

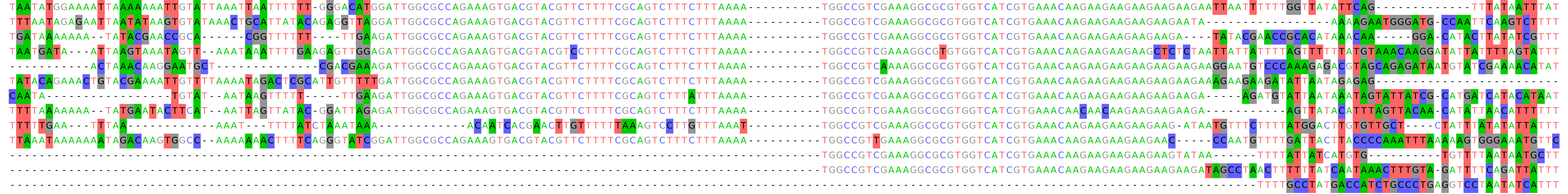
Start crop Point



End crop Point



MSA length = 5895



Start crop Point

End crop Point

0

10

0

1000

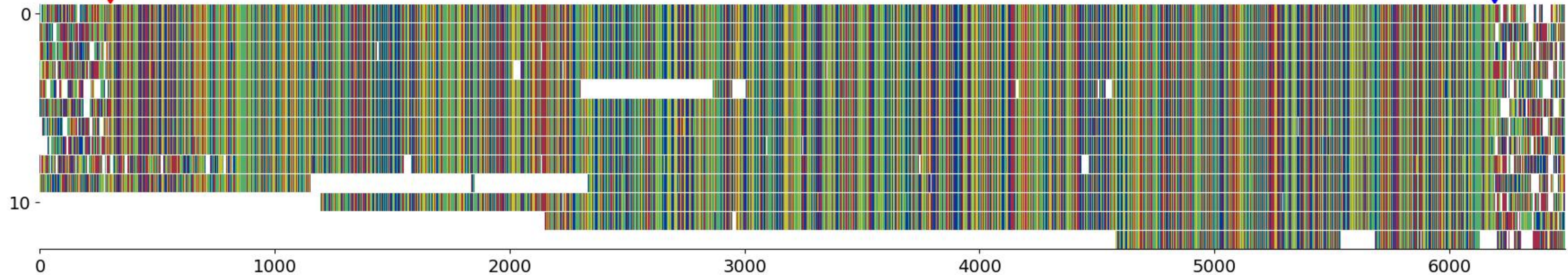
2000

3000

4000

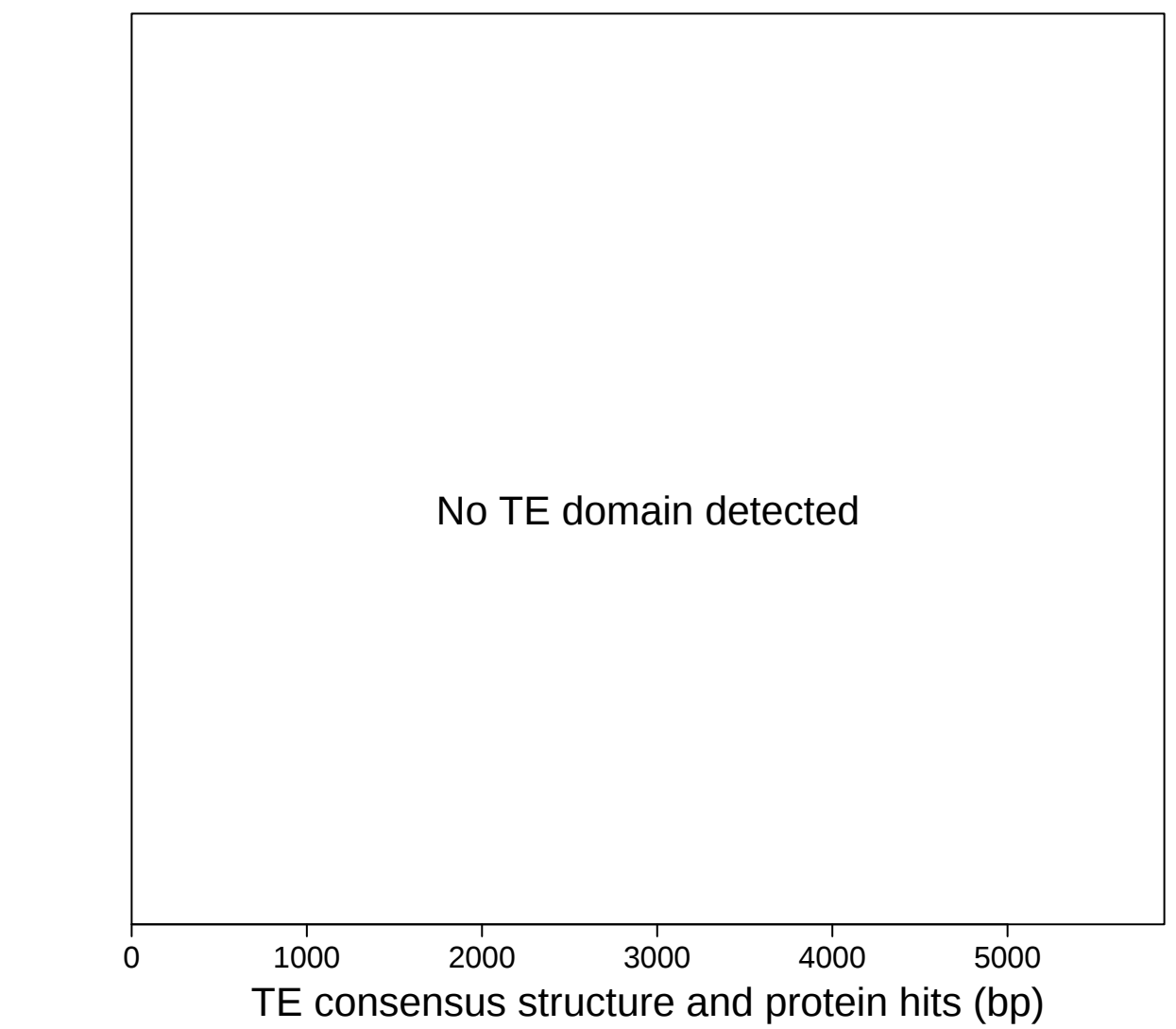
