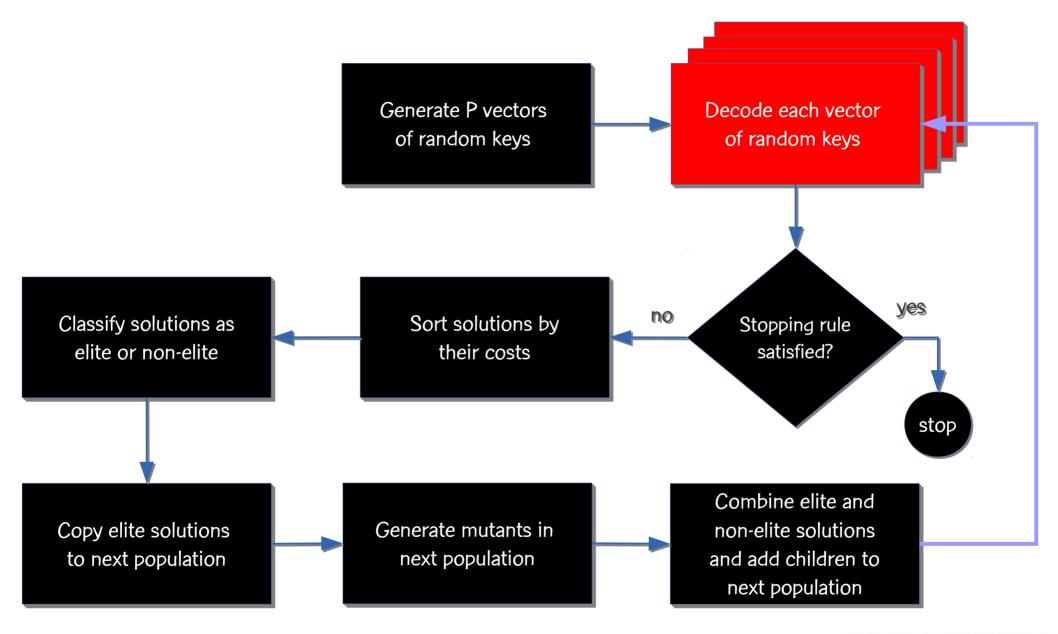
Framework for biased random-key genetic algorithms

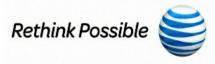


Framework for biased random-key genetic algorithms

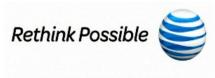


Decoding of random key vectors can be done in parallel

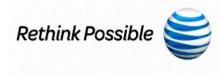




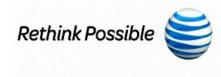
Specifying a BRKGA



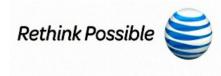
Encoding is always done the same way, i.e. with a vector of N random-keys (parameter N must be specified)



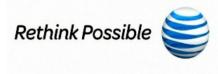
- Encoding is always done the same way, i.e. with a vector of N random-keys (parameter N must be specified)
- Decoder that takes as input a vector of N random-keys and outputs the corresponding solution of the combinatorial optimization problem and its cost (this is usually a heuristic)



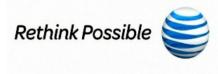
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- Parameters



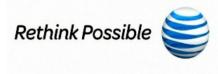
- Size of population
- Size of elite partition
- Size of mutant set
- Child inheritance probability
- Restart strategy parameter
- Stopping criterion



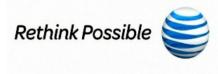
- Size of population: a function of N, say N or 2N
- Size of elite partition
- Size of mutant set
- Child inheritance probability
- Restart strategy parameter
- Stopping criterion



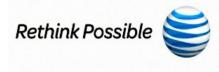
- Size of population: a function of N, say N or 2N
- Size of elite partition: 15-25% of population
- Size of mutant set
- Child inheritance probability
- Restart strategy parameter
- Stopping criterion



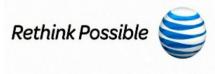
- Size of population: a function of N, say N or 2N
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- Size of mutant set: 5-15% of population
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- Restart strategy parameter
- Stopping criterion



- Size of population: a function of N, say N or 2N
- Size of elite partition: 15-25% of population
- Size of mutant set: 5-15% of population
- Child inheritance probability: > 0.5, say 0.7
- Restart strategy parameter
- Stopping criterion



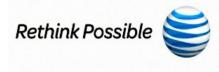
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- Stopping criterion



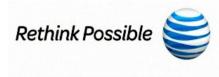
Parameters:

- Size of population: a function of N, say N or 2N
- Size of elite partition: 15-25% of population
- Size of mutant set: 5-15% of population
- Child inheritance probability: > 0.5, say 0.7
- Restart strategy parameter: a function of N, say 2N or 10N
- Stopping criterion: e.g. time, # generations, solution quality, generations without improvement

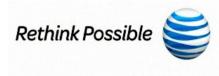
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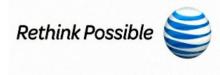
 Efficient and easy-to-use object oriented application programming interface (API) for the algorithmic framework of BRKGA.



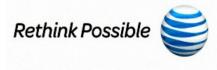
- Efficient and easy-to-use object oriented application programming interface (API) for the algorithmic framework of BRKGA.
- Cross-platform library handles large portion of problem independent modules that make up the framework, e.g.
 - population management
 - evolutionary dynamics



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- Implemented in C++ and may benefit from shared-memory parallelism if available.



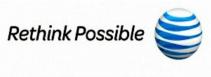
- Efficient and easy-to-use object oriented application programming interface (API) for the algorithmic framework of BRKGA.
- Cross-platform library handles large portion of problem independent modules that make up the framework, e.g.
 - population management
 - evolutionary dynamics
- Implemented in C++ and may benefit from shared-memory parallelism if available.
- User only needs to implement problem-dependent decoder.





Paper: Rodrigo F. Toso and M.G.C.R., "A C++ Application Programming Interface for Biased Random-Key Genetic Algorithms," Optimization Methods & Software, published online 13 March 2014.

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

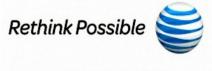


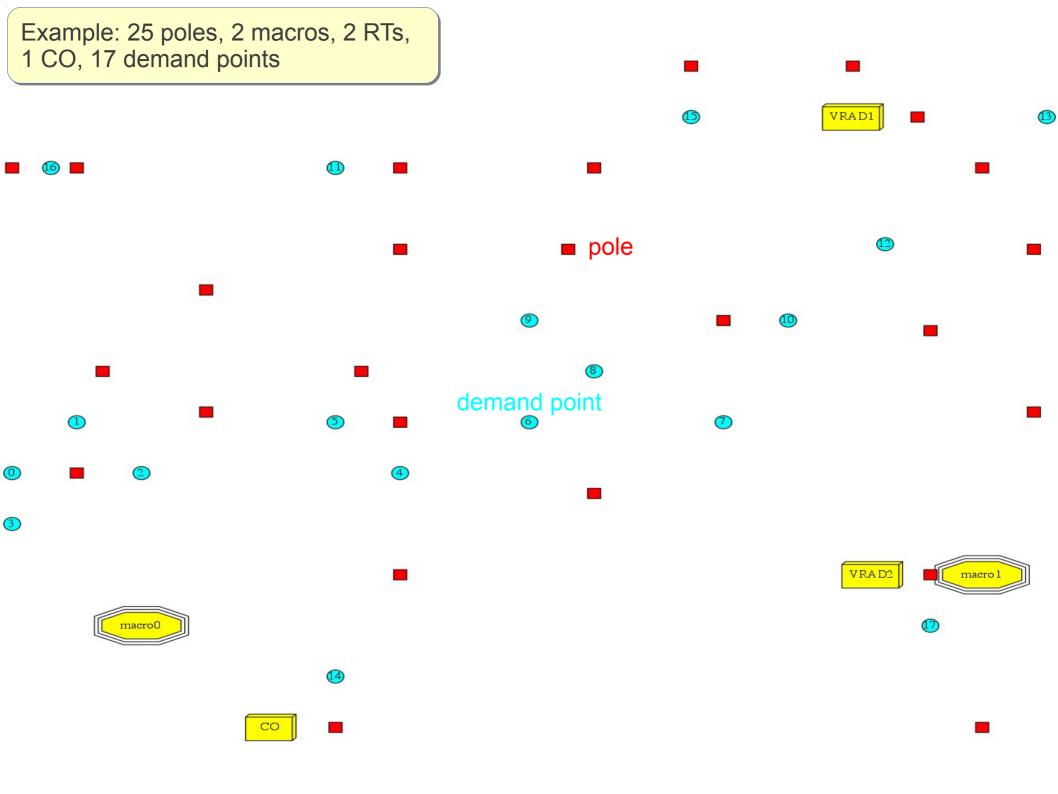
Decoder for wireless backhaul planning

- Biased Random-Key Genetic Algorithm
 - Learn the best network layout and equipment placement
- A solution is encoded by a vector x ∈ [0,1]ⁿ
 - Where $n = 5 \times \#$ of poles

