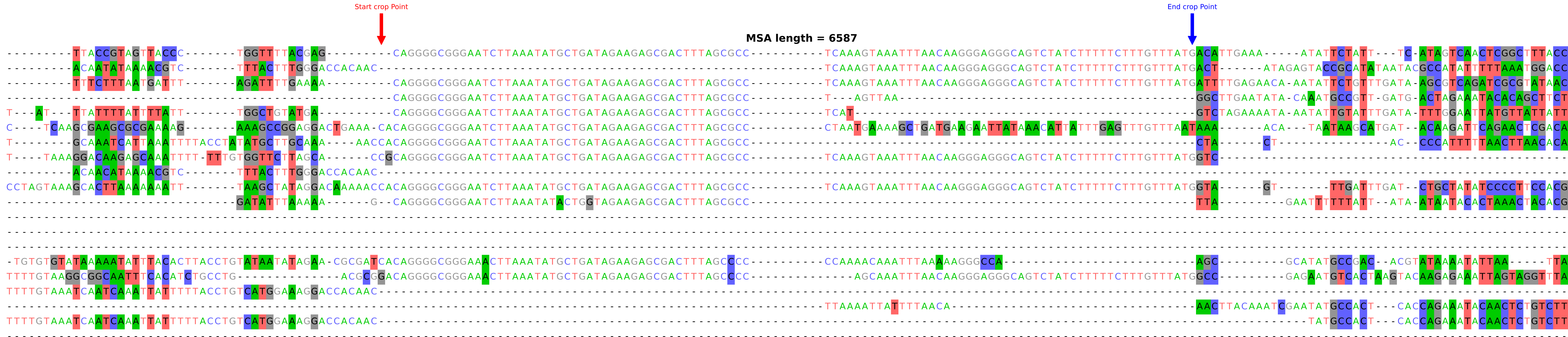




Start crop Point

End crop Point

0

10

0

1000

2000

