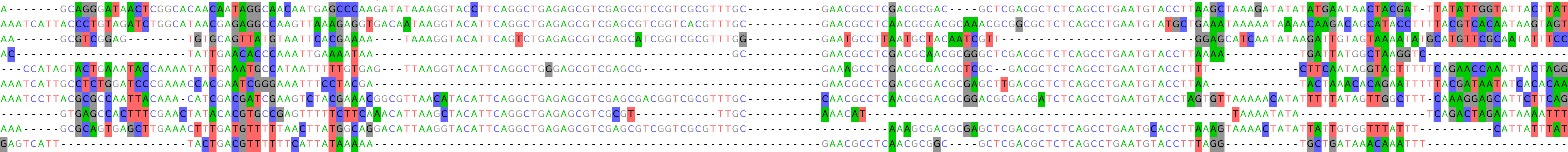


Start Crop Point

End Crop Point

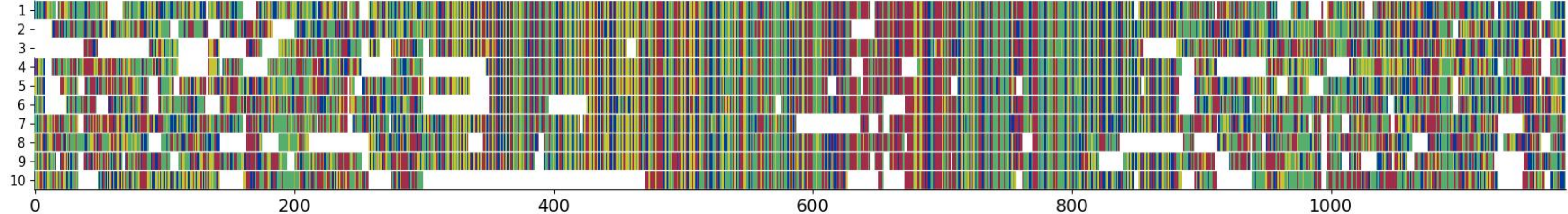
MSA length = 581



