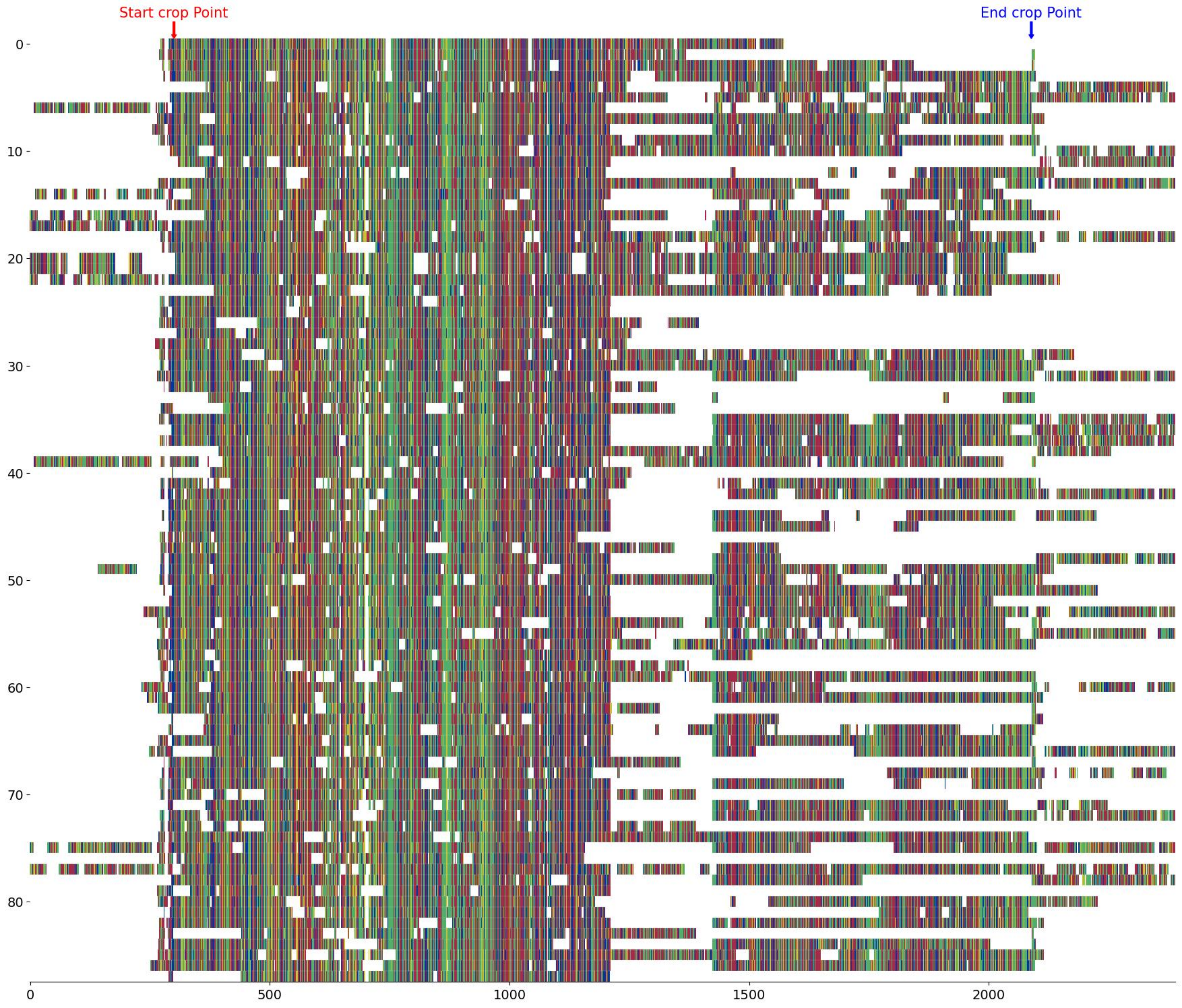


End crop Point



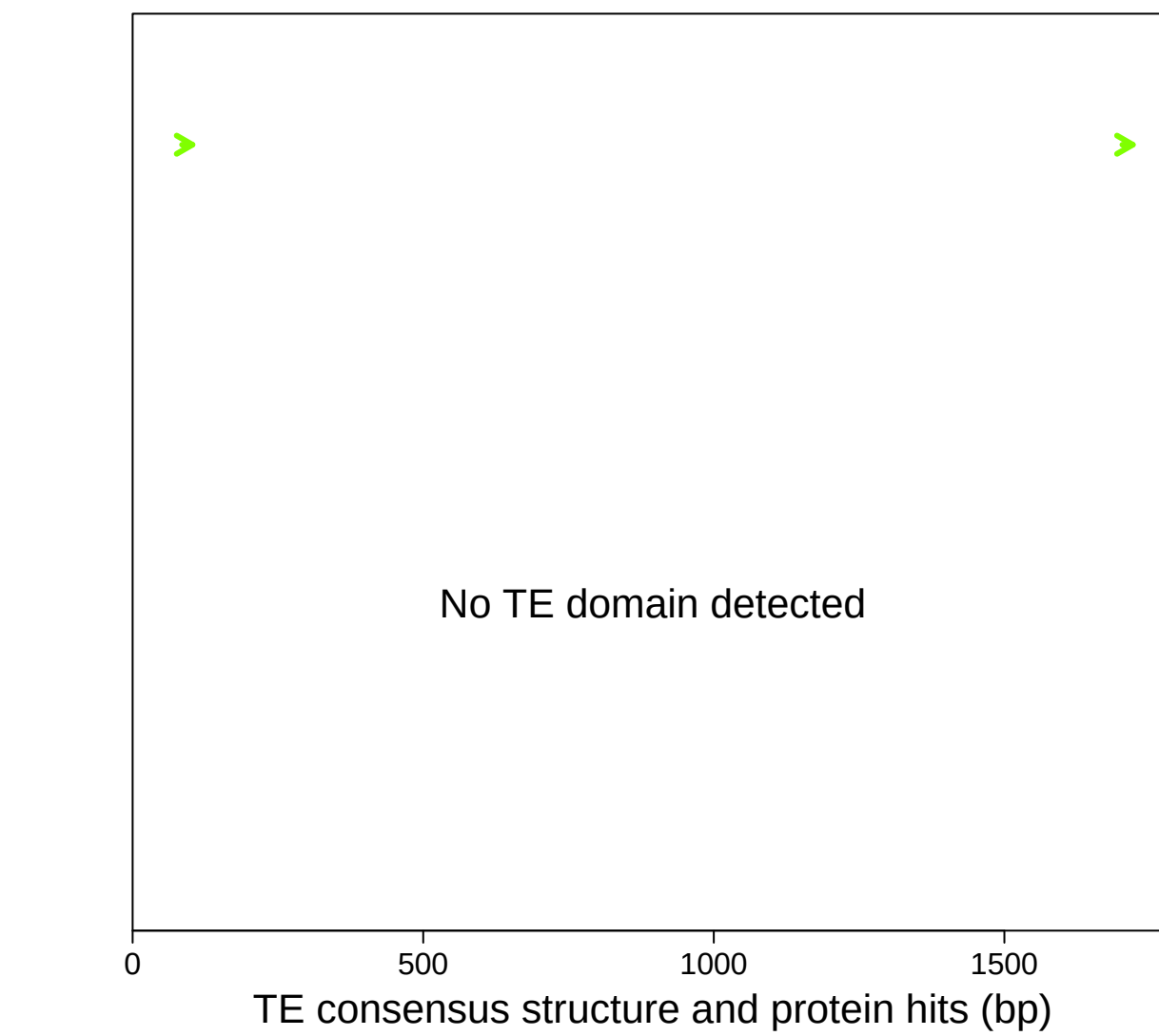
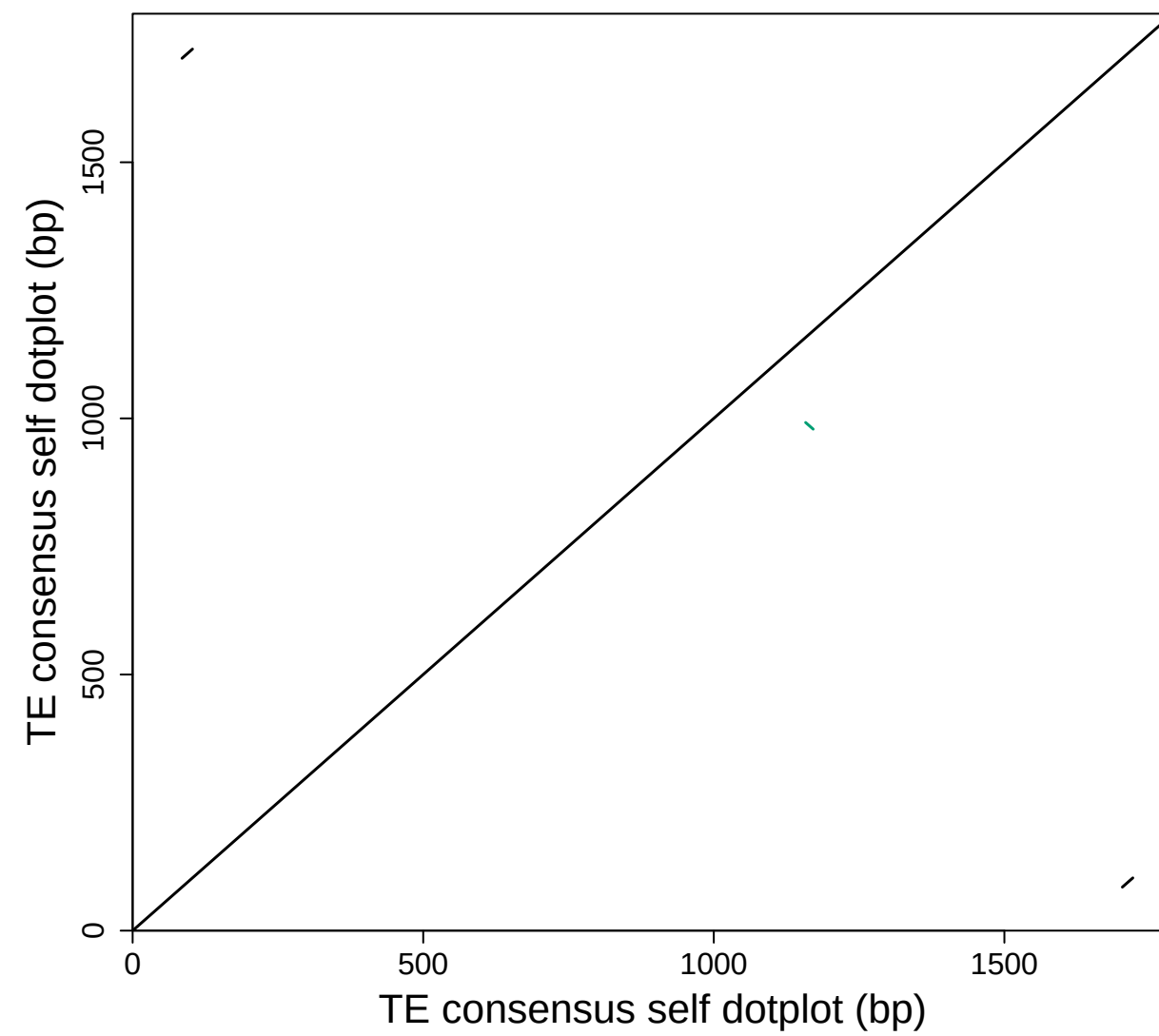
