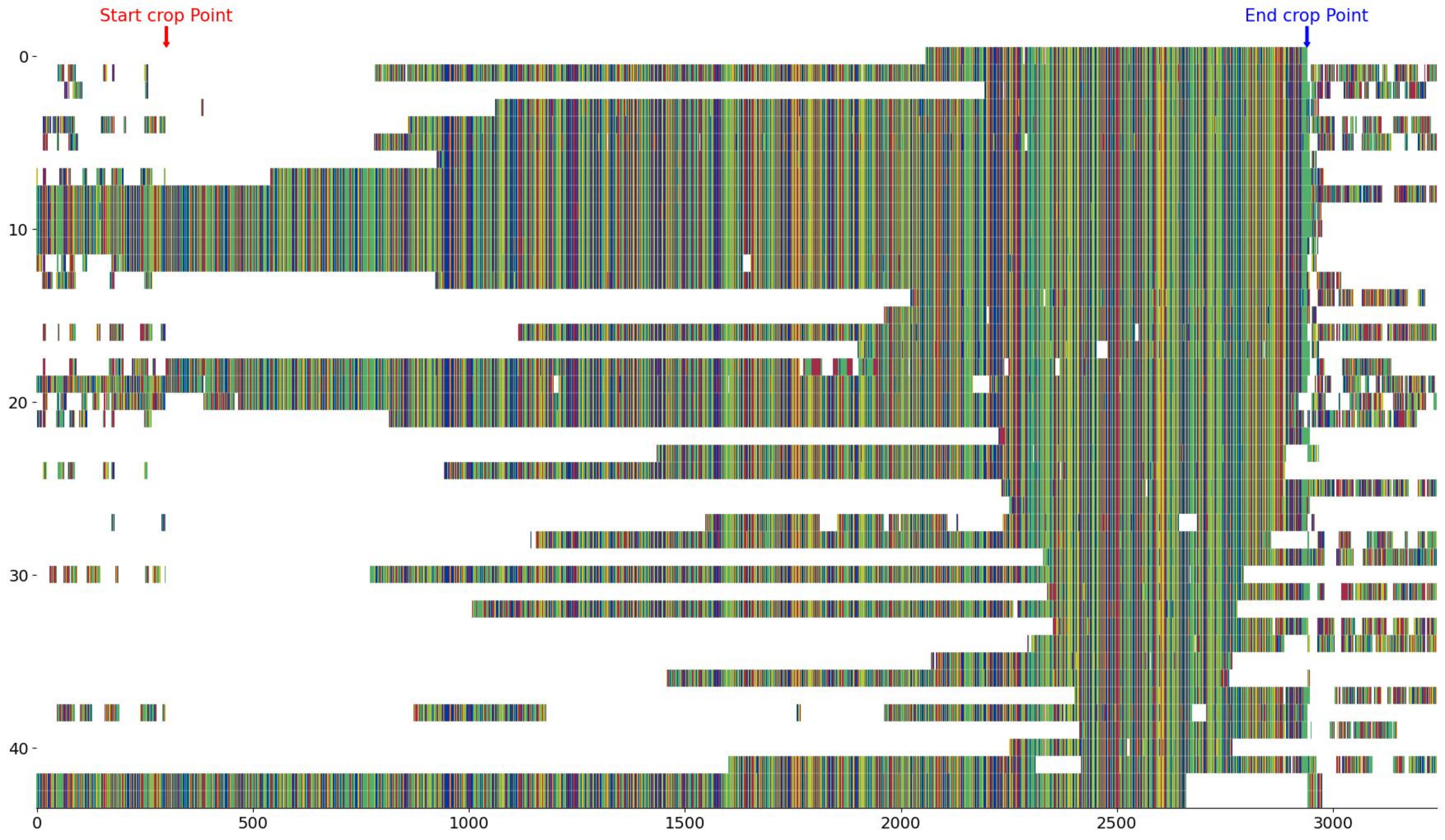
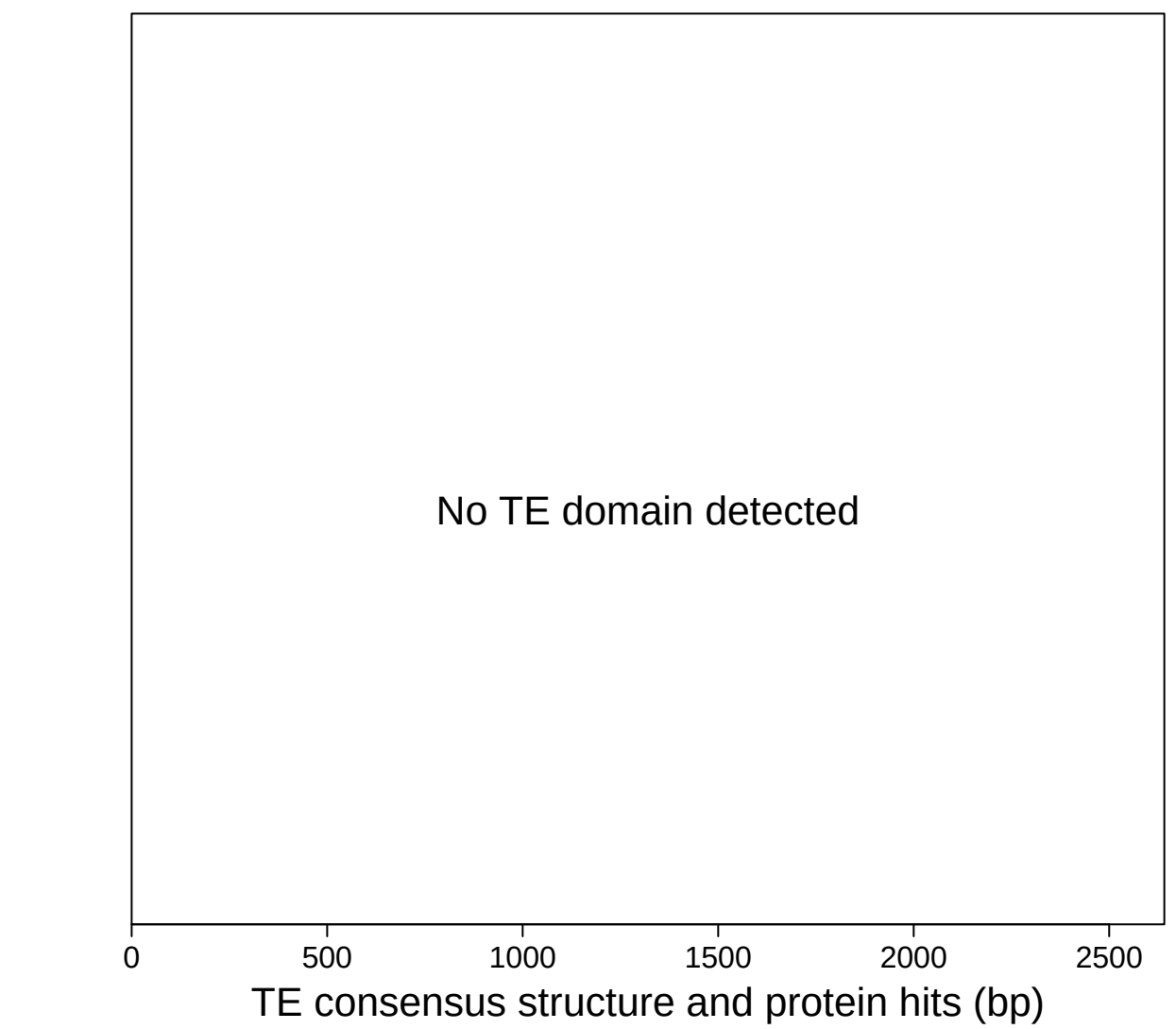
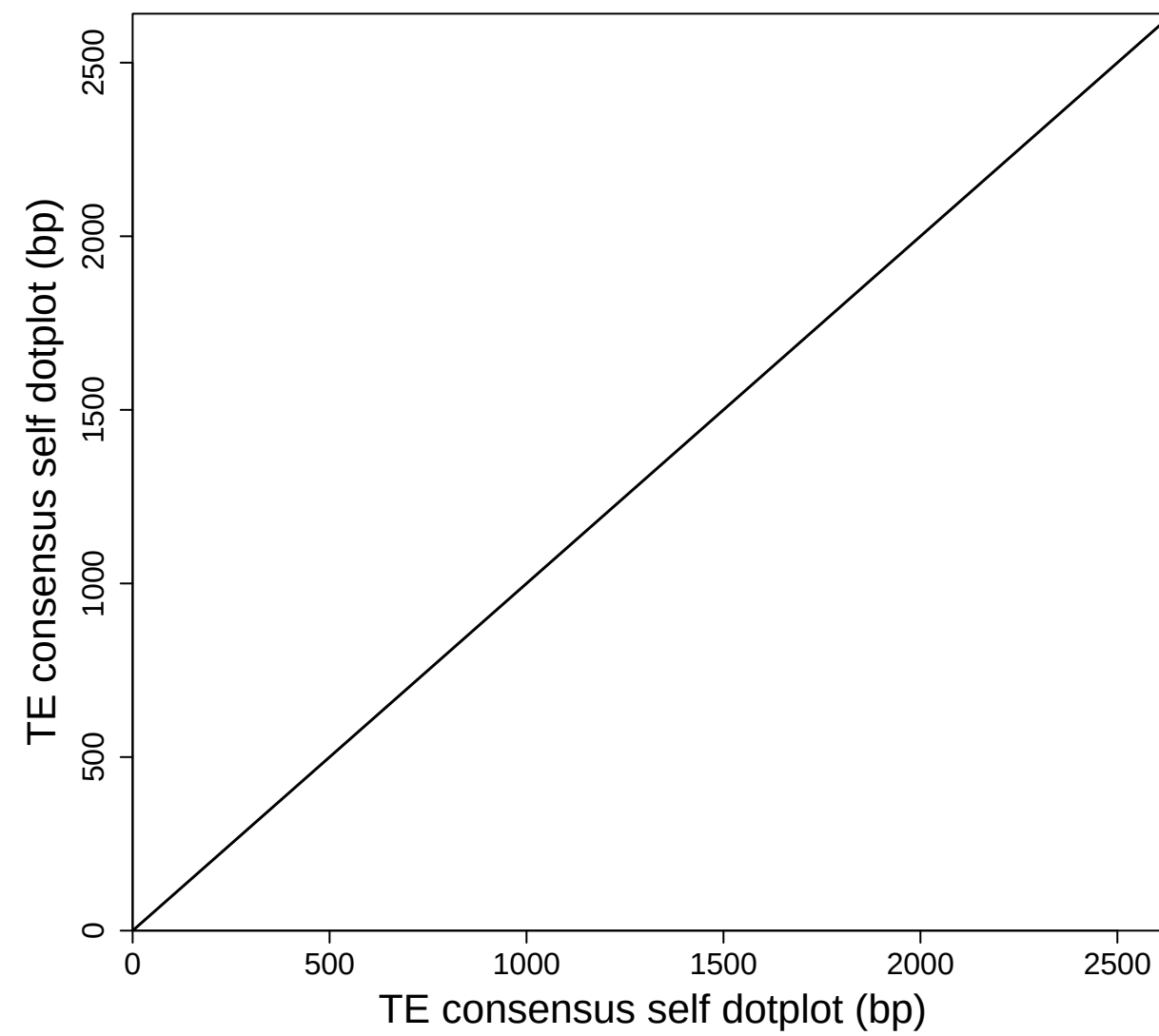
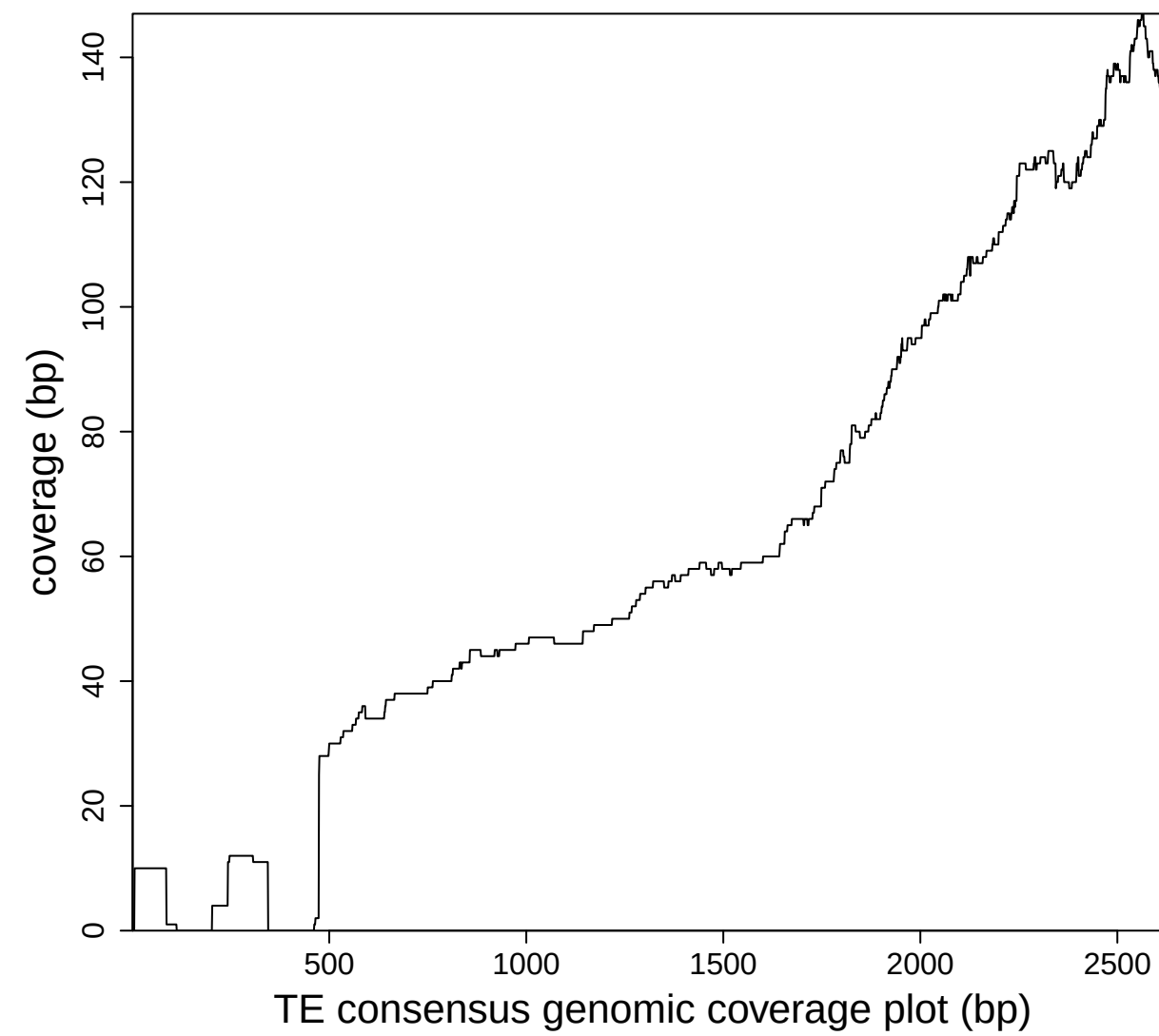
