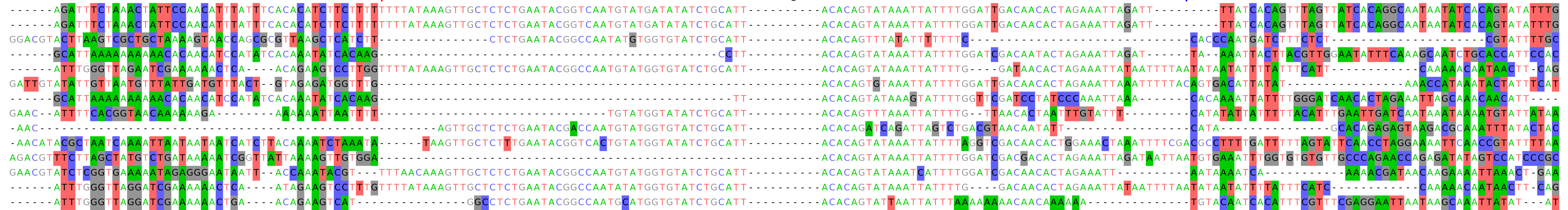
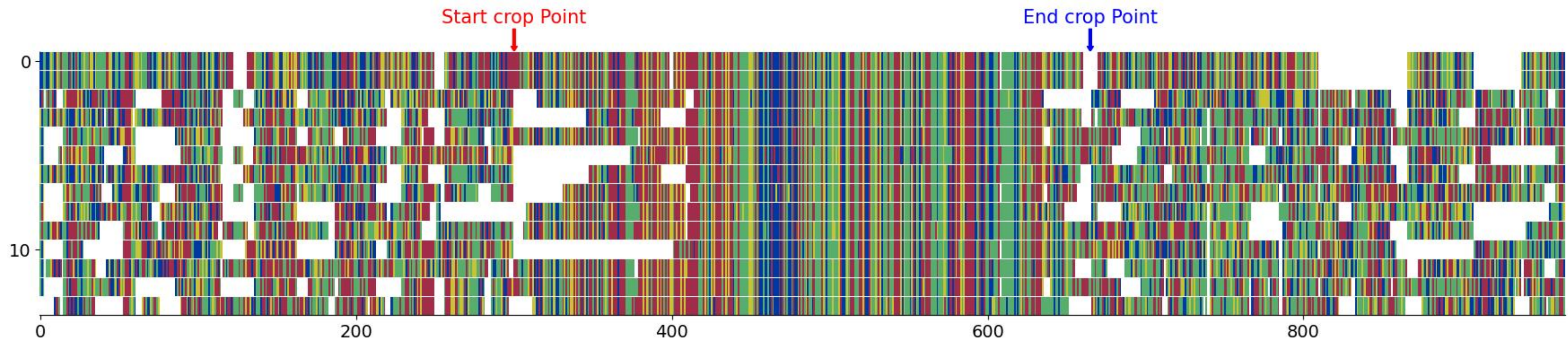


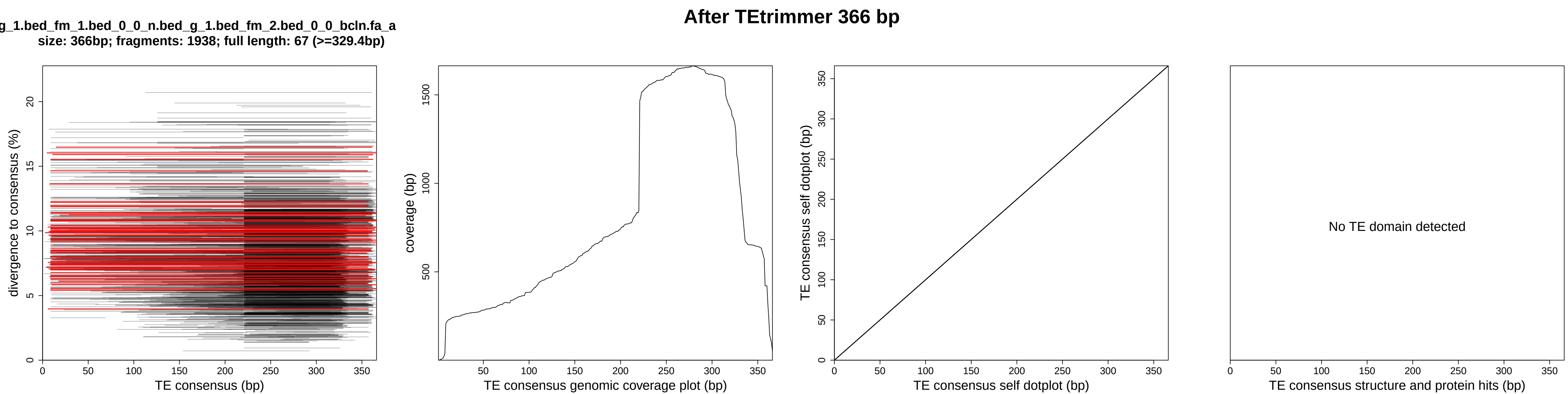
