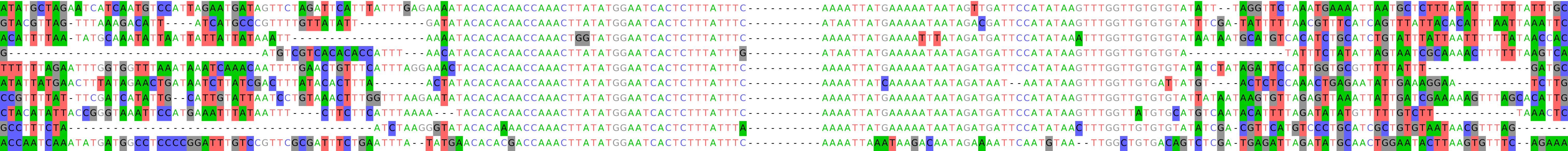


Start Crop Point

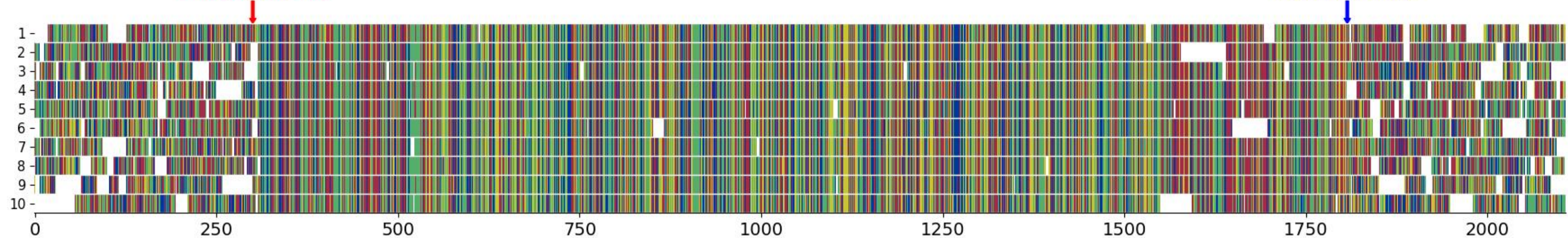
End Crop Point

MSA length = 1508



