

Start crop Point



End crop Point

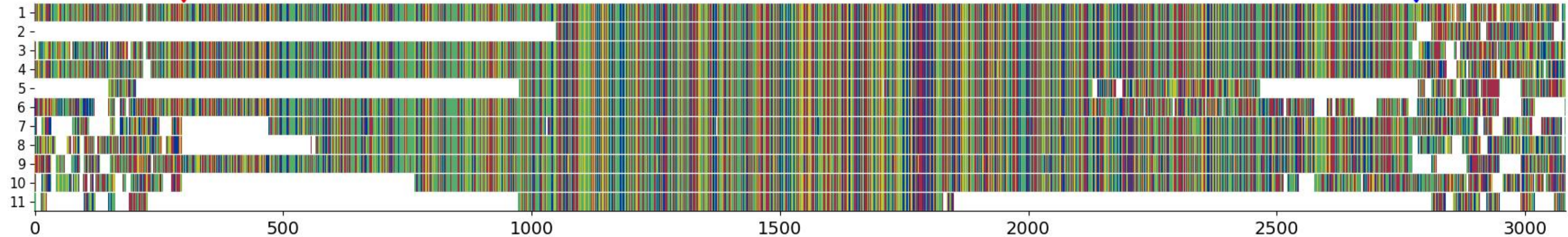


MSA length = 2482



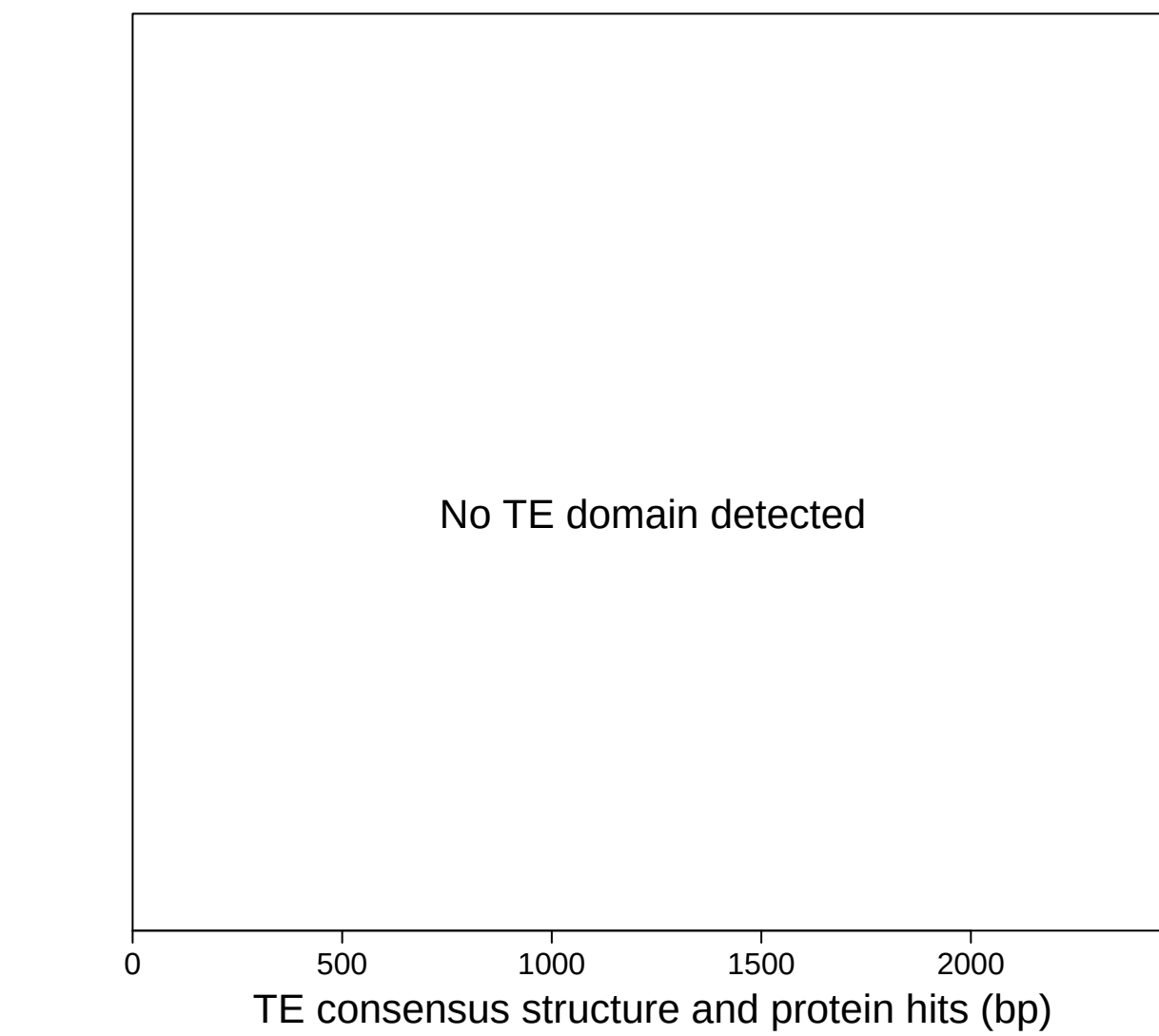
