

# **VISVESVARAYA TECHNOLOGICAL UNIVERSITY**

**“JnanaSangama”, Belgaum -590014, Karnataka.**



## **LAB RECORD**

### **Bio Inspired Systems (23CS5BSBIS)**

*Submitted by*

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*in partial fulfillment for the award of the degree of*

**BACHELOR OF ENGINEERING  
*in*  
COMPUTER SCIENCE AND ENGINEERING**



**B.M.S. COLLEGE OF ENGINEERING**

**(Autonomous Institution under VTU)**

**BENGALURU-560019**

**Sep-2024 to Jan-2025**

**B.M.S. College of Engineering,**  
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(Affiliated To Visvesvaraya Technological University, Belgaum)  
**Department of Computer Science and Engineering**



**CERTIFICATE**

This is to certify that the Lab work entitled “ Bio Inspired Systems (23CS5BSBIS)” carried out by **Shantanu Shrivastav(1BM22CS252)**, who is bonafide student of **B.M.S. College of Engineering**. It is in partial fulfillment for the award of **Bachelor of Engineering in Computer Science and Engineering** of the Visvesvaraya Technological University, Belgaum. The Lab report has been approved as it satisfies the academic requirements of the above mentioned subject and the work prescribed for the said degree.

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Github Link: [https://github.com/Shantanuu3/BIS\\_Lab](https://github.com/Shantanuu3/BIS_Lab)

## Program 1

### Genetic Algorithm for Optimization Problems

DATE: 03/08 PAGE:

#### Genetic Algorithm for Optimization Problems

→ Genetic algorithms are inspired by the process of natural selection and genetics. In GA, a population of potential solution is evolved over generations to find the best solution to a problem. The same idea is to select the fittest individual from the population and use them for reproduction to create the next generation. GAs are widely used for solving optimization and search problems. In this document, we will explore how to implement a genetic algorithm.

Example: Maximizing  $f(x) = x^2$   
 $x \in [0, 1]$   
 $f(x) = x^2$

#### Implementation Steps:

- 1) Define the problem  
The first step is to create a mathematical function that needs to be optimized. This function could represent anything depending on the nature of the problem.



- 2) Initialize Parameters  
Set the necessary parameters for the GA. These include:
  - population size
  - Mutation rate
  - Crossover rate
  - Number of Generation
- 3) Create initial Problem  
Generate an initial problem population of potential solution. This population typically consist of random individuals that represent possible answers.
- 4) Evaluate fitness  
Evaluate the fitness of each individuals in the population. The fitness function is problem-specific and is used to quantify how good a particular solution is.
- 5) Selection  
select individual based on their fitness to reproduce and generate offspring for the next generation.
- 6) Crossover  
Crossover is the process of combining two parent solution.



to create offspring. It mimics biological reproduction where two parent exchange genetic material to produce a child.

#### 7) Mutation

Mutation introduces genetic diversity by making small changes to an individual's structure.

#### 8) Iteration

Repeat the evaluation, selection, crossover, and mutation process.

Applications in Computer Science and Engineering

#### 1) Machine Learning and Neural Networks

- Hyperparameter tuning
- Hyperparameter tuning
- Feature selection

#### 2) Software Engineering

- Automated test case gen
- Refactoring

#### 3) Optimization problem

- Travelling Salesman
- Scheduling



## 4) Robotics :

- Path Planning
- Behavioural Optimization

## 5) Network Design and optimization

- Routing protocols
- Network topology design

## Conclusion :

- OA are a versatile and powerful tool for solving complex optimization problems across many domains in computer science and engineering. Their ability to explore large search spaces

Complete  
Kangaroo Algorithm  
3/10

**Code:**

```
import random

# Set a random seed for reproducibility
random.seed(42)

def fitness(chromosome):
    x = int("".join(map(str, chromosome)), 2)
    return x ** 2

def binary_string_to_chromosome(binary_string):
    return [int(bit) for bit in binary_string]

def generate_population_from_input():
    population = []
    for _ in range(population_size):
        while True:
            binary_string = input("Enter a binary string of size 5 (e.g., '11001'): ")
            if len(binary_string) == 5 and all(bit in '01' for bit in binary_string):
                population.append(binary_string_to_chromosome(binary_string))
                break
            else:
                print("Invalid input. Please enter a binary string of size 5.")
    return population

def select_pair(population, fitnesses):
    total_fitness = sum(fitnesses)
    selection_probs = [f / total_fitness for f in fitnesses]
    parent1 = population[random.choices(range(len(population)), selection_probs)[0]]
    parent2 = population[random.choices(range(len(population)), selection_probs)[0]]
    return parent1, parent2

def crossover(parent1, parent2):
    point = random.randint(1, len(parent1) - 1)
    offspring1 = parent1[:point] + parent2[point:]
    offspring2 = parent2[:point] + parent1[point:]
    return offspring1, offspring2

def mutate(chromosome, mutation_rate):
    return [gene if random.random() > mutation_rate else 1 - gene for gene in chromosome]
```



```

# Parameters
population_size = 4
generations = 20
mutation_rate = 0.01

# Initialize population from user input
population = generate_population_from_input()

for generation in range(generations):
    fitnesses = [fitness(chromosome) for chromosome in population]

    new_population = []

    # Create new population
    while len(new_population) < population_size:
        parent1, parent2 = select_pair(population, fitnesses)
        offspring1, offspring2 = crossover(parent1, parent2)
        new_population.append(mutate(offspring1, mutation_rate))
        new_population.append(mutate(offspring2, mutation_rate))

    # Ensure the new population has the right size
    population = new_population[:population_size]

# Get the maximum fitness
fitnesses = [fitness(chromosome) for chromosome in population]
max_fitness = max(fitnesses)

print(f"Maximum Possible Fitness: {max_fitness}")

```

### Output:

```

Enter a binary string of size 5 (e.g., '11001'): 11011
Enter a binary string of size 5 (e.g., '11001'): 01011
Enter a binary string of size 5 (e.g., '11001'): 11100
Enter a binary string of size 5 (e.g., '11001'): 01101
Maximum Possible Fitness: 841

```

## Program 2

### Ant Colony Optimization

DATE: PAGE:

Ant colony Optimization for travelling salesman problem

→ Ant colony optimization is a nature inspired algorithm based on the foraging behaviour of ants. Specifically the way ants find the shortest path from their colony to a food source. The algorithm is used for solving combinatorial optimization problem including TSP.

→ ACO algorithm for TSP

- 1) Initialization:
  - initialize the pheromone levels  $\tau_{ij}$  for all edges between cities
  - Set the parameters:  $\alpha, \beta$ , number of ants, number of iteration, pheromone
- 2) Ant's tour construction:
  - For each ant, start from a random city
  - at each step, the ant chooses the next city based on the pheromone levels and distance.
  - Repeat until all cities

are visited and the tour is completed.

