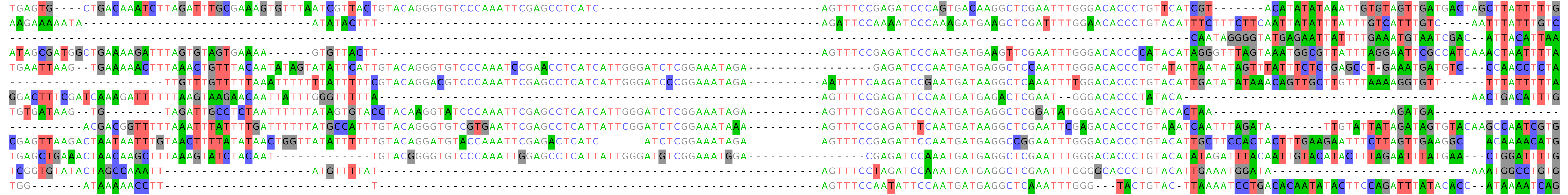


Start crop Point

End crop Point

MSA length = 987



Start crop Point

End crop Point

0

10

0

200

400

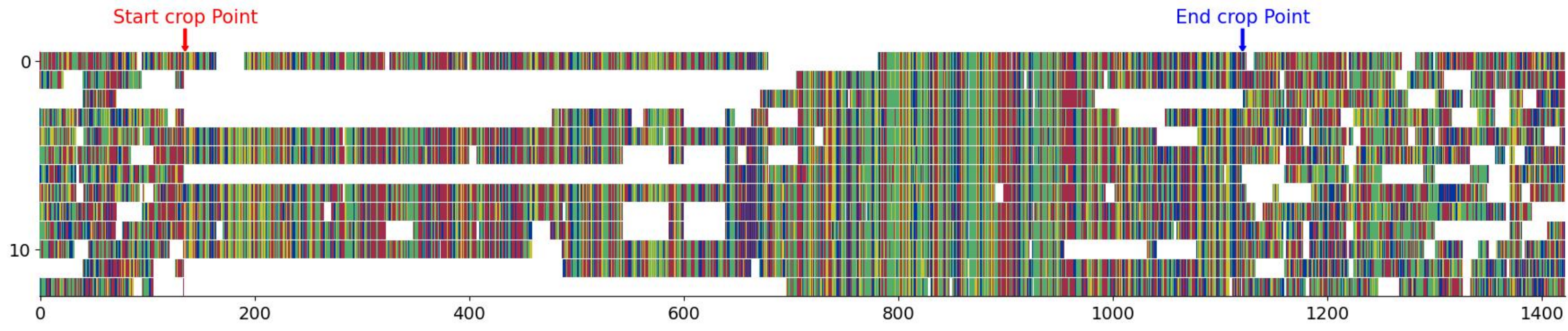
600

