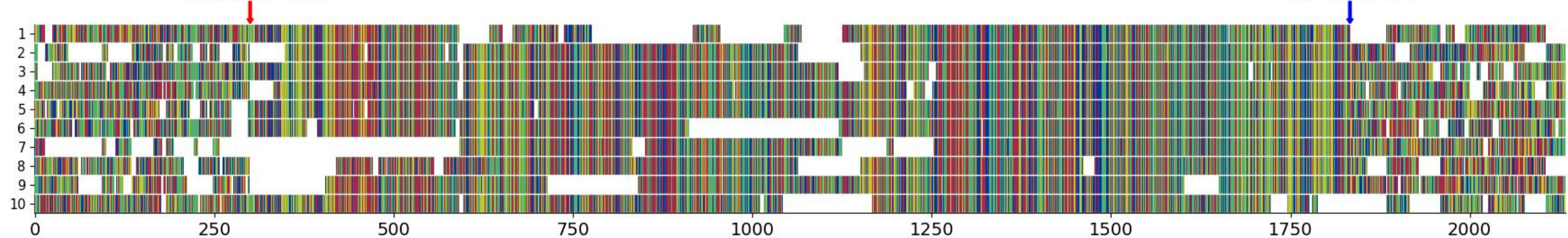


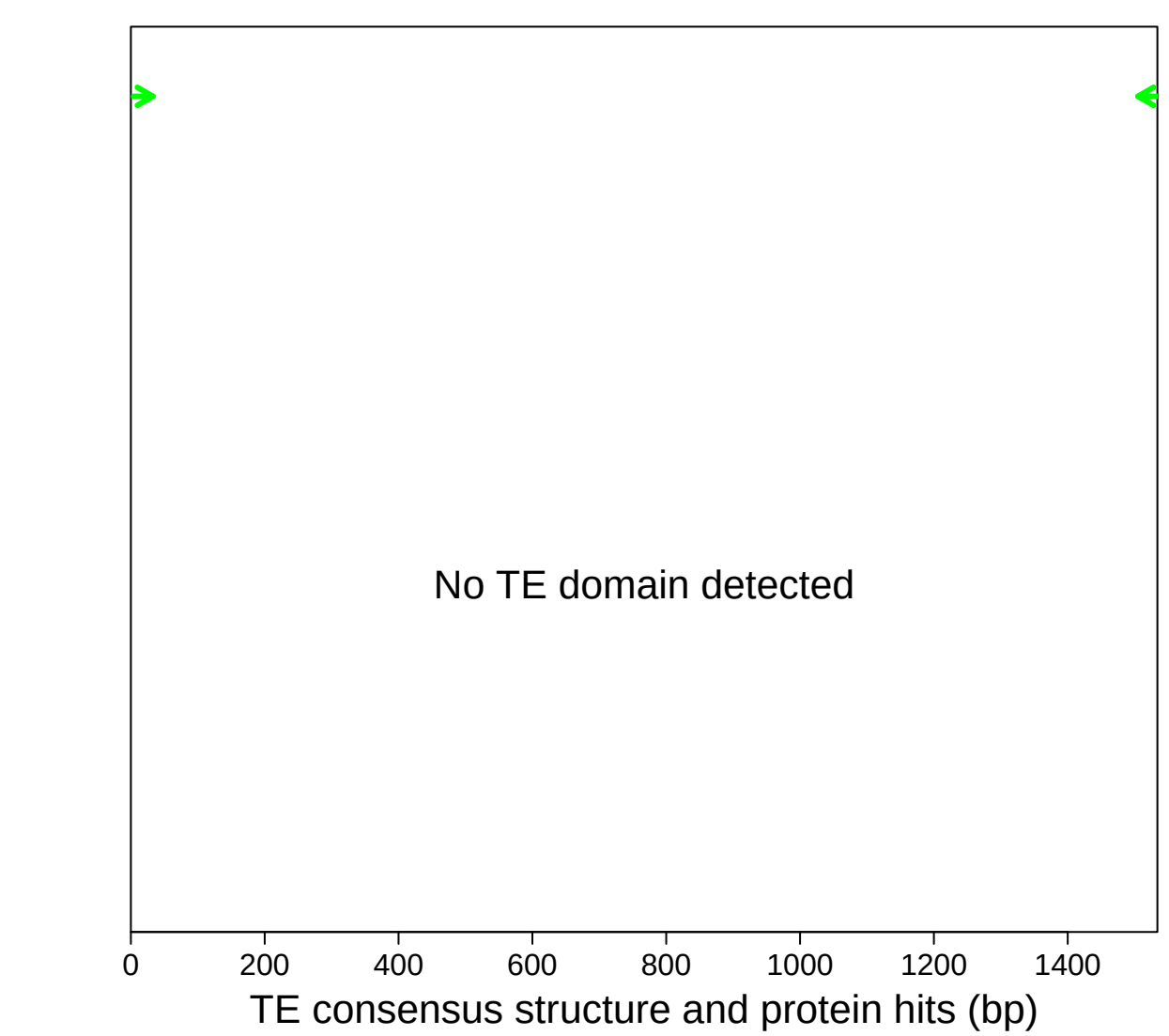
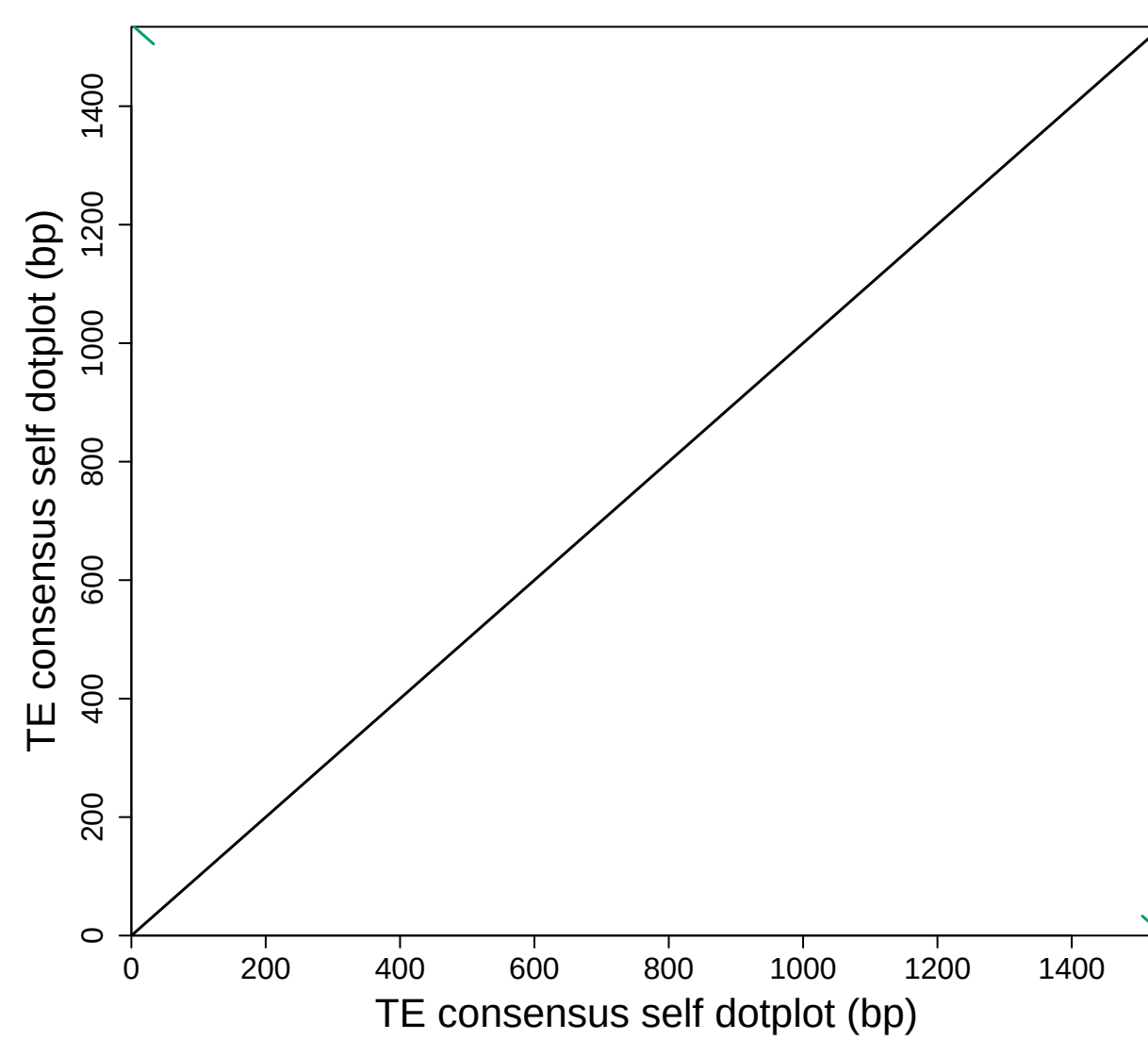
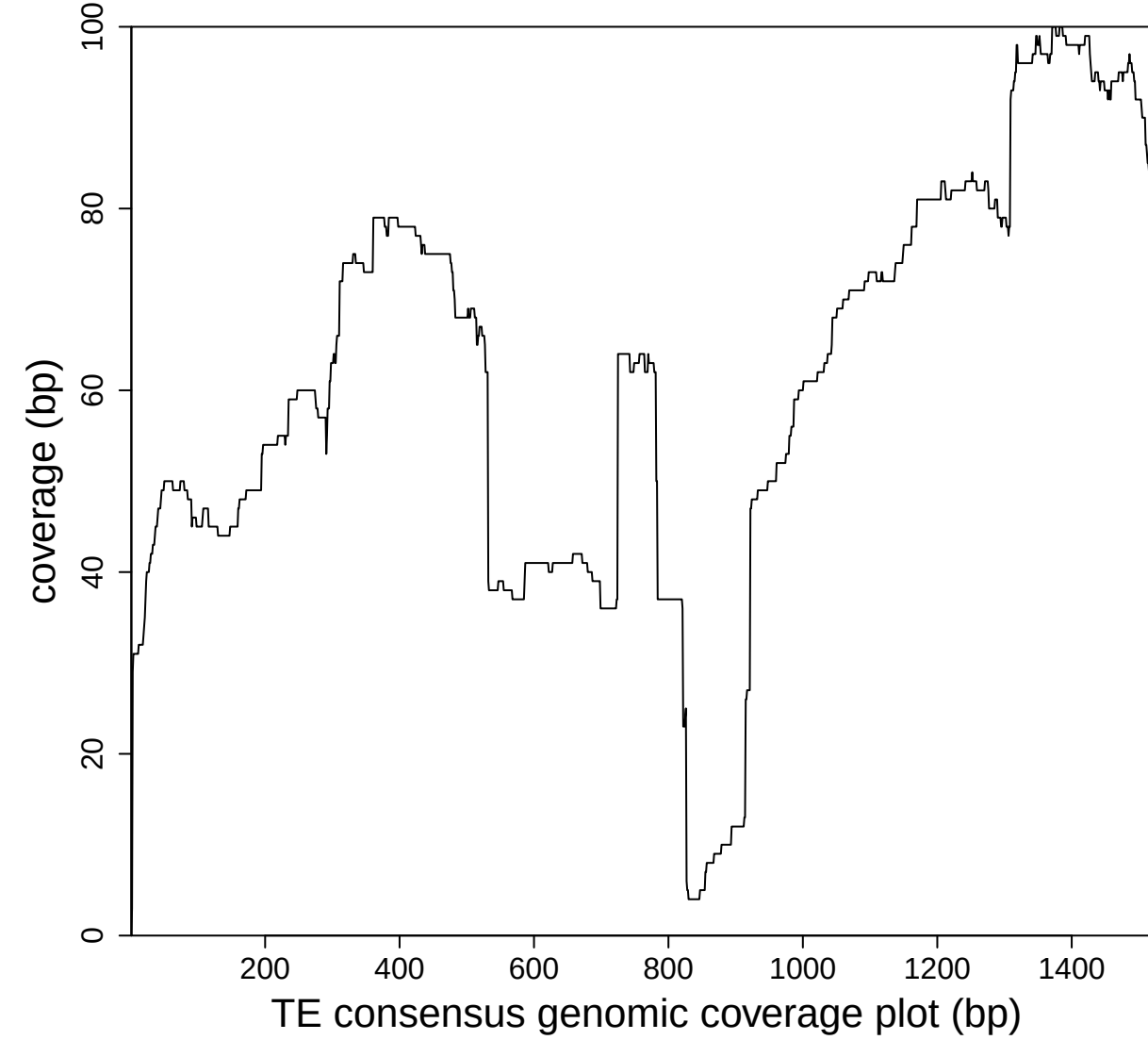