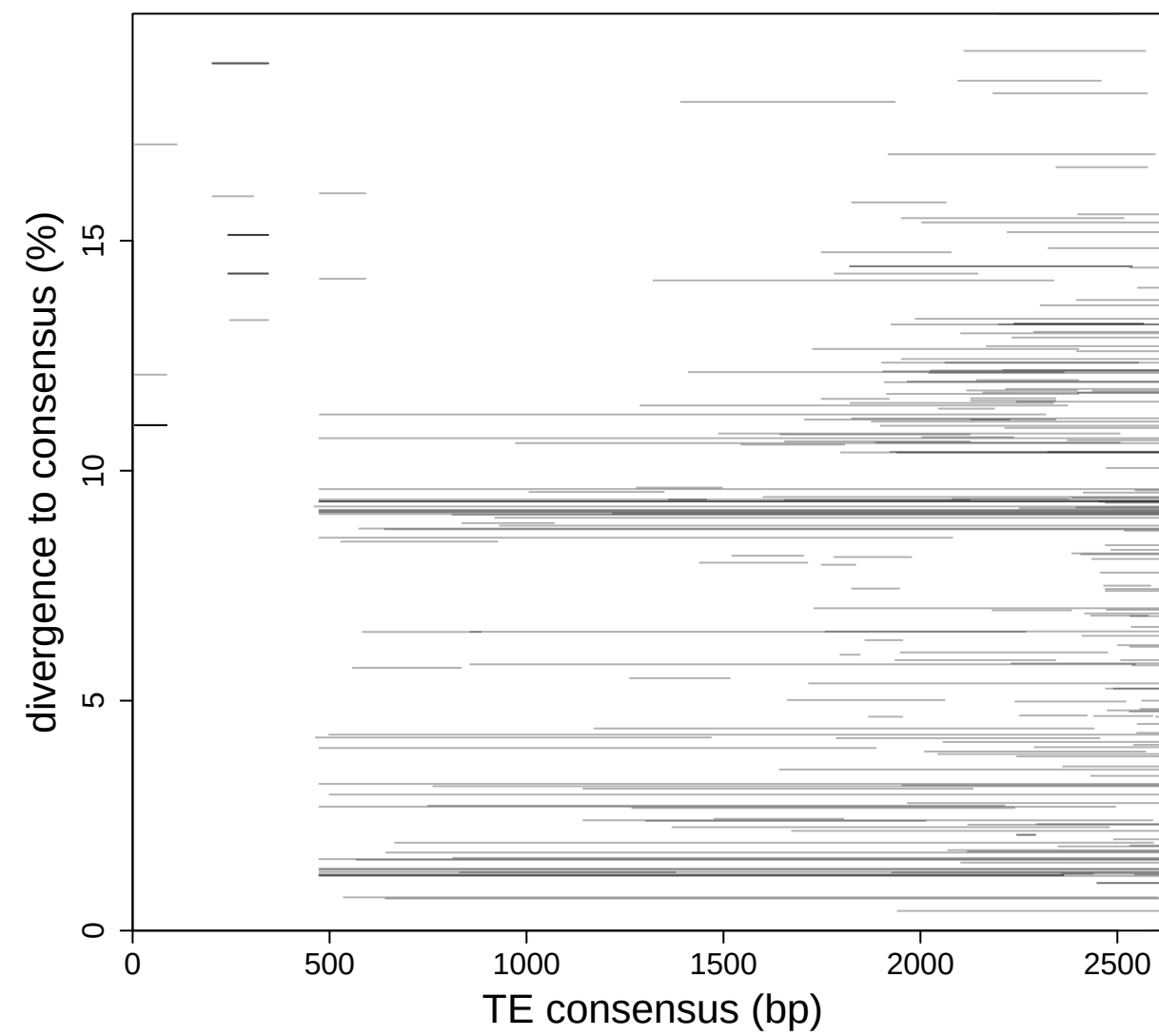






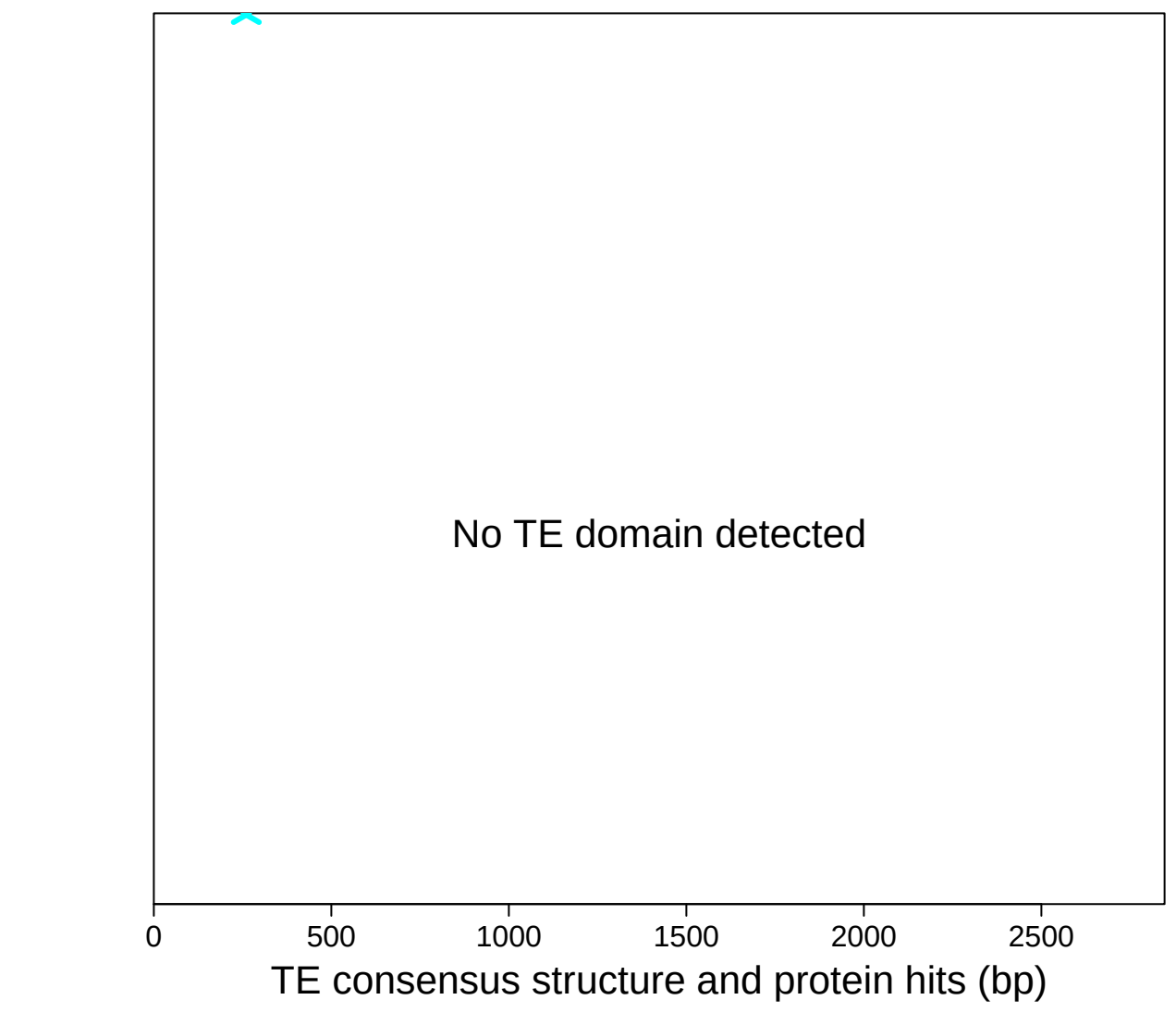
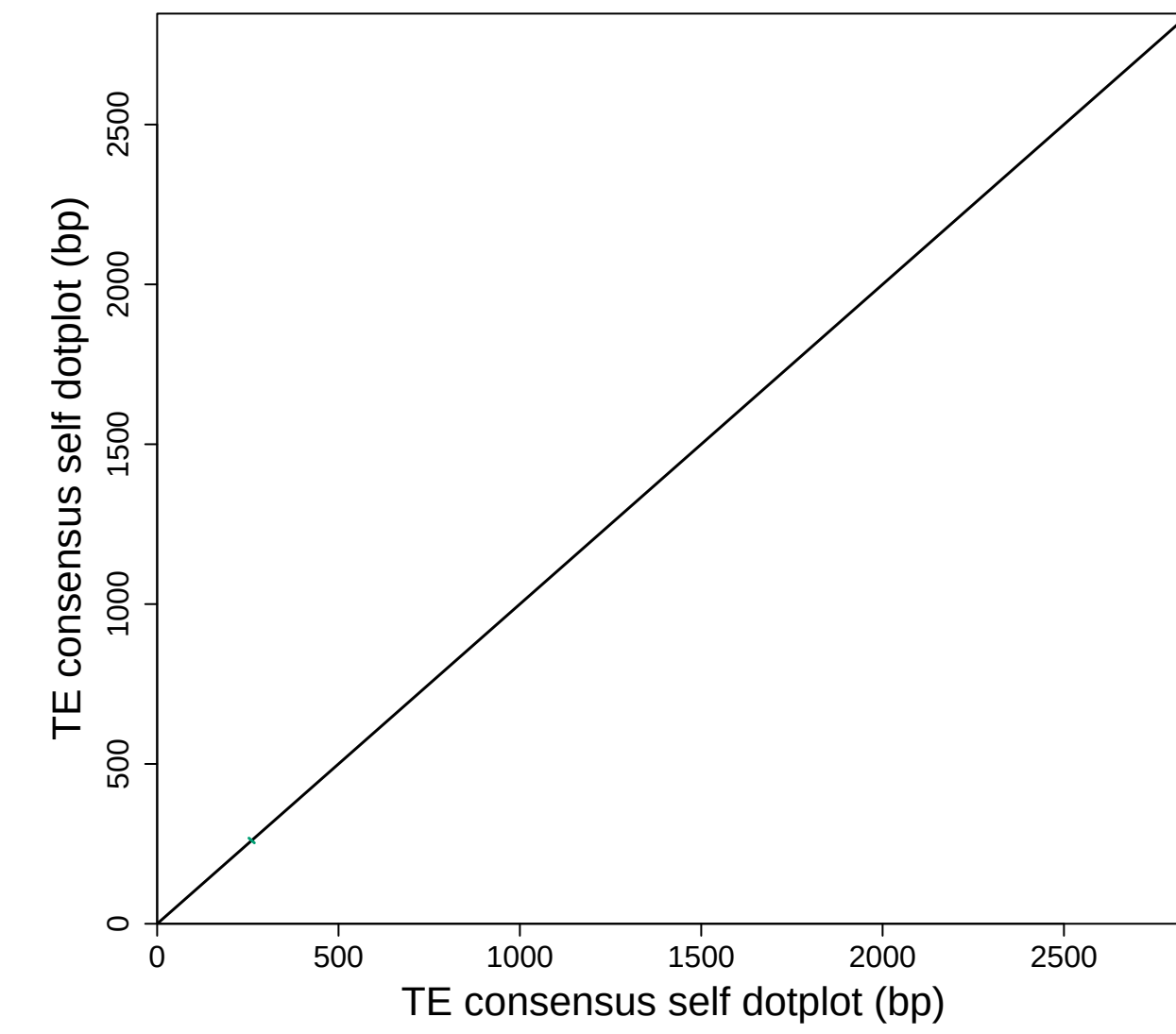