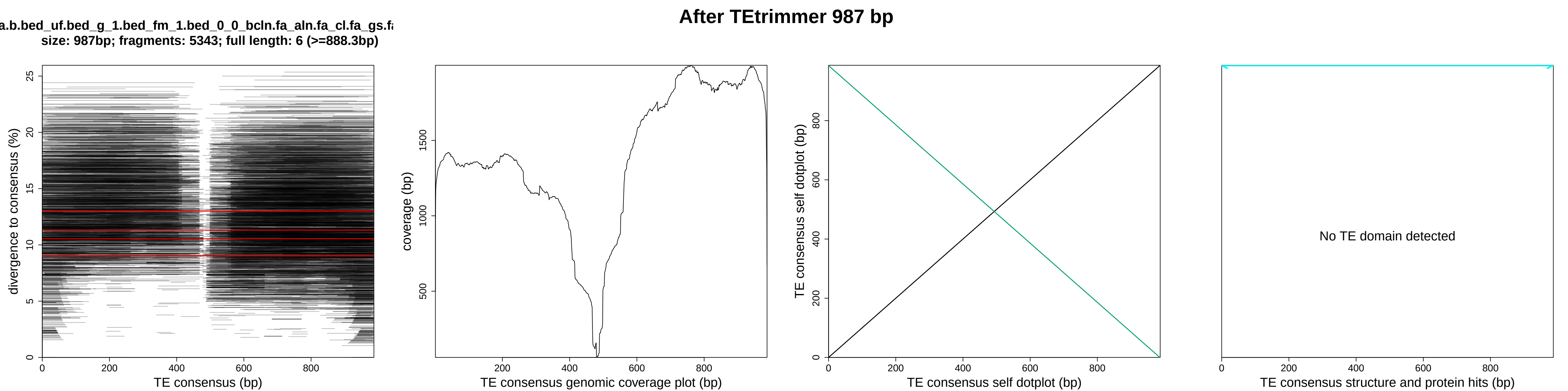
800

1000

1200

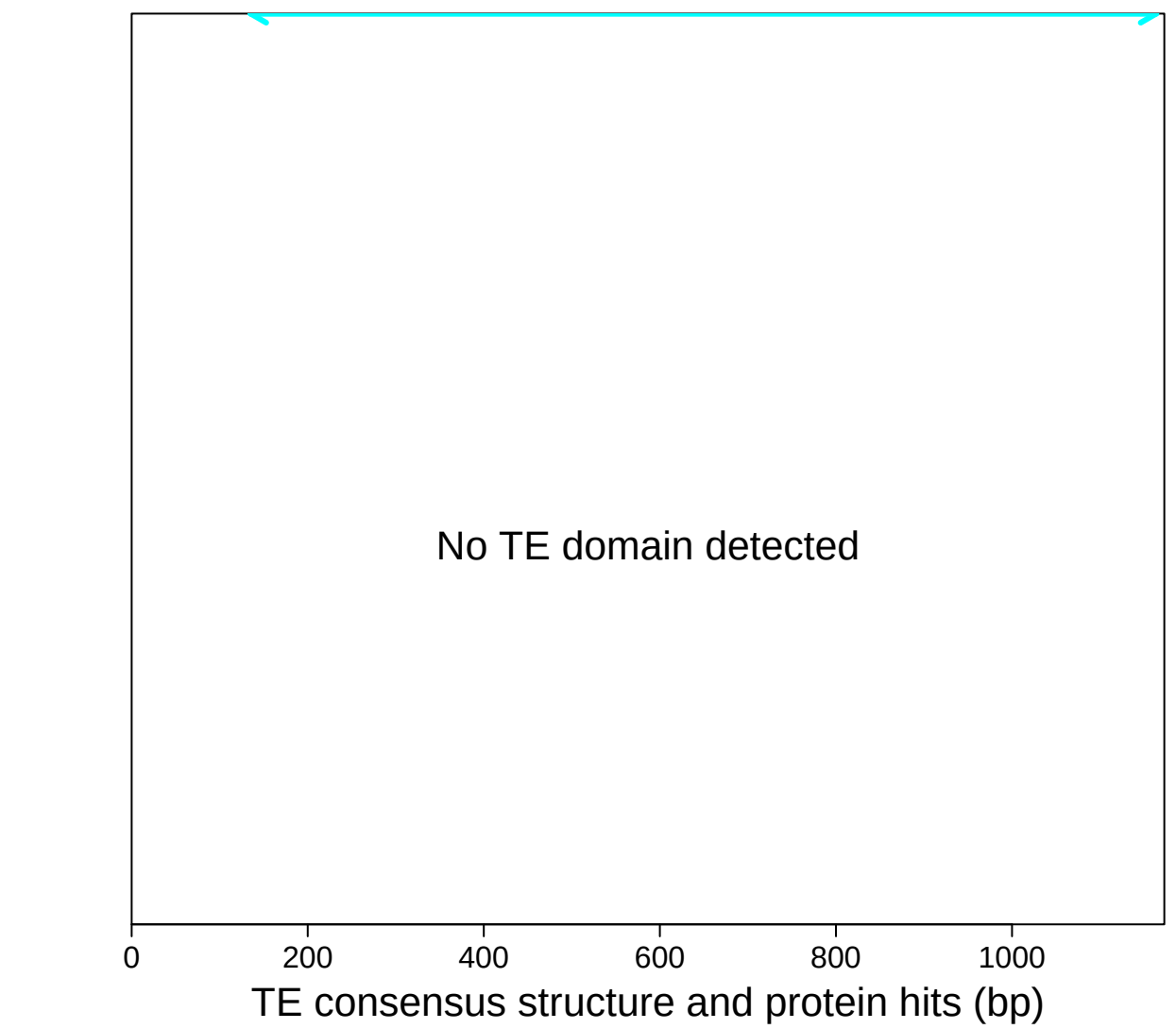
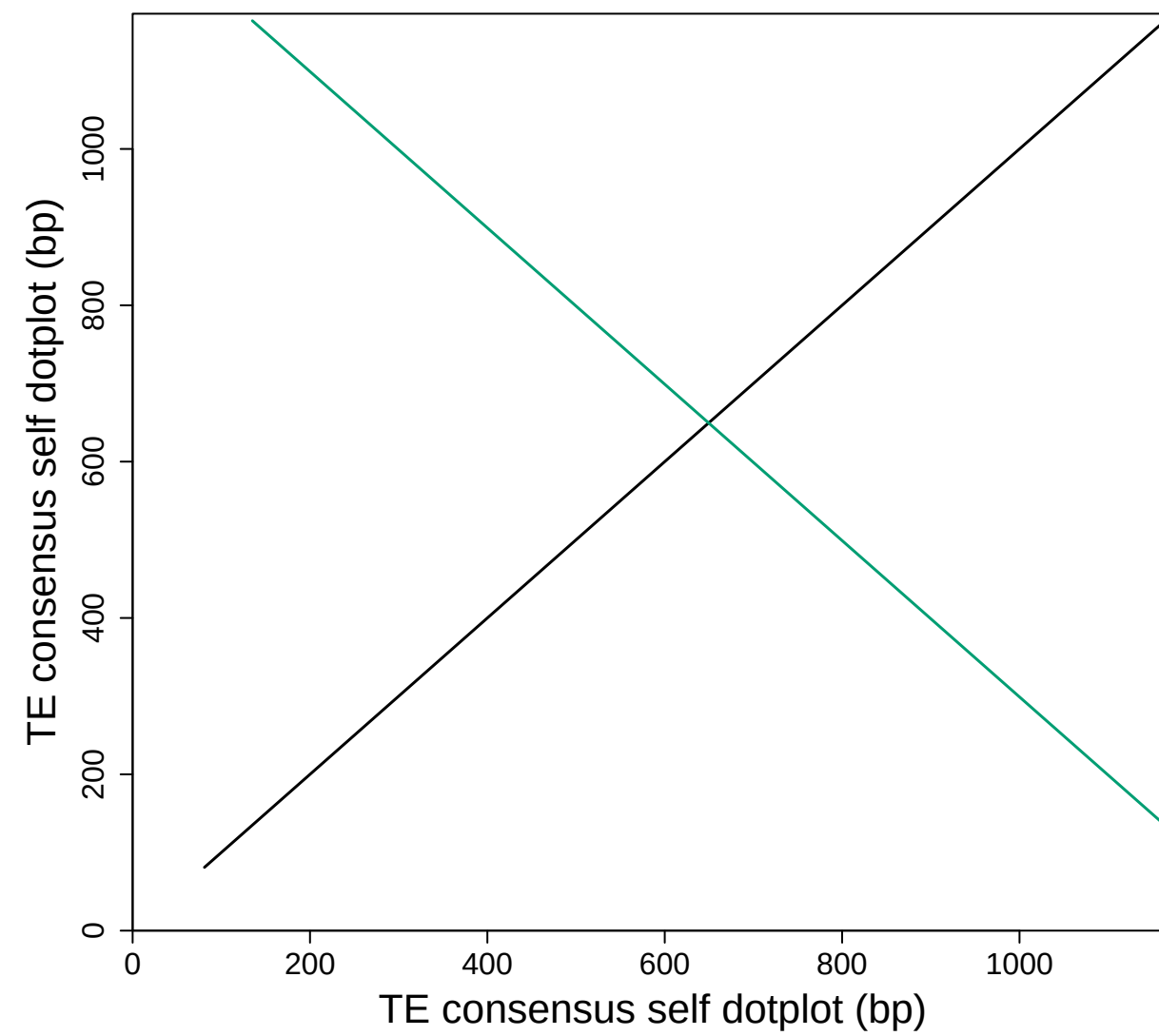
