

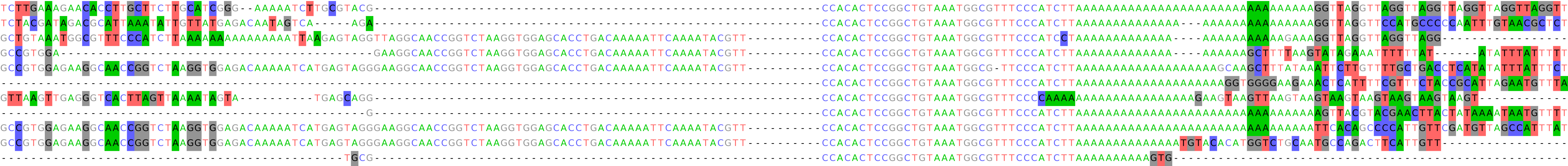
Start crop Point



End crop Point

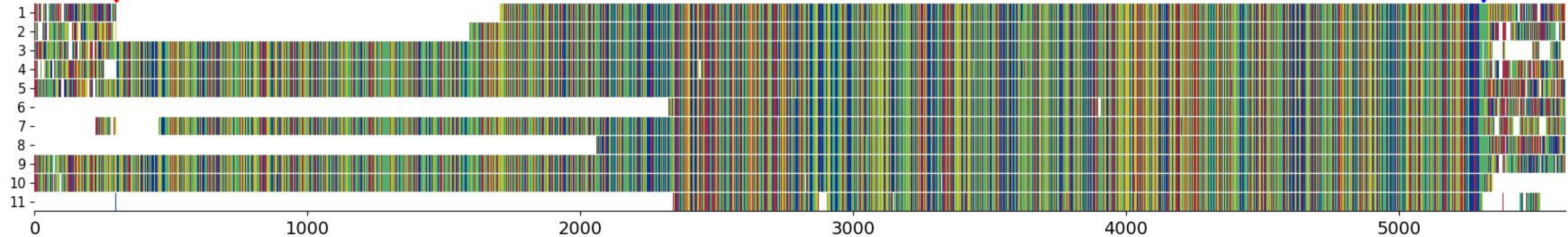


MSA length = 5012



