

Start crop Point



End crop Point

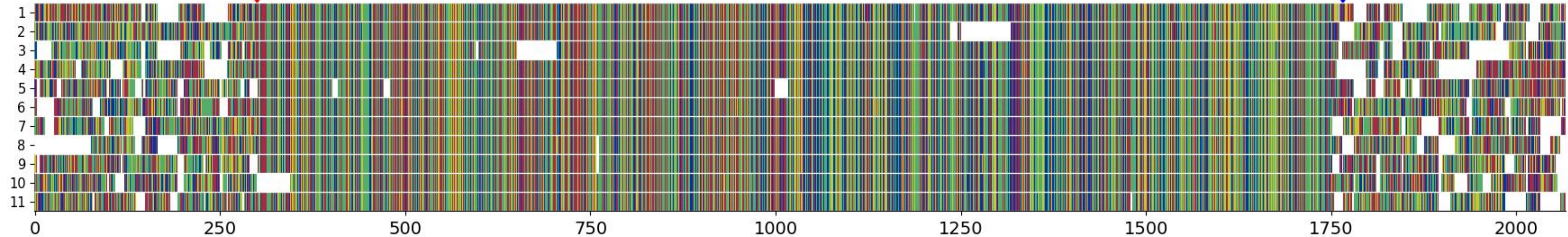


MSA length = 1467



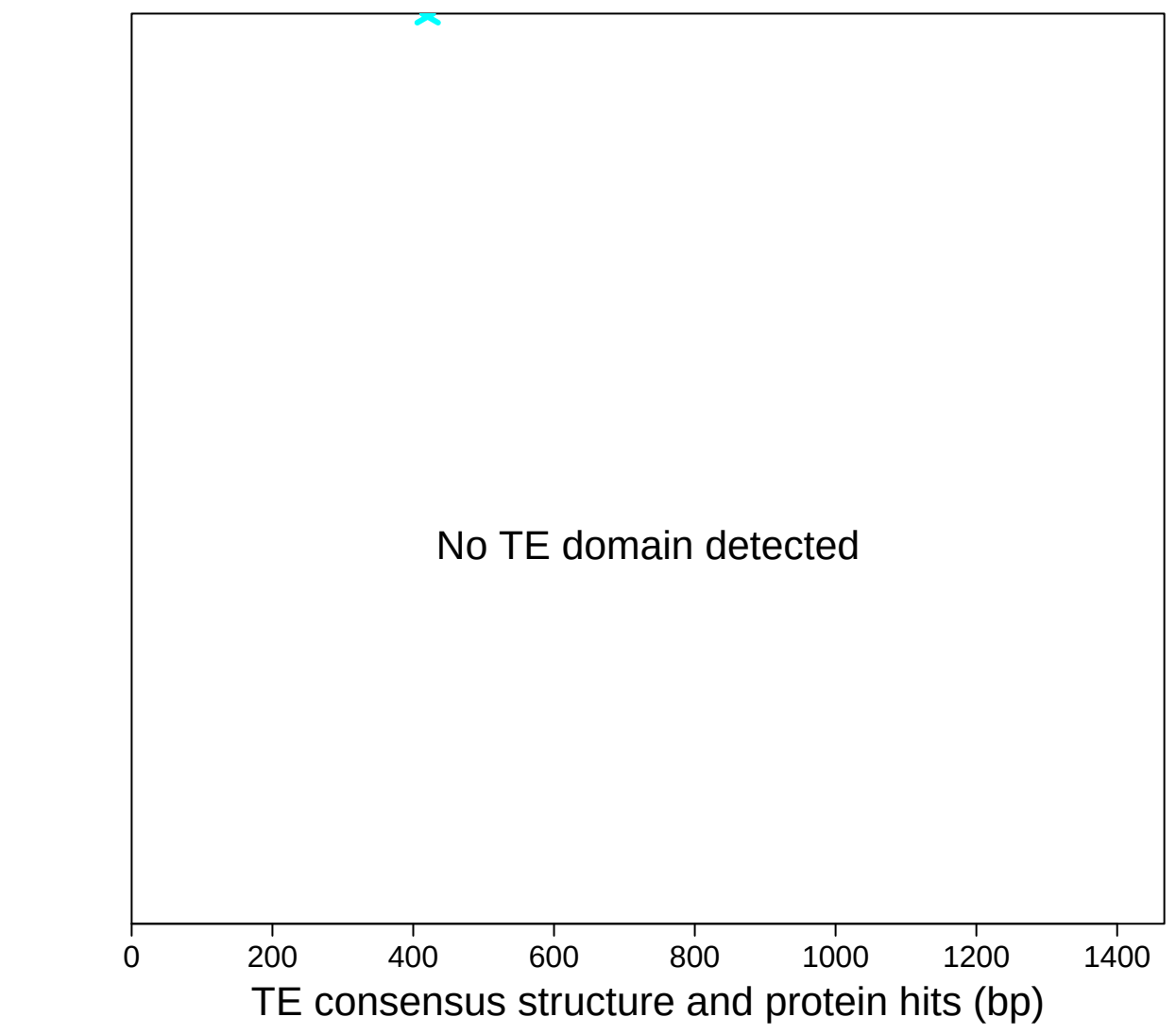