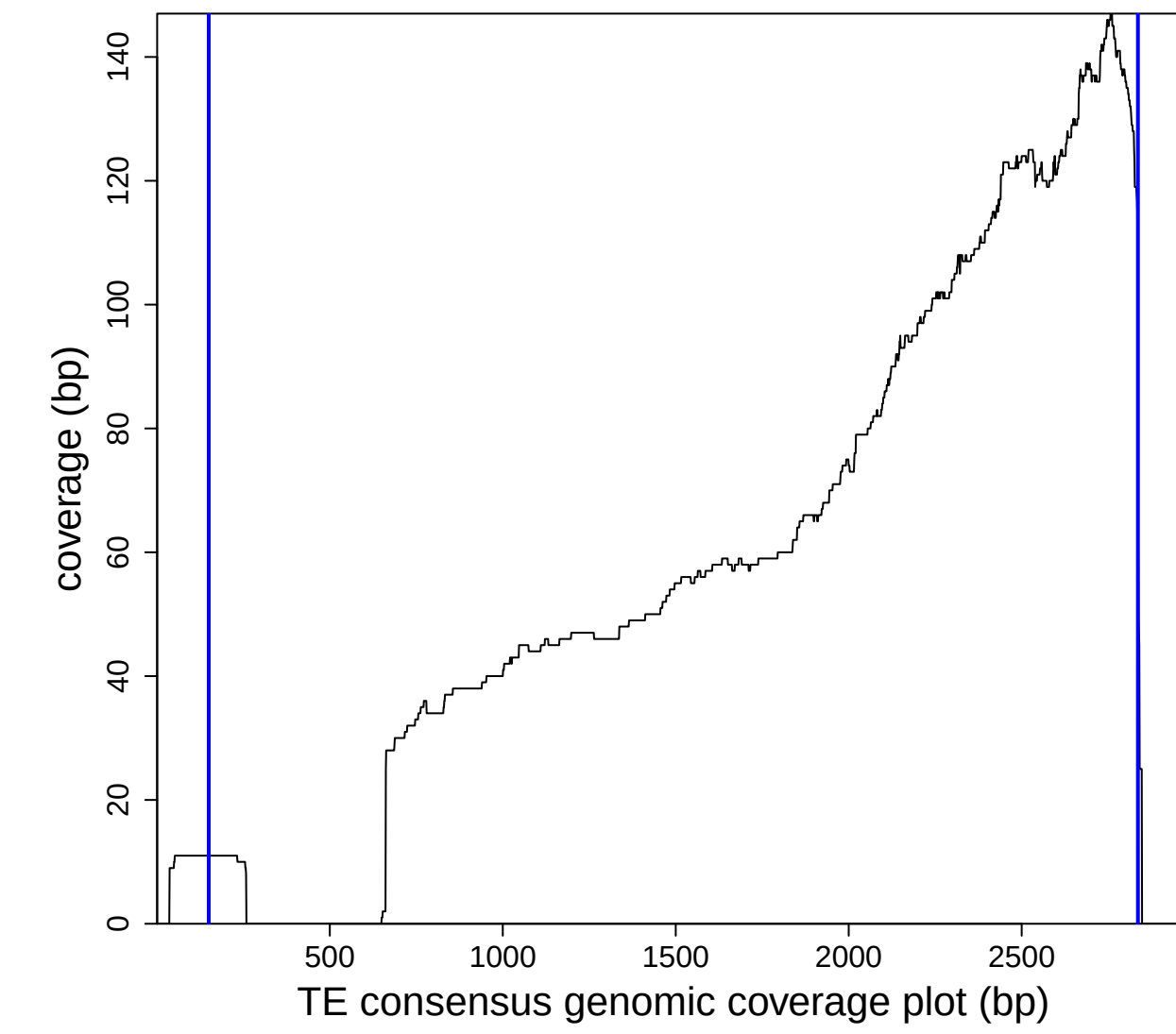
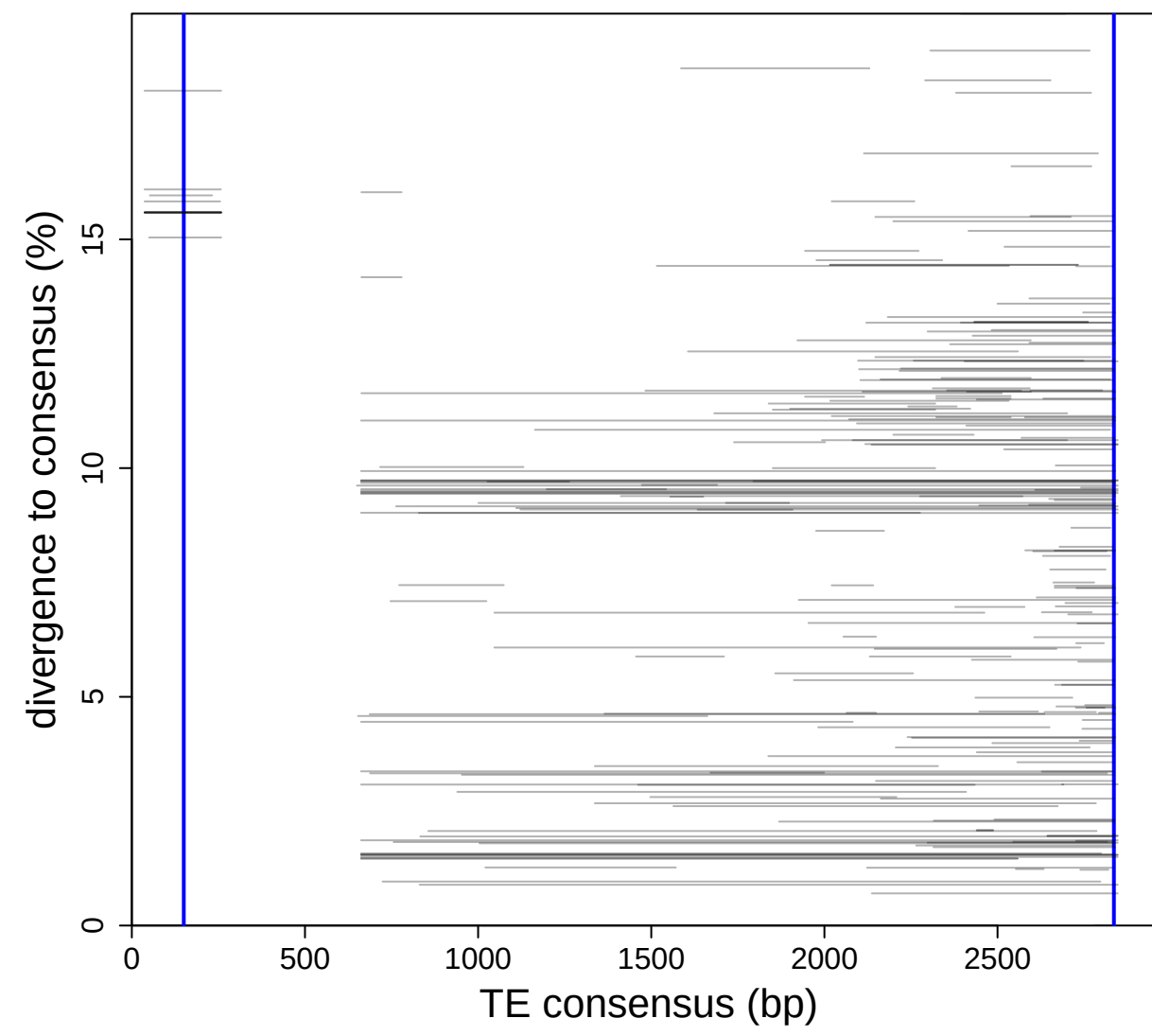
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 2641bp; fragments: 