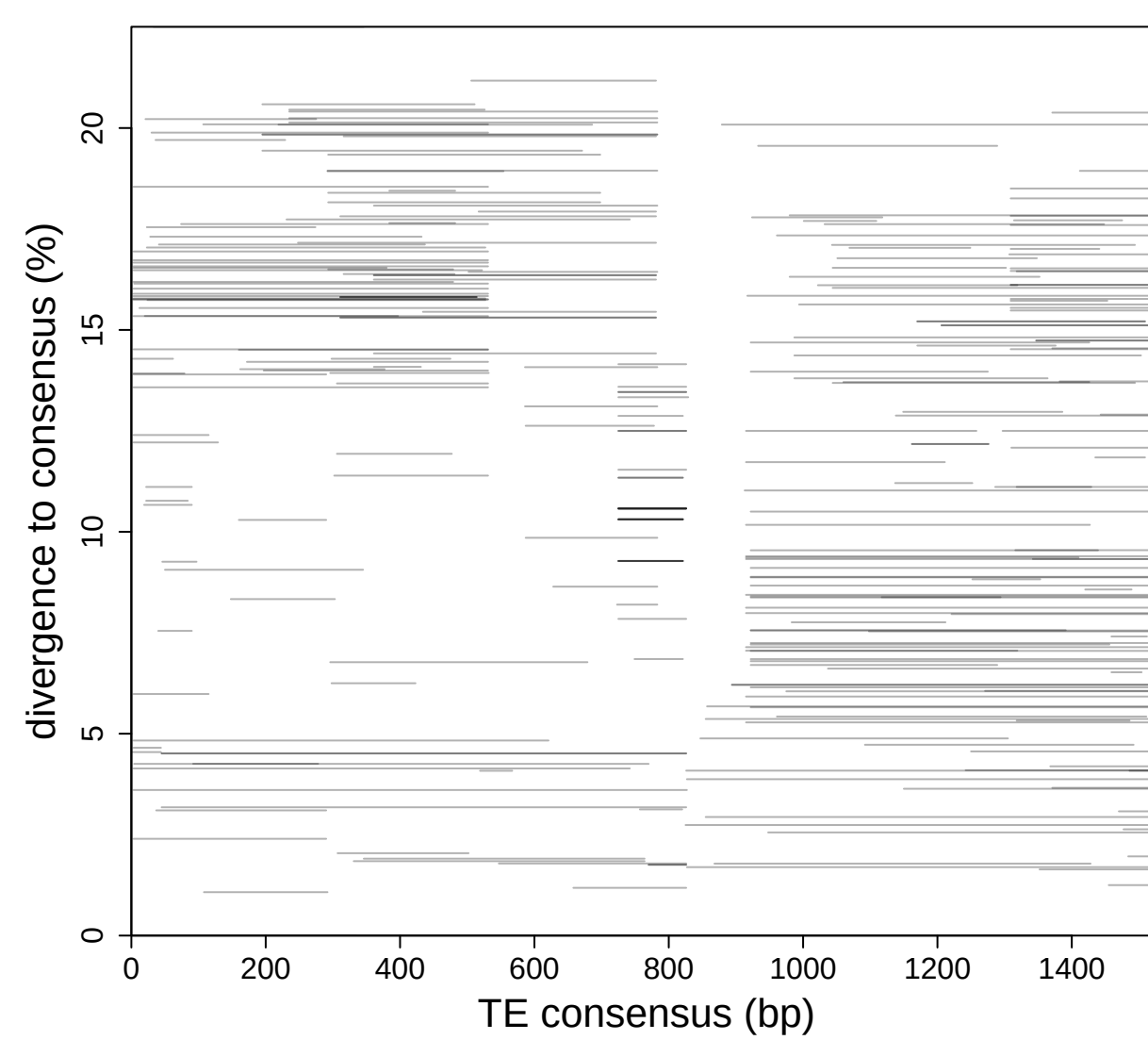
Start crop Point

End crop Point



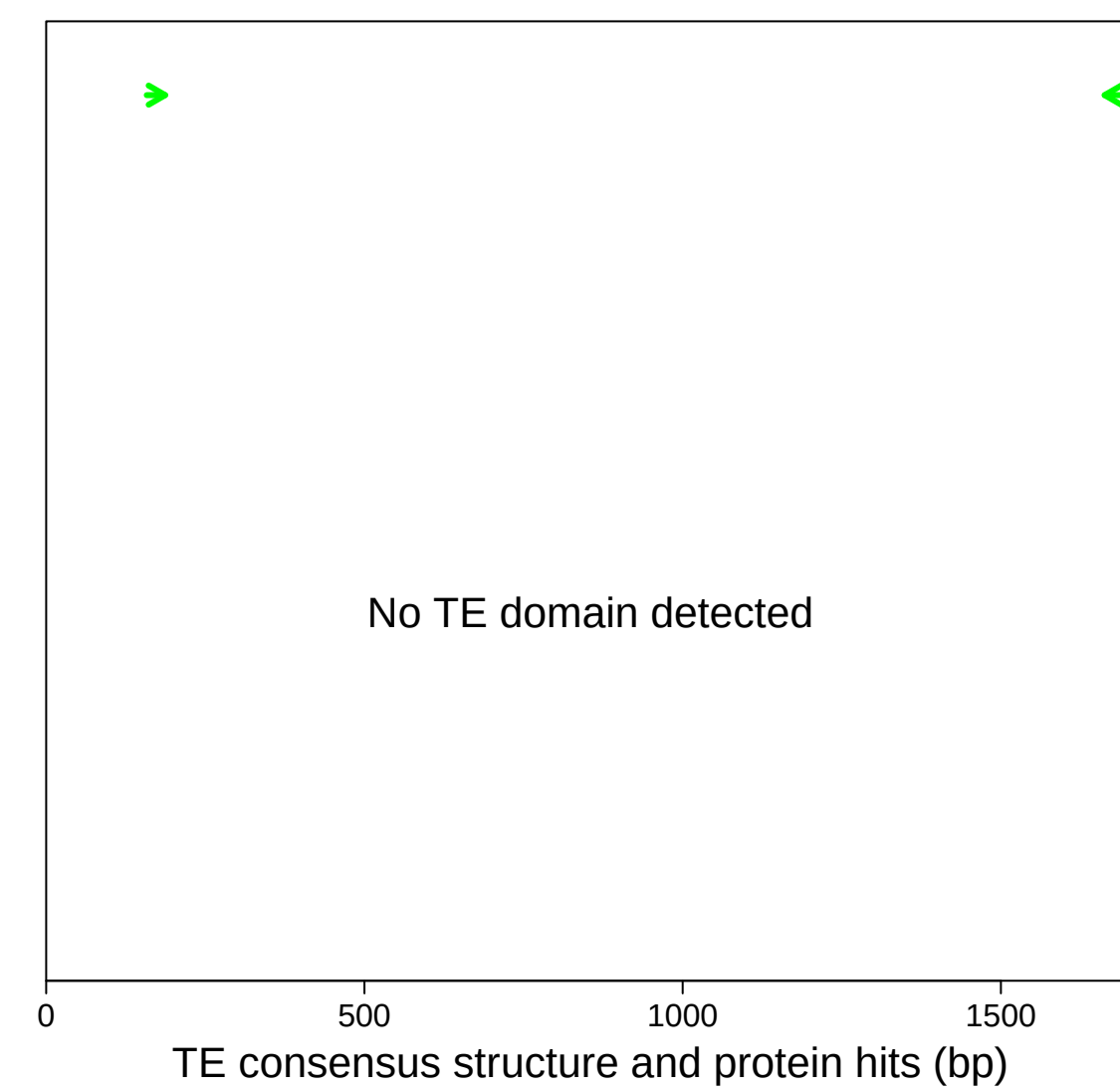
