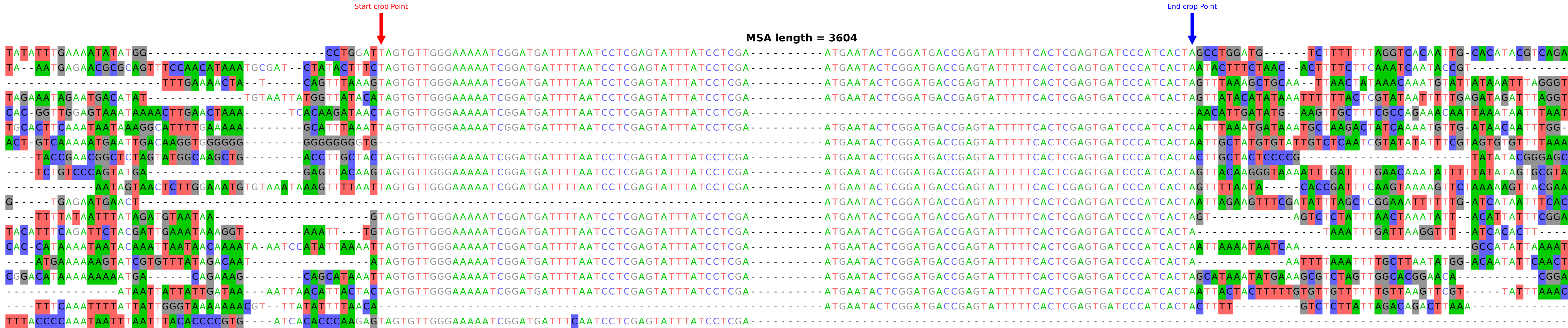
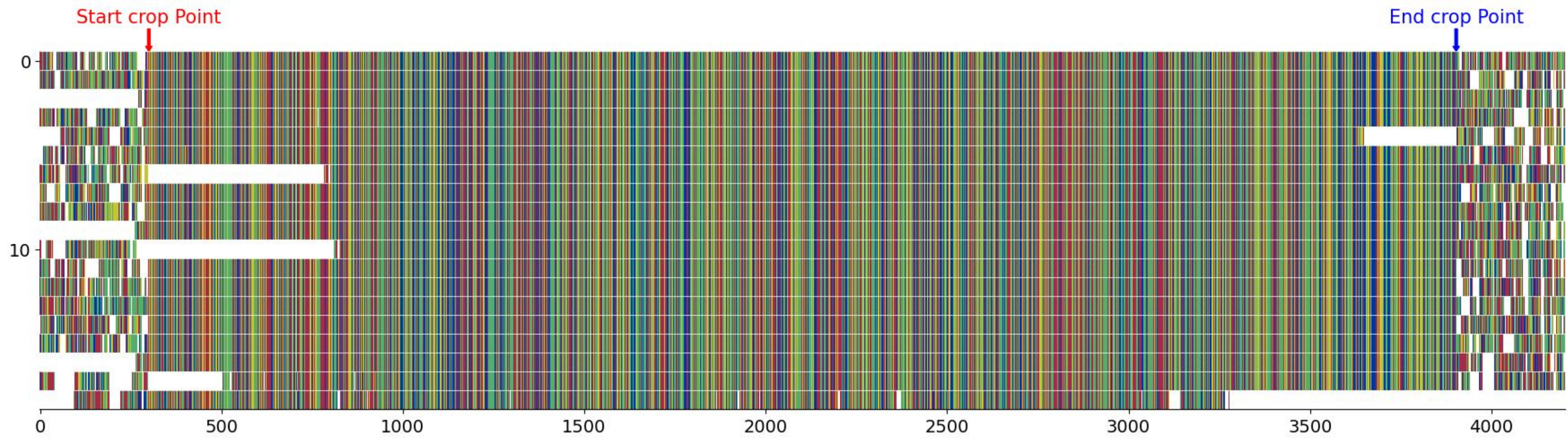




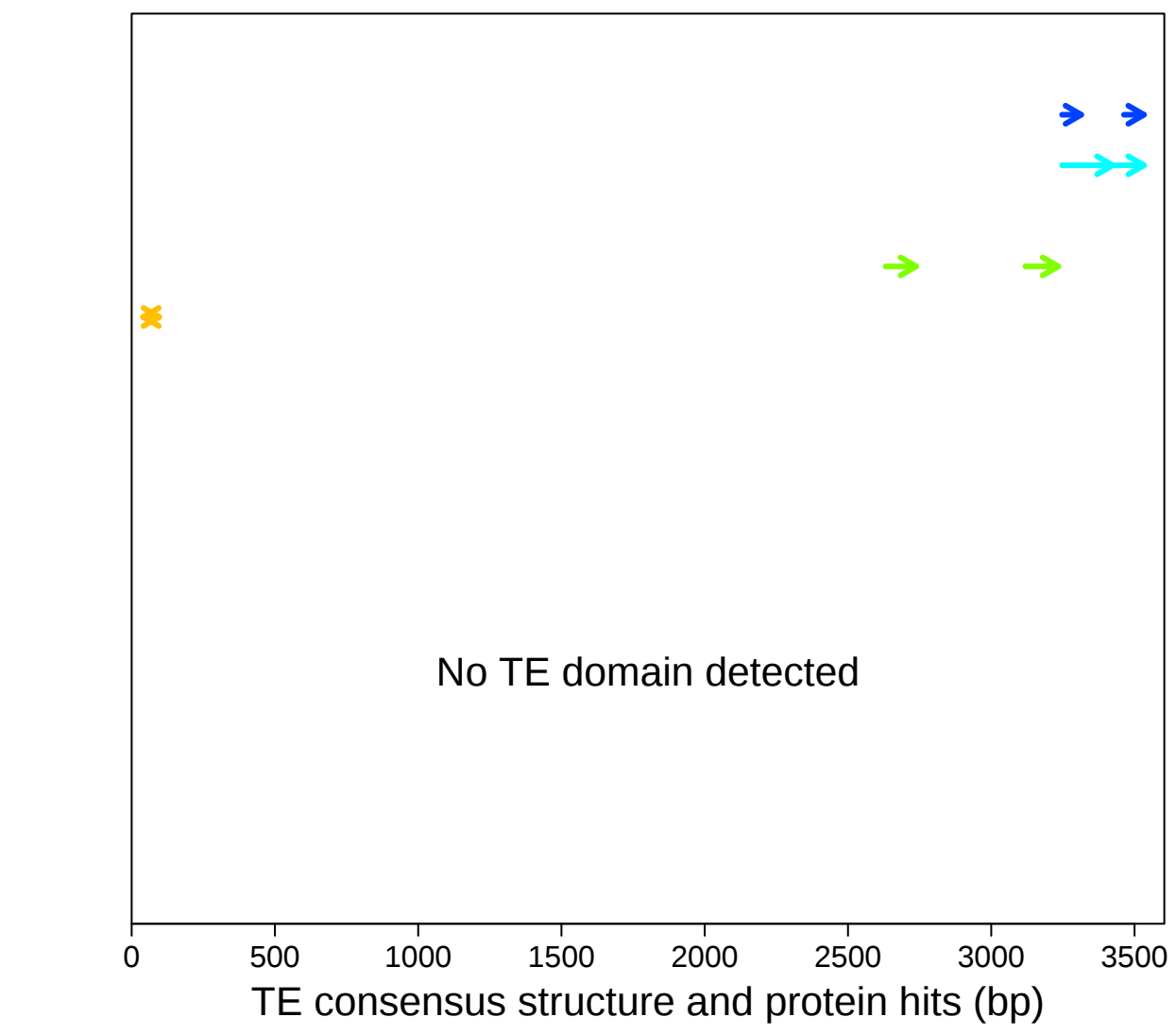
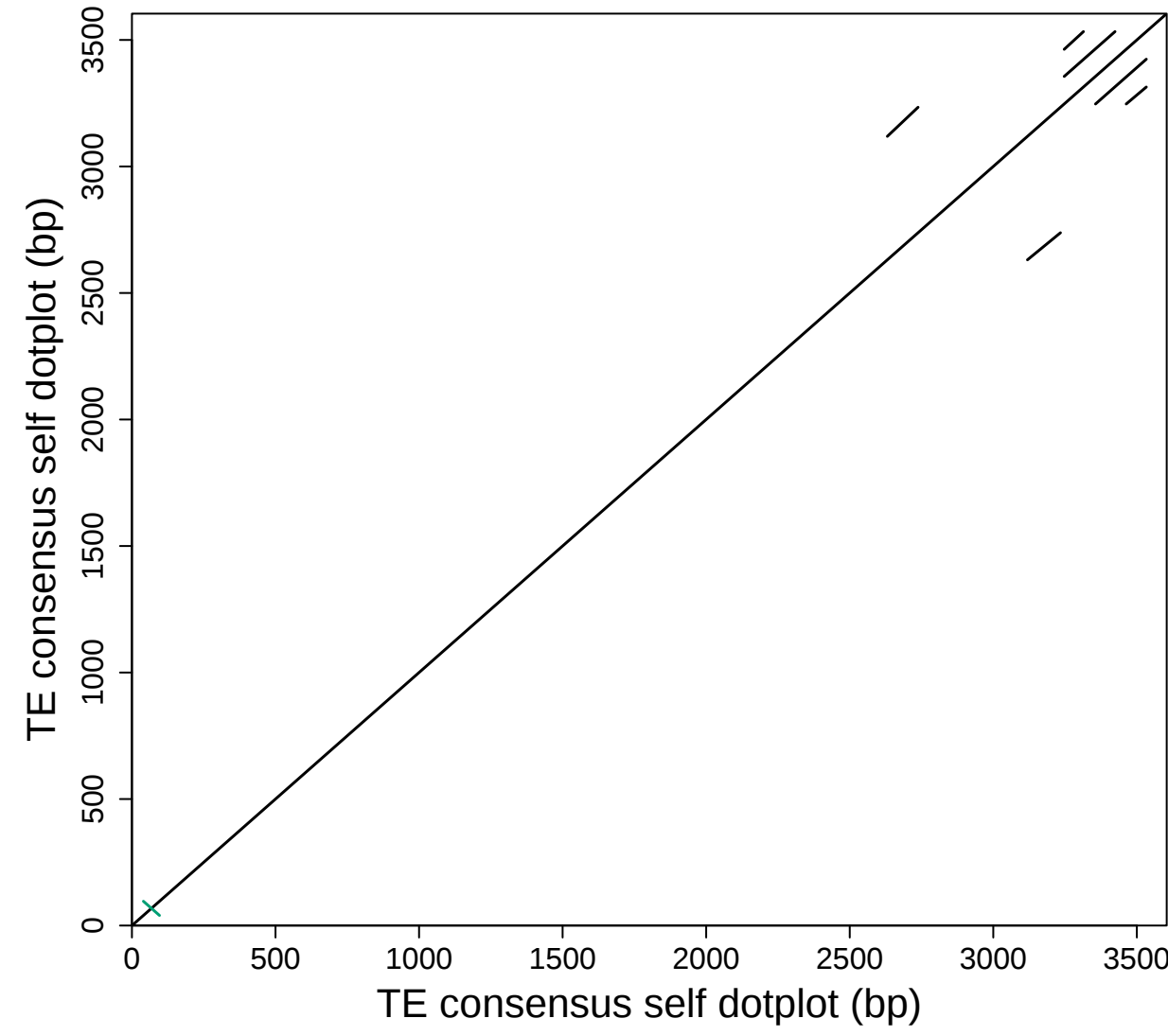