Start crop Point

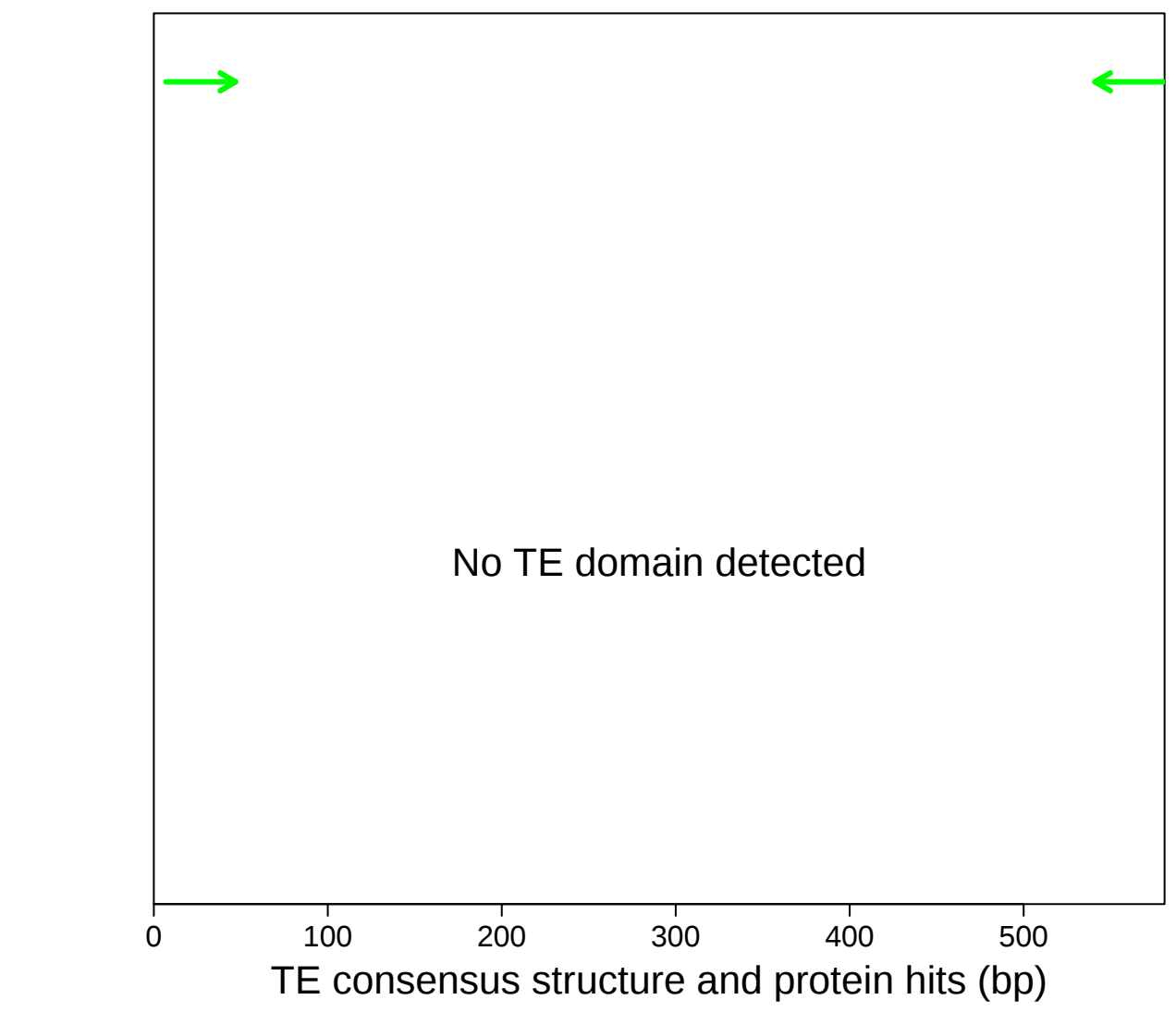
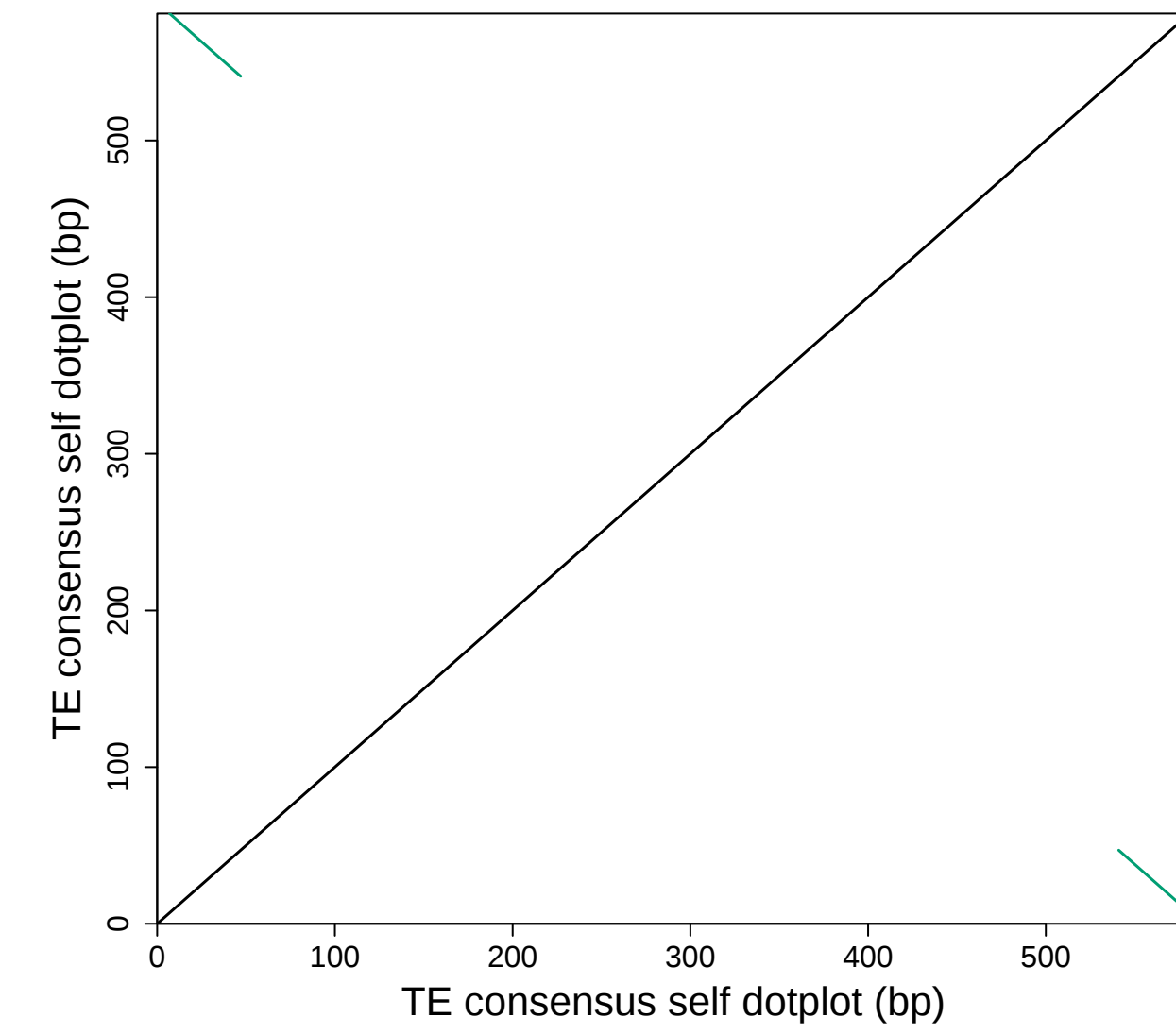
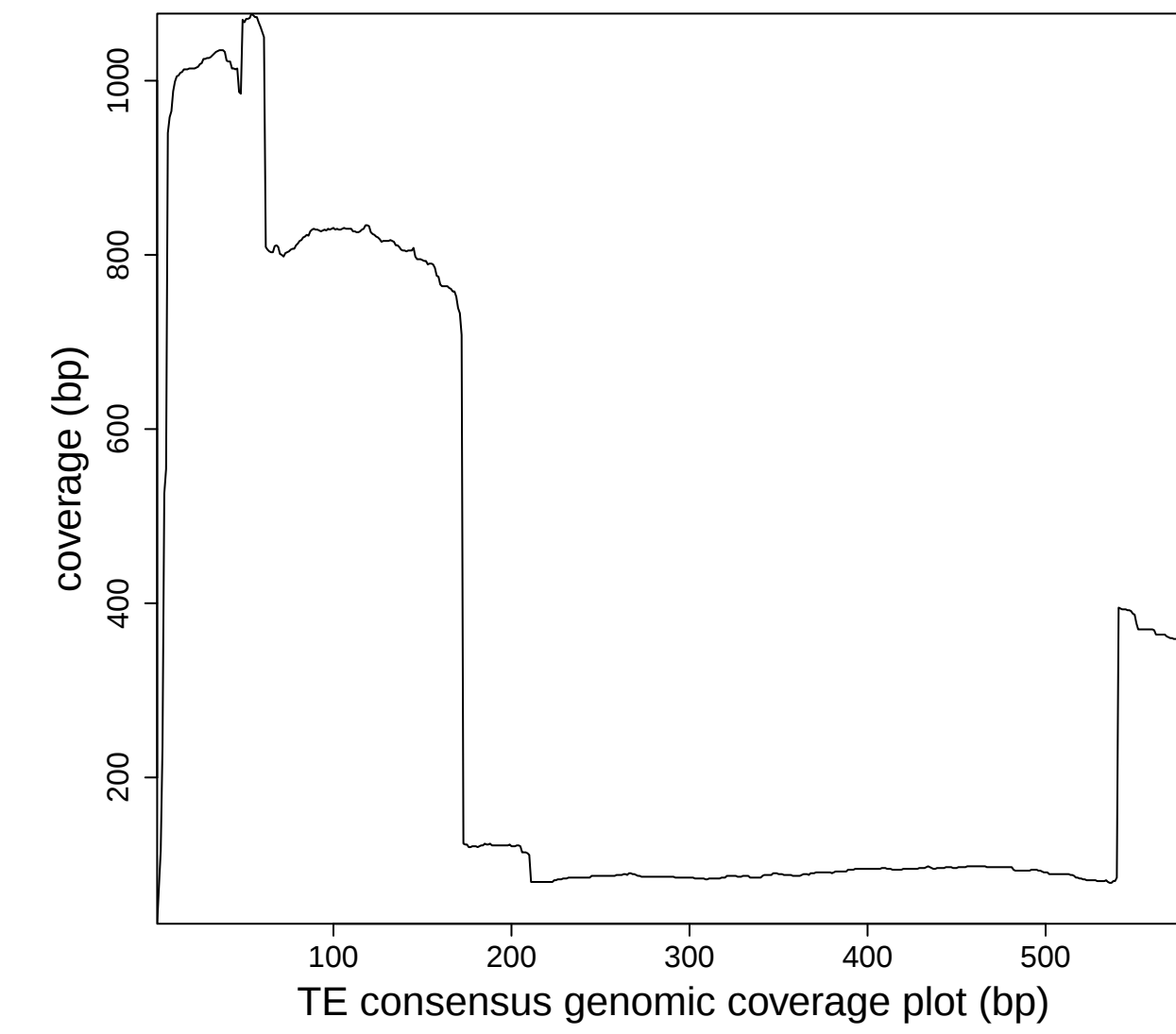
