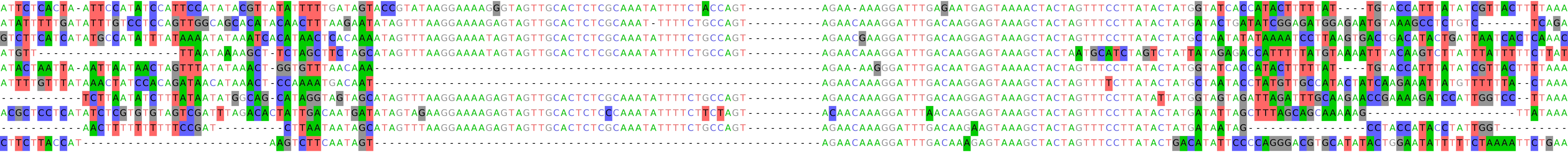


Start Crop Point

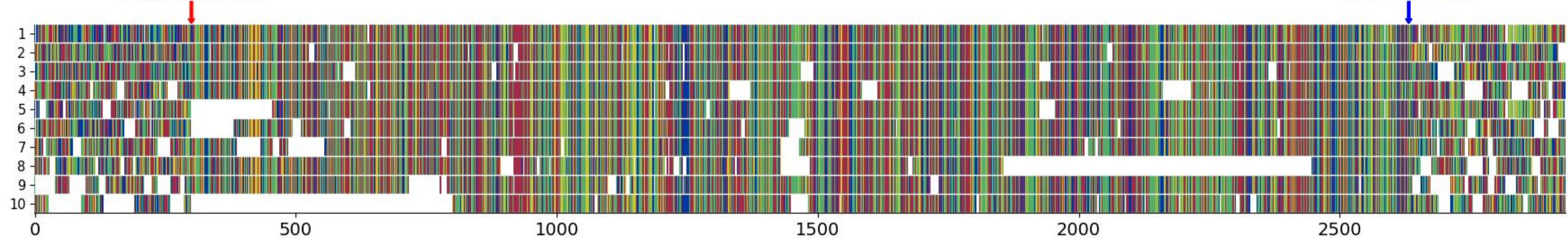
End Crop Point

MSA length = 2333



