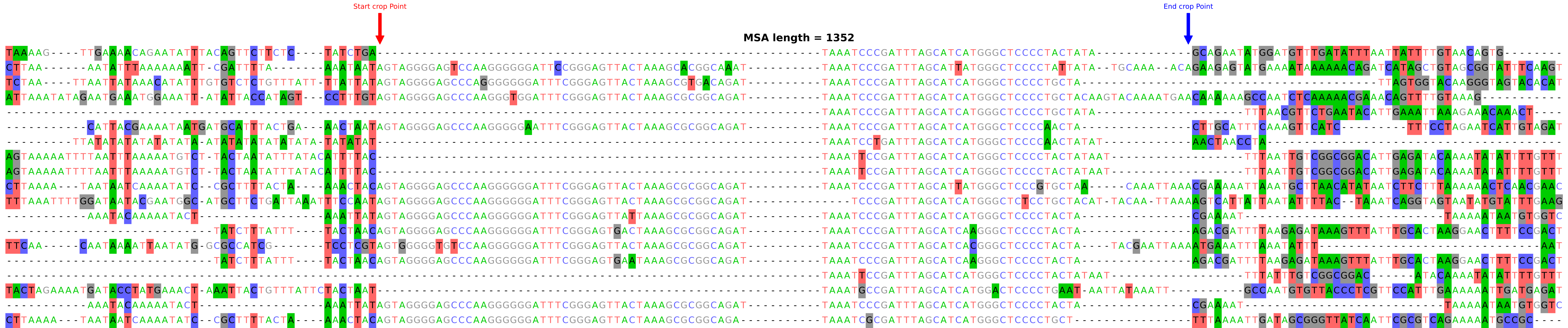
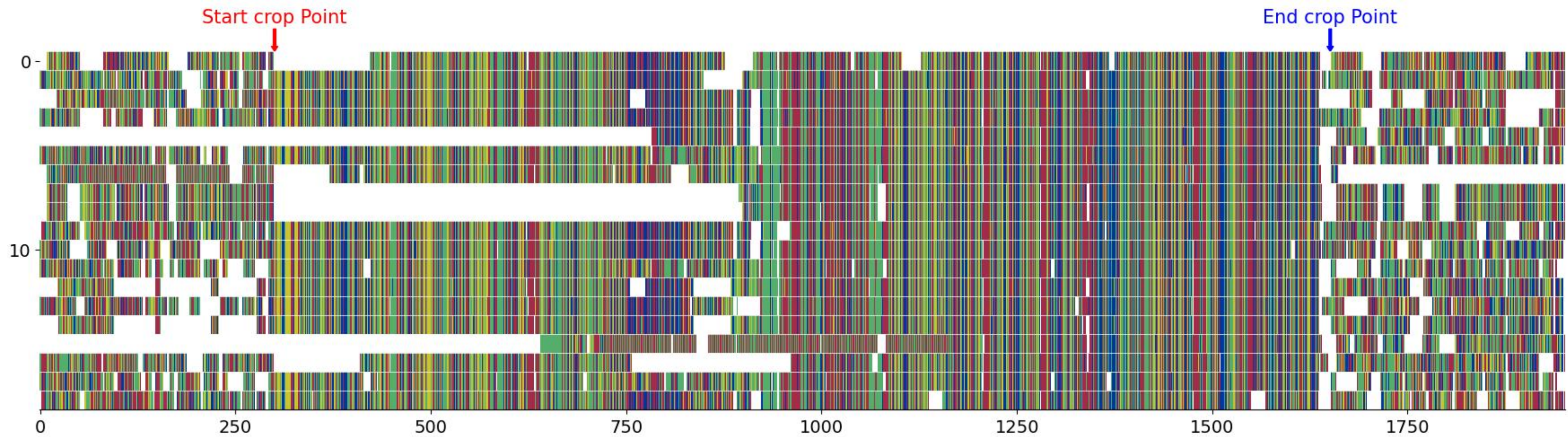




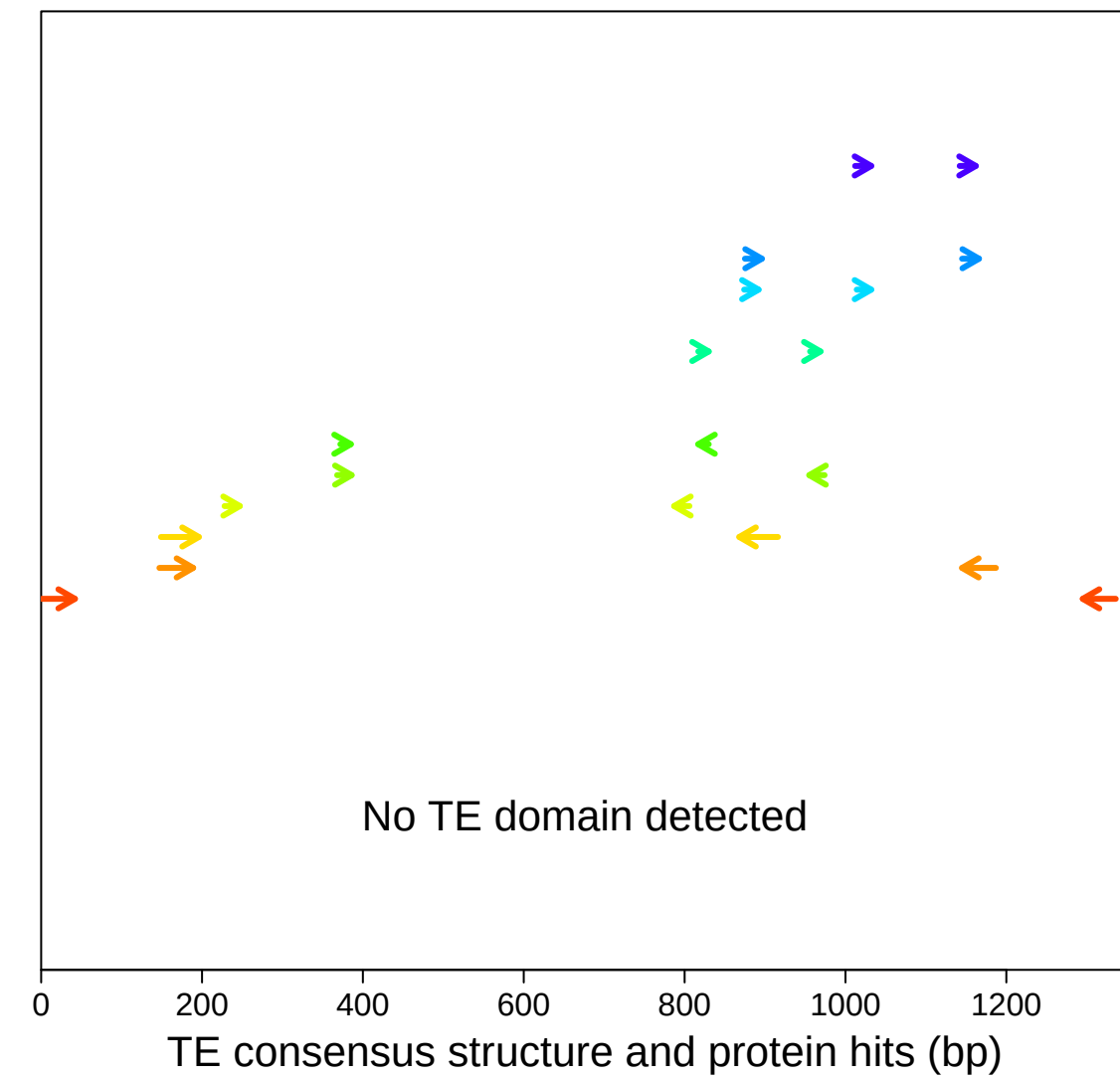
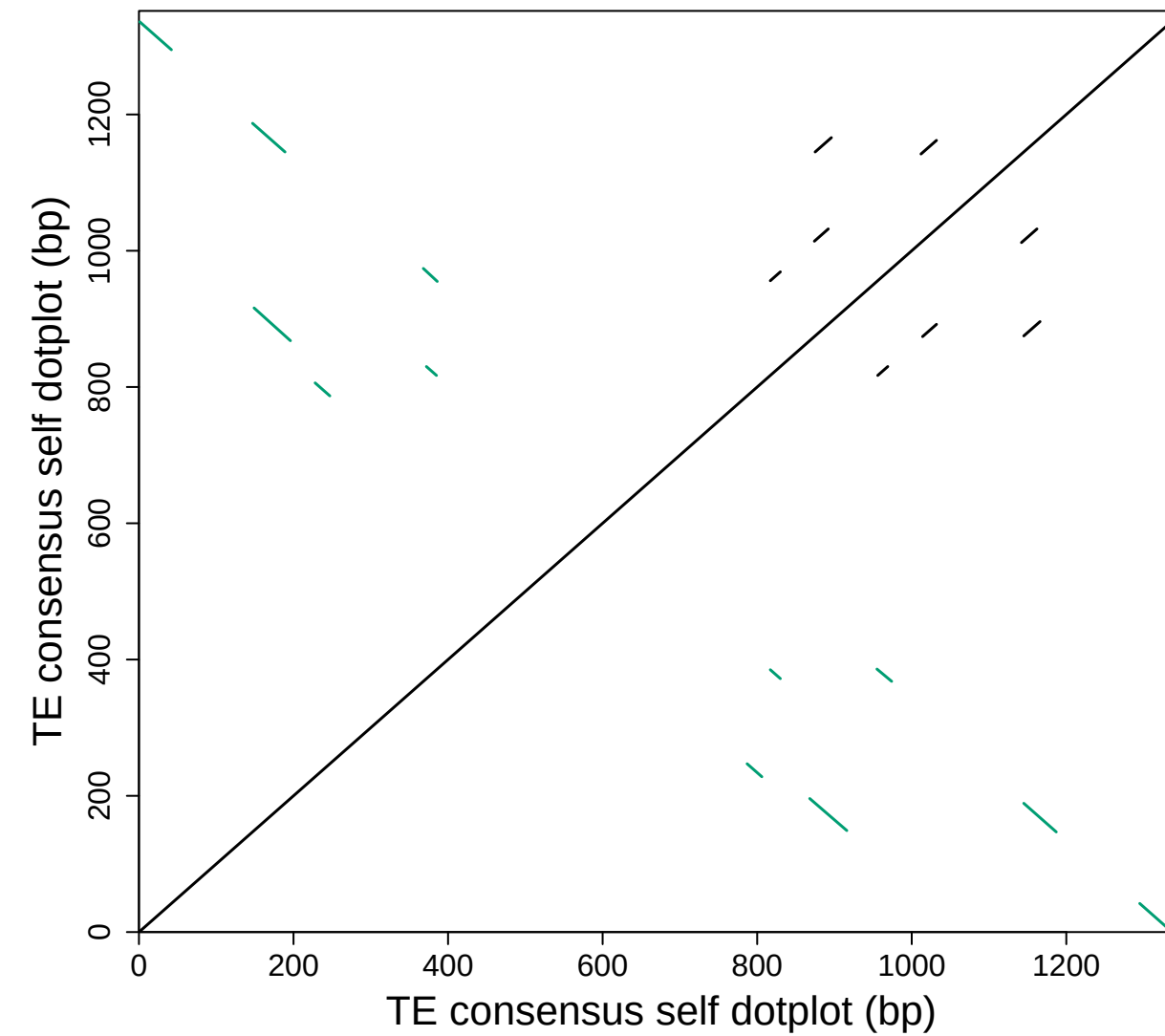
