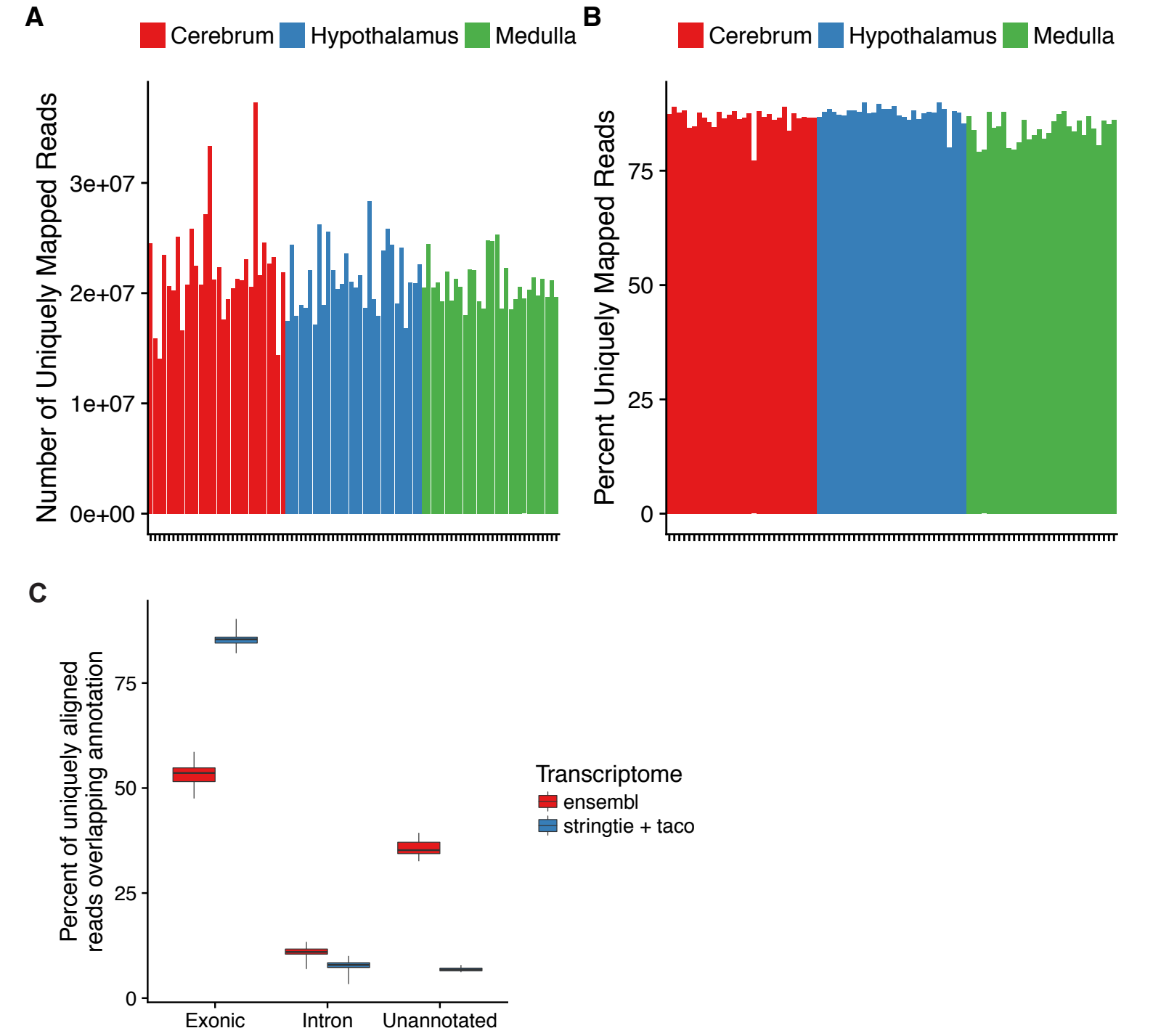
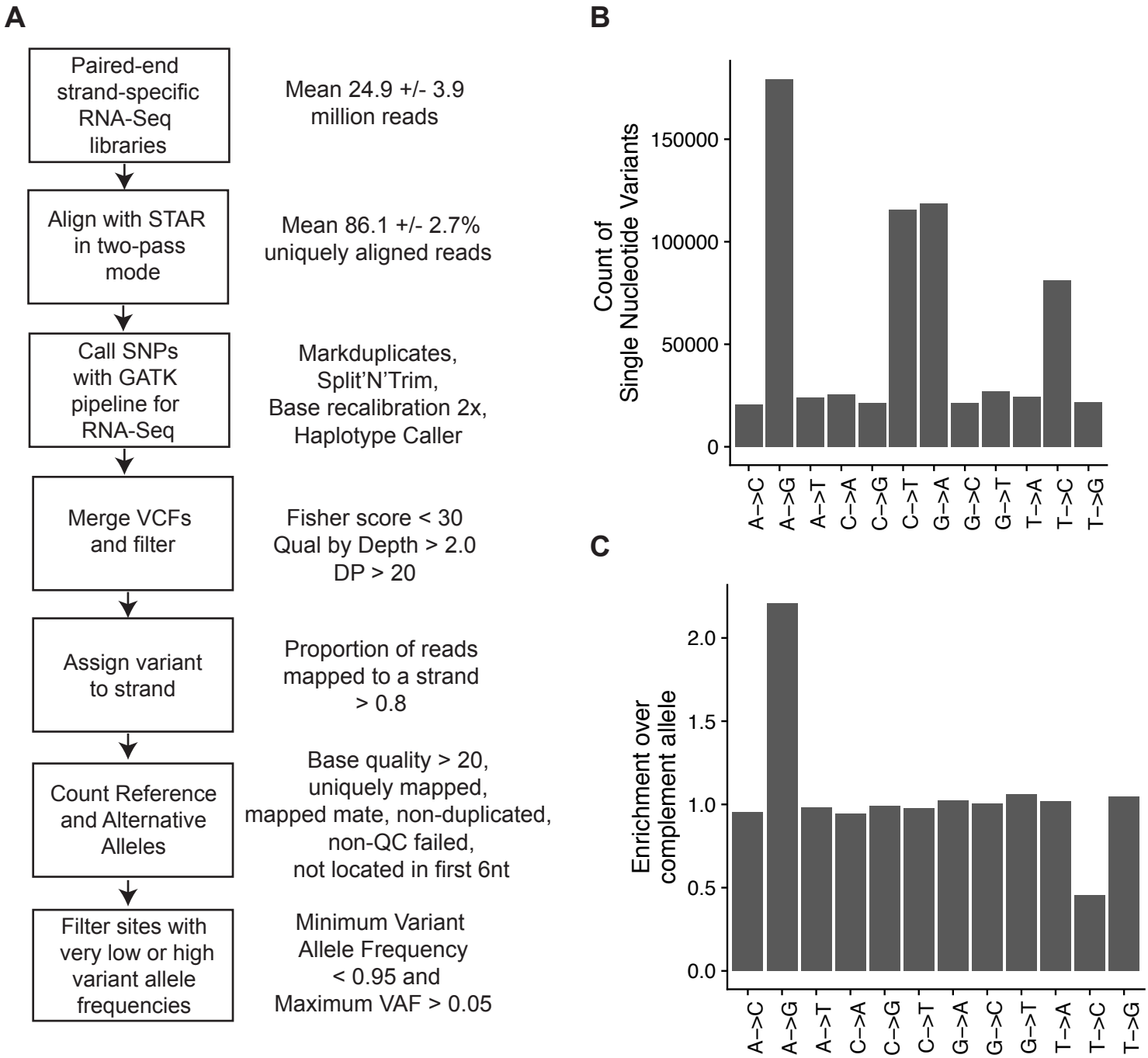
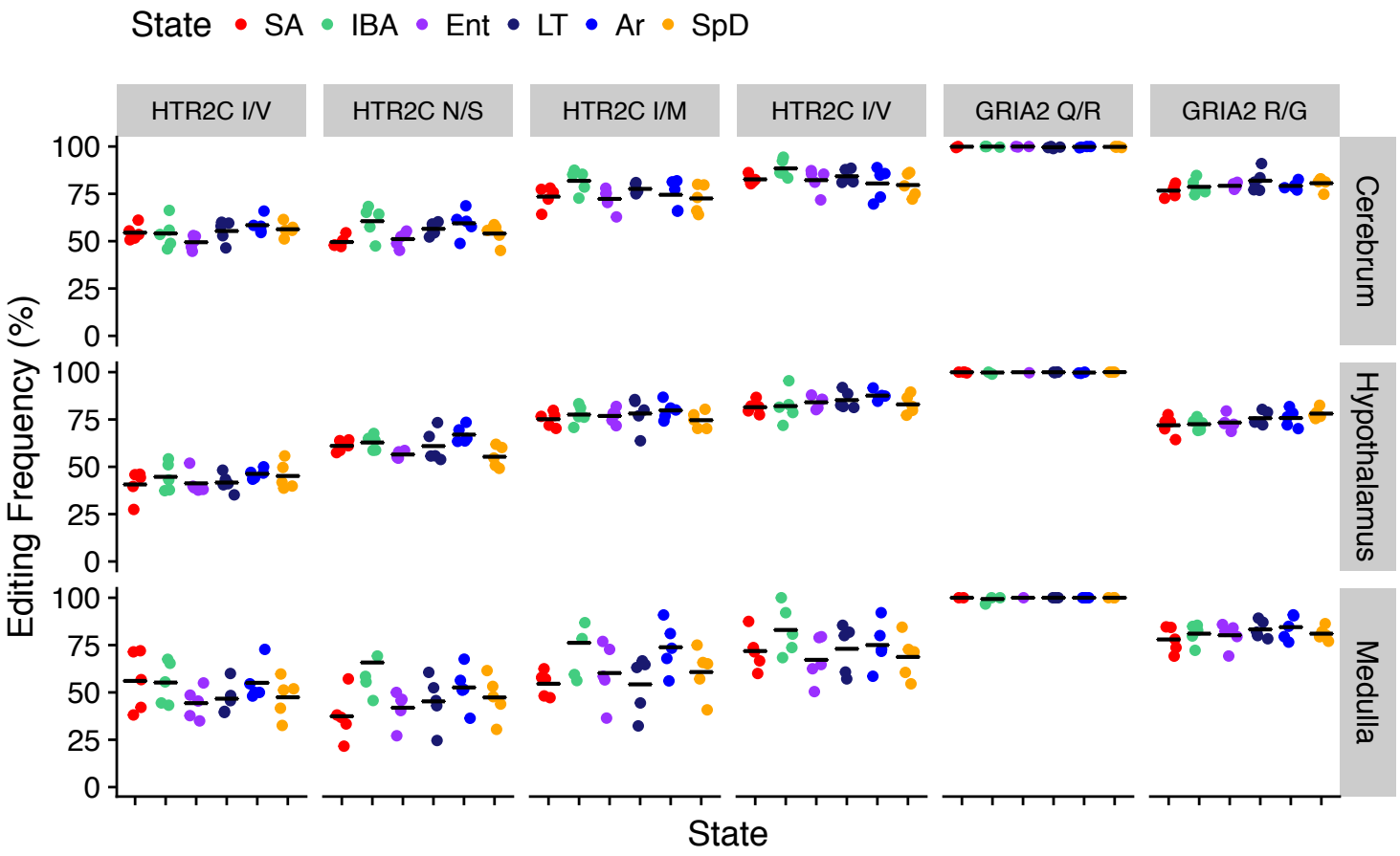
Supplemental Figure 1