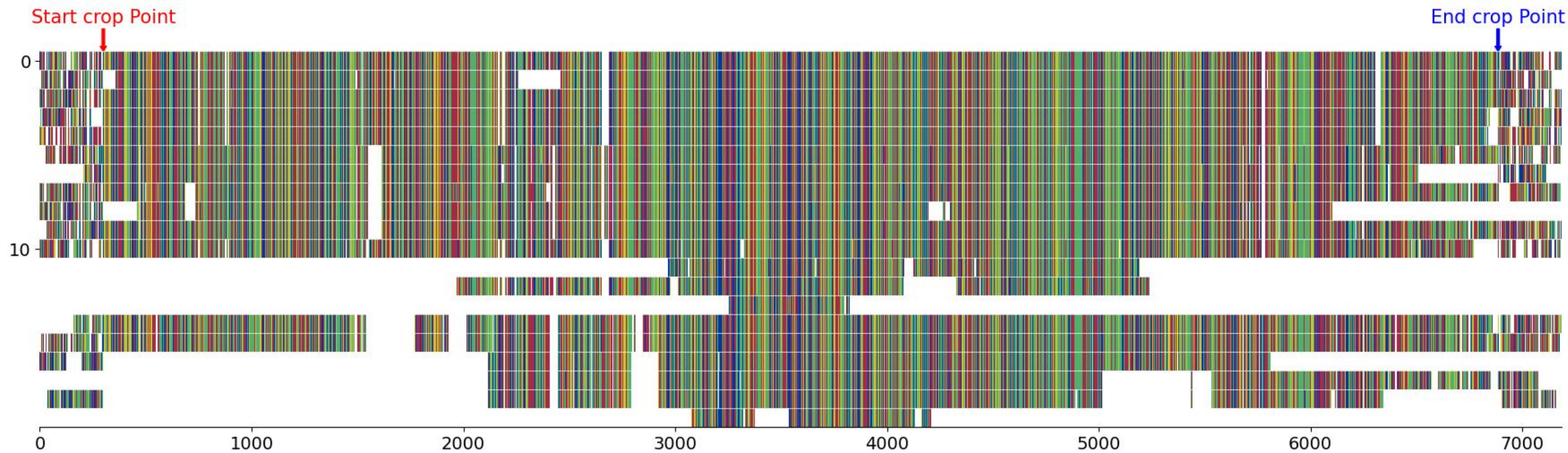
3000

4000

5000

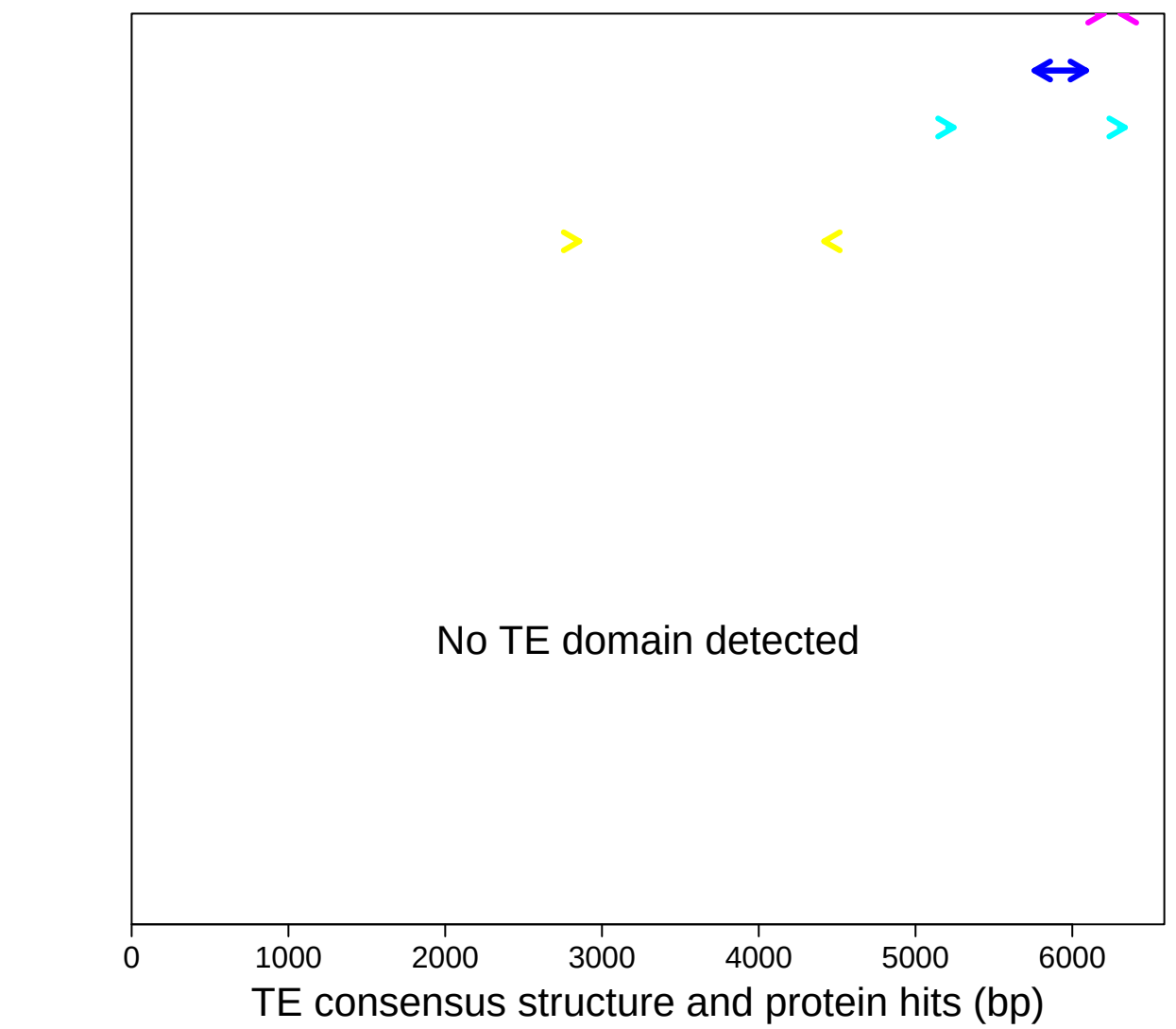
