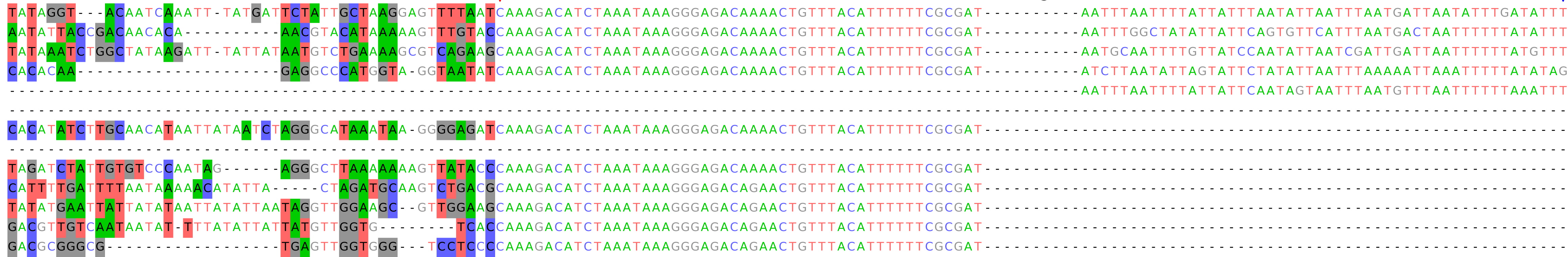


Start crop Point

End crop Point

MSA length = 7126



Start crop Point

End crop Point