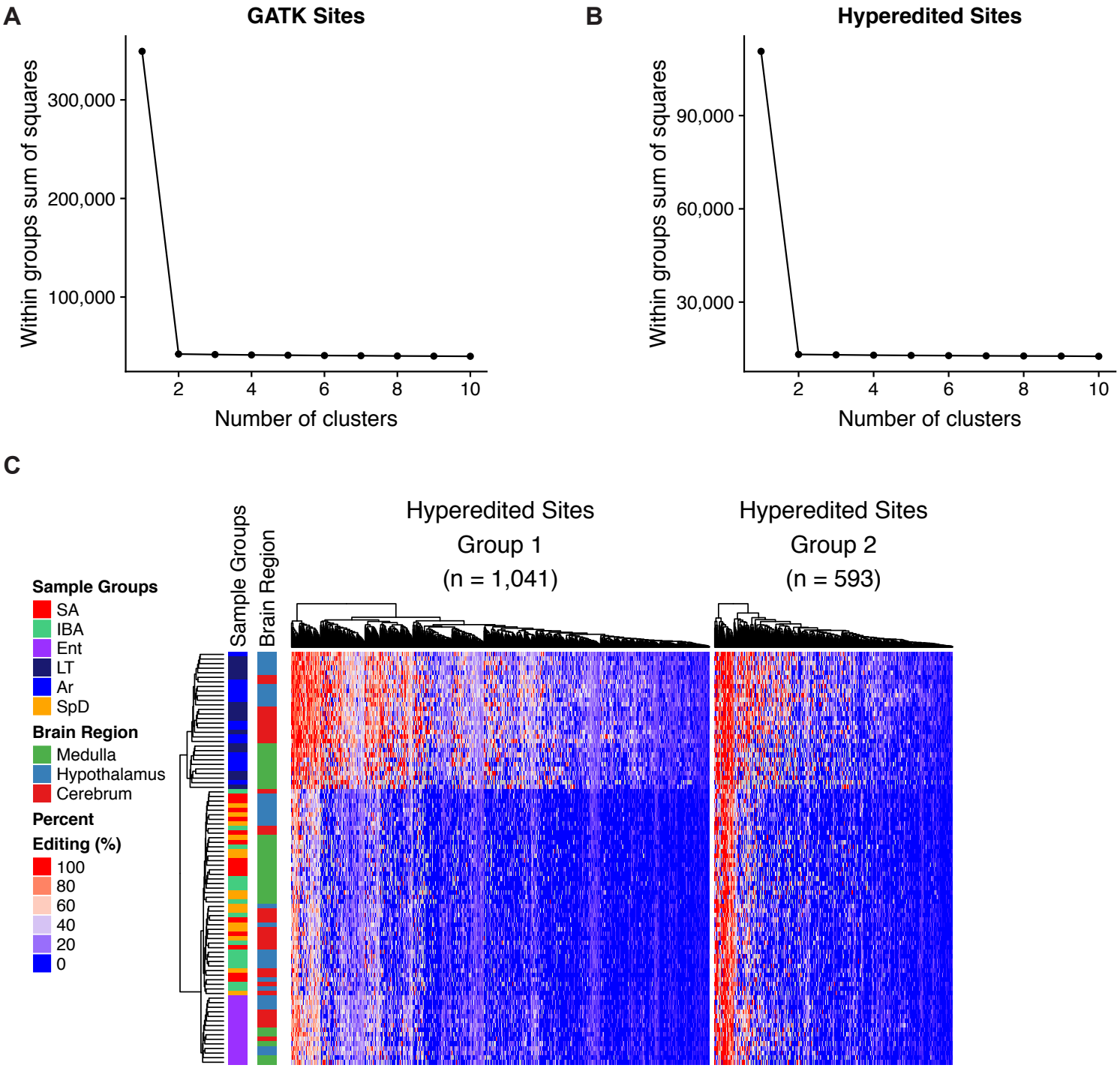
Supplemental Figure 2



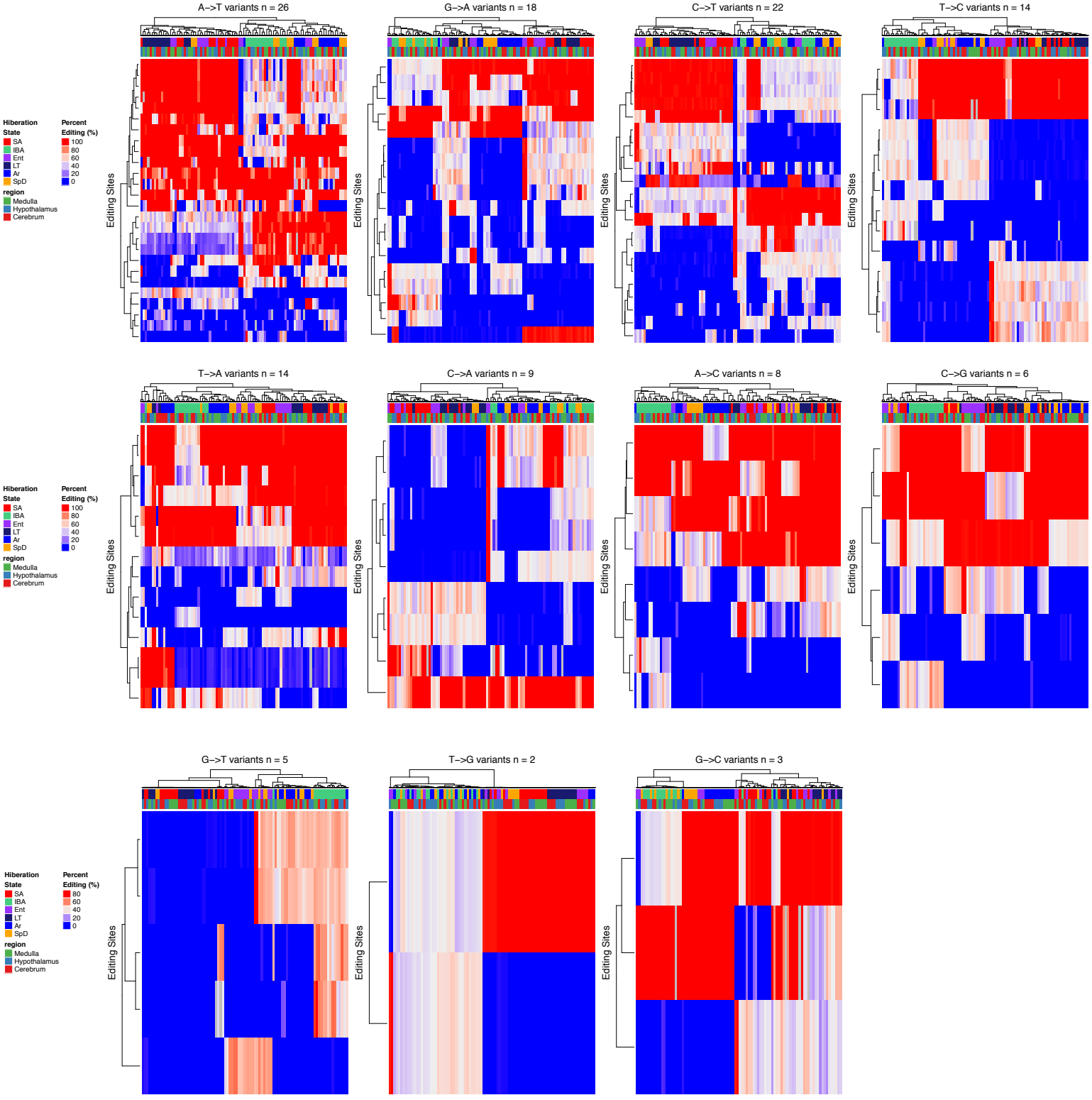
Supplemental Figure 3



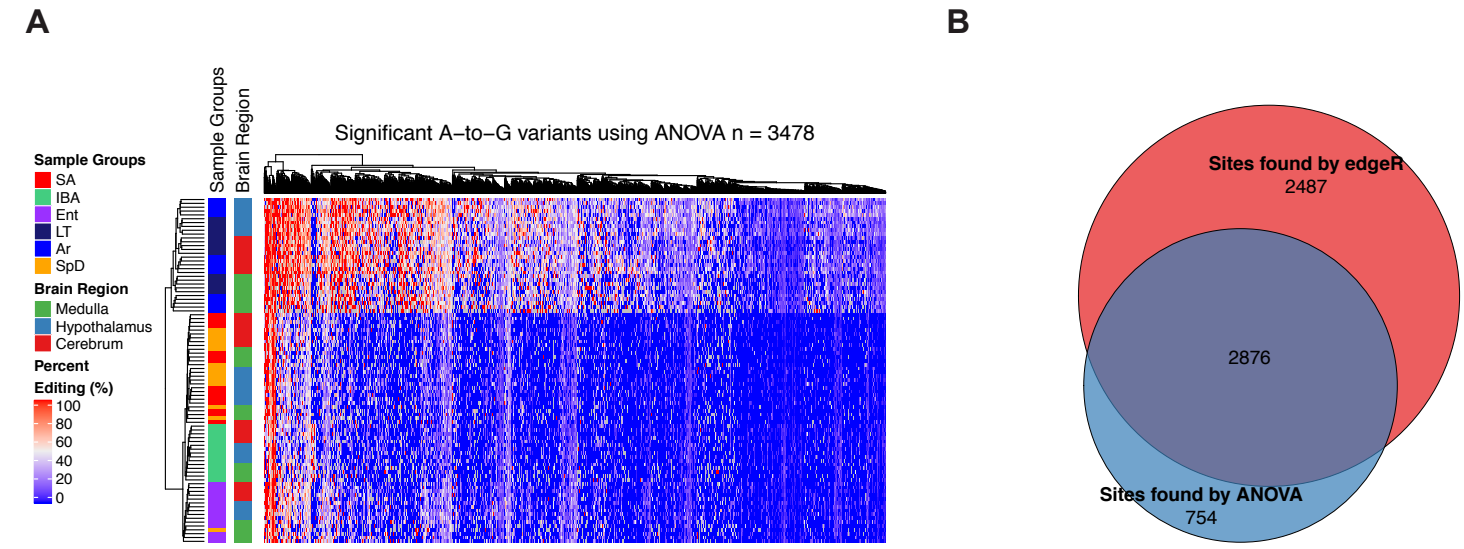
Supplemental Figure 4



