

# RoyalRoads

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# Paper introduction

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# 1. History of the work

Reminder of the Building Block Hypothesis : A genetic algorithm (will be referred as GA in the next slides) seeks

optimal performance through the juxtaposition of short, loworder, high-performance schemata, called the building blocks.

- No detailed description on how combination occurs

  - > Design of fitness landscapes : Royal Blocks functions

- What Makes a Problem Hard for a Genetic Algorithm? Some Anomalous Results and Their Explanation – 1992 by Michel and Forrest.
- The Royal Road for Genetic Algorithms: Fitness Landscapes and GA Performance – 1993 Mitchell, Forrest and Holland.
- When will a Genetic Algorithm Outperform Hill Climbing – 1994 by Michel, Forrest and Holland
- Mitchell Royal Roads, An Introduction to GAs, The MIT Press, 1996

## 2. Major idea in the paper

- Suggestion of two features of fitness landscapes :
  - the presence of short, low-order, highly fit schemas
  - the presence of intermediate “stepping stones” —intermediate-order higher-fitness schemas



- Then, what is  $R1(111\dots1) = ?$
- With this method, GA should outperform simple hill-climbing schemes, no ?

# Testing of GA algo

- Comparison with:
  - Steepest-ascent hill climbing (SAHC)
  - Next-ascent hill climbing (NAHC)
  - Random-mutation hill climbing (RMHC)



# Steepest-ascent hill climbing (SAHC)

- 1. Choose a string at random. Call this string current-hilltop.
- 2. Going from left to right, systematically flip each bit in the string, one at a time, recording the fitnesses of the resulting one-bit mutants.
- 3. If any of the resulting one-bit mutants give a fitness increase, then set current-hilltop to the one-bit mutant giving the highest fitness increase (Ties are decided at random.)
- 4. If there is no fitness increase, then save current-hilltop and go to step 1. Otherwise, go to step 2 with the new current-hilltop.
- 5. When a set number of function evaluations has been performed (here, each bit flip in step 2 is followed by a function evaluation), return the highest hilltop that was found.

# Next-ascent hill climbing (NAHC)

- 1. Choose a string at random. Call this string current-hilltop.
- 2. For  $i$  from 1 to  $l$  (where  $l$  is the length of the string), flip bit  $i$ : if this results in a fitness increase, keep the new string, otherwise flip bit  $i$  back. As soon as a fitness increase is found, set current-hilltop to that increasedfitness string without evaluating any more bit flips of the original string. Go to step 2 with the new current-hilltop, but continue mutating the new string starting immediately after the bit position at which the previous fitness increase was found.
- 3. If no increases in fitness were found, save current-hilltop and go step 1.
- 4. When a set number of function evaluations has been performed, return the highest hilltop that was found.

# Random-mutation hill climbing (RMHC)

- 1. Choose a string at random. Call this string best-evaluated.
- 2. Choose a locus at random to flip. If the flip leads to an equal or higher fitness, then set best-evaluated to the resulting string.
- 3. Go to step 2 until an optimum string has been found or until a maximum number of evaluations have been performed.
- 4. Return the current value of best-evaluated.

# Results of GA against HC algorithms

Table 4.1 Mean and median number of function evaluations to find the optimum string over 200 runs of the GA and of various hill-climbing algorithms on  $R_1$ . The standard error ( $\sigma / \sqrt{\text{number of runs}}$ ) is given in parentheses.

