

Start Crop Point



End Crop Point

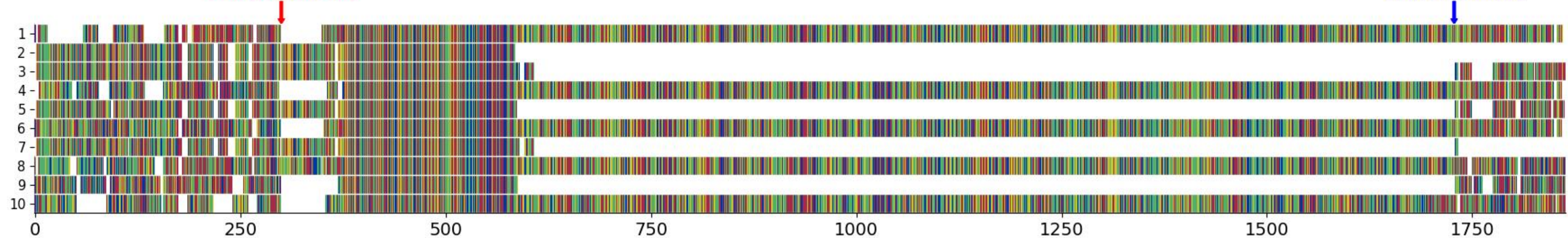


MSA length = 1429



