# Genetic Programming based Constructive Algorithm with Penalty Function for Hardware/Software Cosynthesis of Embedded Systems

Adam Górski and Maciej Ogorzałek

Department of Information Technologies, Jagiellonian University in Cracow, Prof. Stanisława Łojasiewicza 11, Cracow, Poland

- Keywords: Artificial Intelligence, Embedded Systems, Genetic Programming, Penalty Function, Hardware/Software Co-Synthesis.
- Abstract: In this work we present a constructive genetic programming method with penalty function for hw/sw cosynthesis of embedded systems. The genotype is a tree which contains in its nodes system construction options. Unlike existing solutions in this approach individuals which violate time constrains are investigated during evolution process. Therefore the algorithm is even more able to escape local minima of optimizing parameters.

# **1 INTRODUCTION**

Artificial intelligence (AI) (Shastri et al. 2021) is widely used in computer science. Method of AI like: evolution algorithms and multi-agent systems (Jin et al. 2021) were applied to solve many problems. One of those are optimization problems like: hardware design (Dick et al. 1998), traveling salesman problem (Lust and Teghem 2010) Multi-skill resourceconstrained project scheduling problem (Lin et al. 2020), and many others.

Embedded system (Martins et al. 2020) is a computer system consisted of hardware elements optimized to execute appropriate tasks. According to De Michelli and Gupta (De Michelli and Gupta 1997) embedded system design can be divided onto three phases: modelling, verification and implementation. In (Górski and Ogorzałek 2016) authors propose another phase - assignment of unexpected tasks. Hardware/software co-syntesis (Yen and Wolf 1995, Oh and Ha 2002) is a process of concurrent generation of an architecture of embedded system and its software. The goal of the process is to find optimal architecture with optimal task assignment. The architecture can be composed of two groups of resources: Processing Elements (PEs) which are responsible for tasks execution and Communication Links (CLs) which provide communication between connected PEs.. There are two kinds of PEs: Programmable Processors (PPs) able to execute more

than one task and Hardware Cores (HCs) which are specialized to execute only one task. Most of cosynthesis algorithms can be divided on two groups constructive (Deniziak and Górski 2008, Górski and Ogorzałek 2014a, Srinivasan and Jha 1995) and iterative improvement (Oh and Ha 2002, Górski and Ogorzałek 2021). Constructive algorithms build system step by step making decisions separately for each part. They have low complexity but are able to stop in local minima of optimizing parameters. Iterative improvement algorithms start from suboptimal solution and by local changes try to improve the quality of the system. The initial solution in such methods is usually the fastest implementation of all the tasks. In such a solution every task is usually executed by another HC. Genetic algorithms (Conner et. al 2005) were also applied to co-synthesis problem. They can provide an acceptable result in reasonable time and are able to escape local minima of optimizing parameters. However the disadvantage of those methods is that obtained results are sensitive to change of genetic parameters. Very good results were obtained using genetic programming (GP) (Górski and Ogorzałek 2021). In (Deniziak and Górski 2008) authors propose a constructive GP method. One of the biggest disadvantages of the method was that every generated individual could not violate time constrains. Therefore the algorithm had smaller chance to escape local minima of optimizing parameters. The time of computation was also

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increasing. In (Górski and Ogorzałek 2014b) an iterative improvement algorithm for co-synthesis problem was proposed. The algorithm builds initial population by starting from the fastest implementation. The next generations are obtained using genetic operators: mutations, cloning and crossover. However this method also do not use a penalty function. Therefore only valid individuals are used to generate the final solution. In this paper we present a Genetic Programming method to hardware/software co-synthesis of distributed embedded system which uses the penalty function during generation of the results. Unlike other algorithms our method during evolution process investigates also individuals which violate time constrains. Therefore it is a greater chance that algorithm will not stop in local minima of optimizing parameters.

The paper is organized as follows: section 2 describes genetic programming – types and application. In section 3 the representation of embedded system is presented. Section 4 includes the description of the algorithm. Experimental results are given in section 5. The last section summarize the paper and indicates the direction of future work.

### 2 GENETIC PROGRAMMING

Genetic Programming (Suganuma et al. 2020) is an extension of Genetic Algorithm (Nayak and Panda, 2020). The main difference between those methodologies is that in GP genotype is a tree. The tree, in its nodes, includes functions.

In linear genetic programming (Zhang et al. 2020) the tree is represented in linear form.

In cartesian genetic programming (Miller, 2011) the genotype is a graph. Thus the genetic operators needed to be modified.

Multigene Genetic Programming (Riahi-Madvar et al. 2019) evolve strings of genes. However every gene is a tree in which nodes are functions.

