

heavy penalty will be applied for each overlapping block pair. For any two block pairs A and B (Figure 3), one possible way to figure out the overlap is by checking the following condition:

$$\text{overlap} = \text{not} (\\ A_right \leq B_left \text{ or} \\ A_left \geq B_right \text{ or} \\ A_bottom \geq B_top \text{ or} \\ A_top \leq B_bottom \\)$$

where right, left, and top, bottom can be derived from the bottom-left x,y coordinate values of both blocks and the width and height of the corresponding blocks, respectively. The idea is that no overlap exists if one rectangle is completely to the left, right, above, **or** below the other.

Task 1 Instructions

Start with a random initial population of 6 members, where each chromosome represents a candidate layout. Suppose the layout is specified by all the **bottom-left coordinates** of the 6 components (outlined in section 2 in that exact order). Check section 7 for a sample input analysis.

1. Chromosome Representation / Encoding:

- a. Devise a **suitable encoding scheme** to represent all the block components specified in the question. Note that the placement coordinate values will be in the range of [0, 25]. Be as concise as possible in your encoding; that is, try not to keep redundant information in your encoded version.
- b. Then generate an initial population of 6 chromosomes to start the GA loop, where each chromosome represents a placement strategy.

2. Fitness Function Implementation:

- a. Based on the 3 fitness objectives outlined above, design a **suitable fitness function** that focuses on penalizing bad placement combinations. Implementation hints are outlined in detail in section 4. Follow a prioritized order of these objectives as mentioned below, and reflect that in your fitness calculation.
 - i. Overlap counts should be considered the least desirable, hence penalized way more than the other two
 - ii. The total wiring distance and the total bounding area should be considered next

- b. Calculate the fitness of the chromosome pool using the hints specified above. A **weighted sum** can be a good fitness function to start with. Check the sample input output section for some ideas to start with.
- 3. Parent Selection:**
 - a. Choose two parents based on **random selection** at each generation. Tournament selection, Roulette Wheel selection are some of the other popular selection techniques.
- 4. Cross-over:**
 - a. Perform **single-point crossover** to create 2 new offspring from each pair of selected parents.
 - b. To achieve this, pick a random split point from the chromosome representation of the parents and swap the complementary pieces to generate offspring.
- 5. Mutation:**
 - a. Mutation ensures genetic diversity and prevents the algorithm from getting stuck in local optima.
 - b. Implement the following mutation strategy to introduce random changes: pick any arbitrary component block from the chromosome representation and introduce **random coordinate (i.e., the value of x and y) changes** to that single particular block with a low probability (**5-10% mutation rate**).
- 6. New Generation Creation:**
 - a. Remember, the population size should be the same for all generations. Apply **elitism** to select new individuals (allow 1 or 2 elites to be carried forward to the next generation). Select the remaining candidates from the created offspring for the next generation.
- 7. GA Loop:**
 - a. Run genetic algorithms on such a population until the **best fitness** (a plateau) is achieved or the maximum number of **15 iterations** is reached. Keep this generation count as a variable input for further experiments.
- 8. Required Output:**
 - a. Output the best possible placement strategy found once the algorithm halts. Deliverables include the **best total fitness value**, the corresponding **total wiring length**, **the total bounding box area**, and **the total overlap counts**. Also, decode the best chromosome representation to reflect the optimal placement of bottom-left coordinates.

Task 2

Take a look at **Step 4 of Task 1**. Can you incorporate the **Two-point crossover mechanism** into it?

- To implement this task, randomly select two parents from your initial population. Then perform a two-point crossover to generate two children. The two points have to be chosen randomly, but it has to be ensured that the second point always comes after the first point.

Sample Input and Output

A sample initial population set of 6 chromosomes, where each line below represents the bottom-left position coordinate of 6 chip component blocks:

P1 → (9,3), (12, 15), (13, 16), (1,13), (4,15), (9, 6)

P2 → (8, 0), (7,12), (4,11), (1,13), (14,10), (9,11)

P3 → (6, 5), (12, 9), (9, 7), (8, 6), (2, 7), (3, 1)

P4 → (3,11), (11, 12), (14, 11), (6, 10), (3,11), (3,0)

P5 → (10, 12) (8, 16), (10, 4), (13, 6), (6, 0), (3, 7)

P6 → (0, 2), (0, 0), (14, 12), (4, 5), (12, 4), (3, 10)

For **P1**,

Pairwise block overlap count = 3

Total wiring distance (center-to-center) of the specified connected pairs = 12.91 + 12.98 + 12.18 + 10.61 + 9.01 + 9.43 = 67.1

Total bounding box area = (x_max - x_min) * (y_max - y_min) = (19-1) * (20-3) = 306

Total fitness value = - (alpha * overlap penalty + beta * wiring length penalty + gamma * bounding area penalty)

Considering, alpha = 1000, beta = 2, gamma = 1:

Fitness for generation 1 chromosome 1 = - (1000 * 3) - (67.1 * 2) - 306 = -3440.23