

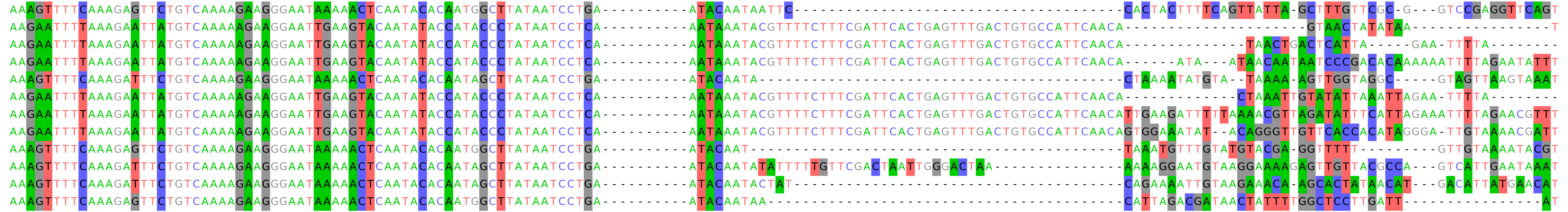
Start crop Point

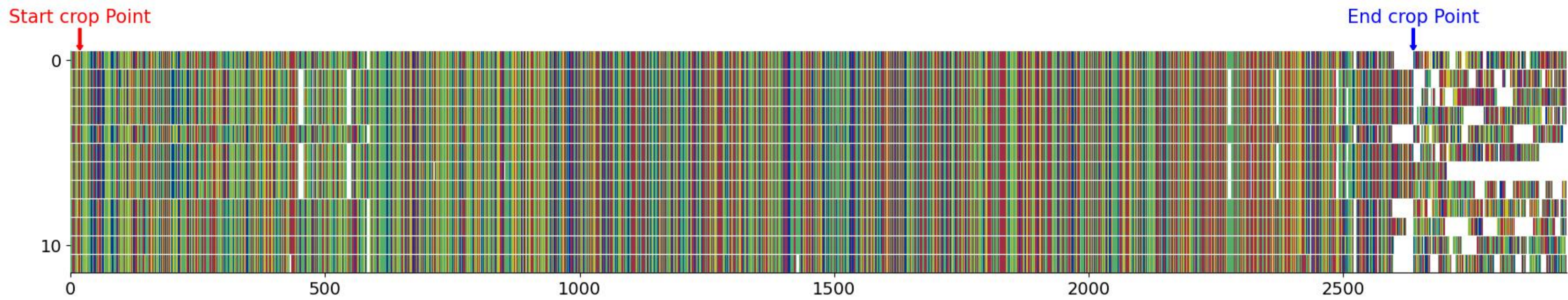


End crop Point



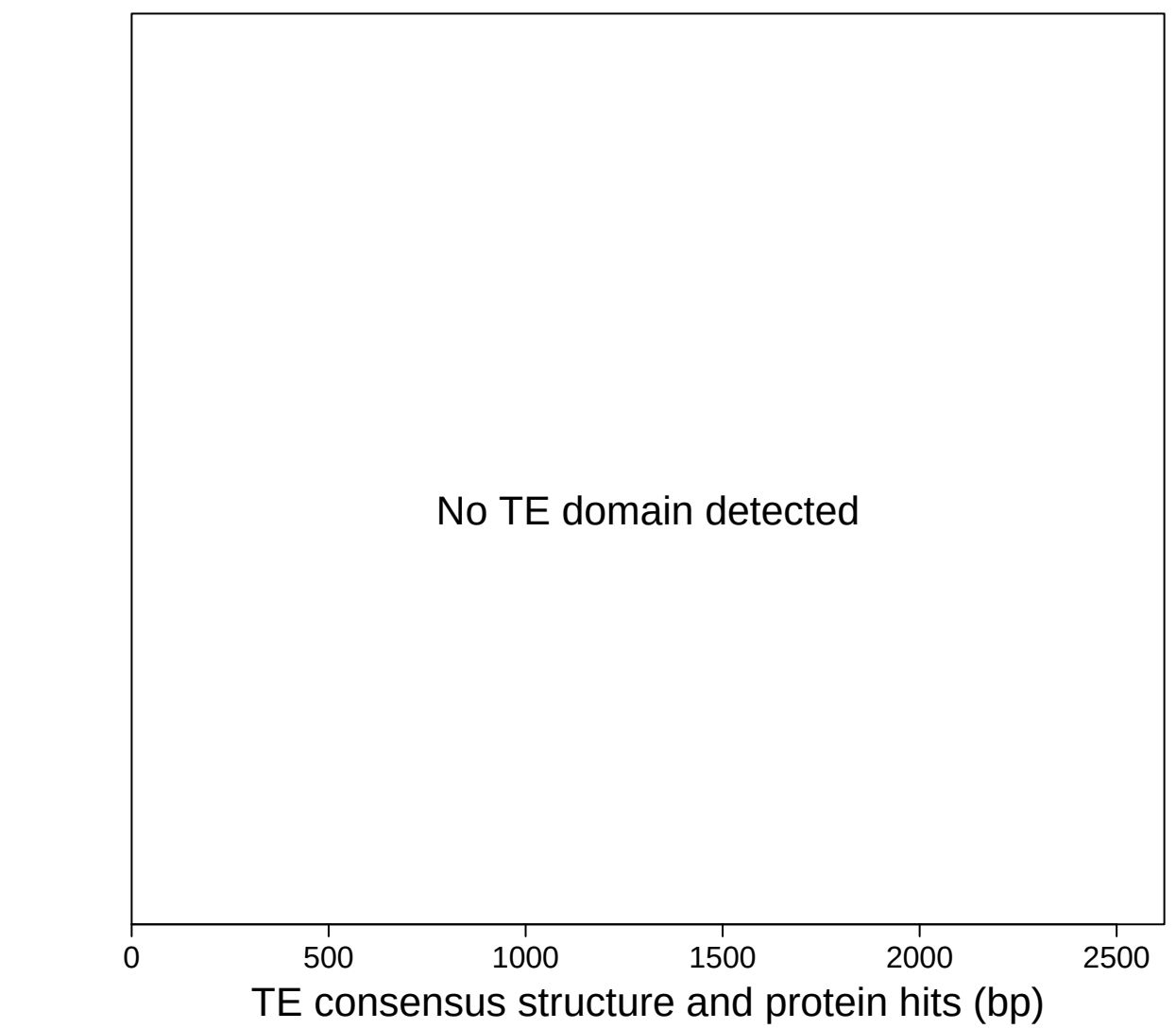
