

MSA length = 7403

Start crop Point

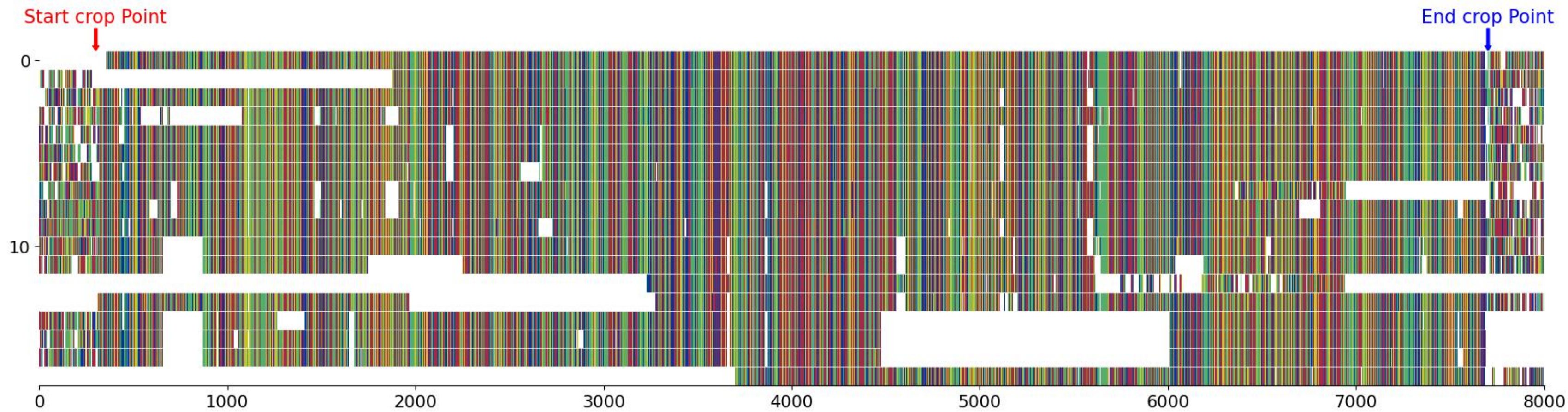


End crop Point



