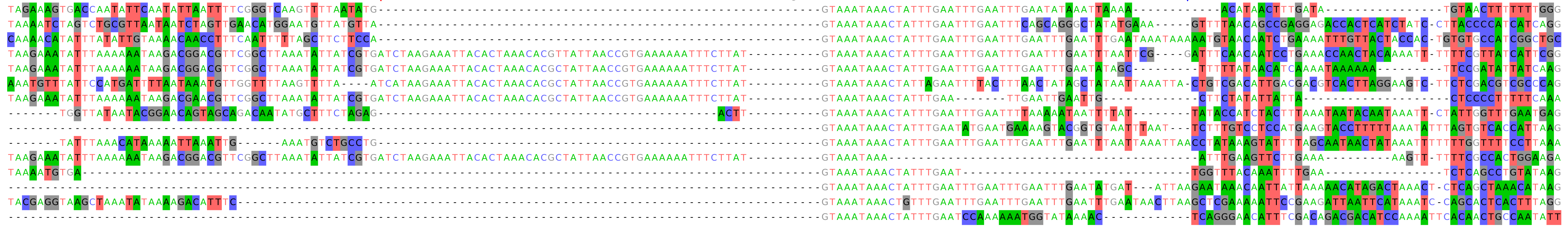
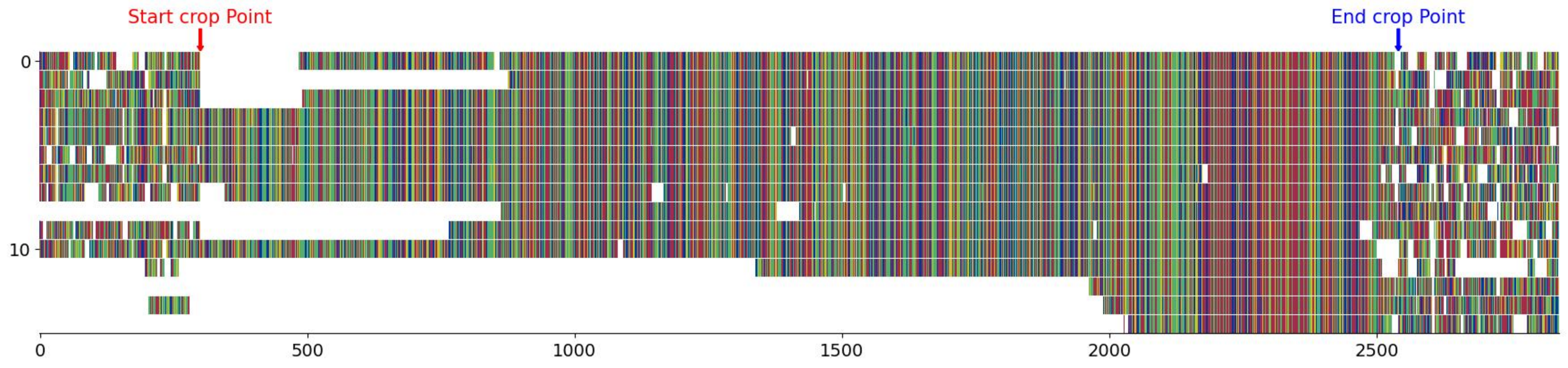


MSA length = 2241



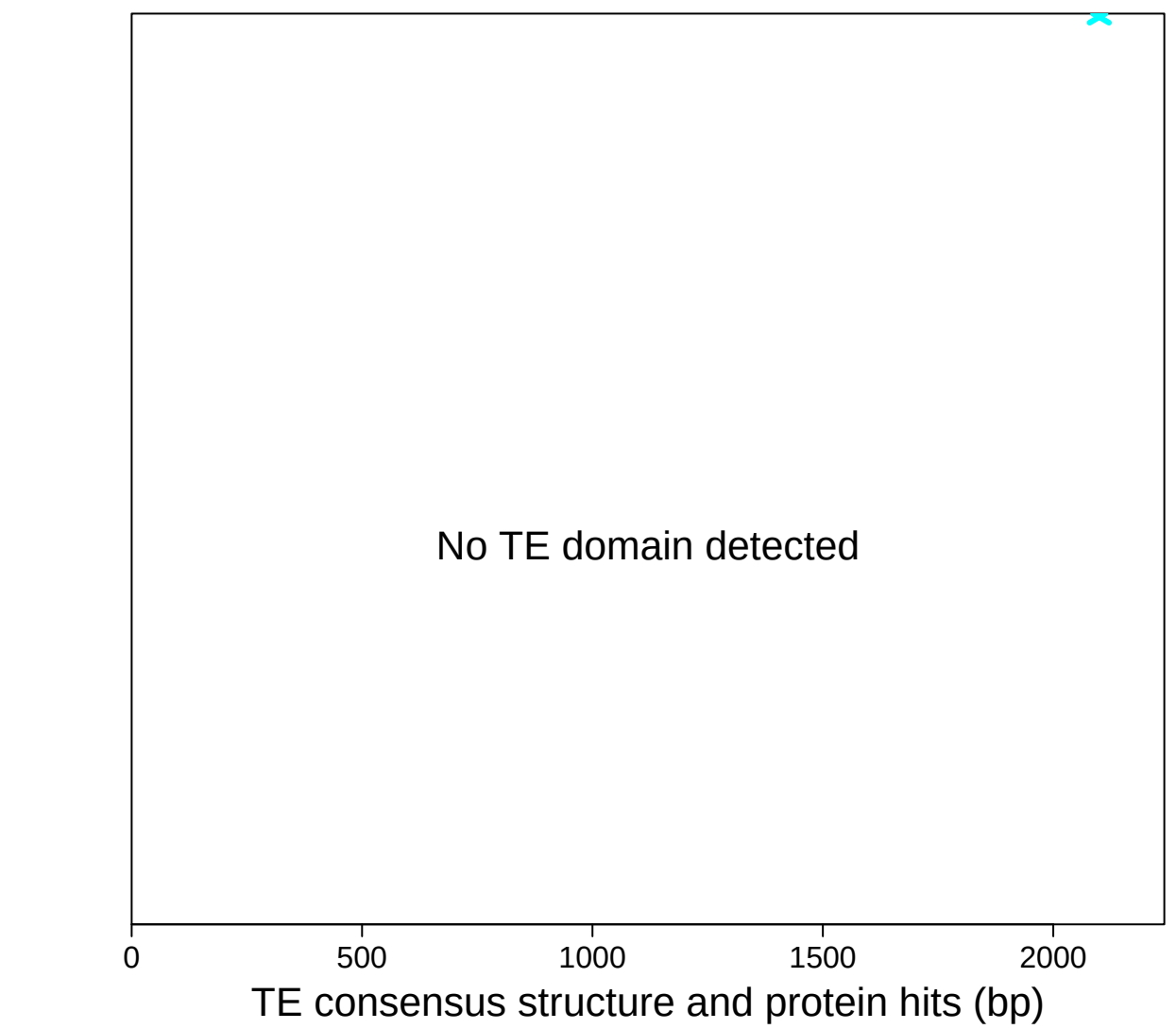
