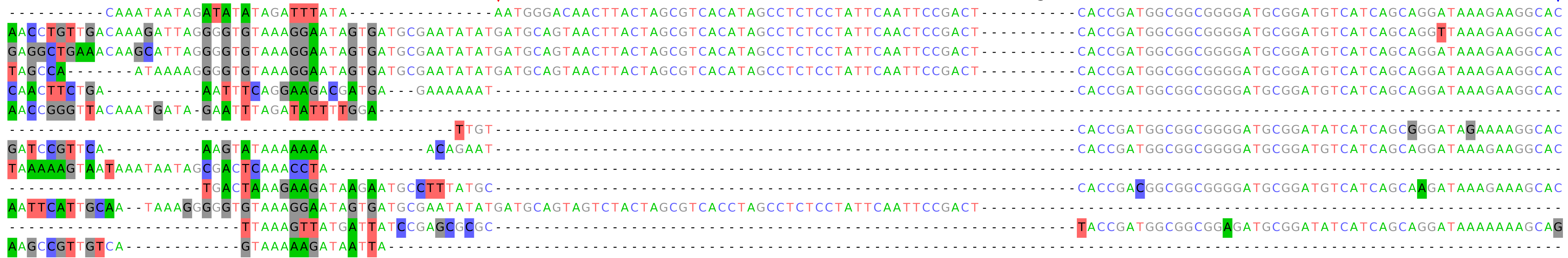
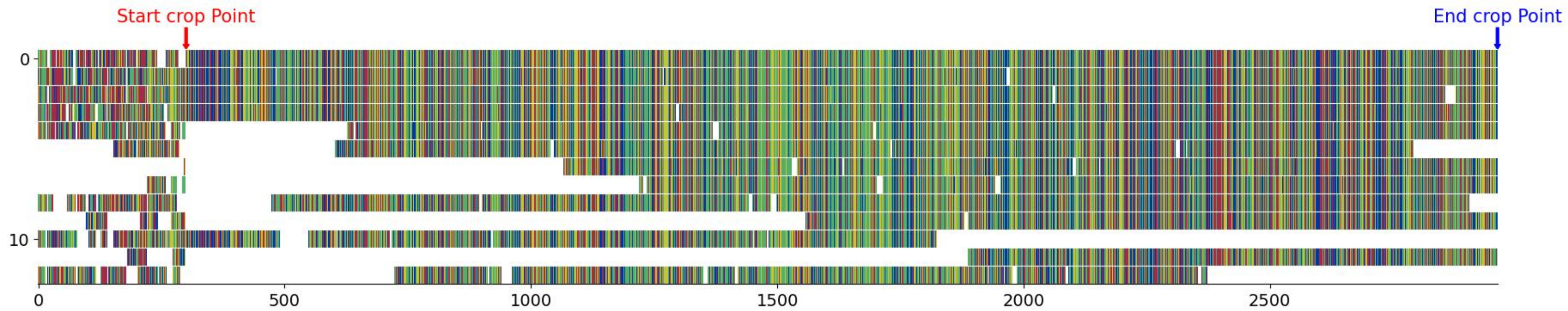


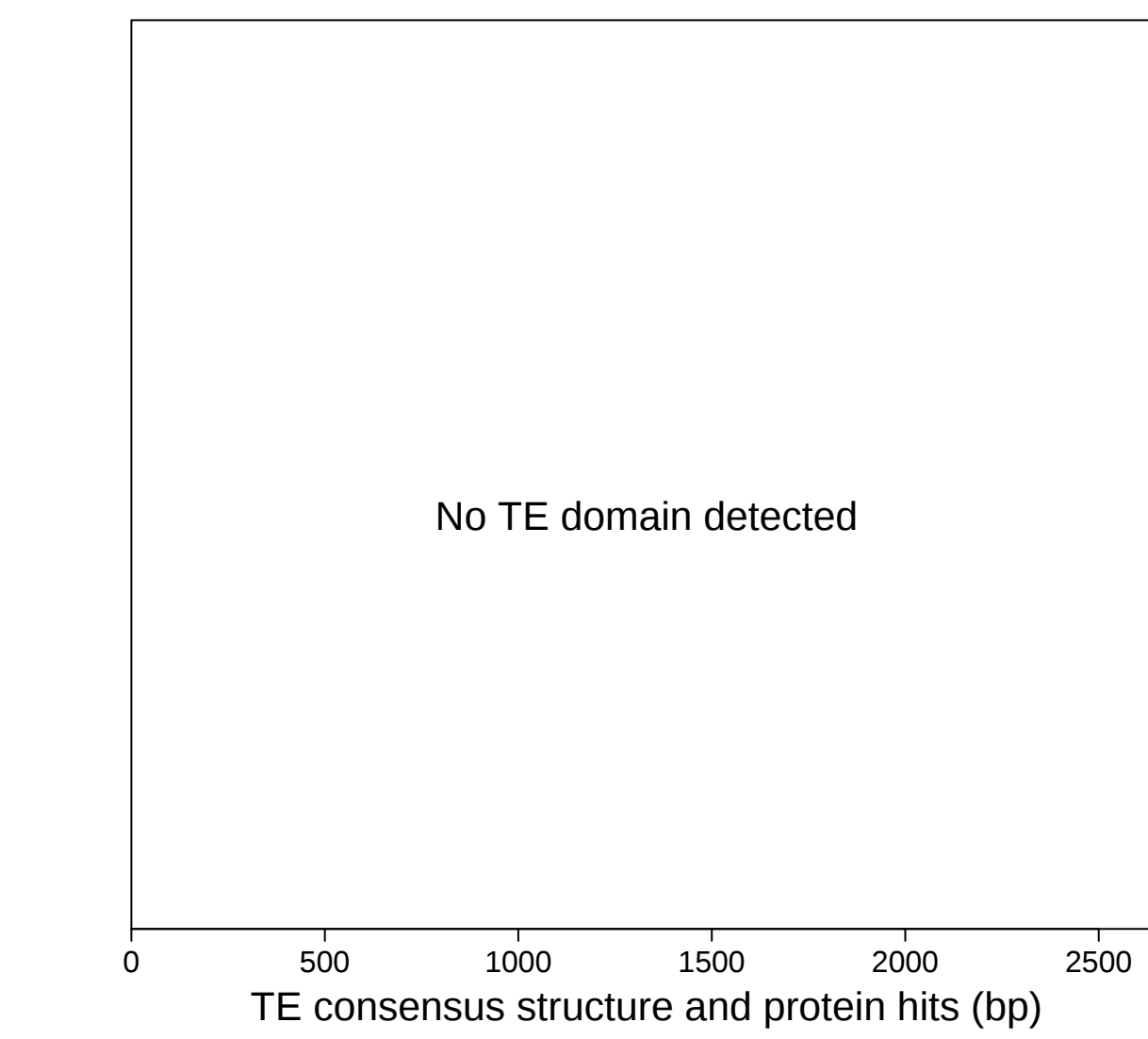
