Order of scripts

1. BC\_BetaDeviationCalculation
2. RC\_BetaDiversityCalculation
   1. Uses RaupCrickFunction\_Chase\_etal2011
3. NMDS&PERMANOVA\_Fig3
4. dbRDA\_VariancePartitioning\_LMM\_Table1Fig4
5. GowerPhysBiotVar
6. Bootstrap&Graph\_Fig2
7. SppRichness\_AppS2\_FigS4
8. IndicatorSpecies&TraitAnalysis\_Table2Fig5