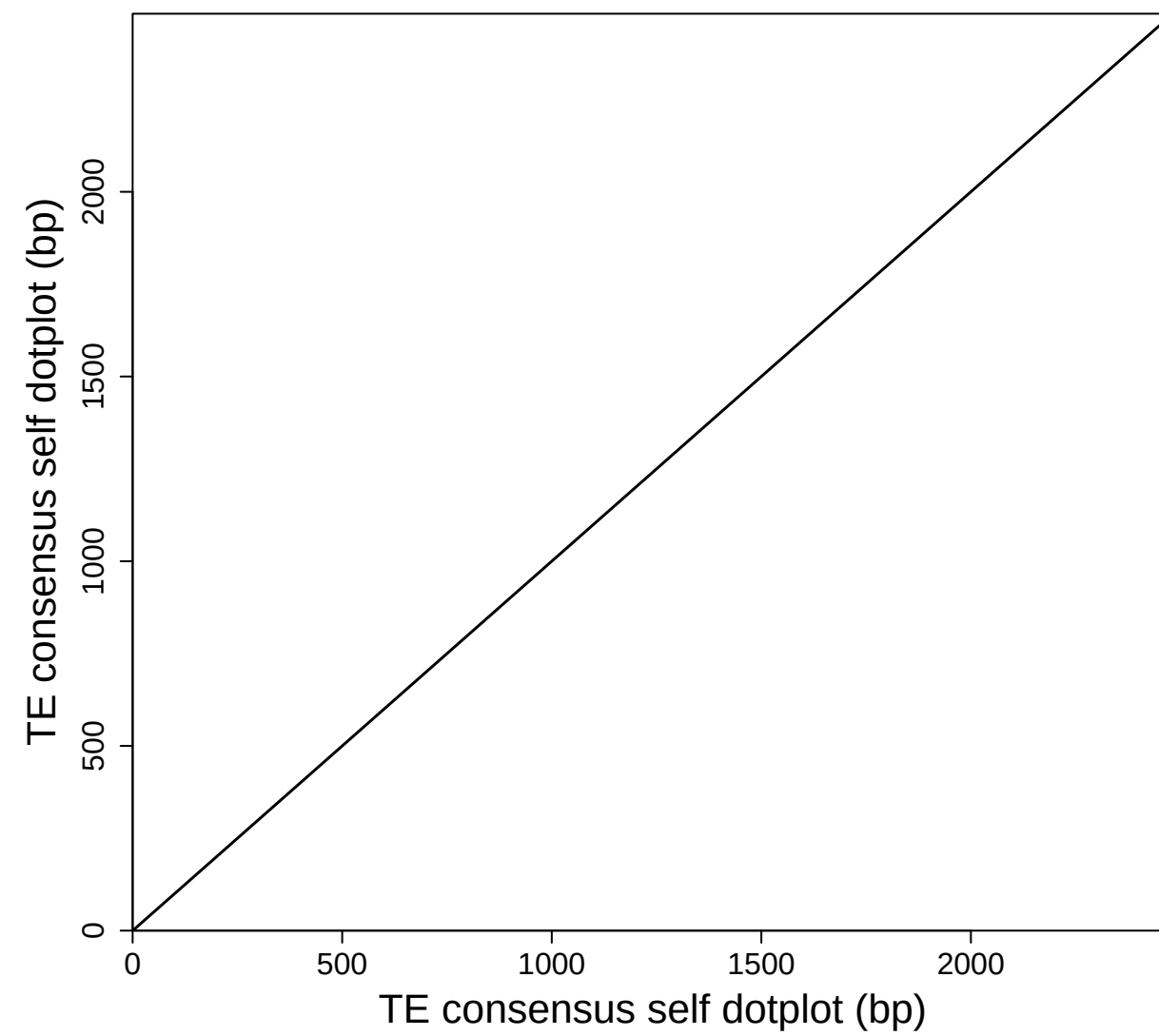
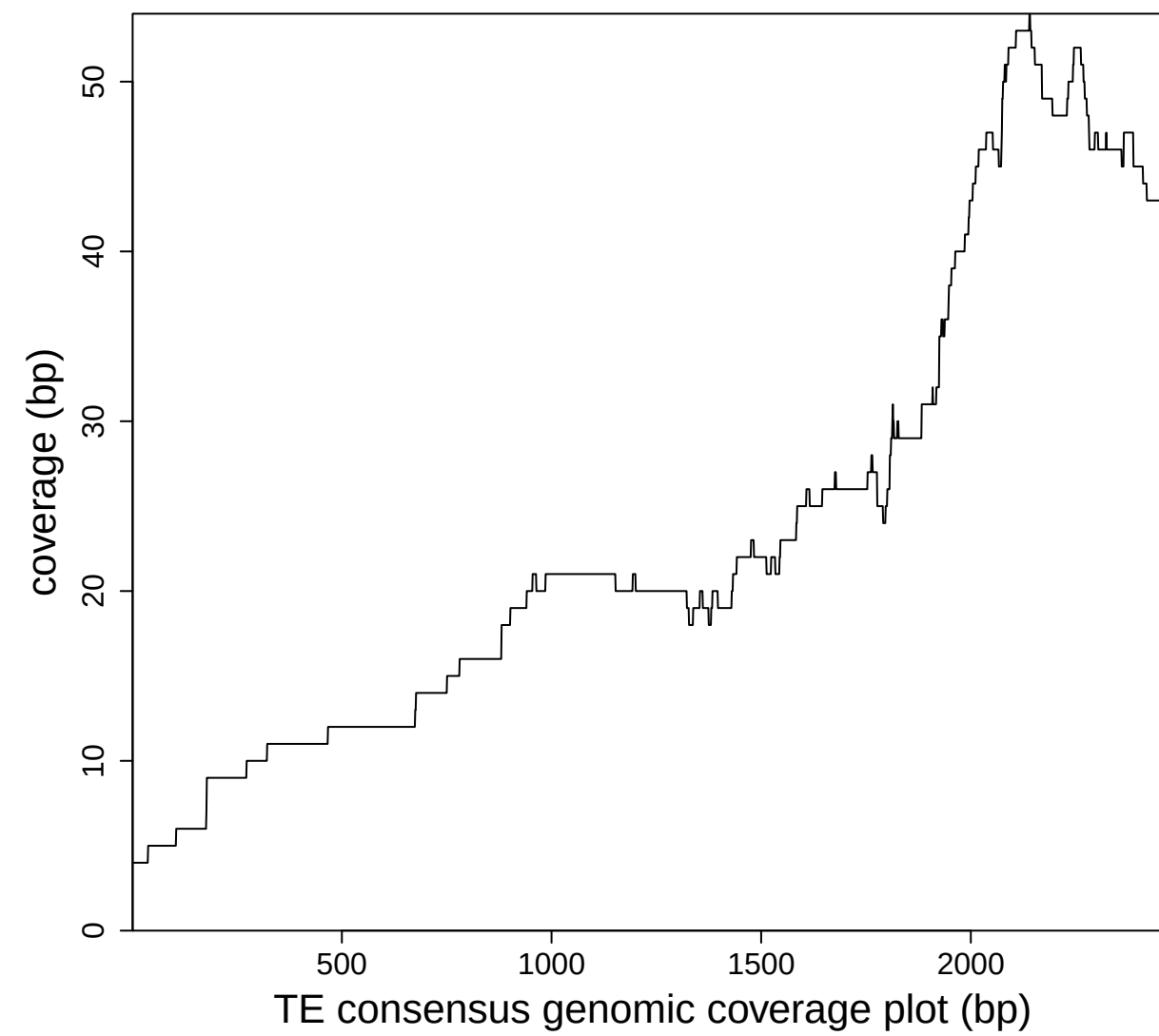
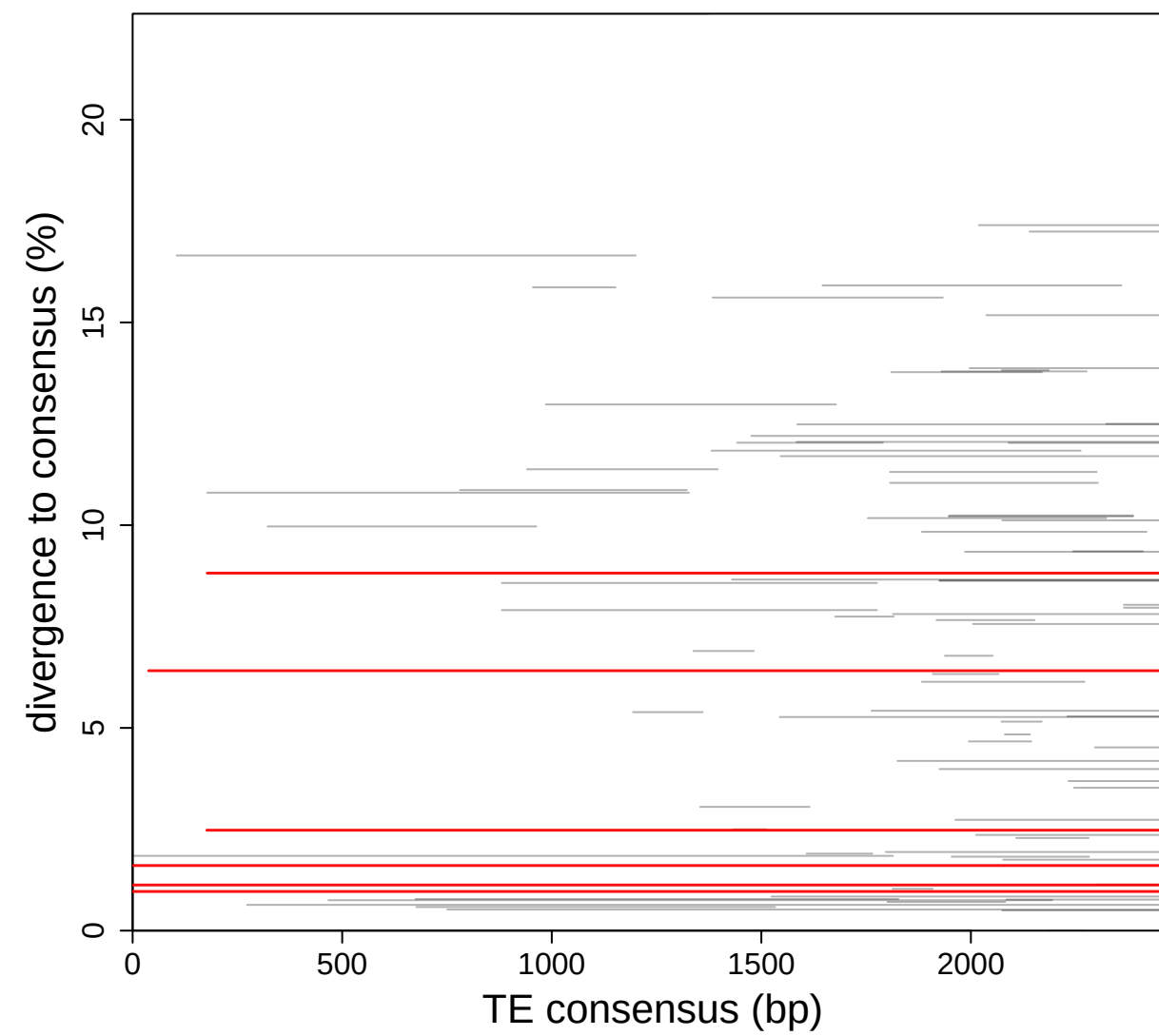
Start crop Point

End crop Point