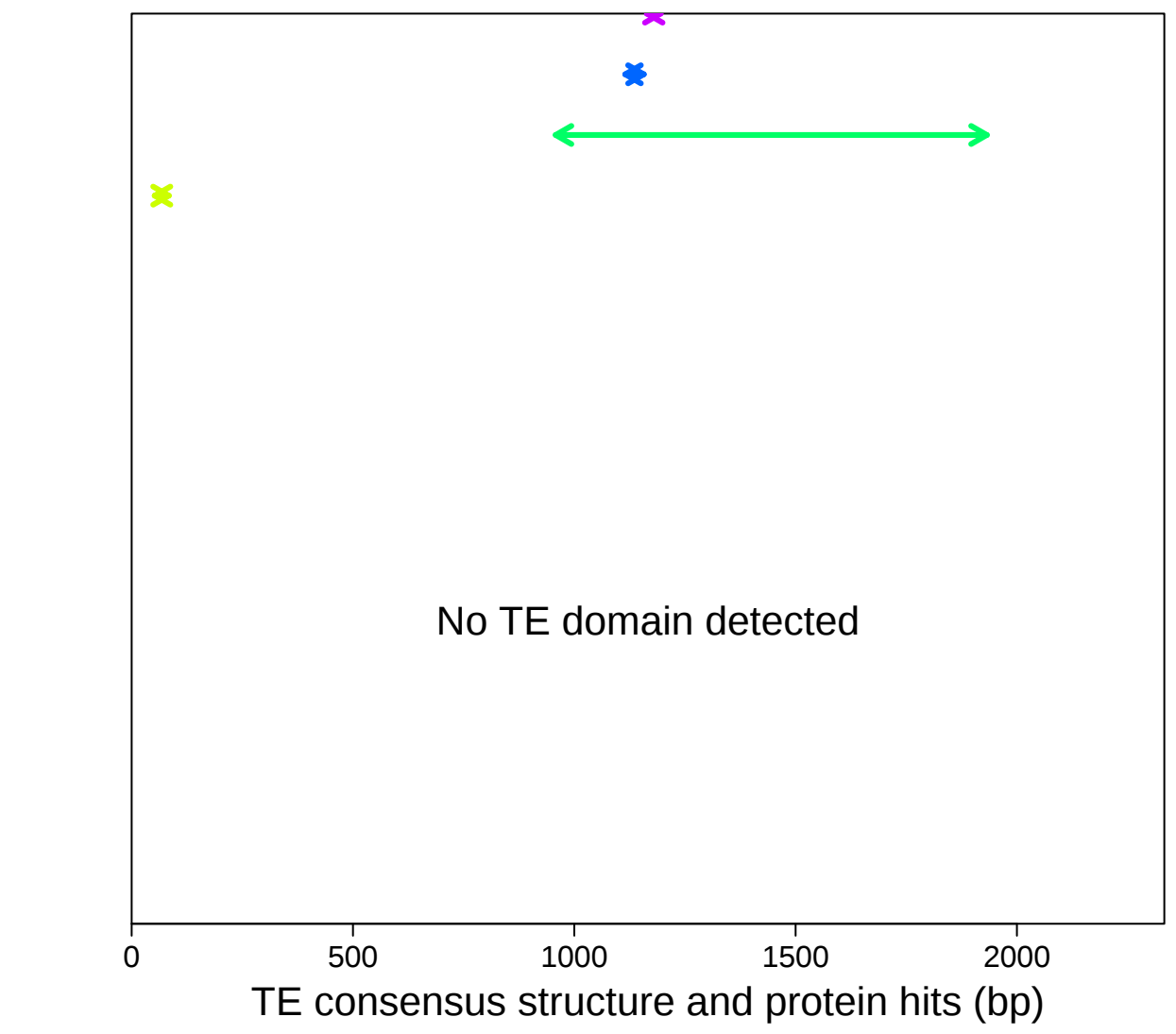
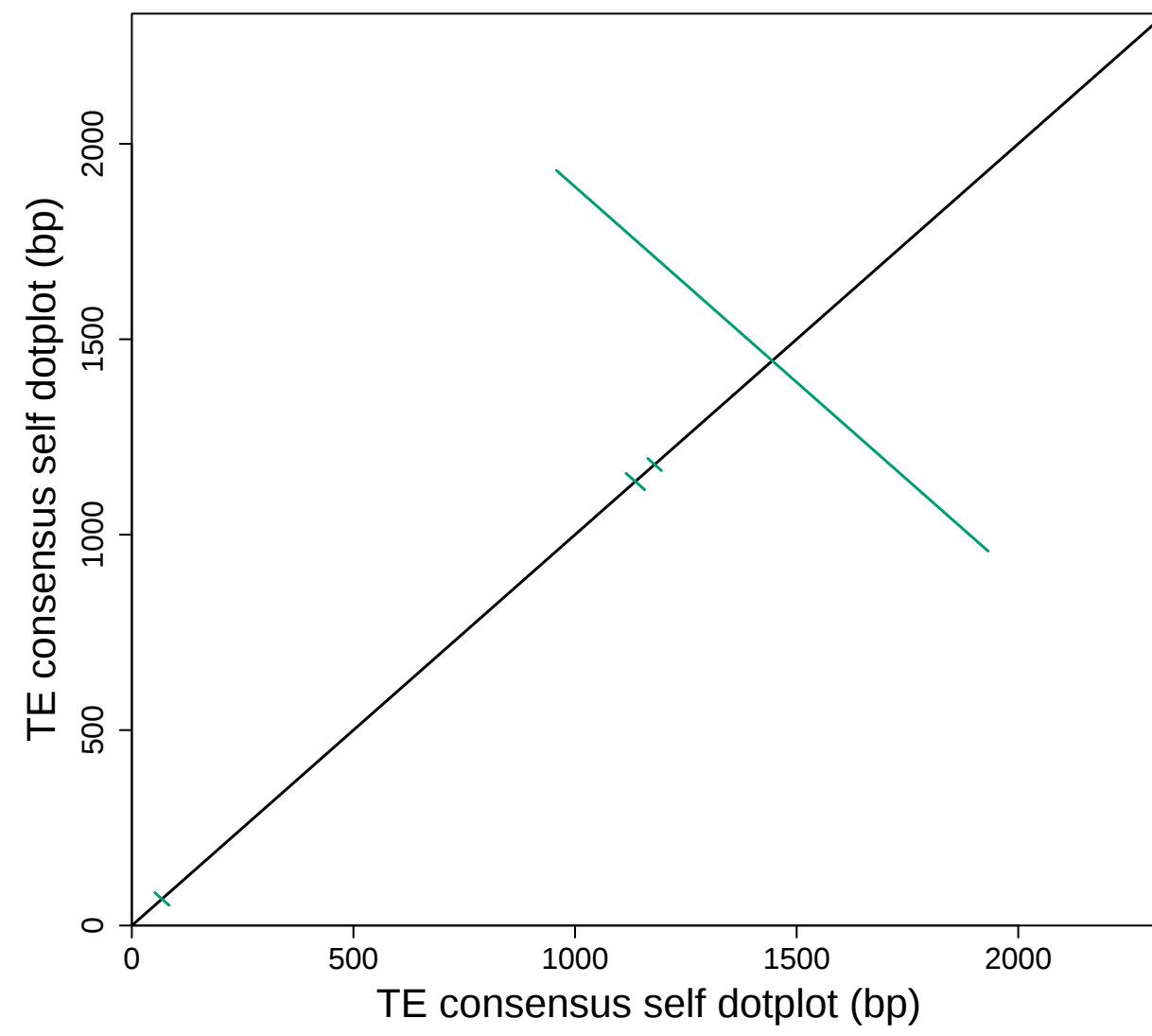
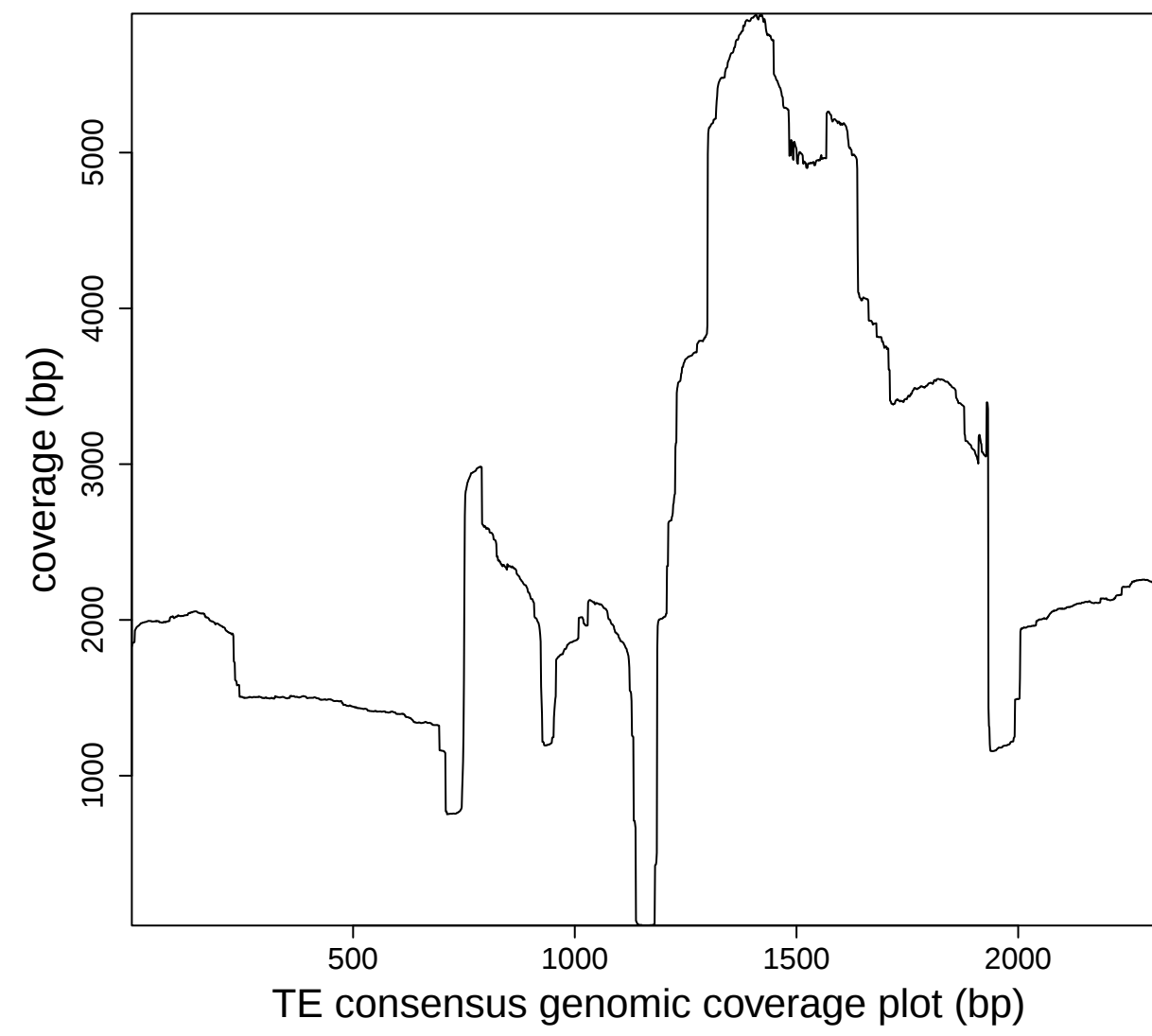
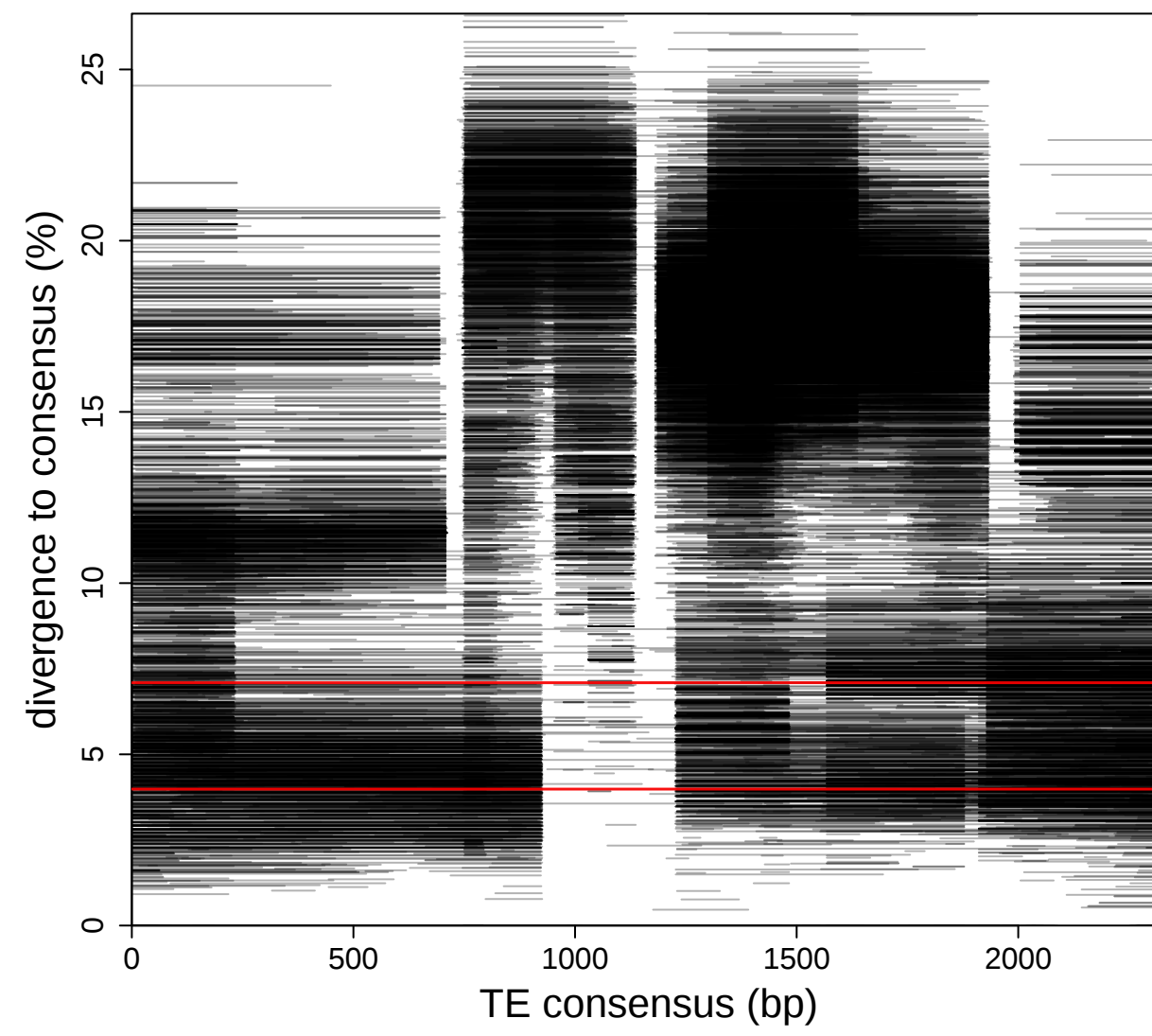
