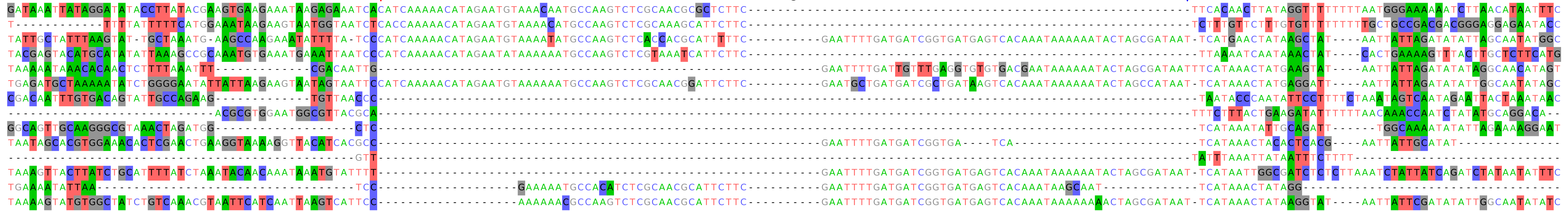


Start crop Point

End crop Point

MSA length = 1769



Start crop Point

End crop Point

0

10

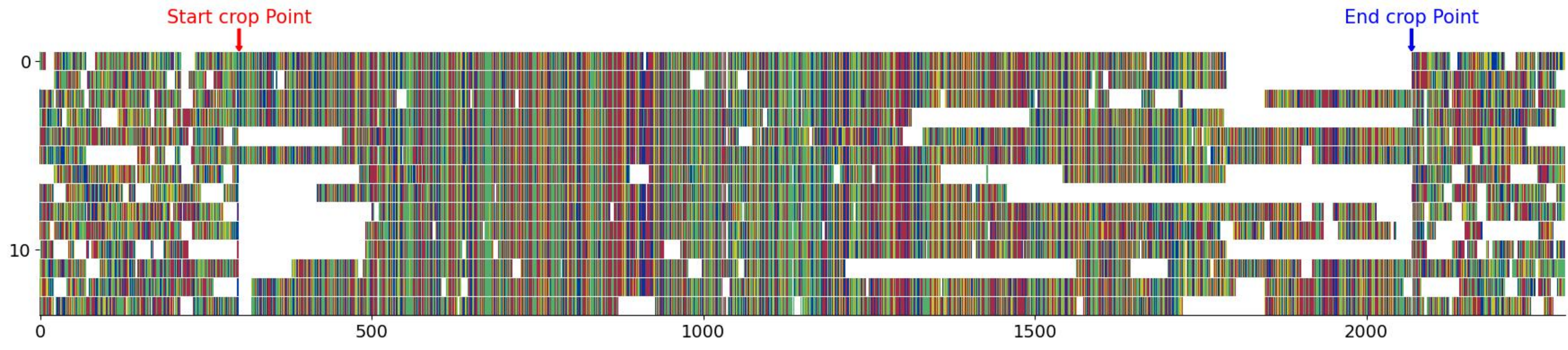
0

500

1000

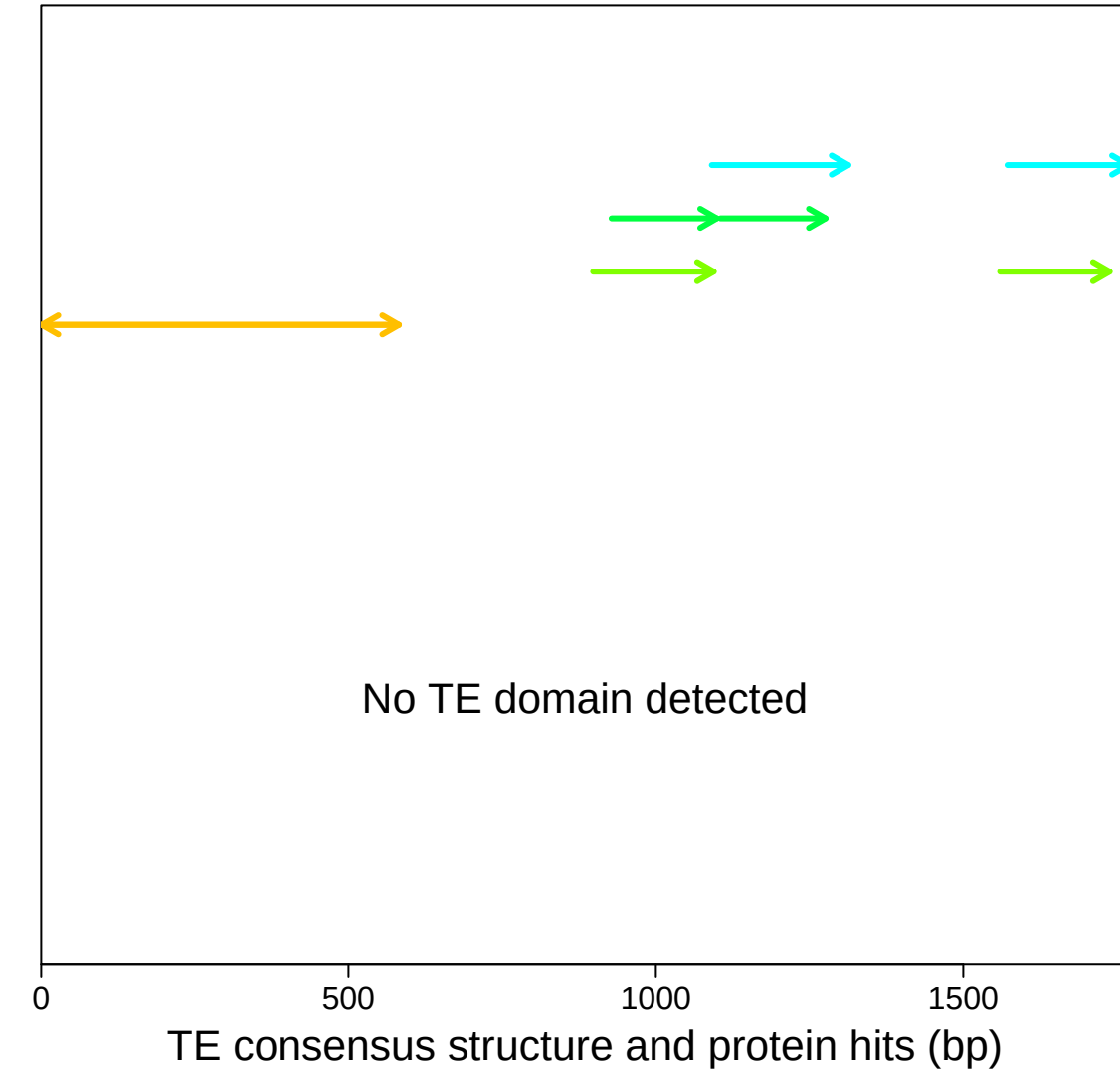