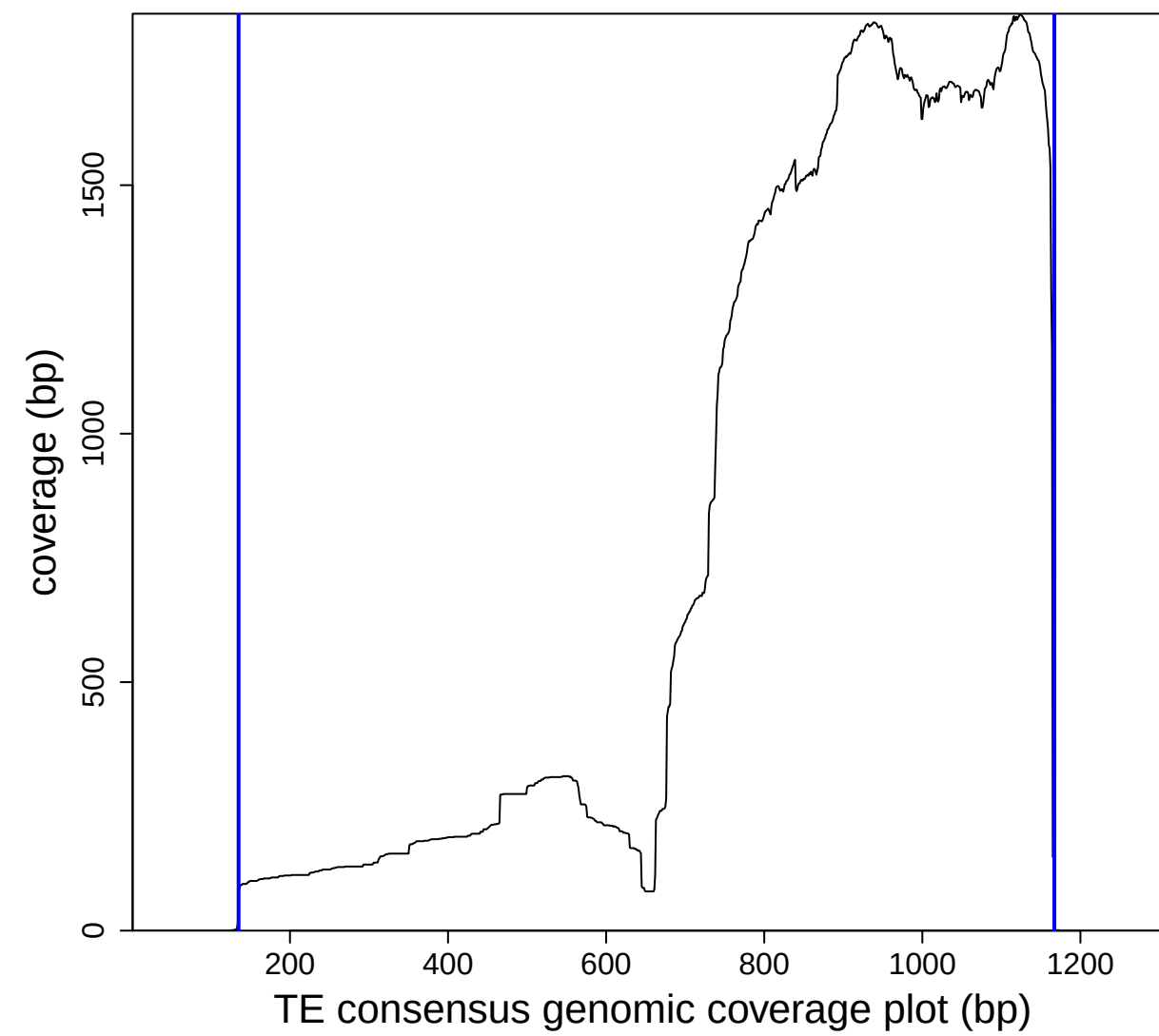
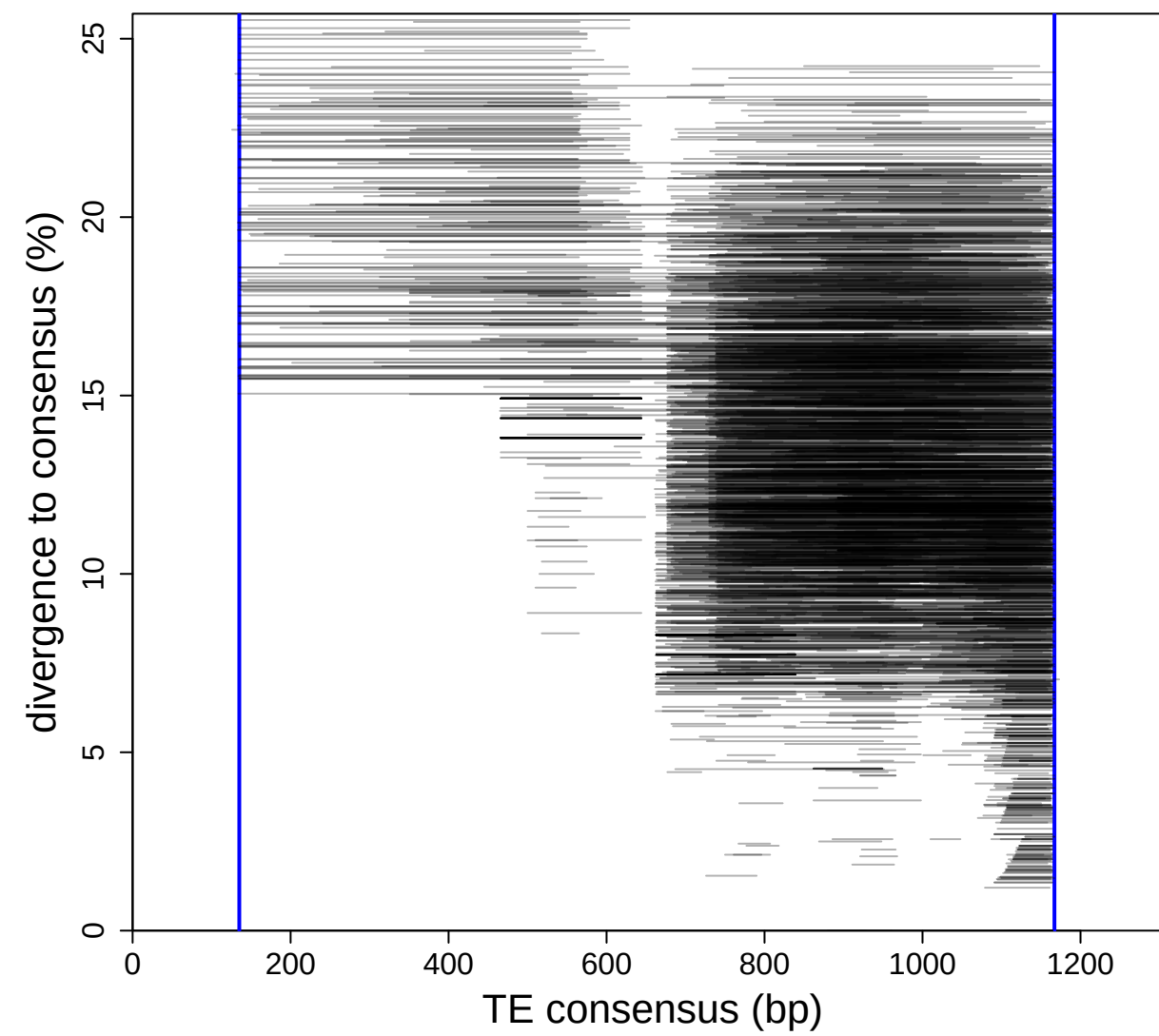
1400



