

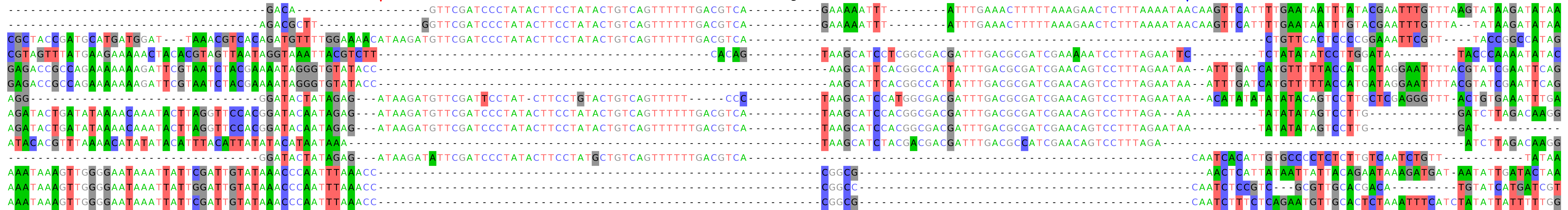
Start crop Point



End crop Point

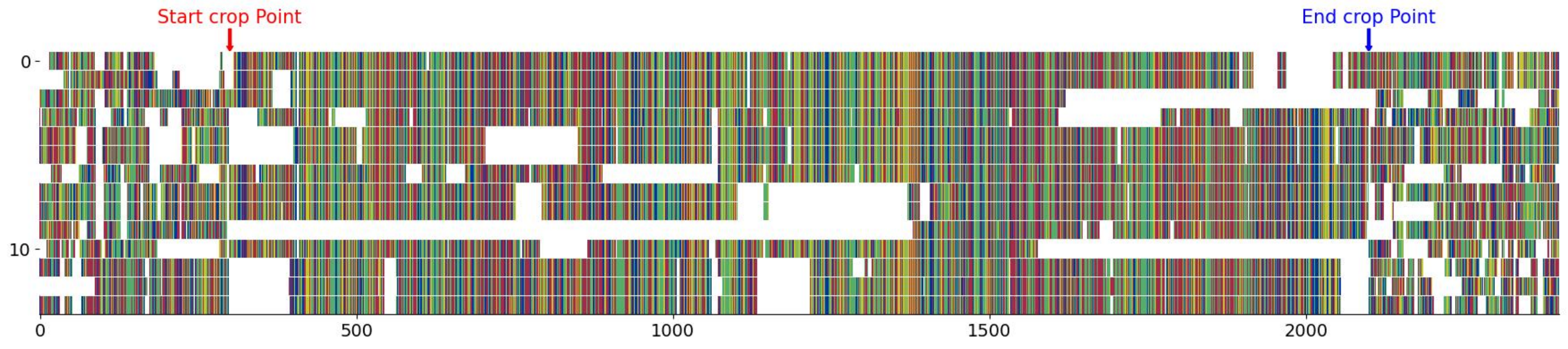