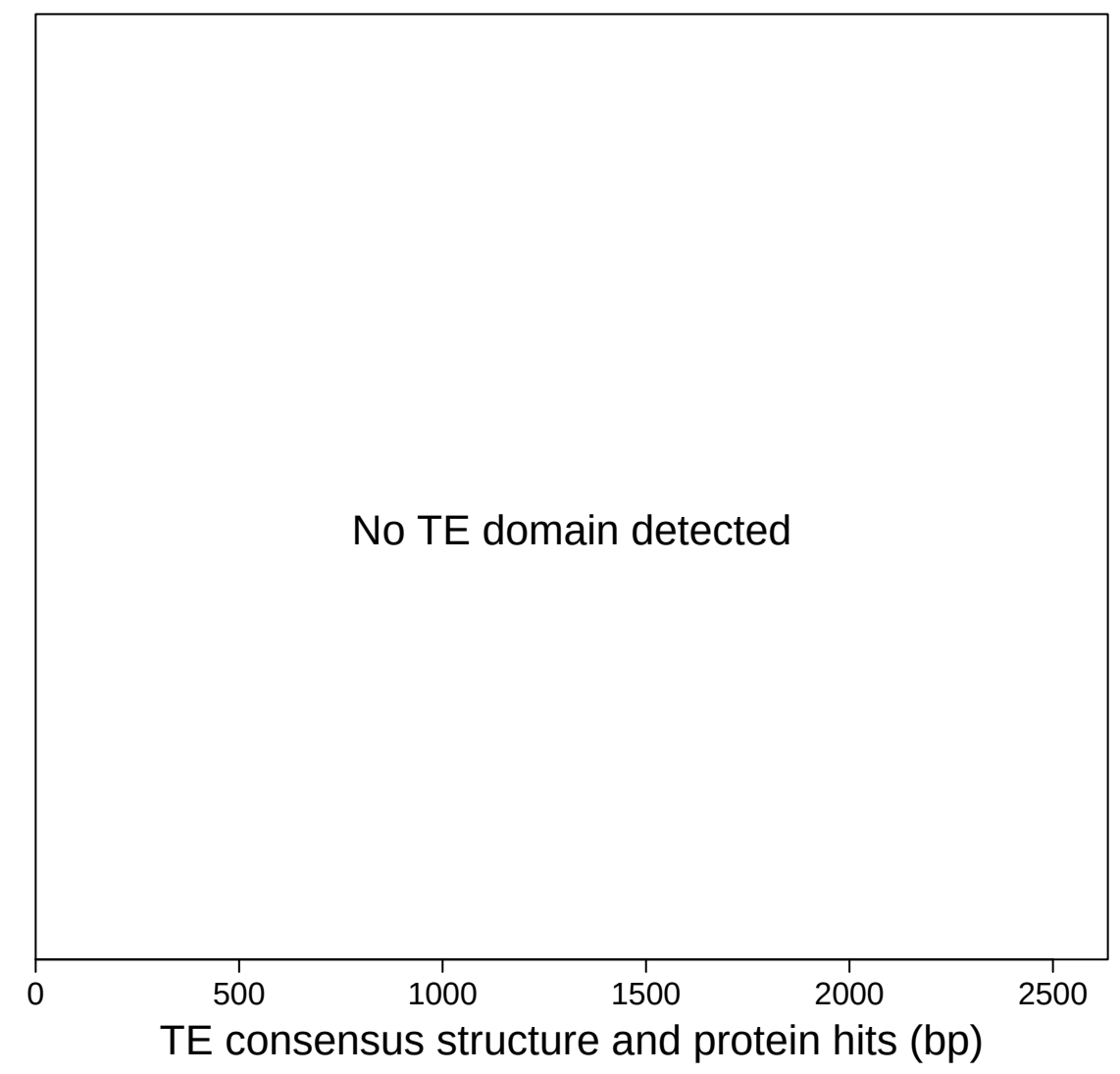
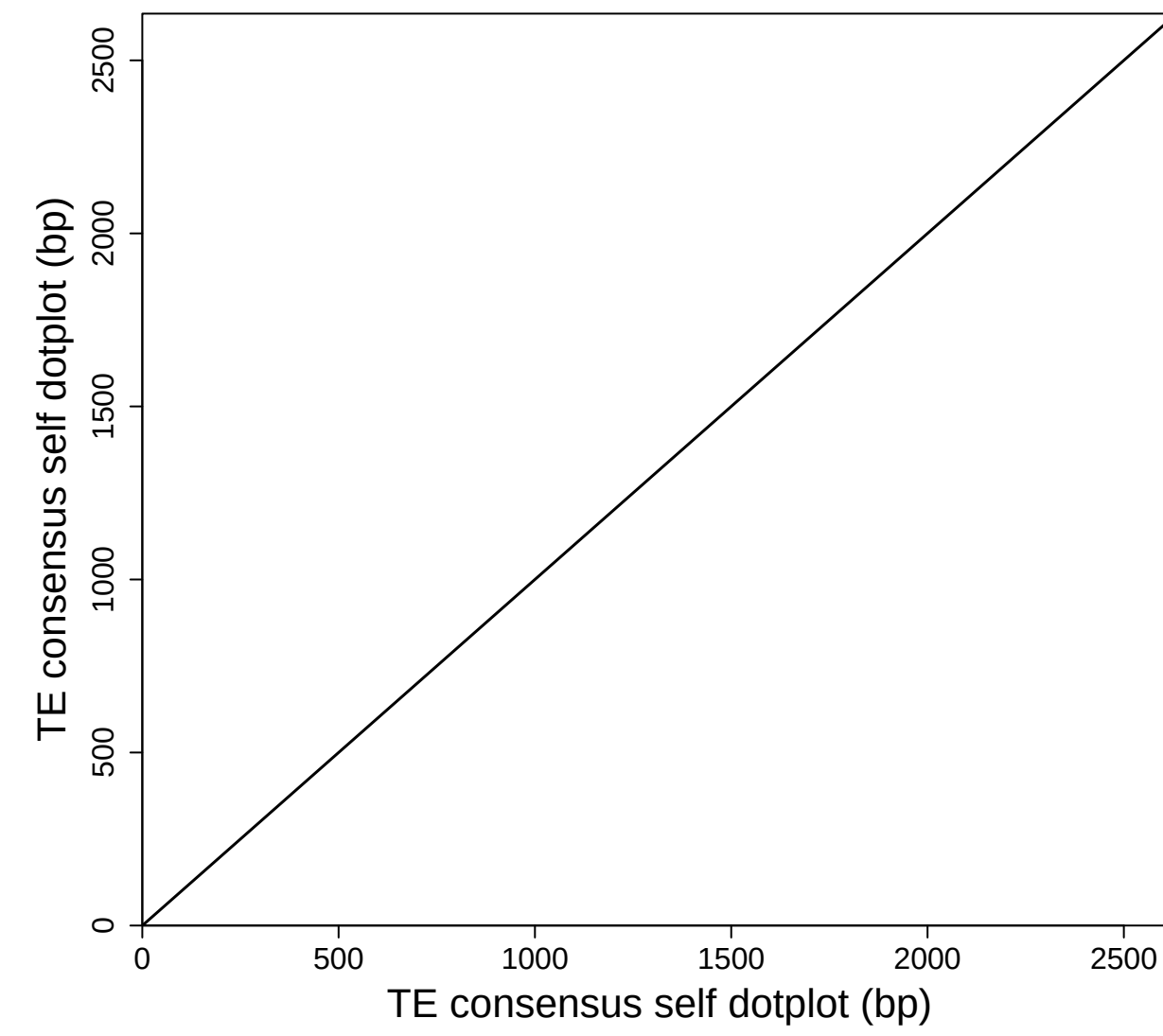
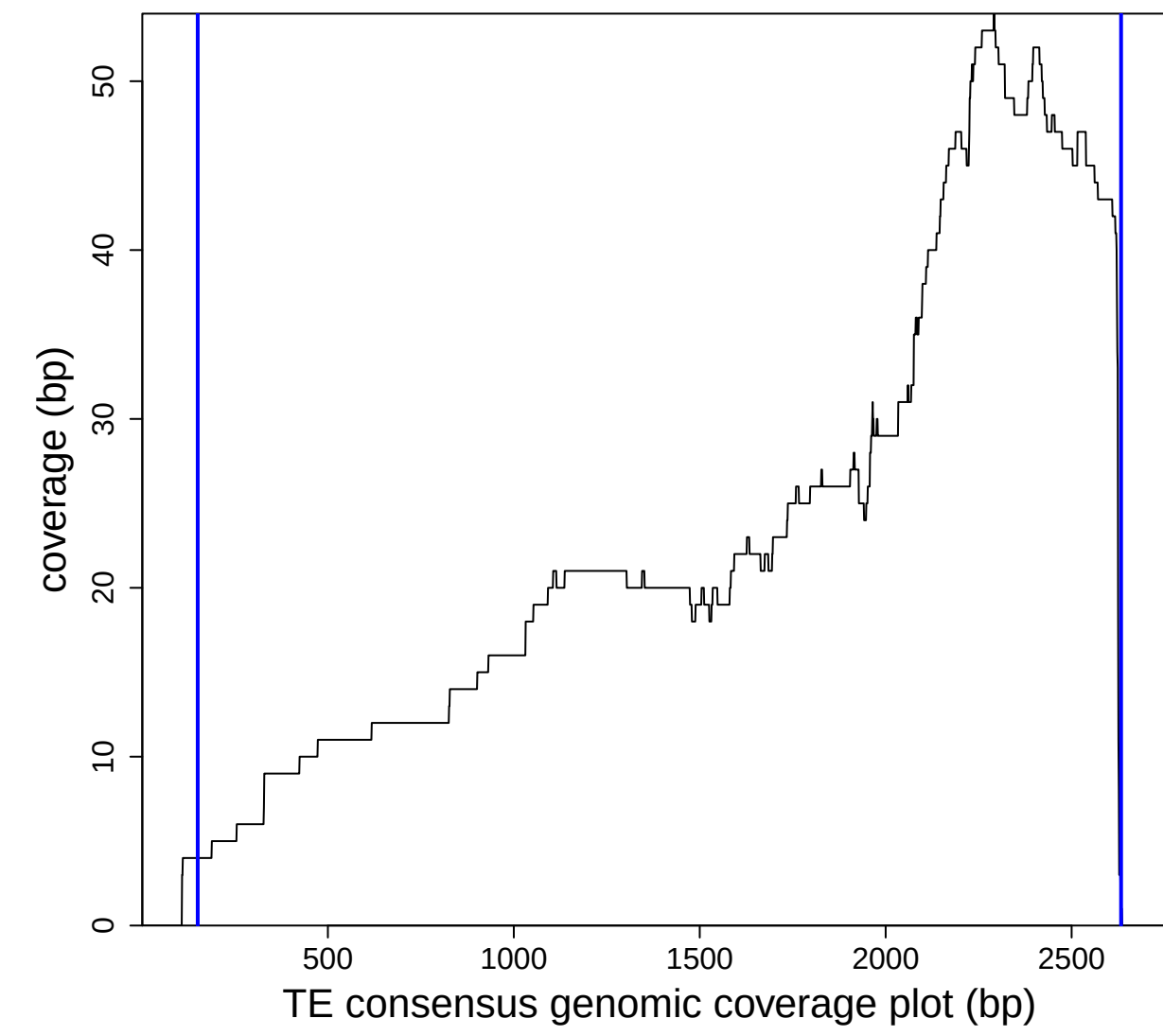