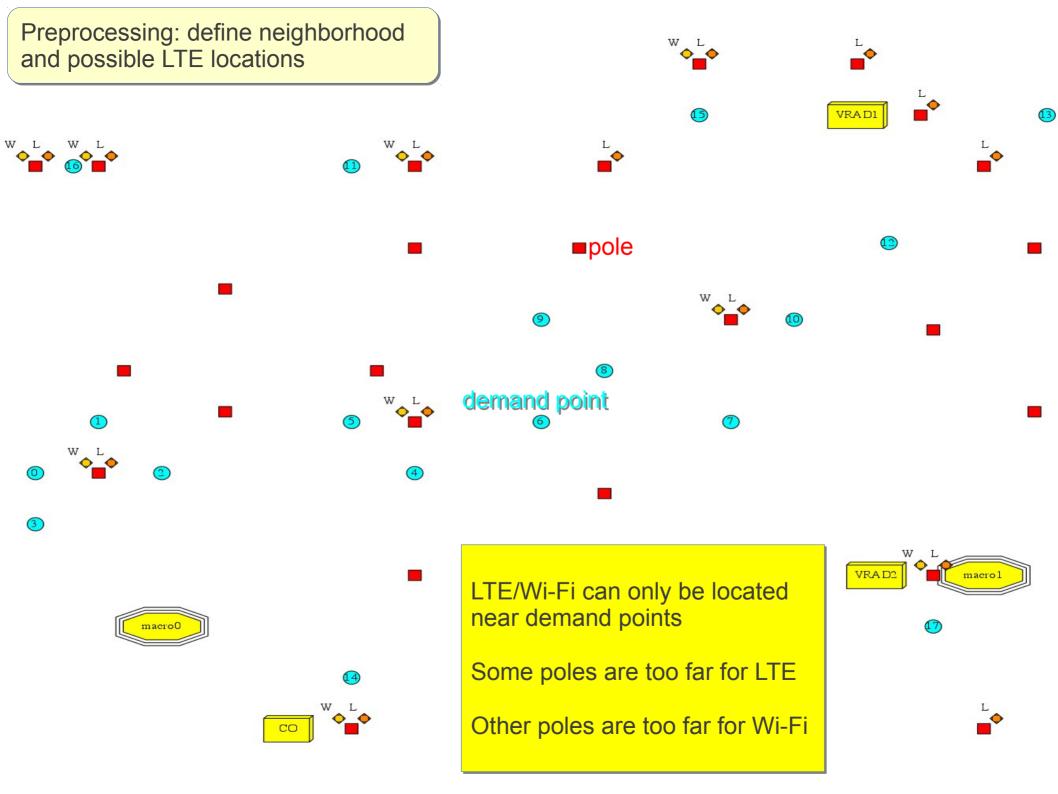
Decoder

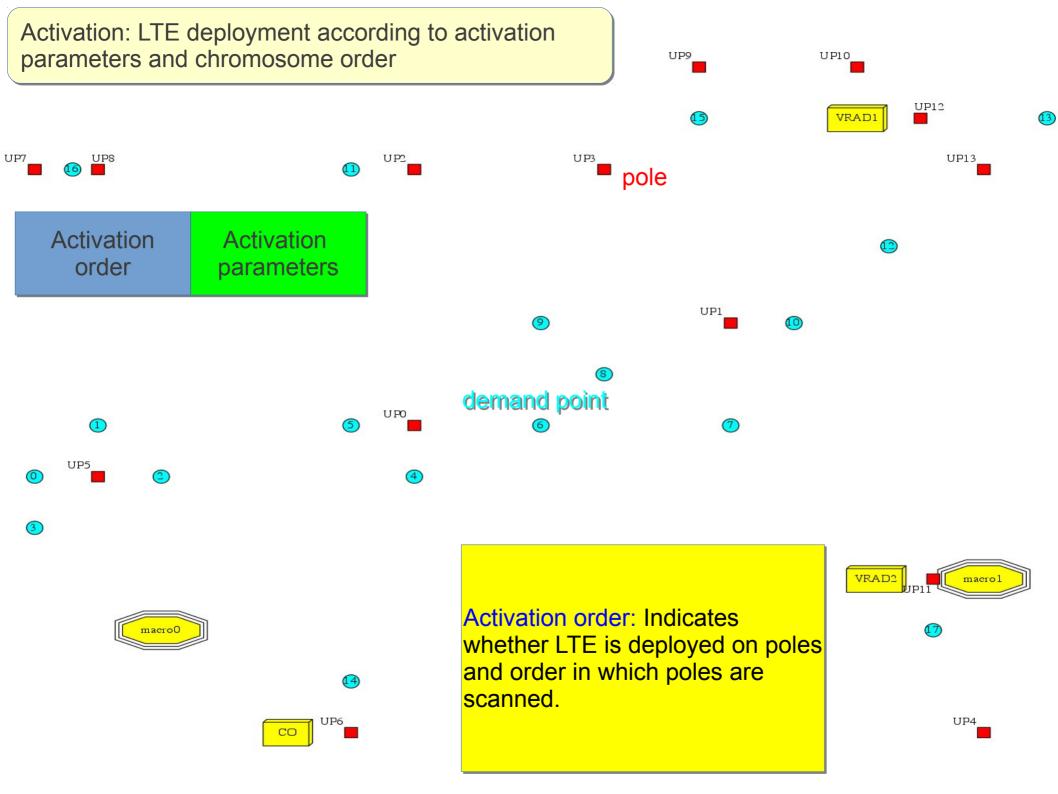
- Define activation order & install LTE on poles
- Build backhaul graph
- Remove unused equipment
- Compute maximum flow from demand points to FAPs
- Remove unused equipment & poles
- Compute cost and revenue and return objective function value

