CT421 Project 1: Evolutionary Search

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Contributions: Equal split of work, Alessio and Kamil have both worked on separate implementations for part A and eventually decided to go with Alessio's approach that can be seen on the github. Part B was developed through pair programming sessions over several days both in person and screen share sessions where we have both contributed to the final solution. The final report was concatenated by Kamil while Alessio gathered graphs and results for the report.

https://github.com/kkcavan/Alassignment1/tree/master

1.1

1. Description of your representation; fitness function; selection, crossover, mutation, and any other operations. [2]

```
#fitness function
def fitness(individual):
    return sum(int(bit) for bit in individual)
```

Simply summing each bit in a string to get its fitness out of 0-30.

```
def russian_roulette_wheel_selection(population, fitness_scores):
    fitness_addedup = sum(fitness_scores)
    selection_likelyhood = [fitness / fitness_addedup for fitness in fitness_scores]

# randomly select the strings but weigh them more with size
    selected_participants = random.choices(range(len(population)), weights=selection_likelyhood, k=len (population))

# Return the selected
    selected_population = [population[index] for index in selected_participants]
    return selected_population
Python
```

A Roulette Wheel Selection procedure that can be used within a GA algorithm. The way this procedure works can be pictured as a roulette wheel with the size of each slice being represented by the fitness of that string.[3]

```
def mutate(individual):
    mutated_individual = list(individual)
    for i in range(len(mutated_individual)):
        if random.random() < mutation_rate:
            mutated_individual[i] = '1' if individual[i] == '0' else '0'
        return ''.join(mutated_individual)</pre>
```

Mutating function that will flip a bit based on a RNG. If the randomly generated number is smaller than the mutation rate the bit will flip causing a mutation.

```
def crossover(parent_one, parent_two):
    if random.random() < crossover_rate:
        crossover_index_point = random.randint(1, len(parent_one) - 1)
        child_one = parent_one[:crossover_index_point] + parent_two[crossover_index_point:]
        child_two = parent_two[:crossover_index_point] + parent_one[crossover_index_point:]
        return child_one, child_two
    else:
        return parent_one, parent_two</pre>
```

A crossover genetic operator. It is used in this GA to combine "genetic" material from two parents. The rate at which this crossover happens depends on the crossover_rate. This is a likelihood of the crossover happening. If it happens that the RNG is smaller than the crossover rate, a point is randomly selected in the string. The offspring will be created and will get the traits of one parent up to the point in the crossover string and the rest will come from the other parent. Otherwise if the rng did not cause a cross over we will return the parents without any modifications.[5]

```
# Main loop
best_fitness_list = []

for generation in range(generations):
    # Calculate fitness for each individual
    fitness_scores = [fitness(individual) for individual in population]

# Calculate best fitness
best_fitness_list.append(max(fitness_scores))

# Select parents based on fitness values scores from using Roulette Wheel Selection
selected_population = russian_roulette_wheel_selection(population, fitness_scores)

# Create the next generation
new_population = []
for i in range(0, len(selected_population), 2):
    parent_one, parent_two = selected_population[i], selected_population[i+1]
    child_one, child_two = crossover(parent_one, parent_two)
    child_one = mutate(child_one)
    child_two = mutate(child_two)
    new_population.extend([child_one, child_two])

population = new_population
```

This is where the core of the GA happens and everything comes together. The GA will run over a predefined number of epochs where a generation will be created. The fitness function is employed to calculate the fitness of each individual string.

The fittest individual strings are found through the max function. This is a representation of the strongest in the herd (almost). We will append these to the most fit list.

Following this we select a number of parents through the roulette wheel, where we hope to mainly land on the fitter individuals and select them as parents.

The selected parents will go through a process of crossover and mutation operations to generate a new and hopefully stronger offspring for the next population.

The offspring will be a combination of genes through crossover followed by a chance of mutation. The offspring then enter the new population.

The old population is replaced with the new population and the cycle can repeat.

In general, high mutation will increase exploration, and high crossover will increase exploitation. So, this means that exploration will come up with new solutions to the problem that have not been seen by the GA before. Whereas exploitation will take the currents solutions and take advantage of solutions.