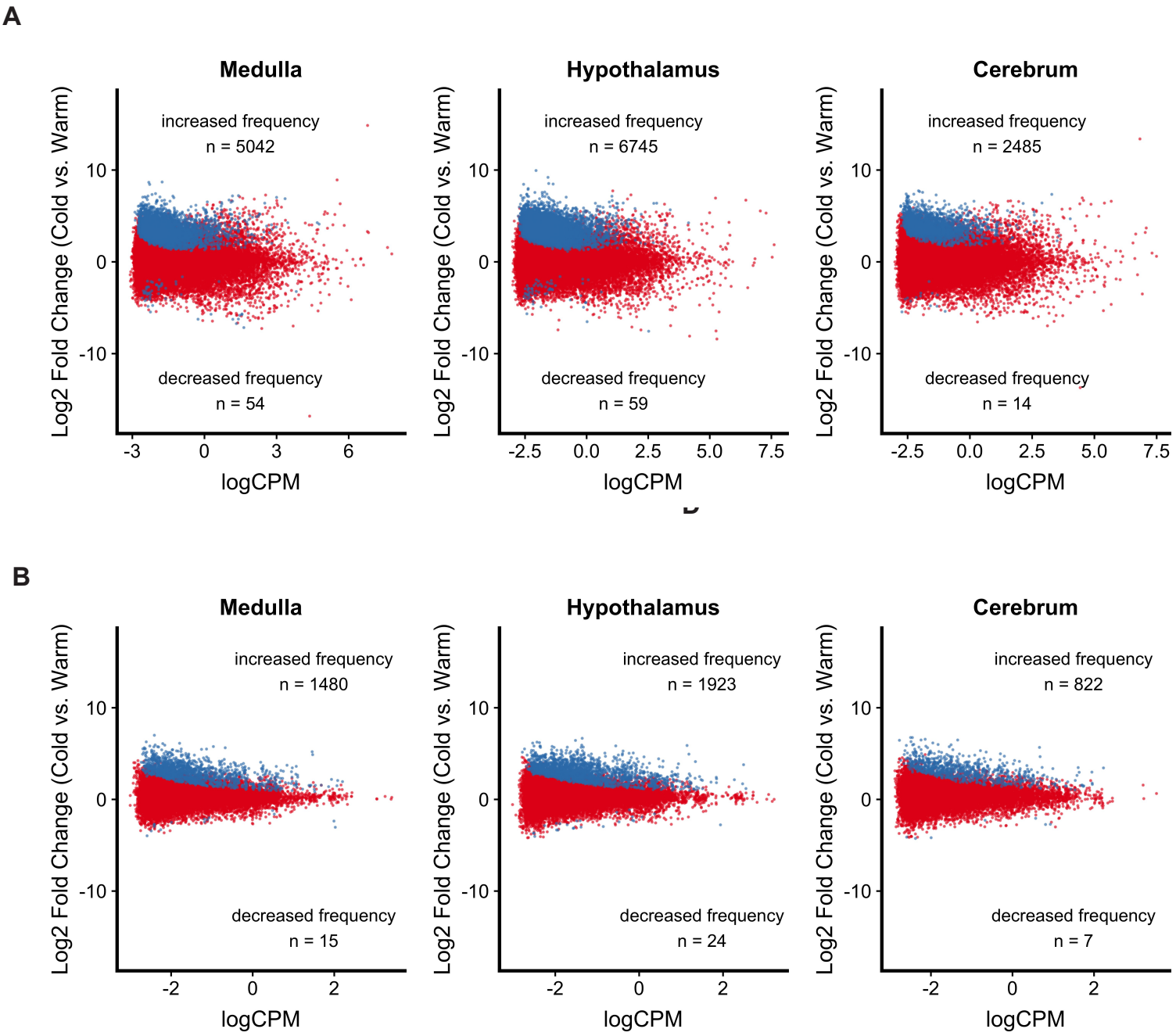
Supplemental Figure 5



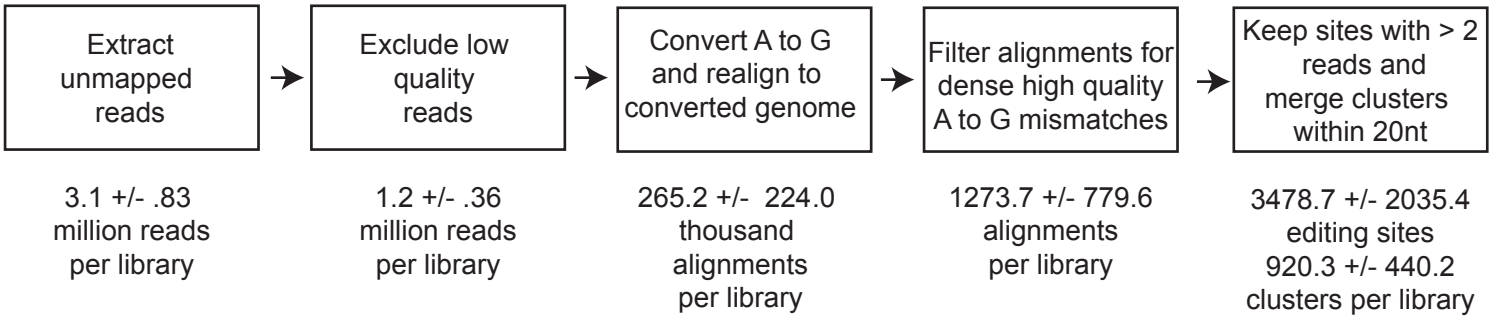
Supplemental Figure 6



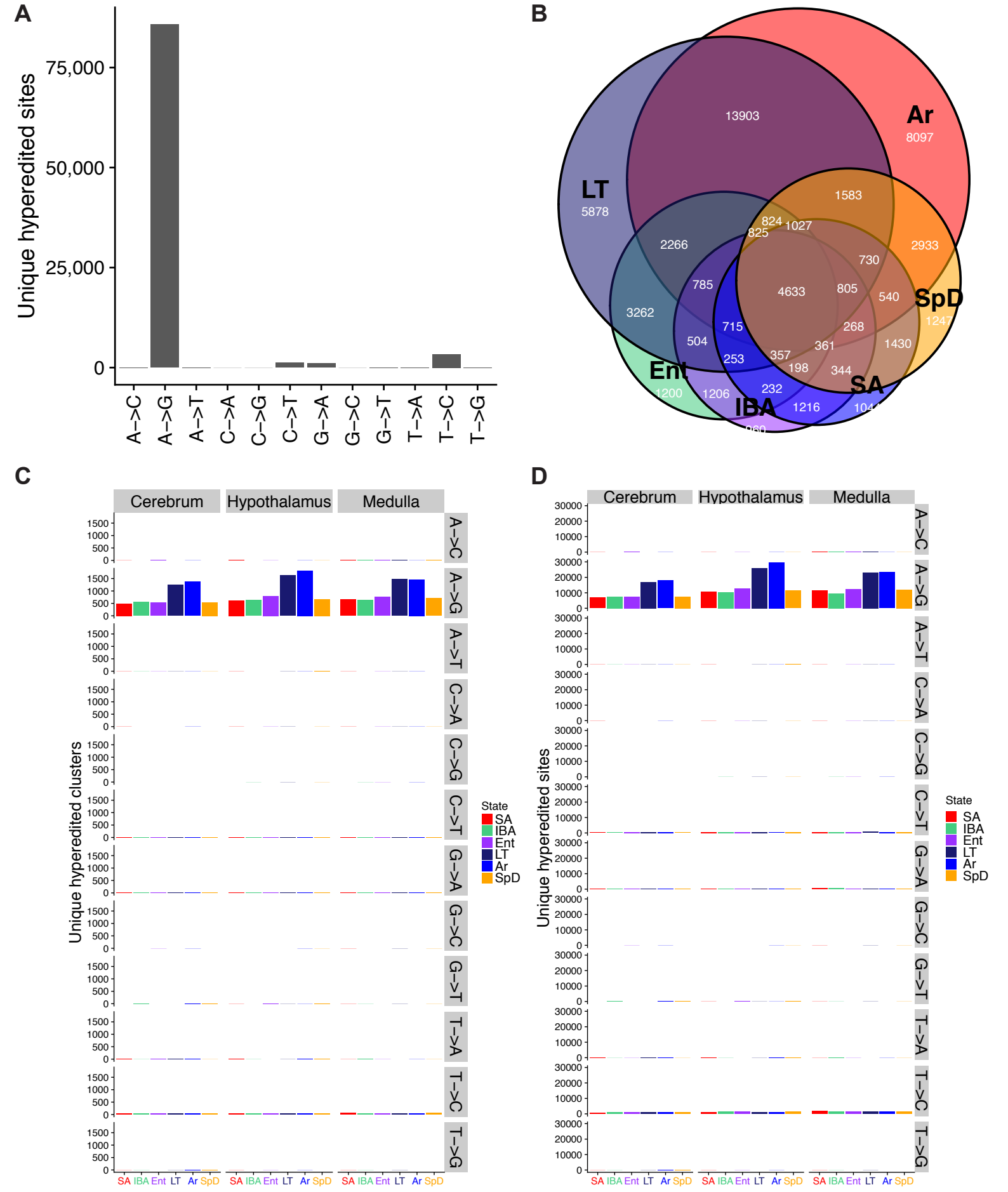
Supplemental Figure 7



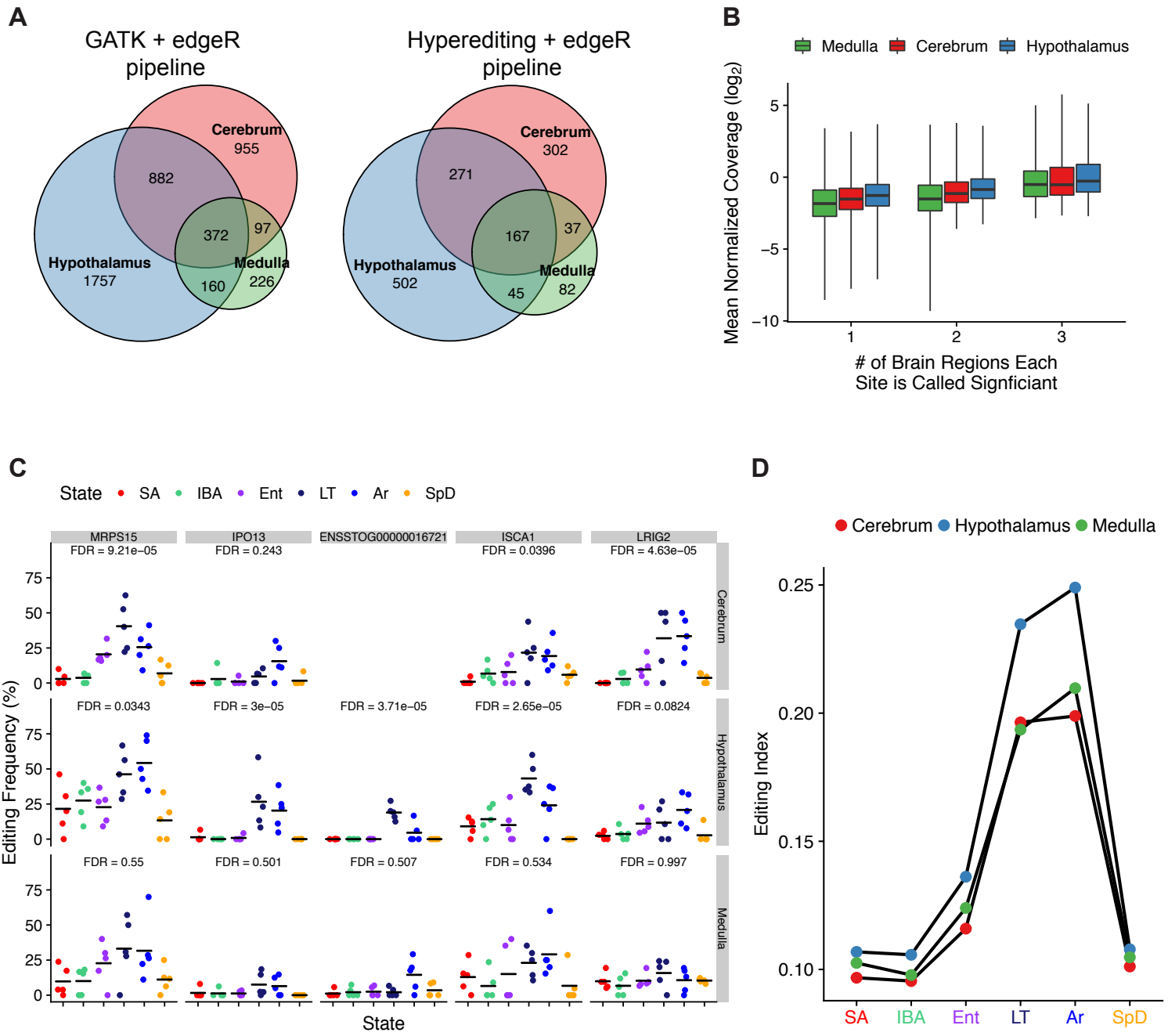
