

## Algorithm 1: Evolutionary Approach using Pure Mutation.

The algorithm was inspired by a paper called a new evolutionary approach to cutting stock problems with and without contiguity [1]. In this approach a population of organisms are created, each one represents a solution.

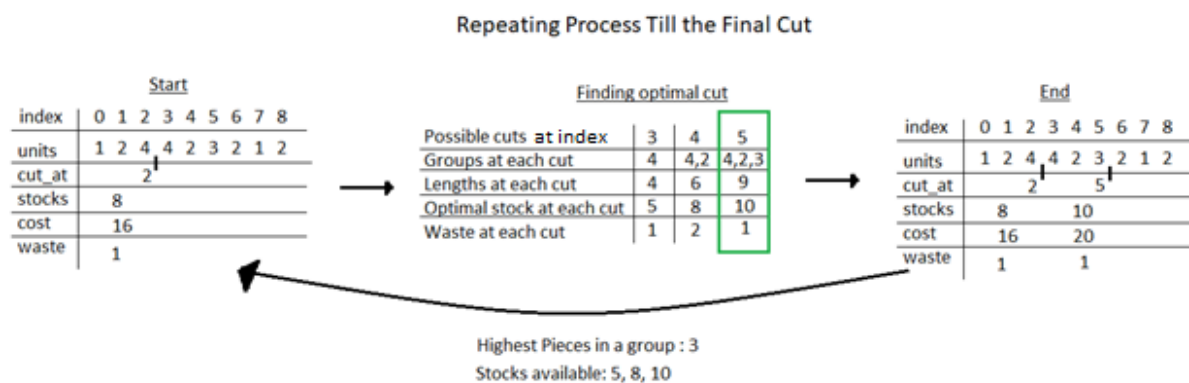
### **Chromosomes of an organism (excluding index and cost)**

In the organism, cost refers to the total cost of the solution, and another property is total waste which refers to the total waste of the solution.

index	0	1	2	3	4	5	6	7	8
units	1	2	4	4	2	3	2	1	2
cut_at			2			5			8
stocks		8			10			5	
cost		16			20			10	
waste		1			1			0	

### **Process of choosing cuts.**

The highest possible pieces in a group are calculated. Starting from the unit after the previously cut one, get the lengths of the combination of the next k units, with k increasing by 1 until the maximum possible pieces in a group have been reached or the group length is larger than the largest stock. Find the optimal stock that goes with each group length and retrieve the waste, the cut with the lowest waste and most units in it is chosen. Due to all stock's length being proportional to its cost, being bias to a certain stock was not needed, selecting the highest stock was used to shorten the number of cuts. This method is beneficial as the best possible cuts can be generated.



### **Evolutionary algorithm**

Each generation has all organisms clone themselves and then have those clones mutate twice. Having a clone mutate instead of the original organism allows the algorithm to allow the population to only improve or stabilise. For survivor selection I used as suggested by the paper, tournament selection. This allows the fittest members of the population to advance. Each organism had a chosen random number

of opponents to face off against, the fitness function which represents the waste of a solution was used to choose the winner. Half of the population with the most wins went on to the next generation.

**Three Point Swap (3PS) is used to mutate an organism.**

A cut represents a group of units that have been cut out of a stock. Mutating an organism involves picking a random unit (A) in a random cut. Two different cuts are selected and a random unit (B and C) for each cut is chosen, cuts with no waste will not be chosen. In the paper it suggests giving cuts with low waste a higher possibility of being chosen (Test 2). After experimenting I have found that giving cuts with high waste a higher probability generates better solutions (Test 1). Lastly, A 3PS is completed where A swaps with B, and B swaps with C.

index	0	1	2	3	4	5	6	7	8
units	1	2	4	4	2	3	2	1	2
cut_at			2			5			8
stocks	8			10			5		
cost	16			20			10		
waste	1			1			0		

