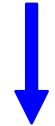


Start crop Point



End crop Point



MSA length = 2111



Start crop Point

End crop Point

0

10

0

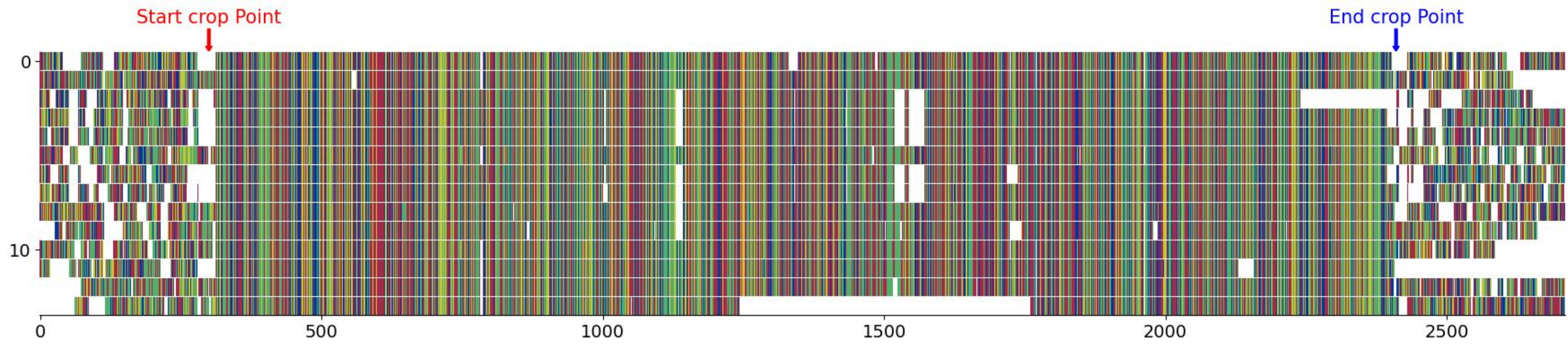
500

1000

1500

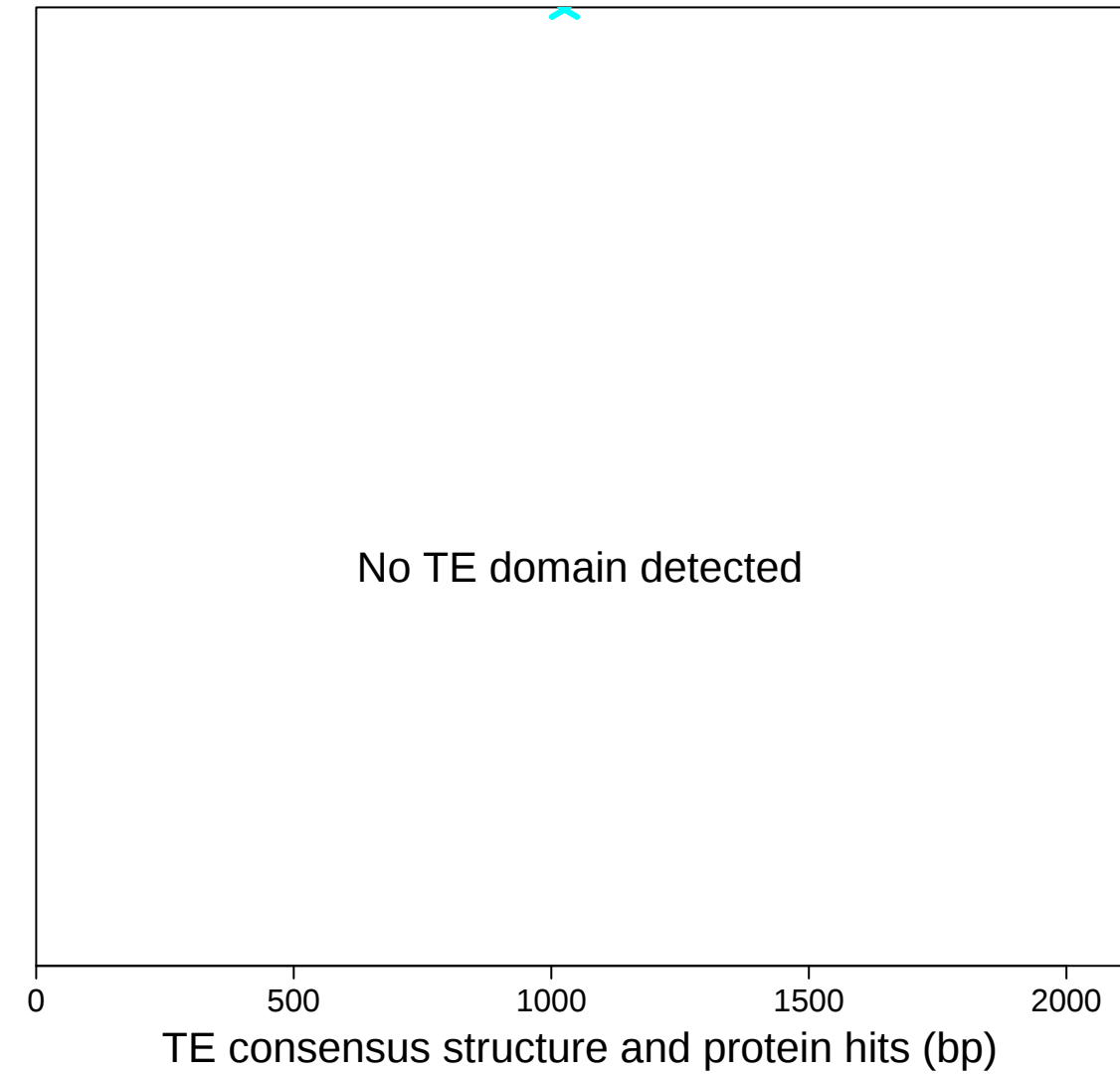