249; full length: 0 (>=2376.9bp)

After TEtrimmer 2641 bp

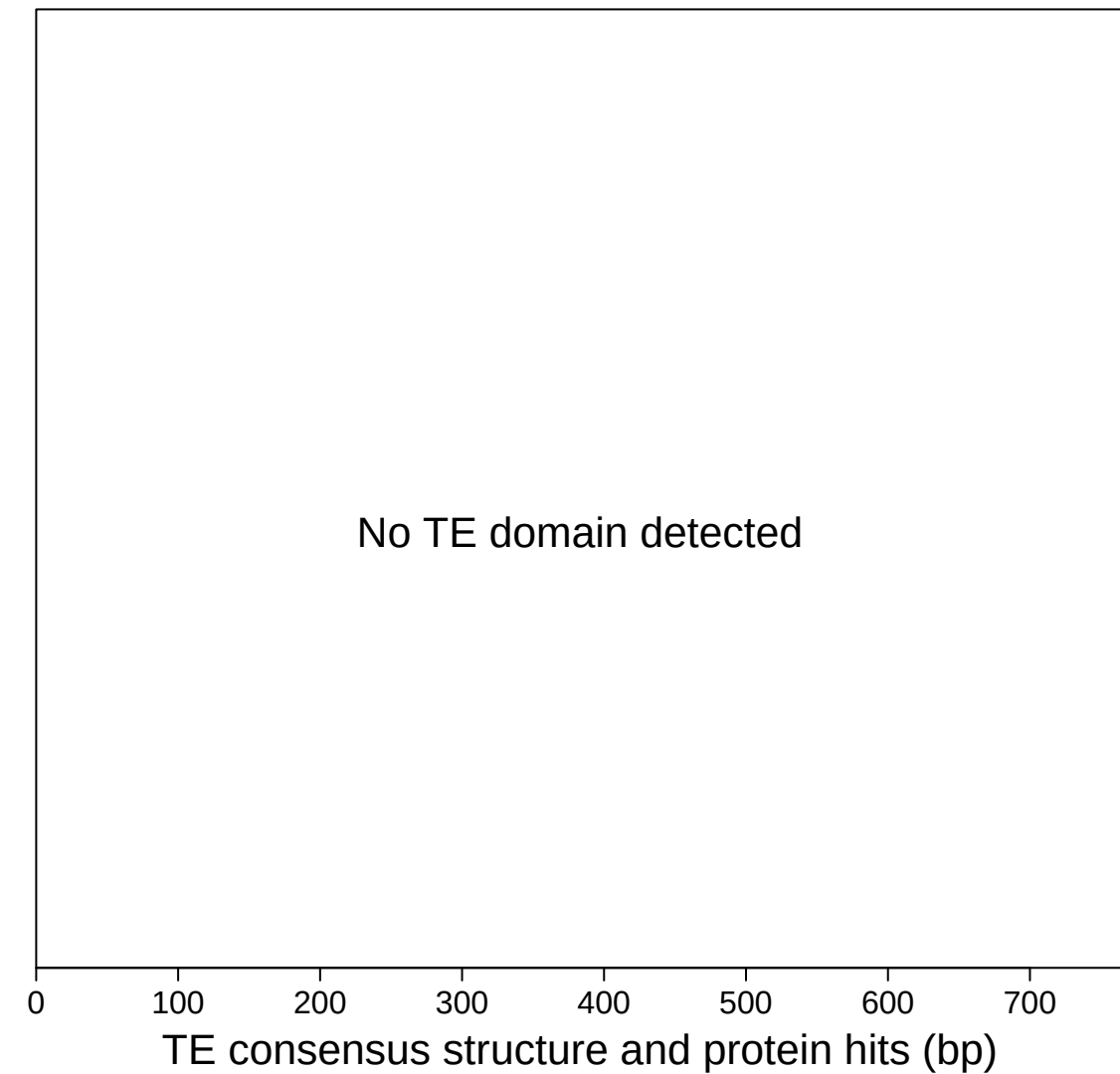
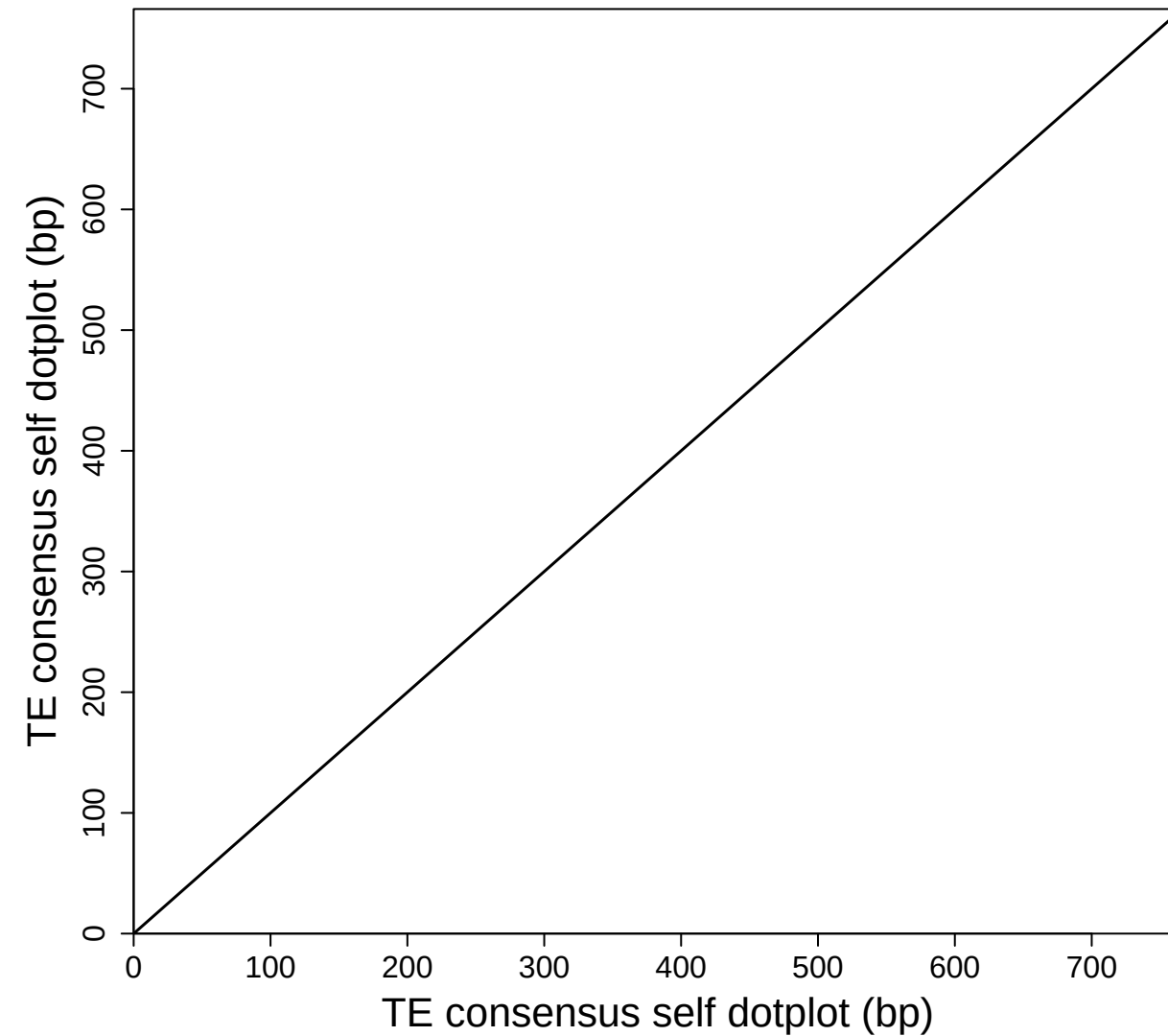