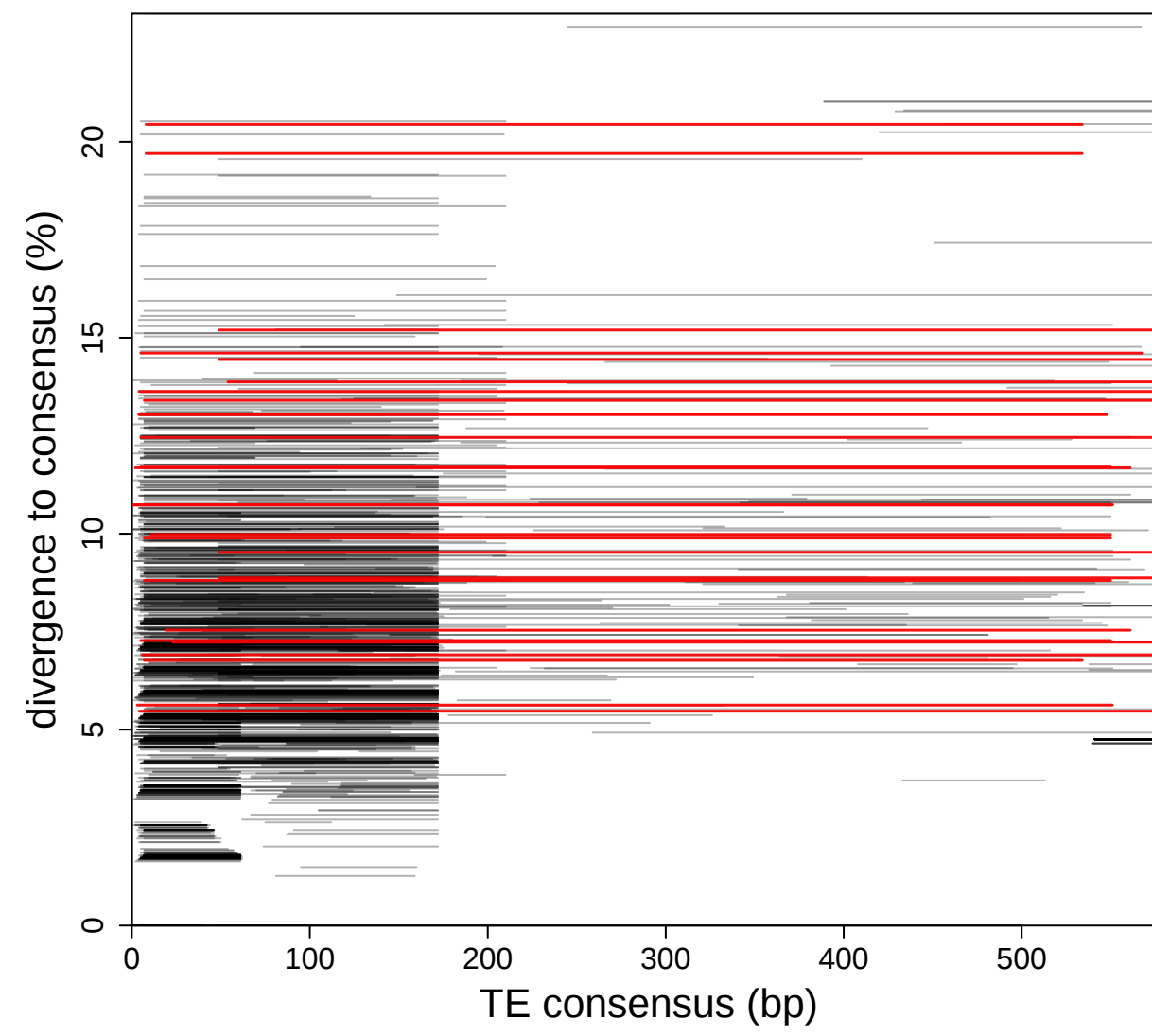


End crop Point



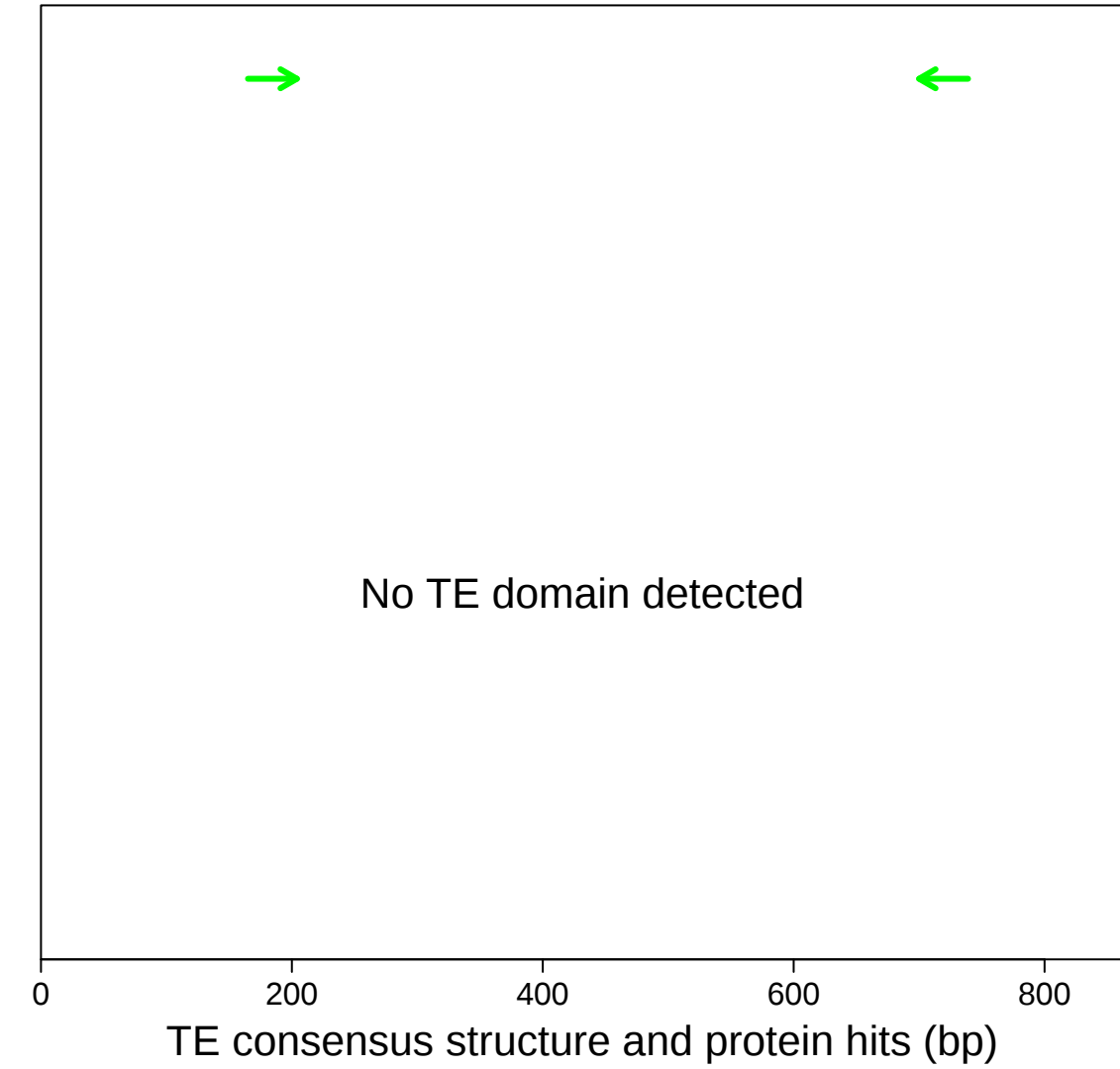
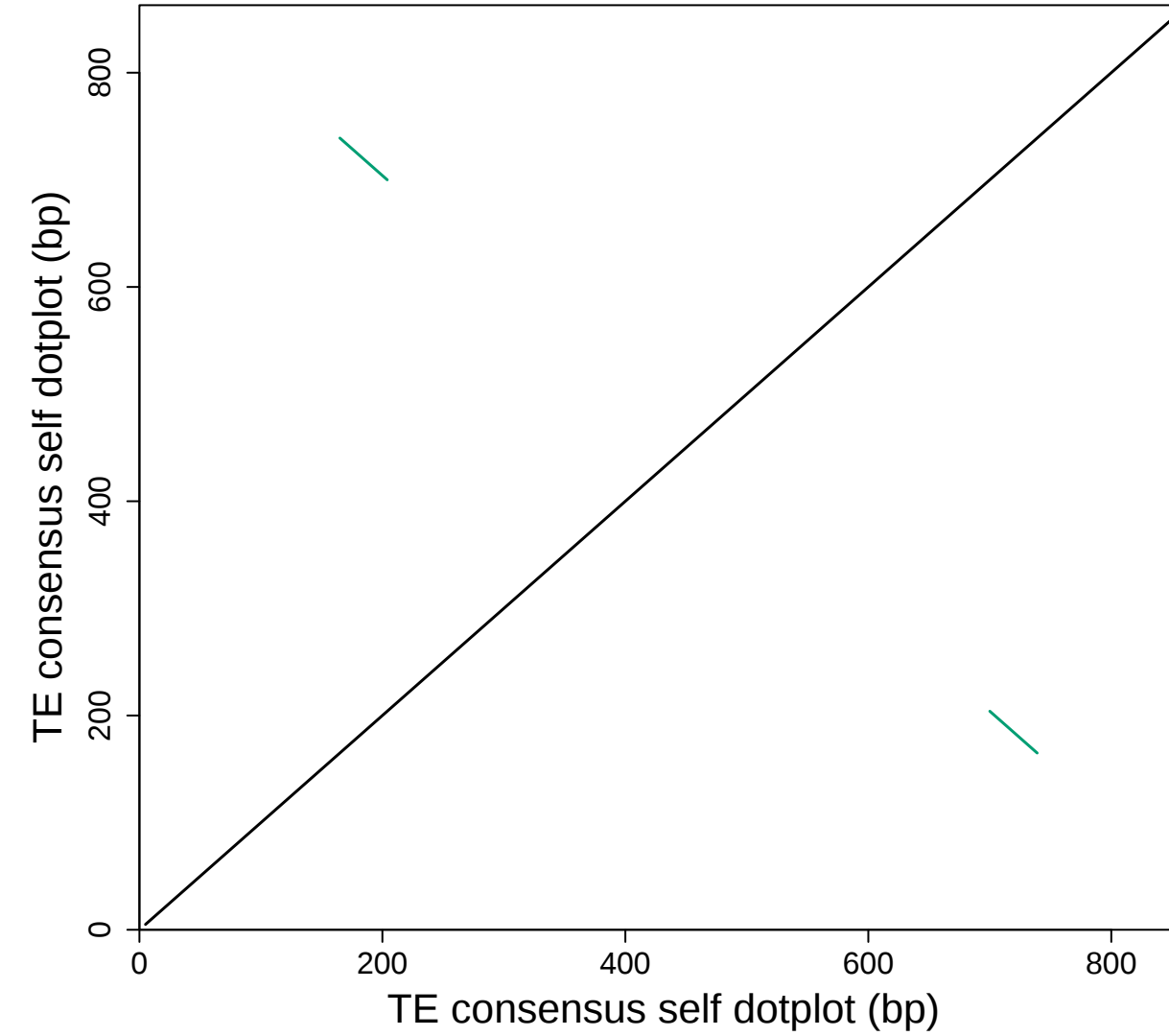
