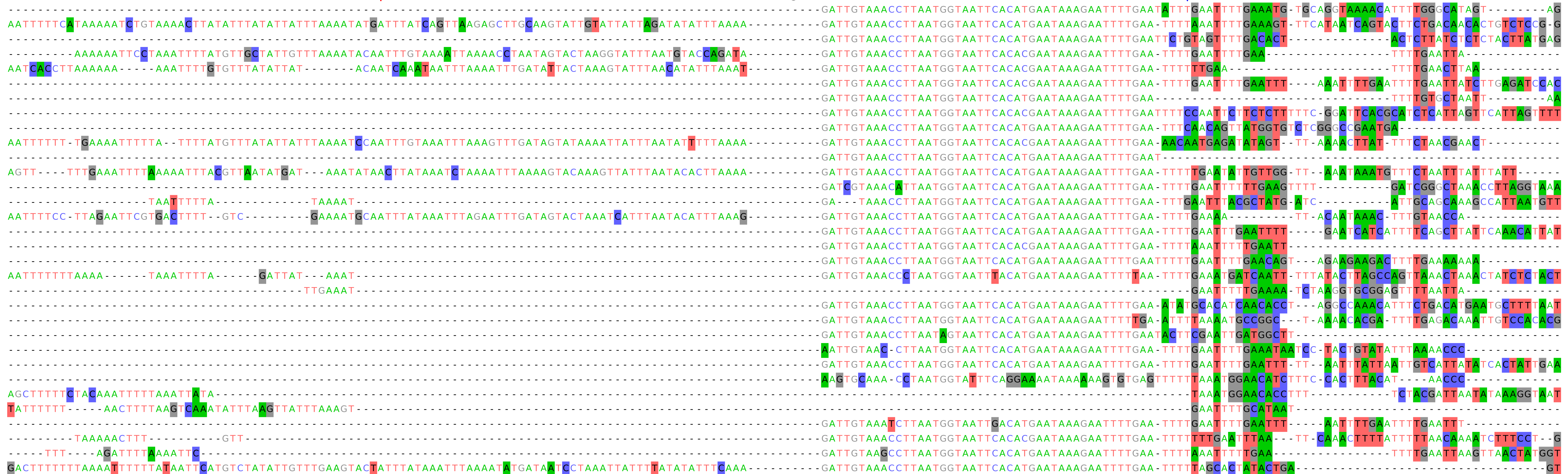
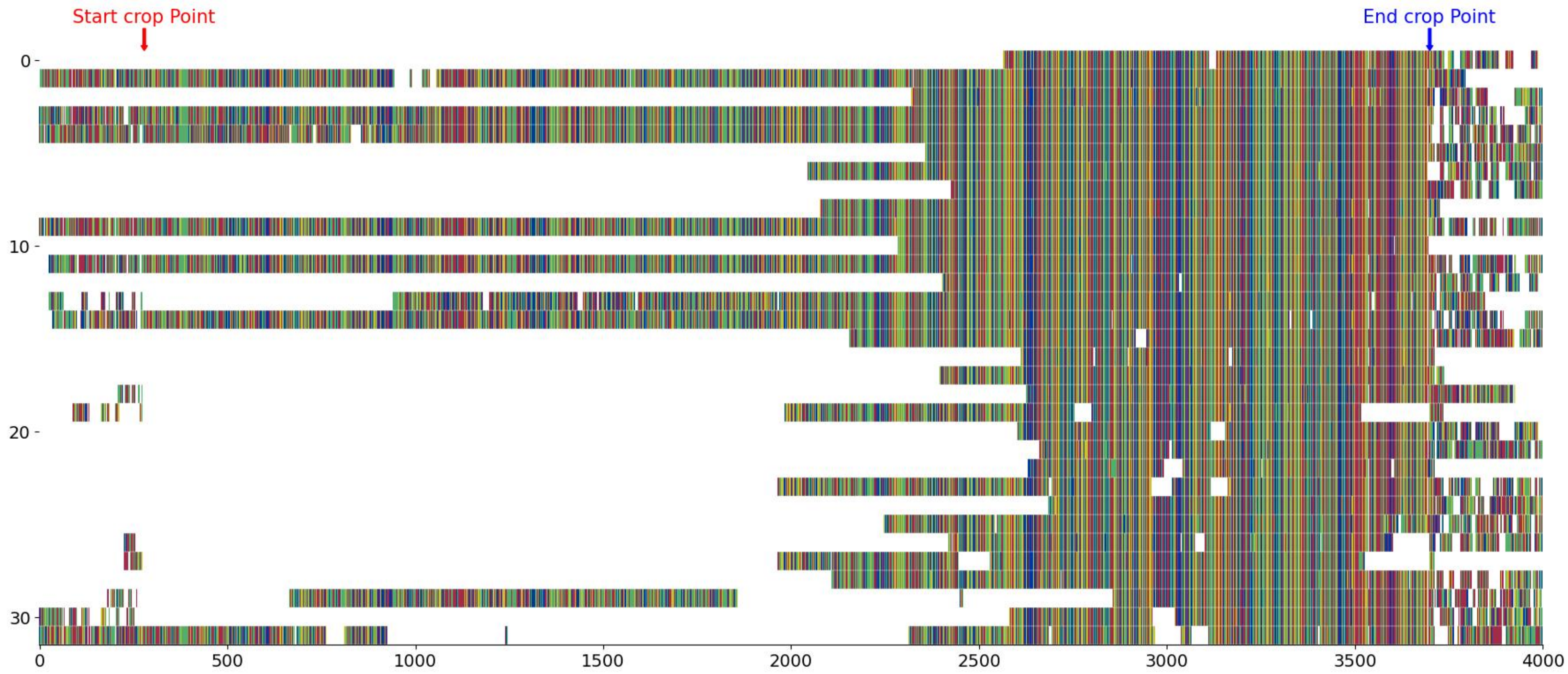


MSA length = 3421



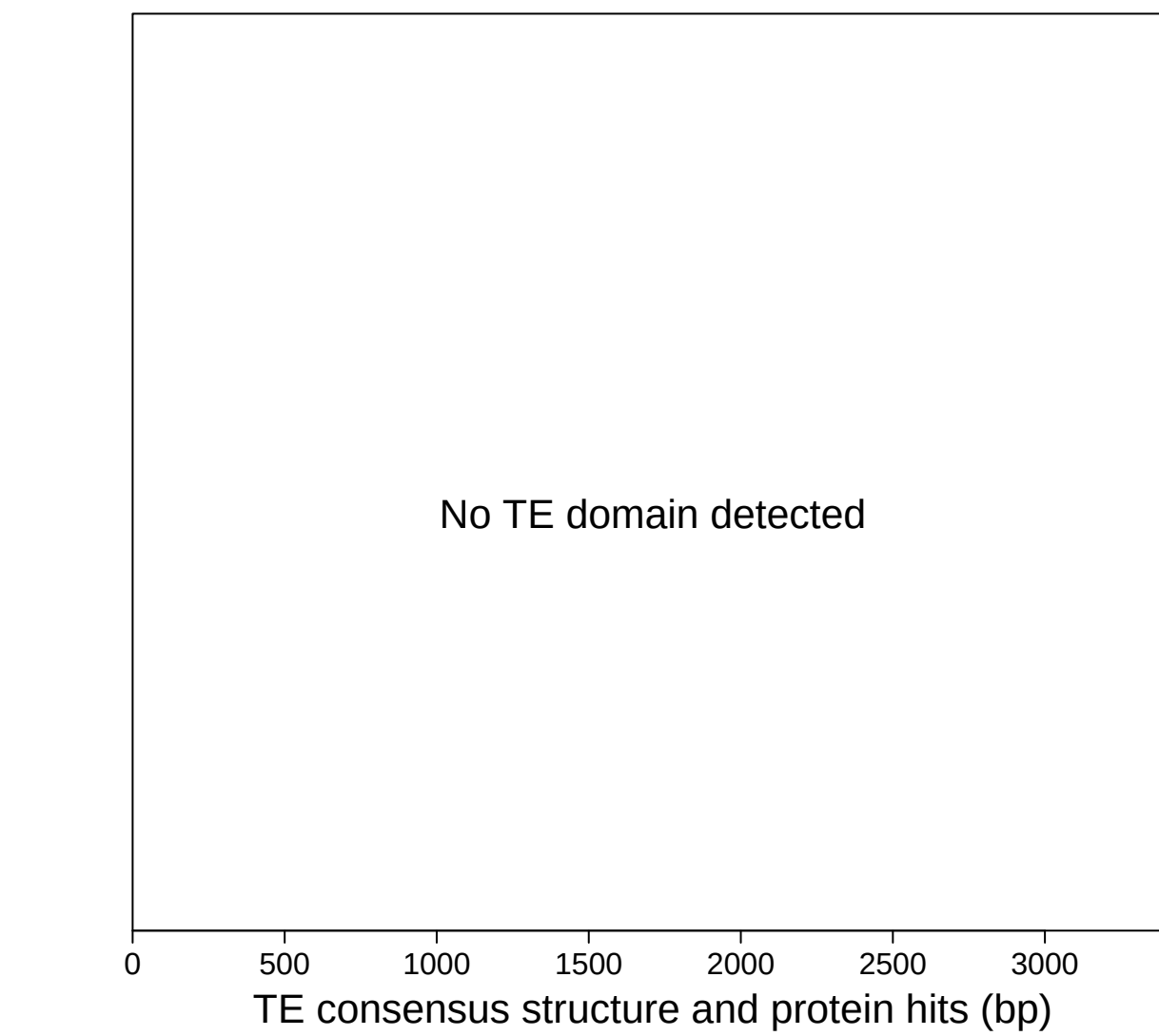
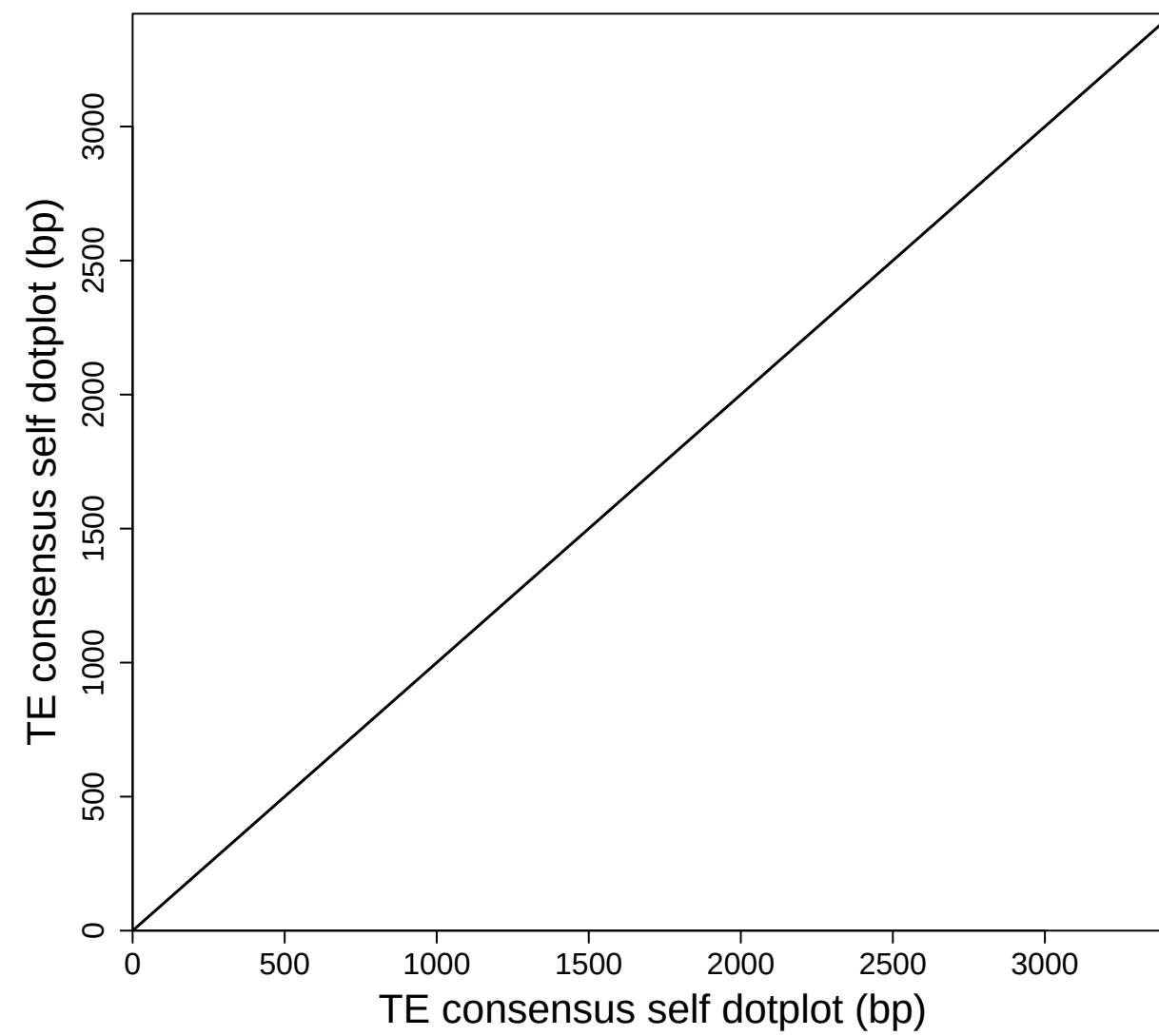
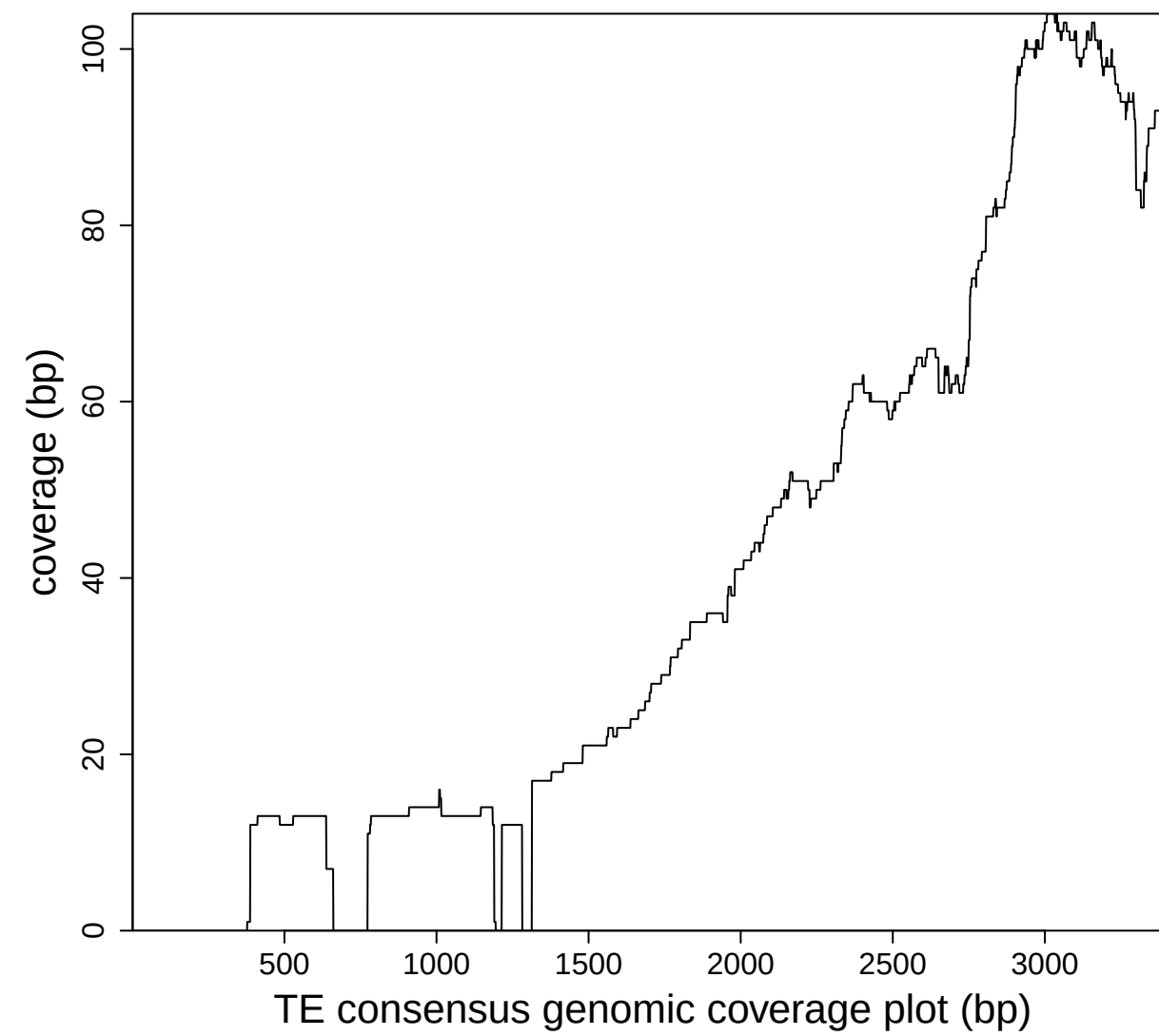
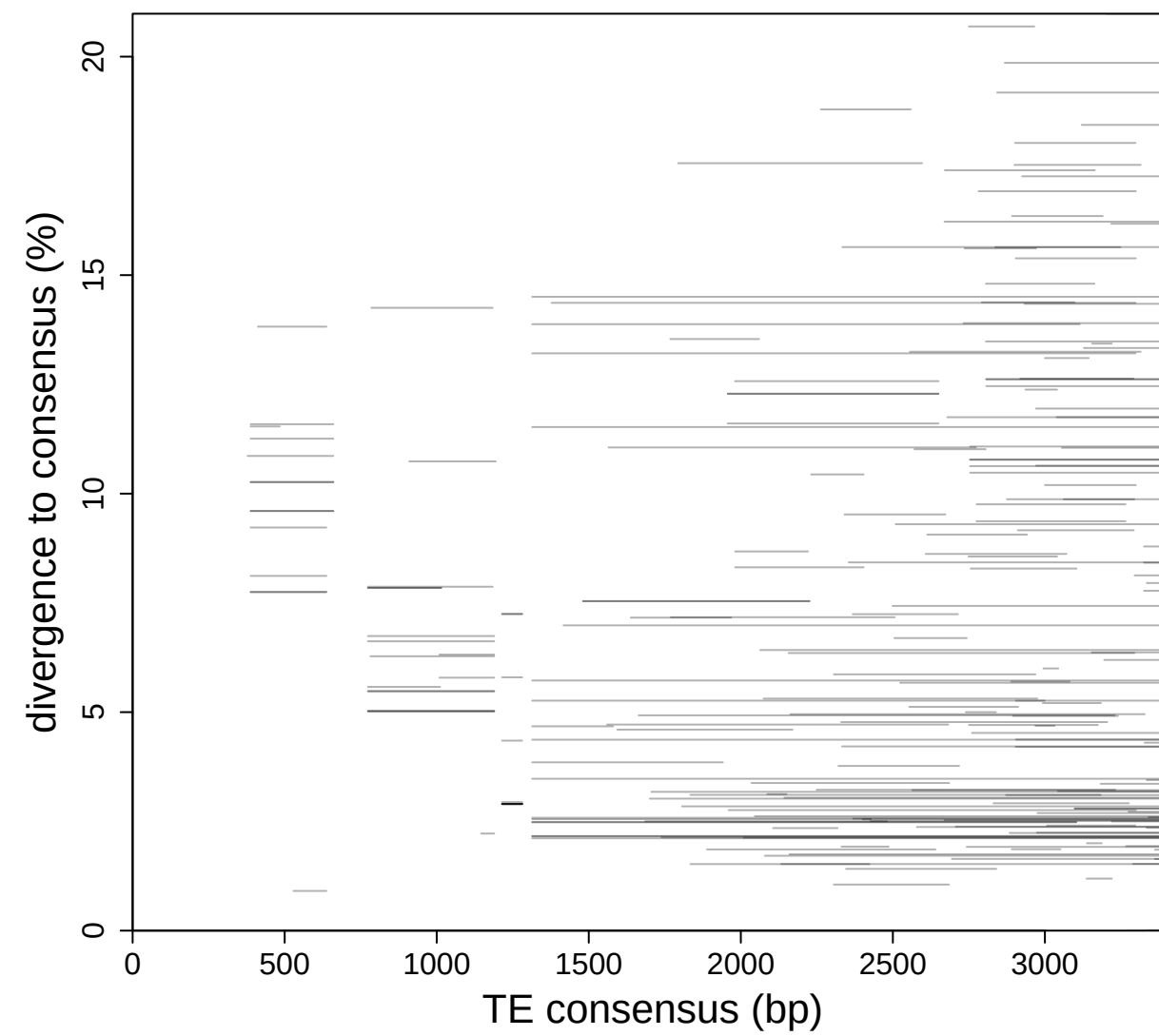
Start crop Point

End crop Point