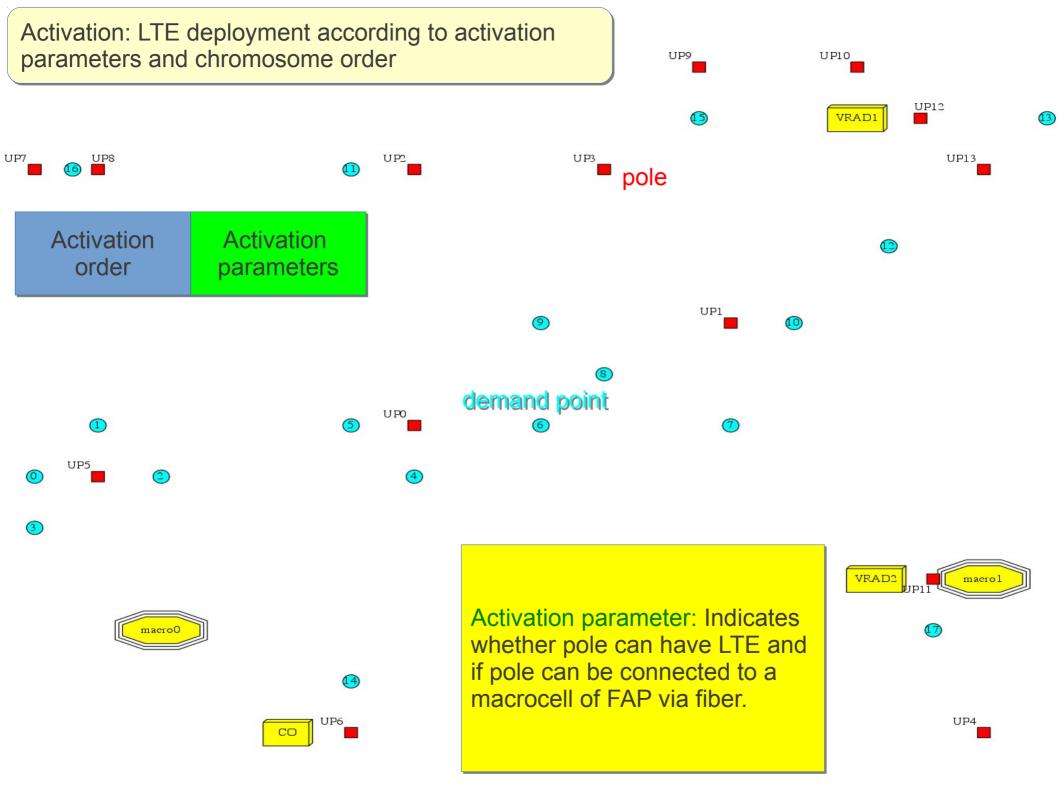


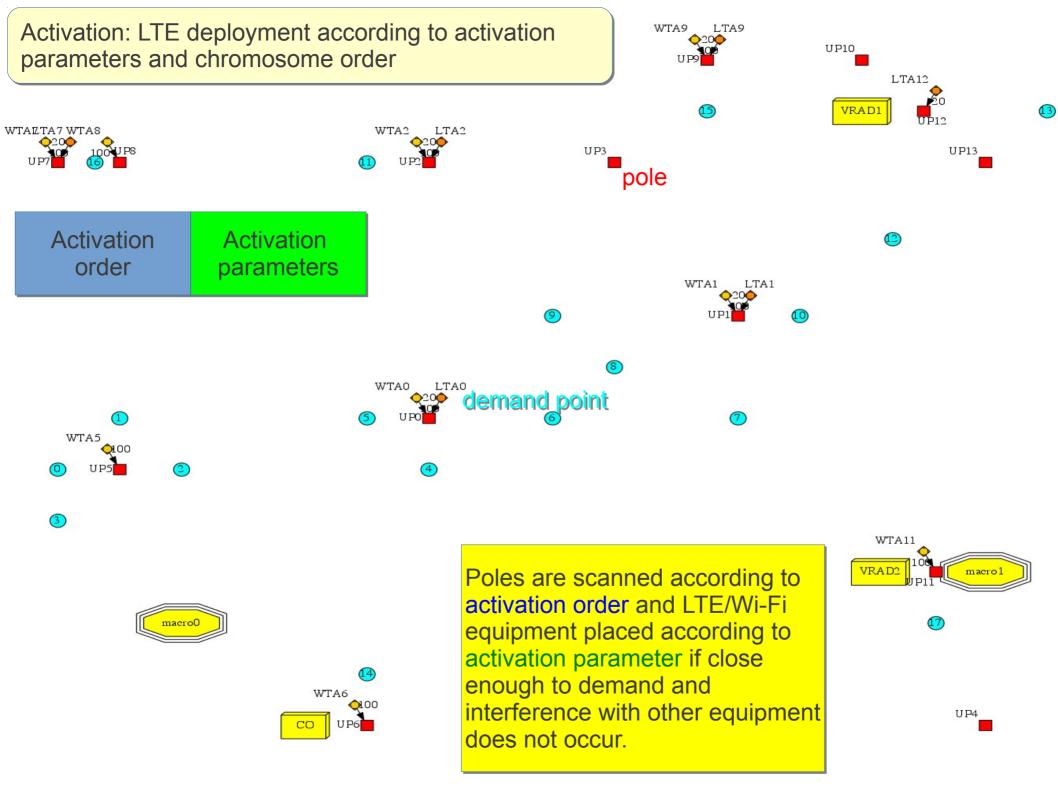


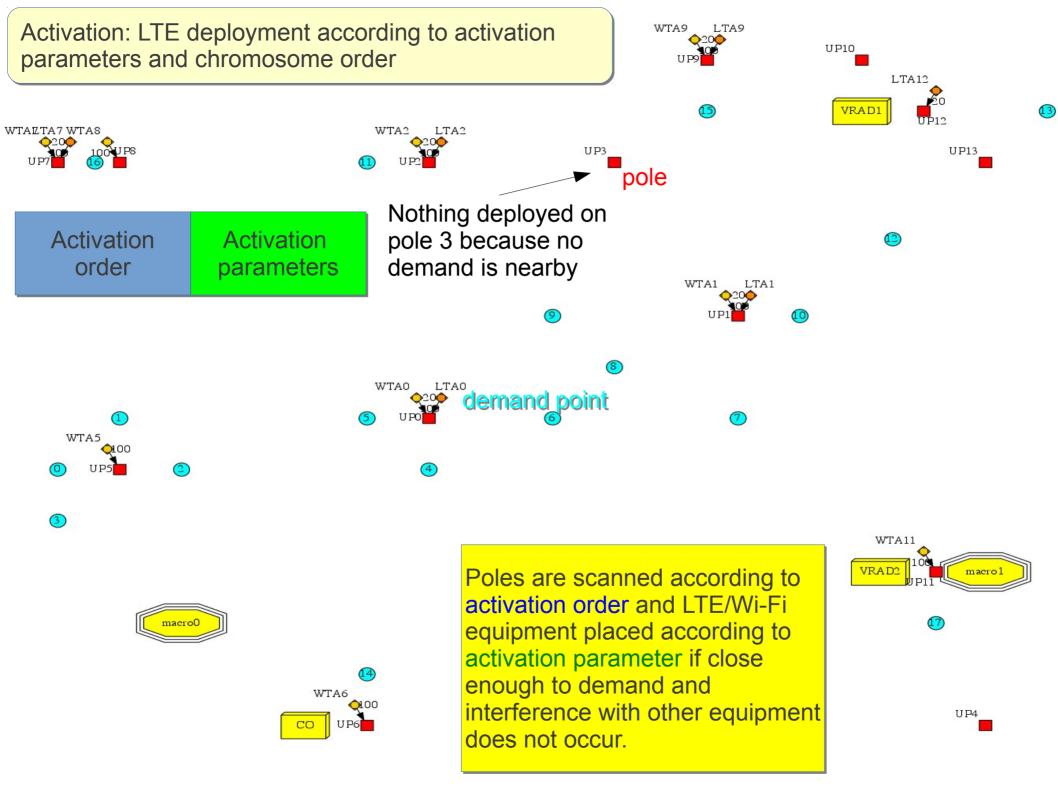
Preprocessing: define neighborhood and possible LTE locations VRA D1 pole demand point macro1 macro0

