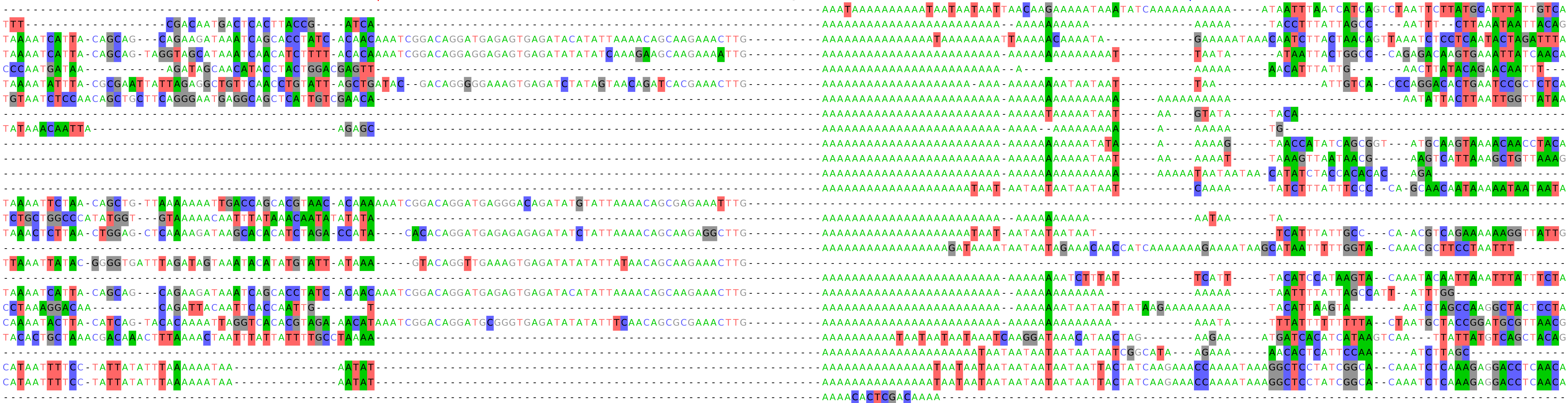


MSA length = 4221

Start crop Point

End crop Point



