

Faculty of Engineering and Technology Electrical and Computer Engineering Department

ARTIFICIAL INTELLIGENCE, ENCS3340 Project #1

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Abstract:

This project aims to use Genetic algorithms to build a "Graduation projects distribution" program.

The program will read inputs from two files:

- Groups selections (Excel File)
- Projects (PDF File).

Then it will assign a project for each group so that:

- The assignment must be closed as much as possible to the order of selection.
- The minimum number of conflicts that can occur between groups selections.

Problem Formalization:

For this project, I used Java language + scene builder.

In genetics algorithm, the chromosome represents a solution,

The solution here is an array of integers, the index of the array represents the group number (id), and the value of that index represents the topic choice for that group.

A population has many chromosomes sorted by fitness.

The fitness of each chromosome = the number of conflicts between group choices.

The population sorts chromosomes ascending by their fitness value.

If two chromosomes have the same fitness value, we compare them by the benefit value.

The benefit of each chromosome = for each index in the array

if it was the first choice add 3

if it was the second choice add 2

if it was the third choice add 1.

We sort them descending.

Code Explanation:

1. Project Class:

Each Project has a project number (pid), project description that has the name and description of that project. we use this class to define Project instances.

2. Group Class:

Each Group has id, Student1, student2, and student3, 3 choices: option1, option2 and option3. we use this class to define Group instances.

3. Chromosome Class:

A chromosome represents a solution,

Each one has an array (int[]) of genes, the index of the array represents the group number (id), and the value of that index represents the topic choice for that group.

Array list of Groups.

Fitness double, Benefit int.

initialize function (initalizeChrom): for every group in the array genes choose a random value between the three options.

Calculate fitness (calcFitness), calculate benefit (calcBenefit),

Compare function that follows the order we explained before.

4. Population Class:

Each population has an array of chromosomes, and Array list of Groups.

initialize function (initalizePopul): for every index in the array define a chromosome and initialize it, then calculate its fitness and benefit. At the end sort the chromosomes in the array by the compare function we define before in chromosome class.

Sort by fitness function (sortByFitness).

5. Genetics Class:

This class represents the algorithm, It has Static final integers:

- Population_size: number of chromosomes in the population.
- Num_of_best_chromosomes: number of chromosomes in the population that will not be changed during reproduction.
- Mutation_Rate: the rate to accept the mutation.
- Tournament_Size: the number of chromosomes selected during tournament selection.

and Array list of Groups.

Functions:

- Population reproduction (Reproduction): apply crossover and mutation over the population.
- ➤ Population crossover(populationCrossover): the first chromosomes are not changed; the remaining chromosomes are replaced by a crossover between two tournament selection chromosomes.
- ➤ tournament selection(tournamentSelection): selects number of chromosomes randomly from the old population and return the best one of them.
- Chromosome's crossover(chromosomeCrossover): create a new chromosome and for its genes array it selects each value randomly between the two old chromosomes.
- Population mutation (populationMutation): the first chromosomes are not changed, the remaining chromosomes we apply chromosome mutation on them.
- Chromosome's mutation (chromosomeMutation): create new chromosome and for each value in the genes array if (math.random) was below the mutation rate we choose the value randomly between the options for that group, if it was above the rate, it has the same value for the old chromosome.

6. Driver Class:

This class just starts the interface from the "layout.fxml" file.

7. Controller Class:

This is the main class.

First, it calls two functions:

Read excel file(readExcelFile): it reads the excel file and for each row, it defines a group instance and adds it to an array list, at the end, it returns the array list.

Read PDF file(readPdfFile): it reads the pdf file and for each row, it defines a Project instance and adds it to an array list, at the end, it returns the array list.

Is numeric (isNumeric): receive a string and check if it's an integer or not.

We initialize a new population and iterations counter. We define the labels and text fields ... etc. for Javafx. Button functions:

- Show projects(showProjects): reset the scroll pane at the top (same for all buttons), for each project in the arraylist print the value on the label.
- > Show Groups(showGroups): for each group in the arraylist print the value on the label.
- ➤ Generate Solutions (generateSolutions): every time we click the button we define a new population, reset the counter, and while (the fitness value is not zero and the counter didn't reach the limit) we apply reproduction on the population and increase the counter, In the end, for each Chromosome in the population print its values, fitness, and benefit on the label.
- > Show Best Solution (showBestSolution): we print the best chromosome values and their fitness on the label.

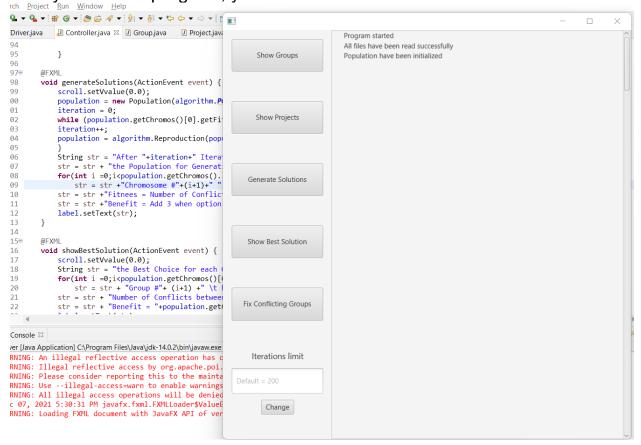
Note: other functions are described in EXTRA.

Compilation

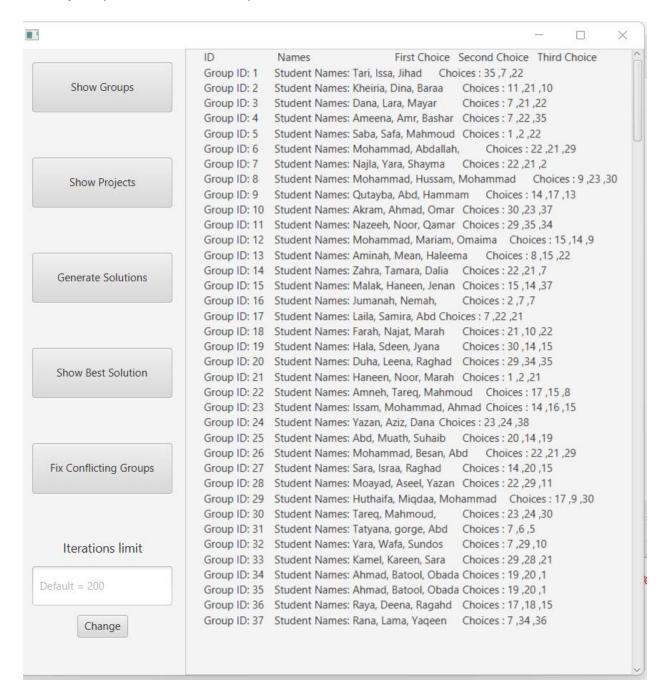
Note that before running the program you need to add these libraries to the project:

- All JavaFX and FXML libraries.
- commons-collections4-4.1
- pdfbox-app-2.0.4
- poi-3.17
- poi-ooxml-3.17
- poi-ooxml-schemas-3.17
- xmlbeans-2.6.0

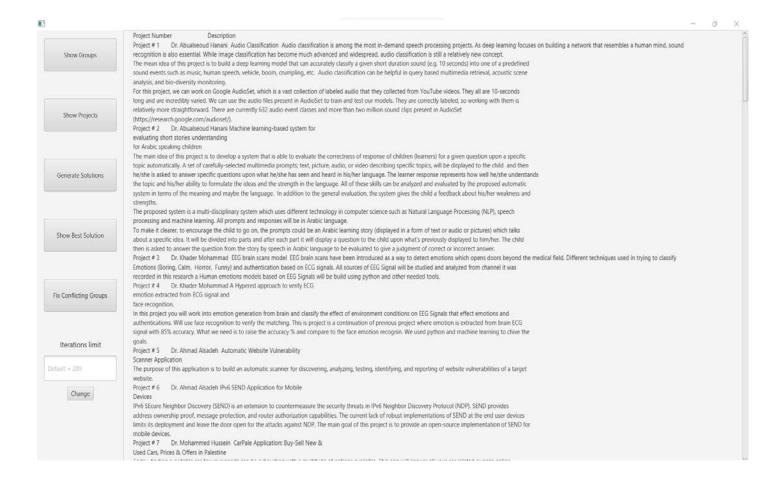
after you run the program, you should have this interface:



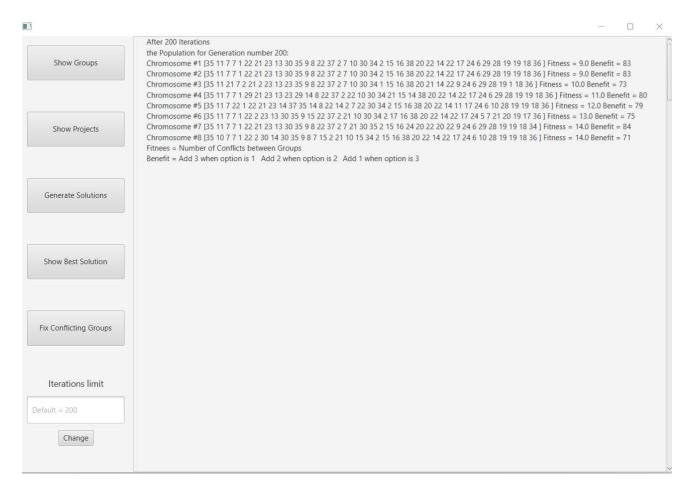
After you press Show Groups:



After you press Show Projects:

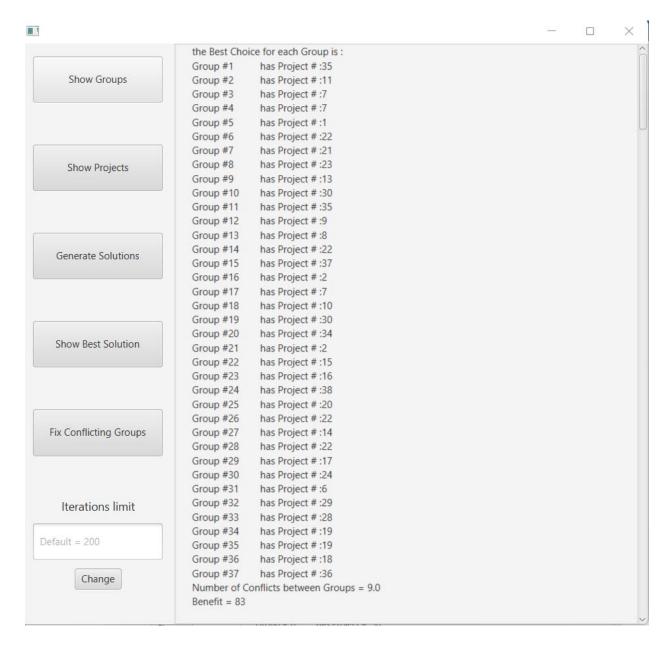


After you press Generate Solutions:



After it enters the while loop this is the output. every time you click the button you will have different output. Note that it shows the population for the current generation, so every time you change the population you will see the last one.

After you press Show Best Solution:



These are the values for the best chromosome in our population.

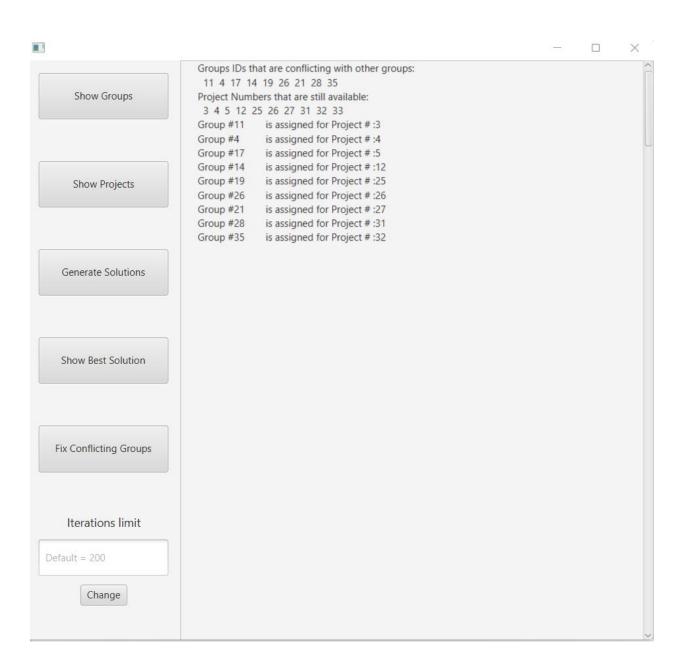
Extra

1) Fix Conflicting Groups:

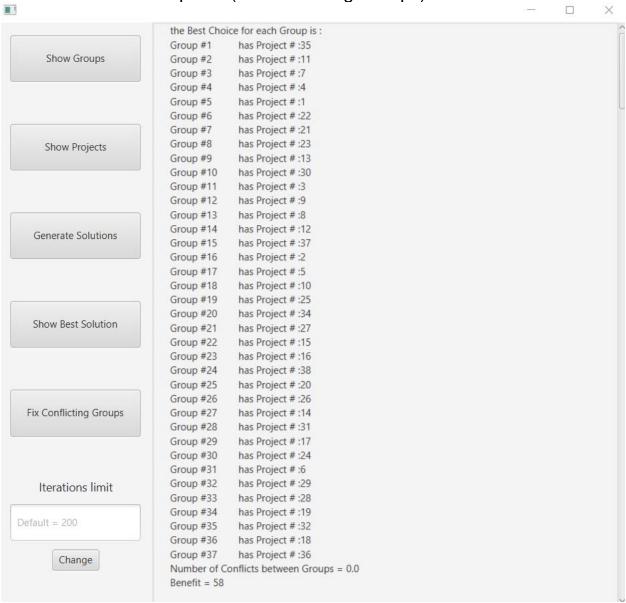
It shows you the group choices that are chosen before by another group (That creates conflict).

Also, it shows you the project numbers that are not chosen by anyone (Still available).

Then it automatically assigns both together so now the solution won't have any conflicts.



The solution after we press (Fix Conflicting Groups):



Note that the benefit value has decreased because we assigned projects for groups that are not part of their choices.

2) Change iterations limit:

The user also can change the number of iterations that reproduce the population with the value he wants.