Start crop Point

End crop Point



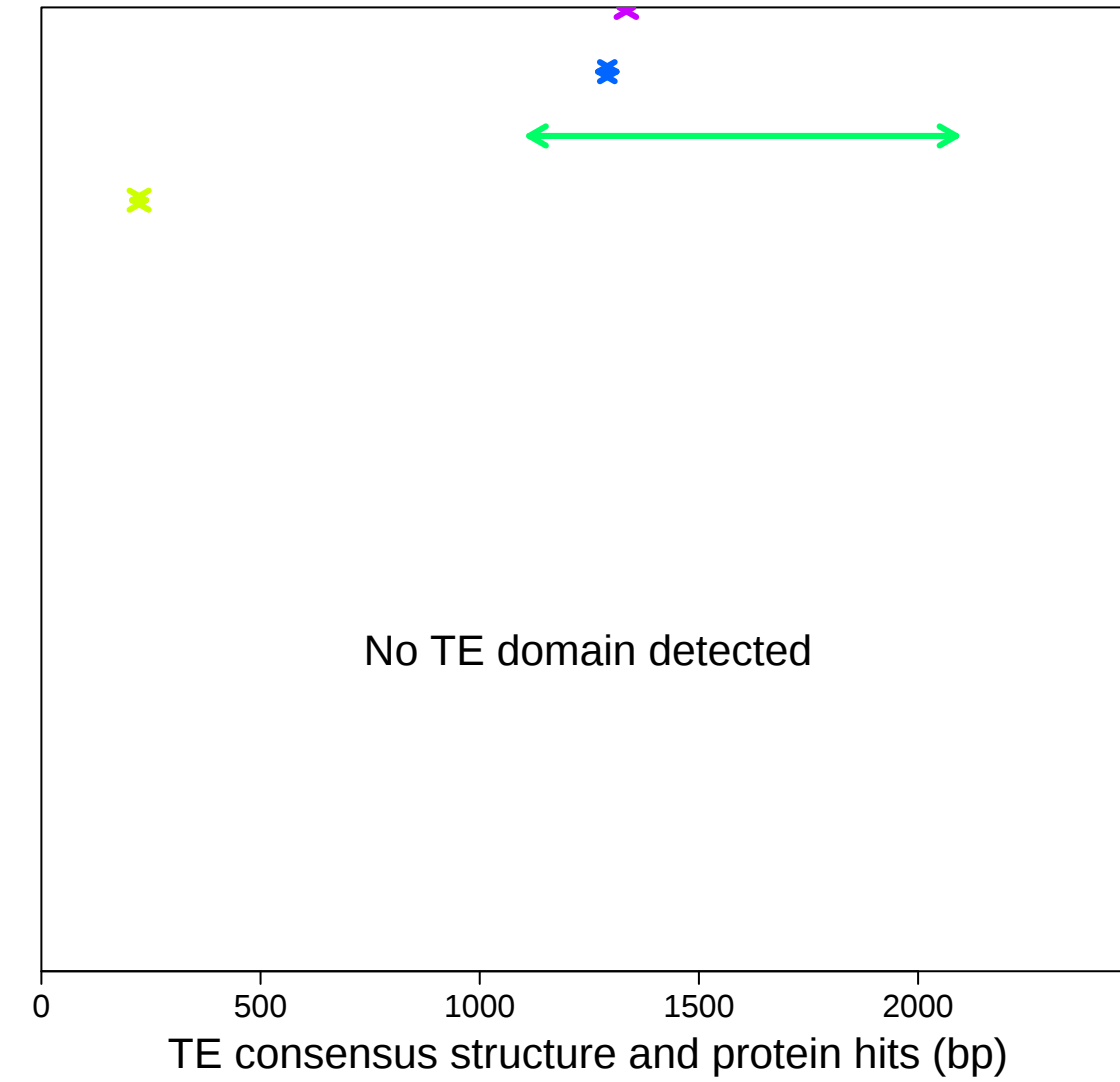
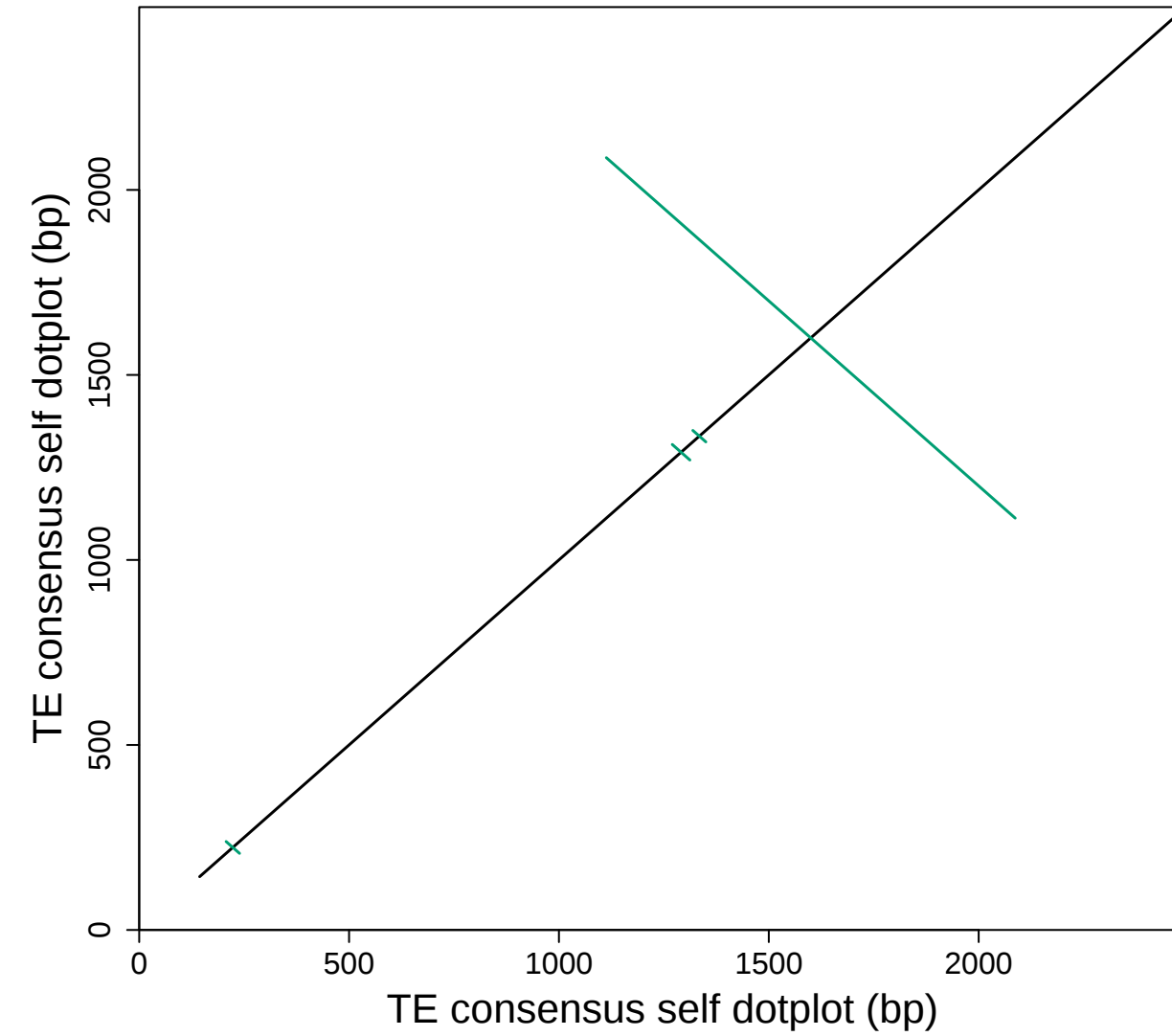
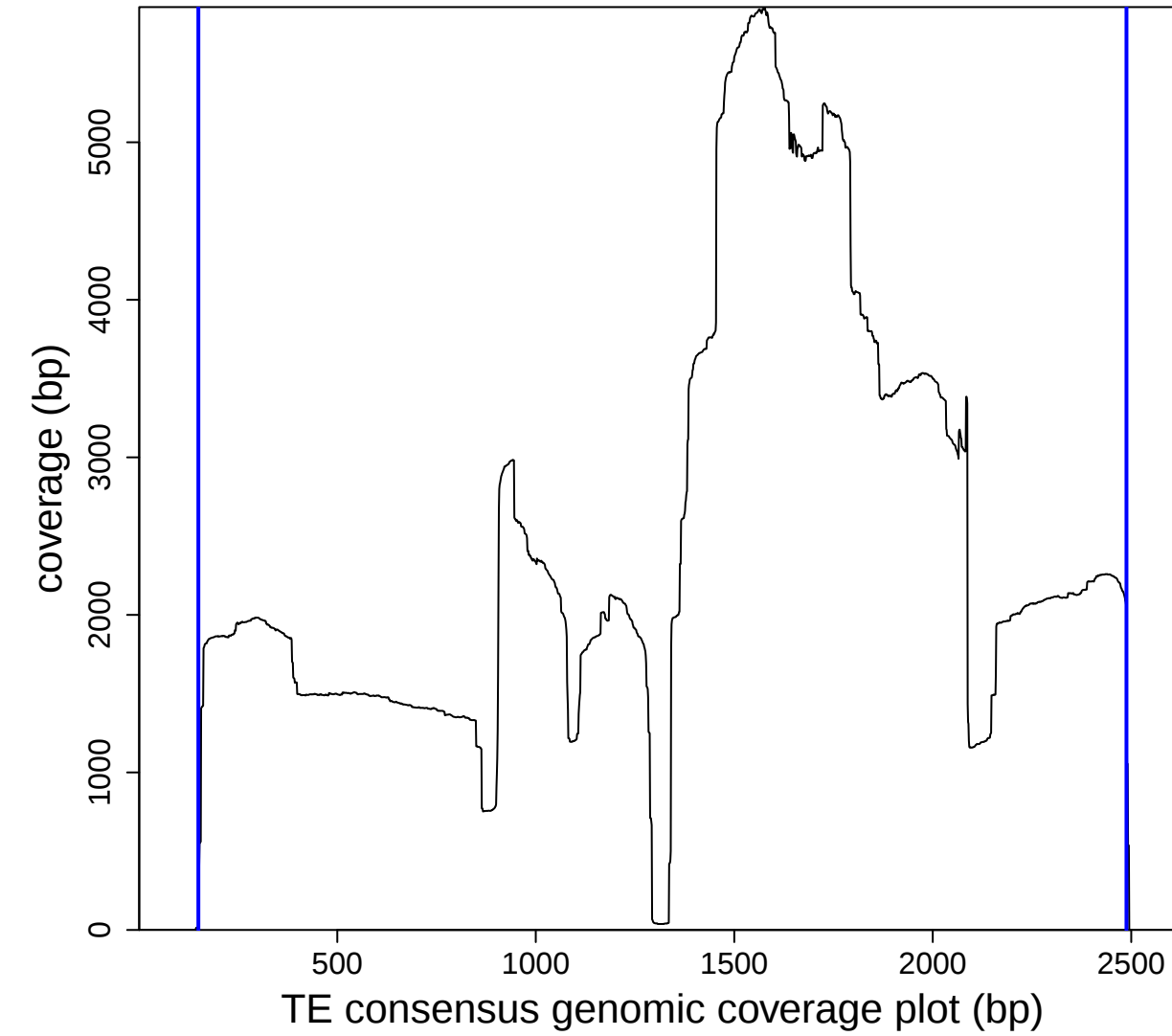
