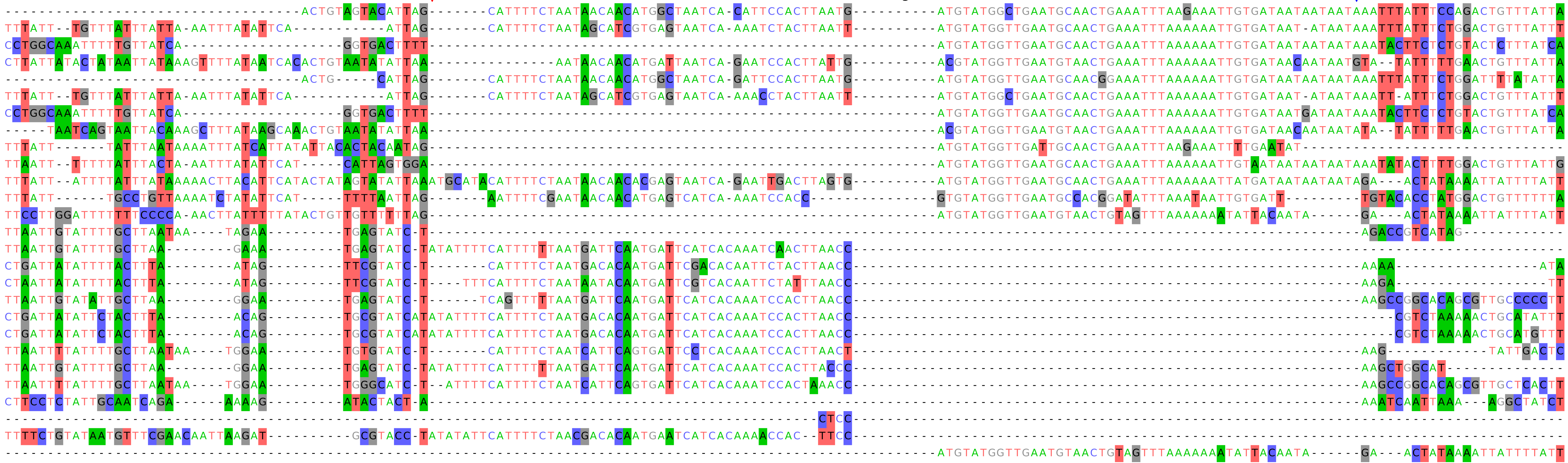


Start crop Point

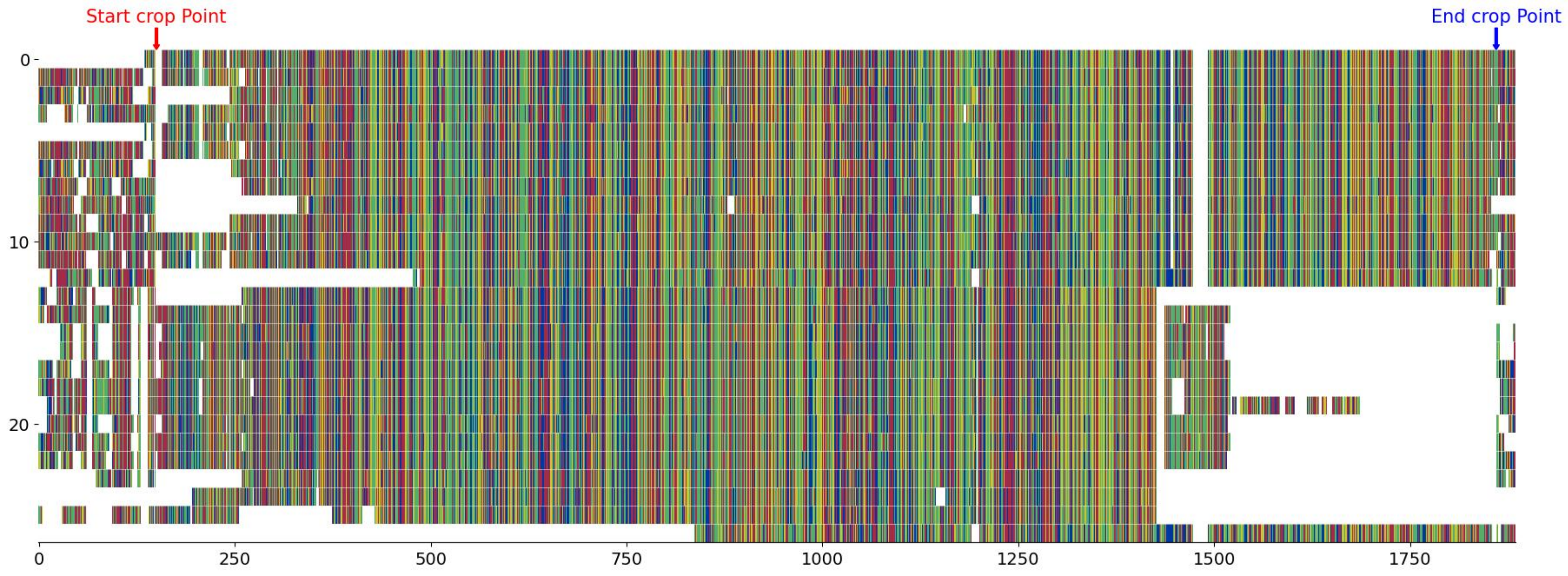
End crop Point

MSA length = 1711