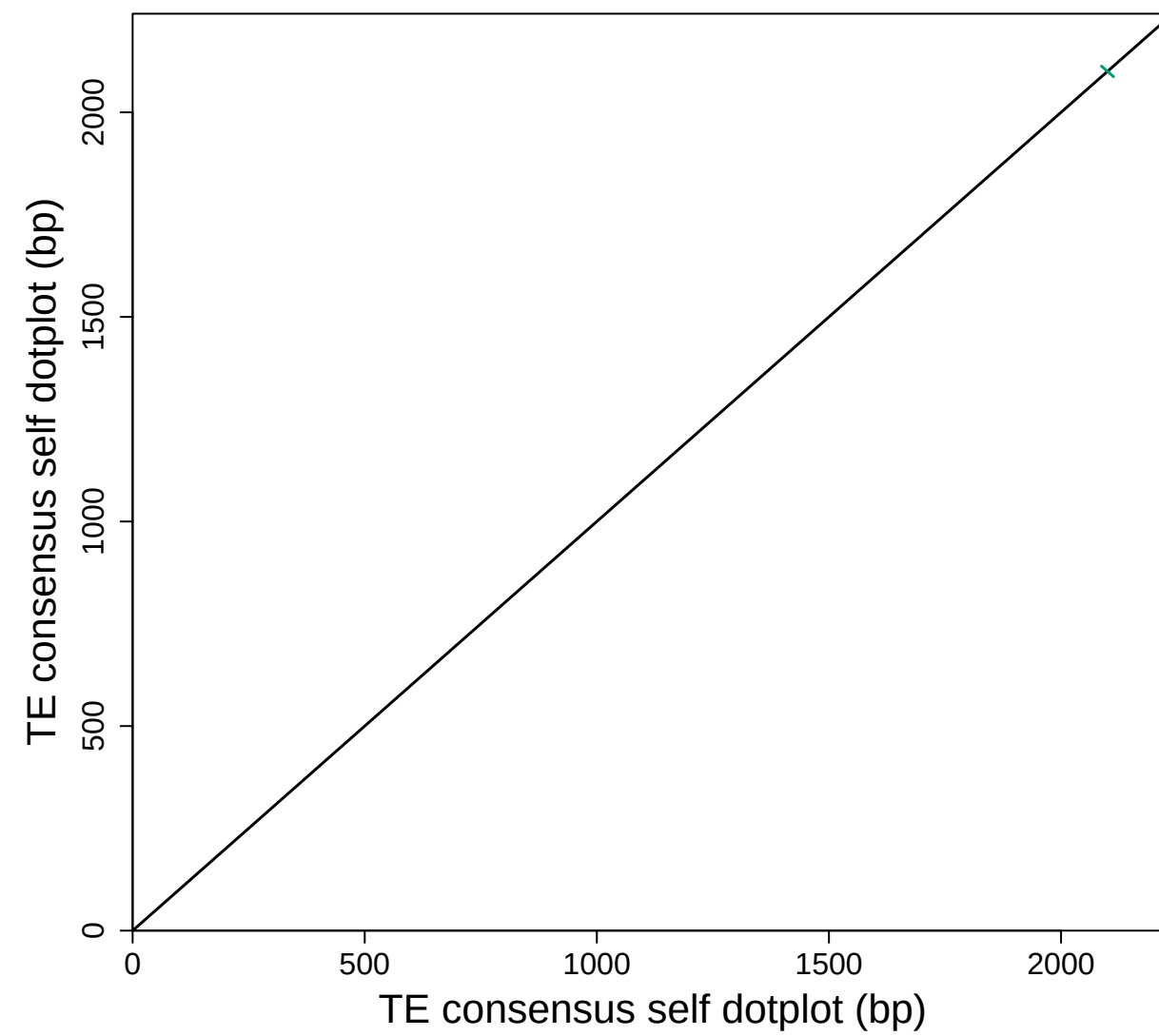
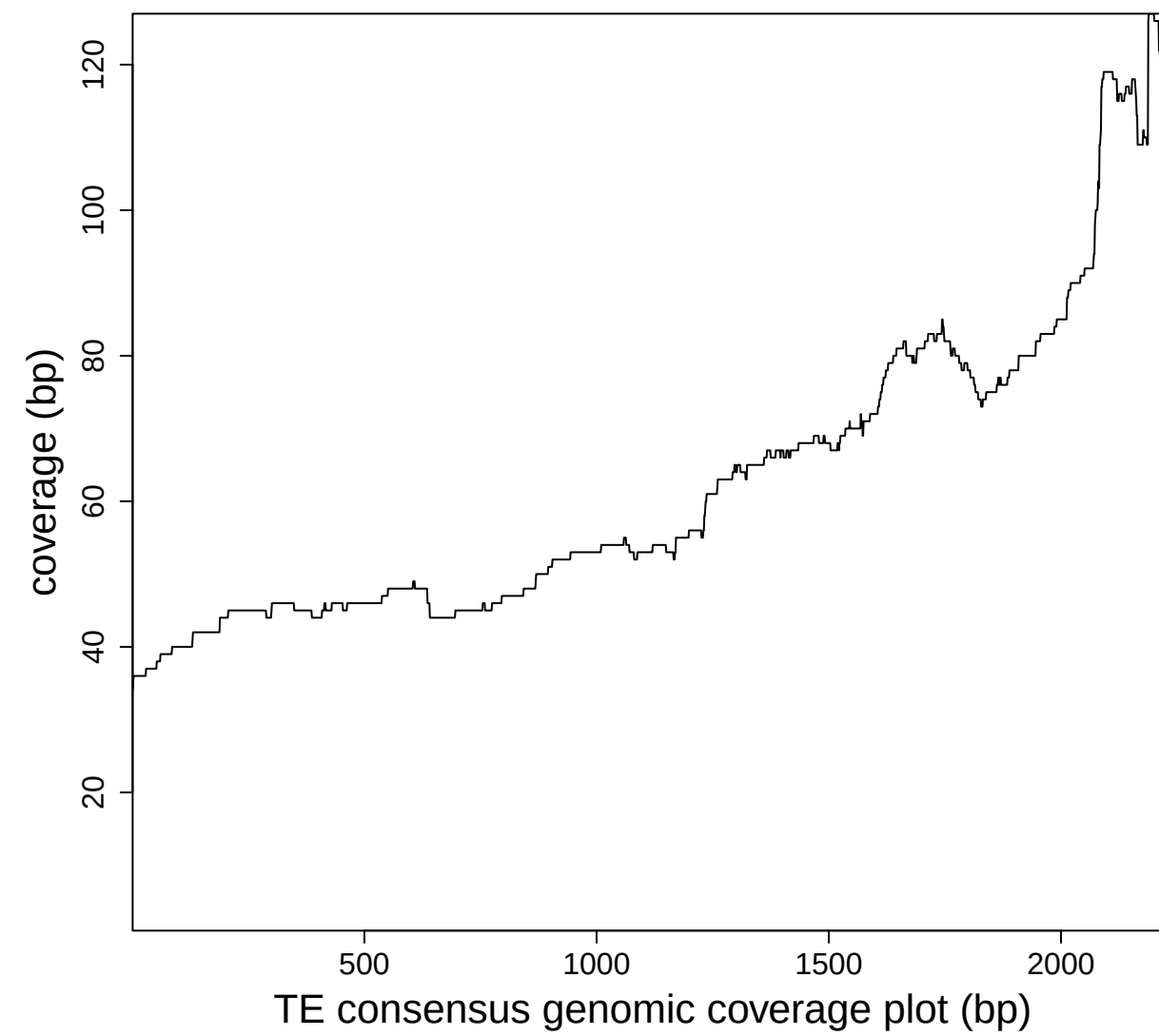
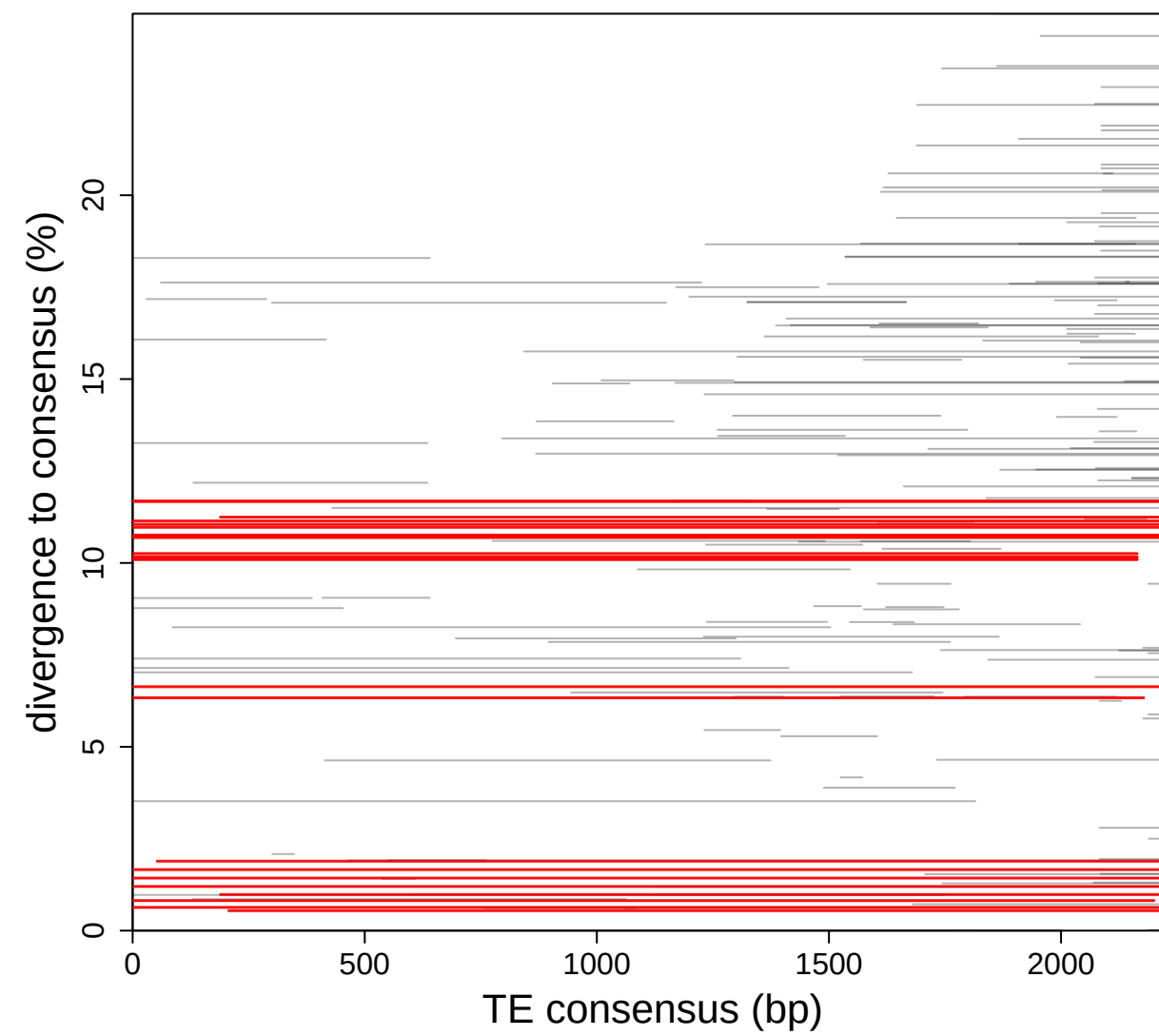
Start crop Point

End crop Point