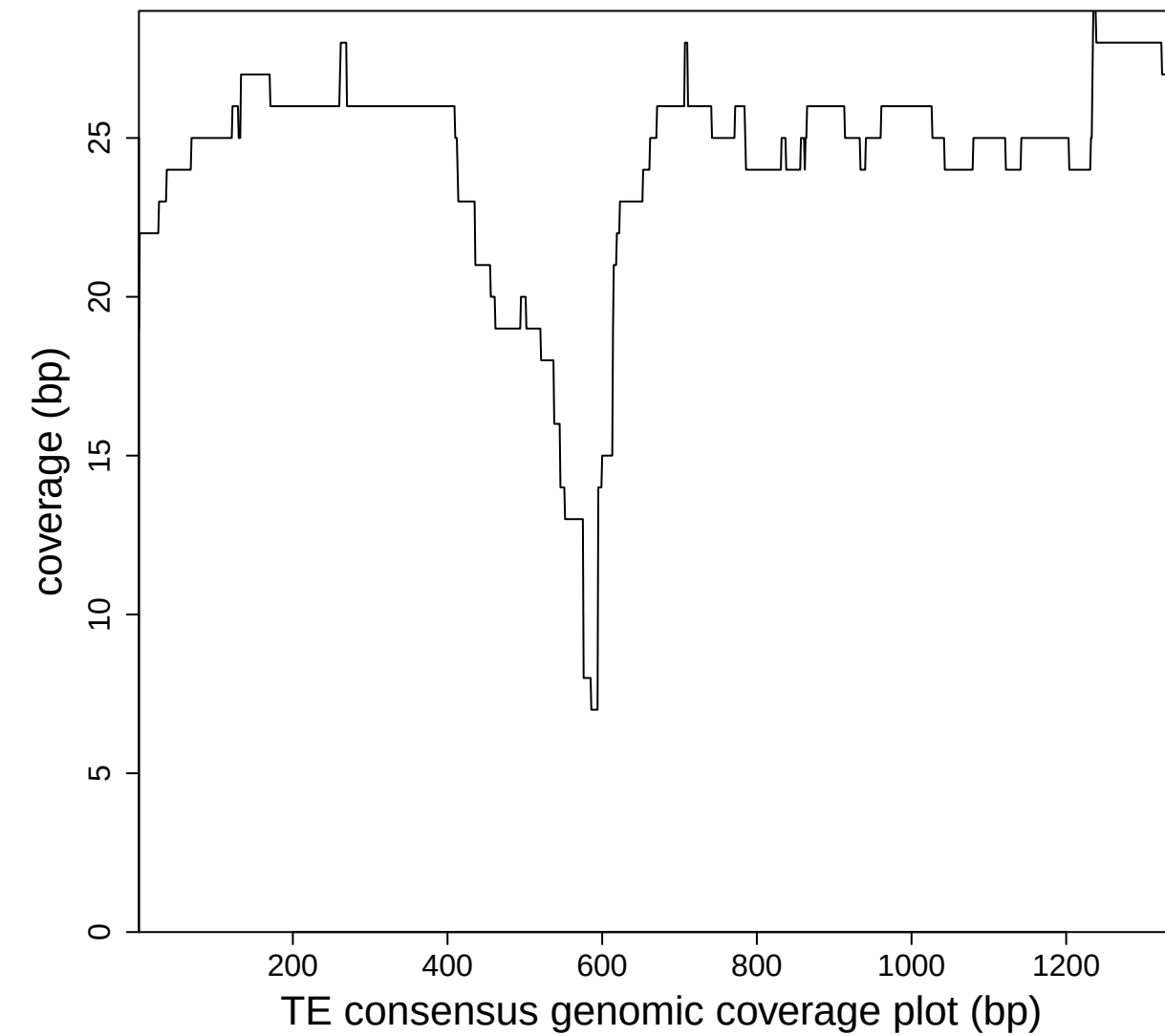
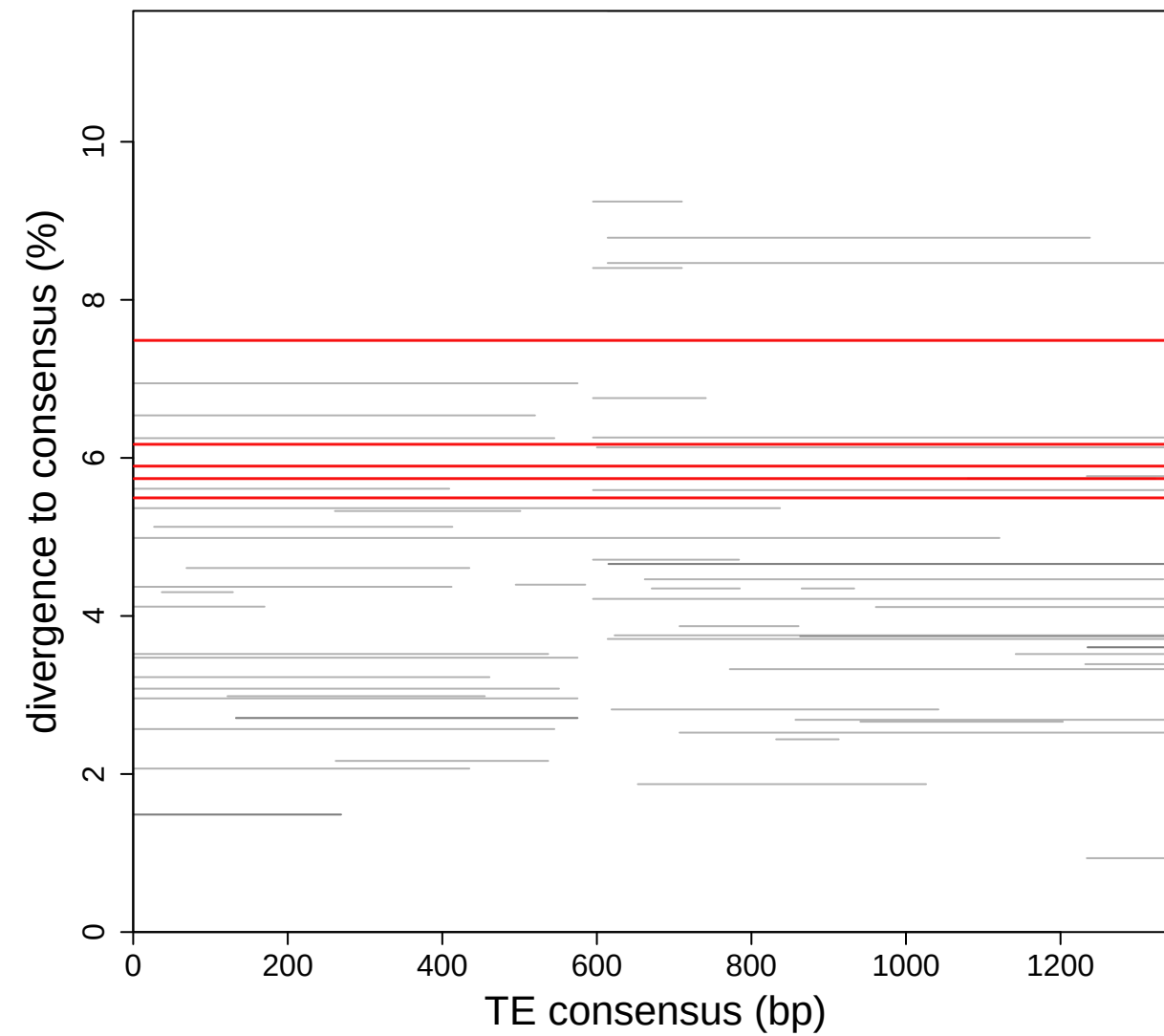
Start crop Point

End crop Point



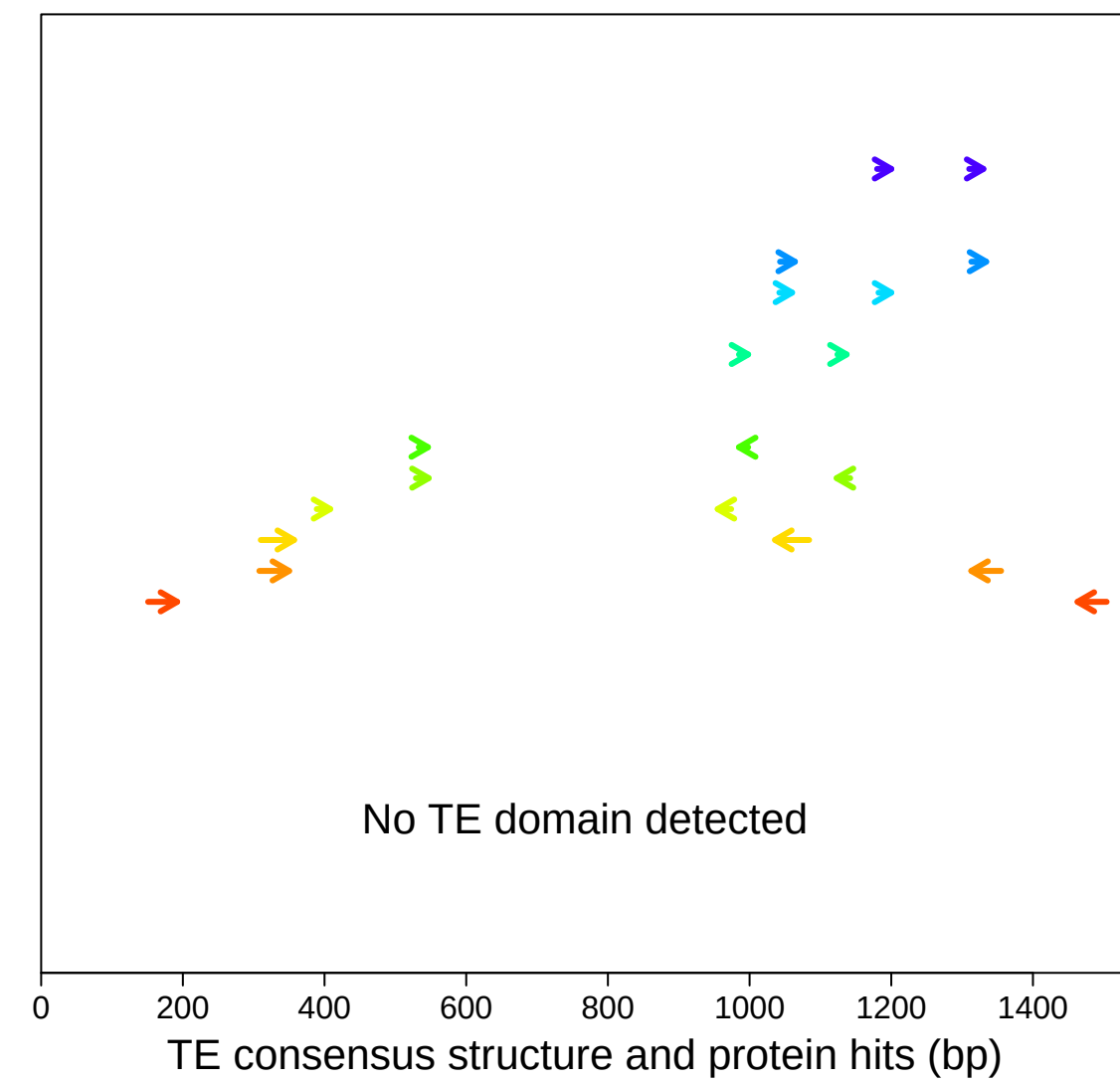
