

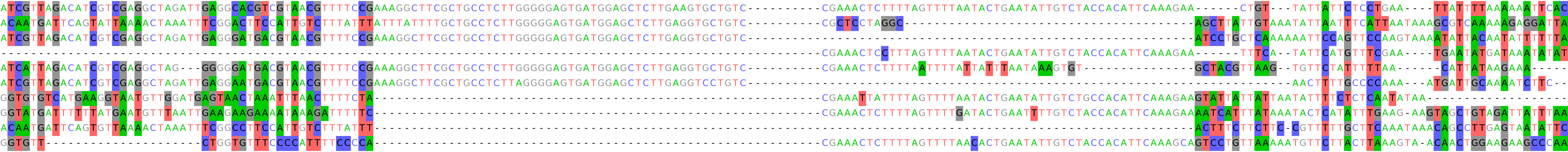
Start Crop Point



End Crop Point



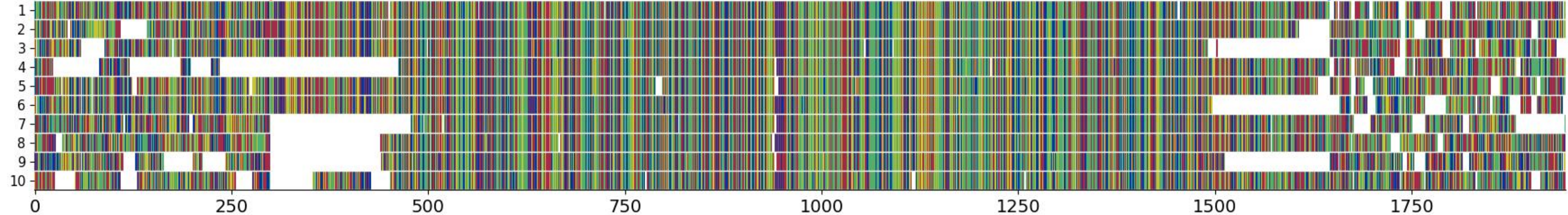