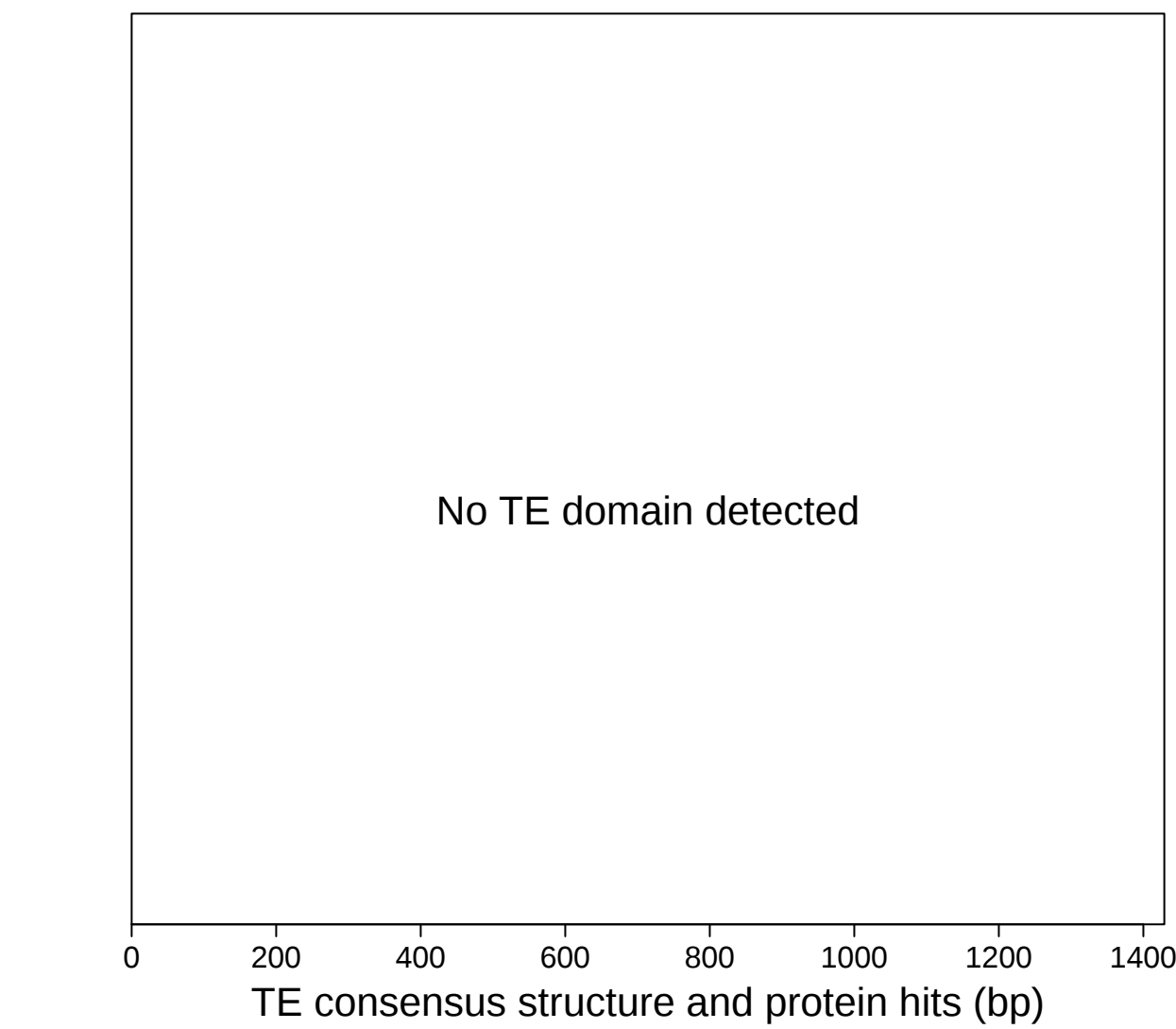
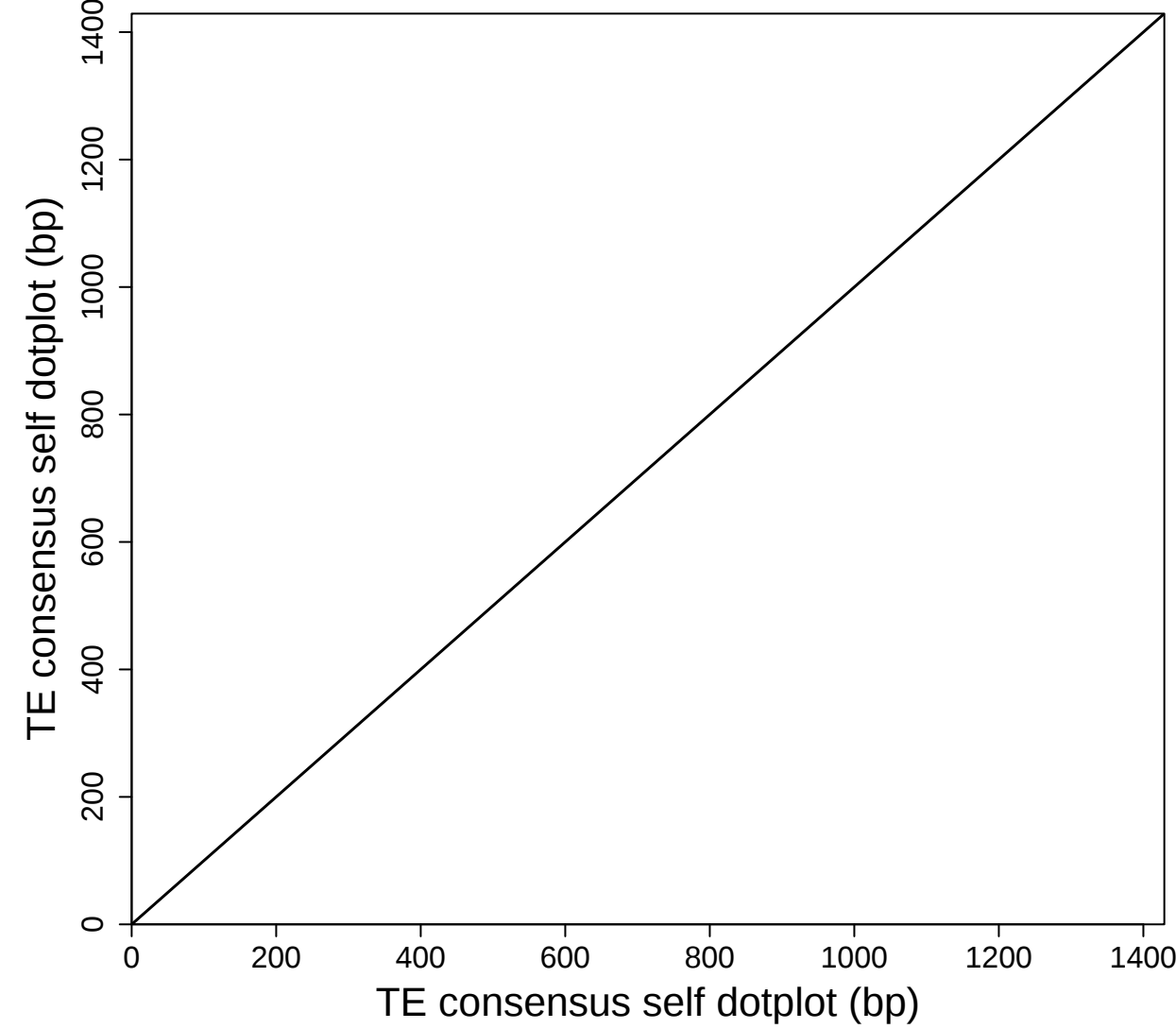
