

Start crop Point



End crop Point



MSA length = 4361



Start crop Point

End crop Point

0 -

10 -

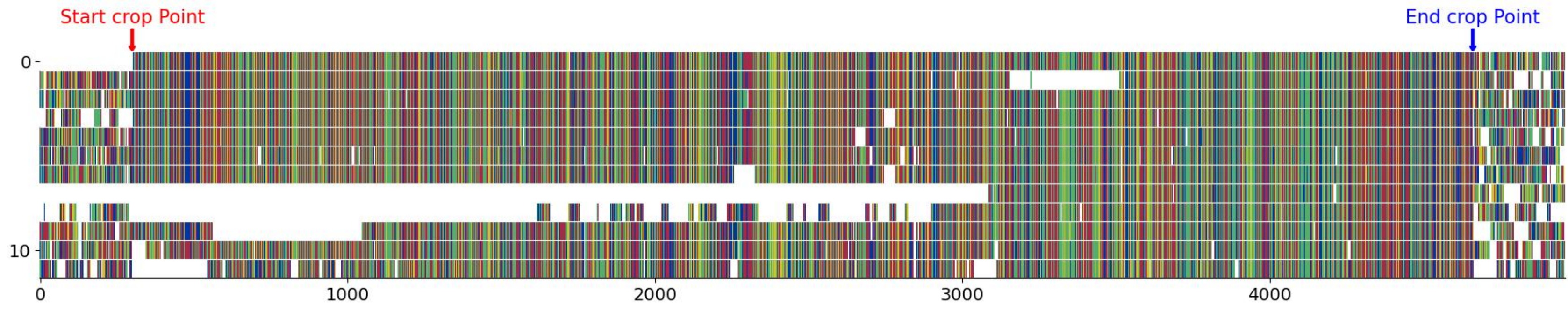
0

1000

2000