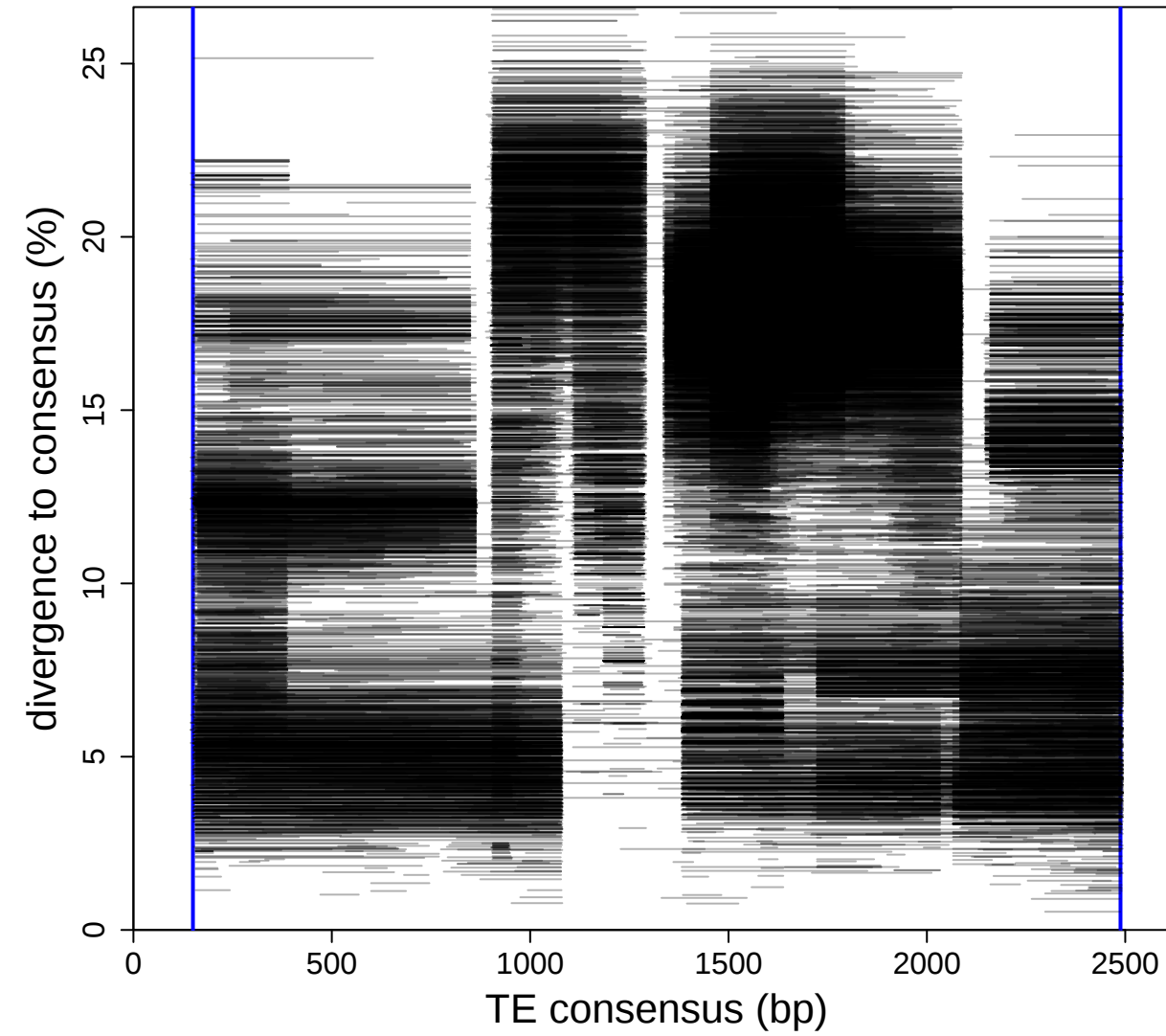
b.bed uf.bed g_3.bed fm_1.bed 0_0 bcIn.fa aln.fa cl.fa gs.fa
size: 2333bp; fragments: 15508; full length: 2 (>=2099.7bp)