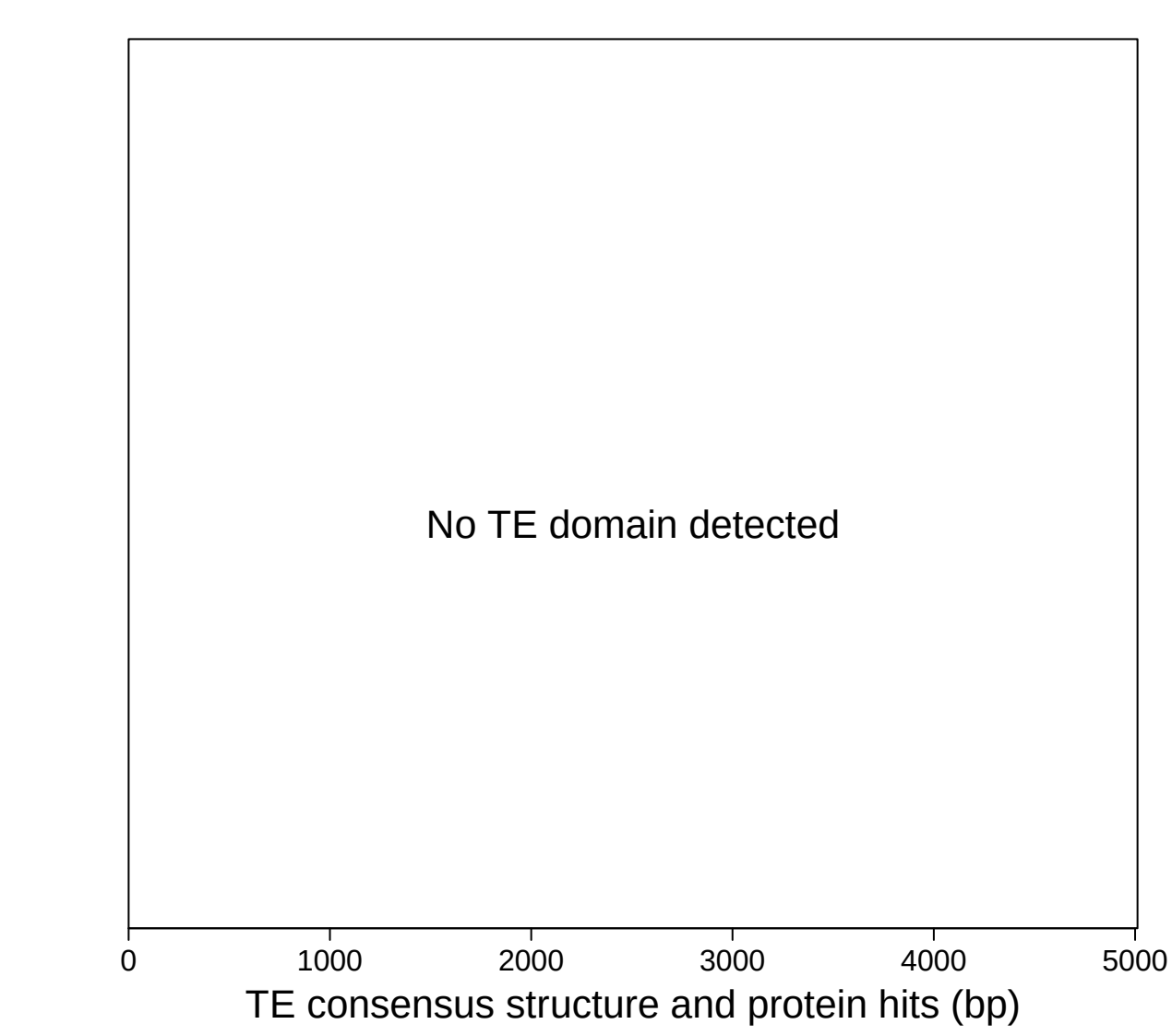
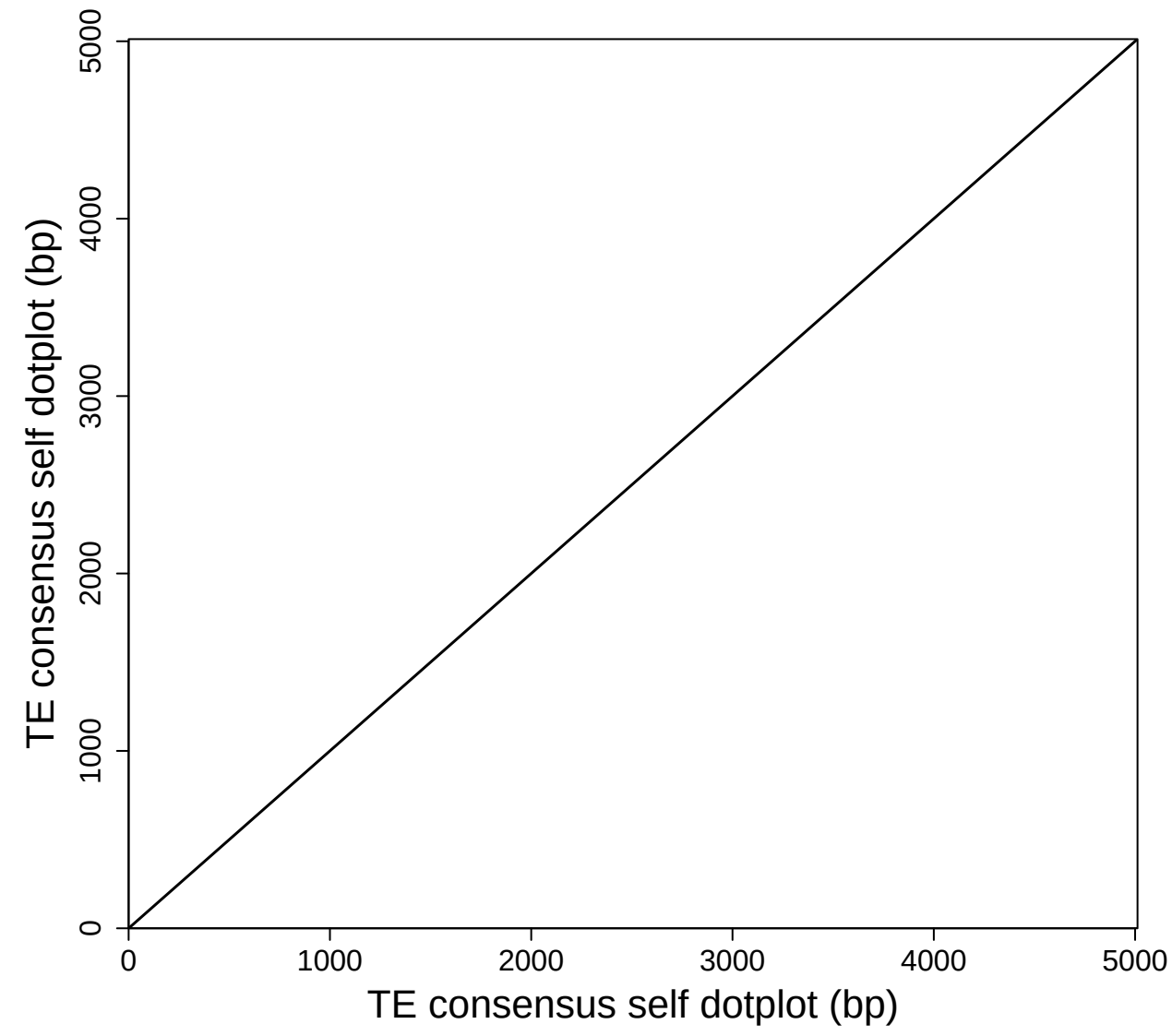
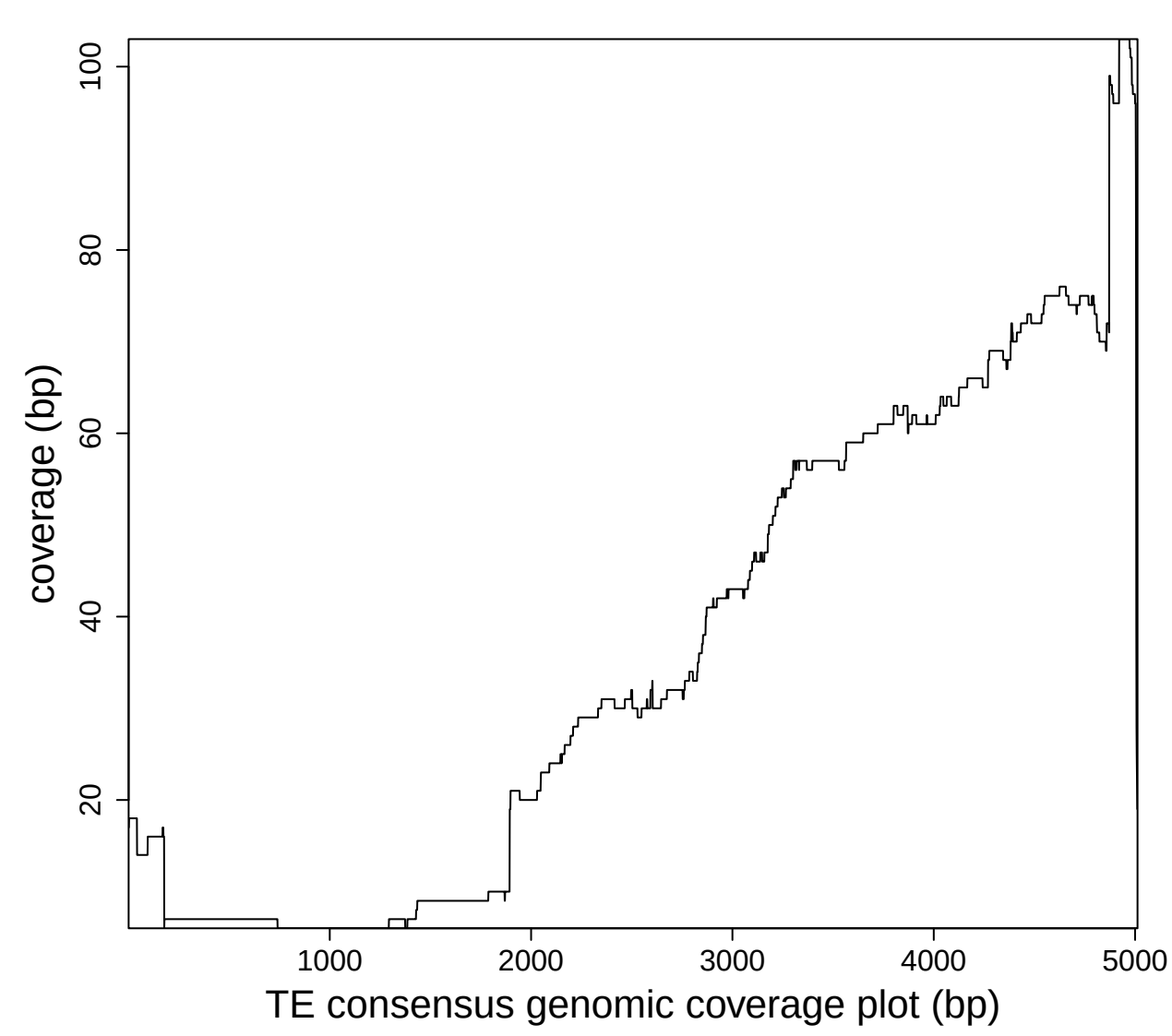
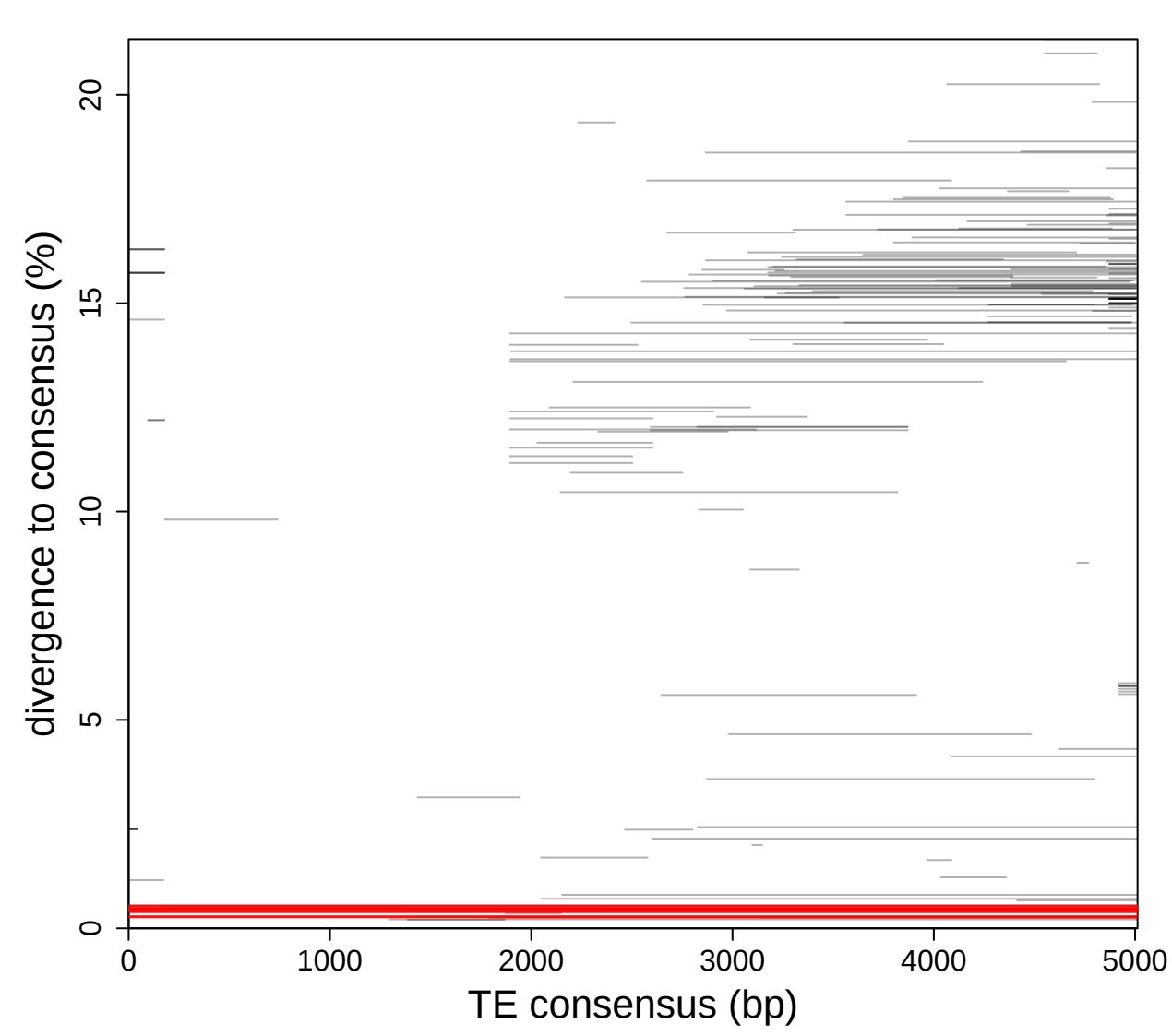
