

Start crop Point

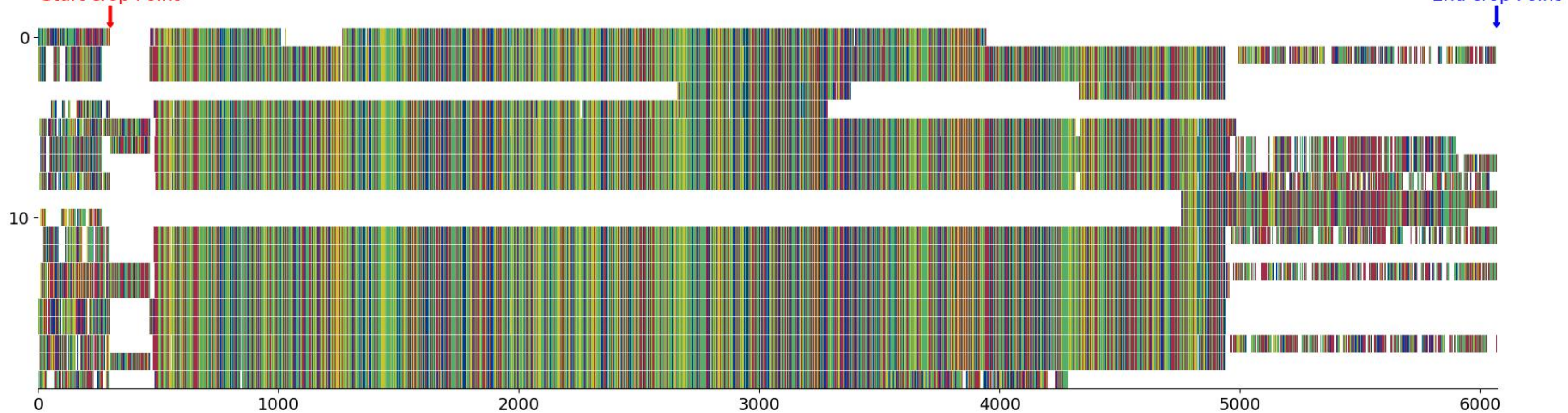
End crop Point

MSA length = 5769



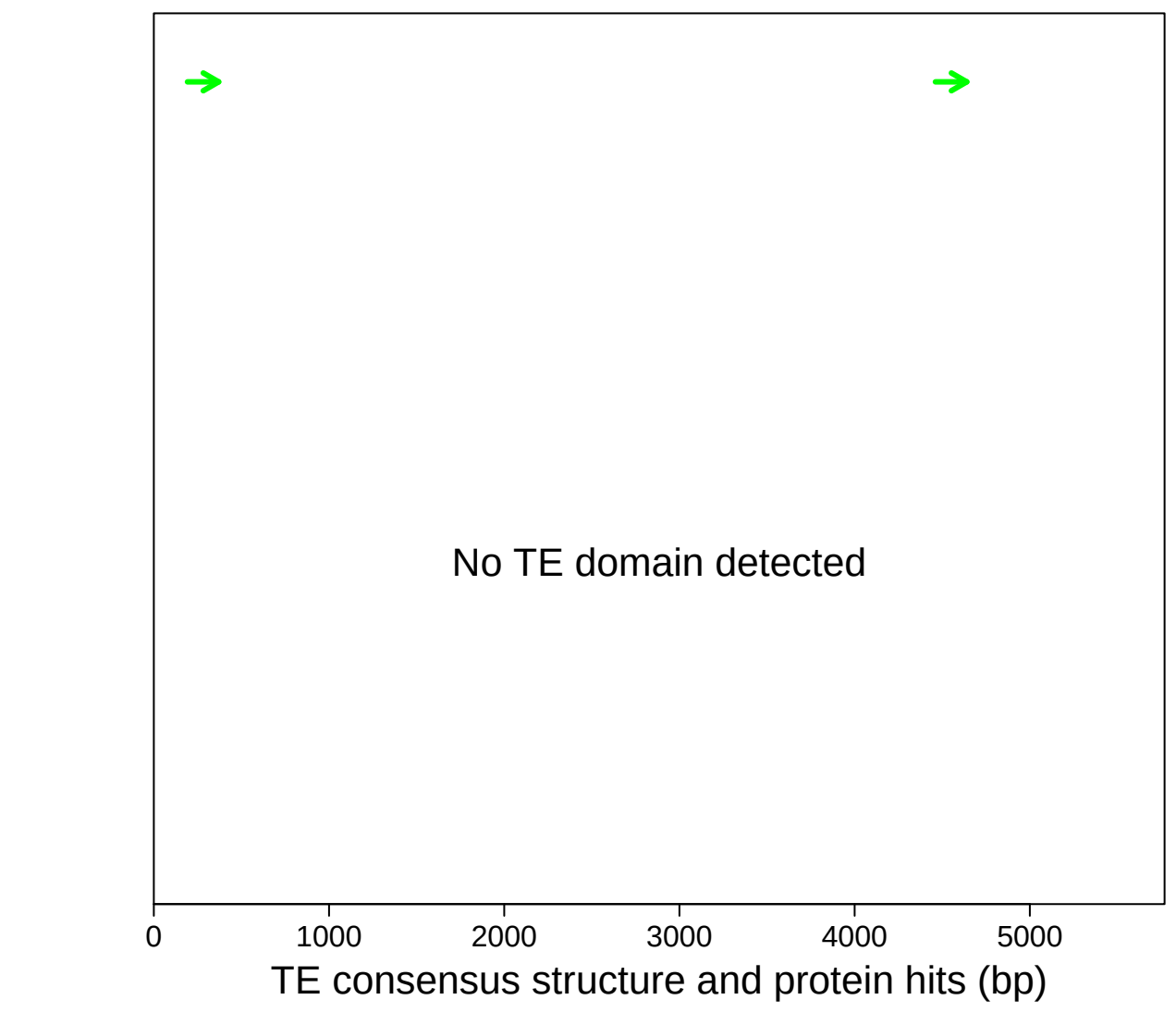