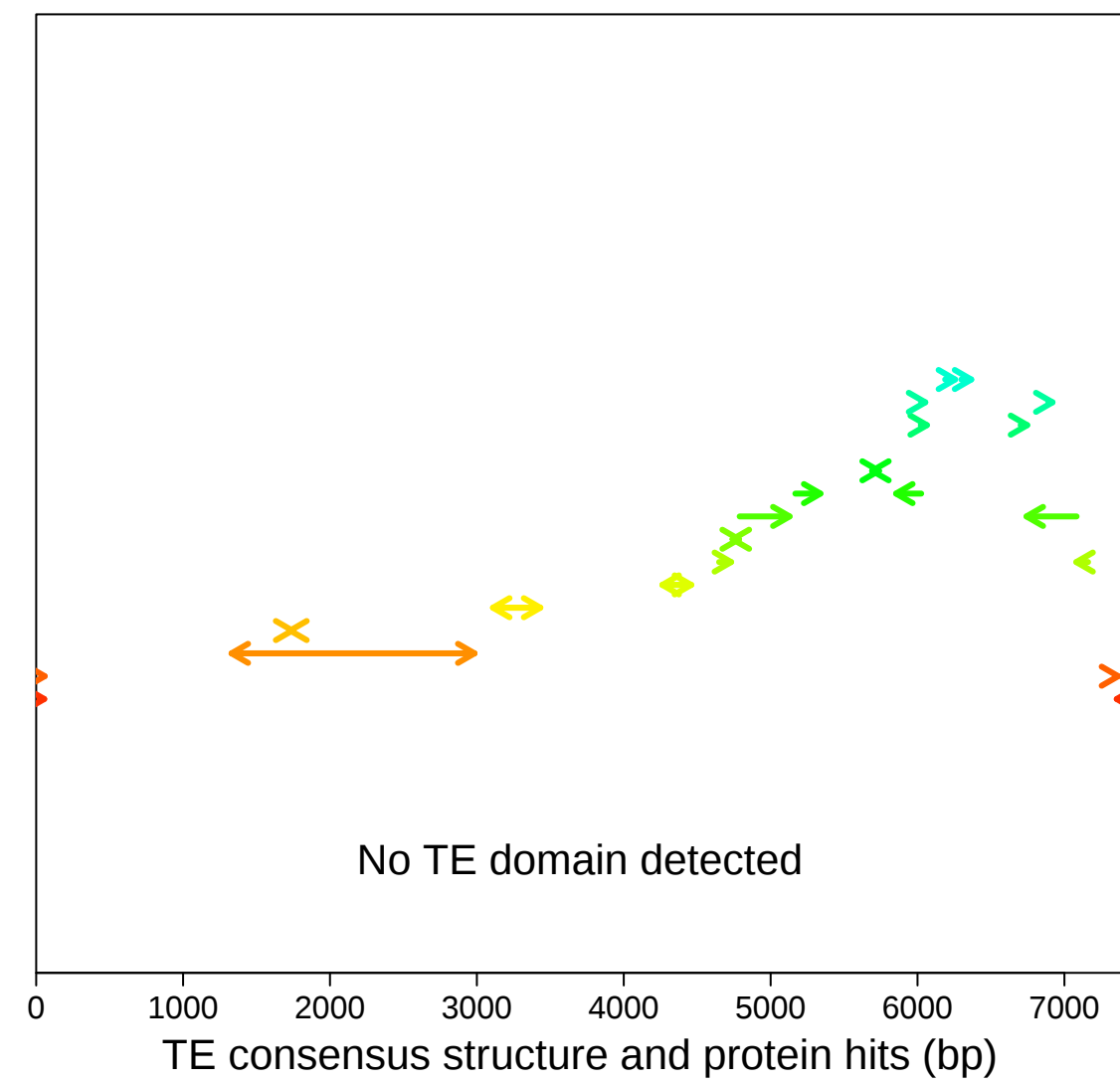
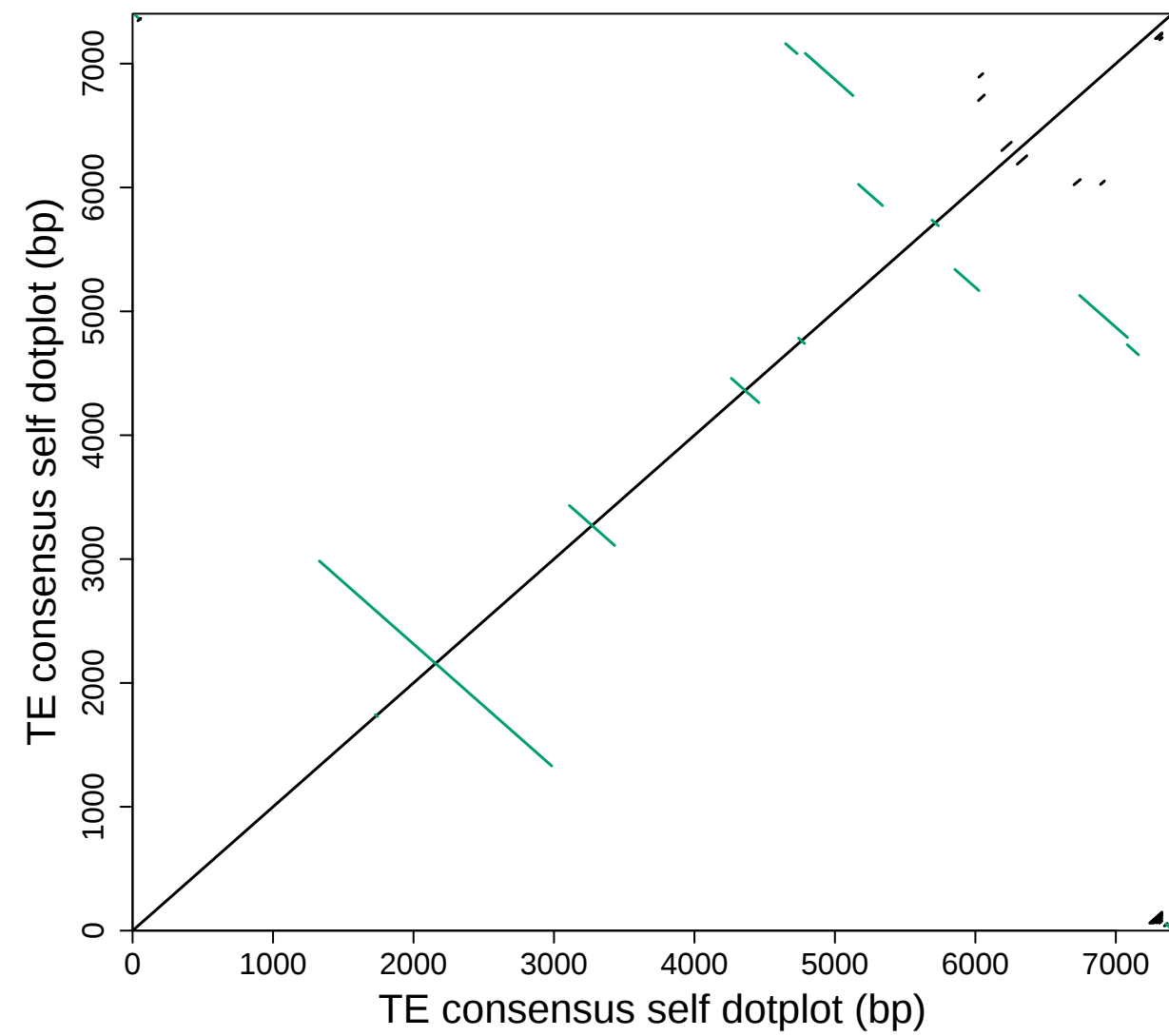
