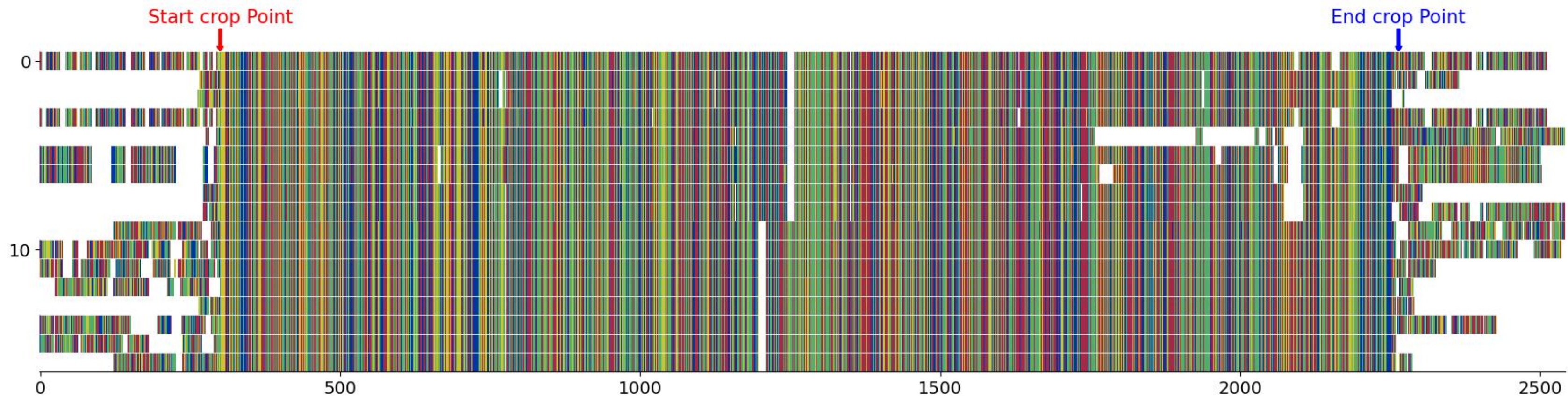
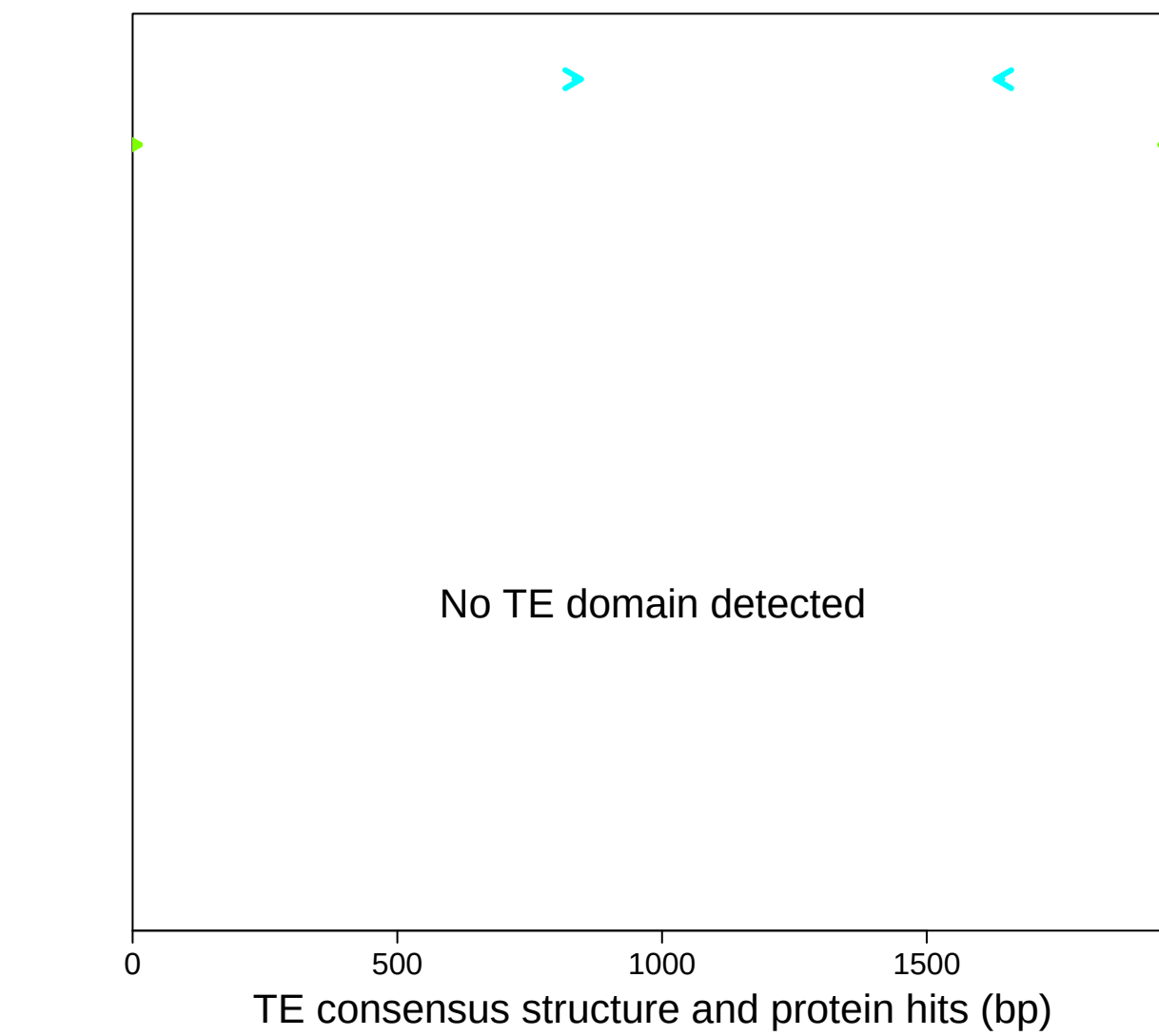
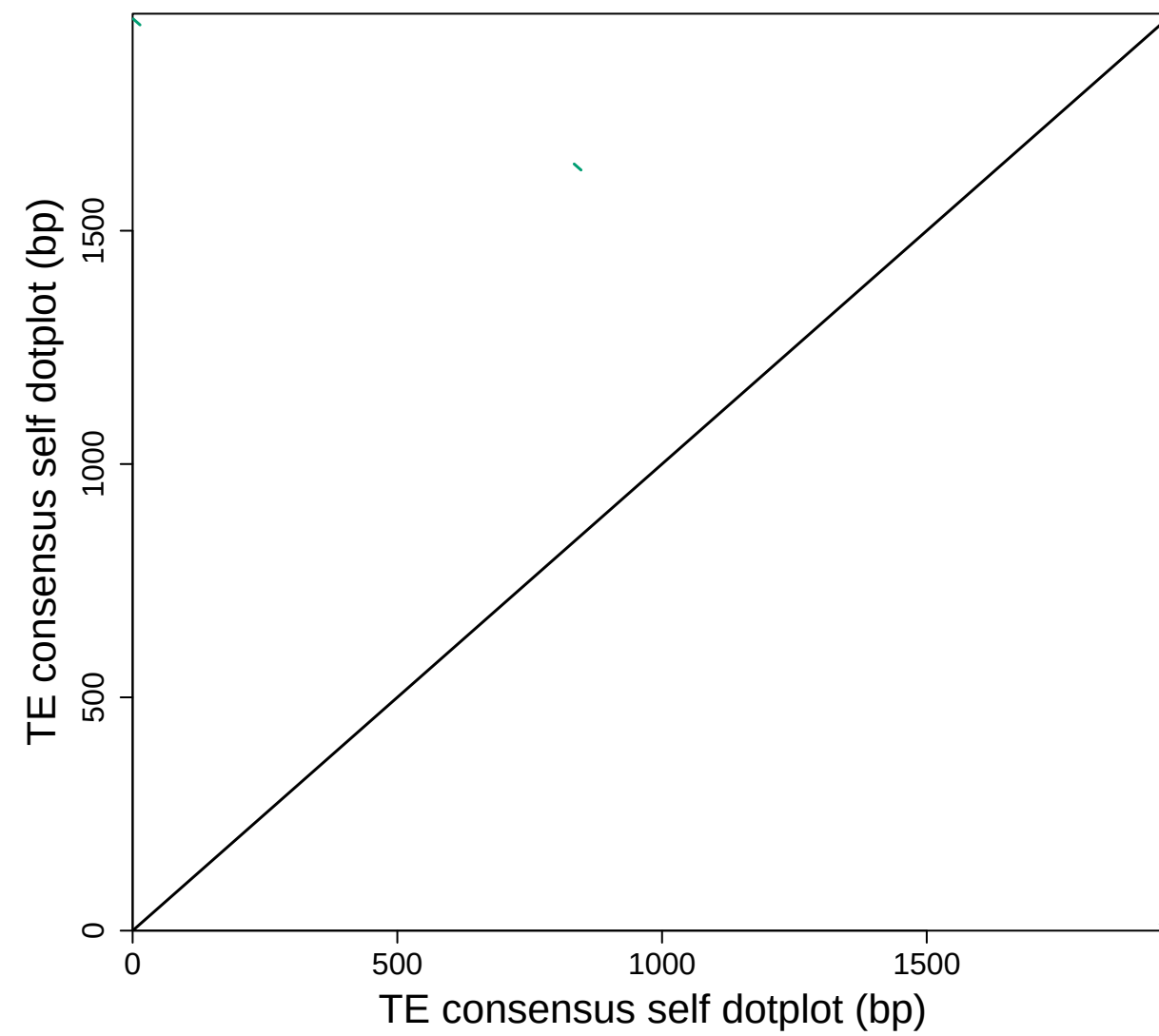
