

Evolution of Sorting Algorithm using Graph Structured Program Evolution

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Abstract—In this paper, we apply Graph Structured Program Evolution (GRAPE) to evolution of general sorting algorithm. GRAPE is a new Automatic Programming technique. The representation of GRAPE is graph structure, therefore it can express complex programs (e.g. branches and loops) using its graph structure. Each program is constructed as an arbitrary directed graph of nodes and *data set*. GRAPE handles multiple data types using *data set* for each type, and the genotype of GRAPE is the form of a linear string of integers. The aim of this work is to evolve a program which correctly sort any sequence of numbers. We demonstrate that GRAPE constructs general sorting algorithm automatically.

I. INTRODUCTION

In recent years a lot of Automatic Programming techniques have developed and applied to a variety of problem domains. A typical example of Automatic Programming is Genetic Programming (GP), and various extensions and representations for GP have been proposed so far. In standard GP, programs are represented as trees containing terminal and non-terminal nodes. Complex programs and hand written programs, however, may contain several branches and loops. We believe that graph representation is the nearest representation of hand written programs. Therefore, we adopt the graph structure as a representation of programs. Our method, named **GRA**ph structured **PR**ogram **EV**olution (GRAPE) [1], uses graph structure as a representation of programs. In GRAPE each program is constructed as an arbitrary directed graph of nodes and *data set*. GRAPE handles multiple data types using *data set* for each type, and the genotype of GRAPE is the form of a linear string of integers. In this paper, we apply GRAPE to evolution of general sorting algorithm. The problem of evolution of general sorting algorithm is considered challenging problem, and it needs to prepare iteration or recursion mechanism.

The next section of this paper is an overview of several related works. In section 3, we describe our proposed method, Graph Structured Program Evolution (GRAPE). Next, in section 4, we apply the proposed method to evolution of general sorting algorithm and show several experimental results. Finally, in section 5, we describe conclusions and future works.

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II. RELATED WORKS

Automatic Programming is a method of generating computer programs automatically. Genetic Programming (GP) [2], [3] is a typical example of Automatic Programming, which was proposed by Koza. GP evolves computer programs, which are usually tree structure, and searches a desired program using Genetic Algorithm (GA).

Various representations for GP have been proposed so far. GP with index memory [4], [5] was introduced by Teller and was proven that the system is Turing complete. This means that, in theory, GP with indexed memory can be used to evolve any algorithm. Linear Genetic Programming (LGP) [6], [7] uses a specific linear representation of computer programs. Instead of the tree-based GP expressions of a functional programming language (like LISP), programs of an imperative language (like C) are evolved. A LGP individual is represented by a variable-length sequence of simple C language instructions. Instructions operate on one or two indexed variables (registers) r or on constants c from predefined sets. The result is assigned to a destination register, e.g. $r_i = r_j * c$. Grammatical Evolution (GE) [8], [9] is an evolutionary algorithm that can evolve computer programs in any language, and can be considered a form of grammar-based genetic programming. GE uses a chromosome of numbers encoded using eight bits to indicate which rule from the BNF (Backus Naur Form) grammar to apply at each state of the derivation sequence, starting from a defined start symbol.

There are various representations using a graph. Parallel Algorithm Discovery and Orchestration (PADO) [10]–[12] is one of the graph based GPs instead of the tree structure. PADO uses stack memory and index memory, and there are *action* and *branch-decision* nodes. The execution of PADO is carried out from the start node to the end node in the network. PADO was applied to the object recognition problems. Another graph based GP is the Parallel Distributed Genetic Programming (PDGP) [13]. In this approach the tree is represented as a graph with functions and terminals nodes located over a grid. In this way it is possible straightforward to execute several nodes concurrently. Cartesian Genetic Programming (CGP) [14], [15] was developed from a representation that was used for the evolution of digital circuits and represents a programs as a graph. In certain respects, it is similar to the graph-based technique PDGP. However, CGP adopts the genotype-phenotype mapping. The genotype is an integer string which denotes a list of node connections and functions. This string is mapped into phenotype of an index