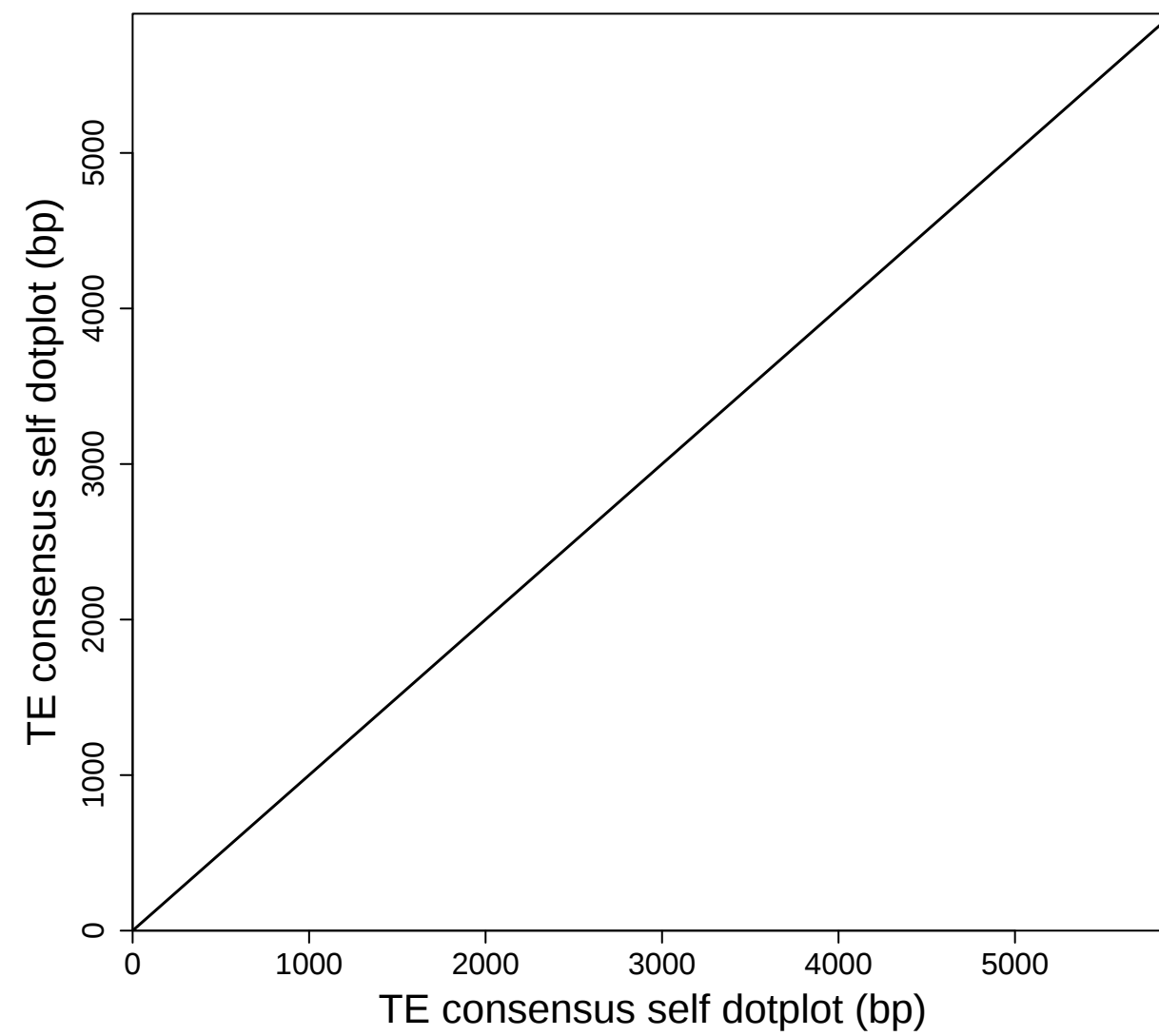
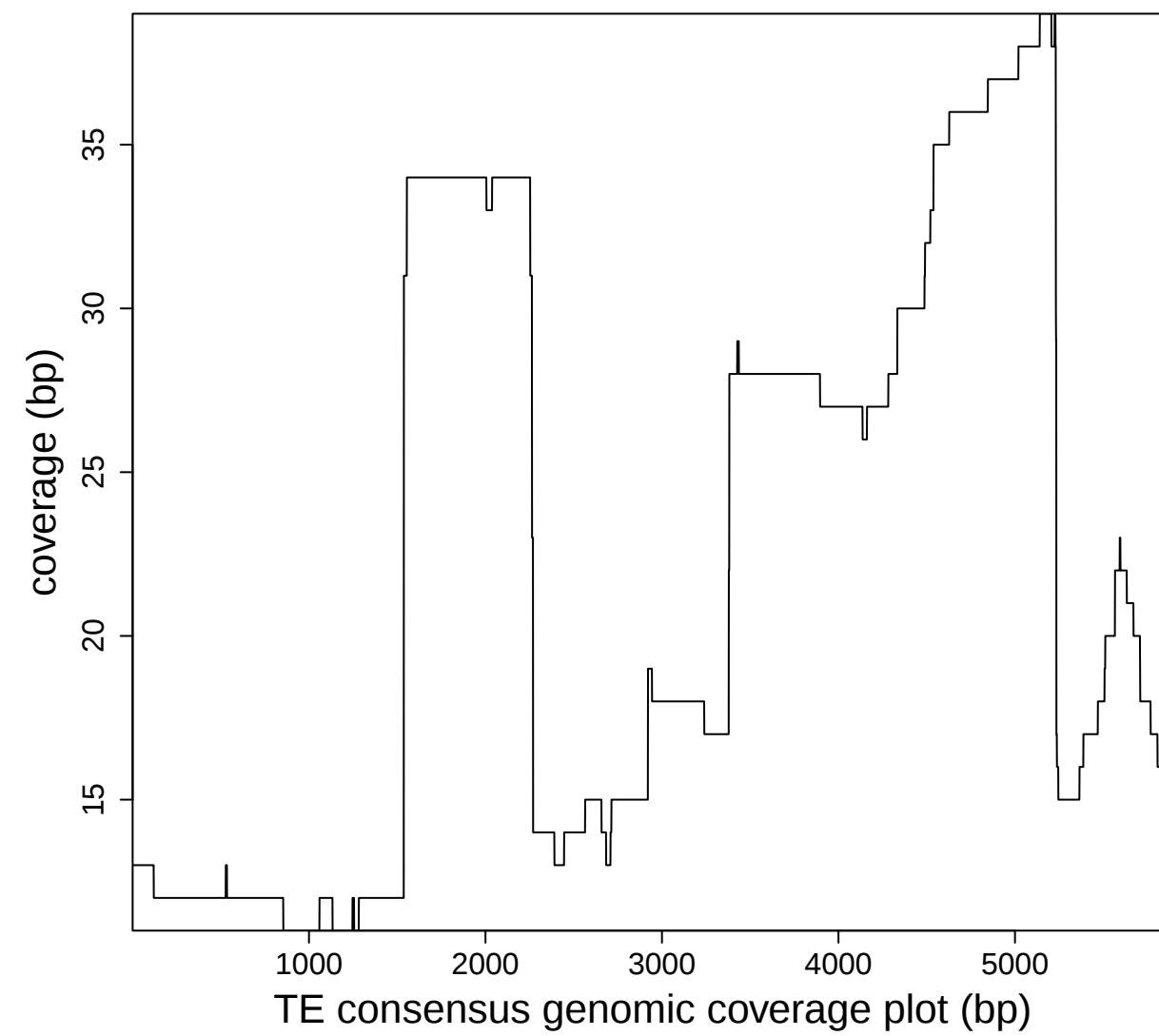
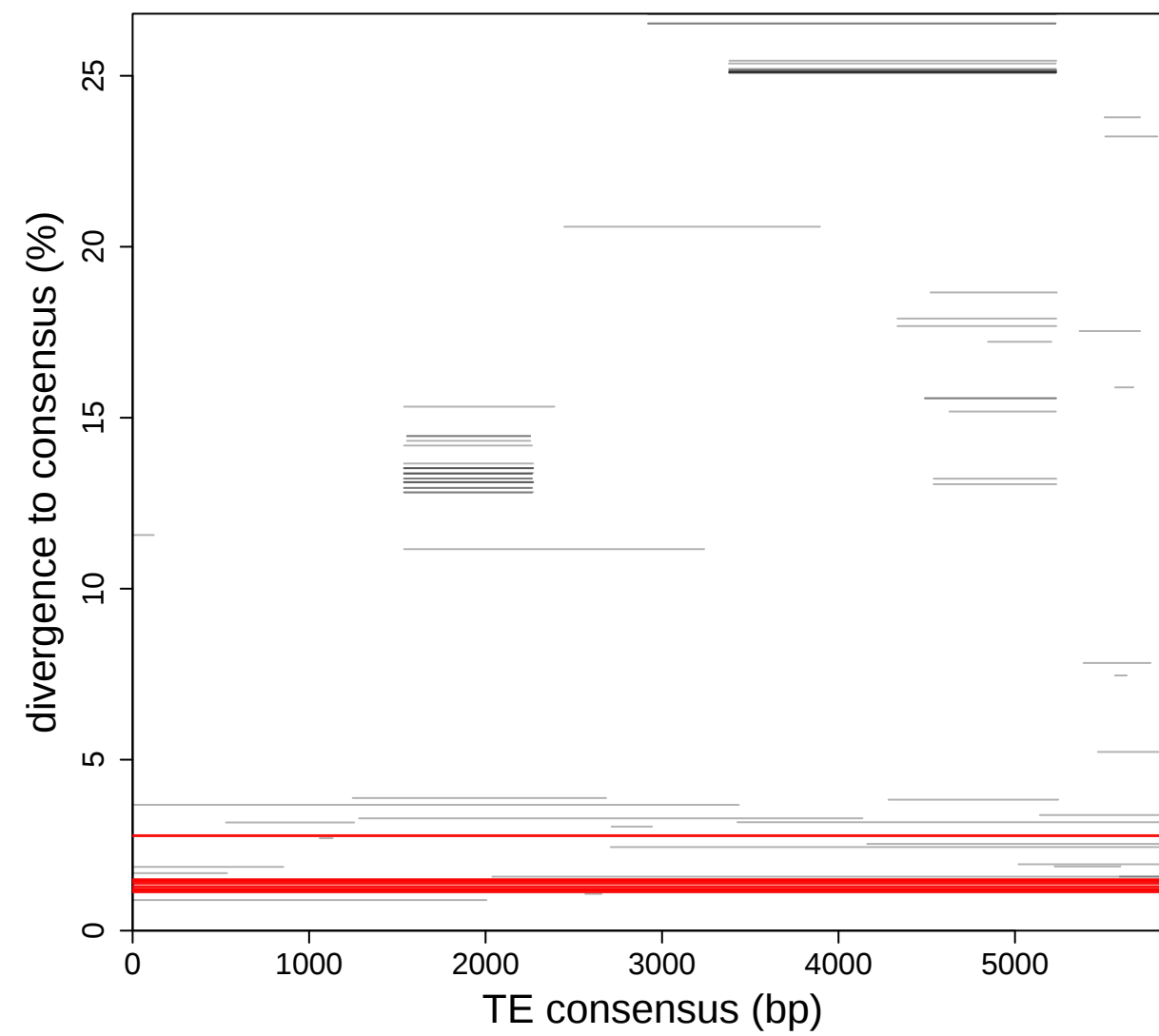
5000

