Preliminary Evaluation of Symbolic Regression Methods for Energy Consumption Modelling

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Abstract: In the last few years, energy efficiency has become a research field of high interest for governments and industry. To understand consumption data and provide useful information for high-level decision making processes in energy efficiency, there is the problem of information modelling and knowledge discovery coming from a set of energy consumption sensors. This paper focuses in this problem, and explores the use of symbolic regression techniques able to find out patterns in data that can be used to extract an analytical formula that explains the behaviour of energy consumption in a set of public buildings. More specifically, we test the feasibility of different representations such as trees and straight line programs for the implementation of genetic programming algorithms, to find out if a building consumption data can be suitably explained from the energy consumption data from other similar buildings. Our experimental study suggests that the Straight Line Programs representation may overcome the limitations of traditional tree-based representations and provides accurate patterns of energy consumption models.

1 INTRODUCTION

The implementation of mechanisms to reduce energy consumption is one of the main objectives at national, European and international levels (Maros et al., 2016). In fact, in (Yu et al., 2010) it is shown that energy consumption in Europe and North America had an increase of 1.5% and 1.9% per year (respectively) from 1994 to 2004. Therefore, one of the most important requirements to address the improvement of energy efficiency is to perform an effective monitoring of buildings using sensors, and to use this information for high-level decision-making processes.

There is a plethora of research proposals in the literature for extracting knowledge from the sensed data. Due to paper length limitations, we cannot afford to make an extended revision of the state-of-the-art in this topic, so we refer the reader to the survey in (Pan et al., 2014) for a complete review. However, we emphasize that most of these efforts in knowledge discovery for energy consumption data might be classified in two main topics: Energy demand prediction (Molina-Solana et al., 2014) and creating user profiles to classify the consumption (Molina-Solana et al., 2014). However, in order to perform a more accurate high-level decision-making processes, it is also necessary to understand the data using abstraction models, find hidden relationships in the consumption data, and extract knowledge useful for the decision making.

In this paper, we address the problem of energy consumption data modelling and knowledge discovery. More specifically, we test the feasibility of the use of symbolic regression and genetic programming to infer an analytical formula that explains the relationship between the energy consumption data of similar buildings in a compound, to analyze and discover hidden knowledge in energy consumption data. Our experiments are targeted at testing of different representations of algebraic expressions, to know the computational power of each model in practice for the specific problem of energy consumption modelling. For that purpose, we select a simple problem in a real dataset, where a general counter value should be modelled as aggregation of the energy consumption data from other 5 different buildings. In the experiments, we show the computational power of two algebraic formula representation mechanisms to infer the previous relationship, and also the capability of the learning stage to distinguish between relevant and non-relevant data for the general counter modelling.

This paper is organized as follows: Section 2 describes the fundamentals of symbolic regression. Sections 3 and 4 introduce the representations used in this research and the genetic operators for each represen-
utation, respectively. Section 5 show the experimentation of the proposals over a set of benchmark functions and a real energy consumption dataset. Finally, conclusions and future research are discussed in section 6.

2 FUNDAMENTALS OF SYMBOLIC REGRESSION

Regression analysis (M., 2007) is one of the basic tools of scientific research. It is used to fit a functional model that represents a relationship between independent and dependent variables. Traditionally, this kind of problems has been solved with algebraic methods, where the researcher provides a hypothesis about a functional model with a set of parameters, and the goal is to optimize these parameters for the studied dataset. Equation 1 shows the parametric model of regression analysis, where \( \hat{x} = (x_1, x_2, \ldots, x_n) \) stands for the set of independent variables of the data, \( f \) is the functional model hypothesis, \( \tilde{w} = (w_1, w_2, \ldots, w_m) \) are the parameters of the model, and \( \hat{y} = (y_1, y_2, \ldots, y_l) \) and the dependent variables of the problem. In these cases, since \( \hat{x} \) and \( \hat{y} \) are the problem data, and \( f \) is a function established as model hypothesis, the regression problem is solved by finding the best values for the parameters \( \tilde{w} \), which are not known in advance. As an example, the most simple case in regression analysis is the well known linear regression, whose functional model can be written as \( y = f(<w_1, w_2 >, x) = w_1 * x + w_2 \).

\[
\hat{y} = f(\tilde{w}, \hat{x})
\]  

A limitation of classical regression arises when the data properties are unknown in advance, it is difficult to find a pattern that explains the dataset, and therefore it is hard to establish a suitable hypothesis for the function \( f \). This problem is even less tractable in the multivariate case, where graphical analyses lack of enough expressiveness to show relations between multidimensional data.

