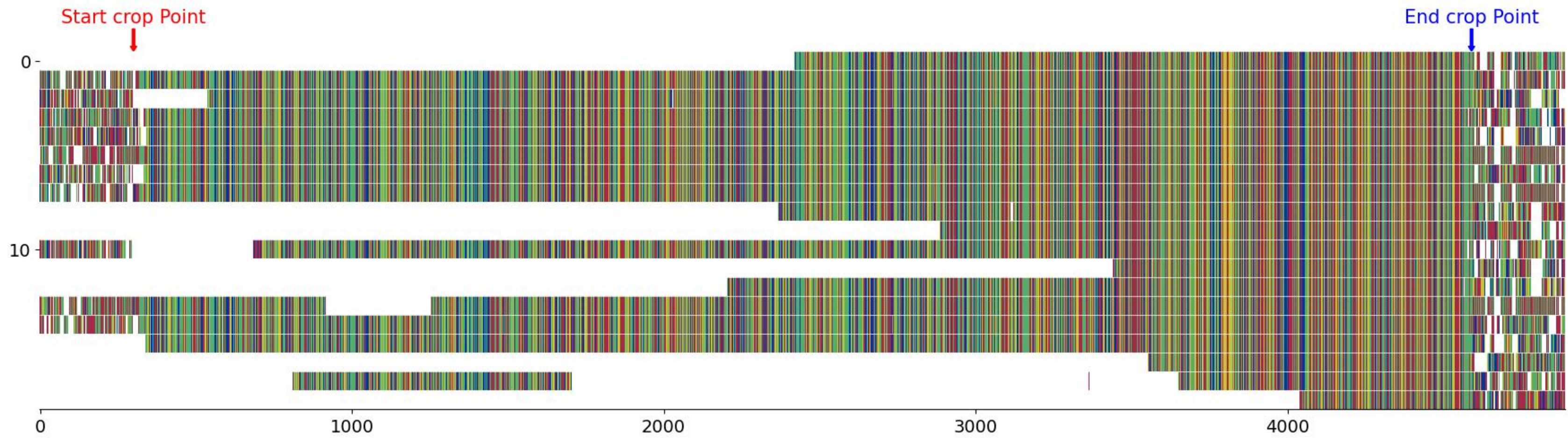


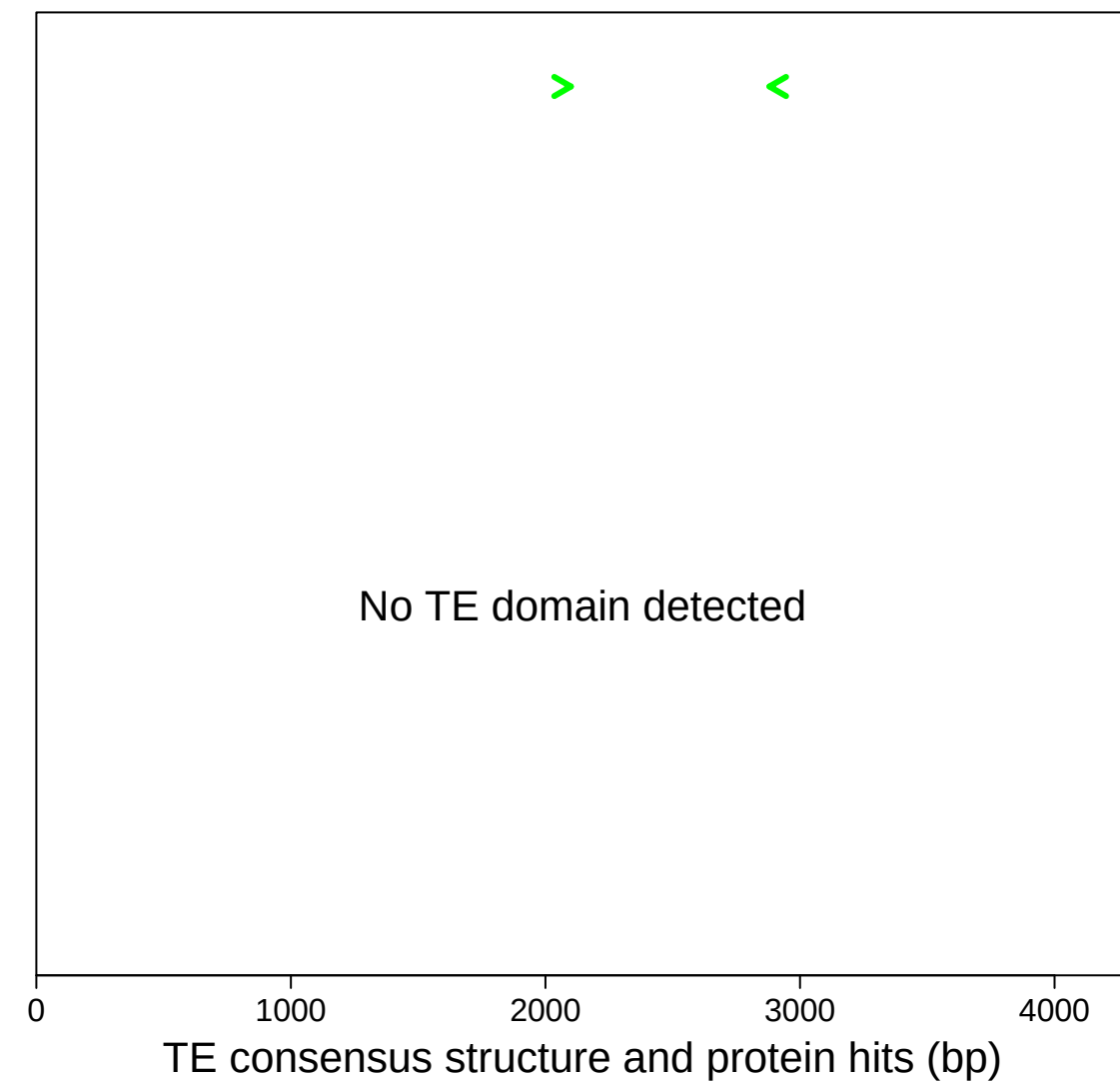
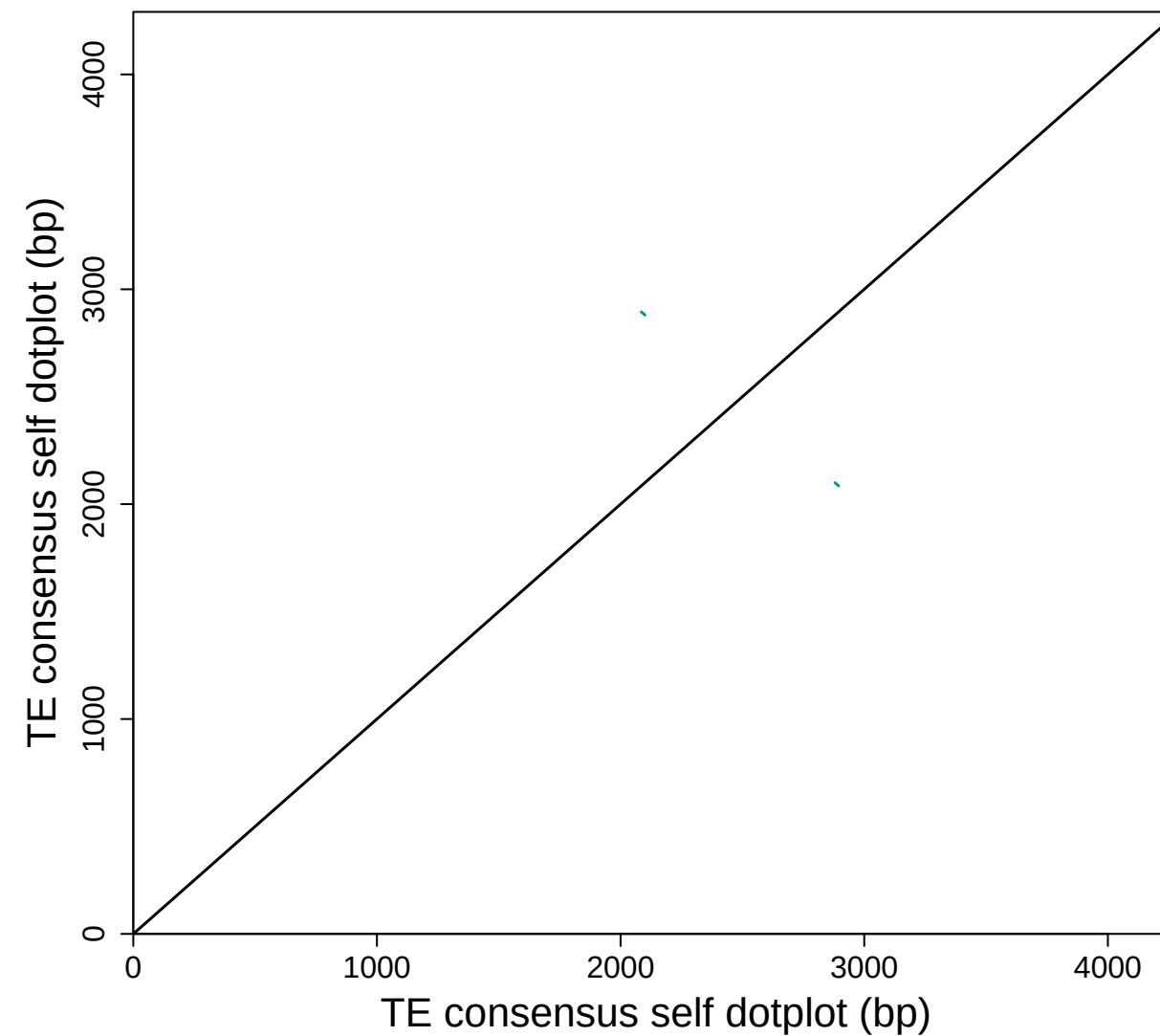
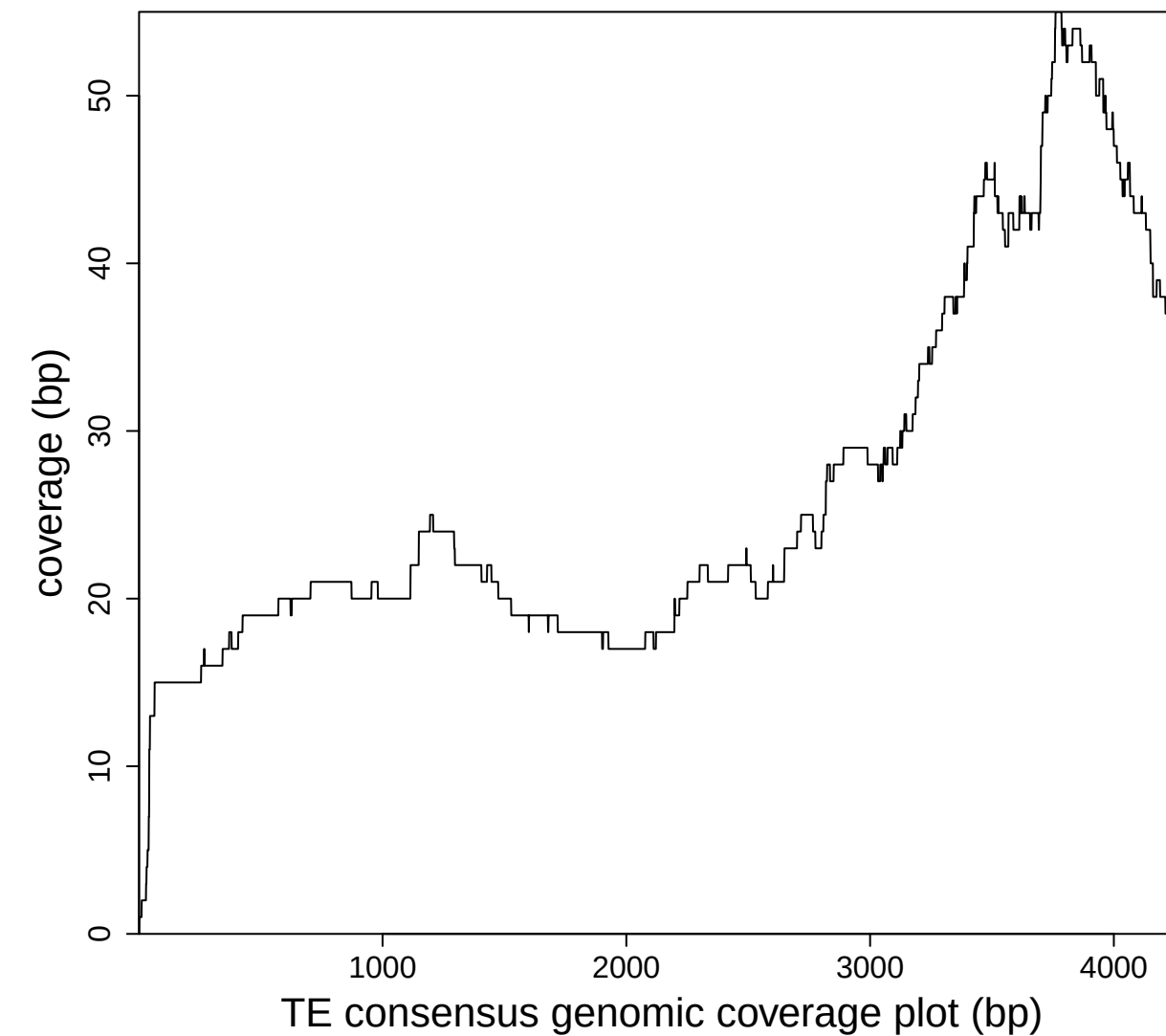