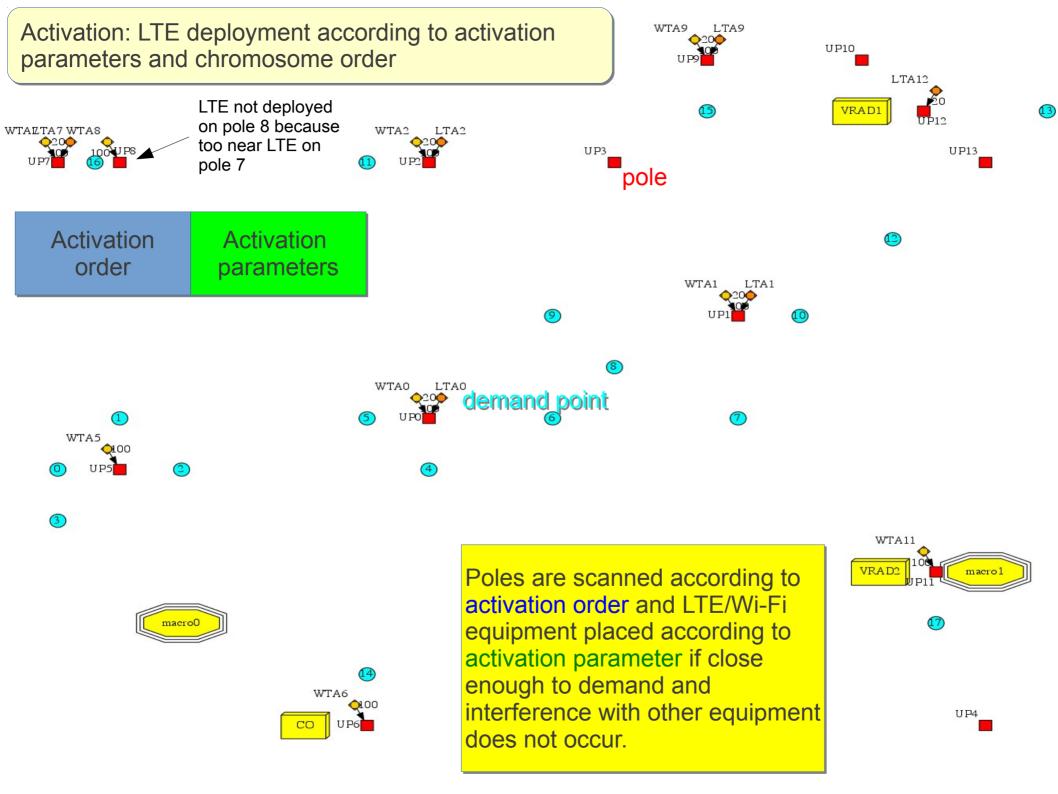


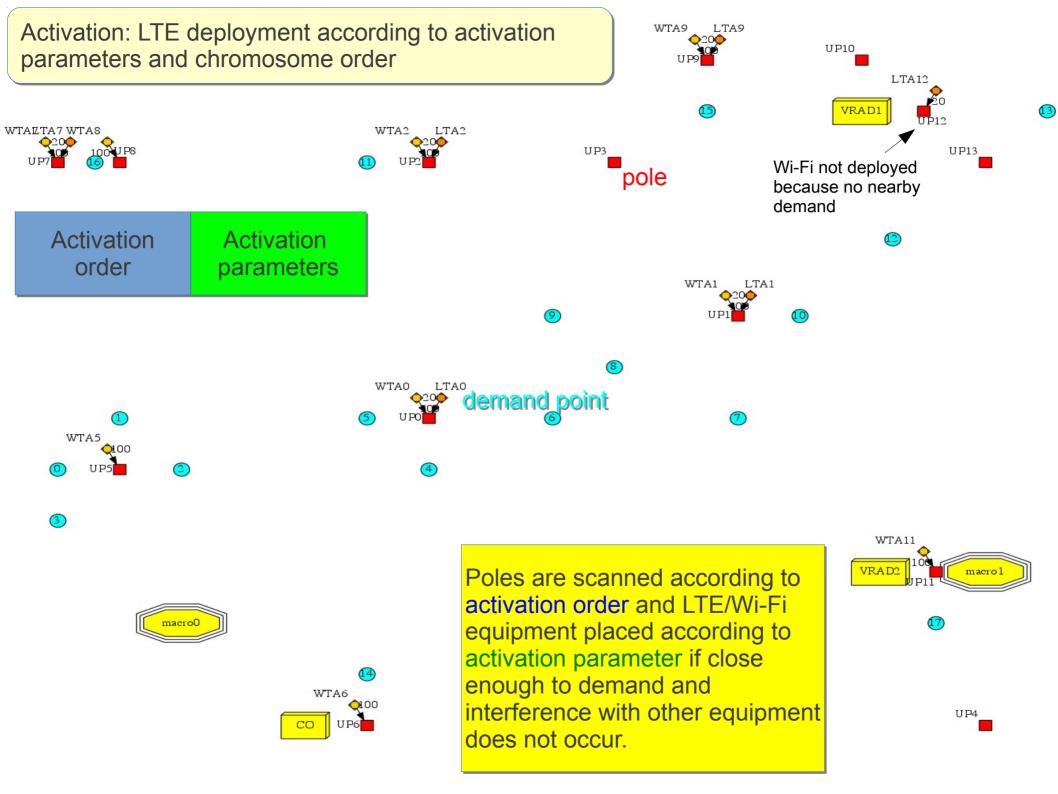


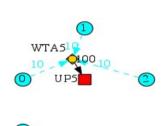




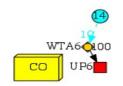


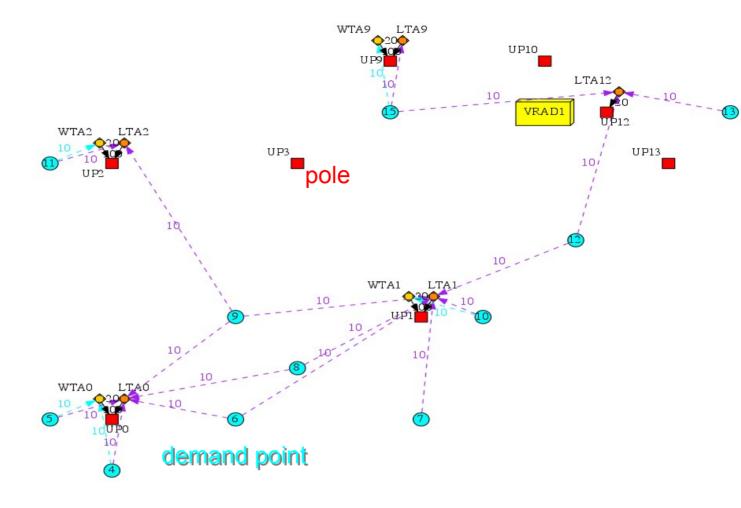






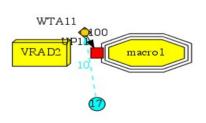




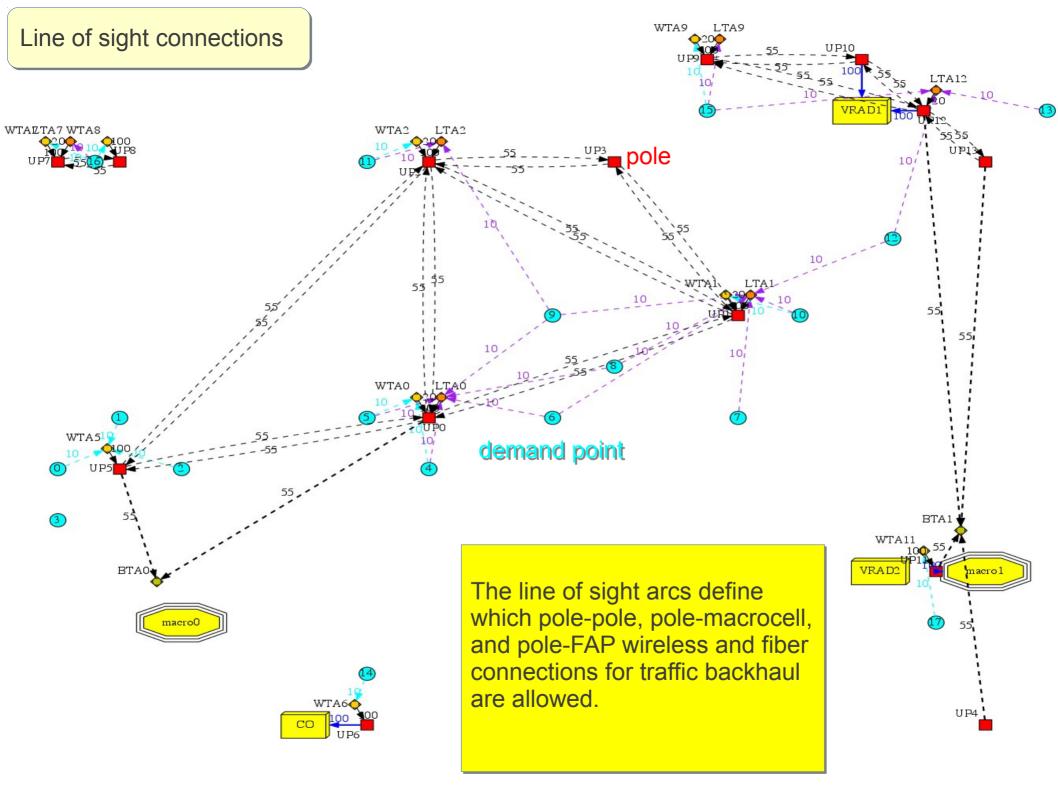


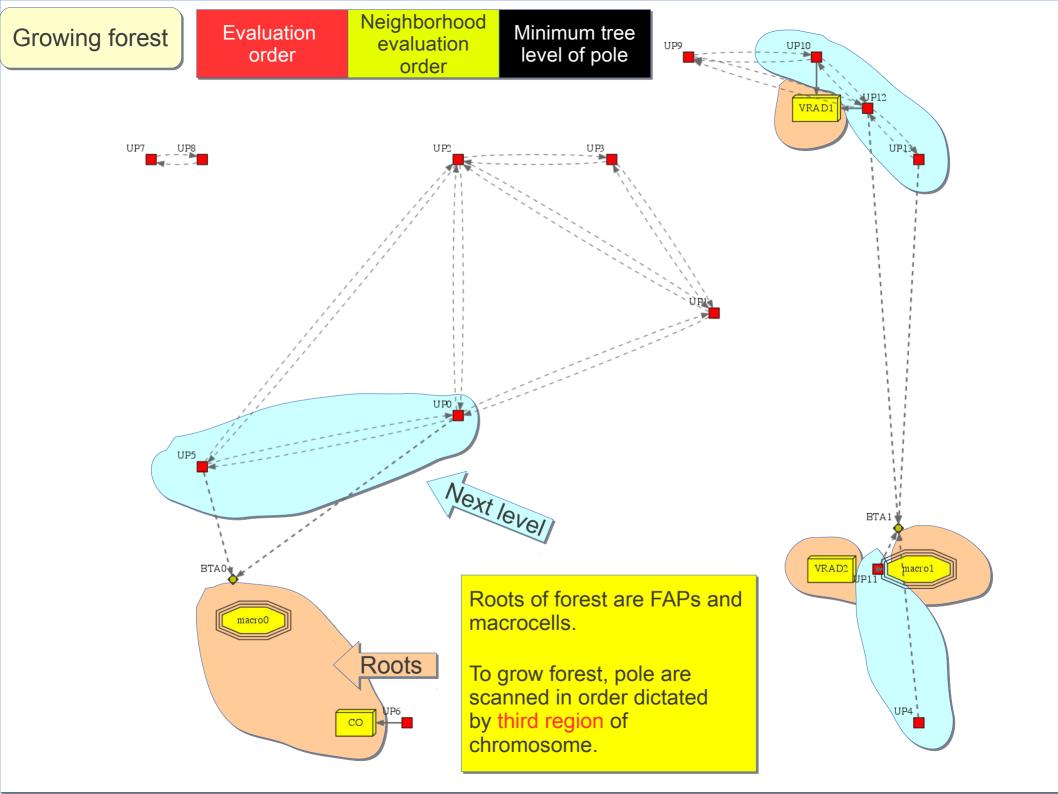
Demands are homed-in on poles that are within range.

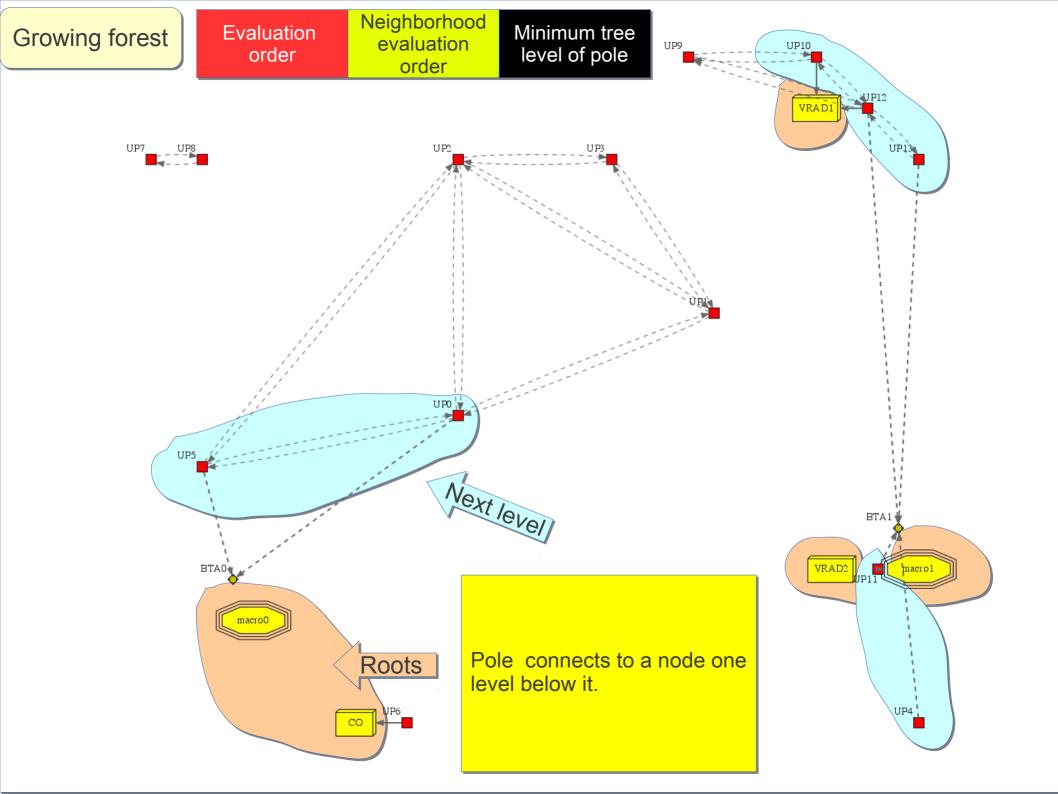
If demand is near more than one pole, its traffic is allowed to be split among these poles.