To solve these limitations, the use of symbolic regression attempts to generalize the traditional problem of regression analysis by assuming that \( f \) is unknown, and developing techniques targeted at finding a suitable model \( f \) and parameters \( \tilde{w} \) that minimizes an error expression such as \( ||\hat{y} - f(\tilde{w}, \hat{x})|| \). In symbolic regression it is assumed that \( \hat{y} \) and \( \hat{x} \) are the only components known in advance. Therefore, the goal of symbolic regression is to find an algebraic expression that models the behaviour of the dependent data as function of independent data. Techniques like genetic programming (Langdon, 1998) have been developed to solve this problem.

Symbolic regression techniques have been applied traditionally in a wide variety of real applications. Besides of being used to solve mathematical optimization problems, they have been of practical application in decision making in economics, chemical processes optimization, etc. For instance, in (Duffy and Engle-Warnick, 2002) it is described how can be used symbolic regression to uncover simple data generating function that have the flavor of strategic rules in economic decisions. In the work (McKay et al., 1997), symbolic regression has been used to model chemical processes systems, to solve problems about vacuum distillation column and a chemical reactor system. On the other hand, in (Schmidt and Lipson, 2010) they explore the use of symbolic regression to perform unsupervised learning by searching for implicit relationships, specifically they present a successful method based on implicit derivated. In (Davidson et al., 2001) the authors use symbolic regression in two real-world problems, approximating the Colebrook-White equation and rainfall-runoff modelling.

In this work, we test the feasibility of the use of symbolic regression to model energy consumption in a set of public buildings, under the hypothesis that the resulting models obtained from the symbolic regression approach will be useful for high-level decision making processes regarding energy efficiency. The following subsection describes in depth the basis of our approach, which is based in genetic programming.

2.1 Introduction to Genetic Programming

Genetic programming (Langdon, 1998) can be seen as a supervised learning method based on biological evolution. Genetic programming fundamentals are inspired in genetic algorithms, and it has been used in previous works to solve optimization problems like symbolic regression (Alonso et al., 2009), digital signal processing (Alcazar and Sharman, 1996), solving differential equations (Tsoulos and Lagaris, 2006), tasks of evolving robotic behaviours (Lazarus and Hu, 2001), grammatical inference (Lankhorst, 1994), automatic program generation (Koza, 1994), etc. If we focus in the problem of symbolic regression, the goal of genetic programming to evolve a set of algebraic expressions encoded as chromosomes, according to Darwinian evolution principles of genetic algorithms and, as fitness measure, the minimization of an error function that explains the behaviour of dependent variables regarding the independent variables in a specific dataset.
Since the basic structure of genetic programming is similar to genetic algorithm, figure 1 shows the evolutionary process. Firstly, a set of individuals, where each individual represents an algebraic expression. Then, a selection operator is applied to obtain a subset of individuals or parents which will be used for genetic recombination. In this work, we use the tournament selection operator (Pandey, 2016). In generational evolutionary scheme, there are selected as parents as many individuals as the size of the initial population. On the other hand, in stationary scheme, we select just 2 parents. After that, the crossover operator is applied to generate as many children individuals as parents. In this work, two parents are recombined to obtain two new children solutions. These children preserve genetic material from both parents. After the crossover, a mutation operator takes place over the children with a probability, imitating biological evolution. Once the previous genetic cycle is finished, the new children are evaluated according to a fitness measure. In the case of symbolic regression, since it is also necessary to optimize the function parameters $\bar{w}$, in this work, we use the Simplex method (K. and K., 1989) to obtain an approximation of $\bar{w}$ before the algebraic expressions encoded into the children are evaluated. Finally, the initial population is replaced with the new children and an evolutionary generation is finished. In a generational scheme, all children replace the whole population, while in the stationary scheme the children replace their parents.

In addition to this basic evolutionary scheme, we also use an elitism factor, to ensure that the best solution found during the evolution remains in the population. In case that the replacement does not include the best solution in the new population, this one replaces the worst solution in it.

