

Start crop Point



End crop Point

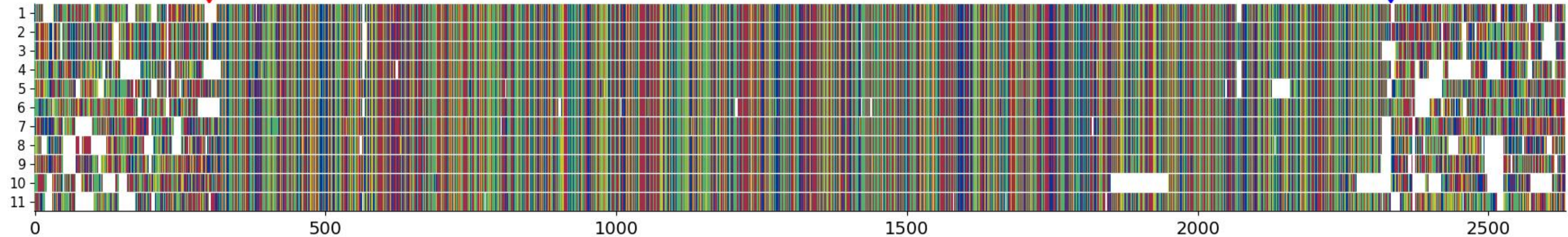


MSA length = 2033



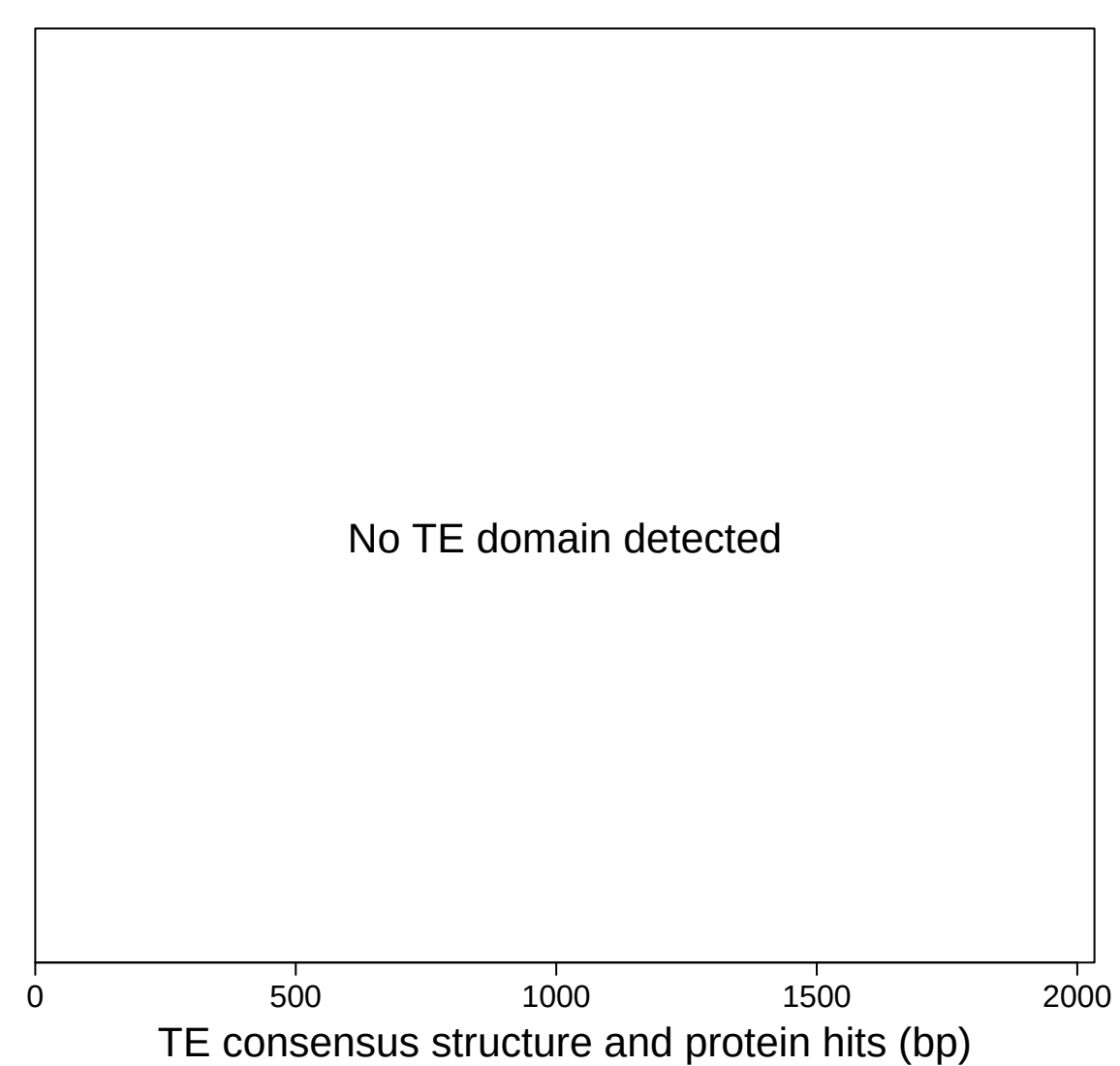