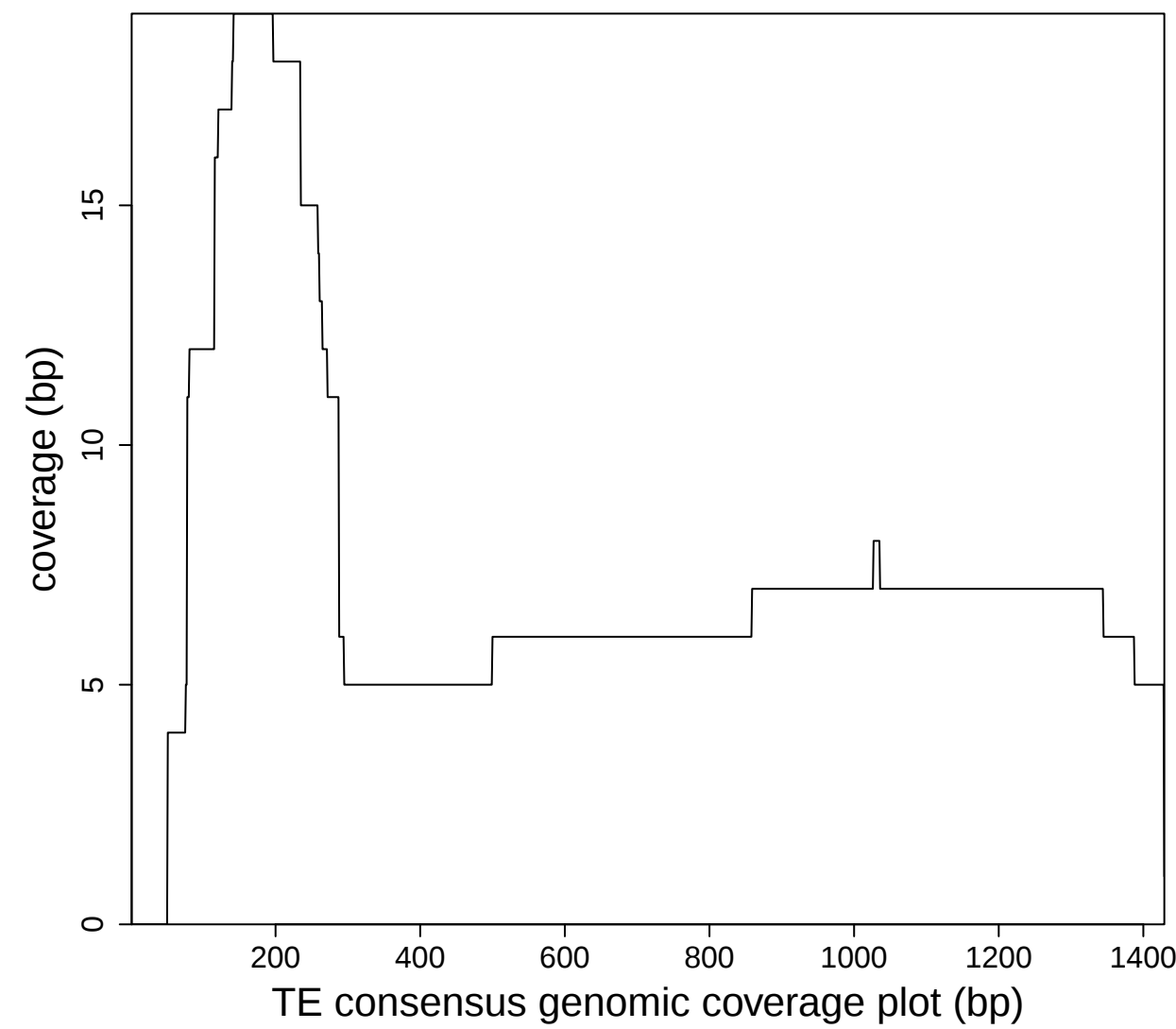
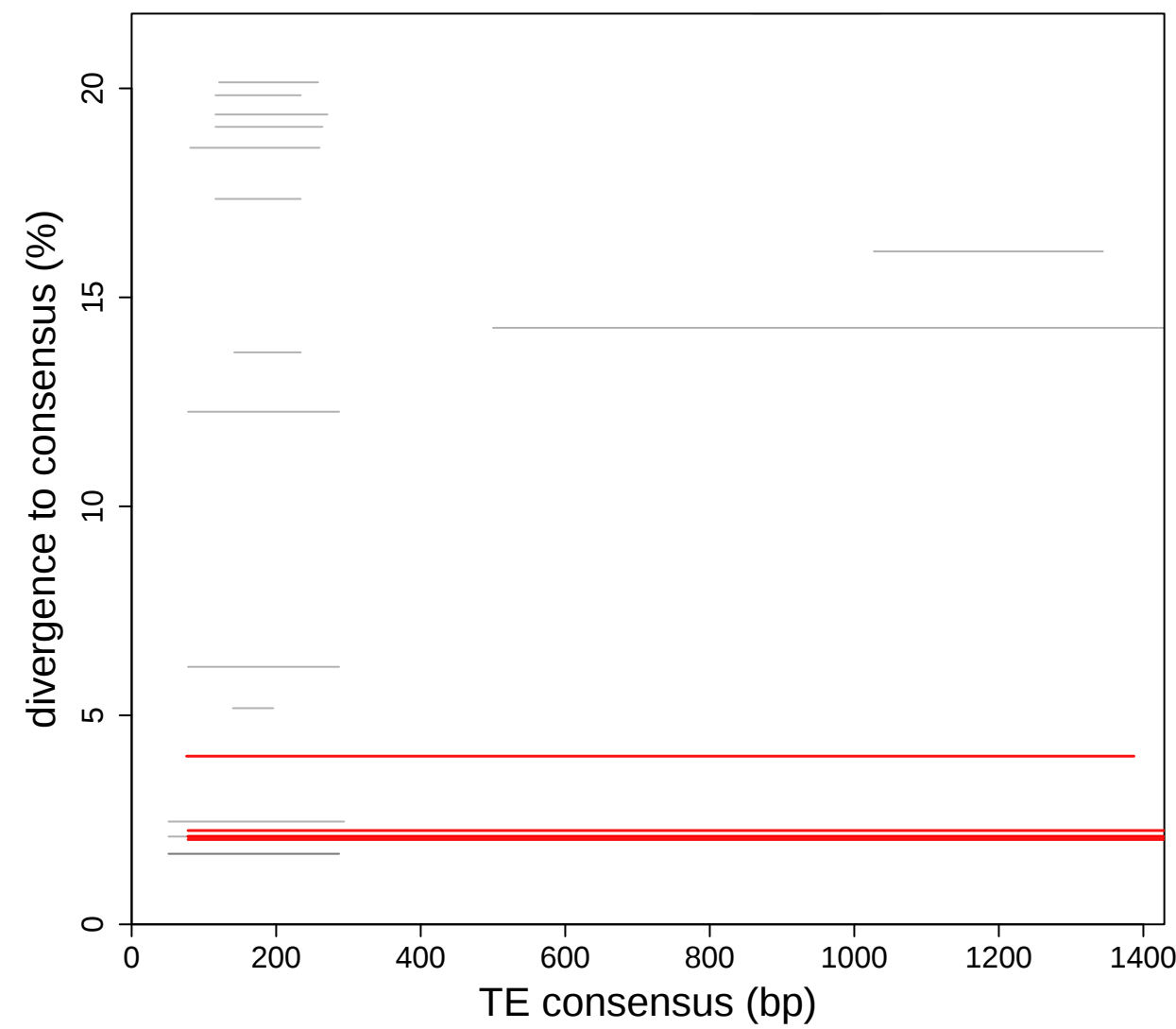
Start crop Point