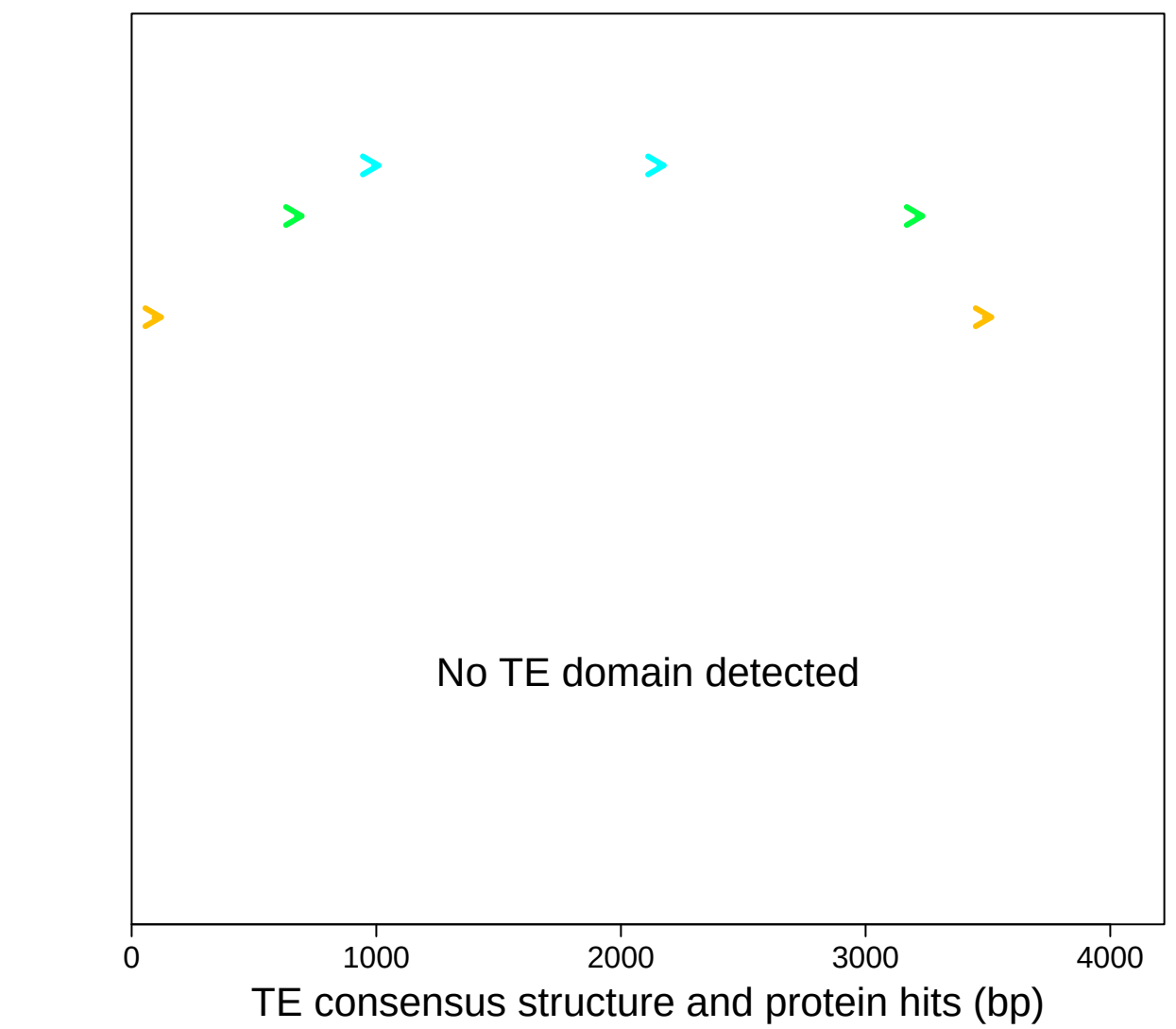
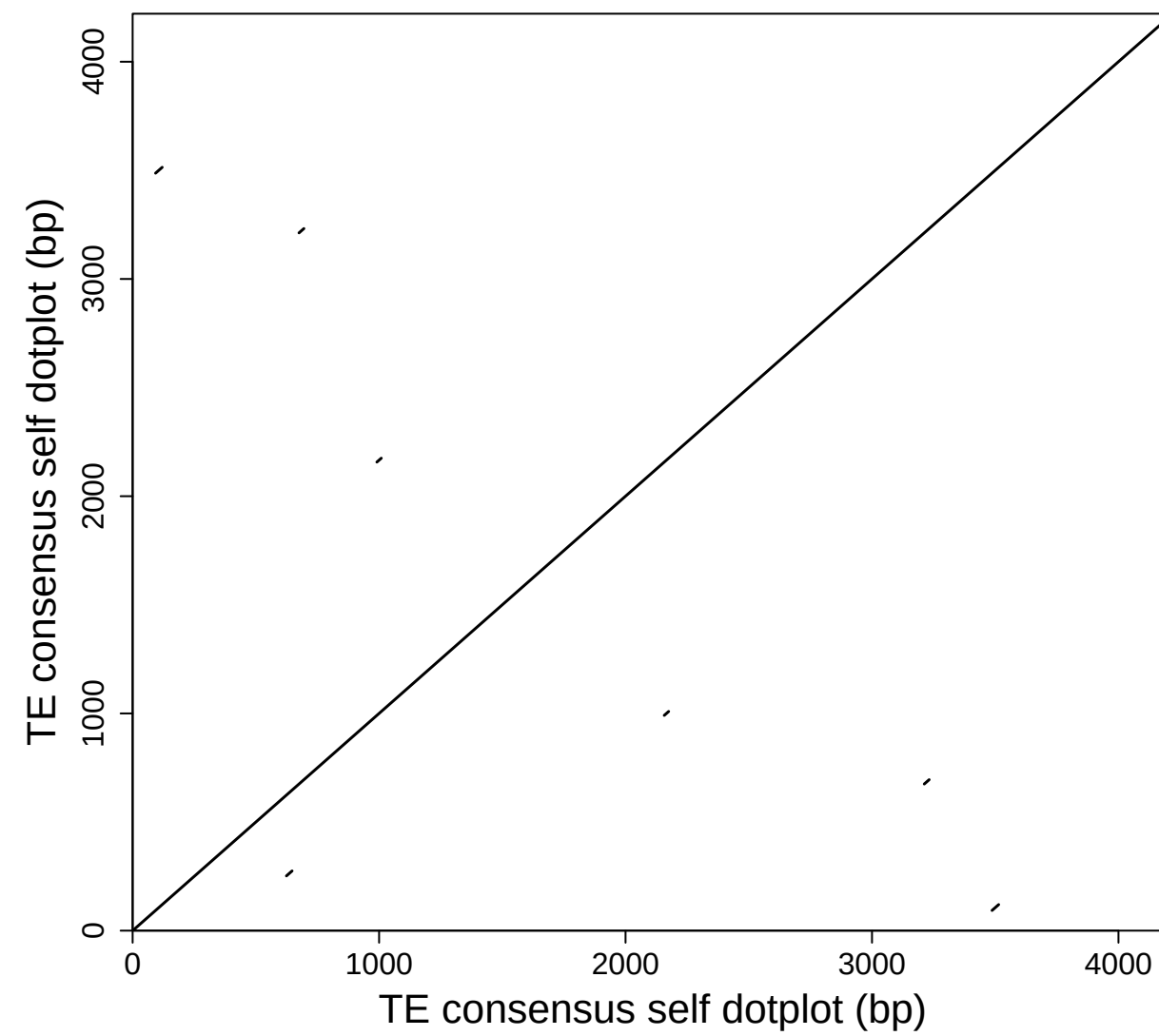
