

MSA length = 1667

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A--CGACCAGAAATTTAATCATTGA--ATT--CT--GAAC-CAAAGTTACGTTTTCATAATGGAAATACT--
CTCTCGTCTCCTTTTGACTC-GTGCTTGACCGCCTTC-AGAC--ACGCGAGTGGCGGGCTCGTCAGGGTAGACAAGTTAGGGCGGGCGCCT--

Start crop Point

End crop Point

0 -

10 -

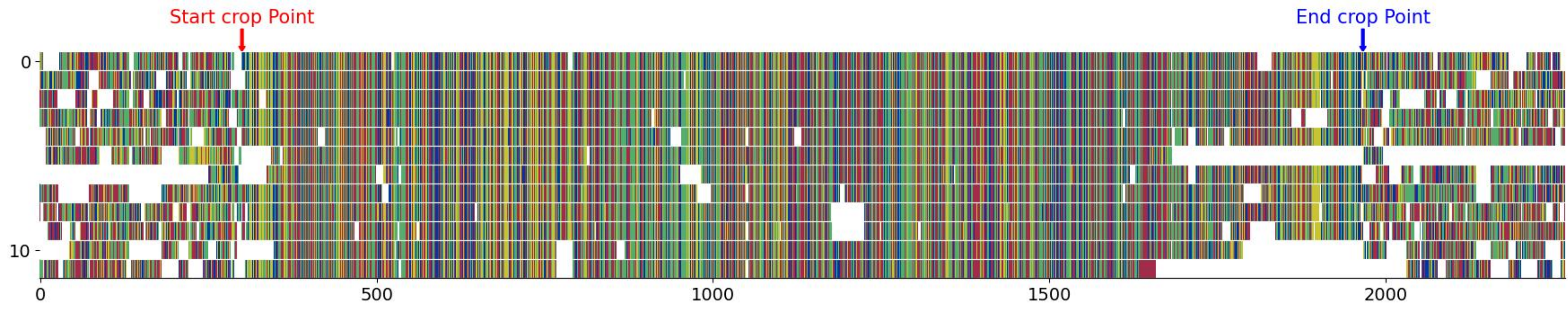
0

500

1000

