

Generating Random Telecommand Test Data Using Genetic Algorithms

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Data Systems in Aerospace DASIA 2018

May 29th-31st, 2018, Oxford, UK

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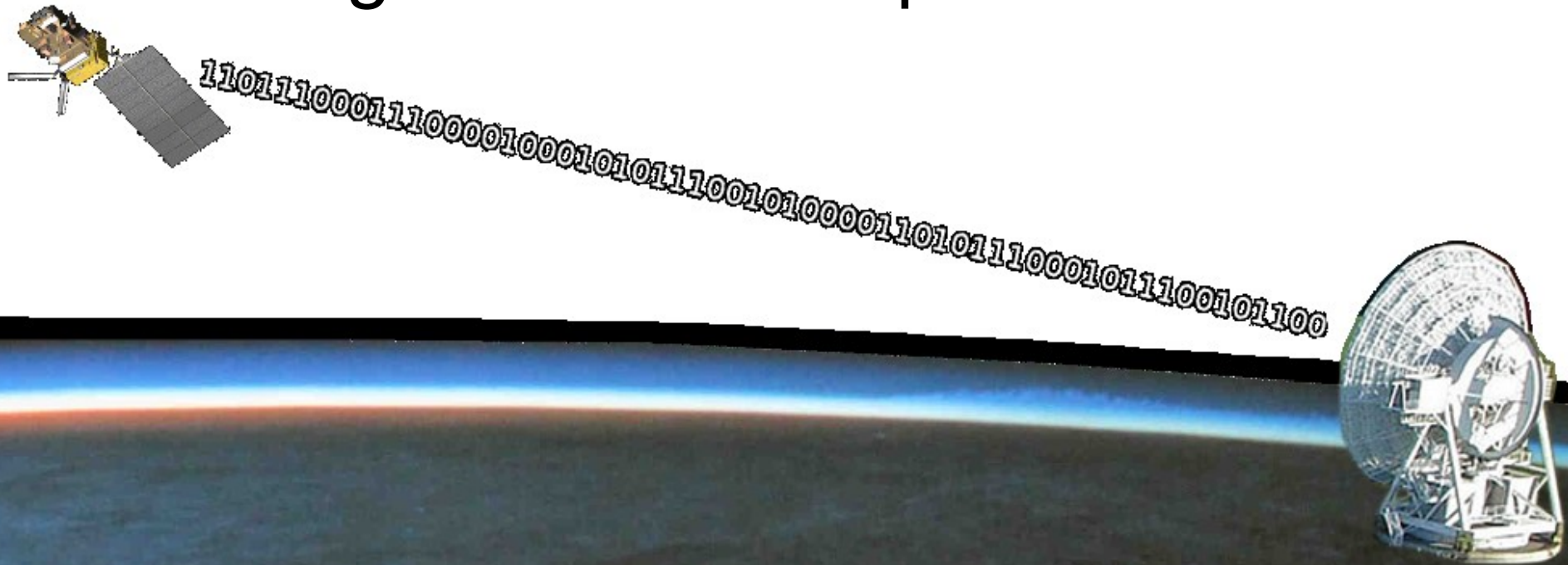
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This work is supported by a grant from the
German Federal Ministry for Economic Affairs and Energy,
based on a decision of the German Bundestag,
grant No. 50PS1601.

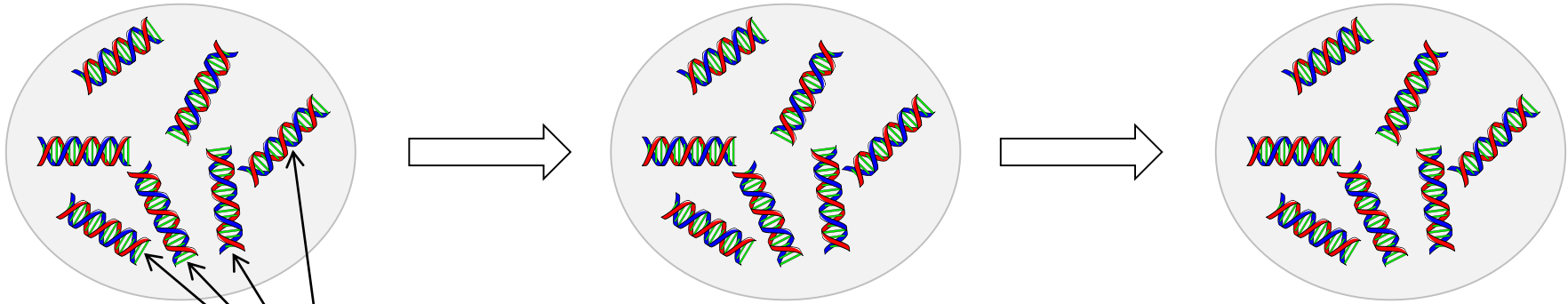
- Introduction to Genetic Algorithms
- Chosen Approach
- Evaluation
- Conclusions and Outlook