Start crop Point

End crop Point



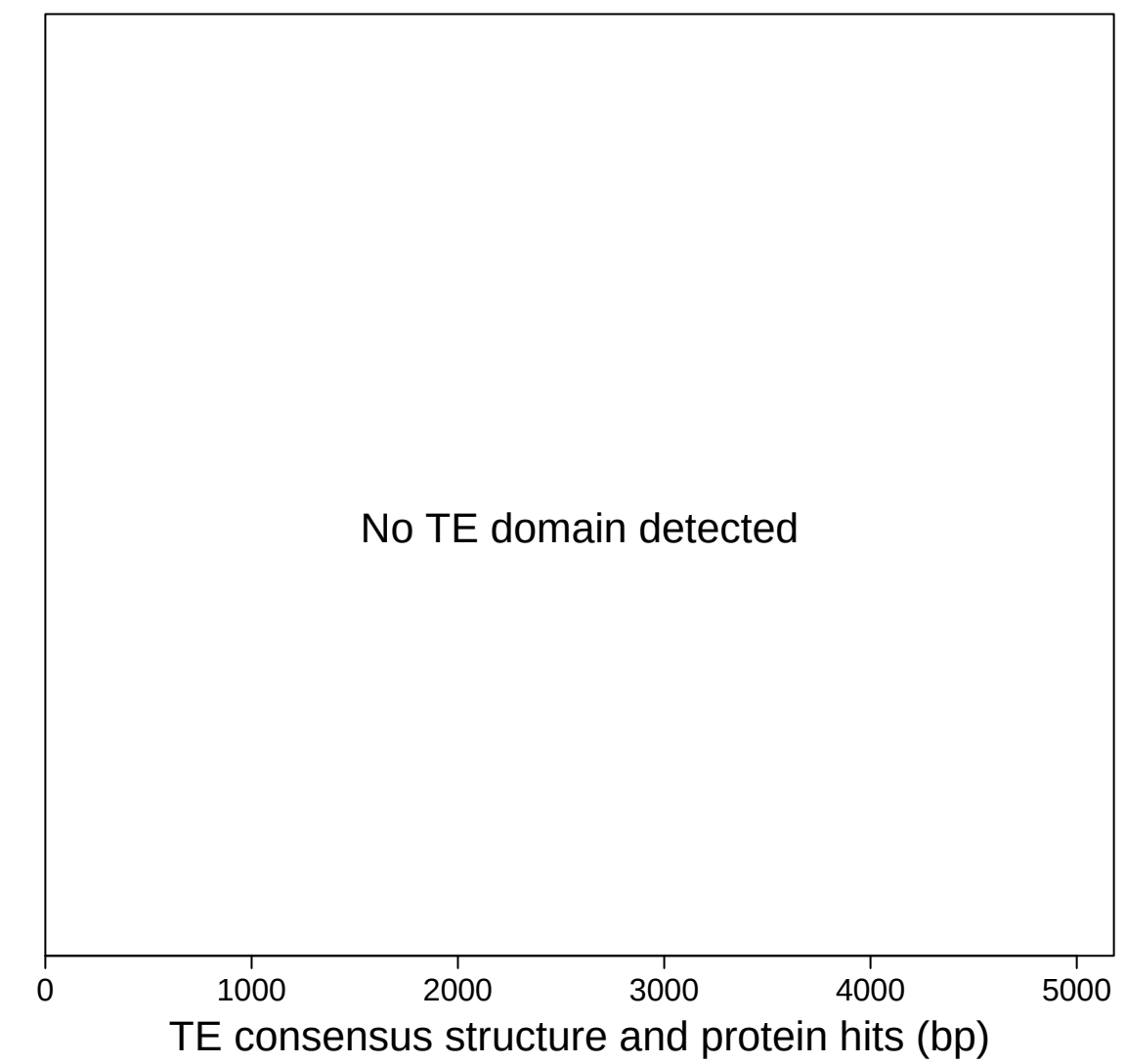
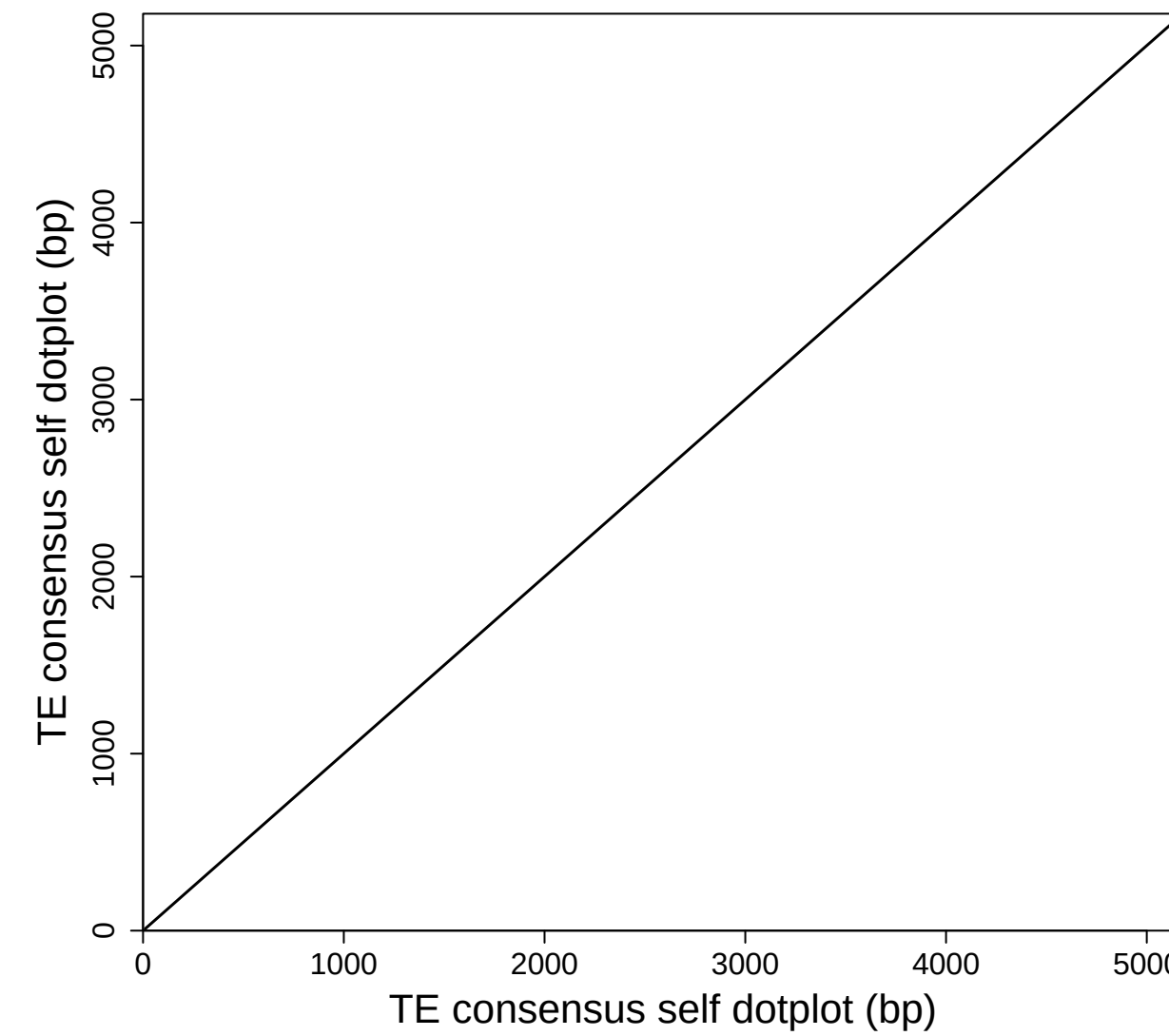
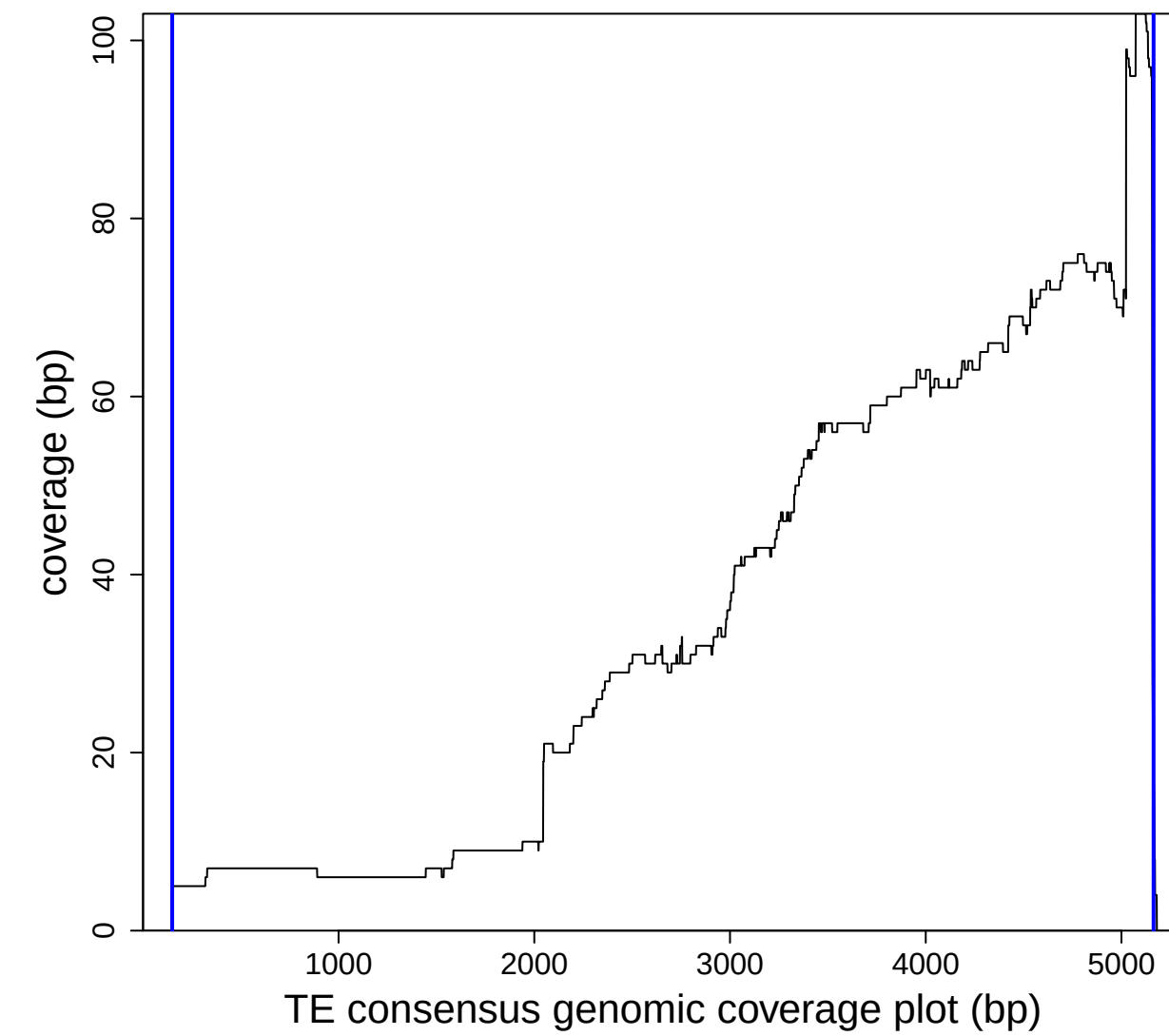
