

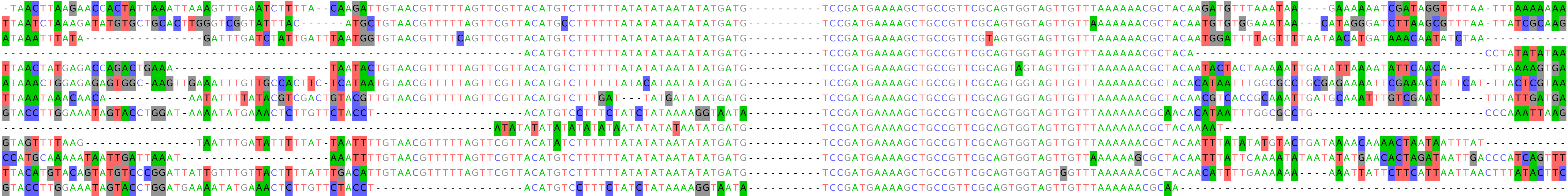
Start crop Point



End crop Point



MSA length = 1649



Start crop Point

End crop Point

0

10

0

250

500

750

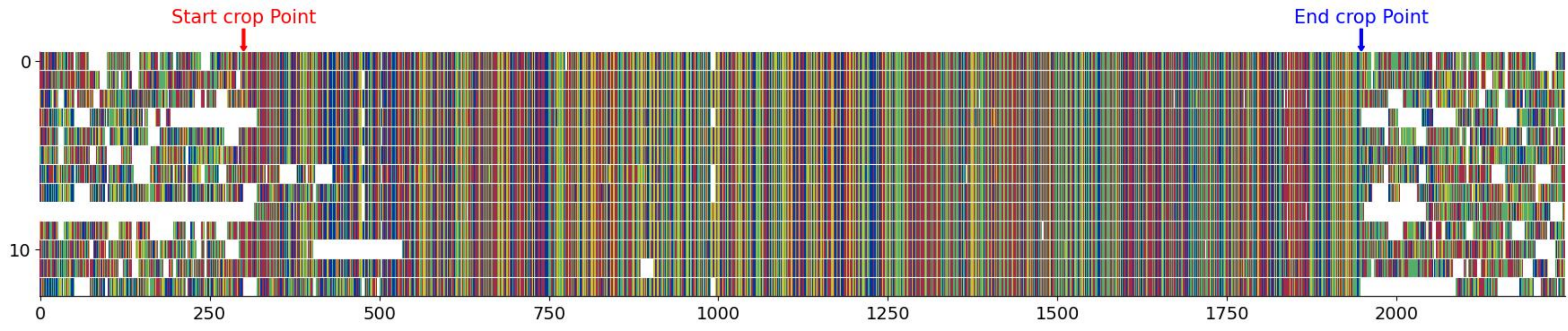
1000

1250

1500

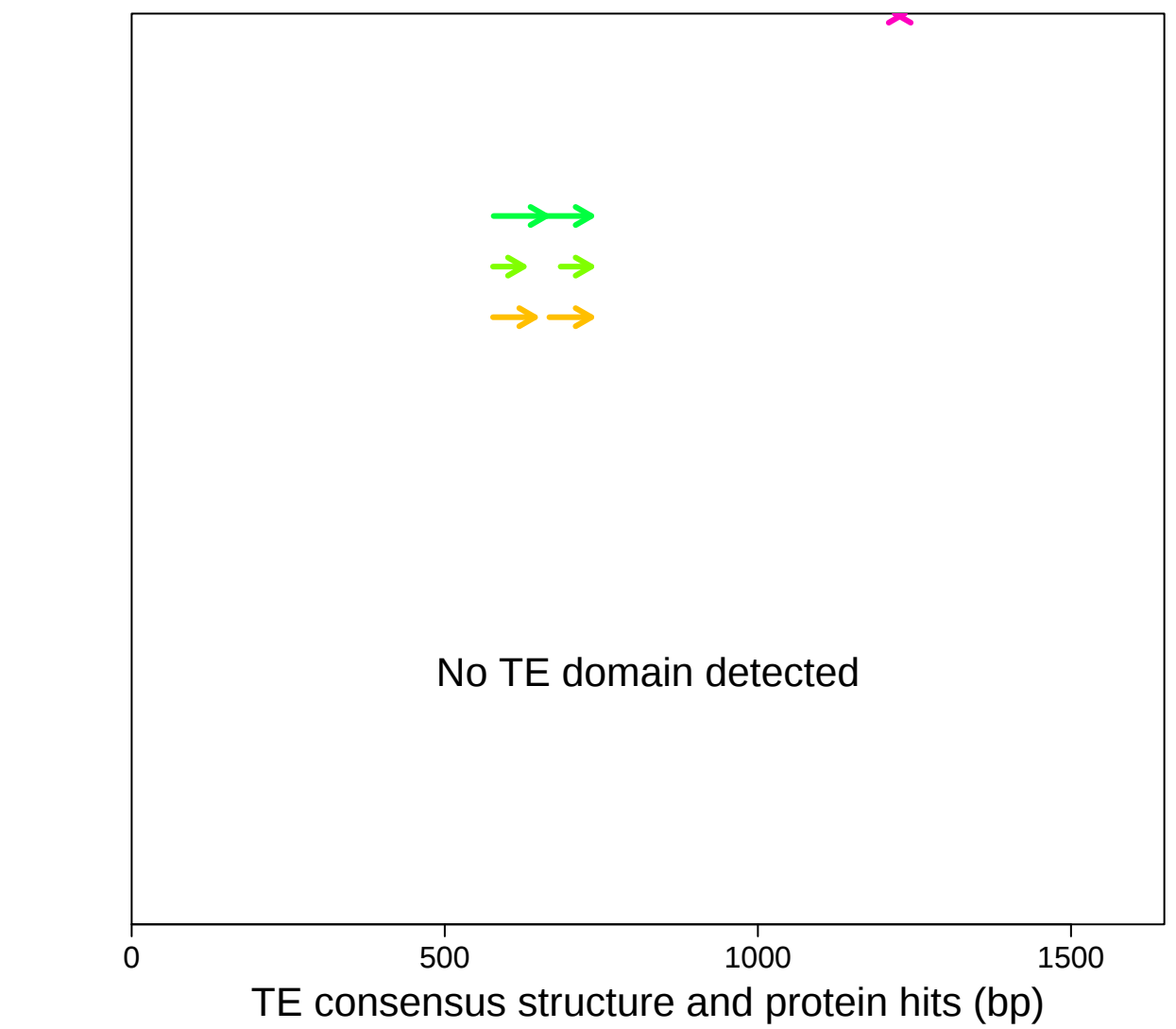
