* **Fac\_comp simulation have a speciation with divergence, creating two new mutants, whereas foodweb only creates one. I need to add the speciation type I do for fac and comp to the foodweb model.**
* **Add the ancestry table computation to the foodweb simulation -> test that it works to compute the phylog signal with the results from the simulation once implemented.**
* **After having the foodweb model with the ancestry table on it and the good speciation form, I need to build the Neutral version, and the 20-80, 50-50, 80-20 neutral vs selection versions.**
* **Do this for fac and comp too.**
* **Added ancestry table computation to function simulation fw but now error when computing phylogenetic signal – ancestry table and presence df doesn’t match.**
* **When computing phylosign, I not only discard the timesteps for which I have less than 3 species. I discard the last timestep for which I don’t have 3 species. I need to modifcy this so that it simply subsets those with 3 species but not eliminate all of them until the one that has more, otherwise Im missing some.**