

MSA length = 4231

Start crop Point



End crop Point



Start crop Point

End crop Point

0

10

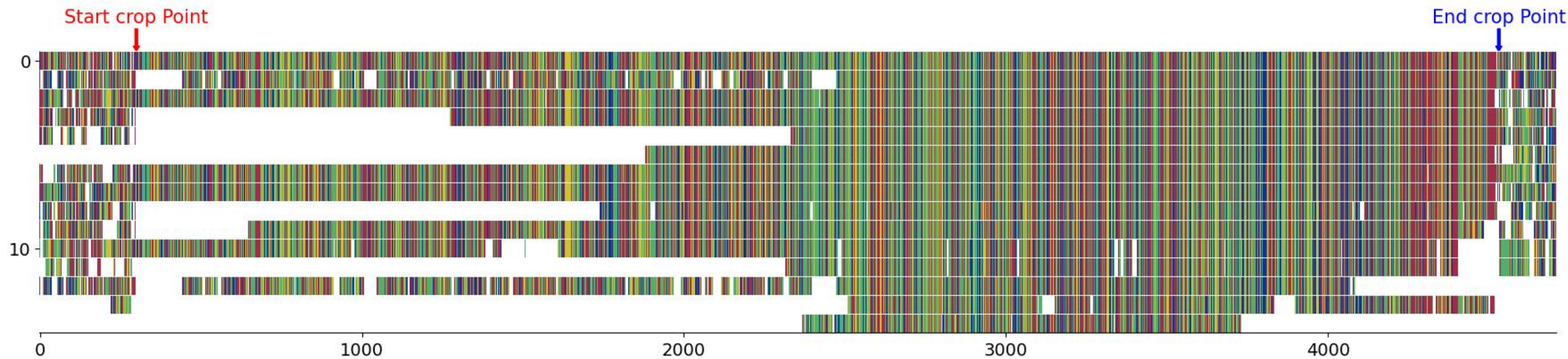
0

1000

2000

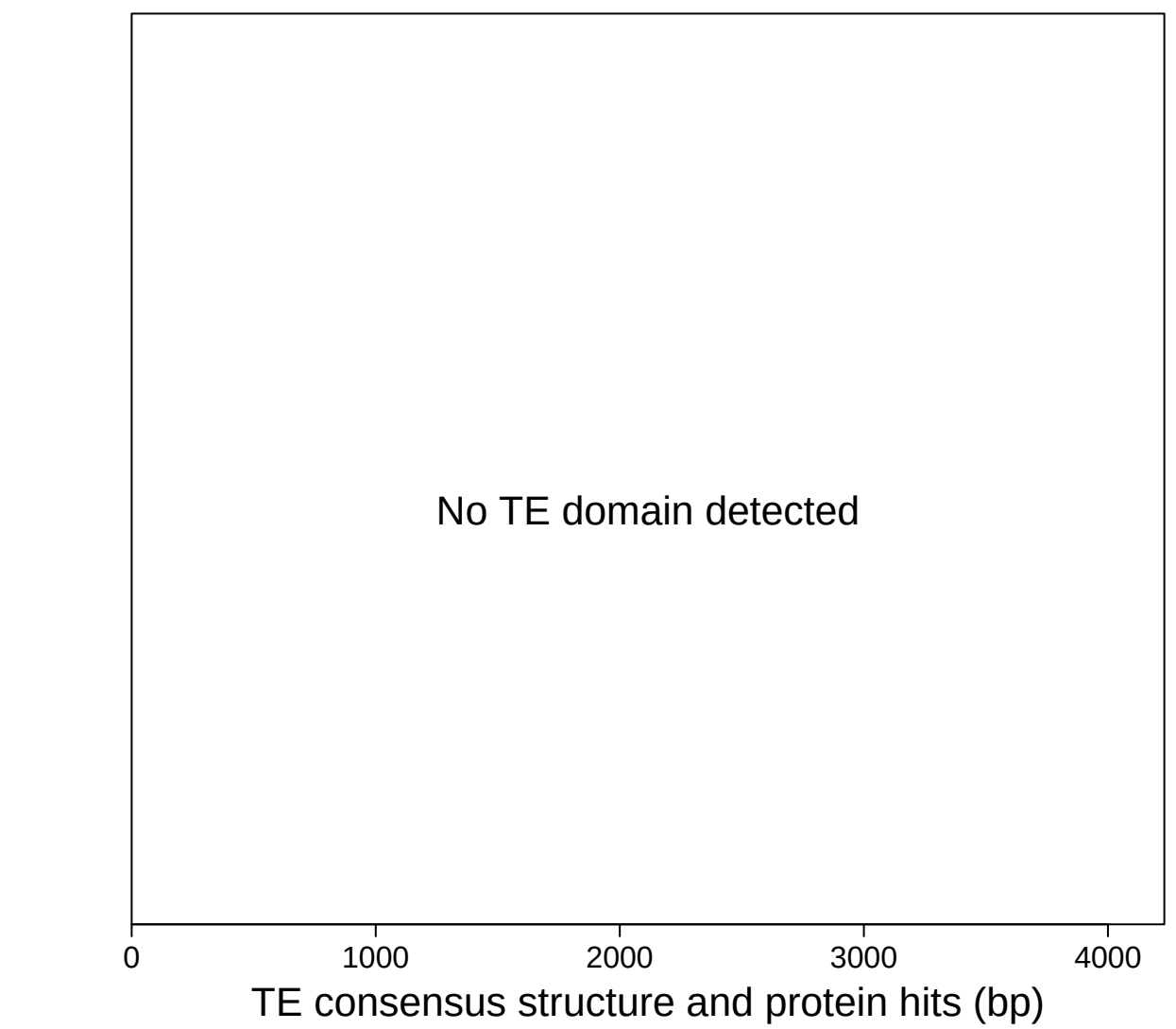