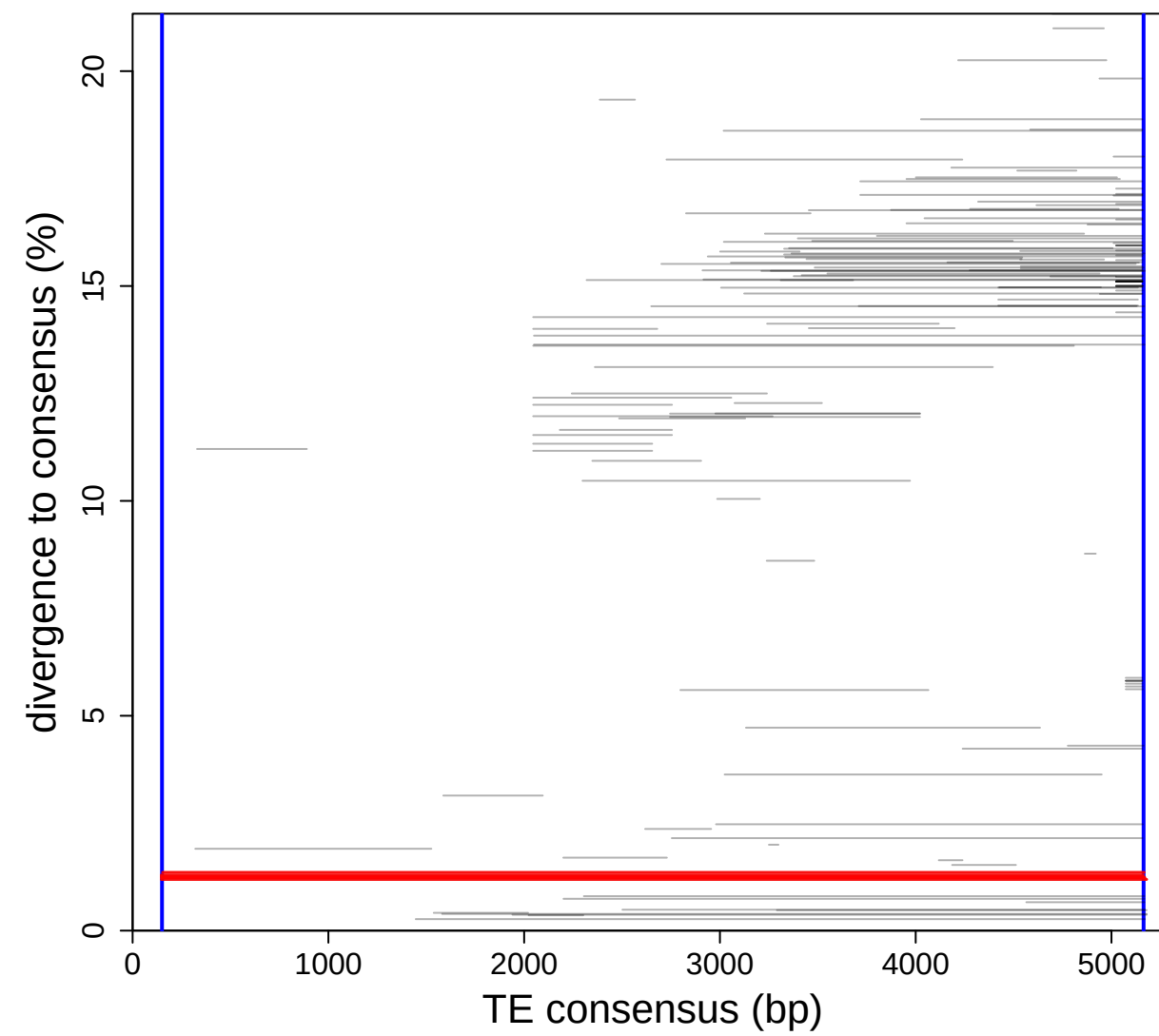
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 5012bp; fragments: 174; full length: 5 (>=4510.8bp)