After TEtrimmer 2333 bp



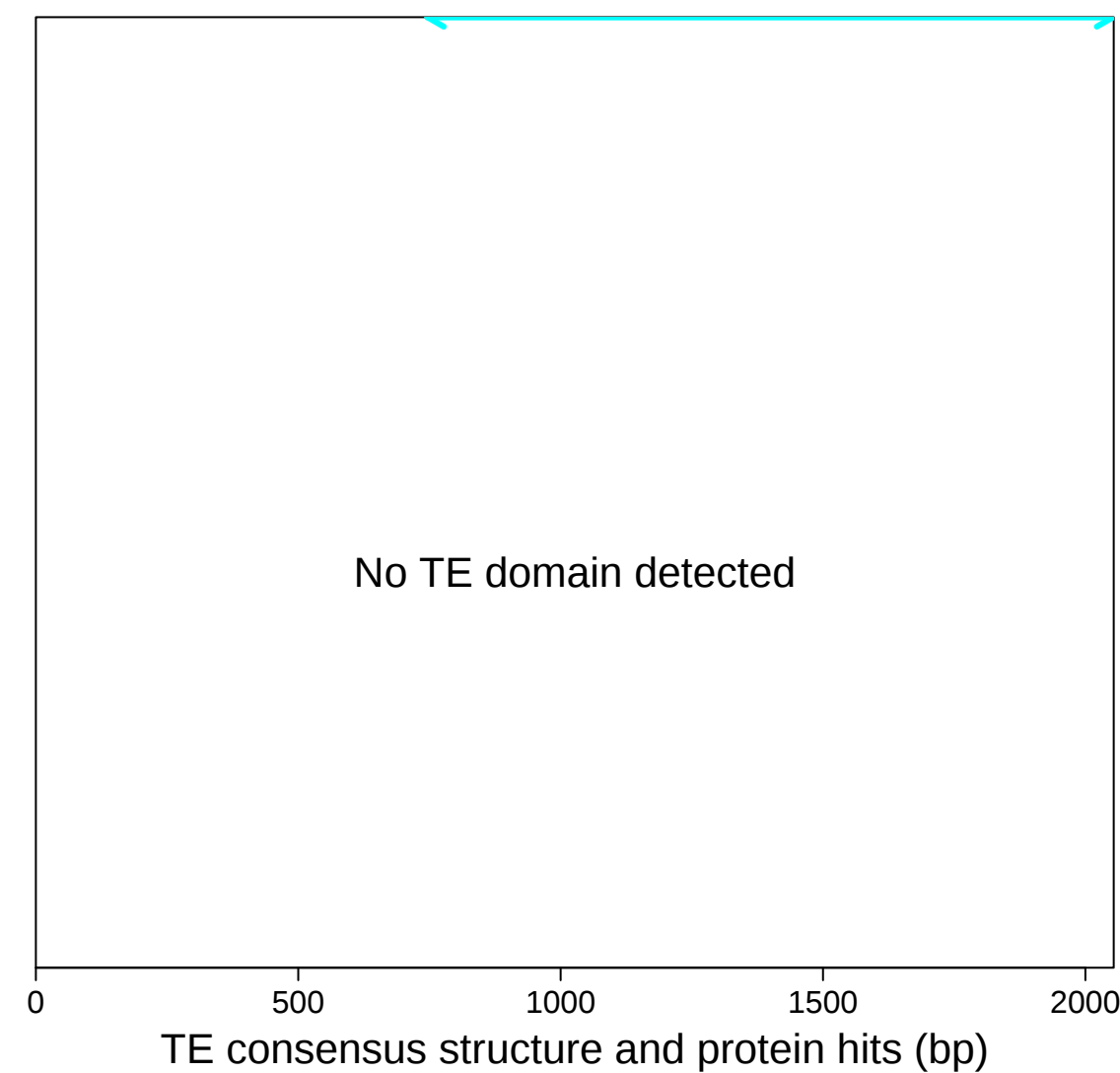
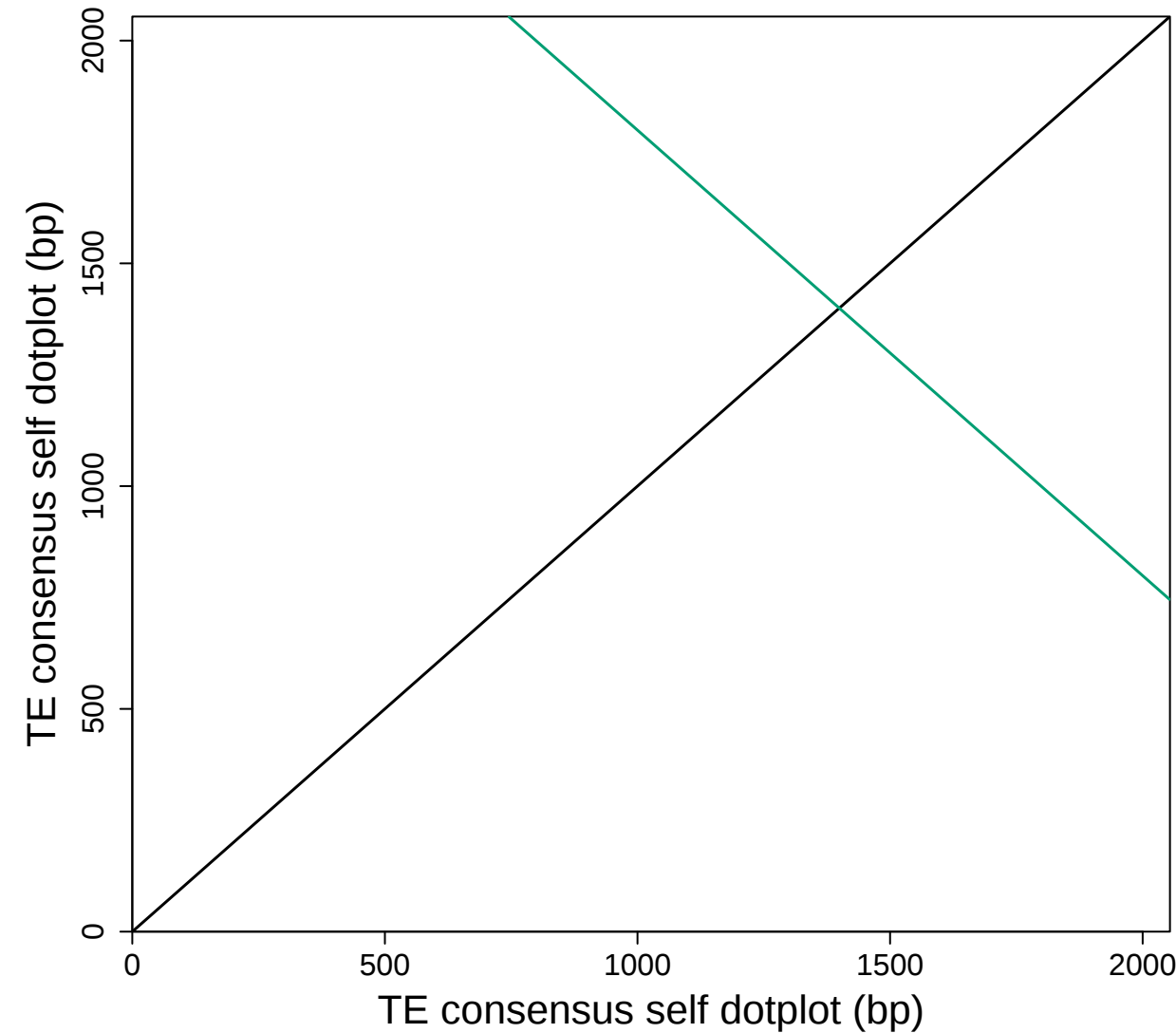
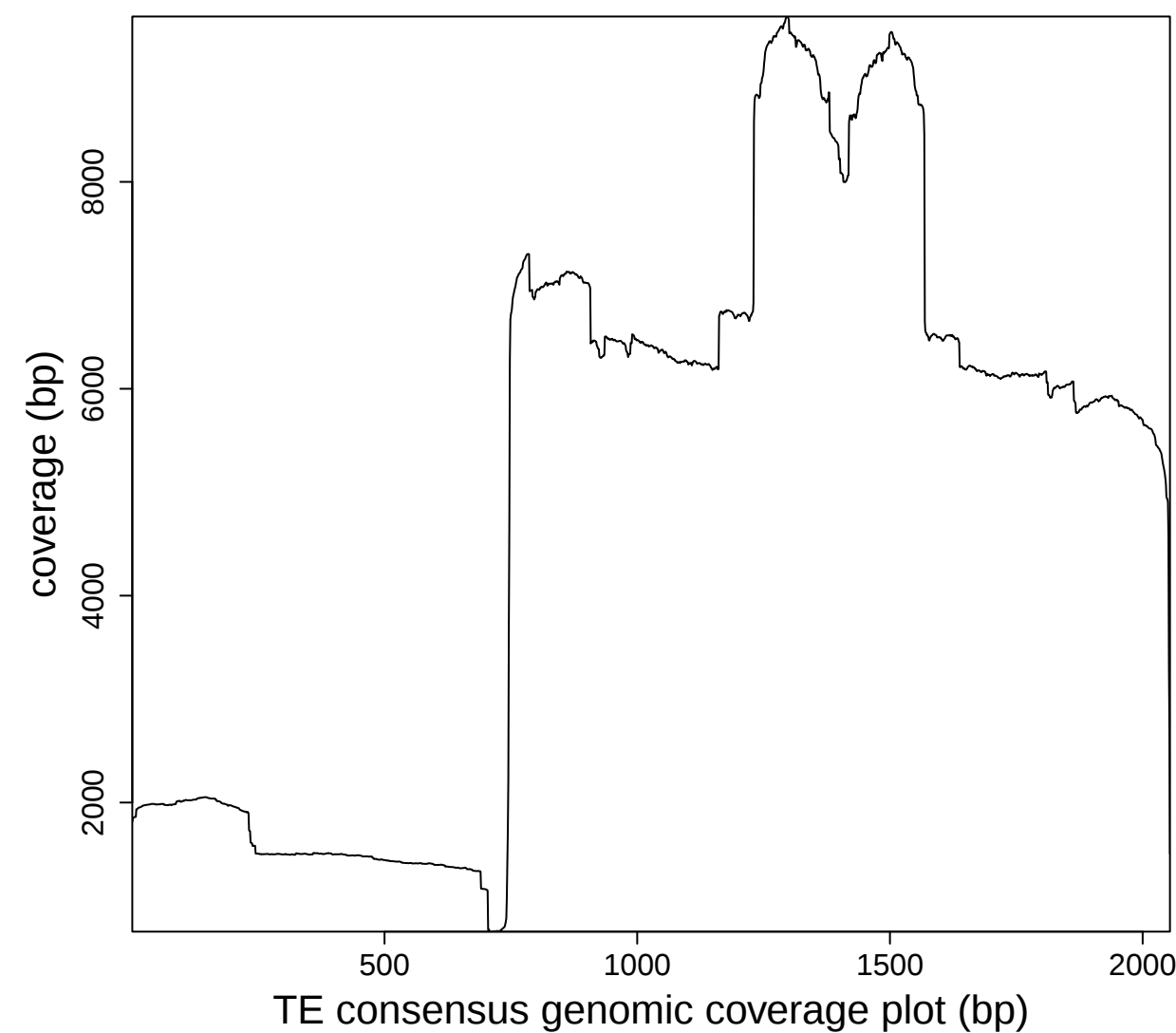