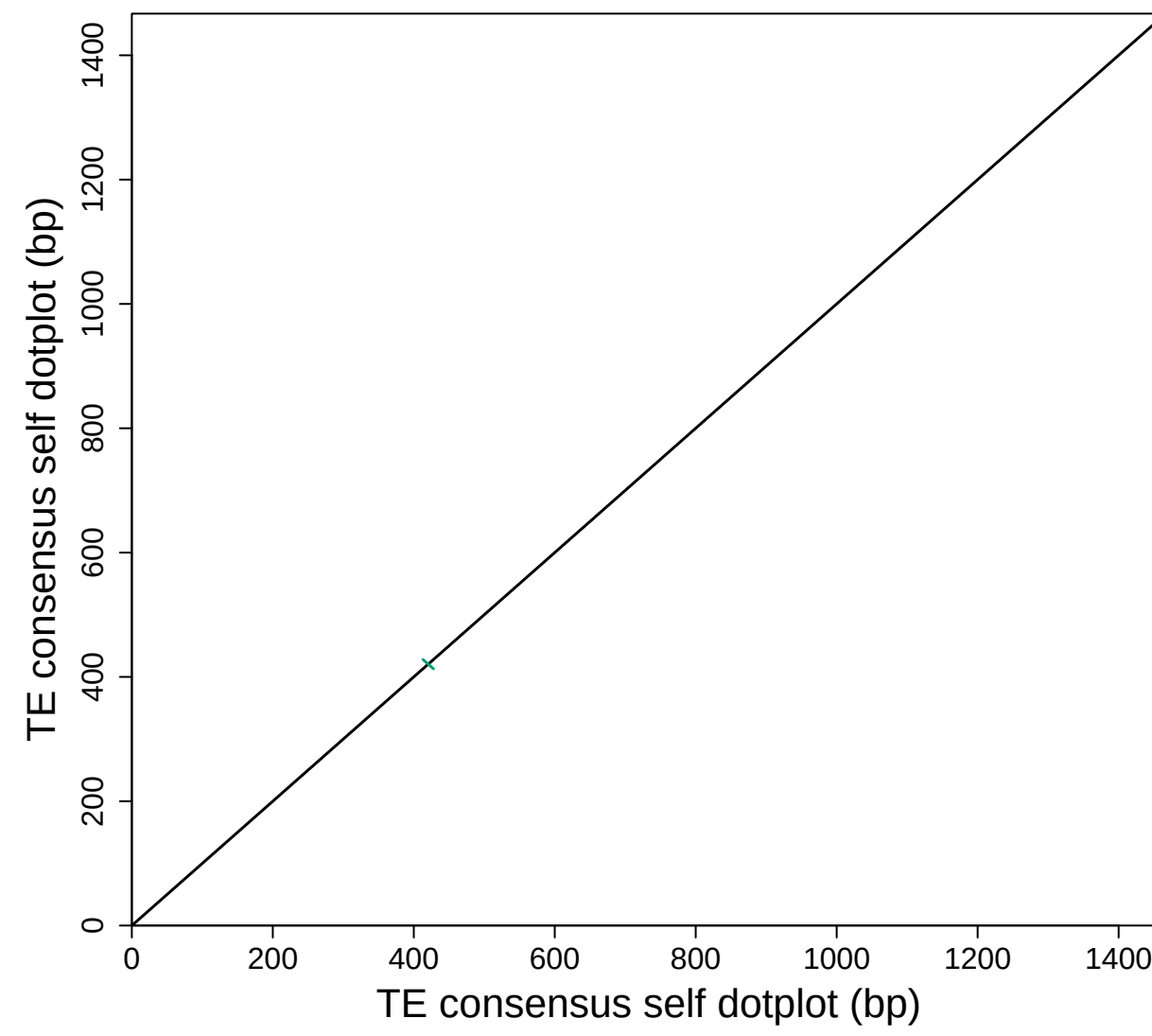
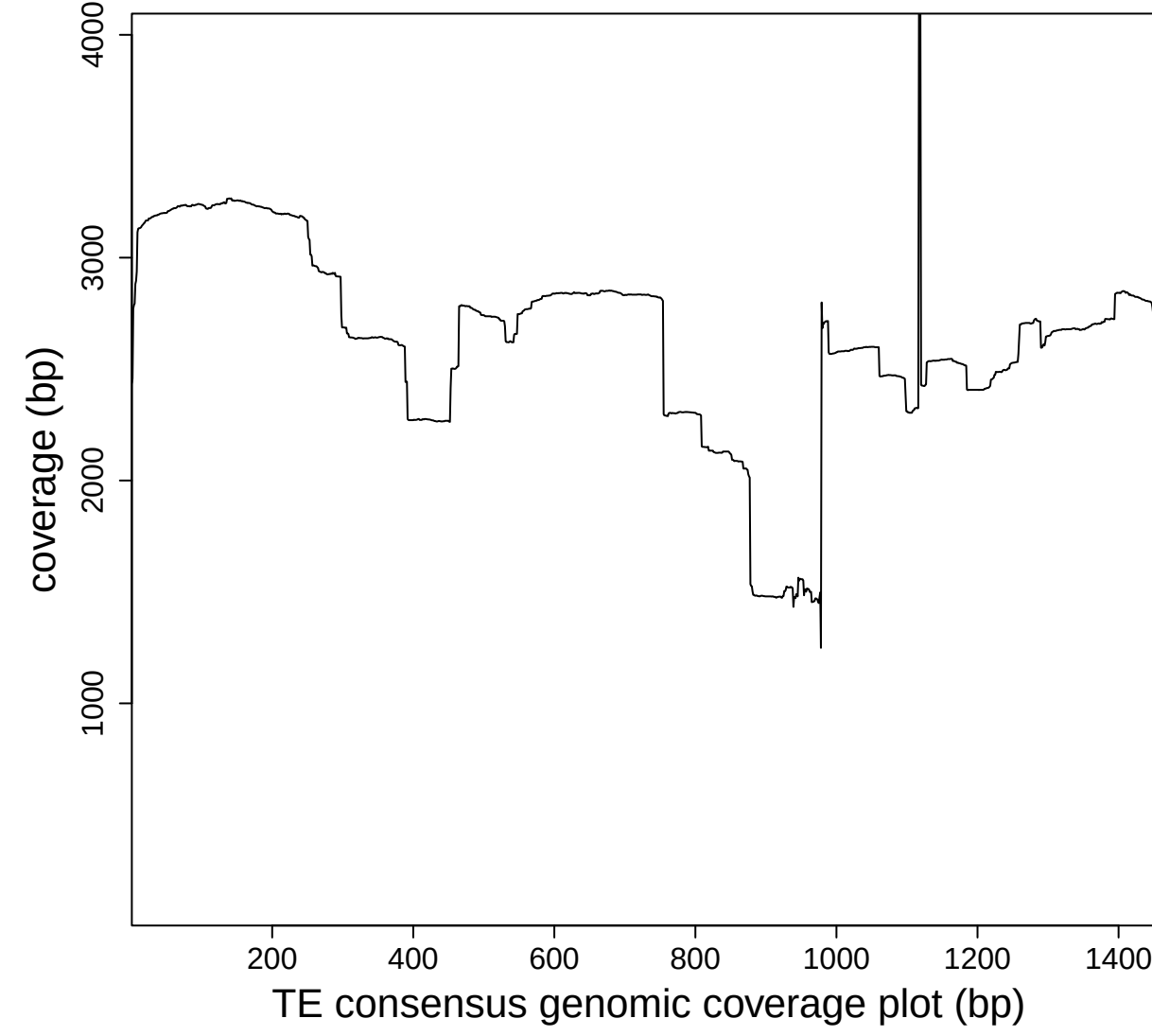
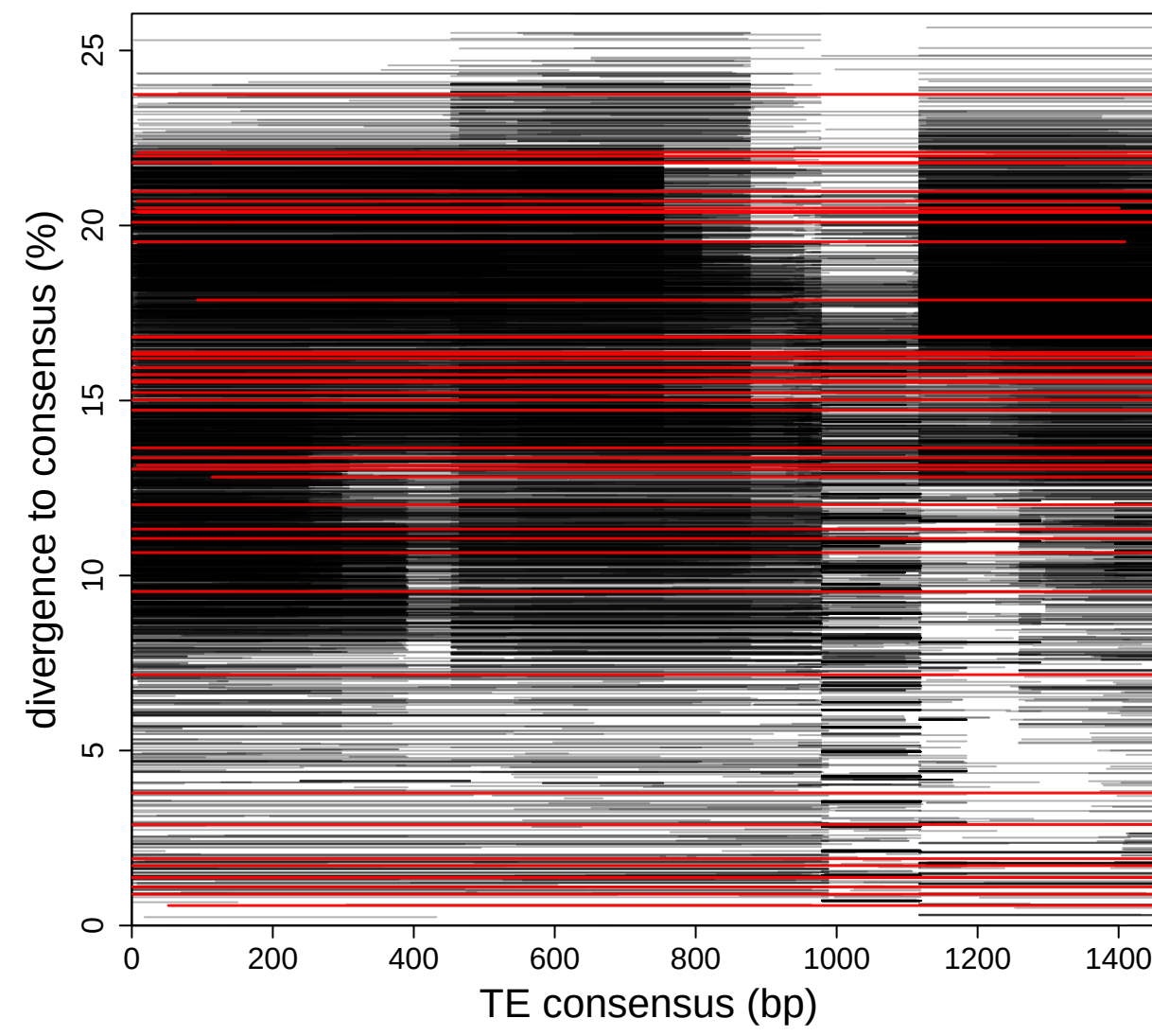
Start crop Point