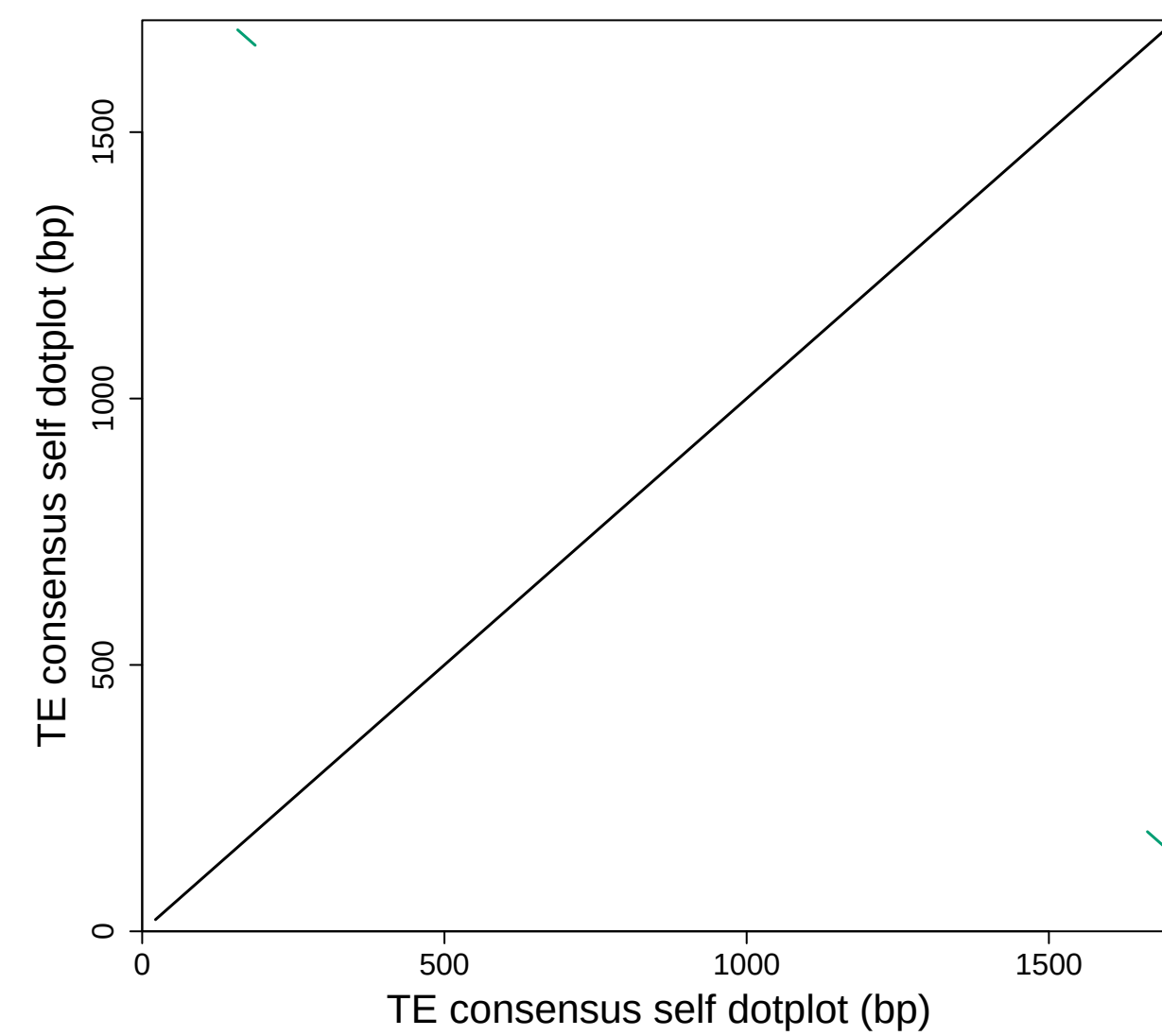
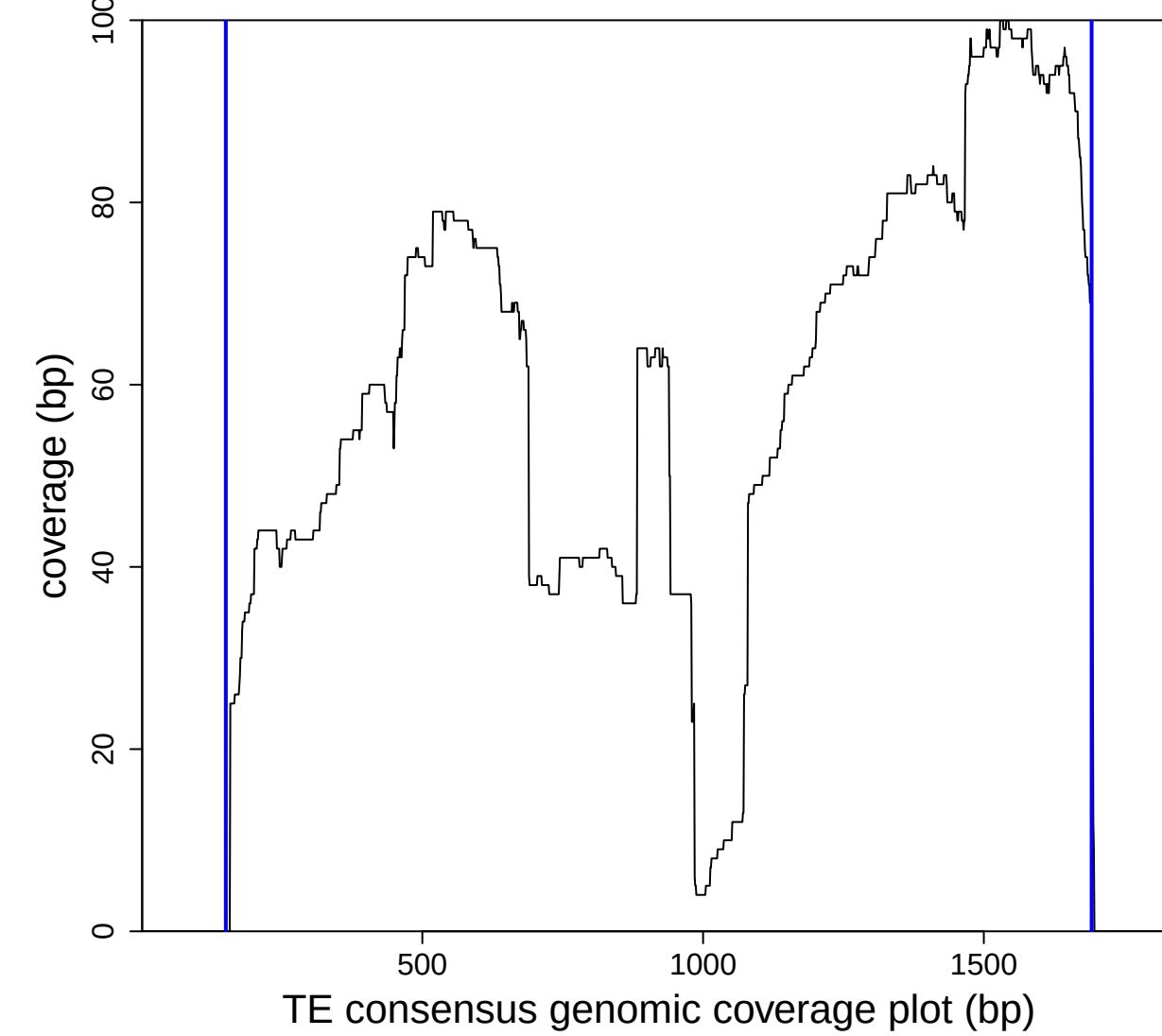
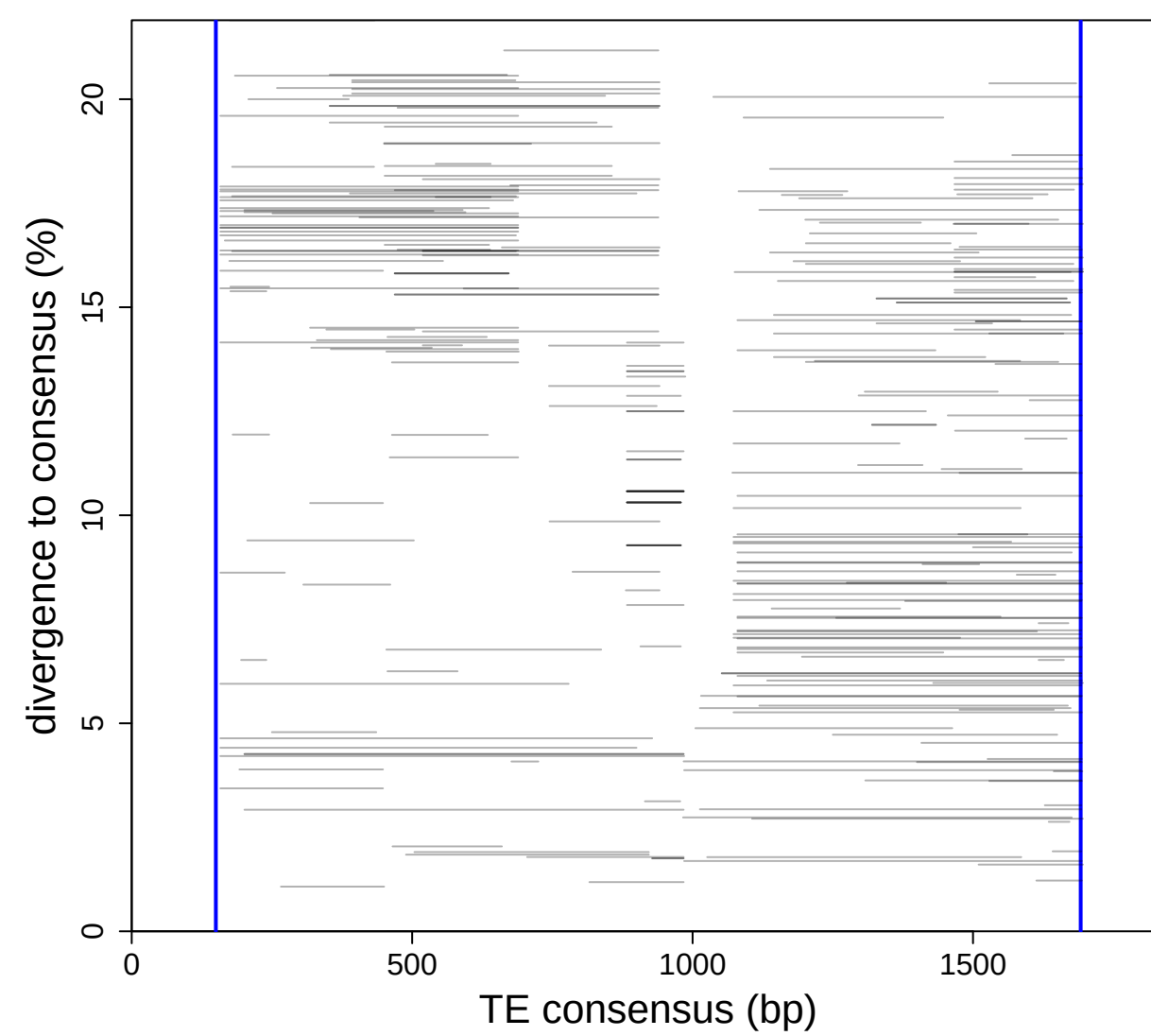
a.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