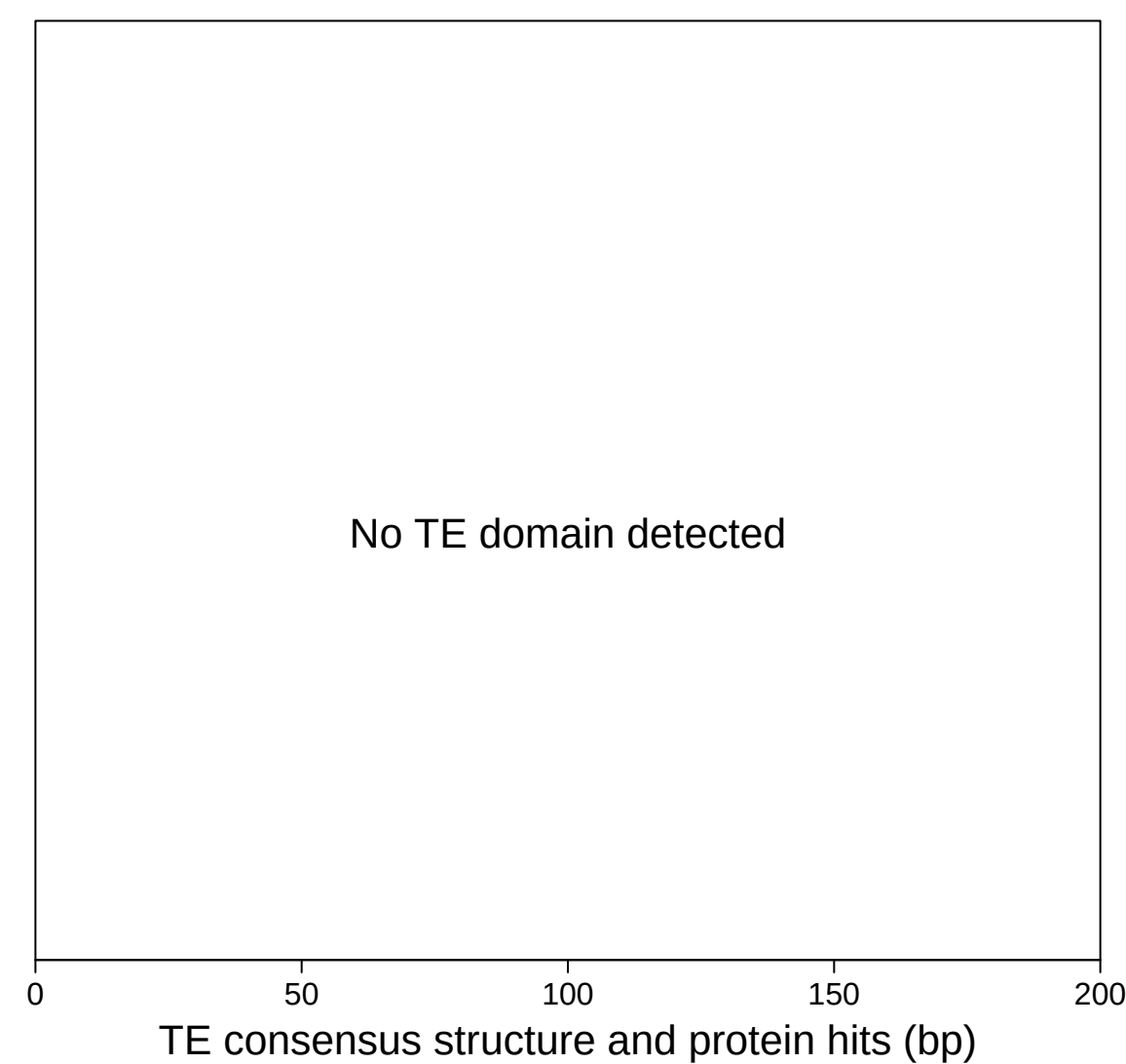