Developmental Genetic Programming (DGP) (Koza et al. 1997) starts with embryo. Embryo is the first node in genotype tree. Every other node contains function which modifies the embryo. This type of GP is often used for hardware design (Deniziak and Górski 2008, Górski and Ogorzałek 2017, Górski and Ogorzałek 2021). This kind of GP was also used in this paper.

#### **3 PRELIMINARIES**

Embedded system is represented as a task graph G = (T, E). This representation is one of the most popular. In the representation each node  $T_i$  in a graph is a task executed by the system. Each edge describes an amount of data  $(d_{ij})$  that need to be transferred between two connected tasks  $T_i$  and  $T_j$ . On figure 1 the example of task graph is presented.



Figure 1: Example of task graph.

The example consists of eight tasks: T0, T1, T2, T3, T4, T5, T6 and T7. Tasks T1, T2 and T4, T5, T6, T7 are parallel. Tasks T1 and T2 can start their execution after finishing of T0 execution. Tasks T3 and T5 can be started only if T2 is finished. Tasks T6 and T7 can be executed after T3.

The transmission time  $t_{i,j}$  is dependent on the bandwidth  $(b_u)$  of a communication link used to connected PEs. It is described by the following formula:

$$t_{i,j} = \frac{d_{i,j}}{b_{i,j}} \tag{1}$$

The execution of a task is characterized by the cost (c) and time (t). We assume that a database which include all times and cost of execution for every tasks on every PEs is given. The database also contains cost of each PE and cost of connection PEs using each CLs. In table 1 the example of database for a graph from figure 1 is presented. In the example there are four kinds of possible PEs. Two of them are PPs and two of them are HCs. The cost (C) of PP1 is 100, the cost of PP2 is 300. The cost of HCs is added to a cost of tasks' execution. PEs can be connected using one CL. The bandwidth of the CL is 13. Cost of connection of PP1 to CL1 is equal to 6. The cost of

connection of PP2 is equal to 3. The costs of connection of every HCs to CL1 are equal to 25. If one or more PEs execute more than one task, then those tasks need to be scheduled. We decided to use list scheduling.

Table 1: Example of resource database.

Task	PP1 C=100		PP2 C=300		HC1		HC2	
	t	с	t	с	t	с	t	с
T0	12	6	15	4	10	200	3	160
T1	11	8	23	10	7	90	12	80
T2	30	2	21	8	6	100	5	140
T3	18	4	20	3	4	150	1	350
T4	26	10	22	5	3	200	2	240
T5	22	5	36	12	6	90	8	70
T6	34	11	44	9	11	88	13	80
T7	45	15	30	16	9	180	10	190
CL1, b=13	c=6		c=3		c=25			

To estimate the quality of individuals we use the fitness function (F). We decided that during evolution process, to avoid stopping in local minima of optimizing parameters, every generated individuals can be used. Therefore every individuals which violate the time constrains must be given a penalty. The penalty is depended on the violation time. The function is depended on time of the solution (t), its  $\cot(C_s)$  and given penalty (PF). It is described by the following formula:

$$F = k * C_s + l * t + p * P_F$$
(2)

Parameters k, l and p are given by the designer. There are no dependencies between the parameters k, l and p. The time of the solution is a time after which the last task finishes its execution. The cost of embedded system can be calculated as:

$$C_{s} = \sum_{z=1}^{m} C_{PE_{z}} + \sum_{y=1}^{n} c_{y} + \sum_{z=1}^{v} \sum_{y=1}^{P_{u}} c_{CL_{u}, PC_{y}}$$
(3)

Where n is a number of tasks in a task graph, m is a number of PP in solution, u is a number of CLs connected to v PEs.

Penalty can be calculated as follows:

$$P_F = (t - t_{\max}) \tag{4}$$

where  $t_{max}$  is a time constrain and t is a time of execution of all the tasks for investigated solution.

## **4** THE ALGORITHM

The algorithm which is presented in this paper is constructive method. Therefore design decisions are made for each task separately. The genotype is a tree. To be sure that every task is executed the genotype is a spanning tree of given task graph. The first node in the tree is embryo. Embryo is a random implementation of the first task. The rest of nodes contains genes. In our solution genes are system construction options. The options are presented in table 2.

Table 2: Options for building system.

Step	Option	Probability		
PE	a. The fastest implementation of the task	0.2		
	b. The cheapest implementation of the task	0.2		
	c. min (t*c)	0.2		
/	d. The same as predecessor	0.2		
	e. The rarest used PP	0.2		
CL	a. The fastest CL	0.2		
	b. The cheapest CL	0.2		
LOGY	c. The rarest used	0.3		
	d. The same as predecessor	0.3		
Task scheduling	list scheduling			

Unlike others constructive methods (Deniziak and Górski 2008) our algorithm do not separate options for used PEs and options for allocating new ones. The algorithm independently makes a decision if it is necessary to allocate a new PE.