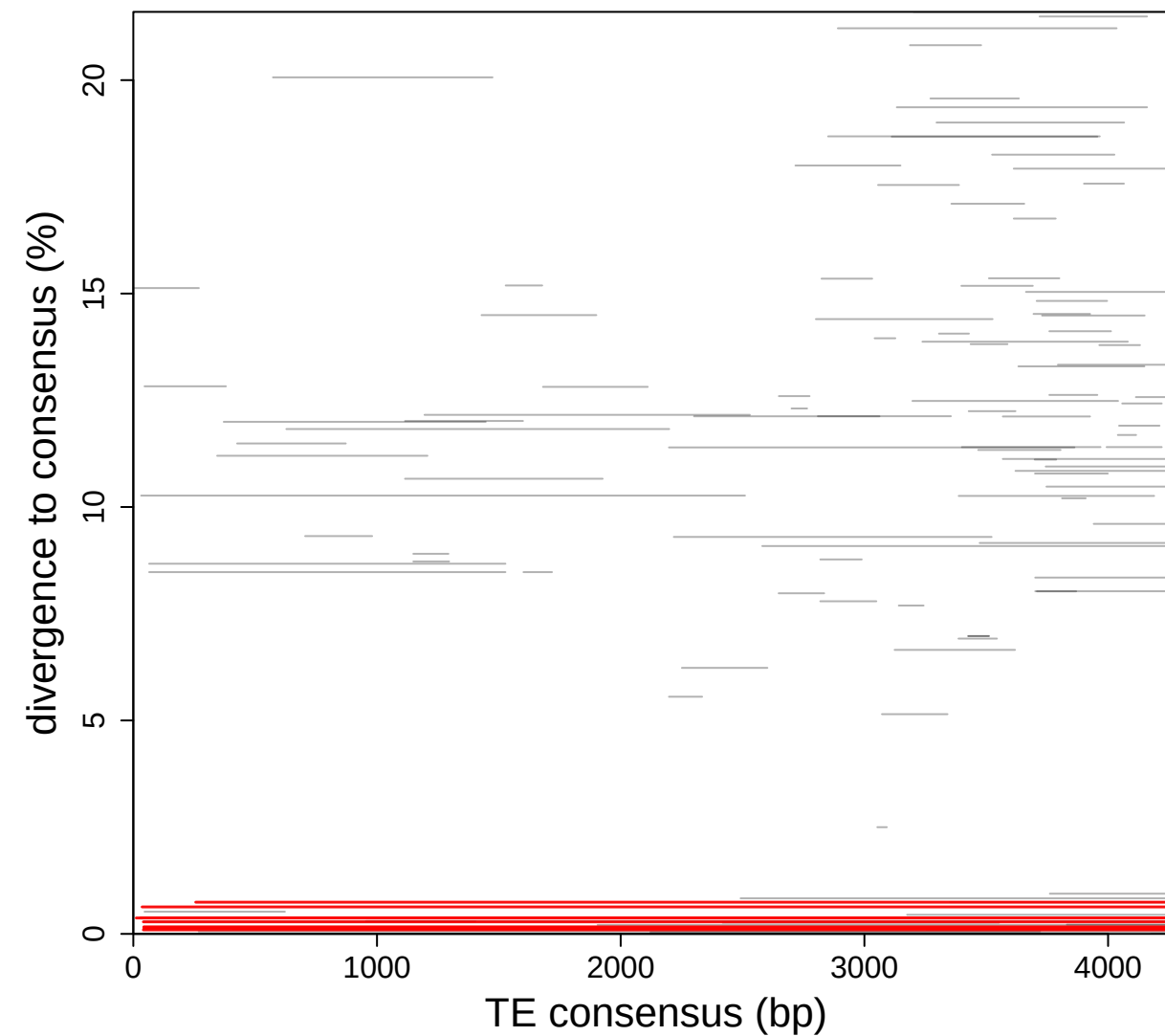
Start crop Point

End crop Point



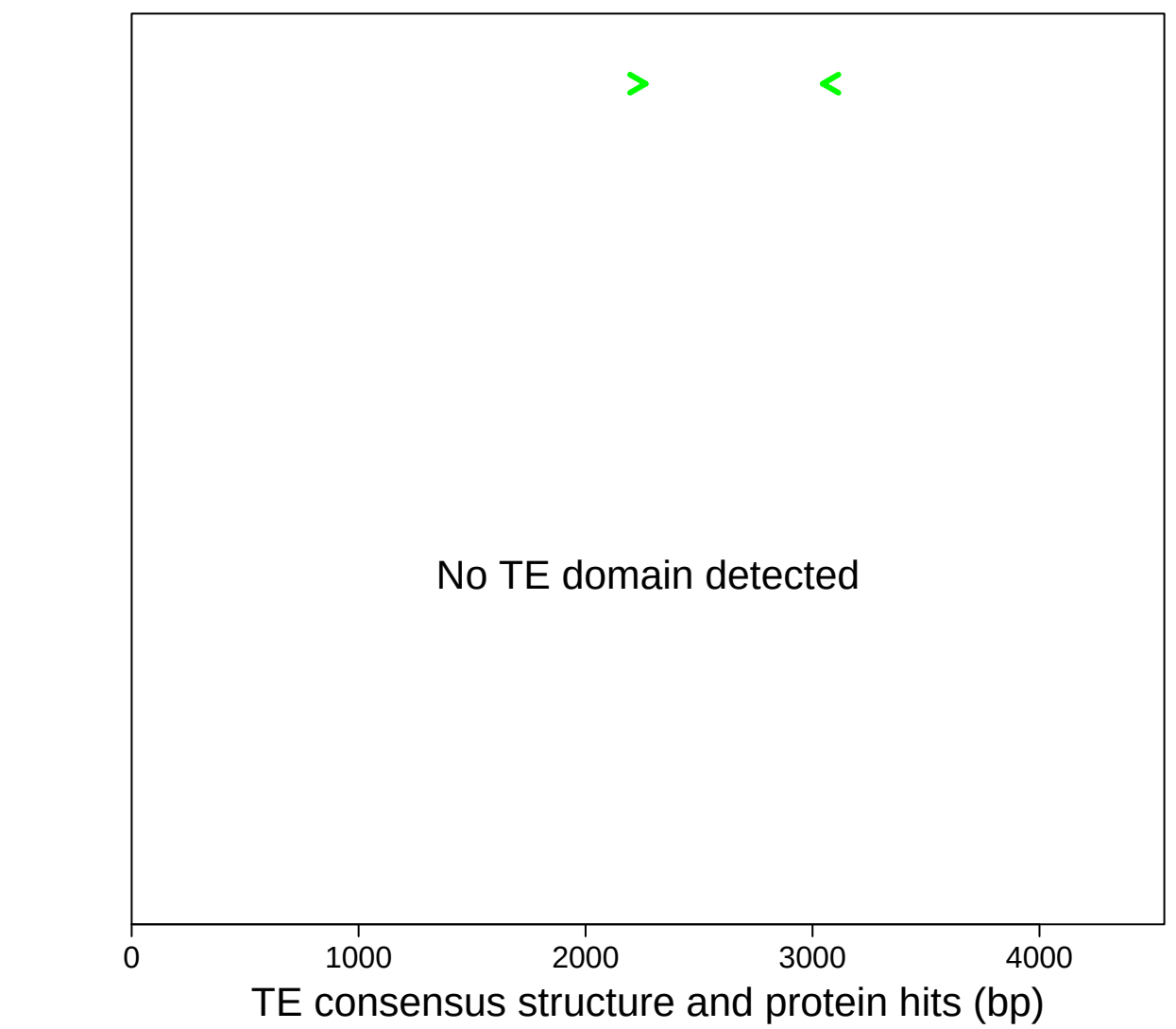
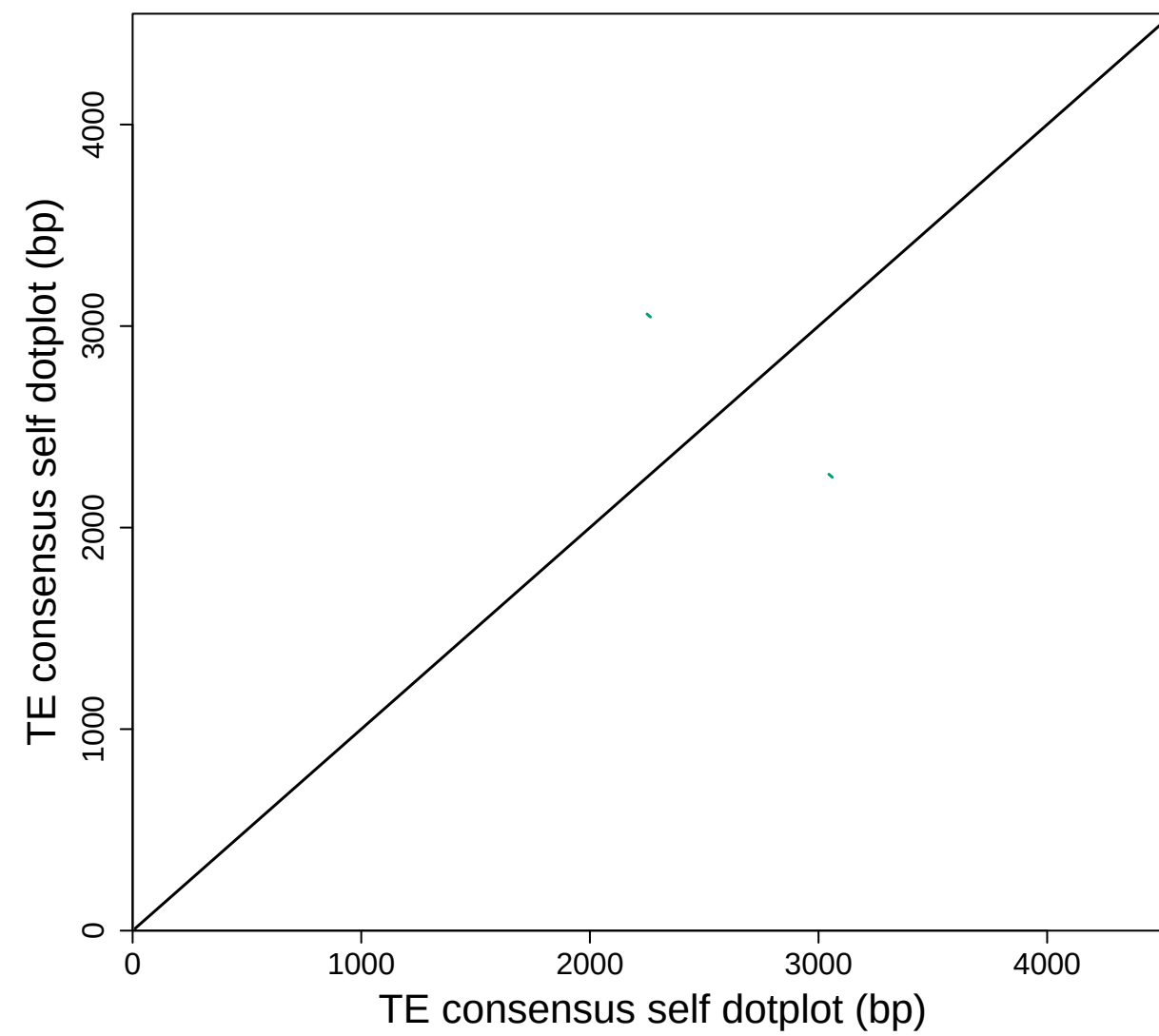