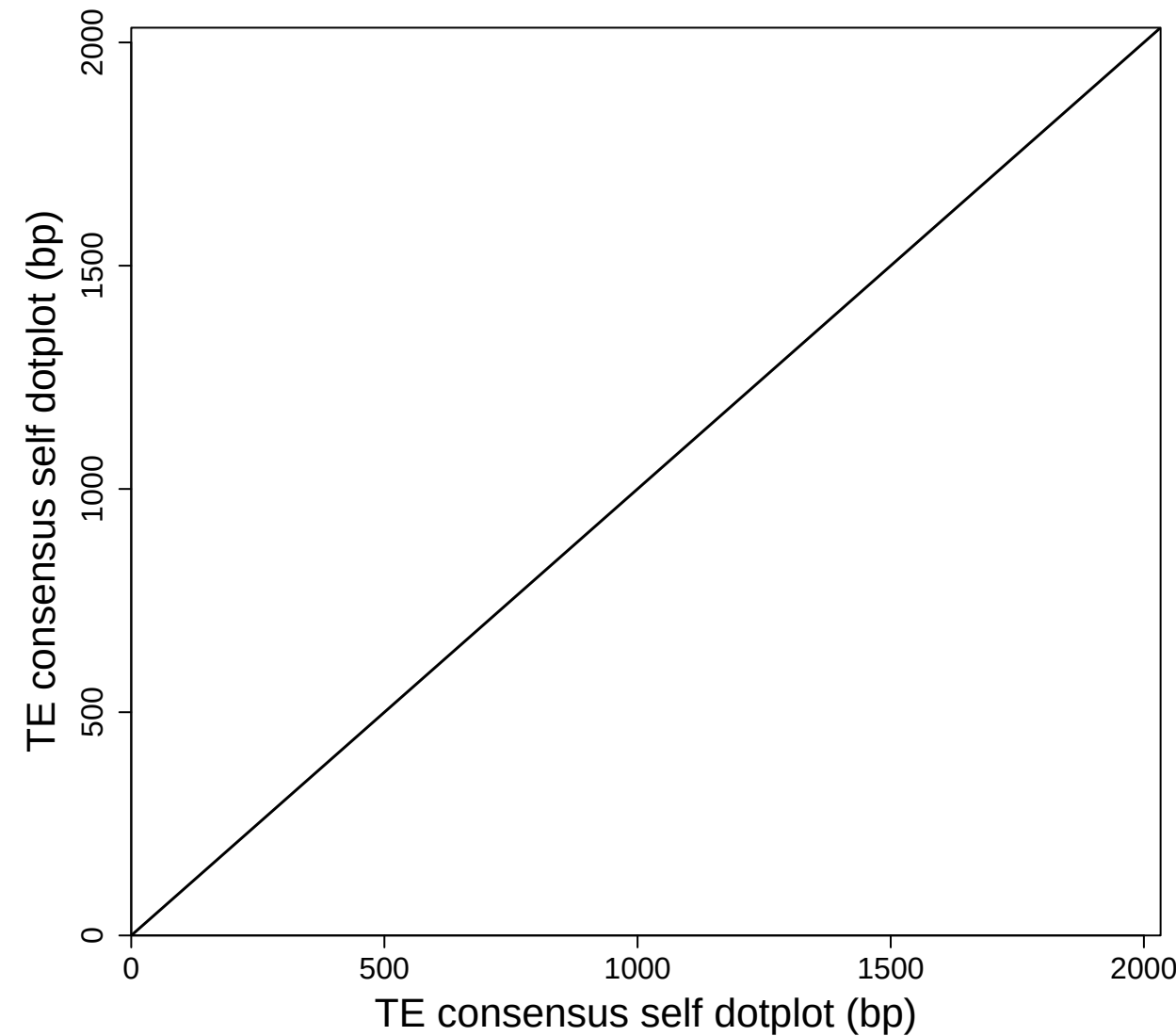
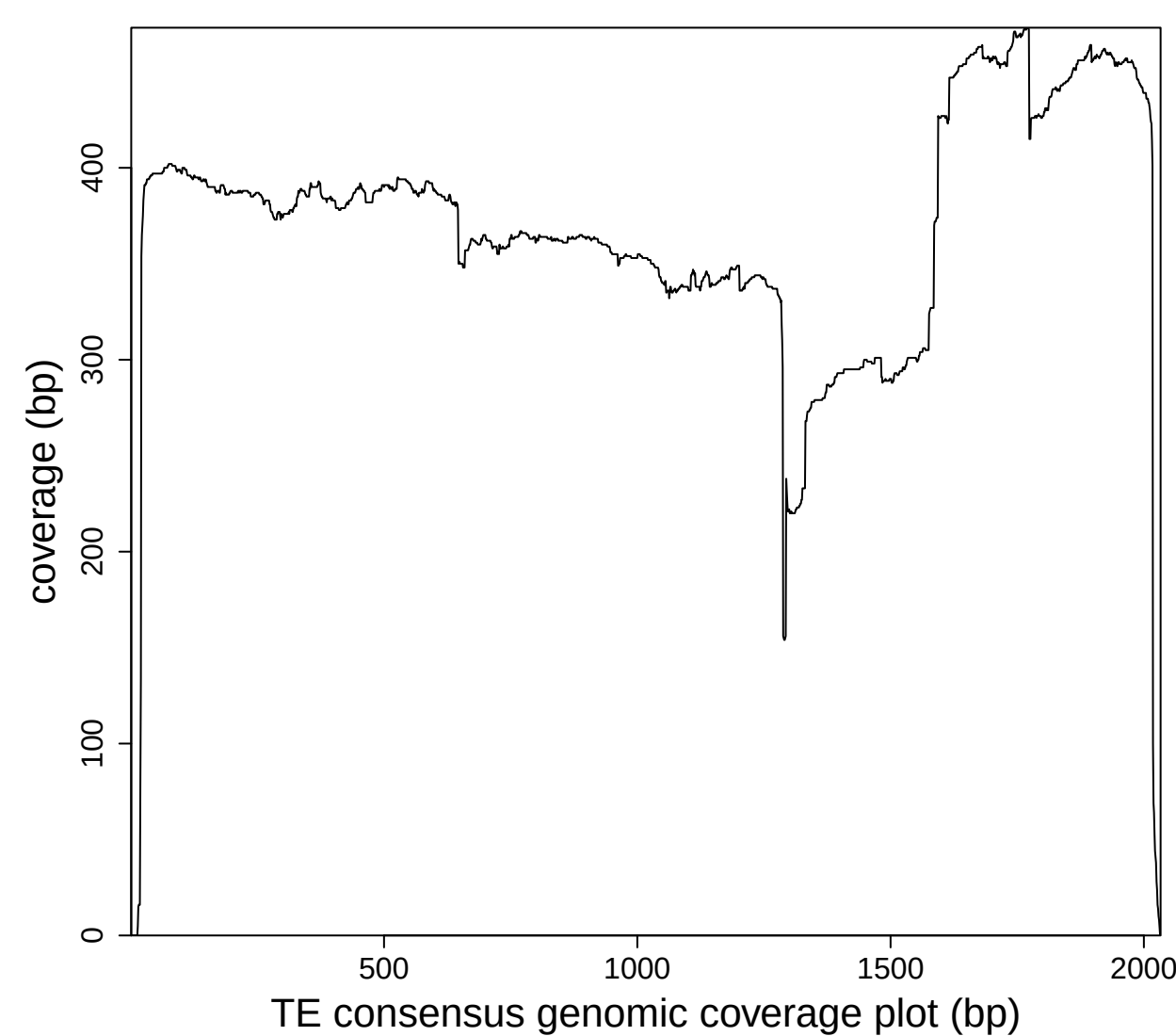
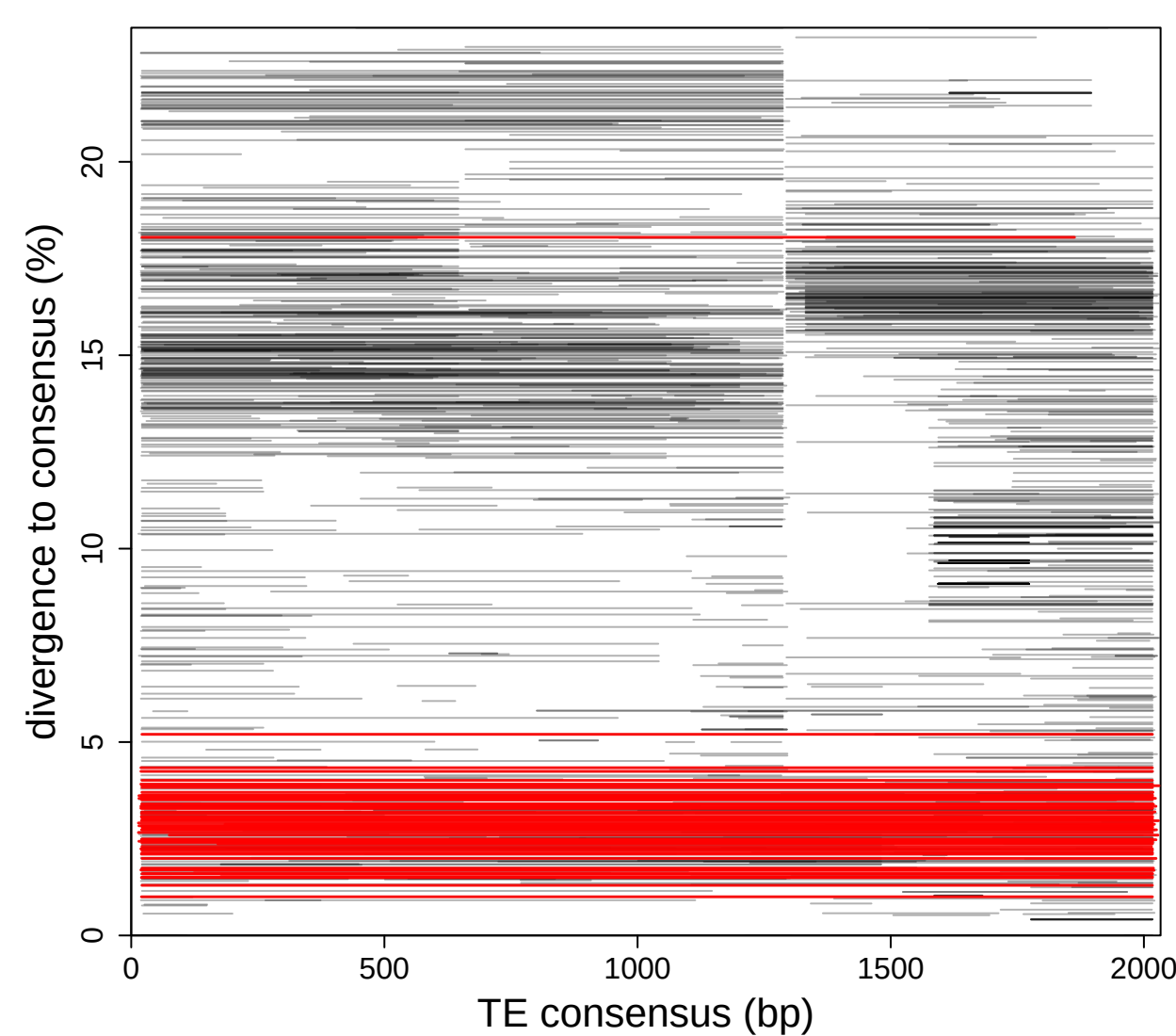
Start crop Point

End crop Point



