



Figure S3. Sense and antisense transcription at TAD borders.

Average of the \log_{10} ratio between sense and antisense transcription for embryonic cells and neuronal cells at (A and D) conserved, (B and E) BG3-specific and (C and F) Kc167-specific TAD borders. (A-C) total mRNA levels (D-F) nascent RNA (GRO-seq in S2 and 3'NT-seq in BG3 as mentioned in method section).