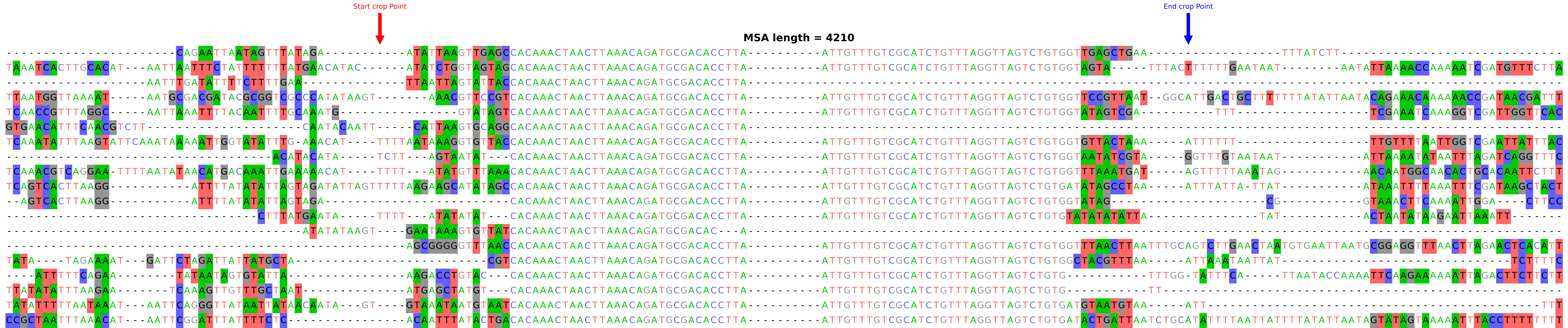
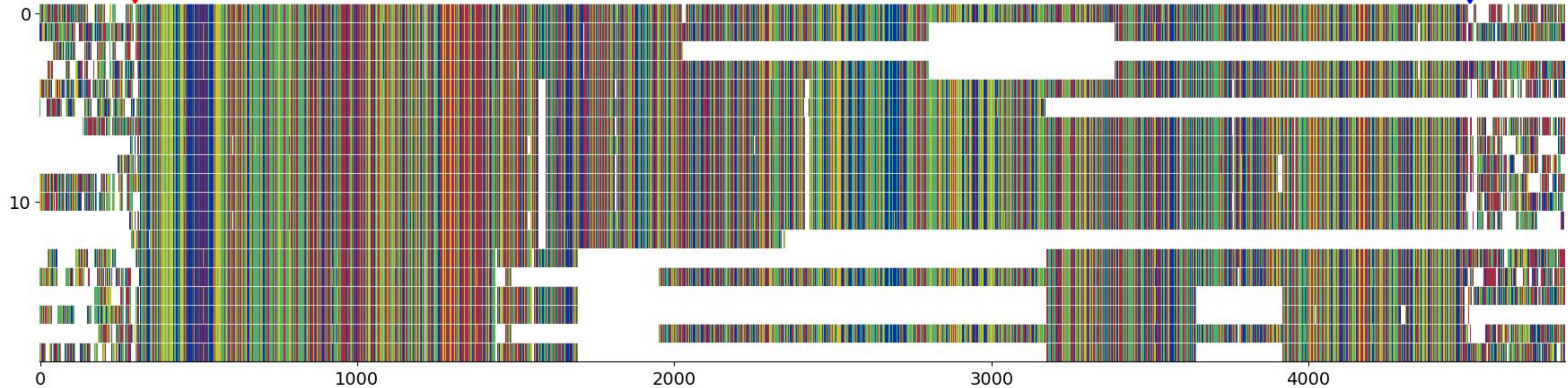




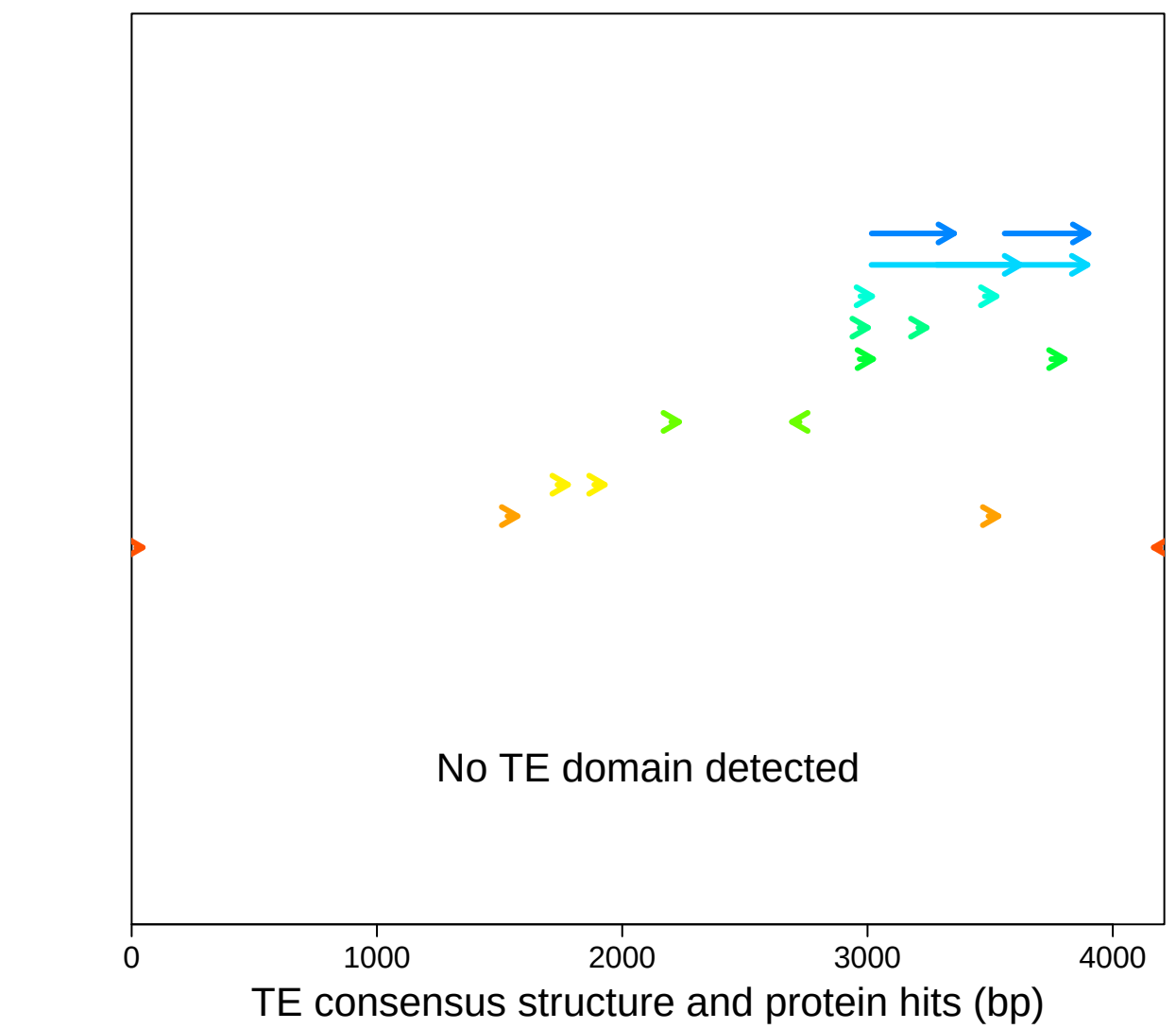
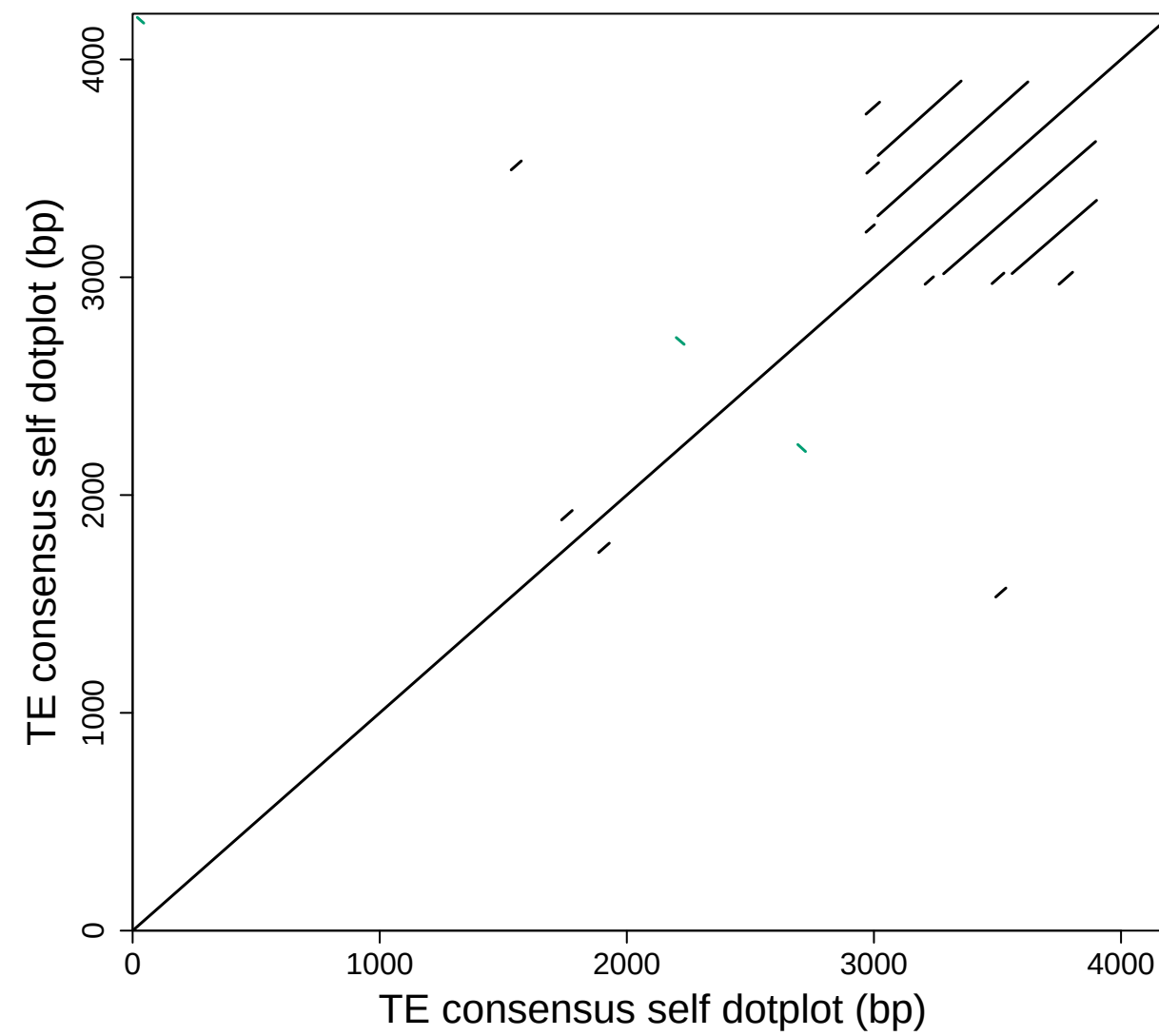