0

10

0

1000

2000

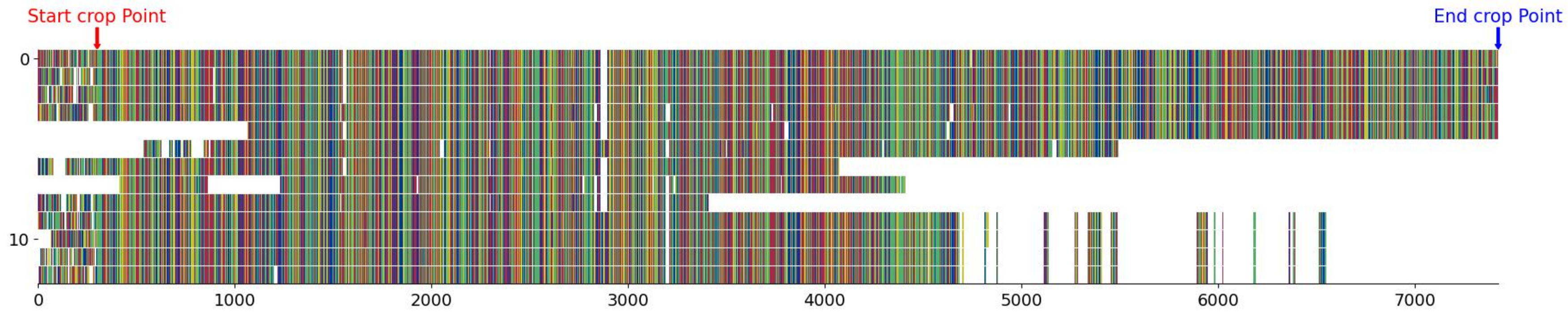
3000

4000

5000

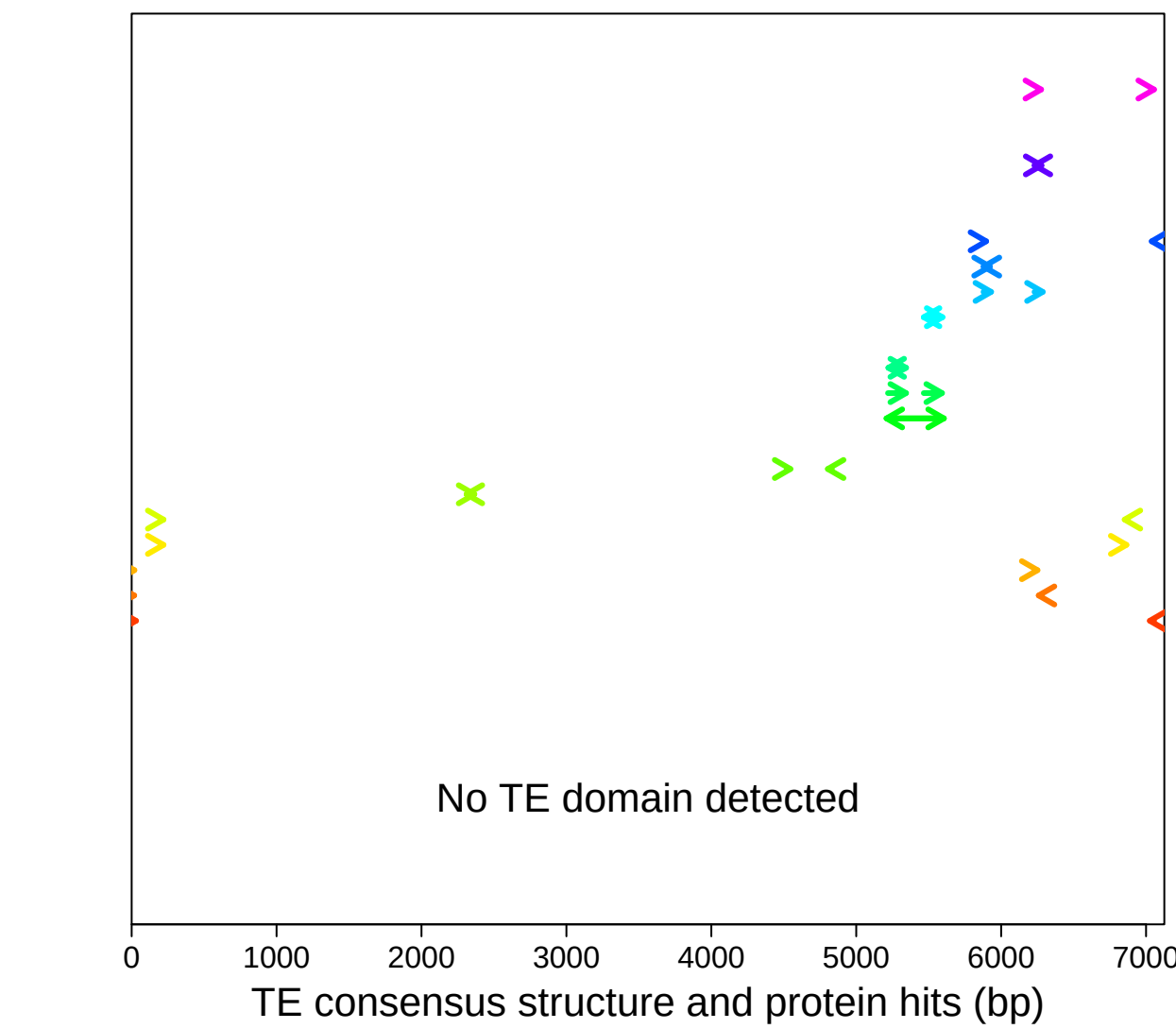