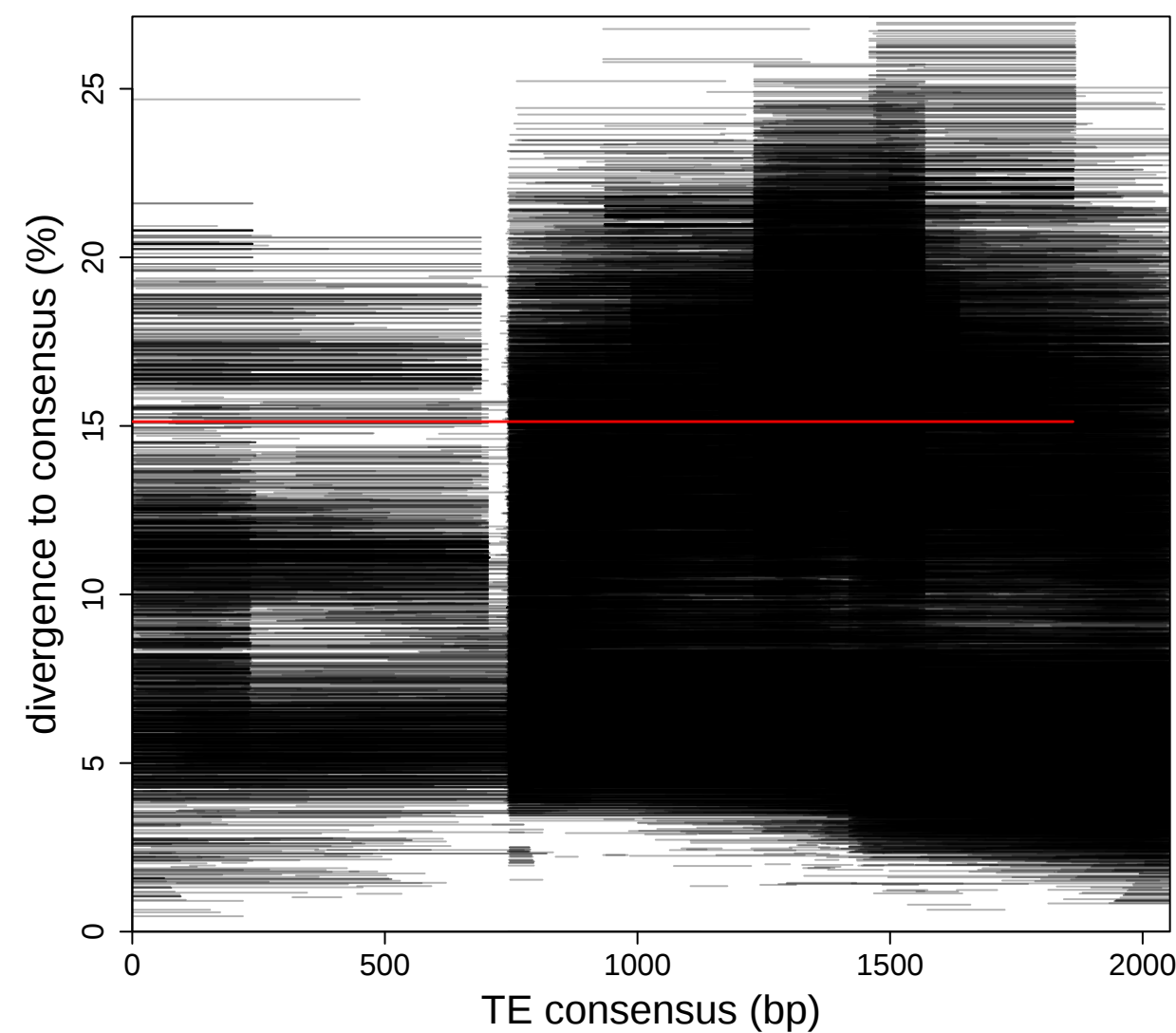
2.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 2638bp; fragments: 15377; full length: 0 (>=2374.