$x_1$   $x_2$   $x_3$

$$P_t = x_1 + x_2 + x_3$$

$$P(x_i) = \frac{x_i}{P_t}$$

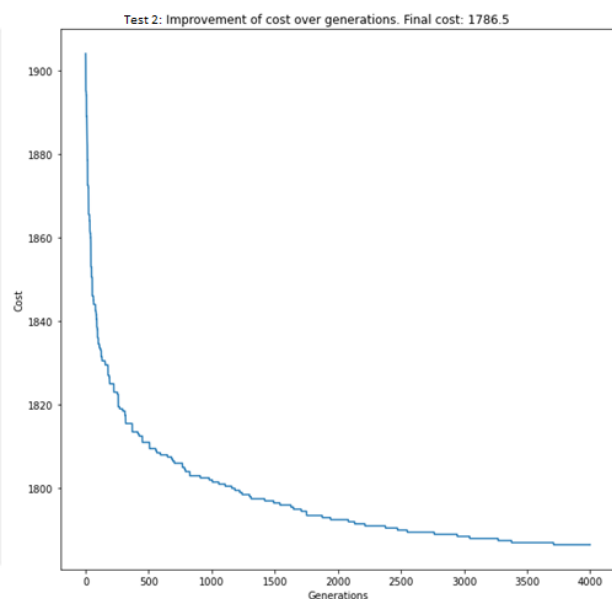
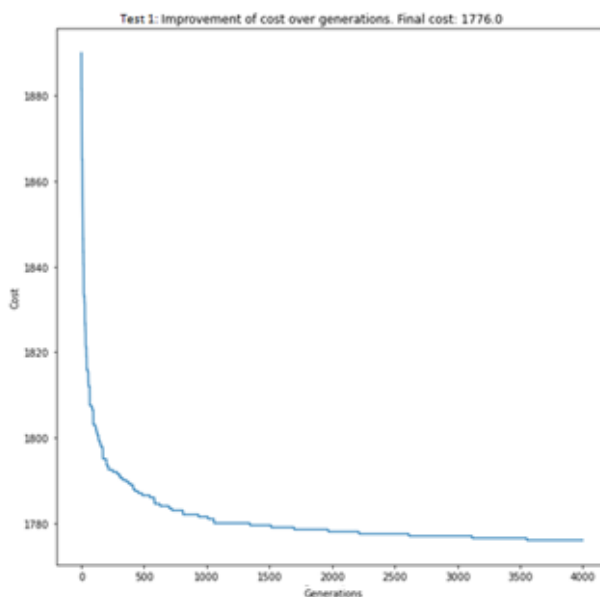
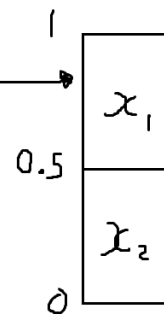
random number: 0.78

$$P_t = 2$$

$$P_{x_1} = 0.5$$

$$P_{x_2} = 0.5$$

$$P_{x_3} = 0$$



Algorithm 2: Evolutionary Approach using Mutation and Recombination.

This algorithm is like algorithm 1, but recombination is introduced which replaces cloning with parents.

### Evolutionary algorithm

A population is initiated and for each generation pairs of parents are selected to create 2 offspring till the population doubles. Proportional roulette wheel selection is used to select the parents. The probabilities of selecting a parent can be seen as spinning a roulette wheel with the size of the segment for each parent being proportional to its fitness. Waste is used as a fitness function over cost as the cost can never reach 0. Solutions which are closer to the optimal cost will be prioritised more as the effect of the inverse of waste is greater than the inverse of cost. As a parent can be picked multiple times, the better parents will be chosen more often than the poorer ones, thus fulfilling the requirements of survival of the fittest [2]. For survivor selection, the same tournament selection from solution 1 is used.

### Recombination

With recombination in this problem, it is not viable to take the first half of the units' chromosome from a parent and the send half of that chromosome from another as the offspring will have duplicated or lost units. When accounting for this, the recombination method I choose to create offspring is Order 1 Crossover. A section from the first parents' unit's chromosome is chosen and placed into the offspring in the same position. Next the remaining units are taken the second parent to fill in the missing places in the offspring. This is one of the fastest recombination operators which means a lot of generations can be processed in small amount of time [3].