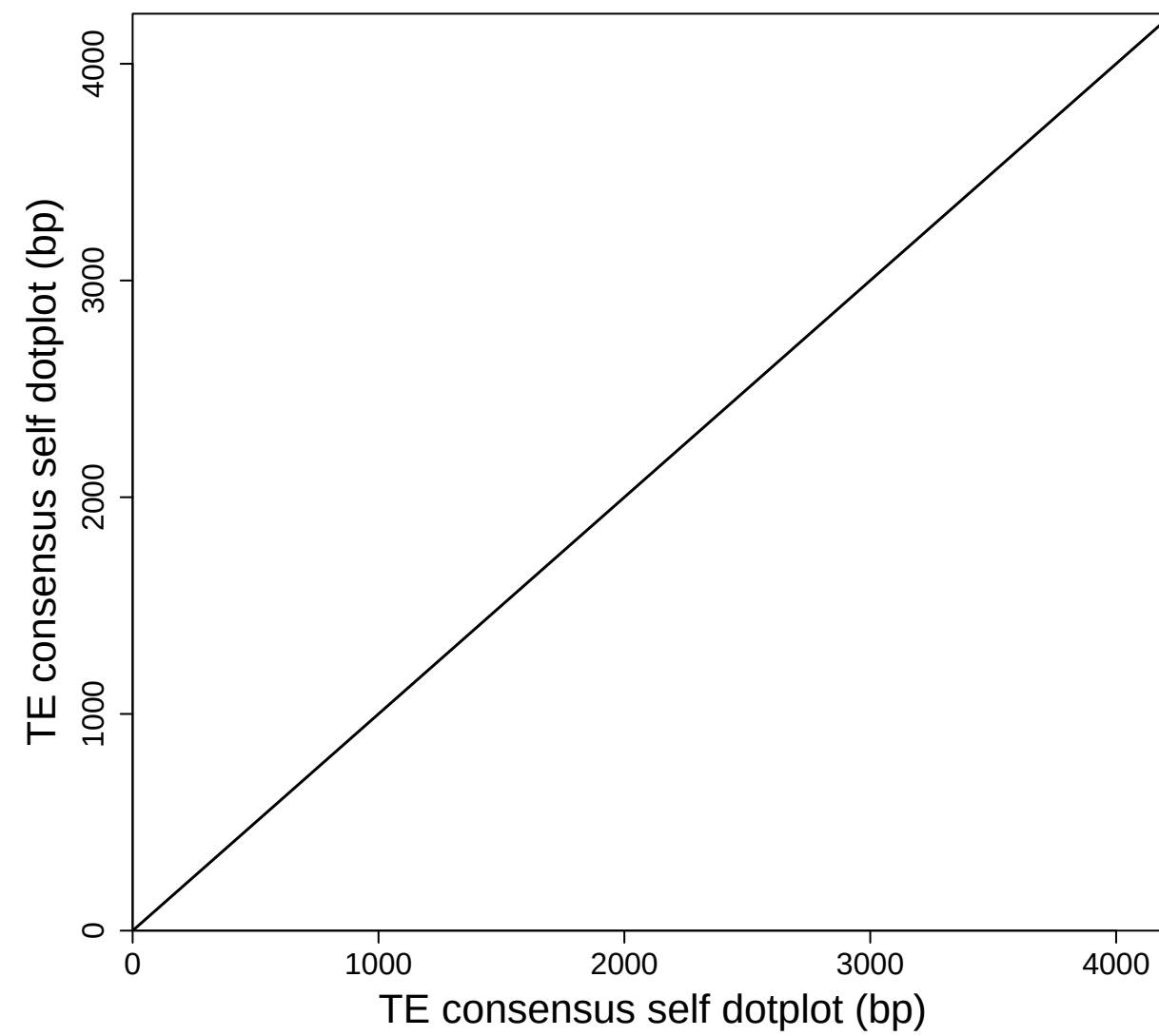
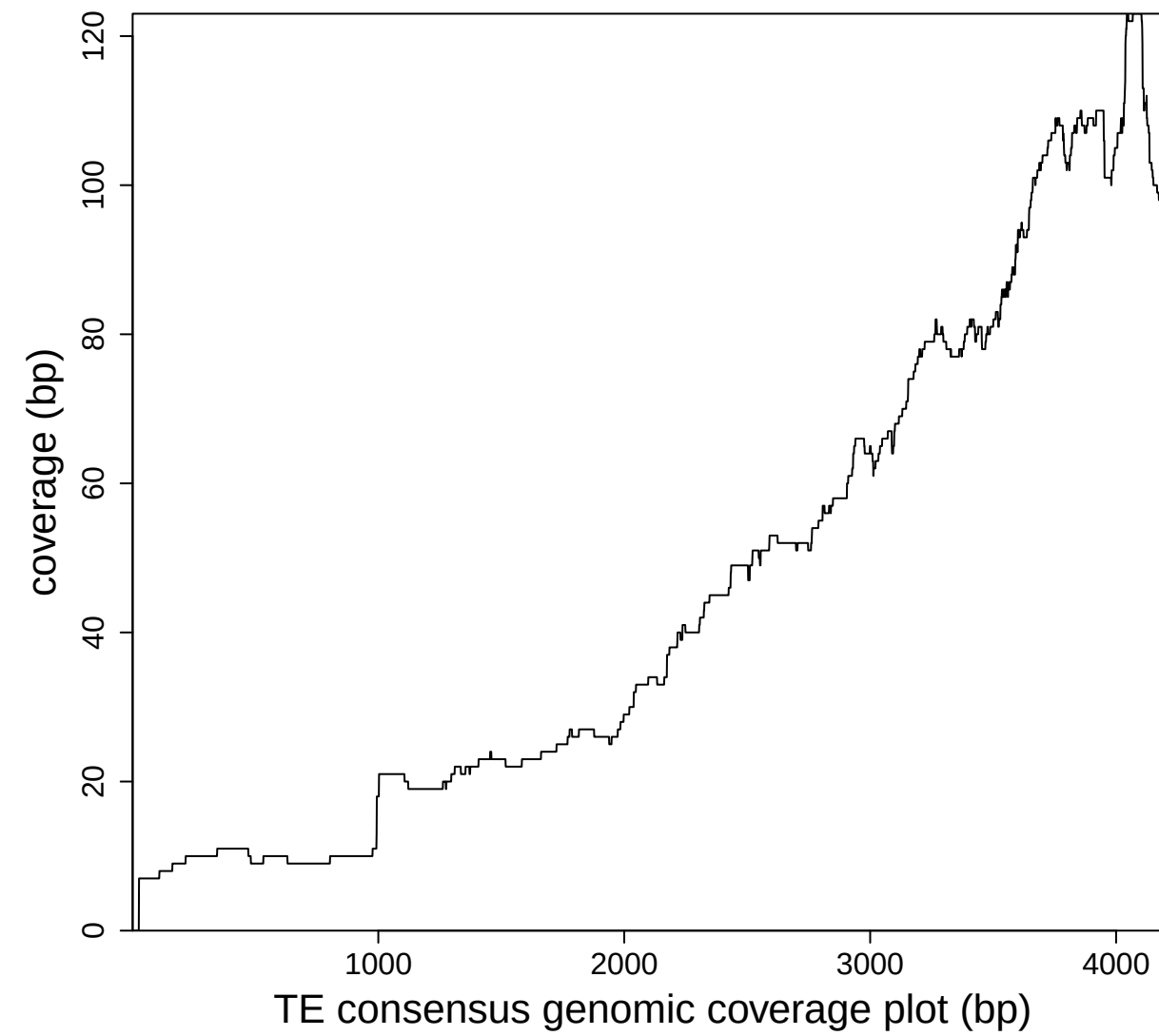
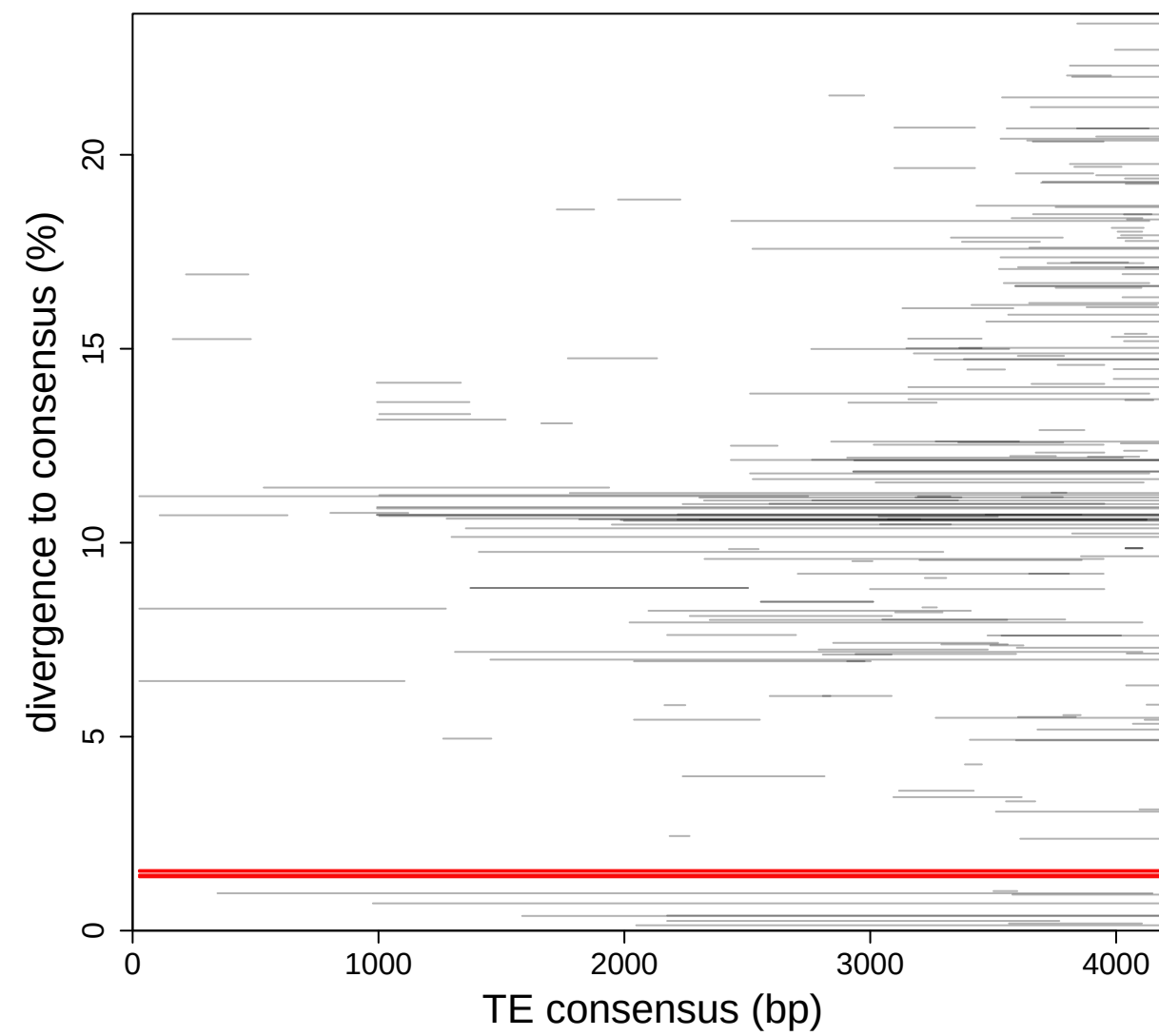
3000