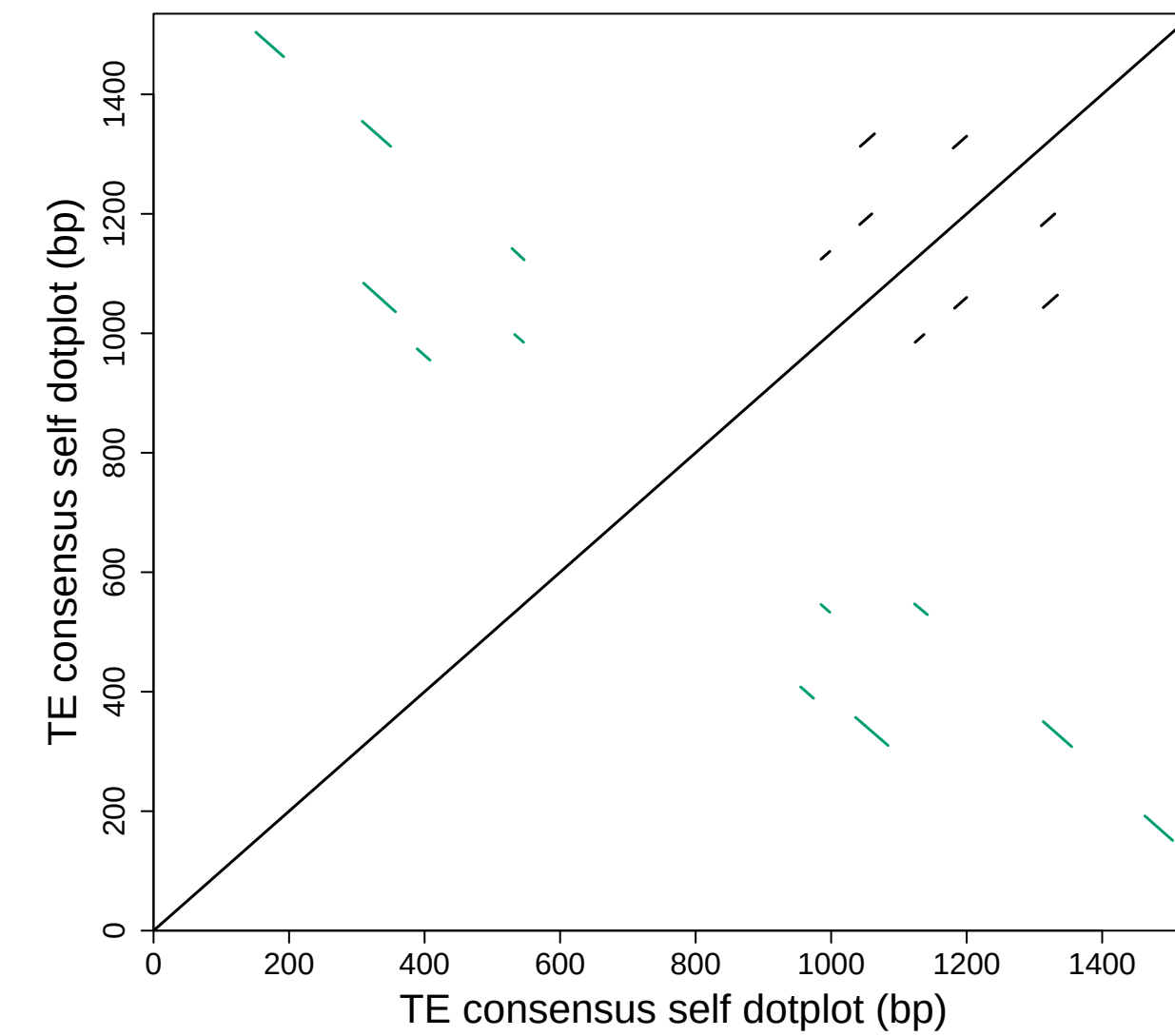
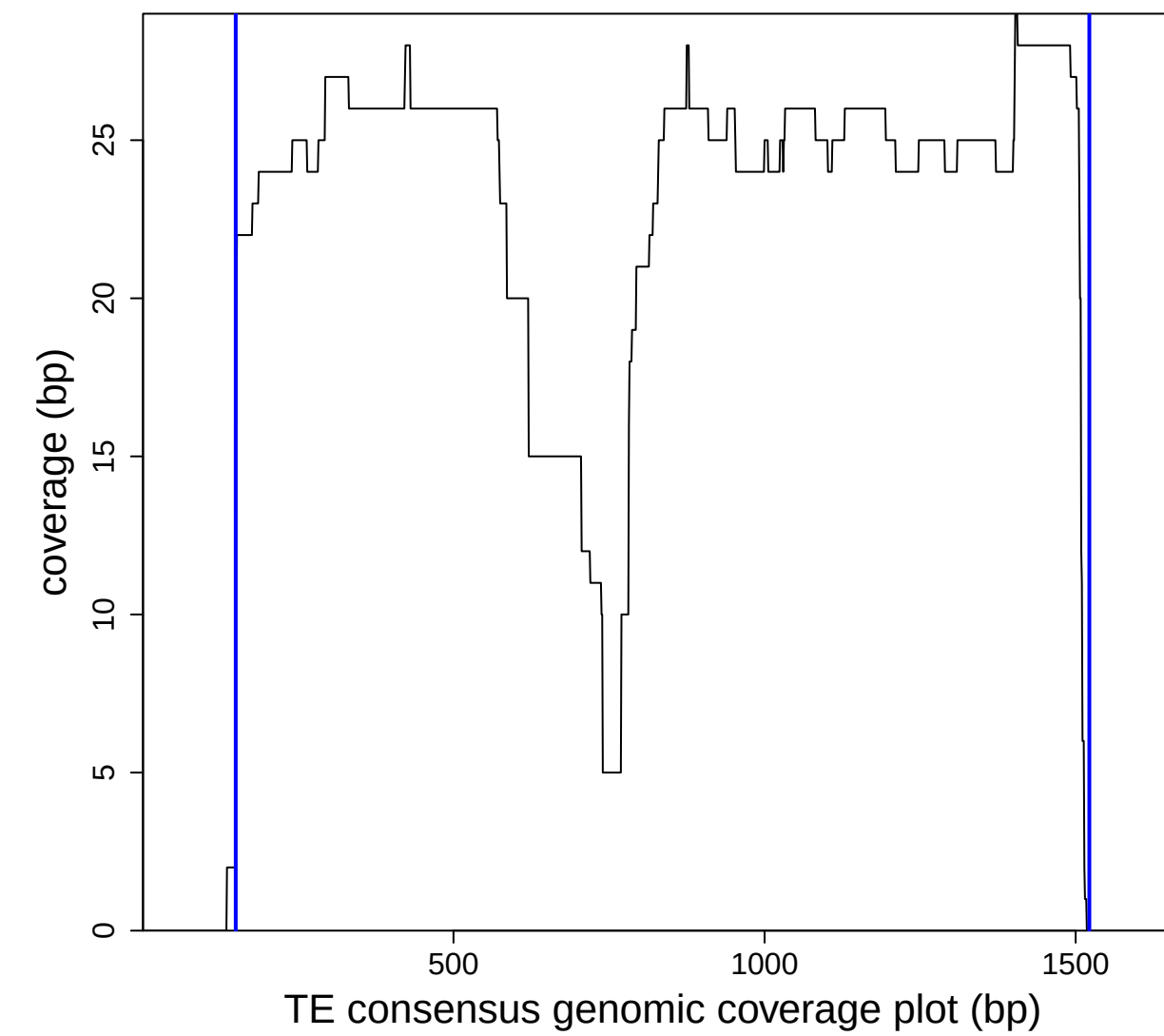
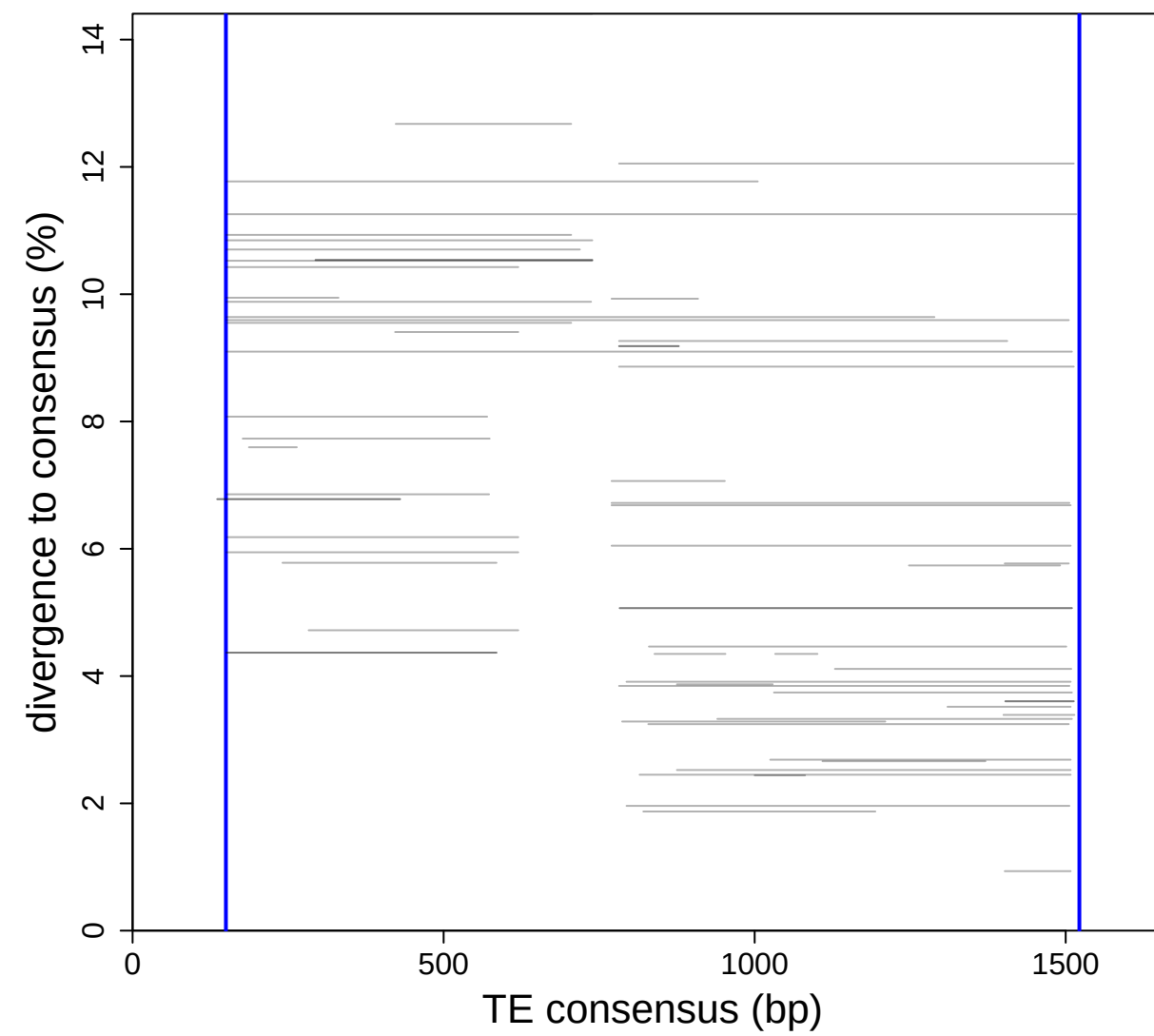
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