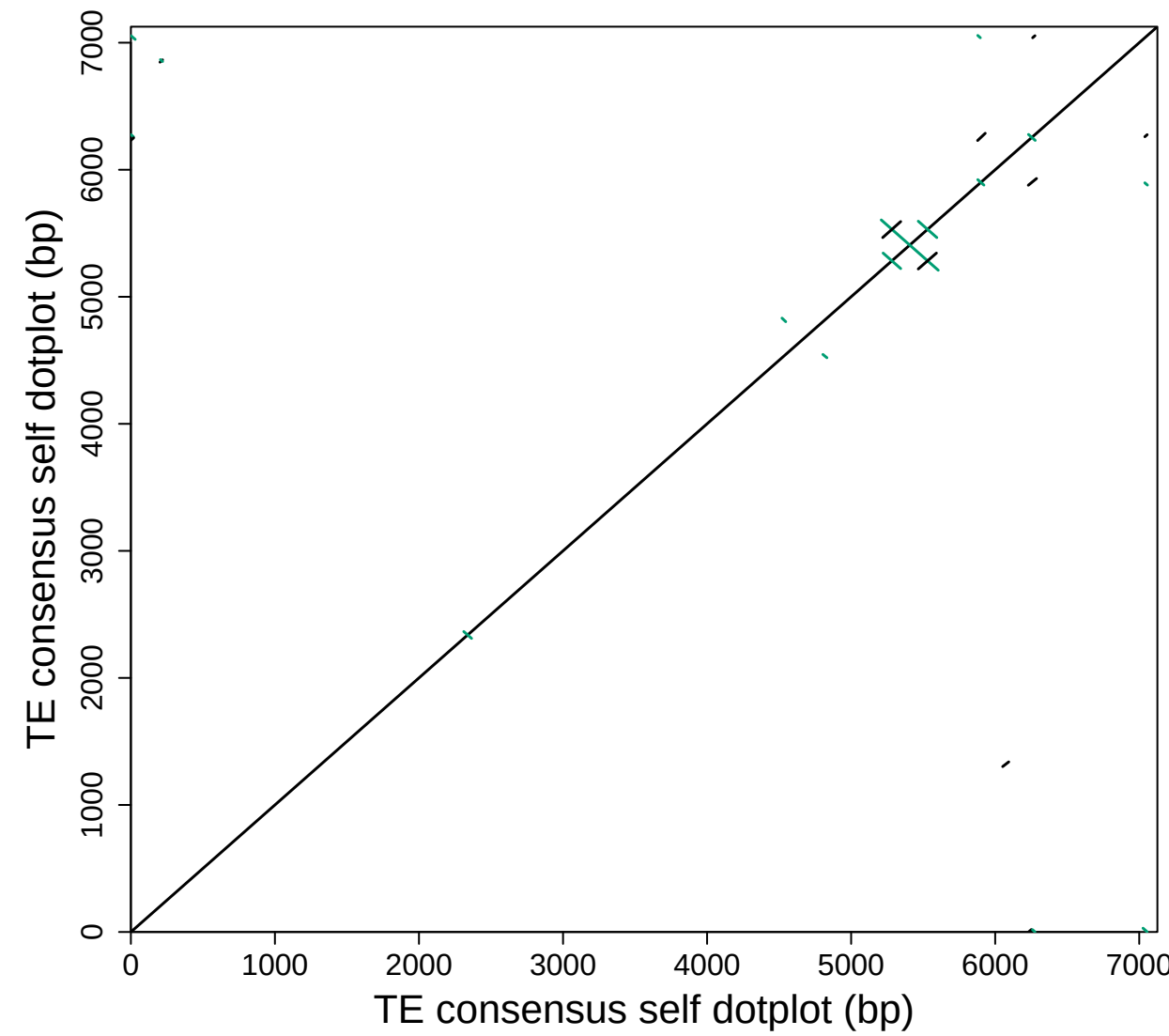
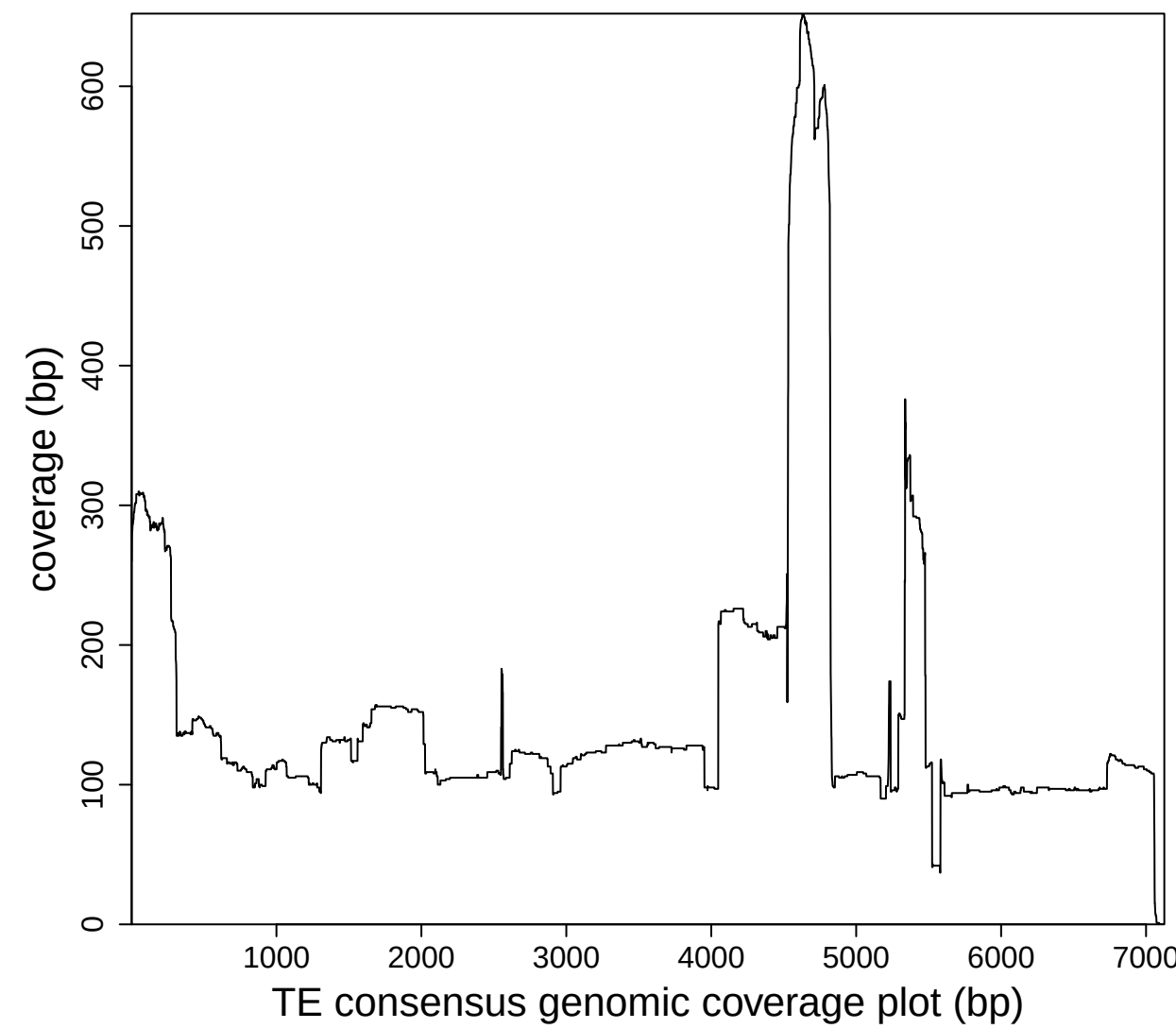
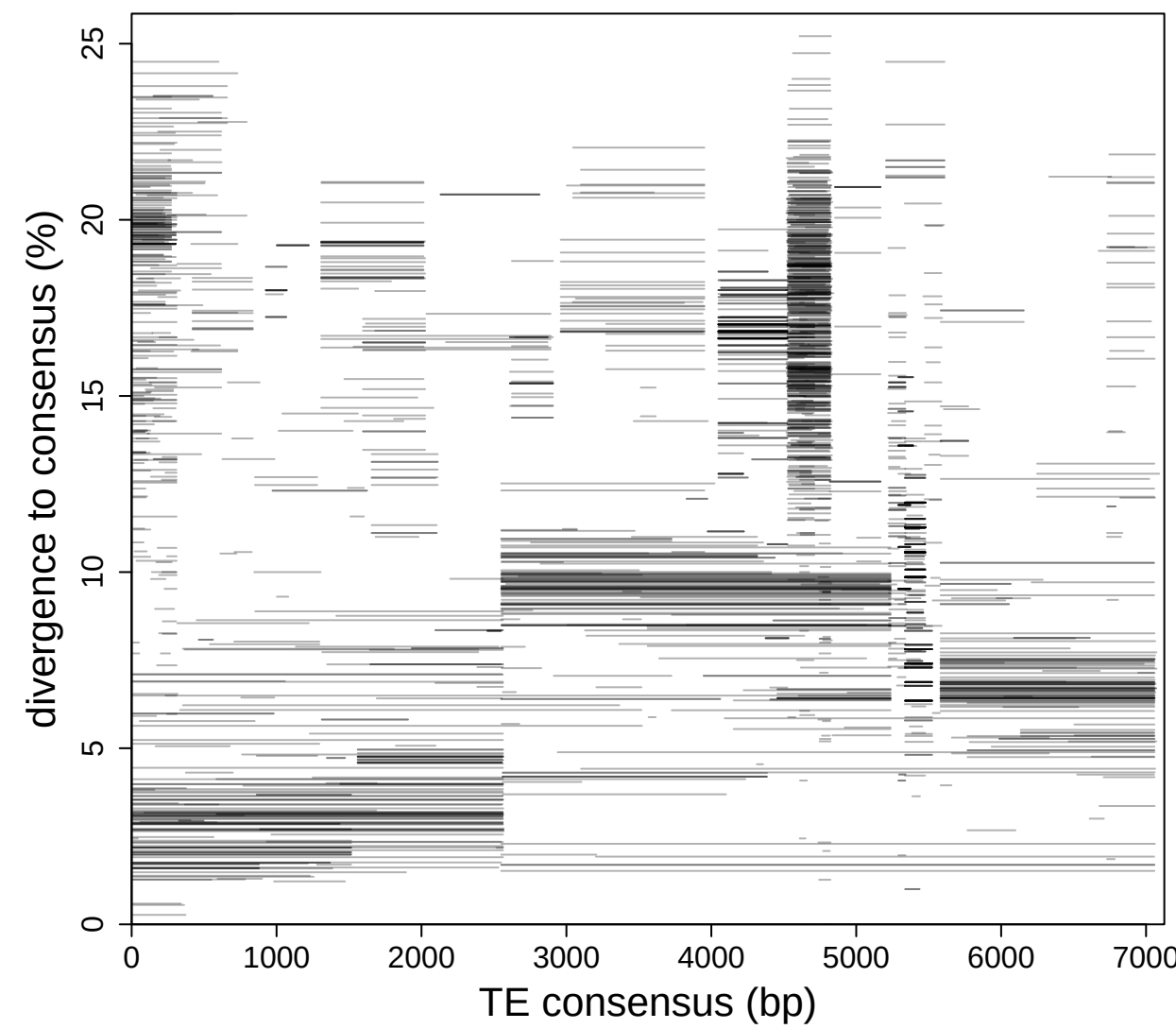
6000