6000



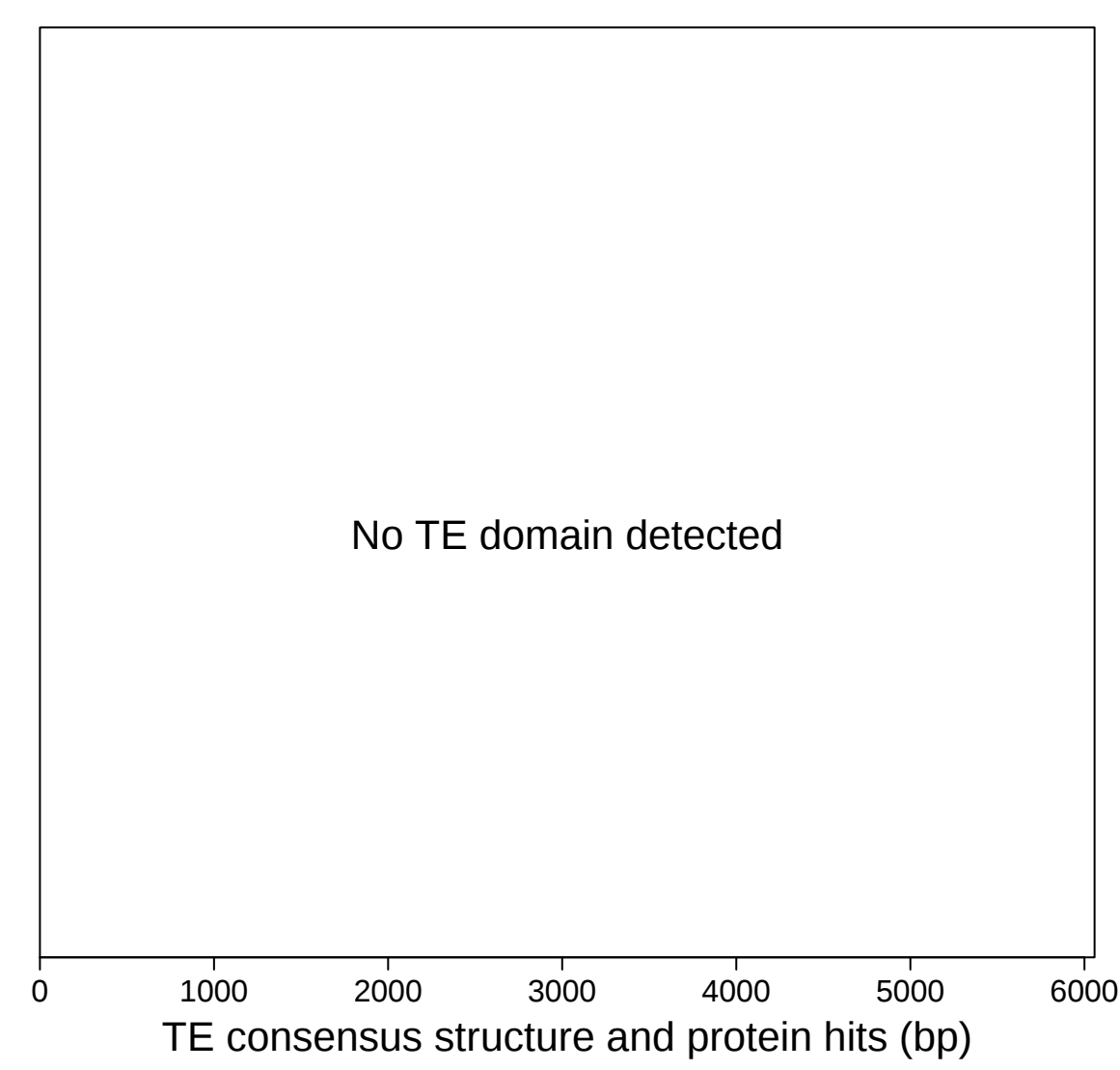
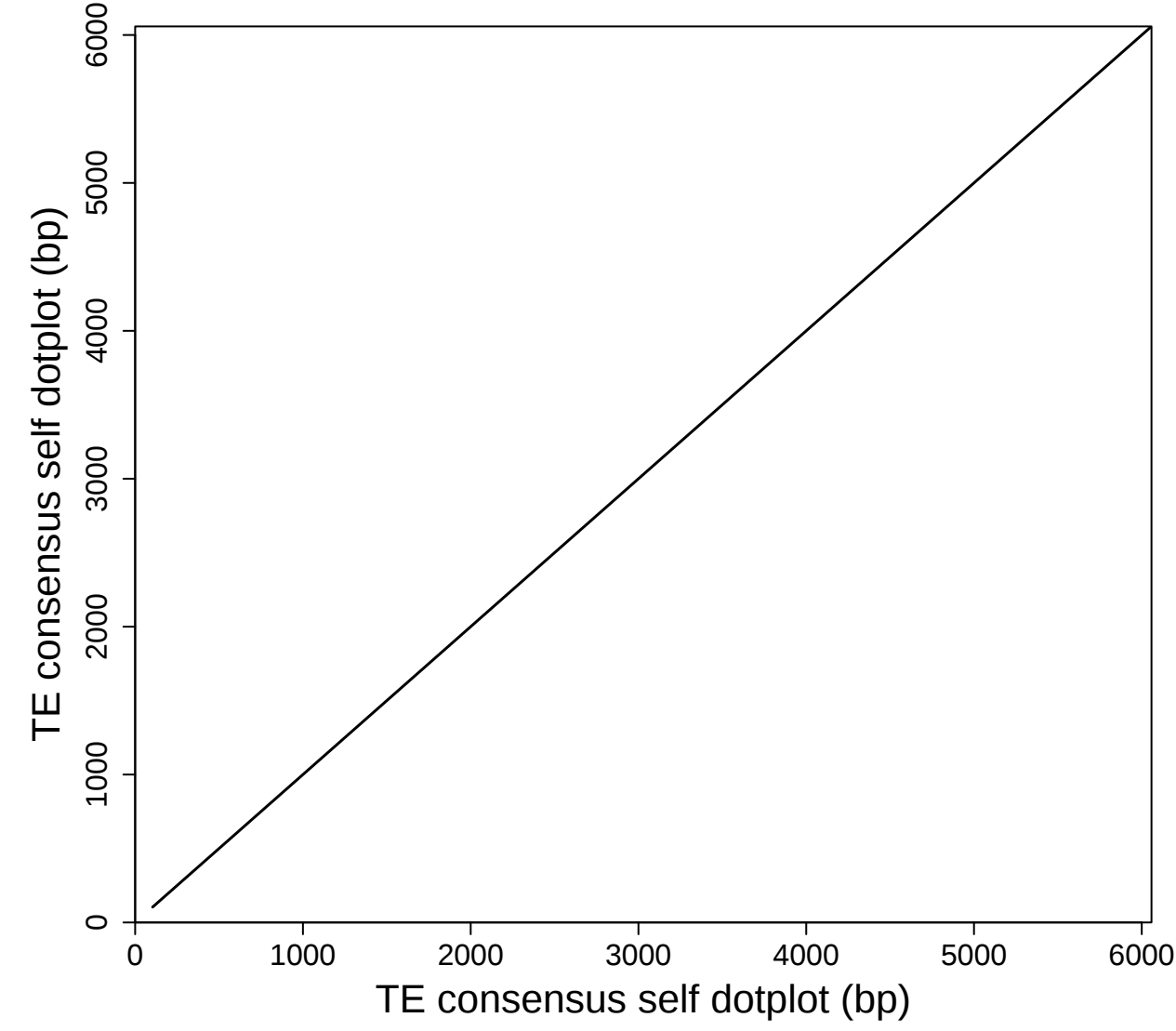
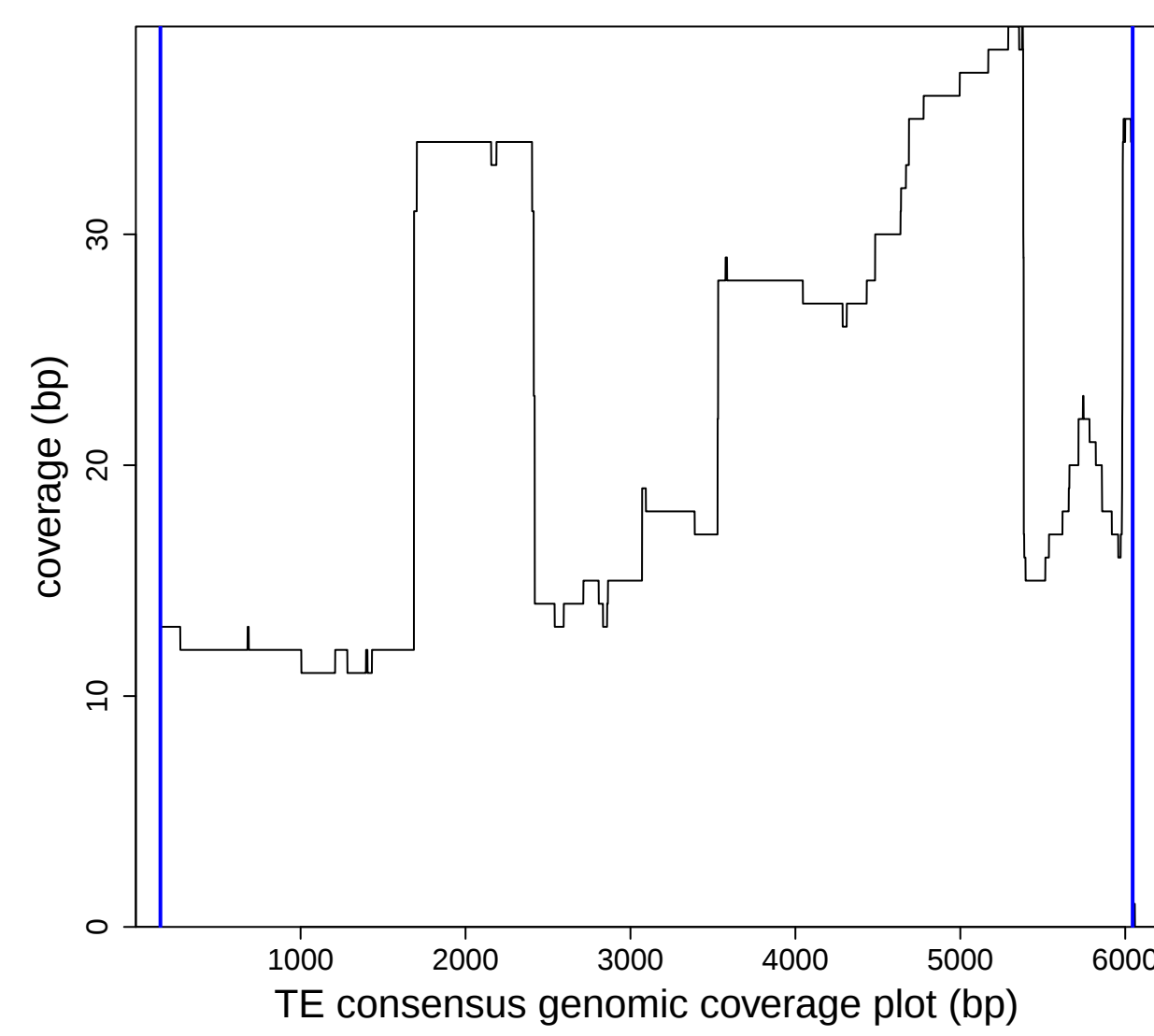