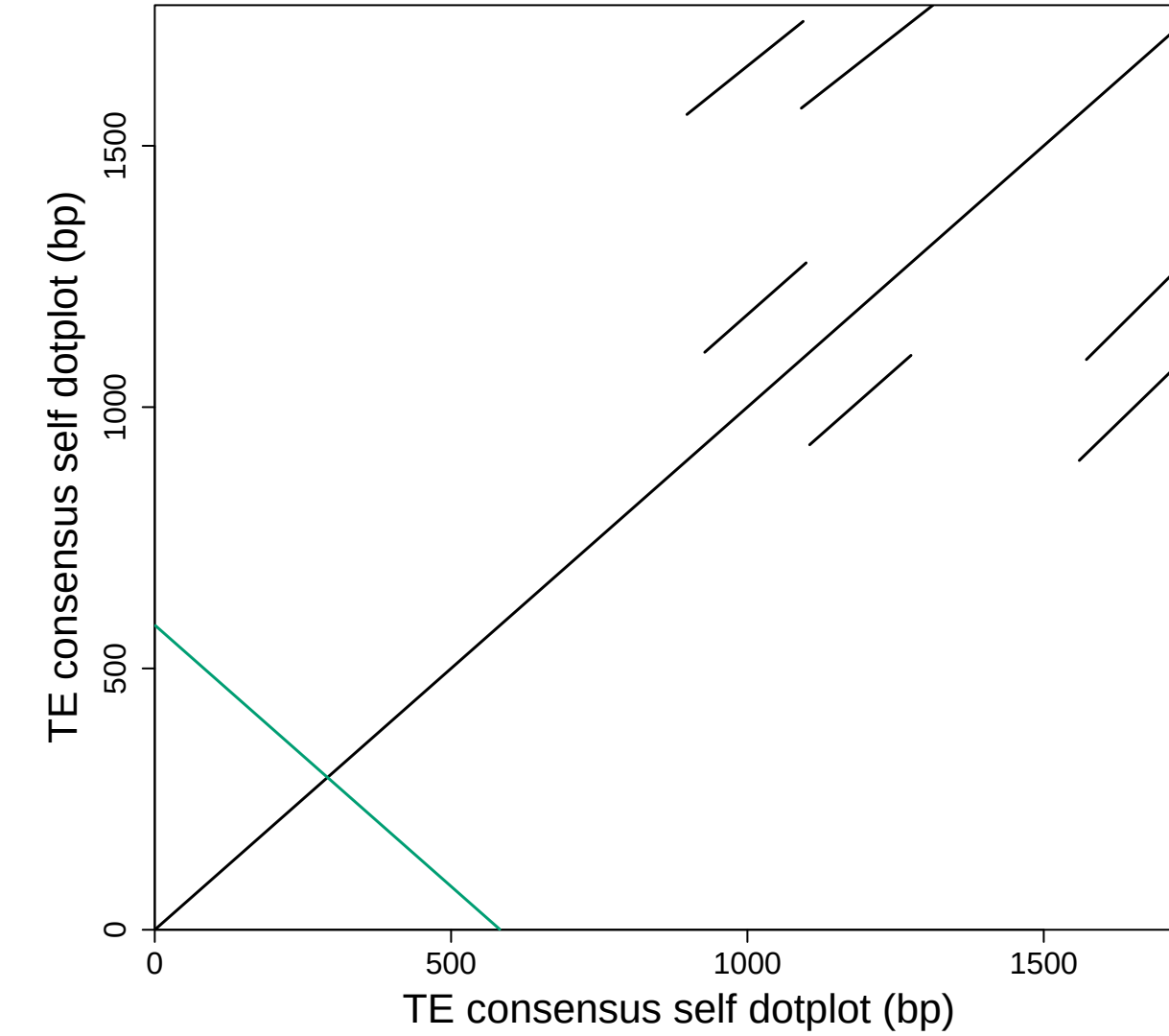
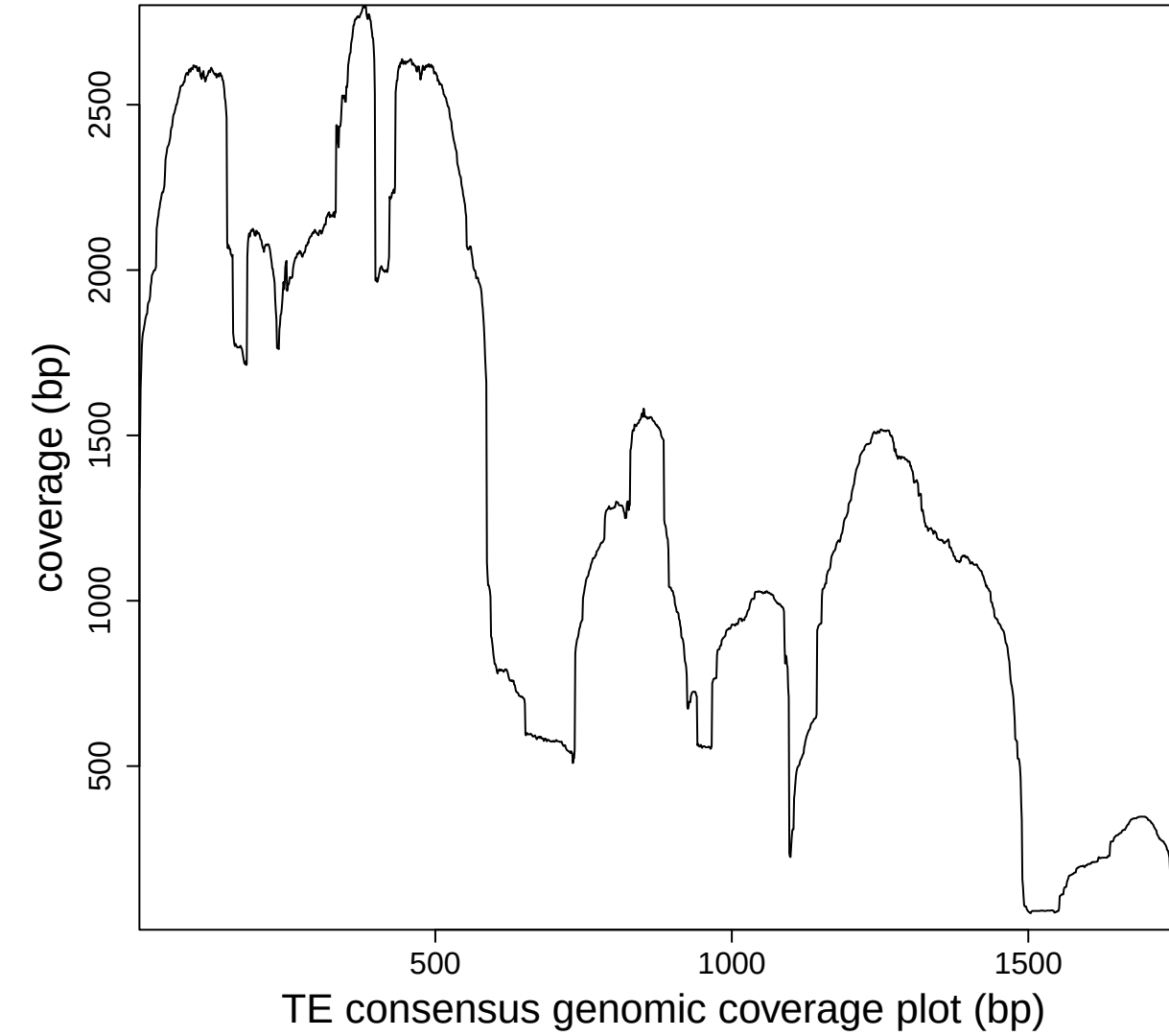
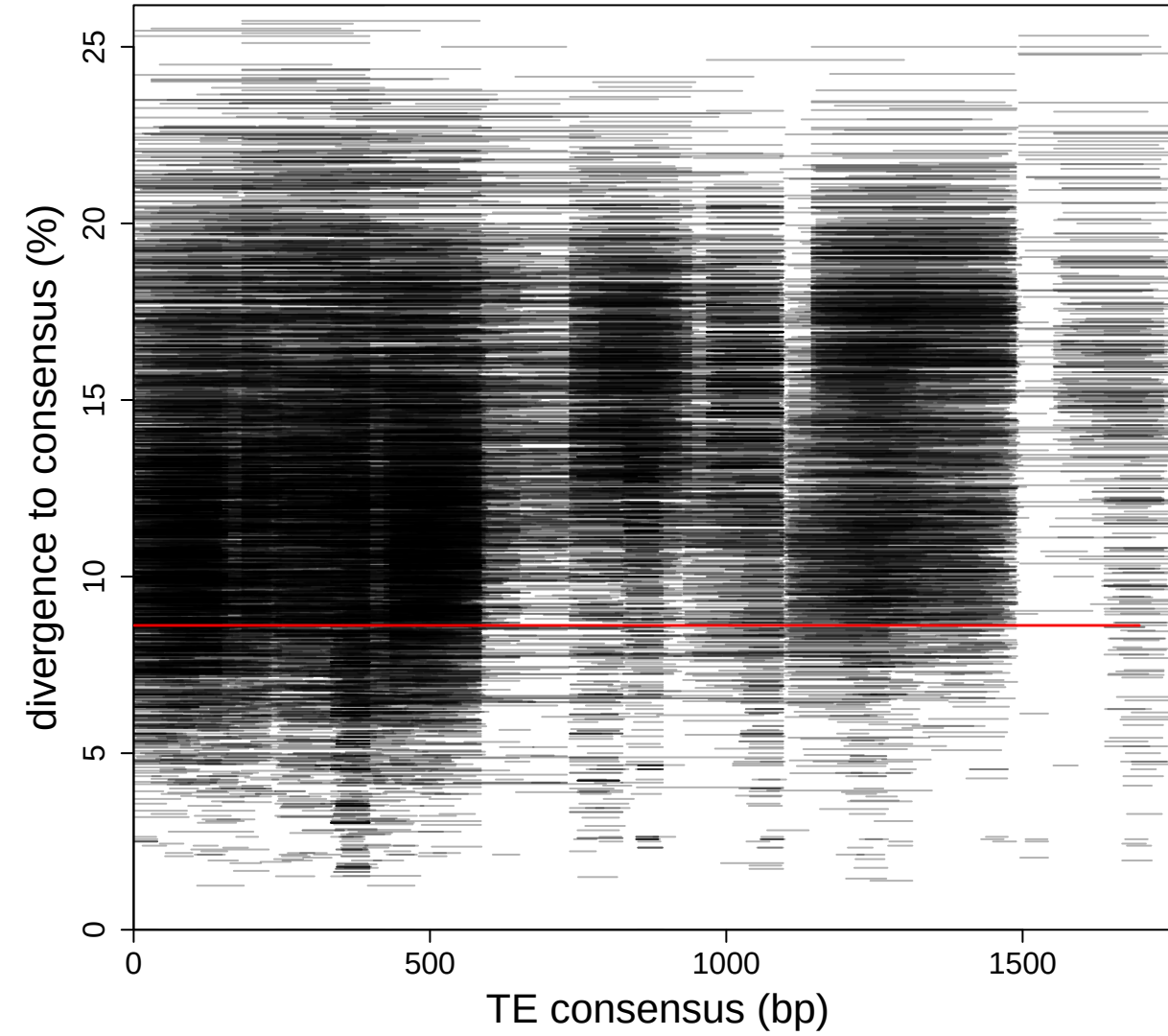
1500

