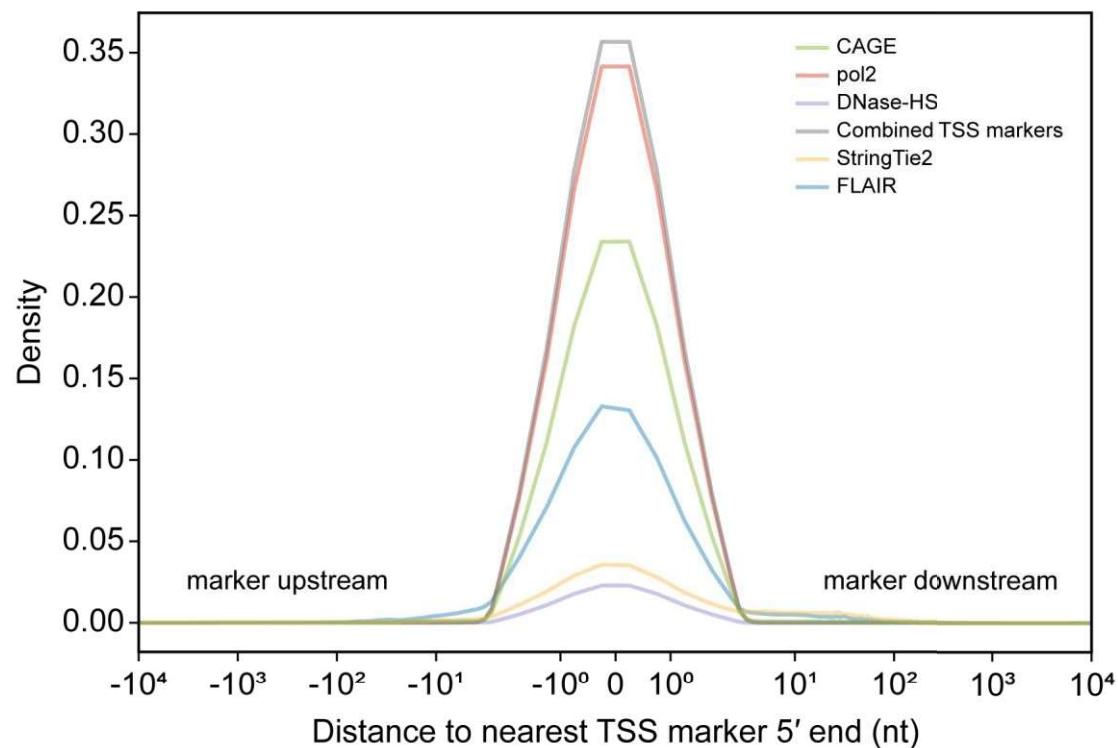




Supplementary Figure 11 Nucleotide distance of orthogonal datasets from high-confidence scaffold 5' ends. The x axis is the number of nucleotides between a nanopore read 5' end (0 position) and the closest signal from the orthogonal dataset. Negative numbers are upstream (5') from the TSS; positive numbers are downstream (3') from the TSS. The y axis is the density at a given distance of the signal from the closest high-confidence scaffold 5' ends.