

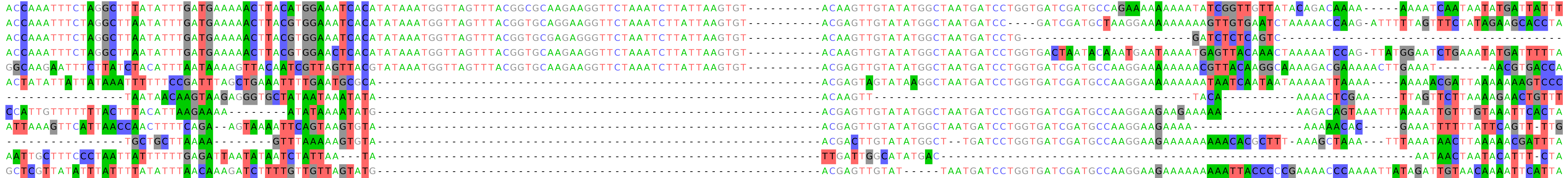
Start crop Point



End crop Point



MSA length = 1626



Start crop Point

End crop Point

0 -

10 -

0

250

500

750

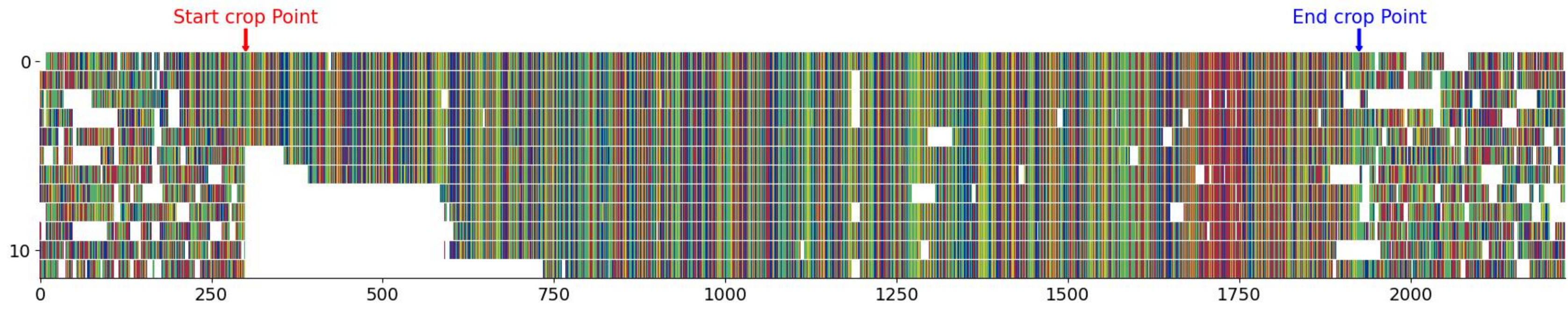
1000

1250

1500

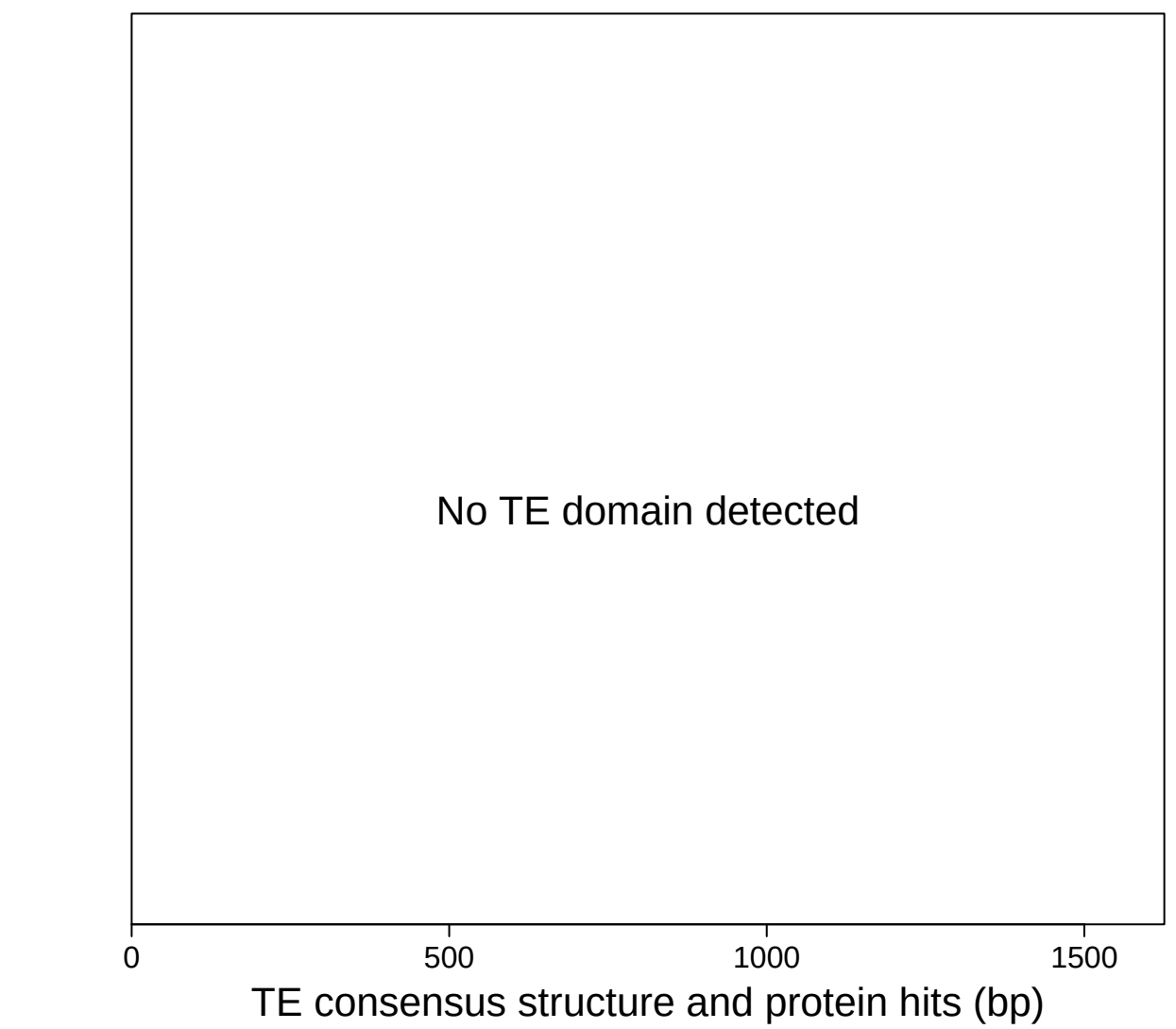