2000



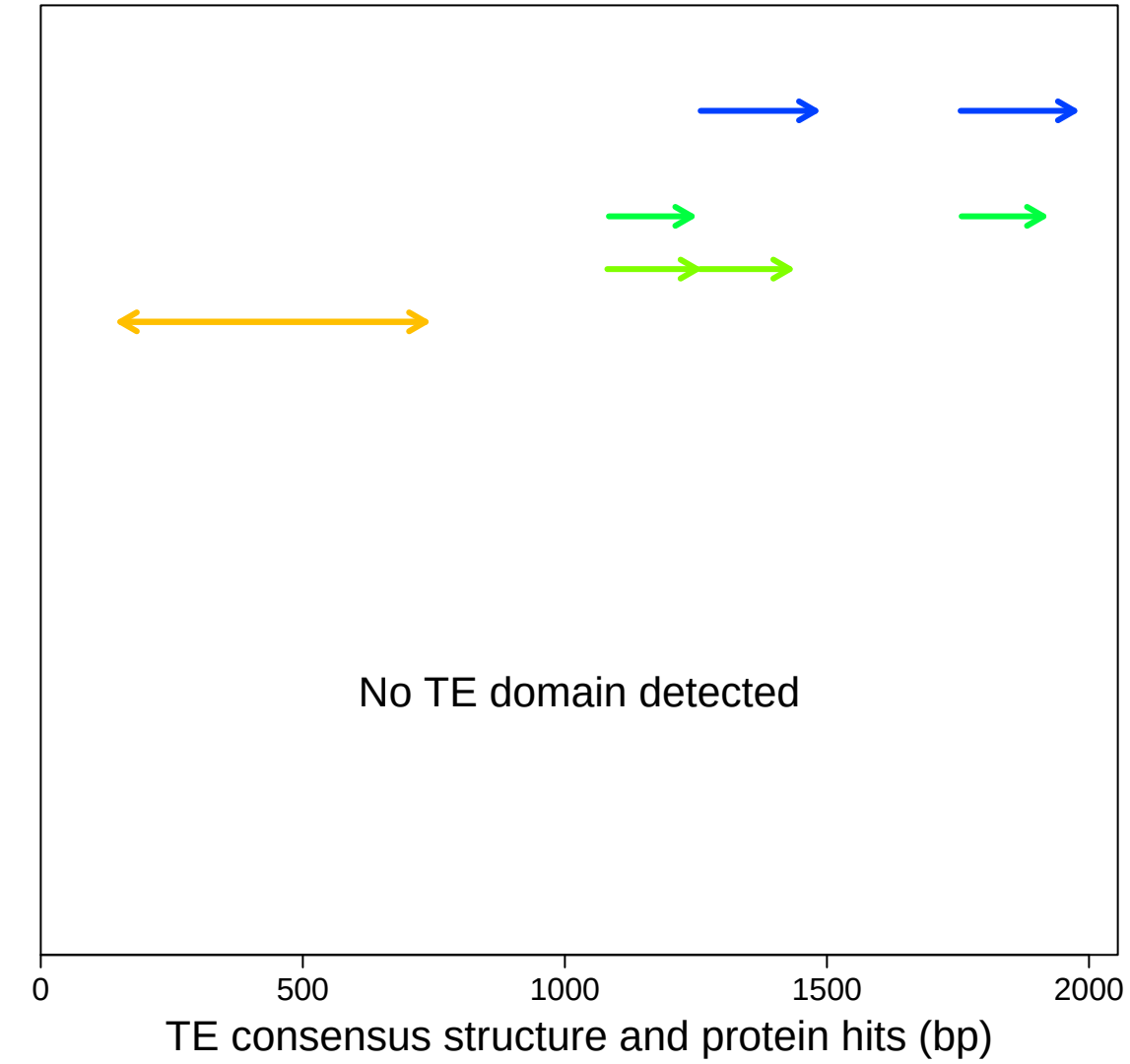
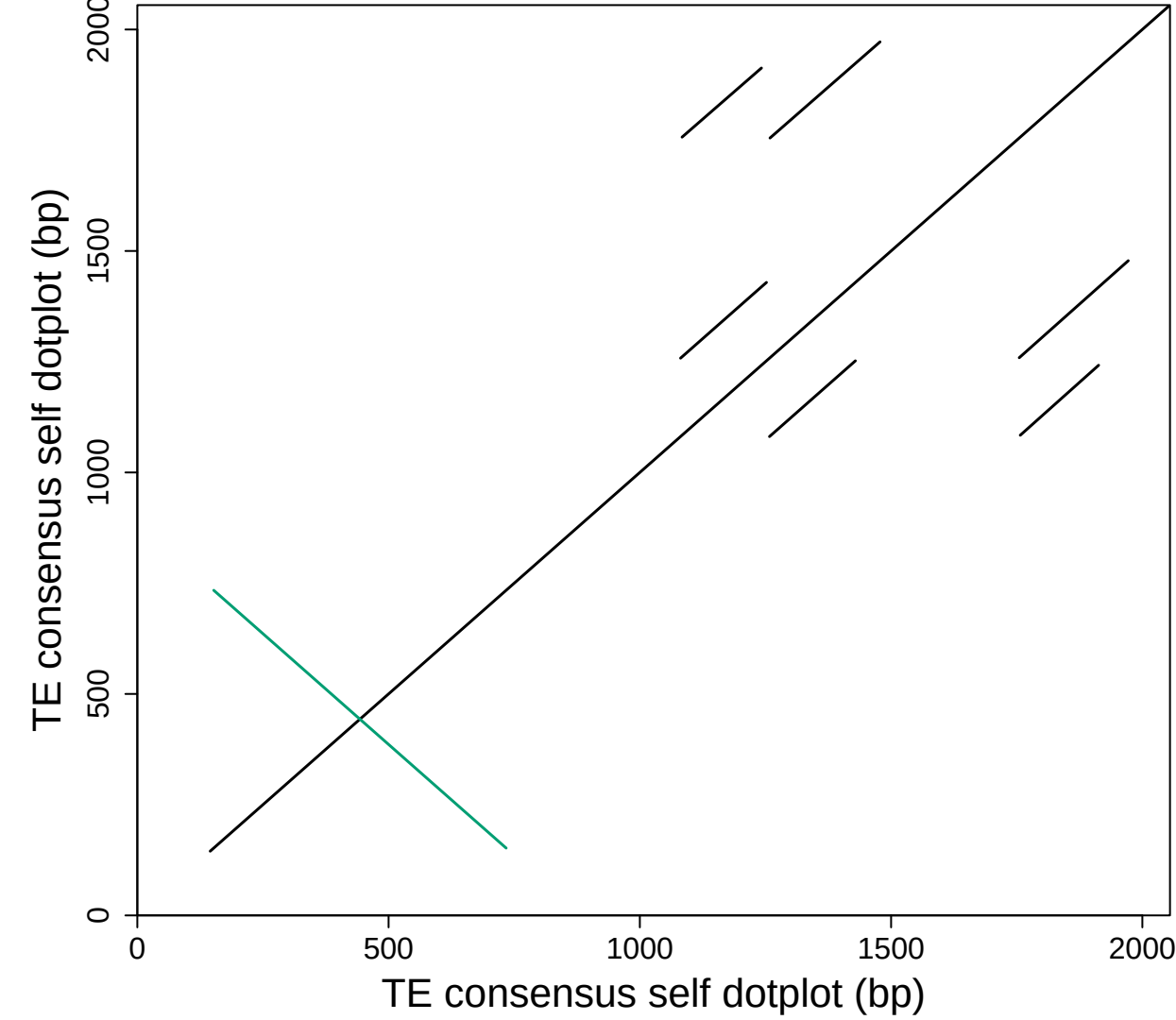
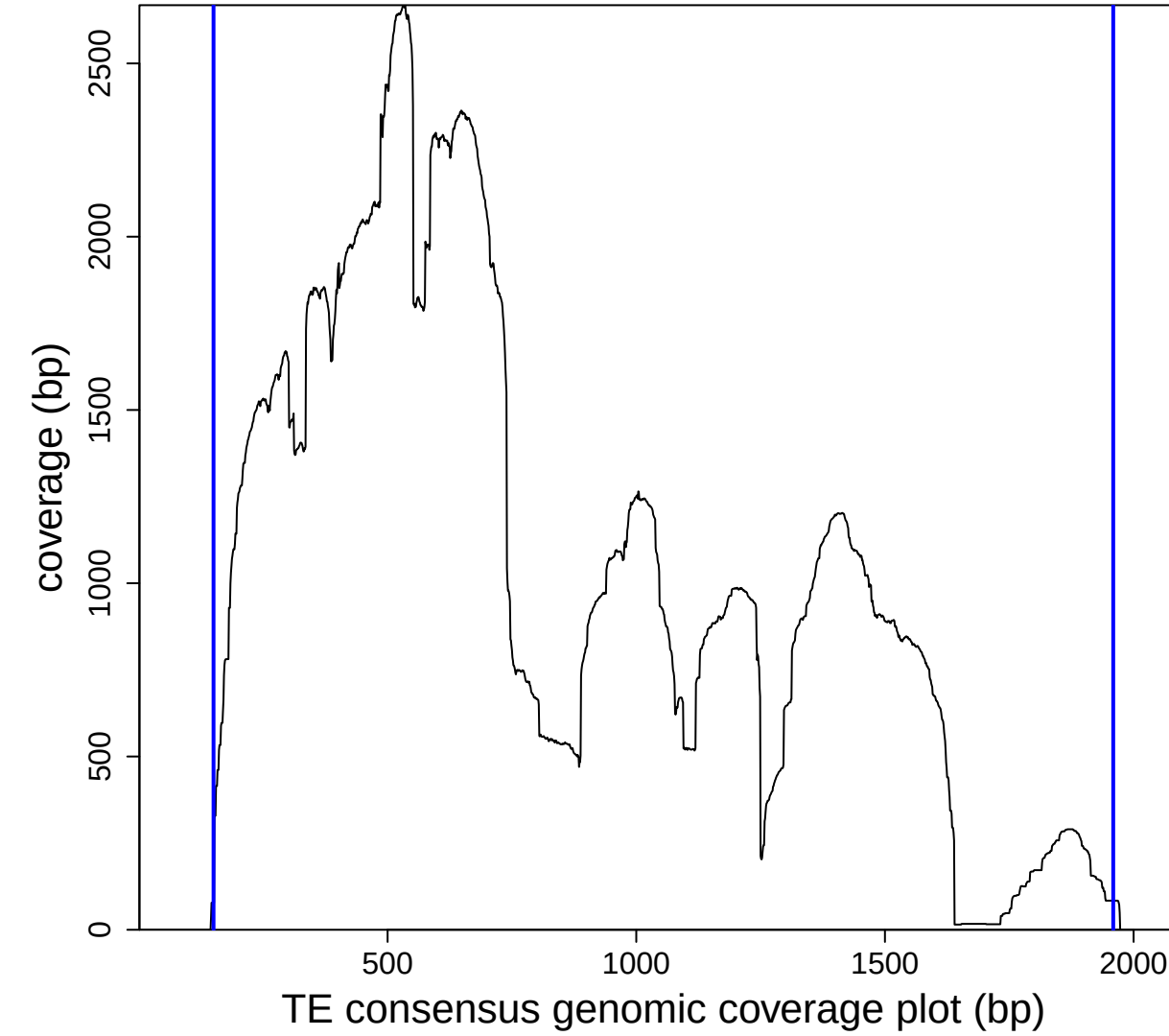