End crop Point



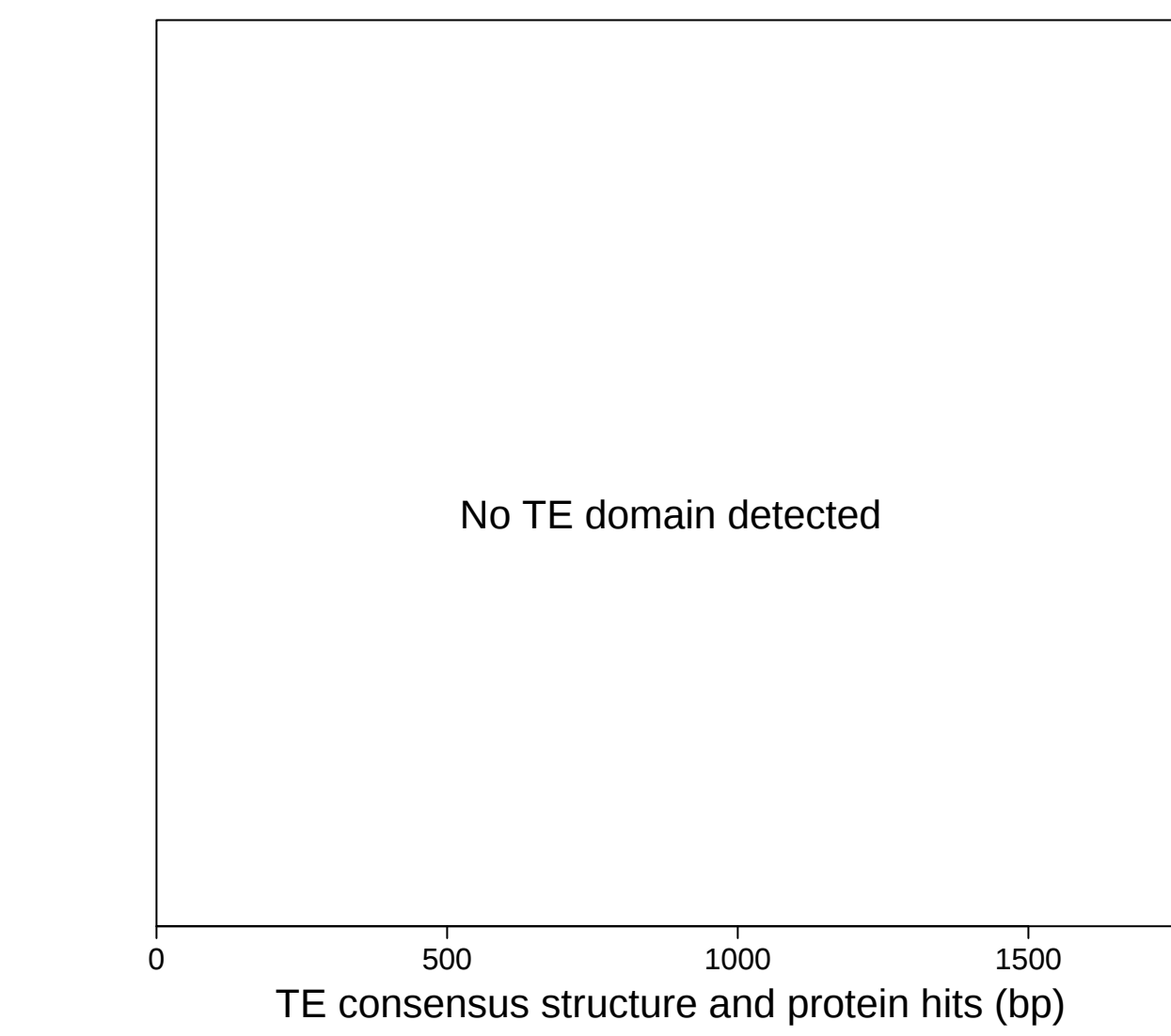
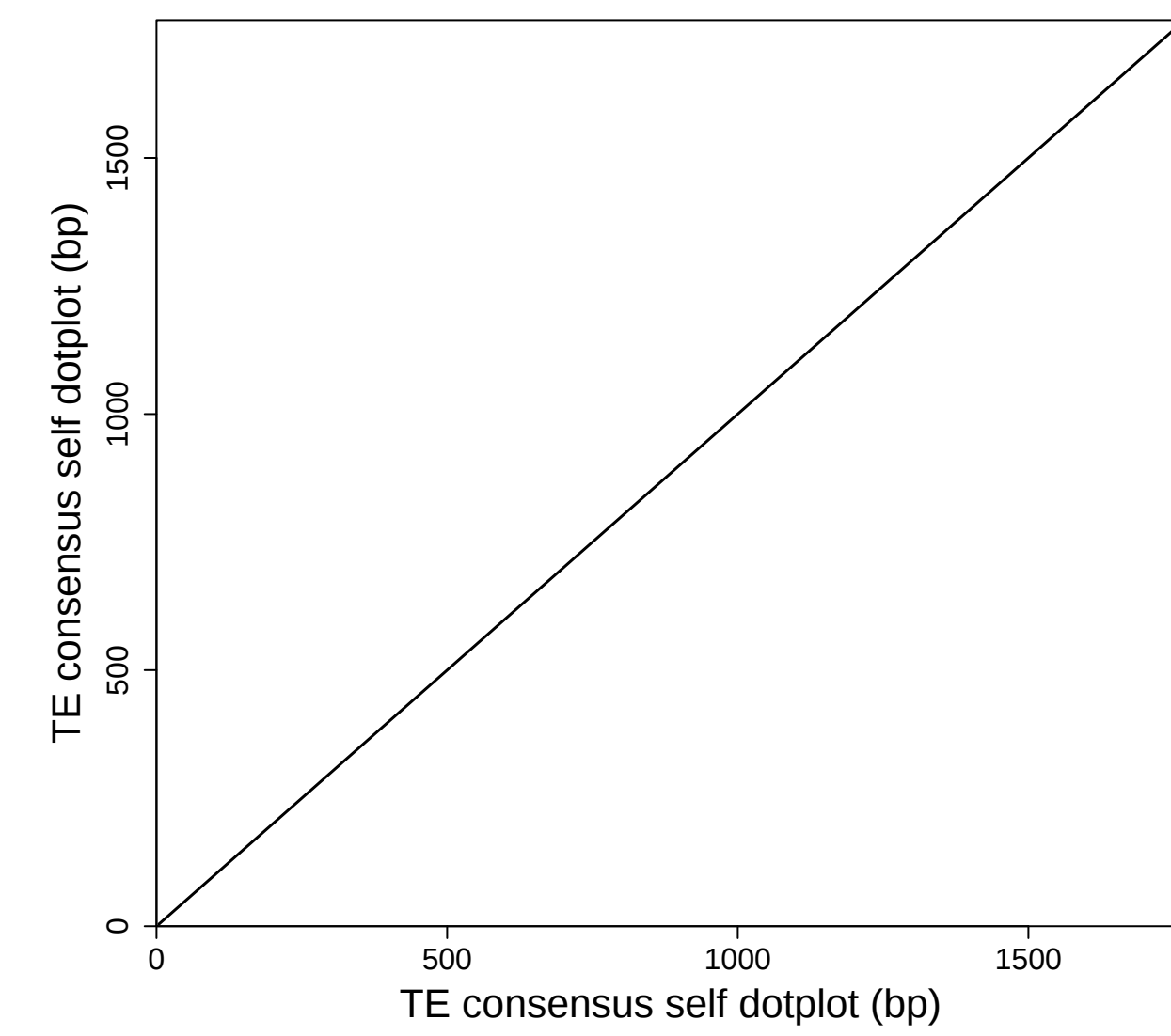
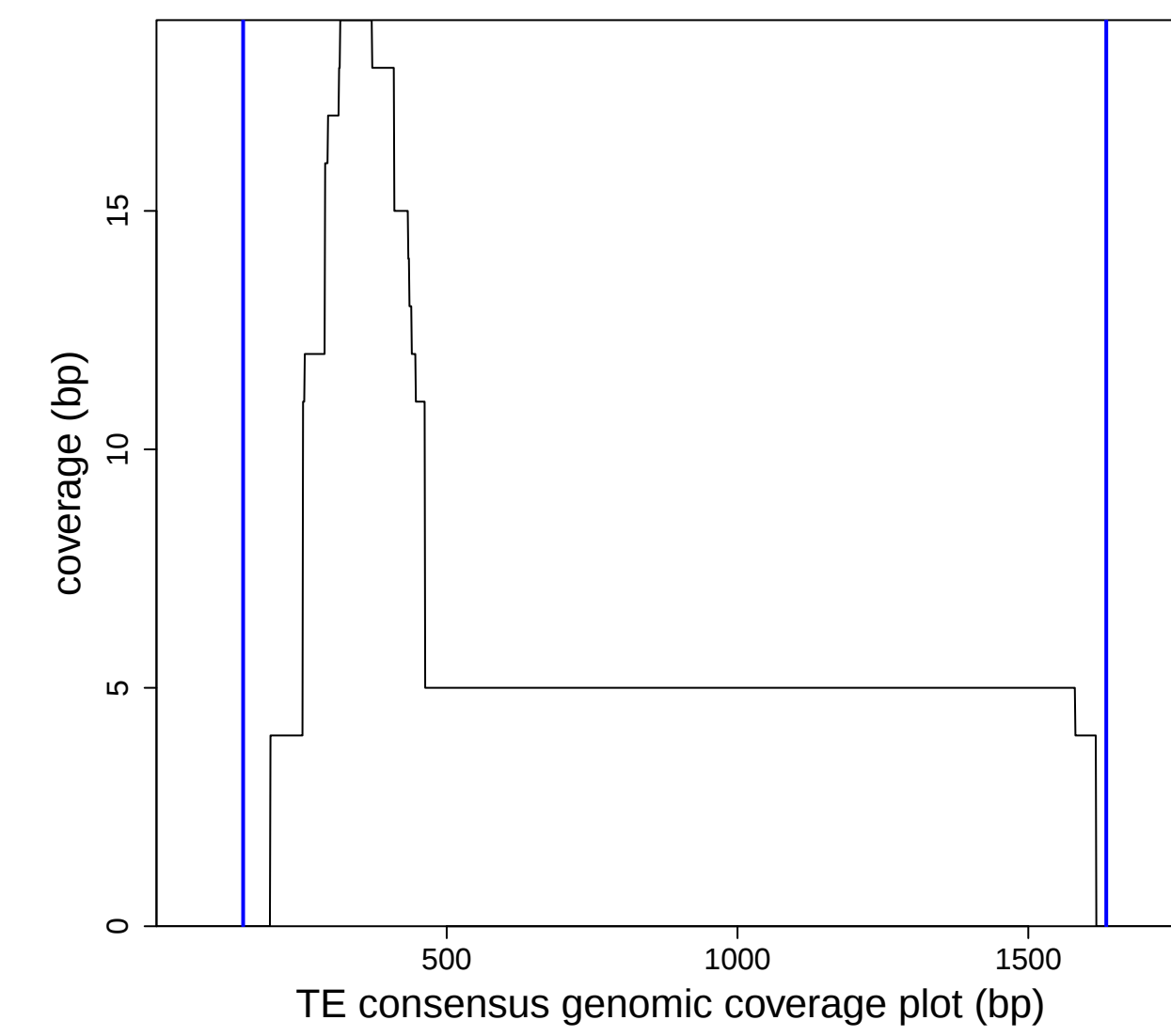