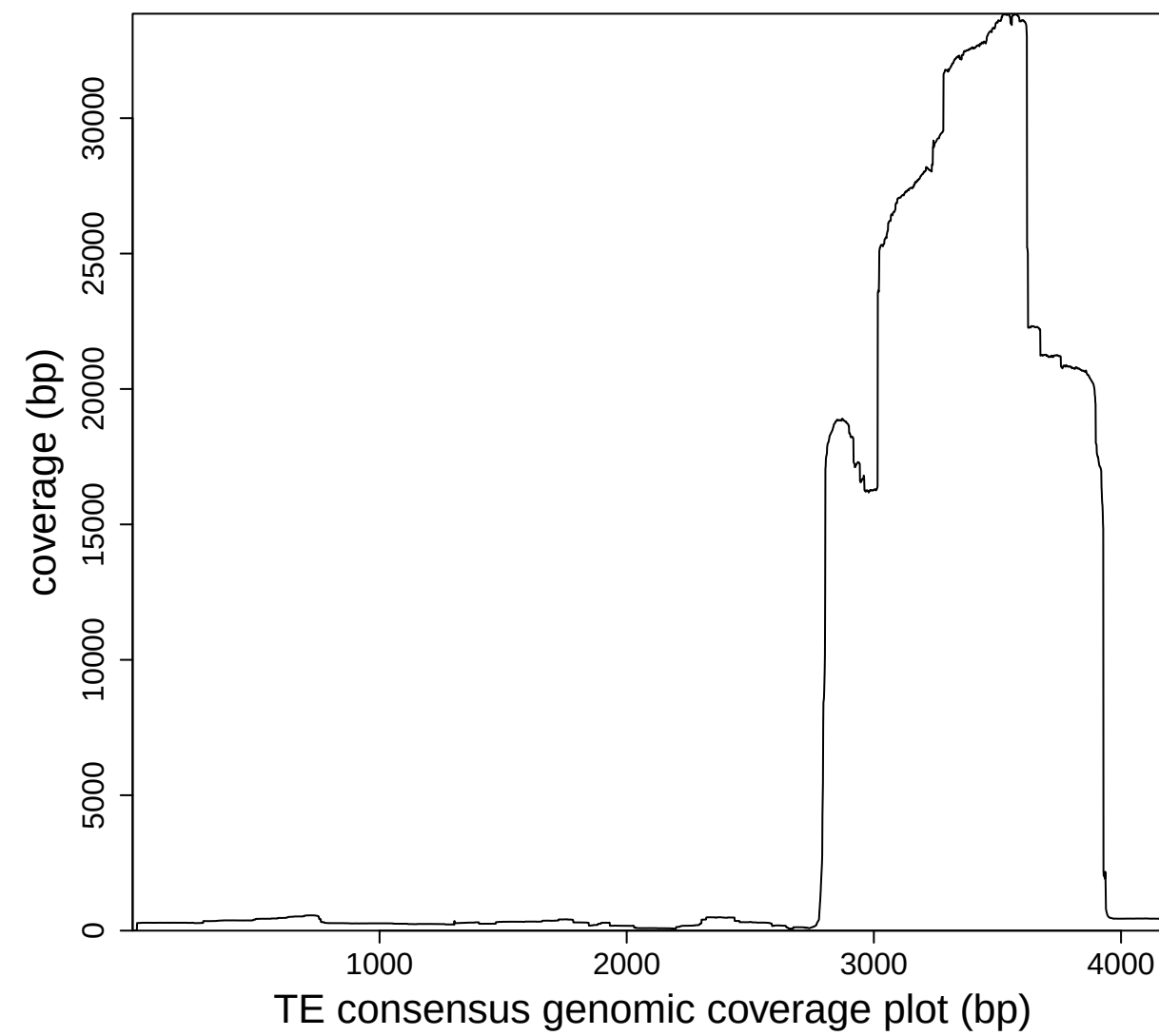
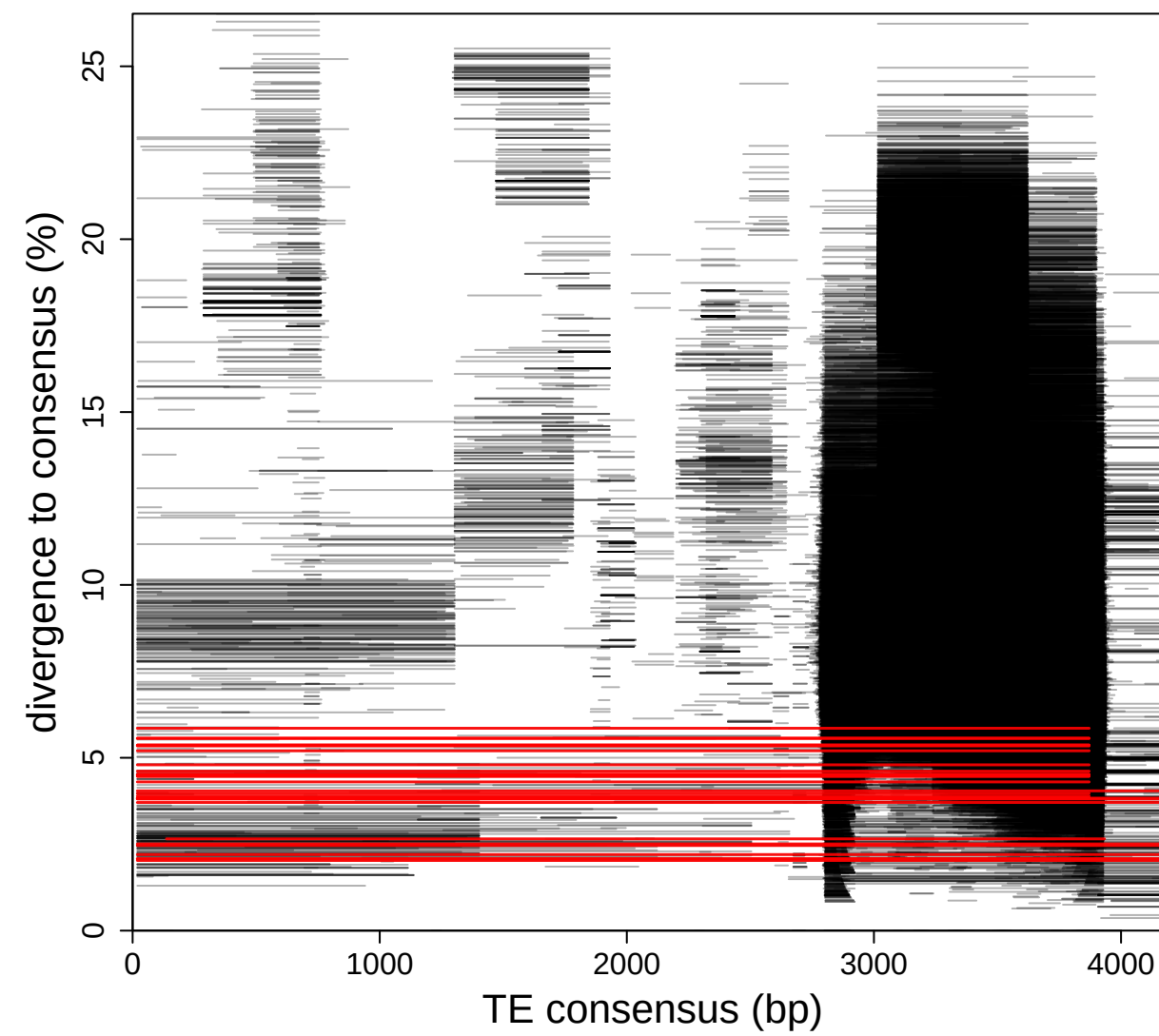
Start crop Point

End crop Point



