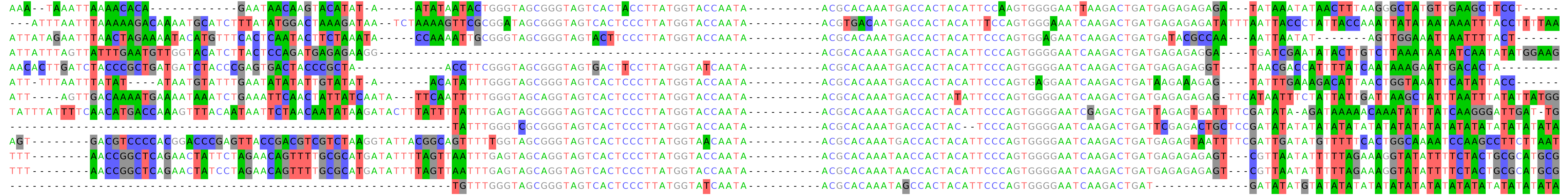


Start crop Point

End crop Point

MSA length = 3428



Start crop Point

End crop Point

0

10

0

500

1000

1500

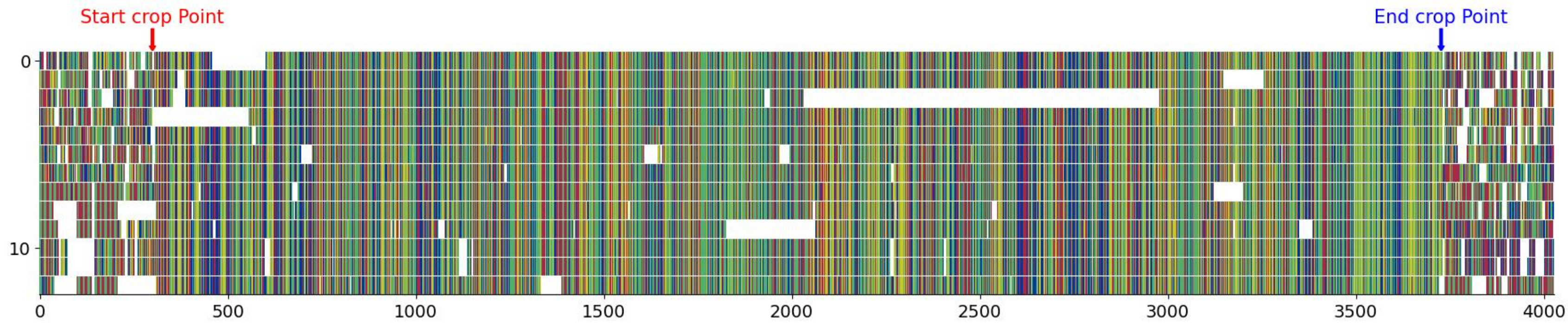
2000

2500

3000

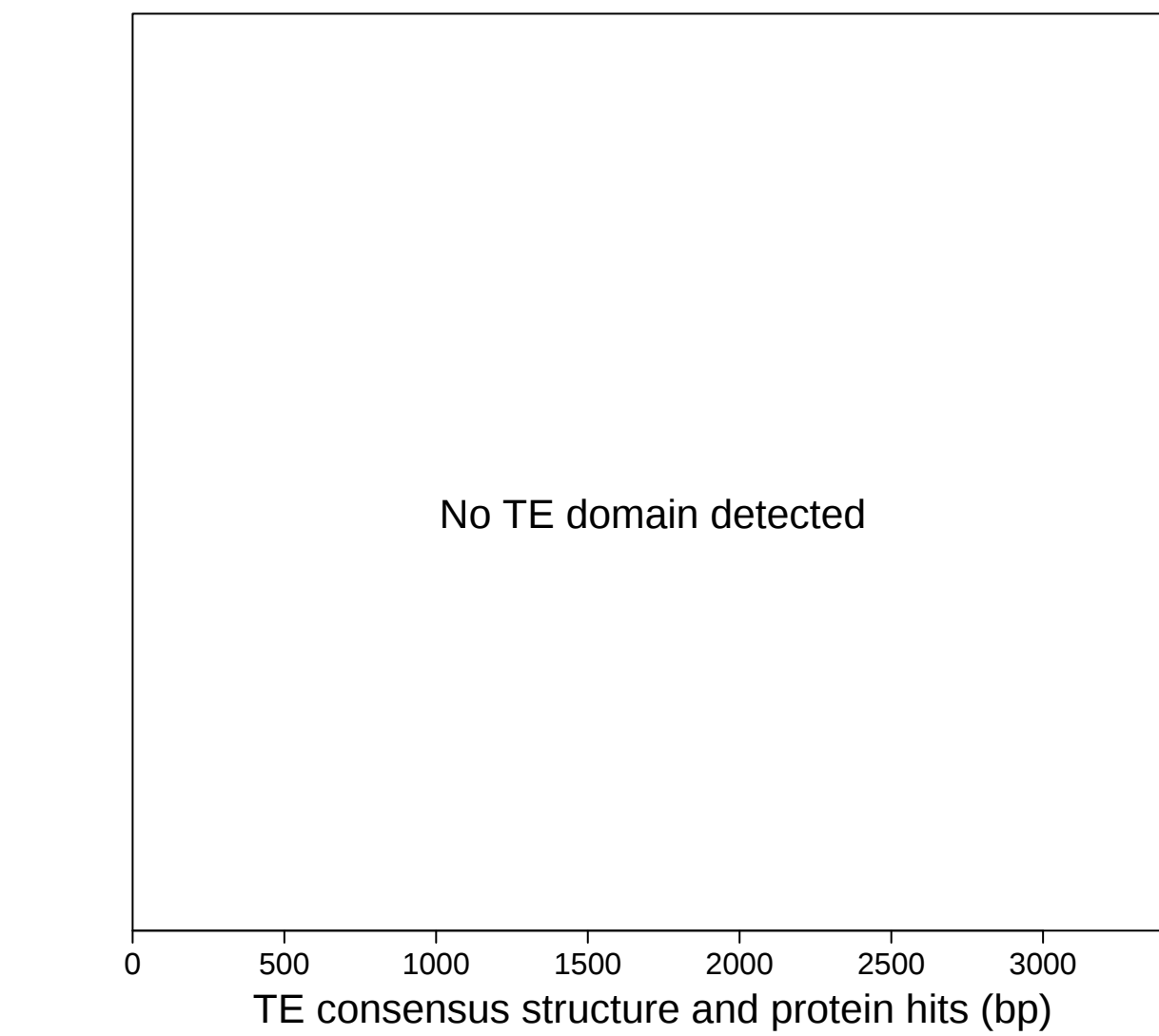
