

IEE/CSE 598: Bio-Inspired AI and Optimization

Mini-Project 1: Implementing a Simple GA with a Base-10 Genotype (30 points)

Description. This first mini-project is a small challenge meant to help solidify your understanding of how to implement a simple GA; it also provides an example of the kind of deliverable that is expected in future mini-projects. This assignment follows from class discussions of Chapter 4 from [Bozorg-Haddad et al. \(2017\)](#) and Chapter 14 from [Passino \(2005\)](#), both of which are available for download on the Canvas site under “Unit 1” in “Modules.”

You may discuss approaches for solving this problem with other students in your class. However, ultimately, this activity is to be completed **individually** without the aid of other students in the class. **Please upload your submission to Canvas by the due date.** Any submissions after the due date will be considered late and may be penalized accordingly.

Submit a **single DOC, DOCX, or PDF file** with solutions to the following questions. **Include any code** you used **in an appendix** in your document, and **upload any source code** as additional files in your submission.

The **next two** questions refer to the following optimization problem. Suppose that you are given the function

$$f(x) = x \sin(10\pi x) + 1$$

and you are asked to:

$$\text{maximize } f(x) \quad \text{subject to } x \in [-0.5, 1]$$

Question 1 (20 points): Implement a simple genetic algorithm that encodes the decision variable x as a string of a sign variable and 4 base-10 digits (e.g., -0.1234 is (-1,1,2,3,4)). You are free to choose whatever GA parameters would like (e.g., crossover operator, mutation operator, population size, number of parents, crossover probability, mutation probability), but your algorithm must have **both** crossover and mutation.

- Describe your GA-implementation choices (5 points)
- Generate two plots:
 - Best, worst, and average fitness for each successive generation of the GA (5 points)
 - Best individual for each successive generation of the GA (5 points)
- Plot the function f and evaluate the quality of the solution found by the GA (5 points)

Question 2 (10 points): Instead of encoding the decision variable as a string of base-10 digits, use a genotype with a **single gene** representing the **continuous** value of the decision variable. Because there is only one gene, there is **no crossover**, and parents selected for crossover are simply copied without any recombination. In this case, the GA is effectively **only doing mutation**.

- Implement this mutation-only GA, and **compare the efficiency** (i.e., number of generations needed to find a solution) of your mutation-only GA to the base-10 discrete GA from Question 1. **For the same number of generations, is one algorithm favorable over the other for some reason? Justify your response.**

Question 3 (BONUS) (4 points): Assume that a base-10 genotype (x_1, x_2, x_3, x_4) represents the number $0.x_1x_2x_3x_4$. While analyzing successive population generations from the operation of a GA, you observe that once all members of the population are within 0.001 of each other, the recombination operator **very rarely** produces offspring that differ from the parent generation, and all new variation in the population is driven by mutation only.

- Explain why this occurs (2 points).
- Why might it be **useful** for the recombination operator to act this way? **HINT:** Think about about “global” and “local” search as well as the **exploration–exploitation tradeoff** when forming your answer. (2 points)

References

Omid Bozorg-Haddad, Mohammad Solgi, and Hugo A. Loáiciga. *Meta-Heuristic and Evolutionary Algorithms for Engineering Optimization*. John Wiley & Sons, 2017. ISBN 978-1119386995.

Kevin M. Passino. *Biomimicry for Optimization, Control, and Automation*. Springer-Verlag, London, 2005. doi:[10.1007/b138169](https://doi.org/10.1007/b138169).