After TETrimmer 5012 bp

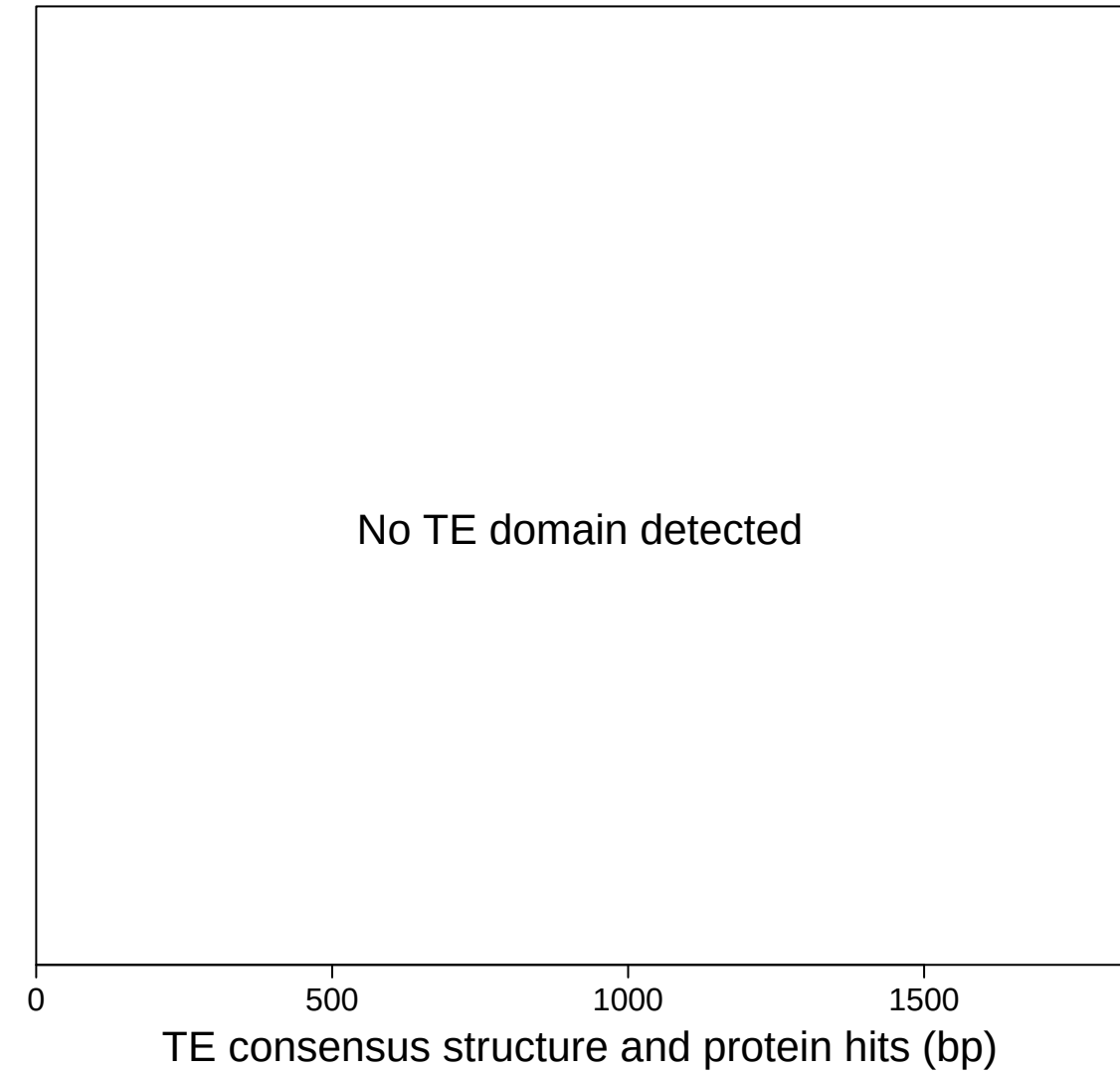
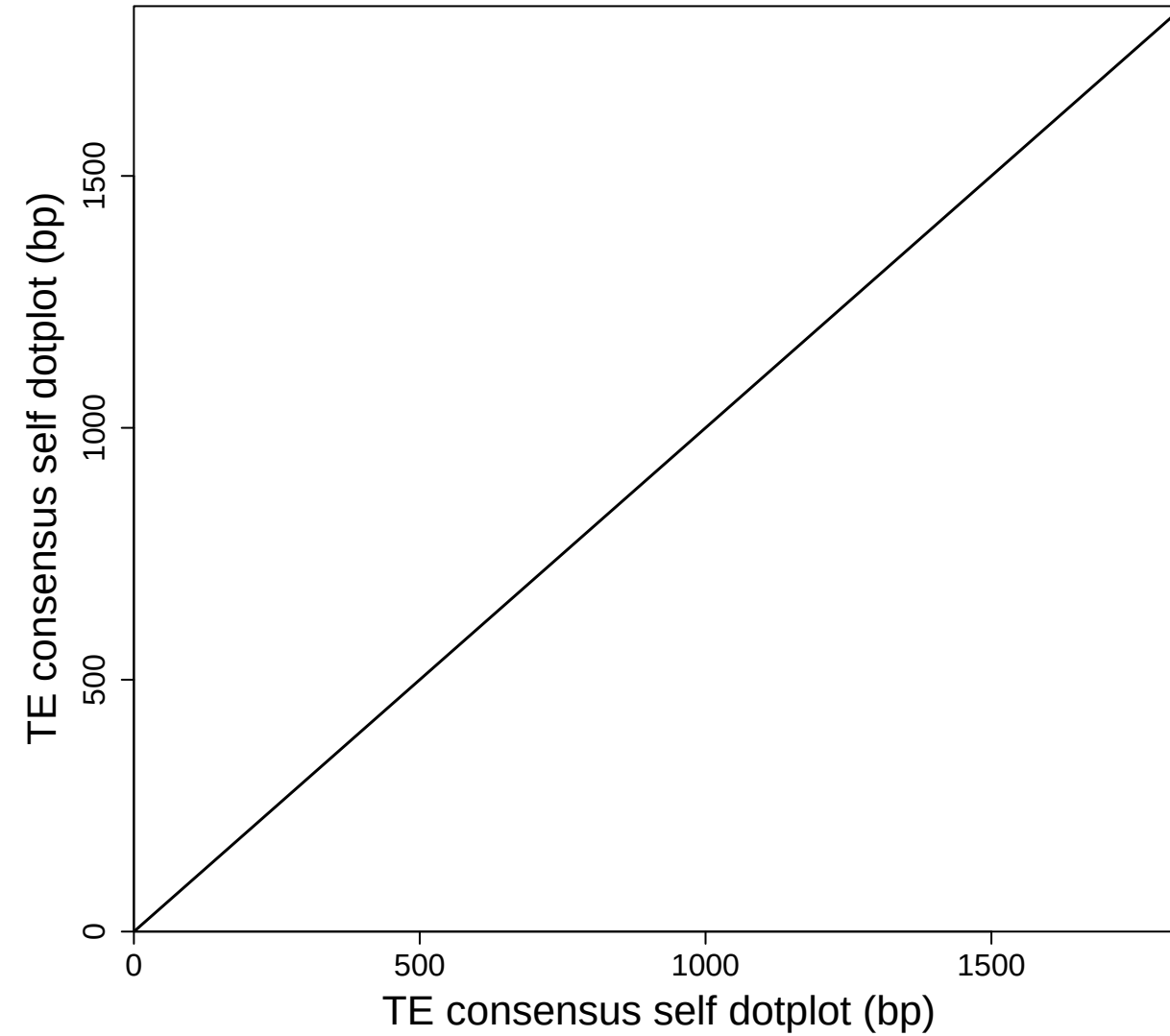
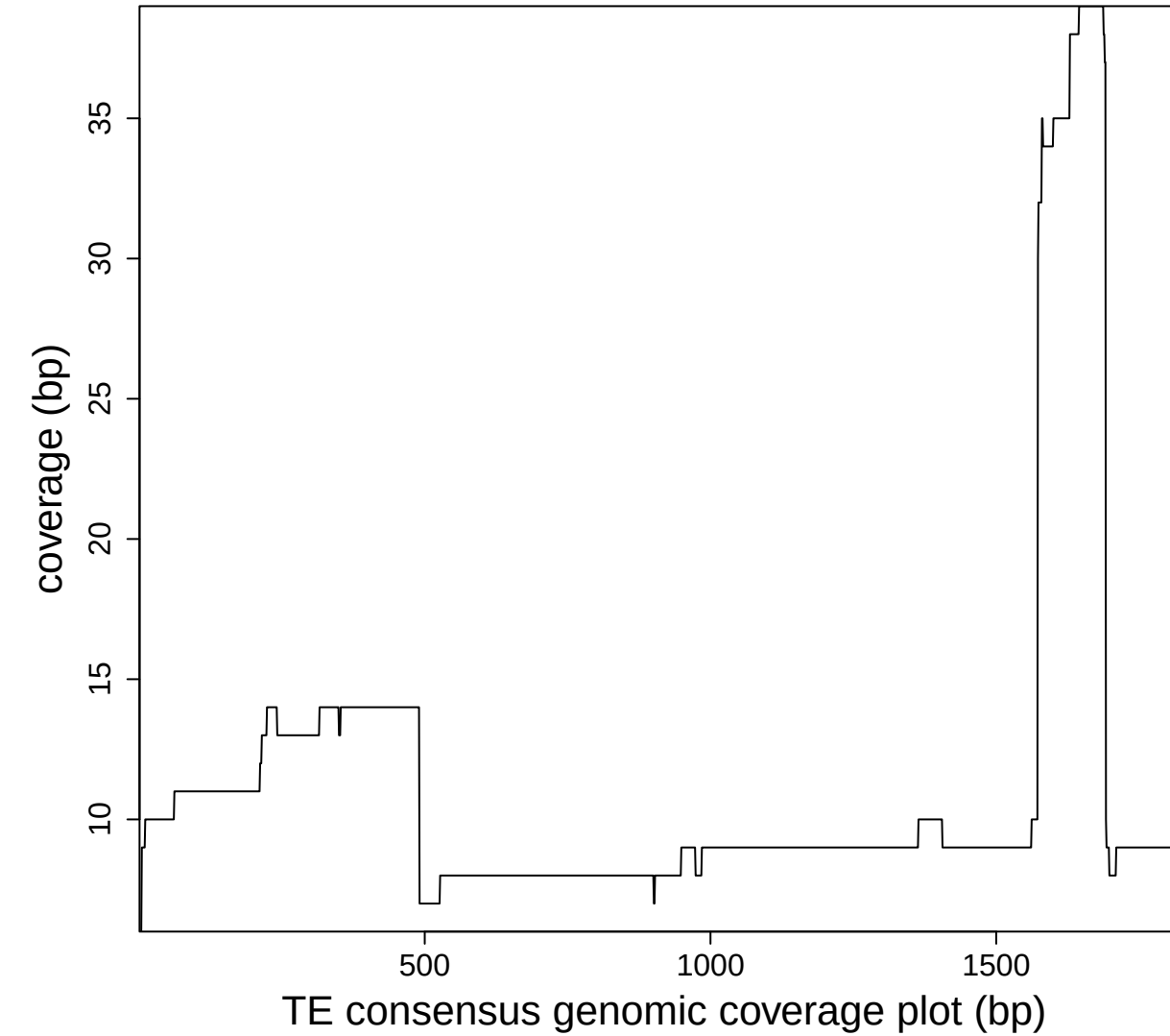
