WEEK 1 Research TODO

Week 1

• Broad goal: Fix code. ✓ Use less filtered dataset Now at 30% with 39 taxa. ✓ Make graphs of parameter values at each iteration - Did a lot of graphs, histograms and changes of betas between loops. Need to do final after changes are made. ✓ Try stepwise update - Using .1 as a constant. Still doesn't work, so likely a more complicated approach would not be helpful. Fix problem where R inverse and others not updating in beta loop Change R-inv function to depend on alpha instead of X and beta since already calculated. Flip order of beta and phi loops. ✓ Look at residuals in each iteration. (in phi step) ☐ Keep something constant through loop and/or try using identity for R. ✓ Try different dataset? Maybe simulated one from previous paper. - Same troubles ☐ Distance matrix argument to dist? - Need both forms, so may or may not be useful \square Check if phi is correct now, and see how is used Figure out large alpha problem - Seems the + and one loop beta fixed this? ☐ Try a different covariate? \square Figure out why geem code uses + instead of - on update I think + is correct. At least it converges. ☐ Identify which covariate is causing the trouble. ☐ Try removing the single OTU that seems to be contributing to problem. \square Save everything to list correctly to look at later. - Saving rho, omega, phi, beta, diff (abs), num iterations - Also do residuals. Which residuals? Cross product ones? Squared ones? Standardized squared ones? Probably standardized but not squared. Should be distributed around zero.? ✓ Plot lists - Plotted rhos, omegas, phis, diffs, not sure what else to plot. ✓ Remove scalar update. - Doesn't work. see journal \square Write up algorithm for Yuan to go through? Ask to check. \square Final run through before meeting of graphs and numbers ✓ Move timing into function

WEEK 2 Research TODO

Week 2

 $\hfill\square$ Wednesday

Main tasks:

•	10-do from sneet
•	Prepare presentation
	Try different covariates (parasite burden/antibiotics/exposure)
	Think about genus level analysis/ phylogenetic tree for taxonomic units
	Choose α automatically (simple line search)
	Make presentation