- Spacecraft processes Telecommands from Ground Station, rejects invalid commands
- Commands arrive as untyped bytestreams
- How to generate test inputs?

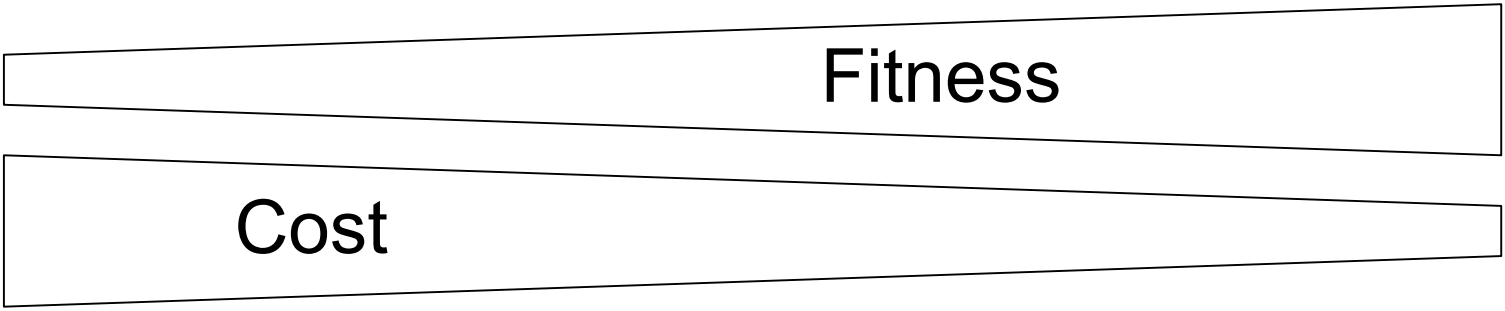


Introduction to Genetic Algorithms

Generation n Generation n+1 Generation n+2