Traditionally, each individual is encoded using a tree structure to represent an algebraic expression in genetic programming. A new alternative to encode algebraic expressions is straight line programs (Alonso et al., 2009) (SLP). A SLP is a lineal structure that represents a non-linear structure as a graph. This alternative allows the SLP representation to reuse subexpressions and represent an algebraic formula in a structure smaller than trees. So, the main advantages of SLP representation against tree representation is the possibility to better explore the search space, being smaller representations of the same formulas in SLPs than in trees. In this paper, we make an experimental evaluation of both approaches to solve our problem. Both representations are described in the next section.

3 GENETIC PROGRAMMING REPRESENTATIONS FOR SYMBOLIC REGRESSION

3.1 Tree Representation

The traditional representation to encode algebraic expressions are tree structures (McKay et al., 1995). This representation allows to generate regular grammars and context-free grammars. On the other hand, this kind of representations could be limited because they are non-linear structures represented by non-linear representations. Thus, the search space is complex, and recombination and mutation operators proposed in the literature may not provide suitable results.

As an example, one limitation is that tree structures cannot reuse algebraic subexpressions. In figure 2 we can see a tree representation of the algebraic expression $x^4 + \cos(8 \cdot x) + 3 \cdot \cos(8 \cdot x)$. We emphasize how we use the expression of $\cos(8 \cdot x)$ in two parts of the tree. Finding such structure with replicated submodules requires a larger exploration of the search space, therefore increasing the computing time in the genetic programming algorithm. In the same way, a new problem to solve symbolic regression problems with genetic programming appears, it is called like bloat (Naoki et al., 2009). This prob-
lem involves the increased size of analytical formula obtained by genetic programming, producing unbalanced trees with high depth in one branch and low depth in the other. This fact also makes difficult to find an optimal solution, and the computing time of the algorithm is also increased. In order to minimize the computational complexity, a new representation called Straight Line Programs is studied in this paper.

3.2 Straight Line Programs

Straight Line Programs are based on Straight Line Grammars (Benz and Kötzing, 2013). A Straight Line Grammar is a non recursive context-free grammar. Straight Line Programs have been used in the literature to represent algebraic operations (Berkowitz, 1984), geometry problems (Giusti et al., 1998), solving polynomial equations (Krick, 2002), document clustering (Sequera et al., 2012), fast sparse matrix-vector multiplication (Neves and Araujo, 2015) etc.

In the previous section we discussed limitations of the tree representation. Using Straight Line Programs as individuals representation in genetic programming, we pursue to improve our results obtained with traditional representations like trees. As a consequence, in our experimentation we show that SLPs representation implies a minimization of the search space and computational time against representation.

The Straight Line Program structure is represented by a table. Each row of the table contains an expression, this expression is built applying a mathematical operator (+,-,*,sin, etc...) to a set of operands. These operands could be variables, constants or references to other rows of the table. Figure 3 shows an example of SLP structure and its graph representation. To evaluate this algebraic expression we must evaluate from last element of the table to the beginning.

![Figure 3: SLP representation.](image)

Figure 3 shows the advantages of this representation. It is a linear structure that models a nonlinear graph structure. Also, this representation allows the reuse of some algebraic subexpressions in the SLP. For example, we use \( \cos(8 \times x) \) in two parts of this representation. Using this structure as representation of each individual in genetic programming, we reduce the search space in respect to the tree representation, and therefore the computing time of genetic programming procedure is also lessen.

4 GENETIC OPERATORS

4.1 Genetic Operators for Tree Representation

In this section we describe the different crossover and mutation operators that we use in genetic programming with a tree representation. We assume that the user has configured a set \( O = \{ o_1, o_2, ..., o_n \} \) of atomic operators (+,-,*,sin,cos,...), and \( T = \{ t_1, t_2, ..., t_m \} \) a set of terminal symbols. These terminal symbols encompass the set of independent variables of the dataset and a set of unknown constants or function parameters \( \vec{w} \). In addition it is required a parameter \( M \) that contains the maximum tree depth allowed. Then, the individuals in the initial population are generated as follows:

Firstly, for each individual, it is selected a random value \( l \) between 2 and \( M \), the current individual tree depth. Then, it is created the root node containing an operator selected randomly. The left and right branches of the operator are also generated randomly with symbols from the operator set \( O \) until level \( l \). Then, the symbols for these leaf nodes are generated randomly from the set \( T \). Using this strategy, we ensure that the initial population contains balanced trees. After all individuals have been generated with this procedure, the evolutionary process begins.