- 3) Pheromone update
- After all ants have completed their tour, update the pheromones on the edges according to the rule:

$$\tau_{i,j} = (1-\rho)\tau_{i,j} + \Delta\tau_{i,j}$$

where:

- $\rho$  is the pheromone evaporation rate
- $\Delta\tau_{i,j}$  is the amount of pheromone deposited by ants on the edge  $(i,j)$  which is inversely prop. to the length

#### 4) Termination

- If the stopping criteria are met stop the algorithm. Otherwise repeat from step 2



Pseudo code :

initialize pheromone matrix  $\tau[i][j] = 1$   
 initialize best\_tour = empty  
 initialize best\_length = infinity

for situation = 1 to max\_iter  
 initialize all ants tours

for each ant k :

ant\_tour = []

current\_city = random\_city()

ant\_tour.append(current\_city)

while tour is not complete  
 (all cities visited) :

next\_city = choose\_city  
 (current\_city, ant\_tour)

ant\_tour.append(next\_city)

current\_city = next\_city

tour\_length = calculate\_tour\_length(ant\_tour)

if tour\_length < best\_length :

best\_tour = ant\_tour

best\_length = tour\_length

apply pheromone evaporation  
 $\tau[i][j] = (1 - \rho) * \tau[i][j]$

for each ant k :

$\text{ant\_tower} = []$   
 $\text{current\_city} = \text{random\_city}()$   
 $\text{ant\_tower.append}(\text{current\_city})$

// function to choose the next city  
 $\text{probability} = []$   
 $\text{total\_probability} = 0$   
 for each city  $j$  not in  
 visited cities:

$\text{pheromone} = \tau[\text{current\_city}][j]^{\beta}$   
 $\text{total\_pheromone} += \frac{\text{pheromone}}{\text{visibility}}$

for each city  $j$  not in  
 visited cities:

$\text{pheromone} = \tau[\text{current\_city}][j]^{\beta}$   
 $\propto$

$\text{visibility} = (1/\text{distance}[\text{current\_city}][j])^{\beta}$

$\text{probability} = \frac{(\text{pheromone} \cdot \text{visibility})}{\text{total\_pheromone}}$

$\text{probabilities.append}(\text{probability})$   
 $\text{next\_city} = \text{select\_city\_based}$   
 $\text{on\_probability}(\text{probabilities})$   
 $\text{return next\_city}$

$\Delta\tau = \frac{Q}{\text{tower\_length}}$   
 $\tau[i][j] += \Delta\tau$

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 21/11/24

**Code:**

```
import random
import numpy as np
import operator

FUNCTIONS = {'+': operator.add, '-': operator.sub, '*': operator.mul, '/': operator.truediv}
TERMINALS = ['x', 1, 2, 3, 4] # x and constants

def random_gene(length=10):
    return [random.choice(list(FUNCTIONS.keys()) + TERMINALS) for _ in range(length)]

def decode_chromosome(chromosome, x):
    stack = []
    for gene in chromosome:
        if gene in FUNCTIONS: # If it's a function, pop arguments and apply
            if len(stack) < 2: # Avoid errors if stack has fewer than 2 elements
                stack.append(0)
                continue
            b = stack.pop()
            a = stack.pop()
            try:
                result = FUNCTIONS[gene](a, b)
            except ZeroDivisionError:
                result = 1 # Avoid division by zero
            stack.append(result)
        elif gene == 'x':
            stack.append(x)
        else:
            stack.append(gene)
    return stack[0] if stack else 0 # Return top of stack as output

def fitness_function(chromosome, target_function, x_values):
    predictions = [decode_chromosome(chromosome, x) for x in x_values]
    targets = [target_function(x) for x in x_values]
    mse = np.mean([(p - t) ** 2 for p, t in zip(predictions, targets)])
    return mse

def selection(population, fitnesses):
    total_fitness = sum(1 / (f + 1e-6) for f in fitnesses) # Avoid division by zero
```



```

probabilities = [(1 / (f + 1e-6)) / total_fitness for f in fitnesses]
return population[np.random.choice(len(population), p=probabilities)]

def mutate(chromosome, mutation_rate=0.1):
    new_chromosome = chromosome[:]
    for i in range(len(new_chromosome)):
        if random.random() < mutation_rate:
            new_chromosome[i] = random.choice(list(FUNCTIONS.keys()) + TERMINALS)
    return new_chromosome

def crossover(parent1, parent2):
    point = random.randint(1, len(parent1) - 1)
    child1 = parent1[:point] + parent2[point:]
    child2 = parent2[:point] + parent1[point:]
    return child1, child2

def ant_colony_optimization(cost_matrix, n_ants=10, n_iterations=100, evaporation_rate=0.5,
alpha=1, beta=2):
    n_nodes = len(cost_matrix)
    pheromones = np.ones((n_nodes, n_nodes)) # Initialize pheromones

    def calculate_probability(i, j, visited):
        if j in visited:
            return 0
        return (pheromones[i][j] ** alpha) * ((1 / cost_matrix[i][j]) ** beta)

    def construct_solution():
        path = [random.randint(0, n_nodes - 1)]
        while len(path) < n_nodes:
            i = path[-1]
            probabilities = [calculate_probability(i, j, path) for j in range(n_nodes)]
            total = sum(probabilities)
            probabilities = [p / total if total > 0 else 0 for p in probabilities]
            next_node = np.random.choice(range(n_nodes), p=probabilities)
            path.append(next_node)
        path.append(path[0]) # Return to start
        return path

    def path_cost(path):
        return sum(cost_matrix[path[i]][path[i + 1]] for i in range(len(path) - 1))

```

```

best_path = None
best_cost = float('inf')

for iteration in range(n_iterations):
    solutions = [construct_solution() for _ in range(n_ants)]
    costs = [path_cost(solution) for solution in solutions]
    for i, cost in enumerate(costs):
        if cost < best_cost:
            best_cost = cost
            best_path = solutions[i]

    pheromones *= (1 - evaporation_rate) # Evaporation
    for i, solution in enumerate(solutions):
        for j in range(len(solution) - 1):
            pheromones[solution[j]][solution[j + 1]] += 1 / costs[i]

    print(f"Iteration {iteration + 1}: Best Cost = {best_cost}")

print("Best Path:", best_path)
print("Best Cost:", best_cost)

cost_matrix = [
    [0, 2, 2, 5, 7],
    [2, 0, 4, 8, 2],
    [2, 4, 0, 1, 3],
    [5, 8, 1, 0, 2],
    [7, 2, 3, 2, 0]
]
ant_colony_optimization(cost_matrix, n_ants=5, n_iterations=20)