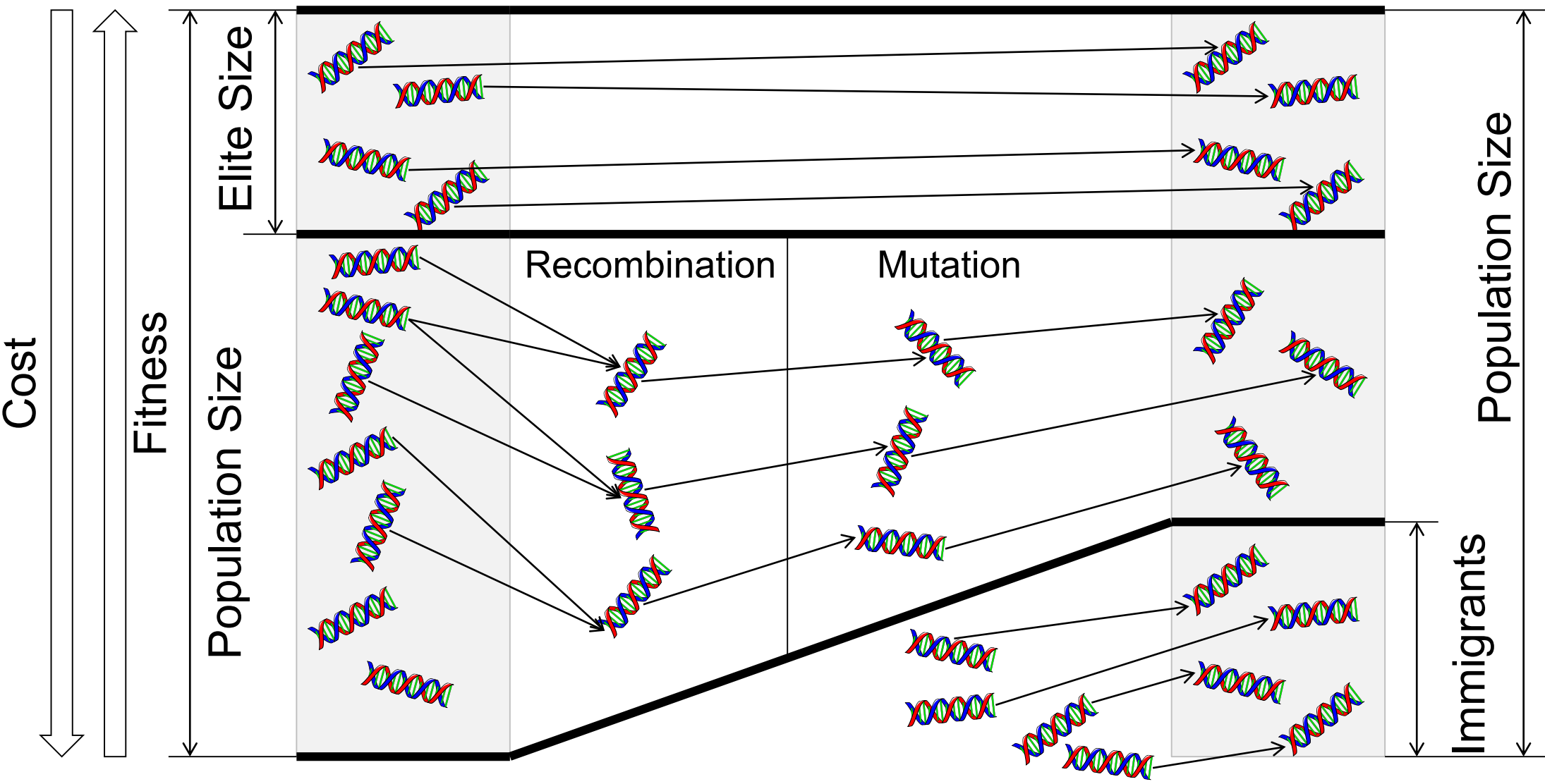


Solution
Candidates

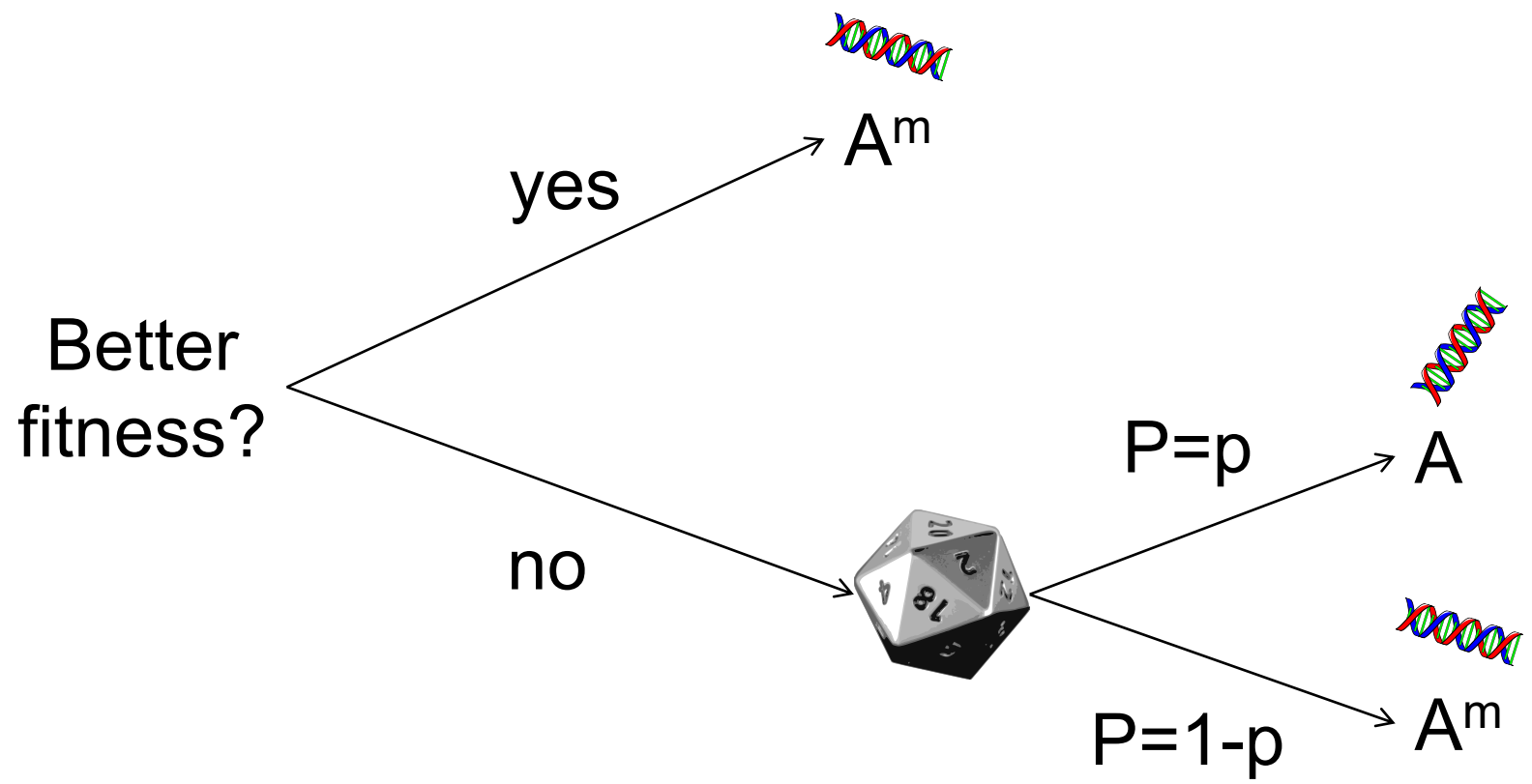
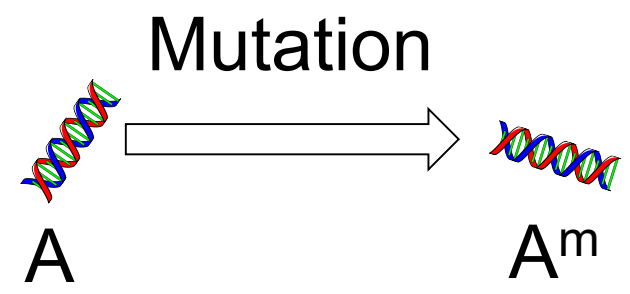