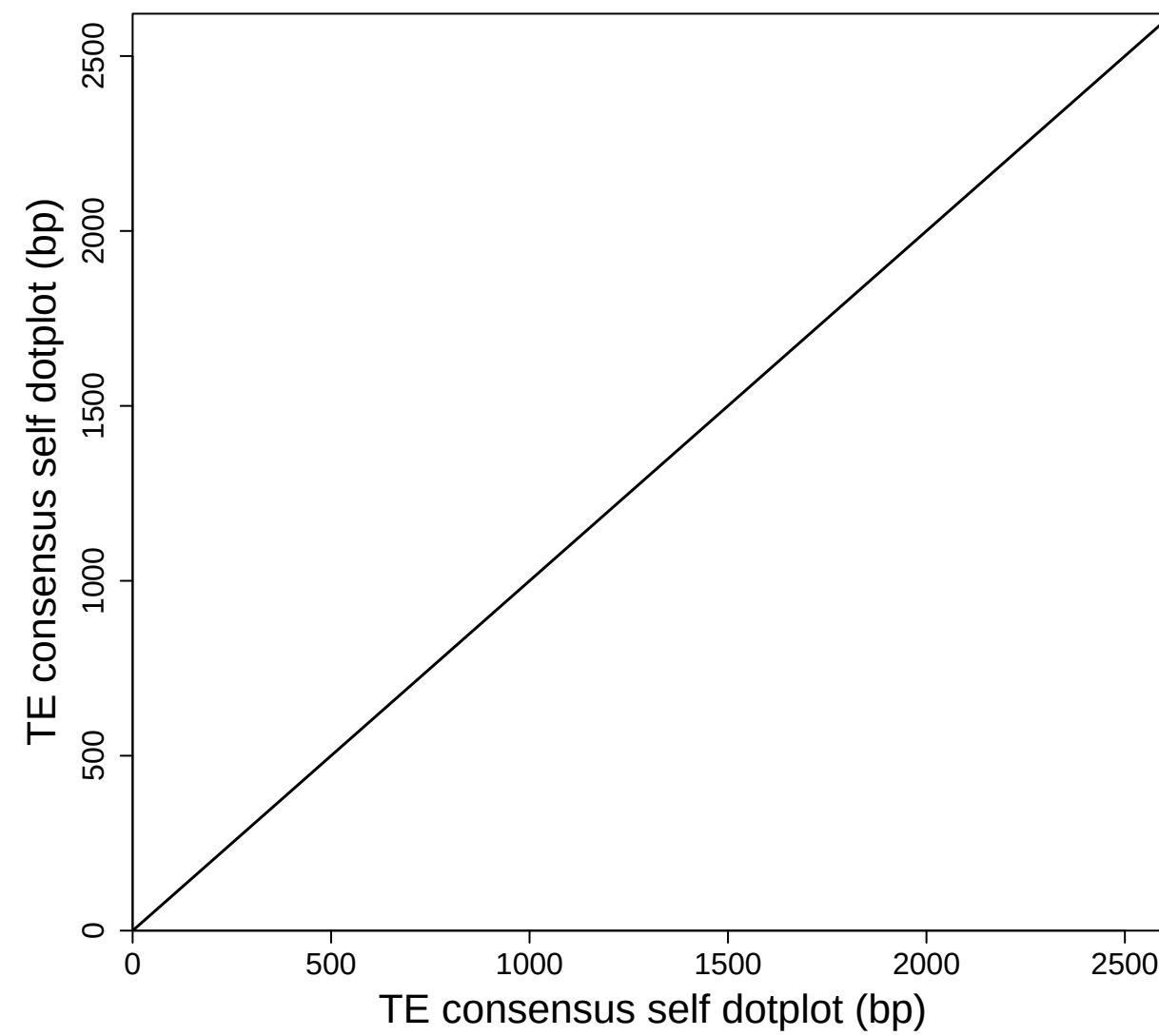
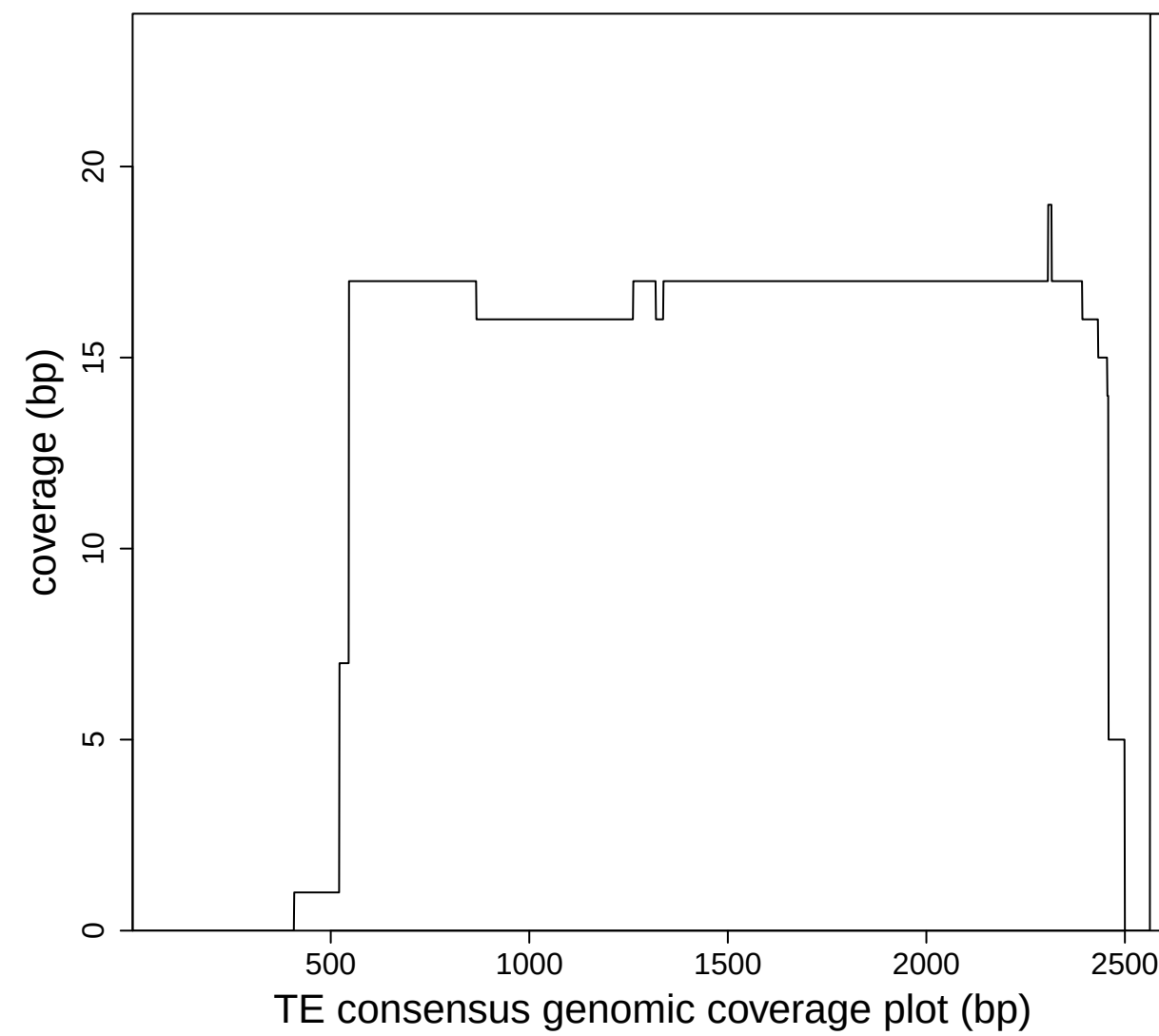
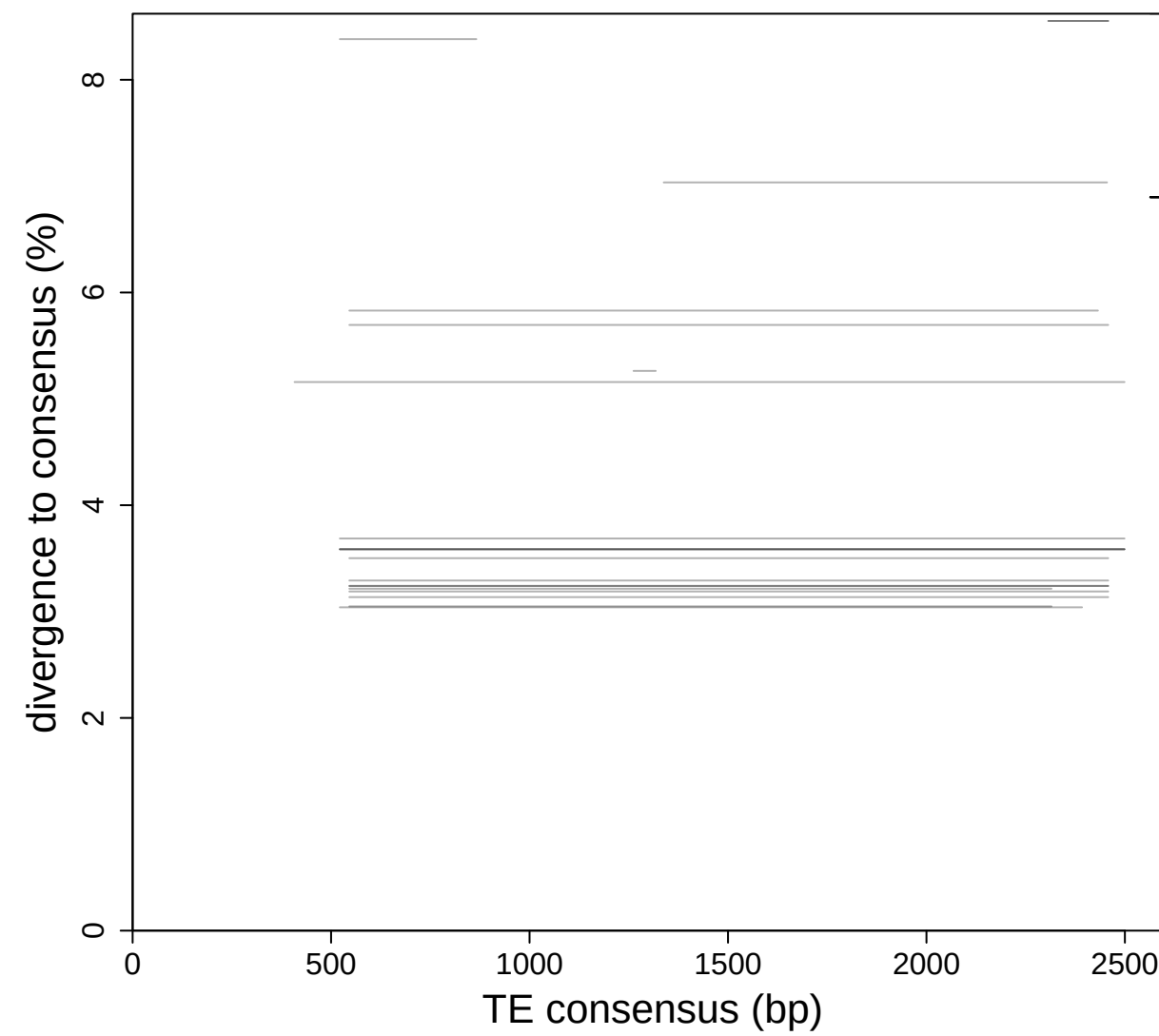
