Folders

* JAGS: JAGS scripts
  + Median: model\_gcm\_de.txt, model\_gmm\_de\_G2.txt, model\_gmm\_de\_G3.txt
  + Normal: model\_gmm\_de\_G2.txt, model\_gmm\_nm\_G2.txt, model\_gmm\_nm\_G3.txt
  + T-dist: model\_gmm\_nm\_G2.txt, model\_gmm\_td\_G2.txt, model\_gmm\_td\_G3.txt
* Rcode: R scripts for running the JAGS scripts
  + Rivanna: main R scripts for running JAGS on Rivanna
  + Sources: R scripts for calculating model comparison indices (DIC, WAIC, LOOCV).
    - Median:
      * GCM: de\_WAIC\_LOOCV\_DIC\_gcm\_RV2.R
      * GMM with 2 classes: de\_WAIC\_LOOCV\_DIC\_gmm\_sameSig\_G2\_RV2.R
      * GMM with 3 classes: de\_WAIC\_LOOCV\_DIC\_gmm\_sameSig\_G3\_RV2.R
    - Normal:
      * GCM: nm\_WAIC\_LOOCV\_DIC\_gcm.R
      * GMM with 2 classes: nm\_WAIC\_LOOCV\_DIC\_gmm\_sameSig\_G2.R
      * GMM with 3 classes: nm\_WAIC\_LOOCV\_DIC\_gmm\_sameSig\_G3.R
    - T distribution:
      * GCM: td\_WAIC\_LOOCV\_DIC\_gcm\_ver3\_RV.R
      * GMM with 2 classes: td\_WAIC\_LOOCV\_DIC\_gmm\_sameSig\_G2\_RV.R
      * GMM with 3 classes: td\_WAIC\_LOOCV\_DIC\_gmm\_sameSig\_G3\_RV.R