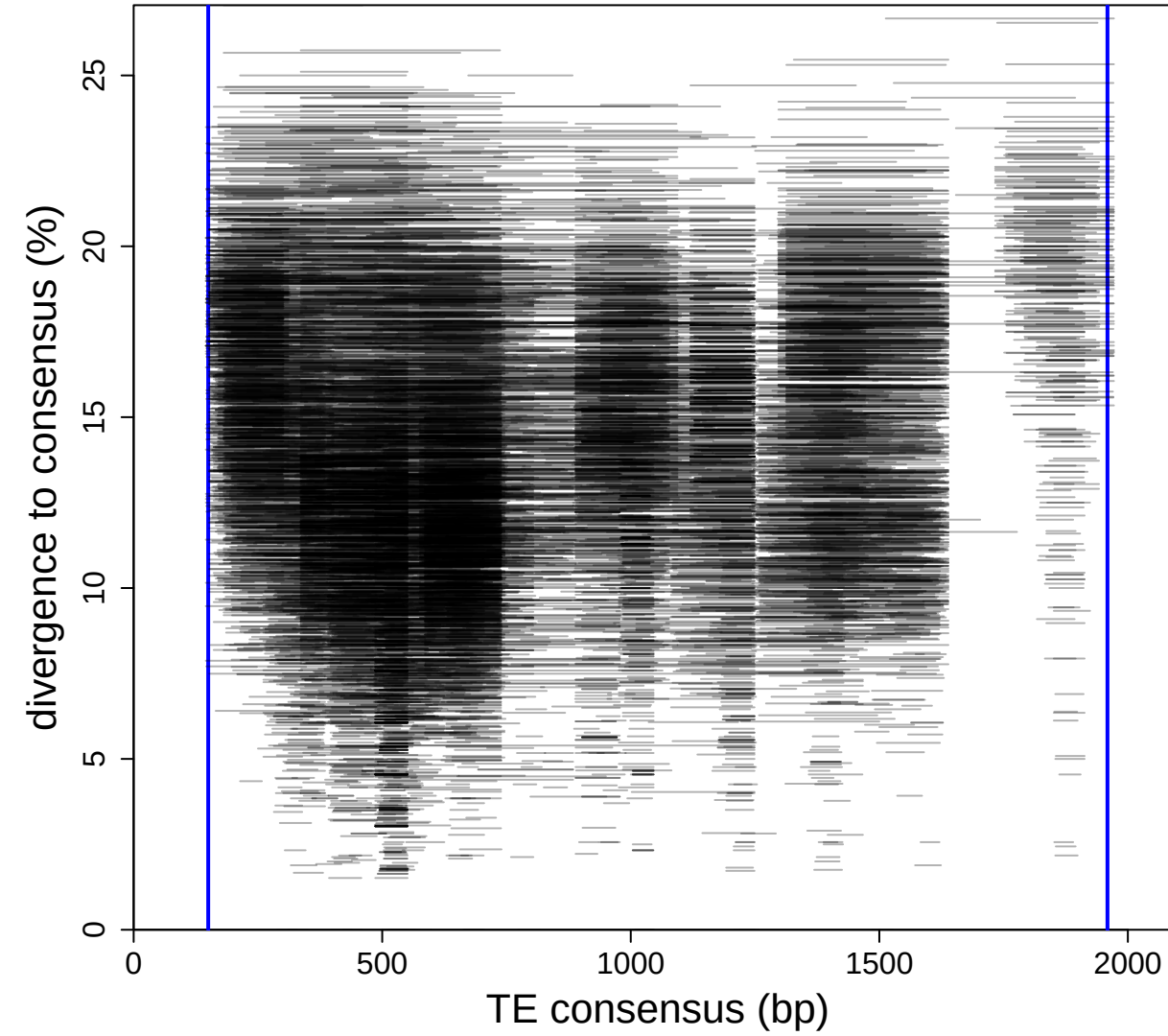
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 1769bp; fragments: 12156; full length: 1 (>=1592.1bp)

