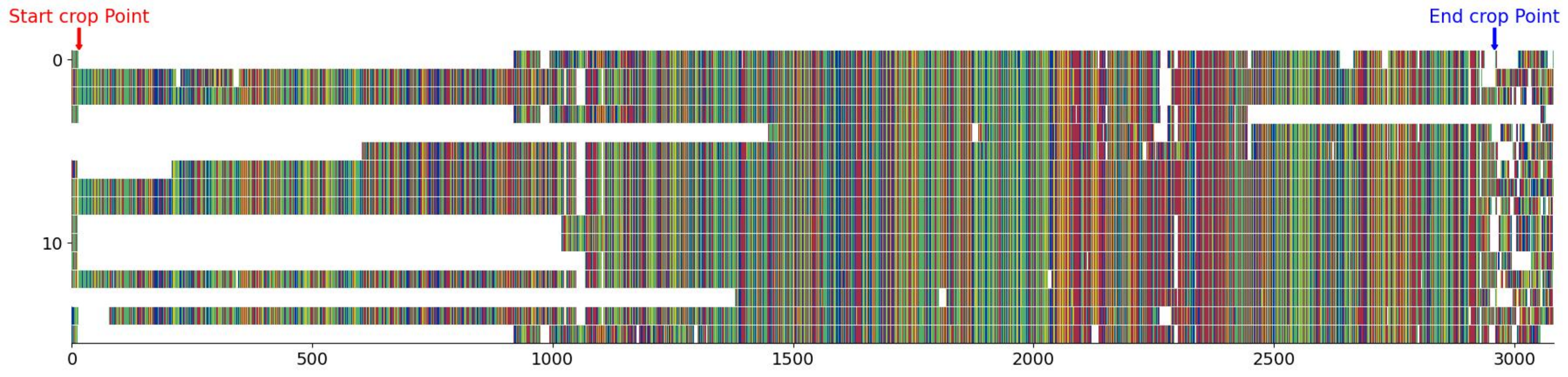


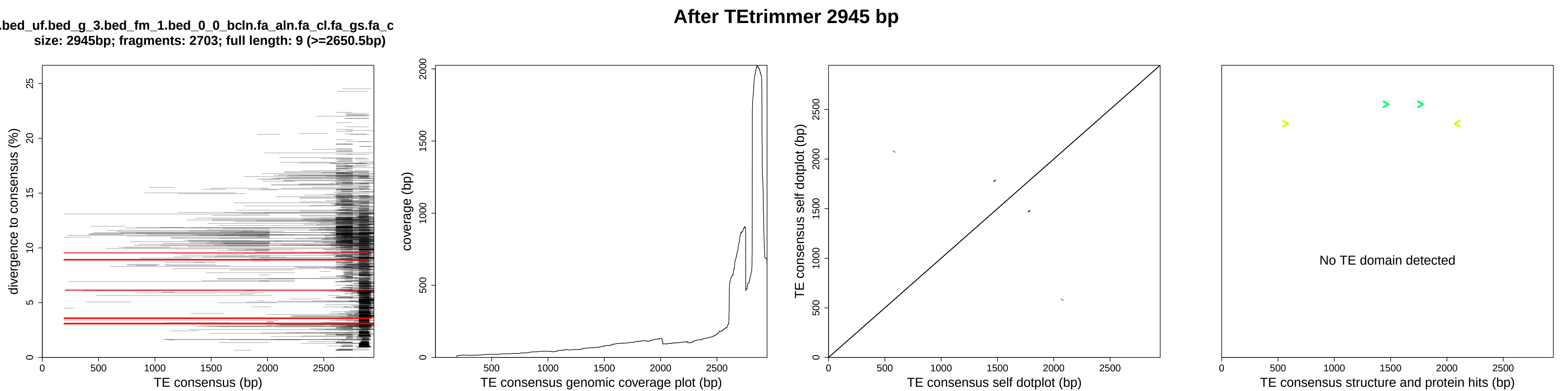
Start crop Point

End crop Point

