

An Ecological Model of Genetic Algorithm for Interpretation of the Data

[Manjit Kaur]

Abstract— Ecological modeling covers a wide collection of methods and ideas of study. The variety of the suitable model is a part of any interpretation of ecological data. That variety is most often made randomly and the selection is frequently closely weighted by the knowledge of the programmer with the special model. Genetic Algorithm, which symbolizes a group of algorithms, intended to search the best alternative solution to a specified problem by using thoughts from natural science, biology, mainly genetics, offer another approach to model selection [3]. This paper presents a review of BEAGLE which is used for interpreting the ecological data behind the application of genetic algorithm.

Keywords— Evolutionary computation, GA: Genetic Algorithms, BEAGLE: Biological Evolutionary Algorithm Generating Logical Expressions.

I. Introduction

The most essential ecological models are concerned with the performance of a particular class, collection of classes and their deviation in time. These can be expressed as predictive functions. These models intend to recognize the biological and evolutionary methods that have formed a practical ecosystem and to discover the basic properties that create variety and arrangement in an idealized scheme. Evolutionary computation is a regulation that formulates use of values from normal development to develop alternative solutions to difficult computational problems [1, 10]. Their ability to handle huge search spaces in a competent mode, and their comparison to biological systems, like genotype (phenotype) mapping, variation and progression makes them perfect nominees for making ecological models.

Genetic Algorithm (GA) is a probabilistic technique encouraged by a conceptual model of our recent understanding of the normal procedure of evolution. Genetic algorithms offer a means of producing logical systems. This paper clarifies the principle underlying GAs and provides an explanation of the means by which systems are produced.

BEAGLE (Biological Evolutionary Algorithm Generating Logical Expressions) is a computer package. It is used for generating decision based system by stimulation from a database. It works on the principle of 'selection' whereby system that fit the information roughly are 'killed off' and swapped by 'mutations' of superior rules or by new generation created by 'mates' two best modified rules.

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II. Genetic Algorithm

A computer program is used to concern GAs to the review facts. GA explore from the population of possible solutions utilize present data to evaluate their value, and use genetic operators that imitate genetic methods in produce new offspring for advance testing [13]. The complete method is then repeated continuous until set of rules has been found. In the given Fig. 1, it described the model of Genetic Algorithm for interpretation of the data. There are four phases of genetic algorithm. These are:

- Population/Chromosomes/Decoded strings
- Evaluation/Fitness/Reproduction
- Selection/Mates/Manipulation
- Genetic Operators/Offspring/New generation

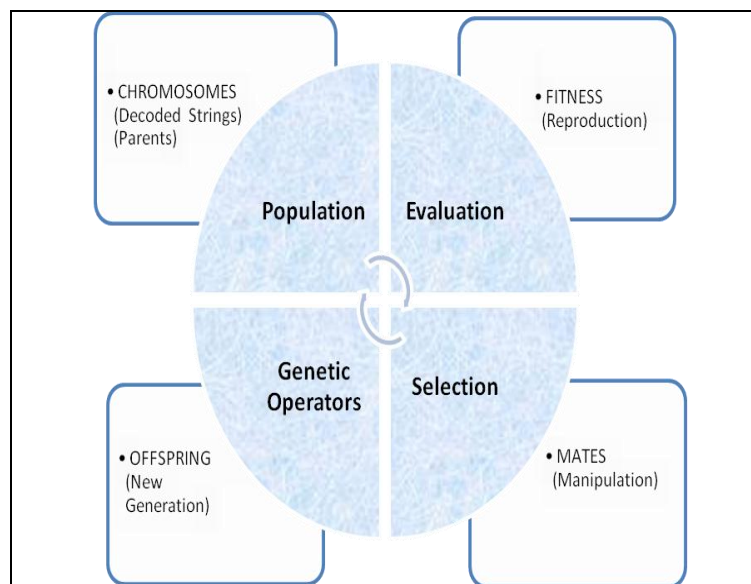


Figure 1. An Ecological Model of Genetic Algorithm

A. Classification of Genetic Algorithm

A GA follows biological evolutionary hypothesis to explain optimization problems. A GA includes a group of individual components and a group of biologically encourages operators defined over the population itself. As shown in the Fig. 2, GA is divided into different categories: Parallel and Sequential GAs. This paper presents a review of BEAGLE which is used for interpreting the ecological data behind the application of GA.