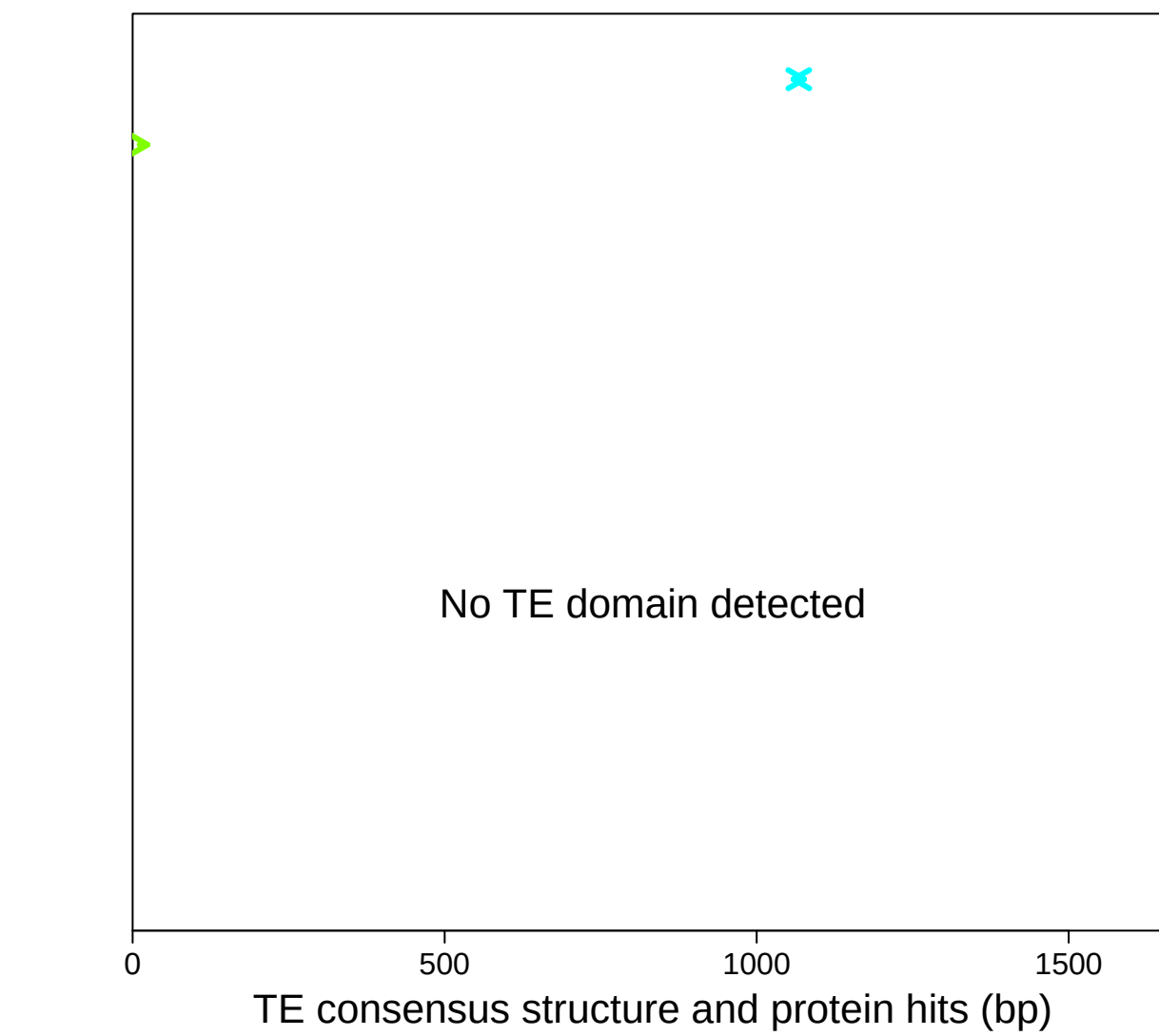
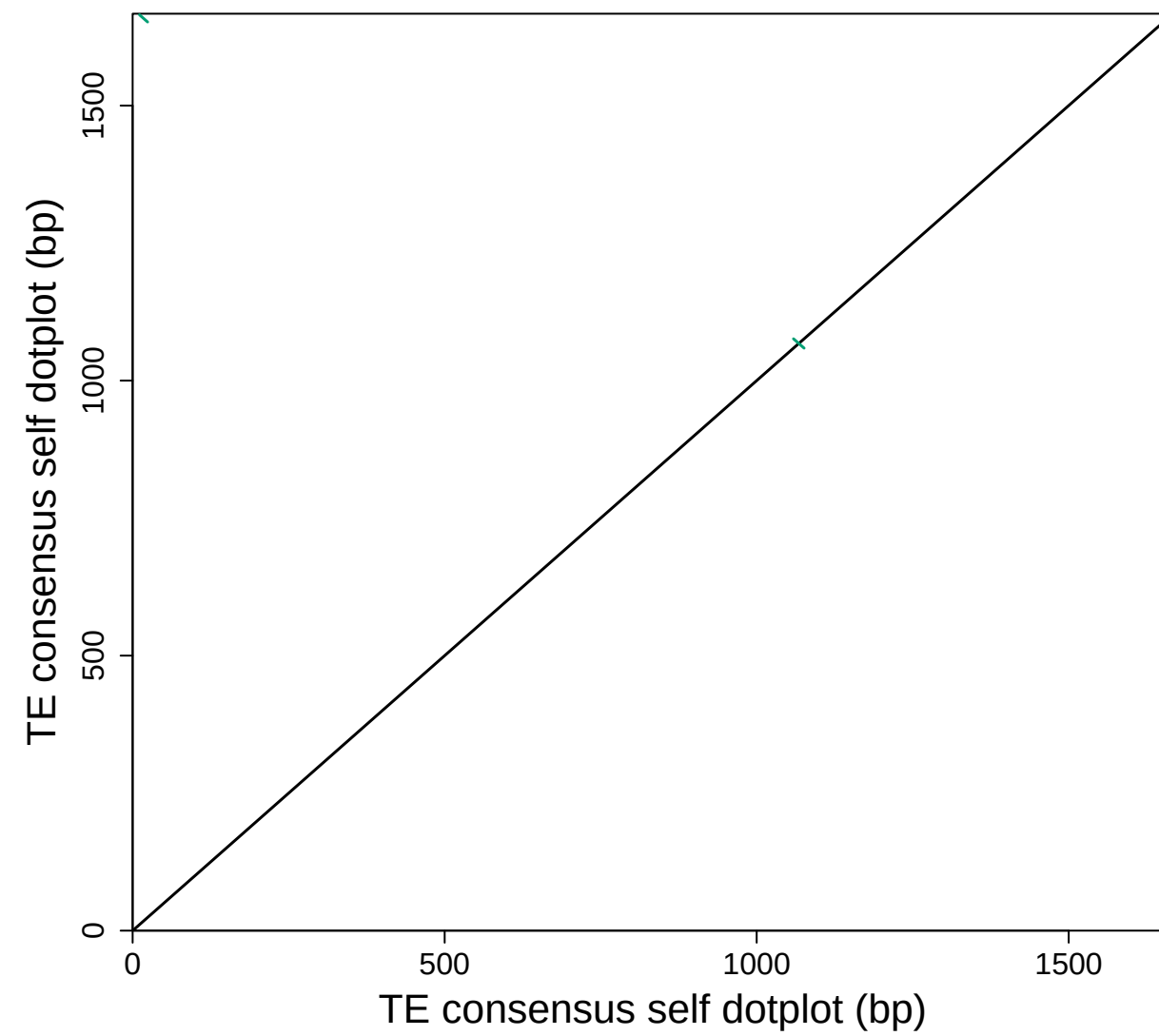
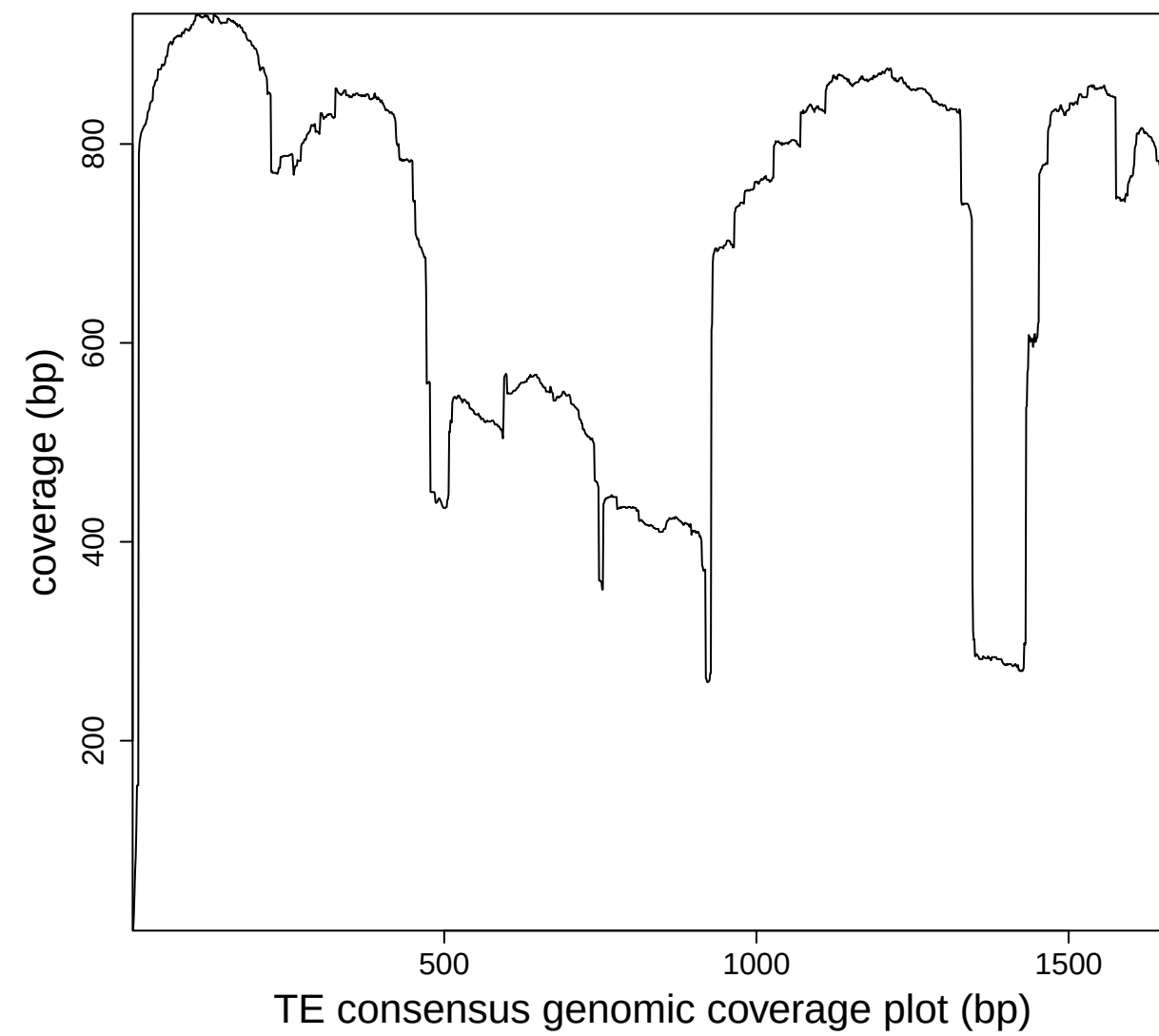
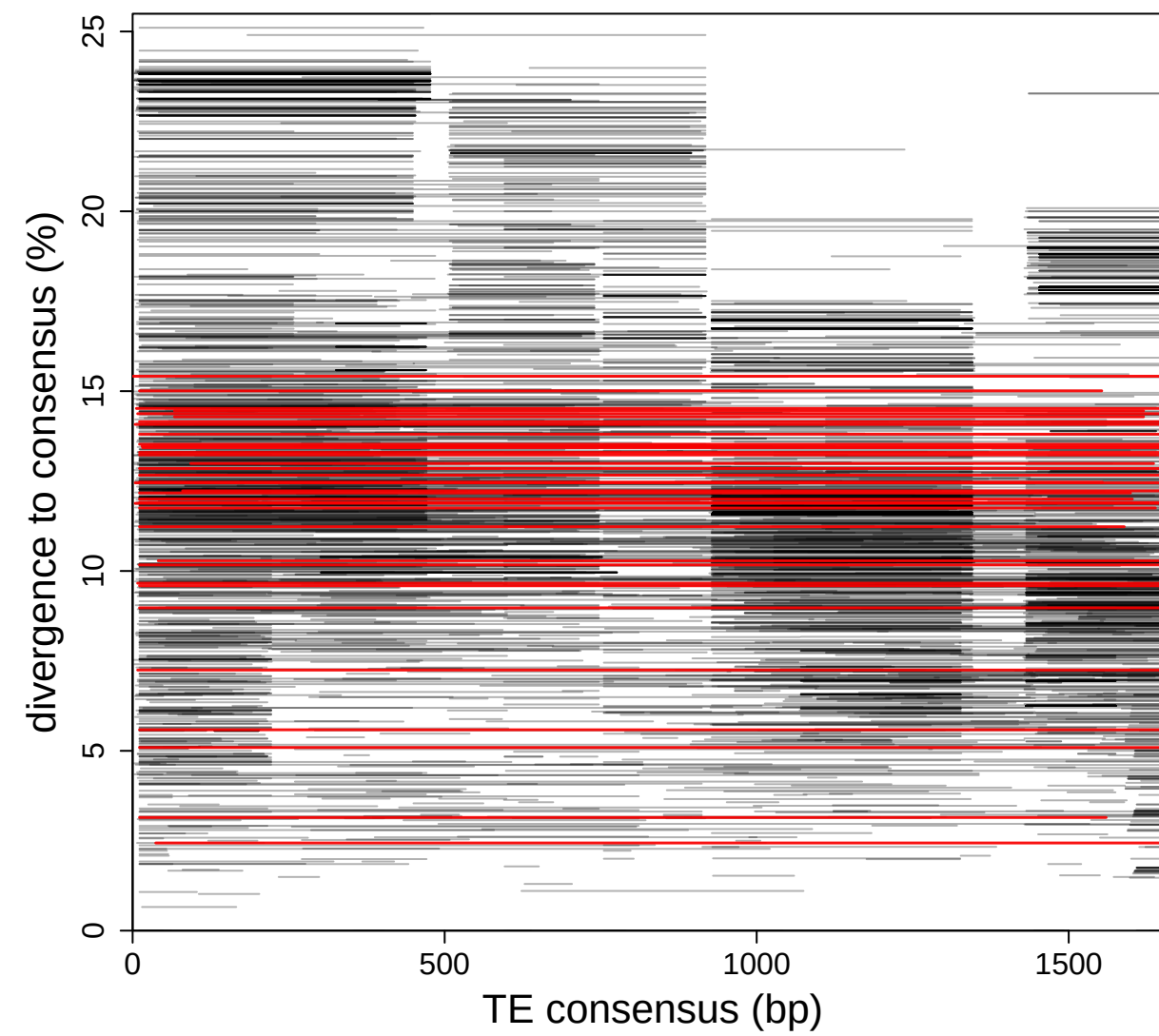
1500

