simplest_code

February 4, 2024

```
[2]: import random
import math
import matplotlib.pyplot as plt
import numpy as np
```

1 Monte Carlo A

```
[3]: def f(x):
    return x**2

def monte_carlo_integration(num_samples, a, b):
    integral_sum = 0

    for _ in range(num_samples):
        x = random.uniform(a, b)
        integral_sum += f(x)

    estimated_integral = ((b - a) / num_samples) * integral_sum
    return estimated_integral

num_samples = 10000 # You can adjust the number of samples for more accuracy
a, b = 0, 1 # Integration limits
estimated_integral = monte_carlo_integration(num_samples, a, b)

print(f"Estimated Integral: {estimated_integral}")
```

Estimated Integral: 0.3315121609624472

2 Monte Carlo Importance Sampling

```
[6]: def f(x):
    return 1 / (x**2 + 1)

def sampling_distribution(x):
    return 2 / (2 + x)

def original_distribution():
```

```
x = random.uniform(a, b)
return x

def importance_sampling(num_samples, a, b):
    integral_sum = 0

for _ in range(num_samples):
    x = original_distribution()
    weight = sampling_distribution(x) / (b - a)
        integral_sum += f(x) / weight

    estimated_integral = integral_sum / num_samples
    return estimated_integral

num_samples = 10000 # You can adjust the number of samples for more accuracy
a, b = -1, 1 # Integration limits
estimated_integral = importance_sampling(num_samples, a, b)

print(f"Estimated Integral: {estimated_integral}")
```

Estimated Integral: 1.5633994889690292

3 Rejection Sampling

```
[8]: # Target distribution (normal distribution)
     def target_distribution(x):
         return (1 / (math.sqrt(2 * math.pi))) * math.exp(-0.5 * x**2)
     # Proposal distribution (uniform distribution)
     def proposal_distribution(a, b):
         return random.uniform(a, b)
     def rejection_sampling(a, b):
         while True:
             x = proposal_distribution(a, b)
             accept_prob = target_distribution(x) / (1 / (b - a))
             if random.uniform(0, 1) < accept_prob:</pre>
                 sample = x
                 break
         return sample
     a, b = -3, 3 # Range for the uniform proposal distribution
     sample = rejection_sampling(a, b)
```

4 Inverse Sampling

5 Metropolis-Hastings

```
[]: # Define the target distribution (a simple discrete distribution)
     def target_distribution(x):
         if x == 1:
             return 0.3
         elif x == 2:
             return 0.4
         else:
             return 0.3
     # Proposal distribution (simple random walk)
     def proposal_distribution(x):
         return x + random.choice([-1, 1])
     def metropolis_hastings(num_samples, initial_sample):
         samples = [initial_sample]
         for _ in range(num_samples - 1):
             current_sample = samples[-1]
             proposed_sample = proposal_distribution(current_sample)
             # Calculate acceptance ratio
             acceptance_ratio = min(1, target_distribution(proposed_sample) /__
      →target_distribution(current_sample))
             # Accept or reject the proposal
```

6 SSA

```
[]: # Initialize the system
     population = 10
     time = 0
     time_points = [time]
     population_counts = [population]
     # Simulation parameters
     end\_time = 100.0
     while time < end_time:</pre>
         # Define reaction constants
         birth_rate = 2.0
         death_rate = 0.02
         # Calculate the propensity functions
         a_birth = birth_rate
         a_death = death_rate * population
         rxn_as = np.array([a_birth, a_death])
         a_total = np.sum(rxn_as)
         if a_total != 0: # can't divide by 0
             # Calculate time step (tau) from exponential distribution
             tau = -np.log(np.random.uniform(0,1)) / a_total
             # Determine the next reaction to occur
             possible_rxns = np.arange(len(rxn_as))
             i = np.random.choice(possible_rxns, p=rxn_as/a_total)
             # Apply rxn
             if i == 0:
                 population += 1
             elif i == 1:
```

```
population -= 1

# Update time and counts
time += tau
population_counts.append(population)
time_points.append(time)
else: # a0 == 0, no more rxns
time = end_time
population_counts.append(population)
time_points.append(time)

# Plot the results
plt.plot(time_points, population_counts)
```