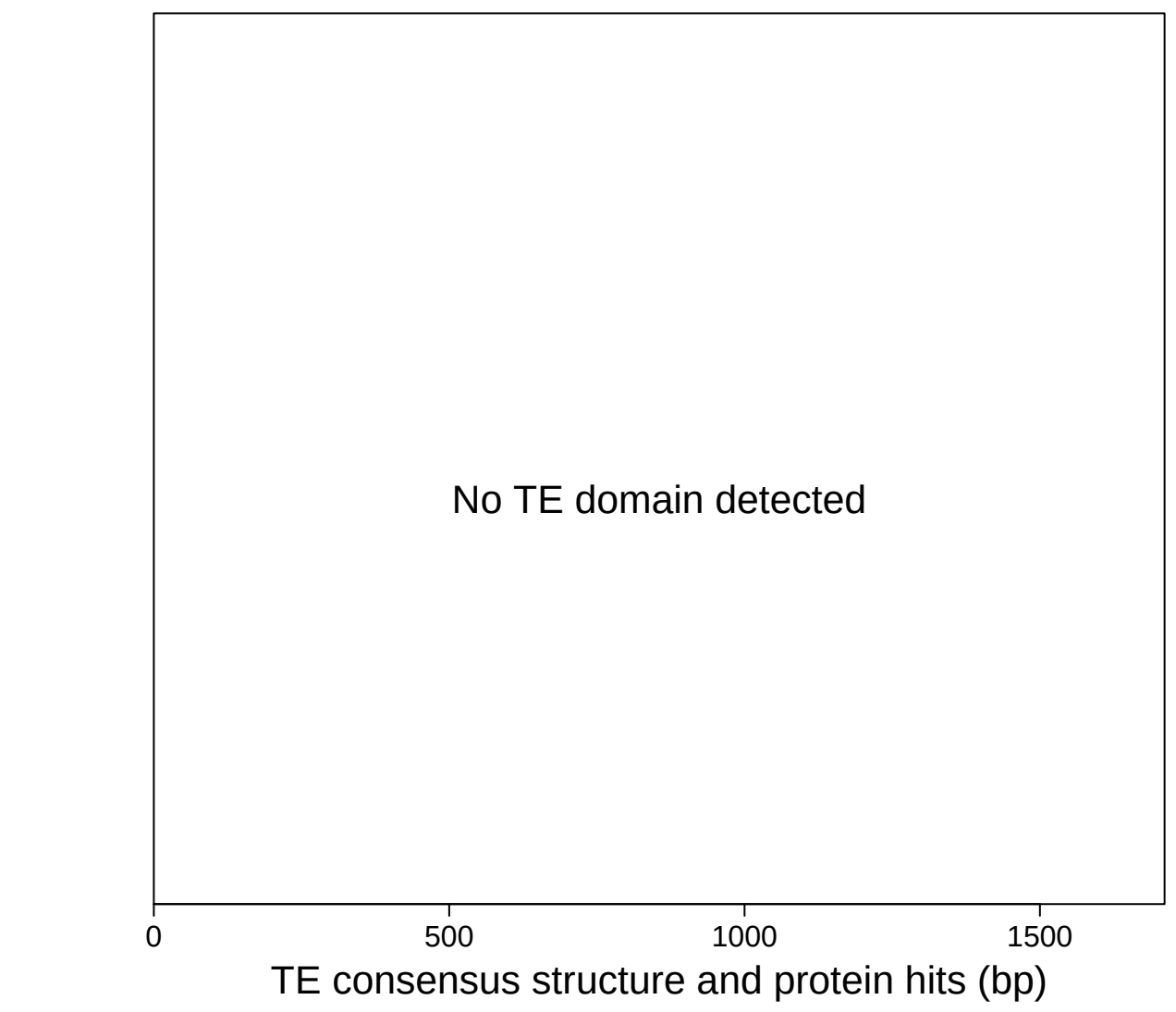
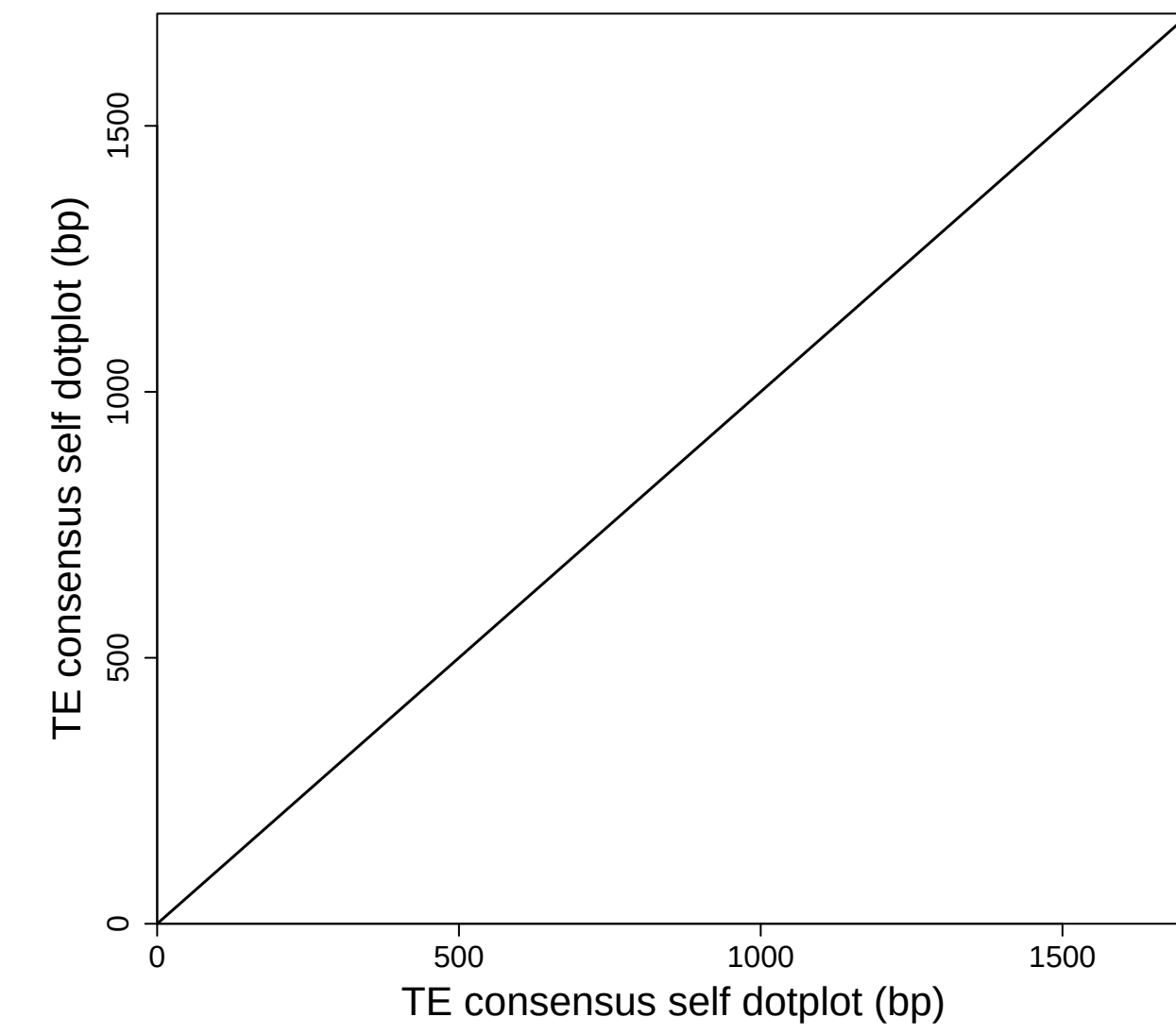
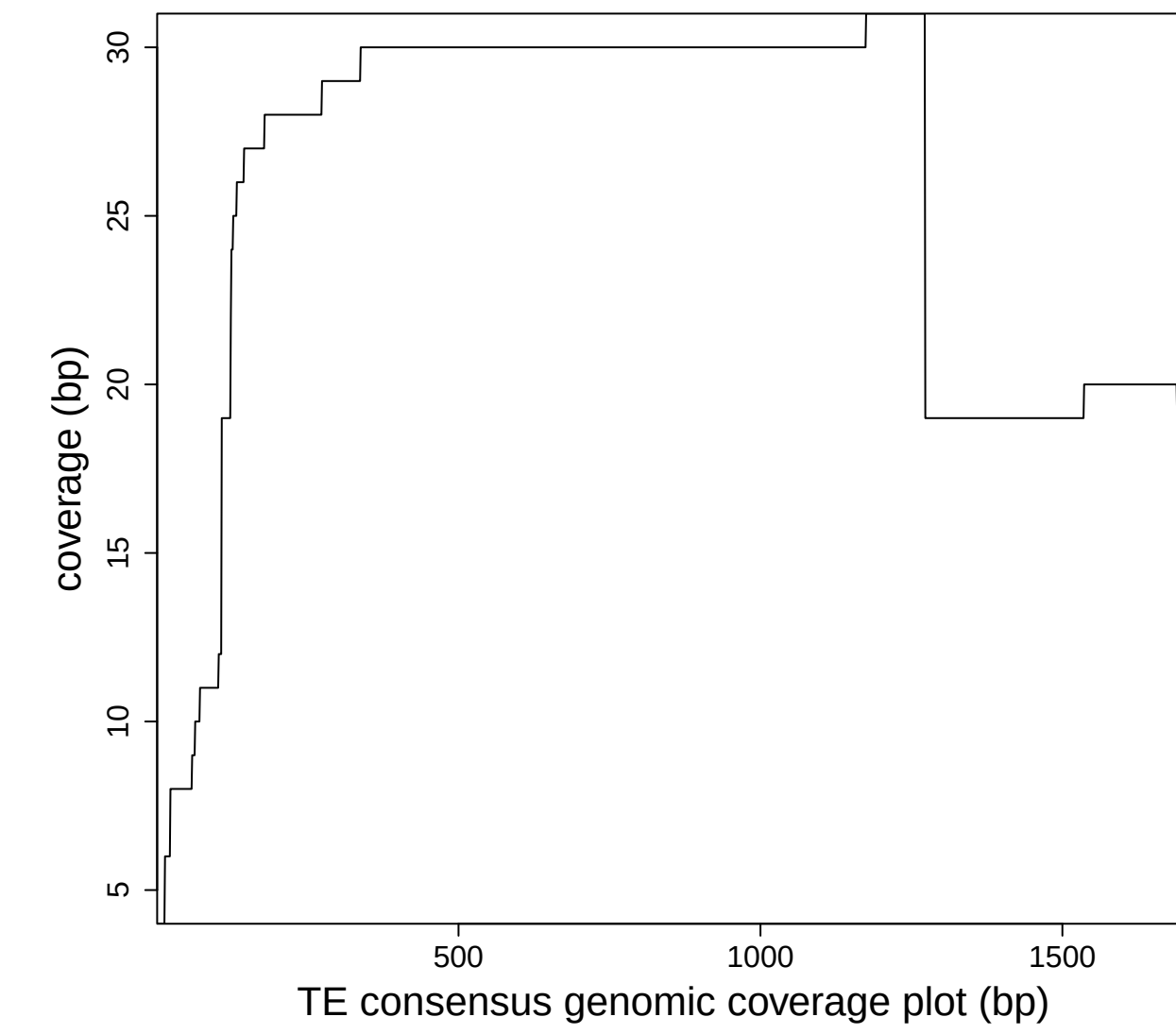
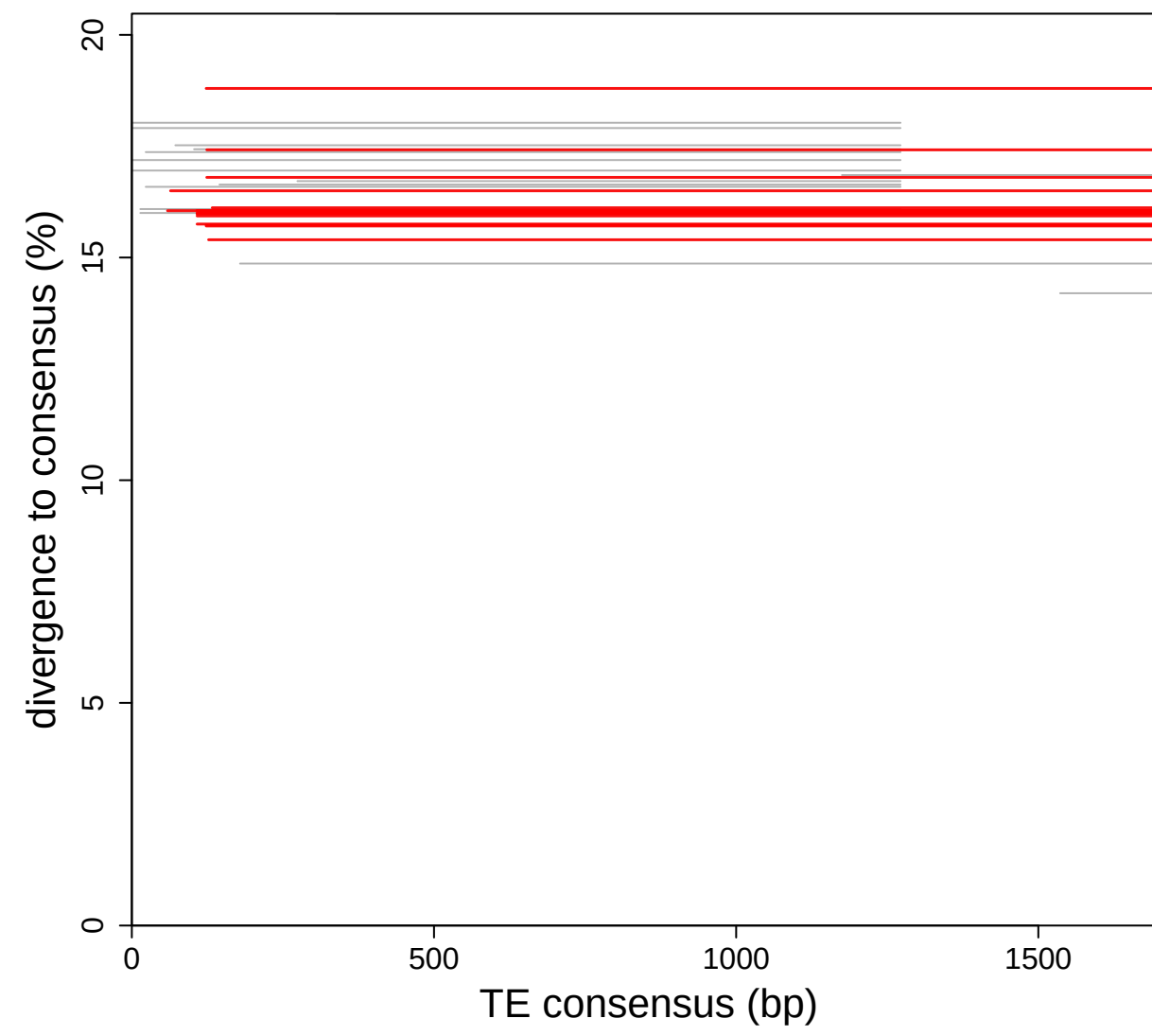