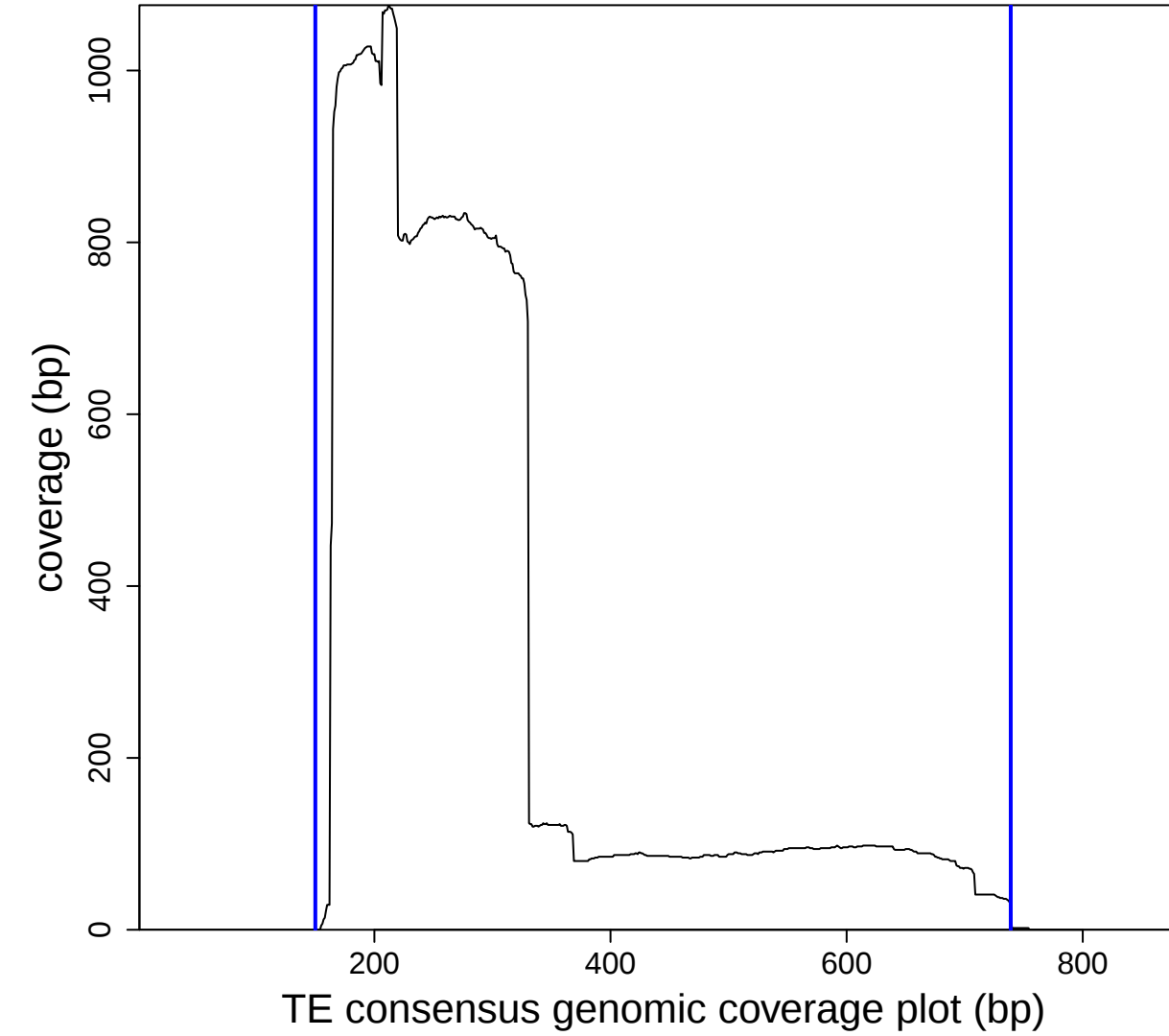
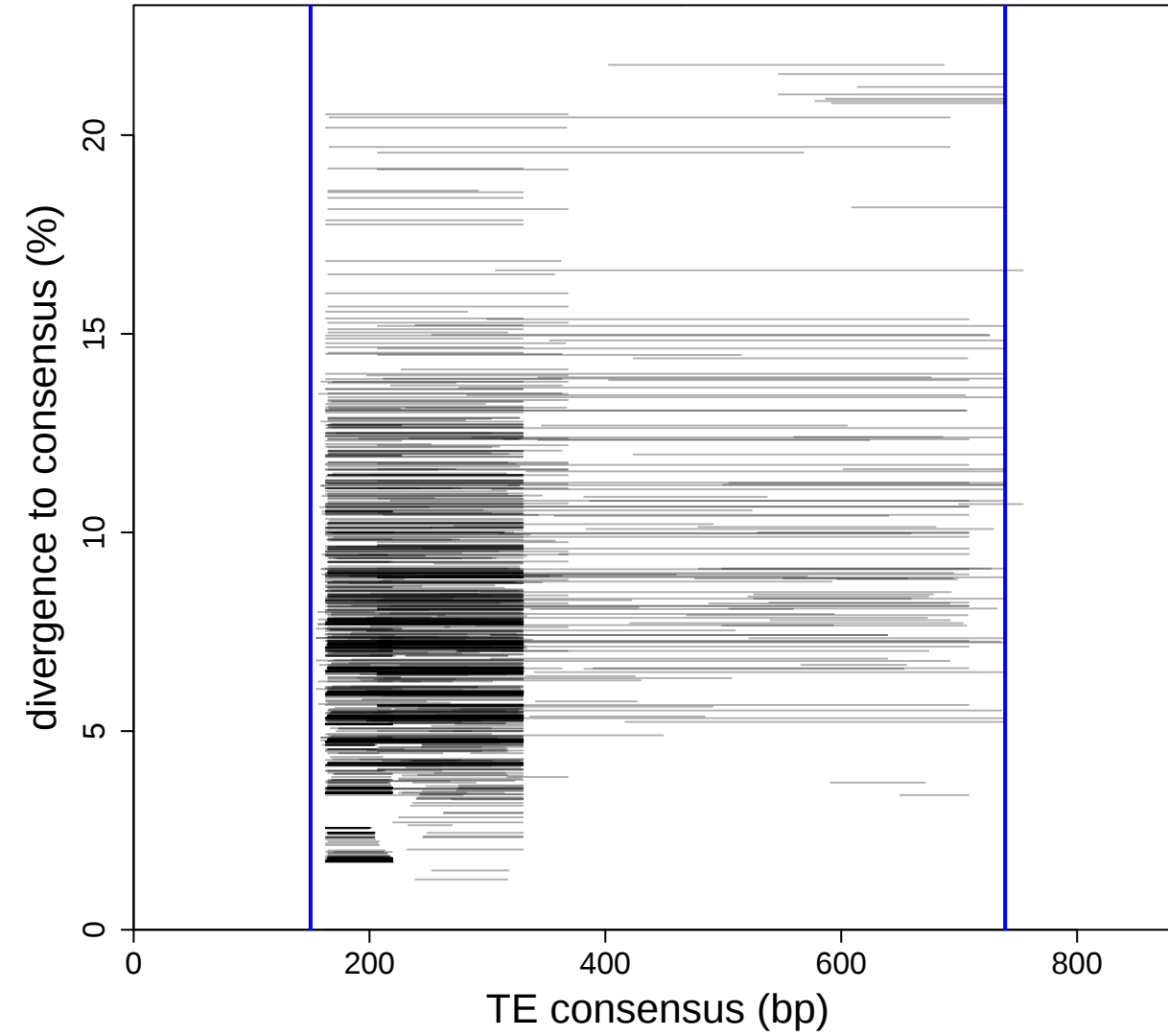
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 581bp; fragments: 1641; full