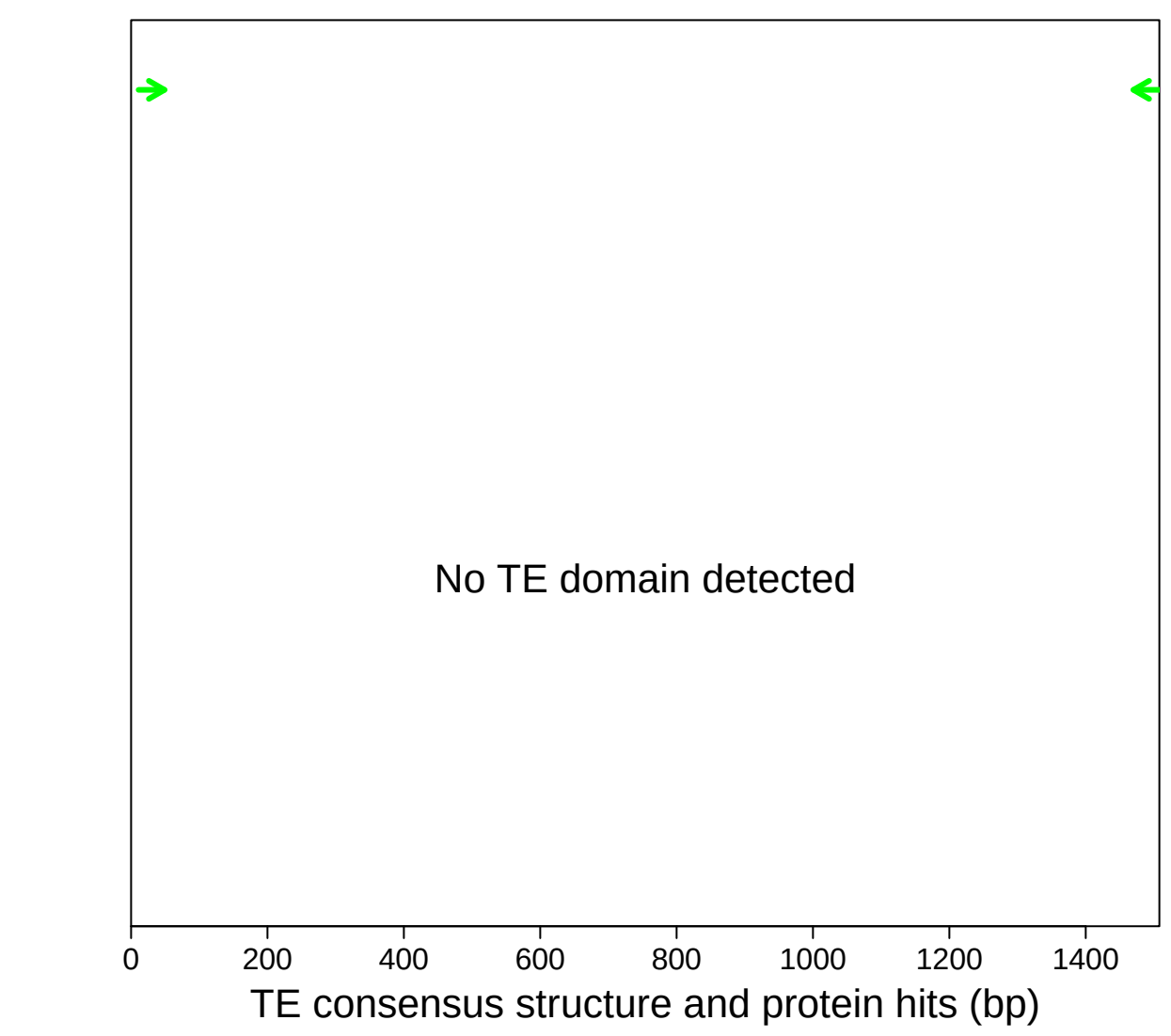
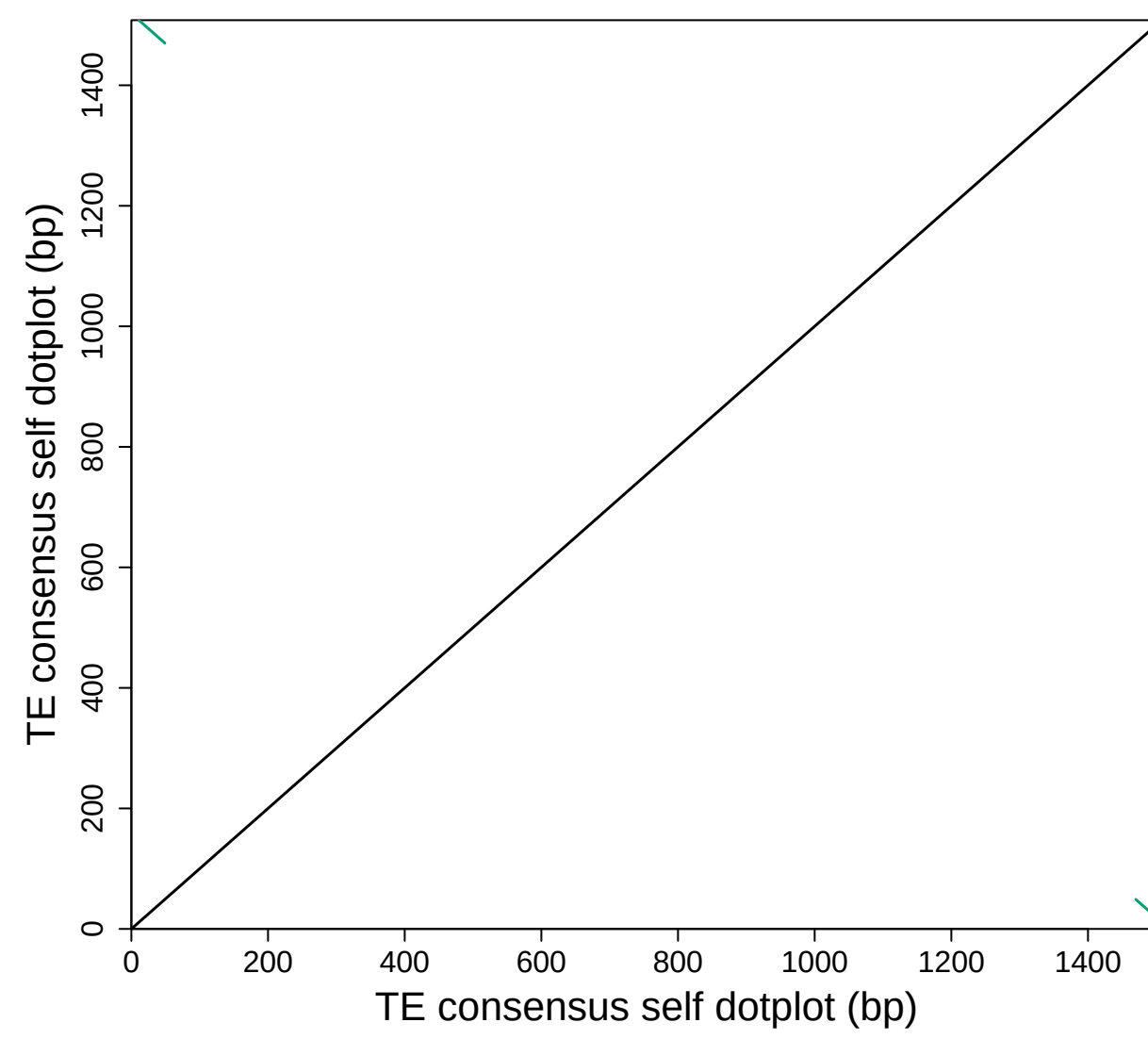
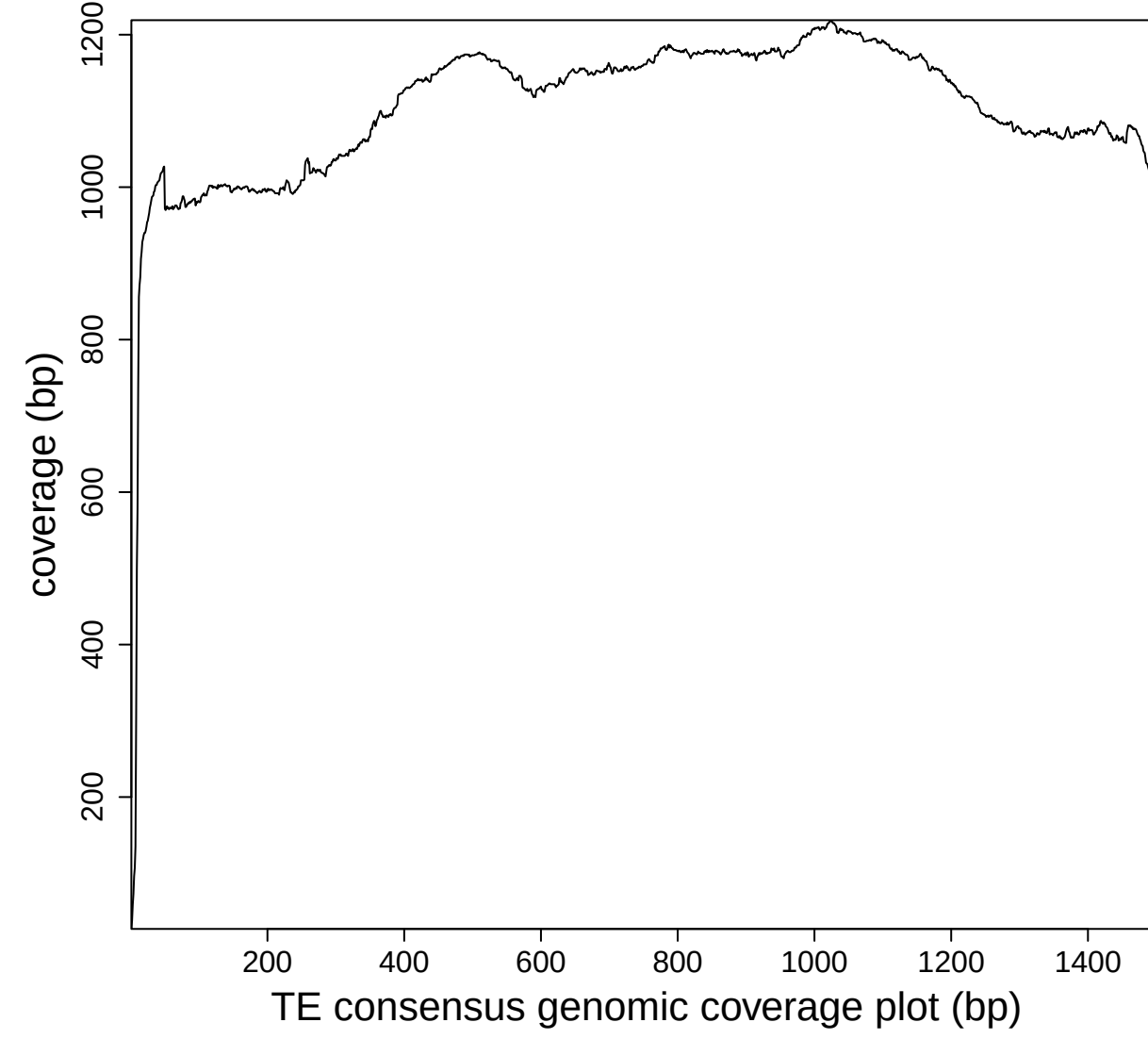
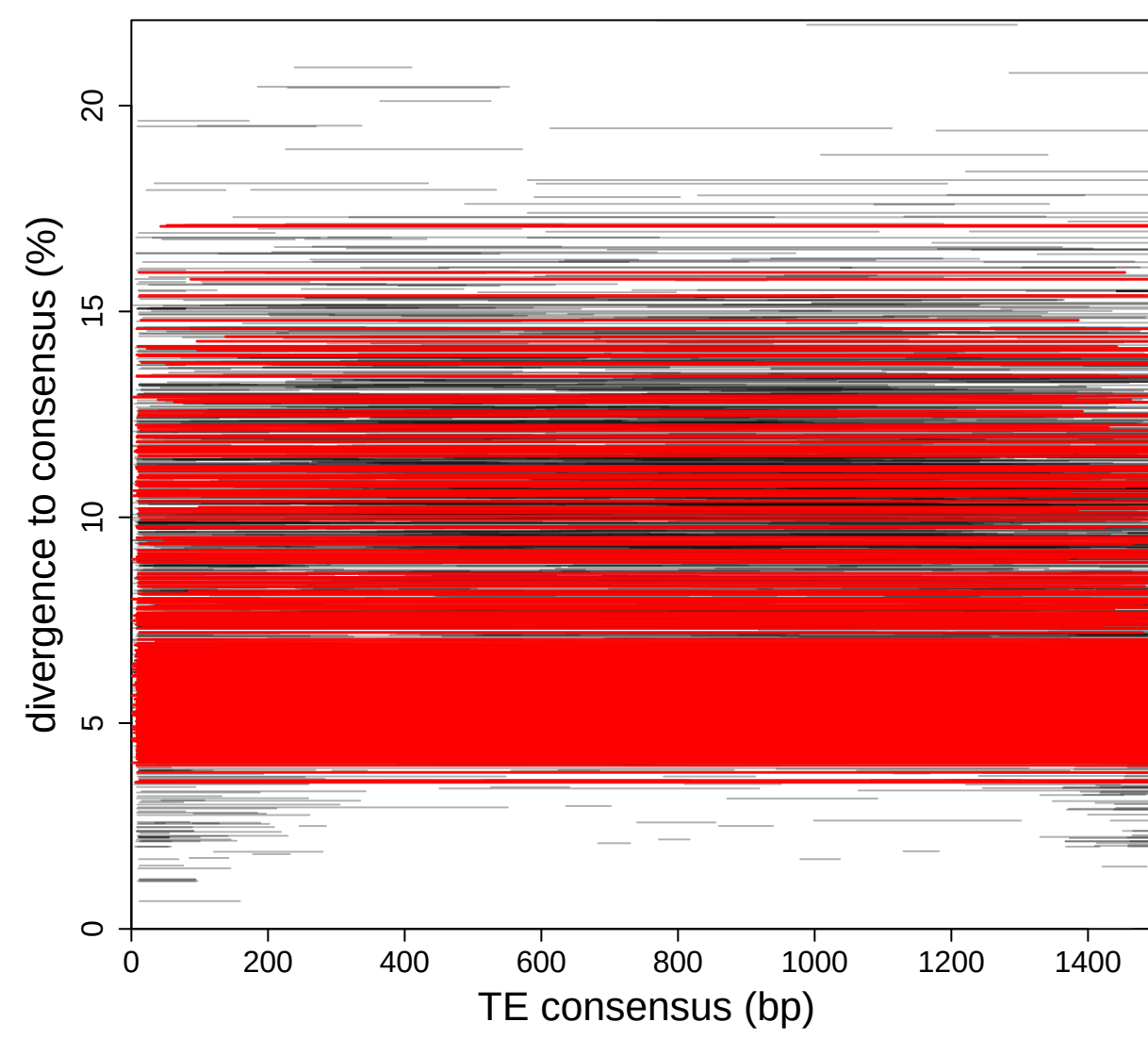
Start crop Point

