

Start crop Point

End crop Point

MSA length = 789



Start crop Point

End crop Point

0

10

0

200

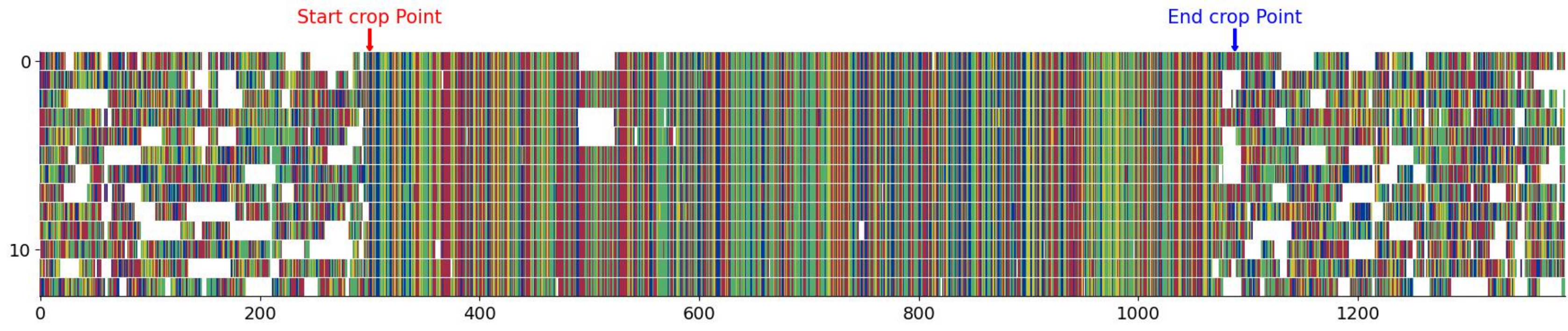
400

600

800

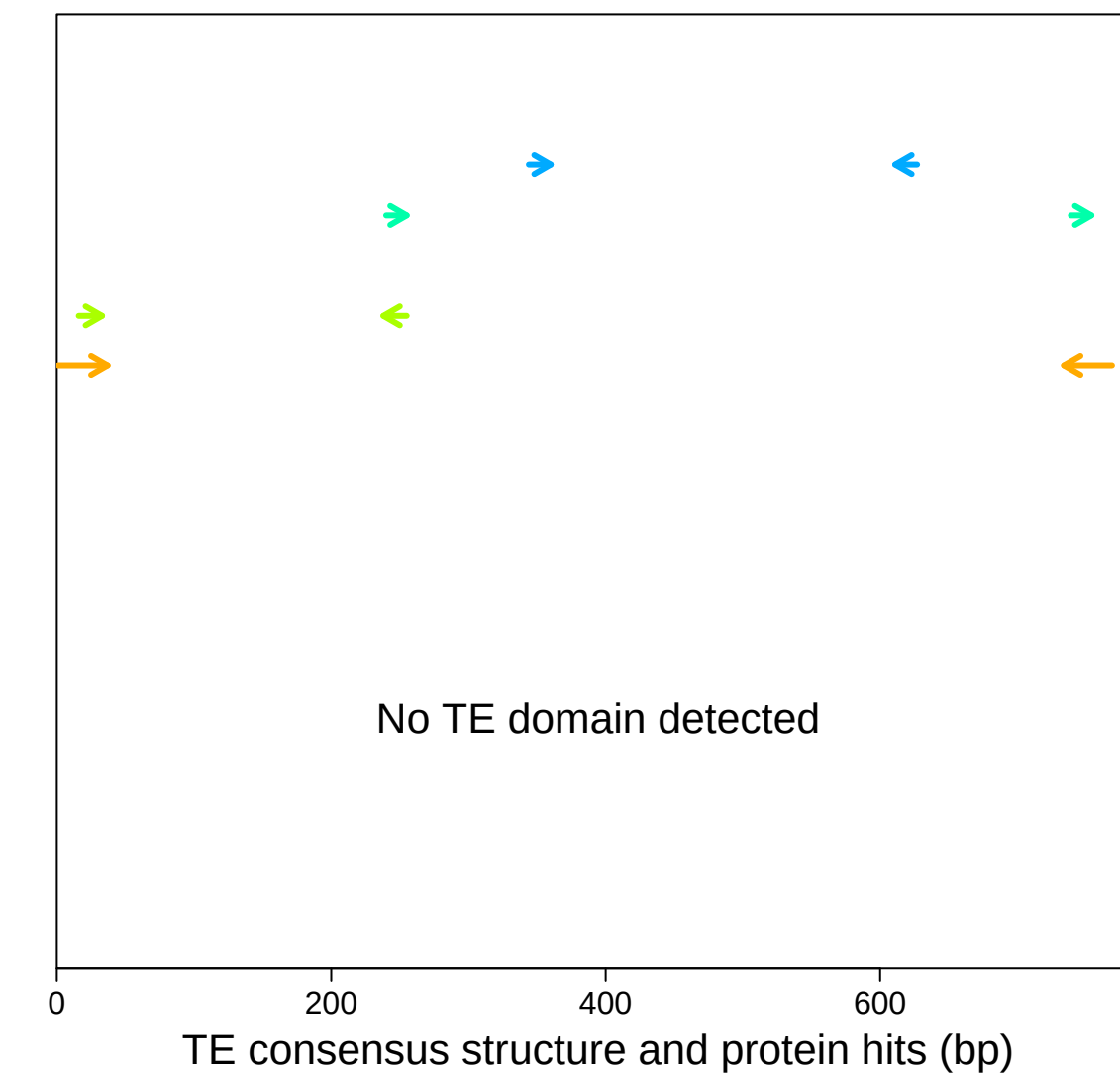