```

### Output:

```

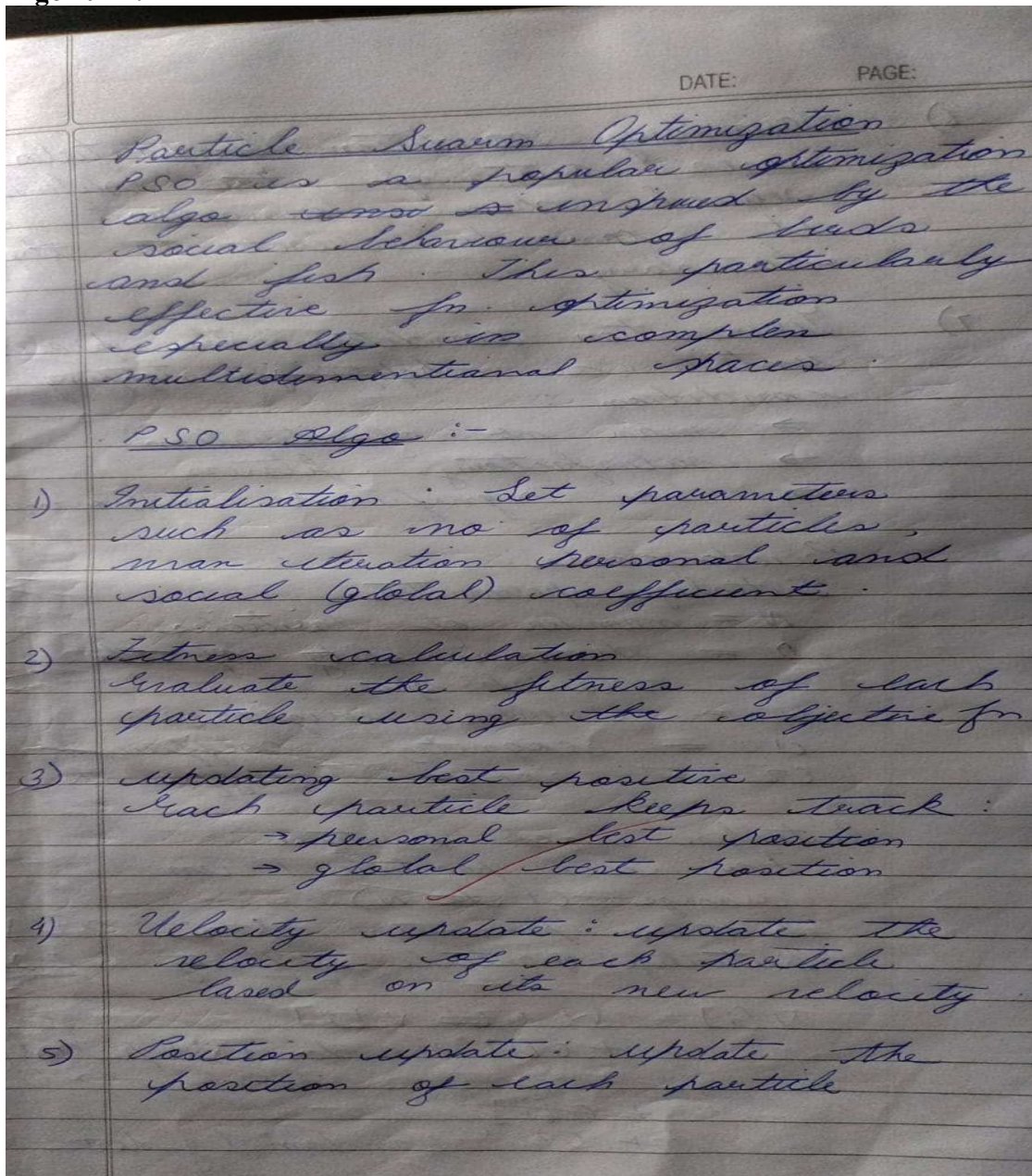
Iteration 15: Best Cost = 9
Iteration 16: Best Cost = 9
Iteration 17: Best Cost = 9
Iteration 18: Best Cost = 9
Iteration 19: Best Cost = 9
Iteration 20: Best Cost = 9
Best Path: [1, 0, 2, 3, 4, 1]
Best Cost: 9

```

### Program 3

#### Particle Swarm Optimization

##### Algorithm:





- 6) Boundary condition: If particles moves out of the defined search space boundaries it can be reset to the boundary
- 7) Termination condition: The process until a stopping condition:
- max no. of iteration
  - satisfactory fitness
  - no significant improvement

Pseudo code:

Initialize parameter

$N$  = number of particles

$D$  = dimensionality of the search space

$W$  = inertia weight

$C1$  = cognitive coefficient

$C2$  = social coefficient

Initialize each particle:

For each particle  $i$  in range (1 to  $N$ ):

Initialize parameter  $pos$ :  $x[i]$

Initialize particle velocity:  $v[i]$

$$v[i] = W * \cancel{v[i]} + C1 * \text{random}() * (p\_best - x[i])$$

$$+ C2 * \text{random}() * (g\_best - x[i])$$

update position

$$x[i] = x[i] + v[i]$$

evaluate fitness

$$\text{fitness}[i] = \text{evaluate}(x[i])$$

update personal best:

$$\text{if } \text{fitness}[i] < \text{fitness}[\text{p\_best}]$$

$$\text{p\_best}[i] = x[i]$$

$$\text{if } \text{fitness}[\text{p\_best}[i]] < \text{fitness}[\text{g\_best}]$$

$$\text{g\_best} = \text{p\_best}[i]$$

return g\\_best as optimal sol

**Code:**

```
import random
import numpy as np
from matplotlib import pyplot as plt
from matplotlib import animation

def fitness_function(x1, x2):
    f1 = x1 + 2 * -x2 + 3
    f2 = 2 * x1 + x2 - 8
    z = f1**2 + f2**2
    return z

def update_velocity(particle, velocity, pbest, gbest, w_min=0.5, max=1.0, c=0.1):
    new_velocity = np.zeros_like(particle)
    r1 = random.uniform(0, max)
    r2 = random.uniform(0, max)
    w = random.uniform(w_min, max)

    for i in range(len(particle)):
        new_velocity[i] = (w * velocity[i] +
                           c * r1 * (pbest[i] - particle[i]) +
                           c * r2 * (gbest[i] - particle[i]))
    return new_velocity

def update_position(particle, velocity):
    new_particle = particle + velocity
    return new_particle

def pso_2d(population, dimension, position_min, position_max, generation, fitness_criterion):
    # Initialization
    particles = np.array([[random.uniform(position_min, position_max) for _ in range(dimension)] for
                           _ in range(population)])
    pbest_position = particles.copy()
    pbest_fitness = np.array([fitness_function(p[0], p[1]) for p in particles])

    gbest_index = np.argmin(pbest_fitness)
    gbest_position = pbest_position[gbest_index]

    velocity = np.zeros((population, dimension))
```



```

images = [] # For animation

for t in range(generation):
    if np.average(pbest_fitness) <= fitness_criterion:
        break

    for n in range(population):
        velocity[n] = update_velocity(particles[n], velocity[n], pbest_position[n], gbest_position)
        particles[n] = update_position(particles[n], velocity[n])

    pbest_fitness = np.array([fitness_function(p[0], p[1]) for p in particles])
    for n in range(population):
        if pbest_fitness[n] < fitness_function(pbest_position[n][0], pbest_position[n][1]):
            pbest_position[n] = particles[n]

    gbest_index = np.argmin(pbest_fitness)
    gbest_position = pbest_position[gbest_index]

    # Plotting the current positions of the particles
    fig = plt.figure(figsize=(10, 10))
    ax = fig.add_subplot(111, projection='3d')
    ax.set_xlabel('x')
    ax.set_ylabel('y')
    ax.set_zlabel('z')

    x = np.linspace(position_min, position_max, 80)
    y = np.linspace(position_min, position_max, 80)
    X, Y = np.meshgrid(x, y)
    Z = fitness_function(X, Y)
    ax.plot_wireframe(X, Y, Z, color='r', linewidth=0.2)

    ax.scatter3D(
        particles[:, 0],
        particles[:, 1],
        [fitness_function(p[0], p[1]) for p in particles],
        c='b'
    )

    # Capture the frame for animation
    plt.title(f'Generation: {t + 1}')
    plt.tight_layout()