MSA length = 1800



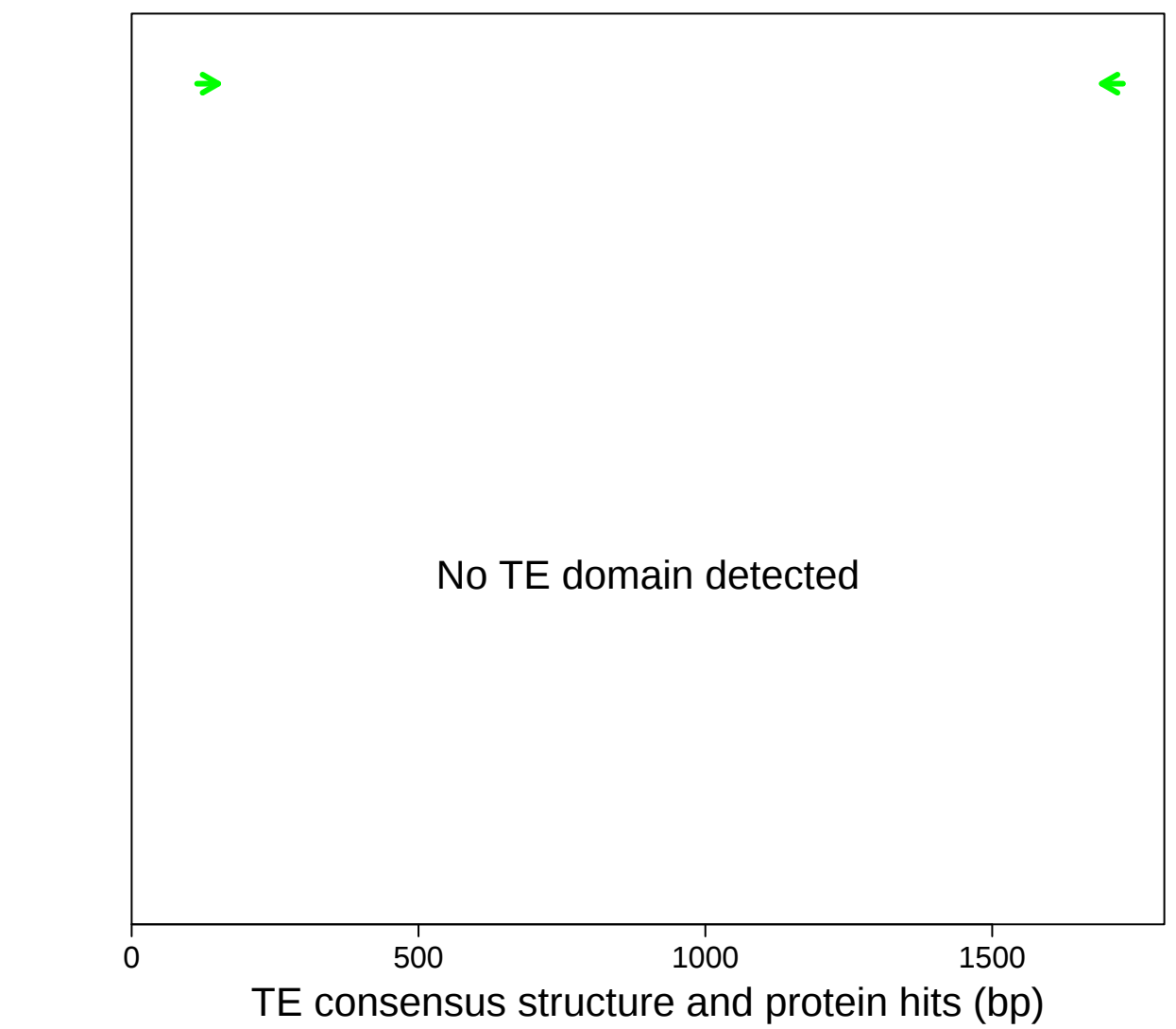
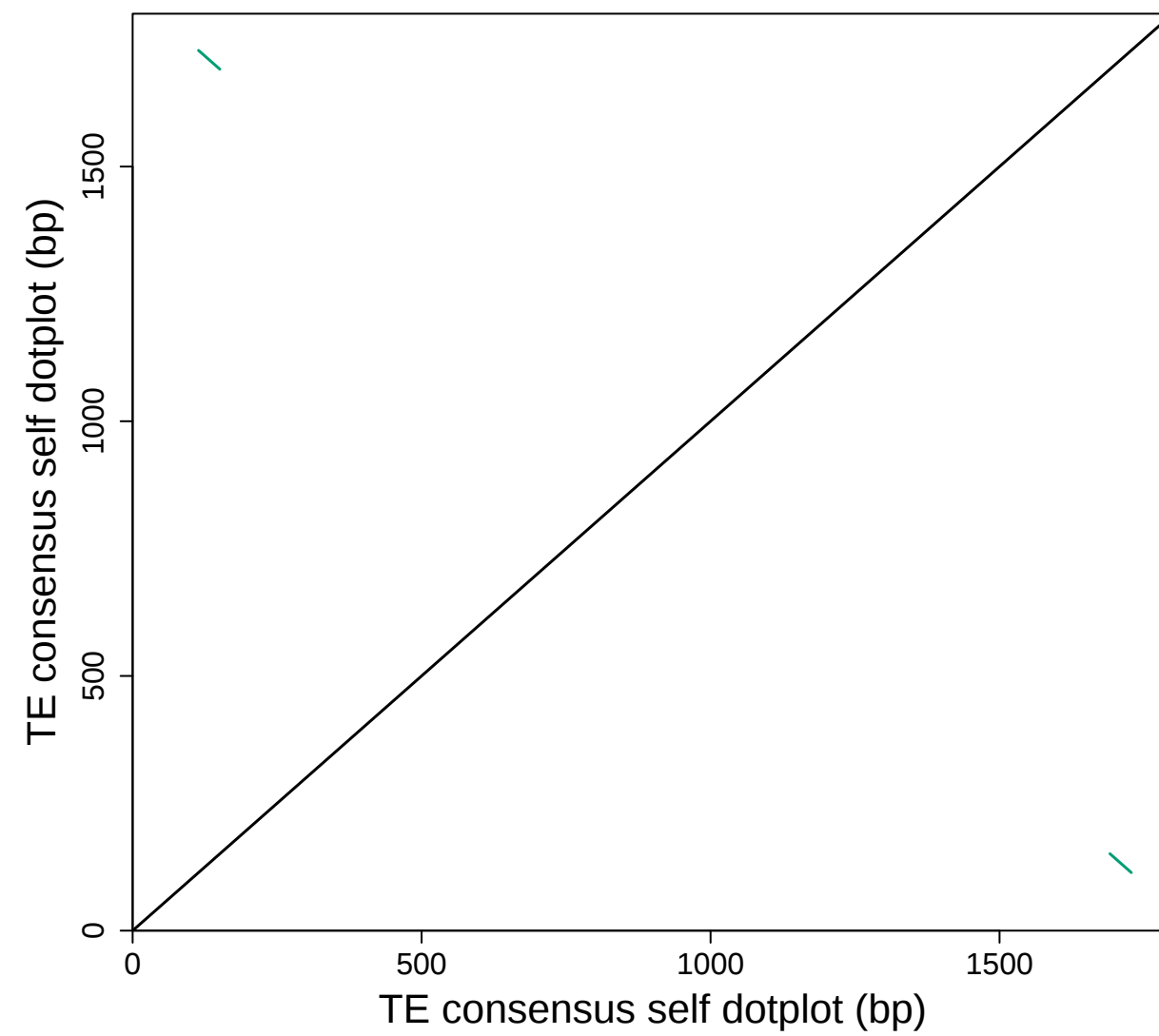
