

# Solving Traveling Salesman Problem using Genetic Algorithms

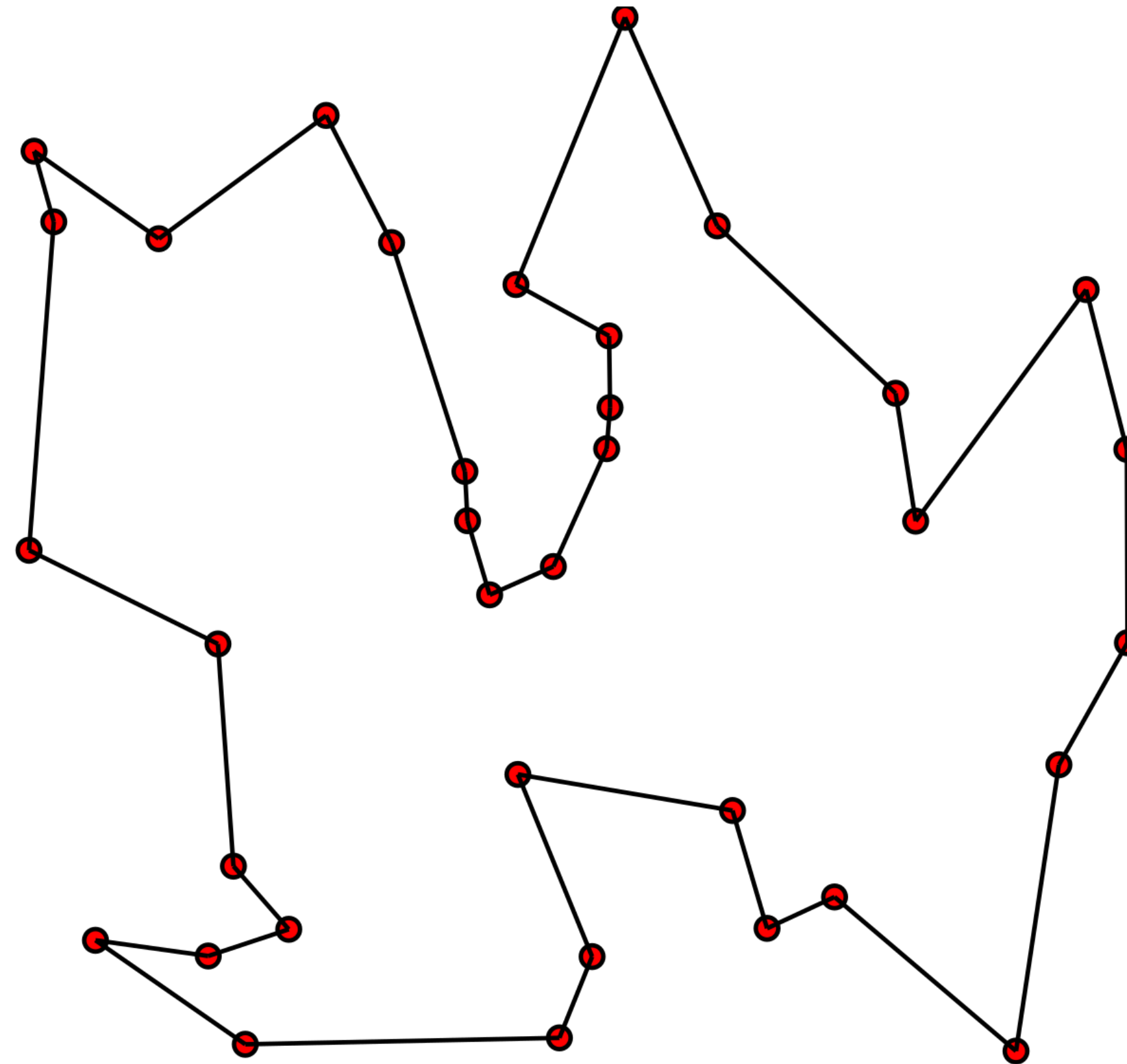
525.770: Intelligent Algorithms

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# Traveling Salesman Problem

- Find the shortest path such that all nodes are visited at least once and come back



# Genetic Algorithm For TSP

- Solving TSP is NP-Hard! No exact solutions in polynomial time
- Genetic Algorithm is a meta-heuristic algorithm
- Gives approximate solutions in reasonable amount of time
- Many different variants are possible (different cross-over, mutate strategies)
- This work compares some of these methods!

# Representation of Chromosome

- A binary chromosome representation is not useful for TSP
- We represent a chromosome as a simple path of nodes visited in order

1 2 3 5 6 4 7 9 8

# Mutation

- Mutation means small perturbations made to the chromosome
- Make mutation with a small probability ( $p_{\text{mutate}}$ )
- Two types considered:
  1. Swap mutation
  2. Insertion mutation

# Swap Mutation

- Two adjacent alleles are swapped.
- The position is randomly sampled

1 2 3 5 6 4 7 9 8

1 2 3 6 5 4 7 9 8

# Insertion Mutation

- alleles are swapped adjacent or not
- The positions are randomly sampled

1 2 3 5 6 4 7 9 8

1 2 3 9 6 4 7 5 8

# Cross Over Operations

- Single point cross over fails in TSP
- Ref: J.-Y. Potvin, The traveling salesman problem

tour (12564387)	:	<b>1</b>	<b>2</b>		<b>5</b>	<b>6</b>	<b>4</b>	<b>3</b>	<b>8</b>	<b>7</b>
tour (14236578)	:	1	4		2	3	6	5	7	8
offspring 1	:	<b>1</b>	<b>2</b>		2	3	6	5	7	8
offspring 2	:	1	4		<b>5</b>	<b>6</b>	<b>4</b>	<b>3</b>	<b>8</b>	<b>7</b>

Figure 5. Application of the one-point crossover on two parent tours.

- Note that the offsprings are not valid paths!



# Partially Mapped Cross-Over (PMX)

- This is a path preserving cross-over (maintains absolute positions of allele)

parent 1	:	<b>1</b>	<b>2</b>		<b>5</b>	<b>6</b>	<b>4</b>		<b>3</b>	<b>8</b>	<b>7</b>
parent 2	:	1	4		2	3	6		5	7	8
<hr/>											
offspring											
(step 1)	:	1	4		<b>5</b>	<b>6</b>	<b>4</b>		5	7	8
(step 2)	:	1	3		<b>5</b>	<b>6</b>	<b>4</b>		2	7	8

Figure 8. The partially-mapped crossover.



# Order Cross-Over (OX)

- This is an order preserving cross-over (maintains the relative order of allele)

parent 1	:	<b>1</b>	<b>2</b>		<b>5</b>	<b>6</b>	<b>4</b>		<b>3</b>	<b>8</b>	<b>7</b>
parent 2	:	1	4		2	3	6		5	7	8
<hr/>											
offspring											
(step 1)	:	—	—		<b>5</b>	<b>6</b>	<b>4</b>		—	—	—
(step 2)	:	2	3		<b>5</b>	<b>6</b>	<b>4</b>		7	8	1

Figure 11. The order crossover.