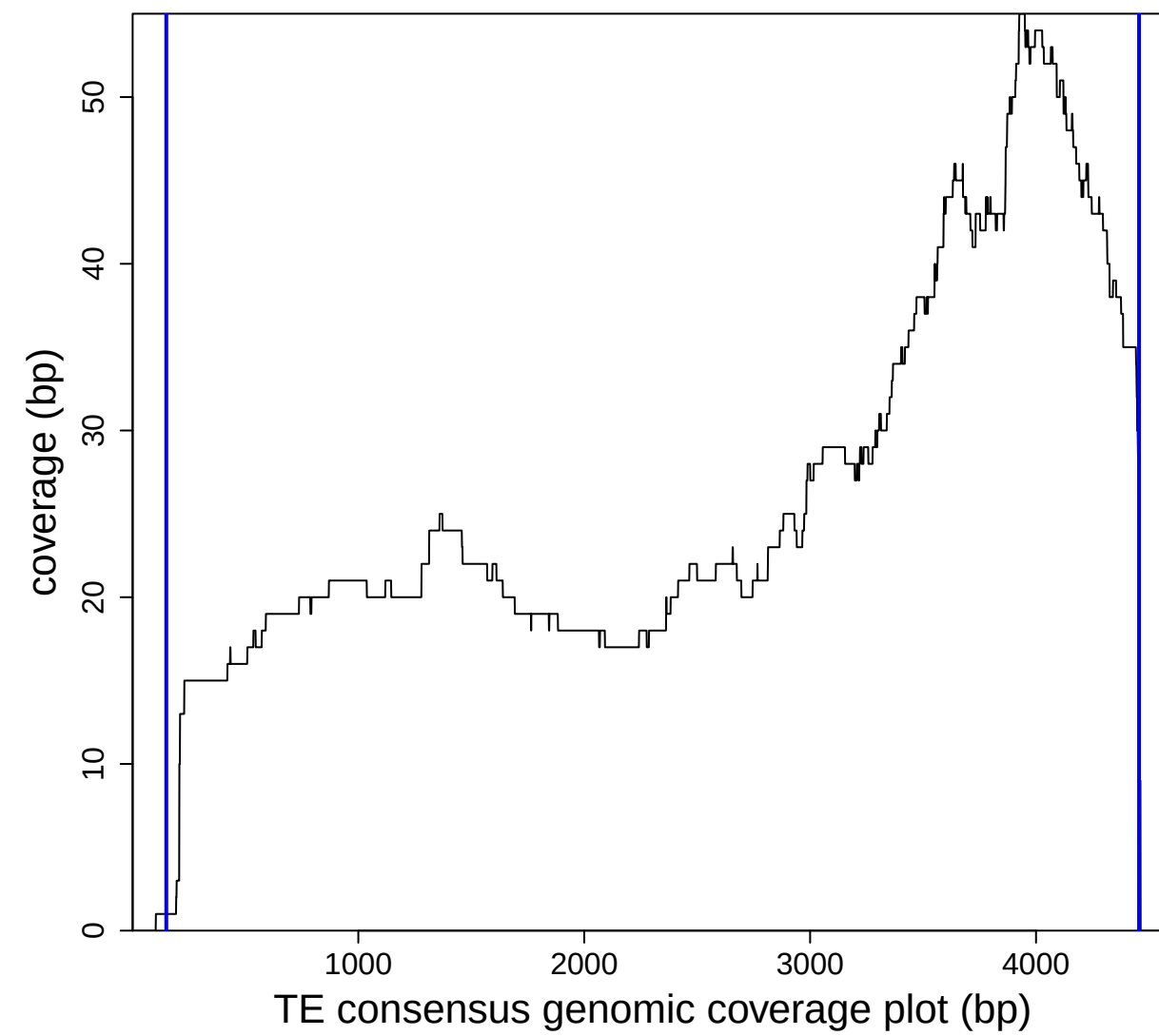
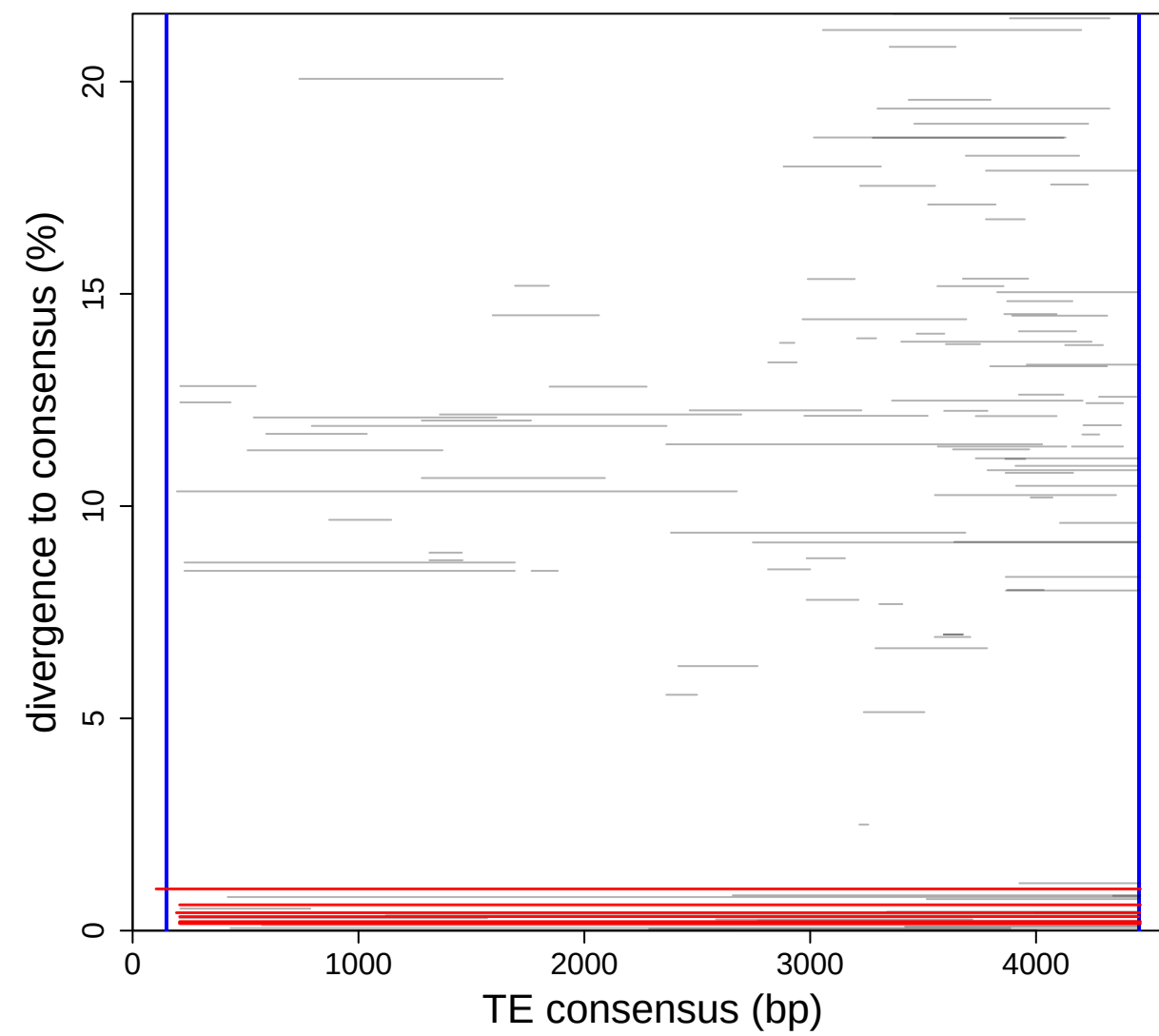
bed_uf.bed g_1.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa c
