

Start crop Point



End crop Point

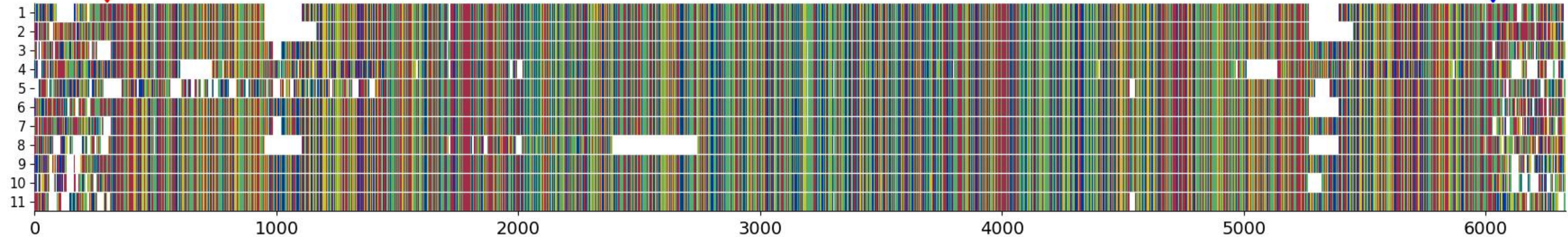


MSA length = 5732



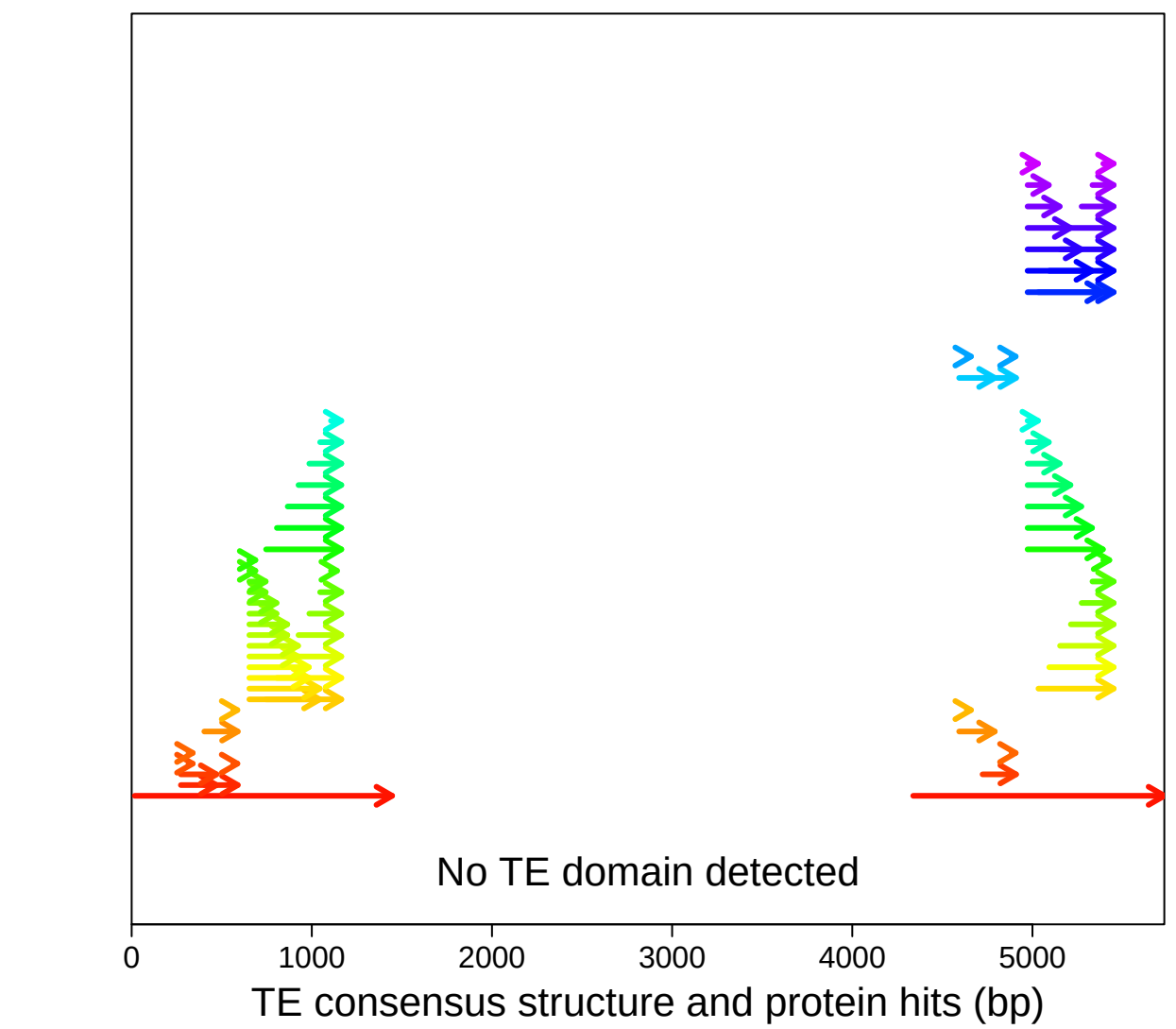
