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Computational Biology

### Project Report

The goal of this project was to implement the cyclopeptide sequencing for an ideal spectrum. I then also add the goal of reaching and implementing the leadership cyclopeptide sequencing with an experimental spectrum as well.

The data was acquired from the Rosalind website. The data is just in the form of list variable that is defined in my main function. This is the same for the leadership function. The data is just in the form of list variable that is defined in my main function. The data I used for the ideal spectrum was [0, 113, 128, 186, 241, 299, 314, 427]. The data I used as the experimental spectrum was [0, 71, 113, 129, 147, 200, 218, 260, 313, 331, 347, 389, 460].

This is the pseudocode for the project, below this I go into detail of the high level steps I have bolded the parts I talk about in the high level steps.

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