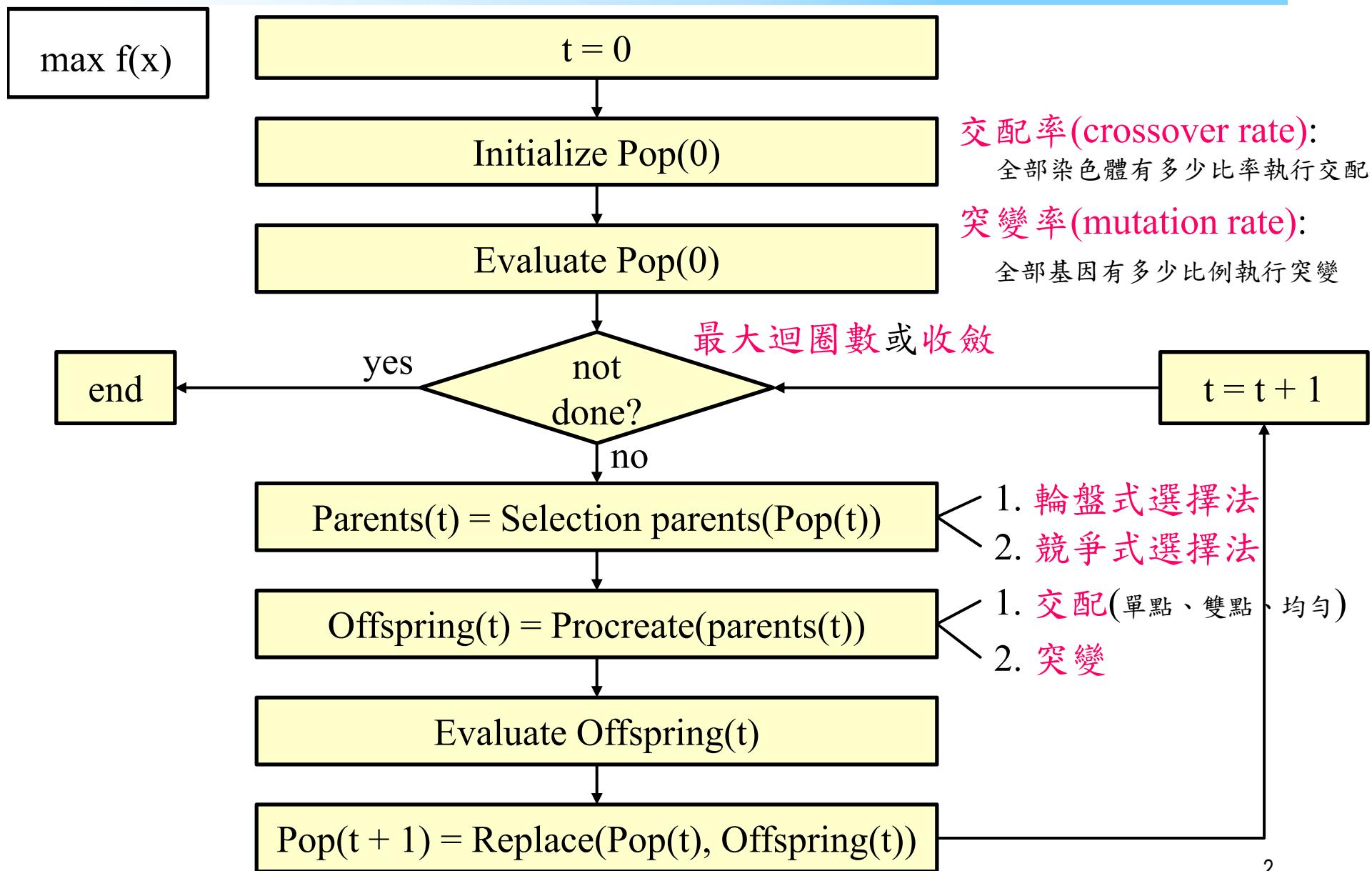
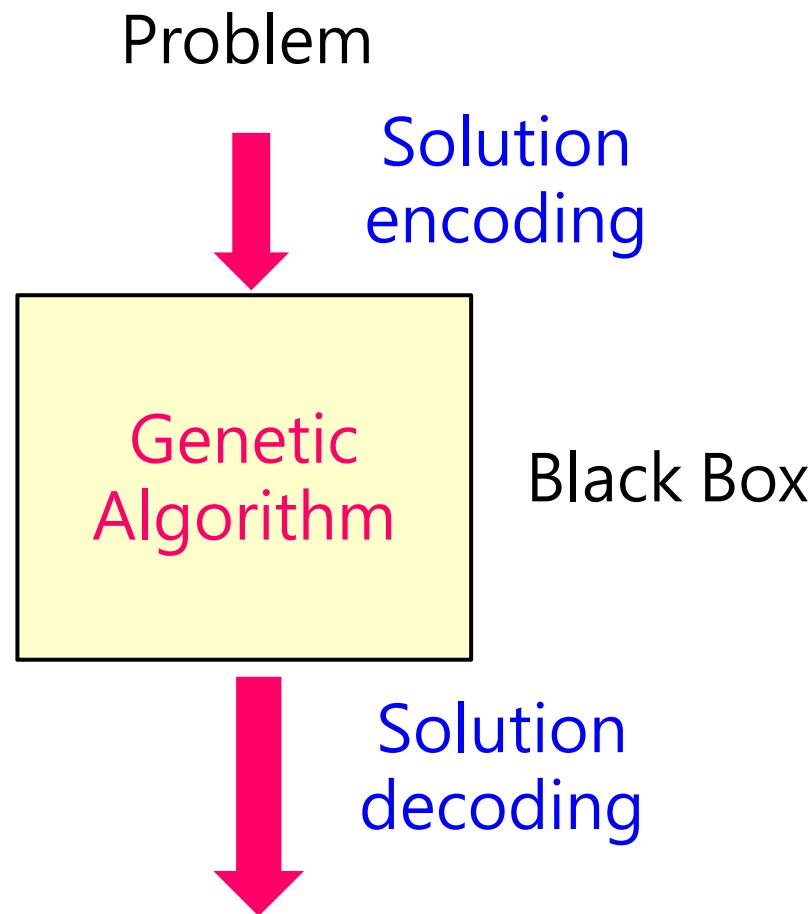


