University of St. Thomas

Graduate Programs in Software

SEIS 610: Software Engineering Spring 2014

Genetic Programming Final Report

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Maryam Alruwaythi Jed Brundidge Tabinda Hasan Kelly Heitz Dawn Young

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Preface

All of the documents referenced here, and historical documentation of the life of this project from beginning to end, are accessible as Google docs in *GP group project* folder. Anyone with this link has access: https://drive.google.com/folderview?id=0BxP-i4iTYfT5LWtKcDFOU1VvbDQ&usp=sharing

Our final genetic programming system was a modification of package called tiny_gp we found online, with guidance from a source called "A Field Guide to Genetic Programming" by Riccardo Poli, William B. Langdon, and Nicholas F. McPhee, copyright March 2008.

Link to the original tiny_gp: http://cswww.essex.ac.uk/staff/rpoli/TinyGP/

Project Summary

Problem Definition

Create a genetic programming application that will automatically evolve based on experiences the system "learns" by evaluating fitness of potential solutions against an established set of training data. The evolution will occur through the use of crossover and mutation methodologies applied probabilistically to the initial and subsequent generations of potential solutions, with the most "fit" solutions having the highest probability of representation in subsequent generations. Specifically, the program will randomly generate an initial set of functions and then repeatedly measure, modify, and regenerate functions until one that is equivalent (or very close) to the target function f(x) = (2x2+4)/3 is achieved within a 15-minute time frame.

The project shall be cooperatively managed, created, and tested by the team, presented on May 12th, and submitted with all supporting documentation and post-project analysis on May 19th, 2014.

Project Requirements

The following requirements include mark-ups to demonstrate how our view the project changed over time. All but one of the requirement changes were based on decisions our group made as we acquired more knowledge and experience along the way. The single requirement change from the customer, Professor Lai, was the target function in requirement 8c.

- 1. Input Expectations
 - a. Training data file (target function implied)

b. Function set

c. Terminal set

d. Population size of each generation

i. 20 100,000

- 2. Binary Tree requirements
 - a. Each Operator node (functional node) must have 2 child nodes
 - b. Each Operand (terminal node) must have 0 child nodes
- 3. Generating the initial population, generation 1, G1
 - a. Use "Full Method" "Grow Method" to randomly generate 20 100,000 functions
 - b. Operators will be randomly selected from the Function Set
 - c. Operands will be randomly selected from the Terminal Set.
 - d. Depth of the initial binary trees will be 43
 - i. Each node in level 1, 2, and 3 must have 2 child nodes.
 - ii. Each node in level 43 must have 0 child nodes

4. Fitness

- a. Calculate the fitness of each function (f1 through f20) in the population in each generation (G1, G2, etc.).
- b. For each function in the population (f1 through f20), calculate its fitness as follows:
 - i. Apply the function to each x in the Training Data Set (x1 through x20) and retain the result as G1f1x1, G1f1x2, etc
 - ii. Subtract the function's result for each x from that x's result in the training set to find the difference, d:
 - 1. For example, referring to the training set function as f0 (the target function):

G1f1x1 - f0x1 = dG1f1x1 (this is the difference d for x1 using function 1 in generation 1)

G1f1x2 - f0x2 = dG1f1x2

G1f1x3 - f0x3 = dG1f1x3

- 2. Note that f0x1 through f0x20 will always be the same in every calculation in every generation because f0 represents the target function applied to each x in the Training Data Set.
- iii. Take the absolute value of every difference
- iv. Fitness of a function is the sum of the absolute values of the differences.
- 1. For example, the fitness of function 1 in generation 1 is calculated as follows: $FG1f1 = |dG1f1x1| + |dG1f1x2| + |dG1f1x3| \dots |dG1f1x20|$
- c. Rank the 20 functions by their fitness:
 - i. Rank 1 will have the lowest fitness value
 - ii. Rank 20 will have the highest fitness value
- d. For each generation, sum the fitness values of the top 5 ranked functions and store the sum as T5G1, T5G2, etc. (This value will be used later to compare the fitness of generations over time.)
- 5. Varying the population from generation to generation
 - a. Start with 20 100,000 functions from the preceding generation (they should be ranked in order of their fitness to the training set).
 - b. Eliminate 10% of the lowest ranked functions, i.e. eliminate functions ranked 19 and 20.
 - c. Reproduce (clone) 10% of the highest ranked functions, i.e. reproduce functions ranked 1 and 2.
 - i. 2 of the 20 functions will be the most fit function from the prior generation
 - ii. 2 of the 20 will be the 2nd most fit function from the prior generation
 - iii. The 16 other functions will be the remaining functions from the prior generation (excluding the 2 least fit functions).
 - d. Use tournament selection.

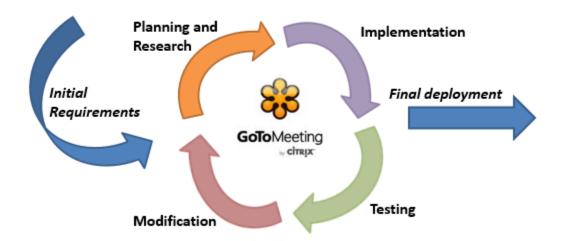
- i. Tournament size = 4
- ii. Negative tournaments are used for the selection of the individuals to be replaced at each steady-state-GP iteration.
- e. Crossover:
 - i. Randomly select 2 functions from list of 20 functions
 - ii. From the higher ranked function, randomly select a node that will be crossed over.
 - iii. In the lower ranked function, randomly select a node that will be replaced.
 - iv. Replace the node in the lower ranked function with the node from the higher ranked function.
 - v. Repeat 10 times
 - vi. The fitness rank of each function will remain static in spite of crossover changes occurring in the current generation.
 - vii. Subtree crossover is used. The selection of crossover points is uniform, so every node is chosen equally likely.
 - e. Mutation
 - i. Apply mutation to the new set of 20 functions with probability of 0.01 (assuming 1 function in 5 generations)
 - ii. Set an indicator to mark "yes" or "no" to each function, indicating whether it was selected for mutation or not.
 - iii. Randomly select any node, a functional node or a terminal node
 - 1. Whether the selected node is functional or terminal, initialize a new sub-tree with a depth of 3 using "Full" method.
 - 2. Apply point mutation to the selected node: i.e., if a functional node is selected randomly replace it with a different functional node, and vice versa.
- 6. Evaluate the general population after each 20th generation
 - a. If T5G20 > T5G10, eliminate lowest 50% fit functions
 - i.—Use strategy we used to generate the initial population to add 10 new functions to the population set
 - ii. At 20th, 40th, 60th generations, etc. the sum of the current top 5 fitness functions should be less than the sum of the top 5 fitness factors from 10 generations prior.
- 7. Output Expectations
 - a. During development phase,
 - i. print after each generation the population of 20 functions that have been generated
 - ii. Rank them by fitness
 - iii. Which functions were eliminated?
 - iv. Which functions were reproduced?
 - v. Which functions were mutated?
 - b. After programming is finalized, print for each generation:
 - i. Print best function
 - 1. Which generation produced it
 - 2. Its fitness
 - 3. The average fitness of the final generation
 - 4. Average function length of the generation

- 5. Length of time the program ran
- 6. A fitness graph comparing the best function against the target function
- 8. Acceptance/Termination criteria
 - a. Terminate program when fitness of any function in the population set = 0 the best function is <= .0009, or
 - b. After 15 minute limit
 - i. Select most fit function
 - c. Function should be equivalent to: $f(x) = (x^2 1)/2 f(x) = (2x^2 + 4)/3$
 - d. Program will run without errors or crashing.