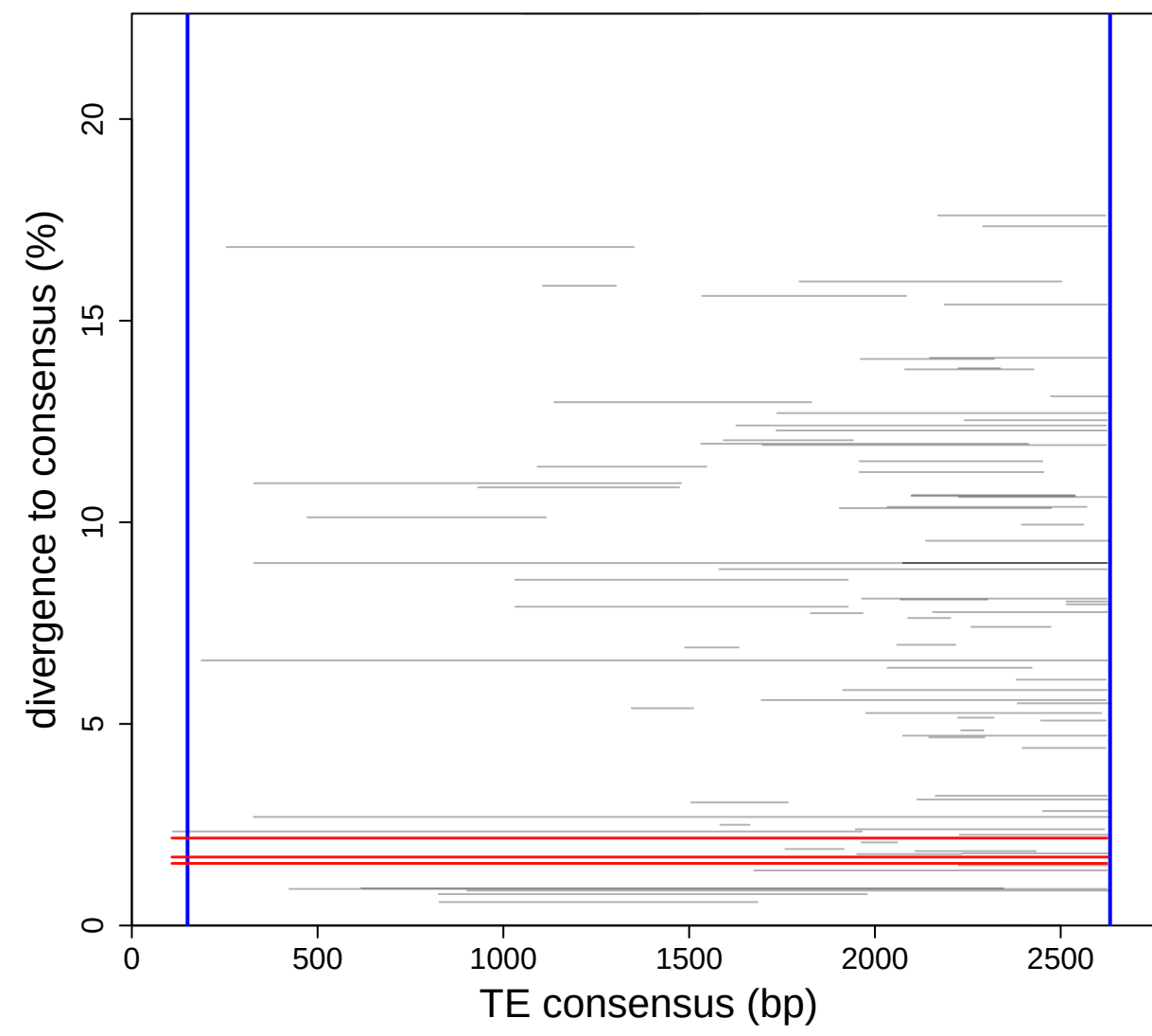
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 2482bp; fragments: 89; full length: 6 (≥ 2233.8 bp)

After TETrimmer 2482 bp