After TEtrimmer 1769 bp

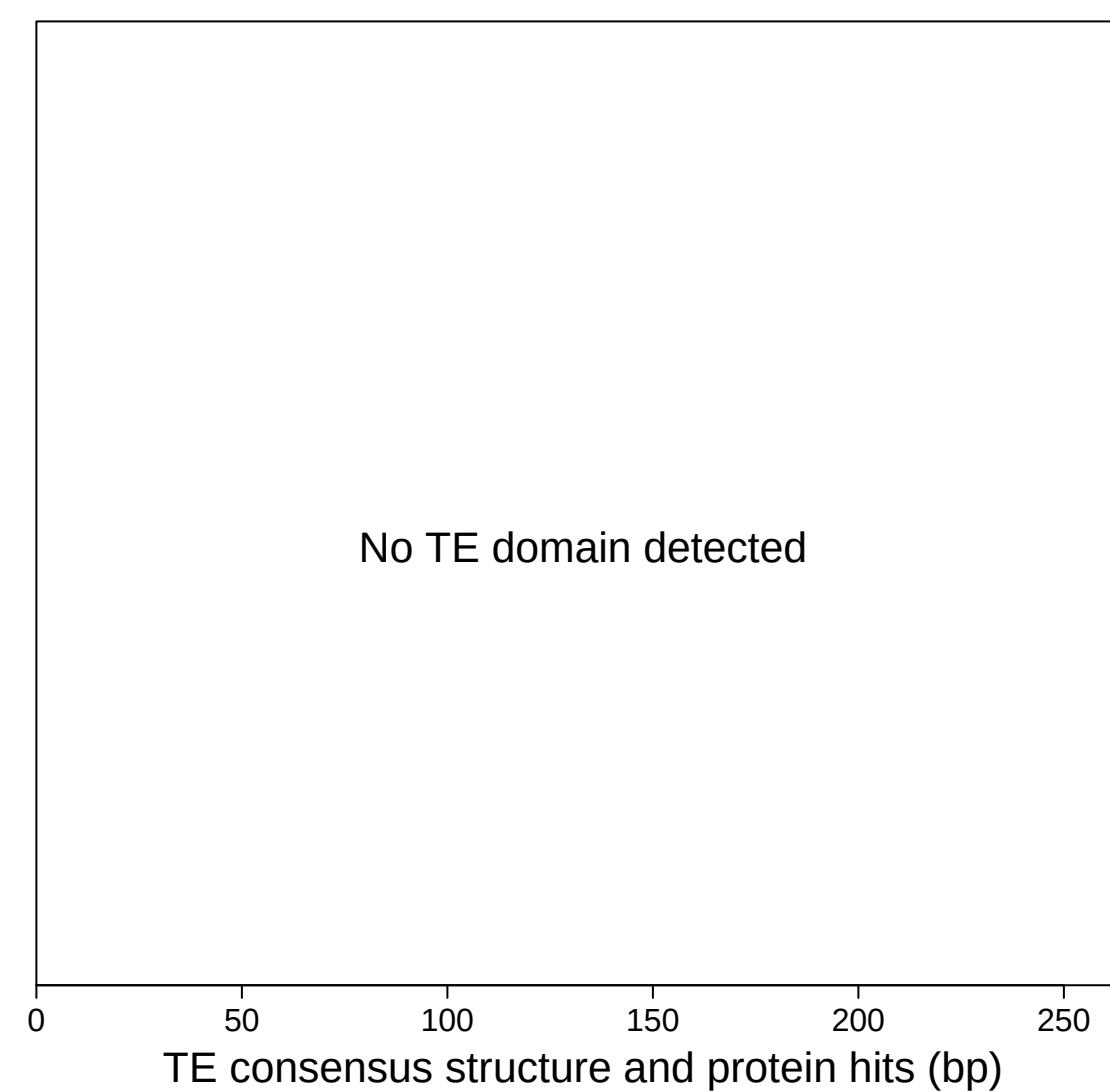
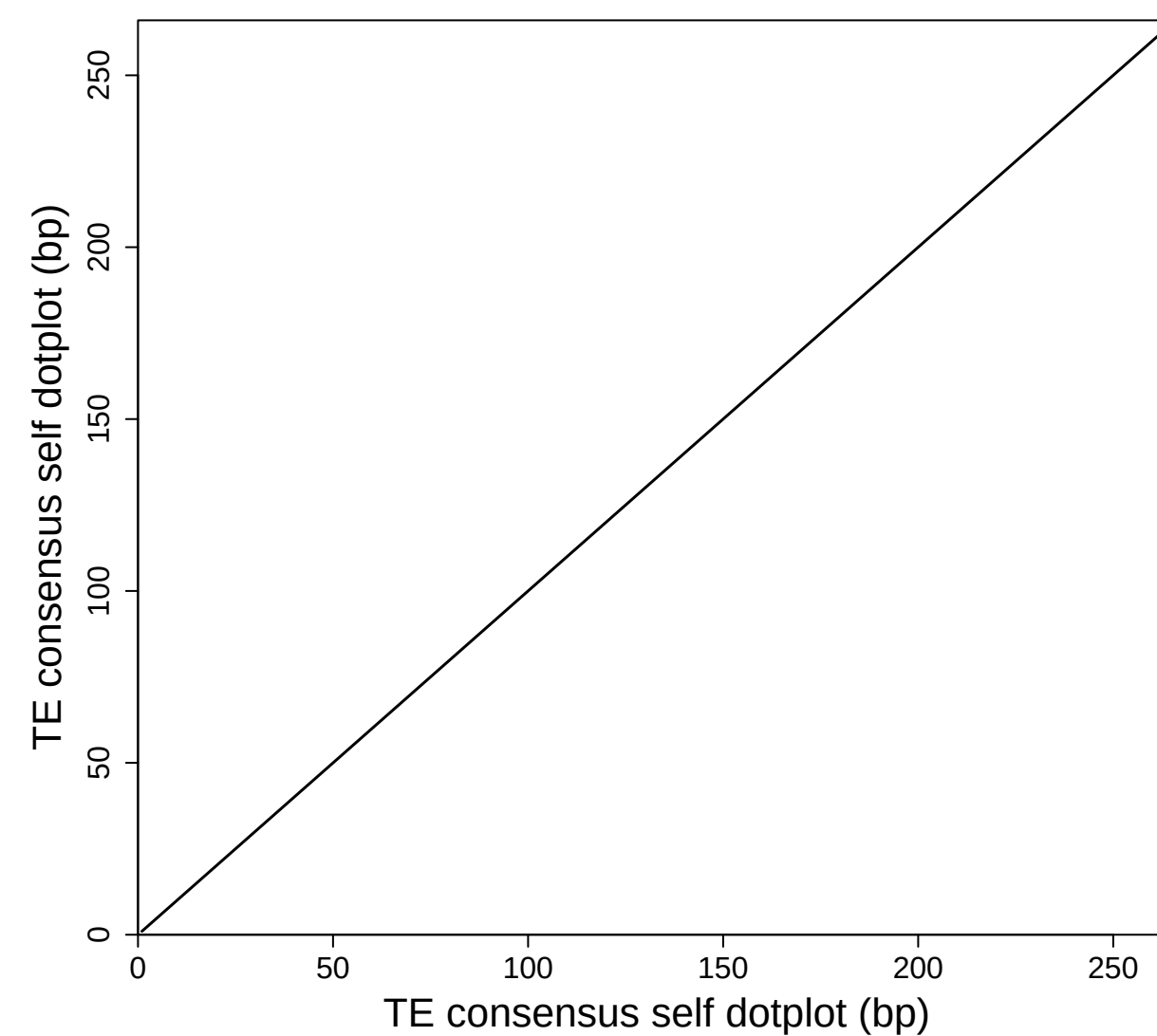