Development Process

Development Process

Detailed initial requirements \rightarrow generalized and modified in response to increased knowledge



Project Implementation

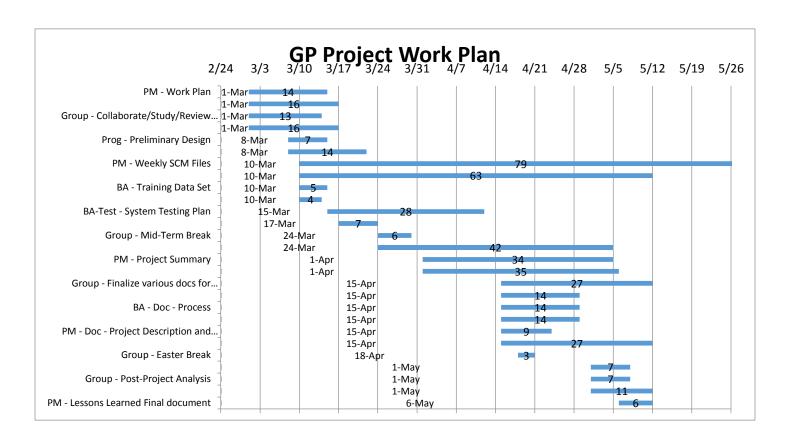
Team Roles

Member	Starting Role	Mid-term Role	End of Project Role
Maryam	Analyze/Develop Reqmts	Analyze/Develop Reqmts	Research Package Code
	Test Code	Analyze Packages	Measure Code
Jed	Develop Code	Analyze/Develop Reqmts	Analyze Package Code
		Develop Code	Develop Test Plan
			Test Code
Tabinda	Analyze/Develop Reqmts	Analyze/Develop Reqmts	Analyze Package Code
	Co-Manage Project	Analyze Packages	Measure Code
Kelly	Develop Code	Analyze/Develop Reqmts	Modify Package Code
		Create UML diagram	Modify Requirements
		Develop Code	Modify UML diagram
			Black box testing
Dawn	Analyze/Develop Reqmts	Analyze/Develop Reqmts	Modify Package Code
	Co-Manage Project	Develop Code	Compile Presentation
		Admin Meetings	Compile Final Report
			Manage Project

Timeline

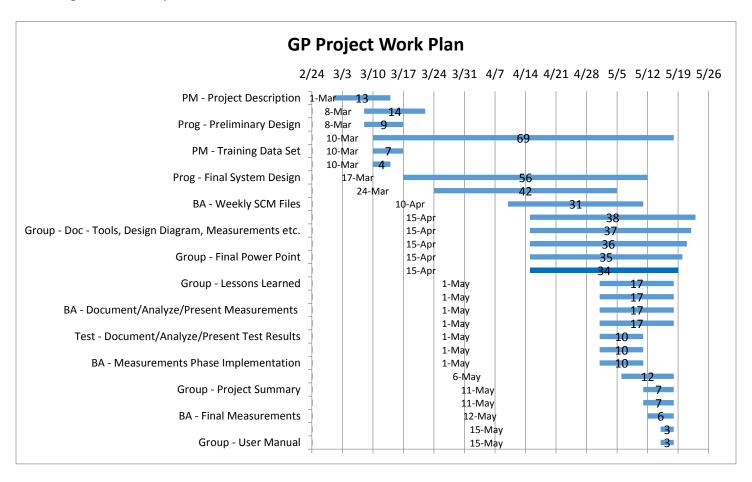
Initial Timeline:

We started the project with the following timeline on when we expected to complete tasks assigned to the group members. Initially we were under the impression that we knew exactly how much time we were expected to (and the amount of time we should/would spend on the assigned tasks). During the course of learning more about the project and the expectations from the Genetic Programming Application we were developing, many of the aforementioned tasks changed, as did the amount we spent on them as a group. We had allocated a large amount of time for meeting and dissecting the requirements & design, and then later for evaluating the program we had started to build.



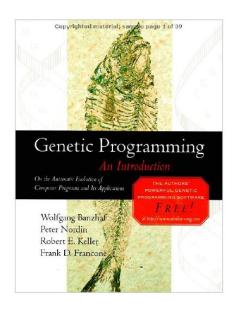
Actual Timeline:

As expected we spent a large portion of our time in finalizing the requirements, design, then developing the program and pulling together all of the documentation and visual materials together. We did eventually decide to go with a package and a lot of the final couple of weeks were spent on getting the code of this new package to return the results we expected from our GP Application, in addition to measuring, testing and then accumulating the visual materials and documentation to support our finished product (i.e. our GP Application). It would have been ideal to spend a lot more time on testing and measuring this new code but with very few requirements and no team experience with packages, the search for an appropriate package for modification took longer than anticipated.