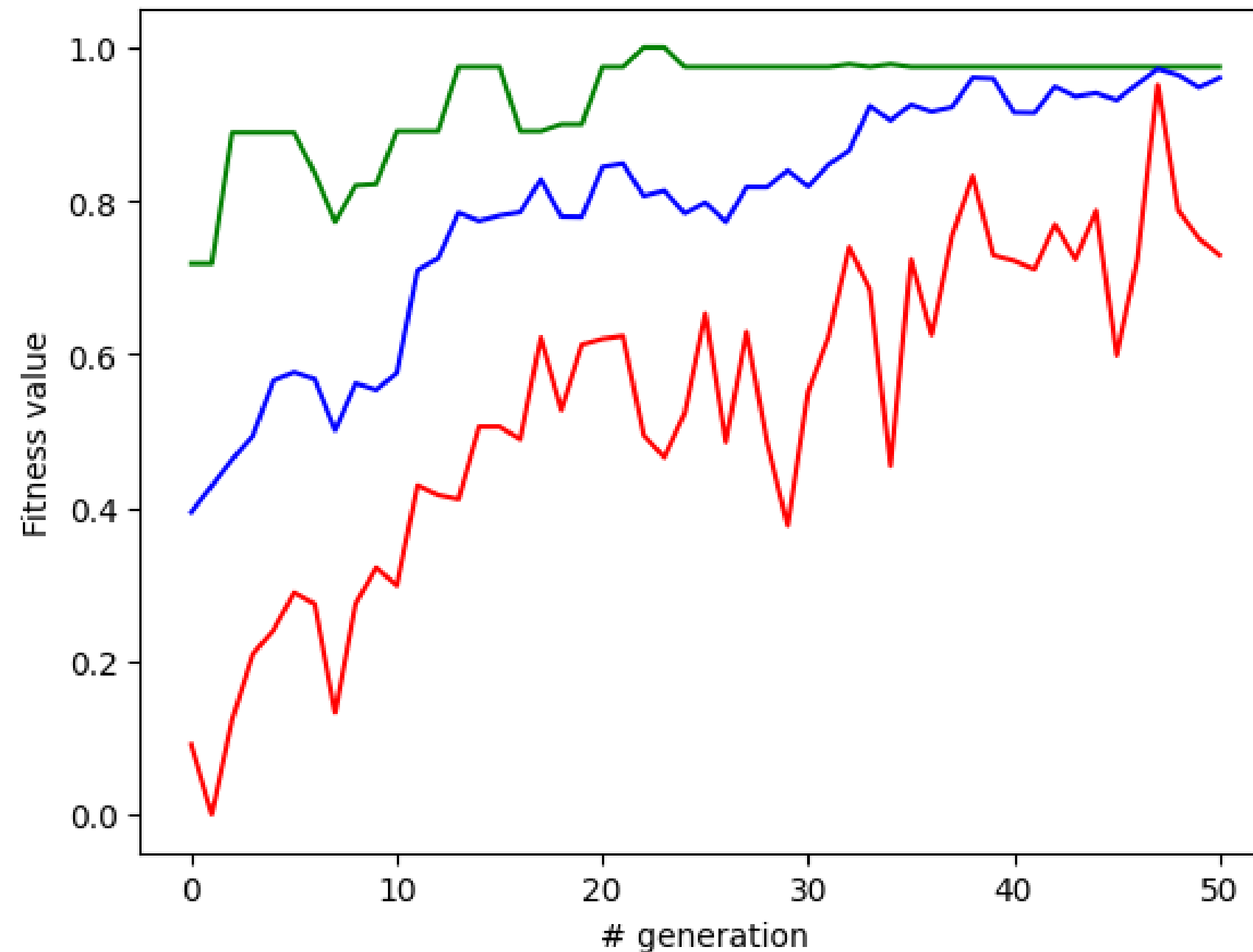


# Dataset

- The datasets are sampled Graphs
- Graph sizes: 10, 50, 100
- The edge weights are randomly sampled from 1 to 99