1534bp; fragments: 283; full length: 0 (>=1380.6bp)

After TEtrimmer 1534 bp

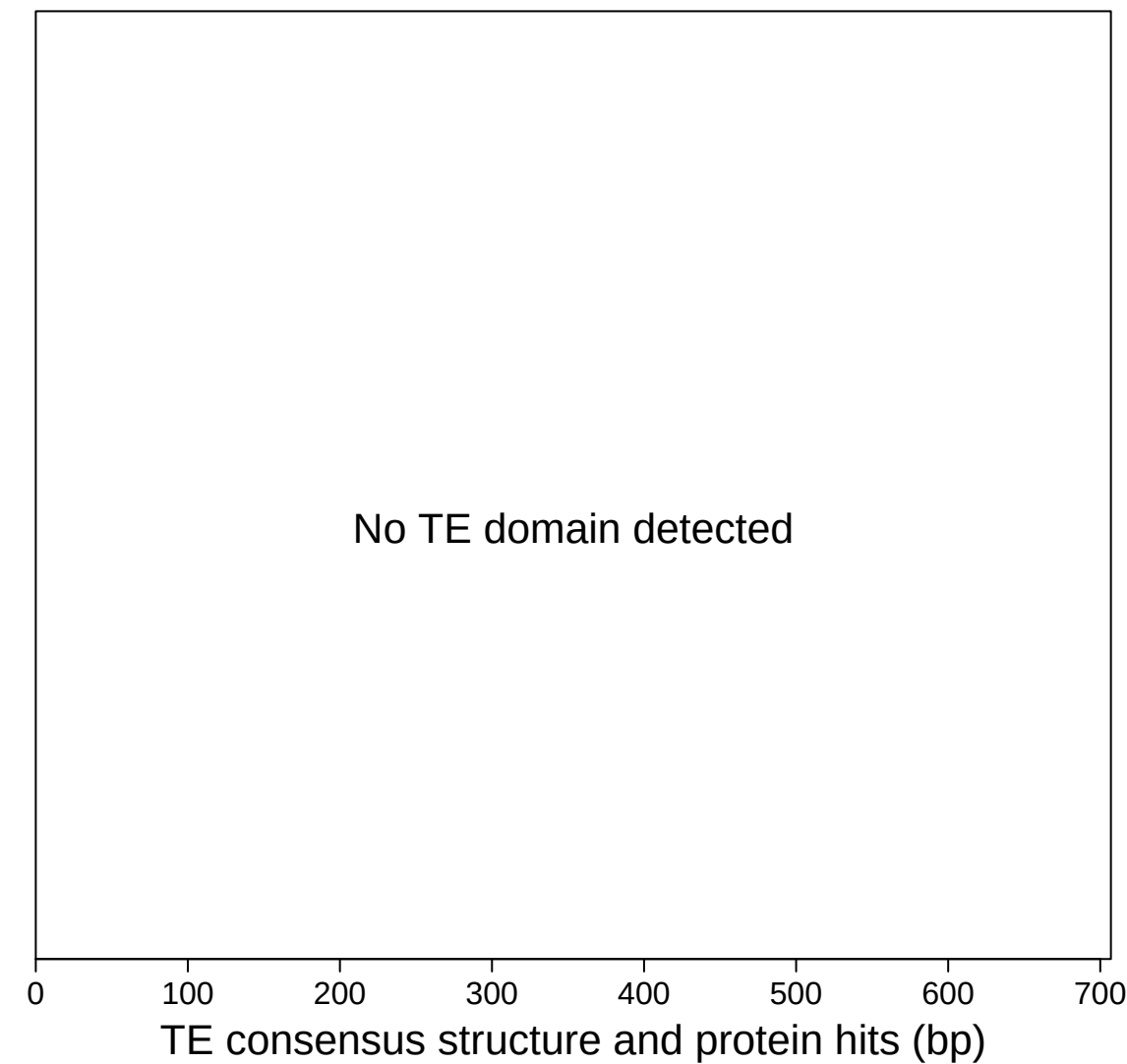
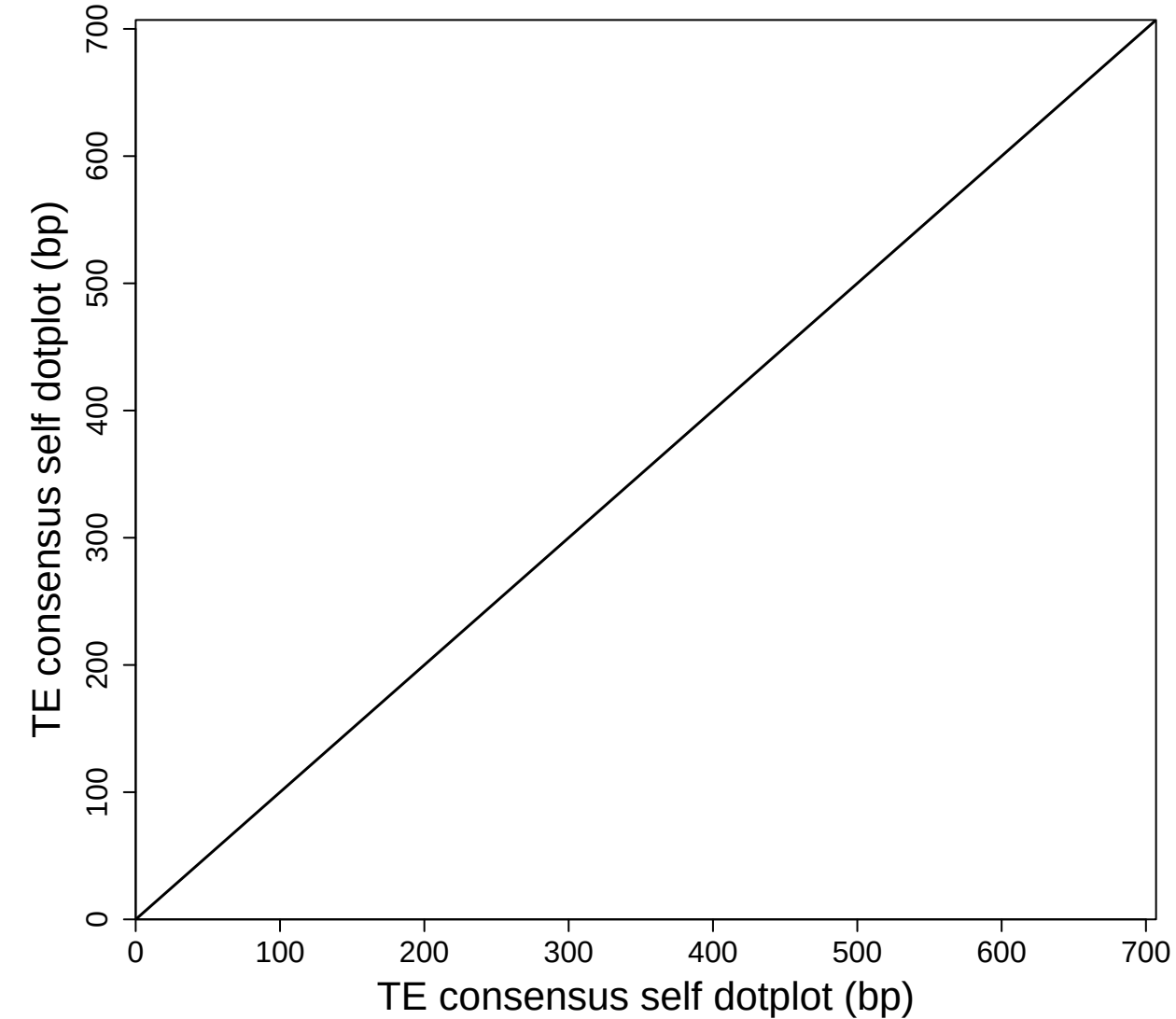