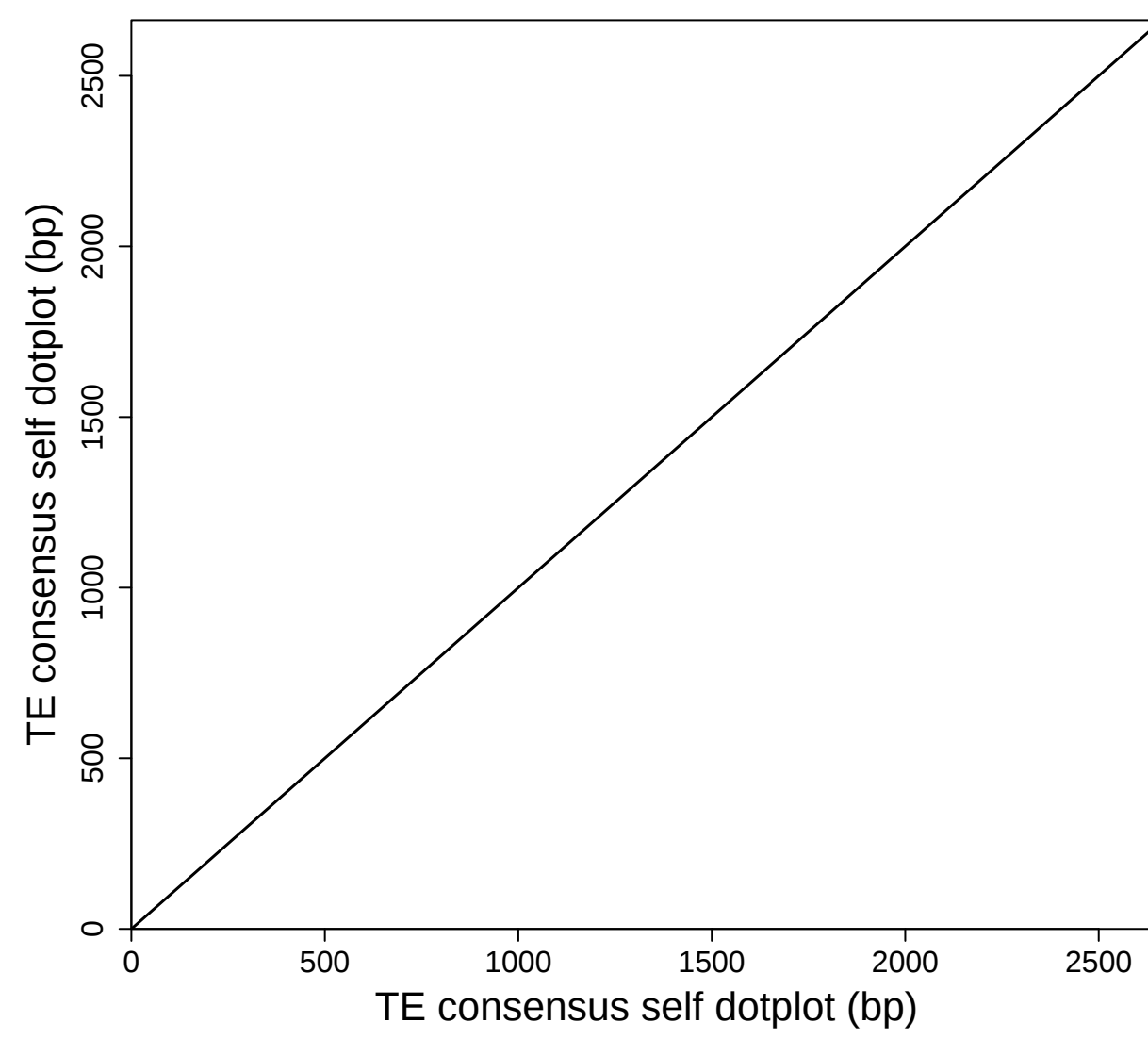
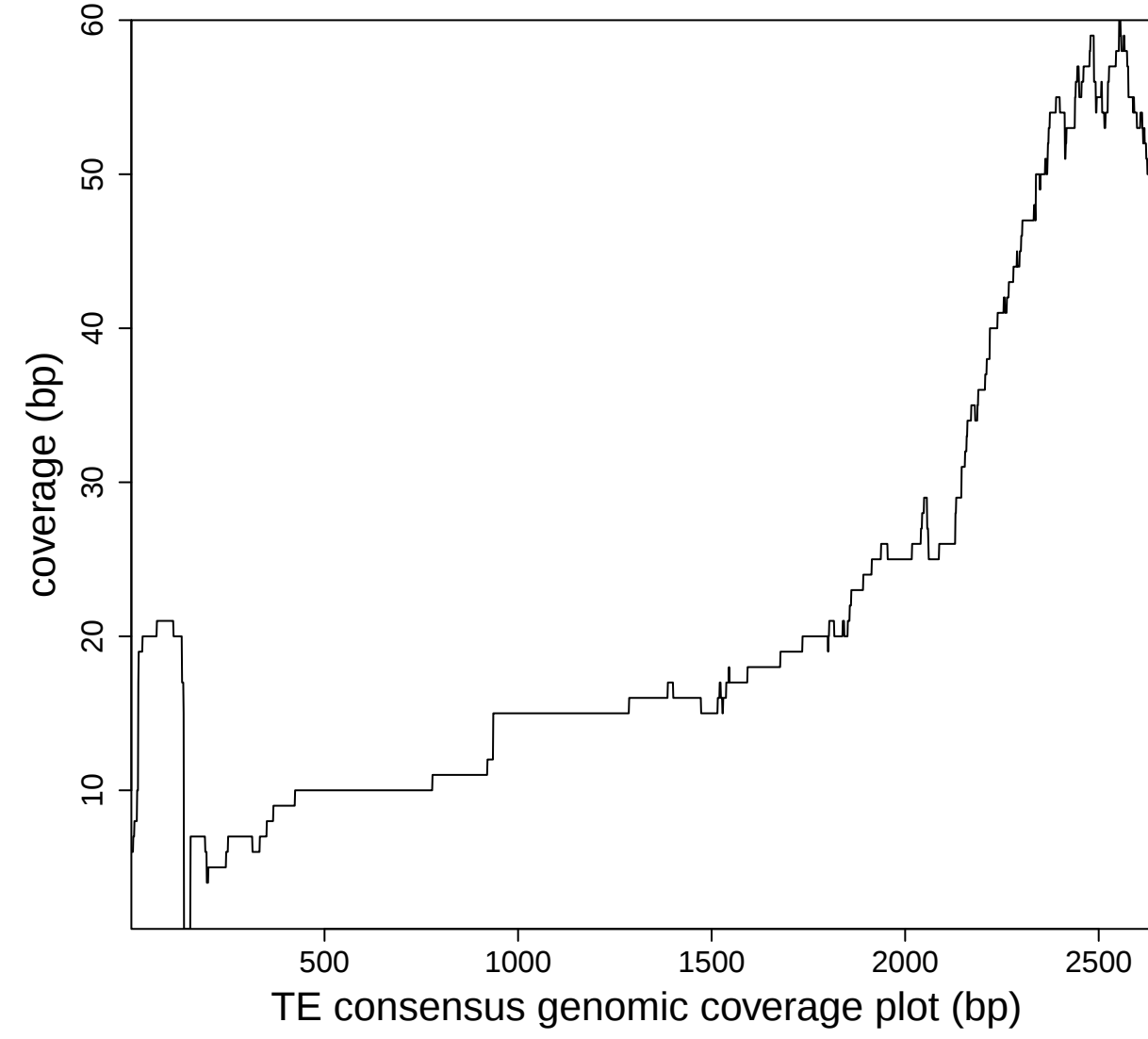
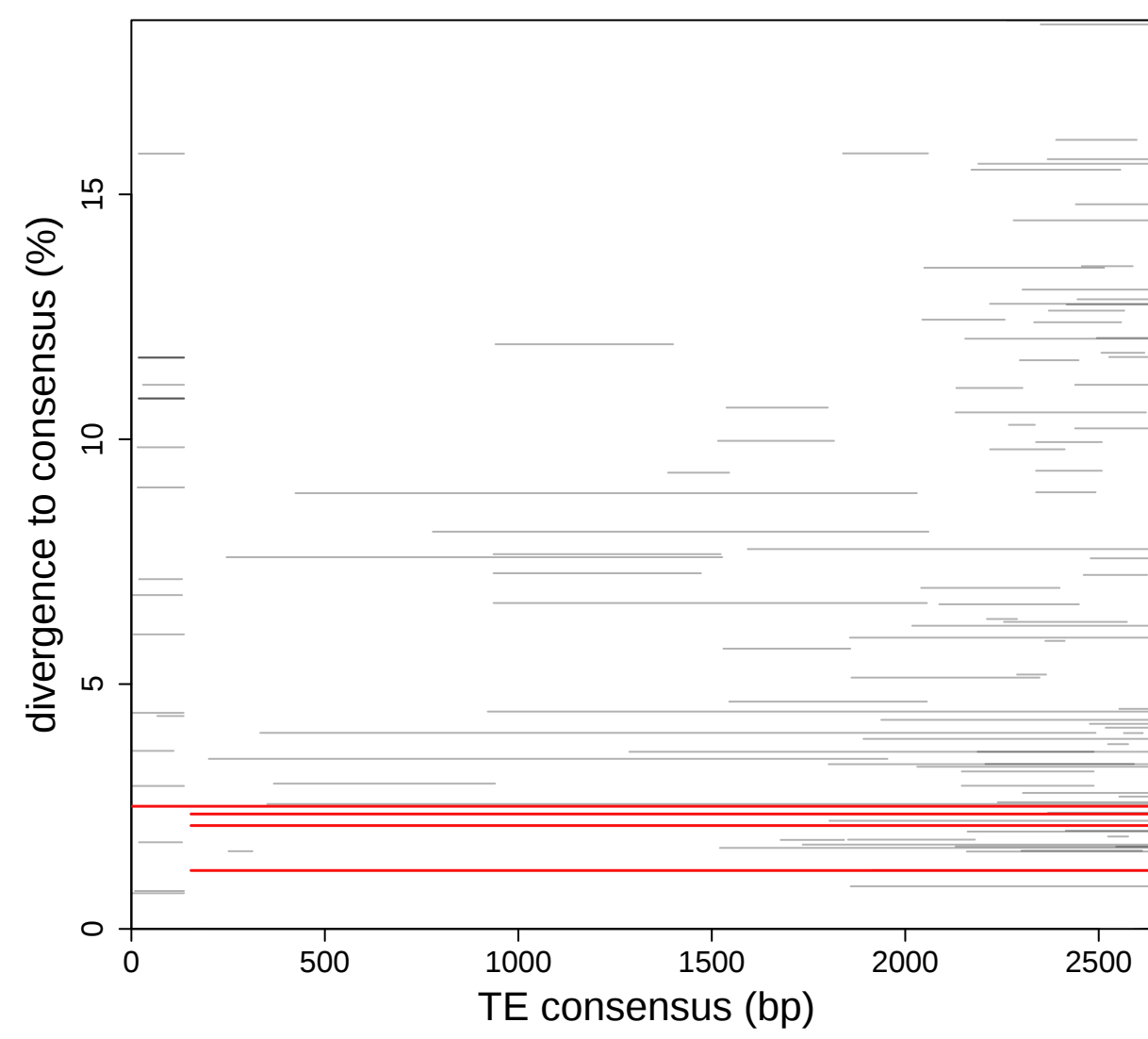
MSA length = 2663