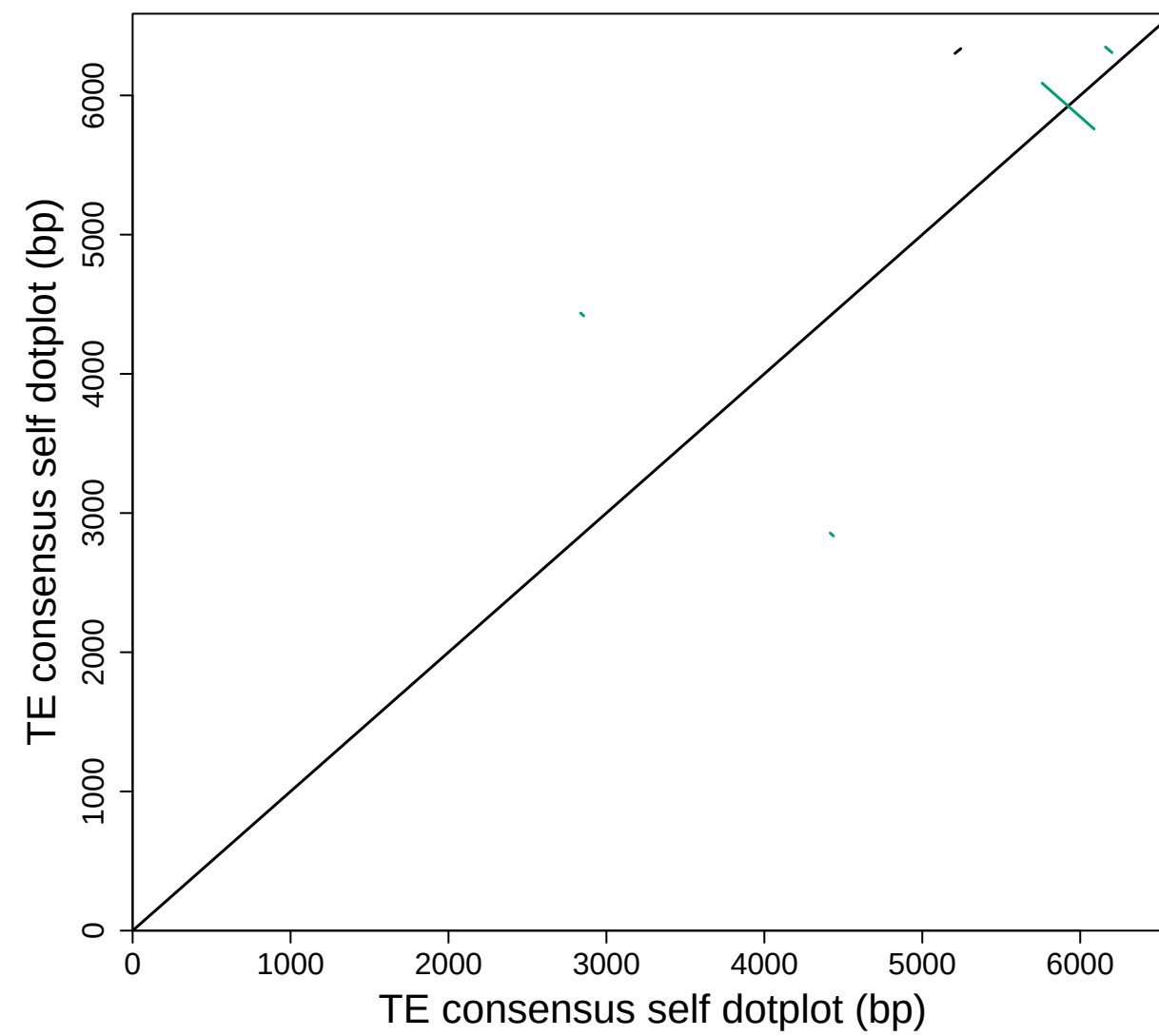
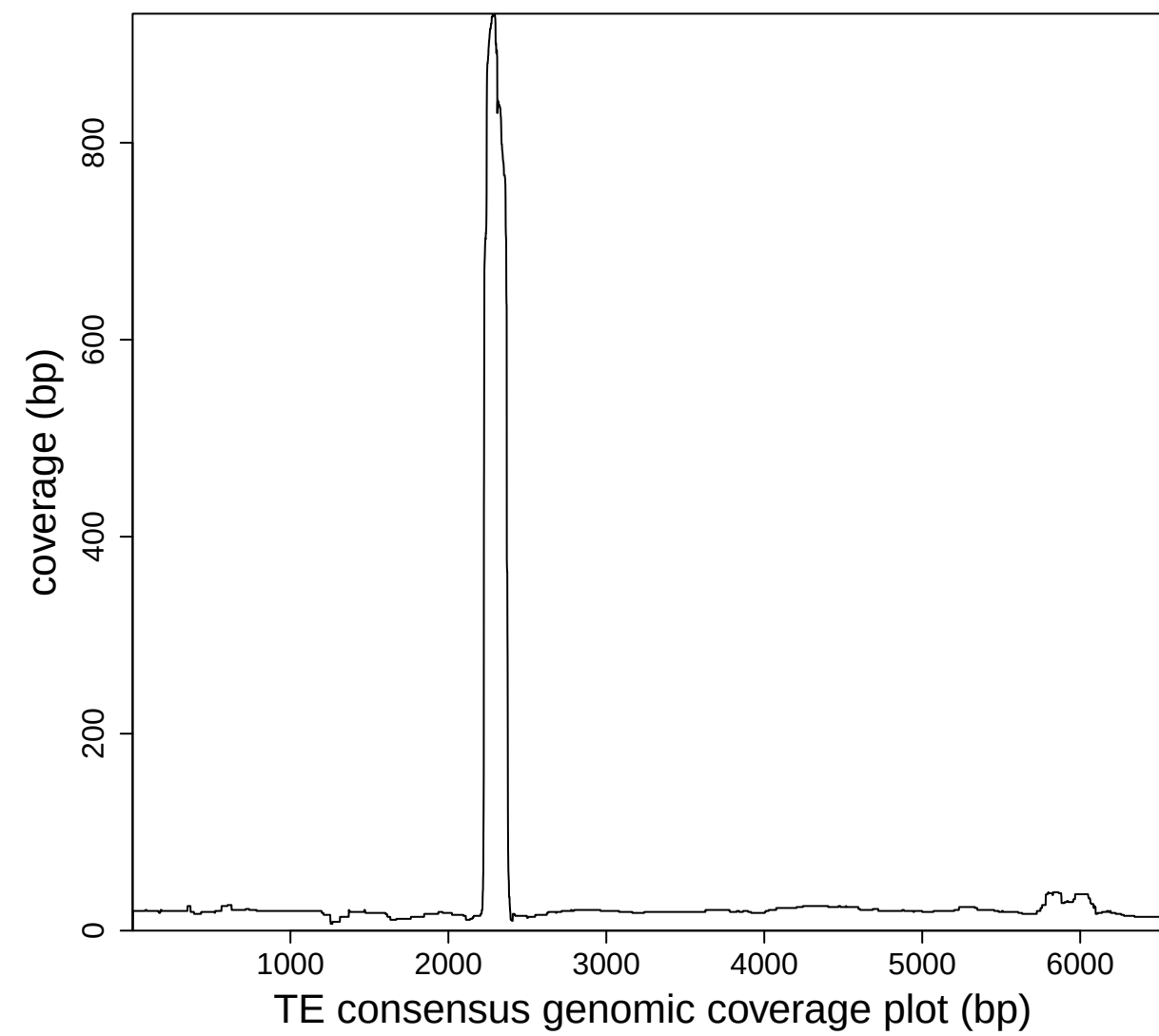
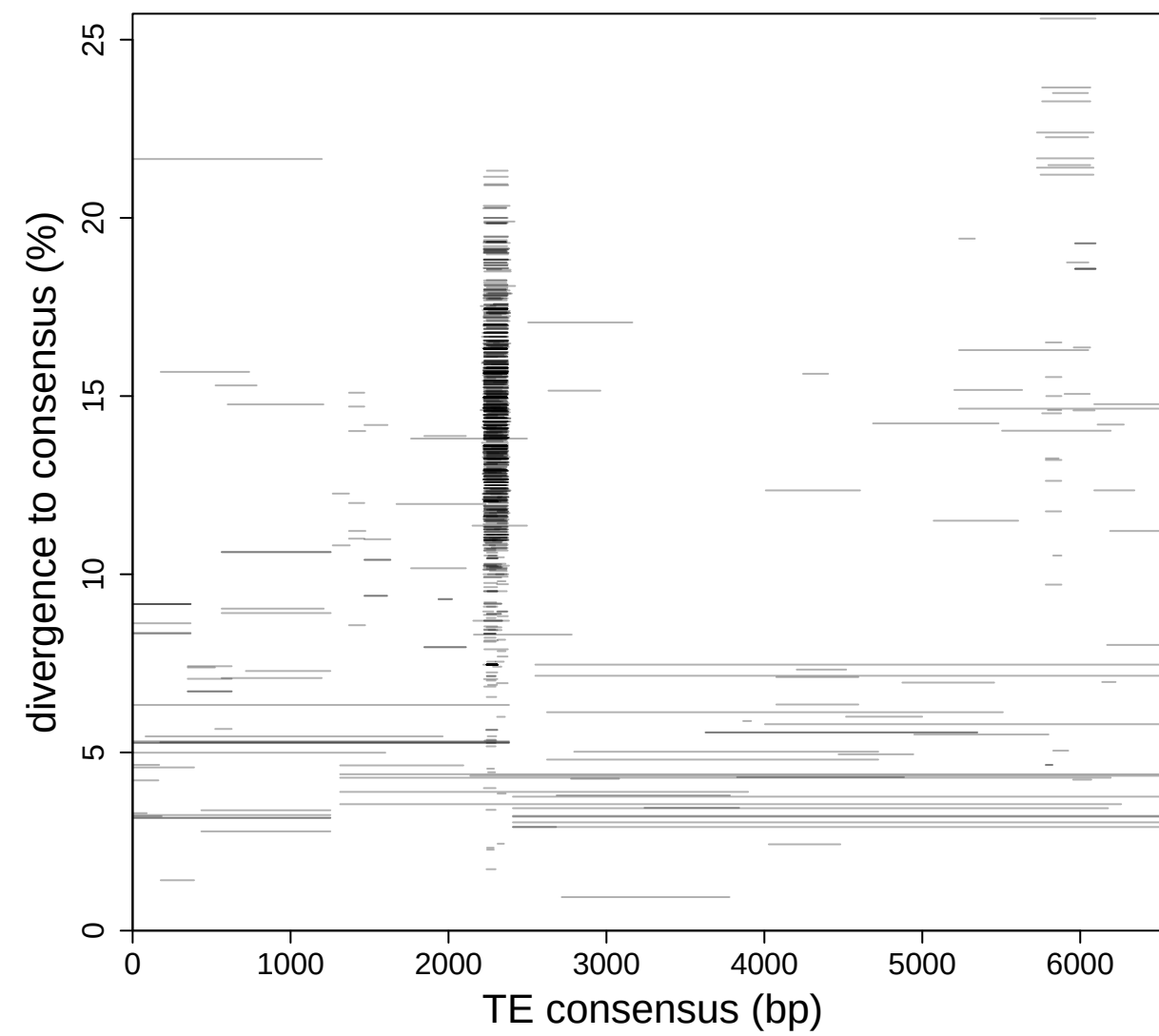
6000