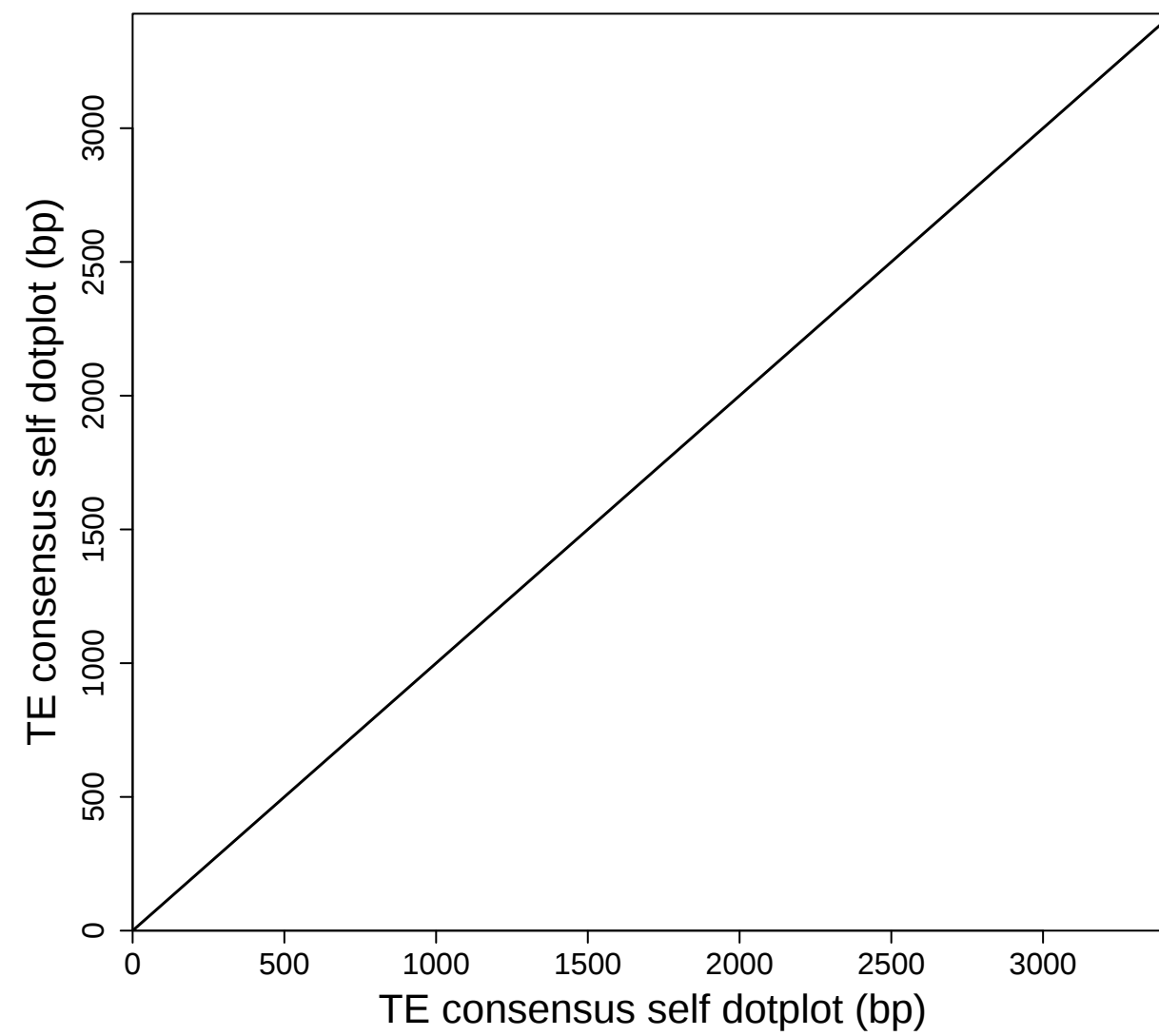
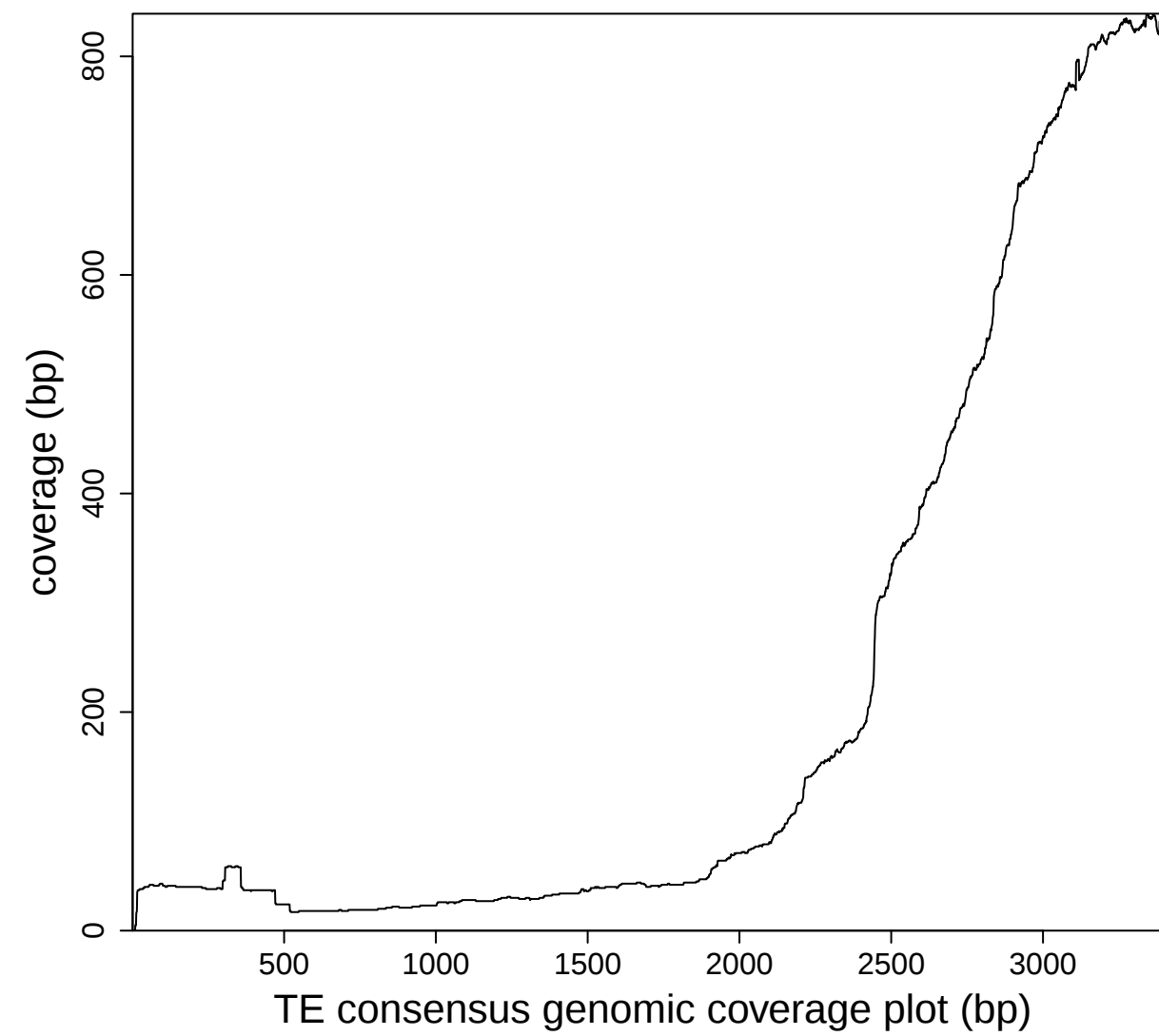
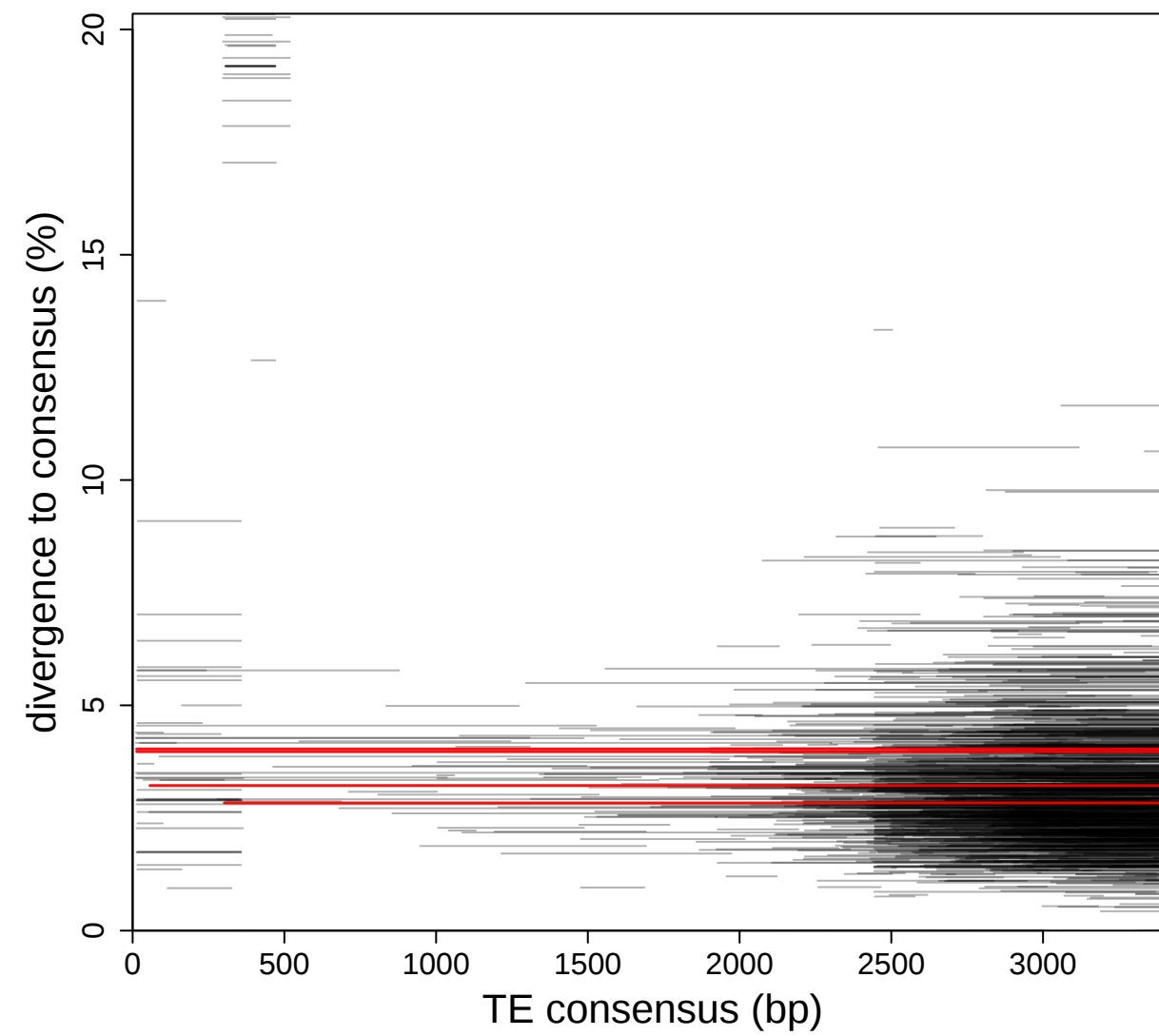
3500

