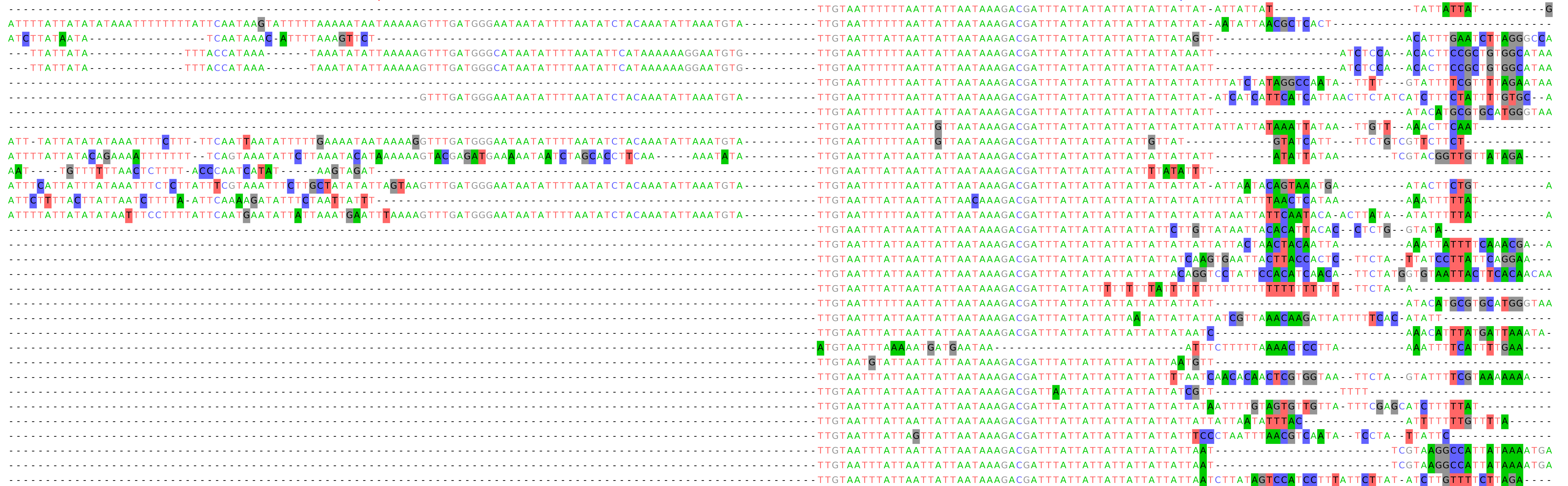
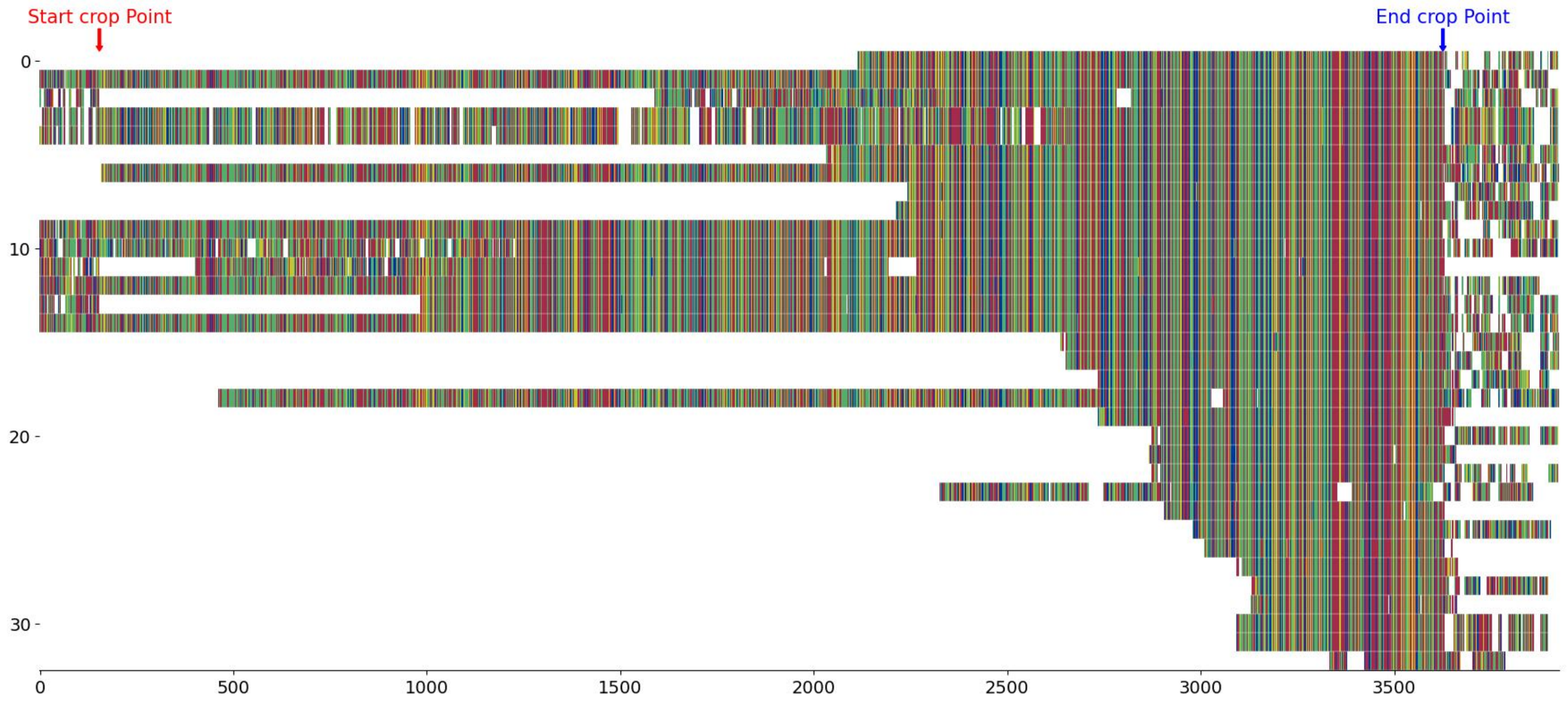


TTCT



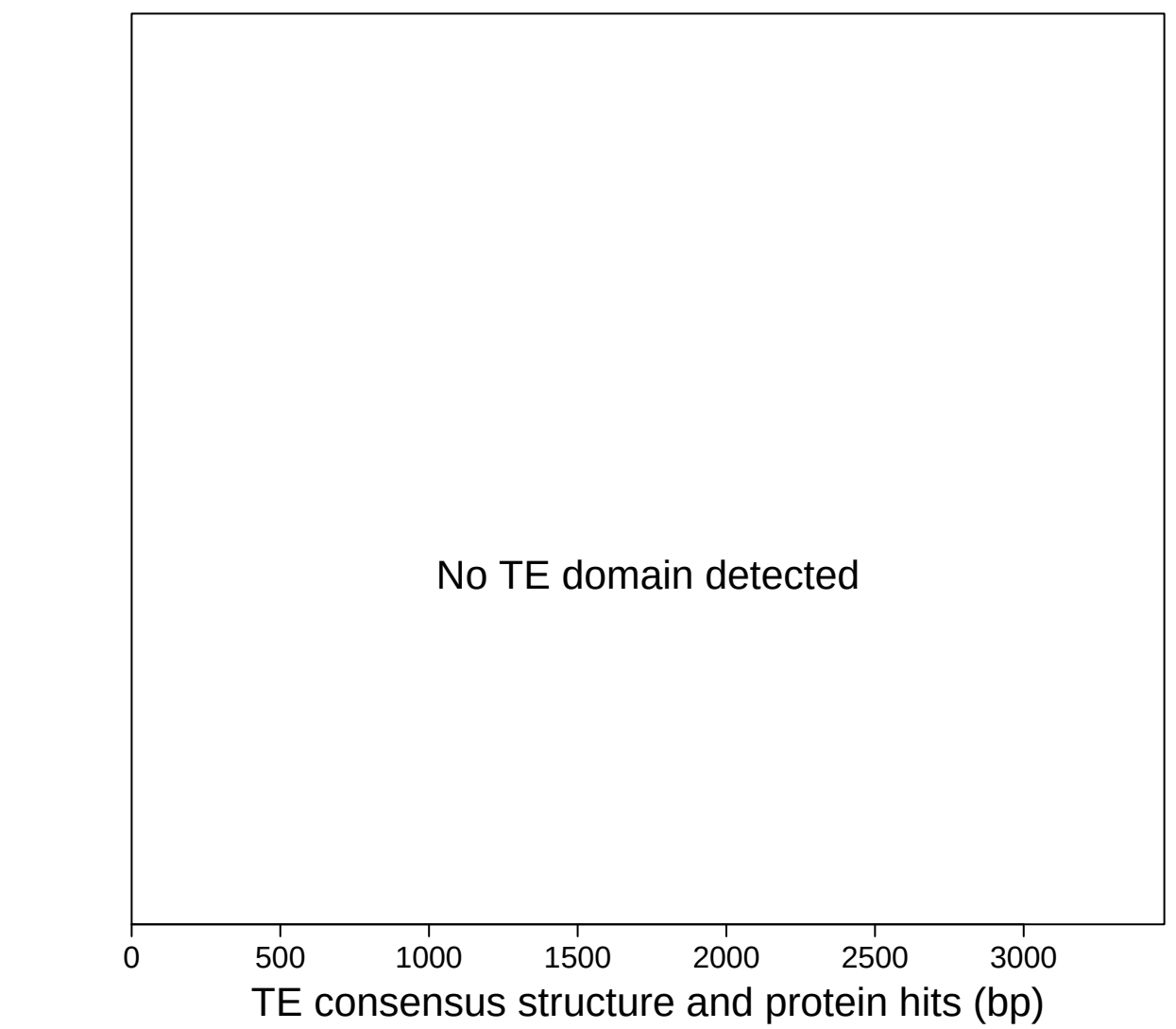
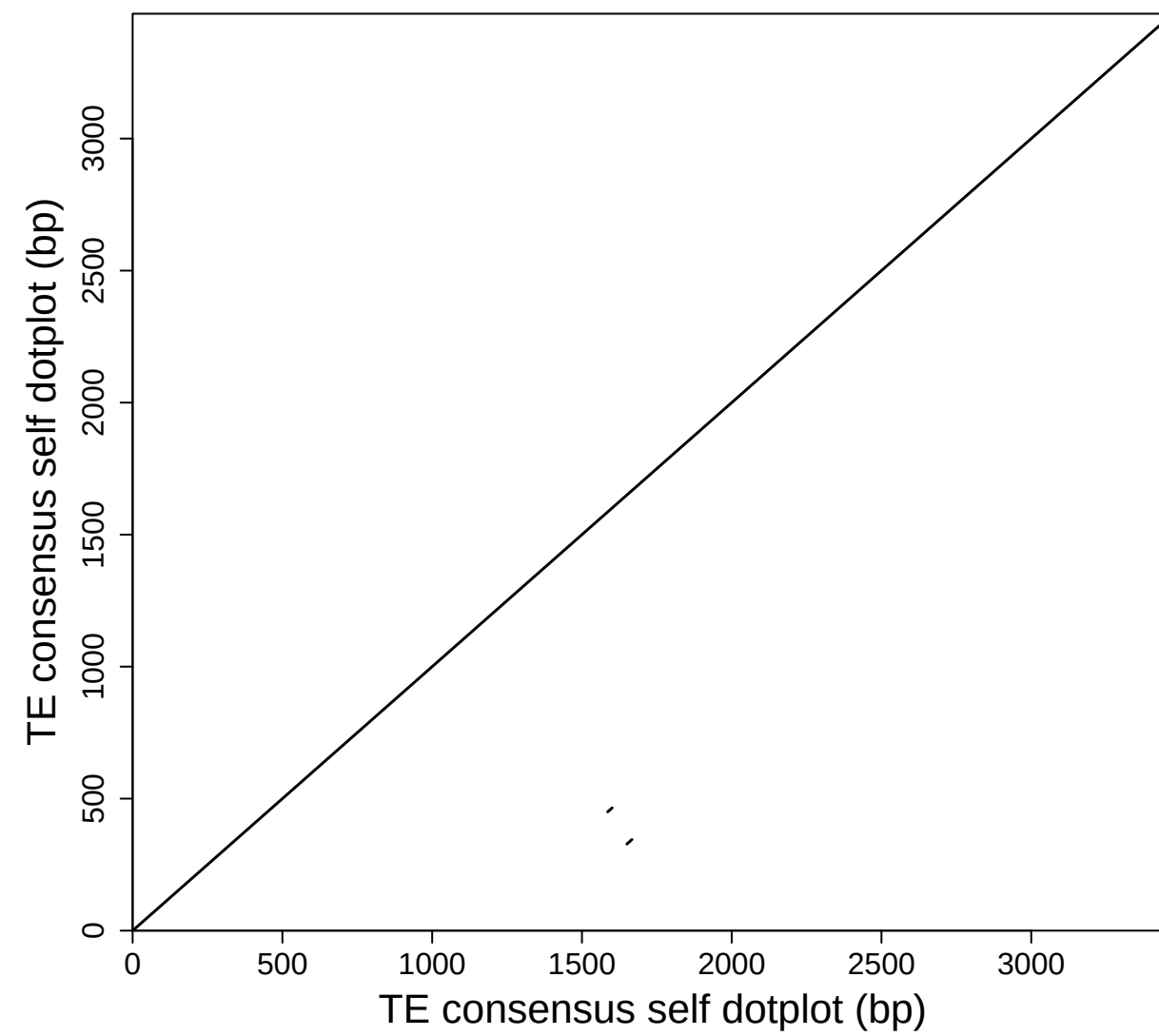