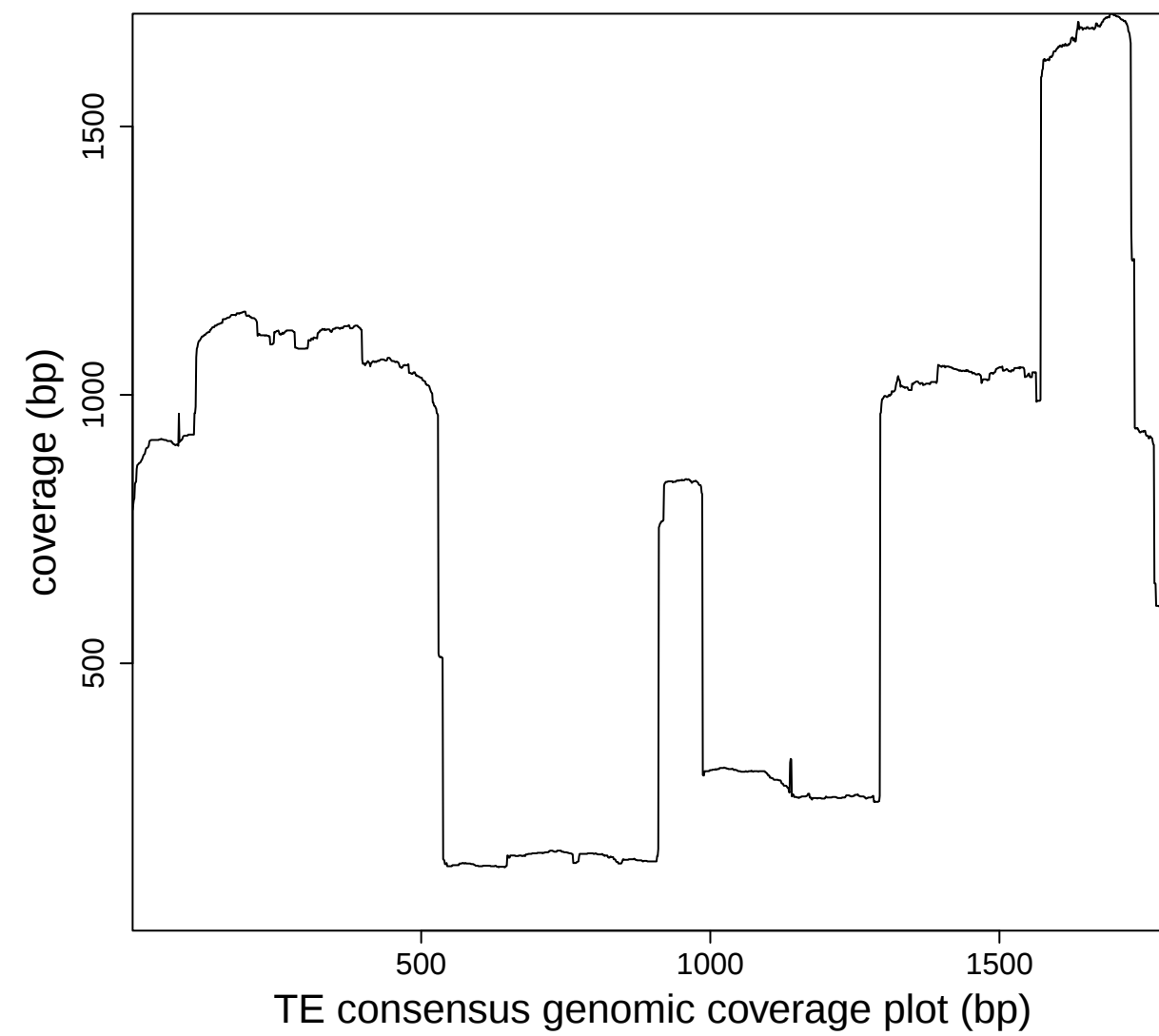
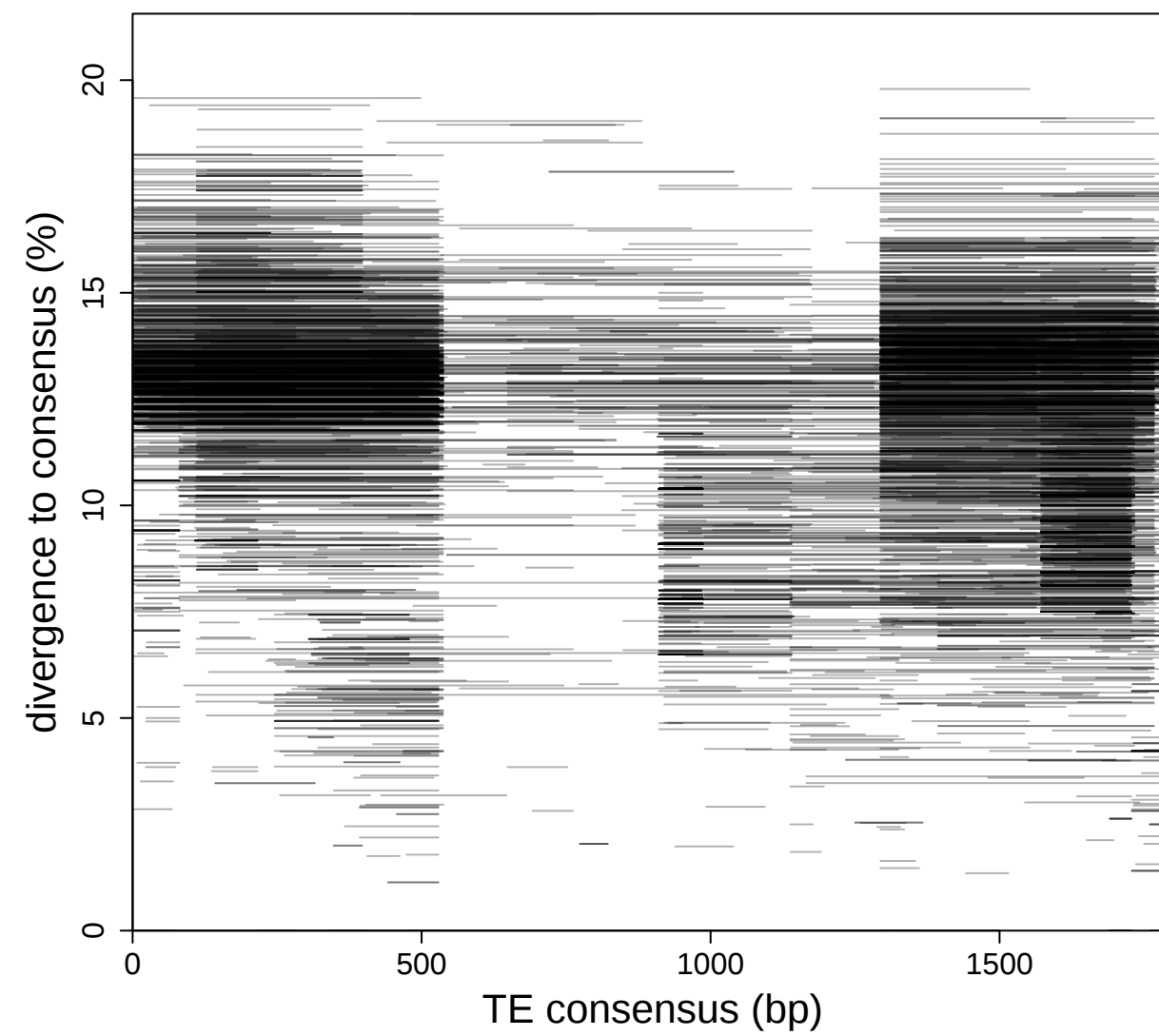
Start crop Point