```

```

plt.savefig(f'frame_{t}.png')
plt.close(fig)

# Create animation
frames = [plt.imread(f'frame_{i}.png') for i in range(t)]
fig, ax = plt.subplots(figsize=(10, 10))
ax.axis('off')
image = ax.imshow(frames[0])

def update(frame):
    image.set_array(frames[frame])
    return image,

ani = animation.FuncAnimation(fig, update, frames=len(frames), interval=100)
ani.save('./pso_simple.gif', writer='pillow')

# Print the results
print('Global Best Position: ', gbest_position)
print('Best Fitness Value: ', min(pbest_fitness))
print('Average Particle Best Fitness Value: ', np.average(pbest_fitness))
print('Number of Generations: ', t)

# Run the PSO algorithm
pso_2d(population=30, dimension=2, position_min=-10, position_max=10, generation=100,
fitness_criterion=1e-3)

```

### Output:

```

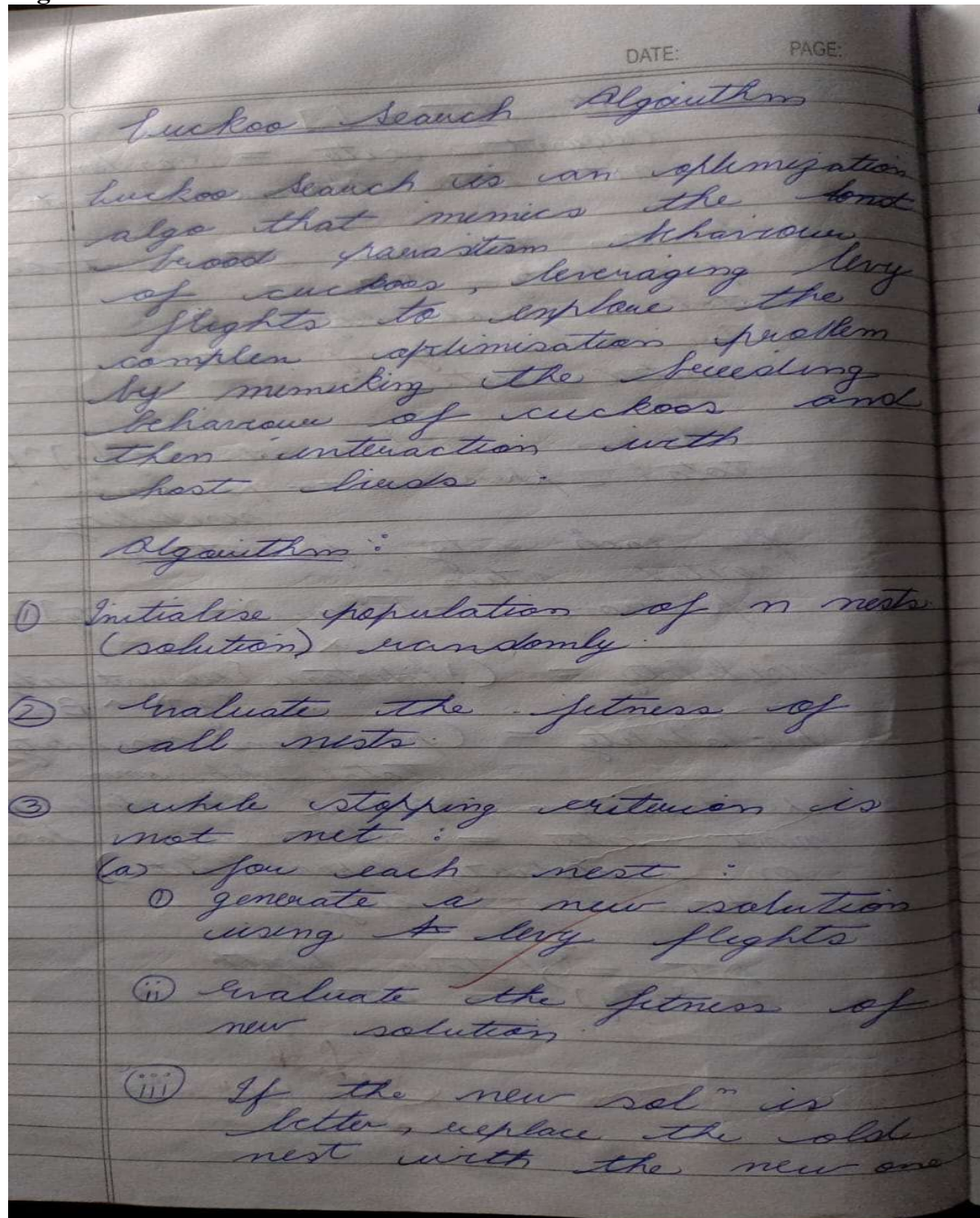
Global Best Position:  [2.59992843  2.79914636]
Best Fitness Value:   3.6691186243893878e-06
Average Particle Best Fitness Value:  0.0007223322365523365
Number of Generations:  45

```

## Program 4

### Cuckoo Search Algorithm

#### Algorithm:



(5) Abandon some nests with probability  $p_a$  and replace them with new random sol<sup>n</sup>.

(4) Return the best sol<sup>n</sup> found

Pseudocode:

① Initialise parameters:

- $N$ : no. of nests (pop<sup>n</sup> size)
- $Max$ : Max<sup>n</sup> number of iterat<sup>n</sup>
- $p_a$ : Probability of discovering alien eggs
- Problem: specific bounds for the search space (lower and upper limits)

② Initialise nests (sol<sup>n</sup>)

- For each nest  $i=1$  to  $N$
- Randomly generate initial sol<sup>n</sup>  $x_i$  within search space

③ Evaluate fitness of each nest

- for each nest  $i=1$  to  $N$
- Evaluate fitness ( $x_i$ ) using the objective funct<sup>n</sup>  $f(x_i)$

④ Set best sol<sup>n</sup> as the initial best sol<sup>n</sup>:

- Set BestSolution =  $\arg \min (fitness(x_i))$



- ⑤ for each iteration ( $t = 1$  to  $\text{max}$ ):
- (a) Generate new sol<sup>n</sup> (nests) using Levy flights:
- $$x-i' = x-i + \alpha \cdot \text{levy flights}$$
- ensure  $x-i'$  stays within the search space
  - if  $x-i$  is out of bounds adjust it to stay within the boundaries
  - evaluate fitness ( $x-i'$ )
- (b) evaluate a new sol<sup>n</sup> with the current nests:
- if  $\text{fitness}(x-i') < \text{fitness}(x-i)$ :
  - Replace the old sol<sup>n</sup>  $x-i$  with the new sol<sup>n</sup>  $x-i'$
- (c) Randomly nests with a prob.  $P_a$  (stochastic discovery of alien eggs)
- for each nests  $i = 1$  to  $N$ :
  - evaluate fitness ( $x - \text{rand}$ )
- (d) Update the best sol<sup>n</sup>:
- if a better sol<sup>n</sup> is found:
  - update best solution to the best solution among all nests.