200 runs	GA	SAHC	NAHC	RMHC
Mean	61,334 (2304)	> 256,000 (0)	> 256,000 (0)	6179 (186)
Median	54,208	> 256,000	> 256,000	5775

# Under what conditions will a GA outperform other search algorithms, such as hill climbing?

- Why is RMHC better ?
- Hitchhiking :
  - Happens When :
    - an instance of a higher-order schema is discovered
    - Implicates -> its high fitness allows the schema to spread quickly in the population, with zeros in other positions in the string
    - Result -> slow discovery of schema in other positions
- RMHC doesn't lose progress -> GA can with crossover and mutations

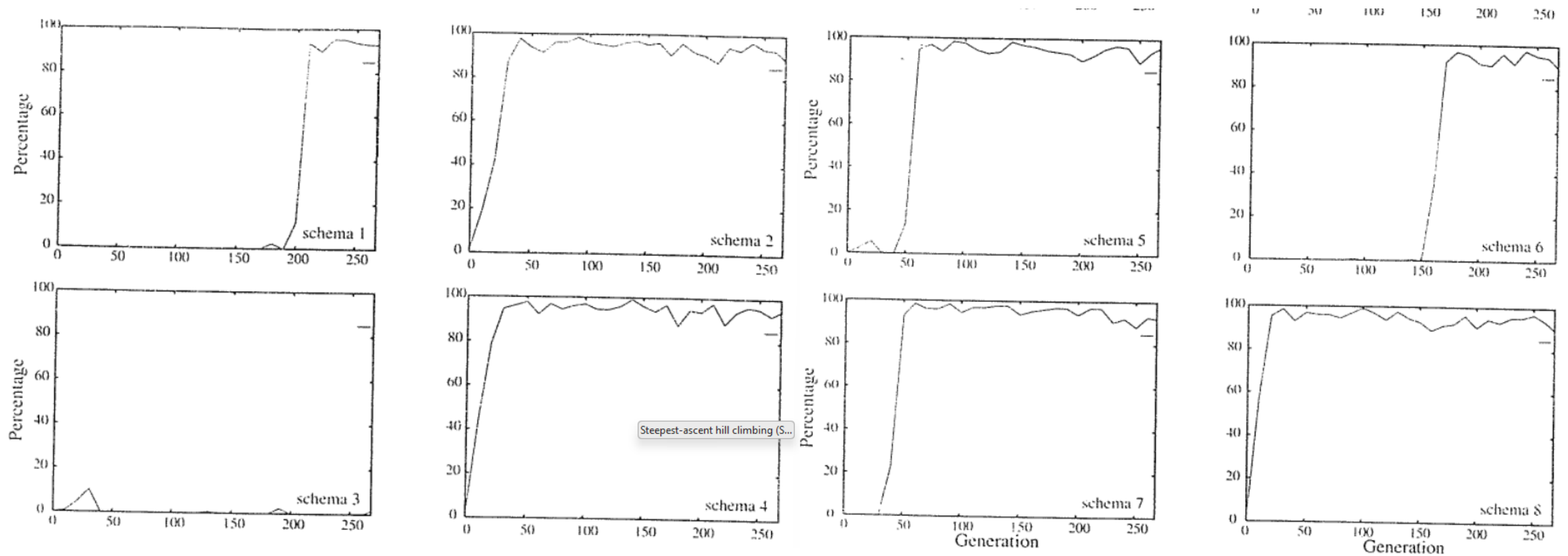


Figure 2 : Mean and median number of function evaluations to find the optimum string over 200 runs of the GA and of various hill-climbing algorithms on &. The standard error ( $\sigma / \sqrt{\text{number of runs}}$ ) is given in parentheses.

# How could we have an idealized GA

- New string random
- If good shema :
  - Keeping good building blocks
- Or:
  - Continue to take new strings
- If new good string found -> crossover with preserved string
  
- Result : IGA
- But what is the problem of this ? Why IGA can still be useful ?