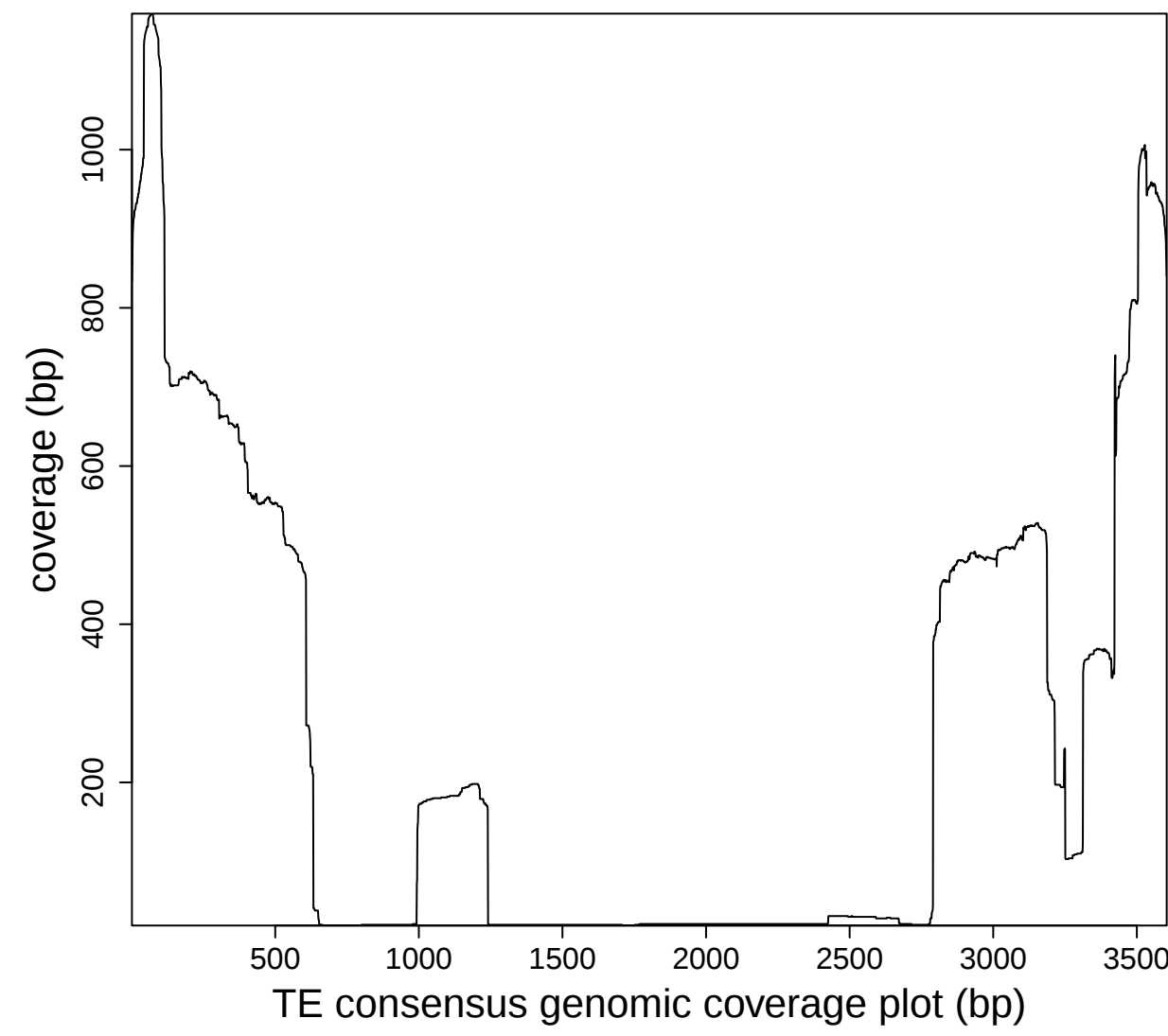
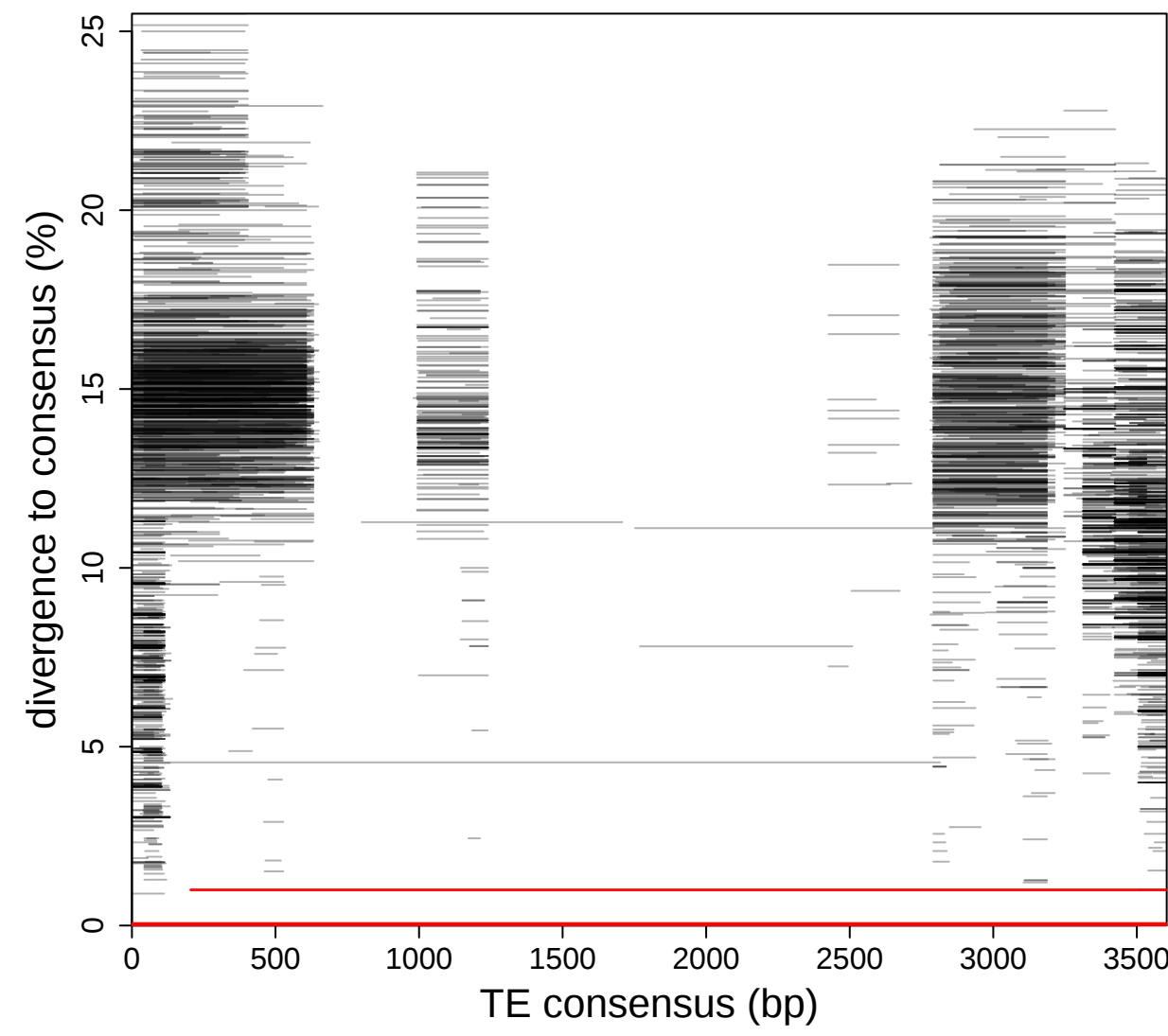
Start crop Point

End crop Point



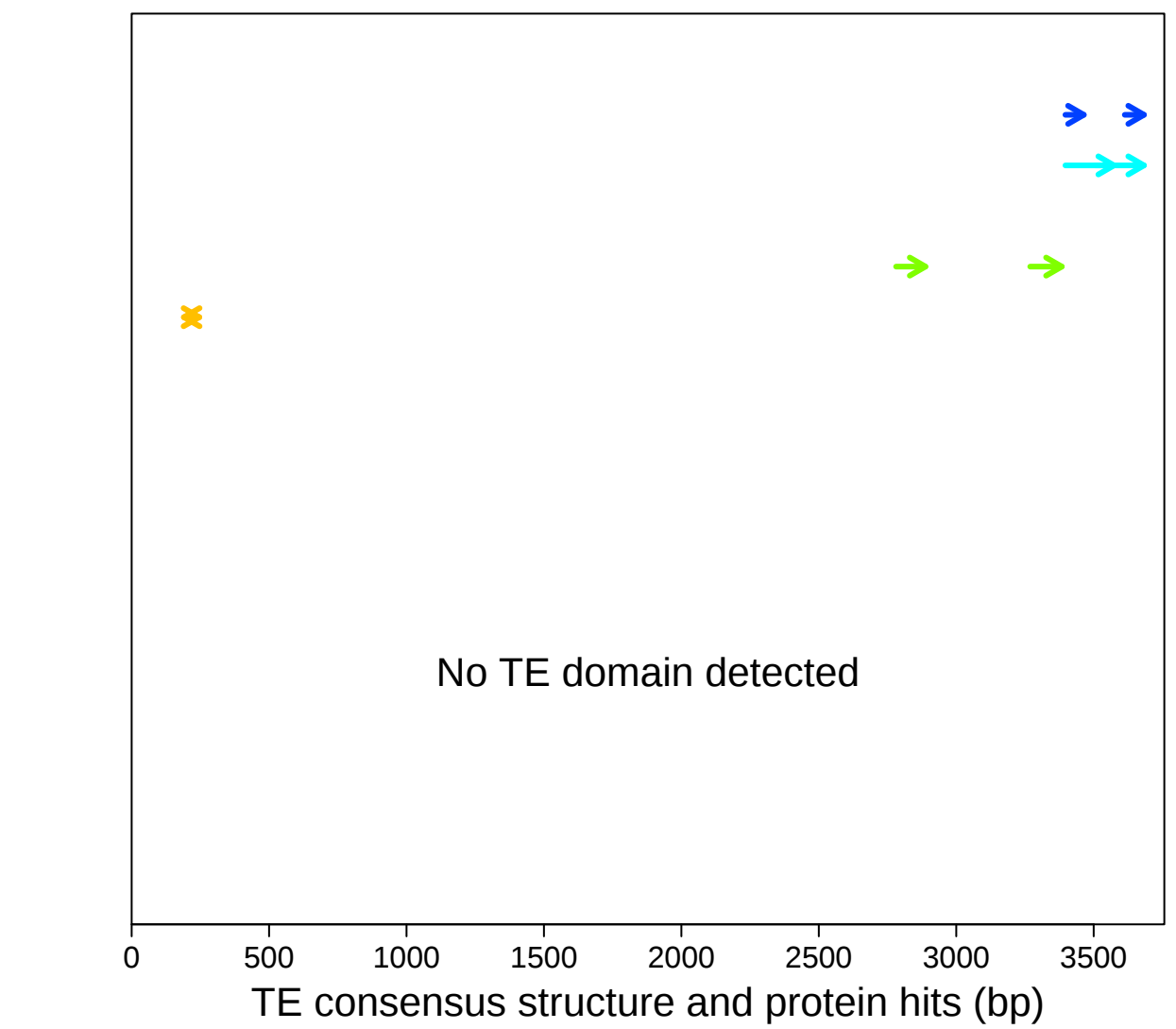
