

Start crop Point



End crop Point

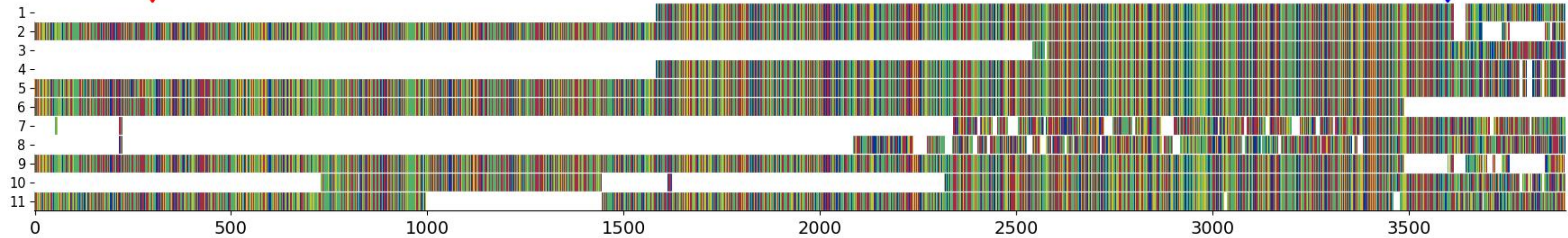


MSA length = 3300



