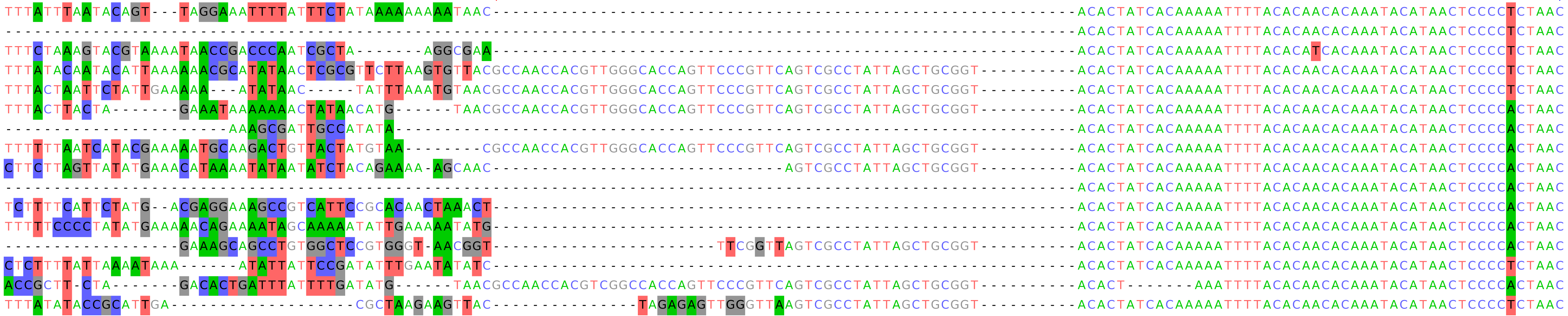


Start crop Point

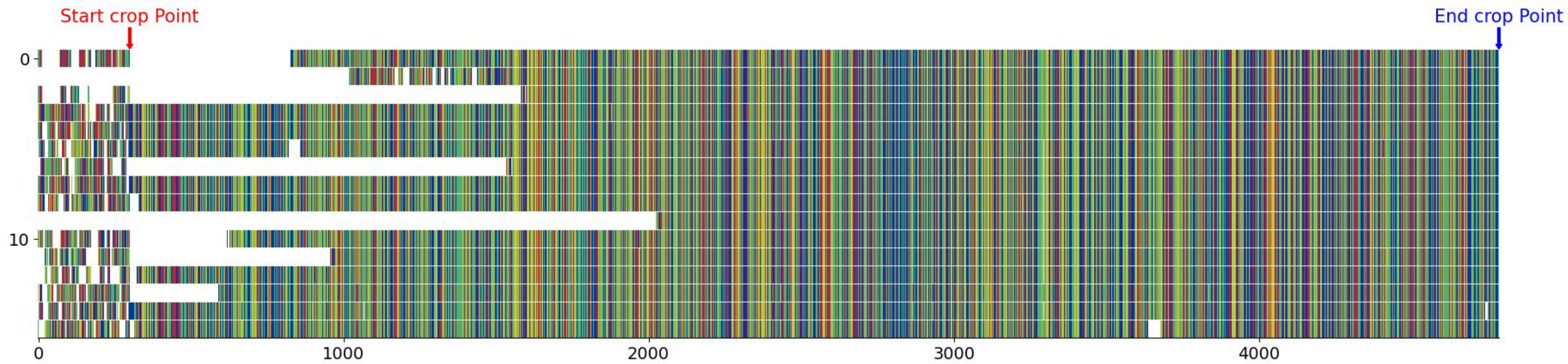
End crop Point

MSA length = 4488