End crop Point



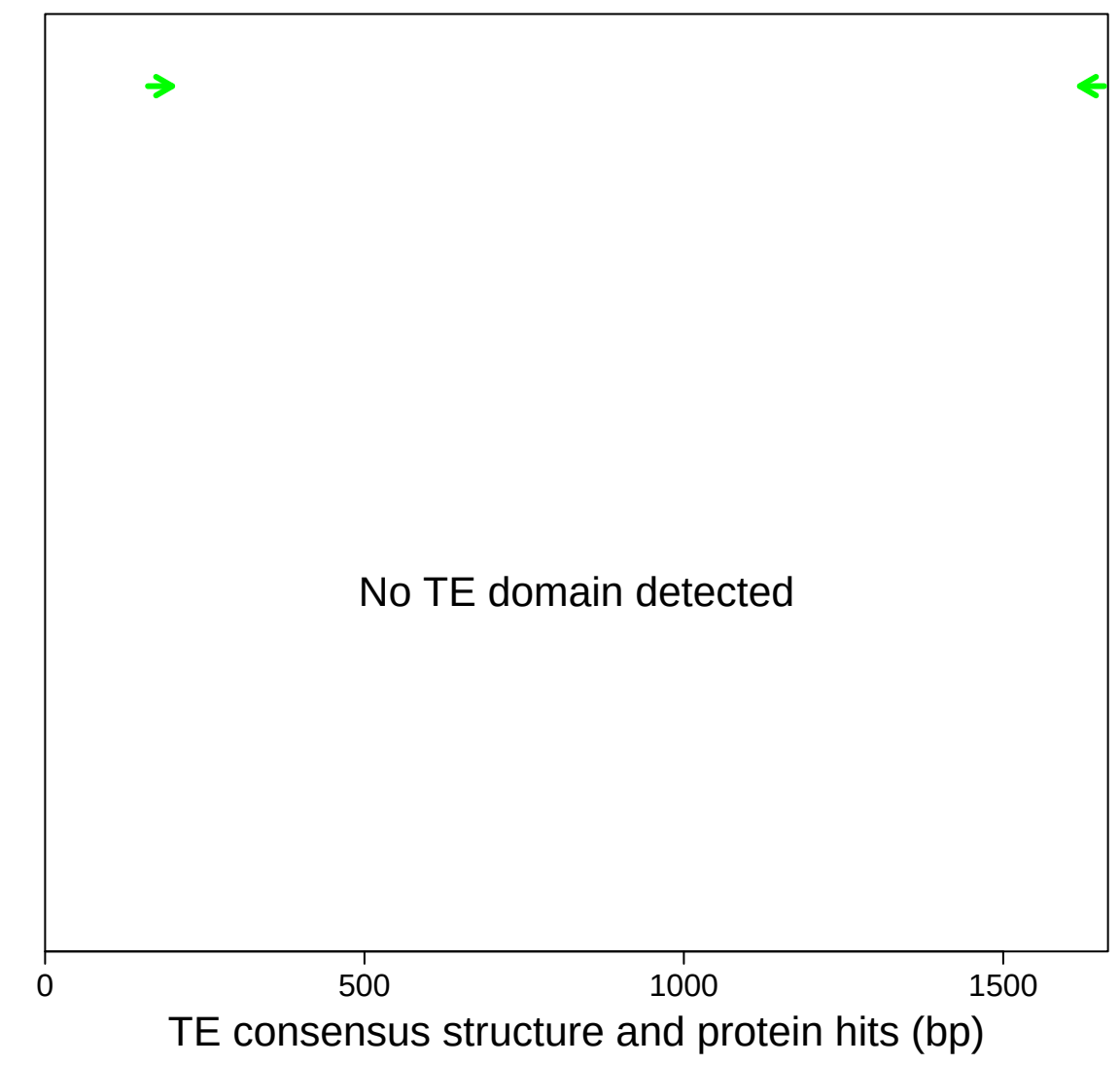
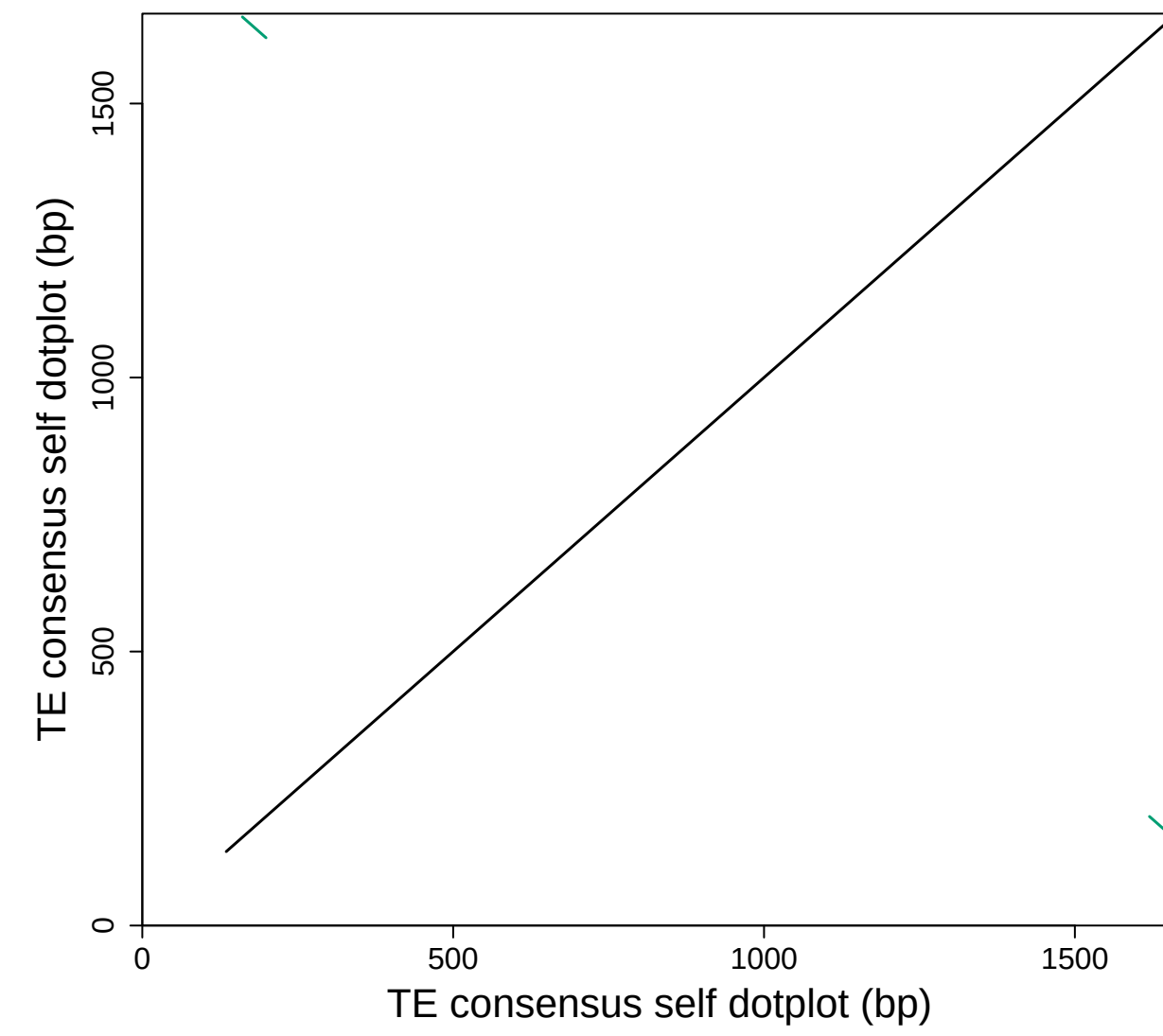
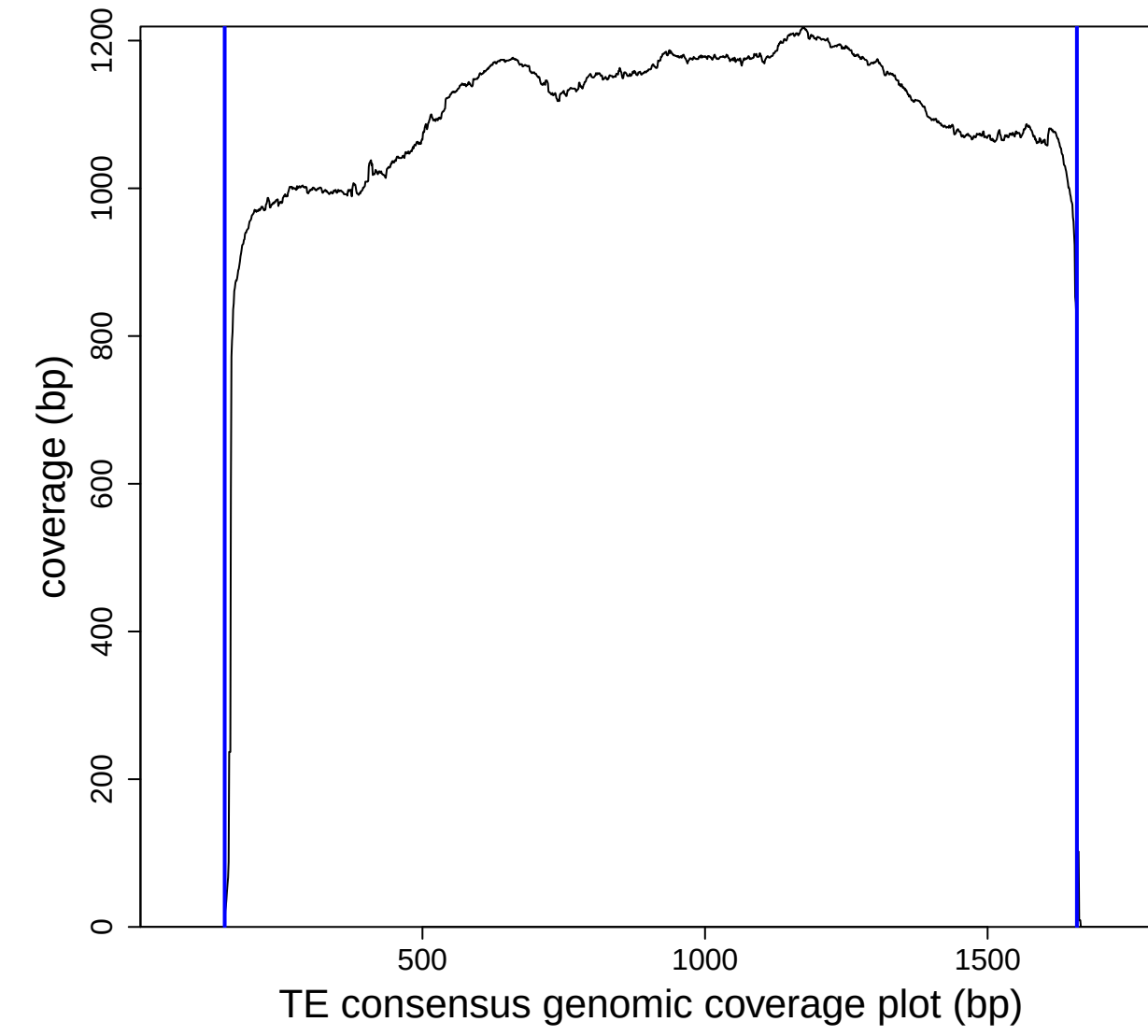