End crop Point



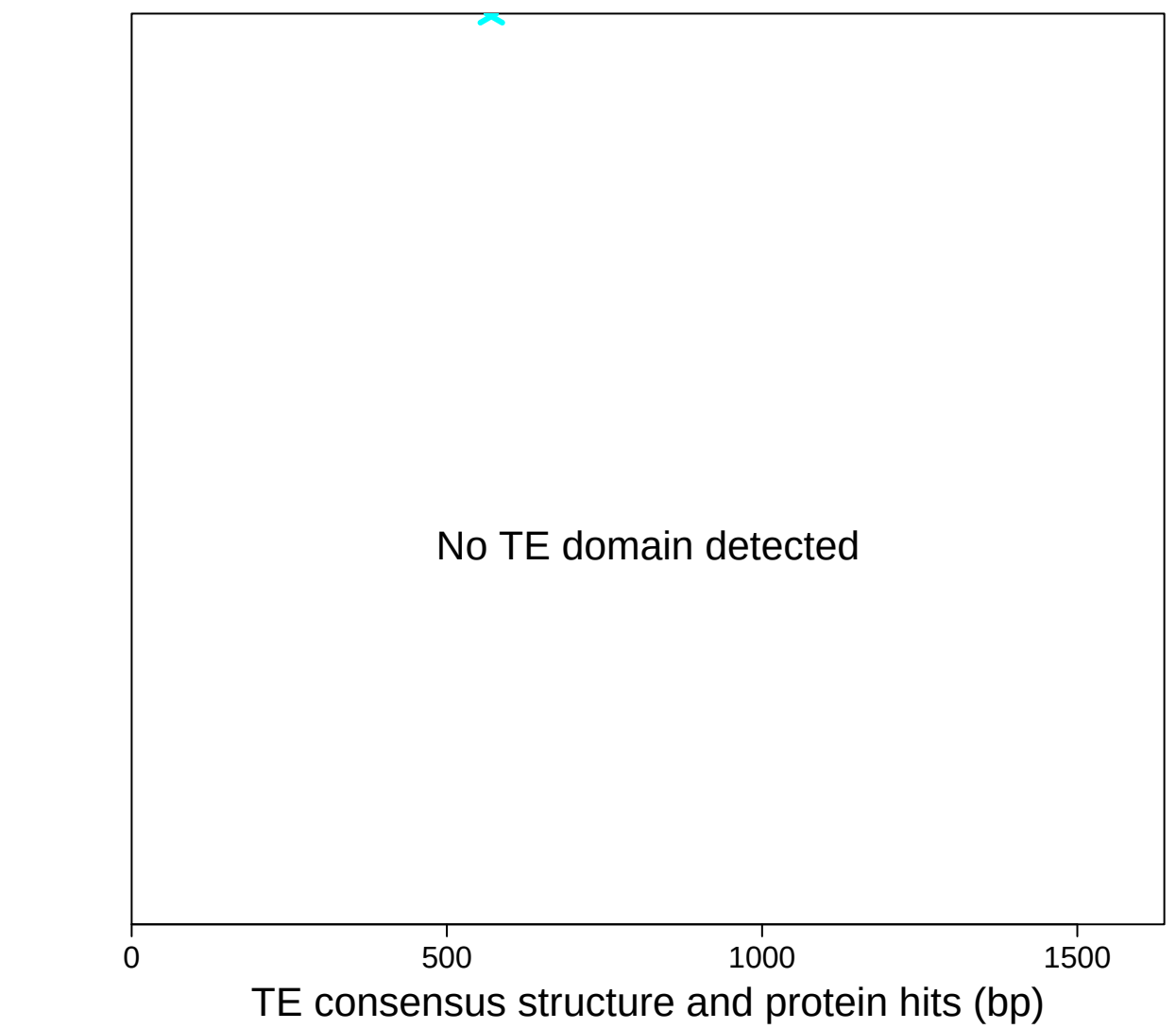
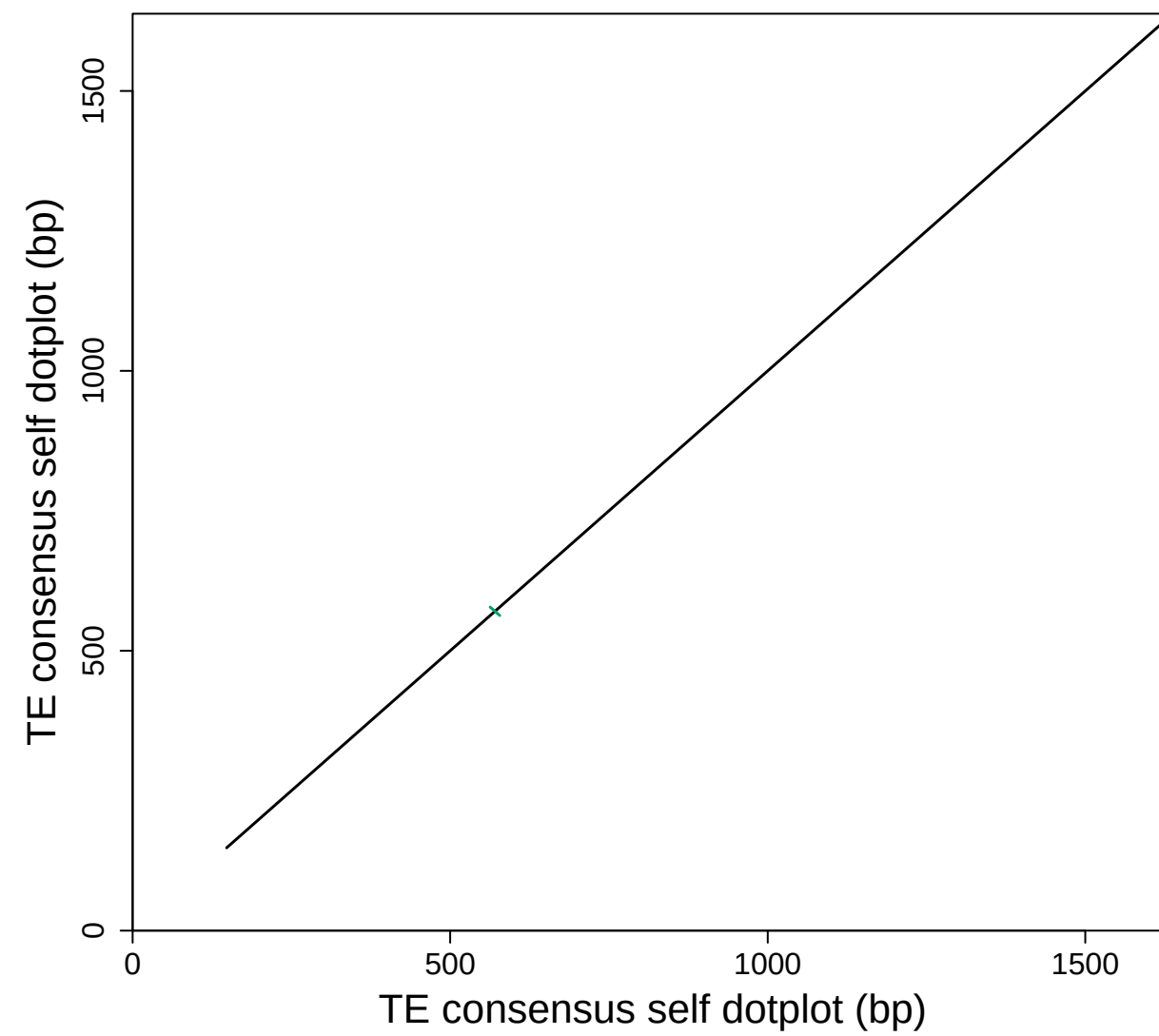
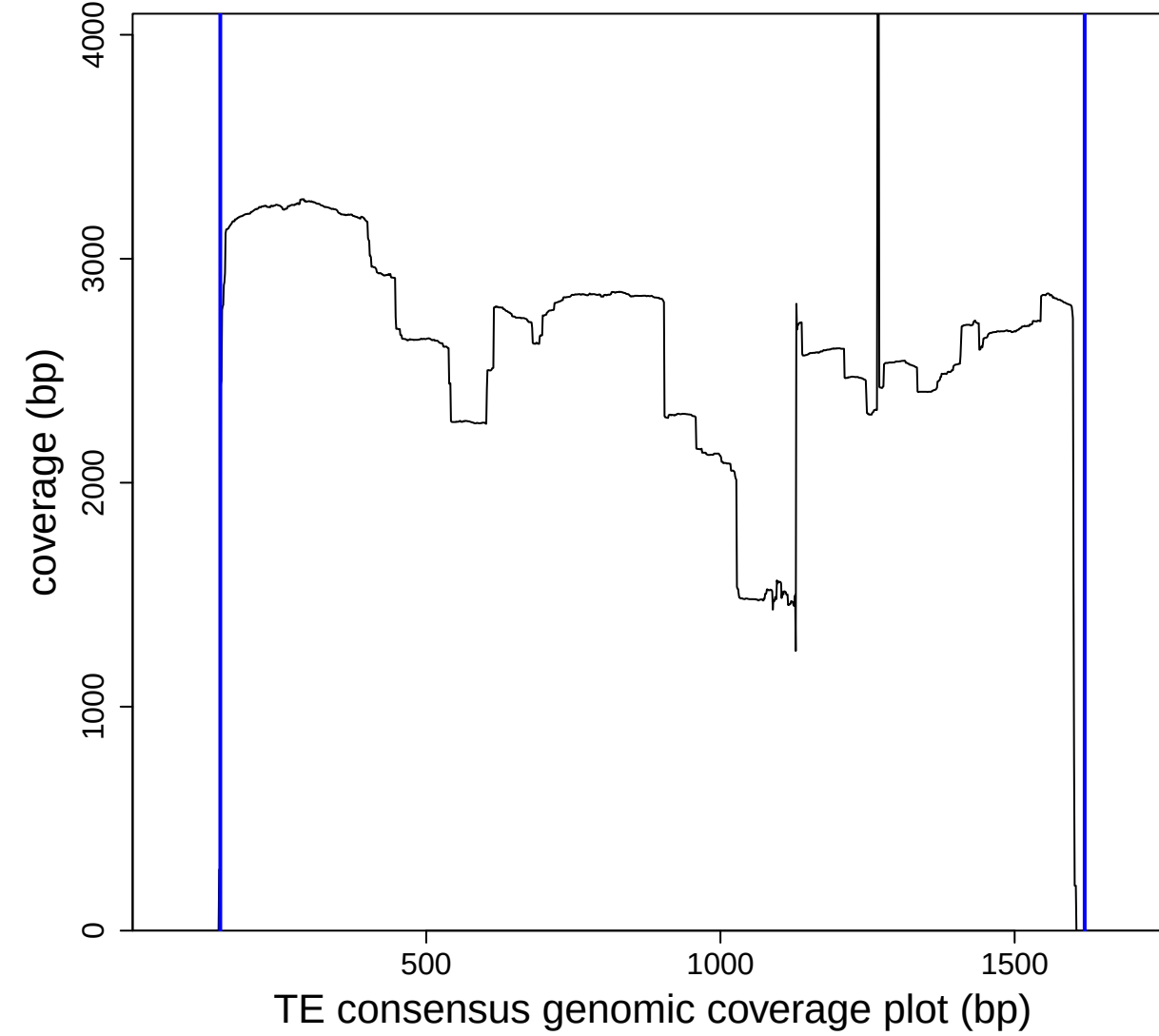
