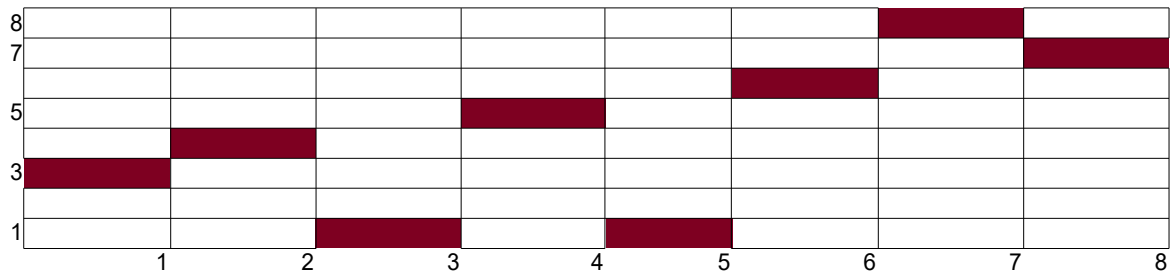


Introductory steps:

- what problem (task formulation): **8 Queens Problem** (place 8 queens on a (8x8) chessboard in a non-attacking configuration)
 - search space: the set of all the chessboard configurations???
 - genome: (slightly) complex
- => we can limit the search space to configurations containing only one queen in one column
- genome: 8-digit sequences, as:

3 4 1 5 1 6 8 7 (genome)
- - - - - - - -



(individual)

- fitness function: **28 - attacks**

(fitness f. Should be non-negative for the roulette wheel method)

We decided to take **N=6** individuals to the population (too few for the computer implementation)

GENERATION OF THE INITIAL POPULATION

We use www.random.org to generate 6 8-digit genomes

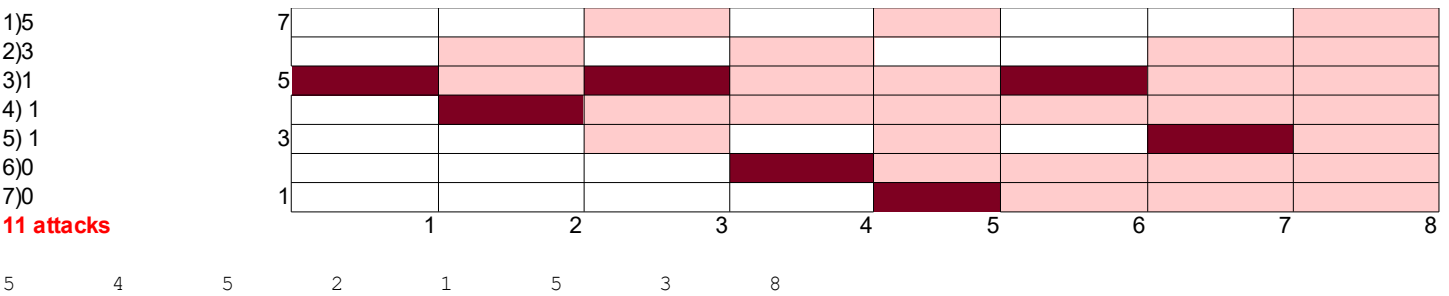
Genome of individual 1 (Ind_1):

5	4	5	2	1	5	3	8
Genome of Ind_2:							
5	6	7	7	2	6	3	6
Genome of Ind_3:							
1	3	1	2	6	3	4	3
Genome of Ind_4:							
3	2	4	5	5	2	3	7
Genome of Ind_5:							
8	5	8	7	4	5	7	4
Genome of Ind_6:							
8	6	7	8	3	5	2	6

