

# Genetic Programming & Bloat

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Substantial part of this material is based on the article  
Sean Luke and Liviu Panait: A Comparison of Bloat Control Methods for Genetic Programming,  
See <http://portal.acm.org/citation.cfm?id=1182892.1182897>



<http://cw.felk.cvut.cz/doku.php/courses/a0m33eoa/start>







## Theories of Code Bloat Based on Introns

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**Hitchhiking** – based on genetic algorithms, where unfit building blocks propagate in the population because they join highly fit building blocks.

- There is no real need to get rid of hitchhikers that do not damage fitness of the program.  
Introns are hitchhikers in GP.
- The theory only suggests a propagation method.

It does not explain why it is more likely that the introns become attached in the first place than to be removed eventually.



## Theories of Code Bloat Based on Introns

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**Removal Bias** – to maintain fitness, the removed subtree must be contained within the inviable region. Since the inserted subtree can have any size, offspring are bigger than average while retaining the fitness of their parents.

- Inactive code in a GP tree tends to be low in the tree, residing therefore in smaller-than-average subtrees.
- Inserted subtree is bigger than the excised ones.

## Non-Intron Theories of Code Bloat

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**Fitness Causes Bloat** – there are many more longer ways than shorter ways to represent the same program, so a natural drift occurs to longer programs.

- Beyond a certain program length, the distribution of fitness does not vary with size.
- Since there are more longer programs, the number of long programs of a given fitness is greater than the number of short programs of the same fitness.
- Over time, GP samples longer and longer programs simply because there are more of them.



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## **Modification Point Depth**

- When a genetic operator modifies an individual, the deeper the modification point the smaller the change in fitness.
- Small changes are less likely to be disruptive, so there is a preference for deeper modification points, and consequently a preference for larger trees (removal bias).





in favor of the larger ones.



# Classification Bloat Control Methods

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Bloat control is possible at different levels of the evolutionary process

- **Evaluation** – Parametric Parsimony Pressure, The Tarpeian
- **Selection** – Multi-objective Optimization, Special Tournaments
- **Breeding** – Special Genetic Operators
- **Survival** – Size/Depth Limits, Operator Equalization
- **Others** – Code Editing, Dynamic Fitness



## Comparison of Methods: Experimental Setup

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Four problems and their characteristics:

- **Symbolic Regression** ( $x^4 + x^3 + x^2 + x$ ) – small improvements can always be made through additions to the existing tree.
- **Artificial Ant** (Santa Fe trail) – strong relationship between nodes throughout the tree, where changes in the left part of the tree have a dramatic effect on the operation in the right part of the tree due to execution order.
- **11-Multiplexer** and **5-bit Even Parity** – both have large-sized solutions.
  - Multiplexer is generally the most difficult of the four problems.
  - Parity is among the easier ones.

**Measure of bloat** – the mean tree size of all individuals generated during the course of an experimental run.

Statistical significance was determined by pairwise t-test:

- at 95% confidence when comparing fitnesses,
- at 99.995% confidence when comparing tree sizes.

This setting was chosen in order to make sure that the mean tree size is much smaller to be considered significantly better (while the fitness did not deteriorate).

# The Tarpeian Method

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**Idea:** Some individuals with above-average size are made uncompetitive by assigning some very poor fitness.

**Realization:**

1. Before the evaluation process, new individuals with above-average size are assigned a very bad fitness with probability  $W$ .
2. These individuals are not evaluated further.
3. Evaluate remaining individuals.
4. Use tournament selection to select individuals that will take part in breeding.

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**Characteristics:**

- The individuals that are marked uncompetitive are not evaluated – this reduces the number of fitness evaluations calculated.

**More samples of the solution space can be tried!**

- The method is overly aggressive – the big individuals are rejected by size before considering their fitness even though the uncompetitive individuals still have a very small chance of being selected thanks to the tournament selection.



- The method is very sensitive to parameter  $W$ .  
If it is high, the method will tend to reject an individual no matter how fit it is.
- The setting  $W = 0.3$  was consistently good across all four problems.

## Lexicographic Parsimony Pressure Method

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**Idea:** Two objectives with fixed priorities assigned are used in the selection procedure

- fitness – a primary objective,
- tree size – a secondary objective.

**Realization:** Uses a modified tournament selection rule of the form

- A)** An individual is considered superior to another if it is better in fitness.
- B)** If they have the same fitness, then an individual is considered superior if it is smaller.
- C)** If they have the same fitness and they are of the same size, the superior individual is determined at random.



## Lexicographic Parsimony Pressure Method: Direct Bucketing

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**Realization:** The number of buckets,  $b$ , is specified beforehand, and each is assigned a quality rank from 1 to  $b$  (the bucket with rank 1 contains the worst-fit individuals).

1. The population of size  $p$  is sorted by fitness.
2. The bottom  $\lceil p/b \rceil$  individuals are placed in the worst bucket.

All individuals remaining in the population with the same fitness as the best individual in the bucket are placed in the bucket as well.

This is to guarantee that all individuals of the same fitness fall into the same bucket (they have the same rank).

