

Start crop Point

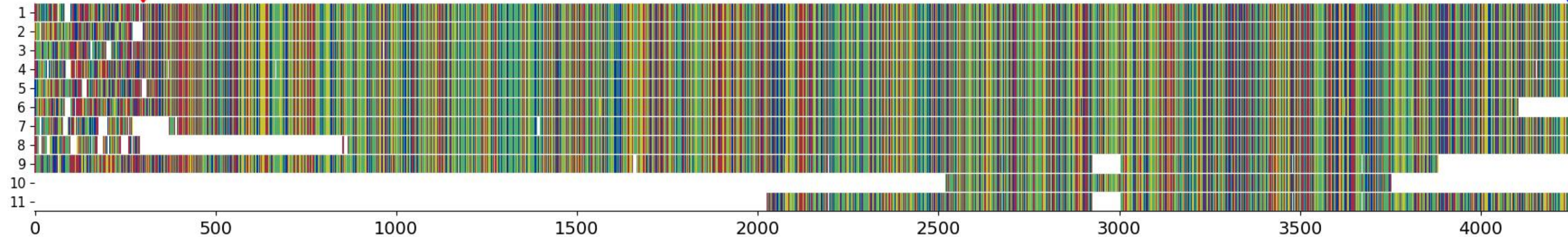
End crop Point

MSA length = 3946



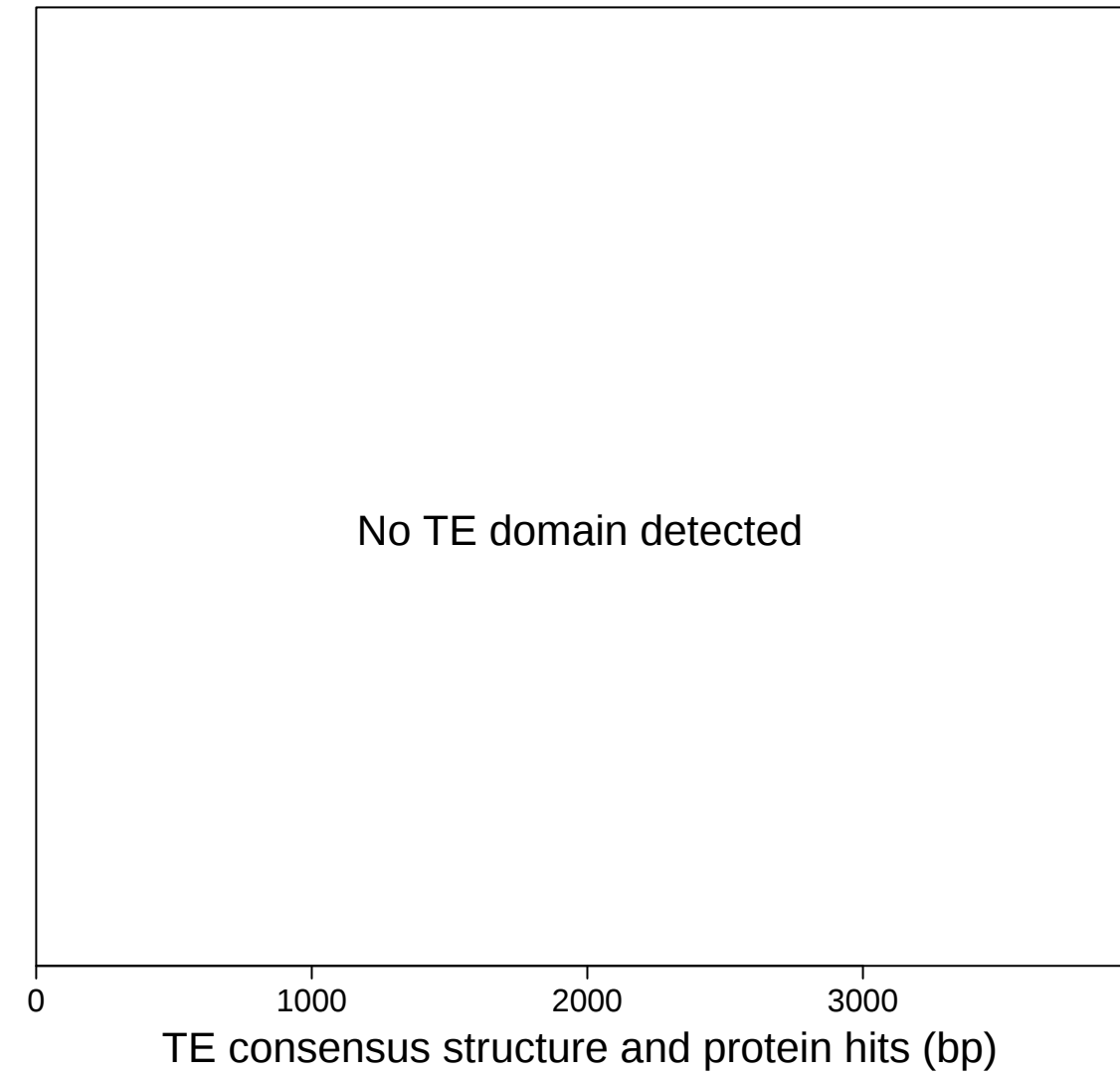