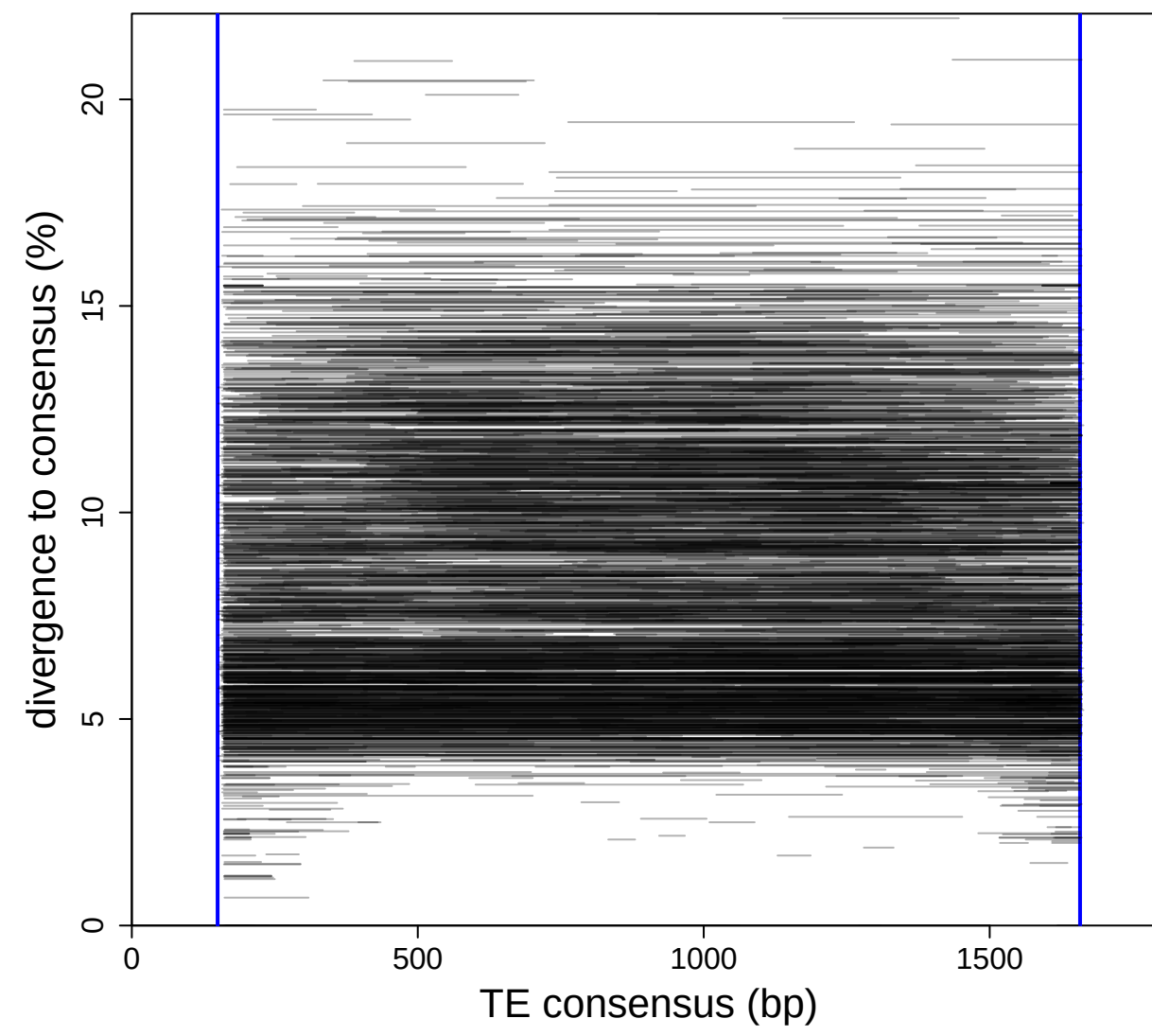
.l.b.bed uf.bed g_4.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 1508bp; fragments: 2719; full length: 377 (≥ 1357.2 bp)