size: 4291bp; fragments: 121; full length: 10 (≥ 3861.9 bp)

After TEtrimmer 4291 bp

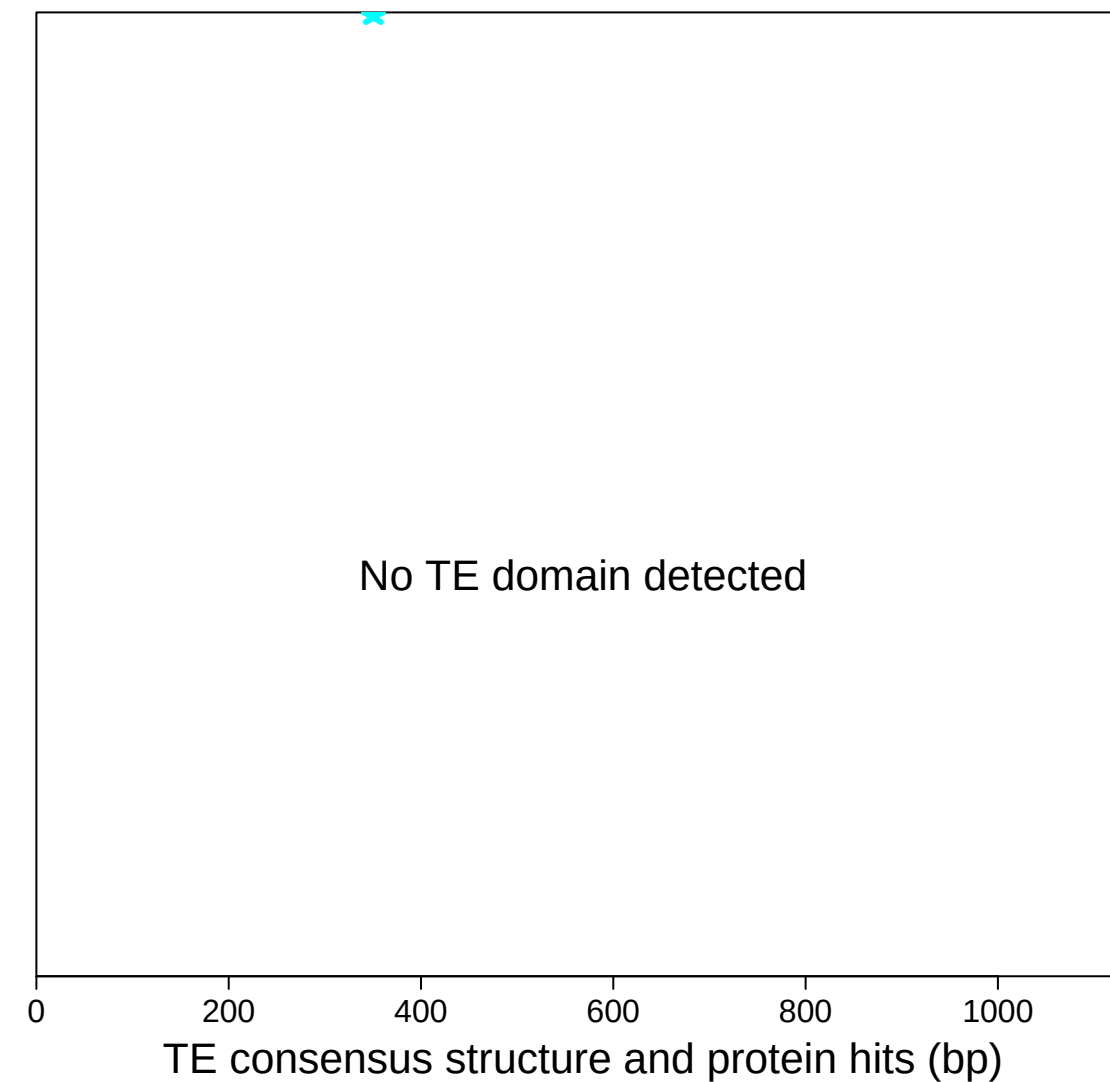