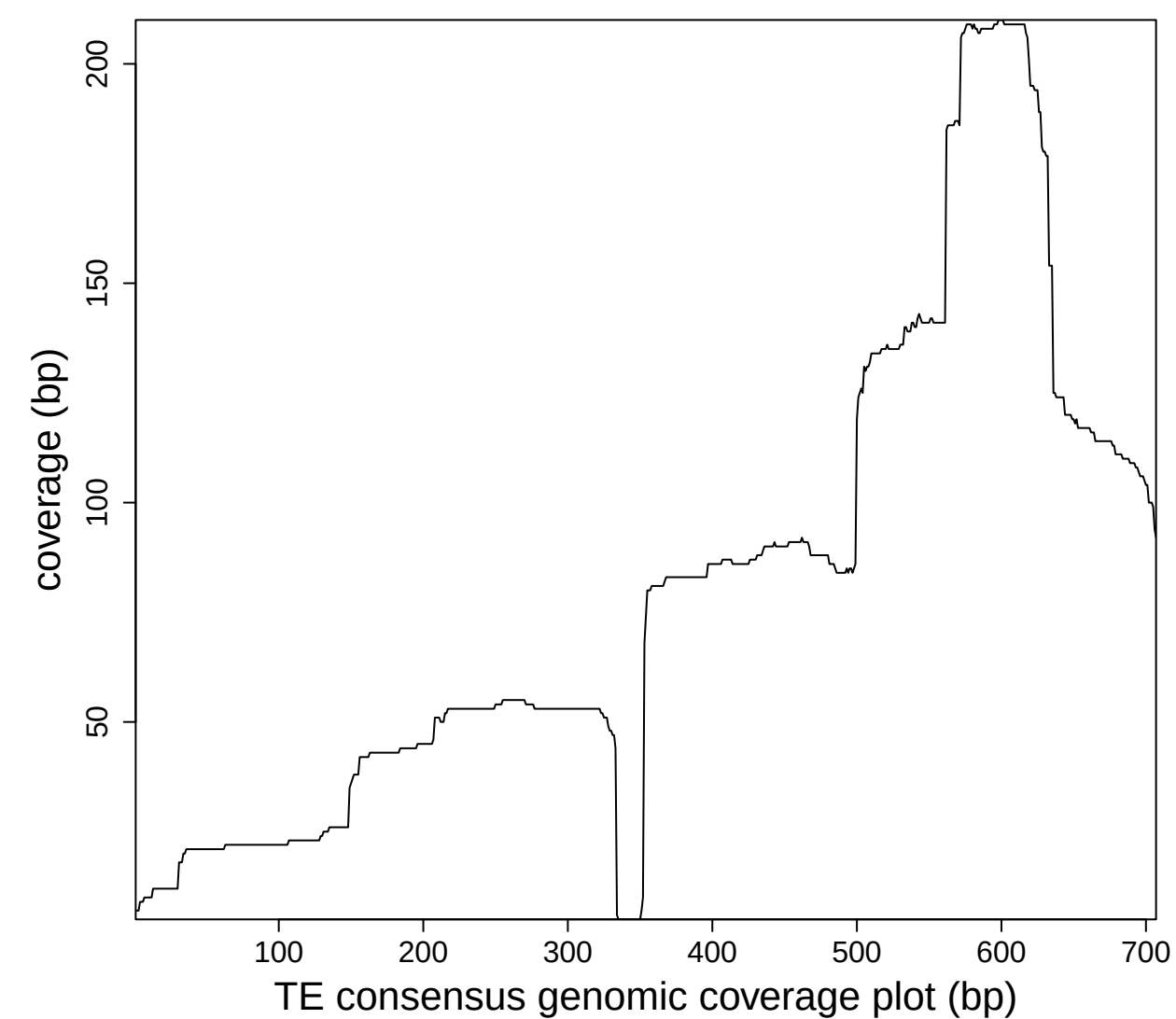
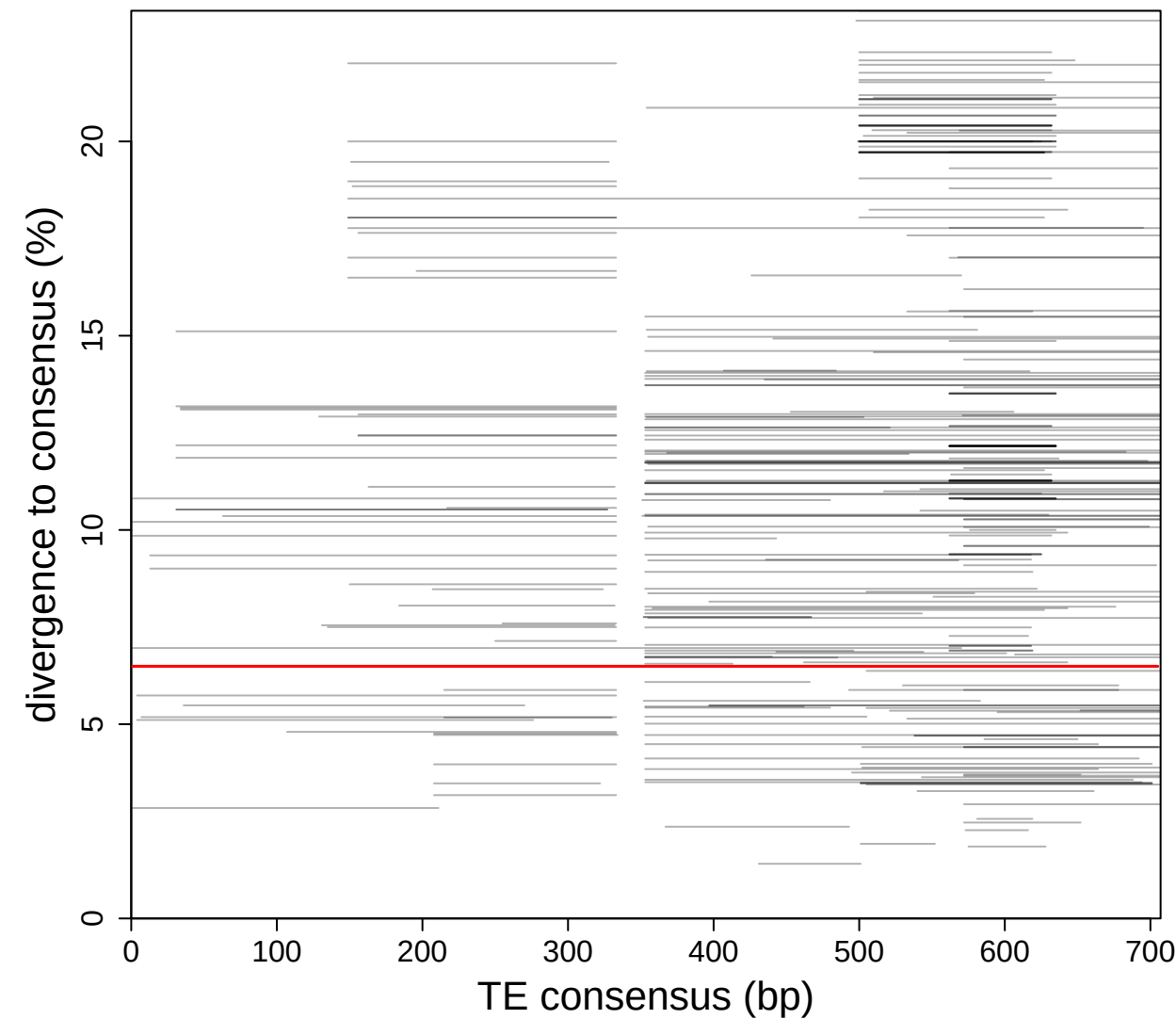


5.fasta.b.bed uf.bed g_4.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