

APPENDIX A:

TSP SOLVER USING GENETIC ALGORITHM

The outline of the GA for TSP

The algorithm consists of the following steps:

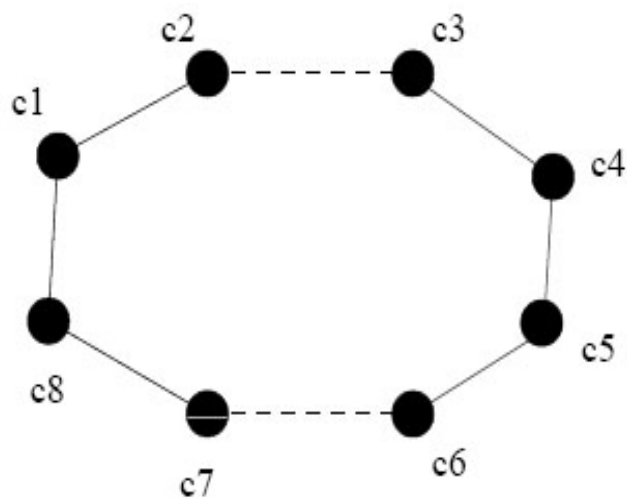
- **Initialization:** Generation of M individuals randomly.
- **Natural Selection:** Eliminate $p_1\%$ individuals. The population decreases by $M.p_1/100$.
- **Multiplication:** Choose $M.p_1/100$ pairs of individuals randomly and produce an offspring from each pair of individuals (by crossover). The population reverts to the initial population M .
- **Mutation by 2-opt:** choose $p_2\%$ of individuals randomly and improve them by the 2-opt method. The elite individual (the individual with the best fitness value in the population) is always chosen. If the individual is already improved, do nothing.

Representation

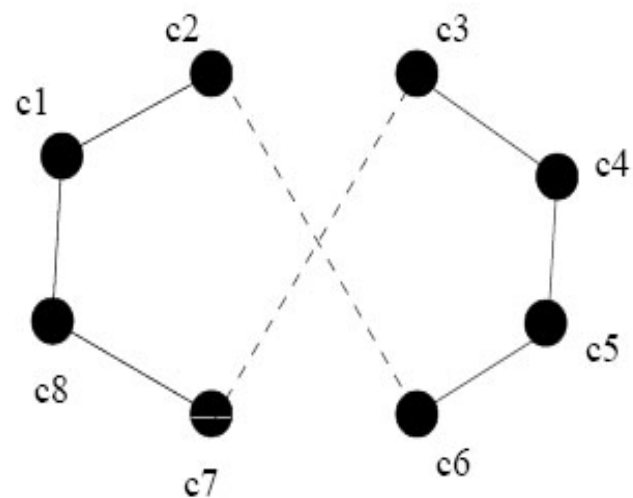
- We use the *path representation* for solution coding.
- EX: the chromosome $g = (D, H, B, A, C, F, G, E)$ means that the salesperson visits D, H, B, A,...E successively, and returns to town D.

Mutation by 2-opt

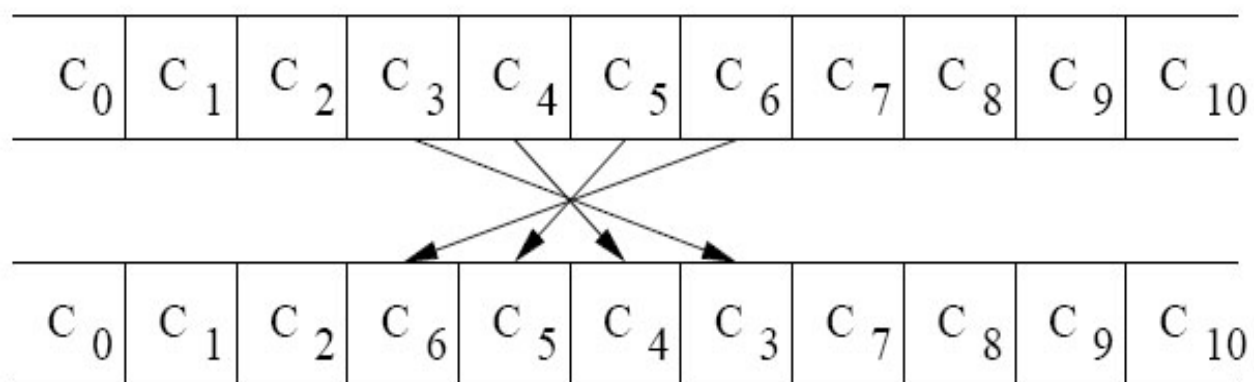
- The **2-opt** is one of the most well-known local search operator (move operator) among TSP solving algorithms.
- It improves the tour edge by edge and reverses the order of the subtour.
- When we apply the 2-opt mutation to a solution, the solution may fall into a local minimum.



