

Start crop Point

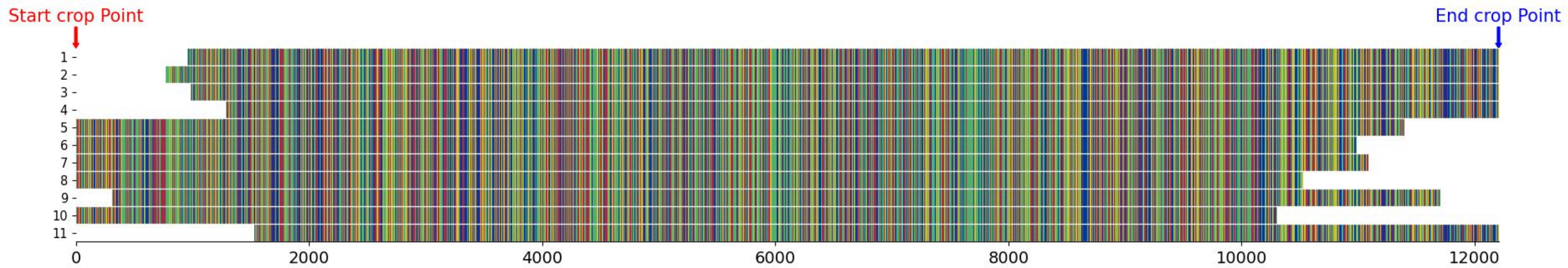


MSA length = 12207