4000



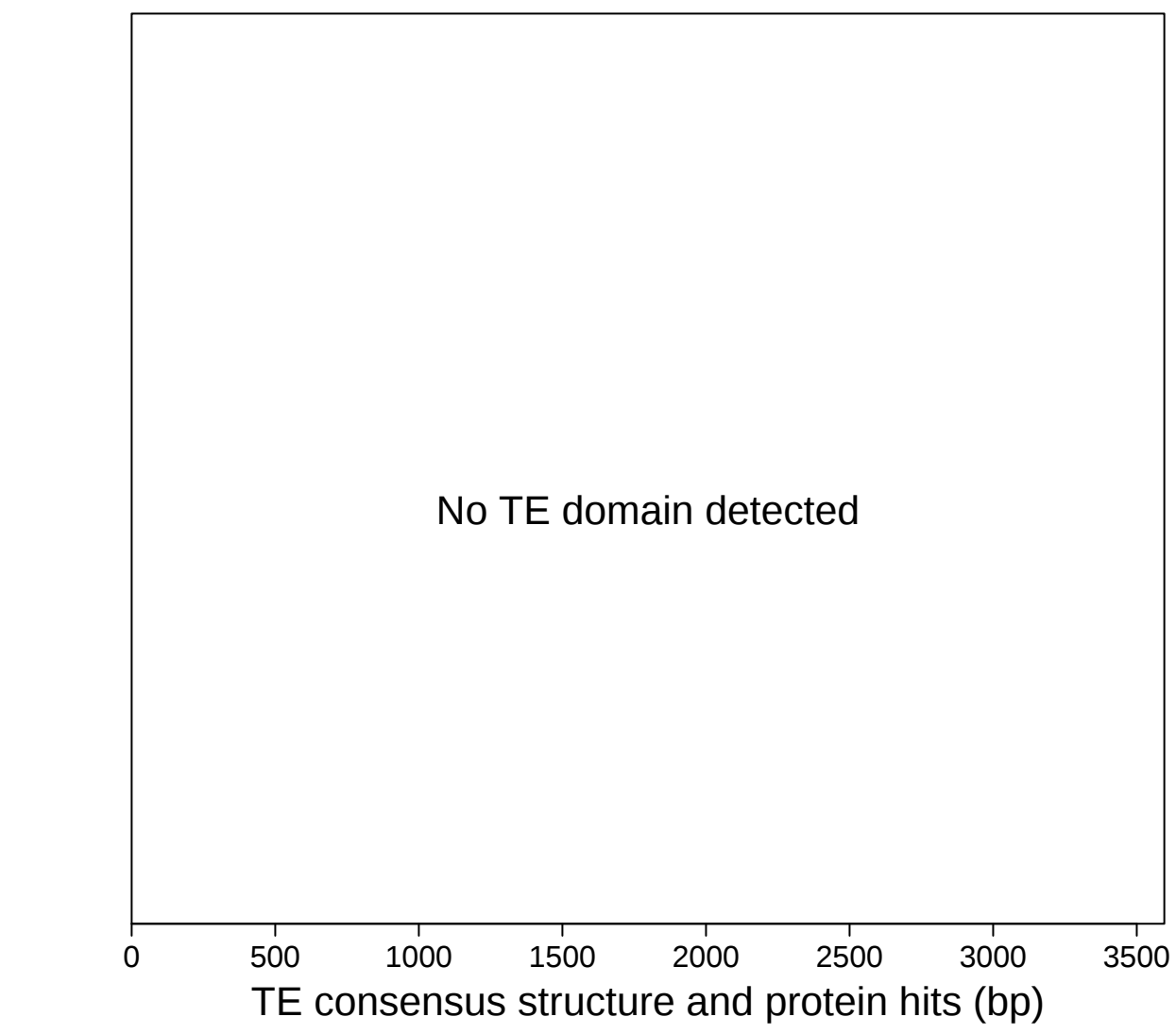
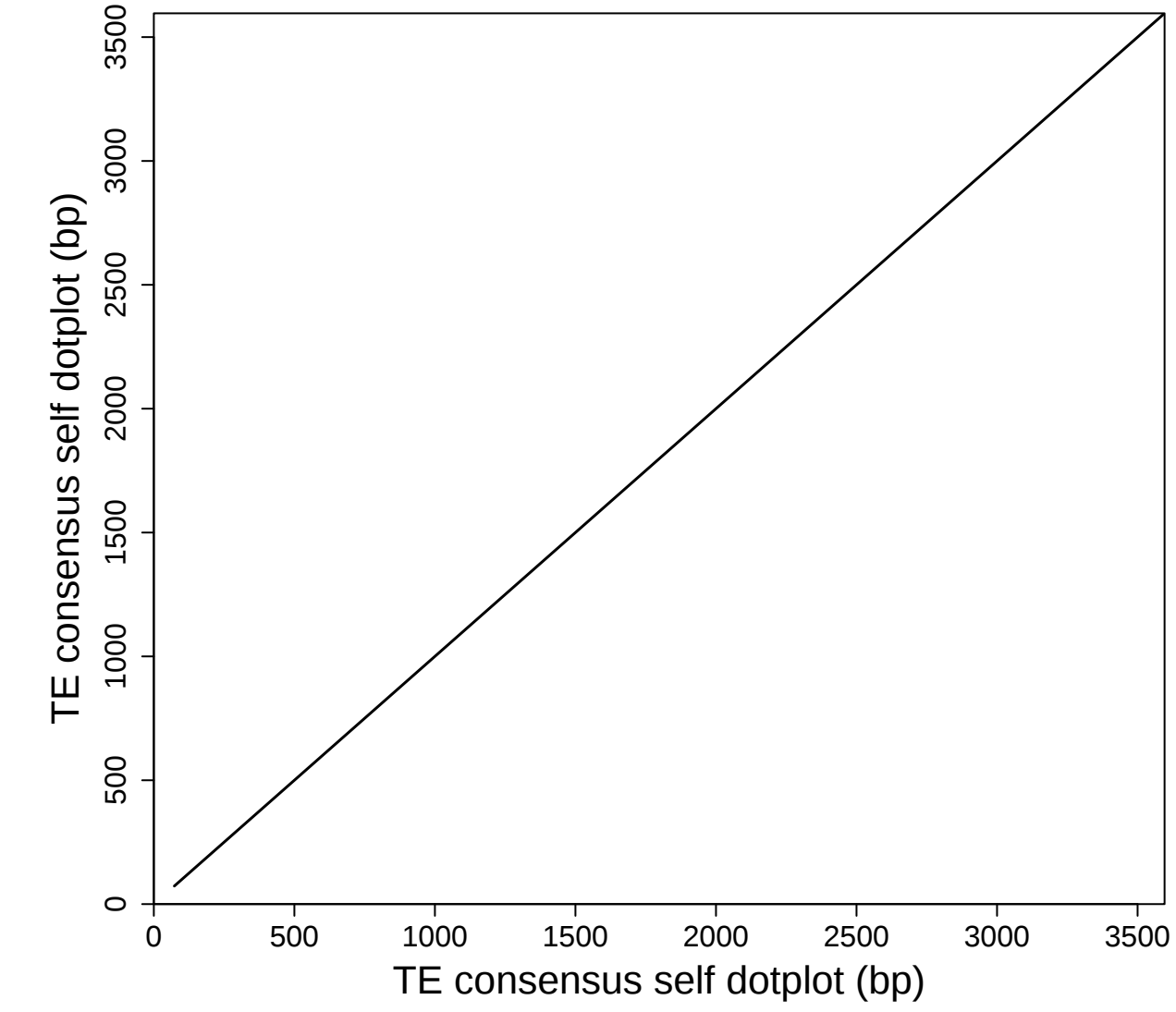
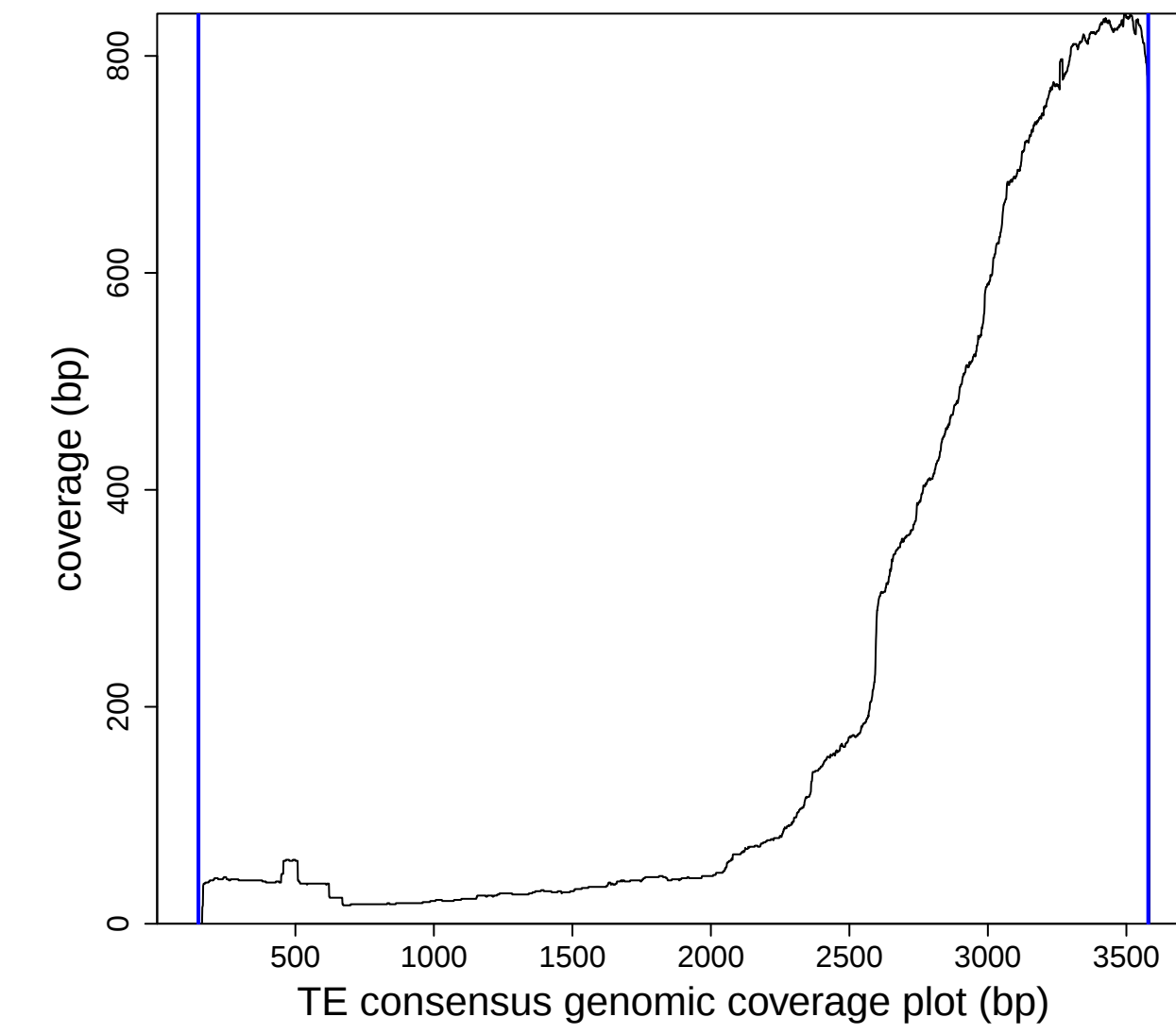