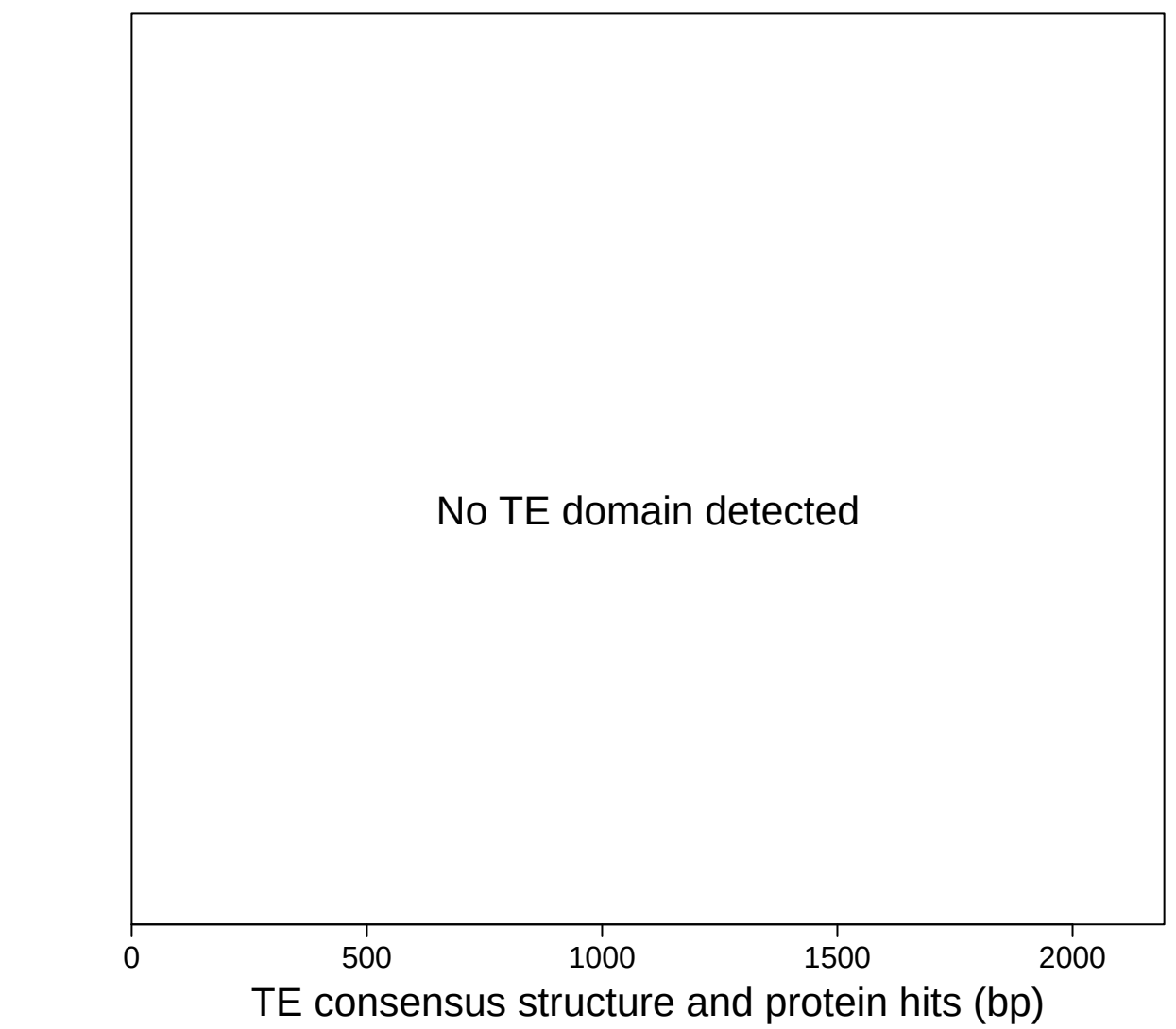
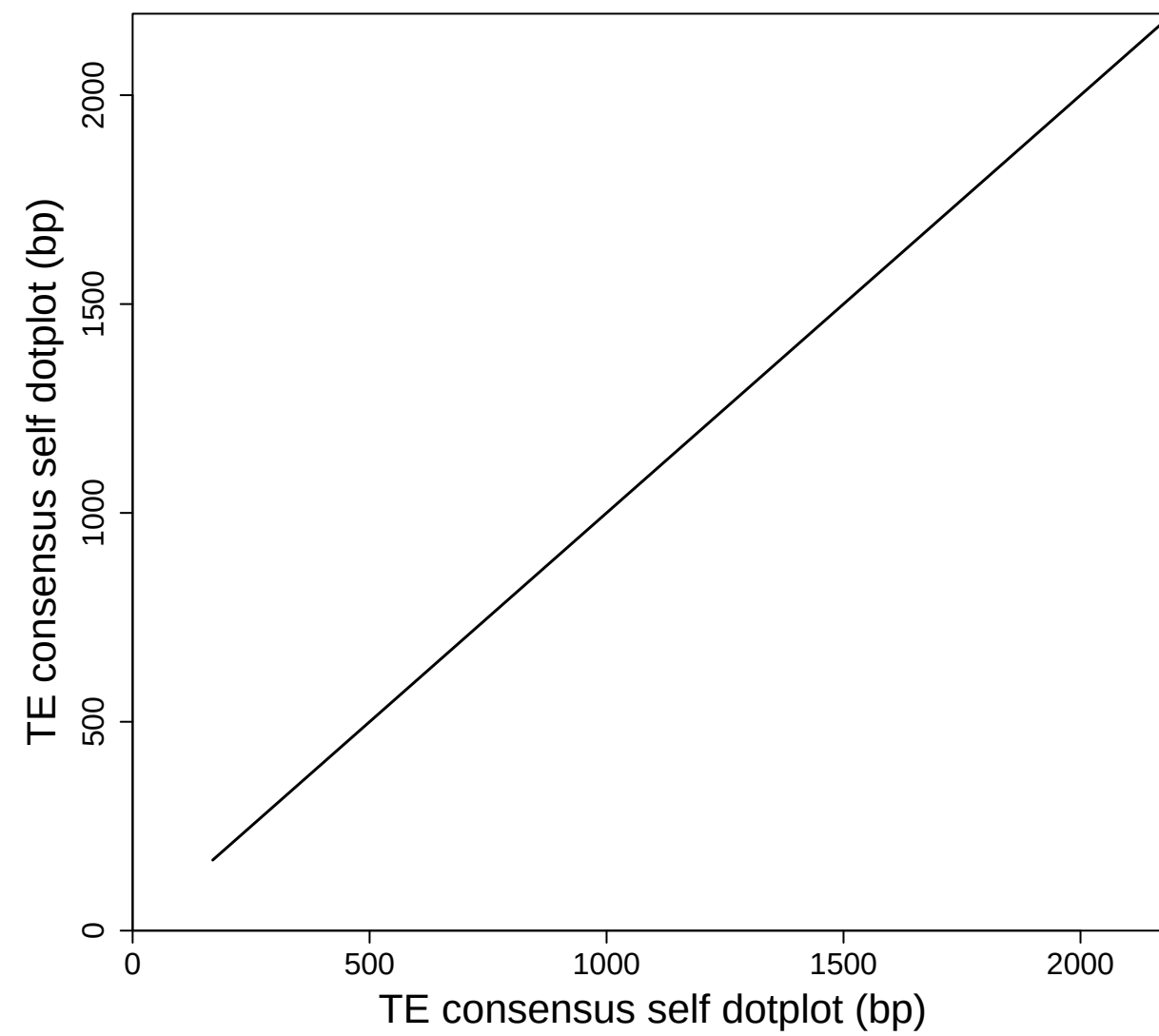
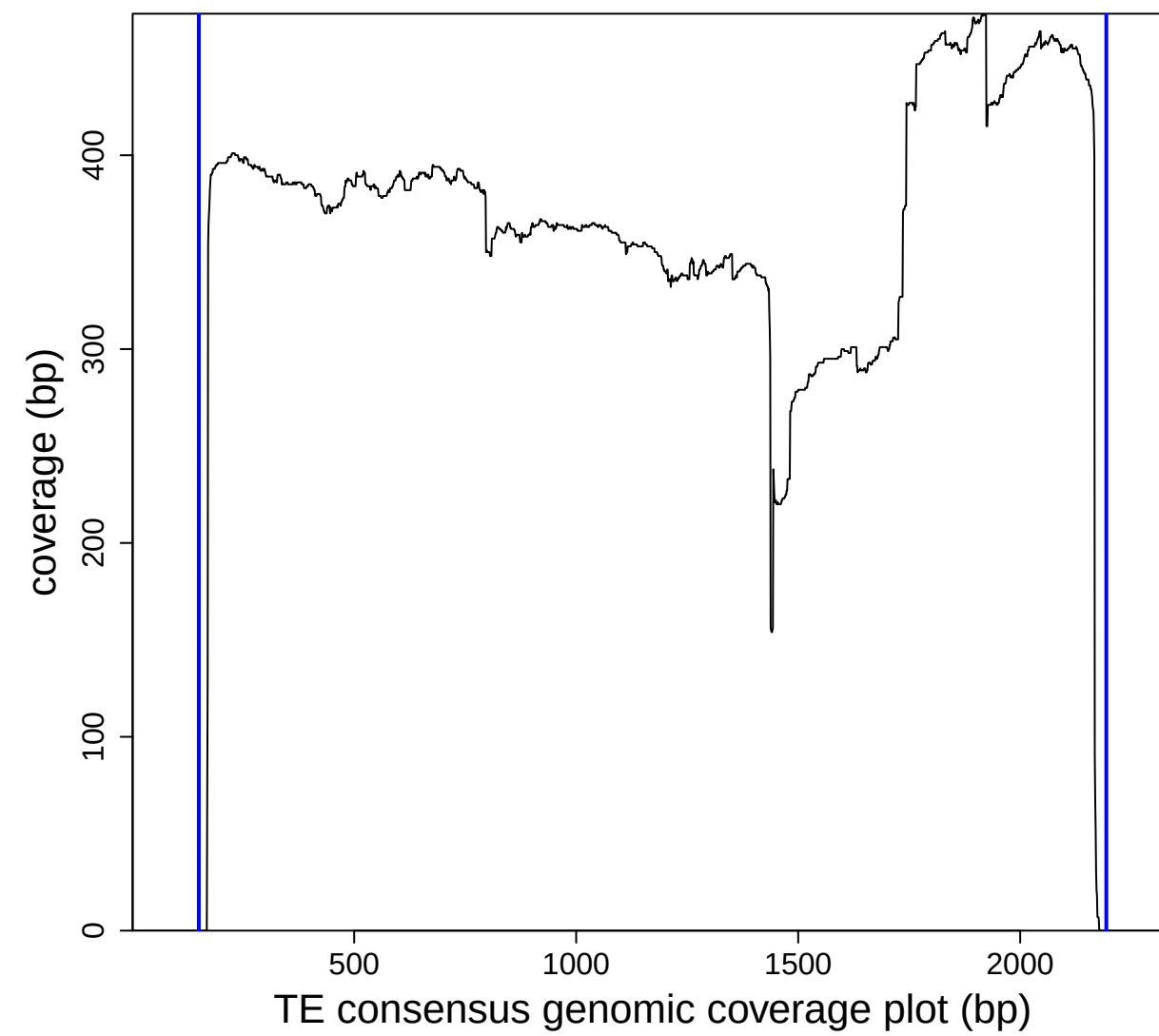
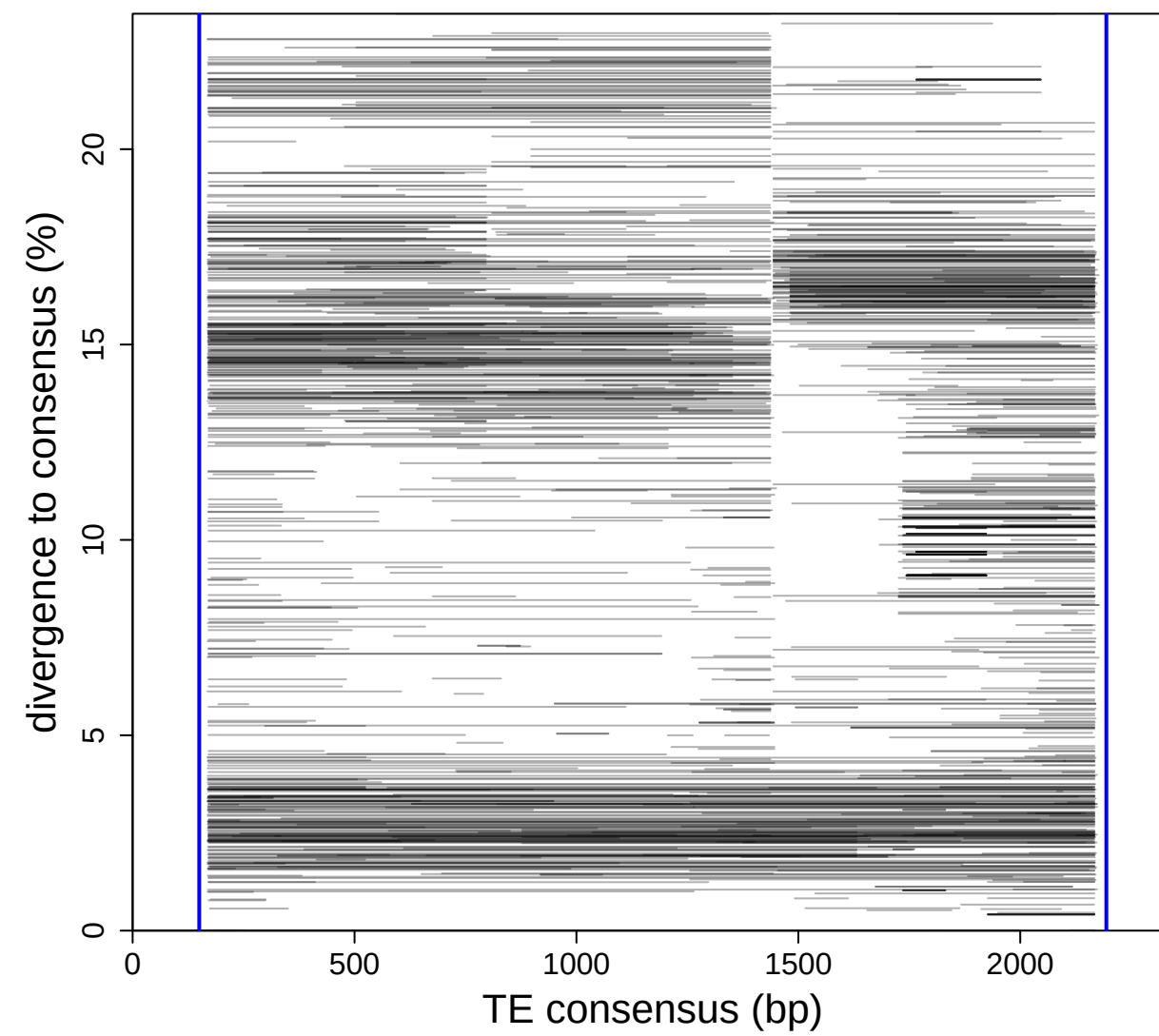