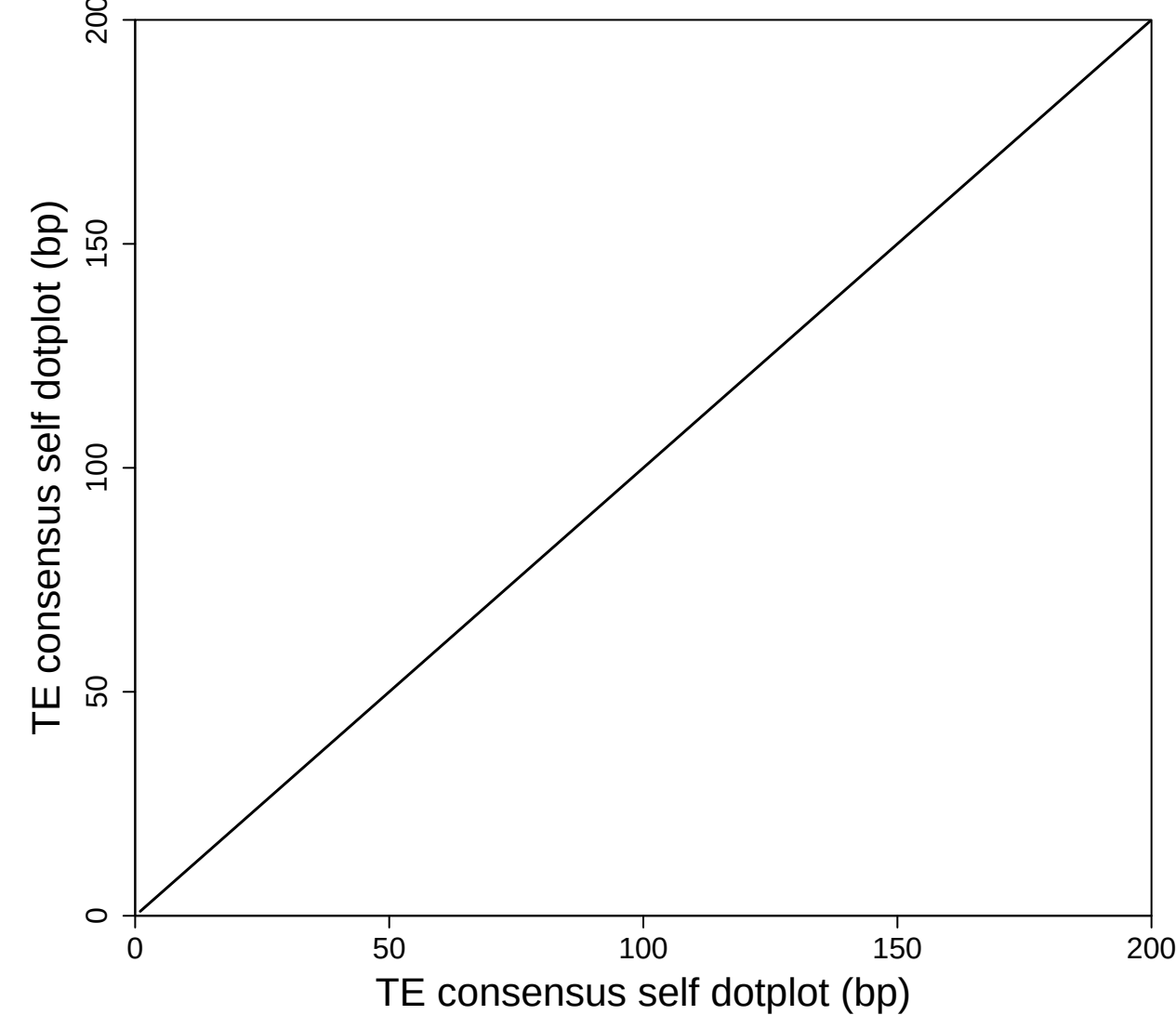
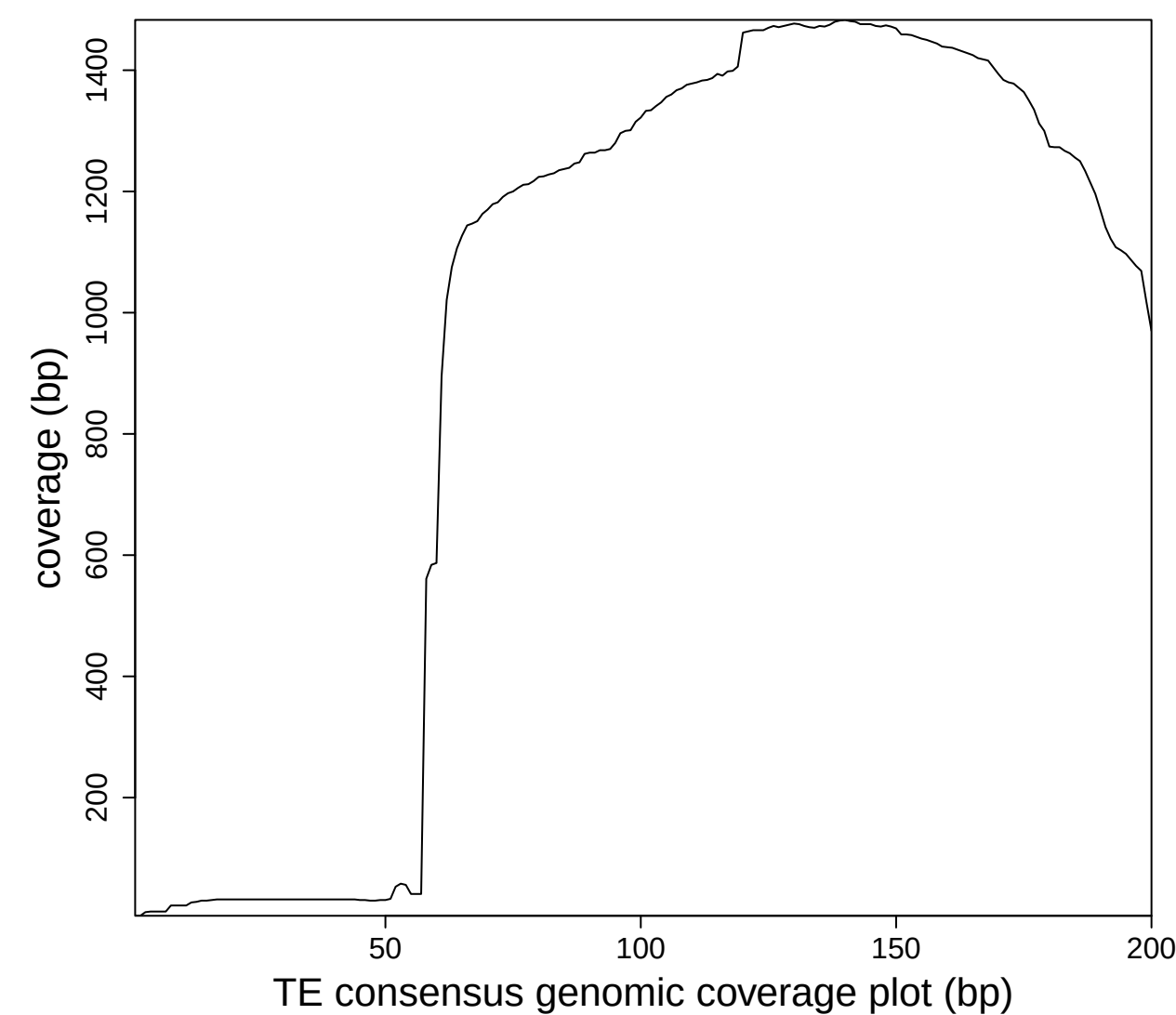