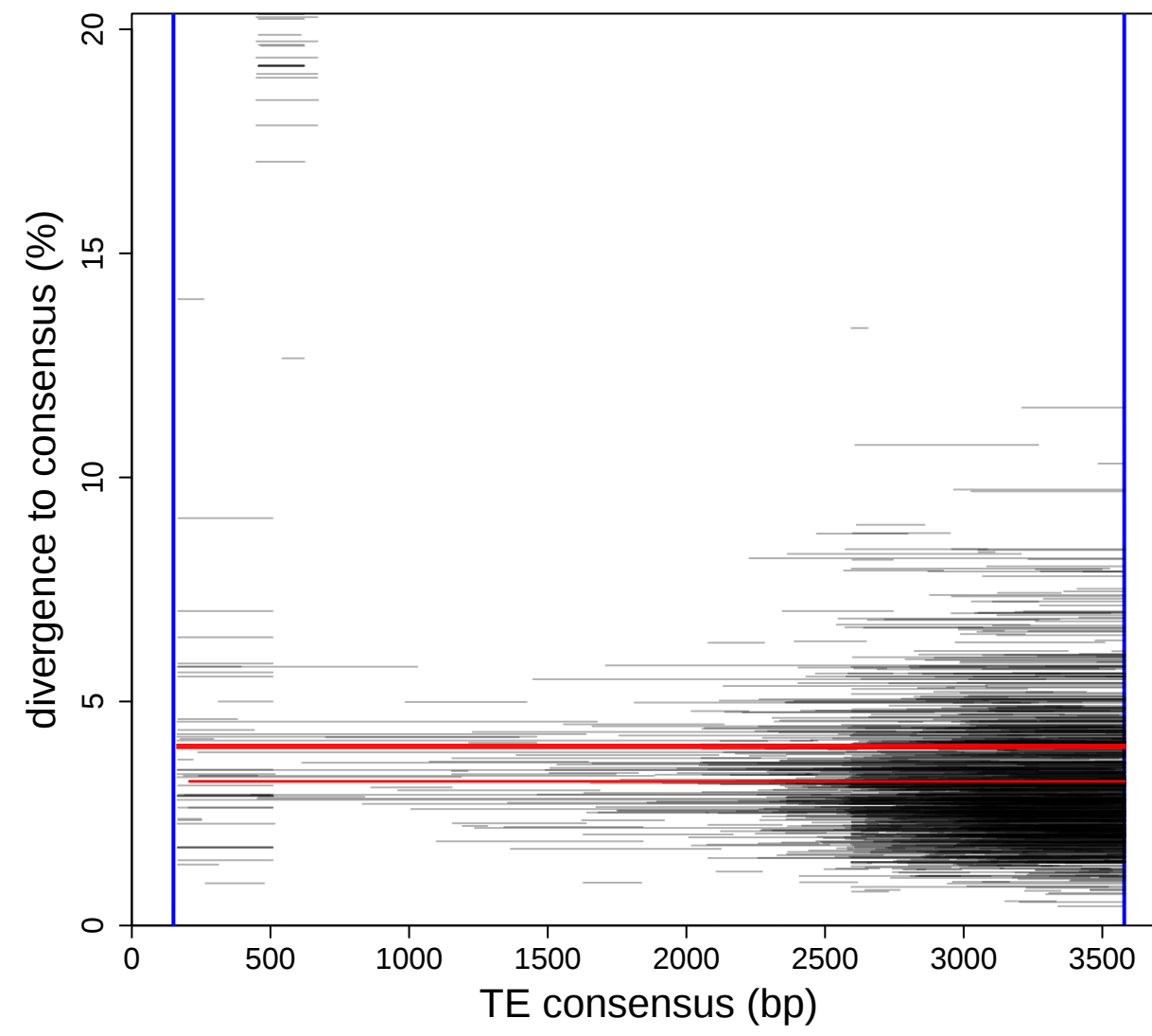
bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.fa
size: 3428bp; fragments: 1267; full length: 4 (≥ 3085.2 bp)