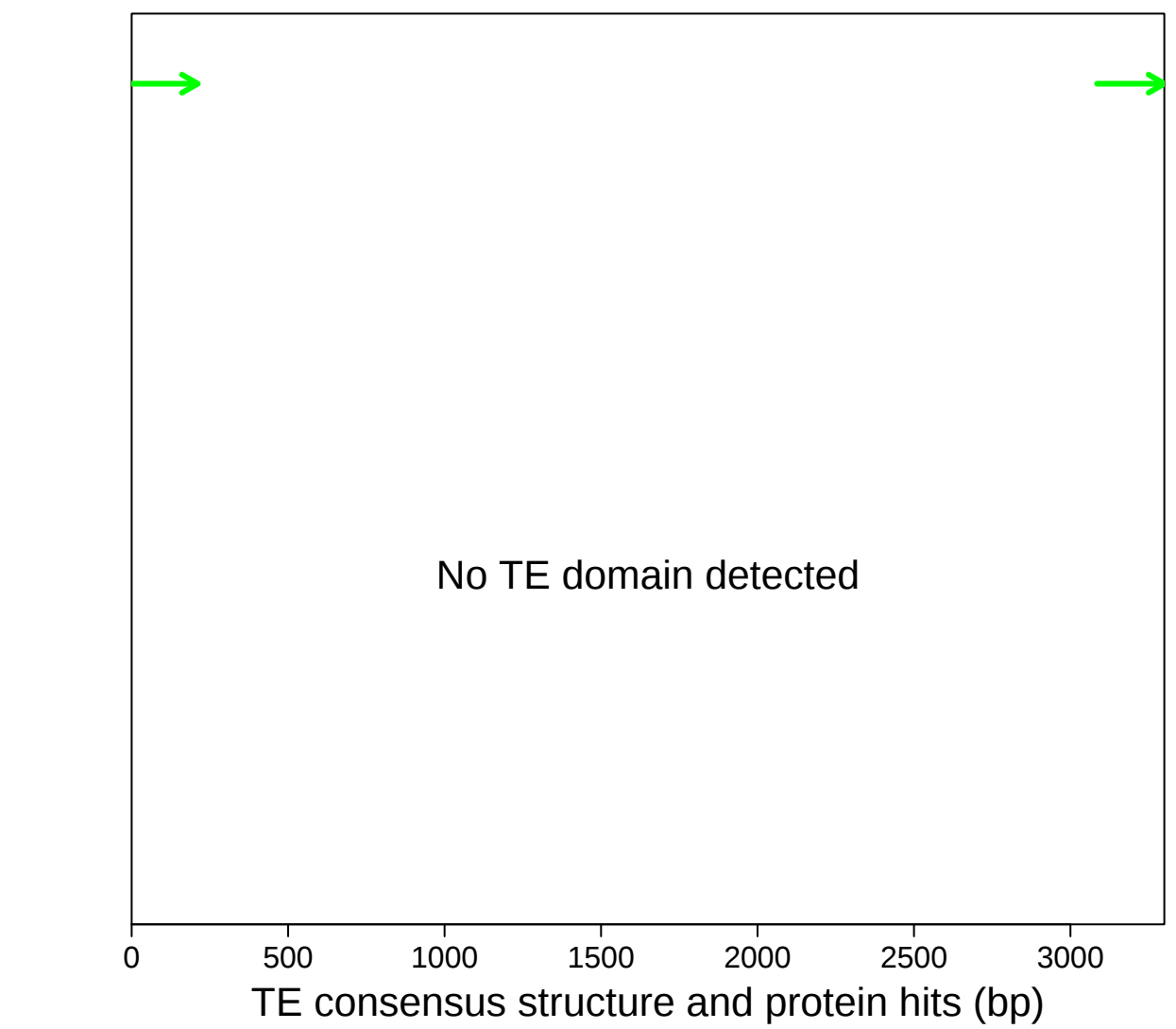
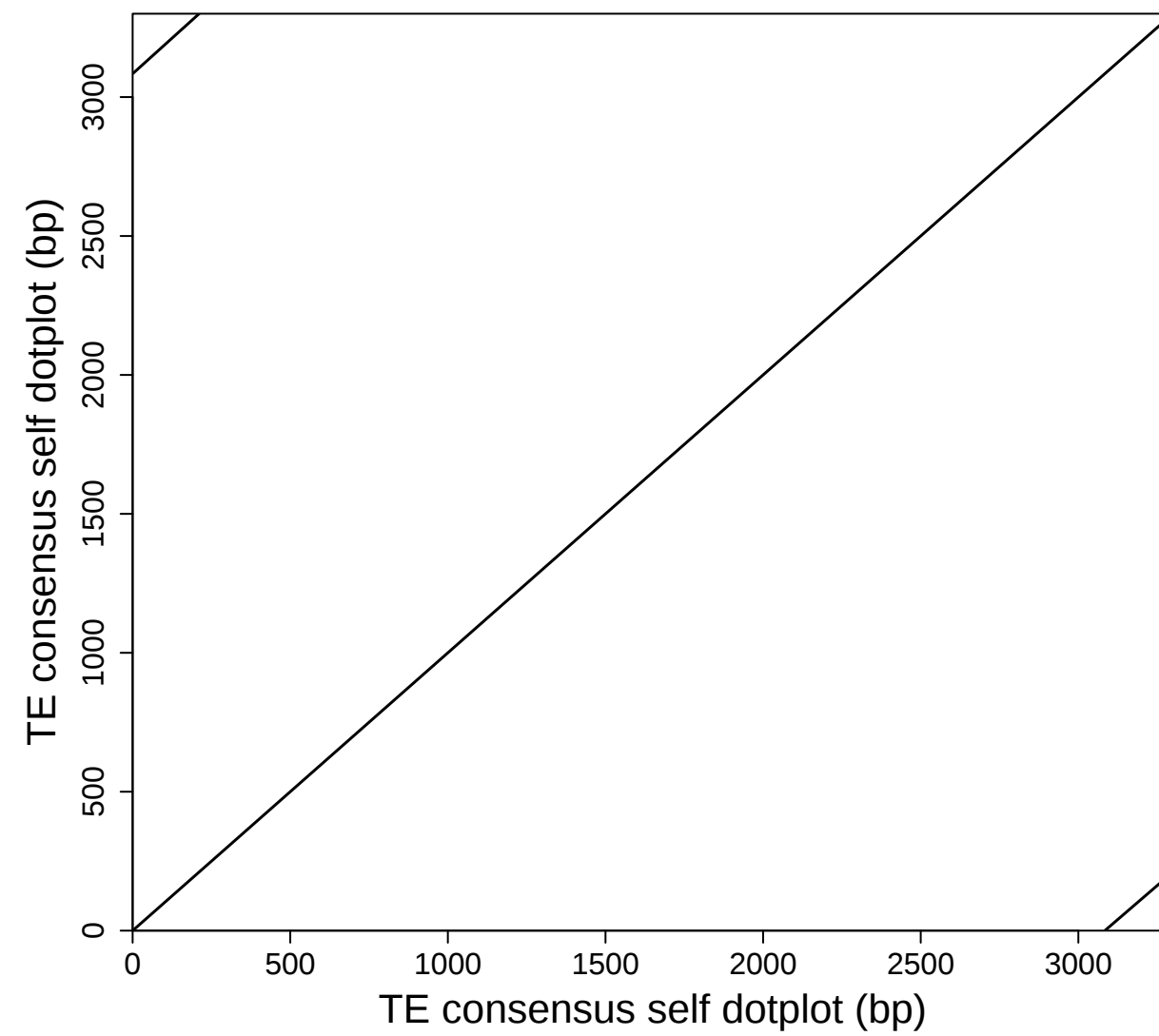