GA使用二元離散解的編碼

基因演算法流程圖



Framework of using the GA



- Optimal solutions
or approximation solutions
- Could solve large-scale problems

Solution encoding/decoding

- Encoding **continuous** decision variables

3.6	7.2	4.9	1.3	2.9
-----	-----	-----	-----	-----

- Encoding **discrete** decision variables

1	0	0	1	1
---	---	---	---	---

3	7	4	3	2
---	---	---	---	---

- Encoding **permutation** solutions

3	1	5	4	2
---	---	---	---	---

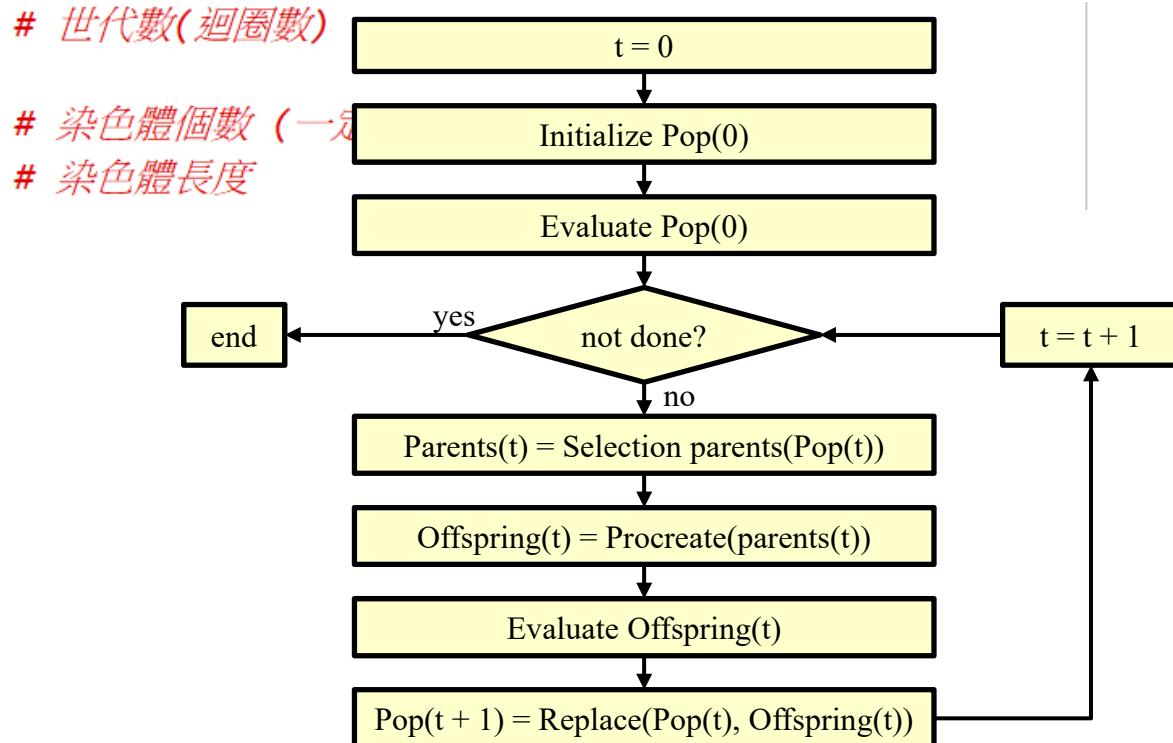
- Mixed encoding

3	1	5	4	2	1	0	0	1	1
---	---	---	---	---	---	---	---	---	---

```

9 NUM_ITERATION = 20
10
11 NUM_CHROME = 20
12 NUM_BIT = 6