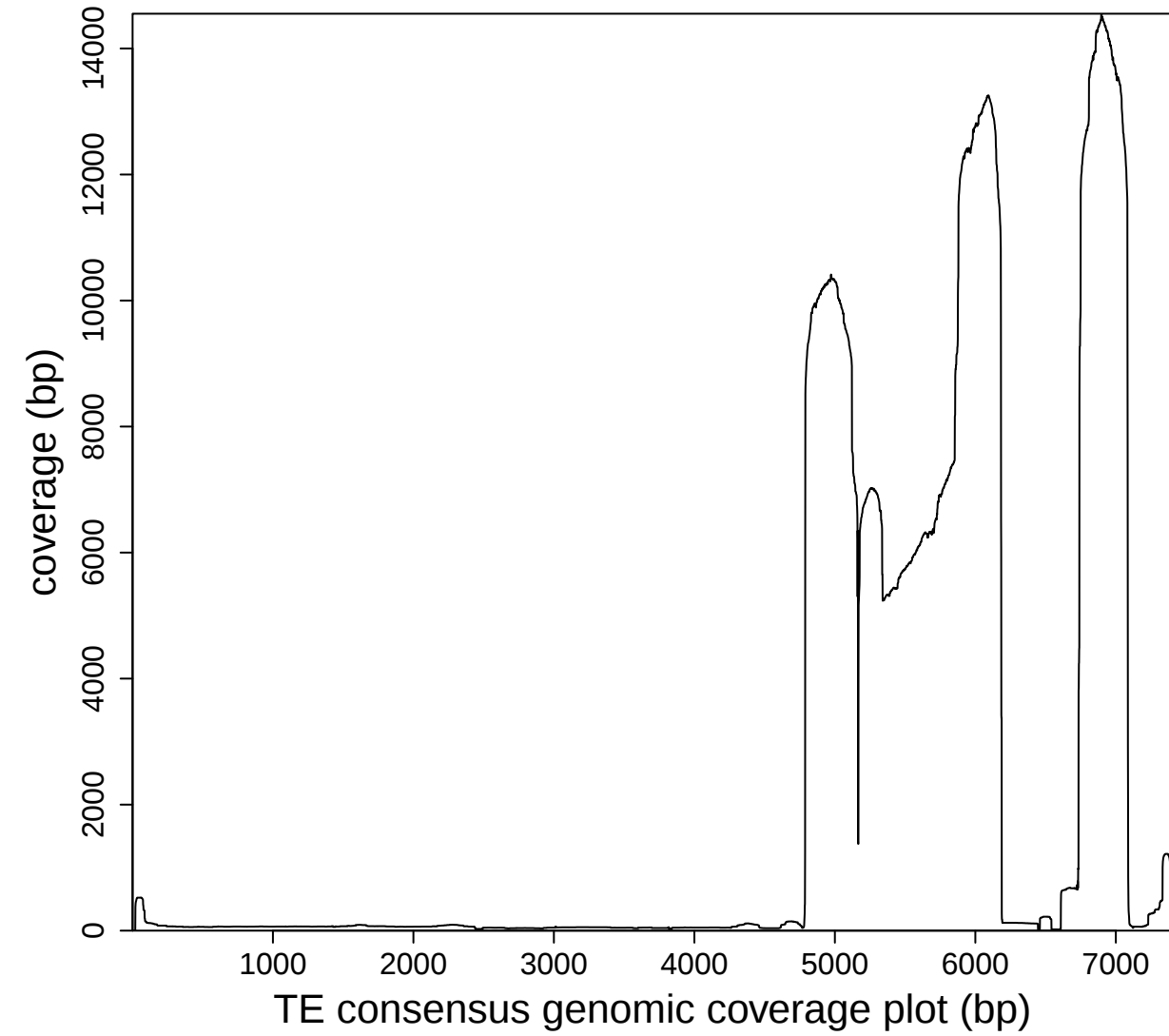
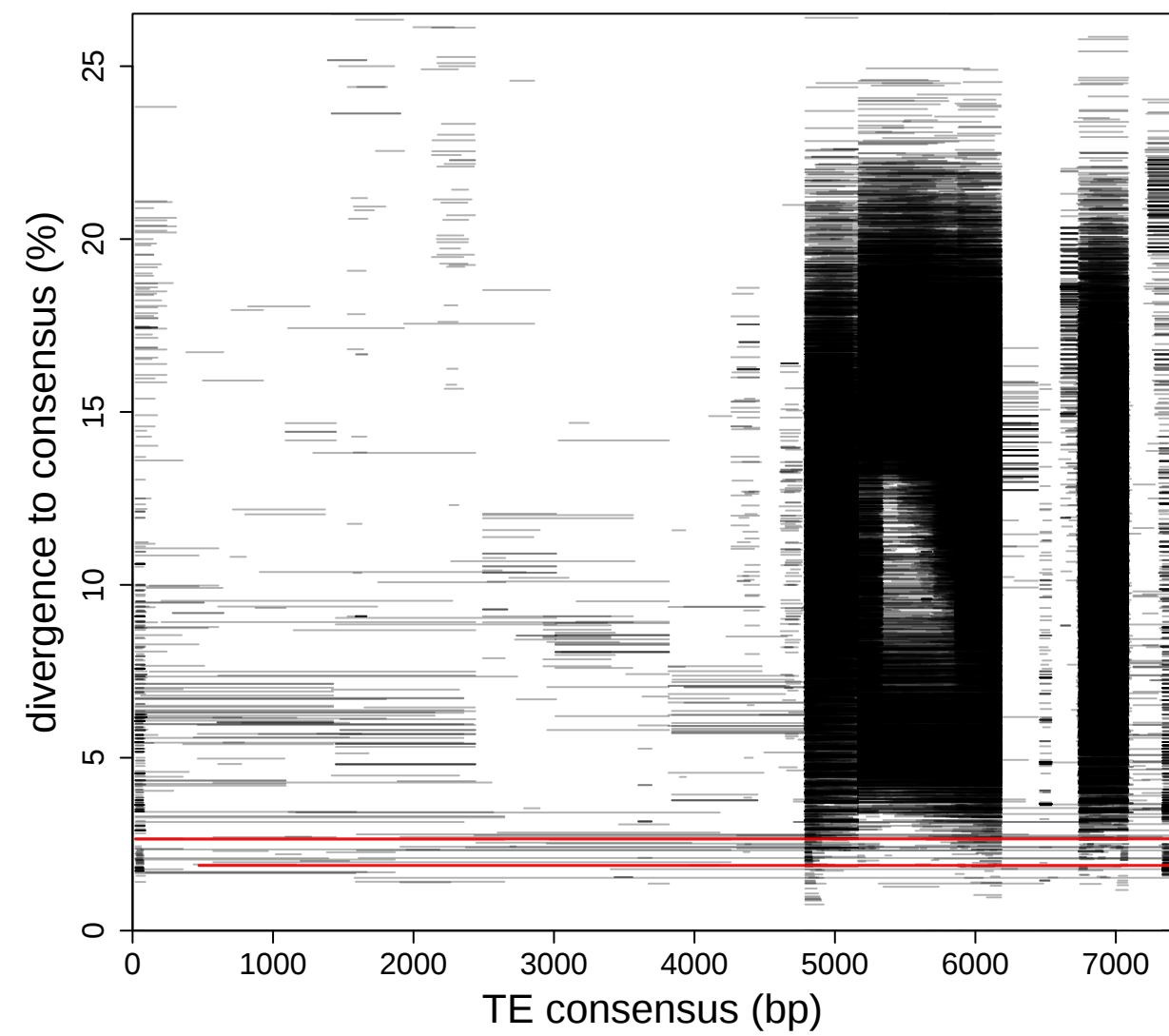
Start crop Point

End crop Point



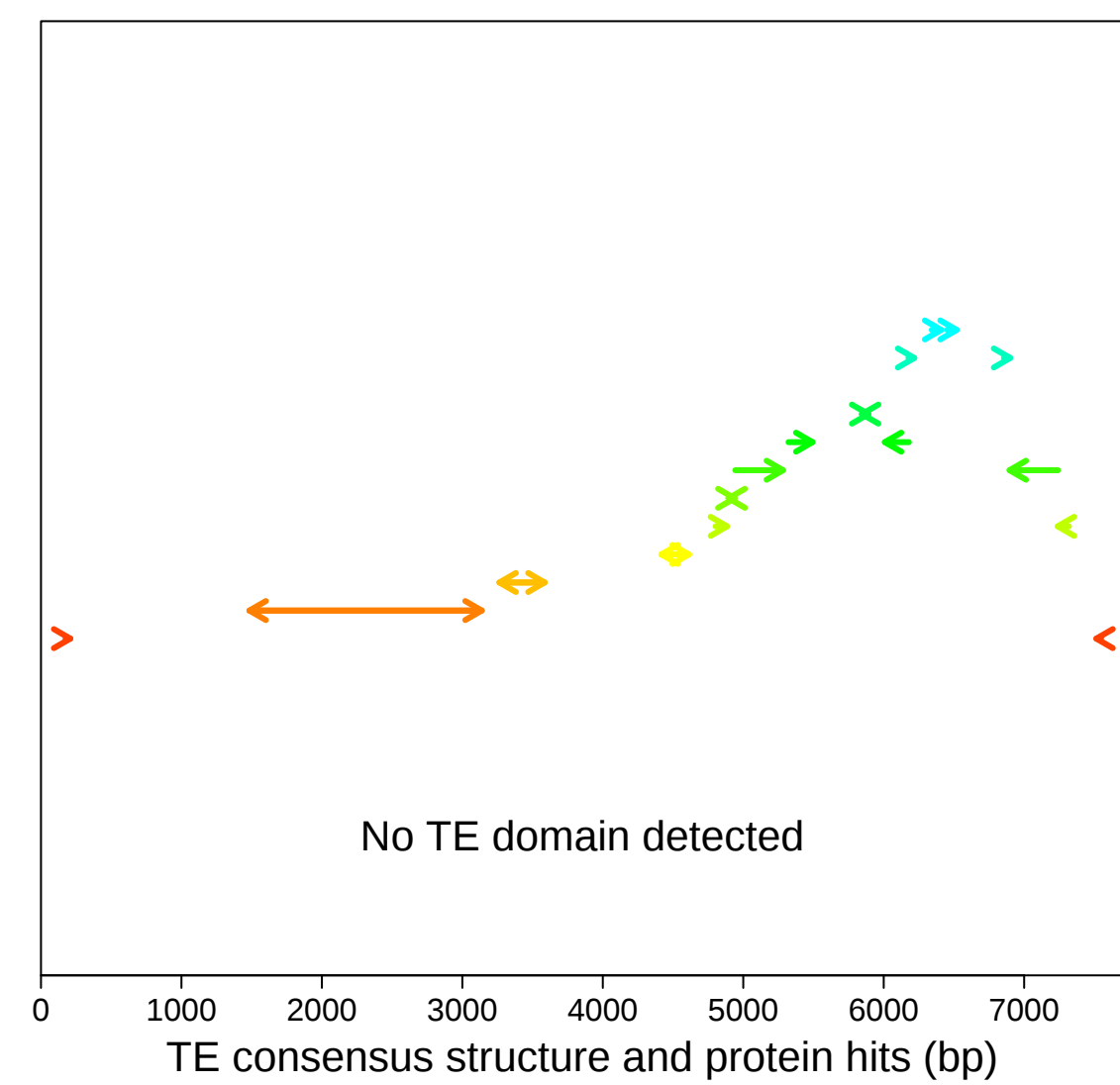
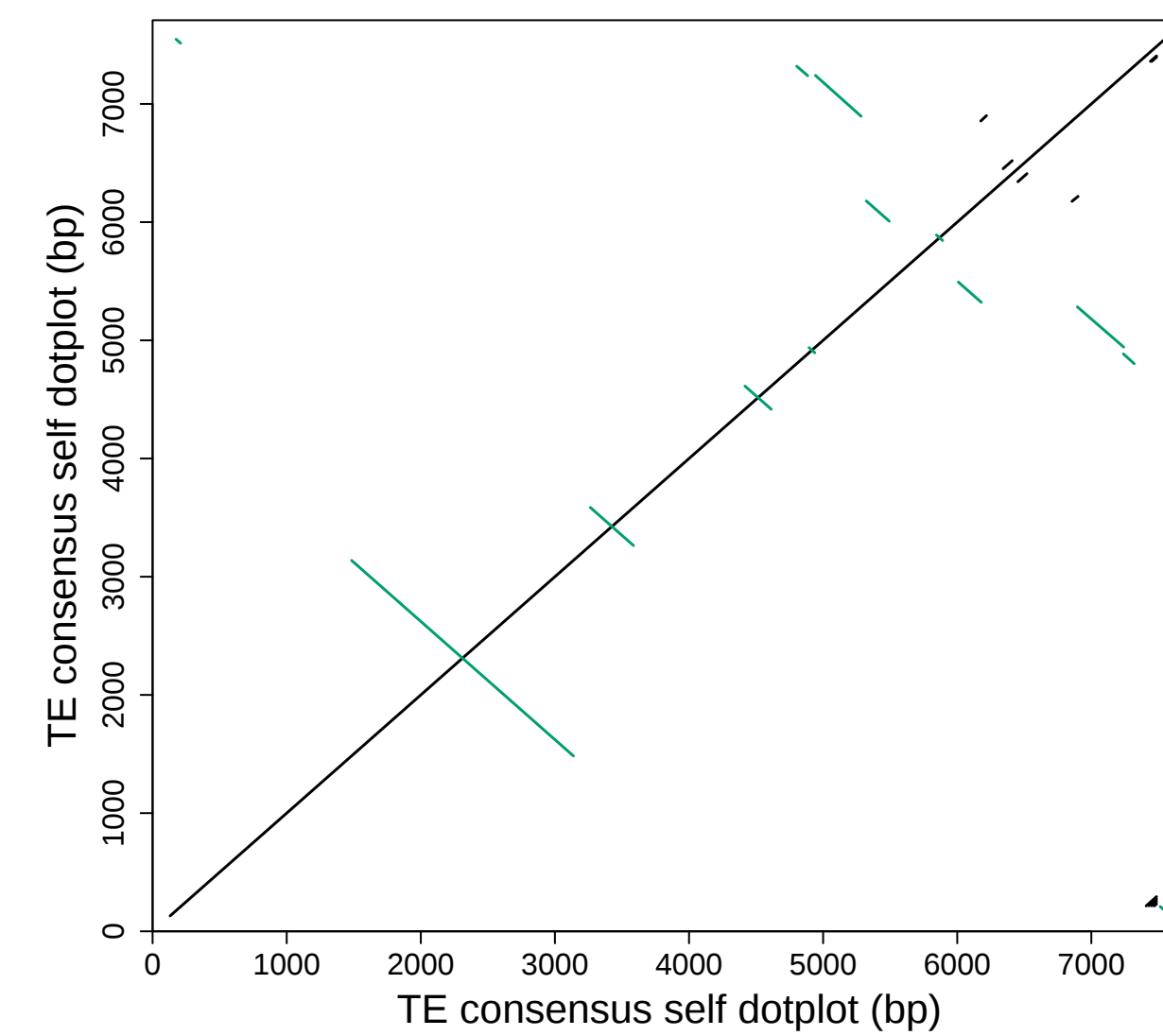
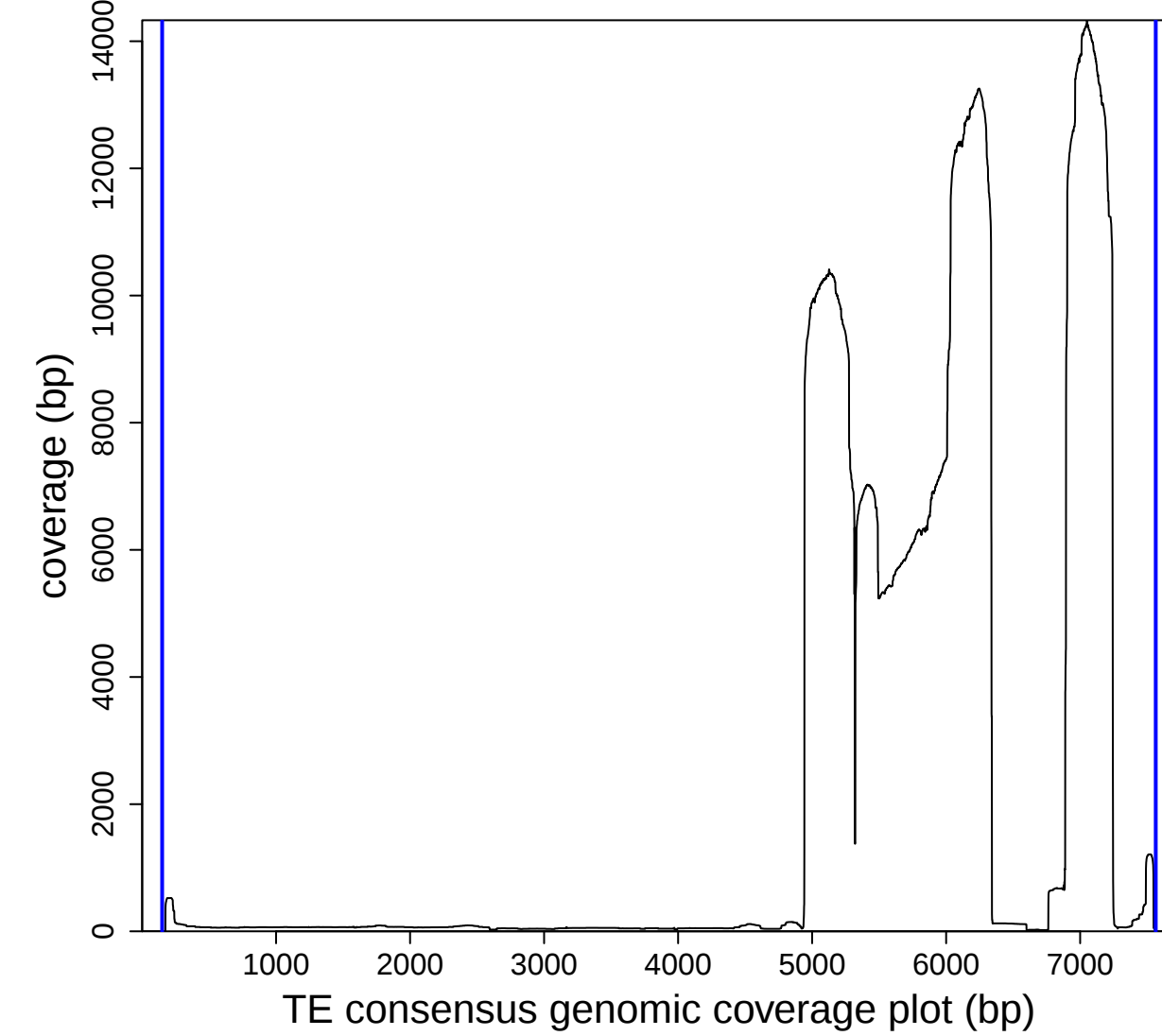
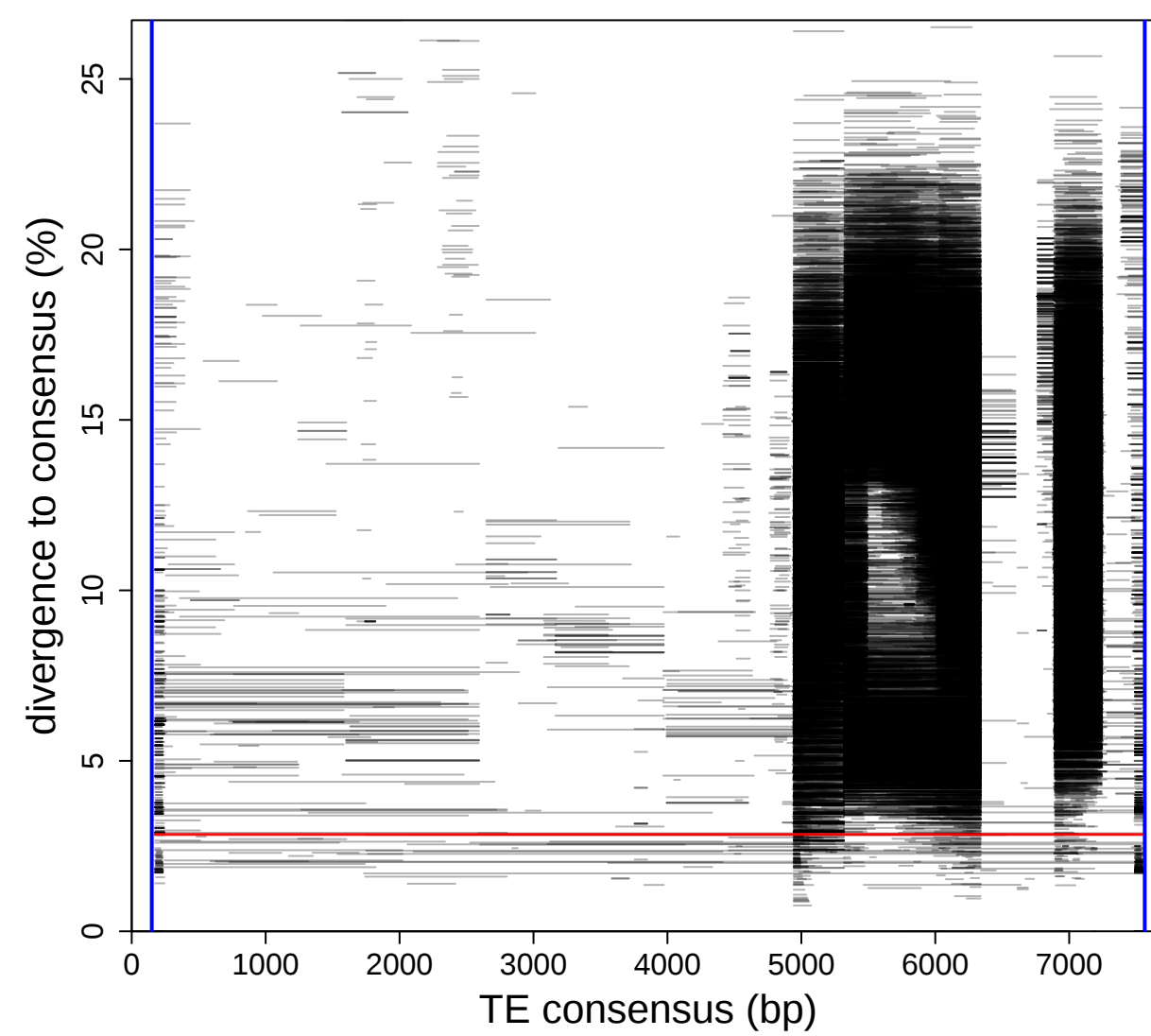
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_al
size: 7403bp; fragments: 49461; full length: 2 (>=6662.7bp)

After TEtrimmer 7403 bp



bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 7714bp; fragments: 48729; full length: 1 (>=6942.6bp)

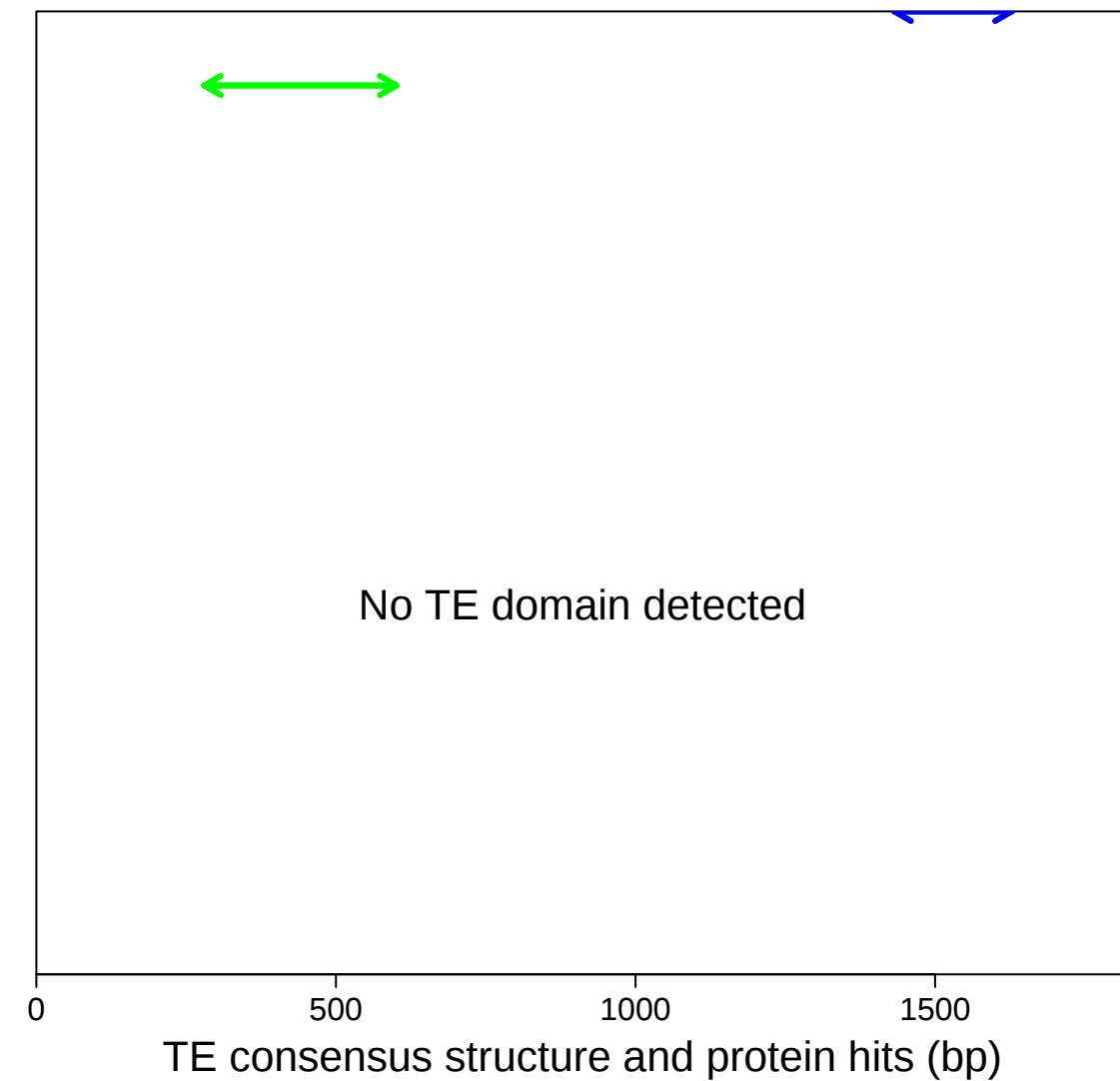
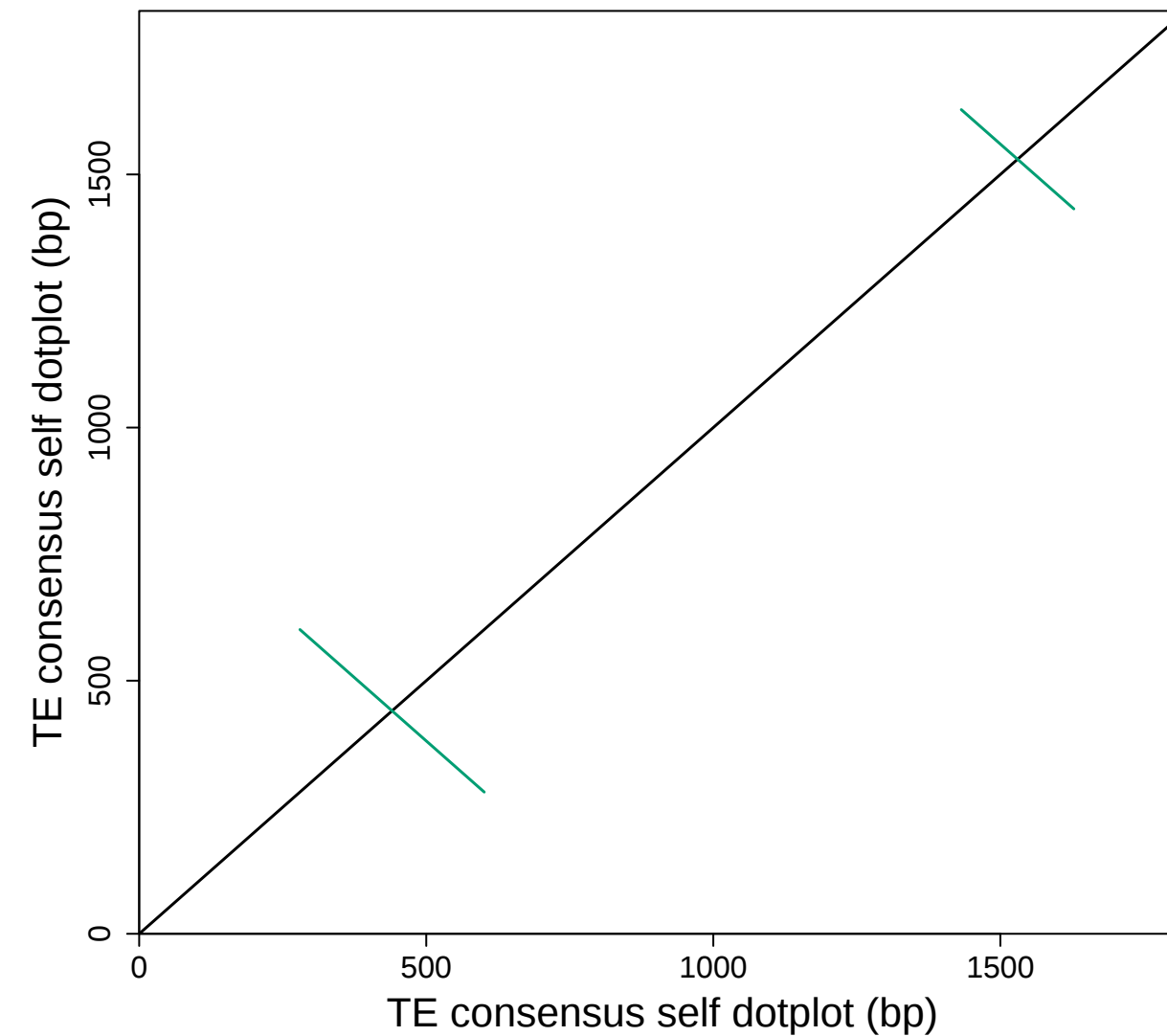
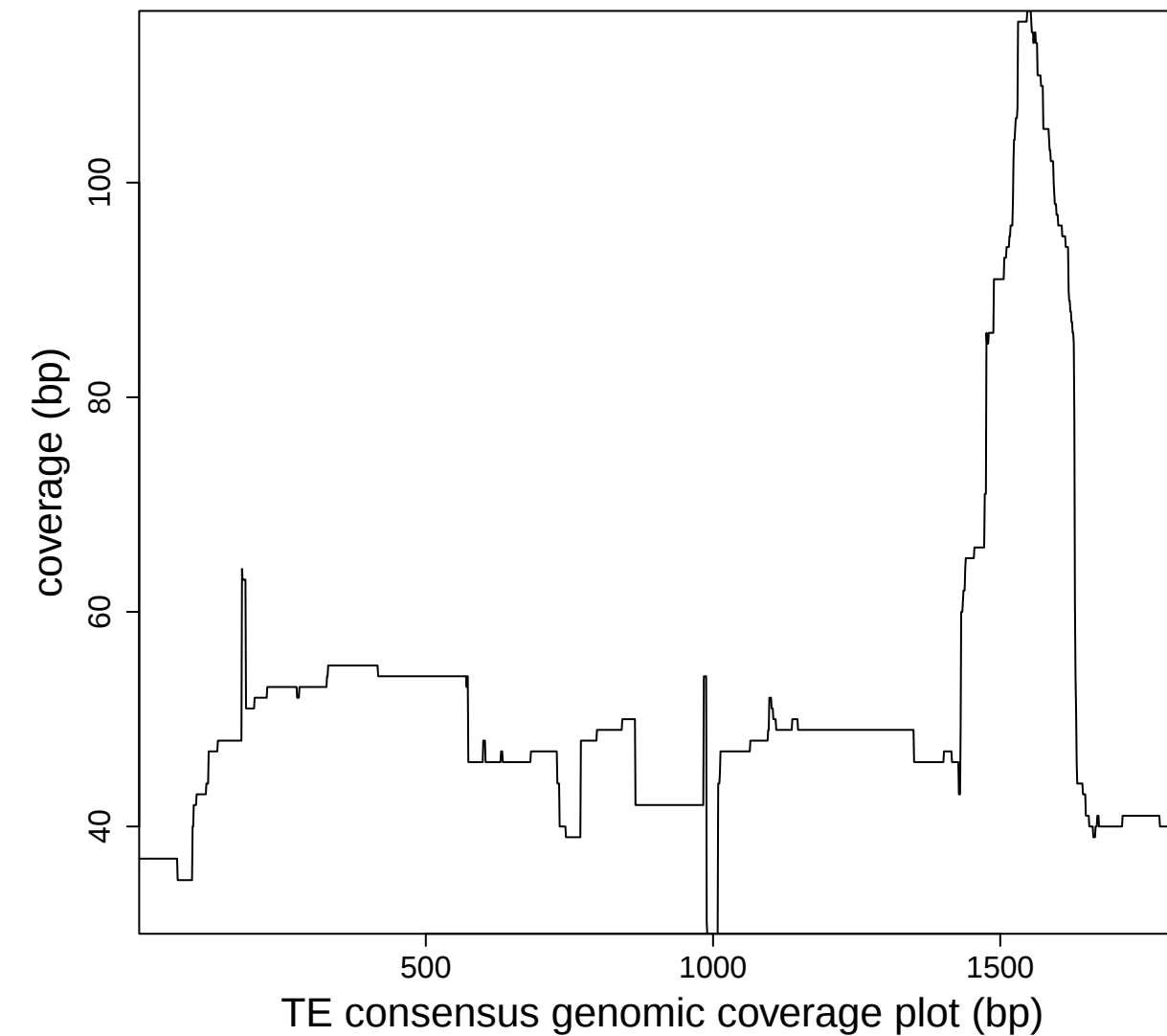
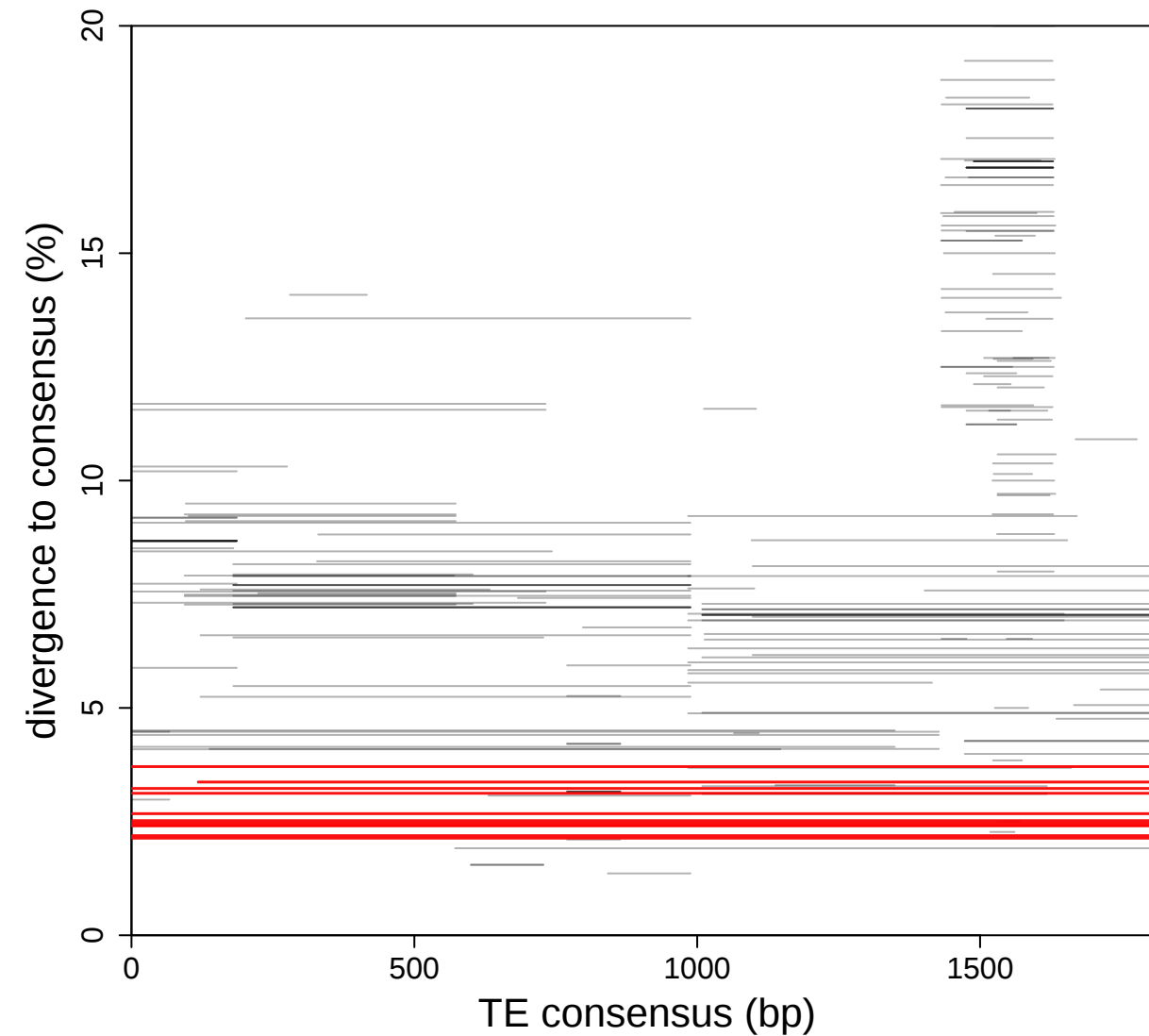
After TEtrimmer Extended plot Blue lines are boundaries