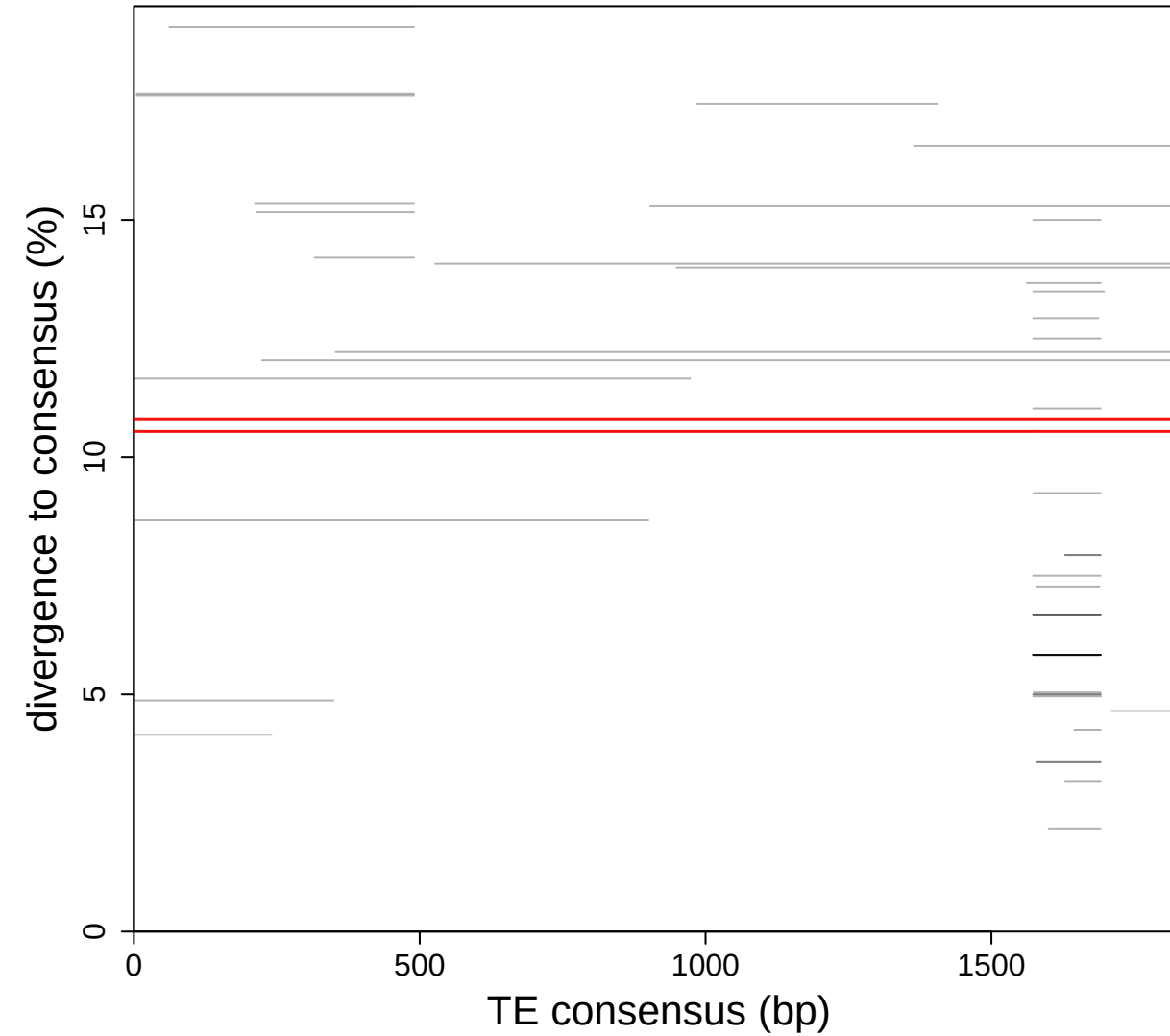


0.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 5314bp; fragments: 159; full length: 5 (>=4782.