End crop Point



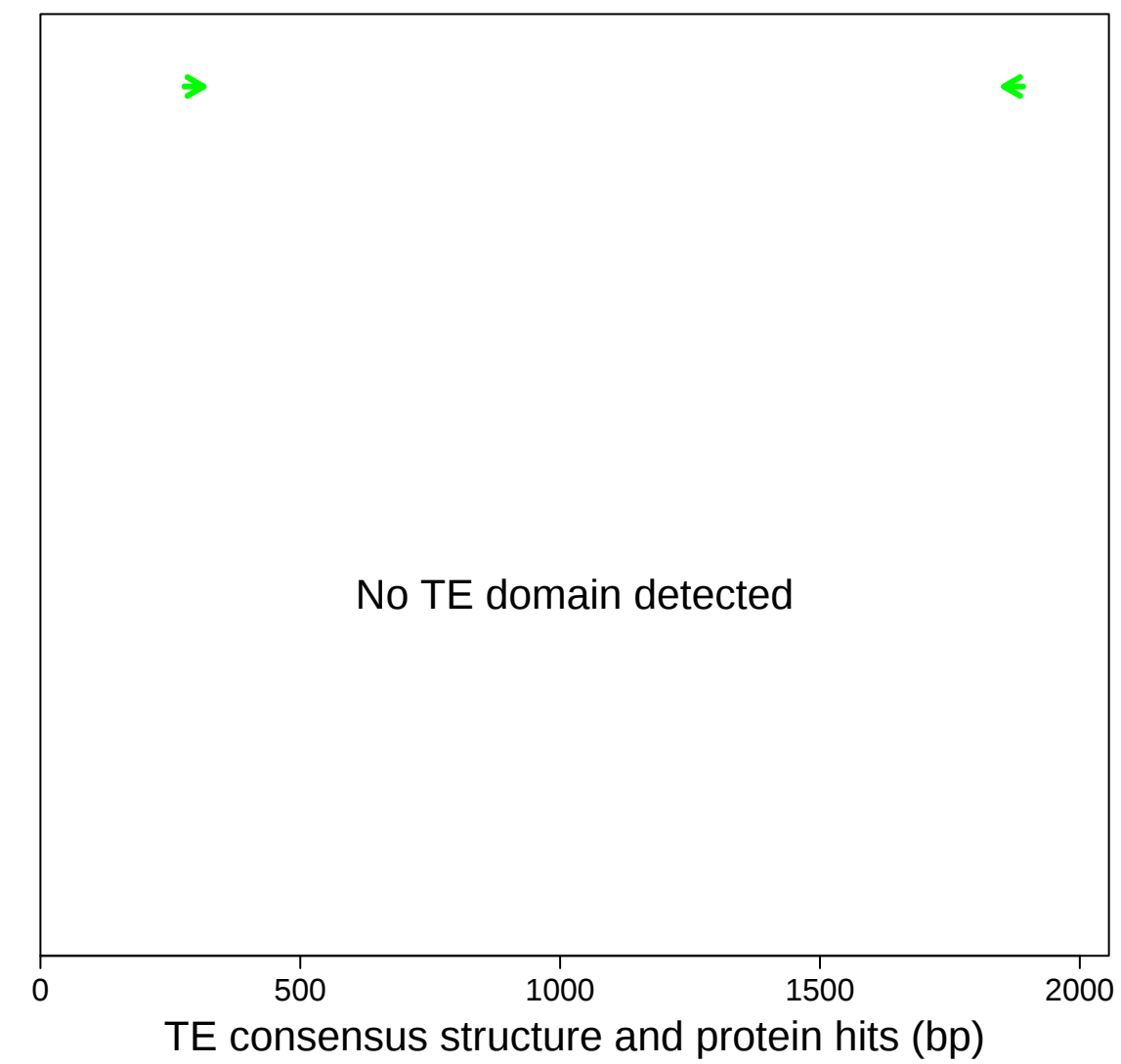
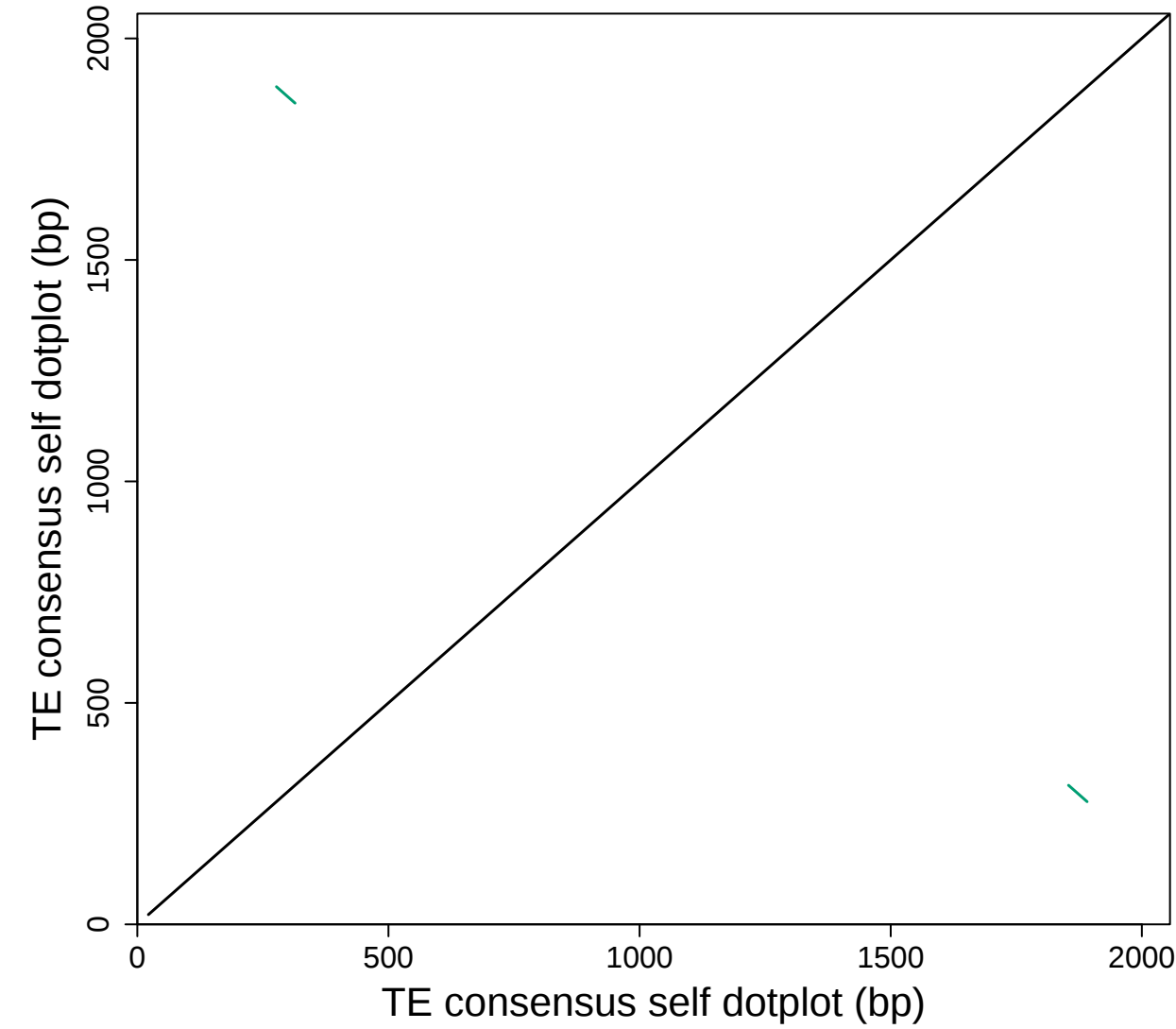
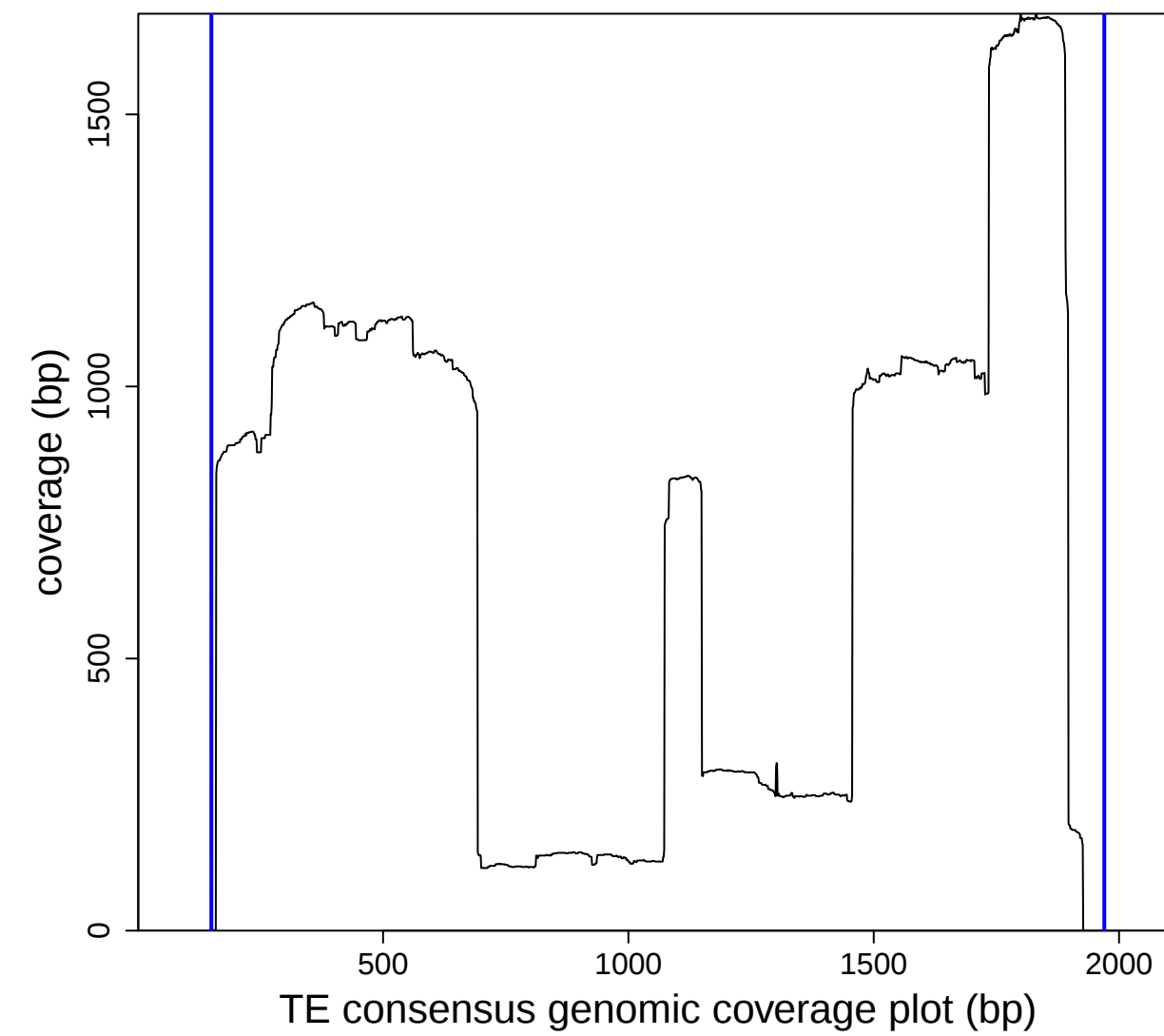