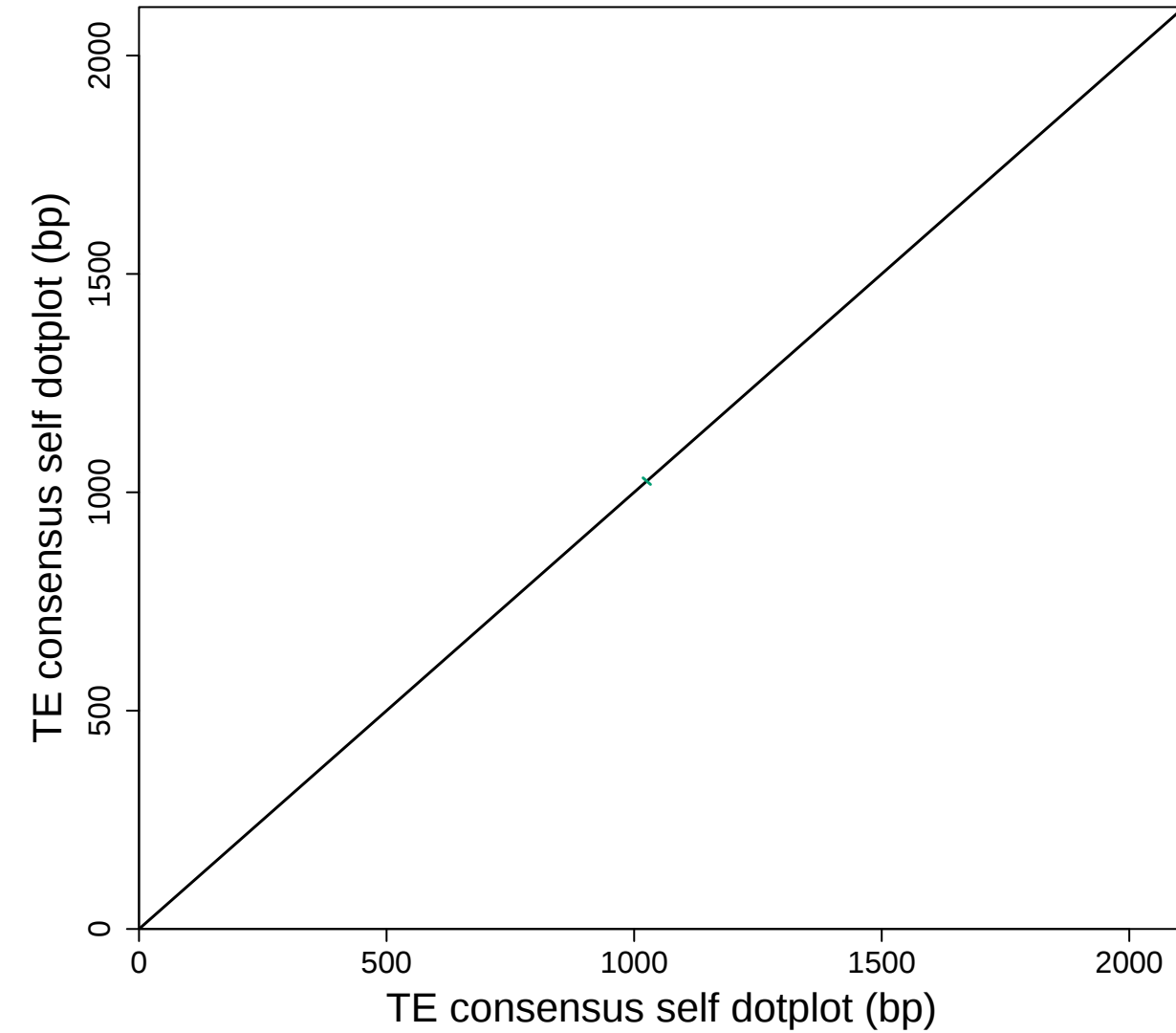
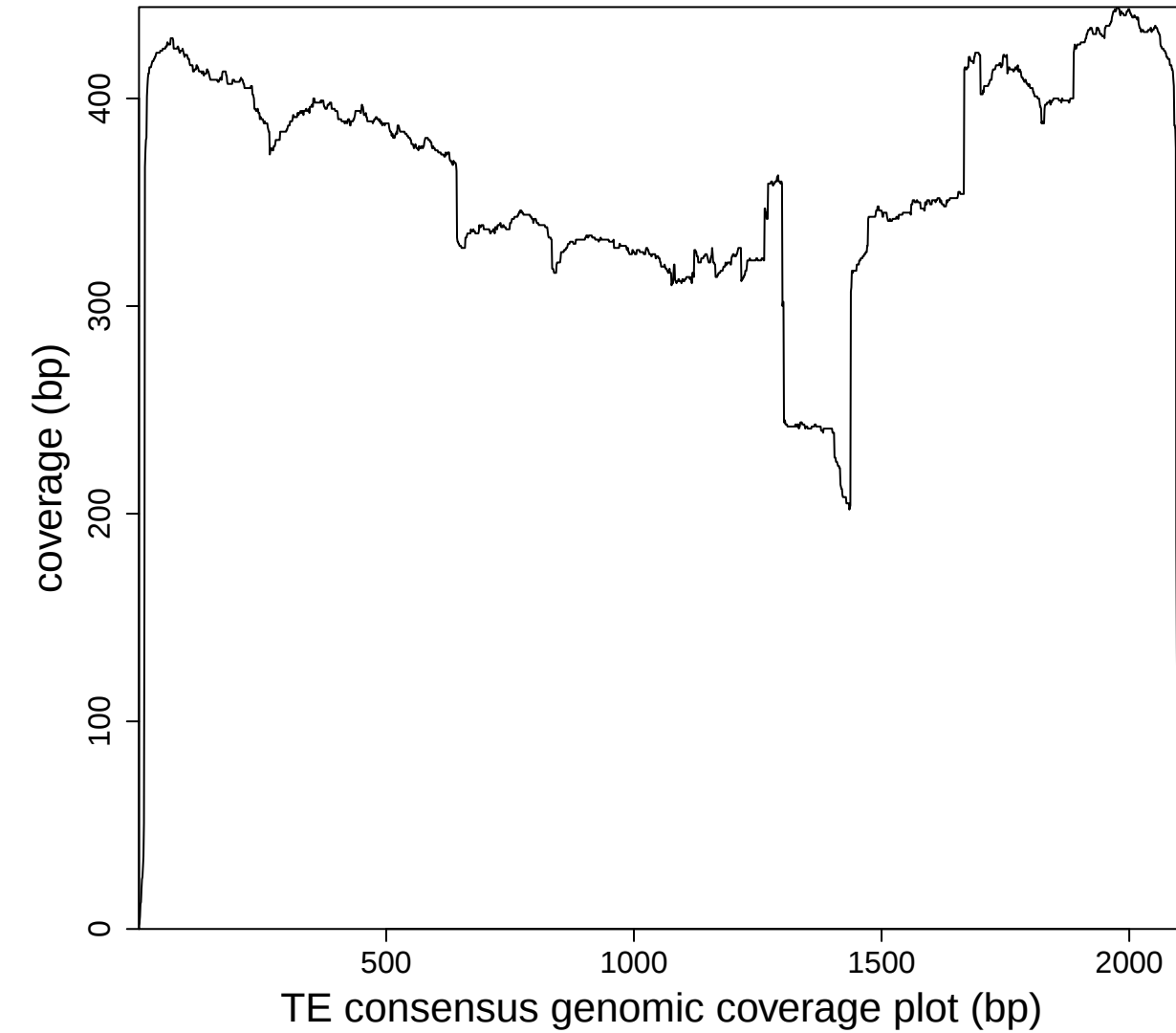
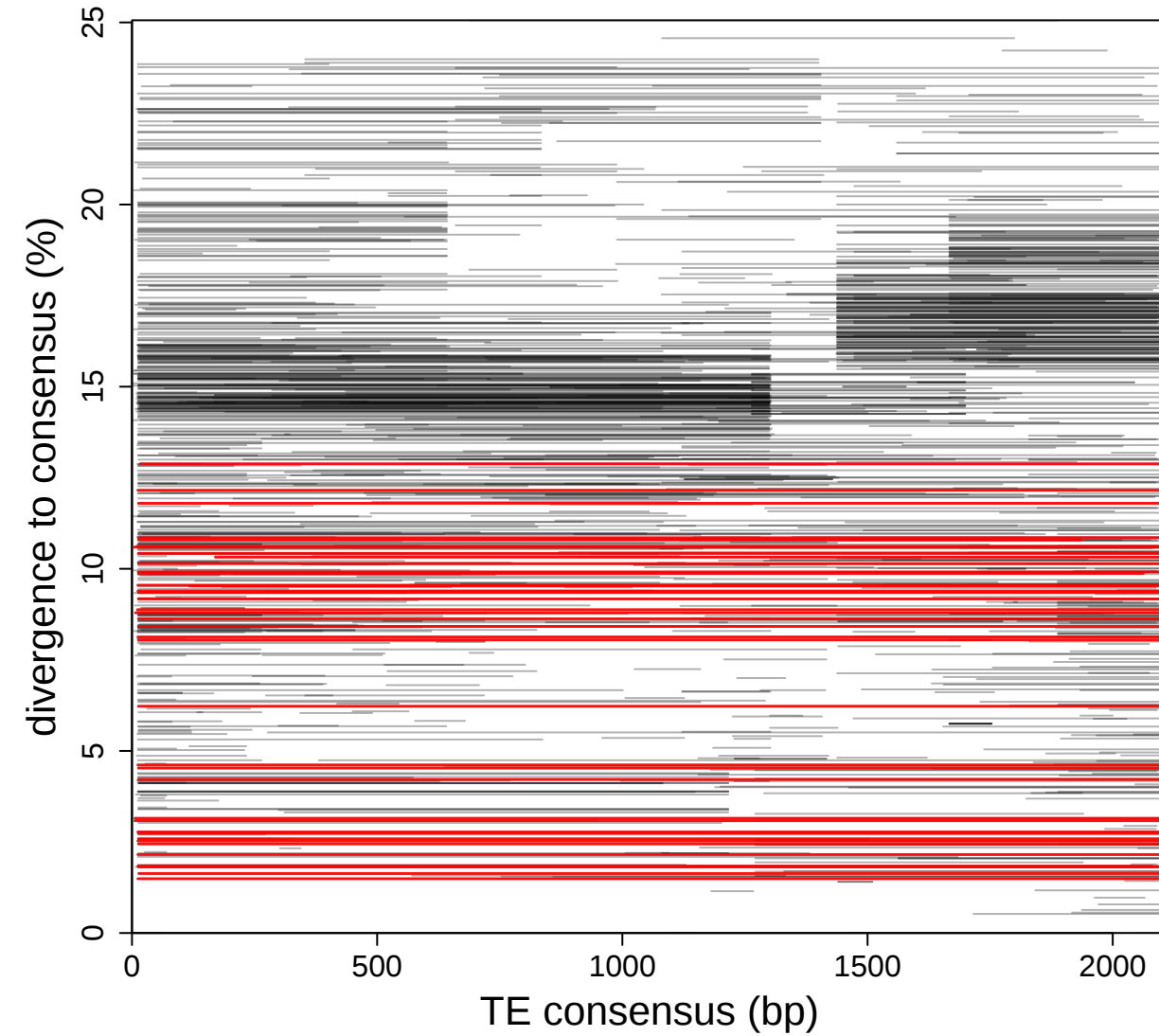
2000