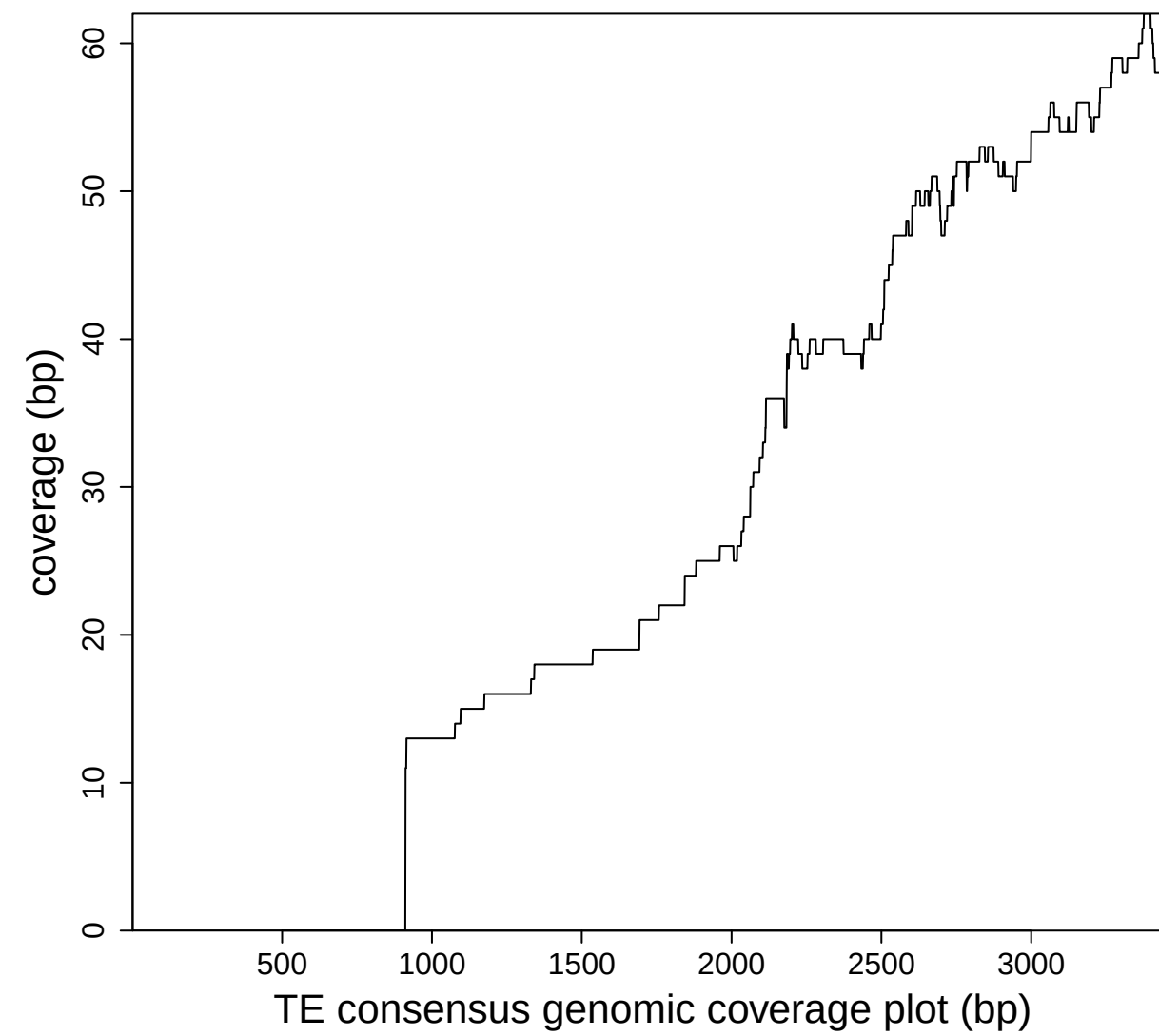
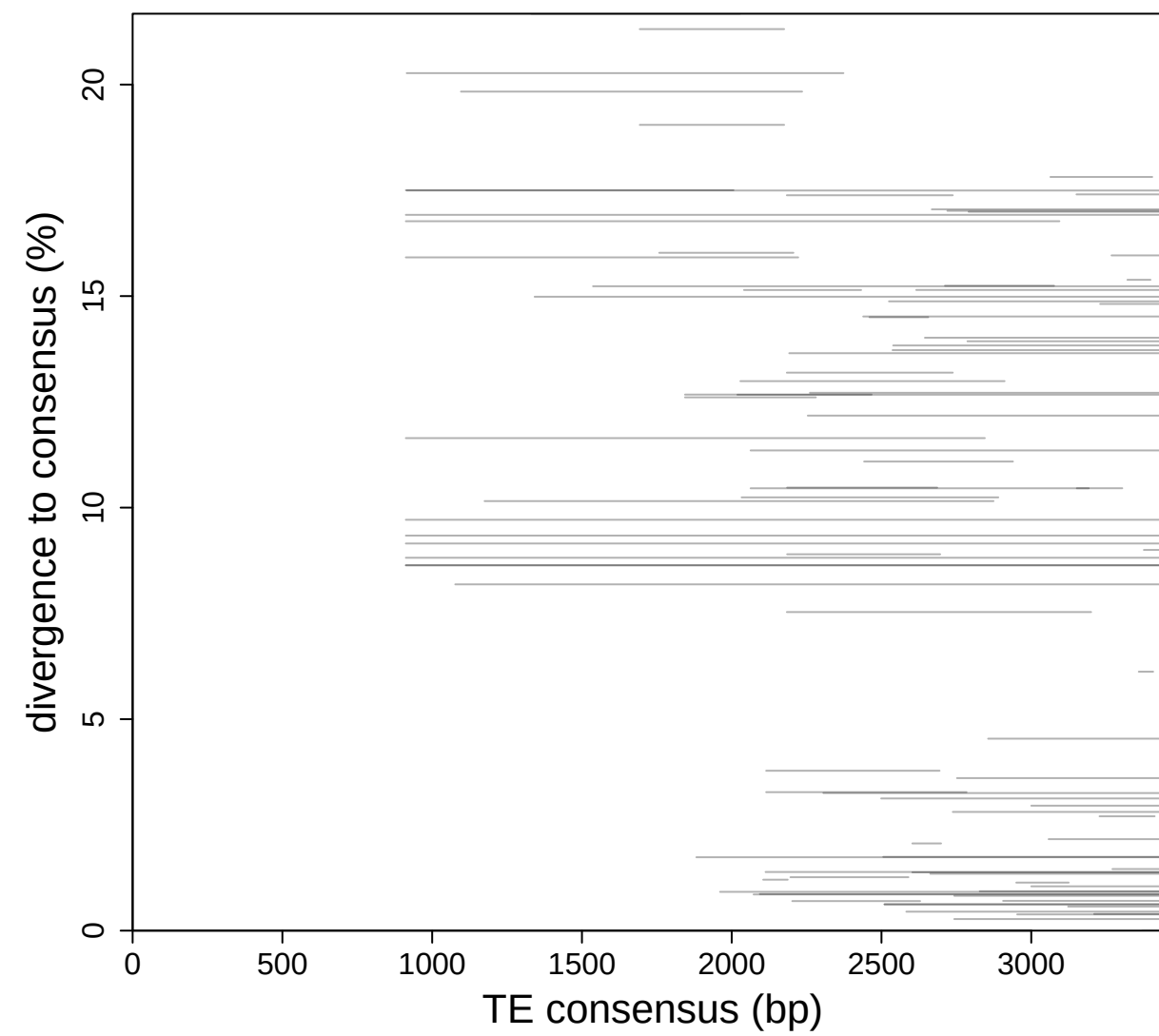
Start crop Point

End crop Point



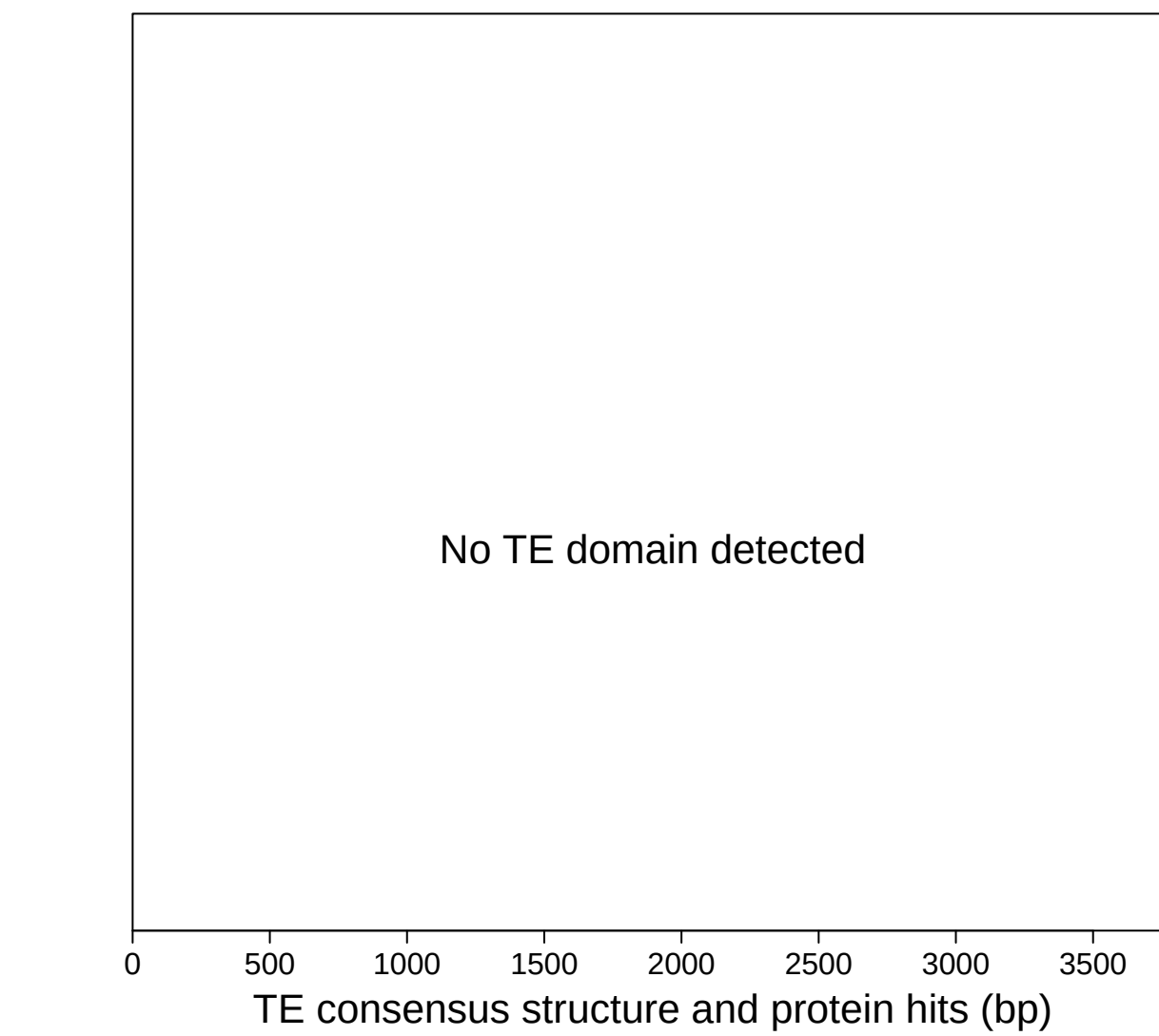