1352bp; fragments: 66; full length: 5 (≥ 1216.8 bp)

After TEtrimmer 1352 bp

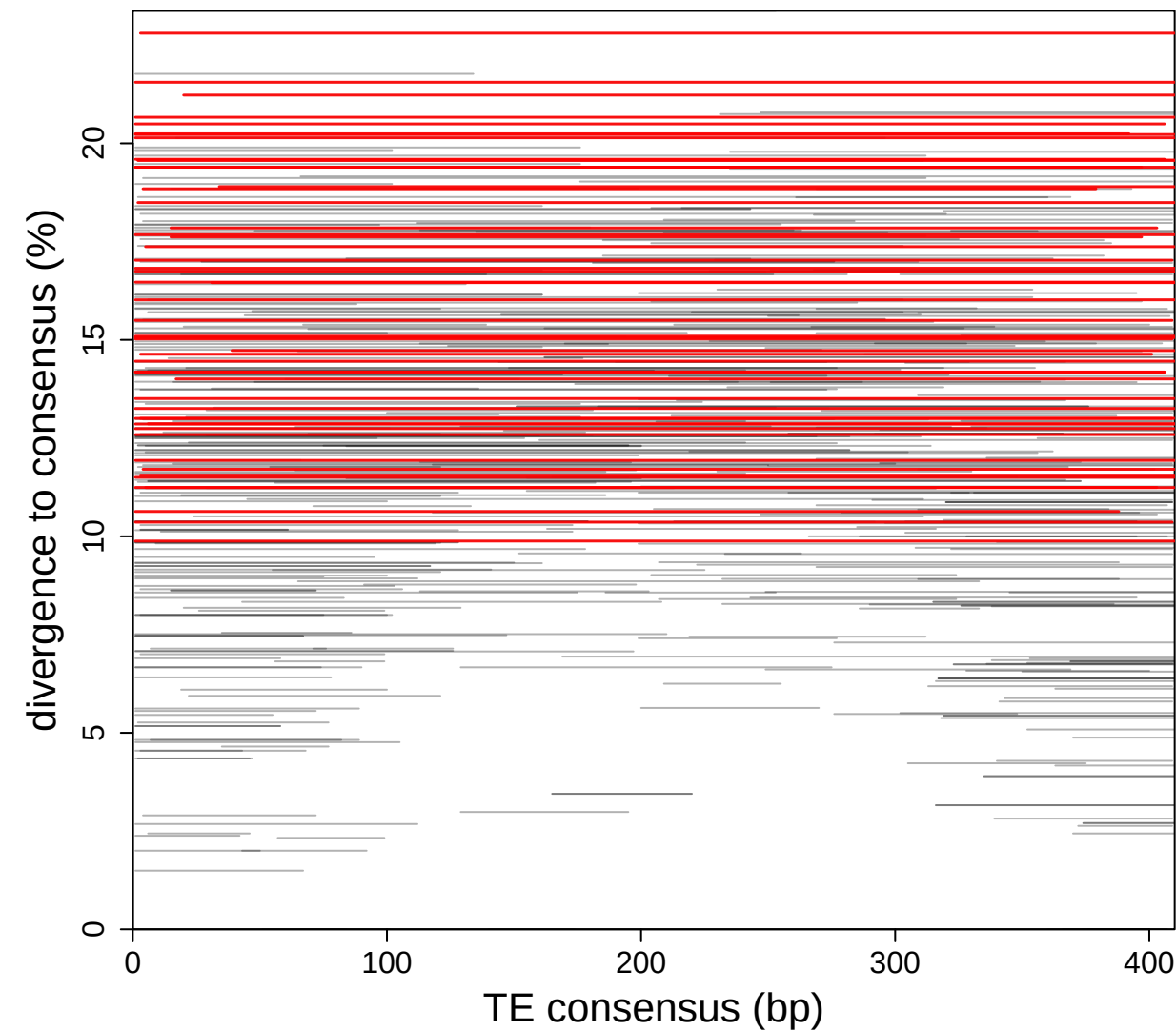


8.fasta.