

Bio Inspired Systems

Name: Sanghamitra R

USN: 1BM22CS237

CODE:

```
import numpy as np

import random

# Define any optimization function to minimize (can be changed as needed)

def custom_function(x):

    # Example function: x^2 to minimize

    return np.sum(x ** 2) # Ensuring the function works for
multidimensional inputs

# Initialize population of genetic sequences (each individual is a
sequence of genes)

def initialize_population(population_size, num_genes, lower_bound,
upper_bound):

    # Create a population of random genetic sequences

    population = np.random.uniform(lower_bound, upper_bound,
(population_size, num_genes))

    return population

# Evaluate the fitness of each individual (genetic sequence) in the
population

def evaluate_fitness(population, fitness_function):

    fitness = np.zeros(population.shape[0])
```

```

    for i in range(population.shape[0]):

        fitness[i] = fitness_function(population[i]) # Apply the fitness
function to each individual

    return fitness

# Perform selection: Choose individuals based on their fitness (roulette
wheel selection)

def selection(population, fitness, num_selected):

    # Select individuals based on their fitness (higher fitness, more
likely to be selected)

    probabilities = fitness / fitness.sum() # Normalize fitness to create
selection probabilities

    selected_indices = np.random.choice(range(len(population)),
size=num_selected, p=probabilities)

    selected_population = population[selected_indices]

    return selected_population

# Perform crossover: Combine pairs of individuals to create offspring

def crossover(selected_population, crossover_rate):

    new_population = []

    num_individuals = len(selected_population)

    for i in range(0, num_individuals - 1, 2): # Iterate in steps of 2,
skipping the last one if odd

```

```

    parent1, parent2 = selected_population[i], selected_population[i +
1]

    if len(parent1) > 1 and random.random() < crossover_rate: # Only
perform crossover if more than 1 gene

        crossover_point = random.randint(1, len(parent1) - 1) #
Choose a random crossover point

        offspring1 = np.concatenate((parent1[:crossover_point],
parent2[crossover_point:]))

        offspring2 = np.concatenate((parent2[:crossover_point],
parent1[crossover_point:]))

        new_population.extend([offspring1, offspring2]) # Create two
offspring

    else:

        new_population.extend([parent1, parent2]) # No crossover,
retain the parents

    # If the number of individuals is odd, carry the last individual
without crossover

    if num_individuals % 2 == 1:

        new_population.append(selected_population[-1])

    return np.array(new_population)

# Perform mutation: Introduce random changes in offspring

def mutation(population, mutation_rate, lower_bound, upper_bound):

```

```

    for i in range(population.shape[0]):

        if random.random() < mutation_rate: # Apply mutation based on the
rate

            gene_to_mutate = random.randint(0, population.shape[1] - 1) #
Choose a random gene to mutate

            population[i, gene_to_mutate] = np.random.uniform(lower_bound,
upper_bound) # Mutate the gene

    return population

# Gene expression: In this context, it is how we decode the genetic
sequence into a solution

def gene_expression(individual, fitness_function):

    return fitness_function(individual)

# Main function to run the Gene Expression Algorithm

def gene_expression_algorithm(population_size, num_genes, lower_bound,
upper_bound, max_generations, mutation_rate, crossover_rate,
fitness_function):

    # Step 2: Initialize the population of genetic sequences

    population = initialize_population(population_size, num_genes,
lower_bound, upper_bound)

    best_solution = None

    best_fitness = float('inf')