MSA length = 2621





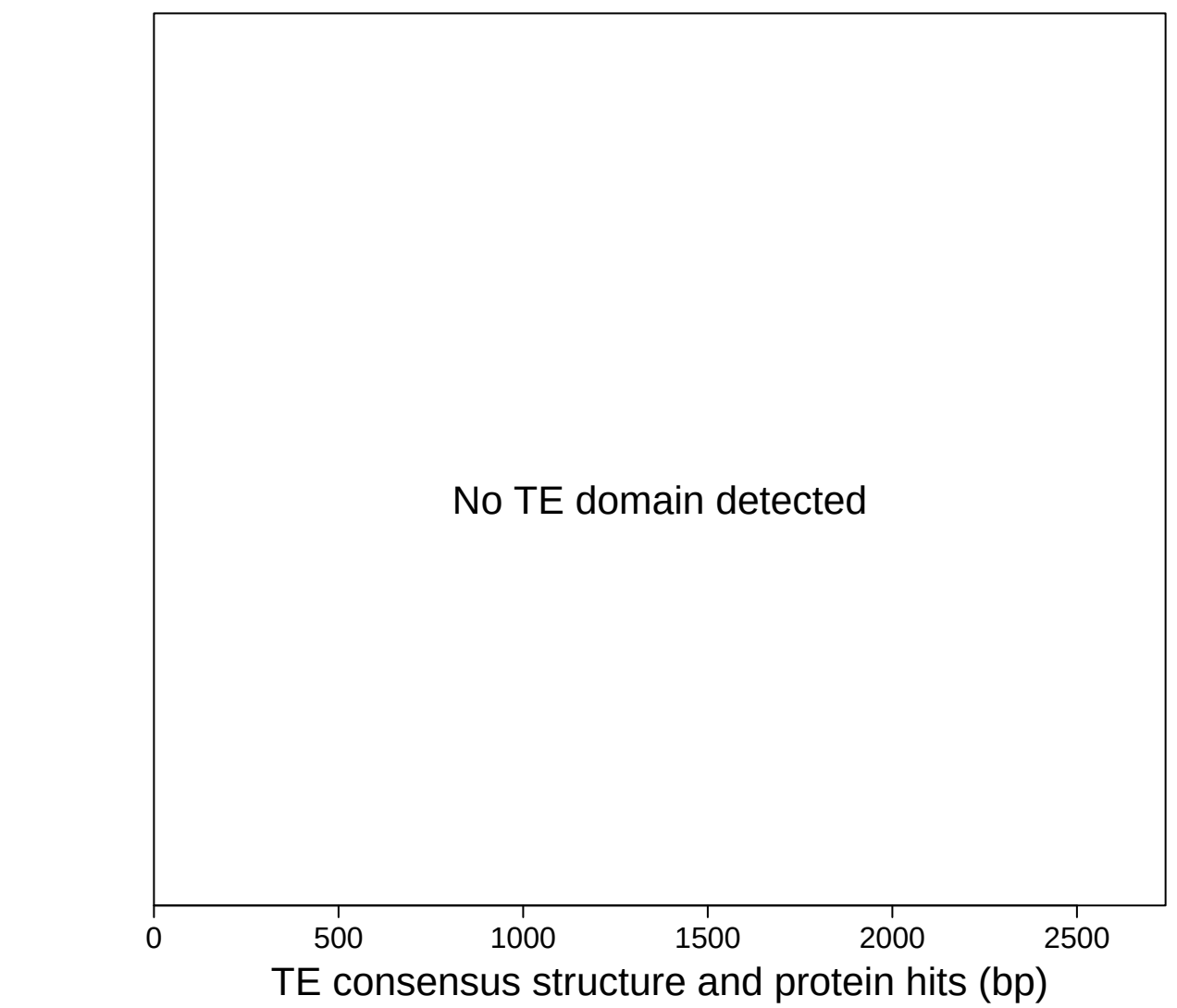
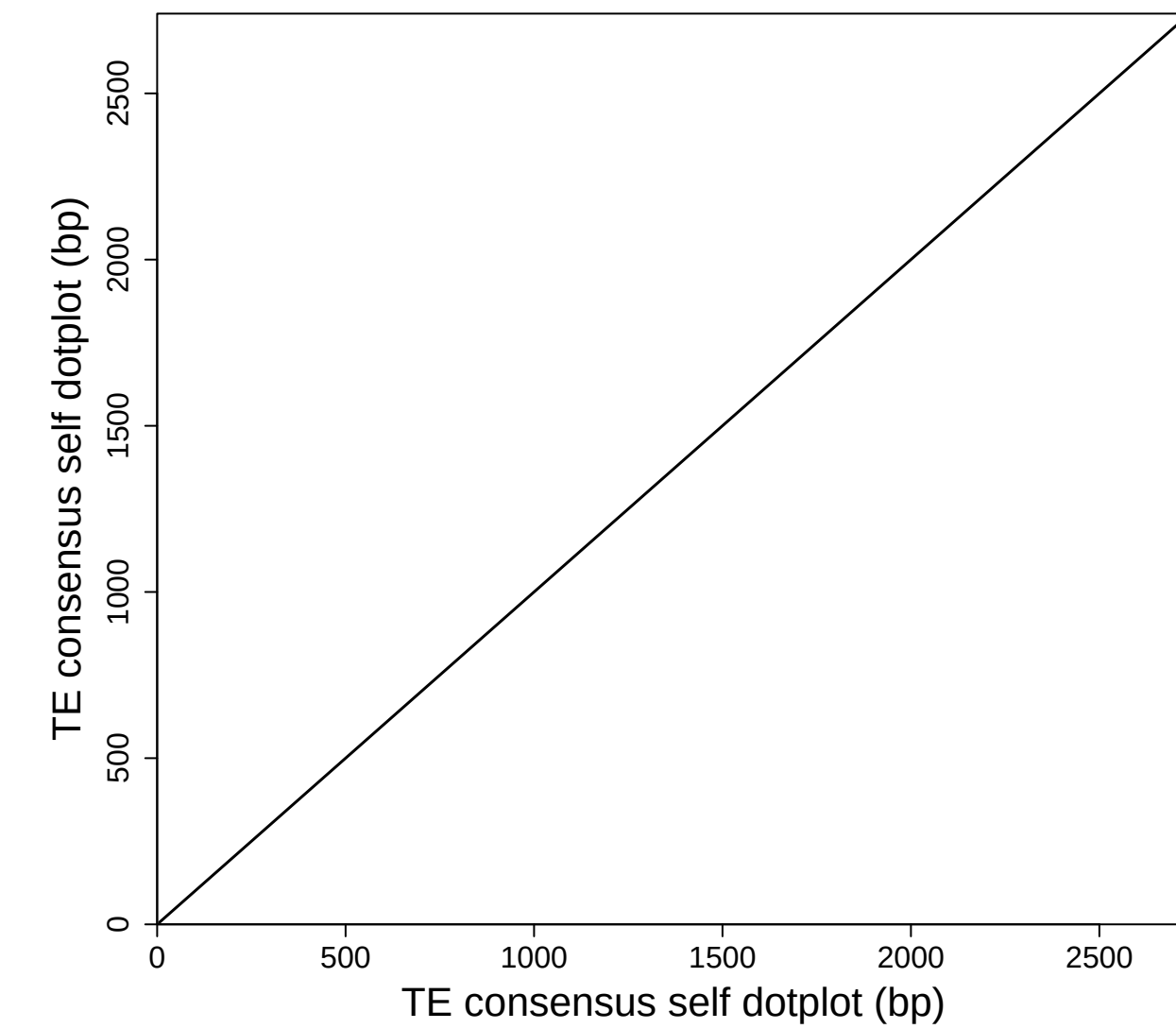
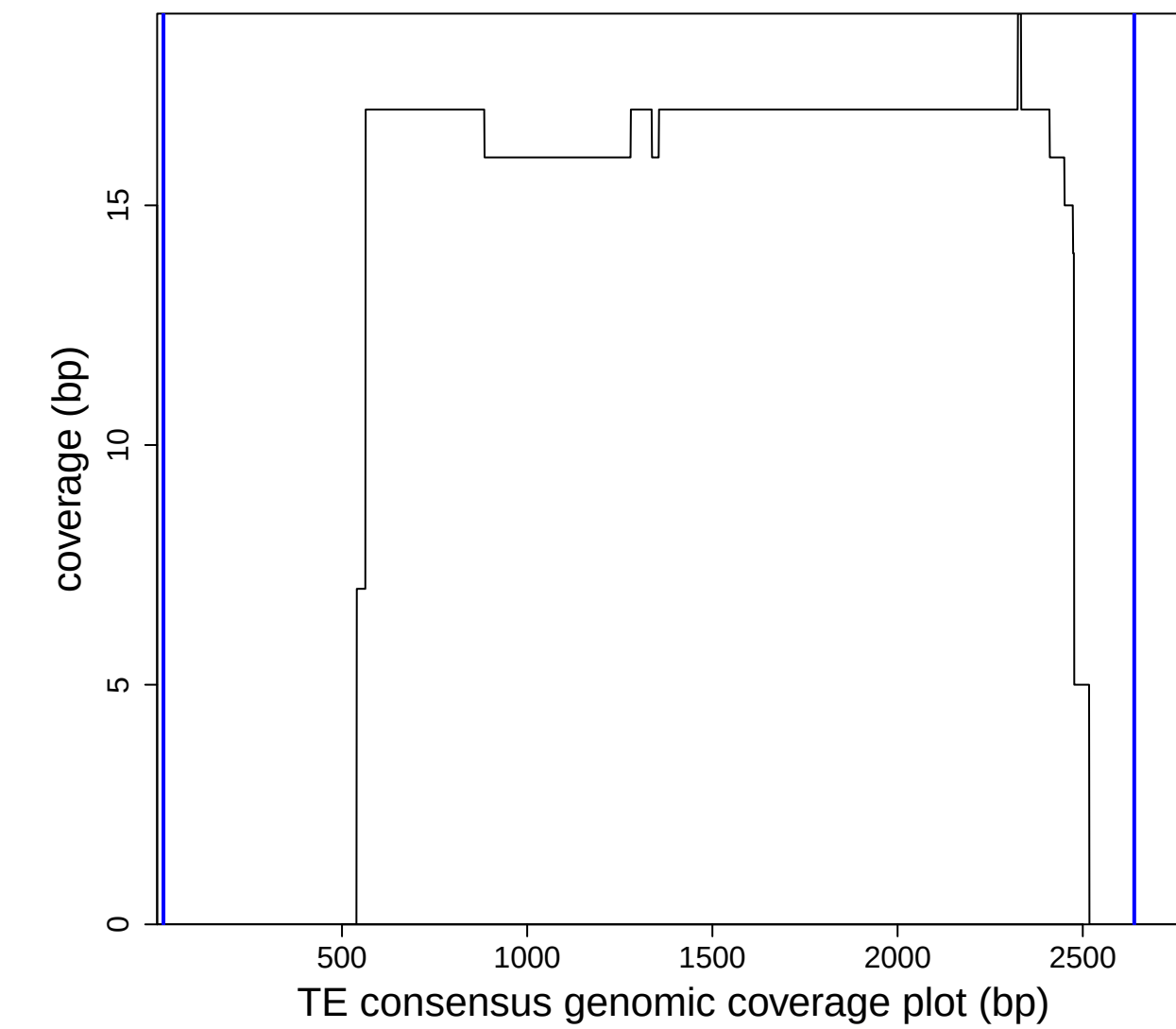
