

MSA length = 4042



Start crop Point

End crop Point

0

10

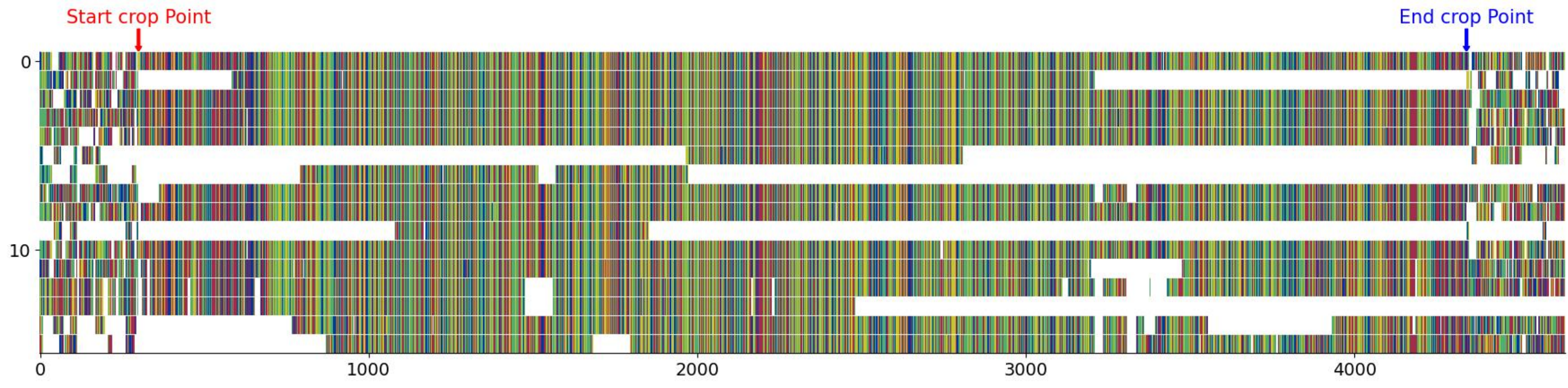
0

1000

2000

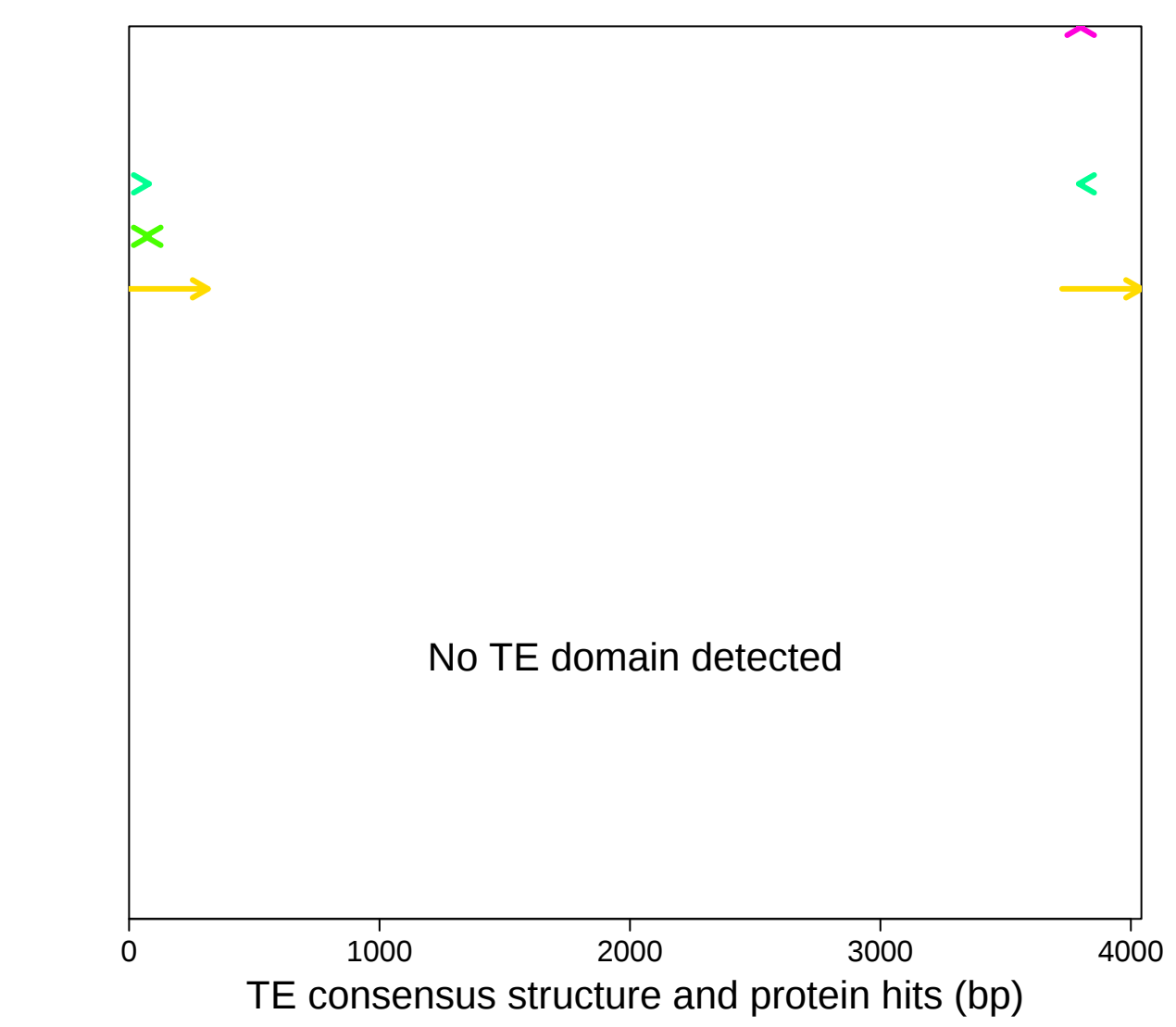
