1. I have downloaded the data from the Chinese database that I was looking at. I am also going to download the data from the pathogen database and it appears that it comes from the SARS database so I might need some help in knowing what exactly I am looking for since I have never downloaded anything from there.
2. At this moment, I don’t need any additional resources.
3. I still feel like my project isn’t enough. I am looking at frequency data from the Chinese database and trying to come up with some kind of test from and then I am going to make a network using Cytoscape to how gut bacteria has a relationship to foodborne bacteria that are from the same enterotype. For this, I obviously can’t compare whole genomes since they are so large (unless I can somehow?), so I think I need help in choosing the right genes to compare.