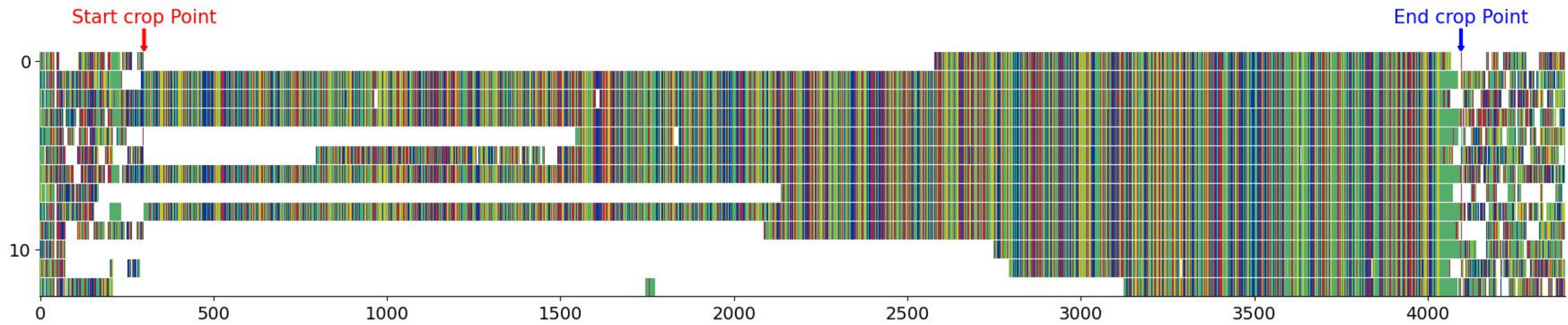


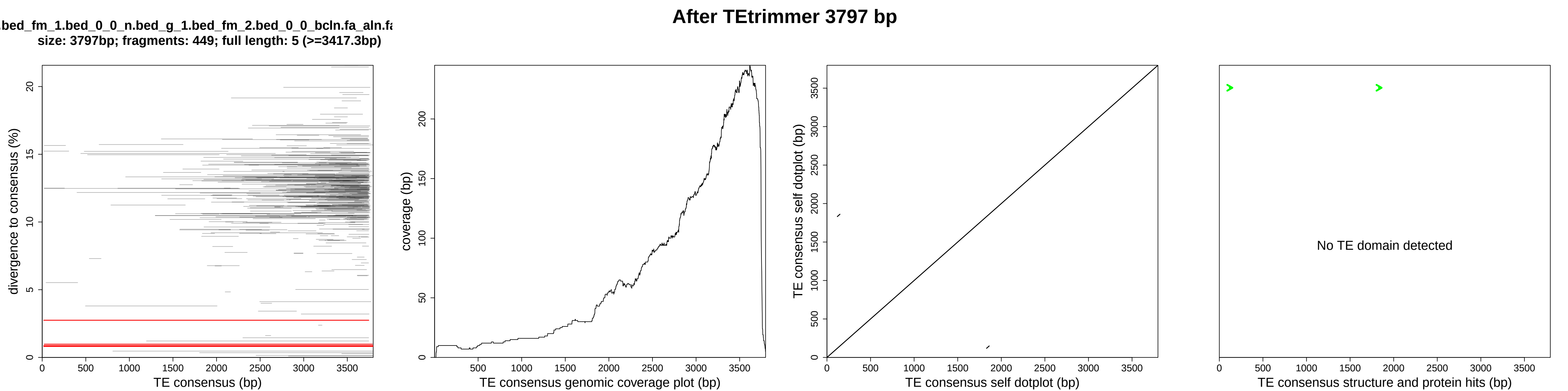
Start crop Point

End crop Point

