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# **Incorporating transposons into Genetic Algorithms**

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# Genetic Algorithms

Genetic Algorithms (GAs) are optimization techniques inspired by **natural evolution**, introduced in the 1960s by John Holland.

- Utilize processes like **selection**, **mutation**, and **crossover** to evolve solutions
- Effective in exploring large solution spaces
- **Premature convergence** is a significant challenge, leading to loss of diversity



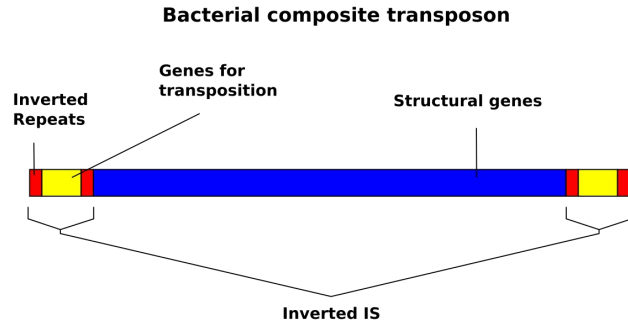
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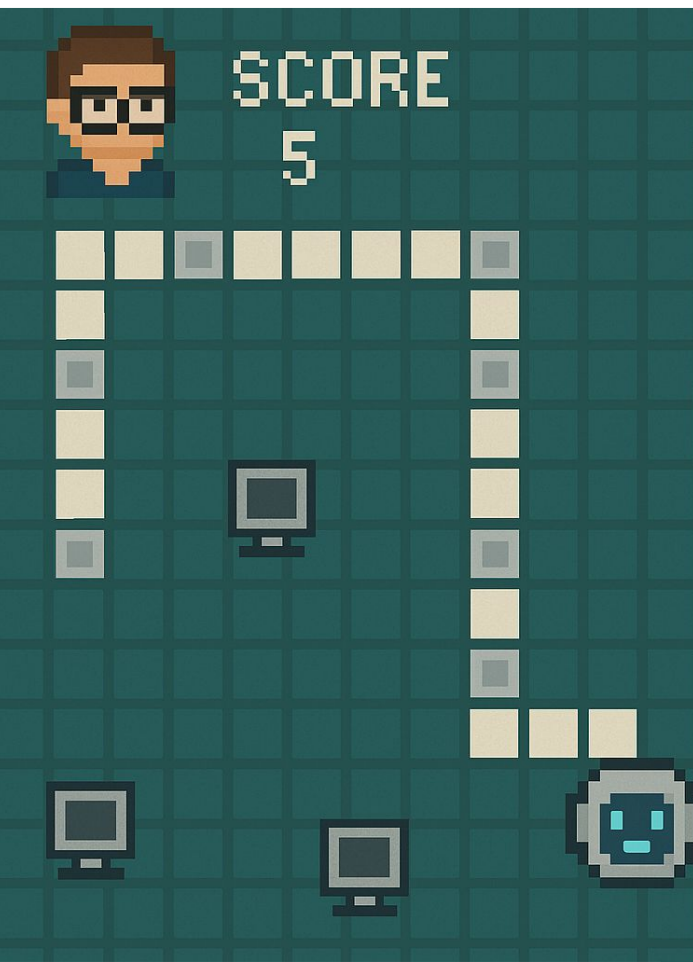


# The Role of Transposons

Transposons are **mobile genetic elements** that can enhance genetic diversity within GAs.

- Help maintain genetic variability to prevent **premature convergence**
- Introduce new genetic combinations, promoting exploration of solution spaces
- Two types: **DNA transposons** (direct movement) and **retrotransposons** (move via RNA)