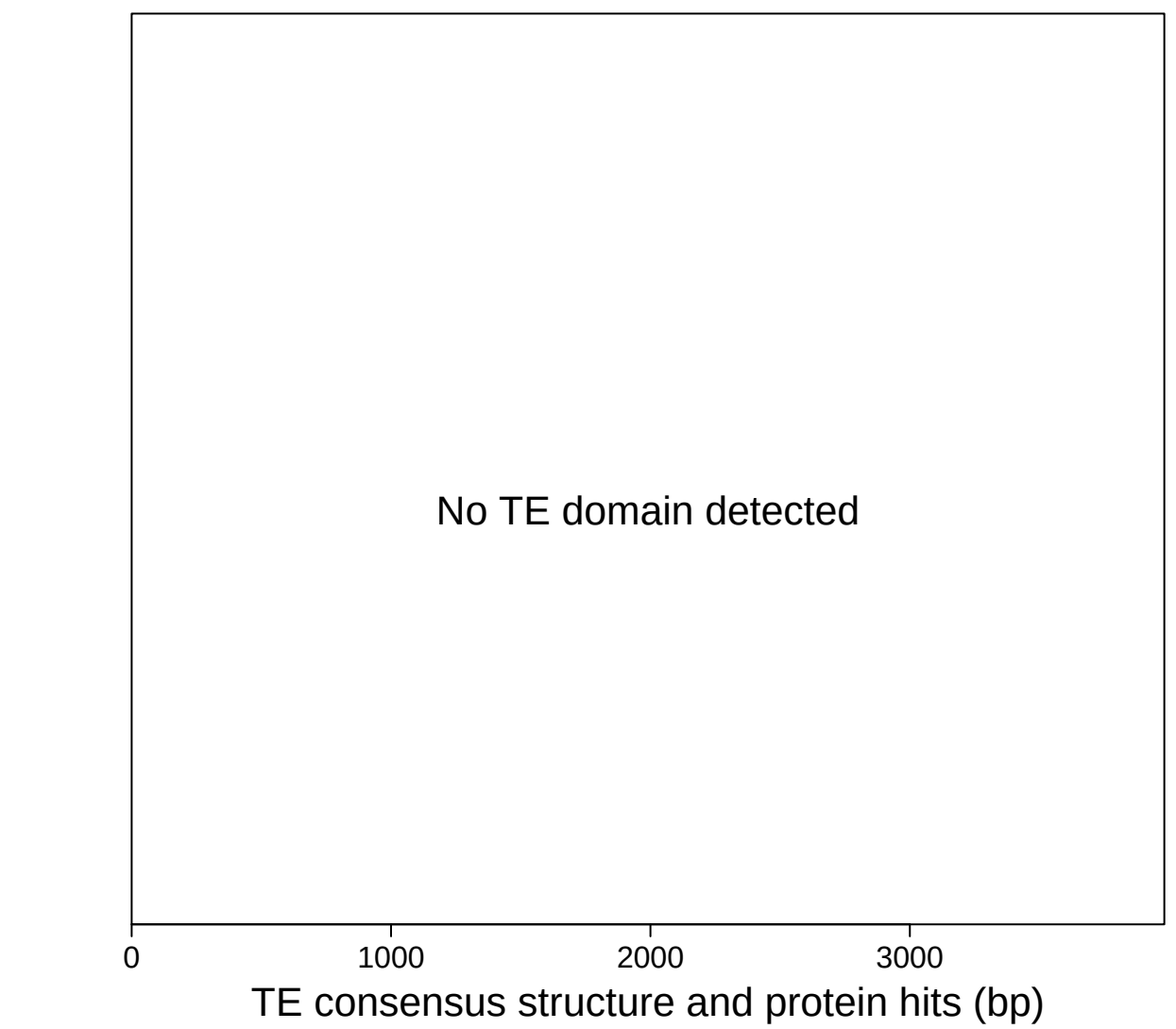
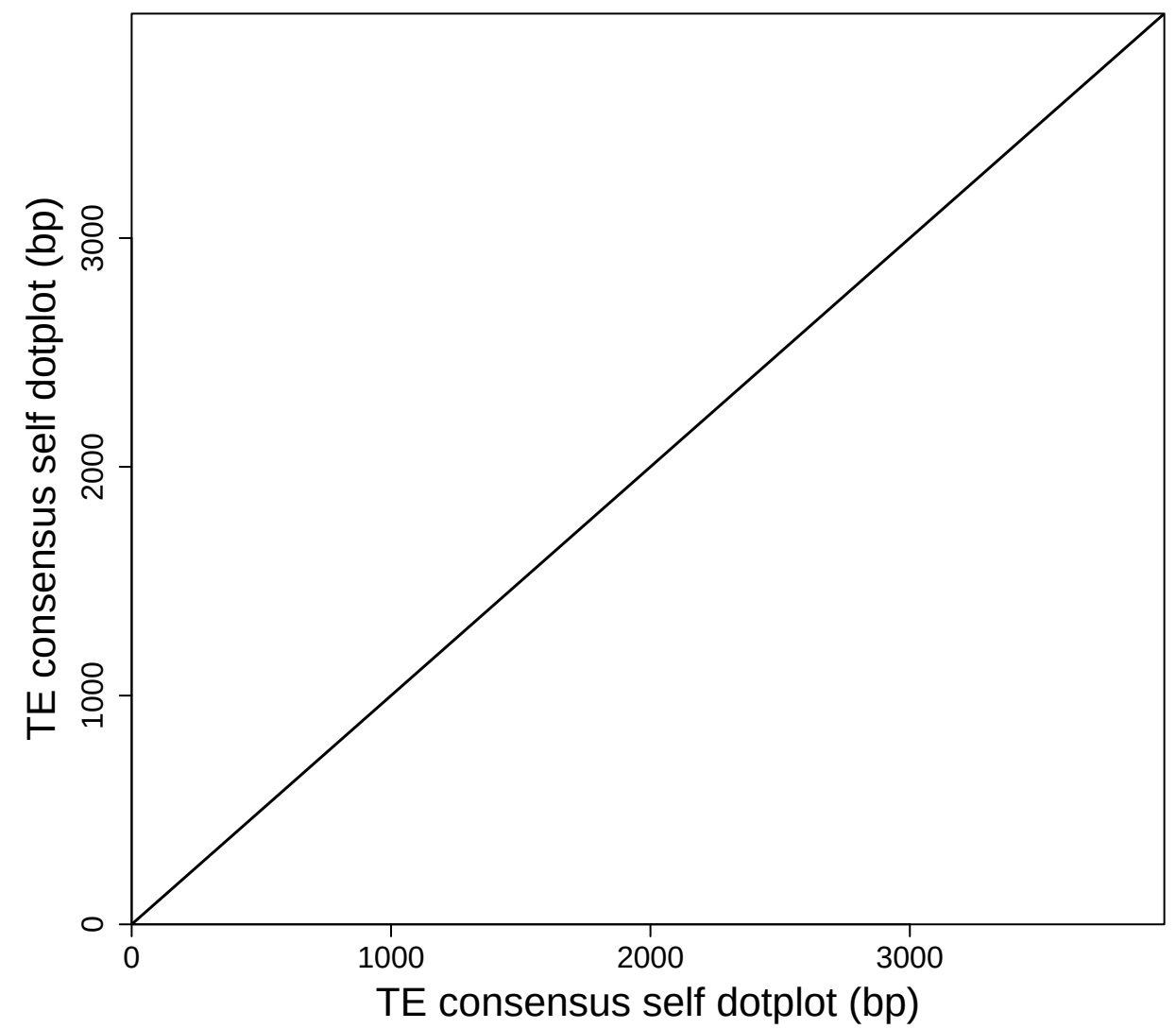
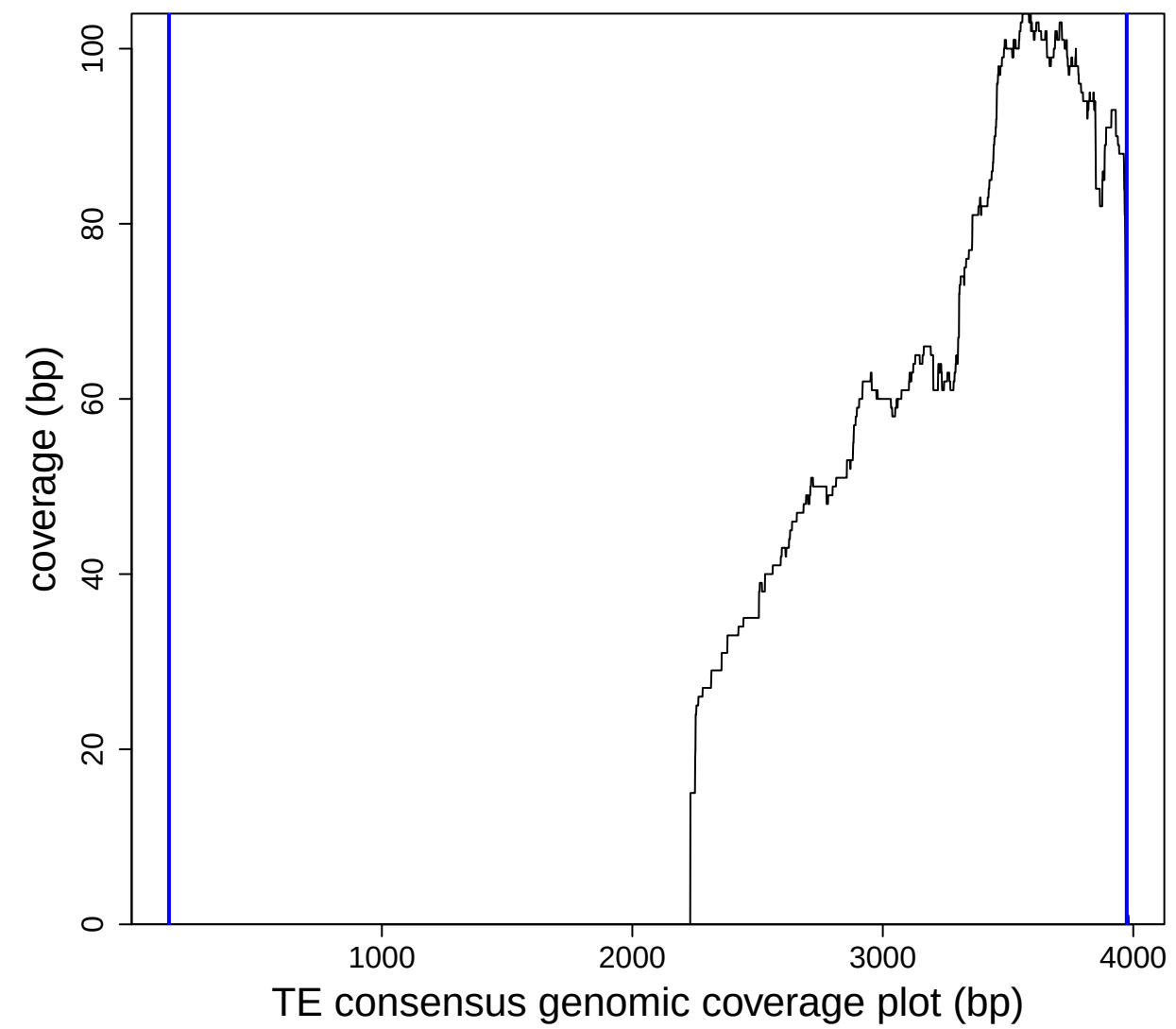
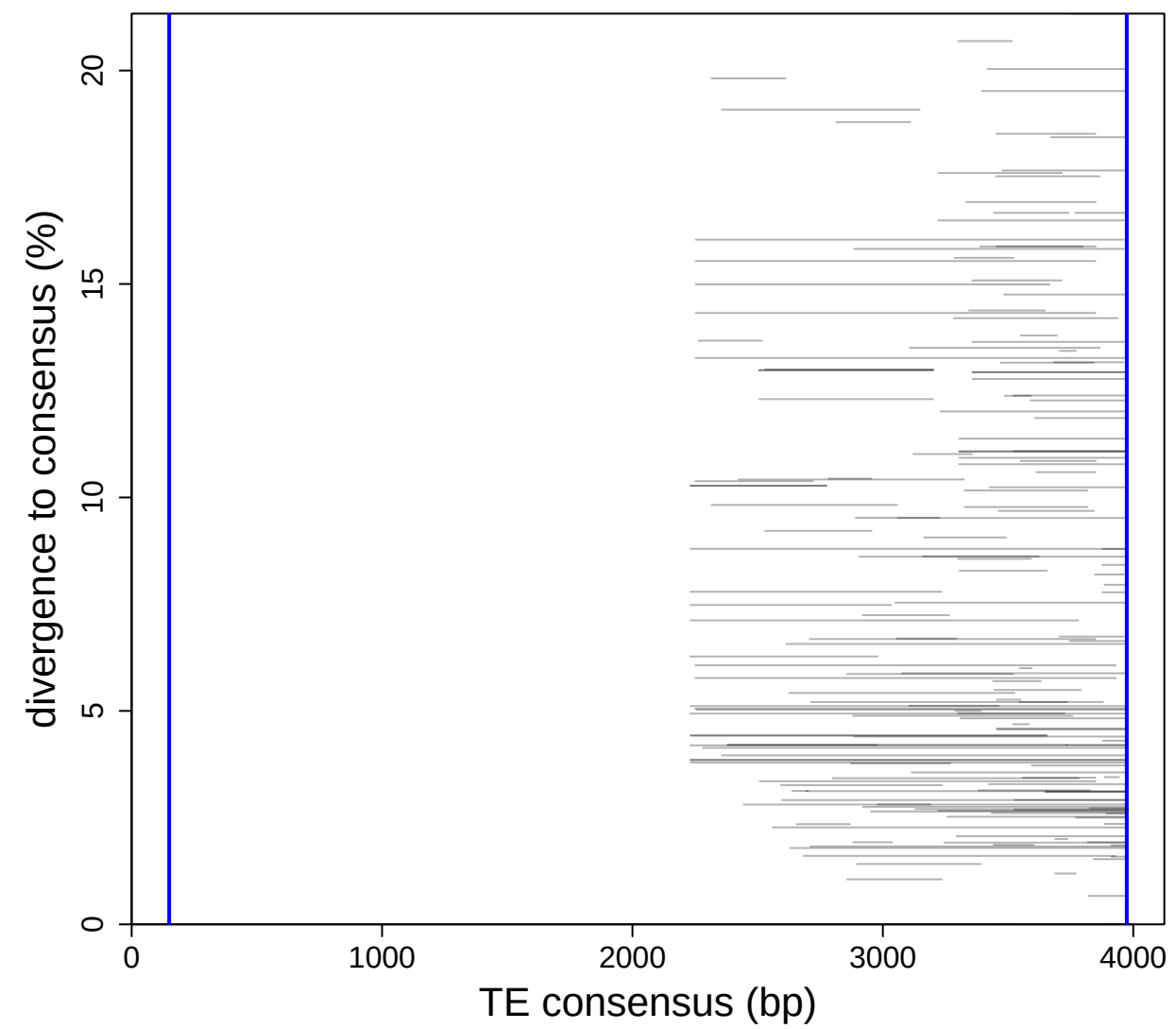
.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.f
size: 3421bp; fragments: 224; full length: 0 (>=3078.9bp)

After TEtrimmer 3421 bp

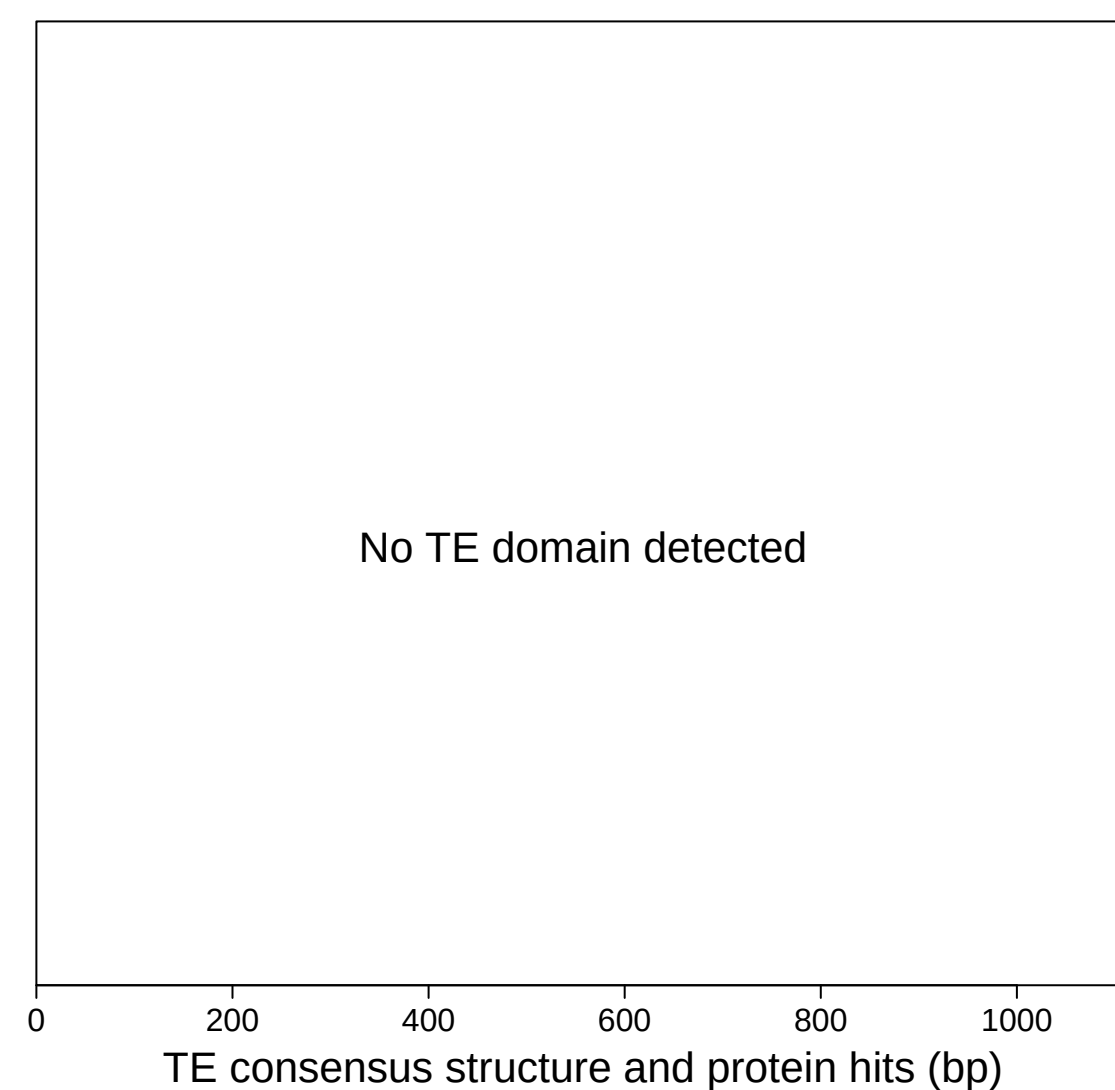
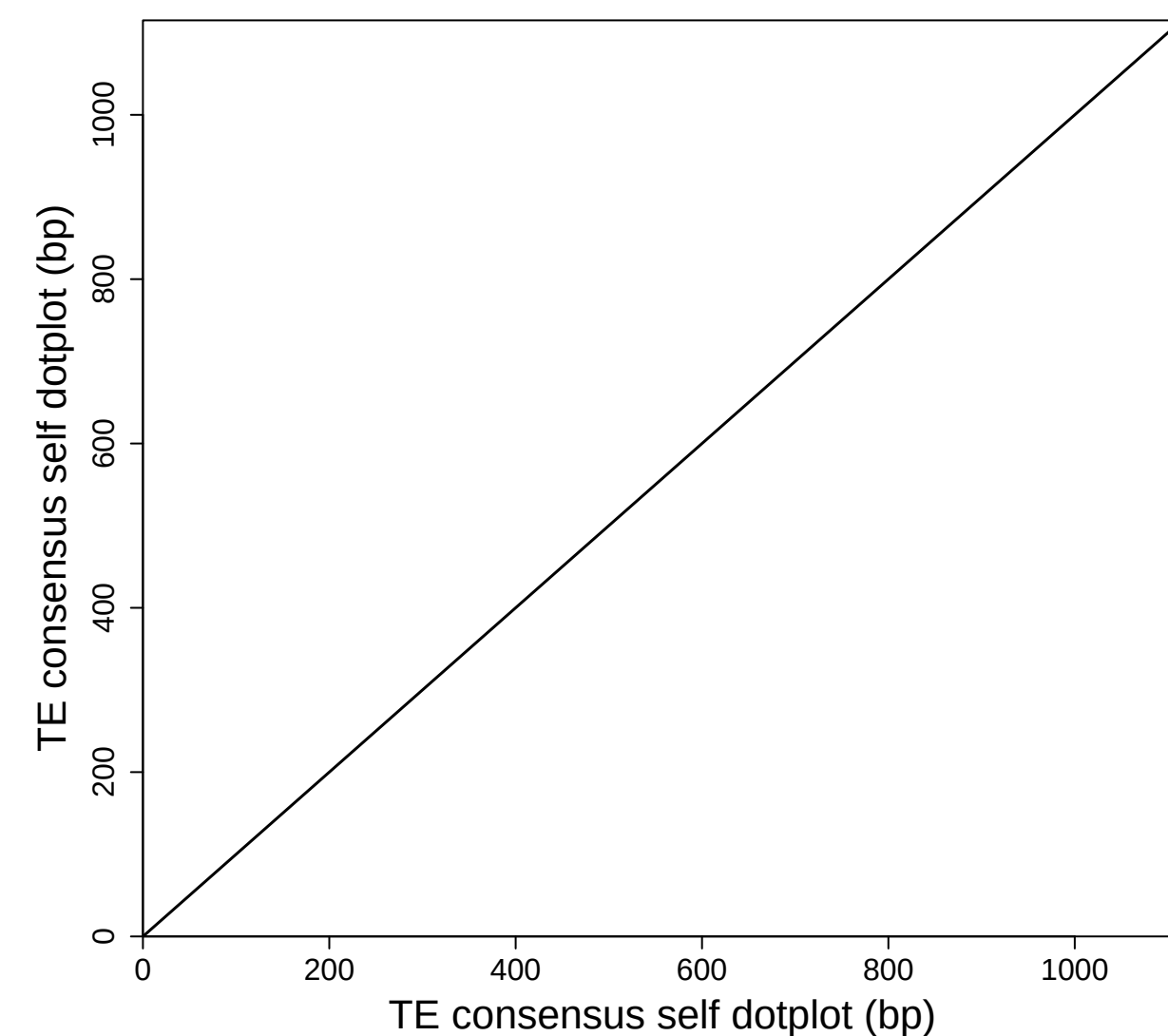
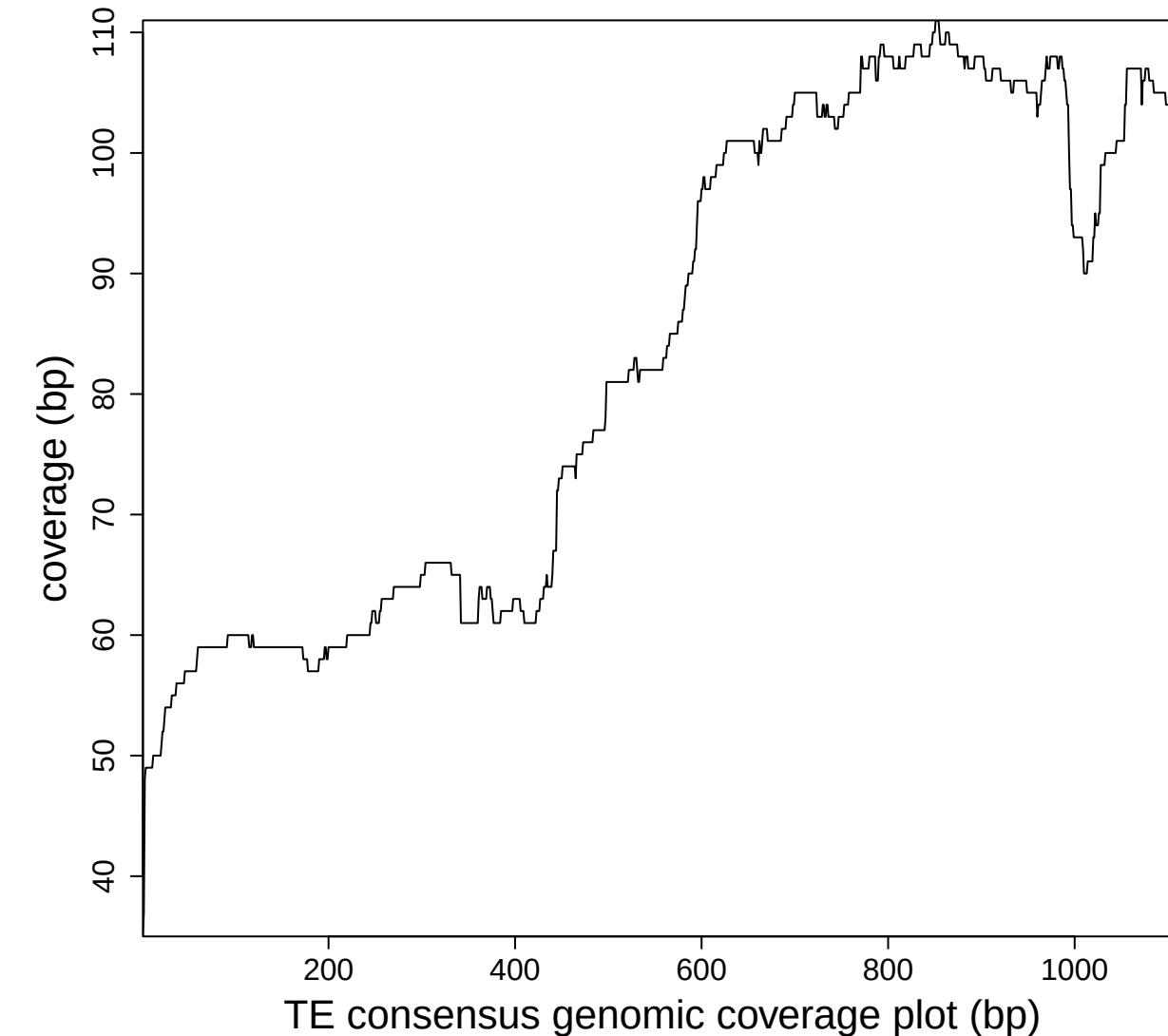
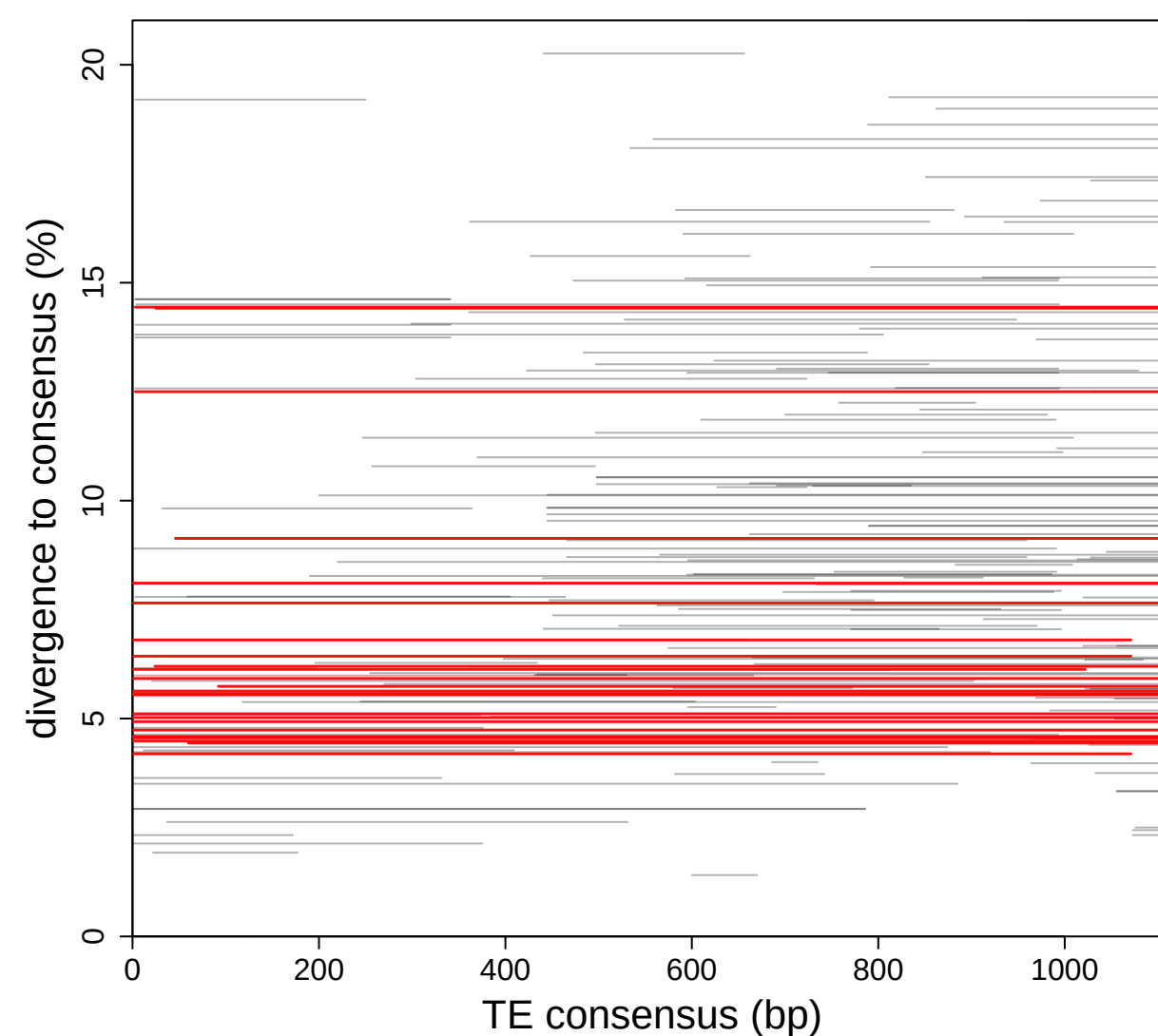


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 4124bp; fragments: 178; full length: 0 (>=3711.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

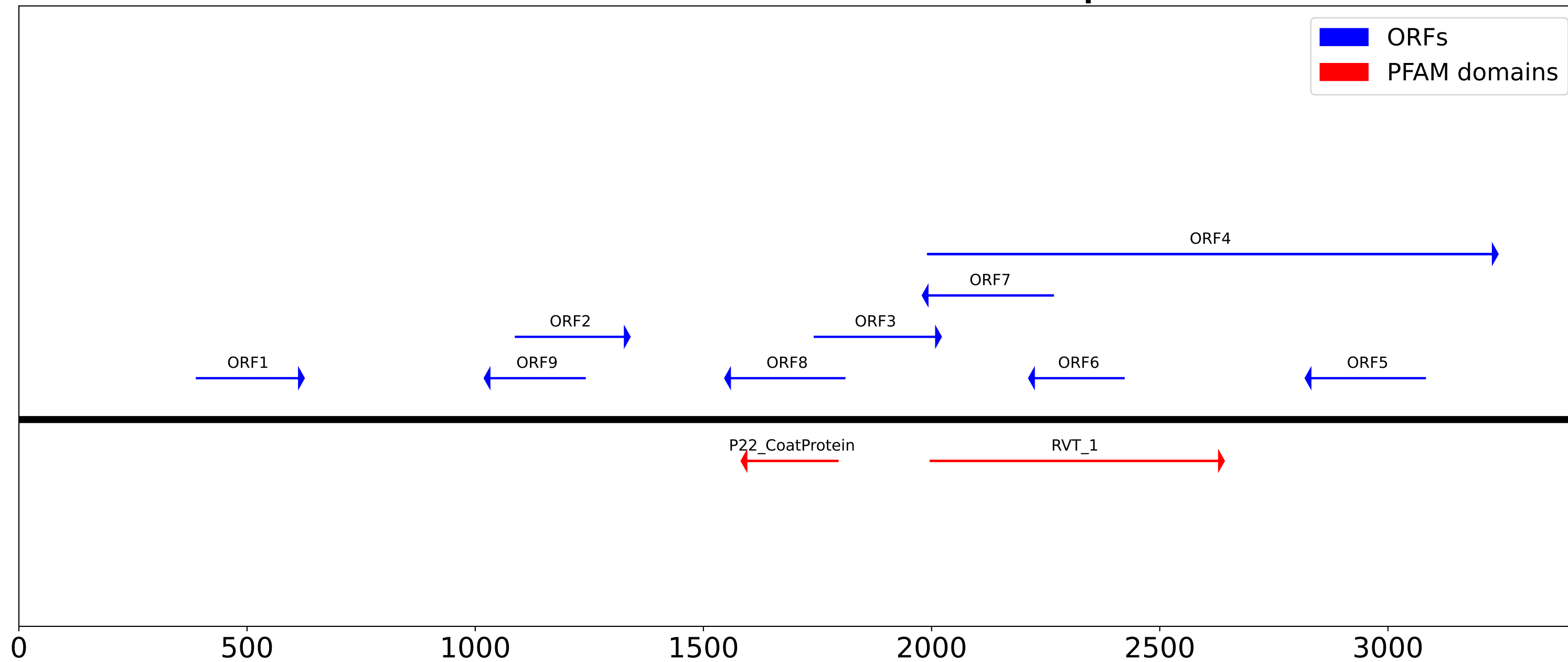


TE: rnd_5_family_3867
size: 1115bp; fragments: 188; full length: 28 (≥ 1003.5 bp)

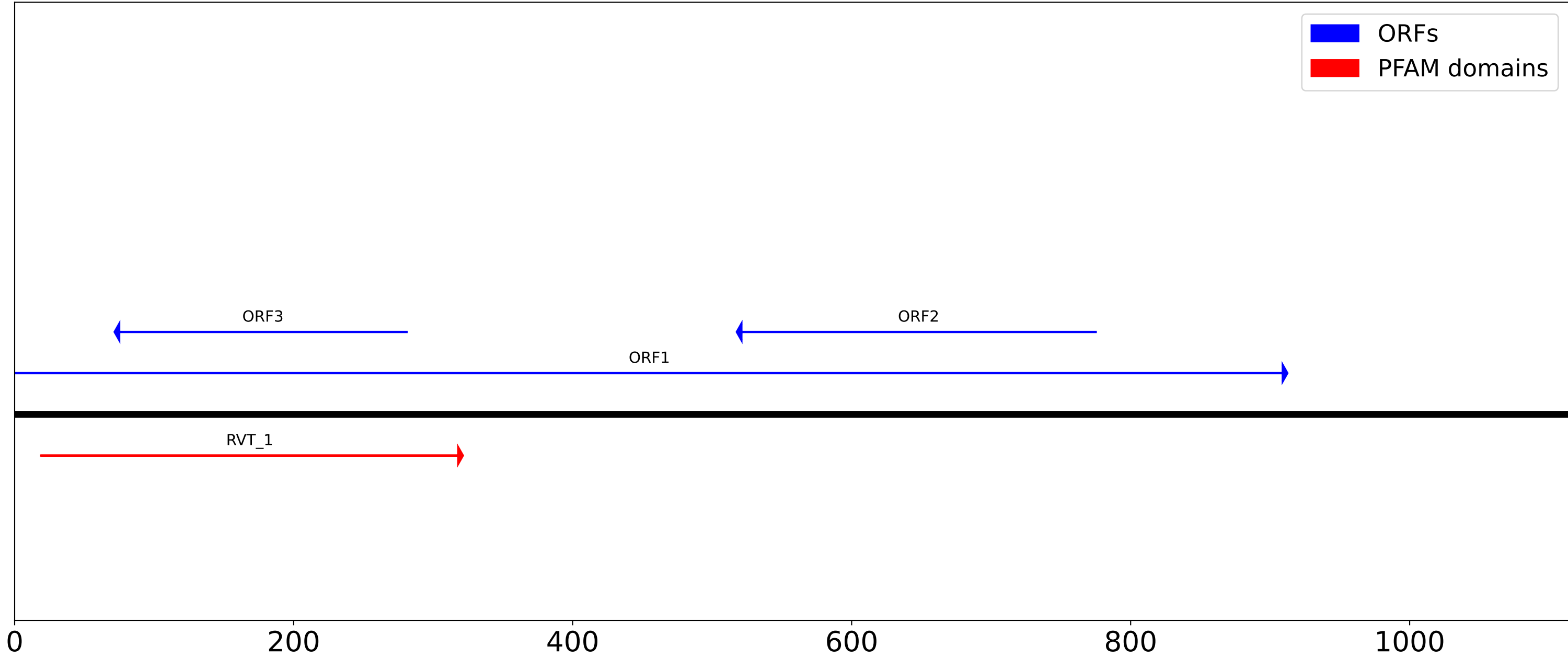


Before TEtrimmer 1115 bp

After TEtrimmer ORF and PFAM domain plot



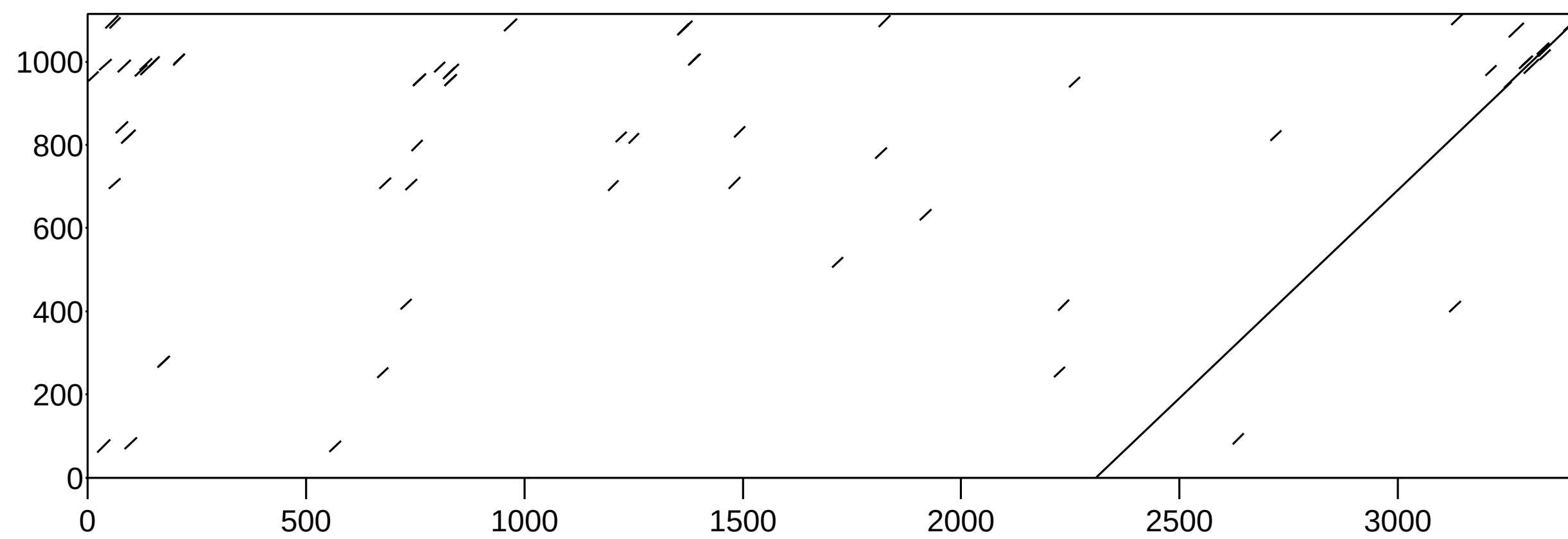
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)