MSA length = 3797

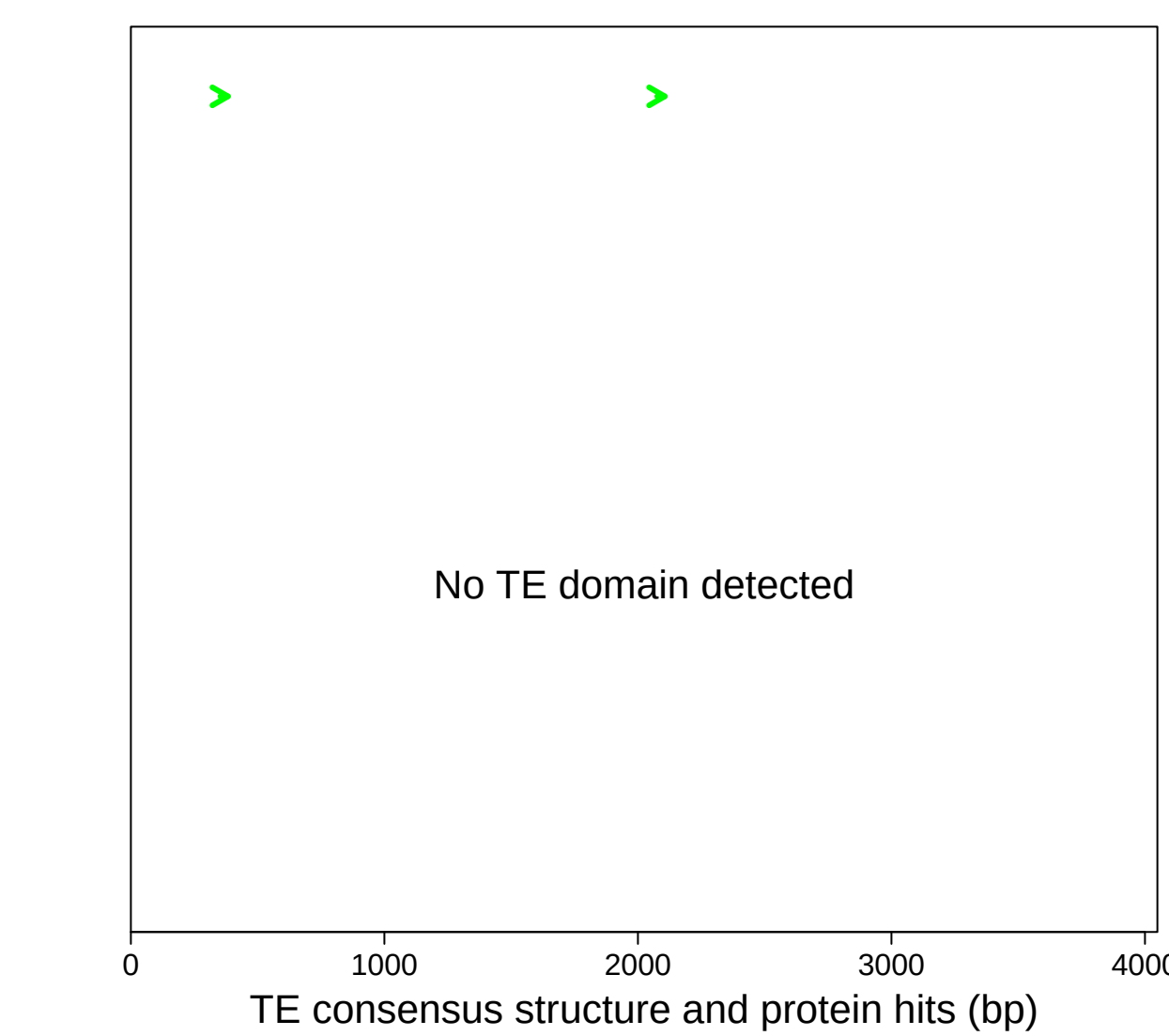
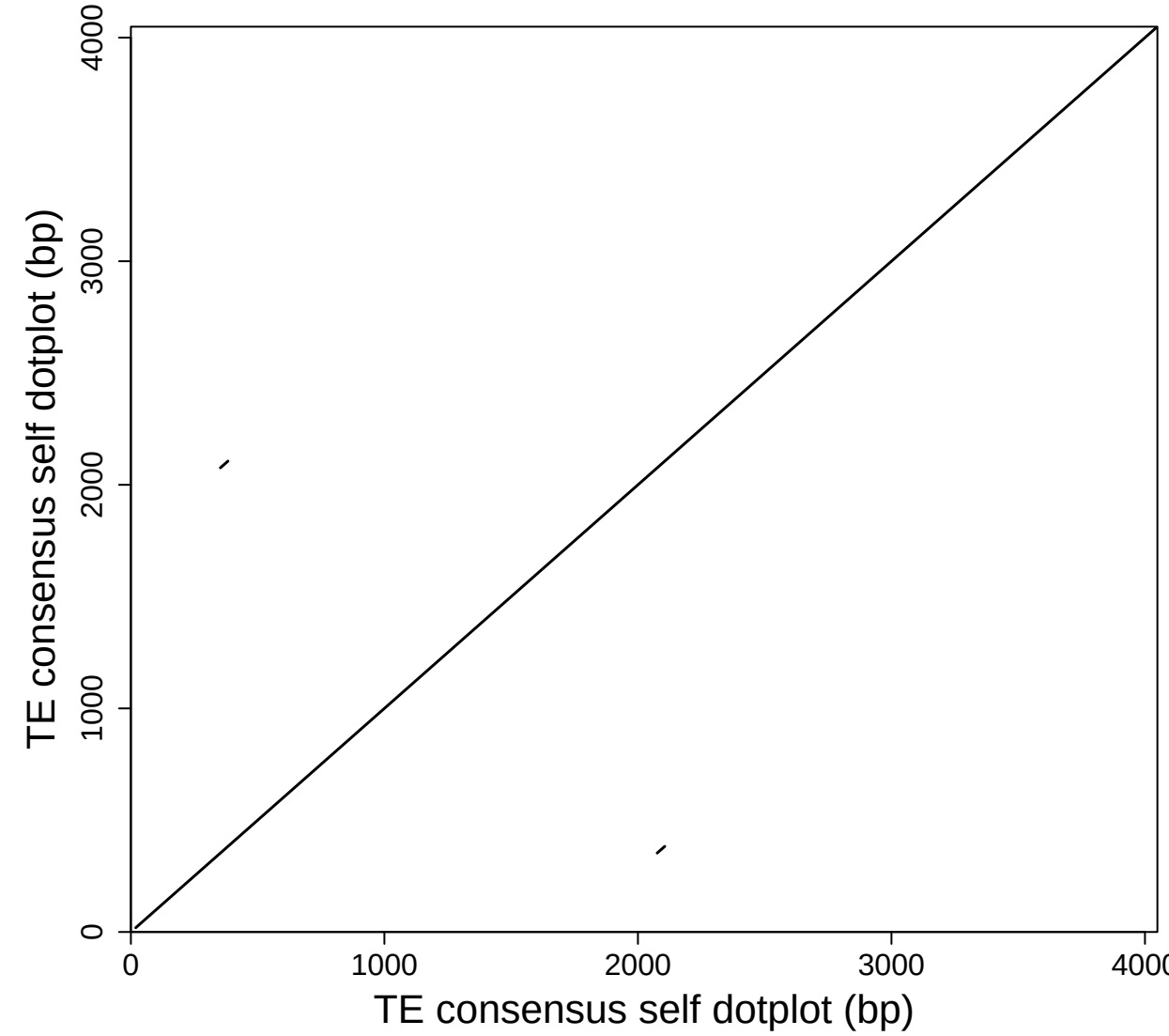
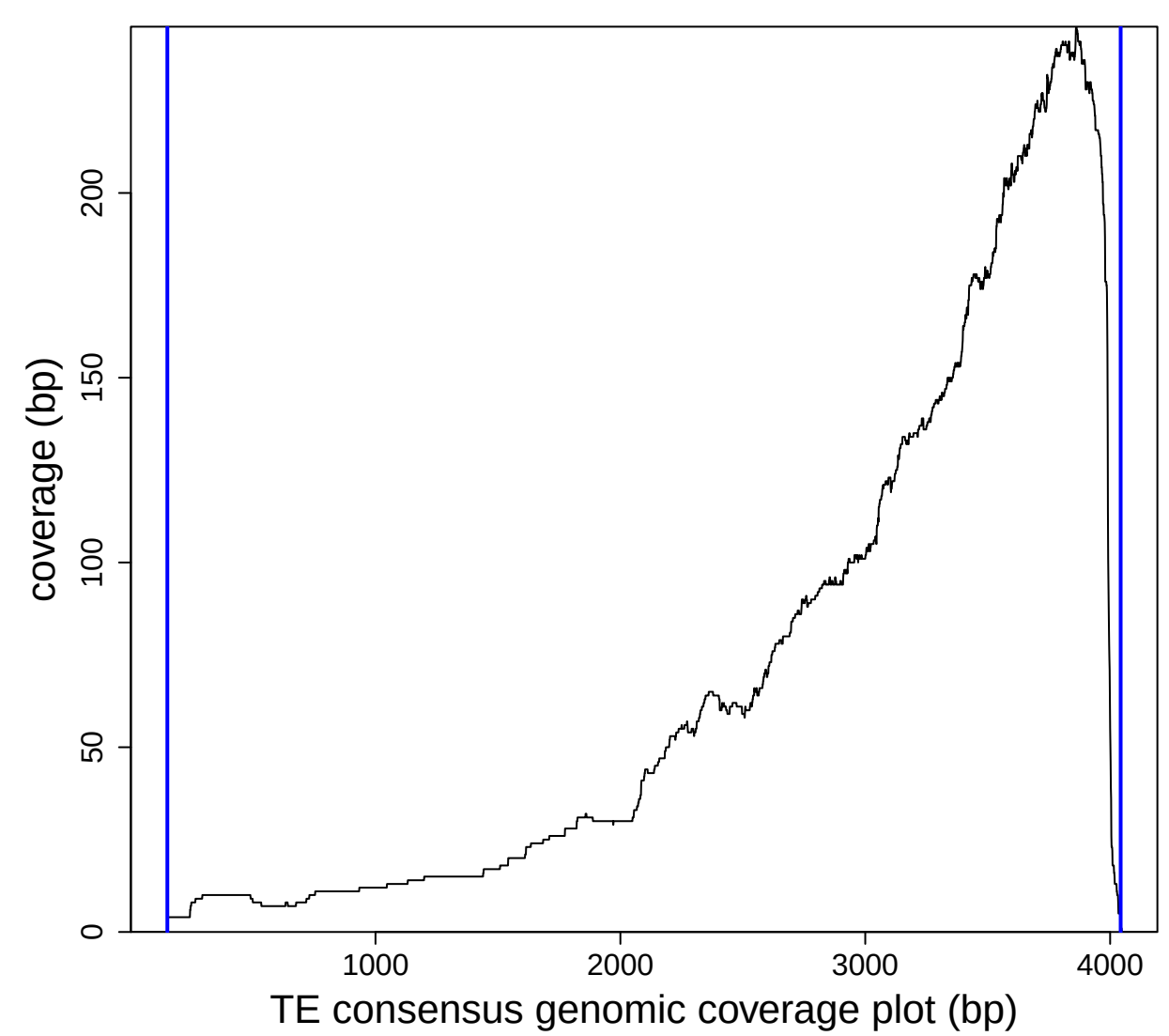