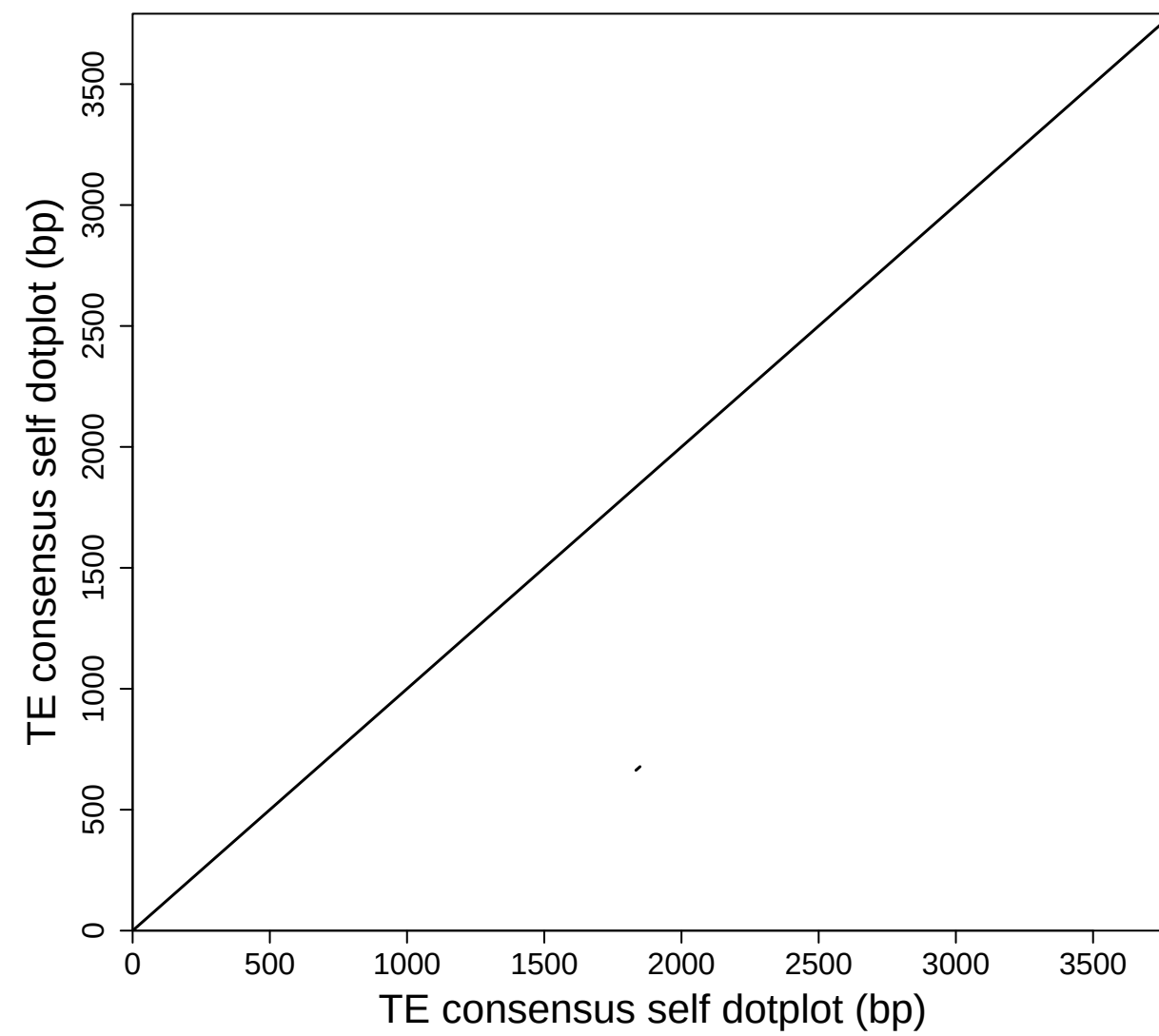
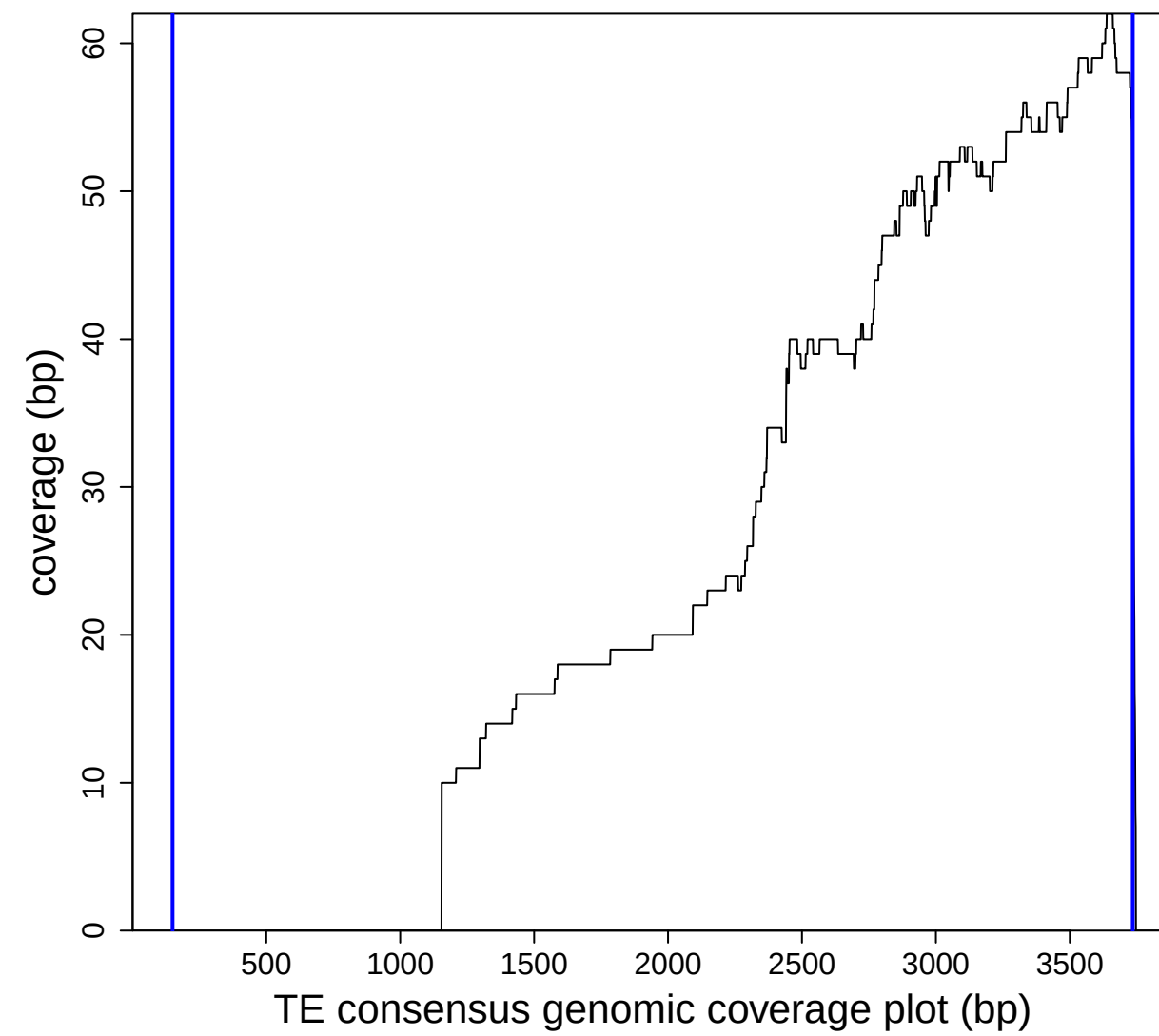
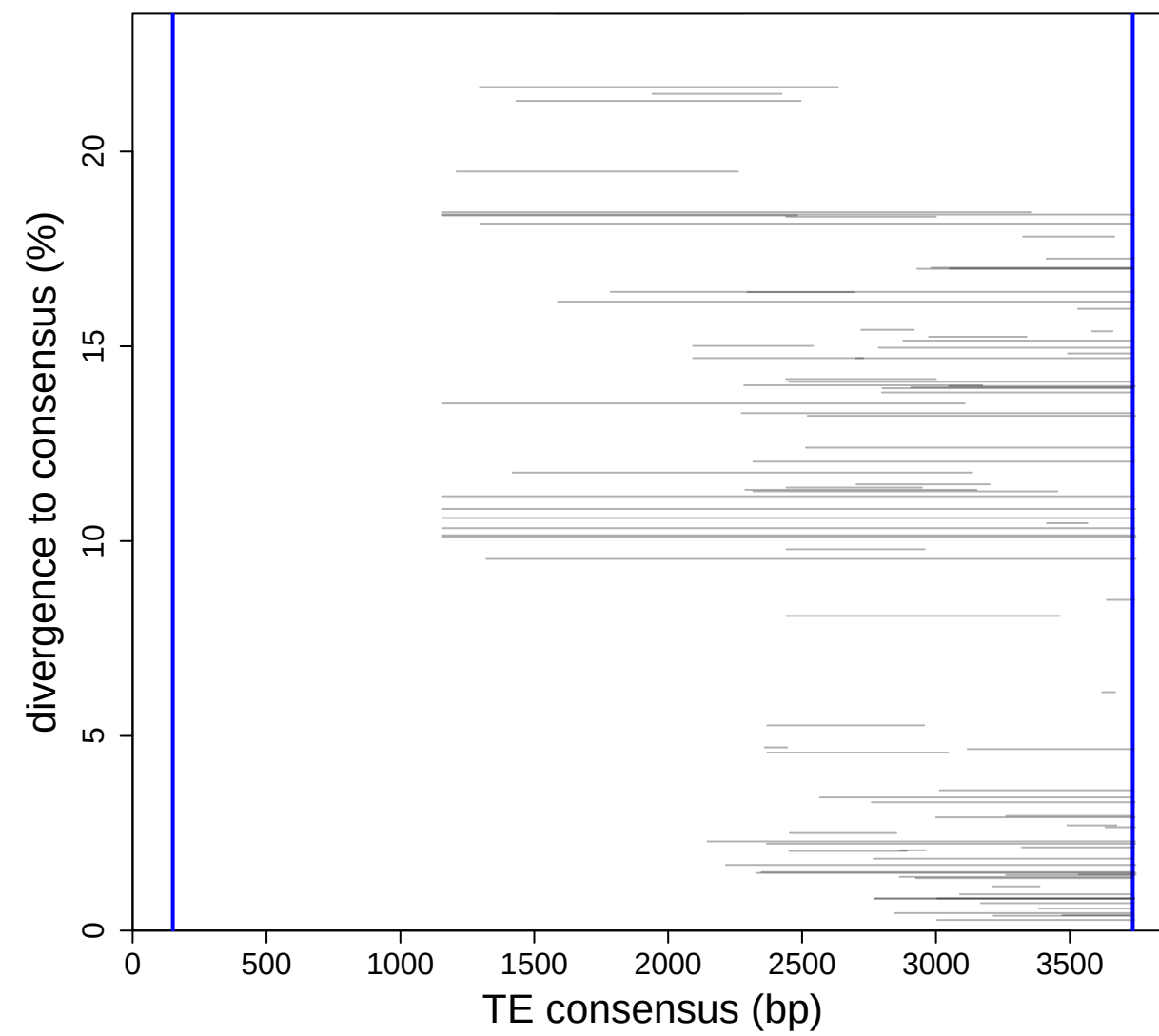