4.1.1 Crossover Operator

The aim of this operator is to genetically recombine two parents to generate two new children. Figure 4 shows two candidates (father on the left, mother on the right) to generate their corresponding children. The procedure of this operator is as follows:

![Figure 4: Example parent trees for crossover.](image)

A node in the tree is selected at random for each parent. In the example of figure 4, the selected nodes...
are named as $k$ and $t$. Then, the two children are generated by exchanging the subtrees with root $k$ and $t$ in both parents. This result is shown in figure 5.

Figure 5: Example children trees for crossover.

### 4.1.2 Mutation Operator

The purpose of this operator is to imitate the behavior of biological evolution, mutating individuals randomly to improve the exploration of the search space.

In this representation we use two different mutation operators: One simple whose aim is to maintain an exploitation of the search space, and another one complex to increase the exploration.

Firstly, figure 6 shows an example of the use of the simple mutation operator. The simple mutation operator selects a random node of the tree; then, if the selected node is a mathematical operator, it is exchanged for another random operator in $O$. On the other hand, if the selected node is a constant or variable, it is exchanged for another random terminal symbol in $T$. Please note that the tree structure is not altered using this mutation operator.

Figure 6: Example of Simple Mutation Operator.

On the other hand, figure 7 shows the complex mutation operator. This operator selects a node of the tree at random, and then this node is pruned and replaced with a new randomly generated subtree. As opposite to the simple mutation operator, the complex one alters the tree structure of the individual.

Figure 7: Example of Complex Mutation Operator.

In each generation of our genetic algorithm, if a mutation must be carried out over an individual, it is selected simple or complex mutation for its application with 50% of probability.

### 4.2 Genetic Operators for SLP Representation

In this section, we describe the evolutionary operators for SLPs. To build the initial population, let $O = o_1, o_2, ..., o_n$ be a set of operators ($+$, $-$, $\ast$, $\sin$, $\cos$, ...), and $T = t_1, t_2, ..., t_m$ be a set of terminals. To build the initial population, figure 8 shows the procedure to build a new individual containing an algebraic expression with SLP representation, where $n$ is the number of rows generated in the SLP during the procedure, and $m$ is the total size of the SLP table. Also, generate expression 1 generates a random operator and two random operands. These operands could be elements of the set $T$. On the other hand, generate expression 2 also generates a random operator and two random operands but, in this case, these operands may be a symbol in $T$ or a call to previous entries of the SLP table representation in the range $\{1..n-1\}$.

For this representation, new crossover and mutation operators have been developed in (Alonso et al., 2009). Since these are the only operators we have found in the literature for SLPs, we will use them in our experimental evaluation. Future work will be targeted at developing new crossover and mutation strategies as part of our research.

### 4.2.1 Crossover Operator

This section shows the crossover operator applied to the straight line program representation. This operator is applied over two individuals or parents. The procedure of this crossover operator is as follows: A random position $k$ and $t$ are selected randomly in the parents’ SLP table. After that, The selected rows and
all those rows involved in the calculation of $k$ (in the second parent) are exchanged to generate the two children. Figure 9 shows an example of the selection of $k$ and $t$ in two parents, and figure 10 illustrates the offspring generation result.

Once two children have been generated using this procedure, the evolutionary process continues to check if a mutation operator should be applied over the offspring.

4.2.2 Mutation Operator

The procedure to apply the mutation operator in straight line programs is to generate a random position between 0 and $m$, where $m$ is the size of the SLP table. This random position is called as $k$. Then, we select a random element of this row (operator or operands). If the selected item is an operand, we replace it with another operand (constant, variable or a position of the SLP table). On the other hand, if the selected item is an operator, it is replaced by another randomly generated operator.

5.1 Preliminary Experiments in Benchmark Data

In order to validate each evolutionary model, and tree and SLP representations, we use a set of classical benchmark functions proposed in (Alonso et al., 2009):

\[
\begin{align*}
    f_1(x) &= x^4 + x^3 + x^2 + x 
    f_2(x) &= \exp(-\sin(3x + 2x)) 
    f_3(x) &= 2.718x^2 + 3.1416x 
    f_4(x) &= \cos(2x) 
    f_5(x) &= \min\{\frac{2}{x}, \sin(x) + 1\}
\end{align*}
\]

For each function we will use each implementation of the genetic algorithm (generational and stationary) and each representation (straight line programs and trees). After that, we will discuss if the SLP structure has potential being compared to the tree representation.