7000



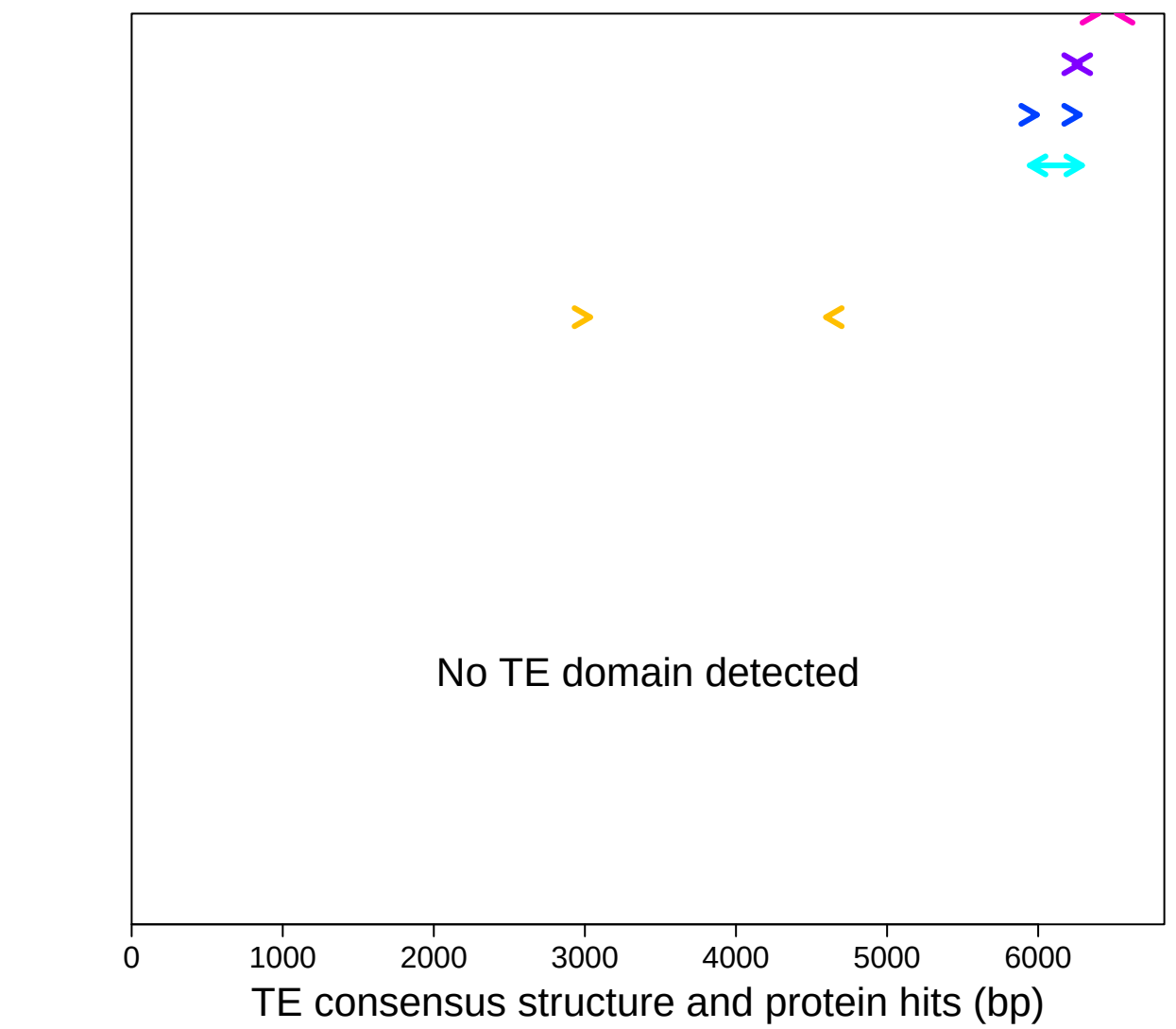
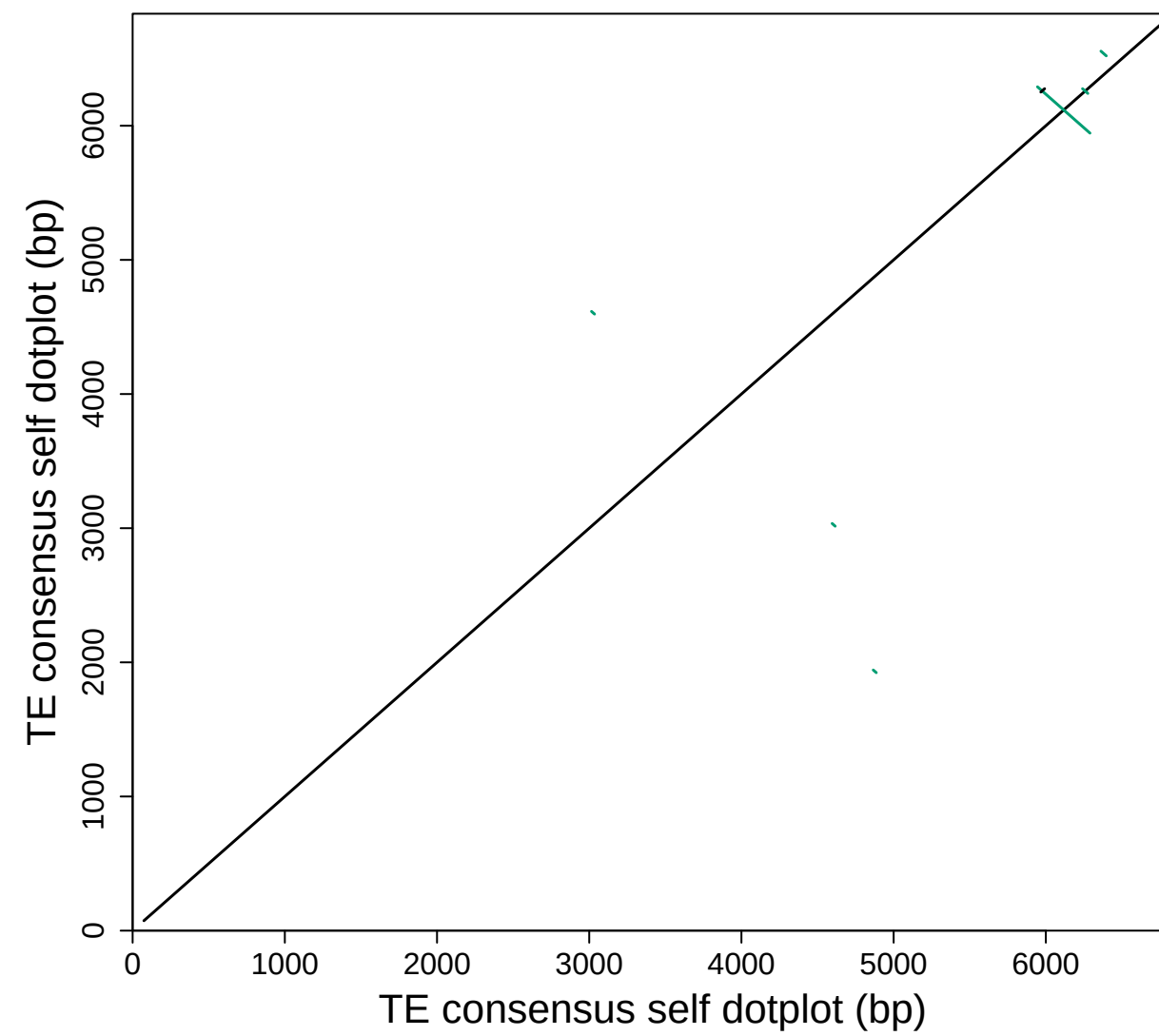
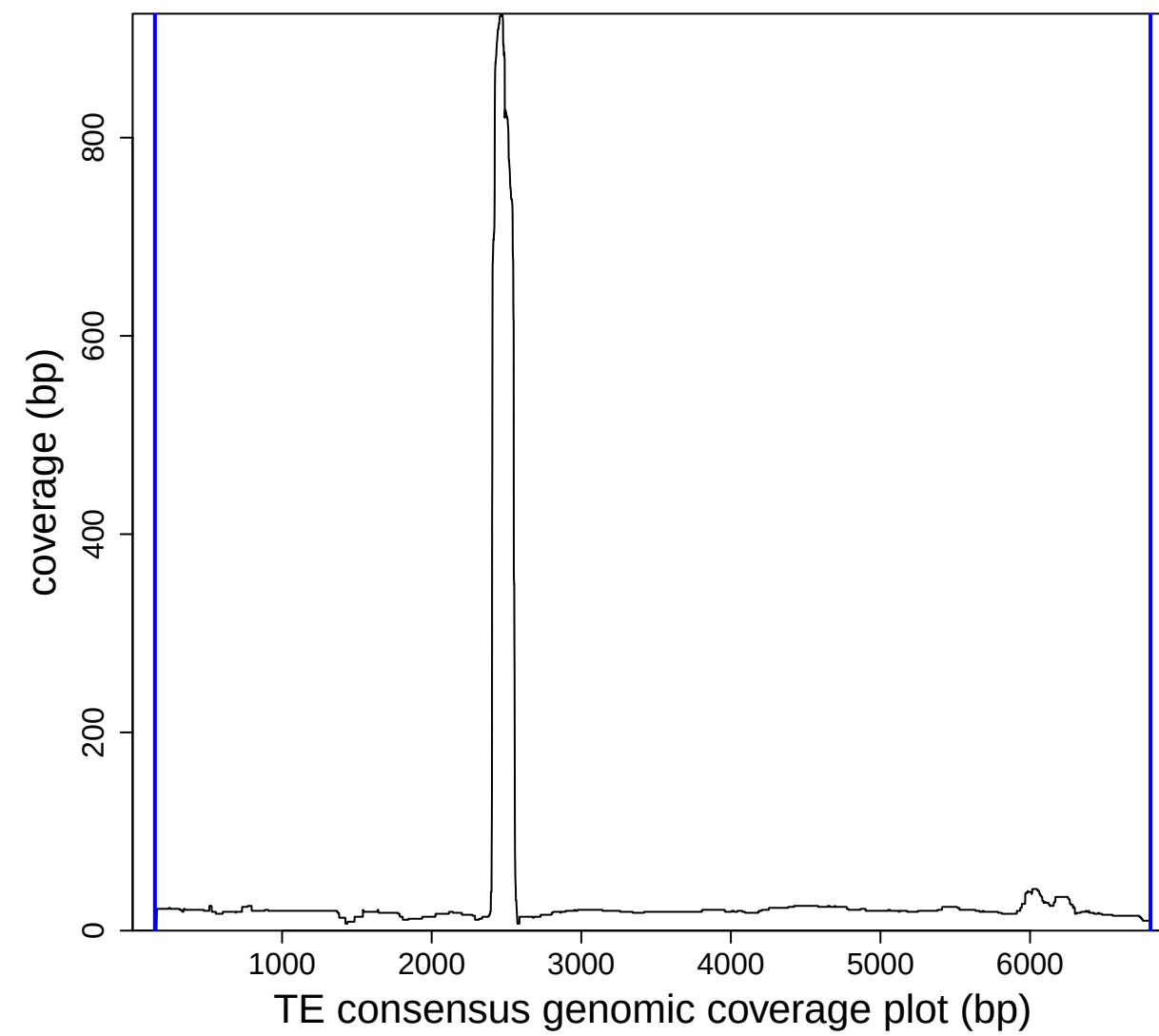