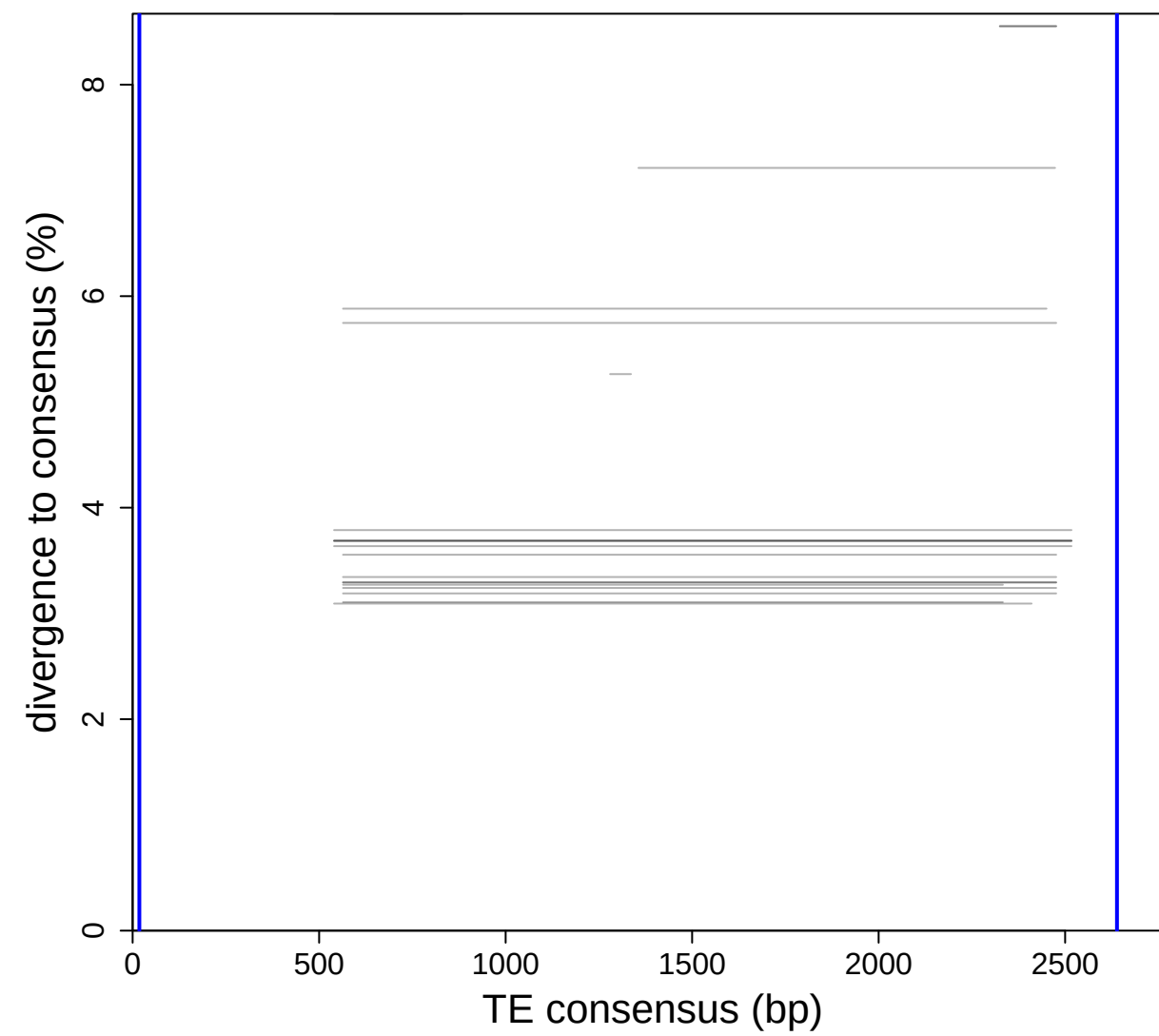
b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 2621bp; fragments: 45; full length: 0 (>=2358.9bp)

