Hackathon 8/21 goals

1. ~~Make sure we have a shared github repo to work on, that we can both push and pull to~~
2. ~~Load in data~~, ~~visualize data~~ and ~~uncertainty~~
3. ~~Define the model~~
4. ~~Define priors~~
   1. ~~Init file added with strain specific values (strain and org column in init file)~~
5. Find ‘initial’ parameter guesses
6. ~~Fit with ODElib and look at posterior distributions, comparing to priors~~

Hackathon sept 5th goals

Katie TODO

Ordering and formatting - priority

1. Structure and comment all code that is being used
2. Remove all code that is not being used into a ‘dump’ folder or similar
3. Decide which figs and formats will be either in paper or supplement. Format these figs and finalize, and delete ALL OTHER figs
4. For main paper or supplement figs, label all axes, properly format in general

Stats stuff (least priority)

1. Visualize residuals

Rest of paper

1. HEPES dynamics
2. Biotic HOOH spike assays