Supplemental Figure 8



Supplemental Figure 9

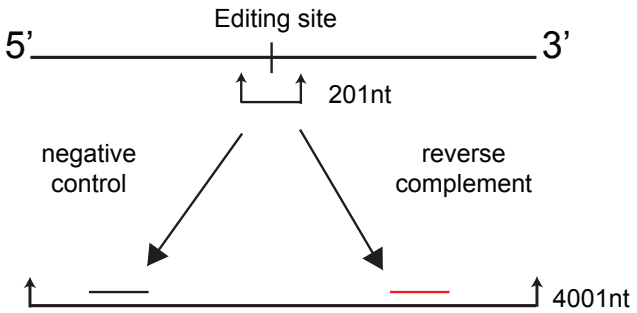


Supplemental Figure 10



Supplemental Figure 11

A

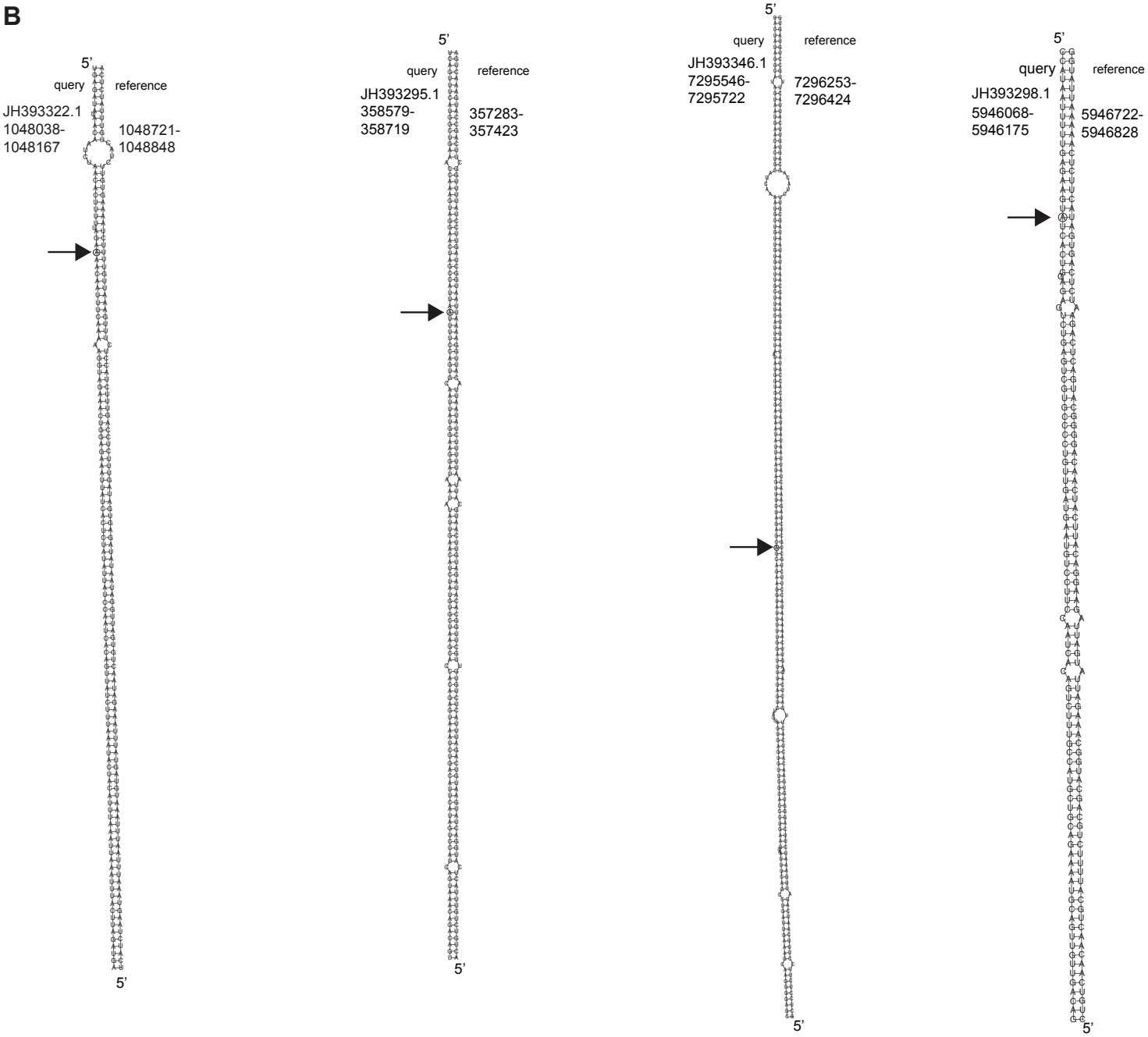