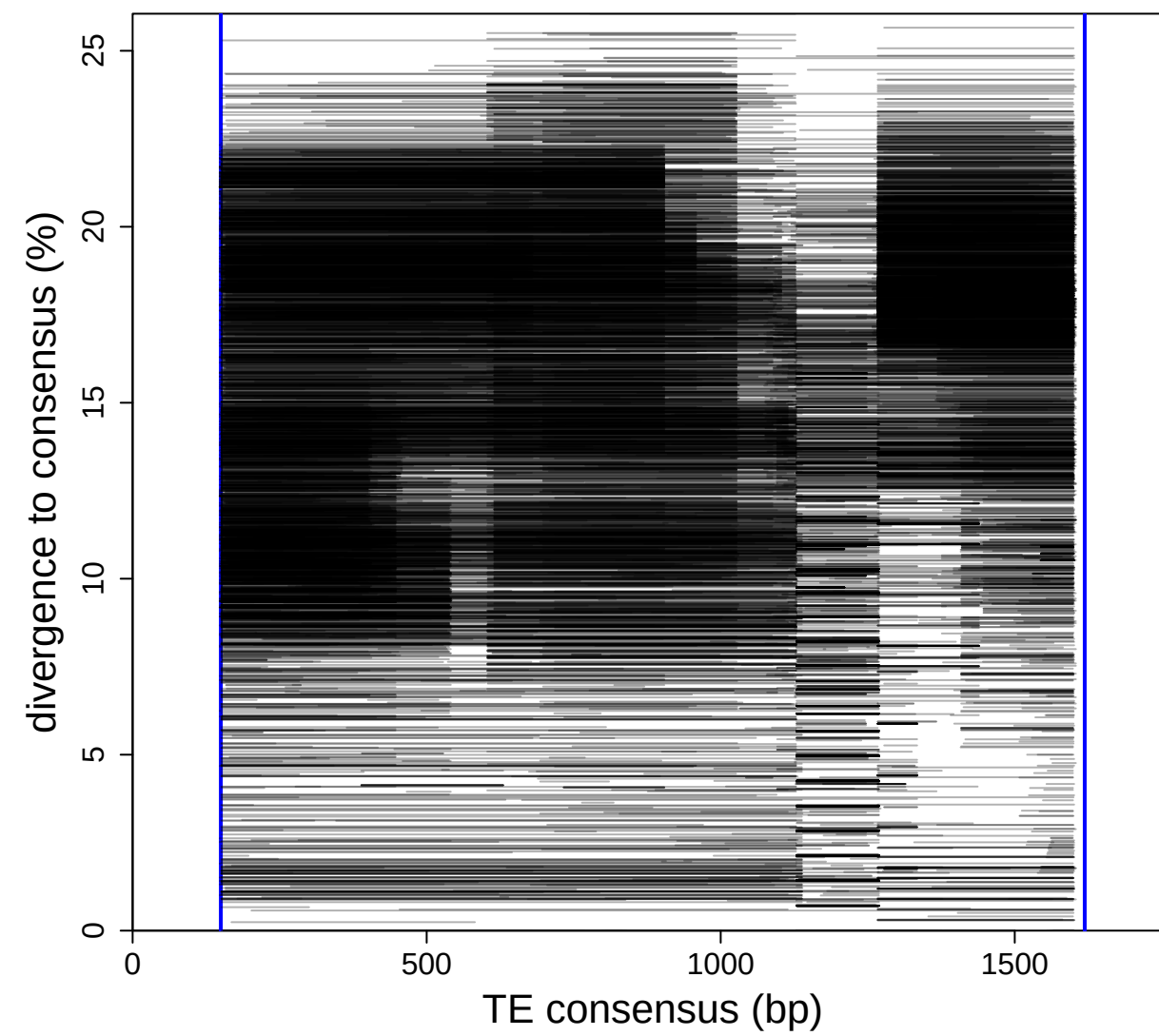
b.bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 1467bp; fragments: 9679; full length: 45 (≥ 1320.3 bp)