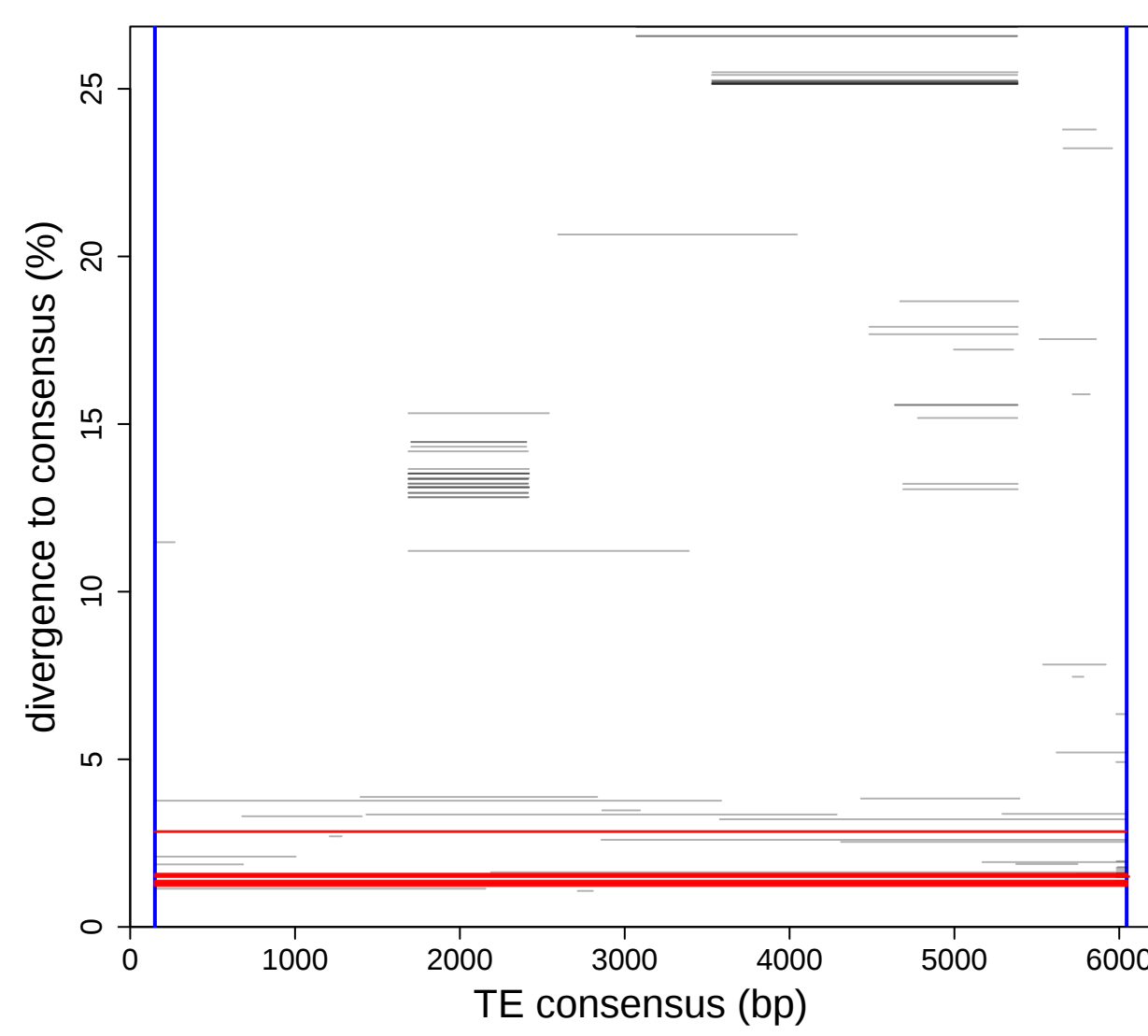
bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 5895bp; fragments: 102; full length: 8 (>=5305.5bp)