```



```

84 # ===== 主程式 =====
85 pop = initPop()          # 初始化 pop
86 pop_fit = evaluatePop(pop) # 算 pop 的 fit
87
88 for i in range(NUM_ITERATION) :
89     parent = selection(pop, pop_fit)      # 挑父母
90     offspring = crossover(parent)          # 交配
91     mutation(offspring)                 # 突變
92     offspring_fit = evaluatePop(offspring) # 算子代的 fit
93     pop, pop_fit = replace(pop, pop_fit, offspring, offspring_fit) # 取代
94
95     print('iteration %d: x = %s, y = %d'    %(i, pop[0], pop_fit[0]))

```

```

9 NUM_ITERATION = 20          # 世代數(迴圈數)
10
11 NUM_CHROME = 20            # 染色體個數(一定要偶數)
12 NUM_BIT = 6                # 染色體長度
13
14 def selection(pop, pop_fit):
15     # 選擇父代
16
17     # 算子代的適應度
18     offspring_fit = evaluatePop(offspring)
19
20     # 取代
21     pop, pop_fit = replace(pop, pop_fit, offspring, offspring_fit)
22
23     print('iteration %d: x = %s, y = %d'    %(i, pop[0], pop_fit[0]))
24
25 def initPop():               # 初始化群體
26     return np.random.randint(2, size=(NUM_CHROME,NUM_BIT)) # 產生 NUM_CHROME 個二元編碼
27
28 def fitFunc(x):             # 適應度函數
29     # 將[1, 2, ..., NUM_BIT-1]的二元數轉成整數(第0數是符號數)
30     fitness = int("".join(str(i) for i in x[1:NUM_BIT]), 2)
31     return 1024 - fitness * fitness
32
33 def evaluatePop(p):         # 評估群體之適應度
34     return [fitFunc(p[i]) for i in range(len(p))]

```

1. 編碼：6個二元編碼，000000~111111
初始化：隨機設為6個二元編碼

2. 適應度：轉成10進位 x_d ，
然後計算 $fit = 1024 - x_d^2$

```

14 Pc = 0.5          # 交配率 (代表共執行Pc*NUM_CHROME/2次交配)
15 Pm = 0.01         # 突變率 (代表共要執行Pm*NUM_CHROME*NUM_BIT次突變)
16
17 NUM_PARENT = NUM_CHROME           # 父母的個數
18 NUM_CROSSOVER = int(Pc * NUM_CHROME / 2)    # 交配的次數
19 NUM_CROSSOVER_2 = NUM_CROSSOVER*2        # 上數的兩倍
20 NUM_MUTATION = int(Pm * NUM_CHROME * NUM_BIT)   # 突變的次數

```

$$0.5 * 20 / 2 \rightarrow 5$$

$$0.01 * 20 * 6 \rightarrow 1$$

```

36 def selection(p, p_fit):      # 用二元競爭式選擇法來挑父母
37     a = []
38     for i in range(NUM_PARENT):
39         [j, k] = np.random.choice(NUM_CHROME, 2, replace=False)  # 任選兩個index
40         if p_fit[j] > p_fit[k] :                                # 擇優
41             a.append(p[j])
42         else:
43             a.append(p[k])
44
45     return a

```

3. 選擇：二元競爭式選擇法

```

47 def crossover(p):            # 用單點交配來繁衍子代
48     a = []
49     for i in range(NUM_CROSSOVER) :
50         c = np.random.randint(1, NUM_BIT)                  # 隨機找出單點(不包含0)
51         [j, k] = np.random.choice(NUM_PARENT, 2, replace=False)  # 任選兩個index
52
53         a.append(np.concatenate((p[j][0: c], p[k][c: NUM_BIT]), axis=0))
54         a.append(np.concatenate((p[k][0: c], p[j][c: NUM_BIT]), axis=0))
55
56     return a