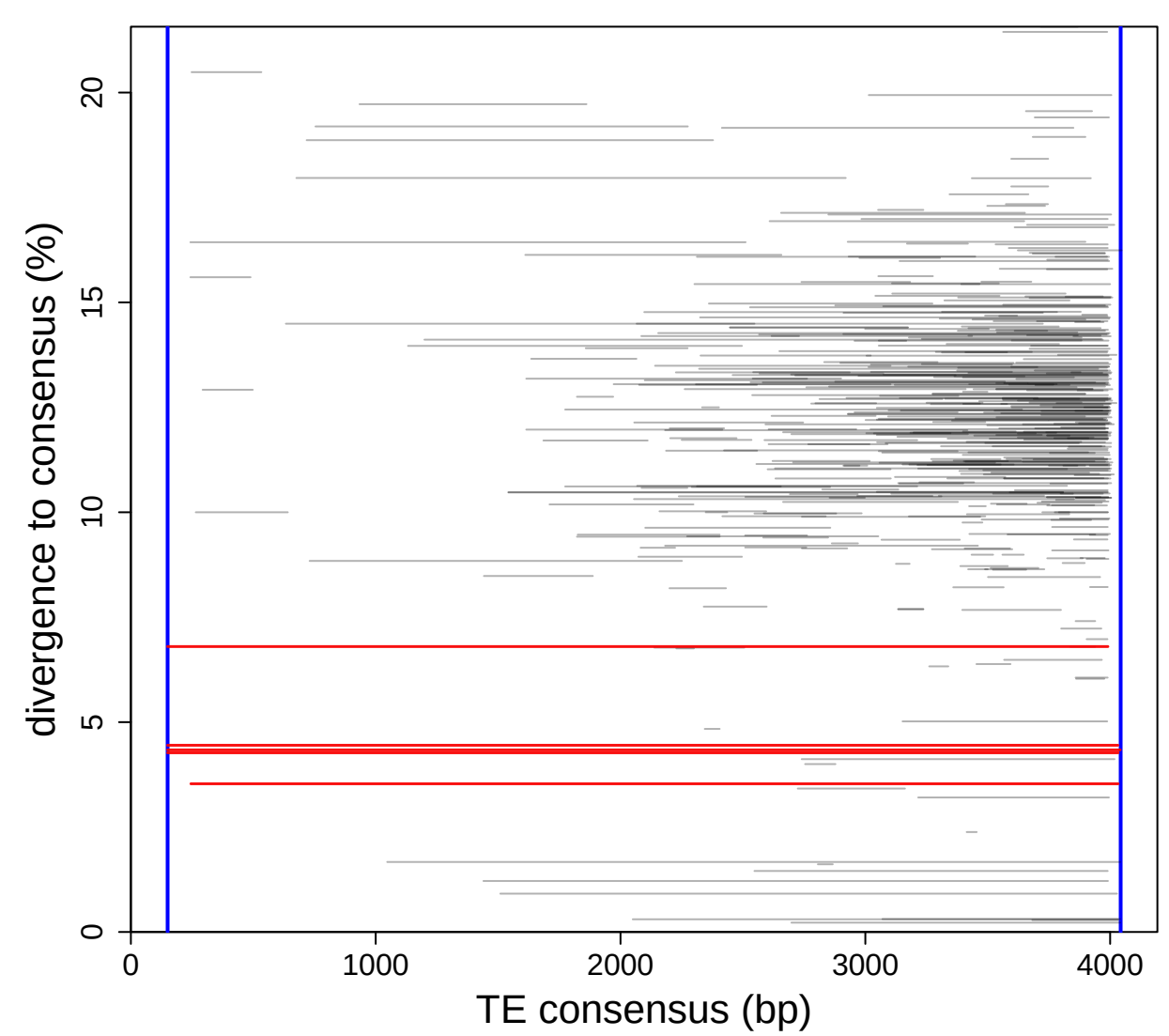




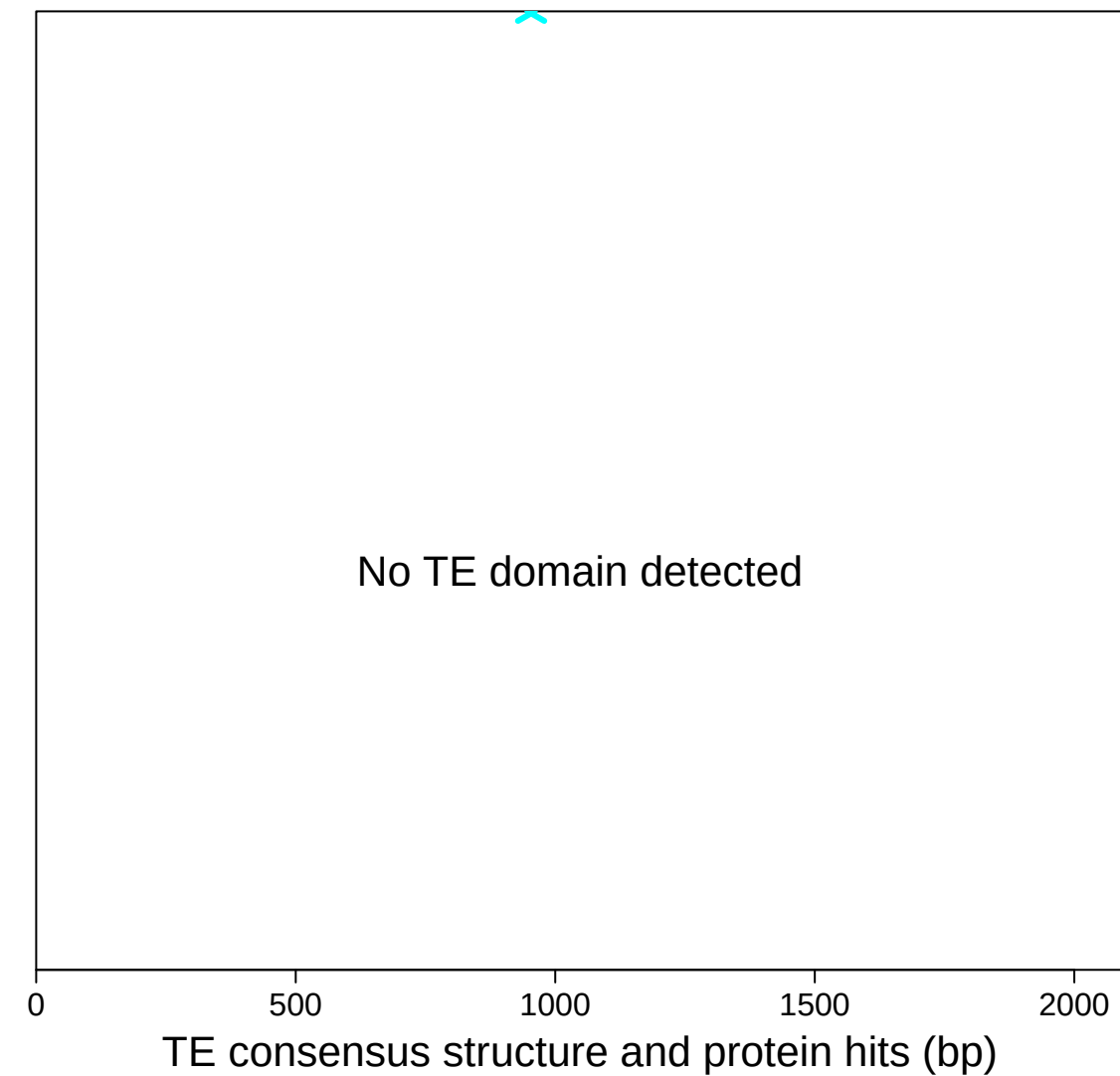