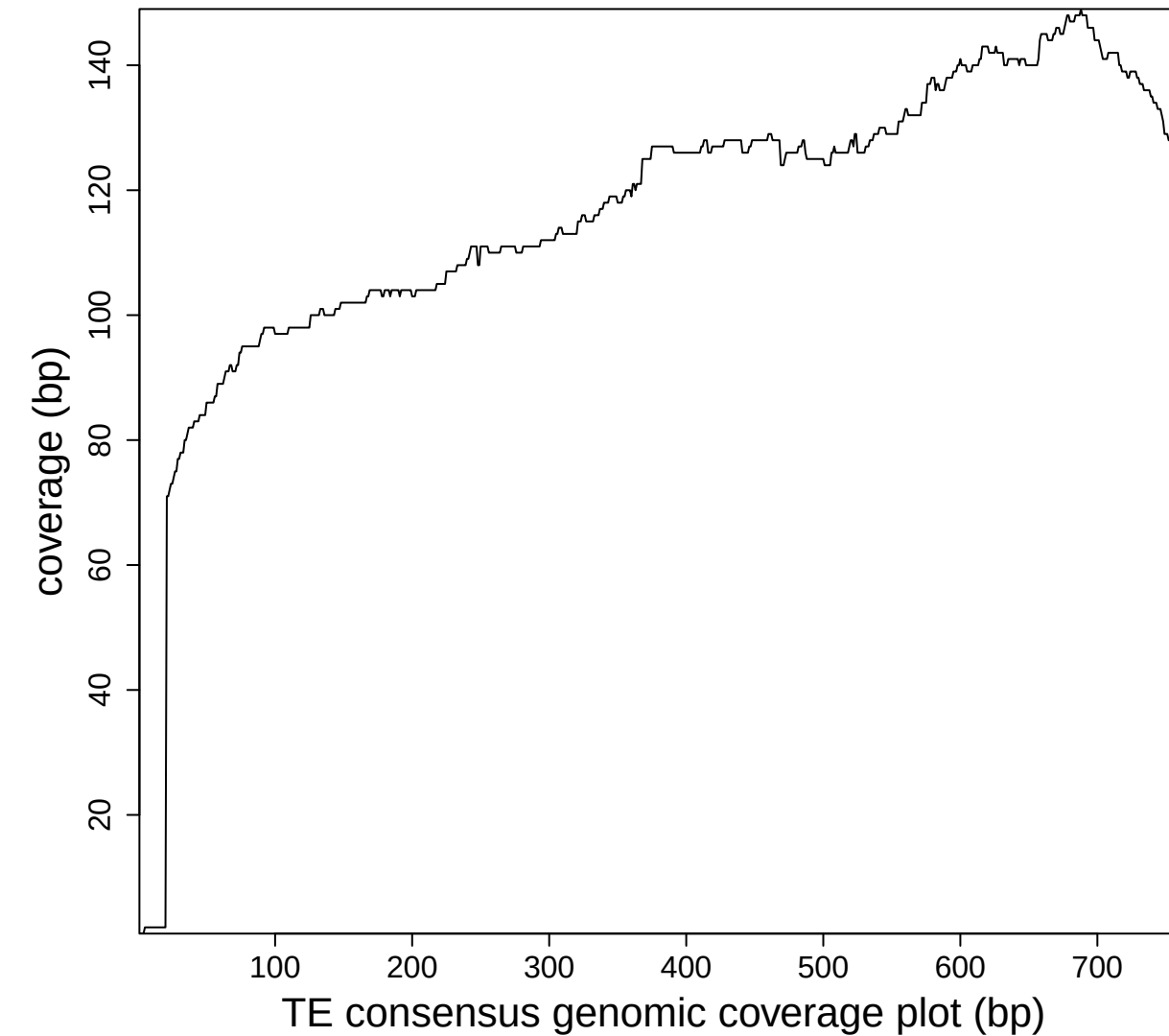
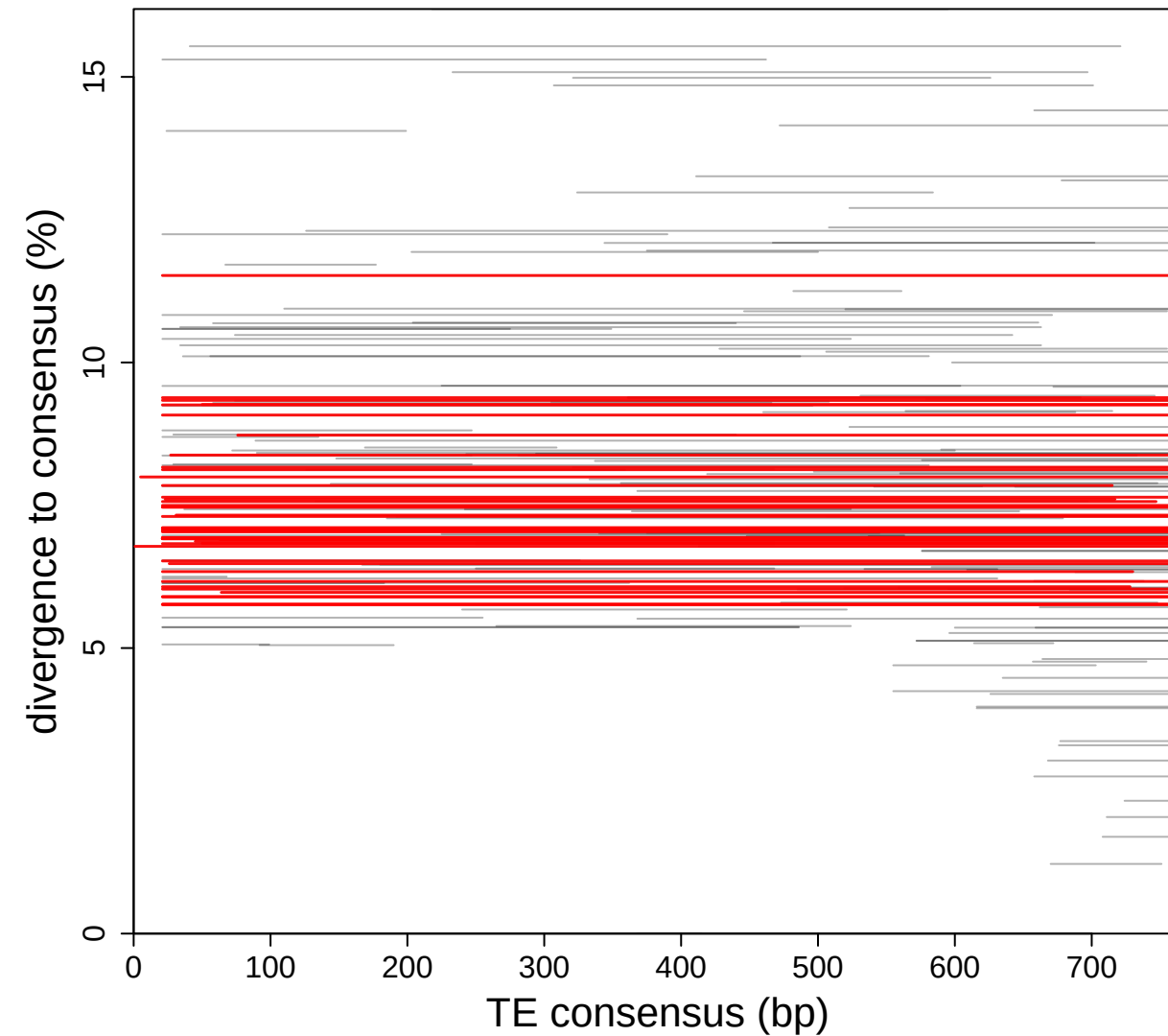


2.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.f
