

MSA length = 5704

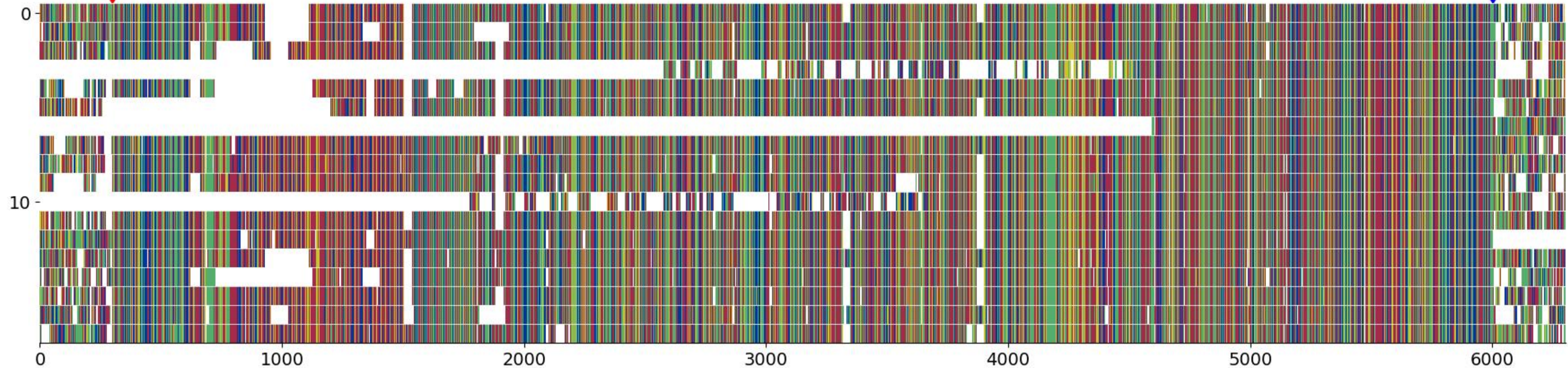
Start crop Point

End crop Point



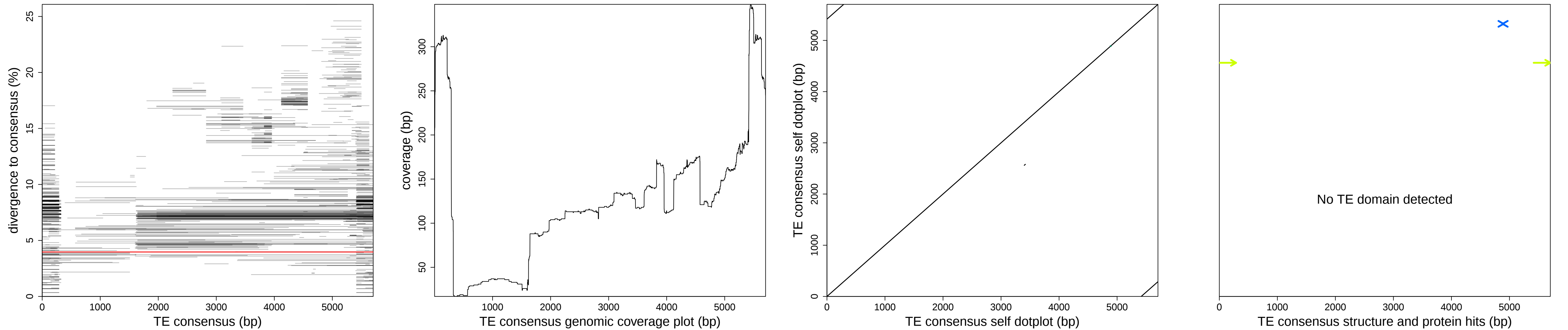
Start crop Point