(a) The current tour.



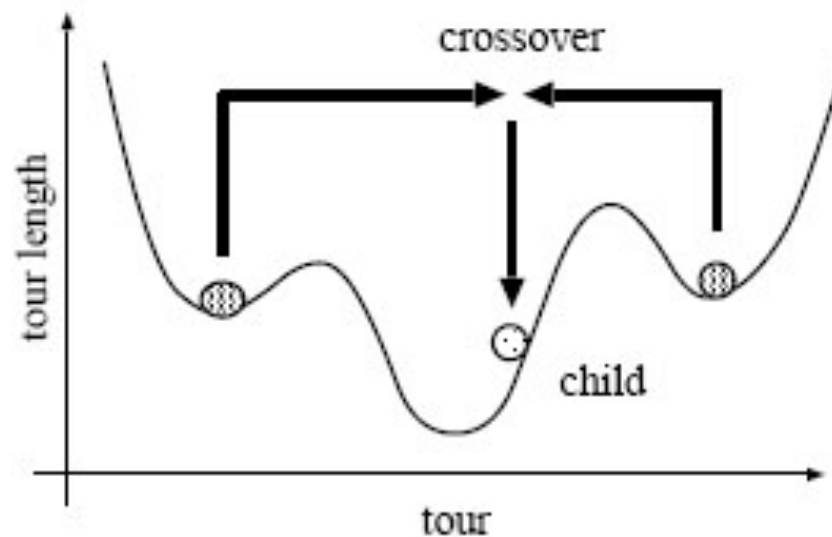
(b) The proposed tour by 2OPT.



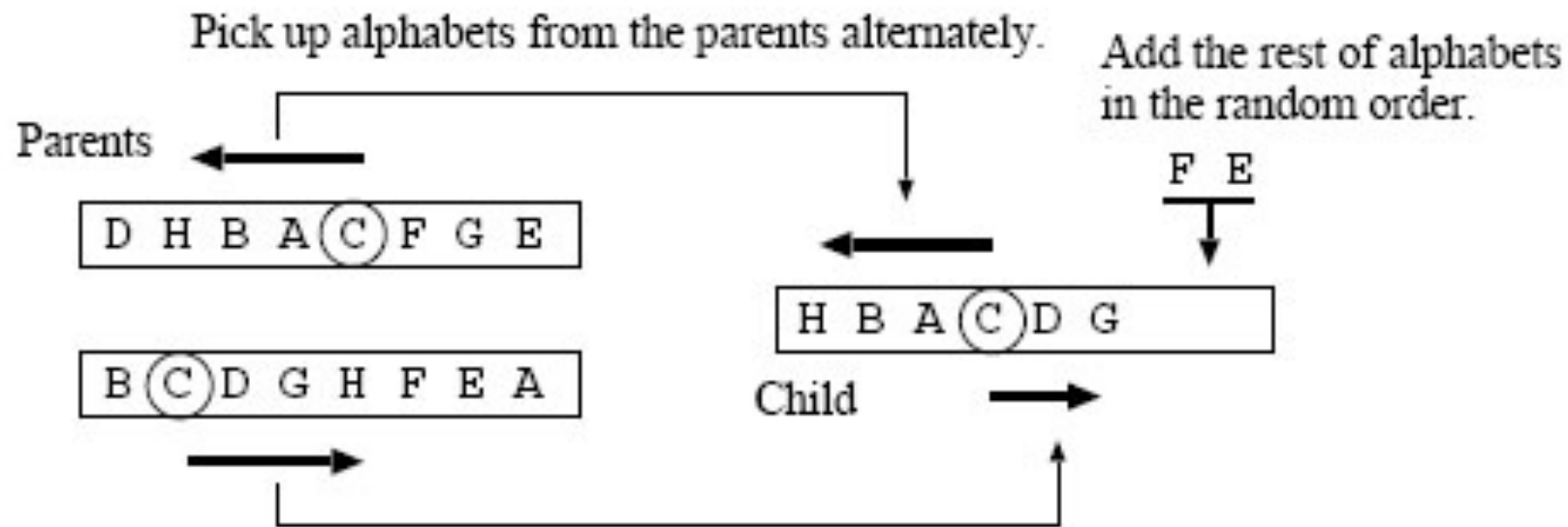
(c) The array structure.