size: 2986bp; fragments: 236; full length: 0 (>=2687.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

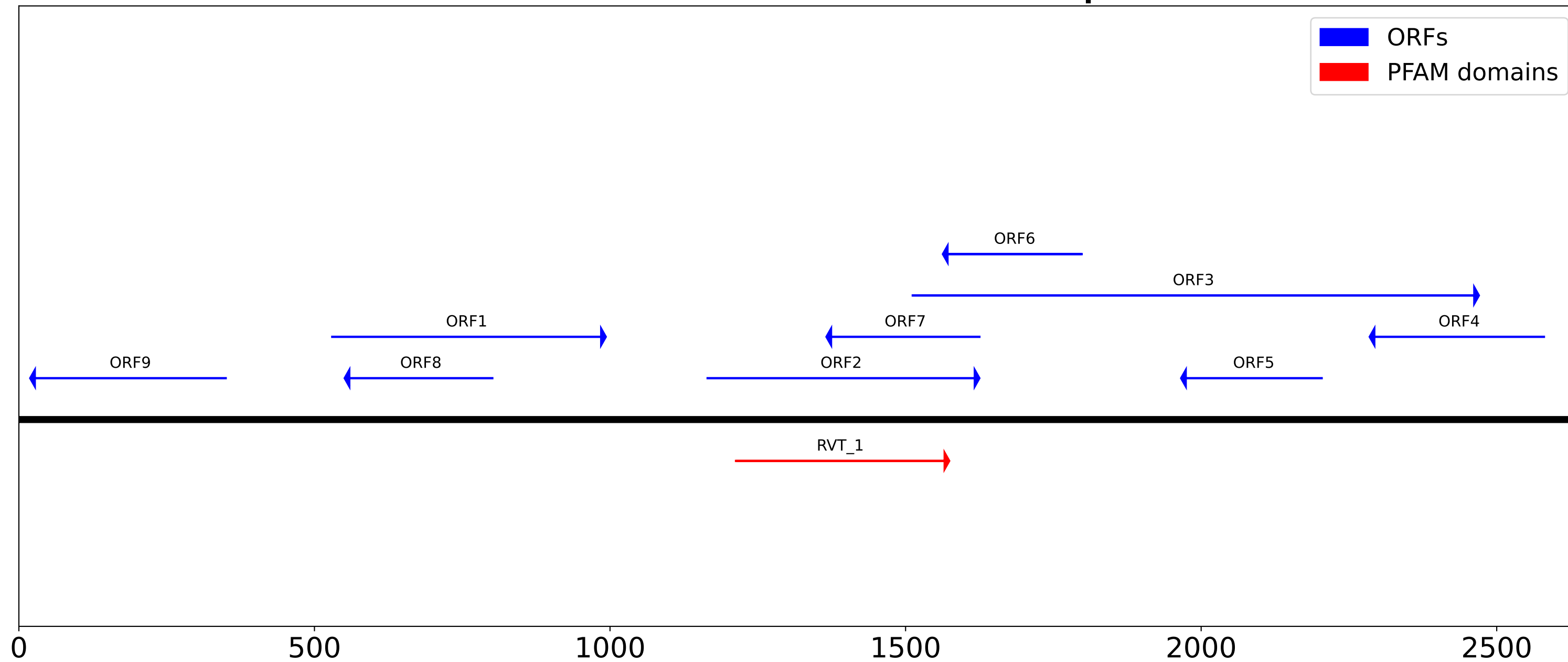