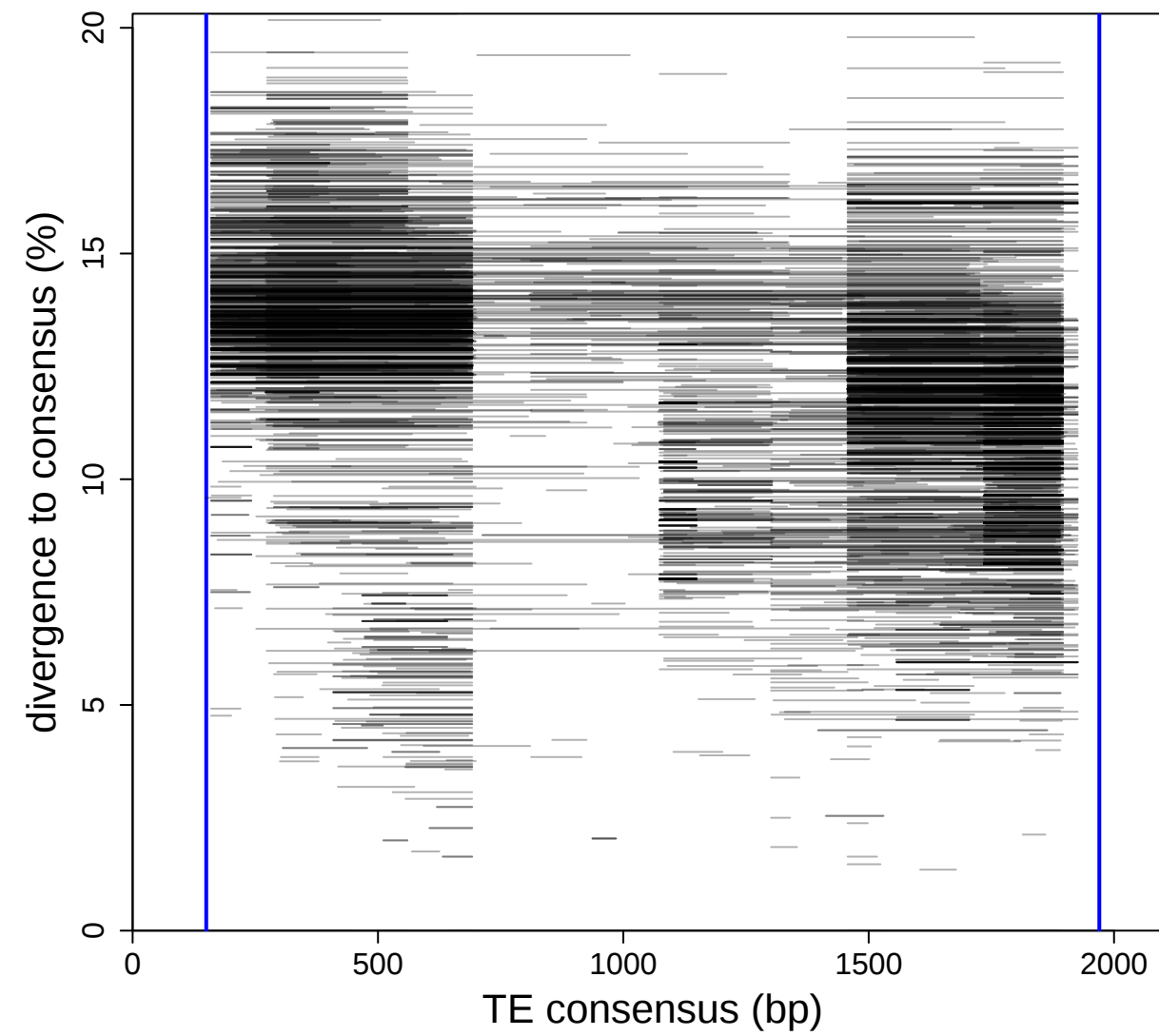
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1800bp; fragments: 4271; full length: 0 (>=1620bp)