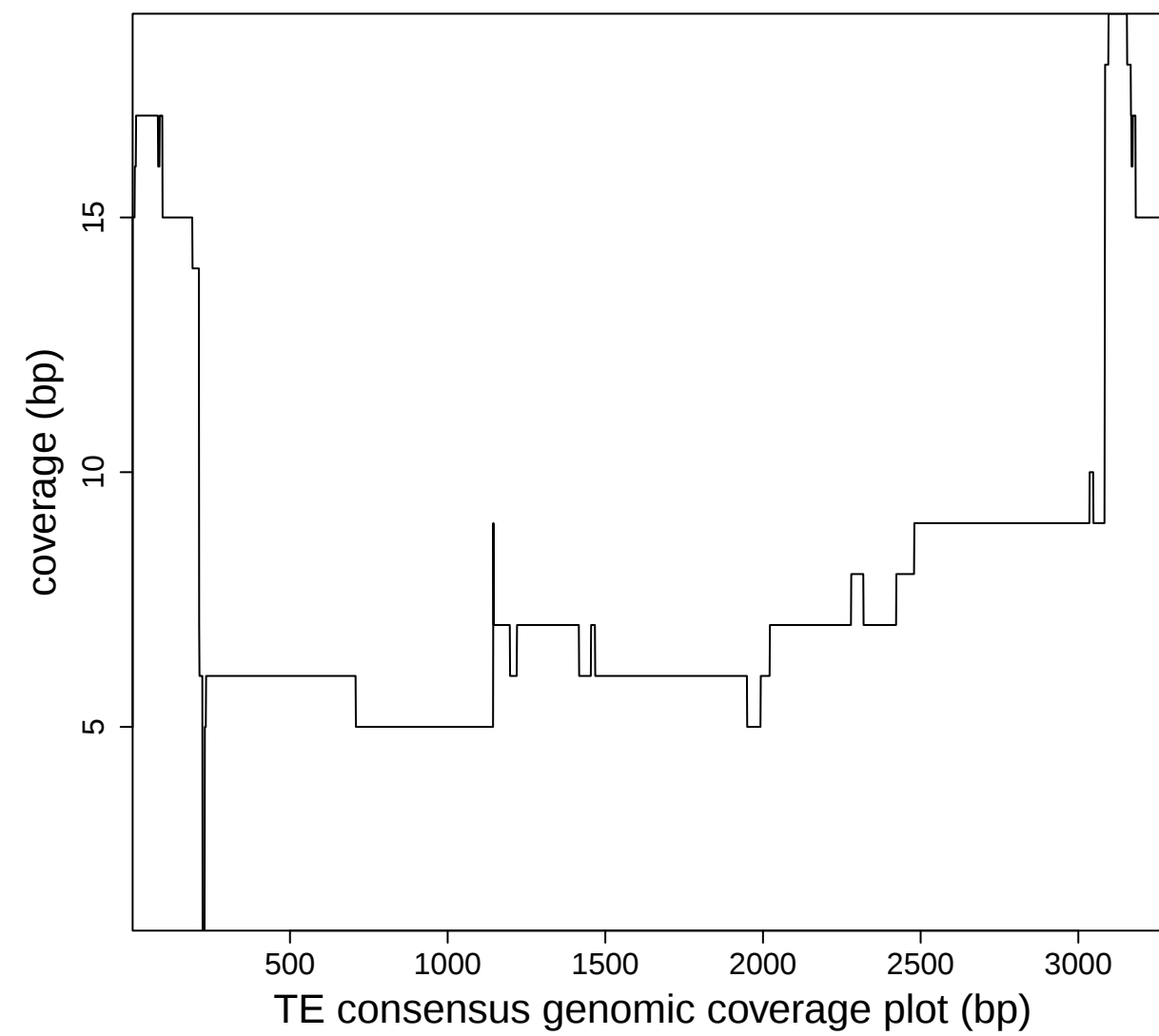
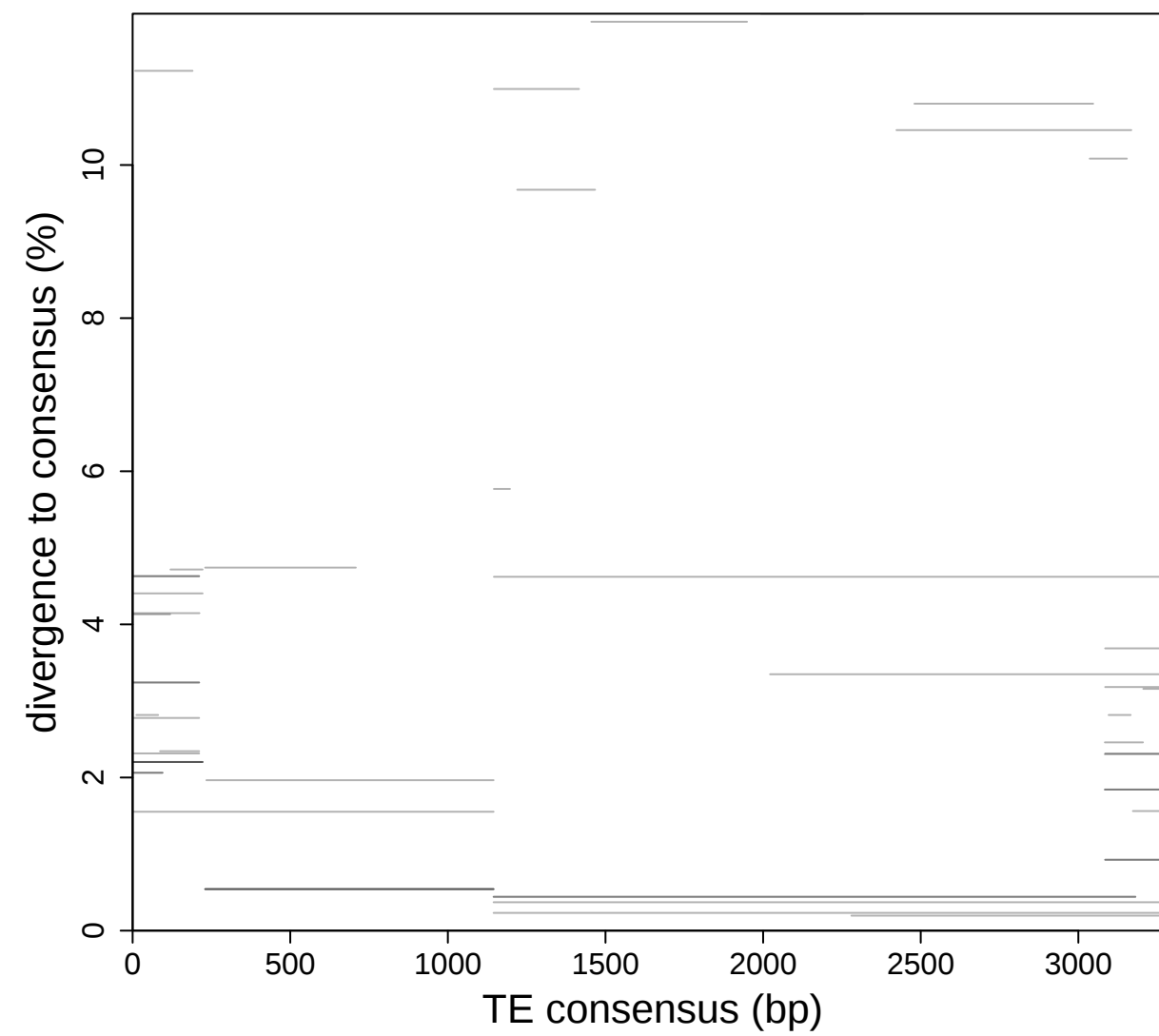
Start crop Point

