User Instructions

- To run the project just:
 - 1. download the contents of the folder "UserDemoPackage"
 - 2. note that the **lib** folder found in the package must be in the same folder as the **GeneticMusic.jar**
 - 3. make sure the jar fie is executable
 - 4. go to the folder of the downloaded content
 - run the project with java -jar "GeneticMusic.jar"
- this runs the project with the default parameters:

```
populationSize = 40;
chromossomeSize = 8;
numGenerations = 500;
selector = "tournament";
tournamentk = 3;
tournamentp = 0.7;
```

- to run the project with different parameters just use the following schema:
 - o **java -jar "GeneticMusic.jar"** <population size> <chromossome size> <number of generations> <[tournamet | bestChromossome]> [tournament k] [tournament p]
- examples:
 - o java -jar "GeneticMusic.jar" 40 8 500 tournamet 3 0.7
 - o java -jar "GeneticMusic.jar" 40 8 500 bestChromossome

Note: you need to provide all the parameters by the order specified if you are using the tournament selection operator. If you want to use the bestChromossome selector ignore the last 2 parameters. Again, if you provide no parameters the project will run with the default configuration.