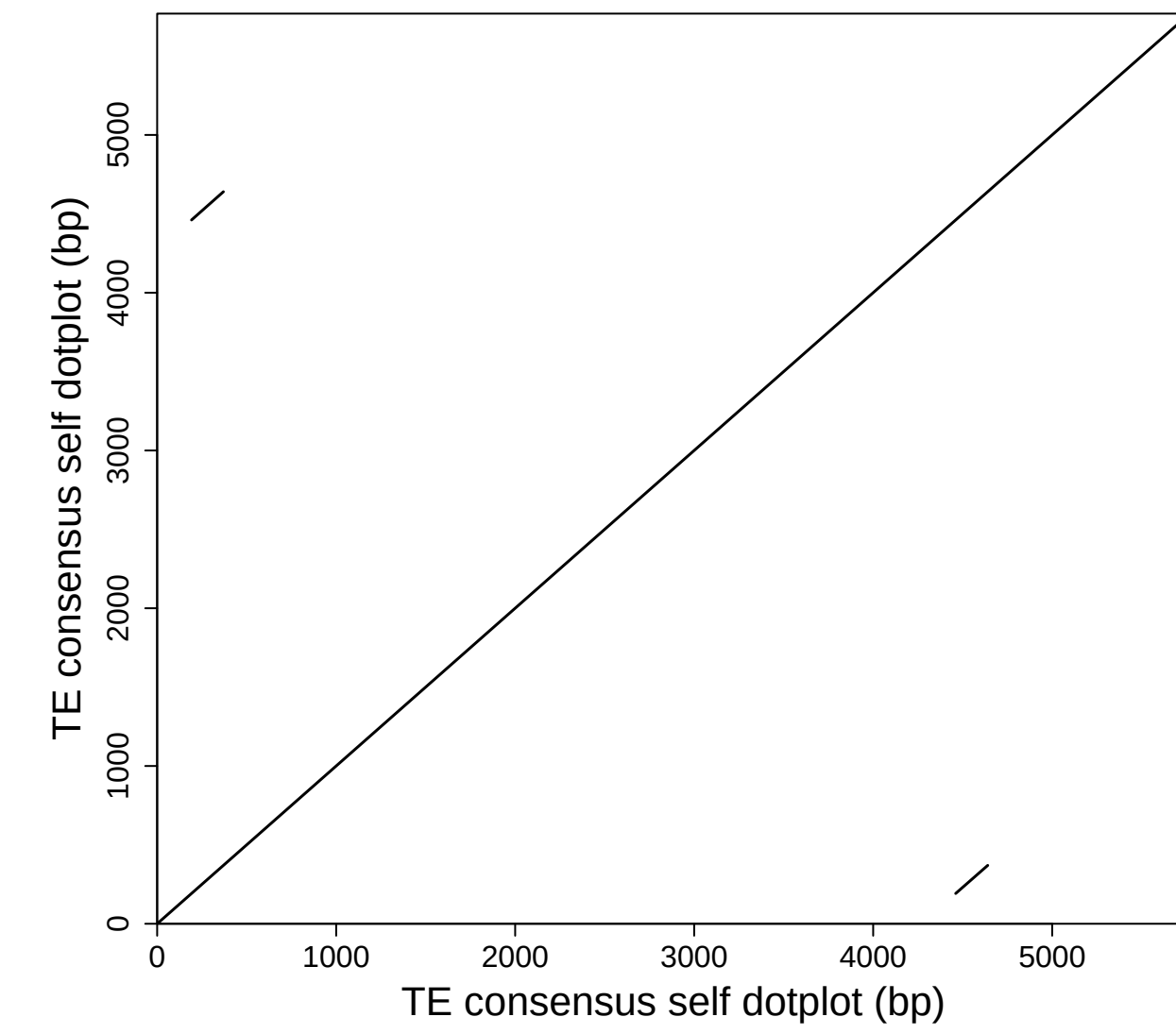
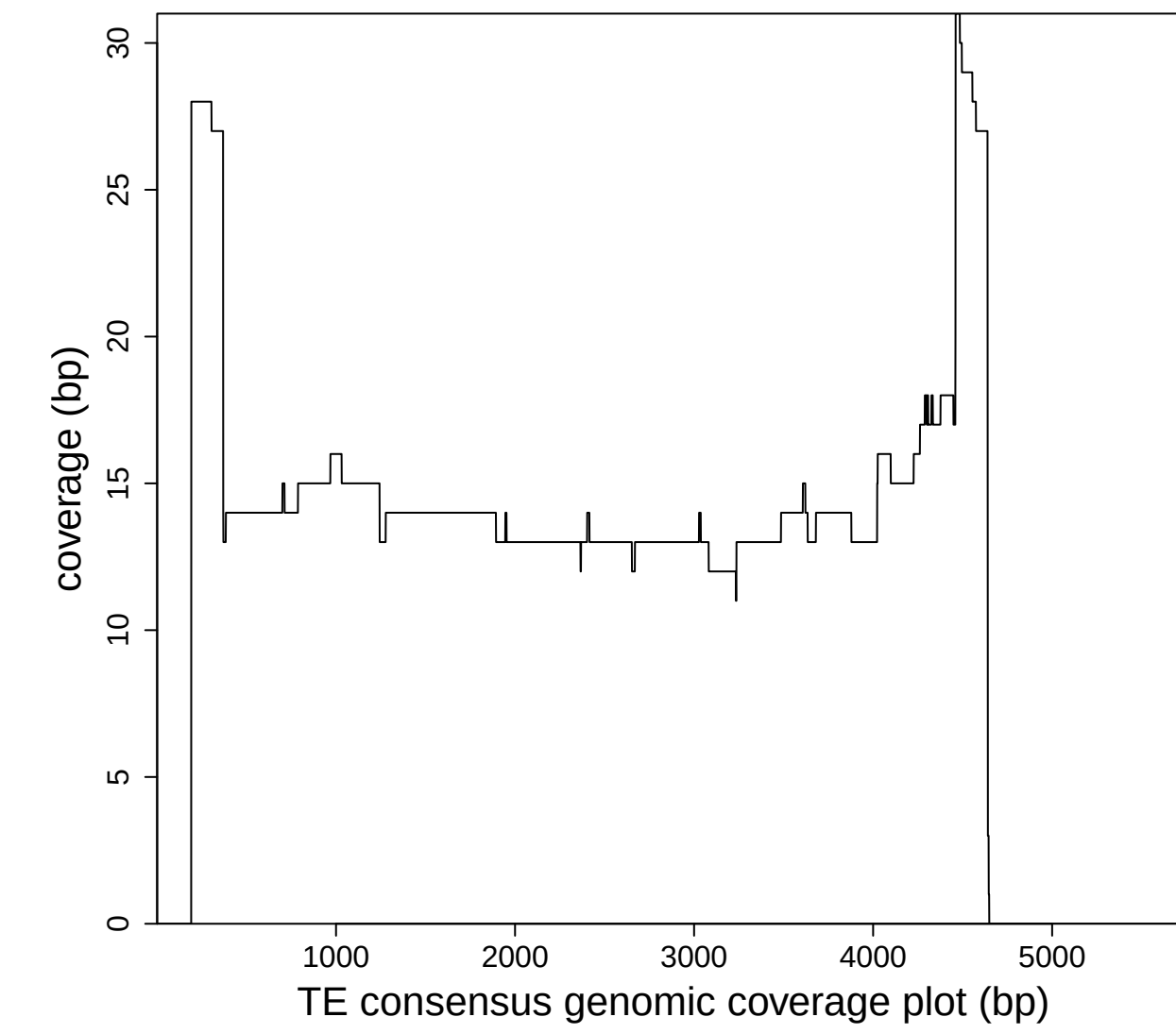
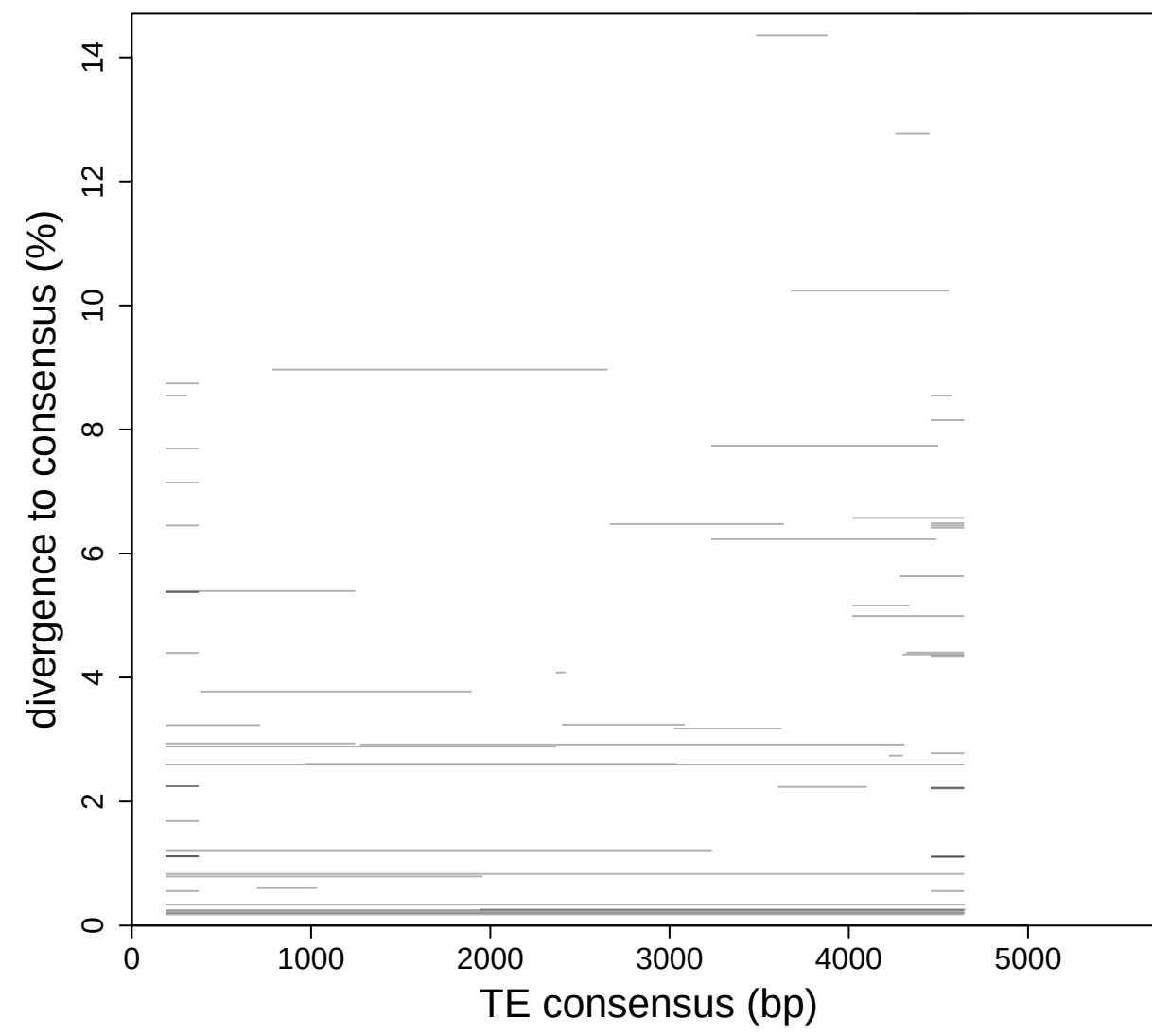
Start crop Point

End crop Point



