# GA FOR SET COVERING PROBLEM

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   <a href="https://github.com/Girasolo/computational\_intelligence">https://github.com/Girasolo/computational\_intelligence</a>
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   <a href="https://github.com/AlessiaLeclercq/ComputationalIntelligence\_S">https://github.com/AlessiaLeclercq/ComputationalIntelligence\_S</a>
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- 302294 Leonardo Tredese https://github.com/LeonardoTredese/s302294-computationalintelligence-2022-2023
- 292113 Filippo Cardanohttps://github.com/Frititati/CI\_2022\_292113

# **WORK GROUP**

- ▶ Idea from the onemax problem:
  - each flag correspond to one of the lists generated by the well known function problem

#### Genome

```
for genome in [tuple([random.choice([1, 0]) for _ in range(len(ALL_LISTS))]) for _ in range(population_SIZE)]:
    population.append(Individual(genome,fitness_function(genome)))
```

# REPRESENTATION

As in the onemax we have used a namedtuple

The fitness function returns a tuple (#coveredEl,-weight), which is assigned to the fitTuple

## REPRESENTATION

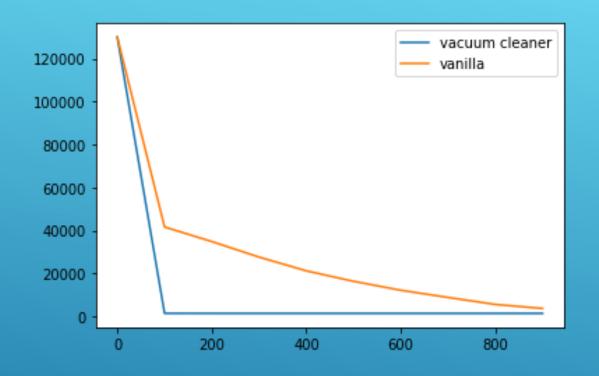
```
def GA(population, NUM_GENERATIONS, OFFSPRING_SIZE, mutation_rate,POPULATION_SIZE,cleaner_rate):
   start = time.time()
                                       #to take note of the computation time
   fittestIndividual=copy(population[0])
   for g in range(NUM GENERATIONS):
       offspring = list()
       for i in range(OFFSPRING SIZE): #generation of the offSprings
            if random.random() < mutation_rate:</pre>
               p = tournament(population)
               if g<NUM GENERATIONS/cleaner rate:
                   o= vacuum cleaner(p.genome)
               else:
                   o = mutation(p.genome)
               #o = mutation(p.genome)
            else:
               p1 = tournament(population)
               p2 = tournament(population)
               o = cross over(p1.genome, p2.genome)
           offspring.append(Individual(o, fitness function(o)))
       population += offspring #offspring added to the population
       #population=deClonizator(population)
       population=list(dict.fromkeys(population)) #deletion of duplicates
       population = sorted(population, key=lambda i: i.fitTuple, reverse=True)[:POPULATION_SIZE] #cut the population
        if(population[0].fitTuple>fittestIndividual.fitTuple): #copy the fittest individual
           fittestIndividual=copy(population[0])
```

# **IMPROVEMENTS**

```
def cross_over(g1, g2):
    cut = random.randint(0, len(ALL_LISTS)-1)
    choice=random.randint(0,2)
    if choice==0:
       return g1[:cut] + g2[cut:]
    else:
       return g2[:cut] + g1[cut:]
```

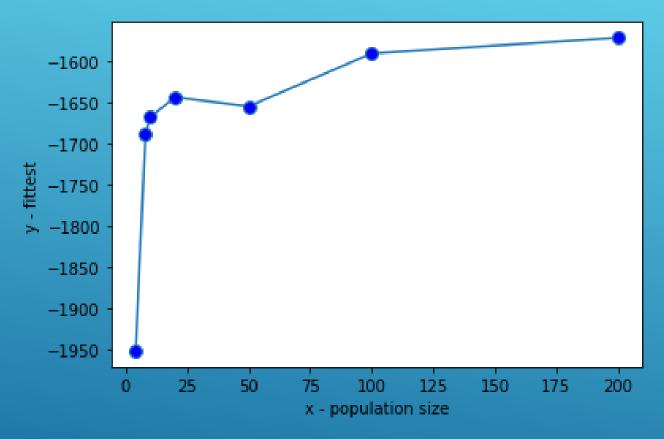
CROSS\_OVER

# VACUUM CLEANER



```
[-129941, -1480, -1460, -1460, -1460, -1460, -1460, -1460, -1460, -1460]
[-129941, -41641, -34870, -27636, -21250, -16377, -12214, -8871, -5626, -3747]
```