length: 27 (>=522.9bp)

After TEtrimmer 581 bp

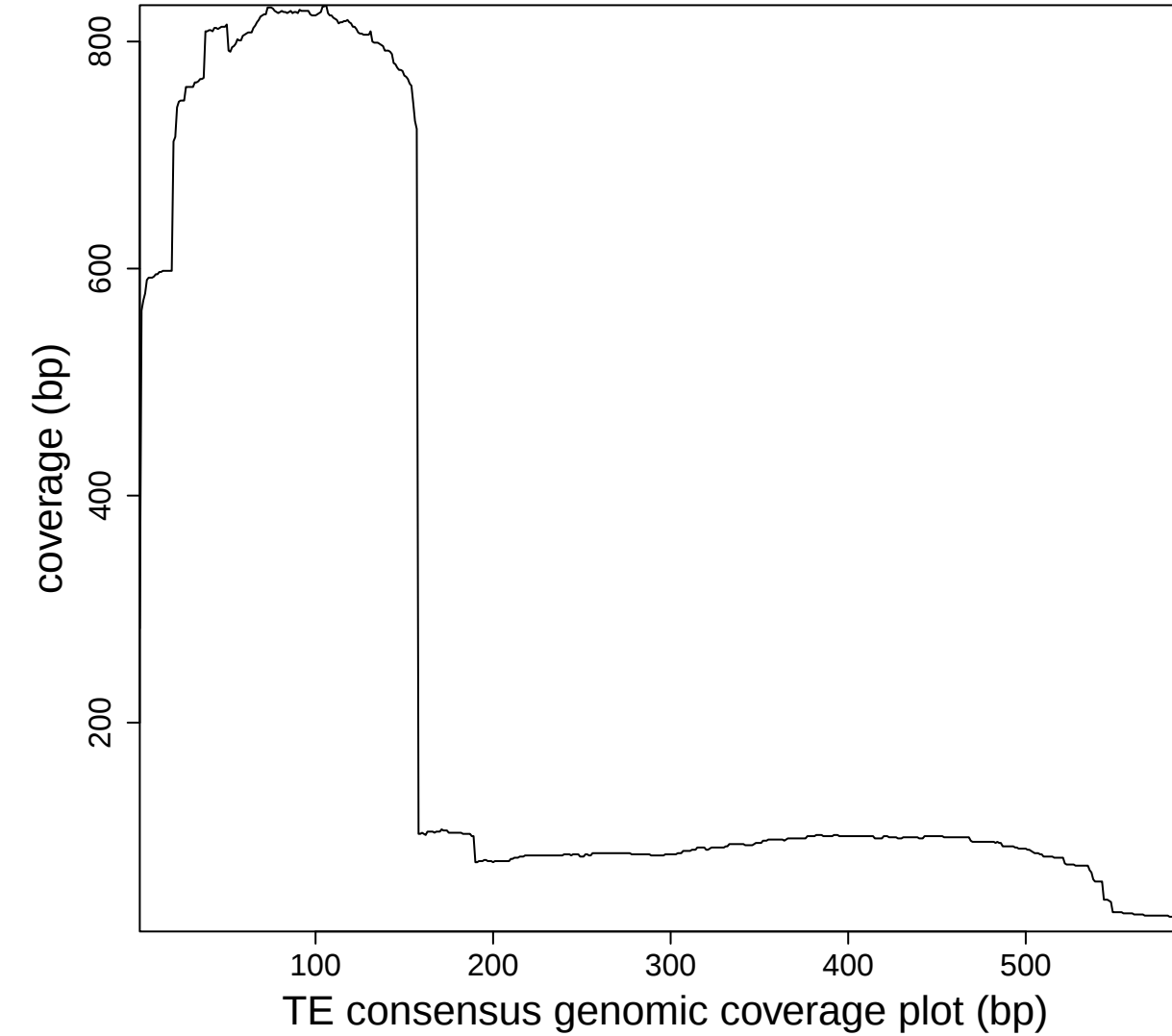
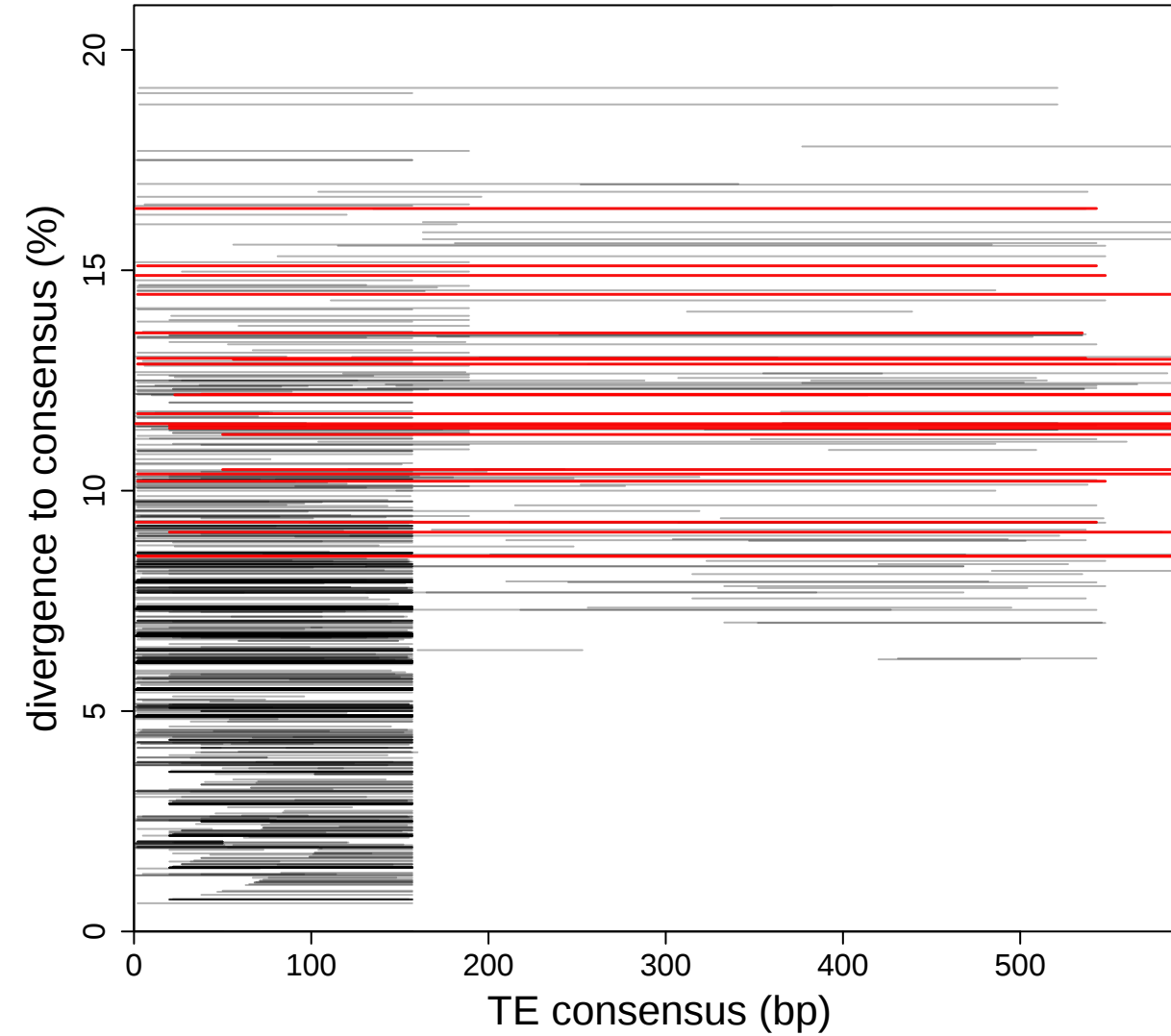