7 Tau leaping

```
[]: # Initialize the system
     population = 10
     time = 0
     time_points = [time]
     population_counts = [population]
     # Simulation parameters
     end time = 10.0
     tau = 0.1
     while time < end_time:</pre>
         # Define reaction constants
         birth_rate = 2.0
         death_rate = 0.02
         # Calculate the propensity functions
         a_birth = birth_rate
         a_death = death_rate * population
         # Sample the number of reactions for each channel from Poisson distribution
         num_births = 0
         num_deaths = 0
         if (a birth*tau > 0):
             num_births = np.random.poisson(a_birth * tau)
         if (a death*tau > 0):
             num_deaths = np.random.poisson(a_death * tau)
         # Update time and population count
         time += tau
         population += num_births - num_deaths
```

```
population_counts.append(population)
   time_points.append(time)

# Plot the results
plt.step(time_points, population_counts)
```

8 PSO

```
[576]: def f(x):
          return x**2
      # Define the PSO parameters
      num_particles = 10
      max_iterations = 10
      c1 = 1.5 # Cognitive parameter
      c2 = 1.5 # Social parameter
      w = 0.7 # Inertia weight
      # Initialize the particles
      particles = []
      for _ in range(num_particles):
          position = random.uniform(-10, 10) # Random initial position
          velocity = random.uniform(-1, 1) # Random initial velocity
          personal_best_position = position
          personal_best_value = f(position)
          particles.append([position, velocity, personal_best_position,_
       →personal_best_value])
      # Initialize the global best-known position and value
      global_best_position = particles[0][0]
      global_best_value = particles[0][3]
      # PSO optimization loop
      for iteration in range(max_iterations):
          for i in range(num_particles):
              particle = particles[i]
              position, velocity, personal_best_position, personal_best_value = __
       →particle
              # 1. Update velocity and position
              r1, r2 = random.random(), random.random()
              velocity = w * velocity + c1 * r1 * (personal_best_position - position)
       position = position + velocity
              # 2. Evaluate the objective function at the new position
```

```
value = f(position)
         # 3. Update personal best if needed
        if value < personal_best_value:</pre>
             personal_best_position = position
             personal_best_value = value
         # 4. Update global best if needed
        if value < global best value:</pre>
             global_best_position = position
             global_best_value = value
         # 5. Update the particle's information
        particles[i] = [position, velocity, personal_best_position,__
  →personal_best_value]
    print(f"Iteration {iteration + 1}: Global Best Value = {global_best_value:.

4f}")
print(f"Optimal Solution: x = {global_best_position}, f(x) = {global_best_value:
  ⇔.4f}")
Iteration 1: Global Best Value = 0.1242
Iteration 2: Global Best Value = 0.1242
```

```
Iteration 1: Global Best Value = 0.1242
Iteration 2: Global Best Value = 0.0003
Iteration 3: Global Best Value = 0.0003
Iteration 4: Global Best Value = 0.0003
Iteration 5: Global Best Value = 0.0003
Iteration 6: Global Best Value = 0.0003
Iteration 7: Global Best Value = 0.0003
Iteration 8: Global Best Value = 0.0003
Iteration 9: Global Best Value = 0.0003
Iteration 10: Global Best Value = 0.0003
Optimal Solution: x = -0.01583582034806863, f(x) = 0.0003
```