2500



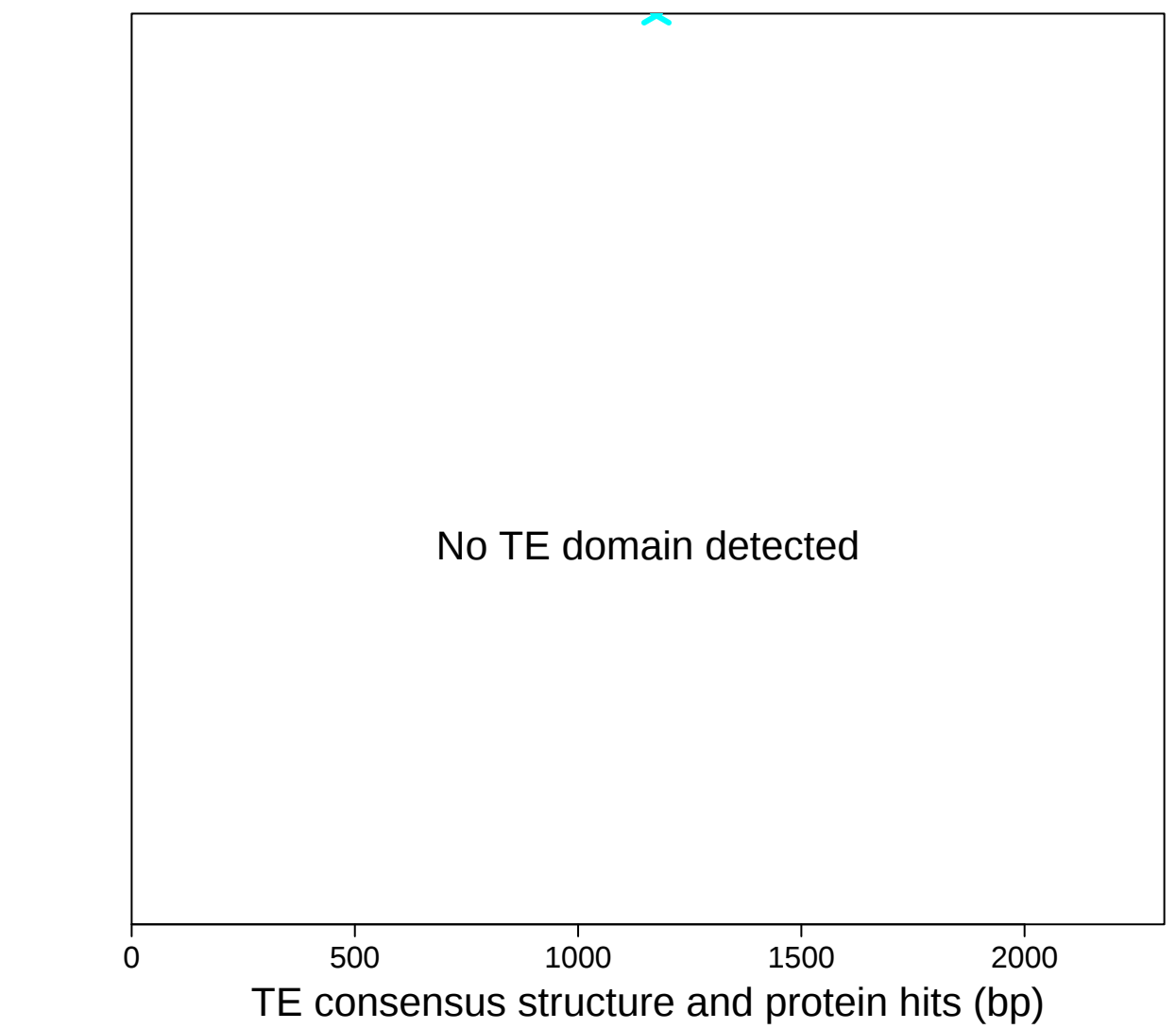
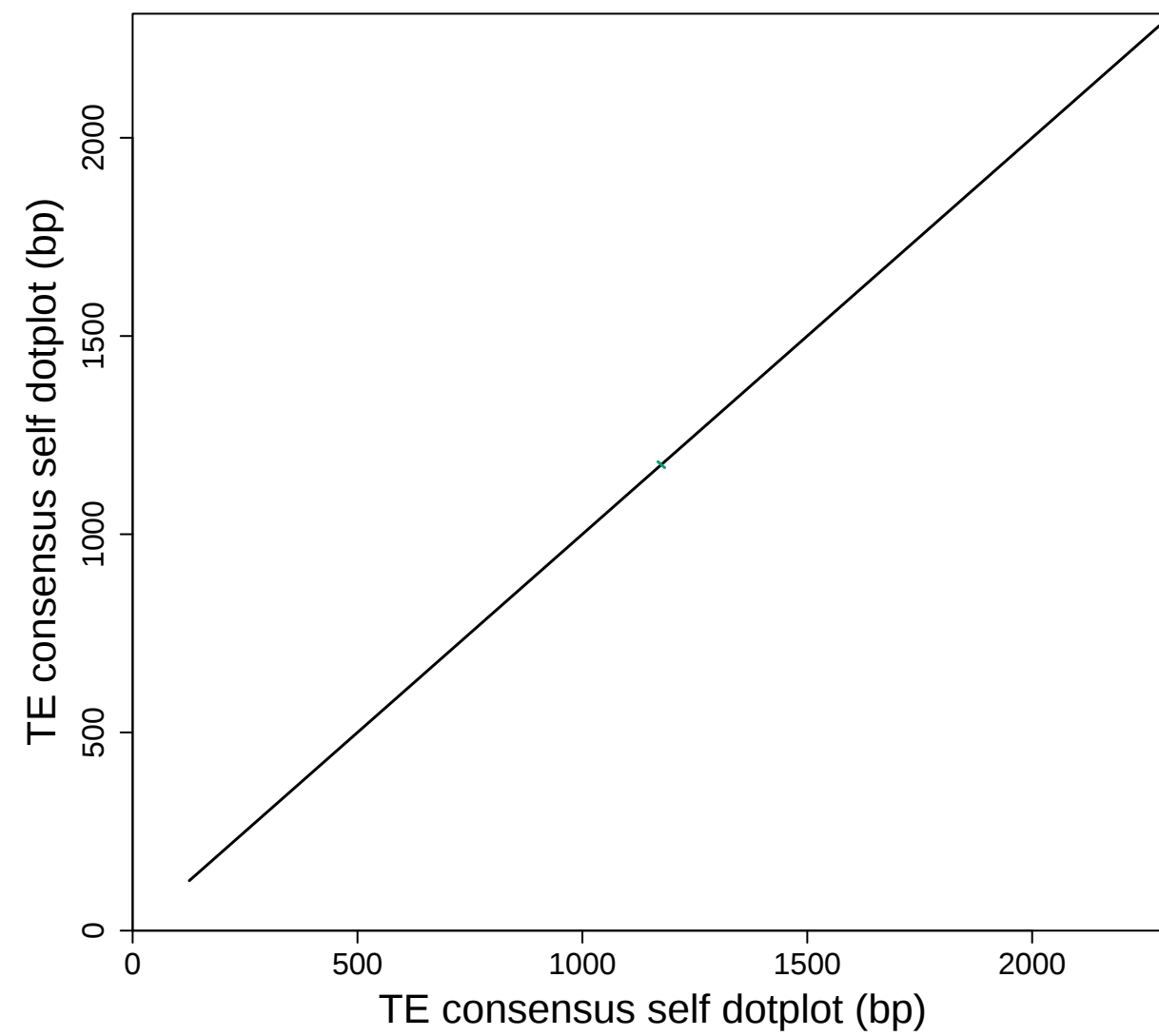
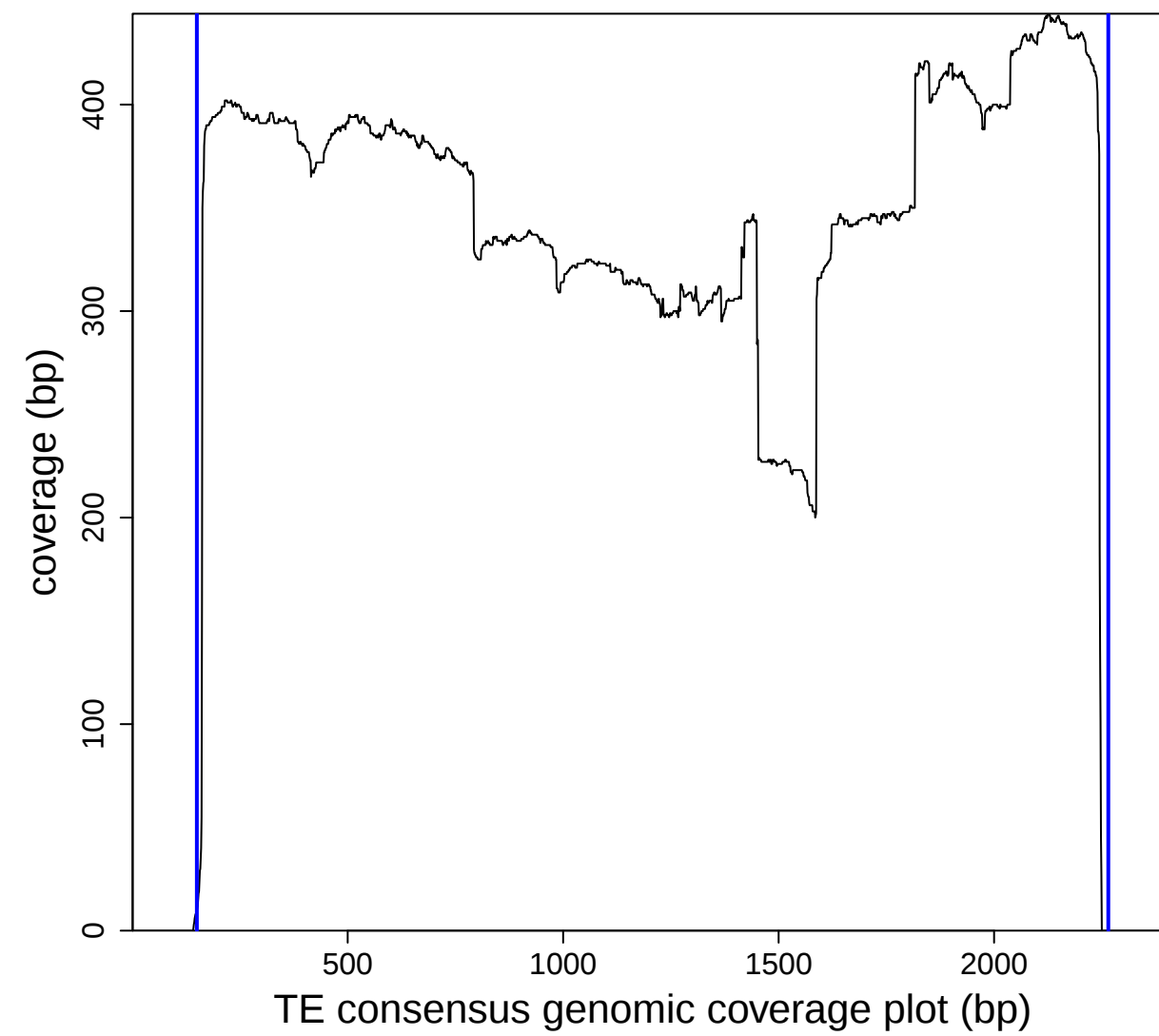