Tools and Resources









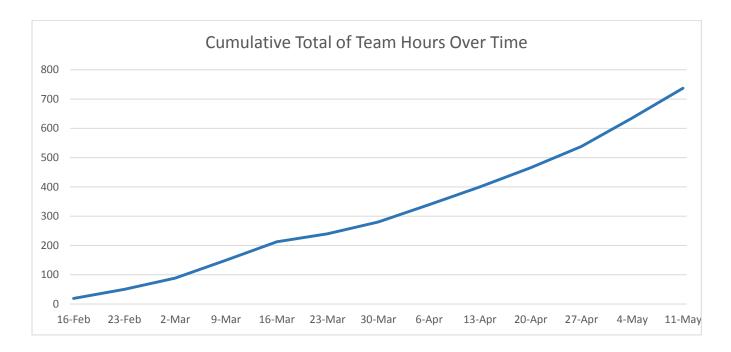


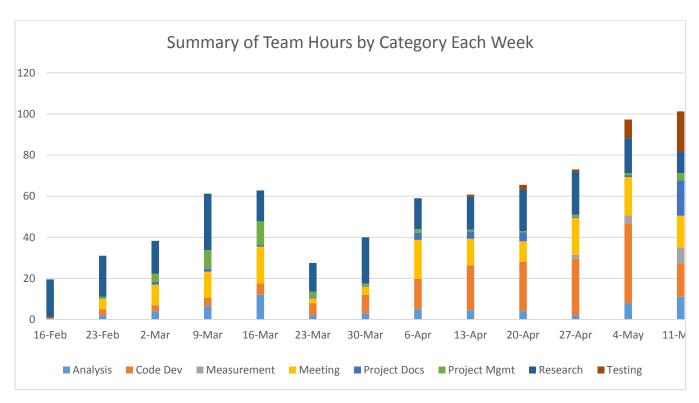


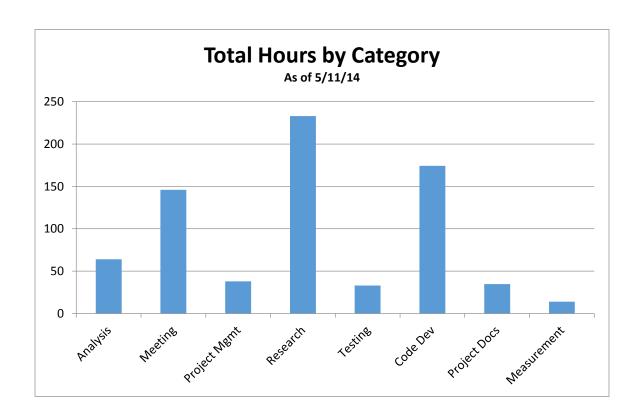


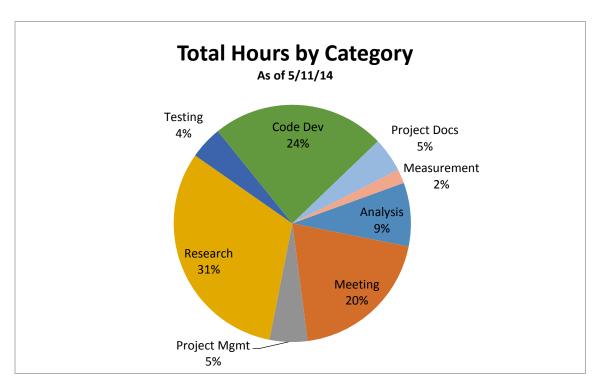
A FIELD GUIDE TO
GENETIC PROGRAMMING
WEB SITE IN SUPPORT OF A FREELY DOWNLOADABLE, CREATIVE COMMONS LICENSED INTRODUCTION TO GENETIC PROGRAMMING

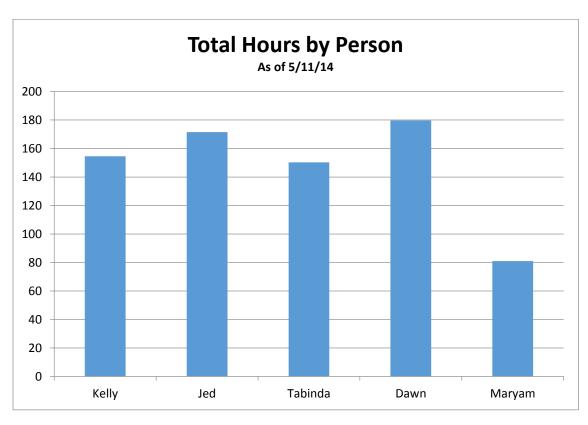
Team Activity Measurements

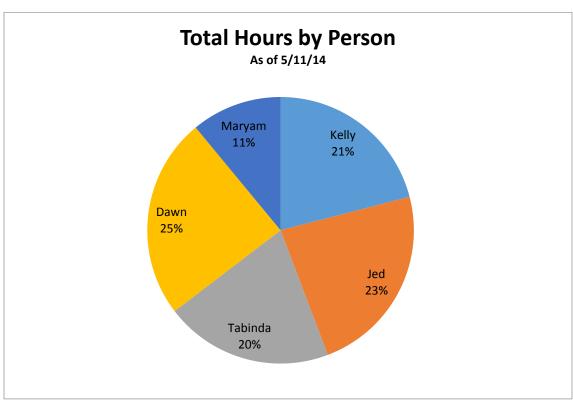


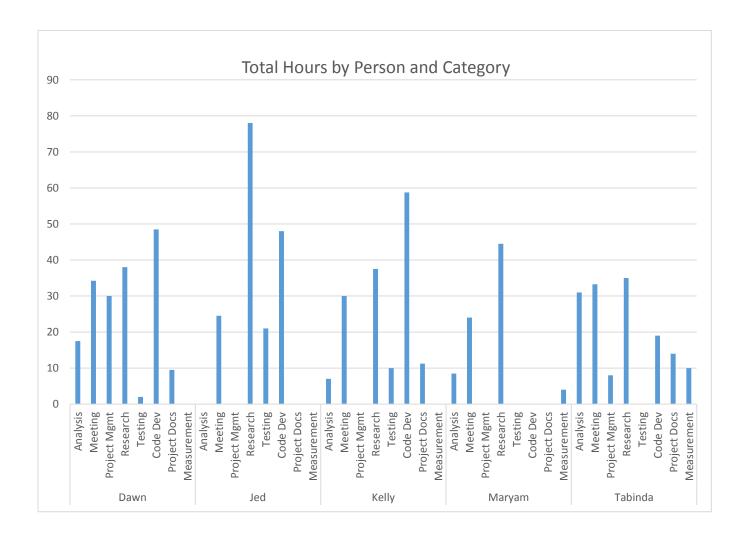












Final System Design

Selecting Training Data Set

We decided our training data set would be comprised of 20 elements. When selecting the range of the "x" values, we considered two options:

Option A: $-5 \le x \le 5$ Option B: $-20 \le x \le 20$

We created a variety of test functions to measure fitness against our proposed data sets, e.g. a constant function, a quadratic function (with x^2 in the numerator), a function with x in the denominator. The resulting fitness values are shown below.

Because of the little variation we saw in our experimental fitness values with Option A, we chose to go with Option B, a range of x values between -20 and +20.





<u>Settings</u>

Initial Operands 0, 1, 2, 3, X

(Used limited operands while developing initial code.)

Final Operands 0, 1, 2, 3, 4, 5, 6, 7, 8, 9, X

(Found it easy to expand set of operands in final code.)