9 ACO

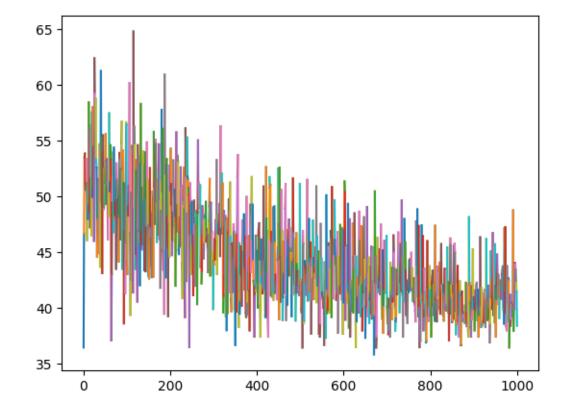
```
[7.211102550927978, 4.123105625617661, 3.1622776601683795, 0, 3.
 →605551275463989, 5.0, 6.324555320336759, 5.830951894845301, 7.
 \hookrightarrow0710678118654755, 5.656854249492381],
    [10.816653826391969, 7.615773105863909, 6.4031242374328485, 3.
 5.0, 6.082762530298219,
    [8.54400374531753, 5.0990195135927845, 7.280109889280518, 5.0, 6.
 4324555320336759, 0, 5.385164807134504, 8.06225774829855, 11.180339887498949, L
 →10.63014581273465],
    [12.806248474865697, 9.219544457292887, 9.486832980505138, 6.
 -324555320336759, 4.123105625617661, 5.385164807134504, 0, 4.242640687119285, u
 →8.602325267042627, 10.198039027185569],
    [13.038404810405298, 9.848857801796104, 8.48528137423857, 5.
 →830951894845301, 2.23606797749979, 8.06225774829855, 4.242640687119285, 0, 4.
 →47213595499958, 7.0710678118654755],
    [13.341664064126334, 11.0, 8.246211251235321, 7.0710678118654755, 5.0, 11.
 ↔180339887498949, 8.602325267042627, 4.47213595499958, 0, 4.242640687119285],
    [10.0, 8.54400374531753, 5.0990195135927845, 5.656854249492381, 6.
 4082762530298219, 10.63014581273465, 10.198039027185569, 7.0710678118654755, u
 →4.242640687119285, 0]
]
# ACO parameters
num_ants = 3
num_iterations = 1000
initial_pheromone = 0.001
evaporation_rate = 0.01
pheromone_deposit_rate = 1.0
alpha = 3.0 # Pheromone Importance
beta = 1.0 # Heuristic Importance
exploitation_prob_q = 0.05 # Exploitation vs Exploration probability
# Initialize pheromone levels
num cities = len(distance matrix)
pheromone_matrix = [[initial_pheromone+1] * num_cities for _ in_
 →range(num cities)]
# ACO main loop
for iteration in range(num_iterations):
   ant_tours = []
   # 1. Loop through ants
   for ant in range(num_ants):
       # 2. randomly initialize ant at random city
       start_city = random.randint(0, num_cities - 1)
       tour = [start_city]
```

```
# 3. Loop through remaining cities
      while len(tour) < num_cities:</pre>
          remaining cities = [i for i in range(num_cities) if i not in tour]
          probabilities = []
          # 4. Decide Exploitation vs Exploration
          if random.uniform(0, 1) > exploitation_prob_q:
              # Explore based on pheromones
              for city in remaining_cities:
                  pheromone = pheromone matrix[start city][city]
                  distance = distance_matrix[start_city][city]
                  attractiveness = (pheromone ** alpha) * ((1 / distance) **__
⇔beta)
                  probabilities.append((city, attractiveness))
              total_attractiveness = sum(attractiveness for _, attractiveness_
→in probabilities)
              probabilities = [(city, attractiveness / total_attractiveness)⊔

→for city, attractiveness in probabilities]
              next_city = np.random.choice([city for city, _ in_
probabilities], p=[prob for _, prob in probabilities])
          else:
              # Exploit based on pure heuristic
              min_distance = max(max(distance_matrix)) # initialize_
⇔min_distance
              for city in remaining_cities:
                  if min_distance >= distance_matrix[start_city][city]:
                      min_distance = distance_matrix[start_city][city]
                      next_city = city
          # 5. Local update pheromone levels to incentivize new trails
          pheromone_matrix[start_city][next_city] = (1 -__
→evaporation_rate)*pheromone_matrix[start_city][next_city] +__
pheromone_matrix[next_city] [start_city] = (1 -__
→evaporation_rate)*pheromone_matrix[start_city][next_city] +

→(evaporation_rate * initial_pheromone)
          tour.append(next_city)
          start_city = next_city
      ant_tours.append(tour)
  # for plotting
  prev_best_distance = best_distance
  # 6. Find the best tour in this iteration
```

Best Tour Found: [3, 1, 0, 2, 9, 8, 7, 4, 6, 5] Total Distance: 38.385810980638034