2000



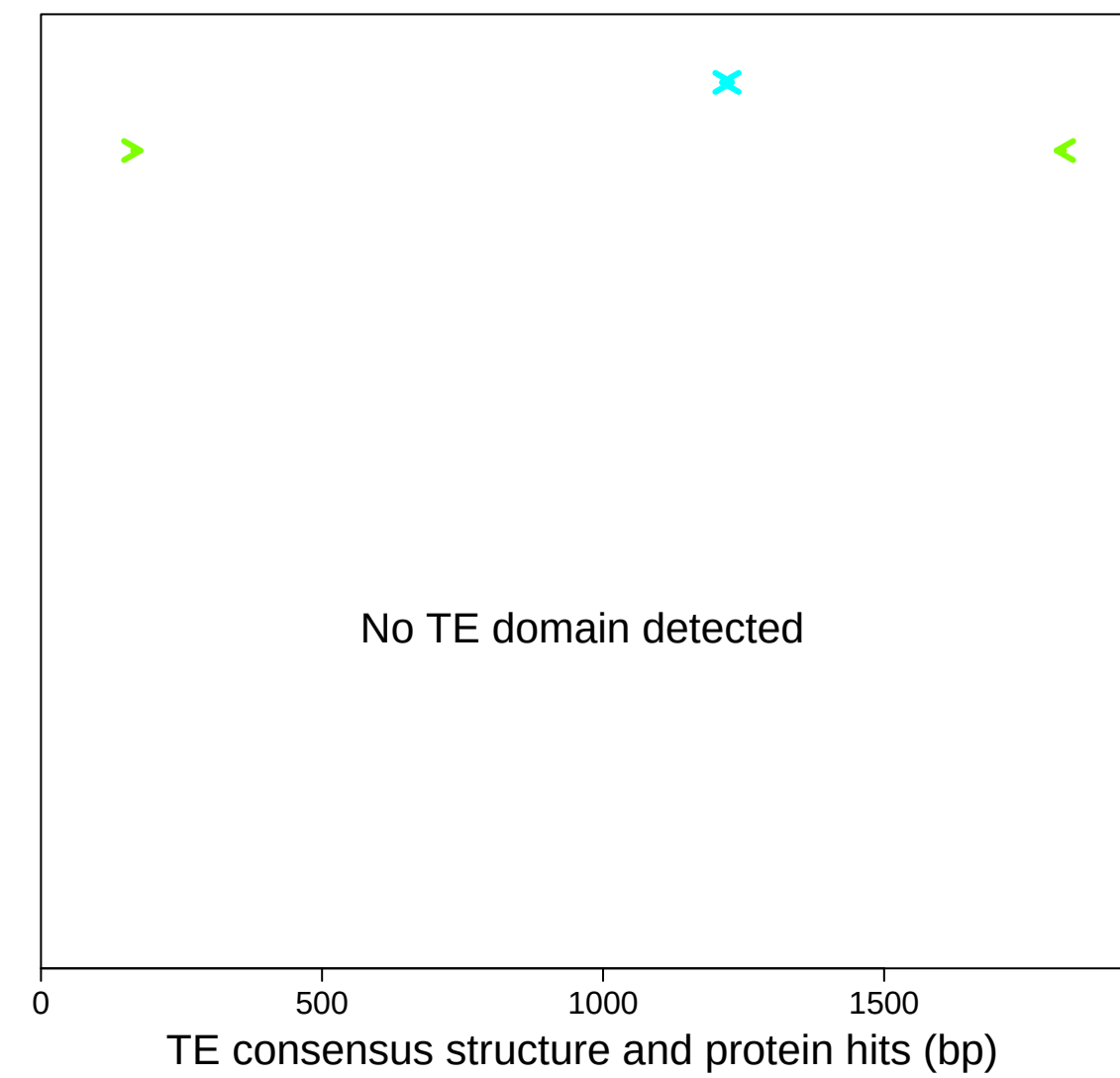
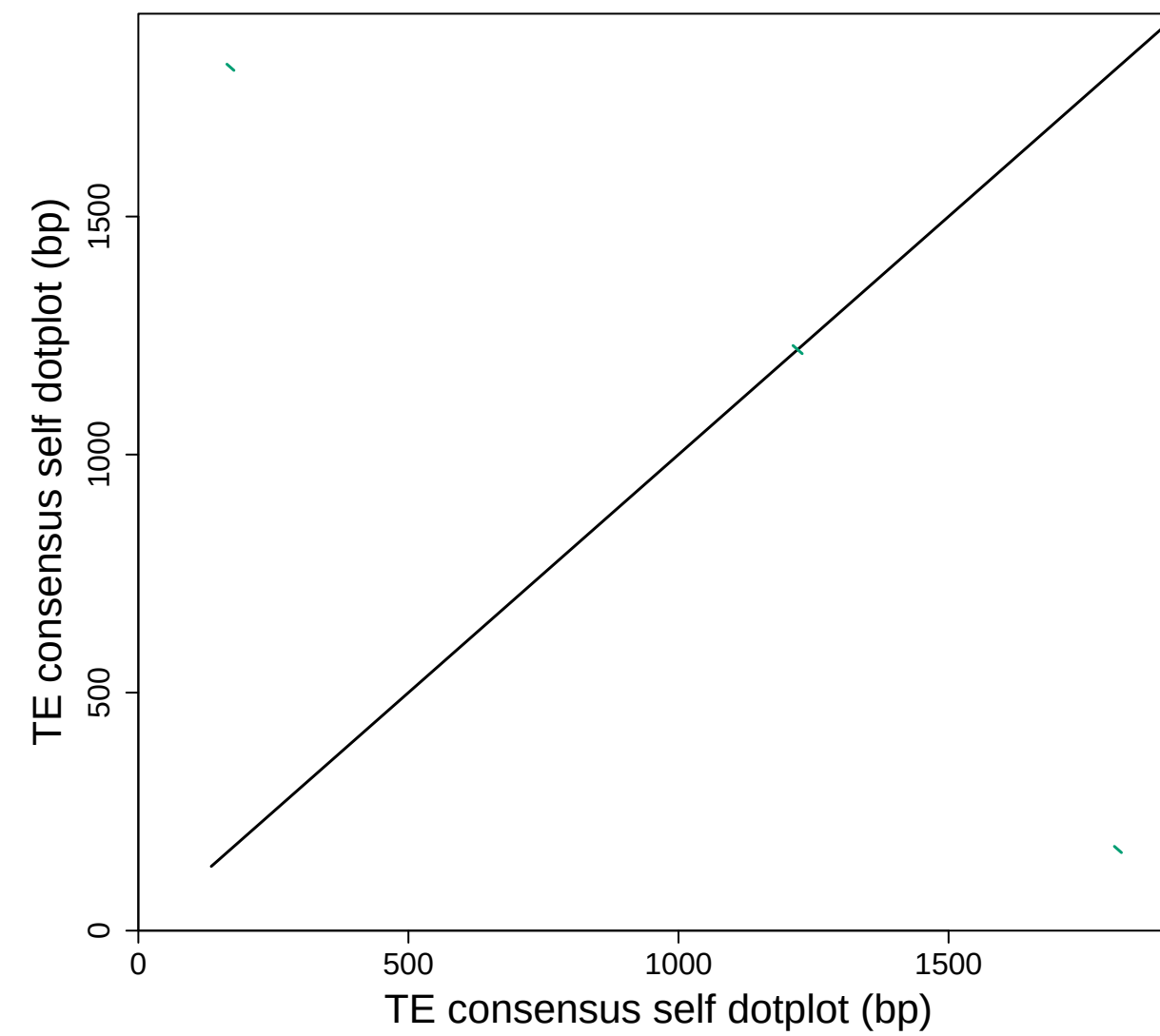
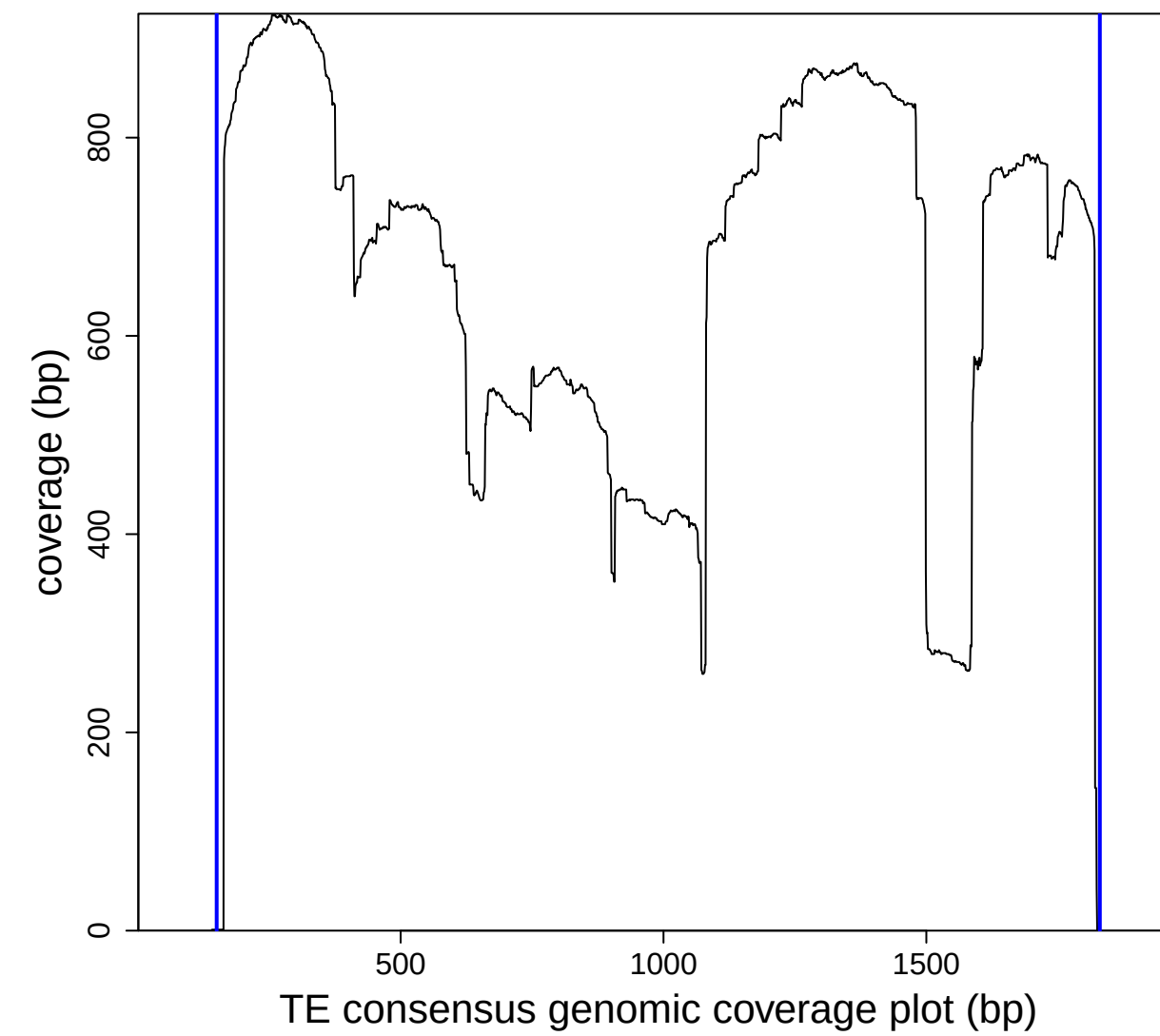