End crop Point



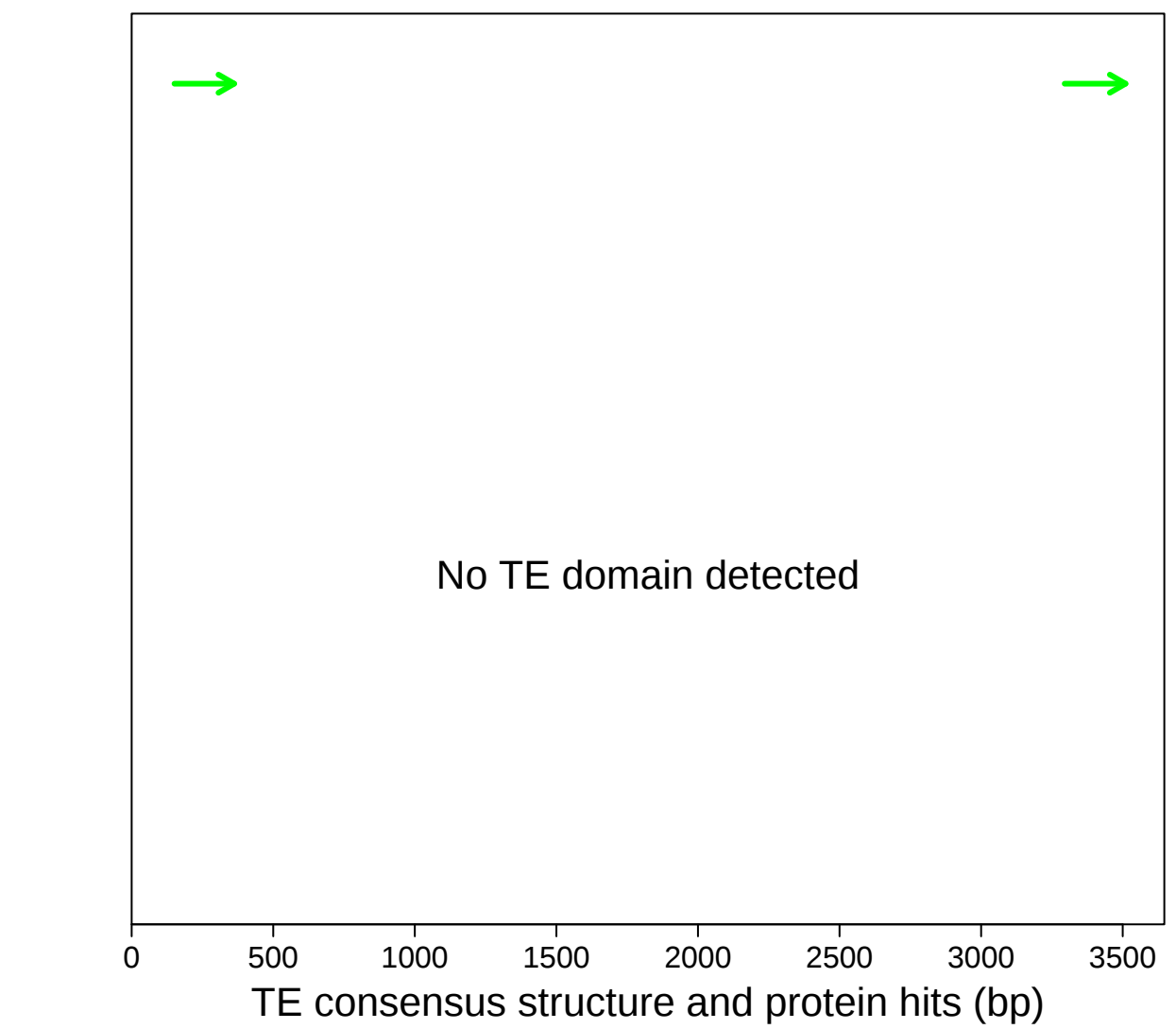
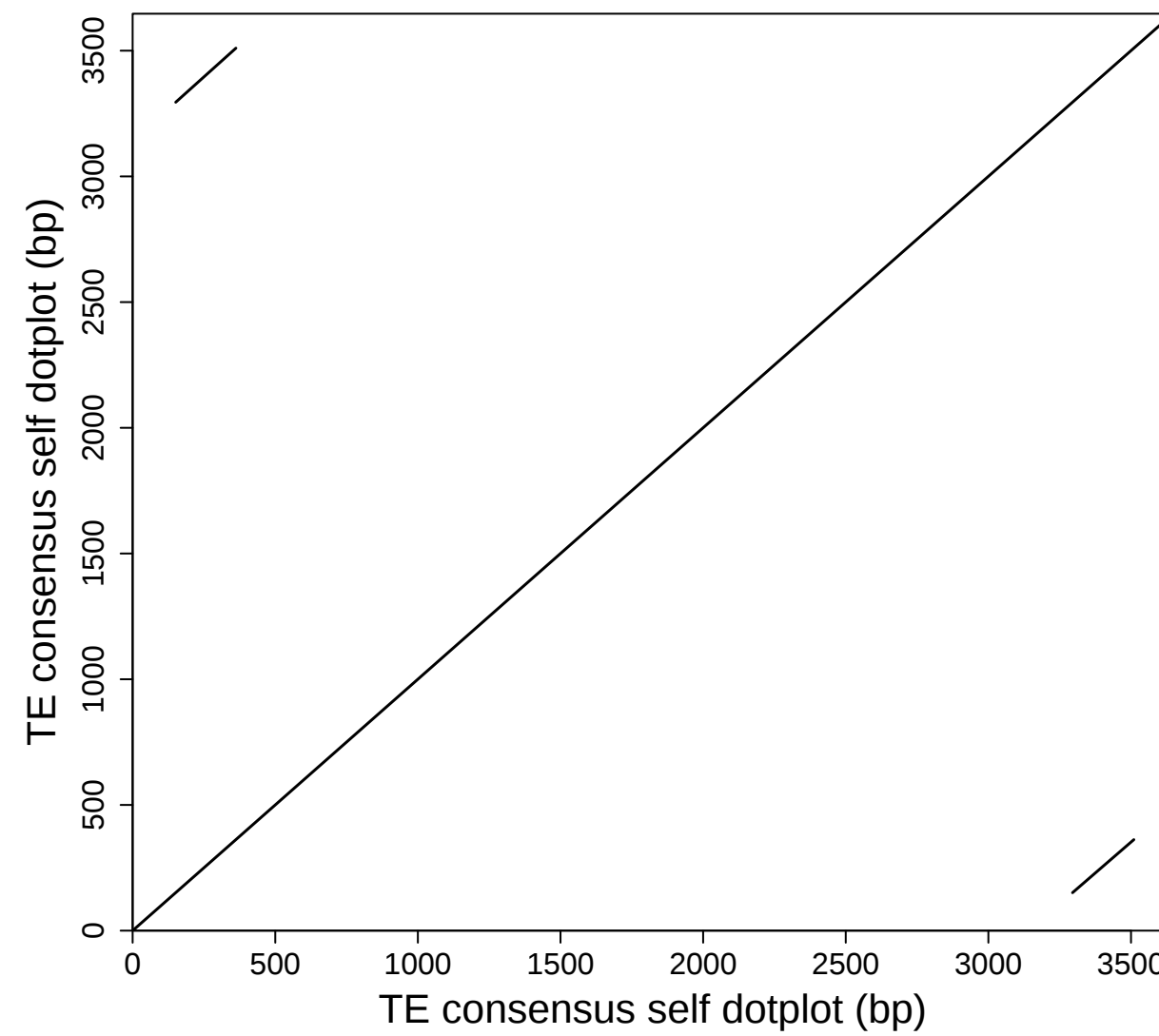
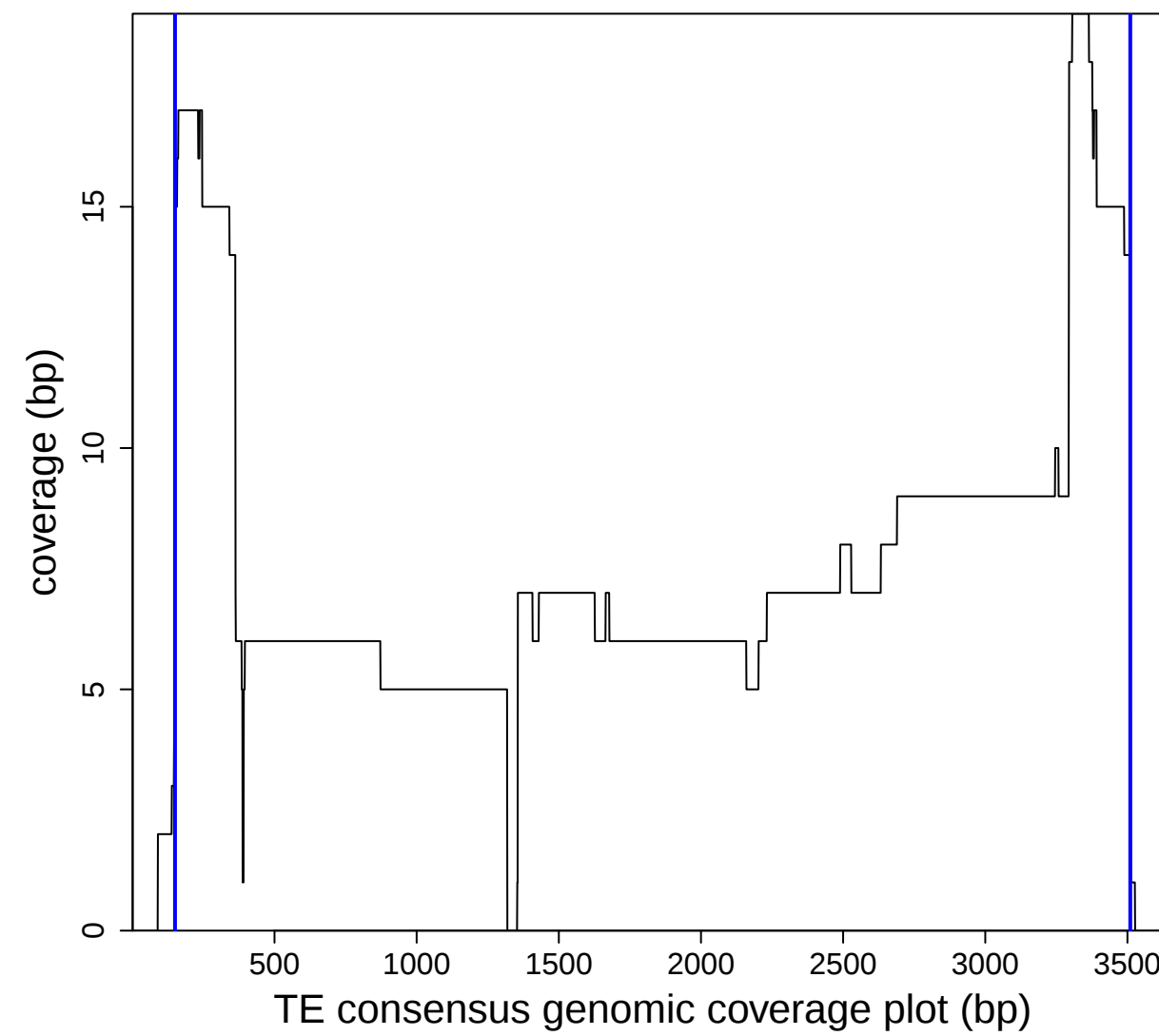