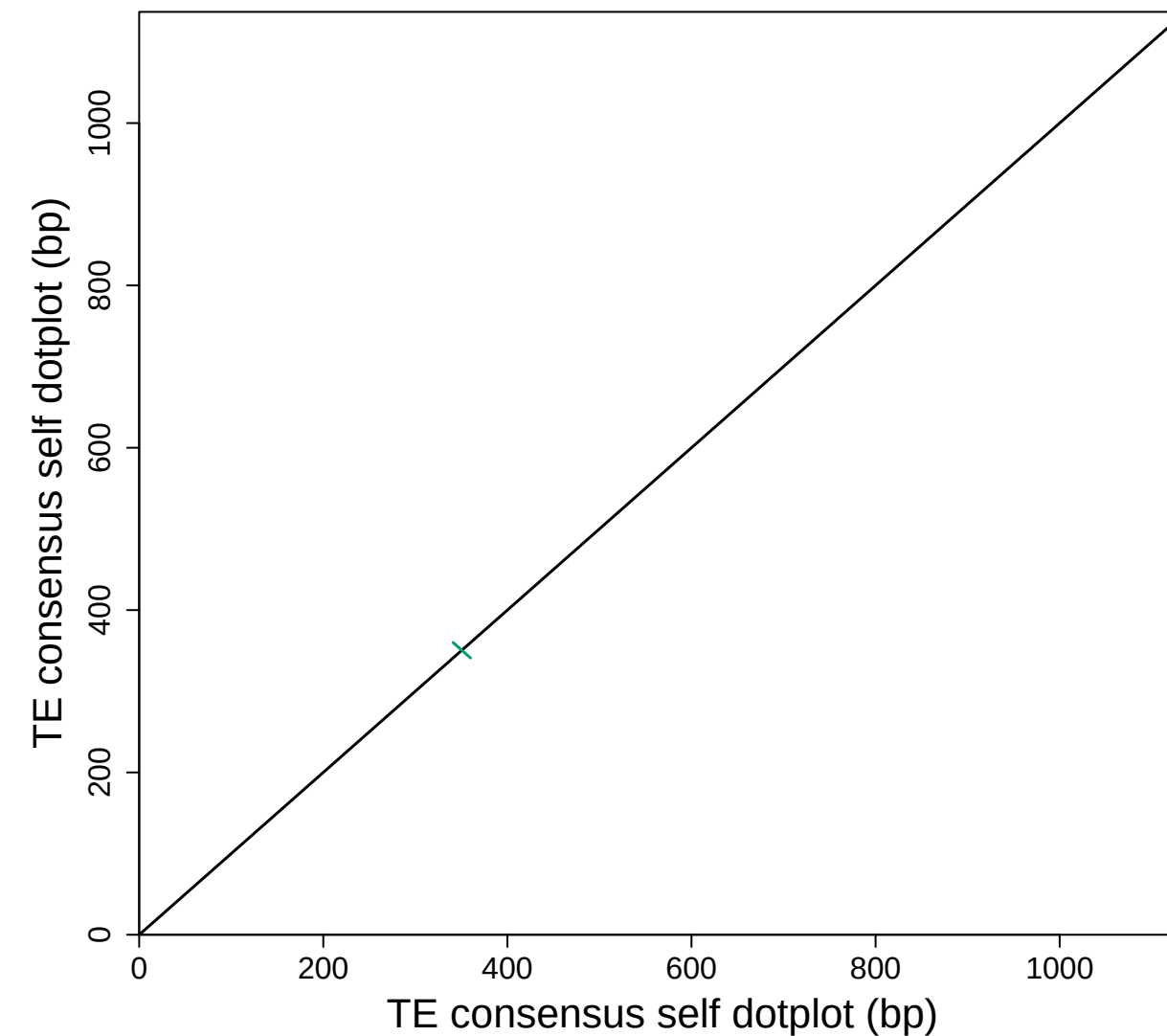
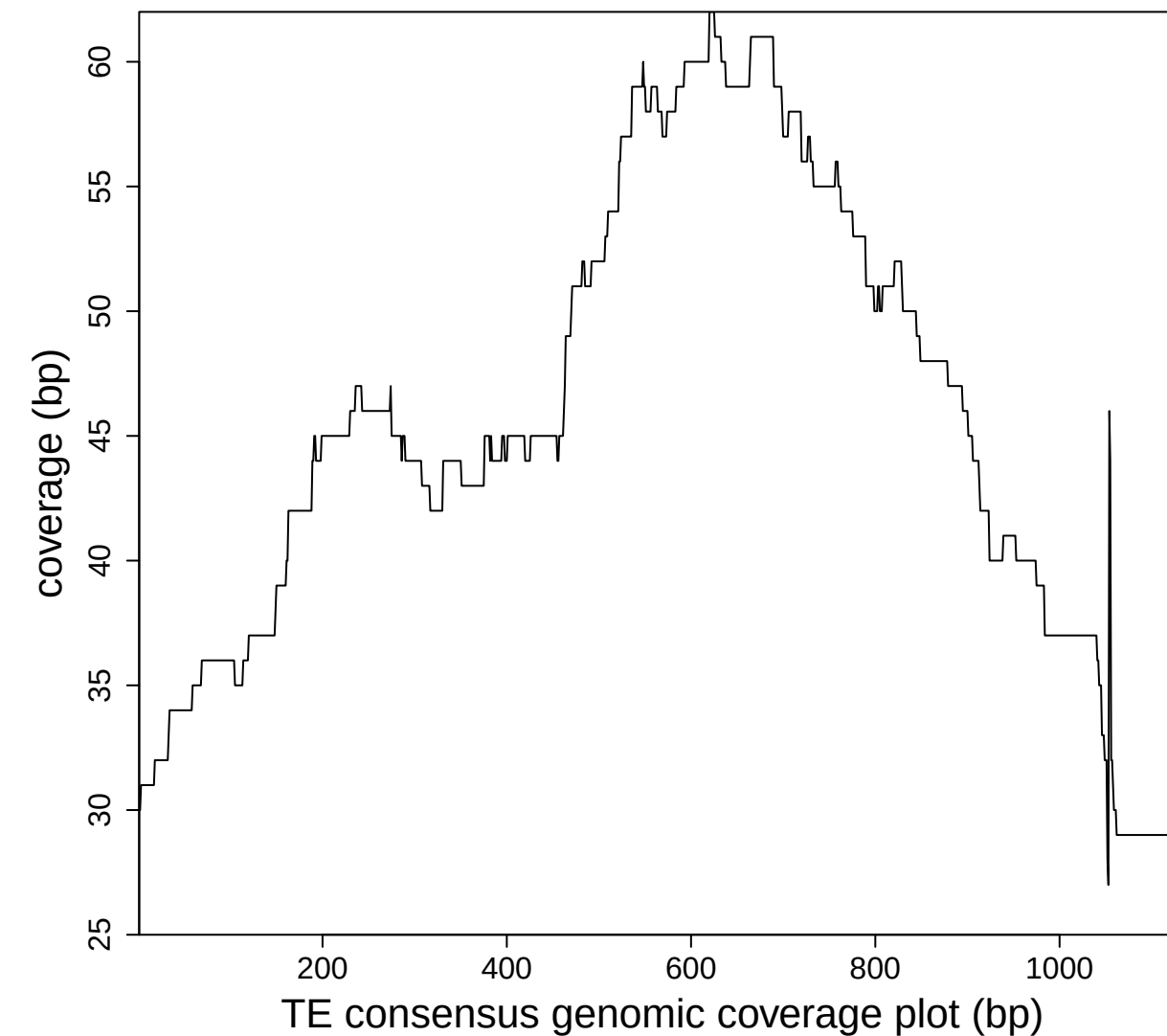
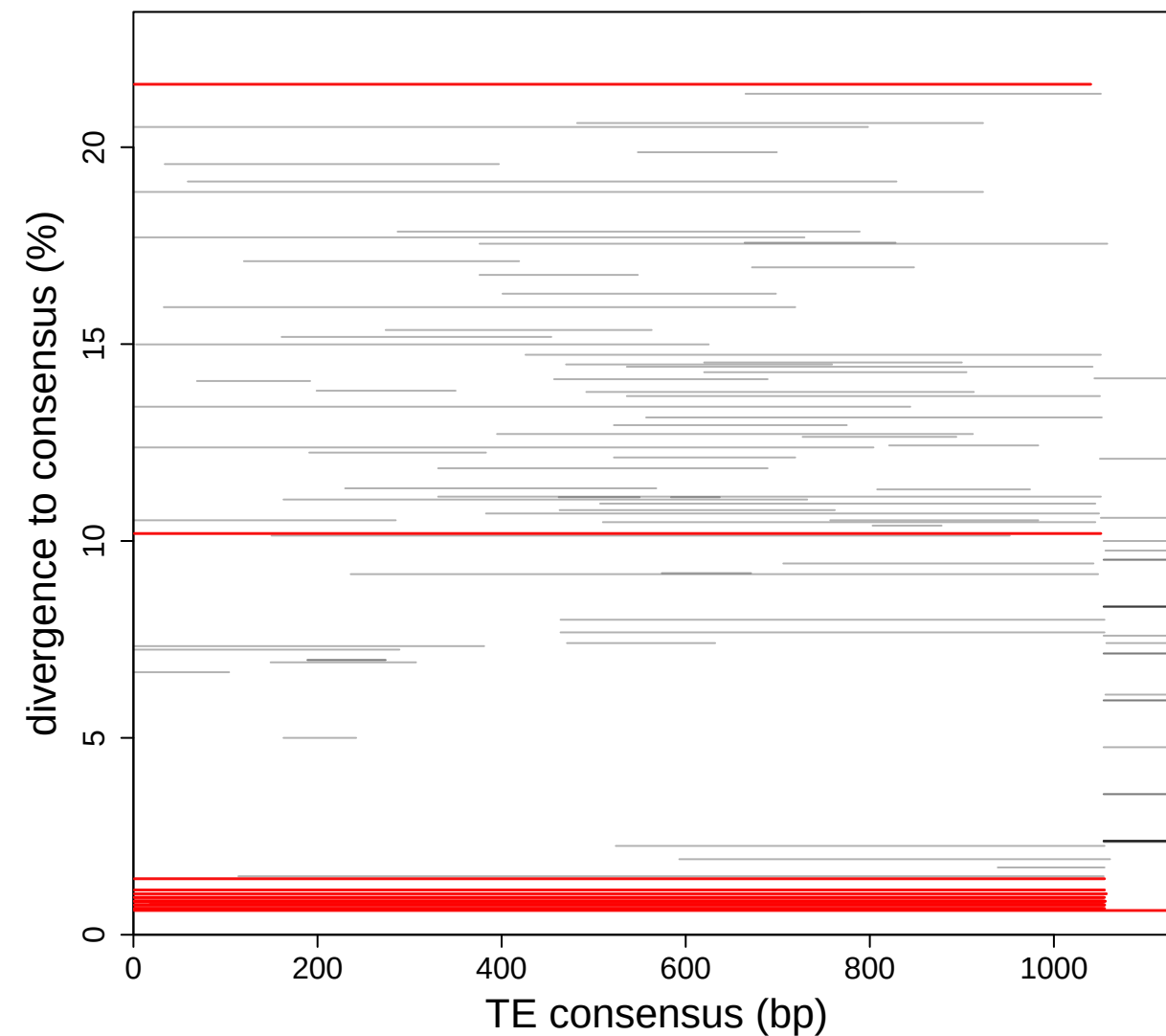


1.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcIn.fa