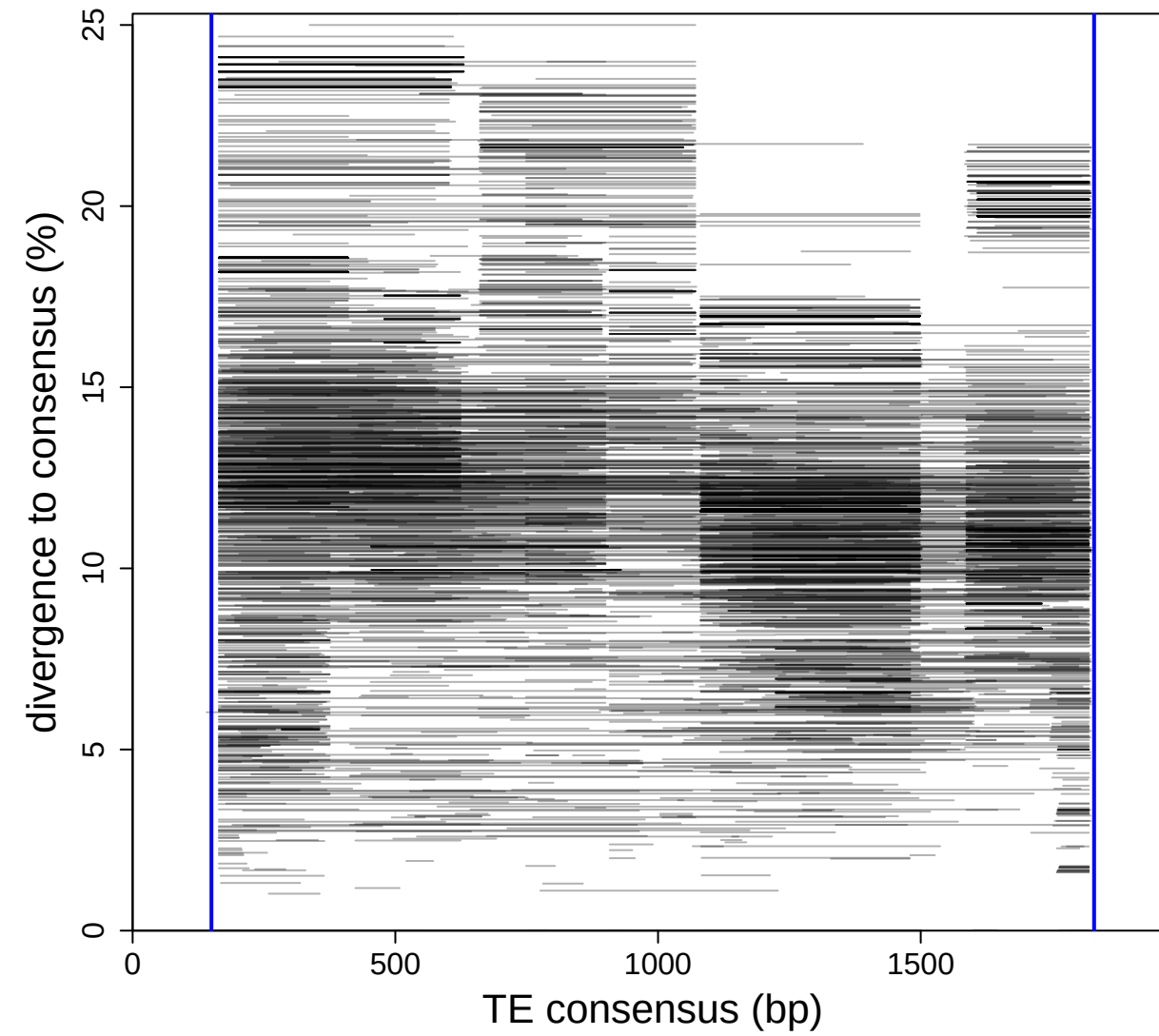
b.bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 1667bp; fragments: 3262; full length: 36 (≥ 1500.3 bp)

