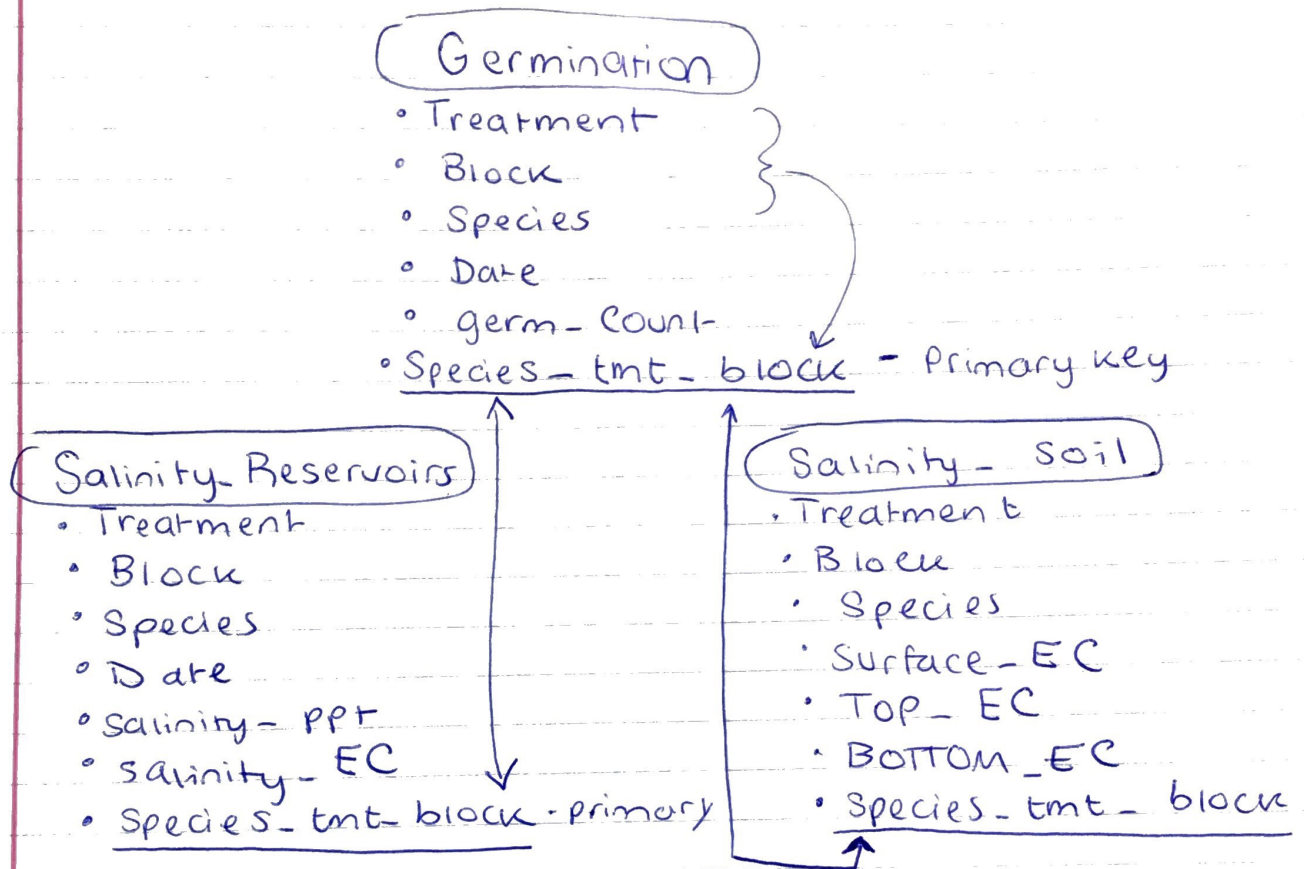


# Database structure sandra's experiment



> germination counts, and salinity-reservoirs are taken weekly. Salinity-soil will only be done once @ the end of the experiment.

> I probably could use some other unique identifier than species-tmt-block, but that would be the clearest way for me to visualize that data point (and I would still want separate columns for species, treatment, block for analysis in R).