

Start crop Point



End crop Point



MSA length = 5062



Start crop Point

End crop Point

0

10

0

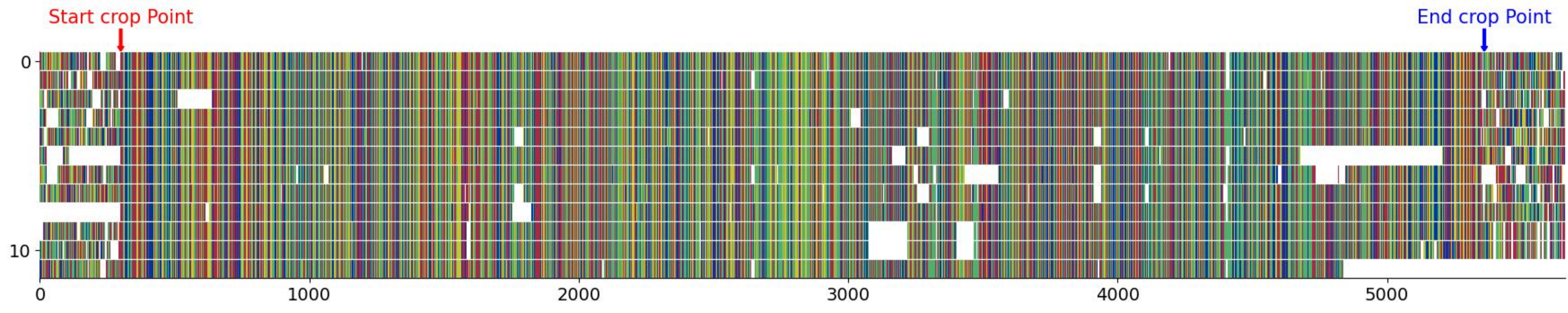
1000

2000

3000

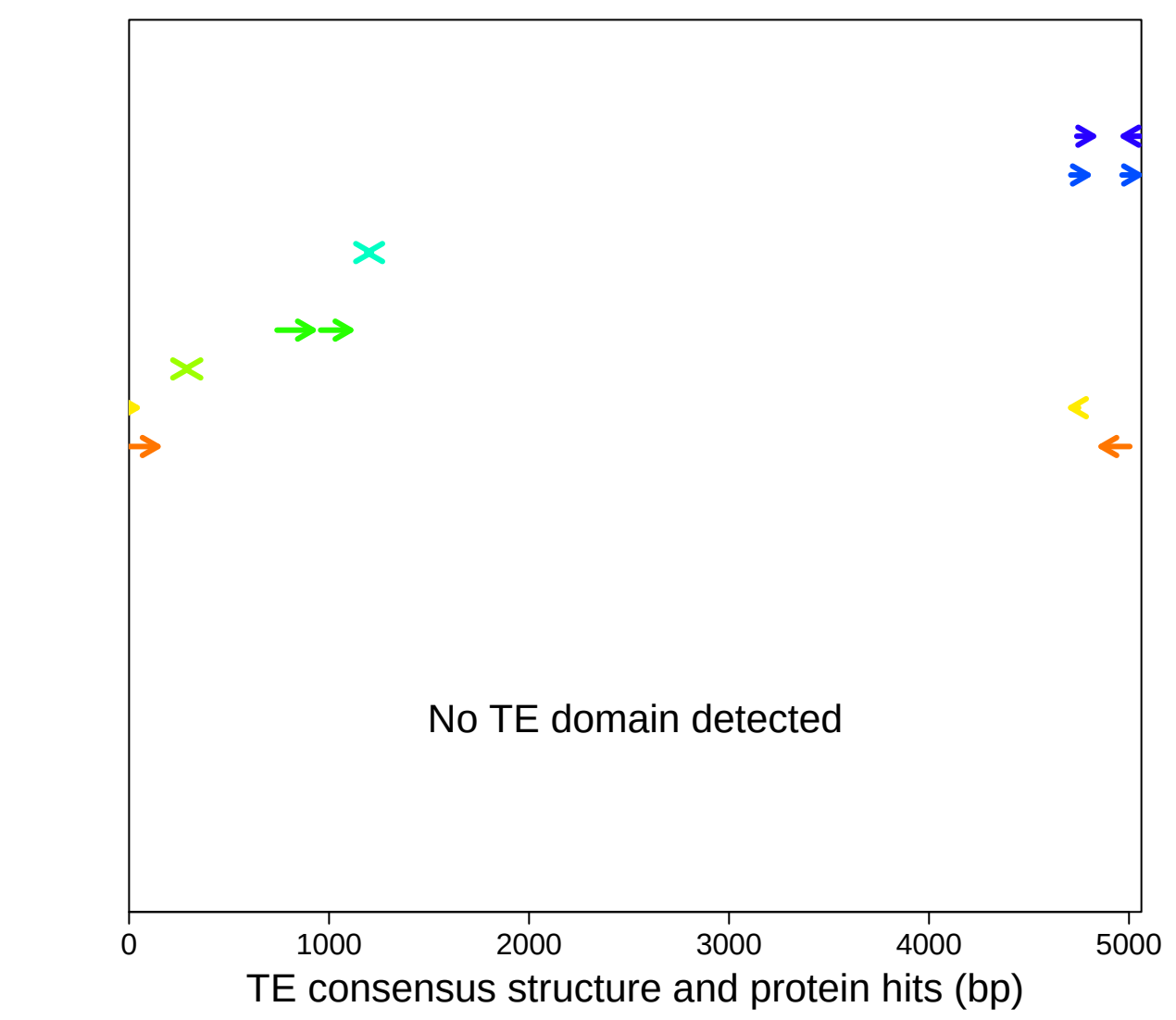
