

Start crop Point

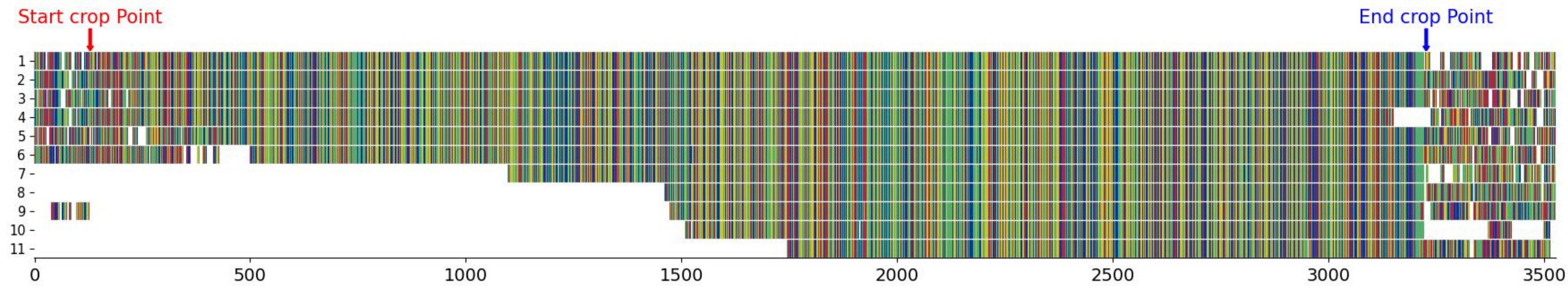


End crop Point



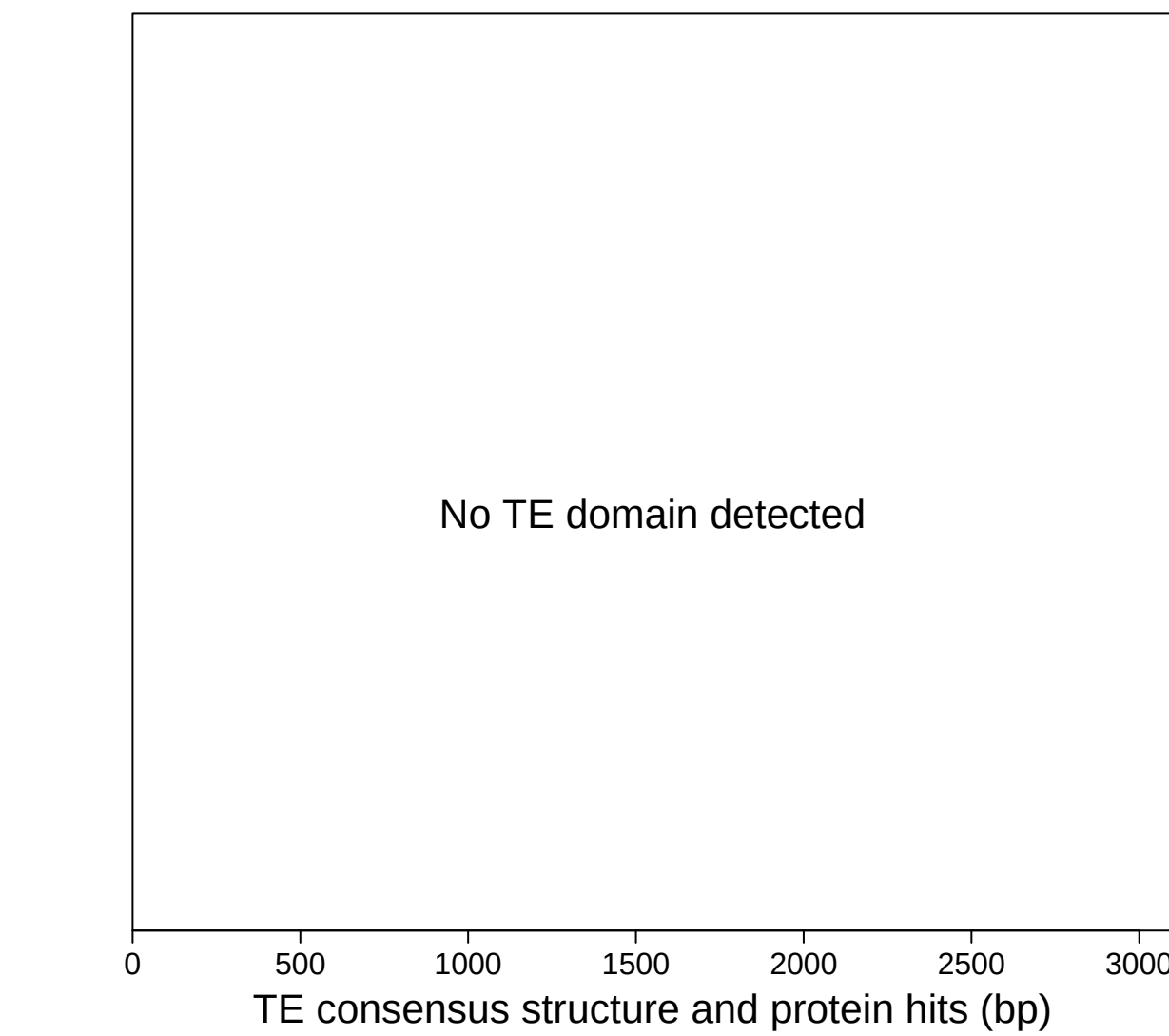
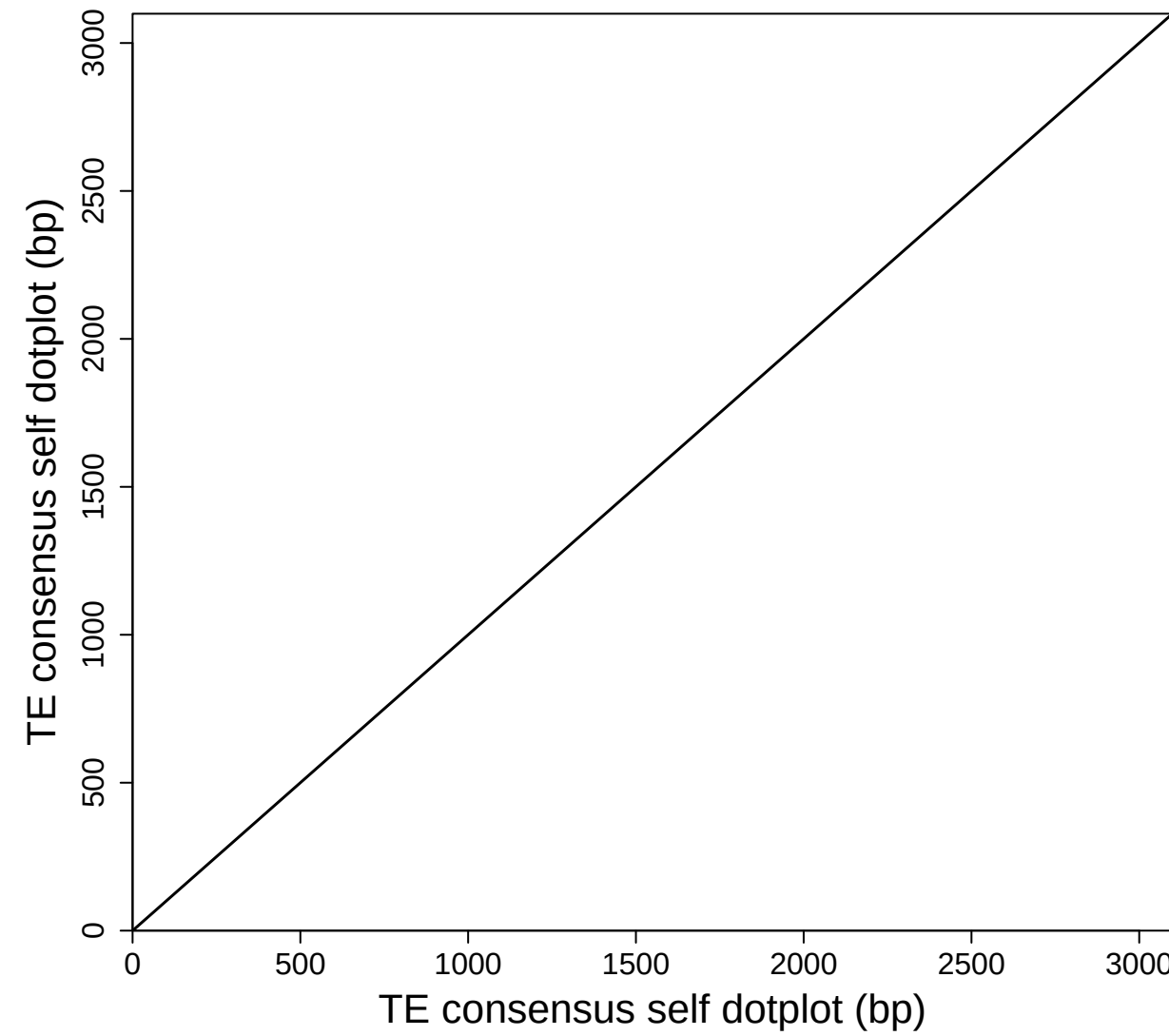