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 1672bp; fragments: 67; full length: 0 (>=1504.8bp)

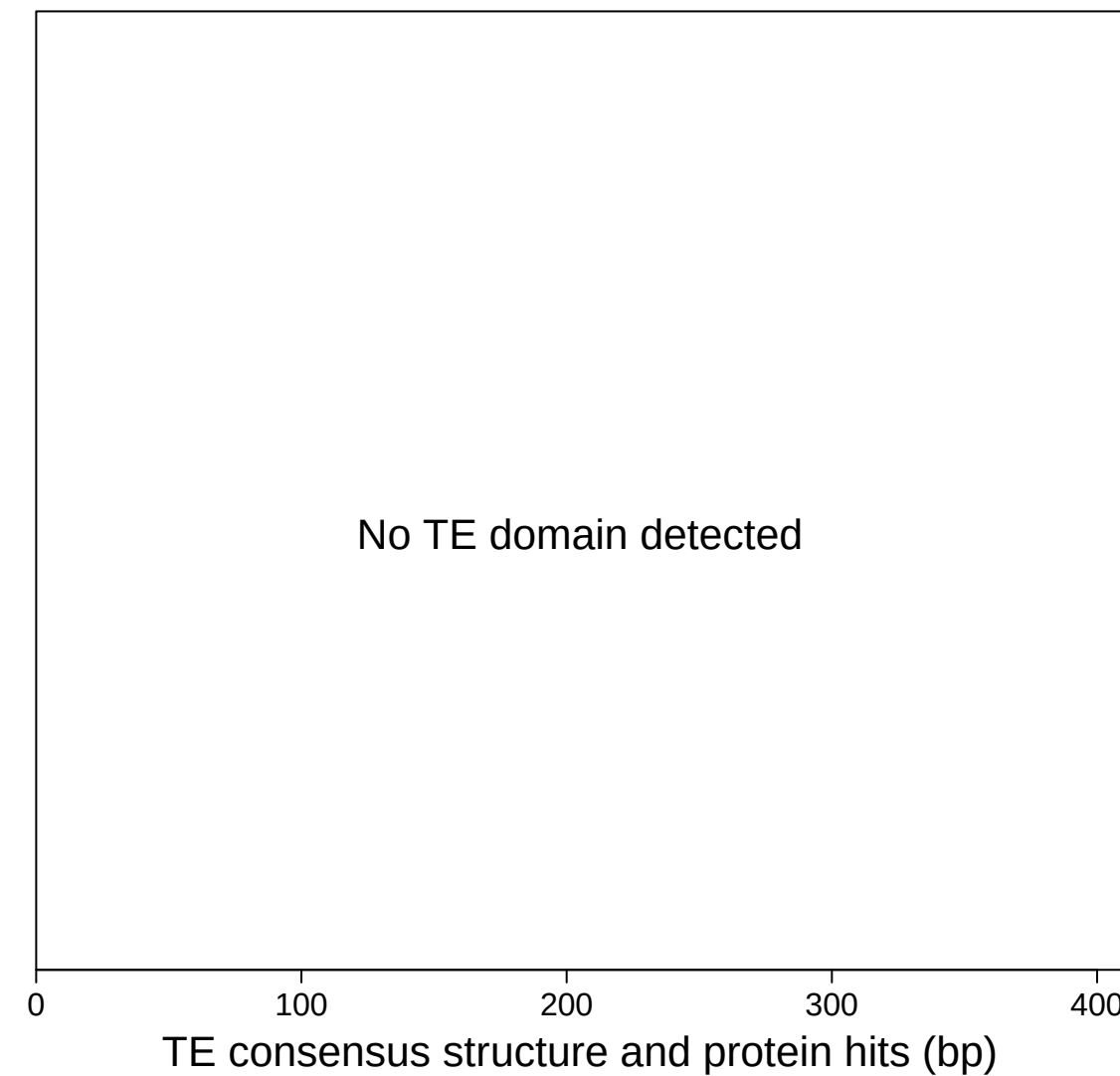
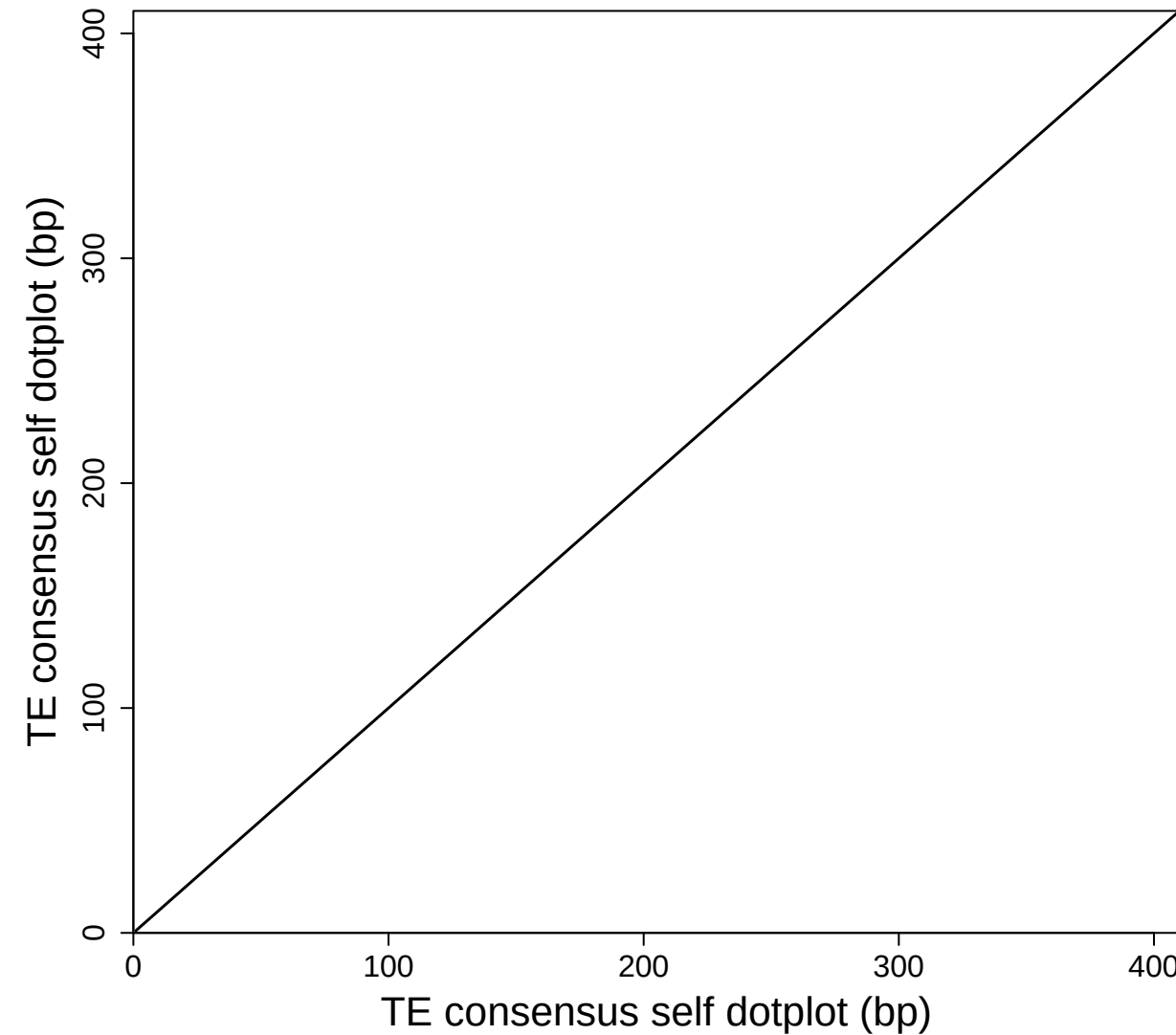