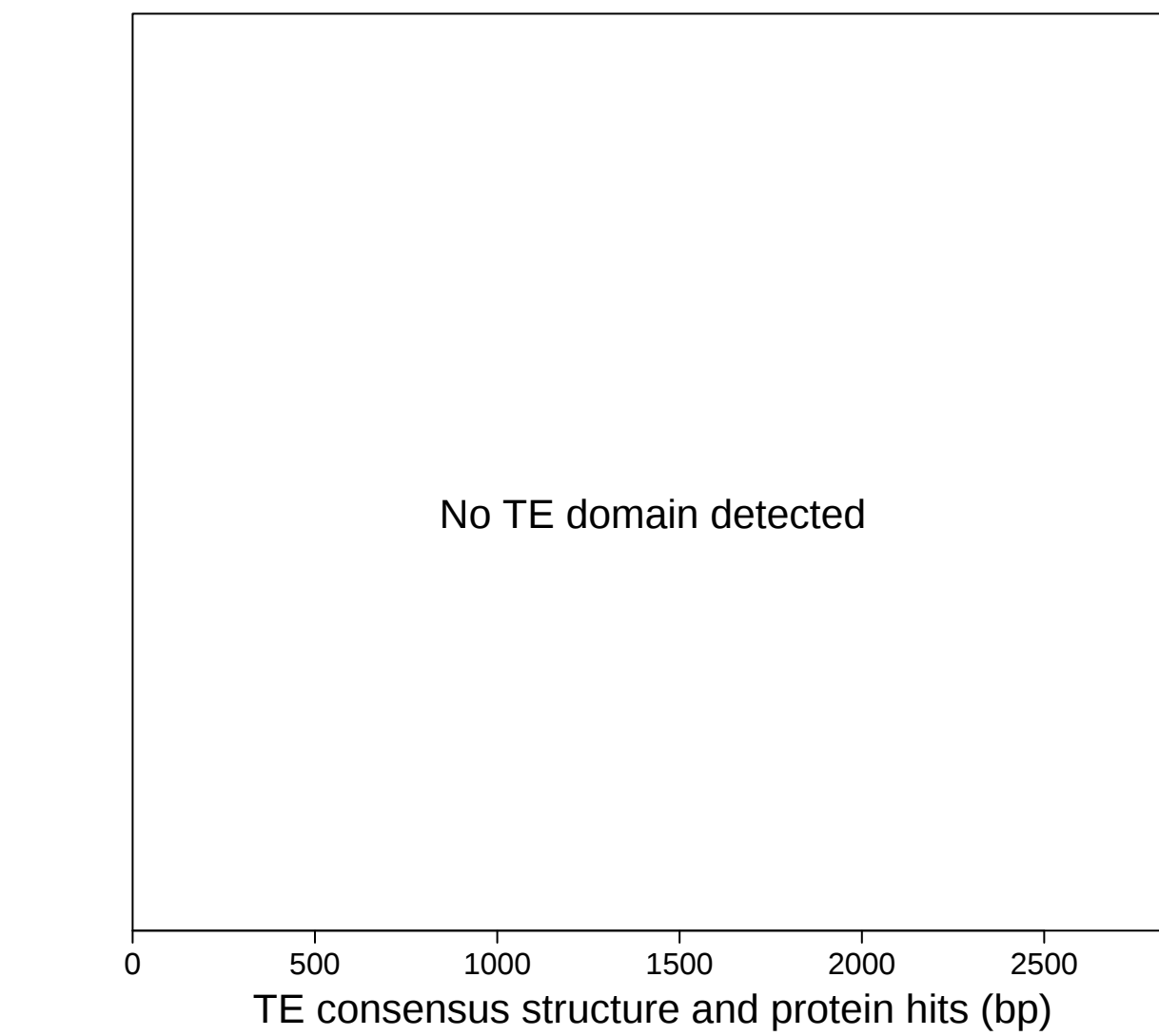
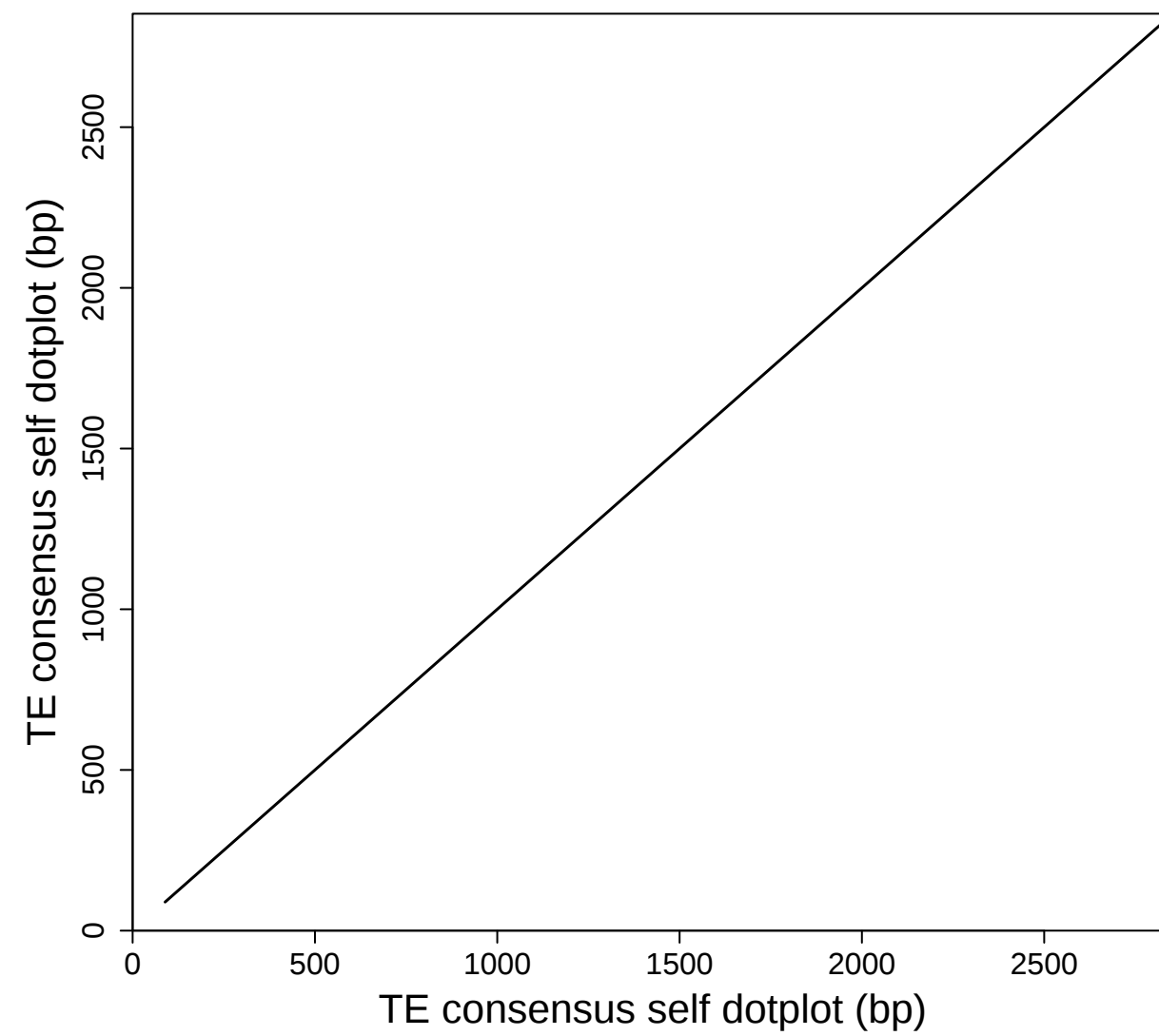
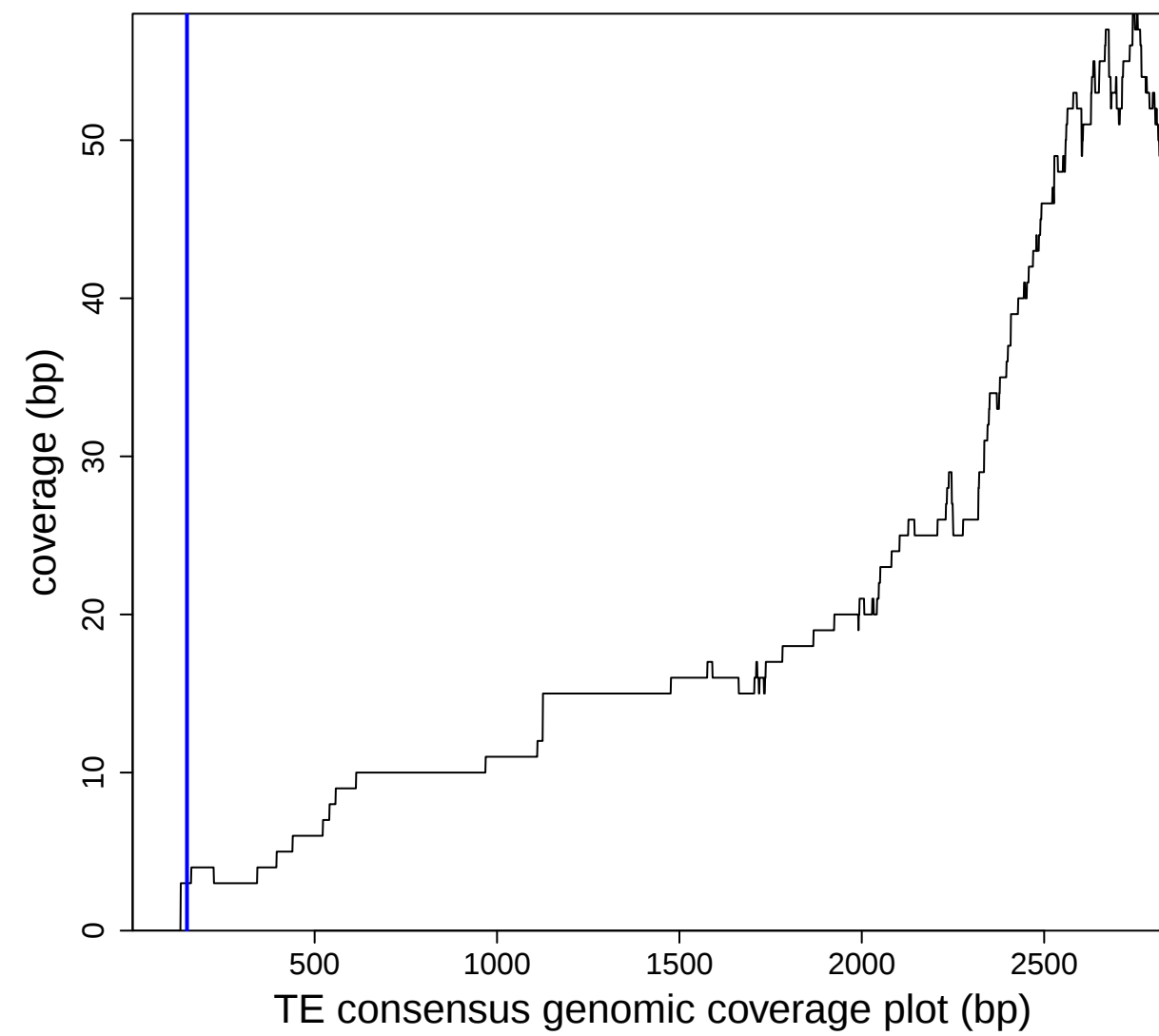
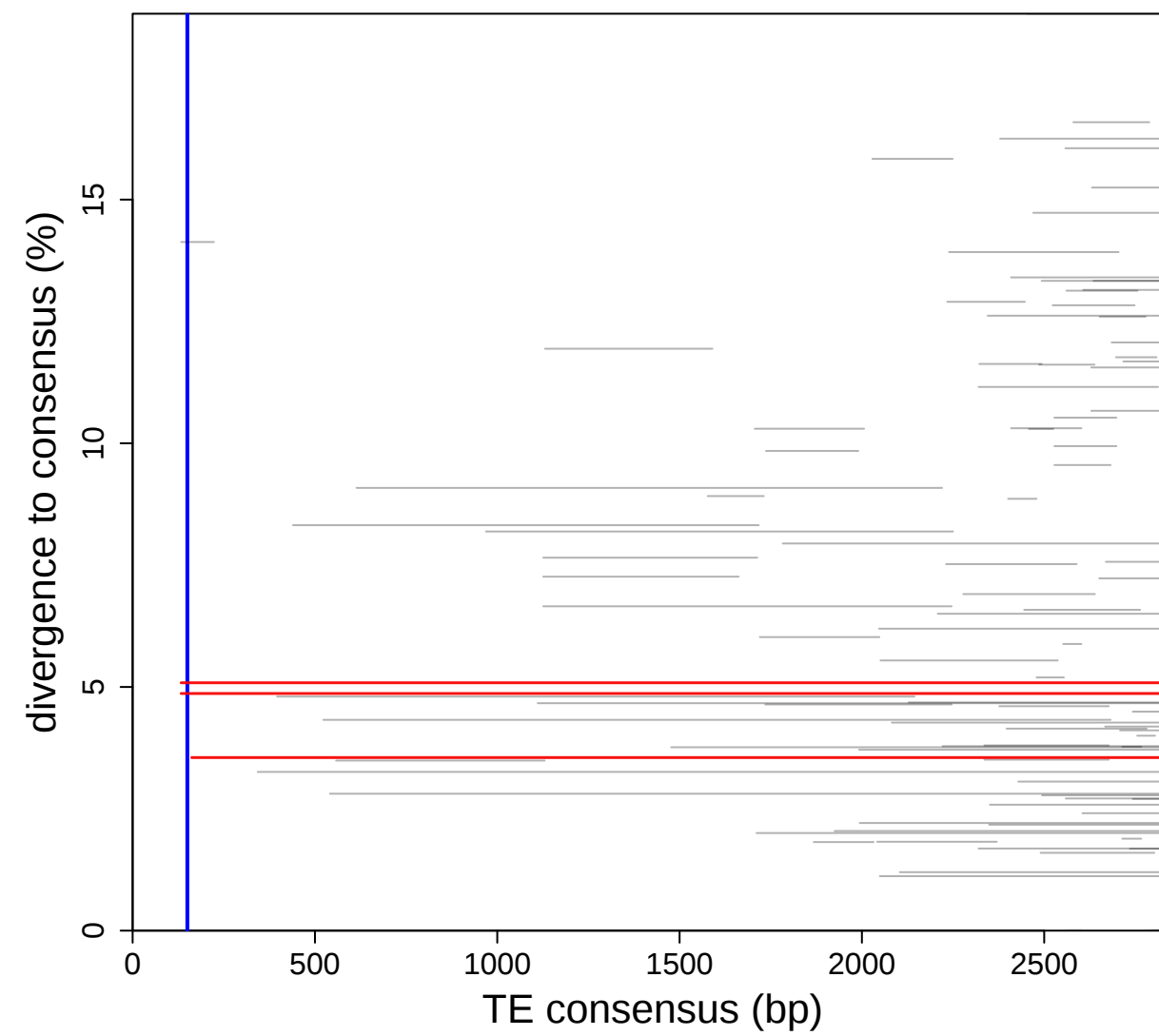