MSA length = 2945

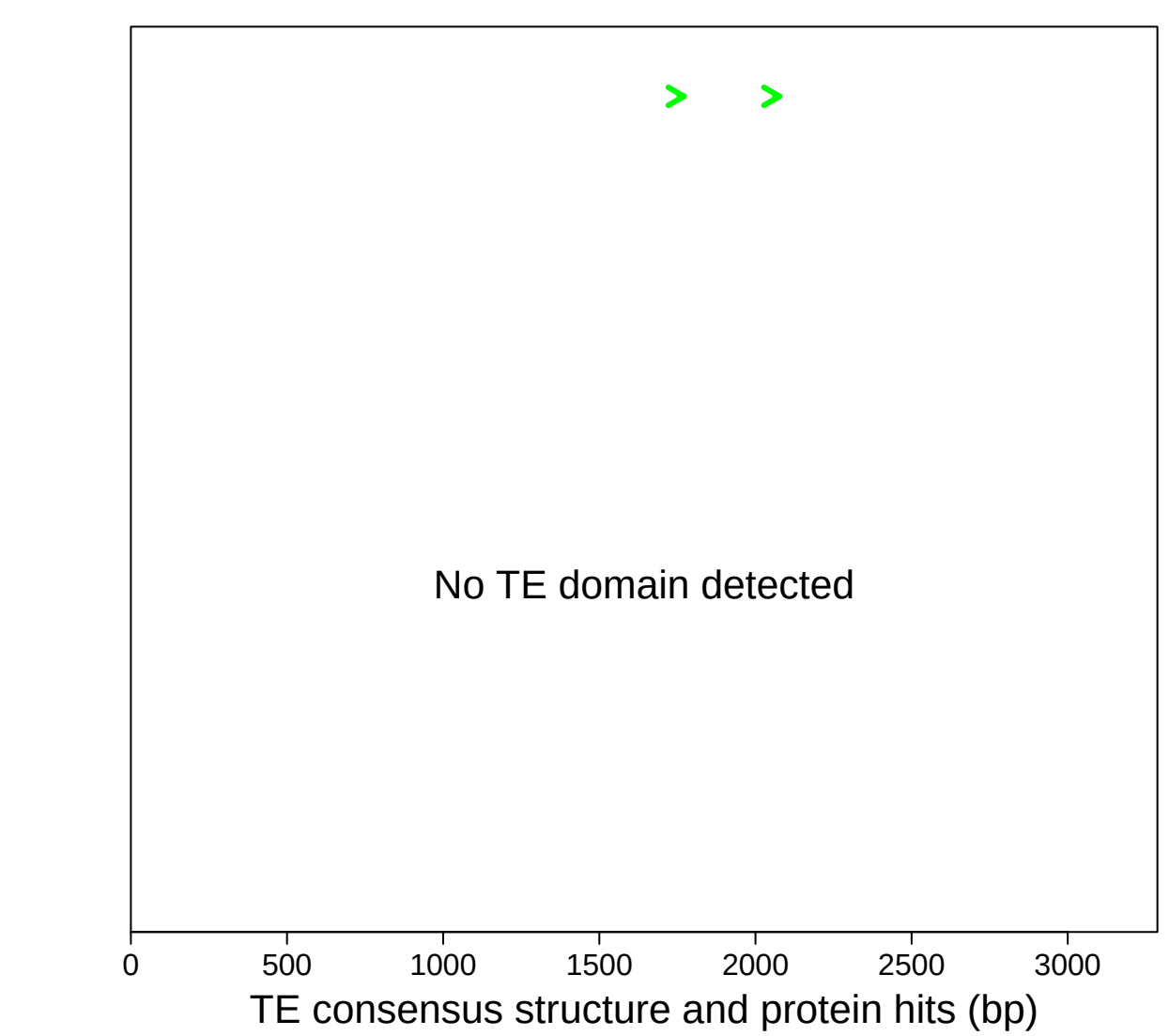
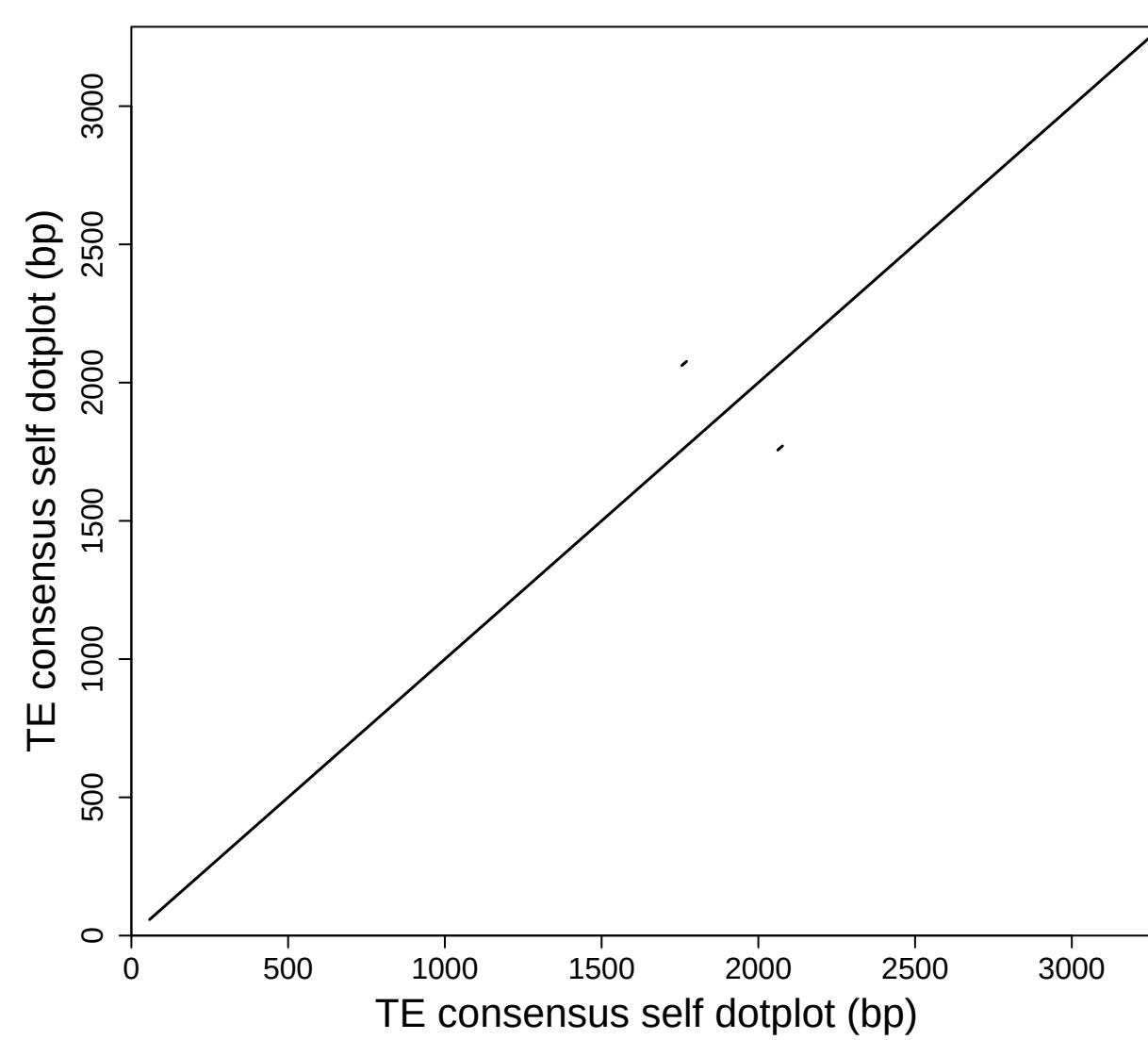
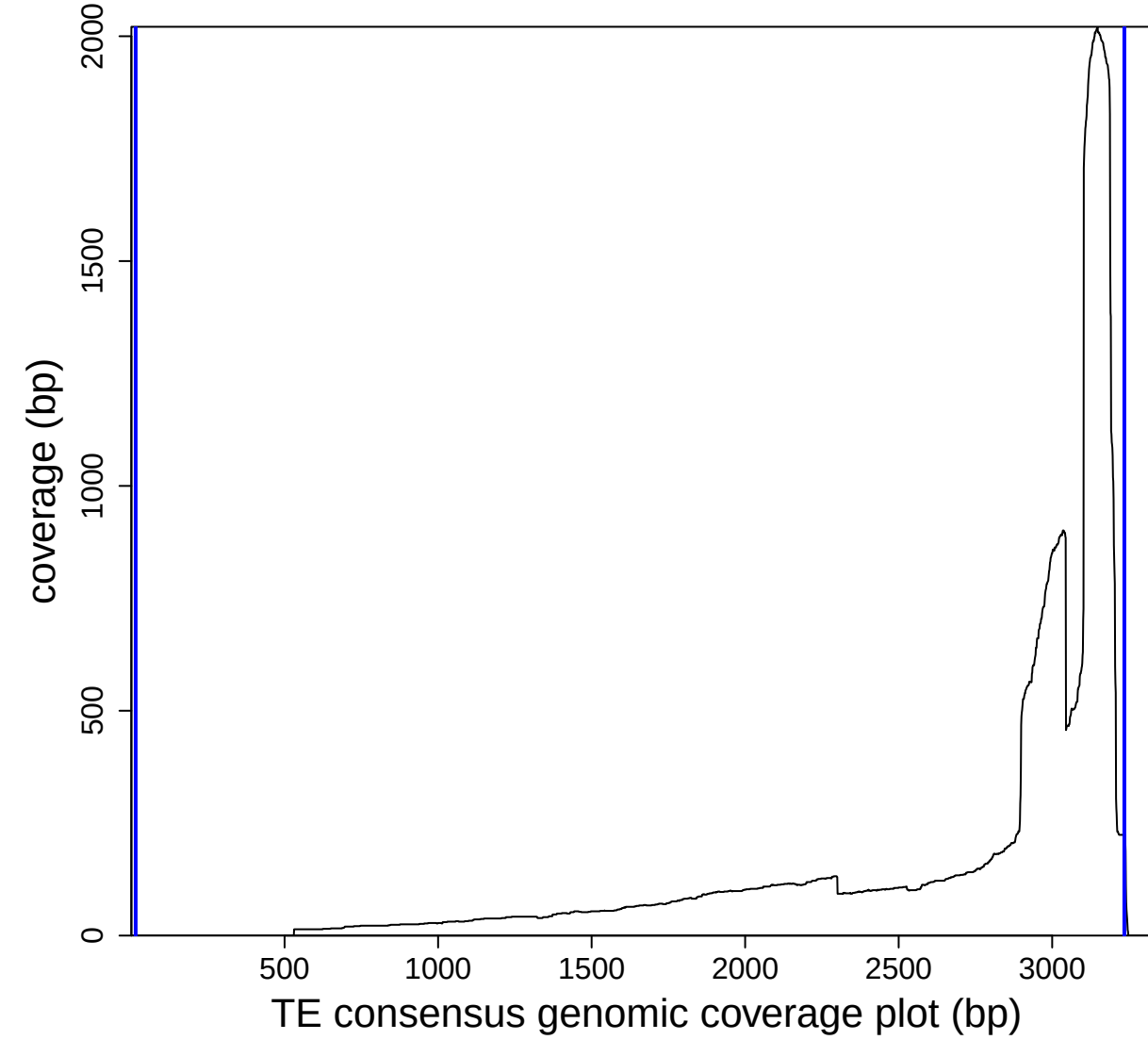