3. The same procedure is used to fill in the second worst bucket, the third one etc.  
This continues until there are no individuals in the population.
4. The fitness of each individual is set to the rank assigned to the bucket holding it.

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## Characteristics:

- It has the effect of trading off fitness differences for size.
- The larger the bucket, the stronger the emphasis on size as a secondary objective.
- The topmost bucket with the best-fit individuals can hold fewer than  $\lceil p/b \rceil$  individuals.

## Lexicographic Parsimony Pressure Method: Ratio Bucketing

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**Realization:** The buckets are proportioned, so that low-fitness individuals are placed into larger buckets than high-fitness individuals. A parameter of the method is the bucket ratio  $1/r$ .

1. The population of size  $p$  is sorted by fitness.
2. The bottom  $\lceil 1/r \rceil$  fraction of individuals are placed into the worst bucket.  
All individuals remaining in the population with the same fitness as the best individual in the bucket are placed in the bucket as well.
3. The same procedure is used to fill in the second worst bucket with the bottom  $\lceil 1/r \rceil$  fraction of the remaining population, etc.  
This continues until every individual of the population has been placed in a bucket.
4. The fitness of each individual is set to the rank assigned to the bucket holding it.



# Lexicographic Parsimony Pressure Method: Performance

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## Plain Lexicographic Parsimony Pressure

- Successful on all problems but the symbolic regression.

The symbolic regression is unusual in that occurrence of individuals of exactly the same fitness in the population is rare since small changes in fitness can be achieved by adding code fragments to the bottom of trees.

## Direct bucketing

- Successful on all four problem, but no single setting of the parameter  $b$  that would be consistently good across all problems was found.

$b \in \{25, 50, 100\}$  is good for the symbolic regression.

$b = 250$  is the common setting for other problems.

## Ratio bucketing

- Nearly uniformly superior in any setting, considering  $r = \frac{1}{2}, \frac{1}{3}, \frac{1}{4}, \frac{1}{5}, \frac{1}{6}, \frac{1}{7}, \frac{1}{8}, \frac{1}{9}, \frac{1}{10}$ .
- Some irregularities in the domain of good values for the multiplexer problem were observed.



## Linear Parametric Parsimony Method

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**Idea:** Parsimony pressure methods consider **size as part of the selection process** – a fitness of the program is a function of its quality and size. A fitness of a program is decreased by an amount that depends on its size. The intensity with which bloat is controlled is determined by a parameter called *parsimony coefficient*.

- If it is too small then the control of bloat is not effective.
- If it is too large then the minimization of tree size will become a primary target and fitness will be ignored.

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### **Realization:**

- Linear Parametric Parsimony Method treats the individual's size as a linear factor in fitness

$$g = x \cdot f + y \cdot s$$

where the parameters  $x$  and  $y$  weight contributions of raw fitness  $f$  and the size  $s$  to the final fitness  $g$ .

- Linear Parametric Parsimony Method with a limit applies the size component only if  $s$  greater than some specified limit  $z$ . Then

$$g = xf, \text{ if } s \leq z$$
$$g = x \cdot f + y \cdot (s - z), \text{ otherwise.}$$









# Double Tournament Method

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**Idea:** Selection procedure applies two layers of tournaments in series

- qualifier tournaments and
- final tournaments.

## **Realization:**

1. Tournament contestants are chosen as winners of qualifier tournaments.
2. Winner chosen in the qualifier tournaments compete in the final tournament.  
Fitness objective can be used in qualifier tournaments and tree size objective in final tournaments or vice versa.

The selection is parameterized by

- fitness tournament size  $F$ ,
- parsimony tournament size  $D$ ,
- *do-fitness-first* – indicates whether fitness tournaments are used in the qualifiers or final tournaments.

## Double Tournament Method

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### Characteristics:

- $D$  should be smaller than 2, otherwise it puts too much pressure on parsimony.  
In order to permit  $D$  to hold real values between 1.0 and 2.0 the following rule was implemented: Given two individuals, the smaller one wins the tournament with probability  $D/2$ , else the larger one wins.  
Thus,  $D = 1$  is random selection, while  $D = 2$  is the same as a plain parsimony-based tournament of size 2.
- An individual passes the double tournament if it is generally low in size and high in fitness.
- $D = 1.4$  was consistently superior on all four problems.









## Death by Size Method

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**Idea:** At each step of an iterative process some individuals are selected to breed children, while other individuals are selected to die and be replaced by the new children.

**Realization:** Steady-state generational model, where at each generation selection of the parents and replacements is done as follows:

- Selection of the parents to breed is based on the fitness.

Tournament selection with size  $S$ .

- Selection of the individuals to die is based on tree size (larger programs are more likely to die).  
Tournament selection with size  $K$ . This value should be very moderate ranging between 1.0 and 2.0.

This is implemented so that given two competing individuals, the larger one wins with probability  $K/2$ , else the smaller one wins.

### Characteristics:

- Under no combination of settings the method was superior on the 11-bit Multiplexer problem. Only one setting ( $S = 7$ ,  $K = 2.0$ ) was successful on the 5-bit Even Parity problem.
- As  $K$  increases, larger individuals are selected to die, resulting in stronger bloat control.
- Steady-state evolution model tends to apply a strong selection pressure (large trees are produced very rapidly).