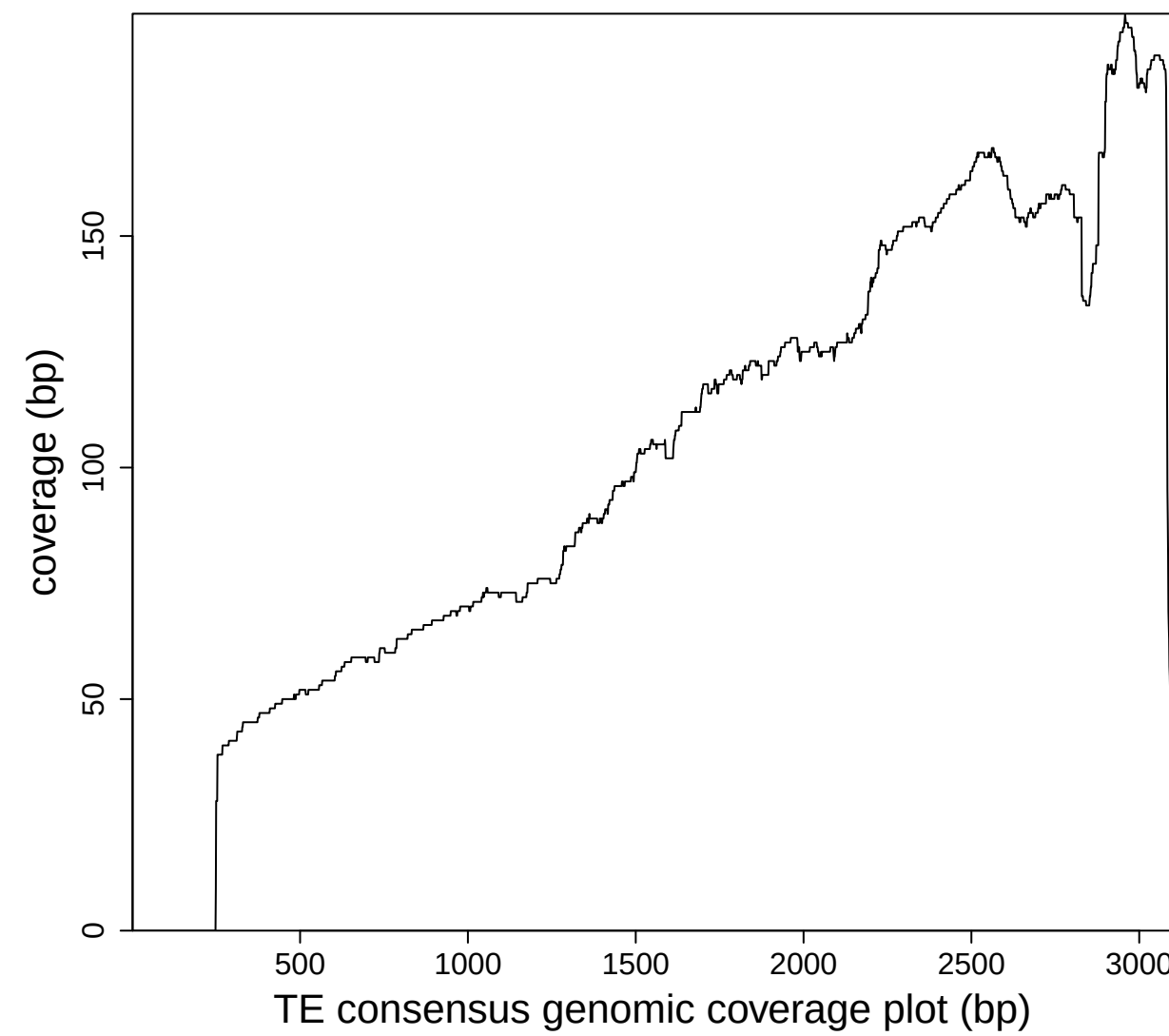
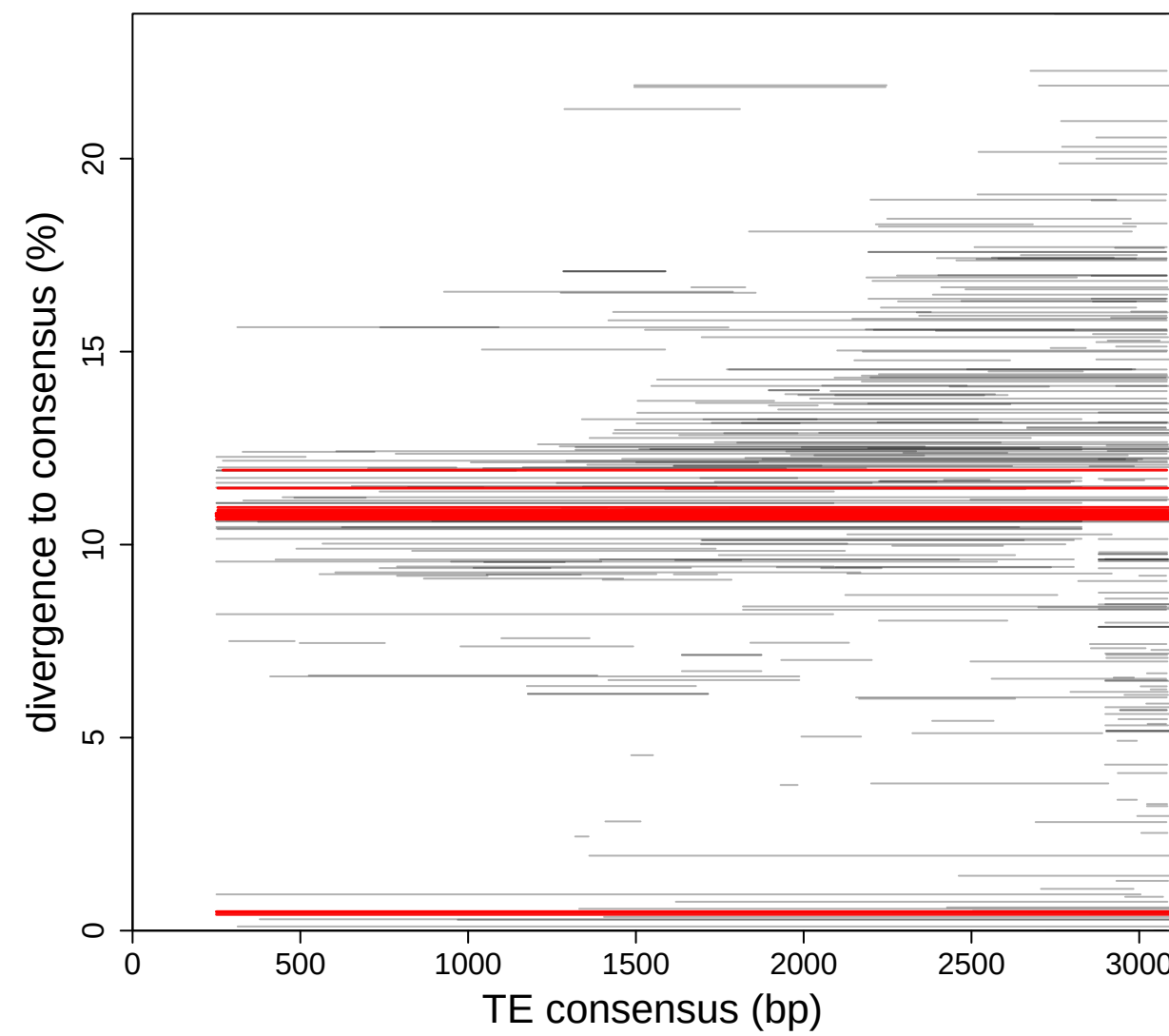
MSA length = 3099





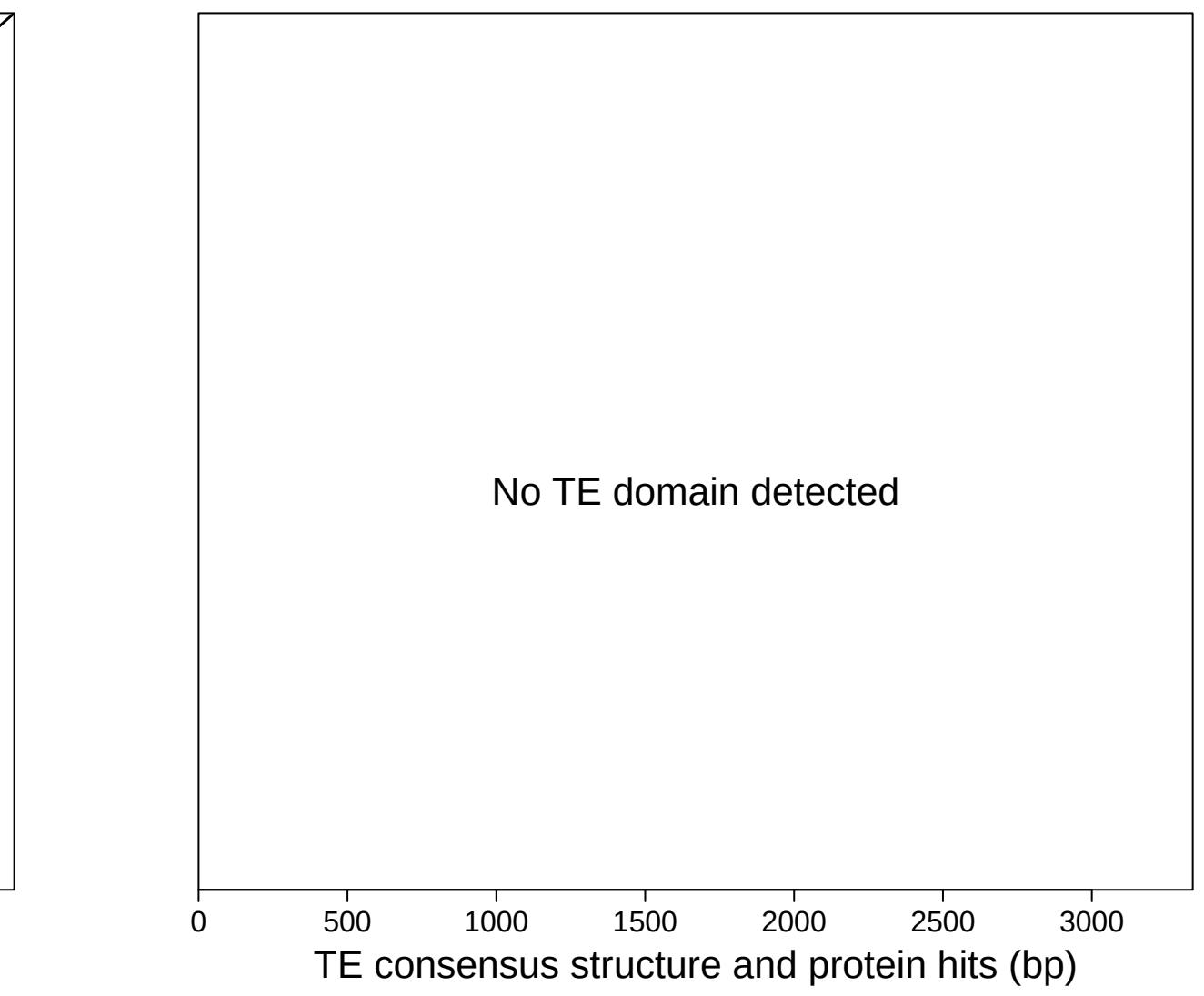