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 3473bp; fragments: 96; full length: 0 (>=3125.7bp)

After TEtrimmer 3473 bp

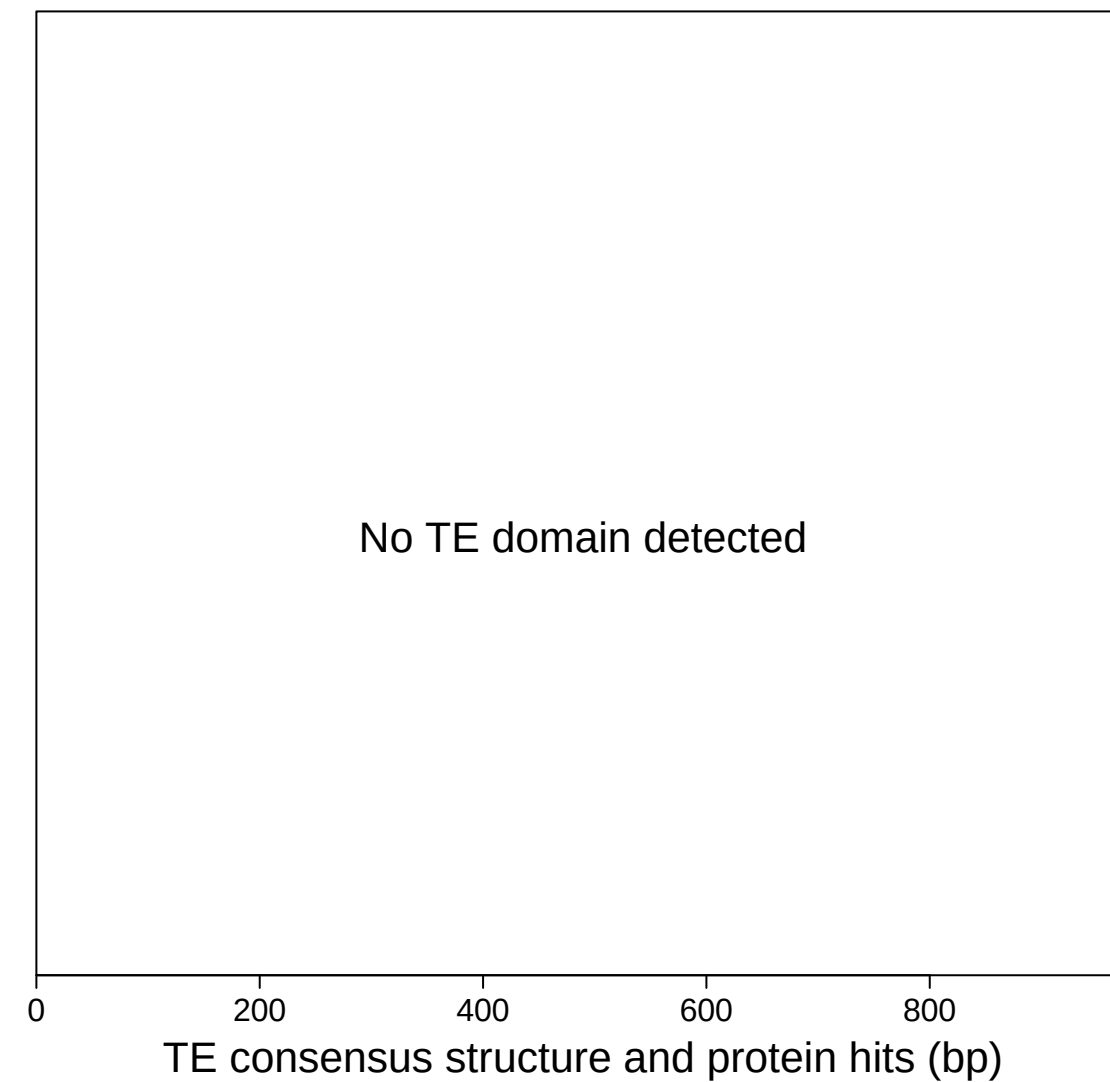