MSA length = 1346



Start crop Point

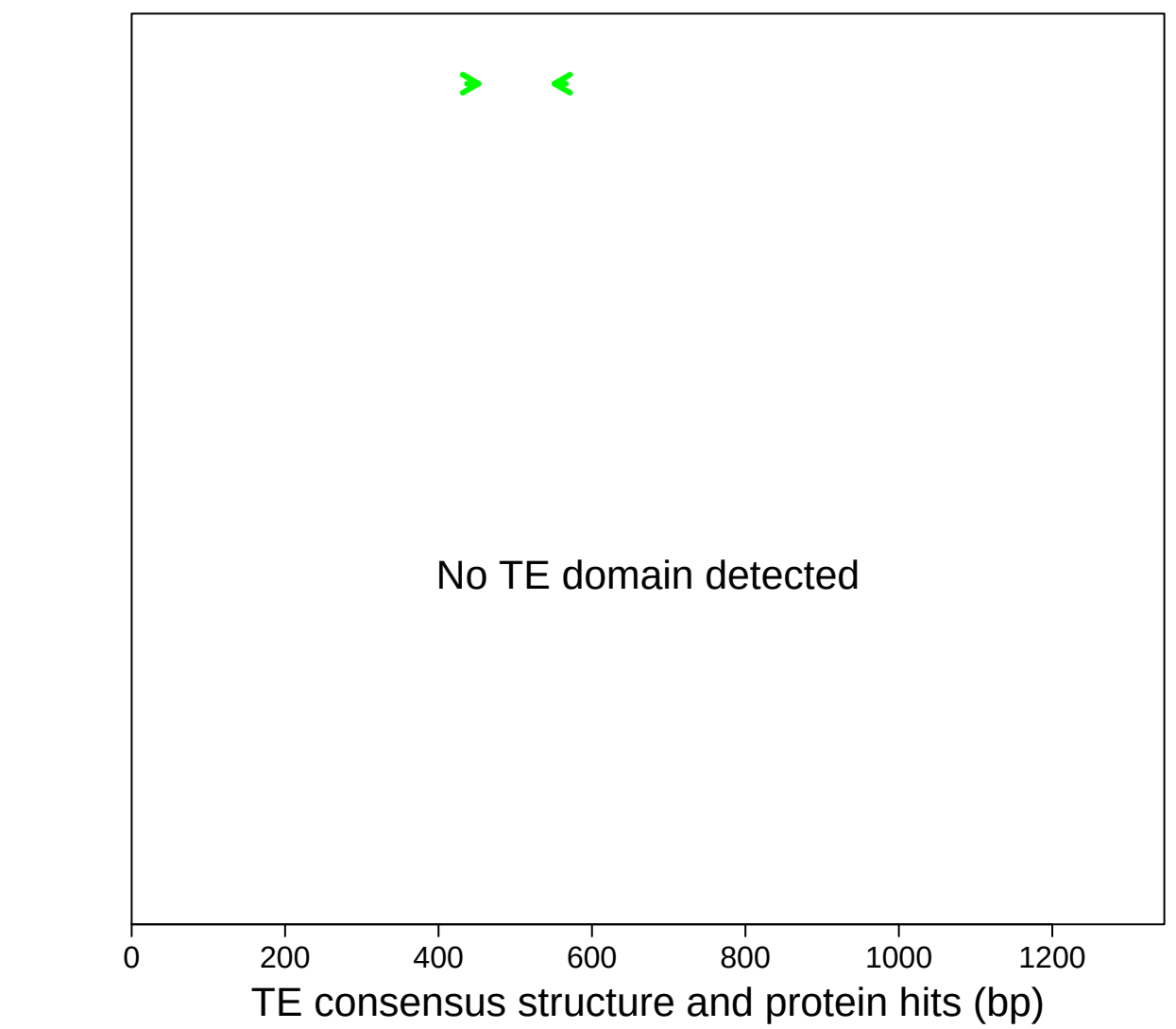