Operators +, -, *, /

Max Length 5

Tree Depth 3

Population Size 100,000

Tournament Size 4

Crossover Probability 0.90

Mutation Probability 0.05

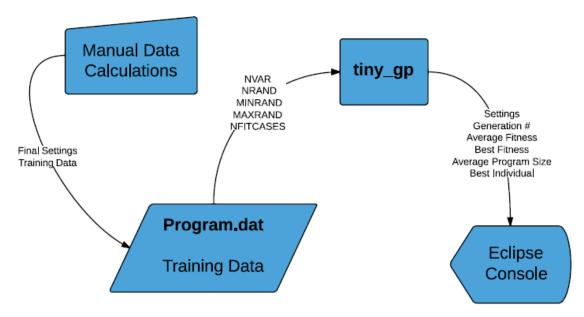
Stop Program Criteria: Fitness <= 0.001

15 minutes

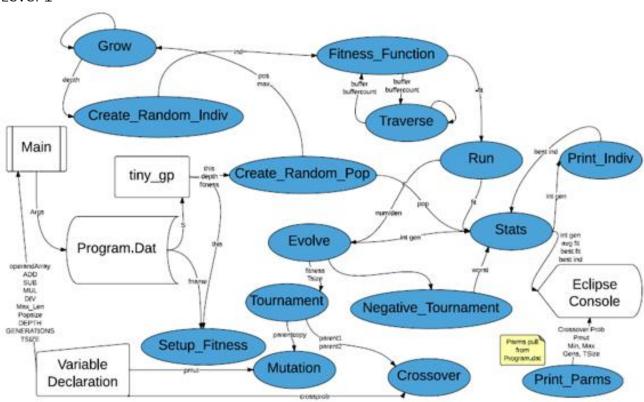
100 generations

Data Flow Diagram

DFD Level 0

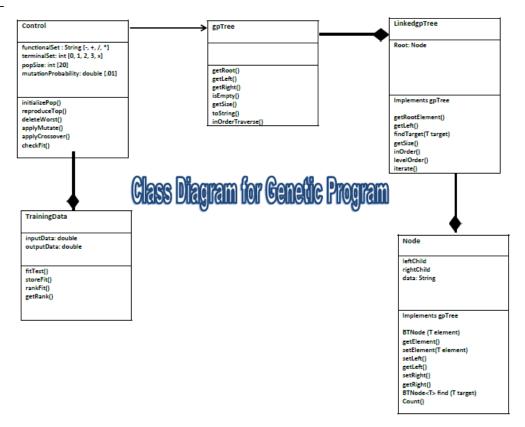


DFD Level 1

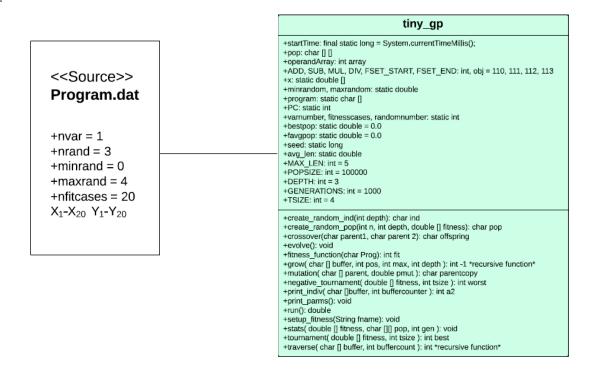


UML

Original UML



Final UML



Division by Zero

Code line to handle a fraction with denominator close to 0:

```
if ( Math.abs( den ) <= 0.001 ) return( num );
else return( num / den );</pre>
```

Translation:

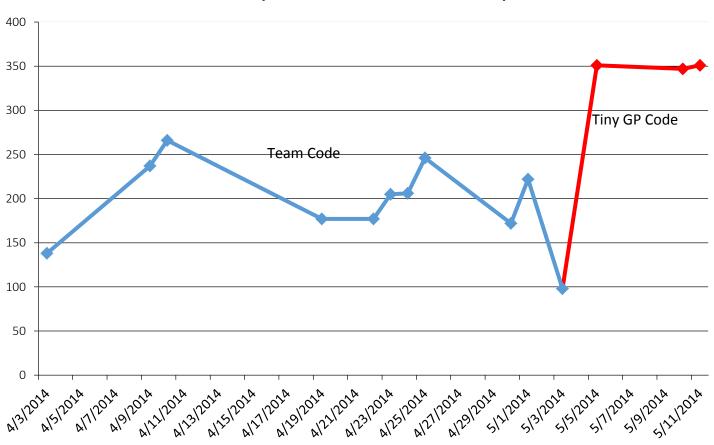
If the denominator of a fraction is very small, (close to zero or equal to zero), return the numerator.

This is the equivalent of dividing the numerator by 1.

Measurements

Lines of Code (LOC)

LOC (based on all SCMs)

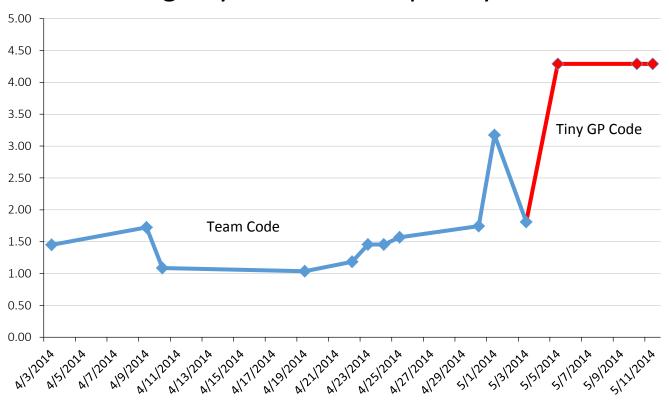


Cyclomatic Complexity

Average Cyclomatic Complexity - Overall

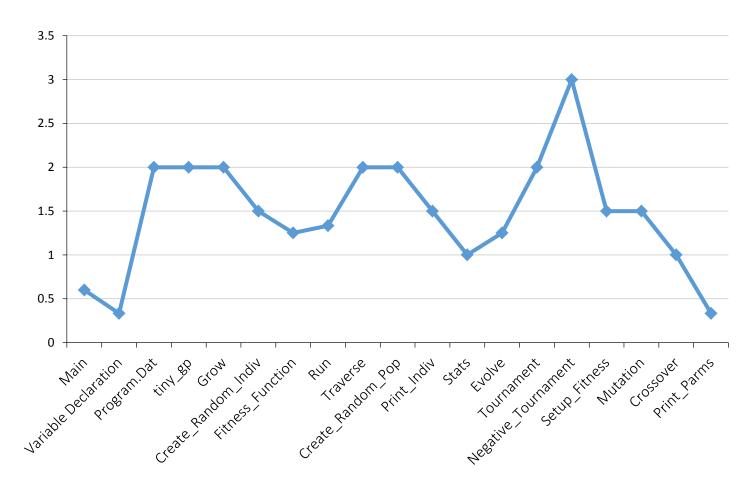
Date:	4/3	4/9	4/10	4/19	4/22	4/23	4/24	4/25	4/30	5/1	5/3	5/5	5/10	5/11
Avg CC:	1.45	1.73	1.09	1.04	1.19	1.46	1.46	1.57	1.74	3.17	1.81	4.29	4.29	4.29