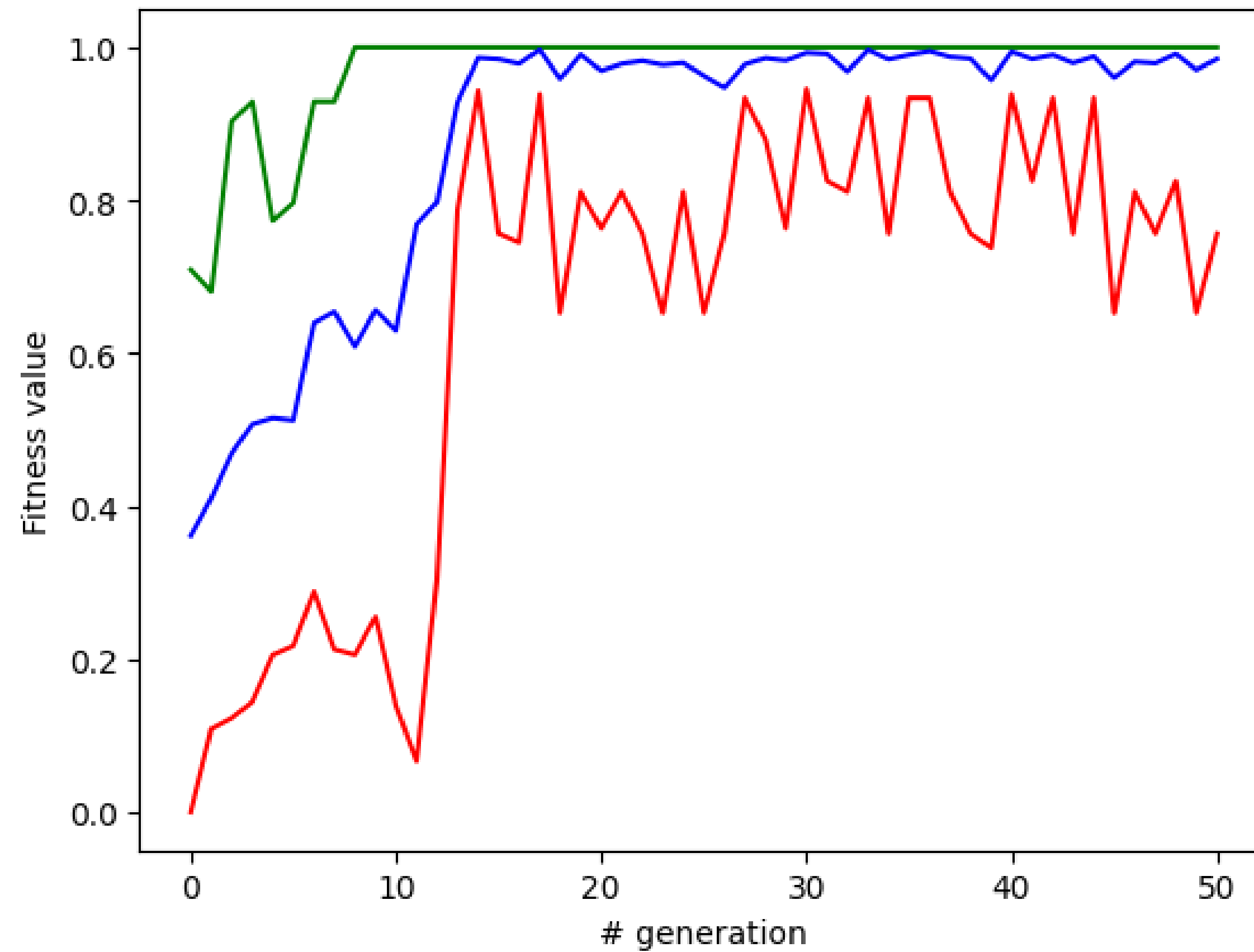
# Experiments

- # parameters: popsize = 20, pmutate = 0.1, pcross = 0.9, ngen = 50
- Mutation: swap, cross-over: Ox