4000



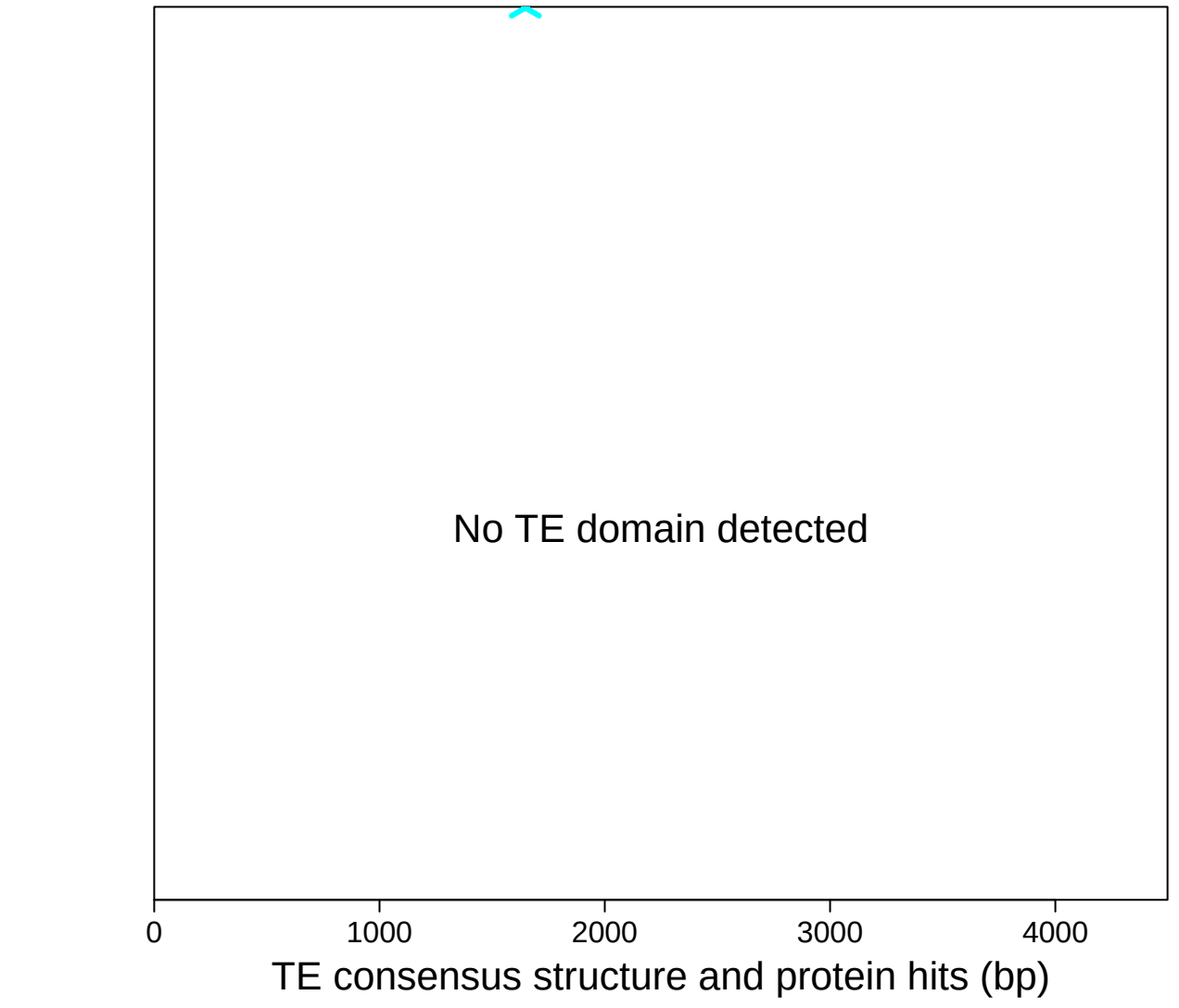
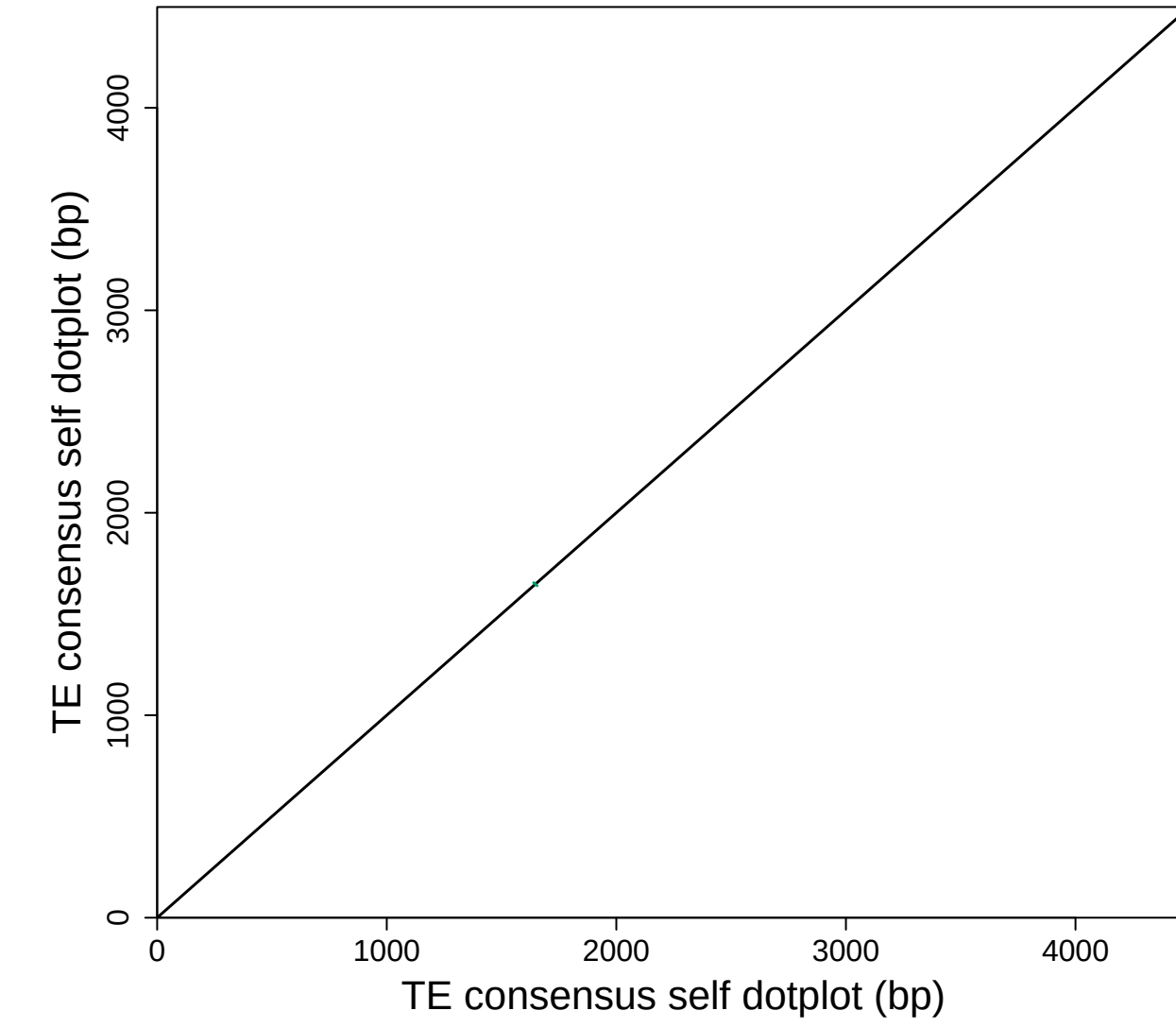
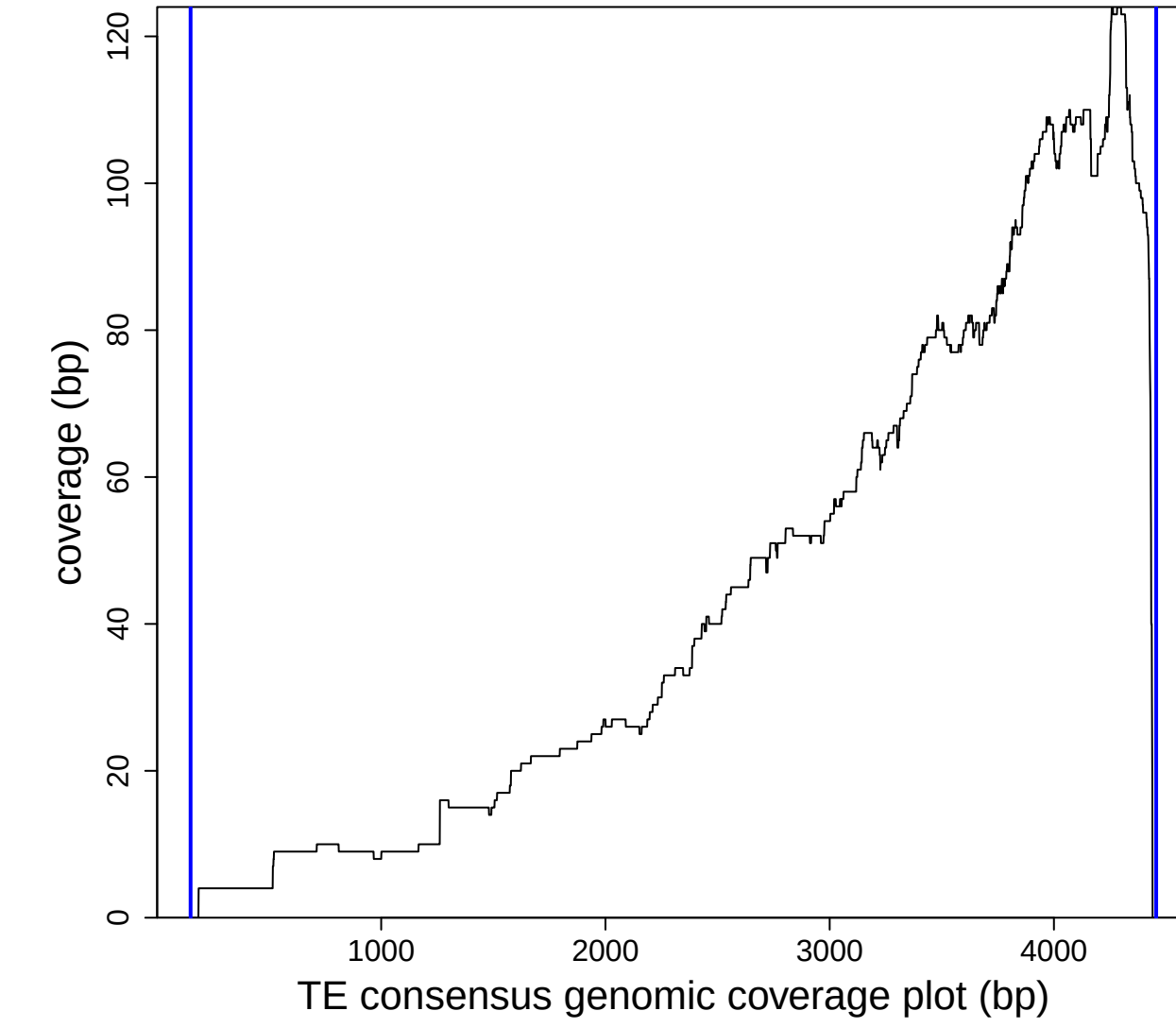