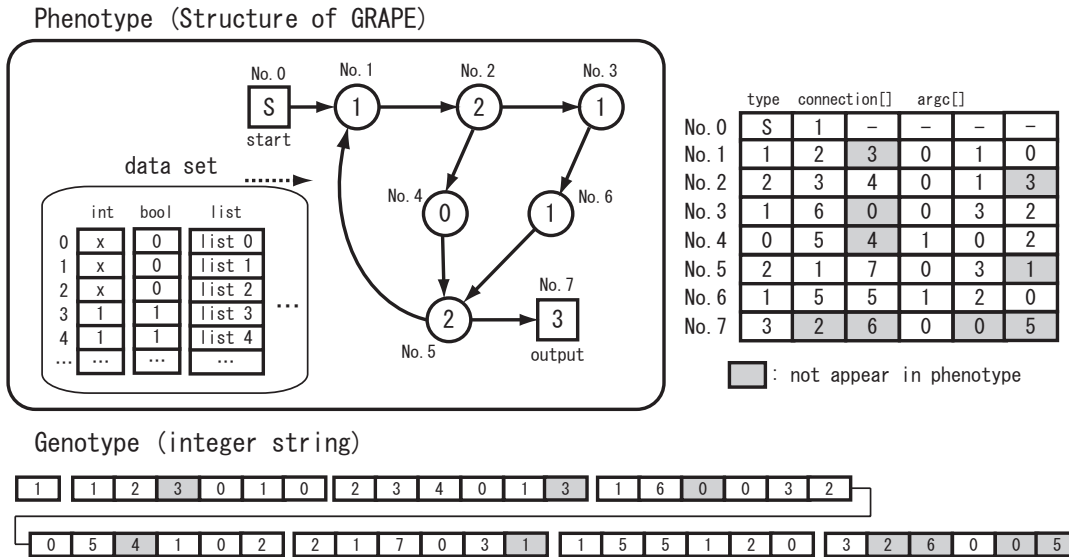


Fig. 1. Structure of GRAPE (phenotype) and the genotype which denotes a list of node types, connections and arguments.

graph. Linear-Graph GP [16] is the extension of Linear GP and Linear-Tree GP [17]. In Linear-Graph GP each program is represented as a graph. Each node in the graph has two parts, a linear program and a branching node. Recently, Genetic Network Programming (GNP) [18], [19] which has a directed graph structure is proposed. GNP is applied to make the behavior sequences of agents and shows better performances compared with GP.

Several researchers have investigated evolution of general sorting algorithm. Kinnear evolved general iterative sorting algorithm using Genetic Programming. He prepared special functions for the sorting problem, such as *Iteration Function* which is an iterative operator.

Recently two interesting Automatic Programming techniques were proposed, PushGP [20]–[22] and Object Oriented Genetic Programming (OOGP) [23]–[25]. The both method tackled the problems of generating recursive programs (e.g. factorial, Fibonacci sequence, exponentiation, sorting a list and so on) and obtained these programs automatically. PushGP evolves programs using a Push language proposed by Spector, et al. Push is a stack-based programming language. They showed that PushGP evolved a general sorting algorithm [21]. OOGP evolves Object Oriented Programs instead of the form of LISP parse tree. OOGP evolved a general recursive sorting algorithm using its recursive mechanism [25].

III. GRAPH STRUCTURED PROGRAM EVOLUTION (GRAPE)

A. Overview

Various extensions and representations for GP have been proposed so far. However, it seems that more improvements are necessary to obtain more complex programs automatically. Graph Structured Program Evolution (GRAPE) [1] constructs graph structured programs automatically. The

graph structured programs is composed of arbitrary directed graph of nodes and *data set*.

