

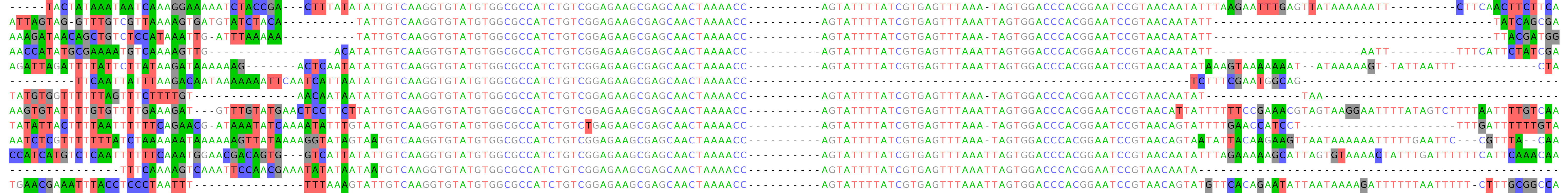
Start crop Point



End crop Point



MSA length = 5963



Start crop Point

End crop Point

0-

10

0

1000

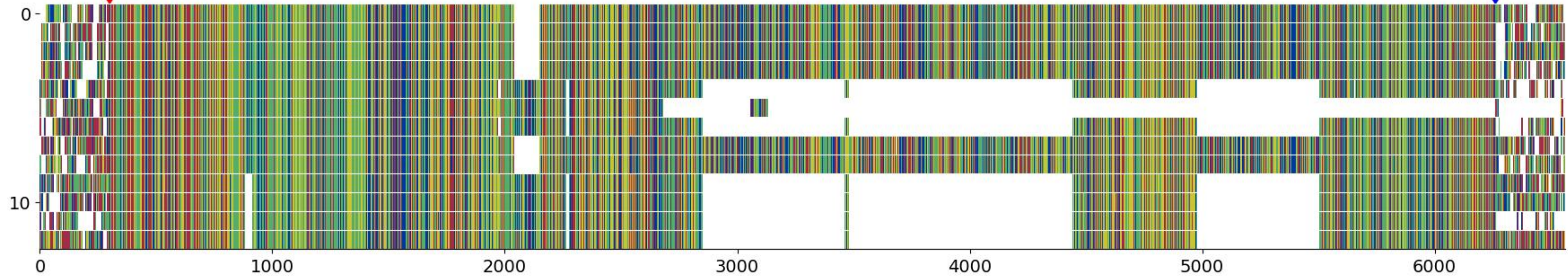
2000

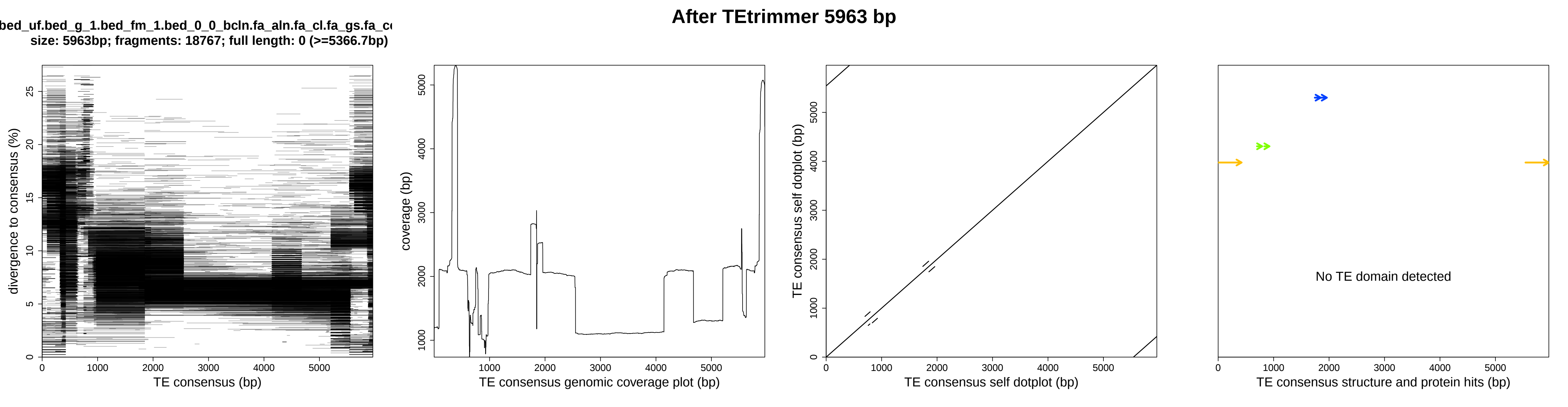
3000

4000

5000

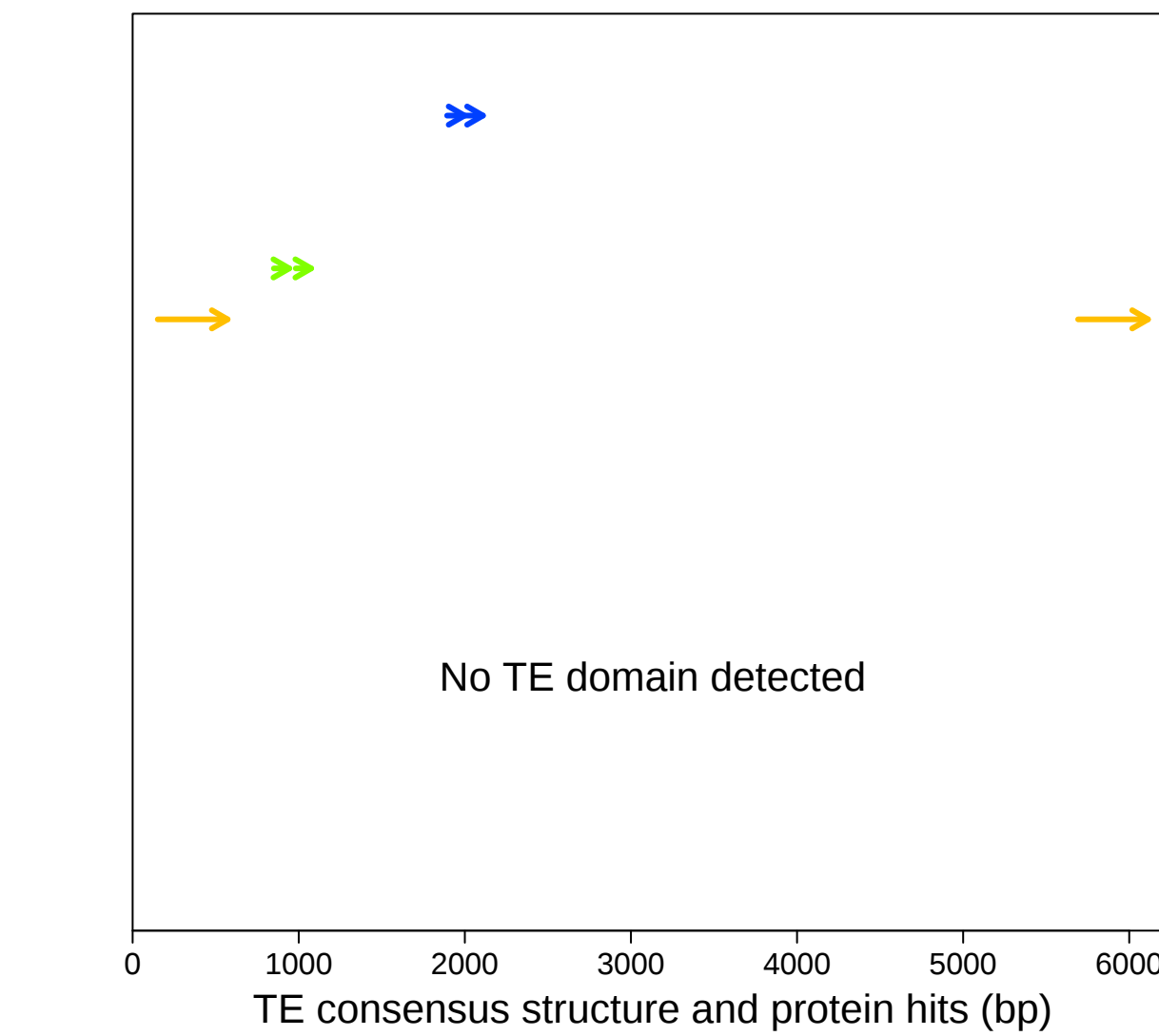