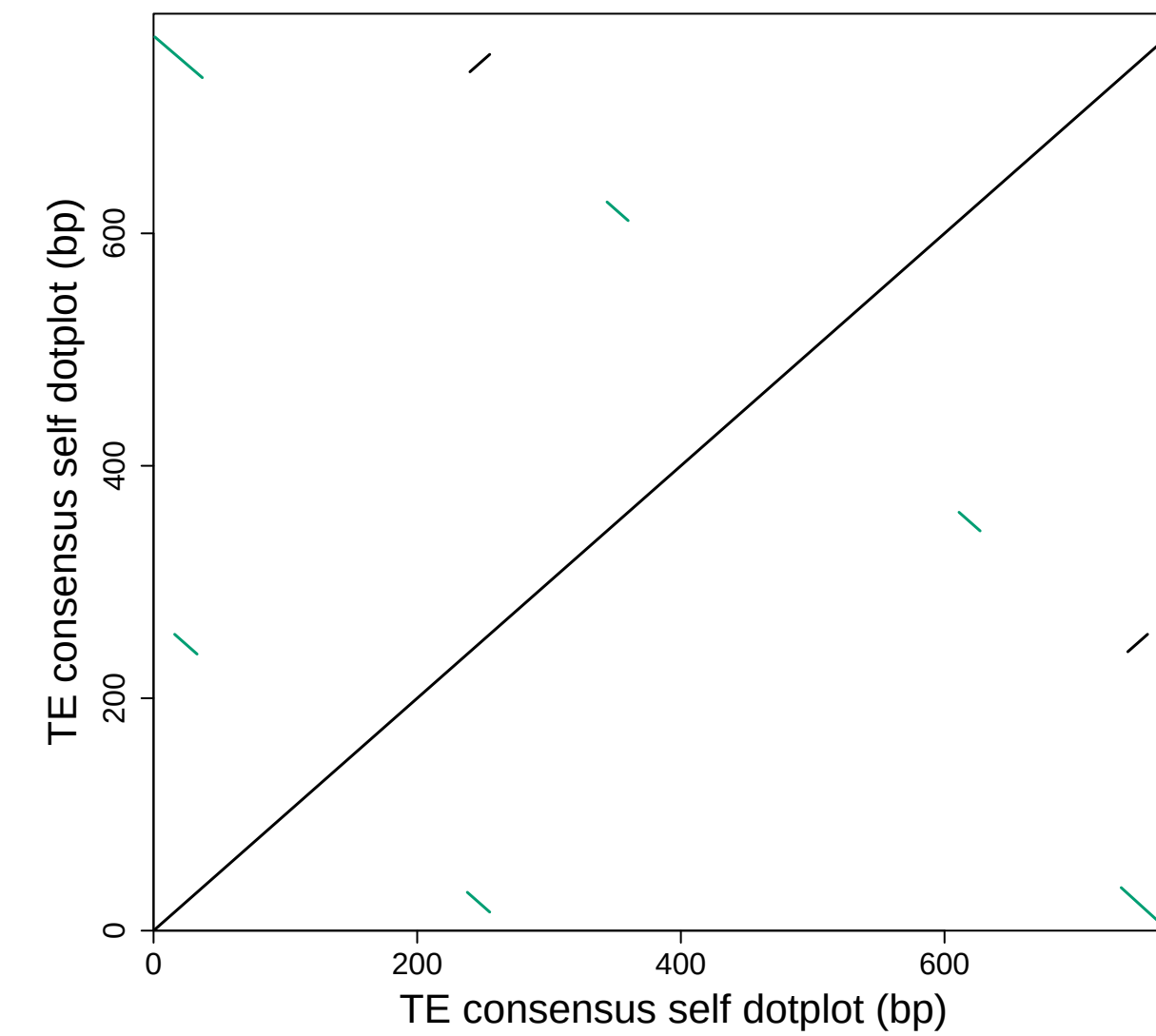
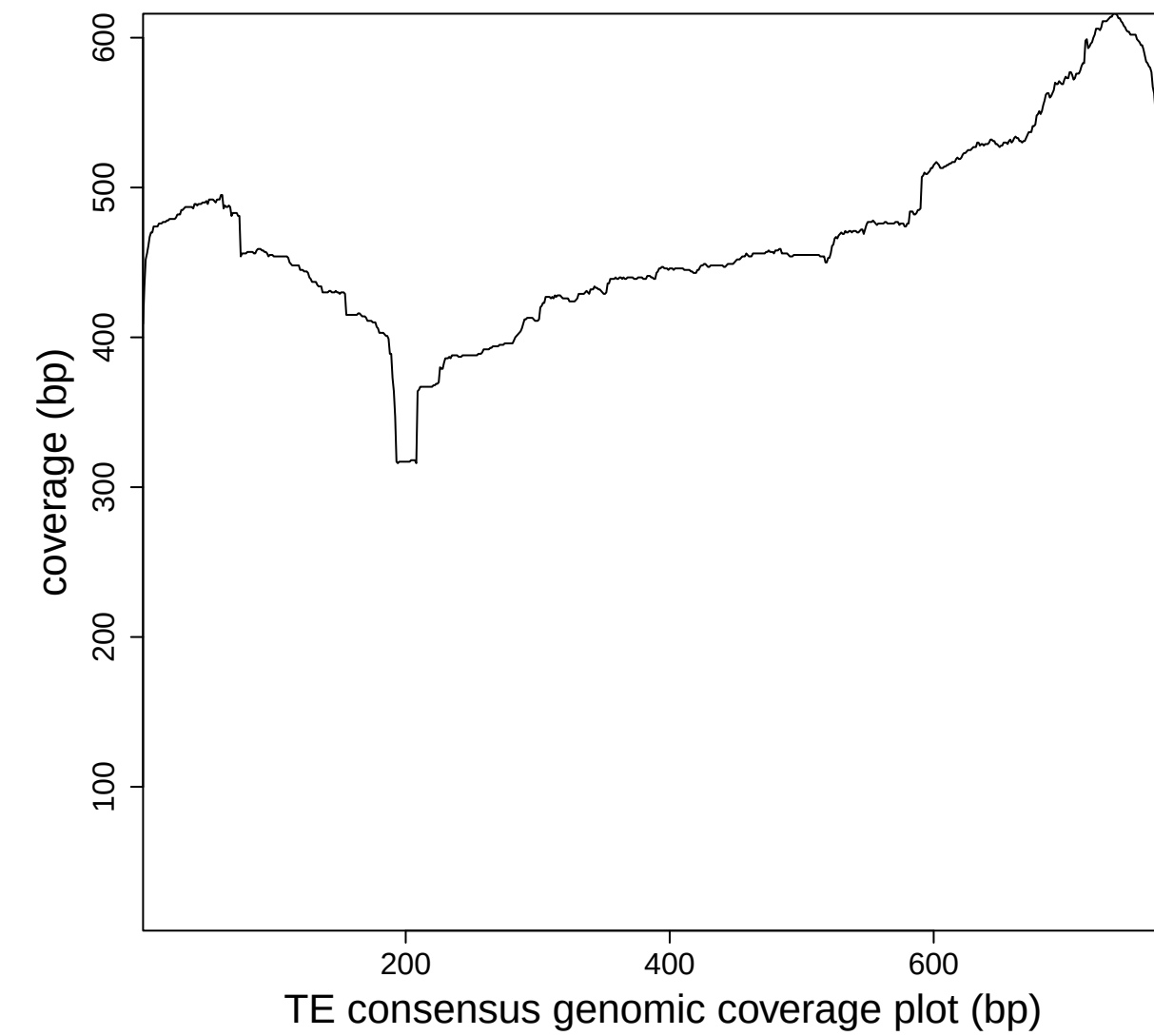
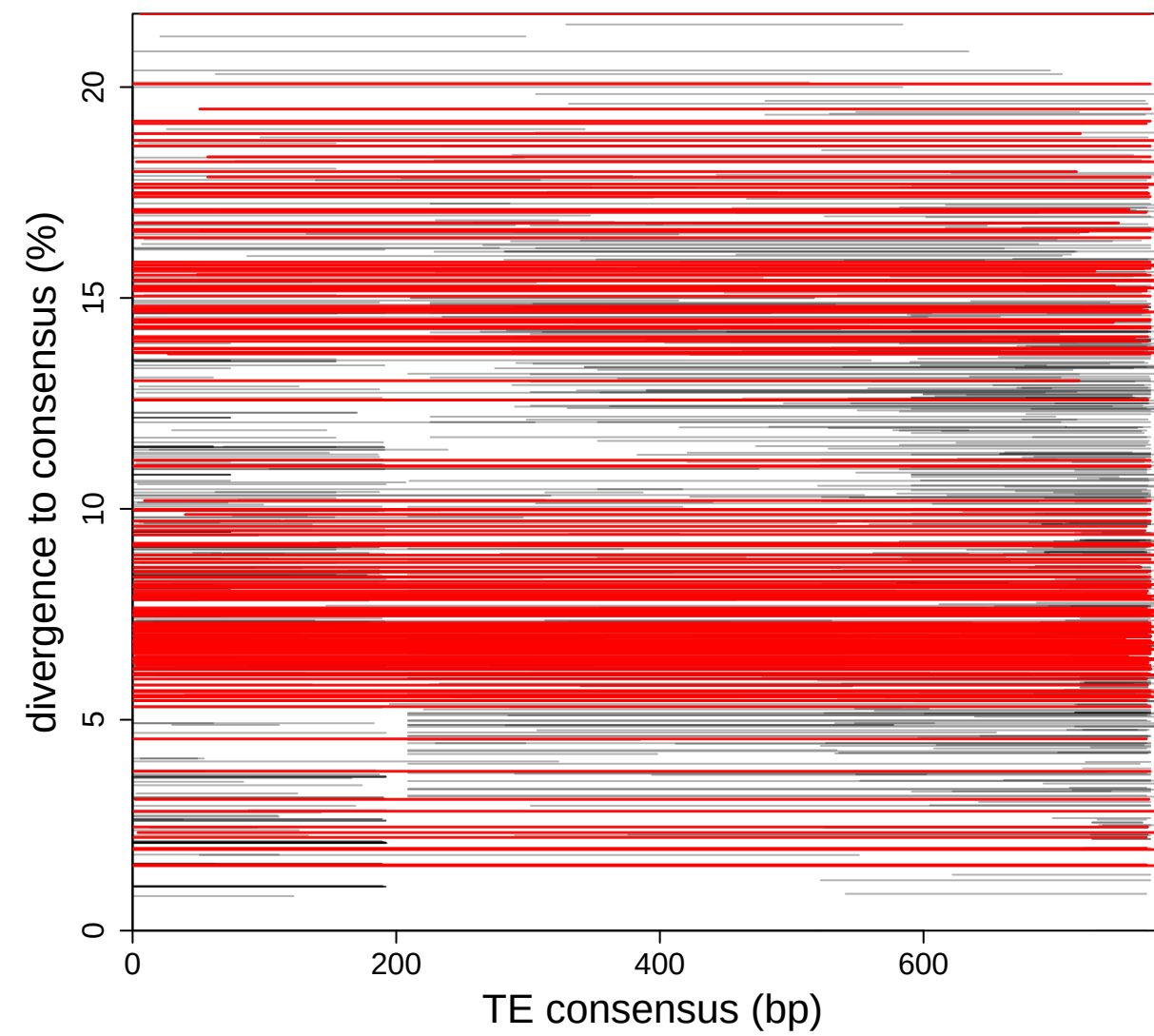
1000