B. Structure of GRAPE

The representation of GRAPE is graph structure. Each program is constructed as an arbitrary directed graph of nodes and *data set*. *Data set* flows the directed graph and is processed at each node. Fig. 1 illustrates an example of structure of GRAPE. Each node in GRAPE program has two parts, a *processing* and *branching*. The *processing* executes several kinds of processing using *data set*, for instance, arithmetic calculation and boolean calculation. After the *processing* is executed, the current node moves. The *branching* decides the next node according to the *data set*.

Examples of node in GRAPE are shown in Fig. 2. “No.1 node” adds *data[0]* to *data[1]* and substitute for *data[0]* using integer data type, and moves to the node “No.2”. “No.2 node” decides the next node using integer *data[0]* and *data[1]*, if *data[0]* is greater than *data[1]*, connection 1 is chosen, else connection 2 is chosen. There are special nodes shown in Fig. 1. “No.0 node” is the *start node* which is the equivalent of root node of GP. It is the first node to be executed when a GRAPE program runs. “No.7 node” is the *output node*. When this node is reached, the GRAPE program outputs data and then the program halts. In Fig. 1 “No.7 node” outputs integer *data[0]*. Although the GRAPE program has only one *start node*, it can have several *output nodes*.

The representation of GRAPE is graph structure, therefore it can represent complex programs (e.g. branches and loops) using its graph structure. There are several data types in GRAPE program, integer data type, boolean data type, list data type and so on. GRAPE handles multiple data types using the *data set* for each type.

To adopt evolutionary method, genotype-phenotype mapping is used in GRAPE system. This genotype-phenotype

Examples of node

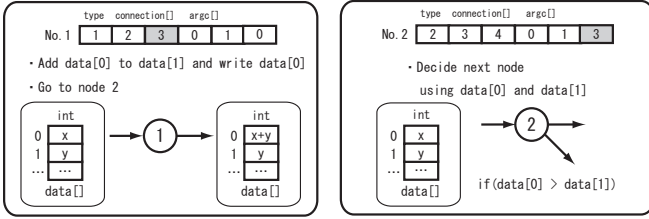


Fig. 2. Examples of node in GRAPE.

mapping method is similar to Cartesian Genetic Programming (CGP). The GRAPE program is encoded in the form of a linear string of integers. The genotype is an integer string which denotes a list of node types, connections and arguments. The connections of nodes are arbitrary, that is different from CGP. The length of the genotype is fixed and equals to $N * (n_c + n_a + 1) + 1$, where N is the number of nodes, n_c is the maximum number of connections and n_a is the maximum number of arguments.

C. Genetic Operators of GRAPE

To obtain the optimum structure of GRAPE, an evolutionary method is adopted. The genotype of GRAPE is a linear string of integers. Therefore, GRAPE is able to use a usual Genetic Algorithm (GA). In this paper we use uniform crossover and mutation as the genetic operators. The uniform crossover operator effects two individuals, as follows:

- Select several genes randomly according to the crossover rate P_c for each gene.
- The selected genes are swapped between two parents, and generate offspring.

The mutation operator effects one individual, as follows:

- Select several genes randomly according to the mutation rate P_m for each gene.
- The selected genes are randomly changed.

D. Features of GRAPE

GRAPE has different representation from PDGP, CGP and Linear-Graph GP. These methods have some restriction of connections (e.g. restrict loops and allow only feed-forward connectivity). The representation of GRAPE is arbitrary directed graph of nodes. PADO is one of the similar methods to GRAPE. PADO has stack memory and index memory, and the execution of PADO is carried out from the start node to the end node in the network. GRAPE is difference from PADO in the fact that GRAPE handles multiple data types using the *data set* for each type and adopts genotype-phenotype mapping.

The features of GRAPE are summarized as follows:

- Arbitrary directed graph structures.
- Handle multiple data types using the *data set*.
- Genotype of integer string.

IV. EXPERIMENTS AND RESULTS

In this section, we apply GRAPE to the problem of automatic construction of sorting algorithm.