Average Cyclomatic Complexity - Overall



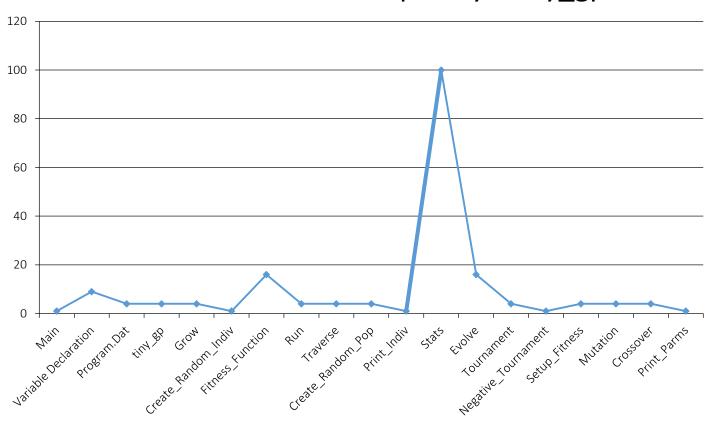
Cohesion

Cohesion – tiny_gp



Information Flow Complexity (IFC)

Information Flow Complexity - tiny_gp



Relative System Complexity (RSC)

Methods	Variables	FI [FanIn]	FO [FanOut]	Information Flow Complexity [IFC]	Functionality Complexity of each Method [H]	
Main	5	1	1	1	1.67	0.6
Variable Declaration	12	-	3	9	3	0.33
Program.Dat	2	1	2	4	0.5	2
tiny_gp	2	1	2	4	0.5	2
Grow	2	2	1	4	0.5	2
Create_Random_Indiv	2	1	1	1	0.67	1.5
Fitness_Function	4	2	2	16	0.8	1.25
Run	3	1	2	4	0.75	1.33
Traverse	2	2	1	4	0.5	2
Create_Random_Pop	2	1	2	4	0.5	2
Print_Indiv	2	1	1	1	0.67	1.5
Stats	8	5	2	100	1	1
Evolve	4	2	2	16	0.8	1.25
Tournament	2	1	2	4	0.5	2
Negative_Tournament	1	1	1	1	0.33	3
Setup_Fitness	2	2	-	4	0.67	1.5
Mutation	2	2	-	4	0.67	1.5
Crossover	3	2	-	4	1	1
Print_Parms	6	-	1	1	3	0.33
				186	18.02	

Coupling, U (Sum of IFCs)	186.00
Total Functionality Complexity, H	18.02
Total System Complexity, C	204.02
Relative System Complexity, RSC	10.74

RSC <= 26 is good!

Additional Measurements

In addition to the charts and graphs provided above, there is much more measurement and analysis located in our Google docs **GP group project > Measurement > Measurements**. Tabinda performed measurement analytics on every class on each day that code was uploaded. Link:

https://drive.google.com/?usp=folder&authuser=0#folders/0BzpM7gg1JvssN0JudFFEc24xSUU

Testing

Summary

PART 1

The initial testing for our system revolved around 8 test cases written and executed by Jed. These test cases manipulated input data in the system to test how the program would react when different inputs were entered. Most of the test cases proved the flexibility of our program in that the program continued to function even with the changes. These test cases proved themselves valuable in allowing us to find which inputs caused the system to react in certain ways. One of the major areas of testing involved finding a function with a fitness value of 0 within our 15 minute time frame. Through our tests we believe we arrived at a synergistic balance which allows our system the perfect opportunity to find the best fitness value within the allotted time.

PART 2

The second aspect to our testing was the code coverage, also conducted by Jed. For this portion of the testing we used CodeCover, which is an open source tool that plugs into Eclipse and generates reports as you run your program. This tool was very easy to use. With a few simple steps Jed was able to get it running and generating reports for our system.

PART 3

The third aspect of our testing was Black box testing was used to confirm the program's ability to find the correct function (or an equivalent expression) in under 15 minutes. This testing, conducted by Kelly, was carried out by running the program with a random seed 30 times. Statistical data from each run was collected from the Eclipse console, and listed by category in an excel spreadsheet for further analysis. The results of this test are summarized by the following statistics:

- Correct function was found 30/30 times: a 100% success rate for this test sample.
- Minimum number of generations to solution: 2
- Maximum number of generations to solution: 14
- Average number of generations to solution: 5.1
- Fastest time to solution (seconds): 5.357
- Slowest time to solution: 23.275
- Average time to solution: 7.8726
- Average length of functions in final generation population: 13.336

Note: Test results were carried out on a Toshiba Satellite E45t-A4300 laptop

Part 1 – 8 Specific Tests

Test Case 1

Test Case ID: TC_01_Correct Training Set_5-10-14 Test Designed by: Jed Brundidge

Test Priority (Low/Medium/High): Low Test Designed date: 5-5-2014

Module Name: Test Executed by: Jed Brundidge

Test Title: Test Training data

Test Execution date: 5-10-2014

Description: Test output with correct training data

Pre-conditions: All inputs have been defined before system build

Dependencies: System data structure

Step	Test Steps	Test Data	Expected Result	Actual Result	Status (Pass/Fail)	Notes
1	Input training set	• • •		Complete Fitness test		System acted as expected with correct inputs

Post-conditions:

The system has made a successful run. When the system is run successfully we will achieve 0 for our fitness.

Test Case 2

Test Case ID: TC_02_Double Test_5-10-14 Test Designed by: Jed Brundidge

Test Priority (Low/Medium/High): Low Test Designed date: 5-5-2014

Module Name: Test Executed by: Jed Brundidge

Test Title: Verify only type double is accepted for

training input

Test Execution date: 5-10-2014

Description: Test that exception is thrown when no integer

Pre-conditions: Define training input.

Dependencies:

Step	Test Steps	Test Data	Expected Result	Actual Result	Status (Pass/Fail)	Notes
	Input type Integer		System should throw exception	Exception		When integer is entered an error occurs
2	Input type Char		System should throw exception	Exception		When char is entered an error occurs
3		"Hello"	System Should throw exception	Exception		When a string is entered an error occurs

Post-conditions:

None.