Start crop Point

End crop Point



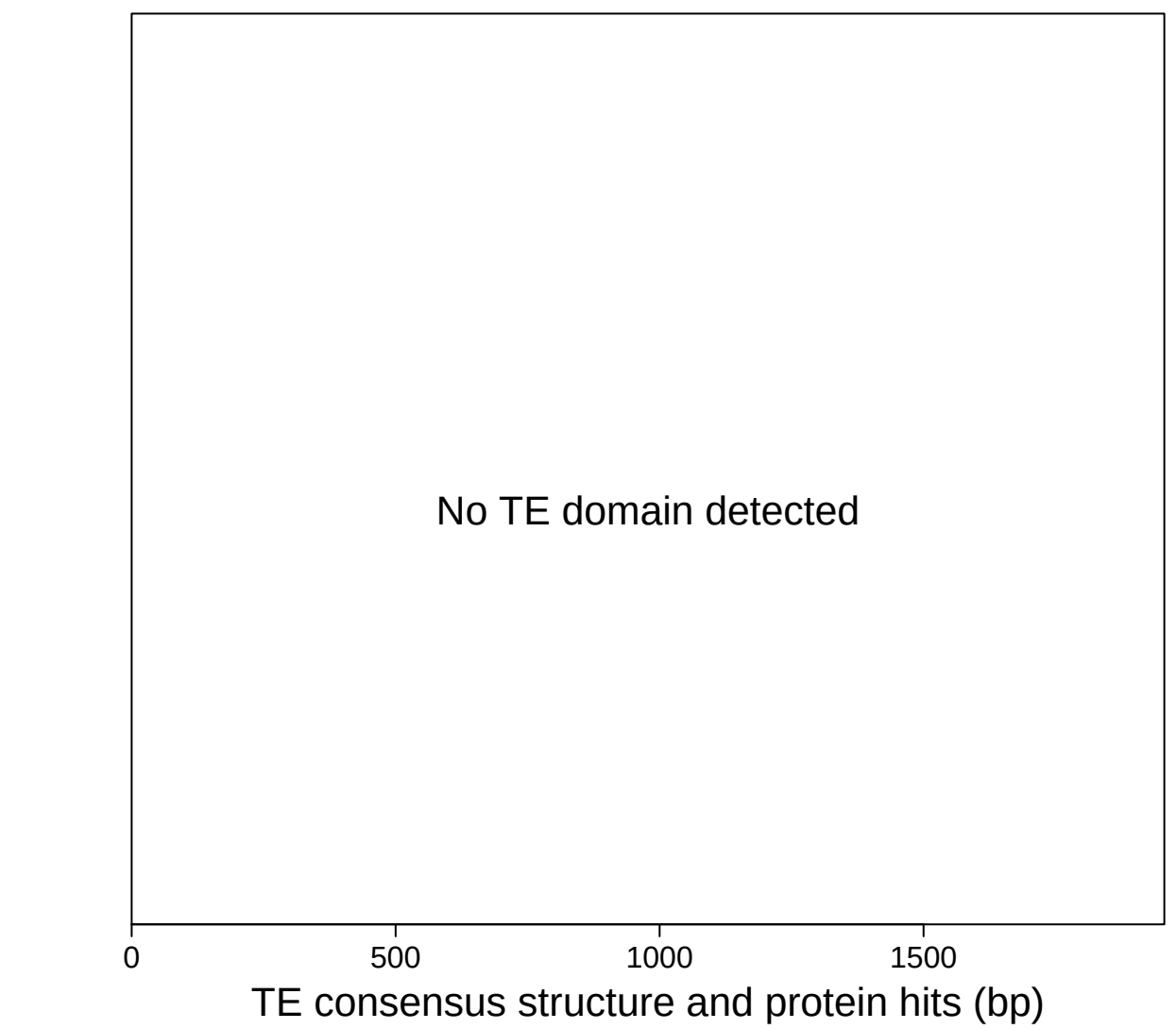
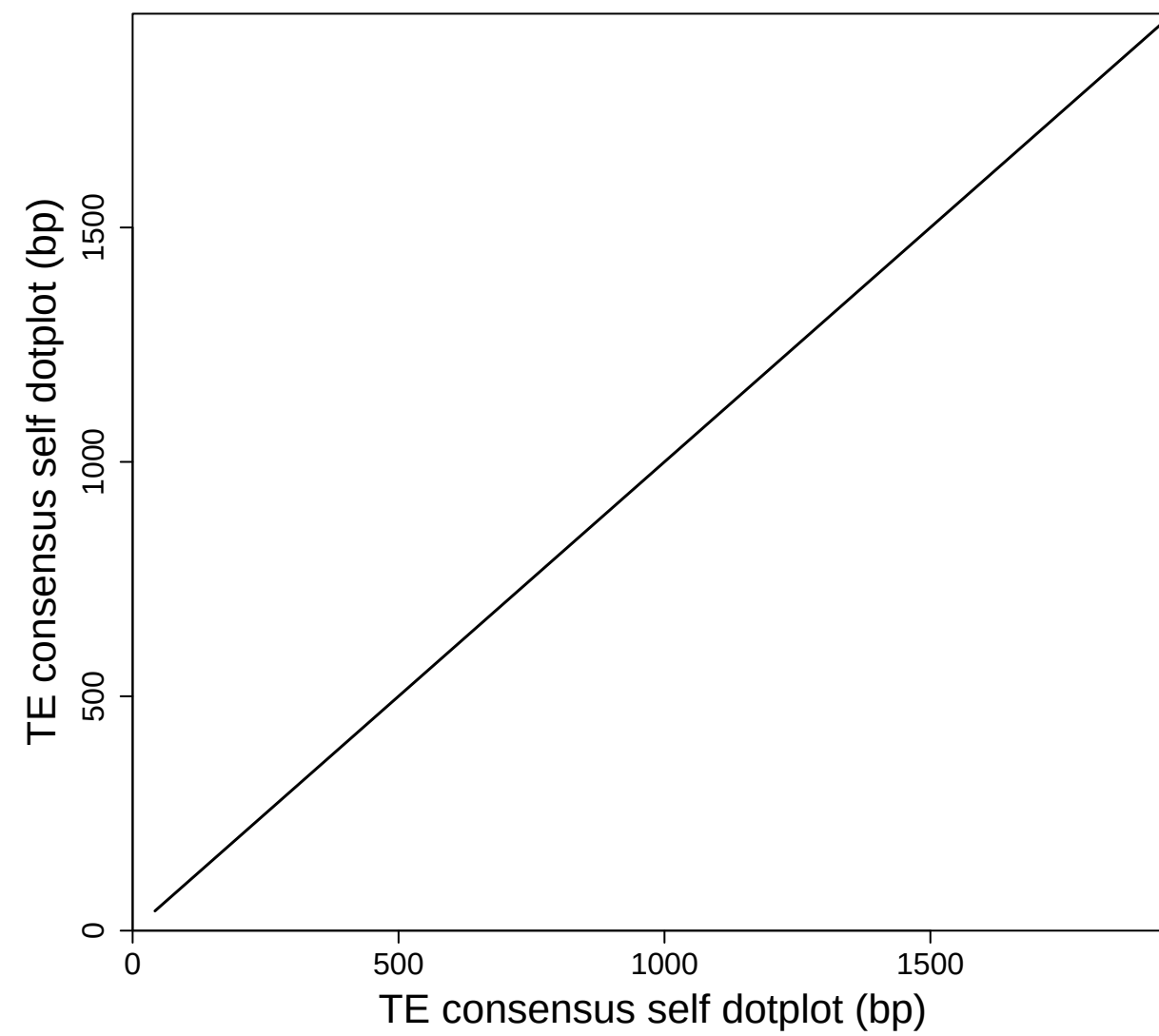
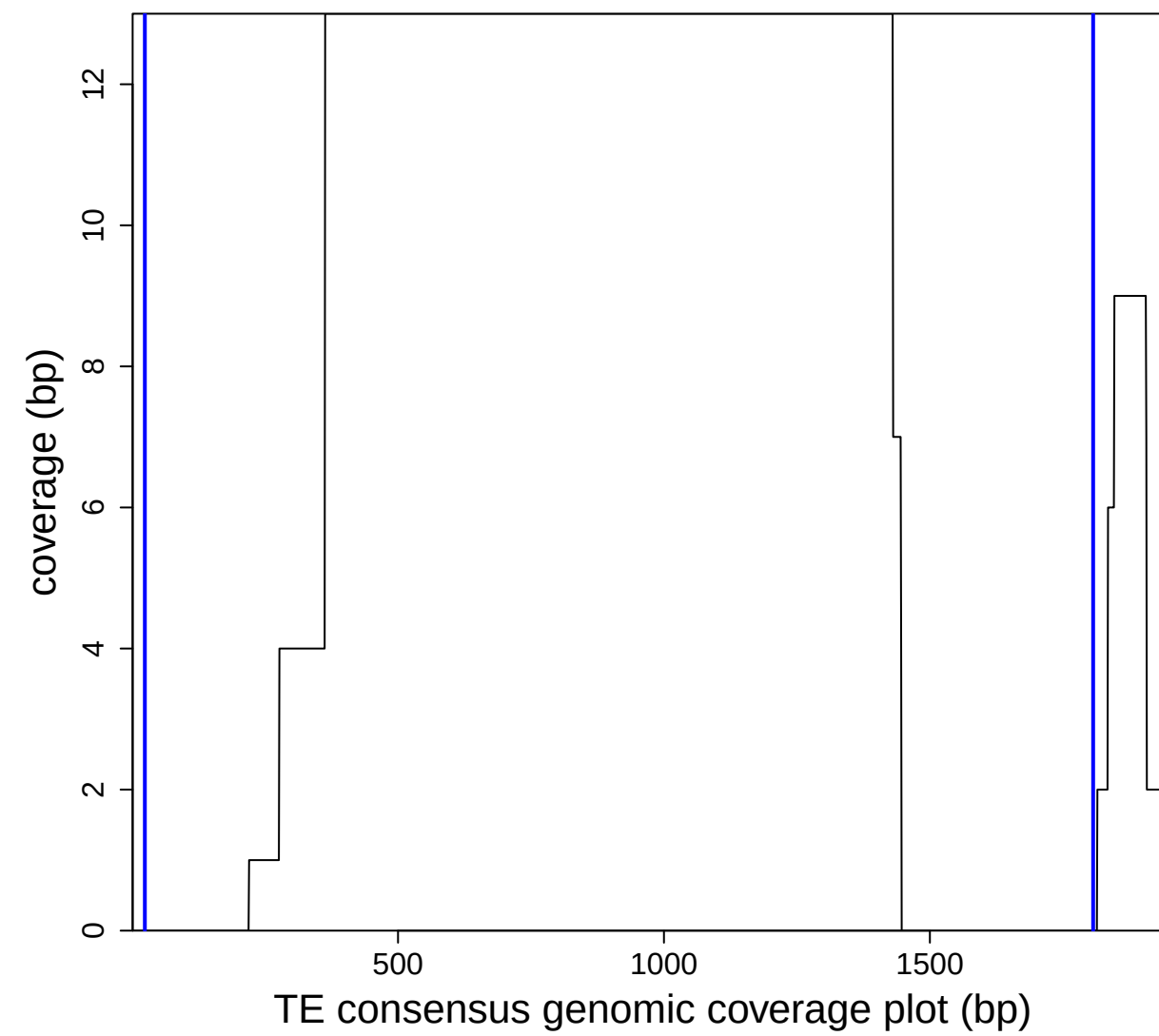