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Parent 1: 8 4 7 3 6 2 5 1 9 0
Parent 2: 0 1 2 3 4 5 6 7 8 9
Child 1:  0 4 7 3 6 2 5 1 8 9
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[3]

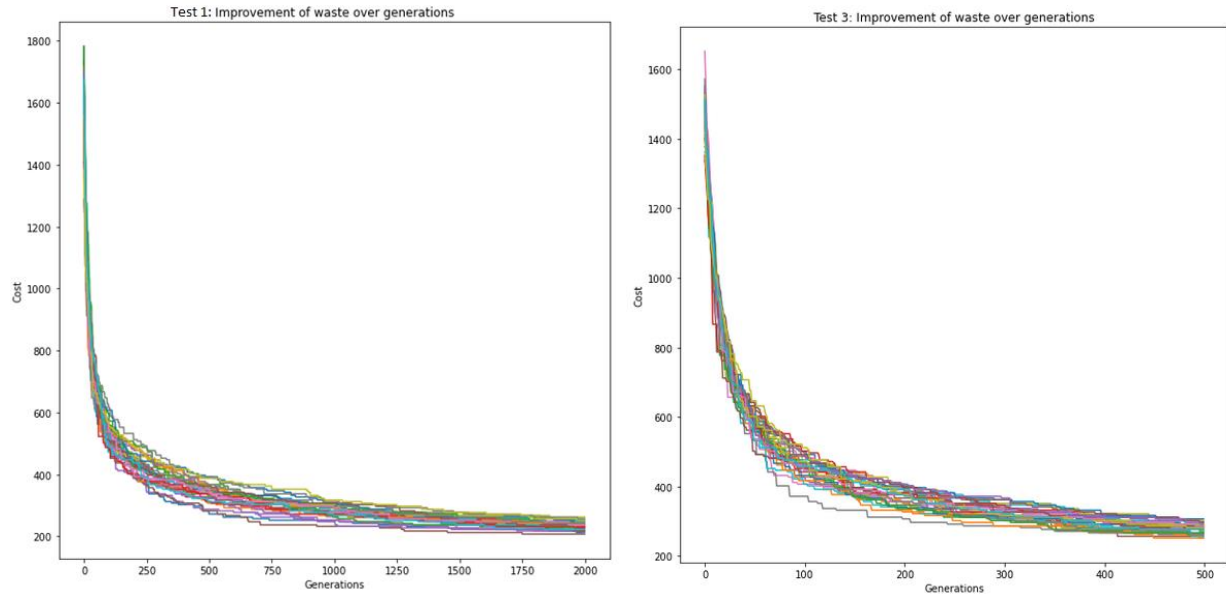
## Methodology

Both algorithm share 3 changeable parameters: generations, size, and opponents. I will be testing to see which combination of values for these parameters produce the best results. As the algorithms could run for days and get an almost optimal solution, a budget of 60 seconds has been set. As the sum of the length of all units is 17633, theoretically, the smallest waste can ever be is 2 (146 stocks of 120, and 1 stock of 115). This will be the target value; the goal would be to get as close of possible to it. Other changeable parameters would be the mutation chance in solution 2. As results may vary when an algorithm with set of parameters is run multiple times, I will be running each set 30 times to get a range of results.

### Algorithm 1

Test	Generations	Size	Opponents	Minimum waste	Mean waste	Maximum waste	Standard deviation	Average seconds to completion
1	2000	10	5	207	232.8	262	14.09	56.86
2	1000	20	10	217	243	267	12.94	55.17
3	500	40	20	252	277.7	307	14.82	58.46

When testing for the best parameters, I discovered that a trend where a small population with a larger number of generations produced the better result in under 60 seconds. The best results came from test 1 as it had the lowest minimum, mean, and maximum waste.

