

# Recent Developments in Autoconstructive Evolution

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# Outline

- Autoconstructive evolution
- AutoDoG (2016): 4 features and evolution evolves!
- 2 new milestones reached via 2.5 new features
- Future

# Motivation

- In nature, the ways in which evolution works ***itself evolves***, through variation and selection of mechanisms for variation and selection
- In evolutionary computation, if the evolutionary process ***can itself evolve***, then it should be capable of solving more and more difficult problems

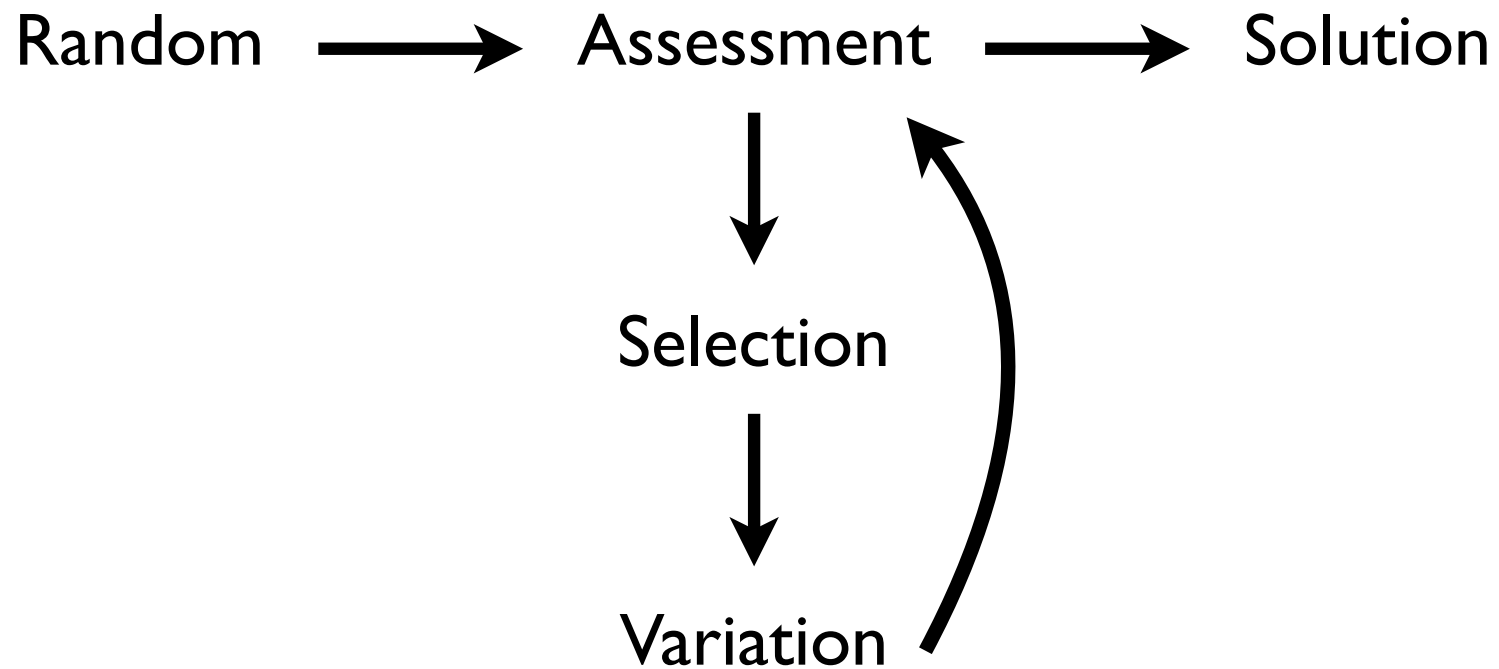
# Meta\*

- Individuals are GA/GP configurations; fitness test includes a full run of a GA/GP system
- Co-evolving populations of problem-solvers and variation operators

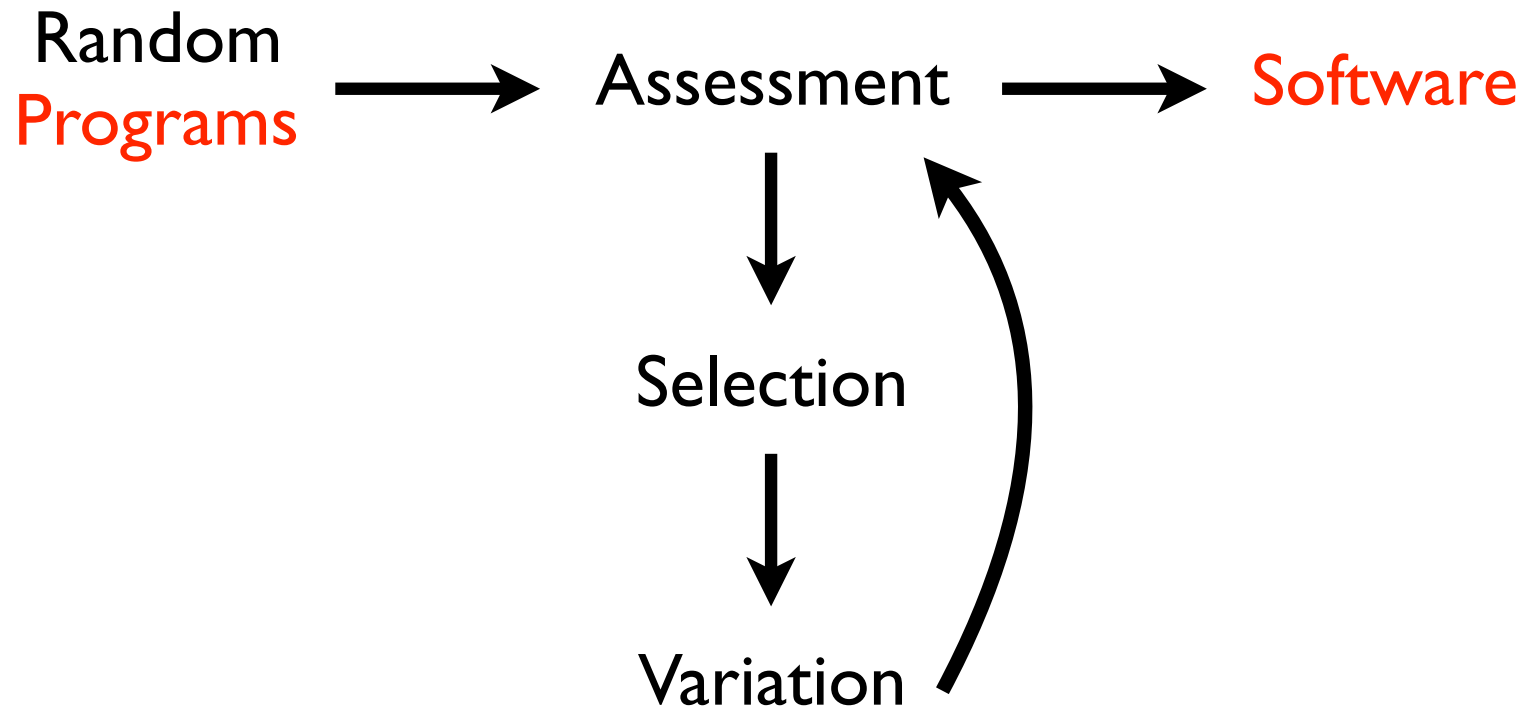
# Autoconstruction

- Individual programs make their own children
- In doing so, they control their own mutation and recombination rates and methods, and in some cases mate selection, etc.
- The machinery of reproduction and diversification (i.e., the machinery of evolution) evolves