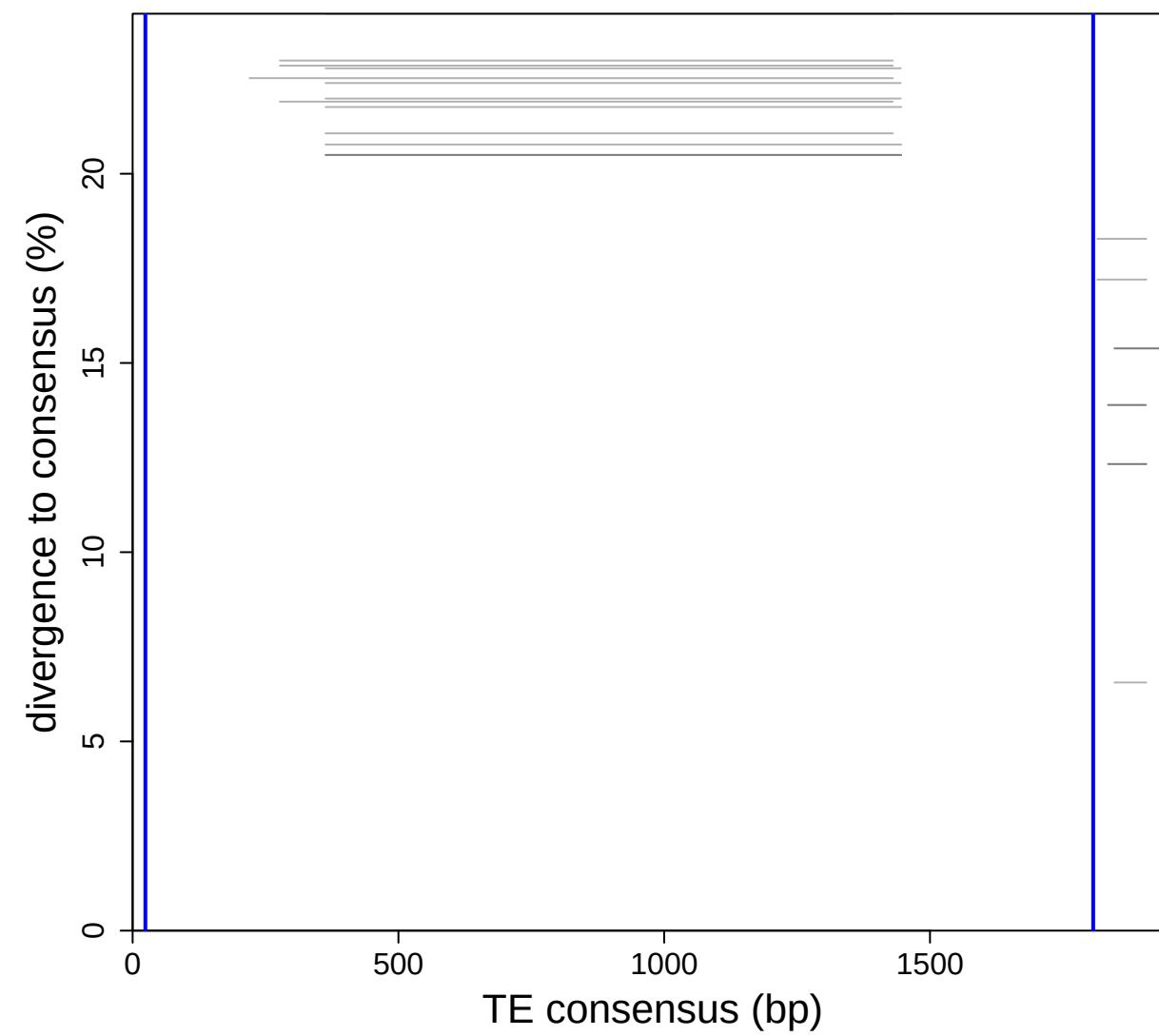
ta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.f
size: 1711bp; fragments: 32; full length: 16 (≥ 1539.9 bp)