After TEtrimmer 1667 bp

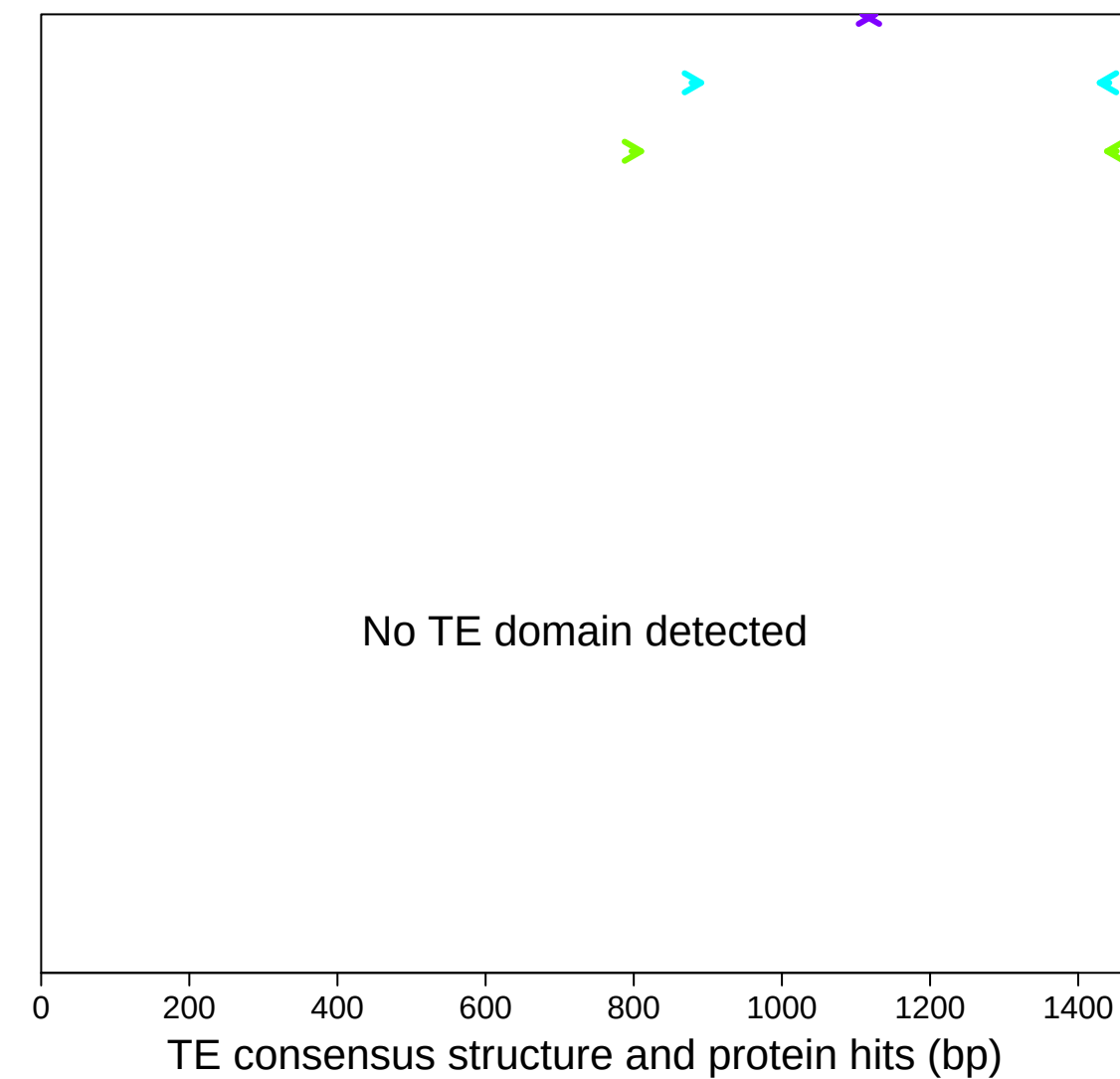
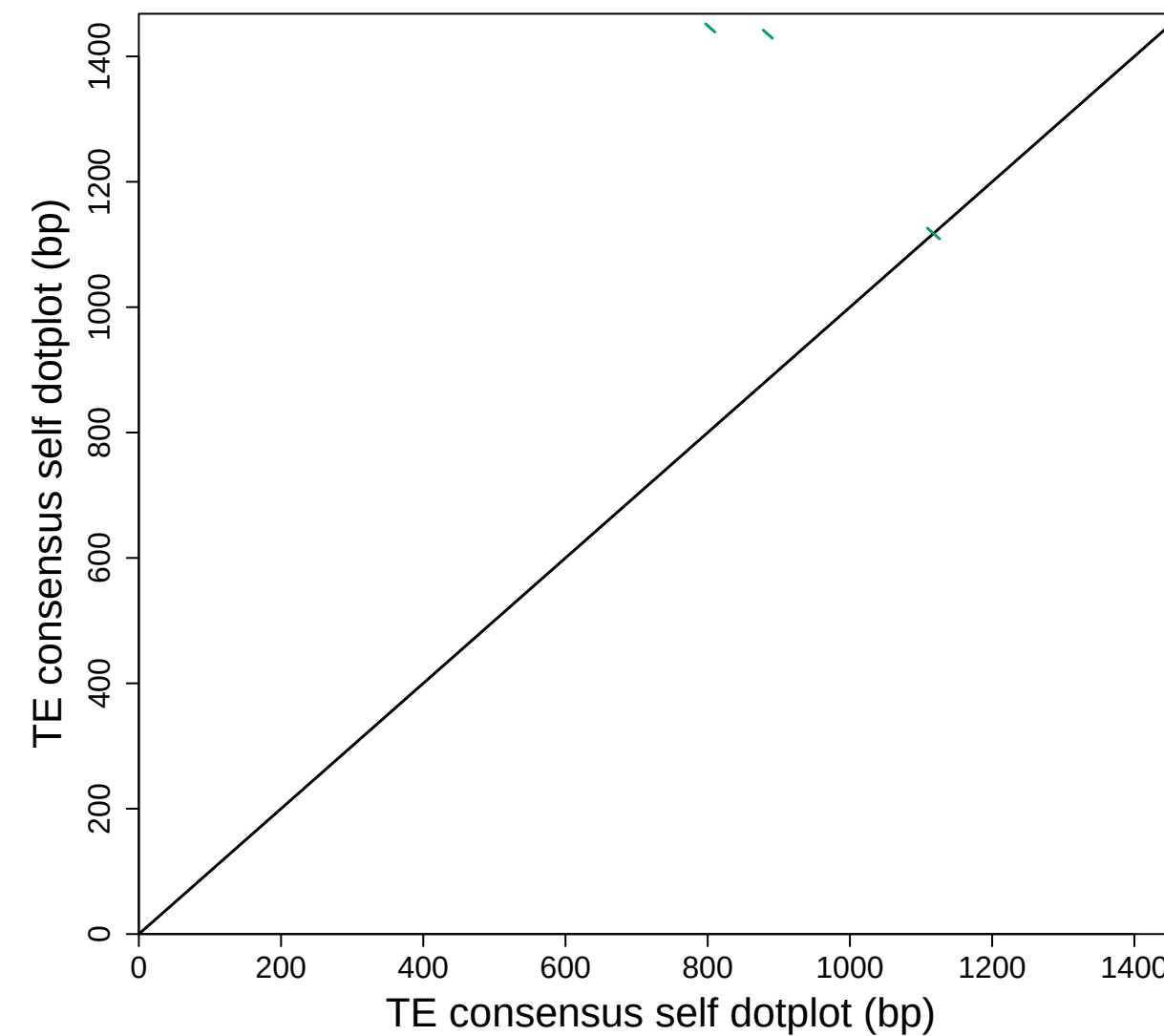
