# Report

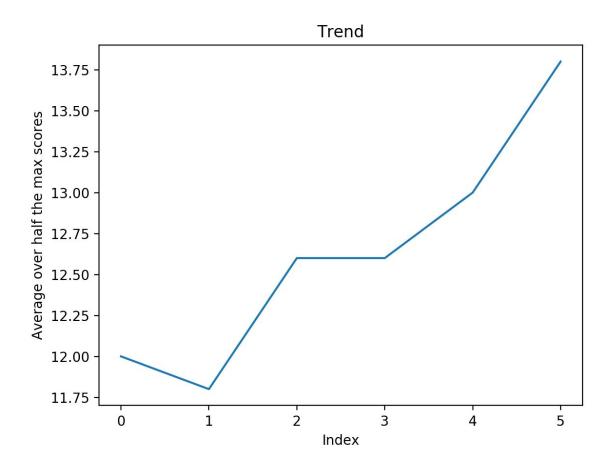
# Genetic Algorithm Project

Pratyusha Karanam & Prerit Pathak Data Science in BioInformatics

University of Florida

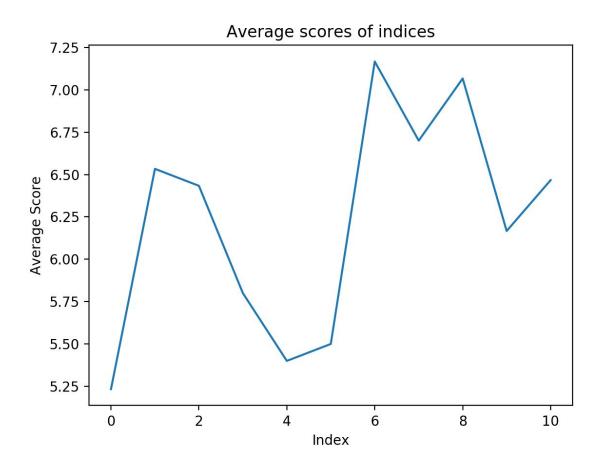
The number of sequences generated is specified as a part of the command while trying the script. When this number is set to be 30, the following graphs are generated:

### TREND ANALYSIS



The first graph is built to analyse the trend. This plot shows the value of average of half the max scores at a particular index. The indices are represented on the x axis whereas the average over half the max scores value is represented on the y axis. We observe that this is overall a rising graph.

# **AVERAGE ANALYSIS**



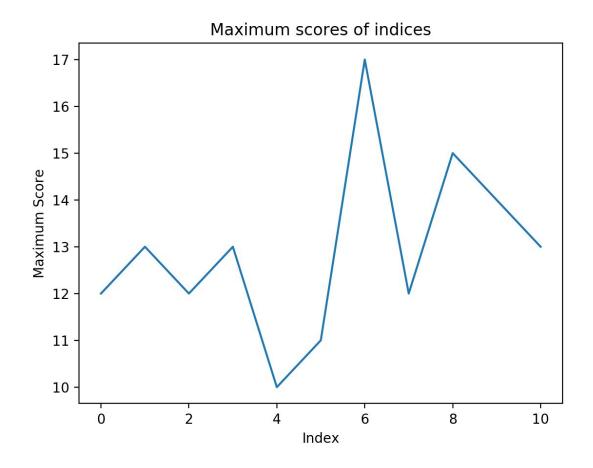
The second graph shows the average score of indices. The indices are represented on the x axis and the average score is represented on the y axis. This graph can be referred to in order to get a basic idea about the average results of our project.

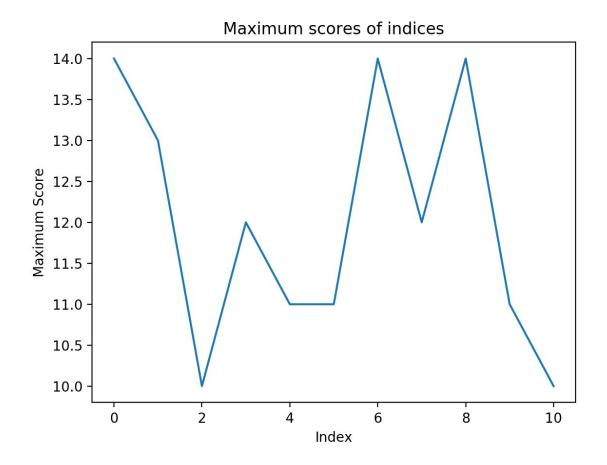
## COMPARISON OF MAXIMUM SCORE PER INDEX

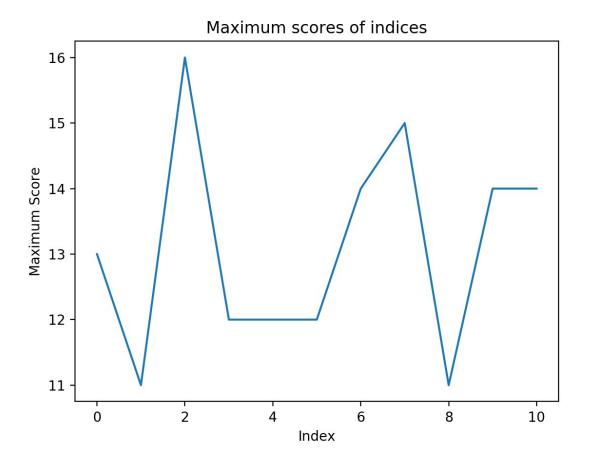
In this part of the visualisation, we generated graphs for various N values; where N = number of sequences generated.

Each graph represents the indices on the x axis and their corresponding max scores on the y axis.

N = 30 & Mutation Probability = 0.01







Hence, we can observe the drastic variation in the graphs when we change the value of N.