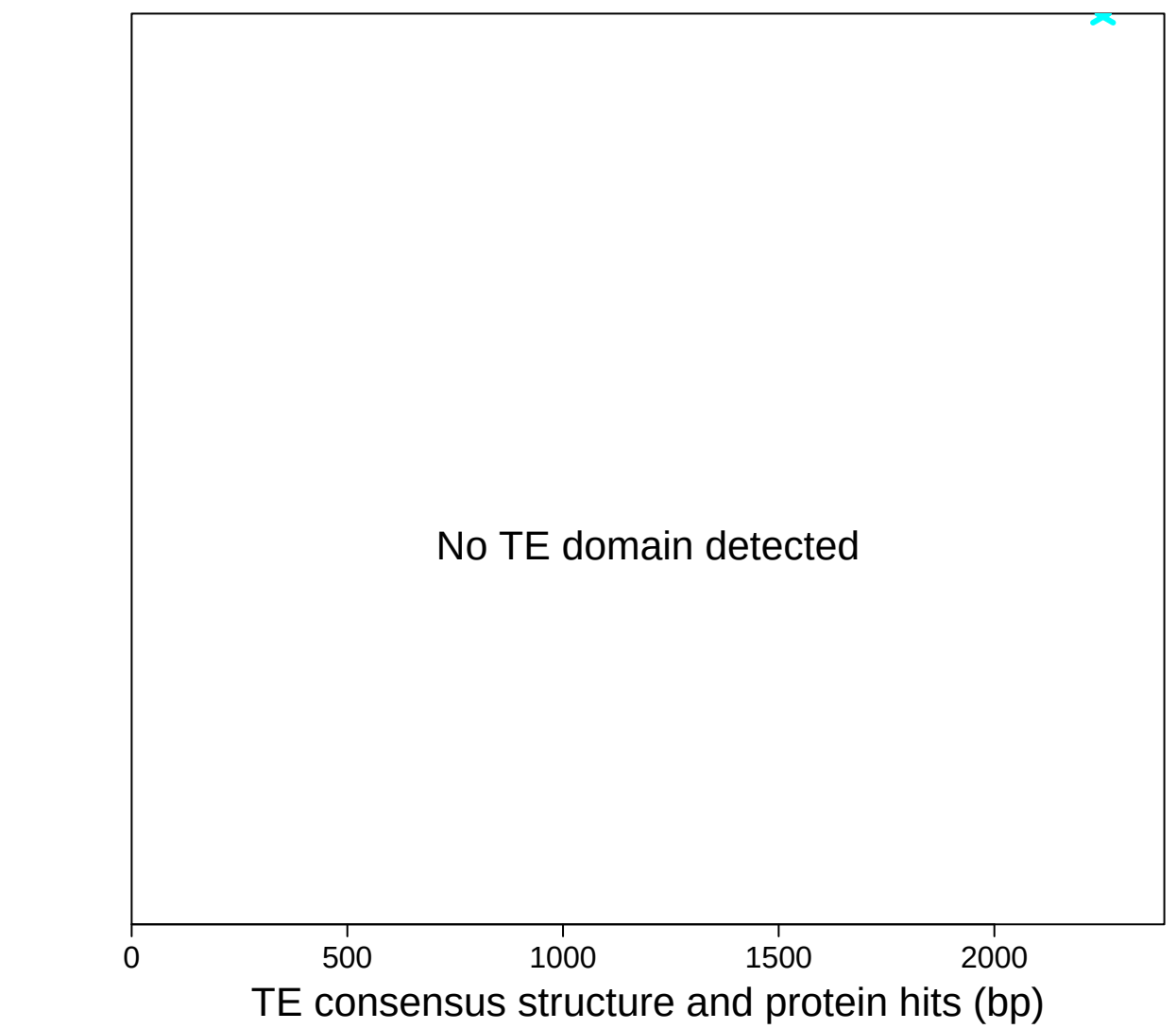
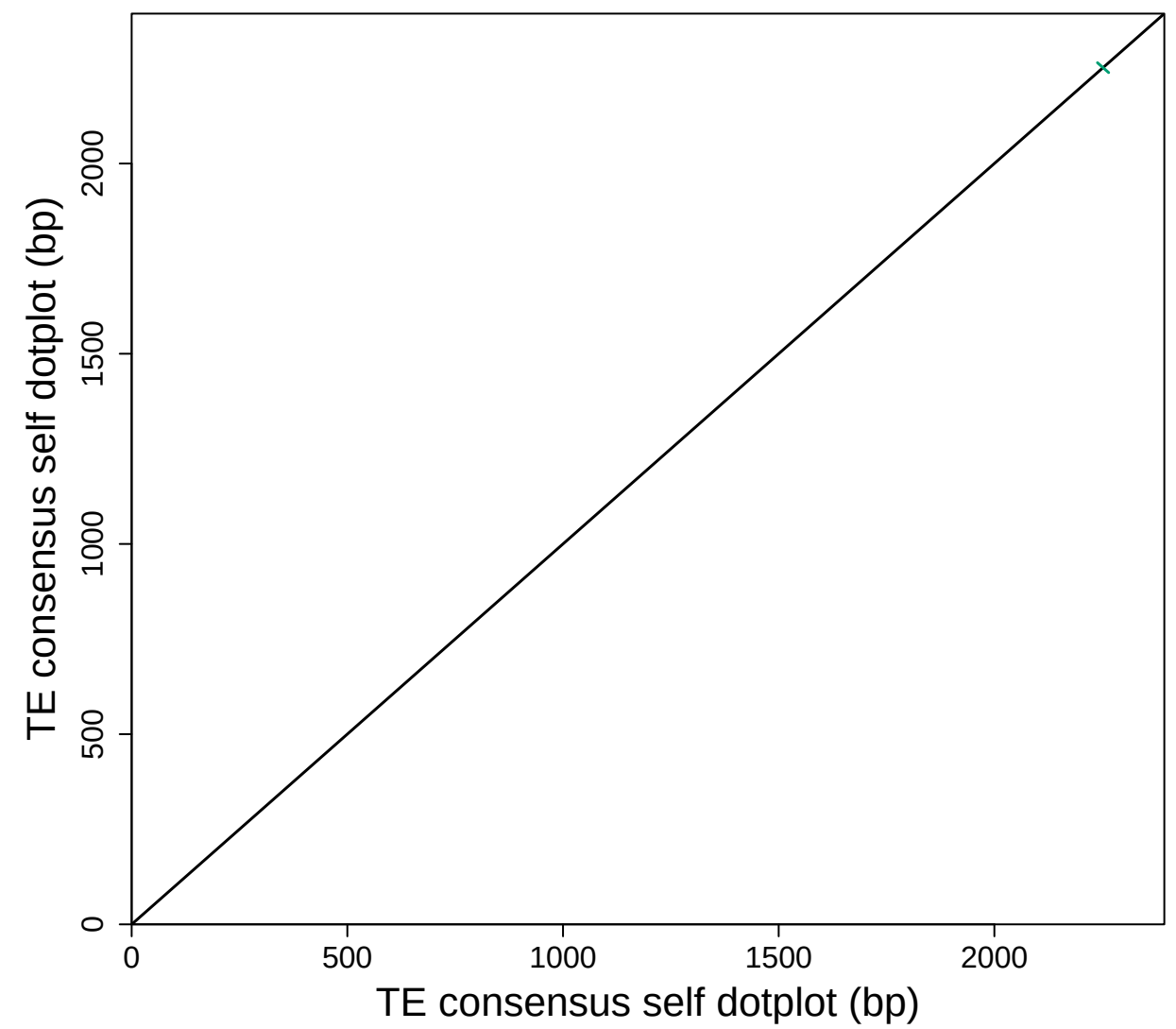
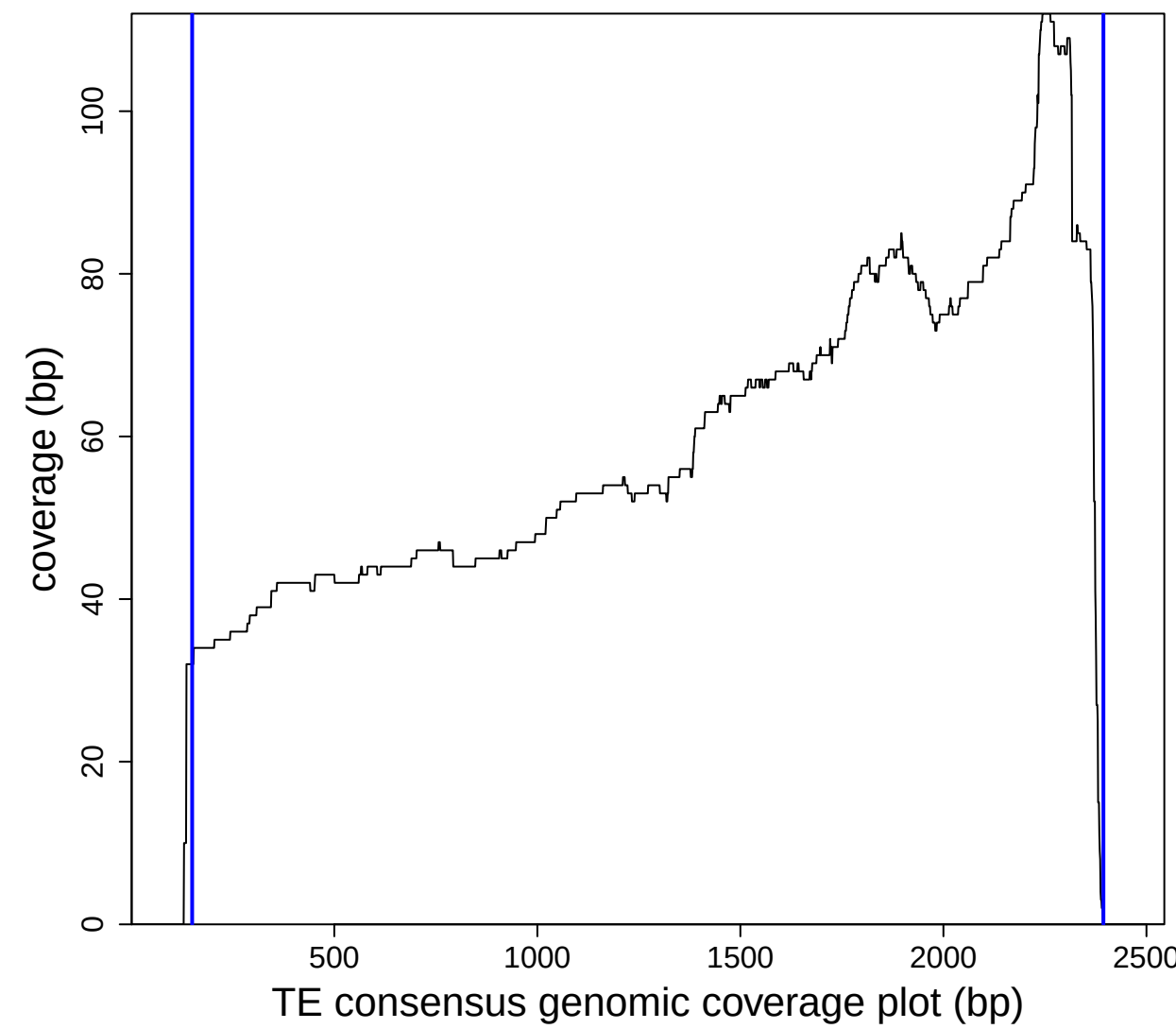