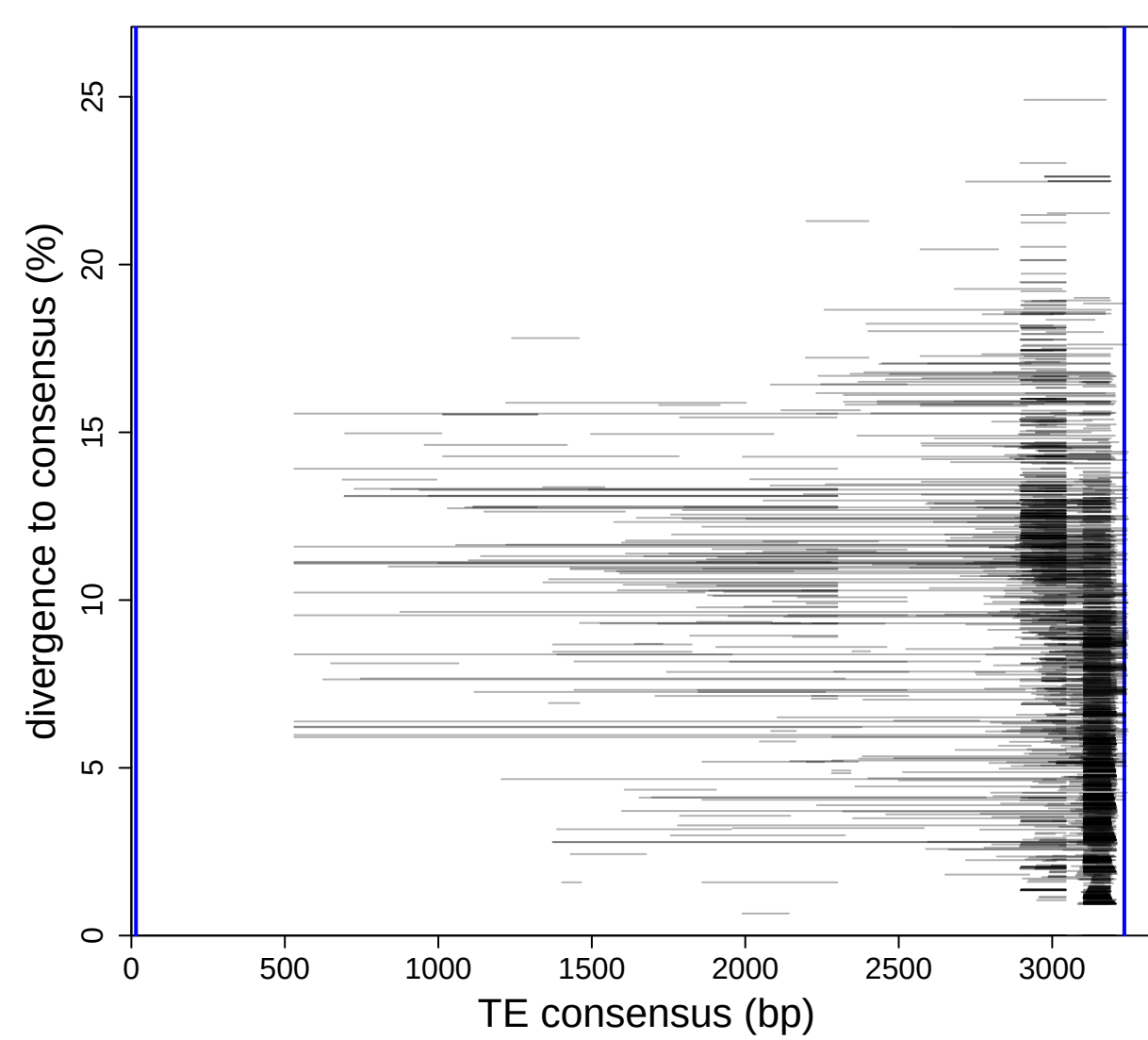






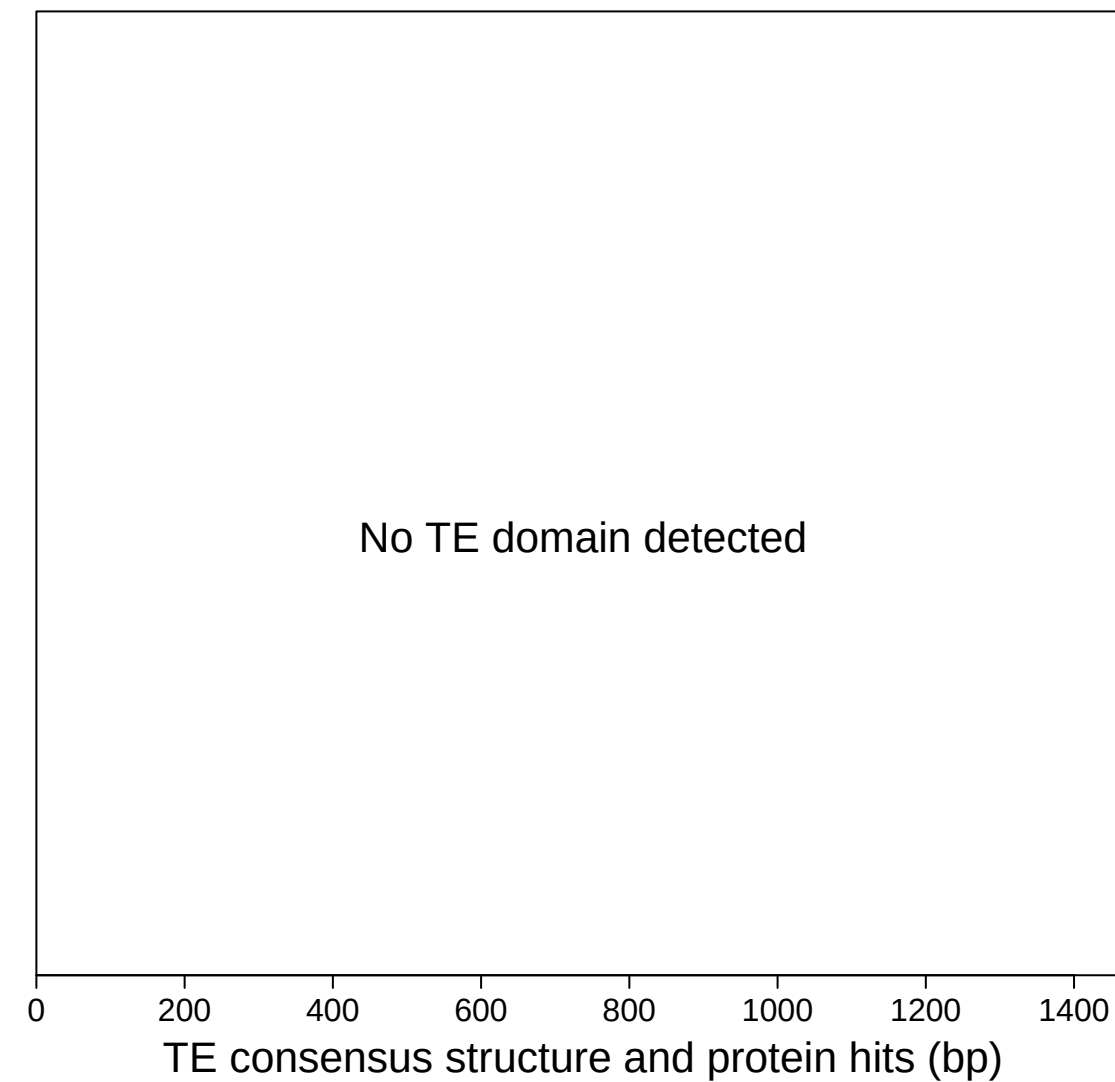