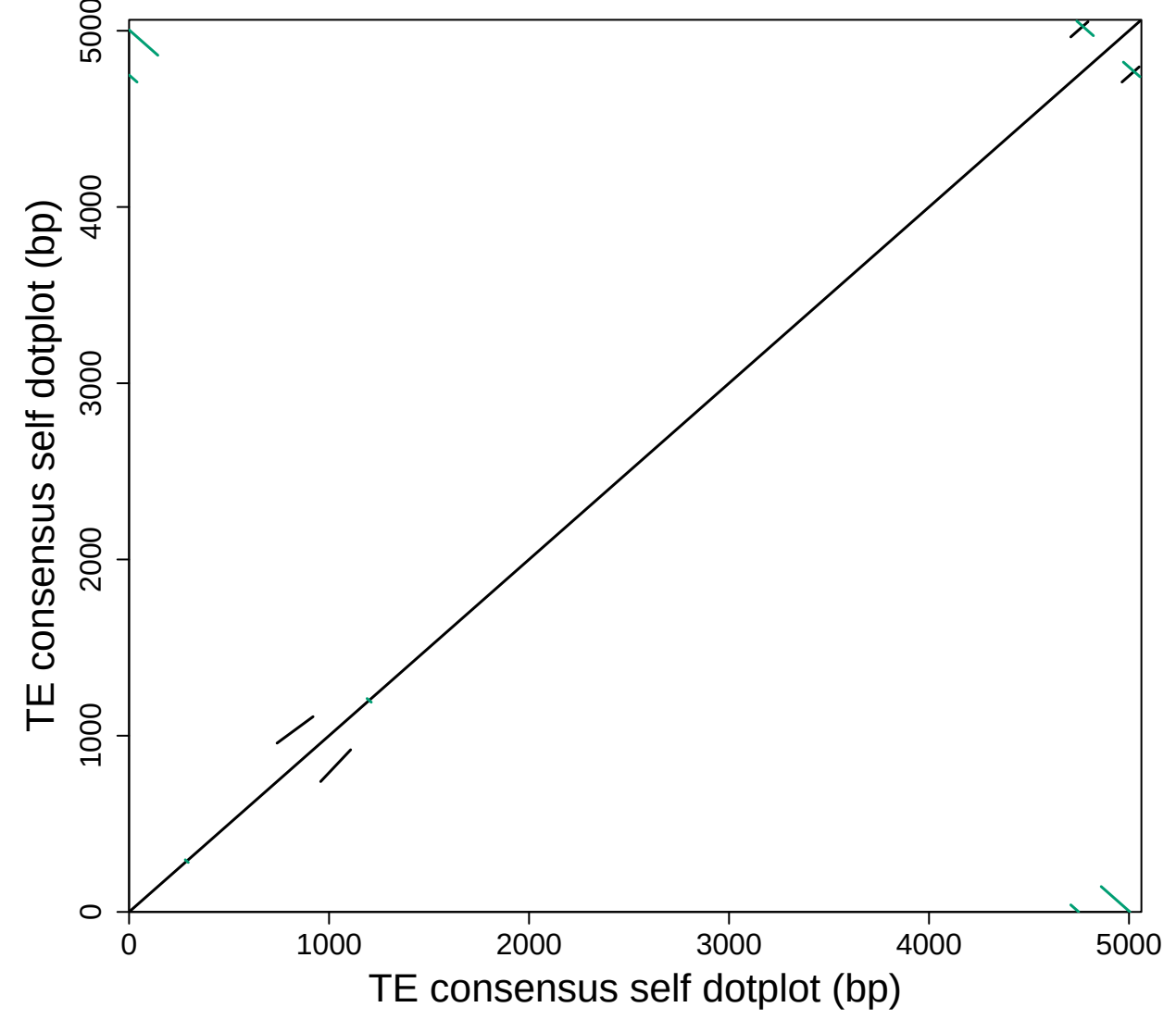
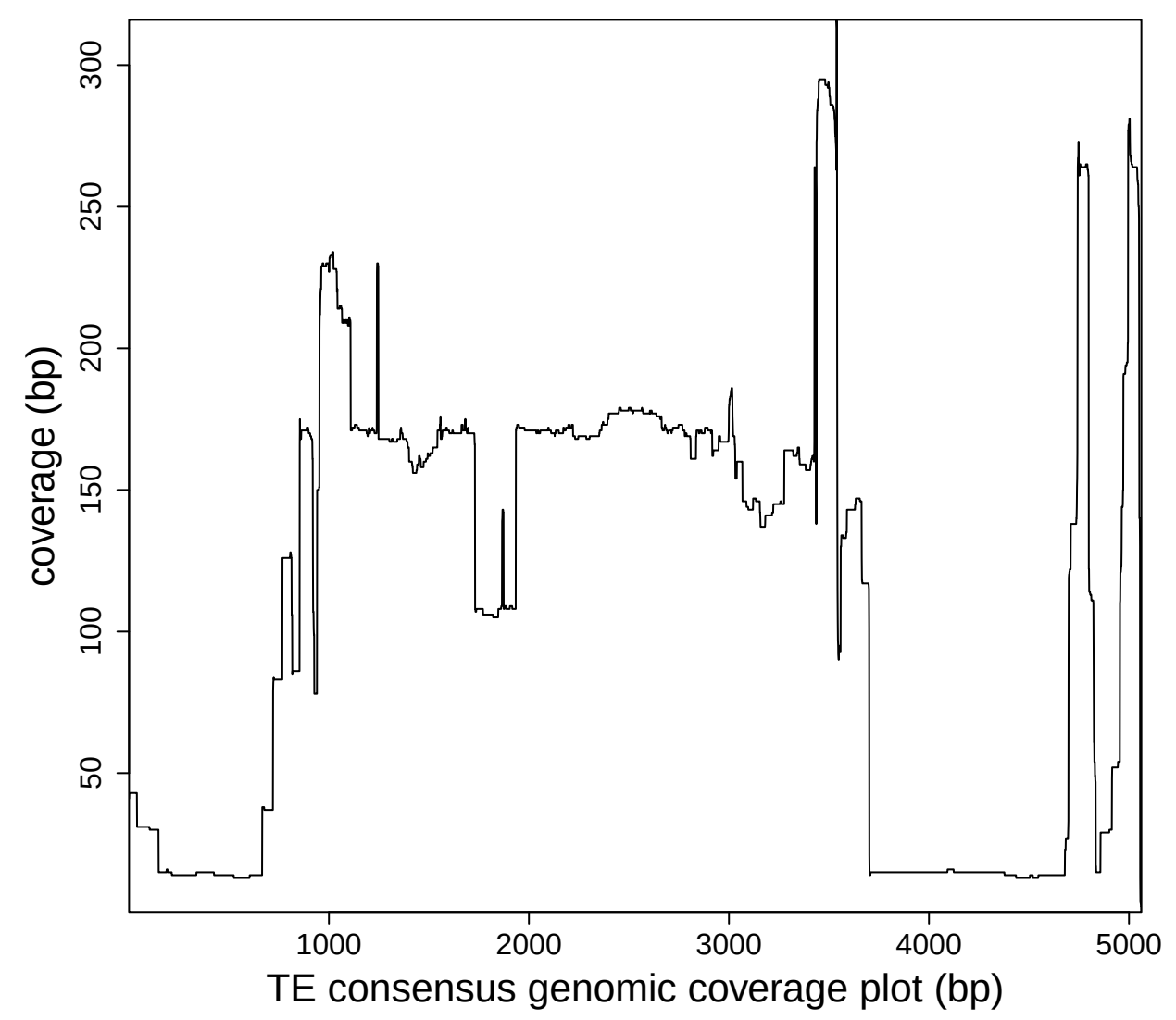
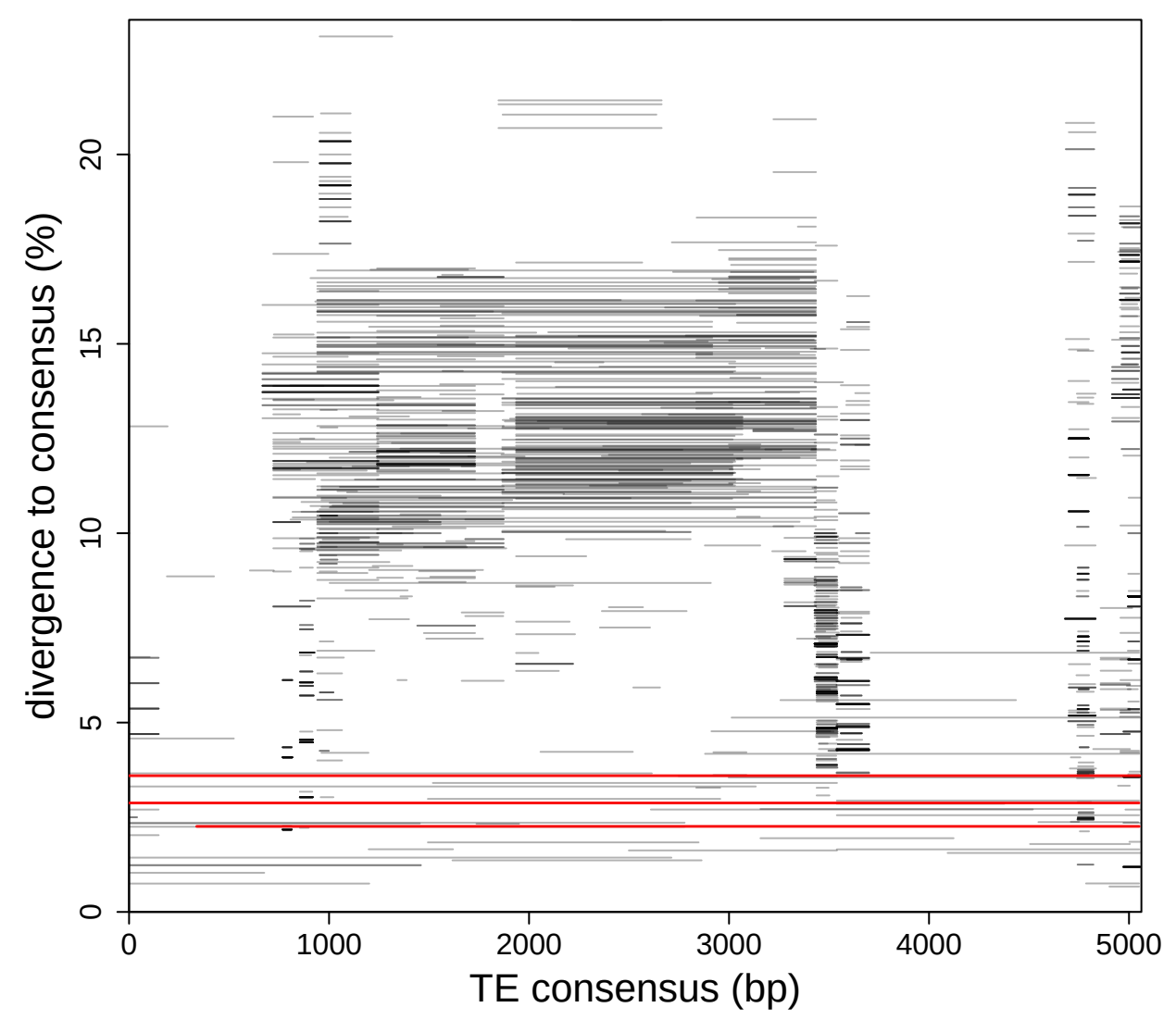
4000