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 2663bp; fragments: 125; full length: 4 (>=2396.7bp)

After TETrimmer 2663 bp

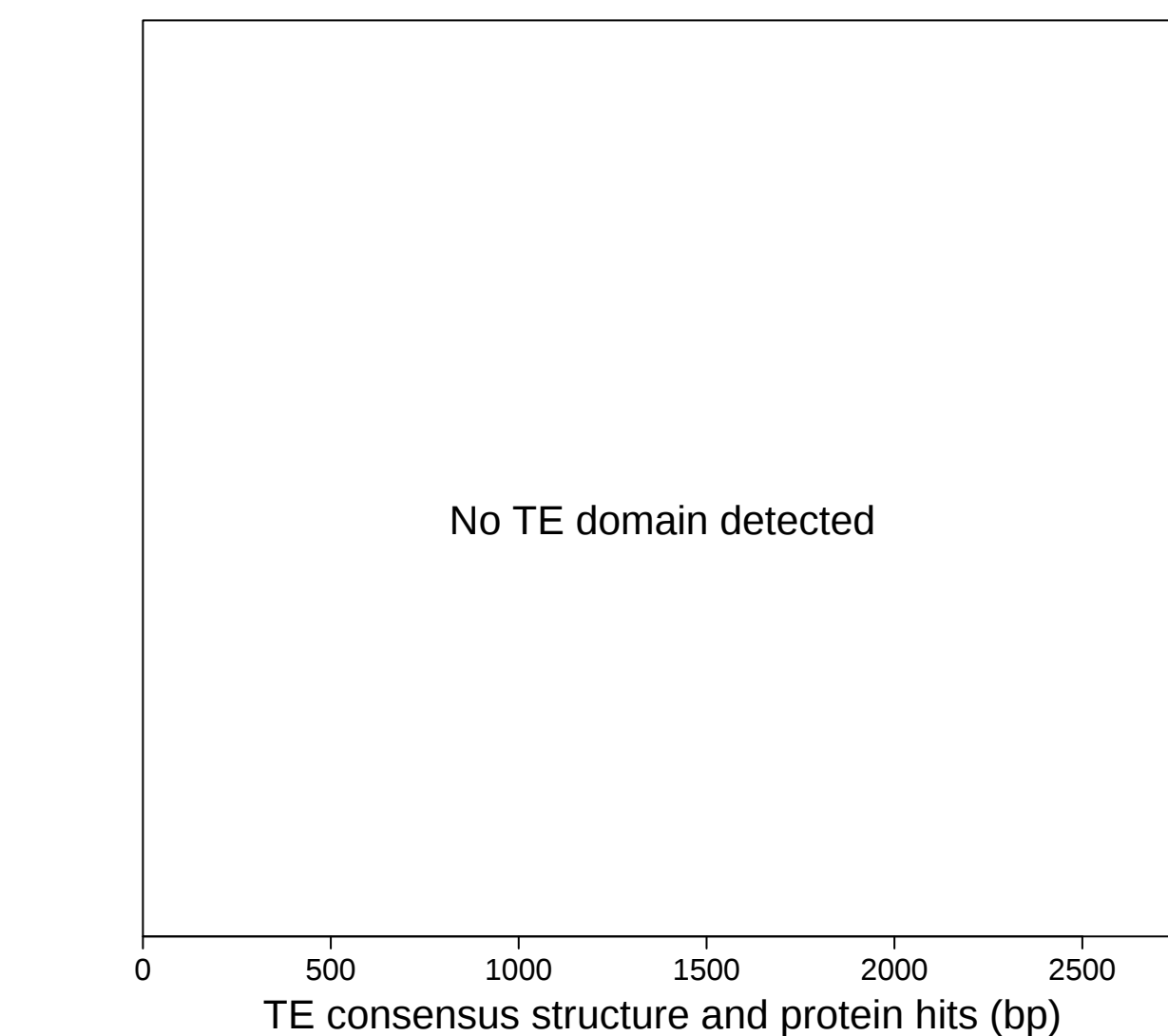
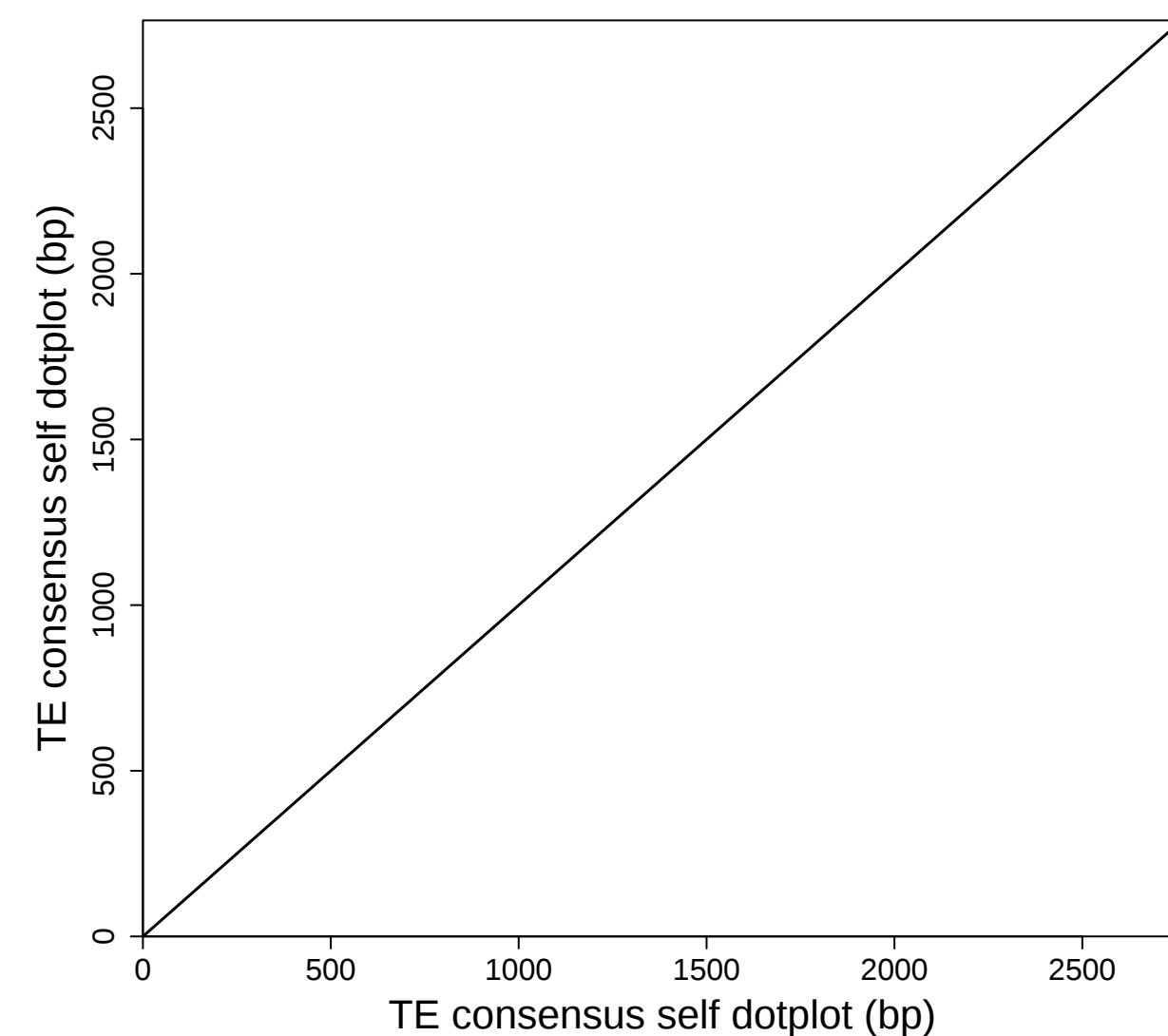
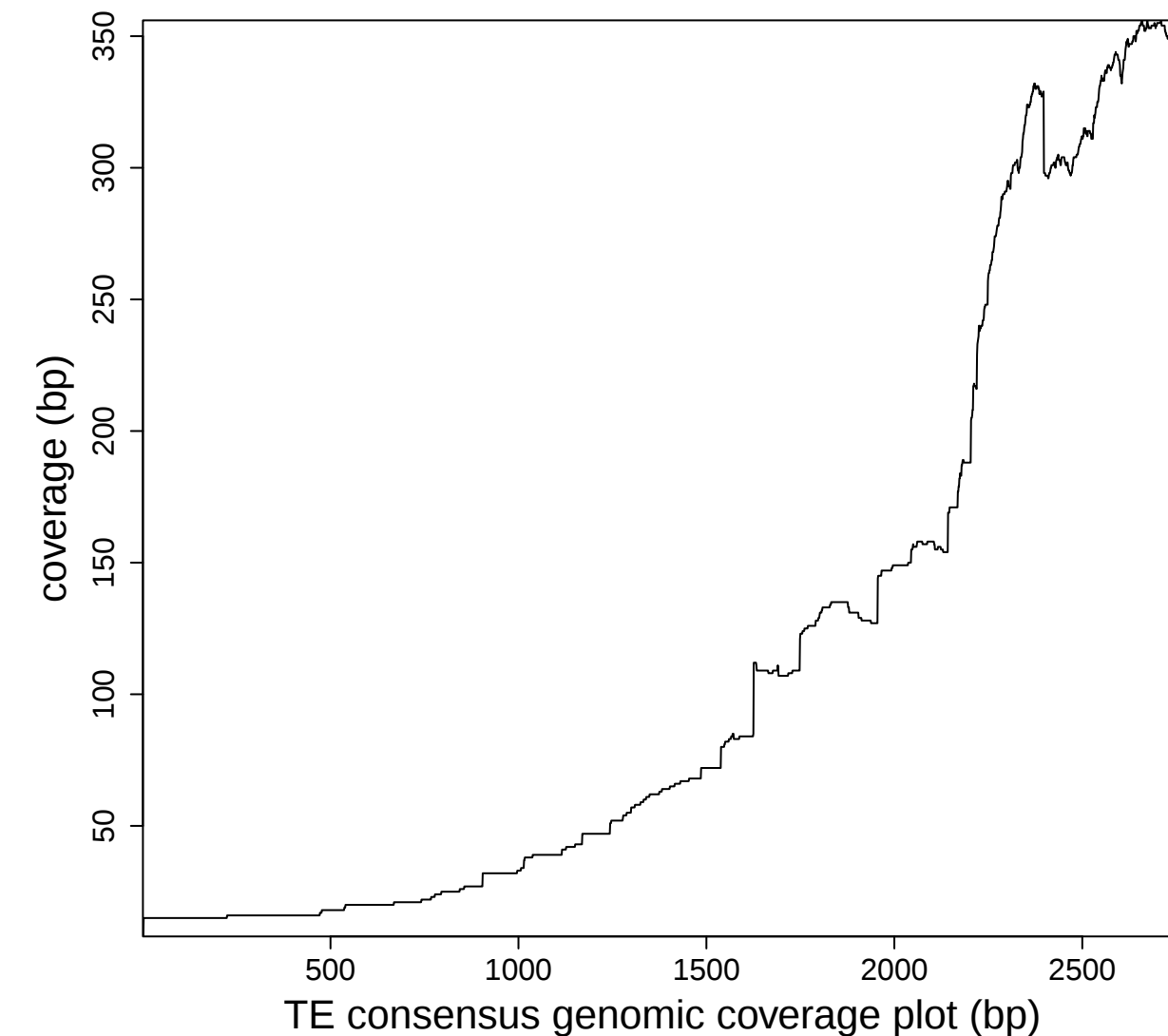