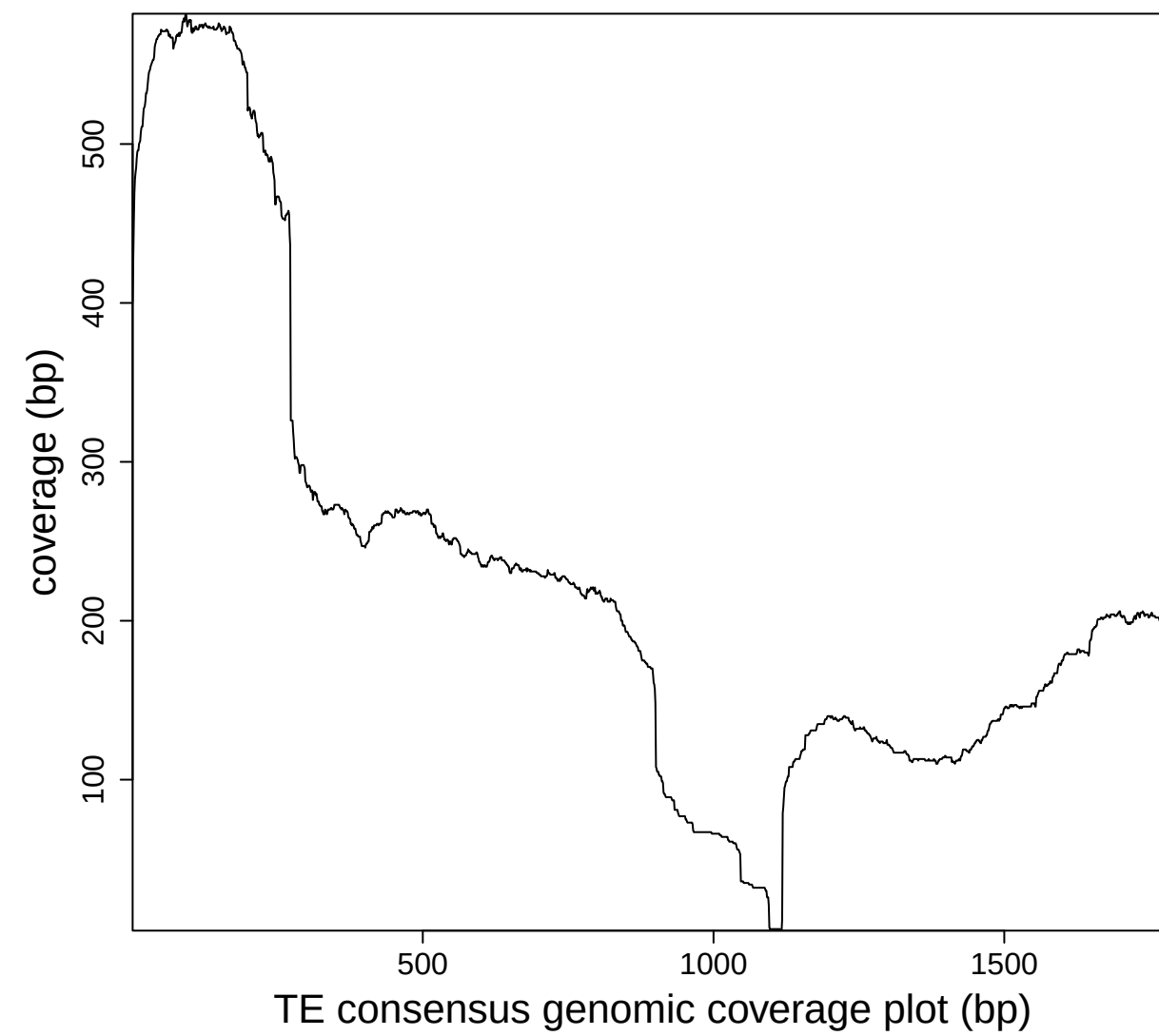
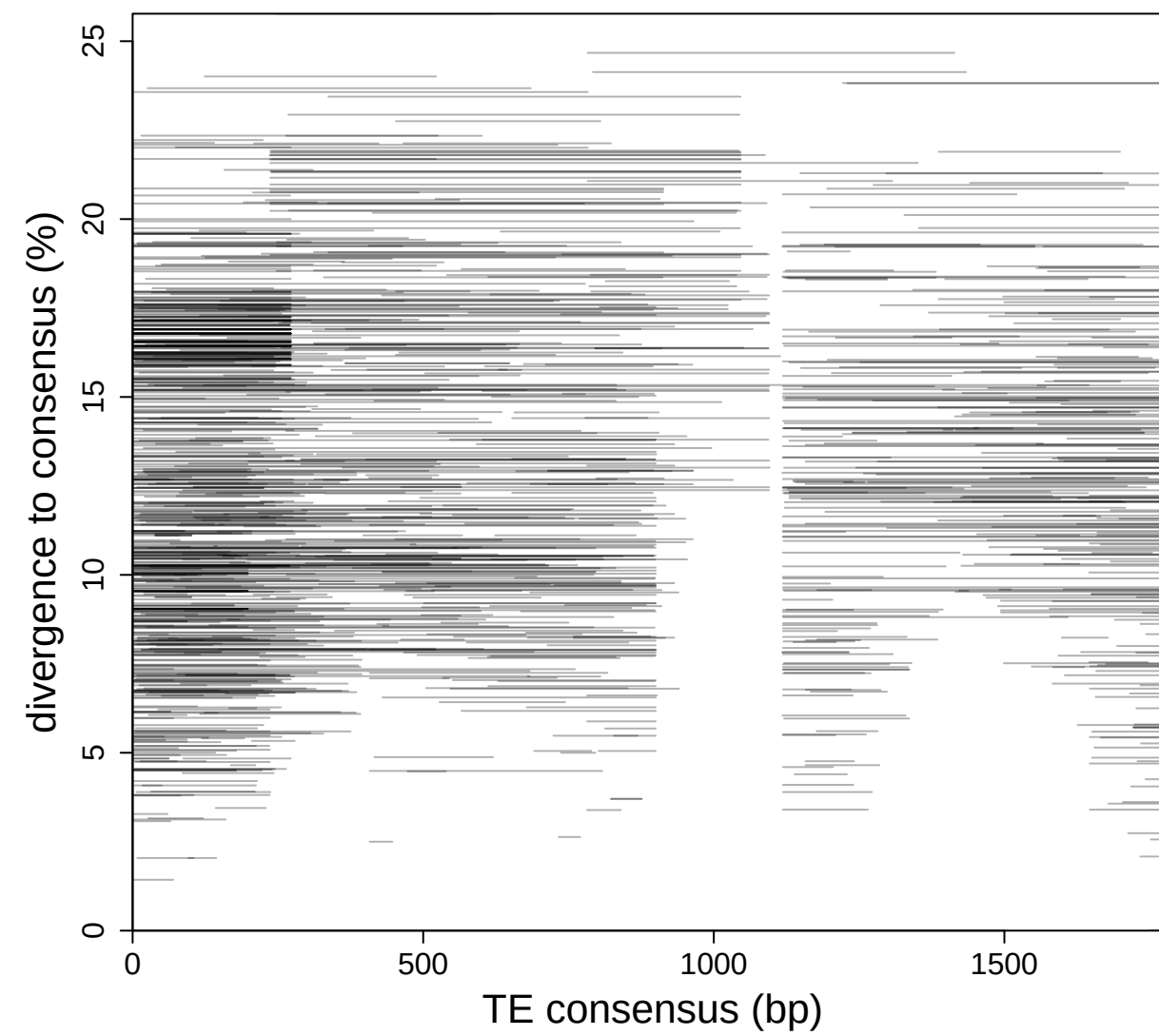
Start crop Point

End crop Point



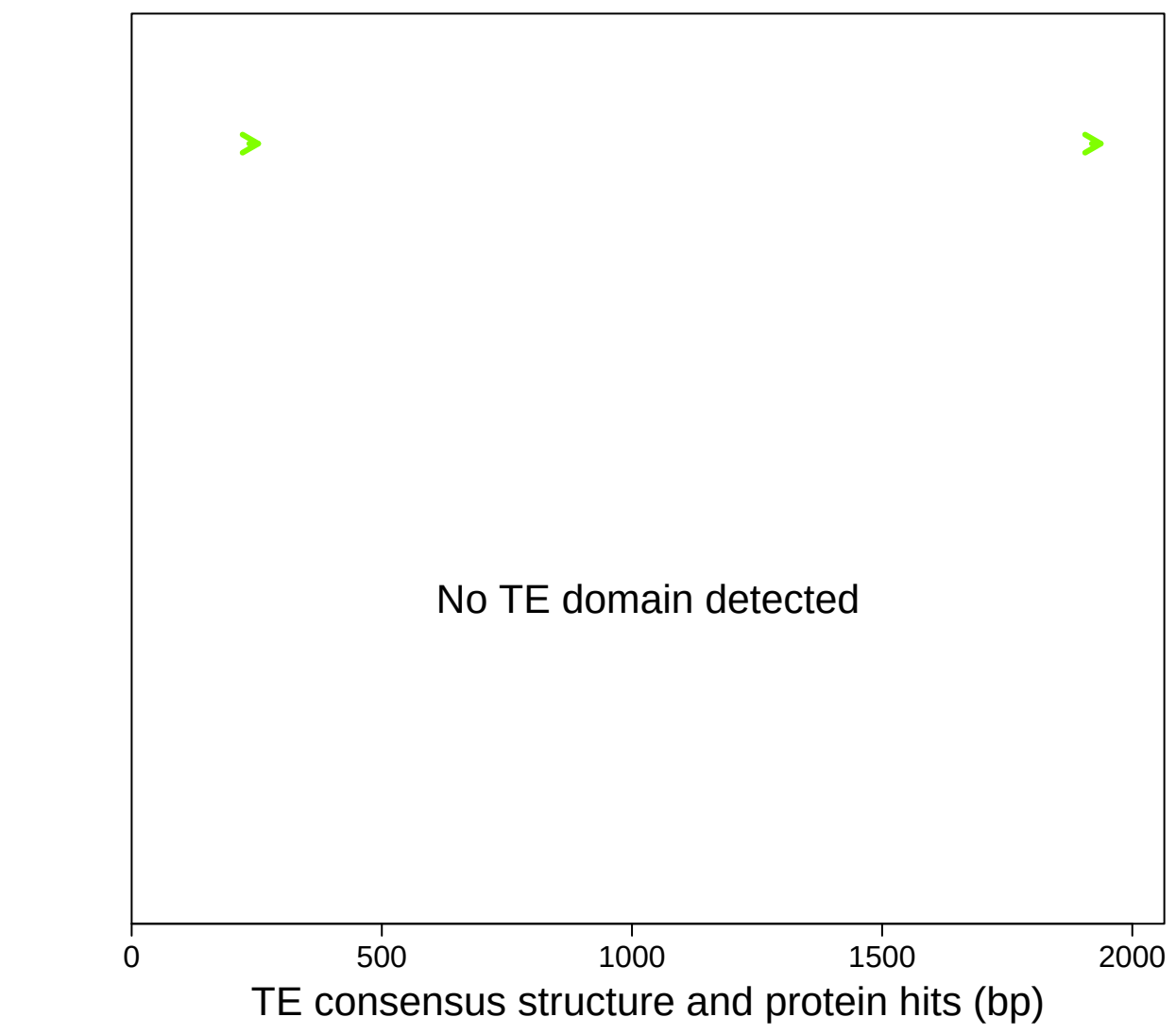