aln.fa cl.f
size: 4606bp; fragments: 121; full length: 8 (>=4145.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

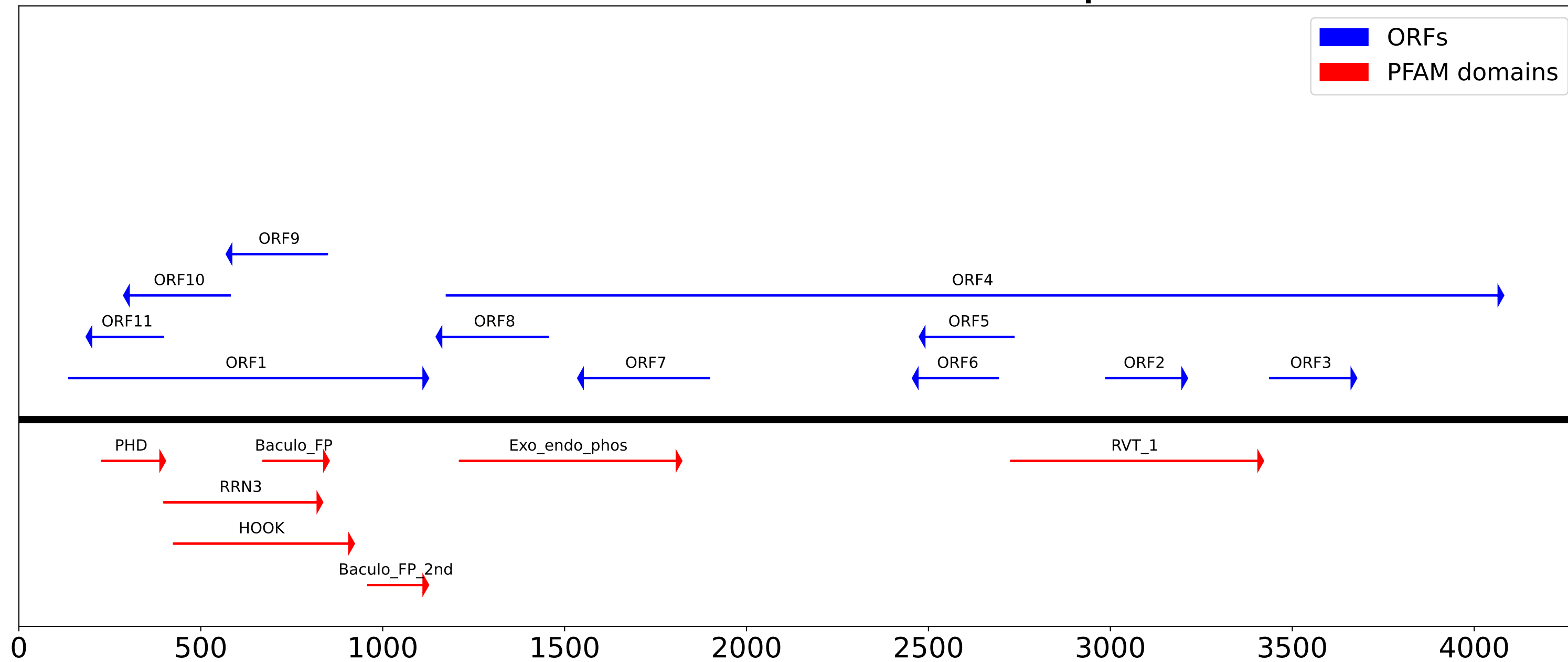