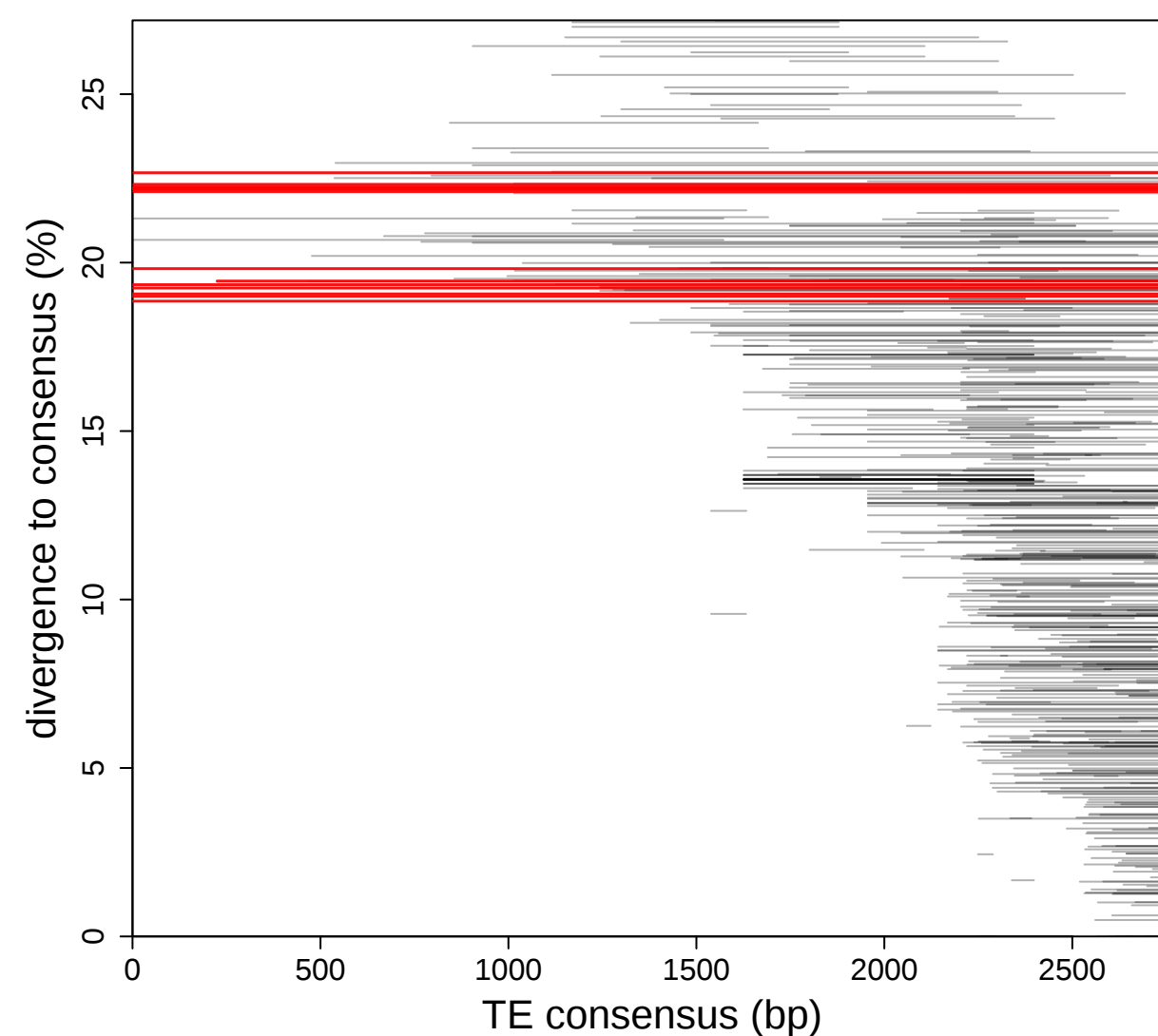


6.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 2853bp; fragments: 100; full length: 3 (>=2567.