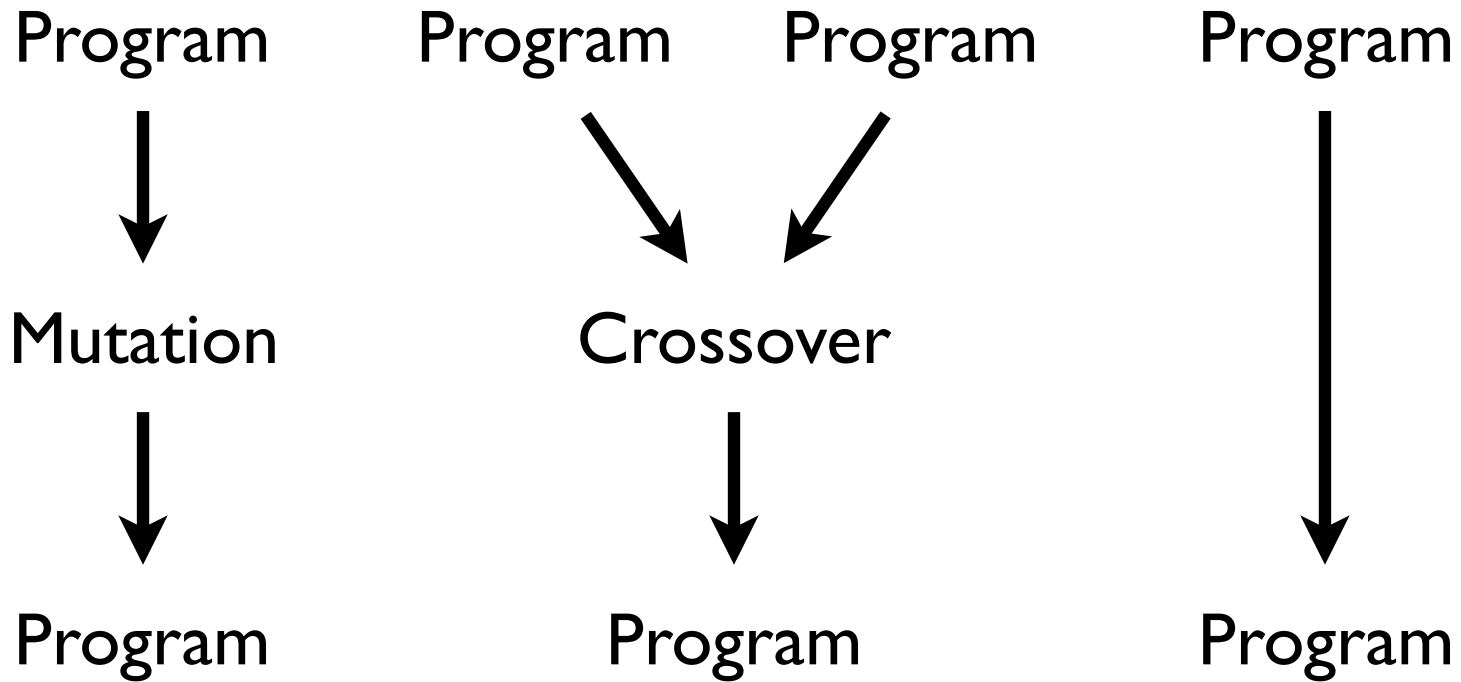
# Evolutionary Computing



# Genetic Programming



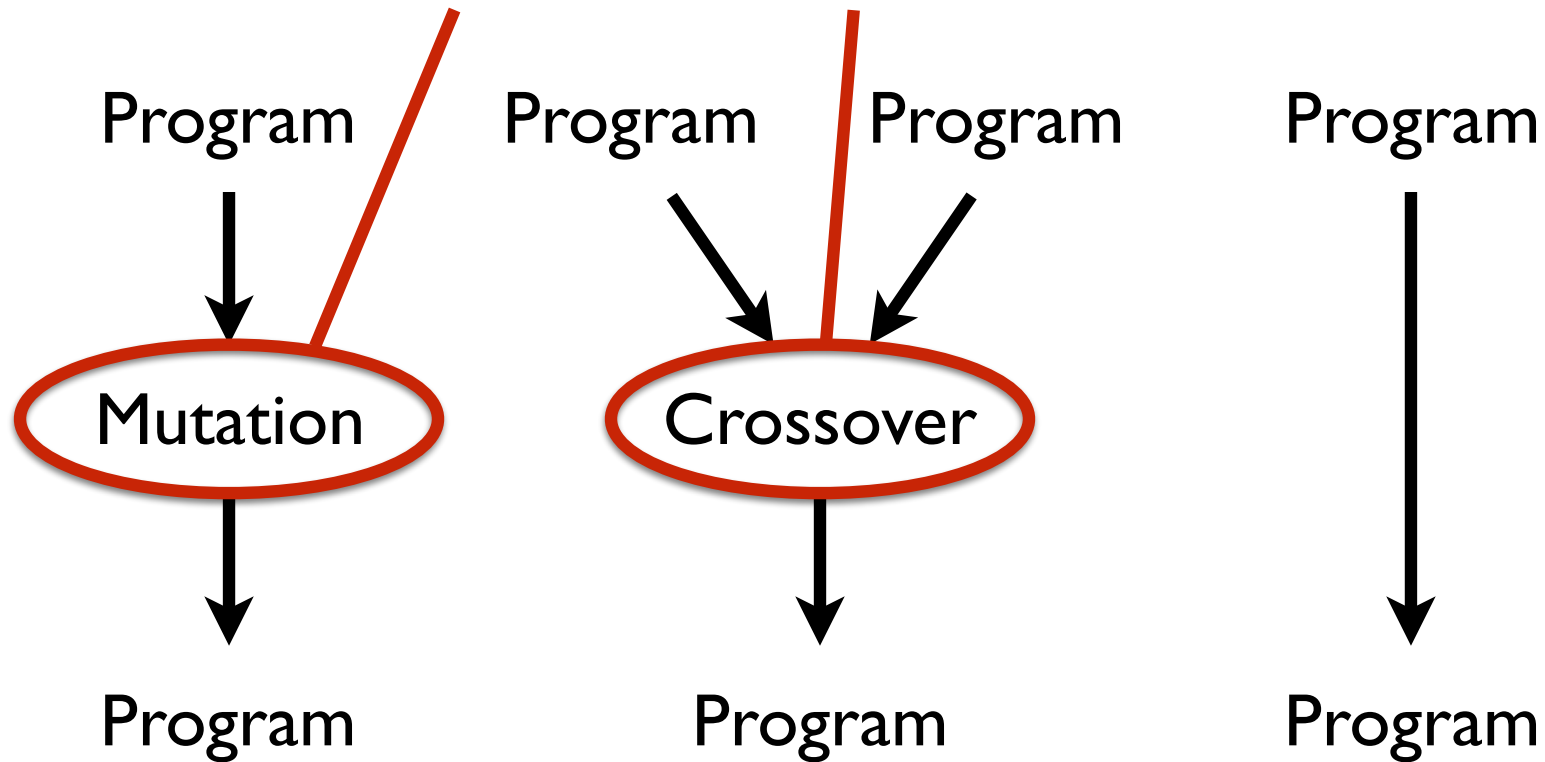
# Variation in GP



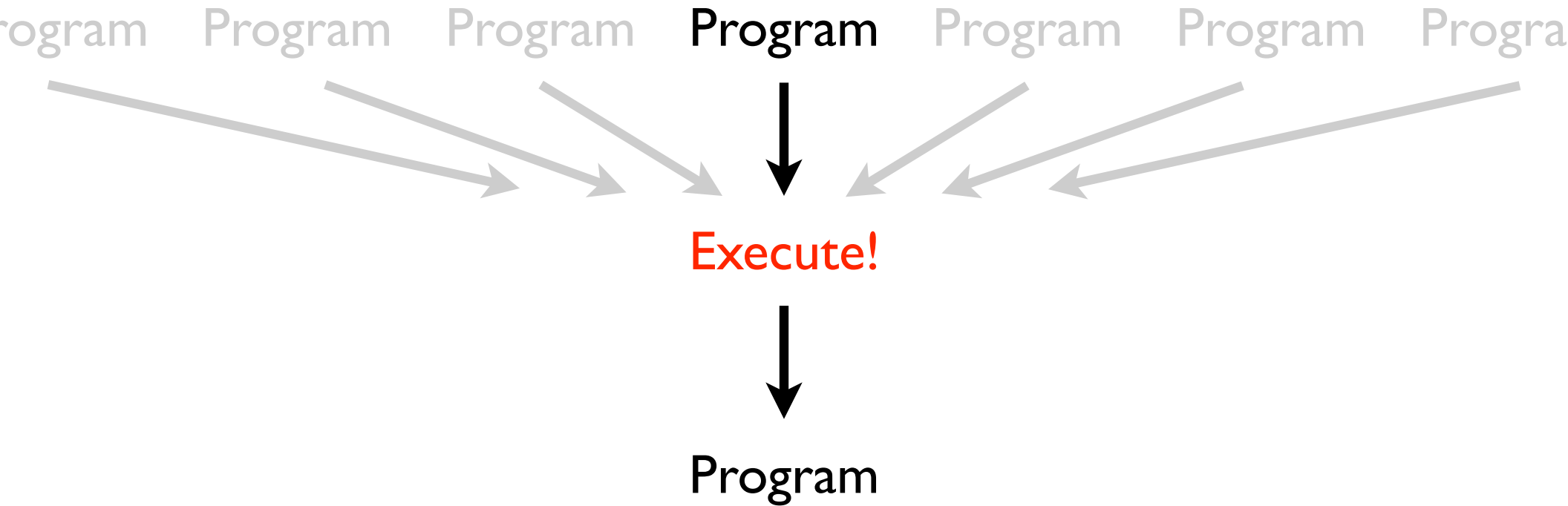


# Variation in GP

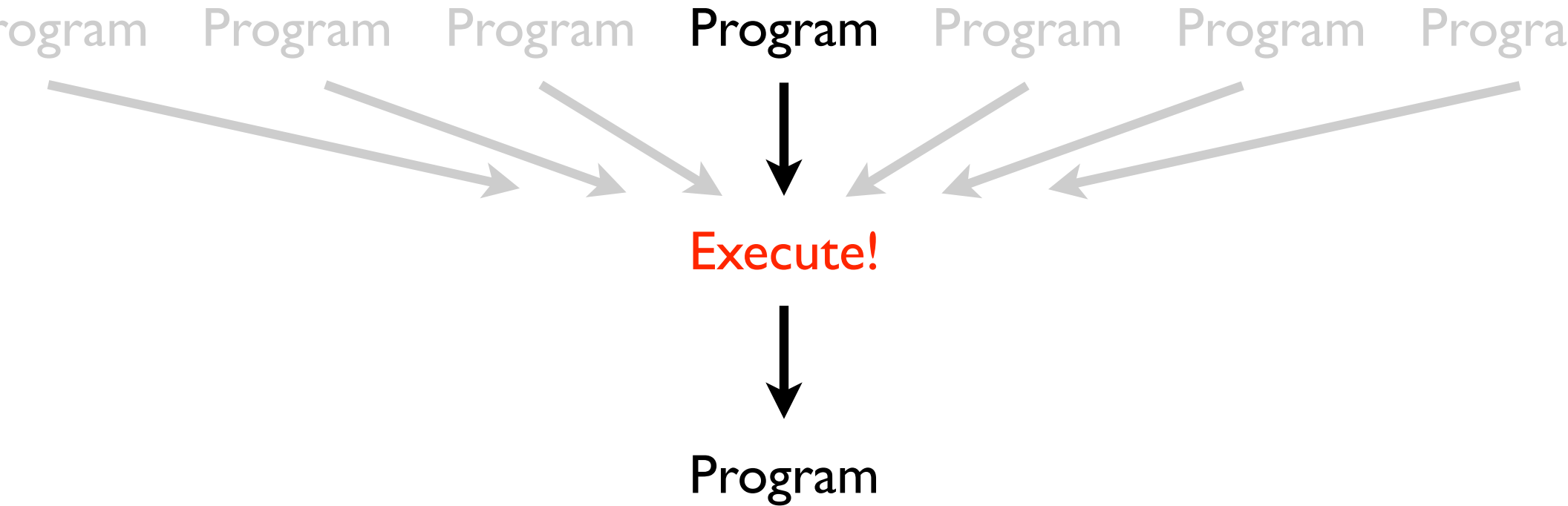
**Written and configured by humans**



# Autoconstruction



# Autoconstruction



A bit more complicated when genomes distinguished from programs

# Autoconstructive Evolution

- Evolve evolution while evolving solutions
- How? Individuals produce and vary their own children, with methods that are subject to variation
- Requires understanding the evolution of variation
- Hope: May produce EC systems more powerful than we can write by hand

# Autoconstructive Evolution

- A 15 year old project (building on older and broader-based ideas)
- Like genetic programming, but harder and less successful! But with greater potential?
- GECCO-2016: AutoDoG, sometimes solve significant problems, intriguing patterns of **evolving evolution**
- **Push** makes it easy and natural

# Push

- Programming language for programs that evolve
- Data flows via per-type stacks, not syntax
- Trivial syntax, rich data and control structures
- PushGP: GP system that evolves Push programs