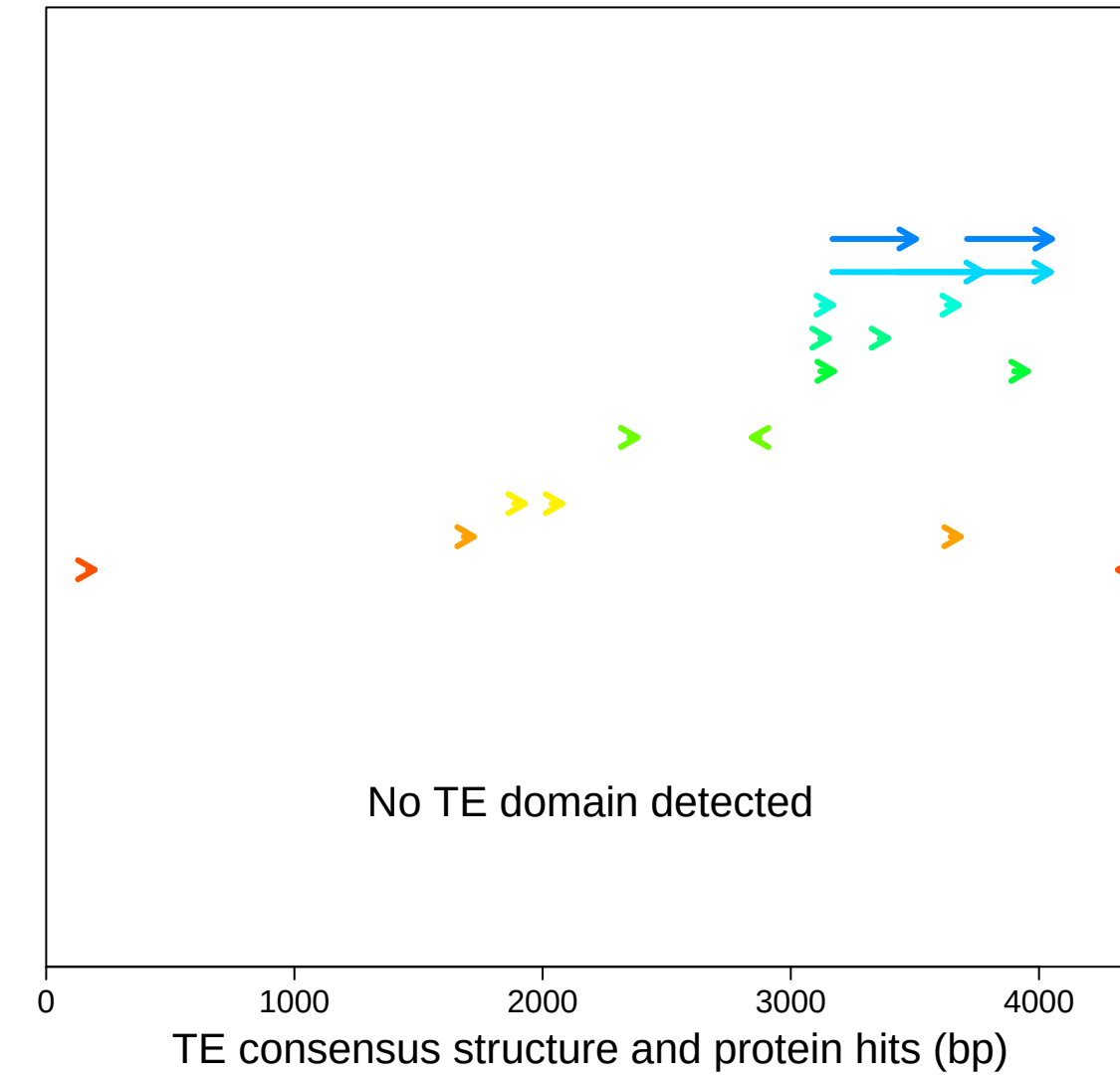
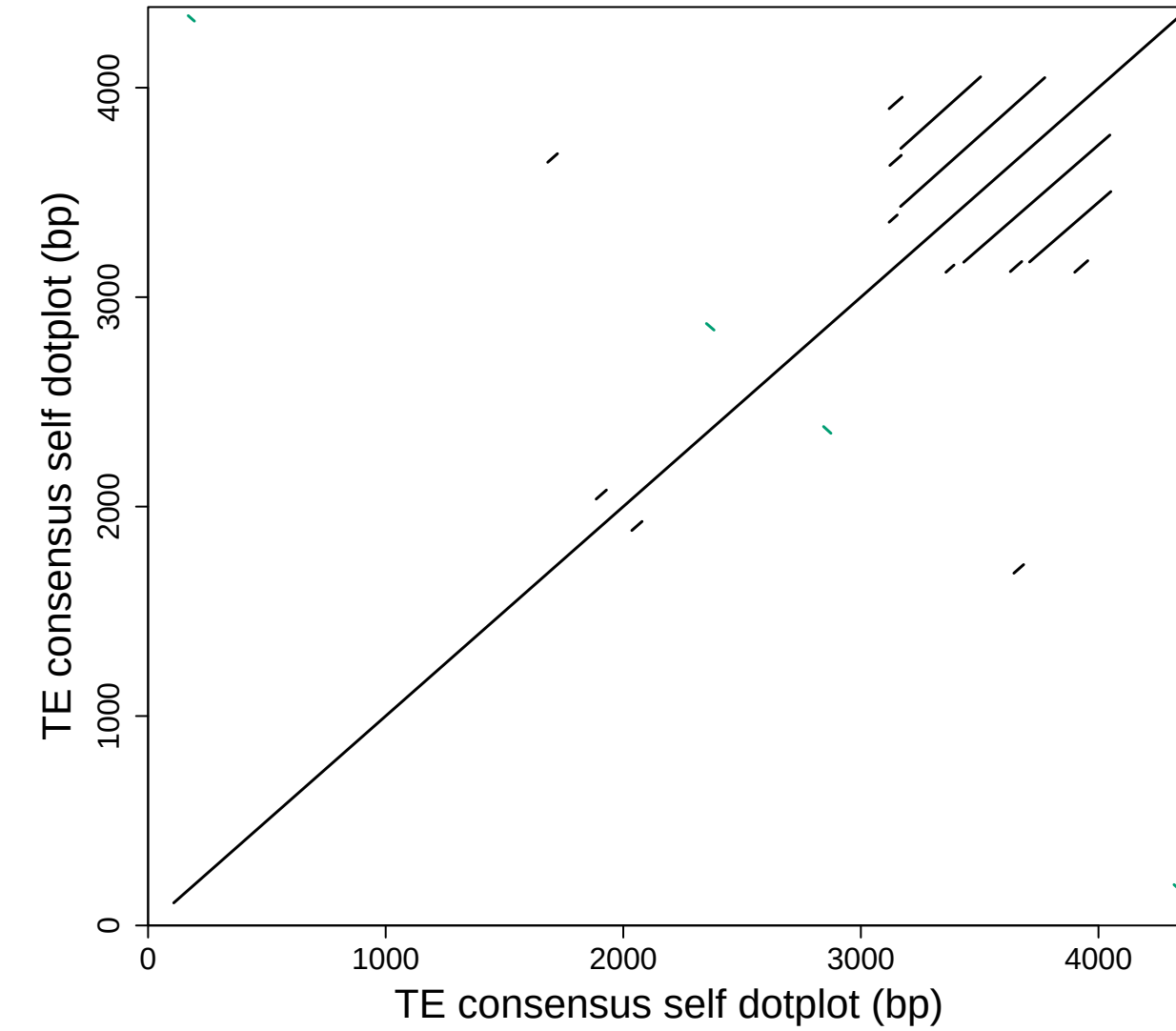
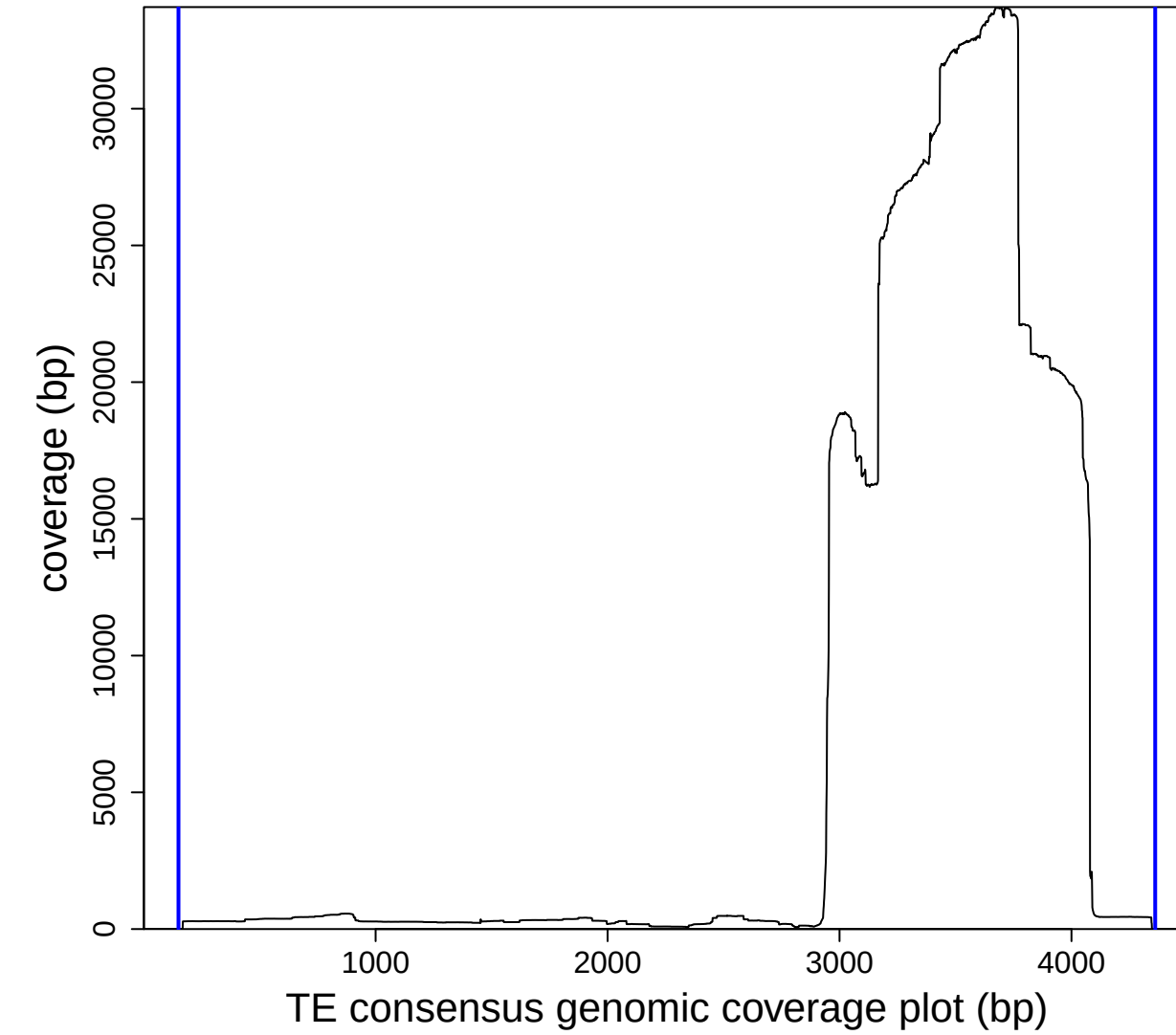
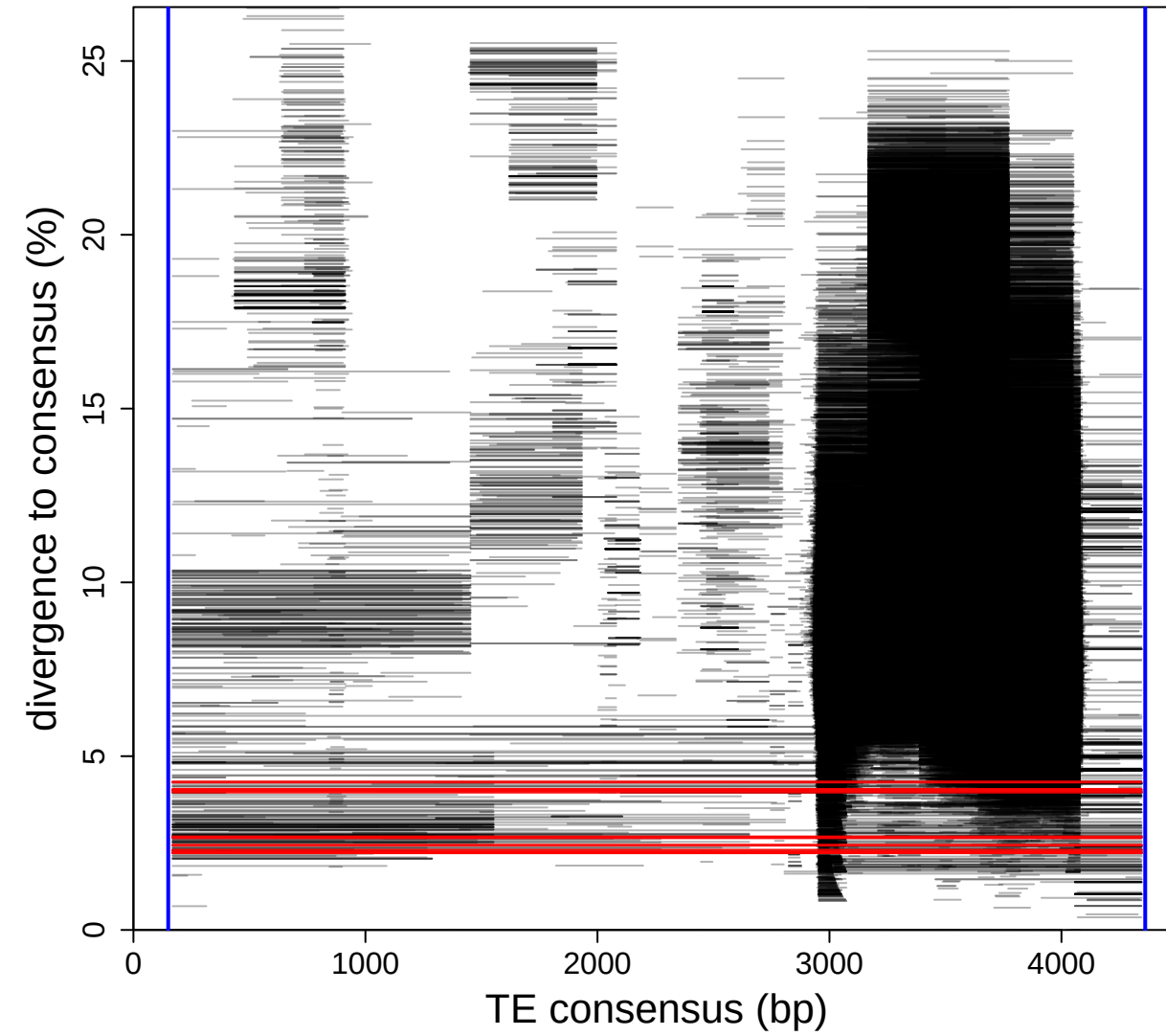