TABLE I

THE PARAMETERS OF GRAPE ALGORITHM.

Parameter	Value
The number of evaluations	5,000,000
Population size	500
Crossover rate P_c	0.9
Mutation rate P_m	0.02
The number of nodes	10, 30, 50
Execution step limits	3,000

A. Settings of Experiments

In this problem we seek to evolve an implementation of the sorting algorithm. We provide a list of integers as input. A correct program returns a sorting input list, of any length (e.g. input: (2 1 7 5 1), output: (1 1 2 5 7)). The training data set is 30 random lists whose lengths are between 10 and 20. Elements are randomly chosen from the range of [0, 255].

The fitness function used in this experiment is given in (1). The range of this fitness function is [0.0, 1.0]. The higher the numerical value indicates the better performance.

$$fitness = 1.0 - \frac{\sum_{i=1}^n \frac{\sum_{j=0}^l (1 - \frac{1}{2^{d_{ij}}})}{l_i}}{n} \quad (1)$$

where d_{ij} is the distance between the correct position and the return value position for the training data i for the element j . l_i is the length of the list for the training data i and n is the size of the training data set. If the fitness in (1) is reached 1.0, the fitness function is calculated using (2).

$$fitness = 1.0 + \frac{1}{S_{exe}} \quad (2)$$

where S_{exe} is the total number of execution steps of the generated program. This fitness function means the less execution step is the better solution.

We used Minimal Generation Gap (MGG) as the generation alternation model. The MGG model [26]–[28] is a steady state model proposed by Satoh et al. The MGG model has a desirable convergence property maintaining the diversity of the population, and shows higher performance than the other conventional models in a wide range of applications. We used the MGG model in these experiments as follows:

- 1) Set generation counter $t = 0$. Generate N individuals randomly as the initial population $P(t)$.
- 2) Select a set of two parents M by random sampling from the population $P(t)$.
- 3) Generate a set of m offspring C by applying the crossover and the mutation operation to M .
- 4) Select two individuals from set $M + C$. One is the elite individual and the other is the individual by the roulette-wheel selection. Then replace M with the two individuals in population $P(t)$ to get population $P(t+1)$.

TABLE II
GRAPE NODE FUNCTIONS FOR THE EXPERIMENT.

Name	# Connections	# Args.	Argument(s)	Description
+	1	3	x, y, z	Use integer data type. Add data[x] to data[y] and substitute for data[z].
-	1	3	x, y, z	Use integer data type. Subtract data[x] from data[y] and substitute for data[z].
*	1	3	x, y, z	Use integer data type. Multiply data[x] by data[y] and substitute for data[z].
/	1	3	x, y, z	Use integer data type. Divide data[x] by data[y] and substitute for data[z].
=	2	2	x, y	Use integer data type. If data[x] is equal data[y] connection 1 is chosen else connection 2 is chosen.
>	2	2	x, y	Use integer data type. If data[x] is greater than data[y] connection 1 is chosen else connection 2 is chosen.
<	2	2	x, y	Use integer data type. If data[x] is less than data[y] connection 1 is chosen else connection 2 is chosen.
SwapList	1	2	x, y	Use integer type and a list data. Swap list[data[x]] for list[data[y]].
EqualList	2	2	x, y	Use integer type and a list data. If list[data[x]] is equal list[data[y]] connection 1 is chosen else connection 2 is chosen.
GreaterList	2	2	x, y	Use integer type and a list data. If list[data[x]] is greater than list[data[y]] connection 1 is chosen else connection 2 is chosen.
LessList	2	2	x, y	Use integer type and a list data. If list[data[x]] is less than list[data[y]] connection 1 is chosen else connection 2 is chosen.
OutputList	0	0	-	Output a list data and then the program halts.

5) Stop if a certain specified condition is satisfied, otherwise set $t = t + 1$ and go to step 2.