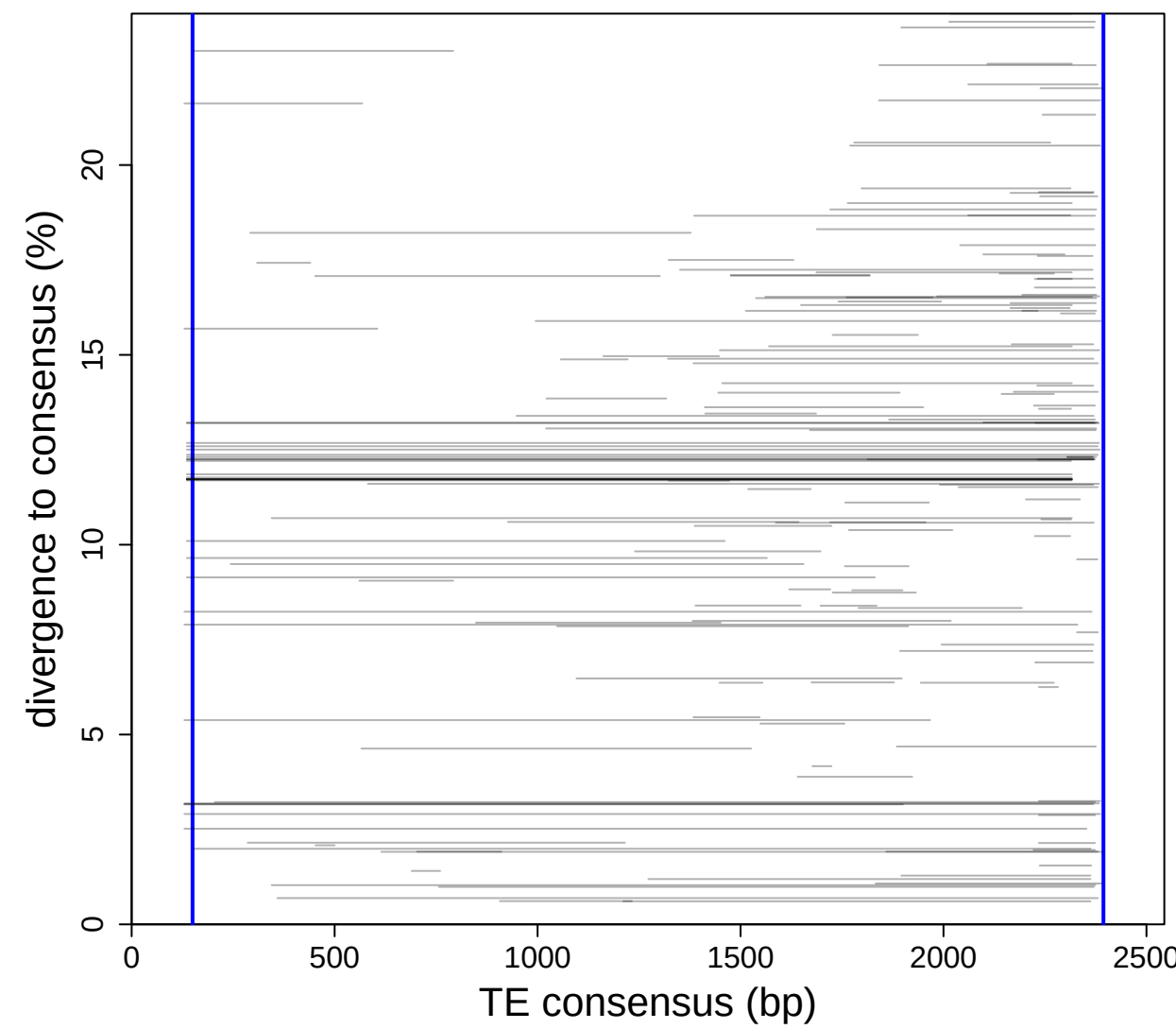
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 2241bp; fragments: 205; full length: 30 (>=2016.9bp)

After TETrimmer 2241 bp

