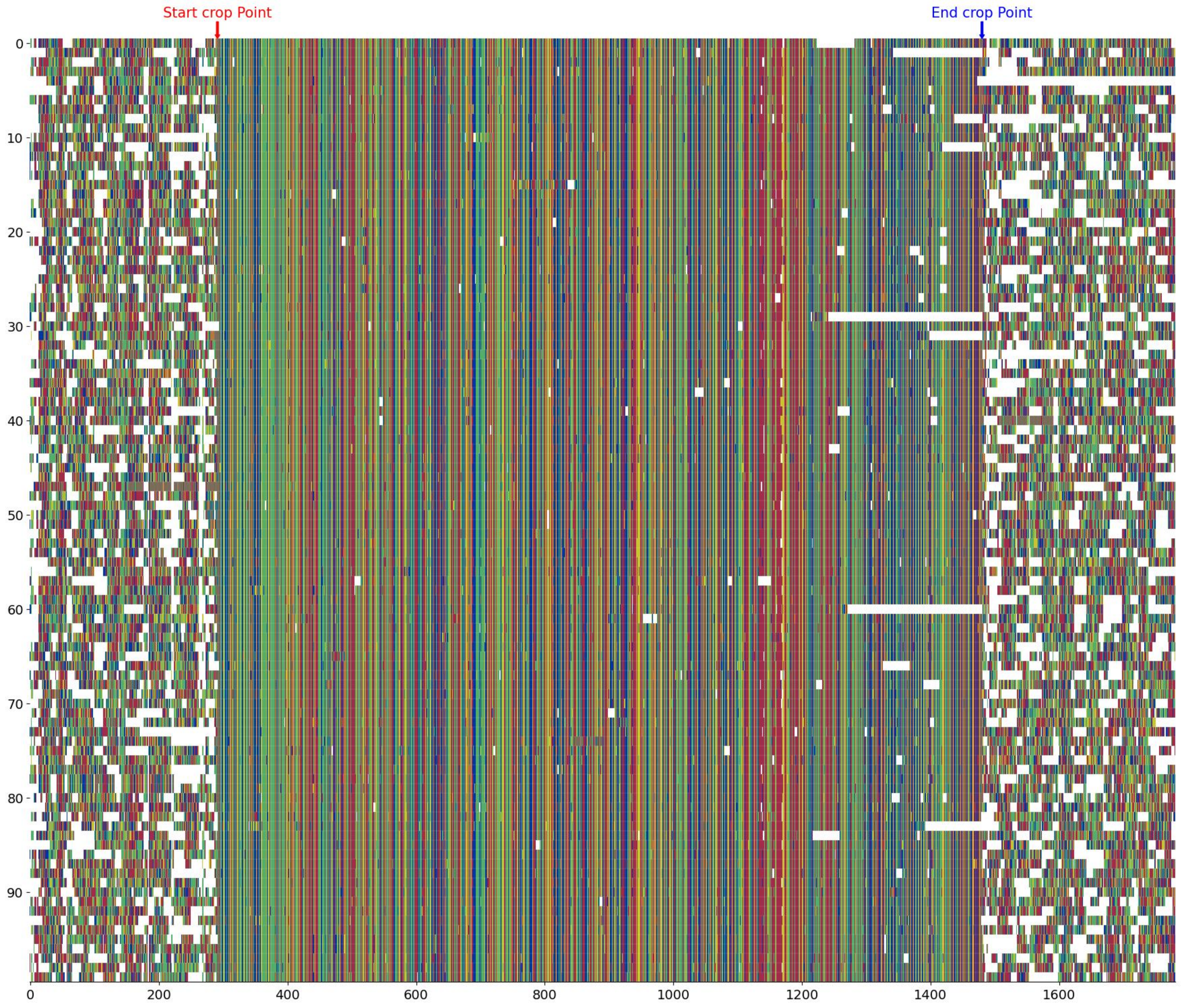




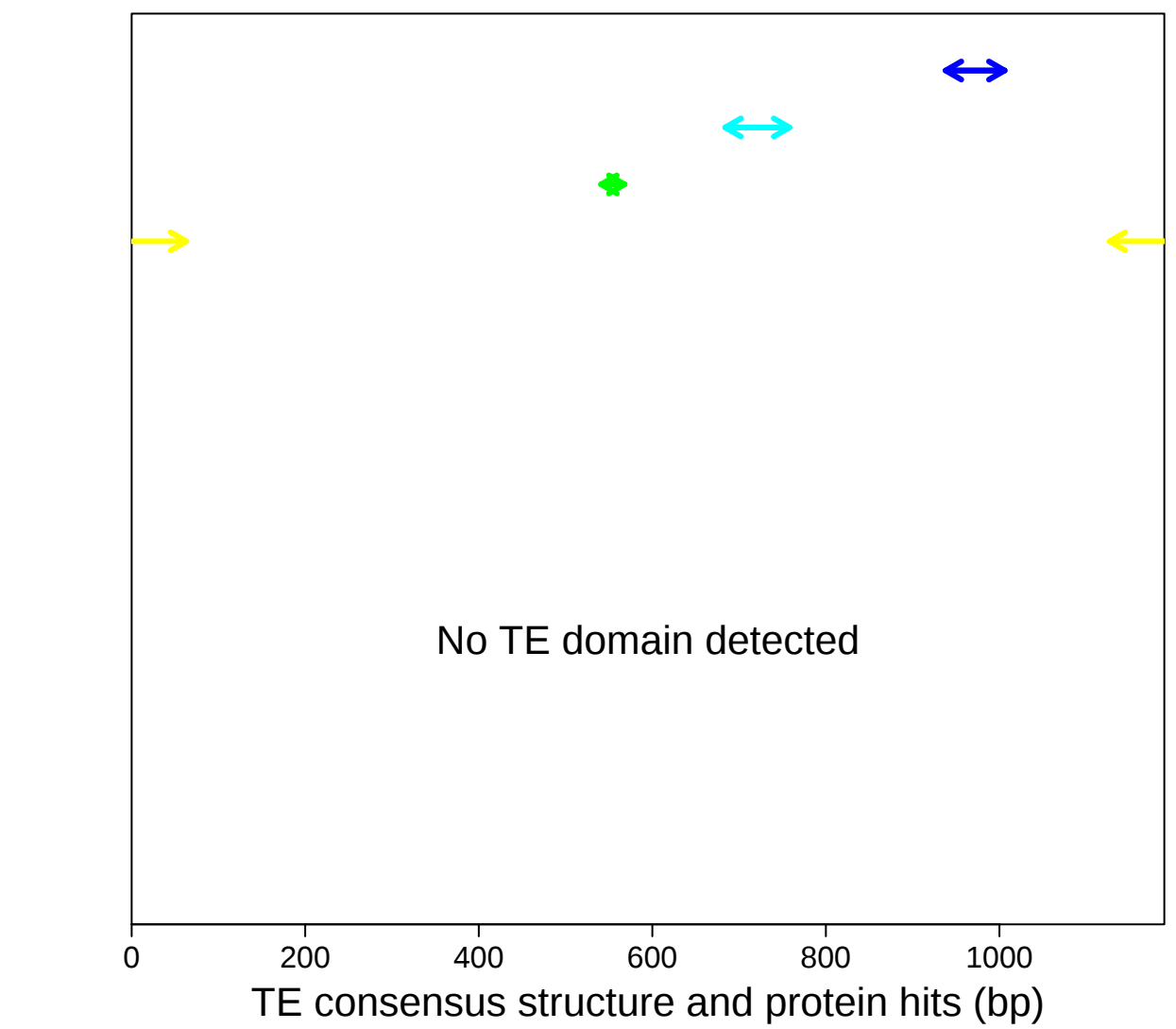
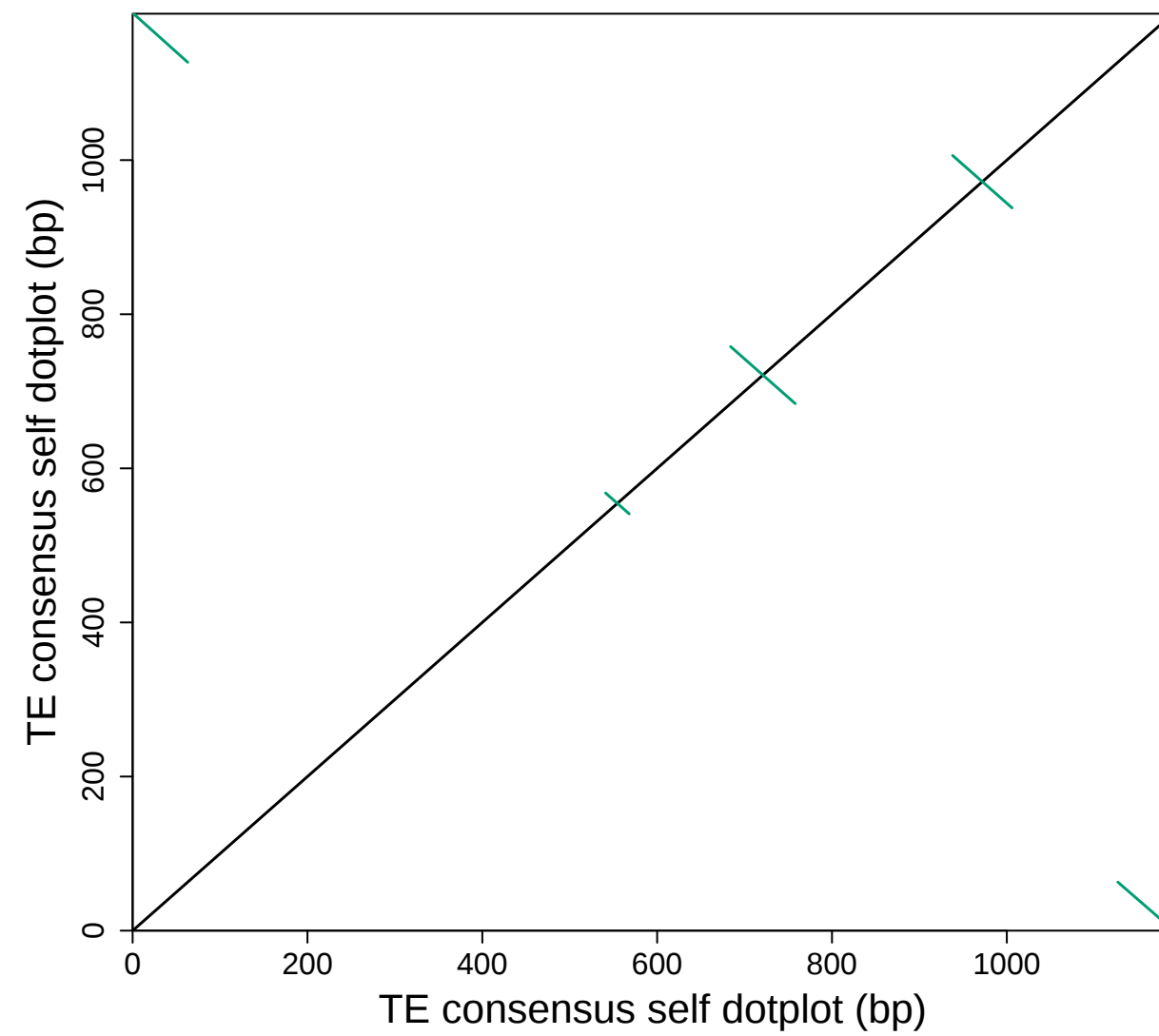
