

MSA length = 2394

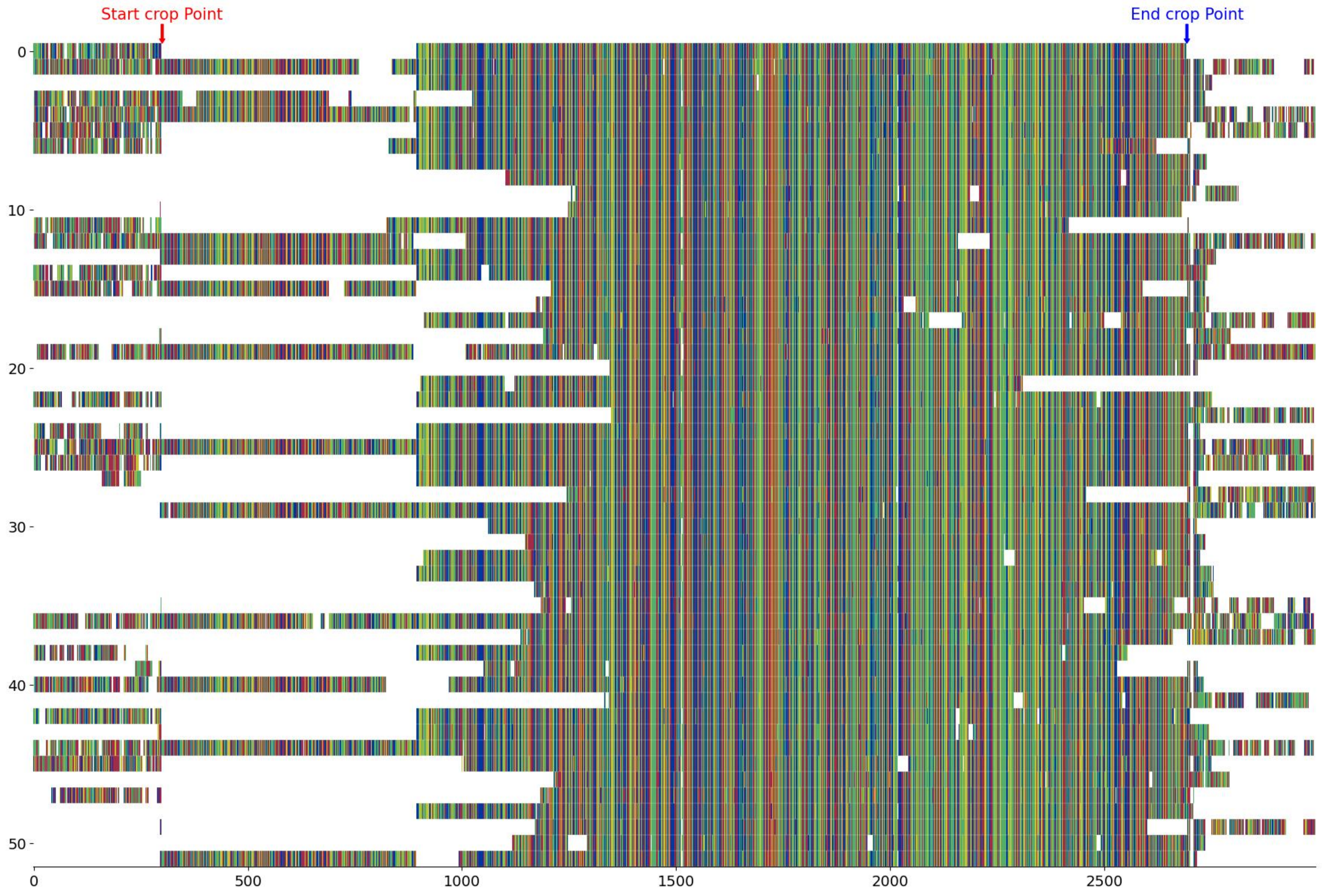
Start crop Point

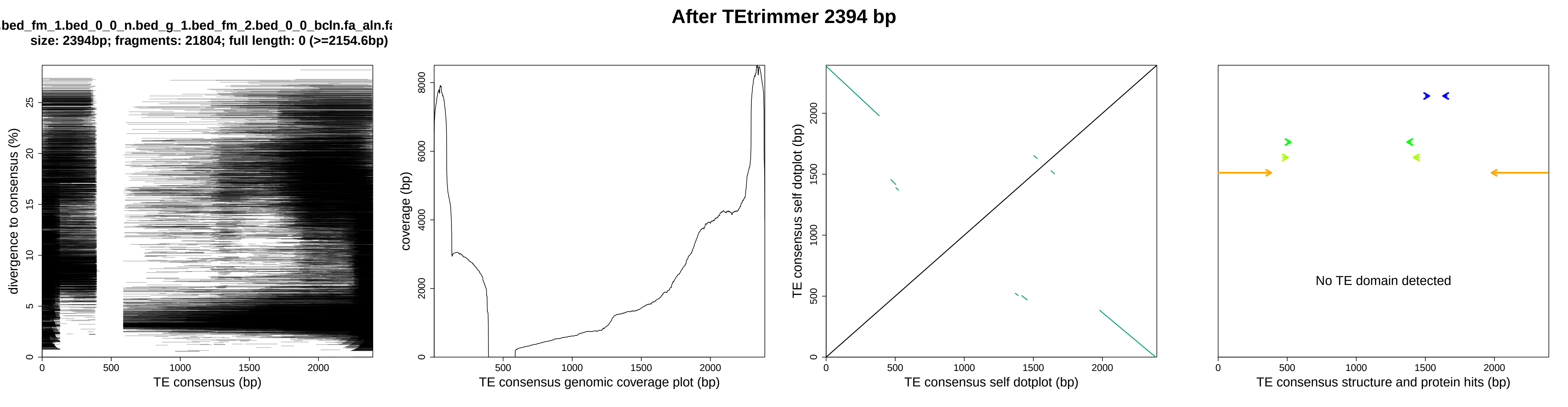
End crop Point



Start crop Point

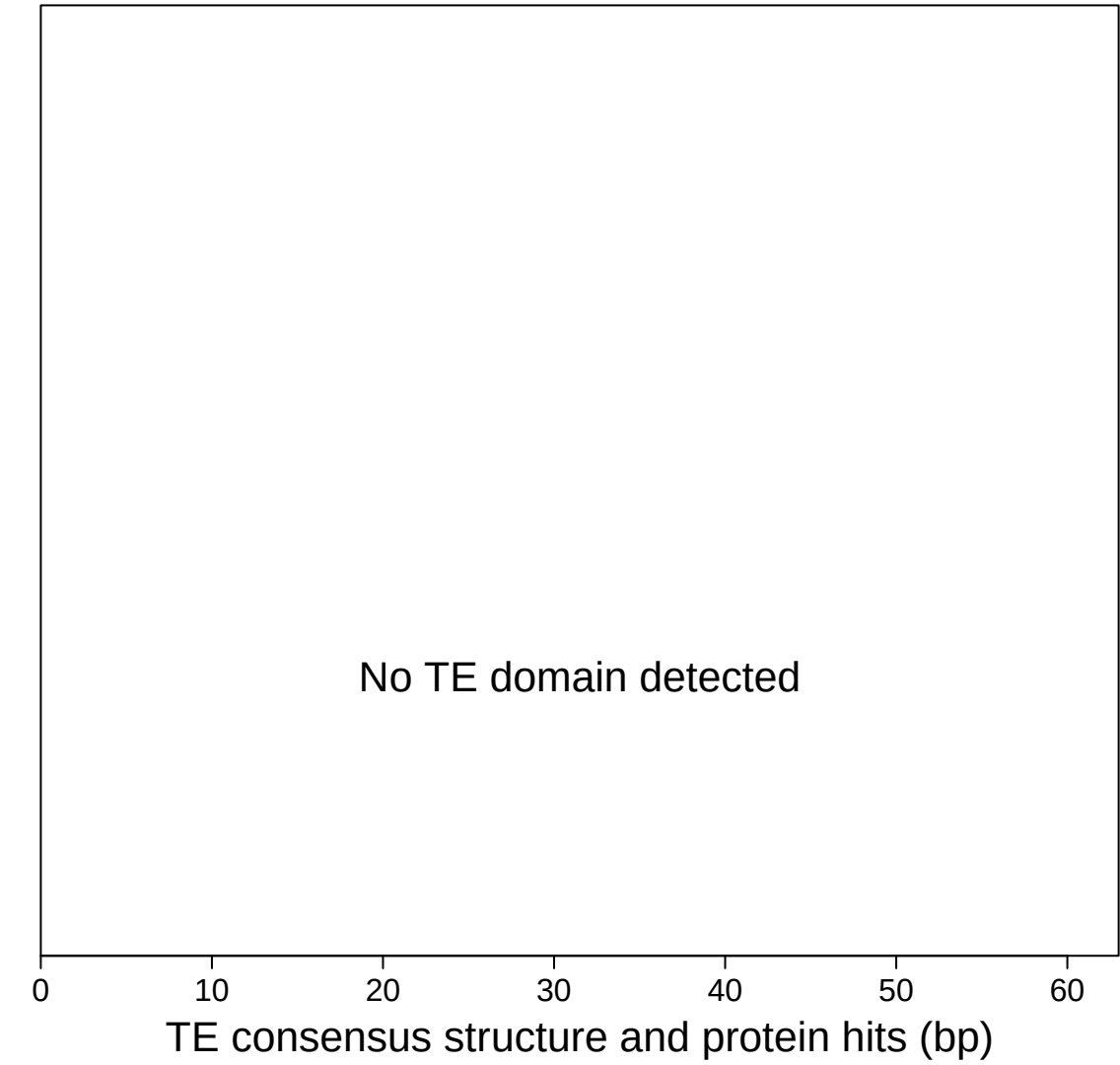
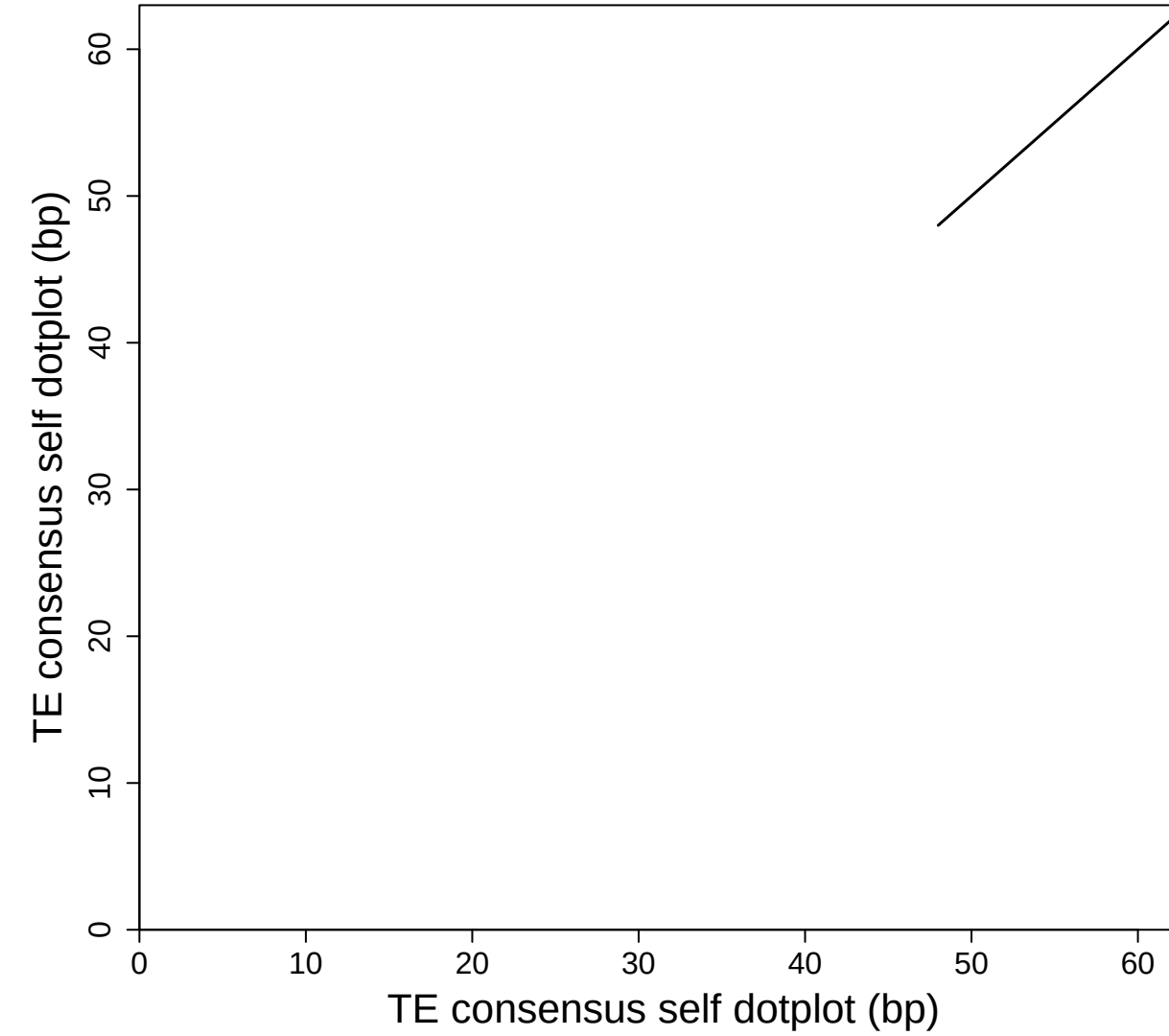
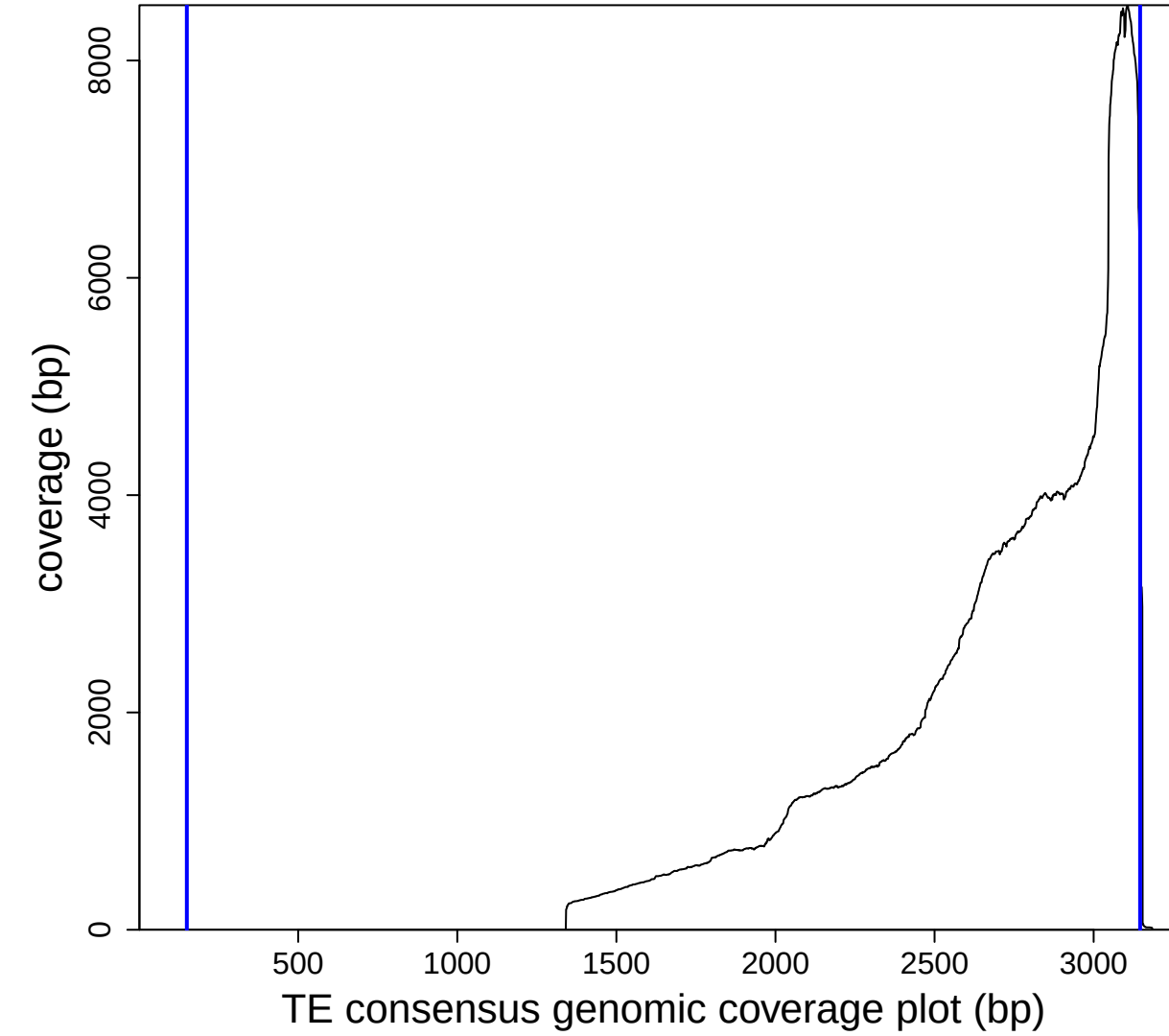
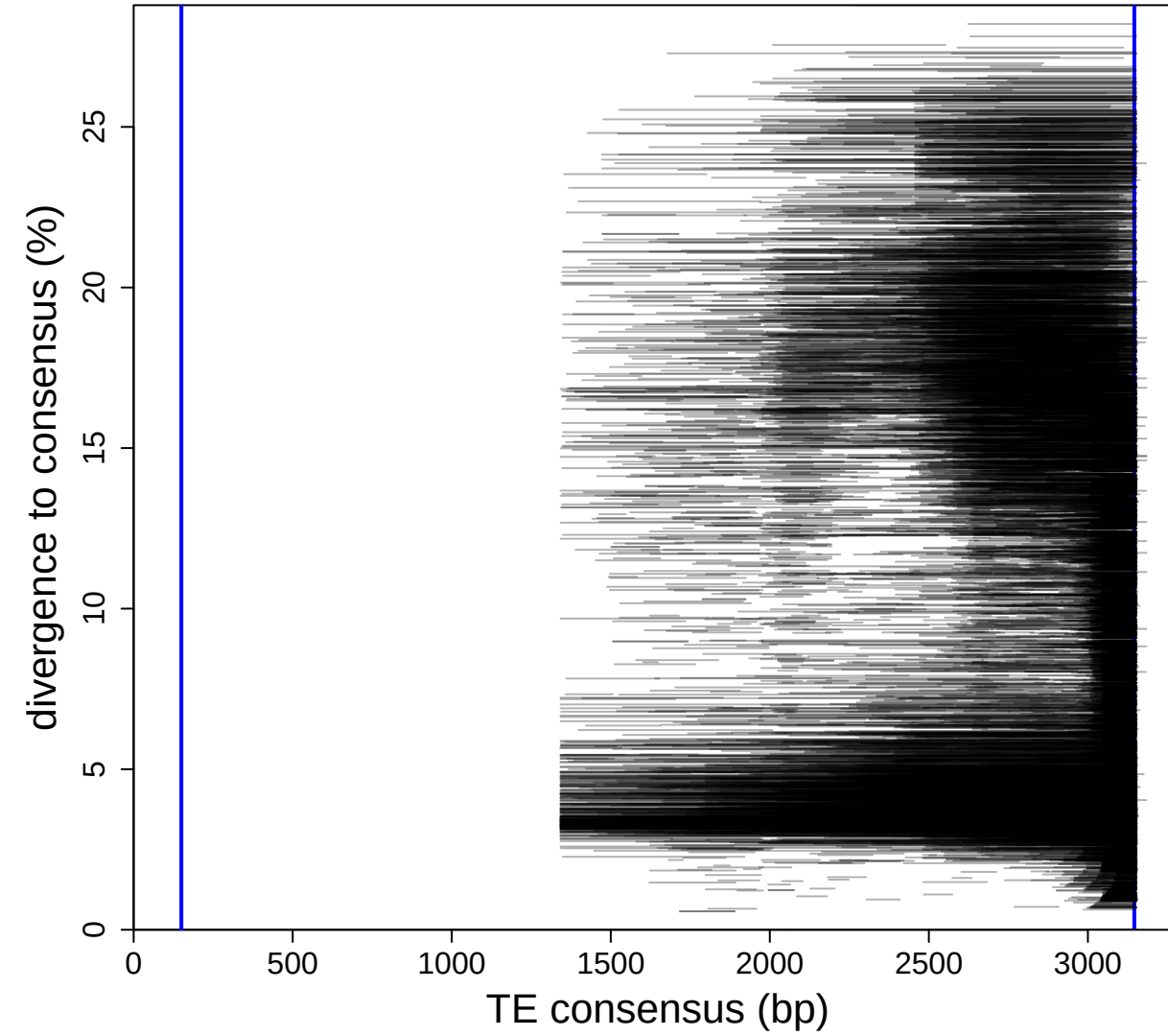
End crop Point





f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 3296bp; fragments: 12430; full length: 0 (>=2966.4bp)

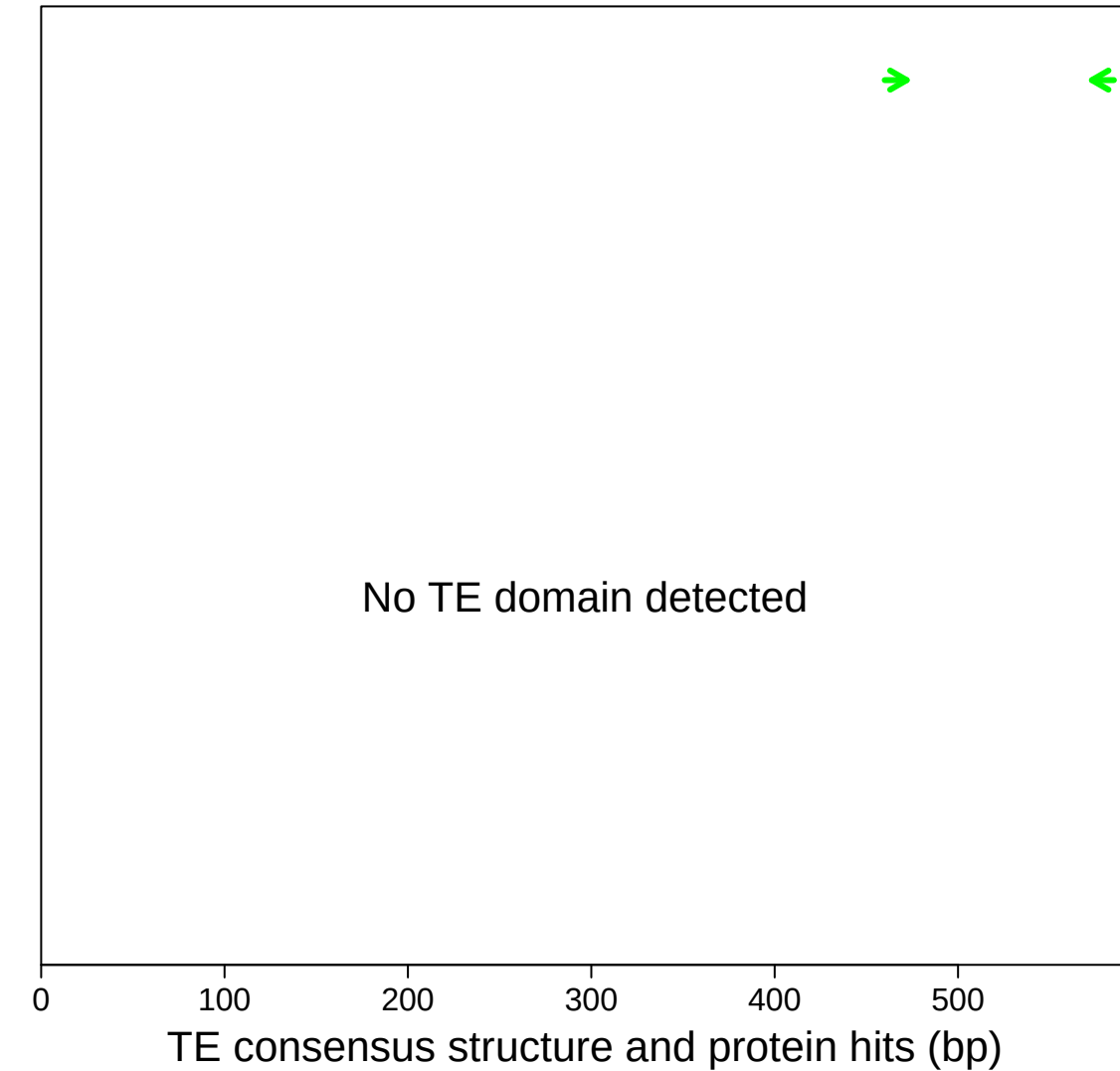
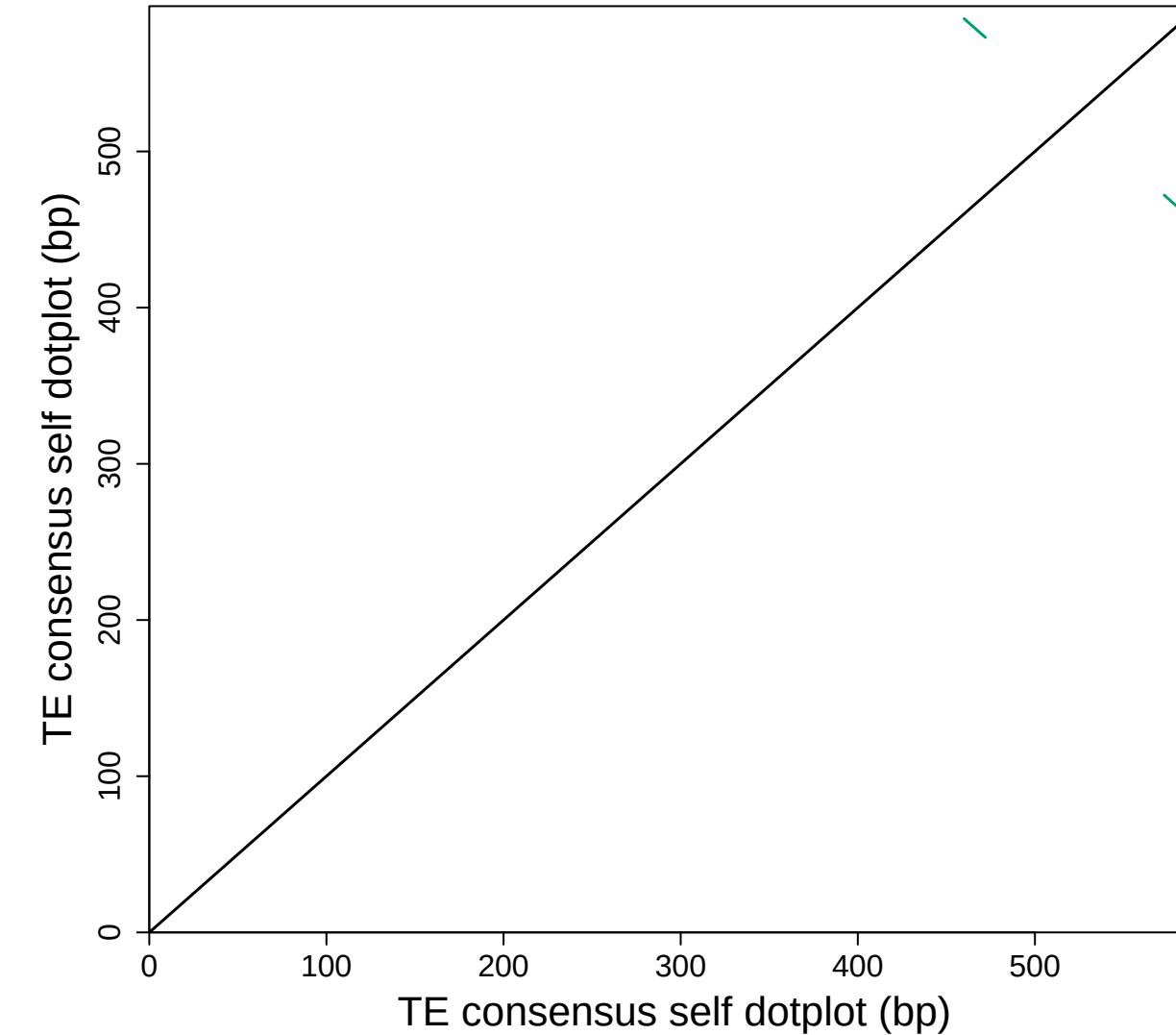
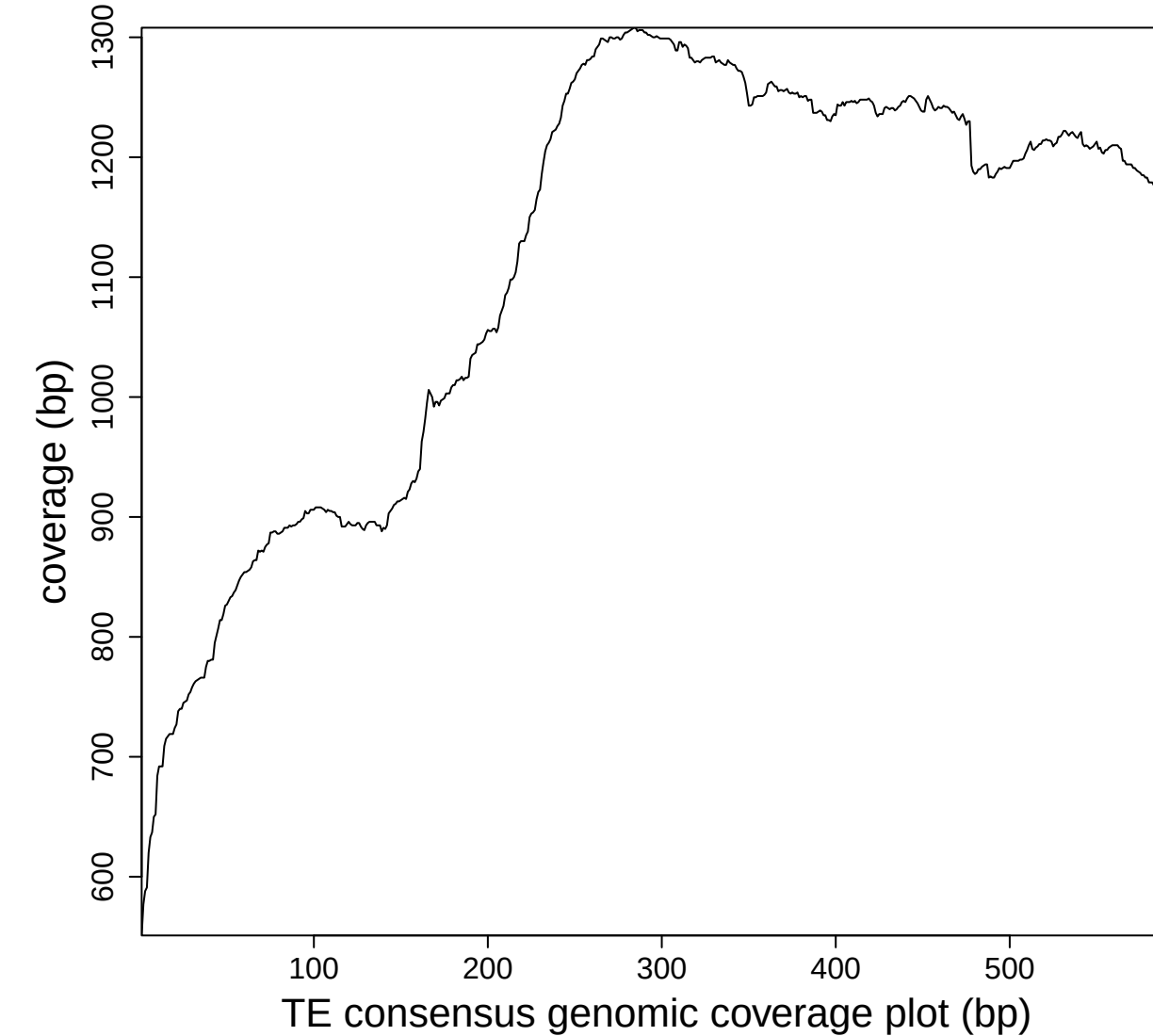
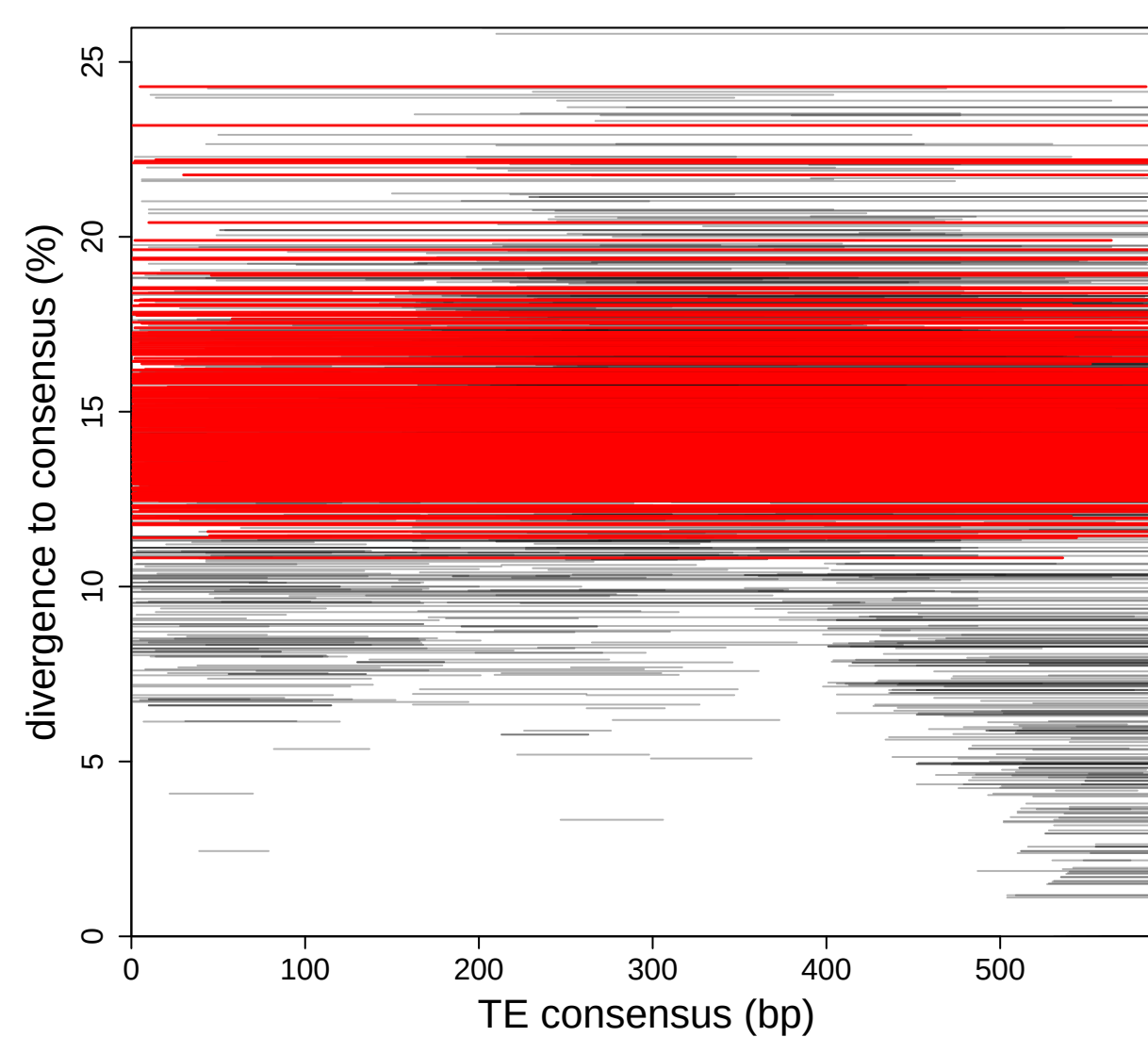
After TEtrimmer Extended plot Blue lines are boundaries



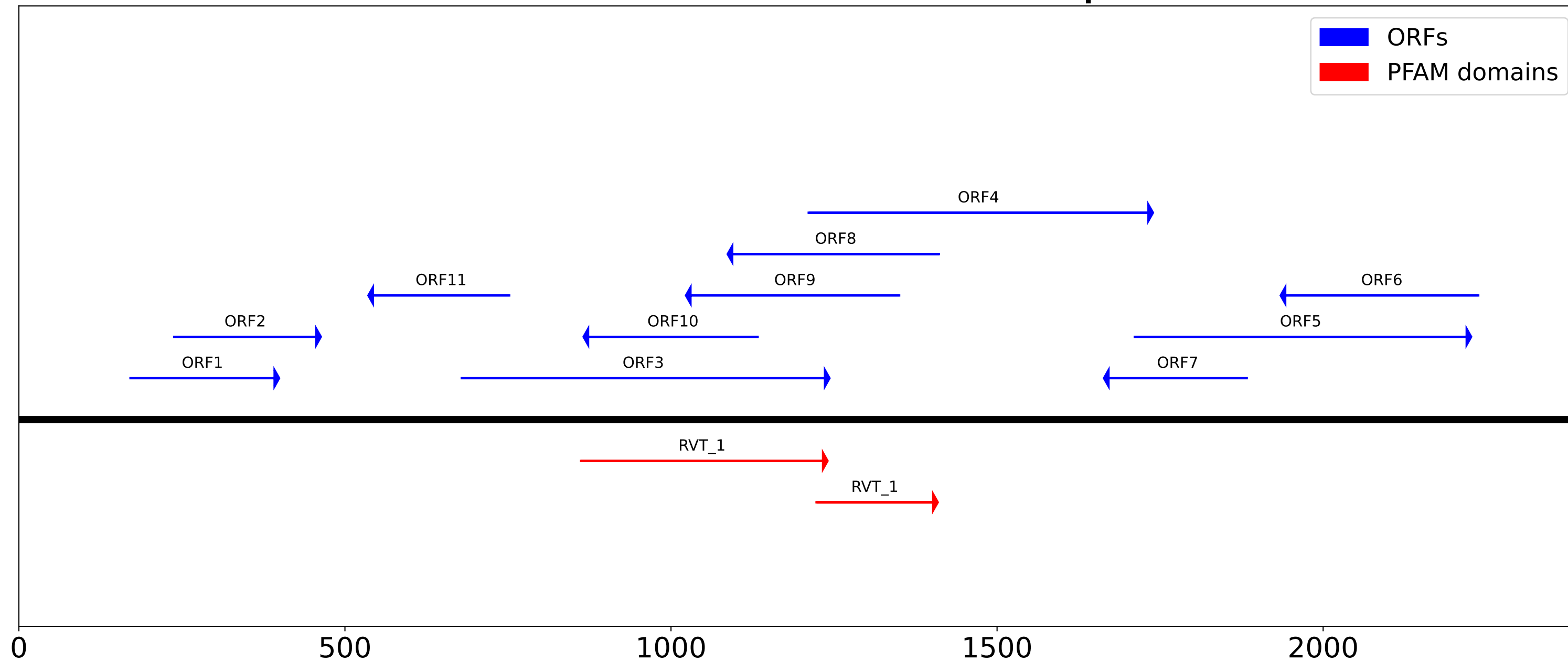
Before TEtrimmer 593 bp

TE: rnd_1_family_208

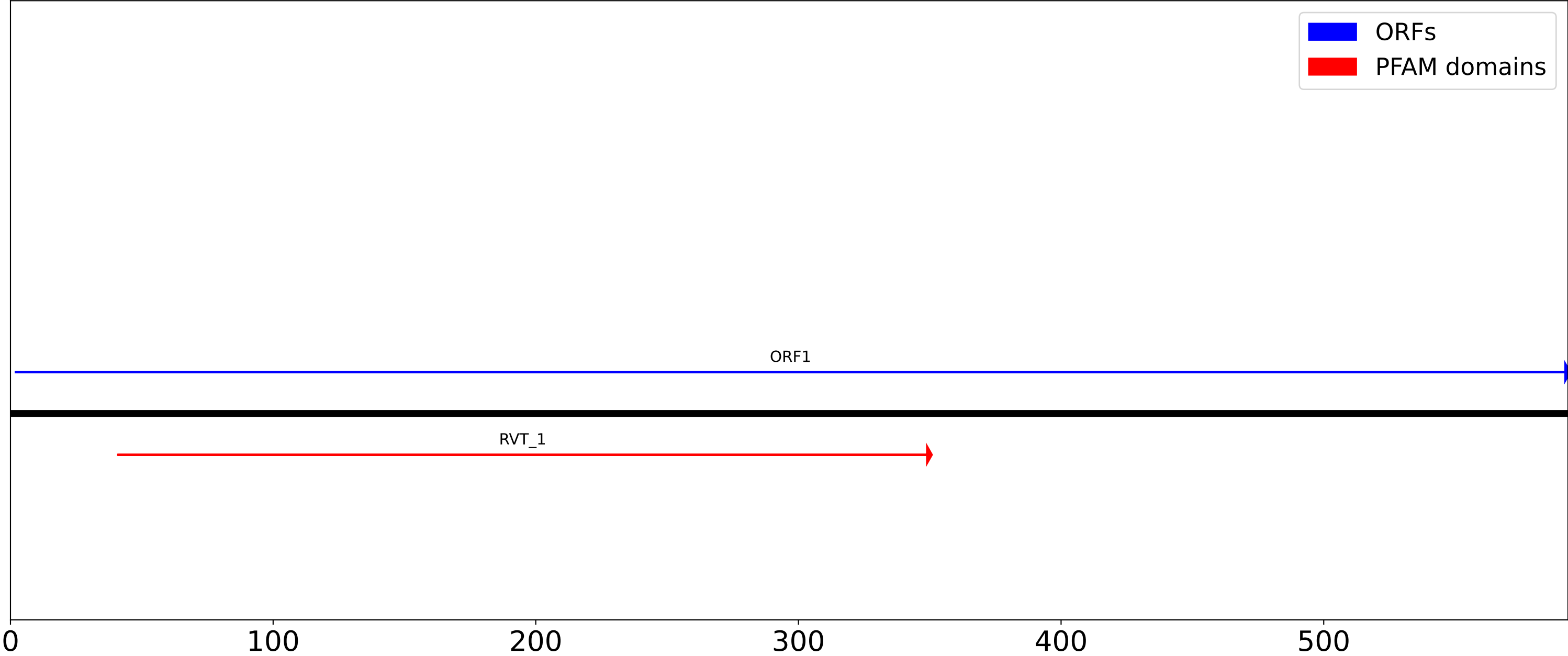
size: 593bp; fragments: 1811; full length: 543 (>=533.7bp)