Progression of Generations

Generation n → Generation n+1

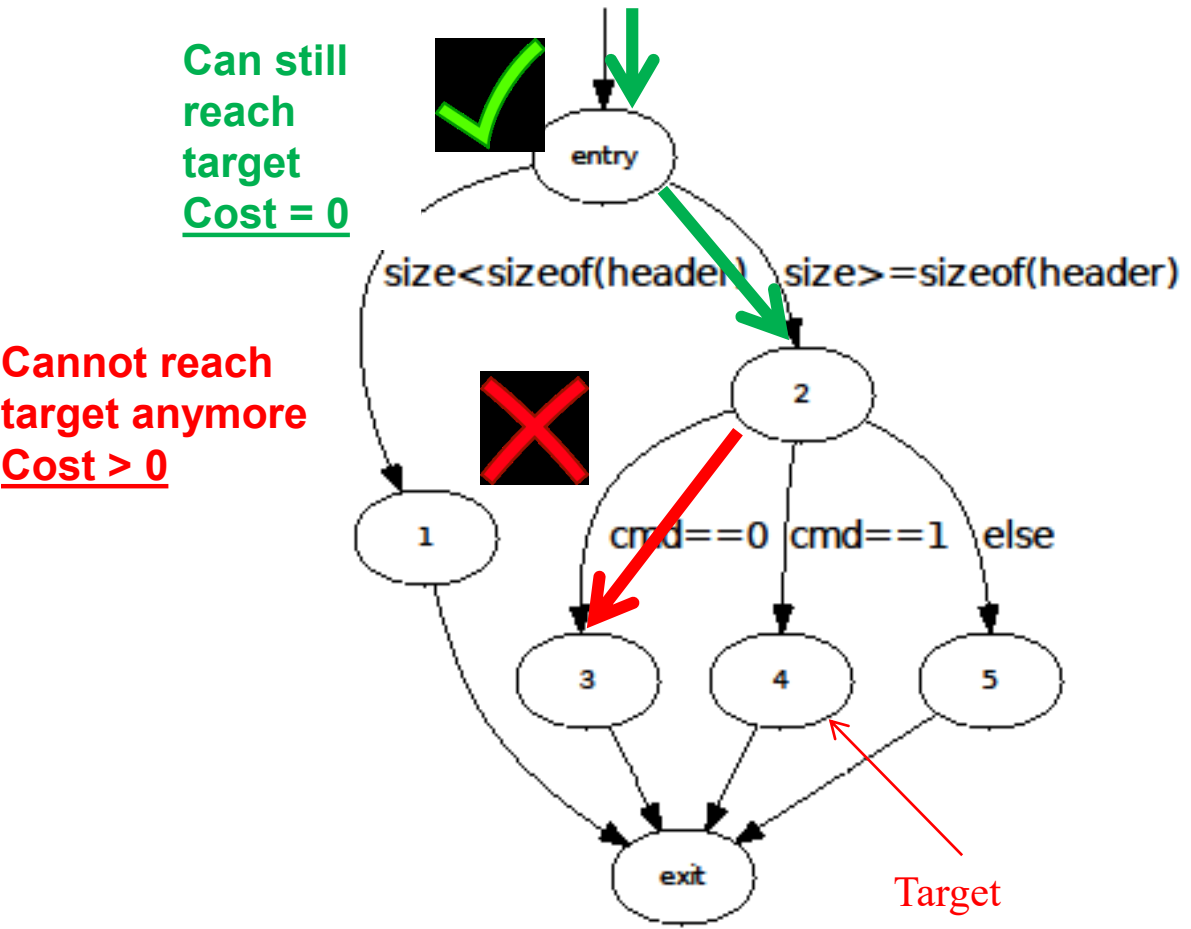


Probabilistic Mutation Reversal



Chosen Approach

Cost-Function

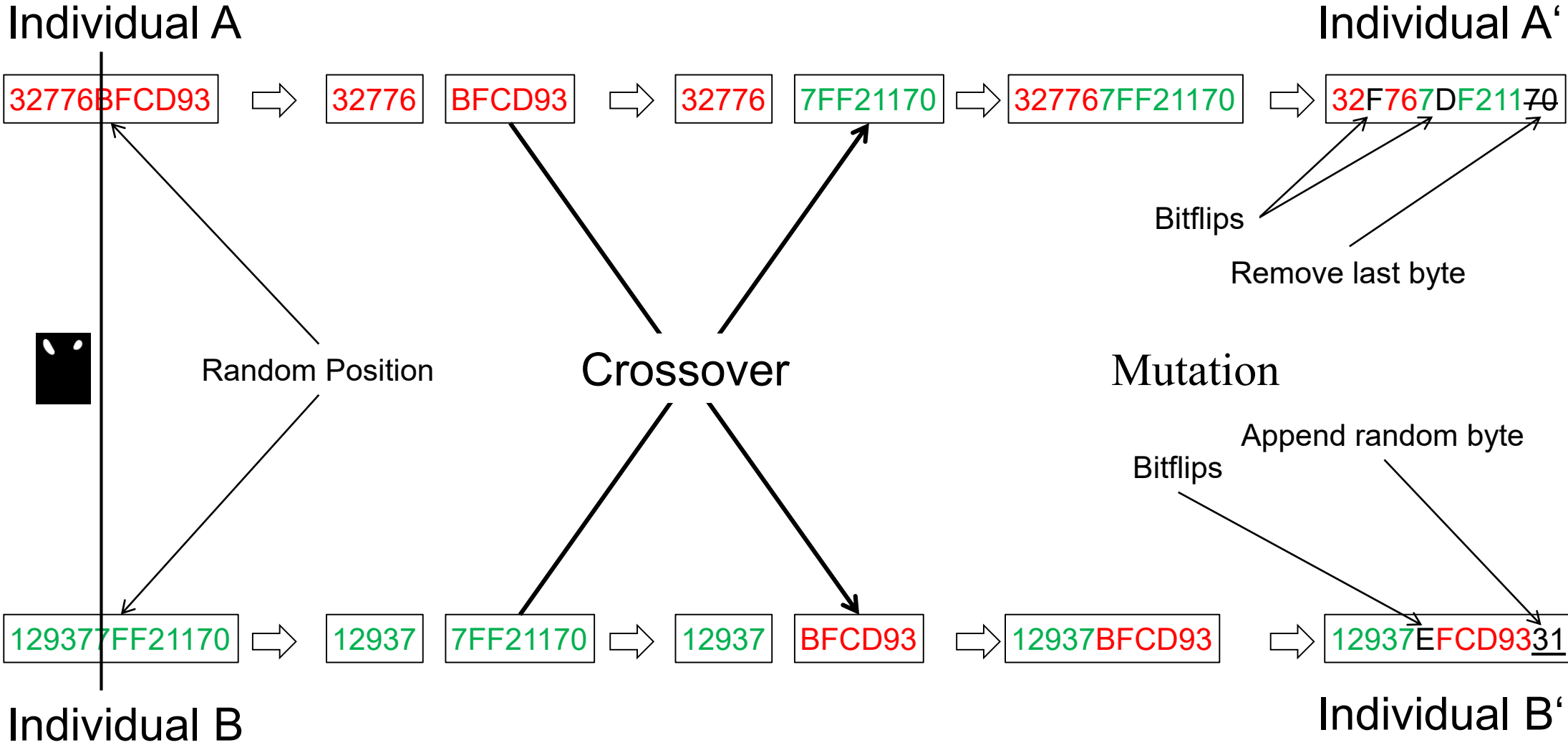


Cost Function describes how far individual is off from reaching target

| Desired Condition | Cost Function |
|--|--|