After TETrimmer 2621 bp

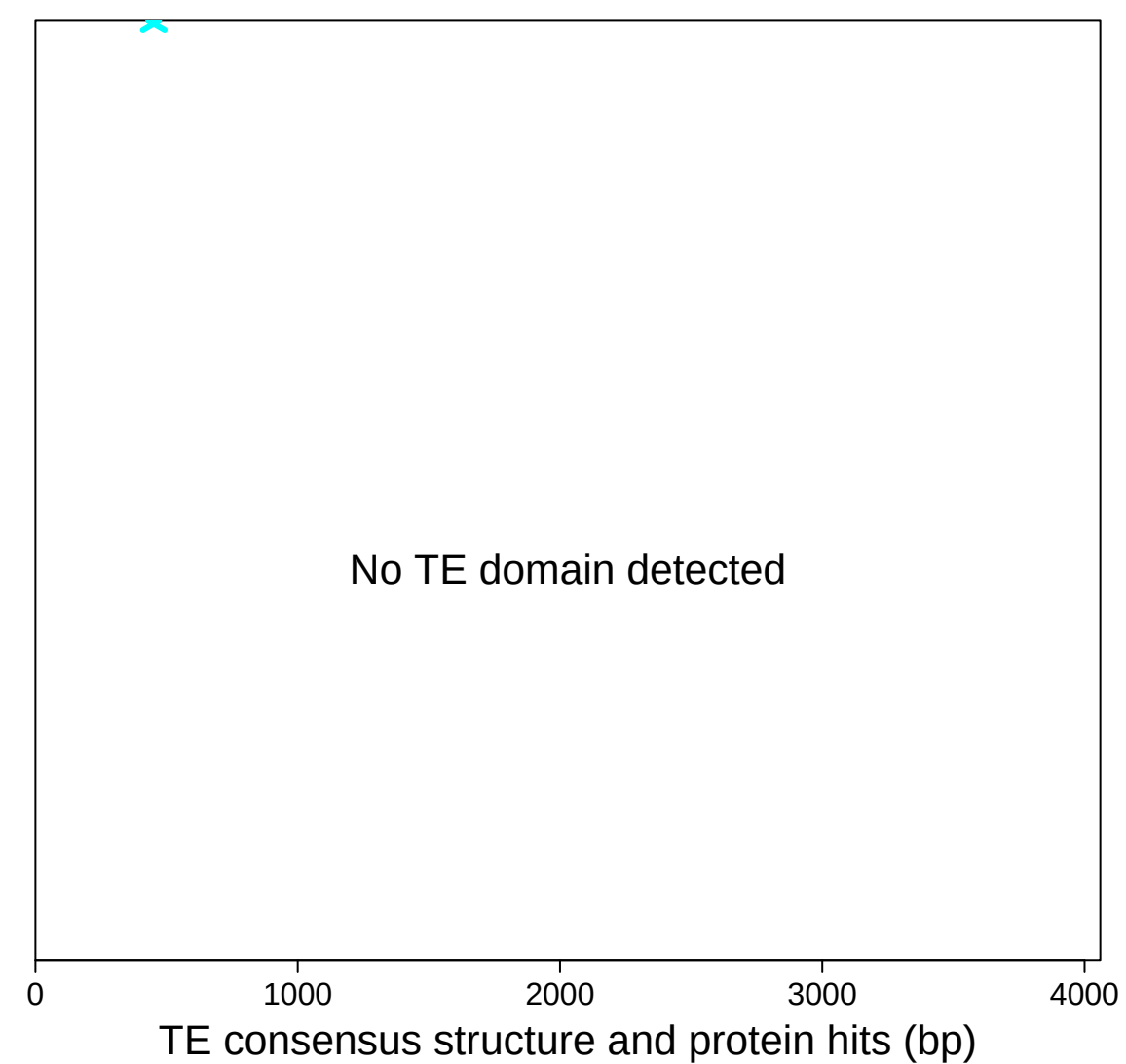
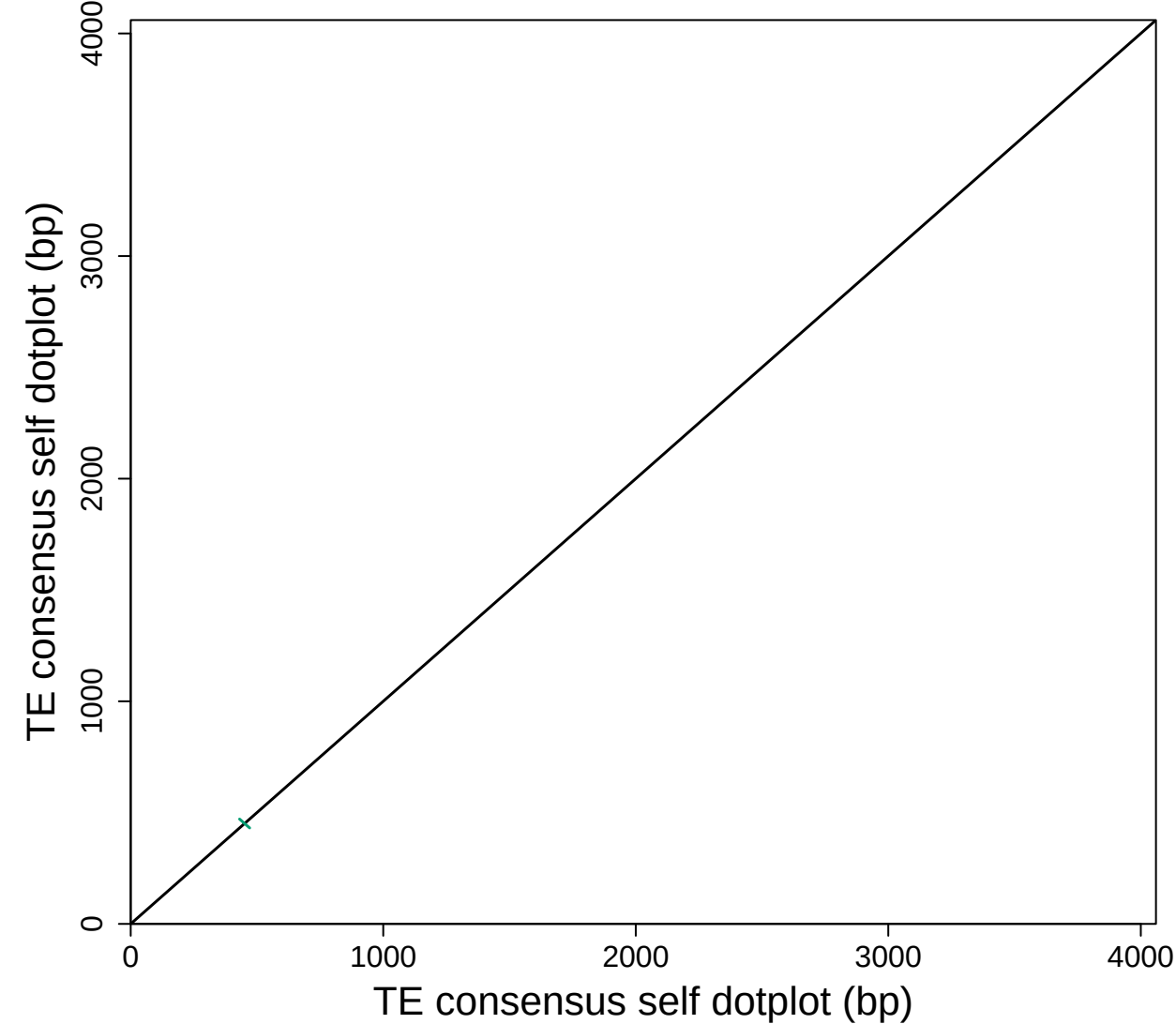
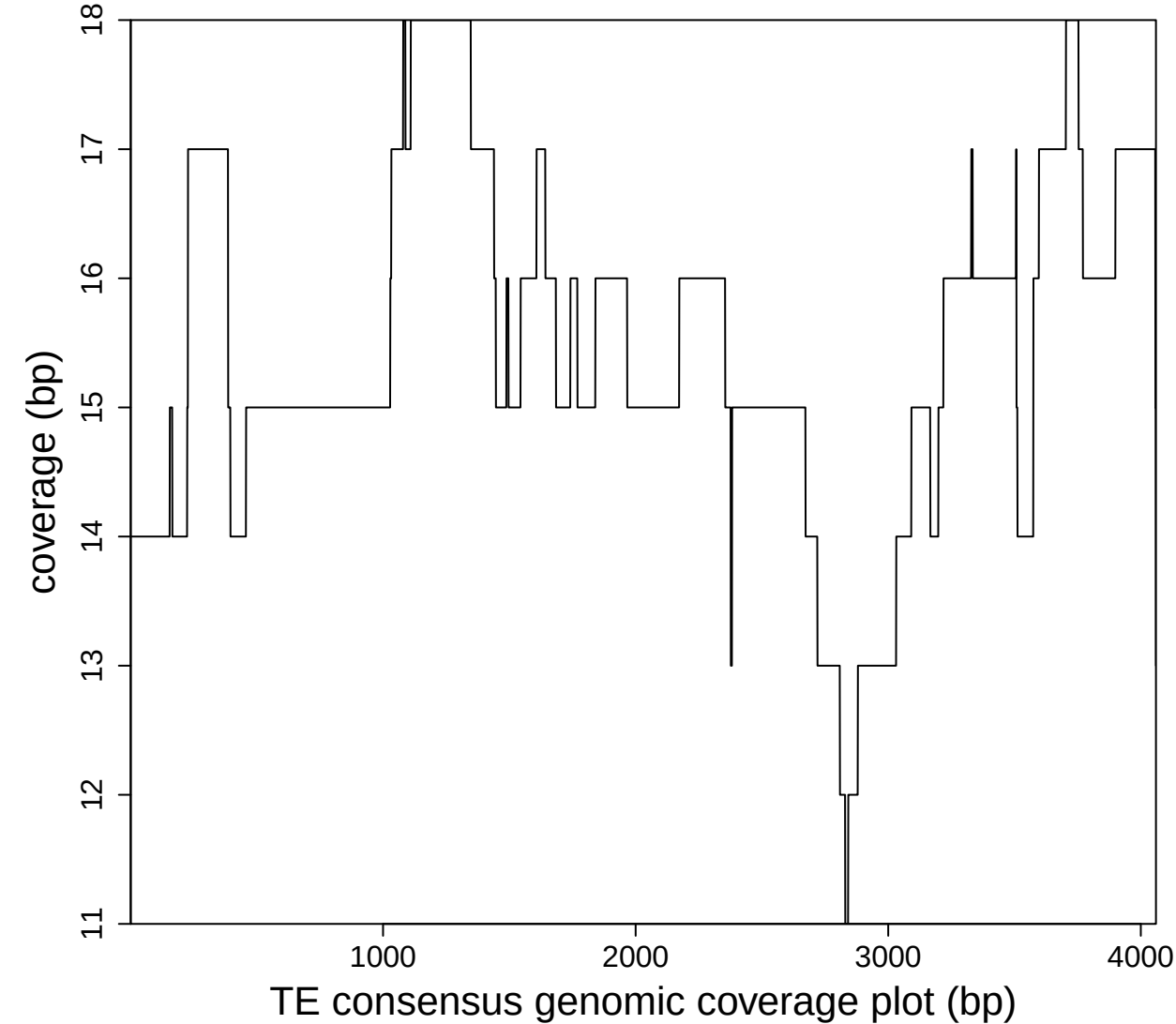