After TEtrimmer ORF and PFAM domain plot



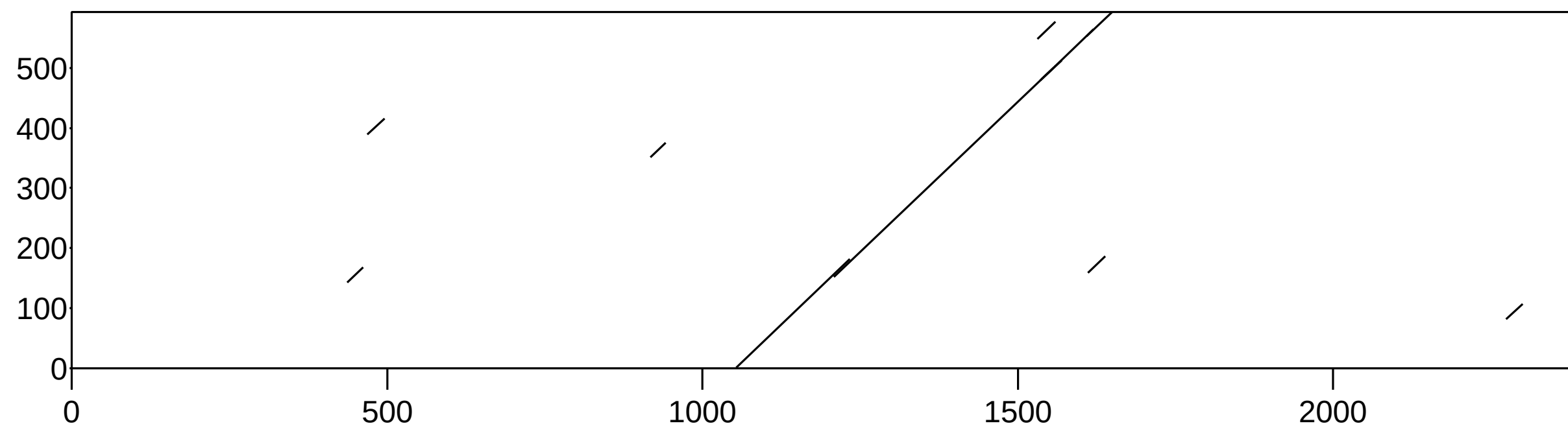
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)