_4.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_al
size: 2033bp; fragments: 1189; full length: 65 (≥ 1829.7 bp)

After TEtrimmer 2033 bp



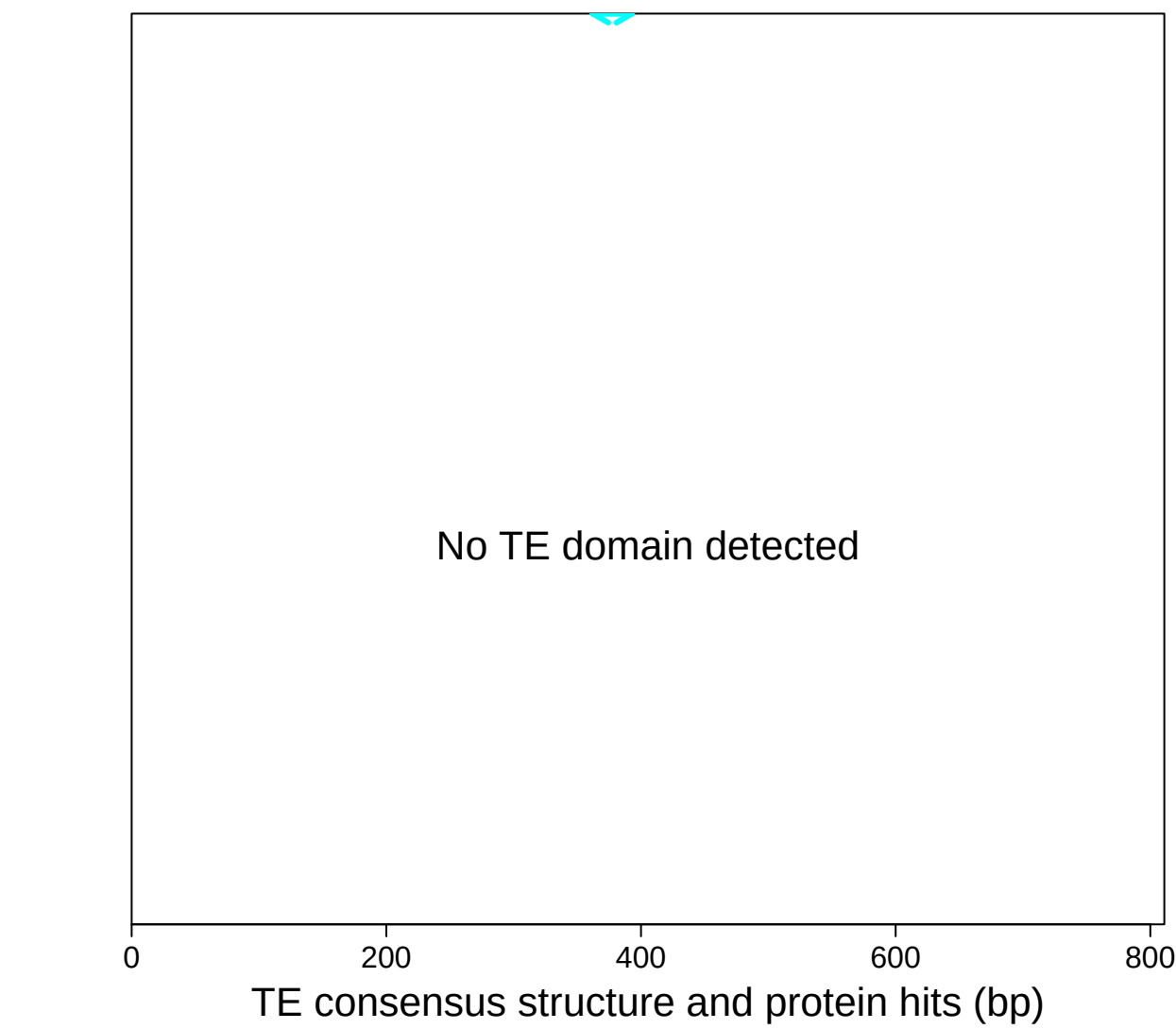
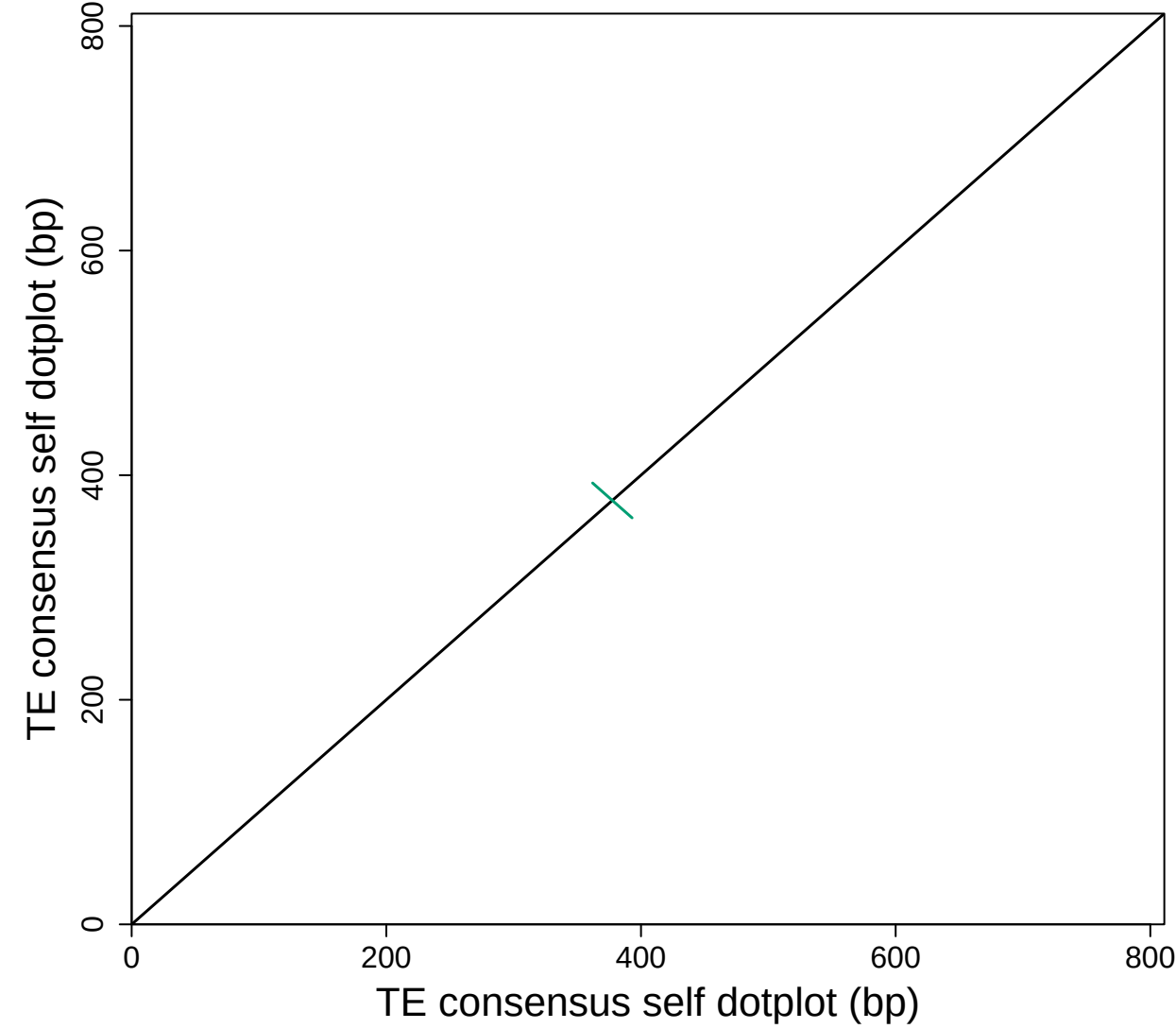
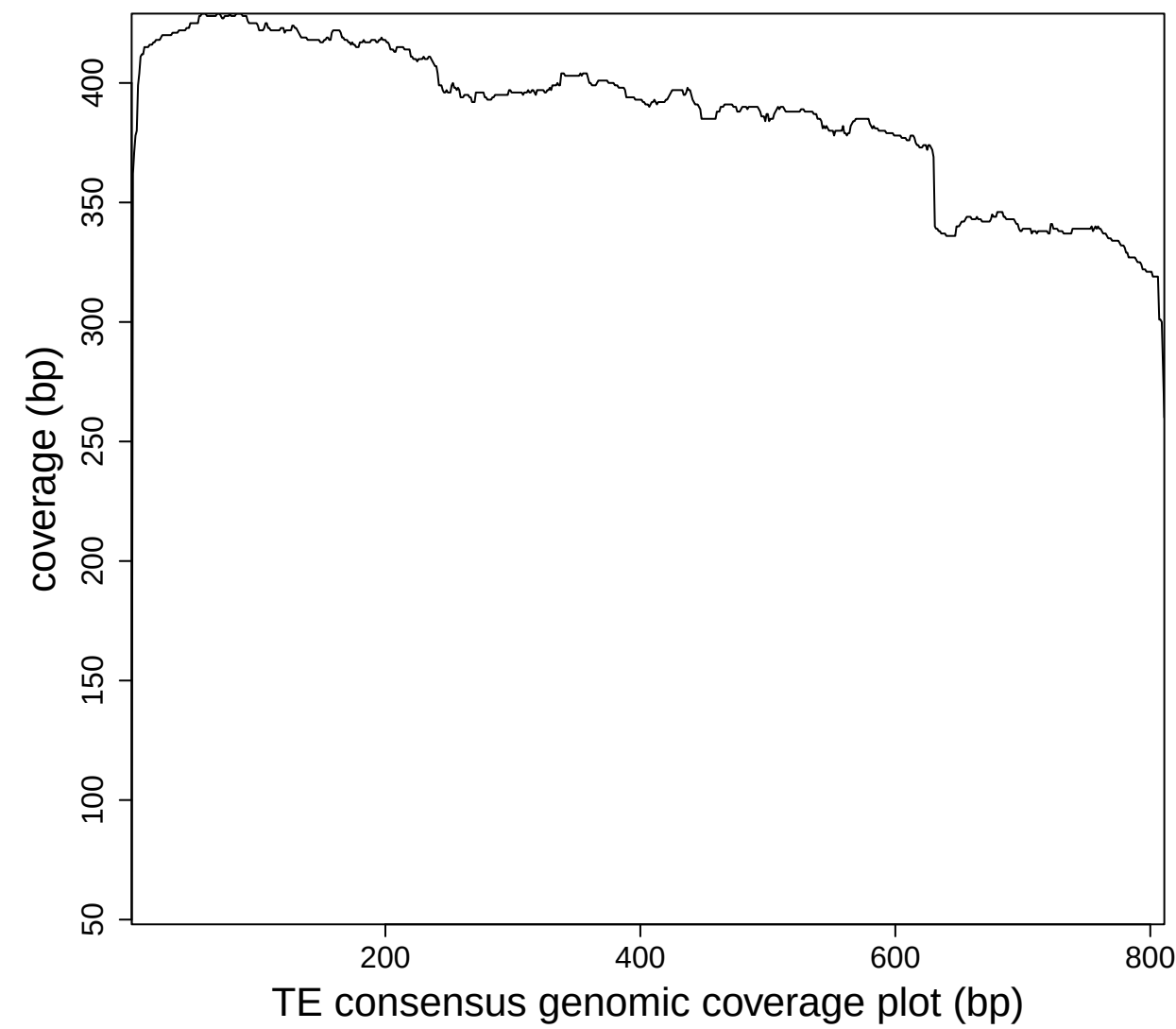