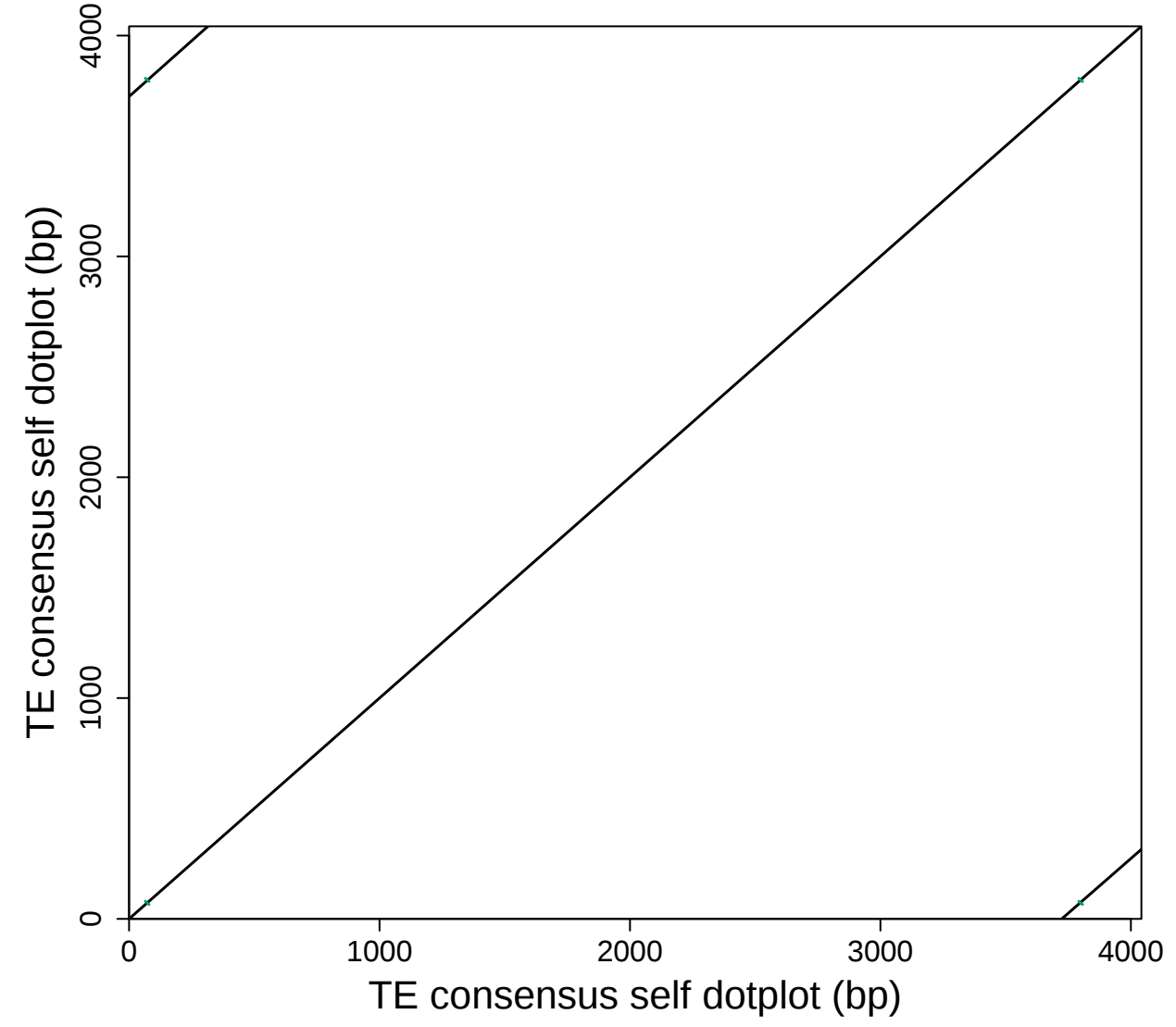
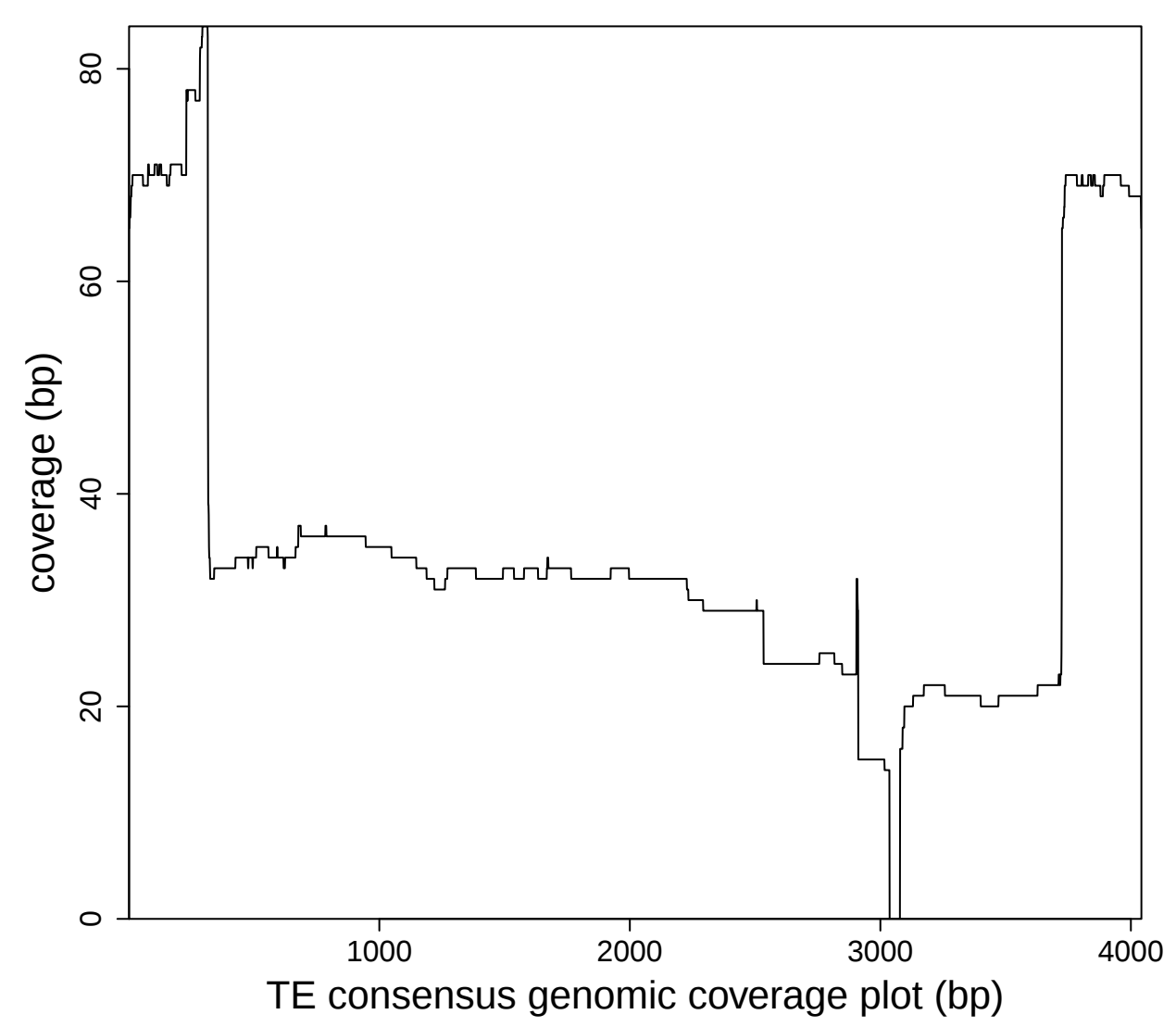
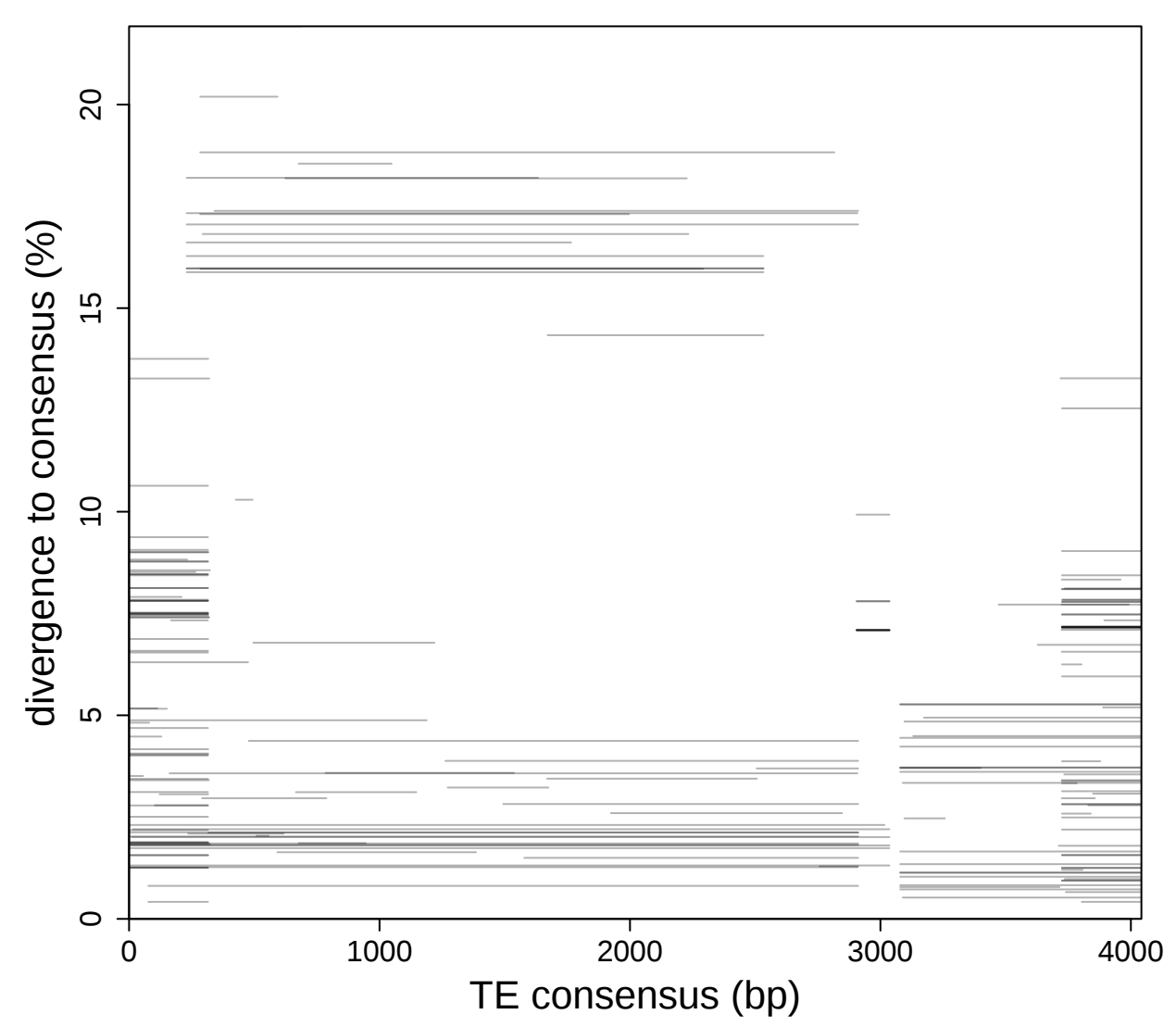
3000