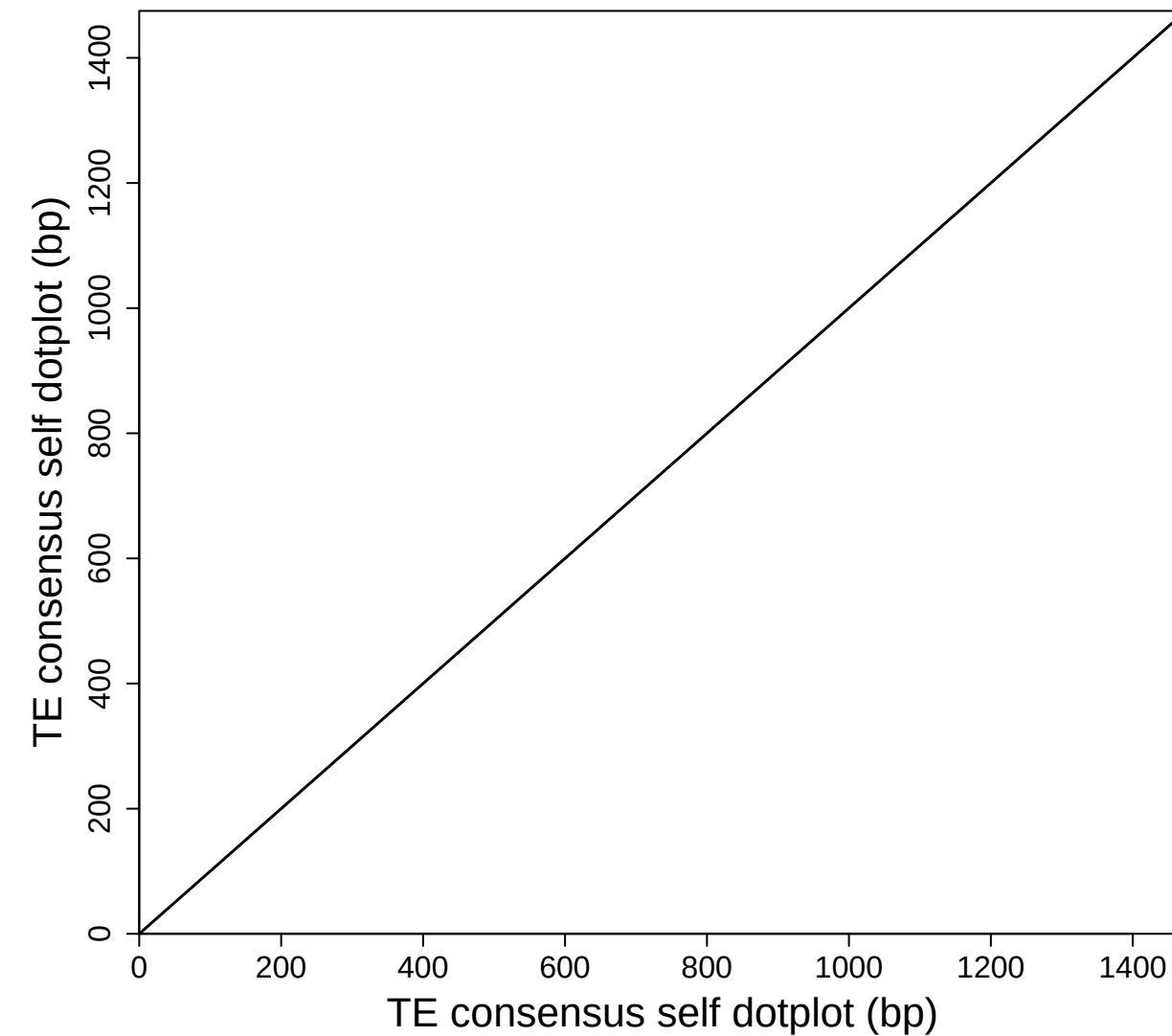
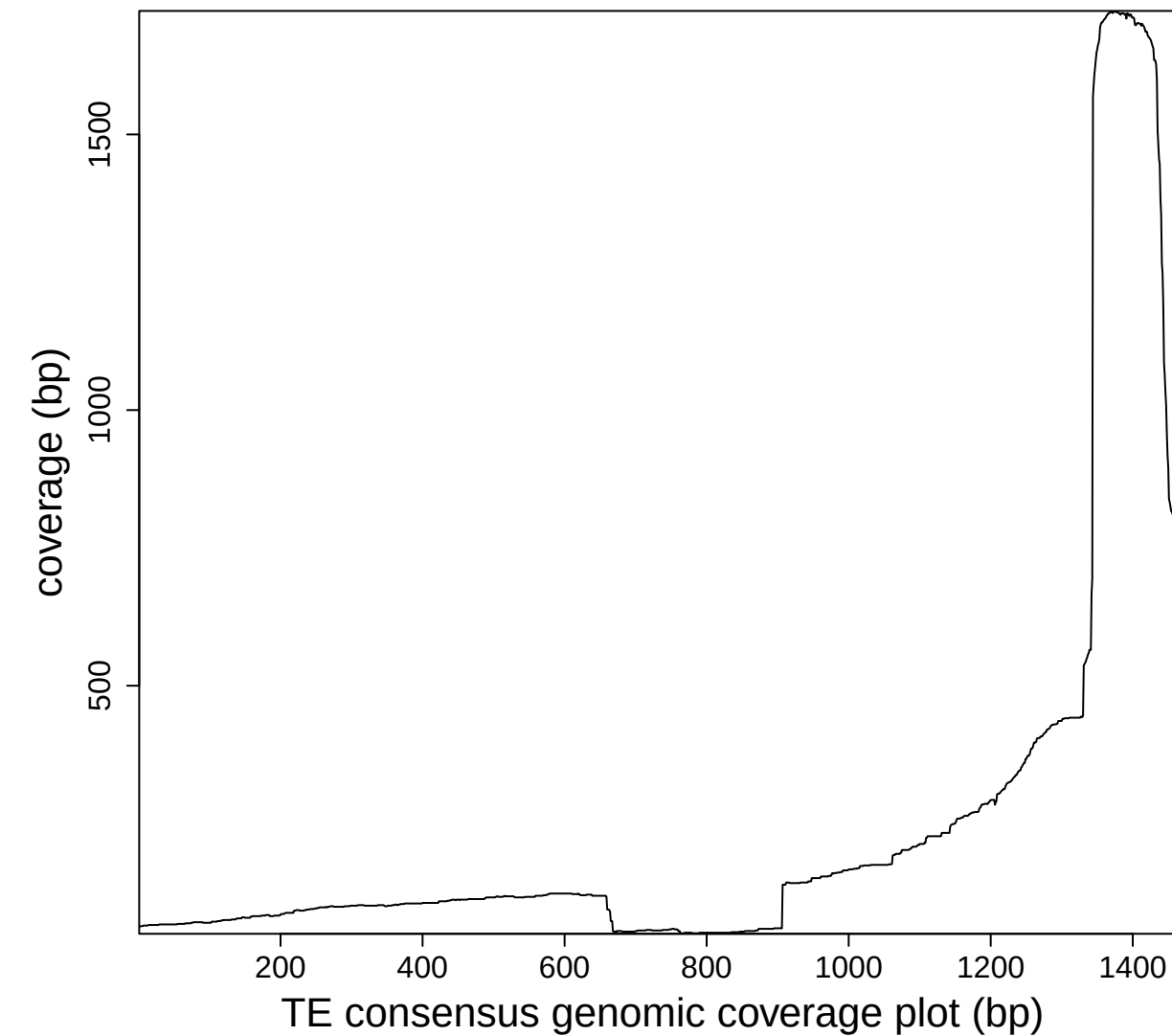
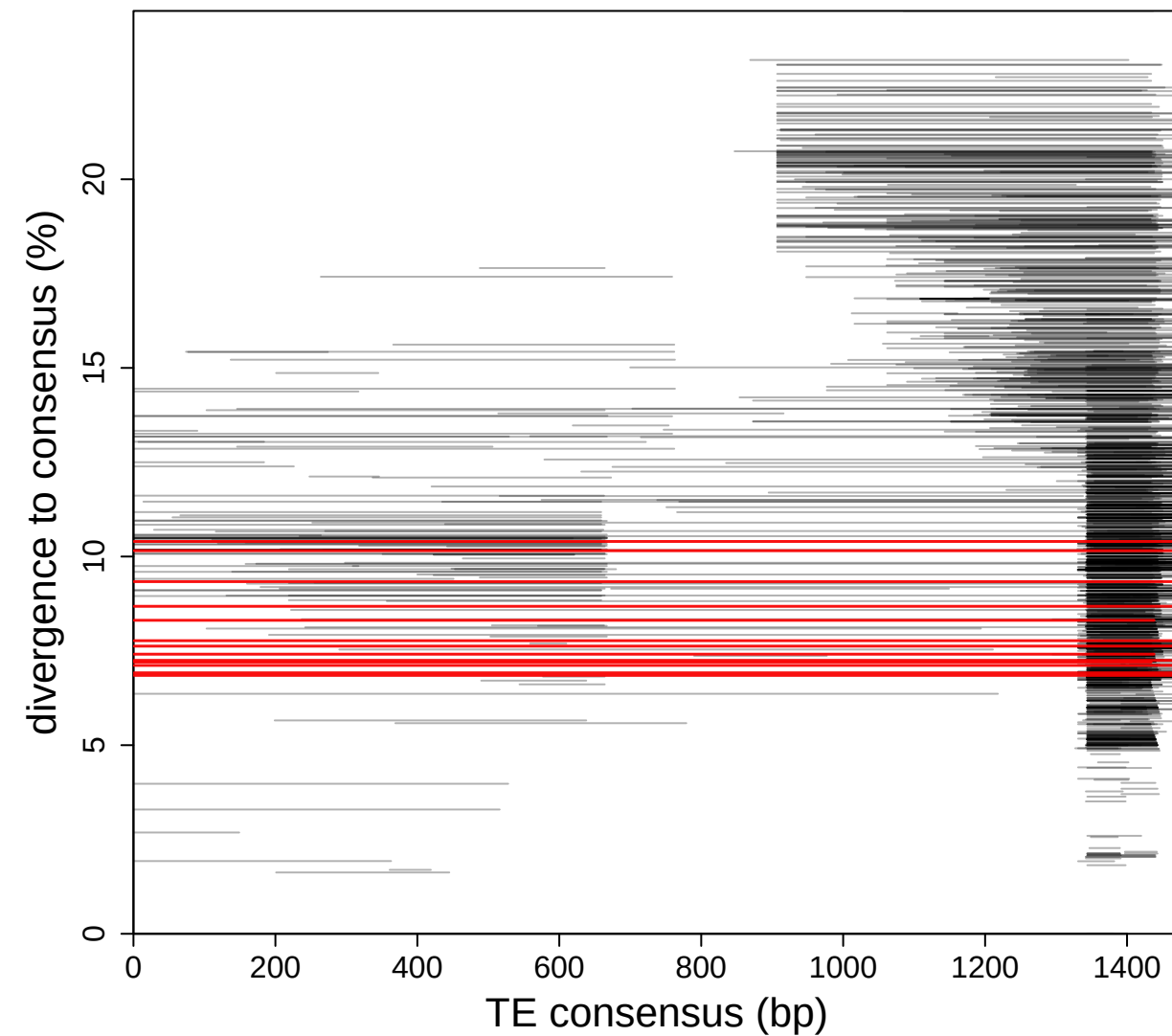