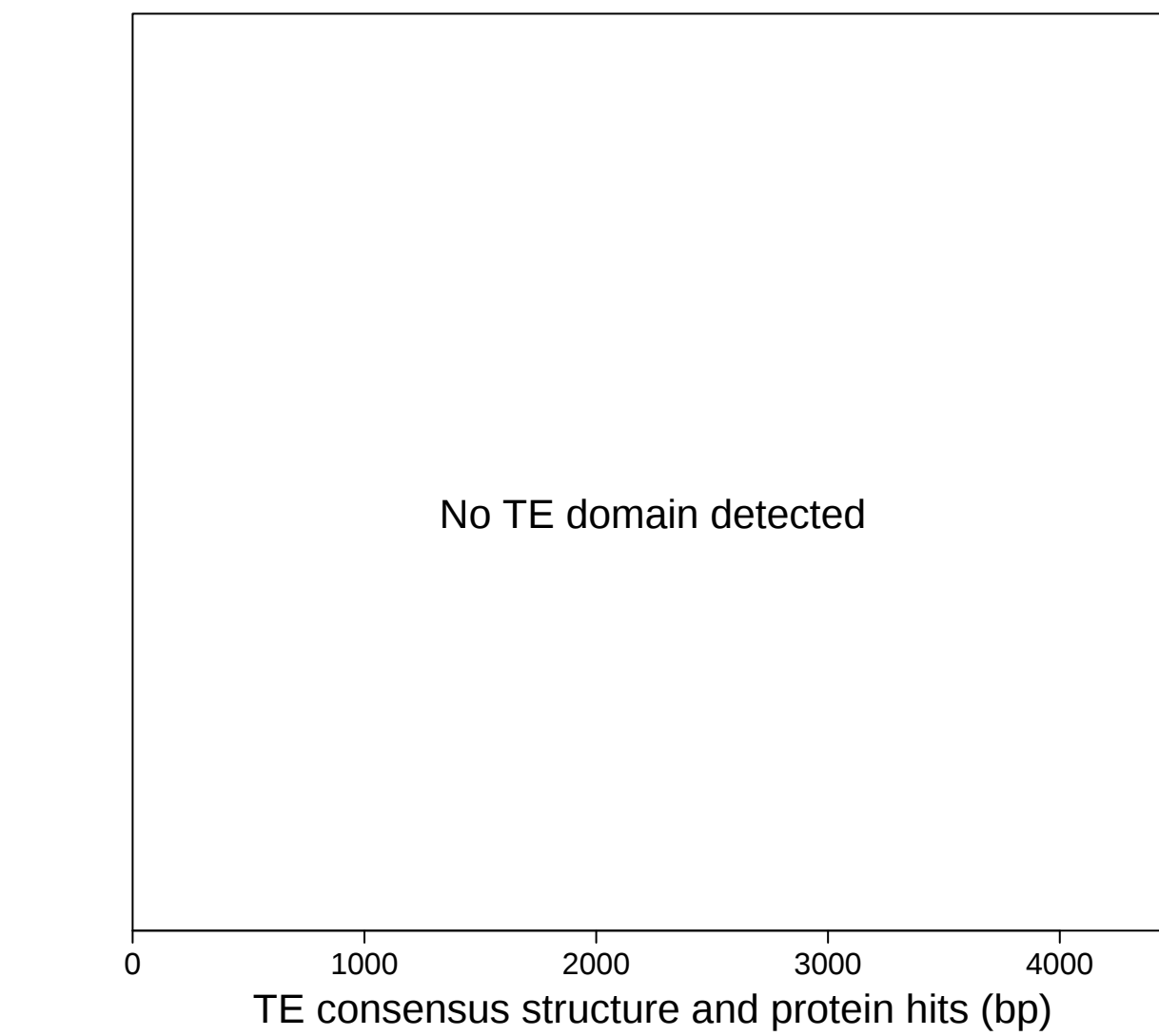
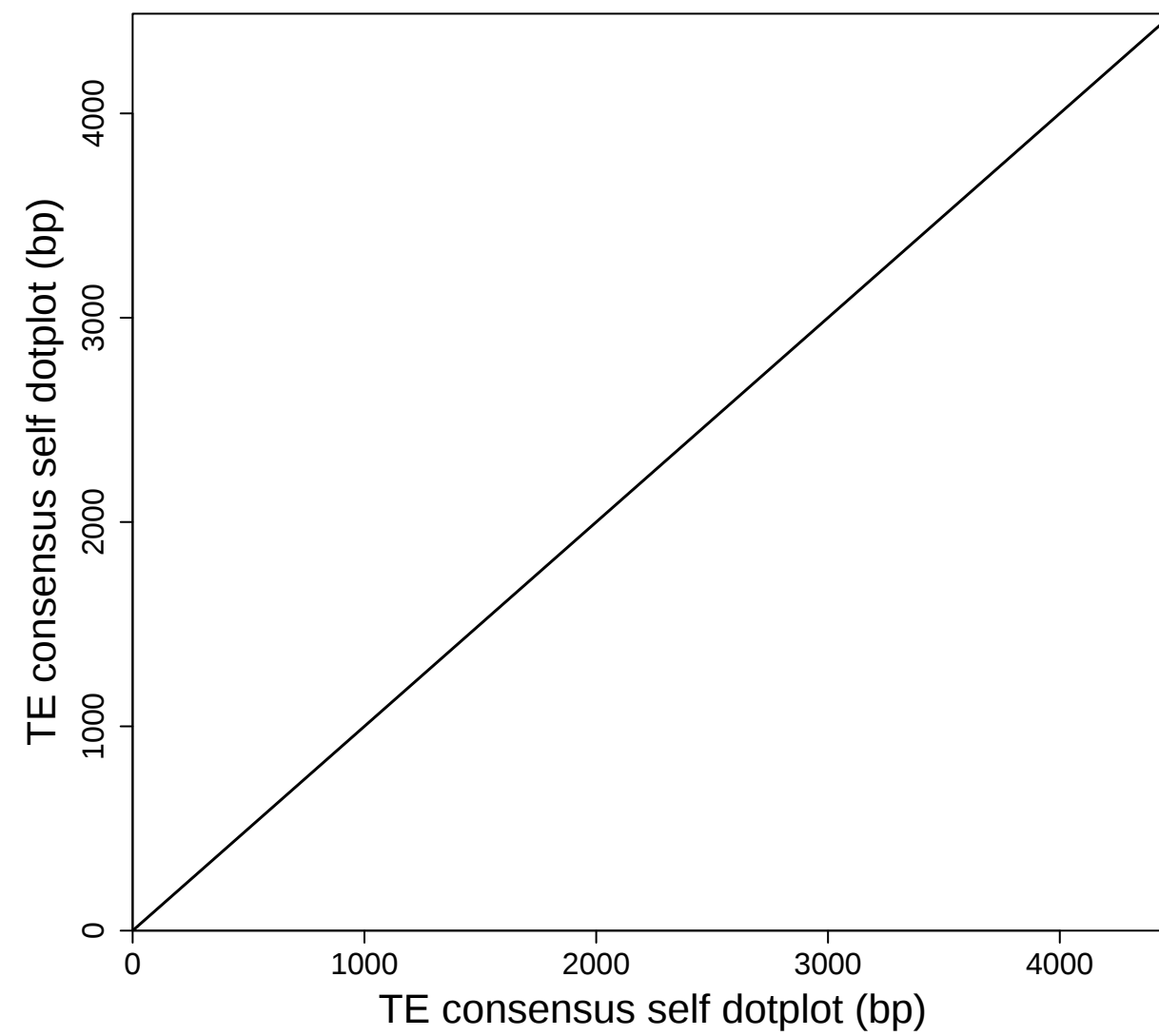
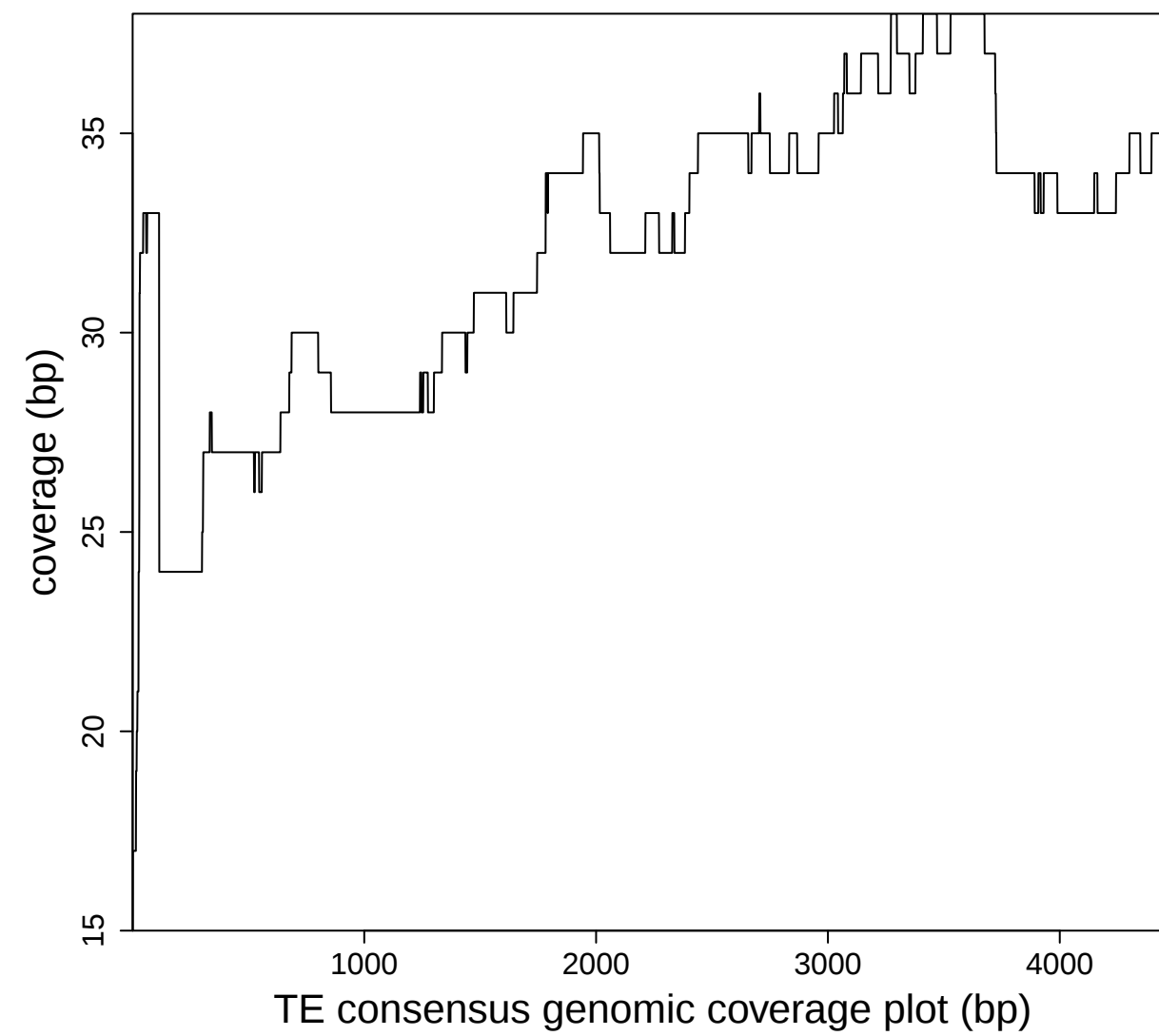
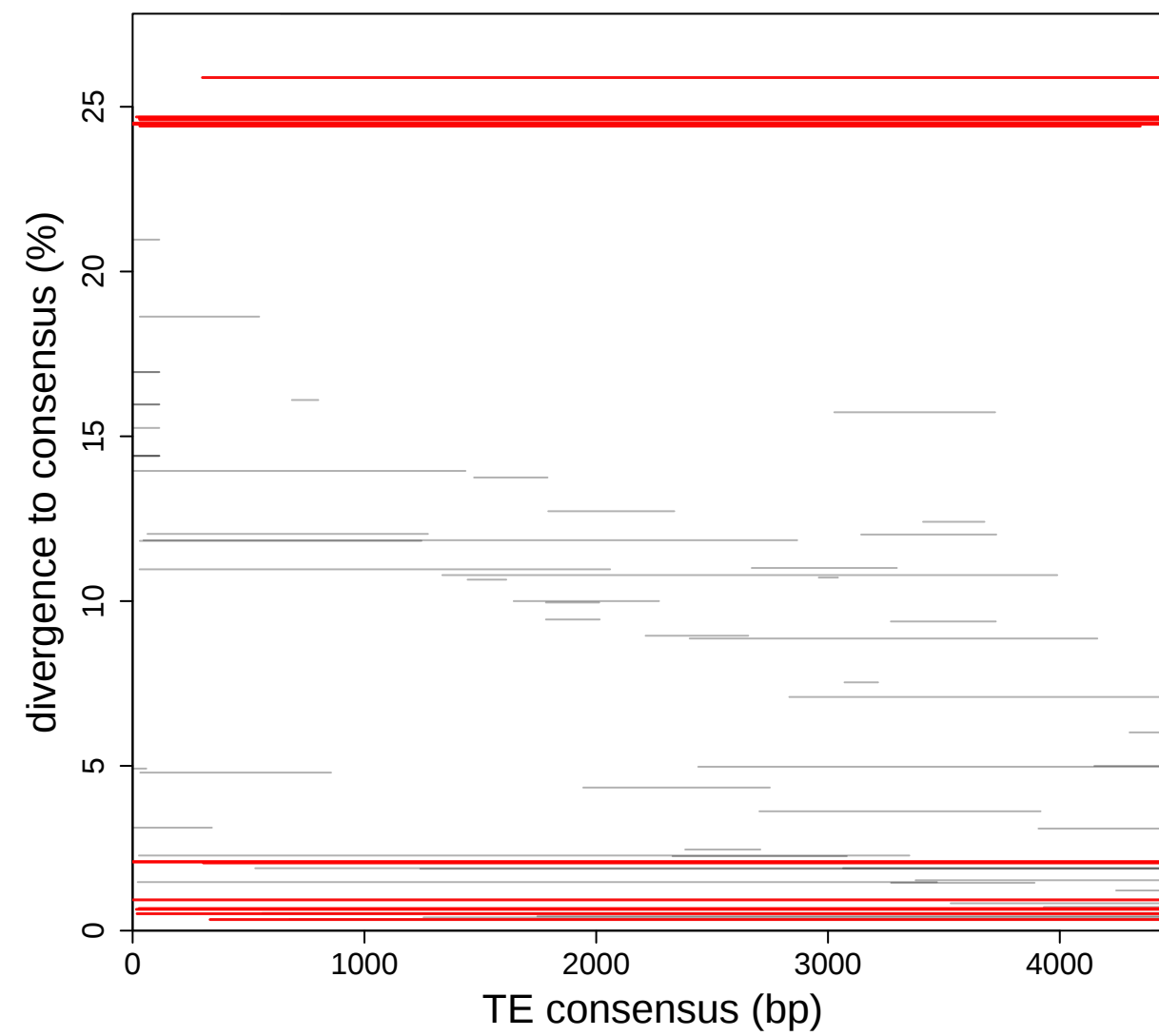