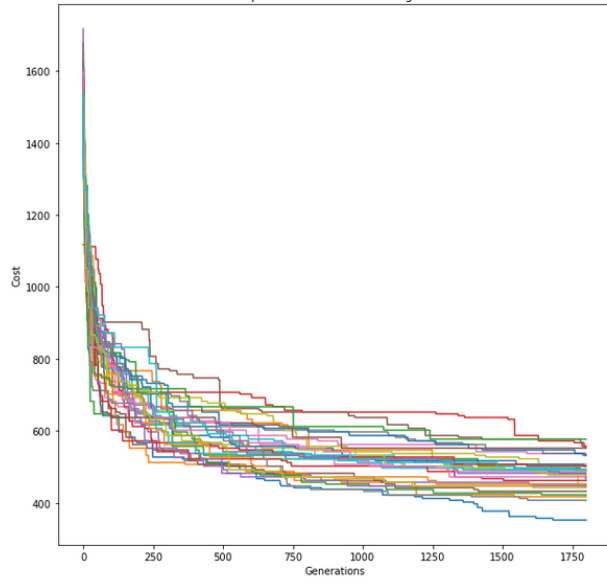


## Algorithm 2

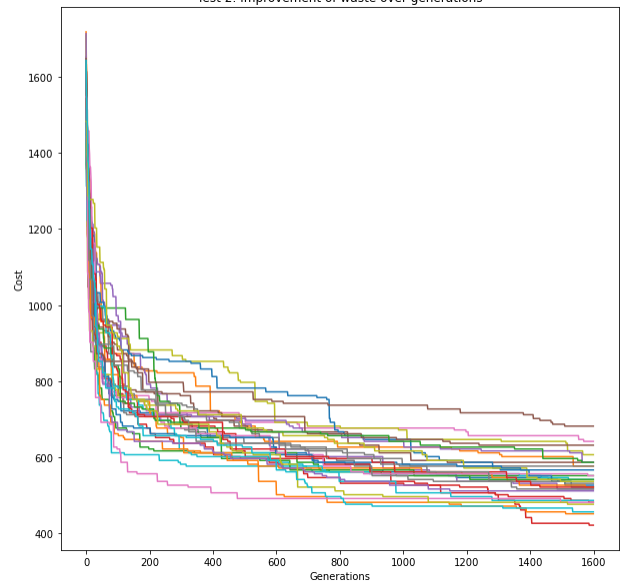
Test	Generations	Size	Opponents	Mutation chance	Minimum waste	Mean waste	Maximum waste	Standard deviation	Average seconds to completion
1	1800	10	5	0.3	352	481.33	577	49.83	61.36
2	1650	10	5	0.7	422	540	682	57.45	58.45
3	800	20	10	0.3	382	478.8	562	41.92	59.91
4	800	20	10	0.7	417	536.67	662	54.65	57.83
5	450	40	20	0.3	387	483.33	612	54.04	62.18
6	450	40	20	0.7	432	535.5	697	64.18	63.72

When testing for the best parameters, I discovered that a trend where a lower mutation chance produced a lower waste, this excludes test 2, 4, and 6 from being selected to represent the algorithm. Tests 1, 3, and 5 had very similar means but test 3 had a much lower standard deviation then the rest.

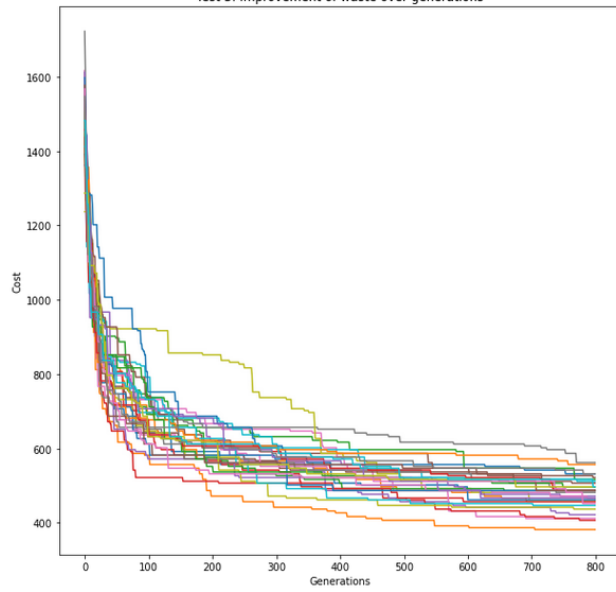
Test 1: Improvement of waste over generations



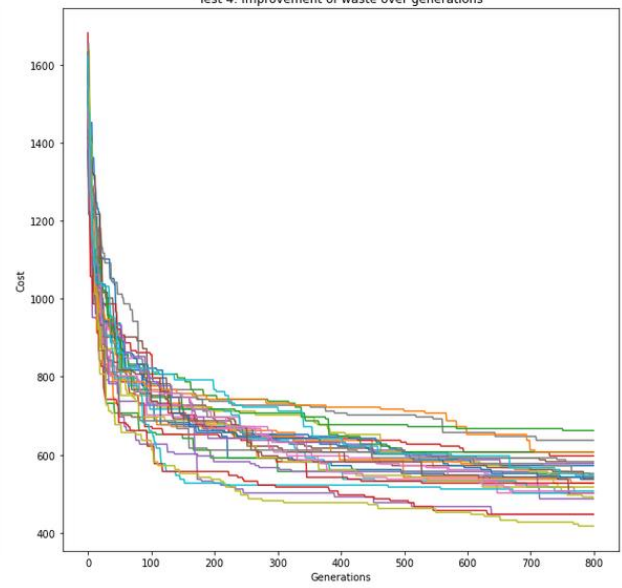
Test 2: Improvement of waste over generations