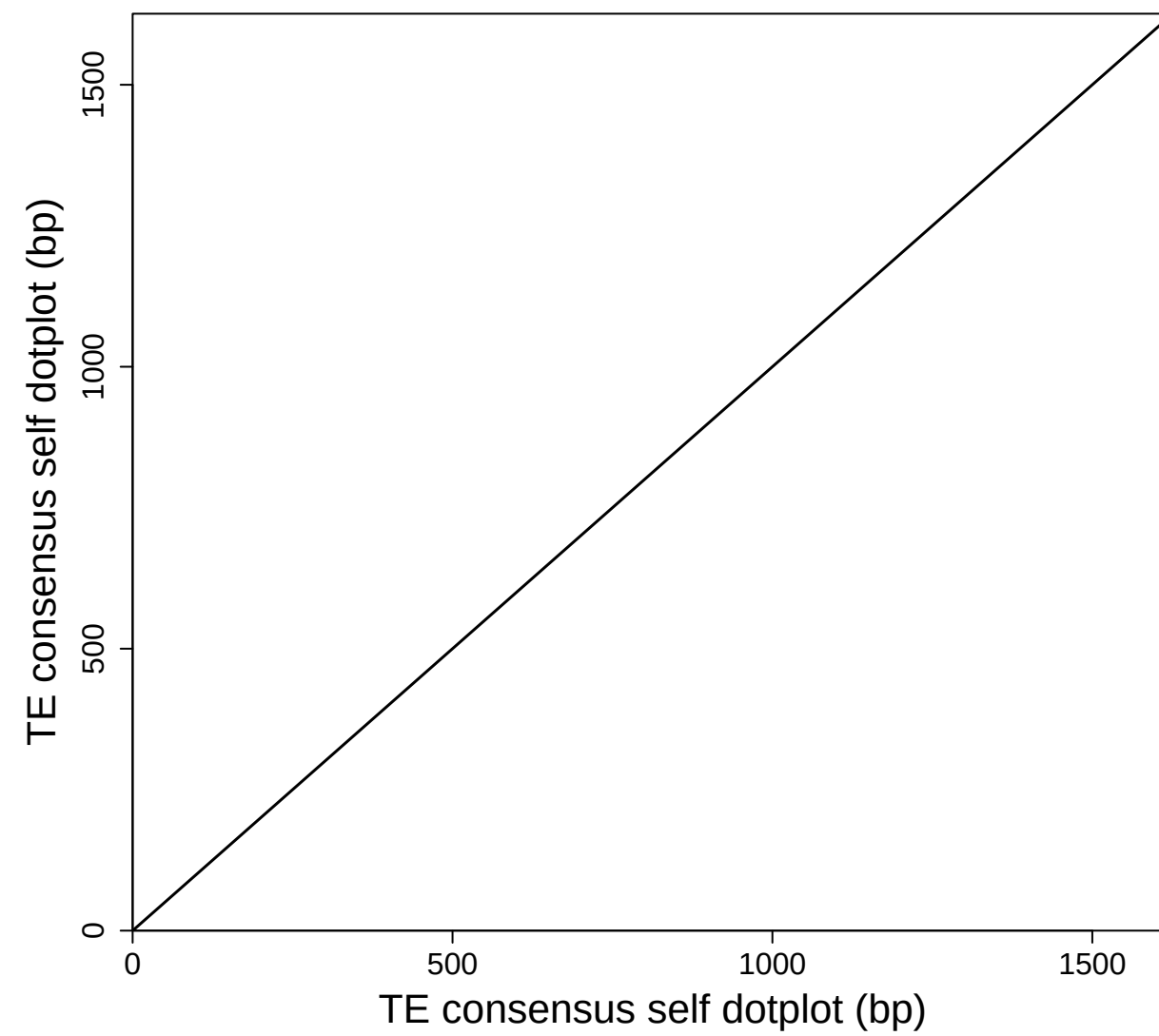
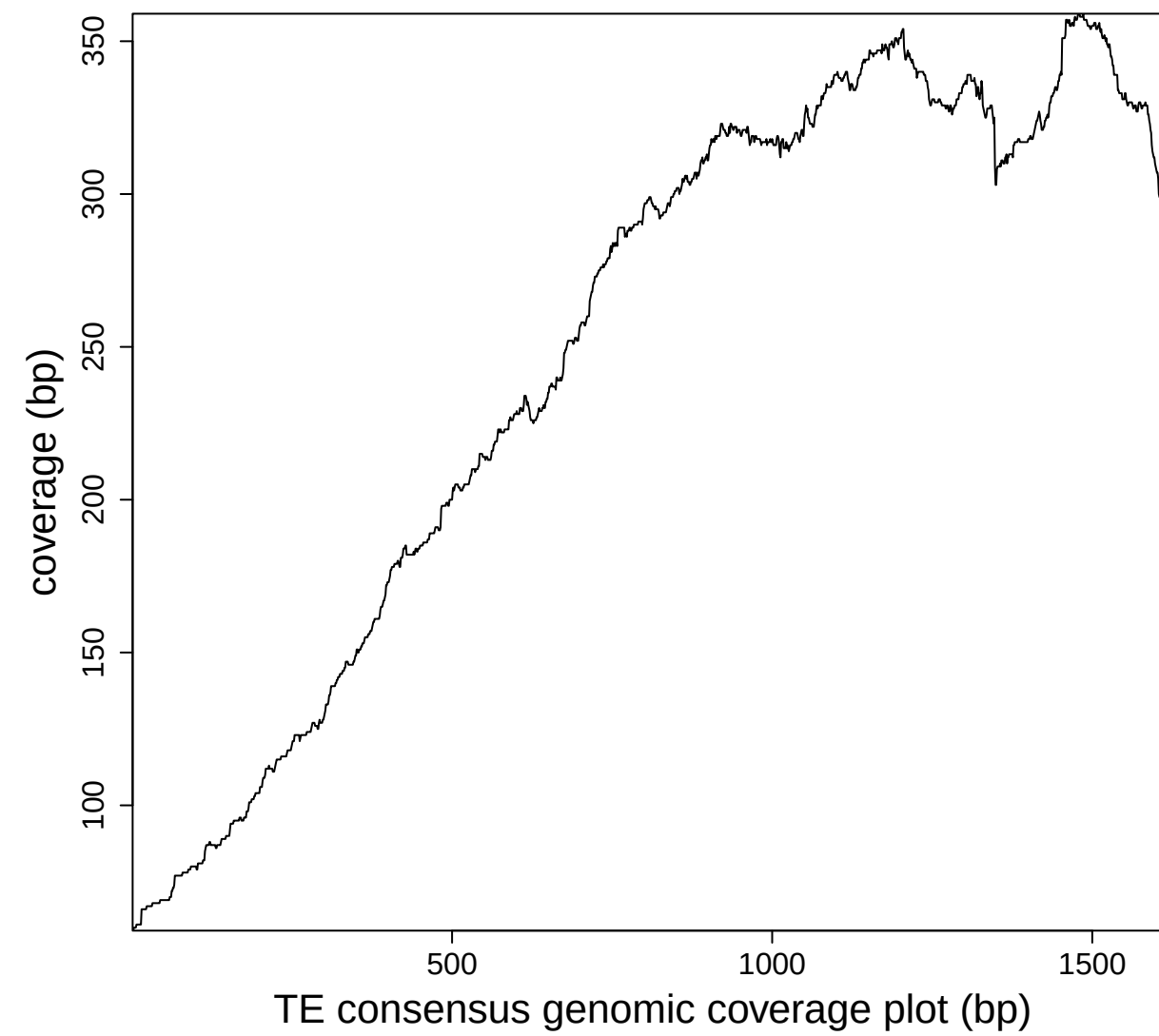
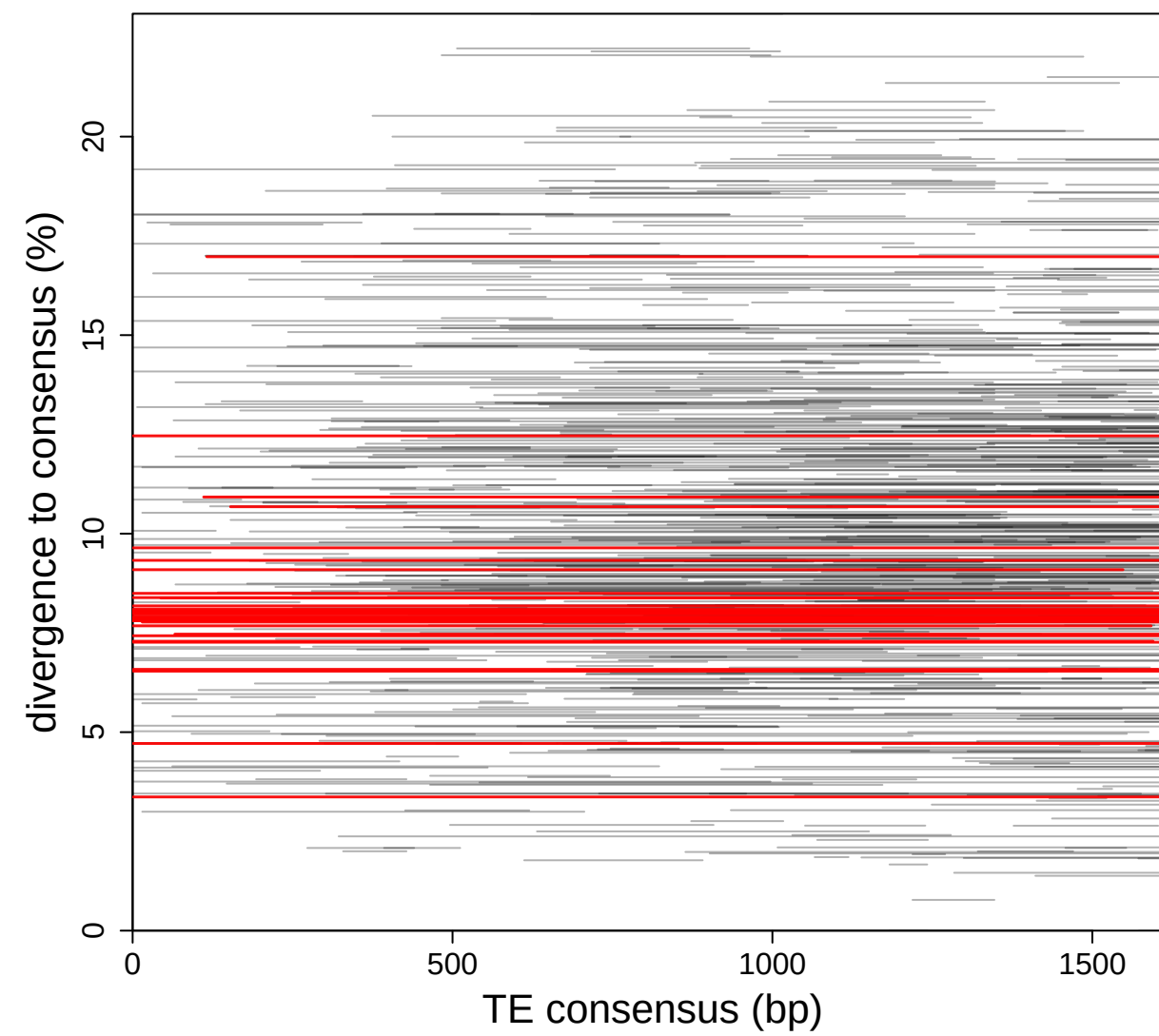
1750

