

Symbolic Regression with Genetic Programming

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Abstract—Symbolic regression is applied to an arbitrary function utilizing genetic programming techniques where the genetic program is represented by an expression tree. The initial population was generated with ramped half and half with randomly constructed trees. Candidates for the next generation were chosen by tournament selection and rank selection based on the fitness of the individual where the fitness was determined by the sum square error (SSE). These individuals were subject to crossover and heavily mutated. It was determined that the best fitness an individual could achieve was an SSE around 9.3, which is much higher than the goal of 0.5.

Index Terms—symbolic regression, genetic programming, expression trees

I. INTRODUCTION

REGRESSION is utilized heavily in all fields when the functional form is known and only the coefficients have to be determined. When the functional form is not known a functional form has to be inferred from the underlying model which can often be difficult to derive and fraught with assumptions. Symbolic regression (function discovery on a multivariate data set) is a form of regression in which the functional form and the coefficients are determined. Symbolic regression is then very different but similar to regular regression; in regression the functional form is known and only the coefficients need to be determined while in symbolic regression the coefficients and the functional form are not known. In regression it is possible to follow the gradient of the error surface for rapid convergence, but for symbolic no such error surface exists, and therefore a different type of search is required. Genetic programming is an automated method for creating solutions to high level statement of the problem by evolving individual solutions in order to optimize a fitness function is then an ideal method to solve this problem. Genetic programming attempts to mimic the evolutionary process by creating an initial population, only letting the best individuals in a population reproduce, and using an analogy of DNA mutations and the crossing of genetic material that occurs during breeding.

II. METHODOLOGY

A symbolic regression individual was modeled using an a recursively defined expression tree. An initial population (forest) of trees was generated and the fitness was computed for each of the individuals. Generations of the forest were computed utilizing genetic operators to evolve the old generation into the new. The following sections explain each part in detail.

A. Expression Tree Representation

Expression trees were used to represent symbolic functional forms due to their ease of use and the ability to easily apply genetic operators such as crossover and mutations. The expression trees were implemented recursively in c as linked nodes. Trees were created recursively in a doubly linked list. Provided a maximum depth, the function would call itself until the the maximum depth was reached. The option to prune the trees to create a more diverse initial tree structure was also implemented, but in practice preliminary pruning made an insignificant difference on the final results and was not used. The algorithm is shown in Algorithm 1. All of the operators were represented as strings, with the evaluation of a tree performed by a series of case statements. The unary functions cos and sin were overloaded to contain an additional competent which represents the magnitude. Trees were constructed in a random manner, with equal probability of picking any operator from the function set. A terminal node was chosen either by max depth or a pruning probability. Terminal node operators (values) were not chosen with a user supplied probability. In addition the terminal set only included two operators, `x` and `value`. The `x` operator represented the value on which to evaluate the expression tree, while the `value` keyword represented a real number uniformly chosen on the interval (-1,1). Table I enumerates the function and terminals sets used.

Table I
FUNCTION AND TERMINAL SET

	Operators
Function Set	$b \sin a, b \cos a, b\sqrt{ a }, a ^b, +, -, \times, /$
Terminal Set	x, \mathbb{R}

B. Initial Population

The initial population was generated using ramped half and half in which half of the population are randomly generated trees to the maximum depth, while the other half of the population was created of full trees ranging from a tree depth of 2 to the maximum depth. The initial population was assured to have no duplicates (by comparing tree structure). It is possible that trees with different structures evaluate to the same result, an example of which is provided in Figure II-B.

C. Genetic Algorithm

The genetic algorithm typically consist of four tasks: creating an initial population, evaluating that populations fitness,

Algorithm 1 Random Tree Creation

```

node *buildTree(node* parent, int depth,
double pruneProb, double constProb){
node *tree = NULL;
if (depth == 0 || (drand(0,1) < pruneProb
&& parent != NULL)) {
tree = leafNode(constProb);
tree->parent = parent;
return tree;
}else{
node *tree = funcNode();
tree->parent = parent;
tree->left = buildTree(tree, depth-1,
pruneProb, constProb);
tree->right = buildTree(tree, depth-1,
pruneProb, constProb);
return tree;
}
}

```

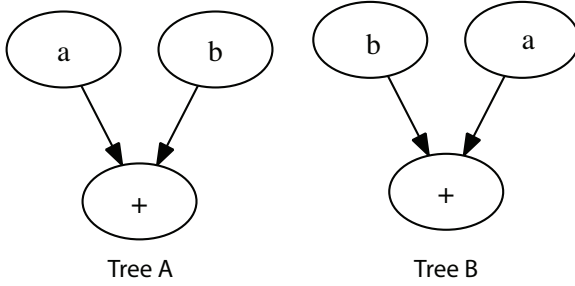


Figure 1. Semantically different trees can mathematically be equivalent

selecting members of the current population to breed, and then applying genetic operators to the selected members to breed the new population. This is completed for either a maximum generation is reached or the desired fitness is achieved. In the symbolic regression problem the fitness of an individual is defined as the sum squared error with lower fitness being considered desirable.