After TEtrimmer 1711 bp

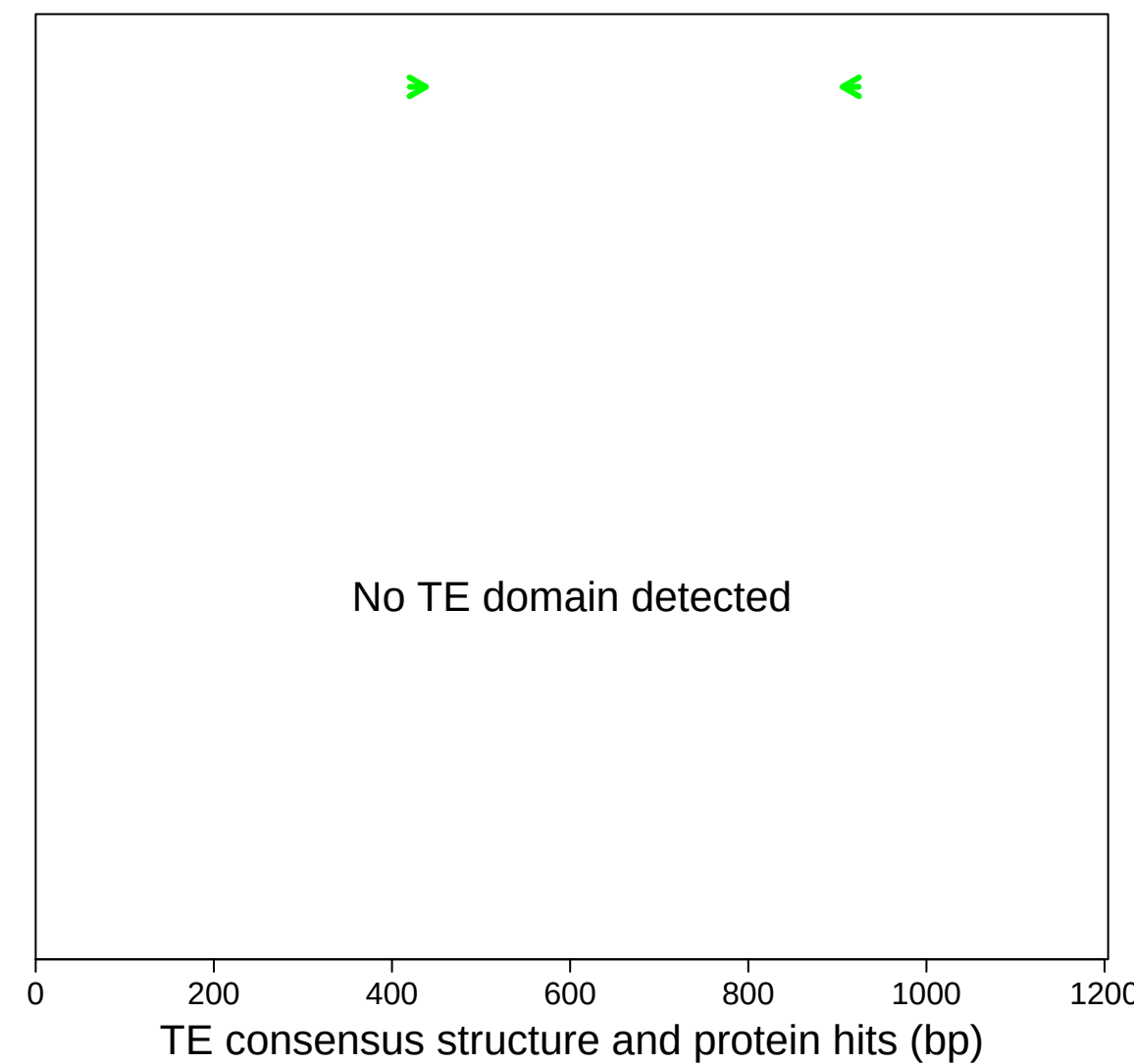
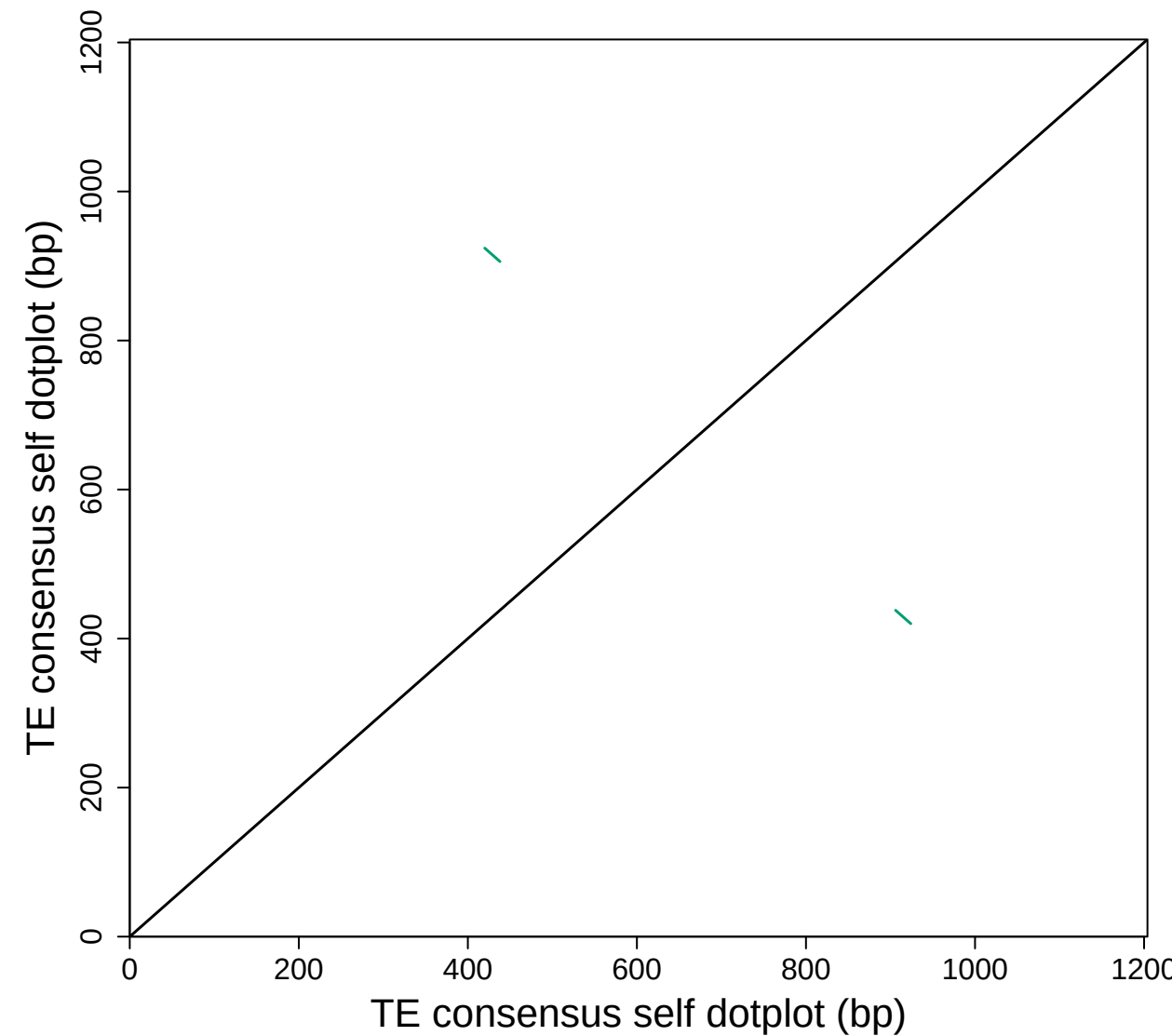
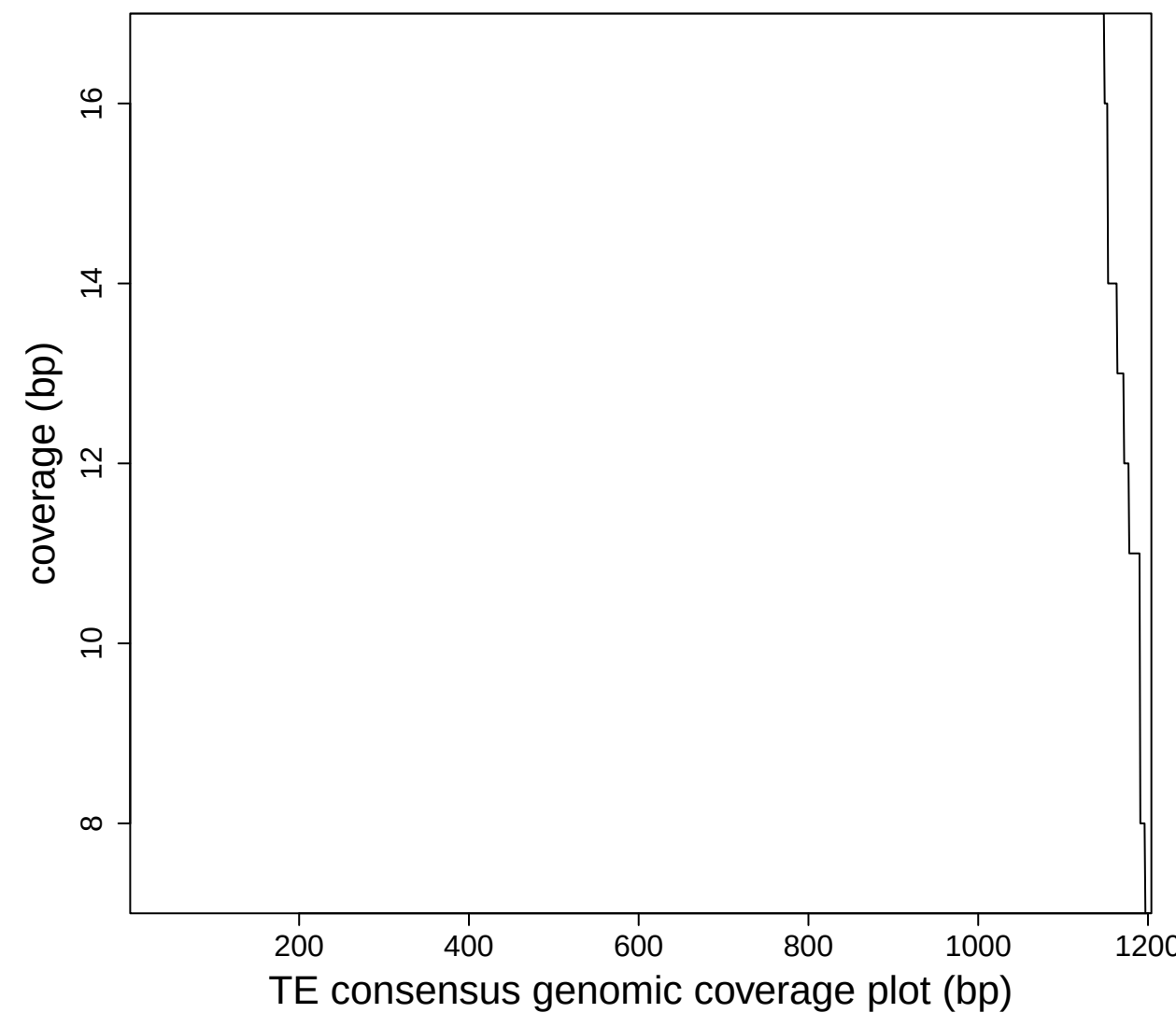