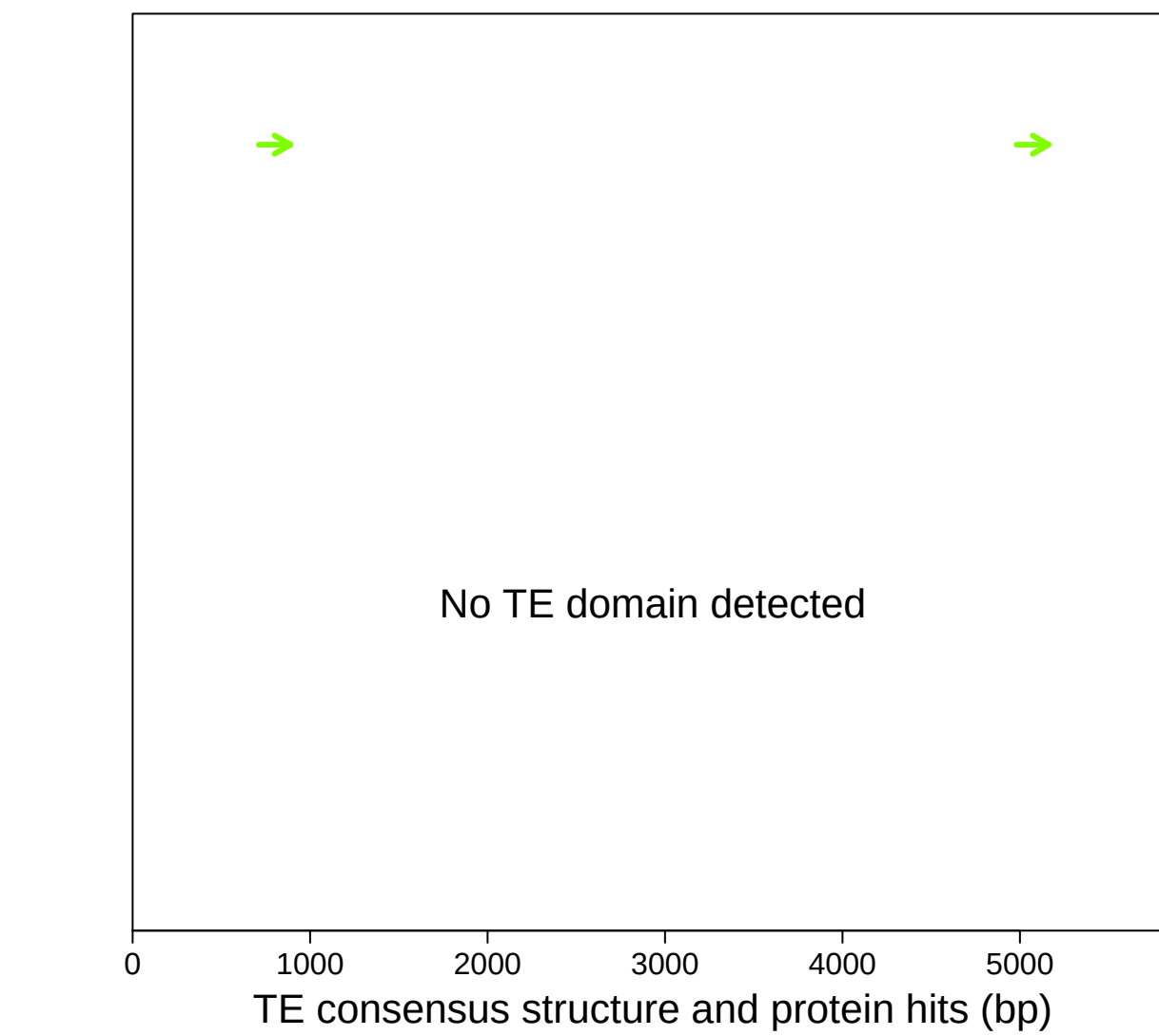
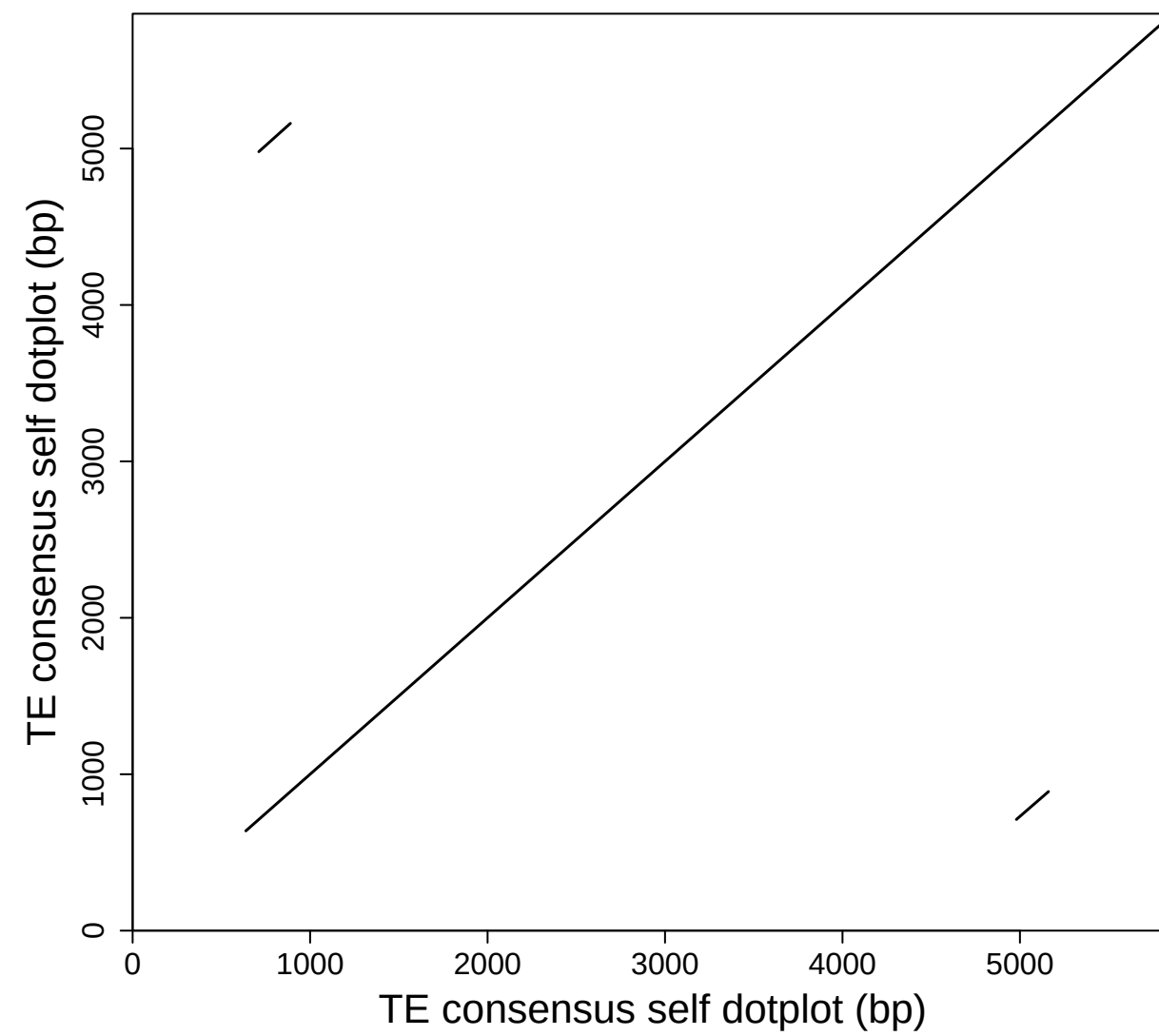
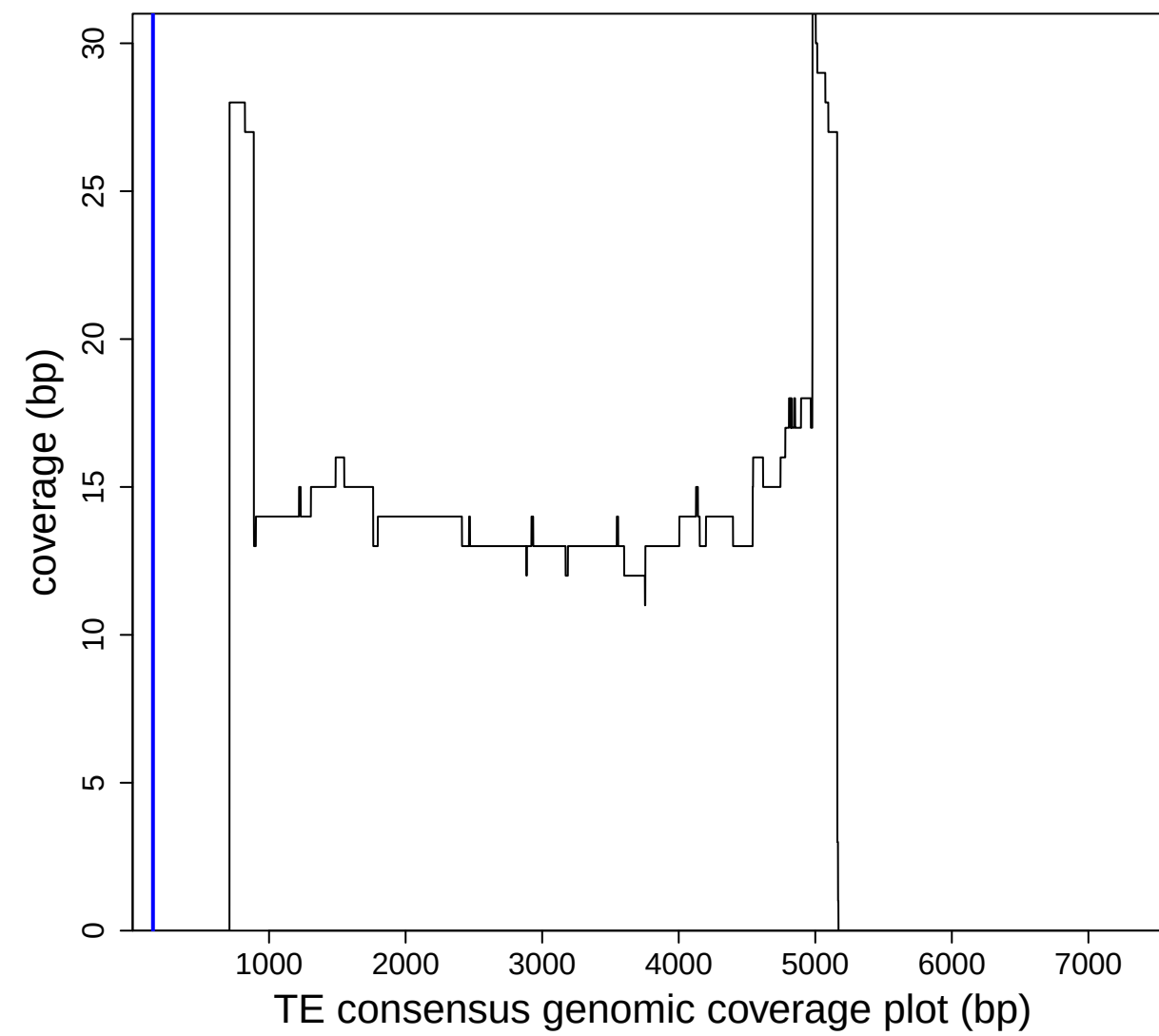
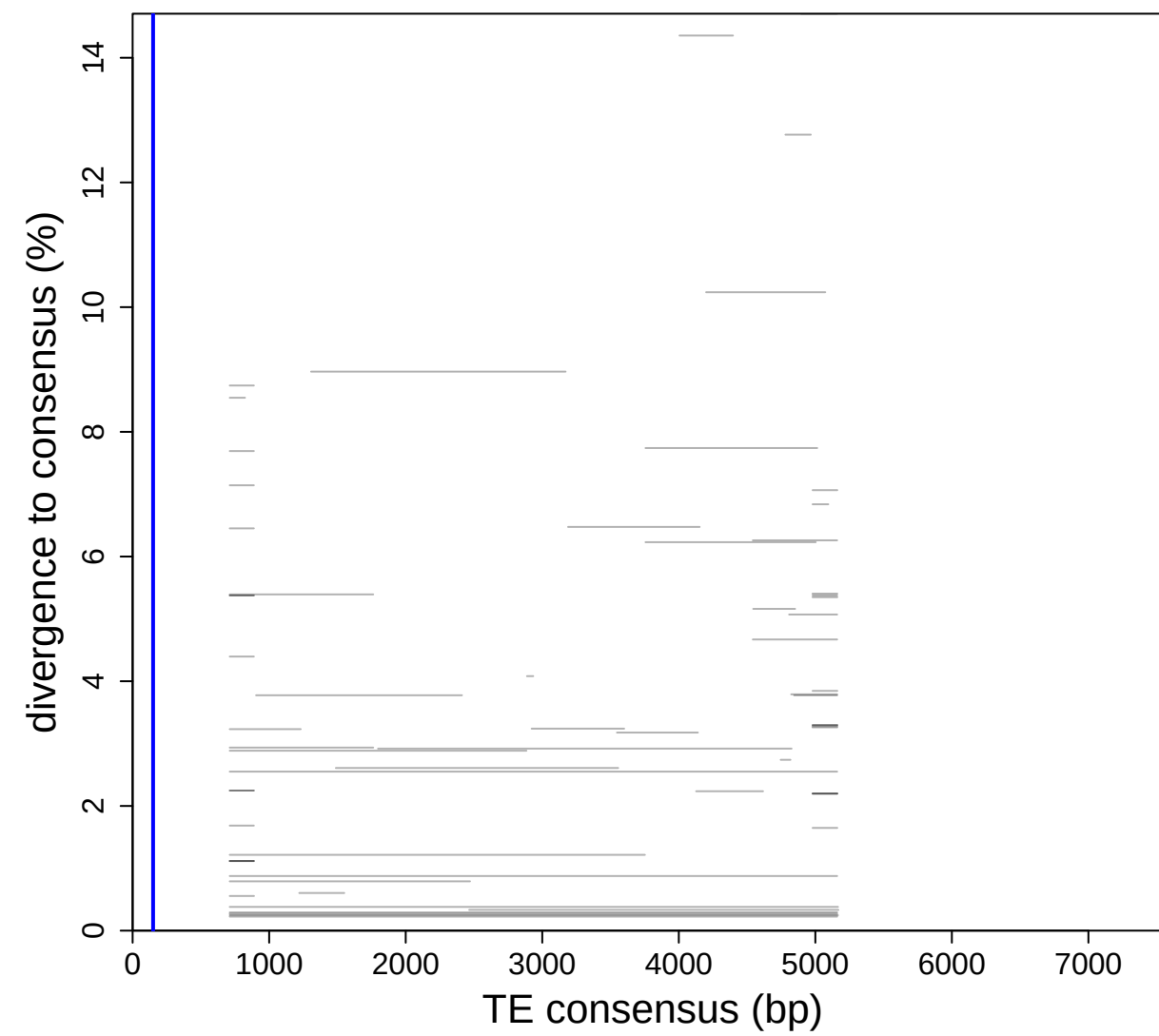