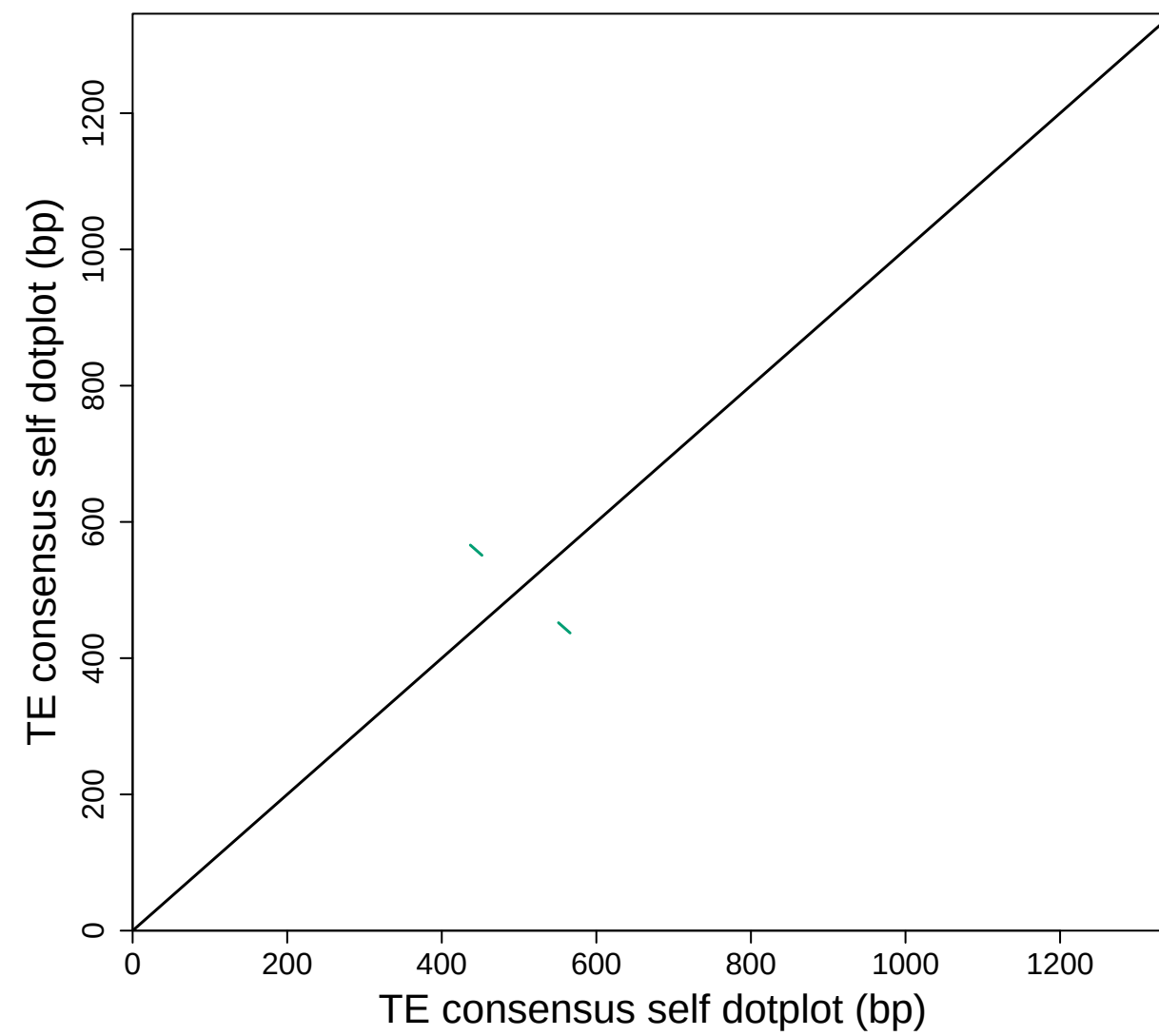
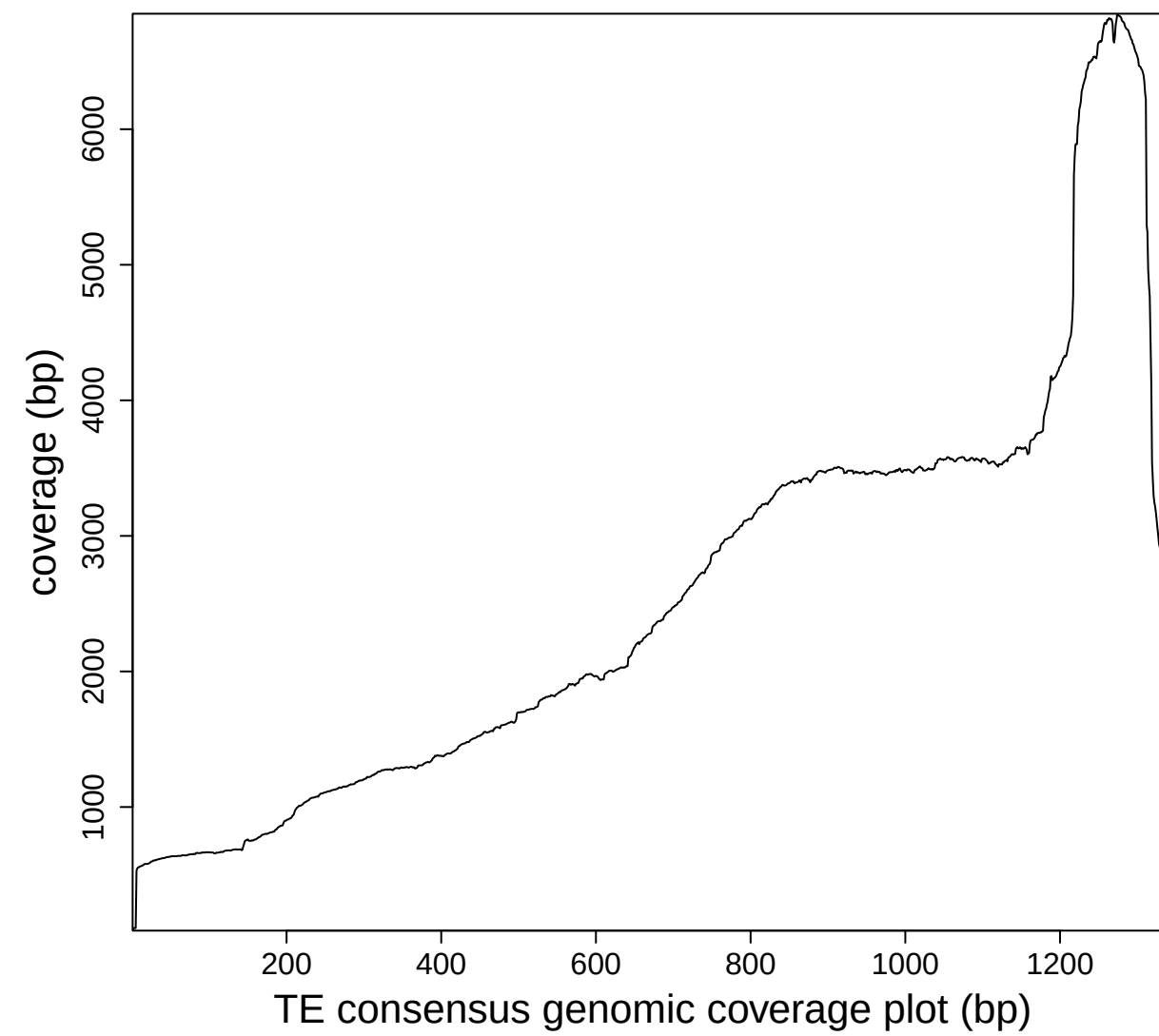
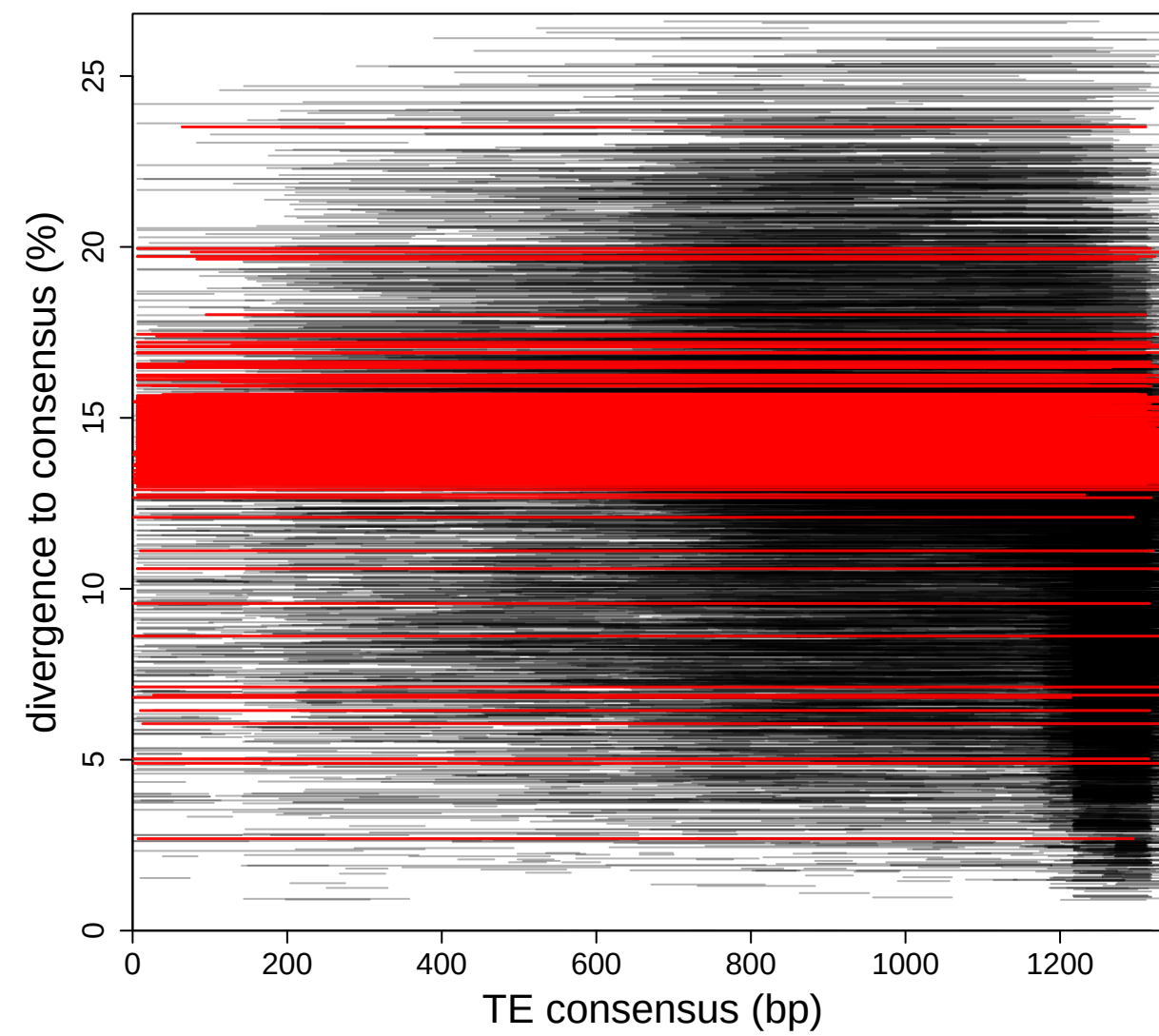


