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Final Project Update

1. My project is now more oriented towards a focus on multiple alignment, and construction of synteny blocks for genome rearrangement. It will still have a heavy usage of GraphSpace, but it will be much less focused on pure alignment and instead as an alternate visualization for displaying synteny blocks and genome rearrangement.
2. I have not yet collected data.
3. I have not yet collected data.
4. All of it.
5. Most likely figuring out multiple alignment and genome rearrangement - adaptation to GraphSpace will likely be easier, but will require a significant amount of brainstorming for graphical usage. Multiple alignment itself, as well as determining how to do the reversals necessary for genome rearrangement will be difficult as well.