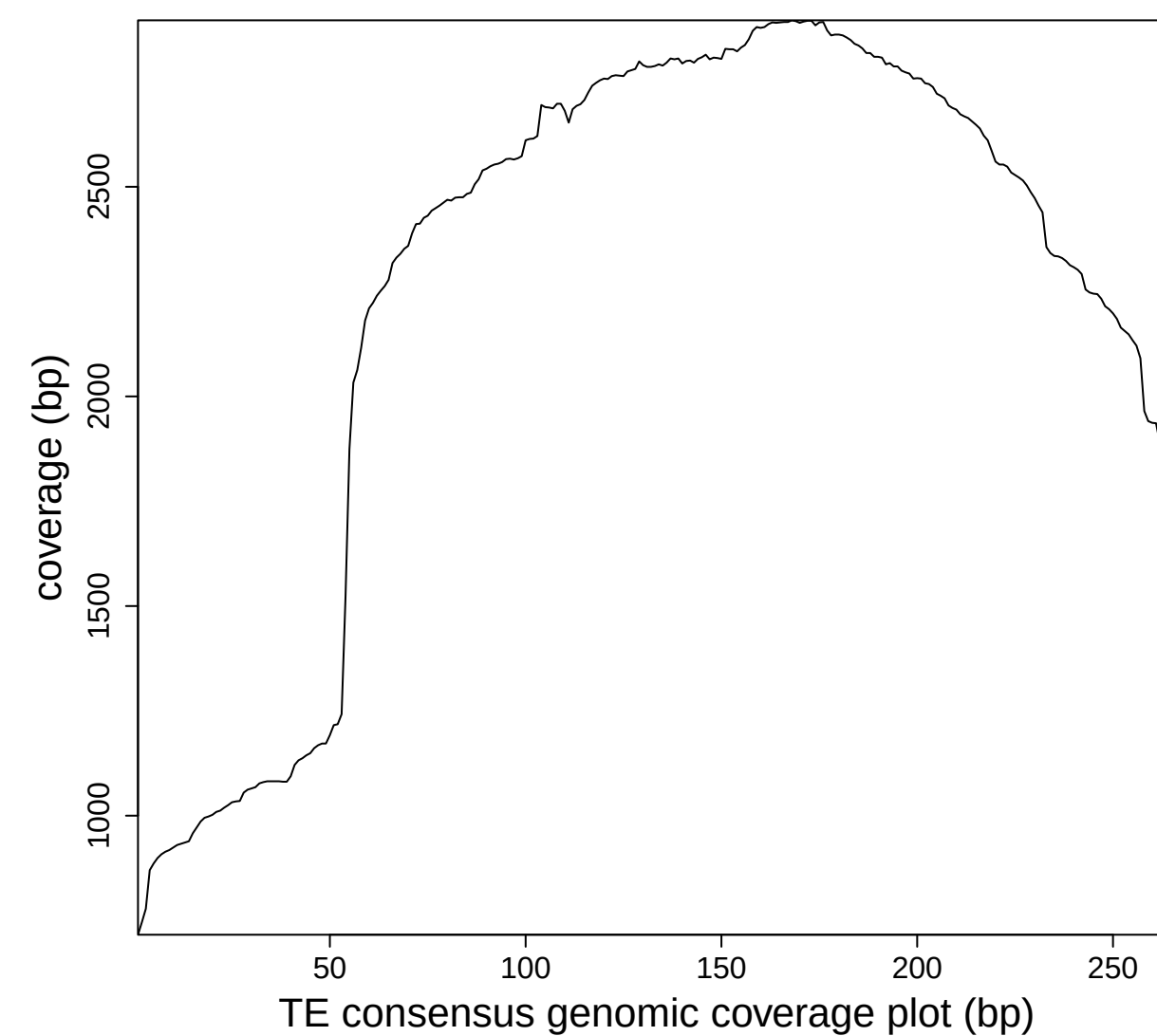
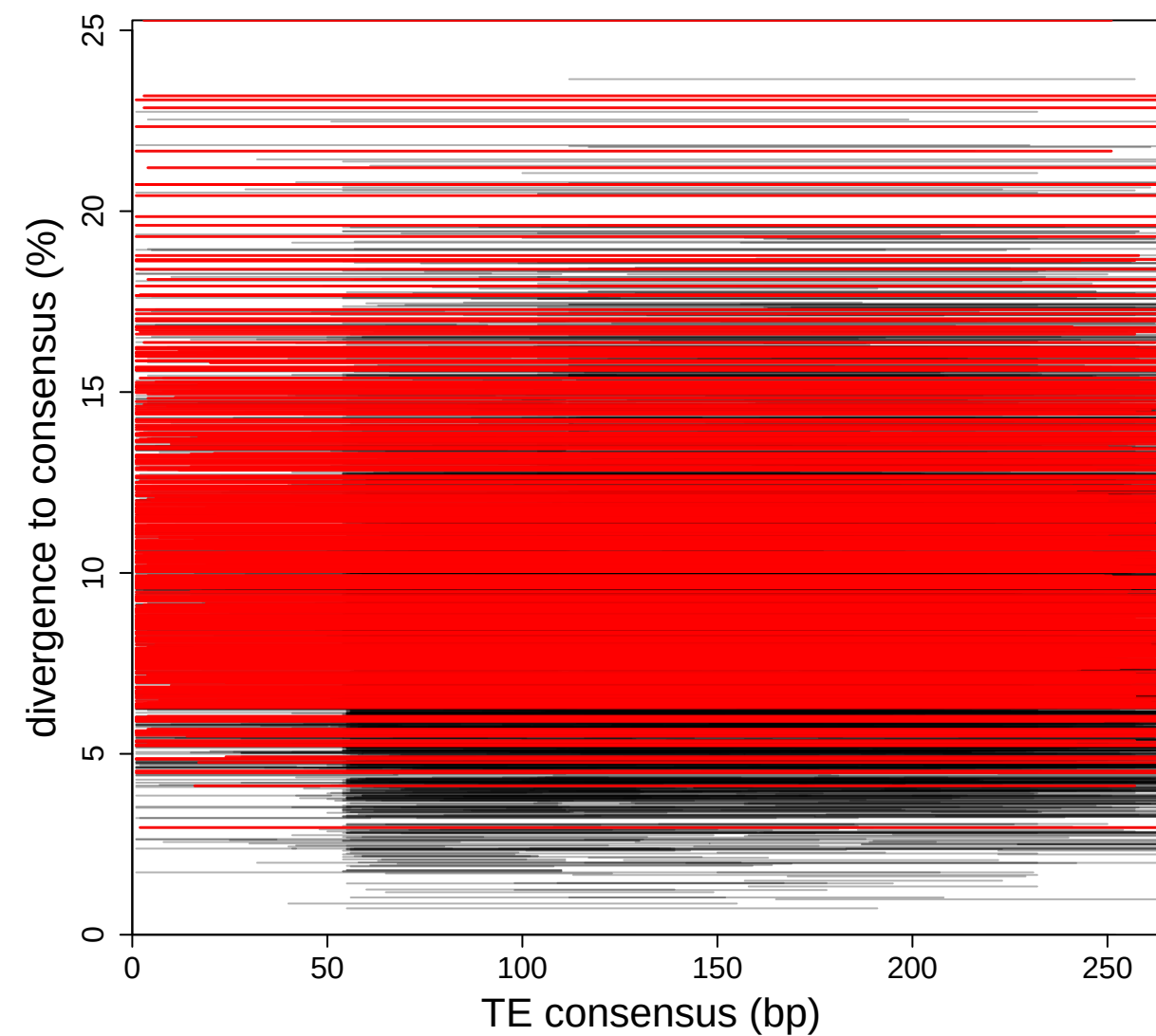