1200



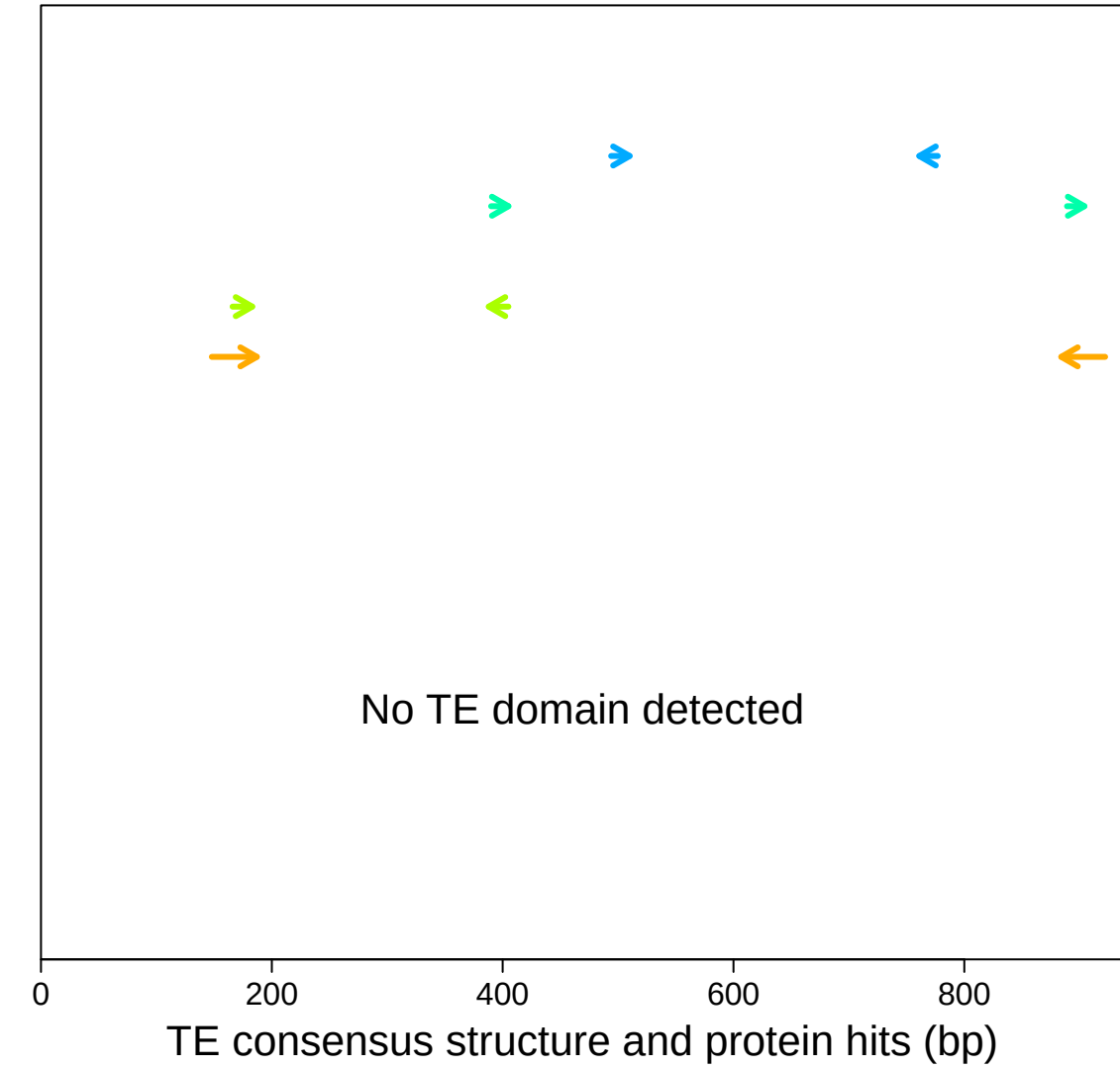
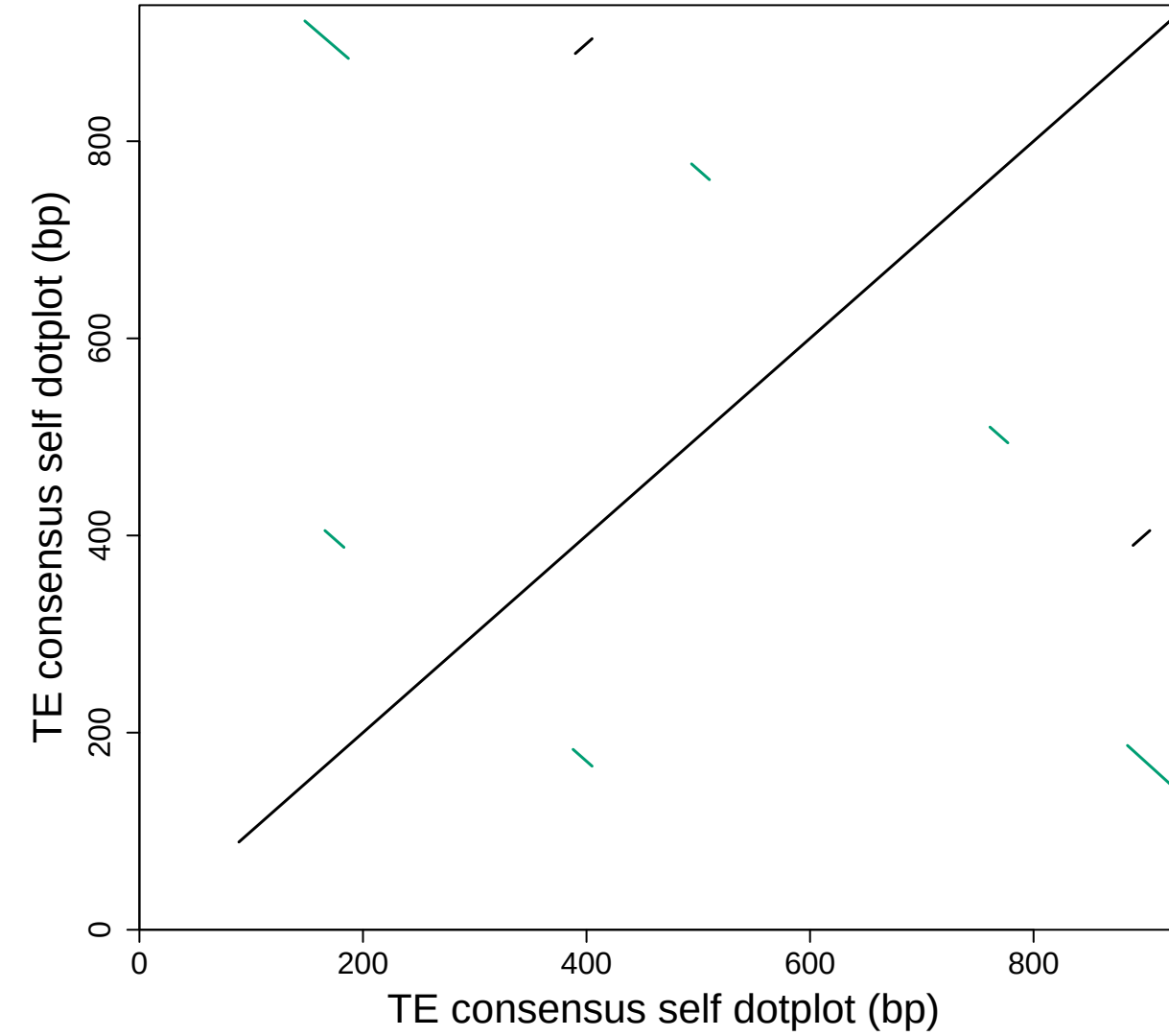
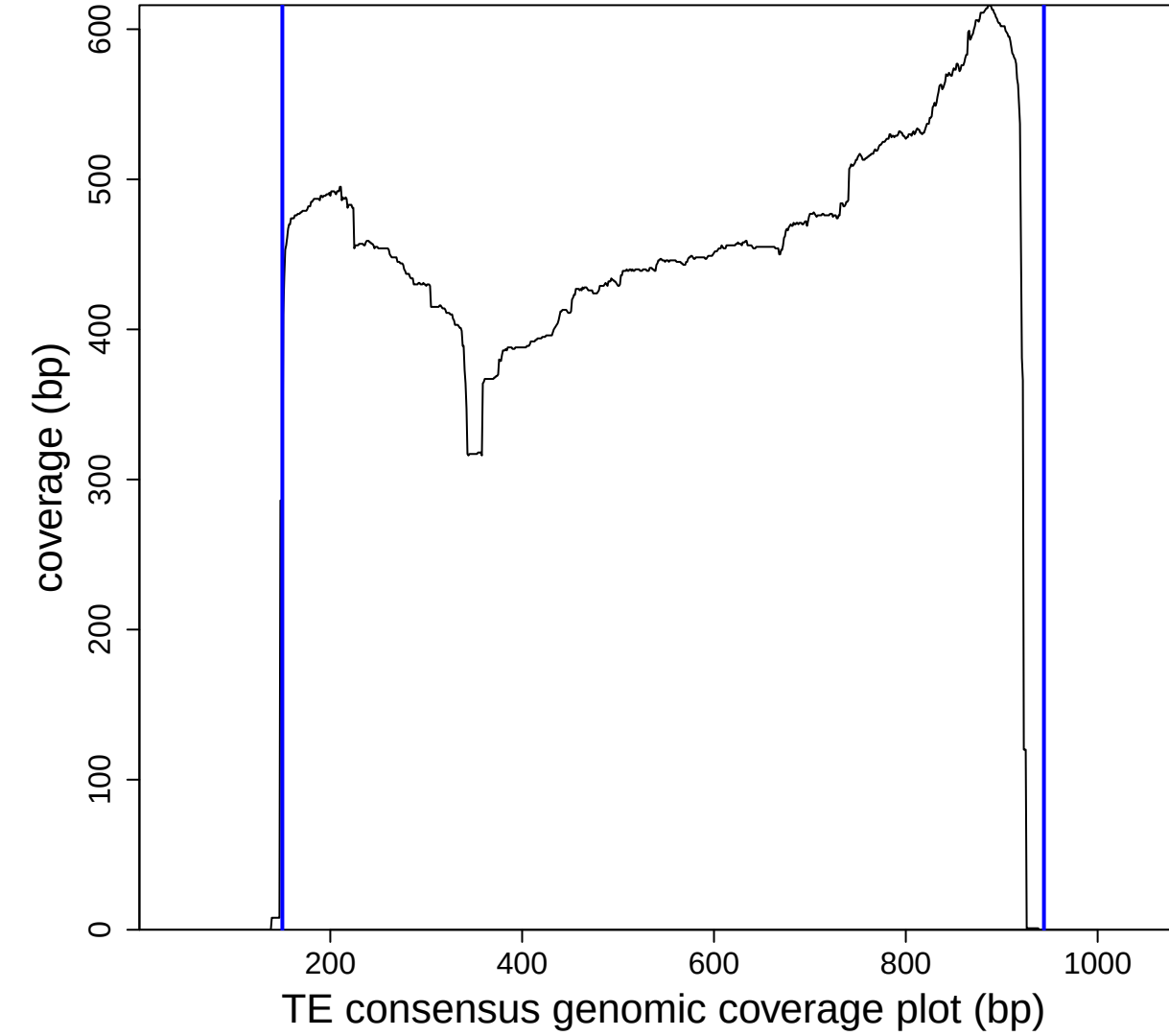