10 DE

```
[3]: def f(x):
         return sum(x**2)
     # DE Parameters
     population_size=5 # Size of the population
     min_bound=-10  # Lower bound for each parameter
     max_bound=10 # Upper bound for each parameter
     num_generations = 10
     mutation=0.8 # Mutation Factor
     crossover_probability=0.7 # Crossover_probability
     dimensions = 2 # Number of dimensions of an individual
     # 1. Initialize population with random solutions and their fitnesses
     population = min_bound + np.random.rand(population_size,_

→dimensions)*(max bound-min bound)
     fitnesses = np.array([f(individual) for individual in population])
     # 2. Loop through generations
     for generation in range(num_generations):
         # 3. Loop through population
         for individual in range(population_size):
             # 4. Choose 3 different individuals out of the population
             # Select indices of all individuals except the current individual
             idxs = [idx for idx in range(population_size) if idx != individual]
             a, b, c = population[np.random.choice(idxs, 3, replace = False)]
             # 5. Generate a mutant donor using the mutation factor and ensure still_{11}
      ⇒in bounds
             mutant_donor = np.clip(a + mutation * (b - c), min_bound, max_bound)
             # 6. Crossover Operation
             # randomly select dimension points for crossover based on crossover
      \hookrightarrowprobability
             points_to_crossover = np.random.rand(dimensions) < crossover_probability</pre>
             # Ensure at least one dimension/parameter is modified
             if not np.any(points_to_crossover):
                 points_to_crossover[np.random.randint(0, dimensions)] = True
             # Do Crossover
             trial_individual = np.where(points_to_crossover, mutant_donor,_
      →population[individual])
             trial_individual_fitness = f(trial_individual)
             # 7. Individual is replaced is trial_individual has better fitness
             if trial_individual_fitness < fitnesses[individual]:</pre>
```

Best Solution: [4.41175621 -0.52758748] Best Fitness: 19.74194138421374 Best Solution: [4.41175621 -0.52758748] Best Fitness: 19.74194138421374 Best Solution: [-0.49708186 -0.0597067] Best Fitness: 0.25065526941514615 Best Solution: [-0.49708186 -0.0597067] Best Fitness: 0.25065526941514615 Best Solution: [-0.49708186 -0.0597067] Best Fitness: 0.25065526941514615 Best Solution: [0.11870417 0.48221743] Best Fitness: 0.24662432951451072 Best Solution: [-0.14440907 0.01058759] Best Fitness: 0.020966077069033456 Best Solution: [-0.00122927 0.10352288] Best Fitness: 0.010718498212355987 Best Solution: [-0.00122927 0.10352288] Best Fitness: 0.010718498212355987 Best Solution: [-0.05115475 -0.08012017] Best Fitness: 0.009036049255503473