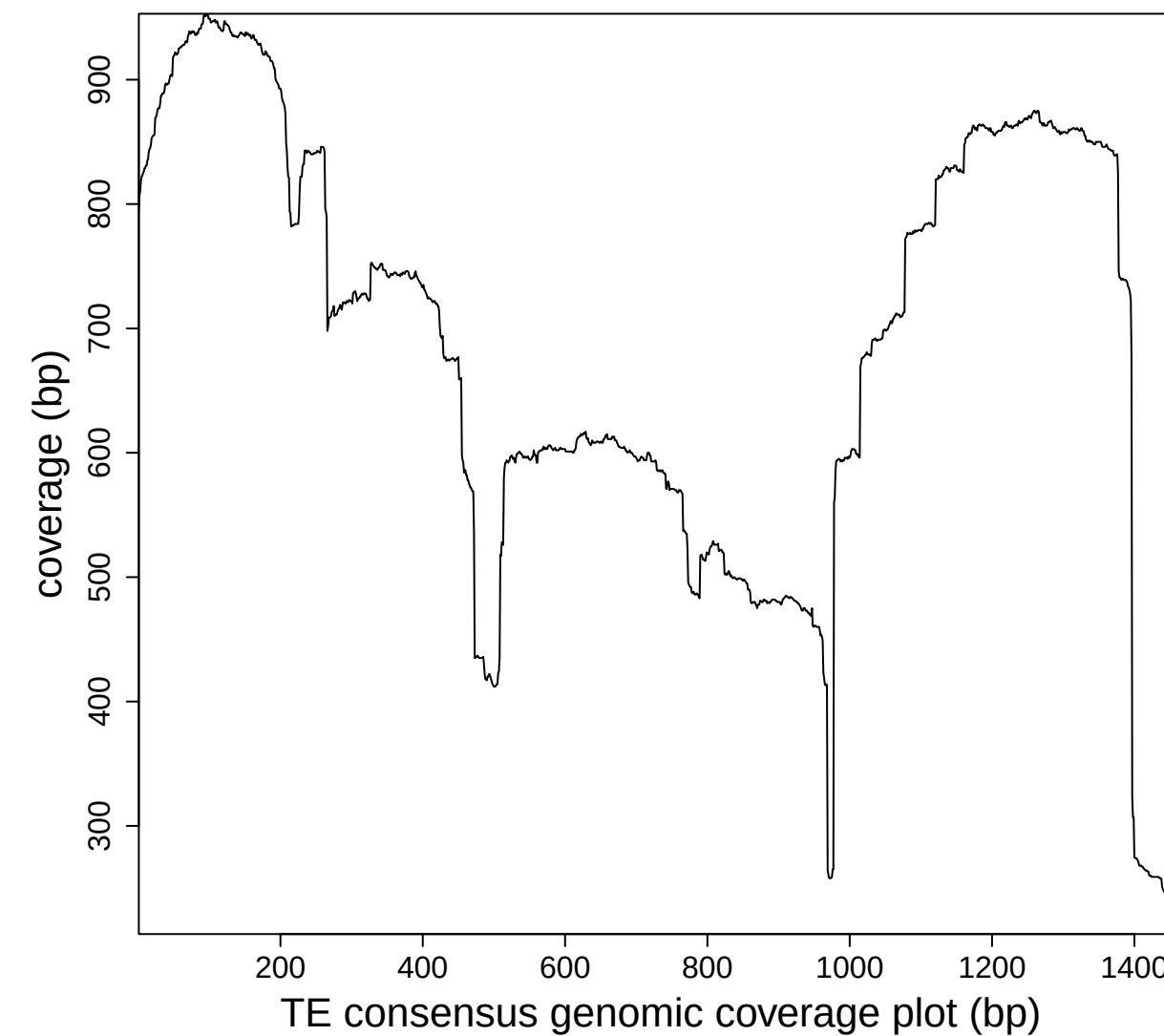
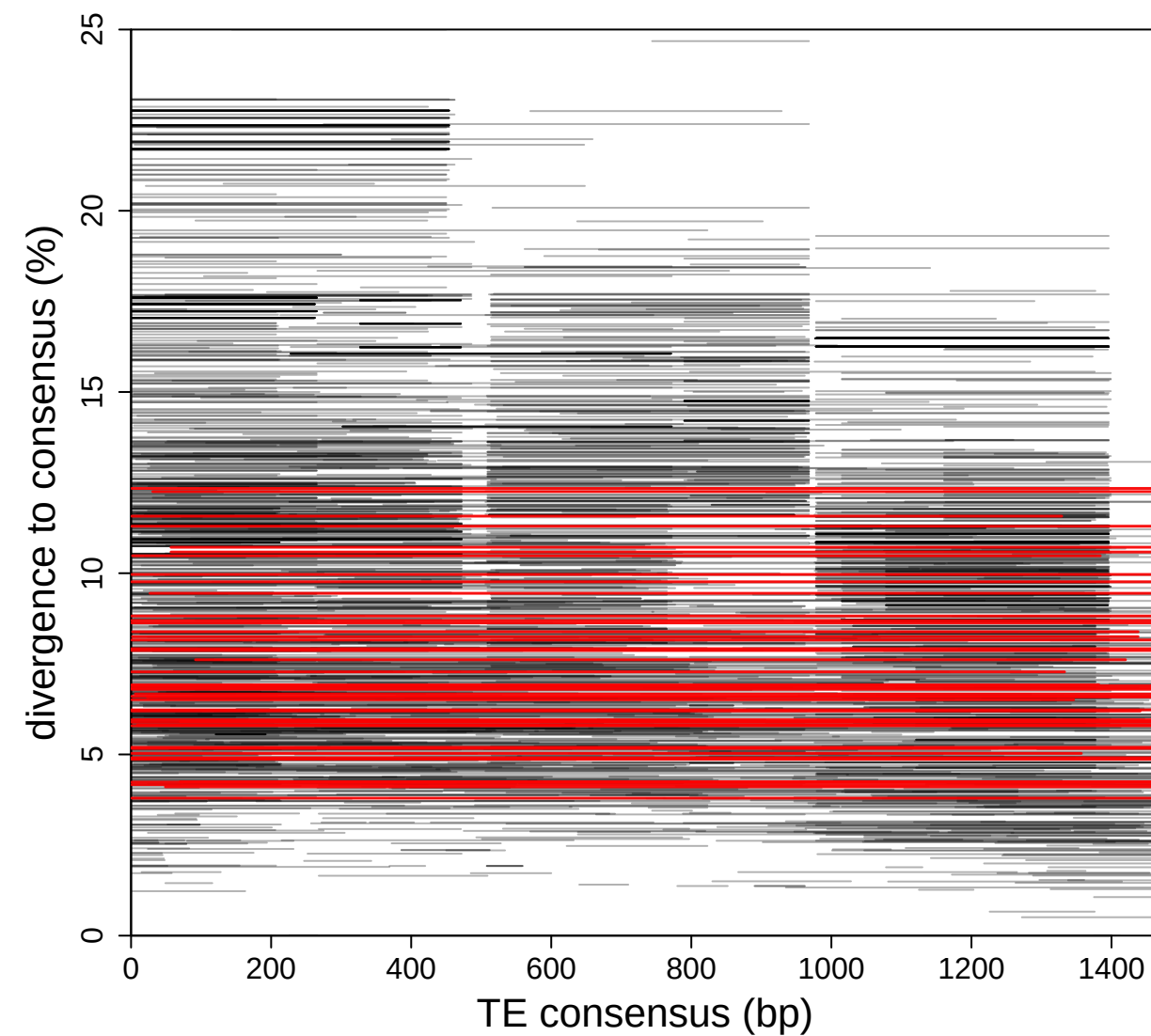