After TEtrimmer 3428 bp



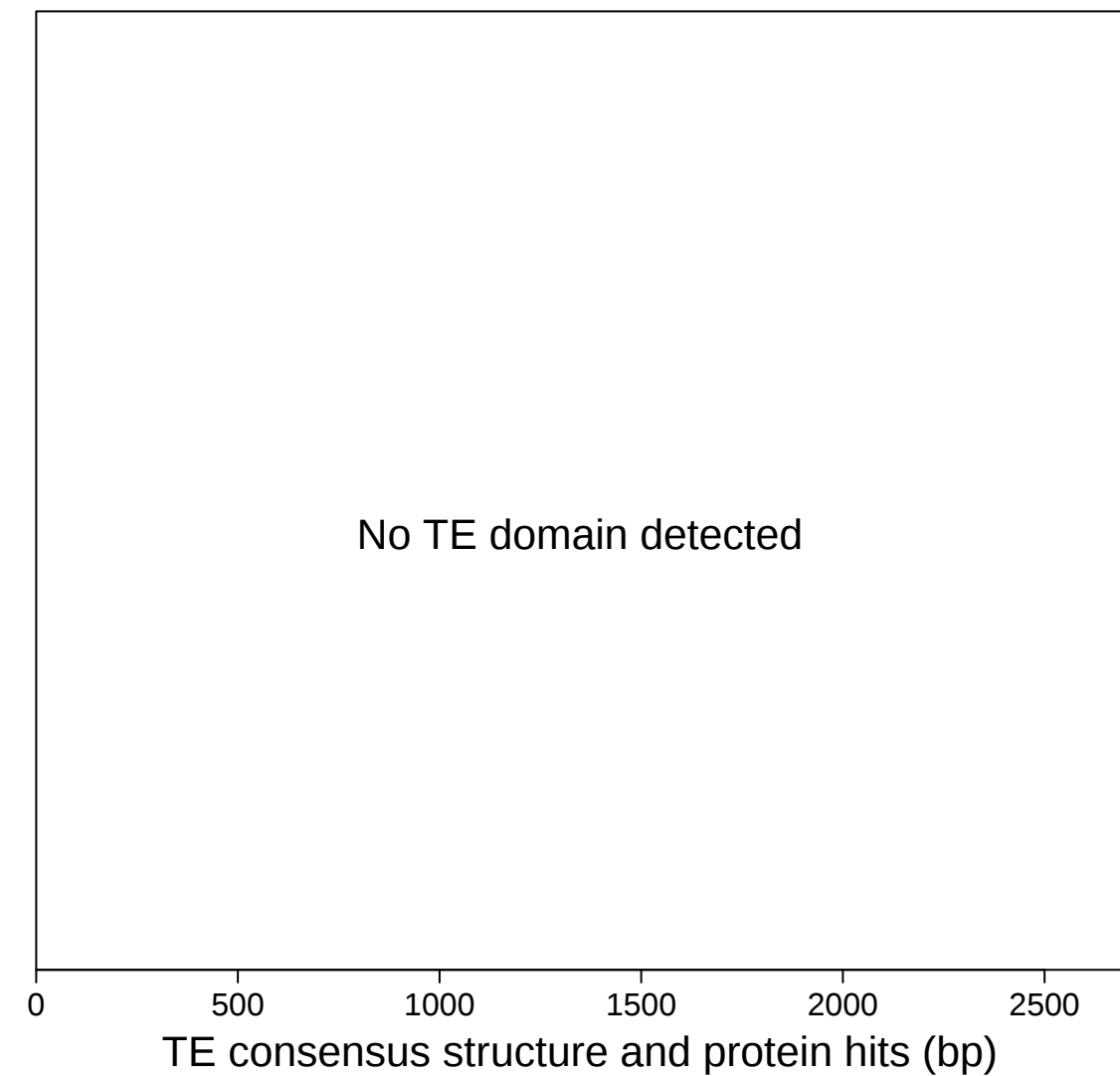
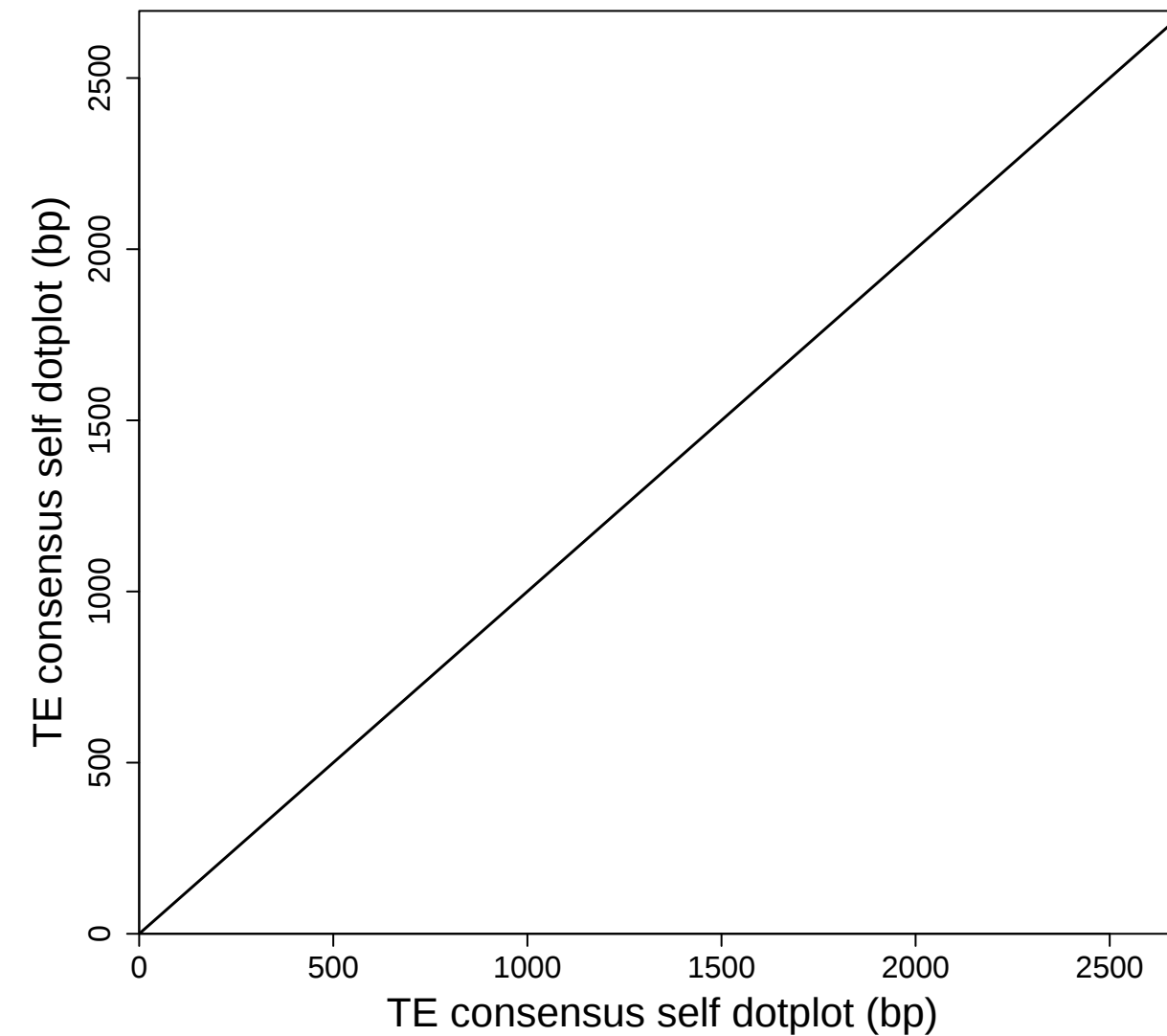
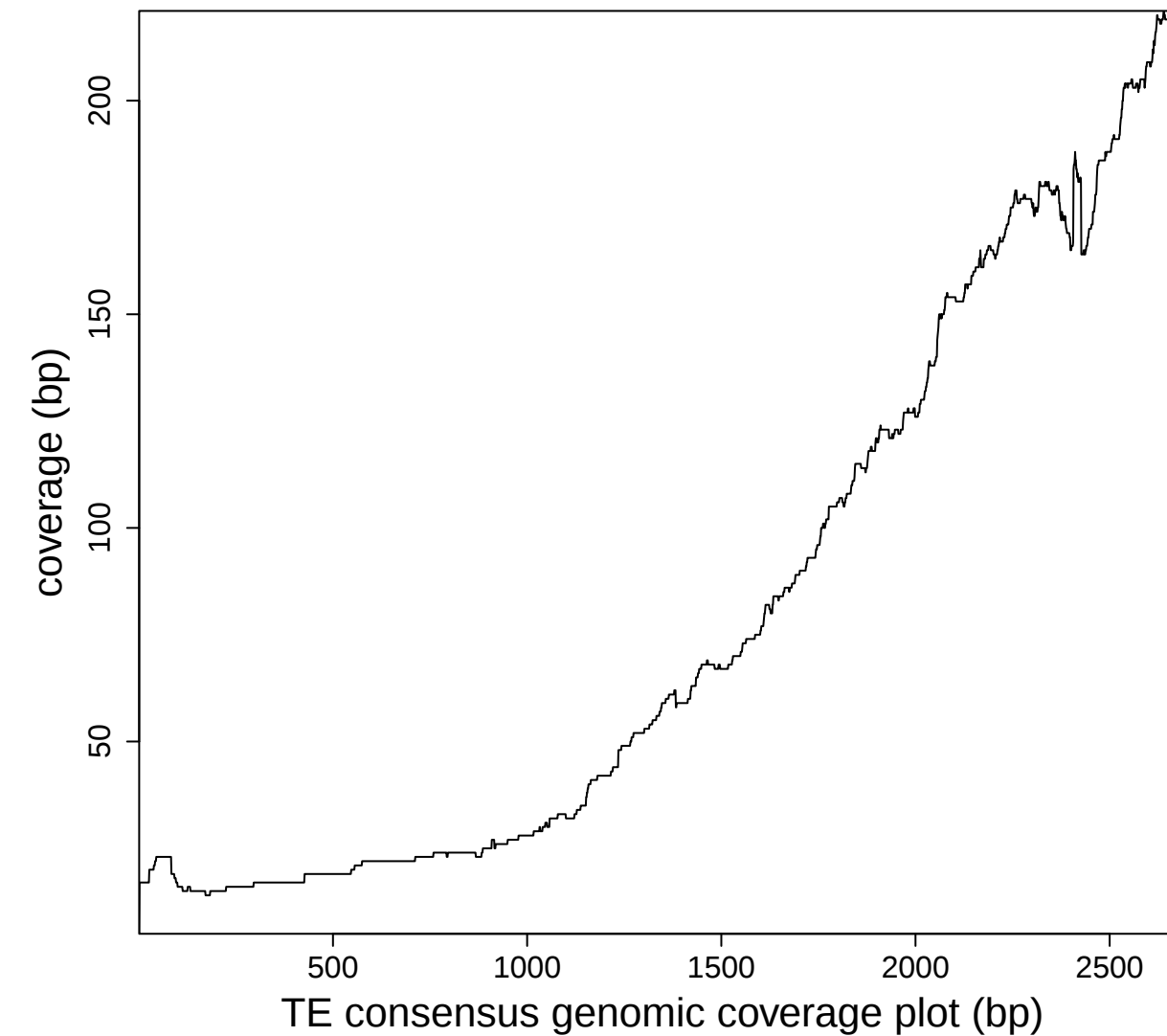
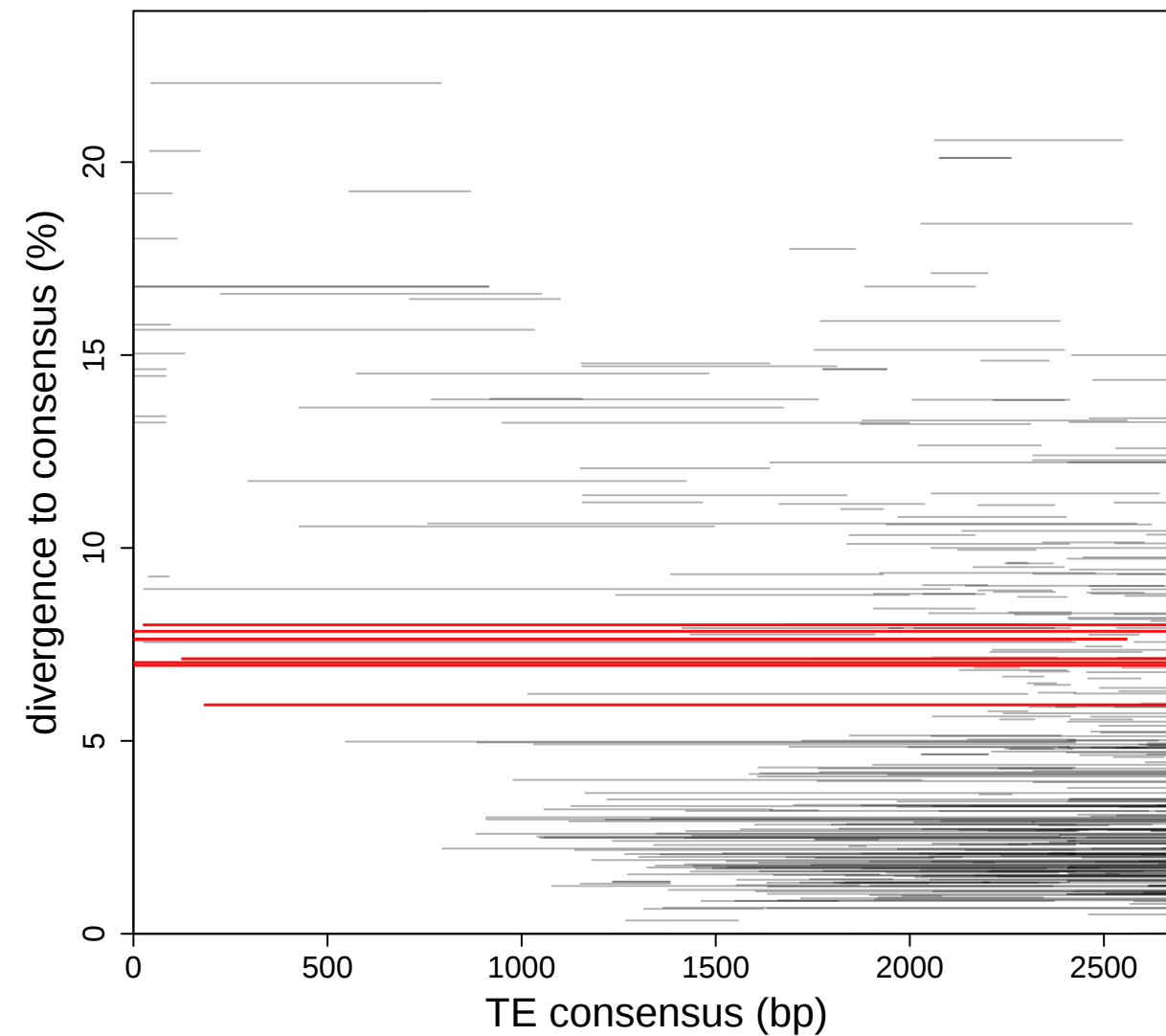
f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 3729bp; fragments: 1270; full length: 3 (>=3356.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