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 4210bp; fragments: 50159; full length: 31 (>=3789bp)

After TETrimmer 4210 bp



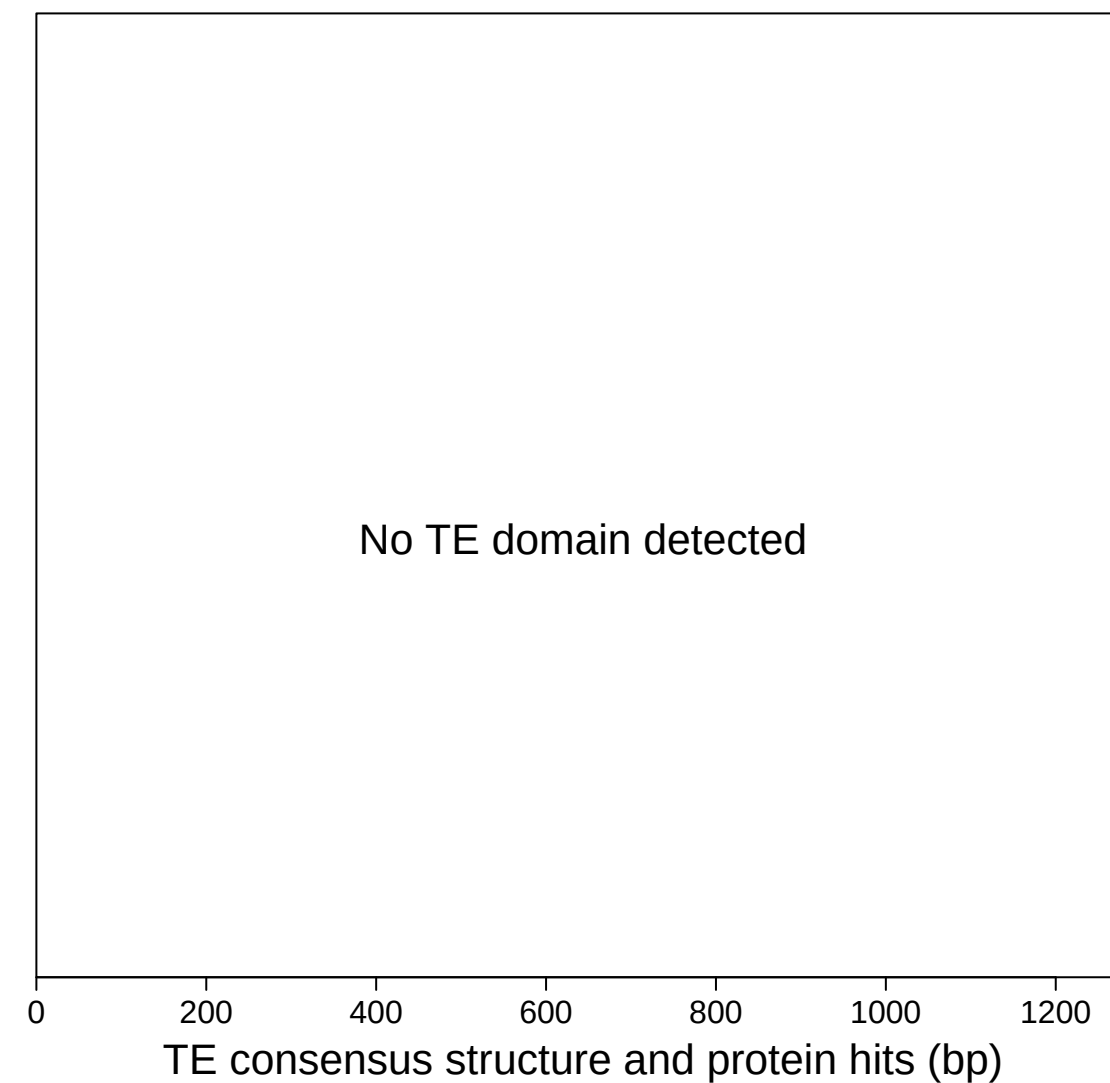
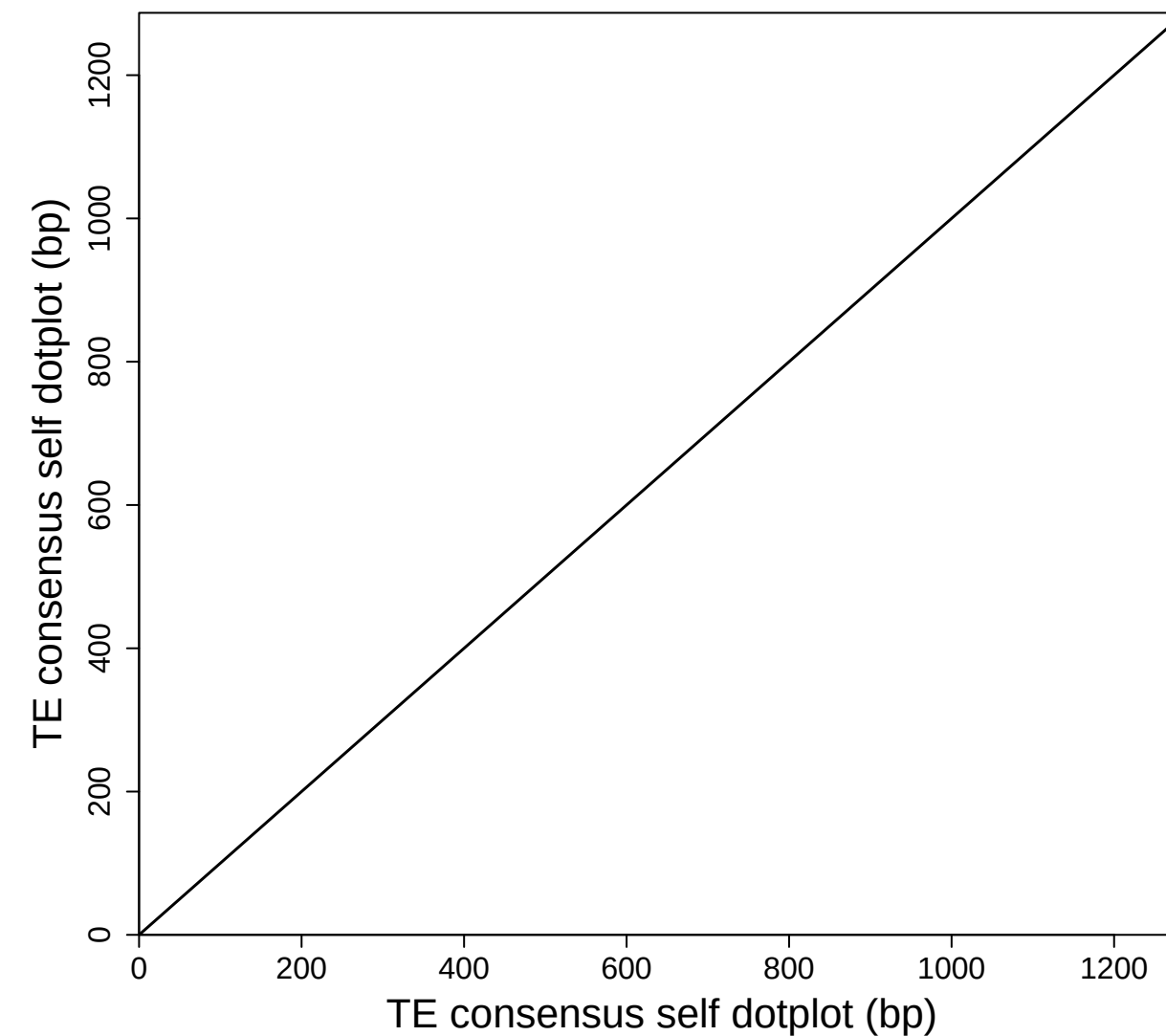
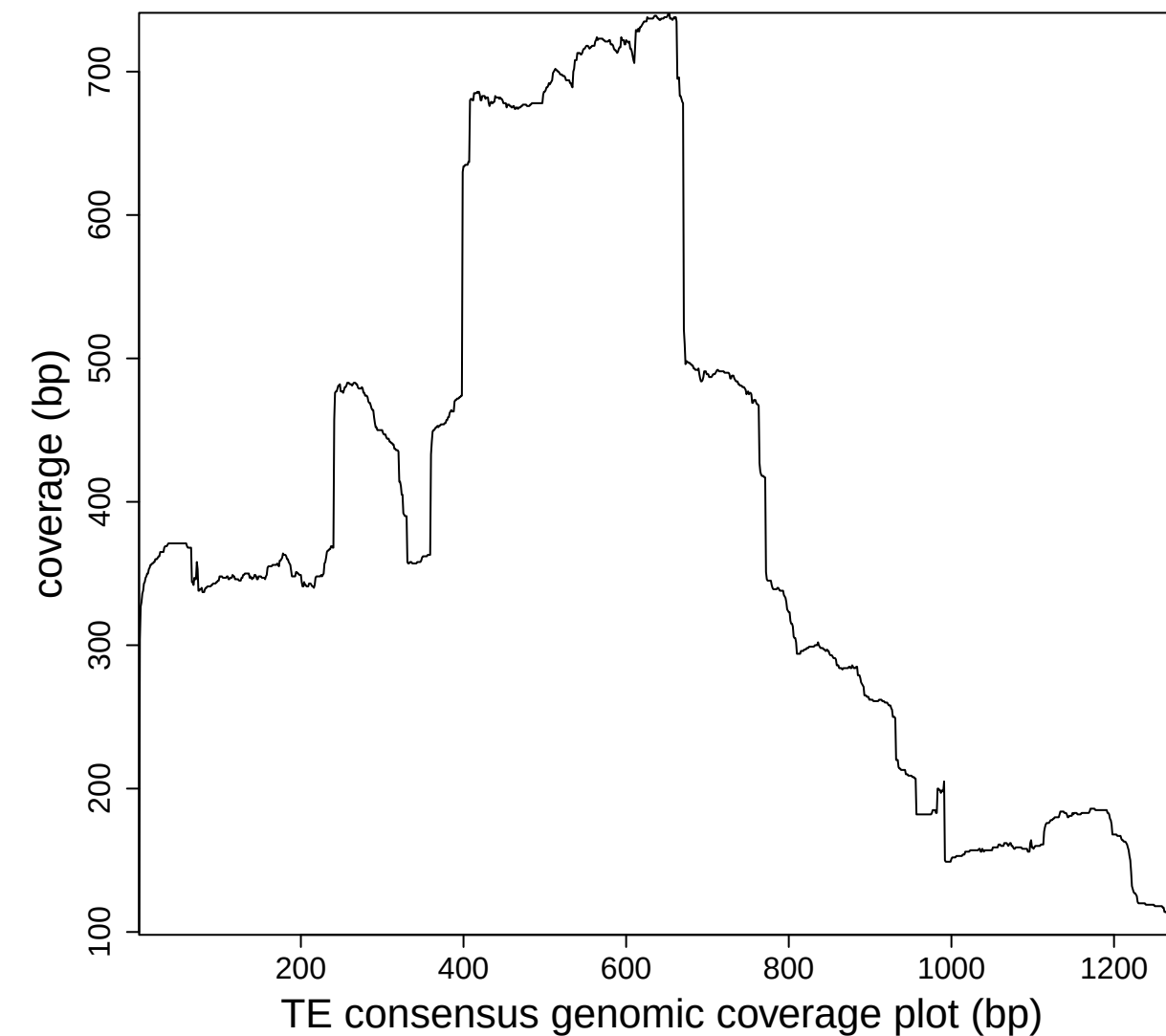
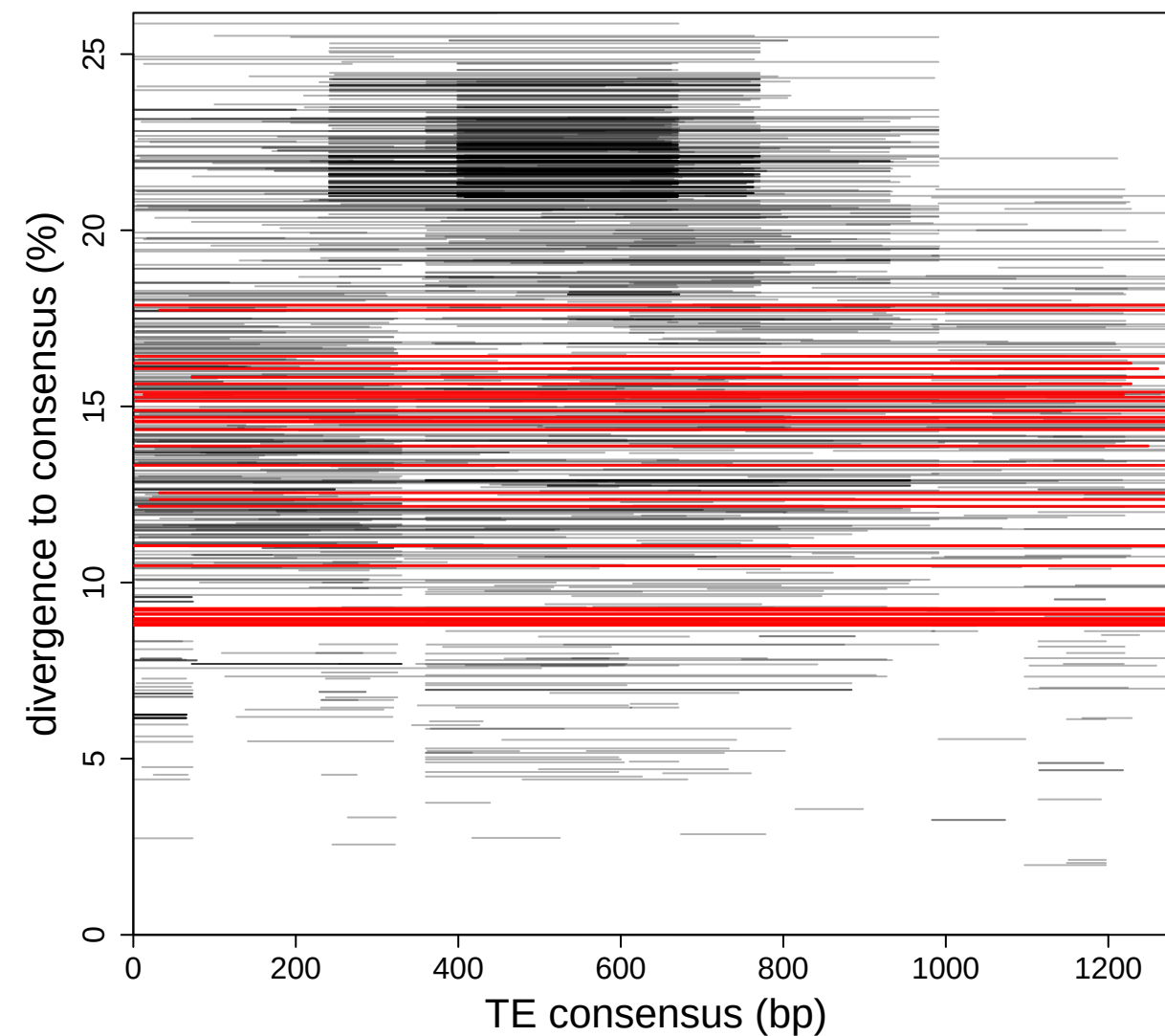
