

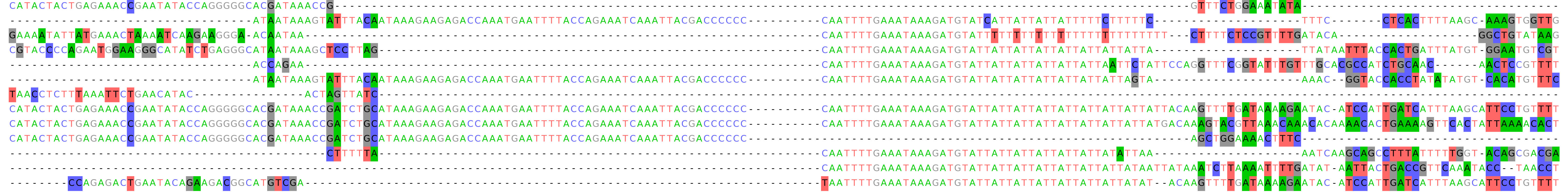
Start crop Point



End crop Point

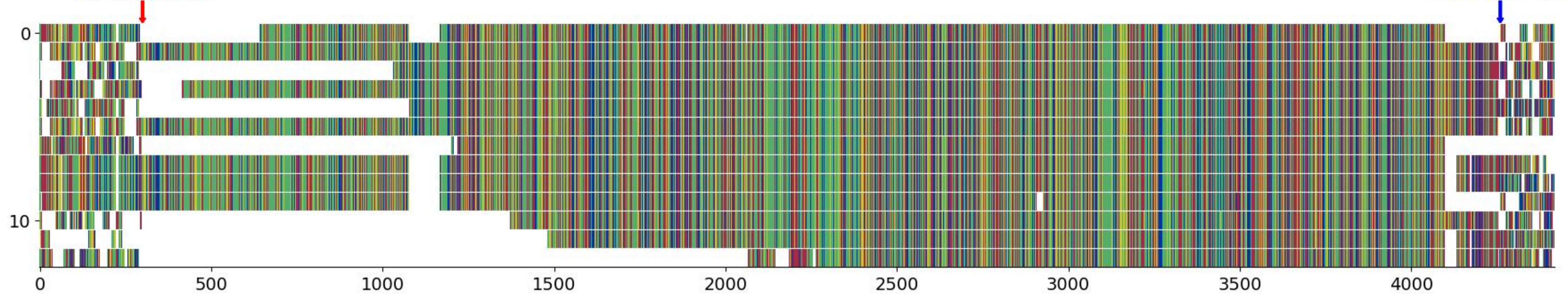


MSA length = 3961



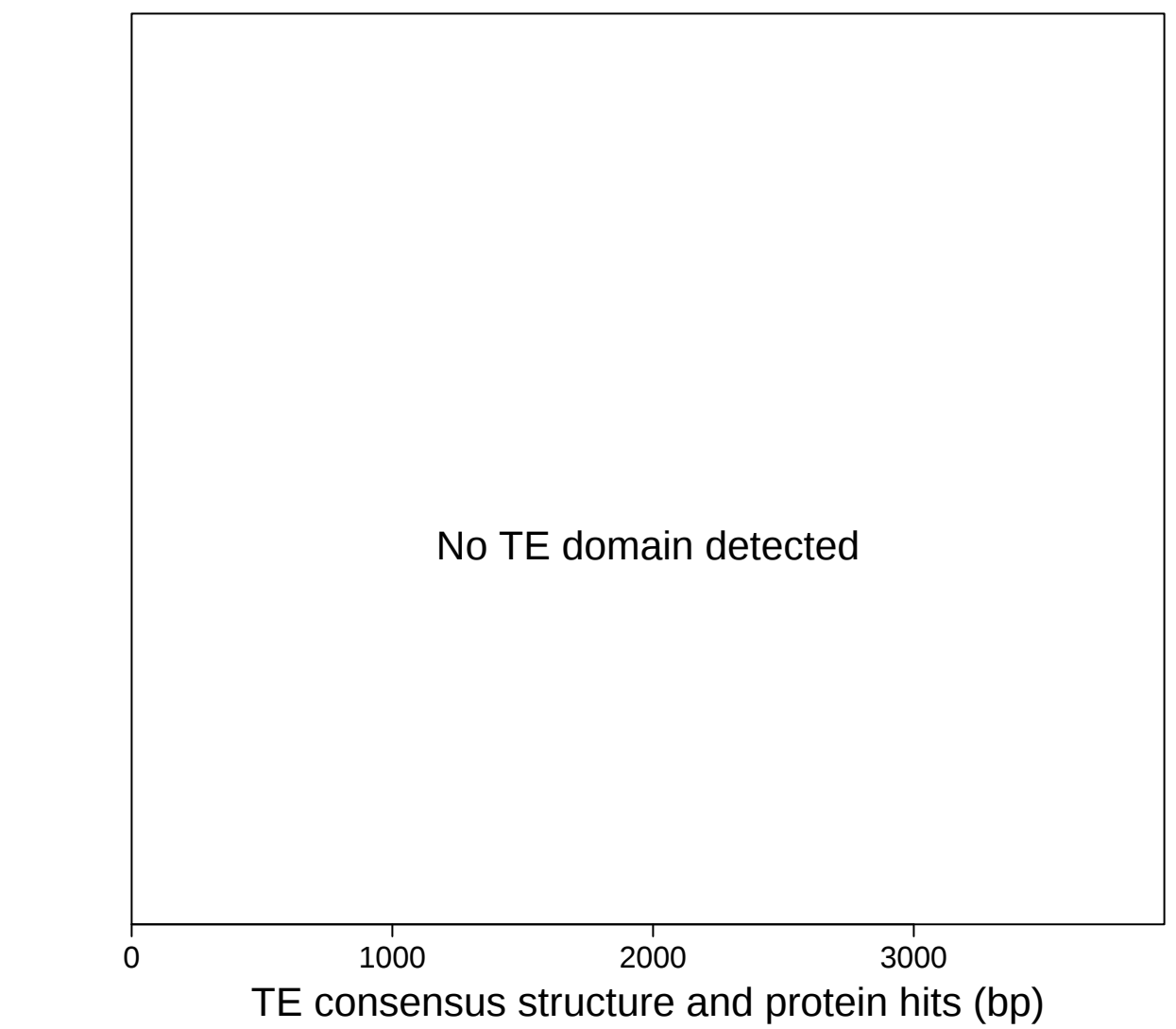
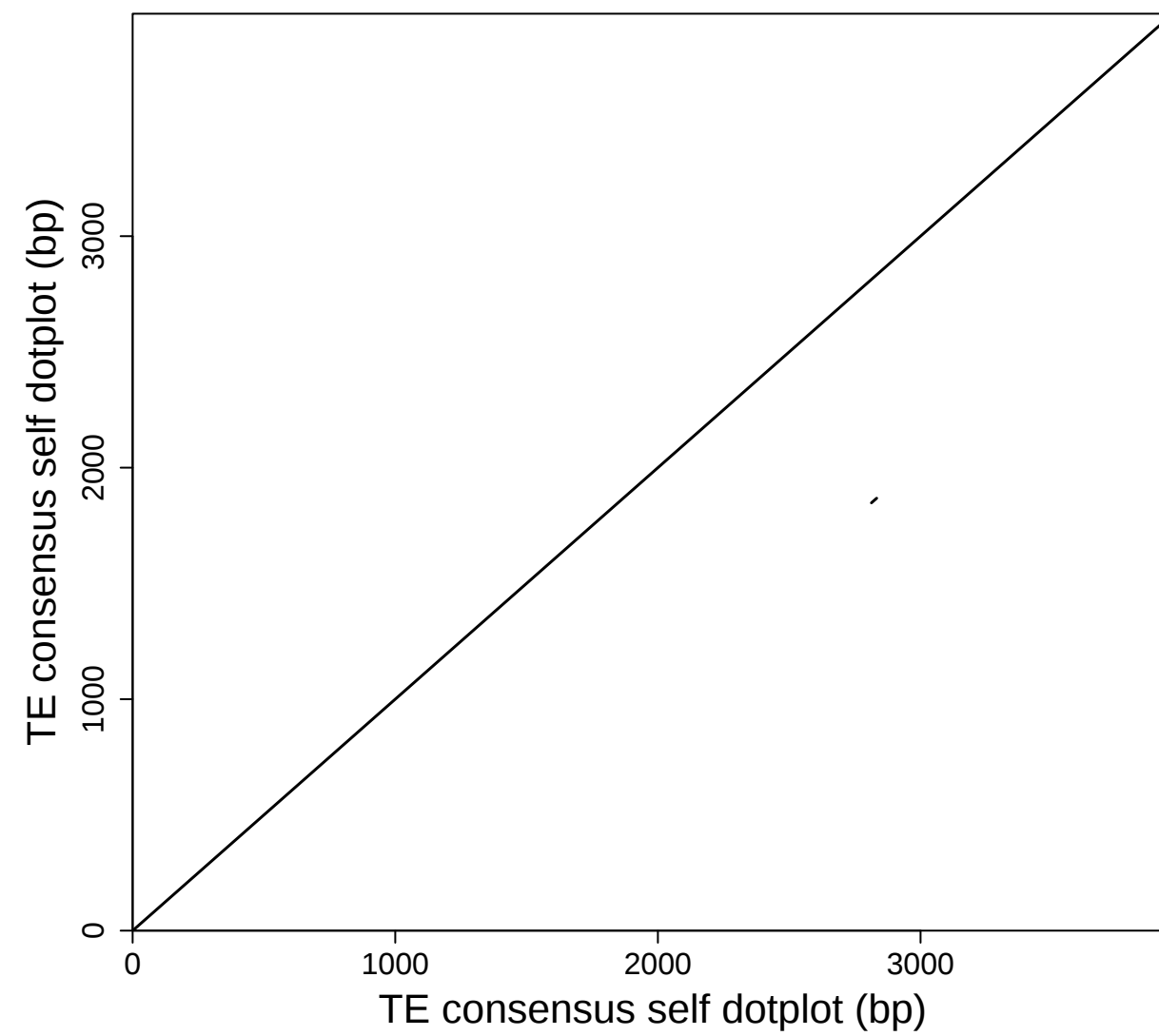
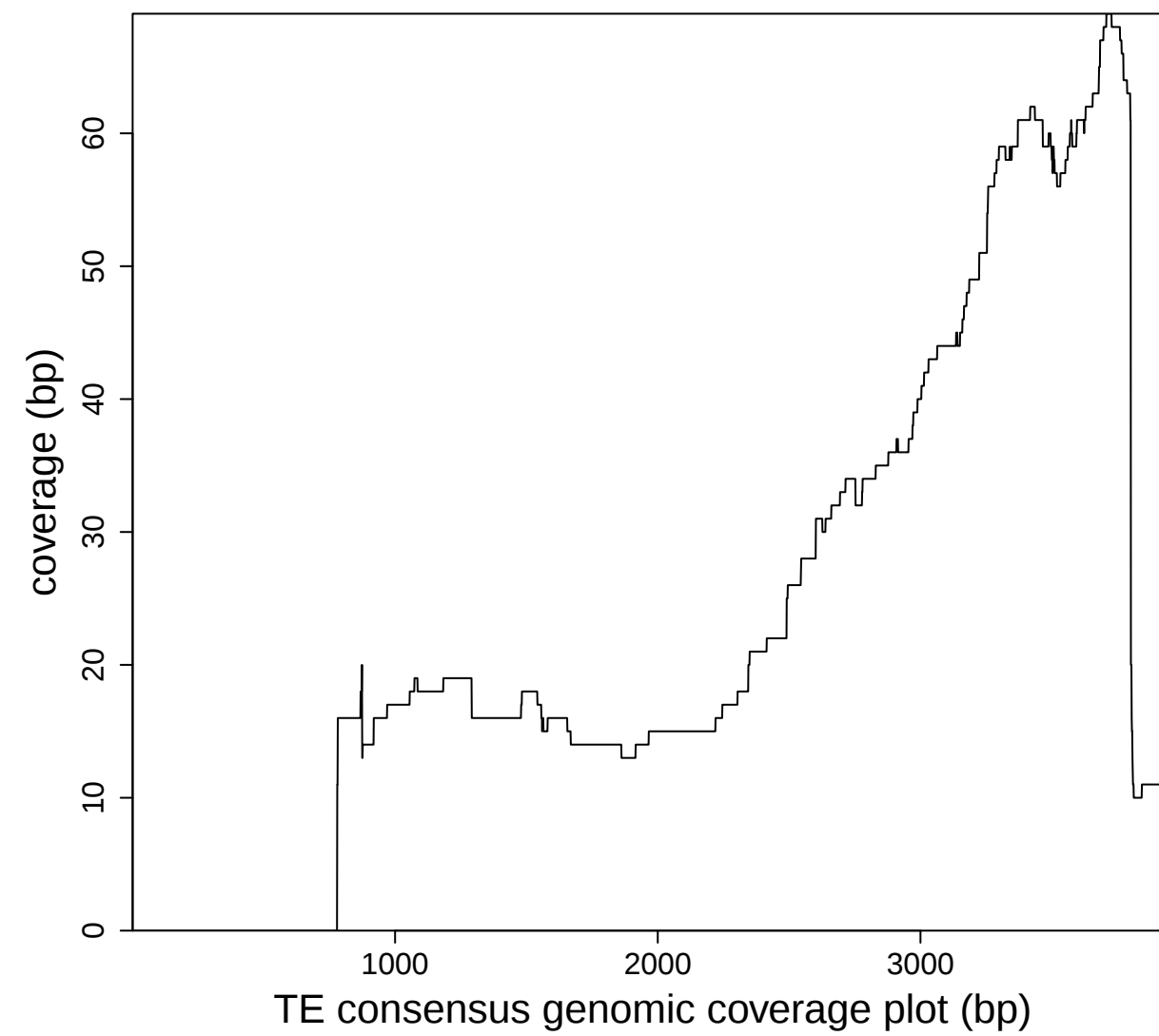
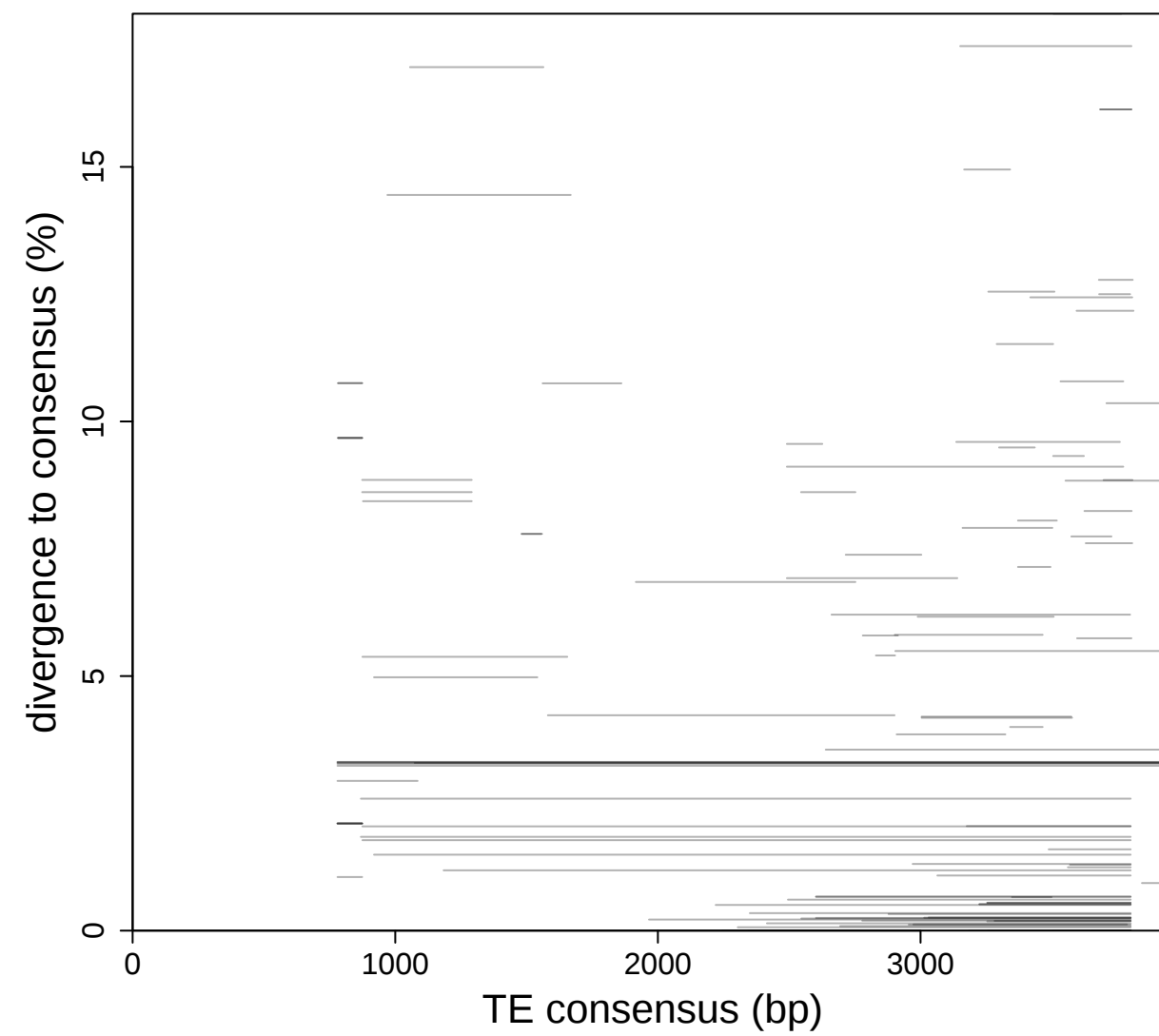
Start crop Point