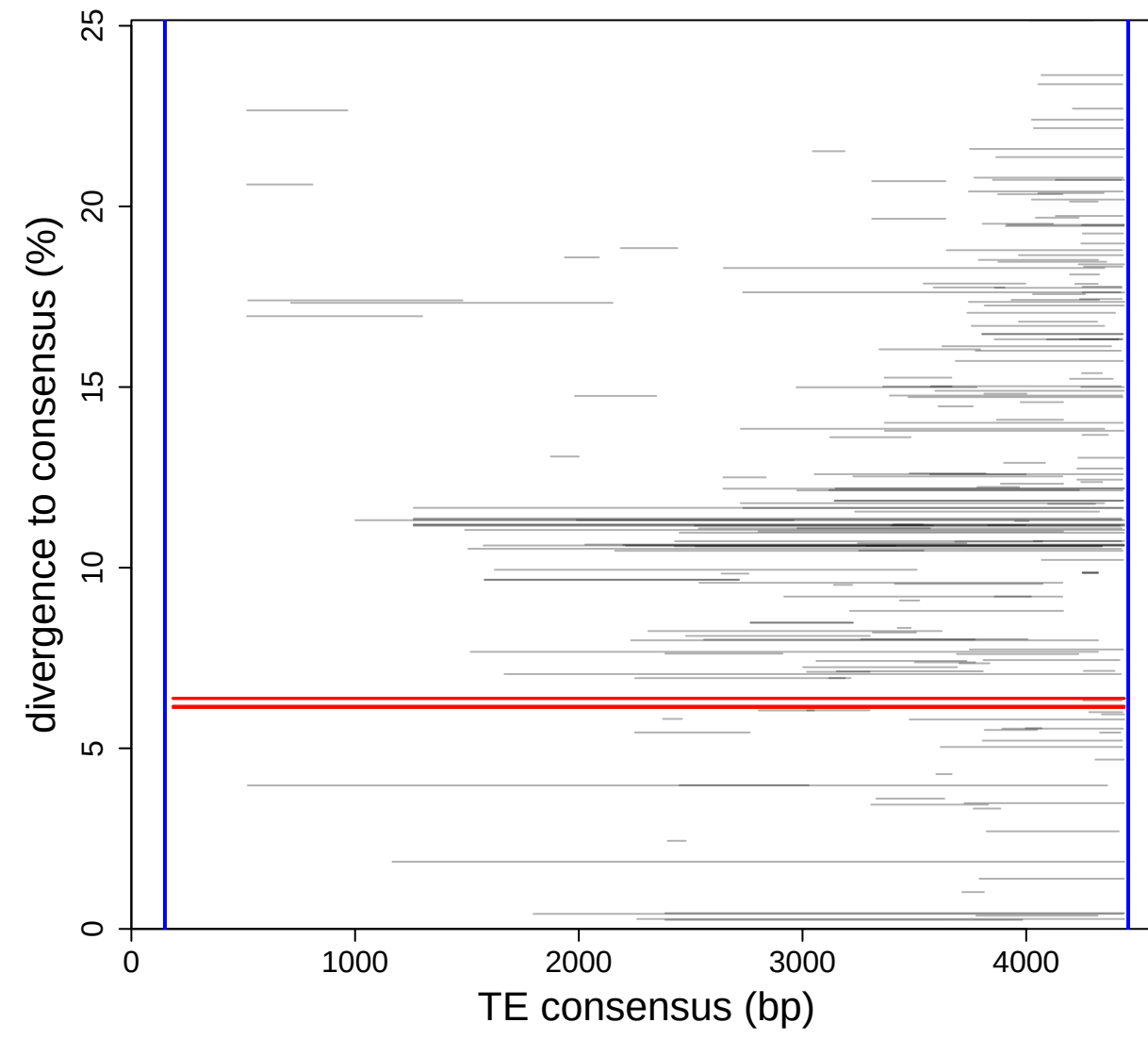
bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 4231bp; fragments: 225; full length: 4 (>=3807.9bp)