Algorithm 2 Genetic Program Outline

```

1: while error > goal do
2:   for all  $t \in F$  do
3:     Compute fitness
4:   for all  $t \in F$  do
5:     Choose individuals based on fitness
6:     Select individuals for next population
7:     Crossover selected individuals
8:     Mutate selected individual

```

D. Population Selection

Population selection was achieved using a hybrid approach of tournament selection and rank selection. In rank selection the individuals were selected based on their rank, where the rank was determined by their fitness (lower SSE correspond to

a higher fitness). Tournament selection was used to generate the other component of the population for breeding. Two trees were chosen at random from the entire population and the tree with the highest fitness was added to the population for breeding. There was no restriction on trees being selected more than once. Algorithm 3 outlines this approach.

Algorithm 3 Candidate selection for the next population

```

1: while numAdded < populationSize do
2:   Determine tournament selection fraction
3:   Determine rank selection fraction
4:   Rank population members based on fitness
   Preform Rank Selection
5:   while i < TotalRankNumber do
6:     Copy ranked tree i
7:   while j < TotalTournamentNumber do
8:     draw a random tree, t1
9:     draw a random tree, t2
10:    if fitness(t1) > fitness(t2) then
11:      keep t1
12:    else if fitness(t1) < fitness(t2) then
13:      keep t2

```

E. Genetic Operators

Individuals selected for reproduction are subjected to genetic operators to breed the next generation. Genetic programs generally contain two genetic operators, crossover and mutation. Crossover serves to create new members of the population by interchanging the genetic material (tree structure) of two parents in which significant changes in the tree solutions are achieved. Mutation serves to slightly modify an existing solution.

1) *Crossover*: Crossover is defined in genetic programming as the swapping of sub-trees between individuals in the population, which is meant to mimic the crossover of genes that occurs during evolution. As the solutions are recursively defined, crossover was implemented by selecting randomly selecting a two nodes from different trees and swapping their parents.

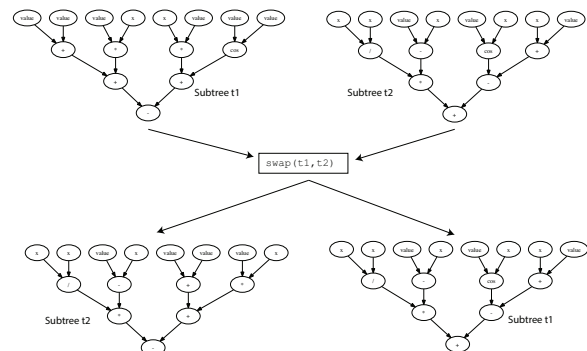


Figure 2. Example of Crossover

2) *Mutation*: Mutation, unlike crossover, does not involve changes to the structure of the solution but rather operators within the solution. For example, in symbolic regression mutation will randomly (defined by the mutation rate) change the node function and leaf values, but it will not add or prune branches to the tree structure. Mutation was implemented as a recursive traversal of the tree, switching the function types with other functions types according to a user supplied probability. The terminals mutations were implemented in a similar fashion, but if the node was a value leaf the leaf was randomly set to a new random variable. An example of mutation is shown in Figure II-E2.

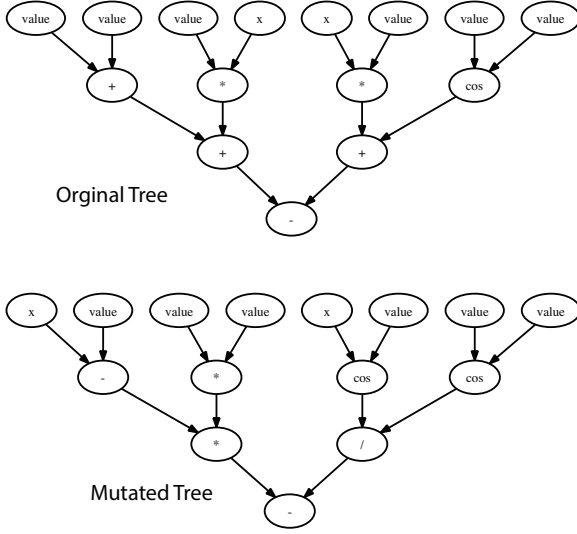


Figure 3. Example of mutation

III. RESULTS

Preliminary trials were completed at various depths in order to determine the optimal tree depth for the problem. Population sizes were set to be 500 individuals, and the genetic operators were completed for 50 generations. The initial populations were generated using a ramped half and half method. Tree depths were chosen to be 6, 8, 10, 12 and 14. As shown in Figure 4 smaller tree sizes tended to preform better than larger tree sizes. It was then determined that the best tree size would be around a tree depth of 8.