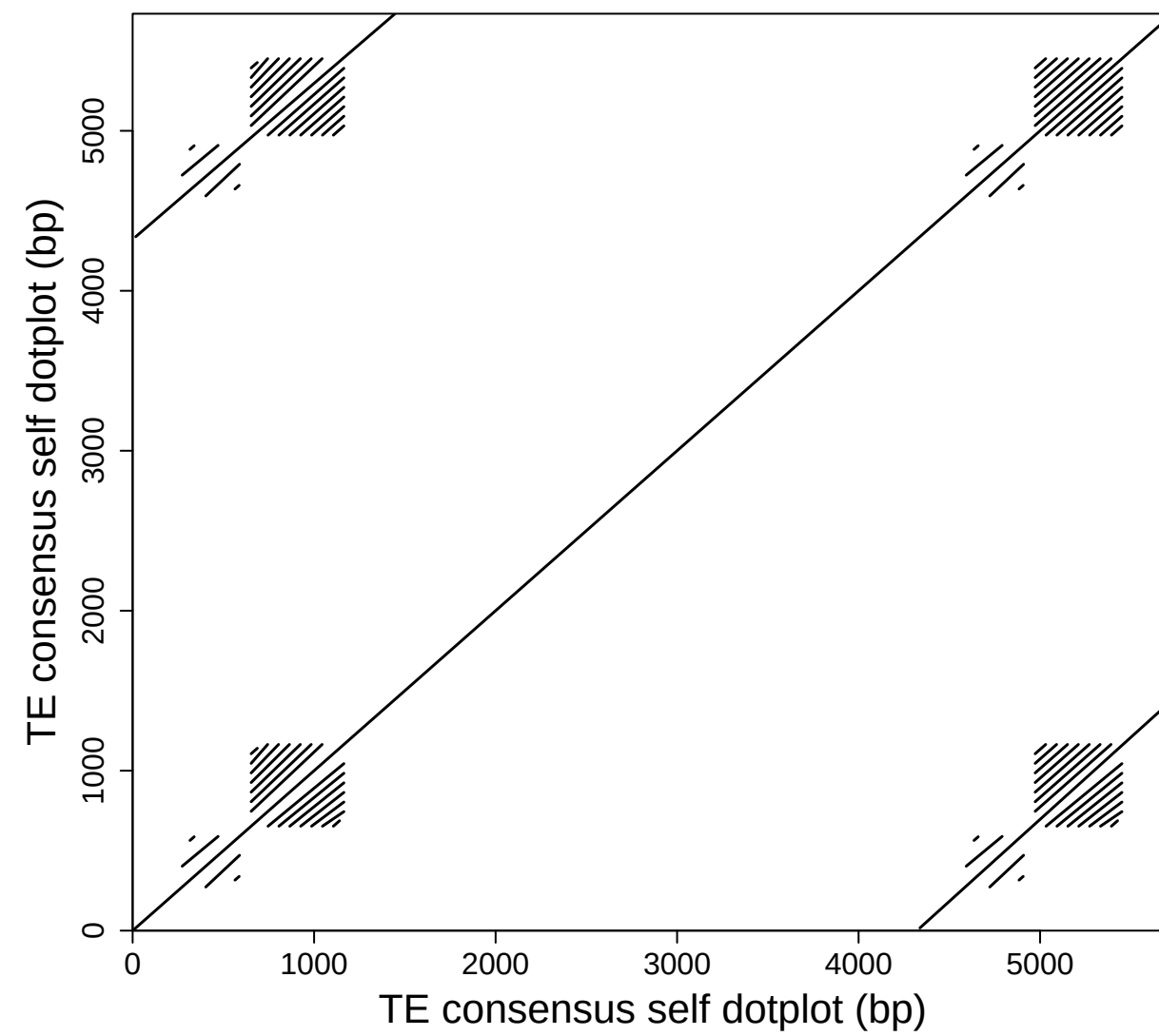
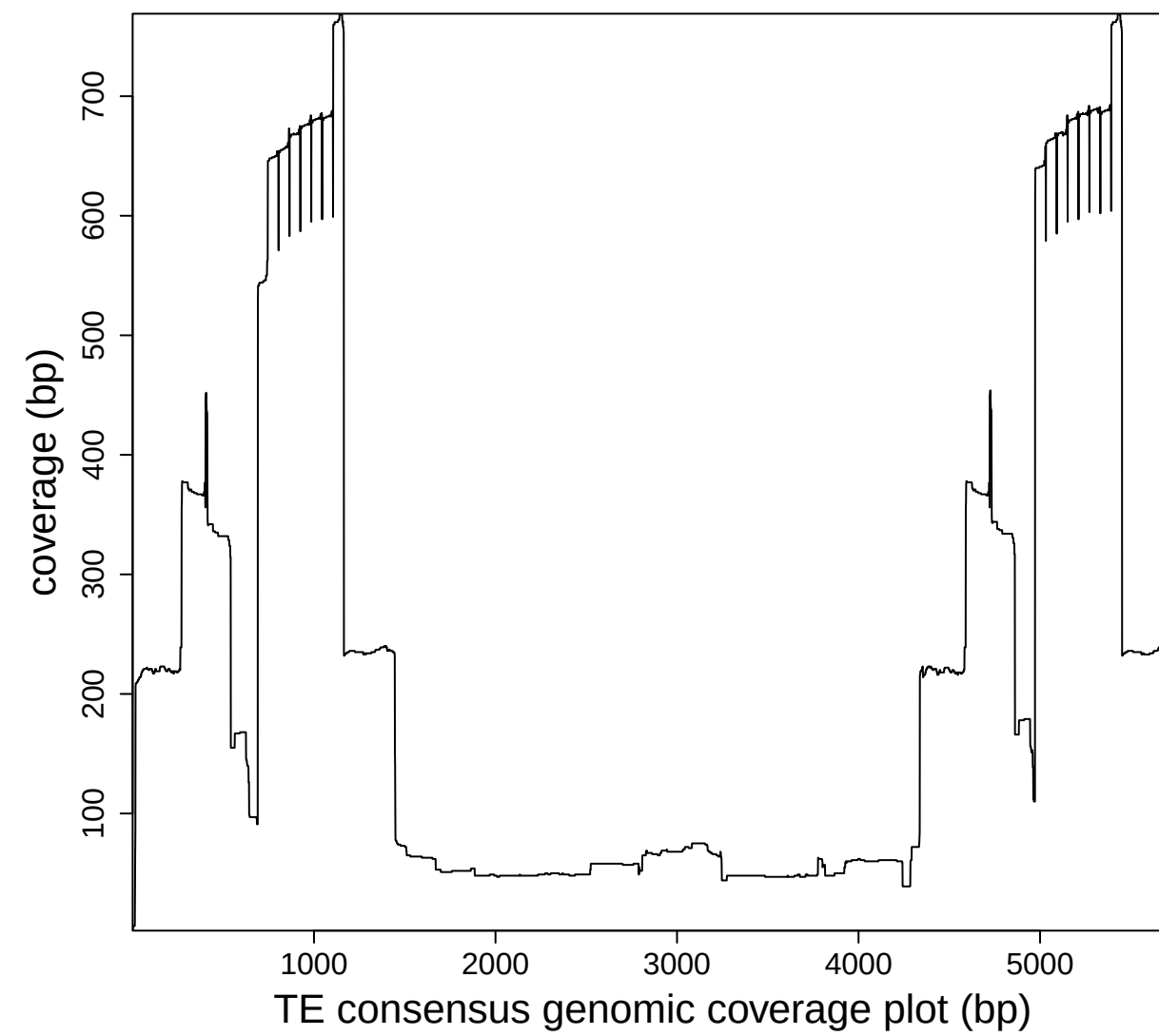
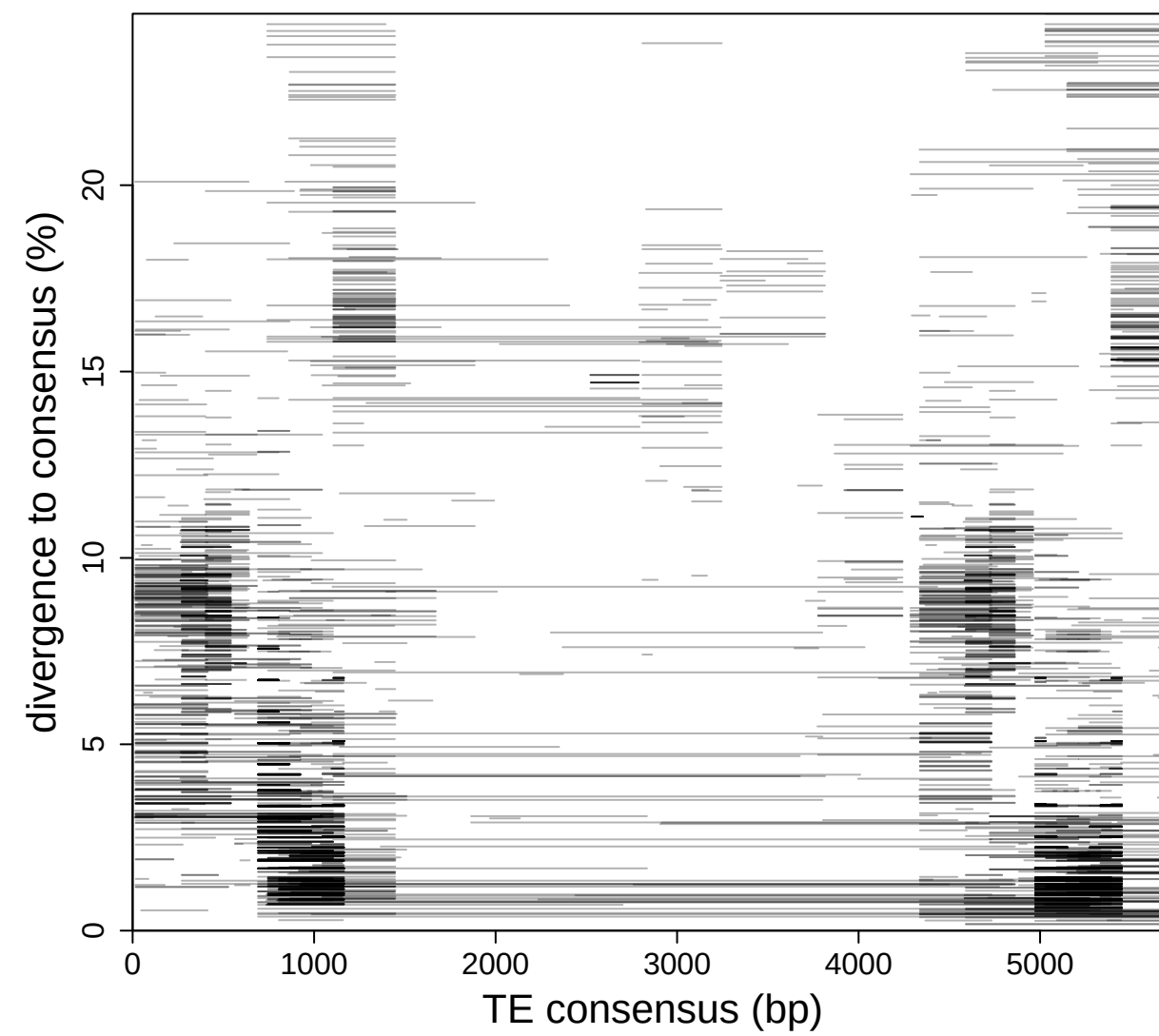
Start crop Point

End crop Point



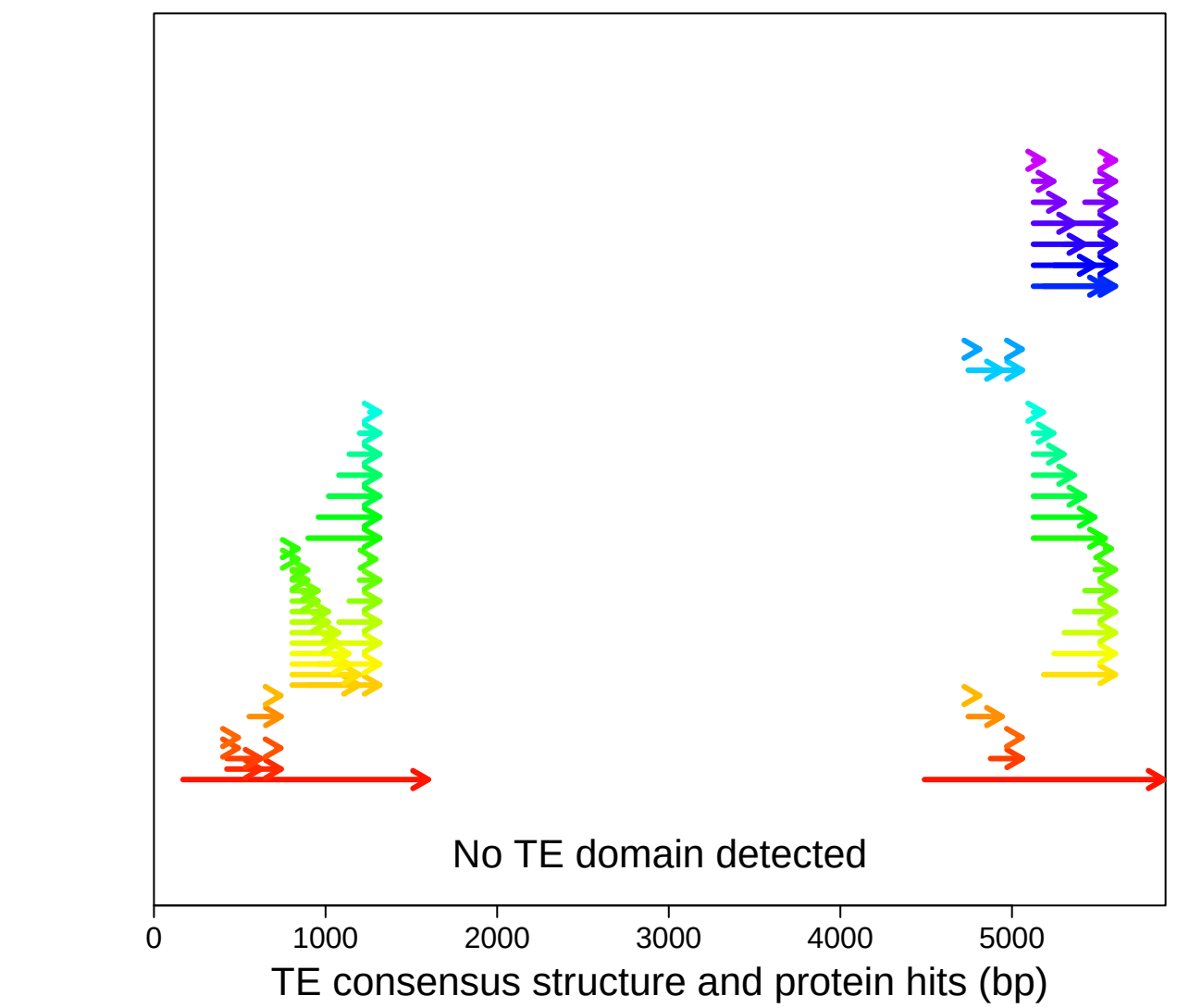
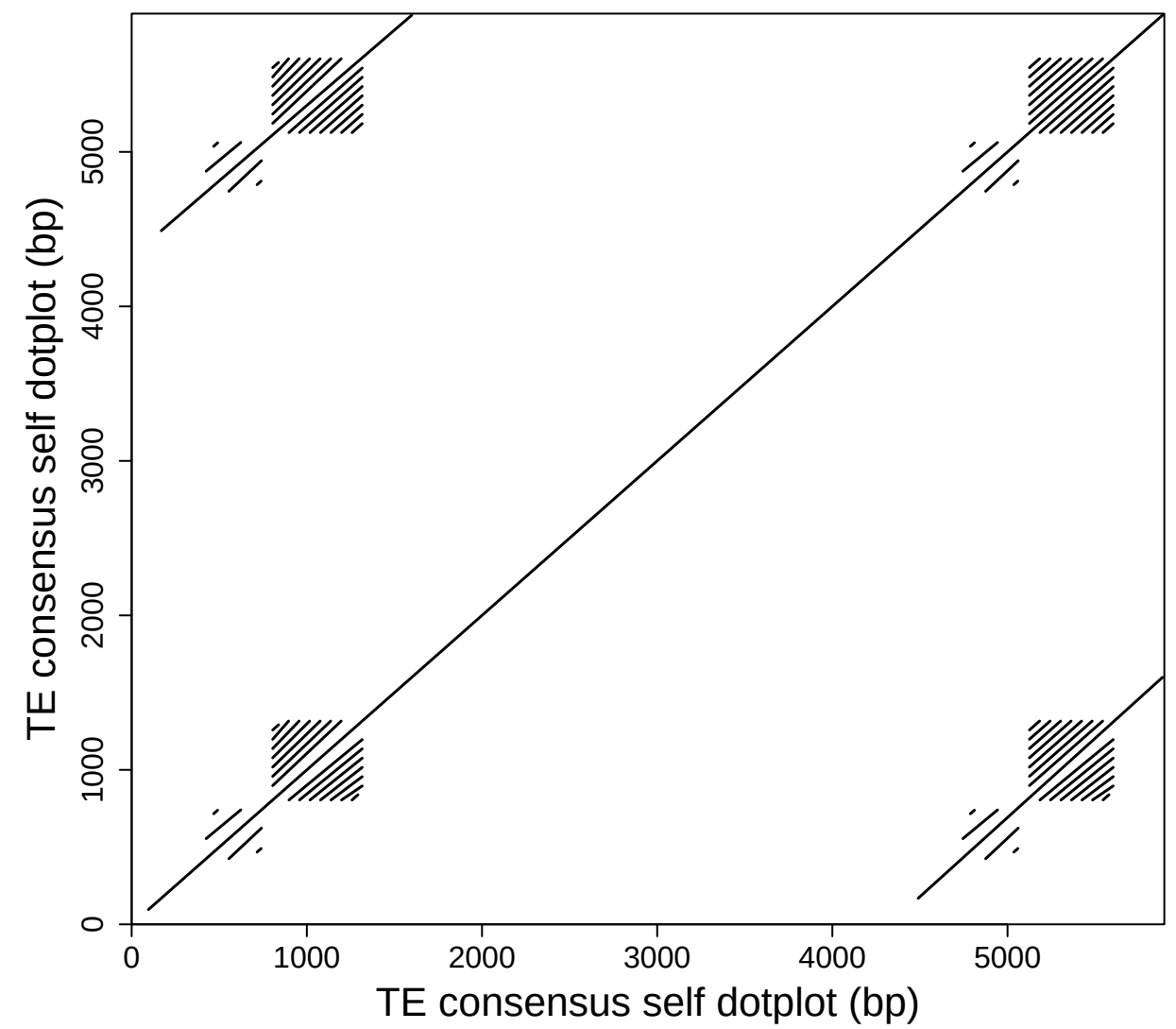
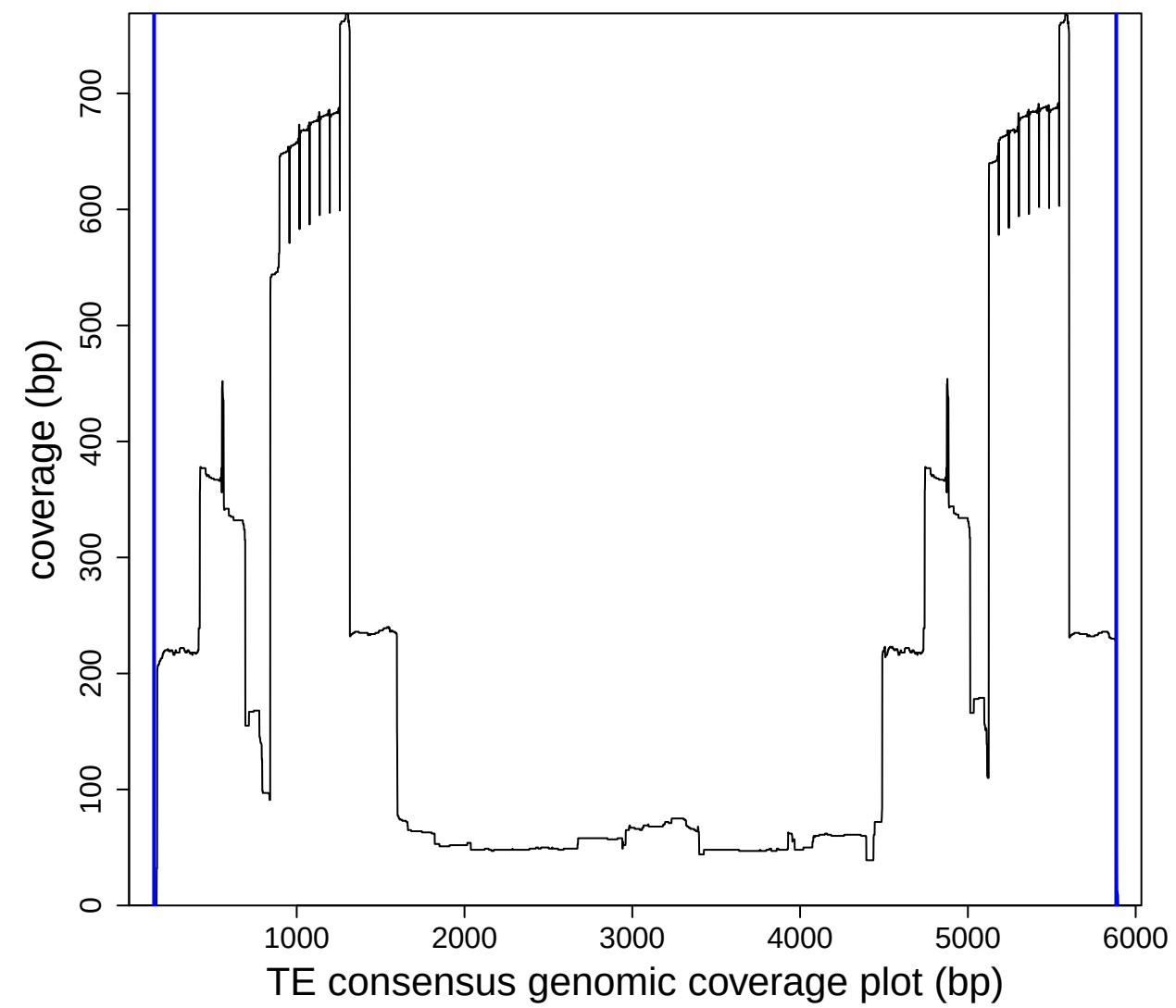
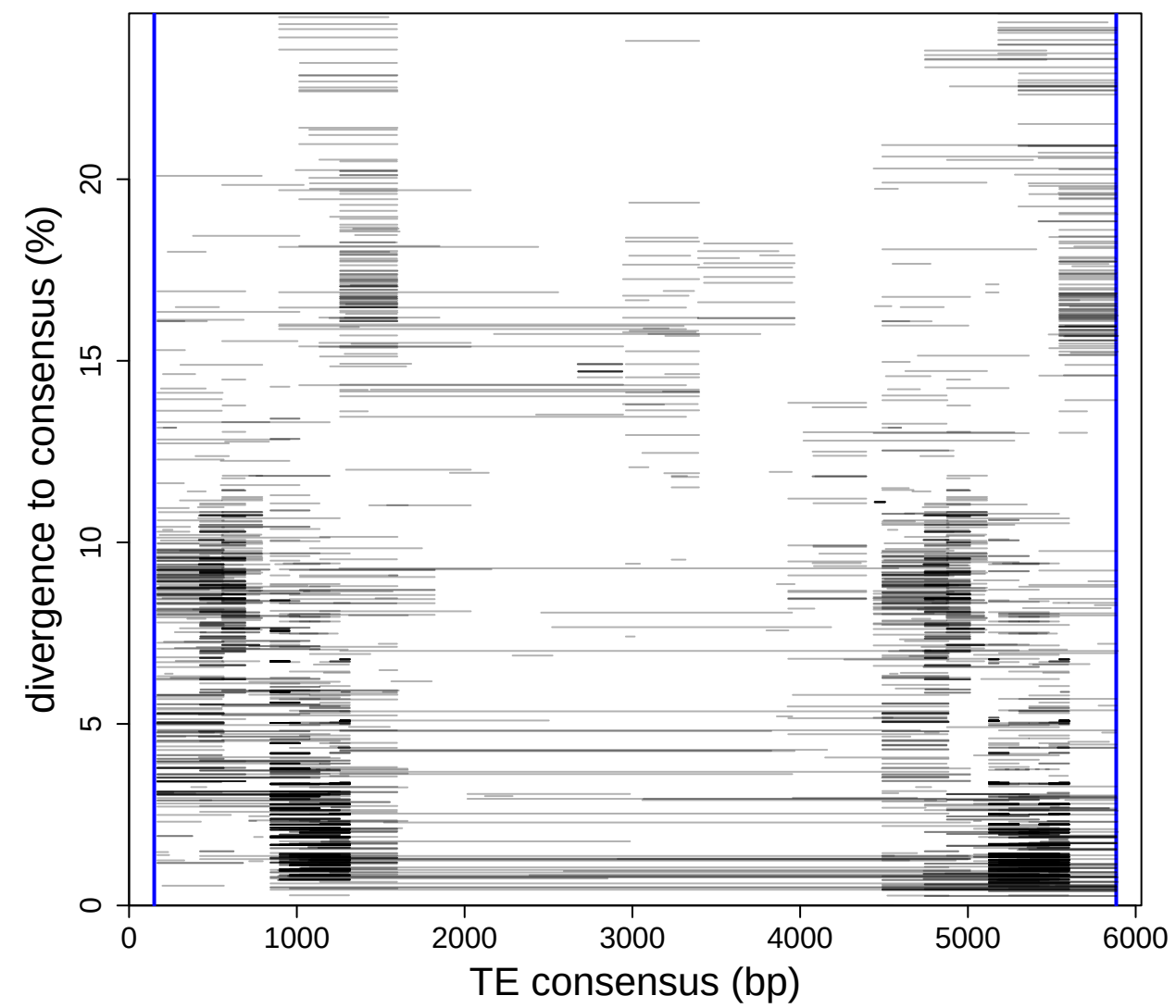
g_3.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln.fa ε
size: 5732bp; fragments: 3685; full length: 0 (>=5158.8bp)

After TEtrimmer 5732 bp



f.bed_g_3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 6034bp; fragments: 3679; full length: 0 (>=5430.6bp)

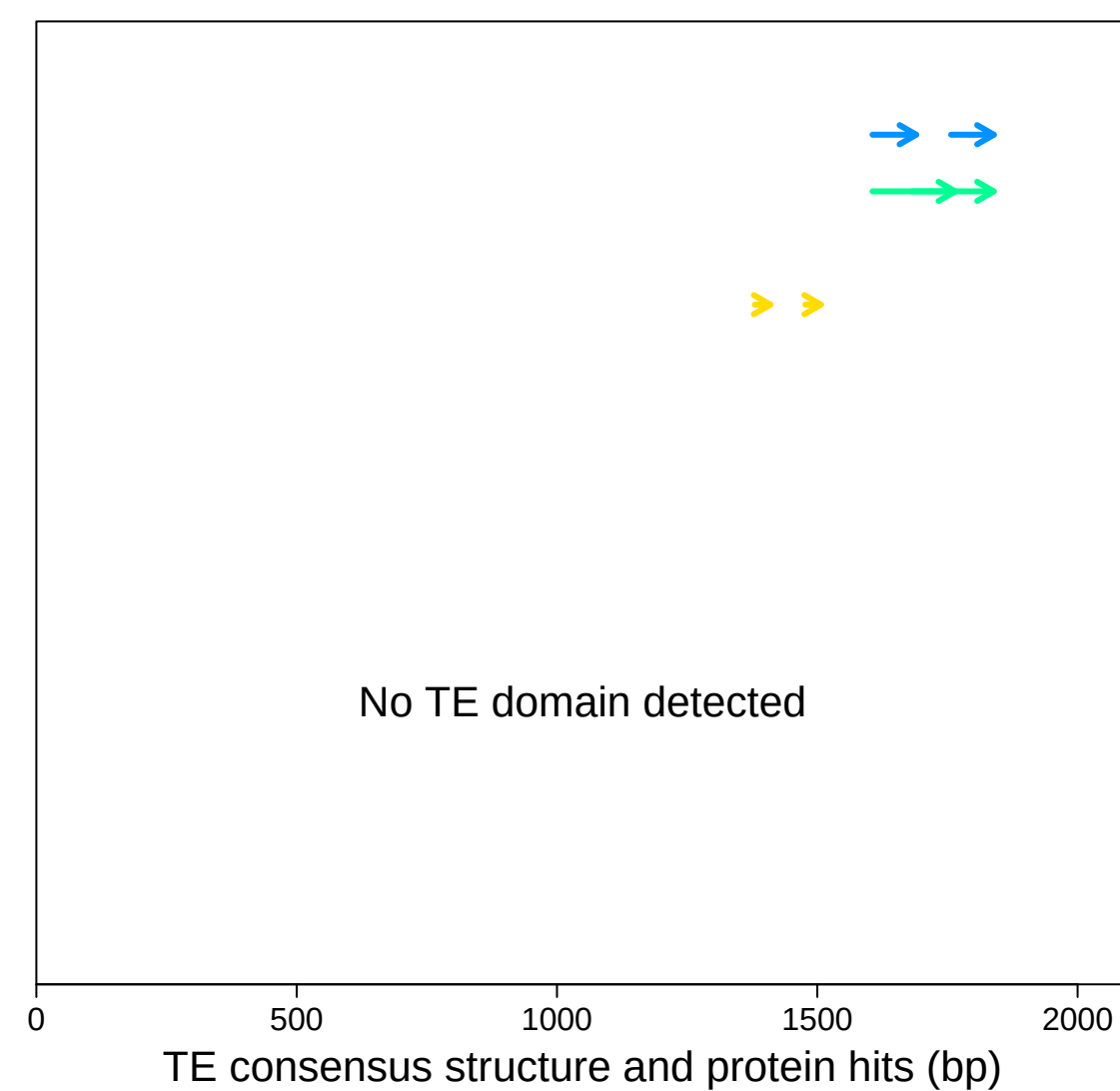
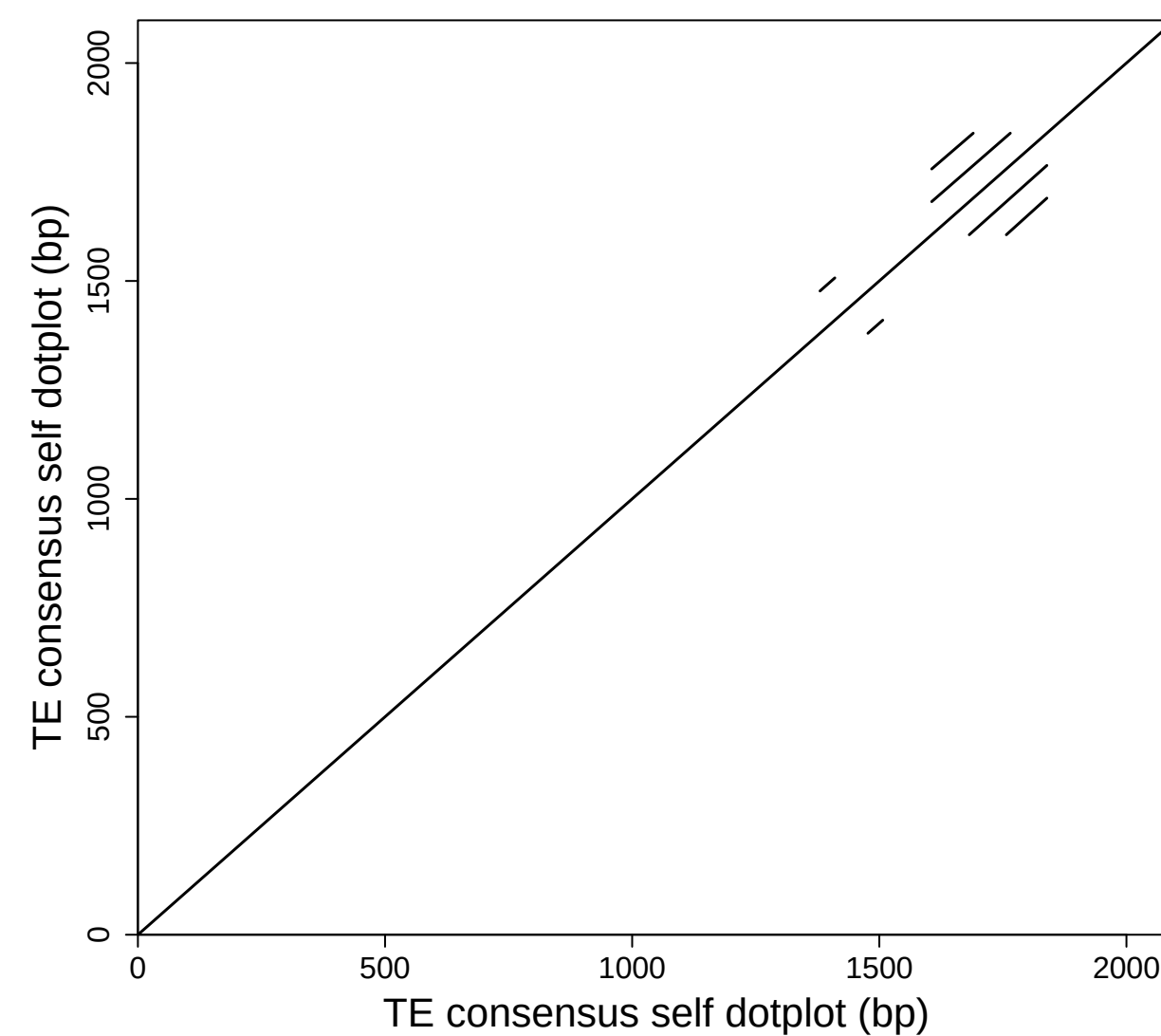
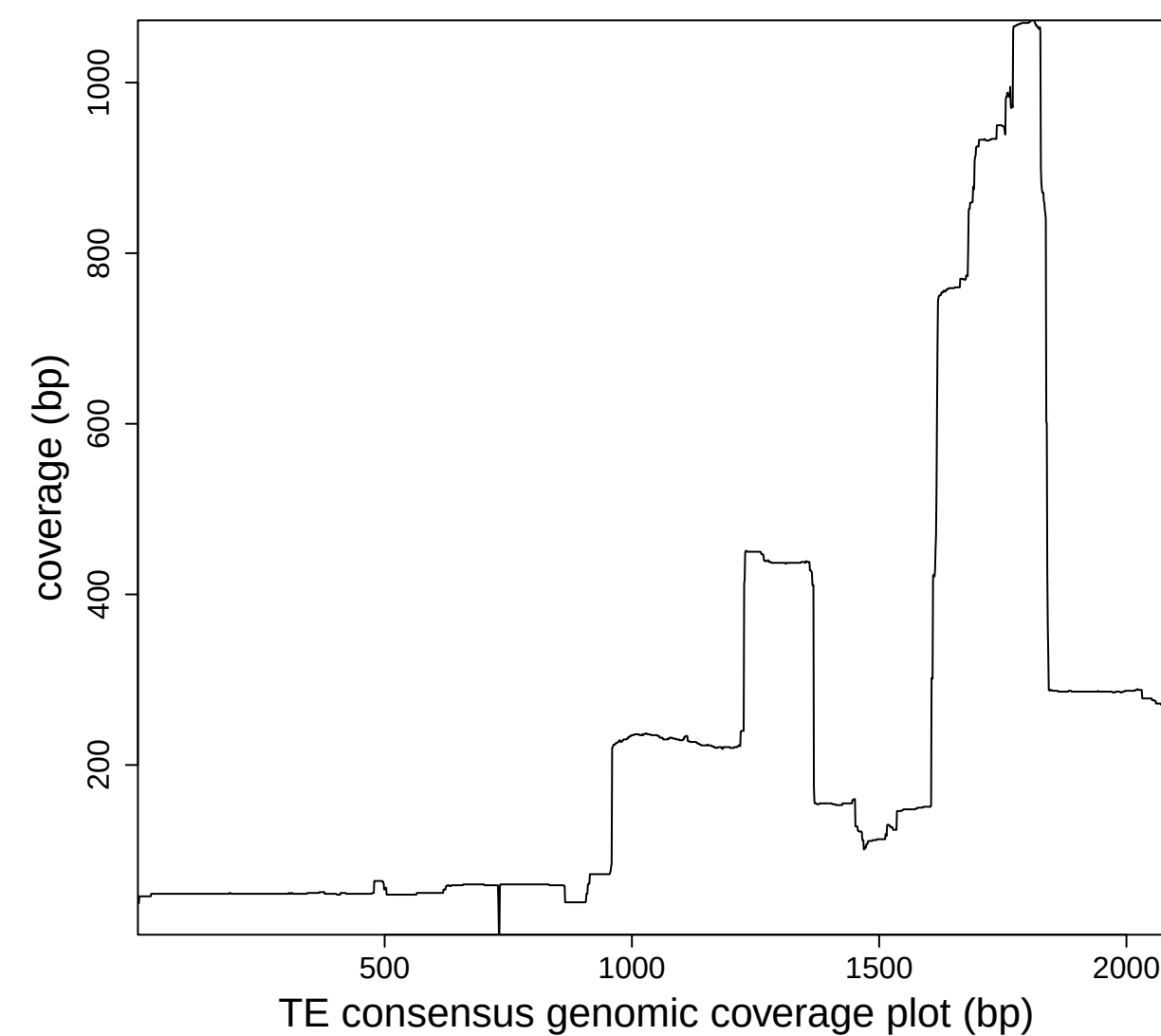
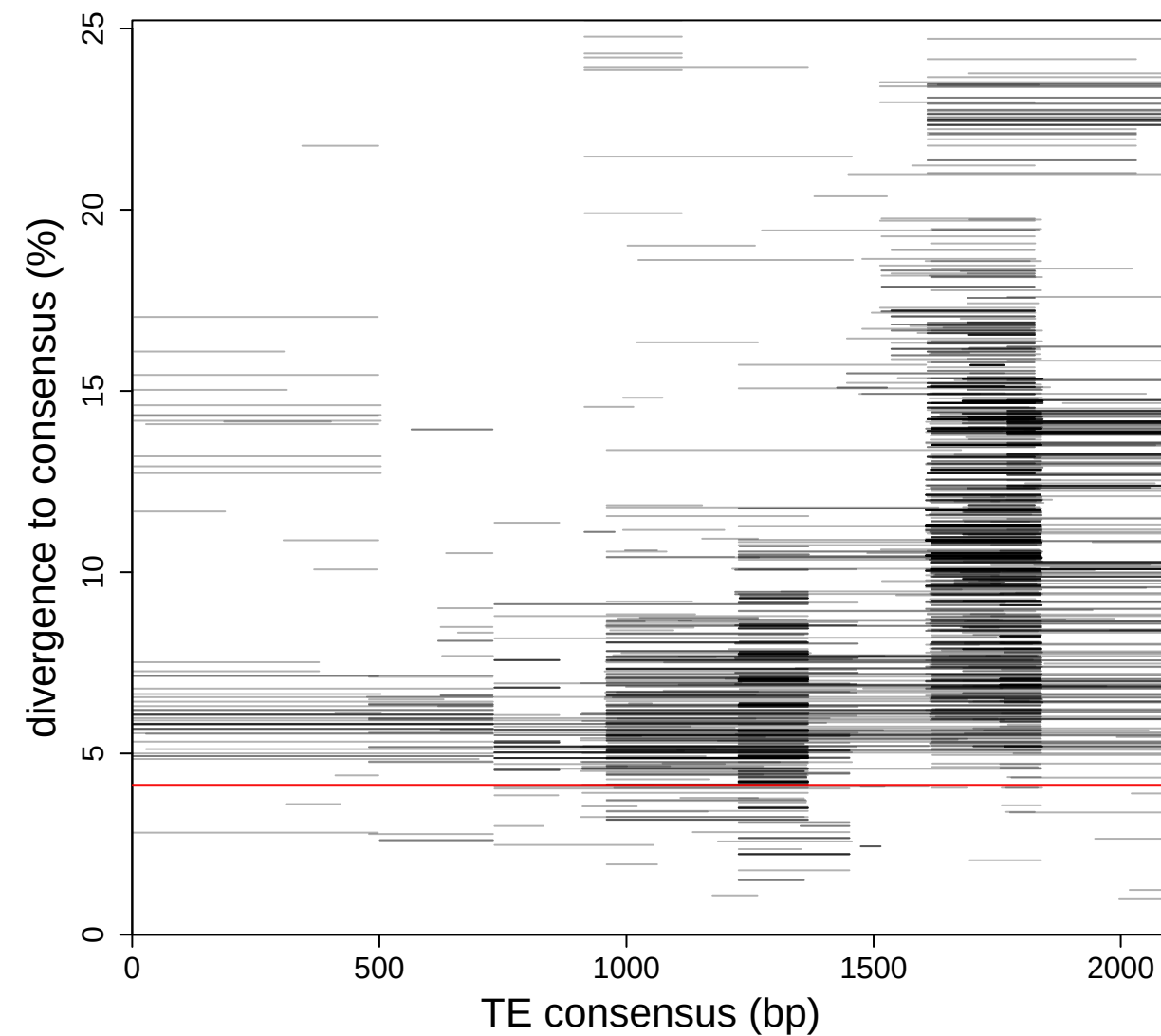
After TEtrimmer Extended plot Blue lines are boundaries



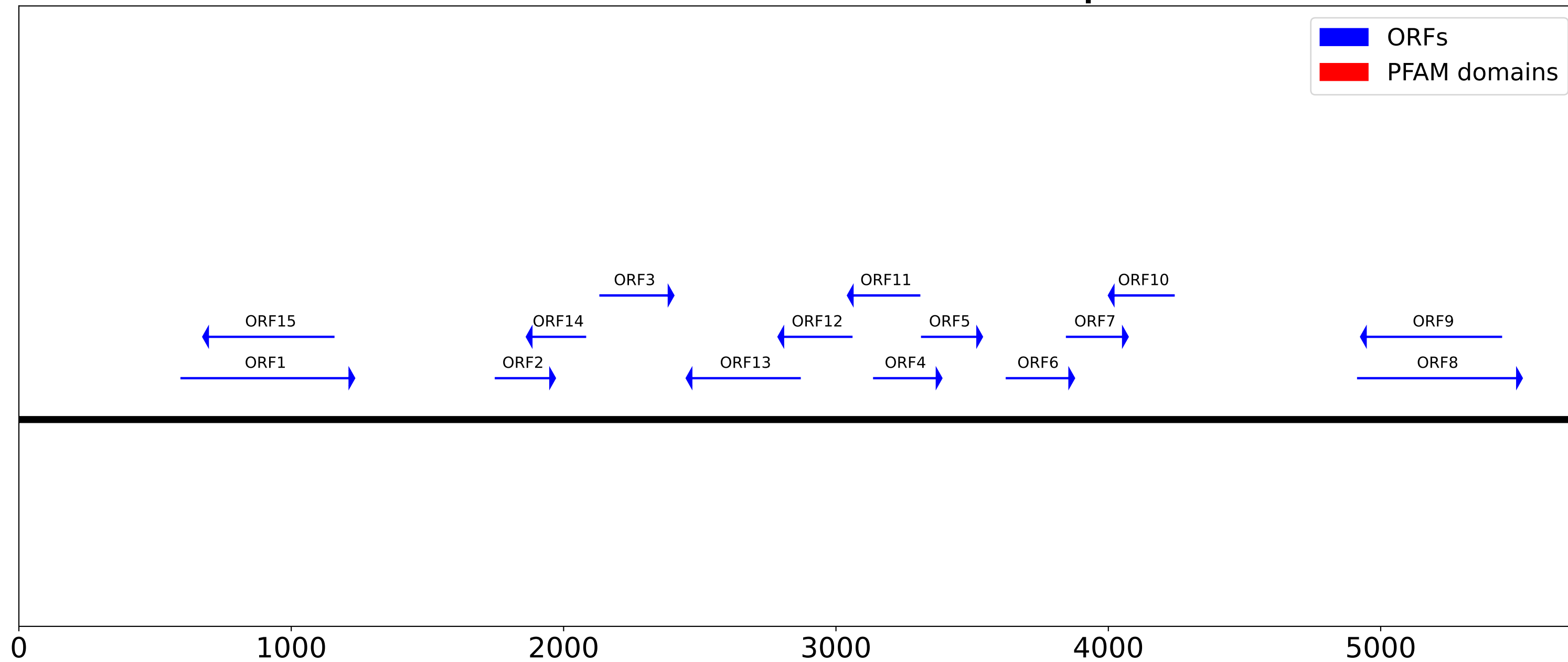
Before TEtrimmer 2098 bp

TE: rnd_5_family_5679

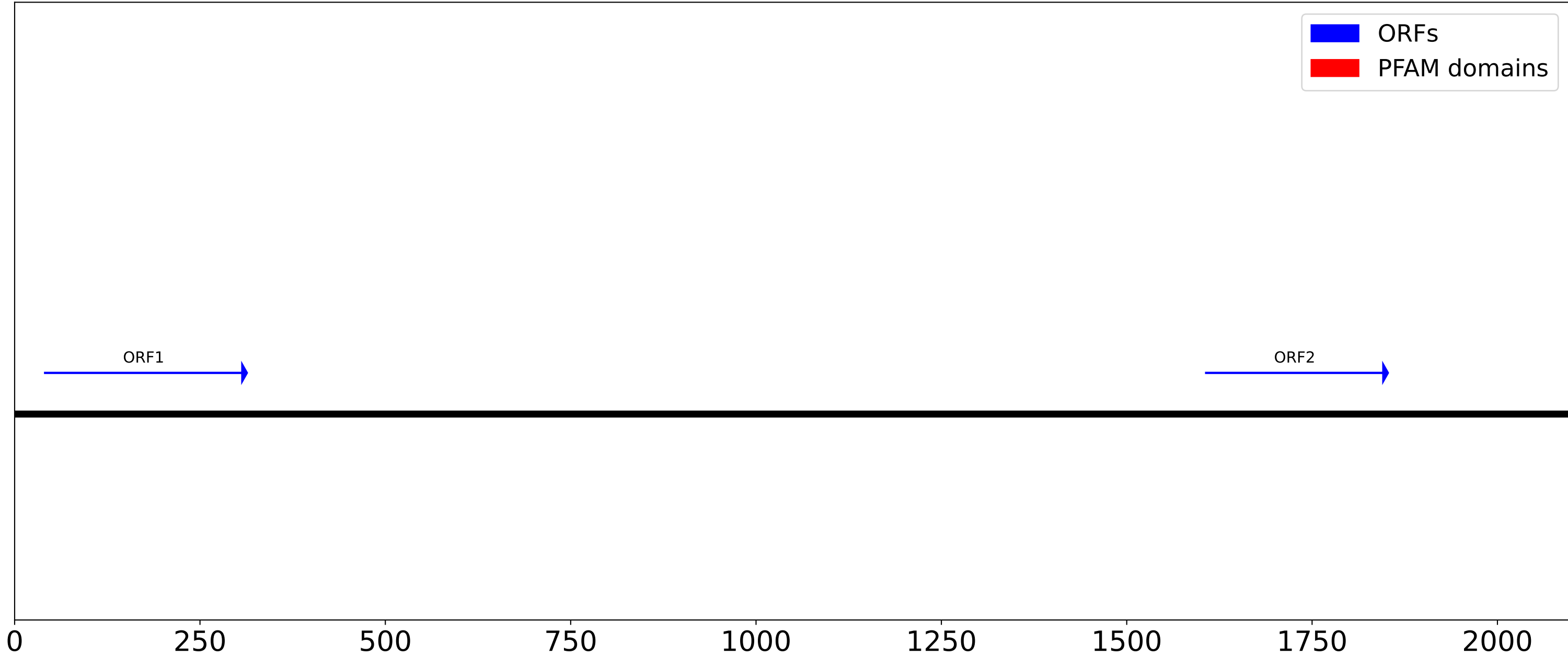
size: 2098bp; fragments: 1664; full length: 1 (≥ 1888.2 bp)



After TEtrimmer ORF and PFAM domain plot



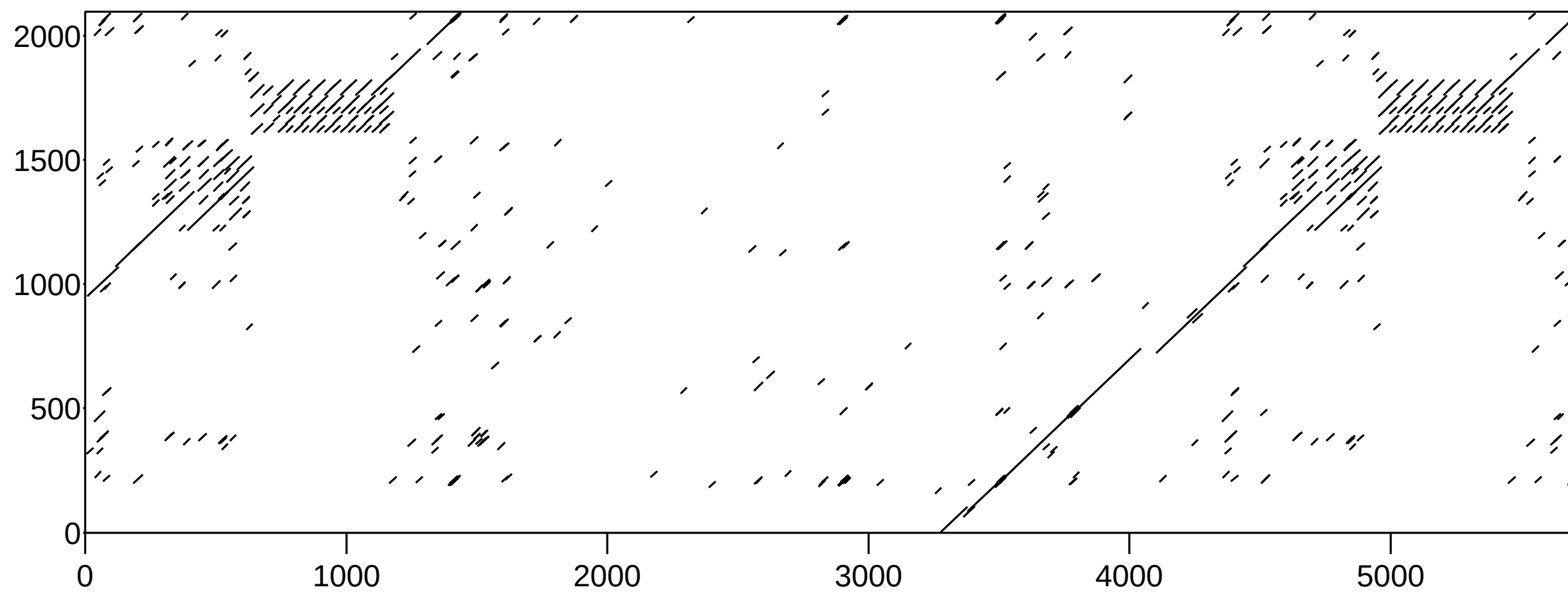
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)