```

```
# Step 9: Iterate for the specified number of generations

for generation in range(max_generations):

    # Step 4: Evaluate fitness of the current population

    fitness = evaluate_fitness(population, fitness_function)

    # Track the best solution found so far

    min_fitness = fitness.min()

    if min_fitness < best_fitness:

        best_fitness = min_fitness

        best_solution = population[np.argmin(fitness)]

    # Step 5: Perform selection (choose individuals based on fitness)

    selected_population = selection(population, fitness,
population_size // 2)  # Select half of the population

    # Step 6: Perform crossover to generate new individuals

    offspring_population = crossover(selected_population,
crossover_rate)

    # Step 7: Perform mutation on the offspring population
```

```

        population = mutation(offspring_population, mutation_rate,
lower_bound, upper_bound)

        # Print output every 10 generations

        if (generation + 1) % 10 == 0:

            print(f"Generation {generation + 1}/{max_generations}, Best
Fitness: {best_fitness}")

        # Step 10: Output the best solution found

        return best_solution, best_fitness

# Parameters for the algorithm

population_size = 50 # Number of individuals in the population

num_genes = 1 # Number of genes (for a 1D problem, this is just 1,
extendable for higher dimensions)

lower_bound = -5 # Lower bound for the solution space

upper_bound = 5 # Upper bound for the solution space

max_generations = 100 # Number of generations to evolve the population

mutation_rate = 0.1 # Mutation rate (probability of mutation per gene)

crossover_rate = 0.7 # Crossover rate (probability of crossover between
two parents)

# Run the Gene Expression Algorithm

```

```

best_solution, best_fitness = gene_expression_algorithm(

    population_size, num_genes, lower_bound, upper_bound,

    max_generations, mutation_rate, crossover_rate, custom_function)

# Output the best solution found

print("\nBest Solution Found:", best_solution)

print("Best Fitness Value:", best_fitness)

```

OUTPUT:

```

⇌ Generation 10/100, Best Fitness: 1.9980464857154846e-05
    Generation 20/100, Best Fitness: 8.798327298656325e-06
    Generation 30/100, Best Fitness: 8.798327298656325e-06
    Generation 40/100, Best Fitness: 8.798327298656325e-06
    Generation 50/100, Best Fitness: 8.798327298656325e-06
    Generation 60/100, Best Fitness: 8.798327298656325e-06
    Generation 70/100, Best Fitness: 8.798327298656325e-06
    Generation 80/100, Best Fitness: 8.798327298656325e-06
    Generation 90/100, Best Fitness: 8.798327298656325e-06
    Generation 100/100, Best Fitness: 8.798327298656325e-06

    Best Solution Found: [0.0029662]
    Best Fitness Value: 8.798327298656325e-06

```

Application Problem

Question: Use GEP to model and predict environmental systems(pollution levels, climate change)by optimising parameters and equations based on historical data and simulation results.

Code:

```

import numpy as np
import random
import math

```

```

import pandas as pd
from sklearn.metrics import mean_squared_error

# Define the objective function based on the environment data (pollution,
temperature, etc.)
def environmental_objective(individual, data):
    """
    The objective function for environmental prediction.
    It evaluates how well the model (equation) represented by 'individual'
predicts actual data.
    """
    predictions = []
    for row in data:
        prediction = evaluate_equation(individual, row) # Evaluate the
individual (equation)
        predictions.append(prediction)

    # Calculate the fitness (mean squared error between predictions and
actual data)
    true_values = data[:, -1] # Assuming last column is the actual value
    mse = mean_squared_error(true_values, predictions)
    return mse # Lower MSE is better

def evaluate_equation(individual, row):
    """
    Evaluate the equation represented by the individual (a sequence of
genes).
    'individual' will be a mathematical expression encoded in genes.
    """
    result = 0
    for i, gene in enumerate(individual):
        result += gene * row[i] # A simple linear model: gene *
feature_value

    return result

# Initialize the population of genetic sequences (individuals)
def initialize_population(population_size, num_genes, lower_bound,
upper_bound):

```



```

    population = np.random.uniform(lower_bound, upper_bound,
(population_size, num_genes))
    return population

# Perform roulette wheel selection
def selection(population, fitness, num_selected):
    probabilities = fitness / fitness.sum()
    selected_indices = np.random.choice(range(len(population)),
size=num_selected, p=probabilities)
    selected_population = population[selected_indices]
    return selected_population

# Perform crossover to generate offspring
def crossover(selected_population, crossover_rate):
    new_population = []
    num_individuals = len(selected_population)

    for i in range(0, num_individuals - 1, 2):
        parent1, parent2 = selected_population[i], selected_population[i +
1]

        if random.random() < crossover_rate:
            crossover_point = random.randint(1, len(parent1) - 1)
            offspring1 = np.concatenate((parent1[:crossover_point],
parent2[crossover_point:]))
            offspring2 = np.concatenate((parent2[:crossover_point],
parent1[crossover_point:]))
            new_population.extend([offspring1, offspring2])
        else:
            new_population.extend([parent1, parent2])

    if num_individuals % 2 == 1:
        new_population.append(selected_population[-1])

    return np.array(new_population)

# Perform mutation: introduce random changes
def mutation(population, mutation_rate, lower_bound, upper_bound):
    for i in range(population.shape[0]):
        if random.random() < mutation_rate:
            gene_to_mutate = random.randint(0, population.shape[1] - 1)