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_al
size: 5769bp; fragments: 66; full length: 0 (>=5192.1bp)

After TEtrimmer 5769 bp

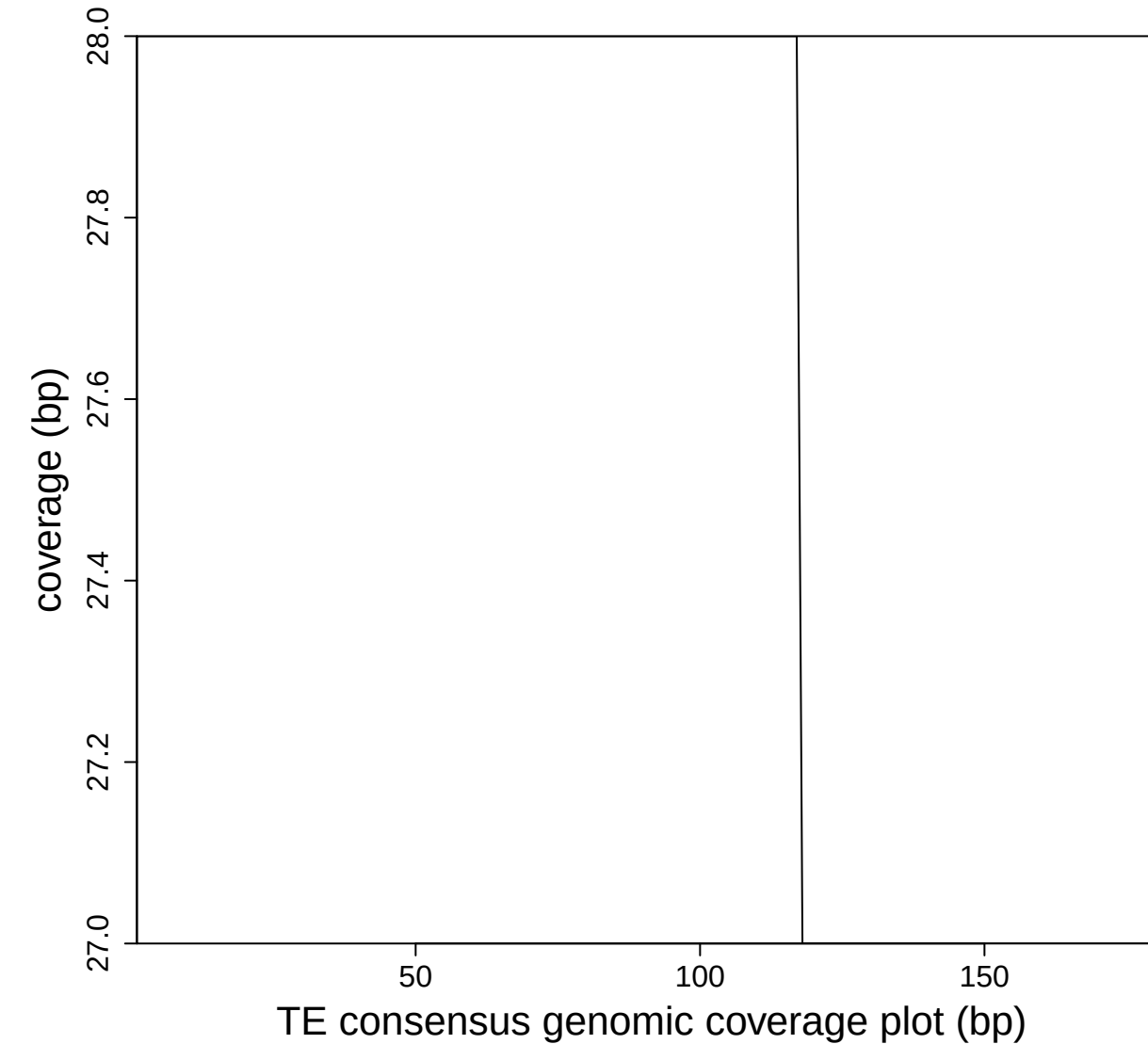
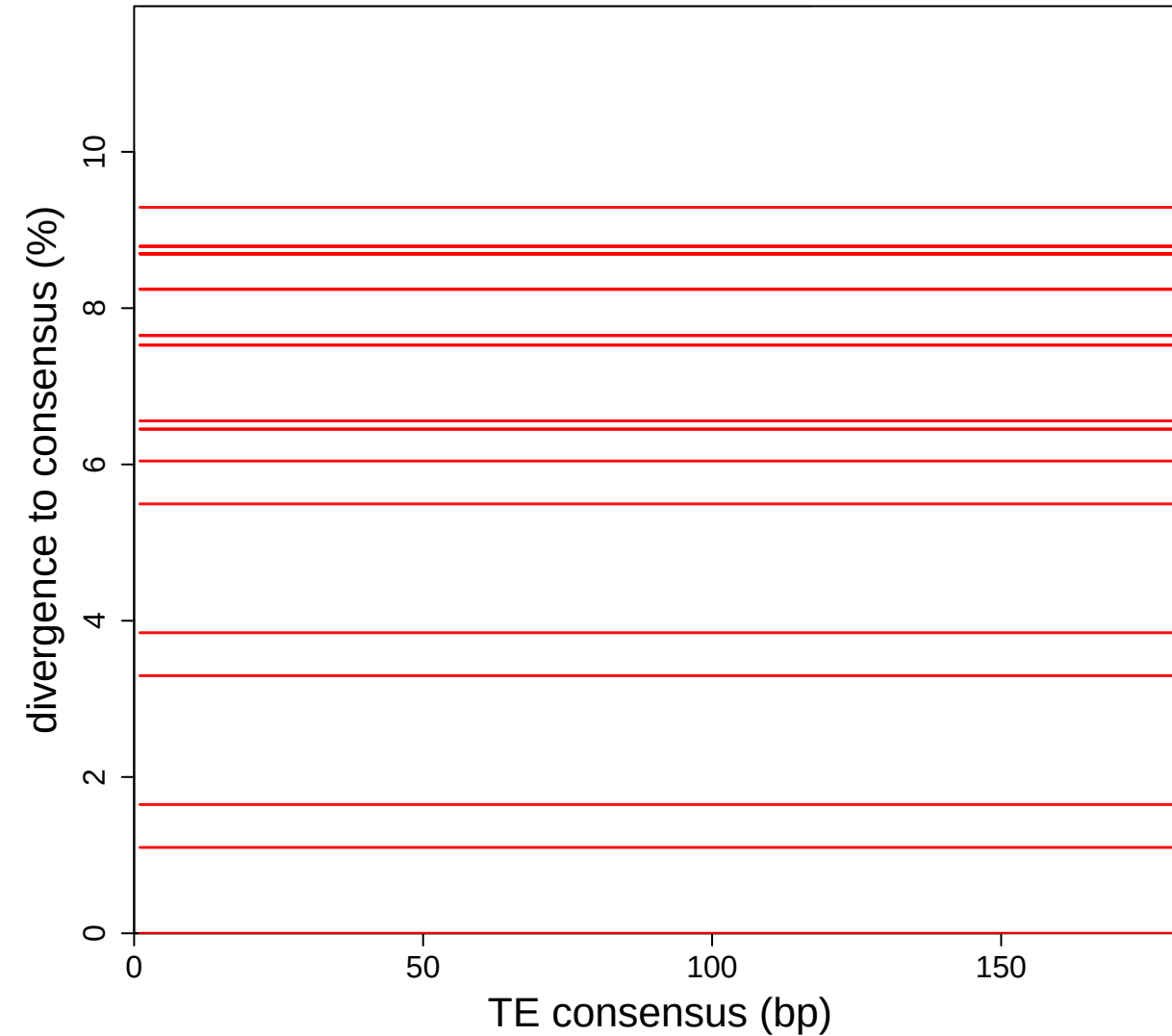


bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 