End crop Point

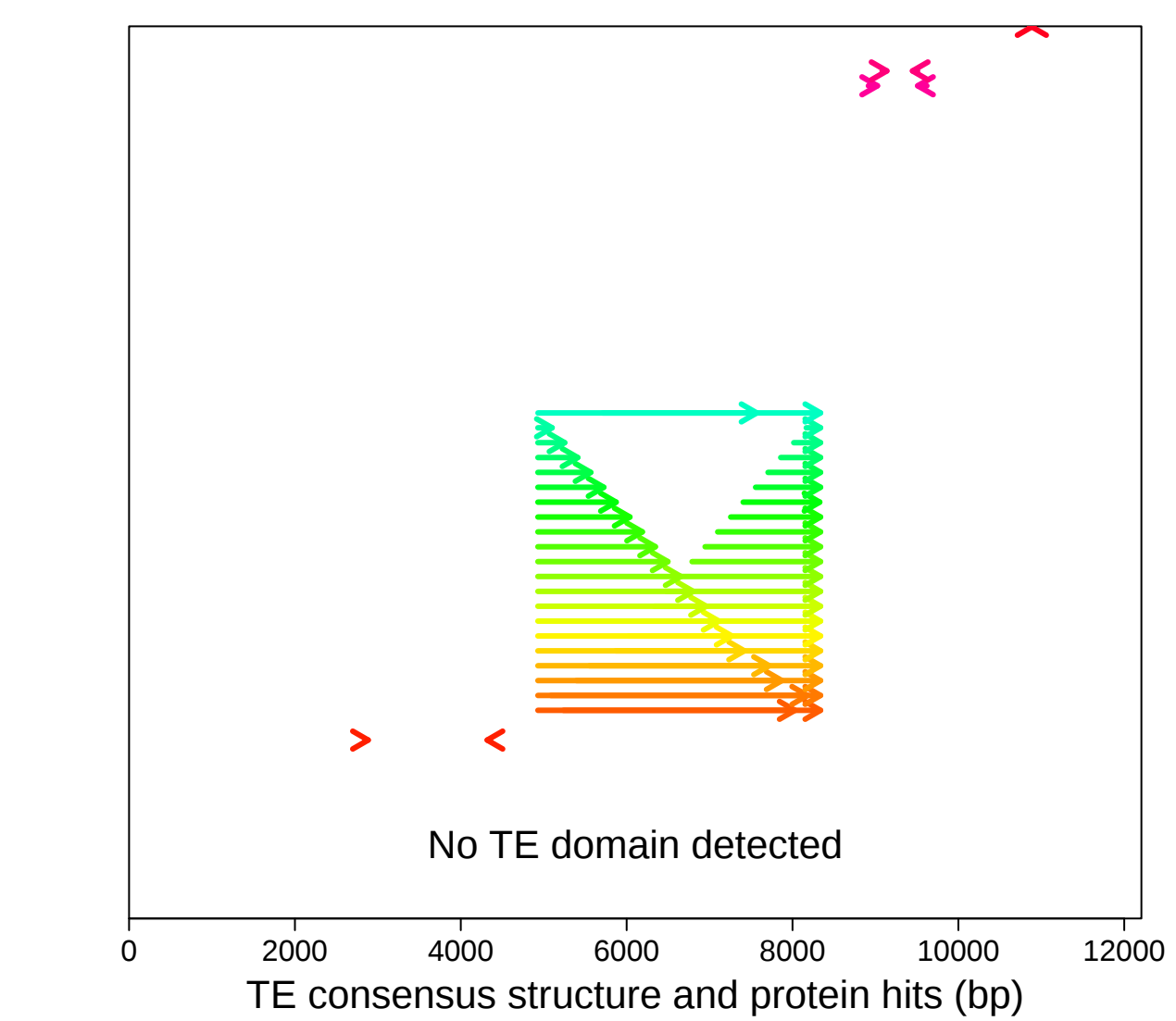
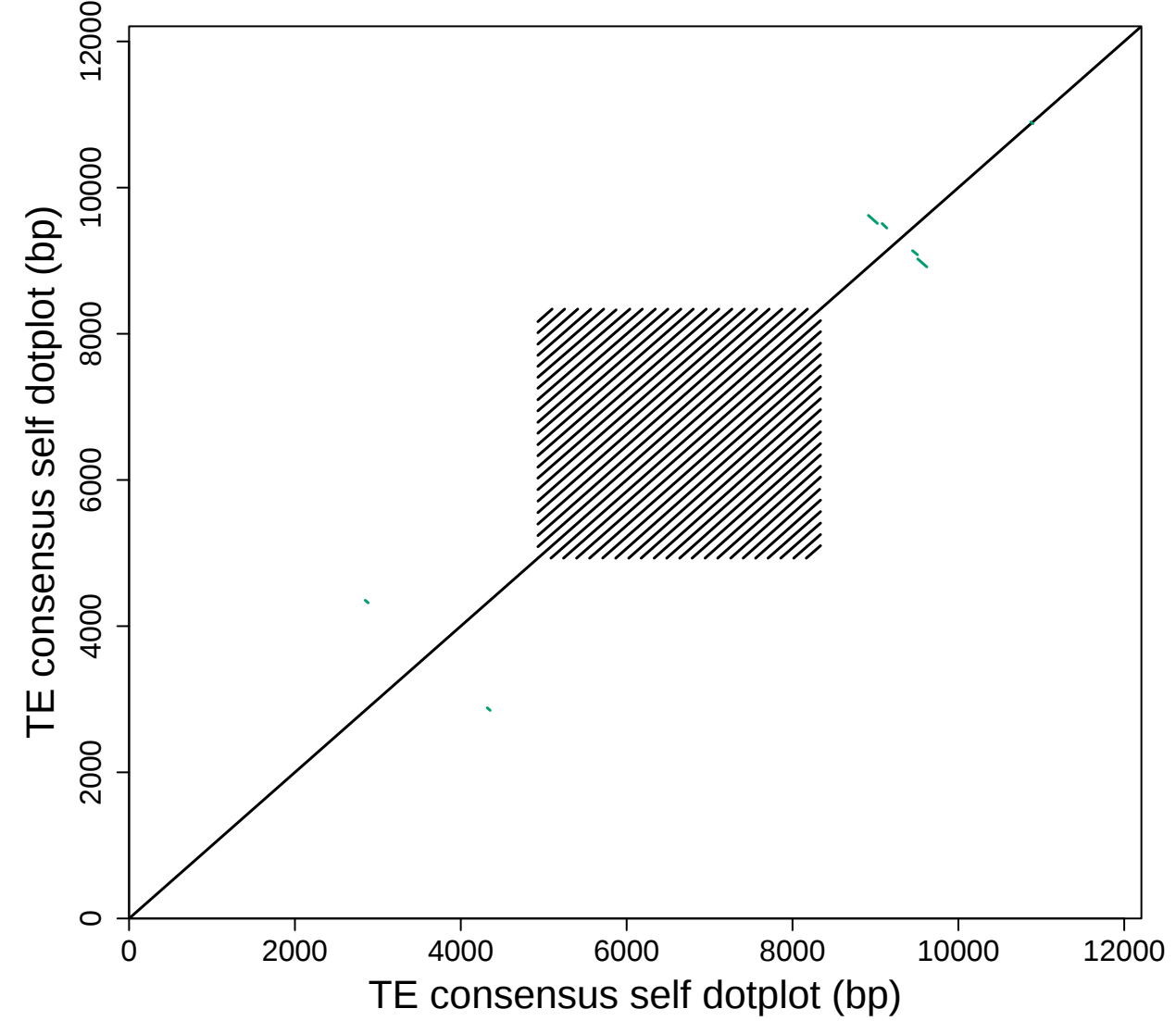