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

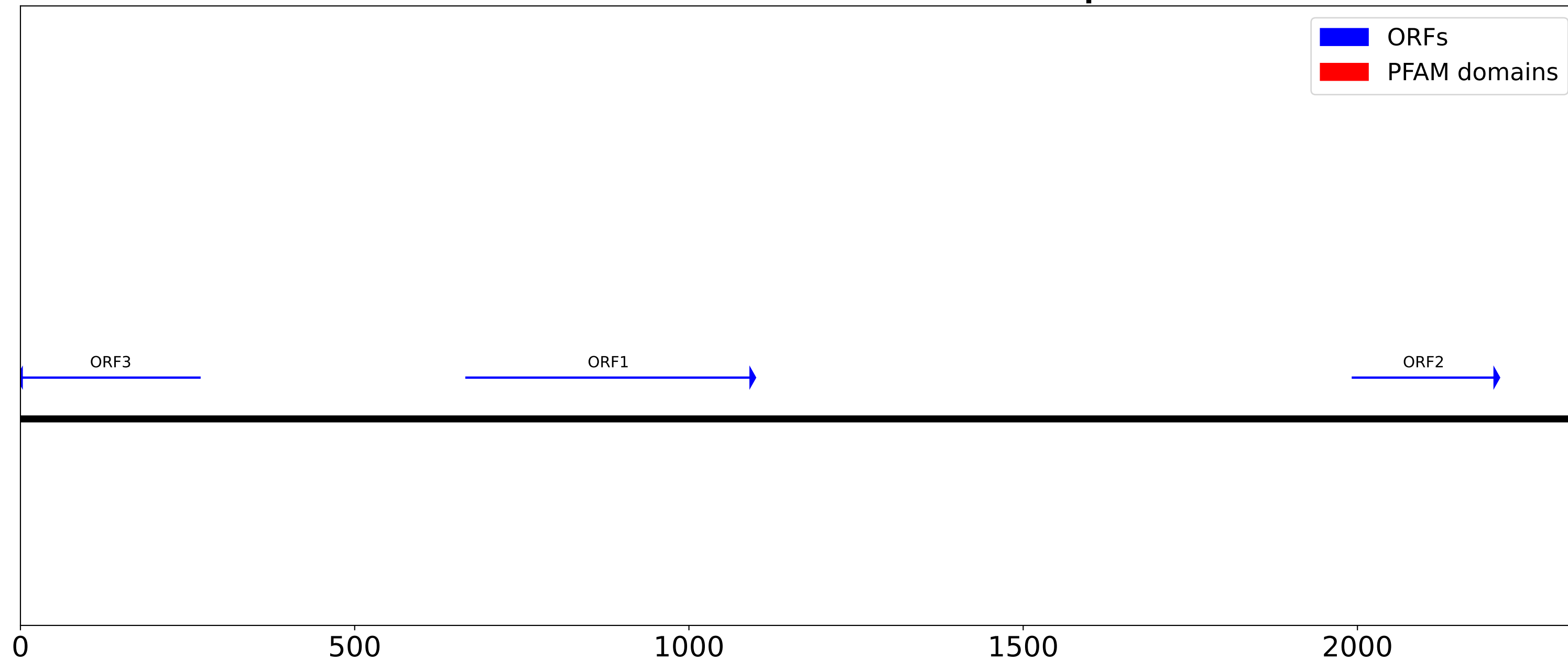


TE: rnd_1_family_22
size: 2054bp; fragments: 