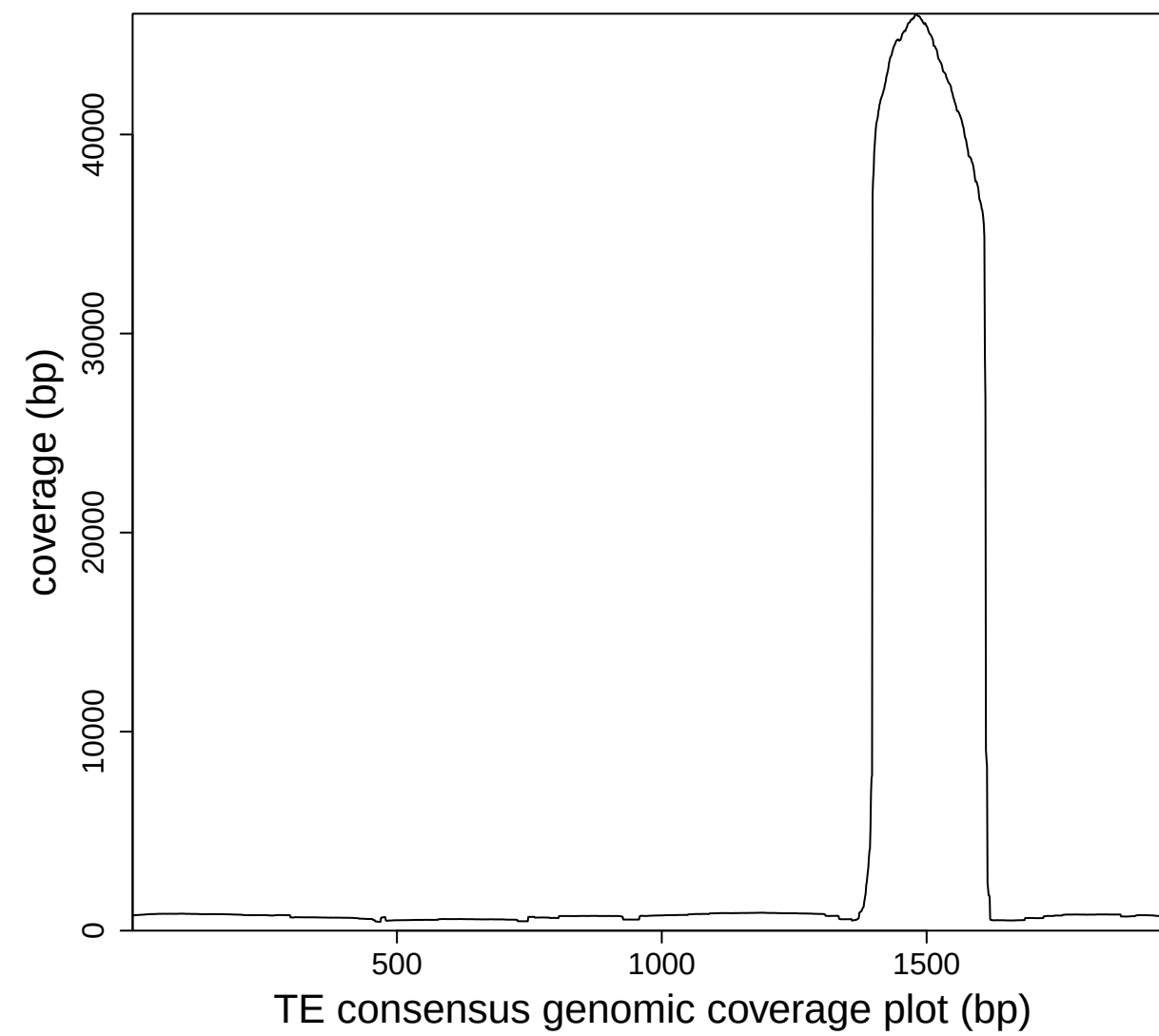
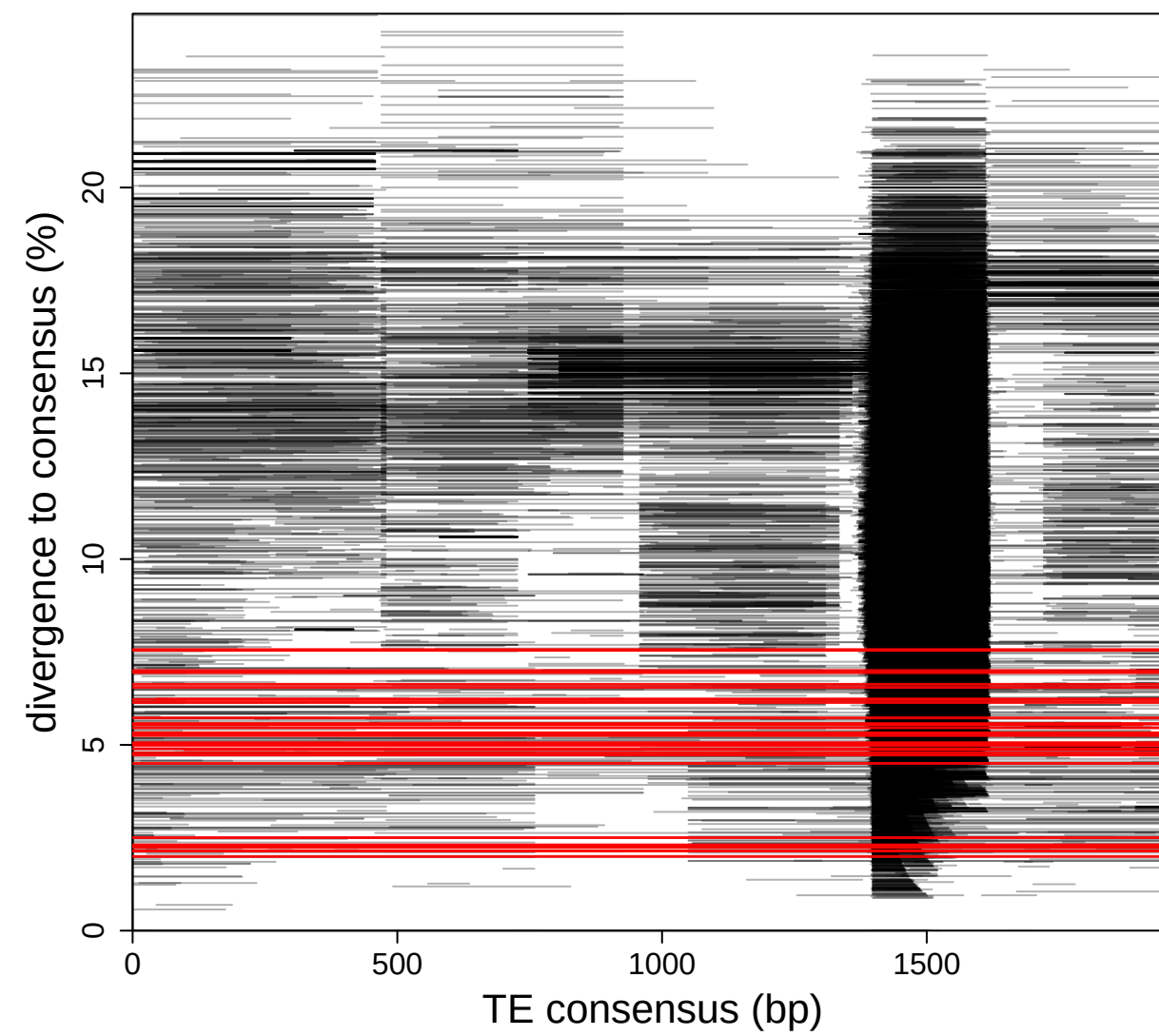




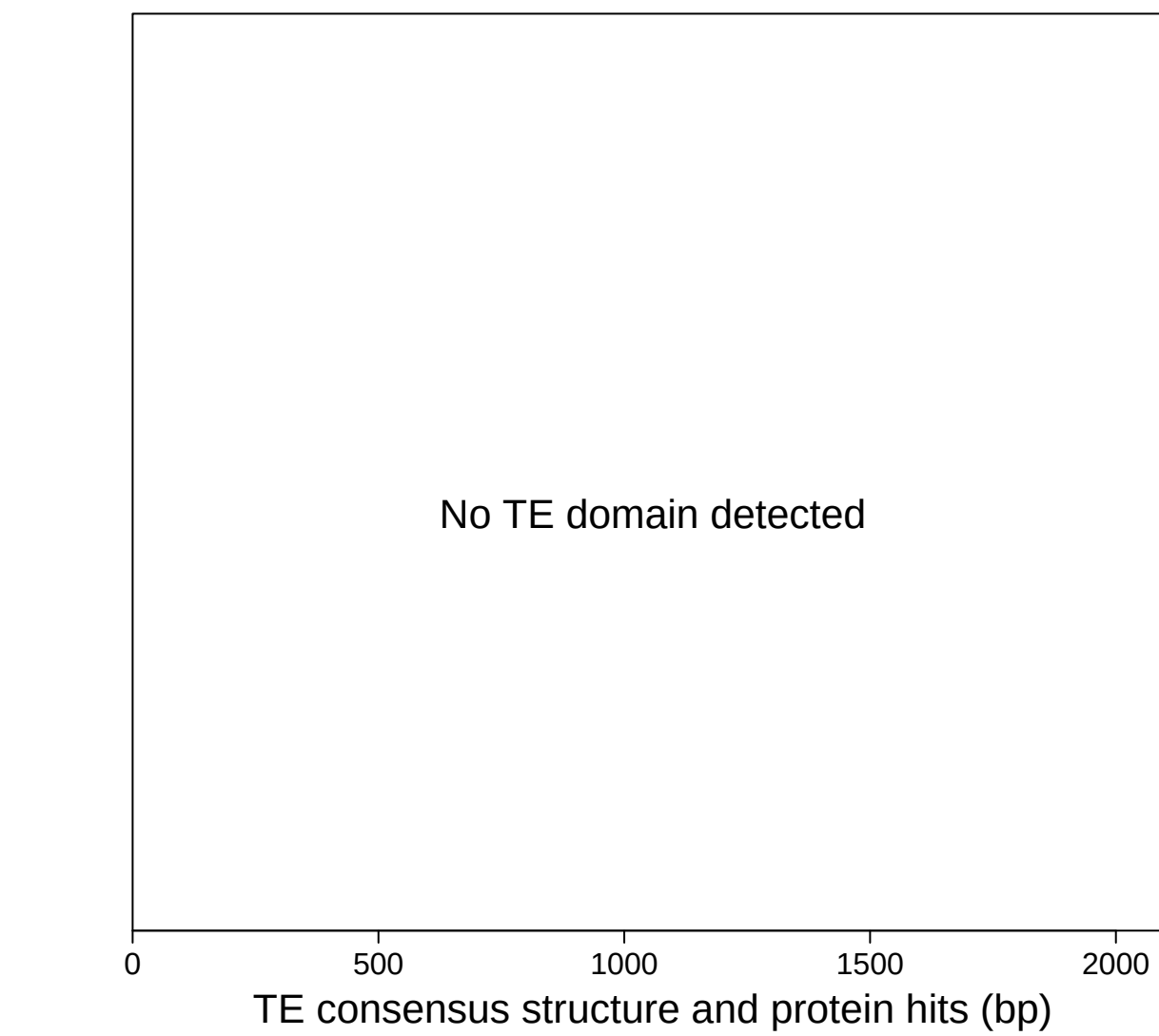