0.fasta.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 4511bp; fragments: 49278; full length: 10 (>=4059.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

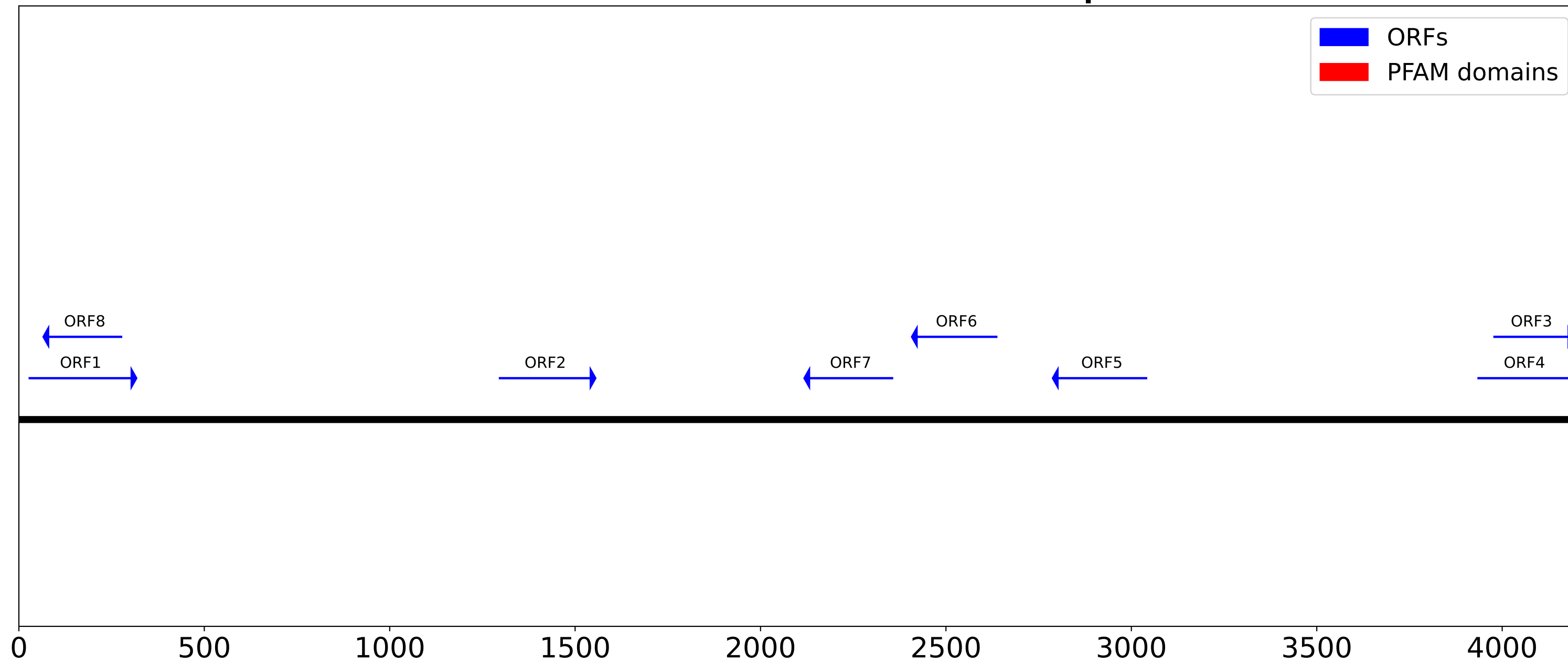


TE: rnd_4_family_1830
size: 1287bp; fragments: 1272; full length: 38 (≥ 1158.3 bp)