TE: rnd_5_family_1991
size: 1137bp; fragments: 120; full length: 19 (≥ 1023.3 bp)

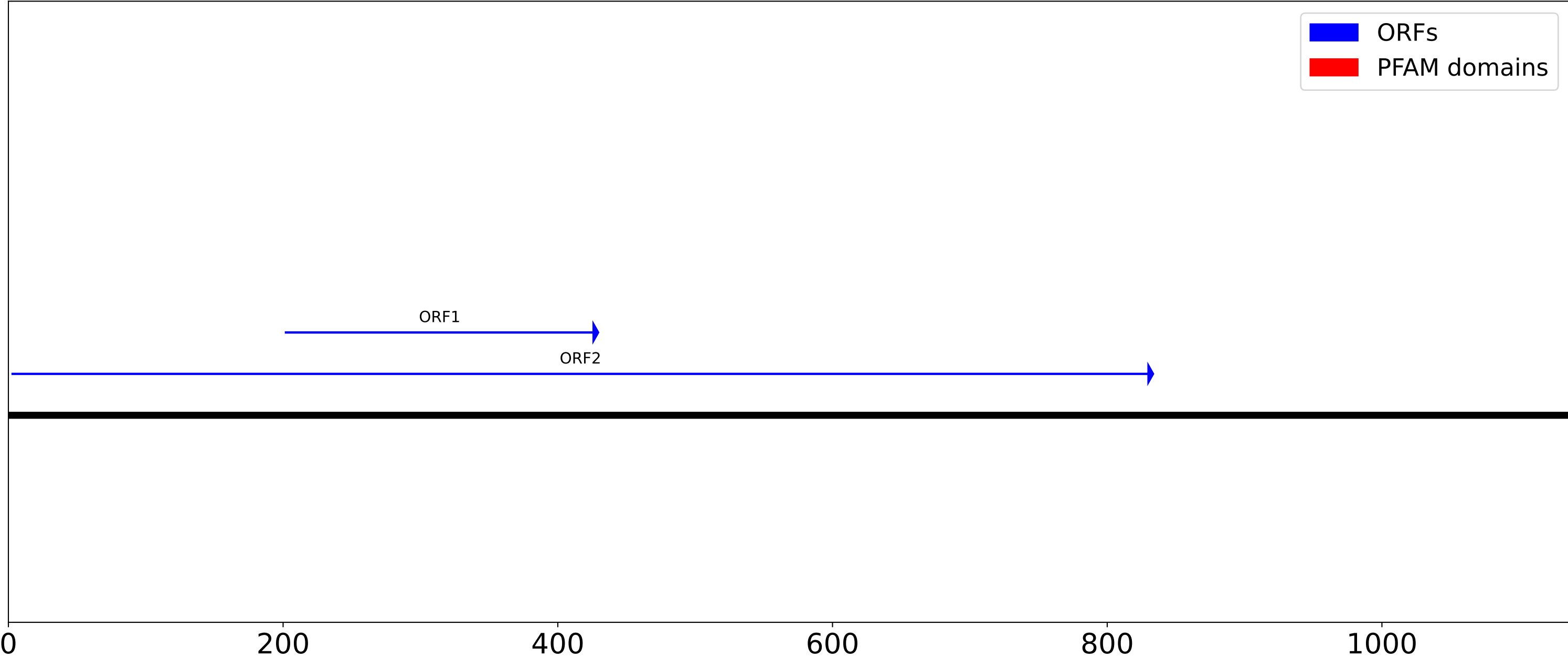