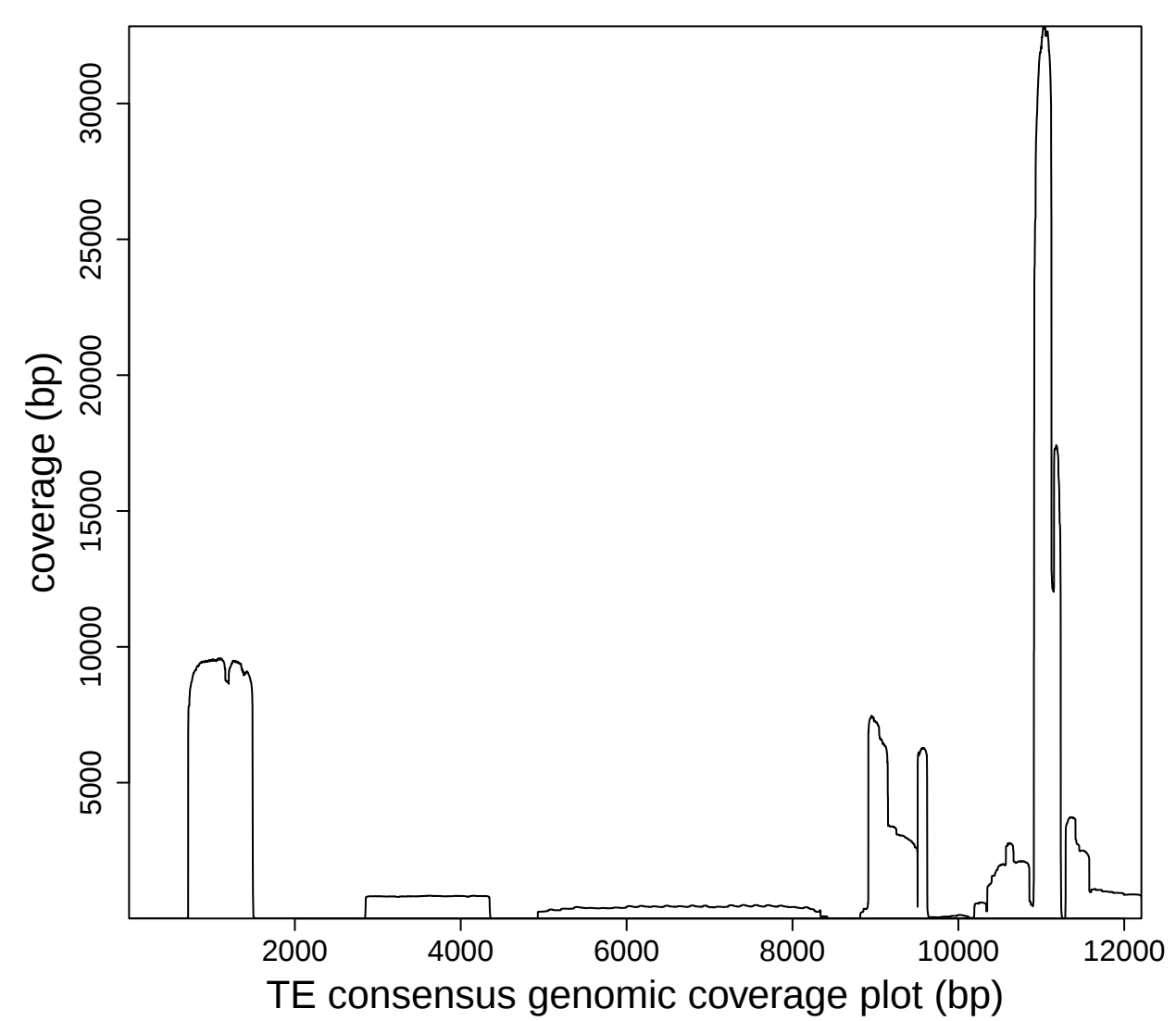
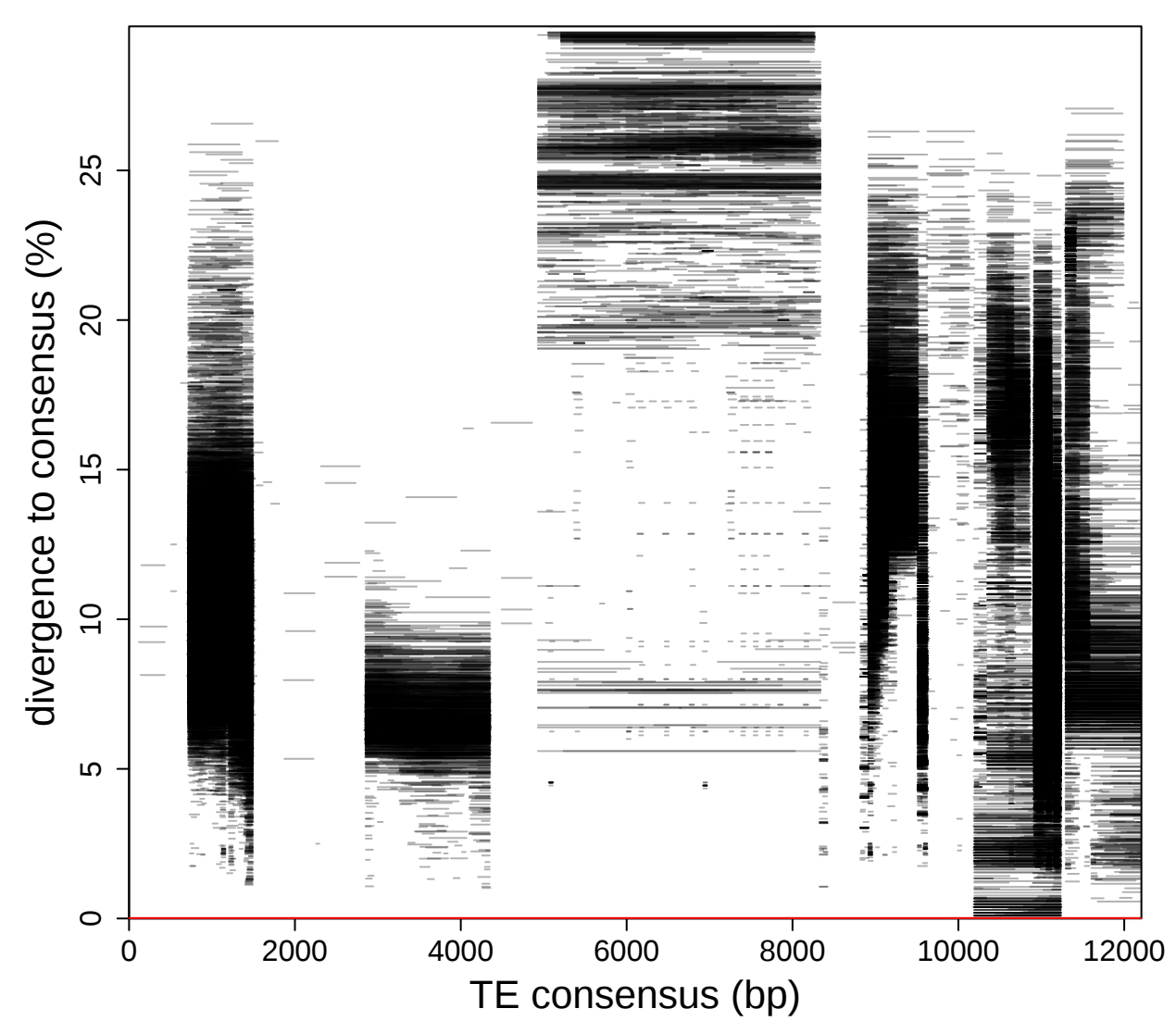