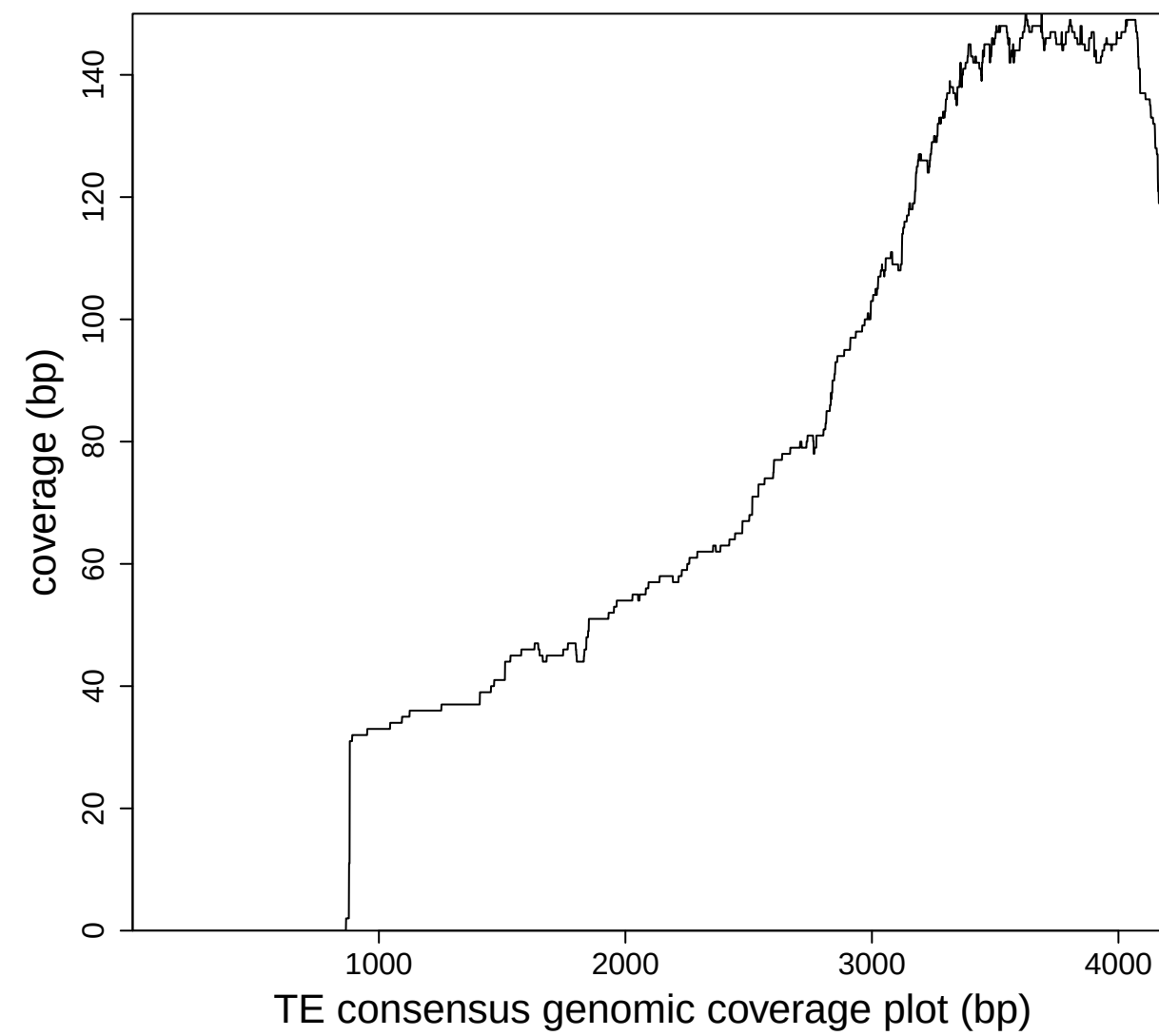
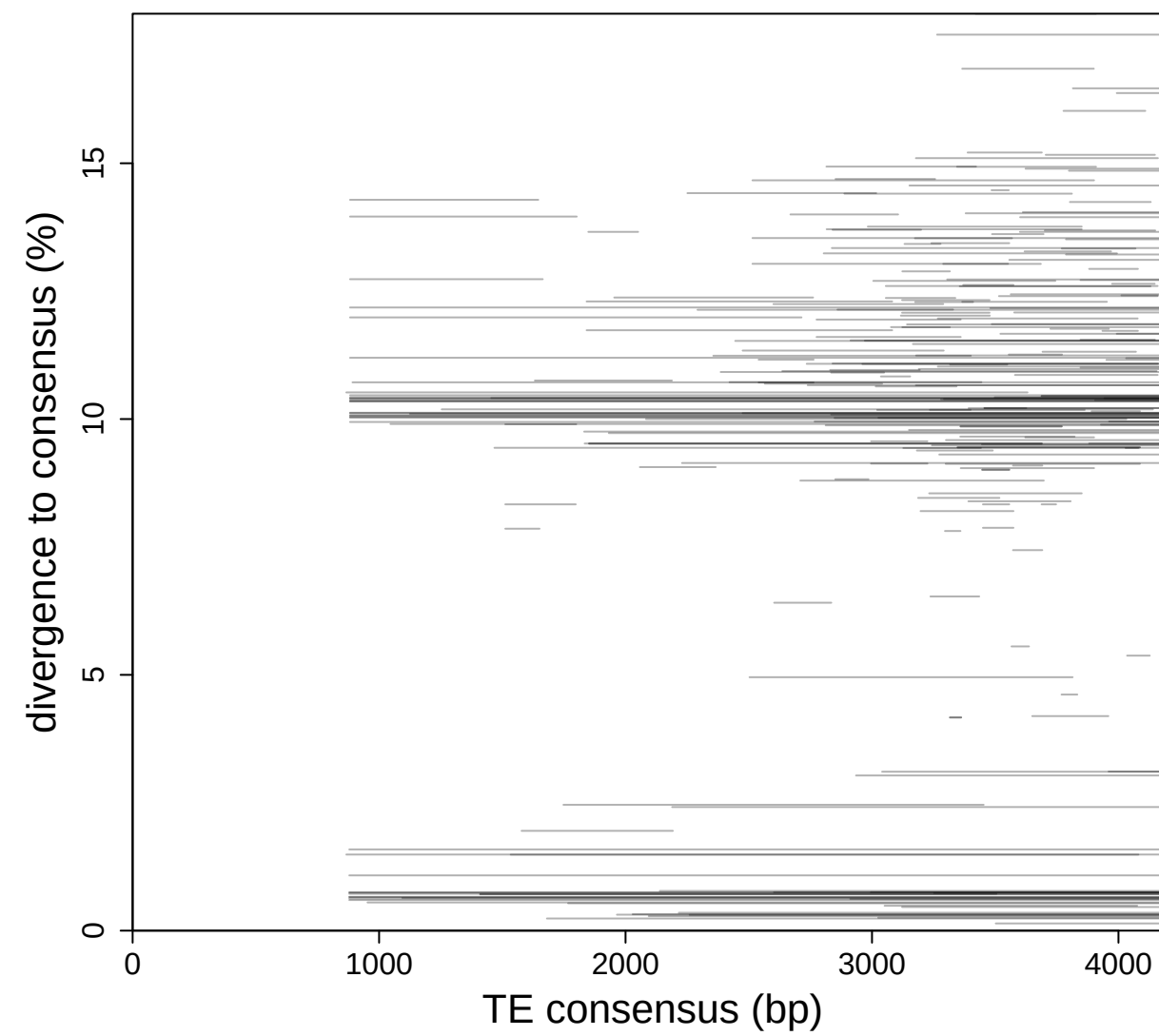
Start crop Point

End crop Point



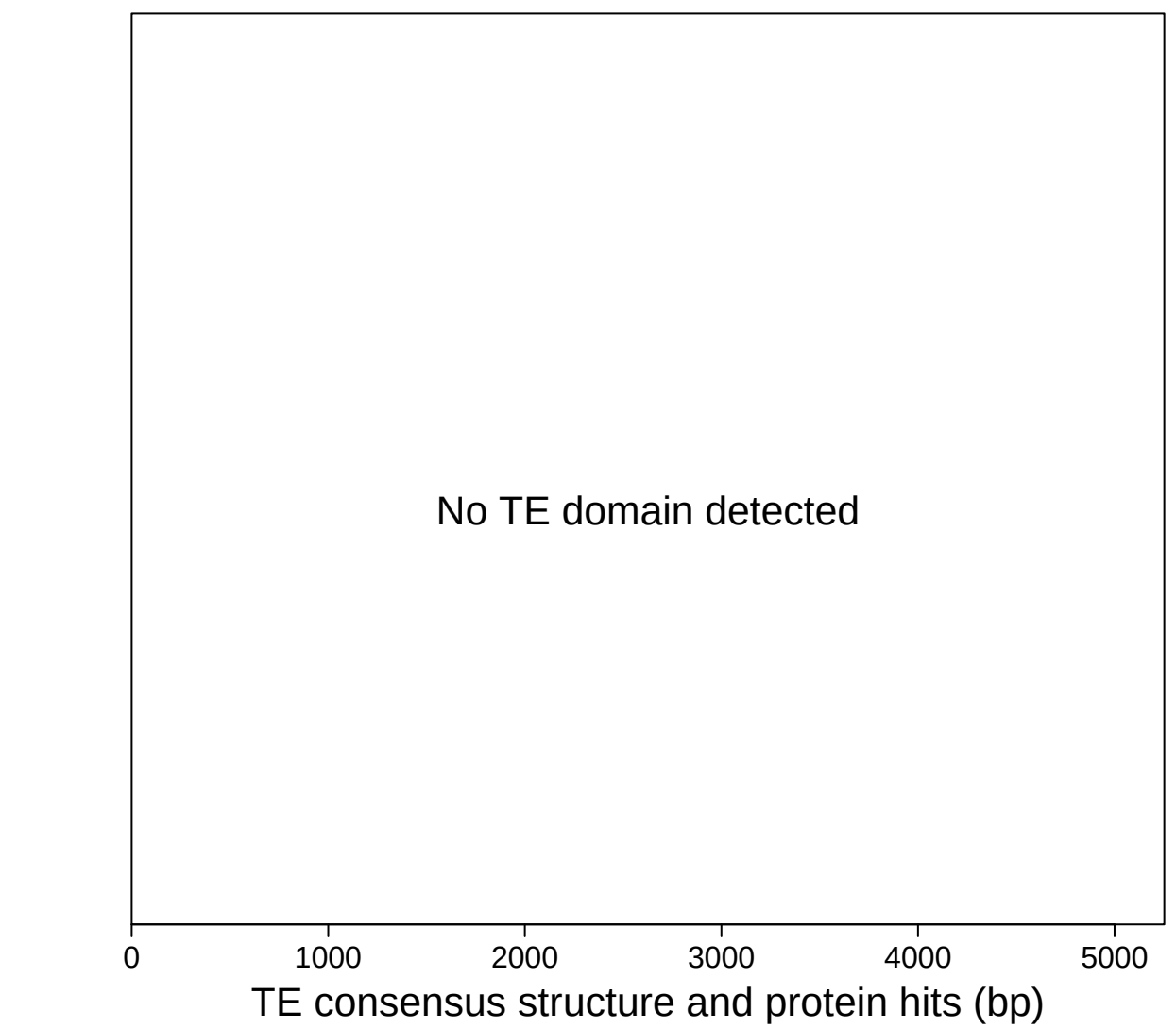
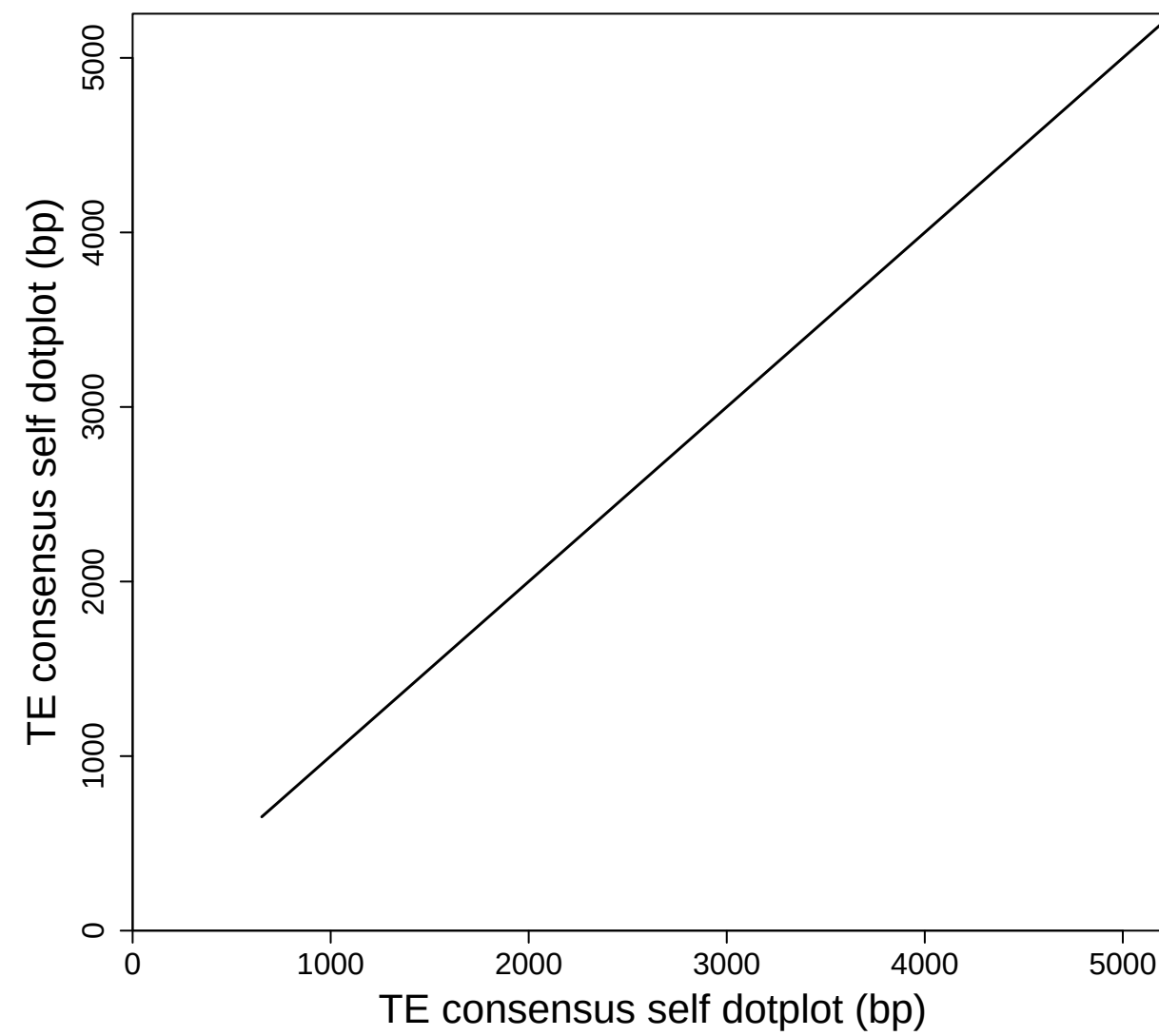
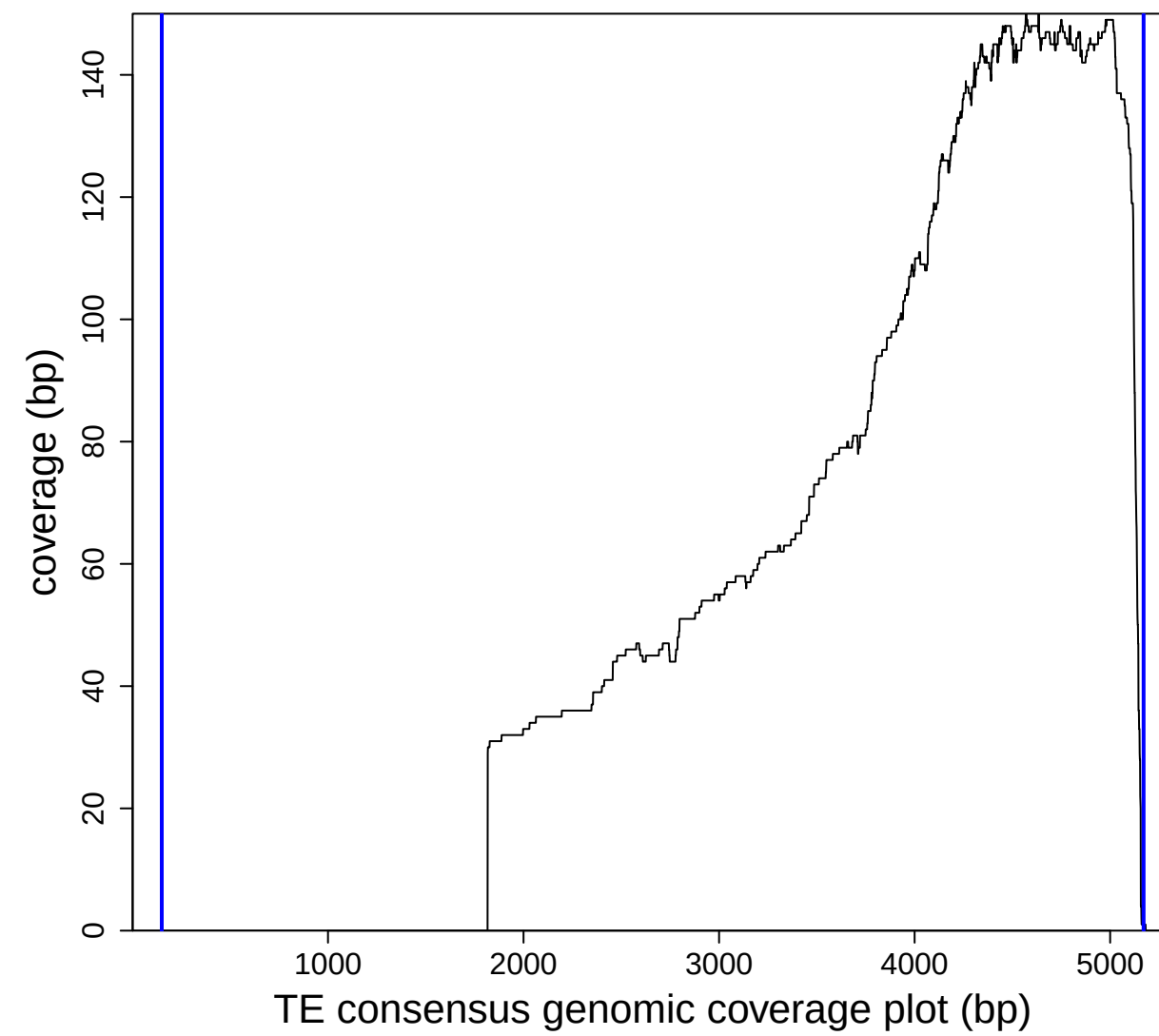
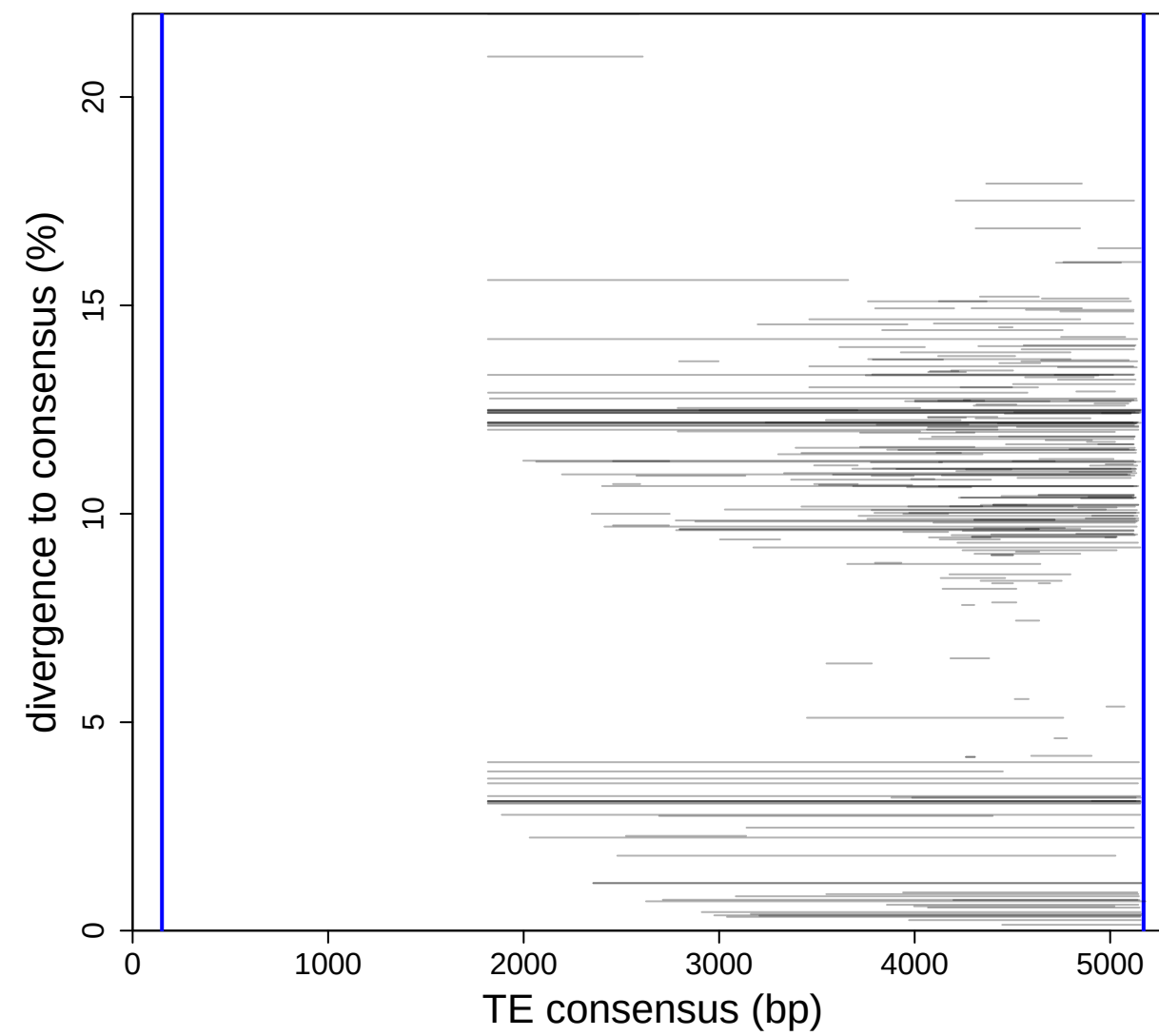
.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.f
size: 4221bp; fragments: 253; full length: 0 (>=3798.9bp)

After TETrimmer 4221 bp



f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 5321bp; fragments: 253; full length: 0 (>=4788.9bp)

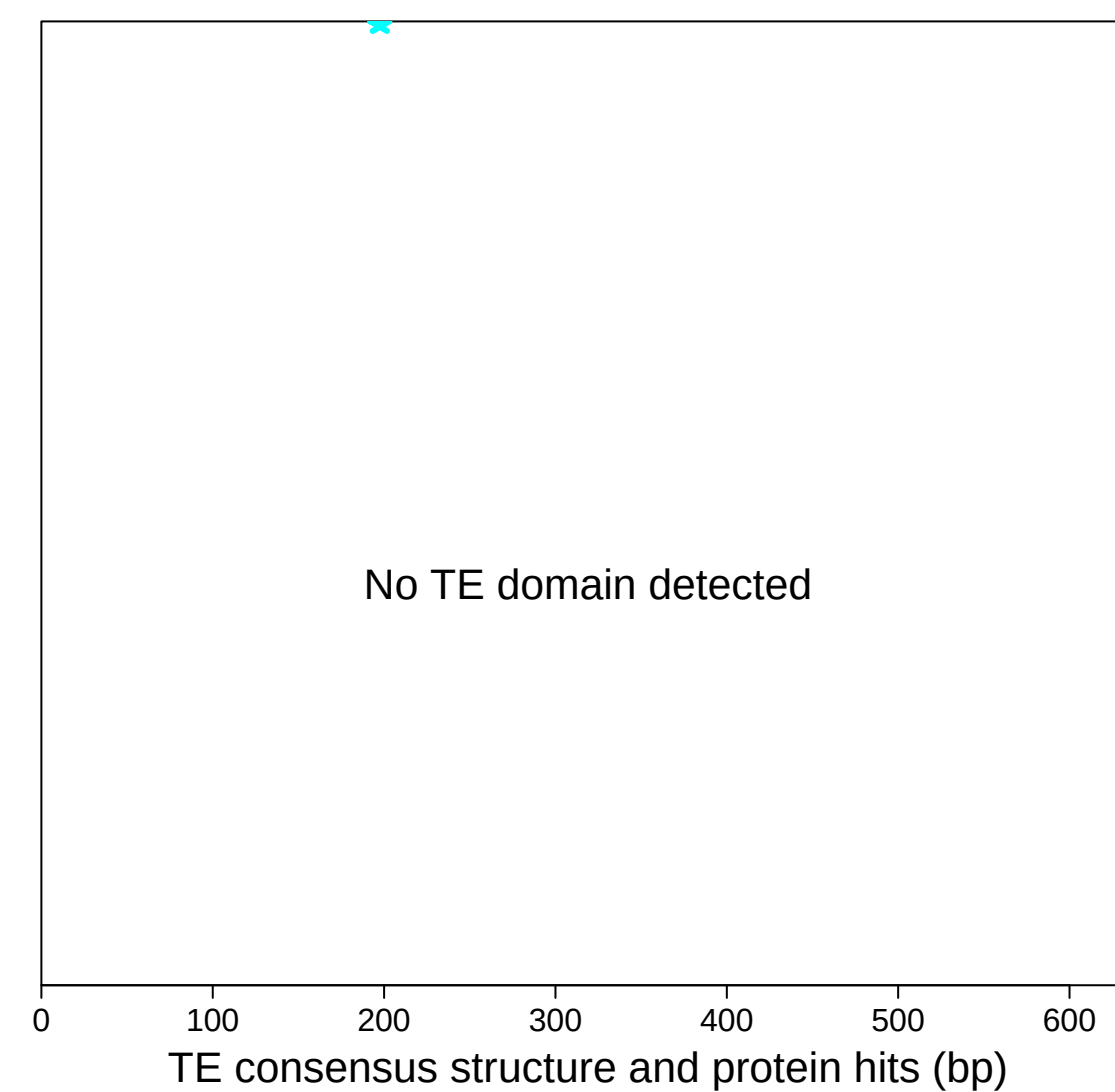