```

```

        population[i, gene_to_mutate] = np.random.uniform(lower_bound,
upper_bound)
    return population

# Main function to run the Gene Expression Algorithm (GEP)
def gene_expression_algorithm(population_size, num_genes, lower_bound,
upper_bound, max_generations, mutation_rate, crossover_rate, data):
    population = initialize_population(population_size, num_genes,
lower_bound, upper_bound)

    best_solution = None
    best_fitness = float('inf')

    # Iterate over generations
    for generation in range(max_generations):
        fitness = np.array([environmental_objective(individual, data) for
individual in population])

        min_fitness = fitness.min()
        if min_fitness < best_fitness:
            best_fitness = min_fitness
            best_solution = population[np.argmin(fitness)]

        # Select individuals based on fitness
        selected_population = selection(population, fitness,
population_size // 2)

        # Perform crossover
        offspring_population = crossover(selected_population,
crossover_rate)

        # Perform mutation
        population = mutation(offspring_population, mutation_rate,
lower_bound, upper_bound)

        if (generation + 1) % 10 == 0:
            print(f"Generation {generation + 1}/{max_generations}, Best
Fitness (MSE): {best_fitness:.6f}")

    return best_solution, best_fitness

```

```

# Sample data: Replace this with actual environmental data (pollution
levels, temperature, etc.)
# Each row represents a data point with features and the last column is
the target value (e.g., pollution level)
data = np.array([
    [1.0, 2.0, 3.0, 4.0, 5.0, 15.0], # Example row (features followed by
actual value)
    [2.0, 3.0, 4.0, 5.0, 6.0, 20.0],
    [3.0, 4.0, 5.0, 6.0, 7.0, 25.0],
    [4.0, 5.0, 6.0, 7.0, 8.0, 30.0],
    [5.0, 6.0, 7.0, 8.0, 9.0, 35.0],
    [6.0, 7.0, 8.0, 9.0, 10.0, 40.0]
])

# Parameters for the GEP algorithm
population_size = 50
num_genes = 5 # Number of genes (equation components)
lower_bound = -10 # Lower bound for the solution space
upper_bound = 10 # Upper bound for the solution space
max_generations = 100 # Number of generations to evolve the population
mutation_rate = 0.1 # Mutation rate
crossover_rate = 0.7 # Crossover rate

# Run the Gene Expression Algorithm
best_solution, best_fitness = gene_expression_algorithm(
    population_size, num_genes, lower_bound, upper_bound, max_generations,
    mutation_rate, crossover_rate, data)

print("\nBest Solution Found:", best_solution)
print("Best Fitness (MSE):", best_fitness)

```

Output:

```
↔ Generation 10/100, Best Fitness (MSE): 11.422478  
Generation 20/100, Best Fitness (MSE): 11.422478  
Generation 30/100, Best Fitness (MSE): 11.422478  
Generation 40/100, Best Fitness (MSE): 11.422478  
Generation 50/100, Best Fitness (MSE): 11.422478  
Generation 60/100, Best Fitness (MSE): 11.422478  
Generation 70/100, Best Fitness (MSE): 11.422478  
Generation 80/100, Best Fitness (MSE): 11.422478  
Generation 90/100, Best Fitness (MSE): 11.422478  
Generation 100/100, Best Fitness (MSE): 11.422478
```

```
Best Solution Found: [ 5.51001406 -3.91428818  7.50820505 -9.77928146  5.96757112]  
Best Fitness (MSE): 11.42247784545849
```