| $E \text{ op } F$ ($\text{op} \in \{=, <, >, \leq, \geq\}$) | $\begin{cases} 0 & \text{if } E \text{ op } F \\ F - E + 1 & \text{otherwise} \end{cases}$ |
| $E \neq F$ | $\begin{cases} 1 & \text{if } E = F \\ 0 & \text{otherwise} \end{cases}$ |

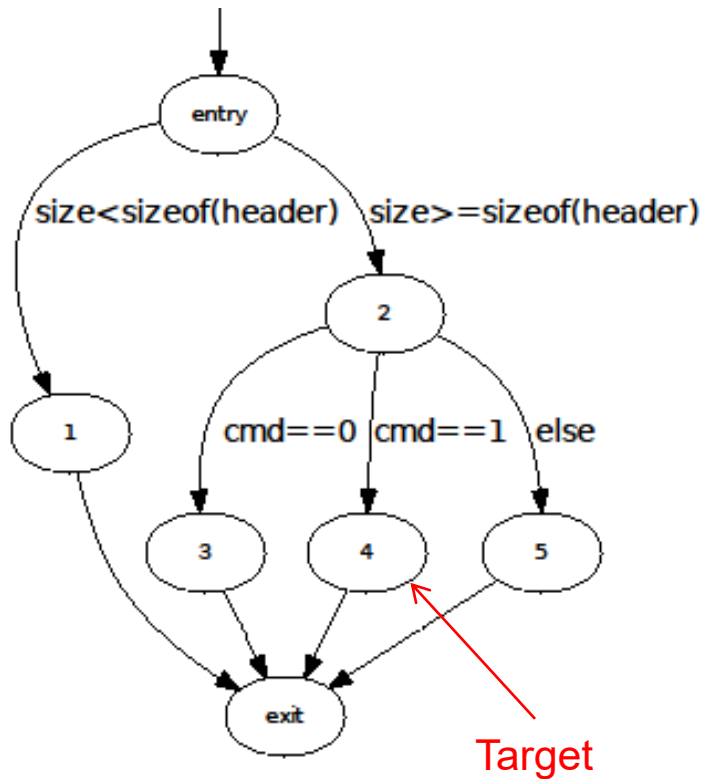
Singularity: No gradient!
Issue for boolean variables as well!