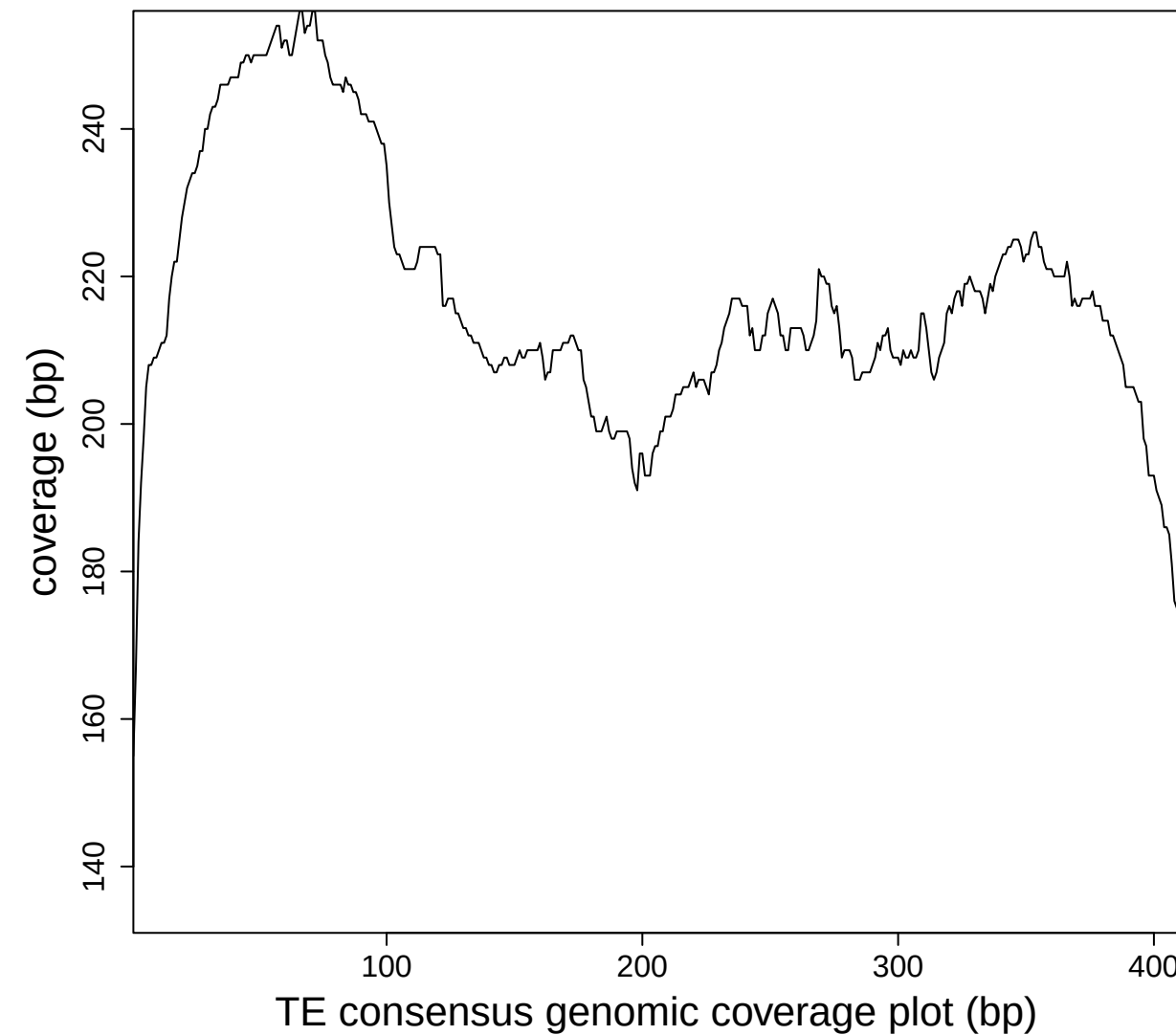
After TETrimmer Extended plot Blue lines are boundaries



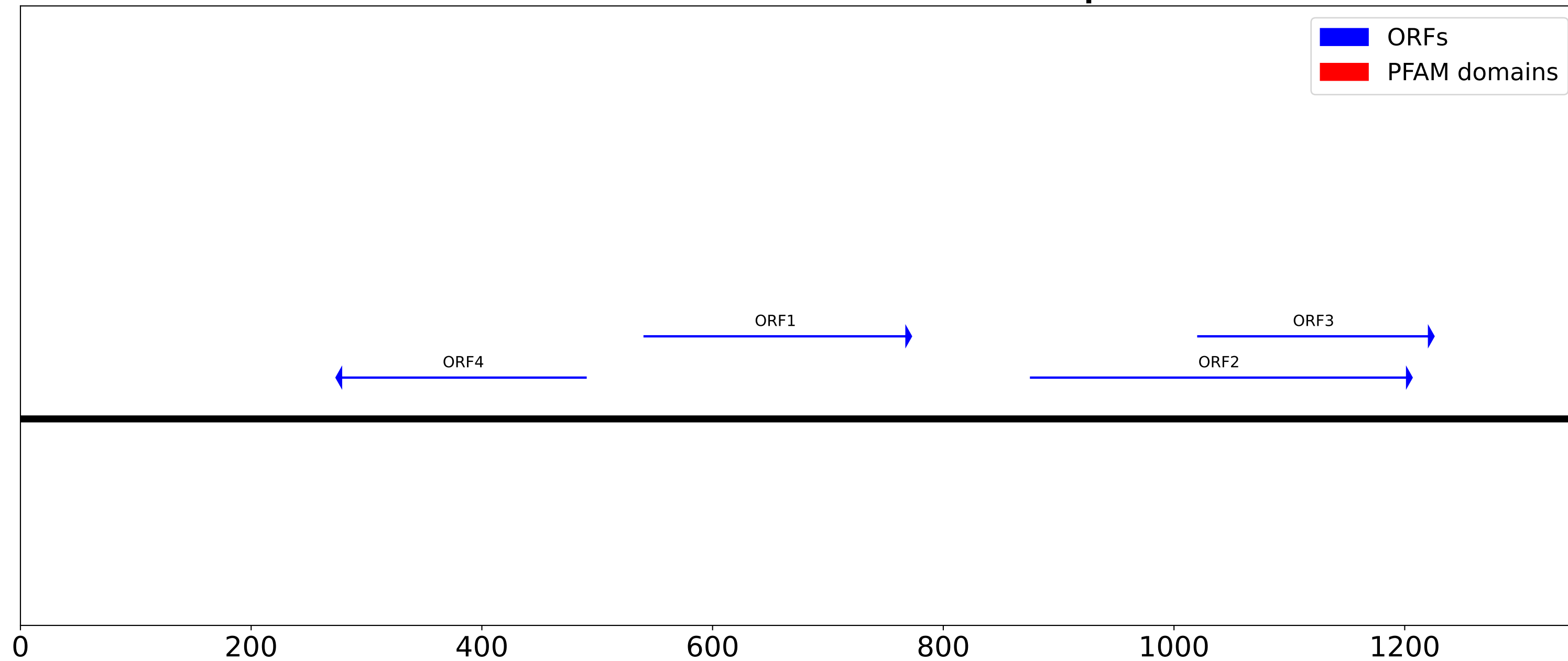
TE: rnd_5_family_7658
