

Lab 5 EI Farol Bar Problem

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Exercise 4

Algorithm 1: Coevolution

Input: pop size, h, weeks, max_t

Output: result

Pop = initial (pop size)

Fit, flag, attendance = fitness

For iteration in range(max_t):

 For w in range(weeks):

 For I in range(len(pop)):

 Next_ds = strategy

 d_s.append(next_ds)

 attendance_situation = d

 state = s

 new_fit, flag, attendance = fitness(attendance_situation)

 Record result

 m = 0, s = 0, count = 0

 For I in range(len(fit)):

 If fit[i] > fit[s]: m = i

 Else if fit[i] > fit[s]: s = i

 Count = count + fit[i]

 Parent1, parent2 = pop[m], pop[s]

 P1, p2, a1,a2,b1,b2 = getm

 P1, a1, b1 = mutation

 P2, a2, b2 = mutation

 Pop = crossover

Return result

Algorithm 2: initial

Input: pop size, h

Output: pop

For I in range (pop size):

 Add h to individual

 For I in range(h):

 Add random number in individual

 For I in range(h):

 a = random dirichlet

 Extend individual with a

 For I in range(h):

 b = random dirichlet

 Extend individual with b

 Pop.append(individuals)

Return pop

Algorithm 3: fitness

Input: result
Output: fit, flag, attendance
Fit = []
L = len(result)
Attendance = result.count(1)
If l == 0: crowded = 0
Else: crowded = attendance/l
If crowded < 0.6:
 For I in range(l):
 If result[i] == 1: fit.append(1)
 Else: fit.append(0)
 Flag = 0
else
 For I in range(l):
 If result[i] == 0: fit.append(1)
 Else: fit.append(0)
 Flag = 1
Return fit, flag, attendance

Algorithm 4: mutation

Input: h, p, a, b
Output: p, a, b
Prob = random number
If prob > mutation prob:
 For I in range(h):
 p.append(random)
index_a = int(random* h)
Prob_a = random number
If prob_a > mutation prob:
 a = random dirichlet
index_b = int(random* h)
Prob_b = random number
If prob_b > mutation prob:
 b = random dirichlet
Return: p, a, b

Algorithm 5: crossover

Input: pop size, h, p1, p2, a1, a2, b1, b2
Output: new pop
For I in range (pop size):
 For j in range(len(p1)):
 Prob = random number
 If prob > crossover prob: p1[j], p2[j] = p2[j], p1[j]

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For j in range(0, len(a1), h):
    Prob = random number
    If prob > crossover prob: a1[j:j + h], a2[j:j + h] = a2[j:j + h], a1[j:j + h]
For j in range (0, len(b1), h):
    Prob = random number
    If prob > crossover prob: b1[j:j + h], b2[j:j + h] = b2[j:j + h], b1[j:j + h]
new1.append(h); new2.append(h)
For I in range(h):
    new1.append(p1[i])
    new2.append(p2[i])
    new1.extend(copy (a1[i * h:(i * h) + h]))
    new2.extend(copy (a2[i * h:(i * h) + h]))
    new1.extend(copy (b1[i * h:(i * h) + h]))
    new2.extend(copy (b2[i * h:(i * h) + h]))
Return new pop
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Exercise 5

The aim of co-evolutionary algorithm which leads the population to as efficient utilisation of the bar as possible (i.e., close to 60 %), while still not being overcrowded. So can use the number of attendances to judge how good the algorithm works. I chose three parameters to turning.

1. Population Size

The initial setting of the crossover probability is 0.45, the mutation probability is 0.45, the number of states in the strategies is 5, the number of weeks to simulate per generation is 100, the number of generations to run the simulation is 100, the set of population size is 10, 50, 100, 500, 1000. The result is shown as box plot in figure 1, it can be seen that when the population size is 1000, the performance of the algorithm is best.