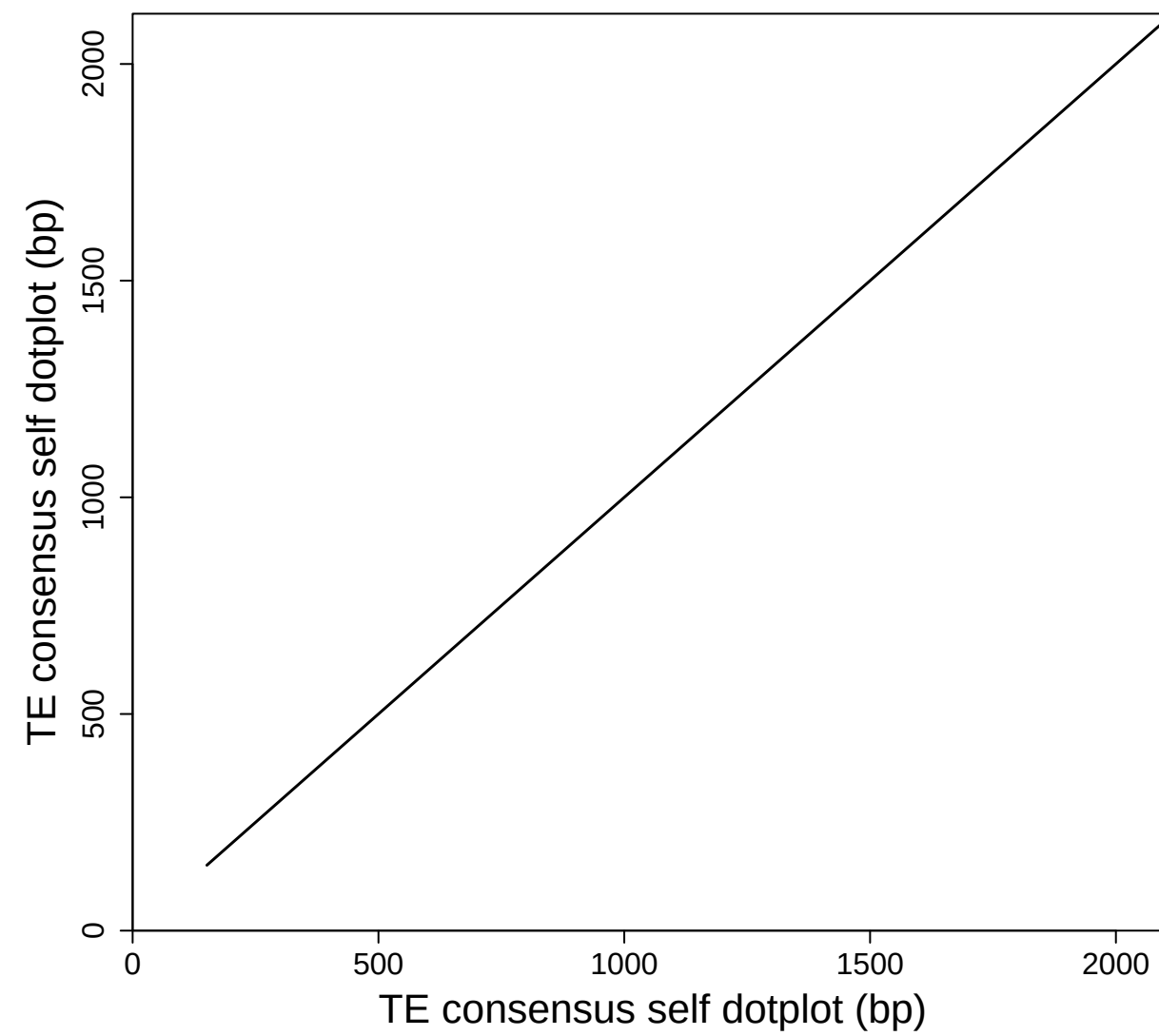
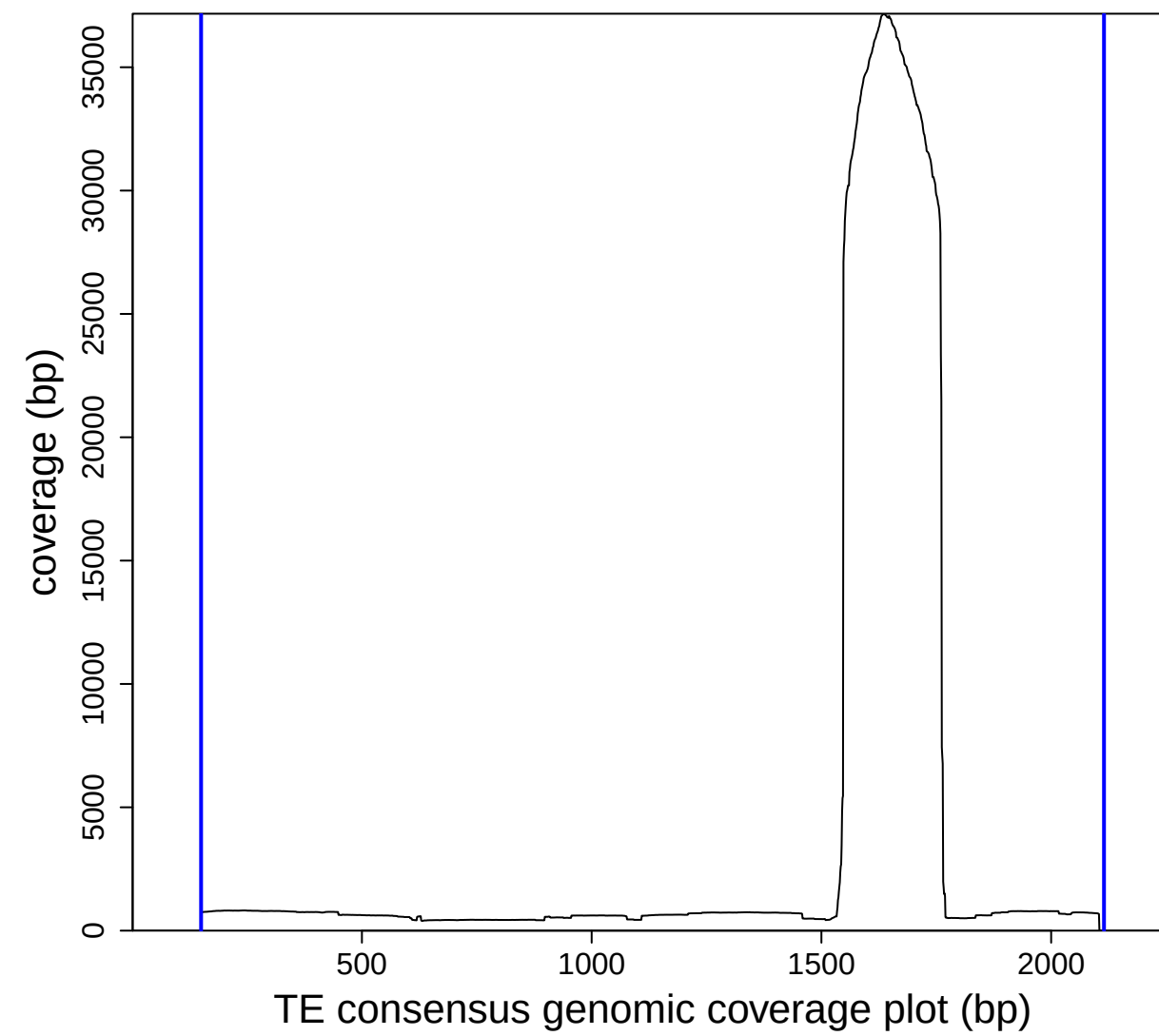
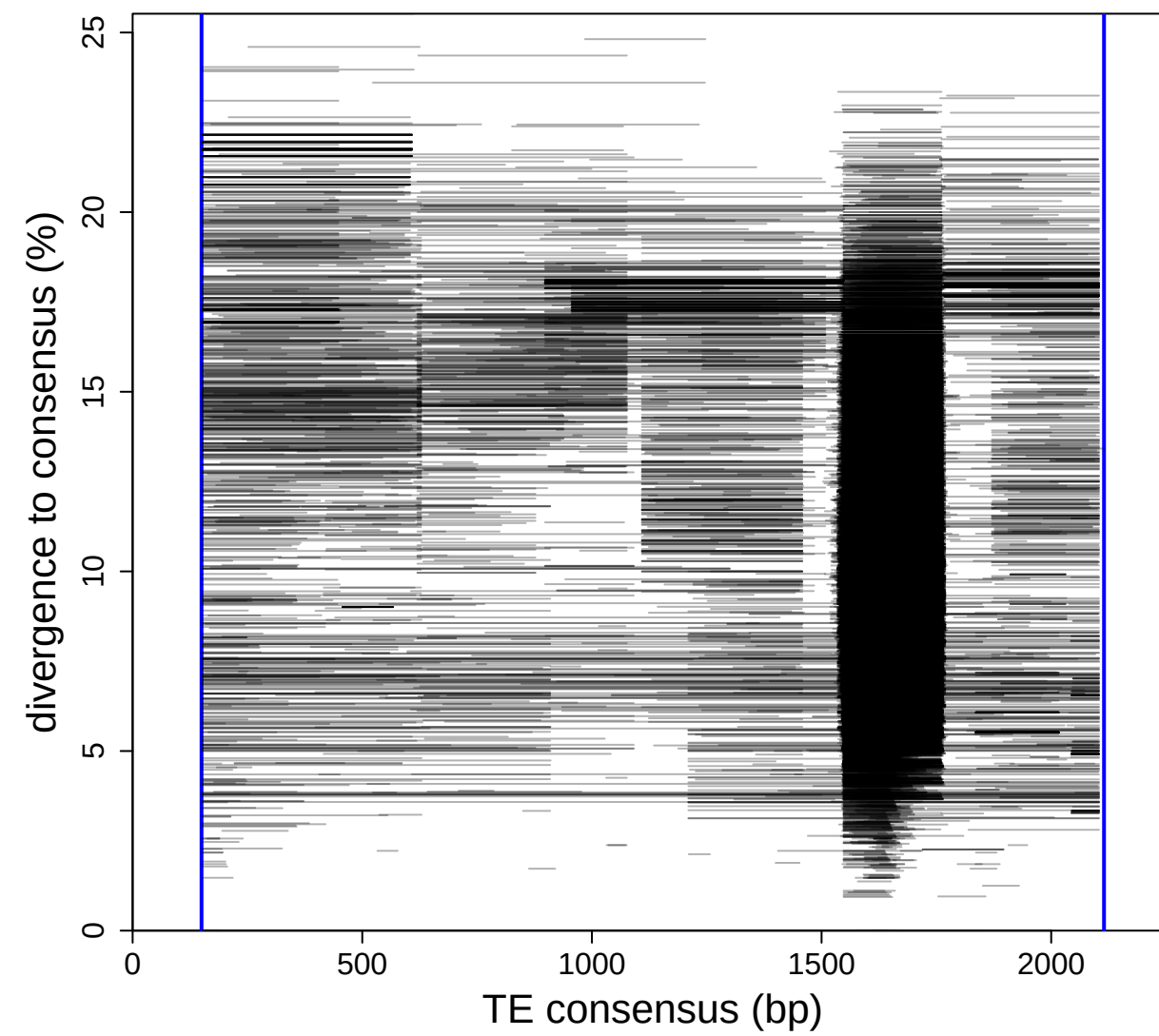
b