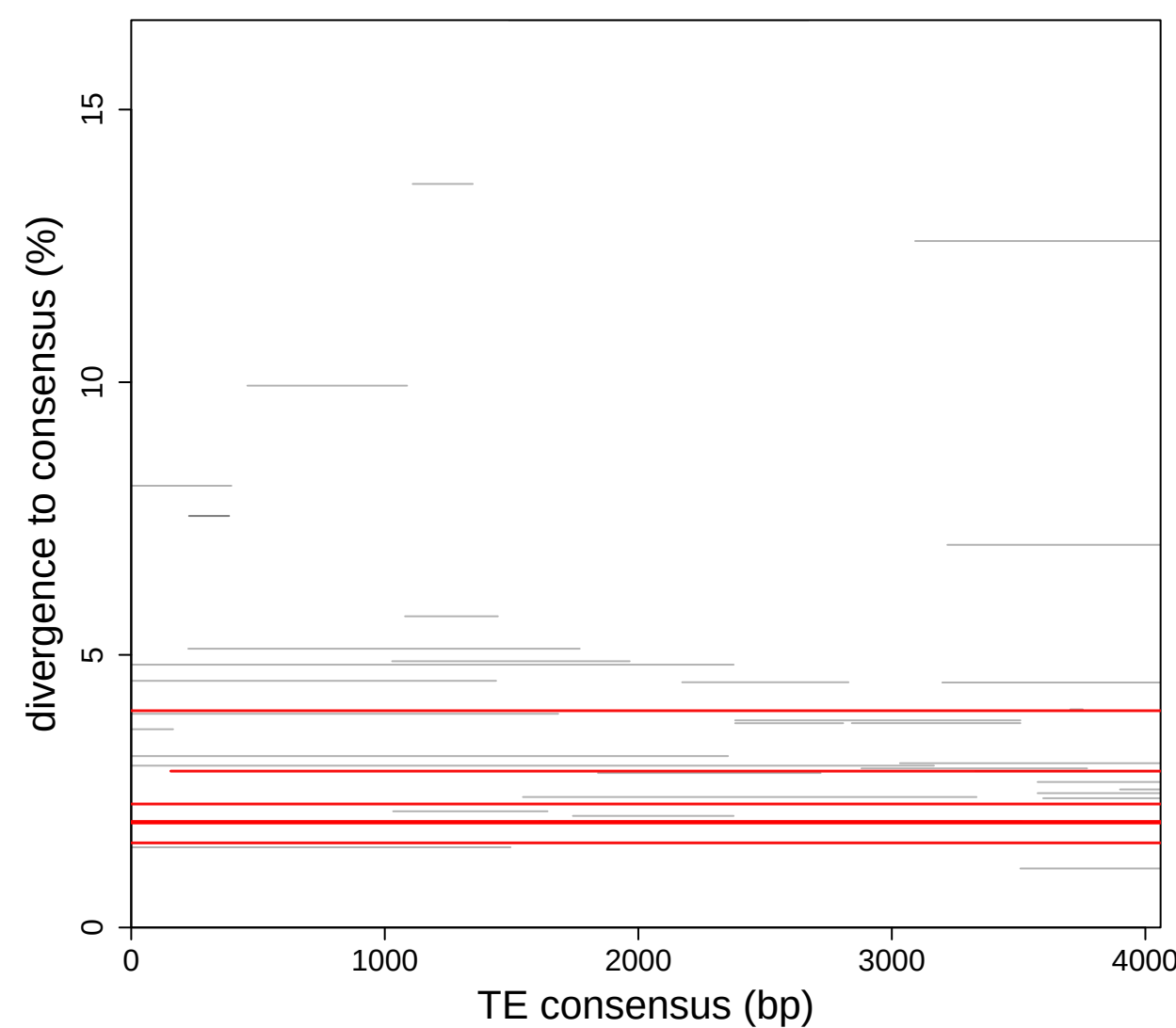


3.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 2789bp; fragments: 21; full length: 0 (>=2510.1bp)