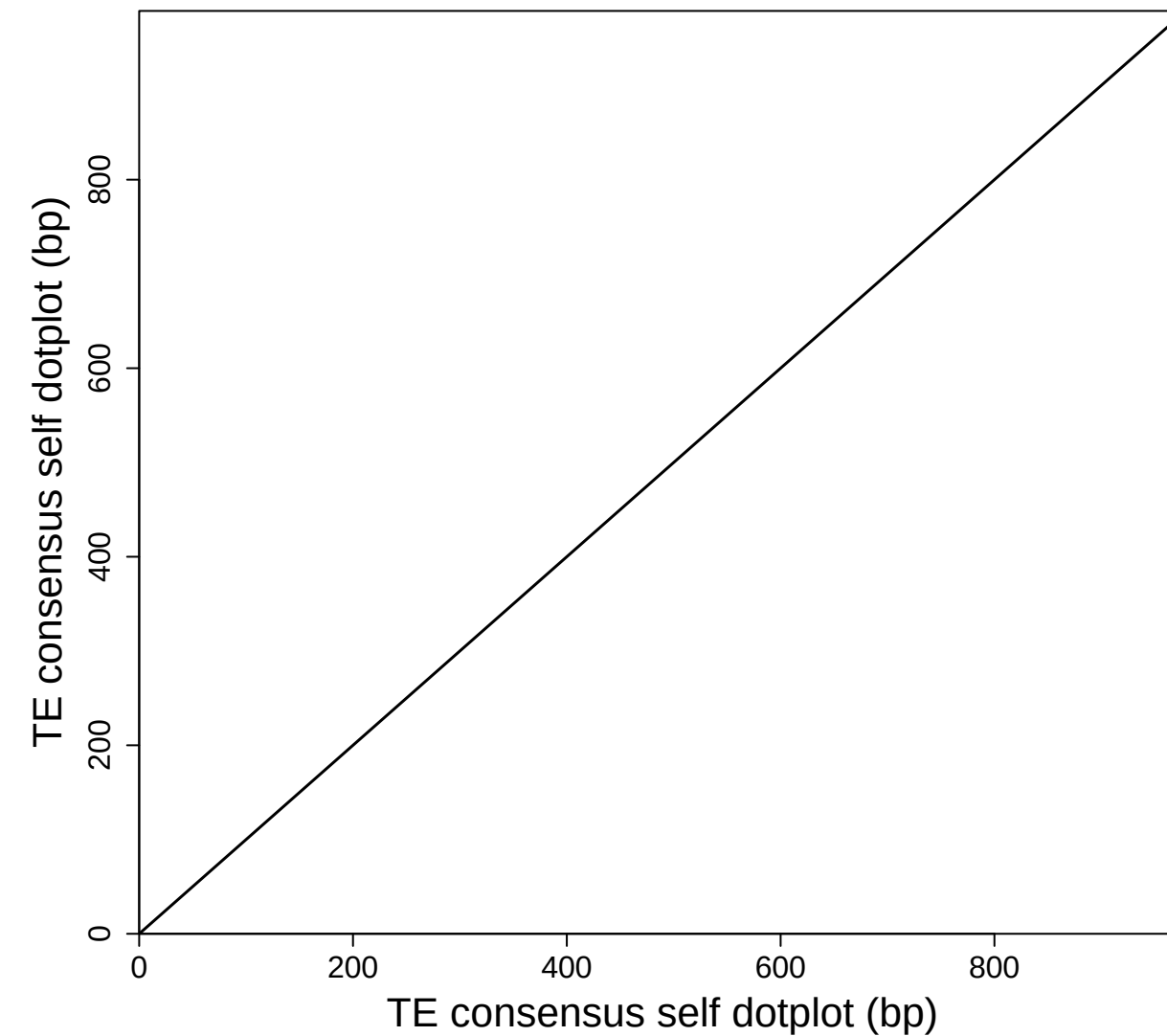
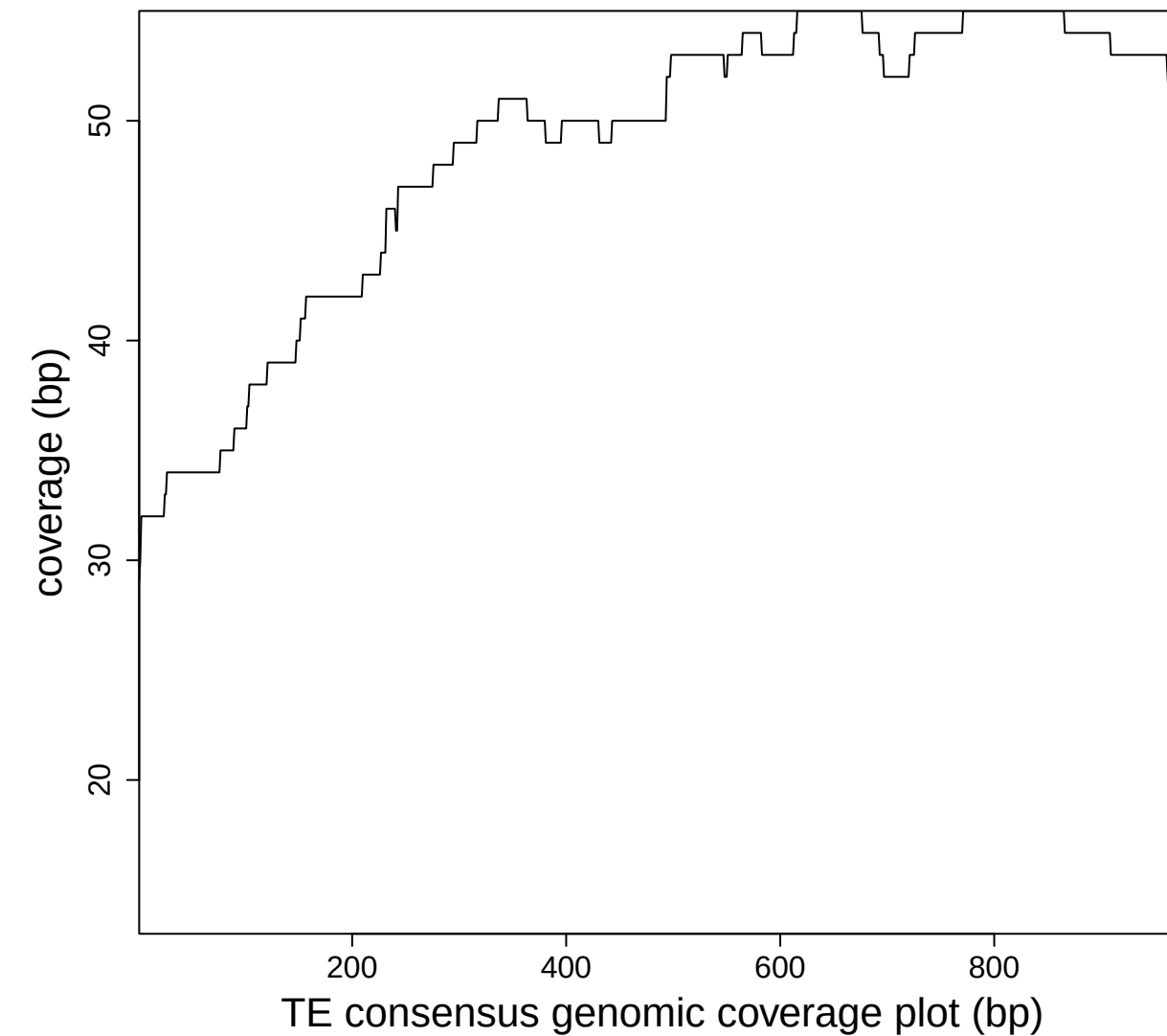
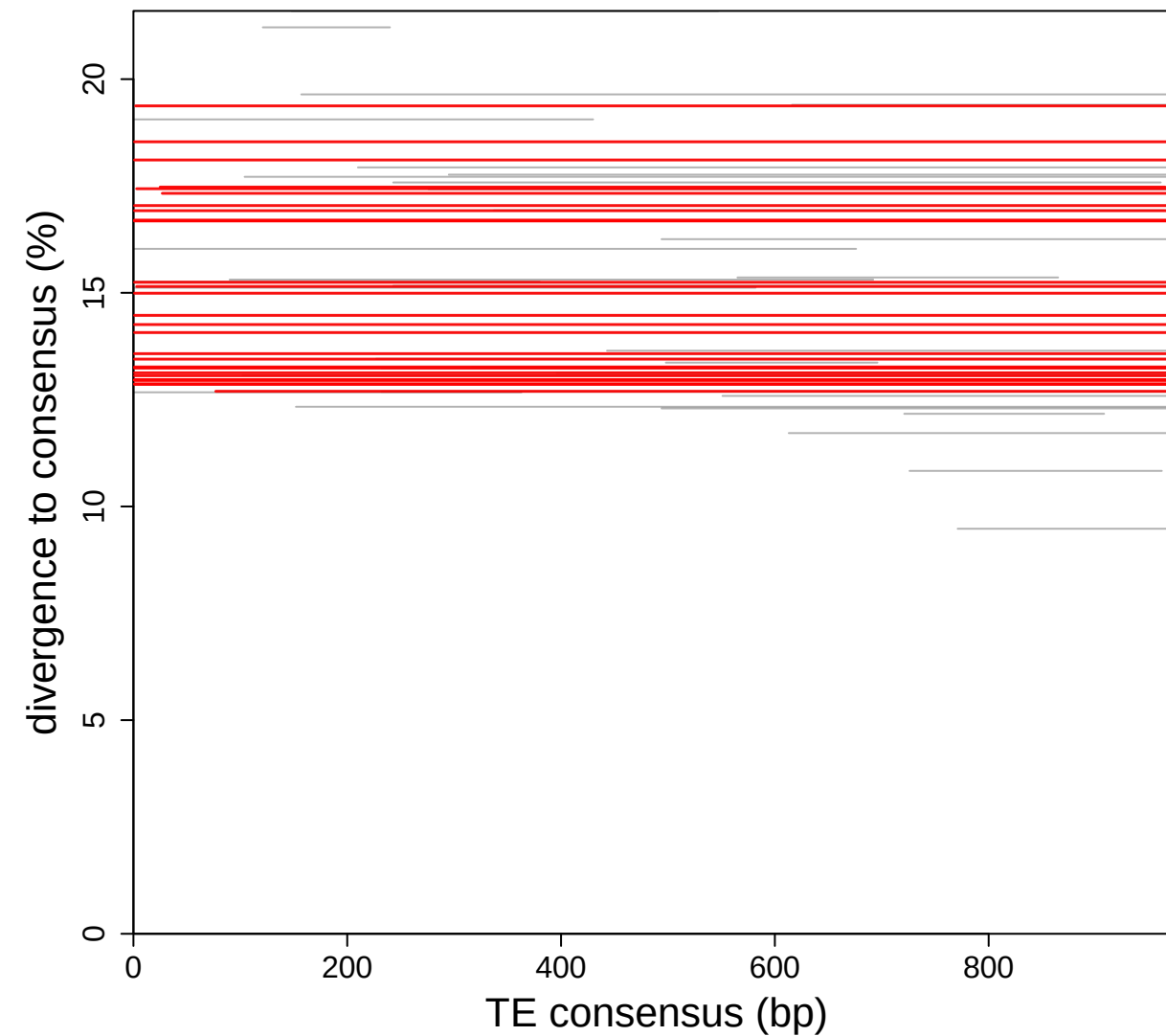


9.fasta.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcIn.fa aln.fa cl.f
size: 3885bp; fragments: 94; full length: 0 (>=3496.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

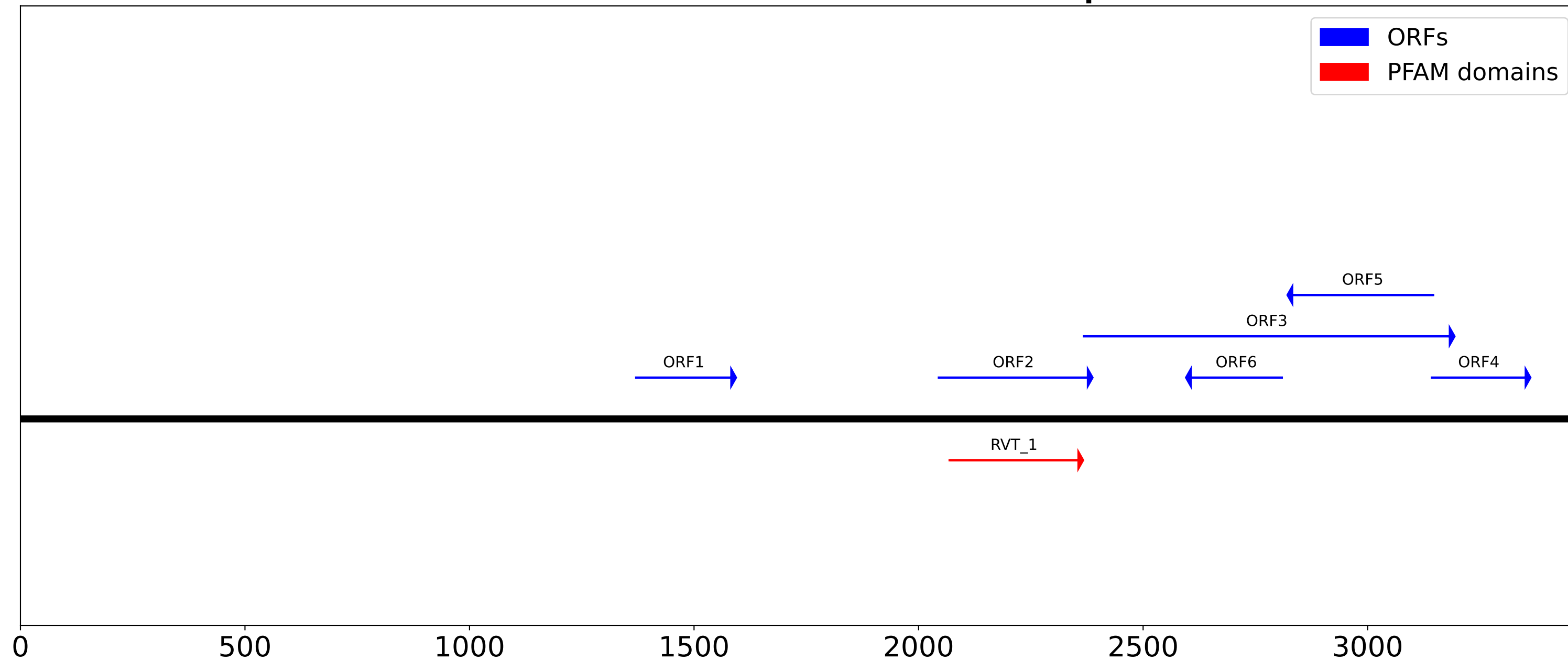


TE: rnd_4_family_2479
size: 979bp; fragments: 64; full length: 30 (≥ 881.1 bp)

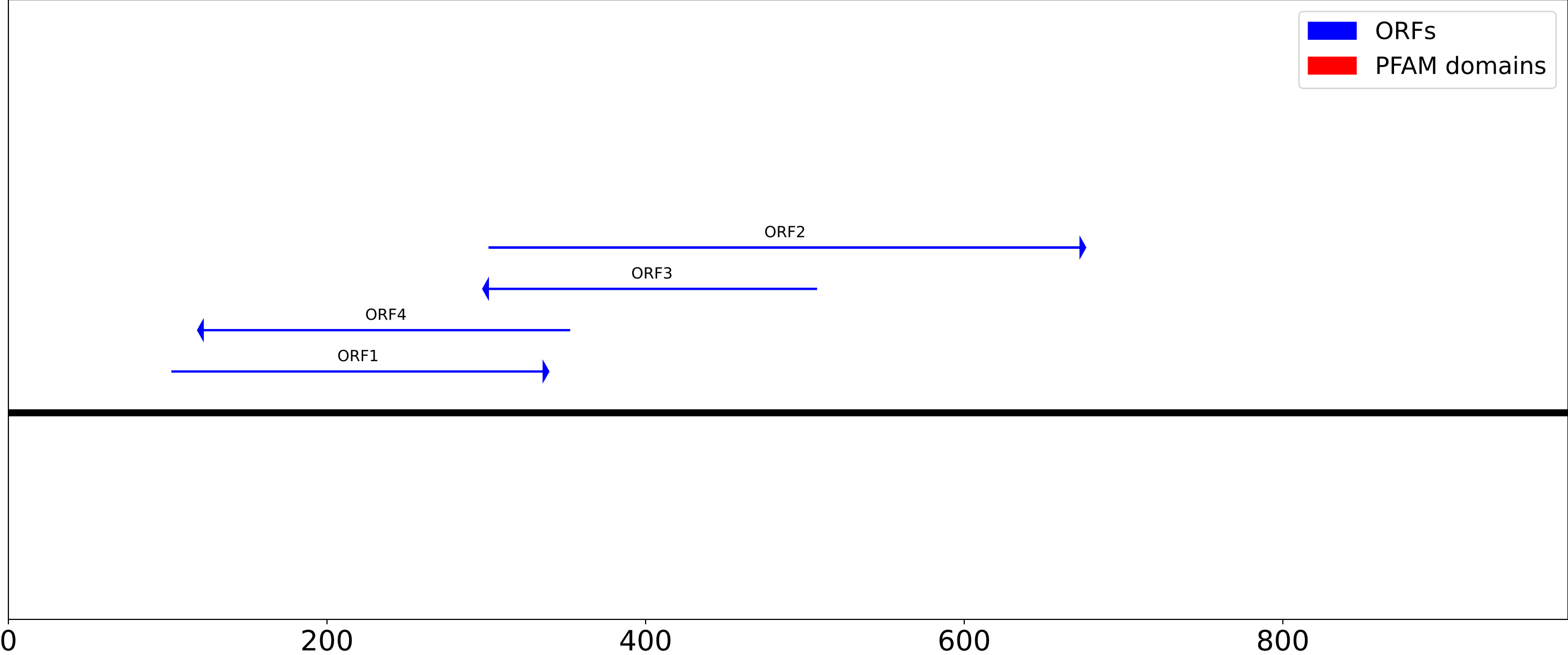