2000



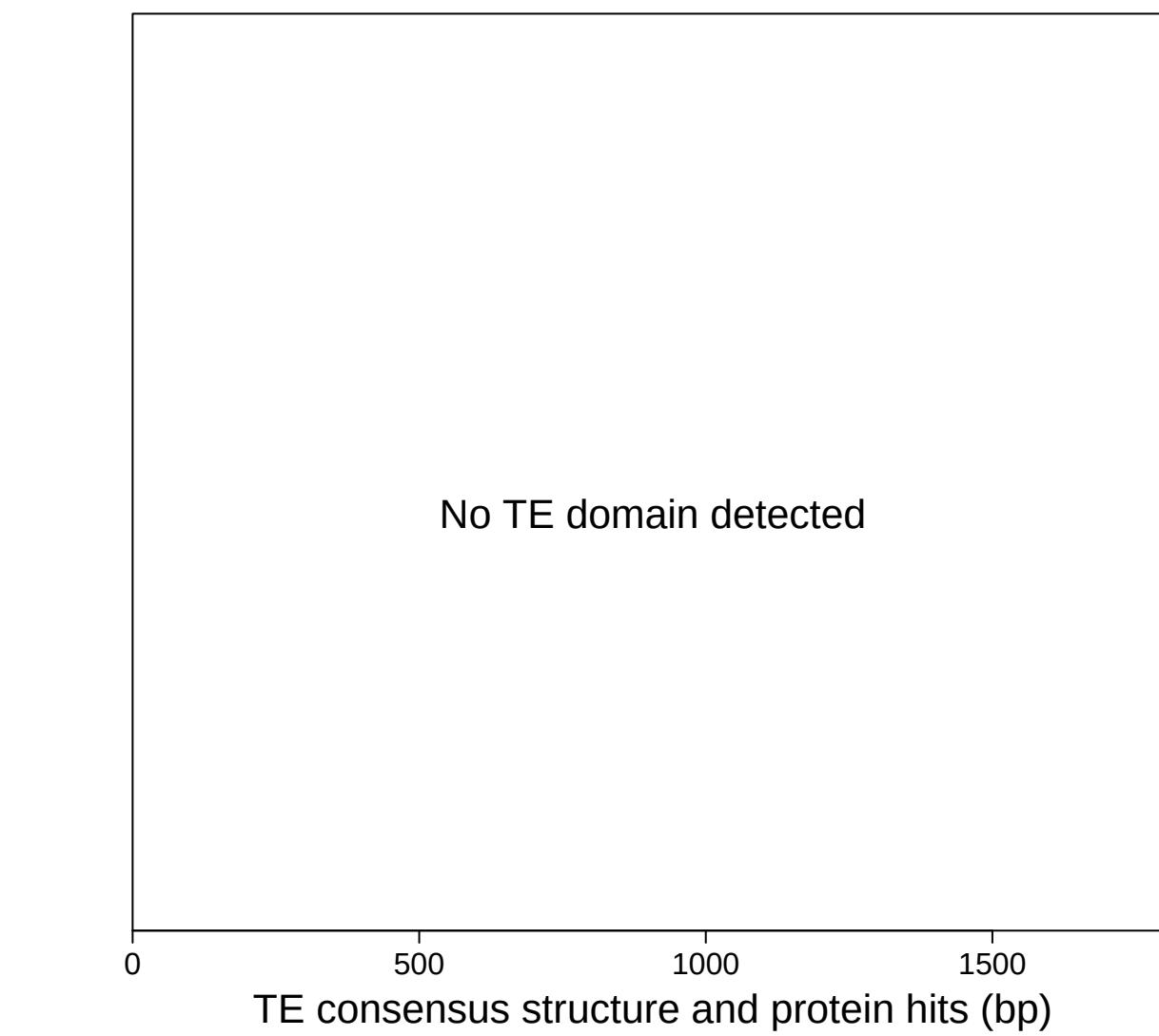
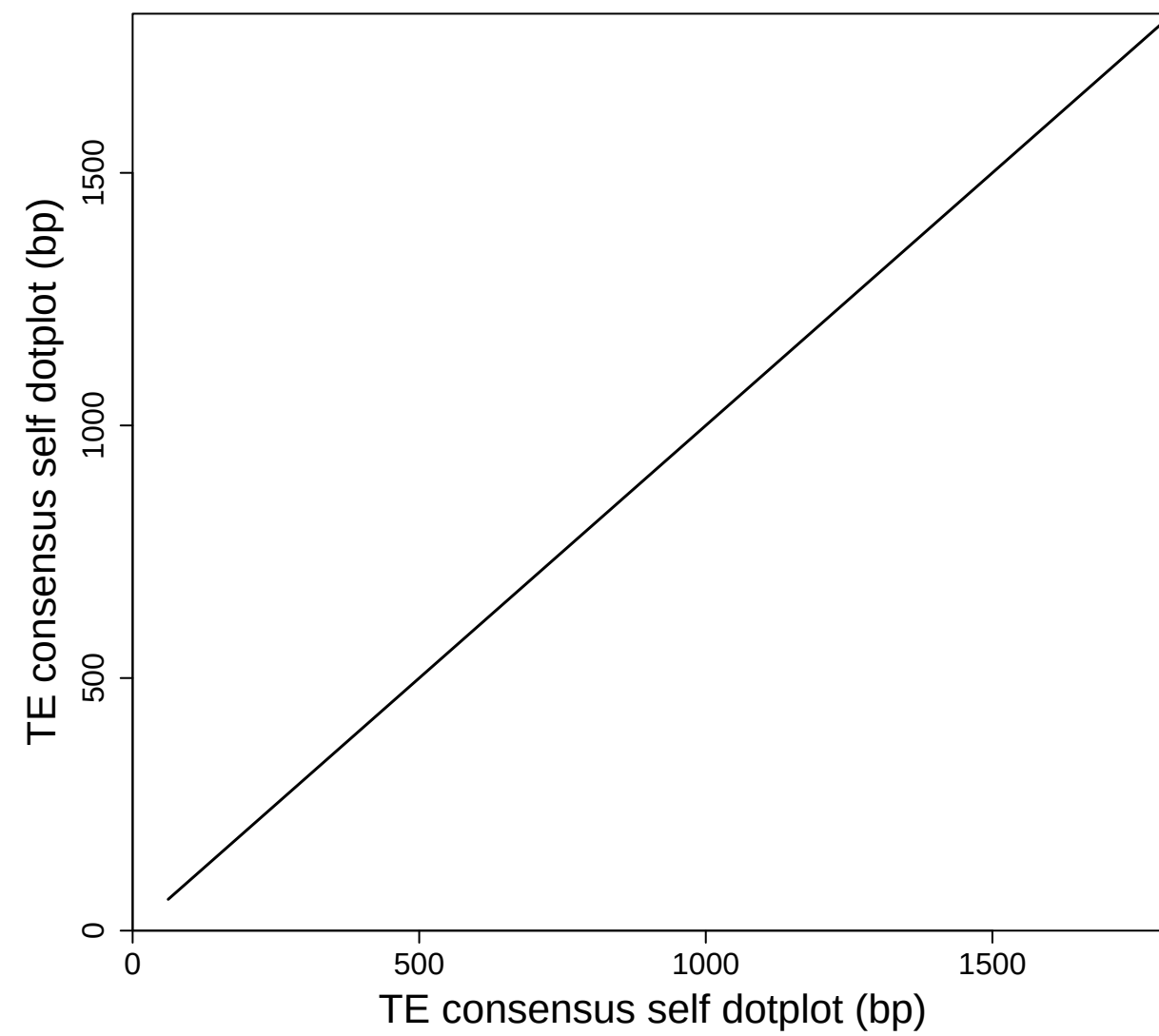
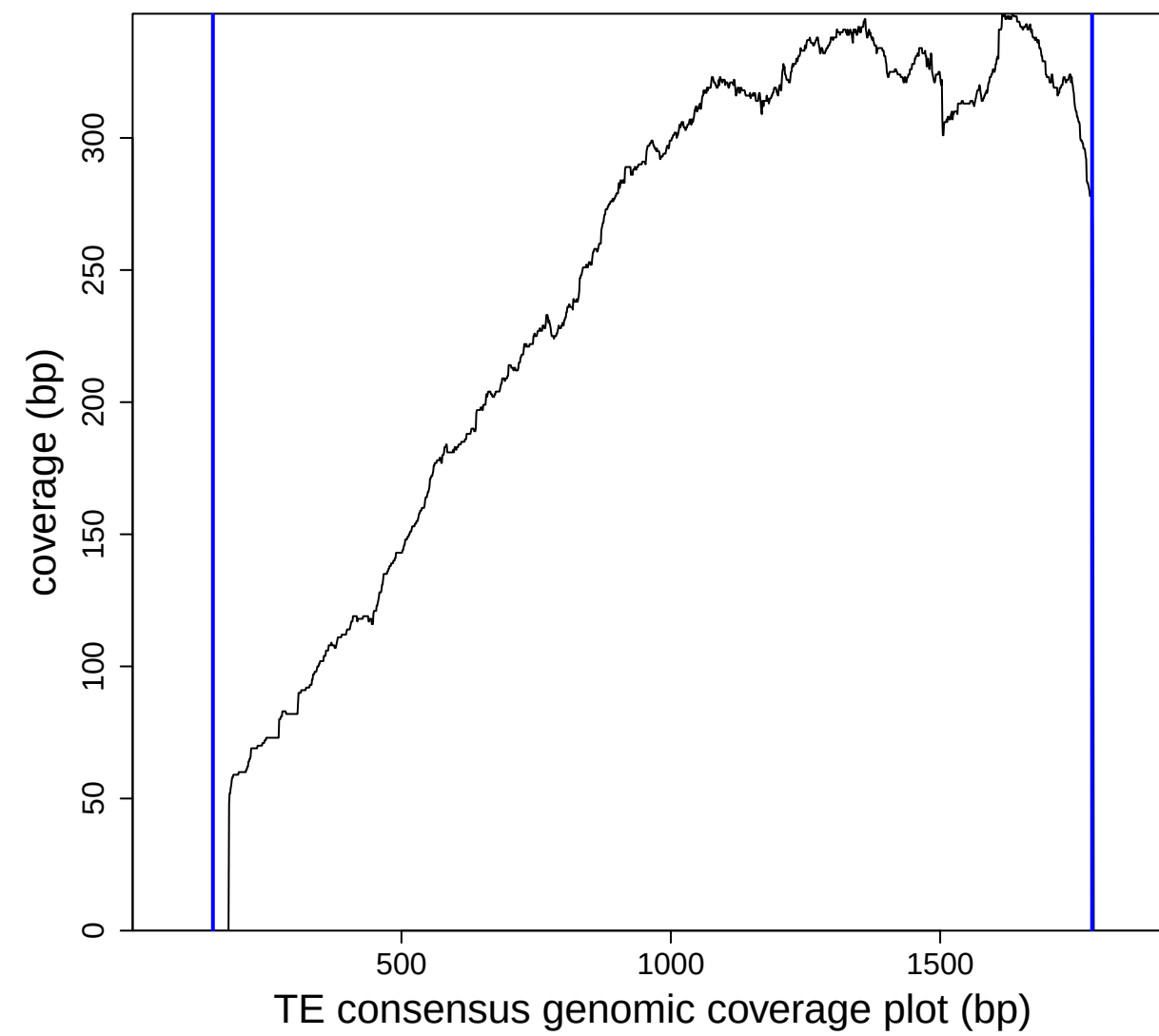