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-----TTCCTAATTACTATTTTCTTCTTCACATTTGCATTTGCTTAATACCGTGC
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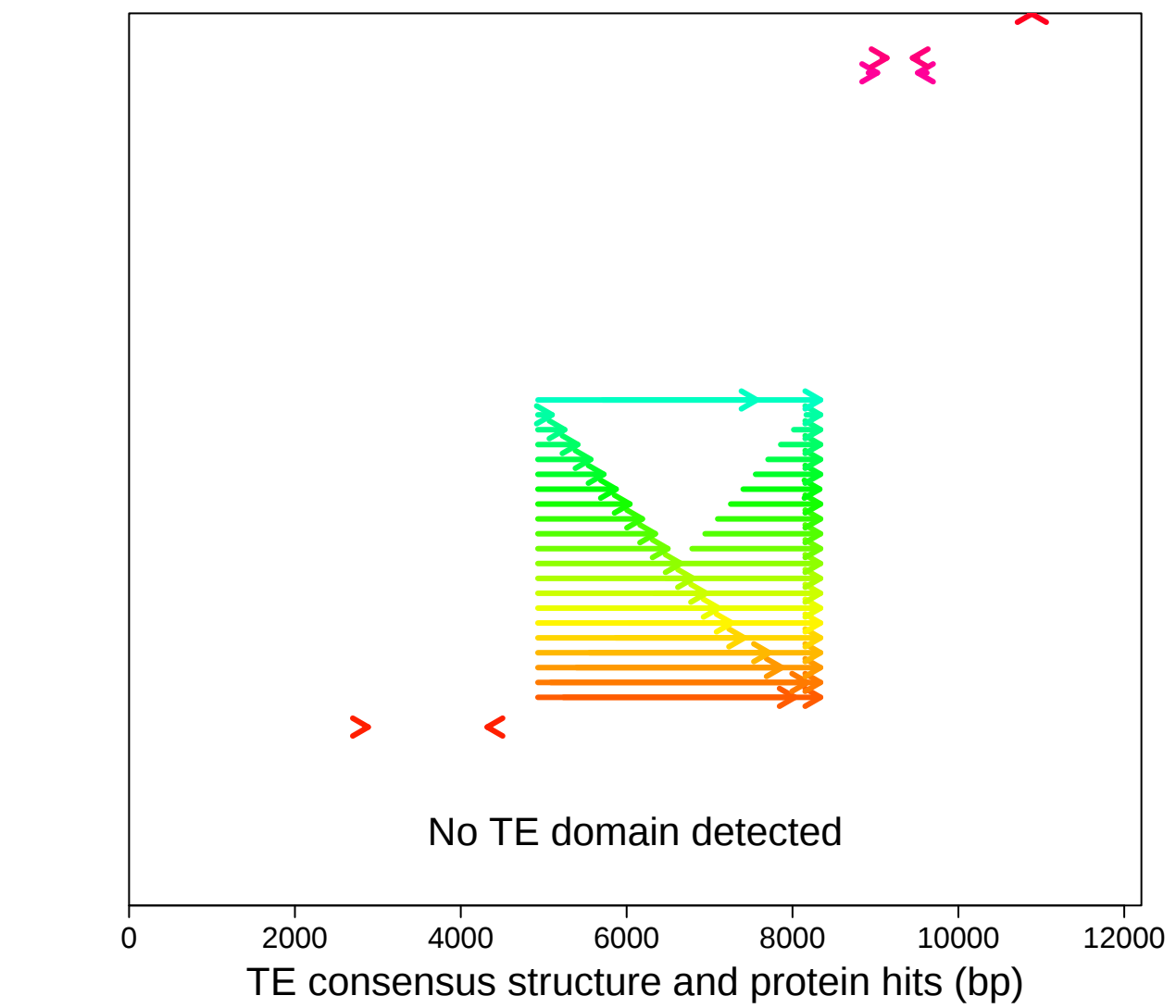