Start crop Point

End crop Point



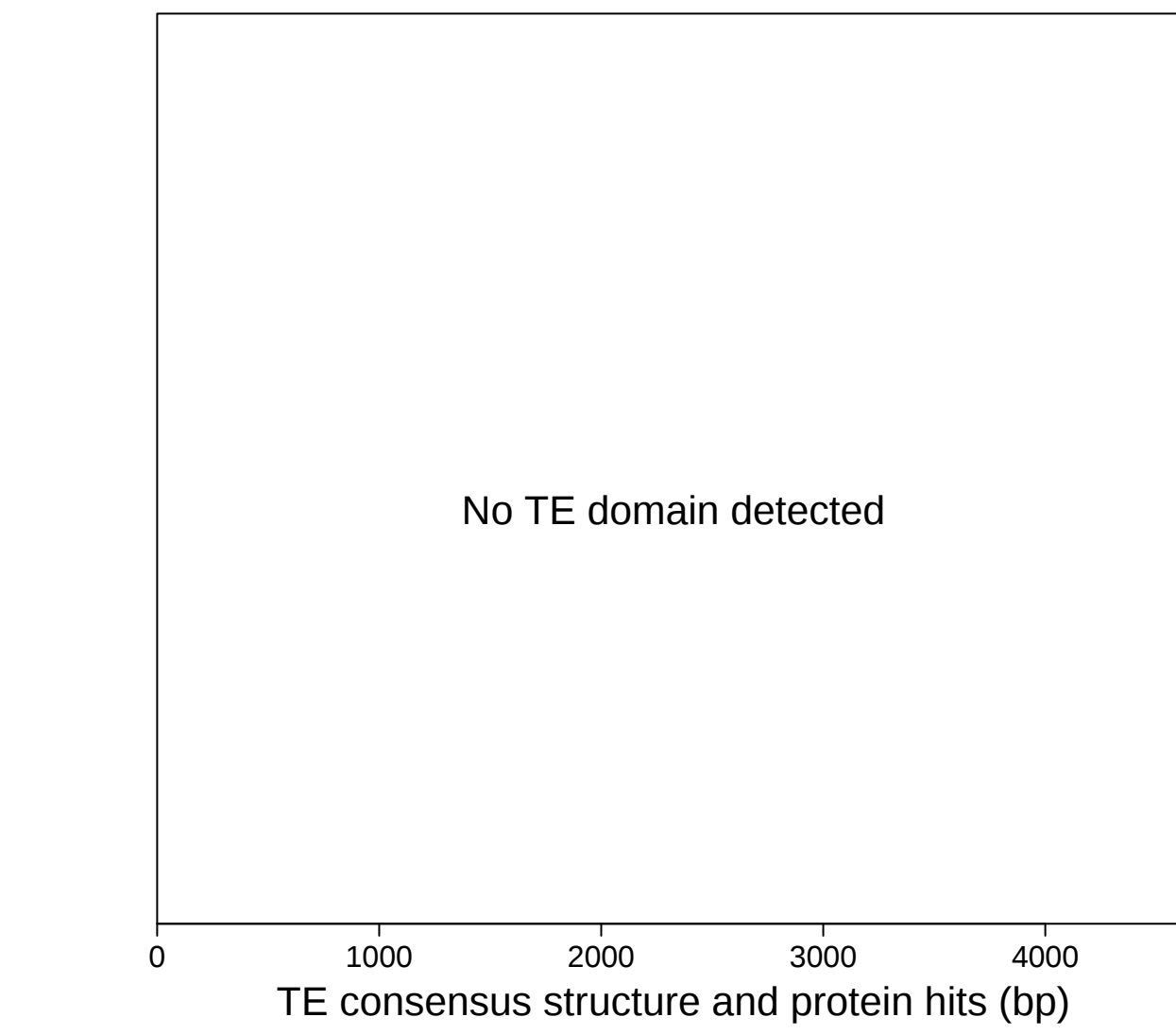
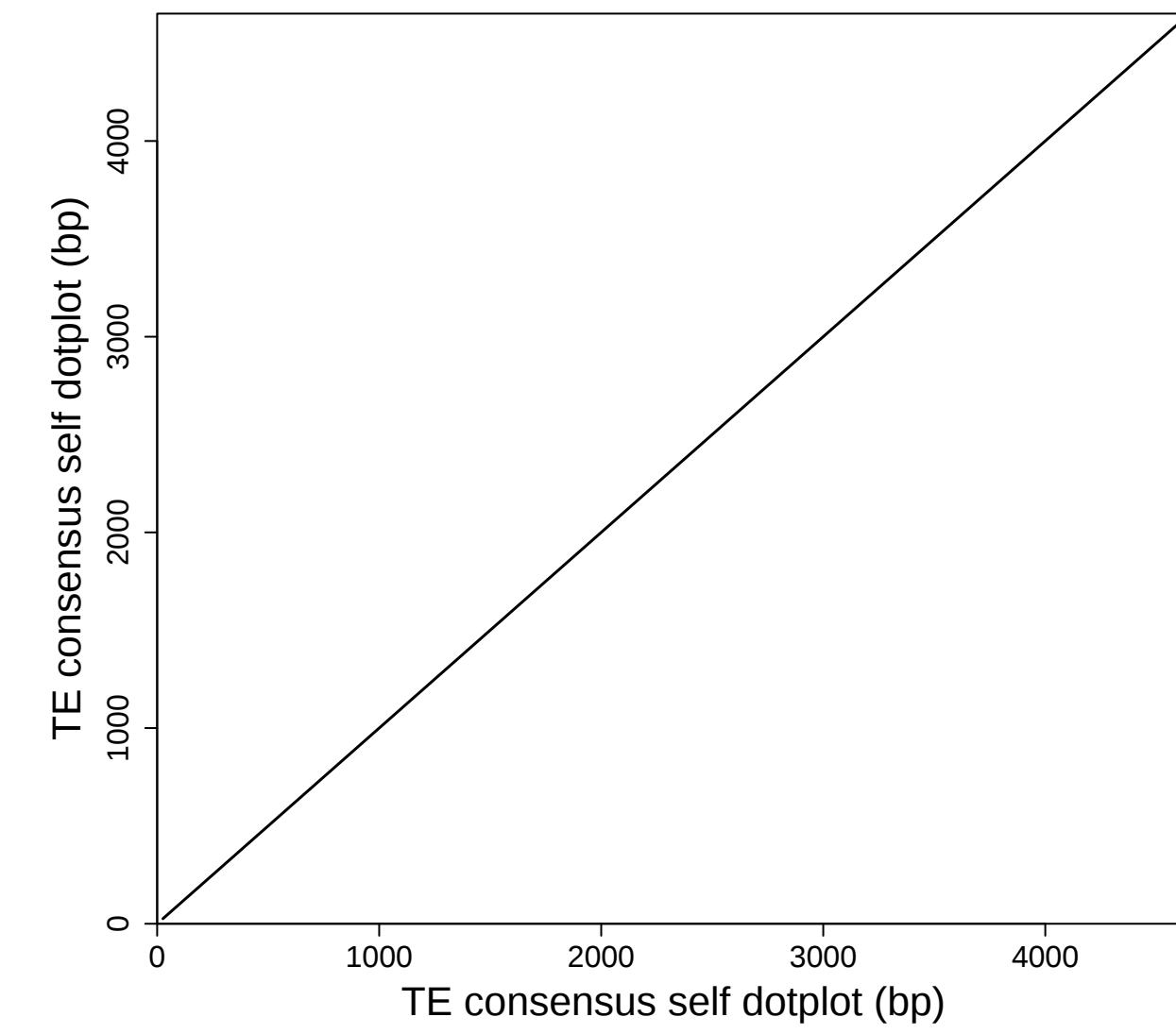
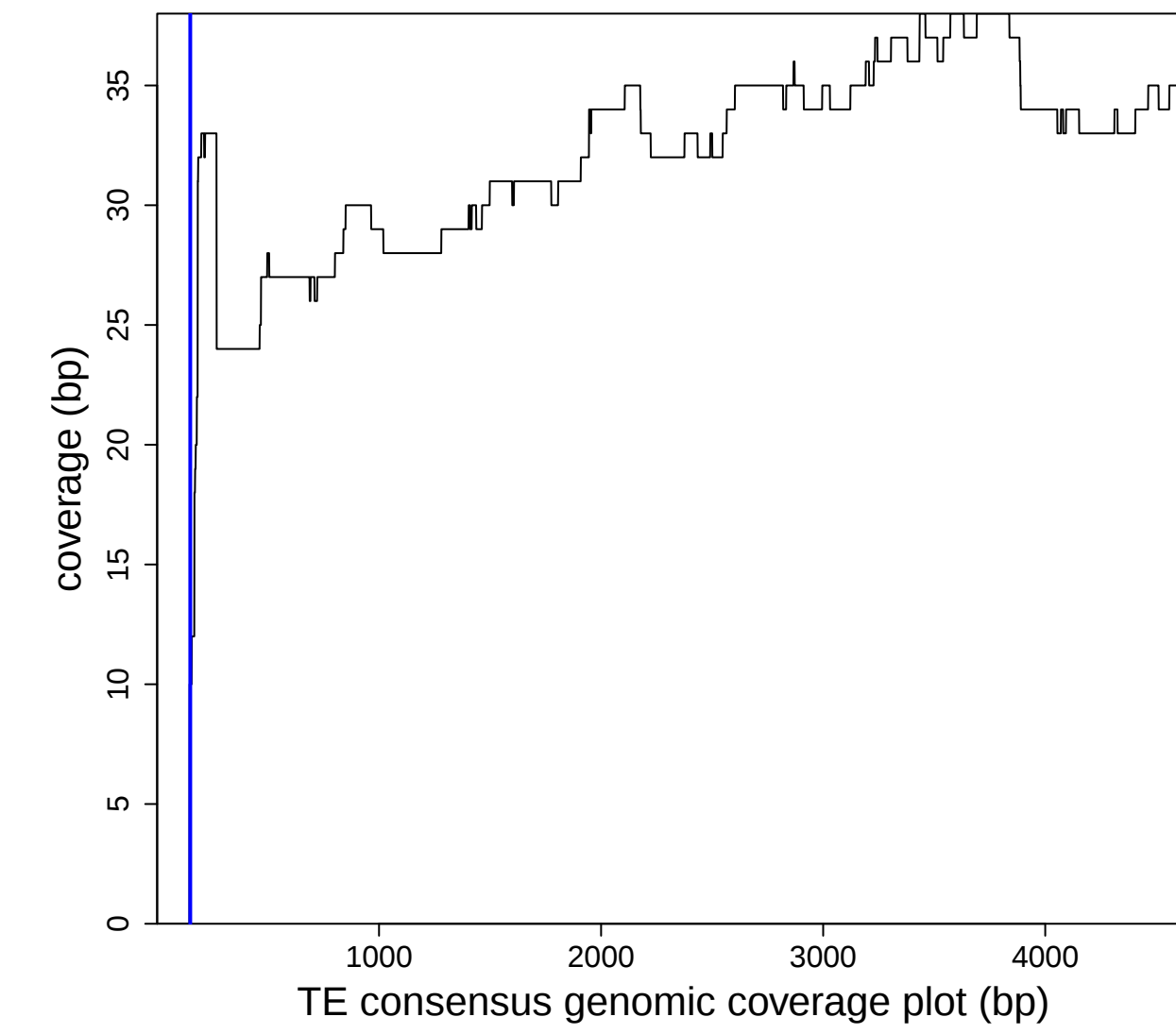