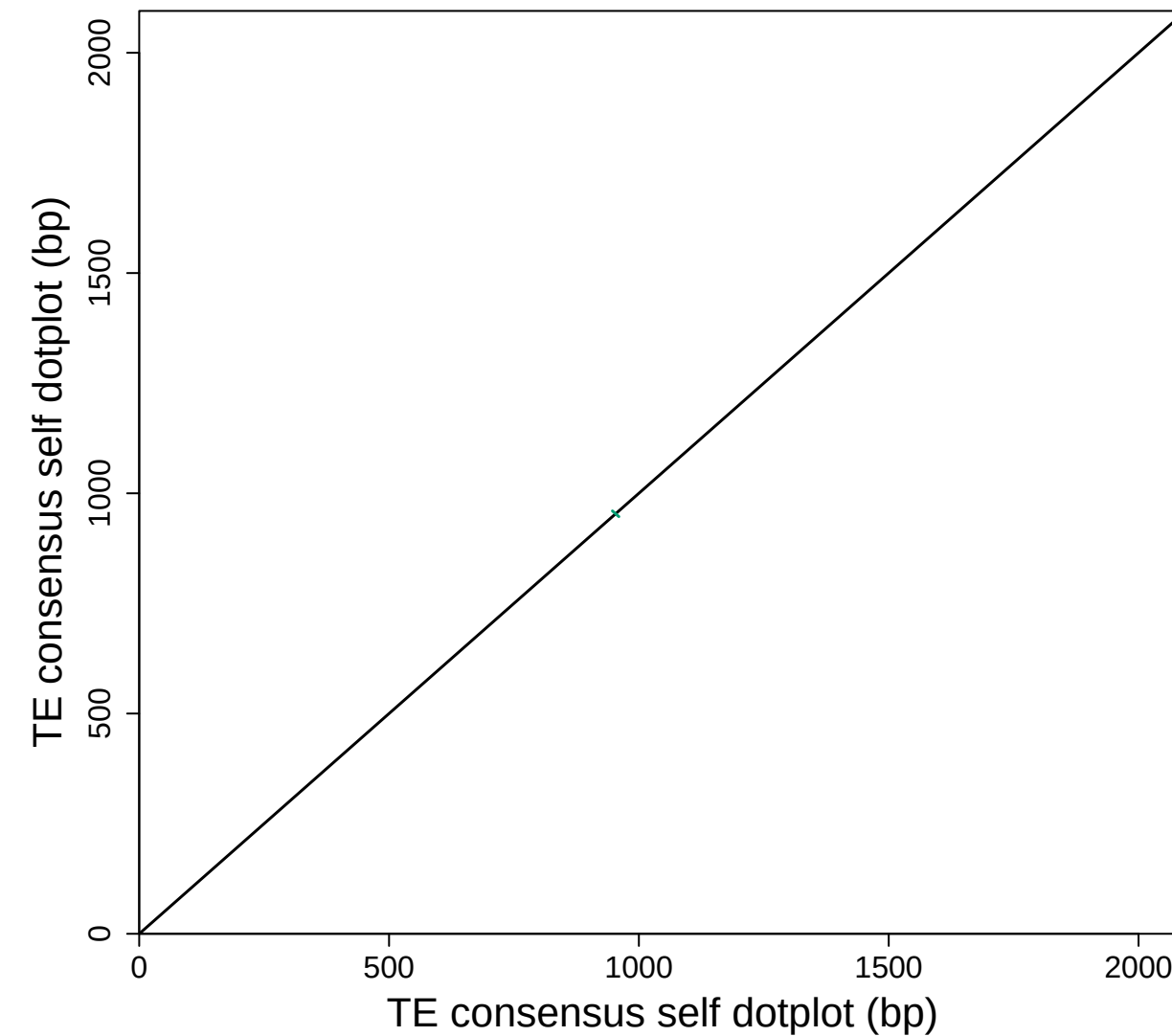
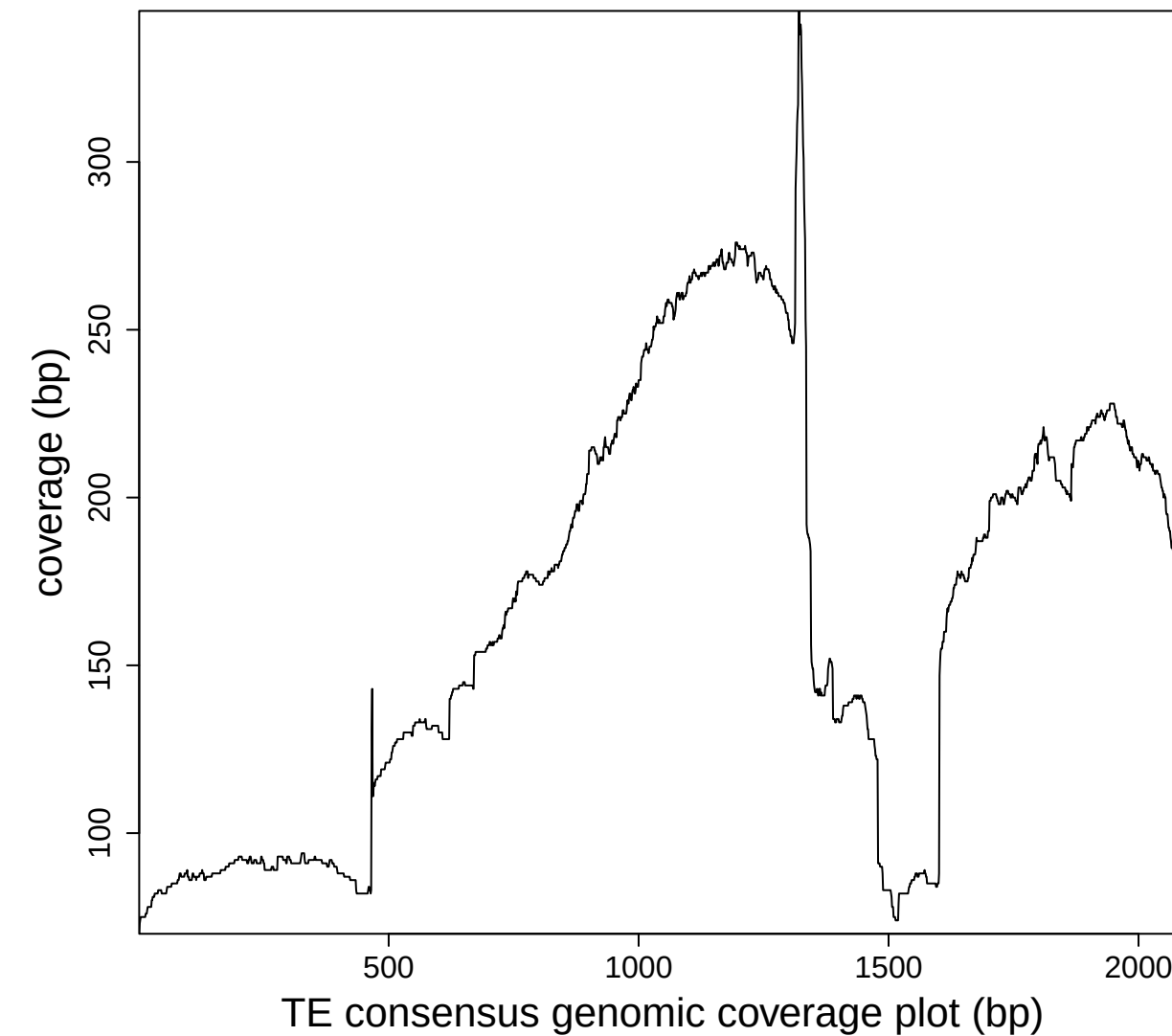
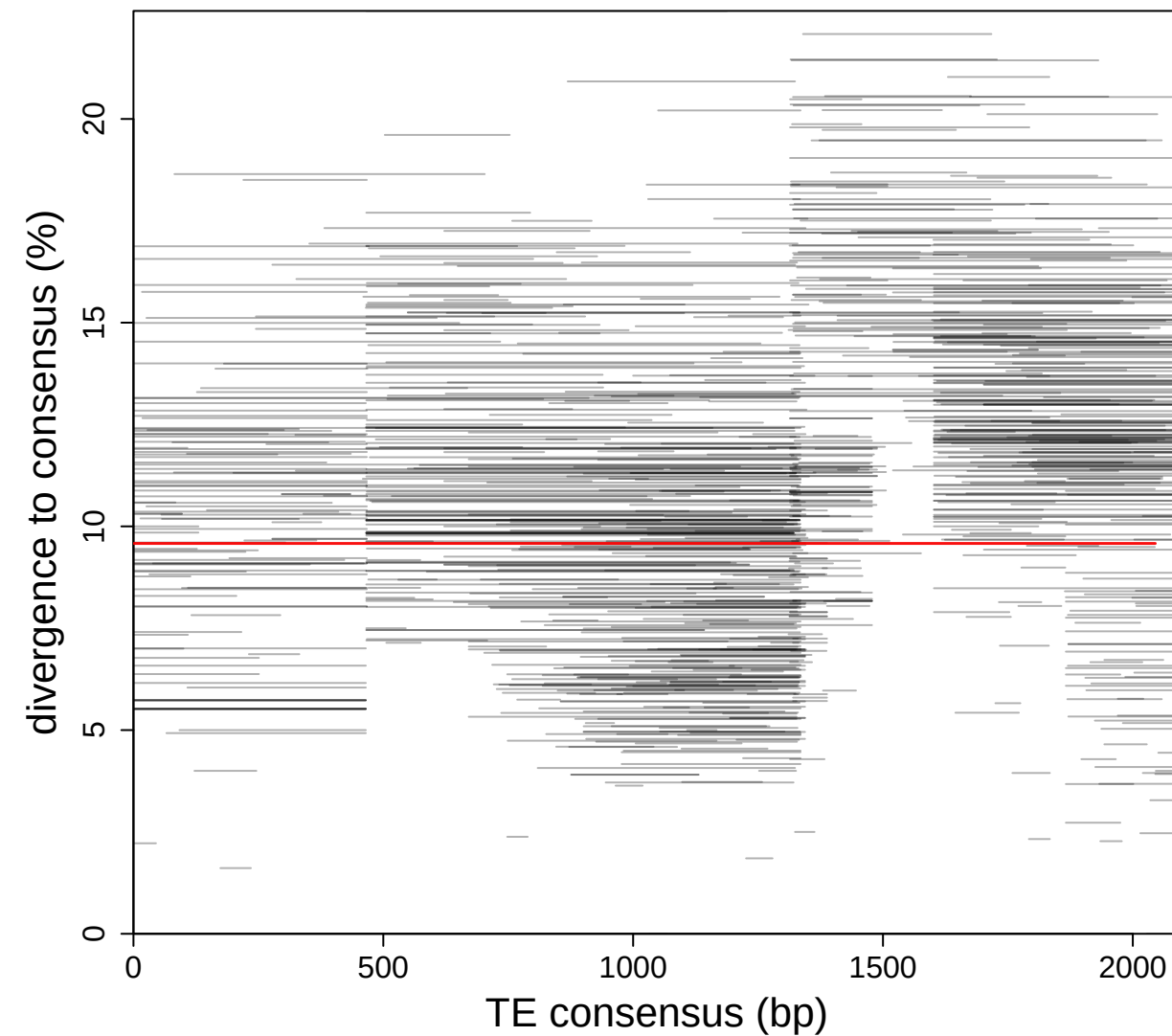