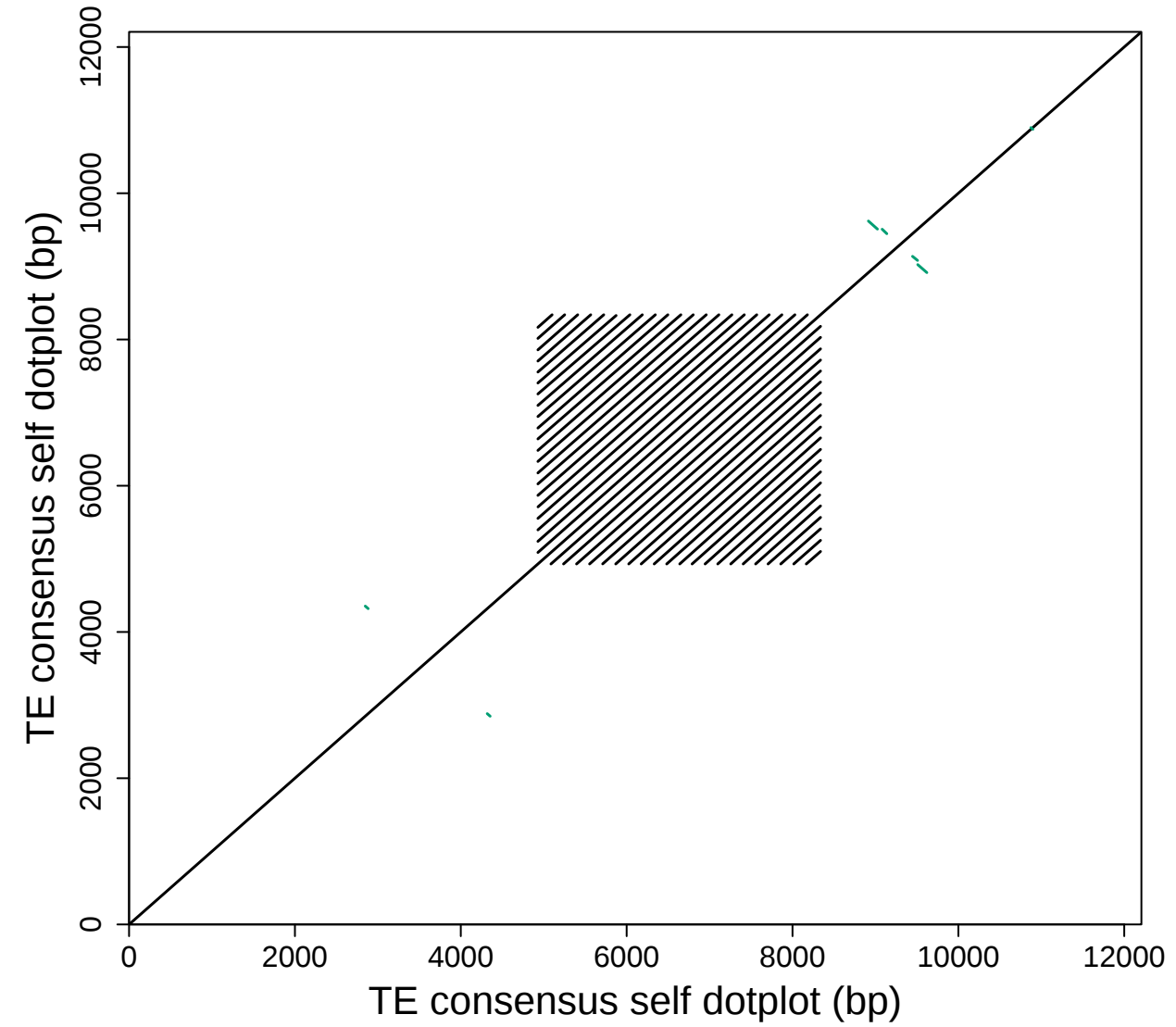
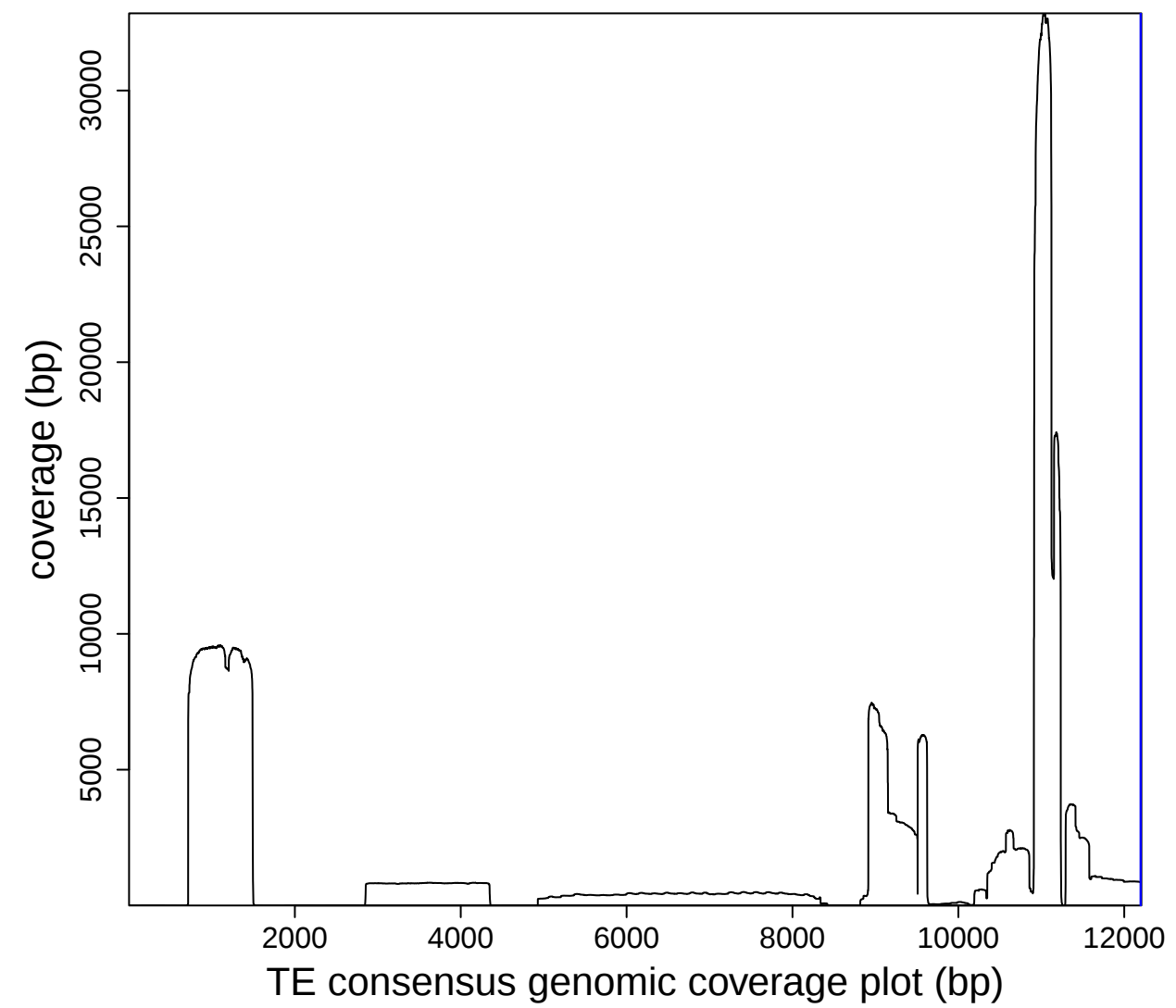
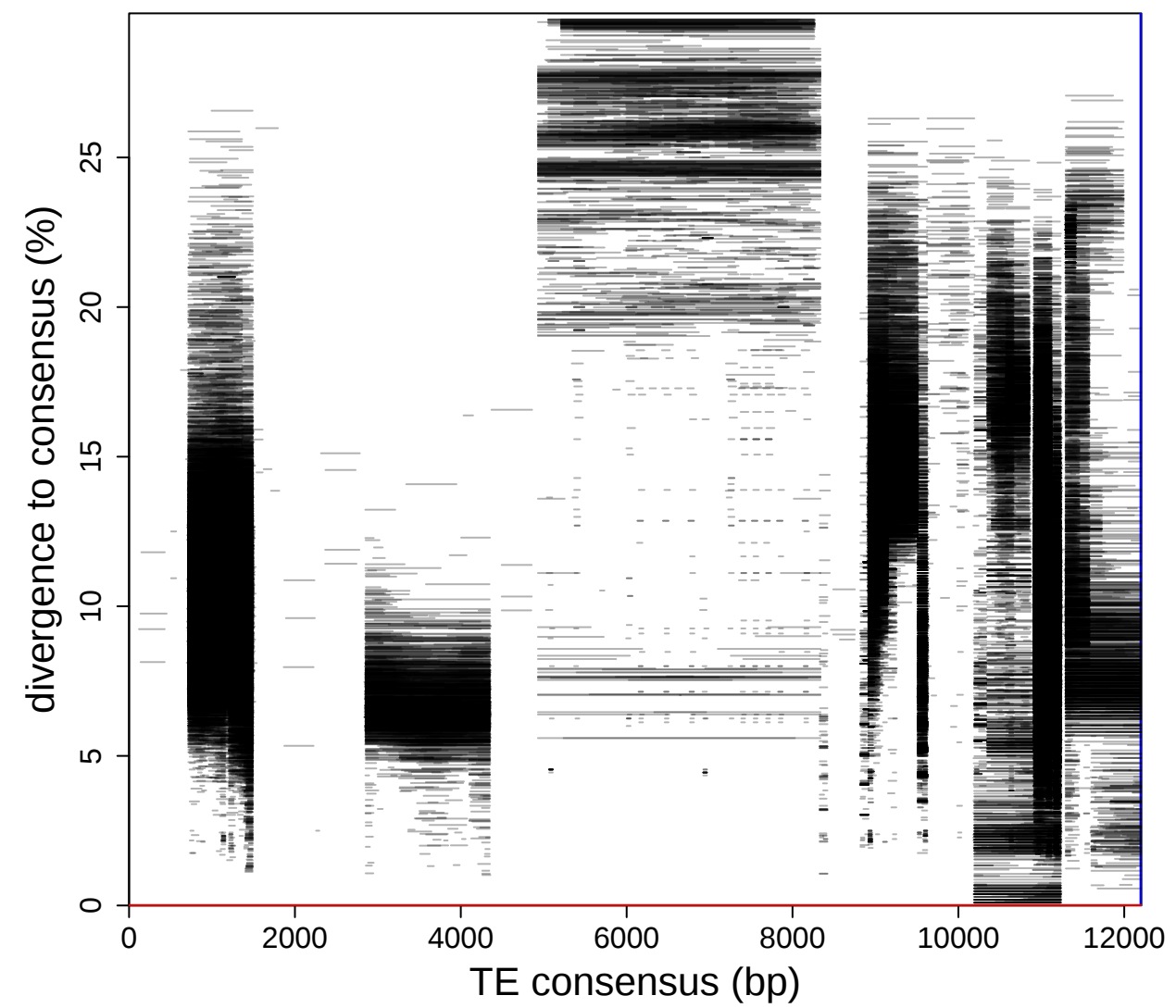
l.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 12207bp; fragments: 80795; full length: 1 (>=10986.3bp)