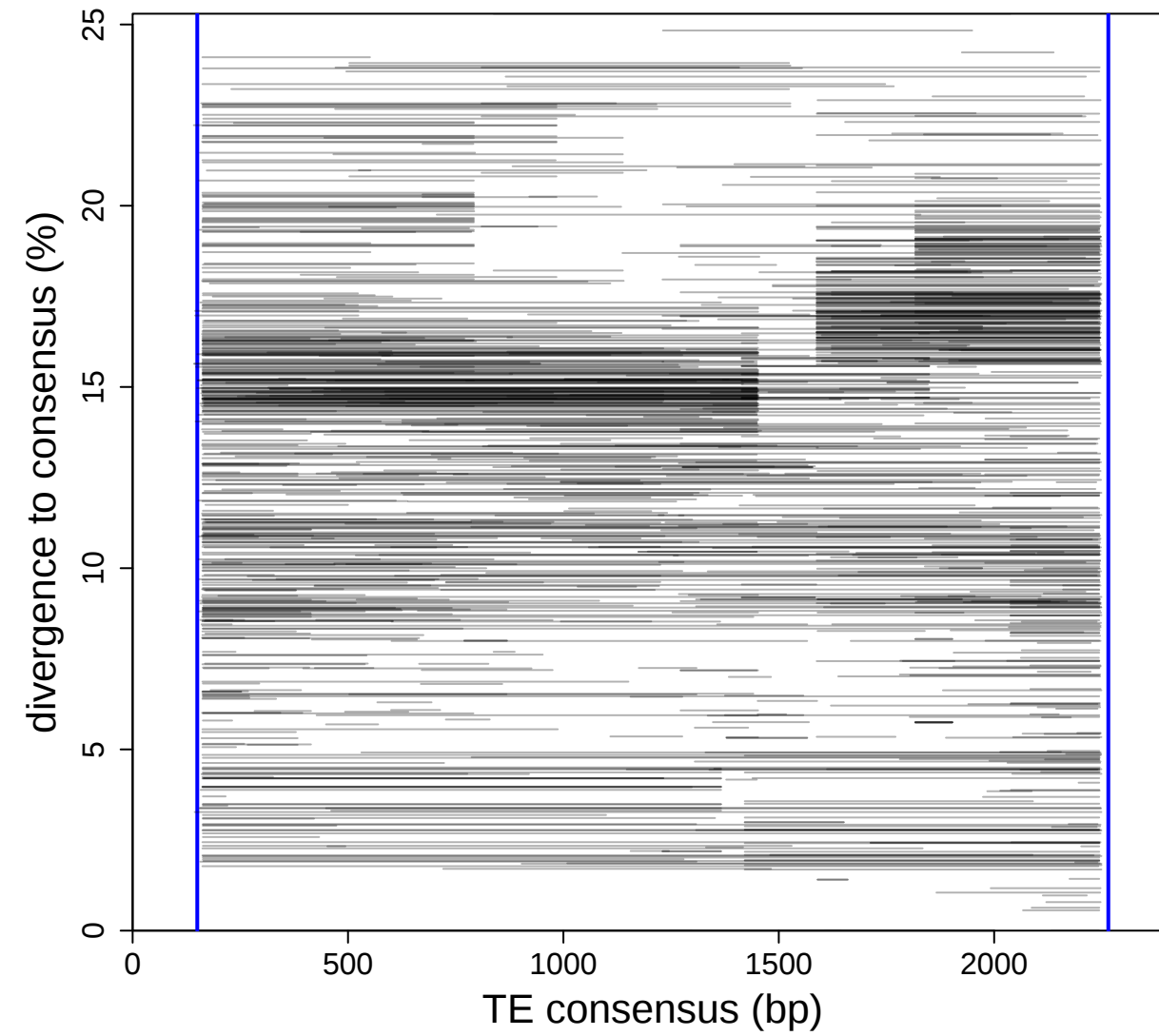
b.bed uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa
size: 2111bp; fragments: 1190; full length: 40 (≥ 1899.9 bp)

After TETrimmer 2111 bp

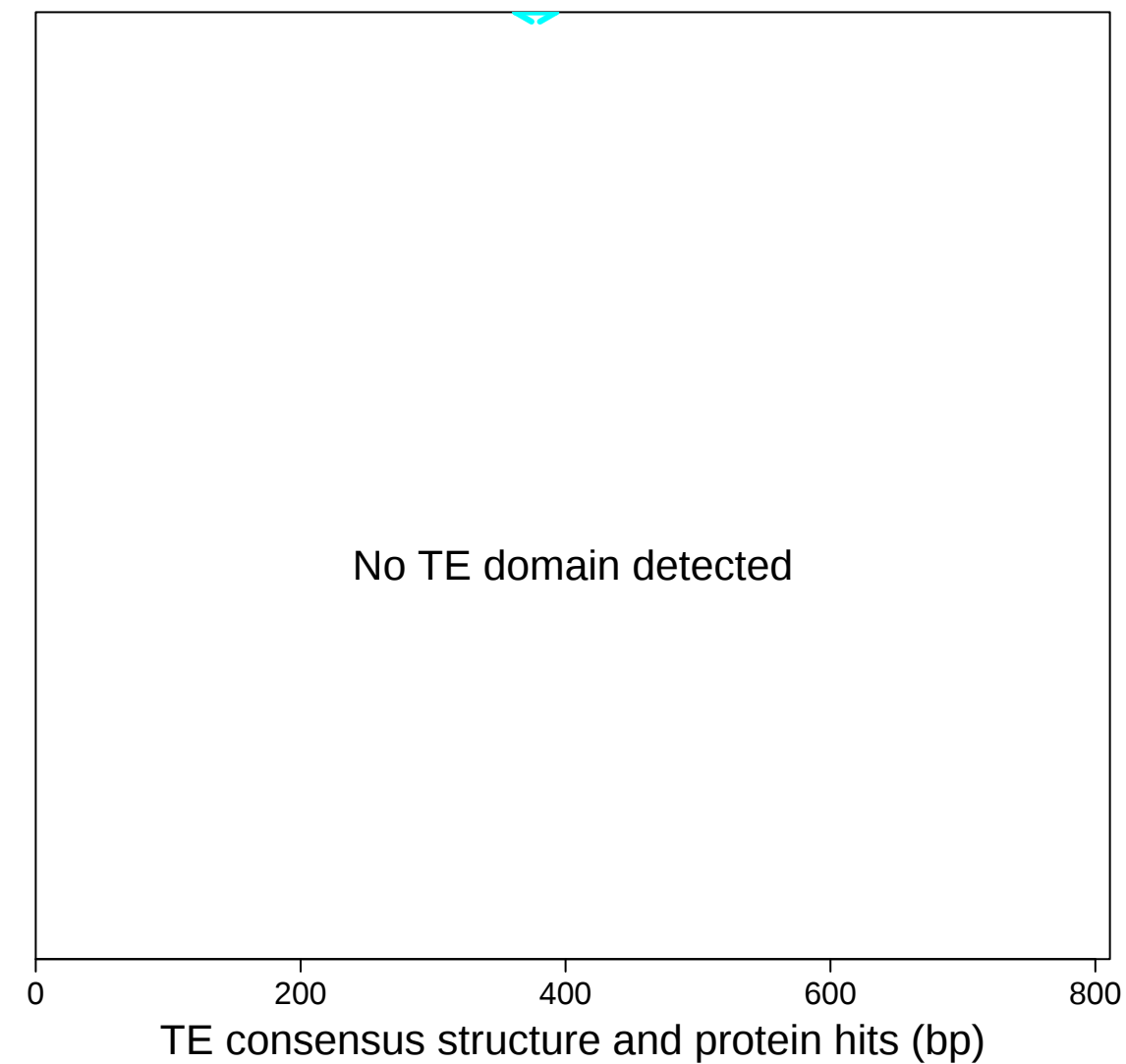
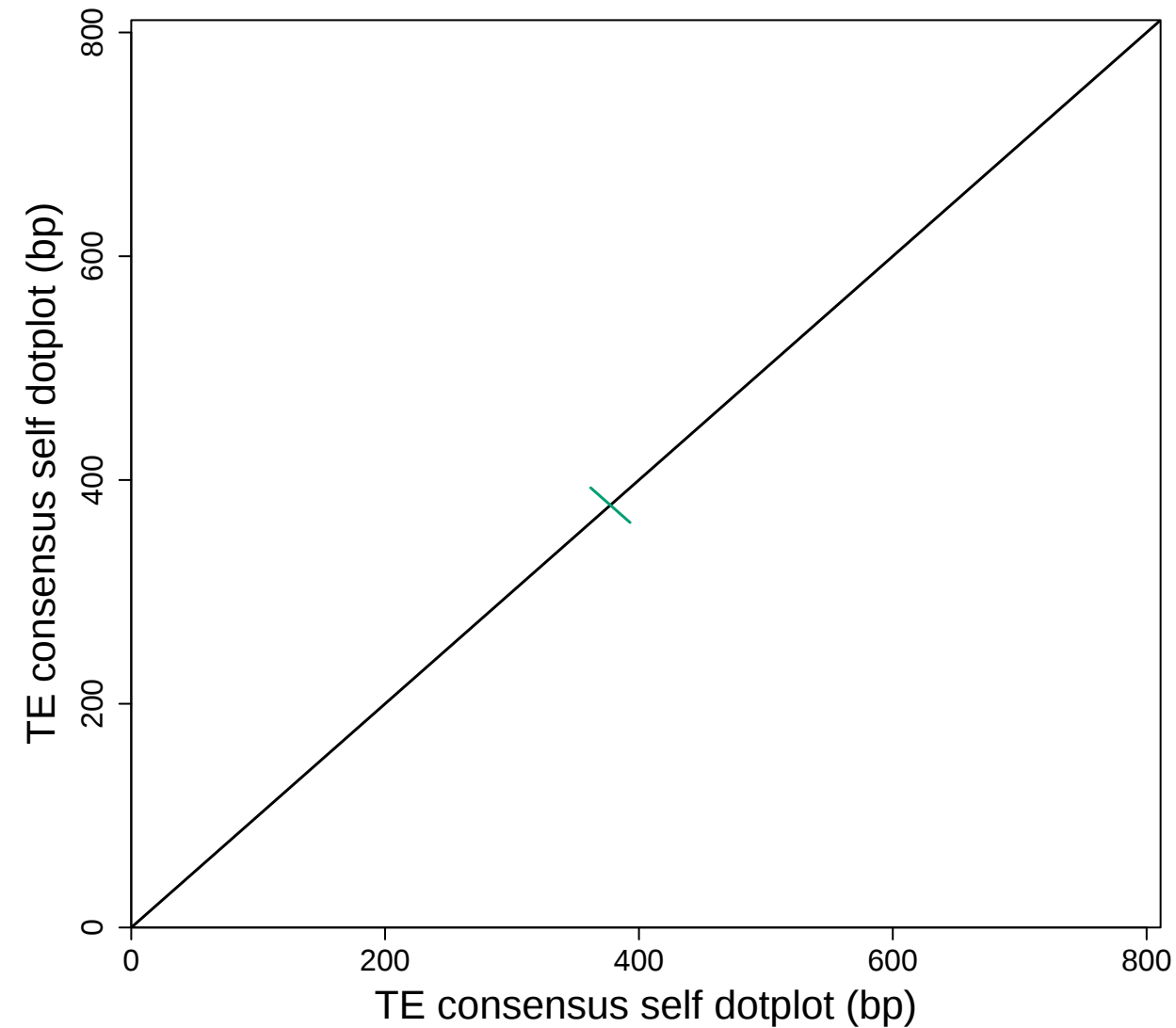