7619bp; fragments: 66; full length: 0 (>=6857.1bp)

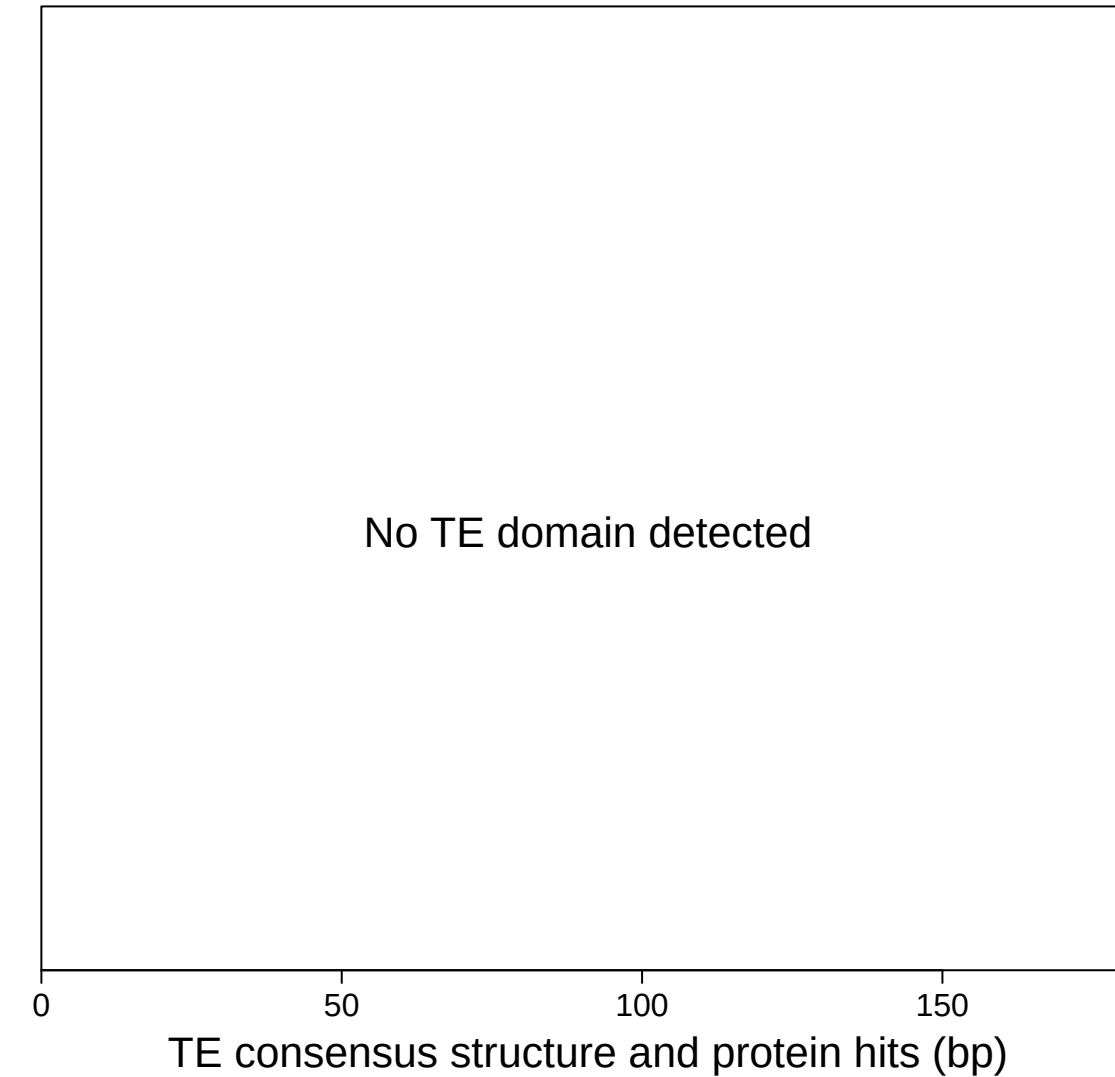
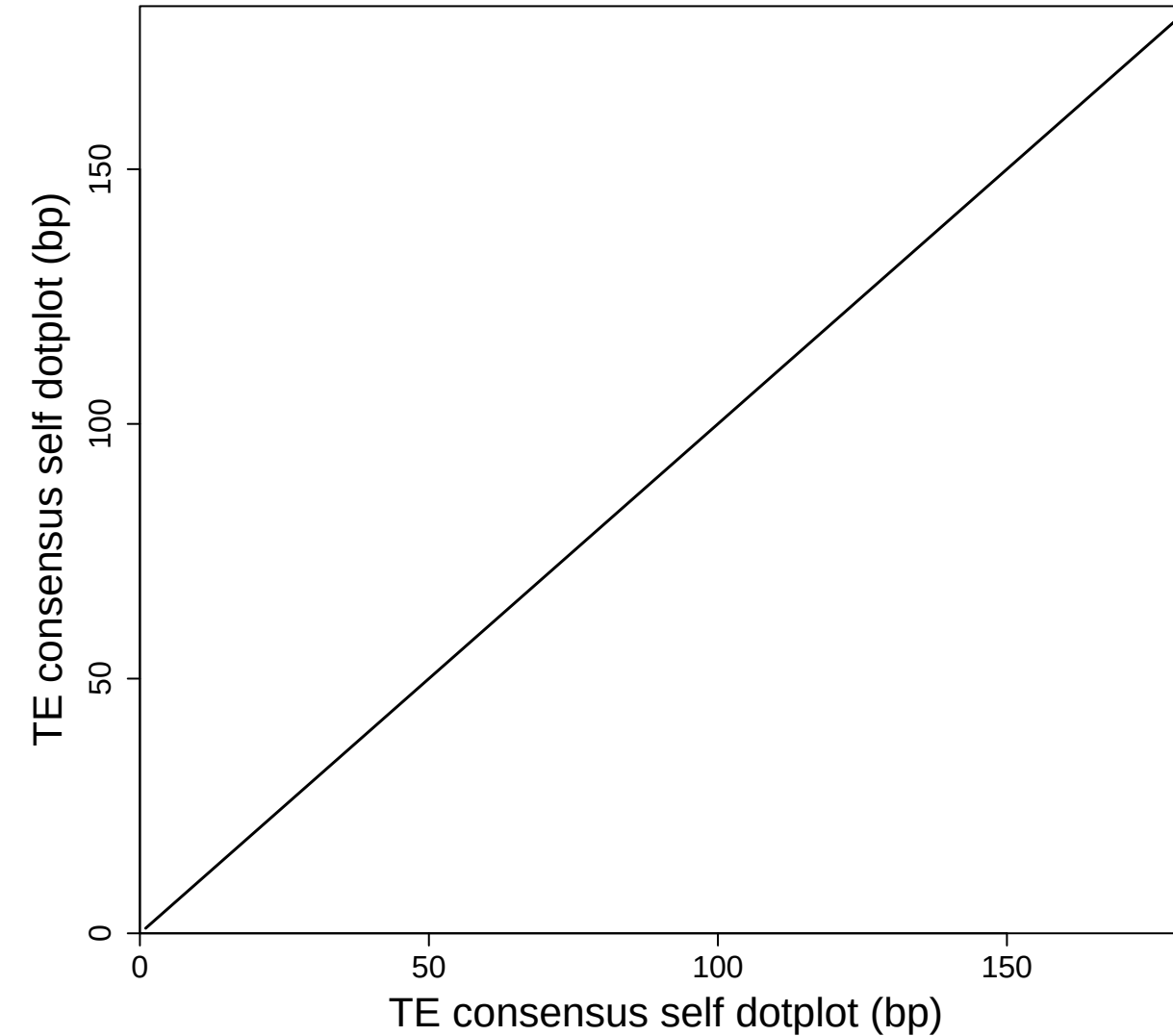