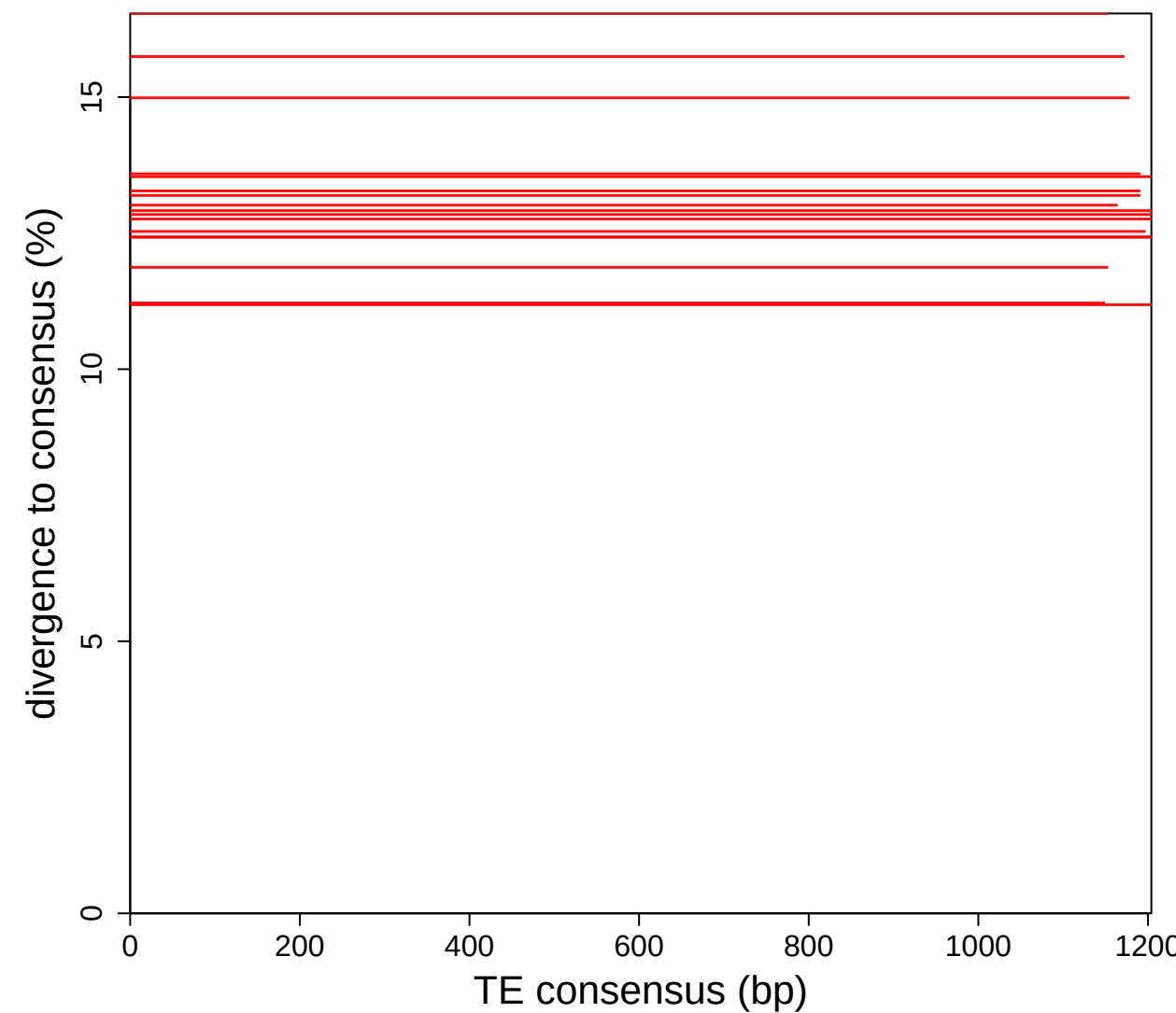


fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_
size: 1957bp; fragments: 22; full length: 0 (>=1761.