- C++, Clojure, Common Lisp, Elixir, Java, Javascript, Python, Racket, Ruby, Scala, Scheme, Swift
- <http://pushlanguage.org>

# Early Autoconstruction

- Demonstrated that selection can promote diversity
- Exhibited dynamics of diversification and adaptation
- Weak problem-solving power
- Difficult to analyze results, compare to ordinary genetic programming, or generalize

# GECCO-2016 (ECADA)

## Evolution Evolves with Autoconstruction

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# AutoDoG (GECCO-2016)

**Auto**constructive **D**iversification **o**f **G**enomes

1. Construct genomes, not programs
2. Distinct mode/phase for construction of offspring
3. Select combinatorially, not on aggregate error
4. Enforce diversification constraints

## **[1. Construct genomes, not programs]**

- Previous: Push programs, on code stacks, Lisp-inspired code-manipulation instructions
- AutoDoG: Plush genomes, linear with epigenetic markers, translated to Push programs prior to running

# Plush

Instruction	integer_eq	exec_dup	char_swap	integer_add	exec_if	
Close?	2	0	0	0	1	
Silence?	1	0	0	1	0	

- Linear genomes for Push programs
- Facilitates useful placement of code blocks
- Permits uniform linear genetic operators
- Allows for epigenetic hill-climbing