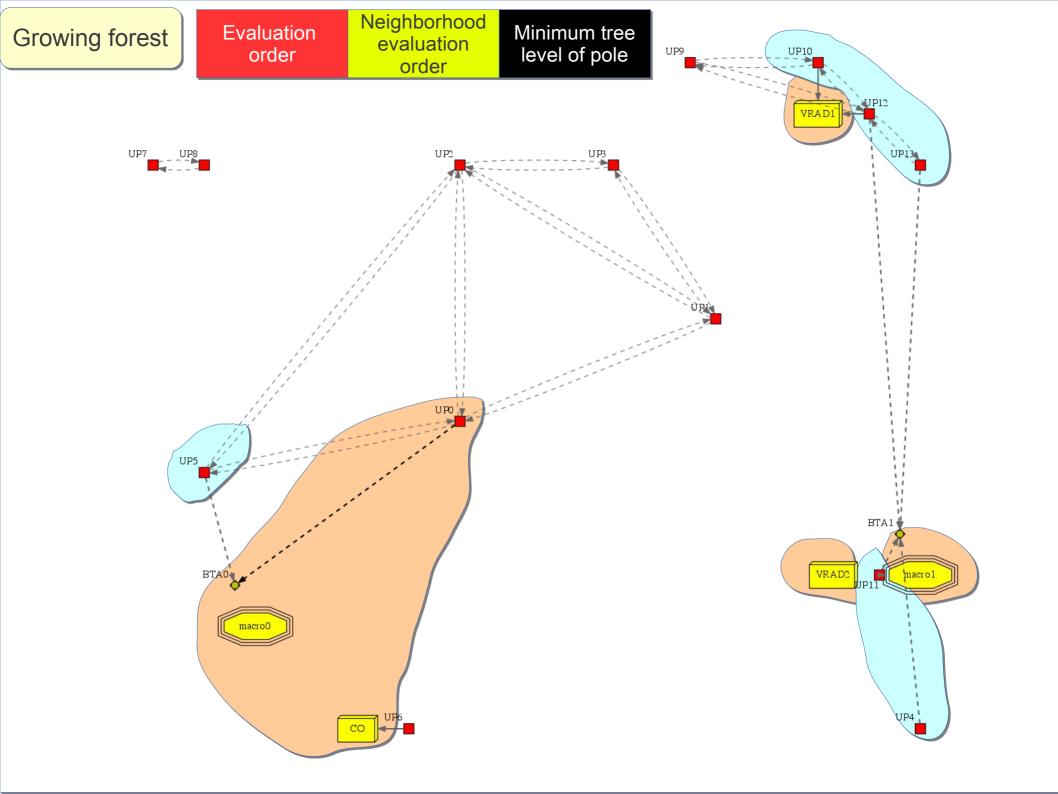


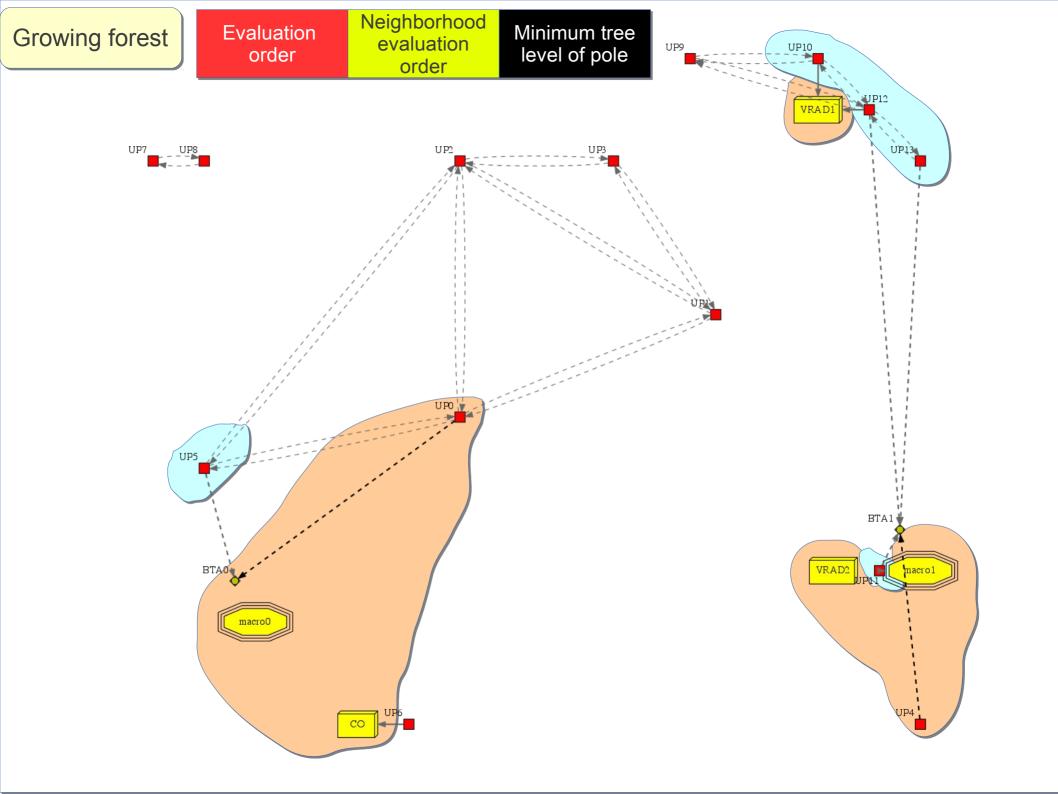


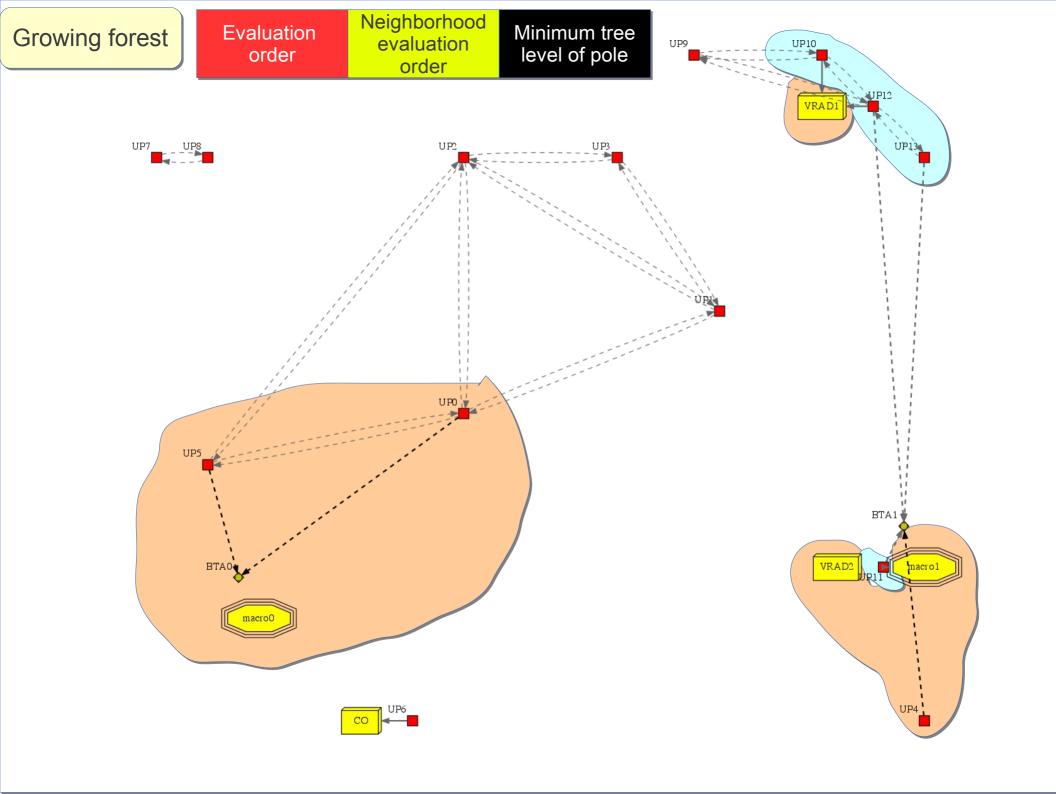


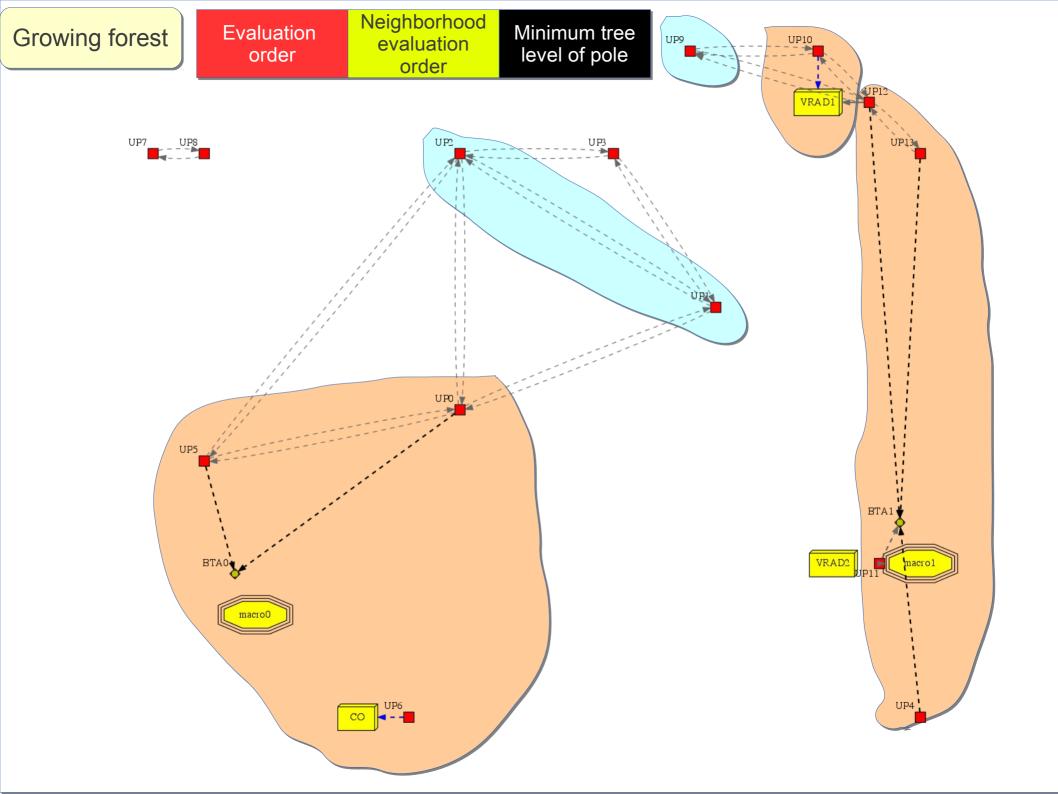


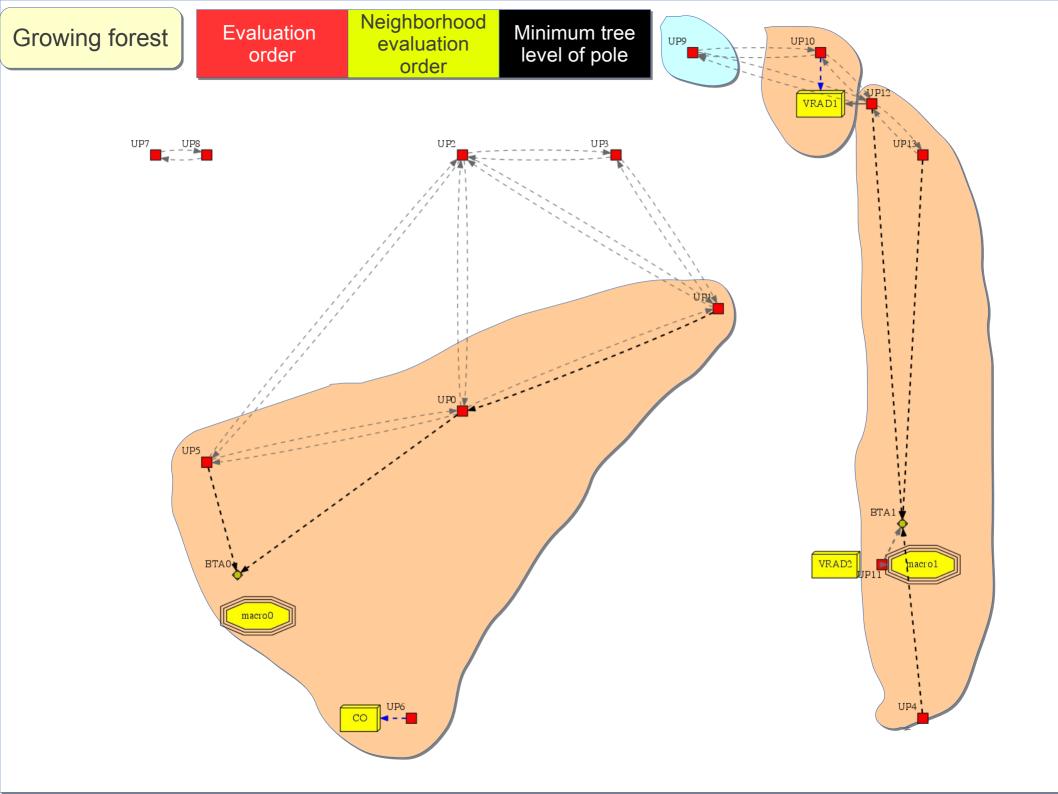


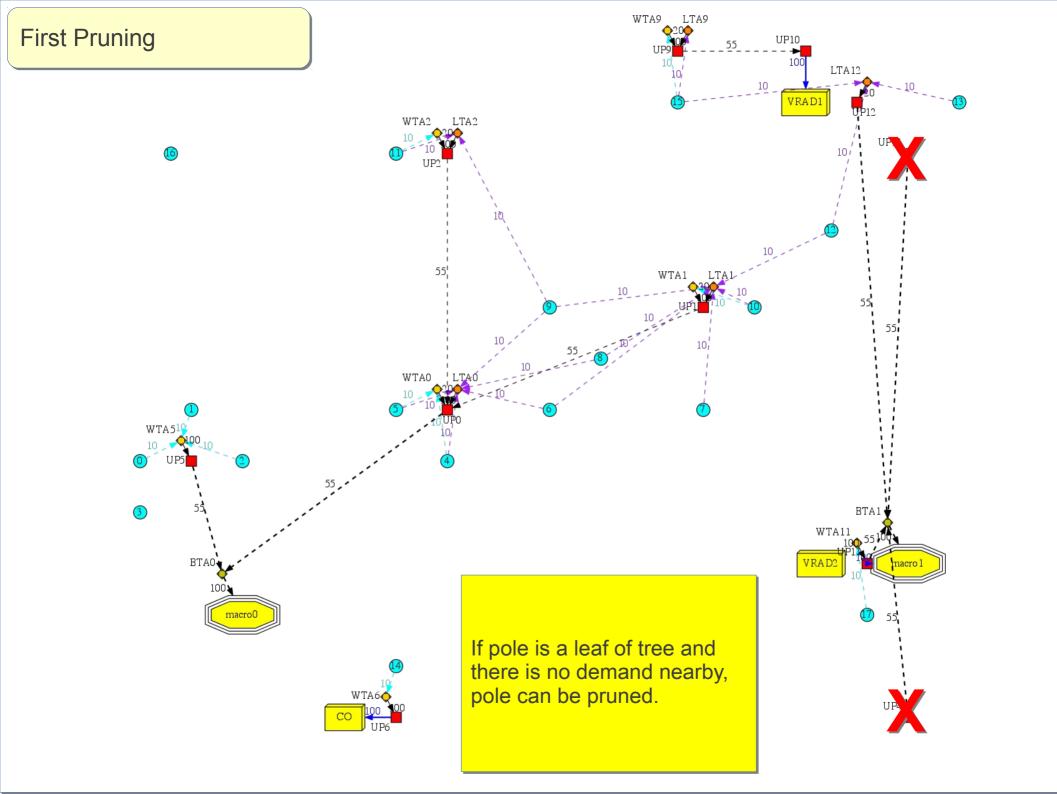


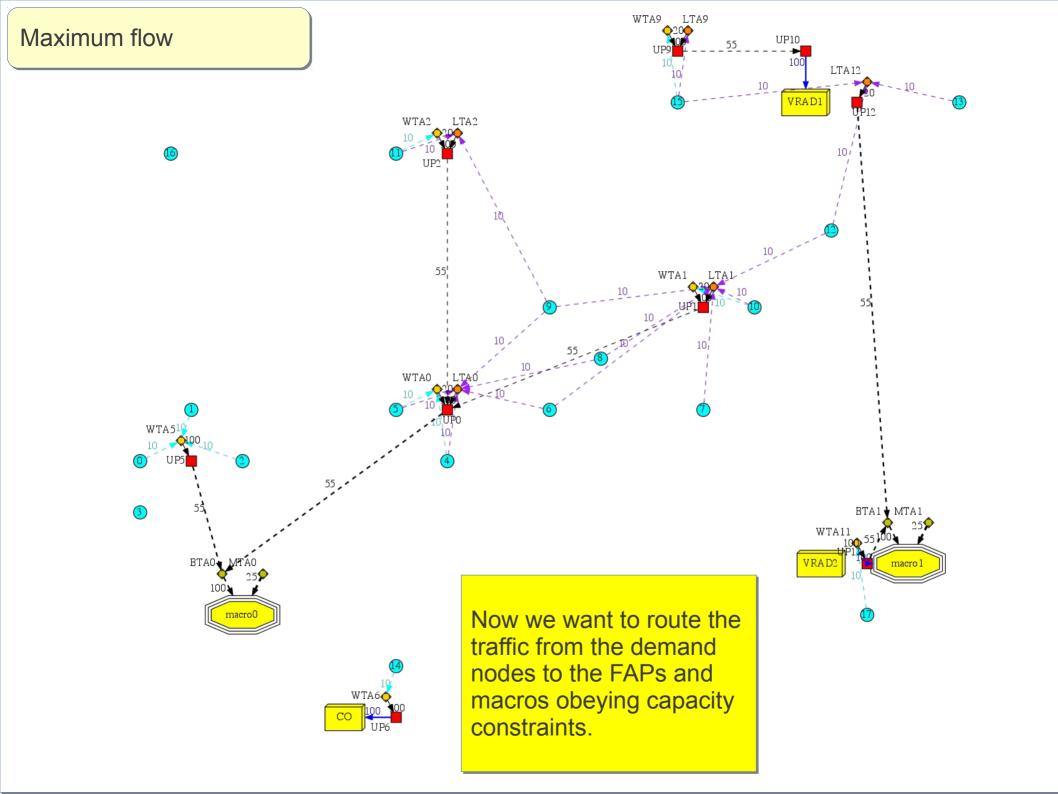


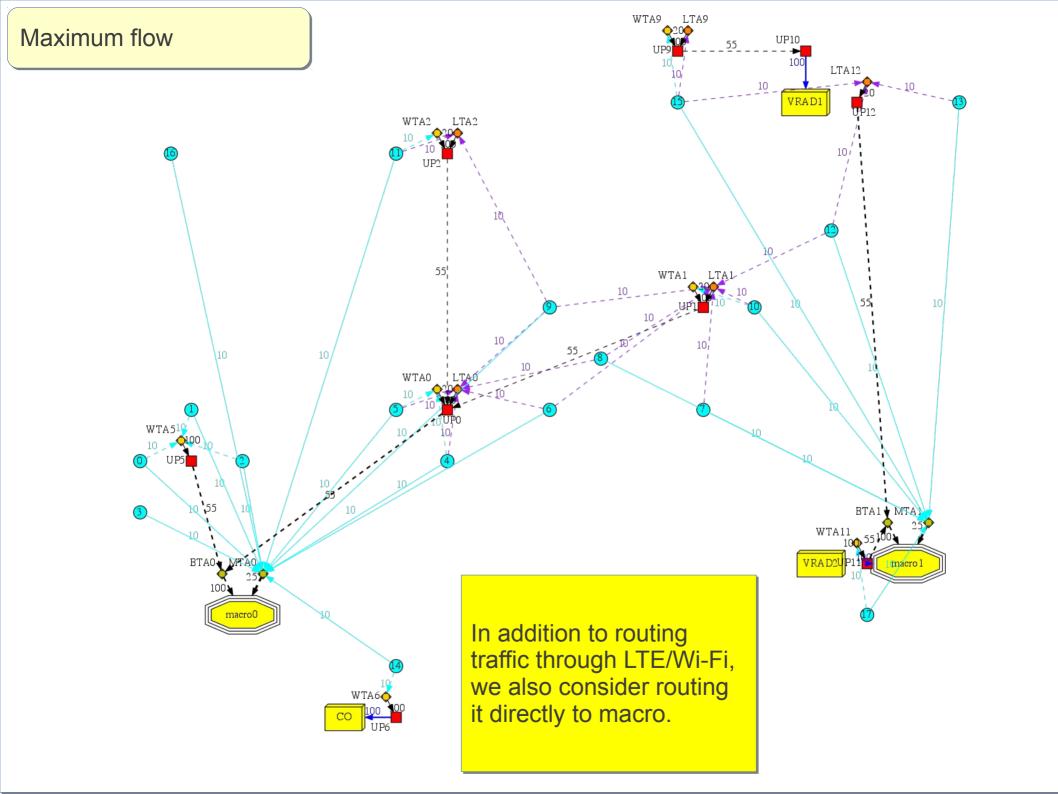


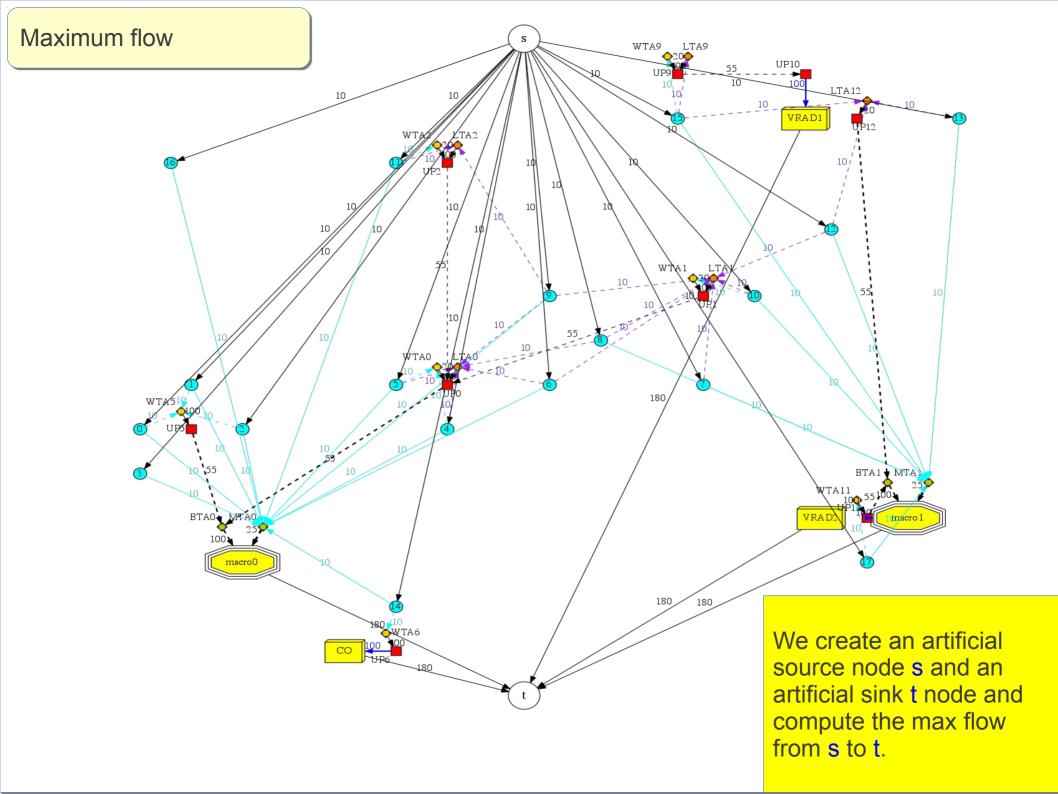


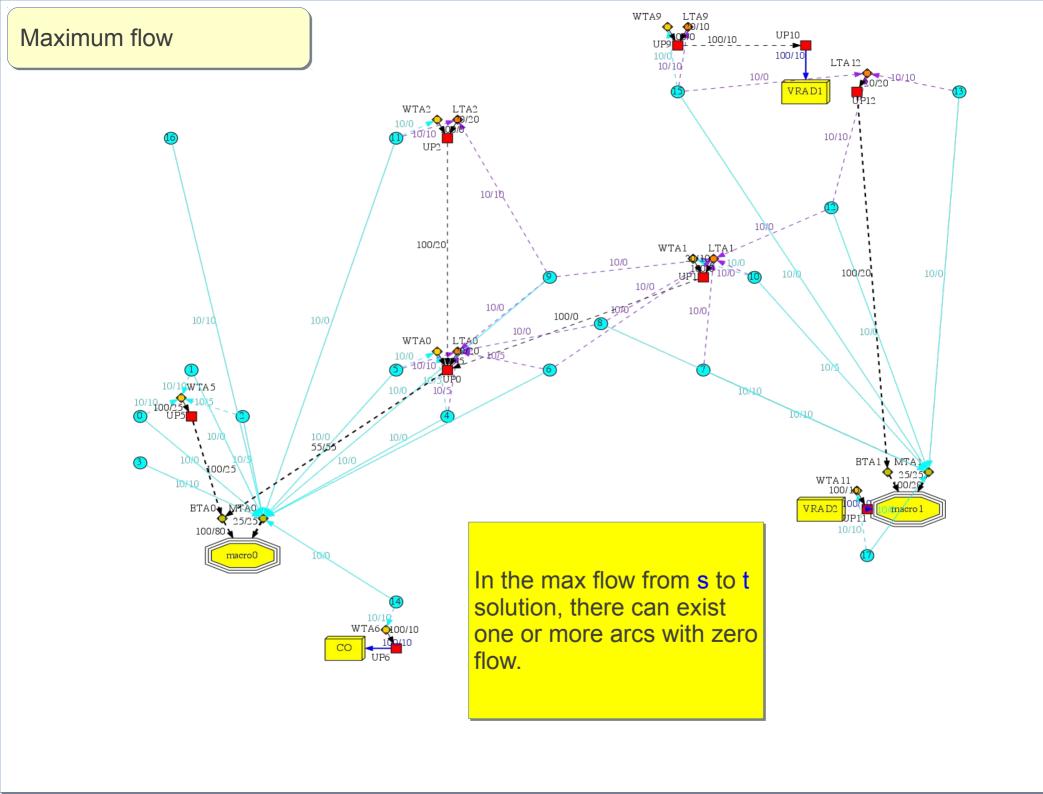