After TEtrimmer 1800 bp

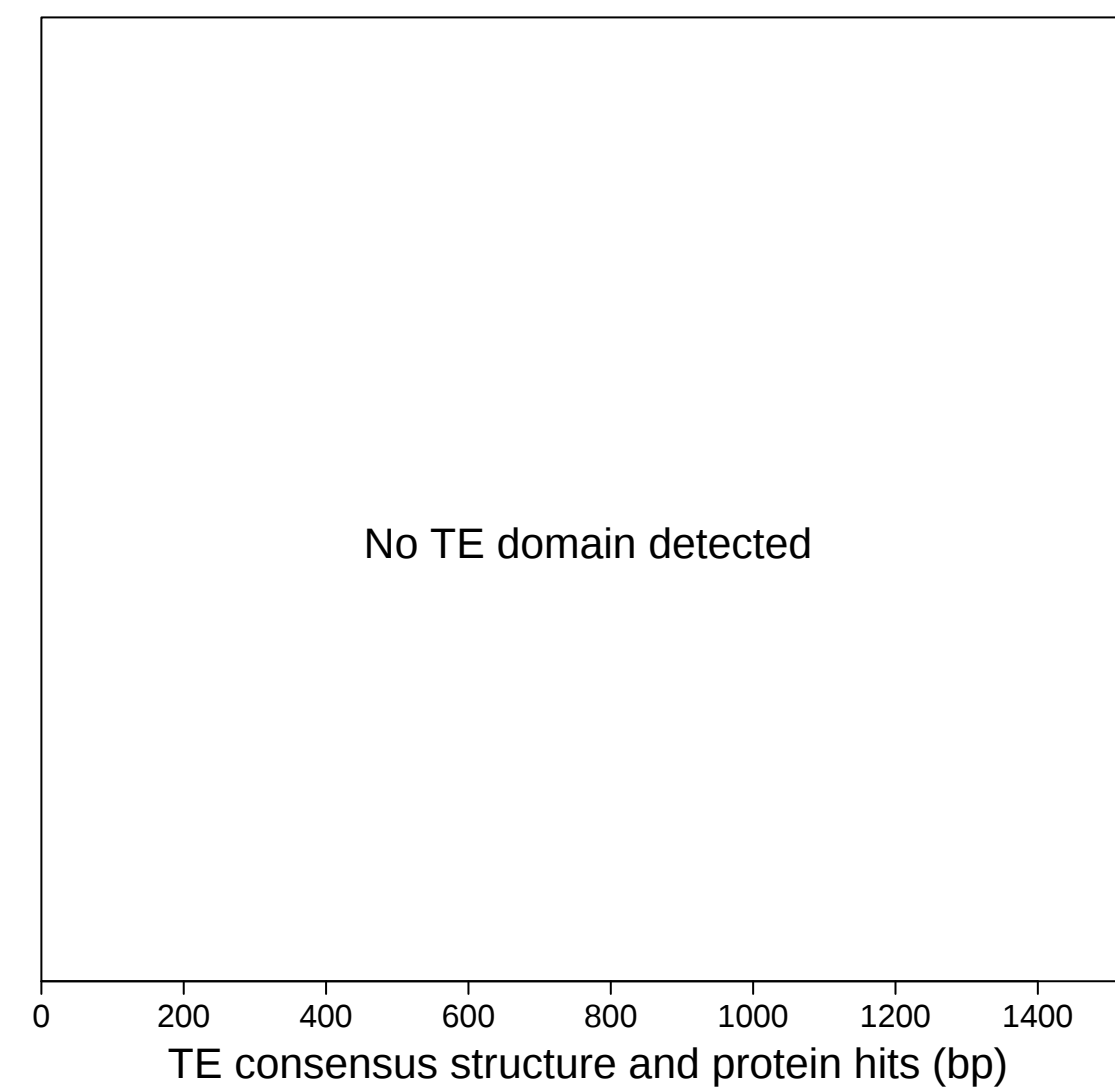
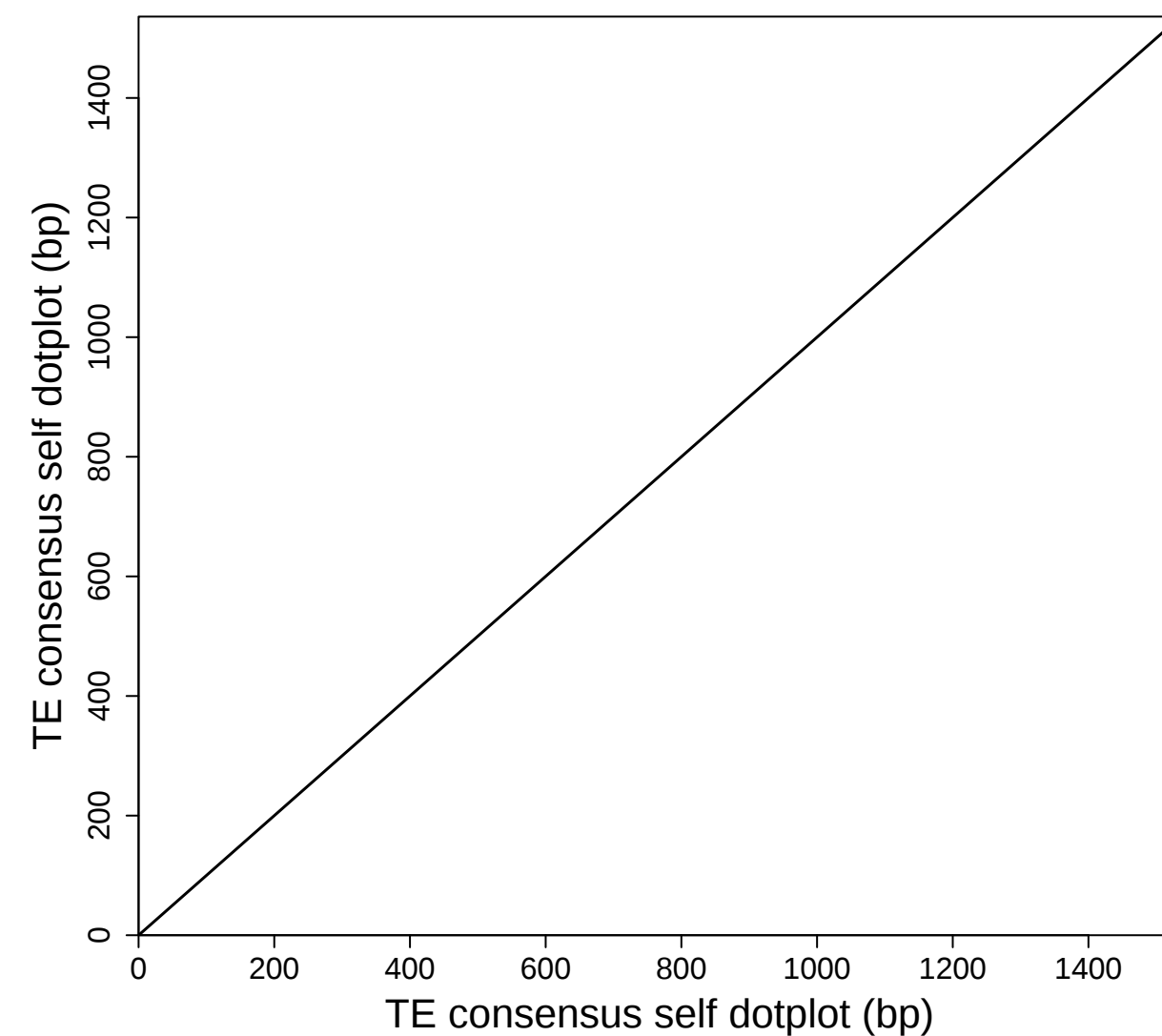