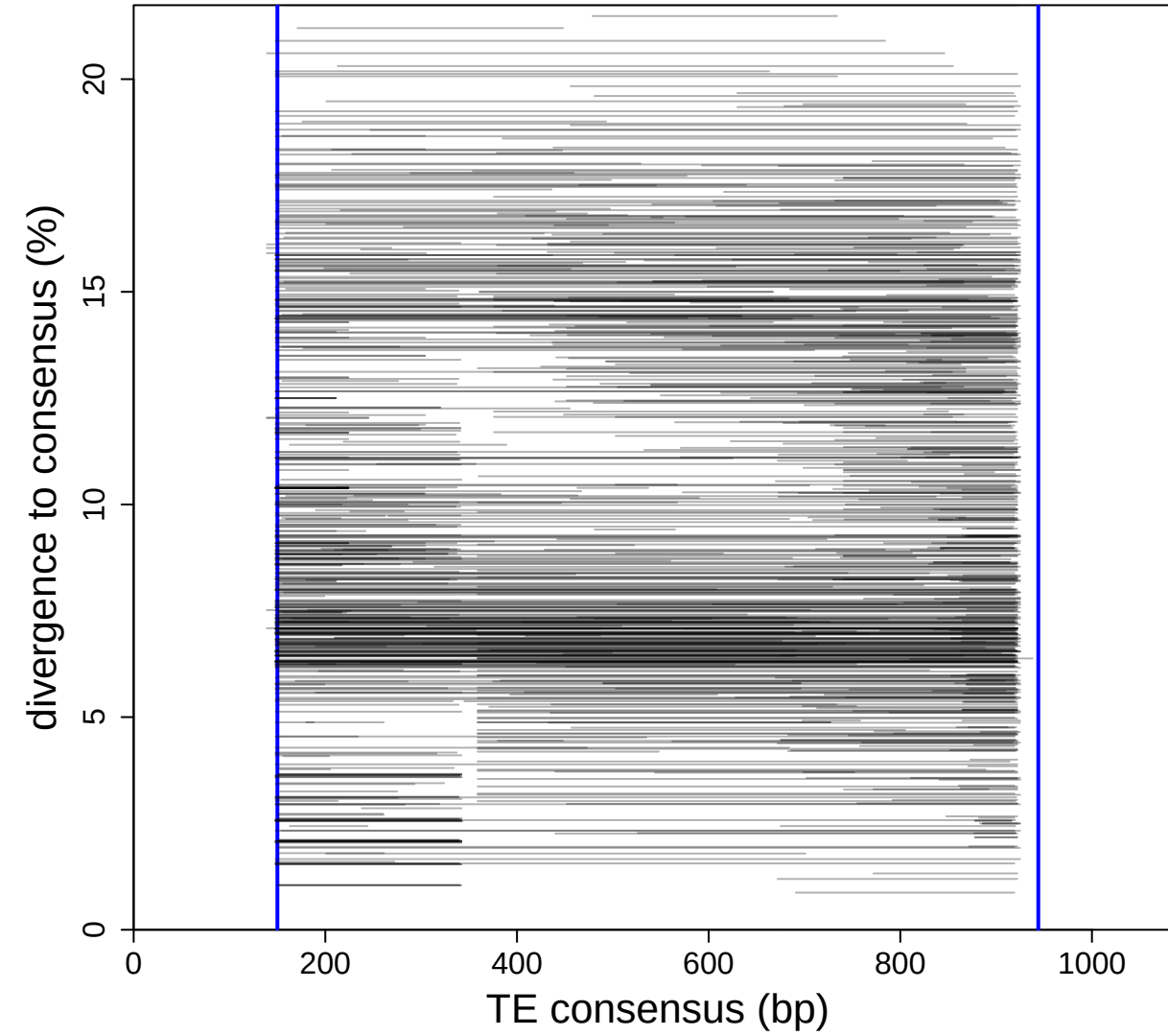
l.b.bed uf.bed g_4.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 789bp; fragments: 993; full length: 215 (≥ 710.1 bp)