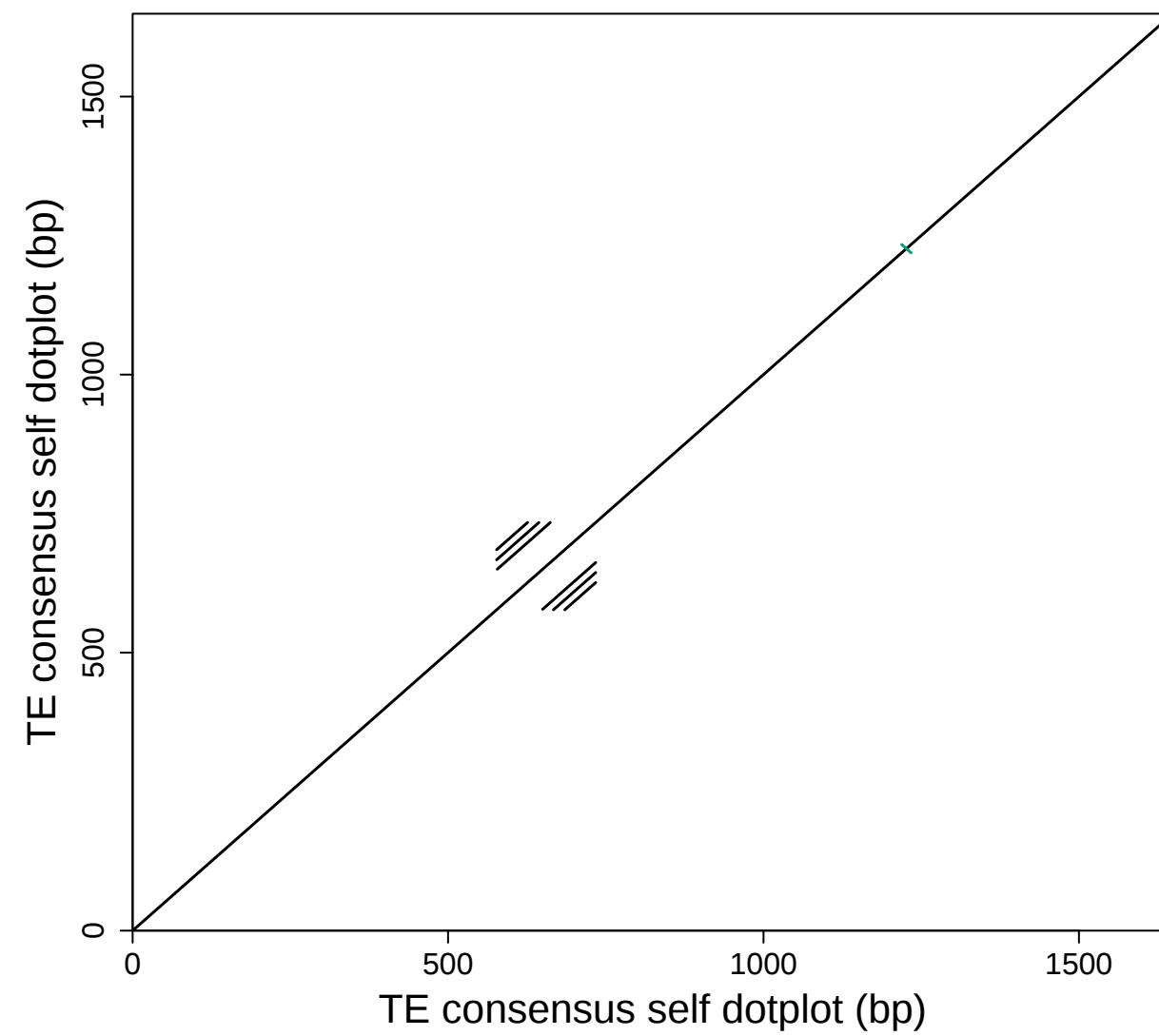
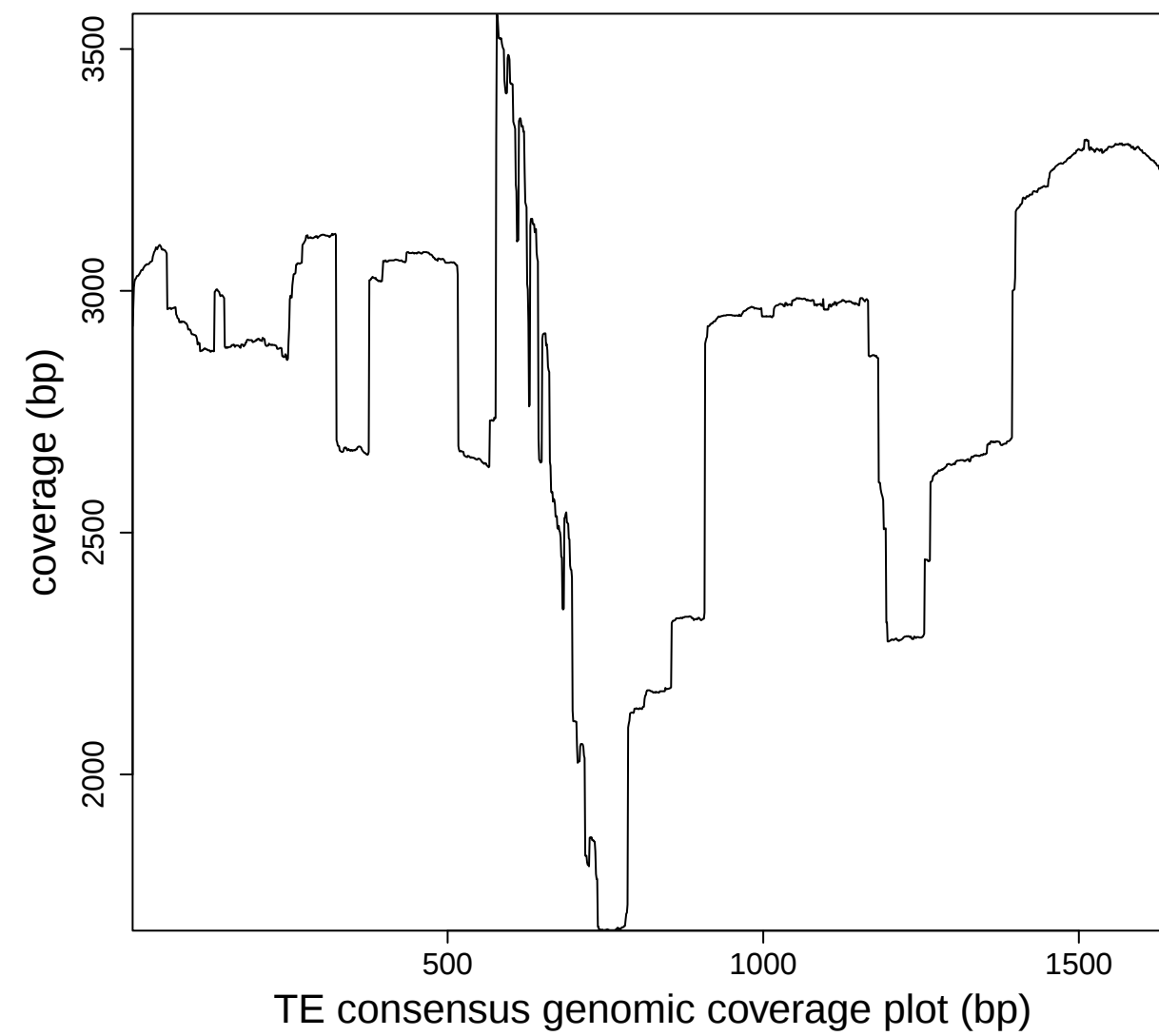
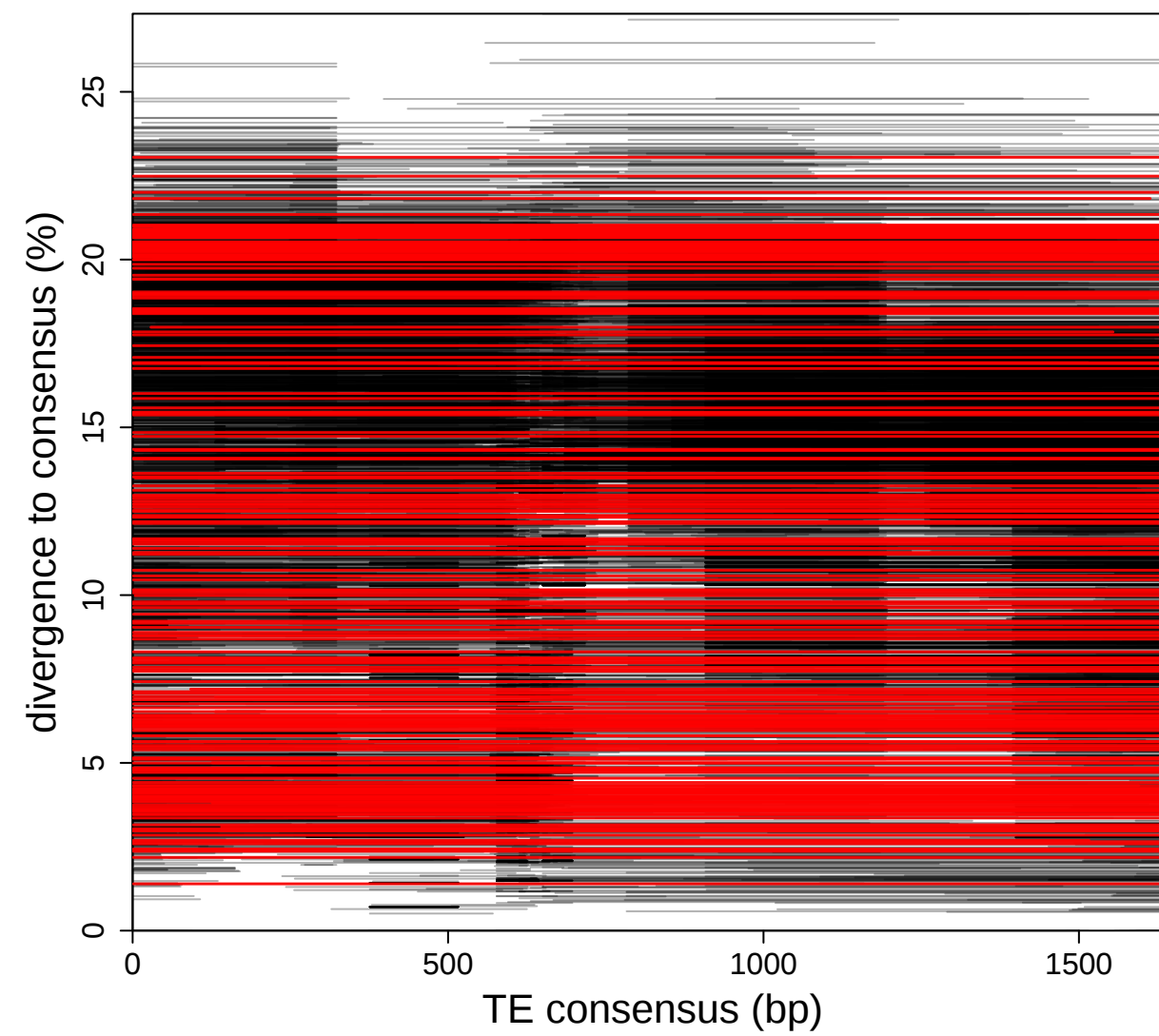
1750