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

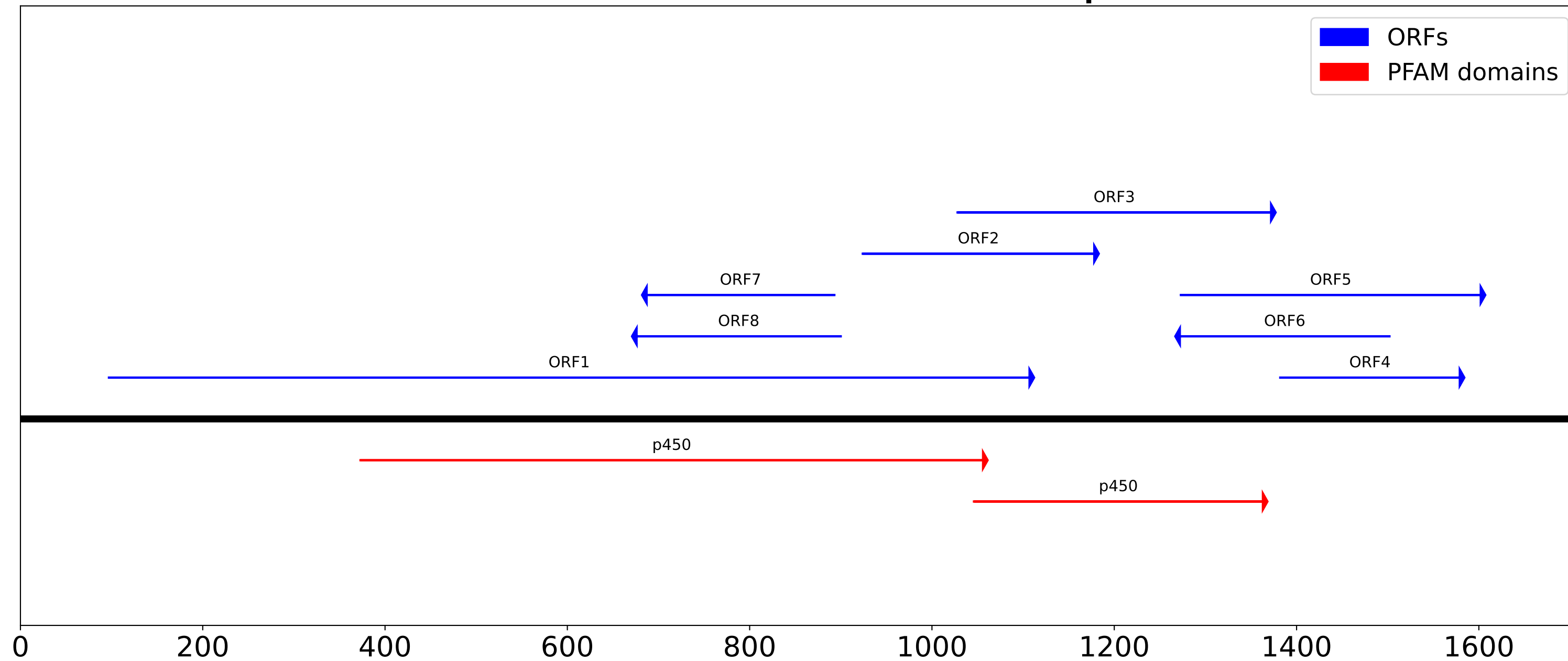


TE: rnd_5_family_12215
size: 1204bp; fragments: 