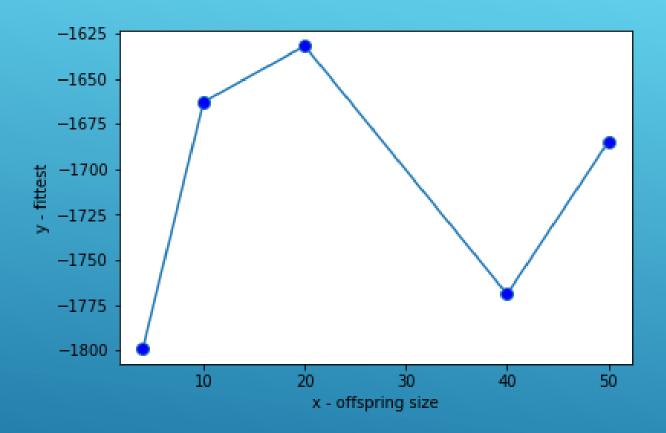
# VACUUM CLEANER



```
population_SIZE=[4,8,10,20,50,100,200]
PROBLEM_SIZE=500
OFFSPRING_SIZE=20
mutation_rate=0.3
cleaner_rate=20
NUM_GENERATIONS=1000
```

```
time: 4.03 fittest: (500, -1951)
time: 4.02 fittest: (500, -1688)
time: 3.94 fittest: (500, -1668)
time: 4.08 fittest: (500, -1644)
time: 4.55 fittest: (500, -1655)
time: 5.26 fittest: (500, -1591)
time: 7.01 fittest: (500, -1572)
```

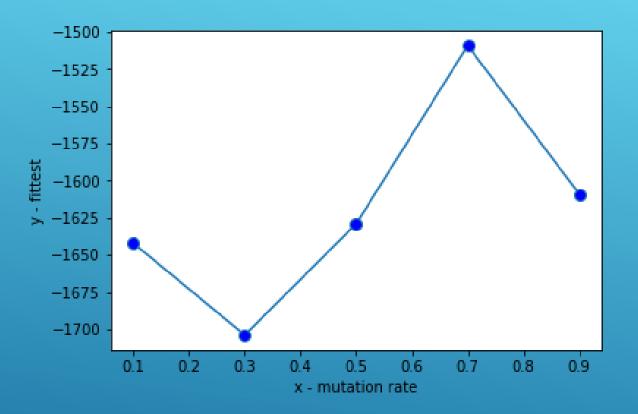
# PARAMETER STUDY:POPULATION SIZE



# PARAMETER STUDY: OFFSPRING SIZE

```
population_SIZE=20
PROBLEM_SIZE=500
OFFSPRING_SIZE=[4,10,20,40,50]
mutation_rate=0.3
cleaner_rate=20
NUM_GENERATIONS=1000
```

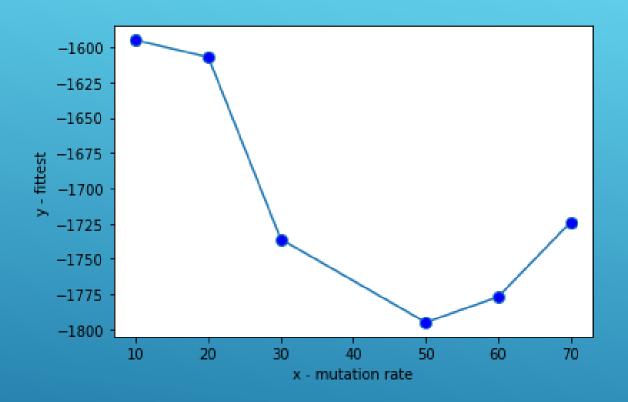
```
time: 1.41 fittest: (500, -1799)
time: 2.33 fittest: (500, -1663)
time: 4.54 fittest: (500, -1632)
time: 9.63 fittest: (500, -1769)
time: 10.69 fittest: (500, -1685)
```



```
population_SIZE=20
PROBLEM_SIZE=500
OFFSPRING_SIZE=20
mutation_rate=[0.1,0.3,0.5,0.7,0.9]
cleaner_rate=20
NUM_GENERATIONS=1000
```

```
time: 4.65 fittest: (500, -1642)
time: 4.40 fittest: (500, -1704)
time: 4.67 fittest: (500, -1629)
time: 4.99 fittest: (500, -1509)
time: 5.12 fittest: (500, -1609)
```