After TETrimmer 1467 bp

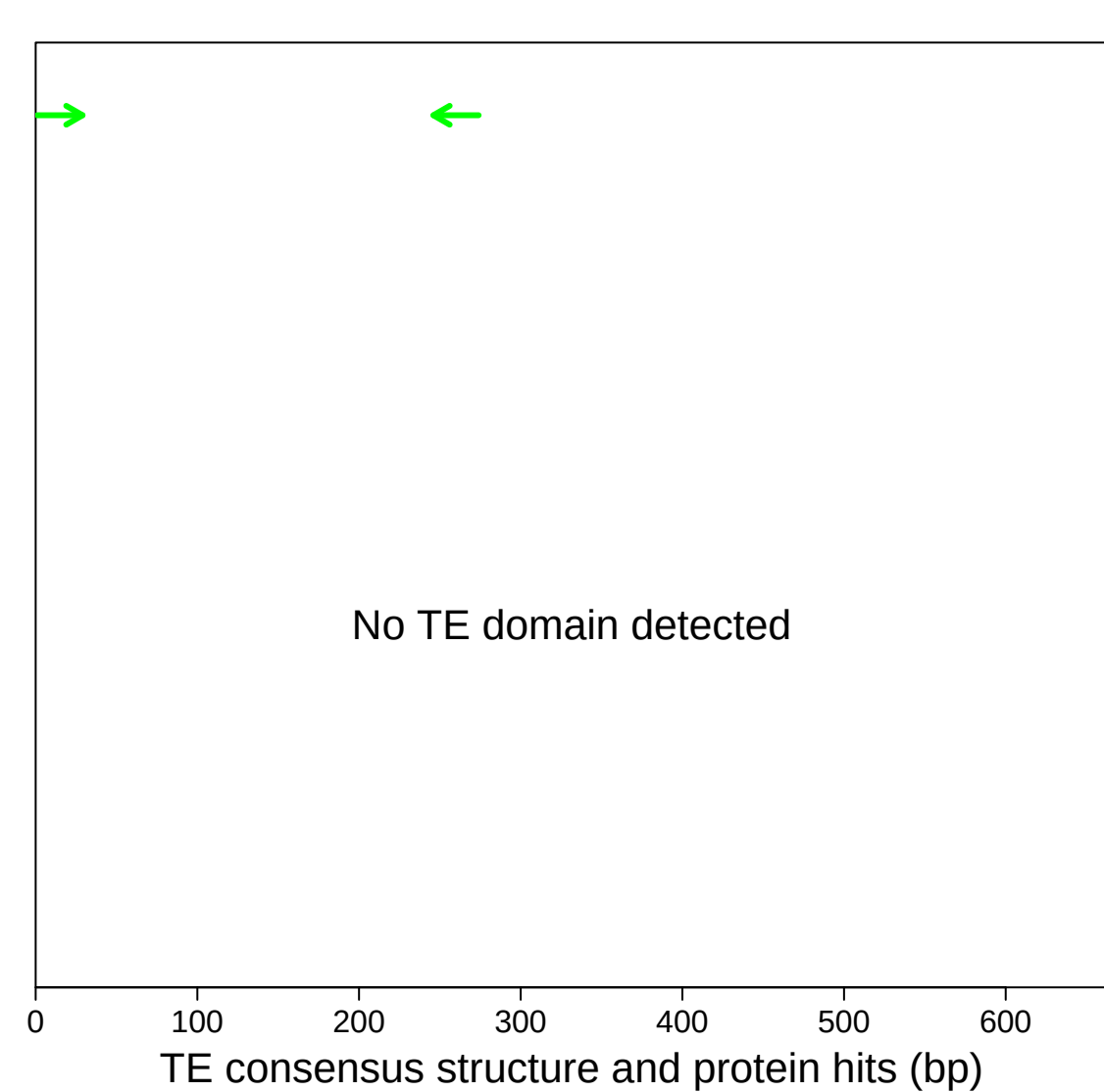
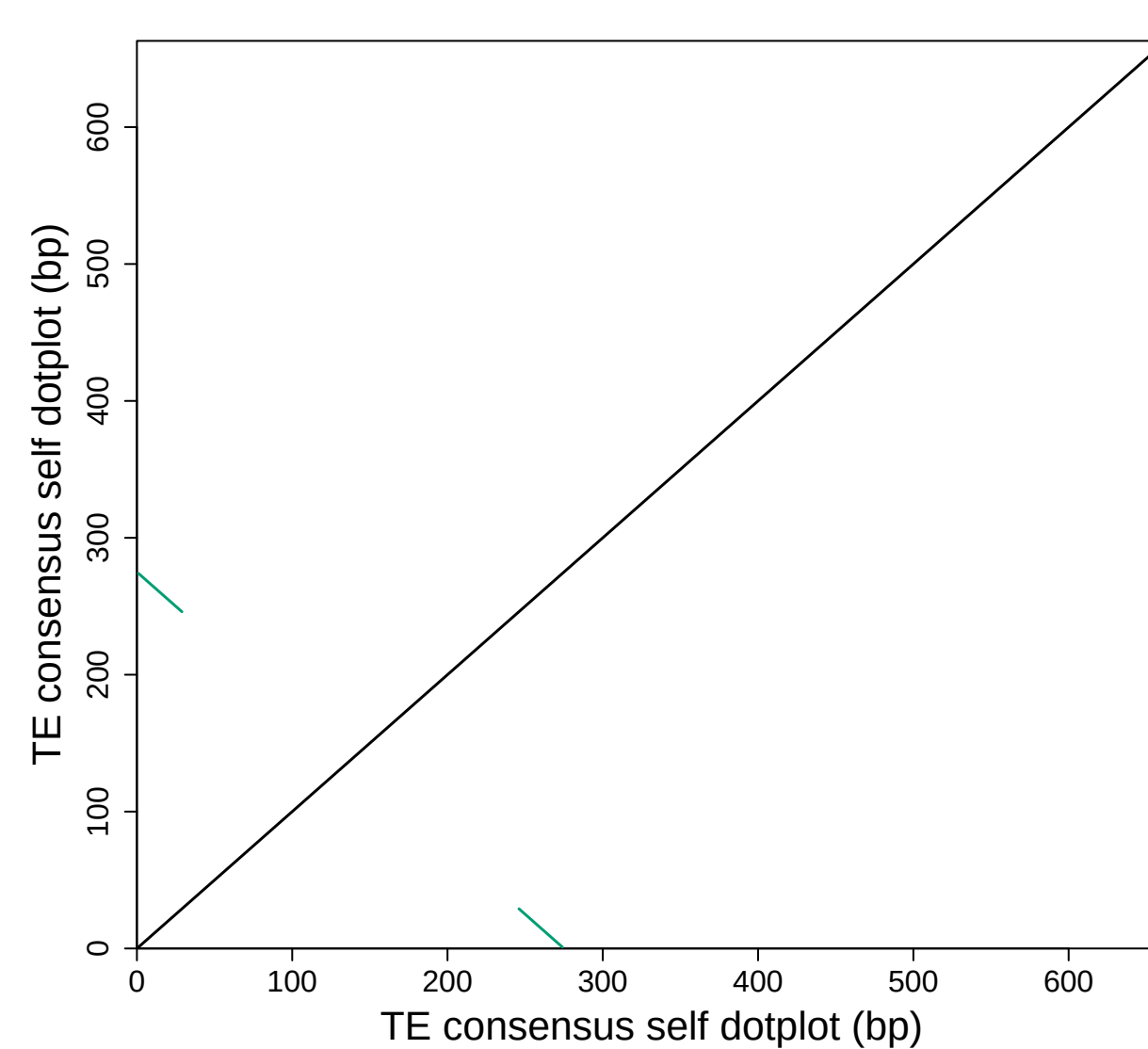