13.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.1
size: 3357bp; fragments: 2680; full length: 0 (>=3021.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

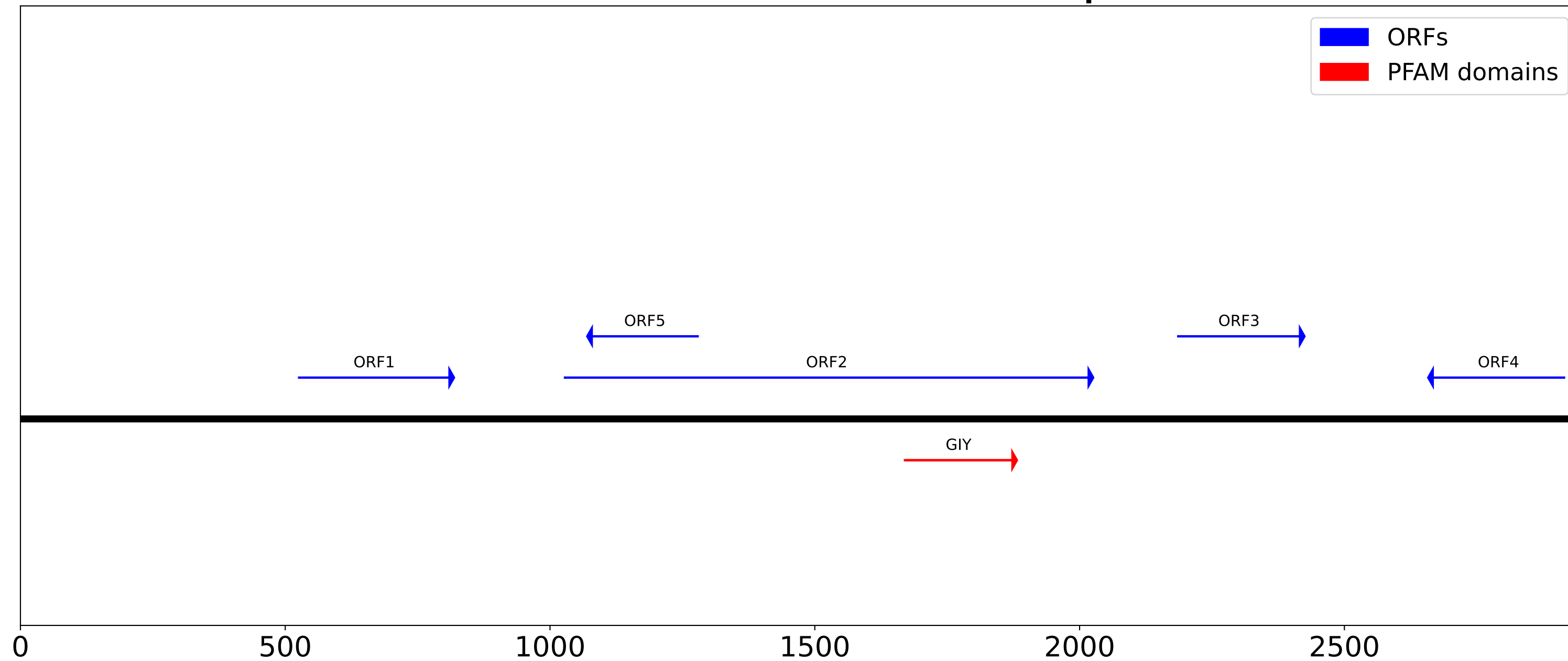


TE: rnd_5_family_10213
size: 1475bp; fragments: 