After TEtrimmer Extended plot Blue lines are boundaries



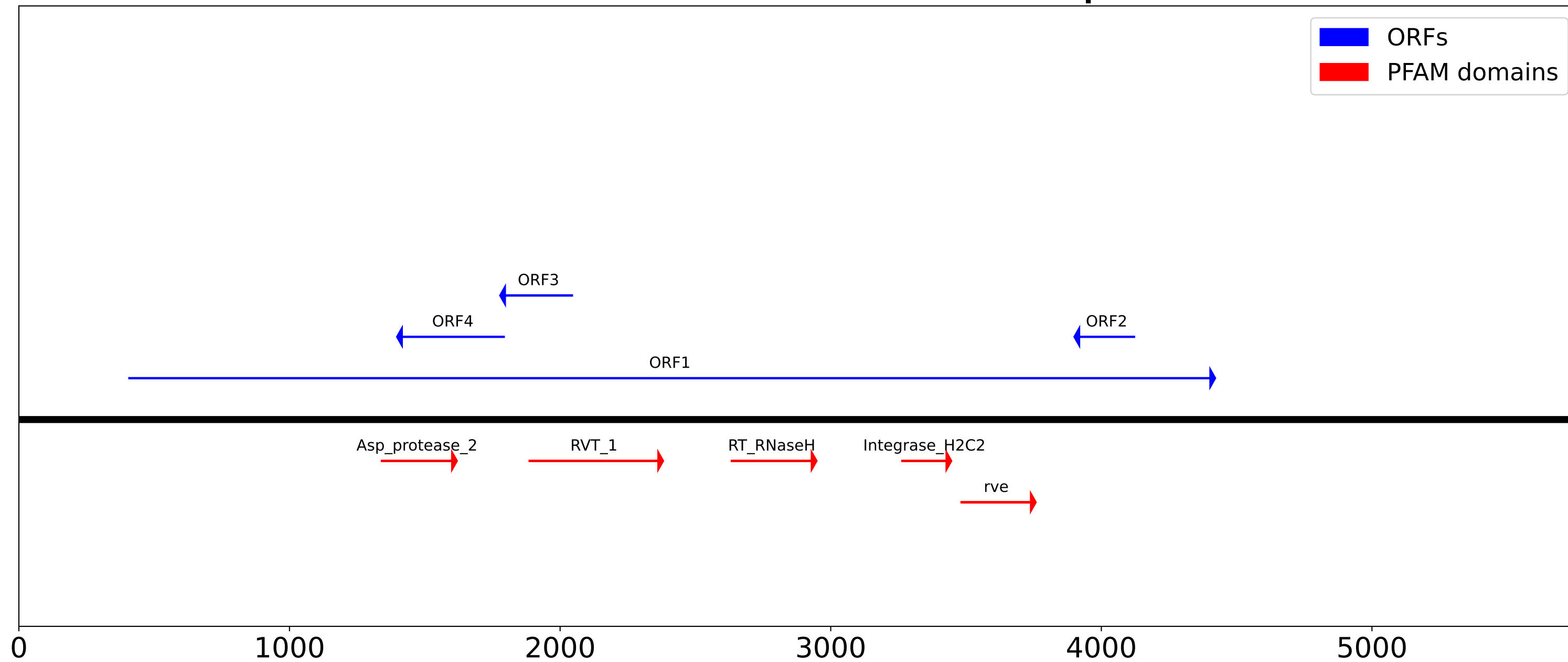
TE: ltr_1_family_413
size: 182bp; fragments: 28; full length: 27 (≥ 163.8 bp)



Before TEtrimmer 182 bp



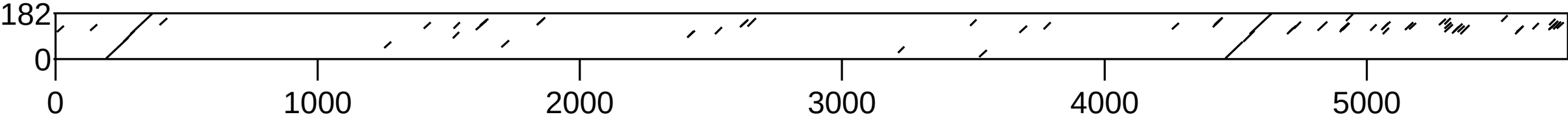
After TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)