End crop Point



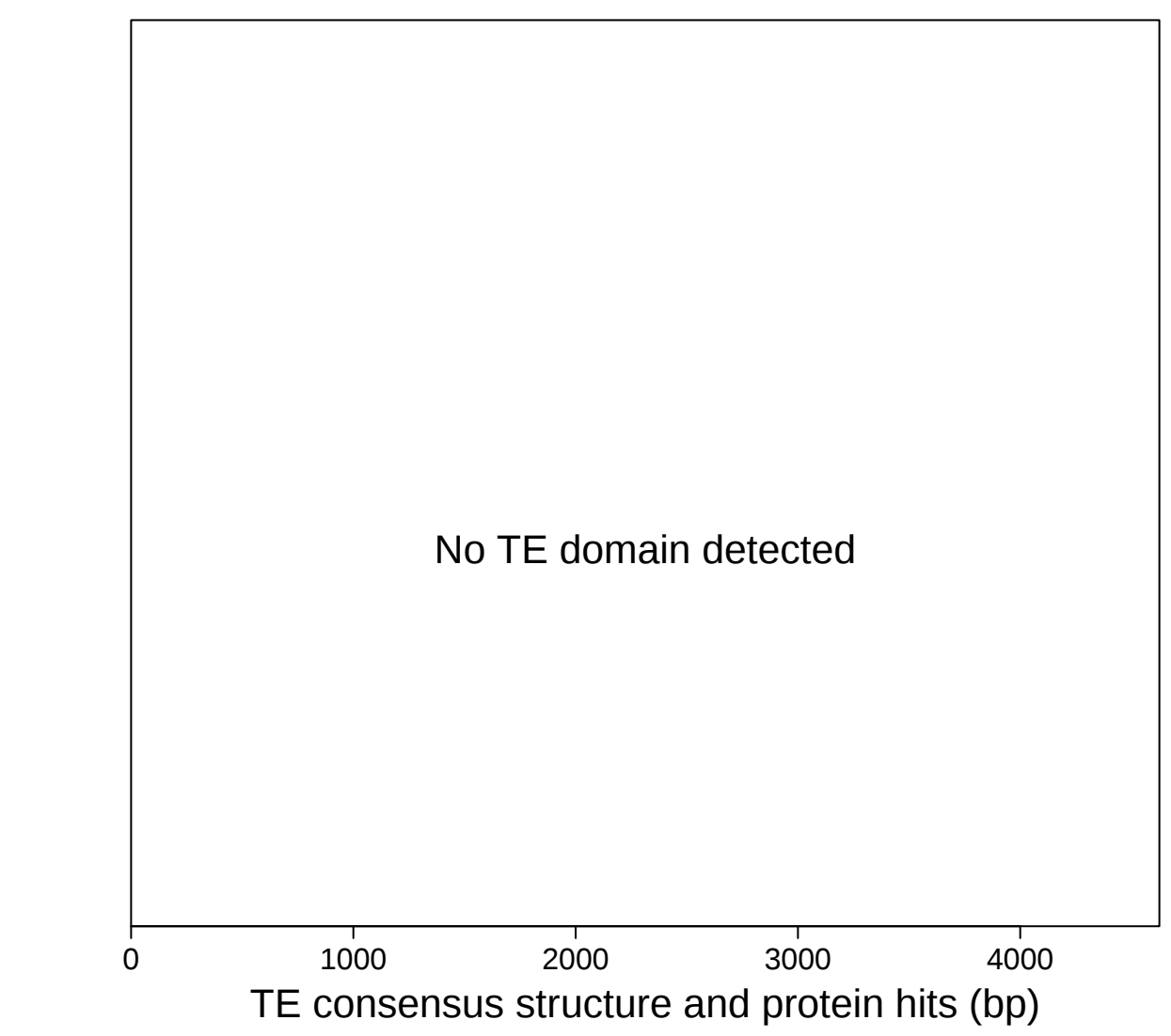
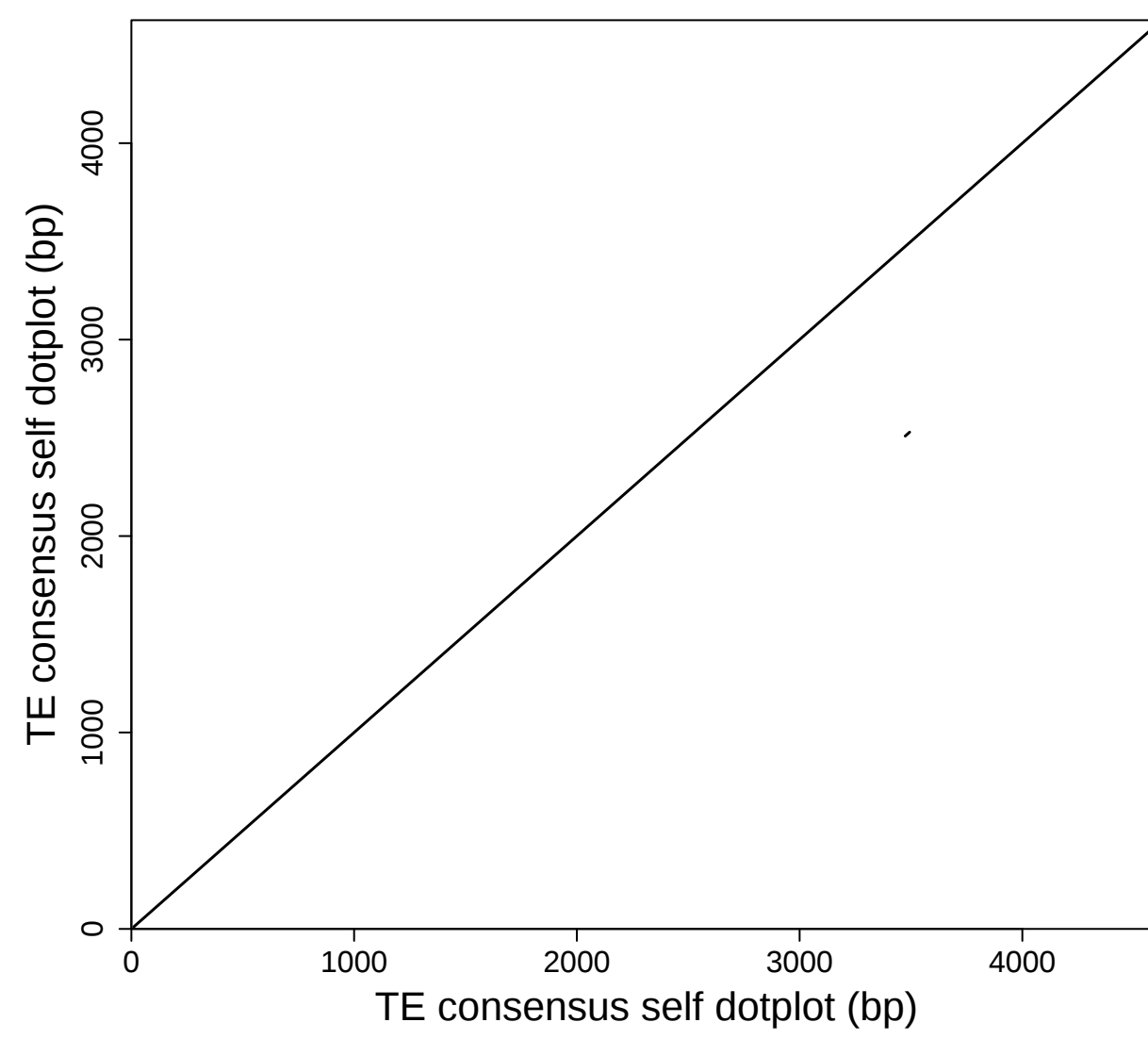
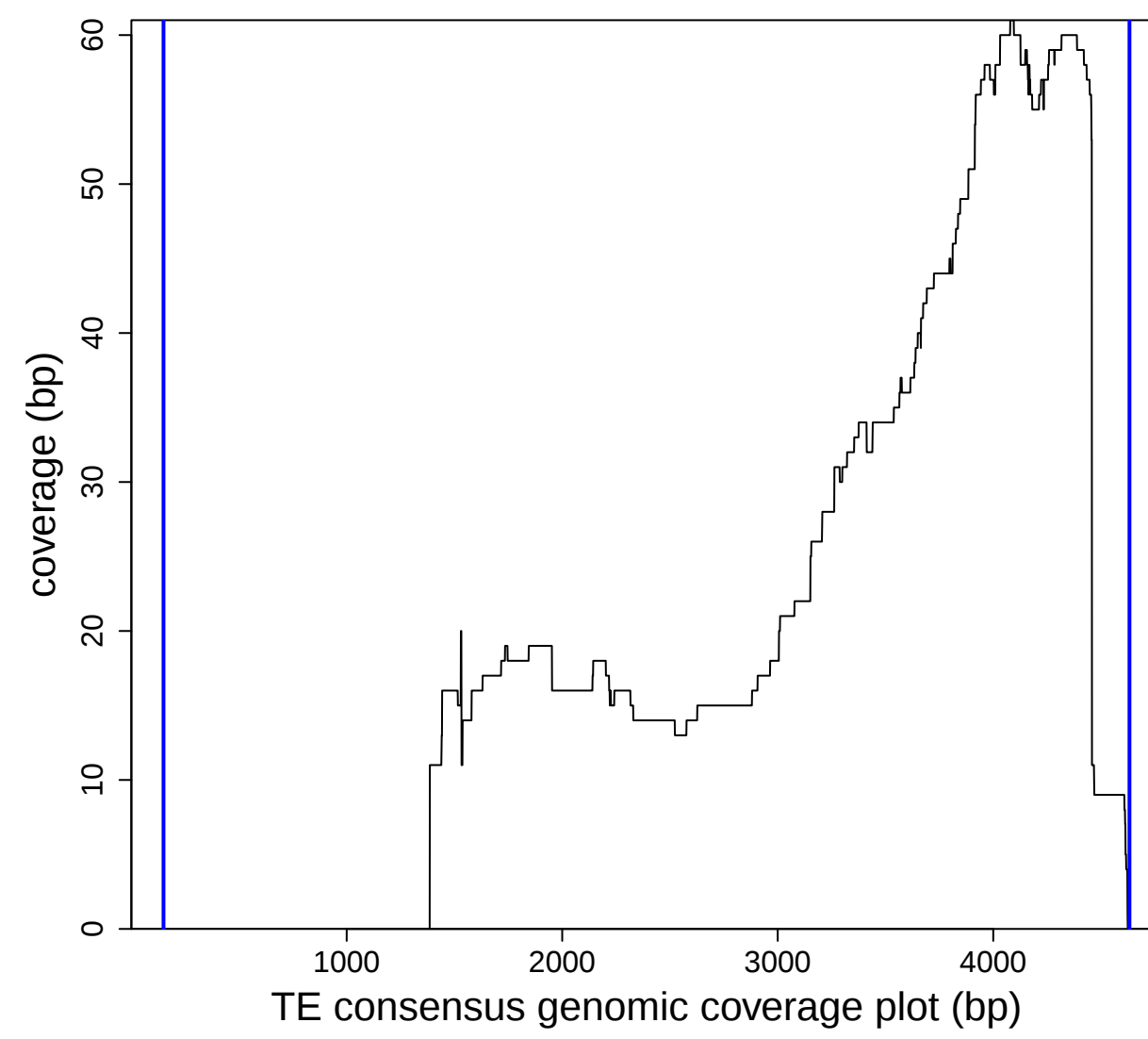
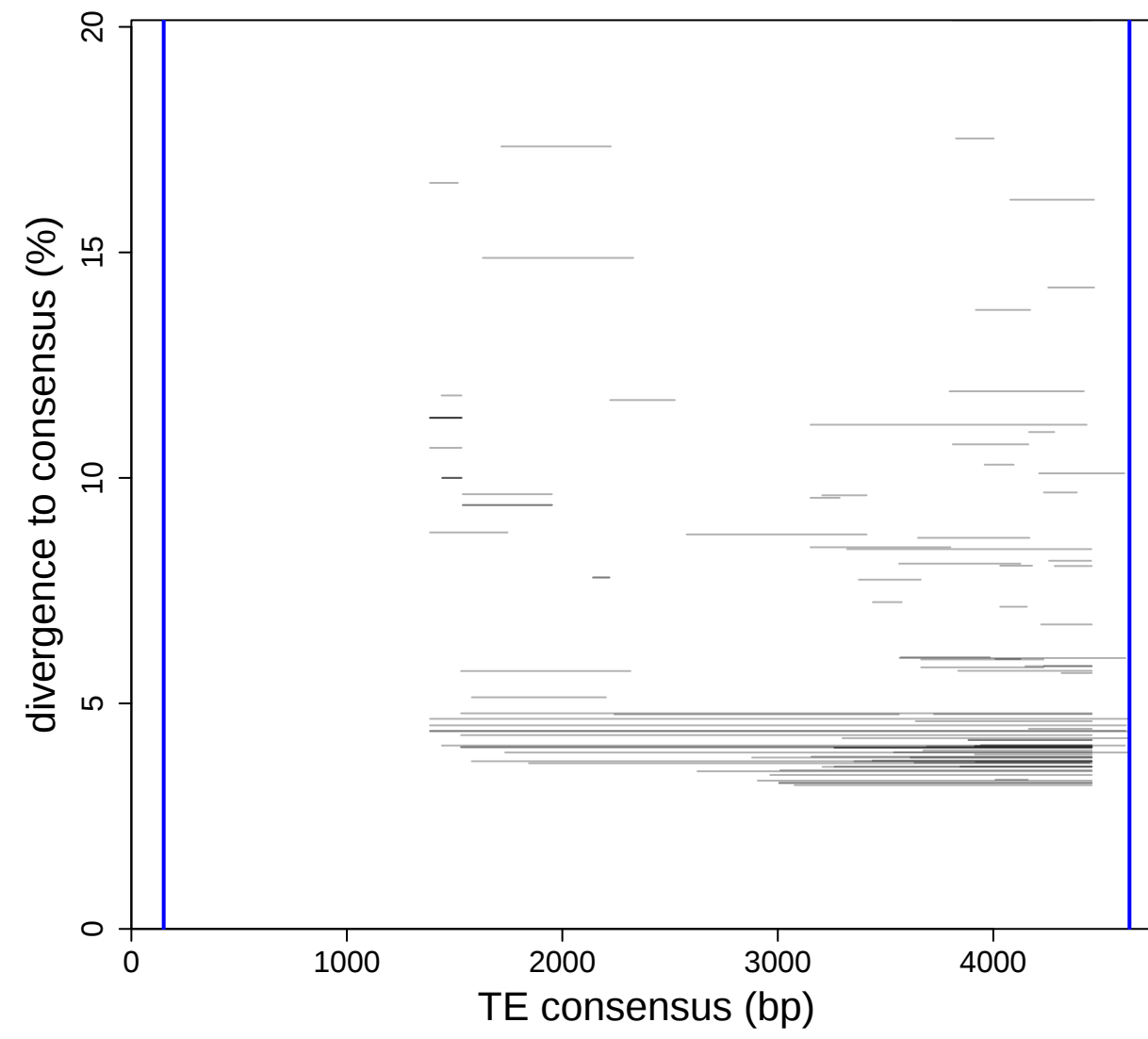
1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.faaln.f
size: 3961bp; fragments: 114; full length: 0 (>=3564.9bp)

After TEtrimmer 3961 bp



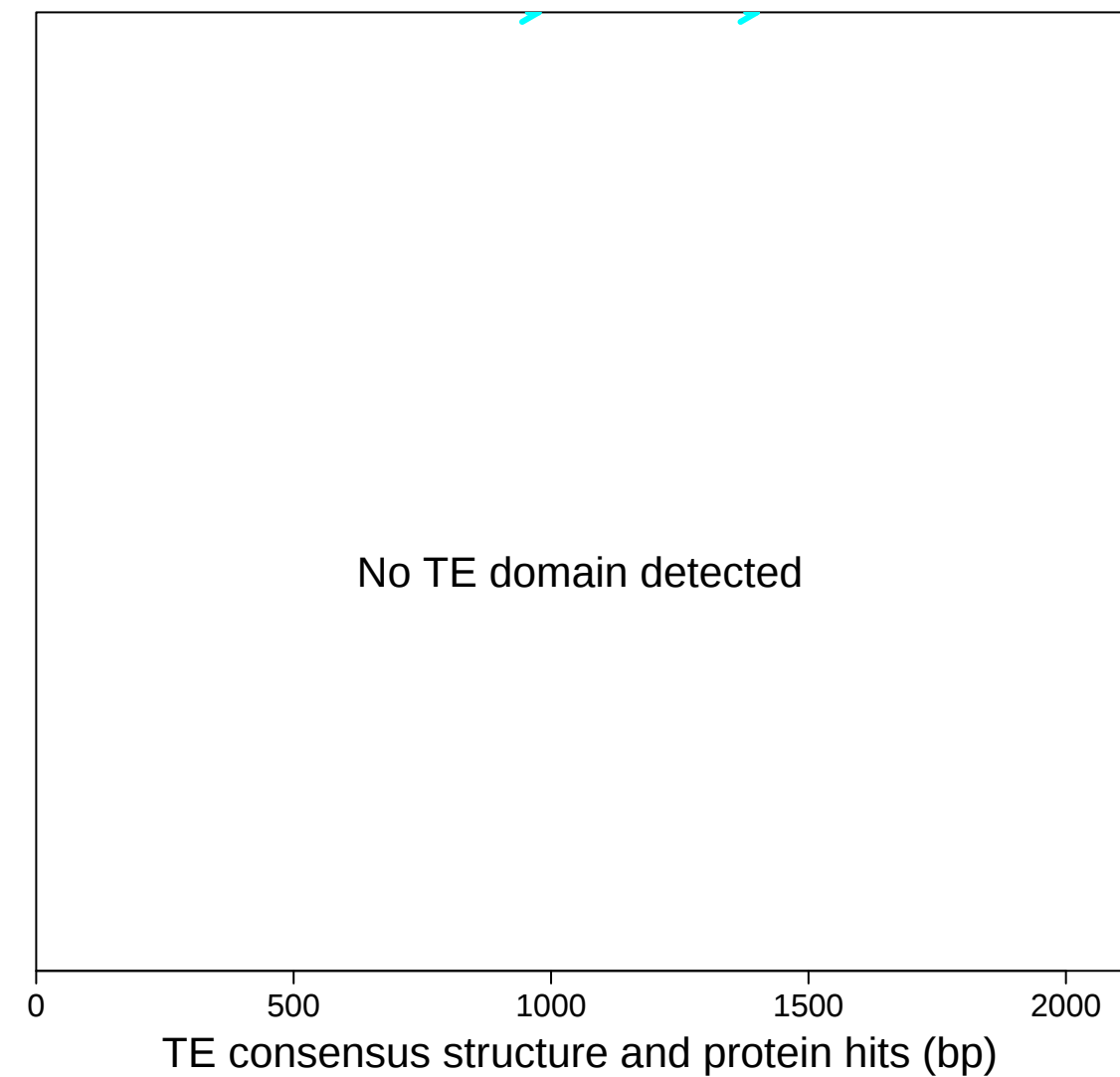
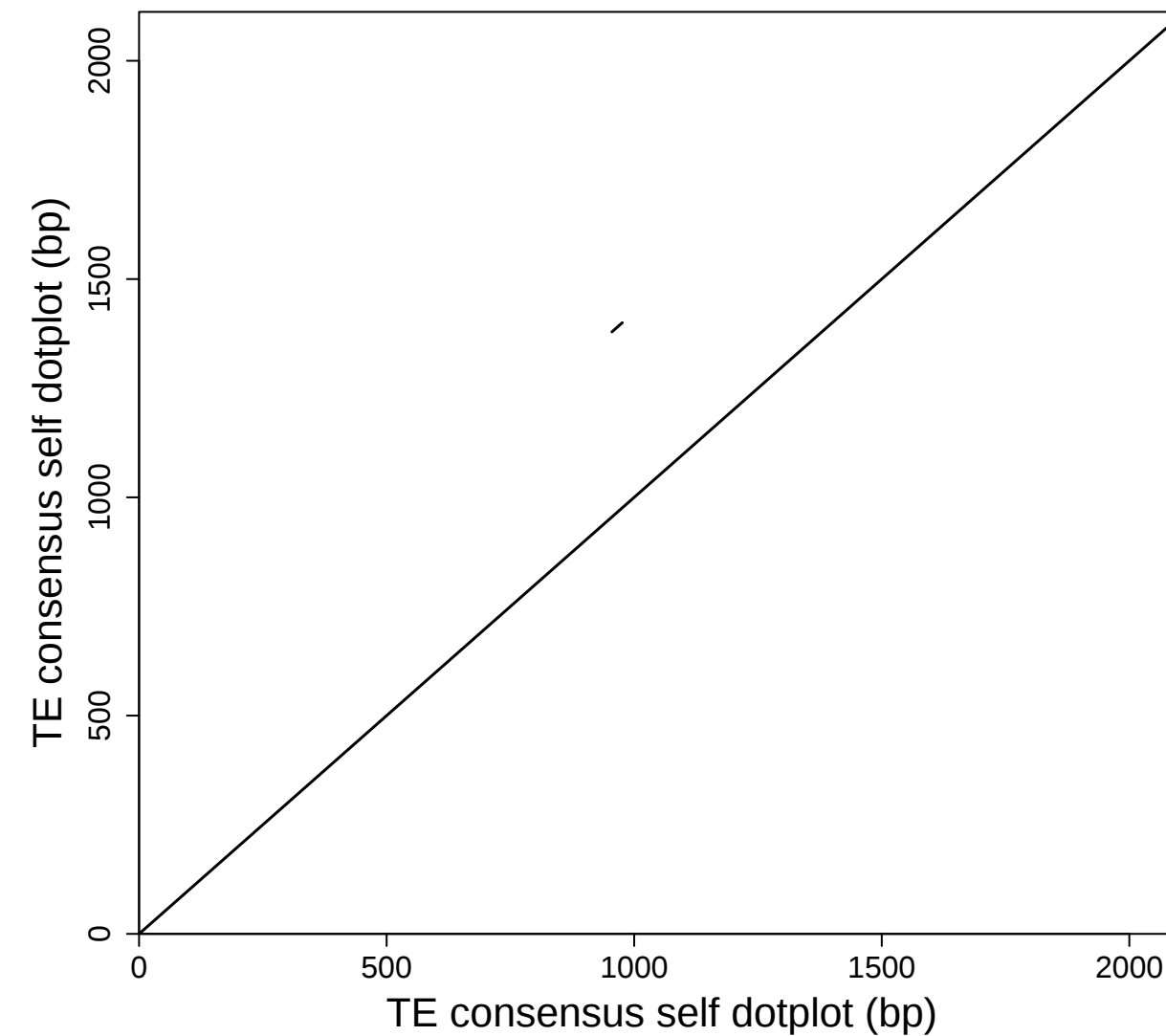
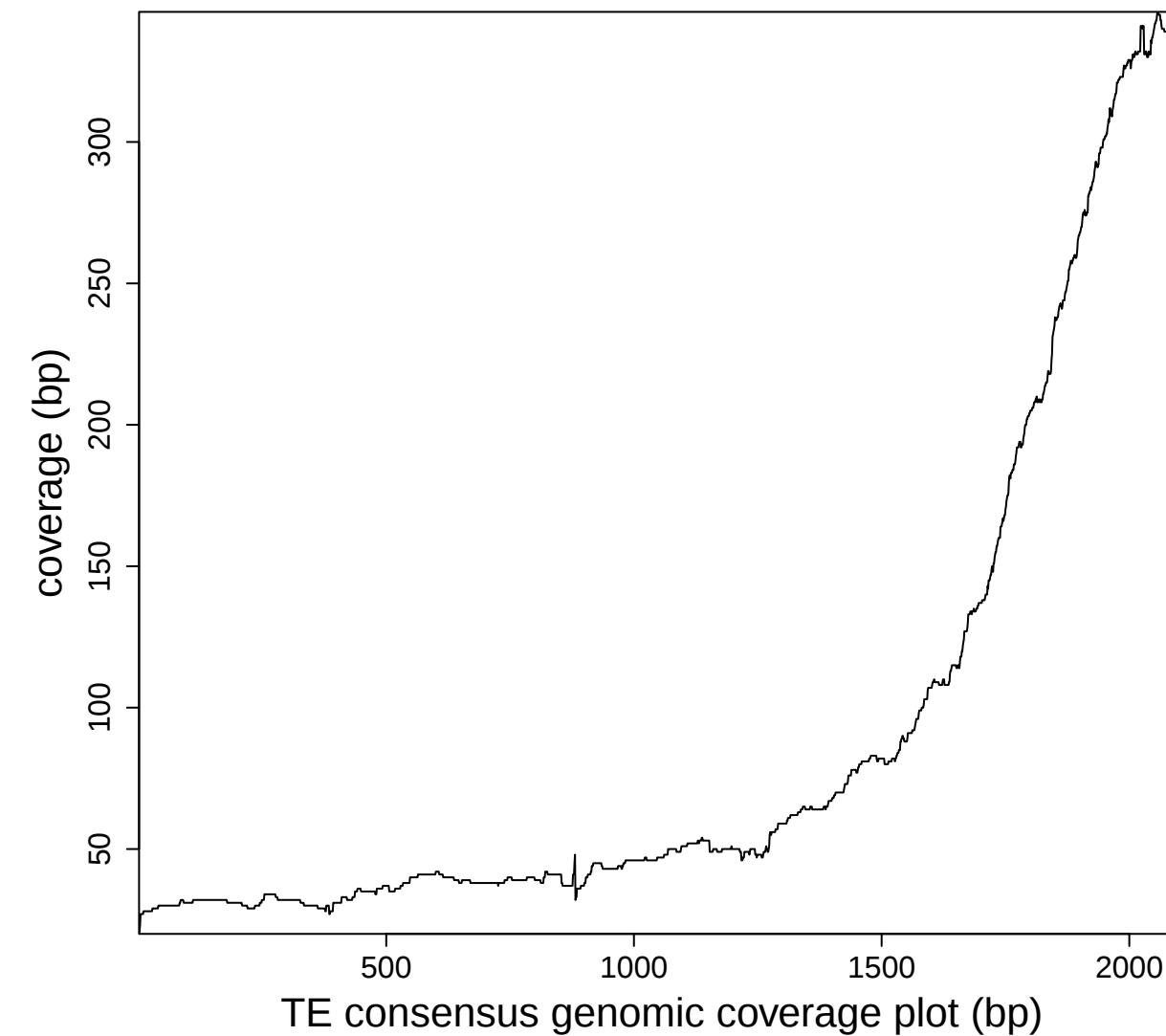
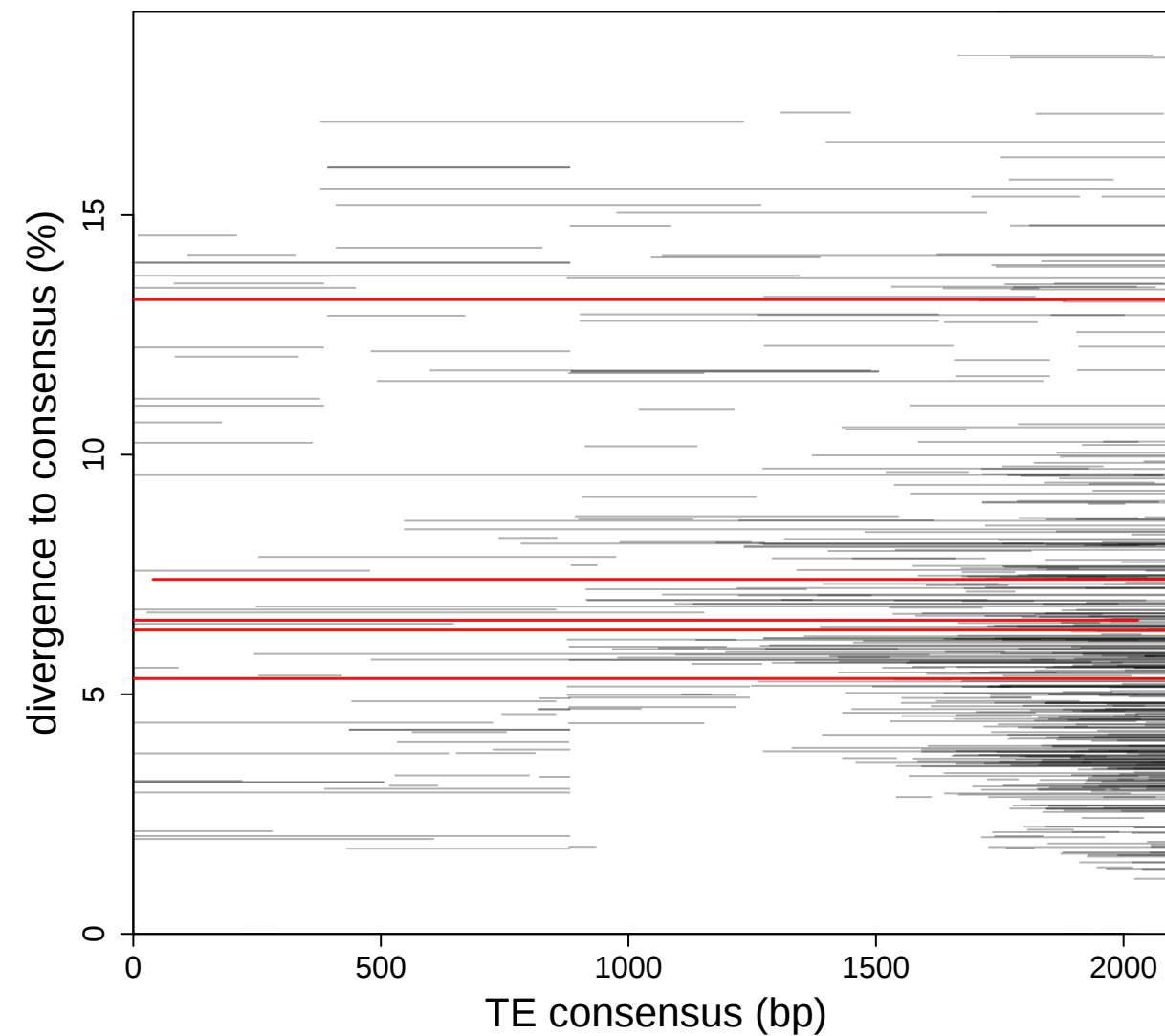
f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 4782bp; fragments: 102; full length: 0 (>=4303.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

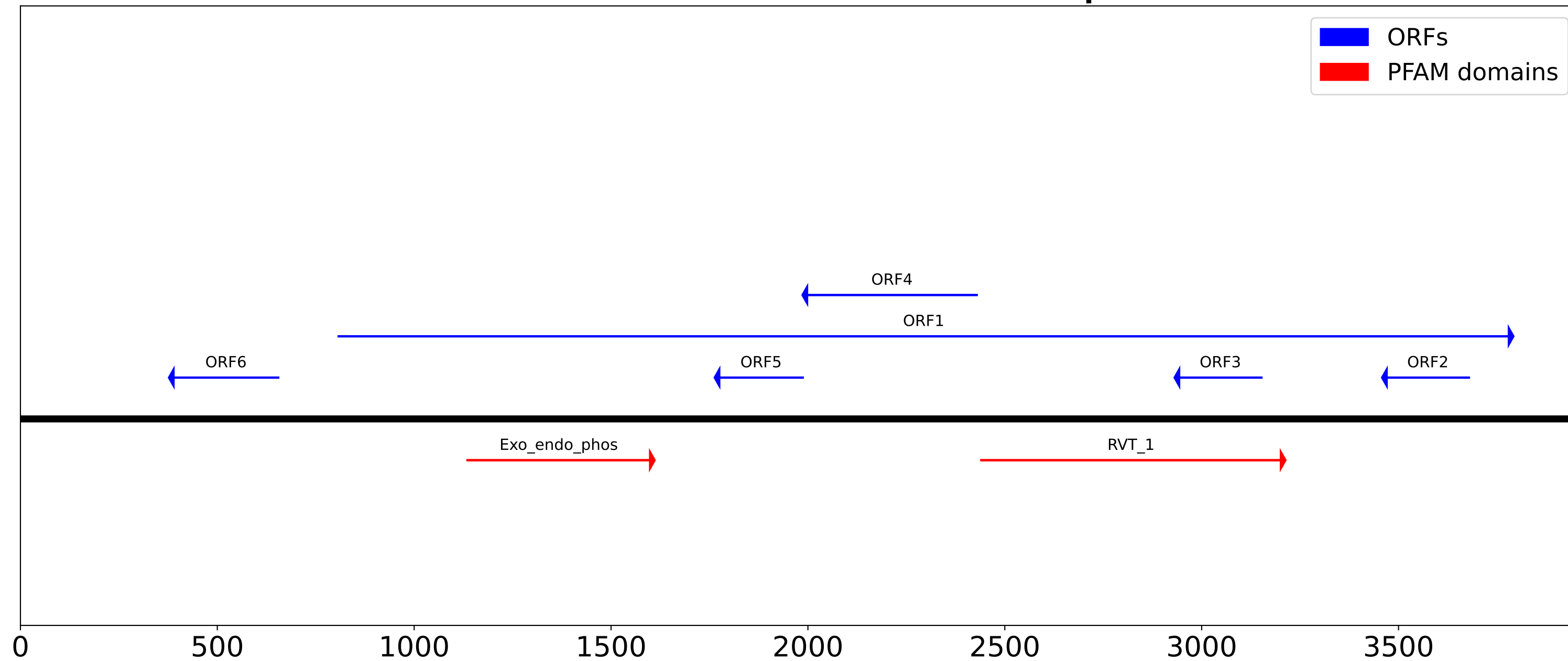


TE: rnd_5_family_4500
size: 2112bp; fragments: 530; full length: 5 (≥ 1900.8 bp)

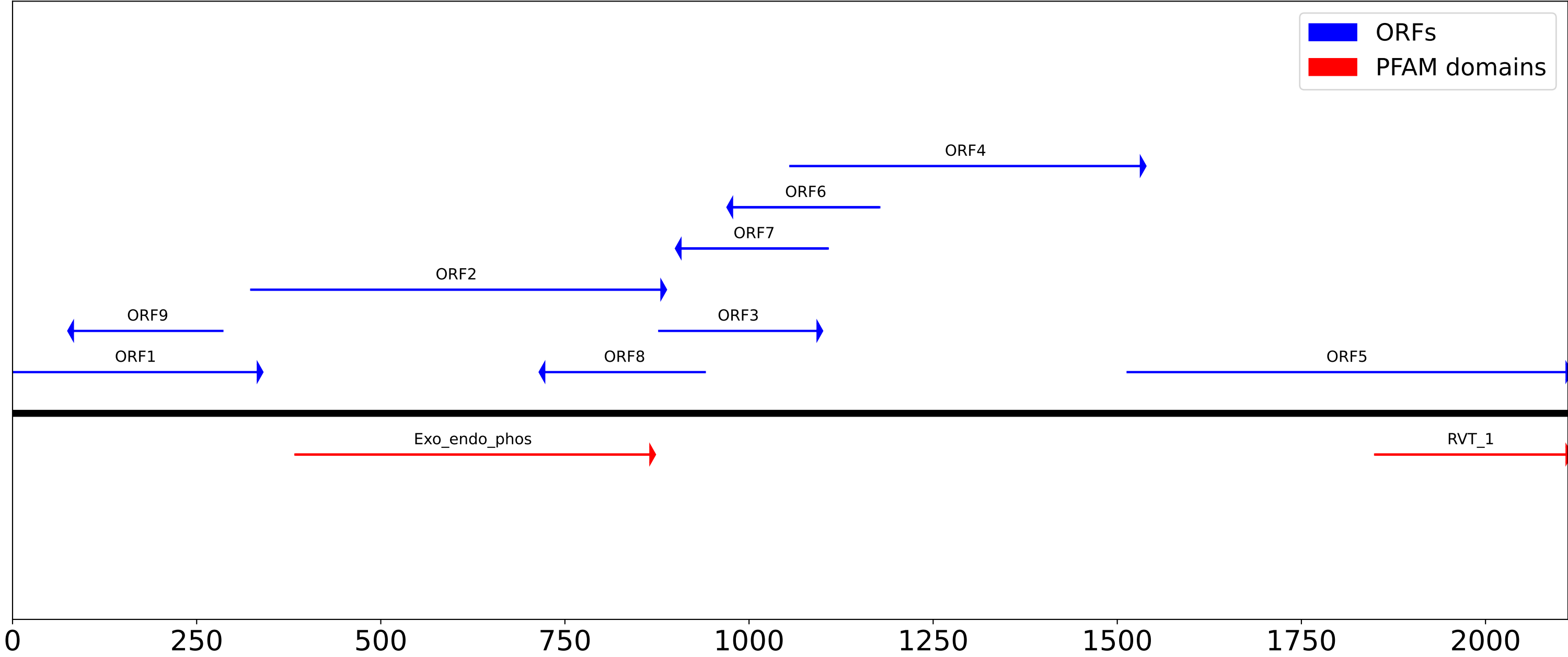