It was discovered that for a large population size (1,000) individuals the diversity only minorly changed from the population being fully diverse. This is to be expected in this implementation of the problem because the value nodes are compared by the node value and it is unlikely for two randomly drawn real values be equal (Figure 5). None of the trees generated had an sum squared error in the target range of 0.5. The best expression tree generated had a sum squared error of 9.30 with the functional form of (1), while the expression tree is shown in Figure III.

$$a \cos x \quad (1)$$

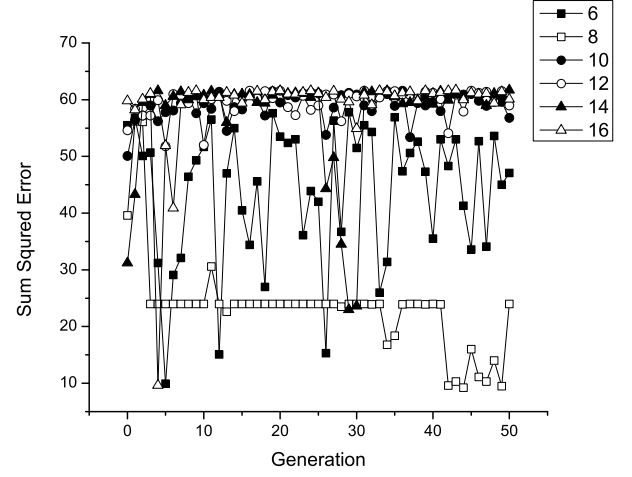


Figure 4. Sum Squared Error per Generation for various tree depths. Populations were maintained at 500 individuals, chosen with ramped half and half.

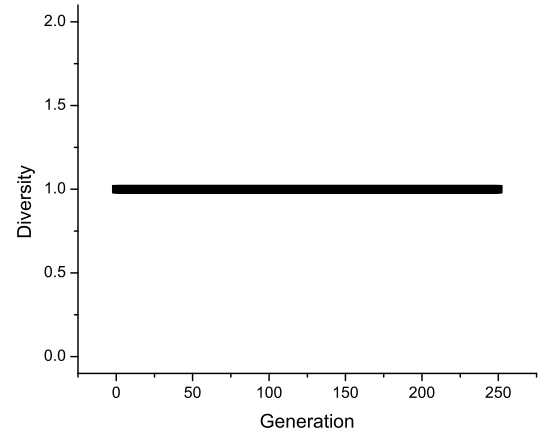


Figure 5. Diversity of the population (1,000 trees of Depth 6)

Various attempts were made to attempt to avoid this local minima but none proved successful. The first attempt was to introduce an very large mutation rate (90%), but without restricting a periodic function from occurring near the root of the tree this only resulted in another tree being switched to a periodic function and taking the previous spartan's place. It was then attempted to introduce large crossover, but if the cosine was the root node in the tree (as it is for a SSE of ~10) this only served to modify the inputs to the cosine, to which the cosine is generally resilient. The final attempt was to increase the minimum depth in the ramped half and half, but this resulted in trees that had SSE's in the 100 after a few hundred generations and was discarded.

IV. CONCLUSIONS

Genetic algorithms have been demonstrated to have the capability of performing simple symbolic regression. It is well known that genetic algorithms can preform complex symbolic

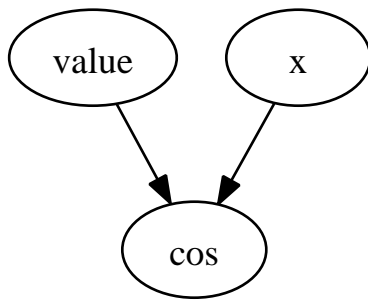


Figure 6. Expression Tree of optimal solution

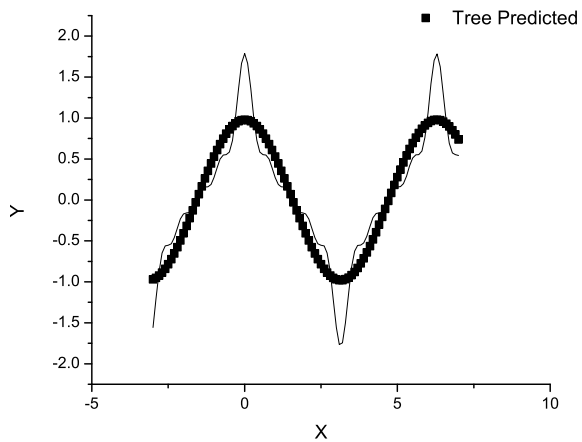


Figure 7. Comparison between data and generated model

regression, but as implemented the algorithm was only capable of building a model that captured the first order behavior of the model (Figure 7). It is thought that unless a way is devised to avoid the local minima provided by the cosine a genetic solution to this problem will have a difficult time converging in a timely manner. This can be cast in evolutionary terms where some mutations might make an individual more fit (such as pigs having wings), but because the species is already pretty well suited the mutation is not very likely. Future work could be directed into casting the problem into a light such that solutions are heavily penalized for being in this local minima. Such a method might be to transform the data into a space (perhaps with a Fourier transform) where only the optimal solution is favored.

ACKNOWLEDGMENT

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