B. BEAGLE Algorithm

GAs offers another approach to model selection. They expand repetitively a bunch of rules which facilitate to clarify the relationships between attributes integrated in the data set [11, 23]. Several genetic algorithms are accessible but they all less or more have the identical features.

This paper describes BEAGLE (Biological Evolutionary Algorithm Generating Logical Expressions) which is a computer system generating decision based system by stimulation from a database. Simultaneously they achieve the assignment of classifying section into one of two or more sorting on the source of the values of a quantity of different factors describing each model [22]. HERB generates and/or changes the categorization rules which LEAF then uses, normally to predict group relationship for samples whose set is unknown.

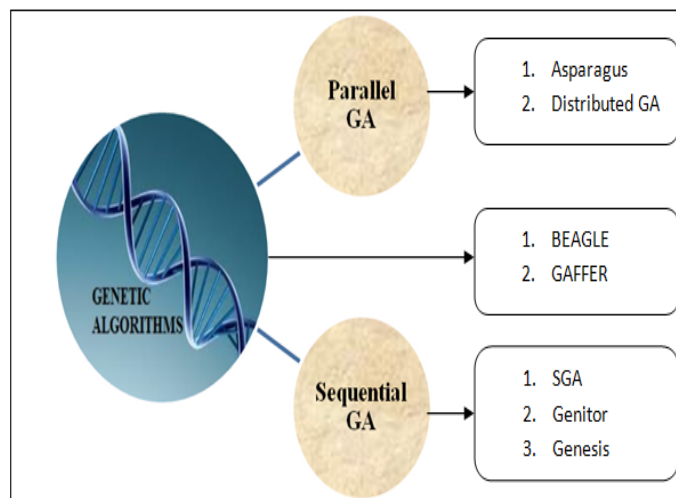


Figure 2. Classification of Genetic Algorithms

BEAGLE consists of six main components:

- SEED stands for Selectively Extracts Example Data. It permits data files to be study in easy formats, including ASCII files. It also executes the subsequent functions: (a) It divides the data into random sub-sets, and (b) It attaches lagging variables to time series.
- ROOT stands for Root-Oriented Optimization Tester. It permits the client to check one or more rules. If successful, these rules will then be used as a initial position for the successive parts. If no preliminary rules are obtainable ROOT will produce the requisite number of preliminary rules at random.
- HERB stands for Heuristic Evolutionary Rule Breeder. It produces original rules for the data file organized by SEED. It estimates all the obtainable rules beside the training data set and then reduces any rule that is not successful. It finally creates a few random modifies to several of the rules, clear out any errors initiated by the rules and executes suitable

syntactic manipulation to simplify the rules and build them more understandable.

- STEM stands for Signature Table Evaluation Module. It uses the rules establish by HERB to build a signature table, reexamines the training data and calculates the number of times each signature arises.
- LEAF stands for Logical Evaluator and Forecaster. It pertains the stimulate rules to an extra data set which has the similar structure as the preparation data. It uses the rules to categorize patterns from a catalog where the accurate relationship may be unknown. It is extreme simpler [22]. The client identifies how many rules to use: these are forever left planned by HERB with the top first. It can be producing: (a) a cataloging of all cases with expected set, and real set and score if known; (b) a review of the performance of each rule and all the rules equally.
- PLUM stands for Procedural Language Utilization Module. It decode the stimulate rules into a Pascal Procedure so that the rules can be exported into new software languages for realistic use.

C. An Example Genetic Algorithm Application

For example, to categorize the GA used in this paper a sequential verdict method that might have been more cost-effective; except the problem with algorithms which defer inequity (filter) net is their vulnerability to noise in the information. They work most excellent in circumstances where there can be no error. HERB, although infrequently if ever best possible, will approximately always rise with great usable.

III. Conclusion

BEAGLE is quiet at an instance phase, and can be significantly enhanced, determined database; with dynamism adaptable sets of individuals sent to consumers; redeployment of information when consumers are covering or not reacting. It would not be complex to execute; it would be to allocate floating-point calculation in addition to integers, however the interface with logical standards would have to be measured first. A solemn necessitate is to construct superior use of information provided by bad systems, rather than removing them and somewhat possibly restoring them later.

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References

- [1] R. Forsyth, "Evolutionary learning strategies", Machine Learning: Applications in Expert Systems and Information Retrieval, Ellis Horwood Ltd., USA, pp.79-95, 1986.

