

Species abundances by each habitat type.

These are all made using Ed’s code, which flipped the similarity values from positive to negative…. So a value of 0 is completely similar and 1 is dis-similar. Only the lm model for edge distance of 100m is significant (p = 0.045, adjust R2 = 0.147). Plots and models by forest type look similar to the first graph (which is the whole forest), just a flat line. I believe X axis is meters, but not totally sure, as I just used Ed’s code.

Full with all sites Forest:



Data Binned by Edge distance:

