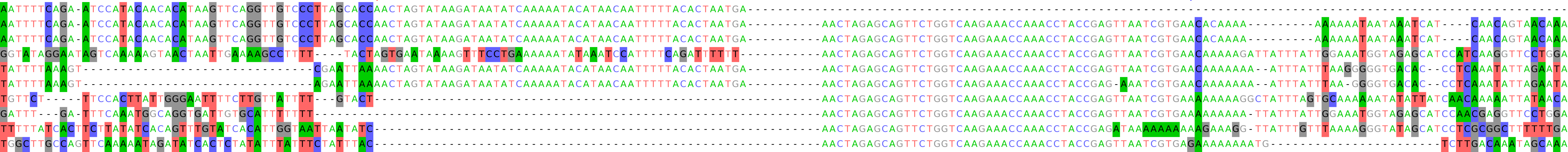


Start Crop Point

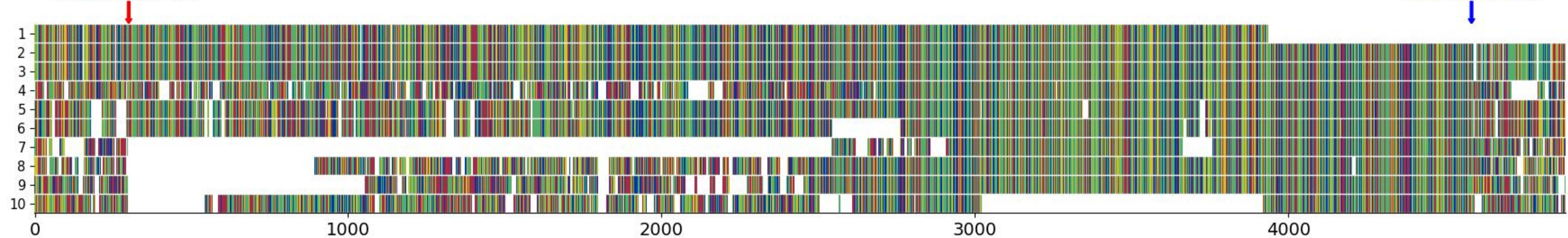
End Crop Point

MSA length = 4285



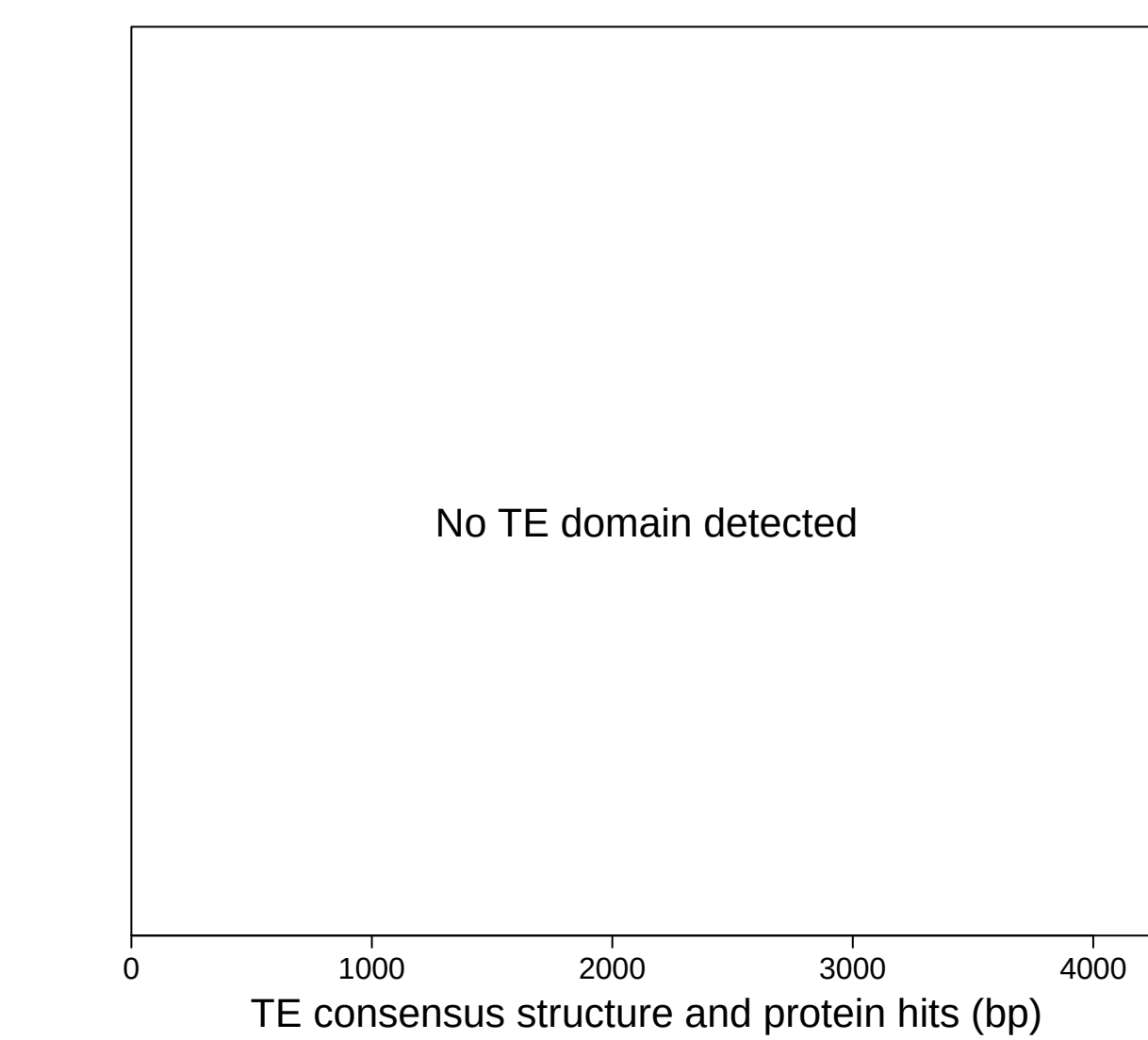
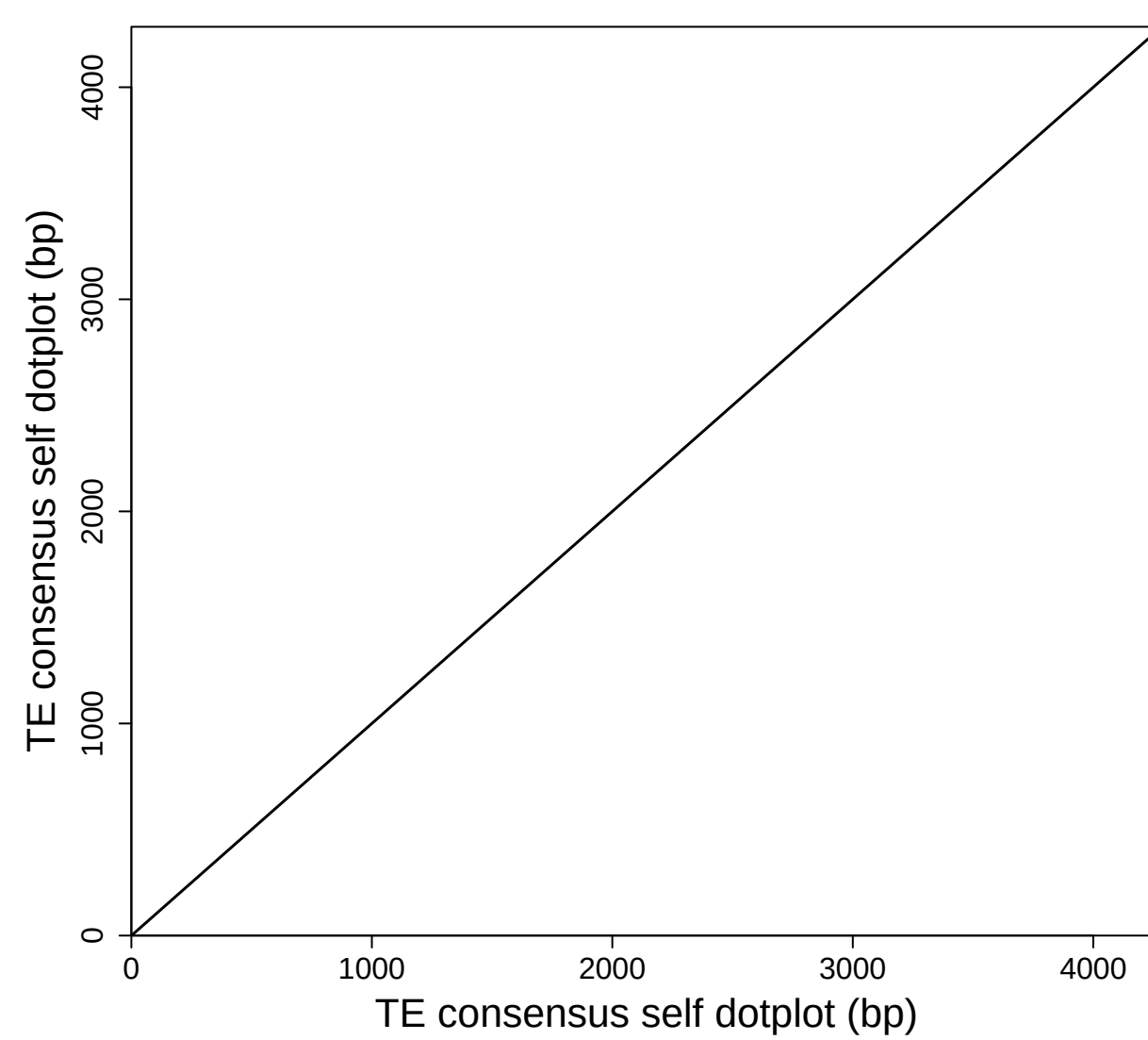
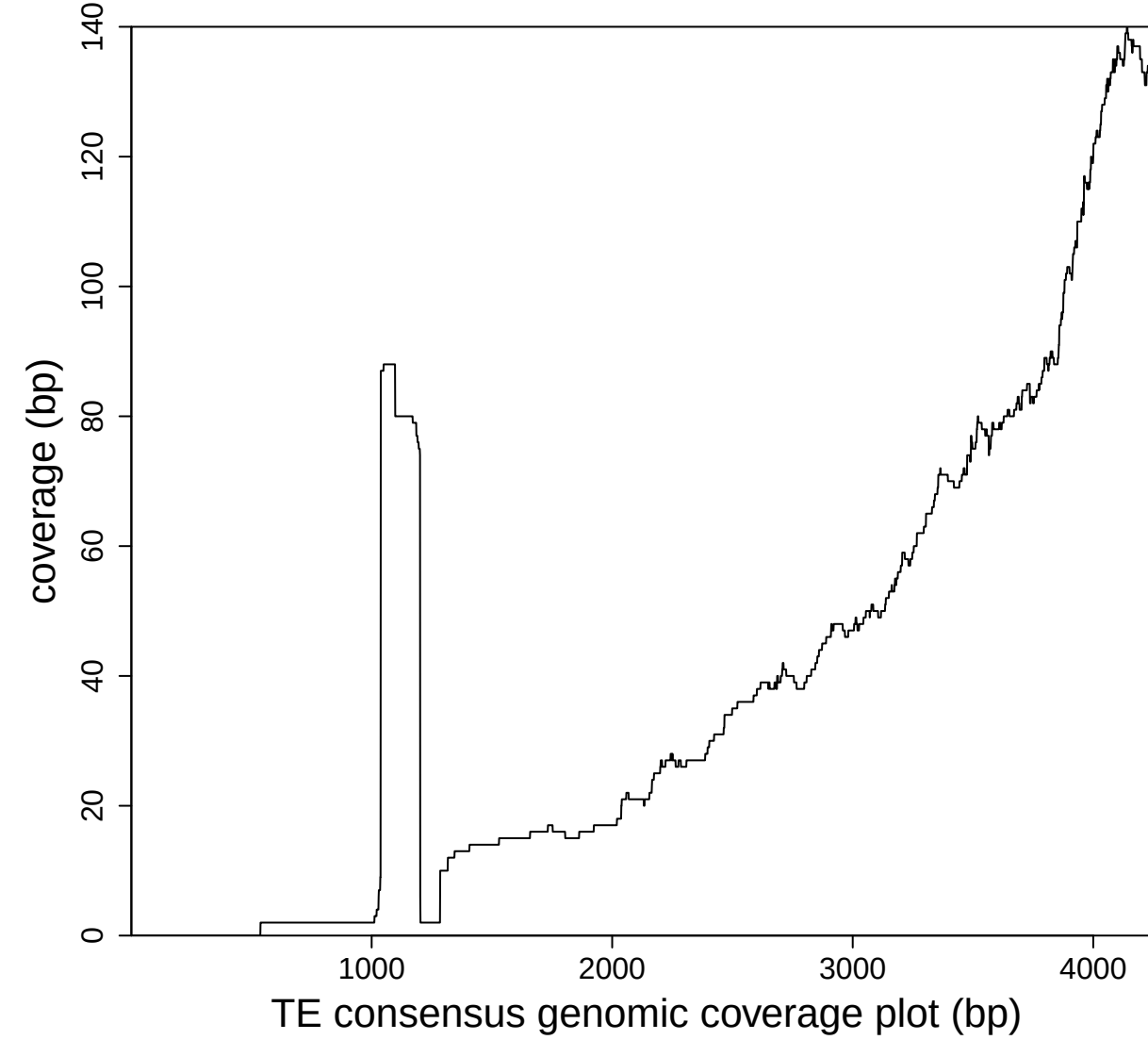
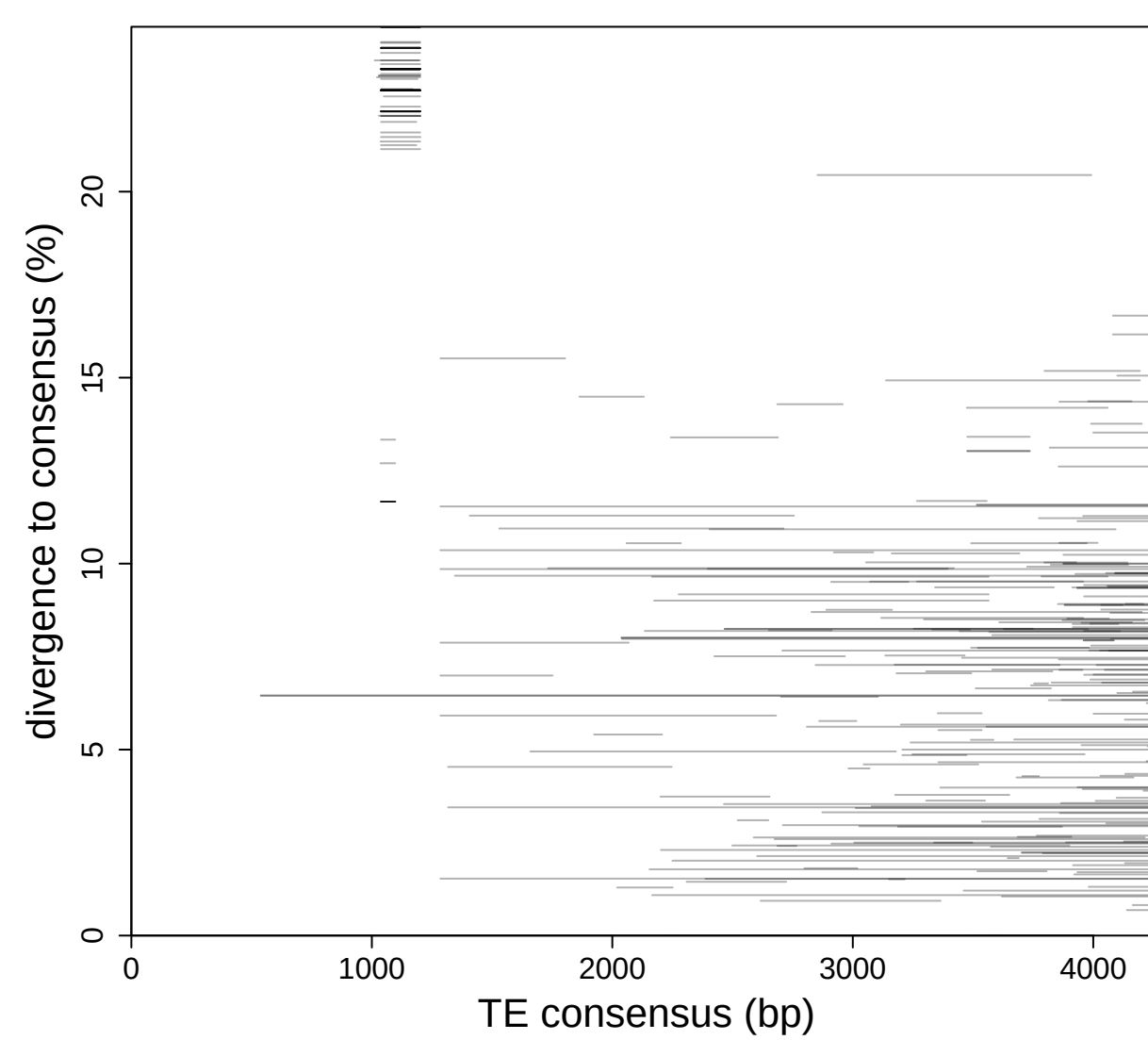
Start crop Point

End crop Point



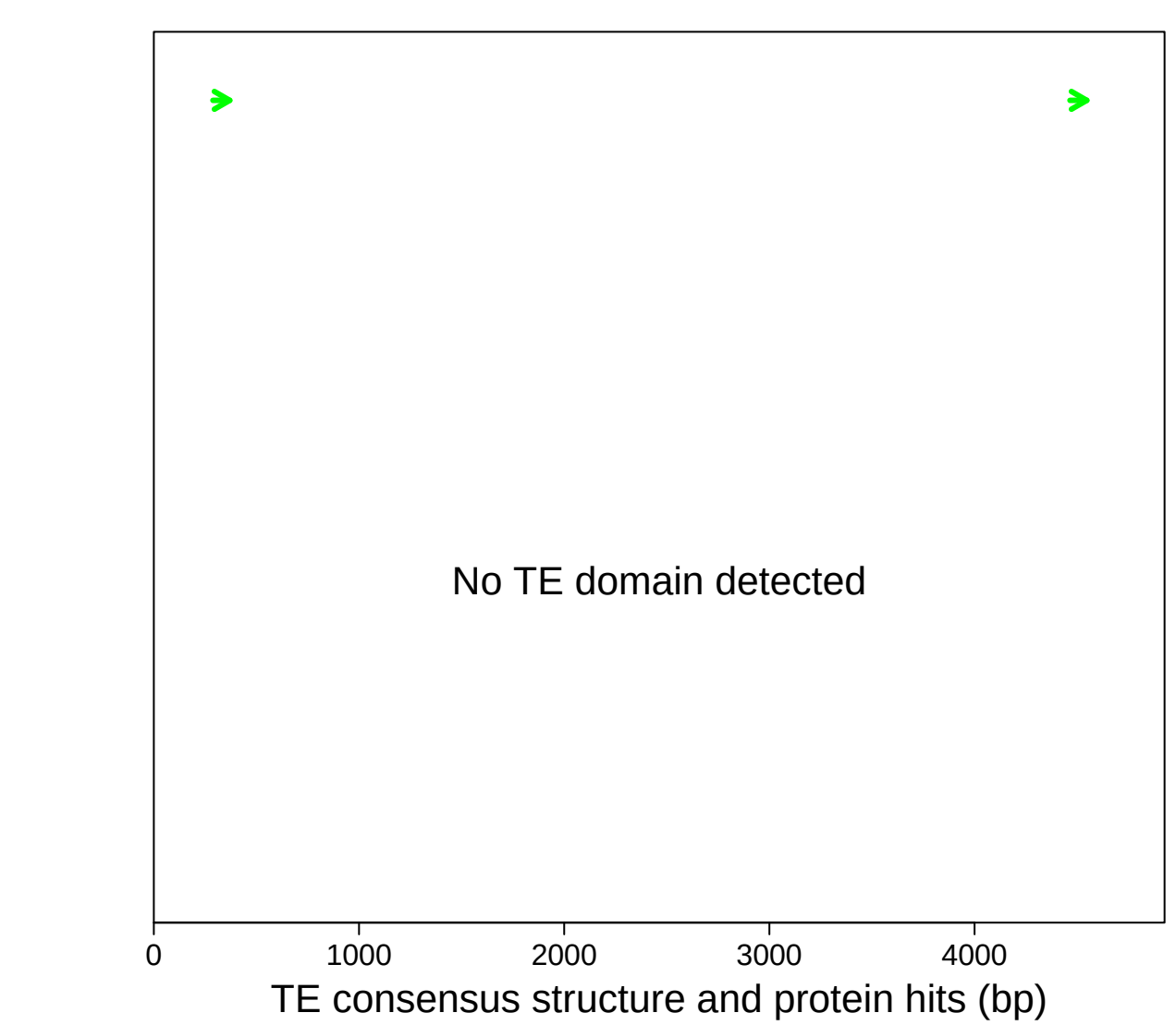
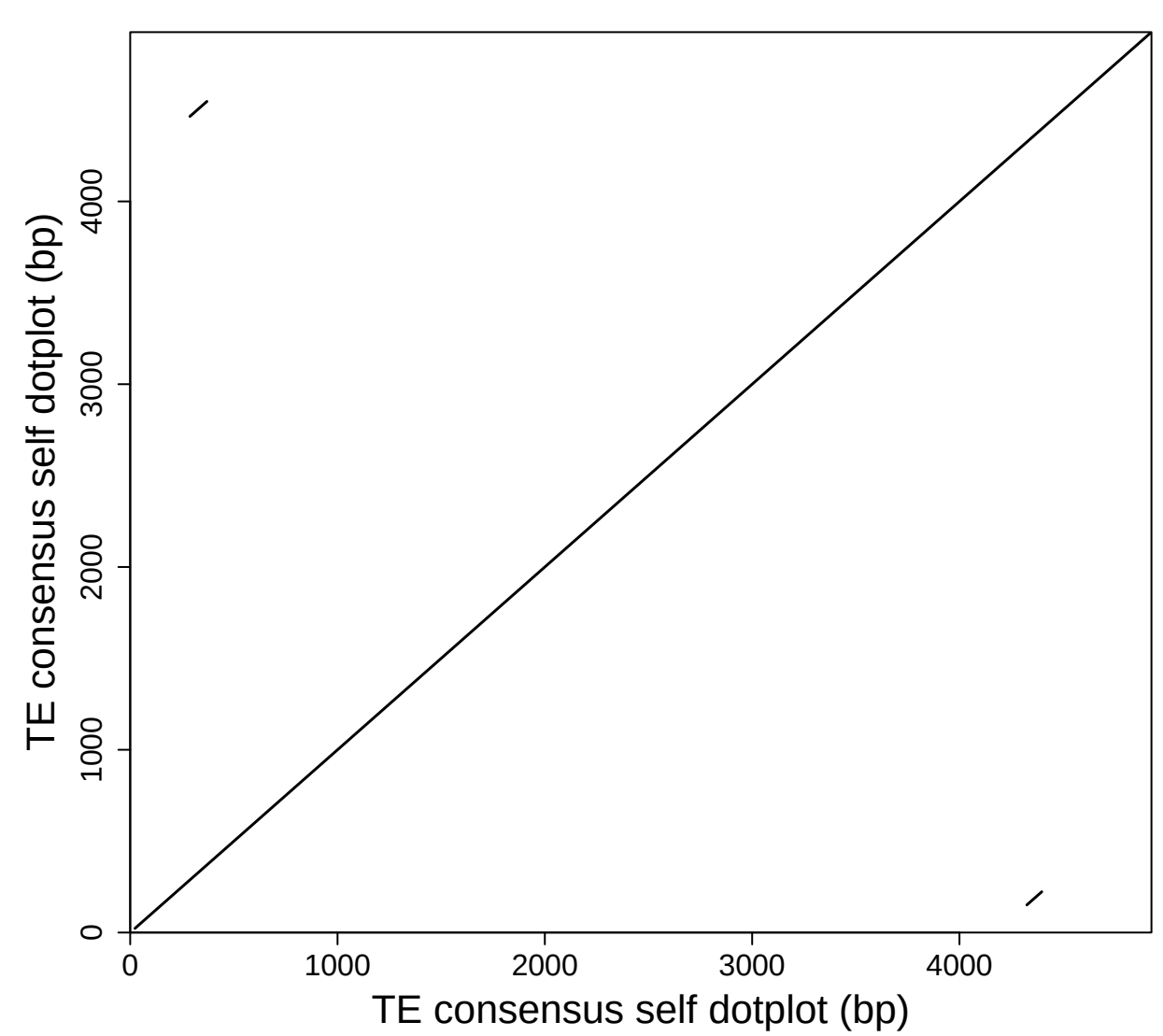
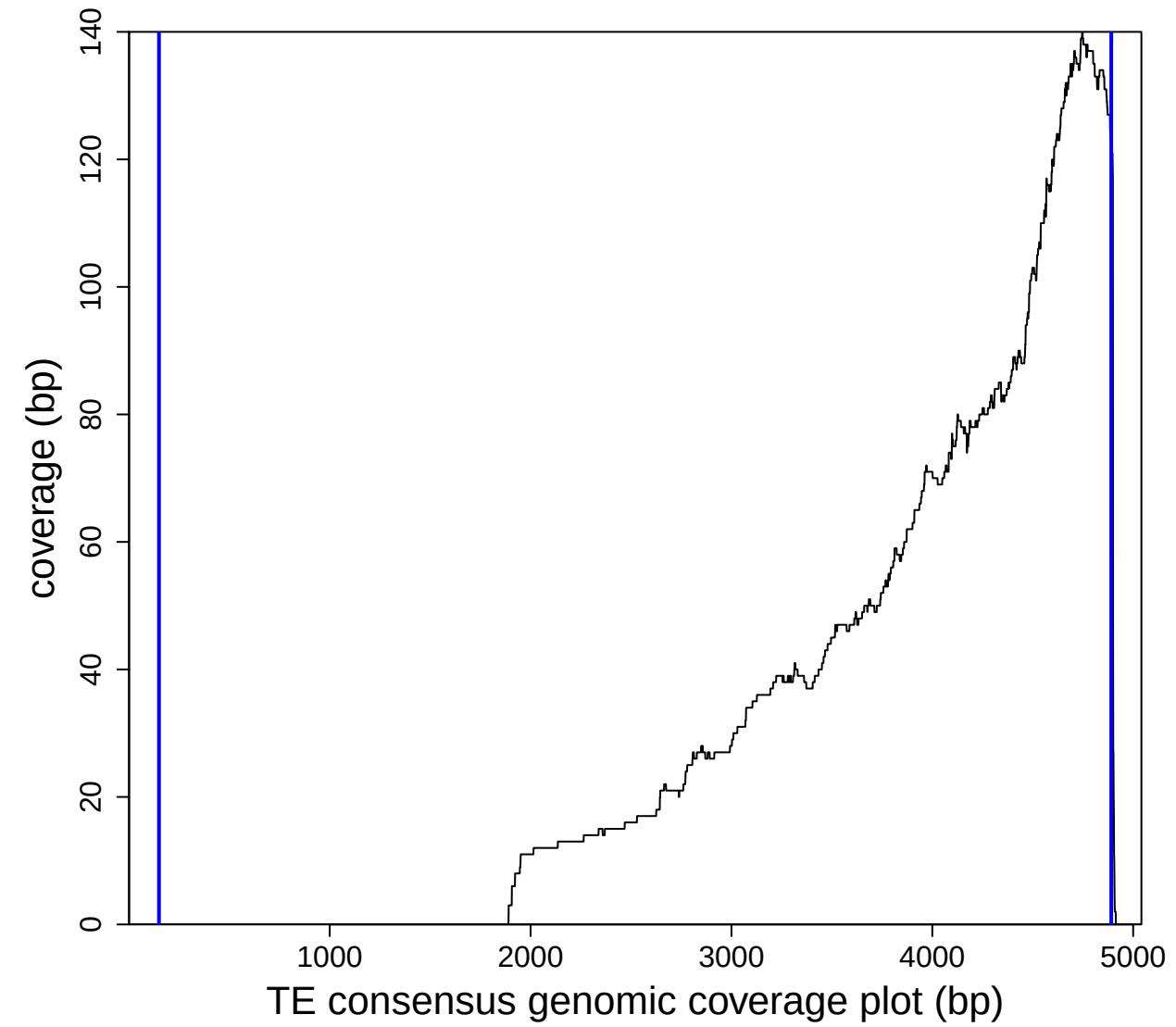
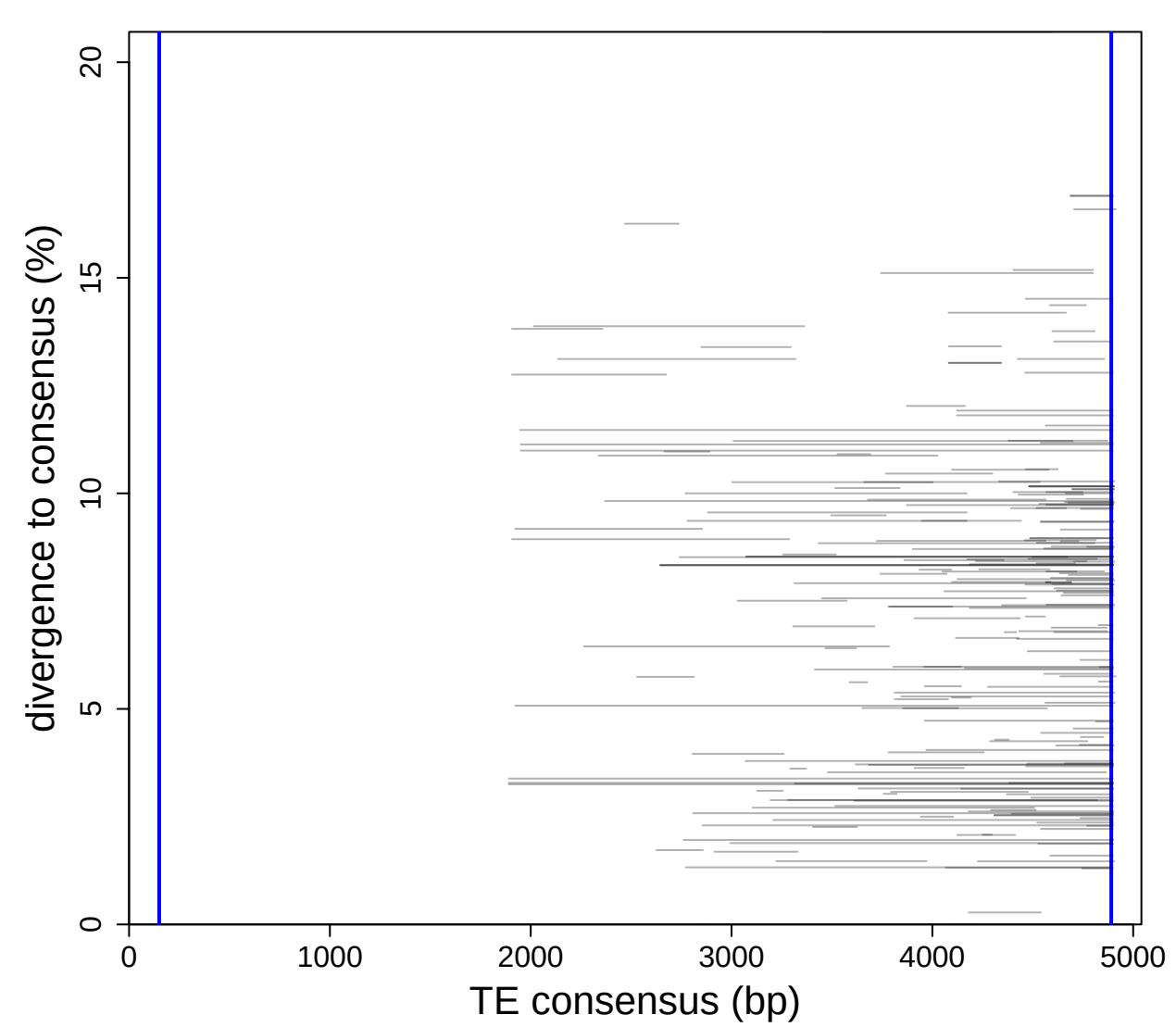
1.bed_fm 1.bed 0 0_n.bed_g 1.bed_fm 2.bed 0 0_bcln.fa_aln.f
size: 4285bp; fragments: 316; full length: 0 (>=3856.5bp)

After TEtrimmer 4285 bp

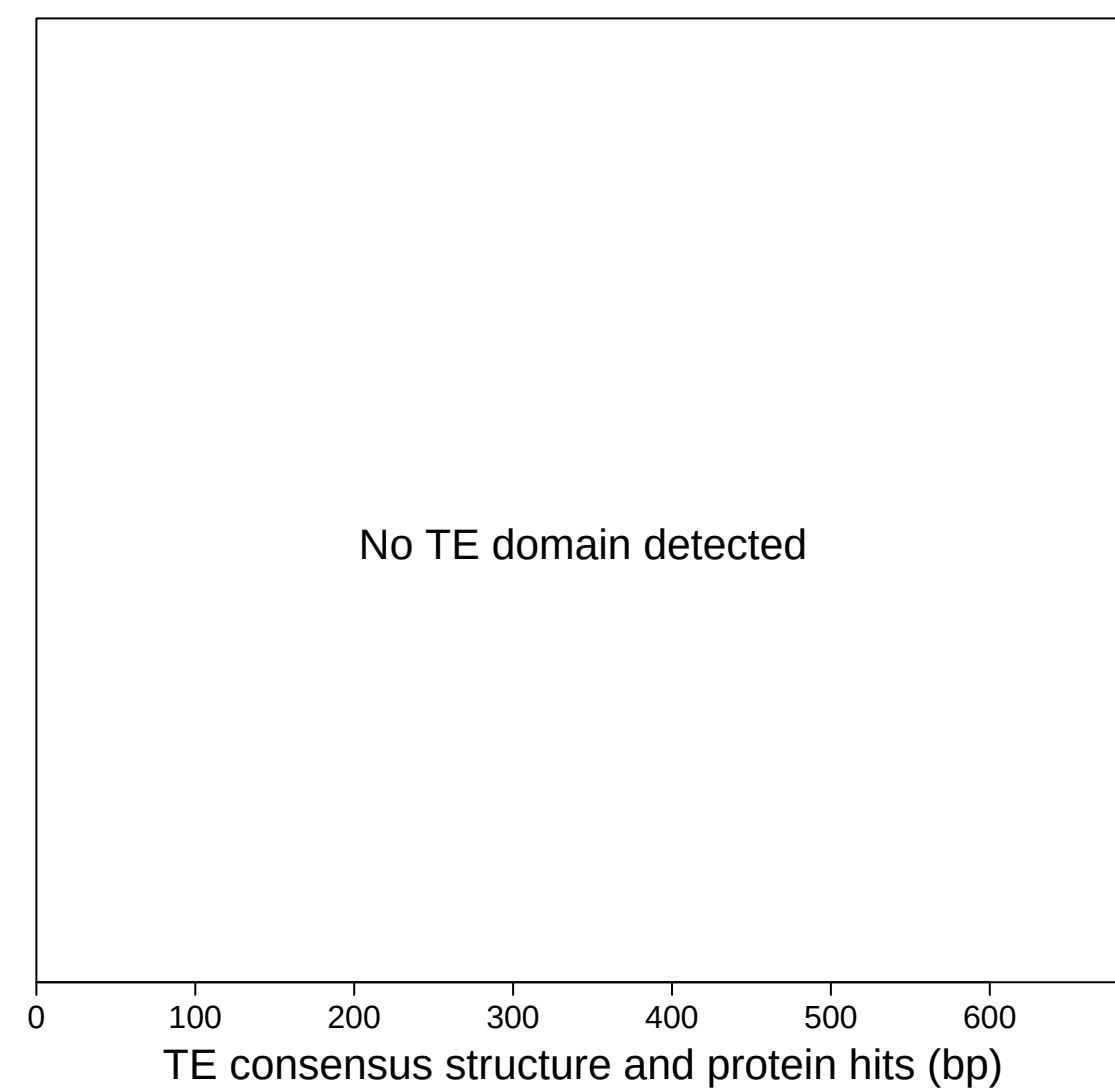
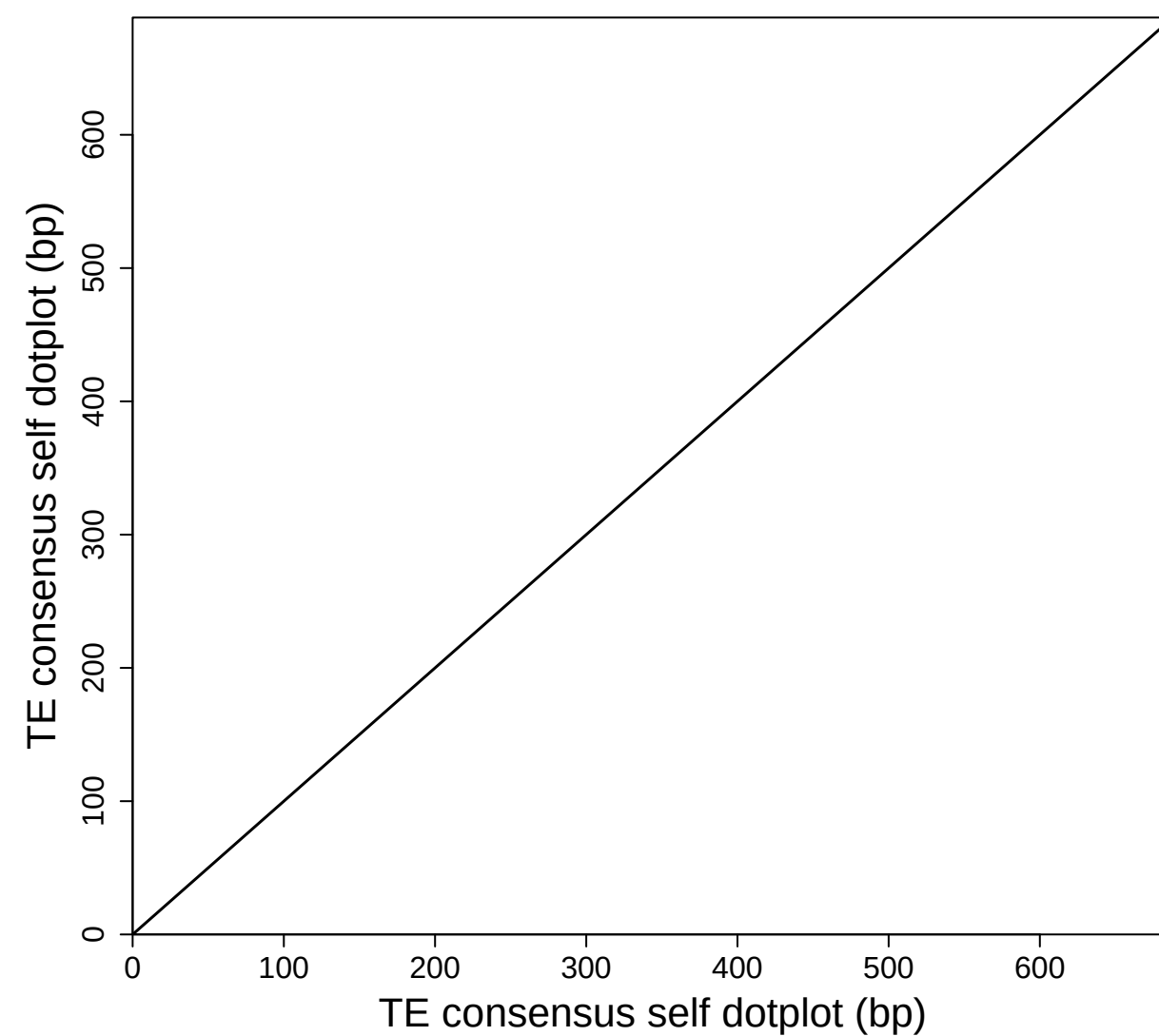
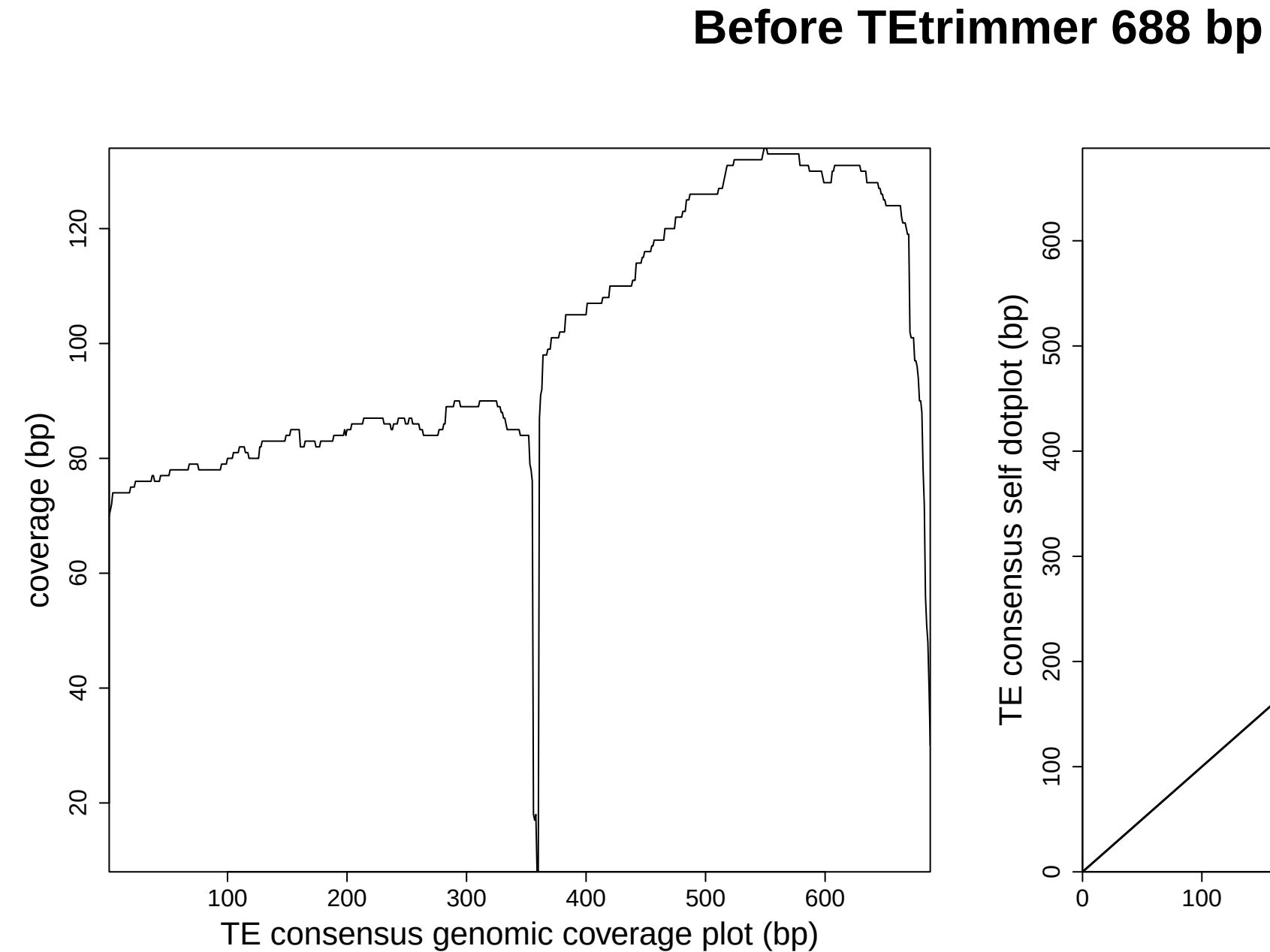
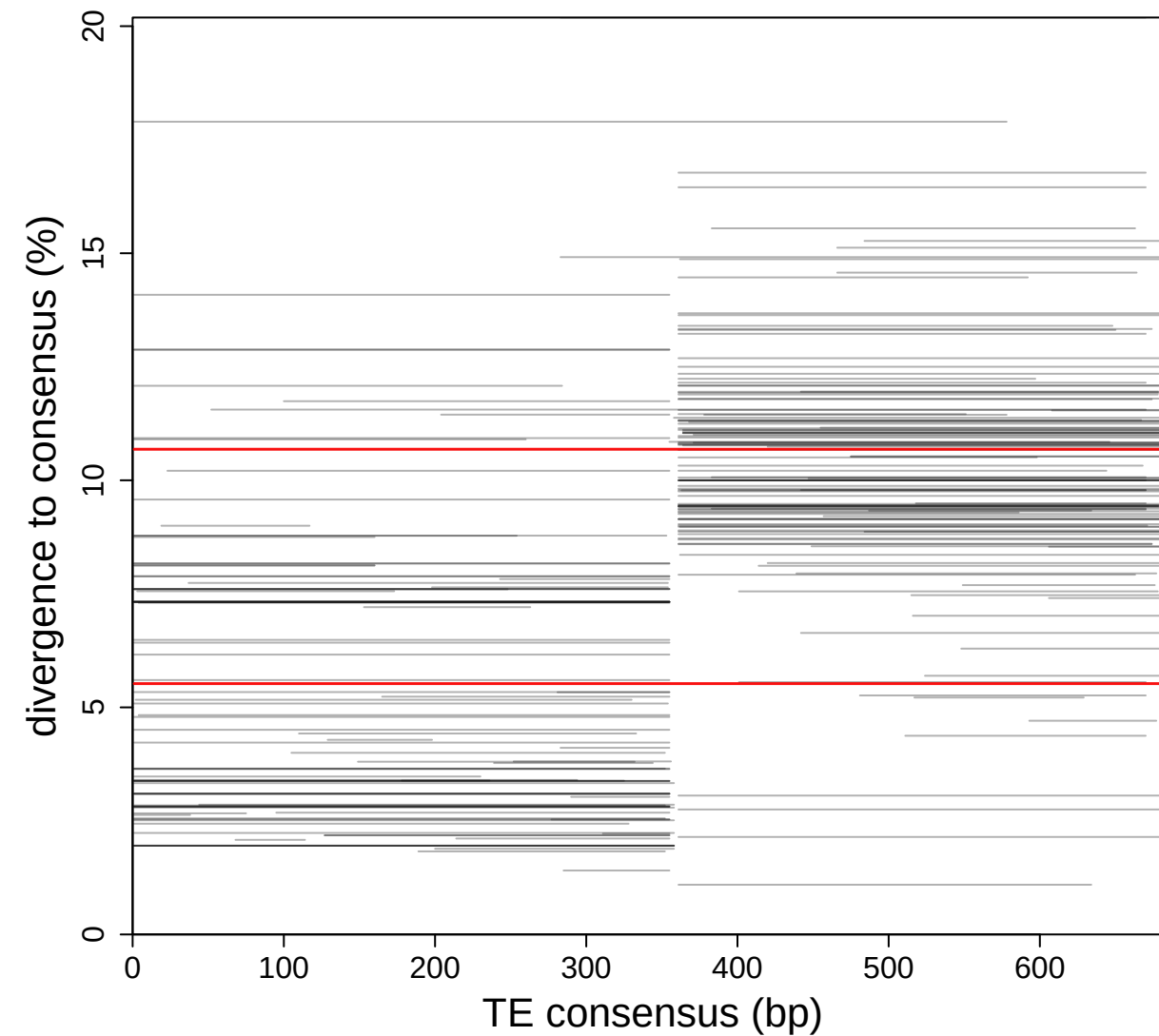


bed_uf.bed_g_4.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0
size: 5041bp; fragments: 228; full length: 0 (>=4536.9bp)

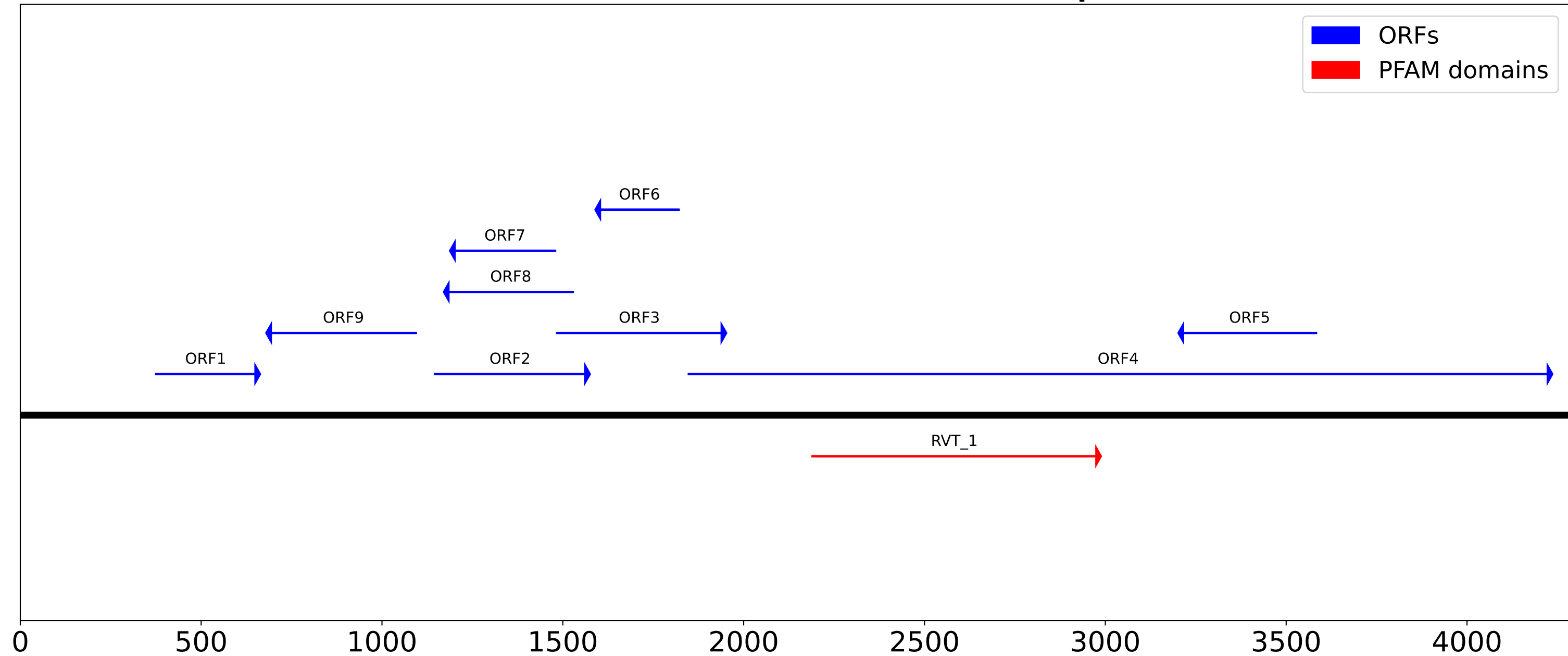