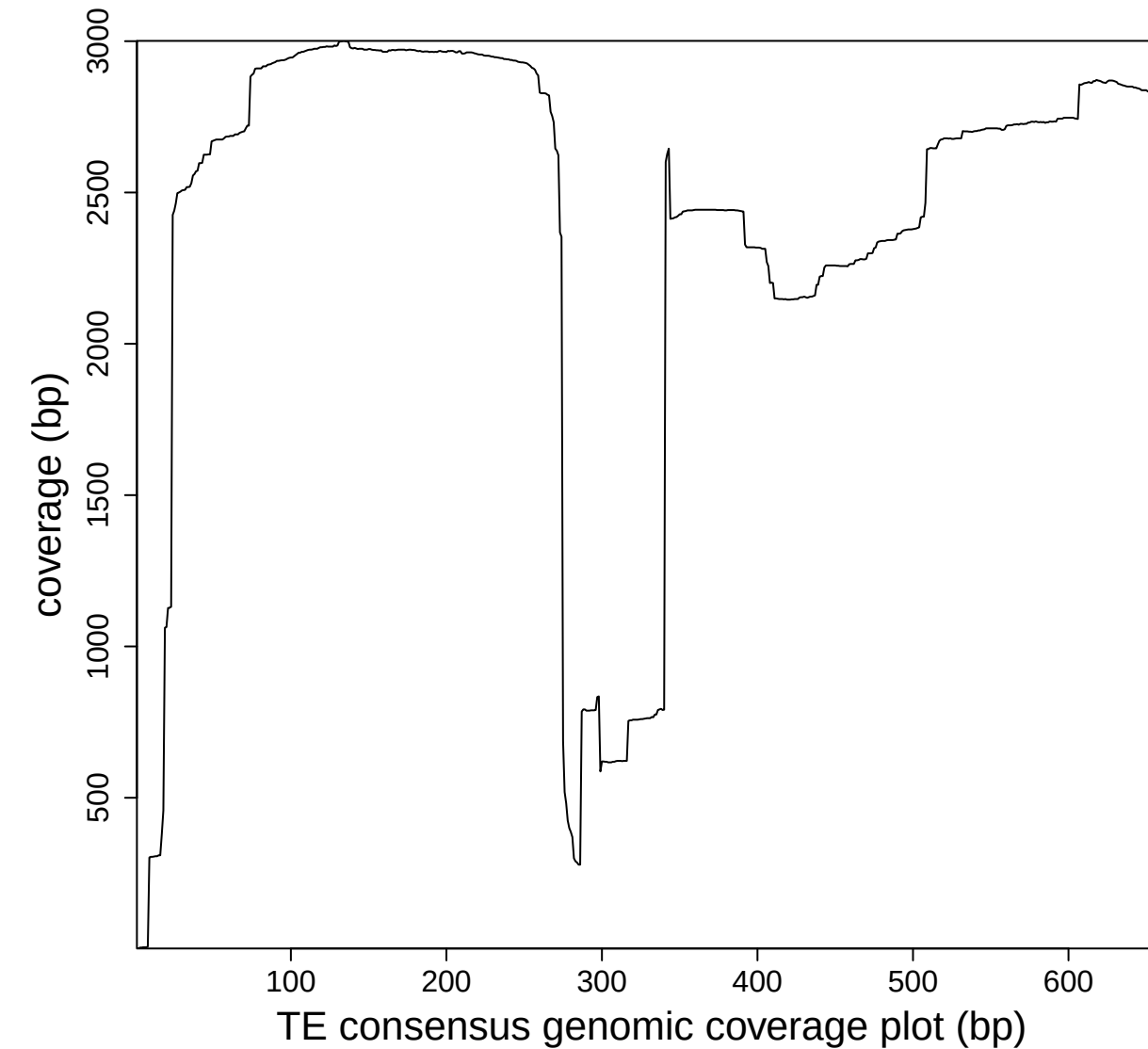
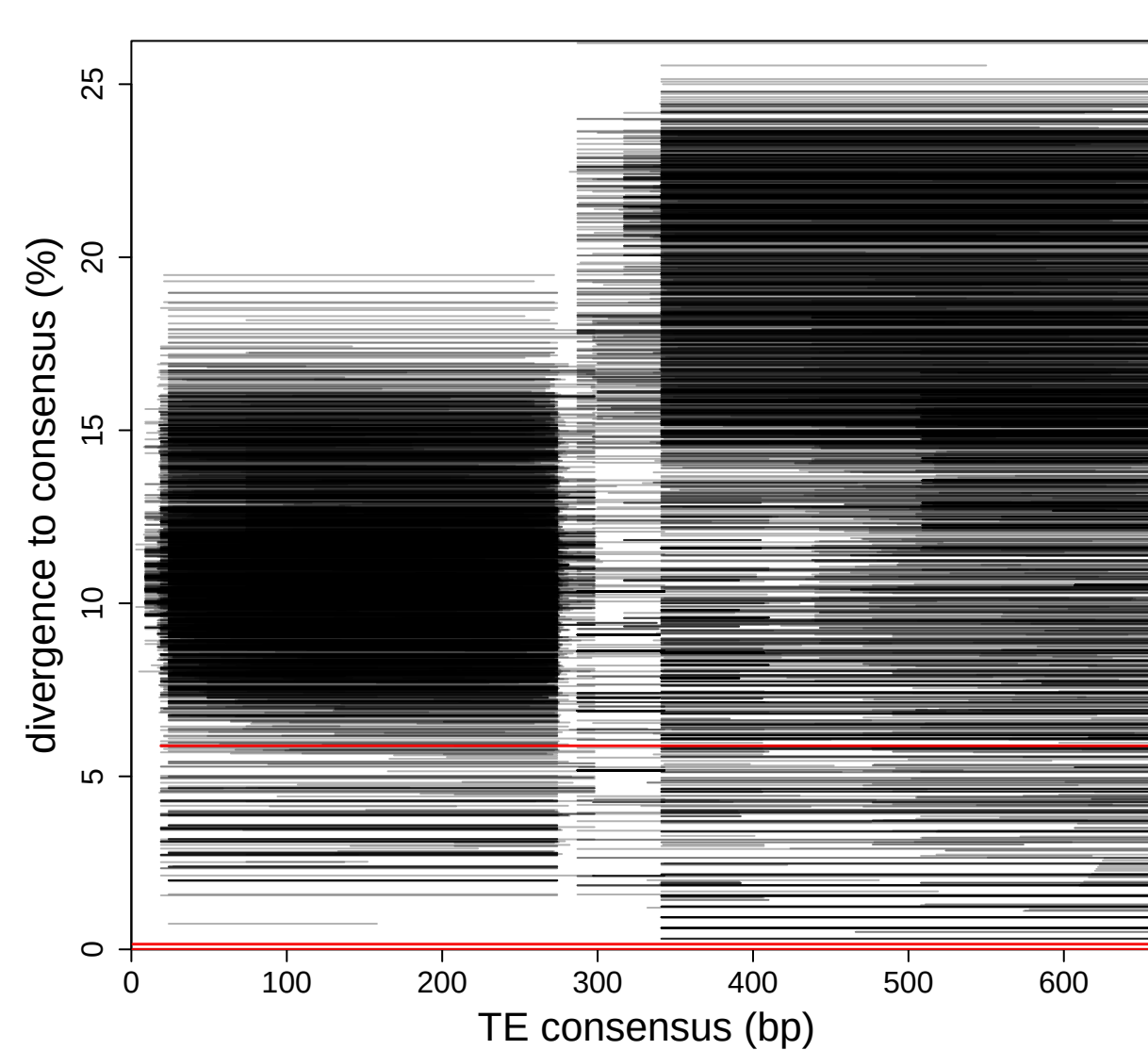