**Code:**

```
import numpy as np
import matplotlib.pyplot as plt

# Objective function: Rastrigin Function
def rastrigin(x):
    A = 10
    return A * len(x) + sum(xi**2 - A * np.cos(2 * np.pi * xi) for xi in x)

# Lévy flight function for generating random steps
def levy_flight(beta=1.5, dim=2):
    sigma_u = np.power(np.math.gamma(1 + beta) * np.sin(np.pi * beta / 2) / np.math.gamma((1 + beta) / 2) / np.power(2, (beta - 1) / 2), 1 / beta)
    sigma_v = 1
    u = np.random.normal(0, sigma_u, dim)
    v = np.random.normal(0, sigma_v, dim)
    return u / np.power(np.abs(v), 1 / beta)

# Cuckoo Search Algorithm
class CuckooSearch:
    def __init__(self, func, dim, population_size, max_generations, pa=0.25, beta=1.5, lower_bound=-5, upper_bound=5):
        self.func = func          # Objective function
        self.dim = dim            # Dimension of the problem
        self.population_size = population_size # Number of nests (solutions)
        self.max_generations = max_generations # Maximum number of generations
        self.pa = pa              # Probability of alien eggs (nest replacement)
        self.beta = beta          # Lévy flight exponent
        self.lower_bound = lower_bound # Lower bound of the search space
        self.upper_bound = upper_bound # Upper bound of the search space

        # Initialize population (nests)
        self.nests = np.random.uniform(self.lower_bound, self.upper_bound, (self.population_size, self.dim))
        self.fitness = np.array([self.func(nest) for nest in self.nests]) # Fitness of each nest
        self.best_nest = self.nests[np.argmin(self.fitness)] # Best solution found
        self.best_fitness = np.min(self.fitness) # Best fitness value

    # Update nests using Lévy flights and objective function evaluations
    def generate_new_nests(self):
```

```

new_nests = []
for i in range(self.population_size):
    step = levy_flight(self.beta, self.dim)
    new_nest = self.nests[i] + step
    # Apply boundary check
    new_nest = np.clip(new_nest, self.lower_bound, self.upper_bound)
    new_nests.append(new_nest)
return np.array(new_nests)

# Main cuckoo search algorithm
def search(self):
    history = [] # To record the best fitness values over generations

    for generation in range(self.max_generations):
        # Generate new nests based on Lévy flight
        new_nests = self.generate_new_nests()
        new_fitness = np.array([self.func(nest) for nest in new_nests])

        # Replace nests with new ones if they are better
        for i in range(self.population_size):
            if new_fitness[i] < self.fitness[i] or np.random.rand() < self.pa:
                self.nests[i] = new_nests[i]
                self.fitness[i] = new_fitness[i]

        # Find the best nest in the current population
        current_best_fitness = np.min(self.fitness)
        current_best_nest = self.nests[np.argmin(self.fitness)]

        # Update the global best solution
        if current_best_fitness < self.best_fitness:
            self.best_fitness = current_best_fitness
            self.best_nest = current_best_nest

        # Record the best fitness for the current generation
        history.append(self.best_fitness)
        print(f'Generation {generation+1}: Best fitness = {self.best_fitness}')

    return self.best_nest, self.best_fitness, history

# Analyze the Cuckoo Search Algorithm
def analyze_cuckoo_search():

```



```
# Set up parameters for Cuckoo Search
dim = 2
population_size = 50
max_generations = 100
cuckoo_search = CuckooSearch(func=rastrigin, dim=dim, population_size=population_size,
max_generations=max_generations)

# Run the Cuckoo Search algorithm
best_nest, best_fitness, history = cuckoo_search.search()

# Plot the convergence curve
plt.plot(history)
plt.title("Convergence Curve of Cuckoo Search Algorithm")
plt.xlabel("Generation")
plt.ylabel("Best Fitness")
plt.show()

print(f"Best solution found: {best_nest}")
print(f"Best fitness: {best_fitness}")

# Run the analysis
analyze_cuckoo_search()
```

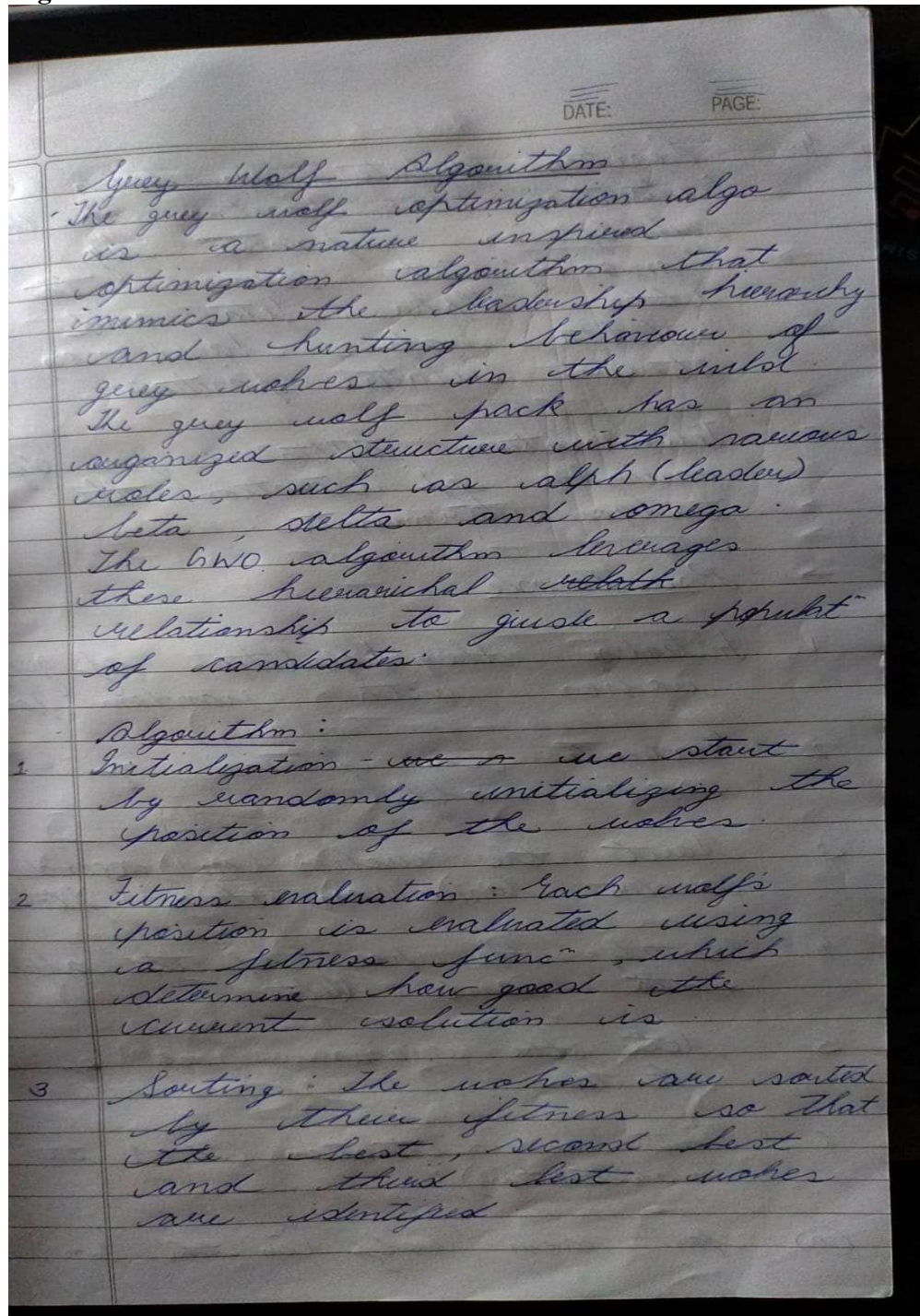
**Output:**

```
Best solution found: [1.30548027 2.02026344]
Best fitness: 0.16306139523513963
```