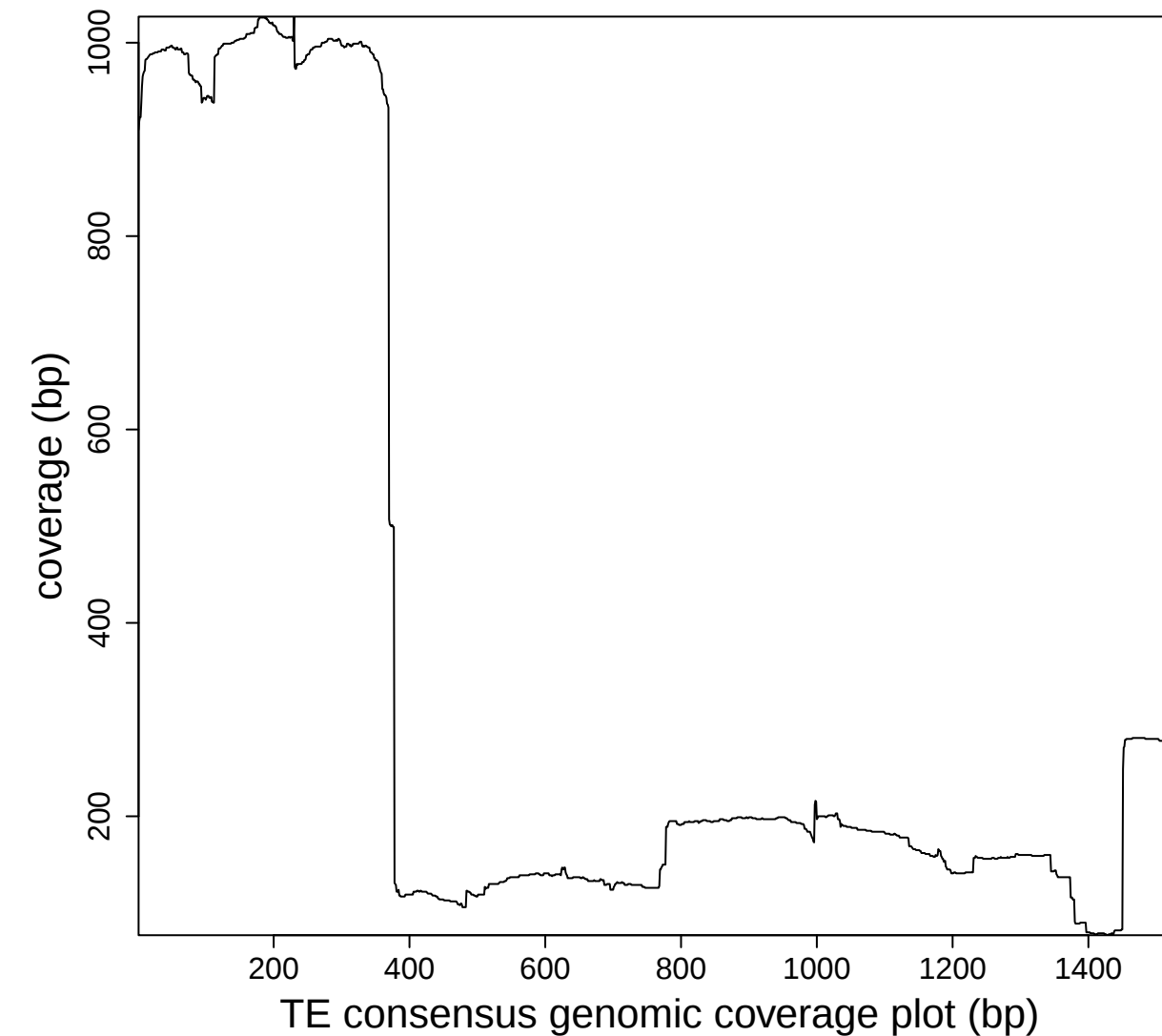
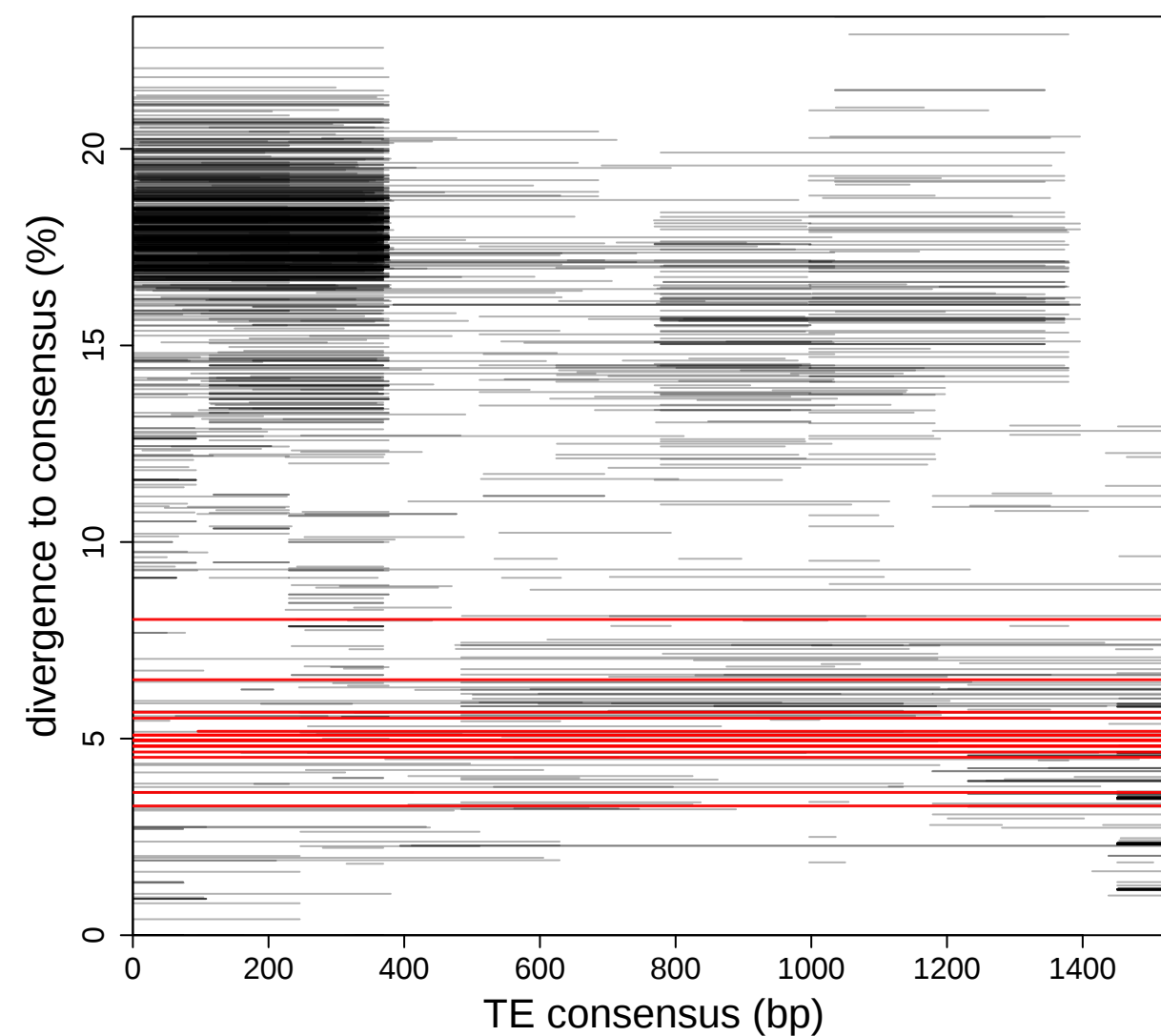