f.bed_g_3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 4193bp; fragments: 448; full length: 5 (>=3773.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

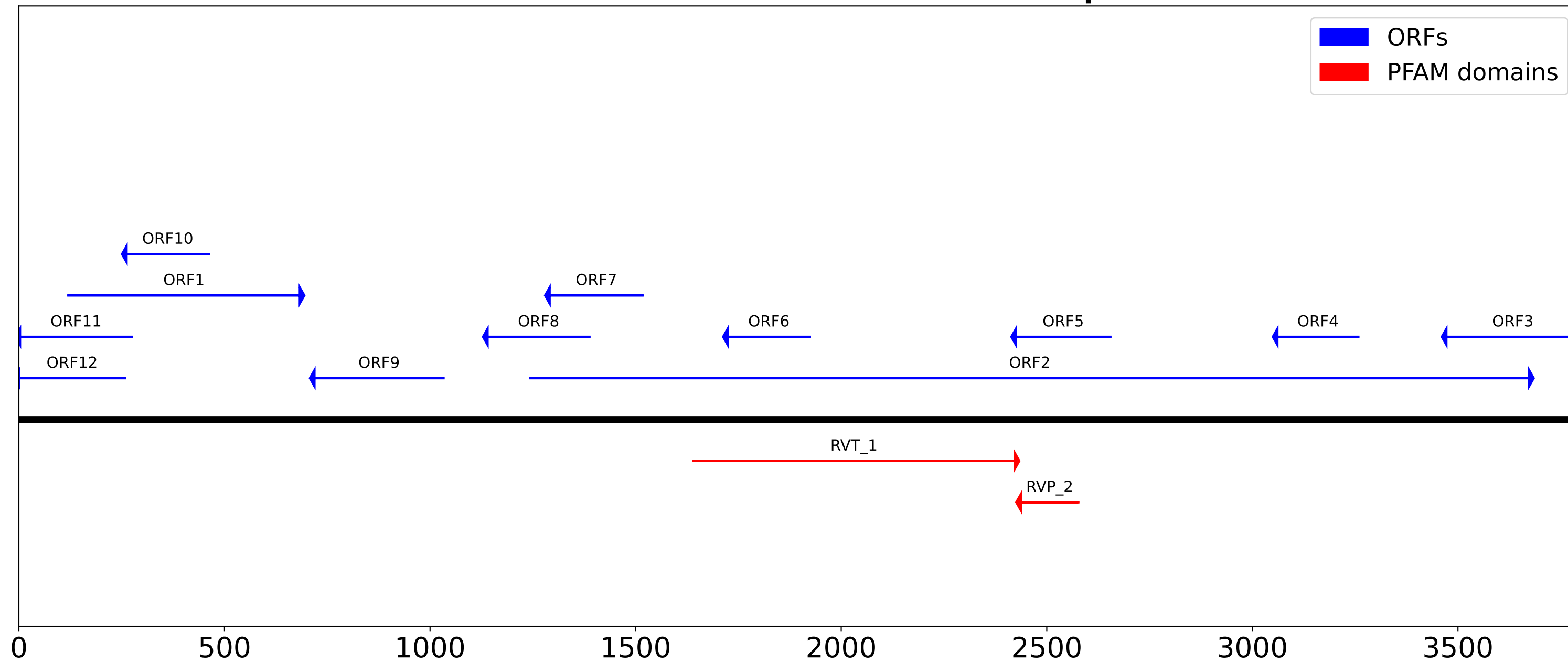


TE: rnd_4_family_594
size: 2095bp; fragments: 