5.1.1 Data Generation and Experimental Settings

For each benchmark function data we generated 100 random data in the domain $[-100, 100]$. The resulting datasets were stored to be used by each genetic algorithm and representation with the aim that all methods run under the same initial conditions.
Thus, we have four proposals to test: Generational scheme with trees (GGA-T) and with SLP (GGA-SLP), and stationary scheme with trees (SGA-T) and with SLP (SGA-SLP). For each configuration, experiments will be executed 30 times for each benchmark function so that statistical analyses can be carried out.

Table 1 contains the parameters for each function described in the previous section, according to the experiments previously carried out in these benchmark functions in the literature (Alonso et al., 2009).

<table>
<thead>
<tr>
<th>Function</th>
<th>Size</th>
<th>Operators</th>
<th>Tree representation</th>
</tr>
</thead>
<tbody>
<tr>
<td>$f_1(x)$</td>
<td>30</td>
<td>$+,-,\cdot,\sqrt{\cdot}$</td>
<td>$f_1(x)$</td>
</tr>
<tr>
<td>$f_2(x)$</td>
<td>20</td>
<td>$+,-,\cdot,\exp,\sin$</td>
<td>$f_2(x)$</td>
</tr>
<tr>
<td>$f_3(x)$</td>
<td>15</td>
<td>$+,-,\cdot,\cos$</td>
<td>$f_3(x)$</td>
</tr>
<tr>
<td>$f_4(x)$</td>
<td>20</td>
<td>$+,-,\cdot,\min,\sin$</td>
<td>$f_4(x)$</td>
</tr>
</tbody>
</table>

For each execution of both genetic algorithms we used a population of 200 individuals and 50000 evaluations as stopping criterion. On the other hand, it has been established a crossover probability of 99% and 1% for the probability of mutation. Finally, equation 7 shows our fitness function. This fitness function is the mean squared error between the output of each benchmark functions ($f(x_i)$) and its real values ($y_i$), where $m$ is the number of samples in each dataset. The objective of the genetic programming algorithms is to minimize $\epsilon(f)$.

$$\epsilon(f) = \frac{1}{m} \sum_{i=1}^{m} (f(x_i) - y_i)^2$$  

5.1.2 Results in Benchmark Data

In this section, we discuss the results obtained for each benchmark function. Table 2 shows the fitness value (MSE) of the best solution found for each genetic algorithm, individuals representation and benchmark function. According to the results, we can see that the best genetic algorithm scheme for tree representation is the stationary, since the mean square error is lower than GGA-T in all functions. On the other hand, the best scheme for SLP representation is generational genetic algorithm. Equations 8 to 12 show the best expressions found by tree representation, and equations 13 to 17 with SLP, for functions $f_1(x)$ to $f_5(x)$, respectively. In addition, Figures 12, 13, 14, 15 and 16 show graphically the results of the best approximation function found. We can see that the best option to model the behaviour data is a SLP representation with generational scheme in all functions but $f_5(x)$ and $f_5(x)$. However, as Figure 14 and 16, SLP also found the right function approximation.

Since in both cases the SLP algebraic representation is larger than the obtained using trees, we think that this increase in the MSE is due to the Simplex method to approximate the function parameters. On the other hand, we can see that in some cases (functions $f_2(x)$ and $f_4(x)$), the tree representation proposals were not able to find an accurate representation. As a partial outcome in the preliminary experimentation carried out in this section, we conclude that SLPs have good potential to overcome local optima and to achieve better solutions than using tree representation, although the final error in the fitness measure might be affected by the parameters $\hat{w}$ estimation algorithm.

$$F_1(x) = (x_1 \times (x_1 + x_1)) \times (x_1 + 1)$$
$$+ ((x_1 + 1) \times (x_1 + 1))$$ (8)

$$F_2(x) = (x_2 - 1425.58)/$$
$$((-1100.38/(-1100.38/(-1100.38))$$
$$- (x_2/(-1425.58)))$$ (9)

$$F_3(x) = ((x_3 + 2.15) - (1.395/1.395))$$
$$\times ((x_3 * (1.395 * 1.395) * 1.395))$$ (10)

$$F_4(x) = (((6.57/((6.57 - x_4))/$$
$$((x_4 + (6.57 + 6.57))) + (x_4/570.08))))$$ (11)