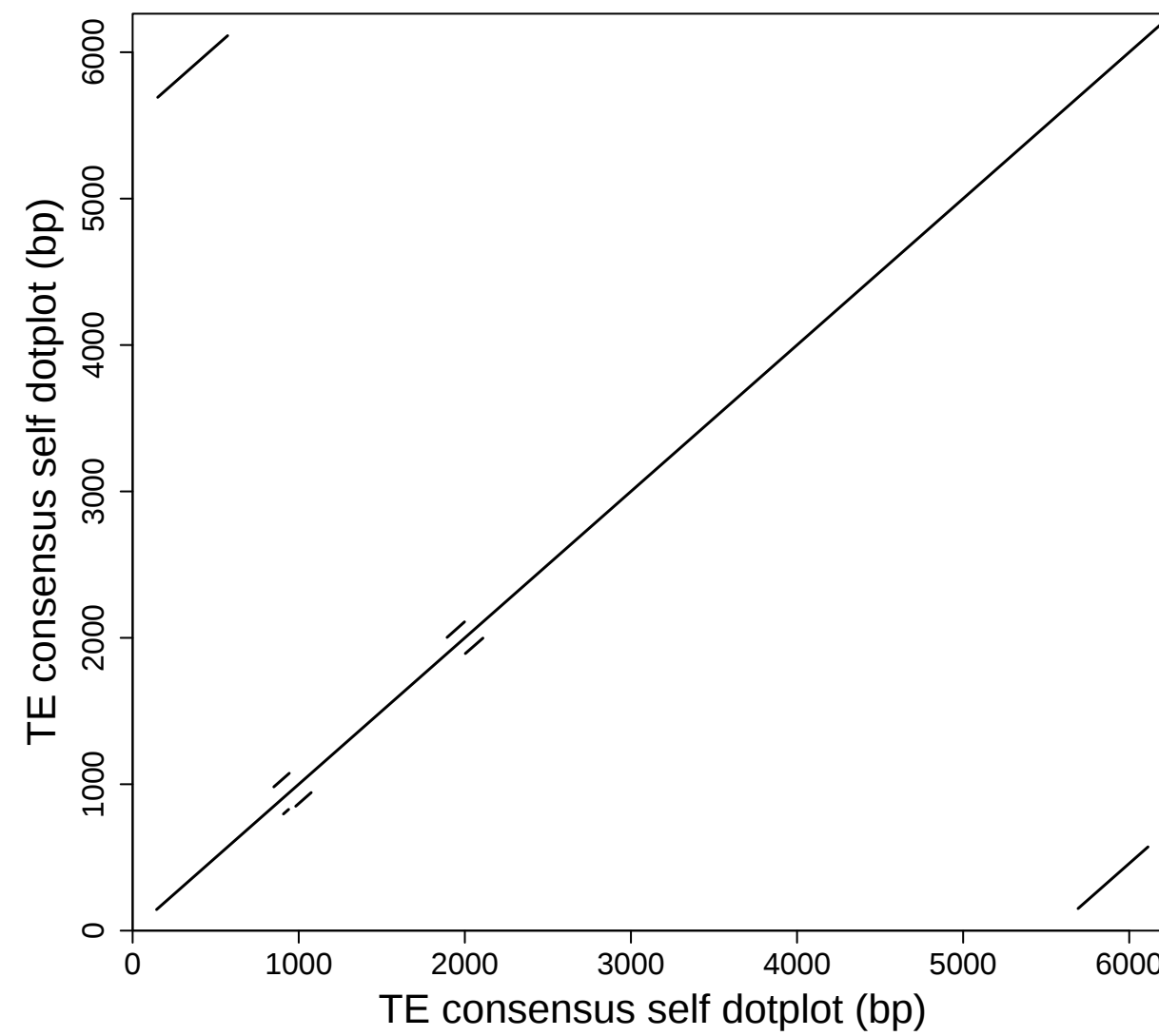
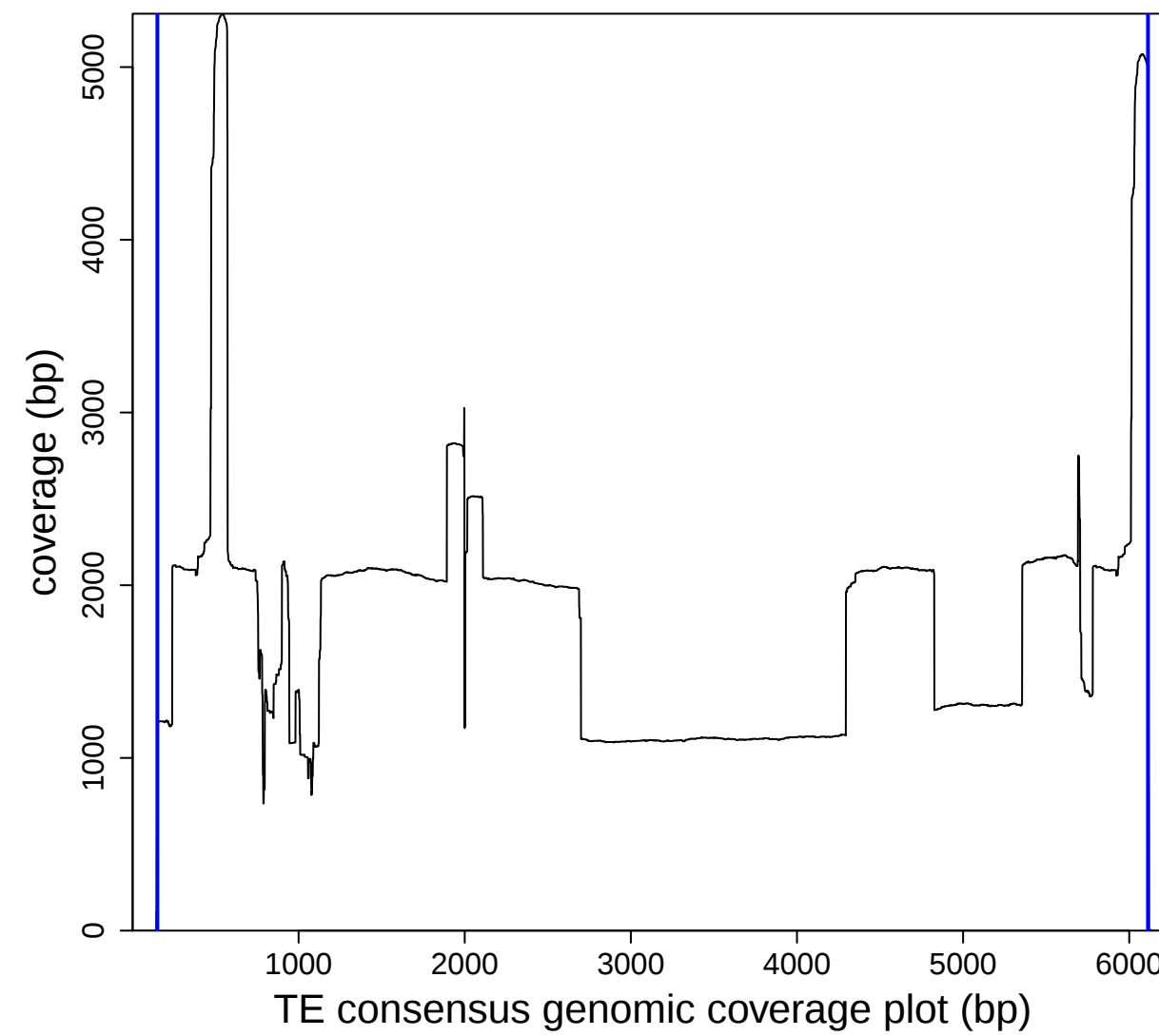
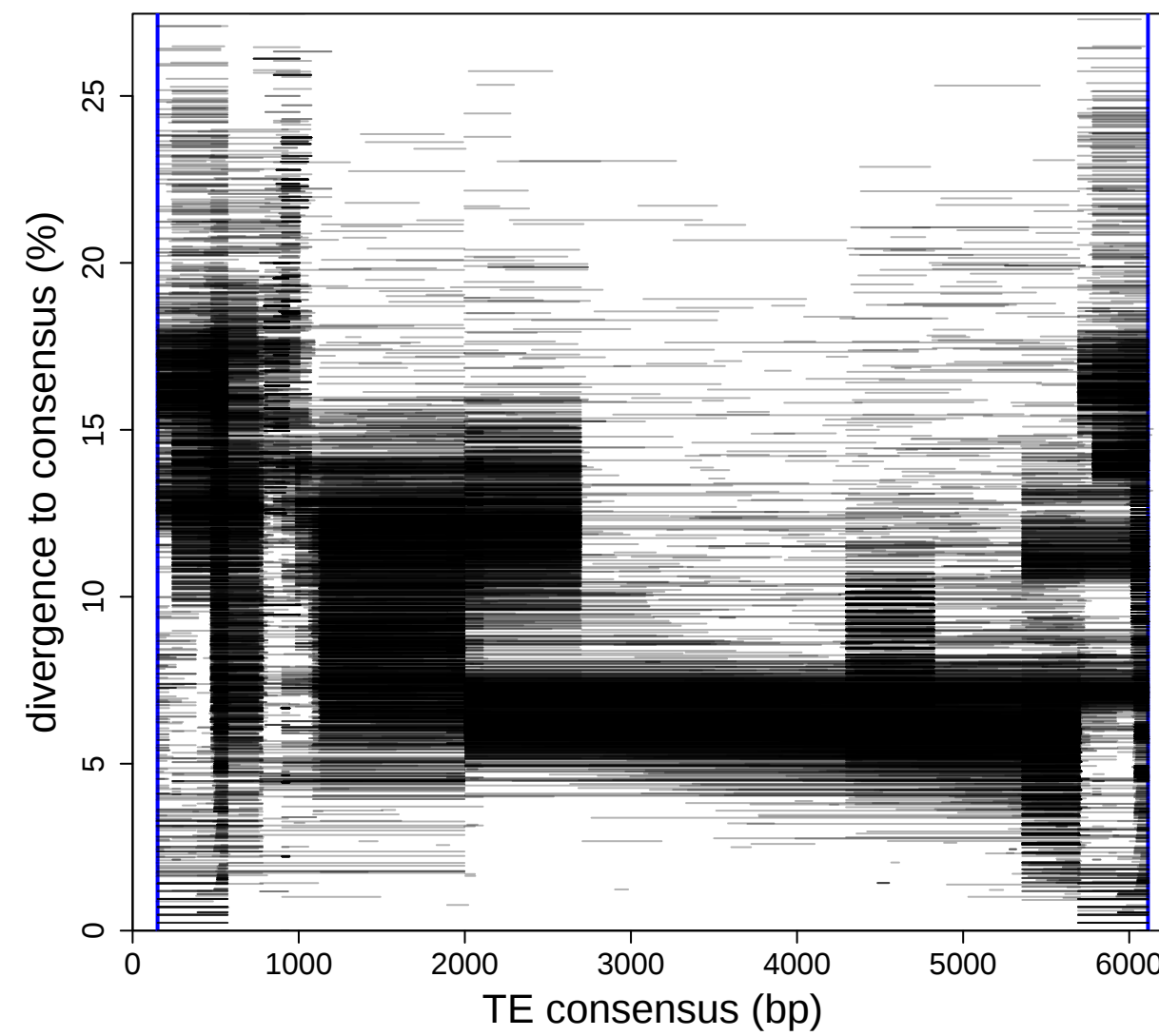
6000