Reverse complement flanking sequencing and align to 4001nt region

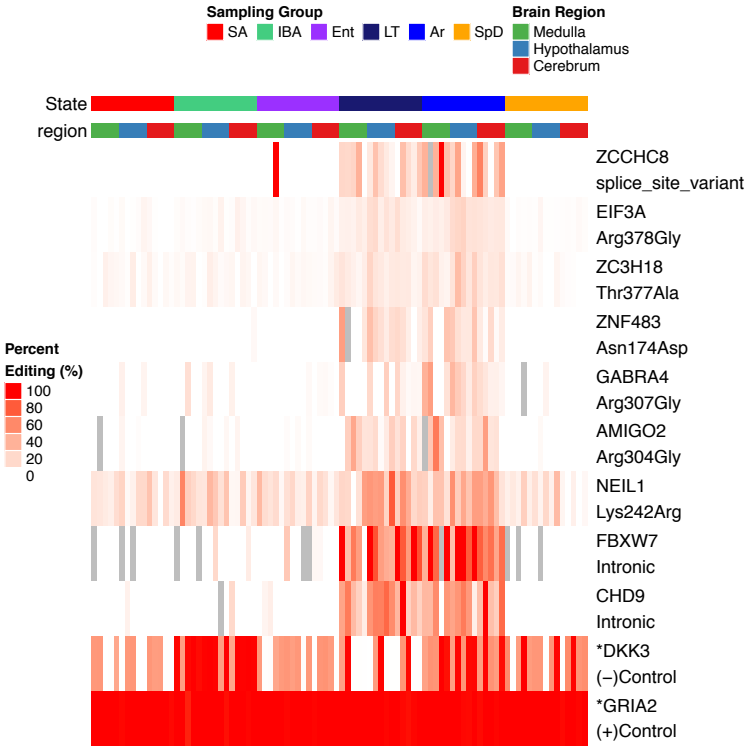
Require an alignment with an e-value < 0.01 that spans the editing site at least 20 nucleotides in length