4000



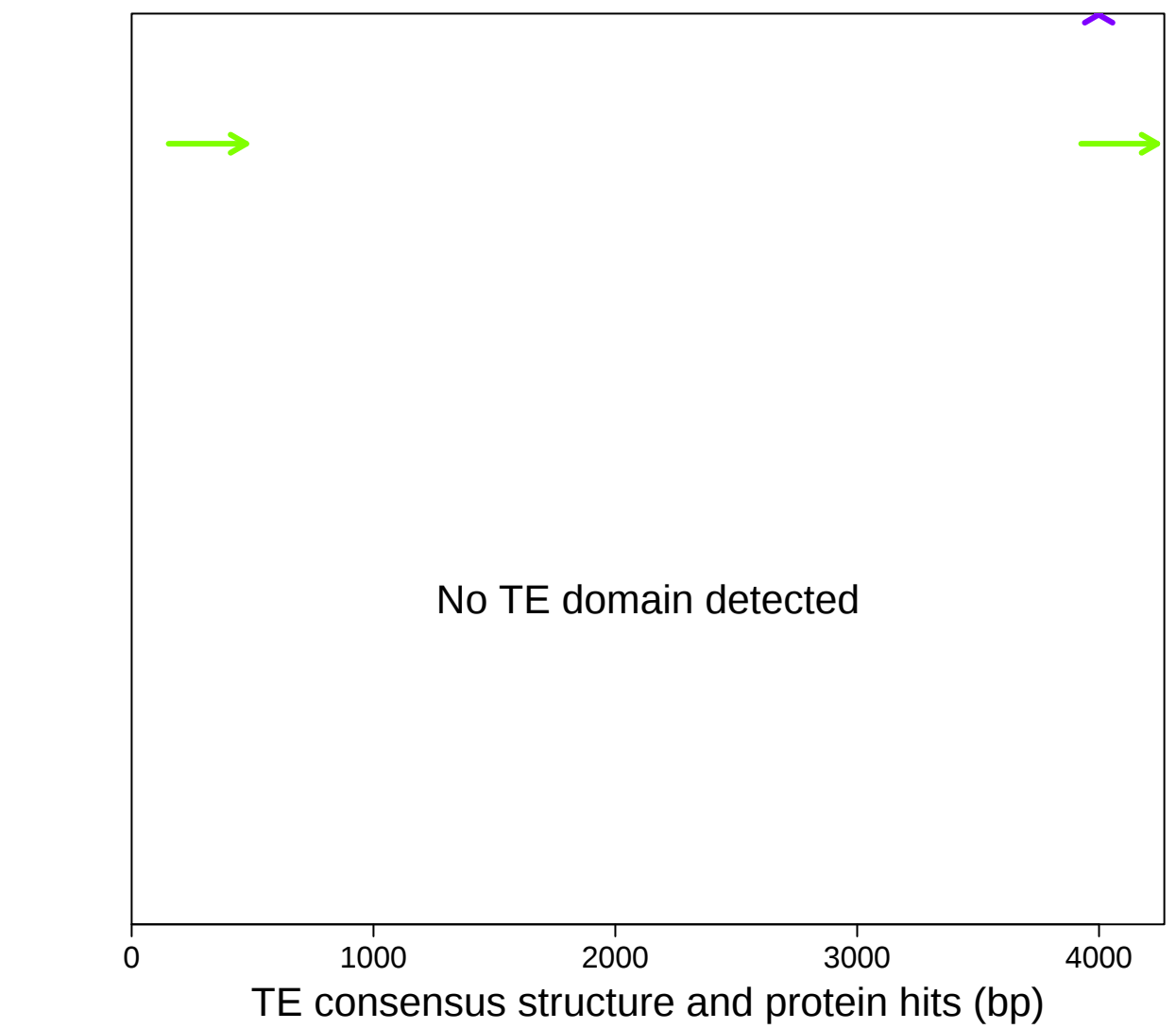
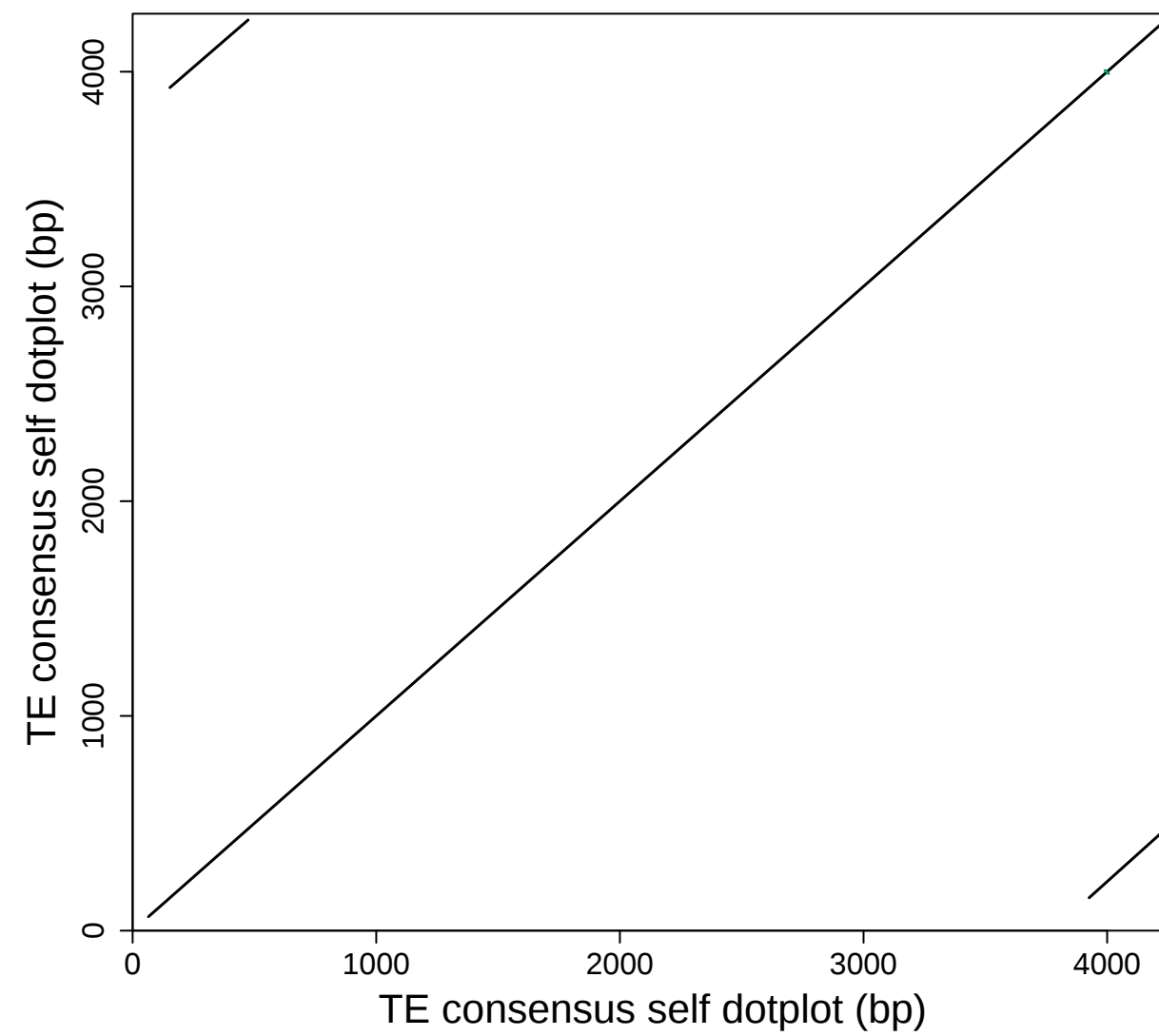
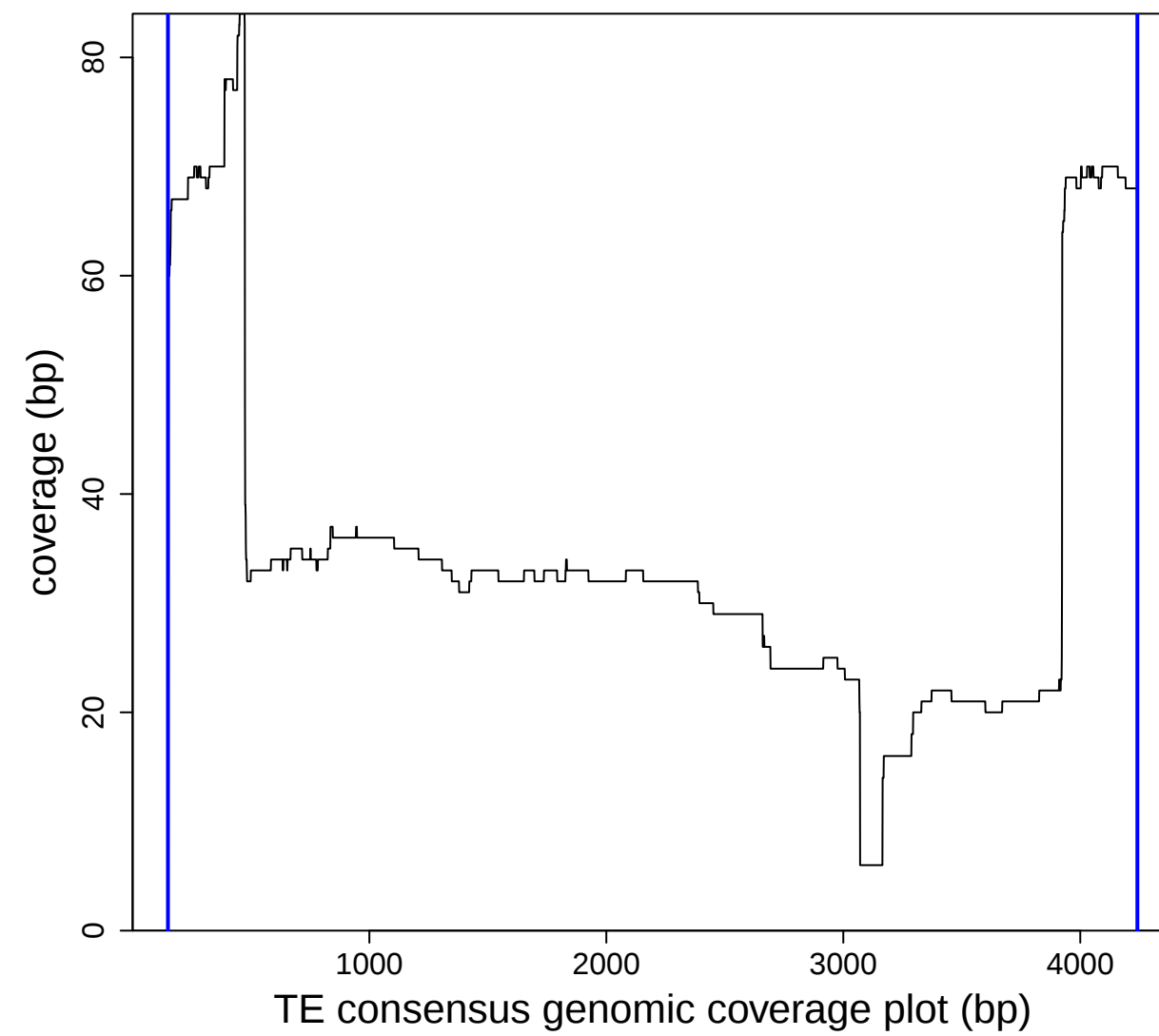