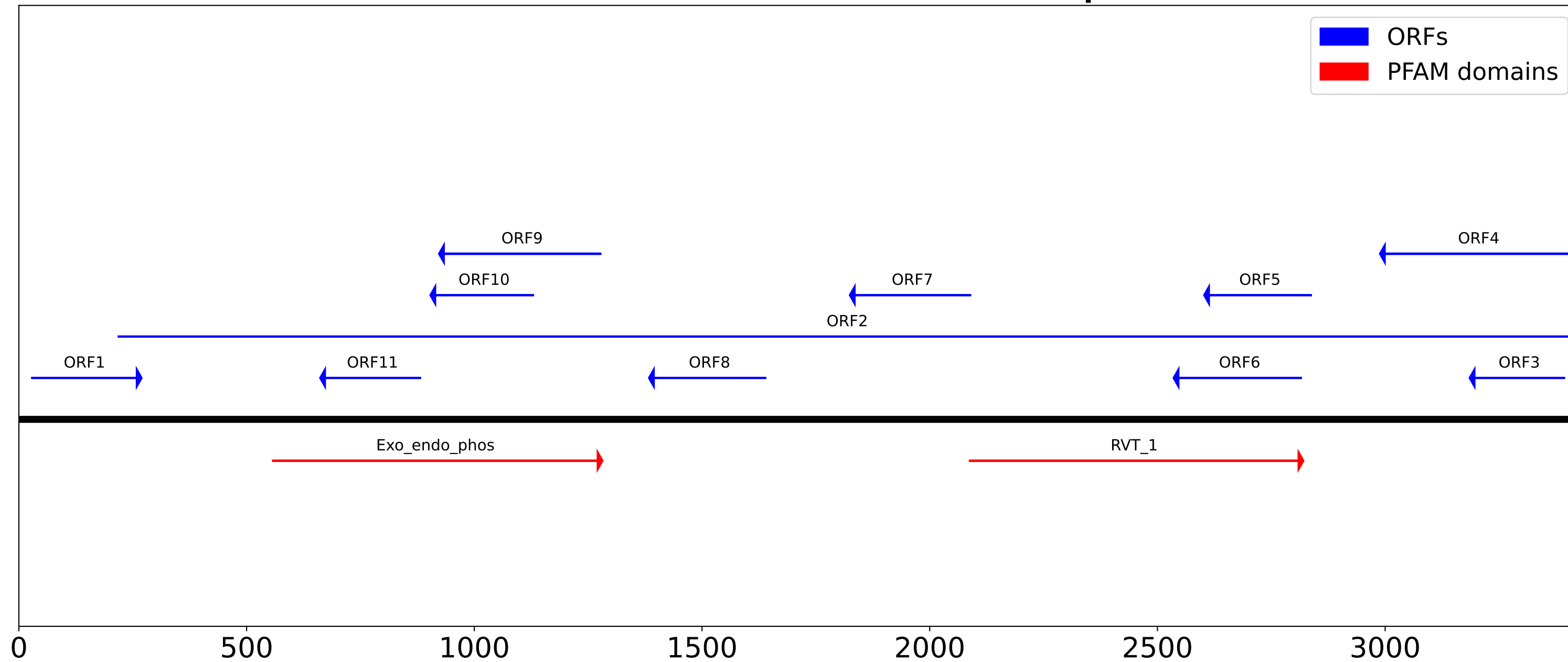


TE: rnd_5_family_565
size: 2696bp; fragments: 397; full length: 7 (≥ 2426.4 bp)

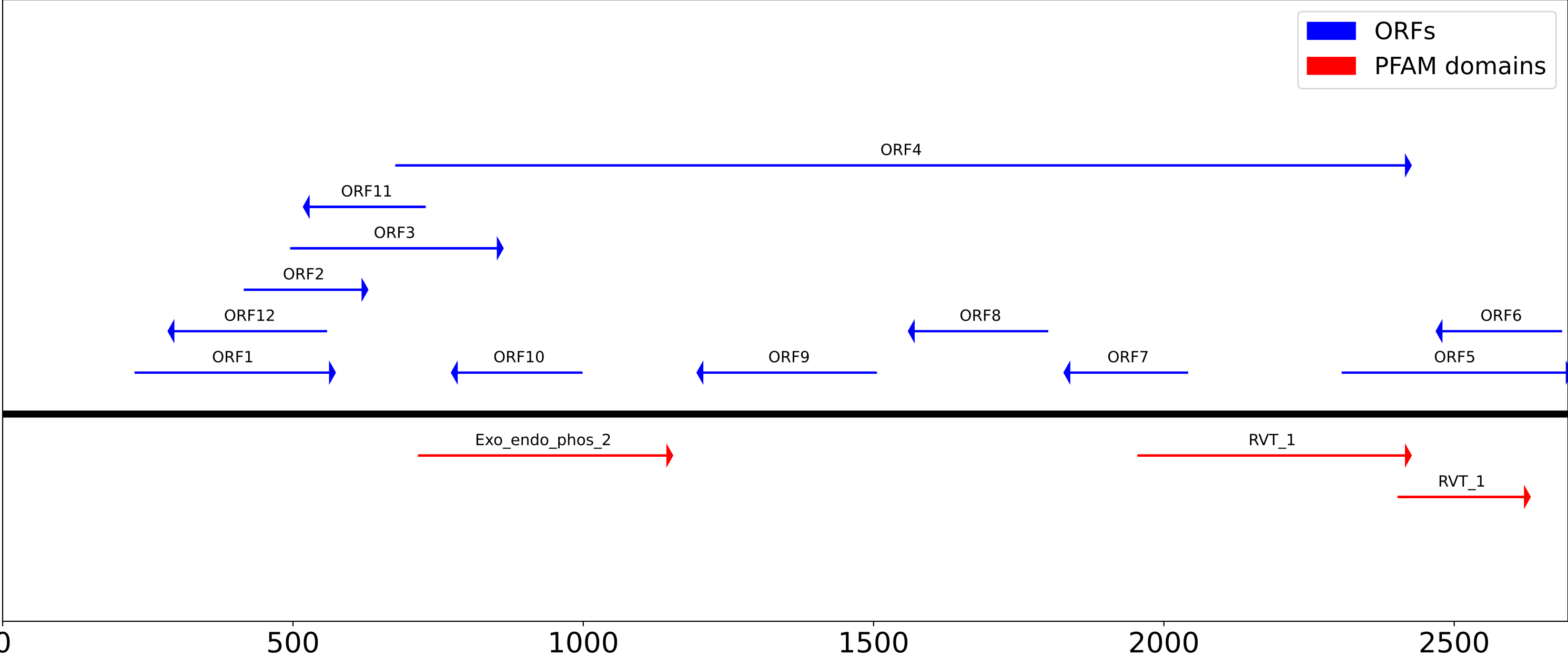
Before TEtrimmer 2696 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