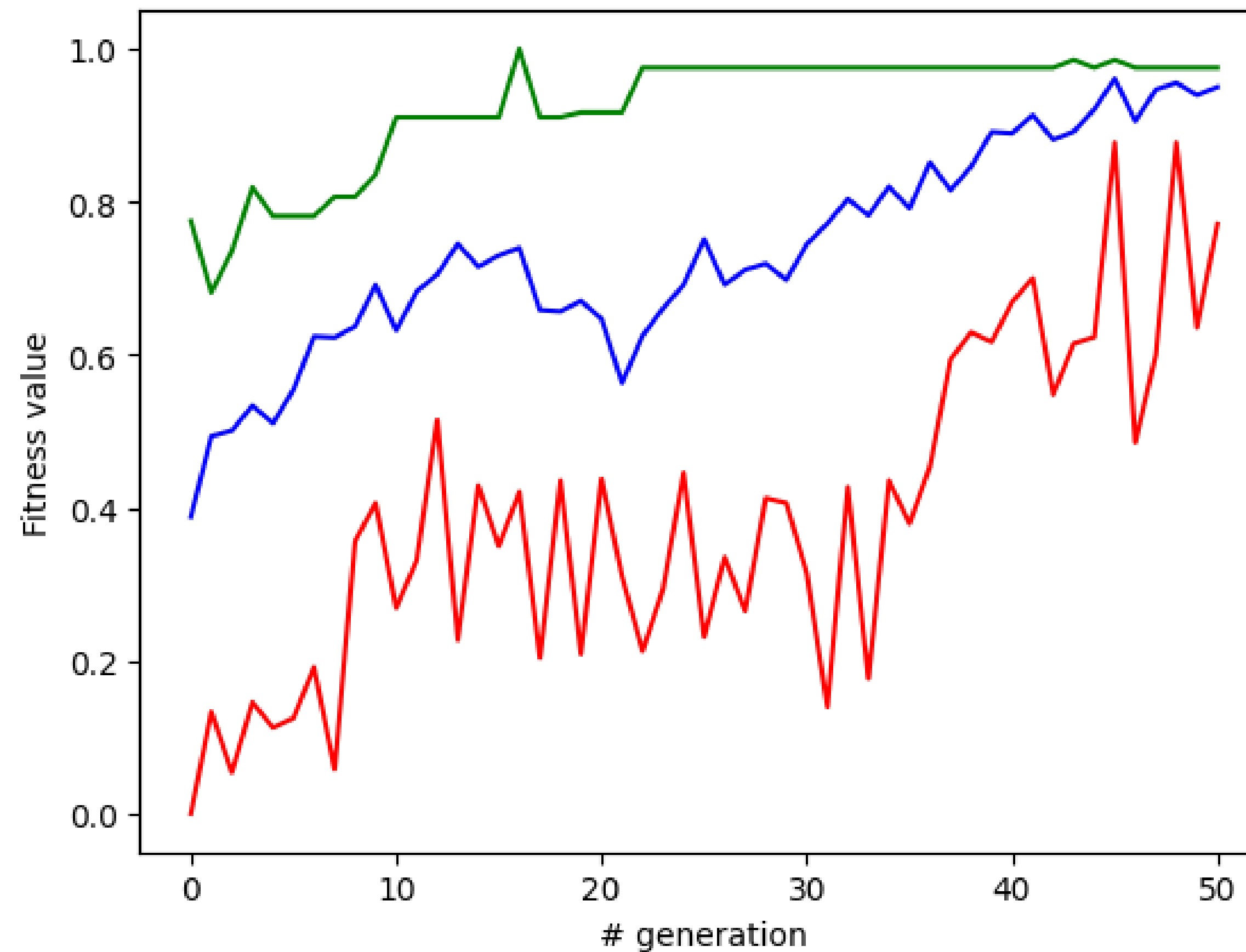
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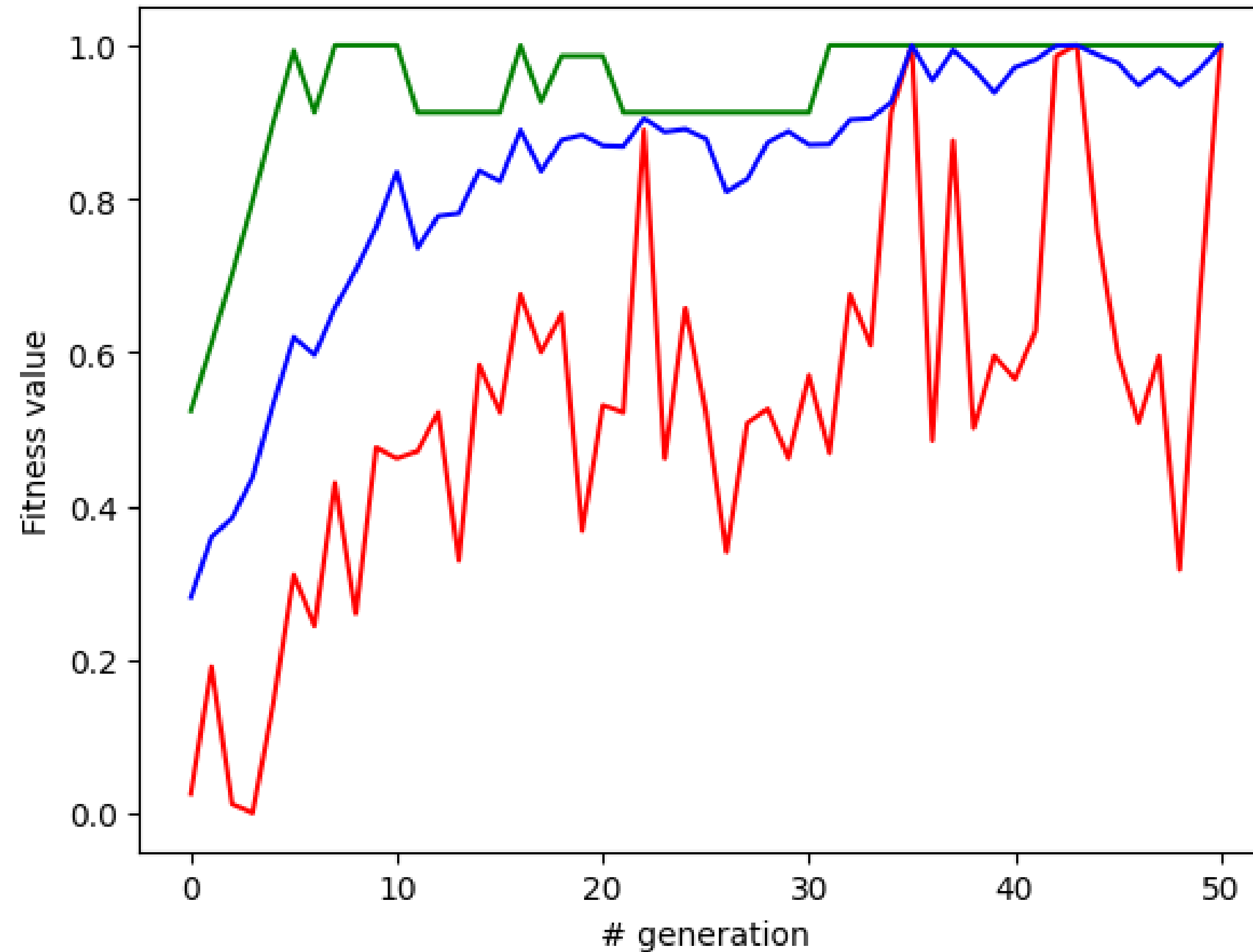
# Experiments