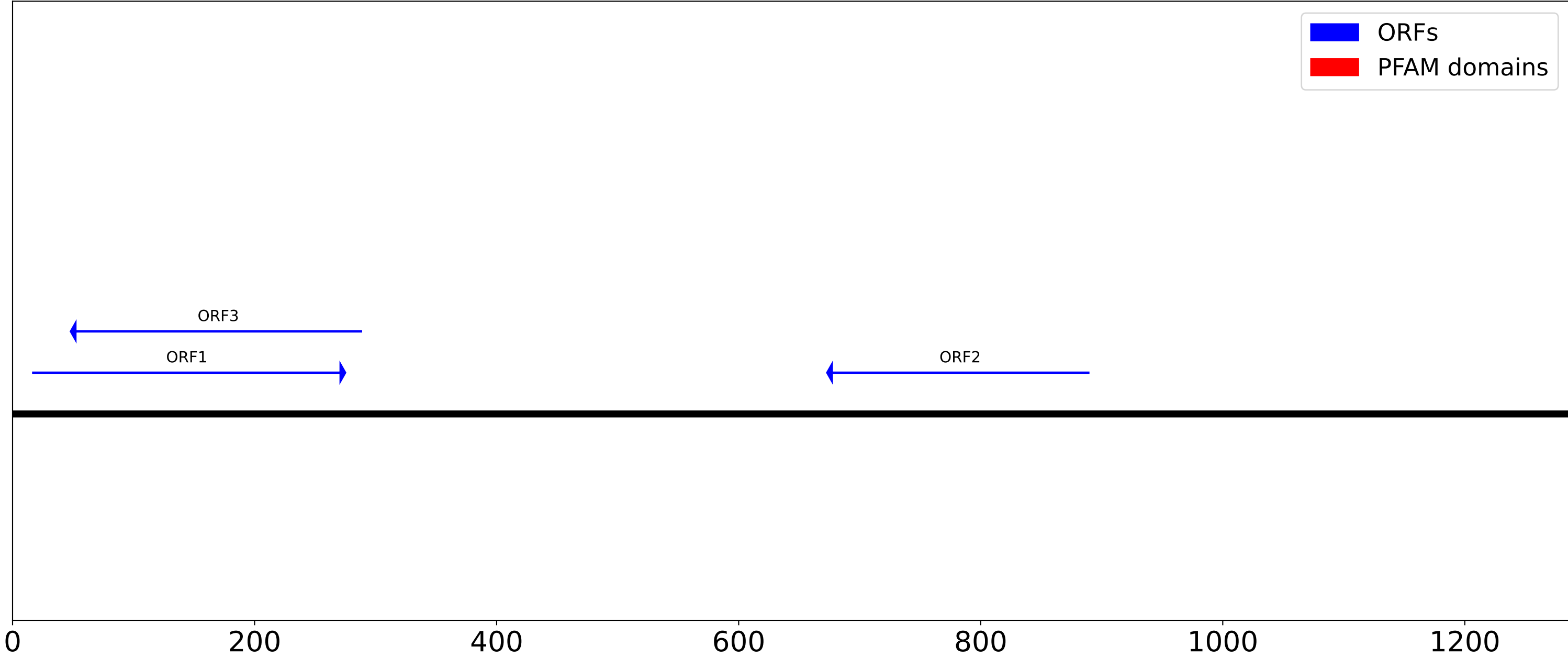
Before TEtrimmer 1287 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

: consensus before TEtrimmer (bp)