## Program 5

### Grey Wolf Optimizer

#### Algorithm:



4. Position update: in each iteration the position of each wolf is updated by "moving" it closer to the alpha, beta and delta wolves based on their relative position.

5. Termination: After a pre defined number of iteration the algo terminates.

→ Pseudocode

1. Initialise the population of grey wolves with random pos.

2. Evaluate the fitness of each wolf.

3. Sort the wolves in ascending order based on their fitness.

4. Identify the pos. of the alpha, beta and delta.

5. Set the maximum no. of iteration.

6. For each iteration ( $t=1$  to maxIteration):

(a) for each wolf in the pos.



- (ii) update the position of the wolf based on their pos<sup>n</sup> of the alpha, beta, delta
- (iii) calculate the new fitness of the wolf.
- (iv) if the new fitness is better than previous
5. update the pos<sup>n</sup> of alpha, beta and delta wolves
- (v) Sort the wolves again based on their fitness.
7. Return the pos<sup>n</sup> of the alpha wolf as the optimal solution.

**Code:**

```
import numpy as np

def objective_function(x):
    return np.sum(x**2)

class GreyWolfOptimizer:
    def __init__(self, objective_function, n_wolves, n_variables, max_iter, lb, ub):
        self.obj_func = objective_function # Objective function
        self.n_wolves = n_wolves # Number of wolves
        self.n_variables = n_variables # Number of variables in the problem
        self.max_iter = max_iter # Maximum number of iterations
        self.lb = lb # Lower bound for the search space
        self.ub = ub # Upper bound for the search space

        self.wolves = np.random.uniform(self.lb, self.ub, (self.n_wolves, self.n_variables))

        self.alpha = np.zeros(self.n_variables)
        self.beta = np.zeros(self.n_variables)
        self.delta = np.zeros(self.n_variables)
        self.alpha_score = float("inf")
        self.beta_score = float("inf")
        self.delta_score = float("inf")

    def update_wolves(self):
        fitness = np.apply_along_axis(self.obj_func, 1, self.wolves)

        sorted_indices = np.argsort(fitness)
        self.wolves = self.wolves[sorted_indices]
        fitness = fitness[sorted_indices]

        # Update alpha, beta, and delta wolves
        self.alpha = self.wolves[0]
        self.beta = self.wolves[1]
        self.delta = self.wolves[2]
        self.alpha_score = fitness[0]
        self.beta_score = fitness[1]
        self.delta_score = fitness[2]

    def optimize(self):
```

```

    for t in range(self.max_iter):
        A = 2 * np.random.random((self.n_wolves, self.n_variables)) - 1 # Random values for
exploration
        C = 2 * np.random.random((self.n_wolves, self.n_variables)) # Random values for
exploitation
        for i in range(self.n_wolves):
            D_alpha = np.abs(C[i] * self.alpha - self.wolves[i]) # Distance to alpha wolf
            D_beta = np.abs(C[i] * self.beta - self.wolves[i]) # Distance to beta wolf
            D_delta = np.abs(C[i] * self.delta - self.wolves[i]) # Distance to delta wolf

            self.wolves[i] = self.alpha - A[i] * D_alpha

            self.wolves[i] = np.clip(self.wolves[i], self.lb, self.ub)

        self.update_wolves()

        print(f'Iteration {t+1}/{self.max_iter}, Best Score: {self.alpha_score}')

    return self.alpha, self.alpha_score # Return the best solution found

n_wolves = 30 # Number of wolves
n_variables = 5 # Number of decision variables
max_iter = 100 # Maximum number of iterations
lb = -10 # Lower bound of the search space
ub = 10 # Upper bound of the search space

gwo = GreyWolfOptimizer(objective_function, n_wolves, n_variables, max_iter, lb, ub)
best_solution, best_score = gwo.optimize()
print("Best Solution Found:", best_solution)
print("Best Score:", best_score)