2000



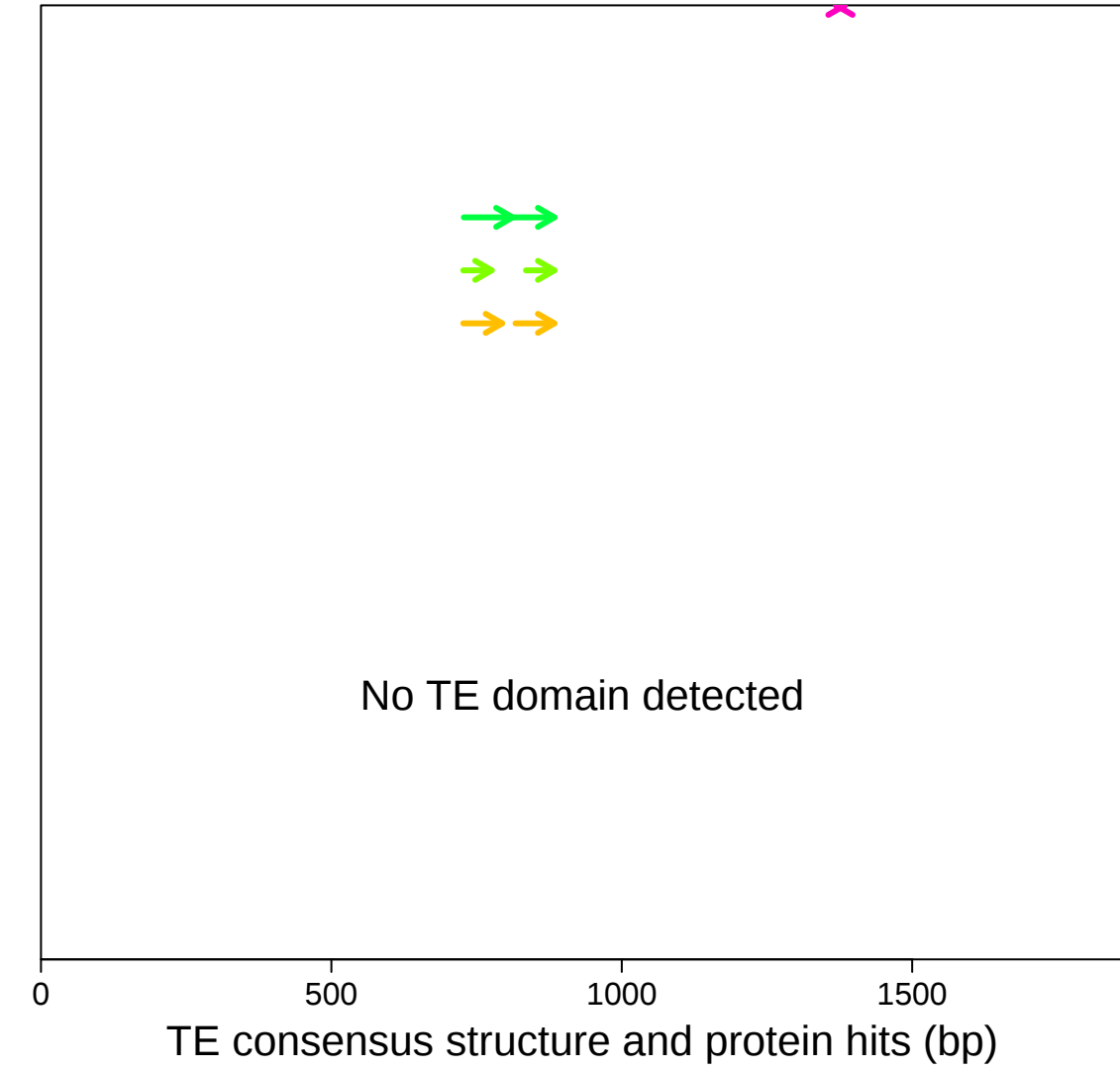
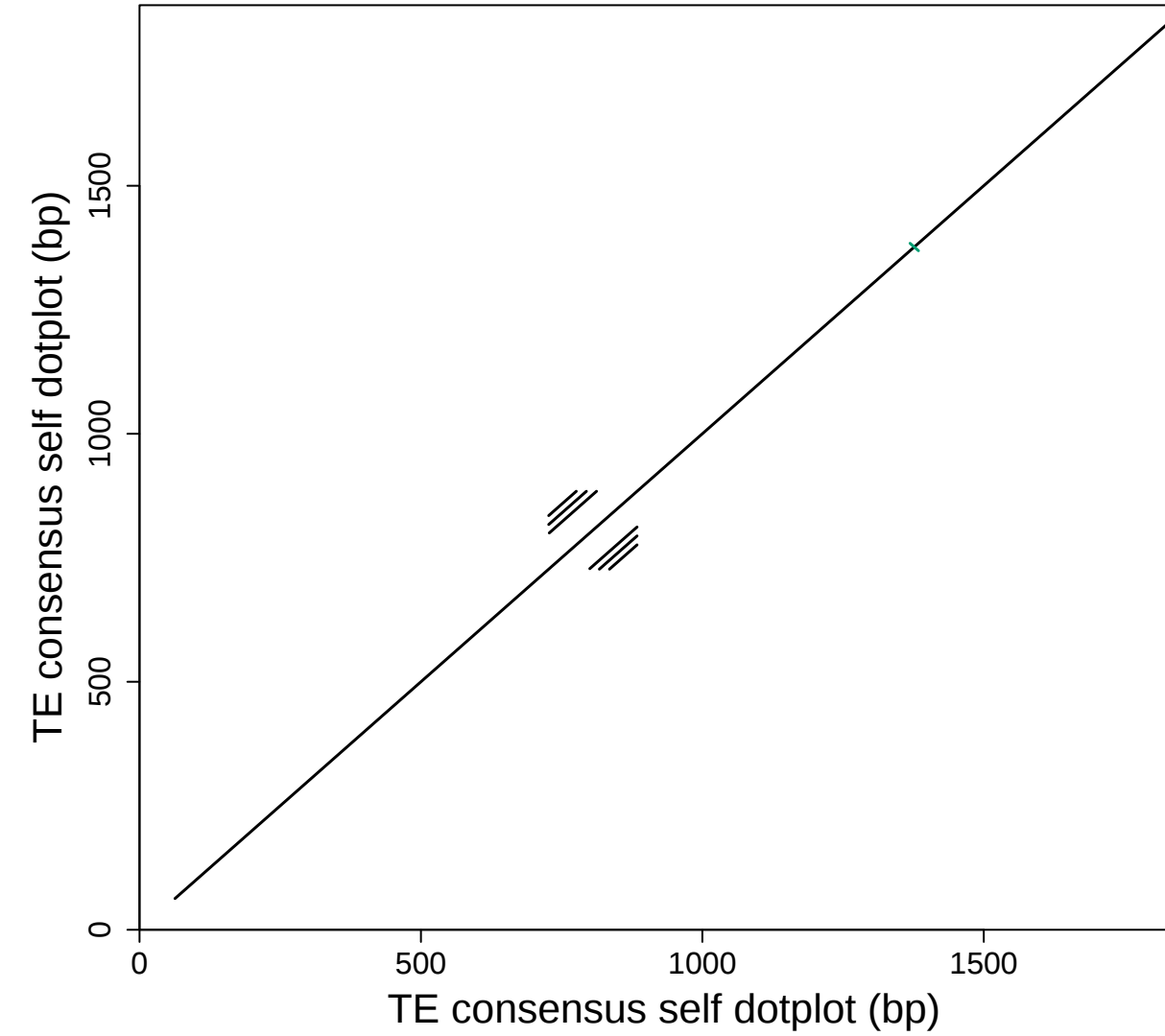
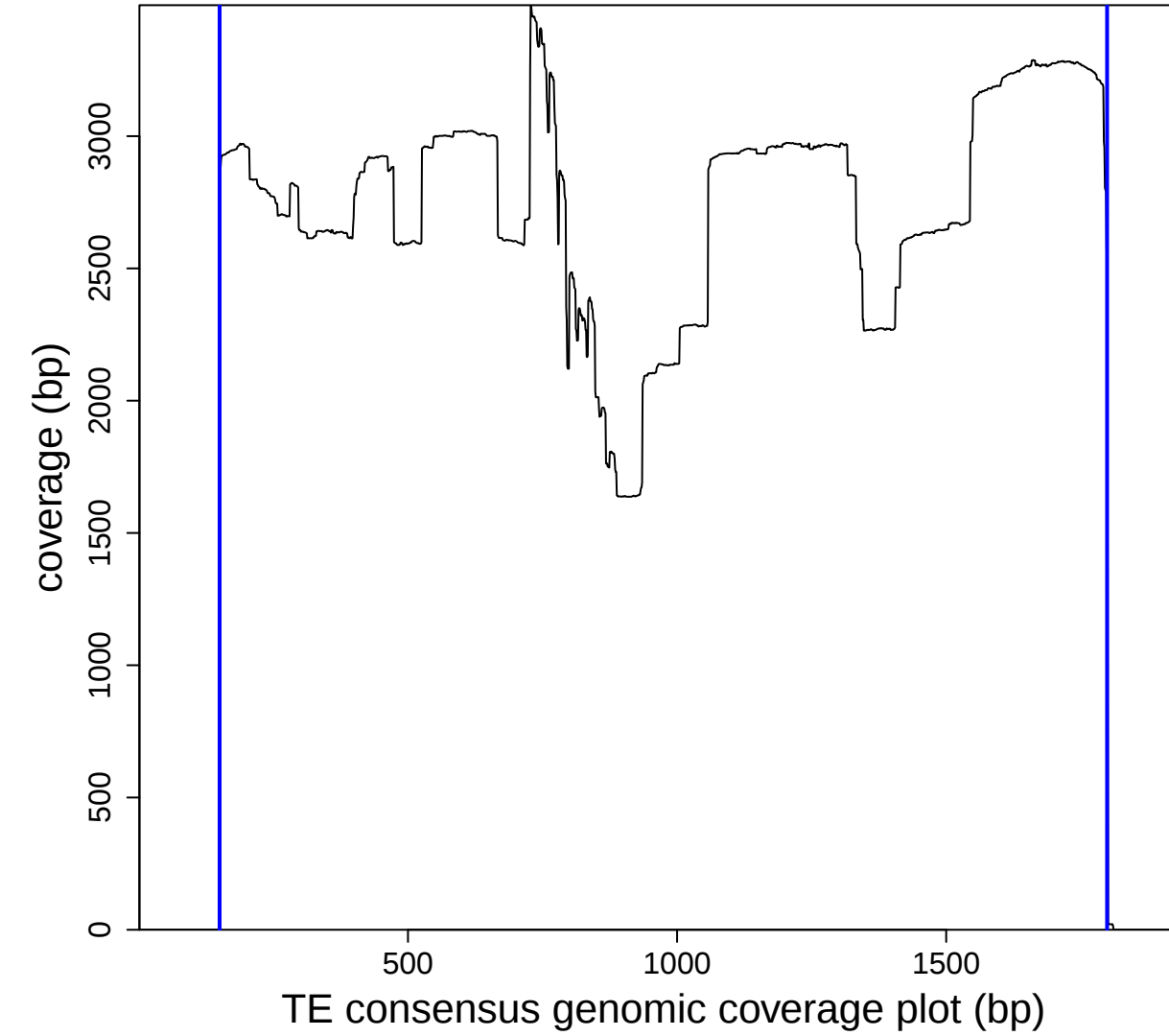