17; full length: 17 (≥ 1083.6 bp)

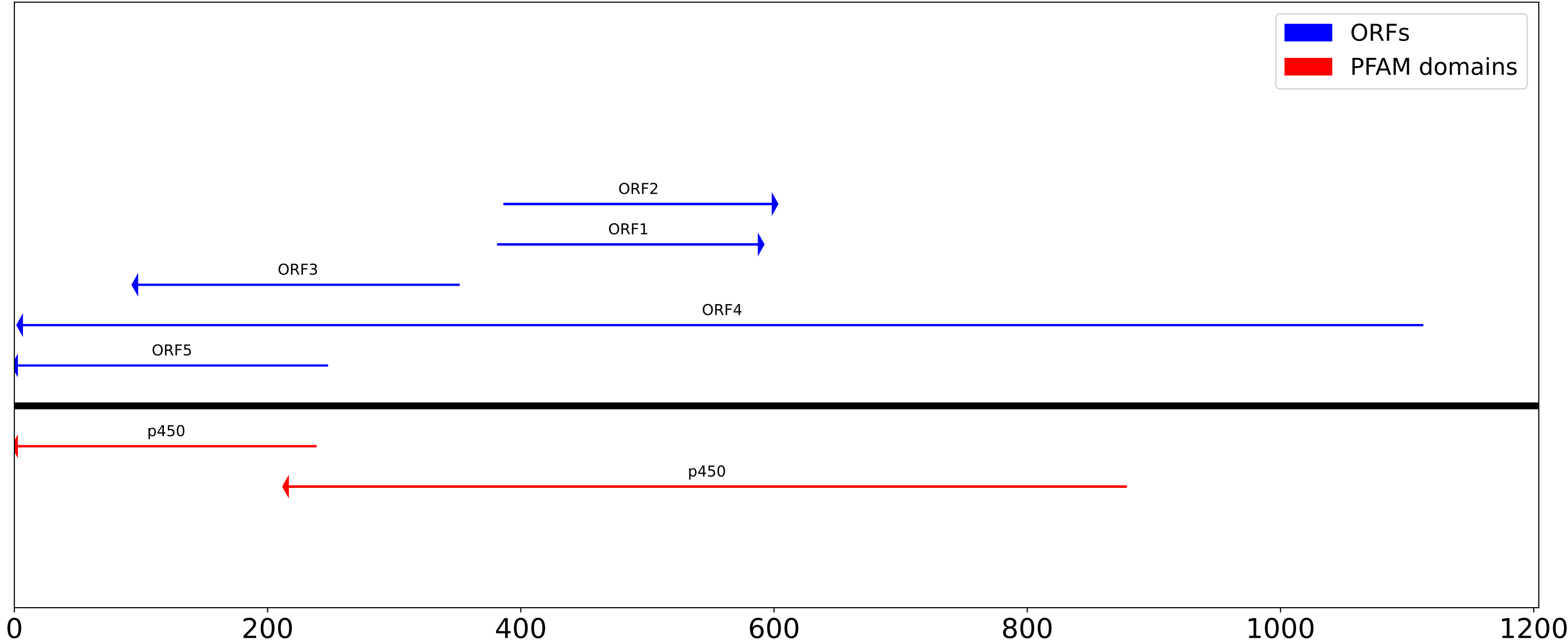


Before TEtrimmer 1204 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