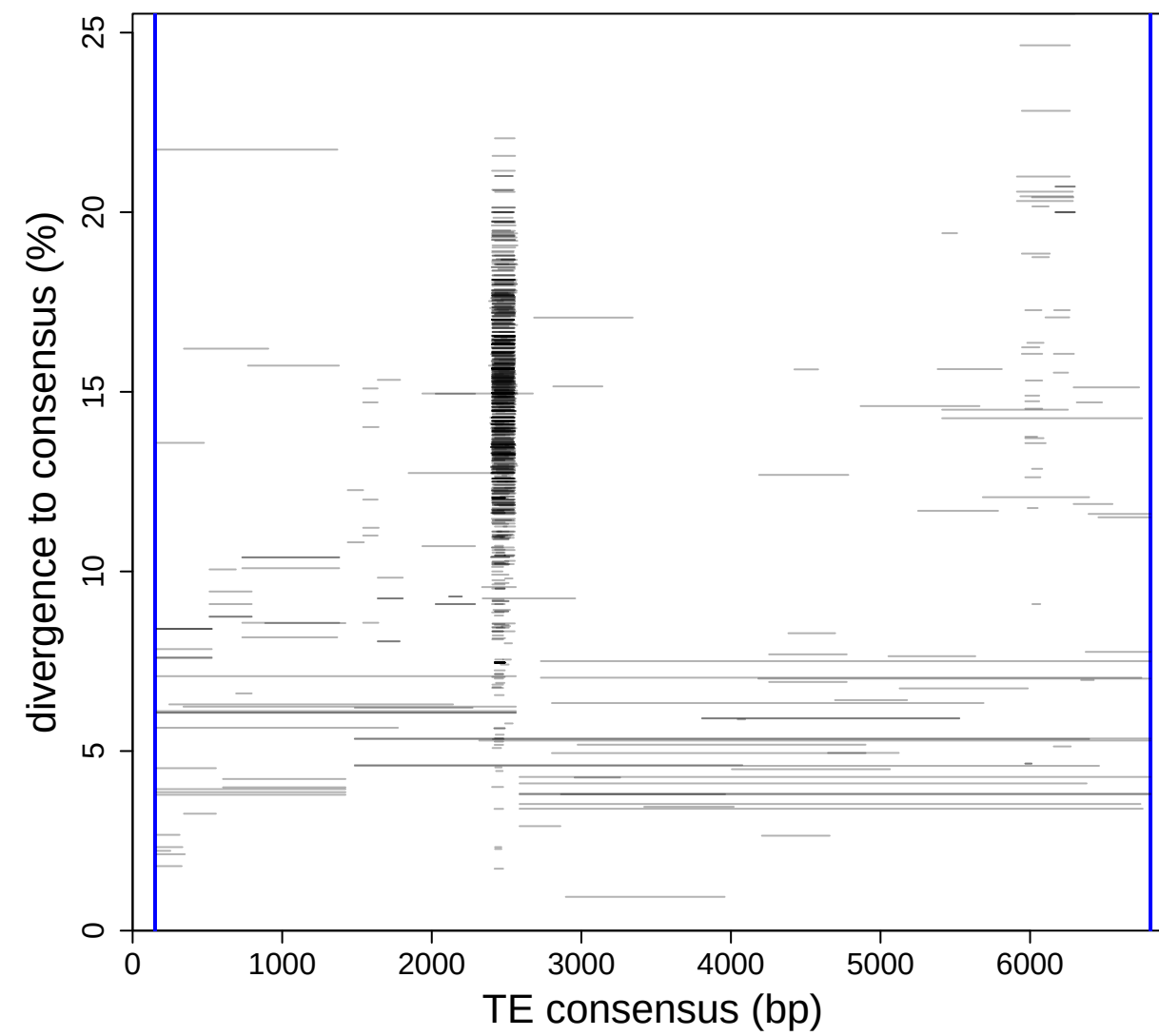
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 6587bp; fragments: 1097; full length: 0 (>=5928.3bp)

After TEtrimmer 6587 bp

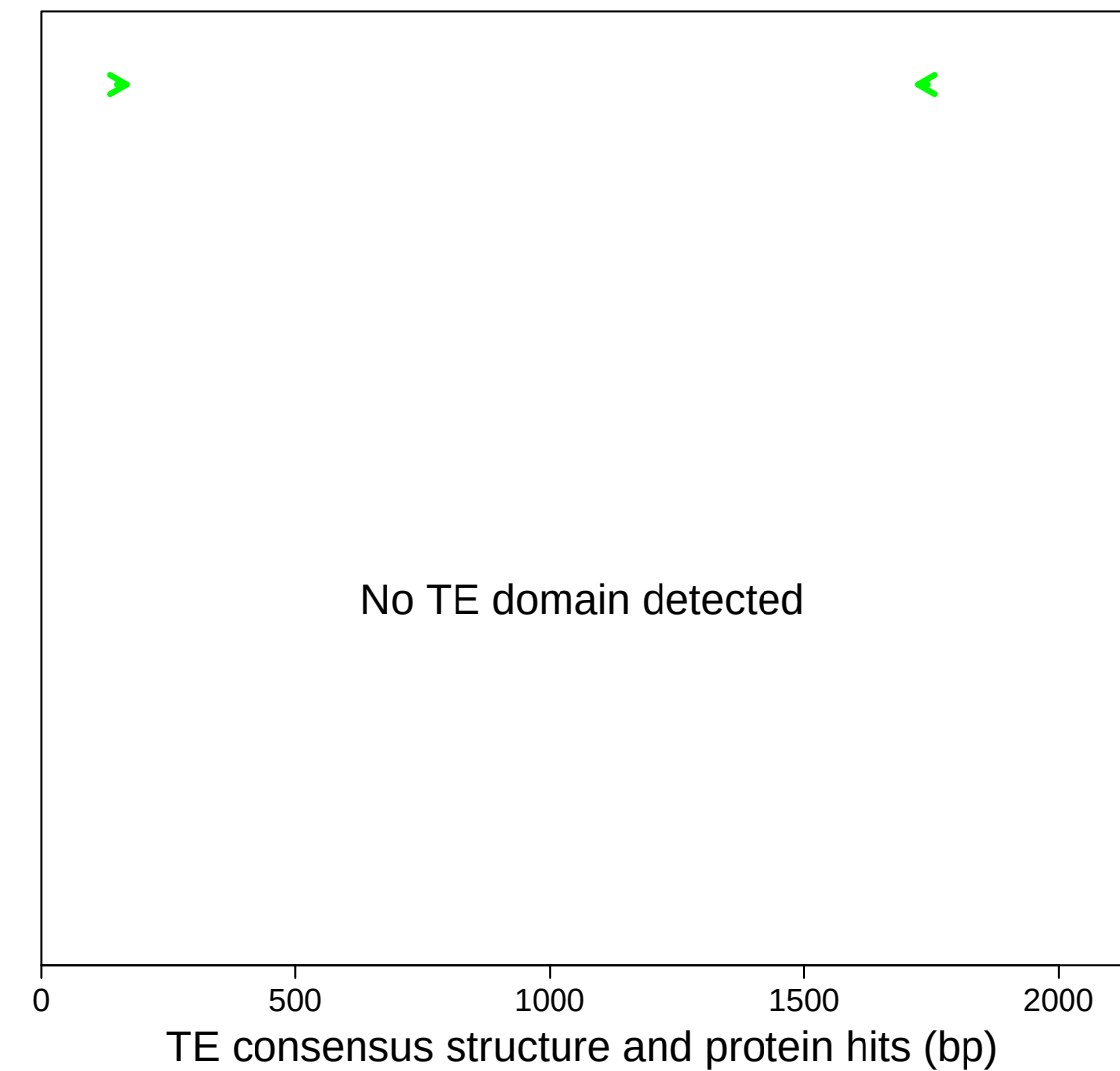
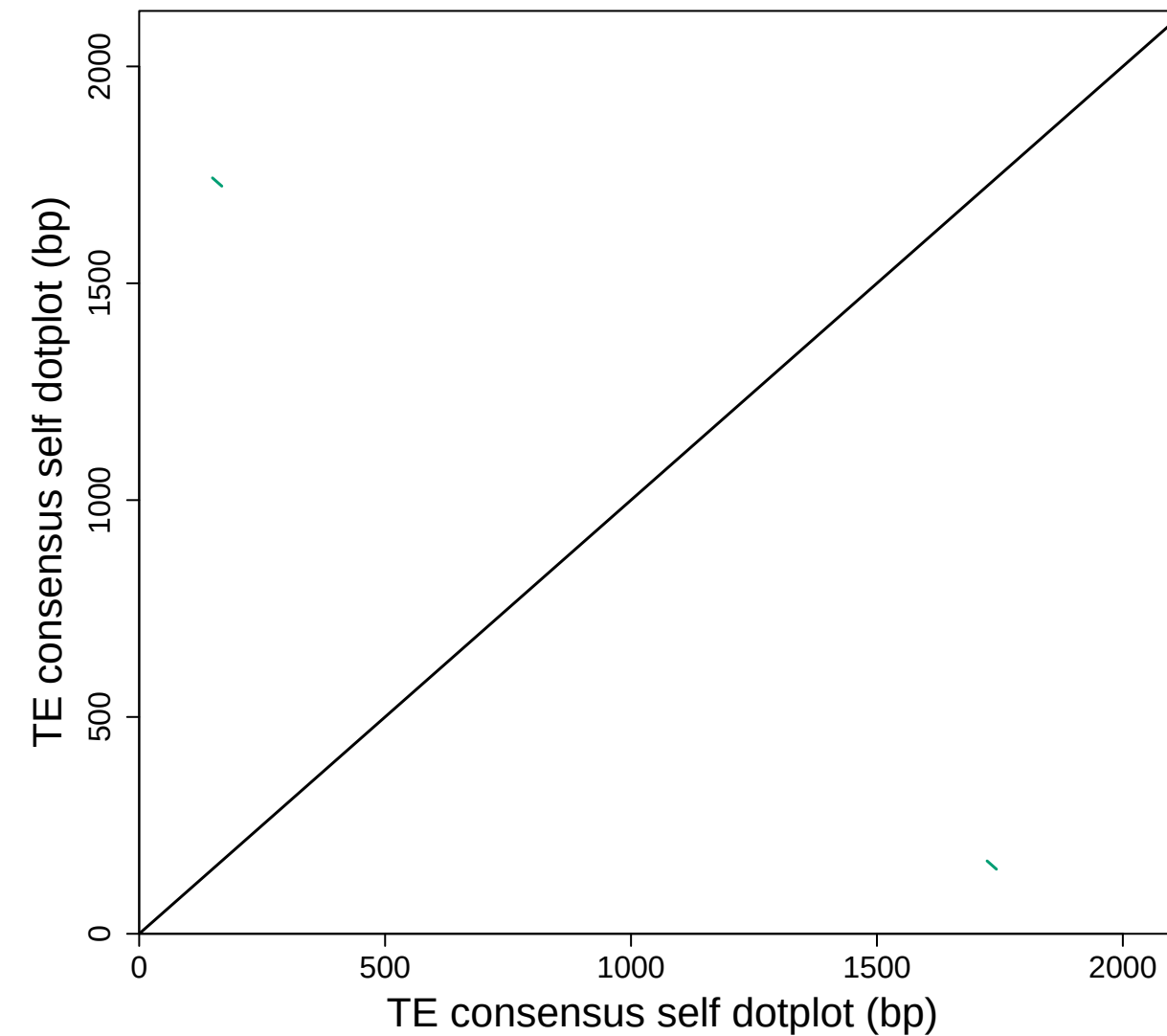