## Comparison of Methods: Experiment Setup

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**General conclusion:** The combination of a method with depth limiting was nearly universally superior to either the method alone or depth limiting alone.

Double Tournament and Biased Multi-objective appeared the best across all problem domains when tuned to their optimal per-problem values.

**Question:** Which method with its single problem-independent setting is the best across all four problems?

# Comparison of Methods: Experiment Setup

Highlighted in blue are methods that have one or more settings with which they are superior to plain depth limiting across all problems.

- shown are the best settings with the lowest mean tree size of run averaged over all four problem domains.

Method	Settings
Double Tournament	$D=1.4, do\text{-}fitness\text{-}first=false$
Proportional Tournament	$P=0.2$
Lexicographic with Direct Bucketing	(not contending)
Lexicographic with Ratio Bucketing	$R=1/2$
Plain Lexicographic	(not contending)
Linear Parametric	$X=32$
Biased Multi-objective	$F=0.95$
The Waiting Room	(not contending)
Death by Size	(not contending)
Tarpeian	$W=0.3$

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# Parsimony Pressure Method

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## Parsimony pressure method

- It is a penalty method that decreases the fitness of programs by an amount that depends on their size. Typically, the penalty is simply proportional to program size.

$$f_p(x) = f(x) - c \cdot l(x)$$

where  $f_p(x)$  is the new fitness,  $f(x)$  is the raw fitness,  $c$  is the **parsimony coefficient** and  $l(x)$  is the size of program  $x$ .

- The value of the parsimony coefficient is typically set once and for the whole run.

However the correct values of this coefficient are highly dependent on

- the problem being solved,
- the choice of functions and terminals,
- and others control parameter settings.

Since very little theory is available to aid in setting the parsimony coefficient, users are forced to proceed by trial and error.

- Yet, it is the simplest and the most frequently used bloat control method.



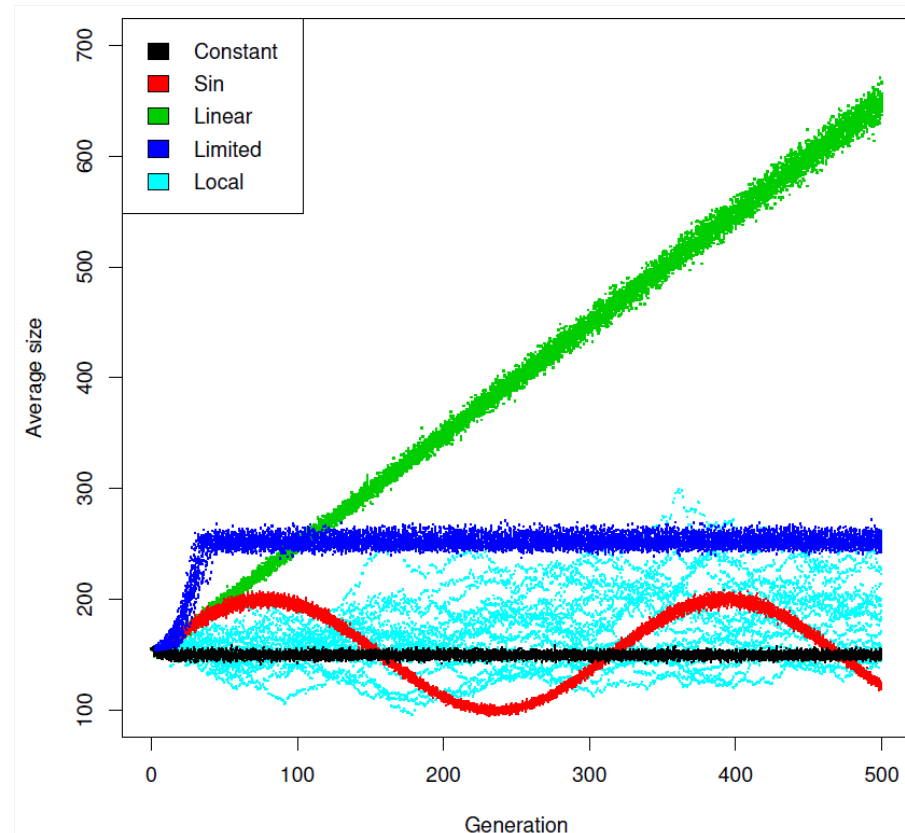


# Covariant Parsimony Pressure Method

Application of the method to the 6-Multiplexor problem.

Five targets for the average program size:

- Constant
- Sin
- Linear
- Limited
- Local



## Reading

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- [Poli08] Poli, R., Langdon, W., McPhee, N.F.: *A Field Guide to Genetic Programming*, 2008.
- [Luke06] Luke, S. and Panait, L.: A Comparison of Bloat Control Methods for Genetic Programming. *Evolutionary Computation*, Volume 14 Issue 3, 2006.  
<http://portal.acm.org/citation.cfm?id=1182892.1182897>