7000



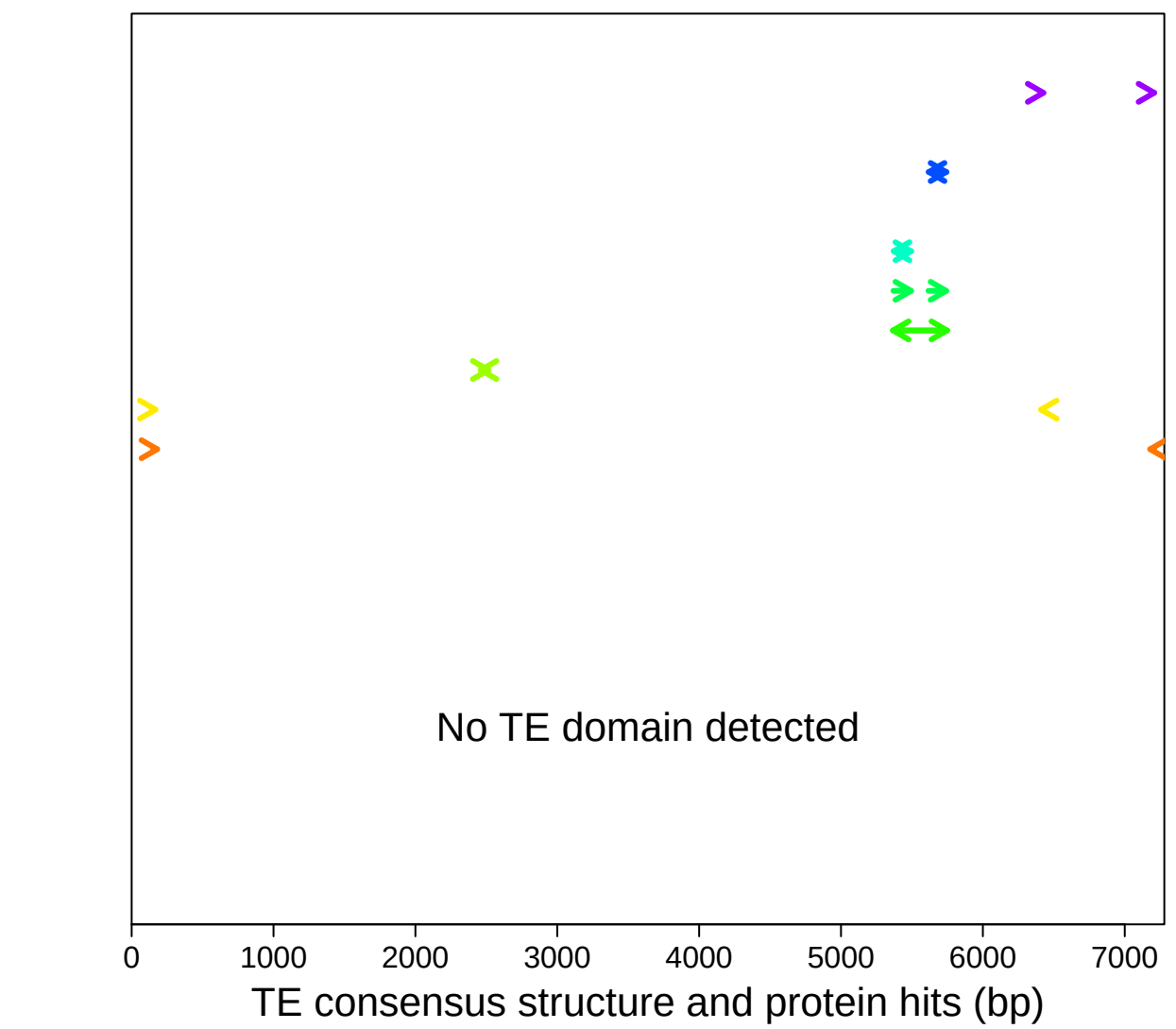
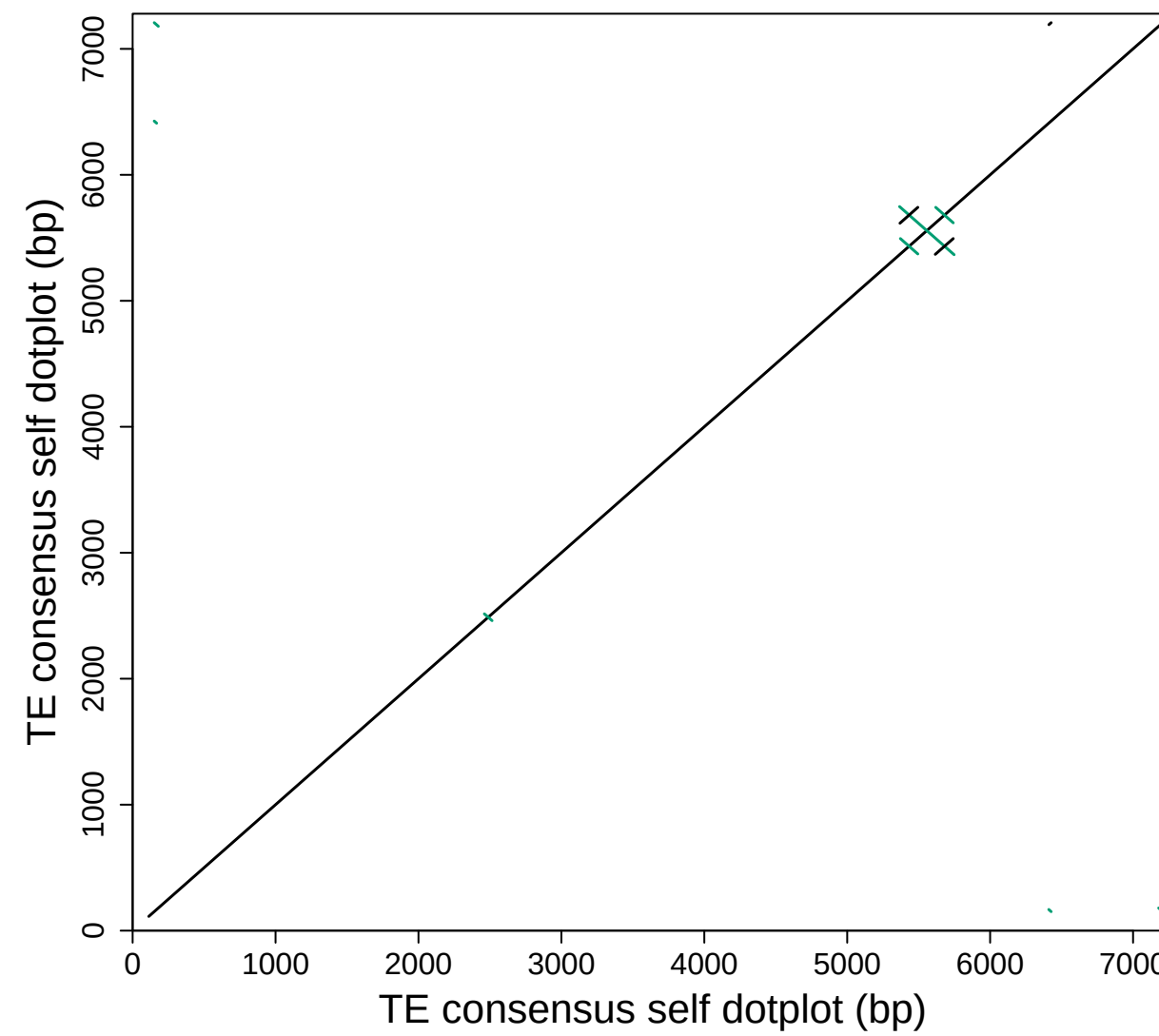
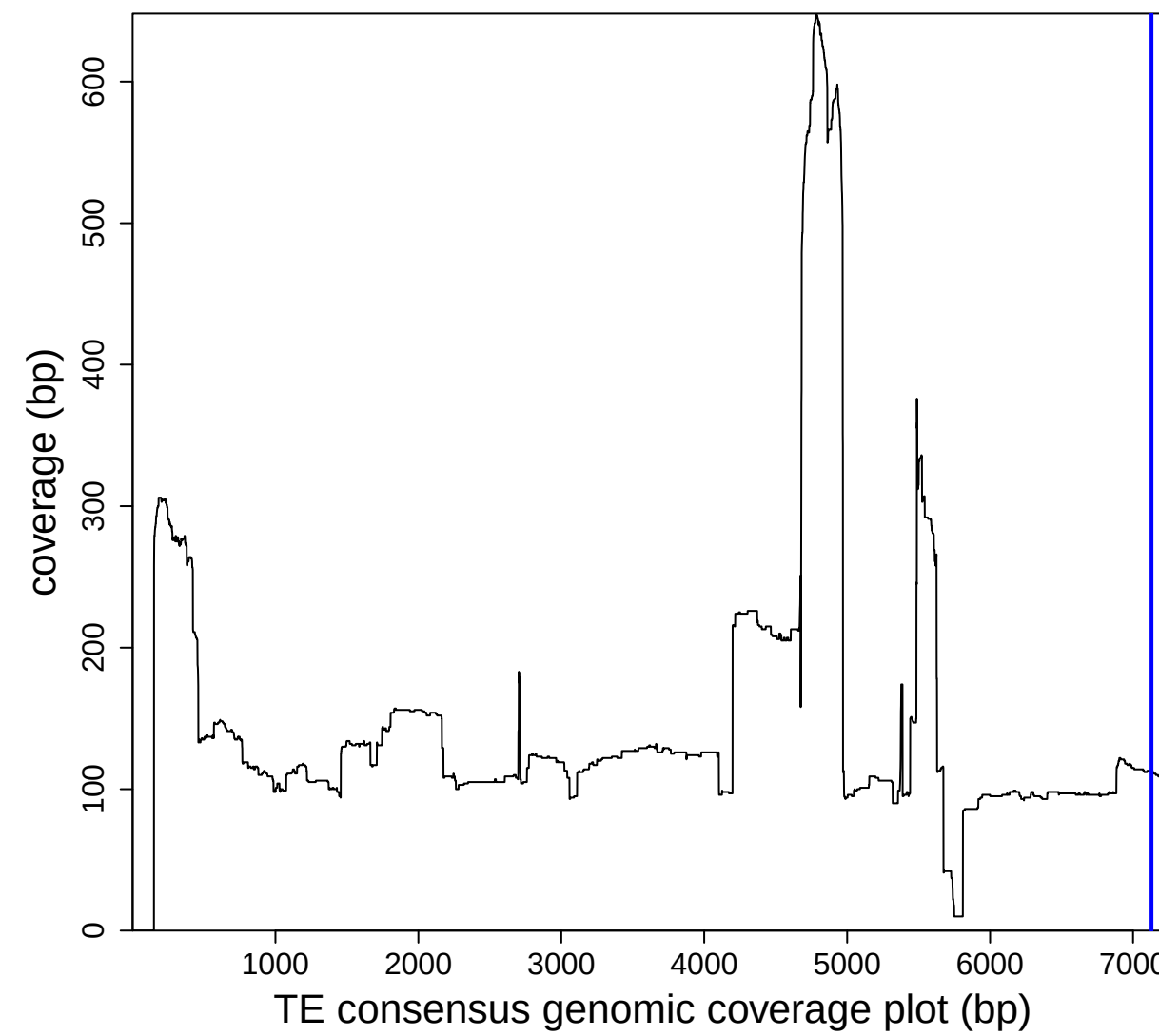
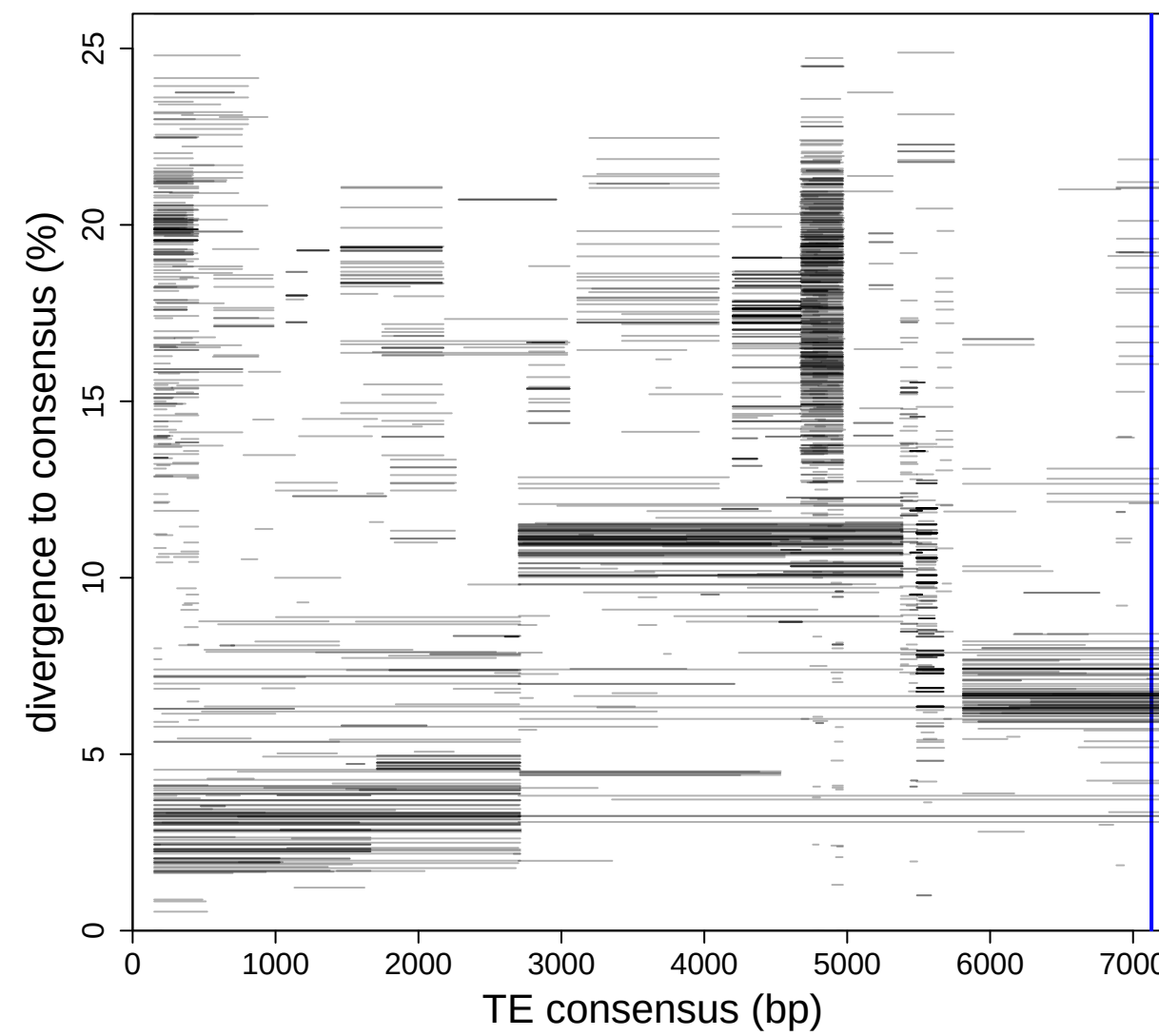
a.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 7126bp; fragments: 2105; full length: 0 (>=6413.4bp)

