Adapting Evolutionary Algorithms to the Concurrent Functional Language Erlang

J. Albert Cruz
Centro de Arquitecturas Empresariales
Universidad de Ciencias Informáticas
La Habana, Cuba
jalbert@uci.cu

Juan-J. Merelo, Antonio M. Mora and Paloma de las Cuevas Departamento de Arquitectura y Tecnología de Computadores Universidad de Granada Granada, Spain {jmerelo,amorag,paloma}@geneura.ugr.es

ABSTRACT

In this paper we describe how the usual sequential and procedural Evolutionary Algorithm is mapped to a concurrent and functional framework using the Erlang language. The design decisions, as well as some early results, are shown.

Categories and Subject Descriptors

D.1.3 [Software]: Programming Techniques—Concurrent Programming; D.2.8 [Software Engineering]: Performance measures; G.1.6 [Mathematics of Computing]: Numerical Analysis—Optimization

General Terms

Algorithms, Languages, Performance, Measurement

Keywords

Evolutionary Algorithms, Functional Languages, Concurrent Languages, Erlang, Algorithm Implementation

1. INTRODUCTION AND STATE OF THE ART

The Evolutionary Computation (EC) field is focused on the widespread use of implementation technologies such as C/C++, Fortran and Java. Getting out of that mainstream it is not normally seen as a land for scientific improvements. Erlang is a programming language with a lot of potential, it supports the functional and concurrent paradigms and it is been used in the scientific community [7].

In this paper we propose to apply this language (and its underlying paradigm) to Genetic Algorithms (GA) [3], which are general function optimizers that encode a potential solution to a specific problem in a simple data structure (e.g. a chromosome). There are only two components of them that are problem dependent: the solution encoding and the function that evaluates the quality of a solution, i.e., the fitness function. The rest of the algorithm does not depend on the problem and could be implemented following the best architecture and engineering practices.

Many of these practices have been proposed on the object oriented paradigm [6], but this is not the case of the func-

Copyright is held by the author/owner(s). GECCO'13 Companion, July 6–10, 2013, Amsterdam, The Netherlands. ACM 978-1-4503-1964-5/13/07. 2. EVOLUTIONARY ALGORITHM IN A CON-CURRENT FUNCTIONAL LANGUAGE

tional side. This work tries to show some possible areas of improvement on that sense by focusing on GAs as a domain

of application and describing how their principal traits can

be modeled by means of Erlang constructs. This will be

done in the next section.

A variety of programming patterns, i.e., paradigms, exist for implementing the algorithms models. GAs are characterized by an intensive use of strings (lists of some kind) for encoding genes and a population that evolves via operators that are applied to all or a part of it. Most modern languages can handle these data structures (and the operators needed to manipulate them).

There is a claim in modern software development for programming languages that help with concurrent programming and simplify coding practice. The functional programming language Erlang could be a good choice since it provides the actor pattern concept for concurrency and the functional paradigm for general modeling, design and coding of solutions. Actors are concurrent execution units which use asynchronous message passing for communication. They are implemented as processes in the Erlang's virtual machine and not as operating system (OS) threads, which means that they are very lightweight in creation and execution. The use of messages eliminates the sharing of state and also many of the typical problems of concurrent development, namely, supporting the emulation of the Object Oriented (OO) paradigm with its modeling facilities. Functional programming, the other main feature of the Erlang language, is defined by the use of functions in program composition and by using lists.

Genetic algorithms, as many other computational models, tend to be described in literature in an operational and imperative way. Their implementation in a functional language must follow a different path, structuring the algorithm model in less imperative and more declarative terms. We are going to use a parallel pool based evolutionary strategy (which was already used for instance in [4]) as use case.

The pool will be an execution entity (an actor acting like a server) that will own the population and also keep a track of the advance in the solution search. The clients, which are concurrent, will do the calculations and will join and leave the system at any time without consequences. Chromosomes

Message	Description
{configPool, NIM}	Initialization, the parame-
->	ter NIM is the initial con-
	figuration.
{requestWork, Pid,	Client requests for a popu-
Capacity} ->	lation to evolve.
{generationEnd,	One client reports its suc-
NewIndividuals,	cessfully end of calculation.
OldIndexes, Pid} ->	

Table 1: The messages that the pool accepts.

Message	Description	
initEvolution ->	Marks the beginning of the	
	processing.	
{evolve, P,	When the pool could assign	
NIndexes} ->	a subpopulation to process.	

Table 2: The messages a client is able to respond.

will be encoded as lists and the different parts of the GA algorithm will be implemented as Erlang functions.

An Erlang actor is implemented by a sequence of pairs pattern/expression defining each message that it could handle. It is close to the OO parlance and a way to organize the code. In this case we use one message per service that pool must provide; Table 1 presents this. Clients are modeled by actors. They are the units of evolution, with the main computation responsibilities; the Table 2 shows its interface.

The two previous components constitute the main features of the architecture of the algorithm; they are general-purpose and could be used for solve many differents optimization problems. In order to solve a particular situation, they must be *injected* by several functions and data structures which define chromosomes, fitness function, mutation operator, selection criteria and replacement policy. All these particularizations must be implemented in an Erlang source file and, in our case, configured in the *configBuilder* module.

The proposed design promotes a clear separation between architecture (the general, constant and paradigmatic foundation part) and problem encoding (the representation and criteria of solution finding) which is good for applying the library to solve others practical problems with GAs using a pool based approach. This design has been implemented and tested with a simple benchmark problem. This will be presented in the next section.

3. EXPERIMENT AND CONCLUSIONS

In this ongoing project we are testing the efficiency and simplicity of implementations of GAs by functional programming. The parallel models of GA are mapped to actors in the Erlang languages obtaining easily to understand architectures. All the code has been released as open source code at https://github.com/jalbertcruz/erlEA/.

The library was tested with *MaxOnes* problem. The chromosomes was 128 elements long, with an initial population of 256 individuals; Table 3 presents the results.

This results shows that in this case there it is not convenient to use many clients in order to obtain the solution, when the number of clients increase there is a lot of bad solutions evaluated.

With this concept test we are showing how simple is to

No. clients	Assignments	Time (seconds)
25	3040	5.339950
50	4301	7.891000
100	5630	11.671210

Table 3: Experiments results.

structure a parallel GA, and now we could proceed with more complex GA models, experiments and problems in order to explore the potential of the technology.

Acknowledgements

This work is supported by project TIN2011-28627-C04-02 (ANYSELF), awarded by the Spanish Mineco and P08-TIC-03903 awarded by the Andalusian Regional Government. It is also supported too by the PhD Program of the AUIP and by project 83, Campus CEI BioTIC.

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