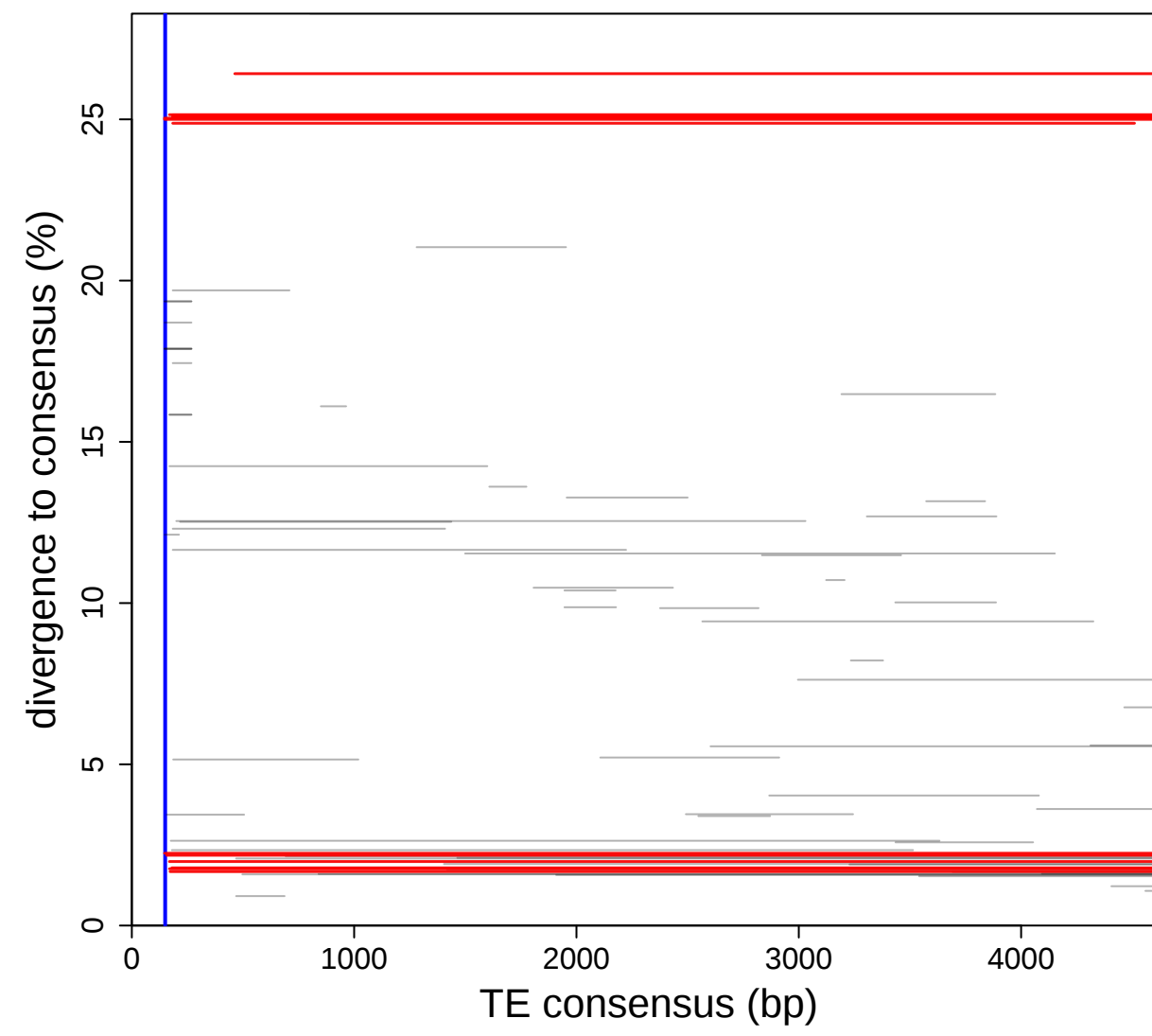
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 4488bp; fragments: 79; full length: 17 (>=4039.2bp)