After TEtrimmer 12207 bp

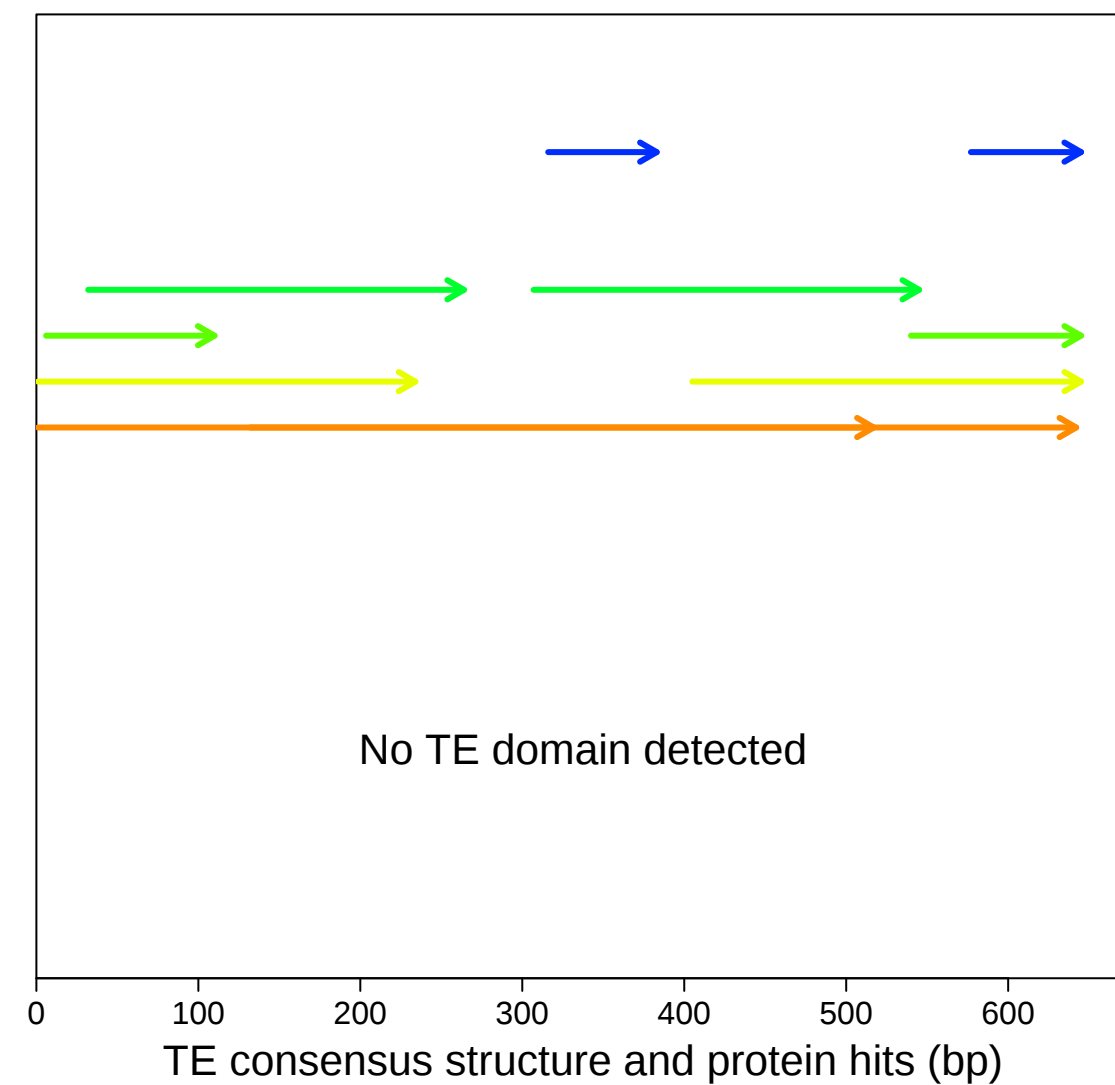
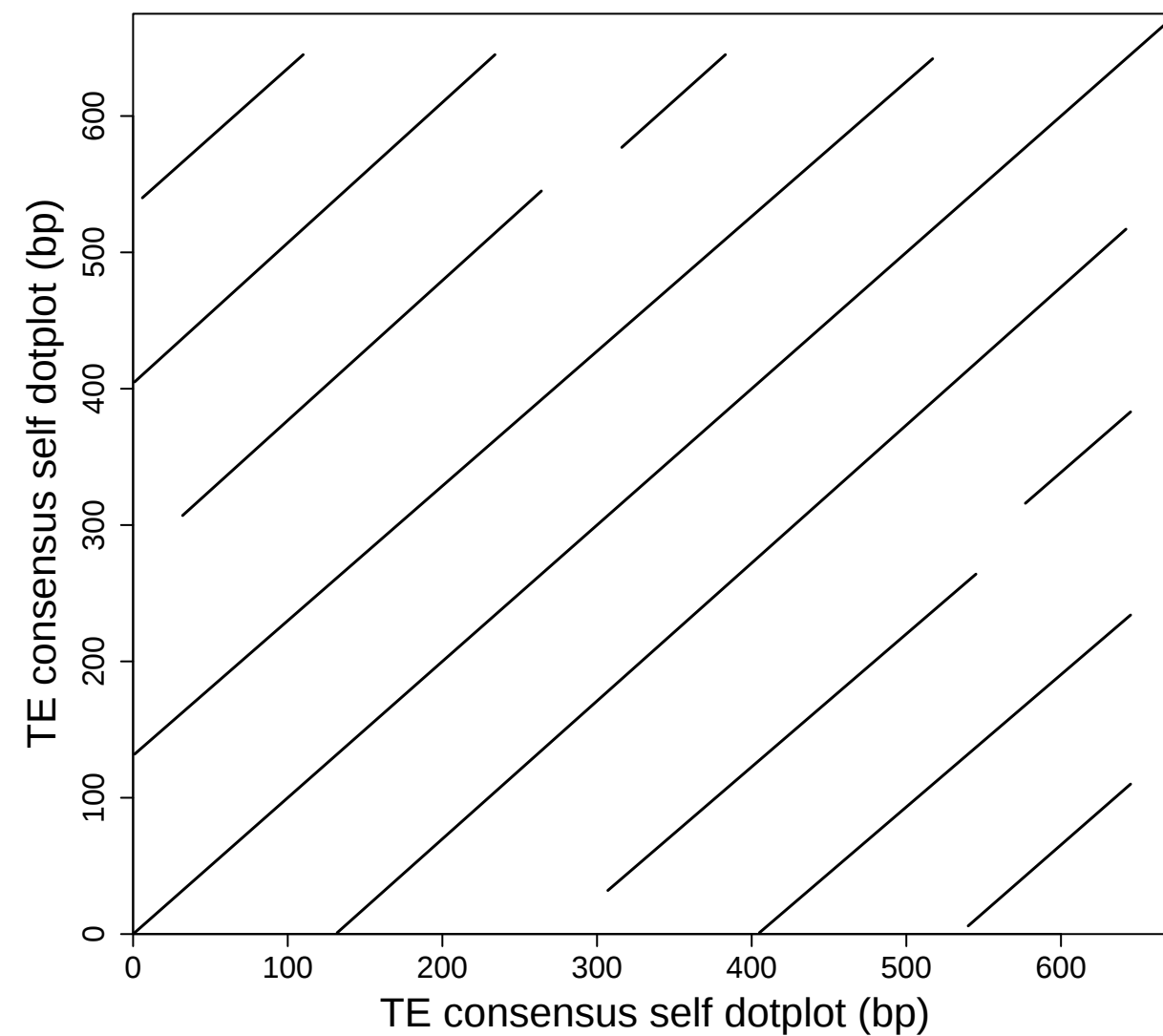