As a control also align non-reverse complemented flanking region and examine second best alignments

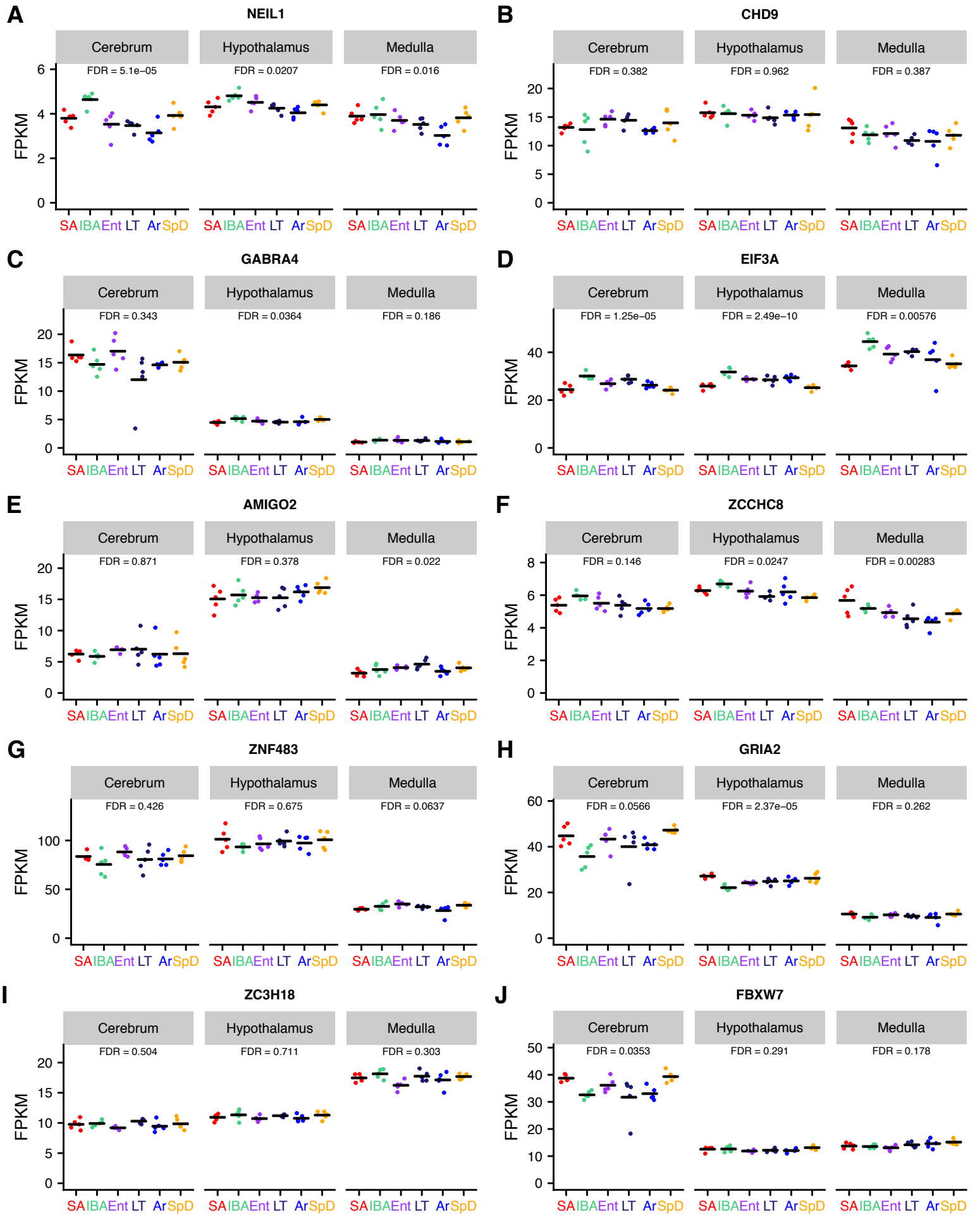
B



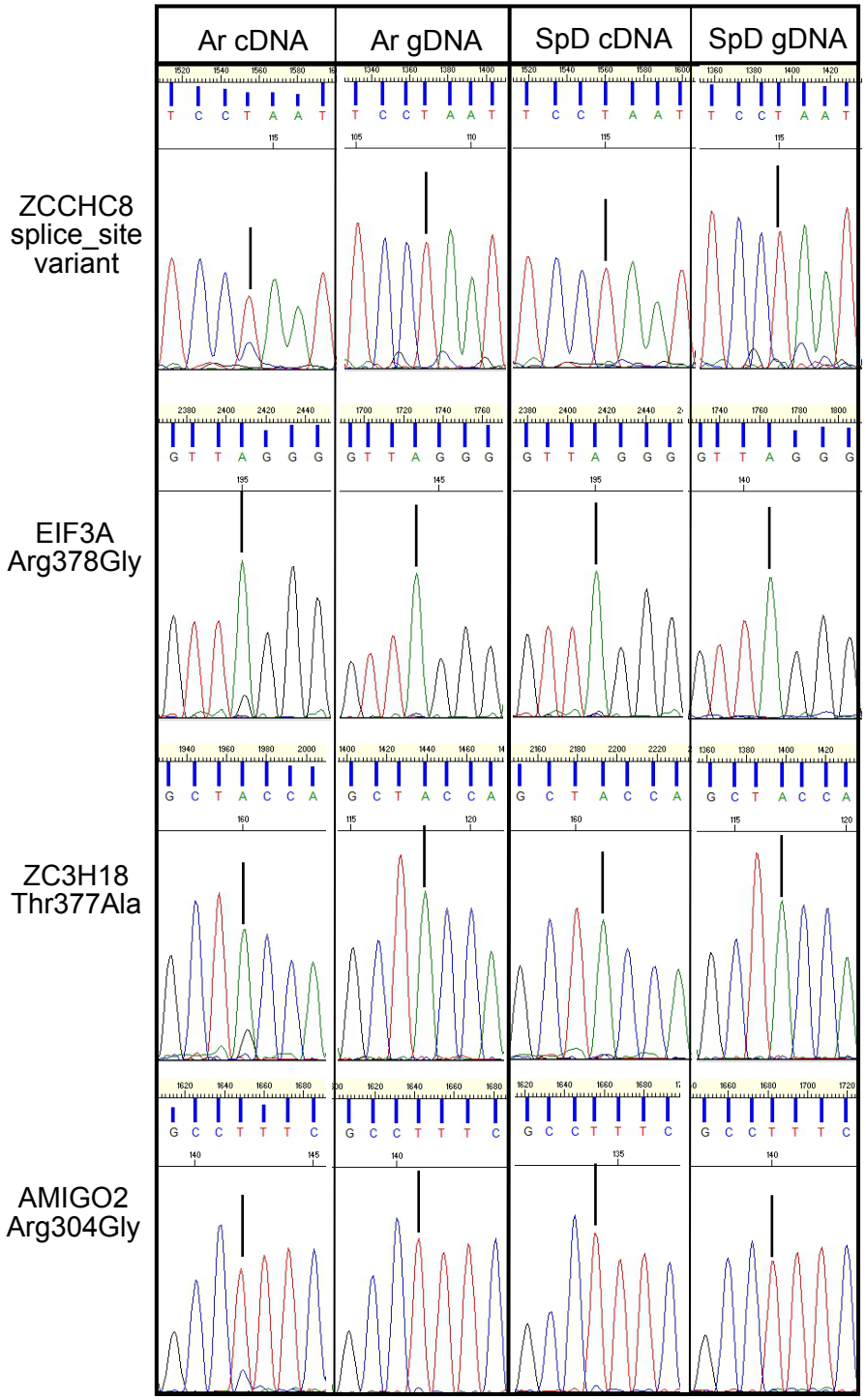