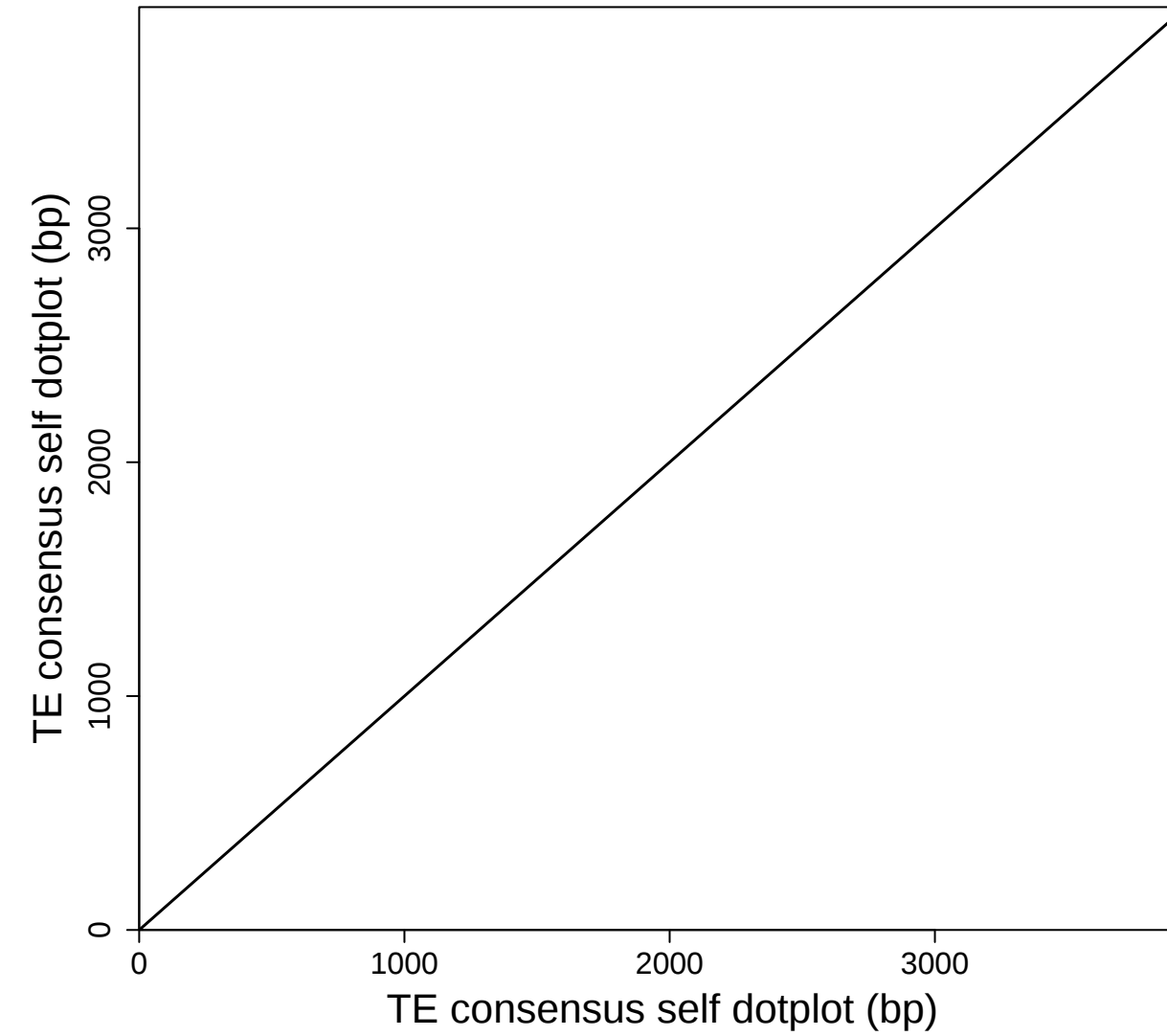
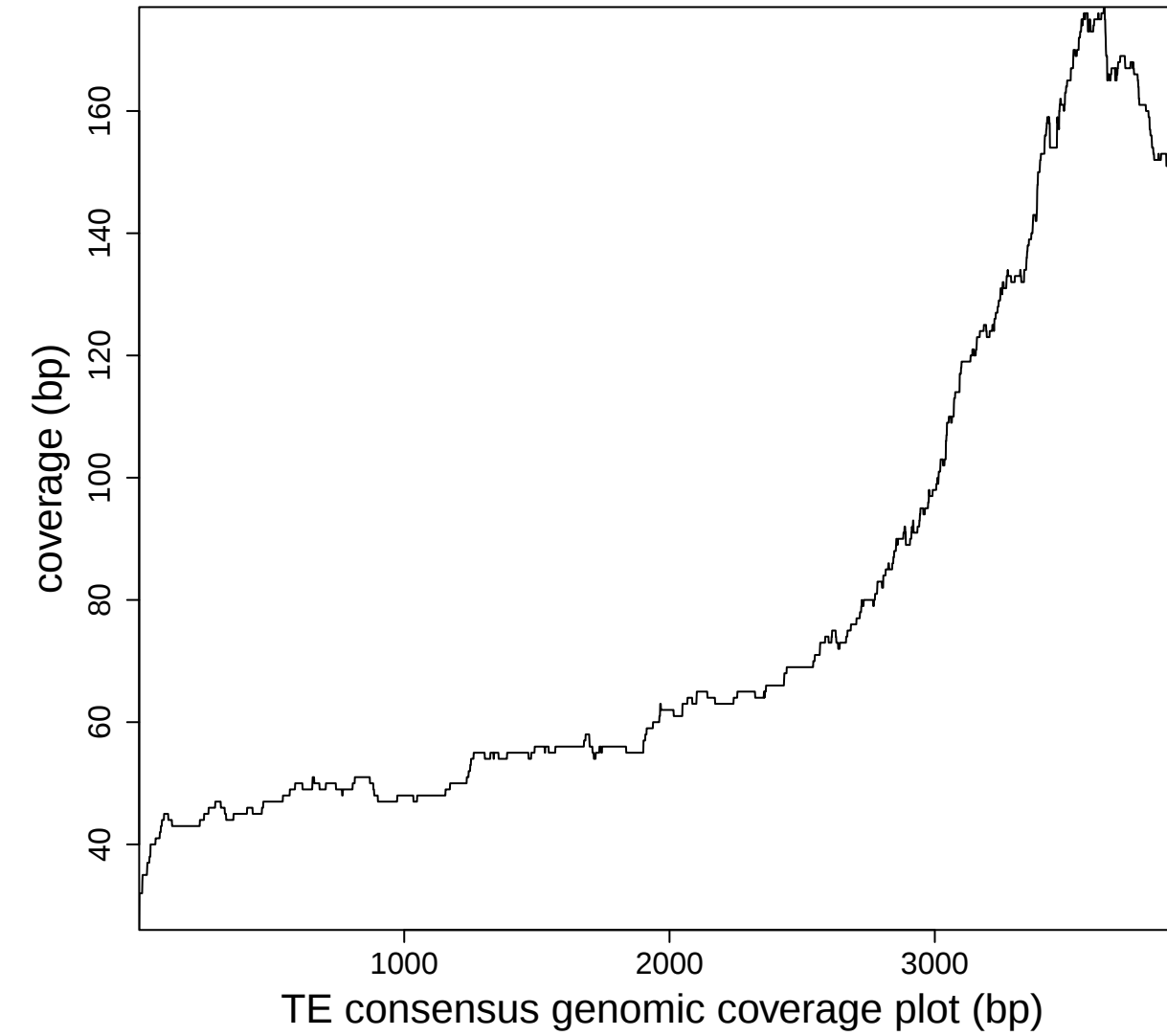
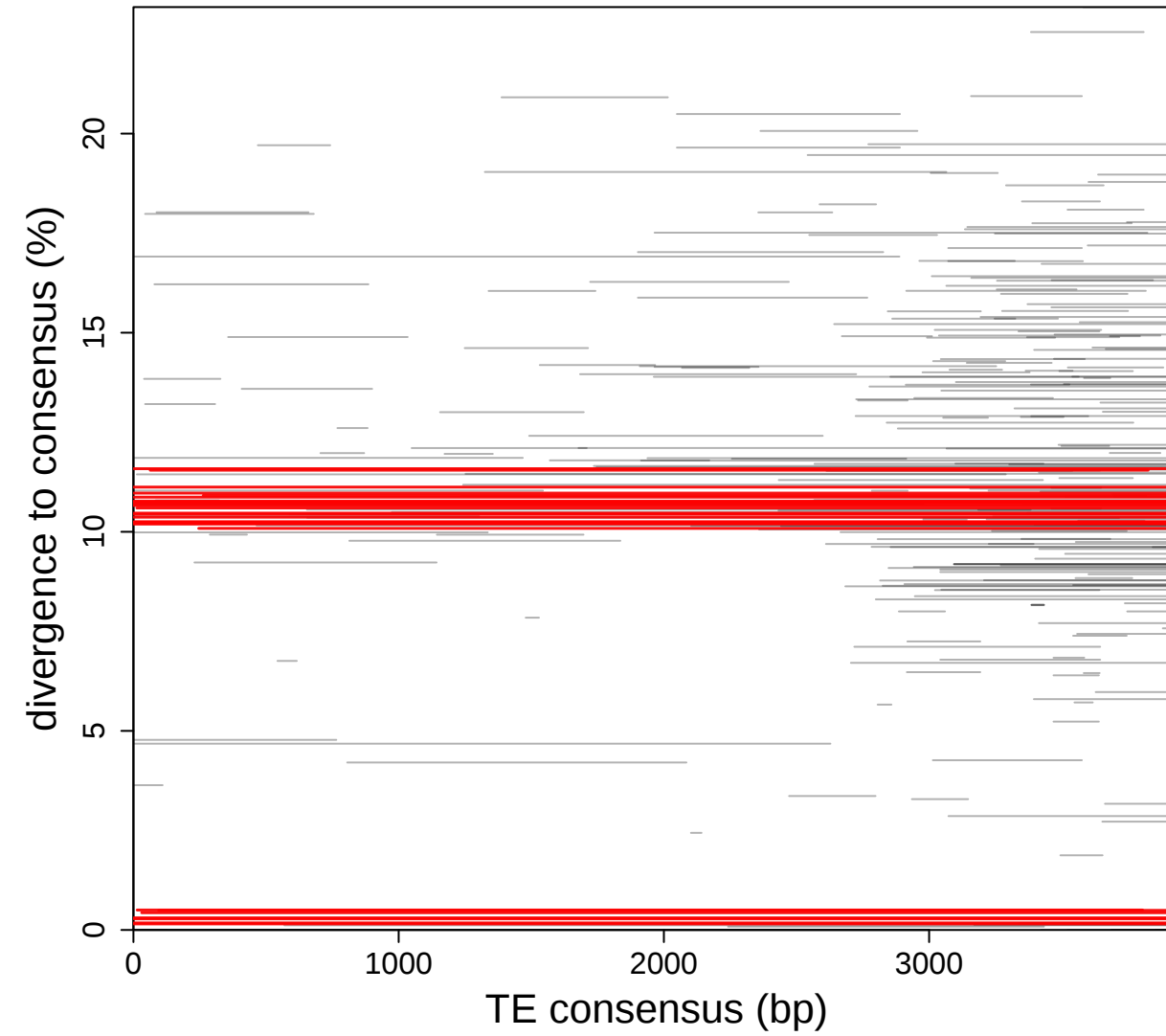
Start crop Point

End crop Point