SELECTION

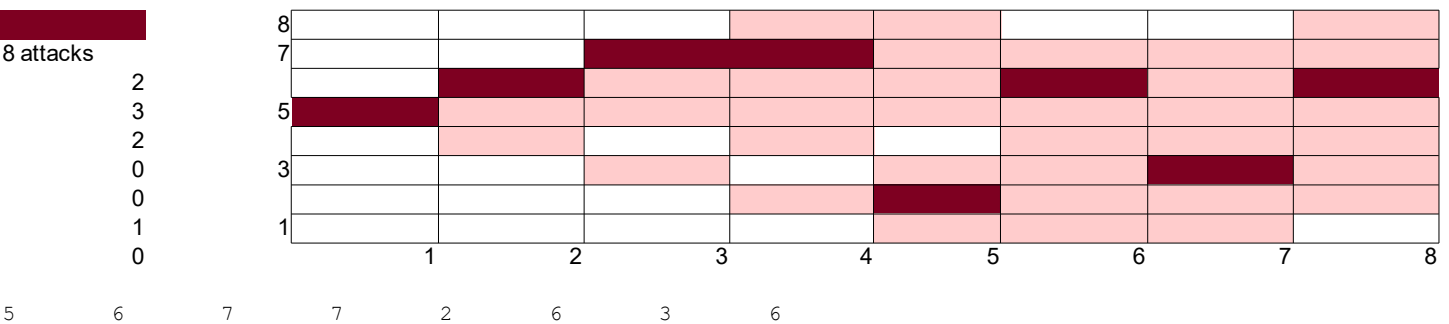
a) Evaluation (if any individual has the maximal fitness, we quit)

Ind_1:



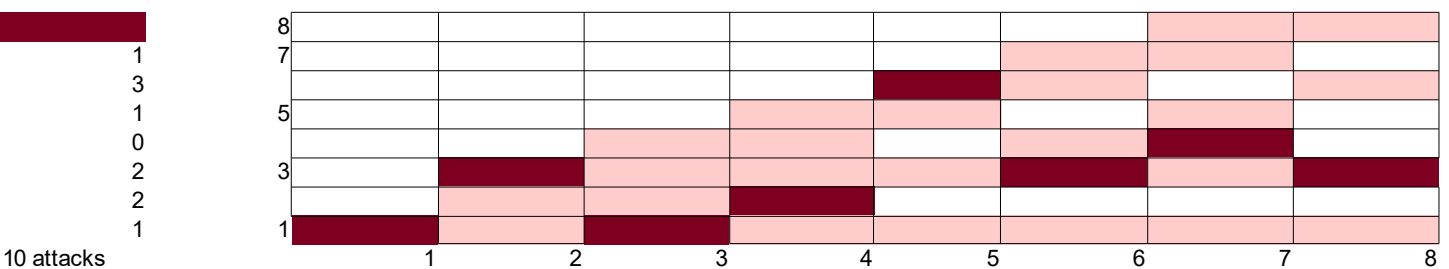
$$F_1 = f(\text{Ind}_1) = 28 - \text{attacks}(\text{Ind}_1) = 28 - 11 = 17$$

Ind_2:



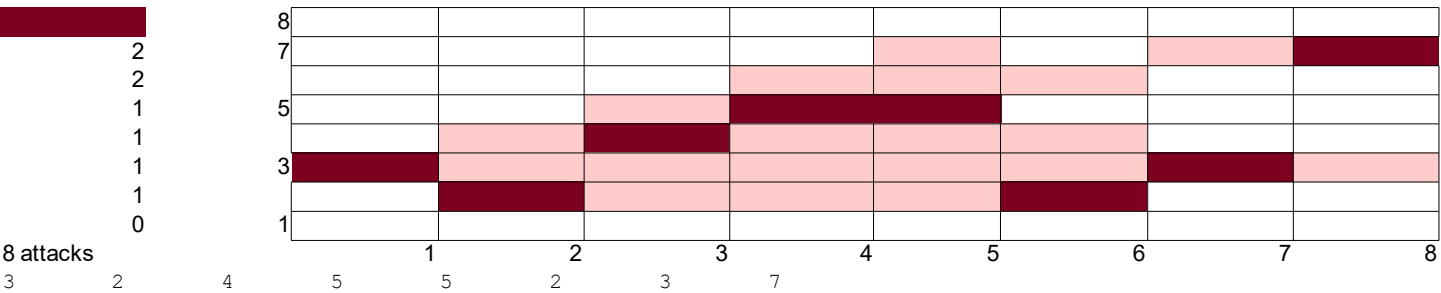
$$F_2 = f(\text{Ind}_2) = 28 - 8 = 20$$

Ind3:



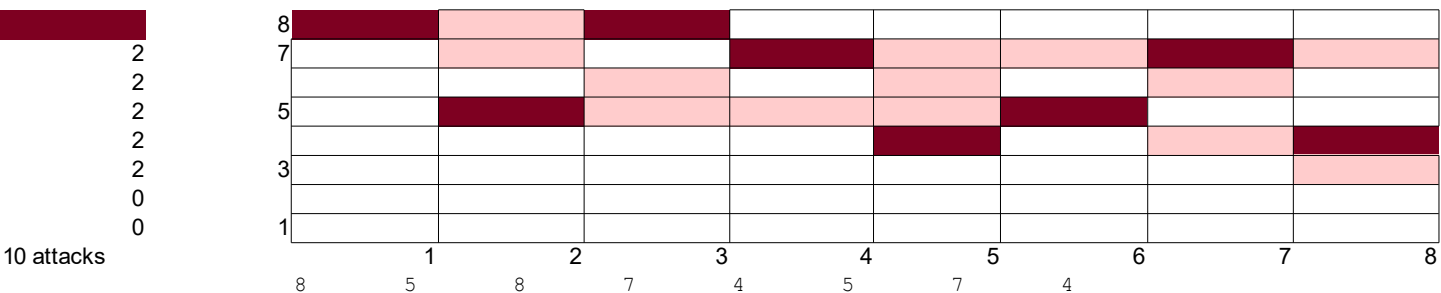
$$F_3 = f(\text{Ind}_3) = 28 - 10 = 18$$

Ind4:



$F4 = f(\text{Ind}_4) = 28 - 8 = 20$

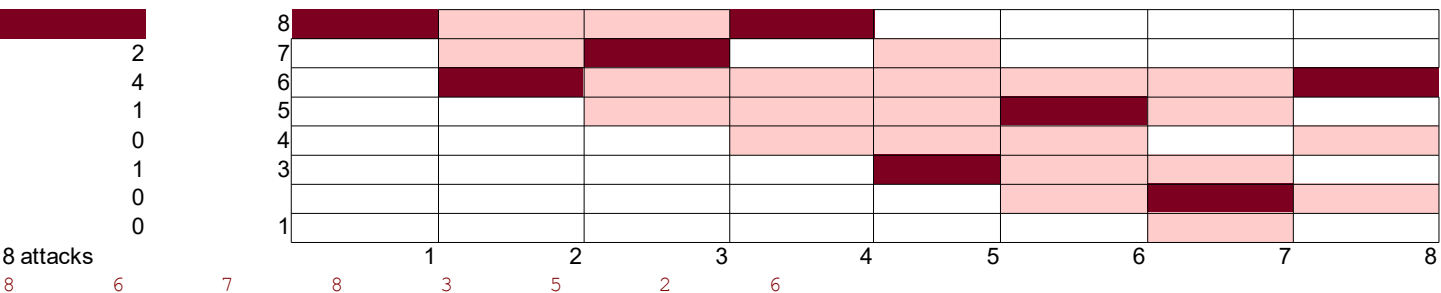
Ind5



$F5 = 28 - 10 = 18$

Ind 6

8 6 7 8 3 5 2 6



$F6 = 28 - 8 = 20$

b) Creating the roulette wheel

F1=17, F2=20 F3=18 F4=20 F5=18 F6=20

$F = F1 + F2 + \dots + F6 = 53 + 60 = 113$ (total fitness)

$S = [0, F) = [0, 113)$ ← total interval

subintervals

$S1 = [0, F1) = [0, 17)$ | $|S1| = F1$ (the length of S1)

$S2 = [F1, F1 + F2) = [17, 37)$ | $|S2| = F2$

$S3 = [F1 + F2, F1 + F2 + F3) = [37, 55)$ | $|S3| = F3$

$S4 = [55, 75)$

$S5 = [75, 93)$

$S6 = [93, 113)$ c) Running the roulette wheel

Draw 6 random numbers $r1, r2, \dots, r6$ from $[0, 1]$

→ $r1=0.76, r2=0.96, r3=0.98, r4=0.29, r5=0.65, r6=1.00$

Then we calculate:

$R1 = r1 * F, R2 = r2 * F, \dots, R6 = r6 * F$ (so $R1, R2, \dots, R6$ are from S)