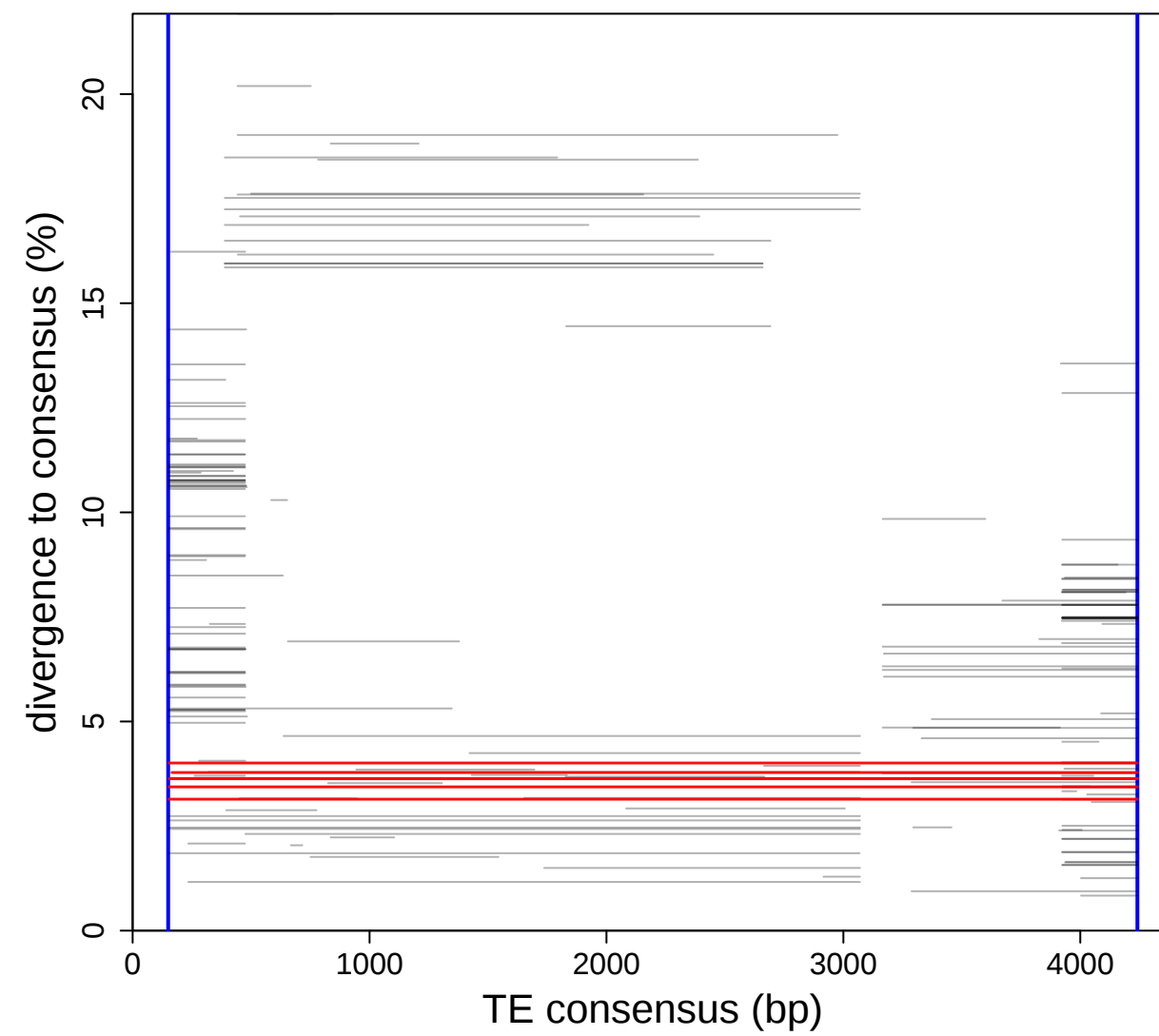
ed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 4042bp; fragments: 201; full length: 0 (>=3637.8bp)