```

### Output:

```

Iteration 100/100, Best Score: 1.985808550535119e-30
Best Solution Found: [-4.38373504e-17 -4.54363691e-16 -1.31663573e-15 -2.05502414e-16
4.09828696e-17]
Best Score: 1.985808550535119e-30

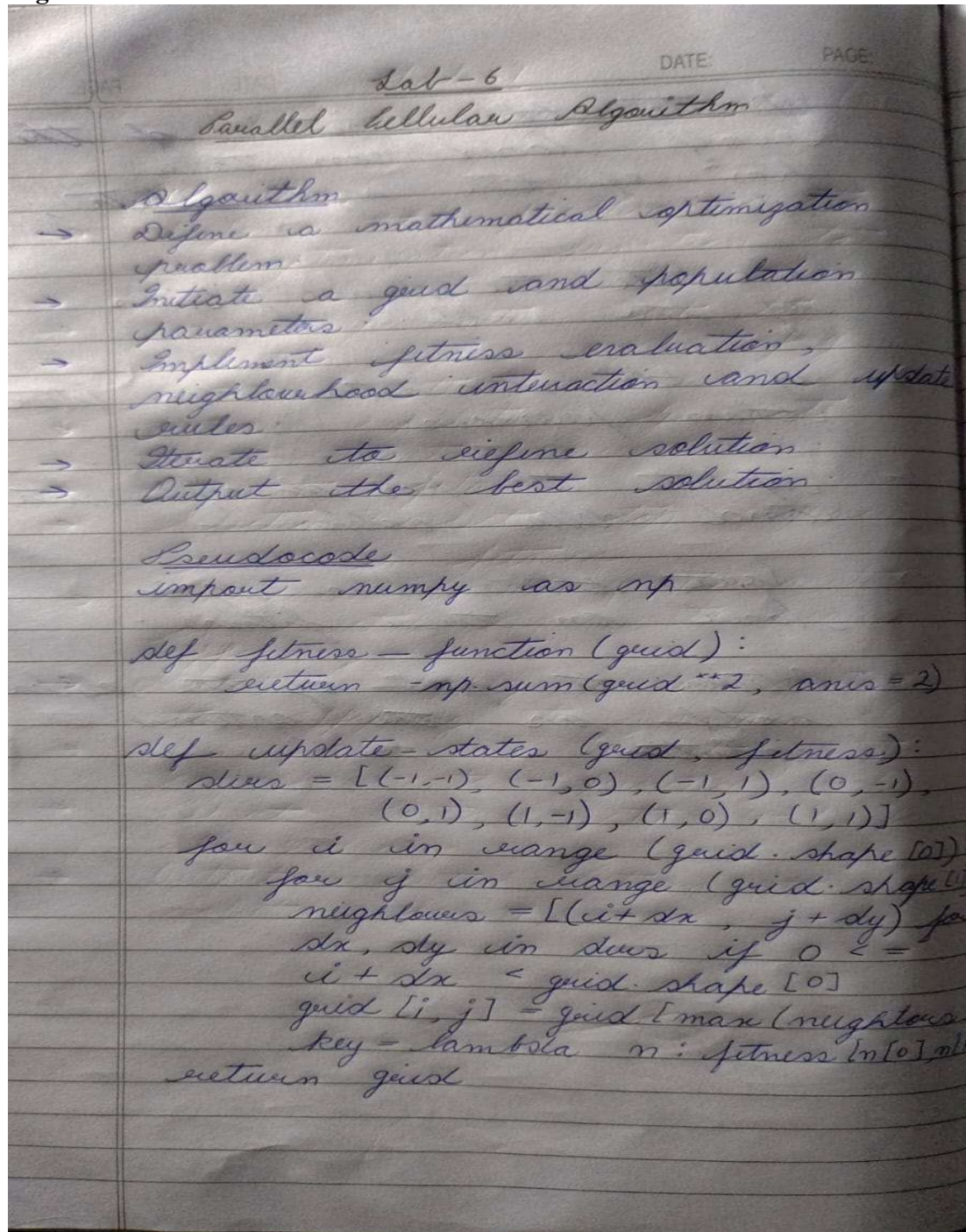
```



## Program 6

### Parallel Cellular Algorithm

Algorithm:



```

def parallel_cellular_algorithm (guid
size, space, iteration):
    guid = np.random.uniform (space[0],
space[1]), (guid_size, guid_size, 2)
    for i in range (iteration):
        fitness = fitness_function (guid)
        guid = update_state (guid, fitness)
    return guid [np.unravel_index
(np.argmax (fitness), fitness.shape),
np.max (fitness)]

```

# Example

```

best_solution, best_fitness = parallel_
cellular_algorithm (10, (-10, 10), 100)
print ("Best Solution: {best_solution}
Fitness: {best_fitness}")

```

**Code:**

```
import numpy as np
from multiprocessing import Pool

def update_cell(cell_index, grid, size):
    x, y = cell_index
    neighbors = [
        ((x-1) % size, y), ((x+1) % size, y),
        (x, (y-1) % size), (x, (y+1) % size)
    ]
    new_state = sum(grid[n[0], n[1]] for n in neighbors) % 2 # example: majority rule
    return (x, y, new_state)

def parallel_update(grid, size, num_iterations):
    pool = Pool(processes=4)
    for iteration in range(num_iterations):
        print(f"Iteration {iteration + 1}:")
        indices = [(x, y) for x in range(size) for y in range(size)]
        result = pool.starmap(update_cell, [(i, grid, size) for i in indices])

        for x, y, new_state in result:
            grid[x, y] = new_state
        print(grid)
    return grid

grid_size = 10
grid = np.random.randint(2, size=(grid_size, grid_size))
print("Initial state:")
print(grid)
num_iterations = 2
updated_grid = parallel_update(grid, grid_size, num_iterations)
```

**Output:**

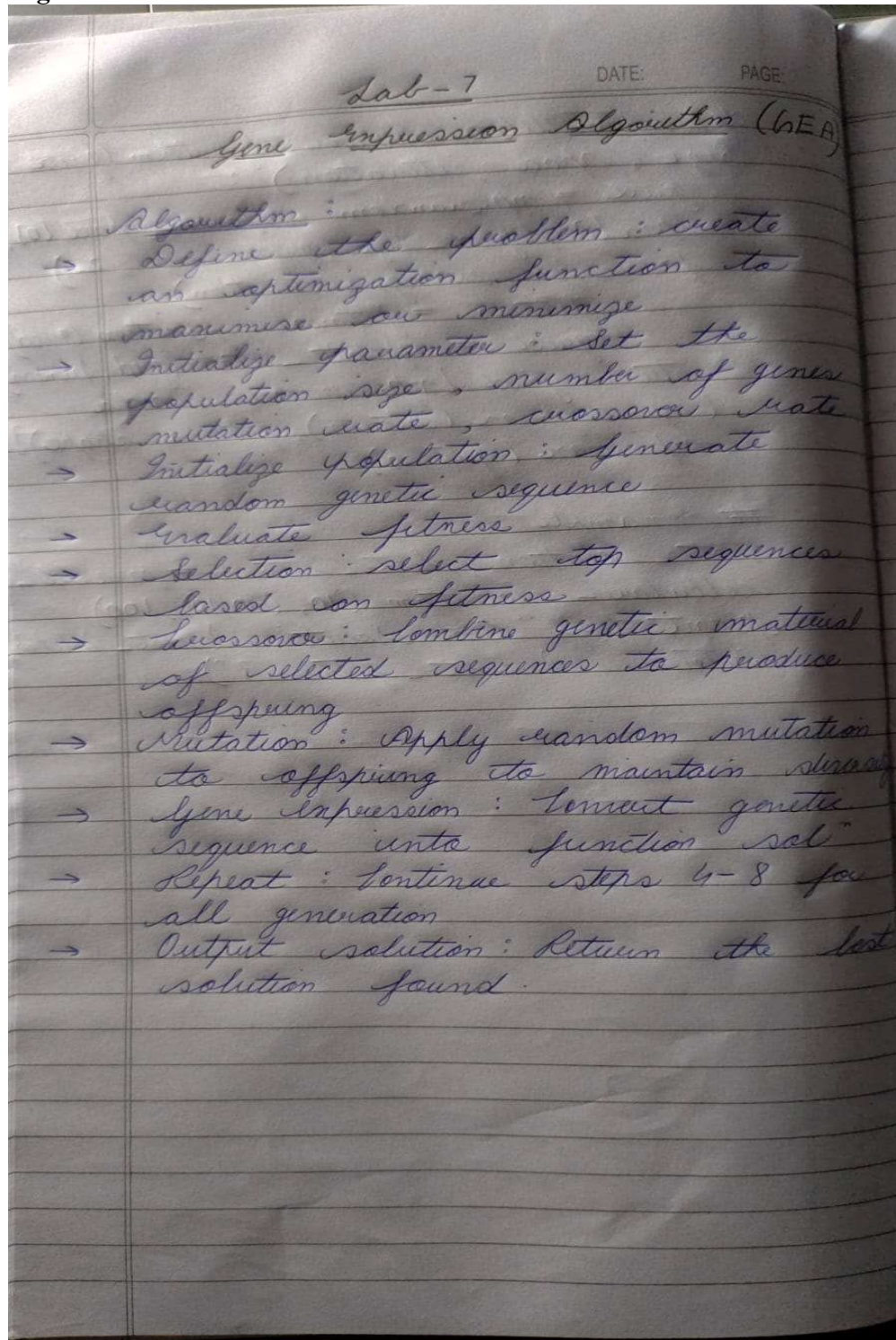
```
Iteration 1:
[[1 0 0 1]
 [1 0 1 0]
 [1 0 0 1]
 [0 1 0 1]]
Iteration 2:
[[0 0 0 0]
 [0 0 0 0]
 [0 0 0 0]
 [0 0 0 0]]
```