In these experiments we used $m = 50$.

The parameters of GRAPE are given in Table I. The number of nodes was 10, 30, and 50. In order to avoid the problem caused by non-terminating structures we limited the execution step to 3,000. When a program reaches the execution limit, the individual is assigned the fitness 0.0. The node functions are shown in Table II. We prepare simple node functions, *arithmetic functions*, *swap the elements of list* and *compare the elements of list*. We do not prepare special functions such as *iteration functions*.

In the experiment a list of integers and integer data type are used, and the size of integer data in GRAPE is 15. Initially, we set the size of input list (the list length) on the $data[0]$ to $data[4]$, constant value 0 on the $data[5]$ to $data[9]$ and constant value 1 on the $data[10]$ to $data[14]$.

B. Results

Results are given for 100 different runs with the same parameter set. Fig. 3 shows transition of average fitness of 100 trials. Fig. 4 shows transition of success rate. The success rate is computed as:

$$Success\ rate = \frac{Number\ of\ successful\ runs}{Total\ number\ of\ runs}. \quad (3)$$

We apply the elite individual generated by GRAPE to the test data set for each run. The test set is 50 random lists. We use the length of the list between 10 and 50 as the test set. Elements are randomly chosen from the range of [0, 255]. The results are shown in Table III.

Fig. 5 is an example of obtained structure which is general sorting algorithm. This GRAPE program has two loop structures and sorts any sequences of numbers.

C. Discussion

GRAPE successfully generates the sorting algorithm automatically, and the obtained structure is unique and solves completely the problems. In Fig. 4 the success rate rises greatly after 2,000,000 fitness evaluations, and finally reaches 67% when the number of nodes is 50. At the beginning of the evolution, the success rate is 0% until about 2,000,000 fitness evaluations. It shows that the evolution of sorting algorithm is very difficult. Although the success is 0% at the beginning of the evolution, the fitness rises little by little in Fig. 3. Therefore, it shows that the evolutionary method is functionally effective.

Table III provides the success rate for the training set and the test set. The results shows high success rate (“node30” is 62% , “node50” is 67%) for the training set. It shows that GRAPE is a powerful Automatic Programming technique. For the test set, “node30” shows best performance (39%), and “node50” is 28% . We can consider that these programs which success for the test set are general sorting algorithm. When the number of nodes is 10, the result is not good (the success rate is 3% for training set and test set). Because it is hard to construct the sorting programs automatically for the training set using “node10”. However, all of the successful individuals (programs) for the training set (3%) also succeed with the test set.

In the experiment, we use (2) as the fitness function. This fitness function means the less execution step is the better solution. For this reason, GRAPE tends to construct the exclusive sorting algorithm for the training set. Thus, the fitness function and selection of the training set are very important for GRAPE.

Fig. 5 is an example of obtained structure which is general sorting algorithm. This GRAPE program has two loop structures and sorts any sequences of numbers. The

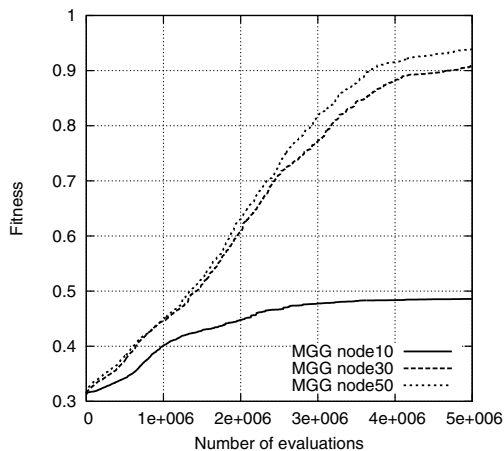


Fig. 3. Average performance over 100 trials by GRAPE with the number of node 10, 30 and 50.