Test Case 3

Test Case ID: TC_03_ChangeTreeDepth_5-10-14 Test Designed by: Jed Brundidge

Test Priority (Low/Medium/High): Med Test Designed date: 5-5-2014

Module Name: Test Executed by: Jed Brundidge

Test Title: Change tree depth Test Execution date: 5-10-2014

Description: Test flexibility of tree depth. Change

from 5

Pre-conditions: Tree class built to specs

Dependencies:

Step	Test Steps	Test Data	Expected Result	Actual Result	Status (Pass/Fail)	Notes
		Set depth to 15	-	Expected output received. System took longer to get the answer as expected		Larger depth means the trees are bigger.
2	Decrease tree depth	Set depth to 3		Expected output received. System took shorter to get the answer as expected.		Shorter depth means tree is smaller.
3			Extremely long run time	Did not finish. Run time took much longer than expected.	Pass	Very long to run with this depth
4	Increase small	•	time	Expected output received. System took a very short time to get answer as expected.	Pass	Very short.

Post-conditions:

Run time for each test went as expected. When the depth of the tree was decreased the program became less complex and took less time to find a solution. When the tree depth was increased the program became more complex making the run time longer.

Test Case 4

Test Case ID: TC_4_Tree pop test_5-10-14

Test Designed by: Jed Brundidge

Test Priority (Low/Medium/High): Med Test Designed date: 5-5-2014

Module Name: Test Executed by: Jed Brundidge

Test Title: Verify that trees have been created. Test Execution date: 5-10-2014

Description: Test population array.

Pre-conditions: All size requirements must be in place.

Dependencies:

Step	Test Steps	Expected Result		Status (Pass/Fail)	Notes
	Create empty array list	1	Populated array list		Trees are being populated correctly

Test Case 5

Test Case ID: TC_5_Change Target Function_5-10-14 Test Designed by: Jed Brundidge

Test Priority (Low/Medium/High): Med Test Designed date: 5-5-2014

Module Name: Test Executed by: Jed Brundidge

Test Title: Alter target function Test Execution date: 5-10-2014

Description: Determine program flexibility by altering the

target function.

Pre-conditions: Program works successfully with initial target function.

Dependencies:

Step	Test Steps	Test Data	Expected Result	Metual Recult	Status (Pass/Fail)	Notes
1	S	Change operators in target function		System found the closest match in under 15 minutes		System passed flexibility test

Test Case 6

Test Case ID: TC_6_Population size_5-10-14 Test Designed by: Jed Brundidge

Test Priority (Low/Medium/High): Med Test Designed date: 5-5-2014

Module Name: Test Executed by: Jed Brundidge

Test Title: Population test. Test Execution date: 5-10-2014

Description: Change population from 1000 to find

best run time.

Pre-conditions: This test is to find out how the system will react when the population size is either increased

or decreased.

Dependencies: Population array.

Step	Test Steps		Expected Result	Actual Result	Status (Pass/Fail)	Notes
	Increase pop to 100000	"		Problem solved quickly		Larger population is increased time
	Decrease pop to 5000	_		Problem solved slowly	Pass	

Test Case 7

Test Case ID: TC_7_15 min run_5-10-14 Test Designed by: Jed Brundidge

Test Priority (Low/Medium/High): Med Test Designed date: 5-10-2014

Module Name: Test Executed by: Jed Brundidge

Test Title: Length of run

Test Execution date: 5-10-2014

Description: Change run parameters

Pre-conditions: We need a fully functional program to test the time of each run when parameters are changed.

Dependencies: Max height of tree, Pop size

Step	Test Steps	Test Data	Expected Result	Actual Result	Status (Pass/Fail)	Notes
	height		Run will be longer than 15 min	Long run	Pass	
	height		Run will be very short	Short run	Pass	

Post-conditions: We can now see that the system is very flexible in allowing us to manipulate the max height of the trees to figure out how long our runs will take and allow us to find a sweet spot to find our target in less than 15 minutes.

Test Case 8

Test Case ID: TC_8_Mutation Test test_5-5-2014 Test Designed by: Jed Brundidge

Test Priority (Low/Medium/High): Med Test Designed date: 5-10-2014

Module Name: Test Executed by: Jed Brundidge

Test Title: Test functionality of mutation Test Execution date: 5-10-2014

Description: Swap trees for mutation

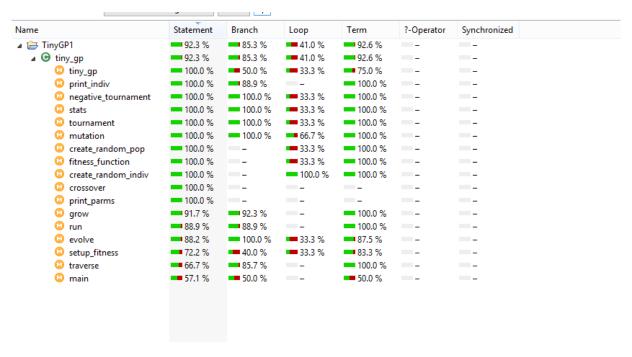
Pre-conditions: Populate trees

Dependencies: Tree class and Population

Step	Test Steps	Test Data	•		Status (Pass/Fail)	Notes
	tree	First tree 3 + 2 / Second tree 1 + x +	·	Mutation was successful	Pass	

Part 2 – Code Coverage

The following display is from CodeCover application, used to test our tiny_GP program.



This diagram demonstrates how CodeCover displayed the testing coverage of our code, line by line, in the Eclipse console. Green indicates full coverage and red indicates no coverage. The full picture is available on Google Docs: **GP group project > Testing > Testing 700pm 5-18-14**. Link:

https://drive.google.com/folderview?id=0B-sYVFgNOuZONkVLbzBuOXY0MGs&usp=sharing&tid=0BxP-i4iTYfT5LWtKcDFOU1VvbDQ

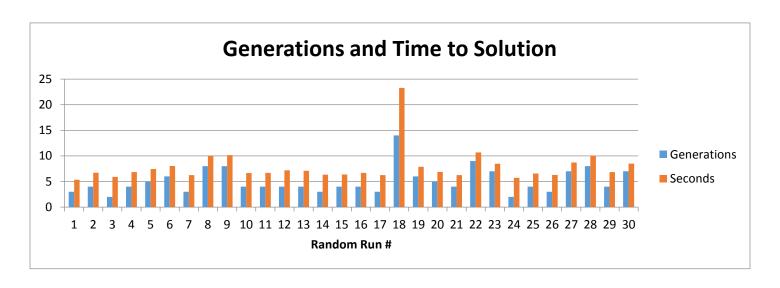
```
static double [][] targets;
public class tiny_gp {
                                                                double run() { /* Interpreter */
  static double [] fitness;
                                                                 char primitive = program[PC++
if ( primitive < FSET_START )
   return(x[primitive]);</pre>
                                                                 switch ( primitive ) {
  static final int
                                                                    case SUB : return( run() - run()
case MUL : return( run() * run()
    SUB = 111.
                                                                     //handle divide by zero
    DIV = 113.
    FSET END = DIV:
  static double minrandom, maxrandom;
  static char [] program;
  static int varnumber, fitnesscases, randomnumber;
                                                                         ( 0.0 ); // should never get here
  static long seed;
static double avg len;
                                                         int traverse( char [] buffer, int buffercount ) {
                                                                 if ( buffer[buffercount] <
  return( ++buffercount );</pre>
    GENERATIONS = 100
                                                                 switch(buffer[buffercount]) {
                                                                    case SUB:
                                                                    case MUL:
```