Before TETrimmer 1137 bp

After TEtrimmer ORF and PFAM domain plot



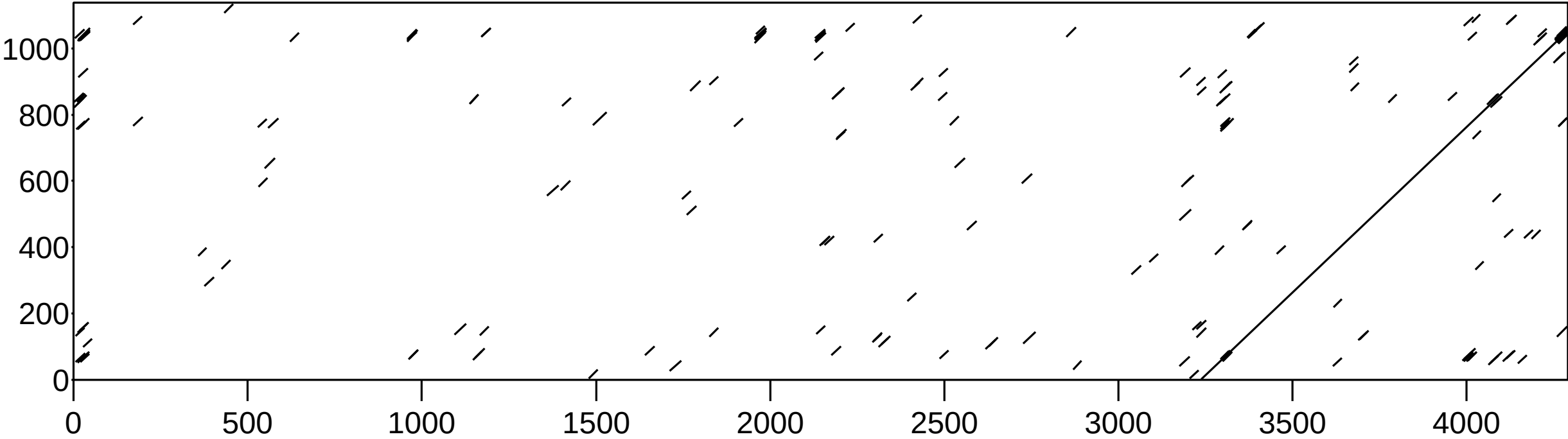
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)