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

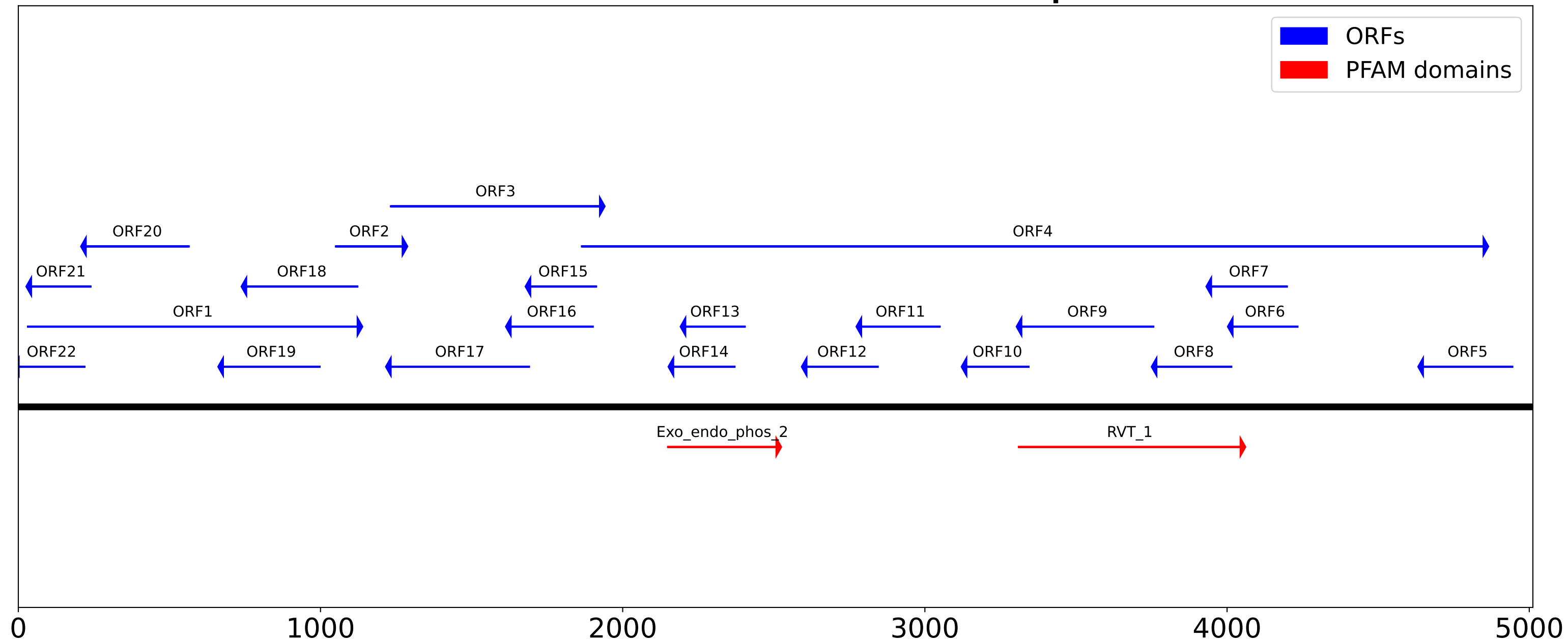


TE: rnd_5_family_240
size: 1837bp; fragments: 