5000



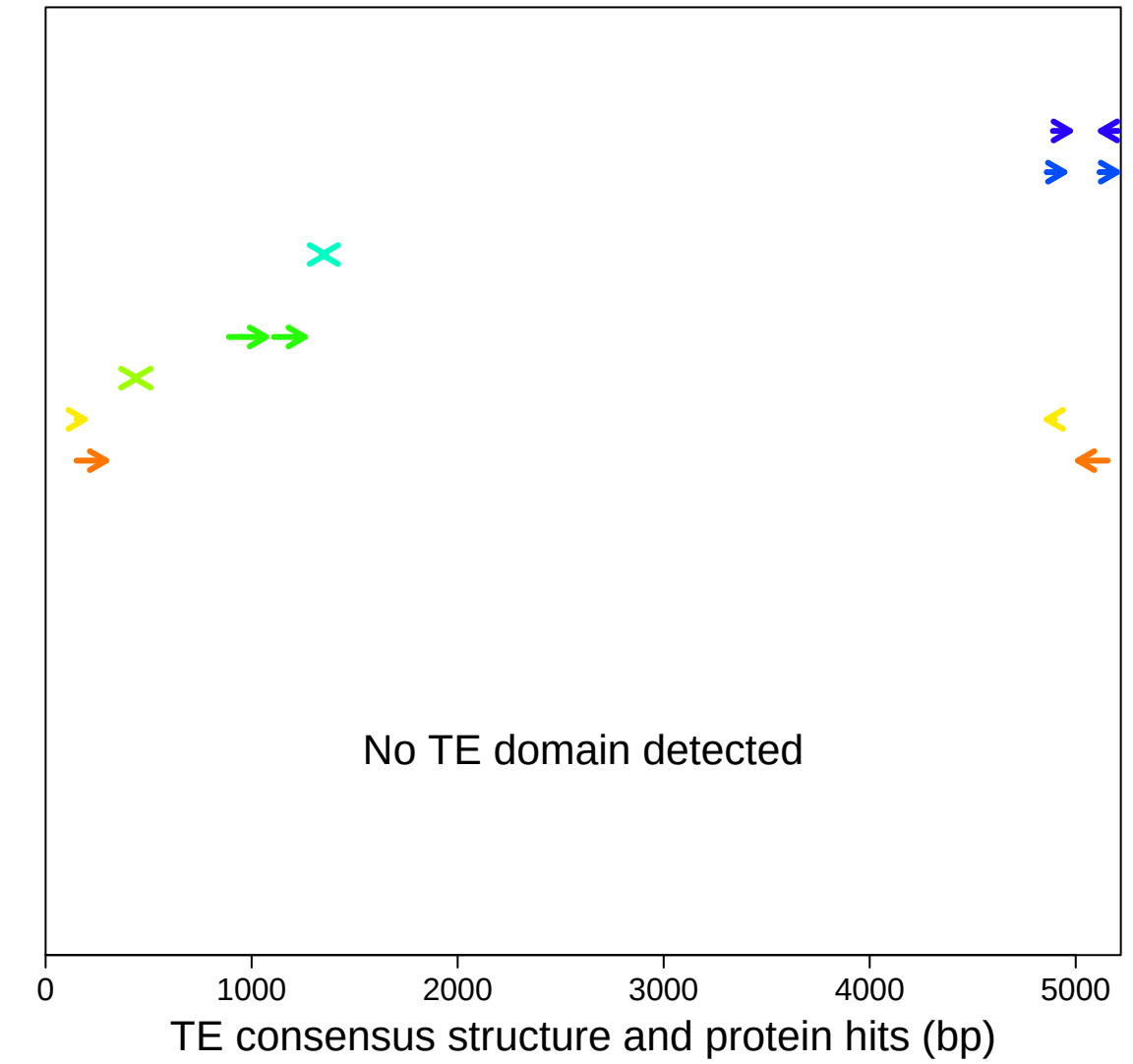
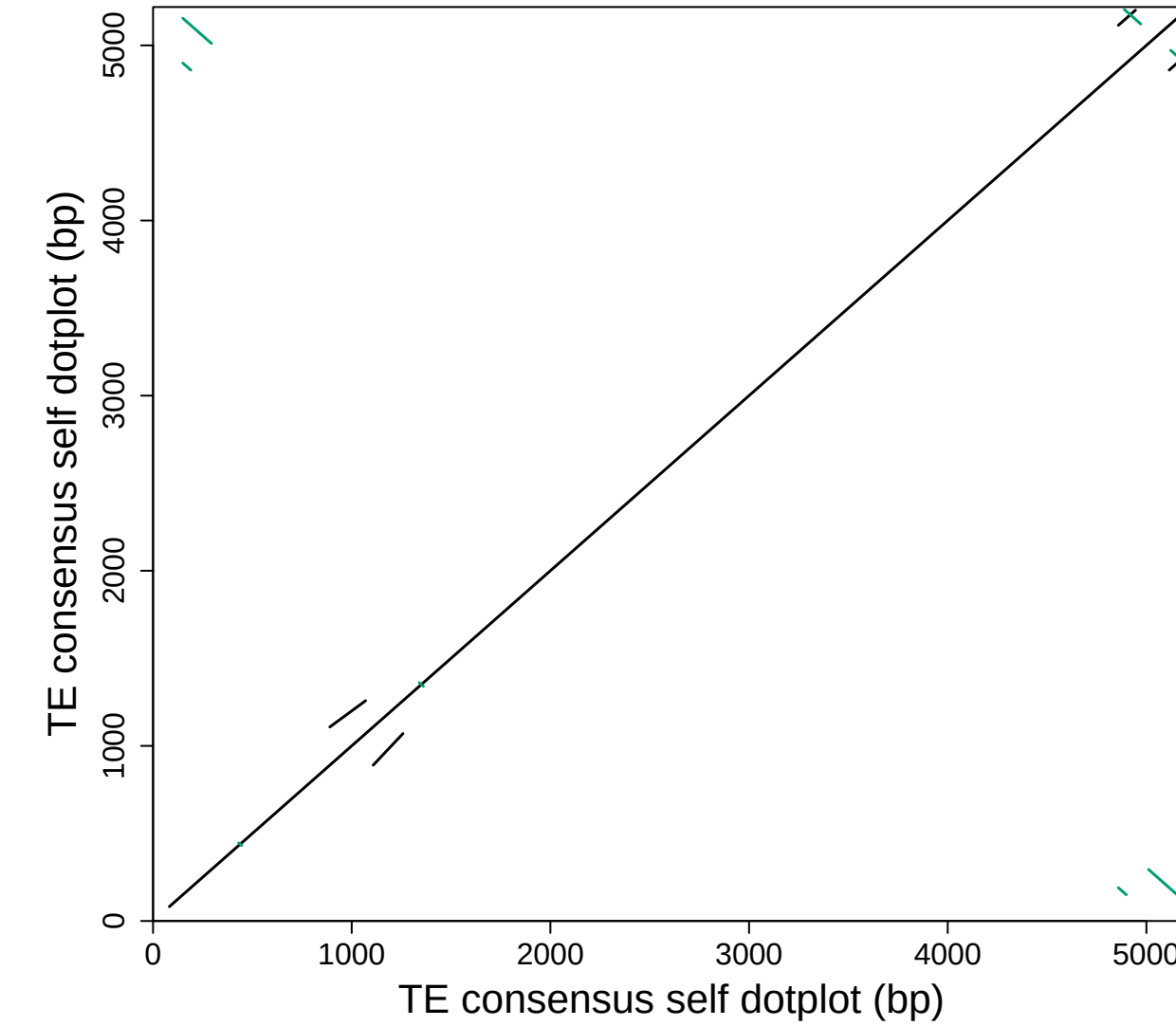
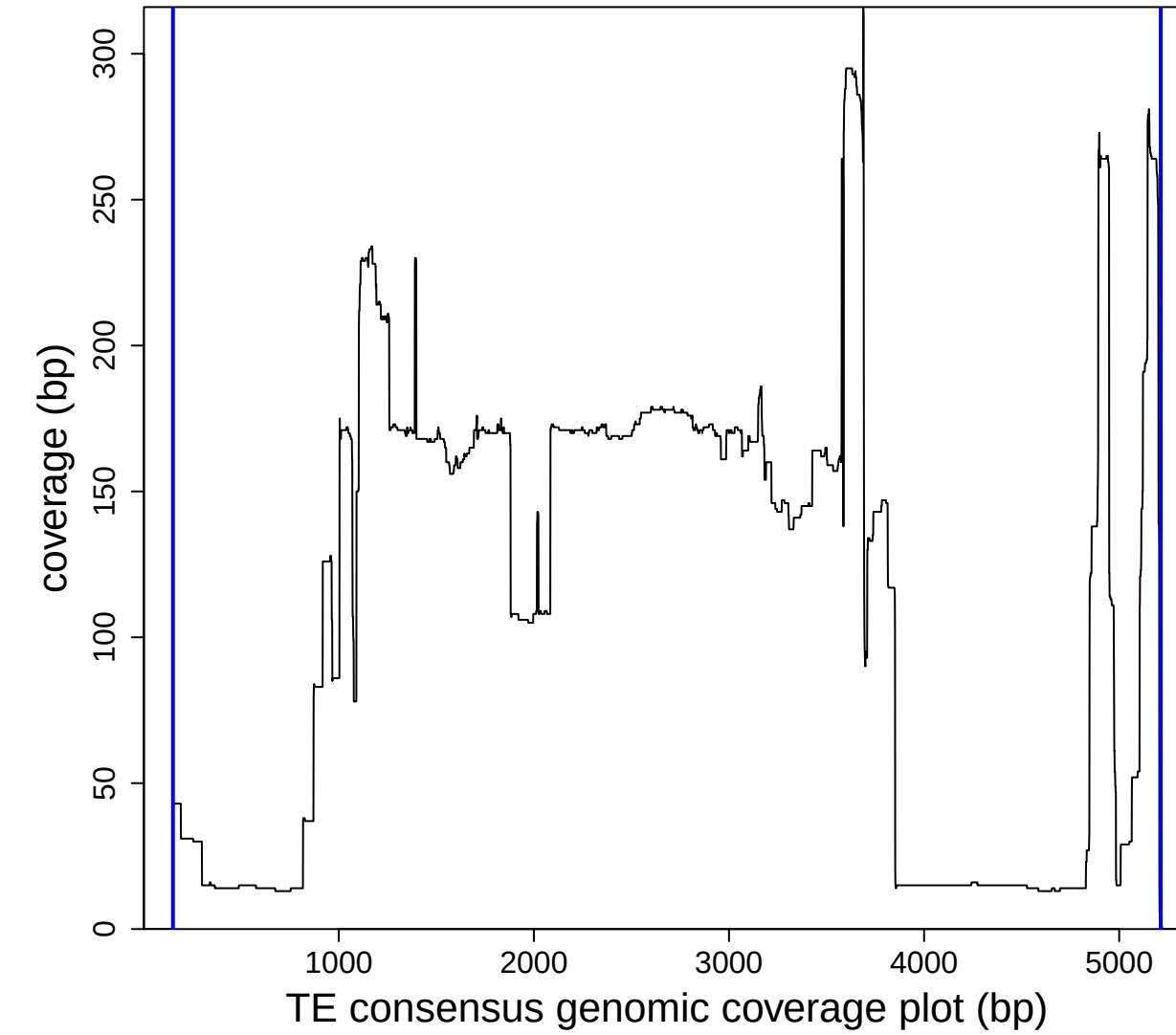