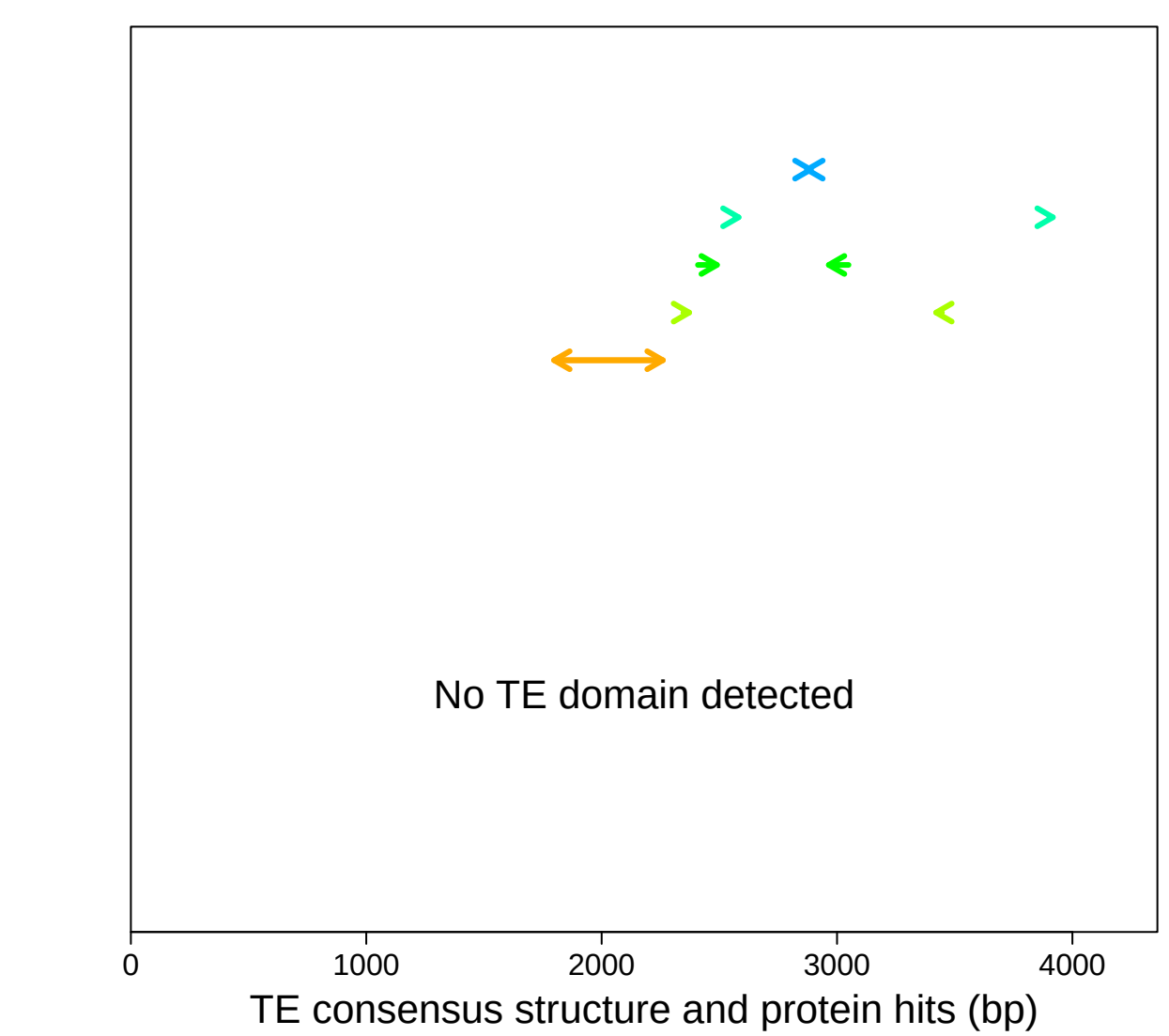
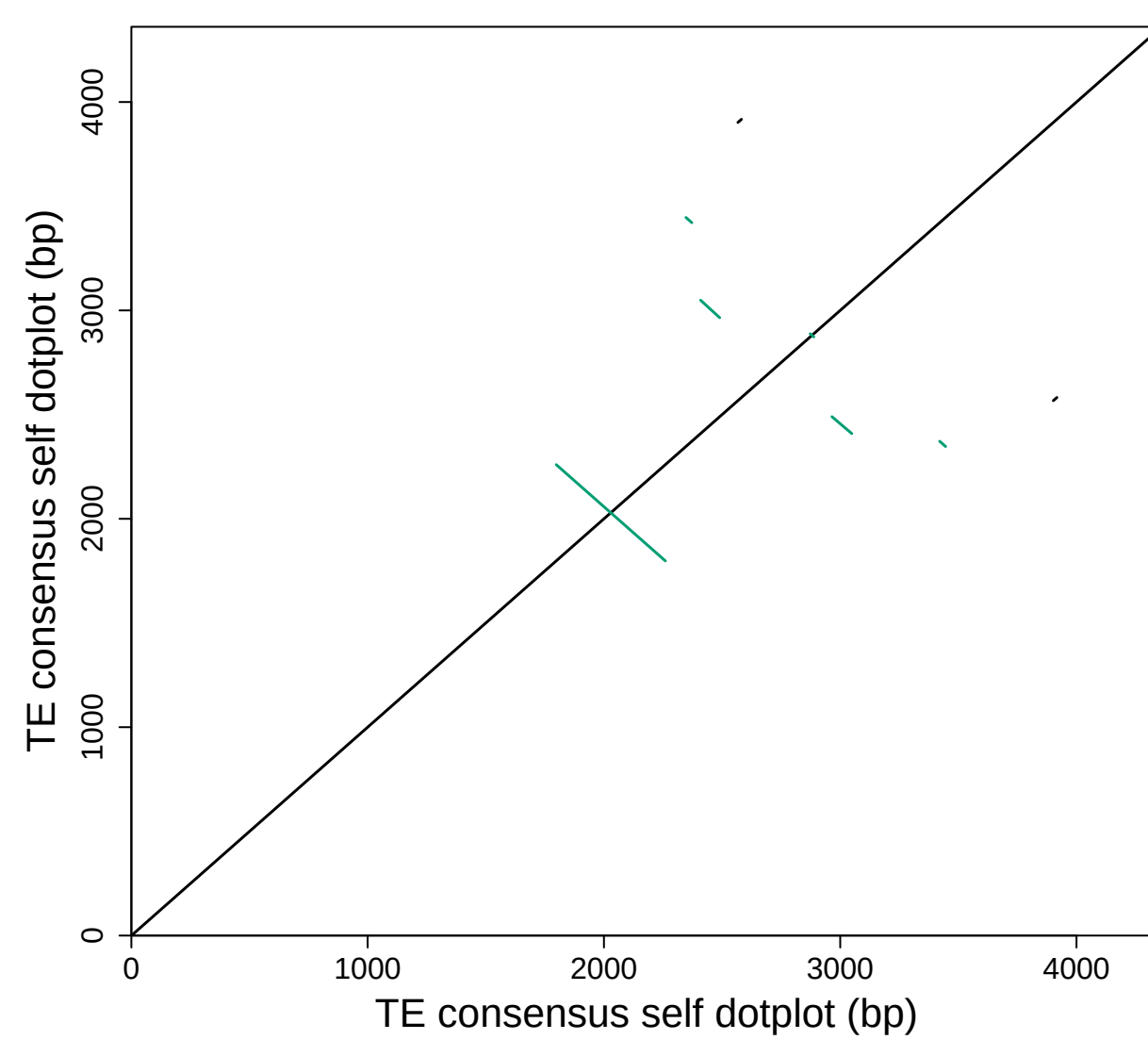