9.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 889bp; fragments: 1315; full length: 0 (>=800.1bp)

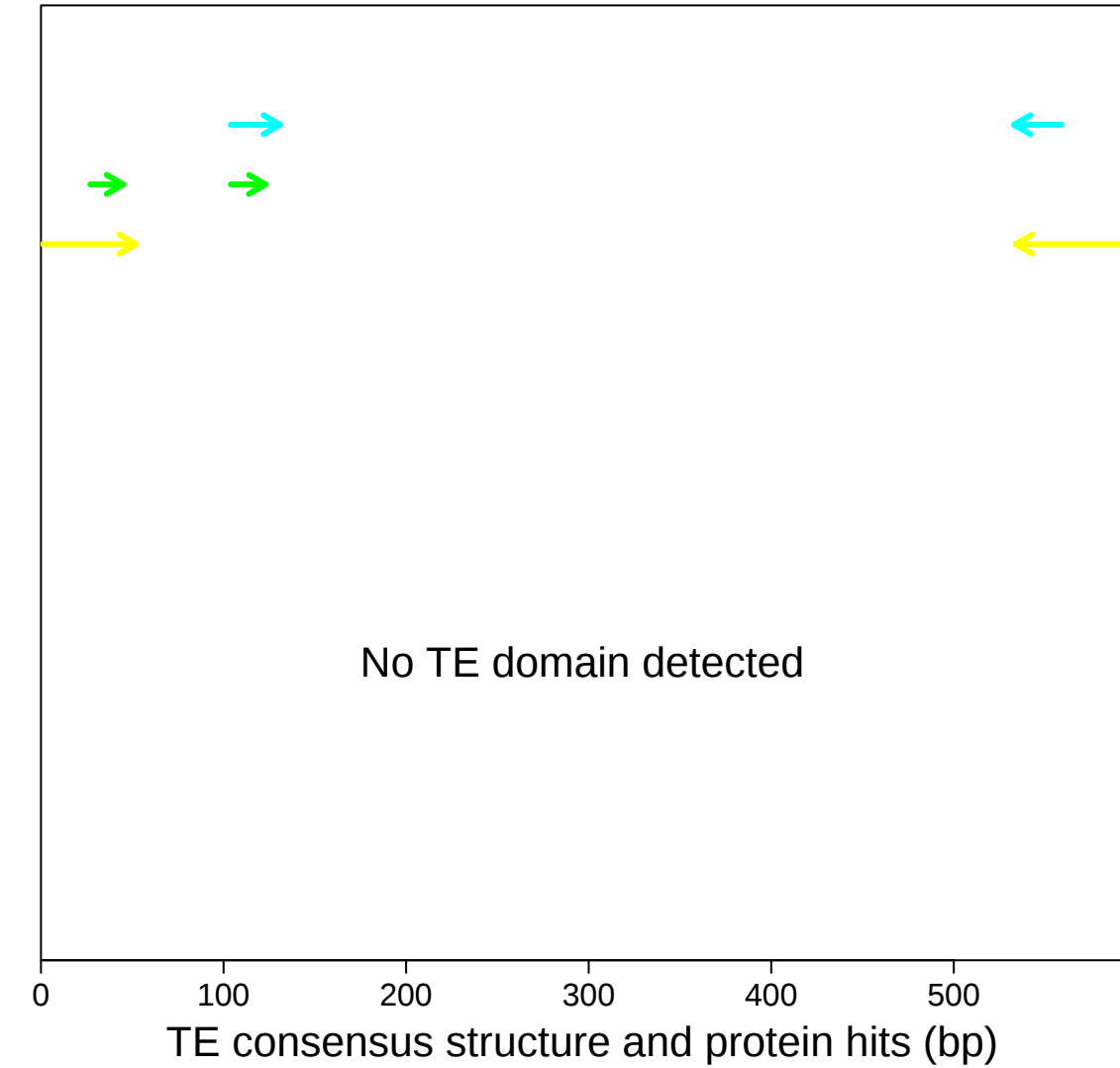