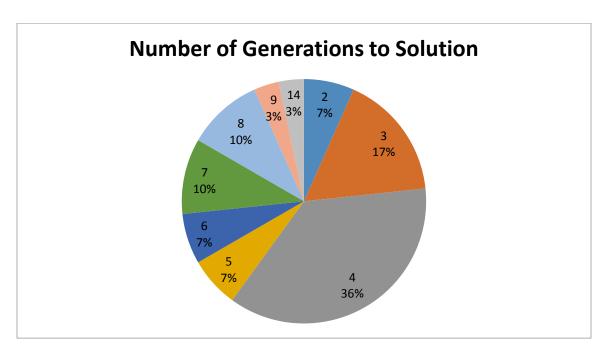
Part 3 – Black Box Testing

Raw data for the following summary is located in Google docs: **GP group project > Testing > 30 Random Runs Output**. Link: https://drive.google.com/folderview?id=0BxP-

i4iTYfT5UUppd19LZjlzcFU&usp=sharing&tid=0BxP-i4iTYfT5LWtKcDFOU1VvbDQ

	# of Gens	Run time	Final Gen	
Run#	to Solution	(seconds)	Avg Size	Best Function
1	3	5.357	5.62092	((2.0 * 2.0) / (6.0 / (2.0 + (X1 * X1))))
2	4	6.719	5.48642	((X1 * X1) + (X1 + ((((4.0 / (X1 / X1)) - (X1 * X1)) / 3.0) - X1)))
3	2	5.912	4.33212	(2.0 * ((2.0 + (X1 * X1)) / 3.0))
4	4	6.848	5.47682	(((2.0 + (X1 * X1)) * 4.0) / 6.0)
5	5	7.452	8.77176	(((((X1 * X1) - 2.0) + 6.0) + (X1 * X1)) / 3.0)
6	6	8.036	12.02732	((((X1 * X1) - 4.0) / ((2.0 + 1.0) / 2.0)) + (4.0 / 1.0))
7	3	6.246	6.11244	((7.0 + (((X1 * X1) - 3.0) + (X1 * X1))) / 3.0)
8	8	10.039	21.53586	((2.0 + (X1 * X1)) - (((X1 * X1) + 2.0) / (8.0 - 5.0)))
9	8	10.155	25.54626	(4.0 / ((2.0 + 4.0) / ((X1 * X1) + 2.0)))
				((X1 * X1) - (((((X1 * X1) - (0.0 + ((1.0 * 0.0) - X1))) - X1) - 4.0) /
10	4	6.668	5.45922	(3.0 / 0.0)))
11	4	6.683	5.82284	((X1 * X1) - (((((X1 + (X1 * X1)) - X1) + (8.0 - 4.0)) - 8.0) / 3.0))
12	4	7.187	5.28024	((X1 * X1) - (((X1 * X1) - 4.0) / 3.0))
13	4	7.107	7.13632	((X1 * X1) - (((X1 * X1) / 3.0) - (8.0 / 6.0)))
14	3	6.33	4.87936	((X1 * X1) + ((4.0 - (1.0 * (X1 * X1))) / 3.0))
				((((8.0 - (((X1 * X1) - X1) - 8.0)) - (((X1 + (X1 * X1)) + 4.0) + 4.0)) /
15	4	6.387	5.37576	6.0) + (X1 * X1))
16	4	6.697	5.9177	((X1 + ((2.0 + ((X1 + (X1 * X1)) - X1)) / (3.0 / 2.0))) - X1)
17	3	6.261	4.5918	((X1 * X1) - (((X1 * X1) - 4.0) / 3.0))
				(((0.0 - (X1 - X1)) / (X1 + (((X1 * 4.0) - ((((((7.0 / 5.0) + (((((X1 * 4.0) + ((((X1 * 4.0) + (((((X1 * 4.0) + (((((X1 * 4.0) + ((((((X1 * 4.0) + ((((((X1 * 4.0) + ((((((X1 * 4.0) + ((((((((((((X1 * 4.0) + (((((((((((((((((((((((((((((((((((
18	14	23.275	111.22718	- X1) - X1) * X1) + 4.0)) / 4.0) / 8.0) - X1) - X1)) * (0.0 + (X1 * X1))))) + (((X1 * X1) + ((X1 * X1) + 4.0)) / (7.0 - 4.0)))
10	14	23.273	111.22/10	((X1 * X1) + (X1 * X1) + ((((X1 * X1) + 4.0))) + ((((X1 * X1) + ((((X1 * X1) + ((((X1 * X1) + (((((X1 * X1) + (((((X1 * X1) + ((((((X1 * X1) + ((((((X1 * X1) + ((((((X1 * X1) + (((((((X1 * (((((((((((((((((((((((((((
19	6	7.87	10.06368	((X1 + X1) + (X1 + (((X1 + X1) + ((((X1 + X1) + ((((X1 + X1) + ((((X1 + X1) + (((((X1 + X1) + (((((((((((((((((((((((((((((((((((
20	5	6.888	9.24008	((3.0 + (X1 * X1)) - (((((X1 + 3.0) * X1) - 1.0) / 3.0) + (2.0 - X1)))
21	4	6.247	5.4528	(((0.0 + ((X1 * X1) + 4.0)) + ((X1 * X1) + 0.0)) / 3.0)
22	9	10.687	29.1852	((2.0 + (X1 * X1)) / ((5.0 - 2.0) / 2.0))
23	7	8.457	14.72522	(((X1 * X1) + (8.0 / 4.0)) * (4.0 / 6.0))
24	2	5.724	4.07976	(2.0 * (((X1 * X1) + 2.0) / 3.0))
25	4	6.577	5.29842	((X1 * X1) - ((4.0 - (X1 * X1)) / ((X1 - 3.0) - X1)))
26	3	6.274	5.5149	(((4.0 - (X1 * X1)) / 3.0) + (X1 * X1))
27	7	8.73	22.27664	(((4.0 + (X1 * (X1 + X1))) / (8.0 - 5.0)) + 0.0)
28	8	10.026	22.58422	((X1 * X1) - (((X1 * X1) - ((4.0 + 4.0) - (X1 * X1))) / 6.0))
29	4	6.844	6.83886	(((7.0 - 3.0) + ((X1 * X1) + (X1 * X1))) / 3.0)
30	7	8.495	14.21442	((((X1 * X1) + 4.0) + (X1 * X1)) / (7.0 - 4.0))
AVG	5.1	7.8726	13.336	





Post Project Analysis

Summary

This genetic programming project was an excellent learning experience for all of us. By the end of the semester we had the tools needed to measure and test the project. However, throughout the semester we lacked the tools necessary to code this project ourselves.