Start crop Point

End crop Point

MSA length = 366

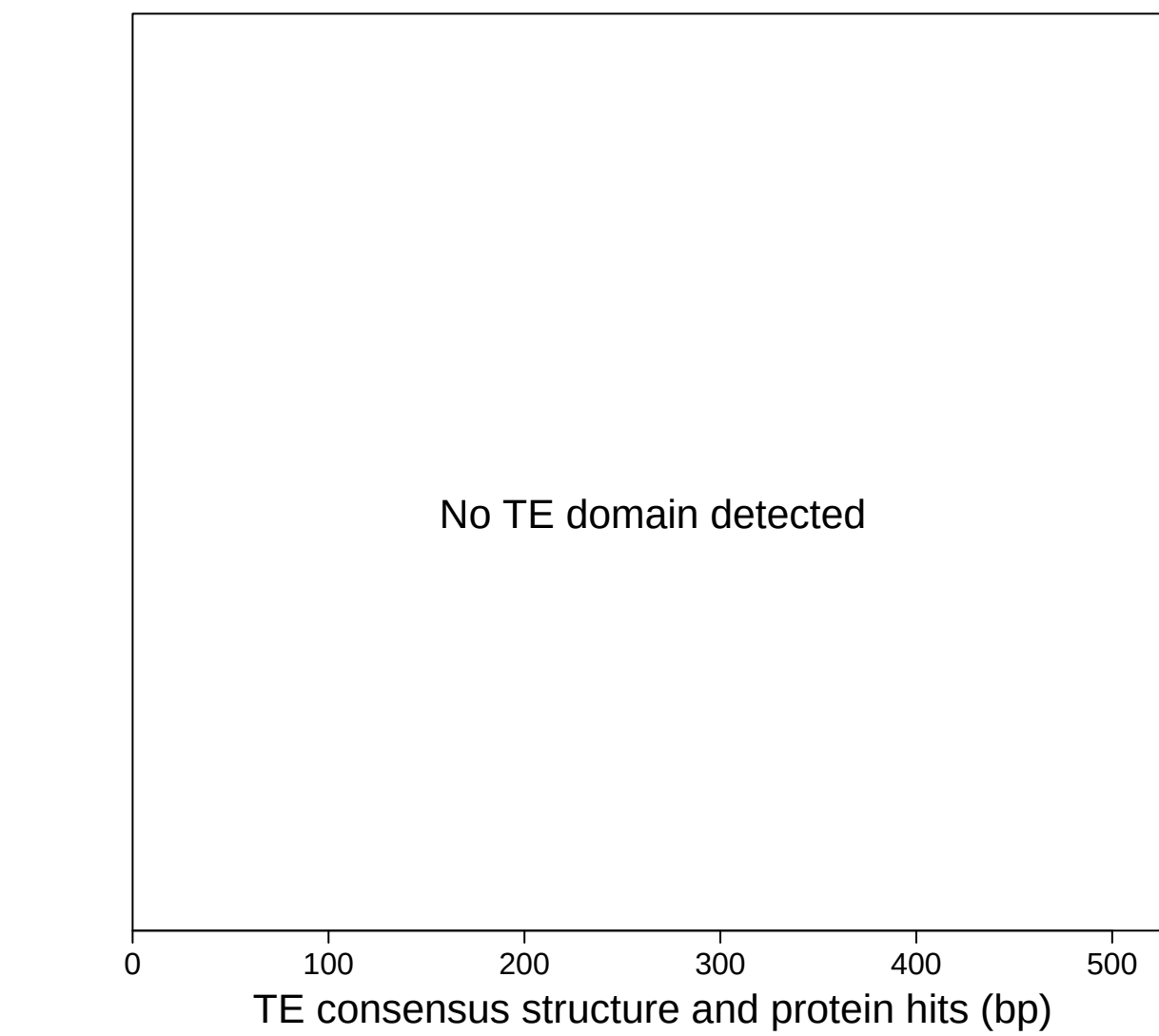