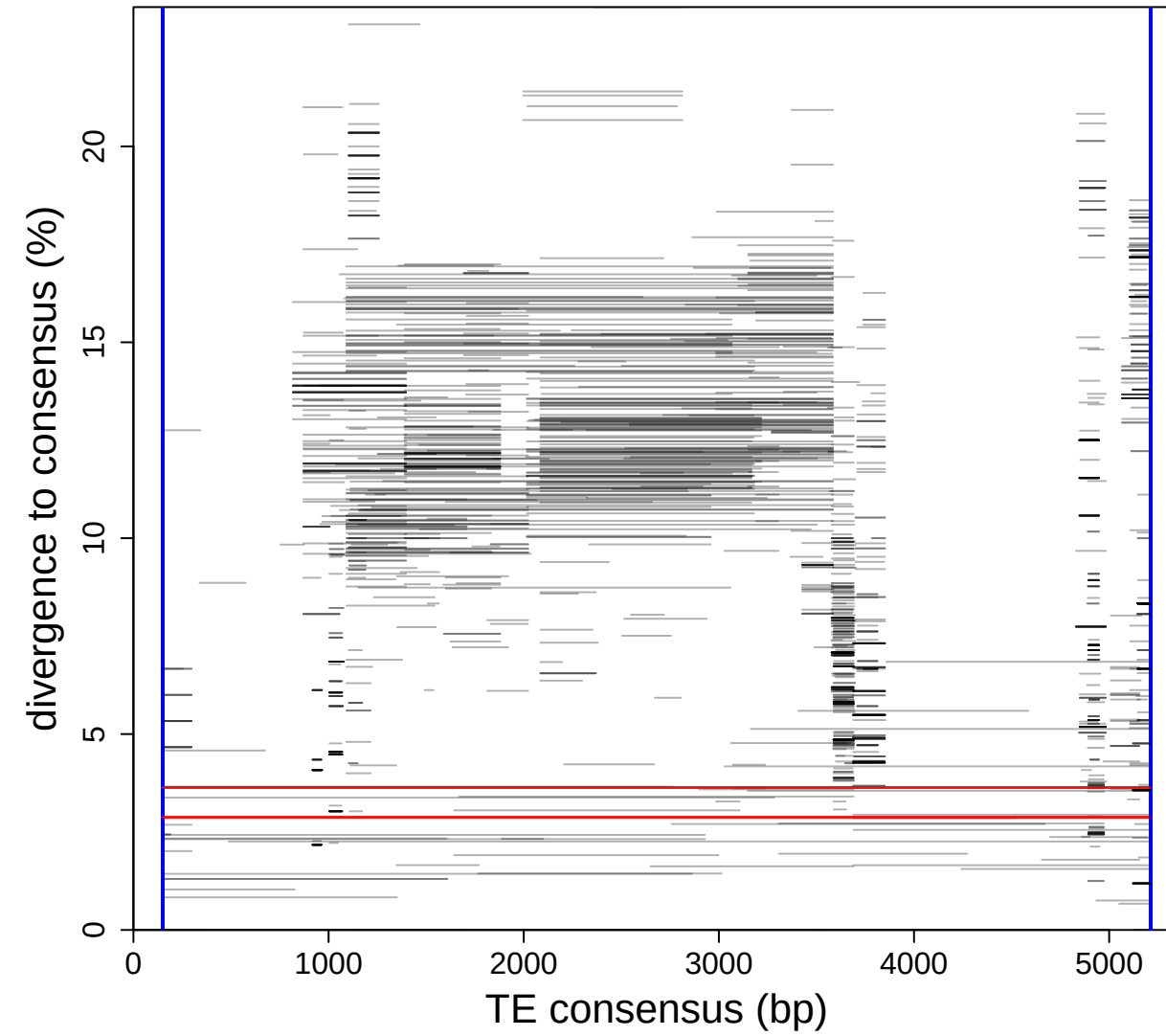
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 5062bp; fragments: 1727; full length: 3 (>=4555.8bp)