End crop Point



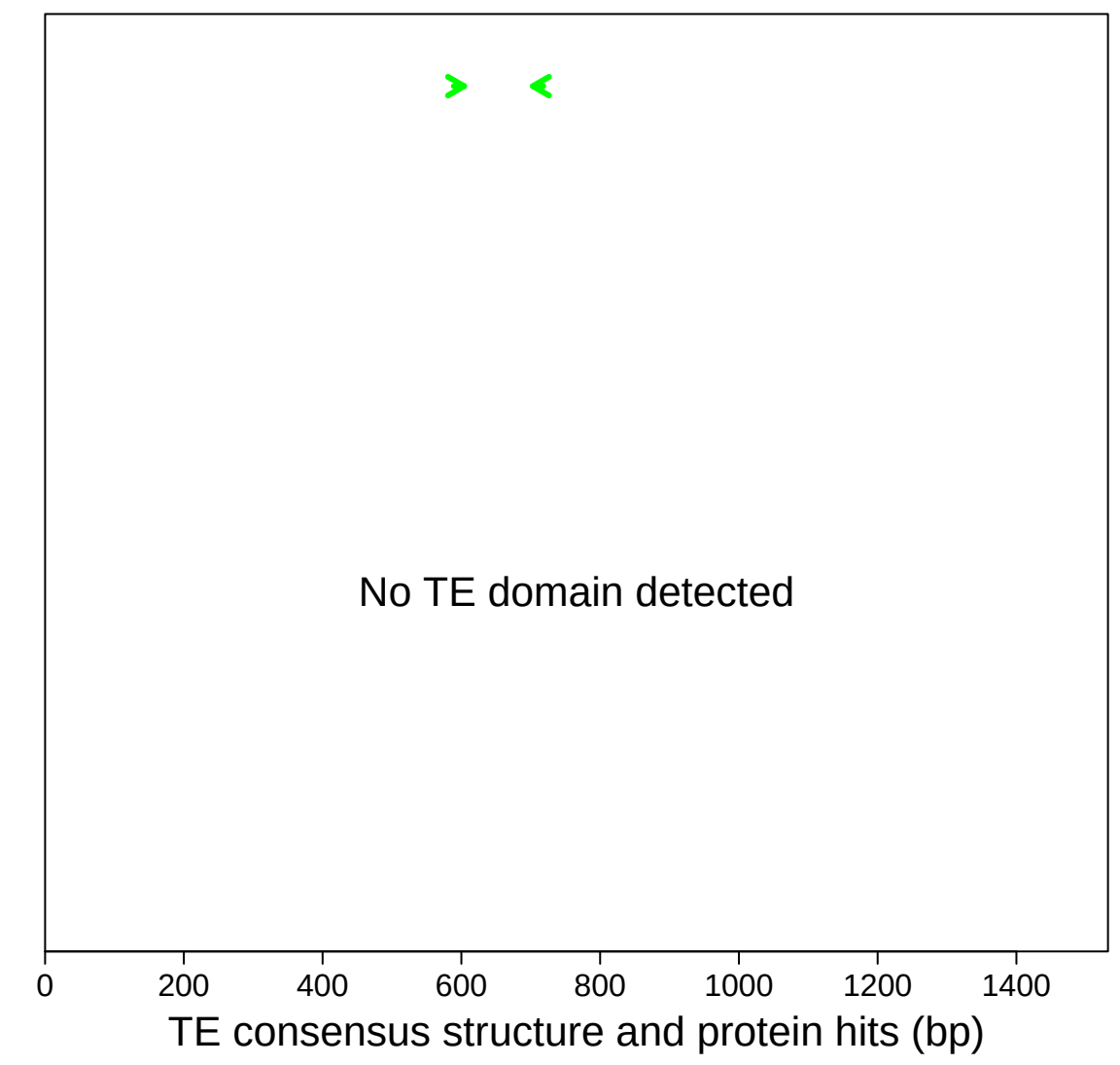
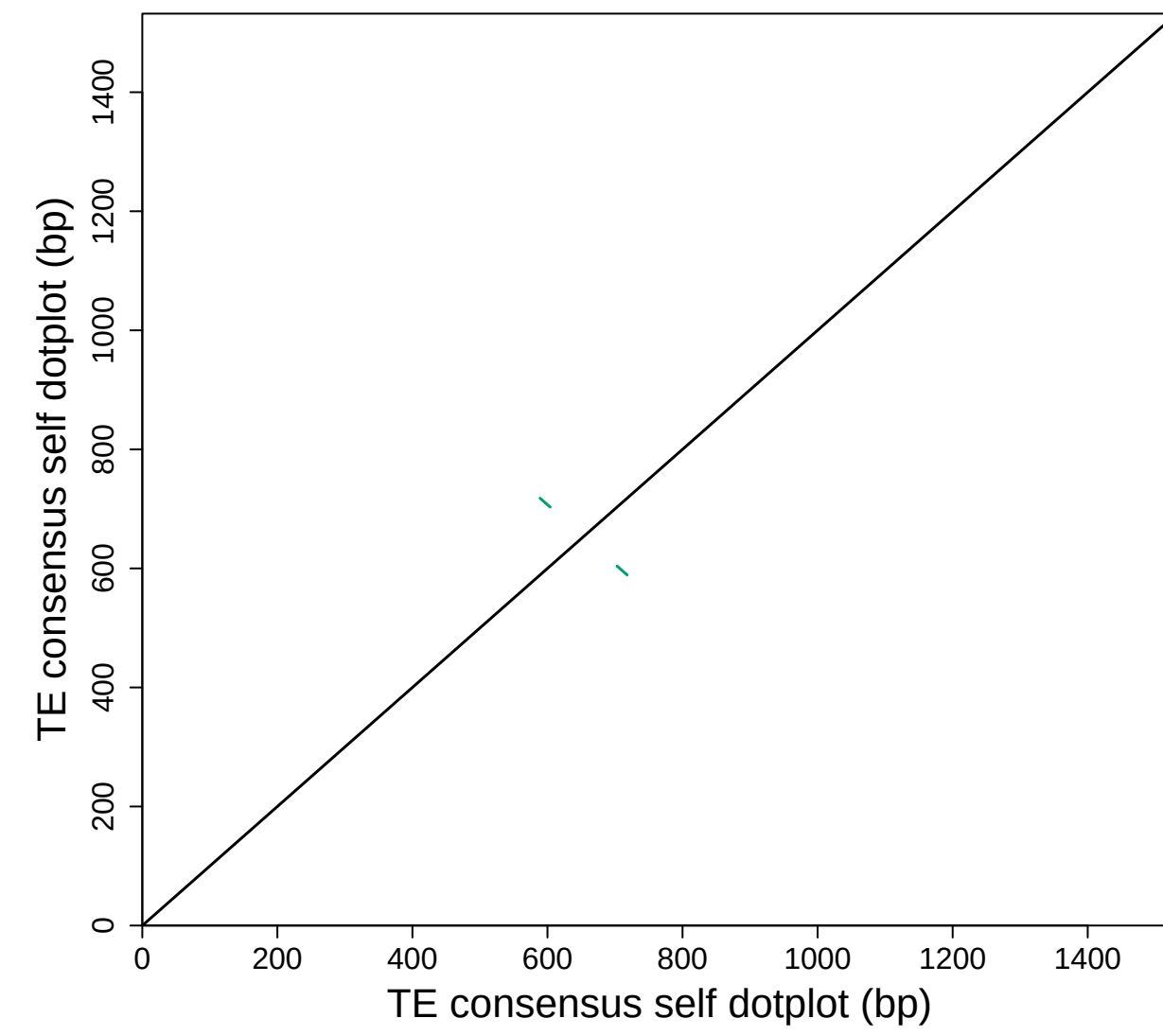
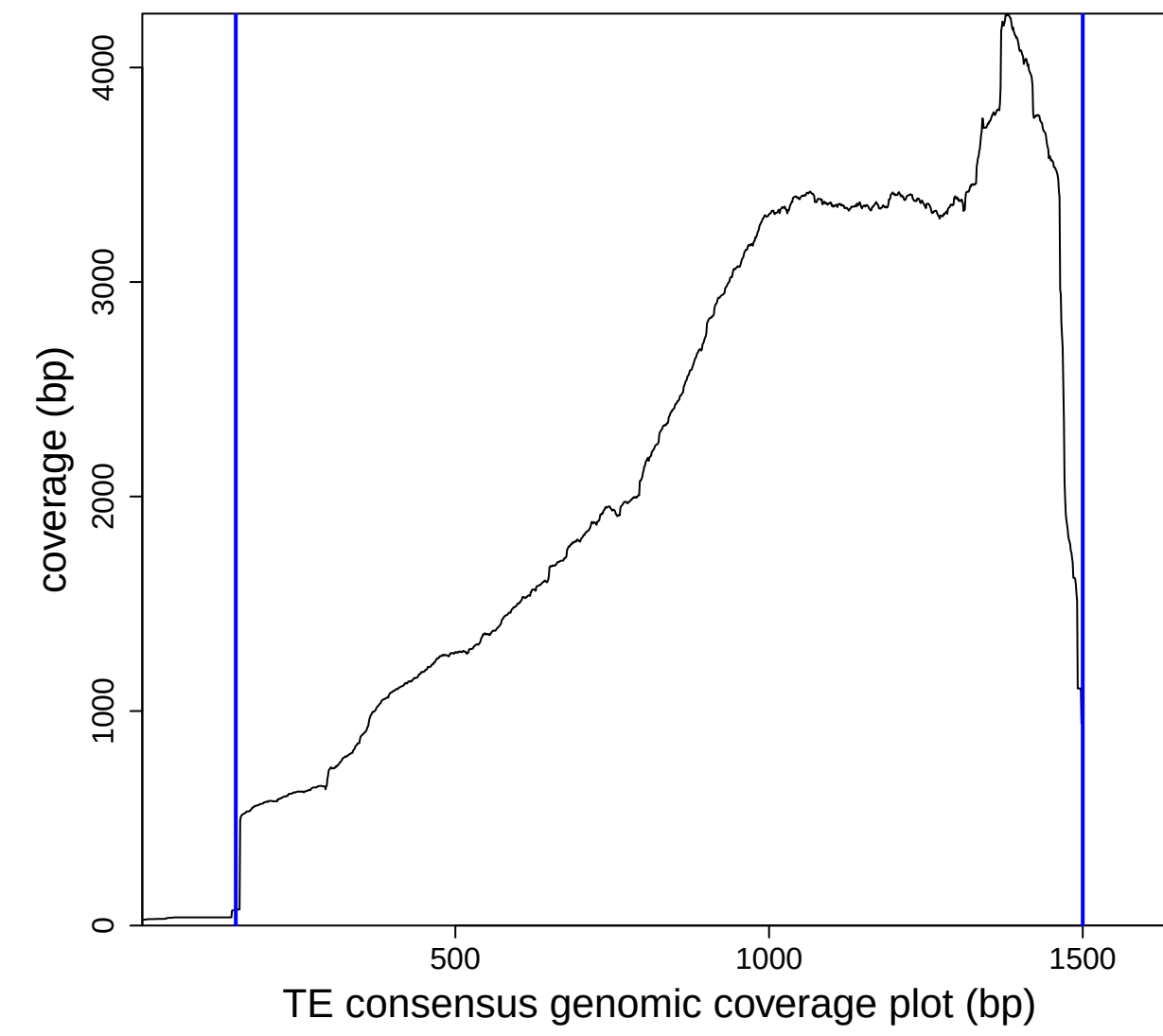