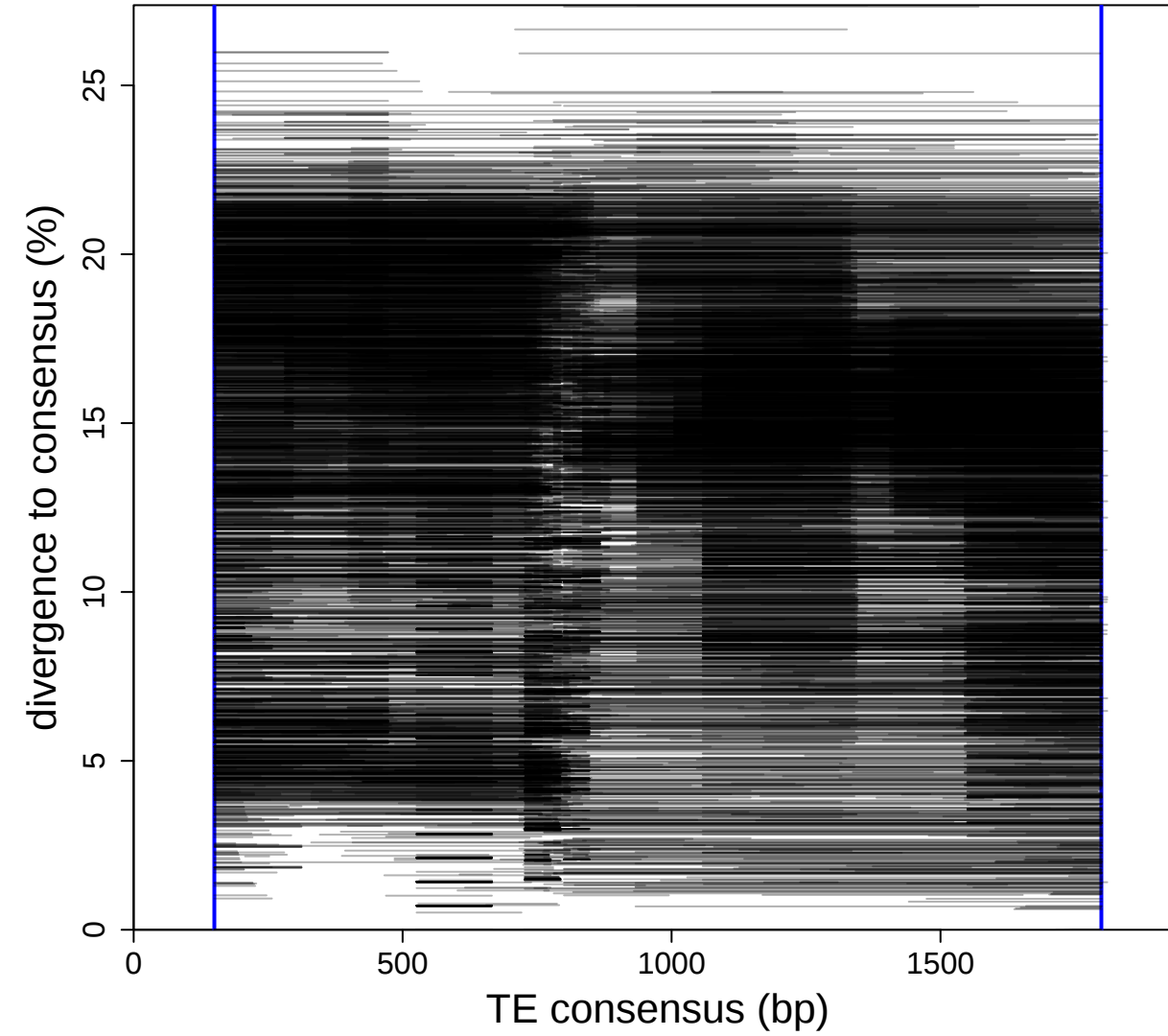
l.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 1649bp; fragments: 9767; full length: 230 (>=1484.1bp)