After TEtrimmer 4488 bp



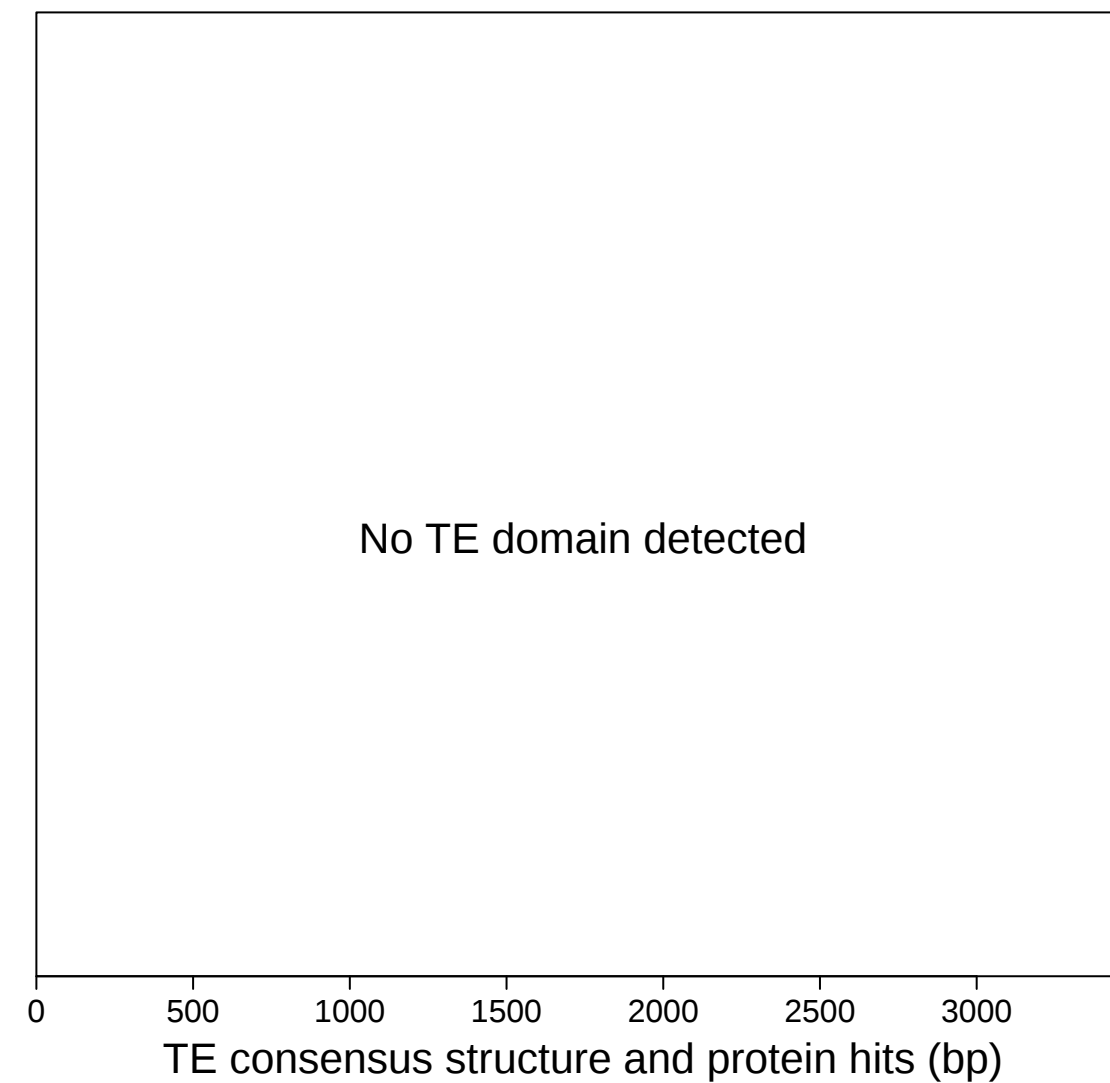
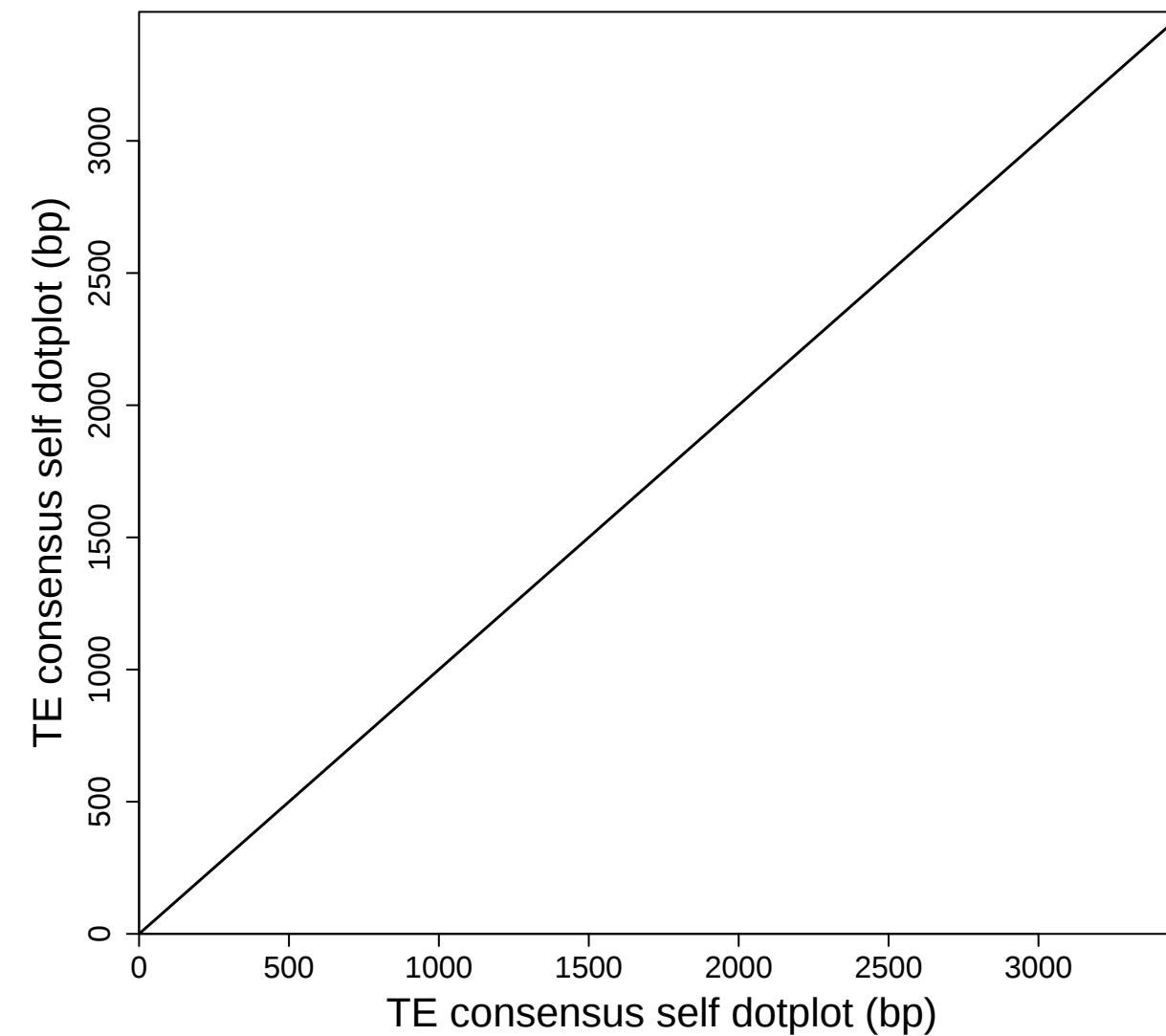
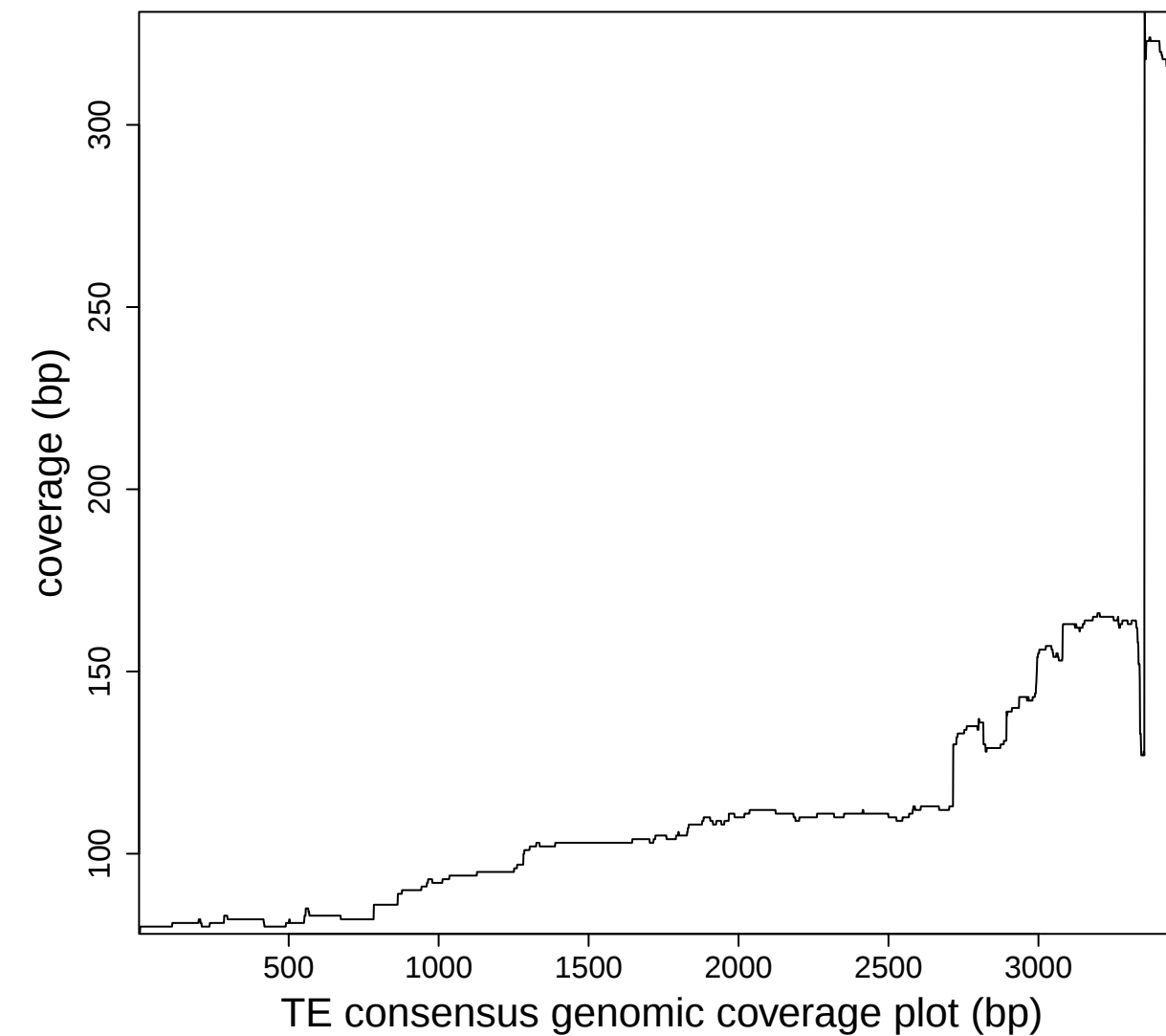