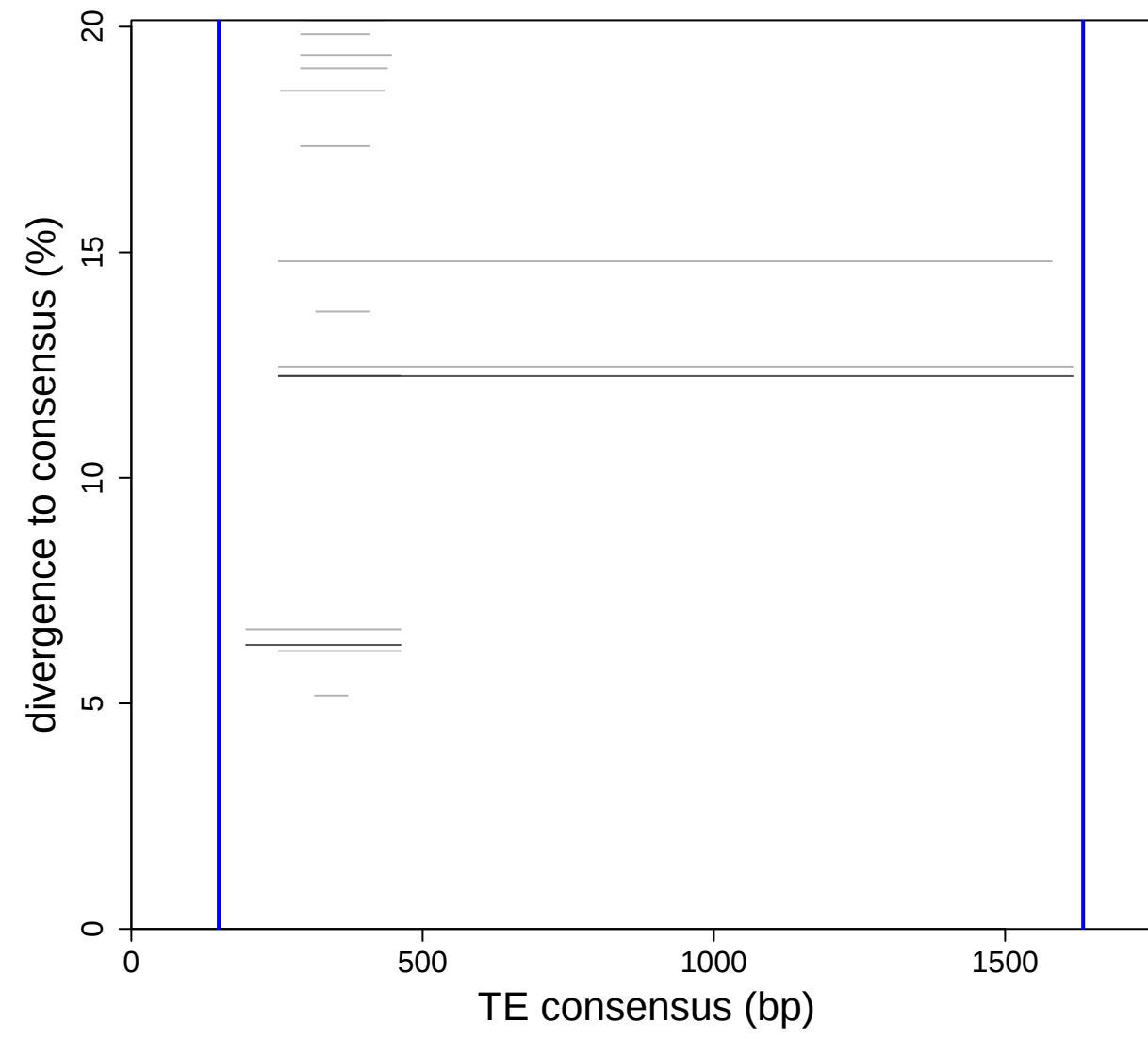
b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.fa_gs.fa
size: 1429bp; fragments: 22; full length: 5 (≥ 1286.1 bp)