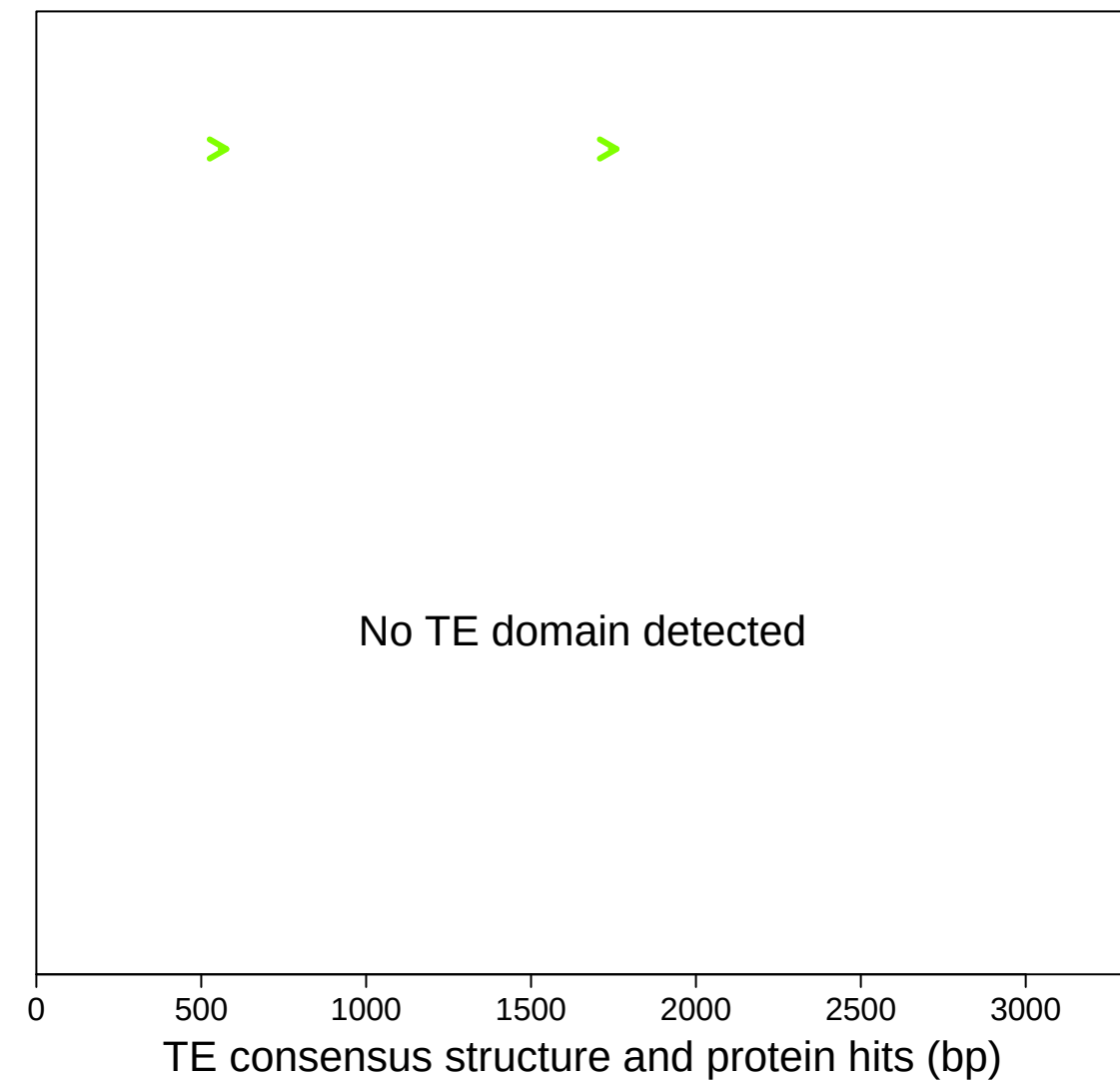
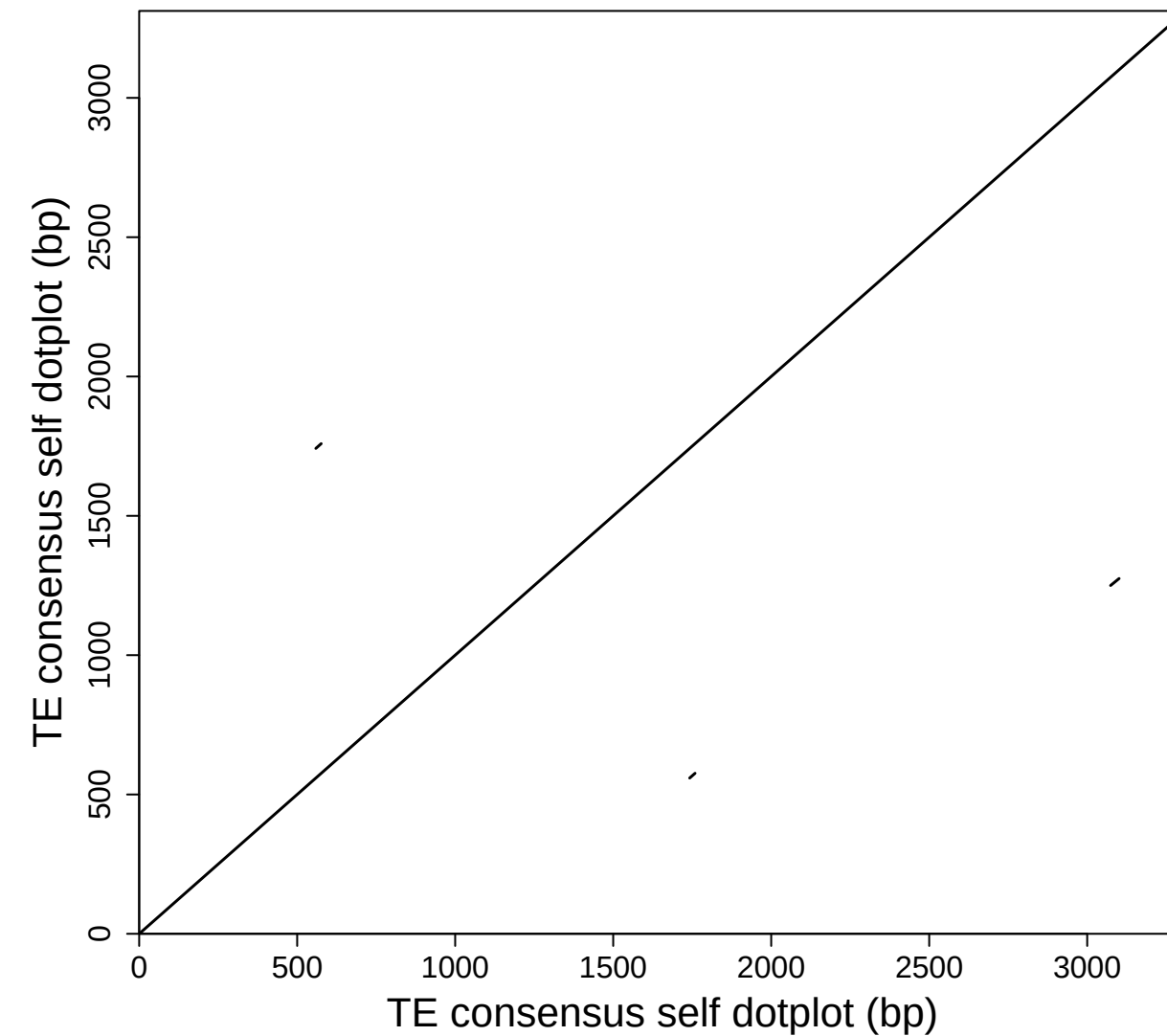
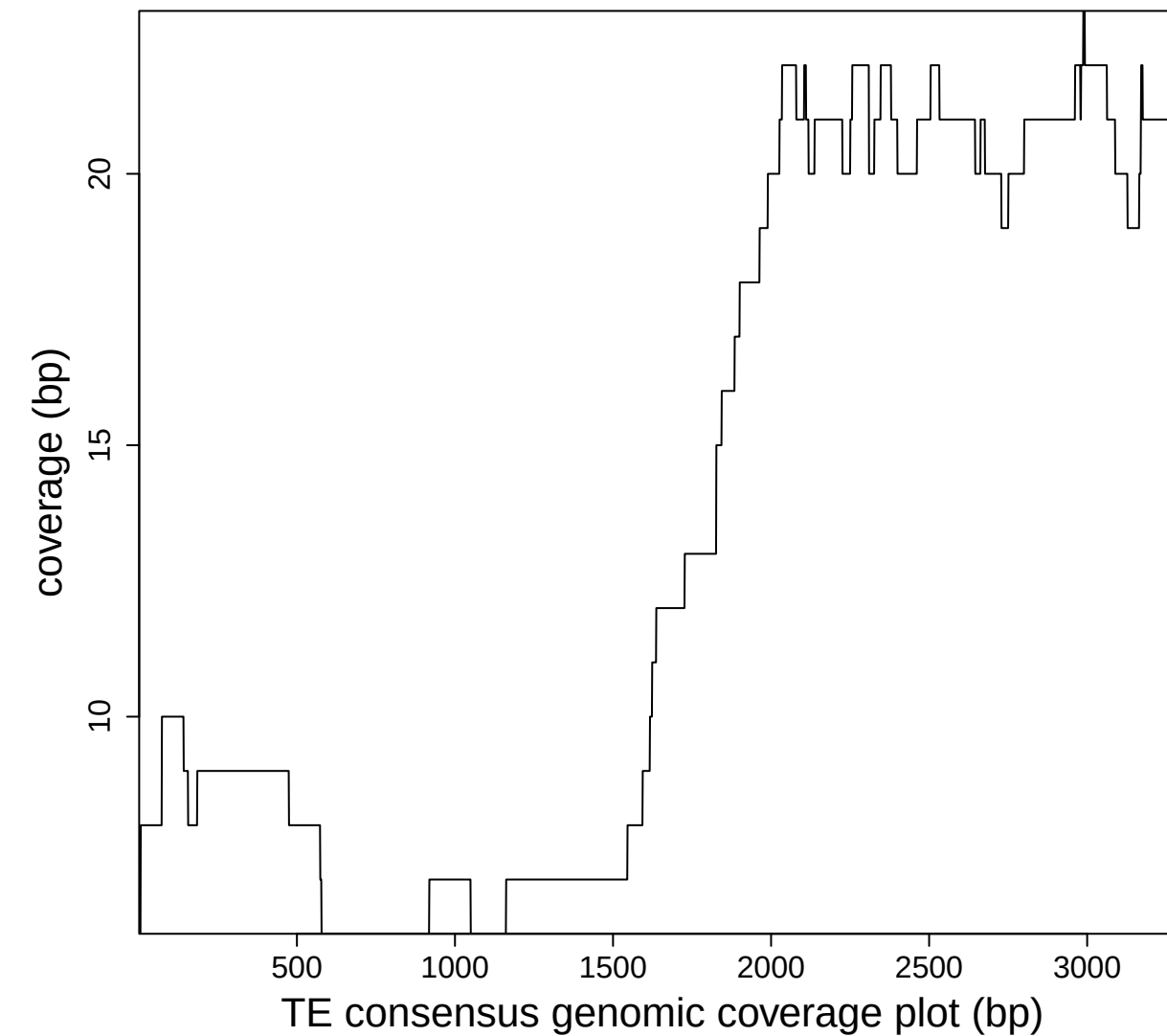