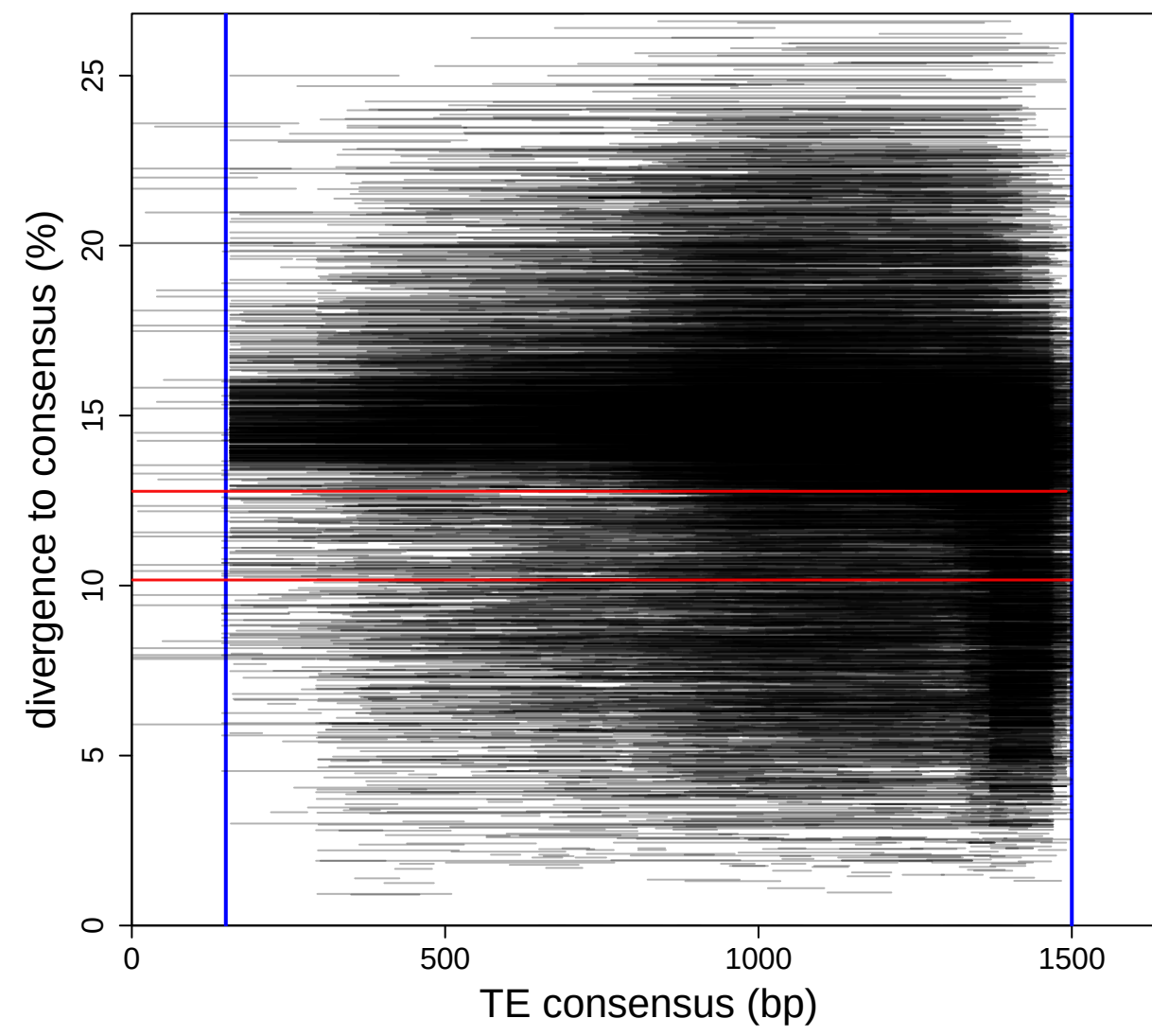
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 1346bp; fragments: 10894; full length: 372 (>=1211.4bp)