End crop Point



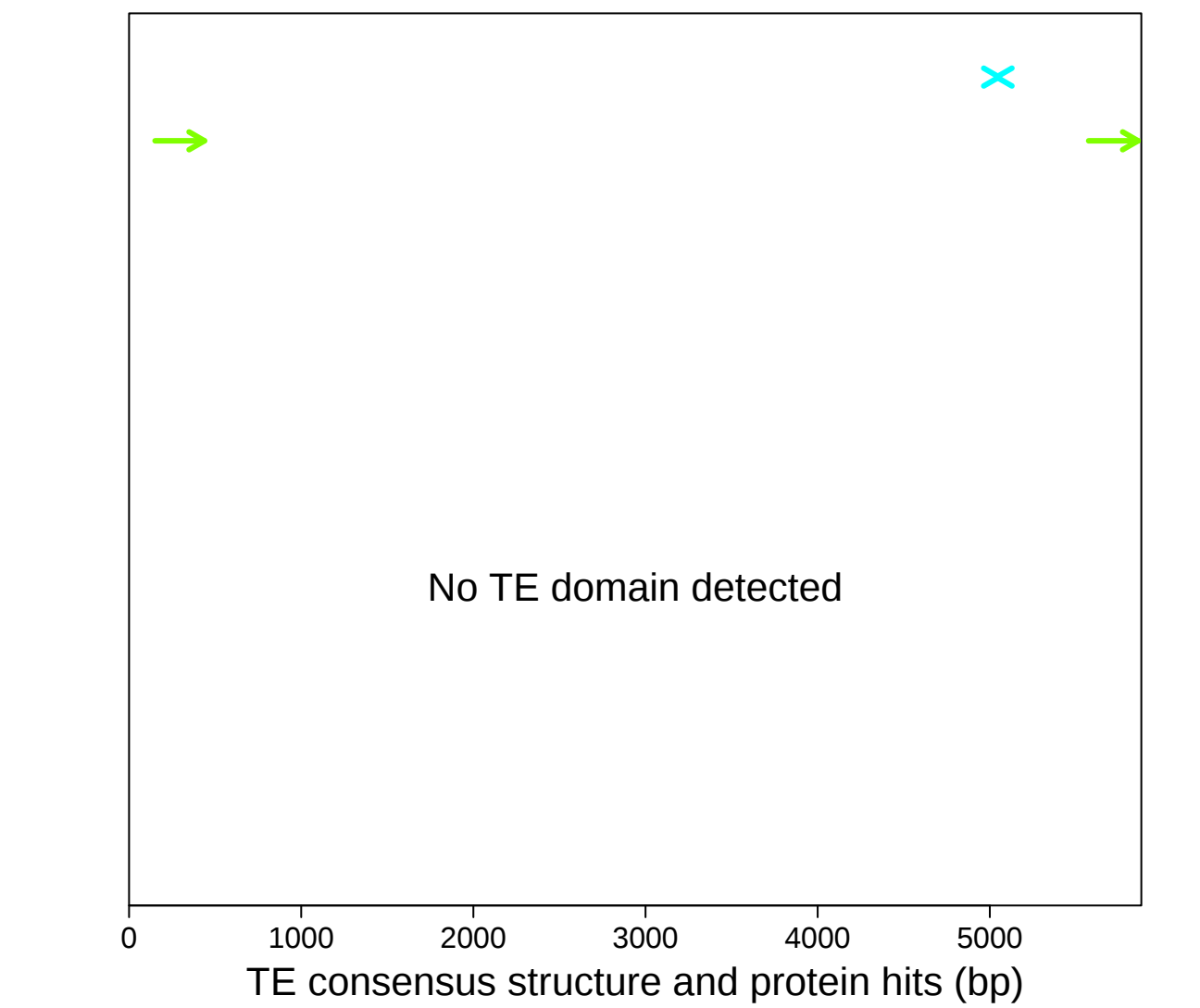
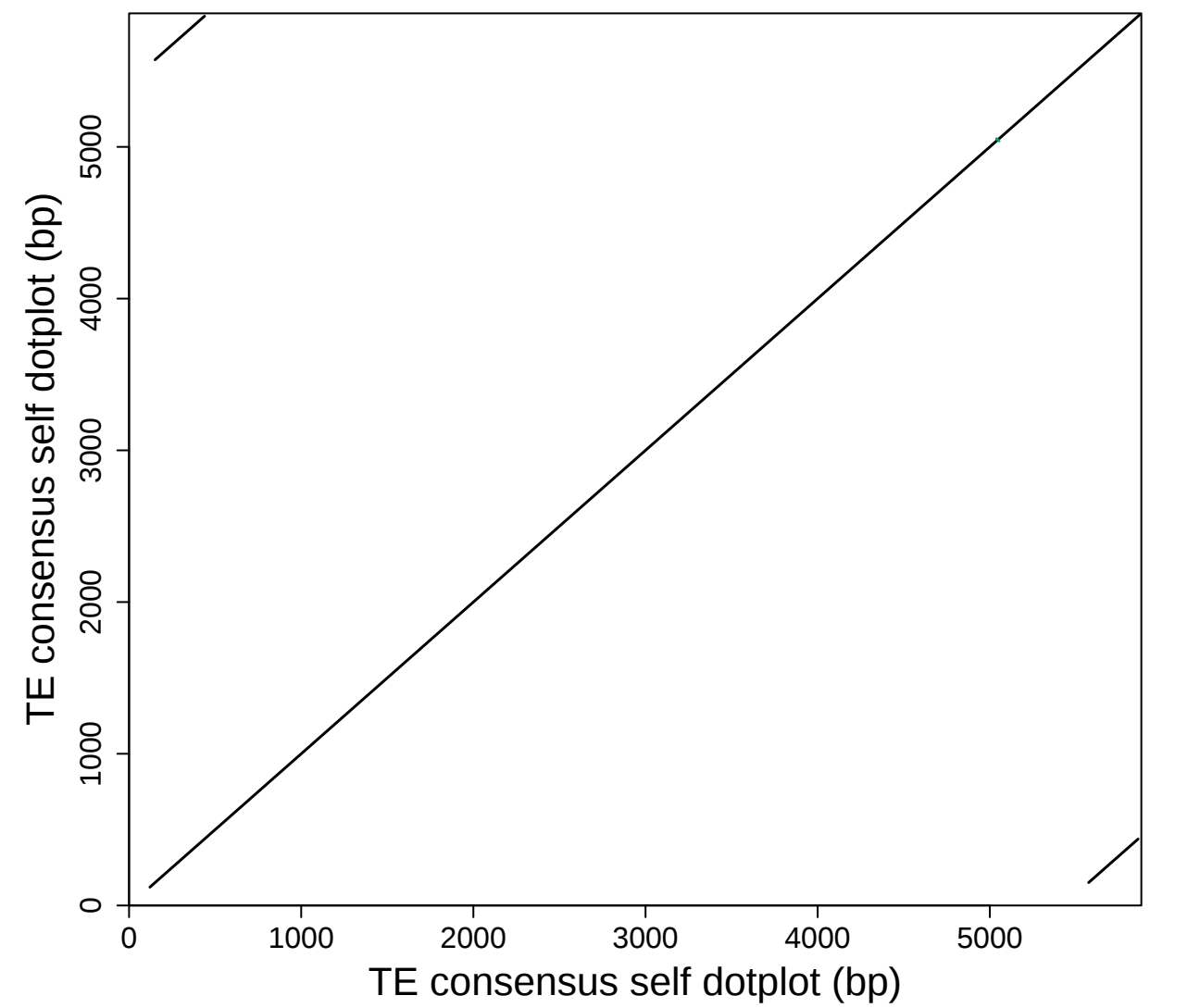
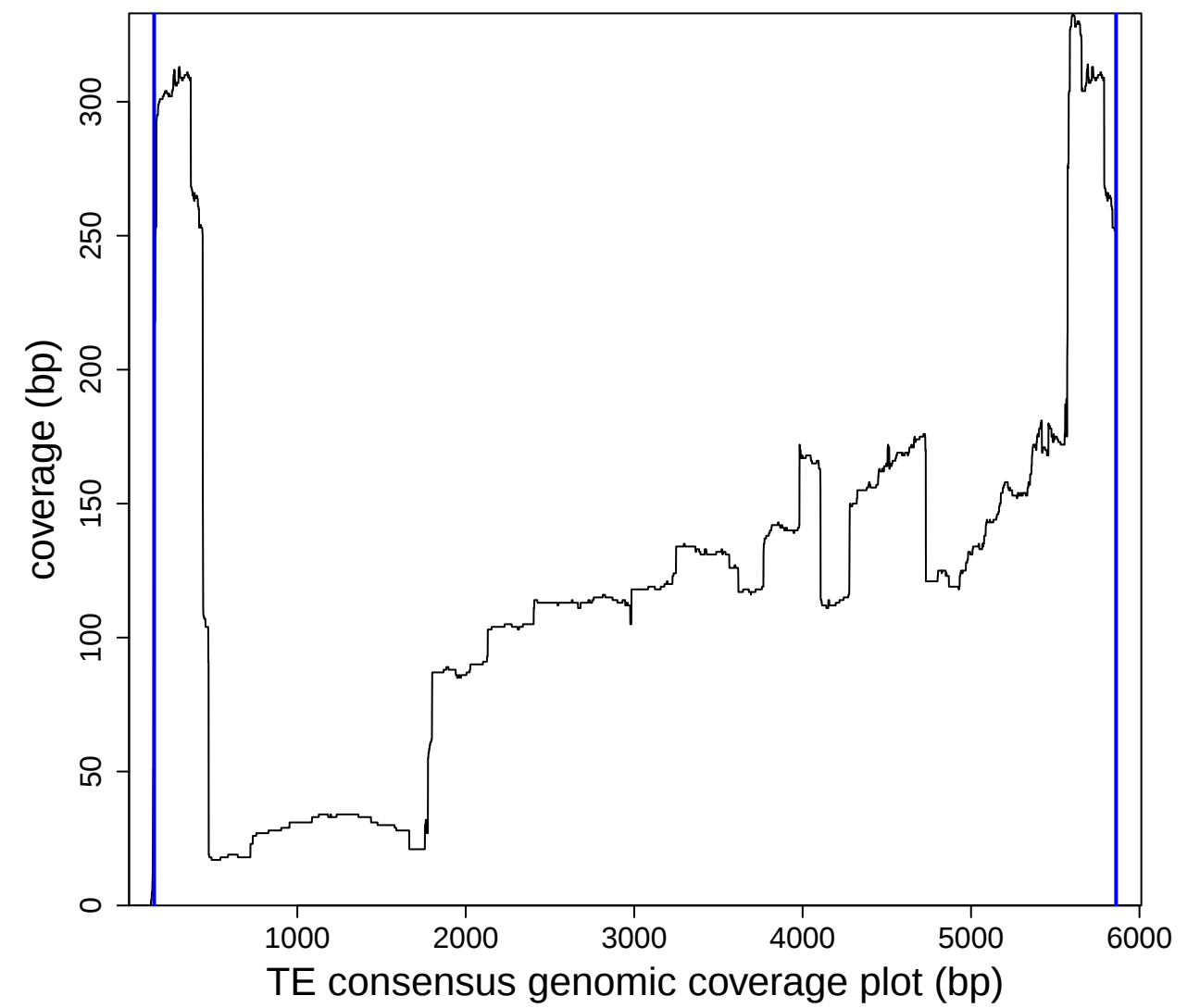