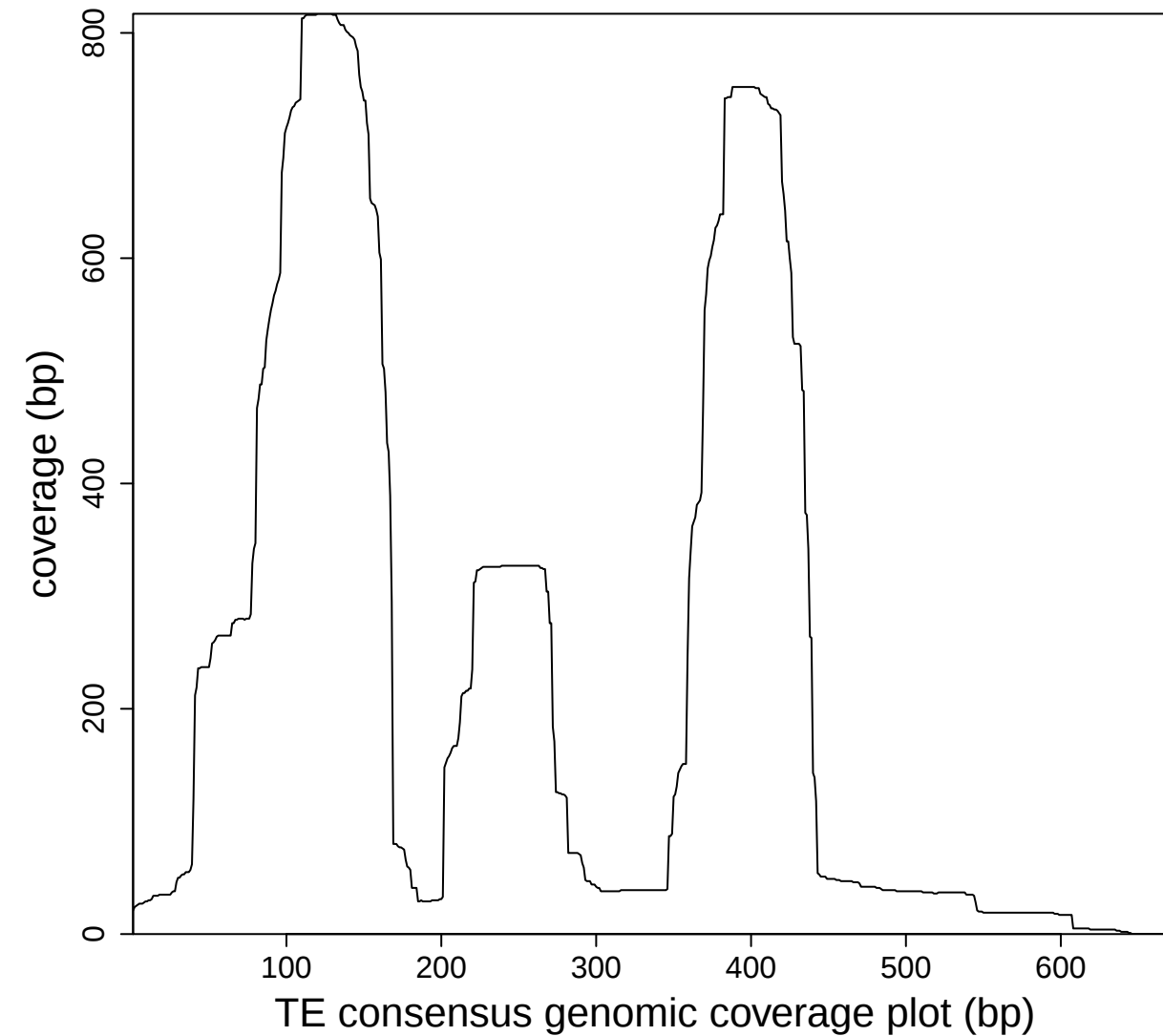
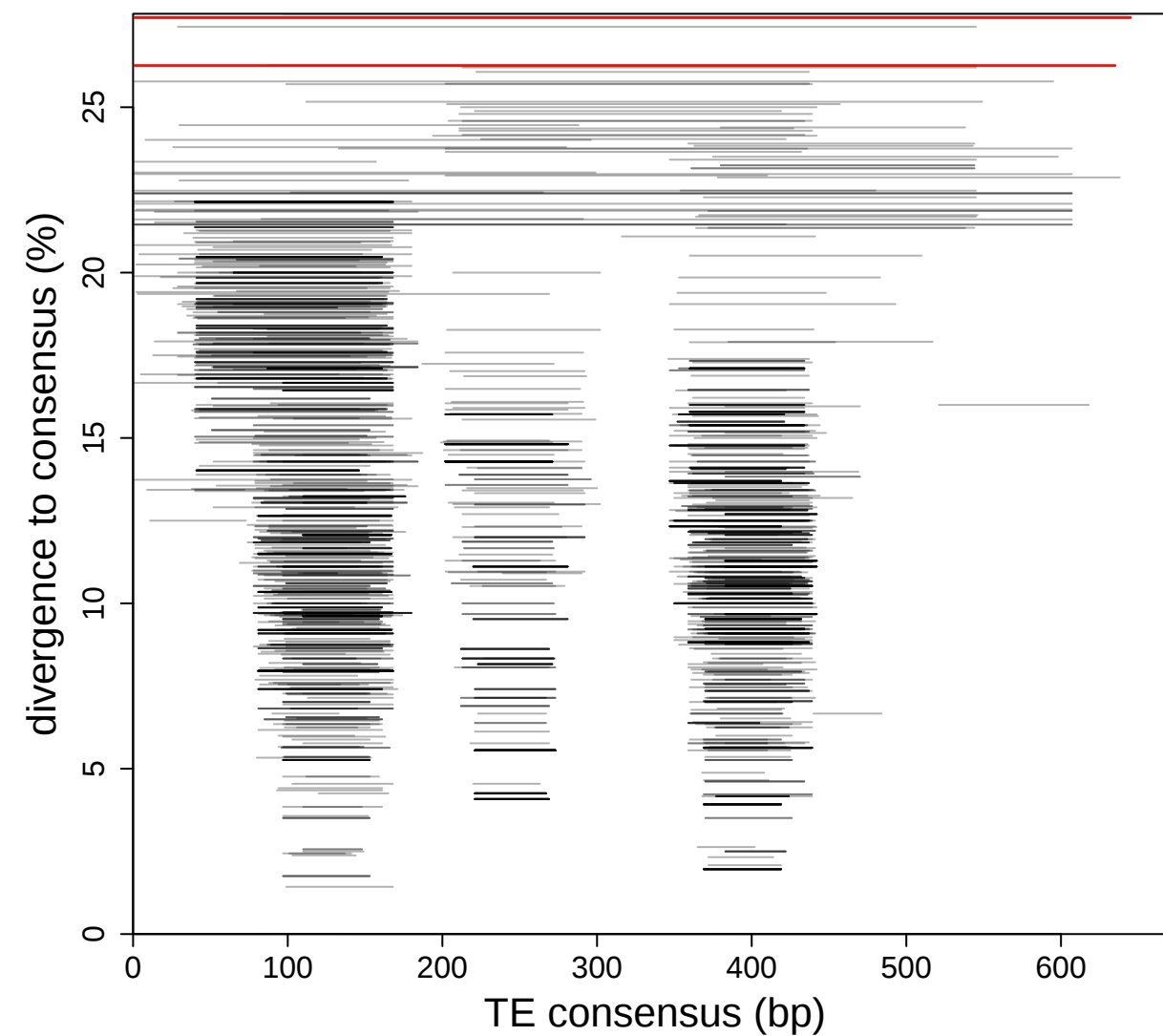