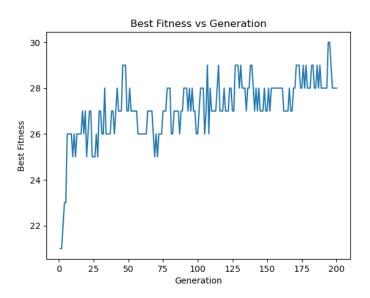


Figure 1: $mutation_rate = 0.01\ crossover_rate = 0.8$. We reach a local optimum around the 50^{th} generation. Due to the mutation rate we introduce novelty into the solution space which caused a decrease in fitness as seen by the low valley at generation 75. Eventually we reach the highest fitness level at generation ~190.

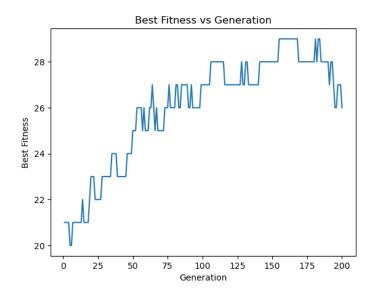


Figure 2: mutation_rate = 0.01 crossover_rate = 0.01.

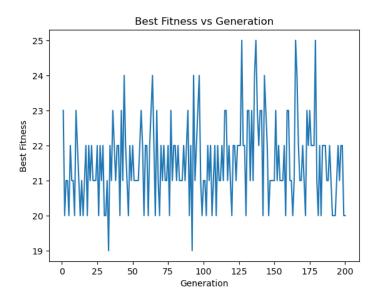


Figure 3: mutation_rate = 0.8 crossover_rate = 0.01

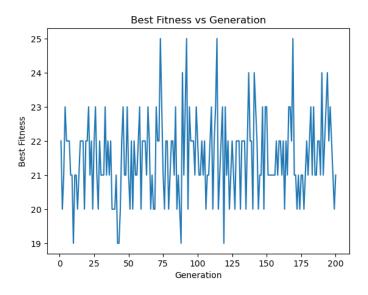


Figure 4: mutation_rate = 0.8 crossover_rate = 0.8

Results: Through trial and testing we have found that the combination of a low mutation rate and a high crossover rate has led to the best performance in this GA problem. This problem solution changes greatly when tweaking the mutation and crossover variables as seen in figure 2,3,4. High mutation rate leads to a very spikey solution with lots of peaks and valleys that indicate that the algorithm has a hard time ending in a plateau.

1.2

1.3

Part B:

Insight into problem landscape: computationally the bin packing problem is known to be np-hard so by this we know that is in a class of very computationally challenging problems in which there is no efficient algorithm that can solve all instances optimally. This is due to the sheer number of combinations that need to be considered. Exact solutions are impractical for such a problem. Thus a way of finding a solution for a problem in which capacities of bins and number of items change, we tend to use genetic algorithms to find viable solutions.[1]

Initially to load the dataset into a jupyter notebook we have split the dataset text file into separate files for each binning problem and converted them to csv files. Using the pandas library we read in each file and create a dataframe out of them, renaming the columns to Weight and num_items respectively. We then flatten the dataframe into one column by multiplying the weight by the num of items.

```
def fitness_function(item_list, max_bin_capacity):
    number_of_bins = 0
    current_capacity_of_bin = 0

/ for item in item_list:
    if current_capacity_of_bin + item <= max_bin_capacity:
        # Add the size of the current item to the current bin capacity
        current_capacity_of_bin += item
    else:
        # If adding another item exceeds the bins capacity we create a new bin
        number_of_bins += 1
        # here we bring back the current_capacity_of_bin to the size of the current item
        current_capacity_of_bin = item

return number_of_bins

</pre>
```

The fitness function: we have defined num_bin and current_capacity. We iterate though each item (weight) in a list of all items. If we add an items weight to the current capacity of a bin and it fits, we add that item to the capacity. Otherwise, we will create a new bin and set the capacity of the bin to the new item.

The mutate function works similarly to the previous problems. If the mutation rate criterion is met, the gene index is mutated to a random index, which in this context means placing an item into a different bin.

```
def crossover(parent_one, parent_two):
    if random.random() < crossover_rate:
        crossover_index_point = random.randint(1, len(parent_one) - 1)
        child1 = parent_one[:crossover_index_point] + parent_two[crossover_index_point:]
        child2 = parent_two[:crossover_index_point] + parent_one[crossover_index_point:]
        return child1, child2
    else:
        return parent_one, parent_two
</pre>
```

A function that is responsible for creating new offspring solutions through combining genetic material from two parents. It works on the same principle as the above problems, we will find a random index in a list and replace it to that point with one parent and the rest with the other parents. So, we are basically combining two solutions by splitting them at some point and giving each child a random proportion of its parent's solution.[4]

We perform a selection process two times to find 2 parents for a crossover. We then randomly sample 2 individual solutions by rng(twice). These 2 solutions will "fight". Leading to the strong individual being selected, This means that most full bins will win and then since we run this twice, we will have 2 parents selected for a crossover.