After TEtrimmer 4231 bp

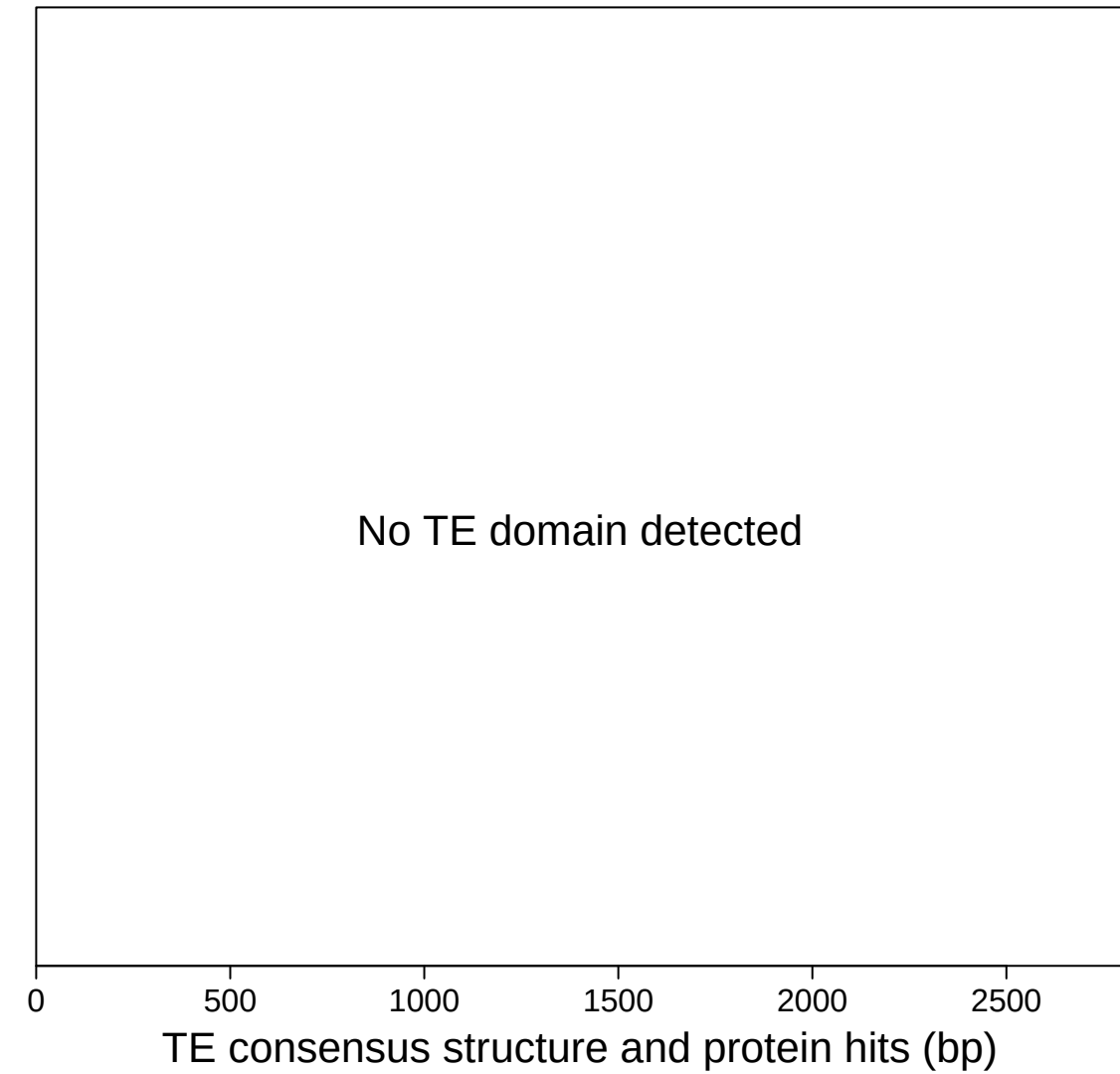
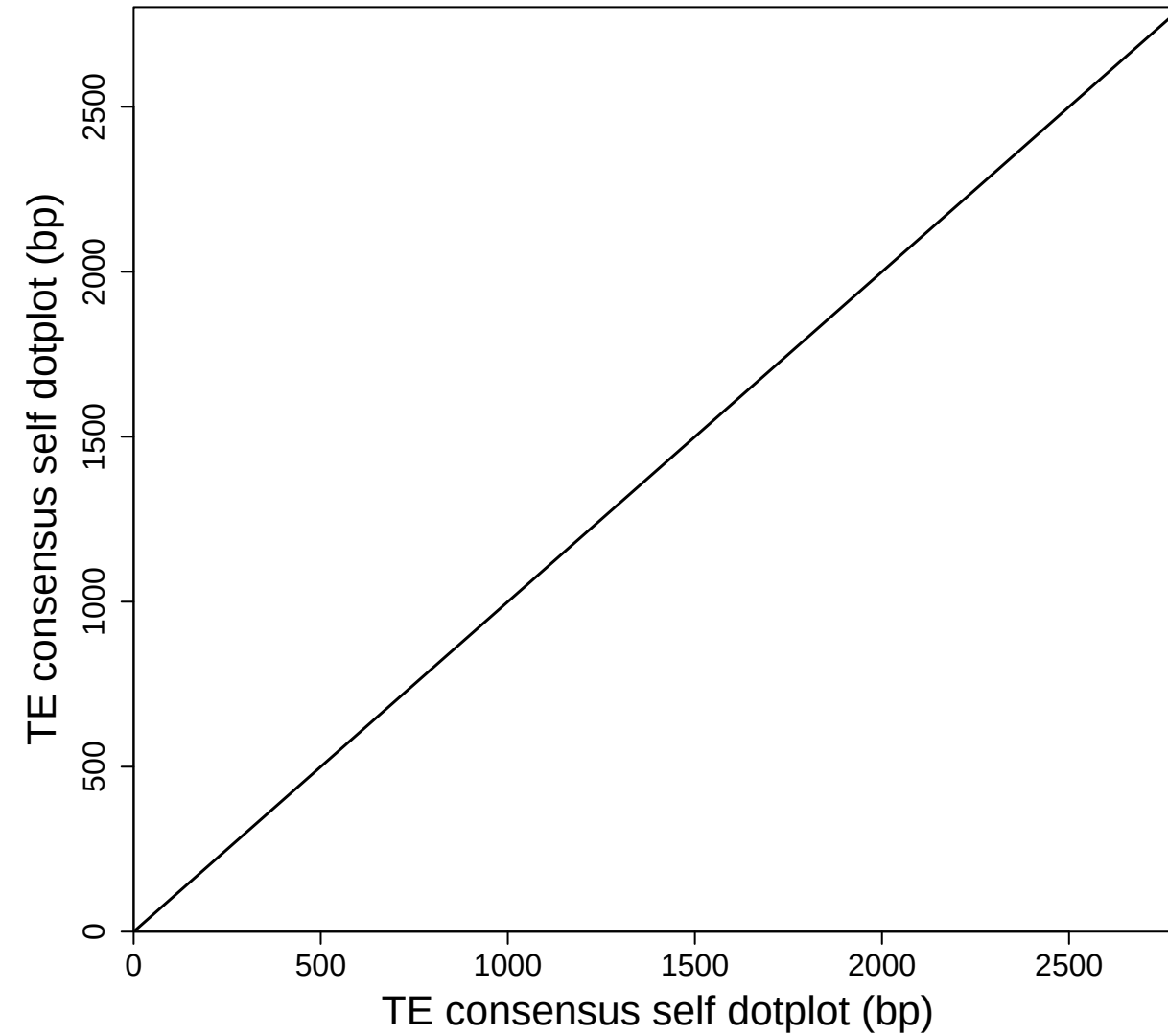