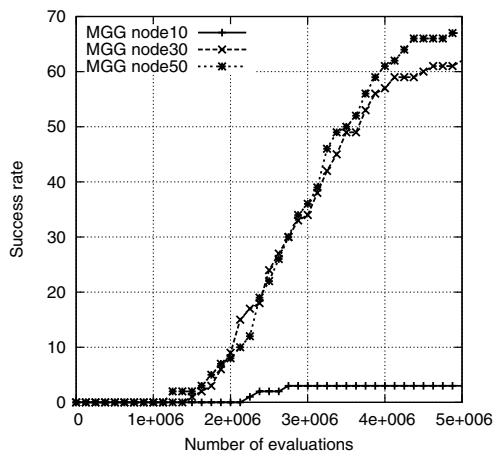


Fig. 4. Transition of success rate over 100 trials by GRAPE with the number of node 10, 30 and 50.

obtained structure briefly translates to the Fig. 6 like C language form. This algorithm is similar to the *selection sort*.

The representation of GRAPE is graph structure, therefore it can easily represent loops and branches. The programs of sorting algorithms can be represented using recursion. A lot of elegant and efficient sorting algorithms (e.g. merge sort, quick sort) are best expressed as recursive functions. Although we have not prepared recursion function in this paper, GRAPE has constructed the best programs using the branches, the loops and the multiple data types. If we introduce recursion functions or modularity mechanisms (like ADFs; Automatically Defined Function [3]) to GRAPE, the performance of GRAPE may improve.

V. CONCLUSIONS AND FUTURE WORKS

In this paper, we apply Graph Structured Program Evolution (GRAPE) to evolution of general sorting algorithm. The representation of GRAPE is graph structure. Each program is constructed as an arbitrary directed graph of nodes and *data set*. The *data set* flows the directed graph and is processed at

TABLE III

THE SUCCESS RATE FOR THE TRAINING SET AND THE TEST SET.

	Training set	Test set
MGG node10	3%	3%
MGG node30	62%	39%
MGG node50	67%	28%

```
List[] (input list)
data[0]=data[4]=ListLength
data[8]=data[9]=0
data[11]=data[13]=data[14]=1
```

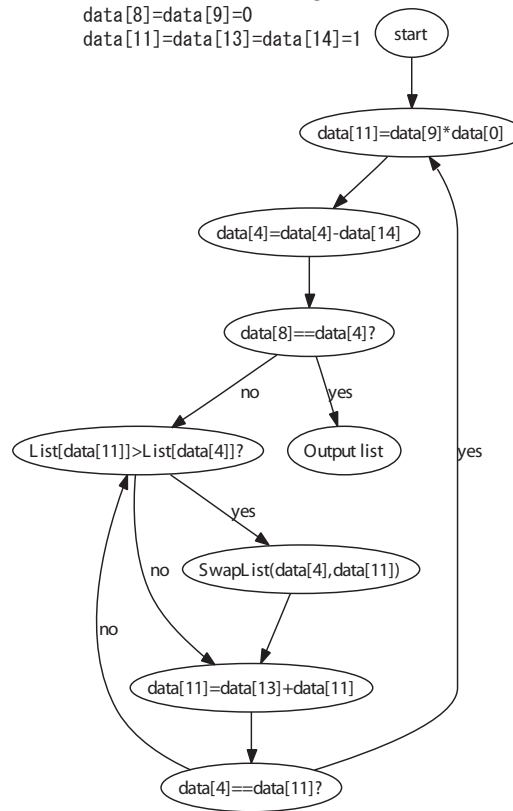


Fig. 5. An example of obtained structure by GRAPE. This GRAPE program is equivalent to general sorting algorithm.

each node. GRAPE adopts the genotype-phenotype mapping. The genotype is an integer string which denotes a list of node types, connections and arguments.

We confirm that the general sorting algorithm is obtained by GRAPE. The obtained structures (programs) of GRAPE are unique and have loops and branches. GRAPE obtain high success rate for the training and the test data set. As a result we showed that the evolutionary method is functionally effective. We show that GRAPE is a powerful Automatic Programming technique.