After TEtrimmer 1508 bp



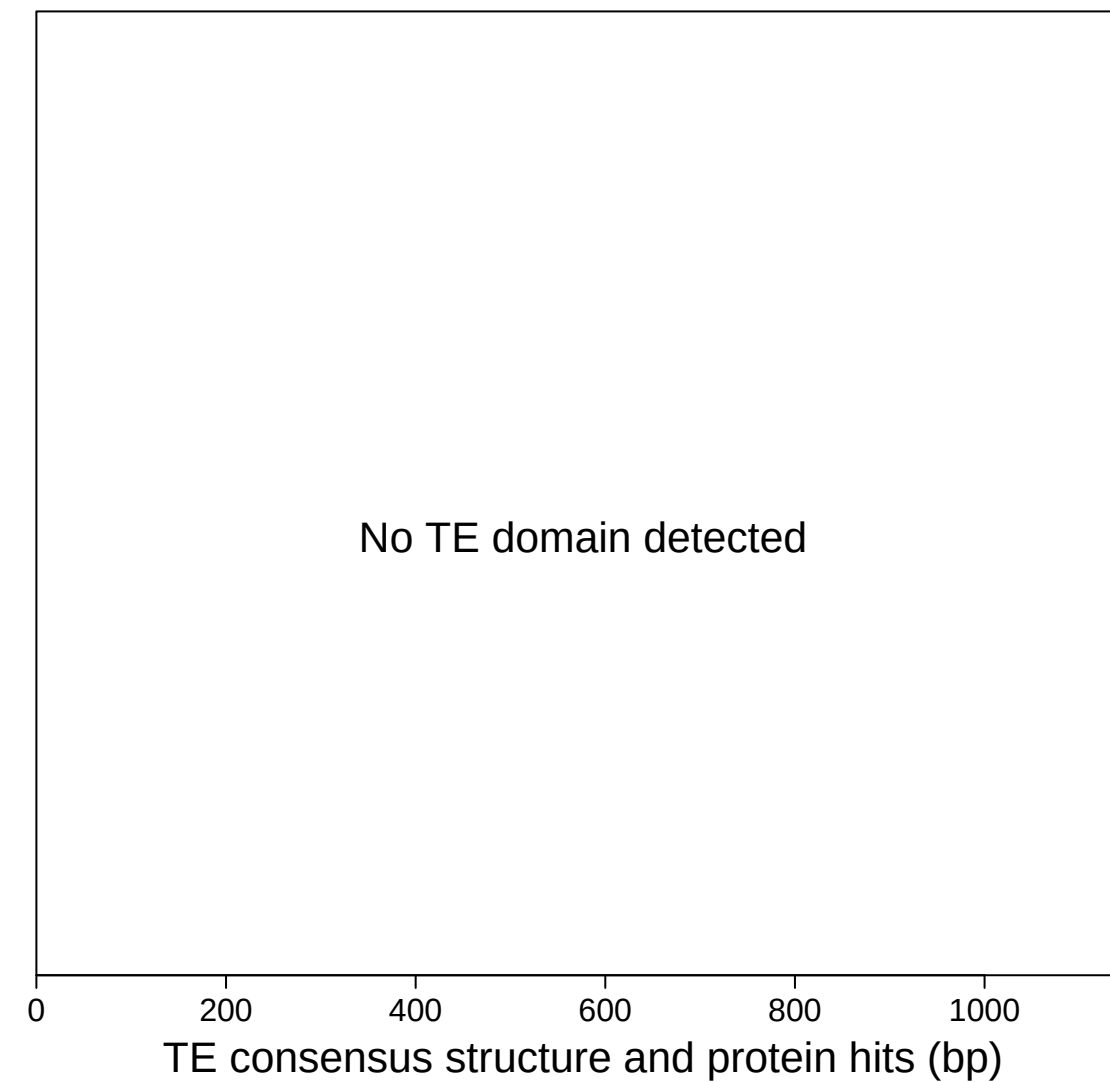
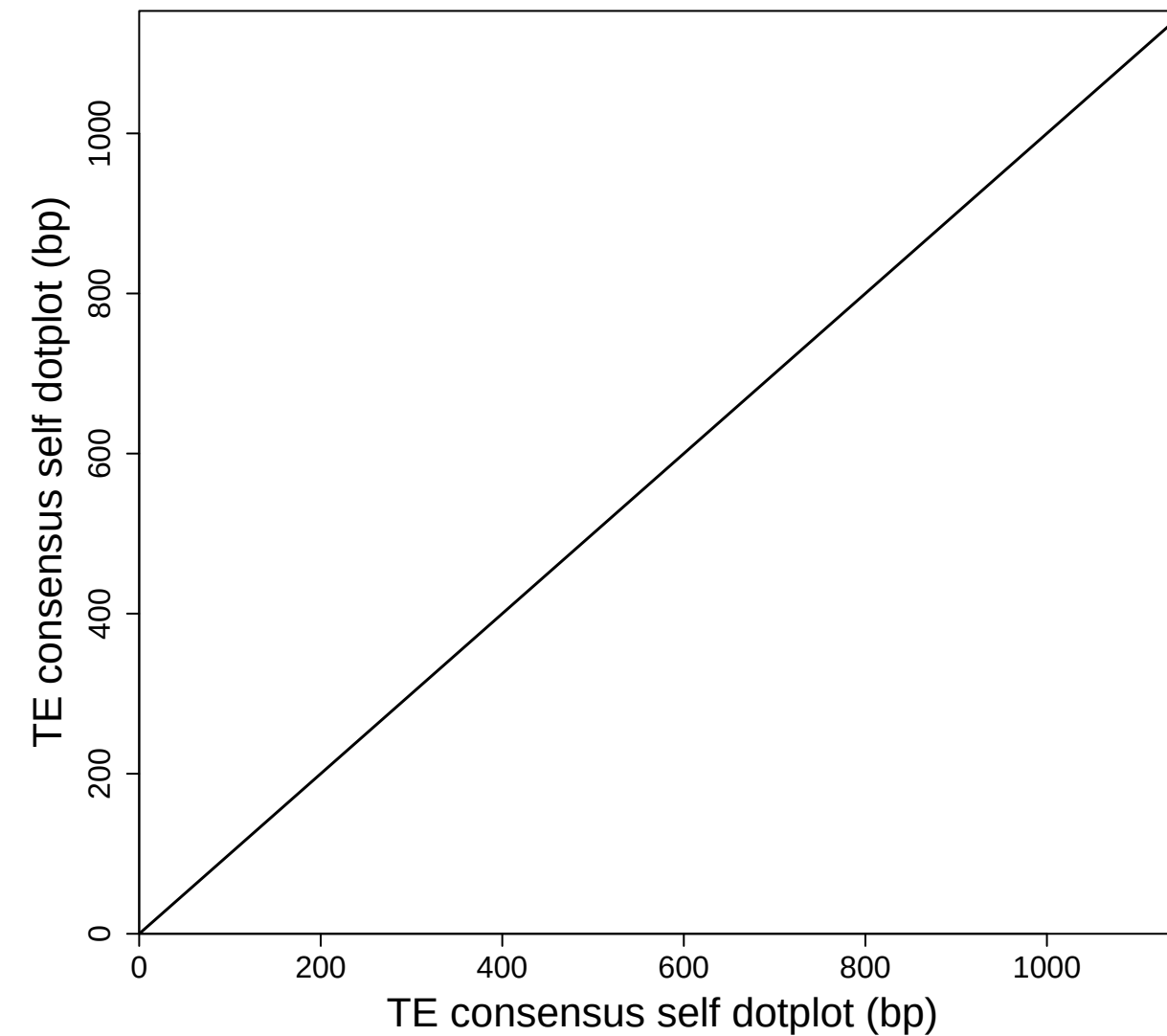
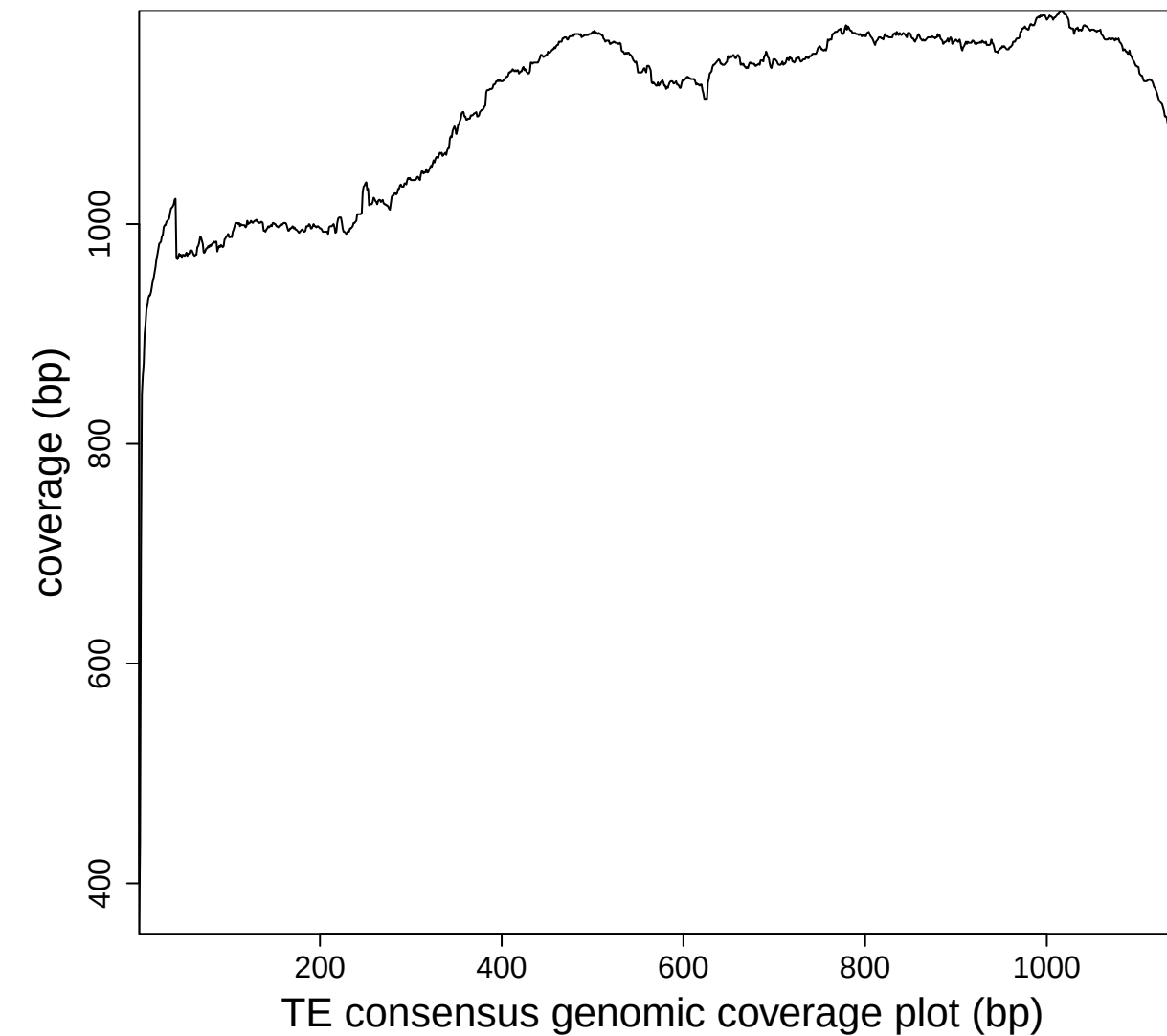