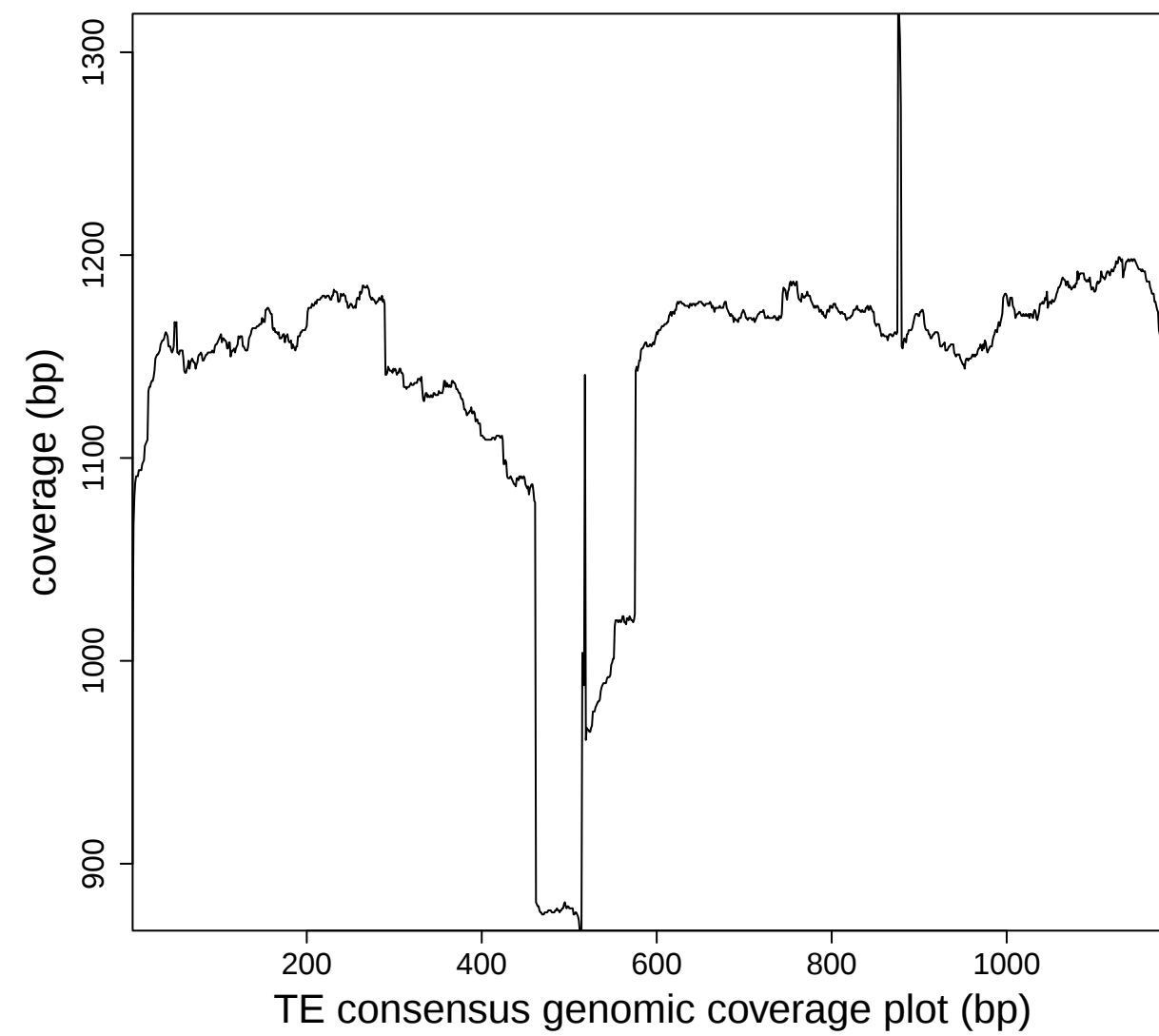
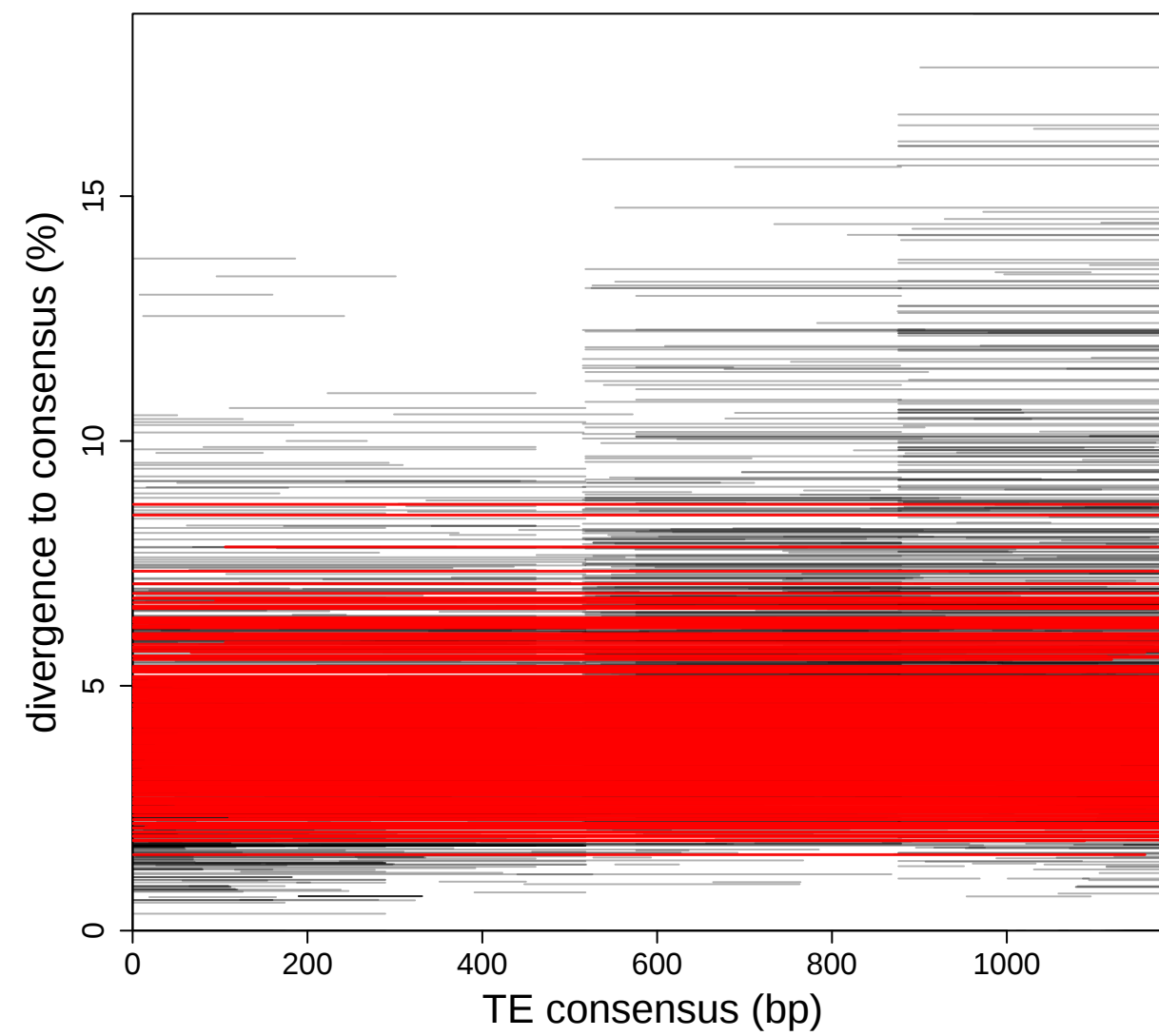
Start crop Point

End crop Point



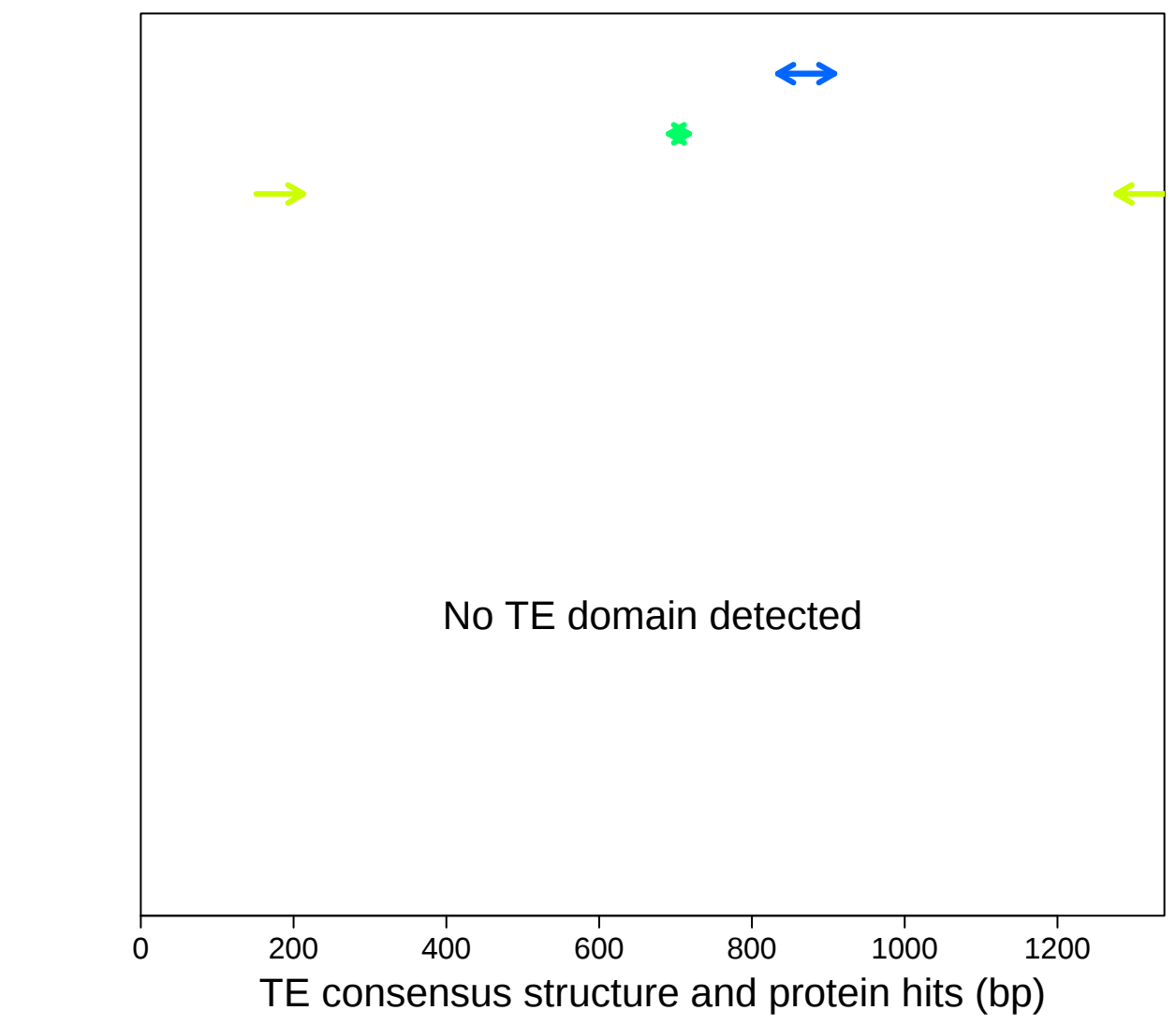