As for further research topics, we will introduce recursion functions or modularity mechanisms to obtain more elegant and efficient sorting algorithm or to solve more complex problems. Moreover, we will plan to apply GRAPE to the problems which are more large scale and require more complex structure, for example, control of autonomous agent, multi agent system and signal processing.

```

data[4]=ListLength;
data[11]=1;
while(1) {
    data[11] = 0;
    data[4] = data[4] - 1;
    if(data[4] == 0) {
        return List[];
    }
    else {
        do {
            if(List[data[11]] > List[data[4]])
                SwapList(data[4], data[11]);
            data[11] = data[11] + 1;
        } while(data[4] == data[11]);
    }
}
}

```

Fig. 6. This C language like program is translated from the GRAPE program in Fig. 5.

REFERENCES

- [1] Shinichi Shirakawa, Shintaro Ogino, and Tomoharu Nagao. Graph Structured Program Evolution. In *Proceedings of Genetic and Evolutionary Computation Conference (GECCO'07)*, volume 2, pages 1686–1693, 2007.
- [2] John R. Koza. *Genetic Programming: On the Programming of Computers by Means of Natural Selection*. MIT Press, Cambridge, MA, USA, 1992.
- [3] John R. Koza. *Genetic Programming II: Automatic Discovery of Reusable Programs*. MIT Press, Cambridge, MA, USA, 1994.
- [4] Astro Teller. Learning mental models. In *Proceedings of the Fifth Workshop on Neural Networks: An International Conference on Computational Intelligence: Neural Networks, Fuzzy Systems, Evolutionary Programming, and Virtual Reality*, 1993.
- [5] Astro Teller. Turing completeness in the language of genetic programming with indexed memory. In *Proceedings of the 1994 IEEE World Congress on Computational Intelligence*, volume 1, pages 136–141, Orlando, Florida, USA, 27-29 June 1994. IEEE Press.
- [6] Markus Brameier and Wolfgang Banzhaf. A comparison of linear genetic programming and neural networks in medical data mining. *IEEE Transactions on Evolutionary Computation*, 5(1):17–26, February 2001.
- [7] Markus Brameier and Wolfgang Banzhaf. *Linear Genetic Programming*. Number XVI in Genetic and Evolutionary Computation. Springer, 2007.
- [8] Michael O'Neill and Conor Ryan. Grammatical evolution. *IEEE Transactions on Evolutionary Computation*, 5(4):349–358, August 2001.
- [9] Michael O'Neill and Conor Ryan. *Grammatical Evolution: Evolutionary Automatic Programming in a Arbitrary Language*, volume 4 of *Genetic programming*. Kluwer Academic Publishers, 2003.
- [10] Astro Teller and Manuela Veloso. Algorithm evolution for face recognition: What makes a picture difficult. In *International Conference on Evolutionary Computation*, pages 608–613, Perth, Australia, 1–3 December 1995. IEEE Press.
- [11] Astro Teller and Manuela Veloso. Program evolution for data mining. *The International Journal of Expert Systems*, 8(3):216–236, 1995.
- [12] Astro Teller and Manuela Veloso. PADO: A new learning architecture for object recognition. In Katsushi Ikeuchi and Manuela Veloso, editors, *Symbolic Visual Learning*, pages 81–116. Oxford University Press, 1996.
- [13] Riccardo Poli. Evolution of graph-like programs with parallel distributed genetic programming. In *Genetic Algorithms: Proceedings of the Seventh International Conference*, pages 346–353, Michigan State University, East Lansing, MI, USA, 19-23 July 1997. Morgan Kaufmann.
- [14] Julian F. Miller and Stephen L. Smith. Redundancy and computational efficiency in cartesian genetic programming. *IEEE Transactions on Evolutionary Computation*, 10(2):167–174, April 2006.