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# Experiments

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- Mutation: insertion, cross-over: pmx



# Comparison

- The minimum path length found by the different combinations of mutation & cross-over functions are

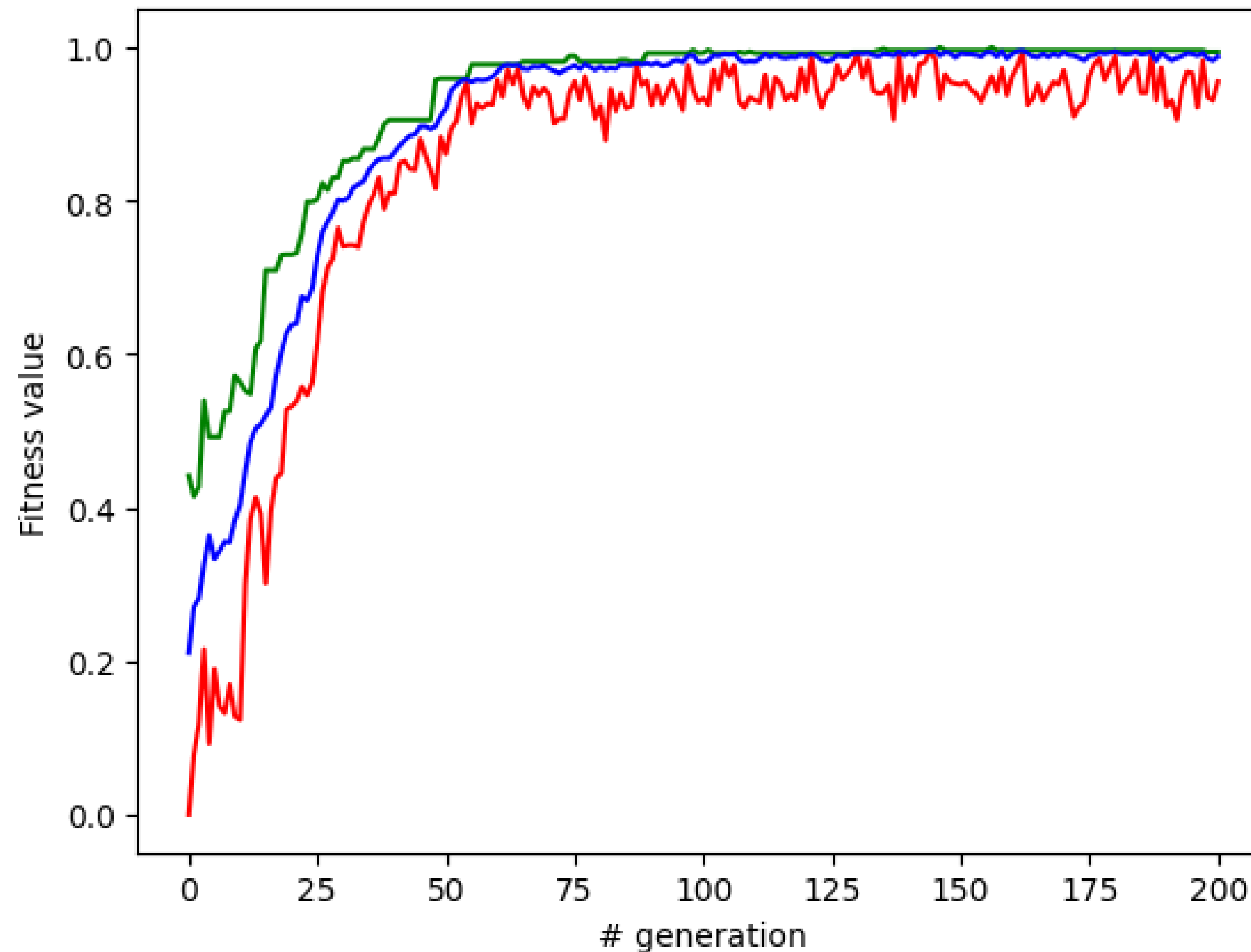
	Swap	Insertion
OX	259	261
PMX	291	261





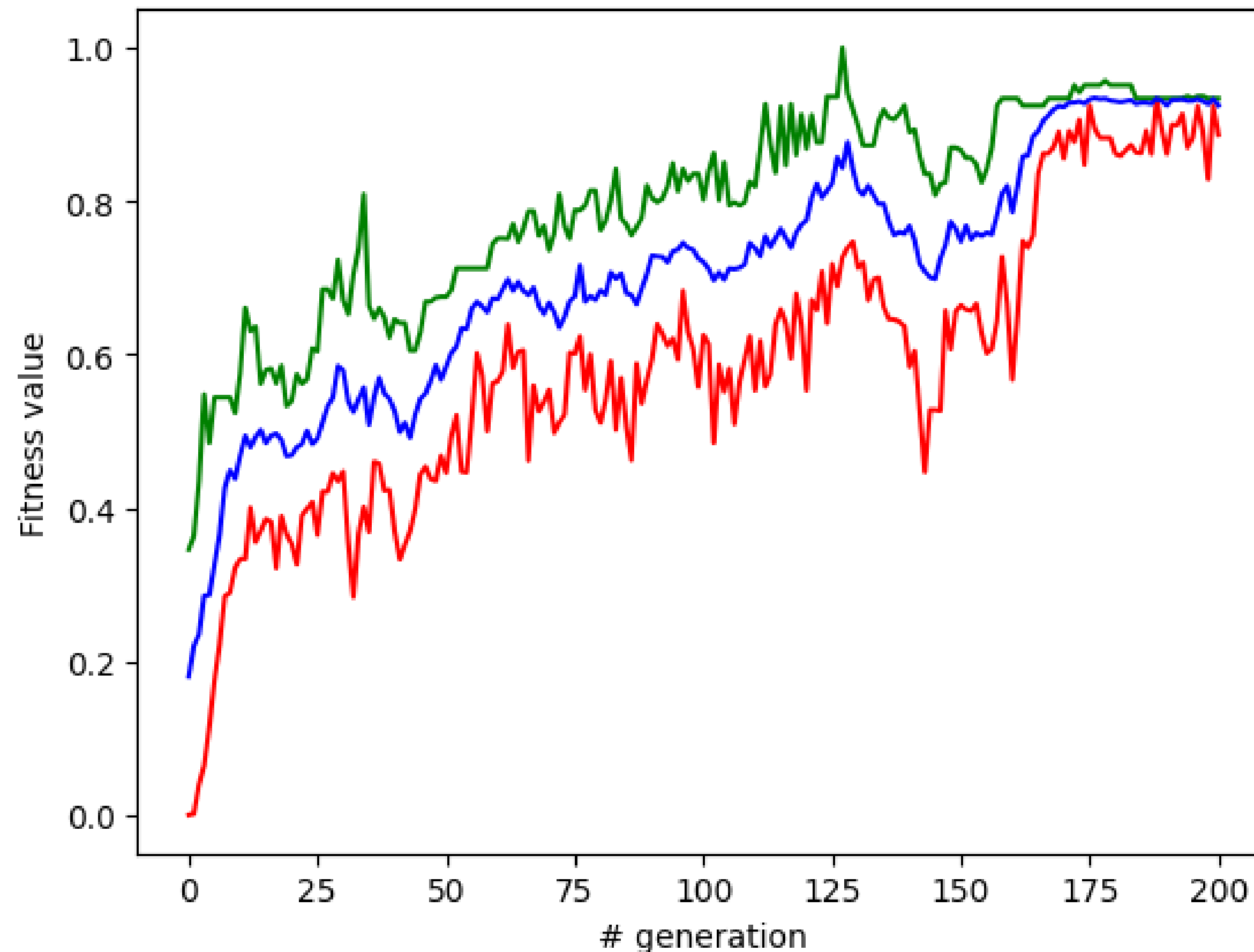
# Larger Graph (N = 50)

- # parameters: popsize = 20, pmutate = 0.1, pcross = 0.9, ngen = 200
- Mutation: swap, cross-over: pmx