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 1965bp; fragments: 50890; full length: 35 (≥ 1768.5 bp)

After TEtrimmer 1965 bp

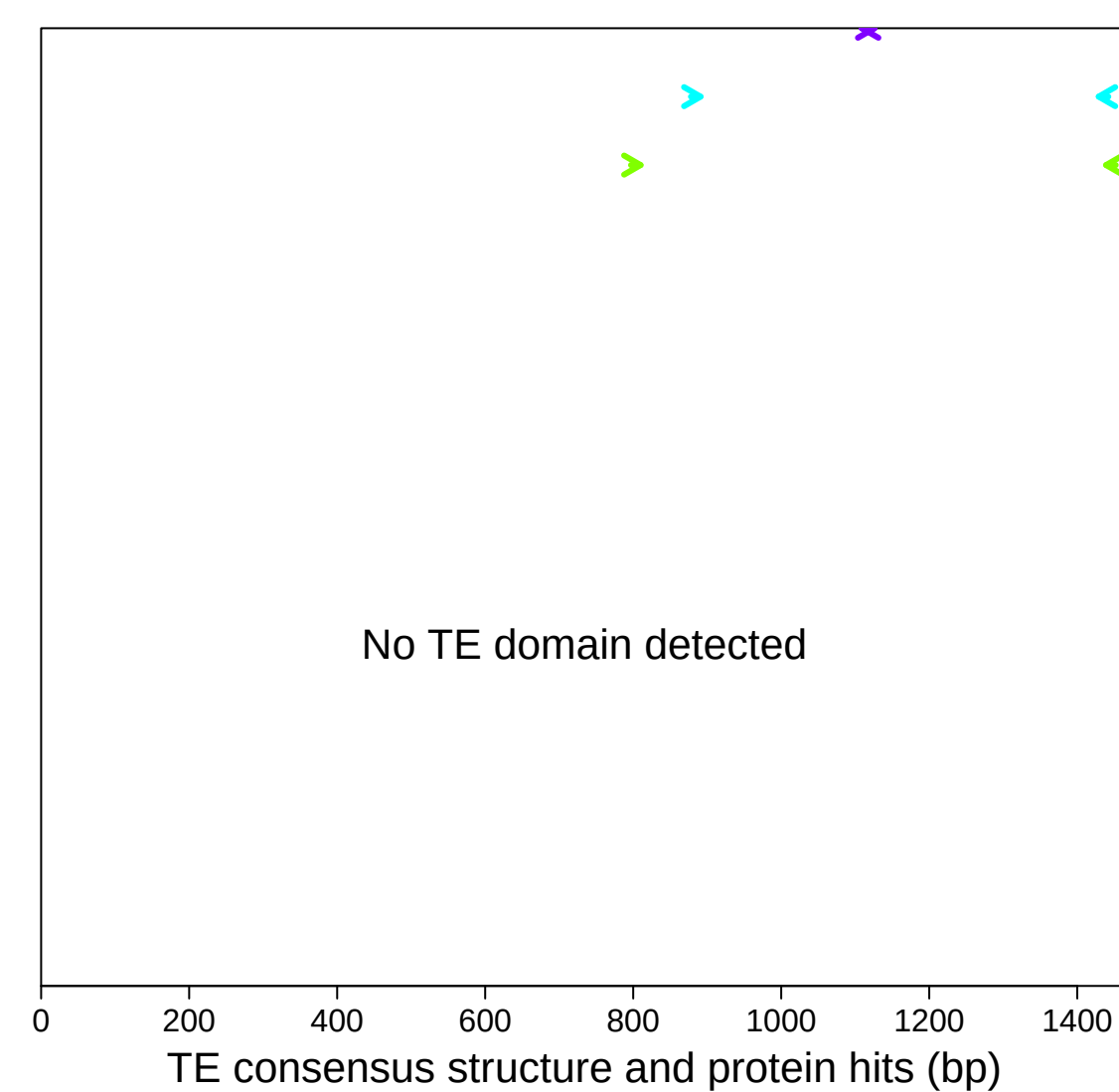