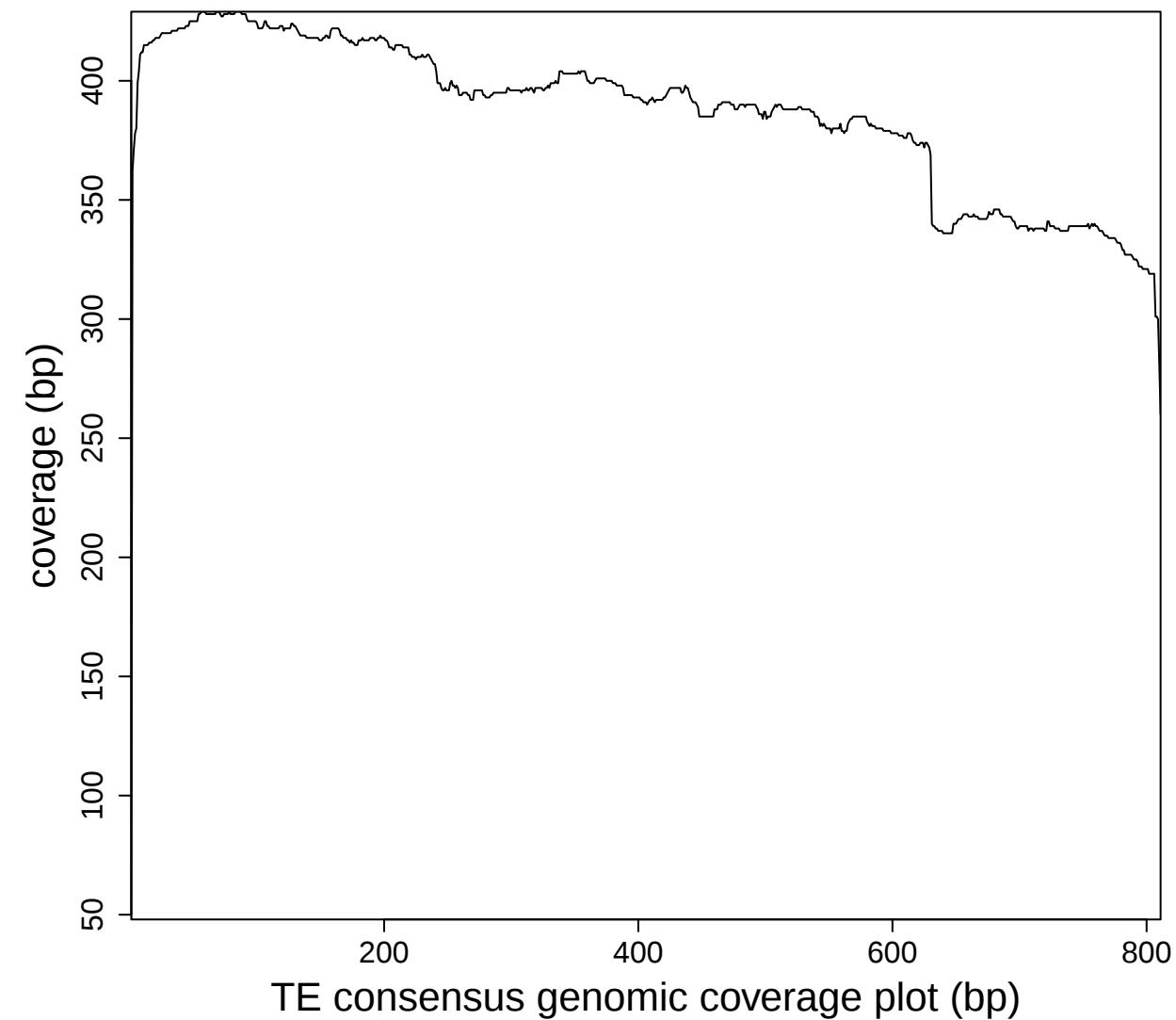
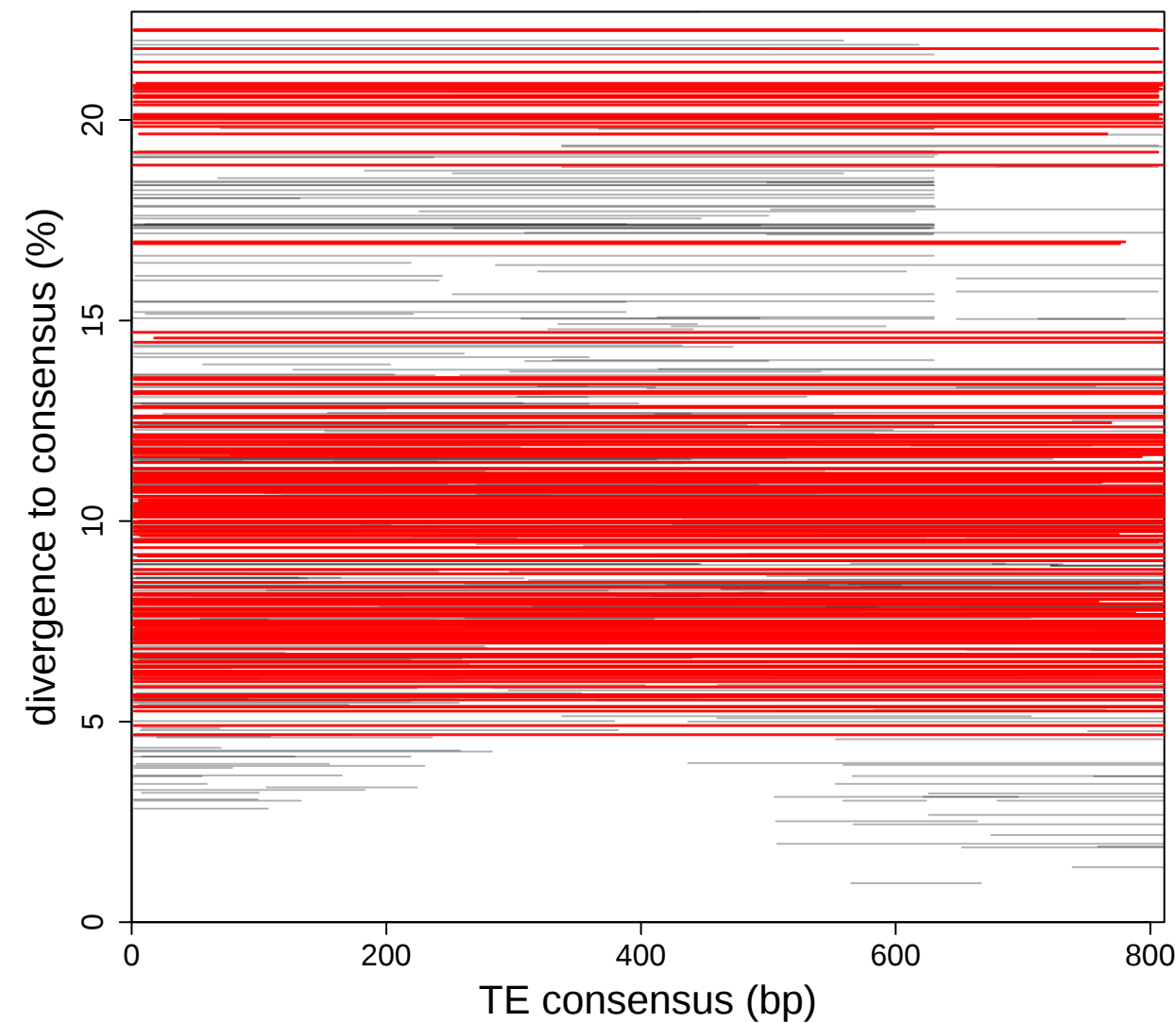