After TEtrimmer 789 bp



7.fasta.b.bed uf.bed g 4.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 1094bp; fragments: 993; full length: 0 (>=984.6bp)

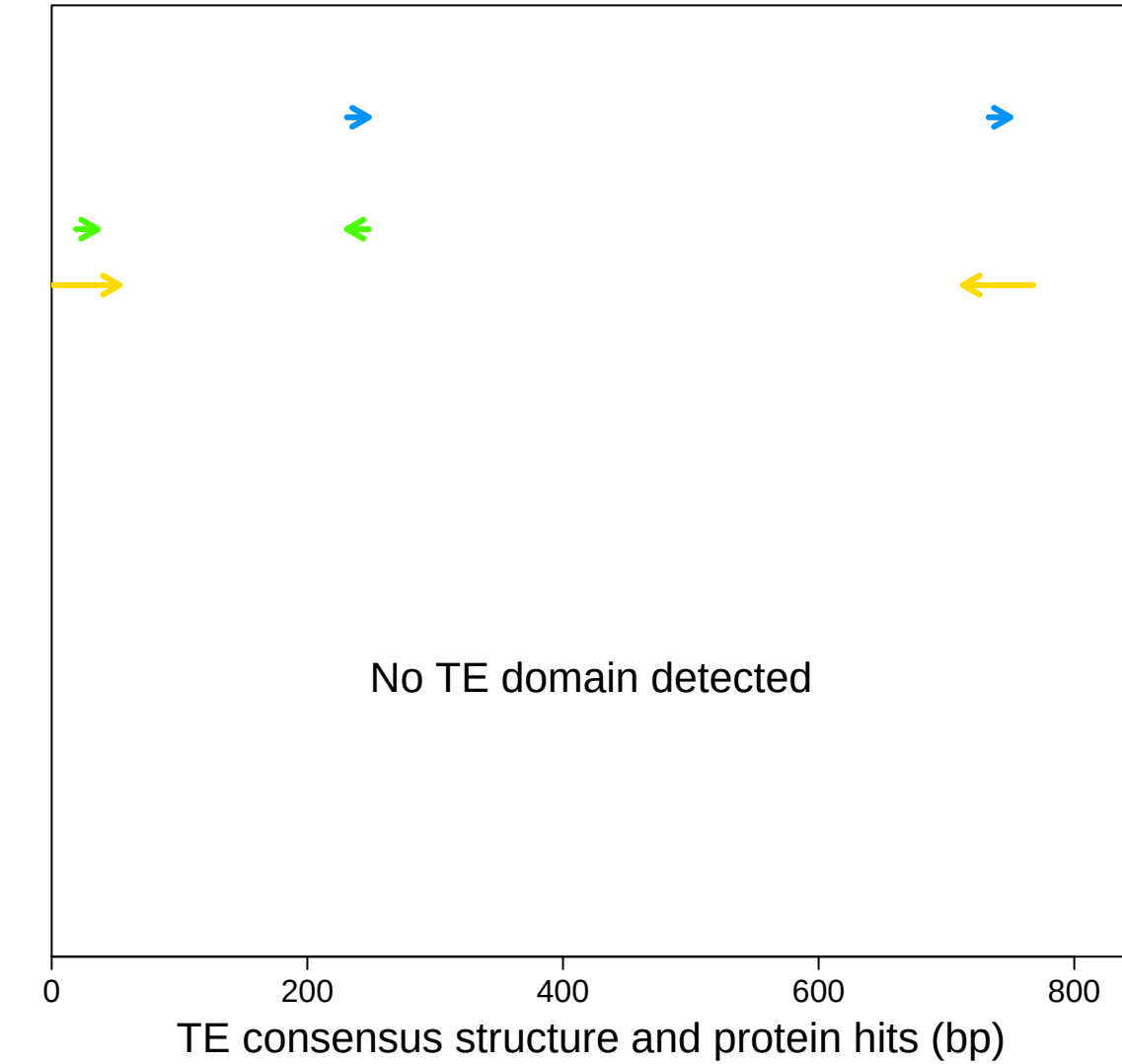