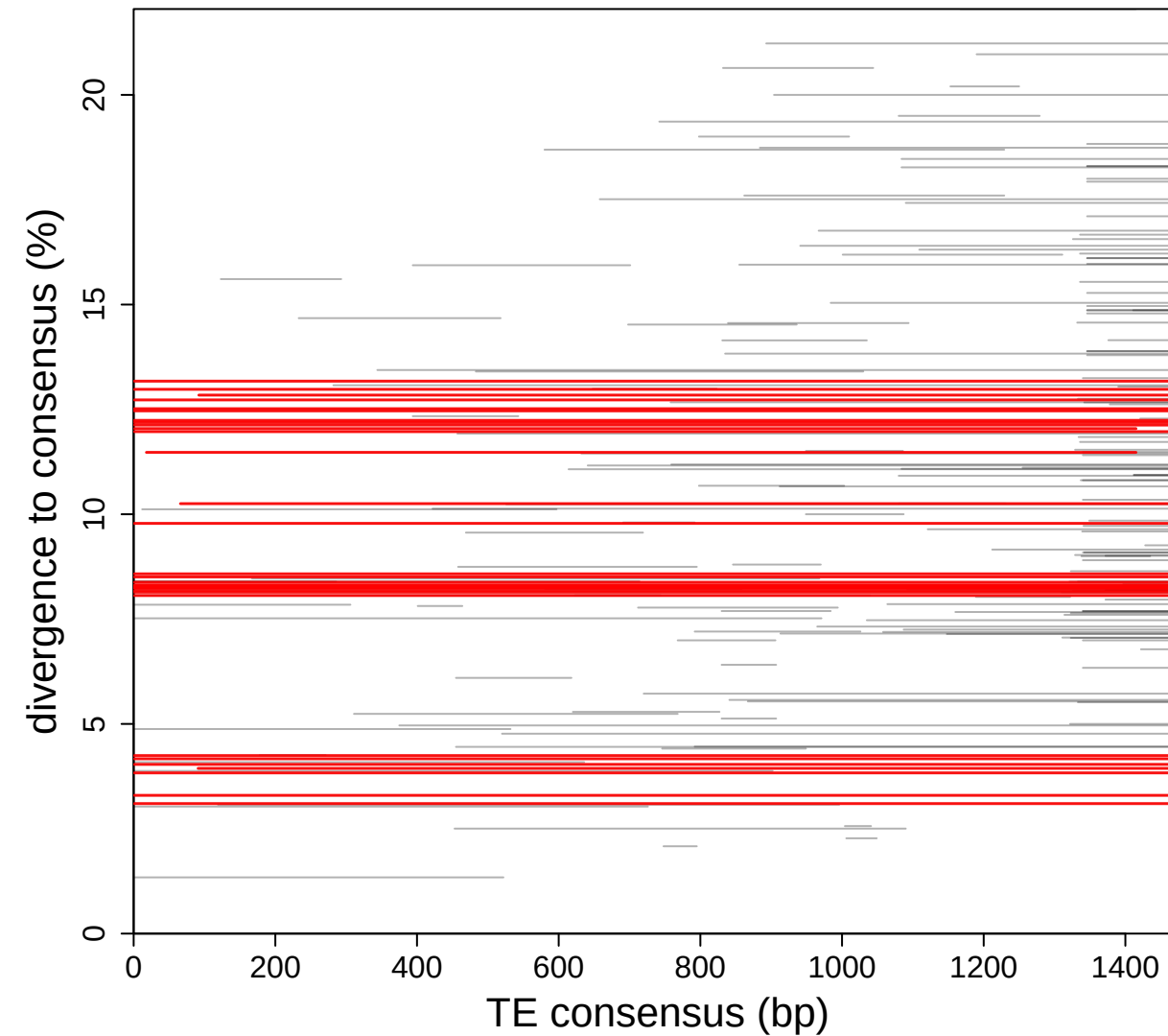


7.fasta.b.bed uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 2544bp; fragments: 175; full