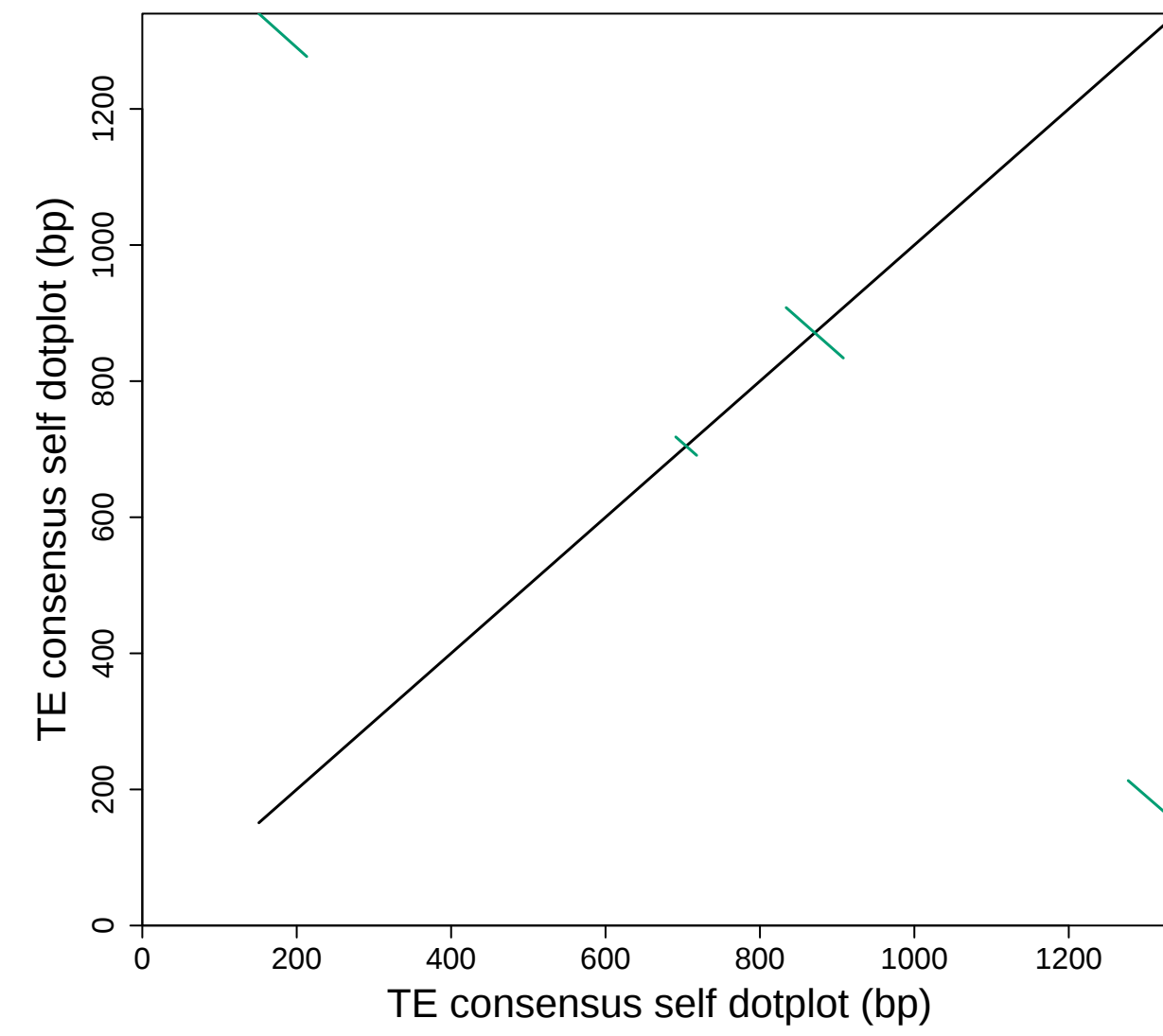
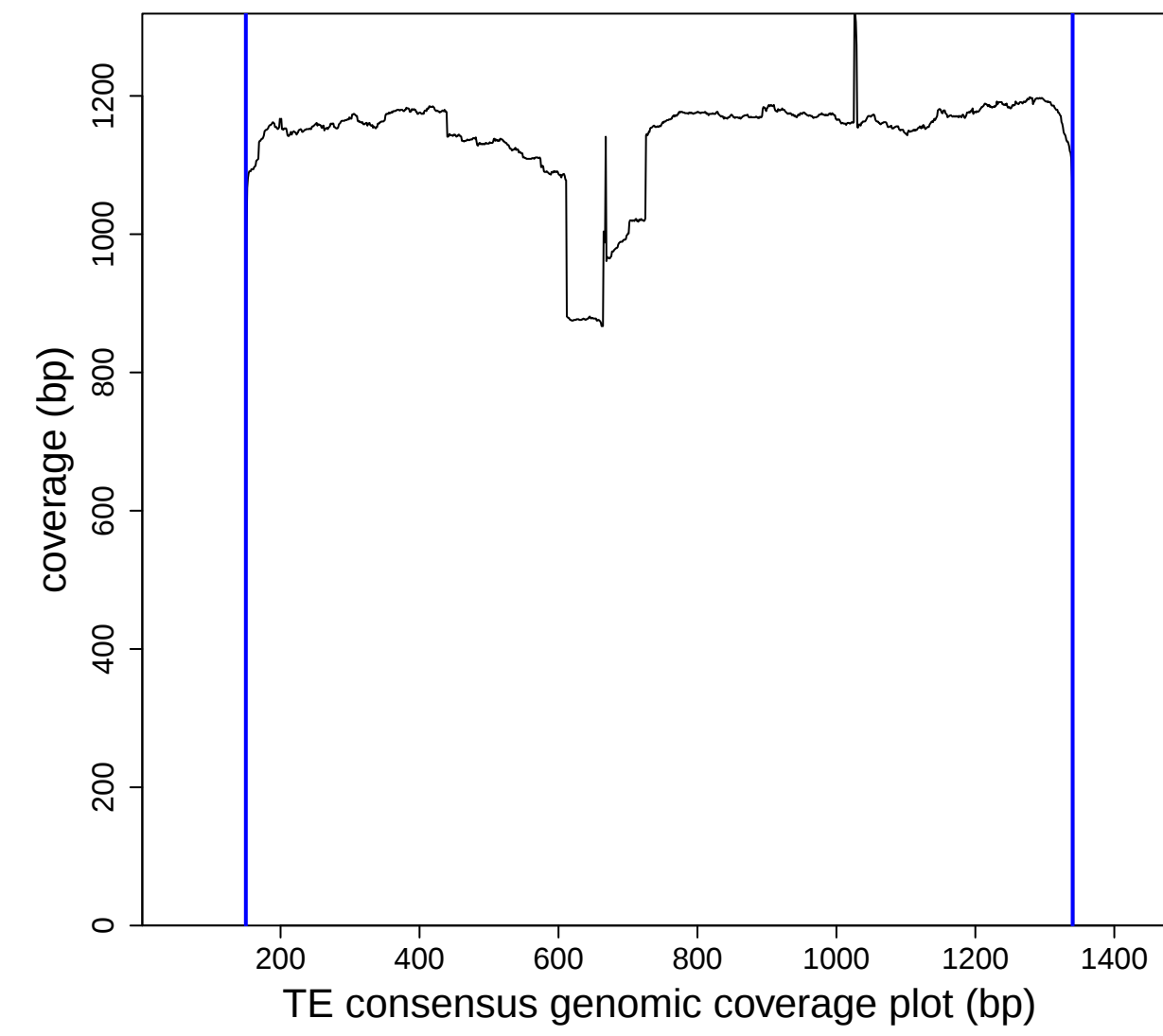
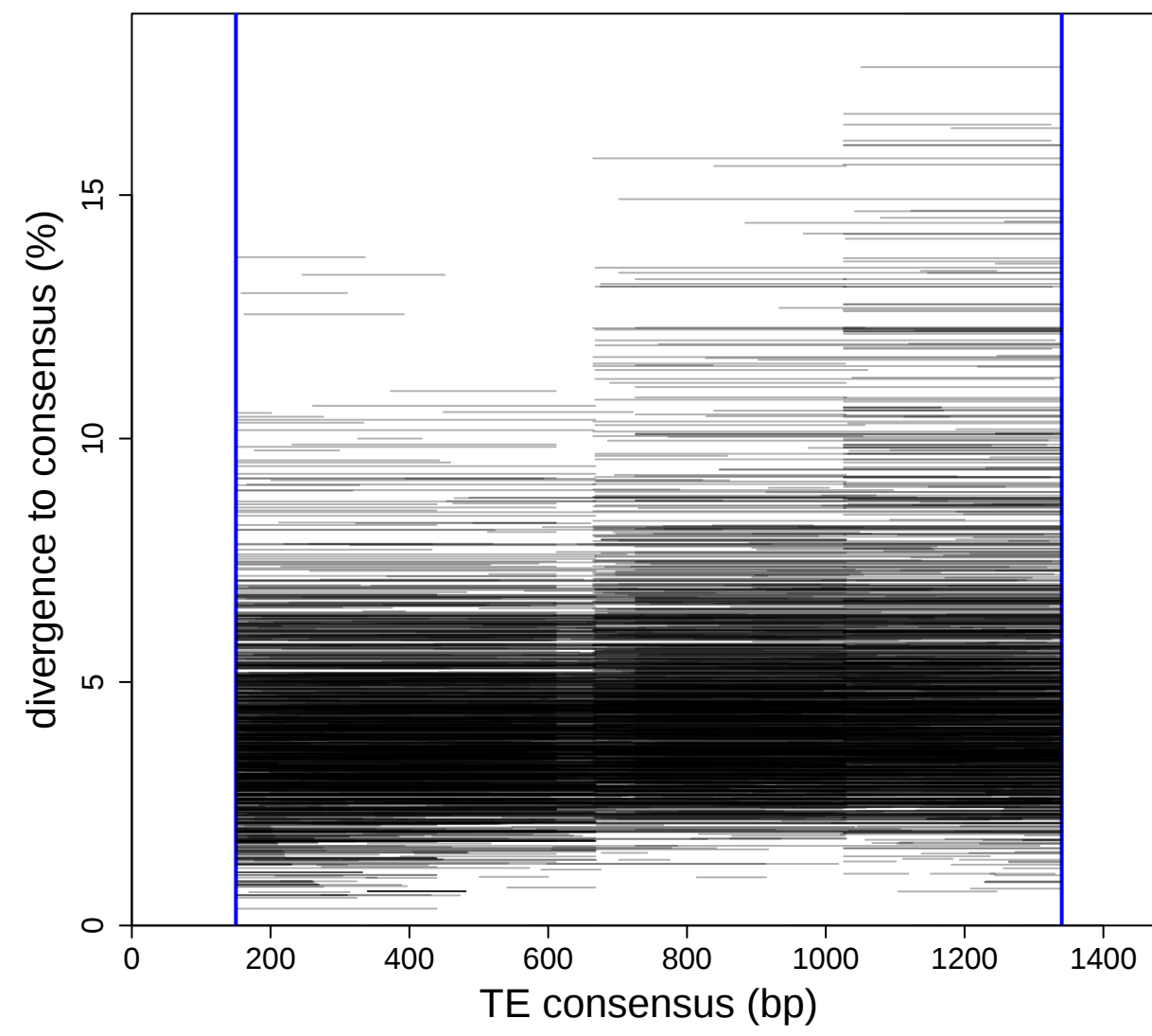
bed uf.bed g_1.