After TEtrimmer 5062 bp



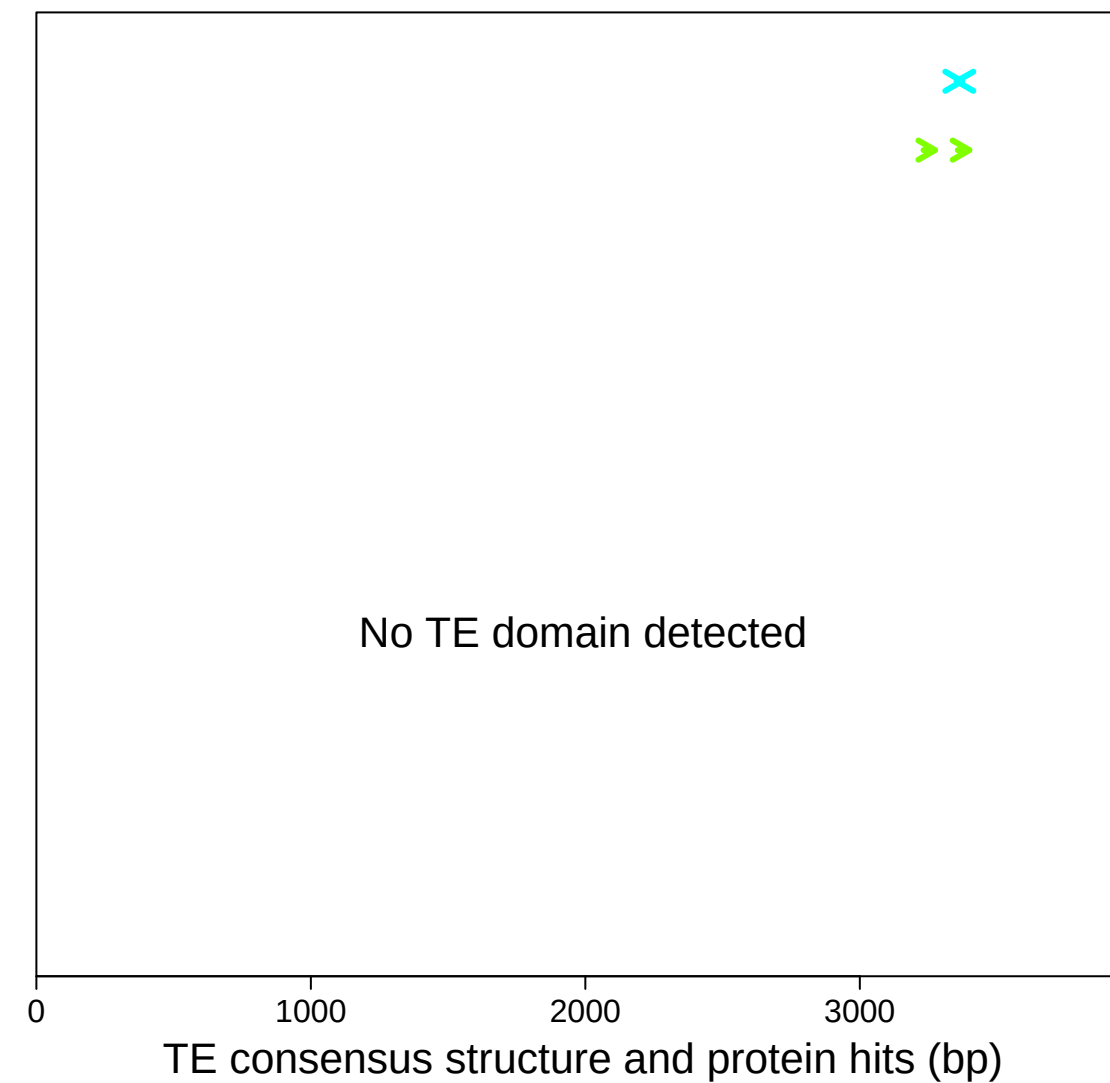
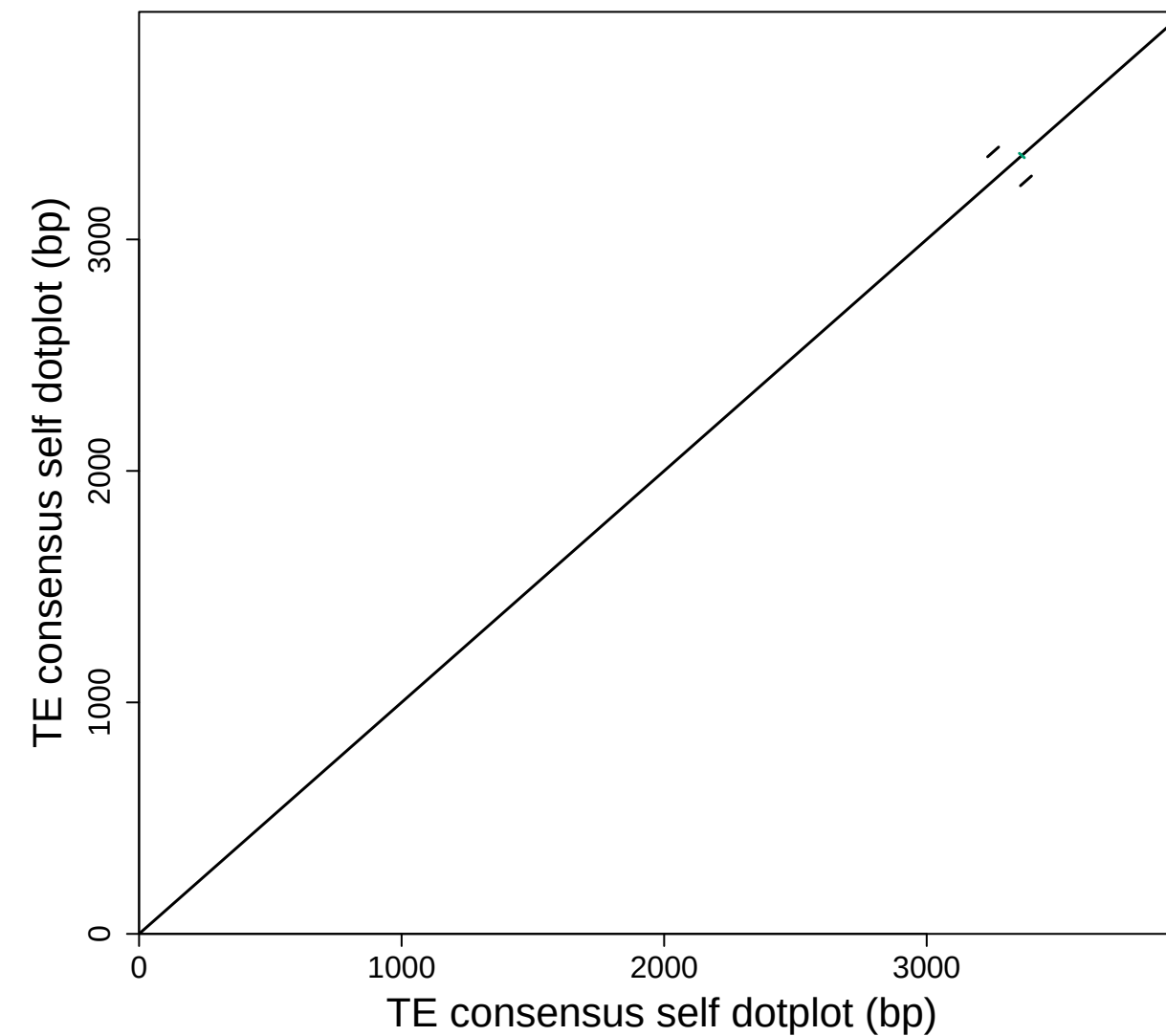
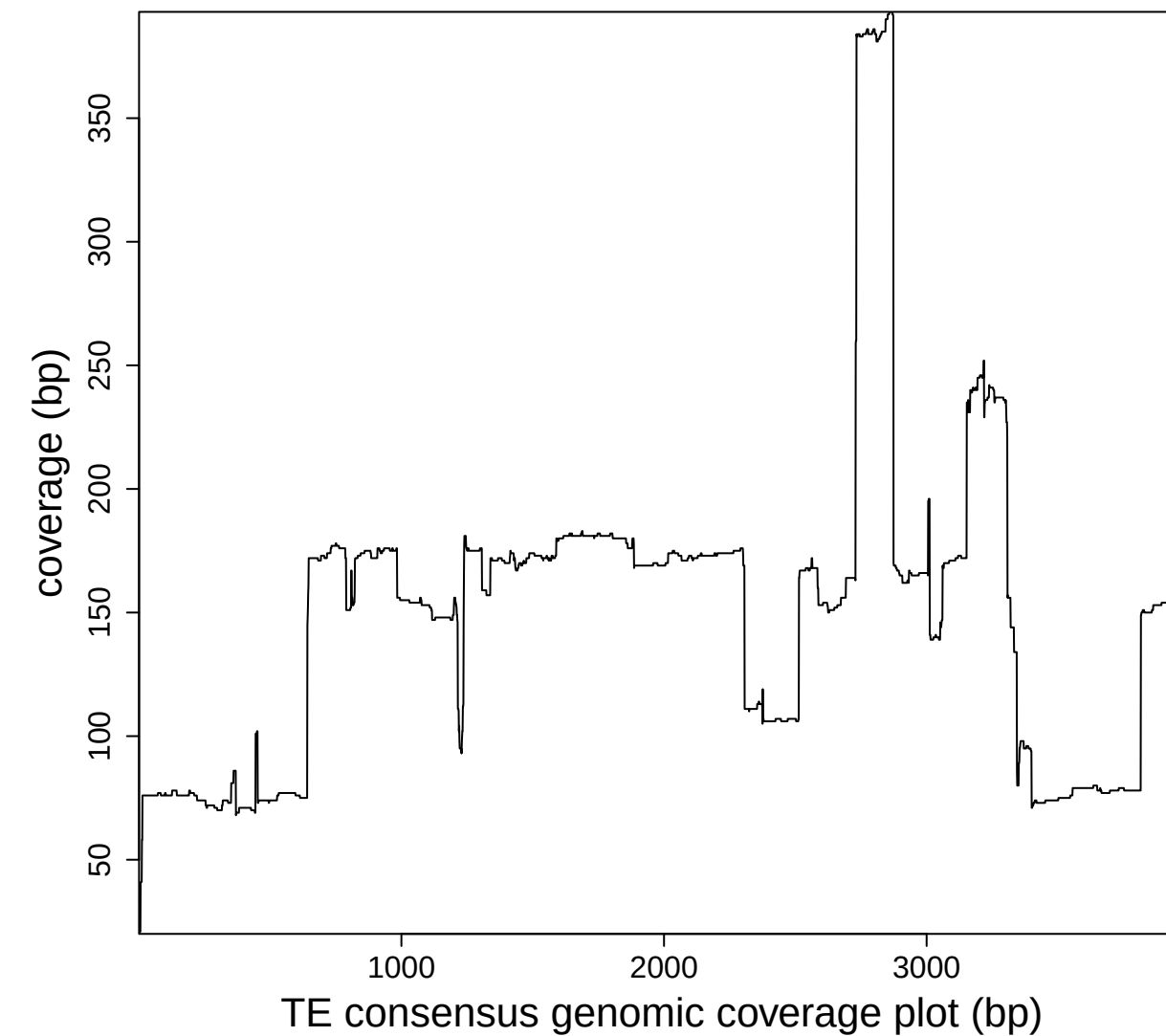
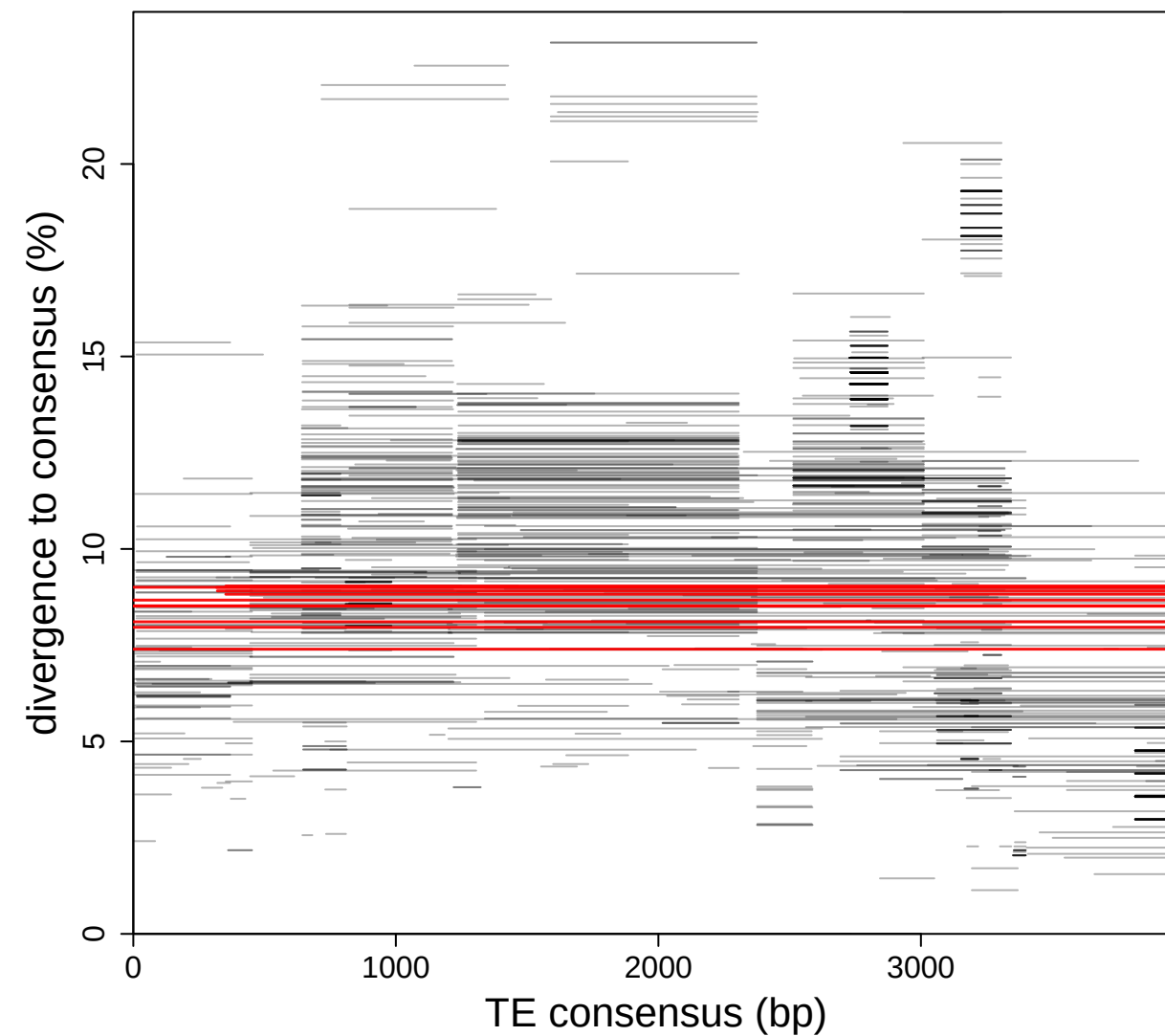
asta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_c
size: 5363bp; fragments: 1727; full length: 2 (>=4826.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