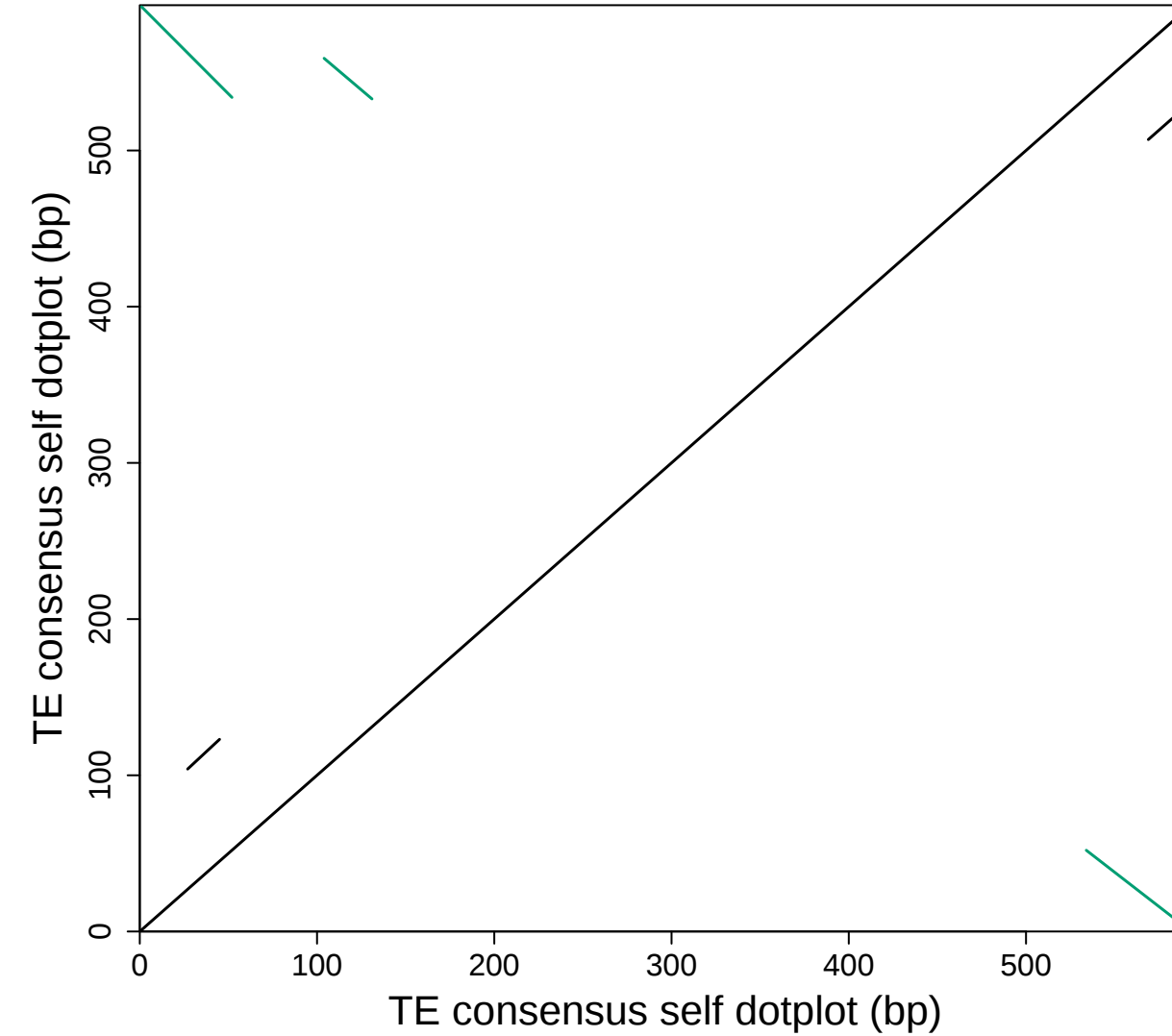
After TEtrimmer Extended plot Blue lines are boundaries



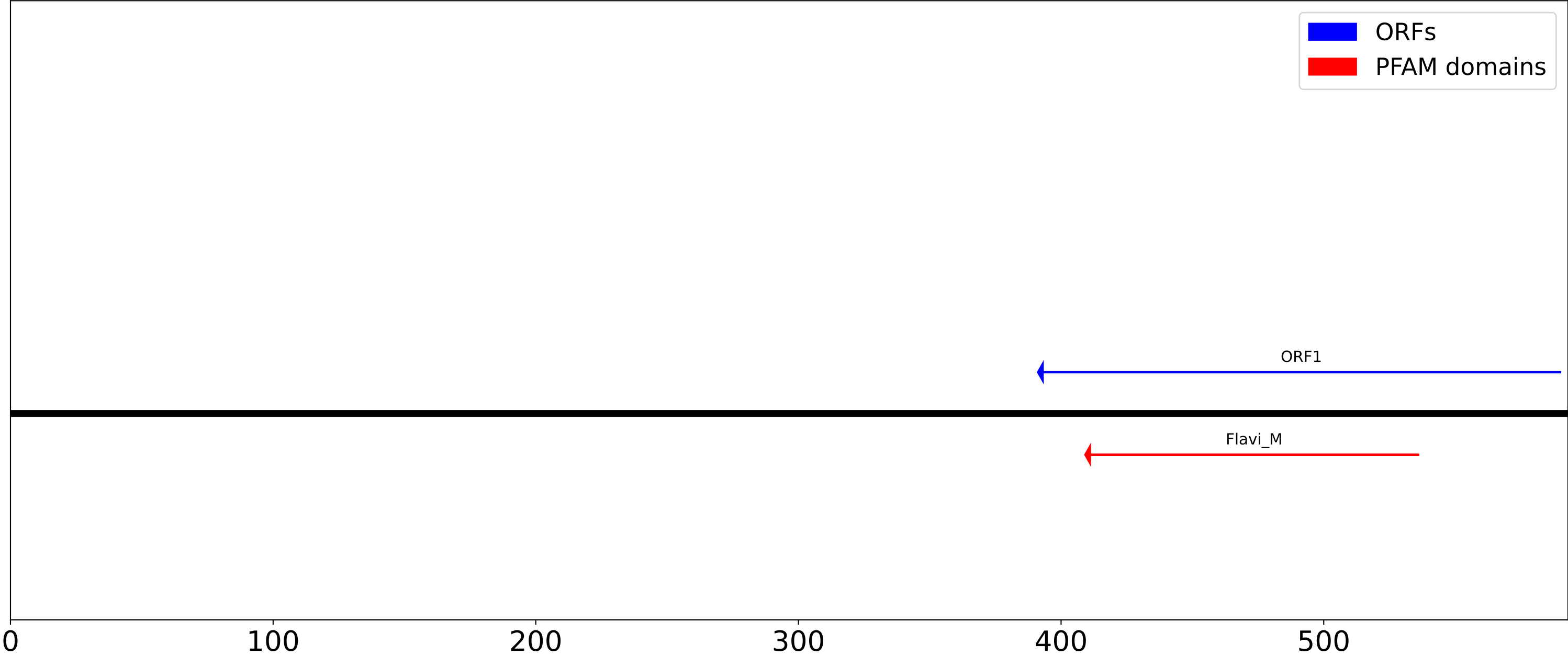
TE: rnd_4_family_1569
size: 593bp; fragments: 975; full length: 22 (≥ 533.7 bp)



Before TEtrimmer 593 bp



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