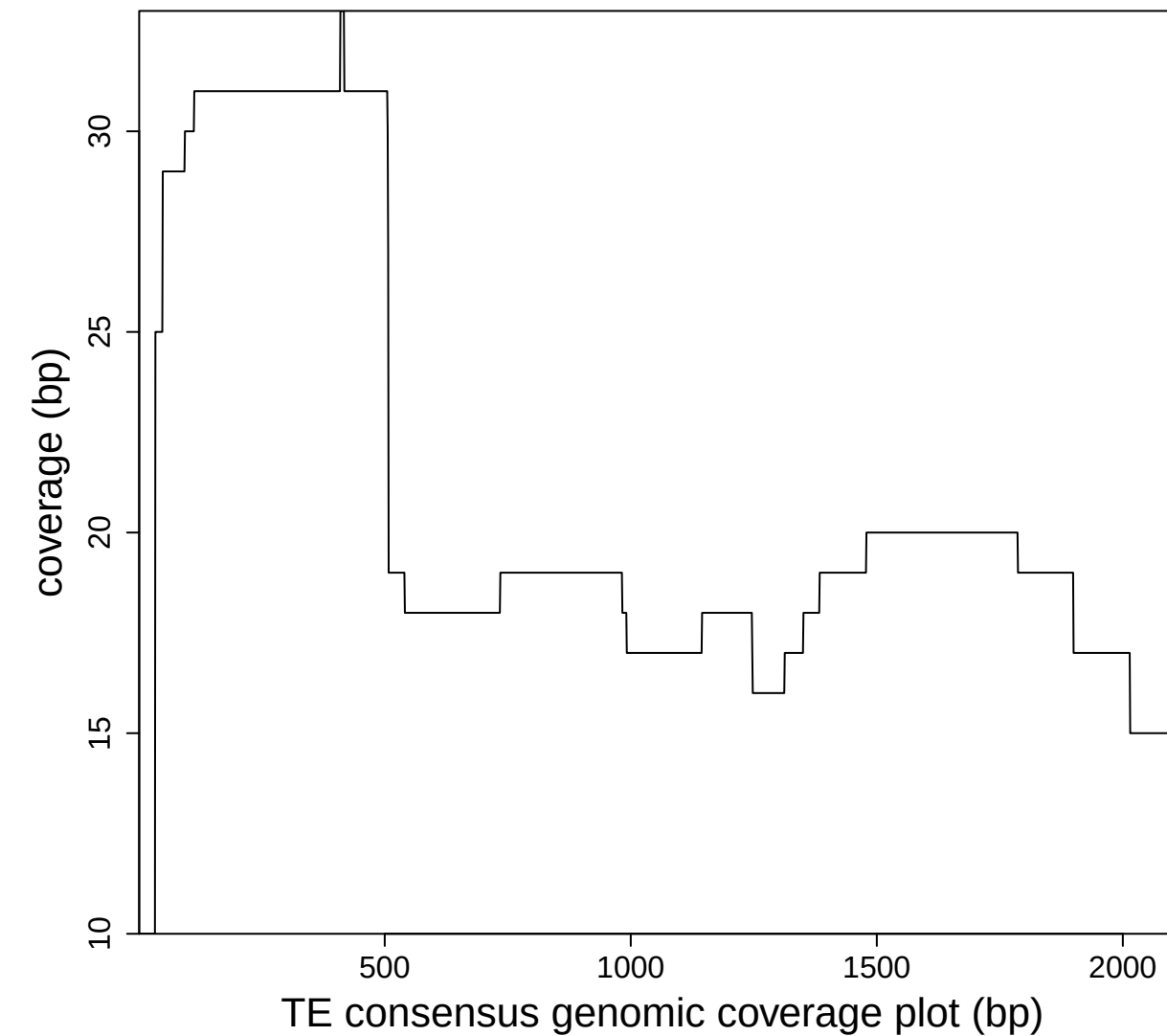
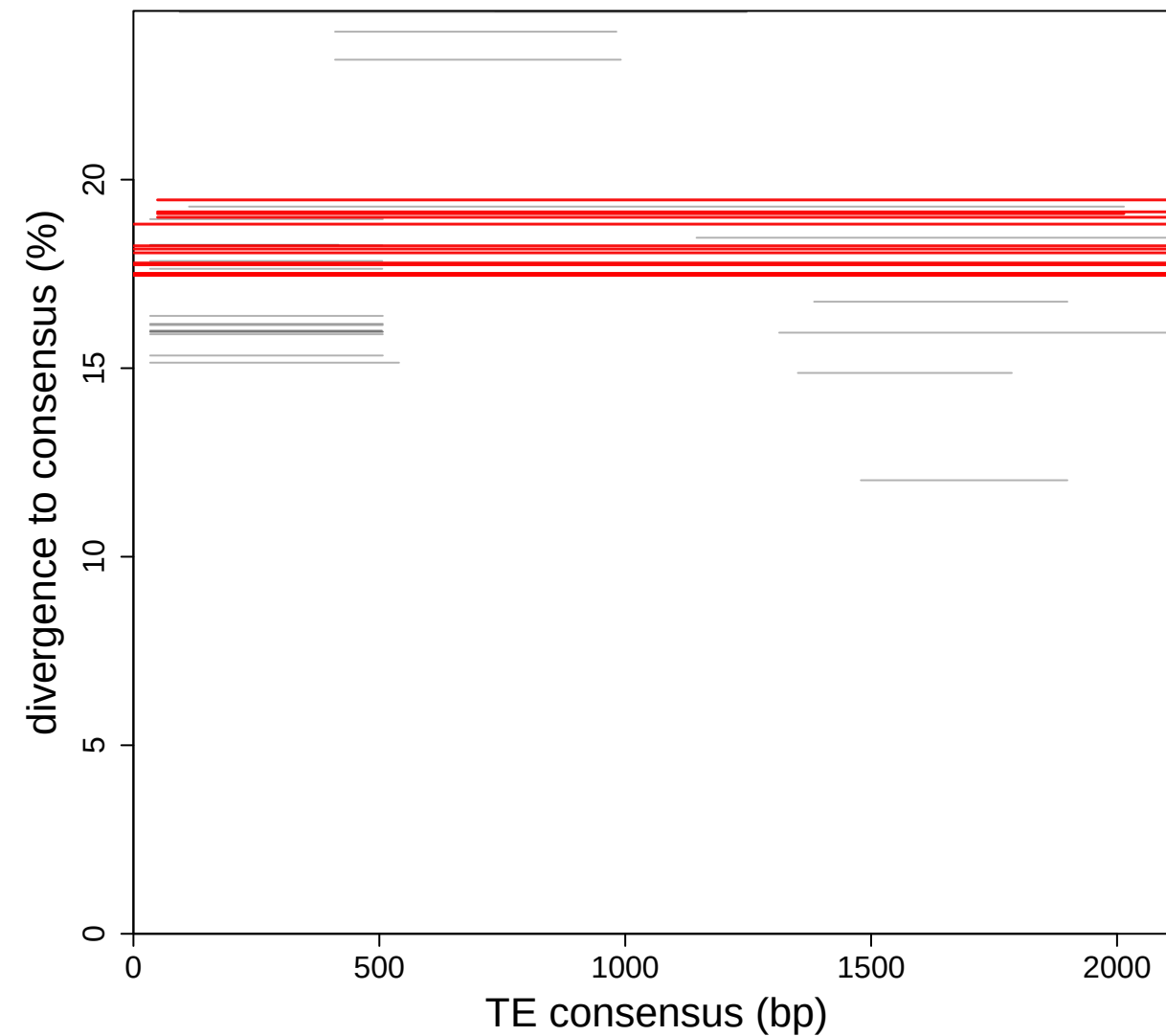