- [15] Julian F. Miller and Peter Thomson. Cartesian genetic programming. In *Genetic Programming, Proceedings of EuroGP'2000*, volume 1802 of *LNCS*, pages 121–132, Edinburgh, 15-16 April 2000. Springer-Verlag.
- [16] Wolfgang Kantschik and Wolfgang Banzhaf. Linear-graph GP—A new GP structure. In *Genetic Programming, Proceedings of the 5th European Conference, EuroGP 2002*, volume 2278 of *LNCS*, pages 83–92, Kinsale, Ireland, 3-5 April 2002. Springer-Verlag.
- [17] Wolfgang Kantschik and Wolfgang Banzhaf. Linear-tree GP and its comparison with other GP structures. In *Genetic Programming, Proceedings of EuroGP'2001*, volume 2038 of *LNCS*, pages 302–312, Lake Como, Italy, 18-20 April 2001. Springer-Verlag.
- [18] Toru Eguchi, Kotaro Hirasawa, Jinglu Hu, and Noriko Ota. A study of evolutionary multiagent models based on symbiosis. *IEEE Transactions on Systems, Man and Cybernetics Part B*, 36(1):179–193, 2006.
- [19] Hironobu Katagiri, Kotaro Hirasawa, Jinglu Hu, and Junichi Murata. Network structure oriented evolutionary model-genetic network programming-and its comparison with genetic programming. In *2001 Genetic and Evolutionary Computation Conference Late Breaking Papers*, pages 219–226, San Francisco, California, USA, 9-11 July 2001.
- [20] Lee Spector. Autoconstructive evolution: Push, pushGP, and pushpop. In *Proceedings of the Genetic and Evolutionary Computation Conference (GECCO-2001)*, pages 137–146, San Francisco, California, USA, 7-11 July 2001. Morgan Kaufmann.
- [21] Lee Spector, Jon Klein, and Maarten Keijzer. The push3 execution stack and the evolution of control. In *GECCO 2005: Proceedings of the 2005 conference on Genetic and evolutionary computation*, volume 2, pages 1689–1696, Washington DC, USA, 25-29 June 2005. ACM Press.
- [22] Lee Spector and Alan Robinson. Genetic programming and autoconstructive evolution with the push programming language. *Genetic Programming and Evolvable Machines*, 3(1):7–40, March 2002.
- [23] Simon Lucas. Exploiting reflection in object oriented genetic programming. In *Genetic Programming 7th European Conference, EuroGP 2004, Proceedings*, volume 3003 of *LNCS*, pages 369–378, Coimbra, Portugal, 5-7 April 2004. Springer-Verlag.
- [24] Alexandros Agapitos and Simon M. Lucas. Learning recursive functions with object oriented genetic programming. In *Proceedings of the 9th European Conference on Genetic Programming*, volume 3905 of *Lecture Notes in Computer Science*, pages 166–177, Budapest, Hungary, 10 - 12 April 2006. Springer.
- [25] Alexandros Agapitos and Simon M. Lucas. Evolving efficient recursive sorting algorithms. In *Proceedings of the 2006 IEEE Congress on Evolutionary Computation*, pages 9227–9234, Vancouver, 6-21 July 2006. IEEE Press.
- [26] H. Satoh, M. Yamamura, and S. Kobayashi. Minimal generation gap model for considering both exploration and exploitations. In *Proceedings of the IIZUKA'96*, pages 494–497, 1996.
- [27] H. Kita, I. Ono, and S. Kobayashi. Multi-parental extension of the unimodal normal distribution crossover for real-coded genetic algorithms. In *Proceedings of the 1999 Congress on Evolutionary Computation (CEC99)*, volume 2, pages 1581–1587, 1999.
- [28] S. Tsutsui, M. Yamamura, and T. Higuchi. Multi-parent recombination with simplex crossover in real coded genetic algorithms. In *Proceedings of Genetic and Evolutionary Computation Conference (GECCO'99)*, pages 657–664, 1999.