# PARAMETER STUDY: MUTATION RATE



# PARAMETER STUDY: CLEANER RATE

```
population_SIZE=20
PROBLEM_SIZE=500
OFFSPRING_SIZE=20
mutation_rate=0.7
cleaner_rate=[10,20,30,50,60,70]
NUM_GENERATIONS=1000
```

```
time: 6.10 fittest: (500, -1595)
time: 5.01 fittest: (500, -1607)
time: 4.94 fittest: (500, -1736)
time: 4.39 fittest: (500, -1795)
time: 4.31 fittest: (500, -1777)
time: 4.27 fittest: (500, -1724)
```

```
population_SIZE=20
PROBLEM_SIZE=[5,10,20,100,500,1000,2000]
PROBLEM_SIZE_=[5000,10000]
OFFSPRING_SIZE=20
mutation_rate=0.7
cleaner_rate=10
NUM_GENERATIONS=1000
```

```
time: 112.67 fittest: (5000, -26718)
```

peak memory: 604.85 MiB, increment: 2.73 MiB

time: 271.91 fittest: (10000, -56751)

peak memory: 2073.34 MiB, increment: 3.61 MiB

#### FINAL TRIAL

```
time: 0.26 fittest: (5, -5)
peak memory: 159.83 MiB, increment: -2.07 MiB
time: 0.22 fittest: (10, -11)
peak memory: 159.84 MiB, increment: 0.01 MiB
time: 0.24 fittest: (20, -24)
peak memory: 159.85 MiB, increment: 0.00 MiB
time: 1.14 fittest: (100, -198)
peak memory: 160.15 MiB, increment: 0.25 MiB
time: 5.27 fittest: (500, -1595)
peak memory: 161.92 MiB, increment: 1.06 MiB
time: 12.28 fittest: (1000, -3810)
peak memory: 195.11 MiB, increment: 1.64 MiB
time: 29.81 fittest: (2000, -8789)
peak memory: 309.75 MiB, increment: 4.27 MiB
```

▶ Use of binary numbers instead of an array of ones and zeros

**...** 

# FURTHER POSSIBLE OPTIMIZATIONS

```
def GAstrategy2(population, NUM GENERATIONS, OFFSPRING SIZE, mutation rate,POPULATION SIZE,cleaner rate):
   start = time.time()
                                        #to take note of the computation time
   fittestIndividual=copy(population[0])
   for g in range(NUM GENERATIONS):
       for i in range(OFFSPRING_SIZE):
           0=0
           if random.random() < mutation rate:</pre>
               p = tournament(population)
               if g<NUM_GENERATIONS/cleaner_rate:</pre>
                   o= vacuum cleaner(p.genome)
                   o = mutation(p.genome)
           if random.random() < 0.5:</pre>
               p1 = tournament(population)
               p2 = tournament(population)
               o = cross over(p1.genome, p2.genome)
           if o:
               population.append(Individual(o, fitness function(o)))
               population = sorted(population, key=lambda i: i.fitTuple, reverse=True)
               del population[-1]
           #in the strategy2 the population is cut every time we add a new individual
       #population=deClonizator(population)
       population=list(dict.fromkeys(population)) #deletion of duplicates
       if(population[0].fitTuple>fittestIndividual.fitTuple):
           fittestIndividual=copy(population[0])
```

```
time: 0.25
            peak memory: 119.12 MiB fittest: (5, -5)
time: 0.27
            peak memory: 119.24 MiB fittest: (10, -14)
            peak memory: 119.26 MiB fittest: (20, -35)
time: 0.40
time: 1.93 peak memory: 119.45 MiB fittest: (100, -257)
time: 7.78 peak memory: 126.46 MiB fittest: (500, -1908)
time: 20.20 peak memory: 154.77 MiB fittest: (1000, -4354)
time: 45.22 peak memory: 272.02 MiB fittest: (2000, -9560)
time: 112.67 peak memory: 604.85 MiB fittest: (5000, -26718)
time: 271.91 peak memory: 2073.34 MiB fittest: (10000, -56751)
 population SIZE=20
 PROBLEM SIZE = [5,10,20,100,500,1000,2000]
 PROBLEM SIZE=[5000,10000]
 OFFSPRING SIZE=20
 mutation rate=0.3
 cleaner rate=10
 NUM GENERATIONS=1000
```

#Strategy2

#### ALTERNATIVE STRATEGY: STRATEGY 2