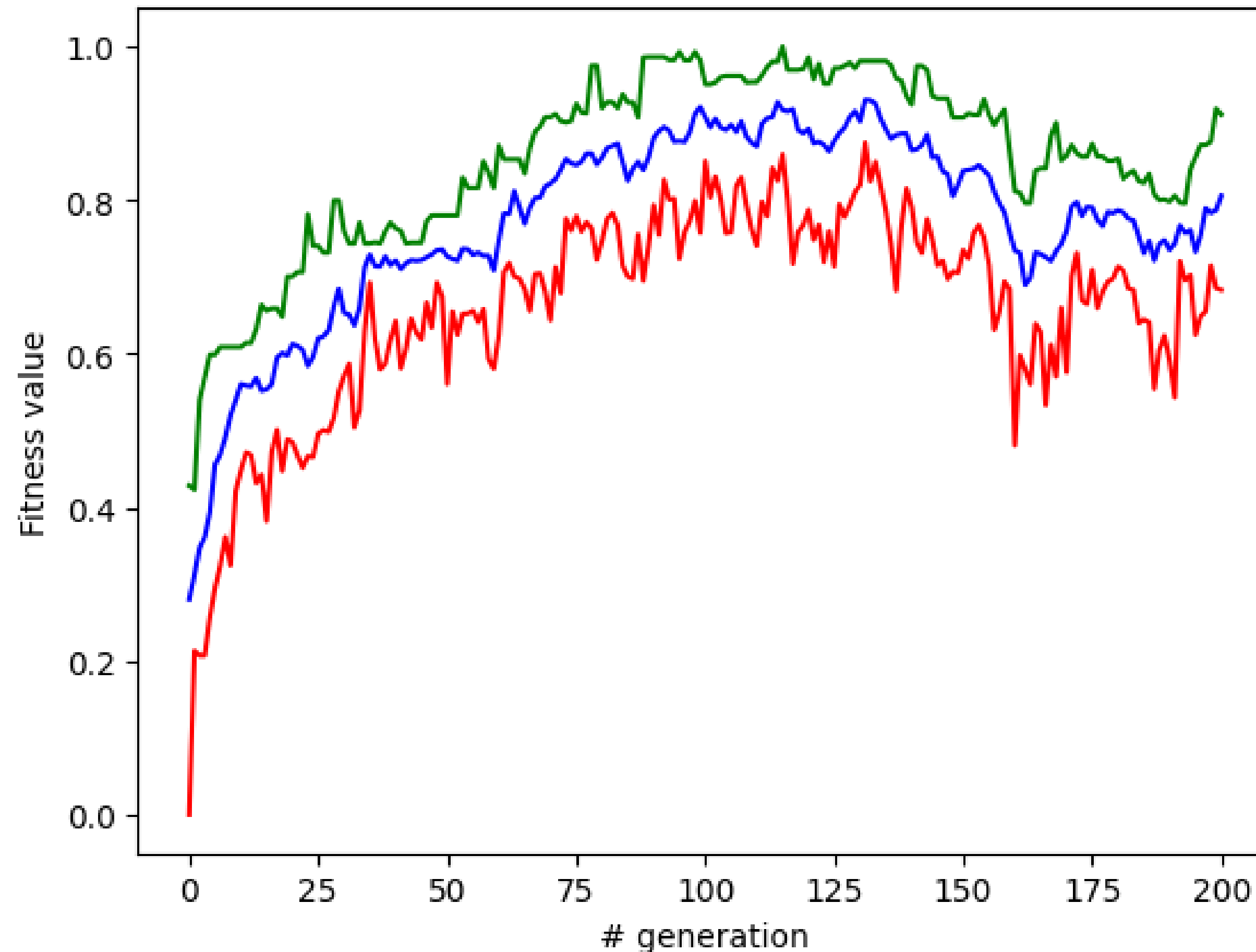
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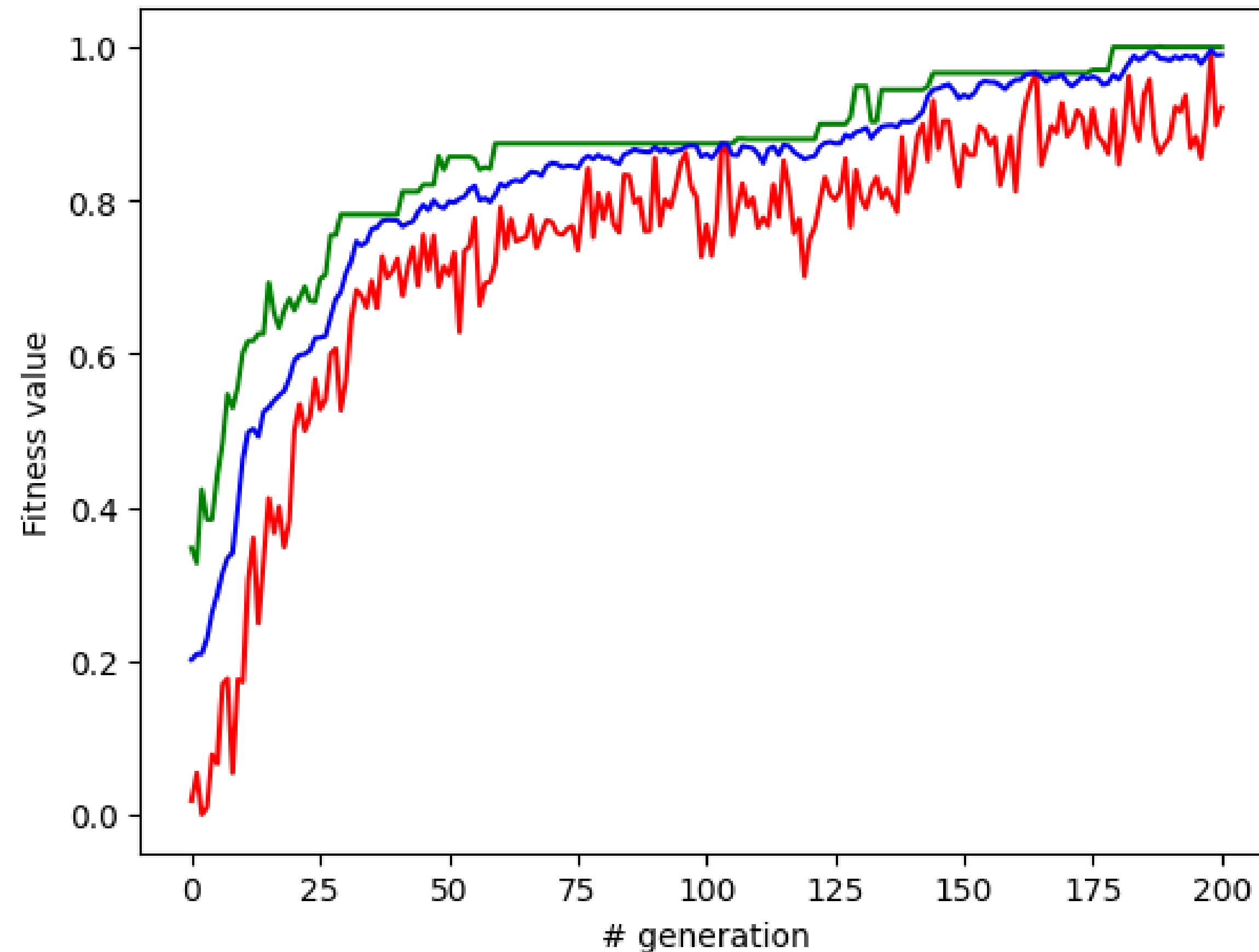
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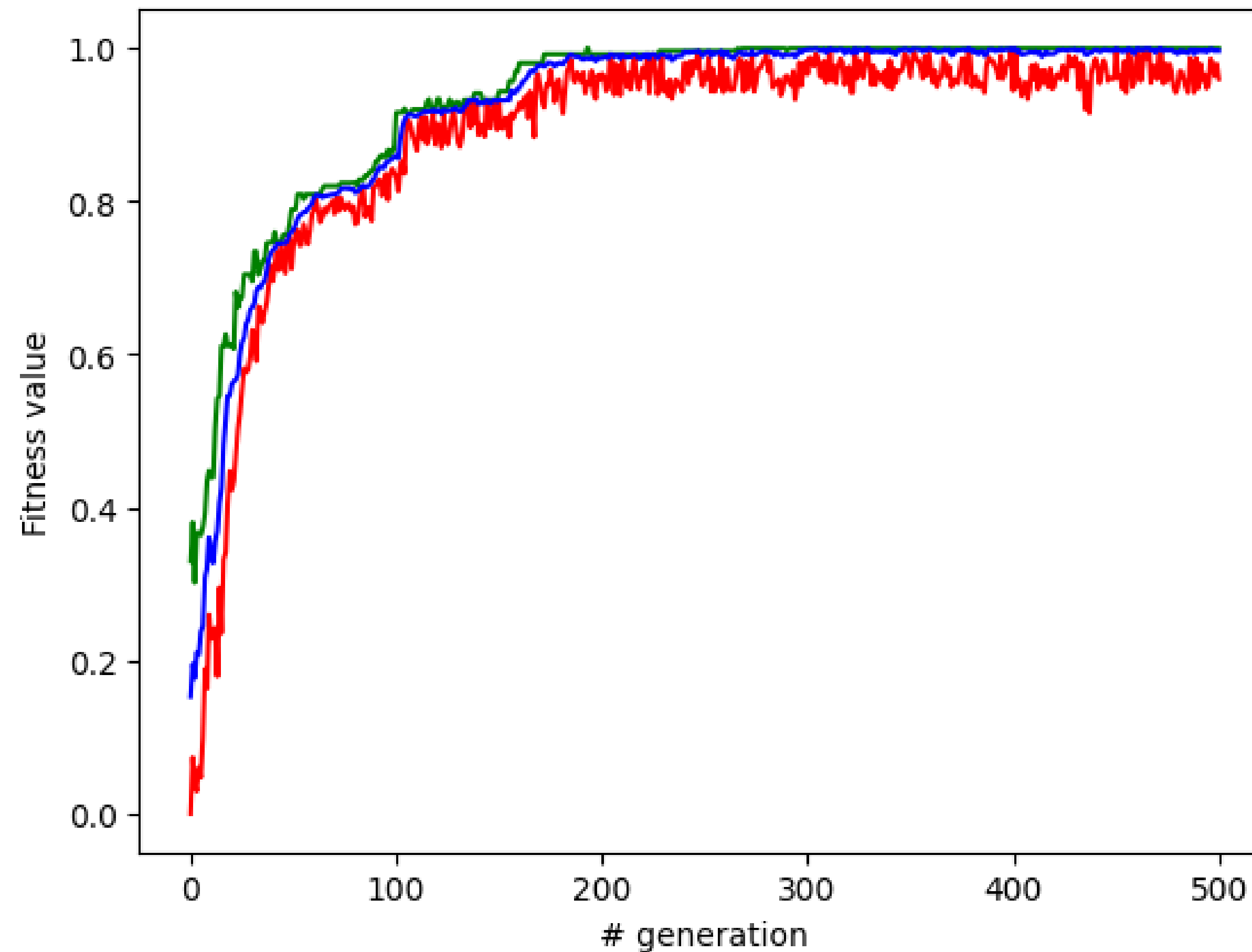
# Comparison