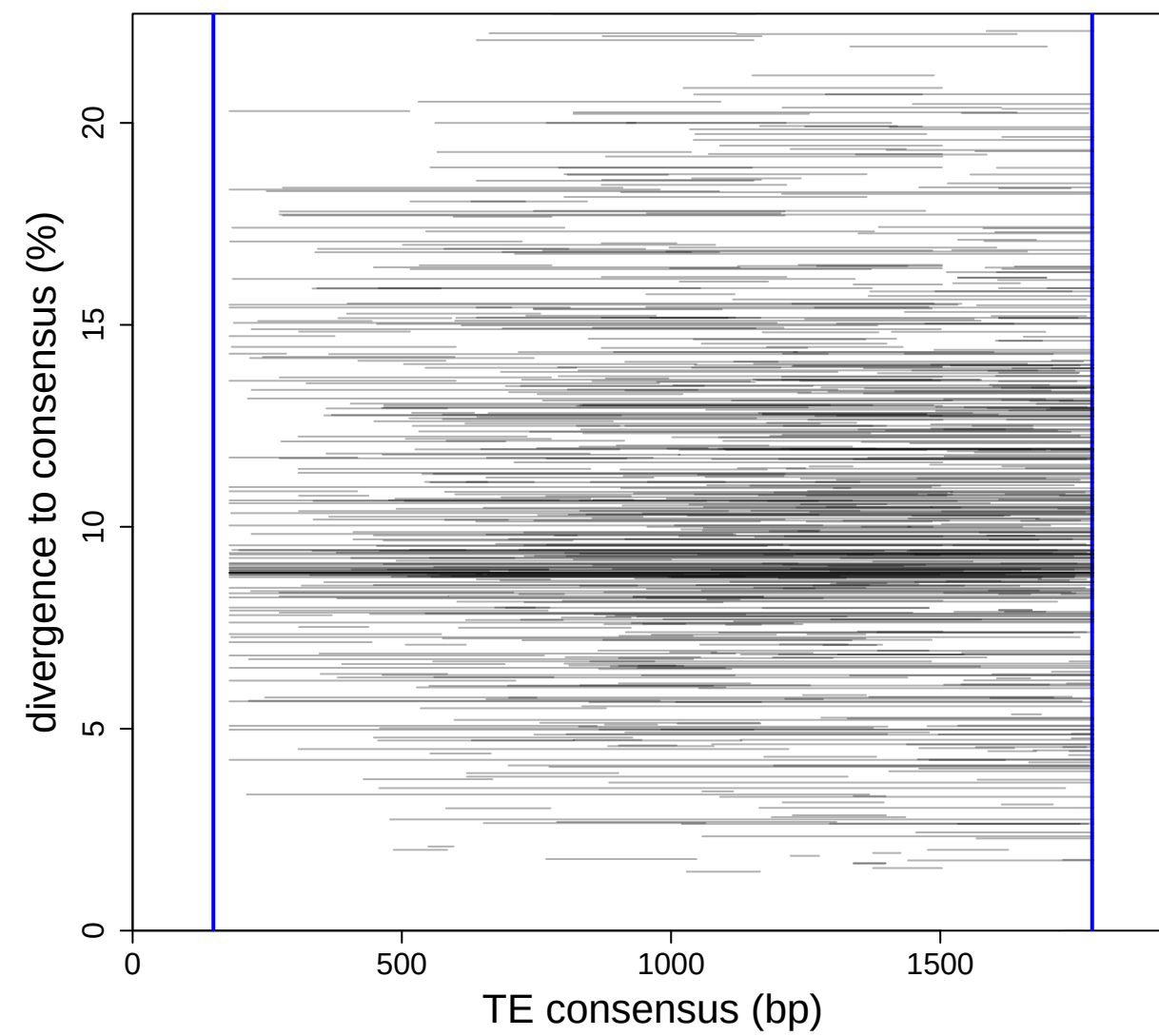
3.bed_fm 1.bed 0 0_n.bed_g 1.bed_fm 2.bed 0 0_bcln.fa aln.f
size: 1626bp; fragments: 777; full length: 36 (>=1463.4bp)