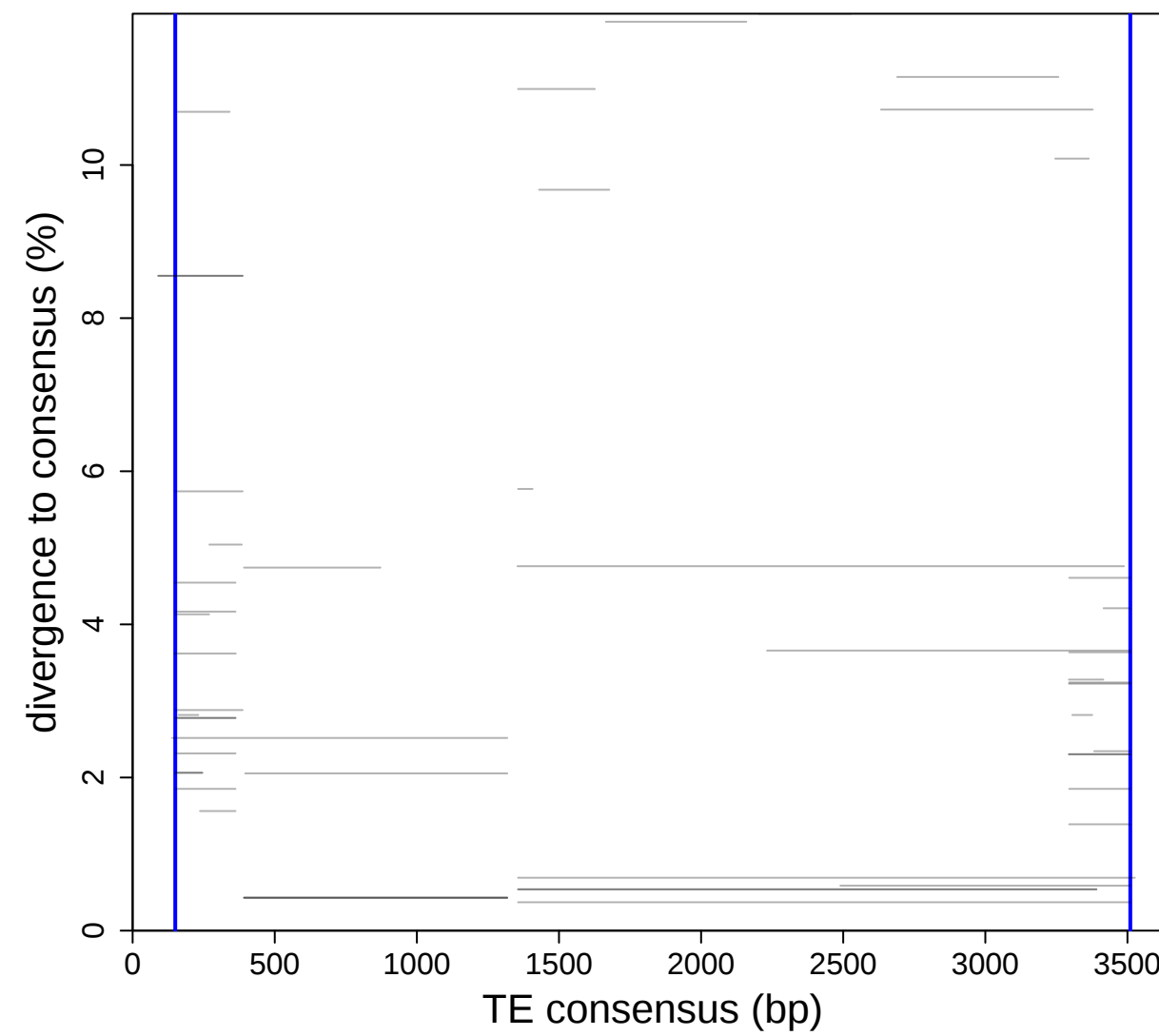
bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.fa
size: 3300bp; fragments: 51; full length: 0 (>=2970bp)