$$F_5(x) = (((2.068/2.068)/2.068) - (2.068/x_5))$$
$$/((x_5 - 4.075)/4.075))$$ (12)

$$F_1(x) = (x_1 + ((x_1 * x_1) +$$
$$(((x_1 * x_1) * x_1) + (x_1 * x_1) * x_1))))$$ (13)

$$F_2(x) = \exp((\sin((x_2 * (-5))))))$$ (14)

$$F_3(x) = (((-1.85) * x_3) + (((x_3 + x_3) +$$
$$((x_3 + x_3) + ((x_3 + x_3) - (((-1.85) * x_3)$$
$$+ ((-1.85) * x_3)) + x_3))$$
$$- x_3)) + x_3))$$ (15)

$$F_4(x) = \cos(x_4 + x_4)$$ (16)

$$F_5(x) = (x_5 - (x_5 - (2.1697 + 2.1697))$$
$$/(((x_5 - (2.1697 + 2.1697)/(x_5 - 2.1697$$
$$+ 2.1697))))/((2.1697 + 2.1697) + x_3 + x_3)))$$ (17)

We emphasize that our goal is to seek an algebraic expression equivalent to initial functions described
above, for this reason we do not use a set of test data to validate our solution, if we find an algebraic expression equivalent to our initial functions, we could verify that the solution found is the right one. While SLP finds algebraic expressions equivalent to the objective functions, trees sometimes fail. On the other hand, we used the Student’s statistical test to check if there are significant differences between the error distributions of the algorithms (GGA-T versus GGA-SLP and SGA-T versus SGA-SLP). After applying the statistical test, we obtained a p-value less than 0.05 in all cases, so that we reject our initial hypothesis (non significant differences) and we can ensure that there are differences between the algorithm for SLP representation.

In the next section, we apply the proposals SGA-T and GGA-SLP over a real dataset of consumption data, to check the behavior of the methods under real problems with low noise data.

Table 2: Benchmark functions results.

<table>
<thead>
<tr>
<th></th>
<th>GGA-T</th>
<th>SGA-T</th>
<th>GGA-SLP</th>
<th>SGA-SLP</th>
</tr>
</thead>
<tbody>
<tr>
<td>( f_1 )</td>
<td>7.32E14</td>
<td>2.87E-17</td>
<td>0</td>
<td>2.324E-15</td>
</tr>
<tr>
<td>( f_2 )</td>
<td>0.619</td>
<td>0.659</td>
<td>1.81E-26</td>
<td>1.81E-26</td>
</tr>
<tr>
<td>( f_3 )</td>
<td>6.94E7</td>
<td>1.74E-24</td>
<td>0.1348</td>
<td>270.62</td>
</tr>
<tr>
<td>( f_4 )</td>
<td>0.4366</td>
<td>0.4414</td>
<td>0</td>
<td>0.0146</td>
</tr>
<tr>
<td>( f_5 )</td>
<td>9.99E-4</td>
<td>1.55E-4</td>
<td>0.0146</td>
<td>0.044</td>
</tr>
</tbody>
</table>

Figure 12: Results for \( f_1(x) \) using trees (left) and SLPs (right). Red lines are the real data, and green lines the approximated data.

5.2 Description of the Energy Consumption Dataset

In this section we analyze the energy consumption data of a set of public buildings. Here, our goal is to know the feasibility of using symbolic regression to infer the relationship between the energy consumption of similar buildings in the same compound, in the case it exists, and provide an analytical formula that explains such relationship. In this work, our test bed data come from a Campus of the University of Granada containing 5 buildings. For confidentiality restrictions, we name those buildings as \( x_1, x_2, x_3, x_4, x_5 \). We use the available data for all buildings, measured from 2012 to September of 2015.

Each building has an energy consumption sensor that measures energy consumption hourly. In addition, there is an aggregation node (general counter) that provides the sum of the energy consumption of the buildings \( x_1, x_2, x_3, x_4 \). Thus, our initial hypothesis to test is that the value of the general counter \( y = x_1 + x_2 + x_3 + x_4 + \epsilon \), where \( \epsilon \) stands for an error coming from data preprocessing. We will use symbolic regression with algorithms SGA-T and GGA-SLP to test if the methods are able to find such relationship, to validate the feasibility of the technique to be used in energy consumption data modeling.
A prior stage of data preprocessing was required before applying the algorithms. This preprocessing is the imputation of missing values (Royston, 2004) for each energy sensor and data alignment (Rhudy, 2014). Finally, as the data are provided hourly, we applied a data aggregation procedure (Patil and Patil, 2010) to model daily consumption data. Section 5.3 shows the results after applying symbolic regression in this real dataset.