After TEtrimmer 1626 bp



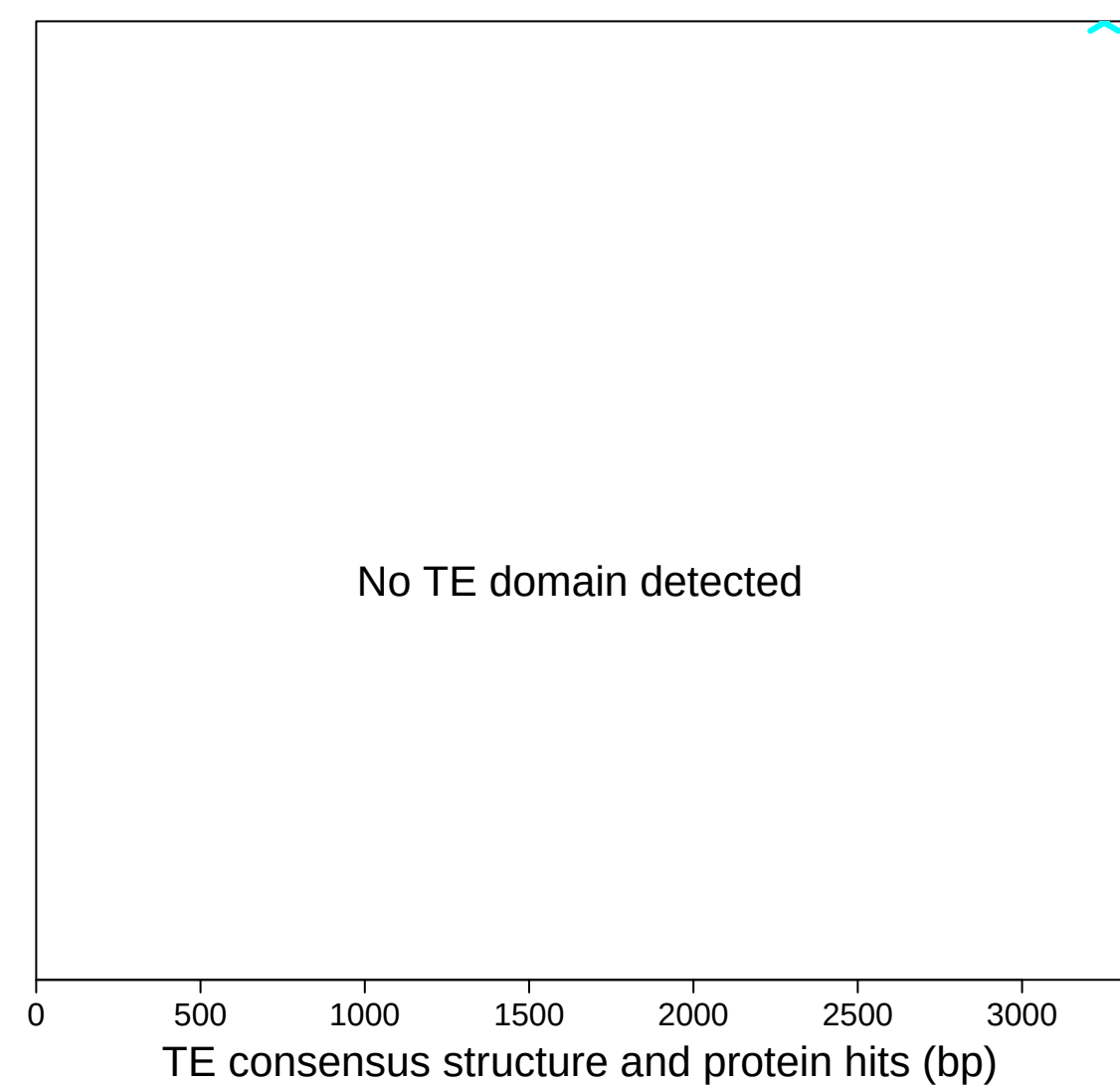
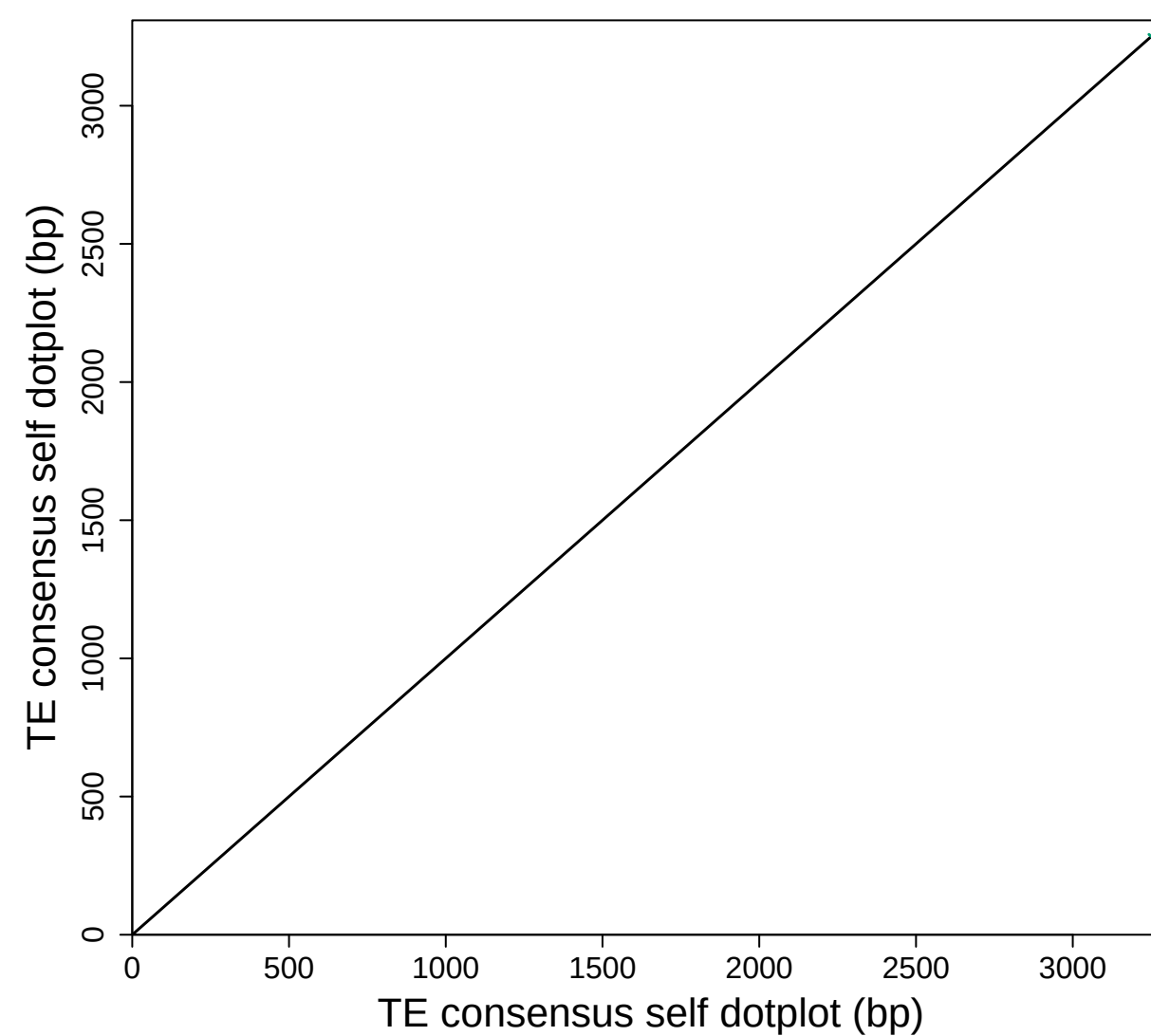
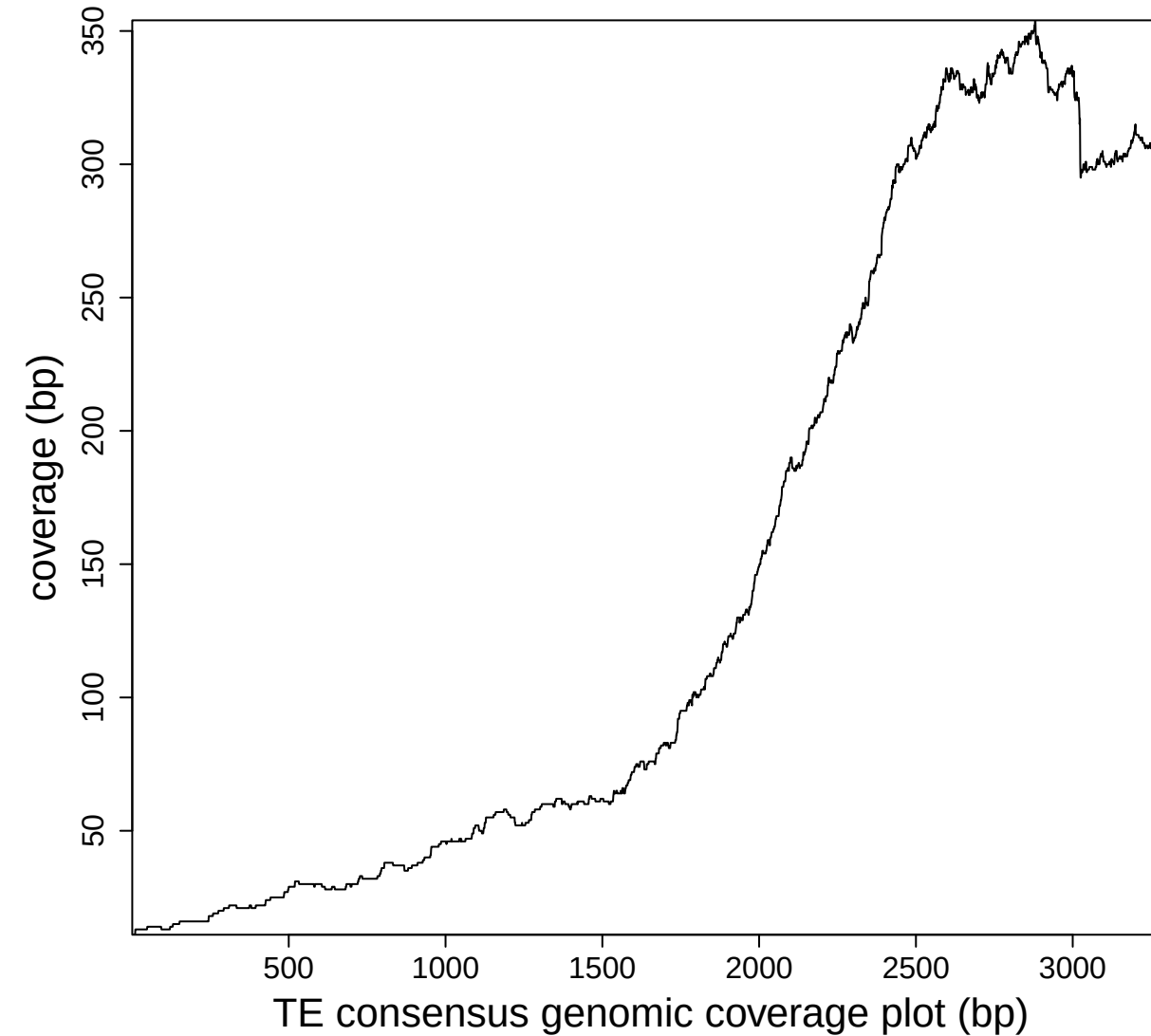
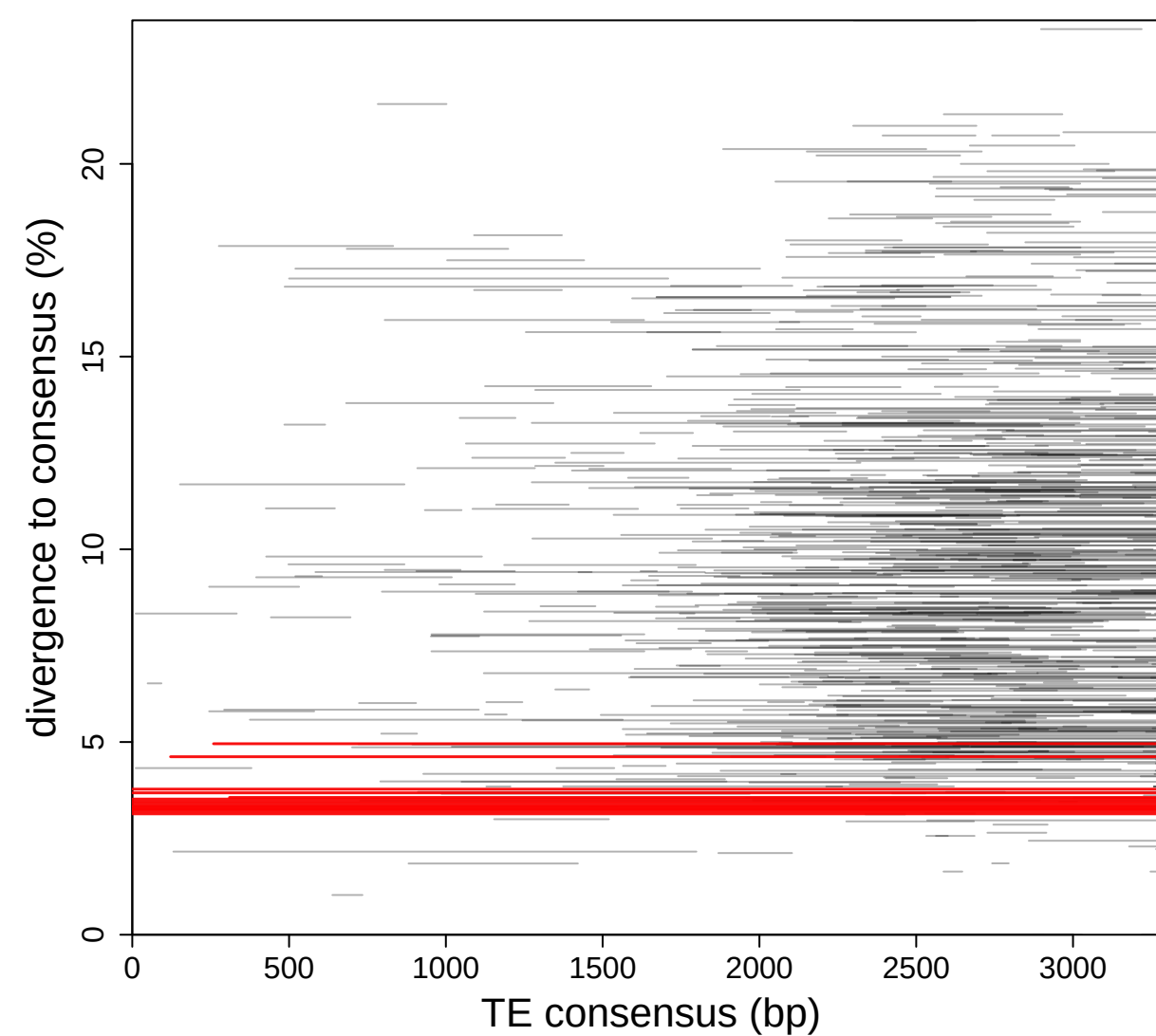
f.bed_g_3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 1932bp; fragments: 753; full length: 0 (>=1738.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