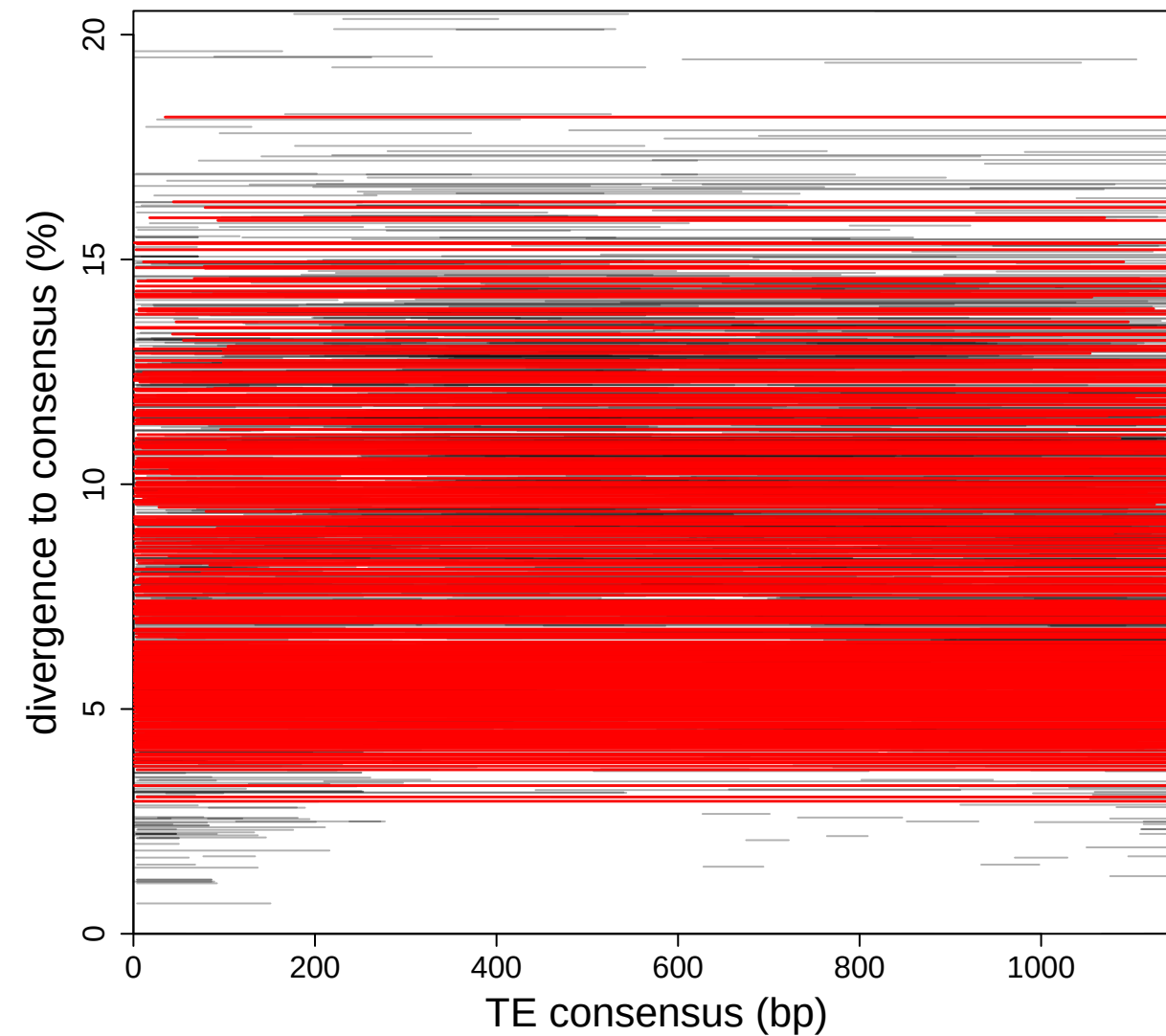
6.fasta.b.bed uf.bed g 4.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 1808bp; fragments: 2656; full length: 0 (>=1627.