1.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1769bp; fragments: 9672; full length: 0 (>=1592.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

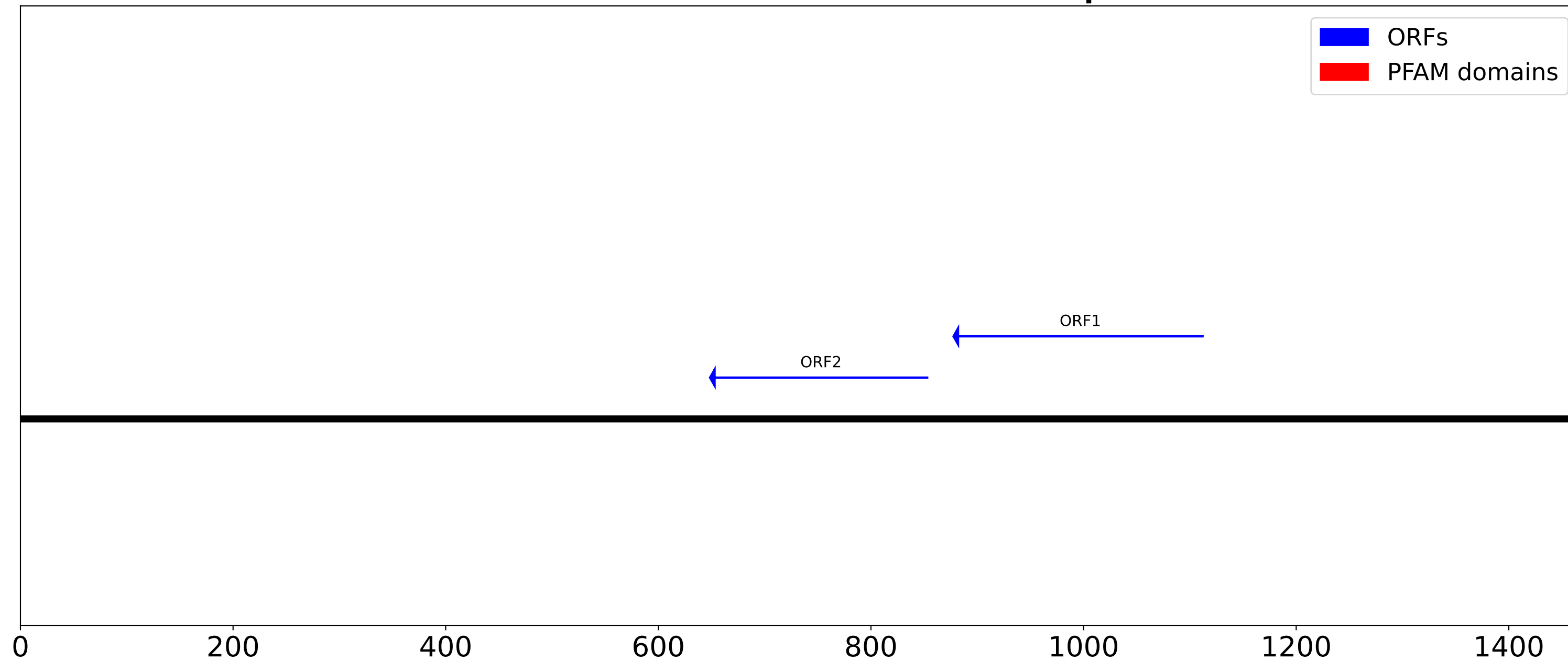


TE: ltr_1_family_257
size: 663bp; fragments: 6624; full length: 3 (≥ 596.7 bp)