asta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.fa_g
size: 12207bp; fragments: 80795; full length: 1 (>=10986.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

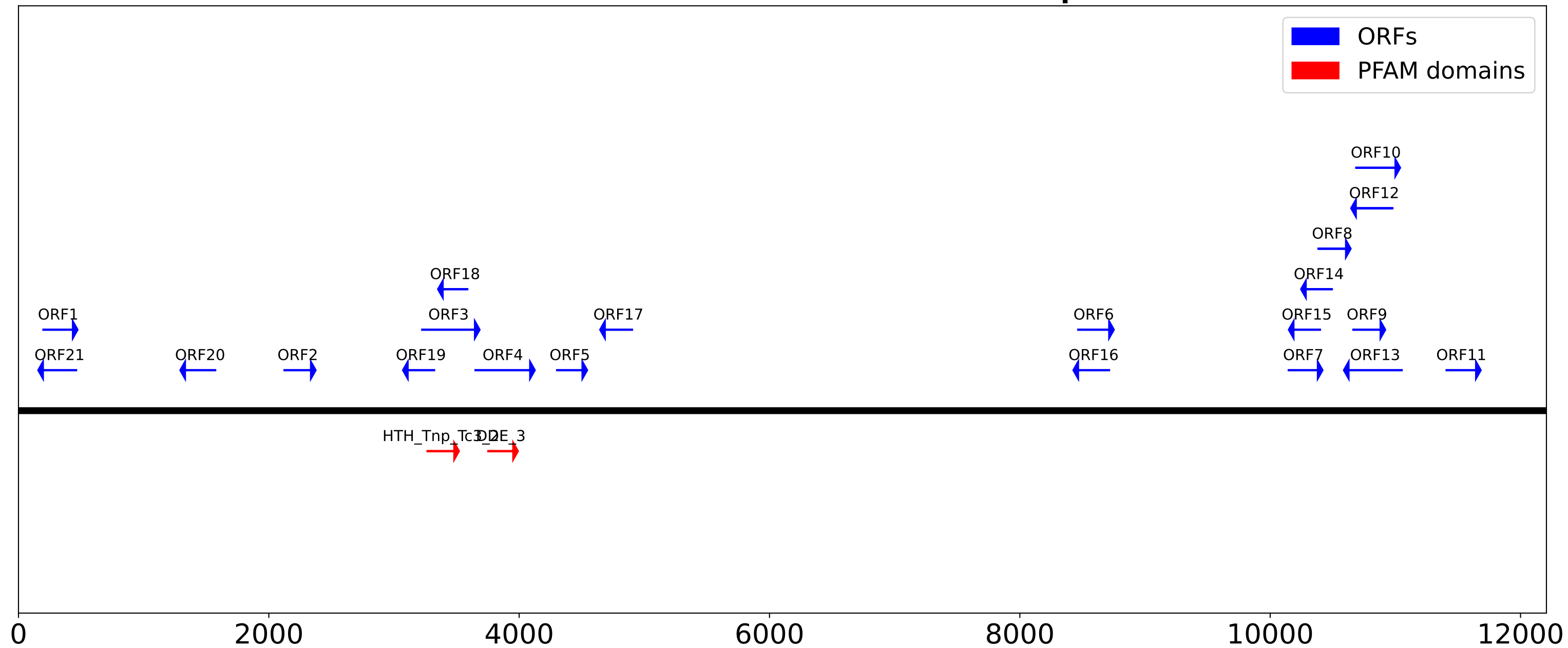


TE: rnd_4_family_318
size: 675bp; fragments: 1835; full length: 2 (≥ 607.5 bp)

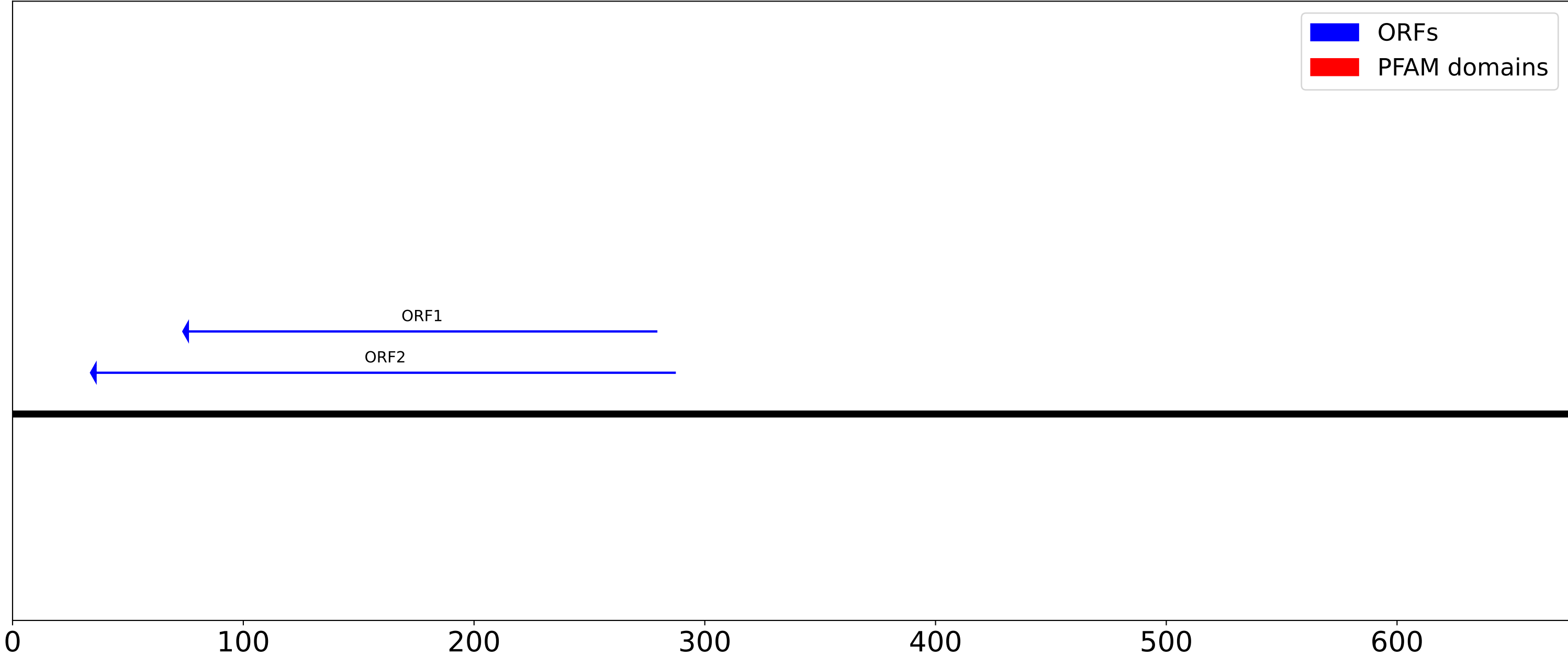


Before TEtrimmer 675 bp

After TEtrimmer ORF and PFAM domain plot



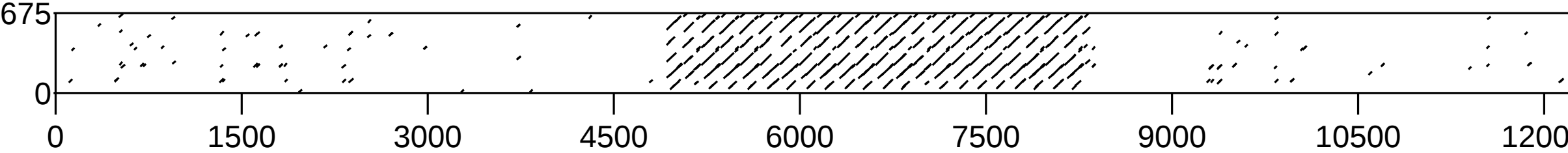
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)