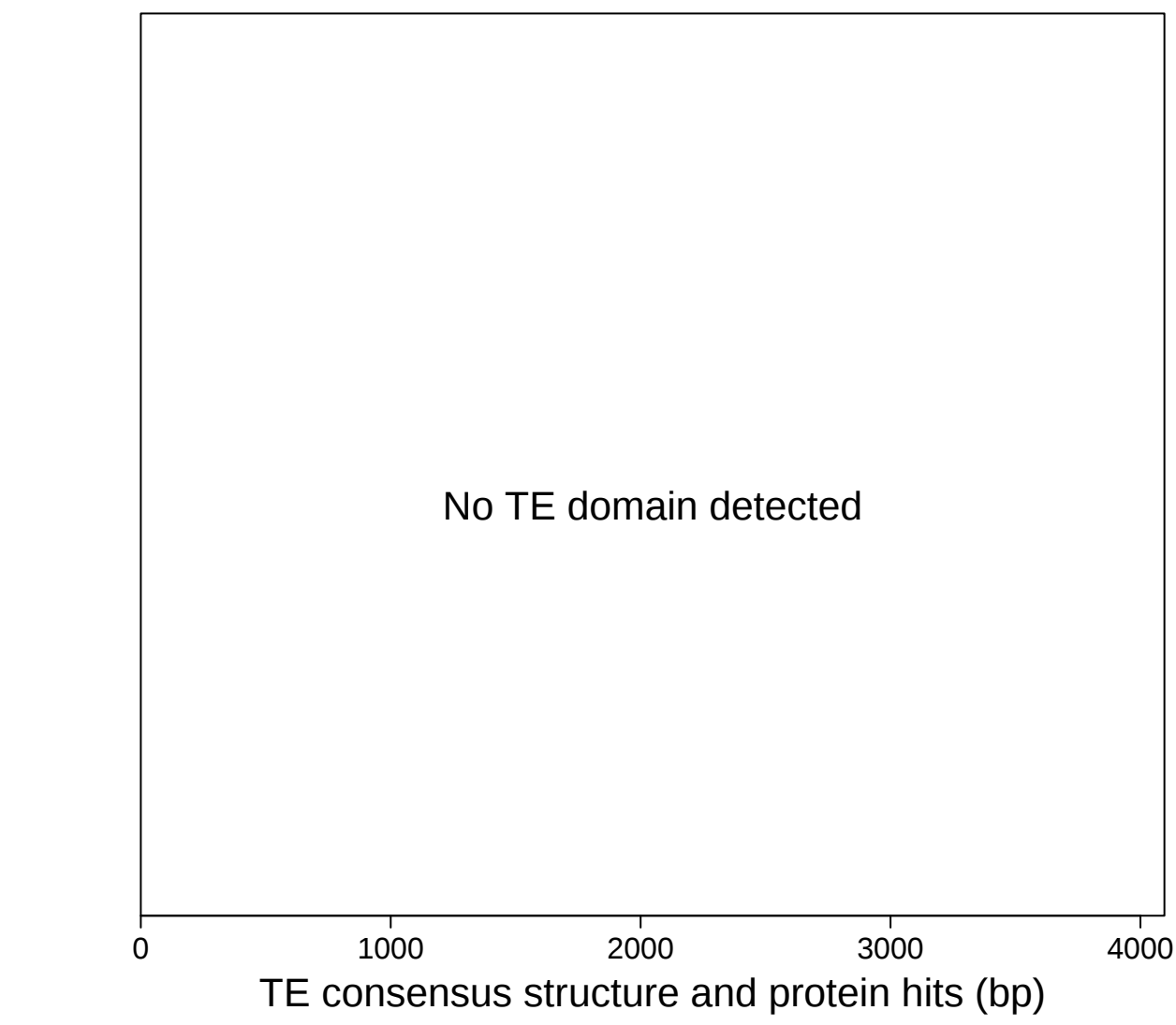
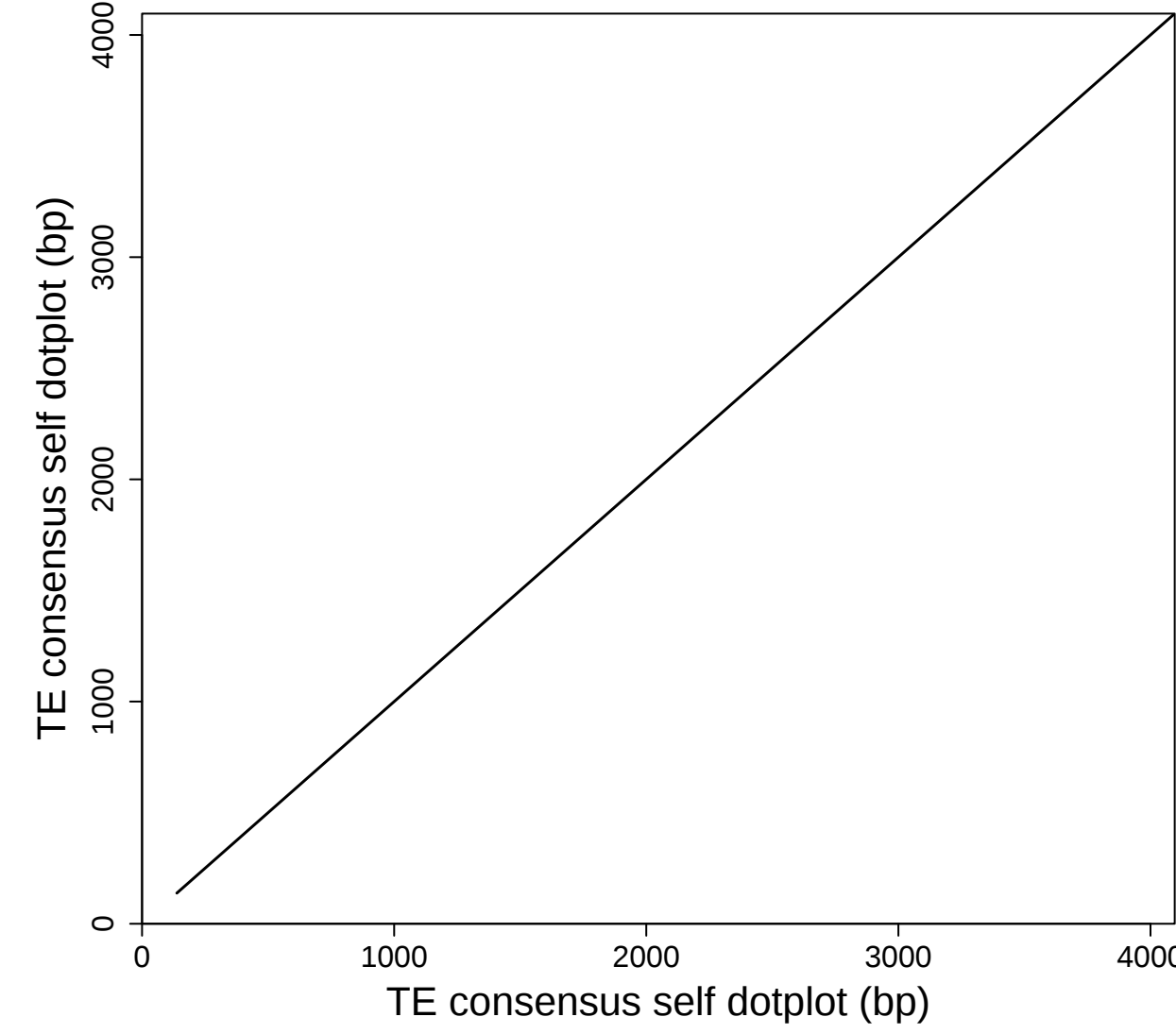
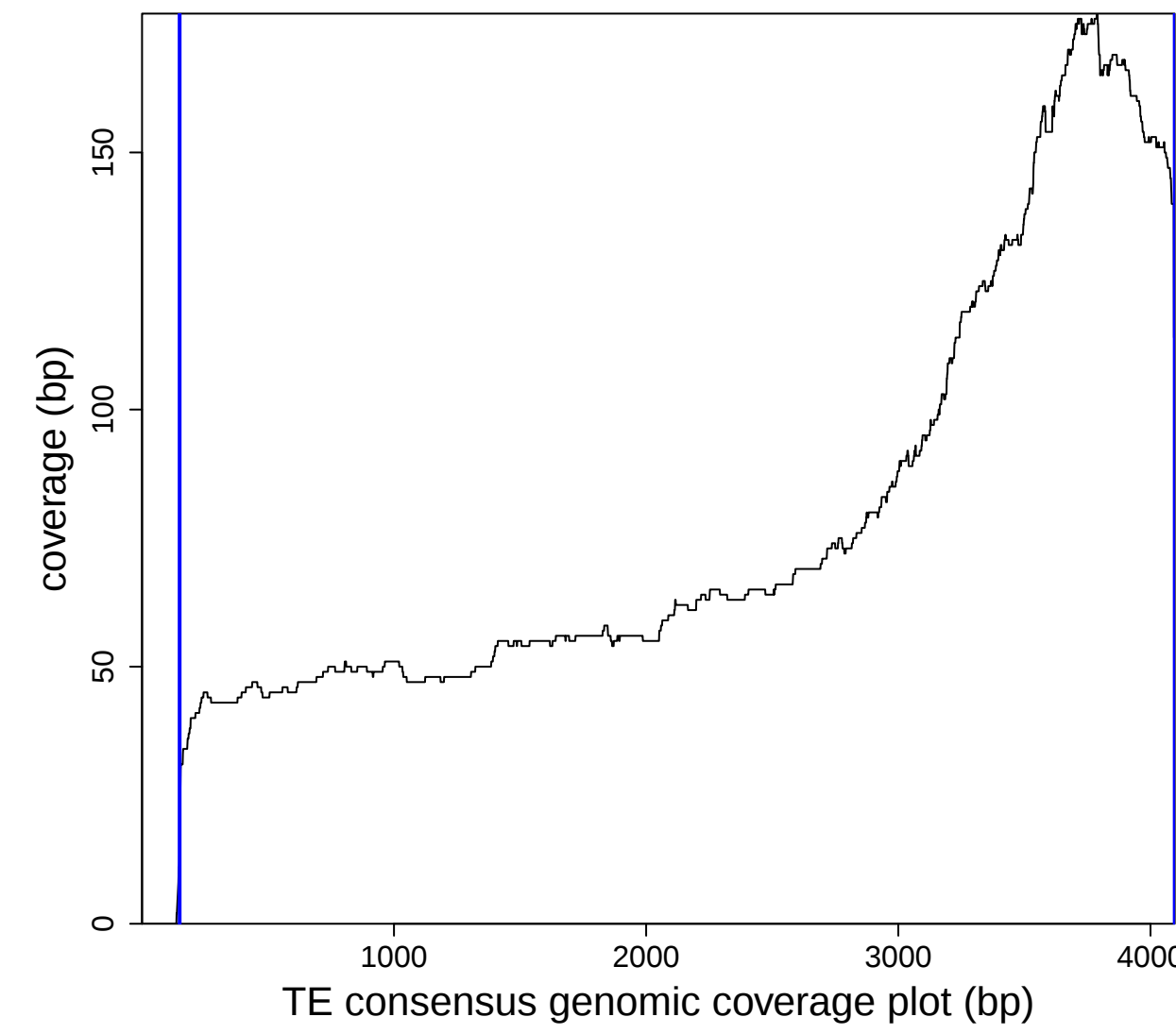
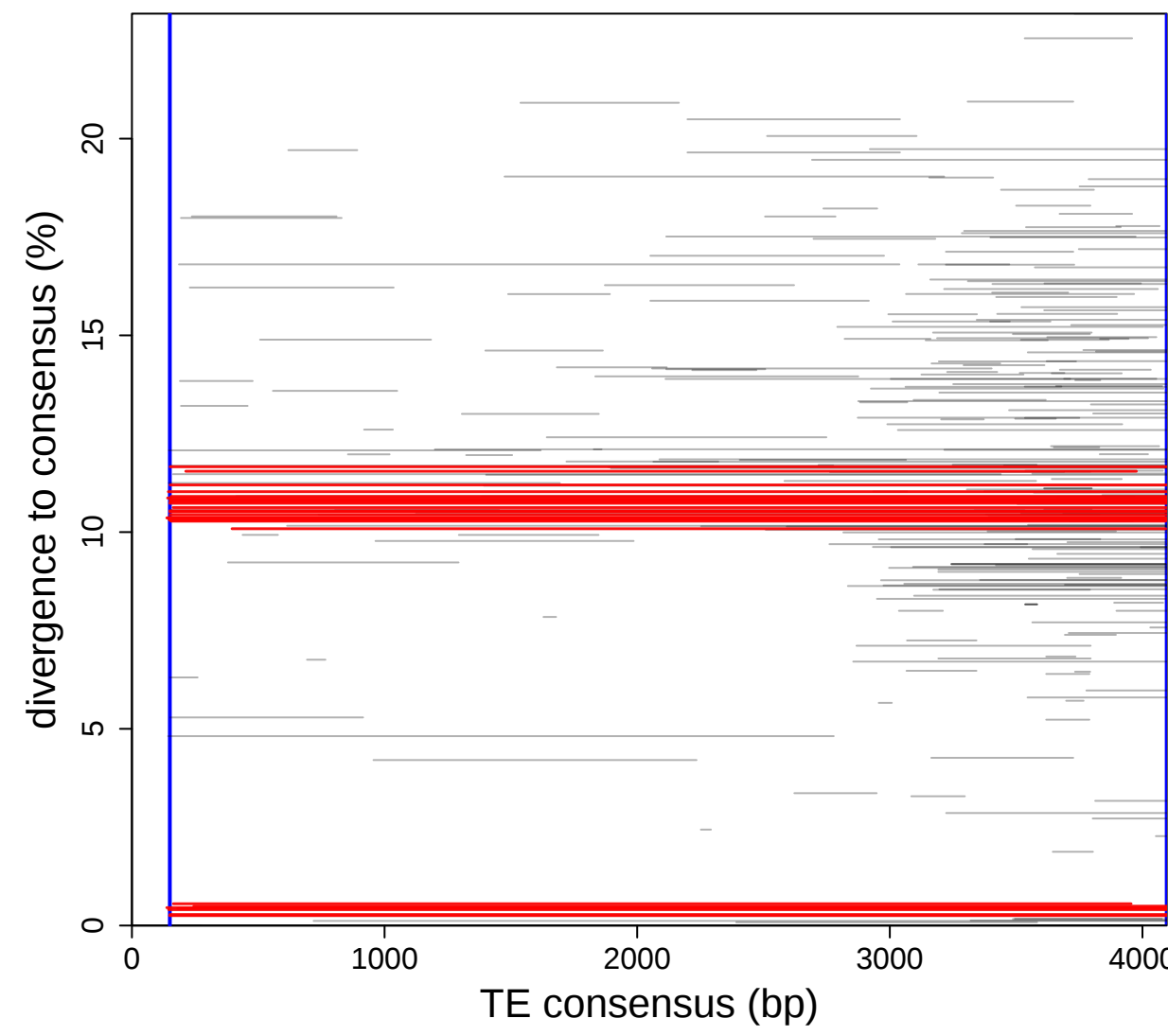
bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 3946bp; fragments: 294; full length: 29 (≥ 3551.4 bp)

After TEtrimmer 3946 bp



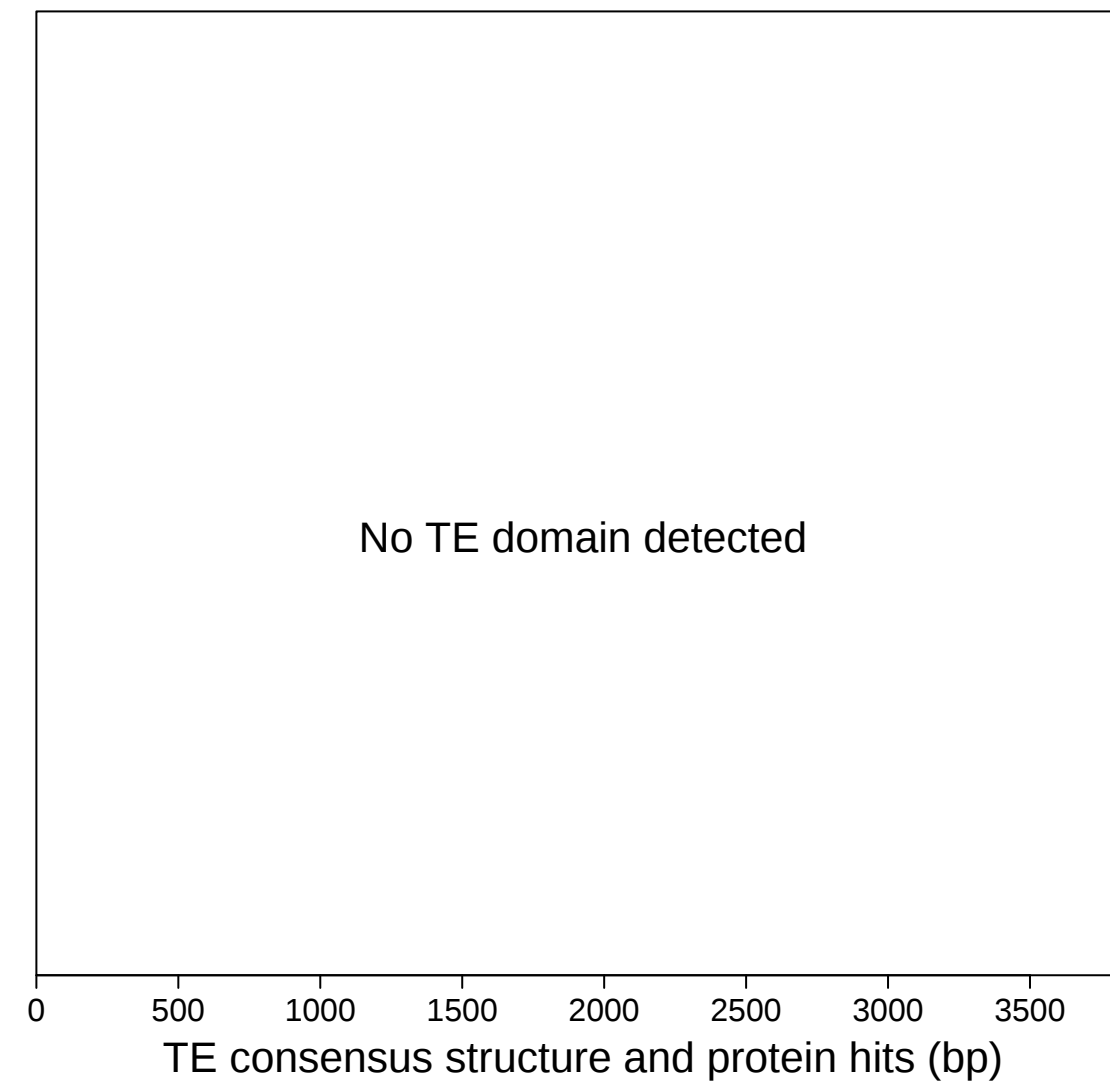
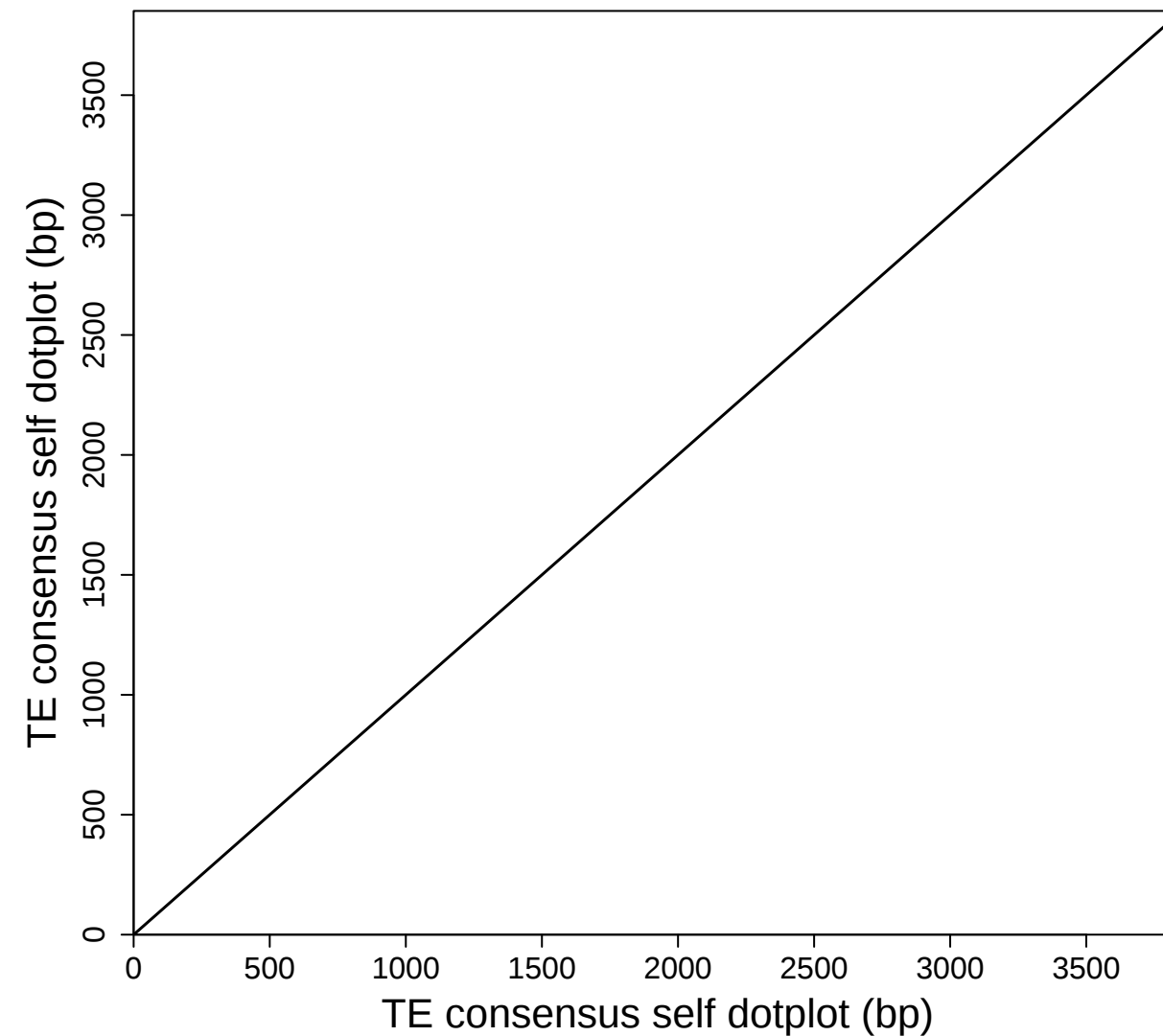
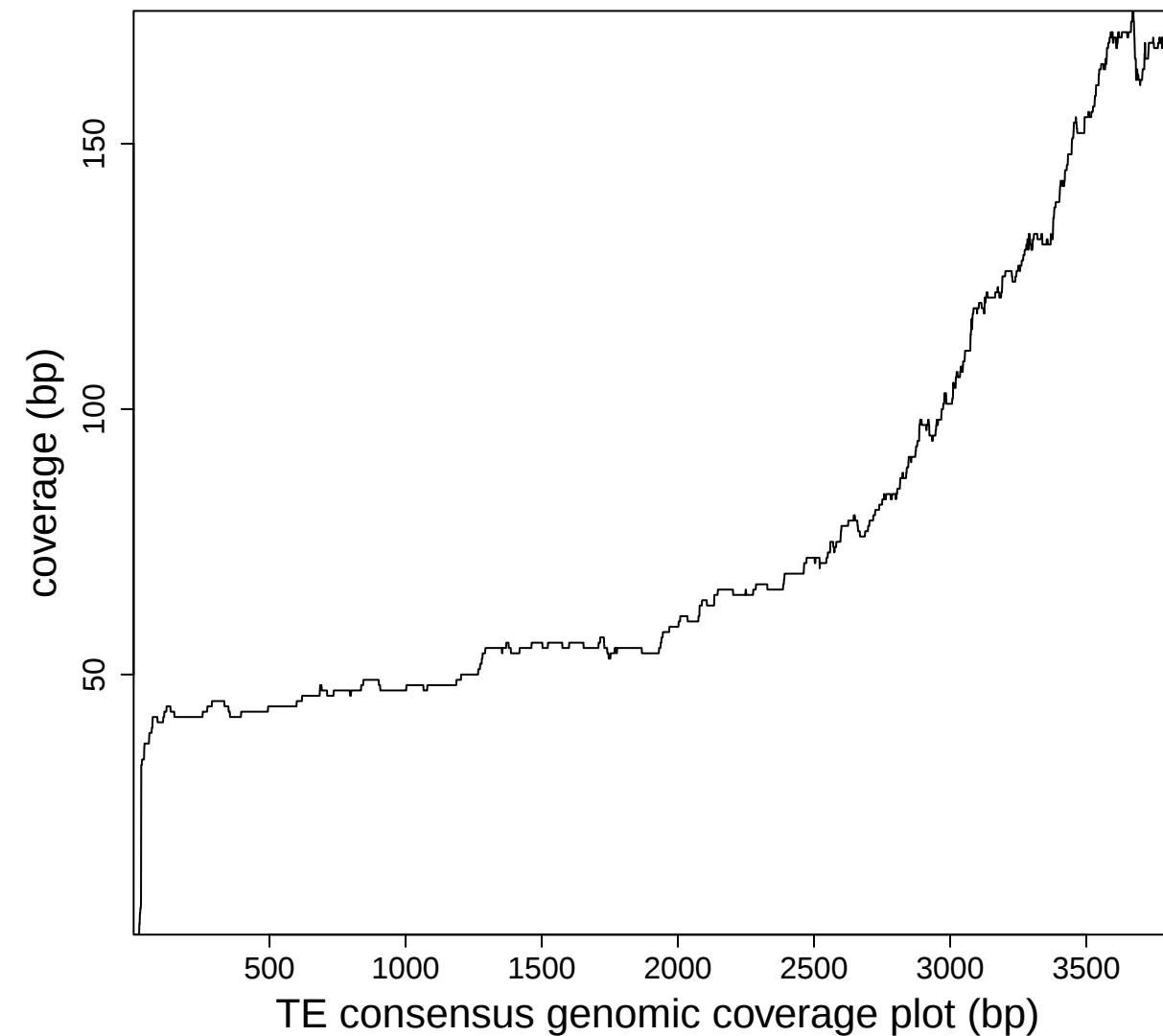
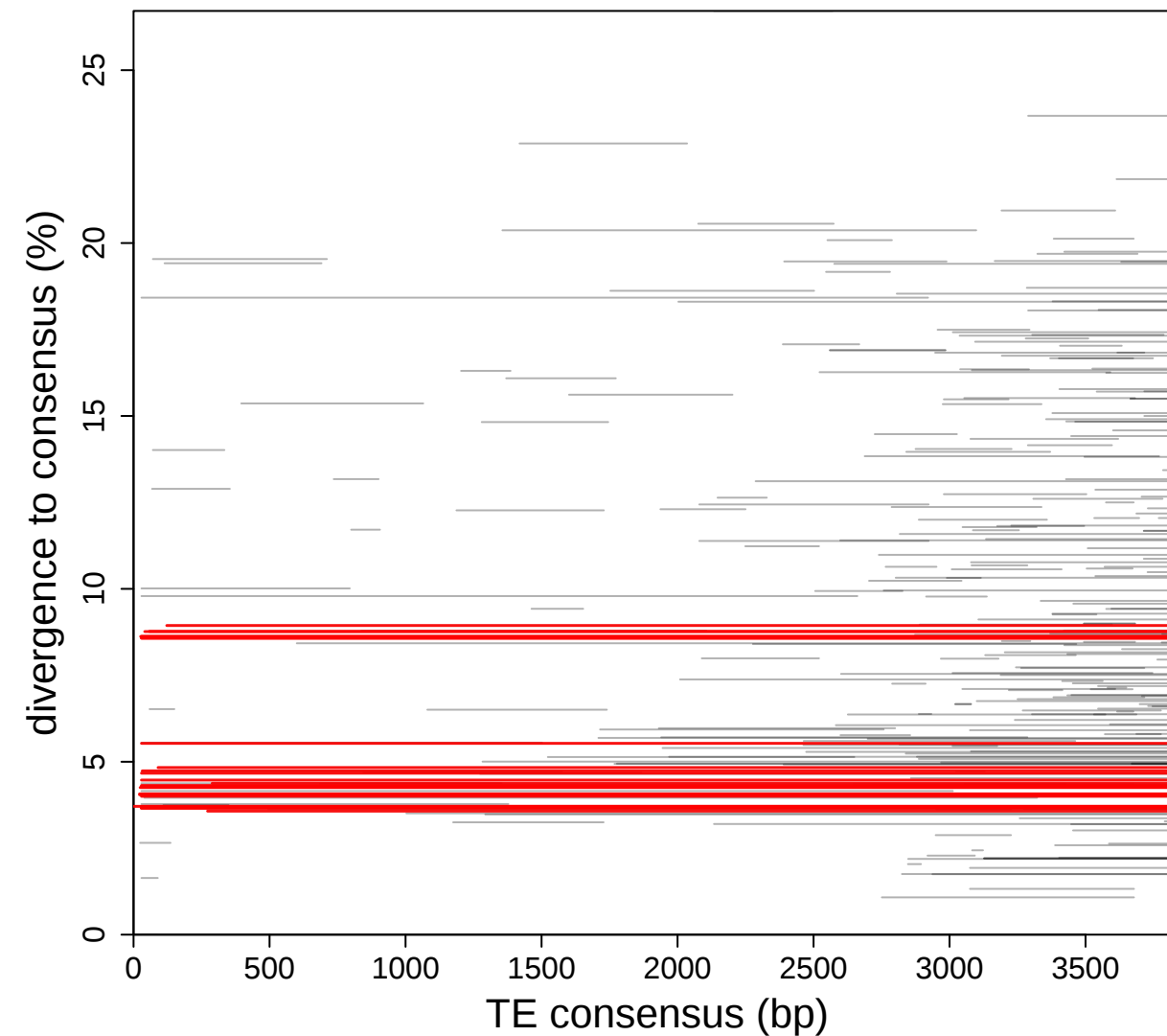