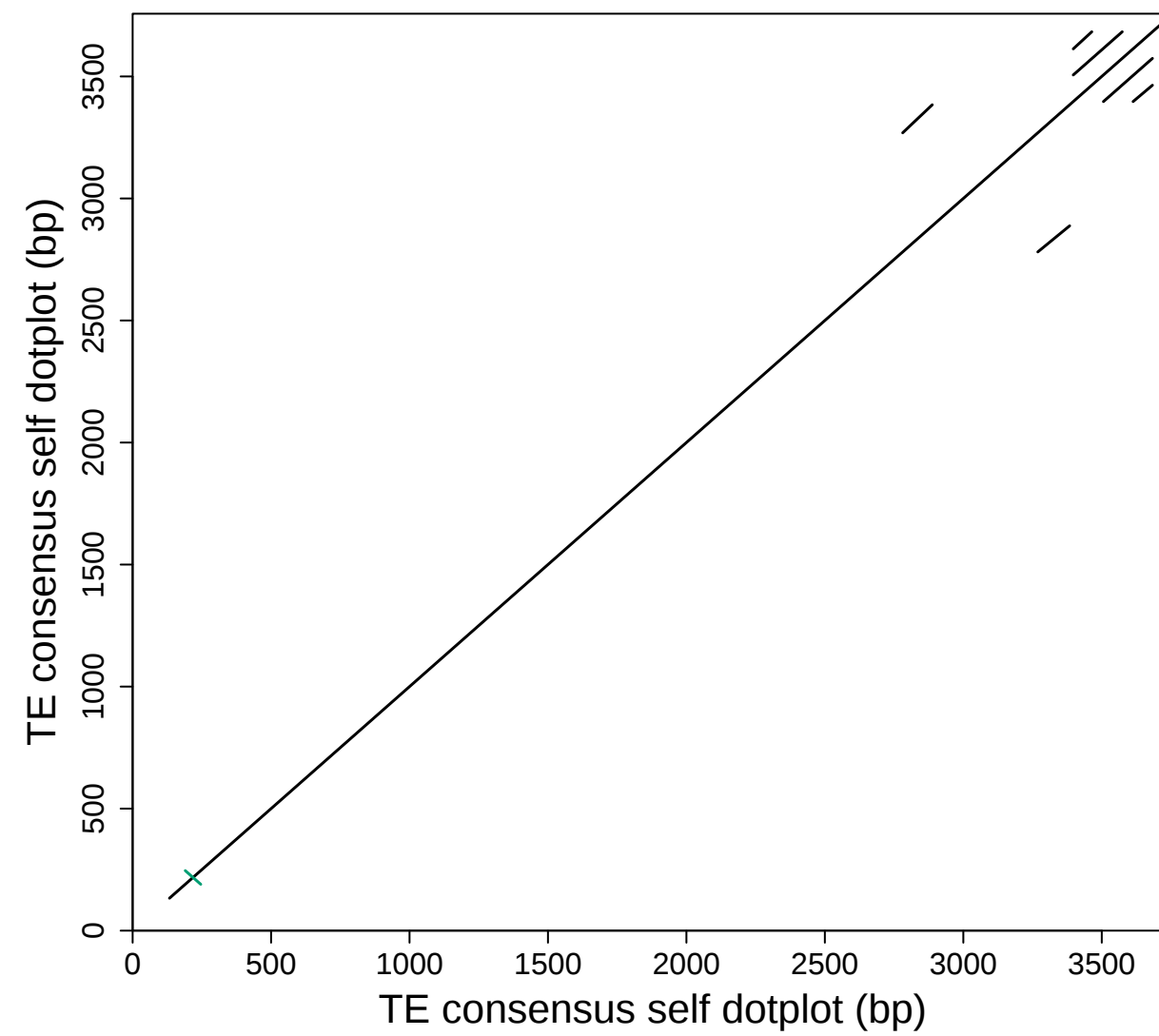
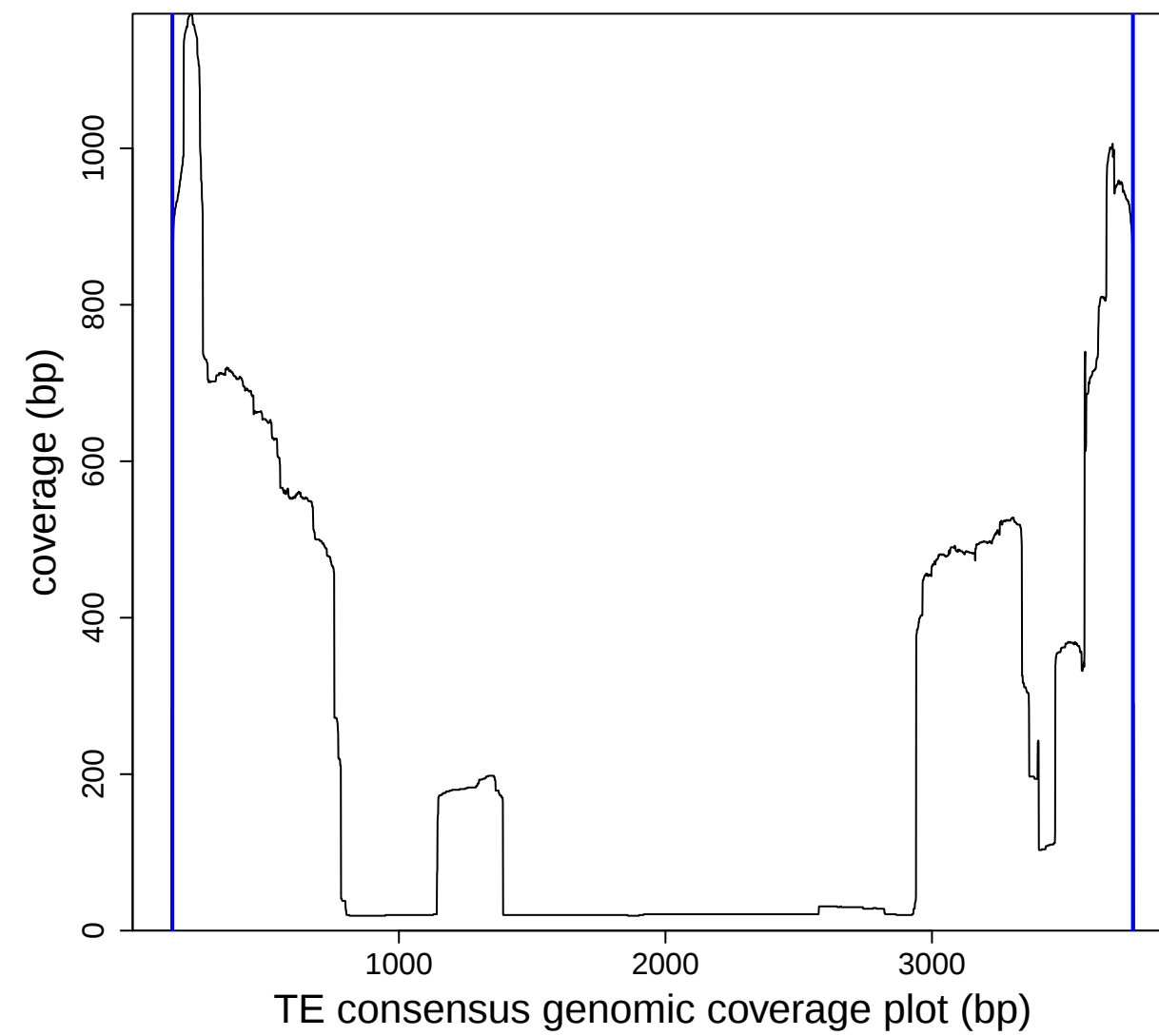
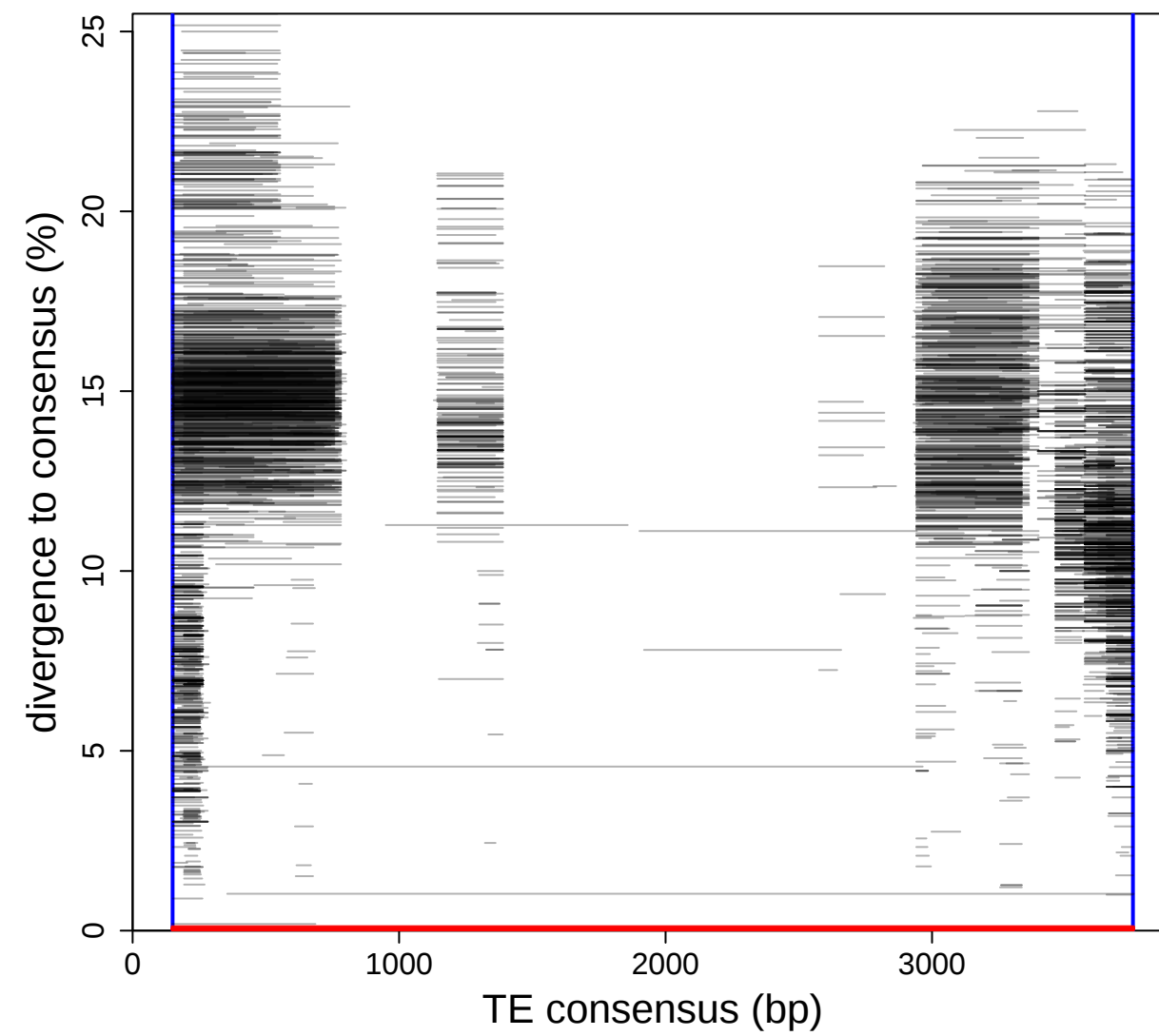
g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 3604bp; fragments: 3265; full length: 16 (≥ 3243.6 bp)

After TETrimmer 3604 bp



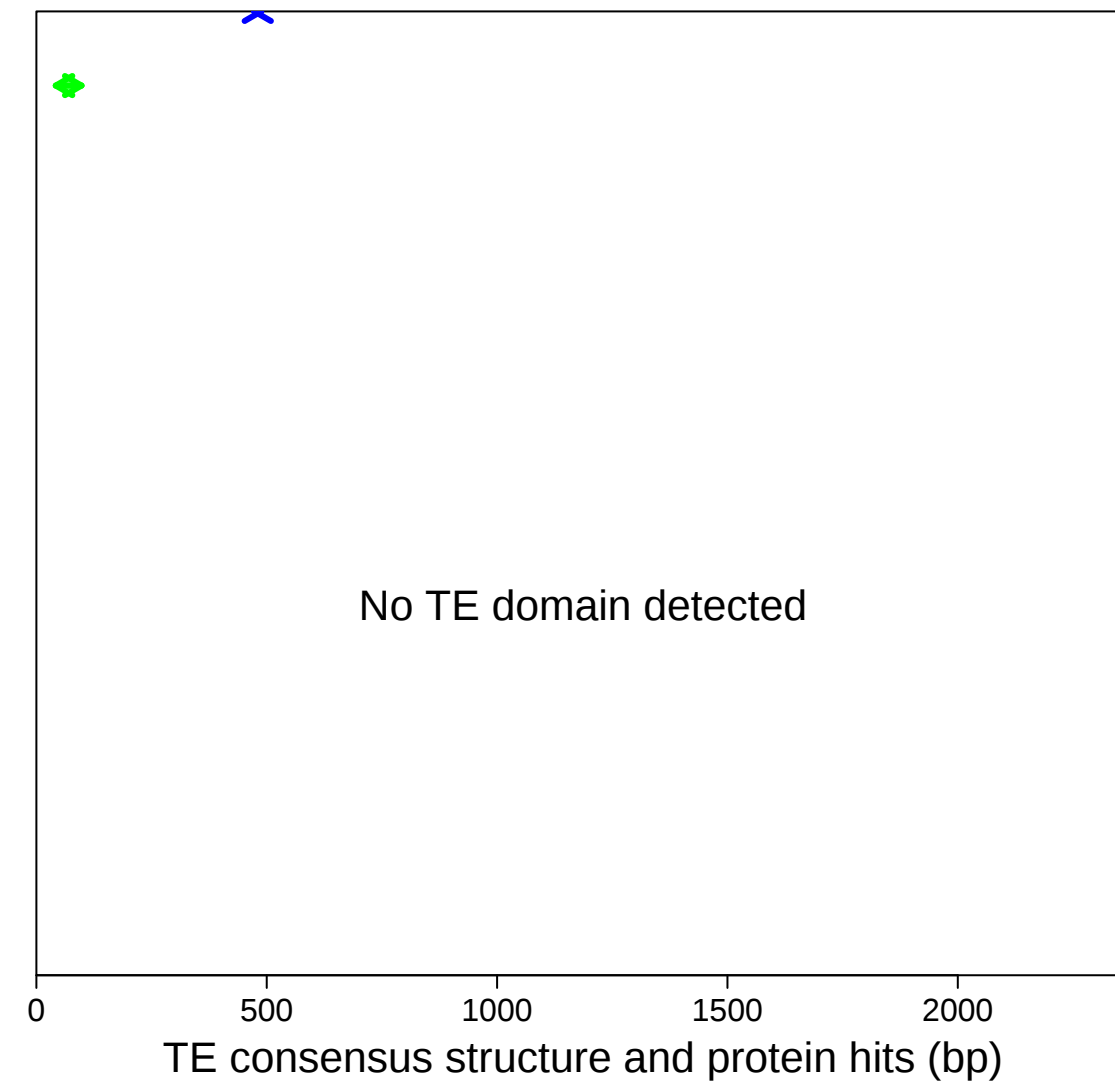