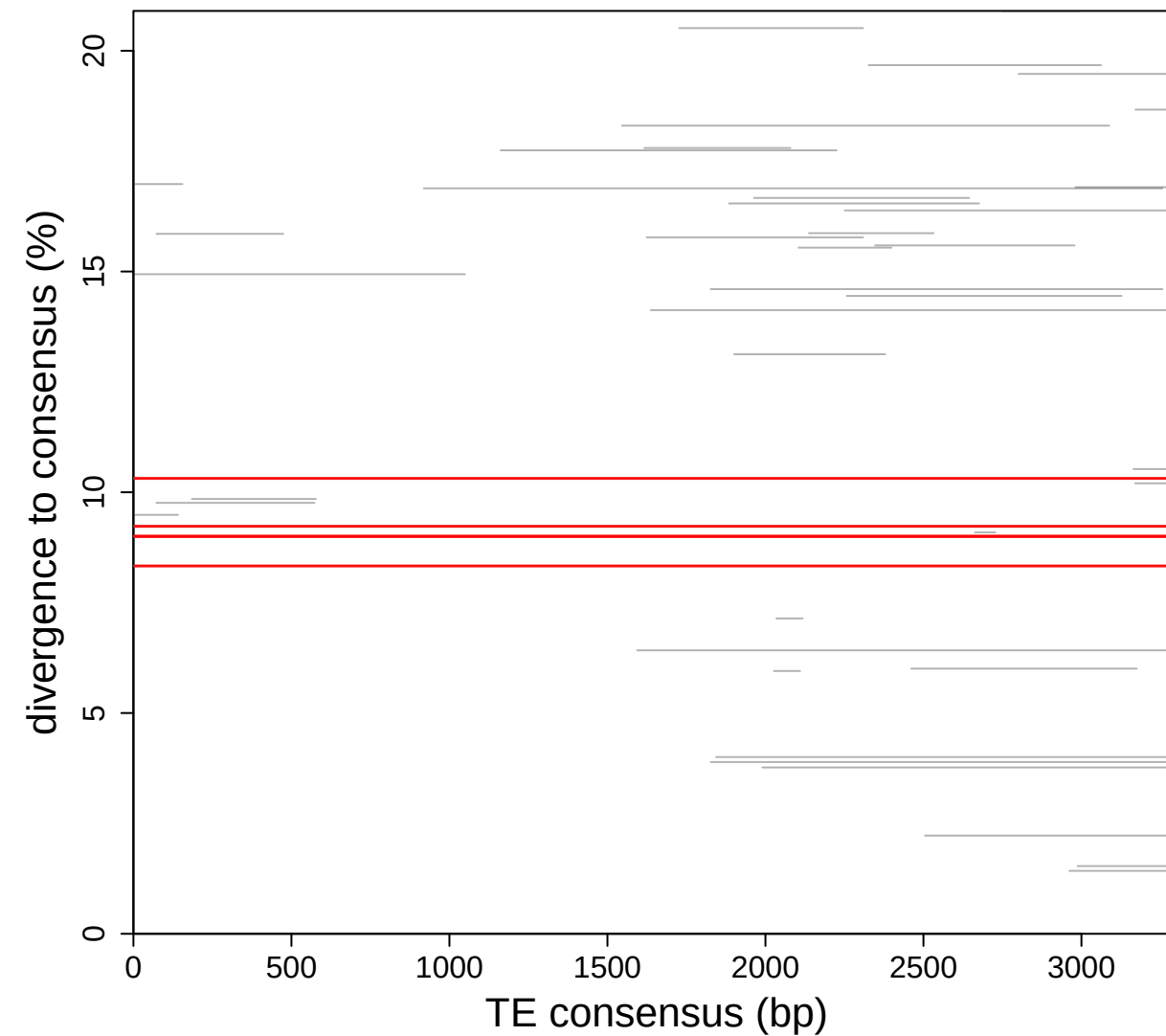


5.fasta.b.bed uf.bed g_1.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 2783bp; fragments: 89; full length: 3 (>=2504.