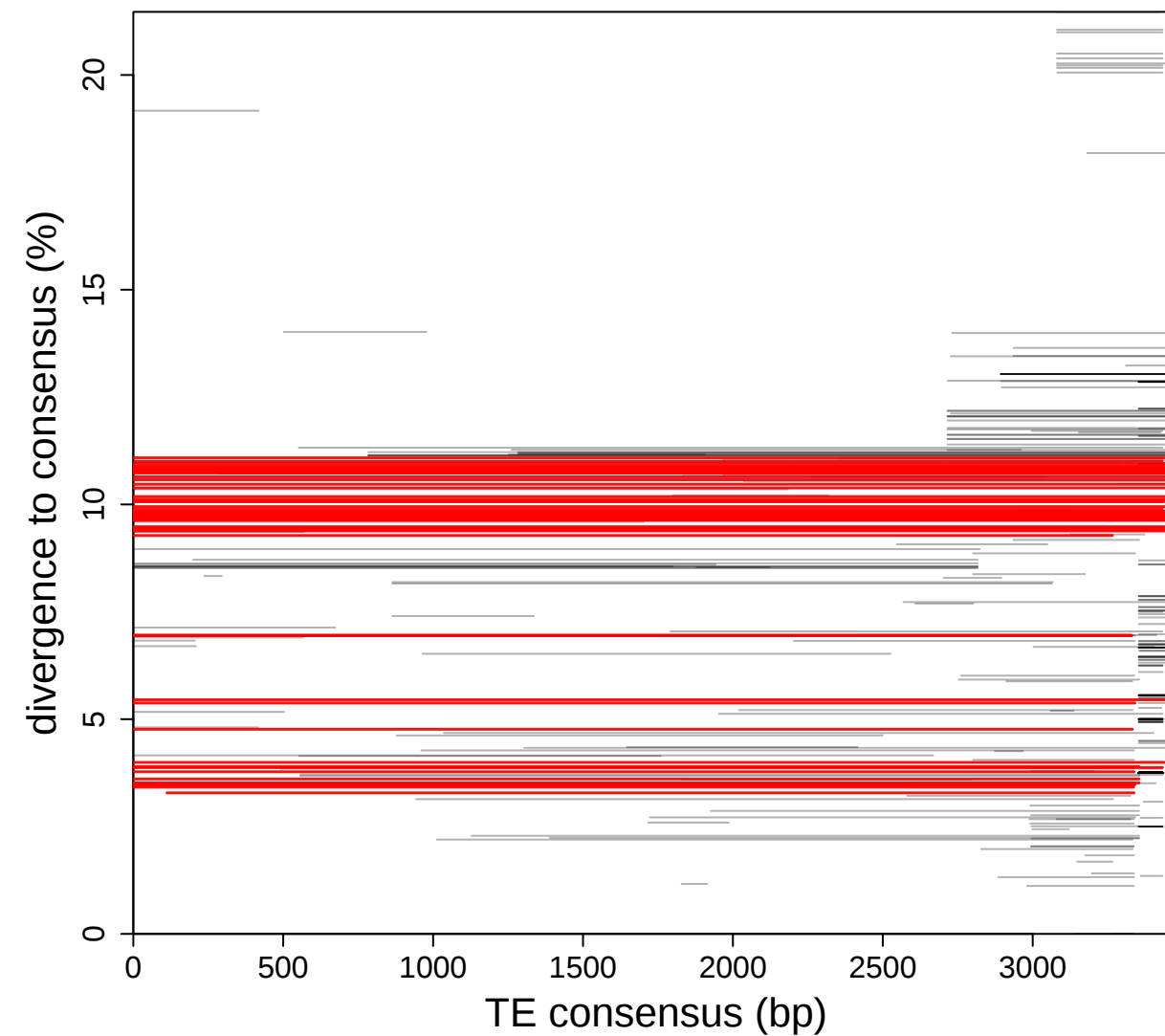
2.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 4651bp; fragments: 79; full length: 15 (>=4185.