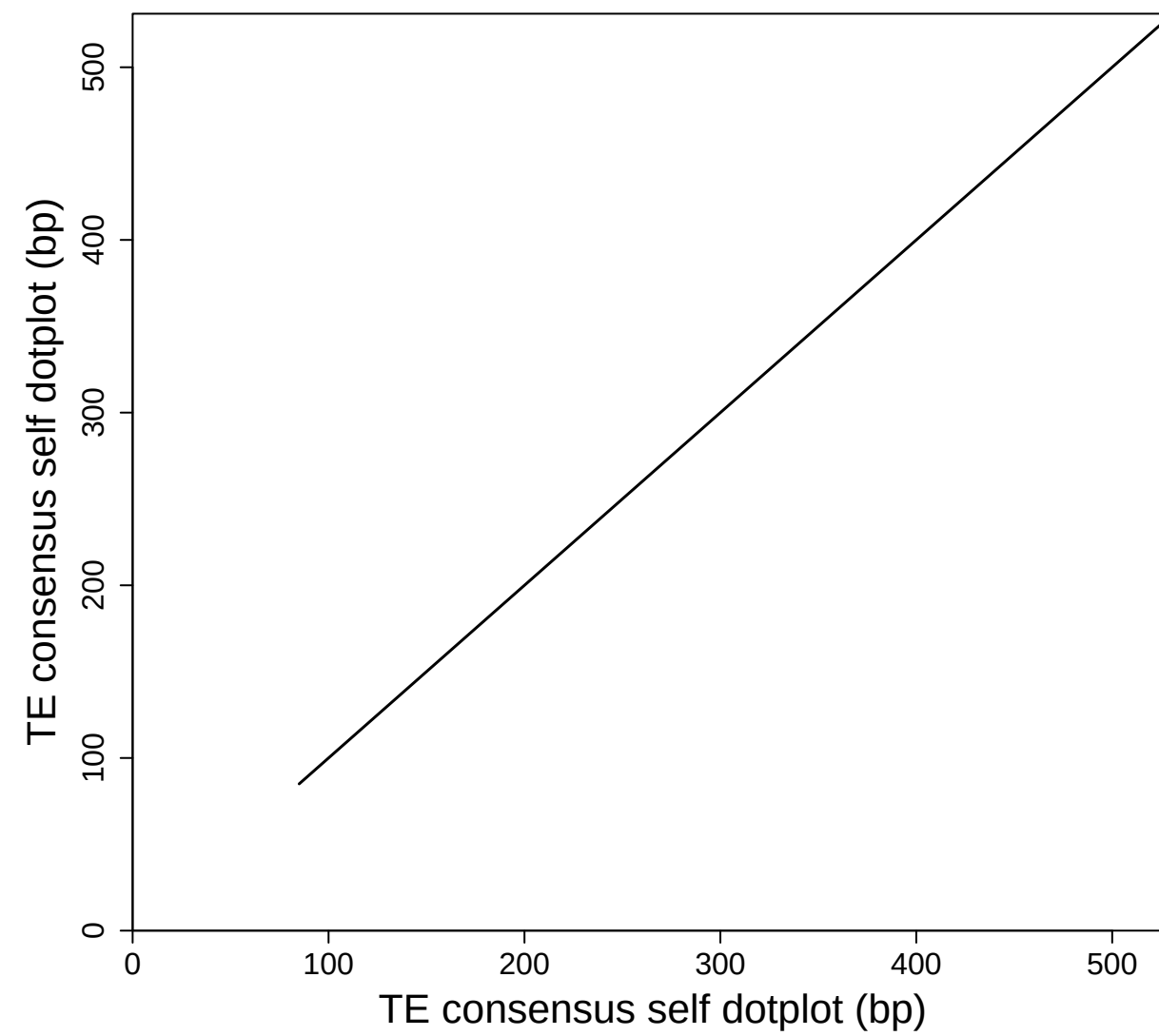
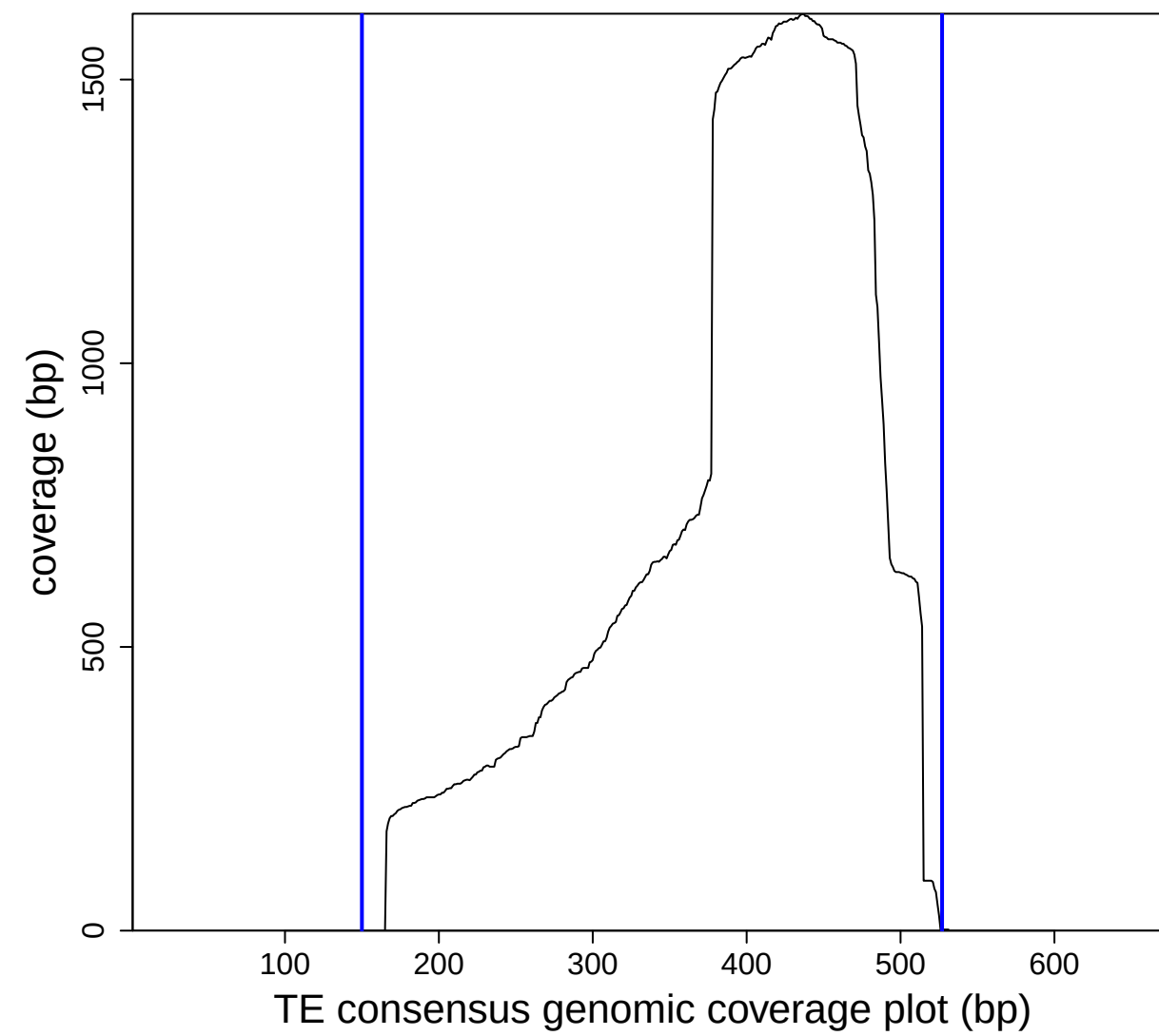
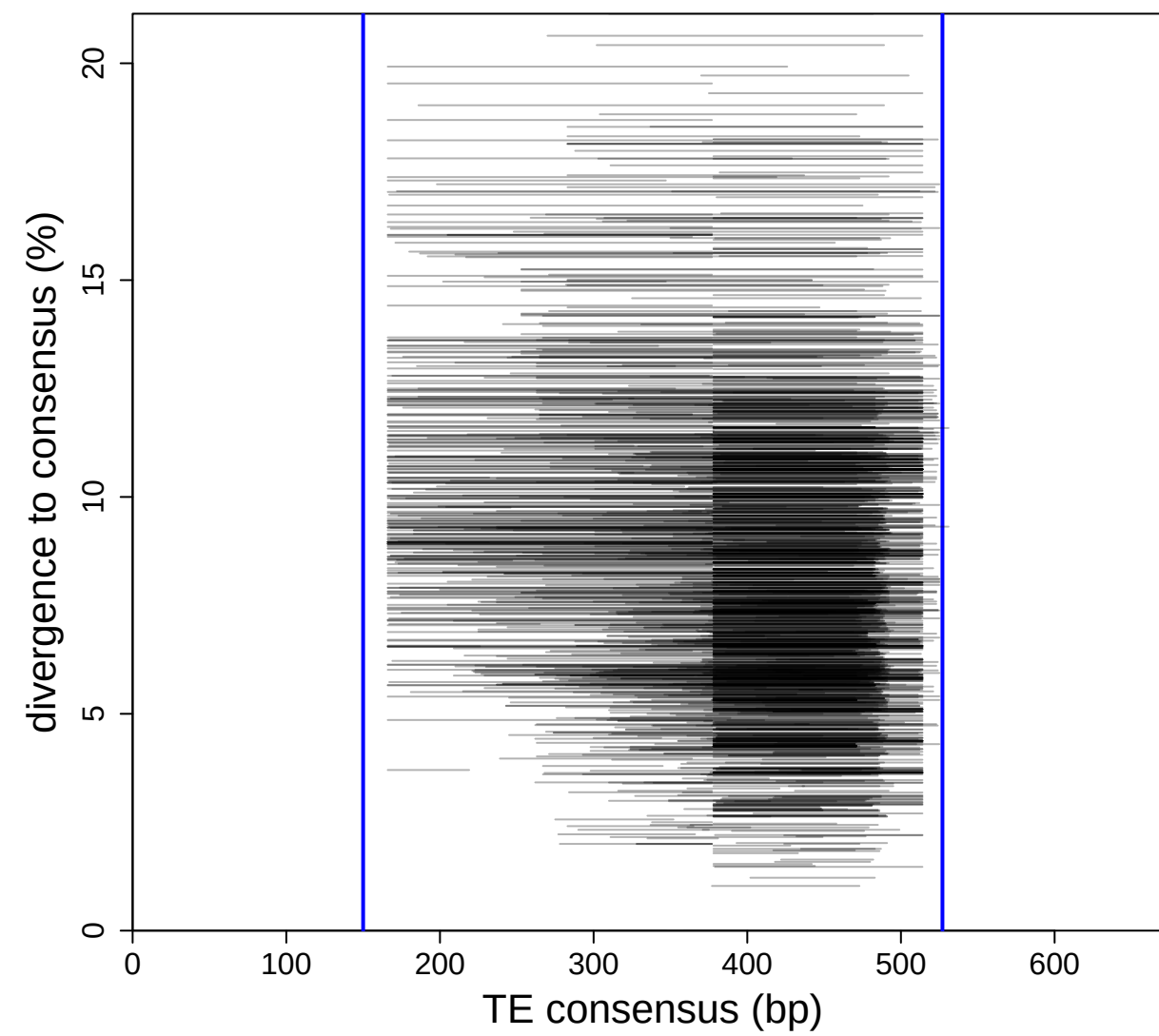