927; full length: 1 (≥ 1885.5 bp)

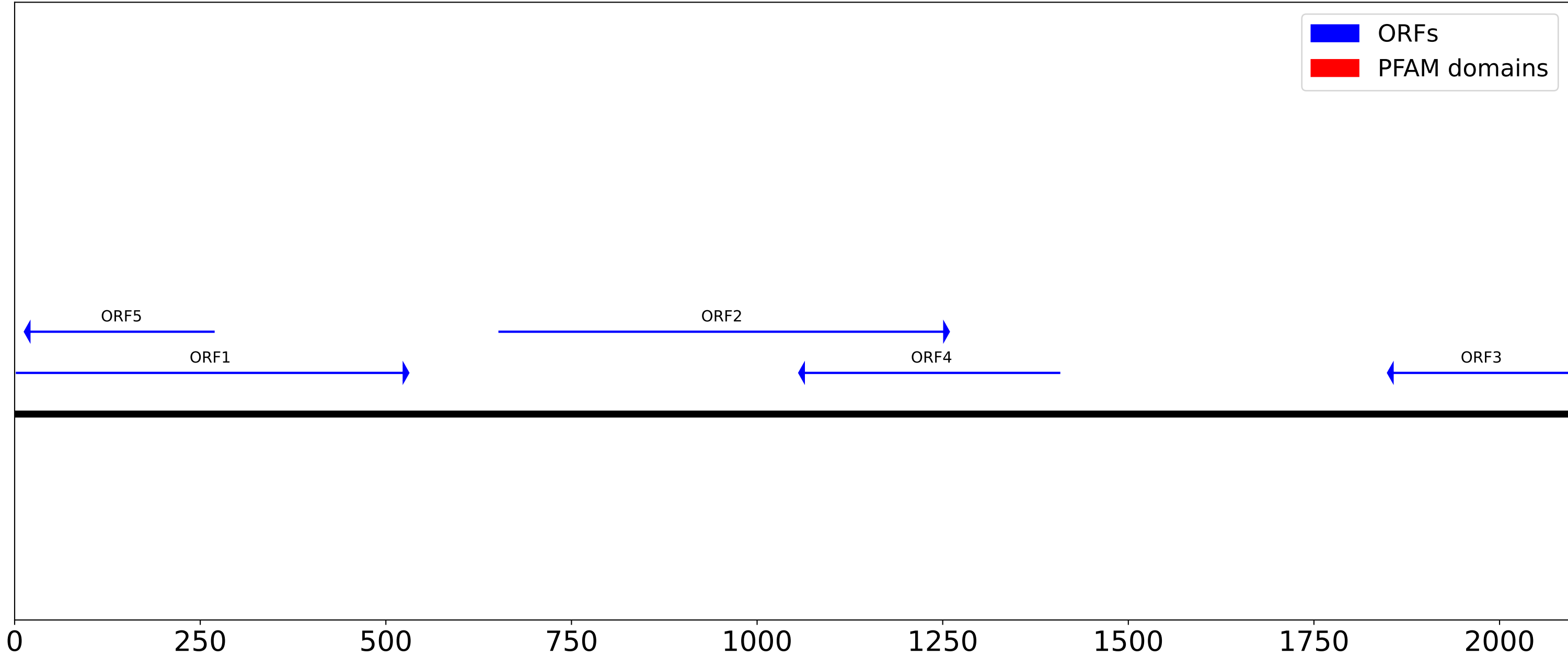


Before TETrimmer 2095 bp