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

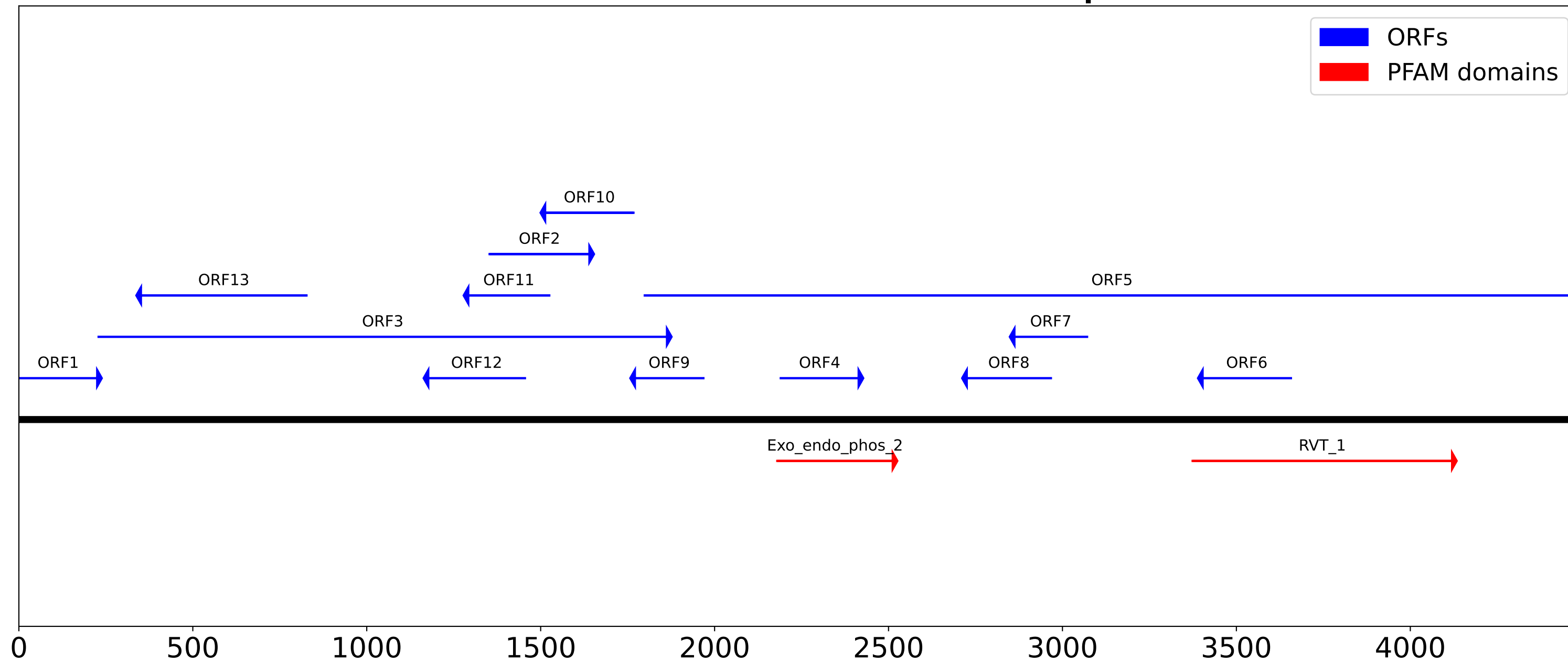


TE: rnd_5_family_3712
size: 3488bp; fragments: 