Crossover operator

- Greedy Subtour Crossover (GSX). It acquires the longest possible sequence of parents' subtours.
- Using GSX, the solution can pop up from local minima effectively.



Greedy Subtour Crossover



Inputs: Chromosomes $g_a = (D, H, B, A, \underline{C}, F, G, E)$ and $g_b = (B, \underline{C}, D, G, H, F, E, A)$.

Outputs: The offspring $g = (H, B, A, \underline{C}, D, G, F, E)$

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procedure crossover( $g_a, g_b$ ) {
     $f_a \leftarrow \text{true}$ 
     $f_b \leftarrow \text{true}$ 
    choose town  $t$  randomly
    choose  $x$ , where  $a_x = t$ 
    choose  $y$ , where  $b_y = t$ 
     $g \leftarrow t$ 
    do {
         $x \leftarrow x - 1 \pmod{n}$ ,
         $y \leftarrow y + 1 \pmod{n}$ .
        if  $f_a = \text{true}$  then {
            if  $a_x \notin g$  then
                 $g \leftarrow a_x.g$ 
            else  $f_a \leftarrow \text{false}$ 
        }
        if  $f_b = \text{true}$  then {
            if  $b_x \notin g$  then
                 $g \leftarrow g.b_y$ 
            else  $f_b \leftarrow \text{false}$ 
        }
    } while  $f_a = \text{true}$  or  $f_b = \text{true}$ 

```

Algorithm: Greedy Subtour Crossover

Inputs: Chromosomes $g_a = (a_0, a_1, \dots, a_{n-1})$ and $g_b = (b_0, b_1, \dots, b_{n-1})$.

Outputs: The offspring chromosome g .

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if  $|g| < |g_a|$  then {
    add the rest of towns to  $g$  in the
    random order
}
return  $g$ 

```


Example:

- Suppose that parent chromosomes $g_a = (D, H, B, A, C, F, G, E)$ and $g_b = (B, C, D, G, H, F, E, A)$.
- First, choose one town at random, say, town C is chosen. Then $x = 4$ and $y = 1$. Now the child is (C).
- Next, pick up towns from the parents alternately. Begin with a_3 (A) and next is b_2 (D). The child becomes $g = (A, C, D)$.
- In the same way, add a_2 (B), b_3 (G), a_1 (H), and the child becomes $g = (H, B, A, C, D, G)$.
- Now the next town is $b_4 = H$ and H has already appeared in the child, so we can't add any more towns from parent g_b .
- Now, we add towns from parent g_a . The next town is $a_0 = D$, but D has already used. Thus, we can't add towns from parent g_a , either.
- Then we add the rest of the towns, i.e., E and F, to the child in the random order. Finally the child is $g = (H, B, A, C, D, G, F, E)$.

Survivor Selection

- Eliminate $R = M.p_1/100$. We eliminate similar individuals to maintain the diversity in order to avoid immature convergence.
- First, sort the individuals in fitness-value order. Compare the fitness value of adjoining individuals. If the difference is less than ε , eliminate preceding individual while the number of eliminated individuals (r) is less than R .
- Next, if $r < R$, eliminate $R - r$ individuals in the order of lowest fitness value.

Other parameters

- Test data: TSPLIB
- gr96.data (666 cities)
 - Population size: 200
 - maximum number of generations: 300
 - $p_1 = 30\%$
 - $p_2 = 20\%$
 - produces the minimum solution
- Conclusion: The solver brings out good solution very fast since GA here utilizes heuristic search (2-opt) in genetic operators. It is a hybrid algorithm.

Reference

- H. Sengoku, I. Yoshihara, “A Fast TSP Solver Using GA on Java”, 1997.