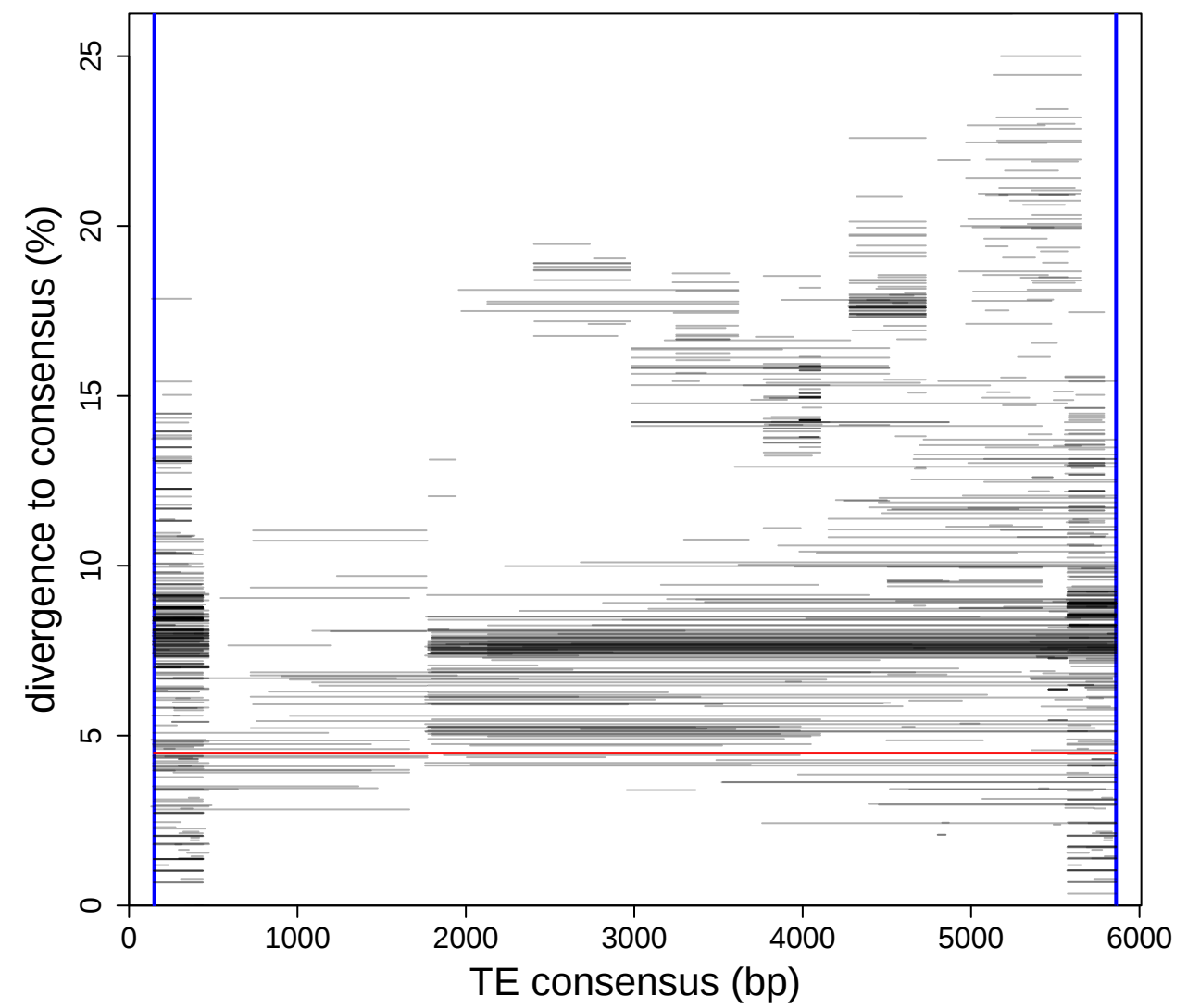
bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.fa
size: 5704bp; fragments: 1034; full length: 1 (>=5133.6bp)

After TETrimmer 5704 bp

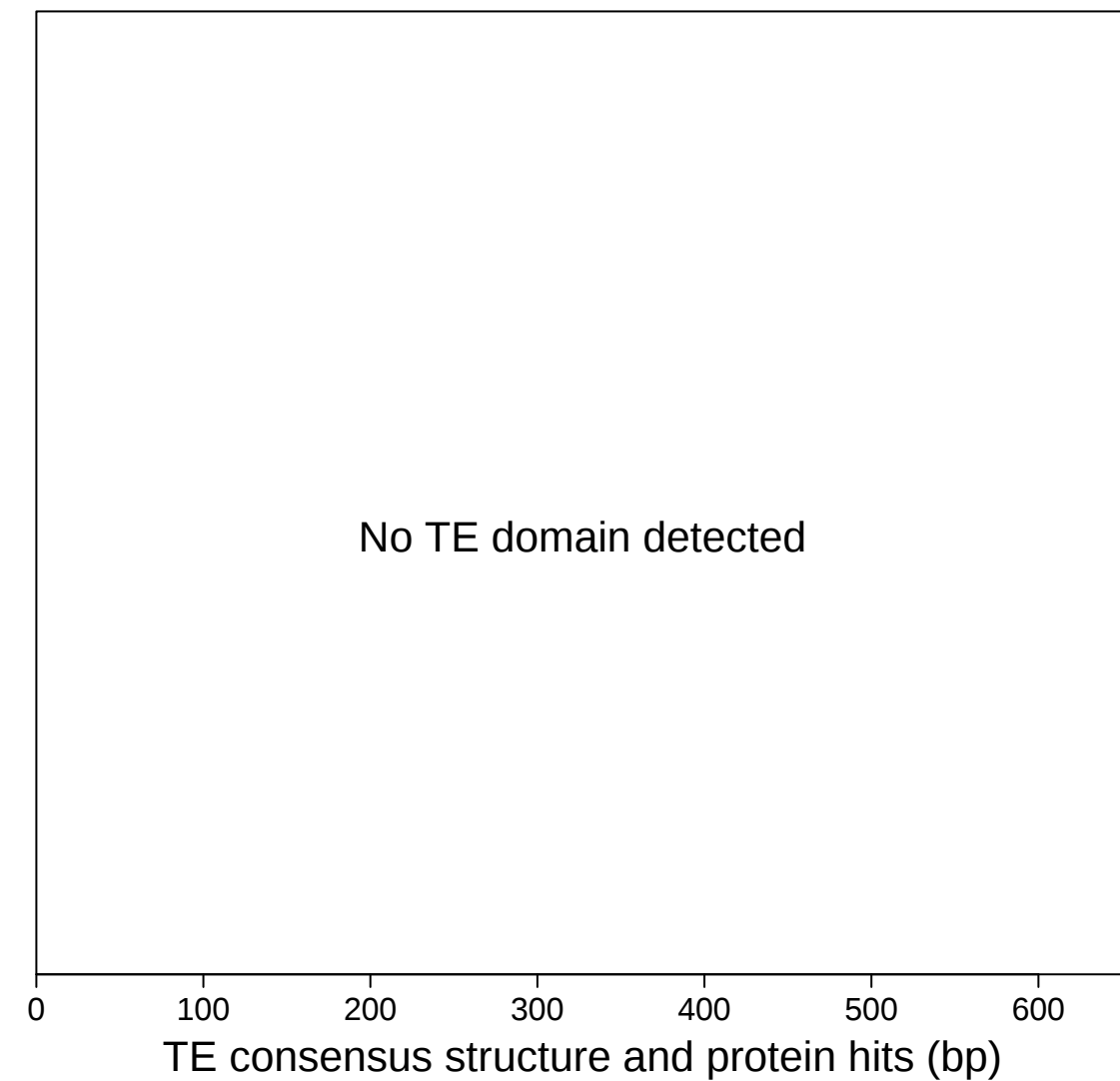
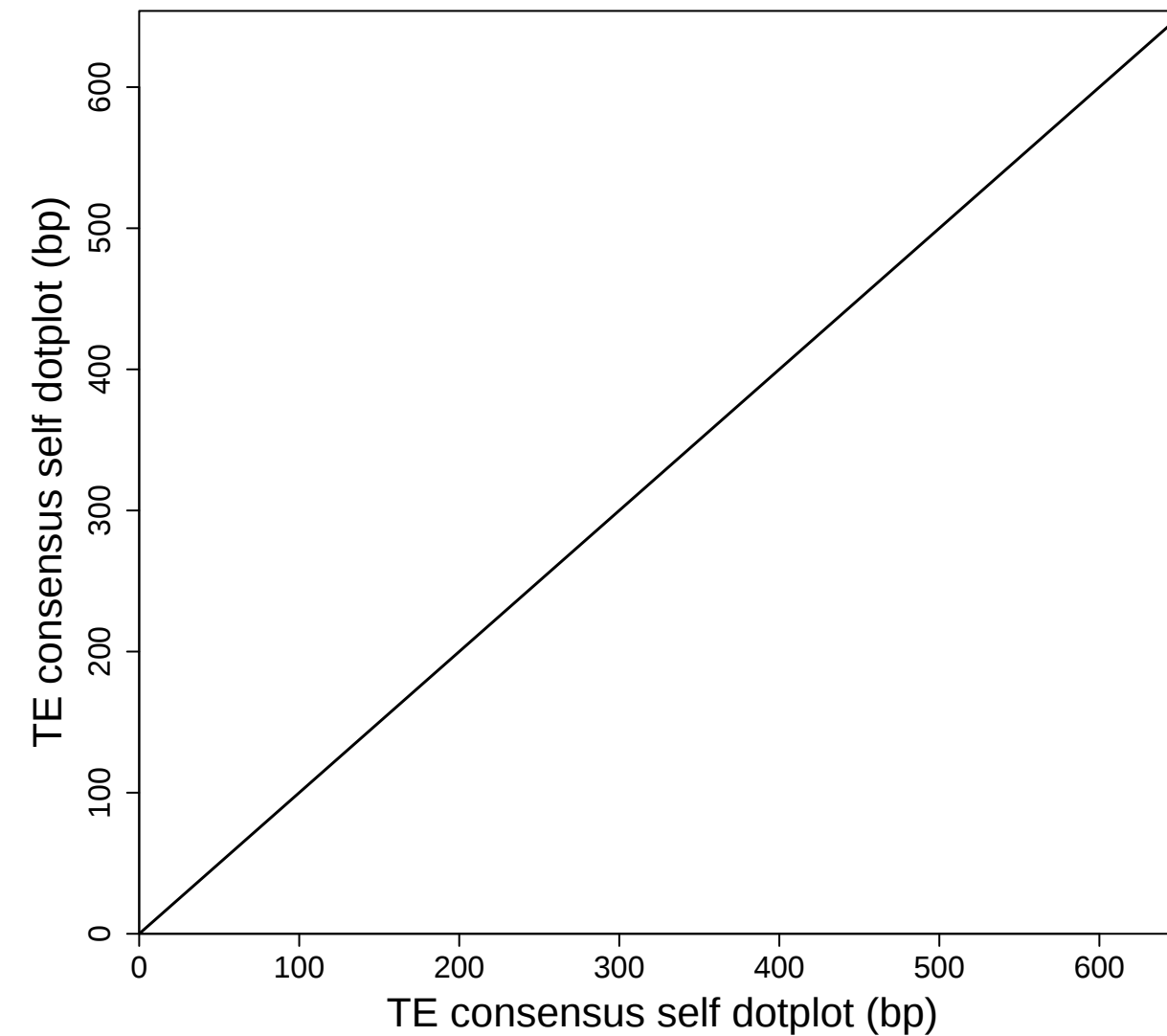
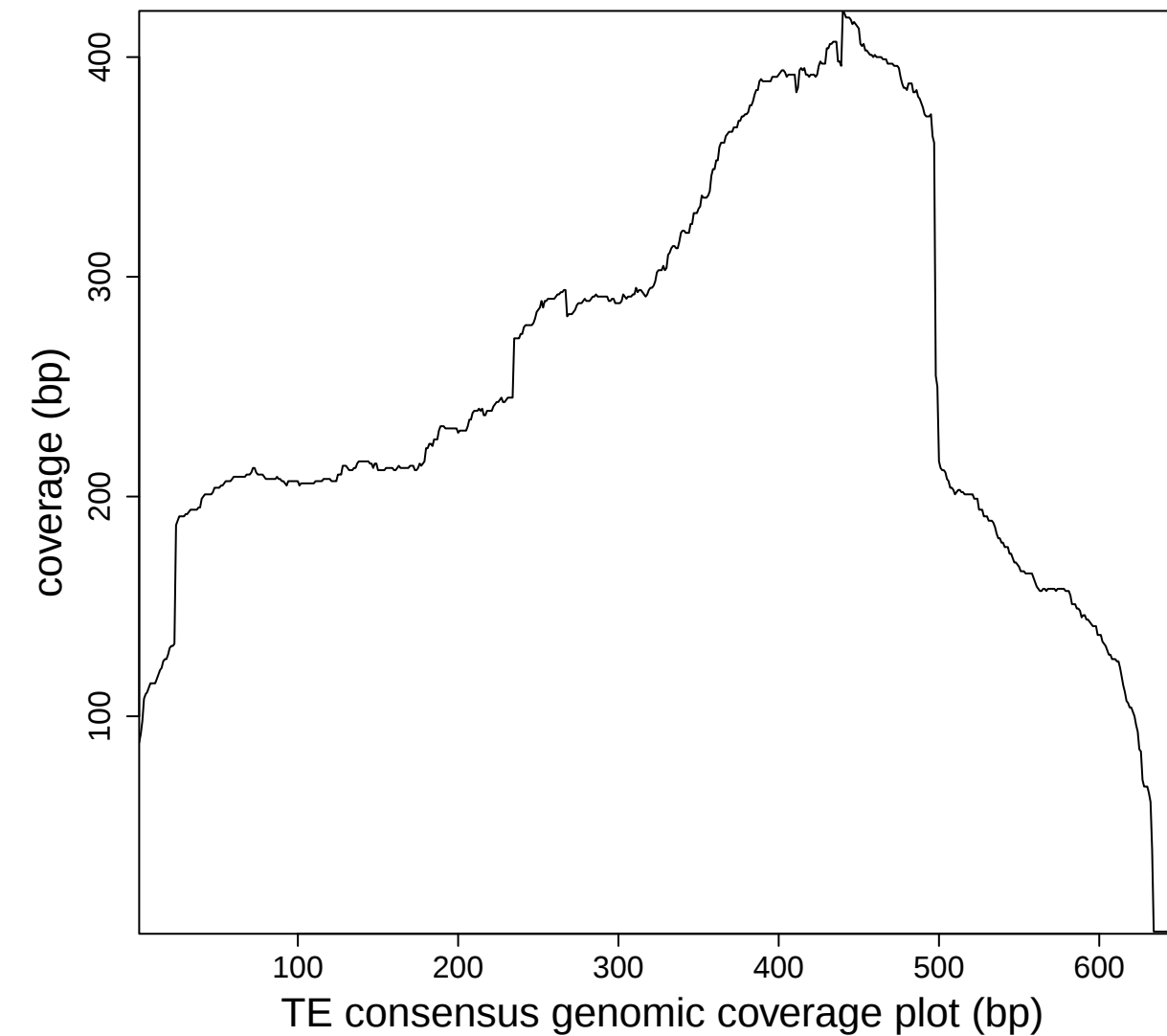
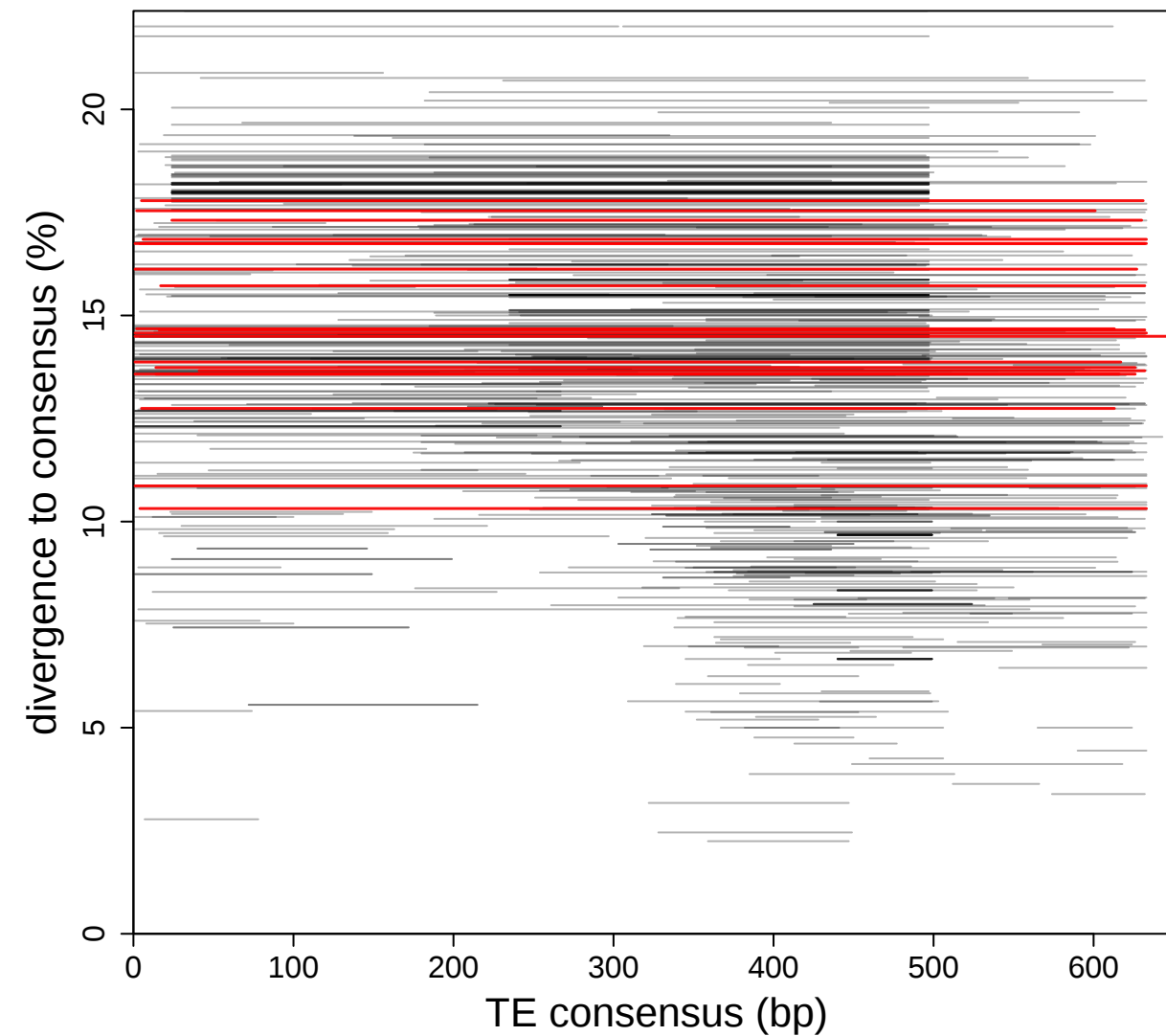