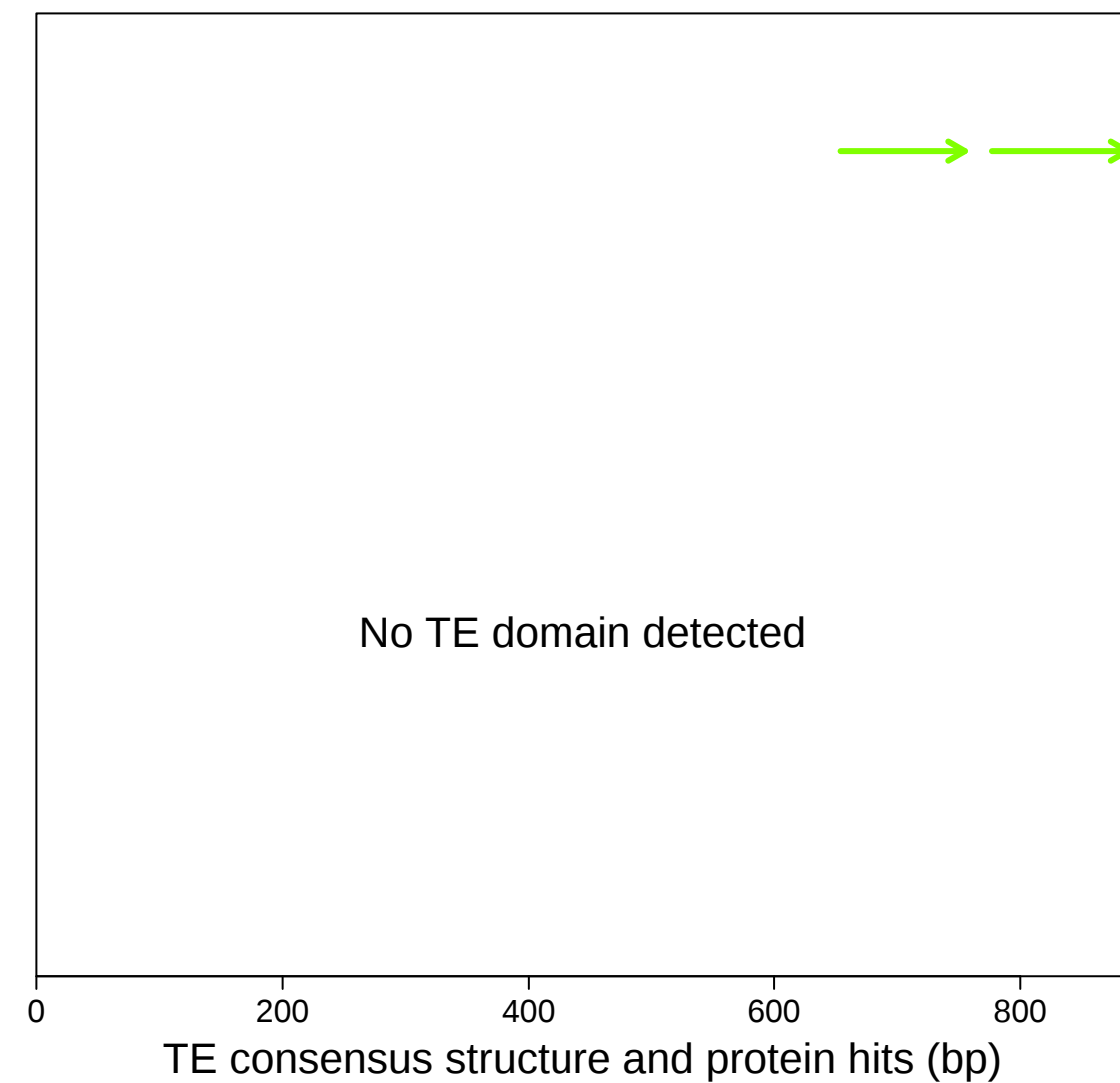