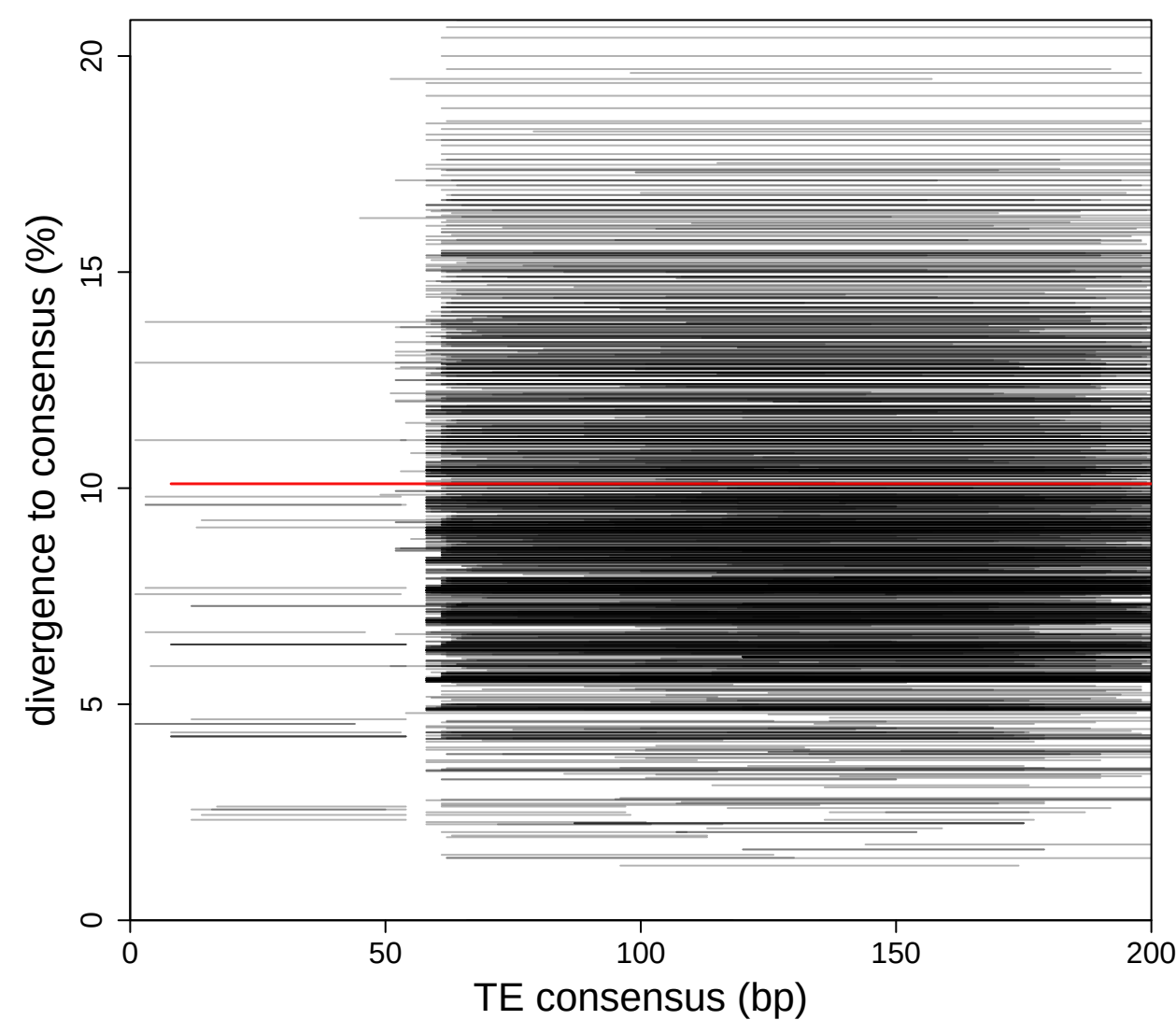