After TETrimmer 4042 bp

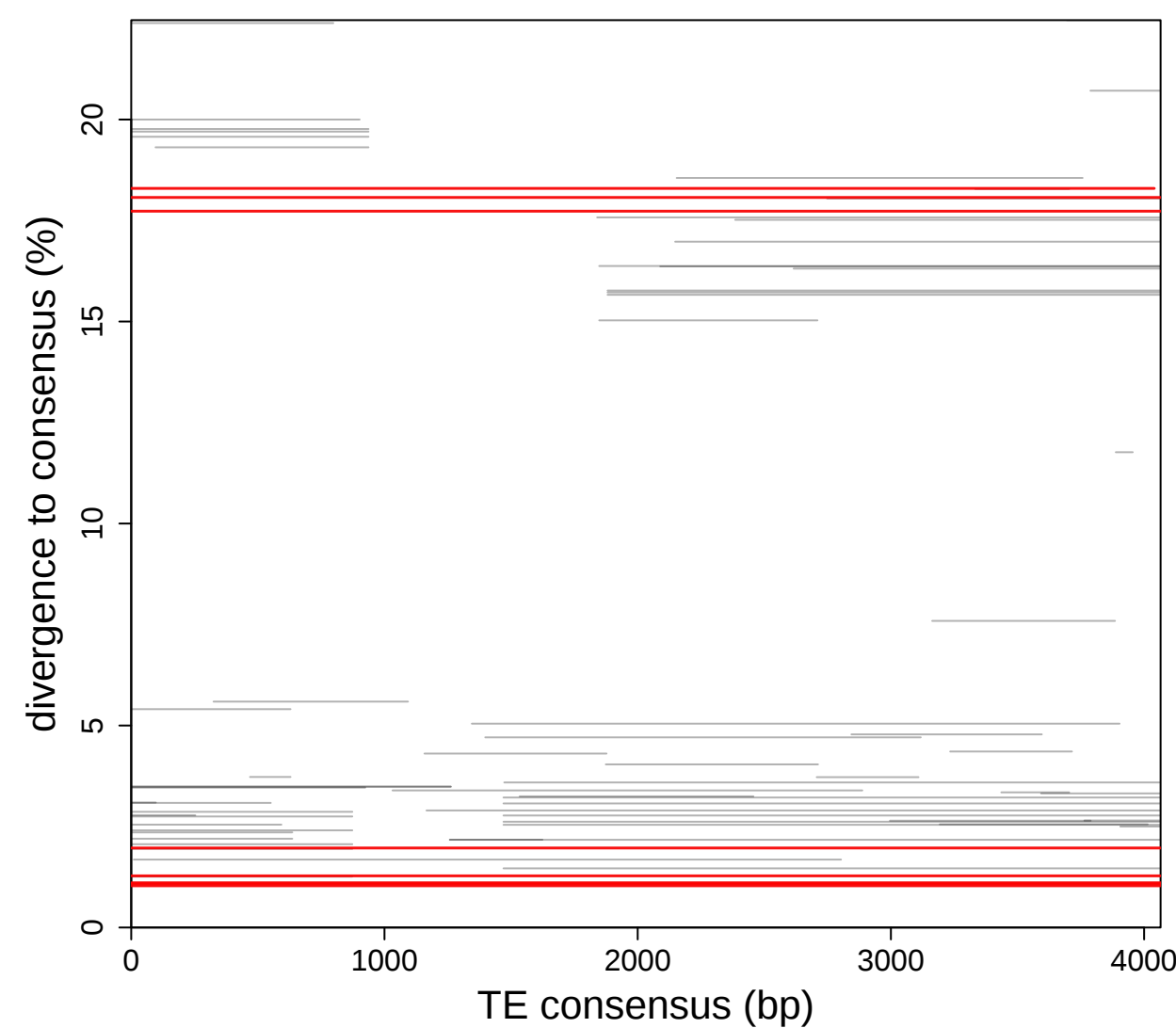


598.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_c
size: 4391bp; fragments: 182; full length: 6 (≥ 3951.9 bp)