- The minimum path length found by the different combinations of mutation & cross-over functions are for  $N = 50$

	Swap	Insertion
OX	1218	1220
PMX	1028	1093

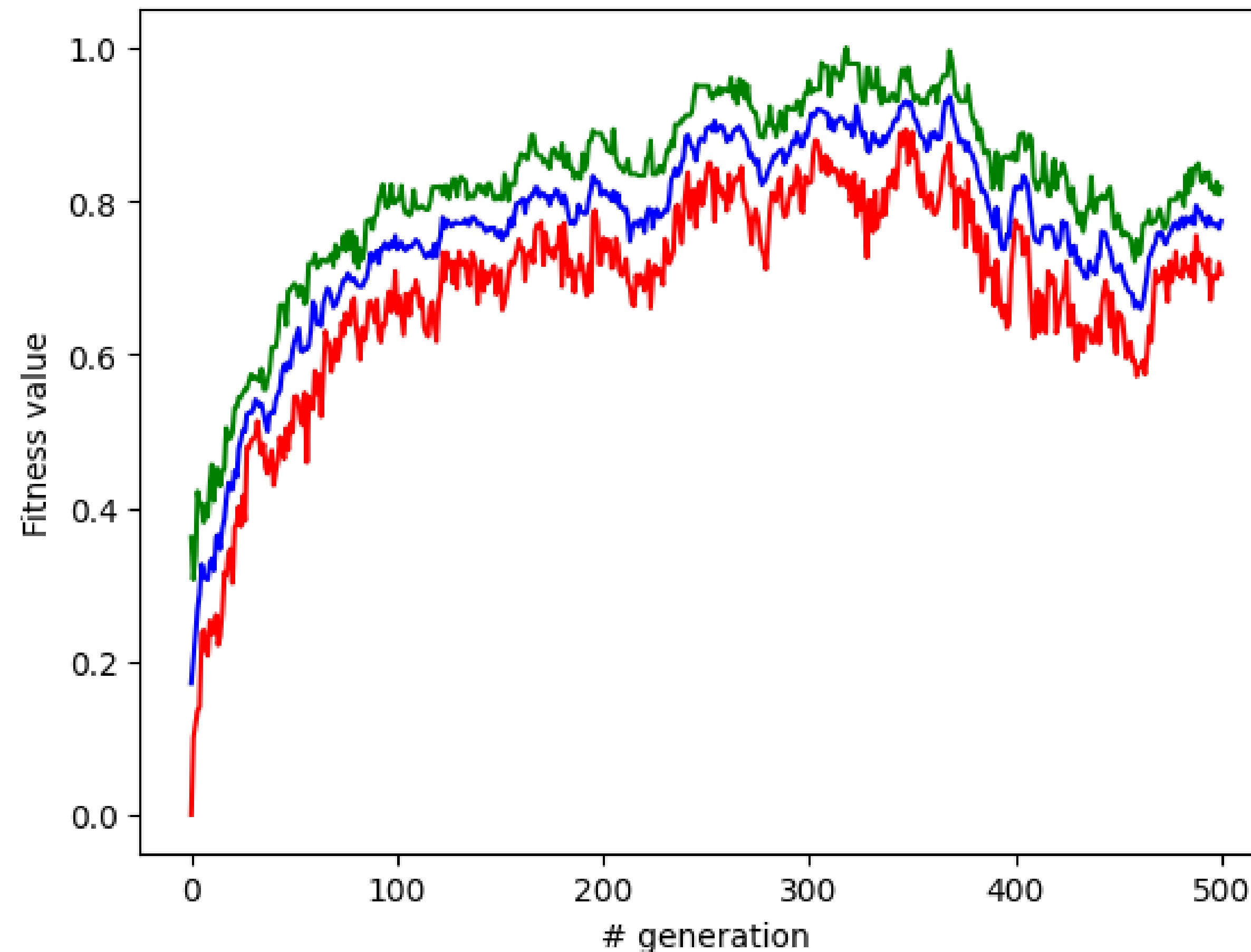
# Larger Graph (N = 100)

- # parameters: popsize = 20, pmutate = 0.1, pcross = 0.9, ngen = 500
- Mutation: swap, cross-over: pmx



