* **Parkinson’s Disease**
  + **Nalls et al. (2019)**
    - Need to add rsids to full SS file (Towfique will do this)
  + **23andMe**
    - LRRK2
      * SusieR identified rs[#####] as the variant with the top PIP, in contrast to the GWAS where the lead variant was identified as rs[#####]. Specifying that there was more than one causal variant (L>1) prevented the model from converging.
* **Alzheimer’s Disease**
  + IGAP
  + Marioni et al . (2018)
  + Posthuma et al. (2018)
* **eQTL**
  + MESA
    - AFA
    - CAU
    - HIS