 1842bp; fragments: 276; full length: 0 (>=1657.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

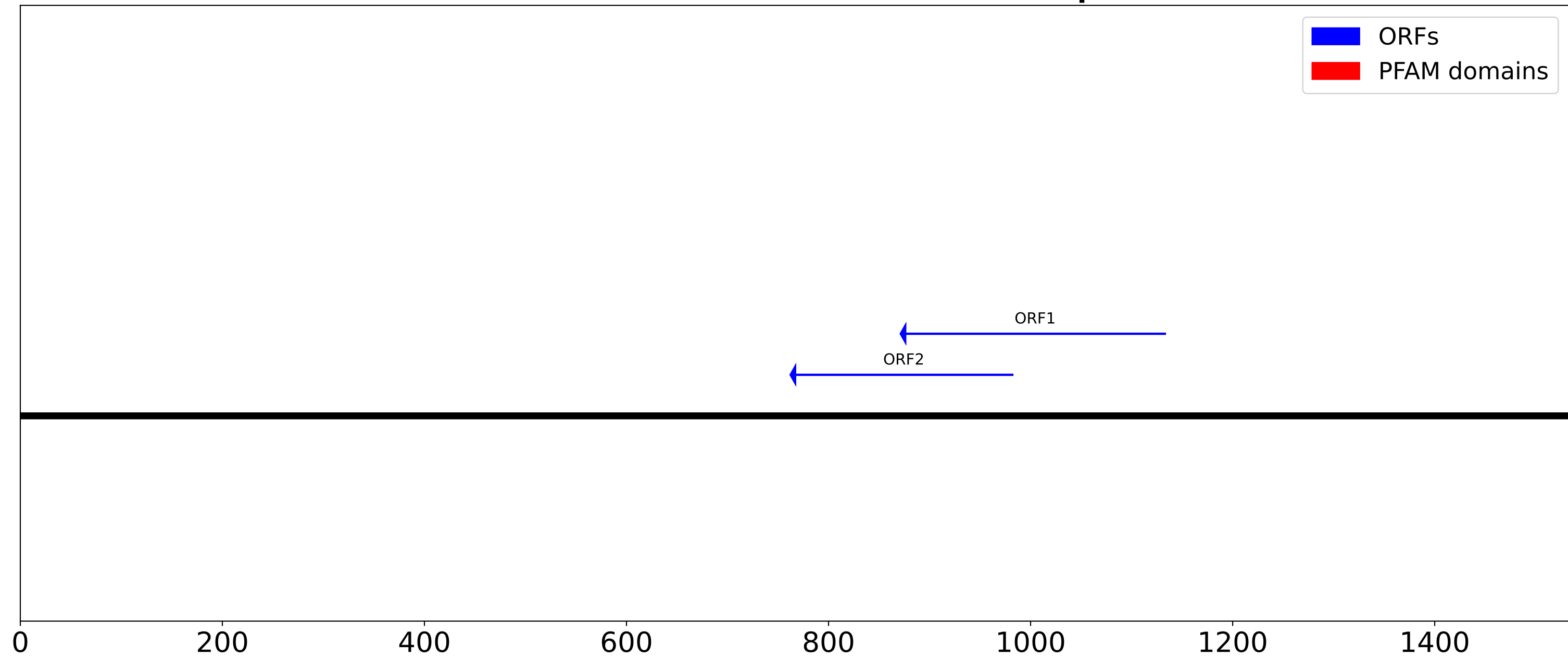