53; full length: 2 (>=1653.3bp)

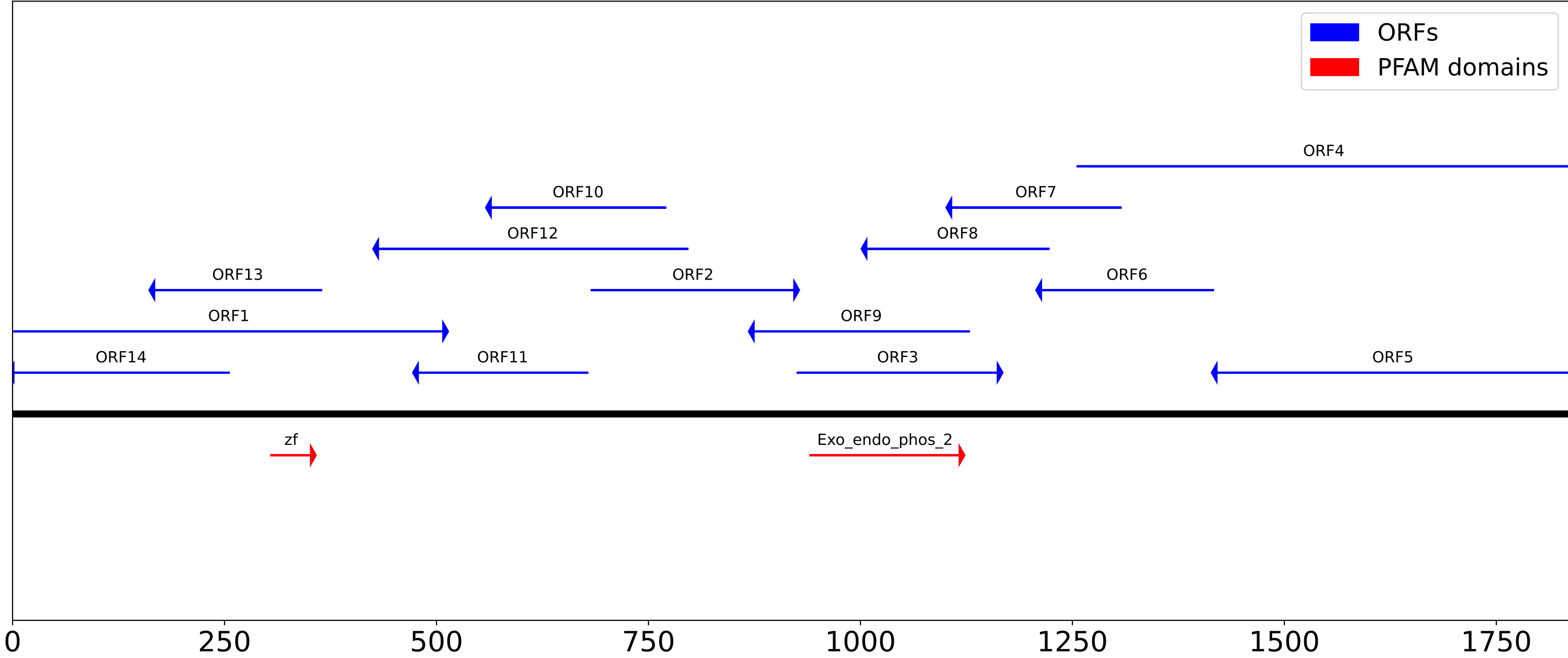


Before TEtrimmer 1837 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)

