Agenda for weeks going forward

Notes 10\_15\_19

* Start collecting sequences from the sodalist clade organisms
* Check for primary organisms and add those to the list of organisms
* Look for genes within the FliE-R operon suing Mauve progressive aligner gbk files as inputs (aligned to the *Praecaptivu*s genome)
* Look for genes within the FliE-R operon using Kegg database
* Look for genes using tBLASTn using the protein sequence for *Sodalis praecaptivus* as input
* Write a methods section defining the methods for each of the three steps listed above. Include parameter decisions and a brief description of the methods advantages/disadvantages.

Agenda 10\_24\_19

* Go over specific +/-/P results from the Kegg search
* Go over how to start writing the introduction/background for the thesis

Agenda 10\_28\_19

* Go over blast results
* Examine Written Methods section
* Examine all Mauve runs with parameter specifications
* Discuss first draft of Introduction specific to be determined