After TEtrimmer 1346 bp

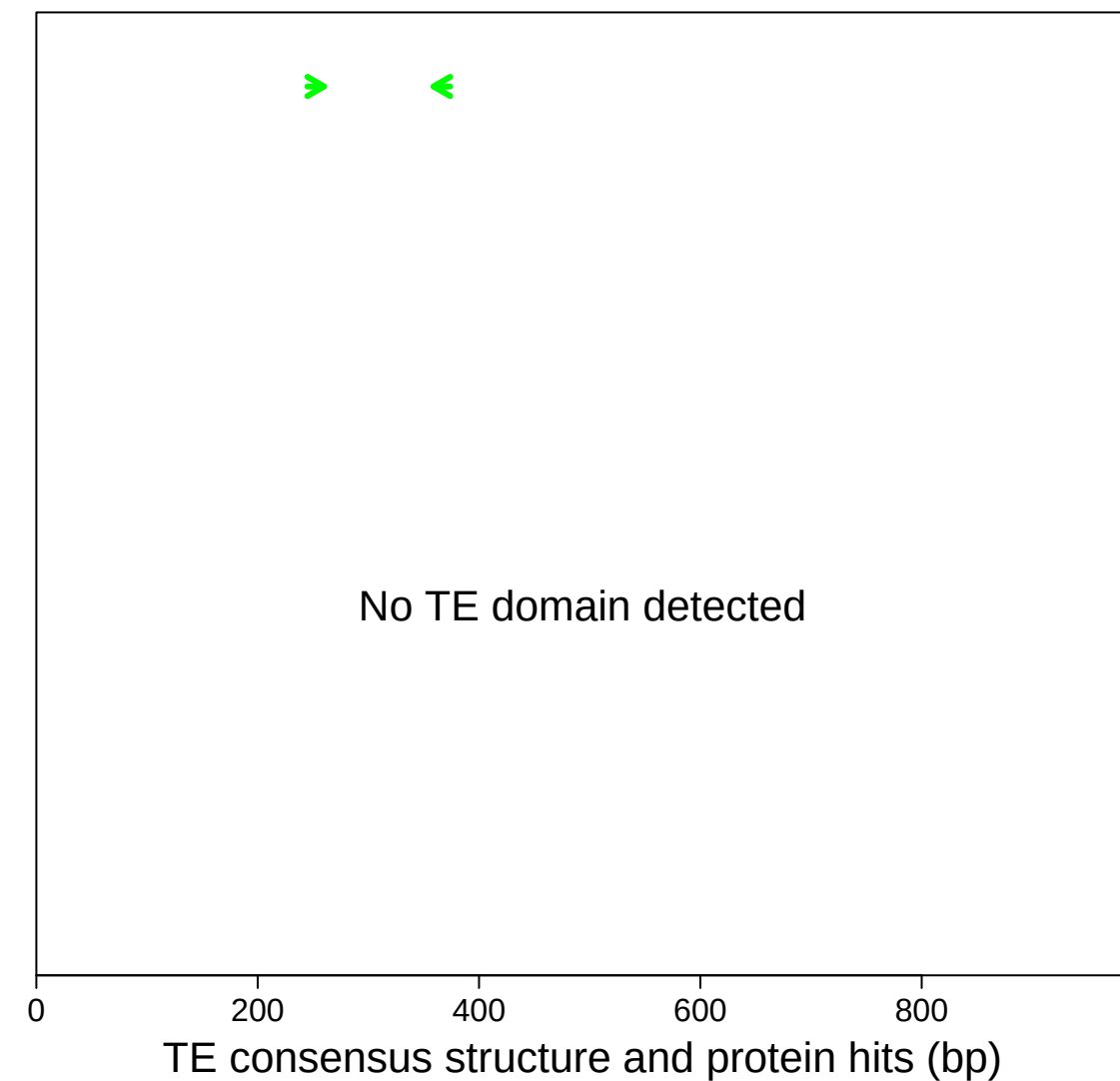
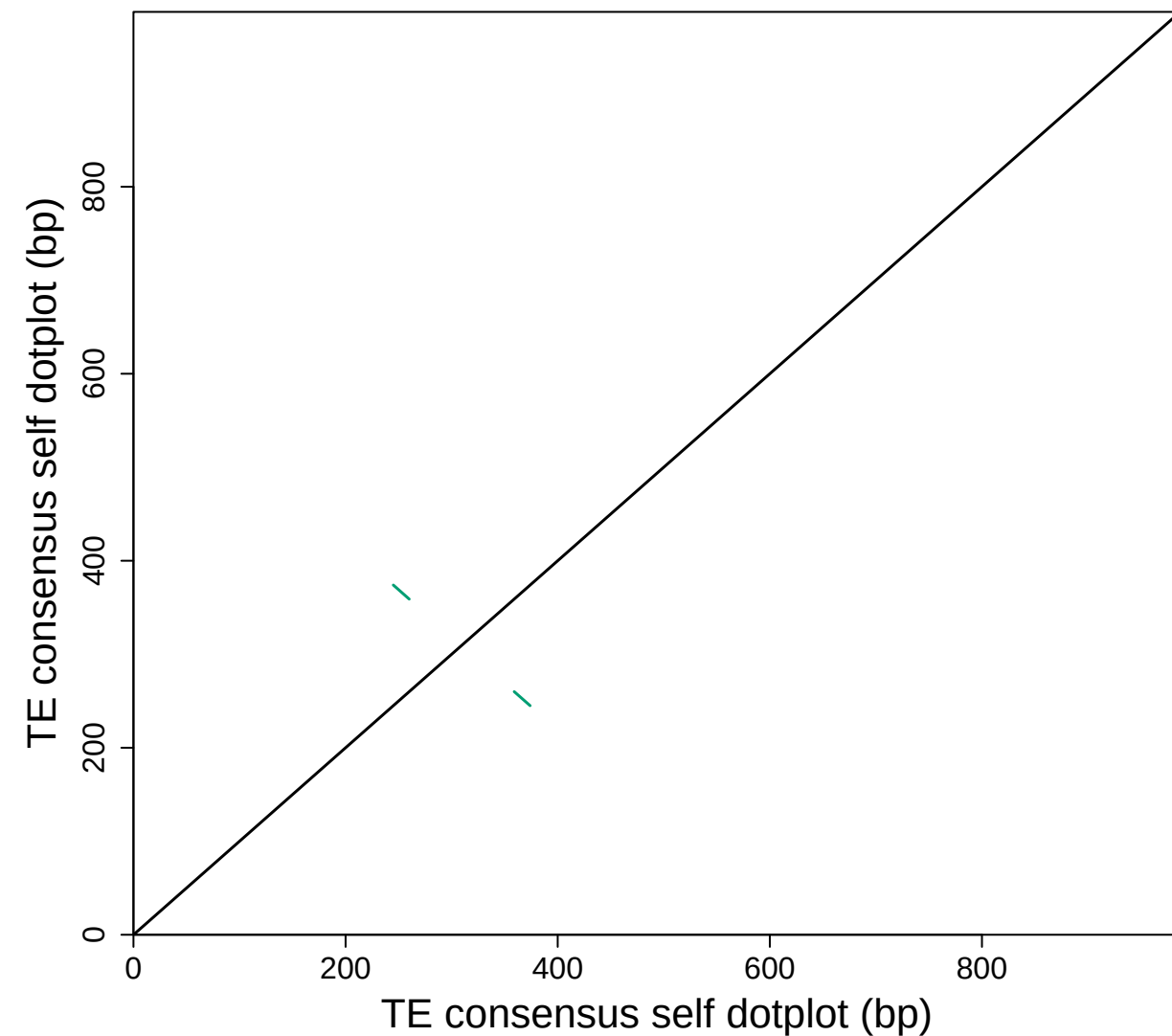