21213; full length: 1 (≥ 1848.6 bp)

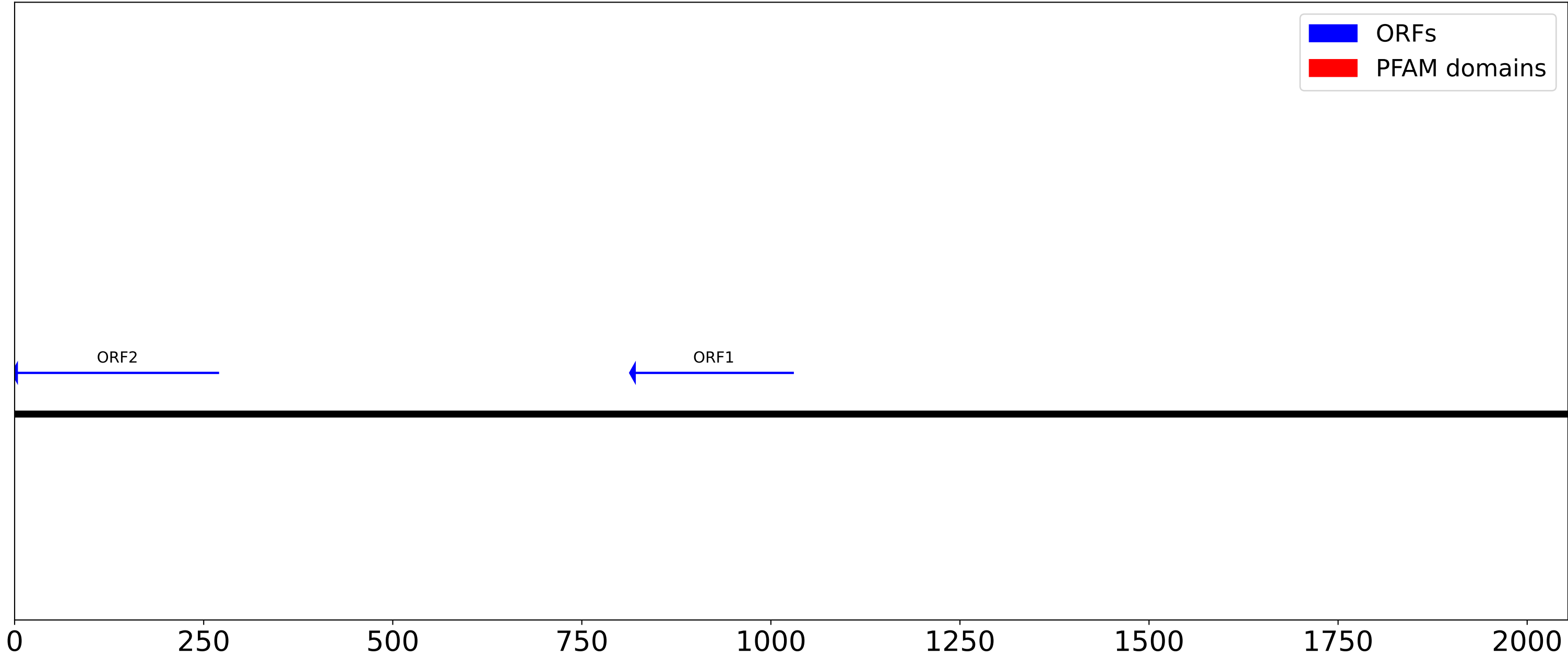
Before TEtrimmer 2054 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