length: 0 (>=2289.6bp)

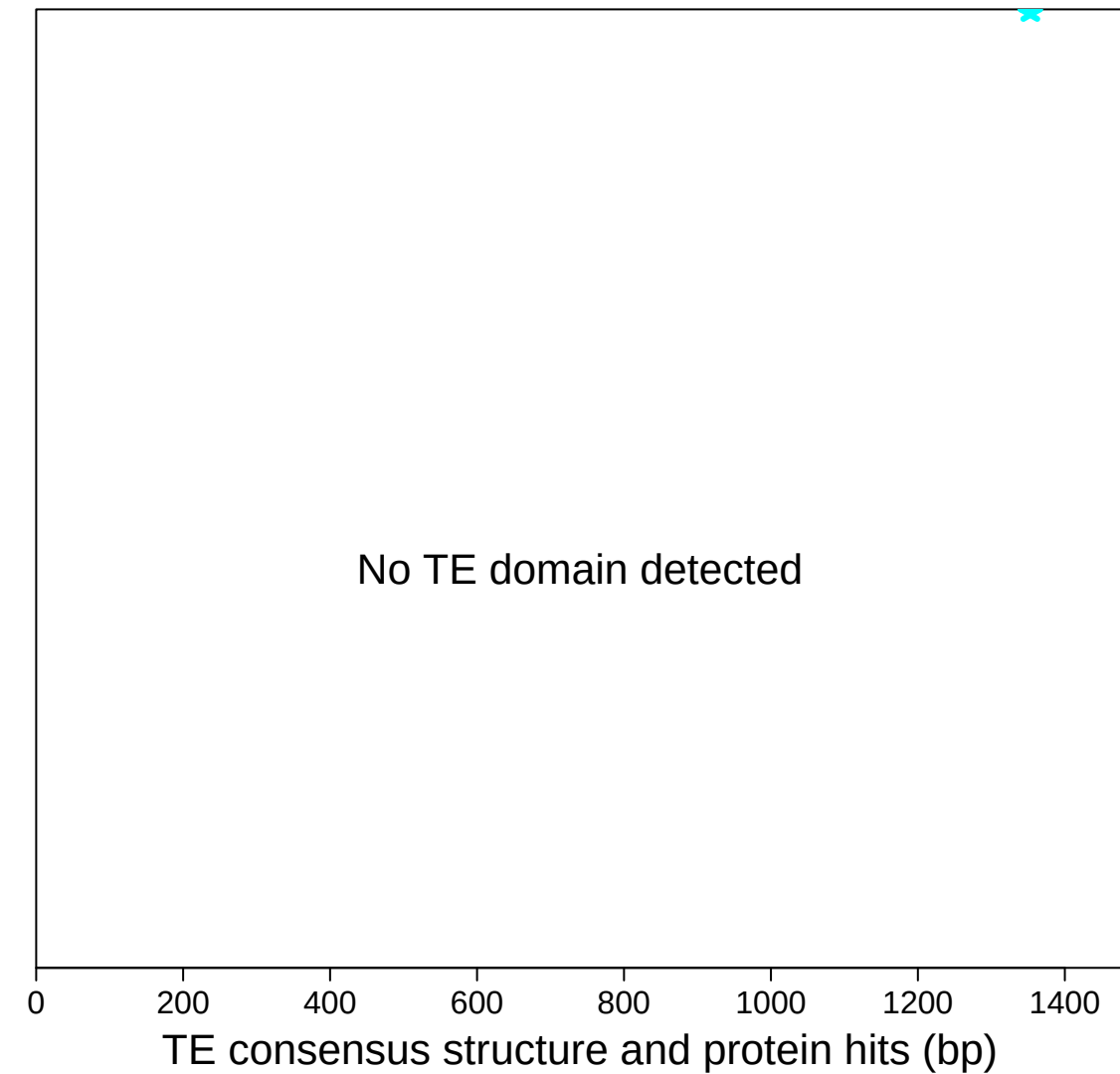
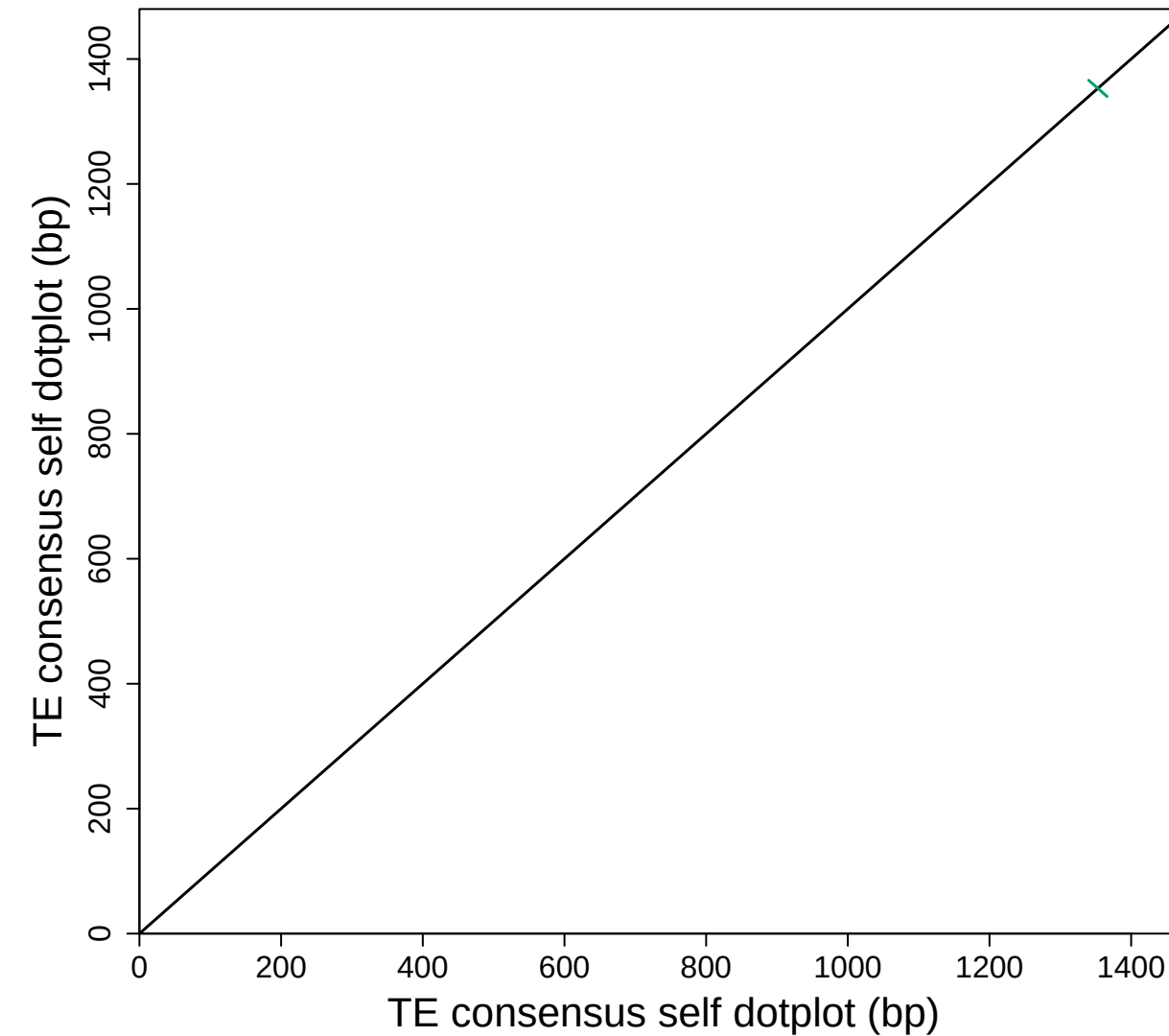
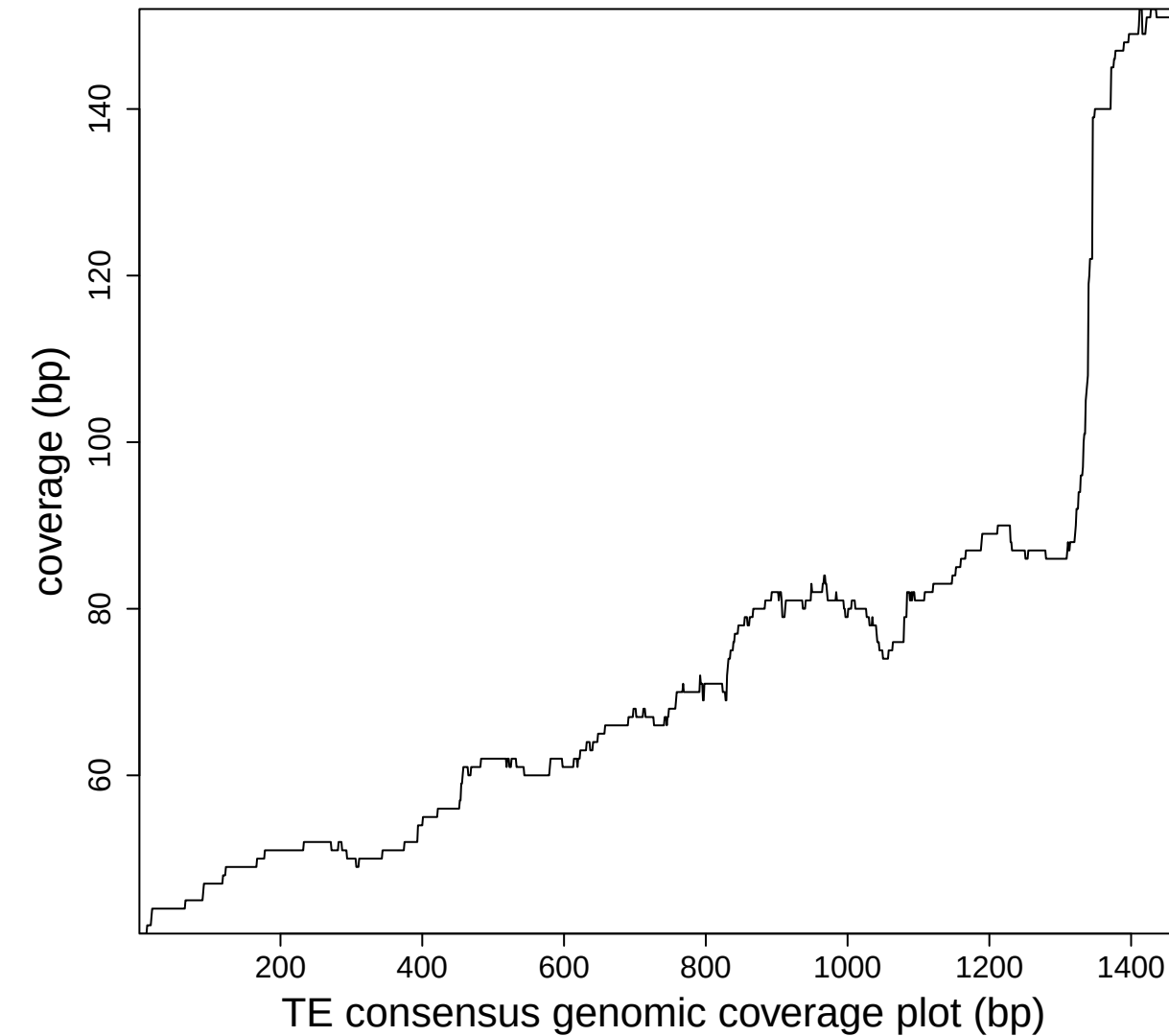