After TEtrimmer ORF and PFAM domain plot



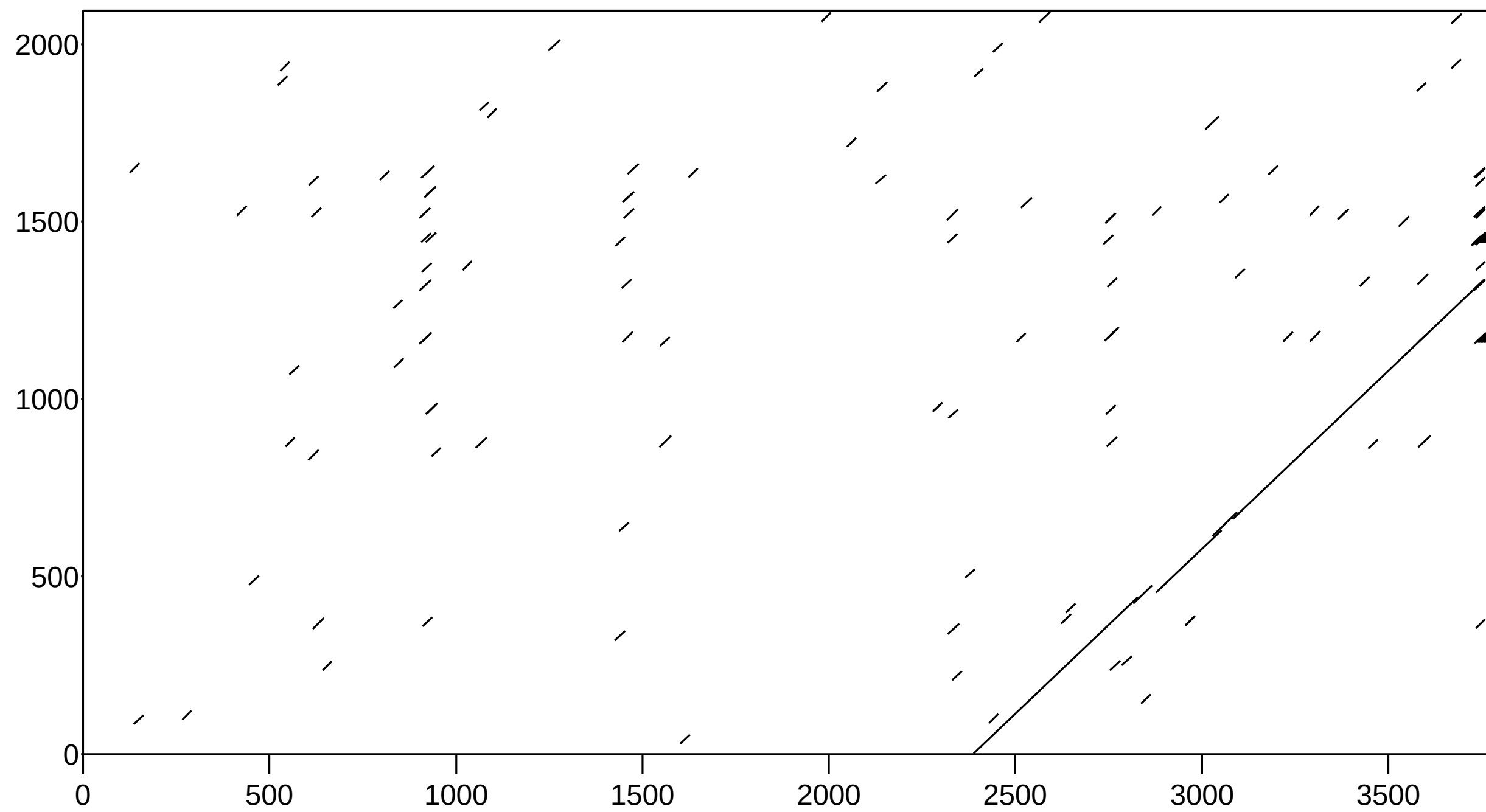
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)