READ ME

Model\_scripts – 3 photosynthesis models and 3 implementation scripts. These codes have been posted to github as part of manuscript submission. Data file for model in data folder, Full\_data\_complilied\_Pleban\_Guagagno\_Summer2017.txt , is 4 genotypes, ACi

* 1. CiCc\_Temp\_Jm\_model\_H.R uses CiCc\_Temp\_Jm\_model\_fixedCCrit\_script\_4\_server\_ACi\_LR\_no\_late.R implementation script following FvCB model of manuscript
  2. GECF\_hier\_model.R uses GECF\_hier\_script\_4\_server\_ACi\_LR\_no\_late .Rimplementation script following Yin model of manuscript
  3. GECFextendedfull\_hier\_model.R uses GECFextendedFull\_hier\_script\_forServer\_ACi\_LR\_no\_late .R implementation script following *βPSII* model of manuscript