size: 410bp; fragments: 478; full length: 53 (≥ 369 bp)



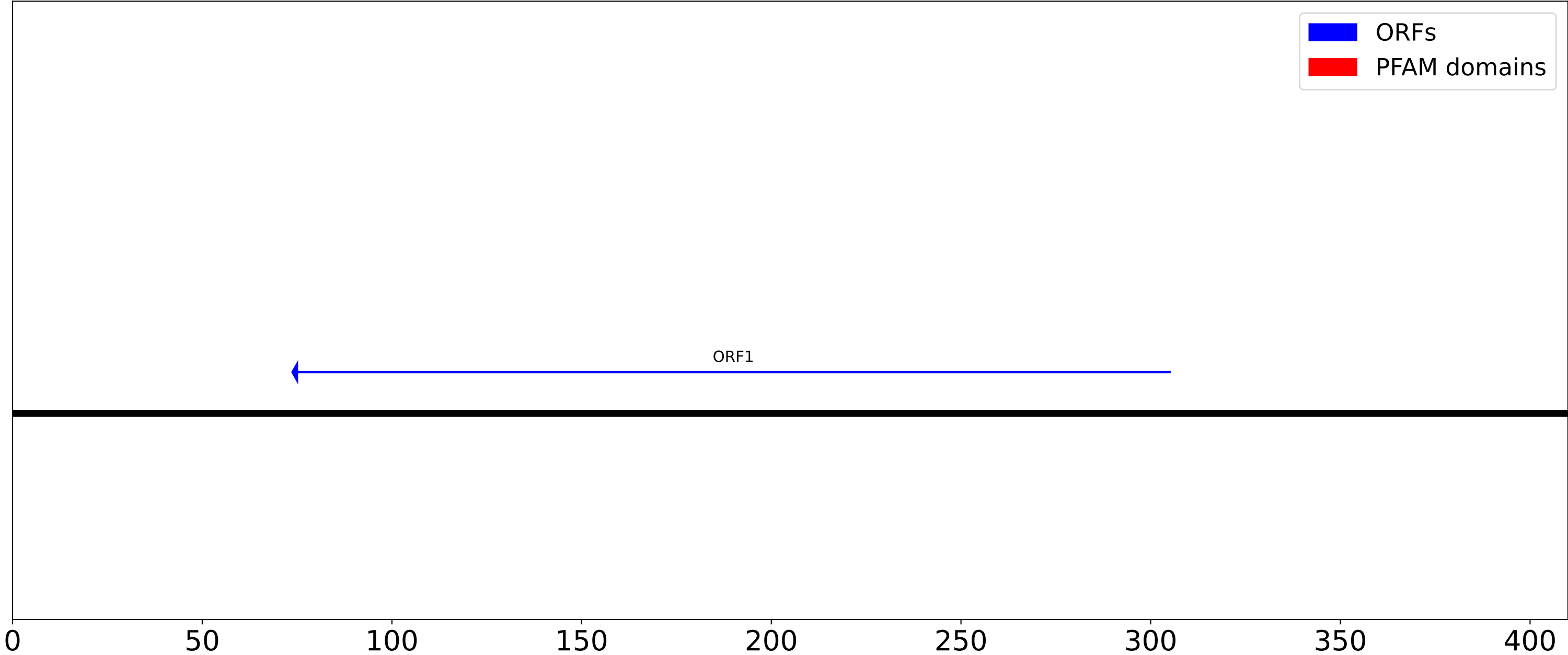
Before TEtrimmer 410 bp



After TEtrimmer ORF and PFAM domain plot