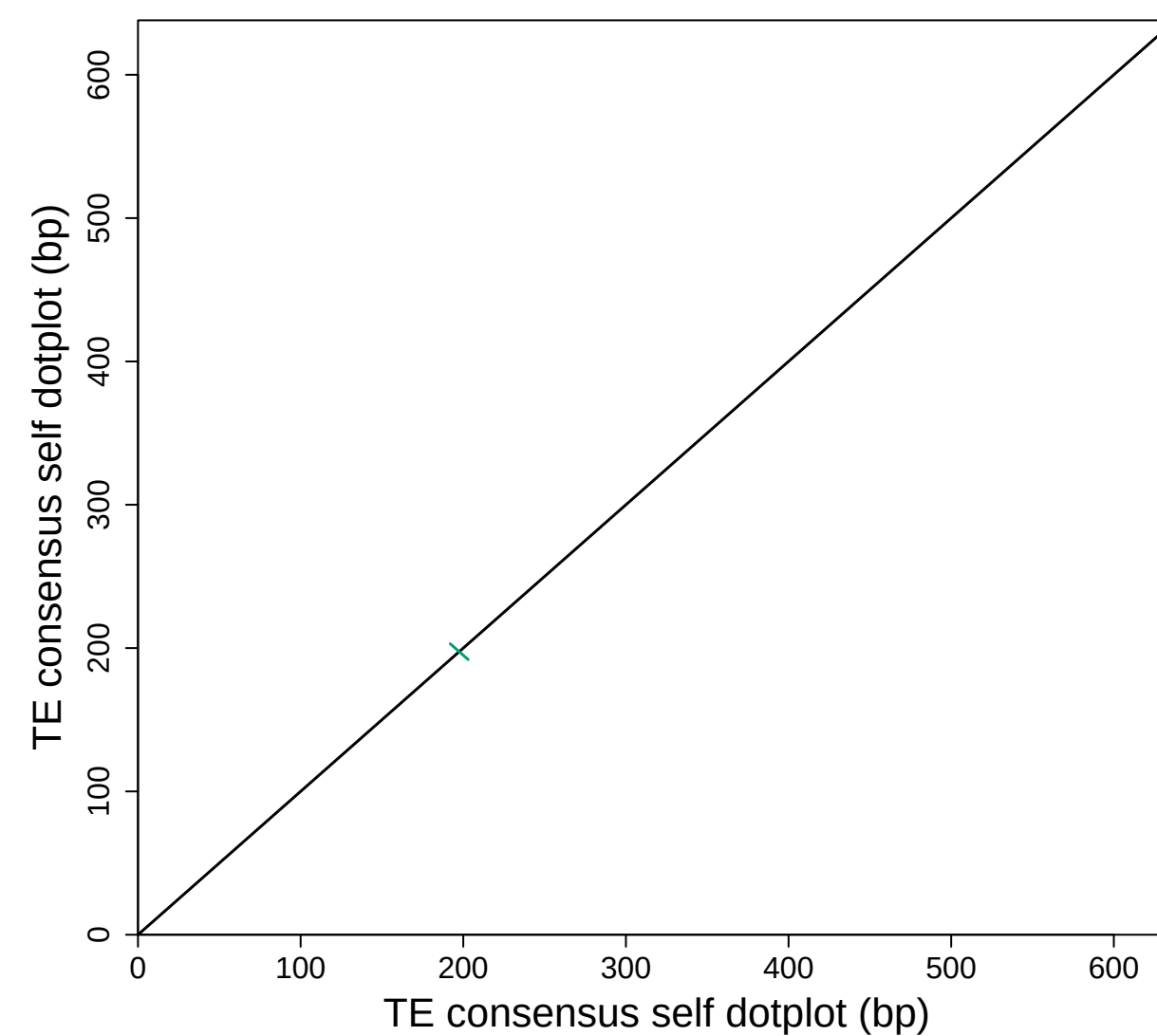
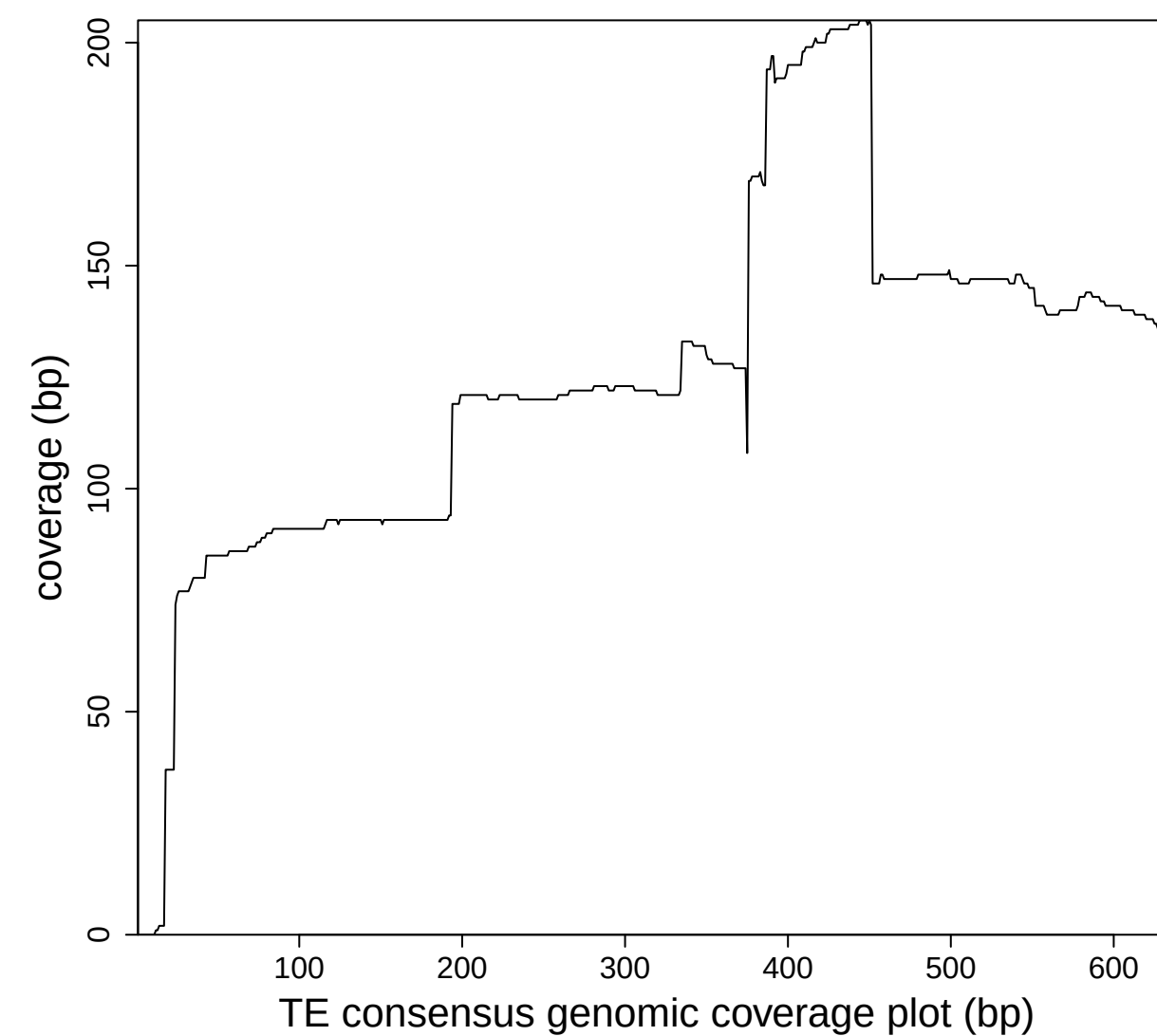
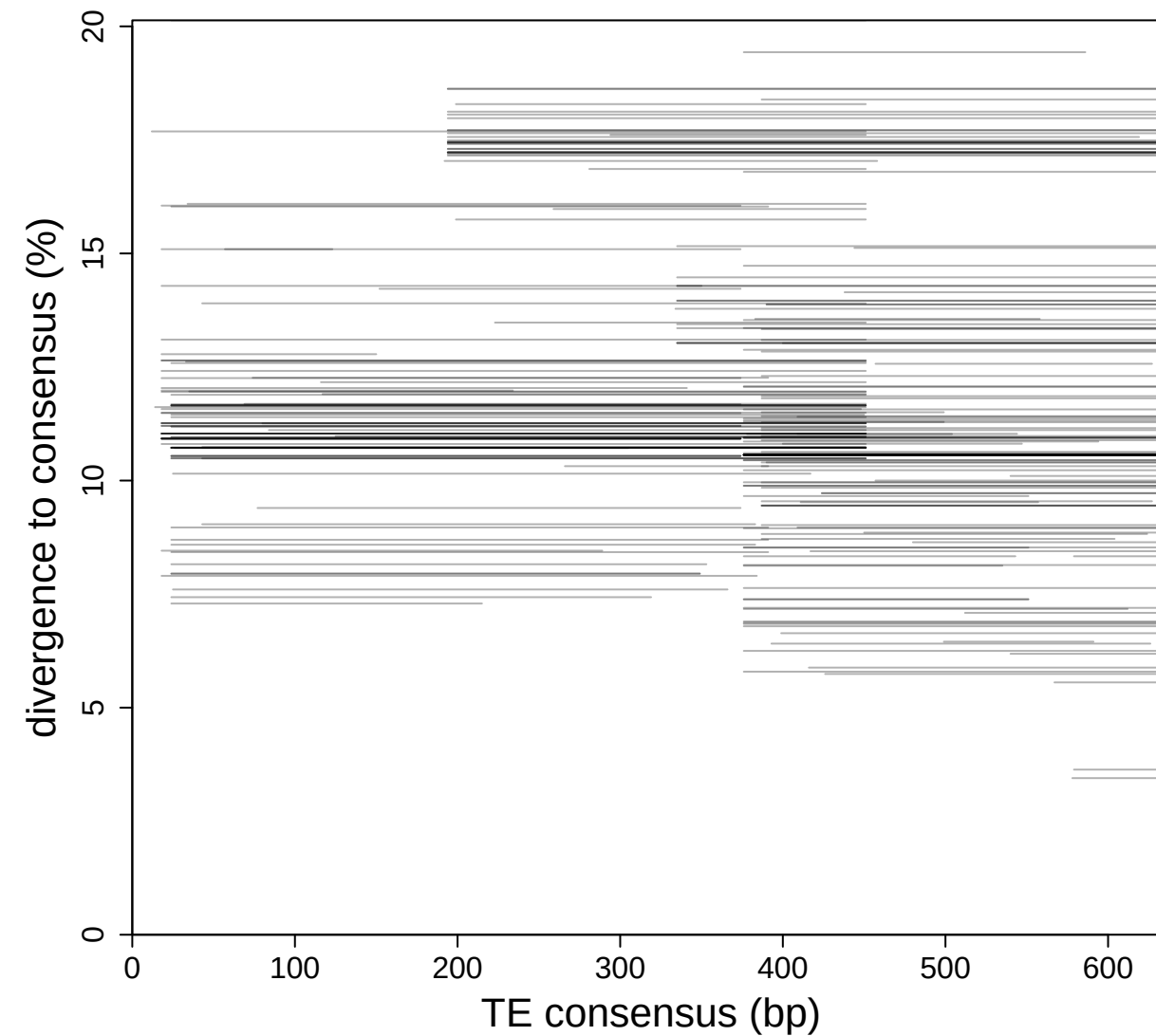
After TEtrimmer Extended plot Blue lines are boundaries