After TEtrimmer 7126 bp



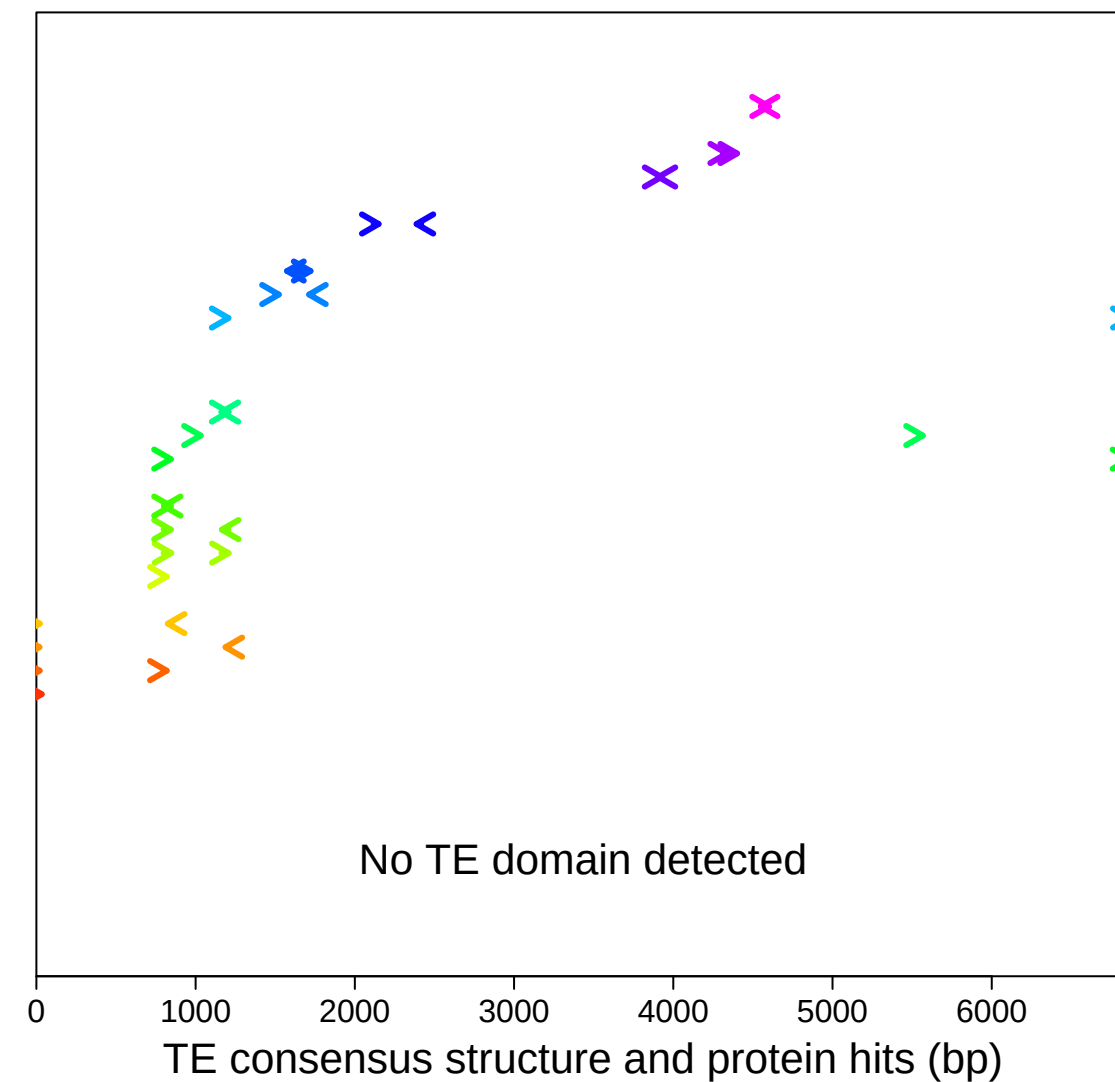
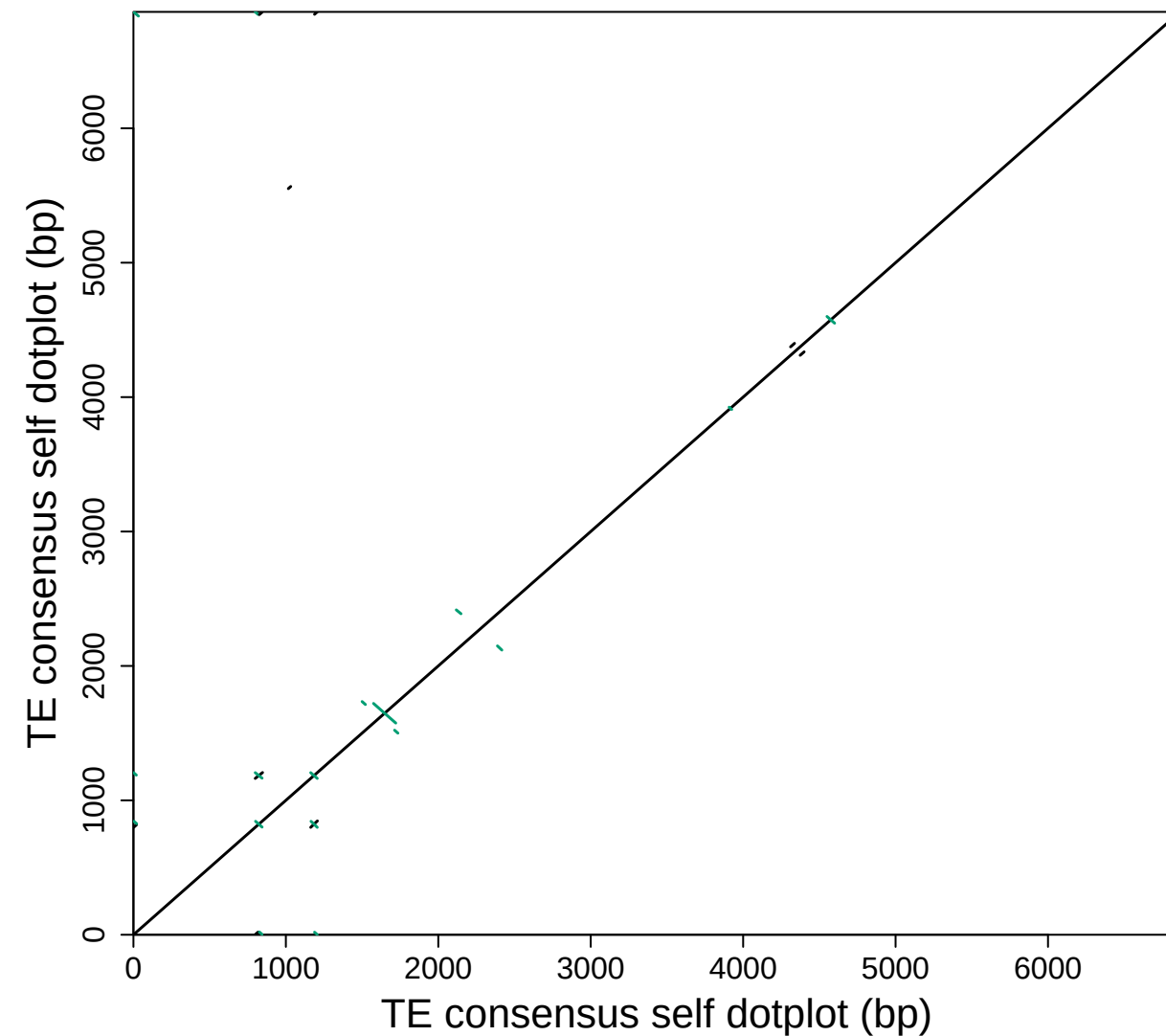
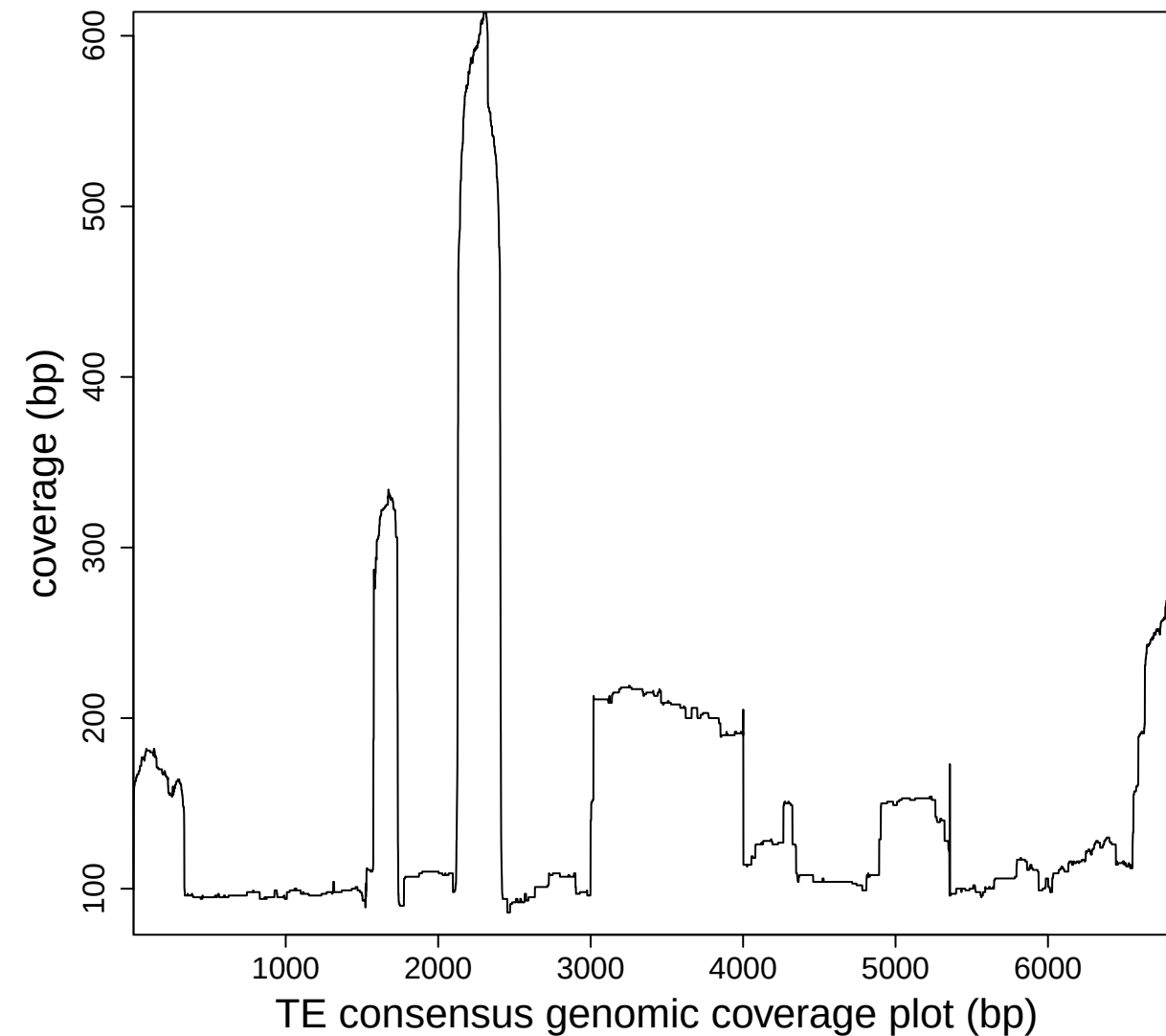
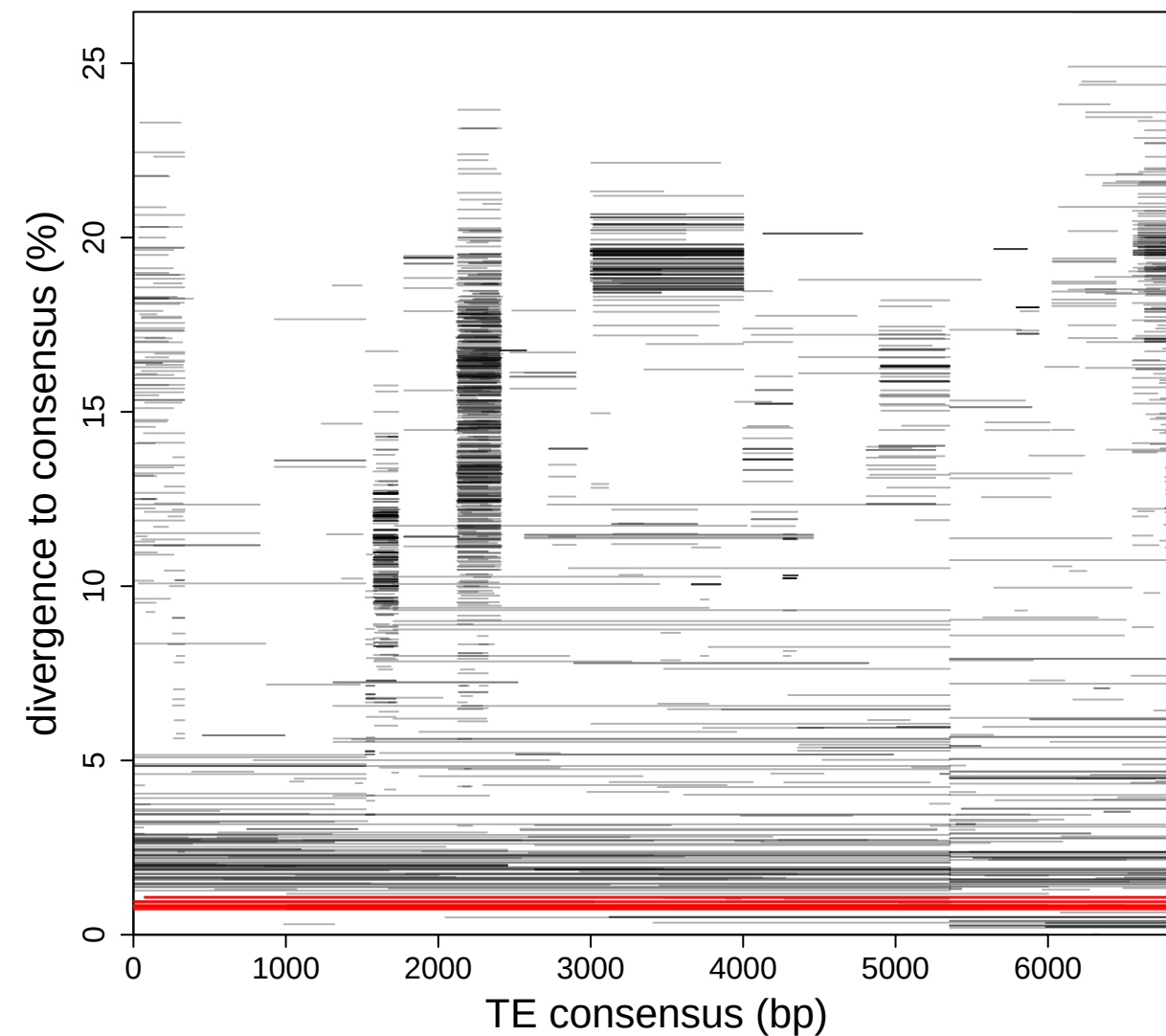
asta.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_c
size: 7279bp; fragments: 2086; full length: 0 (>=6551.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

