Figure S10. A schematic flow of the CAFE pipeline

Shown is the schematic flow of the CAFE pipeline according to the combined, strand-specific support and strand-specific assembly. If there are both stranded and unstranded reads in the same cell type, the MAXIM, COCOA, and BEX steps are all executed. If there are only unstranded reads, the MAXIM step is carried out with the pooled stranded RNA-seq data.