66.fasta.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.1
size: 4096bp; fragments: 294; full length: 28 (≥ 3686.4 bp)

After TEtrimmer Extended plot Blue lines are boundaries

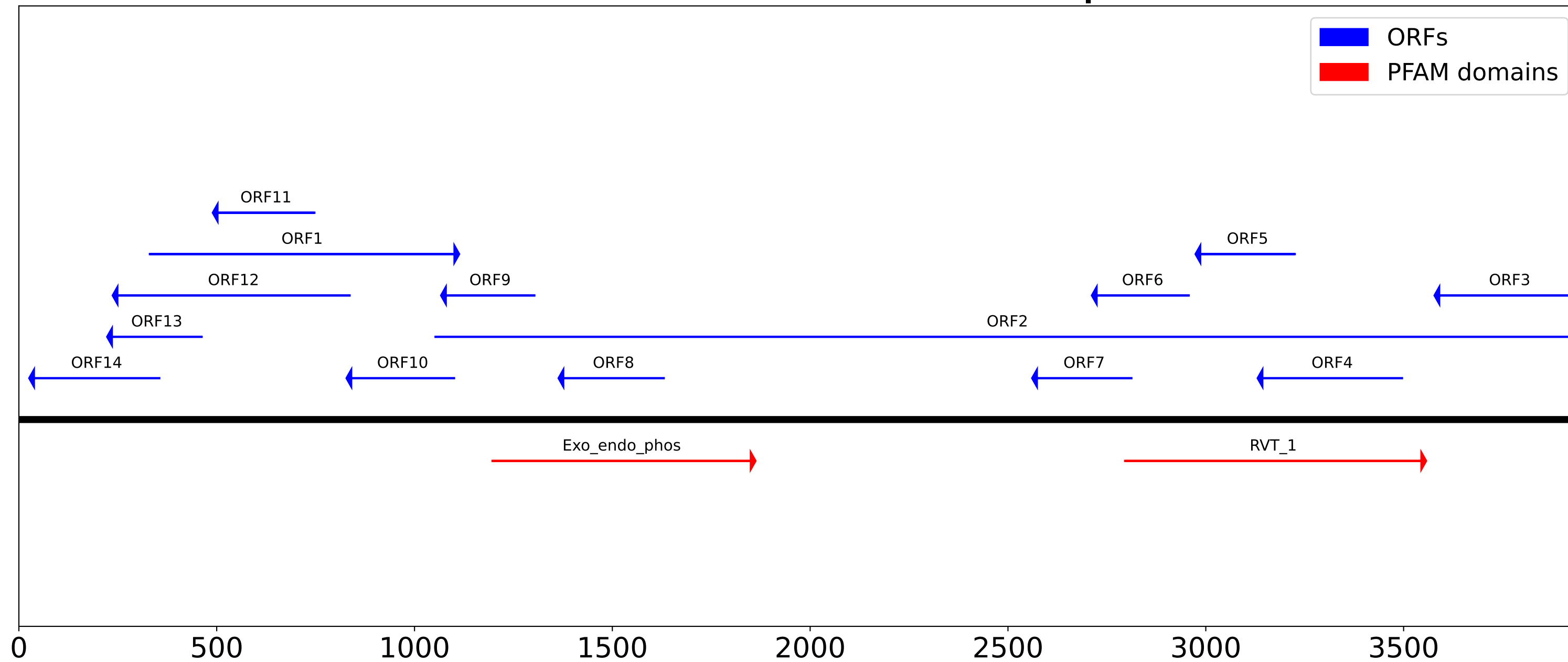


TE: rnd_5_family_12266
size: 3851bp; fragments: 301; full