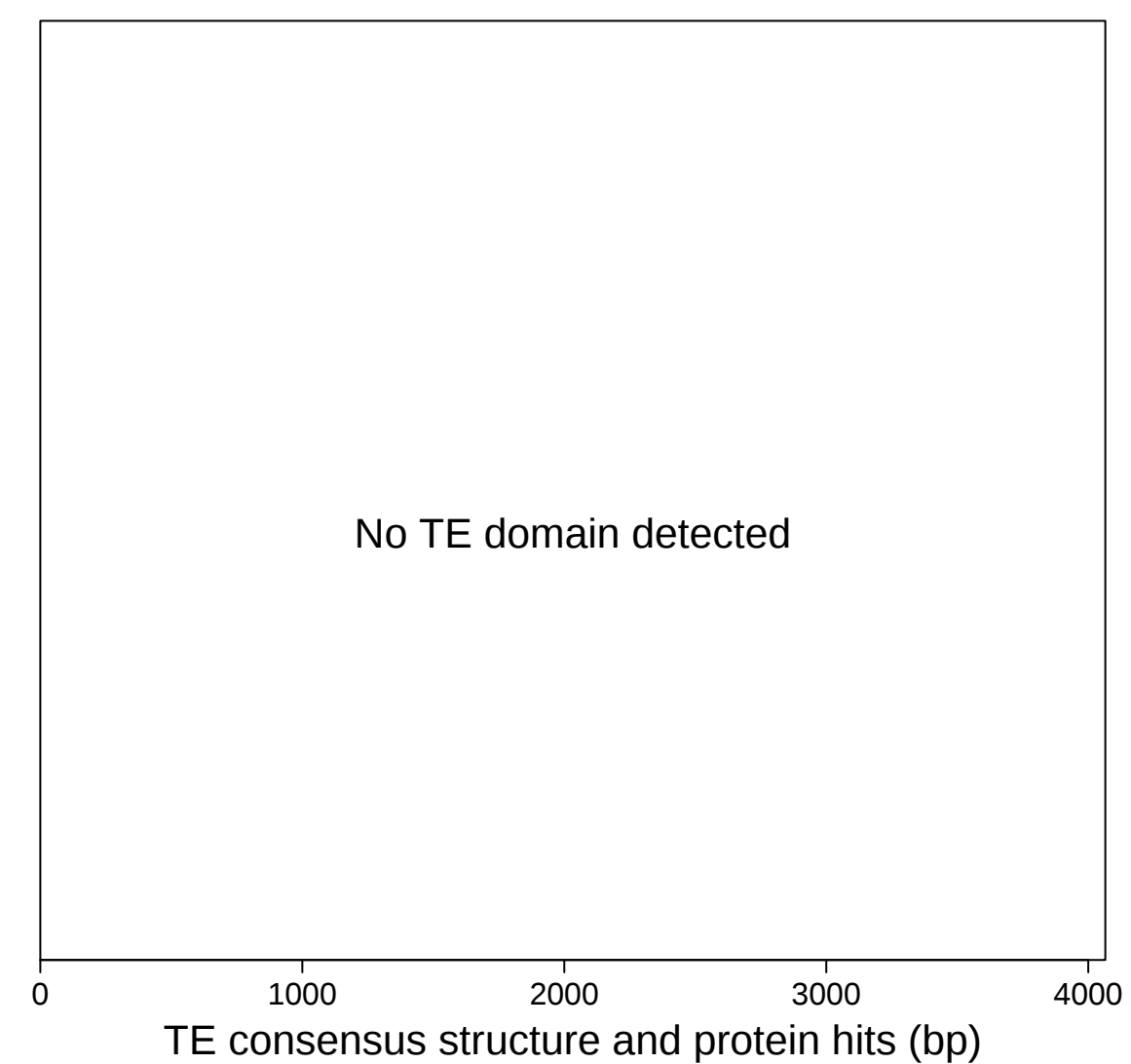
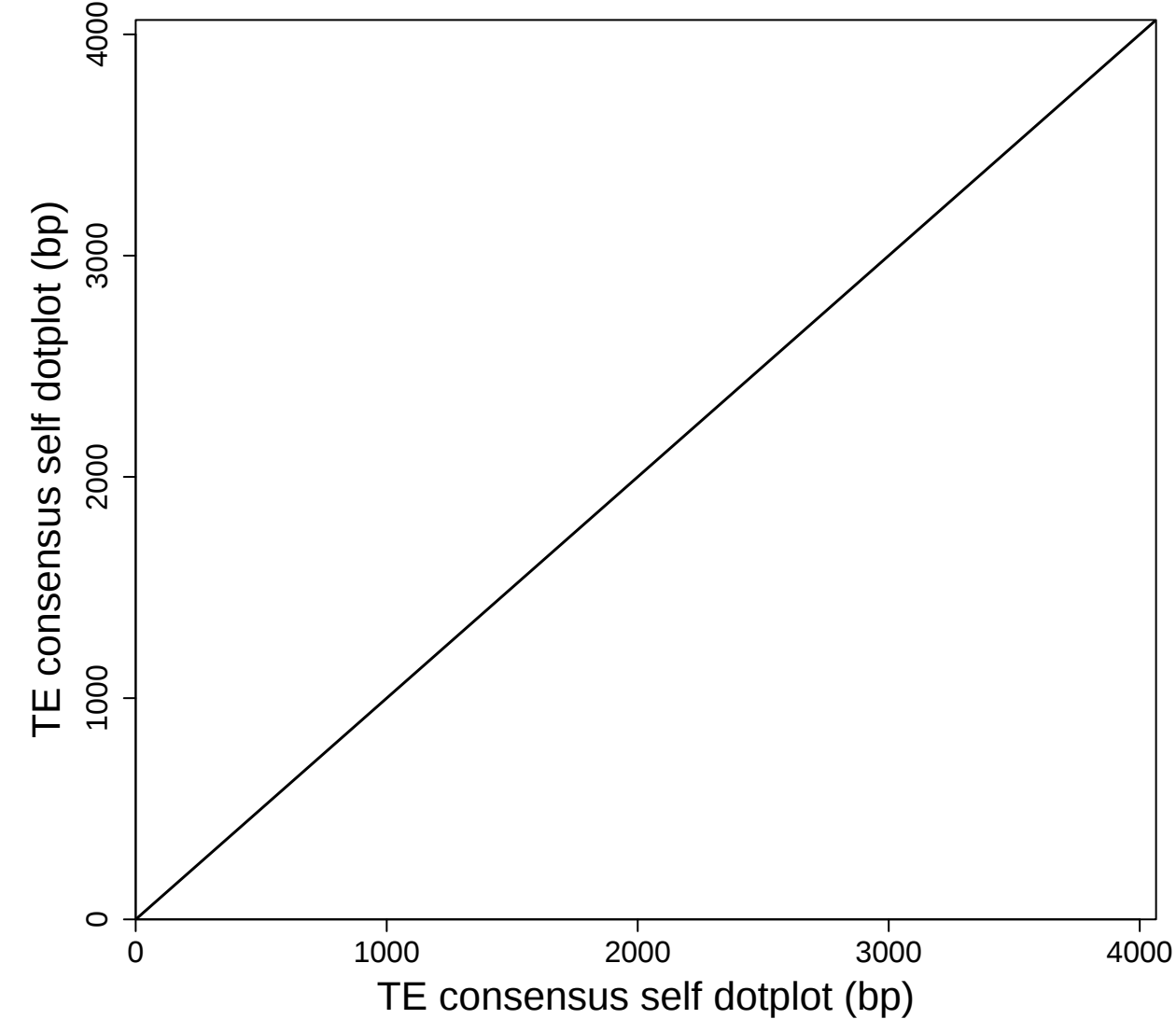