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa ce
size: 1190bp; fragments: 2685; full length: 367 (>=1071bp)

After TETrimmer 1190 bp

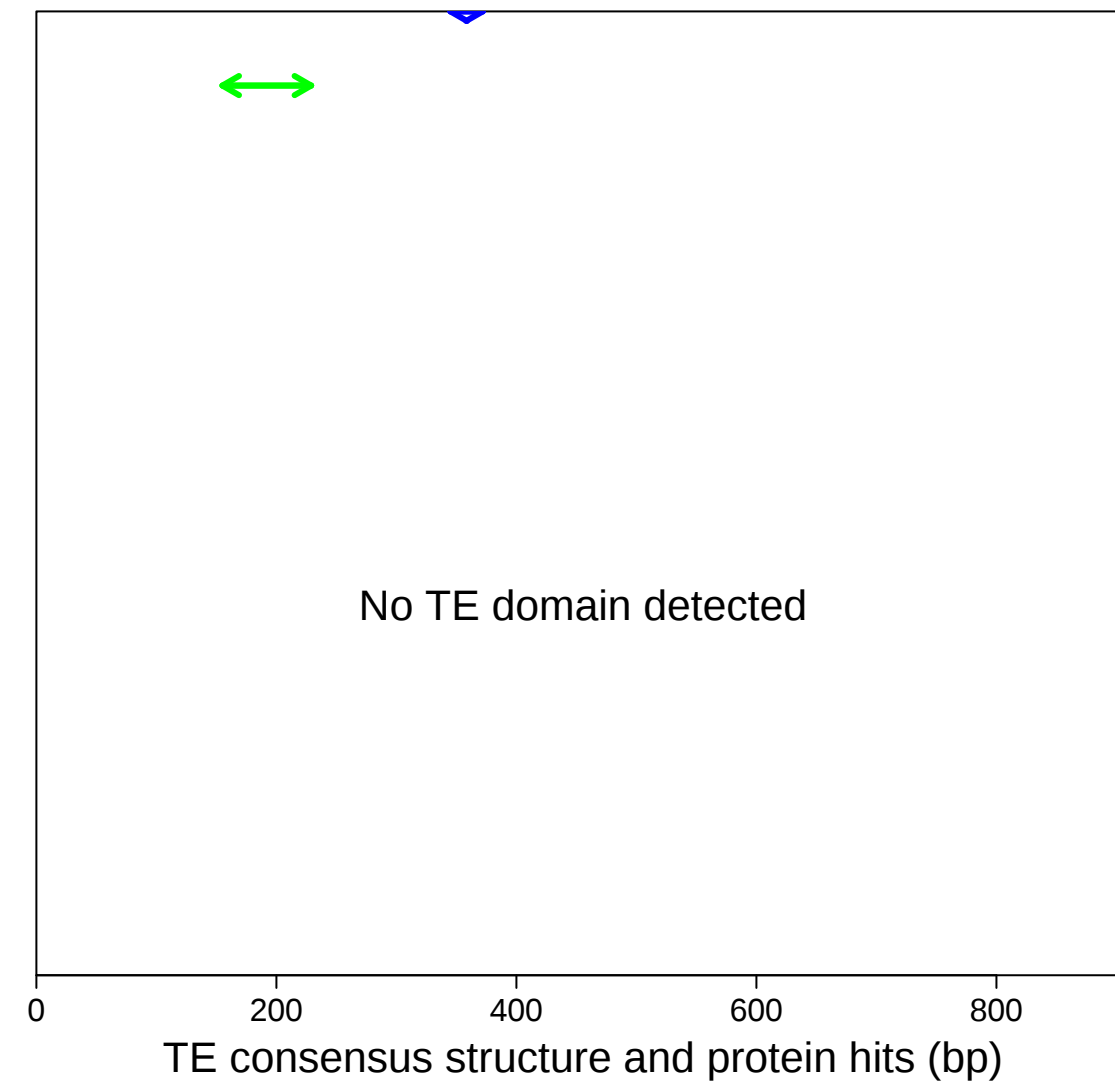
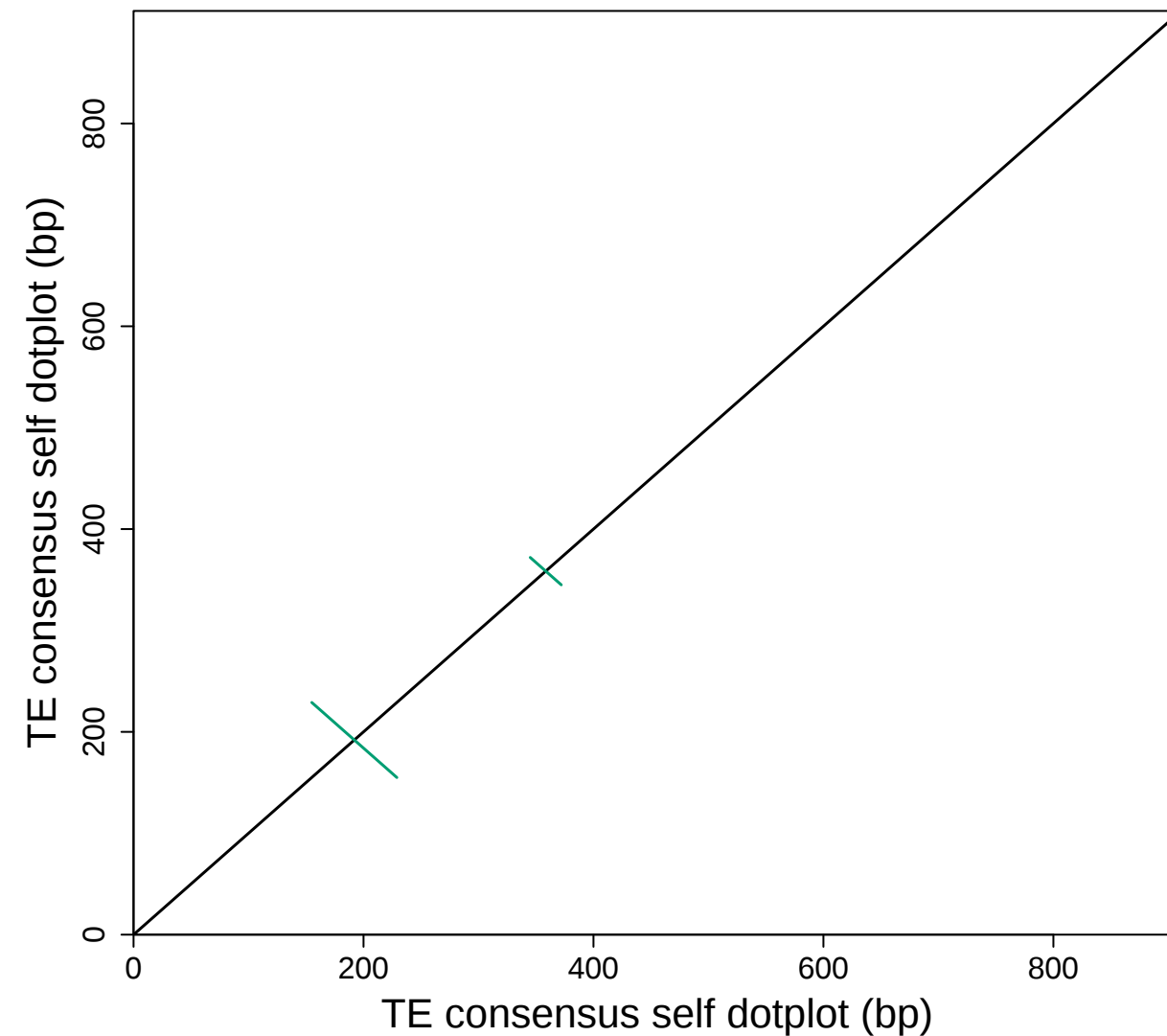
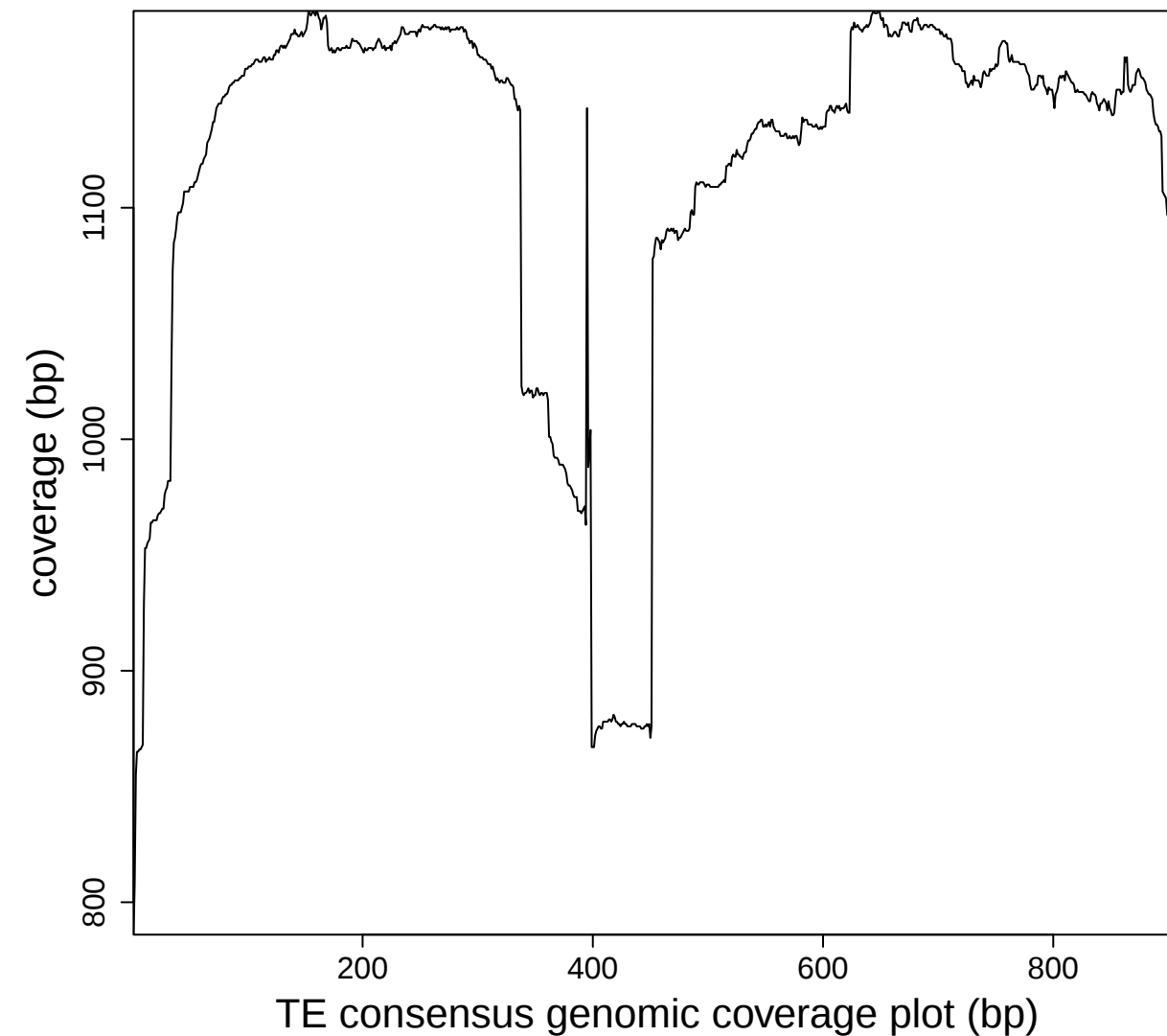
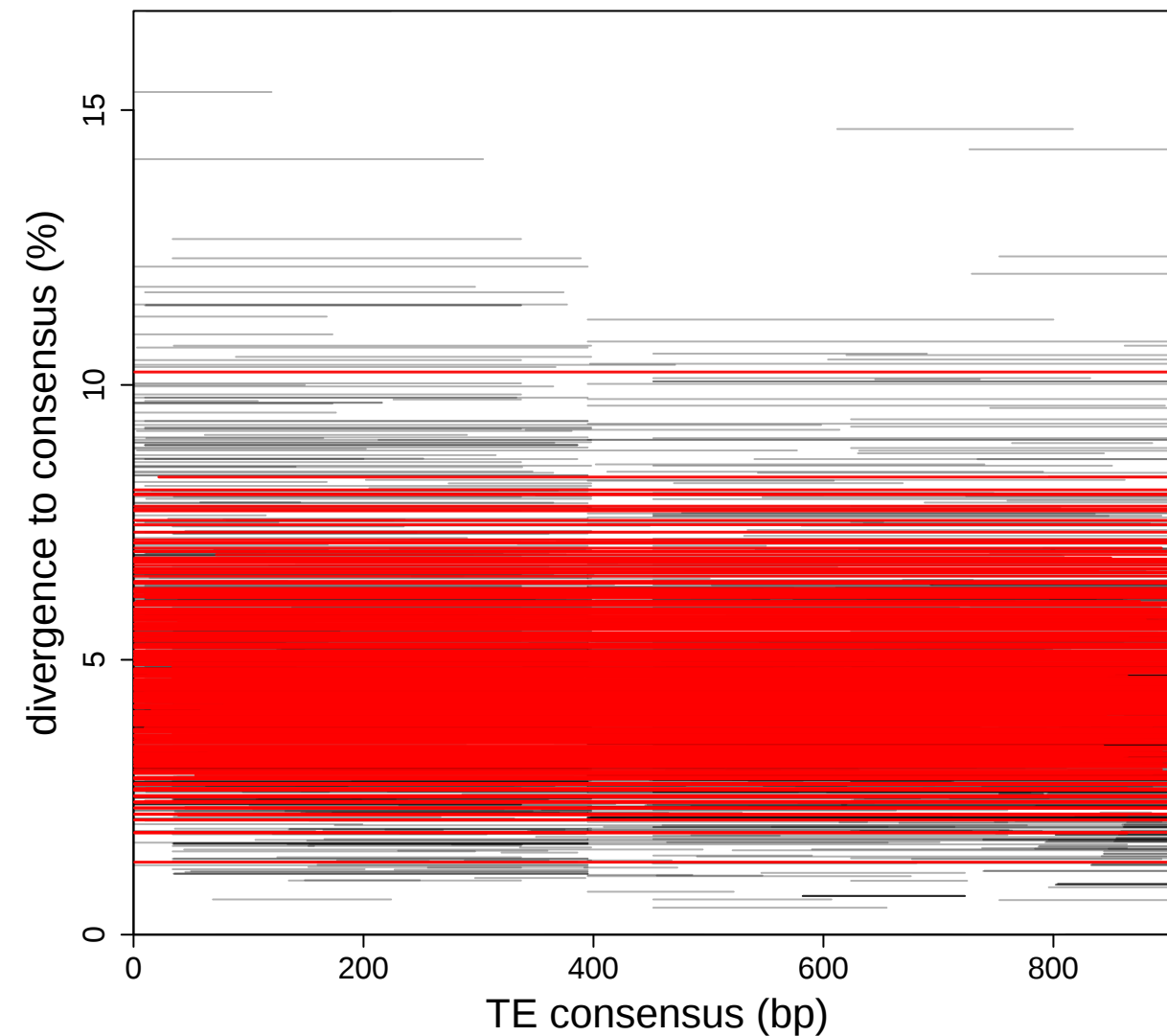


_428.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_c
size: 1490bp; fragments: 2684; full length: 0 (>=1341bp)

After TEtrimmer Extended plot Blue lines are boundaries

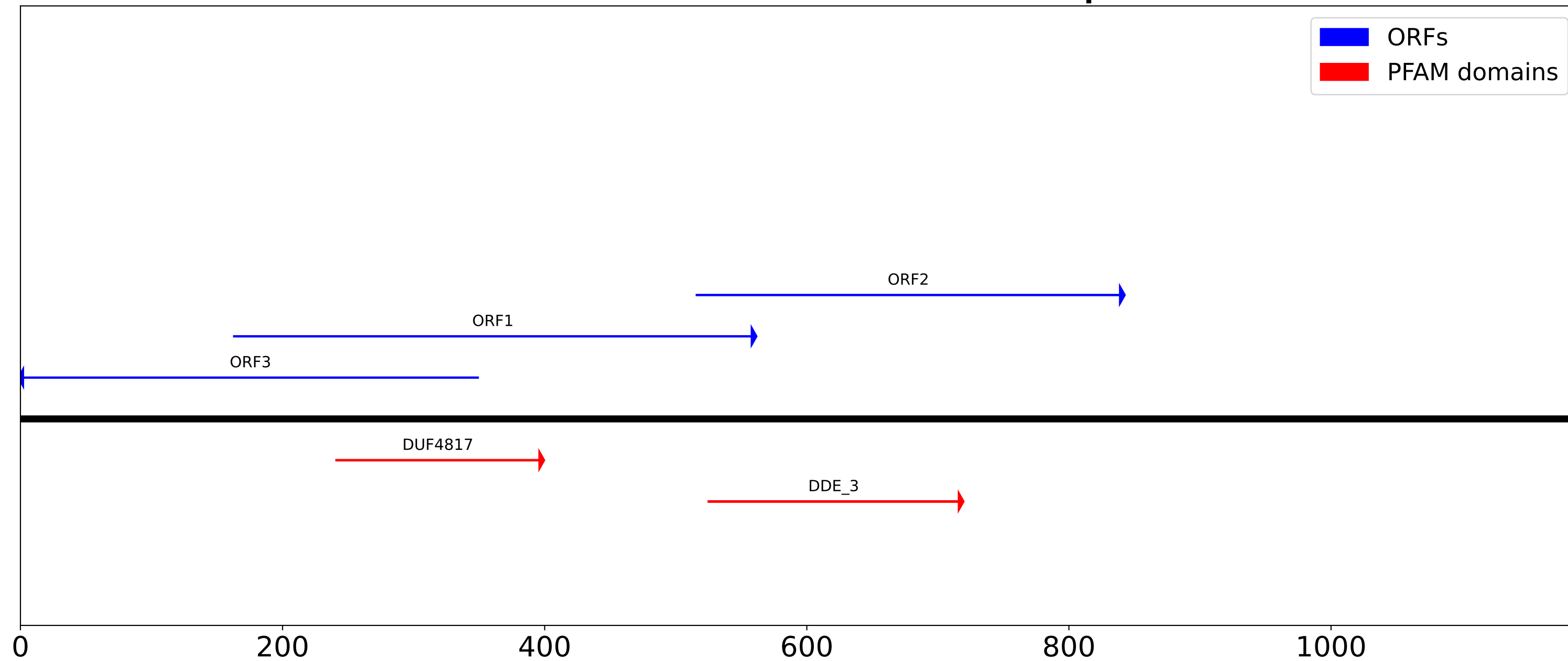


TE: rnd_1_family_428
size: 911bp; fragments: 2188; full length: 444 (≥ 819.9 bp)

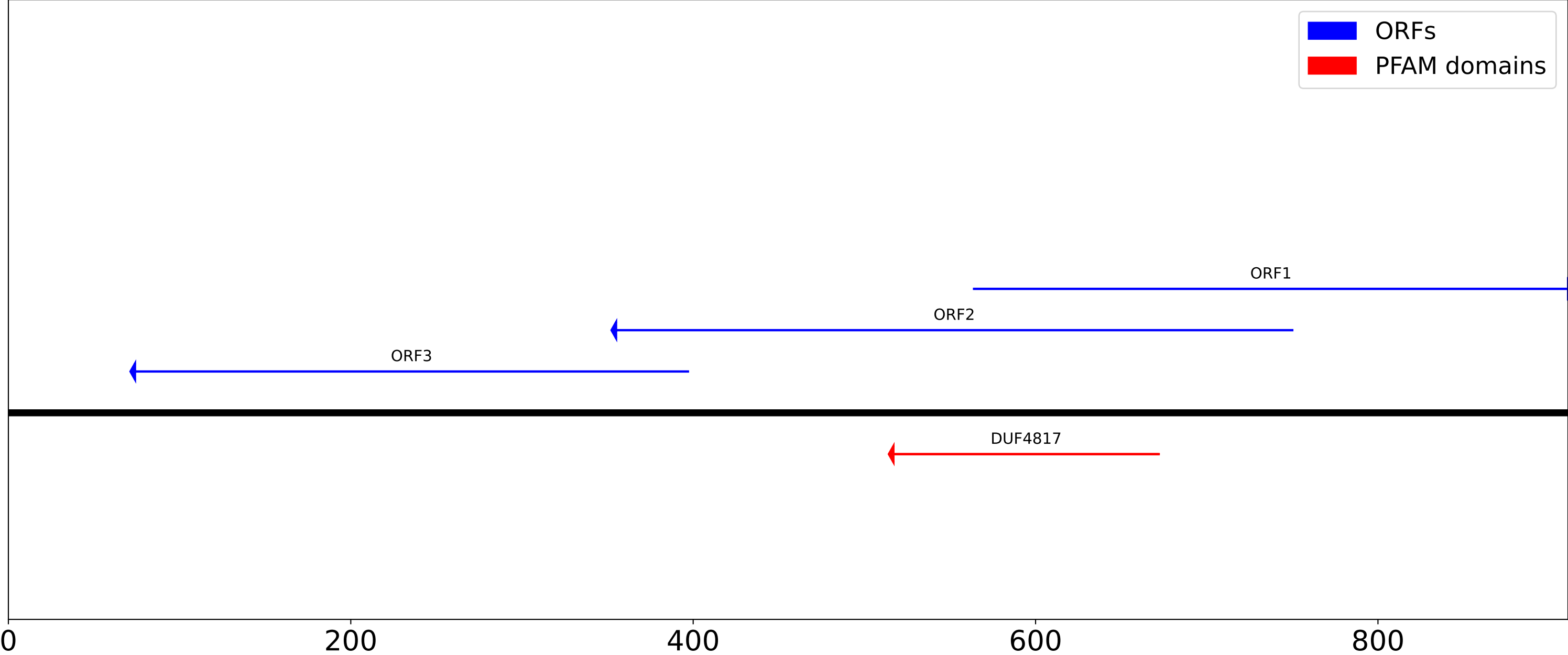


Before TEtrimmer 911 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