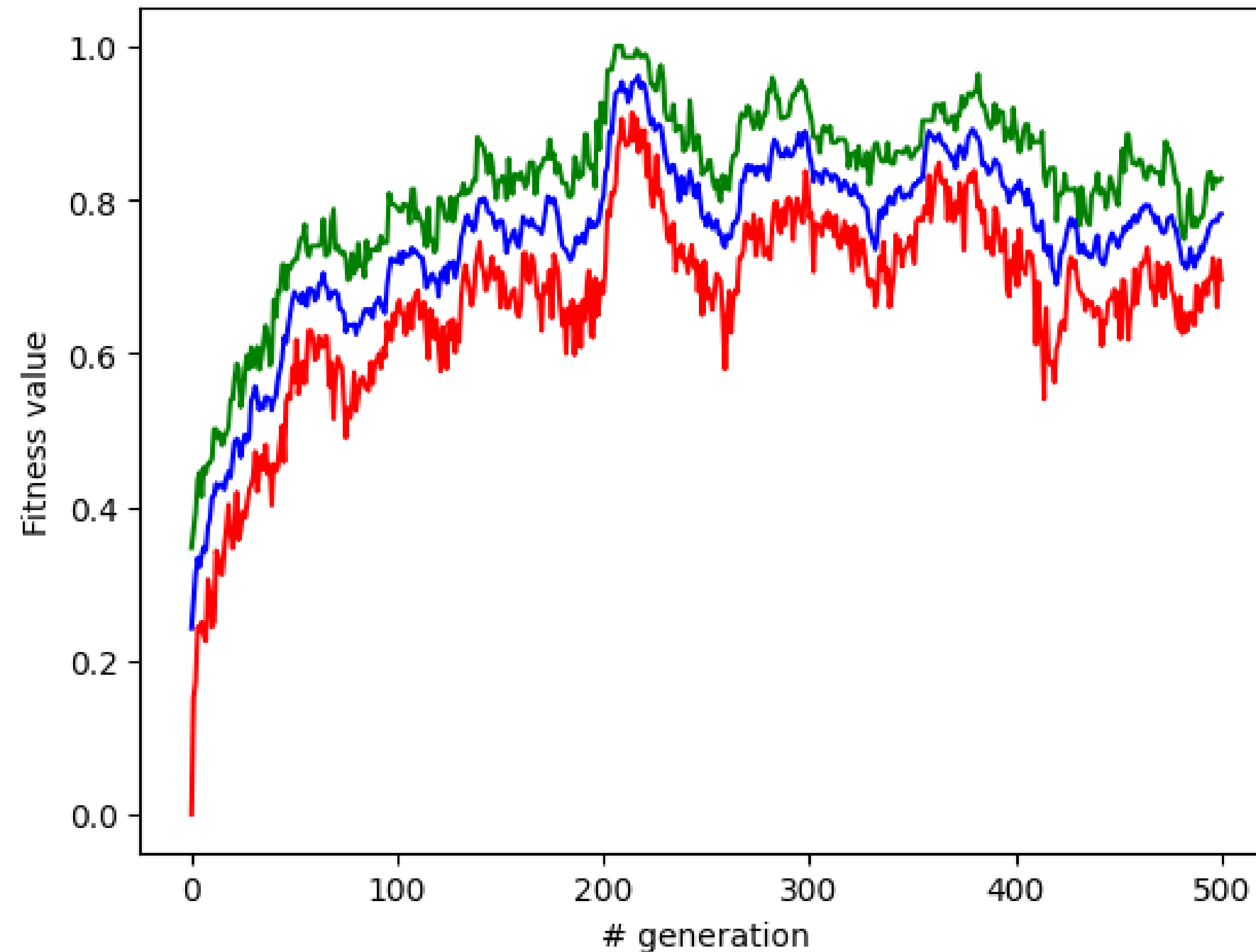
# Larger Graph ( $N = 100$ )

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- Mutation: swap, cross-over: Ox



# Larger Graph (N = 100)

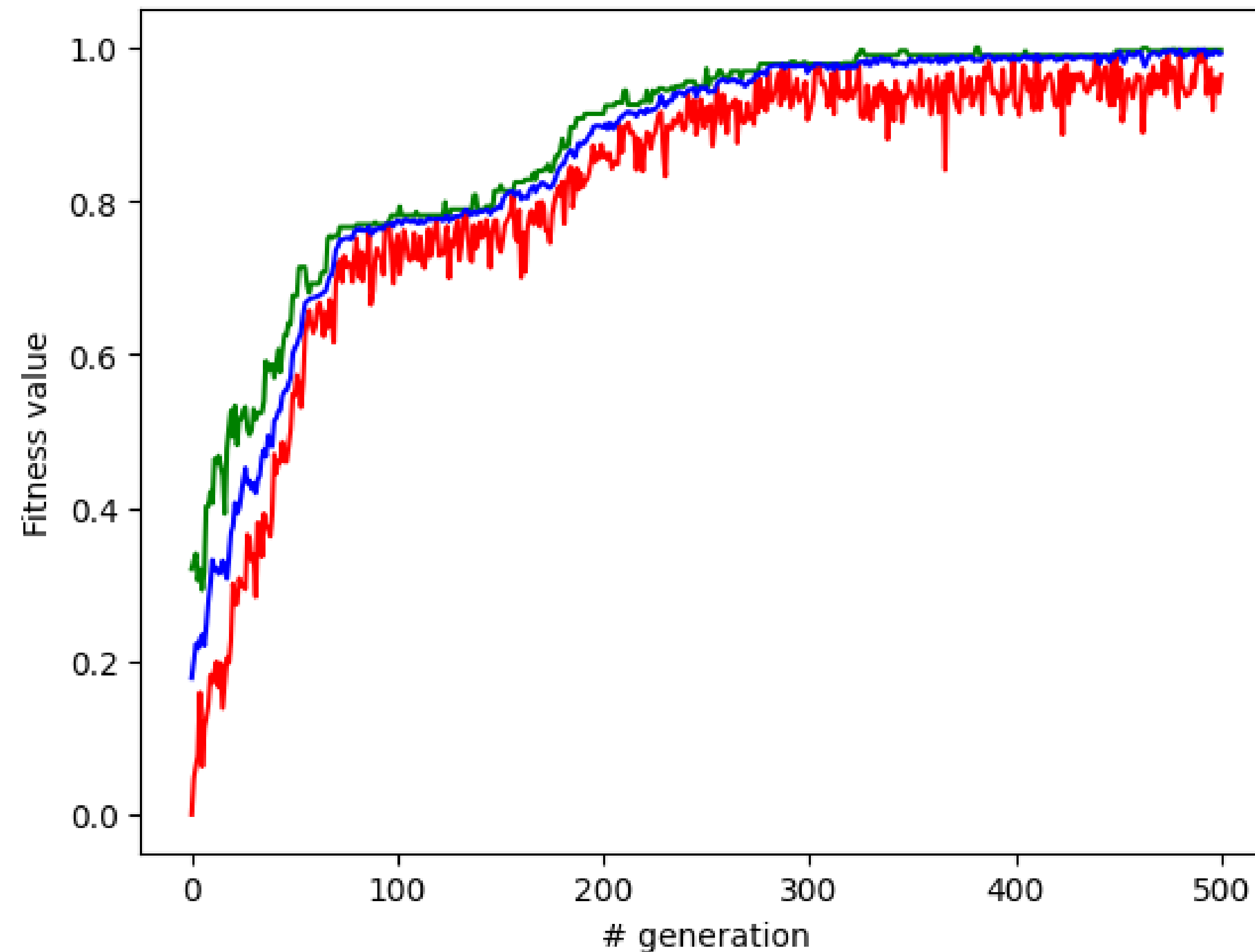
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- Mutation: insertion, cross-over: Ox





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- Mutation: insertion, cross-over: pmx



# Comparison

- The minimum path length found by the different combinations of mutation & cross-over functions are for  $N = 100$

	Swap	Insertion
OX	2652	2830
PMX	2733	2197

# Conclusion & Future Work

- In general PMX cross over seems to be better than OX, especially for large graphs
- This is in contradiction to the claim in paper [2] which claims that OX is better in general.
- There is unlikely to be major difference between swap & insertion mutation
- Running the code takes less than a second  $N = 100$ . Could consider larger graphs.
- Could consider different parameter values for pmutate, pcross, population size etc.
- Could introduce elitism
- Overall Genetic algorithm converged & found good solutions to Travelling salesman problem for varying graph sizes.



# References

- [1] An Improved Genetic Algorithm for Solving the Traveling Salesman Problem by Pen Chang
- [2] Genetic algorithms for the traveling salesman problem Jean-Yves Potvin