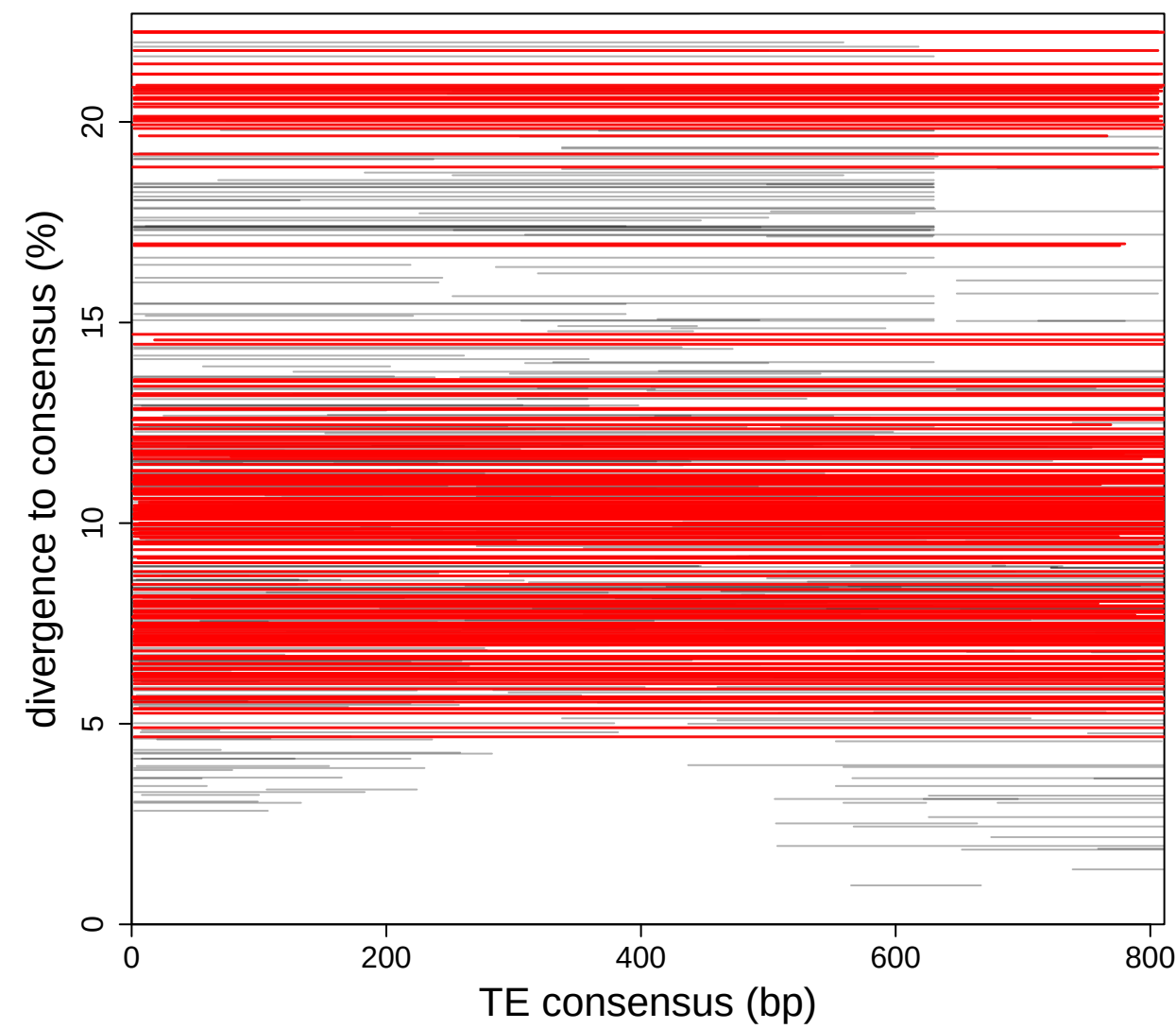
bed_g_4.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 2344bp; fragments: 1188; full length: 0 (>=2109.