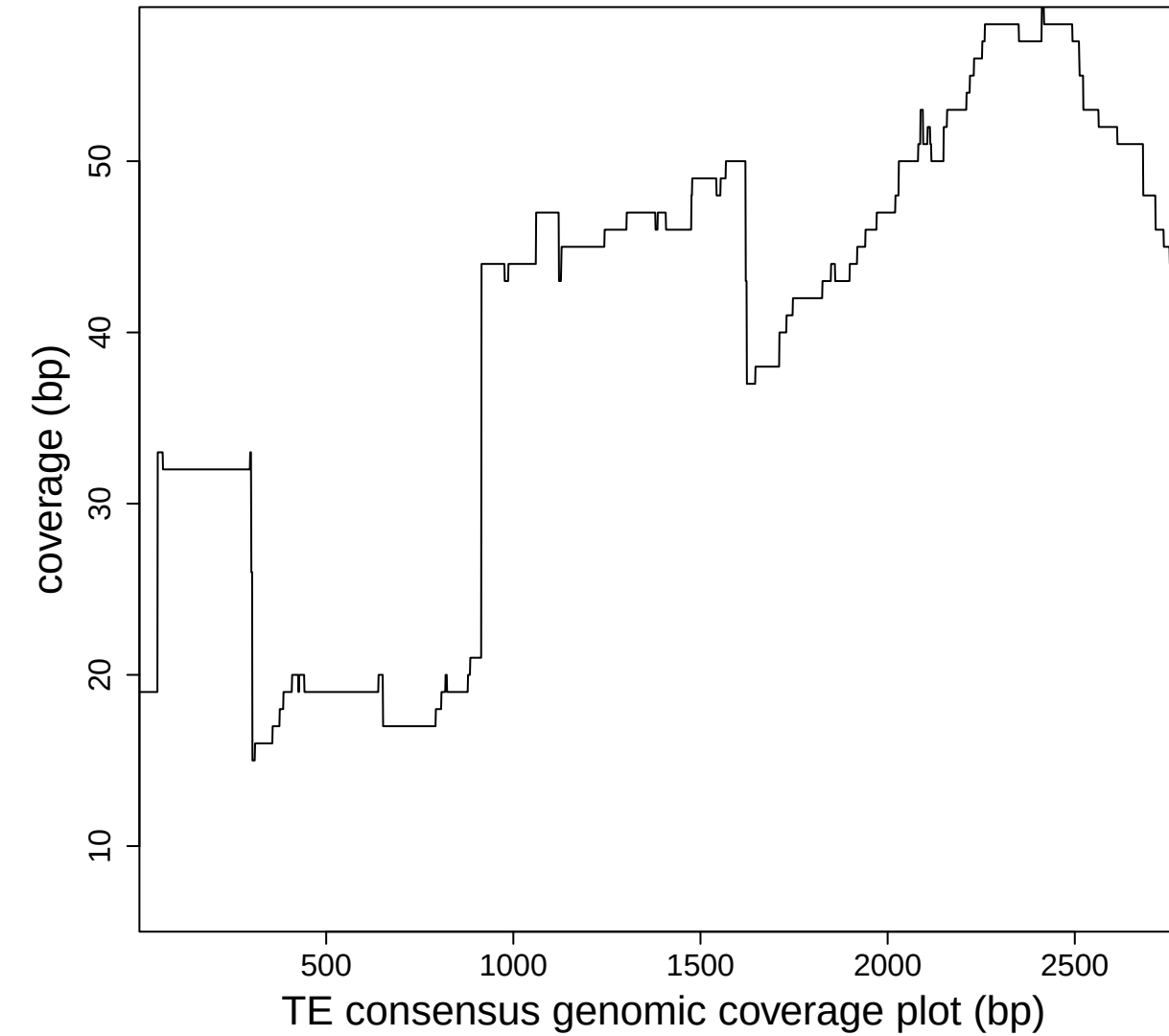
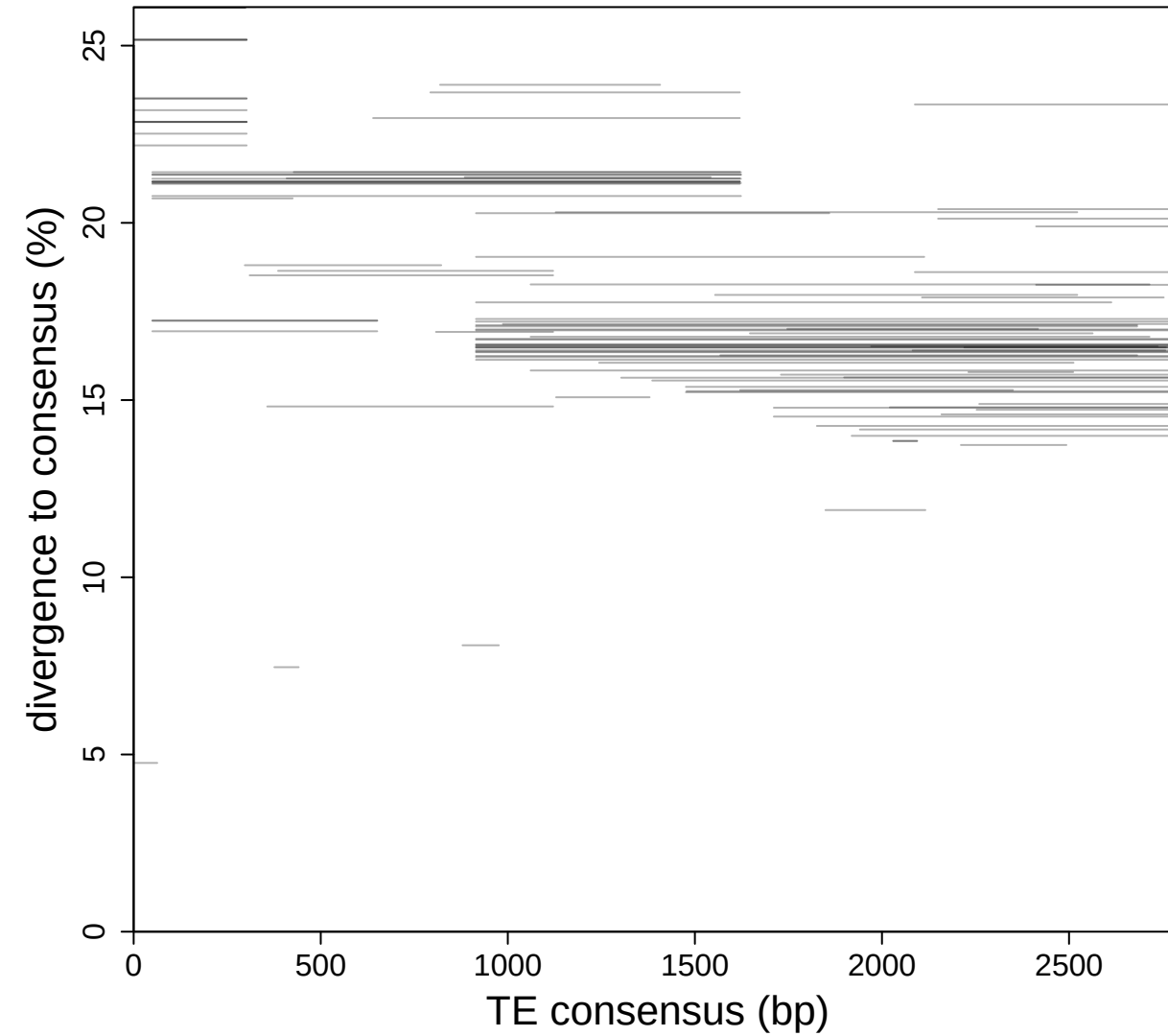