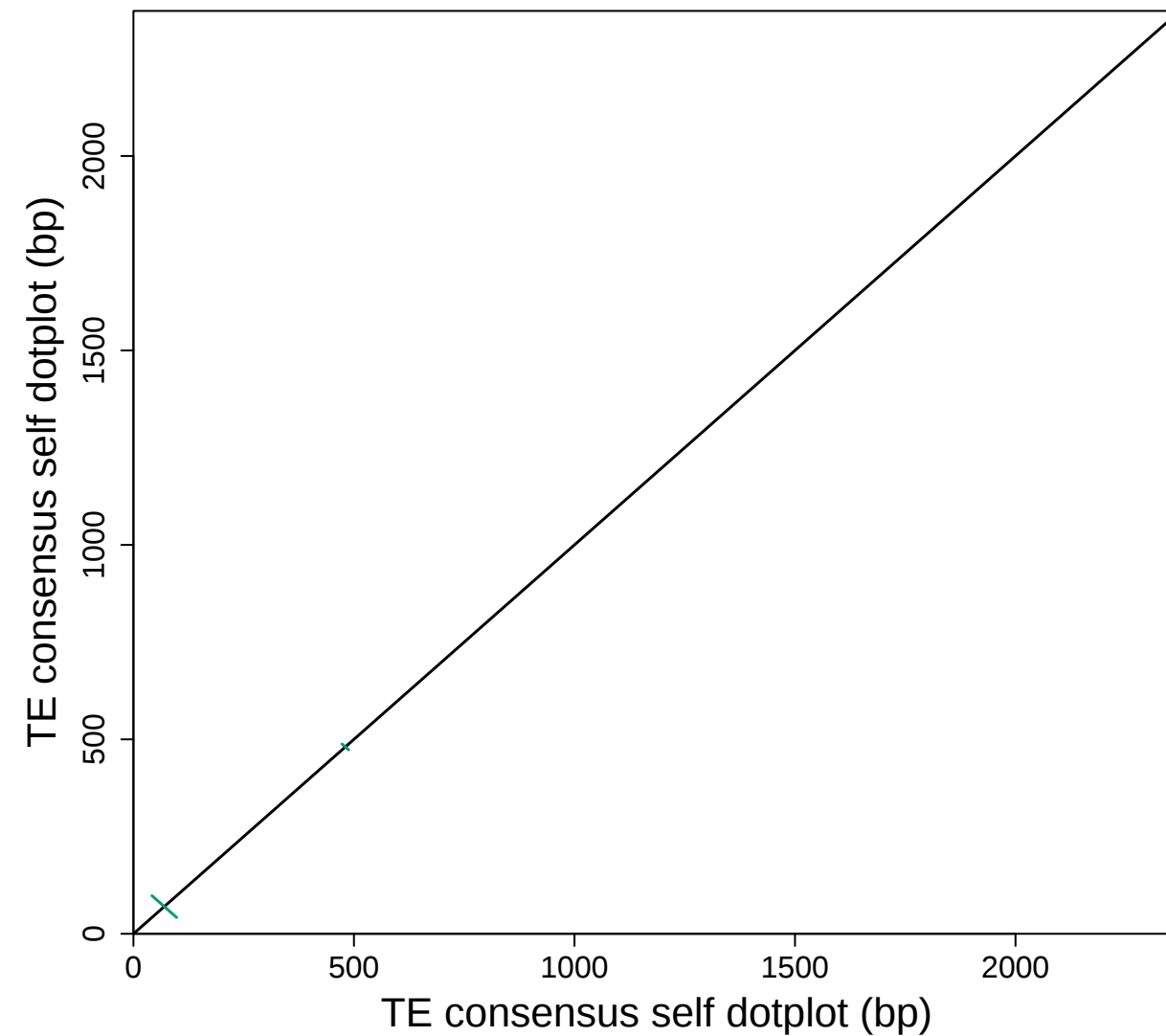
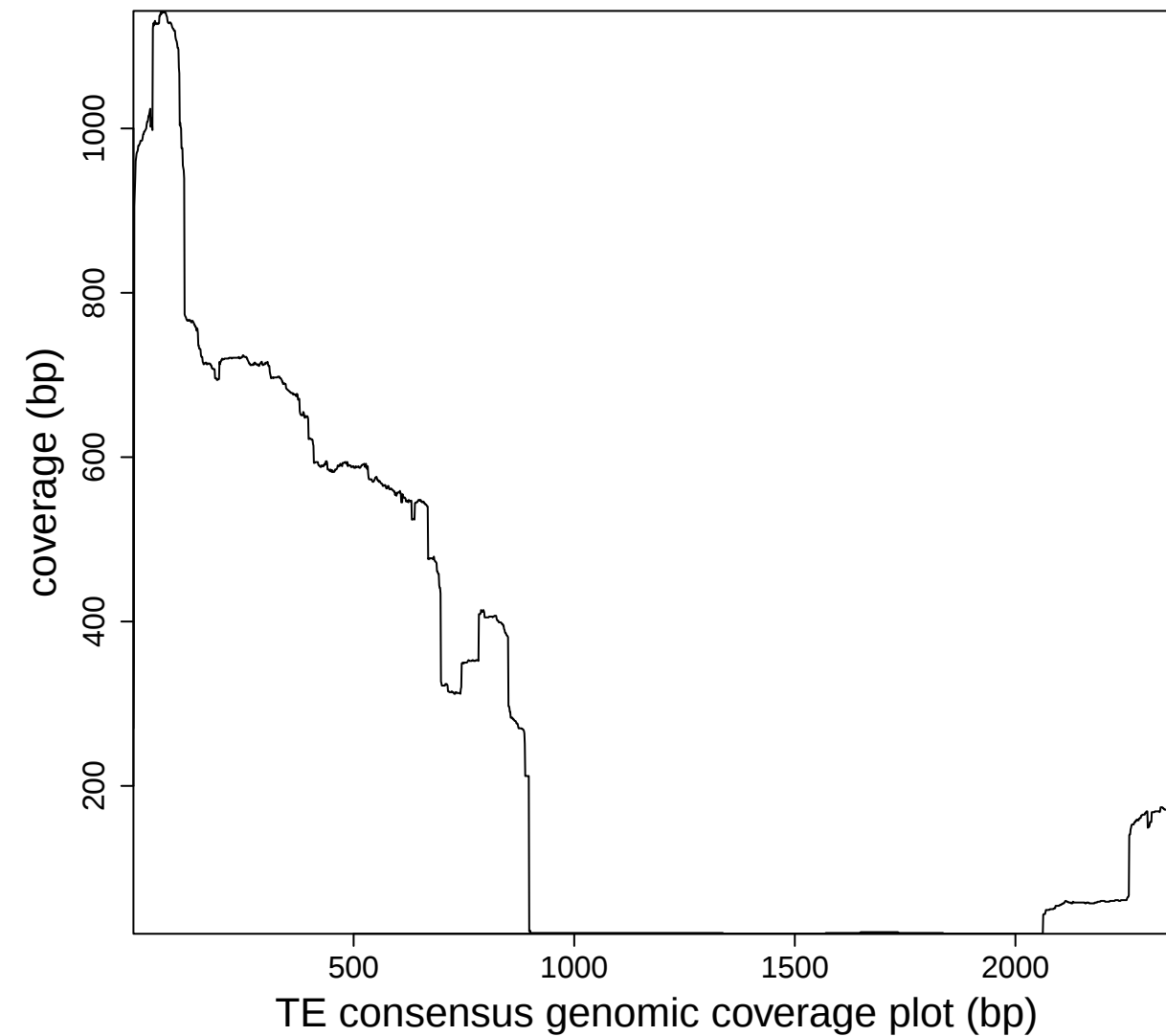
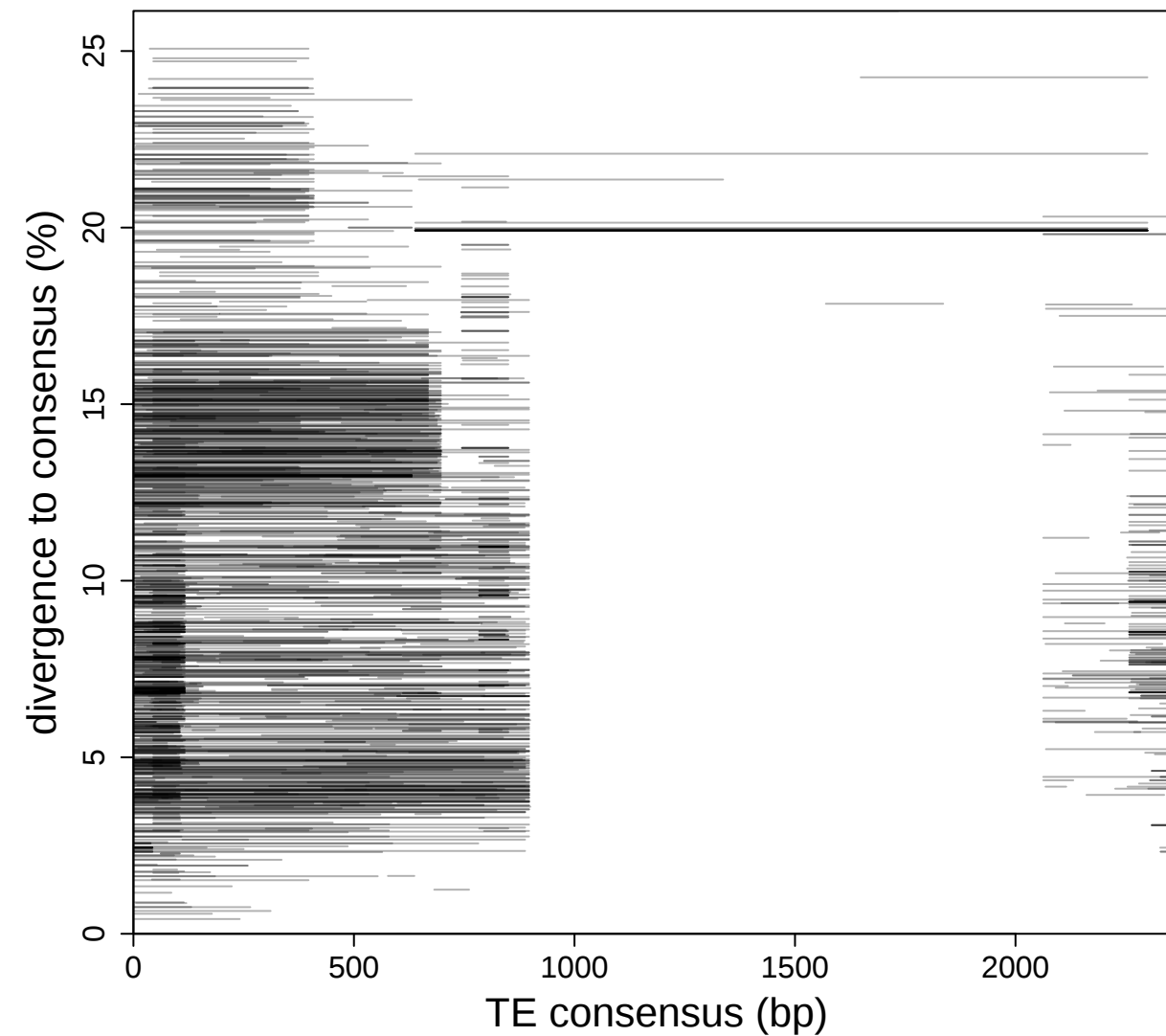
f