$R1 = 85.88$ belongs to $S5 \rightarrow \text{Ind } 5$

$R2 = 108.48$ belongs to $S6 \rightarrow \text{Ind } 6$

$R3 > R2 \Rightarrow$ belongs to $S6 \rightarrow \text{Ind } 6$

$R4 = 34.77$ belongs to $S2 \rightarrow \text{Ind } 2$

$R5 = 73.4$ belongs to $S4 \rightarrow \text{Ind } 4$

$R6 = 113 \rightarrow \text{Ind } 6$

Hence the **Mating Pool** is: **Ind 5 Ind 6 Ind 6 Ind 2 Ind 4 Ind 6**

CROSSOVER

Pair I (Ind 5---Ind 6)

draw random $r1, r2$ from $\{1, \dots, 7\} \rightarrow r1=r2=4 \rightarrow$

parent G5 8 5 8 7 | | 4 5 7 4

parent G6 8 6 7 8 || 3 5 2 6

offspring 8 5 8 7 4 5 7 4

offspring 8 6 7 8 3 5 2 6

Pair I (Ind 6---Ind 2)

draw random r_3, r_4 from $\{1, \dots, 7\} \rightarrow r_3=3, r_4=7 \rightarrow$

parent G6 8 6 7 | 8 3 5 2 | 6

parent G2 5 6 7 | 7 2 6 3 | 6

offspring 8 6 7 7 2 6 3 6

offspring 5 6 7 8 3 5 2 6

pair III (Ind 4---Ind 6)

draw random r_5, r_6 from $\{1, \dots, 7\} \rightarrow r_5=1, r_6=3 \rightarrow$

parent G4 3 | 2 4 | 5 5 2 3 7

parent G6 8 | 6 7 | 8 3 5 2 6

offspring 3 6 7 5 5 2 3 7

offspring 8 2 4 8 3 5 2 6

Hence we've obtained the following set of

Offsprings

O1 8 5 8 7 4 5 7 4

O2 8 6 7 8 3 5 2 6

O3 8 6 7 7 2 6 3 6

O4 5 6 7 8 3 5 2 6

O5 3 6 7 5 5 2 3 7

O6 8 2 4 8 3 5 2 6

Mutation

p_m - mutation probability (for the single gene)

here (for demonstration purposes) we assume $p_m=1/15$ (for computer implementation try p_m from $[0.002, 0.01]$)

Since there are $48=6*8$ genes , we need 48 random numbers

r_1^1	r_1^2	r_1^3	r_1^4	r_1^5	r_1^6	r_1^7	r_1^8
r_2^1	r_2^2	r_2^3	r_2^4	r_2^5	r_2^6	r_2^7	r_2^8
r_3^1	r_3^2	r_3^3	r_3^4	r_3^5	r_3^6	r_3^7	r_3^8
r_4^1	r_4^2	r_4^3	r_4^4	r_4^5	r_4^6	r_4^7	r_4^8
r_5^1	r_5^2	r_5^3	r_5^4	r_5^5	r_5^6	r_5^7	r_5^8
r_6^1	r_6^2	r_6^3	r_6^4	r_6^5	r_6^6	r_6^7	r_6^8

from the set of integers: $\{1,2,3,\dots,15\}$ (as $p_m=1/15$). We assume that if $r_i=1$ then it means that the gene number i will be mutated.

Here are my random numbers

13	5	1	7	12	14	4	7
4	4	9	12	7	7	3	2
6	8	9	7	5	3	7	2
1	1	6	5	11	2	9	1
8	7	1	1	6	15	2	15
14	1	1	7	5	7	14	8

$r_1^3, r_4^1, r_4^2, r_4^8, r_5^3, r_5^4, r_6^2, r_6^3=1$

---> the genes $O1(3), O4(1), O4(2), O4(8), O5(3), O5(4), O6(2), O6(3)$ should be mutated.

We generate new random values for these 8 genes:

2	4	7	2	8	5	3	6
---	---	---	---	---	---	---	---

2

O1	8	5	8	7	4	5	7	4
N1	8	5	2	7	4	5	7	4

4 7 2

O4	5	6	7	8	3	5	2	6
N4	4	7	7	8	3	5	2	2

8 5

O5	3	6	7	5	5	2	3	7
N5	3	6	8	5	5	2	3	7

3 6

O6 8 2 4 8 3 5 2 6

N6 8 3 6 8 3 5 2 6

New population: N1,N2,N3,N4,N5,N6

N1 8 5 2 7 4 5 7 4

N2 8 6 7 8 3 5 2 6

N3 8 6 7 7 2 6 3 6

N4 4 7 7 8 3 5 2 2

N5 3 6 8 5 5 2 3 7

N6 8 3 6 8 3 5 2 6

Then we replace the old population with the new one, that is:

G1=N1, G2=N2, ...,G6=N6

... and repeat the steps (until finding a non-attacking configuration)

„Paper version”: max. 4 points ; take N=6, p_m=1/15; do 5 iterations of the algorithm