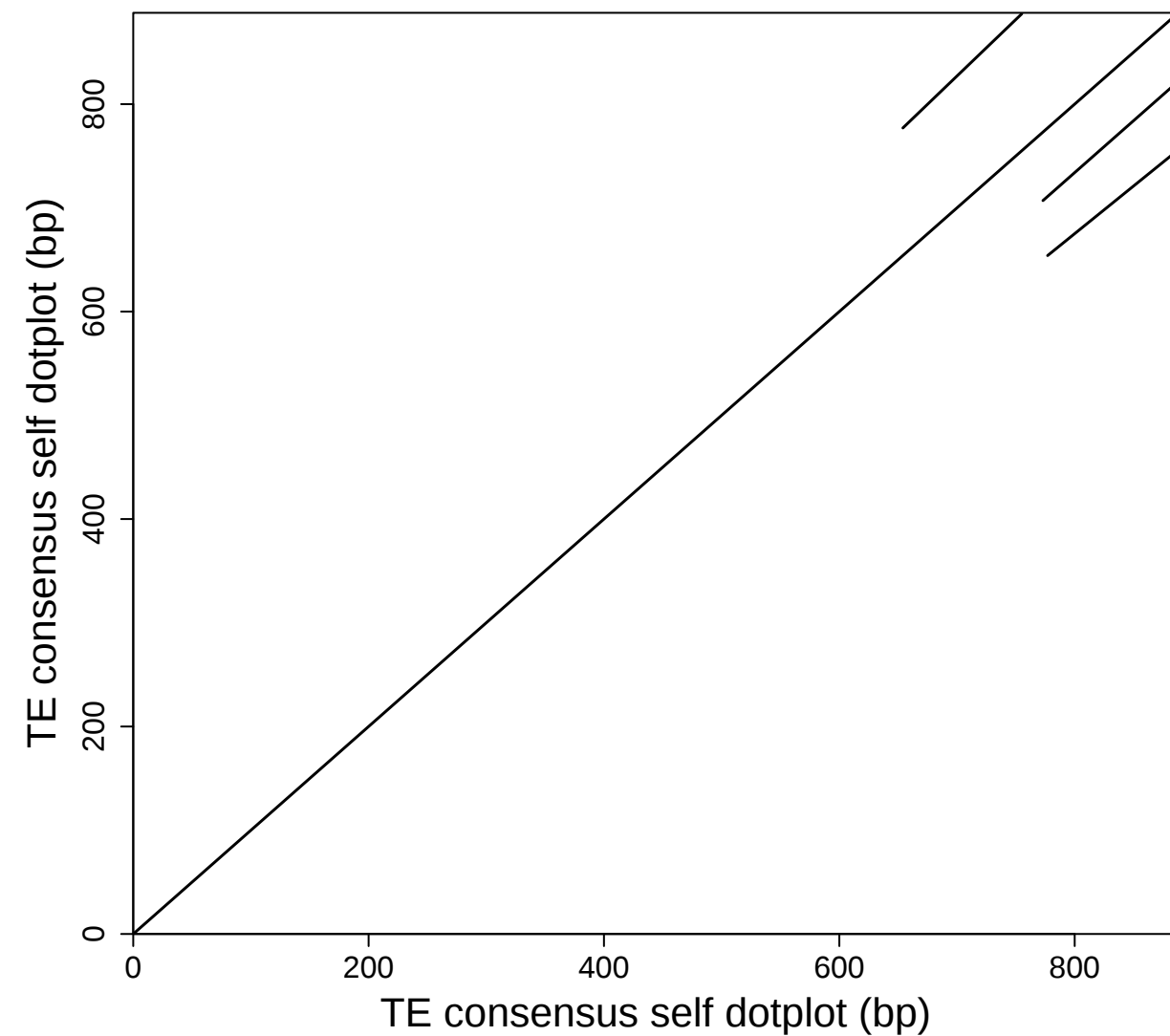
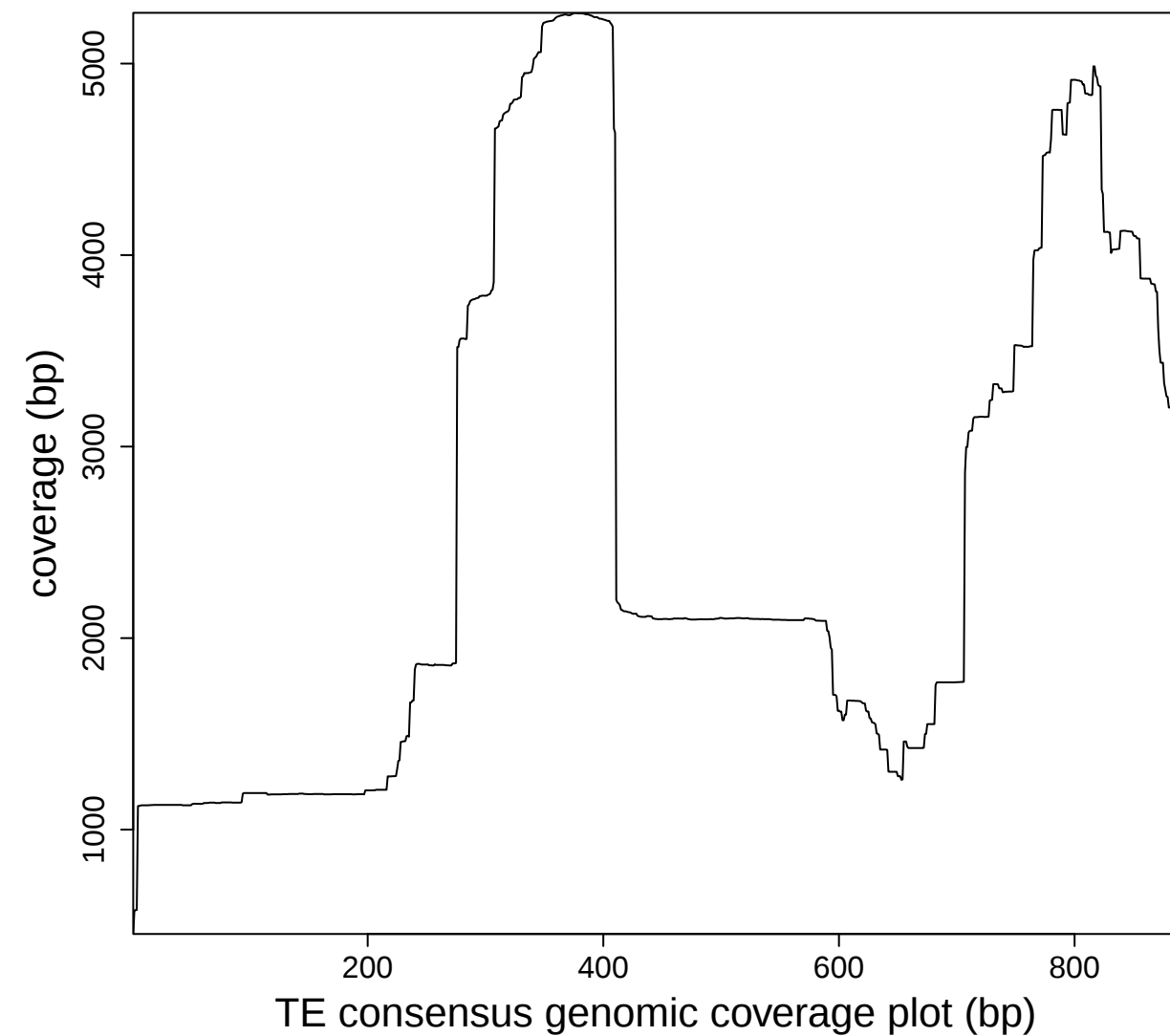