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

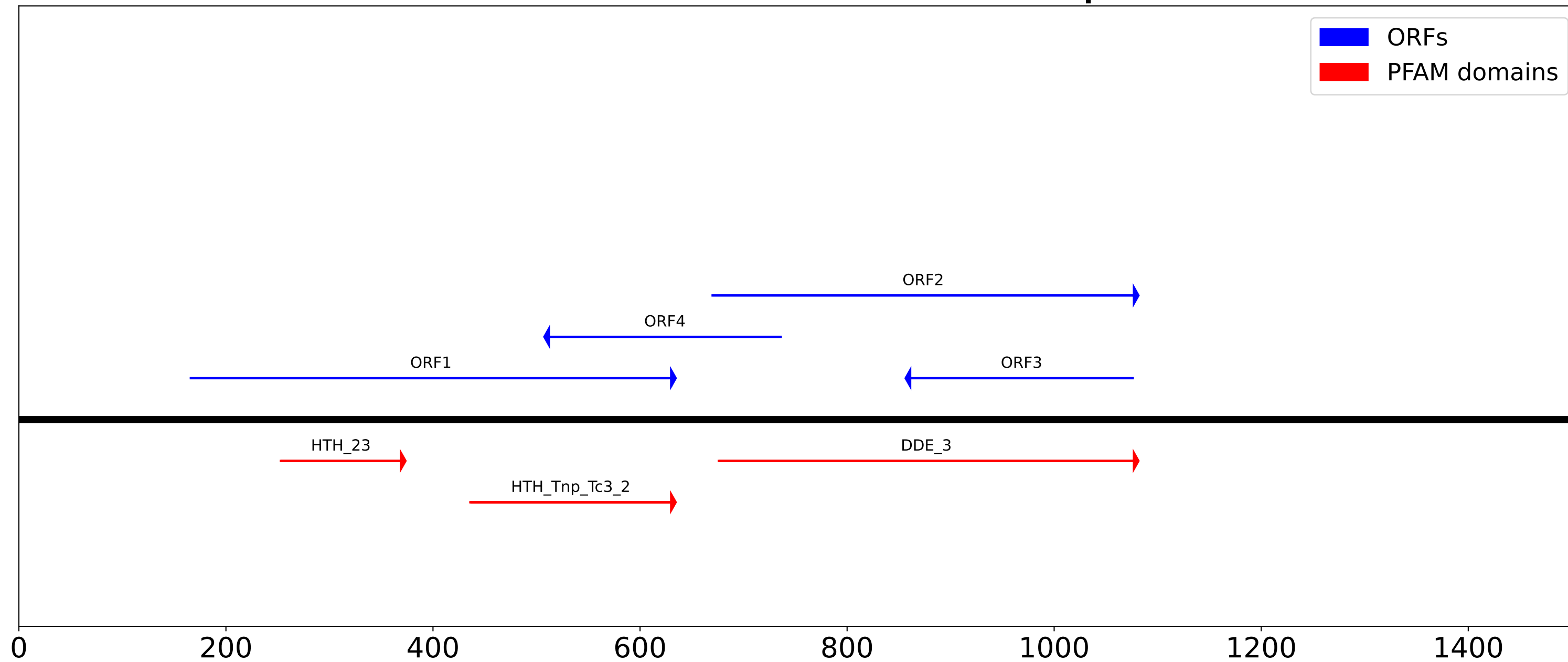


TE: rnd_1_family_466
size: 1153bp; fragments: 