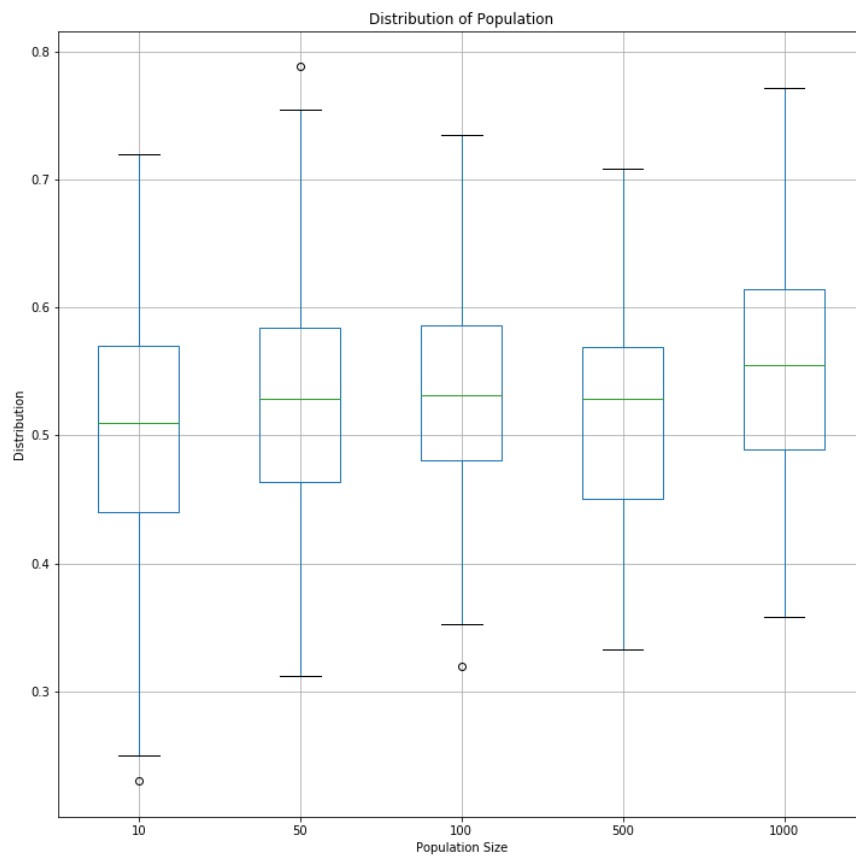


Figure 1. The distribution of population size

2. Crossover Probability

Choose the population size 1000 which perform, keep the other set, change the set of crossover probability. The set of crossover probability is 0.25, 0.45, 0.65, 0.85 and 0.99. The result is shown as box plot in figure 2, it can be seen that when the crossover probability is 0.45, the performance of the algorithm is best. But the change in crossover has little effect on the results.

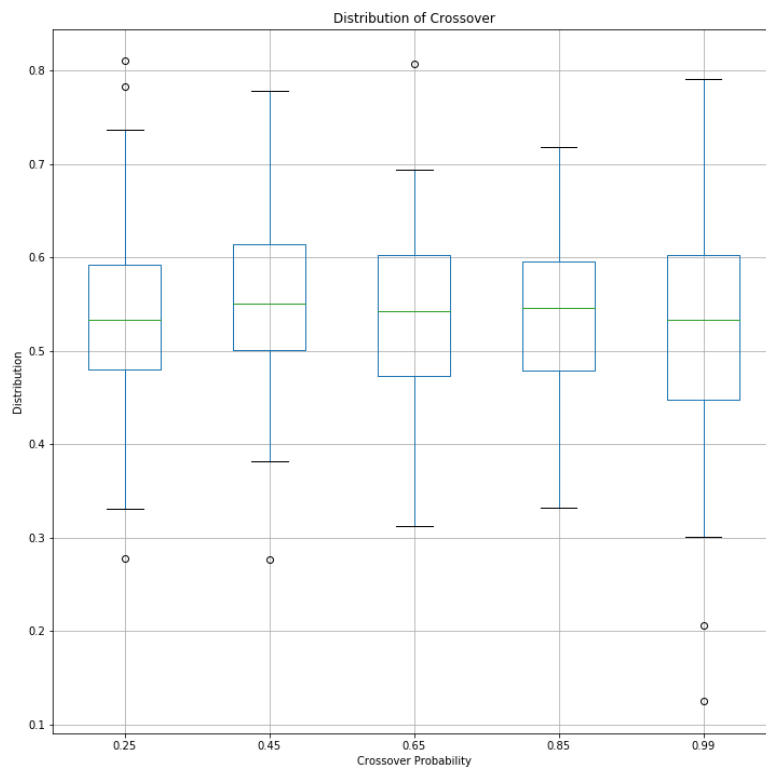


Figure 2. The distribution of Crossover probability

3. Mutation probability

After the first and second experiments, when the set of population size is 100 and the set of crossover probability is 0.45, the performance of the algorithm is better than other settings. Using these two setting and keep the other setting, change the set of mutation probability. The set of mutation probability is 0.25, 0.45, 0.65, 0.85 and 0.99. The result is shown as box plot in figure 3.

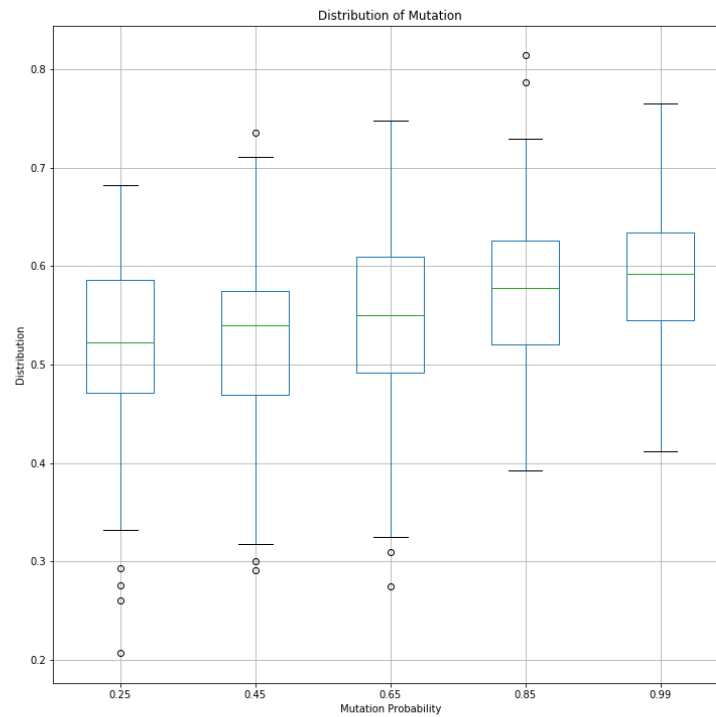


Figure 3. The distribution of Mutation probability