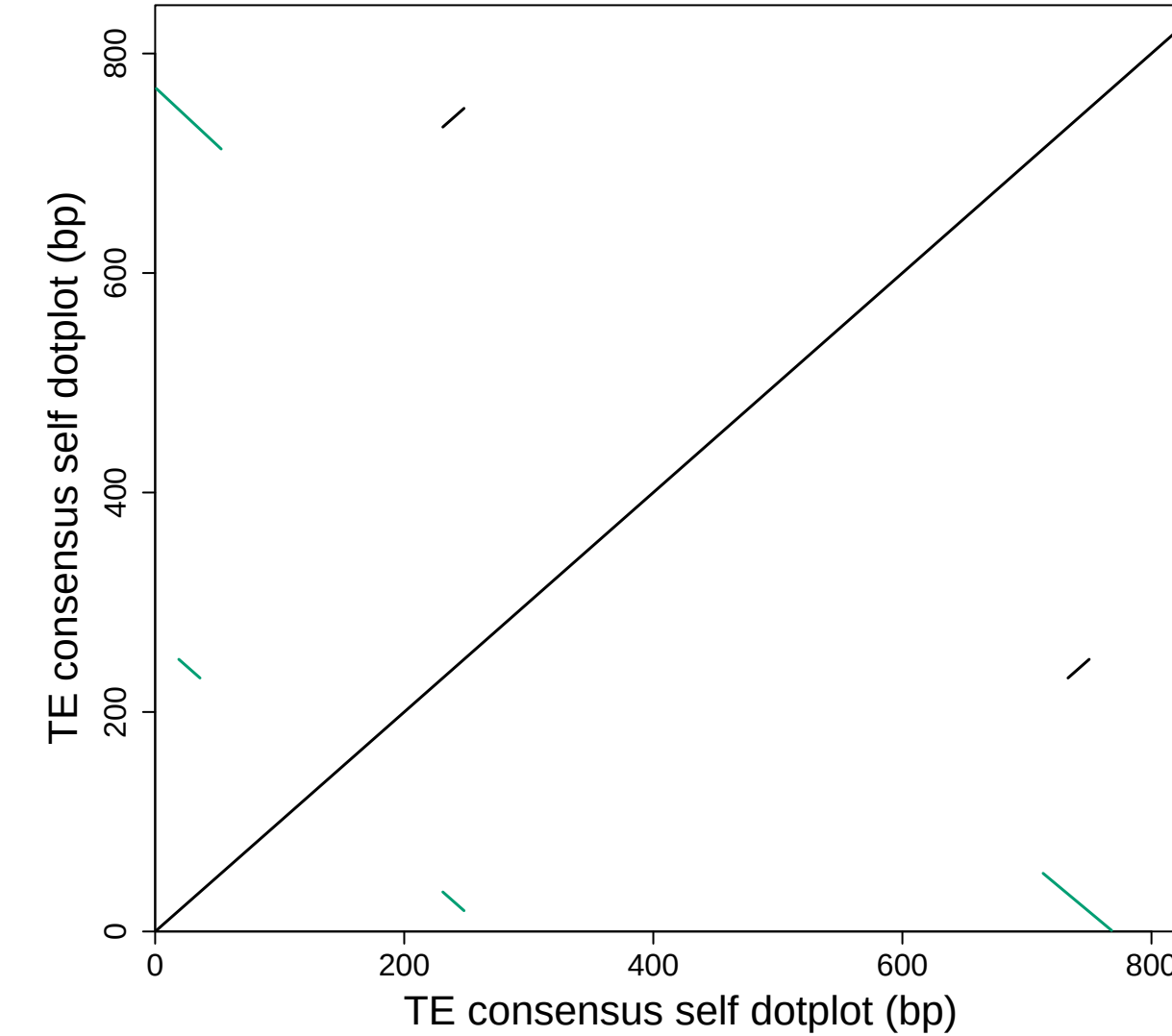
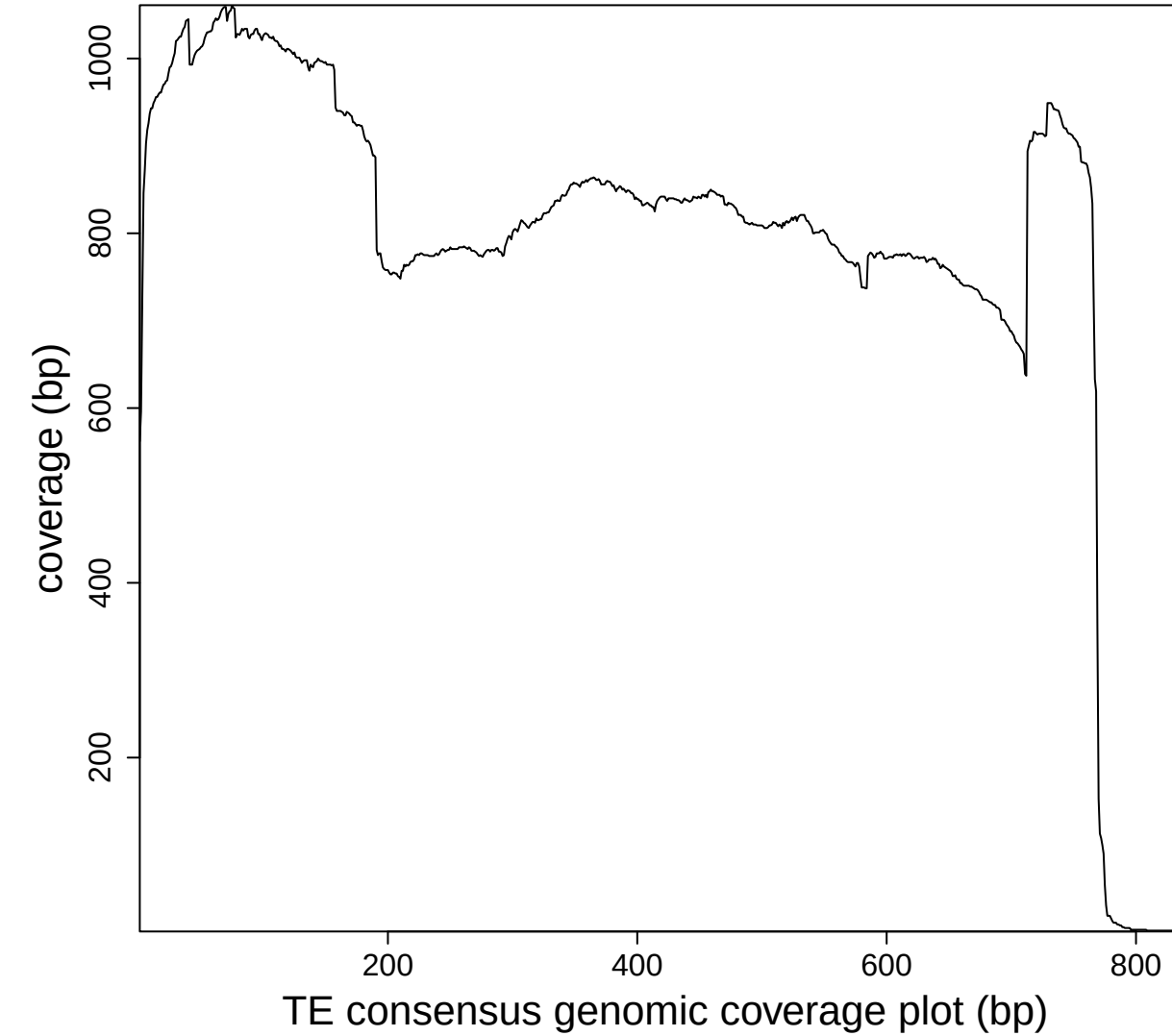
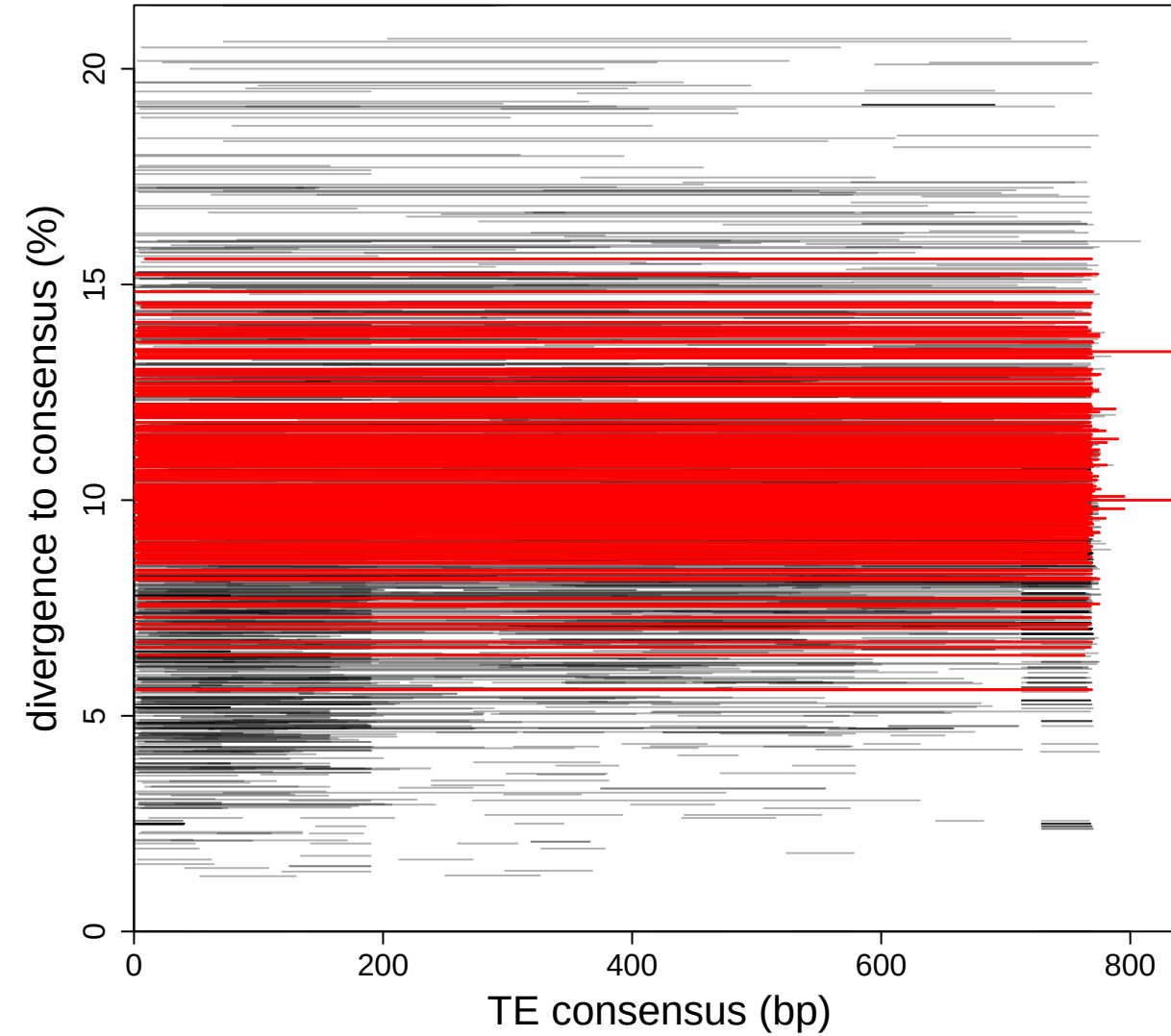
After TEtrimmer Extended plot Blue lines are boundaries



Before TEtrimmer 844 bp

TE: rnd_4_family_477

size: 844bp; fragments: 2232; full length: 235 (≥ 759.6 bp)



Dotplot

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