bed_uf.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0
size: 6011bp; fragments: 1012; full length: 1 (>=5409.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

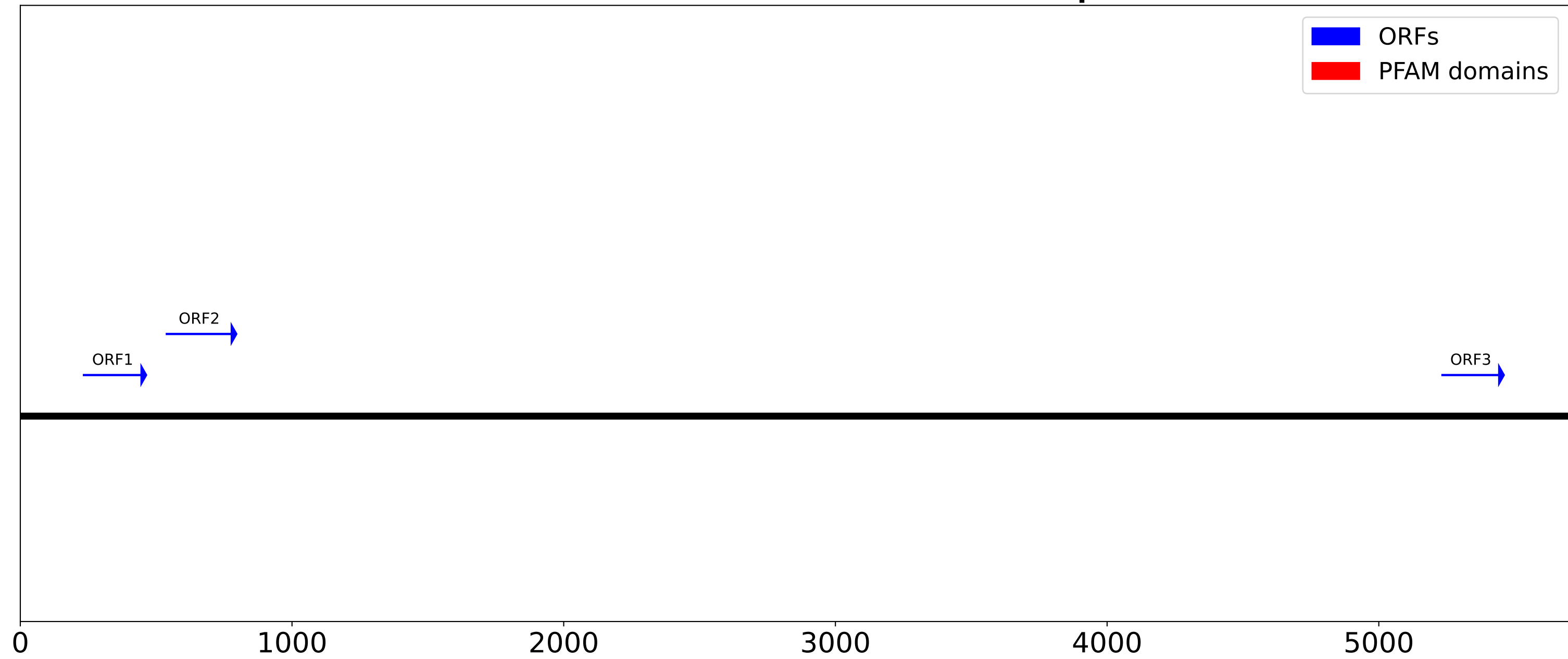


TE: rnd_4_family_778
size: 654bp; fragments: 565; full length: 20 (≥ 588.6 bp)