8.fasta.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 1317bp; fragments: 3101; full length: 0 (>=1185.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

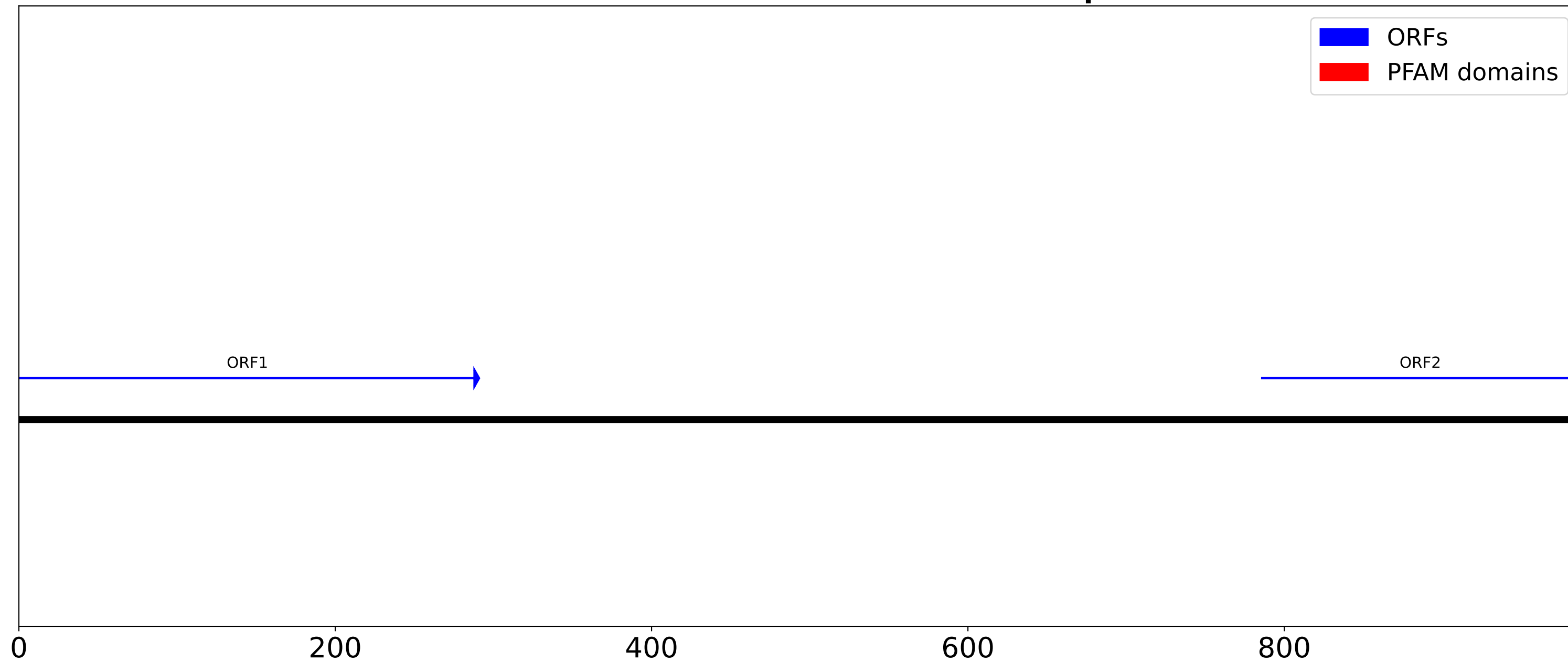


TE: rnd_4_family_1338
size: 200bp; fragments: 1582; full length: 1 (≥ 180 bp)



Before TEtrimmer 200 bp

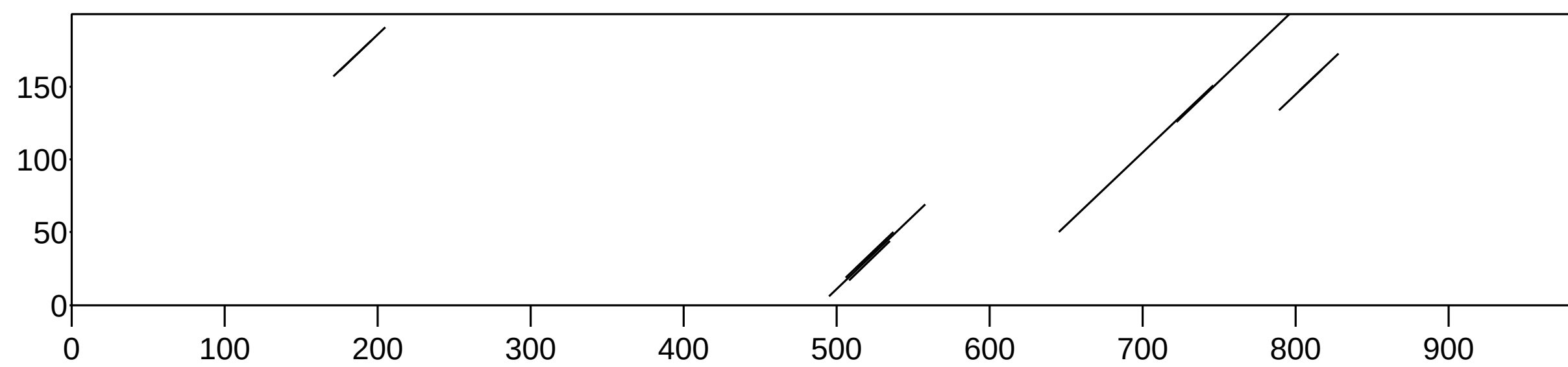
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)