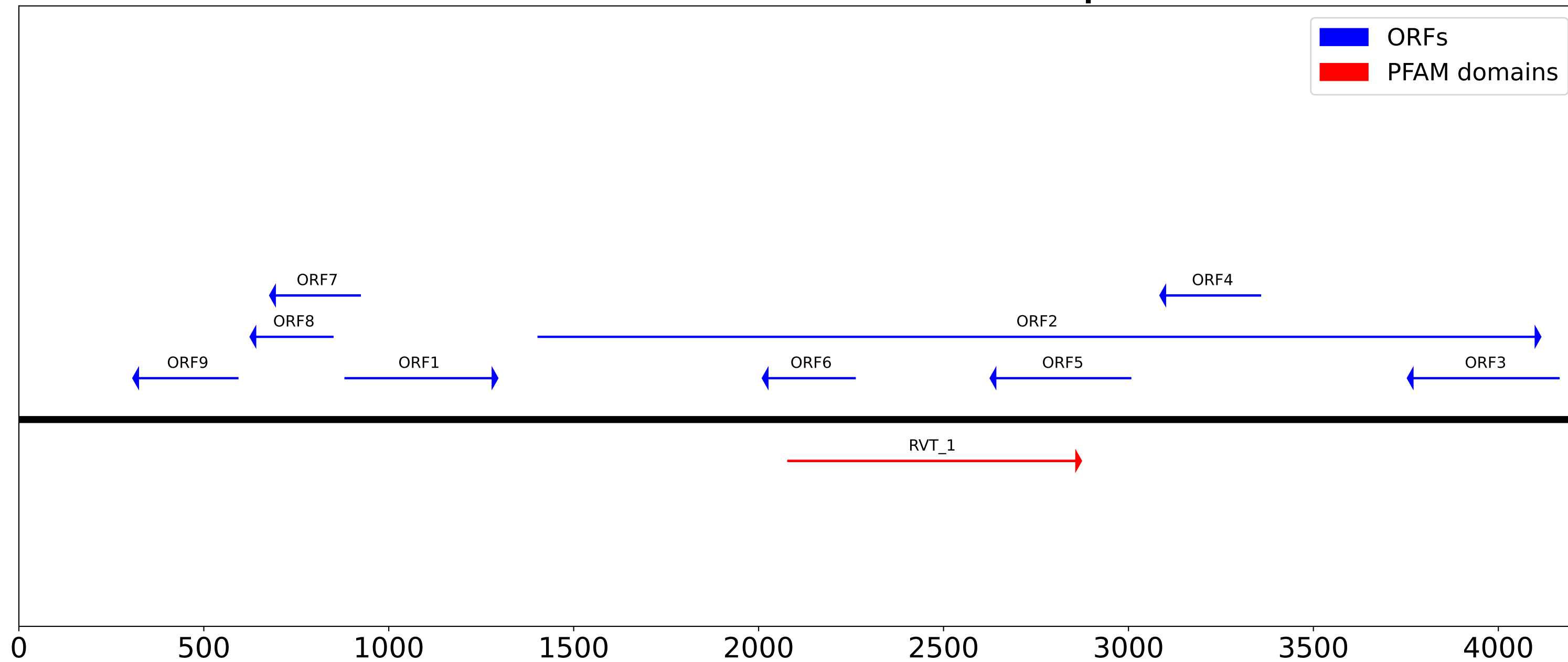
Before TEtrimmer 638 bp

TE: rnd_5_family_3018

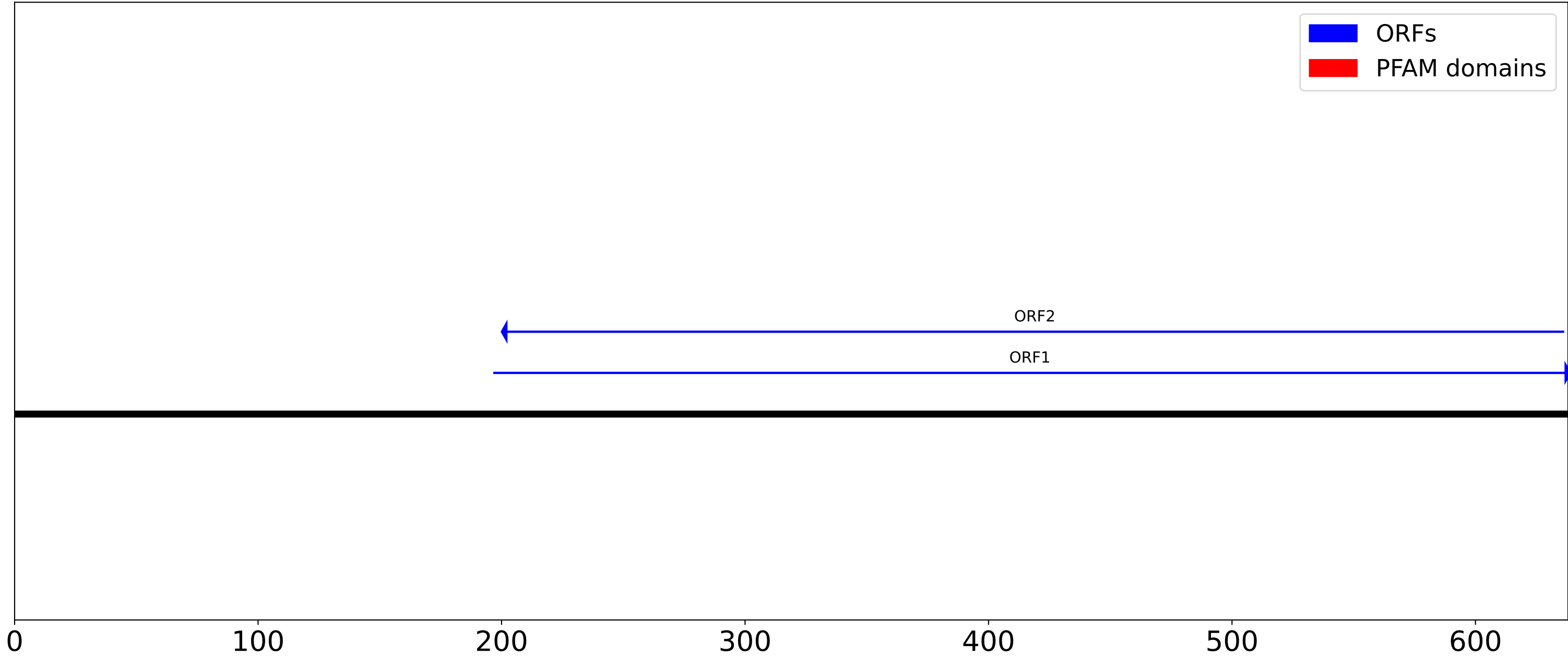
size: 638bp; fragments: 260; full length: 0 (≥ 574.2 bp)



After TEtrimmer ORF and PFAM domain plot



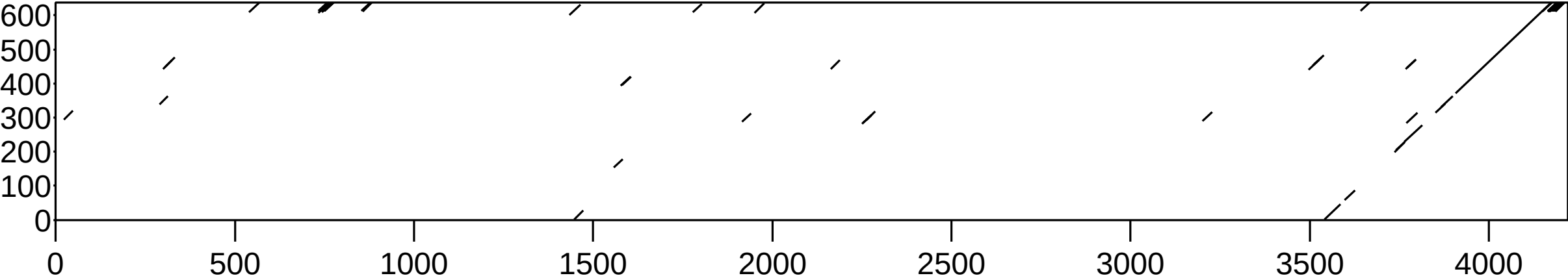
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

Isensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)