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December 2014

Bioinformatics Final Presentation

1. An appropriate, informative project name: Chimera
2. An introductory paragraph describing your project.
3. A detailed description of the interface for your project.
4. Step by step description of the organization of your project from design to final product(code).
5. Specs for each step in 4.
6. The division of labor among the project team members.
   1. We worked side by side for every step of the project. We both took turns writing code and documentation. There was no other division of labor.
7. Any special coding features you use/develop.
   1. We implemented the BioPerl library in Perl which really helped us compute the necessary biological information. There are several interesting modules that allow you to work with sequence alignment and tree building.
8. A graphic of your code organization.
9. Testing results
10. Code
    1. See attached file folder for “Hybrid Species Research”