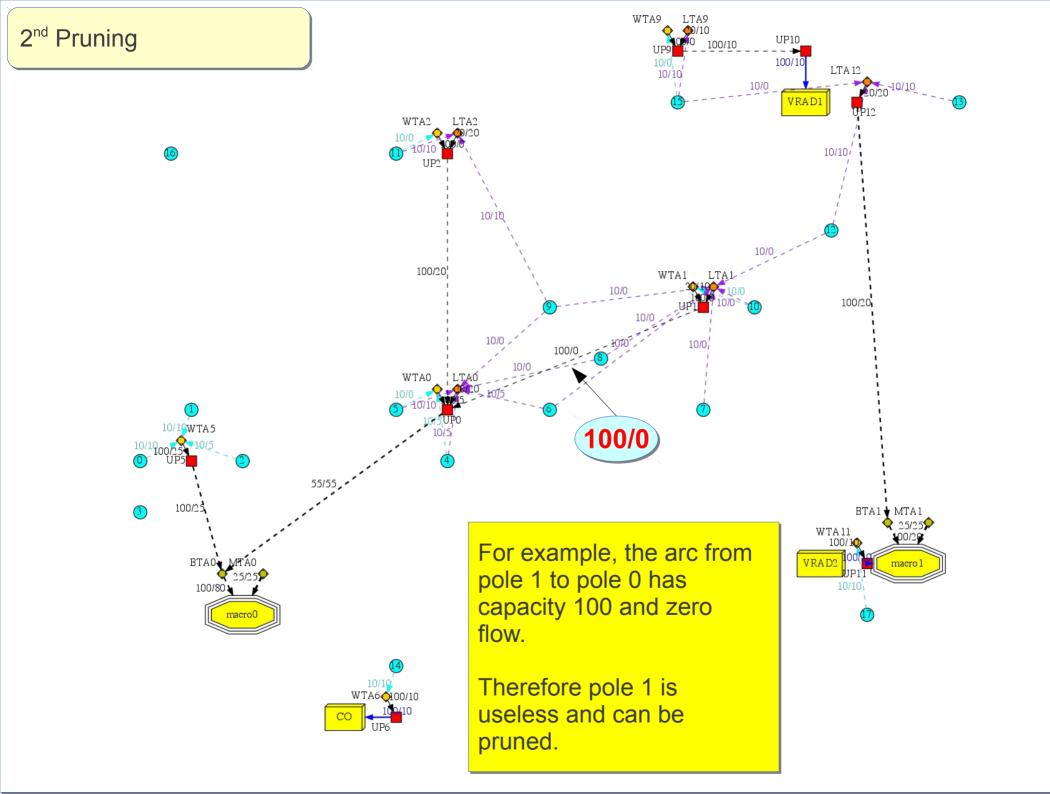


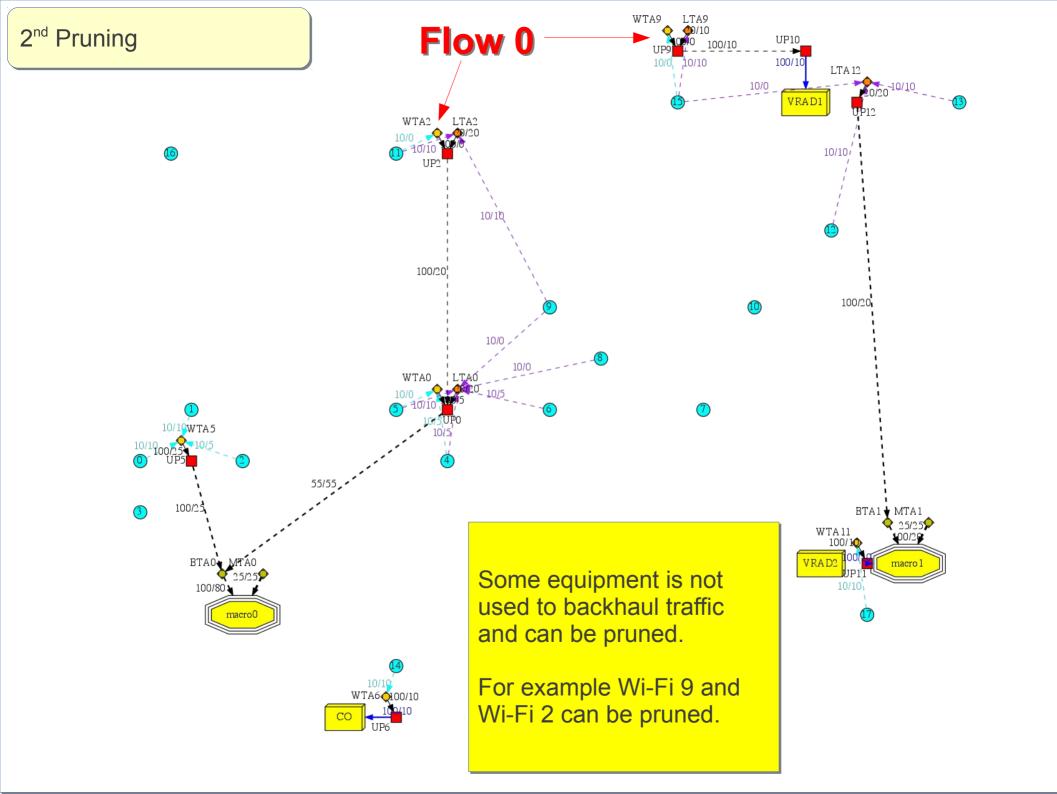




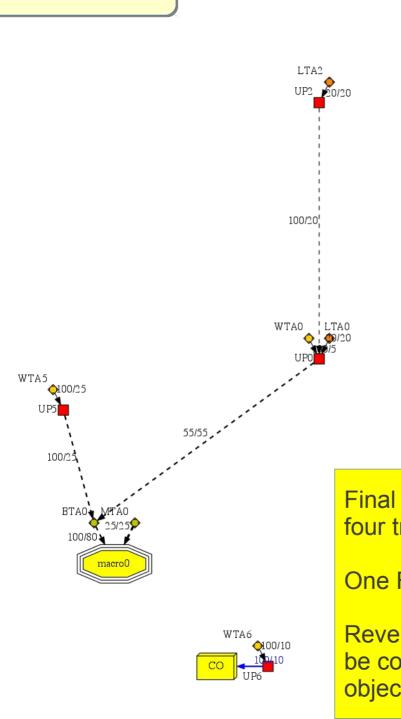








Final backhaul forest



100/10 LTA12 100/20 BTA1 **♦** MTA1 25/25 WTA11 100/10 100/20 macro 1

Final backhaul forest has four trees.

UP9 0/10 100/10

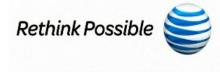
One FAP is not used.

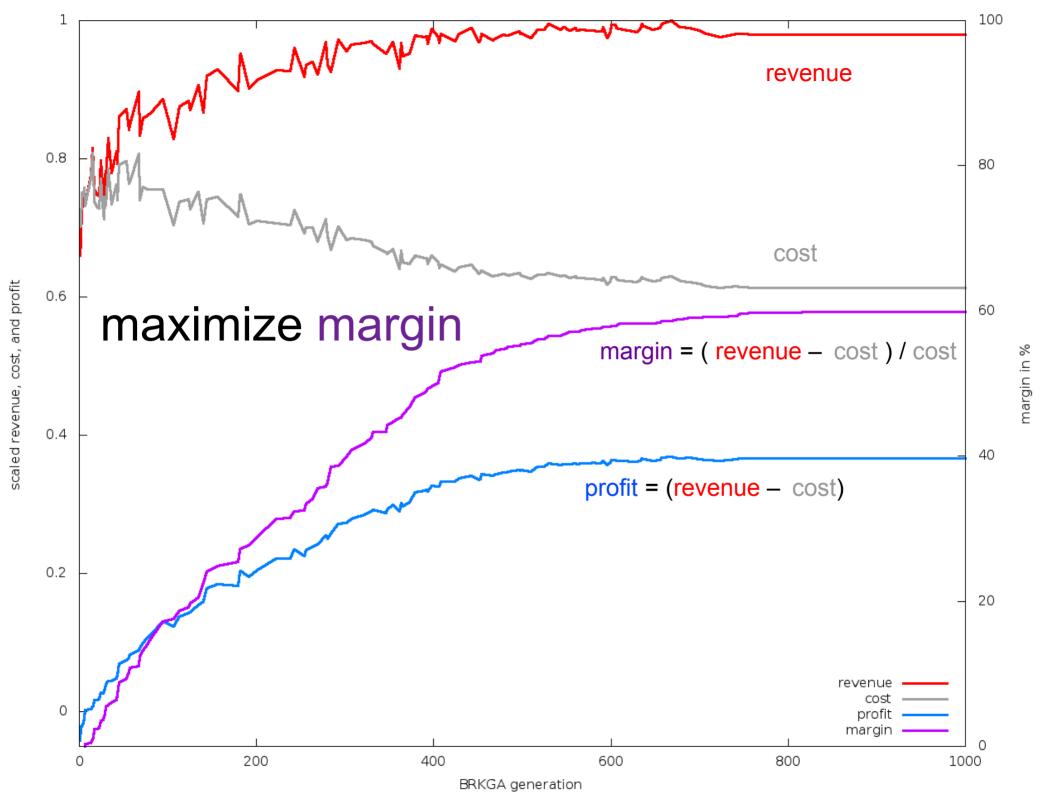
Revenues and costs can be computed to determine objective function value.

An example optimization

We ran the BRKGA on a "real" instance from a large Tier 1 Internet Service Provider

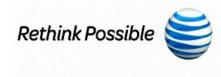
- Number of utility poles: 4395
- Number of demand points: 3236
- Number of fibered access points (FAP) excluding macrocells: 482
- Number of macrocells: 31
- Objective function: maximize margin = (revenue cost)/cost





Concluding remarks

- We described the prize collecting directed k-hop Steiner forest (PCk-HSF) problem
- We modeled a wireless backhaul network planning problem as a PCk-HSF problem with additional constraints
- We described a biased random-key genetic algorithm (BRKGA)
 for the wireless backhaul network planning problem focusing on
 the decoder
- We applied the BRKGA to a "real" instance of the wireless backhaul network planning problem



The End

These slides are available at http://mauricioresende.com