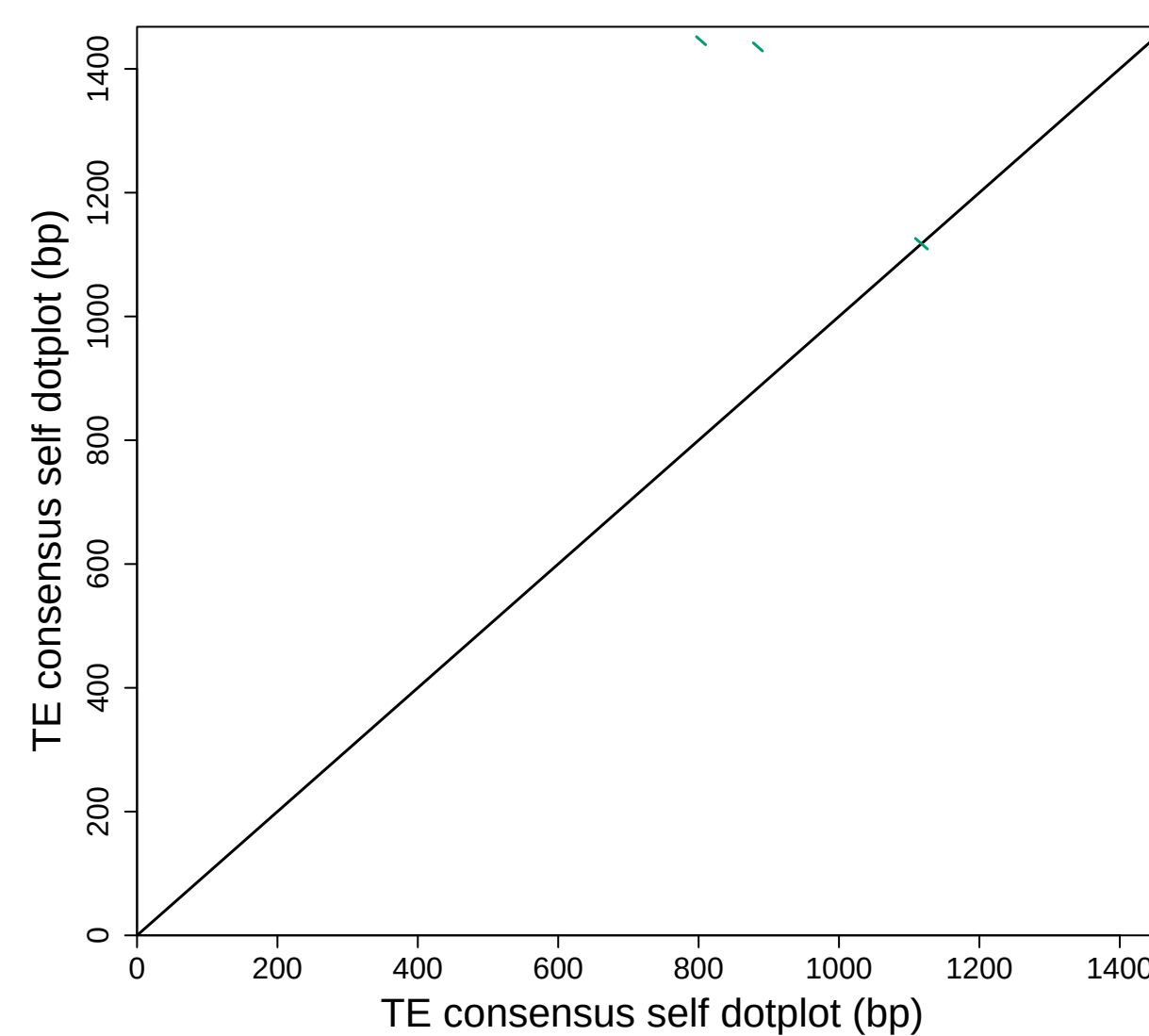
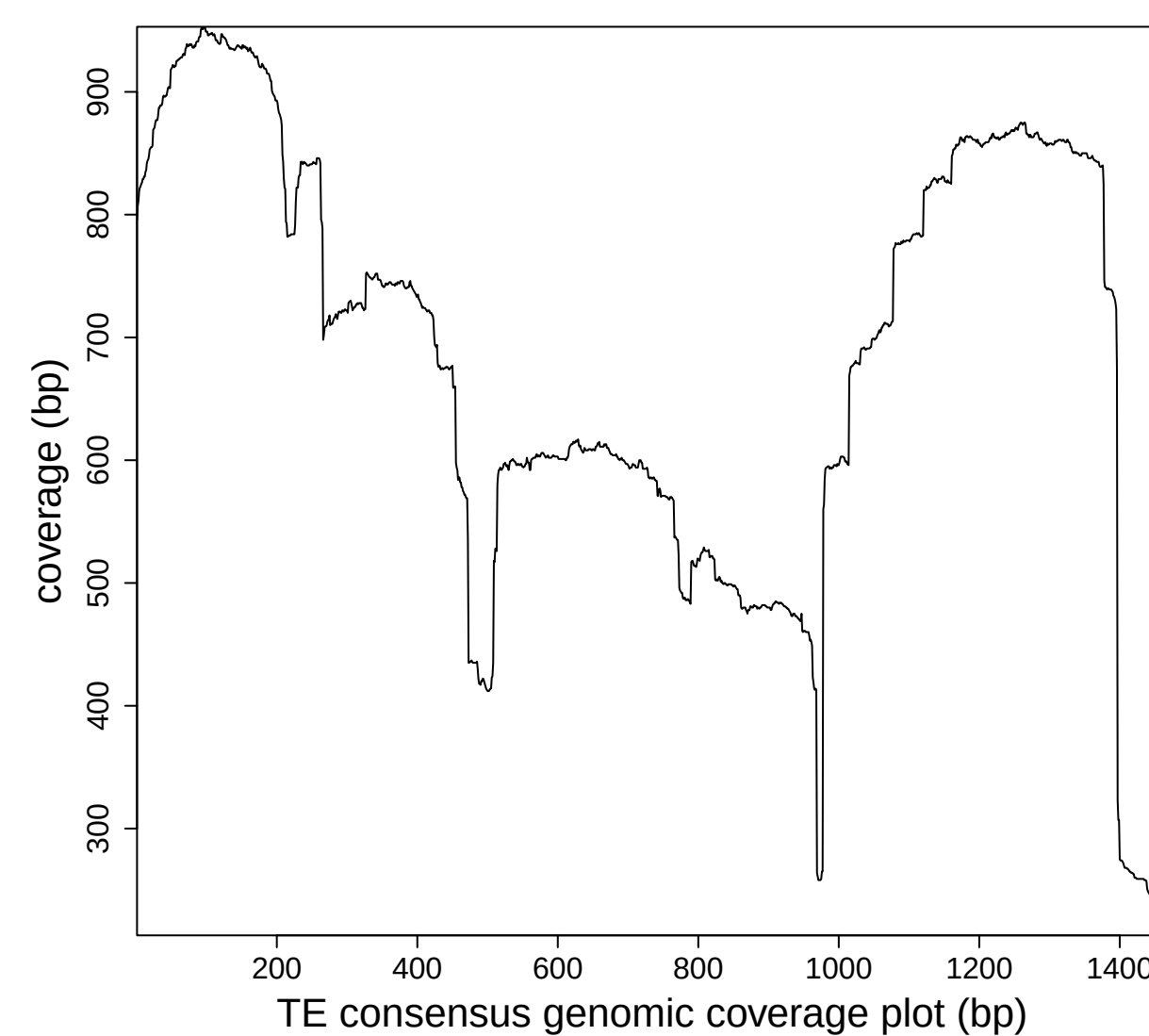
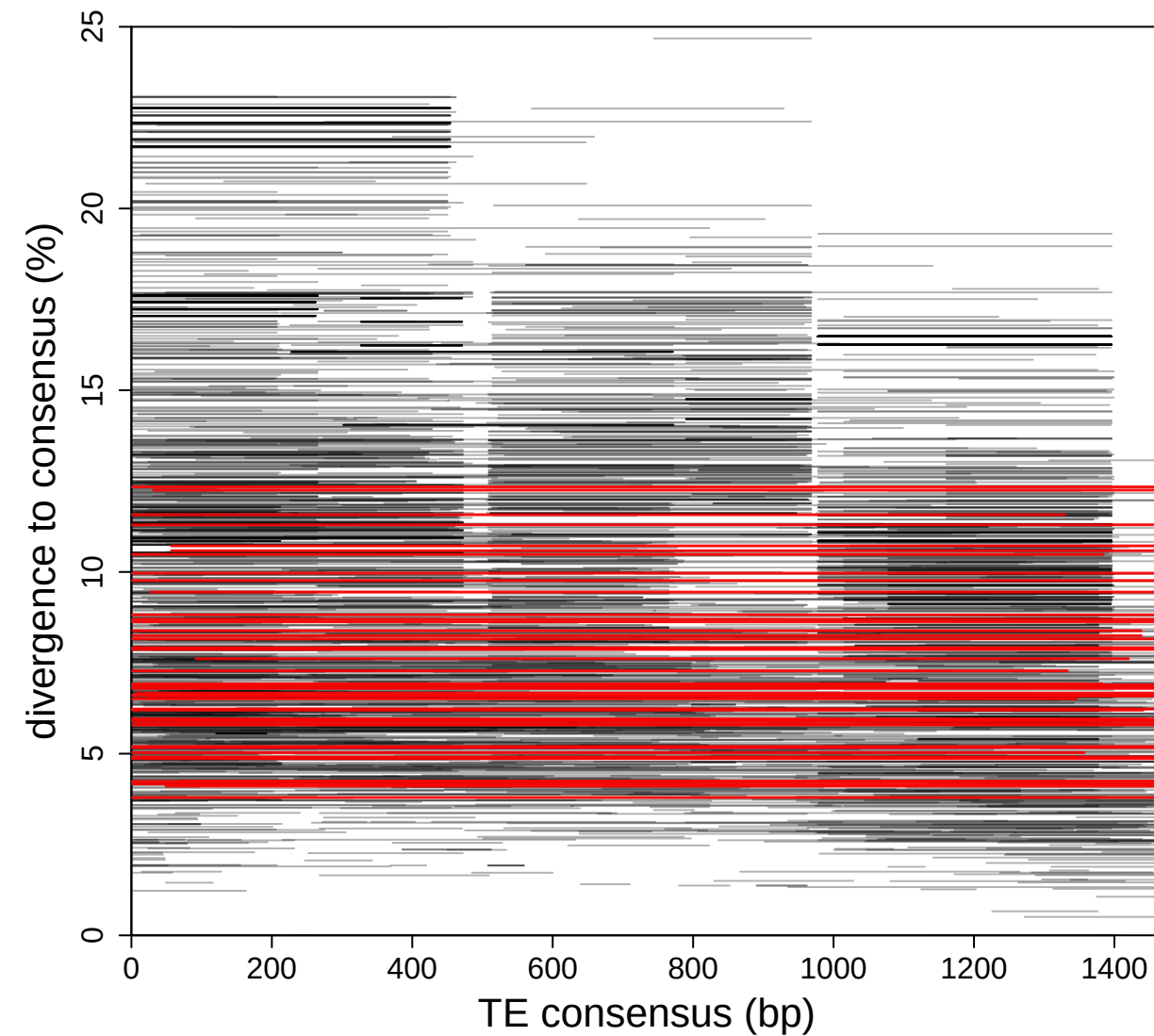