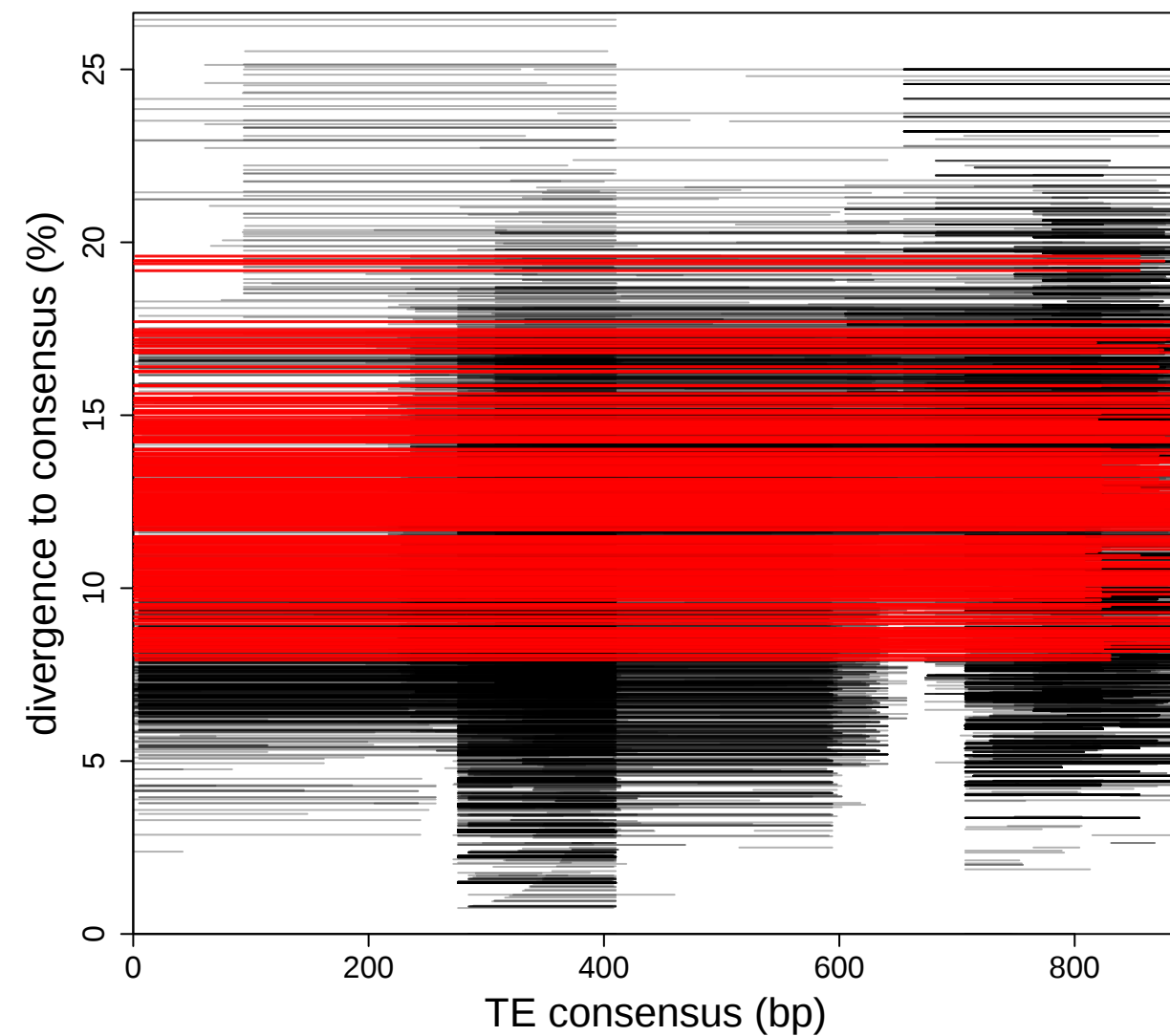


