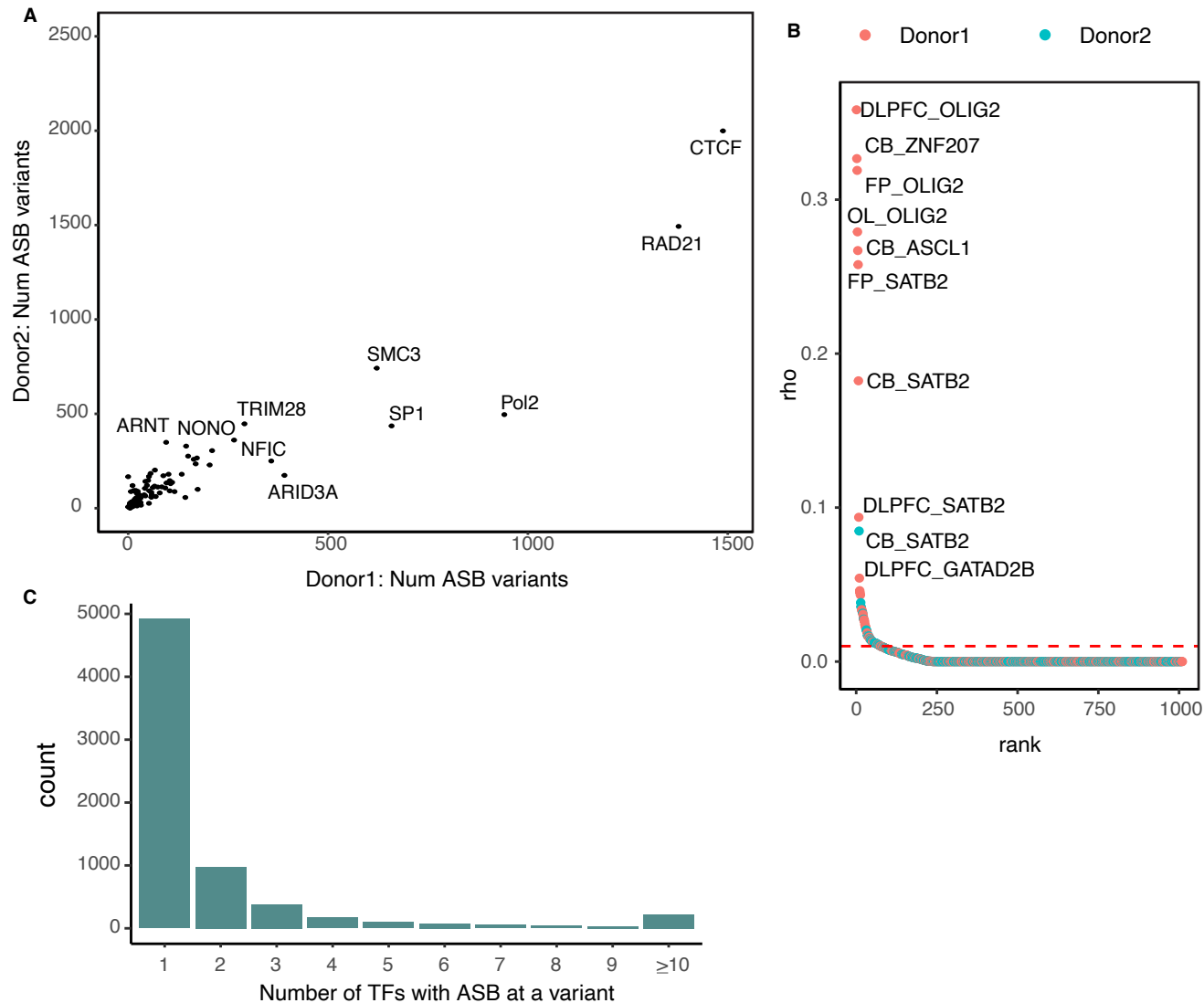




Supplemental Figure 2. Supplemental to Figure 1, part 2. A. The number of total ASB variants identified for each TF by donor. **B.** Rank plot of the ρ estimates of overdispersion from the beta-binomial model for each ChIP-seq dataset. **C.** Distribution of the number of TFs per ASB variant that exhibited a change in binding.