```

4. 交配：單點交配

```

14 Pc = 0.5          # 交配率 (代表共執行Pc*NUM_CHROME/2次交配)
15 Pm = 0.01         # 突變率 (代表共要執行Pm*NUM_CHROME*NUM_BIT次突變)
16
17 NUM_PARENT = NUM_CHROME           # 父母的個數
18 NUM_CROSSOVER = int(Pc * NUM_CHROME / 2)    # 交配的次數
19 NUM_CROSSOVER_2 = NUM_CROSSOVER*2        # 上數的兩倍
20 NUM_MUTATION = int(Pm * NUM_CHROME * NUM_BIT)  # 突變的次數

```

```

60 def mutation(p):      # 突變
61     for _ in range(NUM_MUTATION):
62         row = np.random.randint(NUM_CROSSOVER_2)  # 任選一個染色體
63         col = np.random.randint(NUM_BIT)           # 任選一個基因
64
65         p[row][col] = (p[row][col] + 1) % 2       # 對應此染色體的此基因01互換

```

```

68 def sortChrome(a, a_fit):      # a的根據a_fit由大排到小
69     a_index = range(len(a))      # 產生 0, 1, 2, ..., |a|-1 的 list
70
71     # a_index 根據 a_fit 的大小由大到小運動的排序
72     a_fit, a_index = zip(*sorted(zip(a_fit,a_index), reverse=True))
73
74     # 根據 a_index 的次序來回傳 a，並把對應的 fit 回傳
75     return [a[i] for i in a_index], a_fit

```

```

73 def replace(p, p_fit, a, a_fit): 6. 取代 : Pop(t+1) = {Pop(t) - {worsts}} ∪ {kids}
74     b = np.concatenate((p,a), axis=0)      # 把本代 p 和子代 a 合併成 b
75     b_fit = p_fit + a_fit                 # 把上述兩代的 fitness 合併
76     b, b_fit = sortChrome(b, b_fit)        # b 和 b_fit 運動的排序
77
78     return b[:NUM_CHROME], list(b_fit[:NUM_CHROME]) # 回傳 NUM_CHROME 個為新的一

```

5. 突變：任選一染色體的一個基因 · 01互換

a的根據a_fit由大排到小

產生 0, 1, 2, ..., |a|-1 的 list

6. 取代 : $\text{Pop}(t+1) = \{\text{Pop}(t) - \{\text{worsts}\}\} \cup \{\text{kids}\}$

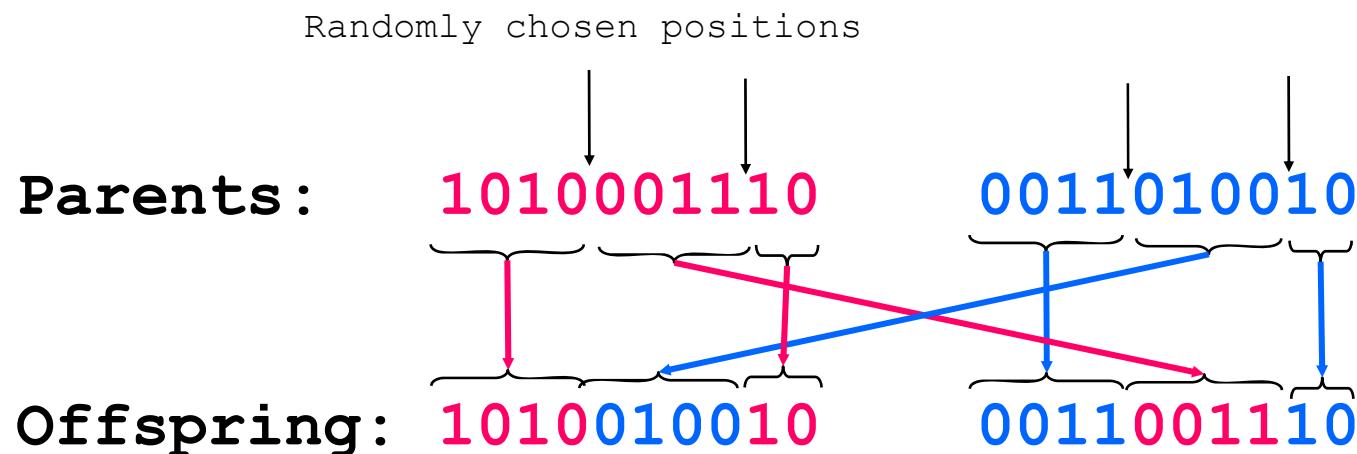
把本代 p 和子代 a 合併成 b

把上述兩代的 fitness 合併

b 和 b_fit 運動的排序

Exercise

- Implement the two-point crossover operation in the sample code “GA.py”.



- (Optional)
Implement the uniform crossover operation.