Supplemental Figure 12

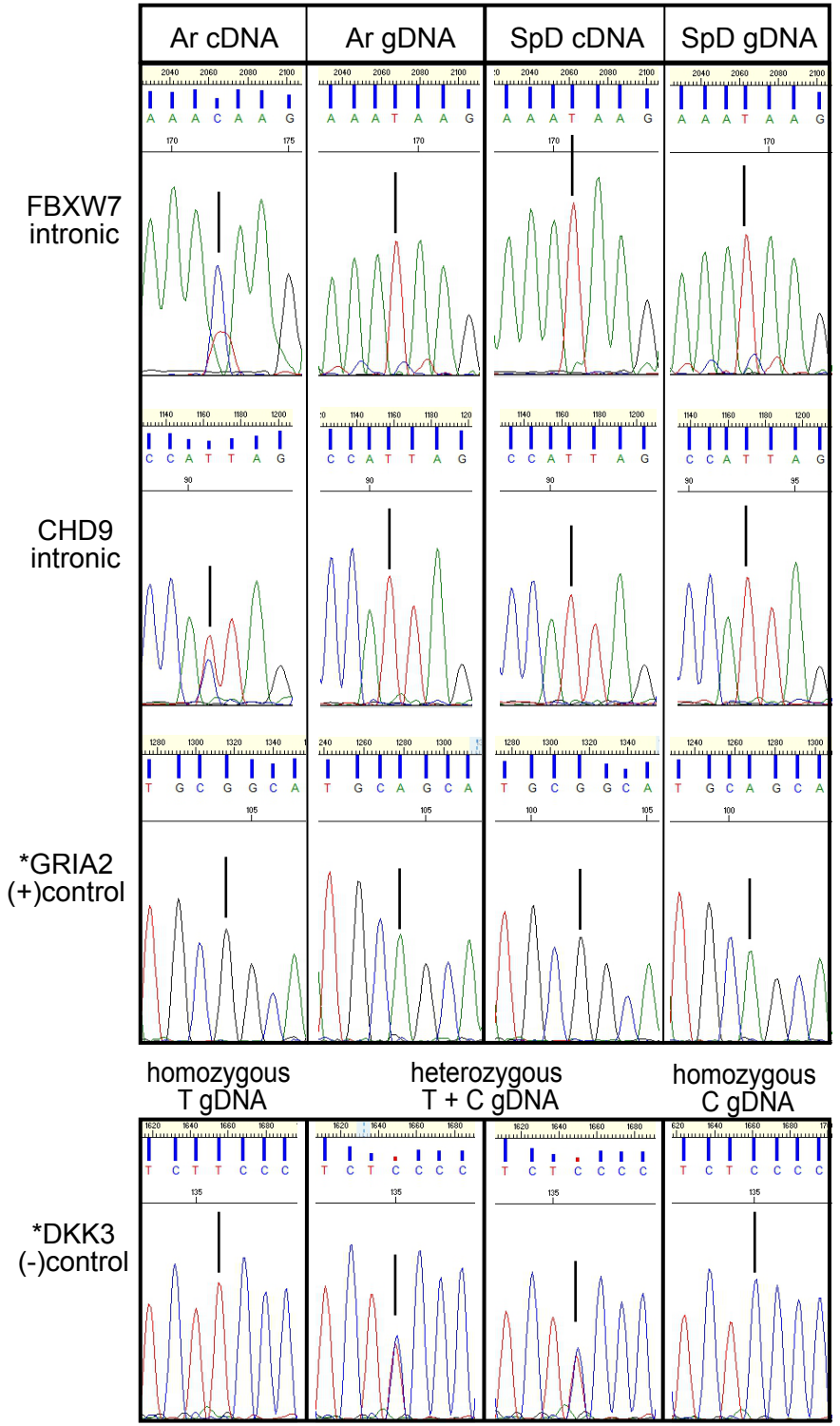


Supplemental Figure 13

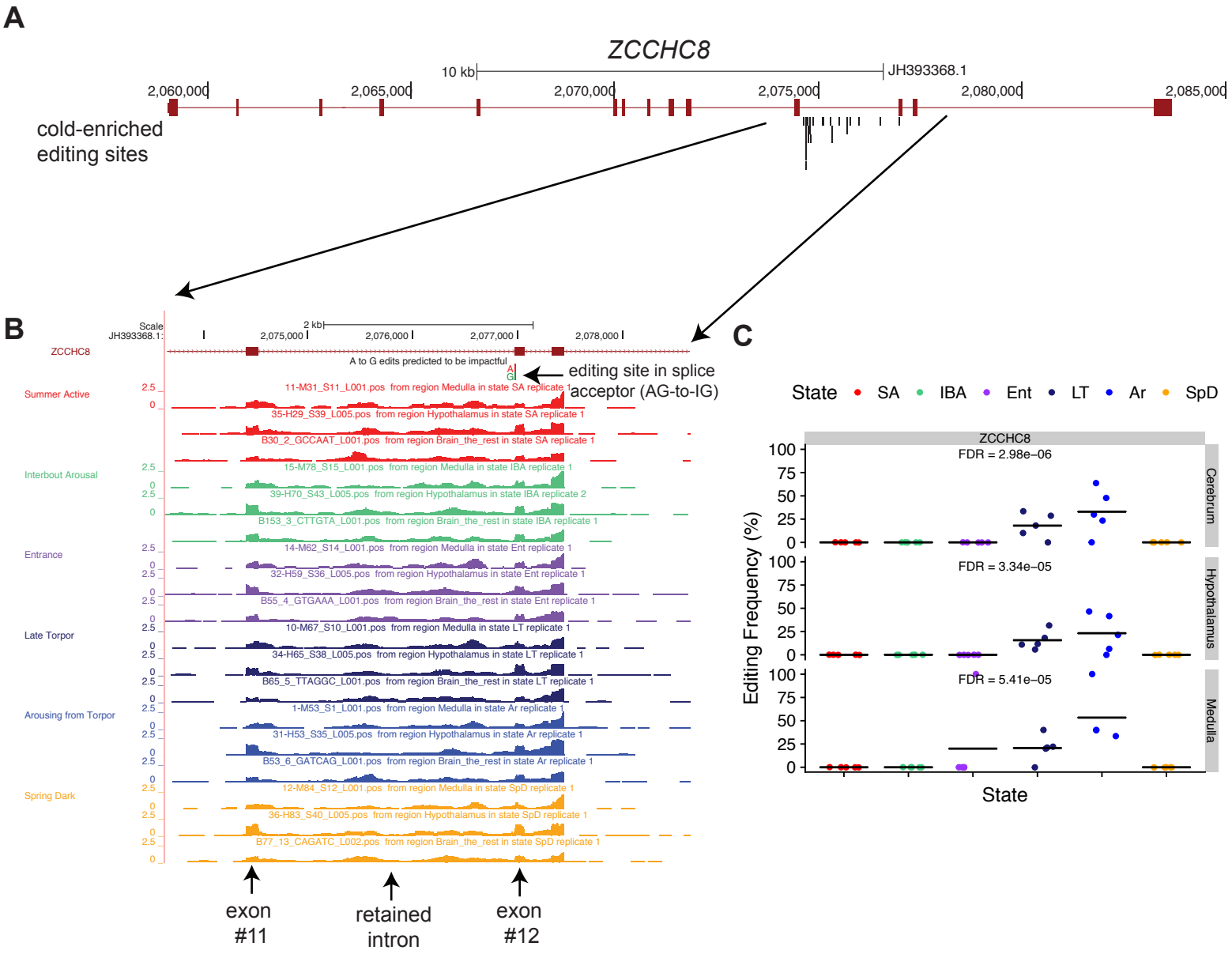


Supplemental Figure 14