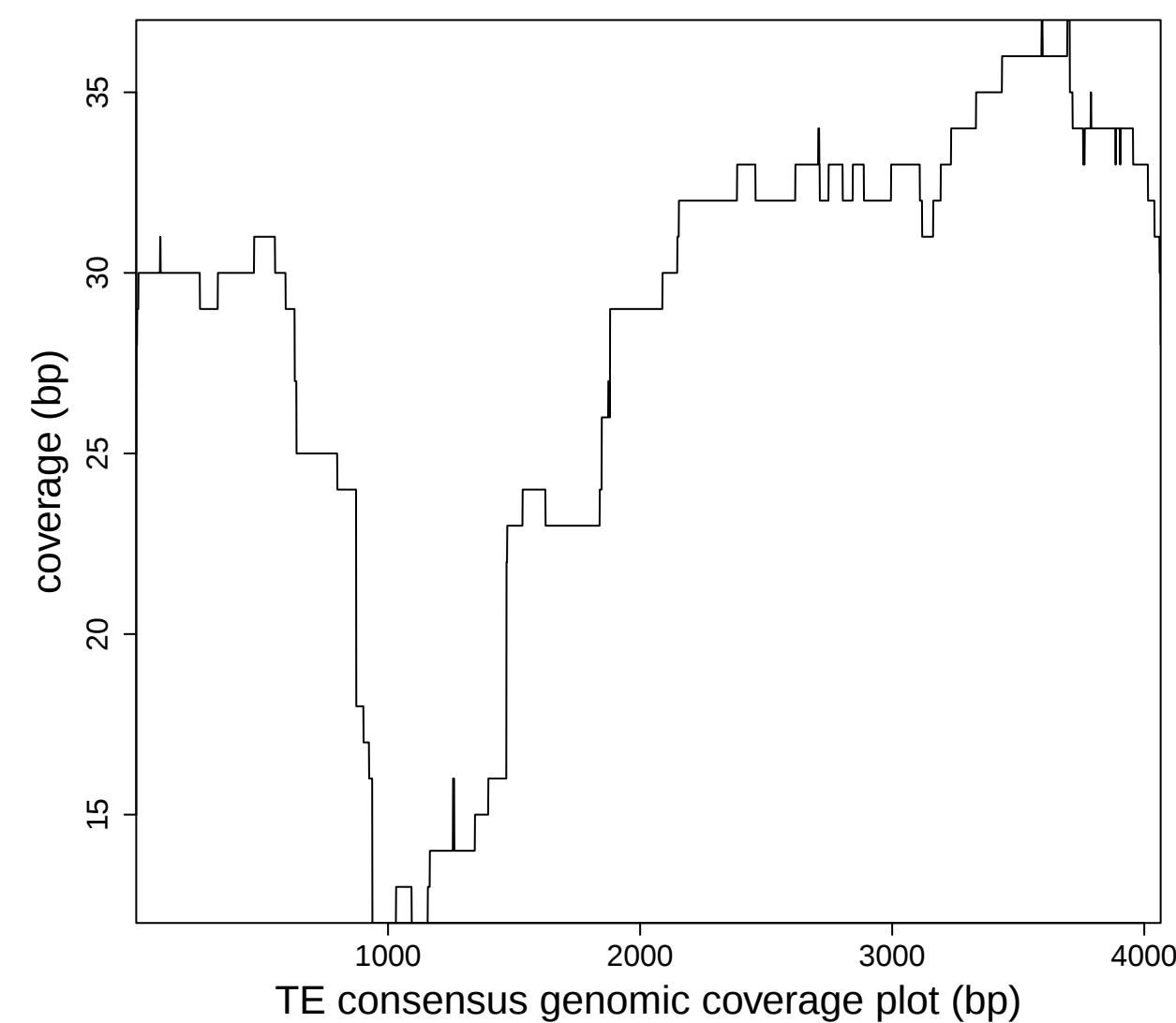
After TEtrimmer Extended plot Blue lines are boundaries