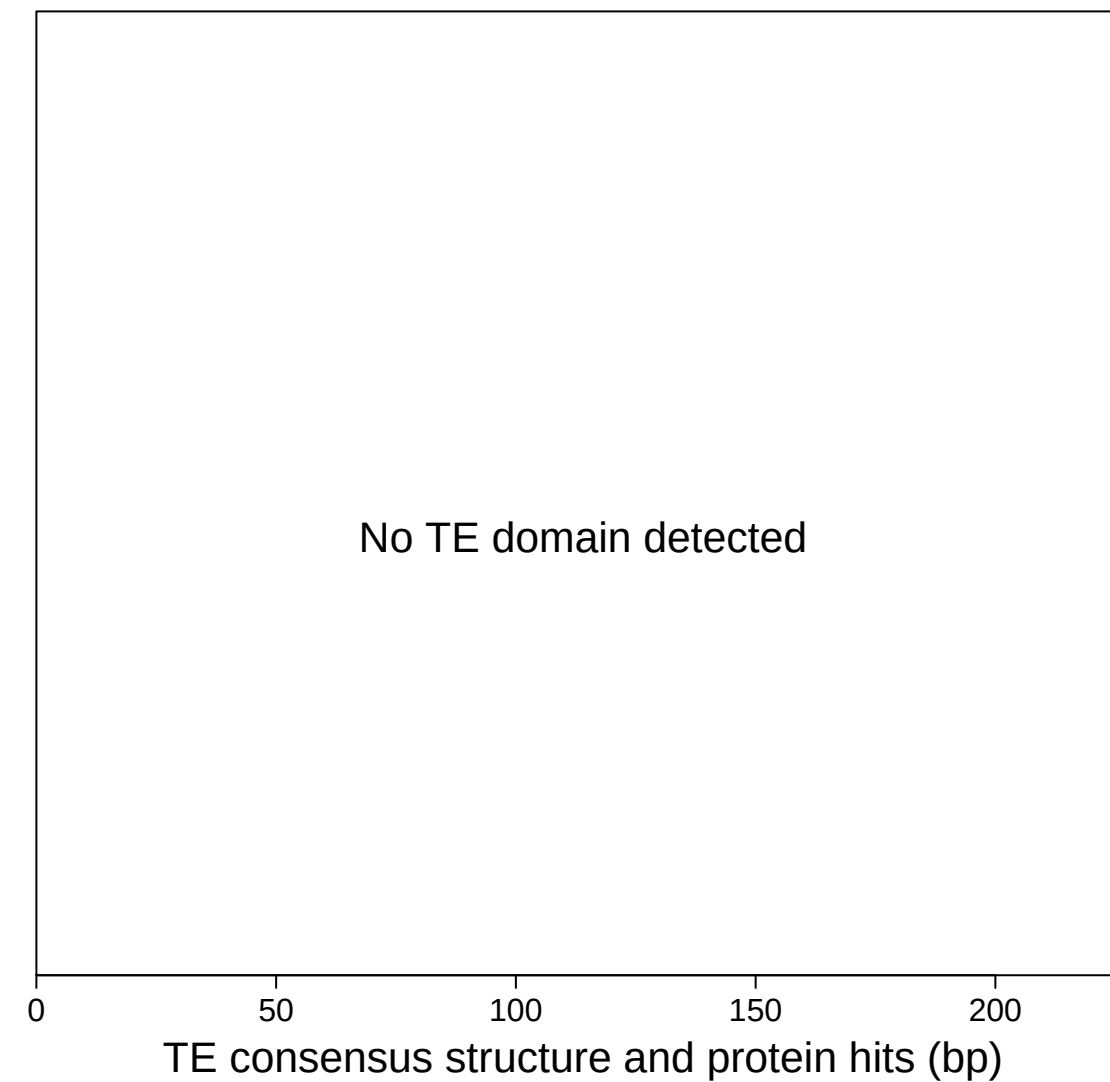
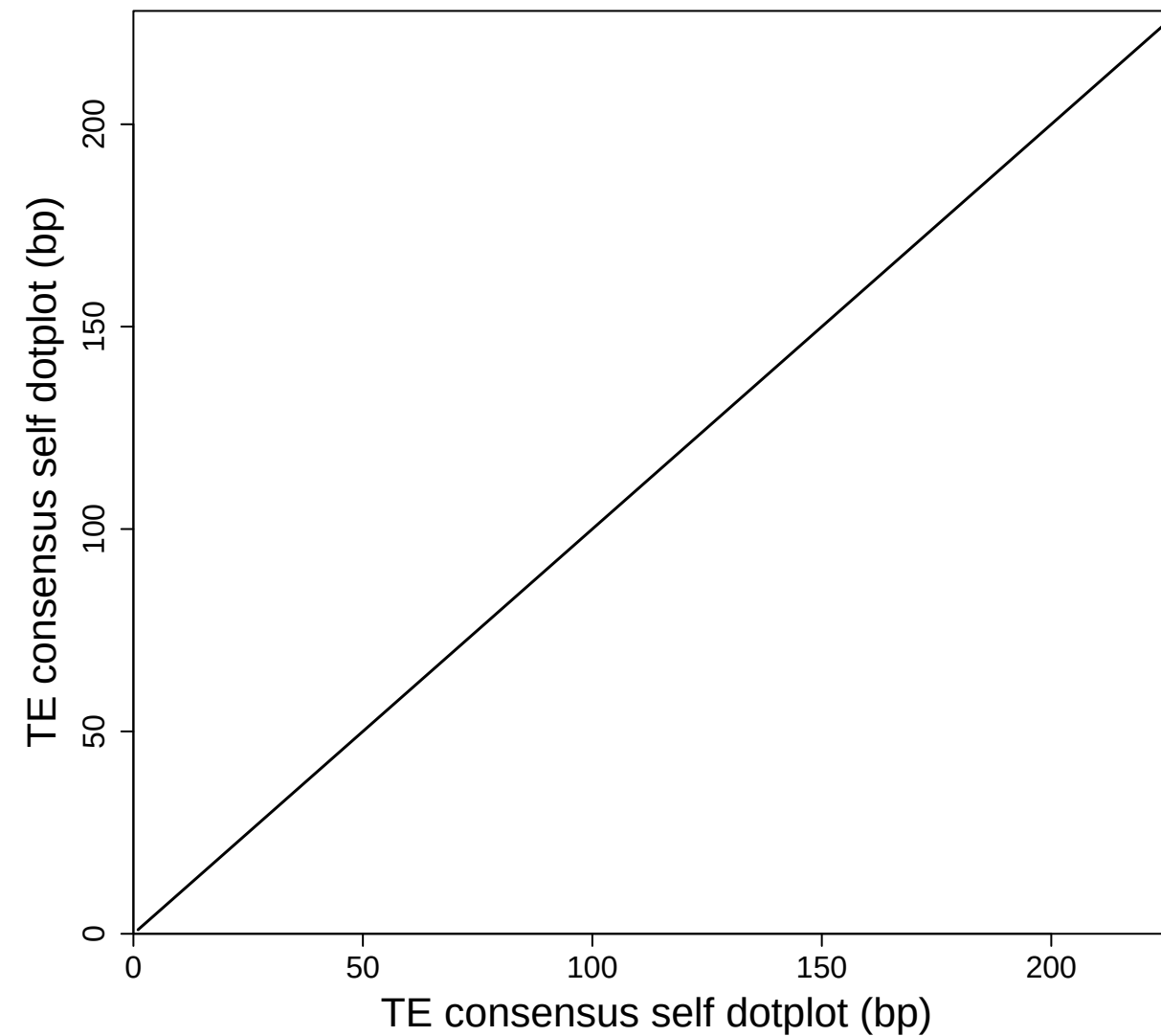
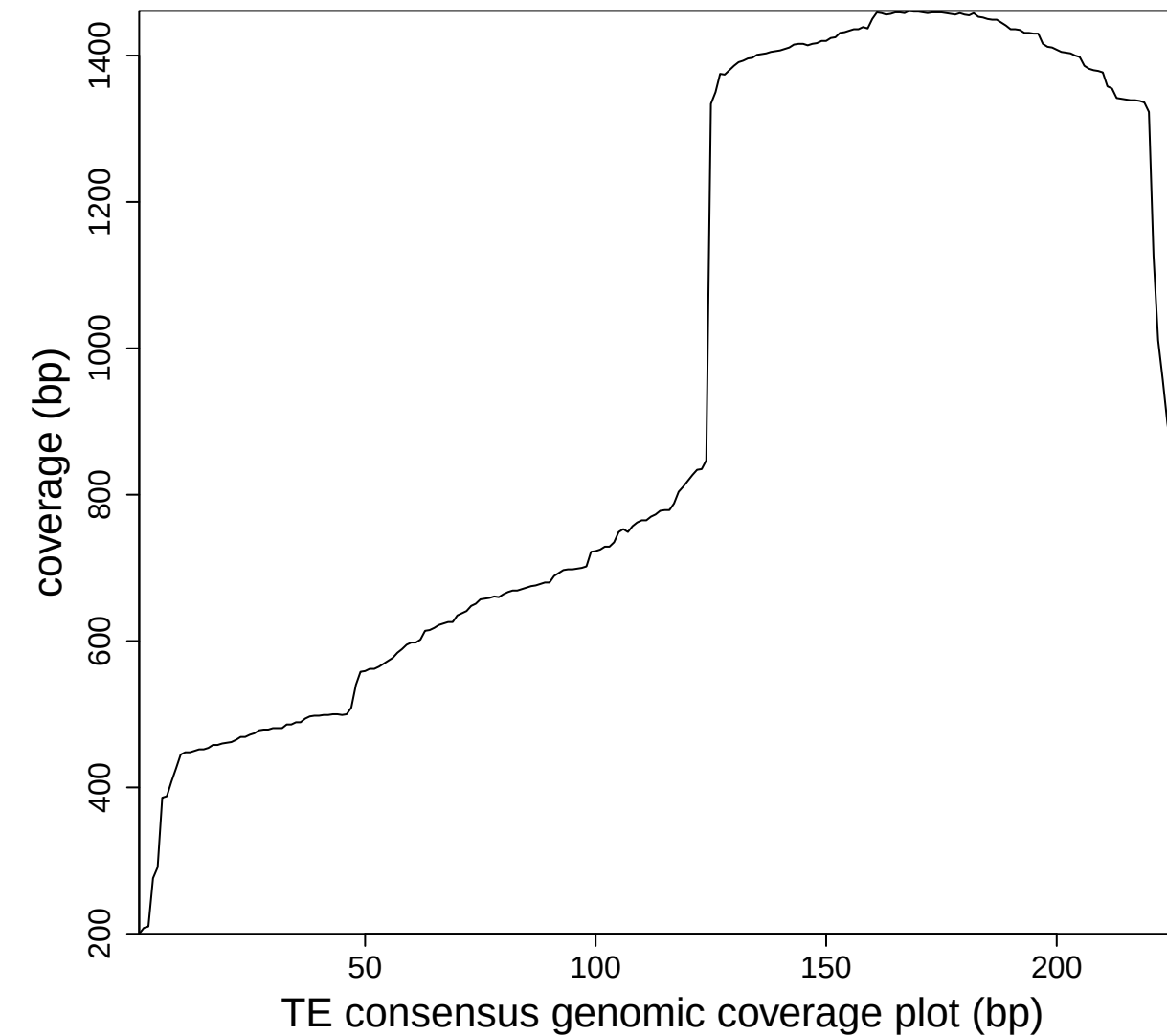