ly_1453.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 6263bp; fragments: 18722; full length: 0 (>=5636.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

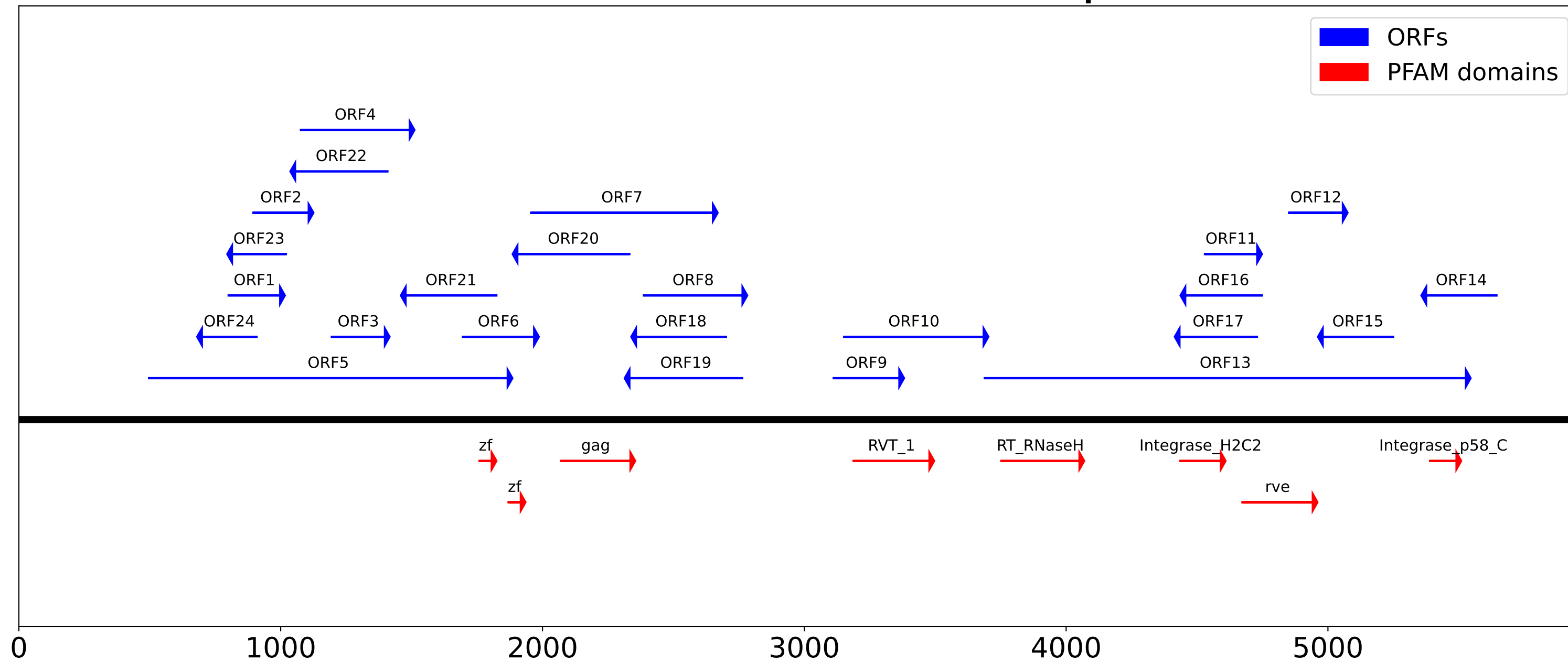


TE: rnd_4_family_1453
size: 888bp; fragments: 9723; full length: 303 (≥ 799.2 bp)