After TEtrimmer 3300 bp

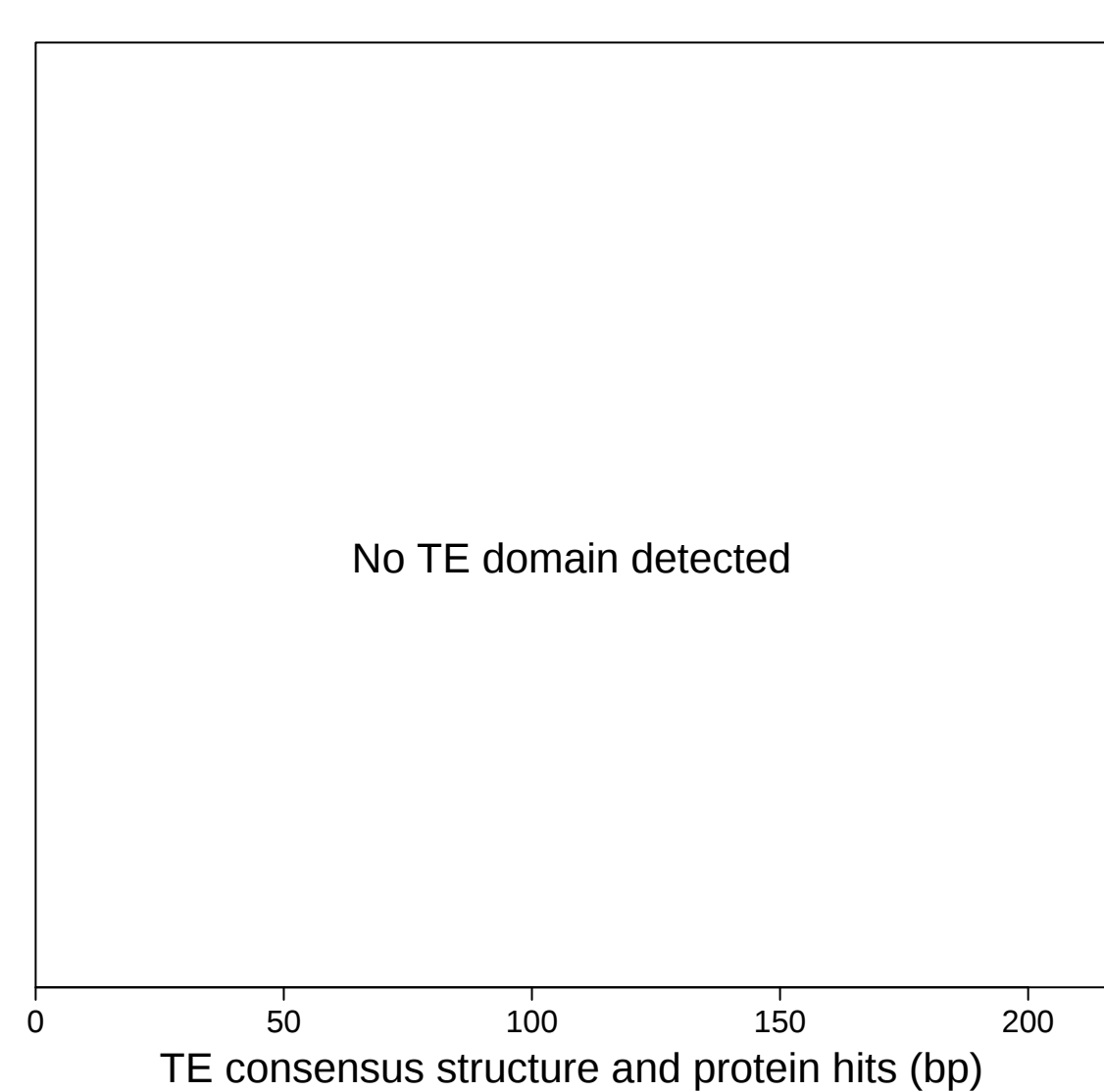
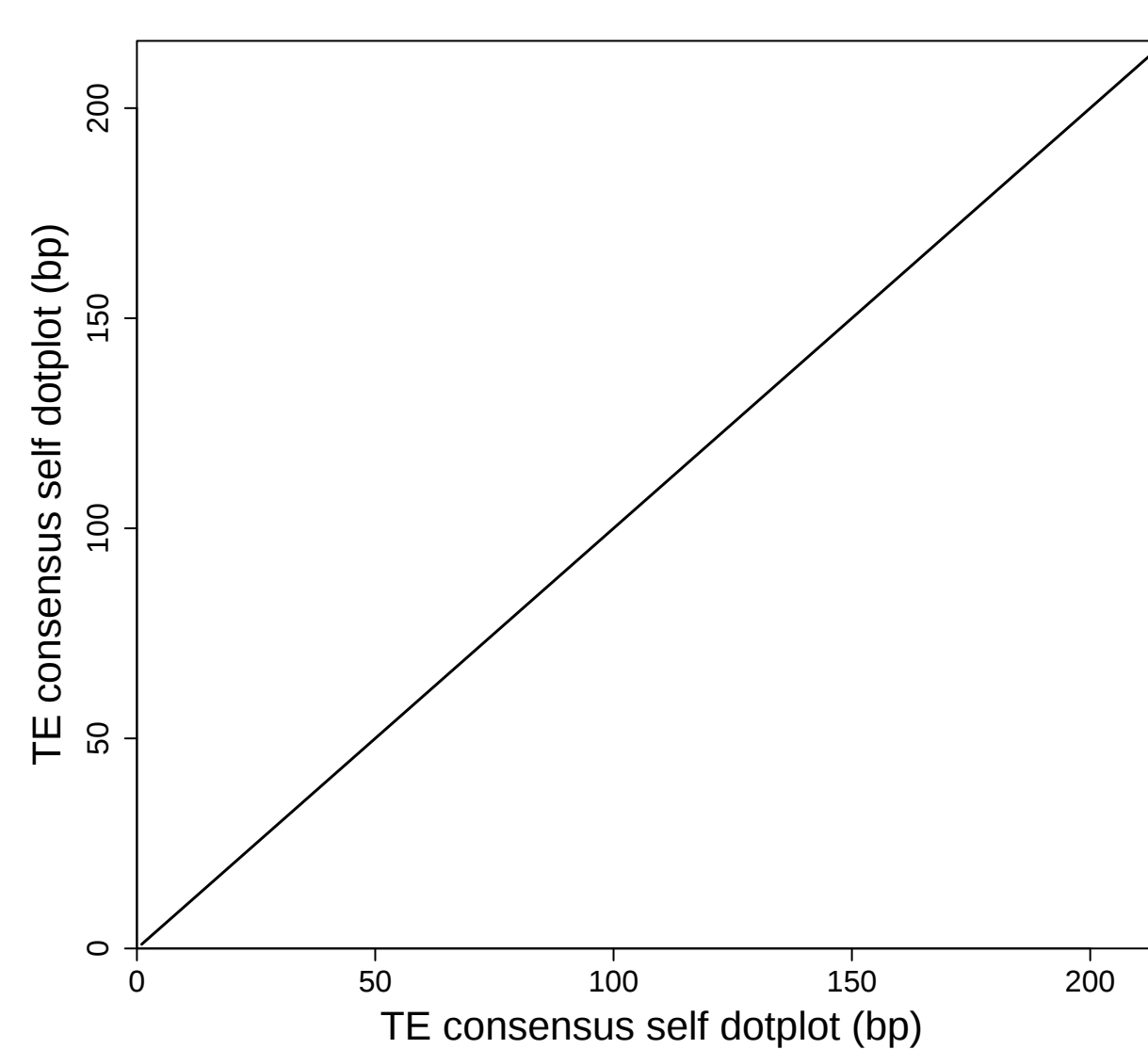