- [2] D. Goldberg, "Computer-Aided Pipeline Operations Using Genetic Algorithms and Rule learning", Part I: Genetic Algorithms in pipeline Optimization, Engineering with Computer 3, 1987.
- [3] D. E. Goldberg, "Genetic Algorithms in Search Optimization and Machine Learning", Addison-Wesley Publishing Company, Inc., Reading, Massachusetts, 1989.
- [4] W. Siedlecki and J. Sklansky, "A Note on Genetic Algorithms for Large-Scale Feature Selection", Pattern Recognition Letters, pp. 335-347, 1989.
- [5] J. Greffentette, "Genetic Algorithms Made Easy", 1991.
- [6] R. J. Hampo, "Genetic programming: A new paradigm for control and analysis", Proc. 3rd ASME Symposium on Transportation Systems, Anaheim, USA, pp.155-163, 1992.
- [7] J. R. Koza, "Genetic Programming", MIT Press, Cambridge, 1992.
- [8] C. M. Fonseca and P. J. Fleming, "Multi objective Genetic Algorithms", IEE Colloquium on Genetic Algorithms for Control Systems Engineering, 6/1-5, 1993.
- [9] W.F. Punch, E.D. Goodman, and Min Pei, "Further Research on Feature Selection and Classification Using Genetic Algorithms", Proc. Fifth Inter. Conf. Genetic Algorithms and their Applications (ICGA), pp. 557, 1993.
- [10] C. A. Grimes, "Application of genetic techniques to the planning of railway track maintenance", Proc. IEE Conf. on Genetic Algorithms in Engng. Systems: Inovations and Applications - GALEZIA'95, Sheffield, UK, pp.467-472, 1995.
- [11] E. Howard and N. Oakley, "Genetic programming as a means of assessing and reflecting chaos", Tech. Report FS-95-01, AAAI Press, pp.68- 72, 1995.
- [12] M. Pei, E.D. Goodman, W.F. Punch and Y. Ding, "Genetic Algorithms for Classification and Feature Extraction", presented at Classification Society Conference, June 1995.
- [13] F. Gruau and K. Quatramaran, "Cellular encoding for interactive evolutionary robotics", Cognitive science research paper, pp. 425, 1996.
- [14] K. S. Tang, K. F. Man and D. W. Gu, "Structured Genetic Algorithm for Robust H_∞ Control Systems Design", IEEE Transactions Industrial Electronics, 43(5) 575, 1996.
- [15] A. H. Watson, I. C. Parmee, "Identification of fluid systems using genetic programming", Proc. of the Second Online Workshop on Evolutionary Computation - WEC2, pp.45-48, 1996.
- [16] K. Rasheed and H. Hirsh, "Using case based learning to improve genetic algorithm based design optimization", In Proceedings of the 7th International Conference on Genetic Algorithms (ICGA97), Morgan Kaufmann, 1997.
- [17] C. M. Fonseca and P. J. Fleming, "Multi objective Optimization and Multiple Constraint Handling with Evolutionary Algorithms – Part 1: A Unified Formulation", IEEE Trans. Sys., Man. & Cyber., 28, 26, 1998.
- [18] P. Marenbach, "Using prior knowledge and obtaining process insight in data based modeling of bioprocesses" Sams, 31, pp. 39-59, 1998.
- [19] S. Chen and S. Smith, "Improving genetic algorithms by search space reduction (with applications to flow shop scheduling)", In GECCO-99: Proceedings of the Genetic and Evolutionary Computation Conference. Morgan Kaufmann, 1999.
- [20] T. D. R. Konig, "Improving genetic algorithms for protein folding simulations by systematic crossover", Biosystems, 50(1), April 1999.
- [21] T. Back, D. B. Fogel, and Z. Michalewicz, "Evolutionary Computation 1: Basic Algorithms and Operators", Institute of Physics Publishing, Bristol, UK, 2000.
- [22] C. Gagn'e and M. Parizeau, "Open BEAGLE: A new versatile C++ framework for evolutionary computation", In Proceeding of GECCO 2002, Late-Breaking Papers, New York, NY, USA, 2002.
- [23] K. E. Nygard and C. H. Yang, "Genetic Algorithm for the Travelling Salesman Problem With Time Window", In Computer Science and Operations Research: New Development in their Interface, O. Balci, R. Sharda and S. A. Zenios Eds, Pergamon Press, pp. 411-423.

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I Manjit Kaur working as Assistant Professor in Lovely Professional University, Phagwara, Punjab, India. I have done Mater of Technology in Computer Science Engineering in 2011. I have more than 4.8 years experience in teaching area. My research interests areas: Data Mining, Data Warehousing, Neural Network and Image Processing.