.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1980bp; fragments: 3171; full length: 0 (>=1782bp)

After TEtrimmer Extended plot Blue lines are boundaries

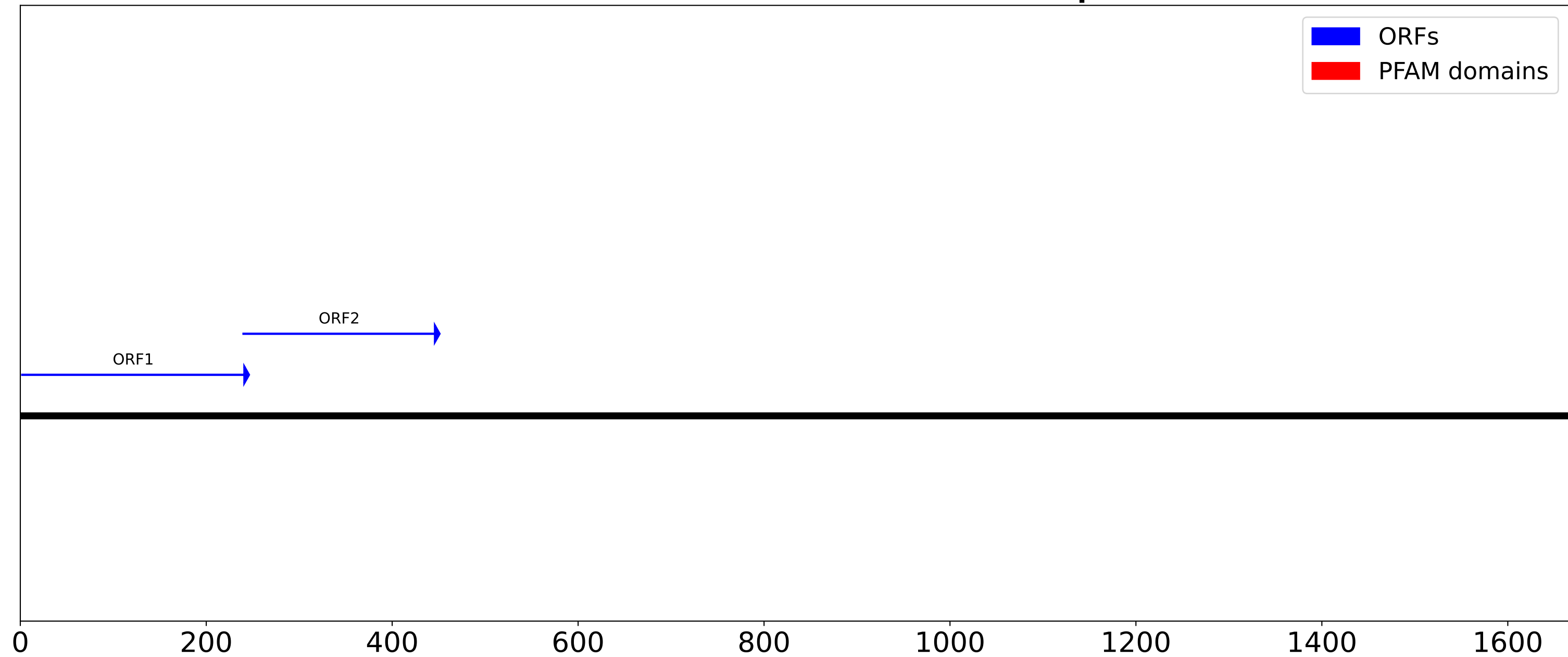


TE: rnd_5_family_79
size: 1468bp; fragments: 2613; full length: 45 (≥ 1321.2 bp)

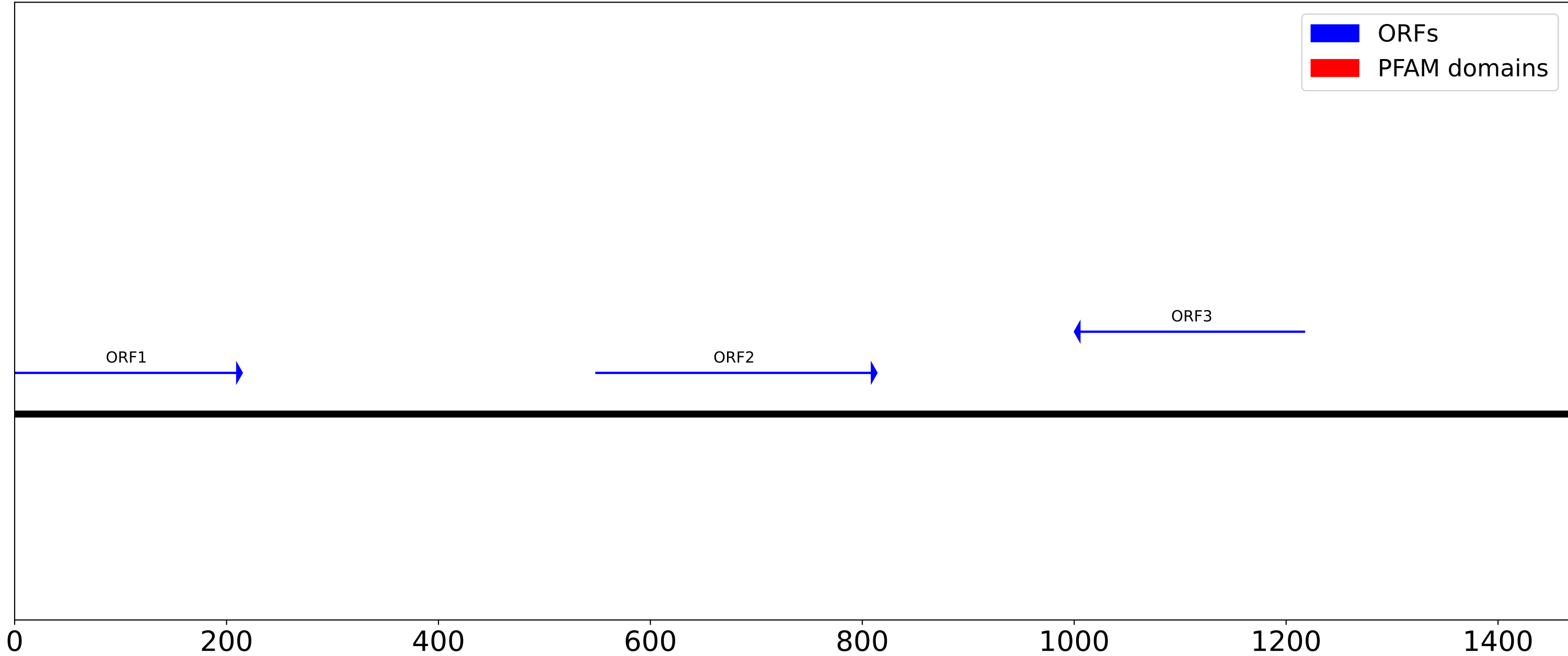


Before TETrimmer 1468 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