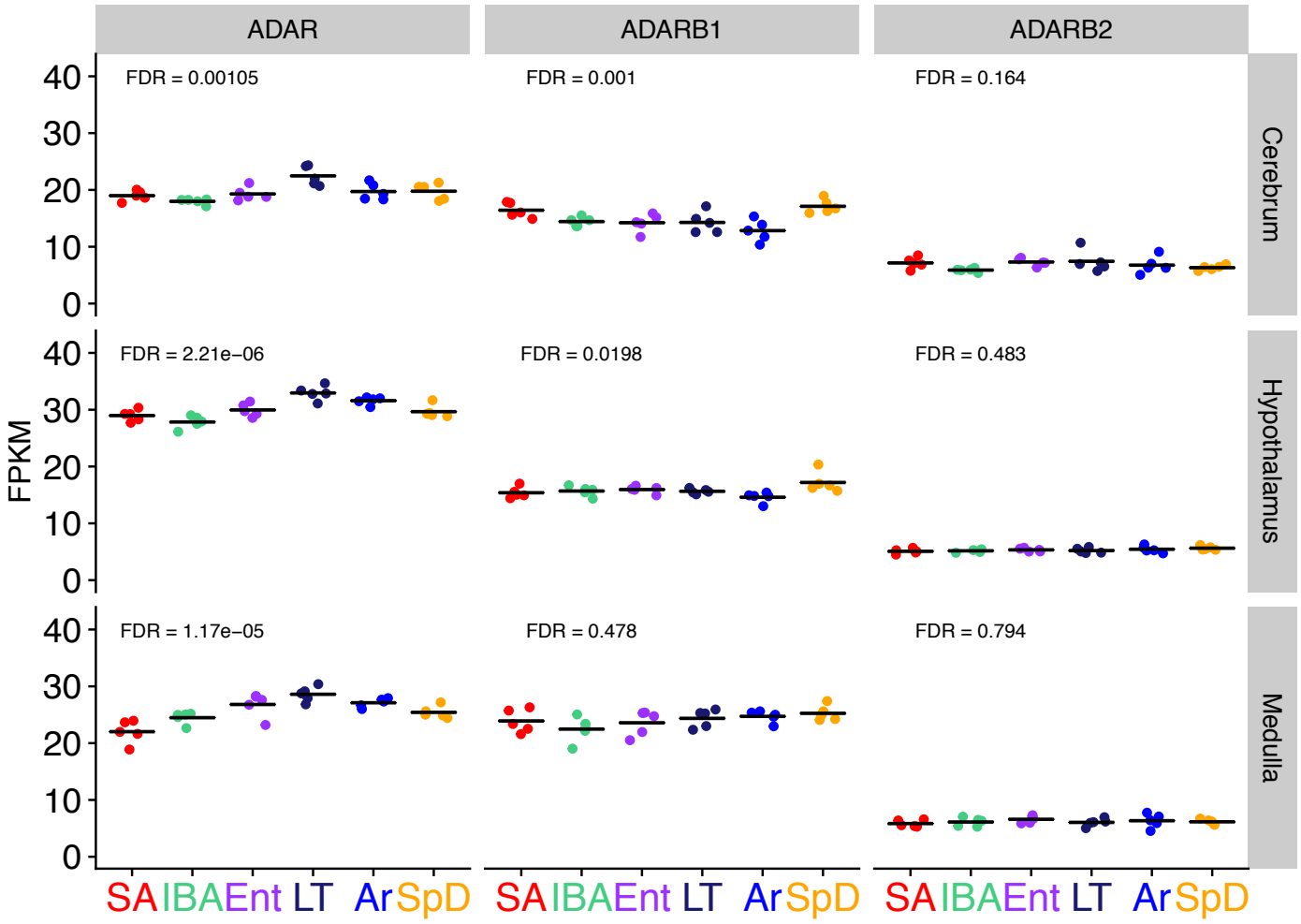
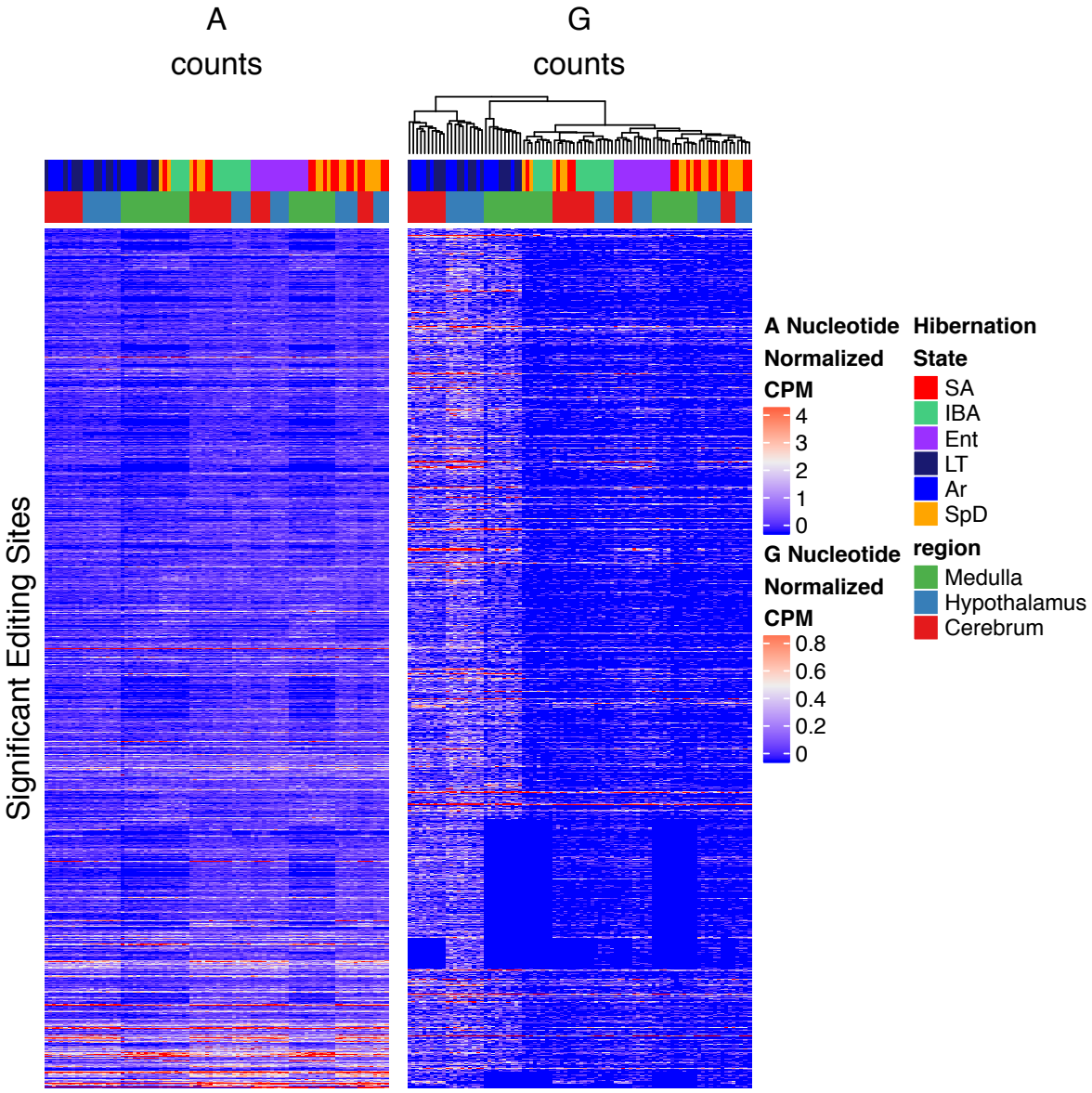




Supplemental Figure 15



Supplemental Figure 16



Supplemental Figure 17