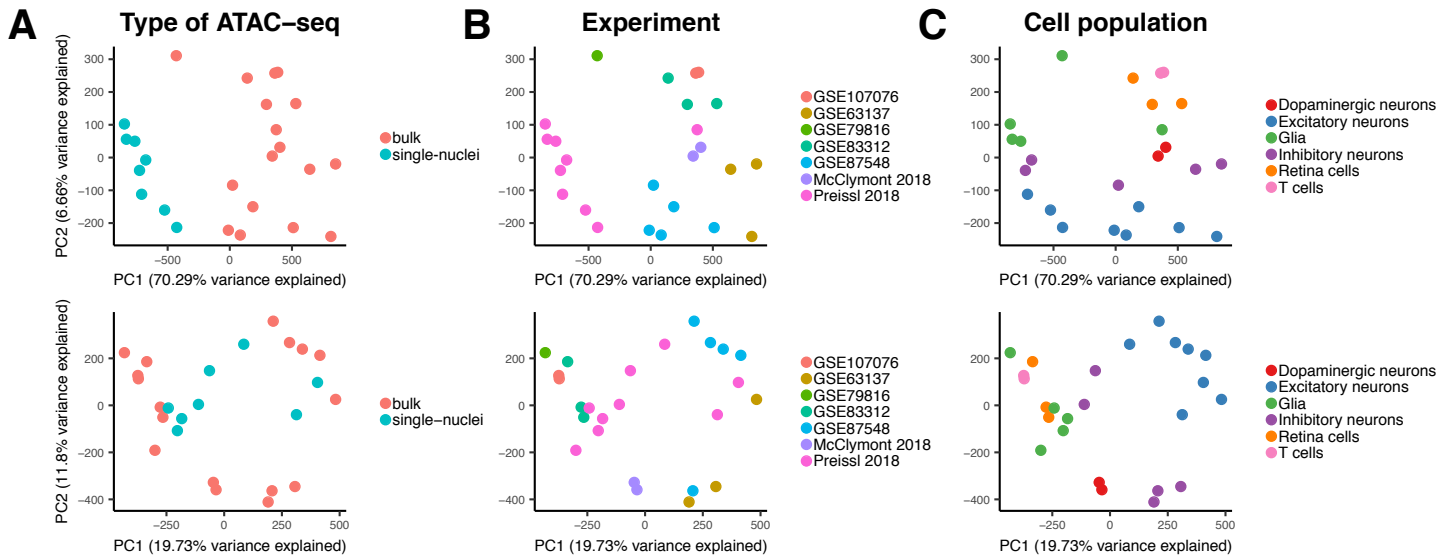


Supplemental Figure S1



Supplemental Figure S1. Summary of principal component analysis of ATAC-seq cell population peak read counts. A, B, C) PC1 vs. PC2 for log₂ (counts + 1) (top) and quantile normalized and batch corrected log₂ (counts + 1) (bottom). A) Cell populations are colored according to type of ATAC-seq performed (“bulk” or “single-nuclei”), B) by experiment from which they came, and C) by broad cell population category. All meta-information can be found in Supplementary Table S1.