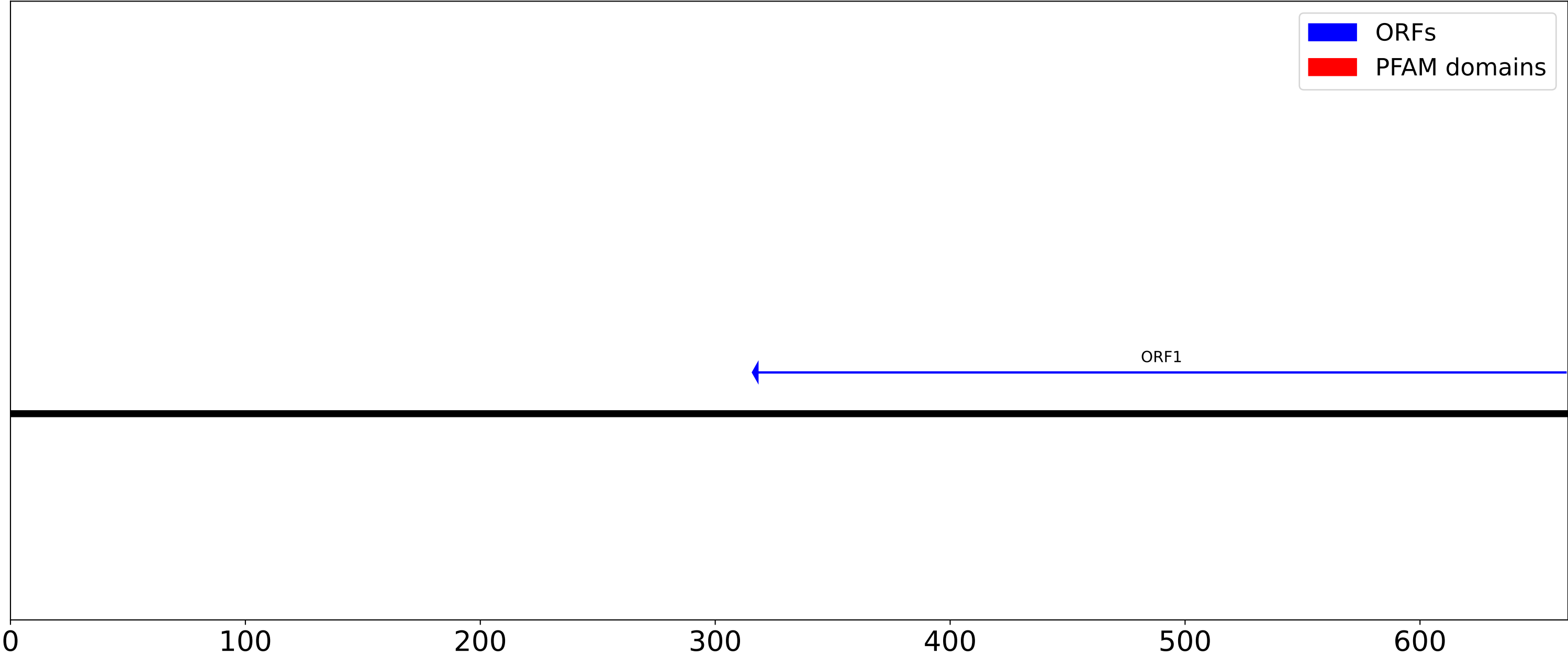


Before TEtrimmer 663 bp

After TEtrimmer ORF and PFAM domain plot



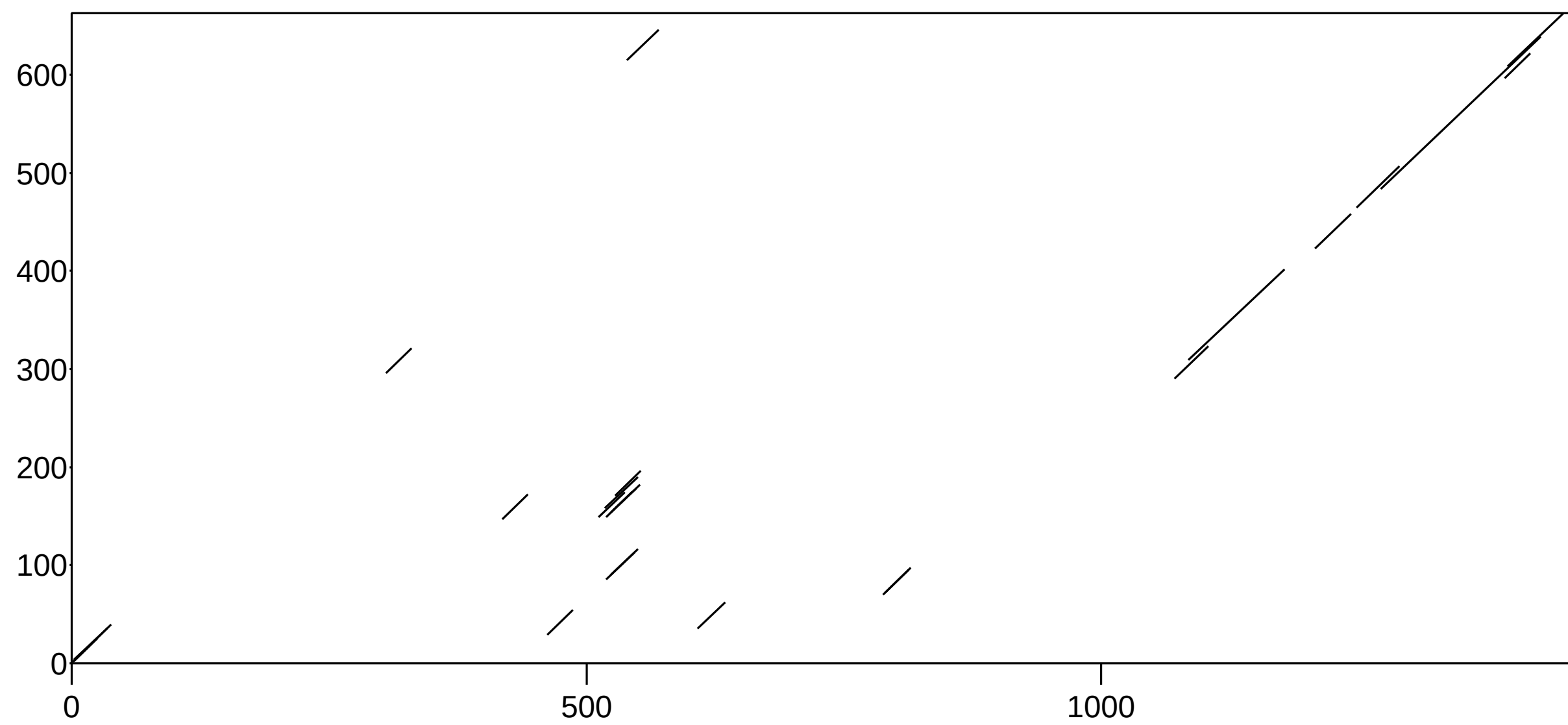
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)