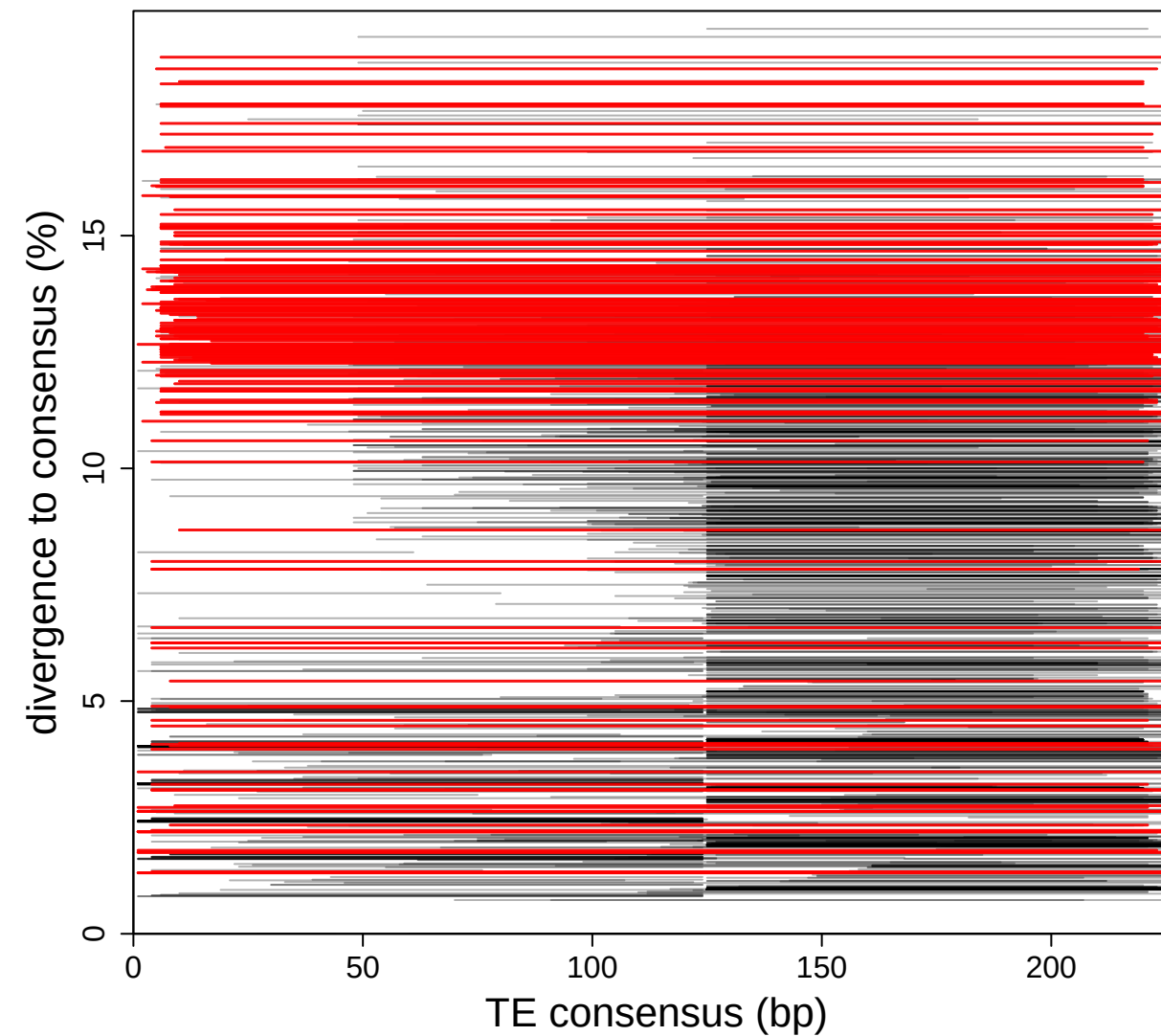
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 677bp; fragments: 