For this experimentation with real data consumption we used the following configuration: 200 individuals for the initial population and 50000 evaluations. Also, we have used the crossover (probability of 99%) and mutation (probability of 1%) previously described. Finally, equation 7 is used as function fitness.

5.3 Results in Real Energy Consumption Data

Using the experimental settings described in the previous section, we run 30 experiments so that statistical analyses could be carried out in the results. Table 3 shows the best fitness values (MSE) obtained for each representation. Again, a generational scheme in our genetic algorithm with SLP representation has found a better solution than the tree representation. Also, the computational time is less in SLP and the average fitness along all 30 runnings is lower. Therefore, here we also prove the power of SLP representation over the tree representation in real data.

<table>
<thead>
<tr>
<th></th>
<th>Trees</th>
<th>SLP</th>
</tr>
</thead>
<tbody>
<tr>
<td>Average Fitness</td>
<td>5.1142e+15</td>
<td>68.9</td>
</tr>
<tr>
<td>Better Fitness</td>
<td>192284.034</td>
<td>43.3801</td>
</tr>
<tr>
<td>Worse Fitness</td>
<td>373236.908</td>
<td>94.42</td>
</tr>
<tr>
<td>Average time (s)</td>
<td>11714.5</td>
<td>8321</td>
</tr>
<tr>
<td>Best solution time (s)</td>
<td>60.5</td>
<td>61</td>
</tr>
</tbody>
</table>

The best solution found by SLP is shown in equation 18. This formula indicates that the general counter is the sum of the following 4 building plus a constant. The term \(\cos(x_3)\) is irrelevant because its range is [-1,1] and we may consider this value as a constant for lower/upper bounds. This constant could be model a part of an error derived from data preprocessing.

\[
Y = (X_1 + (((24.476 + X_4) + (X_2 + X_3)) + (\cos(x_3))))
\]

(18)

Finally, figure 17 shows in blue the real consumption data, and in red the estimated values with the found expression 18. So we accept the initial hypothesis to test the inference capabilities of symbolic regression for energy consumption data. The general counter has been correctly modeled using the information from buildings \(x_1, x_2, x_3, x_4\), and the algorithm has been able to distinguish that \(x_5\) is irrelevant for the problem. This fact opens a future research to test if symbolic regression could be used for both energy consumption modelling and feature selection, if it is applied over a larger dataset and more complex initial hypotheses.

In order to validate the rightfulness of the resulting formula, in case we would have not known the initial hypothesis in advance, Figures 18, 19 and 20 shows the scatterplots of the general counter against the other buildings: These figures contain the real and the estimate data with the resulting formula in 18. In blue color, we can see the values of real consumption of the general counter against other building, and in red color we can see the estimated values by the formula. From these results we can conclude that the
last building (20) is not providing consumption data in the general counter, because we have modeled their consumption with an algebraic expression that does not consider this building, and also that the scatterplot of estimated data is very accurate being compared to the initial real data. Thus, as this behavior is fulfilled in all buildings, we conclude that the resulting formula obtained by symbolic regression is appropriate to model the consumption data.

6 CONCLUSIONS

In this paper, we have tested the feasibility of using a new representation in genetic programming: straight line programs. This alternative to solve the symbolic regression problem allows an improvement in computational time due to their linear structure which facilitates reuse of algebraic subexpressions, so the search space is lower in our genetic algorithm and can overcome local optima. On the other hand, we highlight the good performance in multivariate problems, being able to select the relevant variable of the problem in the real data in our experimentation.

As a general conclusion, we summarize that our findings suggest that SLP representation has a good potential as symbolic regression representation mechanism, against the traditional tree representation. In the benchmark data, we have found out that the method to estimate the parameters during the fitness evaluation stage could be improved, since it affects directly to the fitness performance and might hide good alternative algebraic expressions. In a future work, we will address this problem by testing other mechanisms for parameter estimation, and also reducing the size of the resulting SLP expression. Regarding energy consumption, we will increase our test bed data and make experiments with large datasets containing several buildings where an initial algebraic expression hypothesis cannot be provided in advance to test the results.

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REFERENCES