TE: rnd_5_family_1395
size: 707bp; fragments: 292; full length: 2 (>=636.3bp)

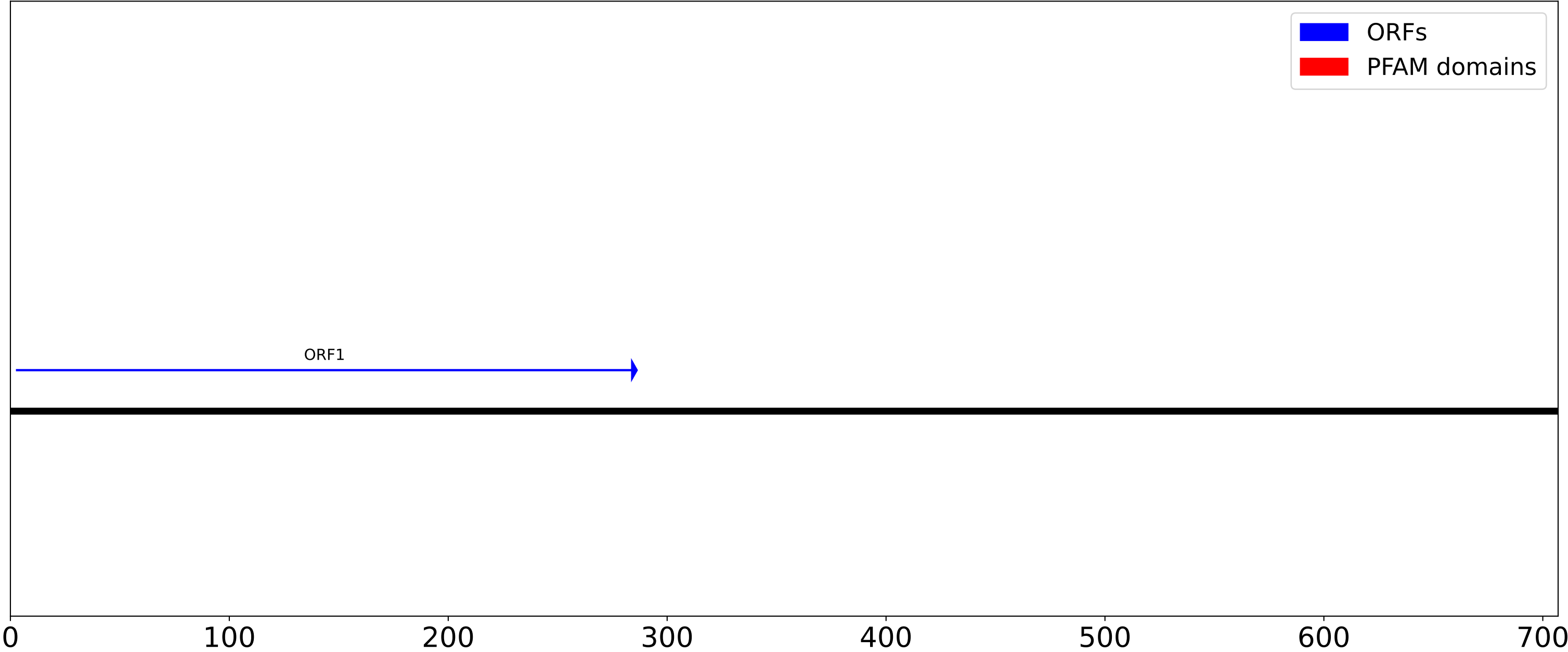