3.fasta.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcIn.fa aln.fa cl.f
size: 6955bp; fragments: 1091; full length: 0 (>=6259.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

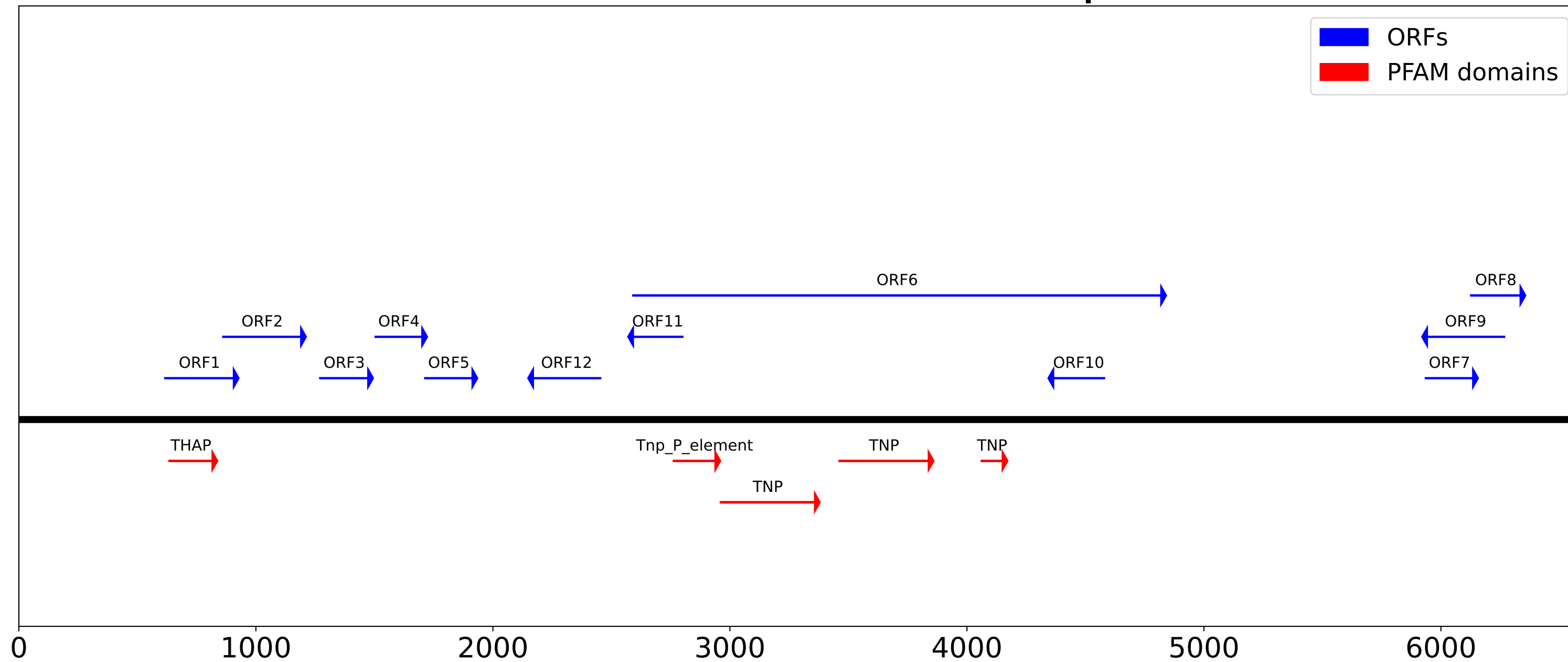


TE: rnd_5_family_7273
size: 2128bp; fragments: 39; full length: 14 (≥ 1915.2 bp)

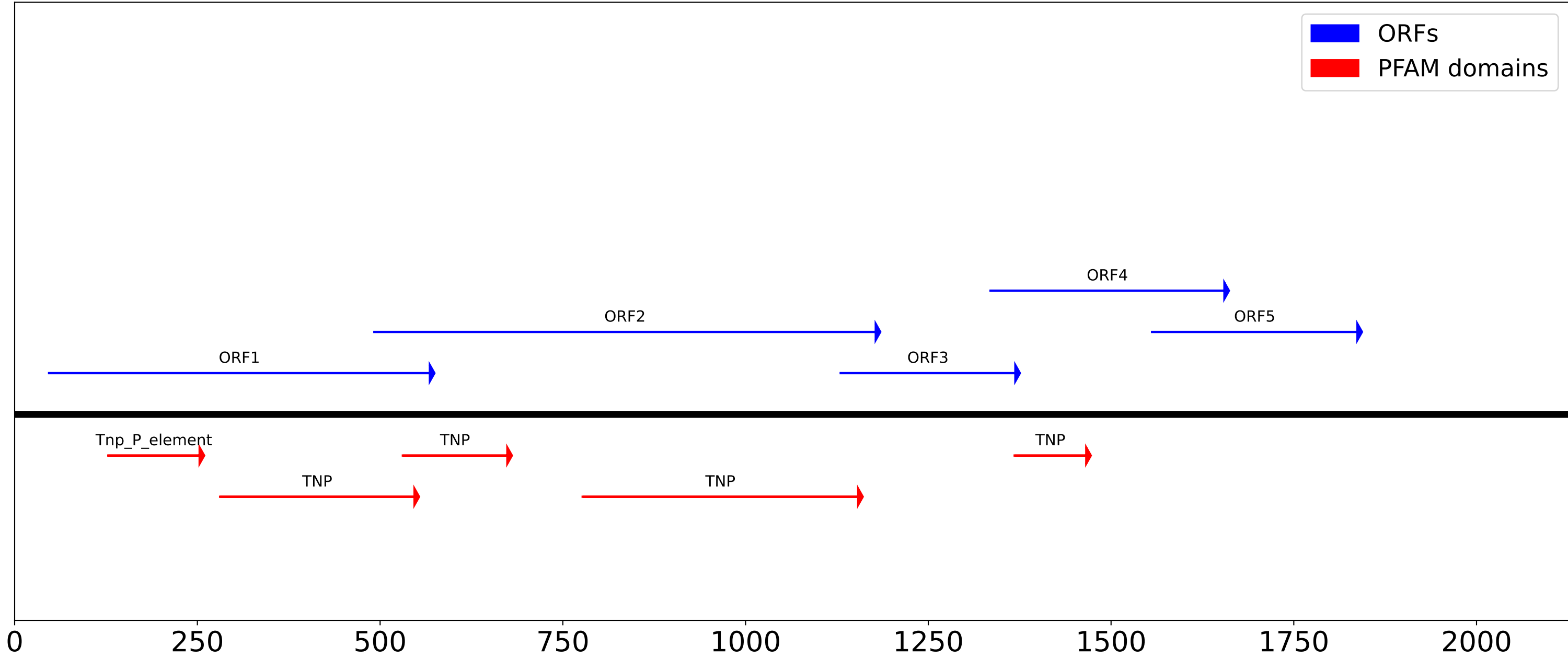


Before TEtrimmer 2128 bp

After TEtrimmer ORF and PFAM domain plot



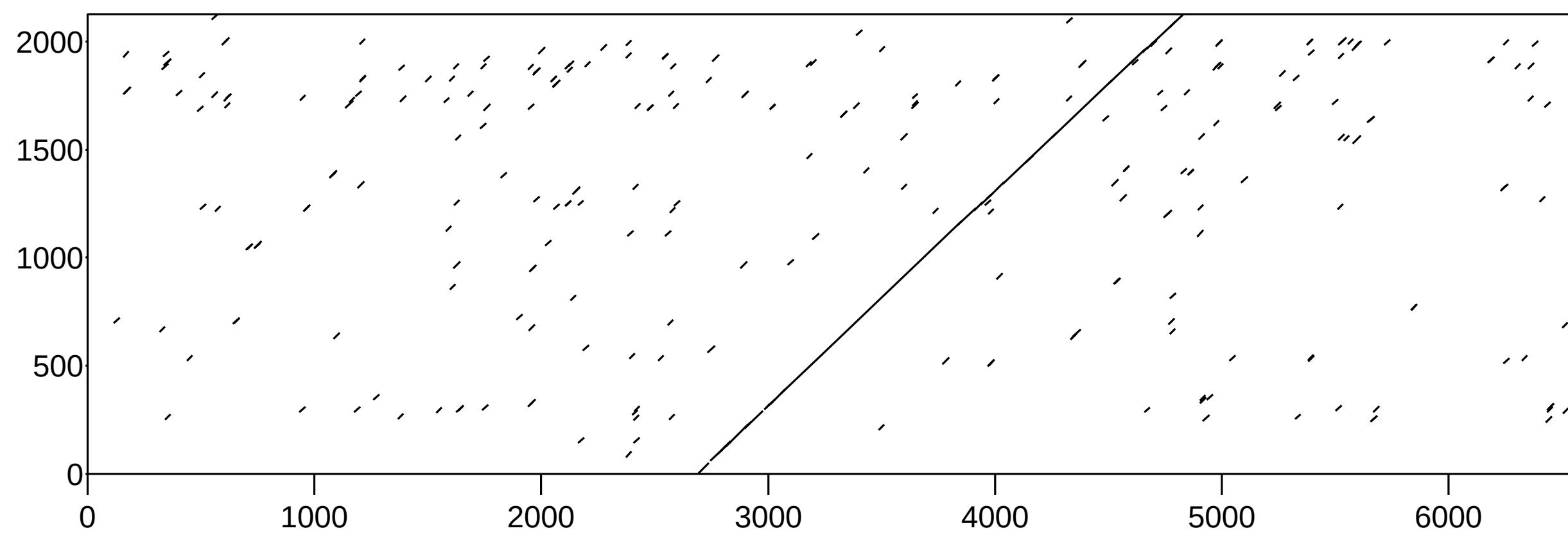
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)