# **GAs To Solve The Travelling Salesman Problem**

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This report investigates the *Travelling Salesman Problem* through the use of a genetic algorithm (GA) employing the *inver-over* crossover operator, a combination of scramble and swap mutations, and a combination of  $\mu + \lambda$  and replacement survival selection operators.

## 1. PROBLEM SPECIFICATIONS

Wolfram MathWorld defines the travelling salesman problem as "a problem in graph theory requiring the most efficient (*i.e.*, least total distance) Hamiltonian cycle a salesman can take through each of n cities," for which "no general method of solution is known." [5] The website also remarks that the problem is assigned to the class of NP-hard (non-deterministic polynomial time) problems.

Given three datasets, for territories Western Sahara, Uruguay, and Canada of size 29, 734, and 4, 663 cities, respectively, the objective has been to design a genetic algorithm from scratch in the Python programming language to solve the TSP for each locale using some advanced technique(s).

The first advanced technique employed in the algorithm is the *inver-over* hybrid crossover/mutation operator.

## 2. THE INVER-OVER OPERATOR

The inver-over operator can be regarded as either a crossover or mutation operator because it takes information from other individuals in the GA's mating pool, yet bases a single offspring off of a single primary parent, much the same way a mutation operator is unary, performing mutation on a single individual [4]. Throughout this report, the inver-over operator

will be referenced as a crossover operator as this was the inver-over algorithm's place in the GA developed.

#### A. Usefulness

As *inver-over* embodies aspects of both crossover and mutation operators, the operator is designed to provide middle-ground to algorithms relying primarily on crossover for variation (as this is computationally expensive) and those relying on mutation for variety, since this is often ineffective in escaping local minima<sup>1</sup> [4].

## **B.** Representation

A set P of individuals of length n, each represented as a sequence

$$i_k = \langle C_{c_k(0)}, C_{c_k(1)}, C_{c_k(2)}, \dots, C_{c_k(n-1)} \rangle,$$

where  $c_k \colon \{0,1,\dots,n-1\} \to \{1,2,\dots,n\}$  is a bijection between indices of  $i_k$ ,  $0 \leqslant k < \operatorname{card}(P)$ , and city indices as provided in the data file, is used to denote the population. Let  $M \subsetneq P$  be the mating pool containing individuals  $i_{m(0)}, i_{m(1)}, \dots, i_{m(\operatorname{card}(M)-1)}$  with  $m \colon \{0,\dots,\operatorname{card}(M)-1\} \to \{0,\dots,\operatorname{card}(P)-1\}$  an injective function mapping individuals in the mating pool M to their equal 'self' in the population P.

Fitness of an individual  $i_k$ , denoted by fitness $(i_k)$  in the Algorithm 1, is then determined by the formula

<sup>&</sup>lt;sup>1</sup>Local minima occur when the natural selection in the GA narrows the gene pool towards a ostensibly optimal solution and eliminates individuals that otherwise would evolve to become the true optimal solution.

in Equation (1).

$$fitness(i_m) = \sum_{j=0}^{n-1} \left\| \overrightarrow{C_j} \overrightarrow{C_{j+1}} \right\|, \qquad (1)$$

where

$$\left\| \overrightarrow{C_j} \overrightarrow{C_{j+1}} \right\| = \sqrt{(x_{C_{j+1}} - x_{C_j})^2 + (y_{C_{j+1}} - y_{C_j})^2}$$

is the Eucliean distance between cities  $C_i$  and  $C_{i+1}^2$ .

## C. Algorithm

The algorithm (Algorithm 1) used mirrors that depicted in the article by Tao and Michalewicz [4].