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 2109bp; fragments: 9611; full length: 0 (>=1898.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

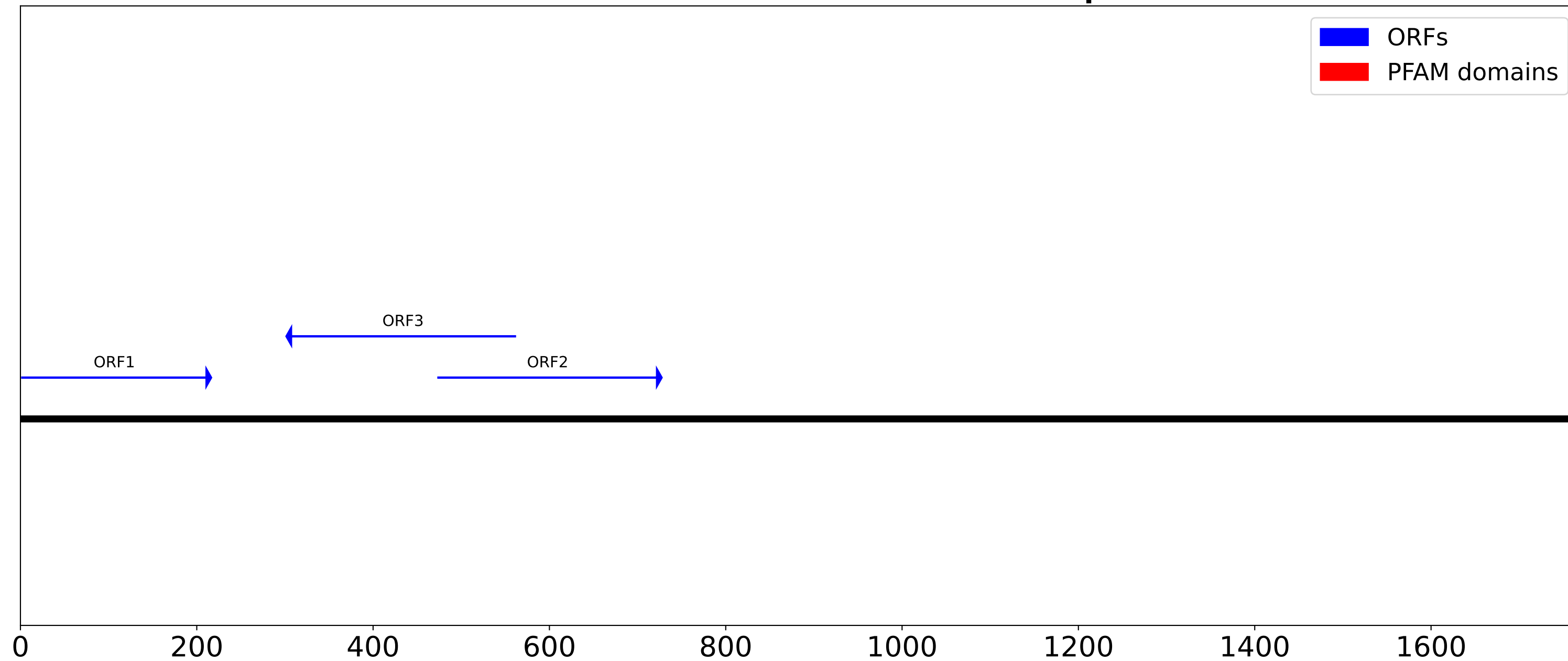


TE: rnd_1_family_150
size: 266bp; fragments: 3663; full length: 531 (≥ 239.4 bp)



Before TEtrimmer 266 bp

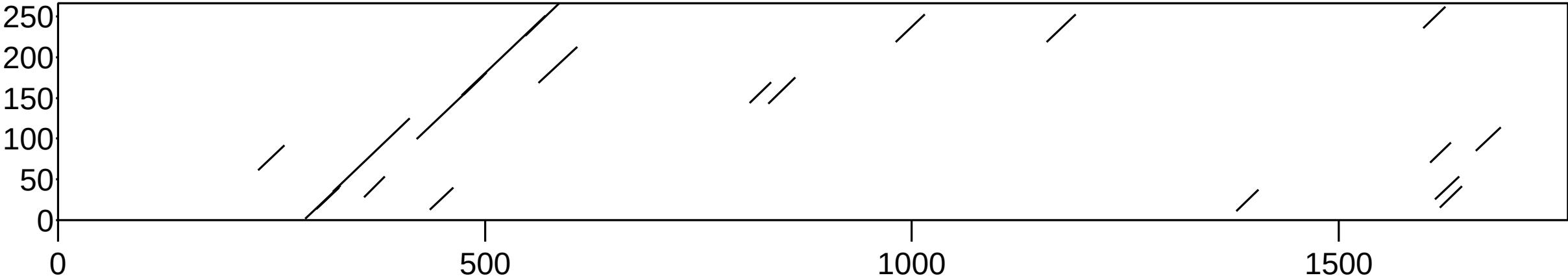
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

Isensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)