Before TEtrimmer 707 bp

After TEtrimmer ORF and PFAM domain plot



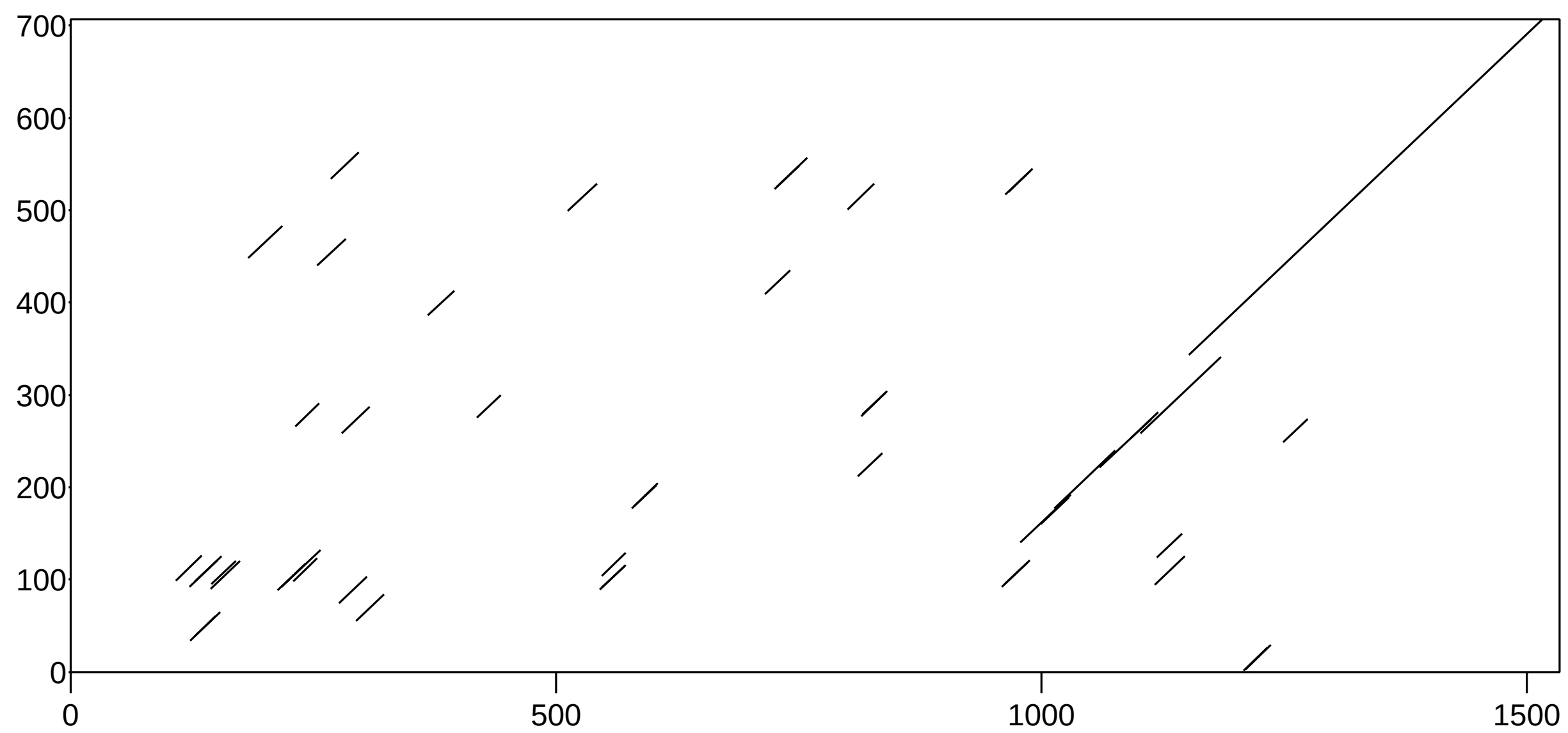
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)