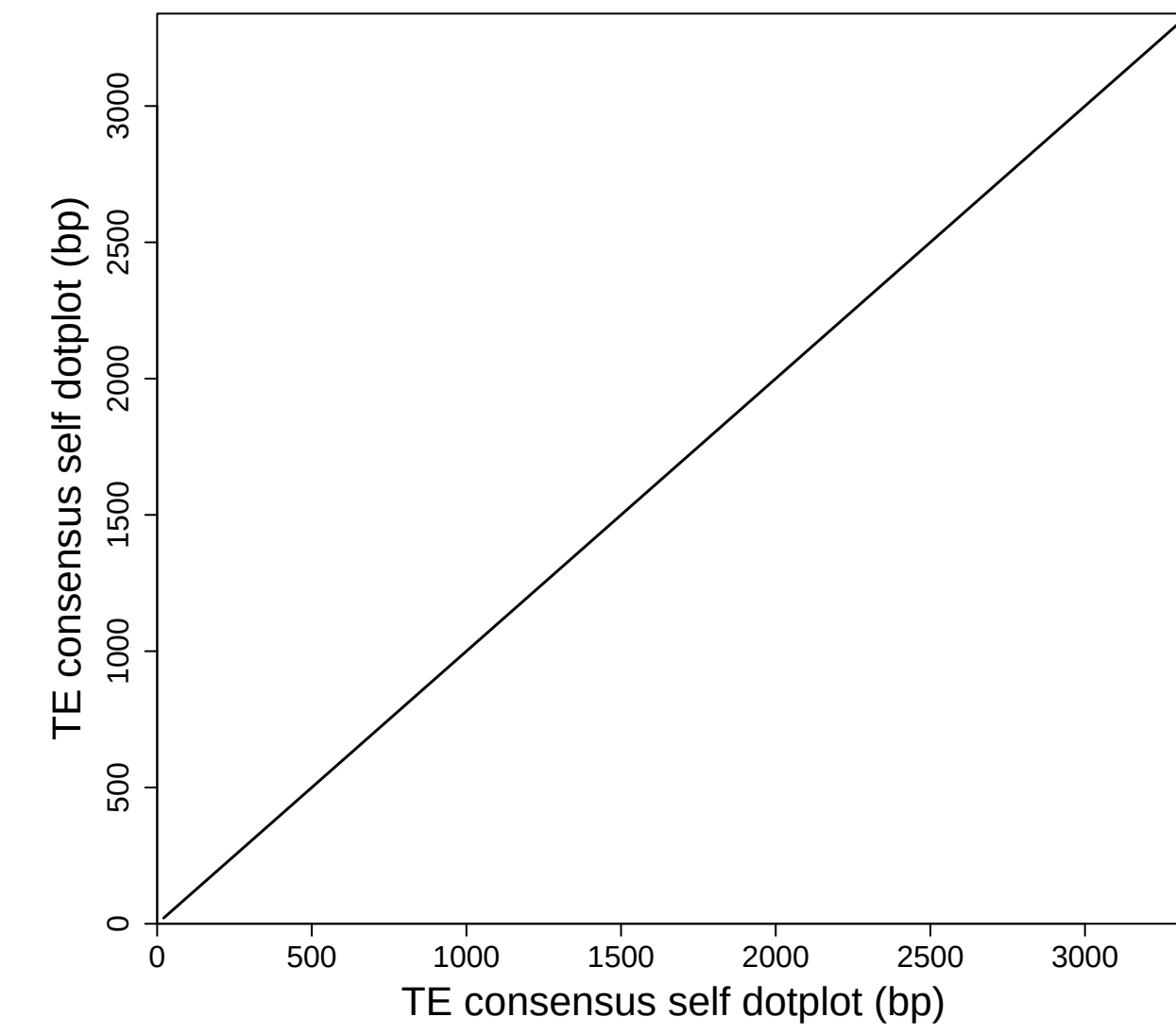
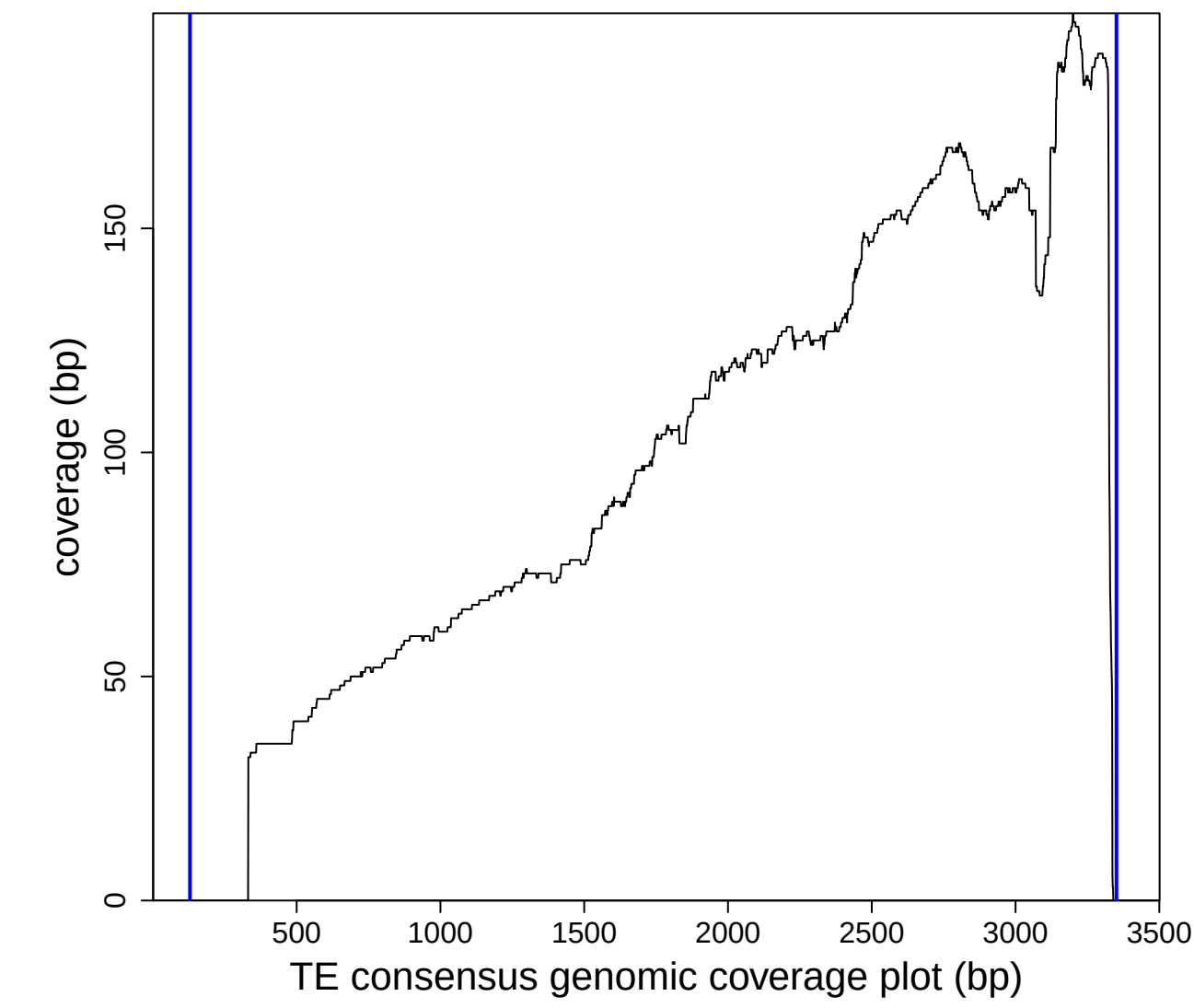
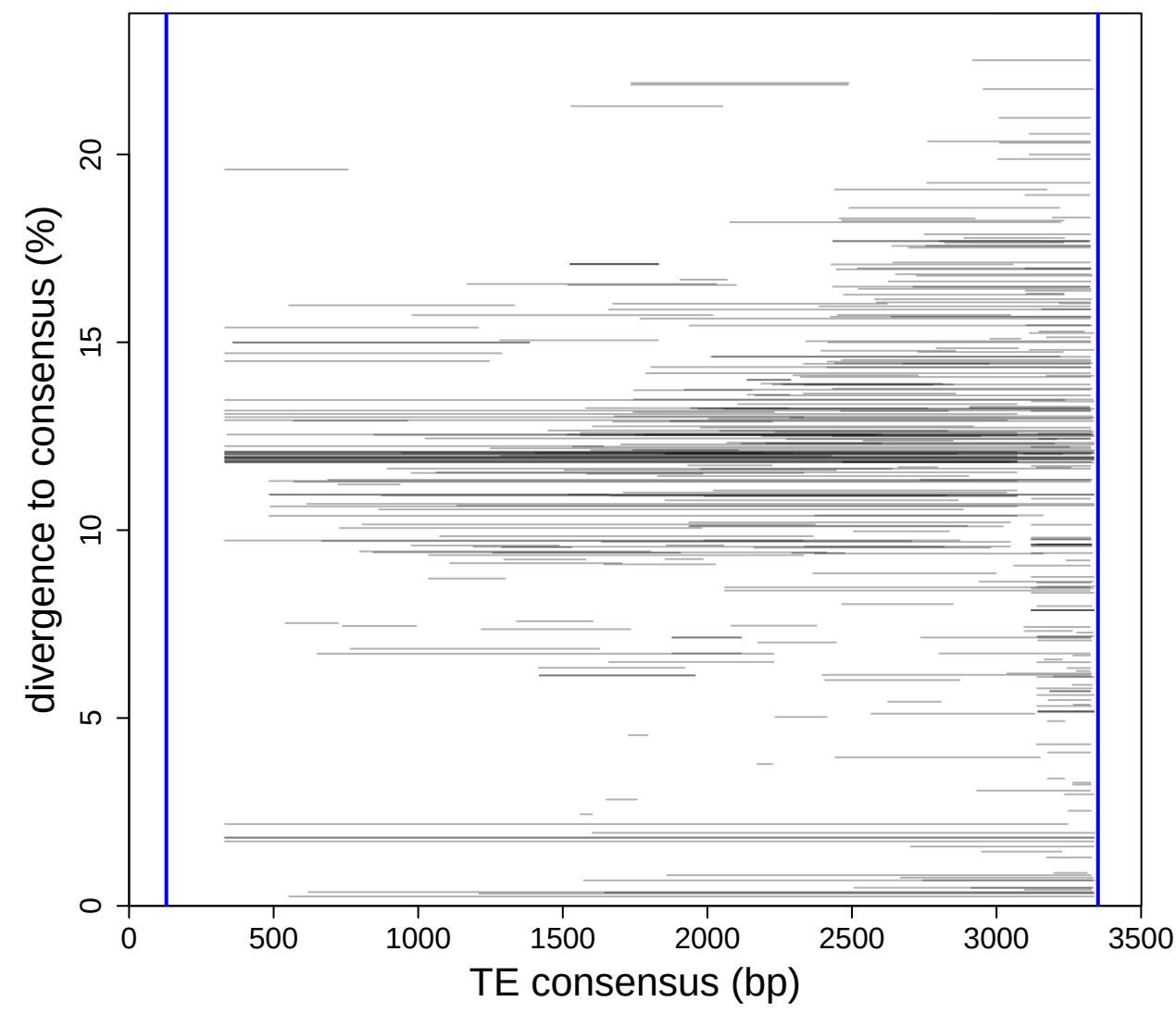