After TEtrimmer 1429 bp



3.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1769bp; fragments: 19; full length: 0 (>=1592.1bp)

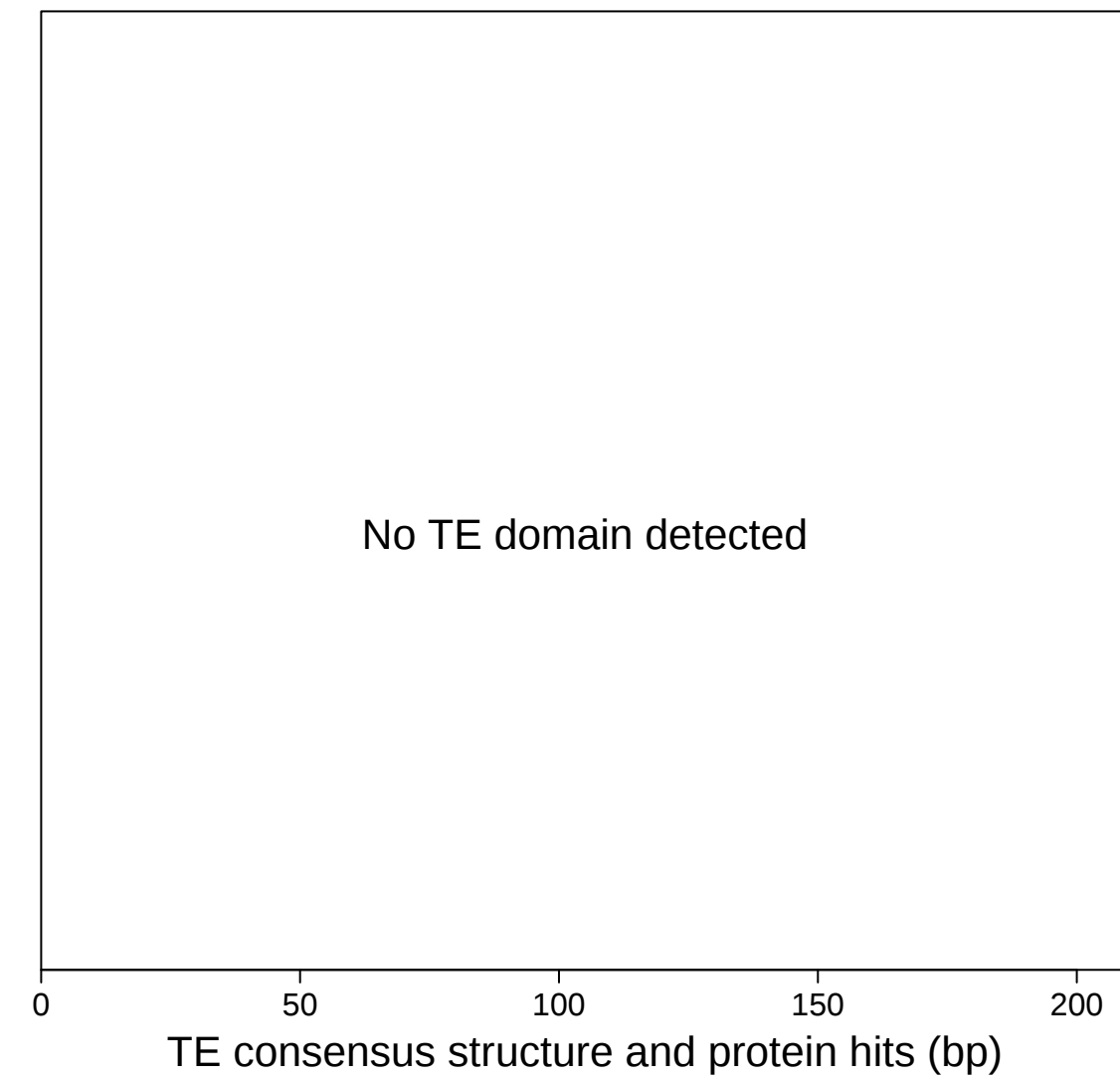