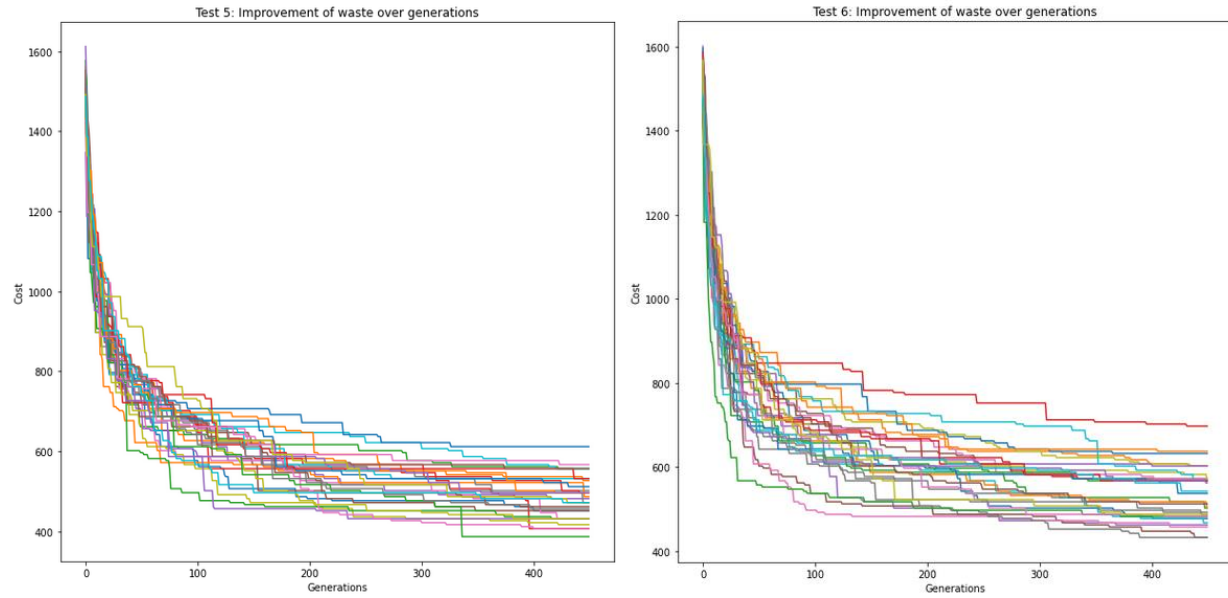


Test 3: Improvement of waste over generations



Test 4: Improvement of waste over generations





## Conclusion

Solution 1 is algorithm 1 with test 1 parameters, solution 2 is algorithm 2 with test 3 parameters, as previously stated, these produced the best results. The two parts of the results to consider are mean and standard deviation. Solution 1 produced the lowest minimum, mean, and maximum waste, the minimum waste for solution 2 did not even come close to the maximum waste of solution 1. A low standard deviation means a solution is very closely related to the average which makes it very reliable. Therefore, a person does not have to chance whether they get an extremely bad or good solution as all runs will be very similar. The standard deviation in solution 1 is about a third of the size of solution 2, which makes it more reliable. With comparison completed, as solution 1 outperformed solution 2 in both categories, it is the best solution.

## References

- [1] K. Liang, X. Yao, C. Newton and D. Hoffman, "A new evolutionary approach to cutting stock problems with and without contiguity", *Computers & Operations Research*, vol. 29, no. 12, pp. 1641-1659, 2002. Available: [https://www.cs.bham.ac.uk/~xin/papers/COR\\_LiangYaoNewtonHoffman.pdf](https://www.cs.bham.ac.uk/~xin/papers/COR_LiangYaoNewtonHoffman.pdf).
- [2] N. Razali and J. Geraghty, "Genetic Algorithm Performance with Different Selection Strategies in Solving TSP", *Proceedings of the World Congress on Engineering*, 2011.
- [3] "Order 1 Crossover Operator", *Rubicite*. [Online]. Available: <https://www.rubicite.com/Tutorials/GeneticAlgorithms/CrossoverOperators/Order1CrossoverOperator.aspx>. [Accessed: 23- Apr- 2021].