Before TEtrimmer 979 bp

After TEtrimmer ORF and PFAM domain plot



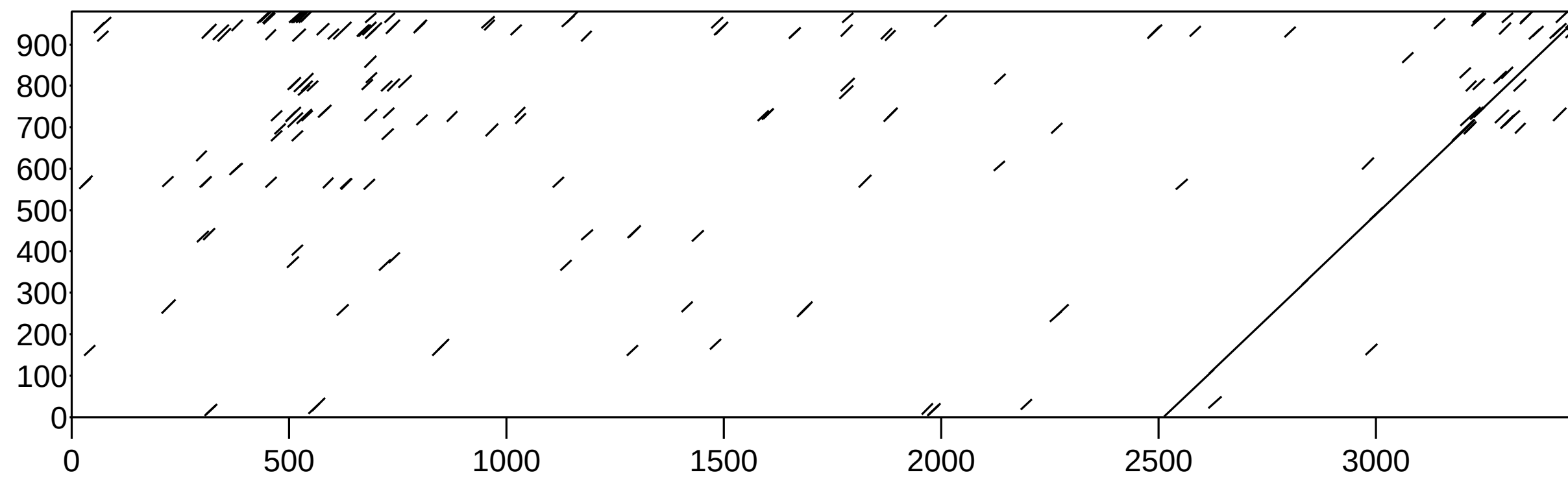
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)