Crossover and Mutation

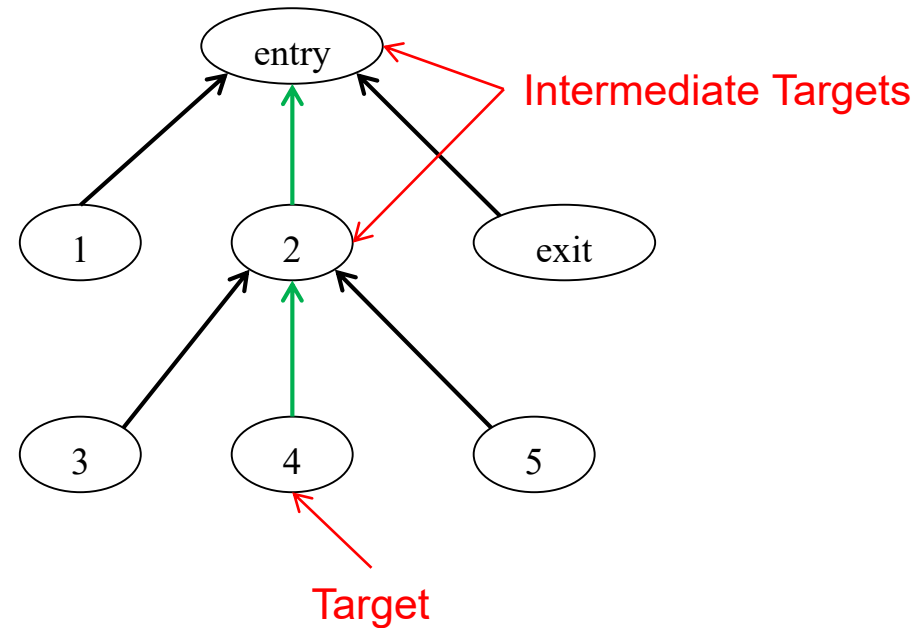


Sequential Approach: Intermediate Target

Control-Flow-Graph



Dominator Graph



- Single Step Approach: Use final target only
- Sequential Approach: Use intermediate targets

Evaluation

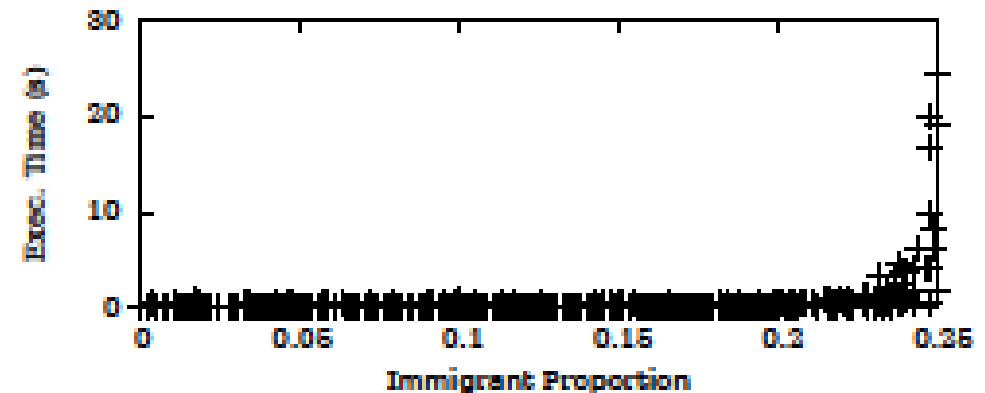
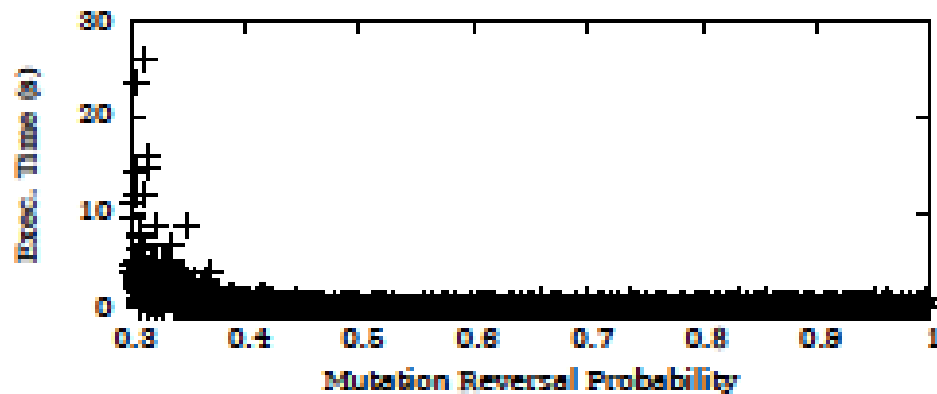
- Analyse Feasibility
- Compare runtime of algorithm variants
- Determine impact of parameters on runtime
 - Population Size
 - Elite Proportion
 - Immigrant Proportion
 - Mutation Probabilities
 - Mutation Reversal Probability
- So far only on simple example