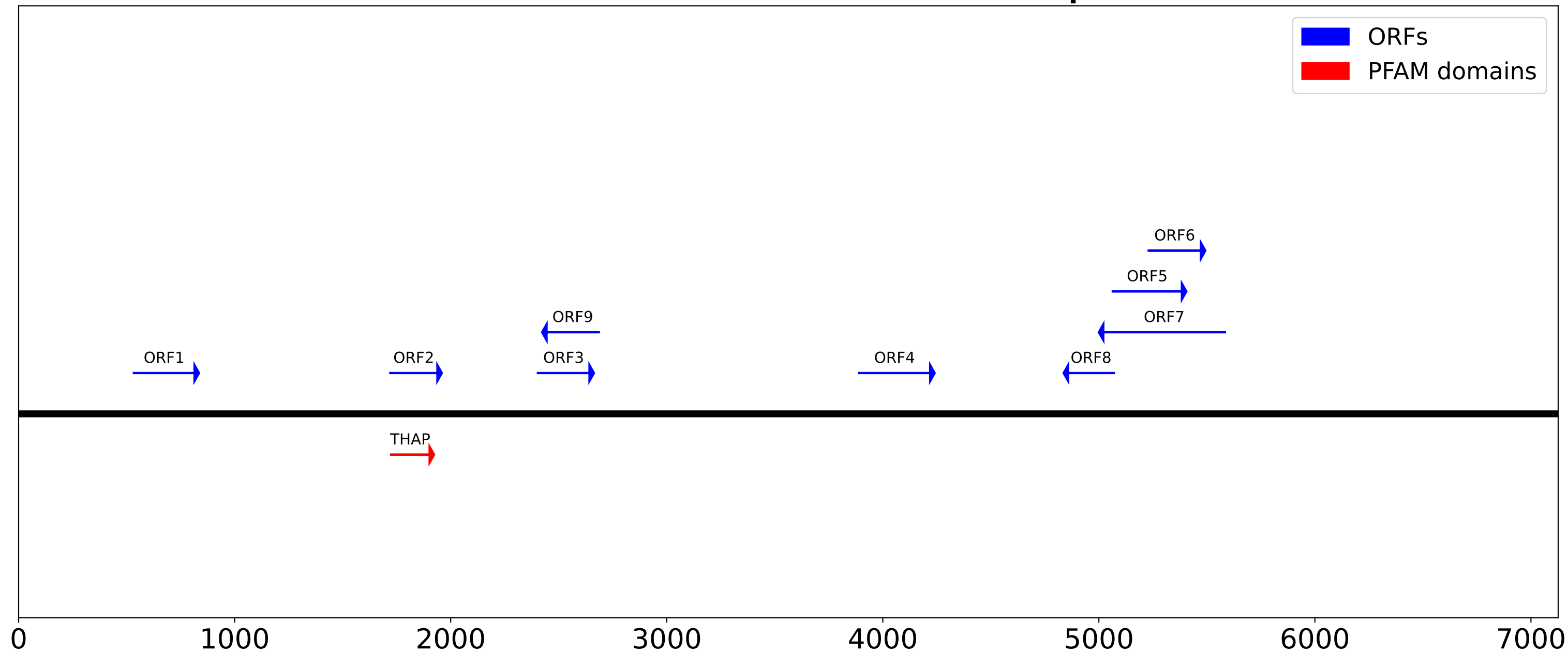


TE: rnd_5_family_3340
size: 6866bp; fragments: 1819; full length: 6 (≥ 6179.4 bp)

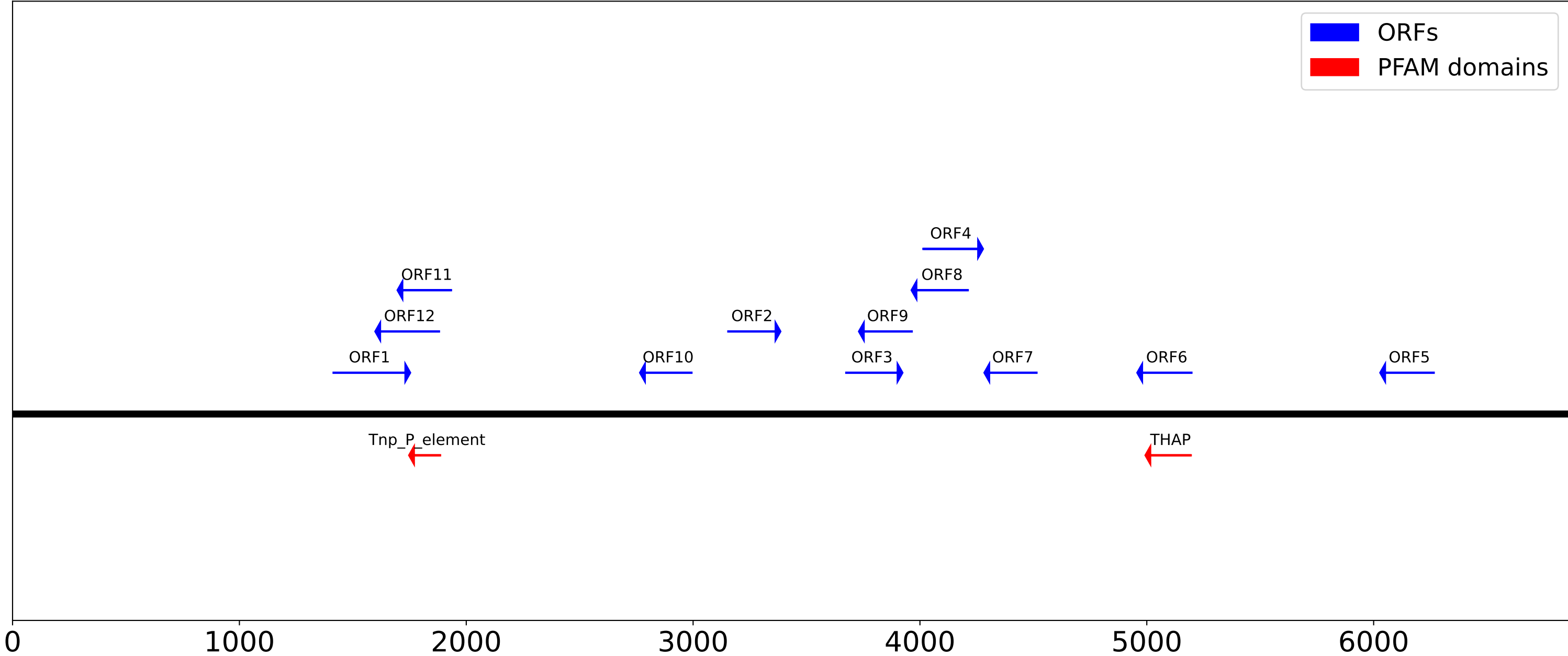
Before TEtrimmer 6866 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