.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 2415bp; fragments: 1148; full length: 0 (>=2173.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

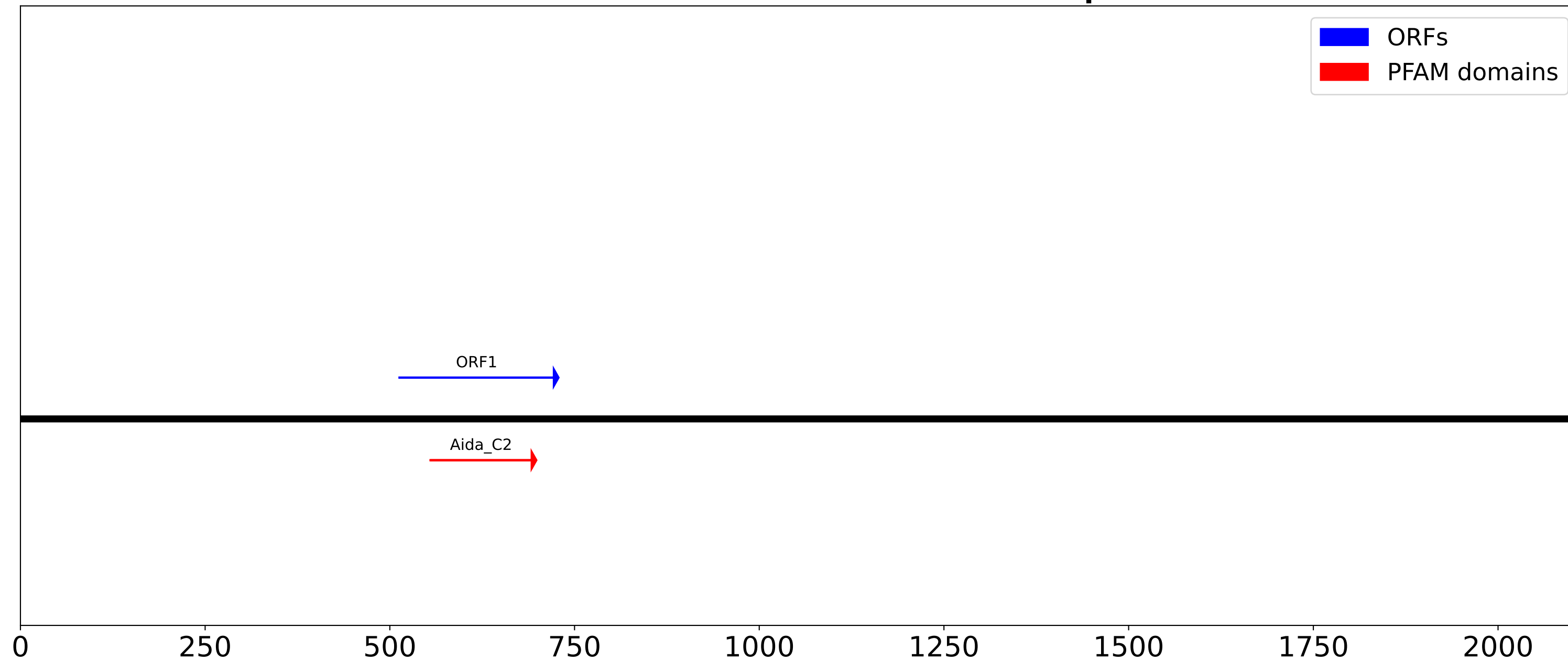


TE: rnd_1_family_71
size: 811bp; fragments: 602; full length: 228 (≥ 729.9 bp)



Before TEtrimmer 811 bp

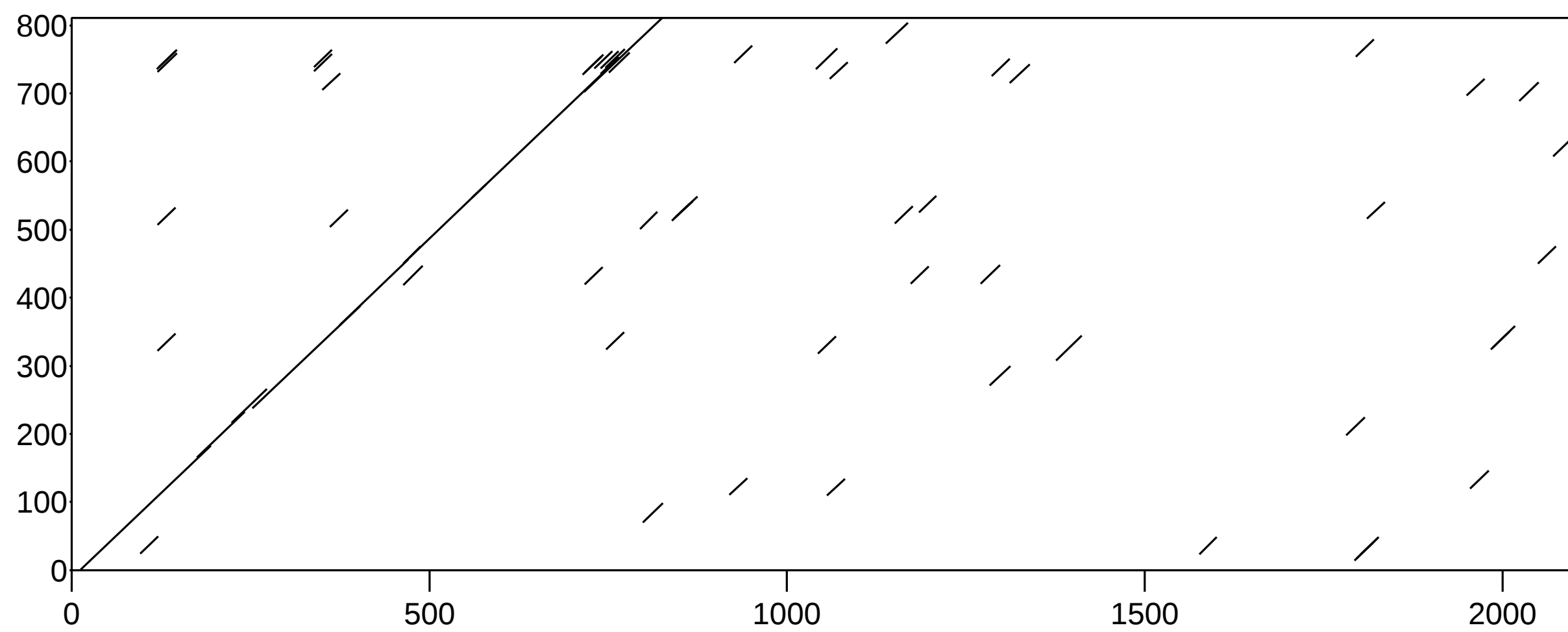
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)