After TEtrimmer 1649 bp



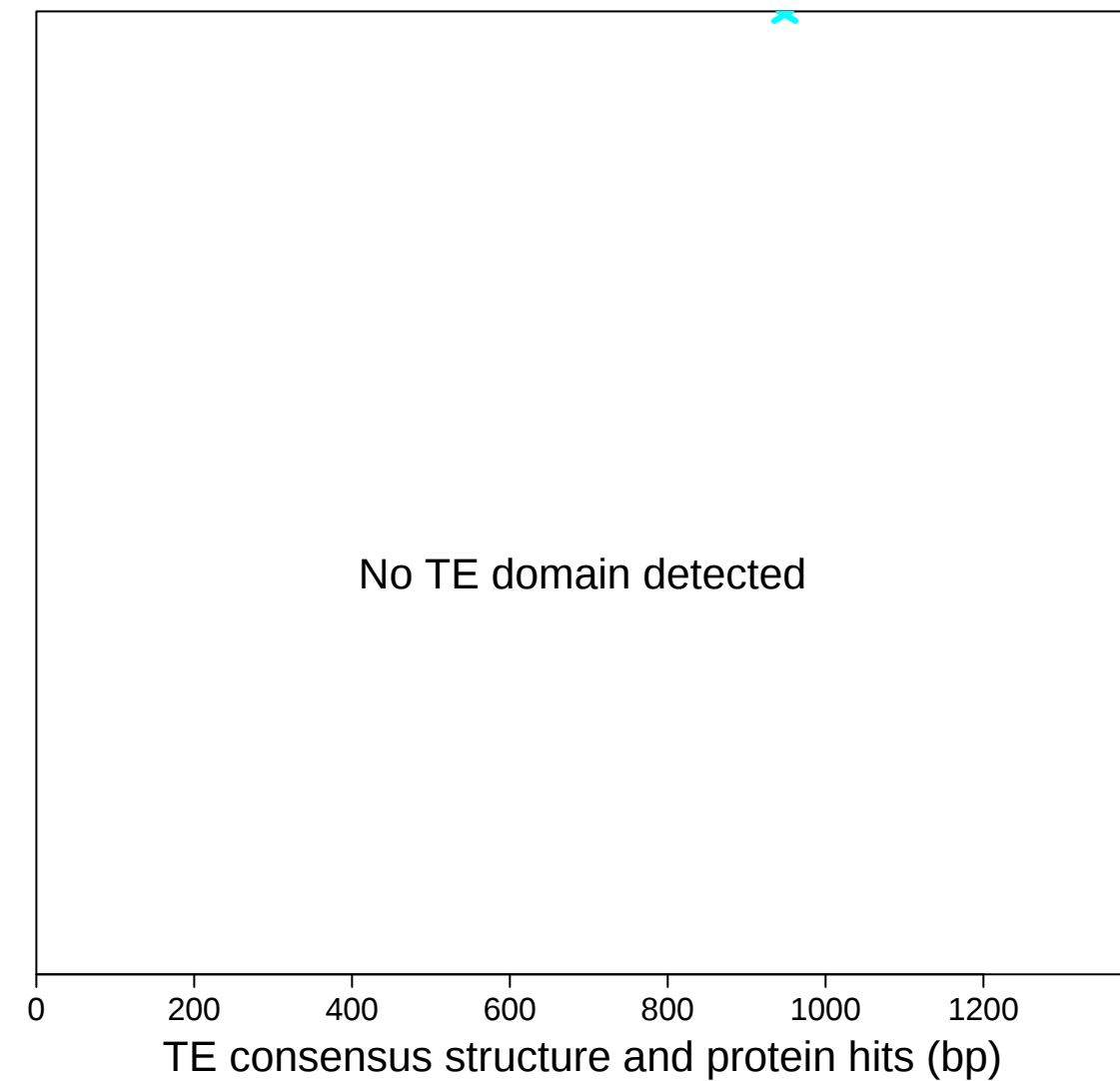
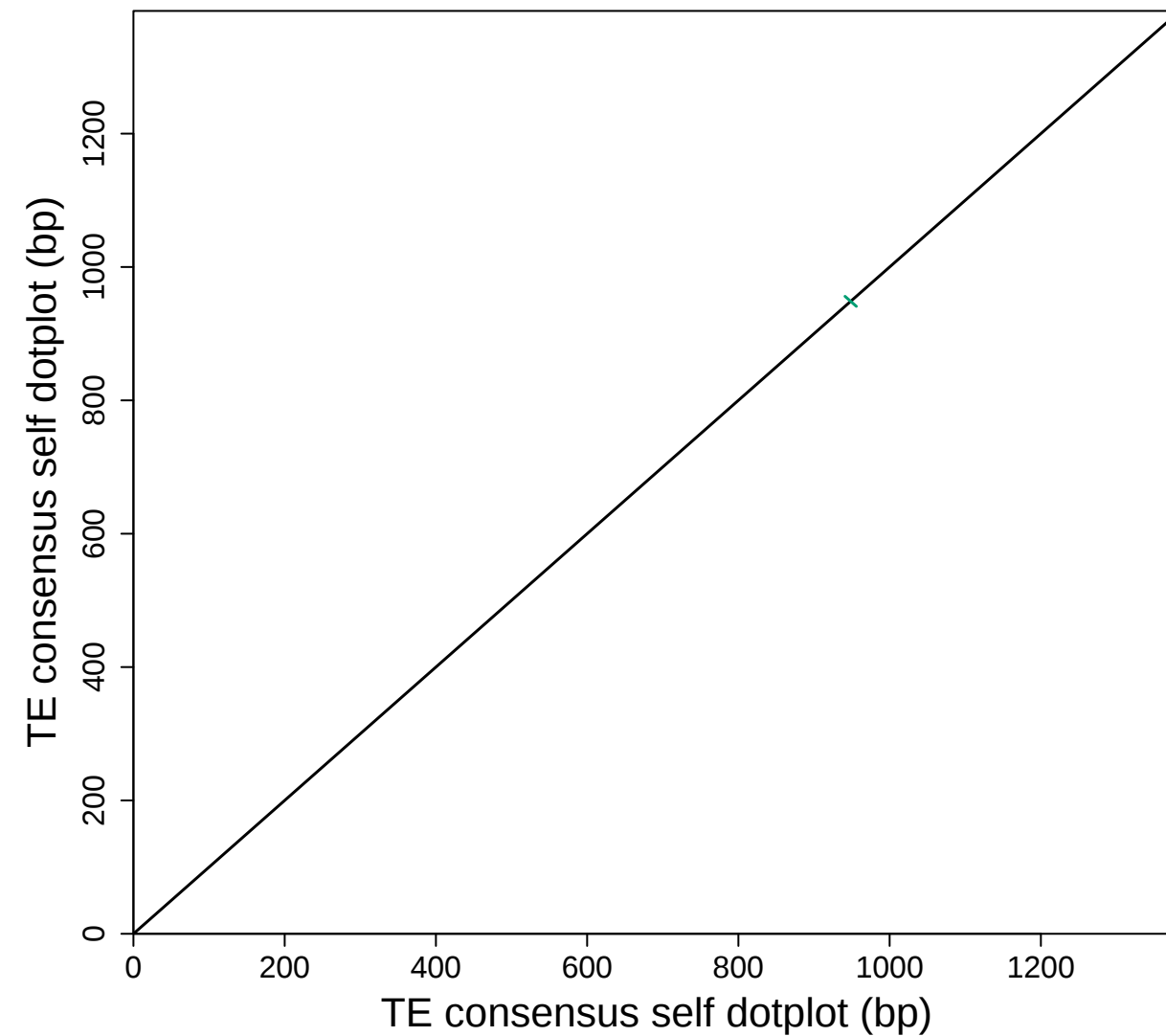
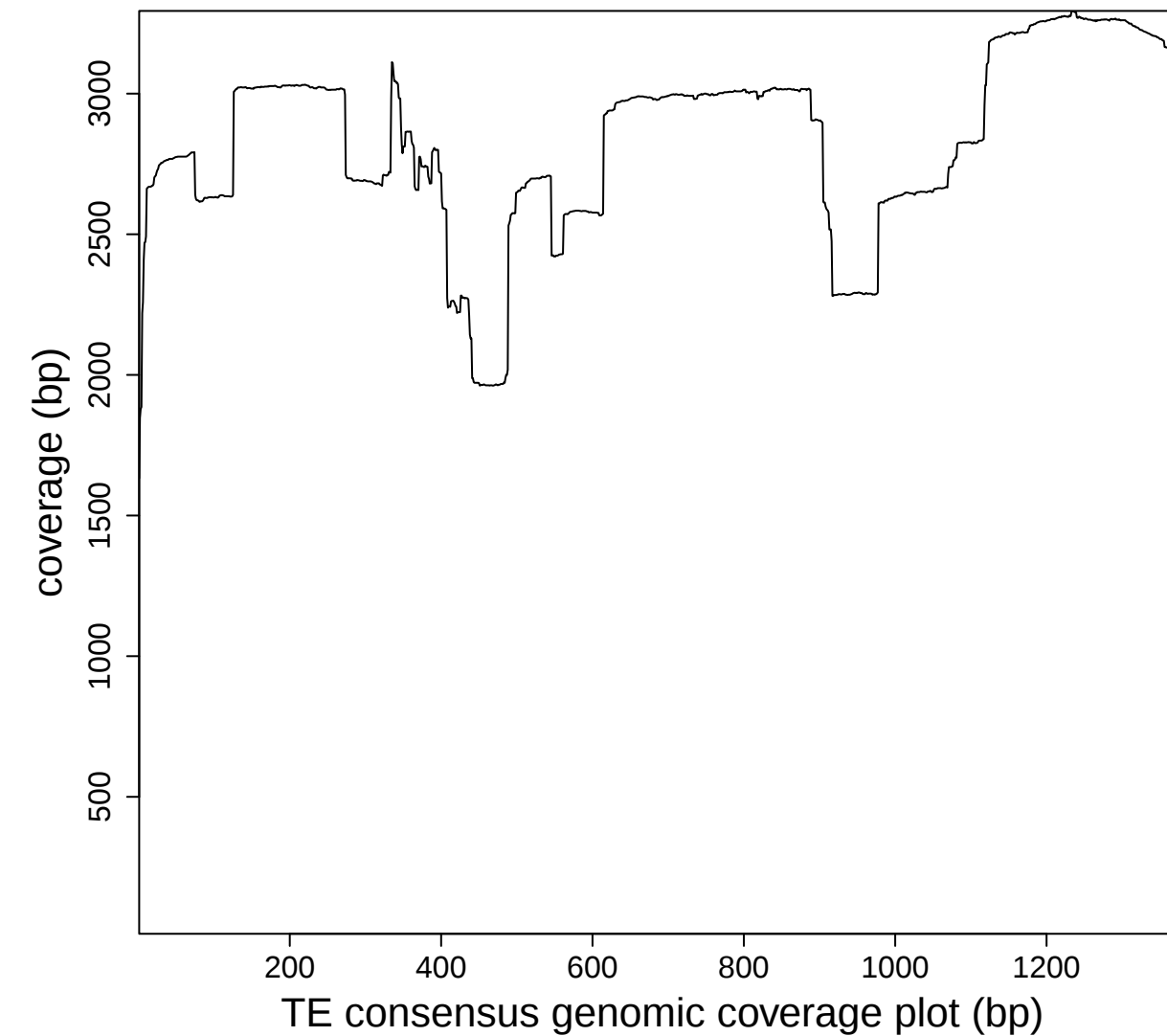
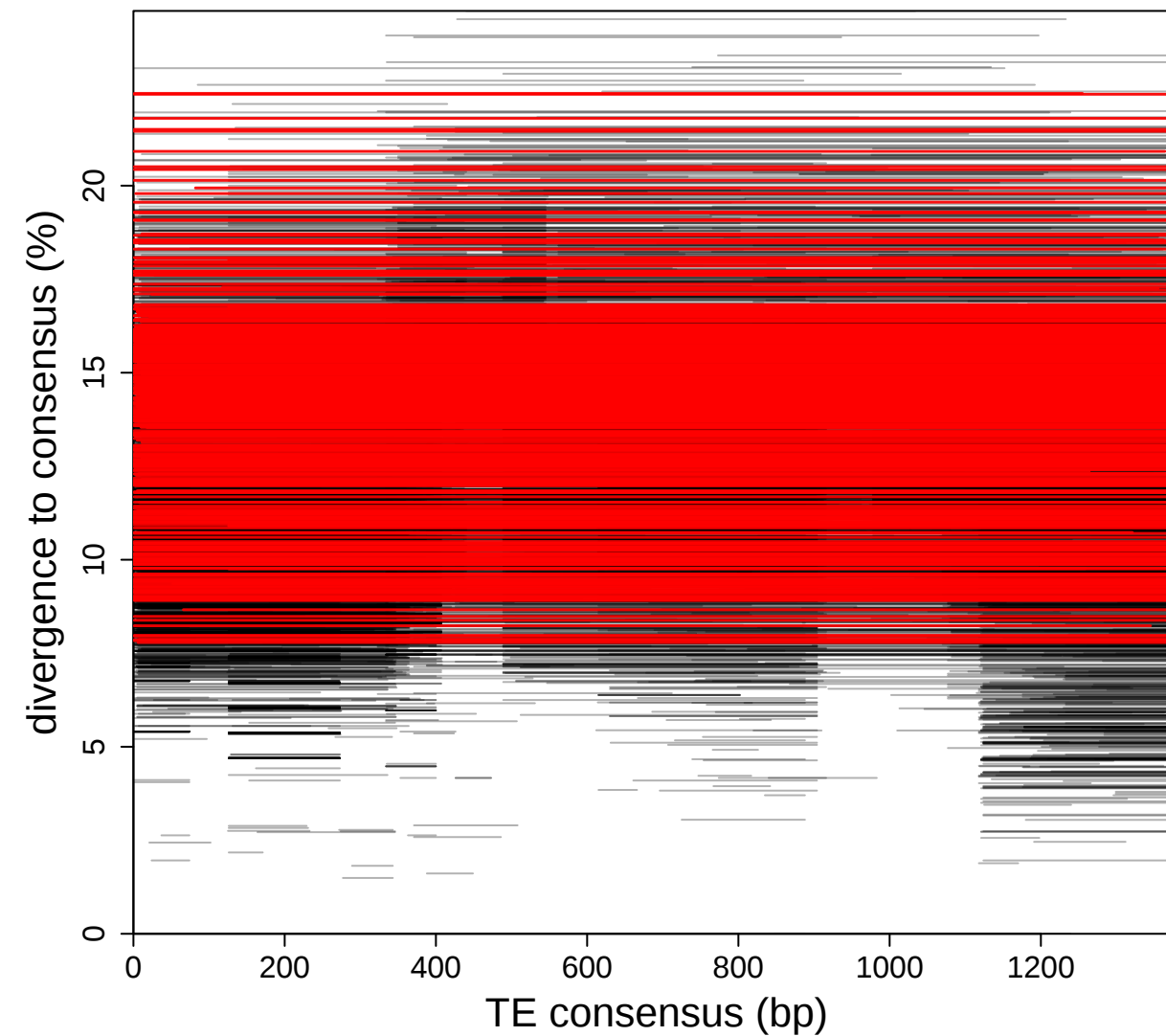
4.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1949bp; fragments: 9548; full length: 0 (>=1754.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