After TEtrimmer Extended plot Blue lines are boundaries



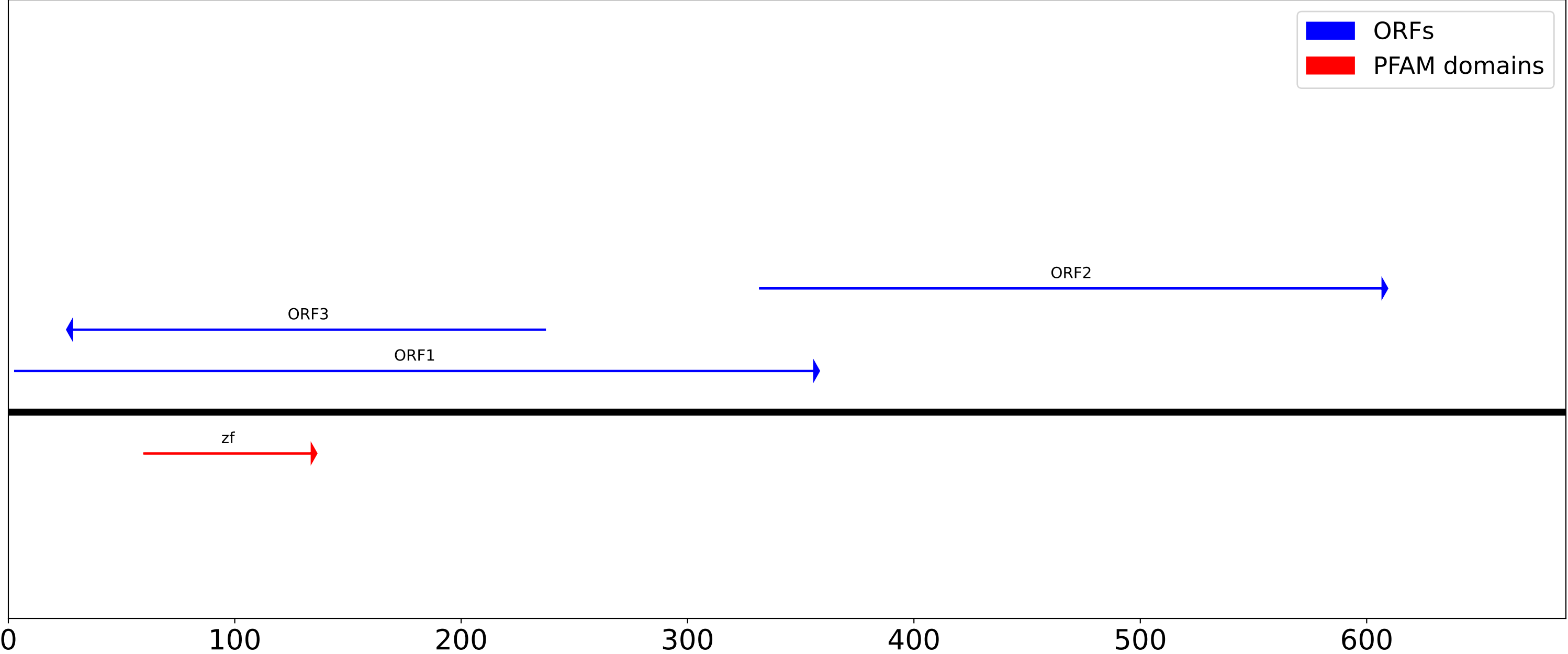
TE: rnd_5_family_7119
size: 688bp; fragments: 239; full length: 2 (≥ 619.2 bp)