.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 2265bp; fragments: 40114; full length: 0 (>=2038.5bp)

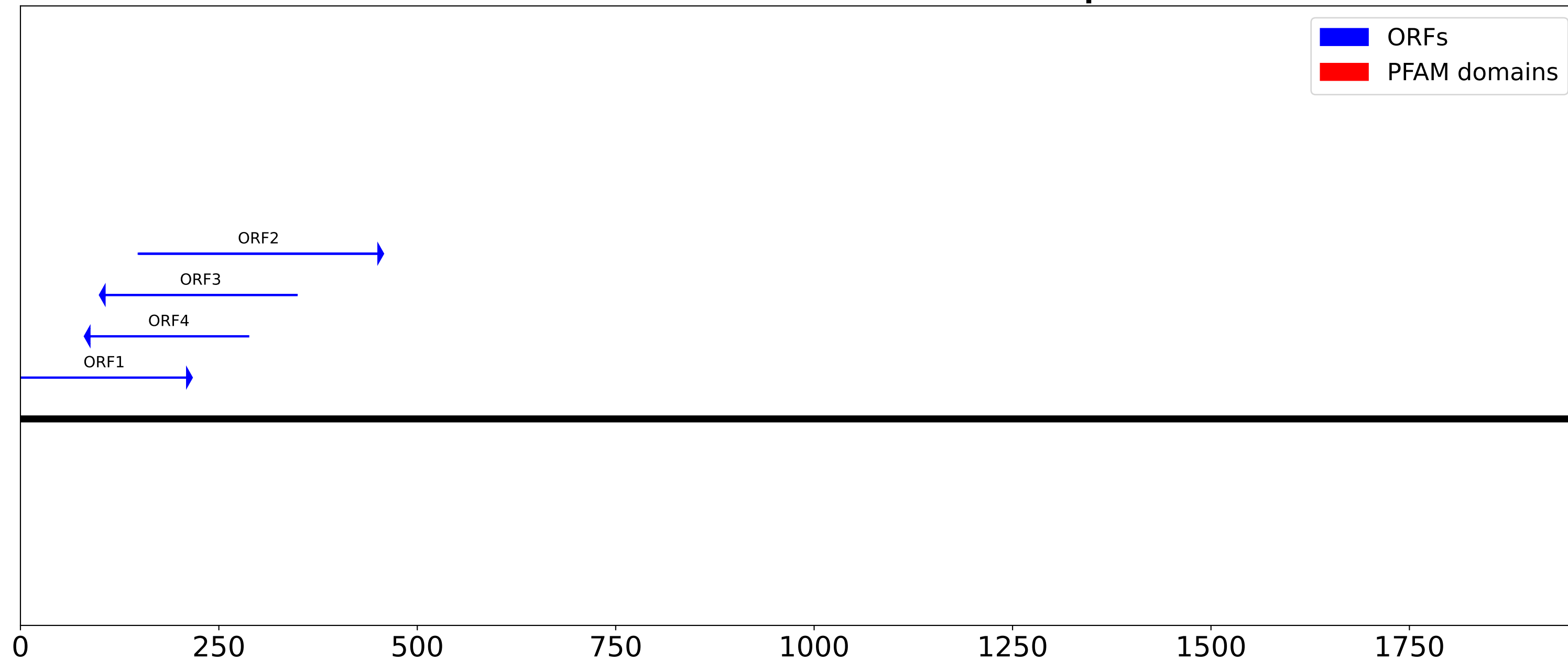
After TEtrimmer Extended plot Blue lines are boundaries



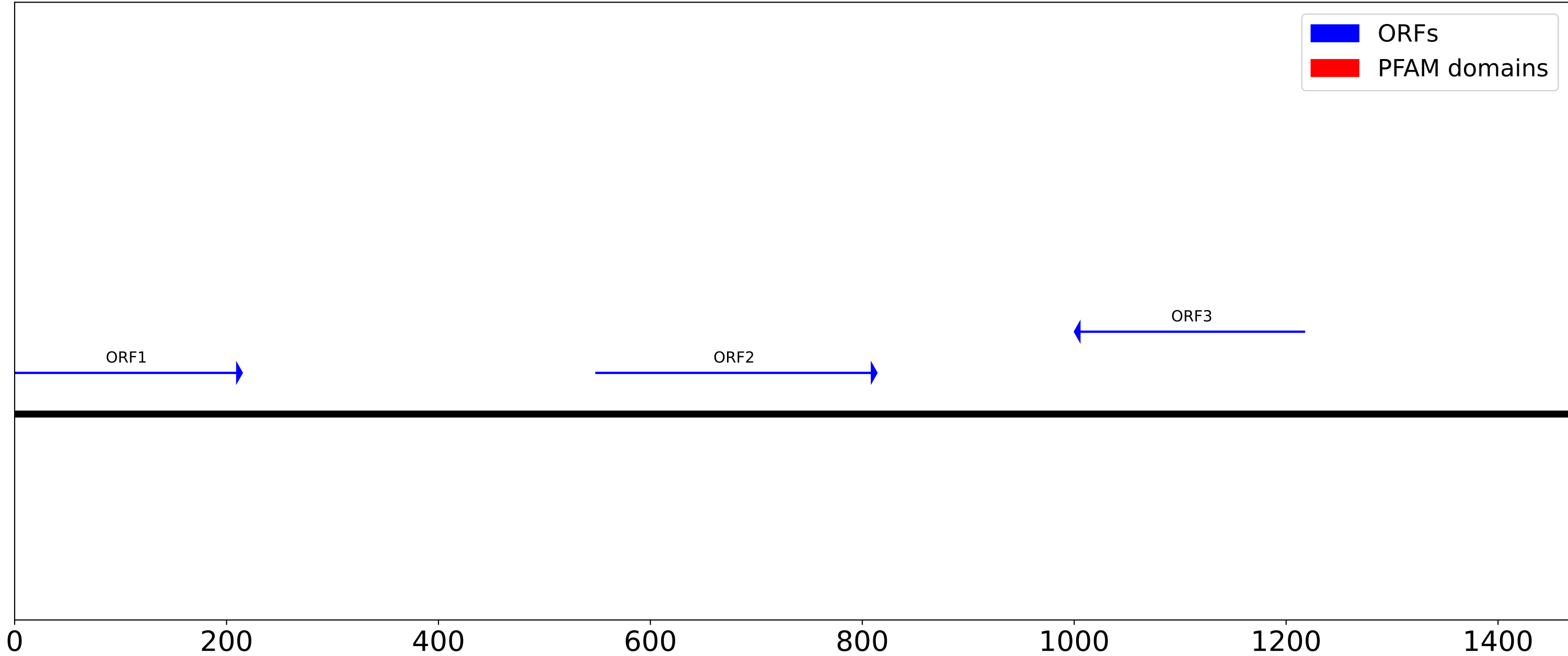
TE: rnd_5_family_79
size: 1468bp; fragments: 2613; full length: 45 (≥ 1321.2 bp)



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