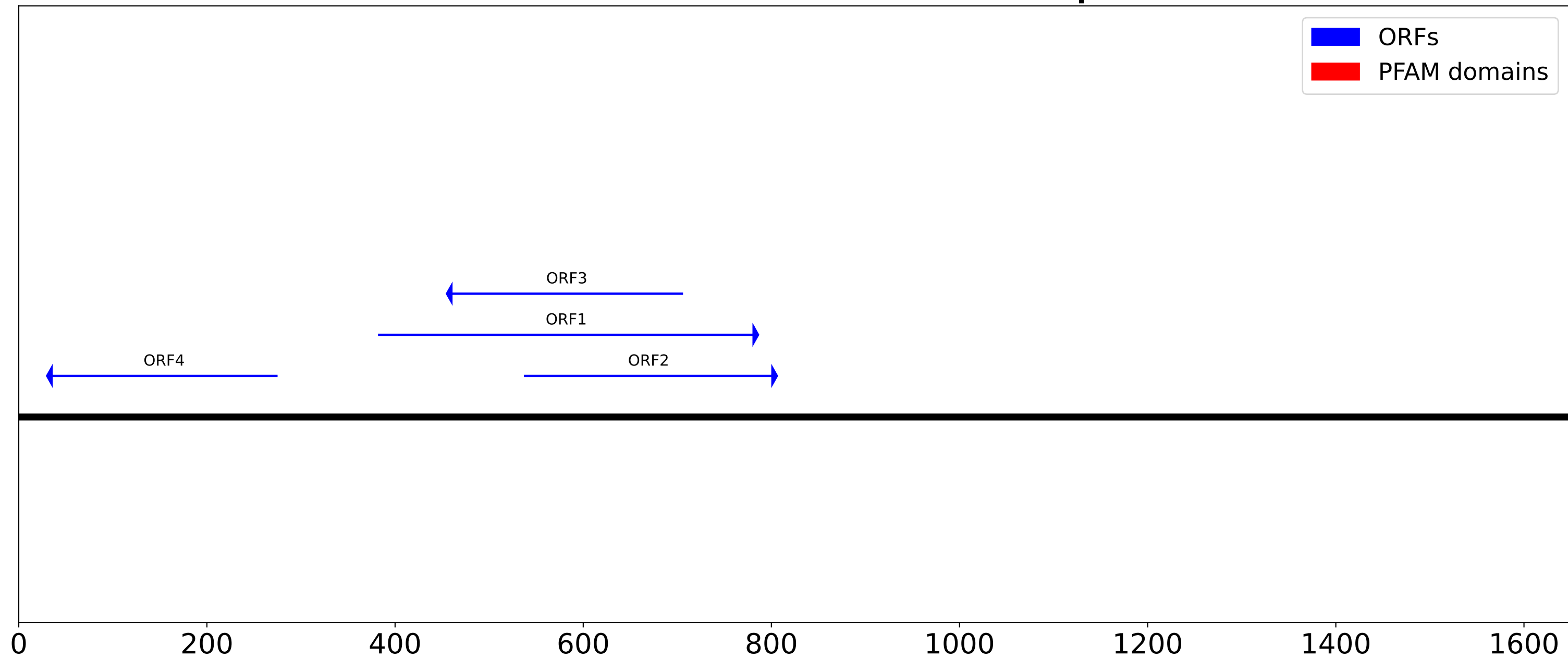


TE: rnd_3_family_414
size: 1384bp; fragments: 7270; full length: 450 (≥ 1245.6 bp)

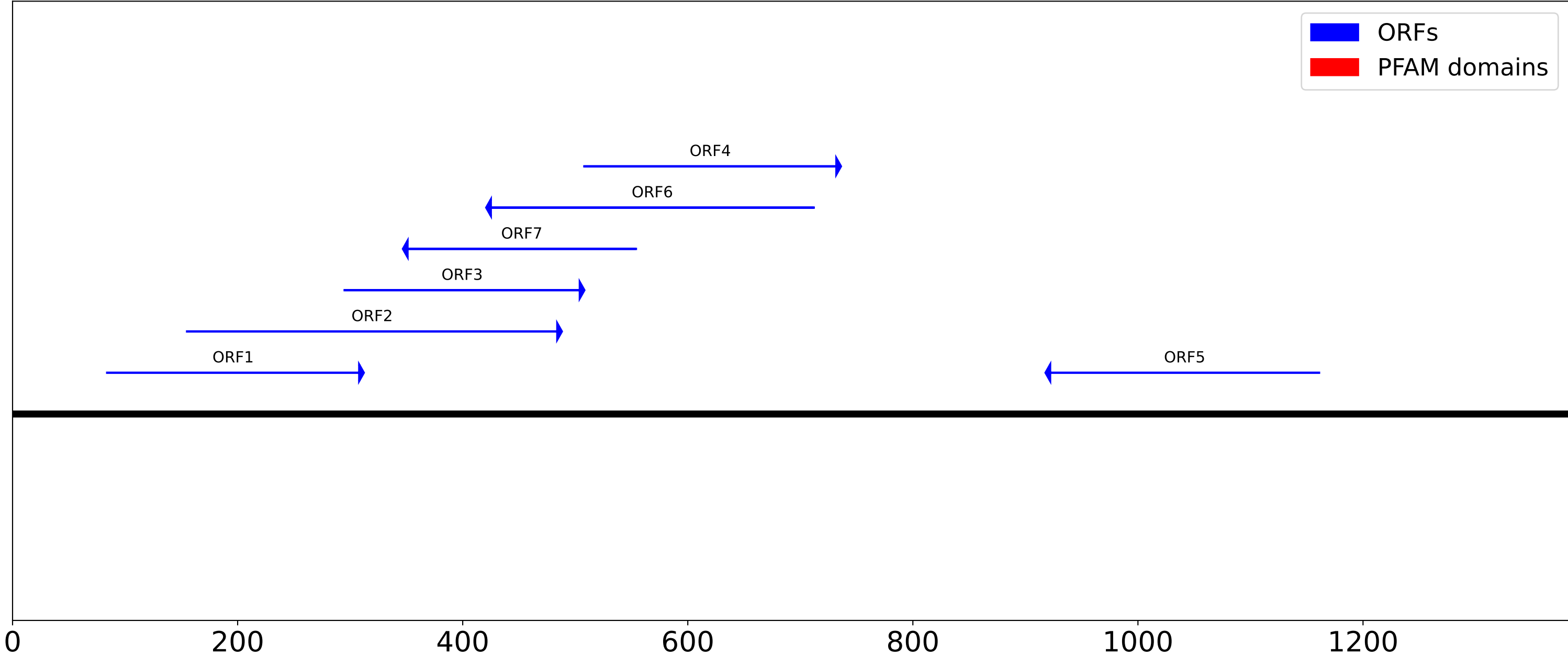
Before TEtrimmer 1384 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