bed uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa ce
size: 3099bp; fragments: 355; full length: 19 (>=2789.1bp)

After TEtrimmer 3099 bp

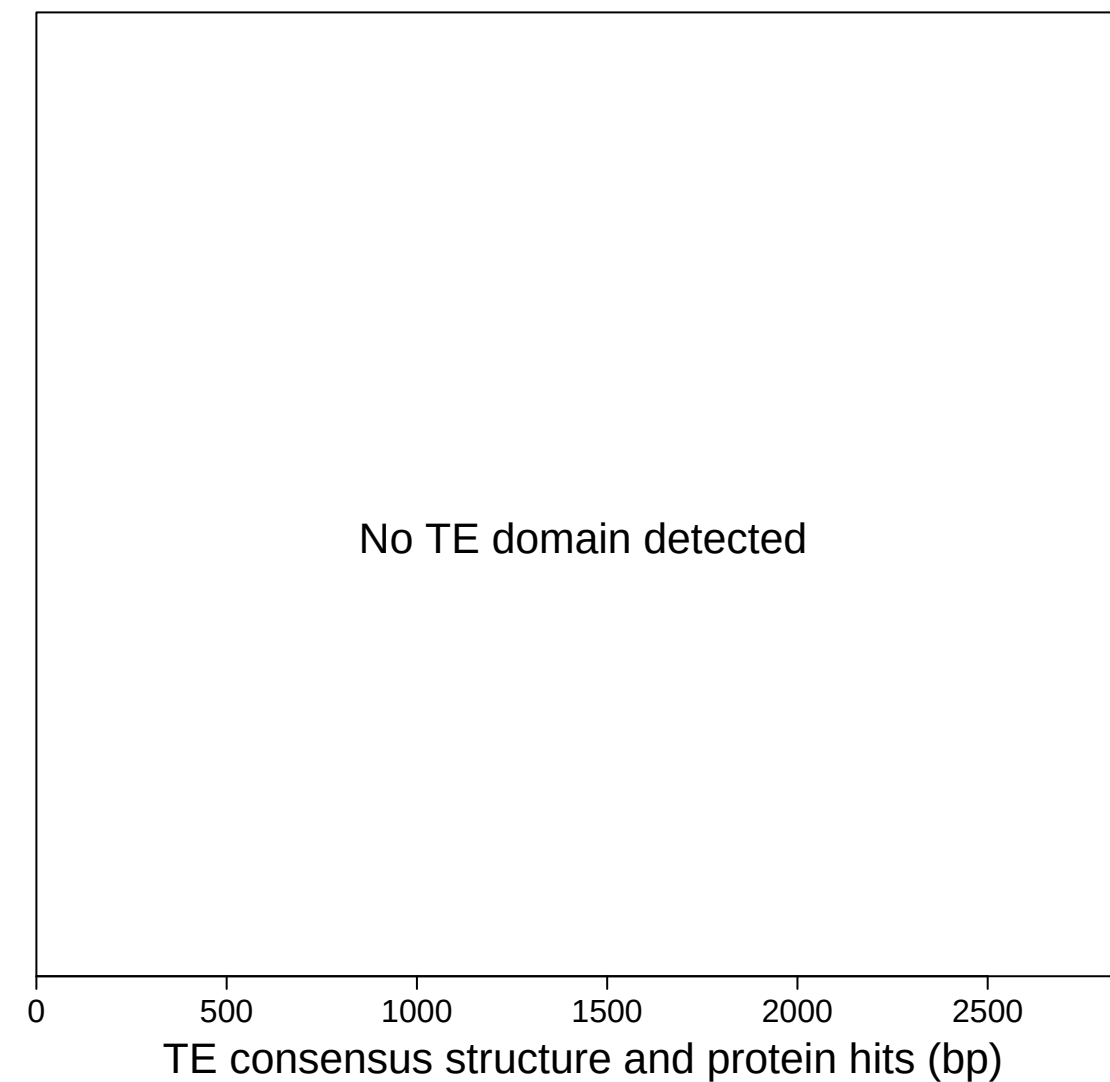
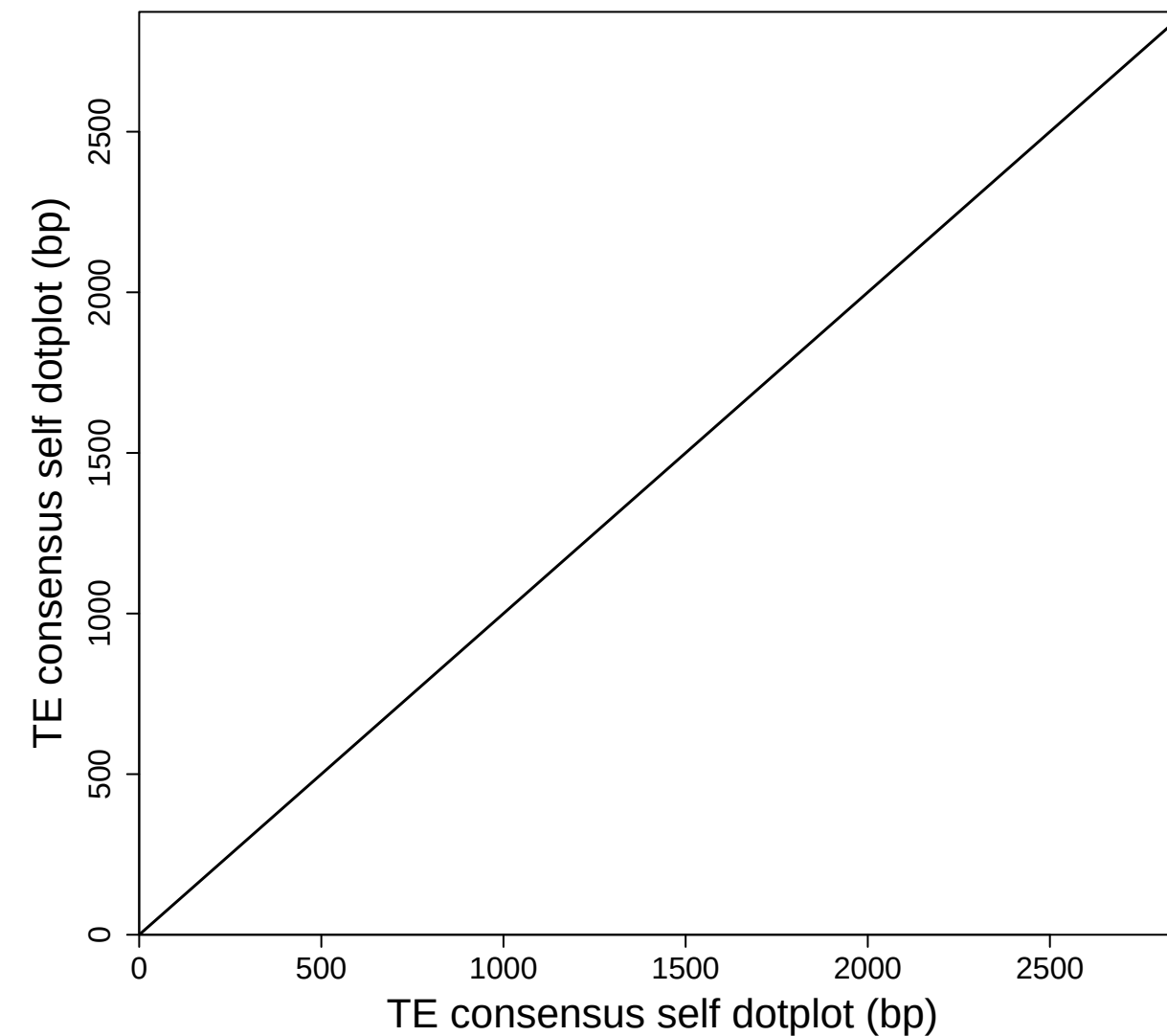
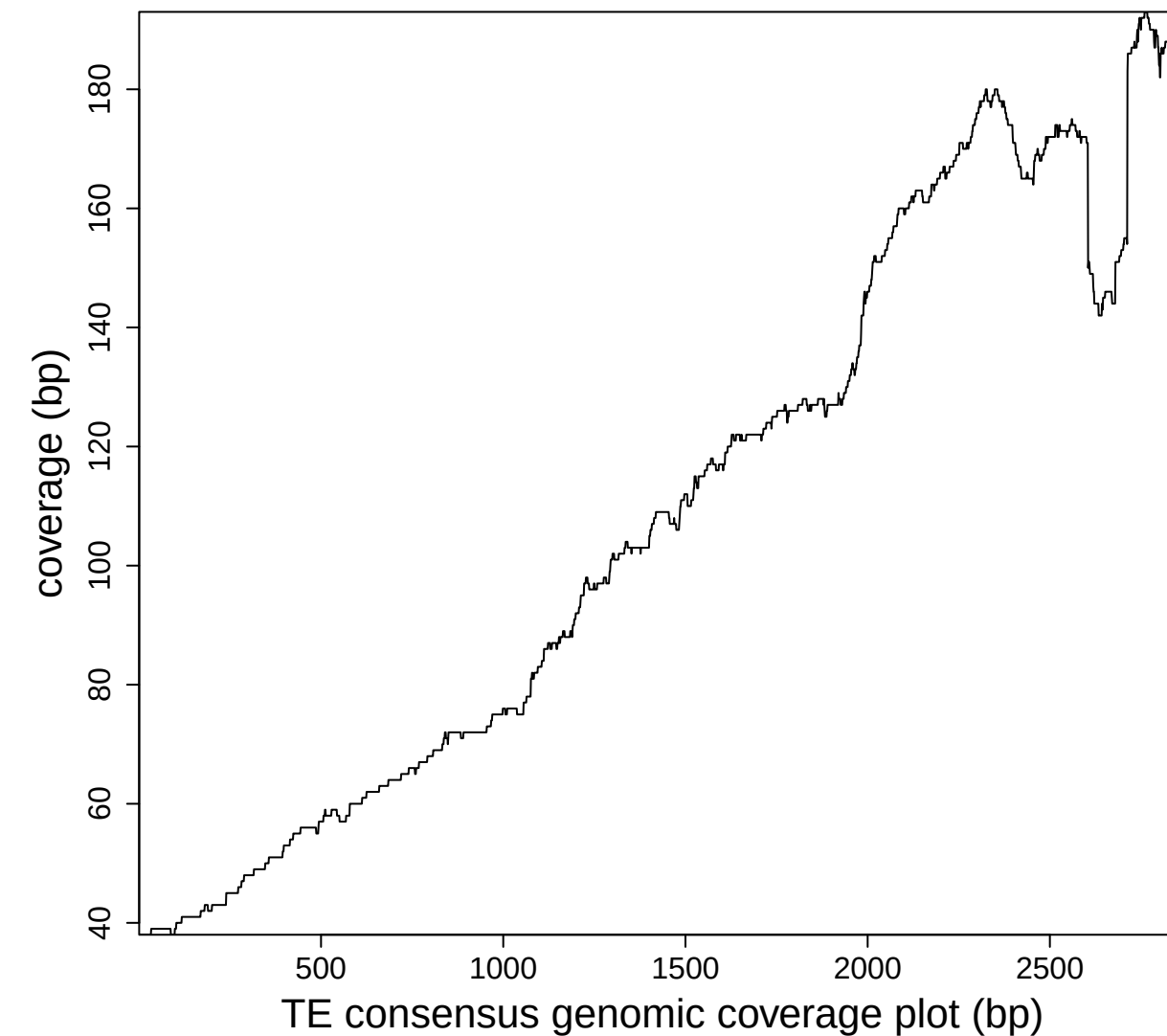
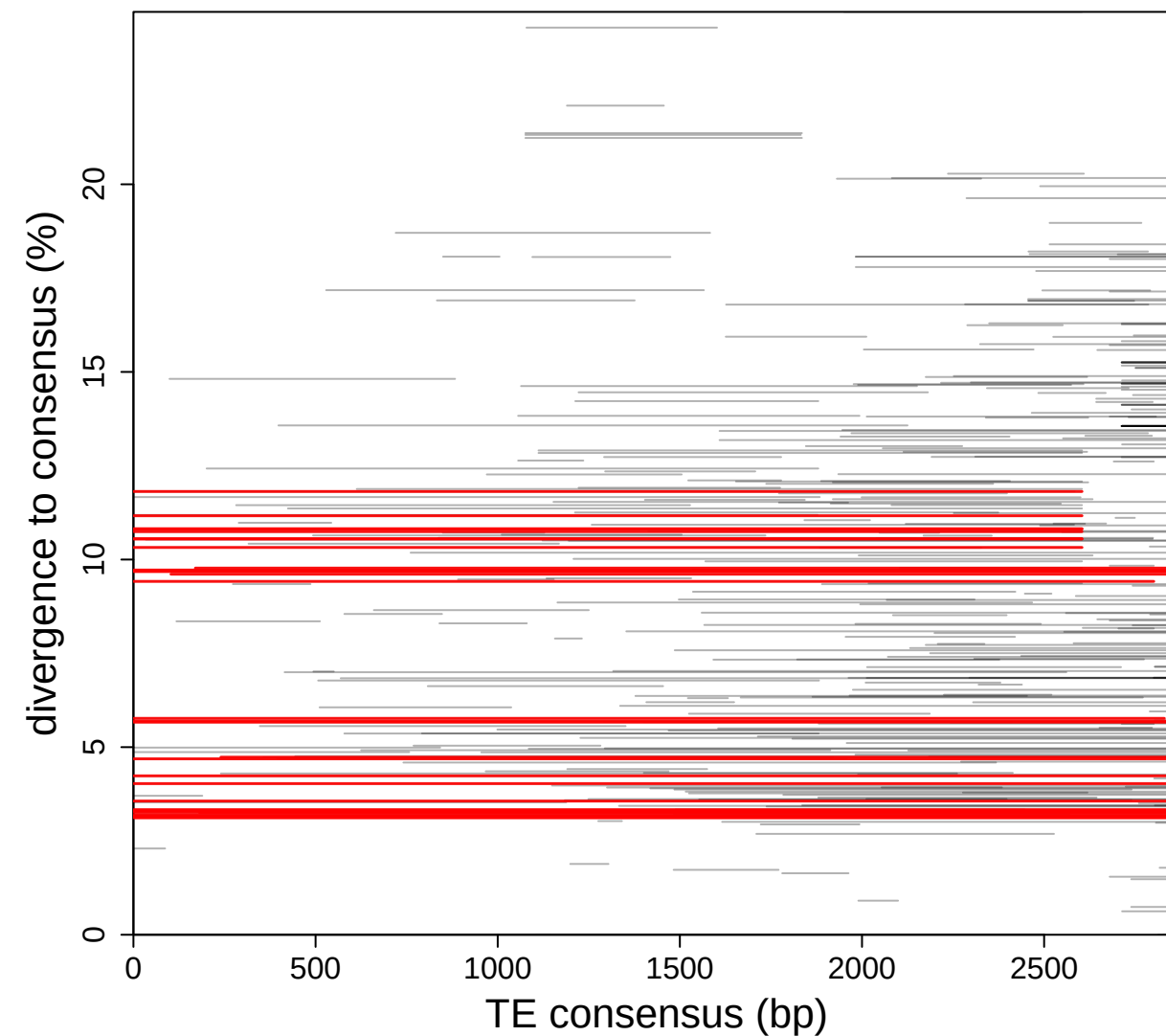