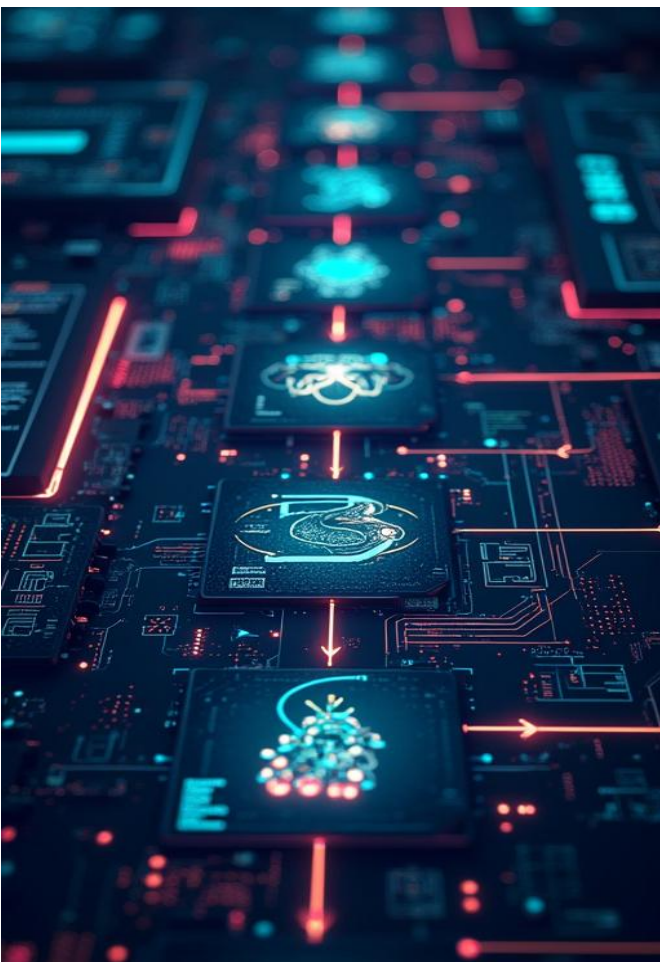




## The Snake Game Problem

The Snake Game serves as a testbed for evaluating GAs with transposons.

- A snake in a 2D board that aims at eating as many apples as possible while avoiding the walls and its own body, since it grows with every apple eaten.
- In each cycle, the snake can either turn left, keep its current direction, or turn right, relative to its head's current orientation.
- This problem is simple enough to focus on transposons, but still complex enough so as to being able to observe a decent diversity of behaviors and hence to compare the different algorithm approaches.



## Research Objectives

This study aims at integrating **transposons** into **GAs** in order to enhance optimization performance.

- Main goal: Improve performance in **GAs** using transposons, and preventing premature convergence in optimization issues.
- Secondary objectives: Analyze effectiveness of **transposons** and their **probability settings**



## ✓ THESIS

> \_\_pycache\_\_

> Documentation

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🔗 SnakeDNA.py

🔗 SnakeGene.py

🔗 Transposon.py

> Results

> Scores

## ✓ Snake

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🔗 Snake.py

🔗 SnakeFarm.py

🔗 SnakeGame.py

> SnakeGenerations

## ✓ Utils

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🔗 Constants.py

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# System Architecture Overview

The system consists of several classes designed to simulate genetic operations in the Snake Game.

- **SnakeGene Class:** Defines an individual gene
- **Transposon Class:** Manages transposon mechanisms and (transposon) mutation probability
- **SnakeChromosome Class:** Wraps a list of SnakeGenes and a list of Transposons inside a class, and enriches them with several important methods.
- **Snake Class:** Represents the agent's behavior, with a body and a brain
- **SnakeFarm Class:** Class in charge of executing the Genetic algorithm.
- **SnakeGame Class:** Environment for agent interactions



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# Evaluating Transposons in GAs

Three configurations were tested to evaluate the performance of **transposon-enhanced GAs**.

- **Traditional GA:** Baseline with no transposons
- **GA with 20% Transposon Probability:** Lower integration rate
- **GA with 40% Transposon Probability:** Higher integration to assess impact on diversity

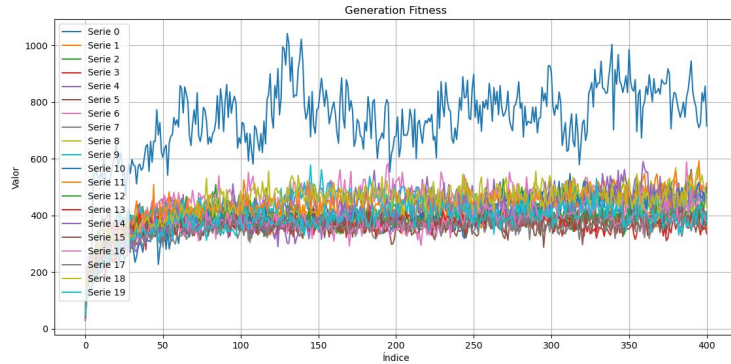


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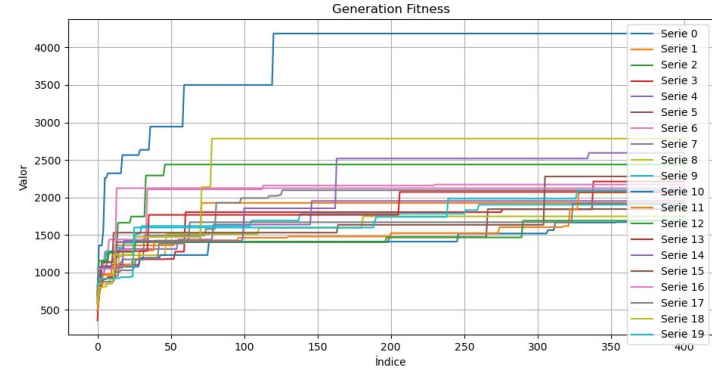