TE: rnd_5_family_9872
size: 766bp; fragments: 212; full length: 55 (≥ 689.4 bp)

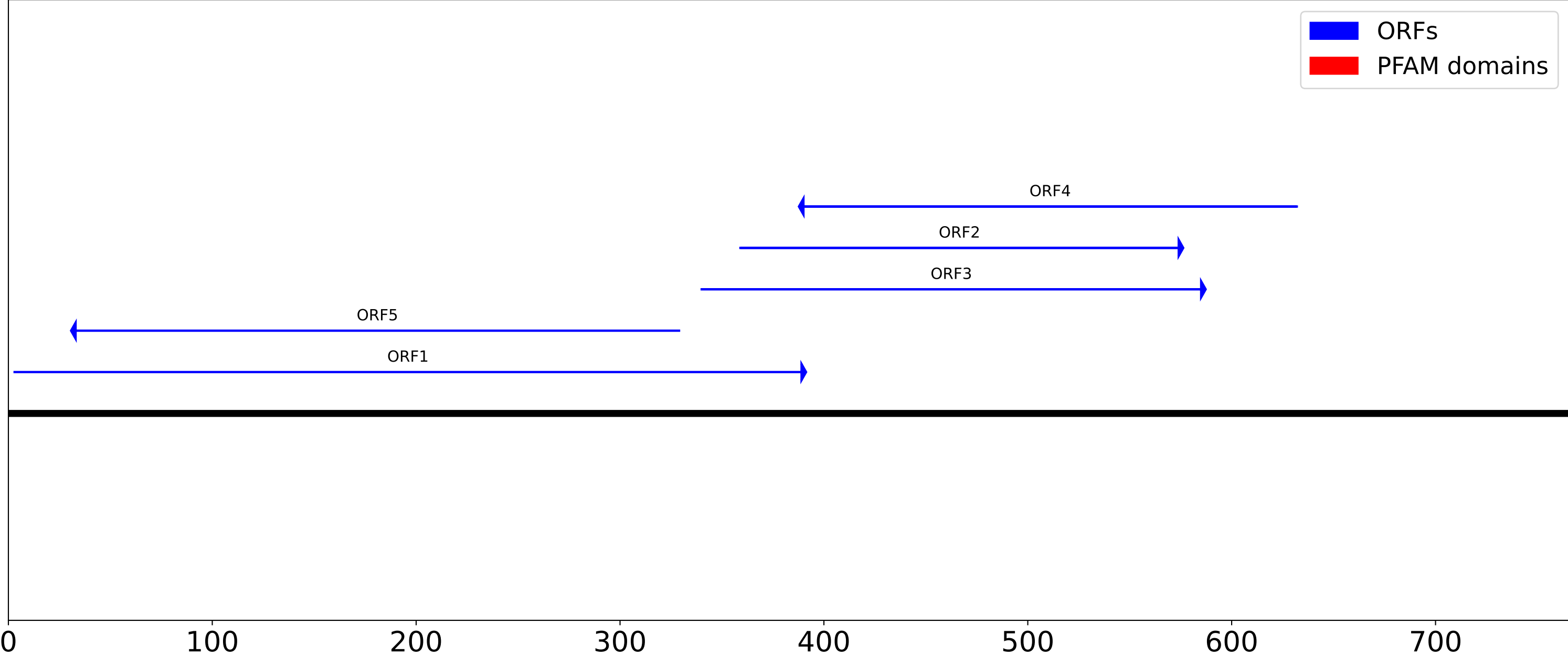


Before TEtrimmer 766 bp