1834; full length: 0 (≥ 609.3 bp)

After TEtrimmer Extended plot Blue lines are boundaries

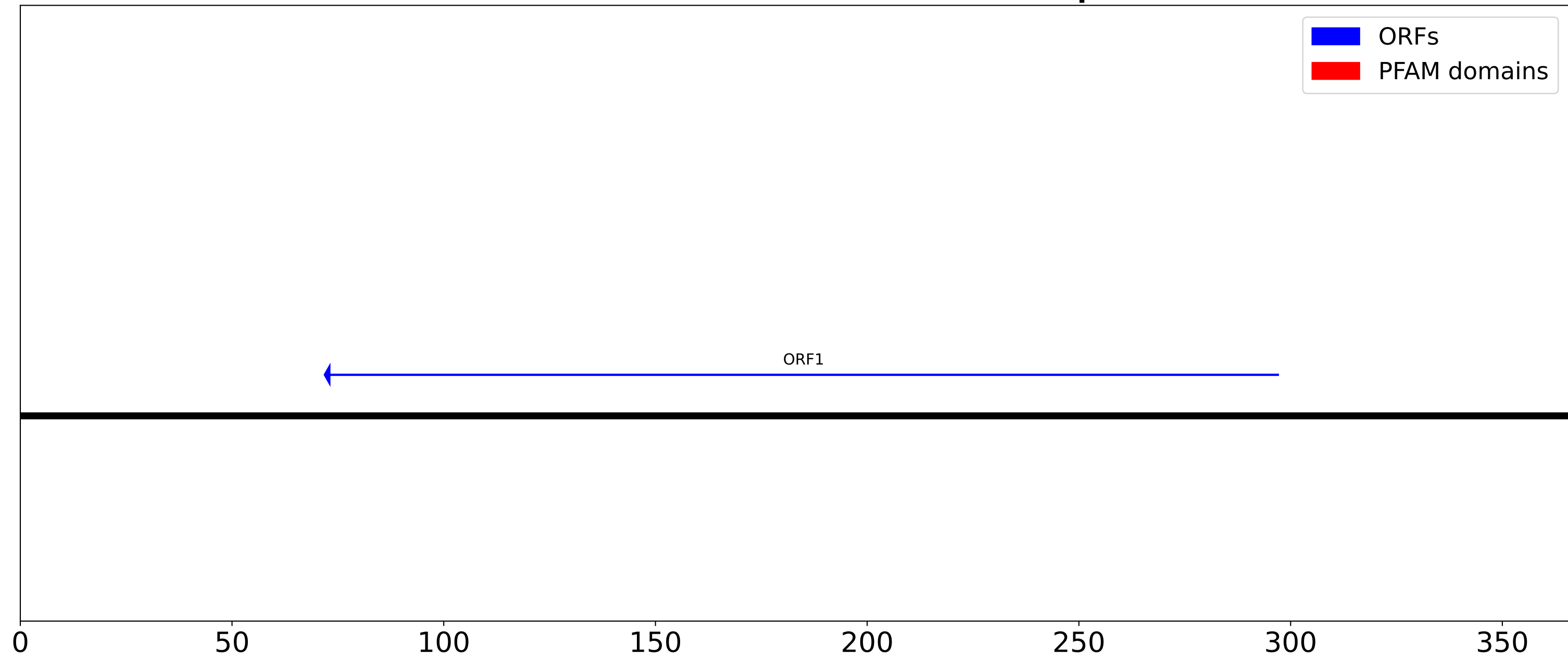


TE: rnd_1_family_700
size: 228bp; fragments: 1788; full length: 182 (≥ 205.2 bp)

