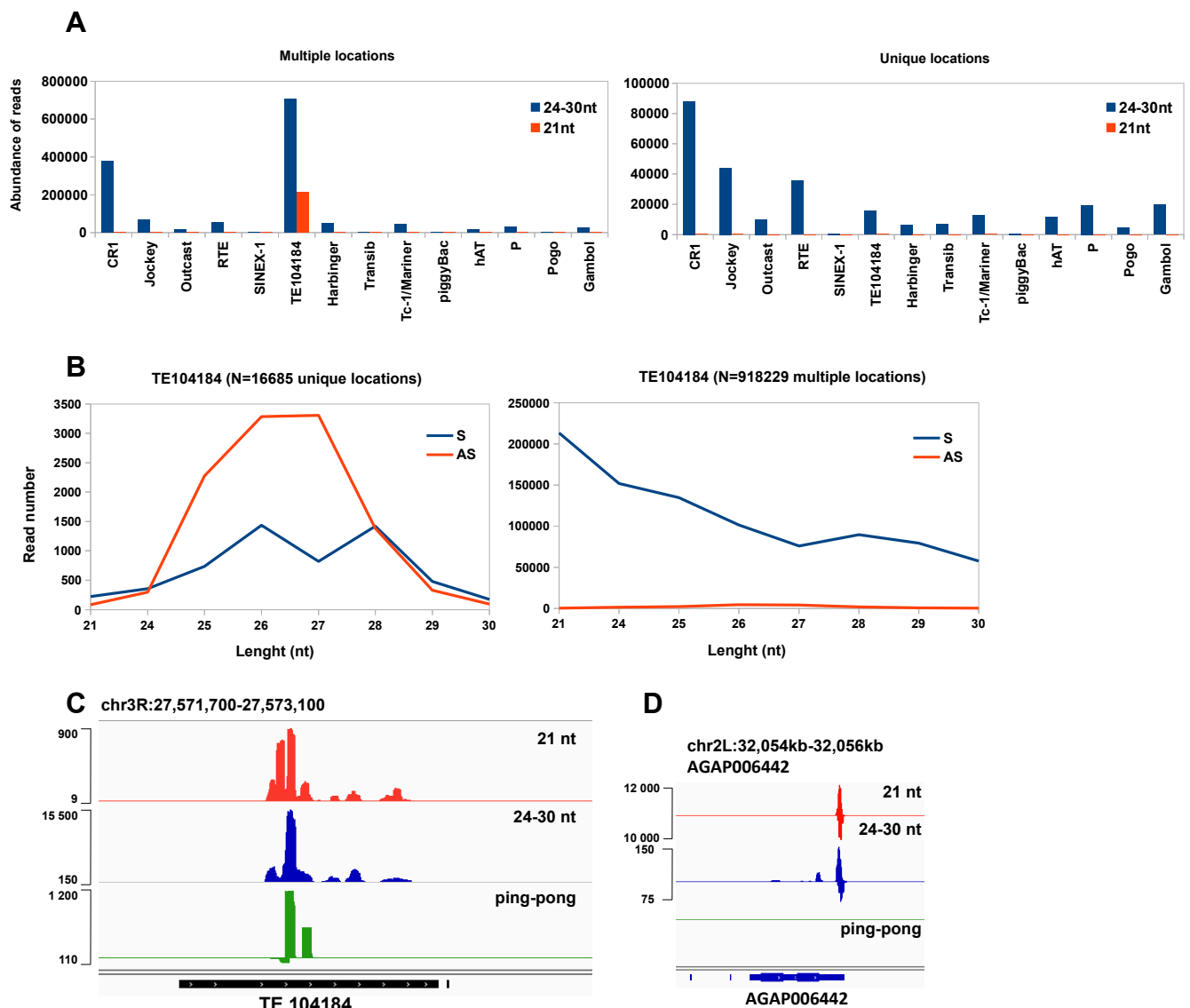




Additional file 4. Figure S1

(A) Read frequency for siRNAs (21 nt) and piRNAs (24-30 nt) mapped to the most abundant NLTR and DNA transposons. (B) Read frequency and distribution mapped to the unknown *TE element 104184* (class II DNA transposons) uniquely mapped to *An. gambiae* genome (left) and multiply mapped reads to the genome (right). (C-D) Density and distribution of small RNAs mapped to *TE104184* transposon (C) and *AGAP006442* (D); siRNAs (red), piRNAs (blue) and piRNA ping-pong pairs (green) are as indicated.