# Expected time to find perfect schema with IGA

- H random schema
- p proba of finding H ( $p = 1/(2^k)$ )
- q proba of not finding  $\rightarrow q = 1 - p$
- P(t) proba finding H in time t
- $\rightarrow P(t) = 1 - q^t$
- Now for more than 1 schema to find:  $P_N(t) = (1 - q^t)^N$  (N number of schema to find)

• Then:

$$\begin{aligned} P_N(t) &= \mathcal{P}_N(t) - \mathcal{P}_N(t - 1) \\ &= (1 - q^t)^N - (1 - q^{t-1})^N. \end{aligned}$$



Via binomial theorem :

$$\begin{aligned}
 \mathcal{E}_N &= \sum_{t=1}^{\infty} t P_N(t) \\
 &= \sum_{t=1}^{\infty} t \left( (1 - q^t)^N - (1 - q^{t-1})^N \right).
 \end{aligned}
 \quad = \quad
 \begin{aligned}
 &\left[ \binom{N}{1} \left( \frac{1}{q} - 1 \right) q^t \right] - \left[ \binom{N}{2} \left( \frac{1}{q^2} - 1 \right) q^{2t} \right] \\
 &+ \left[ \binom{N}{3} \left( \frac{1}{q^3} - 1 \right) q^{3t} \right] - \dots - \left[ \binom{N}{N} \left( \frac{1}{q^N} - 1 \right) q^{Nt} \right].
 \end{aligned}$$

# Transformation of our previous result with math

$$\begin{aligned}
 & \binom{N}{1} \left(\frac{1}{q} - 1\right) \sum_{i=1}^{\infty} i q^i \\
 &= \binom{N}{1} \left(\frac{1}{q} - 1\right) (q + 2q^2 + 3q^3 + \dots) \\
 &= \binom{N}{1} \left(\frac{1}{q} - 1\right) q (1 + 2q + 3q^2 + \dots) \\
 &= \binom{N}{1} \left(\frac{1}{q} - 1\right) q \frac{d}{dq} (q + q^2 + q^3 + \dots) \\
 &= \binom{N}{1} \left(\frac{1}{q} - 1\right) q \frac{d}{dq} \left(\frac{q}{1-q}\right) \quad \text{(using a well-known identity} \\
 & \quad \text{for } 0 \leq q < 1) \\
 &= \binom{N}{1} \left(\frac{1}{q} - 1\right) q \left(\frac{1}{1-q}\right)^2 \\
 &= \binom{N}{1} \frac{1}{1-q}.
 \end{aligned}$$

$$\mathcal{E}_N \approx \frac{1}{p} \left[ \frac{\binom{N}{1}}{1} - \frac{\binom{N}{2}}{2} + \frac{\binom{N}{3}}{3} - \dots - \frac{\binom{N}{N}}{N} \right].$$

K = 8 and N = 8 like example :  
 Time is 696 -> exact result found  
 in  
 paper of 1994 (What Makes a  
 Problem Hard for a Genetic  
 Algorithm? Some Anomalous  
 Results and Their Explanation)

# IGA compared to RMHC and why it's interesting

Can evaluate the previous operation with the following expression : ->

Conclusion:

Order of IGA is  $2^K * \ln N$

Order of RMHC is  $2^K * N \ln N$  (calculated in same way, details in paper)

What conclusions can we take from that ?

$$\begin{aligned}
 \mathcal{E}_N &\approx -\frac{1}{p} \sum_{n=1}^N \binom{N}{n} \frac{(-1)^n}{n} \\
 &= \frac{1}{p} \sum_{n=1}^N \frac{1}{n} \\
 &\approx \frac{1}{p} (\ln N + \gamma) \\
 &= 2^K (\ln N + \gamma).
 \end{aligned}$$

## 4. Major result

- Royal Roads
- What stops a GA from being efficient -> hitchhiking
- Understand how and when the GA will outperform hill-climbing with comparison with IGA
- Goal is to have GA approximate as much as possible IGA
- How ? By taking features of the IGA:
  - Independent samples
  - Sequestering desired schemas
  - Instantaneous crossover
  - Speedup over RMHC
- All of this compatible ? No... everything has to be balanced ! (see The Royal Road for Genetic Algorithms: Fitness Landscapes and GA Performance)