6.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 2120bp; fragments: 4116; full length: 0 (>=1908bp)

After TEtrimmer Extended plot Blue lines are boundaries

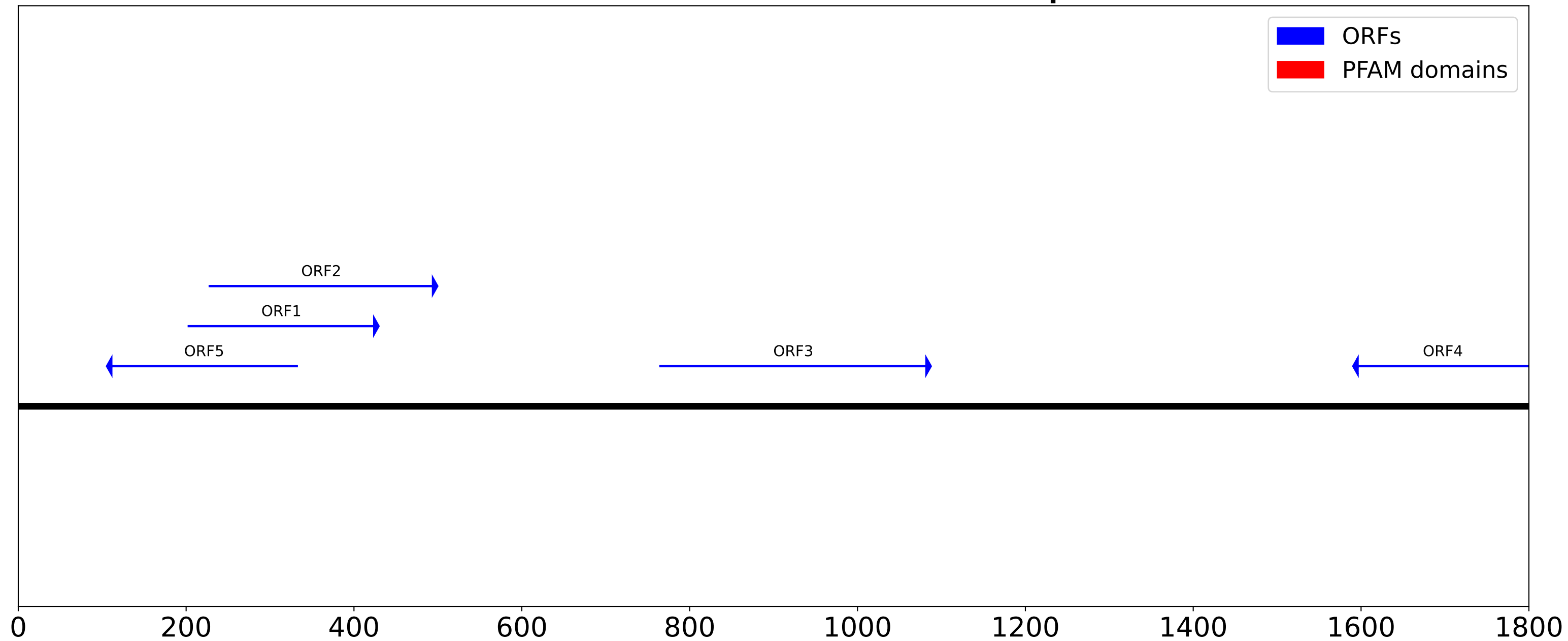


TE: rnd_5_family_5626
size: 1536bp; fragments: 1690; full length: 16 (≥ 1382.4 bp)

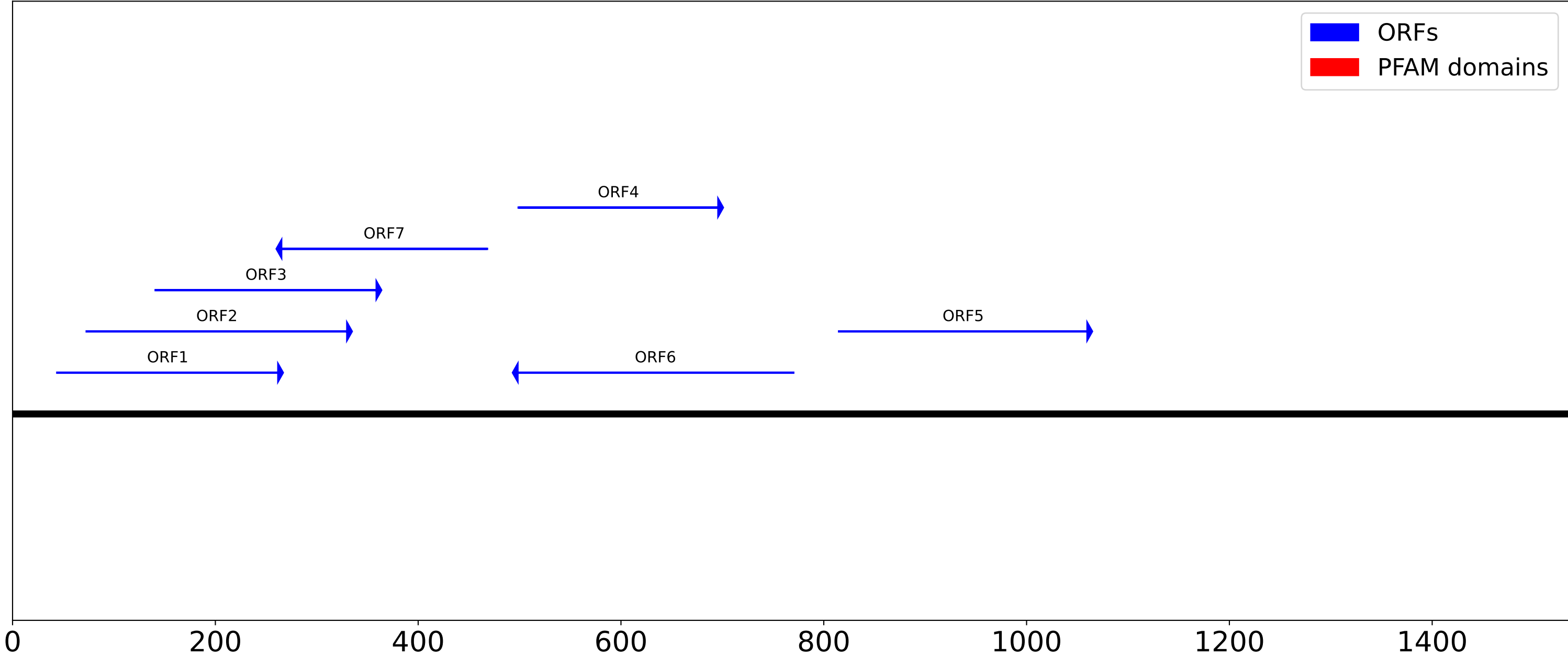


Before TEtrimmer 1536 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