1888; full length: 13 (≥ 1327.5 bp)

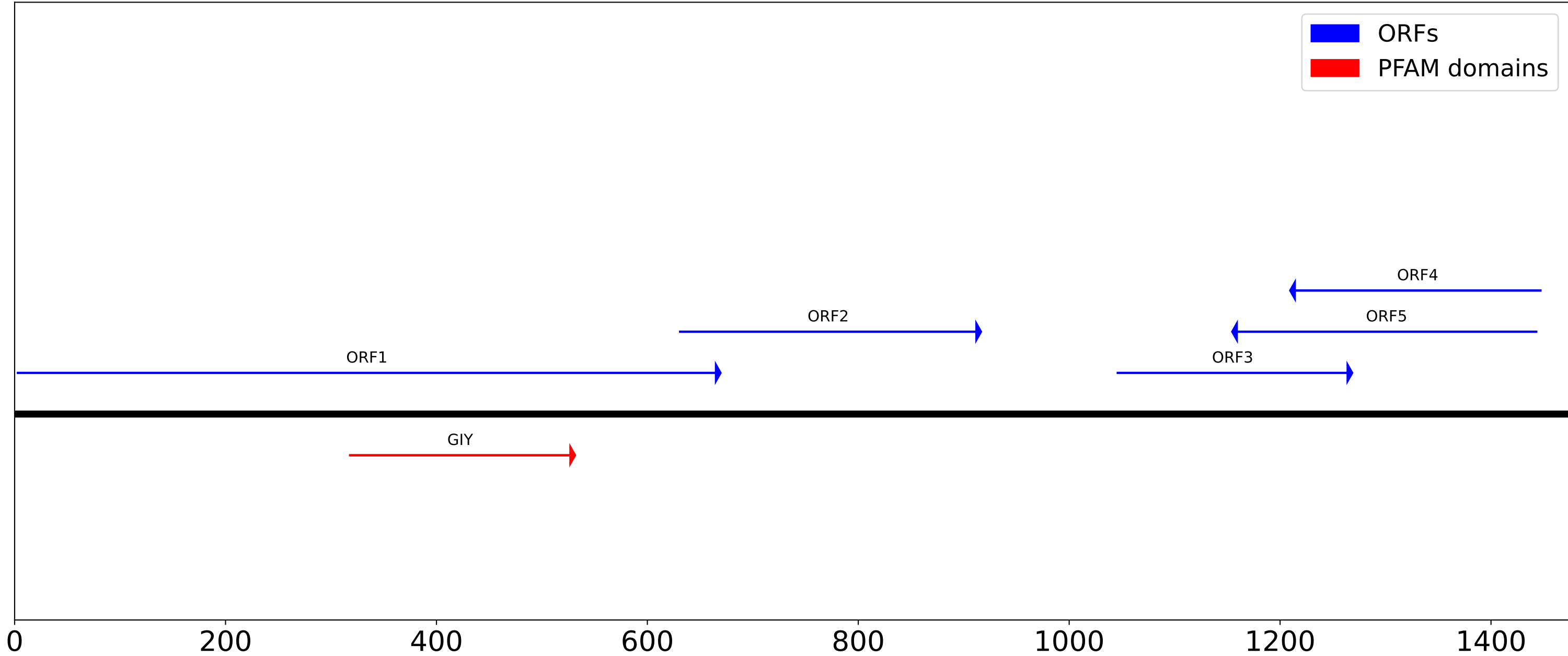
Before TEtrimmer 1475 bp