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

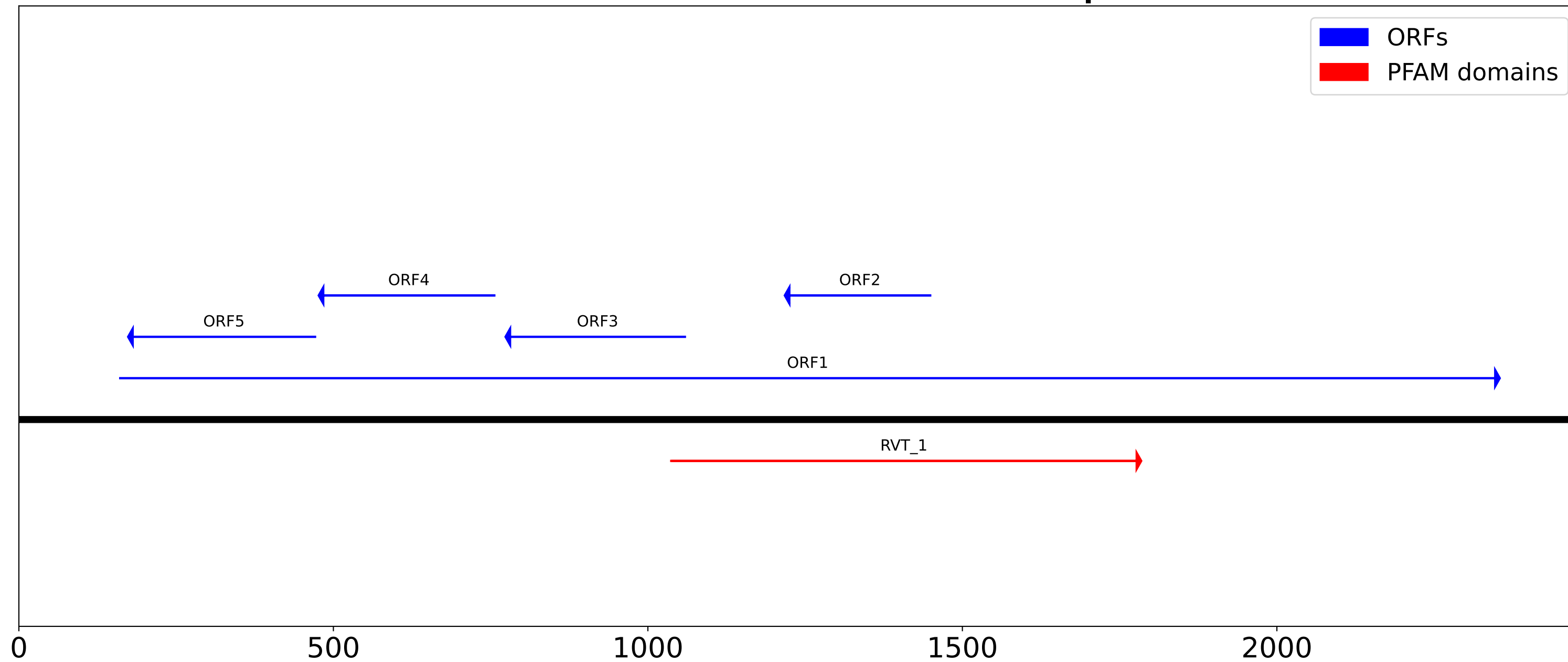


TE: rnd_5_family_2435
size: 3312bp; fragments: 45; full