length: 29 (≥ 3465.9 bp)

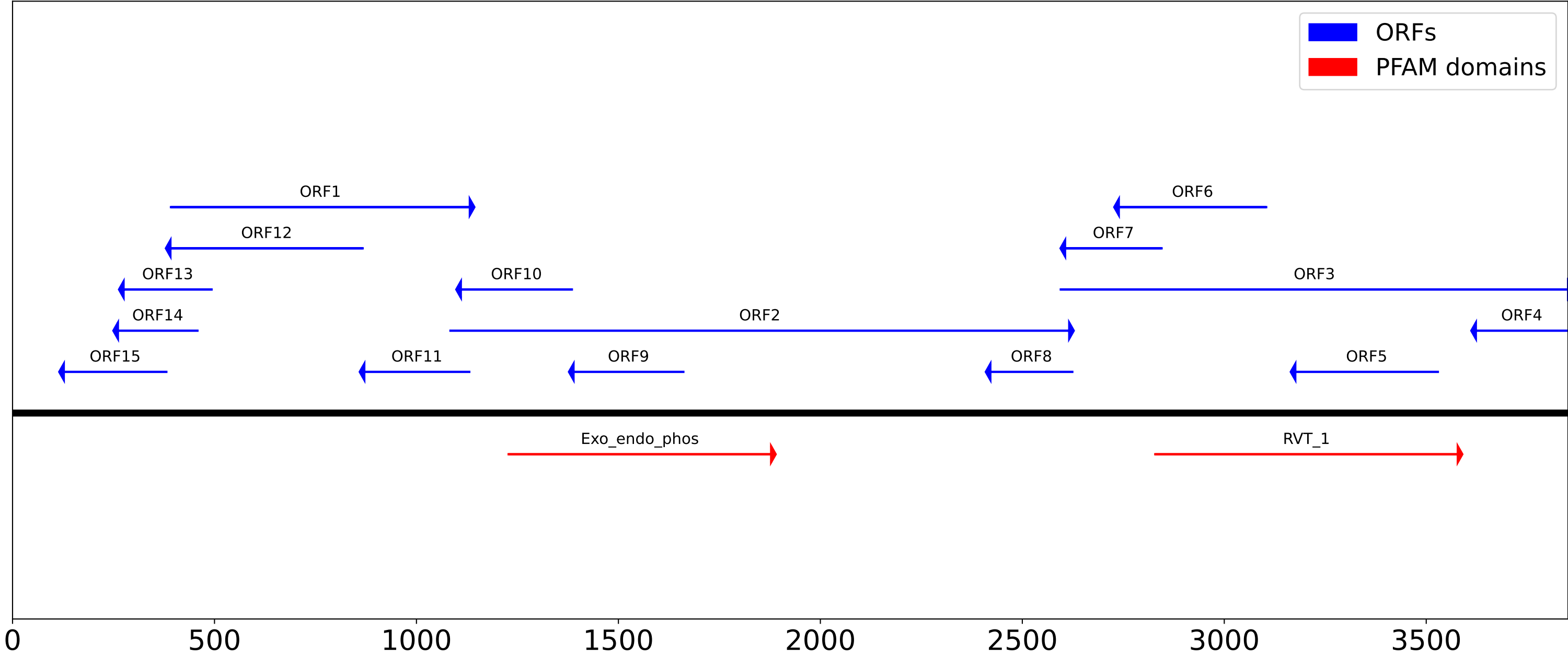
Before TEtrimmer 3851 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