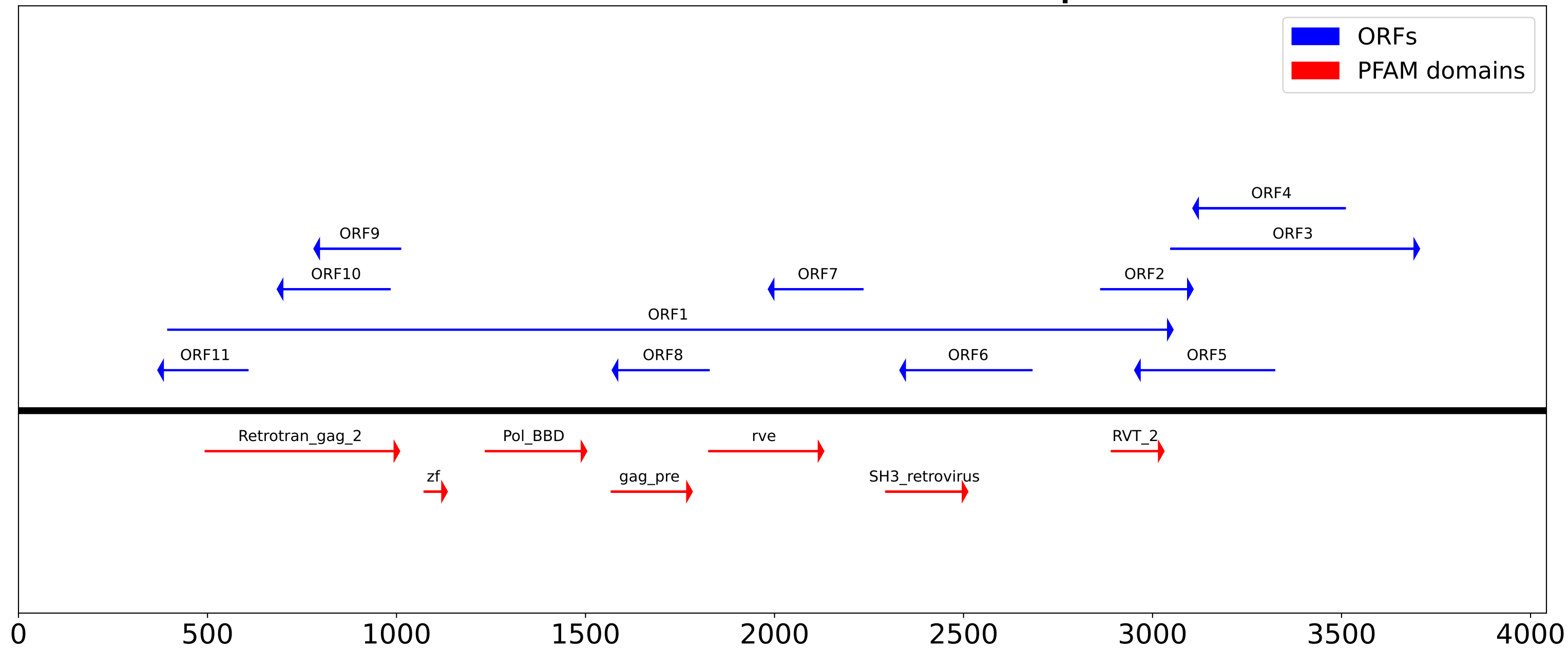
TE: ltr_1_family_598
size: 4065bp; fragments: 75; full length: 8 (≥ 3658.5 bp)



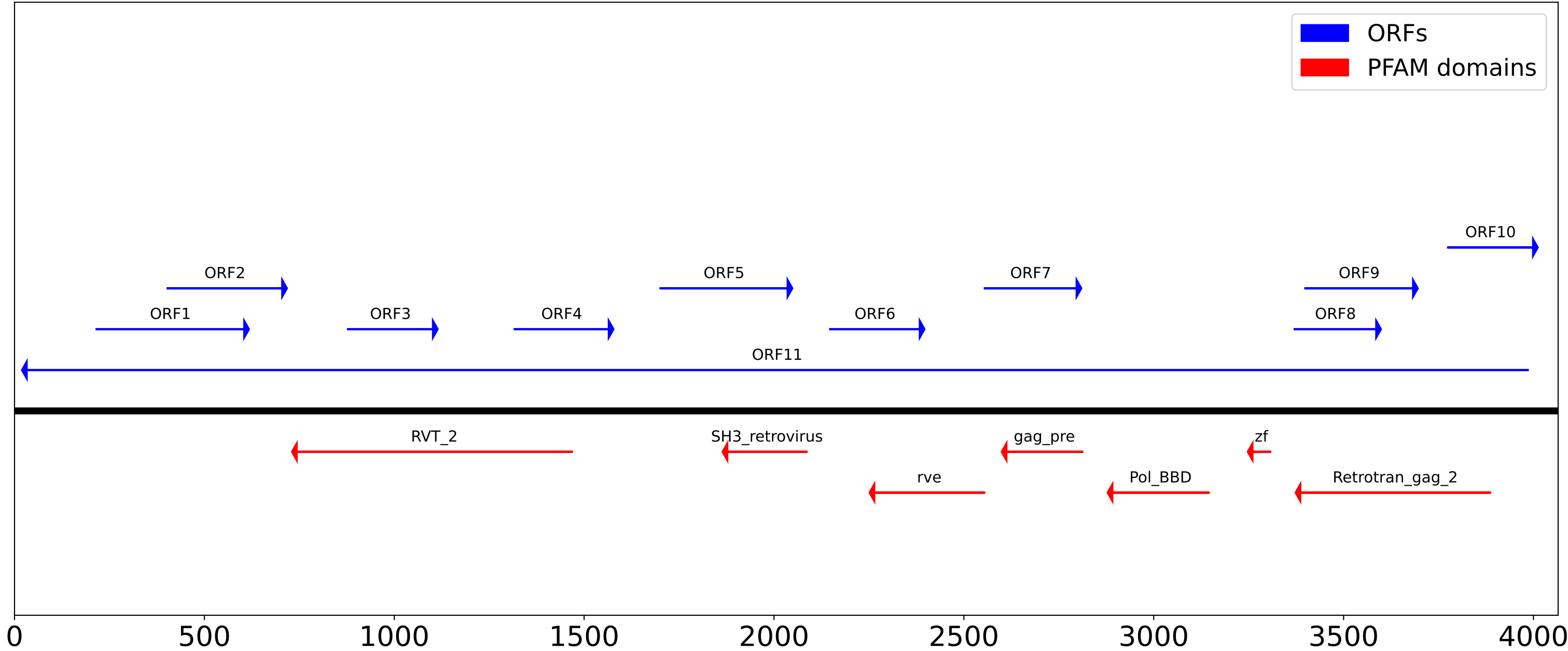
Before TEtrimmer 4065 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