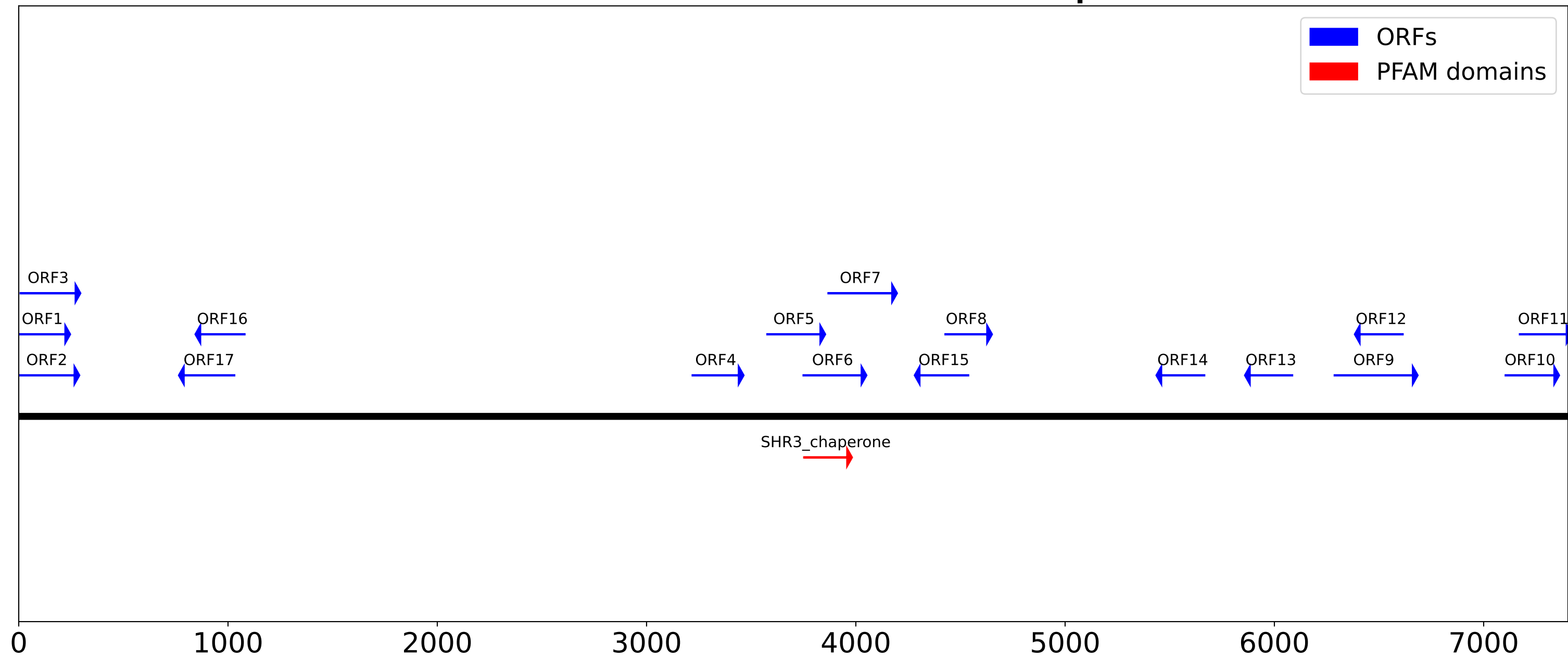
Before TEtrimmer 1823 bp

TE: rnd_5_family_51

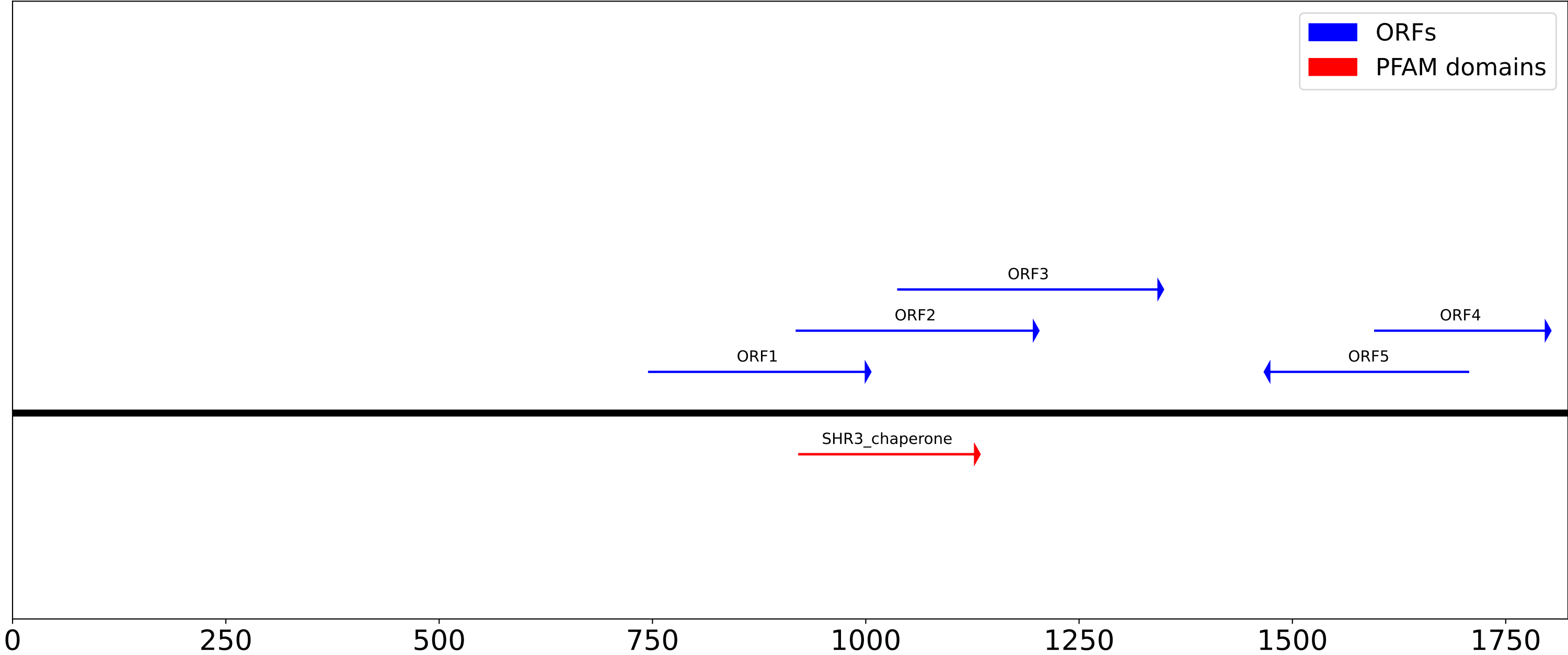
size: 1823bp; fragments: 202; full length: 11 (≥ 1640.7 bp)



After TEtrimmer ORF and PFAM domain plot



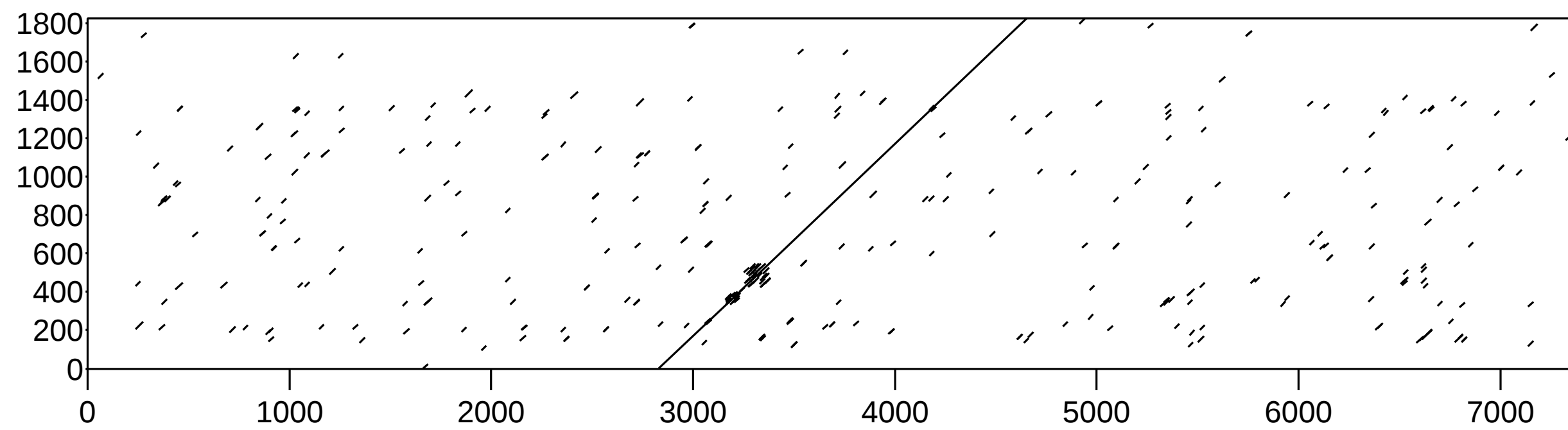
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)