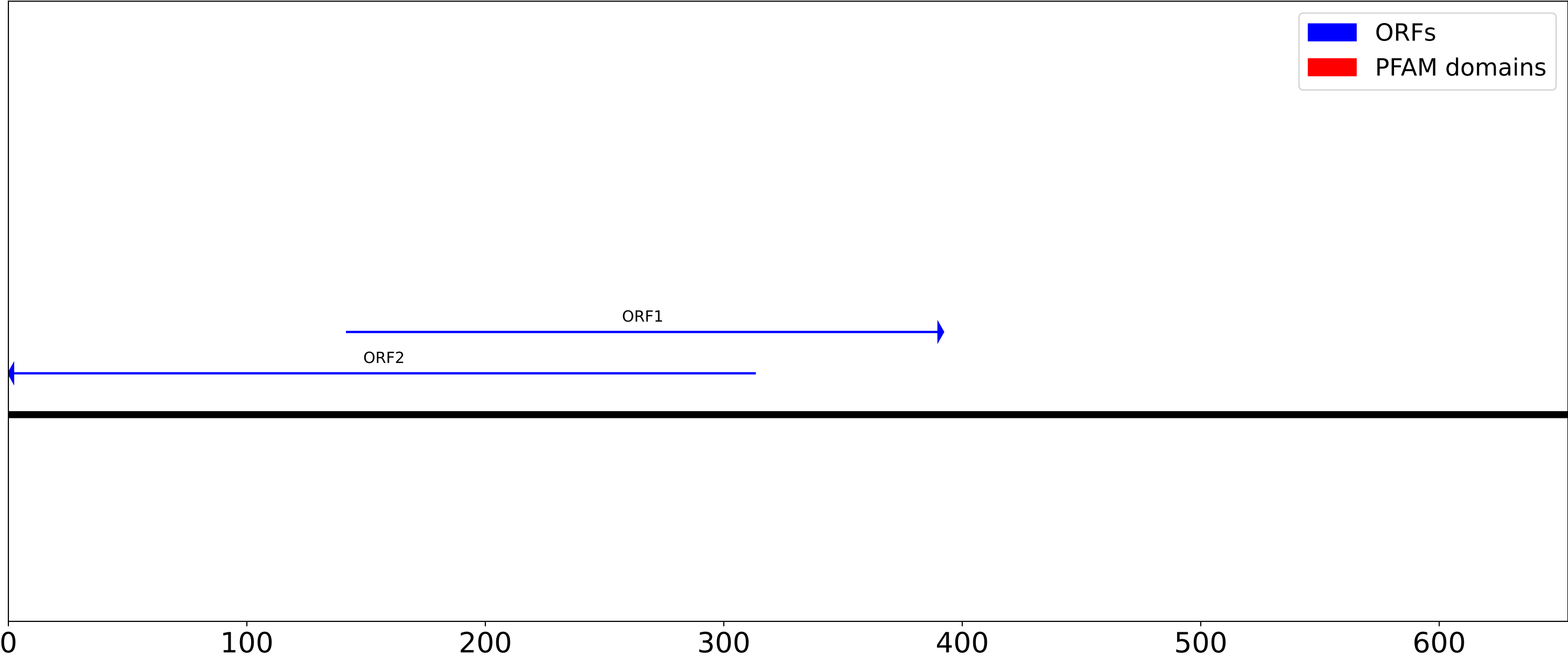


Before TEtrimmer 654 bp

After TEtrimmer ORF and PFAM domain plot



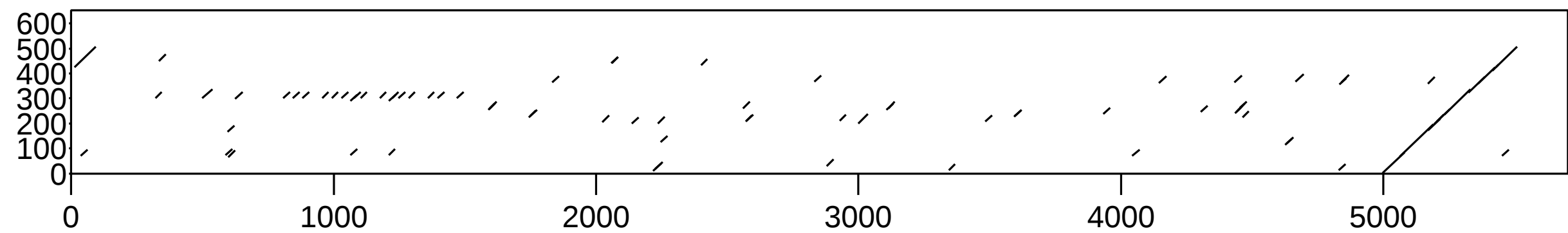
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

sensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)