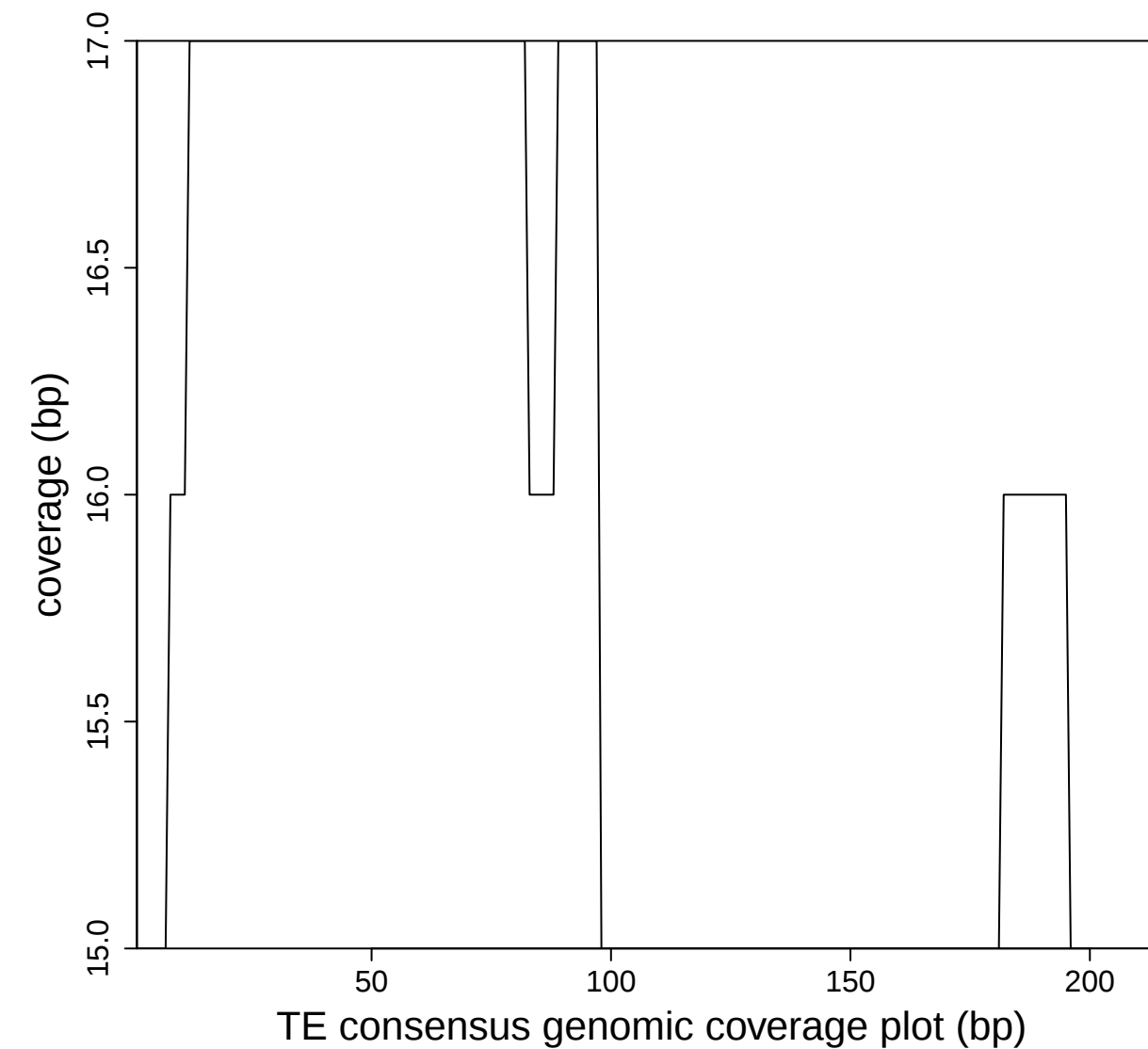
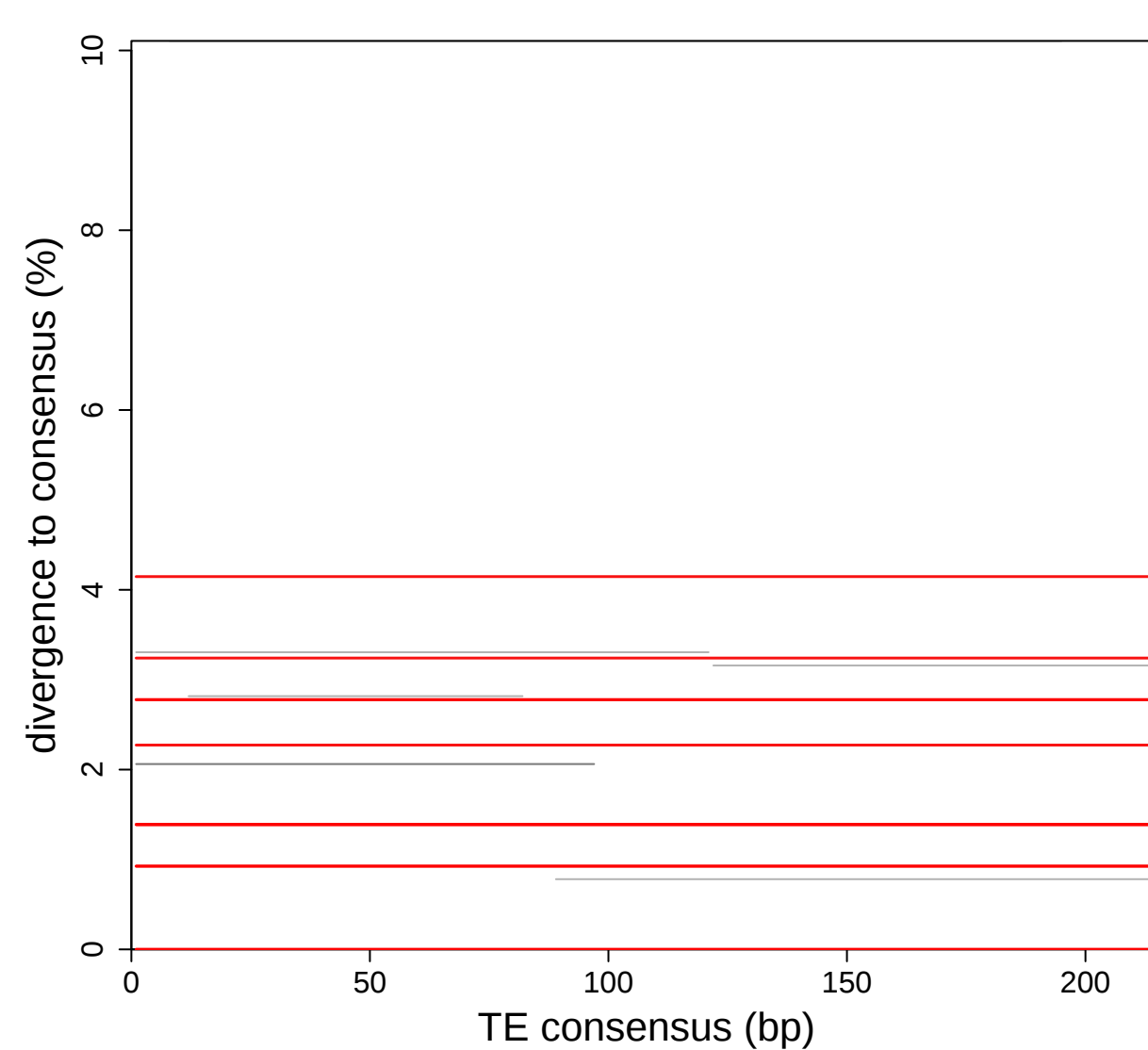