After TEtrimmer 5895 bp



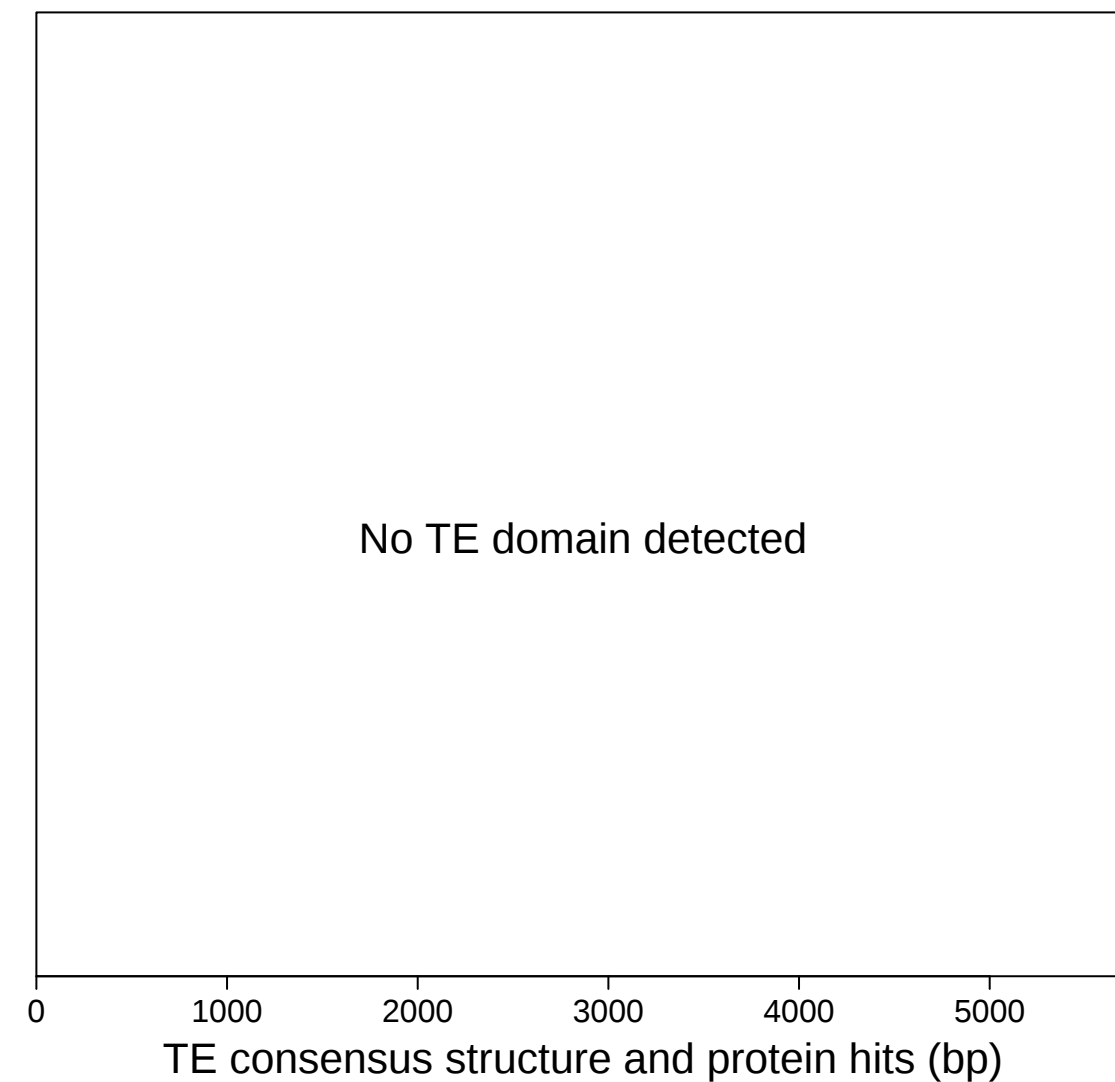
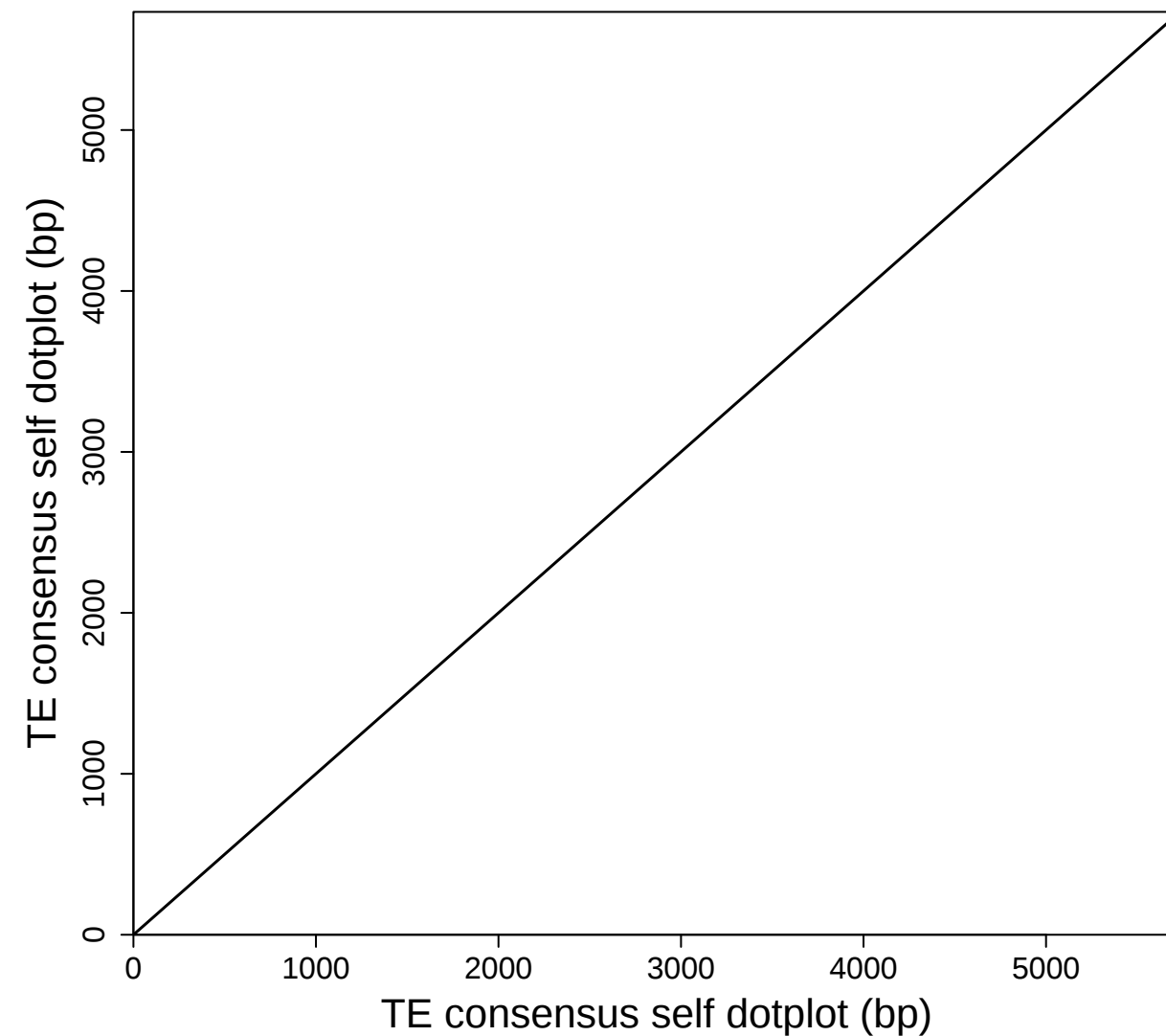
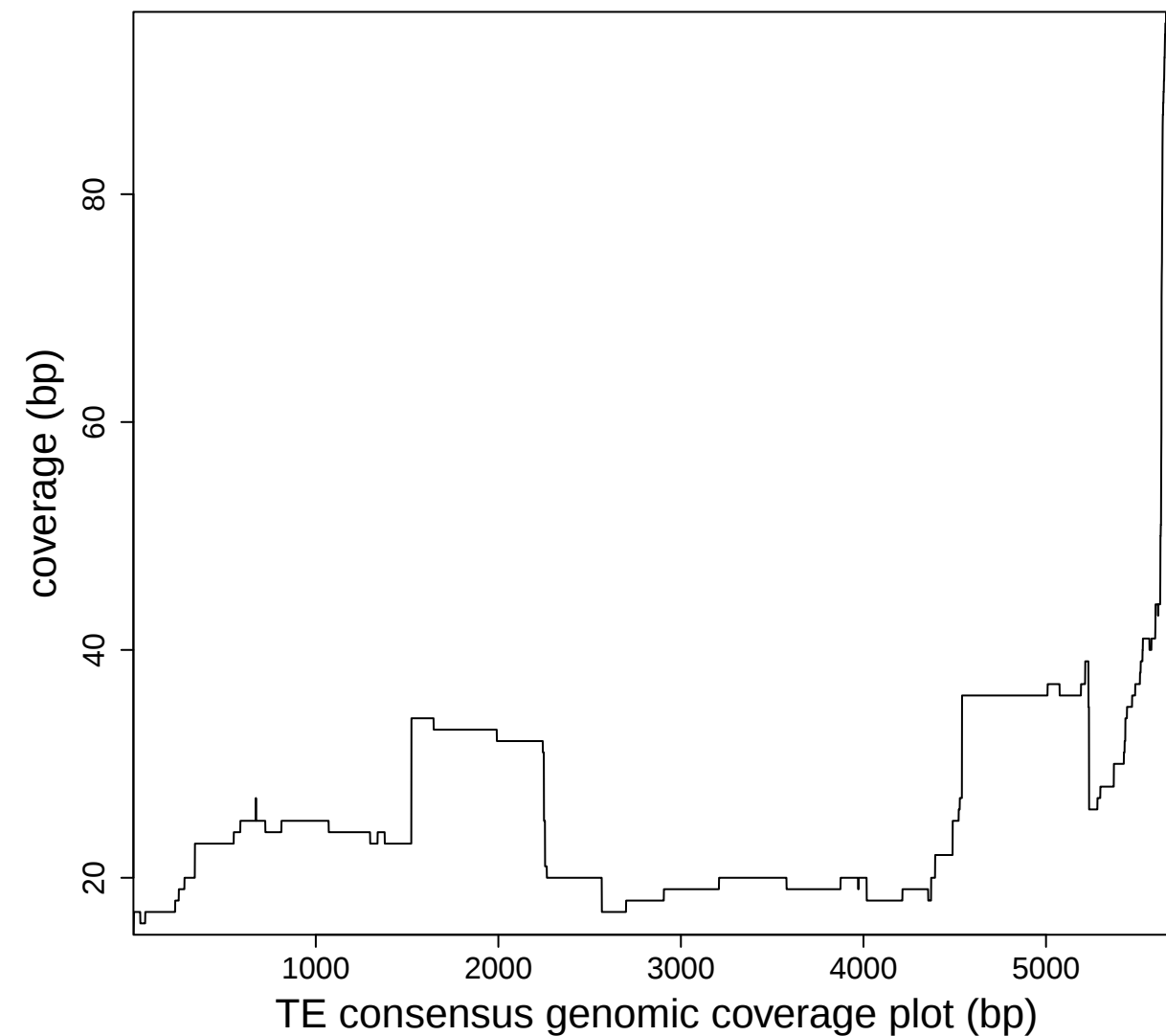
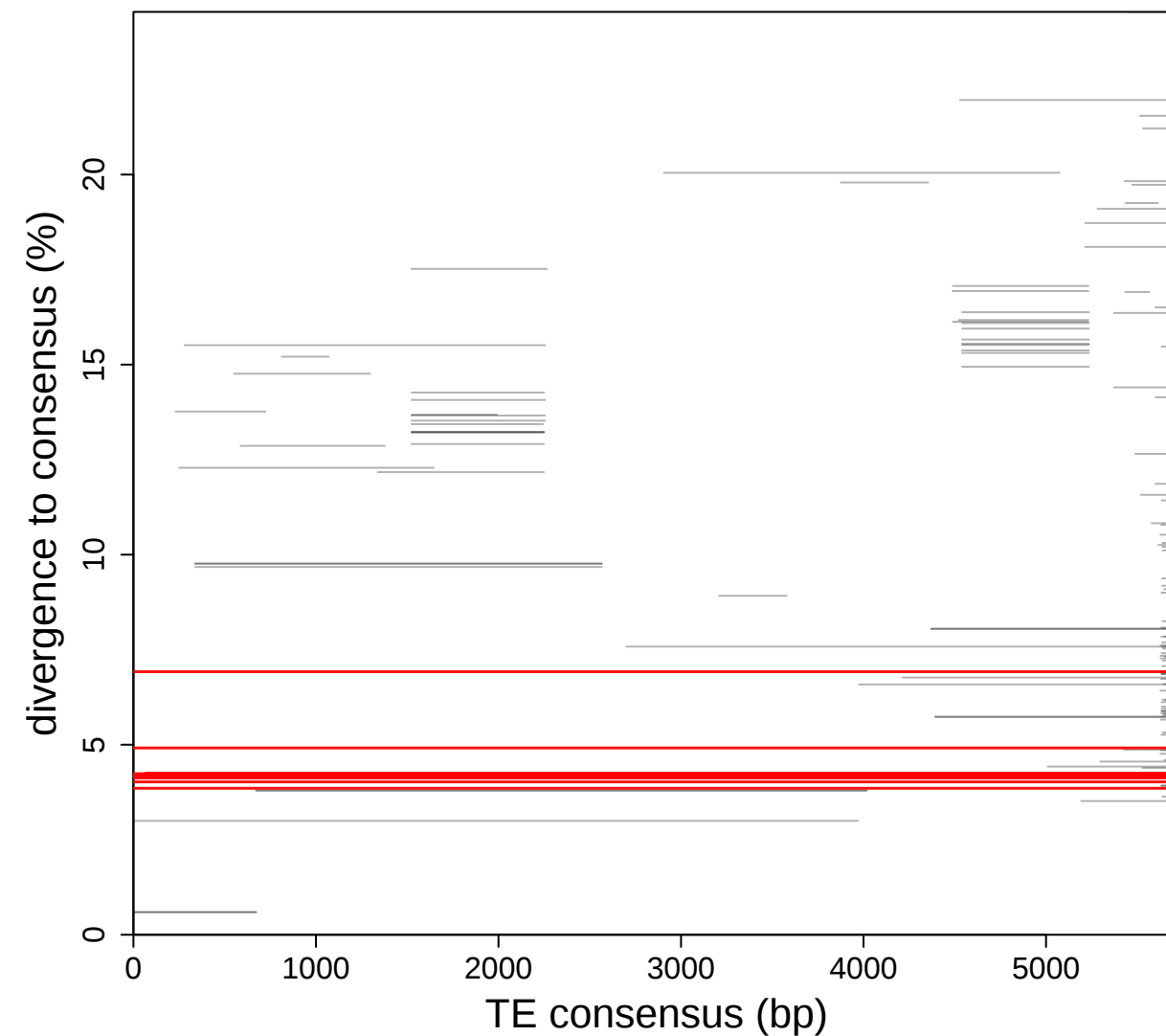
7.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 6195bp; fragments: 102; full length: 8 (>=5575.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