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

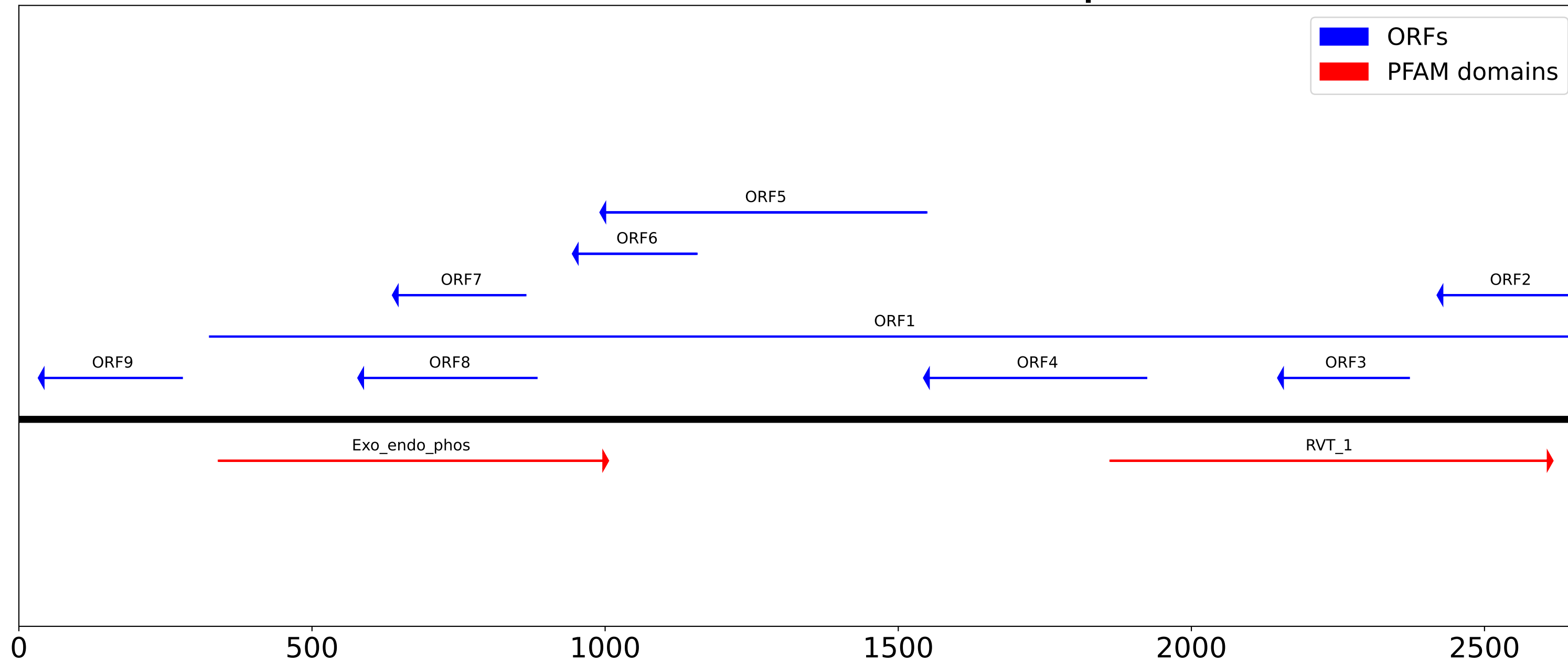


TE: rnd_4_family_1796
size: 2765bp; fragments: 529; full