2295; full length: 470 (≥ 1037.7 bp)

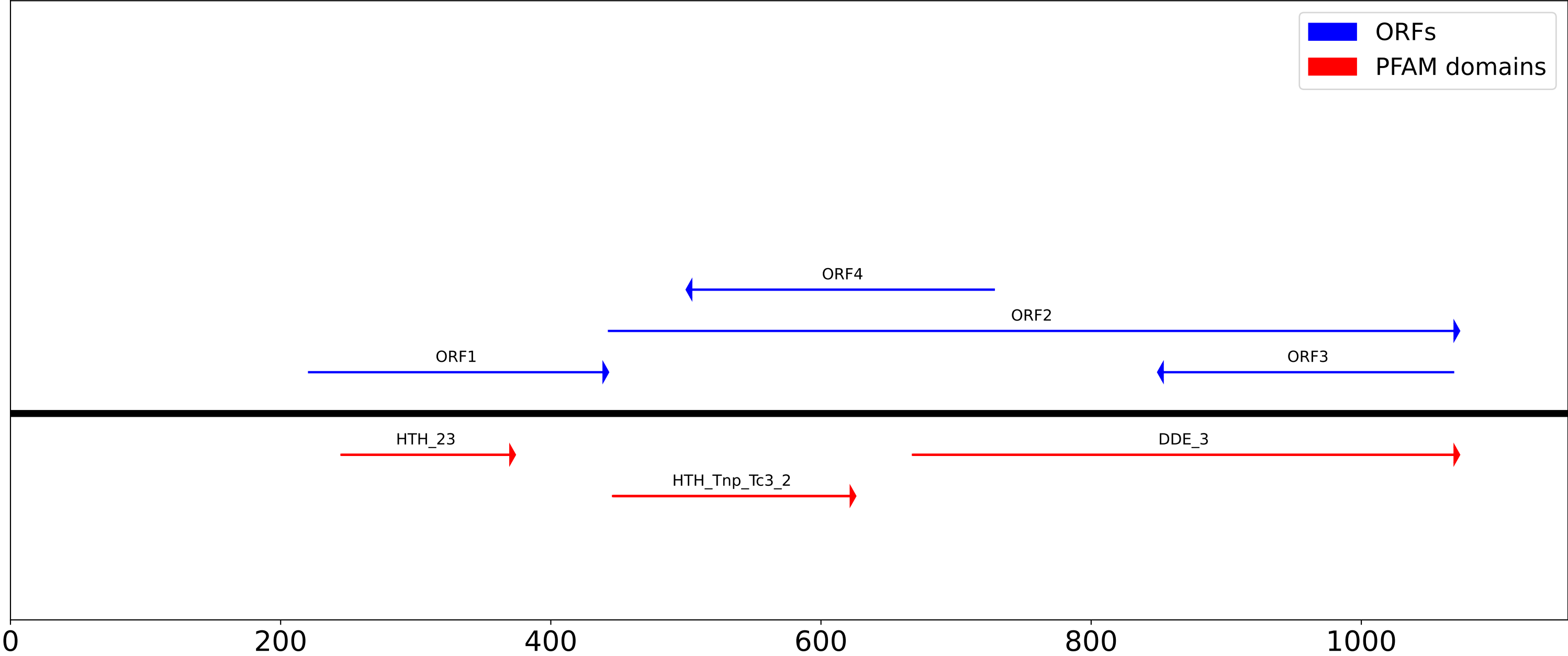
Before TEtrimmer 1153 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