After TEtrimmer ORF and PFAM domain plot



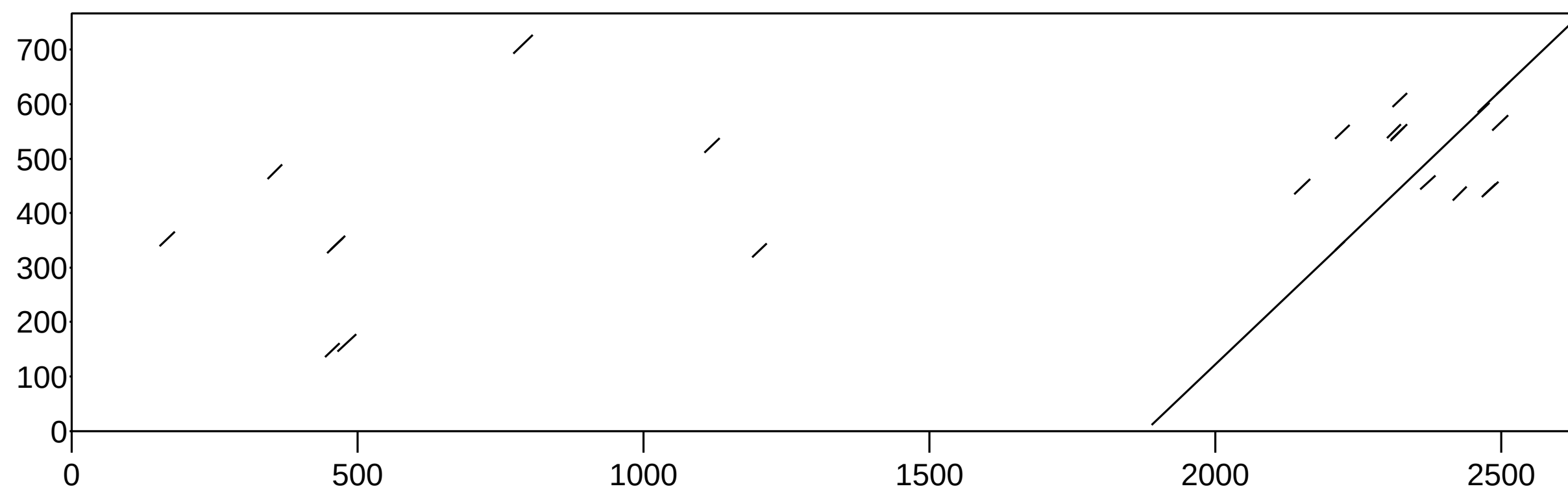
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)