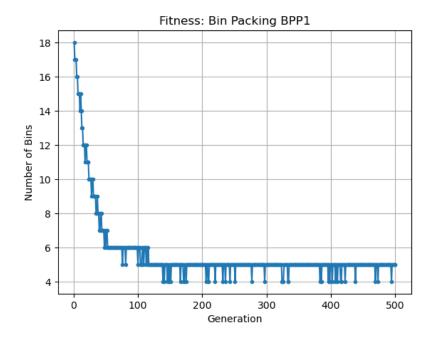
```
def elitism(population, fitness_values, elitism_rate):
    number_of_elite_solutions = int(len(population) * elitism_rate)
    elites_selected = sorted(range(len(fitness_values)), key=lambda i: fitness_values[i], reverse=True)[:number_of_elite_solutions]
    return [population[i] for i in elites_selected]

✓ 0.0s
```

Our solution implementation contains elitism too. This is to carry good generations forward with no change. This will assist us in carrying over good solutions. The function ranks the fitness values of solutions and preserves a list of elite solutions.

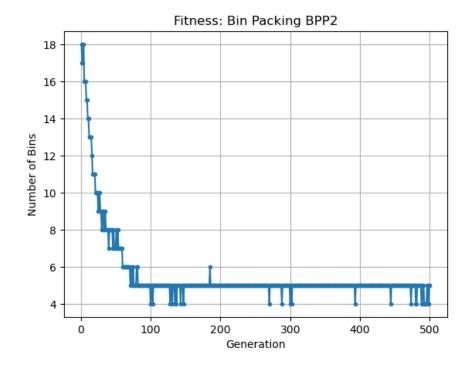
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### Images | Interpretation | Interpreta
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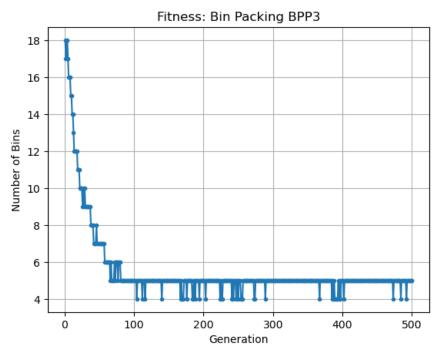
The main function of our genetic algorithm as a function. We can see in what order the GA runs. We start packing one item per bin so we'll have the max number of bins. That's why the high spike at the beginning of the graph. Then we enforce elitism mutation and crossover in order to have a better packing. That's why the decreasing curve. Eventually it will find the local optima then it will go down. This happens from generation 100 onwards.

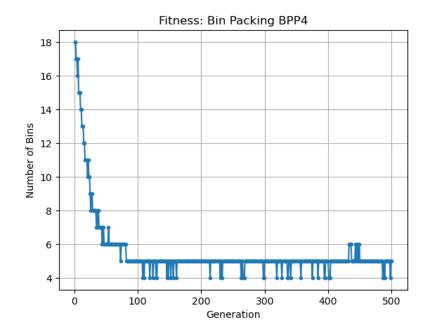


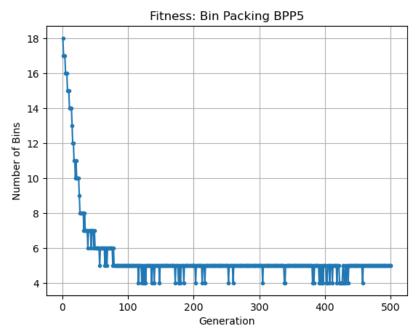
4: min bins is the best solution we can achieve

5.624: average number of bins









References:

- 1. https://en.wikipedia.org/wiki/Genetic algorithm
- 2. https://www.geeksforgeeks.org/genetic-algorithms/
- 3. https://dev.to/rpalo/python-s-random-choices-is-awesome-46ii
- 4. https://www.geeksforgeeks.org/python-single-point-crossover-in-genetic-algorithm/
- 5. https://machinelearningmastery.com/simple-genetic-algorithm-from-scratch-in-python/