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 3904bp; fragments: 3265; full length: 14 (>=3513.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

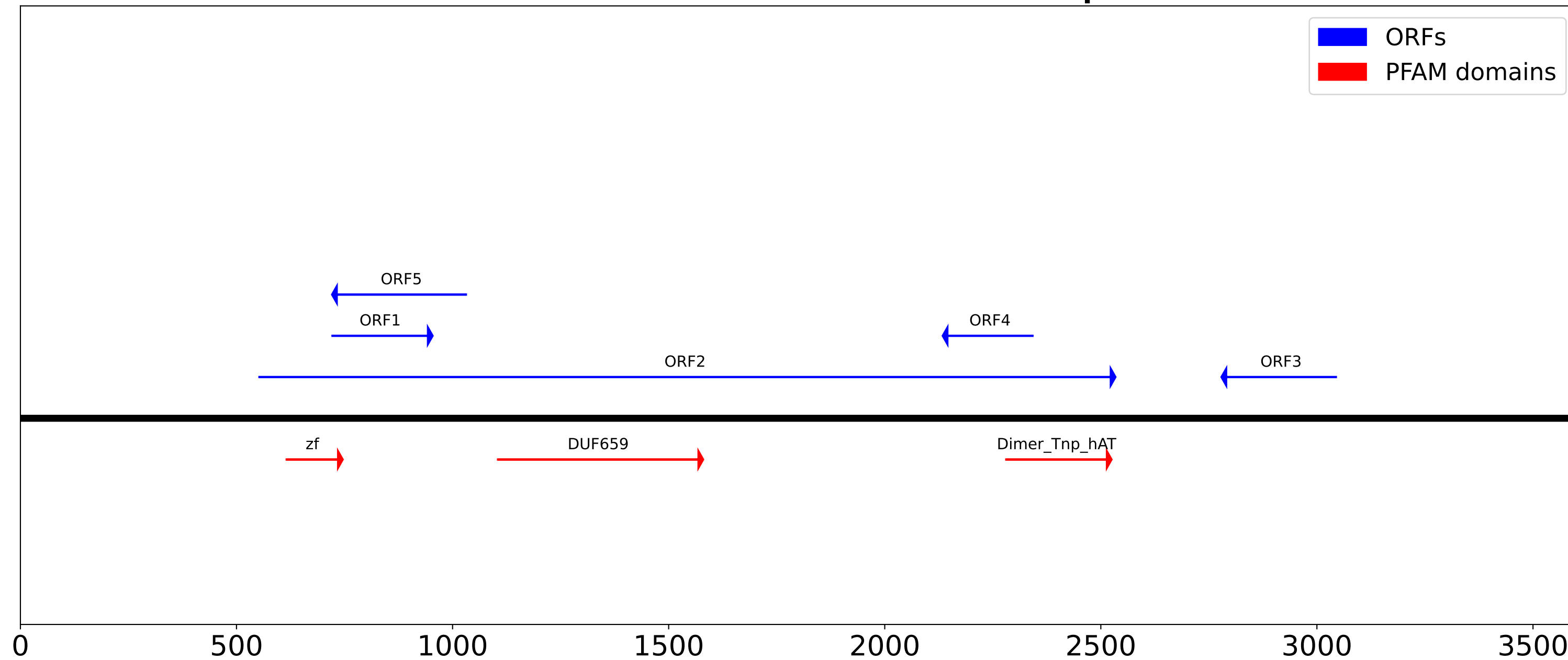


TE: rnd_5_family_482
size: 2373bp; fragments: 1771; full length: 0 (>=2135.7bp)

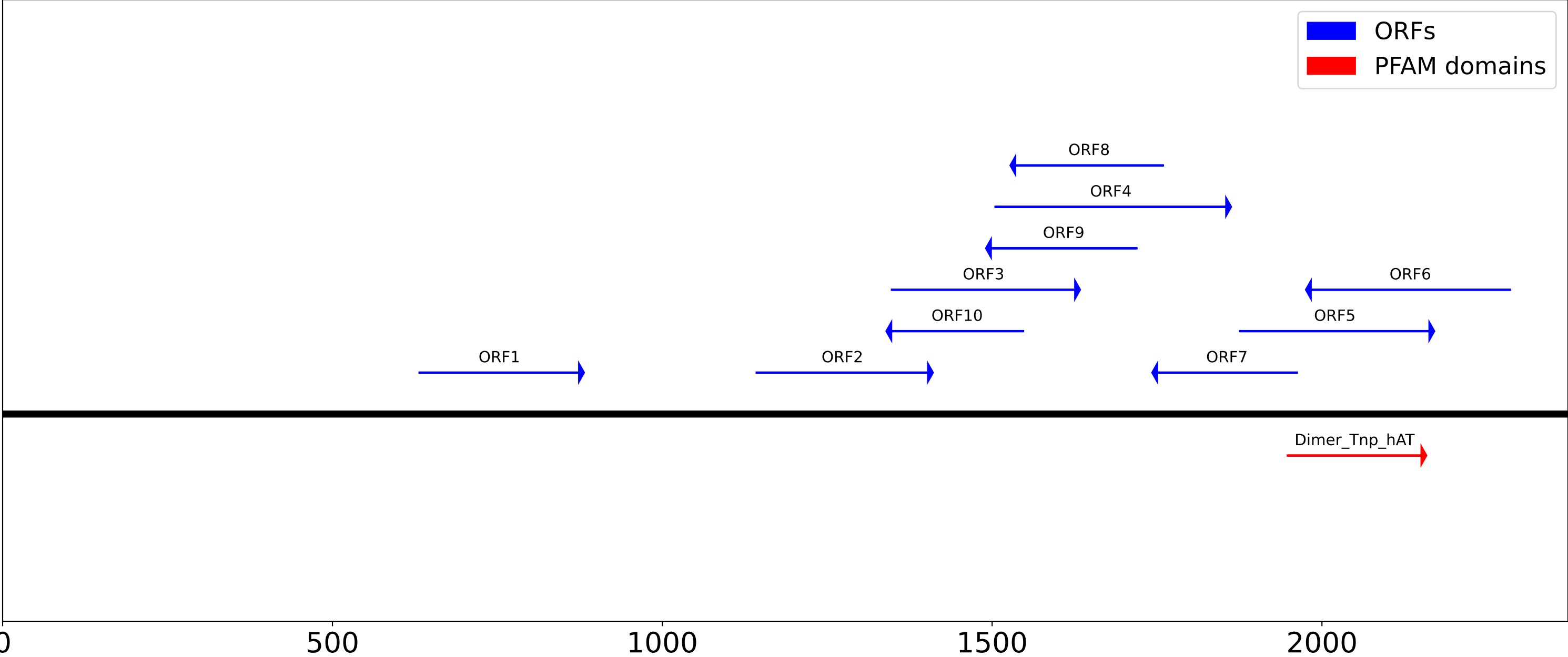