We had three people coding different aspects of the program: Kelly working to generate random binary trees, Jed working to perform crossover and mutation to evolve the population of trees, and Dawn working training data, fitness testing, and ranking. We held onto our optimism until May 1st, when two weeks before the project was due, we didn't have any of the three sections of code working perfectly, and still faced the task of connecting the pieces.

As we shifted gears to find a package, we were rushed. Had we looked for a package sooner, we could have applied measurement software applications against various packages to make an objective decision, and we would have had more time to work with packages that we quickly eliminated because we couldn't modify or enhance to run for our application.

With our final product, a modification of a program called tiny_gp, we had it running for a few days, but could not get a successful outcome. Finally, we had a breakthrough on May 10th, two days before our presentation. We launched into testing and measurement as quickly as we could. The team pulled together, and we felt our final product and presentation were of good quality.

In the week that followed the presentation, we continued to work with tiny_gp, conducting 30 random, timed-tests and expanding the operand set. This additional information is reflected in the Testing section of this, our final report.

Lessons Learned

Time Measurement

It would have been beneficial to:

- (a) Define our time categories more precisely from the beginning
- (b) Report hours weekly rather than after long spans of time, and
- (c) Allocate meeting hours toward specific time categories rather than toward a general "meeting" category.

Each one of our meetings revolved around a specific task such as analysis, coding, or document preparation as a group, as is evident in our meeting minutes. However, we could have given more time to research and analysis of the overall project "together as a team with everyone participating actively in discussions. This could have been achieved by attaching responsibilities to individuals with set timeline expectations earlier on so each person knew what was expected of them and when... rather than do what you can. Our measurement of time-spent is not as accurate as it should be due to weakness in these areas.

Testing

We should have shifted our pre-conceived ideas about testing only the end product (pass or fail), to methodically testing our work in progress so that we could optimize our final product result. For example, we

did not record test results as we modified our package code. We changed settings like population size, tree depth, etc. to achieve a better end result, but had no pattern or record of best combination of settings. It would have been beneficial to use a methodical plan to record those in-process tests.

Package Selection

For package selection we should have (a) started the search sooner, (b) used software measurement applications to compare the package candidates to one another, (c) established weighted criteria for what we would want in a package, and (d) graded each package in each weighted category so that we could assign a numerical/objective value to each package. Subjectivity would still have been evident in the steps of weighting of the criteria and grading, but the other steps in the process would lend some objectivity to the selection process.

Sometimes when you get stuck in code development, it's a sign you need to scrap what you are working on and start over (as painful as that is). Our developer, Kelly Heitz, spent a significant portion of her coding hours attempting to get a recursive insert method working on our binary trees in level order. The original code contains multiple attempts, all commented out with a description of what each attempt did instead. It was only when she "gave up" and began working on the Tiny GP program that Heitz felt she finally understood how nodes could be added to the tree in the fashion desired.

Measurement

We learned how to use tools to do measurement, which greatly improved our team's efficiency toward the end of the project. In order to implement accurate measurements of our code, we searched for and implemented CodeAnalytix as a shared tool. This plug-in allowed us to apply lessons learned in class regarding code cohesion and complexity.

Task Prioritization

Since we became stuck trying to write our own code, our team learned how to prioritize as a group. We worked collaboratively to ensure that tasks of the highest priority were accomplished first, and then worked as possible on additional items that we thought would be beneficial (rather than necessary) within our final program and report.

Software Configuration Management (SCM)

The historical record and evolution of all of the code we developed and our final modified package code for tiny gp are located as .txt files on Google Docs: **GP group project > SCM code files**. Link:

https://drive.google.com/folderview?id=0BzpM7gg1JvssZkdNd3hpX2hPOTQ&usp=sharing&tid=0BxP-i4iTYfT5LWtKcDFOU1VvbDQ

User Manual

Tiny_GP Program User Manual

Pre-Installation:

There are two ways to run our Genetic Program:

Integrated Development Environment

*Eclipse was used for the development of the final product and is the recommended IDE:

Download Eclipse from: https://www.eclipse.org/downloads/

Command Prompt

* Note: Both run methods require installation of the Java Development Kit as a prerequisite.

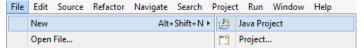
Download JDK from: http://www.oracle.com/technetwork/java/javase/downloads/jdk8-downloads-2133151.html

Installation:

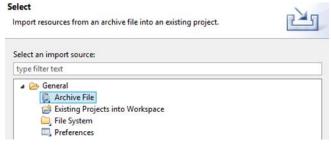
1. Download the tiny_gp folder. Link to this folder:

If using Eclipse as your IDE:

1. Click the **File** tab, select **New**, and create a **Java Project** in Eclipse.



- 2. Right click on your project. A drop down menu will appear.
- 3. Select **Import...** from the list of available options.
- 4. In the pop-up import window which appears, select the following:
 - a. Expand the General file
 - b. Select Archive File



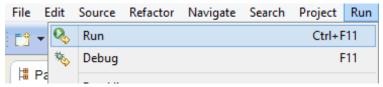
5. Click **Next** at the bottom of the import pop-up screen.



6. Select the **Browse** button to open a file explorer.



- 7. Select the tiny_gp folder from the download location.
- 8. Run the project by clicking the Run tab, and selecting Run (the first option in the drop down menu).



If using the Command Line:

PC

- 1. Open the Command Prompt window by clicking the **Start** button .
- 2. Click All Programs.
- 3. Select Accessories.
- 4. Select Command Prompt.

Mac

- 1. Open Finder. (Finder is available in the Dock.)
- 2. Select **Applications**, then chose **Utilities**.
- 3. Double click on Terminal.

Run the Program

- 1. Type the directory location where you saved the tiny_gp download.
- 2. Append "dir" to the directory name to show the contents to ensure tiny_gp.java is included as a file.
- 3. Enter "javac tiny_gp.java" to compile the code.
- 4. Enter "java tiny_gp" to run the program.