11 PBIL

```
[654]: # Define the MaxOnes fitness function
  def max_ones_fitness(binary_vector):
        return sum(binary_vector)

# PBIL parameters
vector_length = 10  # Length of the binary vector
population_size = 5
  alpha = 0.2  # Alpha rate for updating probabilities (basically learning rate)
  mutation_prob = 0.02  # Probability of mutating a bit
  beta = 1.0  # Mutation strength
  num_generations = 10

# Initialize the probability vector with equal probabilities
probability_vector = np.array([0.5] * vector_length)
```

```
# Main PBIL loop
for generation in range(num_generations):
    # 1. Generate a population based on the probability vector
    population = [[int(random.random() < p) for p in probability_vector] for _u
 →in range(population_size)]
    # 2. Evaluate the fitness of each individual
    fitness_values = [max_ones_fitness(individual) for individual in population]
    # 3. Find the best individual in the population
    best_individual = np.array(population[fitness_values.
  ⇔index(max(fitness_values))])
    # 4. Update the probability vector with alpha
    probability_vector = (1-alpha)*probability_vector + (alpha*best_individual)
    # Mutate each element in probability vector based on mutation prob
    for i in range(vector_length):
        if random.random() < mutation_prob:</pre>
            probability_vector[i] = (1-beta)*probability_vector[i] +__
  \hookrightarrow (beta*random.uniform(0,1))
    # Find the best individual in the final population
    best individual = max(population, key=max ones fitness)
    best_fitness = max_ones_fitness(best_individual)
    print("Best Individual:", best_individual, "Fitness:", best_fitness)
Best Individual: [1, 1, 1, 0, 1, 0, 1, 0, 1, 0] Fitness: 6
Best Individual: [1, 1, 0, 0, 1, 1, 1, 1, 1, 0] Fitness: 7
Best Individual: [1, 1, 1, 0, 1, 0, 1, 1, 1, 1] Fitness: 8
Best Individual: [1, 0, 1, 0, 1, 1, 1, 1, 1, 1] Fitness: 8
Best Individual: [1, 1, 1, 0, 1, 1, 1, 0, 1, 1] Fitness: 8
Best Individual: [1, 1, 1, 1, 1, 0, 1, 1, 1, 1] Fitness: 9
Best Individual: [0, 1, 1, 1, 1, 0, 1, 1, 0, 1] Fitness: 7
Best Individual: [1, 1, 1, 1, 1, 1, 1, 1, 1] Fitness: 10
Best Individual: [1, 1, 1, 1, 1, 1, 1, 1, 1] Fitness: 10
Best Individual: [1, 1, 1, 1, 1, 1, 1, 1, 1] Fitness: 10
    PBILc
12
```

```
[619]: def f(x):
           return -sum(x**2) # Negative since PBIL and PBILc finds the max
       # PBIL parameters
       vector_length = 2  # Length of the parameter vector
```

```
population_size = 10
alpha = 0.1 # Alpha rate for updating the parameter vector (learning rate)
num_generations = 10
# Initialize the parameter vector with random values
parameter_vector = np.array([0.5] * vector_length)
# Standard deviation for normal distribution sampling
standard_deviations = np.array([0.5] * vector_length)
# Main PBIL loop
for generation in range(num_generations):
    # 1. Generate a population based on the parameter vector
    population = [parameter_vector + np.random.normal(0, standard_deviations,_
 ovector_length) for _ in range(population_size)]
    # 2. Evaluate the fitness of each individual
    fitness_values = [f(individual) for individual in population]
    # 3. Find the indices of the two best individuals and the worst individual
    best_indices = np.argsort(fitness_values)[-2:] # Two best individuals
    worst index = np.argmin(fitness values) # Worst individual
    # 4. Update the parameter vector using the two best and subtracting the
  \hookrightarrow worst
    parameter_vector +=_
  alpha*(population[best_indices[0]]+population[best_indices[1]]-population[worst_index])
    # 5. Update standard deviation
    # Find the best individual in the final population
    best_fitness = f(parameter_vector)
    print("Best Parameter Vector:", parameter_vector, "Best Fitness:", __
  ⇔best_fitness)
Best Parameter Vector: [0.46888632 0.54627467] Best Fitness: -0.5182703869594648
Best Parameter Vector: [0.39924071 0.45875045] Best Fitness: -0.3698451271509875
Best Parameter Vector: [0.42085799 0.35316008] Best Fitness: -0.3018434925009673
Best Parameter Vector: [0.43818061 0.237306 ] Best Fitness: -0.2483163874620917
Best Parameter Vector: [0.40625321 0.32200055] Best Fitness: -0.2687260309046179
Best Parameter Vector: [0.33628493 0.28180303] Best Fitness: -0.1925005040925098
Best Parameter Vector: [0.27561629 0.2262637 ] Best Fitness:
-0.12715959984775912
Best Parameter Vector: [0.02939499 0.30611676] Best Fitness:
-0.09457153844125527
Best Parameter Vector: [0.05723213 0.13434918] Best Fitness:
```

```
-0.021325218863756108

Best Parameter Vector: [0.16193774 0.02373301] Best Fitness: -0.026787086895781784
```

13 ES(1+1) with 1/5 rule

```
[]: def f(x):
        return x**2
     # ES parameters
     sigma = 0.1 # Initial mutation step size
     max_generations = 100
     # Success rule parameters
     success counter = 0
     success_rate = 1/5 # The 1/5 rule: Acceptance rate threshold
     # Initial guess for the solution
     x = 5.0 # You can choose any initial value
     # Main (1+1) ES loop
     generations_counter = 0
     for generation in range(max_generations):
         # 1. Create 1 candidate solution by applying a random mutation
         candidate = x + random.gauss(0, sigma)
         # 2. Evaluate the fitness of the candidate solution
         current fitness = f(x)
         candidate_fitness = f(candidate)
         # 3. Decide whether to accept the candidate based on fitness
         if candidate_fitness < current_fitness:</pre>
             x = candidate # Accept the candidate solution
             success_counter += 1
         generations_counter += 1
         # 4. Check the success rate against the 1/5 rule
         if success_counter / generations_counter >= success_rate:
             # Increase mutation step size if the success rate is met
             sigma *= 1.5
             success_counter = 0 # Reset success counter
             generations_counter = 0 # Reset generations counter
         else:
             # If not that many of the new surrounding points are better, decrease,
             sigma *= 0.9
```

```
# Print the best solution in this generation

print(f"Best Solution = {x:.4f}, Best Fitness = {current_fitness:.4f}, 

⇒Sigma = {sigma:.4f}")
```

14 Cov matrix

```
[3]: x = [92, 60, 100]
y = [80, 30, 70]

normed_X = np.array([x - np.mean(x), y - np.mean(y)])

C = np.matmul(normed_X,normed_X.T) / (len(x)-1)
print(C)

[[448. 520.]
[520. 700.]]
```