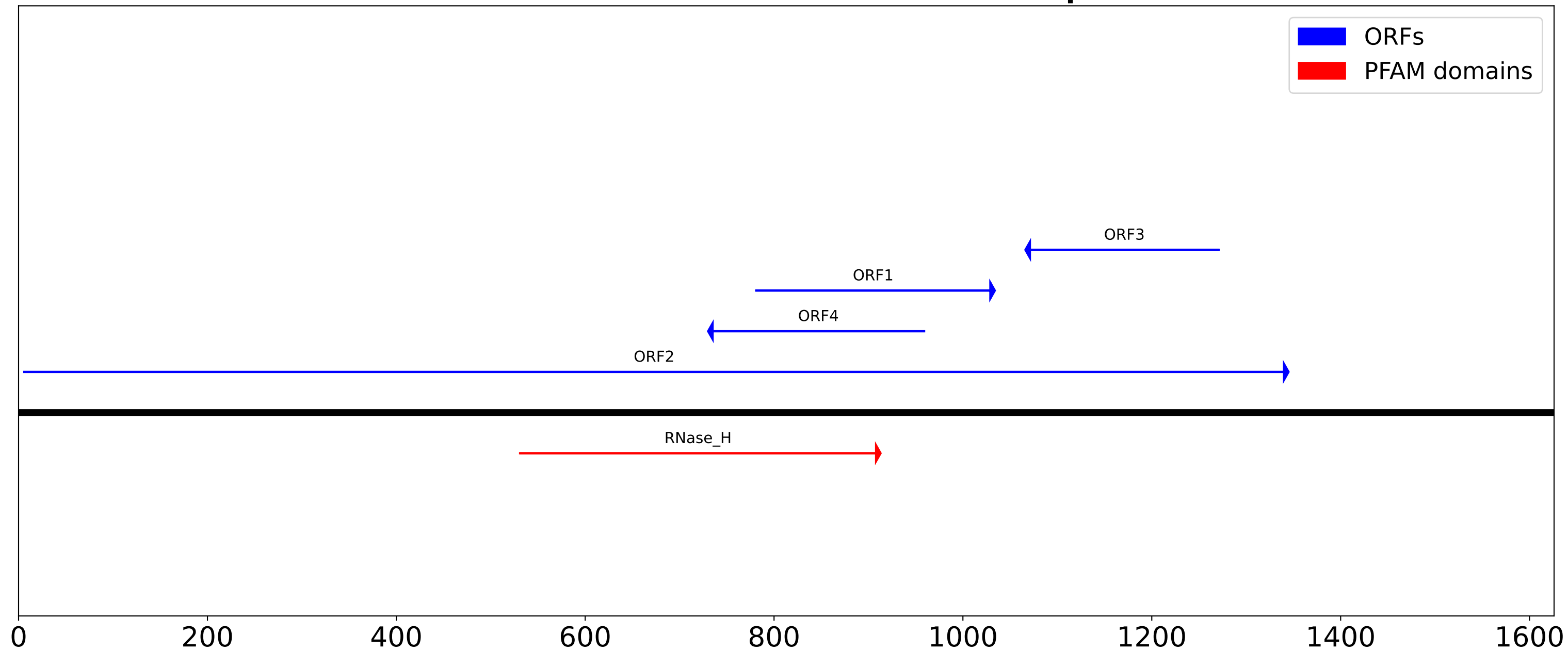


TE: rnd_5_family_3116
size: 3309bp; fragments: 836; full length: 14 (≥ 2978.1 bp)

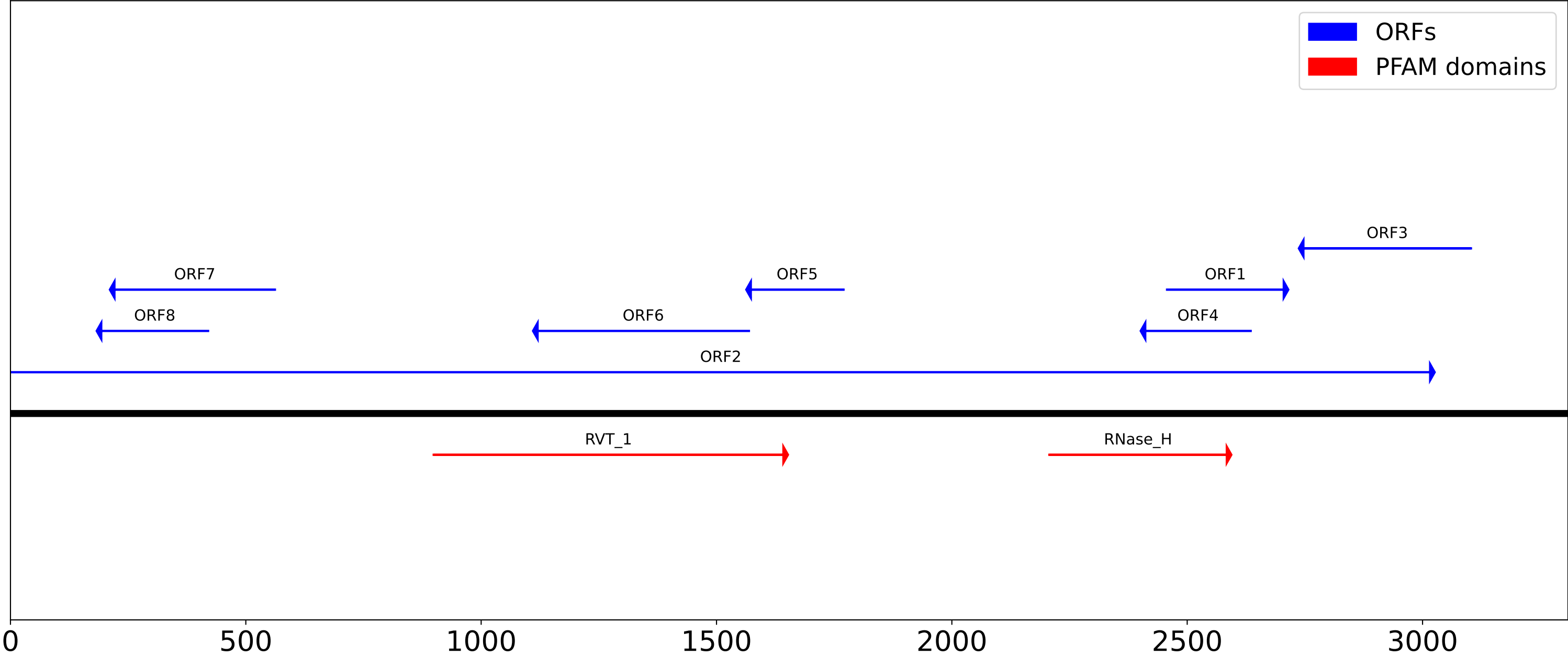
Before TEtrimmer 3309 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