**Table 1: Genome instructions in AutoDoG**

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Instruction	Description
close_dec	Decrement close marker on a gene
close_inc	Increment close marker on a gene
dup	Duplicate top genome
empty	Boolean, is genome stack empty?
eq	Boolean, are top genomes equal?
flush	Empty genome stack
gene_copy	Copy gene from genome to genome
gene_copy_range	Copy genome segment
gene_delete	Remove gene
gene_dup	Duplicate gene
gene_randomize	Replace with random
new	Push empty genome
parent1	Push first parent's genome
parent2	Push second parent's genome
pop	Remove top genome
rot	Rotate top 3 genomes on stack
rotate	Rotate sequence of top genome
shove	Insert top genome deep in stack
silence	Add epigenetic silencing marker
stackdepth	Push integer depth of genome stack
swap	Exchange top two genomes
toggle_silent	Reverse silencing of a gene
unsilence	Remove epigenetic silencing marker
yank	Pull genome from deep in stack
yankdup	Copy genome from deep in stack

---

## [2. Distinct mode/phase for construction of offspring]

- Previous: Various; sometimes during error testing, sometimes with problem inputs, sometimes with imposed but controllable variation
- AutoDoG: Only within the `autoconstruction` genetic operator, entirely by the program itself
  - Construction: inputs are no-ops
  - Error testing: `rand` instructions are constants

### **[3. Select combinatorially, not on aggregate error]**

- Previous: Parents selected using standard, error aggregating methods (tournament selection)
- AutoDoG: Lexicase selection

# Lexicase Selection

- To select single parent:
  1. Shuffle test cases
  2. First test case – keep best individuals
  3. Repeat with next test case, etc.Until one individual remains
- Selected parent may be specialist, not great on average, but lead to generalists later
- Epsilon for floats; **leaky** in experiments below

# Solving Uncompromising Problems with Lexicase Selection

Thomas Helmuth, Lee Spector *Member, IEEE*, James Matheson

**Abstract**—We describe a broad class of problems, called “uncompromising problems,” characterized by the requirement that solutions must perform optimally on each of many test cases. Many of the problems that have long motivated genetic programming research, including the automation of many traditional programming tasks, are uncompromising. We describe and analyze the recently proposed “lexicase” parent selection algorithm and show that it can facilitate the solution of uncompromising problems by genetic programming. Unlike most traditional parent selection techniques, lexicase selection does not base selection on a fitness value that is aggregated over all test cases; rather, it considers test cases one at a time in random order. We present results comparing lexicase selection to more traditional parent selection methods, including standard tournament selection and implicit fitness sharing, on four uncompromising problems: finding terms in finite algebras, designing digital multipliers, counting words in files, and performing symbolic regression of the factorial function. We provide evidence that lexicase selection maintains higher levels of population diversity than other selection methods, which may partially explain its utility as a parent selection algorithm in the context of uncompromising problems.

**Index Terms**—parent selection, lexicase selection, tournament selection, genetic programming, PushGP.

## I. INTRODUCTION

GENETIC programming problems generally involve test cases that are used to determine the performance of programs during evolution. While some classic genetic programming problems, such as the artificial ant problem or lawnmower problem [1], involve only single test cases, others involve large numbers of tests. There are scenarios in which a genetic programming system can consider test cases into consideration during parent selection when determining which individuals to use when producing offspring for the next generation. The best choice may depend on the type of problem.

For some problems it may be better to seek “compromises” among

example, we can imagine a problem involving control of a simulated wind turbine in which some test cases focus on performance in low wind conditions while others focus on performance in high wind conditions. It may not be possible to optimize performance on all of these test cases simultaneously and some sort of compromise may therefore be necessary. Many common parent selection approaches, such as tournament selection, introduce compromises by aggregating the performance of an individual across all test cases into a single fitness value. This may be as simple as summing the squared errors, or as implicit fitness sharing [2].

By contrast, we wish to solve “uncompromising” problems: programs must perform as well as possible on every test case. This is a problem that is often difficult to solve for genetic programming.

# GPTP-2015

Problem name	Lexicase	Tournament	IFS
Replace Space With Newline	57	13	17
Syllables	24	1	2
String Lengths Backwards	75	18	12
Negative To Zero	72	15	9
Double Letters	5	0	0
Scrabble Score	0	0	0
Checksum	0	0	0
Count Odds	4	0	0