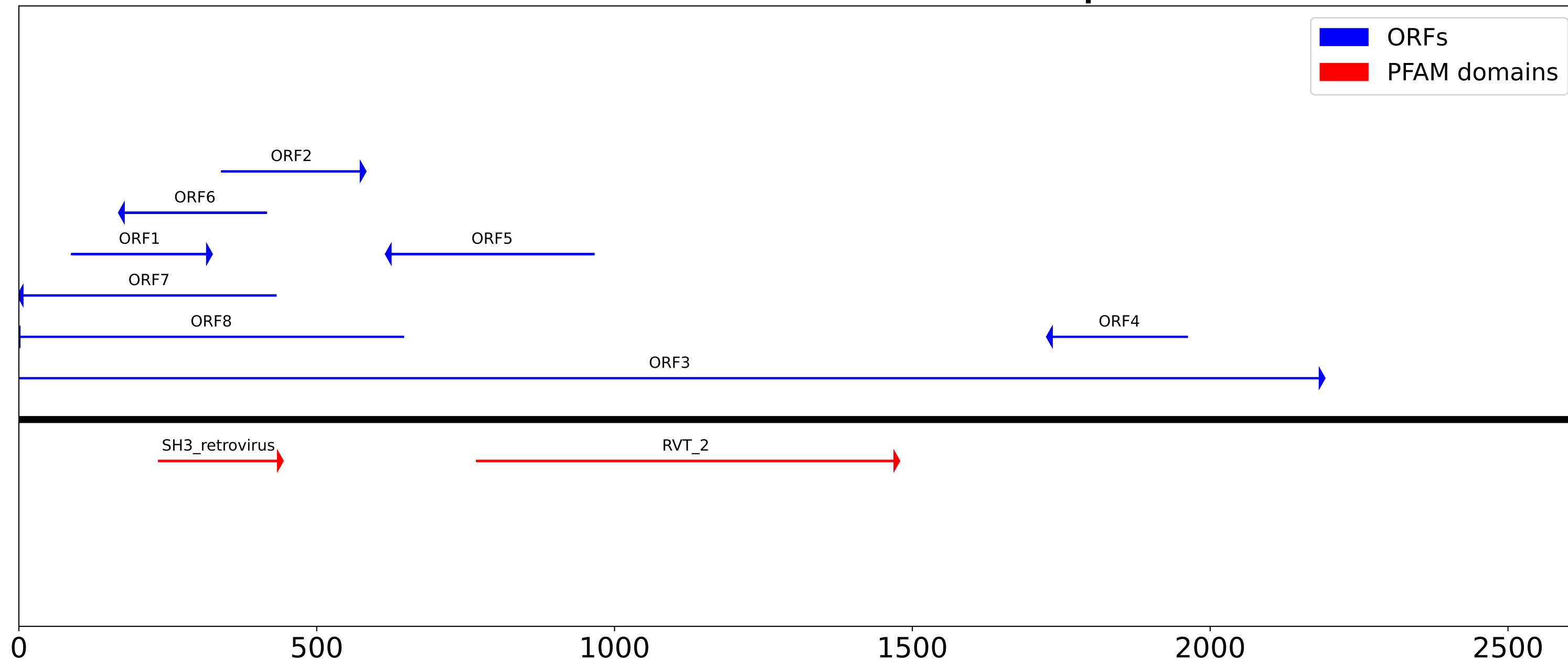
After TEtrimmer Extended plot Blue lines are boundaries



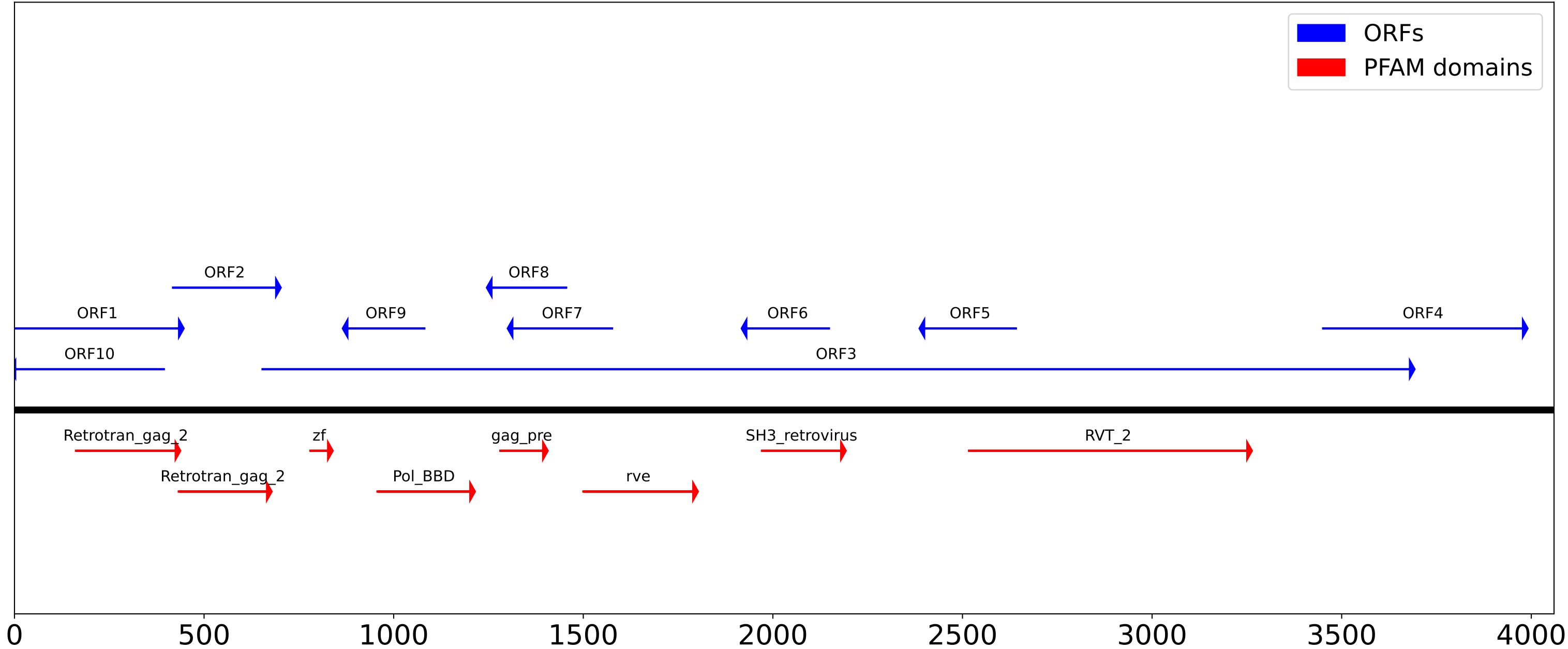
TE: ltr_1_family_668
size: 4060bp; fragments: 44; full length: 7 (≥ 3654 bp)



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