Before TEtrimmer 2112 bp



After TEtrimmer ORF and PFAM domain plot



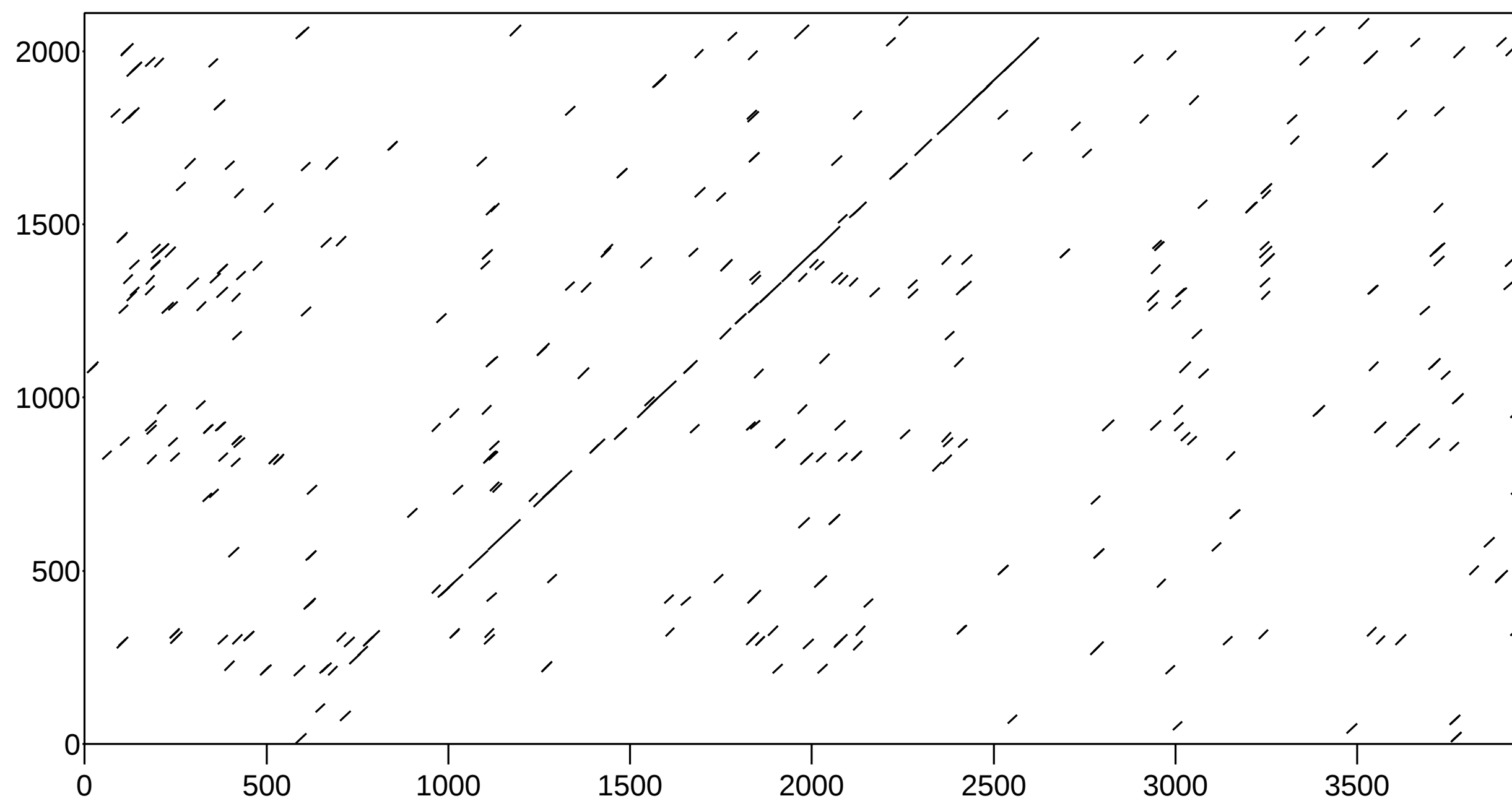
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)