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

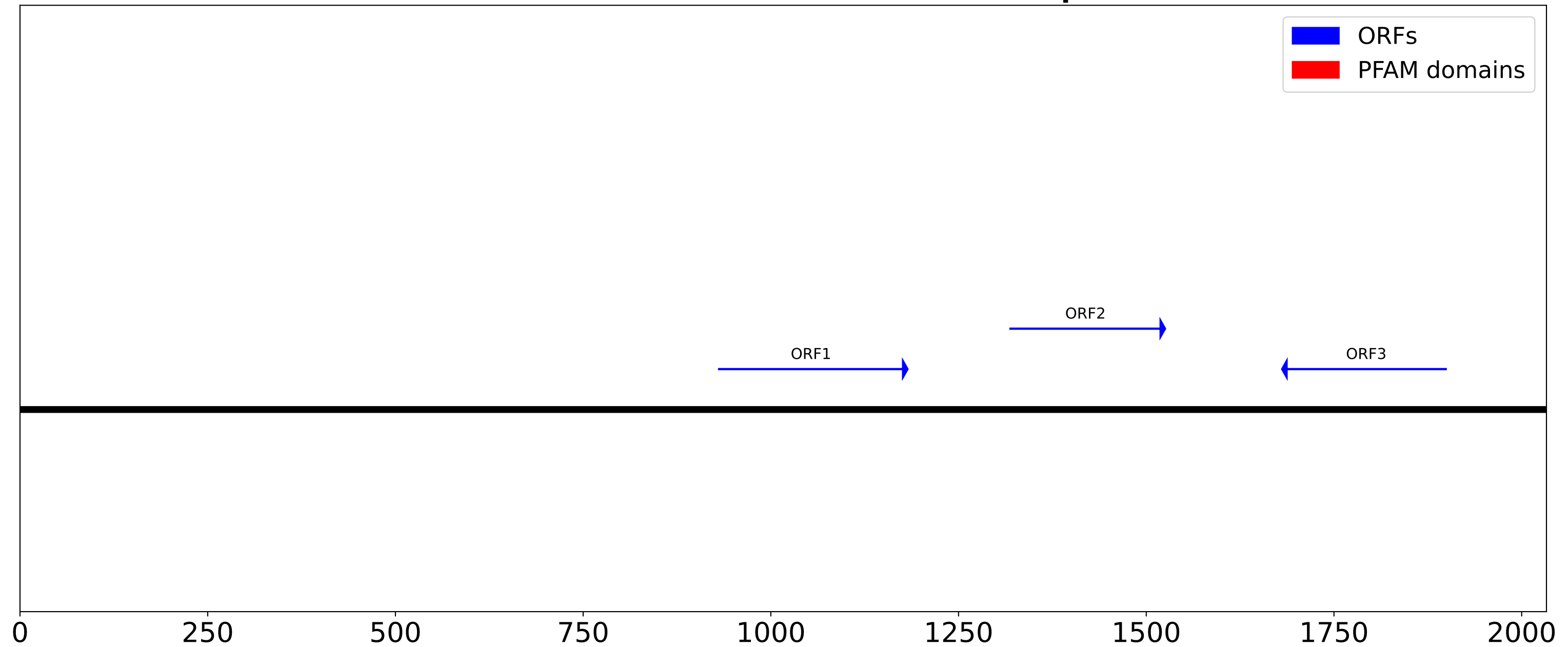


TE: rnd_1_family_71
size: 811bp; fragments: 602; full