length: 5 (≥ 2980.8 bp)

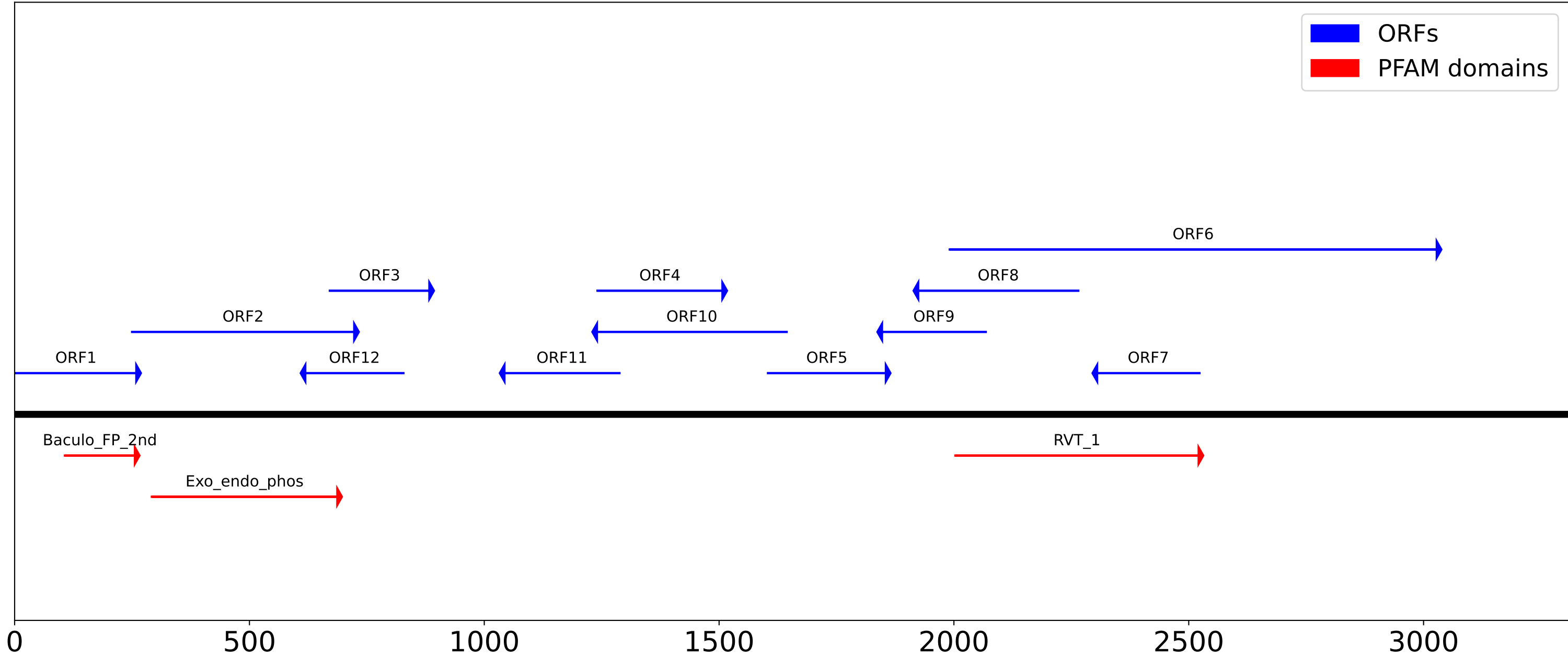


Before TEtrimmer 3312 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

