## Tasks:

* **Assemble scaling richness dataset**
  + **Make code used to assemble richness at scales > 1m2 more robust (don’t use numbering of rows)**
  + Double-check work
* Estimate aggregation of different species across different sampling scales
  + Compile list of aggregation by species
  + Estimate aggregation using different functional groups
    - Forb, legume, grass
    - Native/exotic provenance
* **Associate XY coordinates with each plot/subplot**
  + Check on MCL site2 subplotres2 distribution of subplotres .5
* Construct semi-variograms/correlograms using:
  + Abundance of different species and functional groups
  + **Bray-curtis dissimilarity between communities relative to distance**
* Clean data
* Assemble better species list
  + Correct questionable IDs