length: 228 (≥ 729.9 bp)

Before TEtrimmer 811 bp



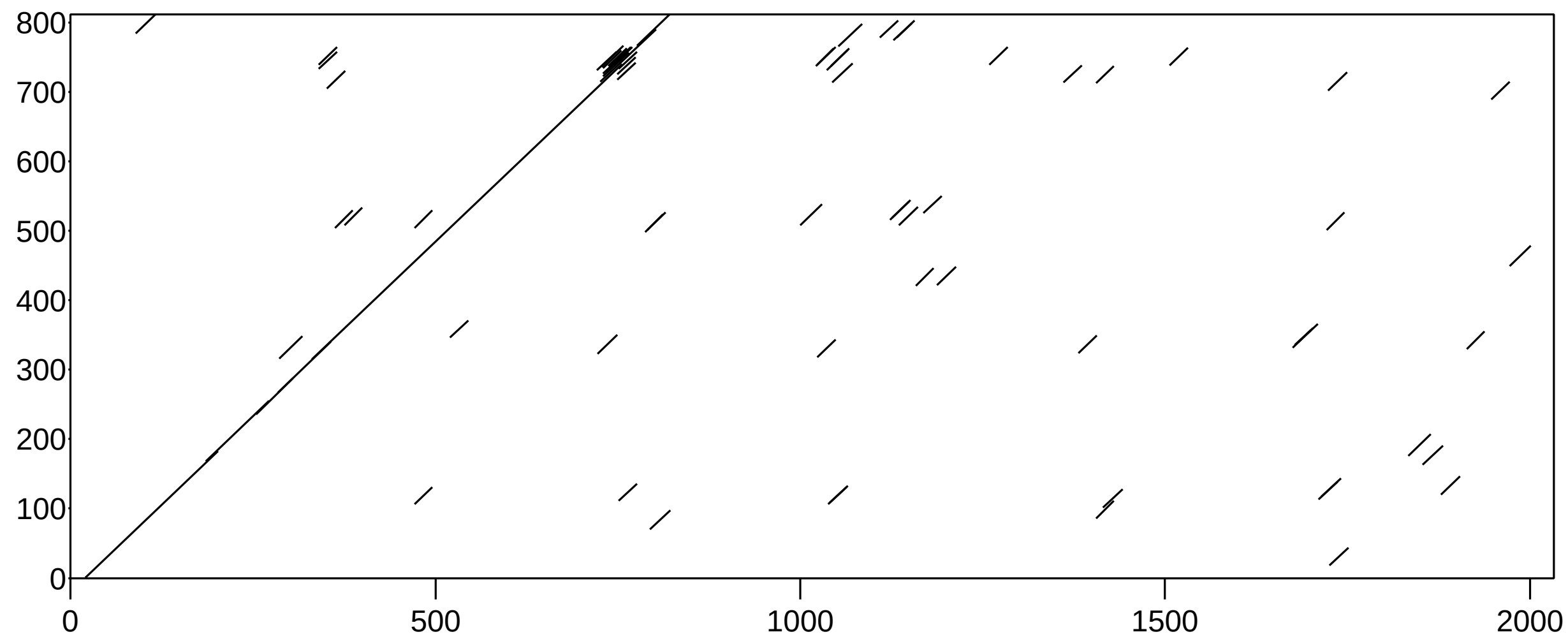
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)