ed_uf.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_
size: 3660bp; fragments: 51; full length: 0 (>=3294bp)

After TEtrimmer Extended plot Blue lines are boundaries

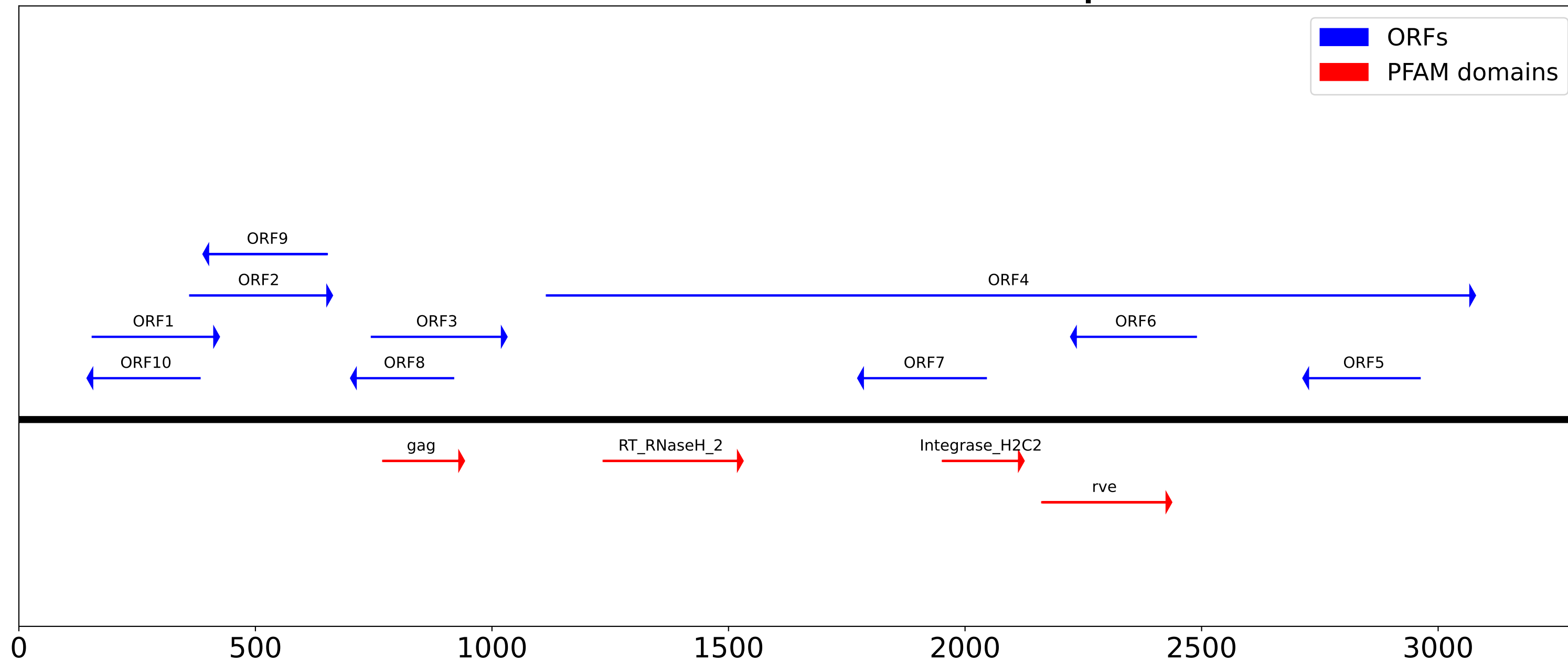


TE: ltr_1_family_235
size: 216bp; fragments: 20; full length: 12 (≥ 194.4 bp)



Before TEtrimmer 216 bp

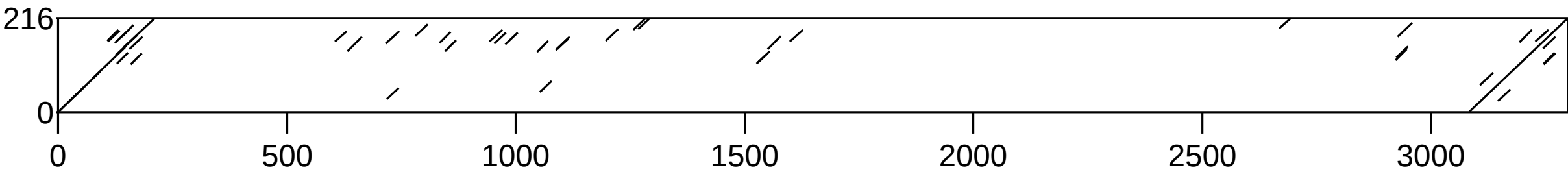
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)