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# Diversity

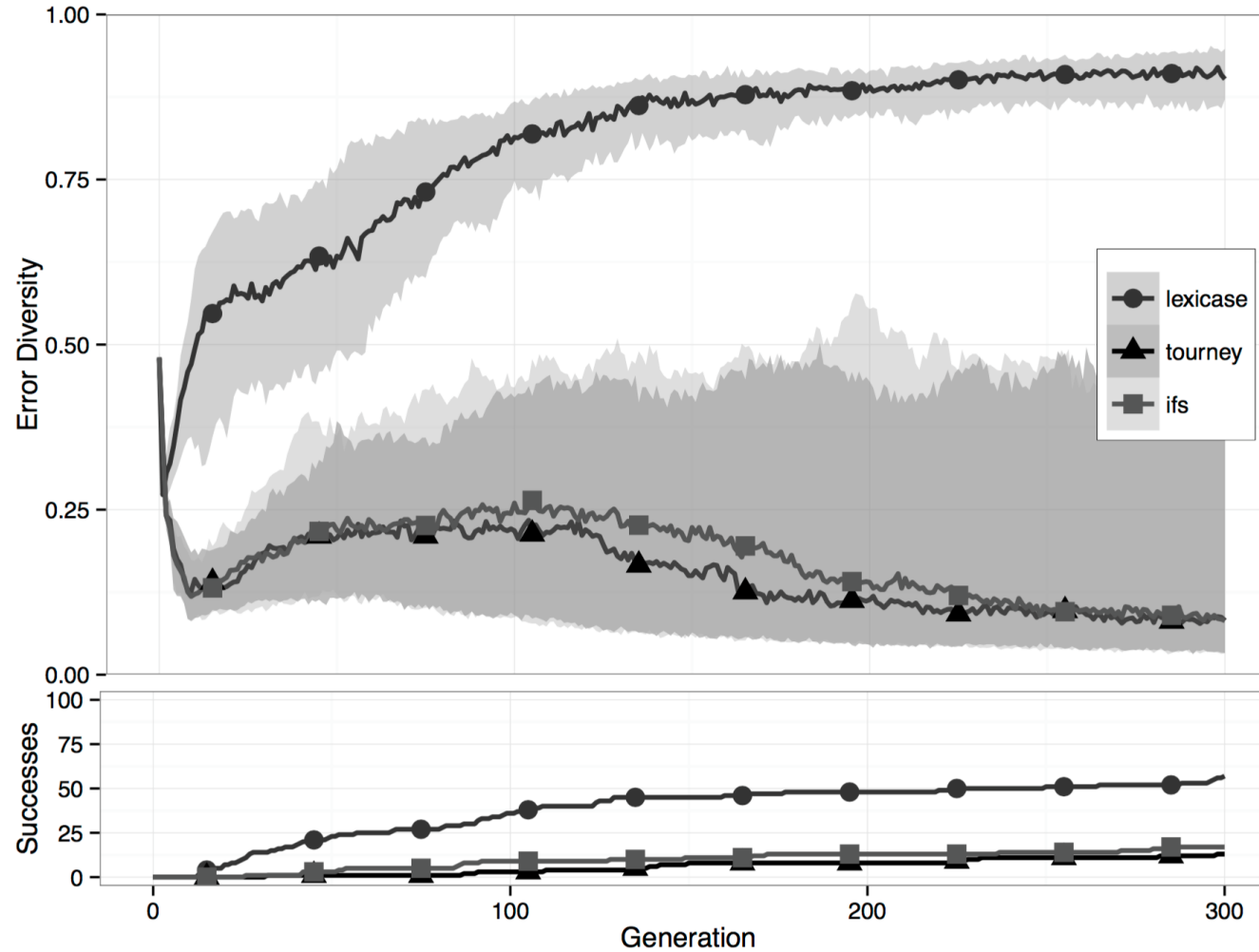


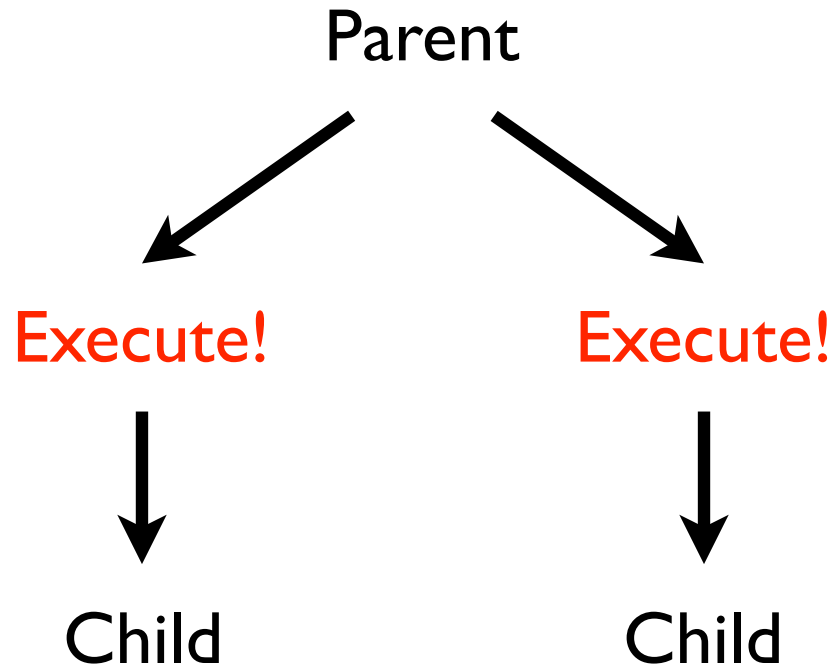
Fig. 1 Replace Space With Newline – error diversity

GPTP-2015

## **[4. Enforce diversification constraints]**

- Previous: Various, including all but clones, or those in lineages making progress
- AutoDoG: Must satisfy diversification constraints on reproductive behavior, determined from a cascade of temporary descendants

# Diversification Constraints



- Parent/child program differences positive; not same
- Many variants possible

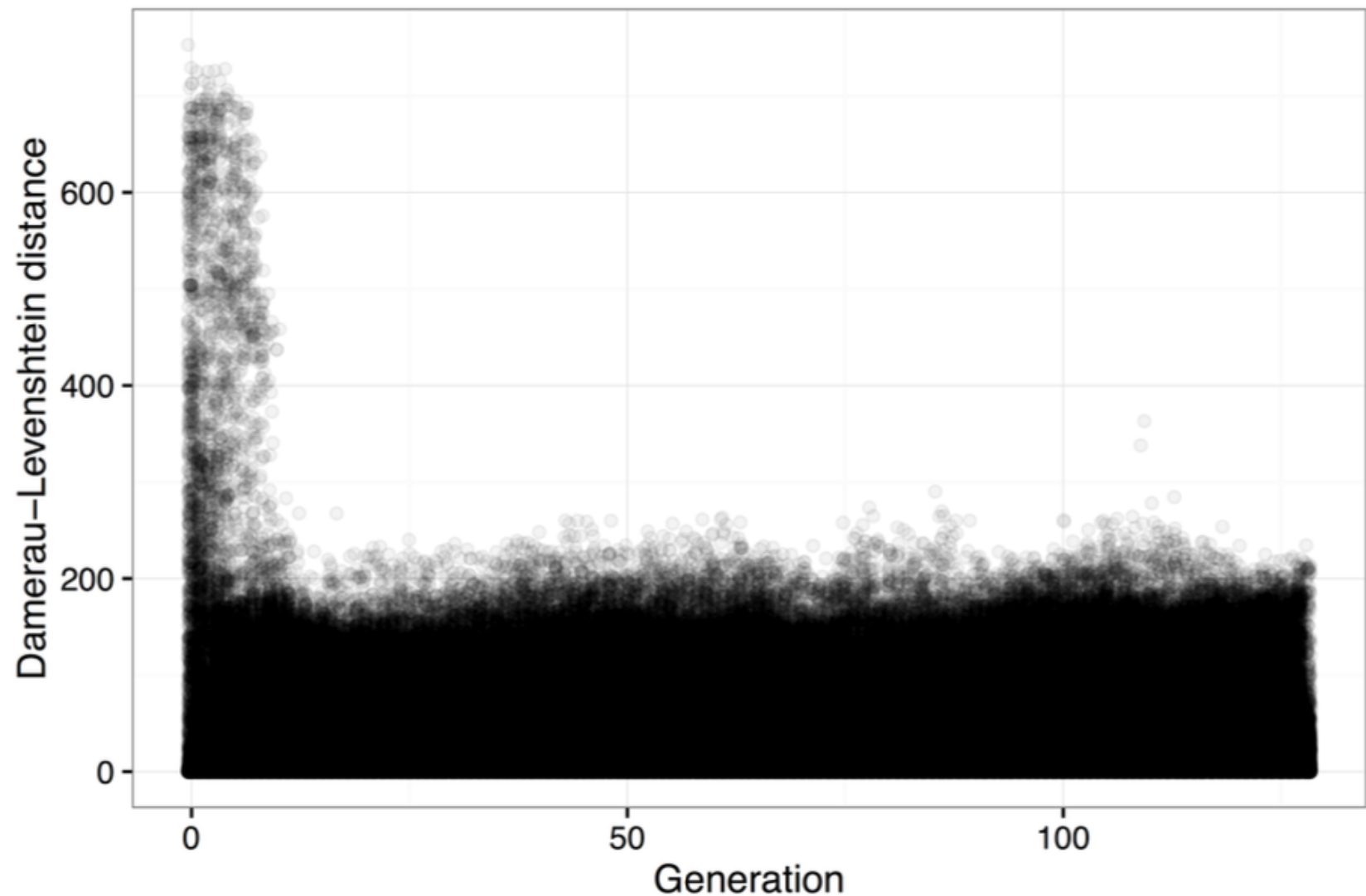
# Software Synthesis Benchmarks (GECCO 2015)

Number IO, Small or Large, For Loop Index, Compare String Lengths, Double Letters, Collatz Numbers, Replace Space with Newline, String Differences, Even Squares, Wallis Pi, String Lengths Backwards, Last Index of Zero, Vector Average, Count Odds, Mirror Image, Super Anagrams, Sum of Squares, Vectors Summed, X-Word Lines, Pig Latin, Negative to Zero, Scrabble Score, Word Stats, Checksum, Digits, Grade, Median, Smallest, Syllables