9.fasta.b.bed uf.bed g 4.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 4606bp; fragments: 218; full length: 4 (>=4145.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

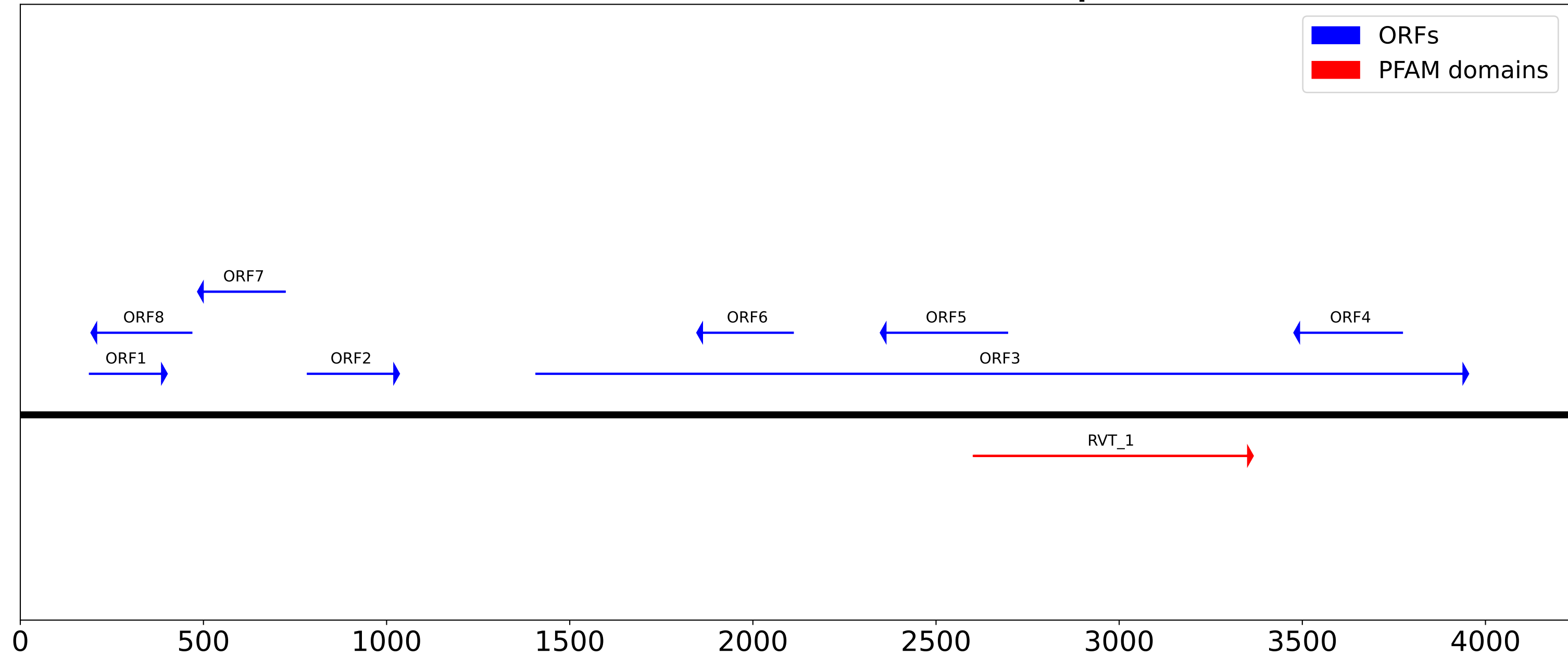


TE: rnd_4_family_309
size: 2802bp; fragments: 112; full length: 0 (≥ 2521.8 bp)

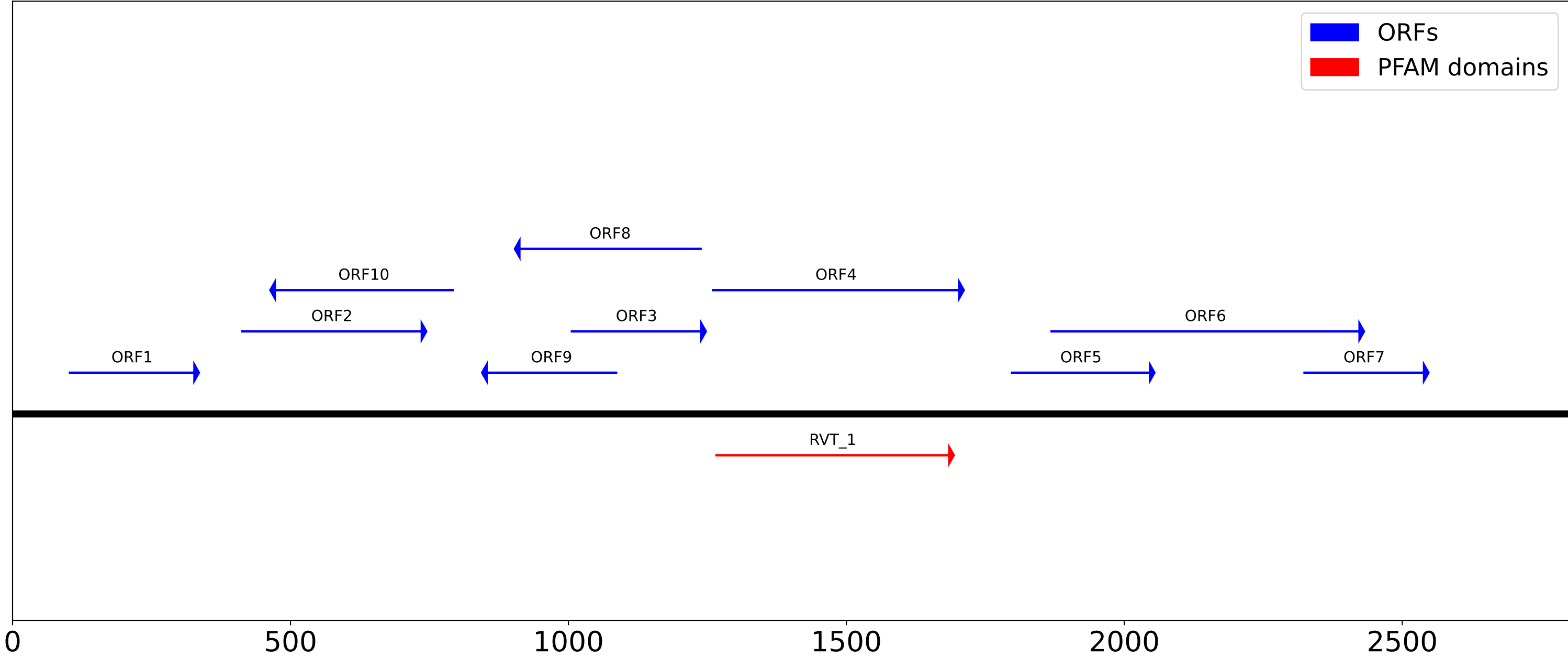


Before TEtrimmer 2802 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