Before TEtrimmer 2373 bp



After TEtrimmer ORF and PFAM domain plot



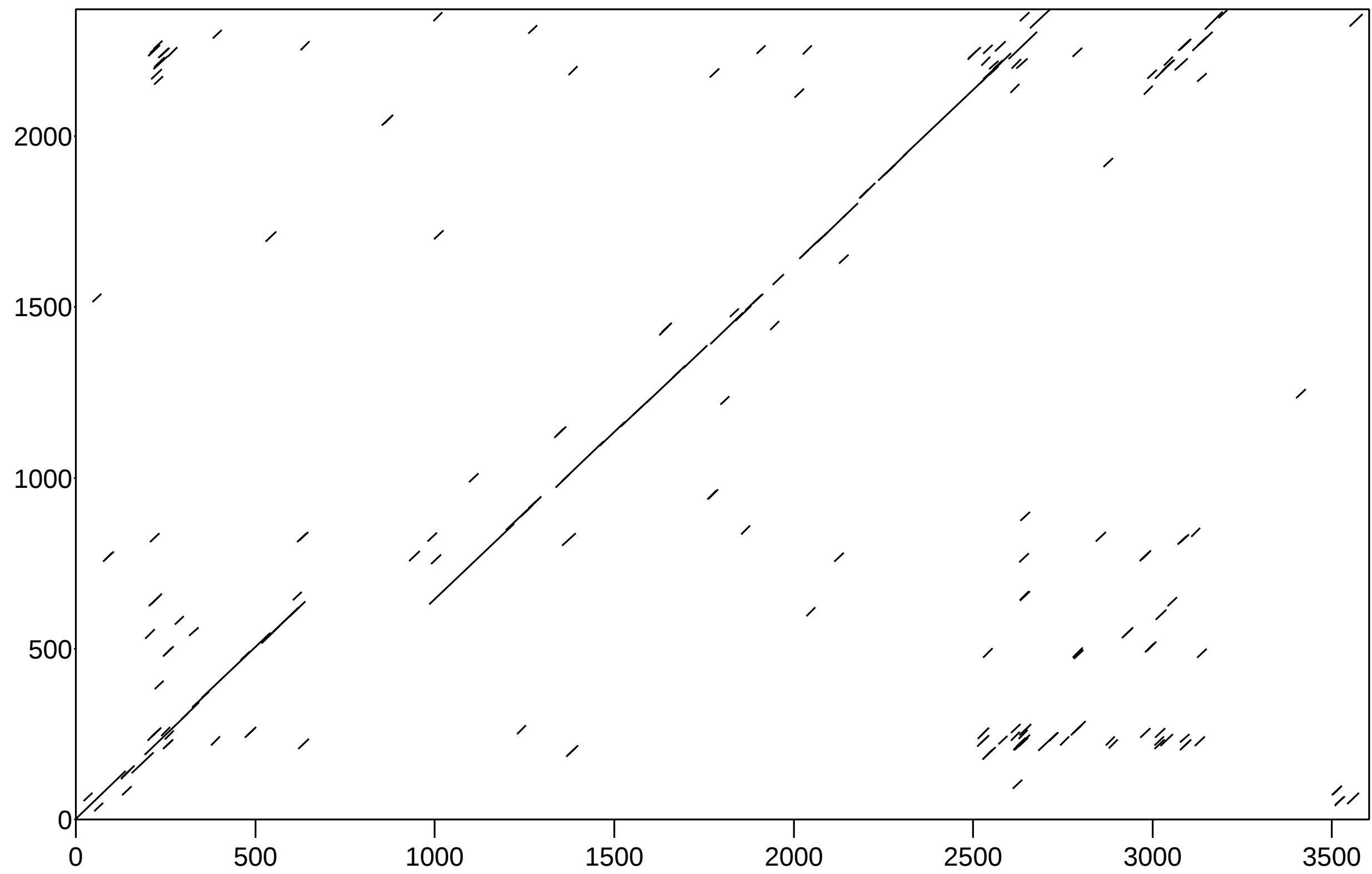
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)