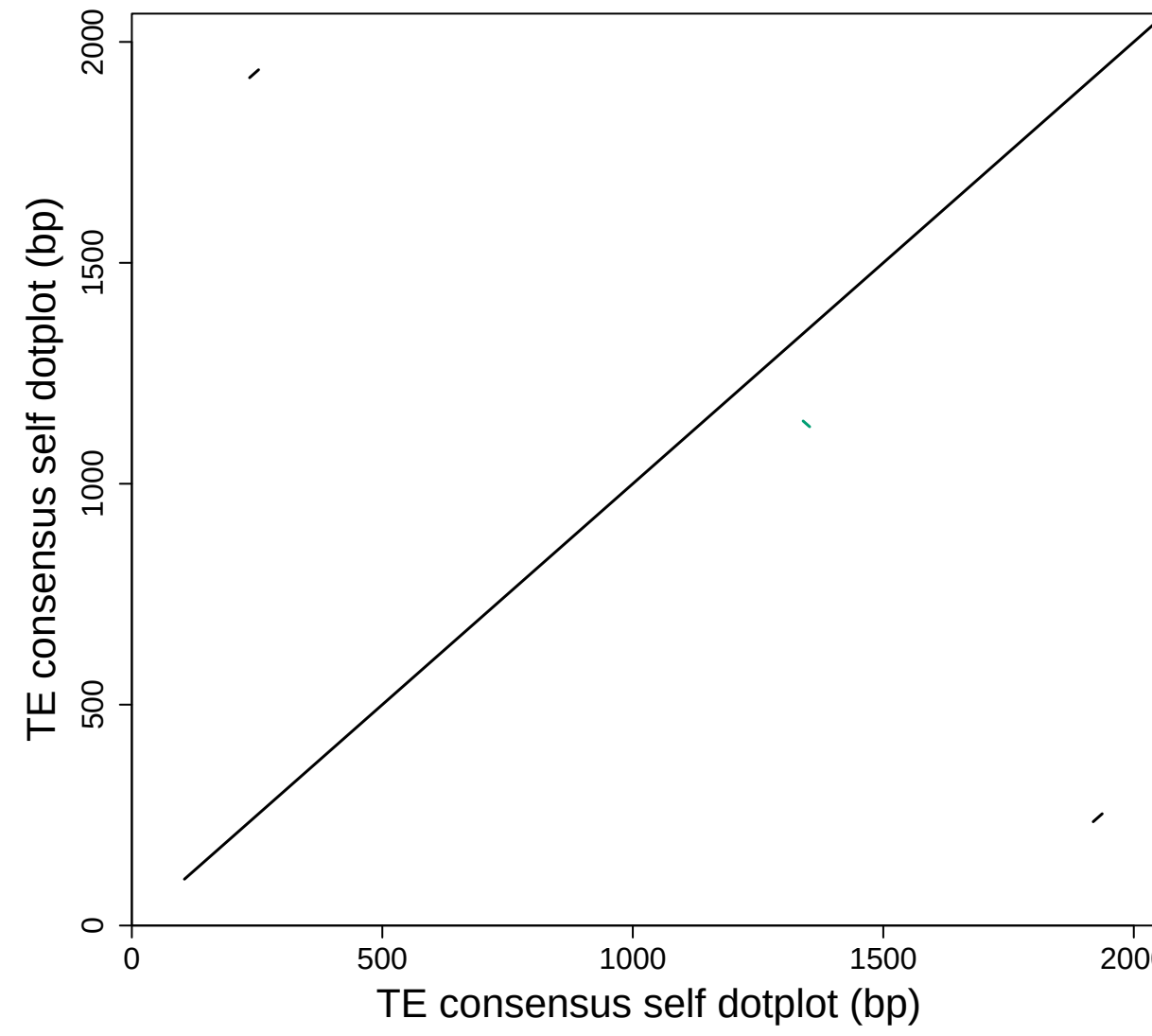
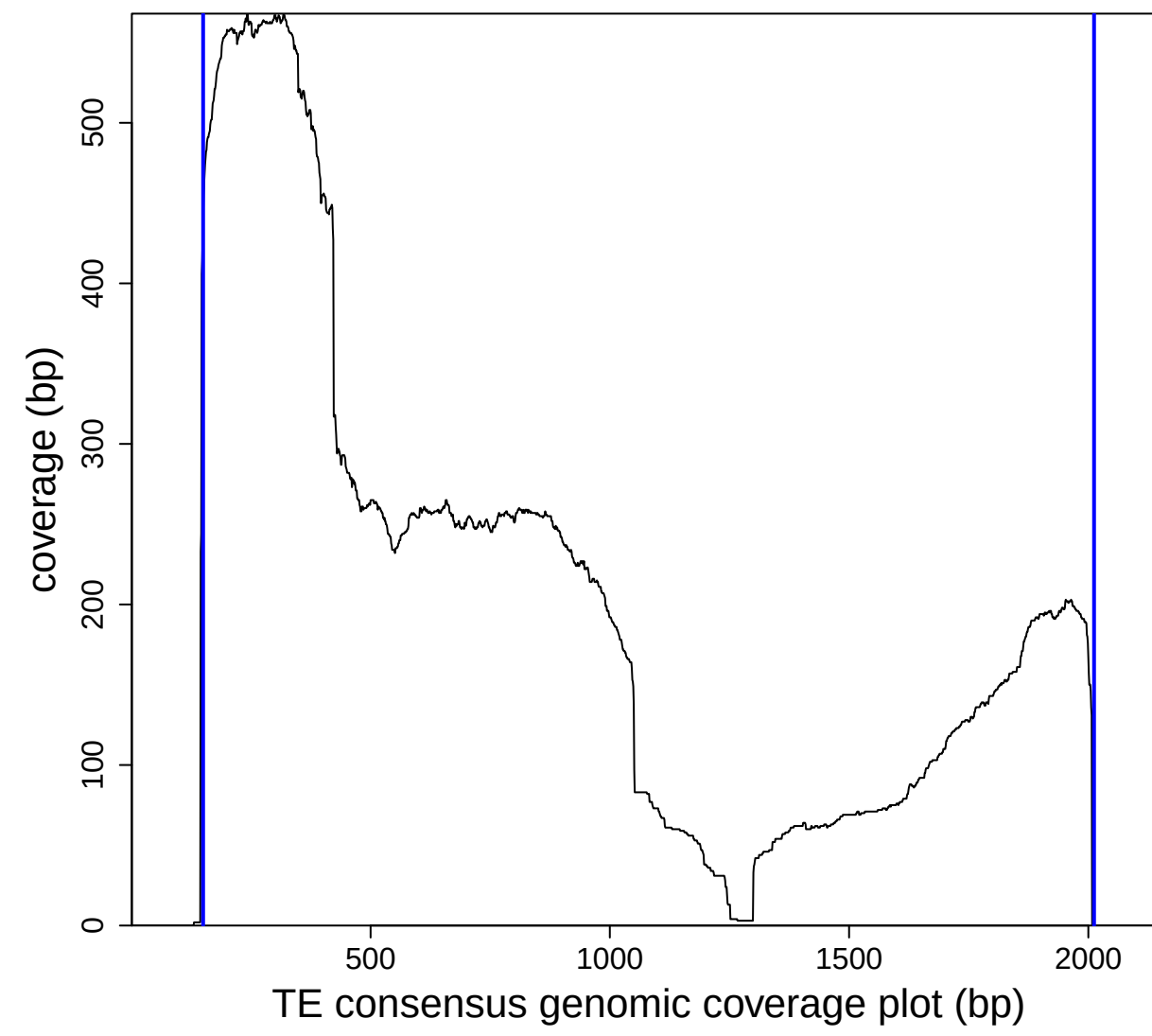
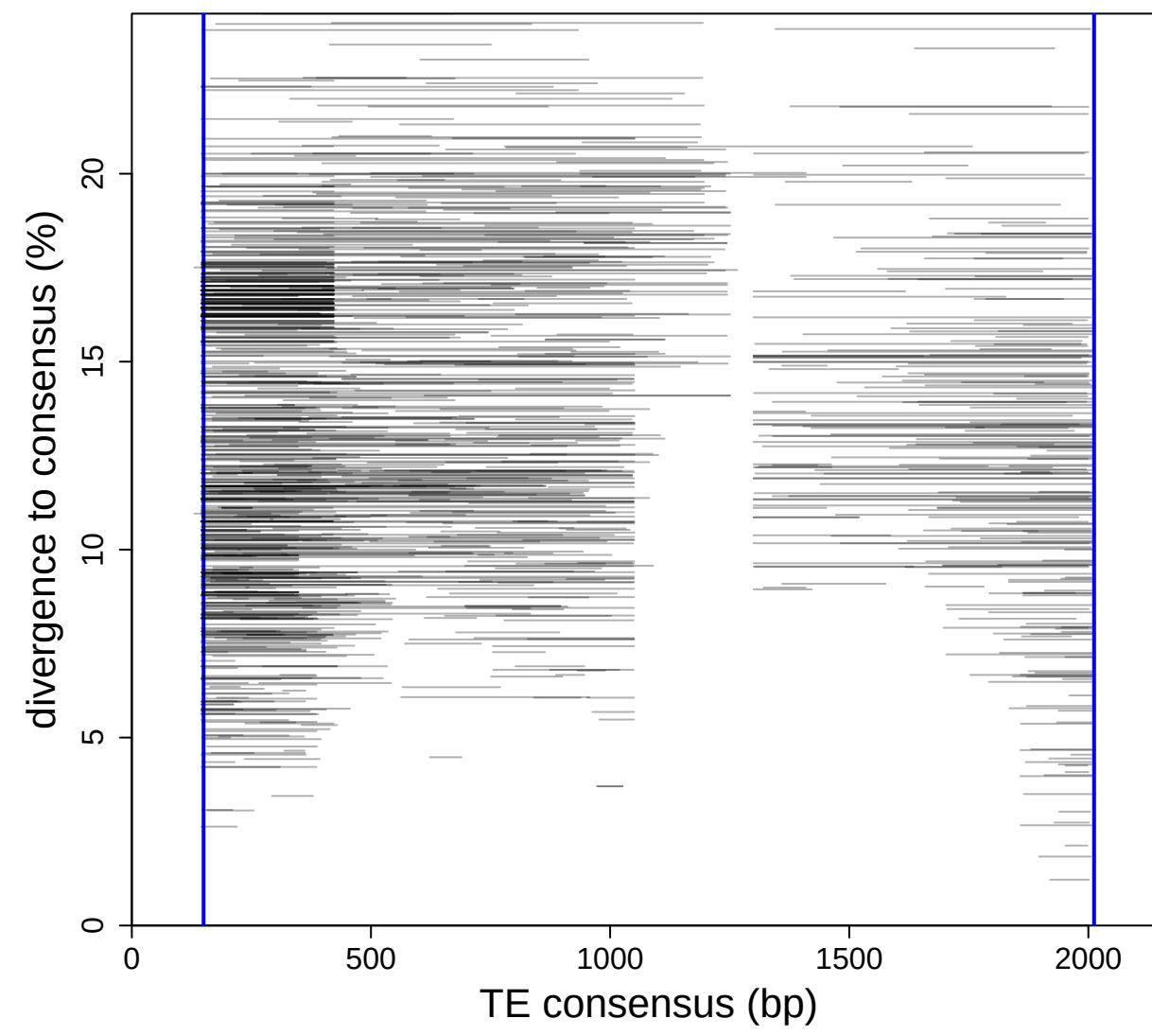
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1790bp; fragments: 1241; full length: 0 (>=1611bp)

After TEtrimmer 1790 bp



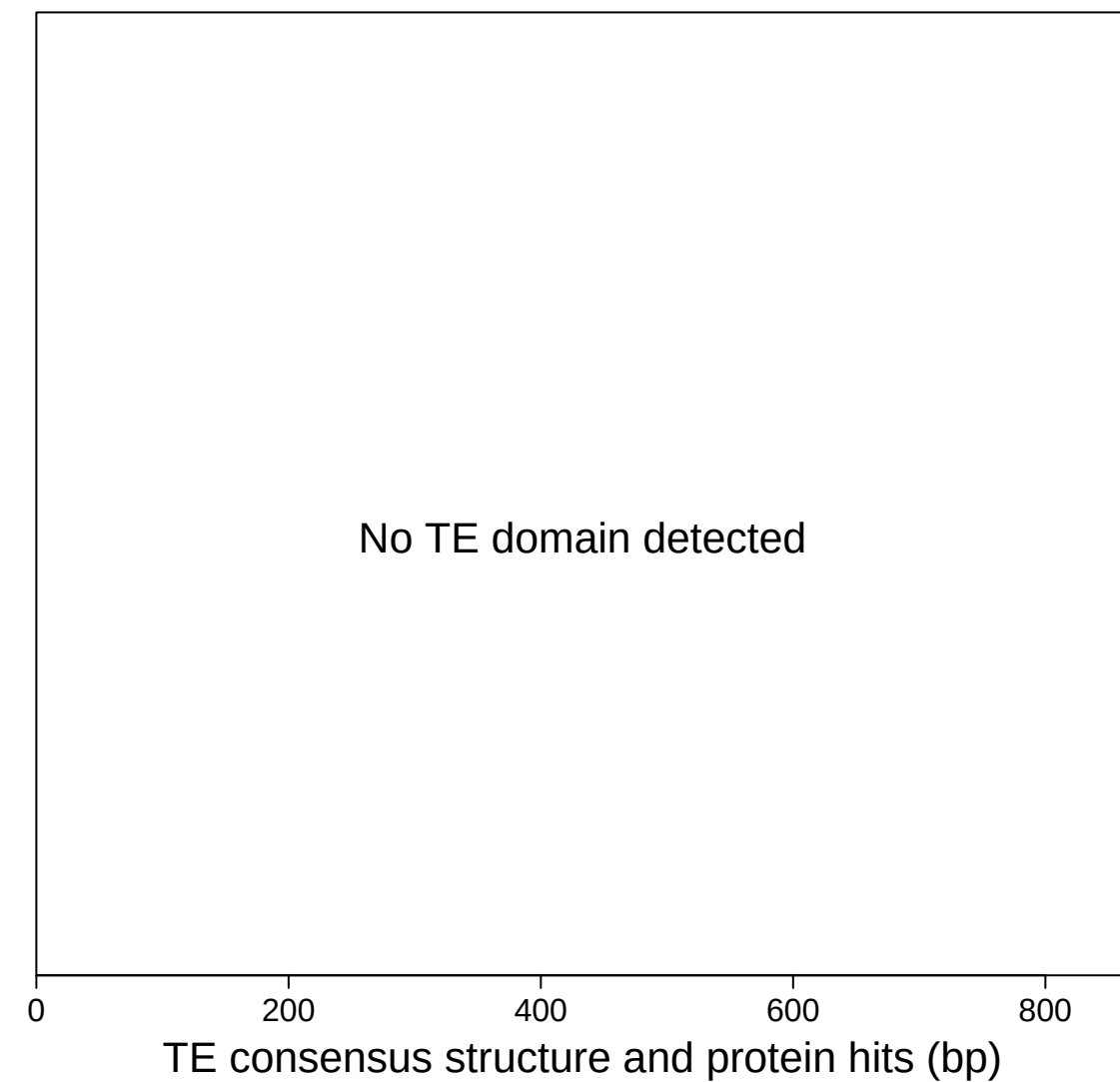
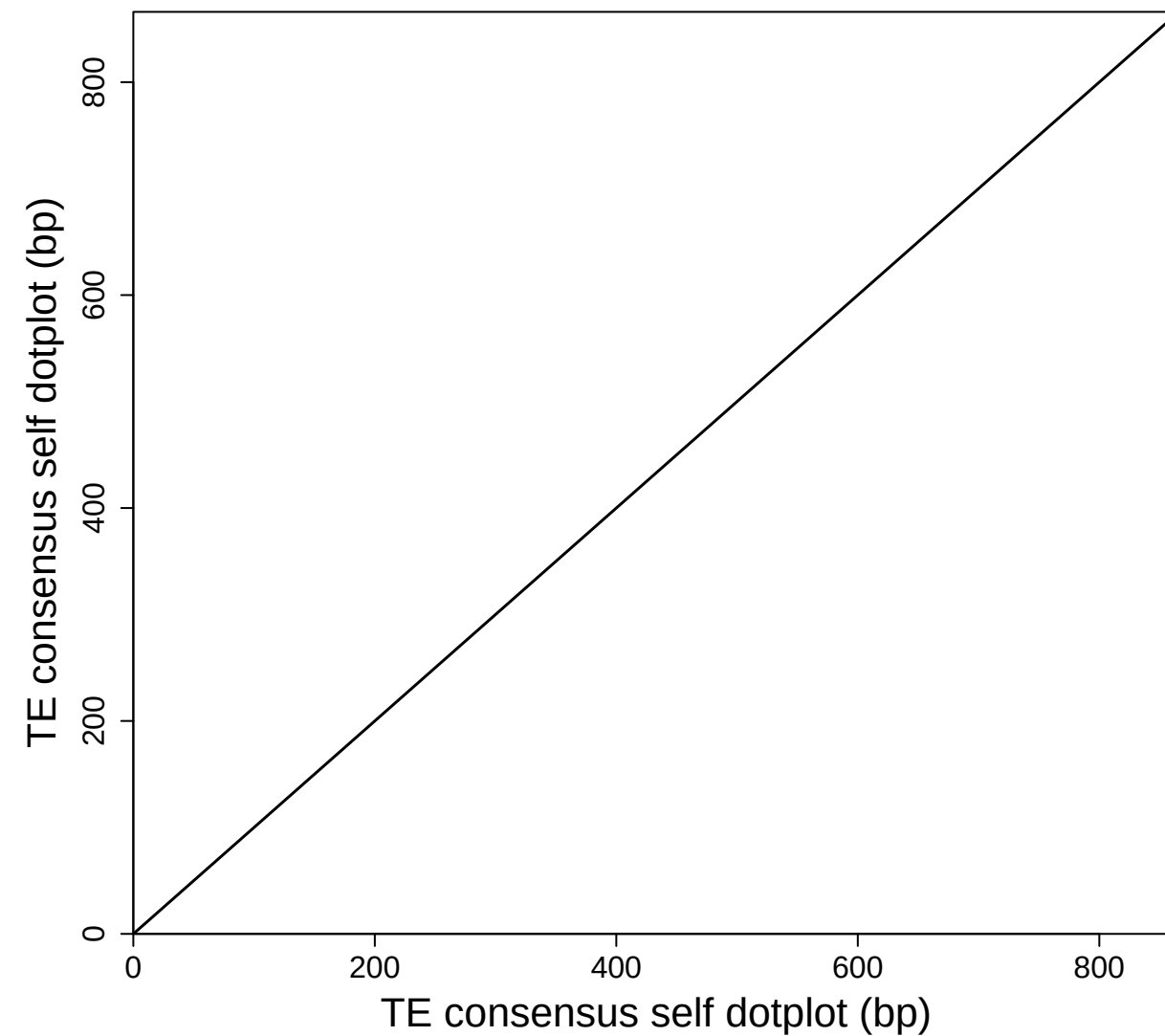
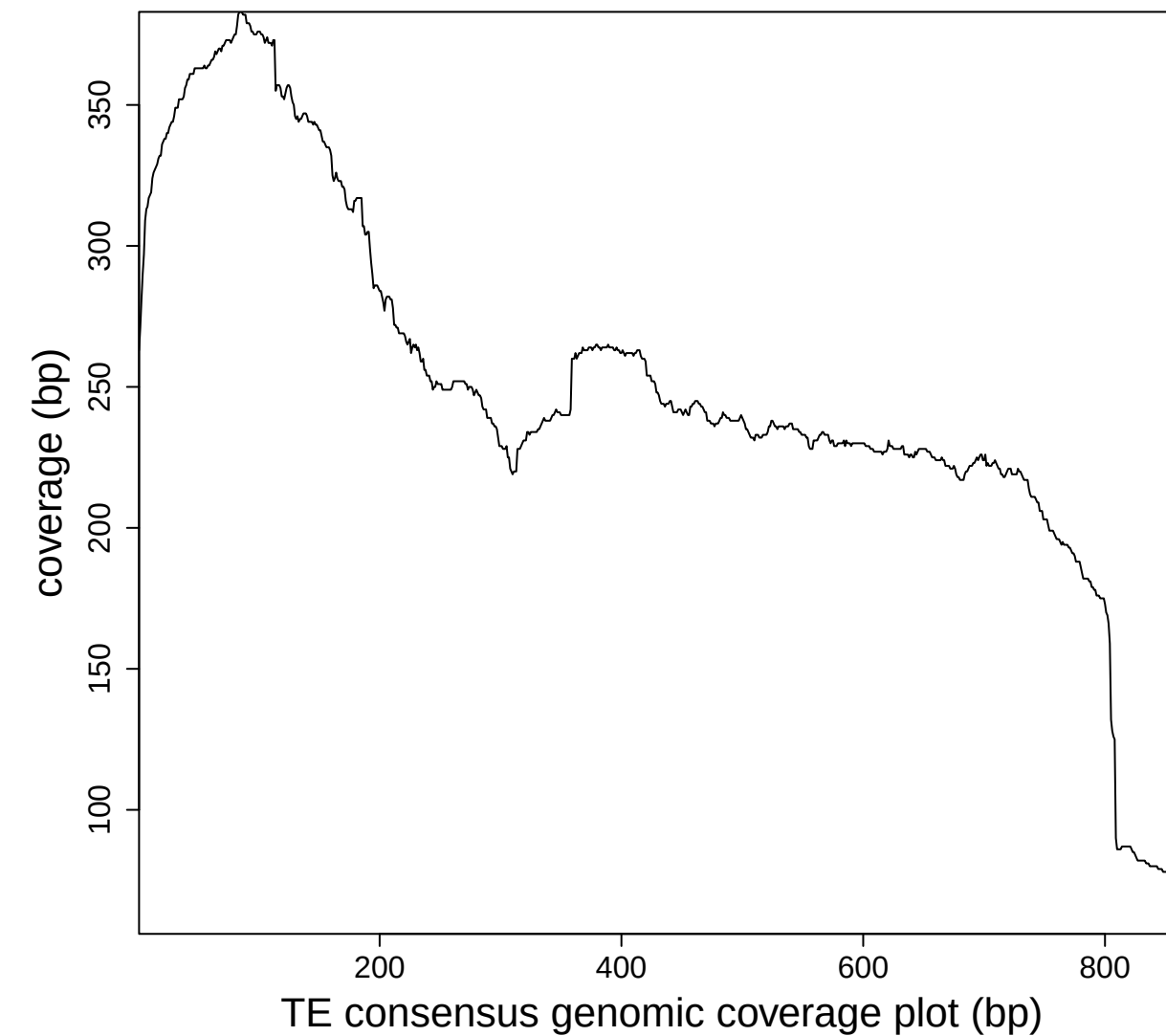
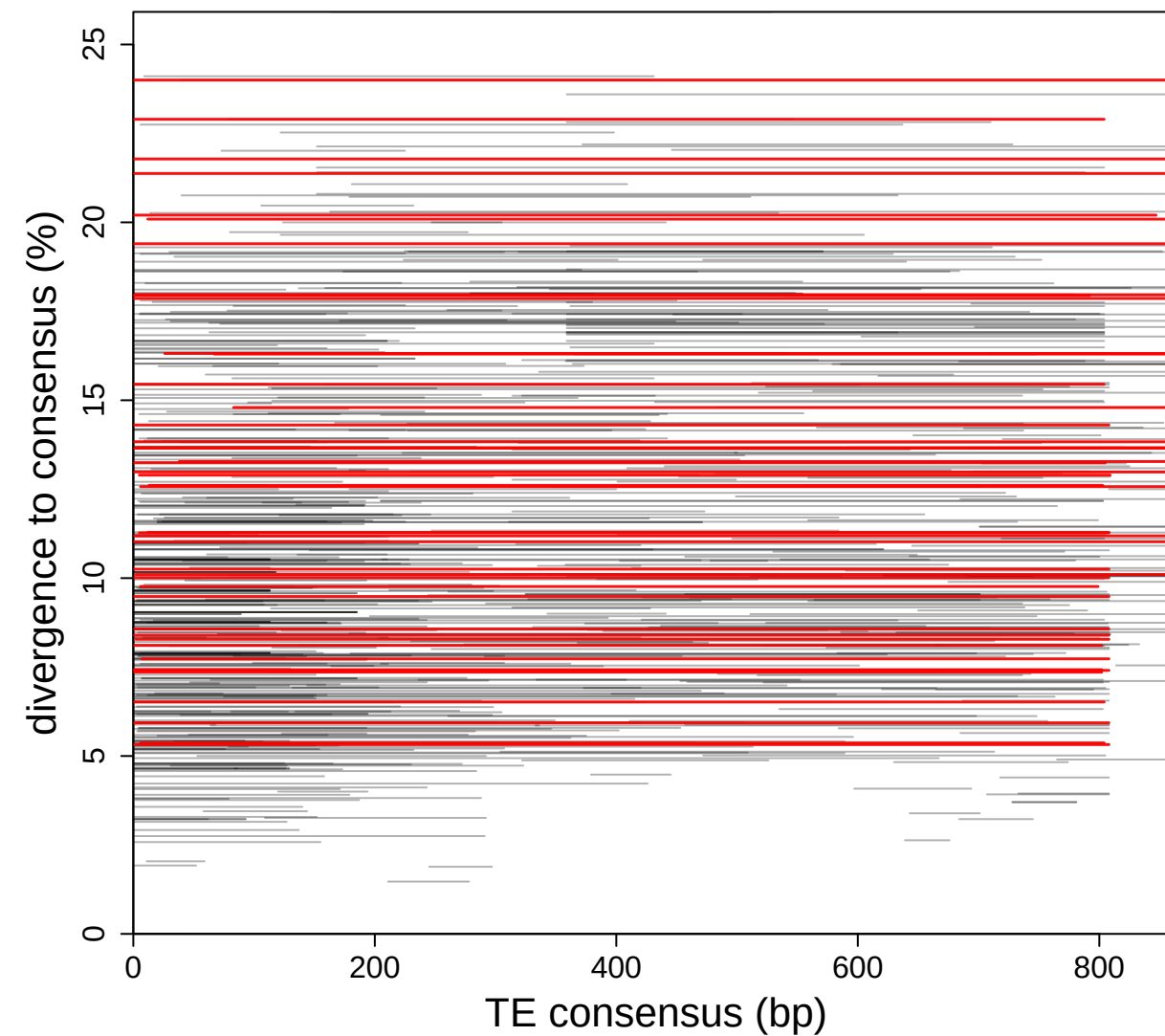
7.fasta.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 2162bp; fragments: 1160; full length: 0 (>=1945.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

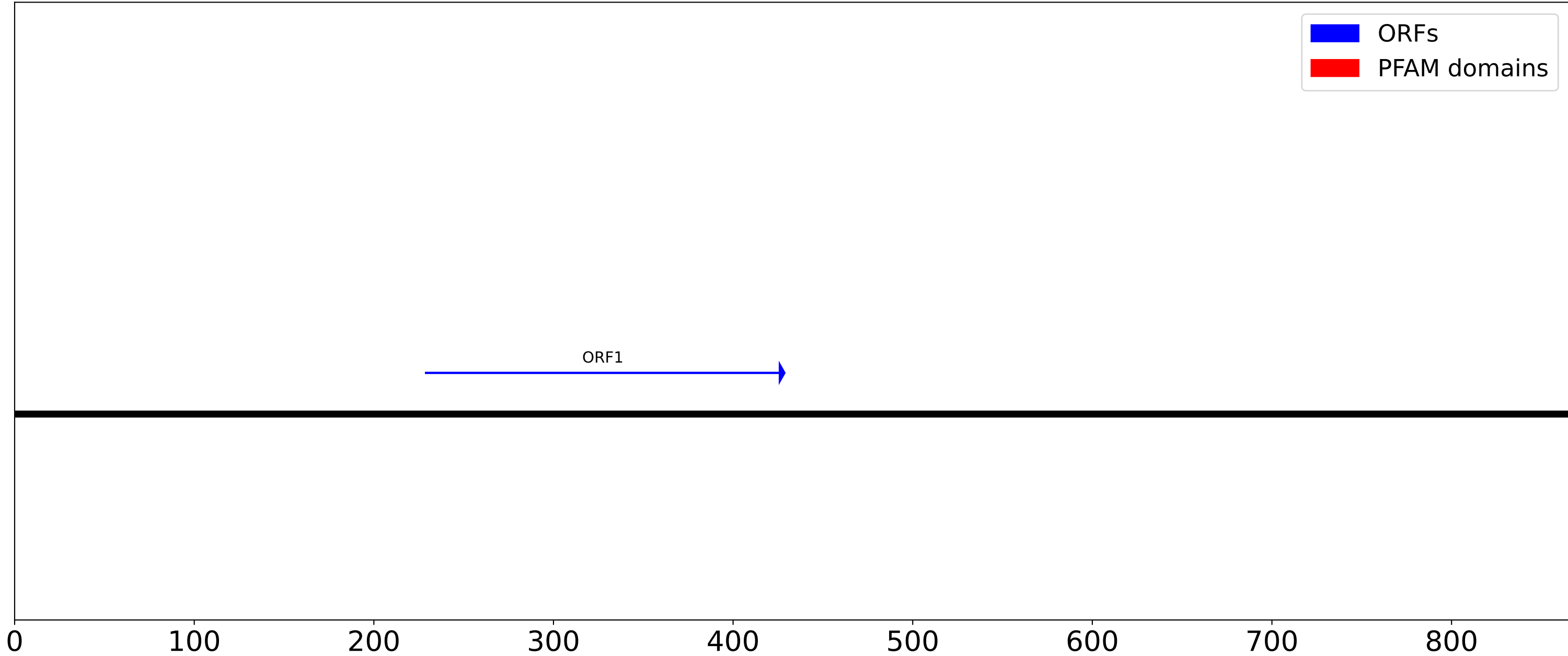


TE: rnd_5_family_1007
size: 866bp; fragments: 649; full length: 45 (≥ 779.4 bp)

Before TEtrimmer 866 bp



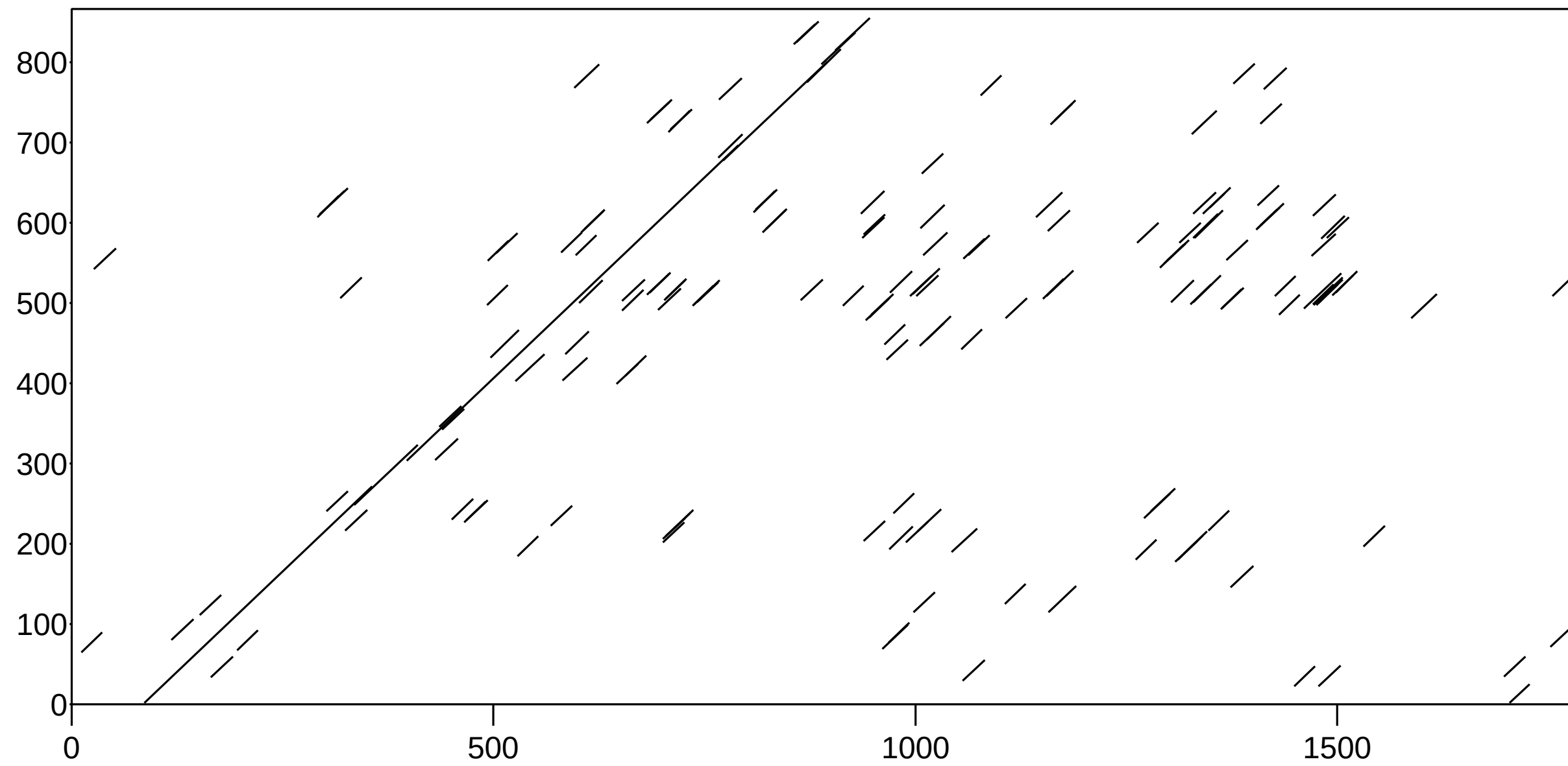
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)