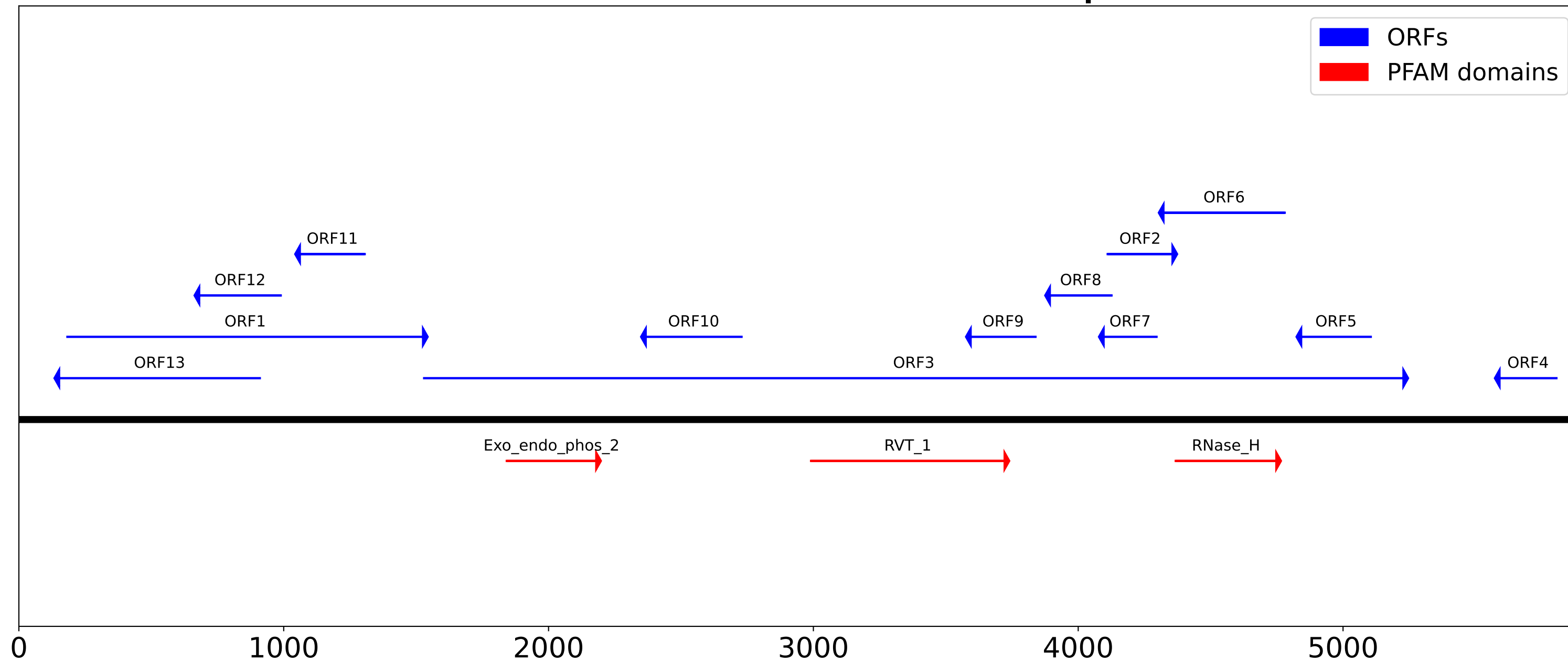


TE: rnd_4_family_257
size: 5734bp; fragments: 141; full length: 14 (≥ 5160.6 bp)

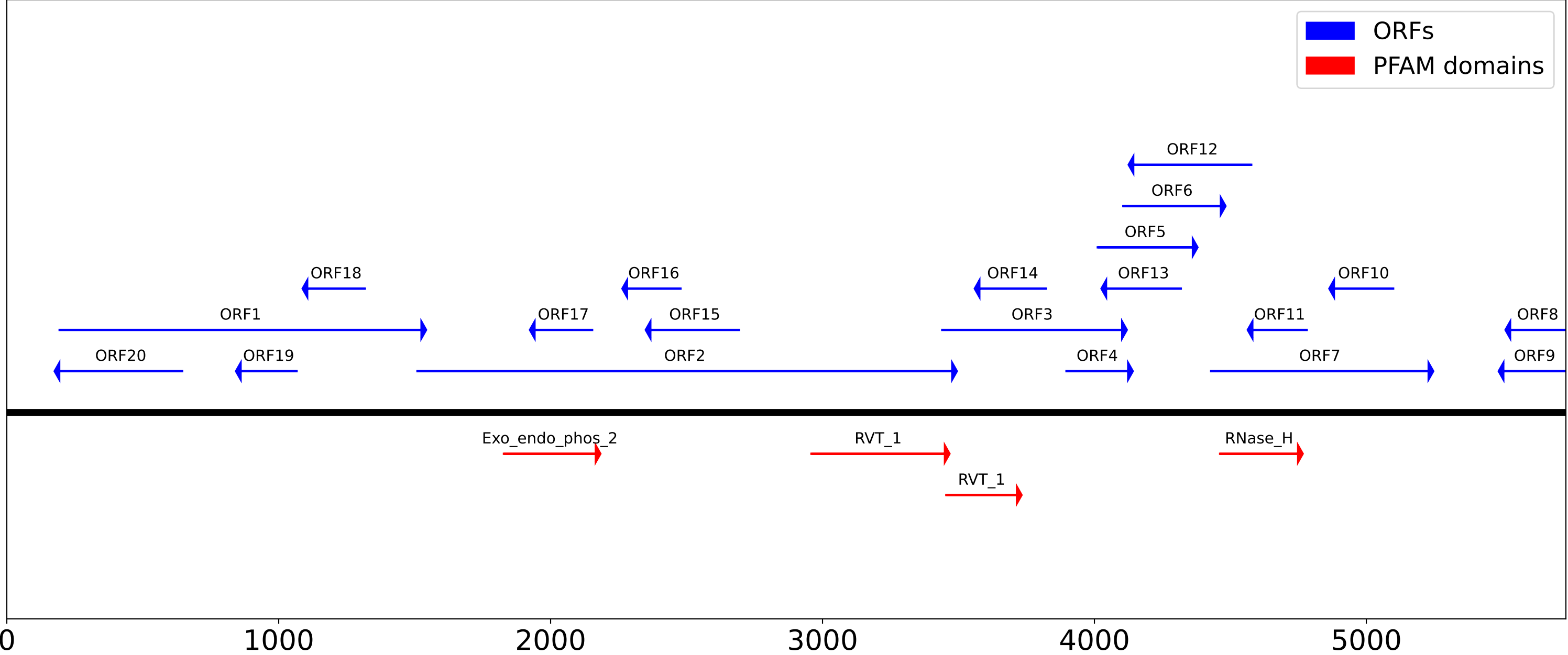
Before TEtrimmer 5734 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