After TEtrimmer Extended plot Blue lines are boundaries



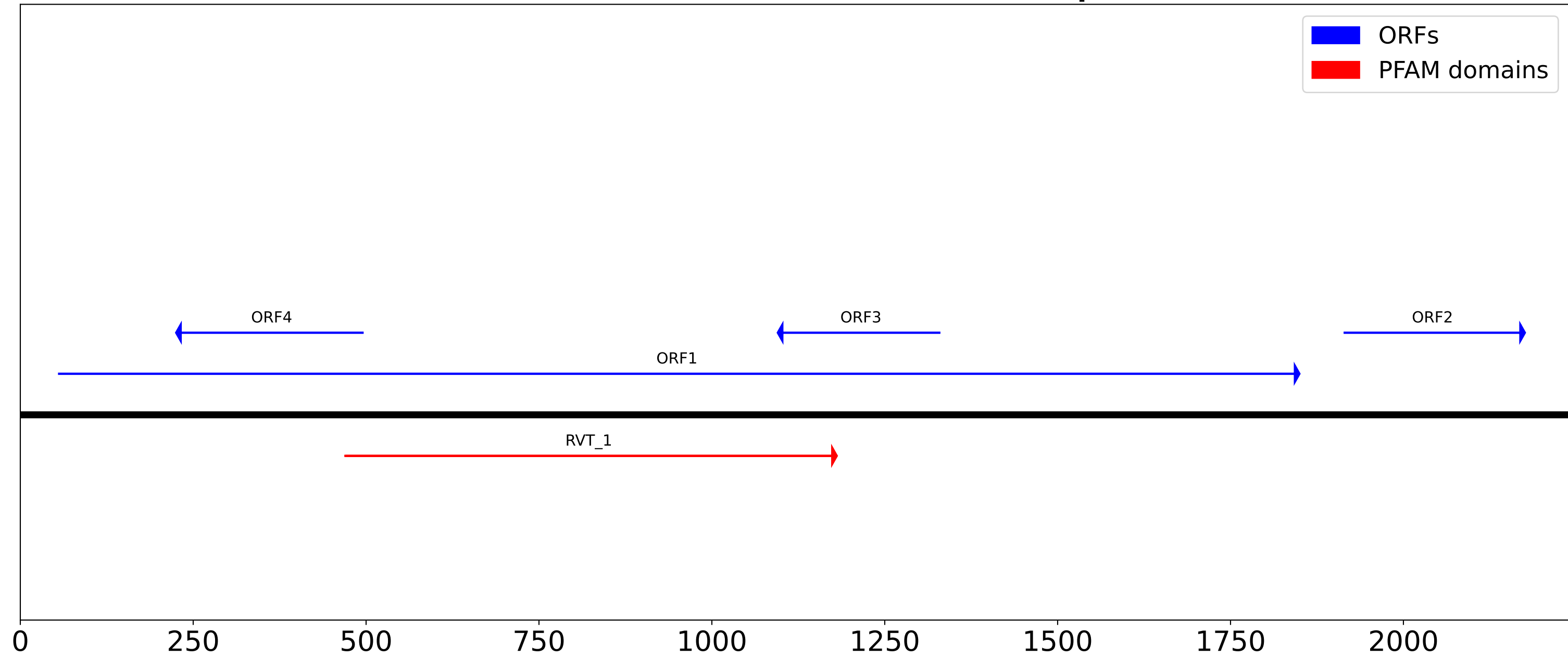
TE: rnd_5_family_9647
size: 1480bp; fragments: 211; full