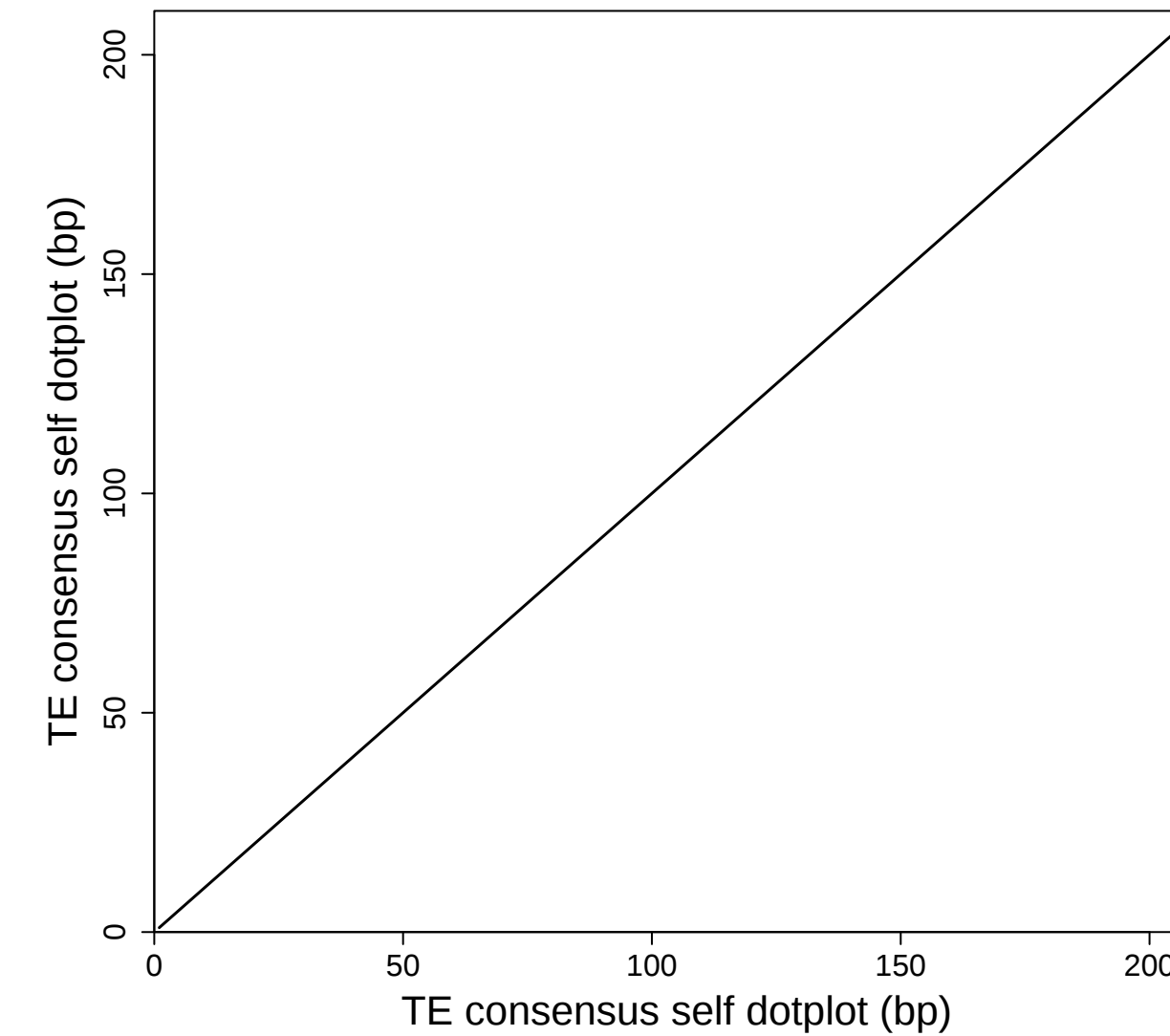
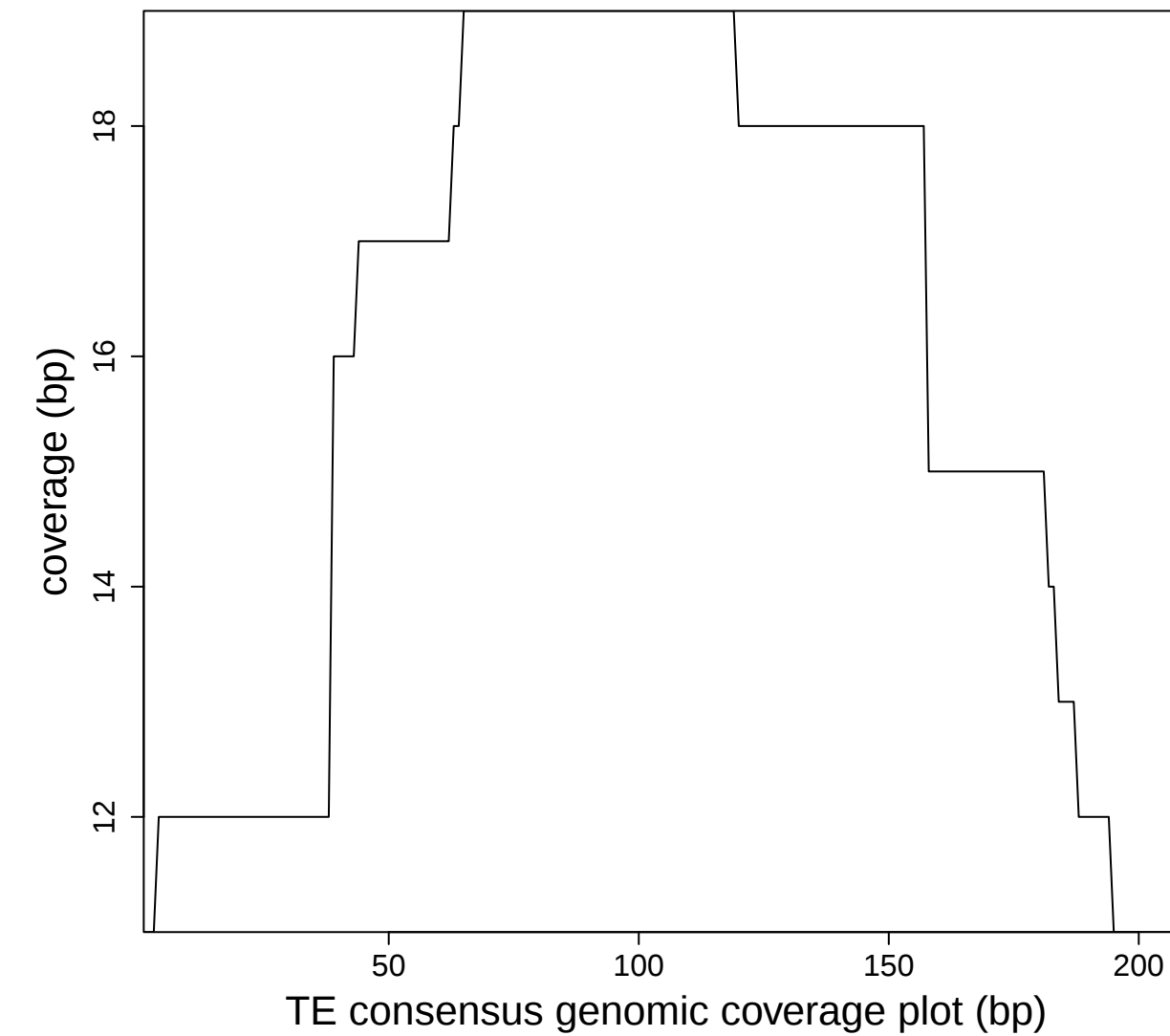
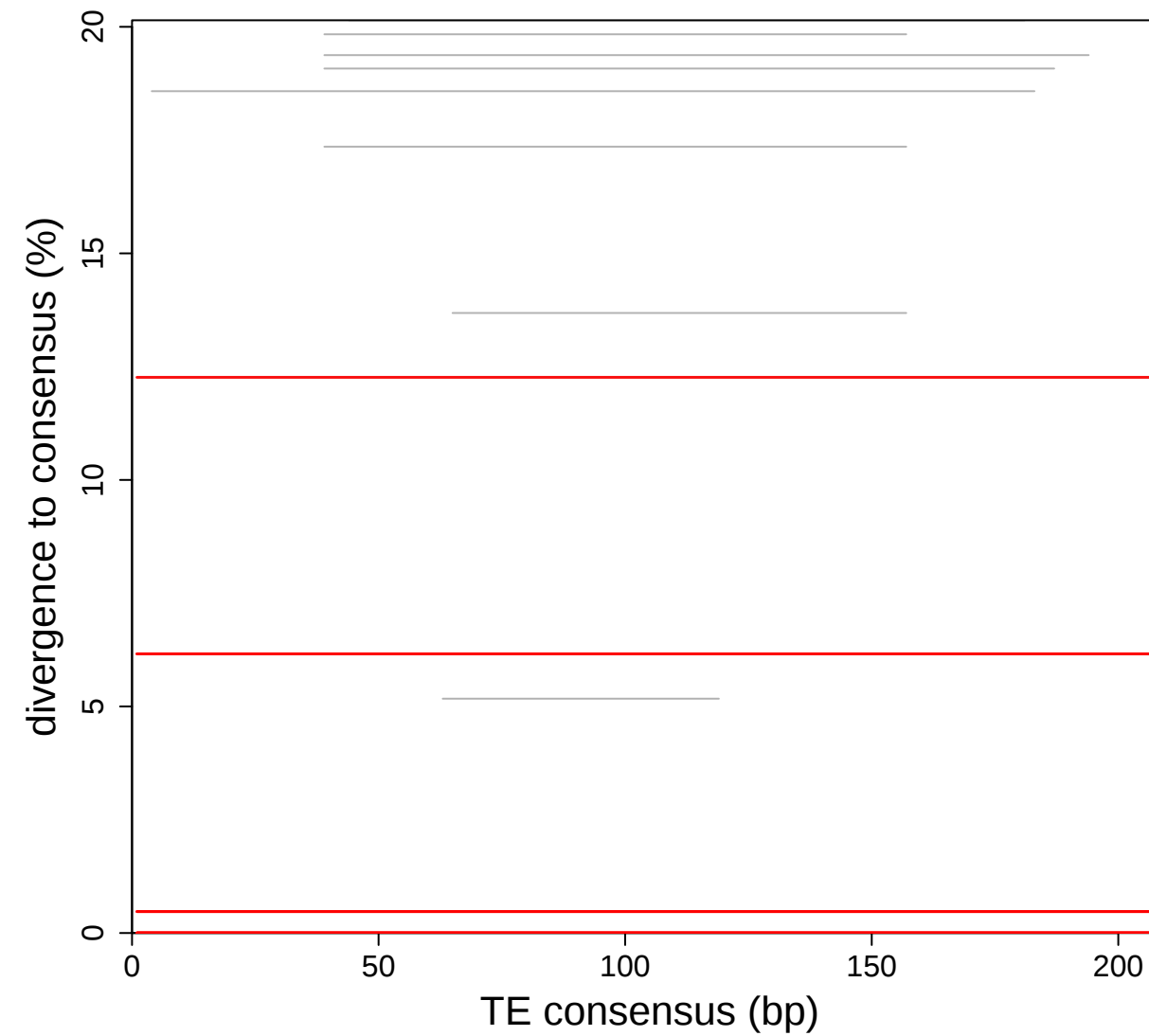
After TEtrimmer Extended plot Blue lines are boundaries