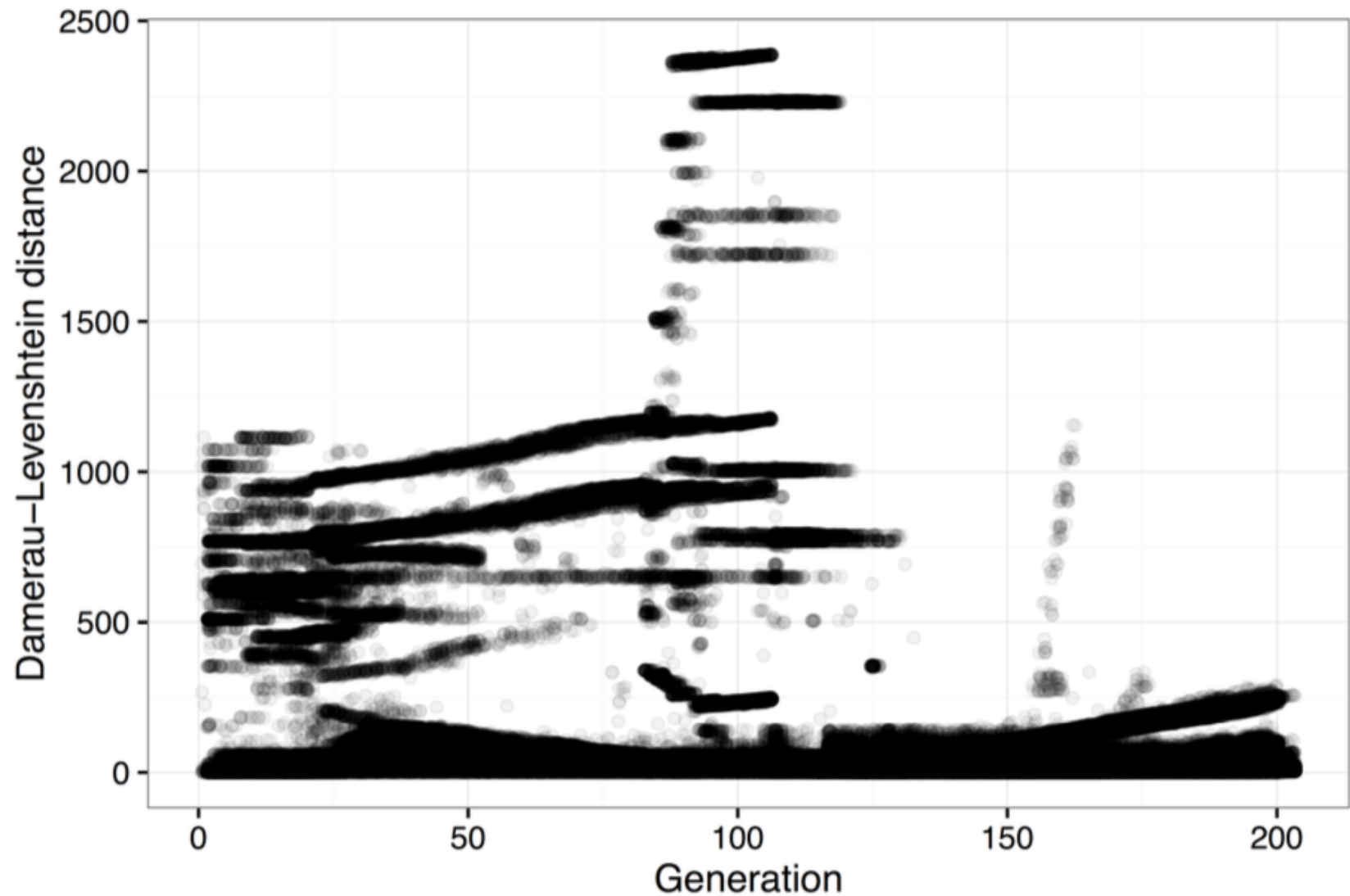
Solved with PushGP; only with autoconstruction

7. **Replace Space with Newline (P 4.3)** Given a string input, print the string, replacing spaces with newlines. Also, return the integer count of the non-whitespace characters. The input string will not have tabs or newlines.

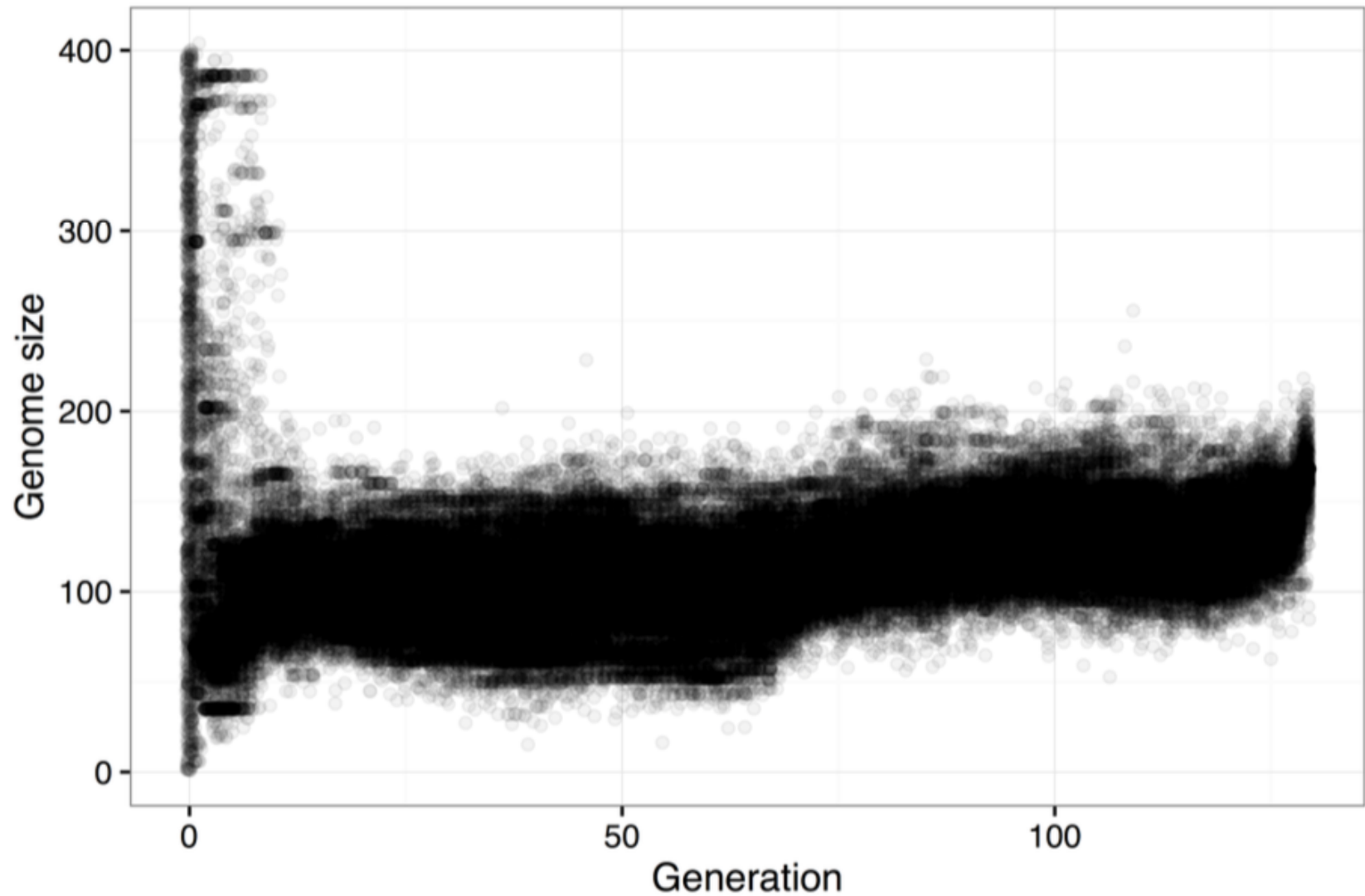
- Multiple types, looping, multiple tasks
- PushGP can achieve success rates up to ~95% in 300 generations
- AutoDoG 2016 succeeded 5-10%