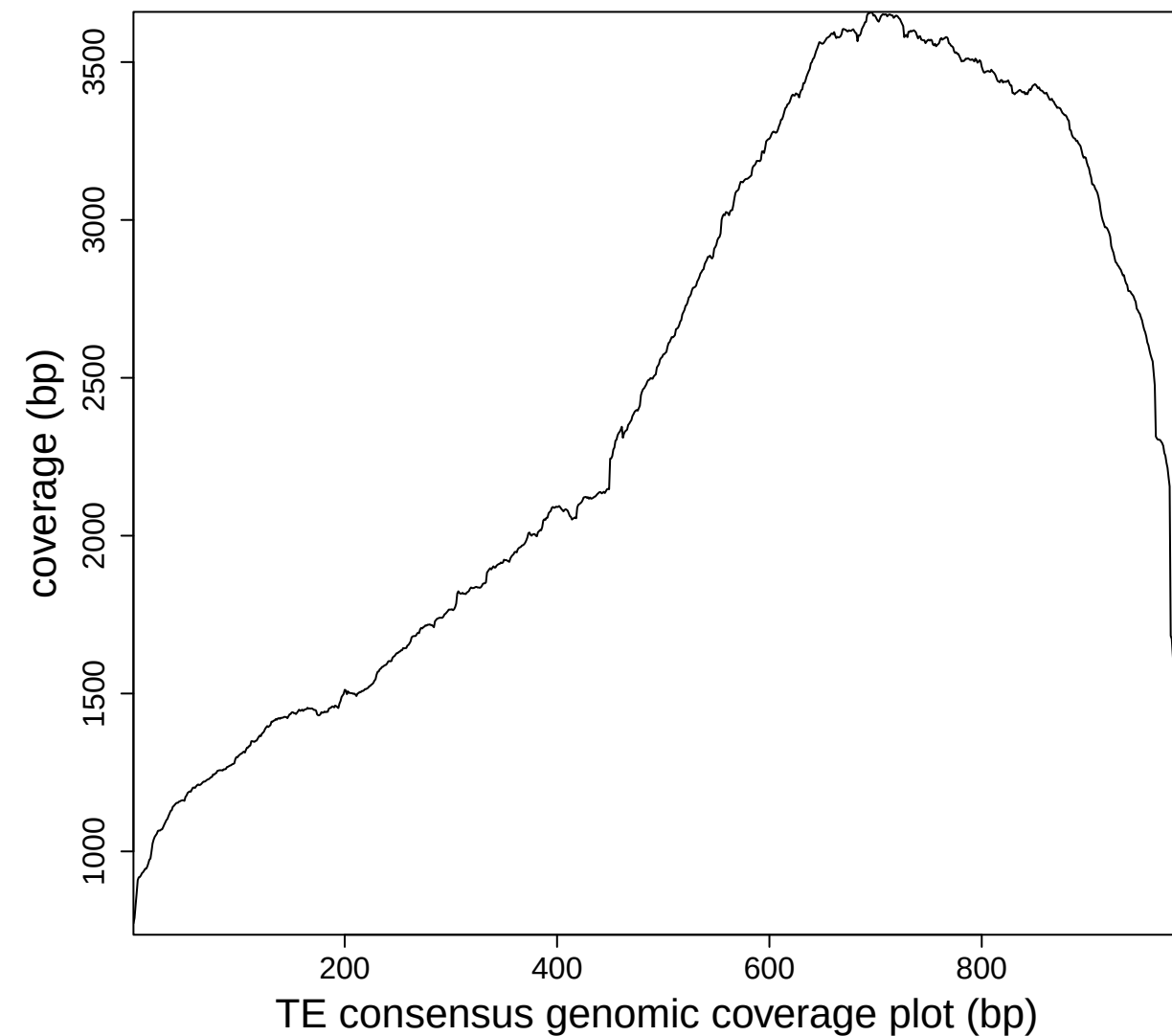
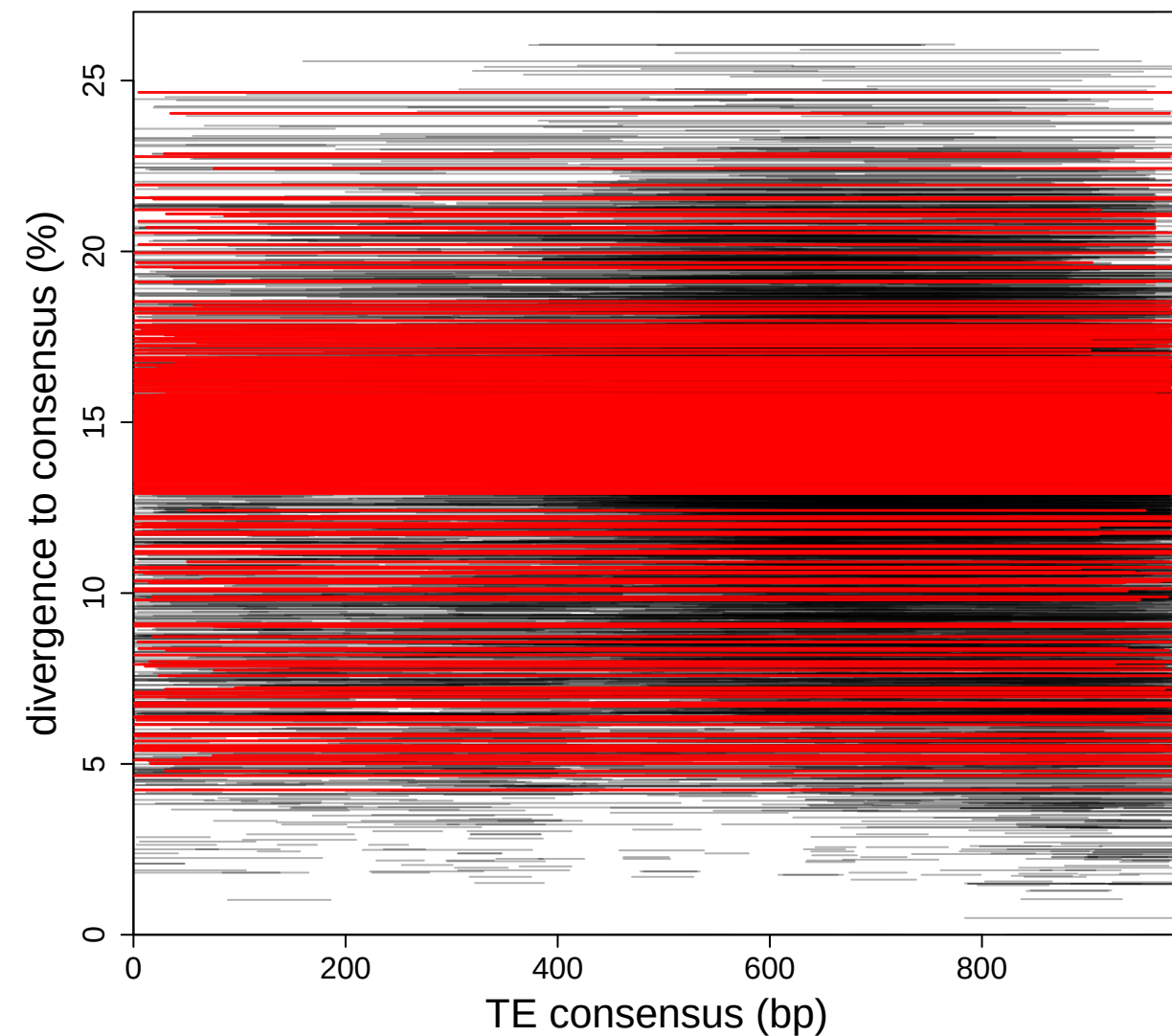