0.fasta.b.bed uf.bed g_3.bed fm_1.bed_0_0 bcln.fa aln.fa cl.fa
size: 3501bp; fragments: 355; full length: 0 (>=3150.9bp)

After TETrimmer Extended plot Blue lines are boundaries

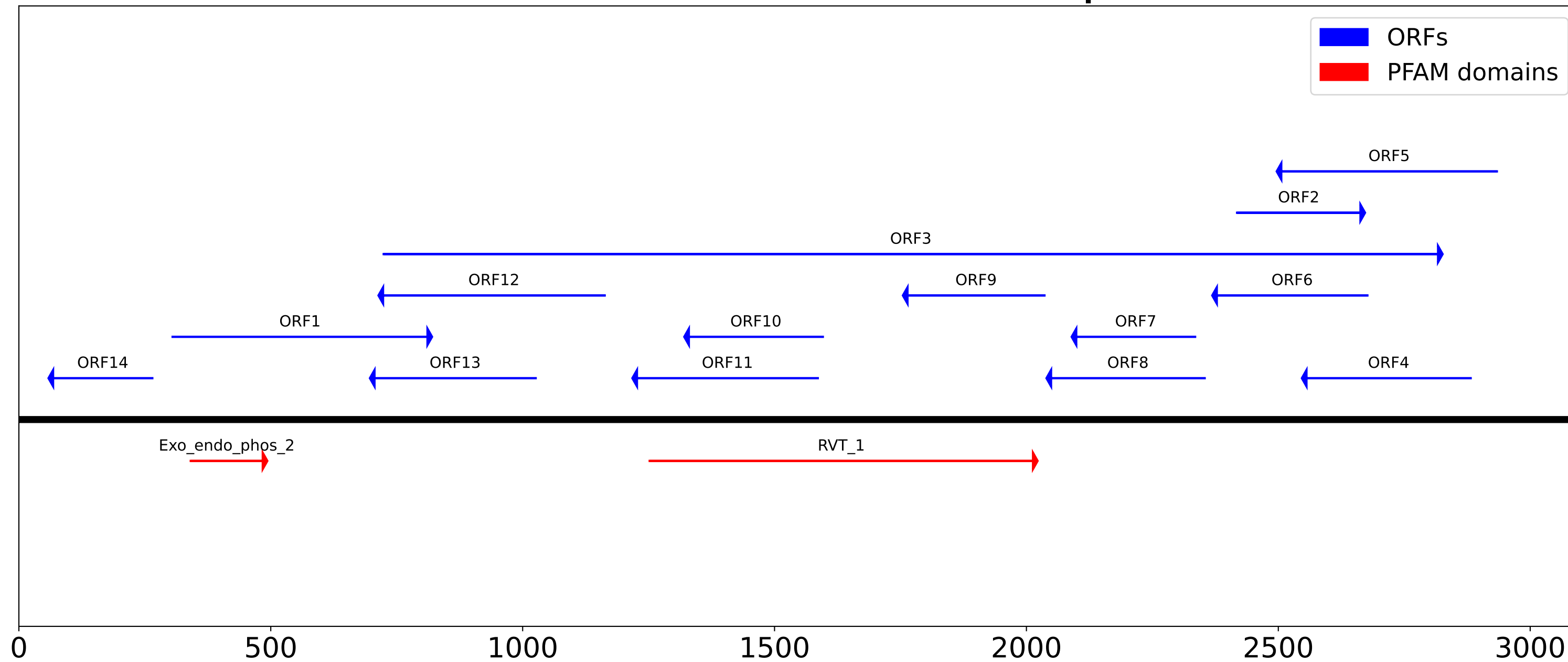


TE: rnd_4_family_950
size: 2873bp; fragments: 361; full