**Figure 1: DL-distances between parent and child during a single non-autoconstructive run of GP on the Replace Space With Newline problem**

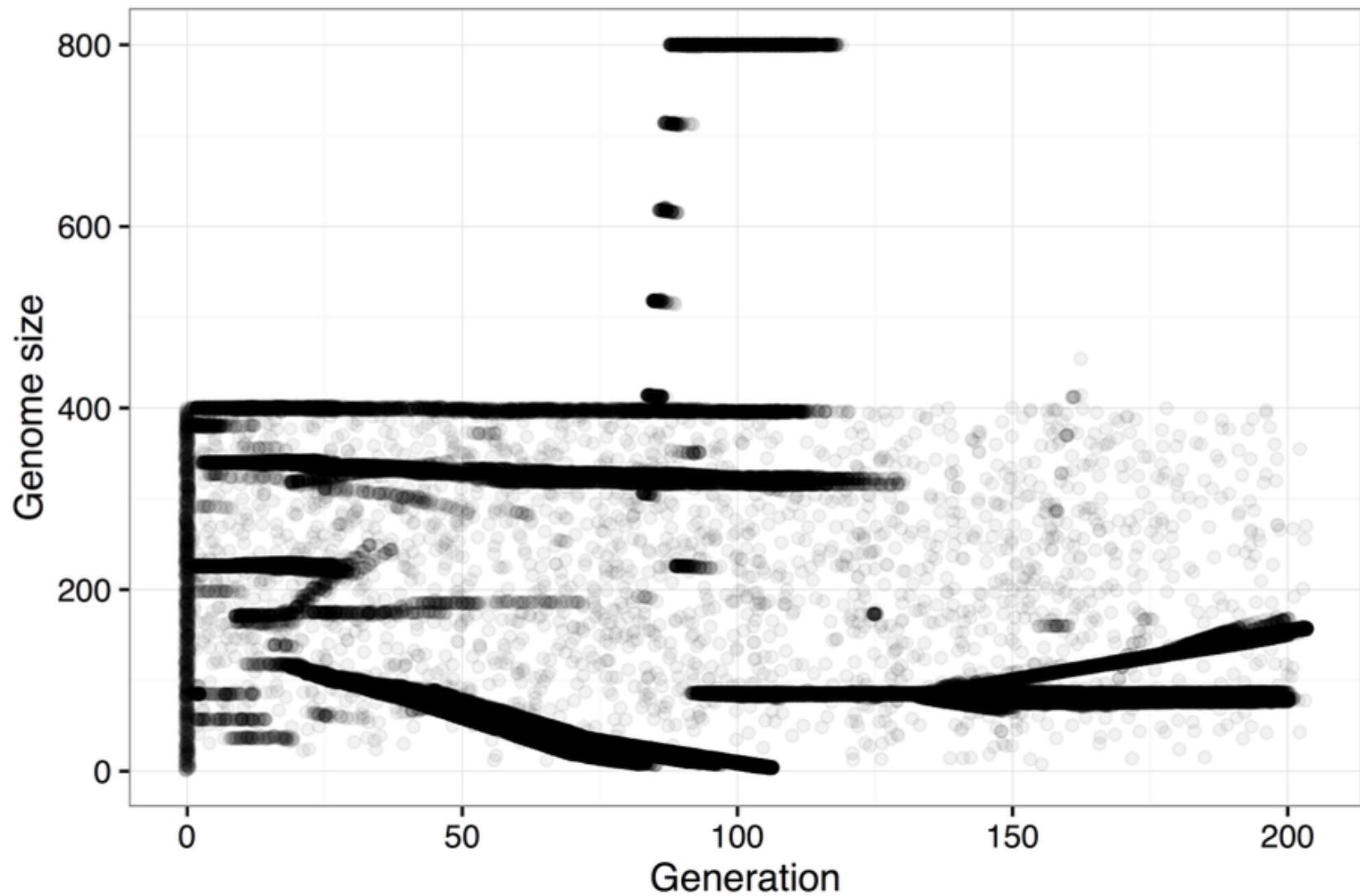


**Figure 3: DL-distances between parent and child during a single autoconstructive run of GP on the Replace Space With Newline problem**