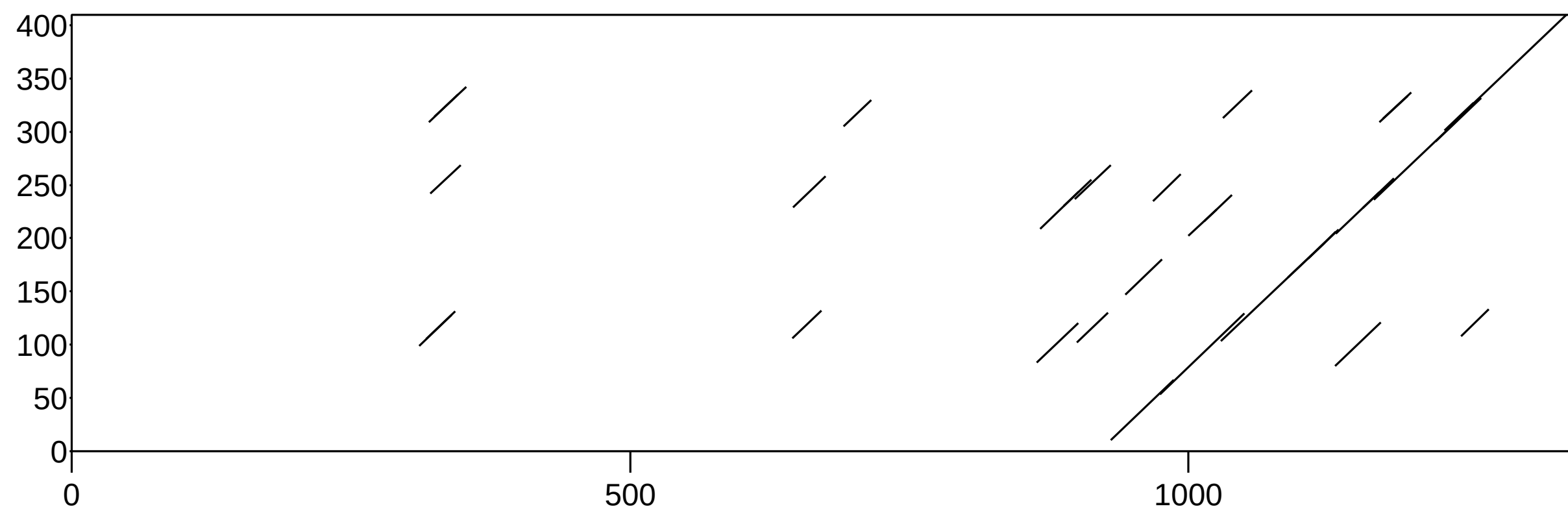
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

: consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)