length: 14 (≥ 2488.5 bp)

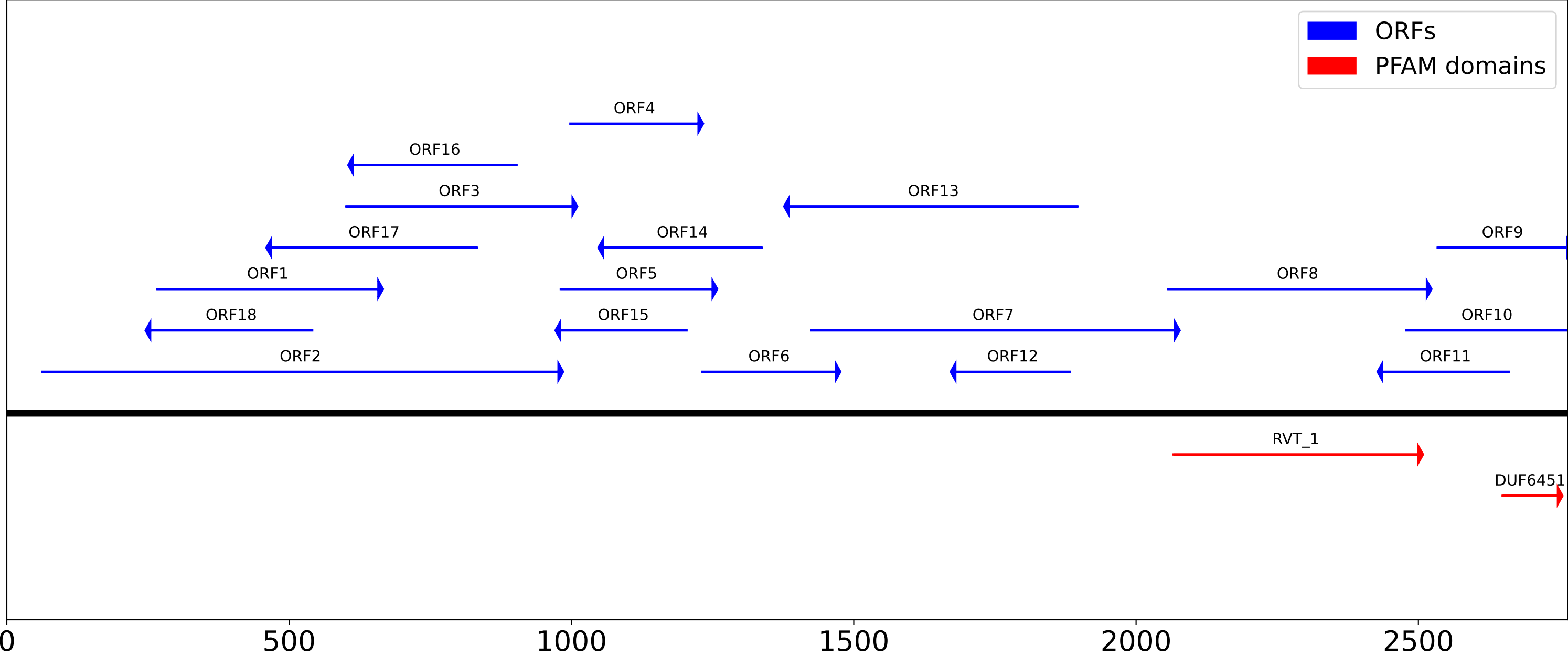


Before TEtrimmer 2765 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