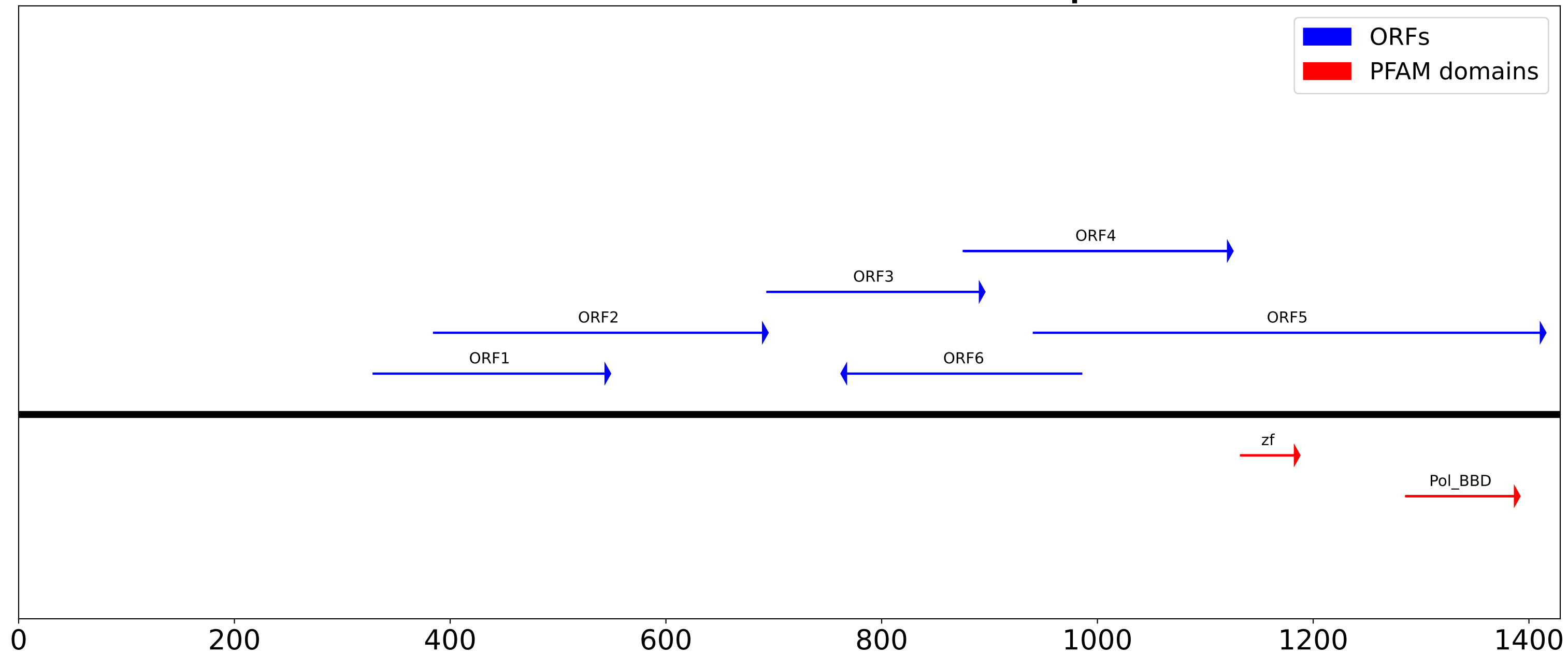
Before TEtrimmer 210 bp

TE: ltr_1_family_338

size: 210bp; fragments: 19; full length: 11 (≥ 189 bp)



After TEtrimmer ORF and PFAM domain plot



Dotplot

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

Isensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)