After TEtrimmer ORF and PFAM domain plot



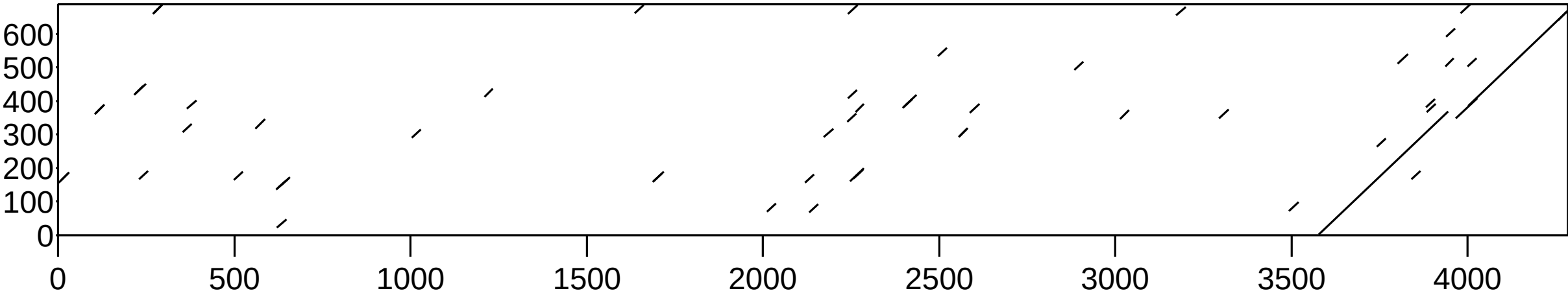
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)