## Program 7

### Gene Expression Algorithm

#### Algorithm:



Pseudocode

import numpy as np

```
def fitness_function(solution):
    return -(solution**2).sum()
```

```
def initialize_population(pop_size,
                           num_genes, space):
    return np.random.uniform
    (space[0], space[1], (pop_size, num_genes))
```

```
def gene_expression_algorithm
(pop_size, num_genes, space,
 generation, mutation_rate, crossover_rate):
    population = initialize_population
    (pop_size, num_genes, space)
    for i in range(generation):
        fitness = -(population**2).sum(axis=1)
        parents = population[np.argsort
        (fitness)[-int(pop_size*crossover
        rate):]]
        offspring = parents[np.random.randint
        (len(parents), size=pop_size -
        len(parents))]
        population = np.vstack((parents, offspring))
        best_idx = np.argmin(-(population**2).
        sum(axis=1))
        return population[best_idx],
        (population[best_idx]**2).sum()
```

**Code:**

```
import random
import numpy as np
import operator

# Function set and terminal set
FUNCTIONS = {'+': operator.add, '-': operator.sub, '*': operator.mul, '/': operator.truediv}
TERMINALS = ['x', 1, 2, 3, 4] # x and constants

def random_gene(length=10):
    """Generate a random chromosome (gene)."""
    return [random.choice(list(FUNCTIONS.keys()) + TERMINALS) for _ in range(length)]

def decode_chromosome(chromosome, x):
    """Decode chromosome into a functional expression tree (phenotype)."""
    stack = []
    for gene in chromosome:
        if gene in FUNCTIONS: # If it's a function, pop arguments and apply
            if len(stack) < 2: # Avoid errors if stack has fewer than 2 elements
                stack.append(0)
                continue
            b = stack.pop()
            a = stack.pop()
            try:
                result = FUNCTIONS[gene](a, b)
            except ZeroDivisionError:
                result = 1 # Avoid division by zero
            stack.append(result)
        elif gene == 'x':
            stack.append(x)
        else:
            stack.append(gene)
    return stack[0] if stack else 0 # Return top of stack as output

def fitness_function(chromosome, target_function, x_values):
    """Calculate fitness based on Mean Squared Error."""
    predictions = [decode_chromosome(chromosome, x) for x in x_values]
```



```

targets = [target_function(x) for x in x_values]
mse = np.mean([(p - t) ** 2 for p, t in zip(predictions, targets)])
return mse

def selection(population, fitnesses):
    """Select individuals based on fitness (roulette wheel selection)."""
    total_fitness = sum(1 / (f + 1e-6) for f in fitnesses) # Avoid division by zero
    probabilities = [(1 / (f + 1e-6)) / total_fitness for f in fitnesses]
    return population[np.random.choice(len(population), p=probabilities)]

def mutate(chromosome, mutation_rate=0.1):
    """Apply mutation to a chromosome."""
    new_chromosome = chromosome[:]
    for i in range(len(new_chromosome)):
        if random.random() < mutation_rate:
            new_chromosome[i] = random.choice(list(FUNCTIONS.keys()) + TERMINALS)
    return new_chromosome

def crossover(parent1, parent2):
    """Perform one-point crossover between two parents."""
    point = random.randint(1, len(parent1) - 1)
    child1 = parent1[:point] + parent2[point:]
    child2 = parent2[:point] + parent1[point:]
    return child1, child2

def gene_expression_algorithm(target_function, x_values, population_size=10, generations=20):
    """Main Gene Expression Algorithm."""
    # Initialize random population
    population = [random_gene() for _ in range(population_size)]

    print("Initial Population:")
    for i, chrom in enumerate(population):
        print(f"Chromosome {i}: {chrom}")

    for generation in range(generations):
        print(f"\nGeneration {generation + 1}:")
        # Calculate fitness for each individual

```

```

fitnesses = [fitness_function(chrom, target_function, x_values) for chrom in population]
for i, (chrom, fit) in enumerate(zip(population, fitnesses)):
    print(f'Chromosome {i}: {chrom}, Fitness: {fit:.4f}')

# Select the next generation
new_population = []
for _ in range(population_size // 2):
    parent1 = selection(population, fitnesses)
    parent2 = selection(population, fitnesses)
    child1, child2 = crossover(parent1, parent2)
    child1 = mutate(child1)
    child2 = mutate(child2)
    new_population.extend([child1, child2])
population = new_population

# Final results
print("\nFinal Population and Fitness:")
fitnesses = [fitness_function(chrom, target_function, x_values) for chrom in population]
for i, (chrom, fit) in enumerate(zip(population, fitnesses)):
    print(f'Chromosome {i}: {chrom}, Fitness: {fit:.4f}')

best_index = np.argmin(fitnesses)
print("\nBest Solution:")
print(f'Chromosome: {population[best_index]}, Fitness: {fitnesses[best_index]:.4f}')

# Target function for regression
def target_function(x):
    return x**2 + 2*x + 1 # Example:  $f(x) = x^2 + 2x + 1$ 

# Input values
x_values = np.linspace(-10, 10, 20)

# Run the algorithm
gene_expression_algorithm(target_function, x_values, population_size=10, generations=10)

```

**Output:**

Best Solution:

Chromosome: [1, 3, '+', 2, 1, 4, '\*', '\*', '\*', 3], Fitness: 1259.2067

<ipython-input-3-6df17022c257>:25: RuntimeWarning: divide by zero encountered in scalar divide  
result = FUNCTIONS[gene](a, b)