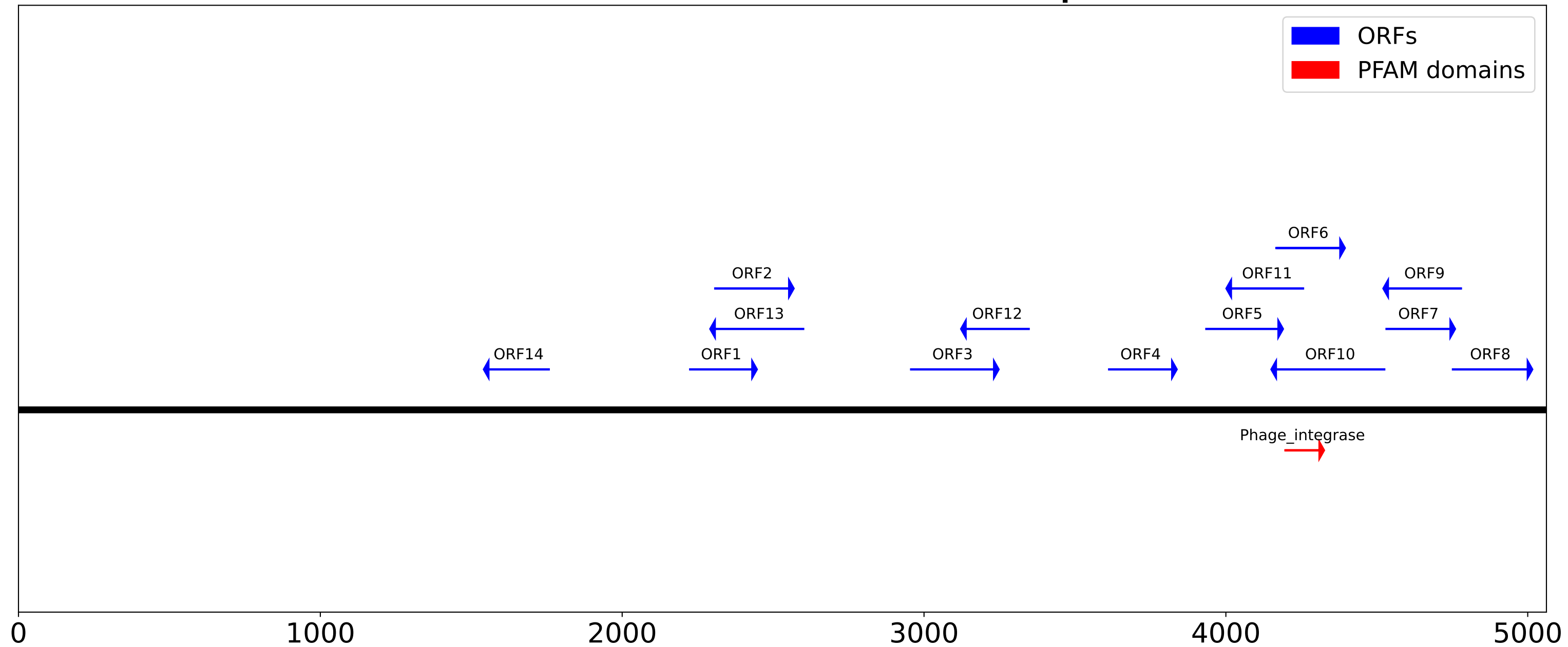


TE: rnd_5_family_1152
size: 3983bp; fragments: 1199; full length: 11 (≥ 3584.7 bp)

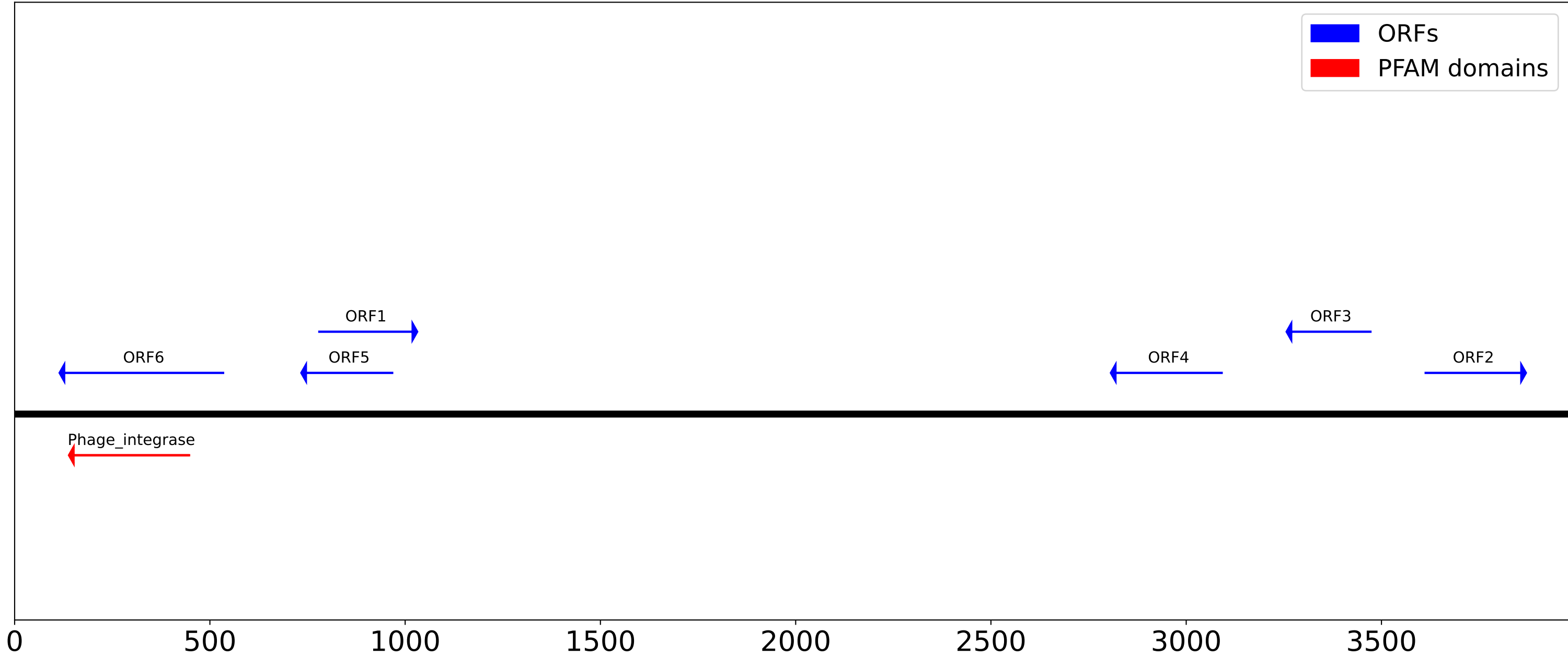
Before TEtrimmer 3983 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