Before TEtrimmer 228 bp



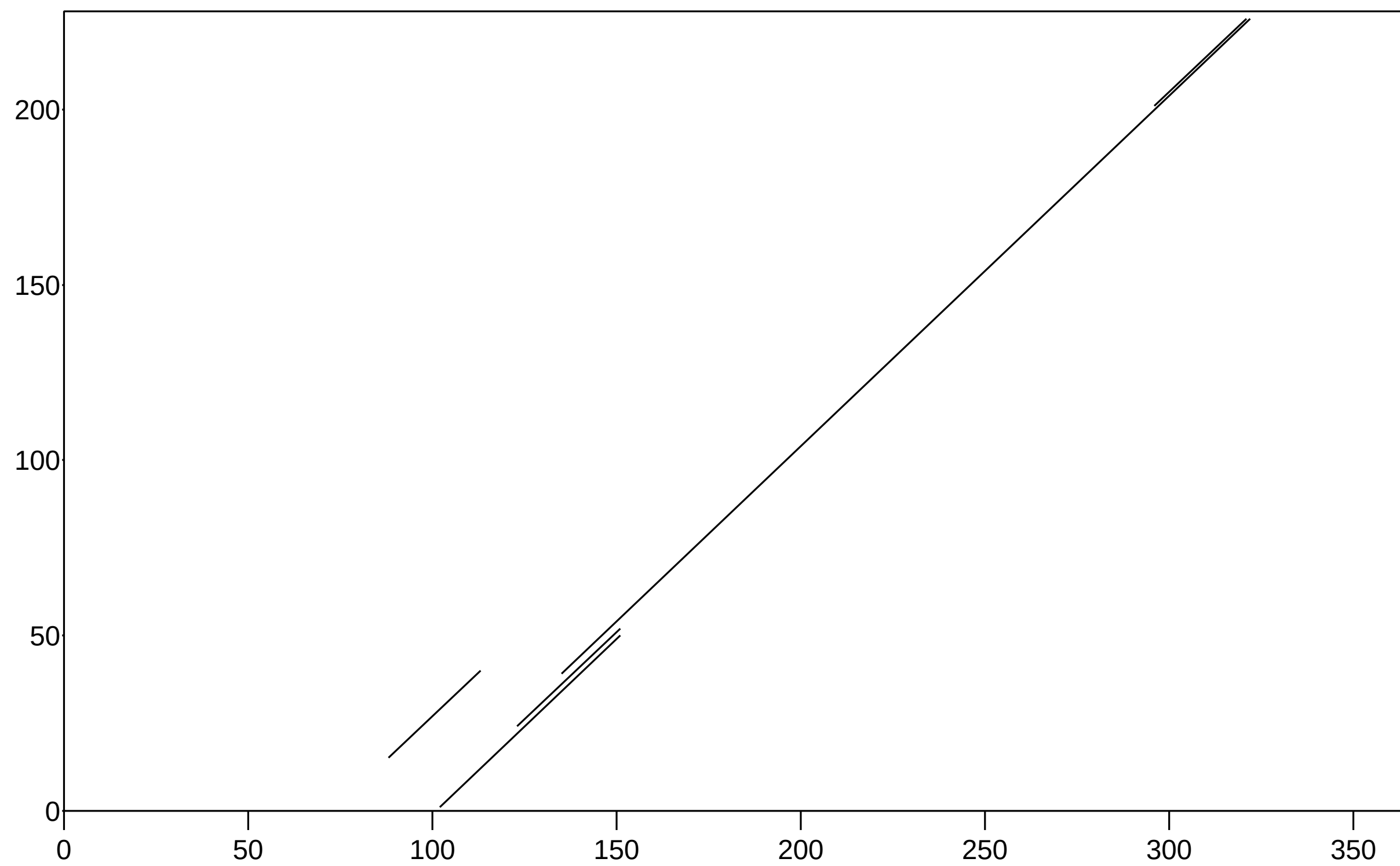
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)