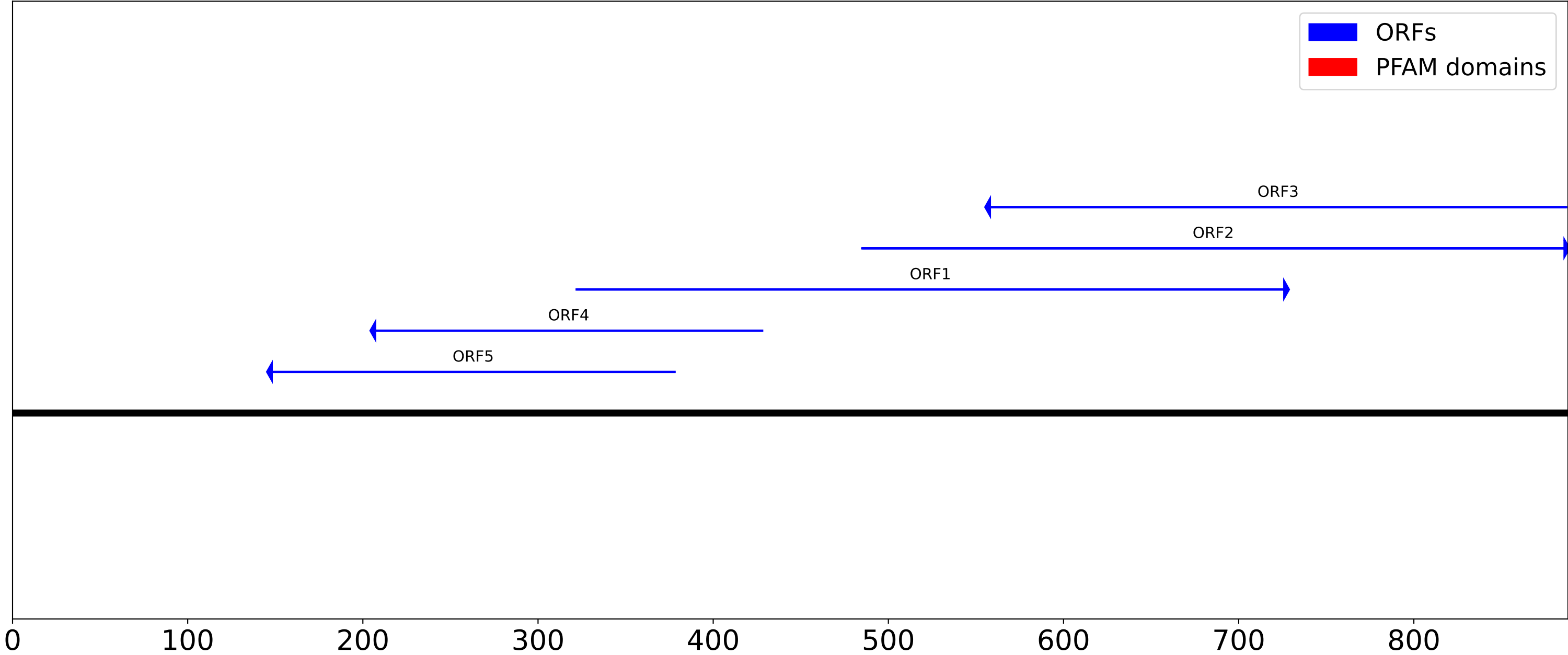


Before TEtrimmer 888 bp

After TETrimmer ORF and PFAM domain plot



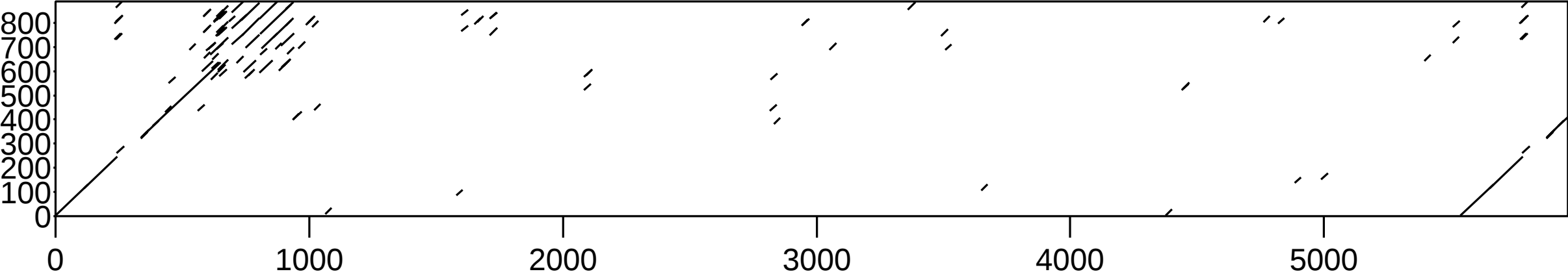
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

Isensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)