Runtime Measurements: Algorithm Variants

| Variant | Min (s) | Mean (s) | Max (s) |
|-------------|---------|----------|---------|
| Sequential | 0.161 | 2.595 | 15.931 |
| Single-Step | 0.268 | 15.553 | 146.180 |

- Sequential is way faster than Single-Step
 - Mean runtime: Factor 6
 - Maximum runtime: Factor 9

Runtime Measurements: Parameters (1/2)



- Mutation Reversal has positive effect on runtime
- Does not seem to extend past $p \gtrsim 0.4$
- Immigrant Proportion has no noticeable effect for $p \lesssim 0.2$
- Negative effect for $p \gtrsim 0.2$

Runtime Measurements: Parameters (2/2)

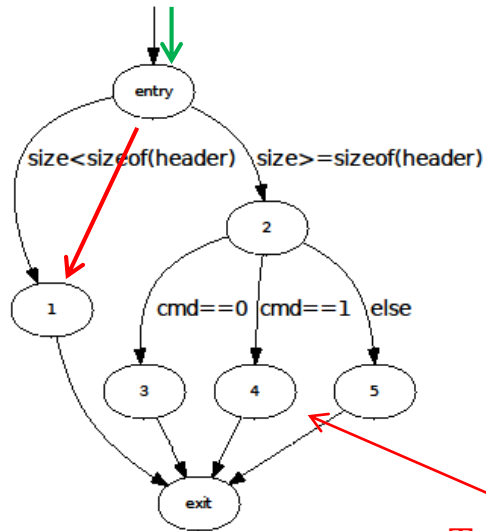
| Byte Extension Prob. | 0.00 | 0.25 | 0.50 | 0.75 | 1.00 |
|-------------------------|------|------|------|------|------|
| Mean Execution Time (s) | 0.46 | 0.13 | 0.12 | 0.12 | 0.13 |

- Optimum Value in the range $0.5 \lesssim p \lesssim 0.75$
- Decrease of Factor 3 from $p=0$ to $p=0.25$
- No further noticeable change for $p \rightarrow 1$

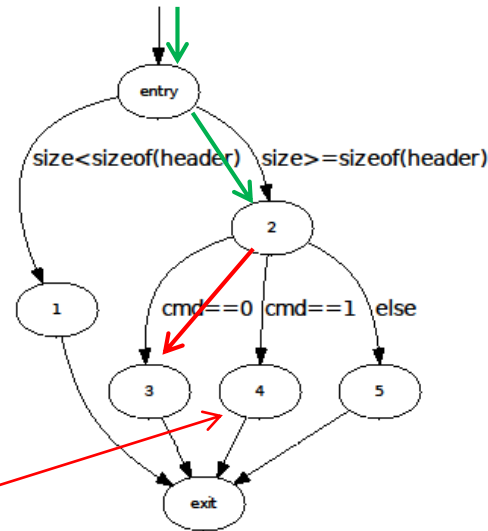
Conclusions and Outlook

Is Cross-Over Necessary?

Individual A



Individual B



Target

- How could A improve B by cross-over?
- Cost-Function does not consider branches not taken!

- Utility of Cross-Over is based on compositionality
- Test Candidates might not be compositional regarding reaching the target
- Multi-candidate gradient descent better?

- Sequential Approach is superior
- Significant positive impact of mutation reversal
- No positive effect from immigration, negative effect for more than 20% of population
- This genetic algorithm seems promising for generating telecommand test data
- Possibility for simplification: drop crossover

- Integration with random test tool (DCRITT) is underway...
- Application to industry-grade code
- Comparison to random testing performance
- More detailed analysis of parameter impacts
 - Multivariate
 - Measurements on more realistic code

Is there a general „good enough“ set of parameters for all applications?

Or do we need to „babysit“ the search?

Thank you for your attention!

Questions?

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Backup