**Figure 2: Genome sizes during a single non-autoconstructive run of GP on the Replace Space With Newline problem**

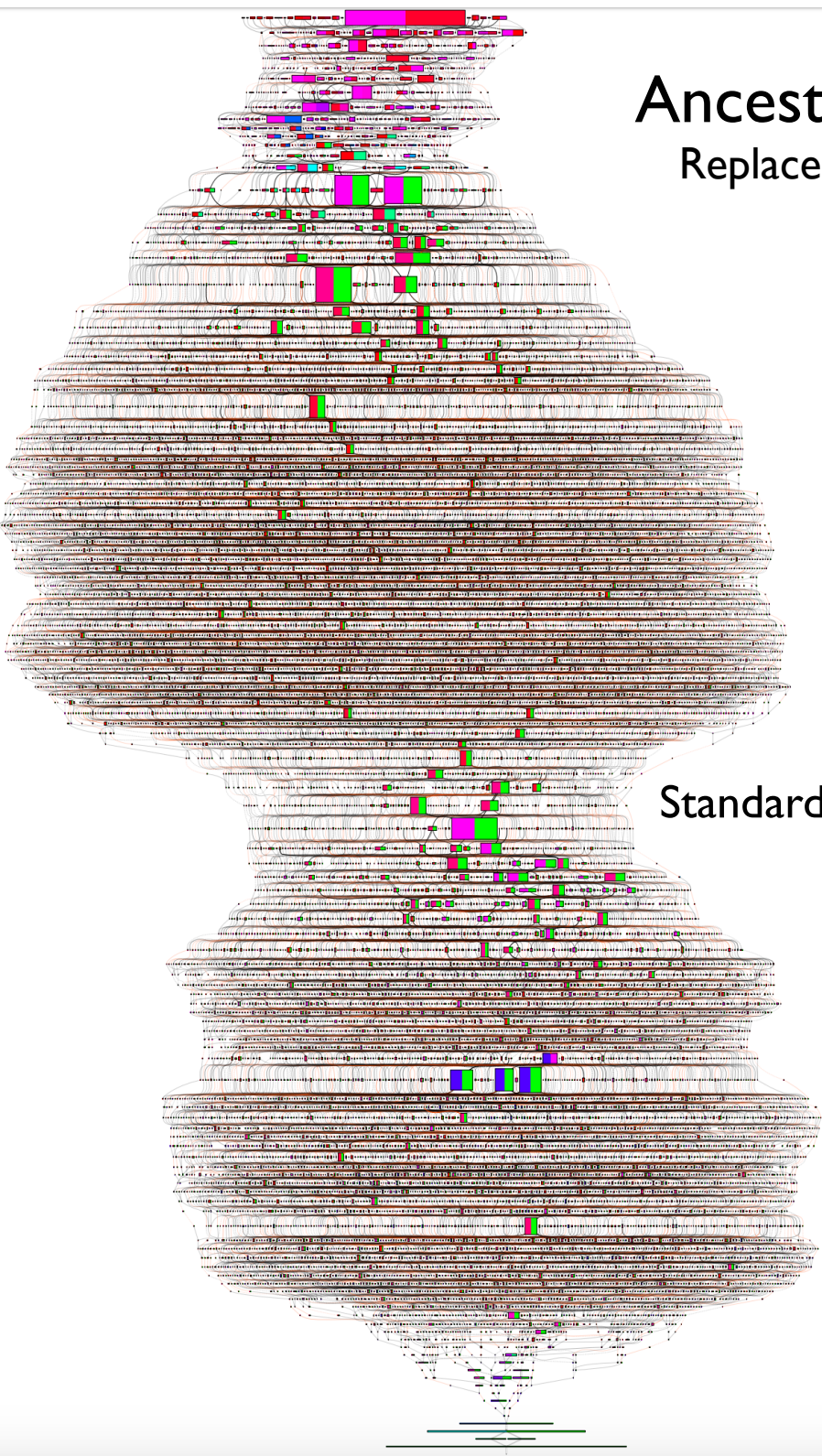




**Figure 4: Genome sizes during a single autoconstructive run of GP on the Replace Space With Newline problem**

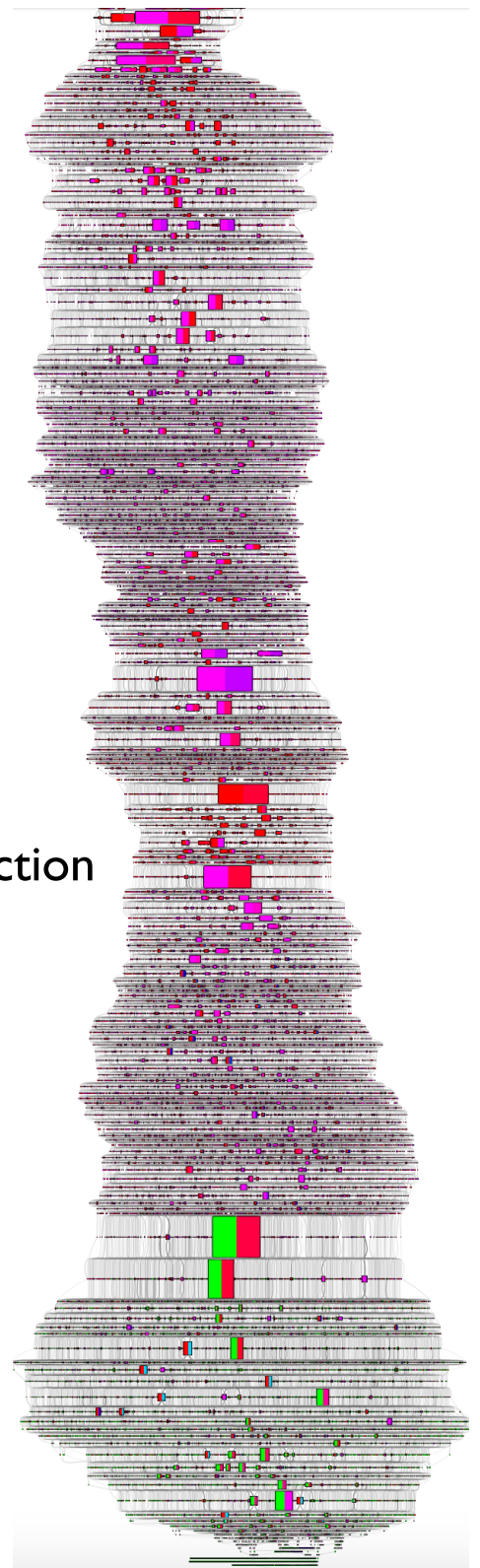
# Ancestors of Solutions

## Replace Space with Newlines



Standard Operators

Autoconstruction



# 2 New Milestones

- Autoconstructive evolution can succeed as much and as fast as non-autoconstructive evolution
- Autoconstructive evolution can solve a problem not yet solved without it

## 2.5 New Features

- DSL for uniform genome manipulation
- Entropy
- Age-Mediated Parent Selection (AMPS)

# DSL for Uniform Genome Manipulation

```
genome_alteration
genome_genesis
genome_new
genome_parent1
genome_parent2
genome_uniform_addition
genome_uniform_addition_and_deletion
genome_uniform_boolean_mutation
genome_uniform_close_mutation
genome_uniform_combination_and_deletion
genome_uniform_crossover
genome_uniform_deletion
genome_uniform_float_mutation
genome_uniform_instruction_mutation
genome_uniform_integer_mutation
genome_uniform_silence_mutation
genome_uniform_string_mutation
genome_uniform_tag_mutation
genome_dup
genome_empty
genome_eq
genome_flush
genome_pop
genome_rot
genome_rotate
genome_shove
genome_stackdepth
genome_swap
genome_yank
genome_yankdup
```

# Entropy

- Random gene deletions after autoconstruction
- Like "cosmic ray mutations" but purely destructive
- All new genetic material must stem from autoconstructive instructions
- Lineages must counteract entropy to survive
- Default rate: 0.1

<https://xkcd.com/1862/>

PARTICLE PROPERTIES IN PHYSICS

PROPERTY	TYPE/SCALE
ELECTRIC CHARGE	-1 0 +1
MASS	0 1 <sub>ks</sub> 2 <sub>ks</sub> →
SPIN NUMBER	-1 ½ 0 ½ 1
FLAVOR	(MISC. QUANTUM NUMBERS)
COLOR CHARGE	R G B (QUARKS ONLY)
MOOD	☹ ☹ ☹ ☹ ☹ →
ALIGNMENT	☐☐☐ GOOD-EVIL, LAWFUL-CHAOTIC
HIT POINTS	0 →
RATING	☆☆☆☆☆
STRING TYPE	BYTESTRING-CHARSTRING
BATTING AVERAGE	0% 100%
PROOF	0 200 →
HEAT	🔥 🔥 🔥 🔥 🔥 →
STREET VALUE	\$0 \$100 \$200 →
ENTROPY	(THIS ALREADY HAS LIKE 20 DIFFERENT CONFUSING MEANINGS, SO IT PROBABLY MEANS SOMETHING HERE, TOO)

ENTROPY	(THIS ALREADY HAS LIKE 20 DIFFERENT CONFUSING MEANINGS, SO IT PROBABLY MEANS SOMETHING HERE, TOO.)
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# Age-Mediated Parent Selection (AMPS)

- Use genealogical age to bias in favor of youth
- Like ALPS (but simpler), and age-fitness Pareto optimization (but for parent selection)
- For each parent, consider only younger than a limit chosen randomly from ages in the population
- Options for age-combining functions; for autoconstruction: age of executing parent + maximum similarity with a parent, scaled to  $[0, 1]$



# Rivaling Ordinary PushGP

- Uniform DSL + Entropy + AMPS
- In 20 runs, 75% success within 300 generations on Replace Space With Newline (100% by generation 628); 80% on Mirror Image
- Surprisingly, rivals ordinary GP on a problem that ordinary GP can solve

8. **String Differences (P 4.4)** Given 2 strings (without whitespace) as input, find the indices at which the strings have different characters, stopping at the end of the shorter one. For each such index, print a line containing the index as well as the character in each string. For example, if the strings are “dealer” and “dollars”, the program should print:

```
1 e o
2 a l
4 e a
```

# Extending the Reach of GP

- Without autoconstruction, string difference not yet solved by GP, despite many efforts/configurations
- 3 autoconstructive solutions so far, with Uniform DSL + Entropy

# First Evolved Solution

- Makes children using uniform addition, with a rate ( $\sim 0.0921$ ) close to the entropy rate (0.1)
- Solves problem in general way, with a few clever tricks (like using the depth of the boolean stack to track the comparison index)

# Future

- Use autoconstruction to solve other previously unsolved problems
- Study how autoconstruction works, to improve it
- Consider implications for study of evolution of biological evolution

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