After TEtrimmer ORF and PFAM domain plot



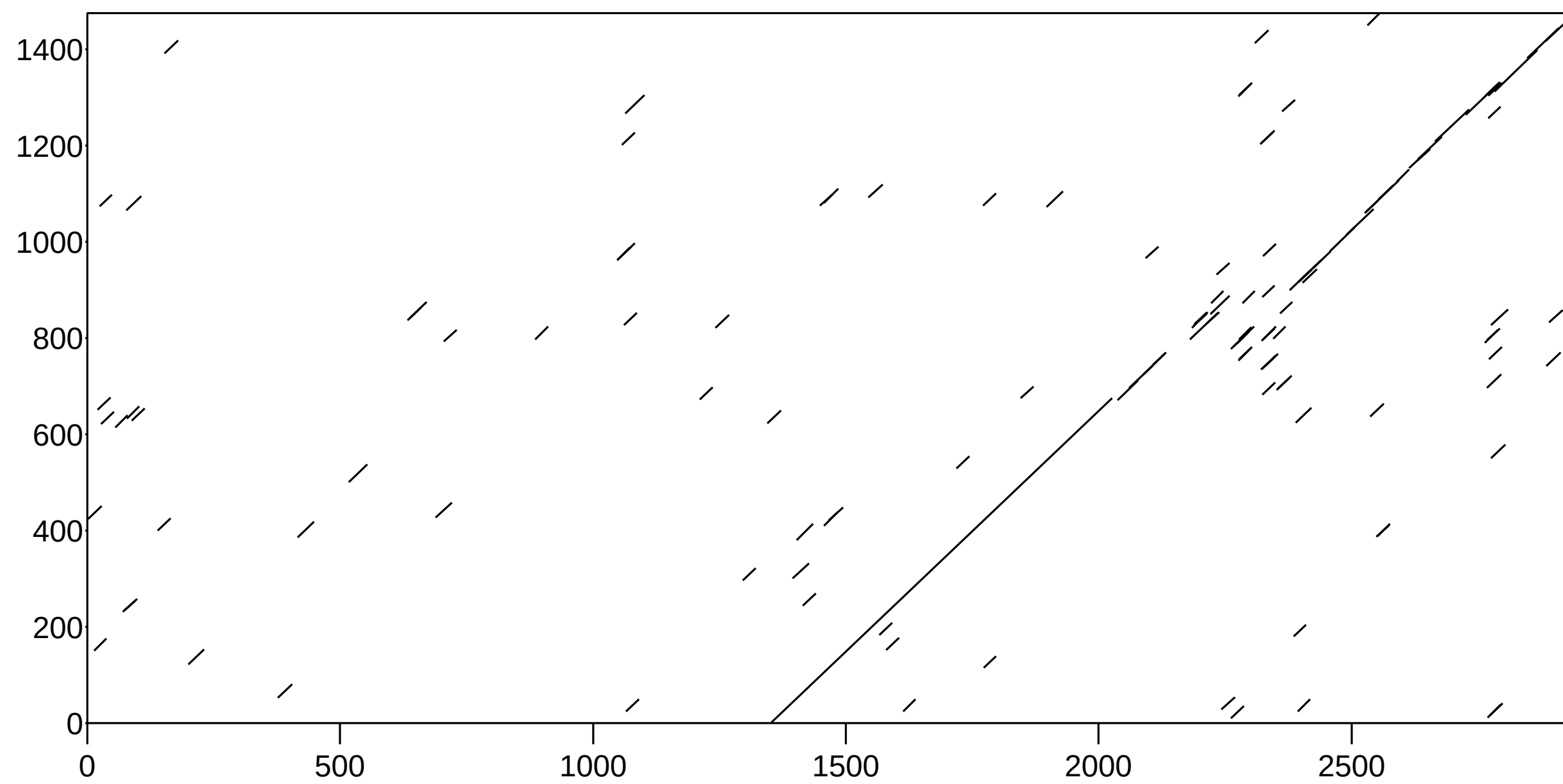
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)