## **Algorithm 1.** The inver-over operator

#### Require:

- $\triangleright M$  be the mating pool
- $\triangleright 0 \leqslant i < \operatorname{card}(M)$  be the index of the initial parent
- $\triangleright n$  be the number of cities in the tour

#### **Ensure:**

▶ New offspring individual created

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1: \mathbf{var}\ child := M[i]
                                             2: var unused := \{x \in \mathbb{Z} \mid 0 \le x < n-1\}
 3: var p := \frac{1}{2}
 4: \operatorname{var} c := \operatorname{rand}(unused)

    ▷ Select randomly

 5: unused := unused \setminus \{c\}
 6: while card(unused) > 0 do
 7:
         if \operatorname{rand}\{x \in \mathbb{R} \mid 0 \leqslant x < 1\} < p then
             c' := rand(unused)
 8:
 9:
         else
10:
             newPar := rand(M)
             newParC := where(newPar = child[c])
11:
                                         \triangleright Index j in newPar
             c' := newParentC + 1
12:
         unused := unused \setminus \{c'\}
13:
         if child[c \pm 1] = child[c'] then
14:
             break from the while loop
15:
         child[c:c'] := child[c':c]
16:
                                                         ▷ Invert
17:
18: if fitness(child) \geqslant \text{fitness}(M[i]) then
         return child
19:
20: else
21:
         return M[i]
```

Here M[i] is the primary parent from which the child is based. One can observe that all changes are then made to this individual, like a mutation, yet they involve other individuals, denoted newPar in the mating pool.

We borrow the following example of a single iteration of the algorithm from Tao and Michalewicz.

- 1. Let  $child = \langle 2, 3, 9, 4, 1, 5, 8, 6, 7 \rangle$  and the current city index c is 1 so child[c] = 3.
- 2. (a) Suppose the random number generated by  $\operatorname{rand}\{x \in \mathbb{R} \mid 0 \leqslant x < 1\}$  does not exceed p. Another city index c' from the child is selected, say c' = 6 so  $\operatorname{child}[c'] = 8$ . The section of  $\operatorname{child}$  after indices c and c' is inverted, leaving  $\operatorname{child} = \langle 2, 3, 8, 5, 1, 4, 6, 7 \rangle$ .
  - (b) Otherwise, another individual is (randomly) selected from the mating pool to become the new parent. Let this be  $\langle 1,6,4,3,5,7,9,2,8 \rangle$ . Here the city next to child[c]=3 is city 5. So, the segment of child to invert is that which starts after city 3 and terminates after city 5. This leaves  $child=\langle 2,3,5,1,4,9,6,7 \rangle$ .

As in Tao and Michalewicz, we remark that in either case the resulting string is intermediate in the sense that the above inversion operator is applied several times before an offspring is evaluated. After a number of iterations, suppose  $child = \langle 9, 3, 6, 8, 5, 1, 4, 2, 7 \rangle$  and c = 2 so child[c] = 6.

- 3. (a) If  $\operatorname{rand}\{x \in \mathbb{R} \mid 0 \leqslant x < 1\}$  is greater than p, the city following  $\operatorname{child}[c] = 6$  is selected from a randomly chosen individual in the mating pool M. Let this city be city 8. As 8 follows 6 in  $\operatorname{child}$ , the algorithm terminates.
  - (b) Otherwise, a randomly selected city is selected. This may also be 8, in which case the algorithm also terminates. If it is not 8, the algorithm continues as described in (2).

<sup>&</sup>lt;sup>2</sup> Take indices j modulo n so when  $j=n-1, (n-1)+1\equiv 0\pmod n$  and we compute the distance from the last city back to the first.

## 3. OPTIMIZATION

## A. The NumPy Module

Before any benchmarking was done, an effort was made to use the NumPy [2] module's interfaces as much as possible over raw Python data types. For example, when possible numpy.ndarray was chosen over Python's list data structure. The SciPy³ documentation states:

... the fact that [Python lists] can contain objects of differing types mean that Python must store type information for every element, and must execute type dispatching code when operating on each element. This also means that very few list operations can be carried out by efficient C loops—each iteration would require type checks and other Python API bookkeeping. [1]

As the problem being solved requires extensive accessing and mutating of arrays containing a single data type (*i.e.*, real numbers), NumPy provided a very easily-implemented speed boost.

## **B.** Cheaply Computing Distance

Suppose that the distance between cities  $C_j$  and  $C_{j+1}$  is not less than 1 and the same applies for cities  $C_k$  and  $C_{k+1}$ . That is,

$$1 \leqslant \left\| \overrightarrow{C_j C_{j+1}} \right\|$$

$$= \sqrt{(x_{C_{j+1}} - x_{C_j})^2 + (y_{C_{j+1}} - y_{C_j})^2}$$

and

$$1 \leqslant \left\| \overrightarrow{C_k C_{k+1}} \right\|$$

$$= \sqrt{(x_{C_{k+1}} - x_{C_k})^2 + (y_{C_{k+1}} - y_{C_k})^2}$$

Then  $\left\|\overrightarrow{C_j}\overrightarrow{C_{j+1}}\right\| < \left\|\overrightarrow{C_k}\overrightarrow{C_{k+1}}\right\|$  if, and only if,  $\left\|\overrightarrow{C_j}\overrightarrow{C_{j+1}}\right\|^2 < \left\|\overrightarrow{C_j}\overrightarrow{C_{j+1}}\right\|^2$ . Hence, by adding an assert statement to our code to ensure that

$$(x_{C_{i+1}} - x_{C_i})^2 + (y_{C_{i+1}} - y_{C_i})^2 \geqslant 1$$

we can omit the expensive square root operation  $\frac{1}{2}(n^2+n)$  times and still maintain accurate distance comparisons and fitness scores.

After the final iteration of the GA, the true distance is calculated using Equation (1).

## C. Maximizing Hardware Utilization

At the time of writing, nearly all modern computers utilize multiple processor cores (or in some cases multiple processors, each with multiple cores) to accomplish tasks<sup>4</sup>. Natively, the Python language uses only *one* core, leaving the rest underutilized.

Immediately, multi-threaded processing using Python's multiprocessing dummy module was investigated to take full advantage of the computer's hardware. Initially, this resulted in *worse* performance. Unfortunately, as QuantStart explains,

...the Python interpreter is not thread safe. This means that there is a globally enforced lock when trying to safely access Python objects from within threads. At any one time only a single thread can acquire a lock for a Python object or C API. The interpreter will reacquire this lock for every 100 bytecodes of Python instructions and around (potentially) blocking I/O operations. Because of this lock CPU-bound code will see no gain in performance when using the Threading library...[3]

- D. Precomputing All Distances
- 4. OUR ALGORITHM
- 5. RESULTS
- 6. CONCLUSION

### REFERENCES

- [1] Frequently Asked Questions. URL: https://www.scipy.org/scipylib/faq.html#what-advantages-do-numpy-arrays-offer-over-nested-python-lists.
- [2] Travis E. Oliphant. Guide to NumPy. Continuum Press, 2015.
- [3] Parallelising Python with Threading and Multiprocessing.

  URL: https://www.quantstart.com/articles/Parallelising-Python-with-Threading-and-Multiprocessing.
- [4] G. Tao and Z. Michalewicz. "Inver-over operator for the TSP". In: Lecture Notes in Computer Science (including subseries Lecture Notes in Artificial Intelligence and Lecture Notes in Bioinformatics) 1498 (1998), pp. 803–812. ISSN: 03029743.
- [5] Eric W. Weisstein. *Traveling Salesman Problem*. URL: http://mathworld.wolfram.com/TravelingSalesmanProblem.html.

<sup>&</sup>lt;sup>3</sup>NumPy is part of the SciPy Stack.

<sup>&</sup>lt;sup>4</sup>Specifications of several of Memorial's LabNet computers used for computation are included in Appendix A.

## A. COMPUTER SPECIFICATIONS