length: 36 (≥ 1332 bp)



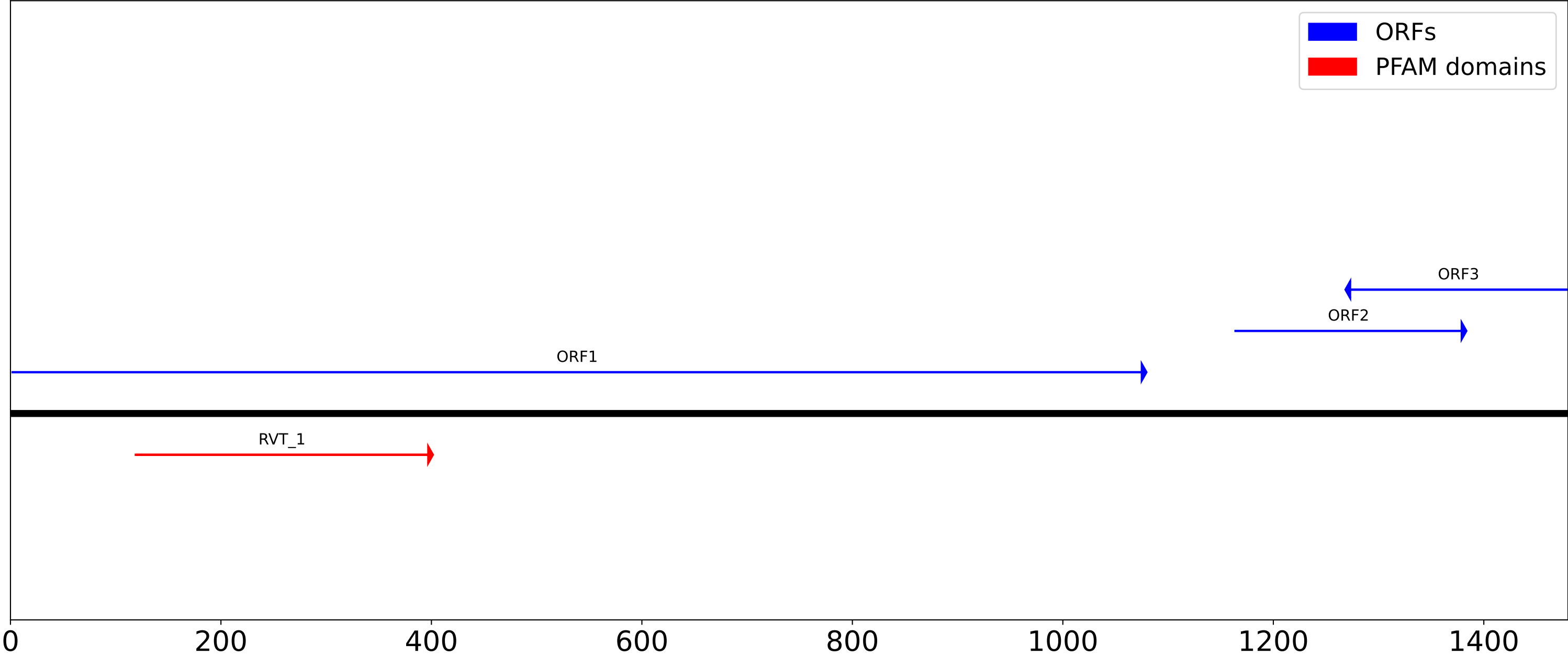
Before TEtrimmer 1480 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