The number of individuals in each population is described by the following formula:

$$\Pi = \alpha * n * e \tag{5}$$

where n is number of tasks (nodes in the task graph), e is possible number of embryos.

After generating an initial population the algorithm builds new individuals using genetic operators: selection, crossover, mutation and cloning. In this paper we decided to use rank selection. Every generated individuals are ranked by the minimum value of fitness function. The individuals have probability (P) depended on the position on the rank list (r):

$$P = \frac{\Pi - r}{\Pi} \tag{6}$$

The probability describes the chance that every individual will be used during evolution process. Individuals with lower value of fitness function have greater probability of being selected, however the solutions with higher value are not necessary rejected.

Mutation randomly selects (using selection operator)  $\Omega$  individuals:

$$\Omega = \delta^* \Pi \tag{7}$$

For each individual one node is selected randomly. Mutation substitutes the option in selected node on another using available options (from table 2).

Crossover chooses  $\Psi$  individuals. Selected individuals are connected in pairs. Then a crossing point is selected randomly. The point is the same for both individuals in each pair. The operator substitutes subtrees between the solutions. The number of individuals created by crossover is equal to the following formula:

$$\Psi = \gamma * \Pi \tag{8}$$

Cloning operator copies  $\Phi$  individuals to the next population. formula:

$$\Phi = \beta * \Pi \tag{9}$$

To have the same number of individuals in each population must be satisfied the condition:

$$\beta + \gamma + \delta = 1 \tag{10}$$

The algorithm stops if in  $\varepsilon$  next generations better individual was not found.

### **5 FIRST RESULTS**

To check the quantity of presented algorithm we decided to use benchmarks with 10, 20 and 30 nodes. The results are presented in table 3 below. For each graph 10 runs were made and the best obtained results were put in table 3. They were compared with Genetic Programming algorithm proposed by Deniziak and Górski (Deniziak and Górski 2008) which is also a developmental genetic programming constructive method. The parameters were set as follows:  $\alpha$ =2,  $\beta$ =0.2,  $\gamma$ = 0.7,  $\delta$ =0.1,  $\epsilon$ =3, k=8, l=1, p=4.

Table 3: The results.

Algorithm	Graph	t	с	Gen.	F
DGP08	10	479	1667	3	13815
	20	970	3575	9	29570
	30	1273	6323	28	51857
DGP2021	10	531	2095	15	17322
	20	937	2749	29	22929
	30	937	6035	44	49217

The time constrains for each graph were set as follows: graph with 10 nodes - 500, graph with 20 nodes -1000, graph with 30 nodes -1300. As it can be observed in table 3 the results obtained by the presented algorithm are much better in most of cases than obtained by DGP08. Only for a graph with 10 nodes the best solution obtained by presented algorithm has fitness function equal to 17322, meanwhile the best solution obtained by DGP08 has fitness function equal to 13815. For graphs with 20 nodes presented algorithm generated results with fitness function equals to 22929. For that graph presented algorithm generated individual with better value of time (937) and cost (2749) than DGP08 (cost equal to 3575, and time equal to 970). What is worth to underline using the presented approach during the evolution process despite the fact that in some of the generations best individuals violated time constrain, the final solution has time under maximum value. The same situation can be observed for graph with 30 nodes. The time of a final solution obtained by presented algorithm (937) does not violate time constrains. It is also lower than time obtained using DGP08 (1273). The solution generated by presented approach has also lower value of cost (6035) than result obtained by DGP08 (6323). This suggest that algorithm presented in this paper has greater ability to escape local minima of optimizing parameters.

## 6 CONCLUSIONS AND FUTURE WORK

In this paper a constructive genetic programming method for hardware/software cosynthesis of distributed embedded system was presented. The method uses a penalty function. Some of the individuals in each generation can violate time constrains. Therefore those individuals can be used to construct next generations. Thus the algorithm is more able to escape local minima of optimizing parameters. The first experimental results indicates bigger effectiveness of presented algorithm than DGP08. Only in one case better results were generated by DGP08 algorithm – for graph with 10 nodes. However more experiments must be performed using presented algorithm to be sure why in such a case the results were worse. Maybe with different value of genetic parameters or another values of probability of system construction options results could be better even for graph with 10 nodes. Therefore more experiments are needed to find the best values of genetic operators. In future work we also plan to check another combination of system construction options and different genetic operators and their impact on final results.

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