# Standard GAs



## Mean scores of populations

Mean scores of the populations in the different executions



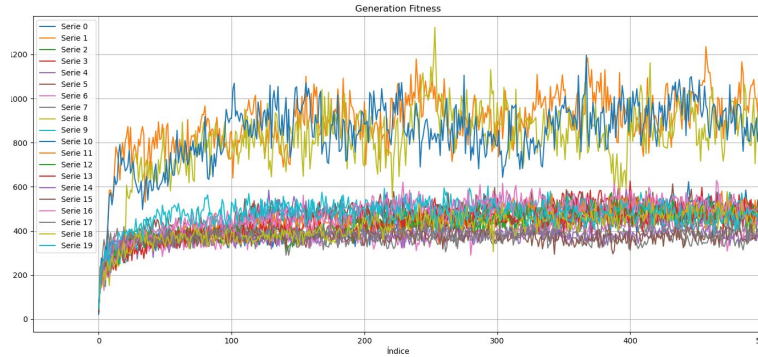
## Best scores

Evolution of best scores of the populations in the different executions



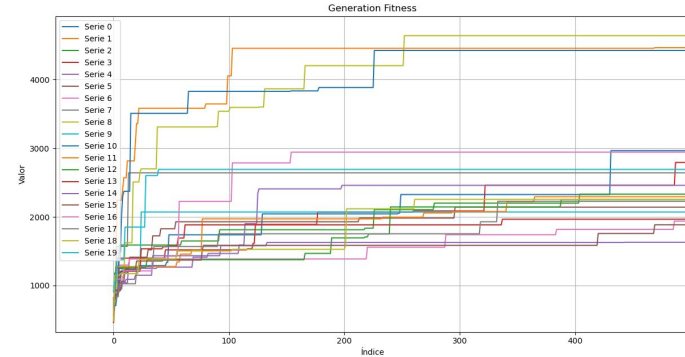
# GAs with transposons

(maximum probability of mutation: 20%)



## Mean scores of populations

Mean scores of the populations in the different executions

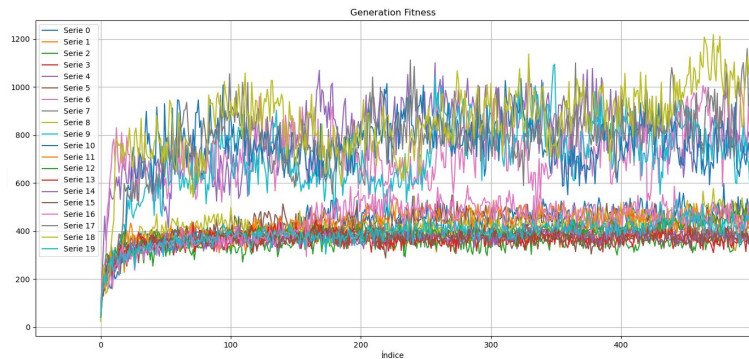


## Best scores

Evolution of best scores of the populations in the different executions

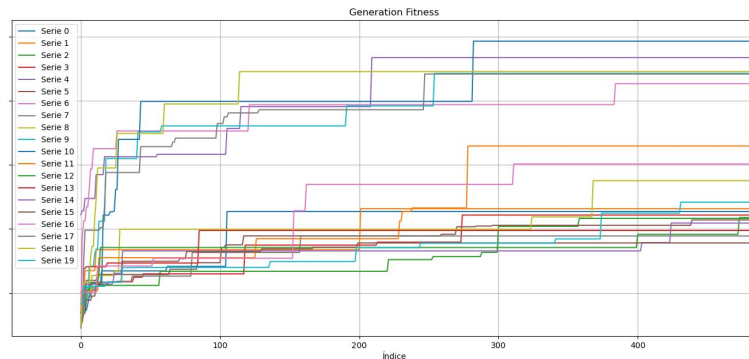
# GAs with transposons

(maximum probability of mutation: 40%)



## Mean scores of populations

Mean scores of the populations in the different executions



## Best scores

Evolution of best scores of the populations in the different executions



14



23



## Key Findings and Conclusions

The research indicates that incorporating **transposons** into **GAs** can significantly enhance performance.

- Transposons help maintain **diversity**
- Improved exploration of solution spaces leads to better **optimization outcomes**
- Demonstrated potential for transposons to enhance traditional GAs effectively



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# Future Work Directions

Future research will explore additional avenues to further validate findings.

- Implementing different features, such as using A\* distances instead of Manhattan, for example.
- Testing in more complex environments to assess **robustness**
- Investigating different types of **transposons** for varied impacts
- Expanding applications of **transposon-enhanced GAs** to other optimization problems



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**Thank you**

Code available at: <https://github.com/Somersault0023/TFM-UAH-Transposons/tree/main>