430; full length: 62 (≥ 3139.2 bp)

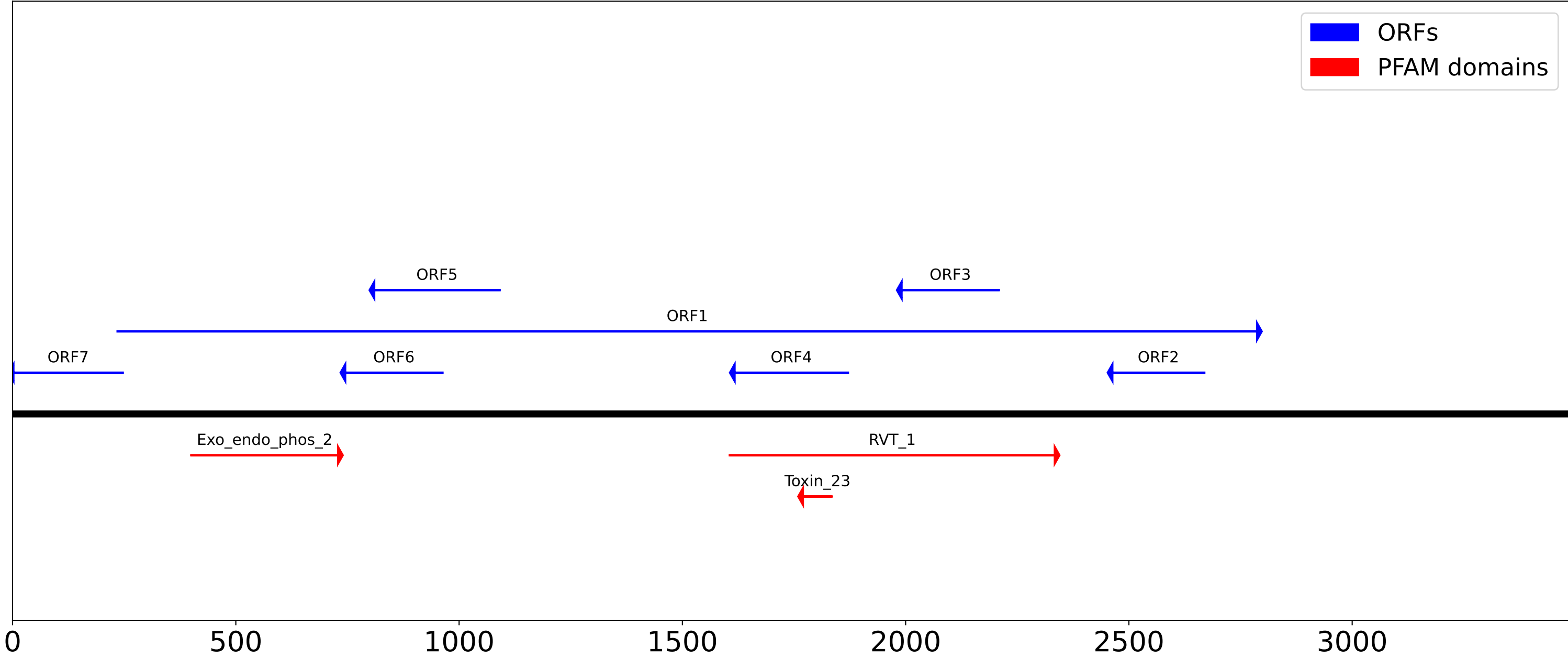
Before TEtrimmer 3488 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