length: 34 (≥ 2585.7 bp)

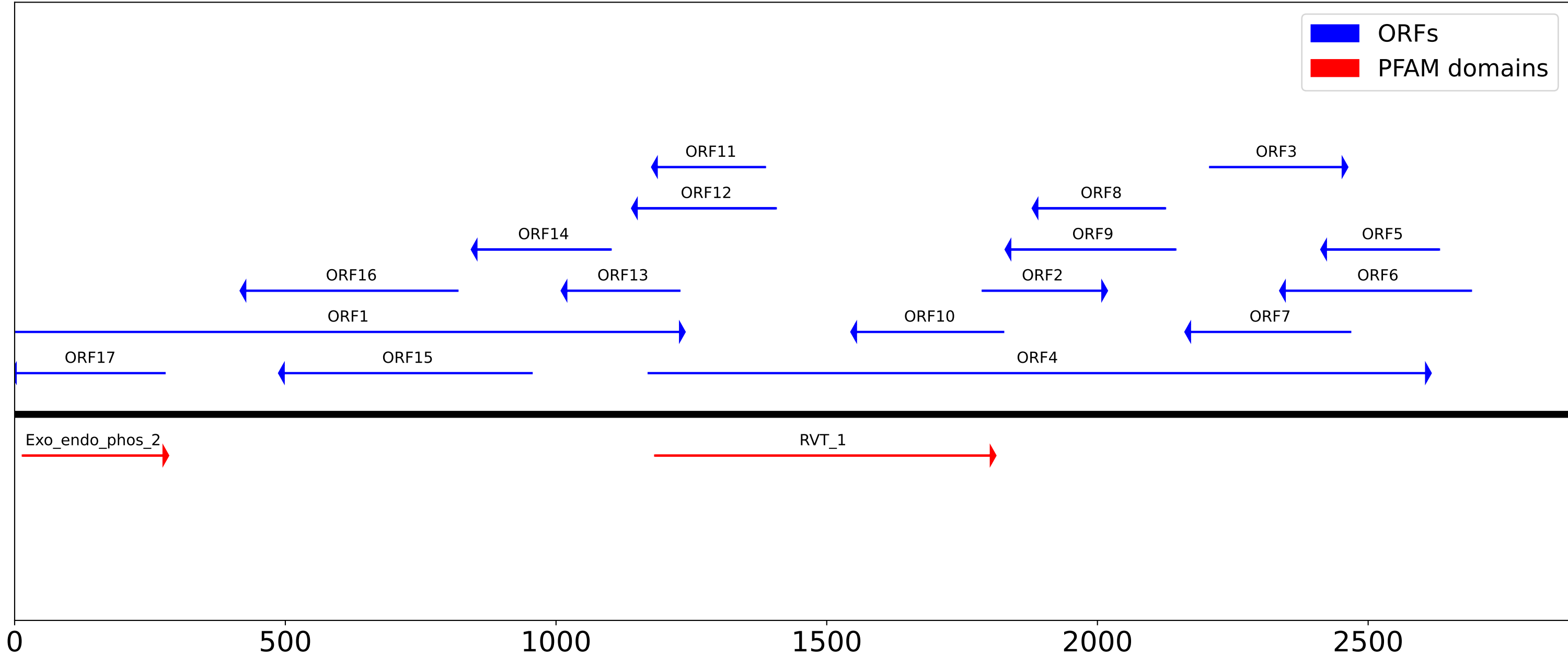


Before TEtrimmer 2873 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