7.fasta.b.bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 1650bp; fragments: 7833; full length: 2 (>=1485bp)

After TEtrimmer Extended plot Blue lines are boundaries

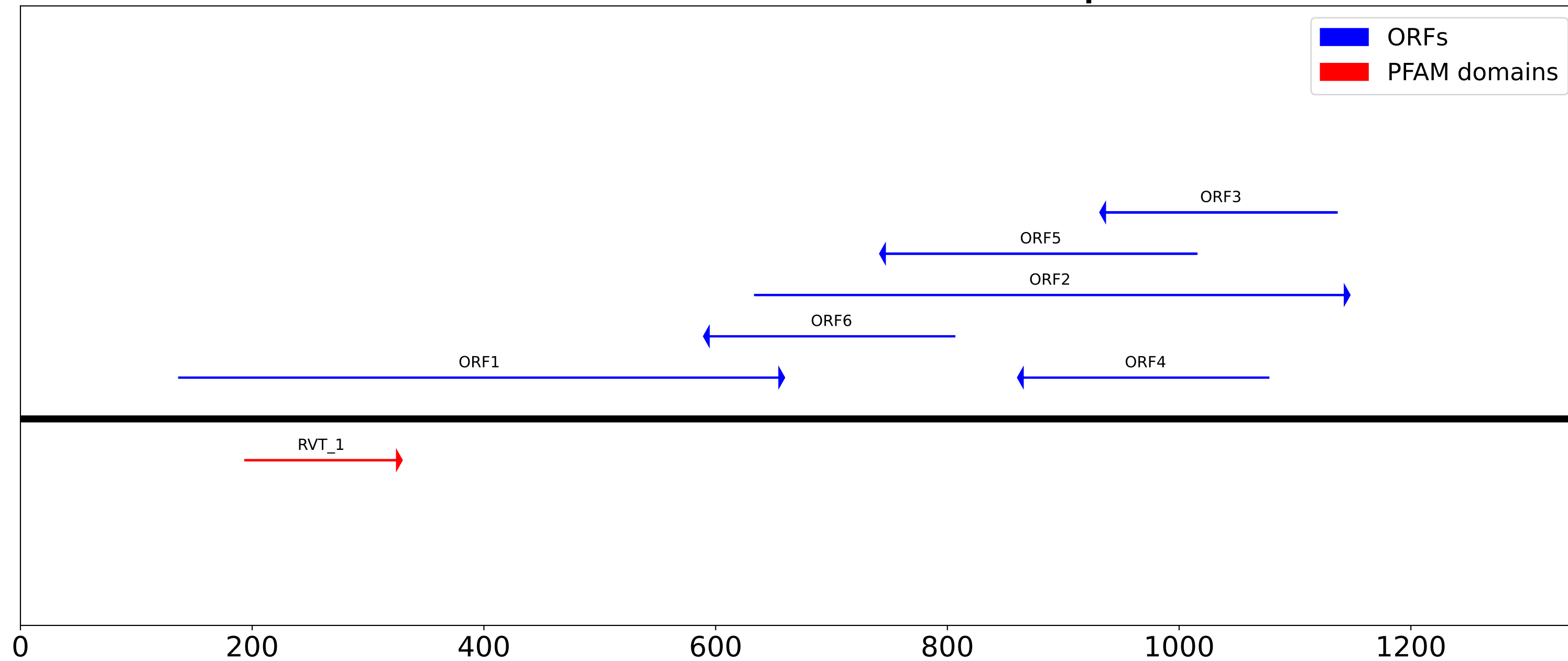


TE: rnd_1_family_317
size: 987bp; fragments: 5990; full length: 595 (≥ 888.3 bp)

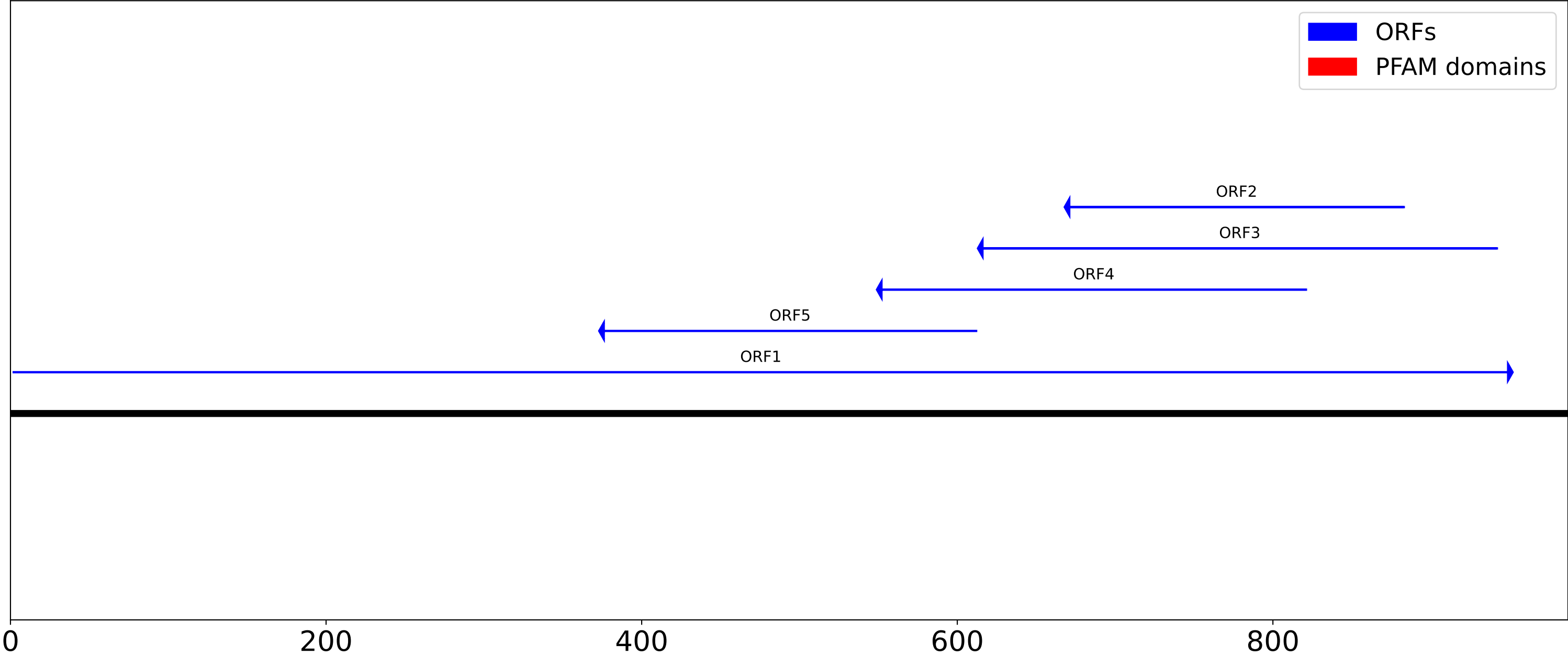


Before TEtrimmer 987 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