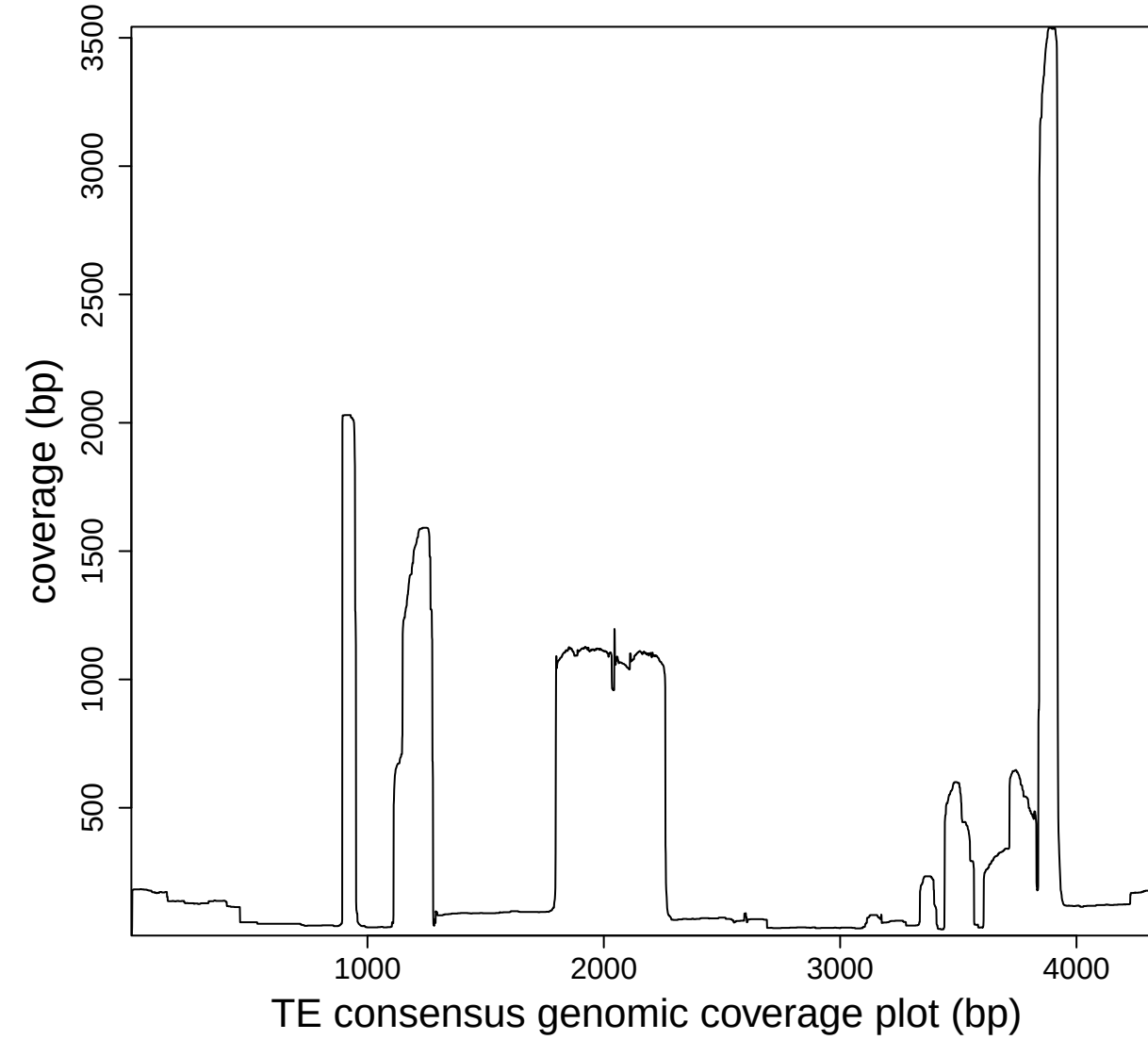
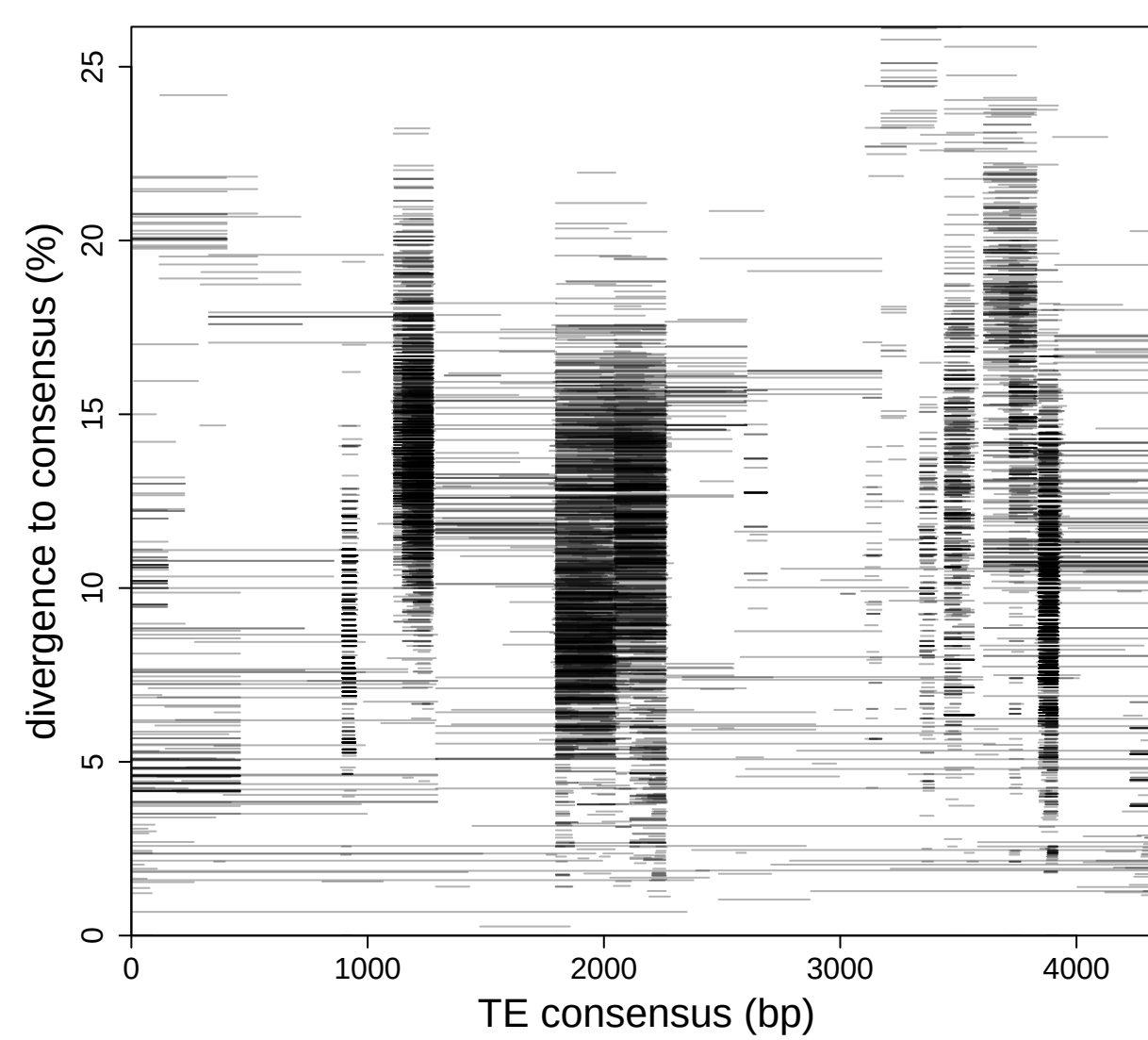
3000

4000



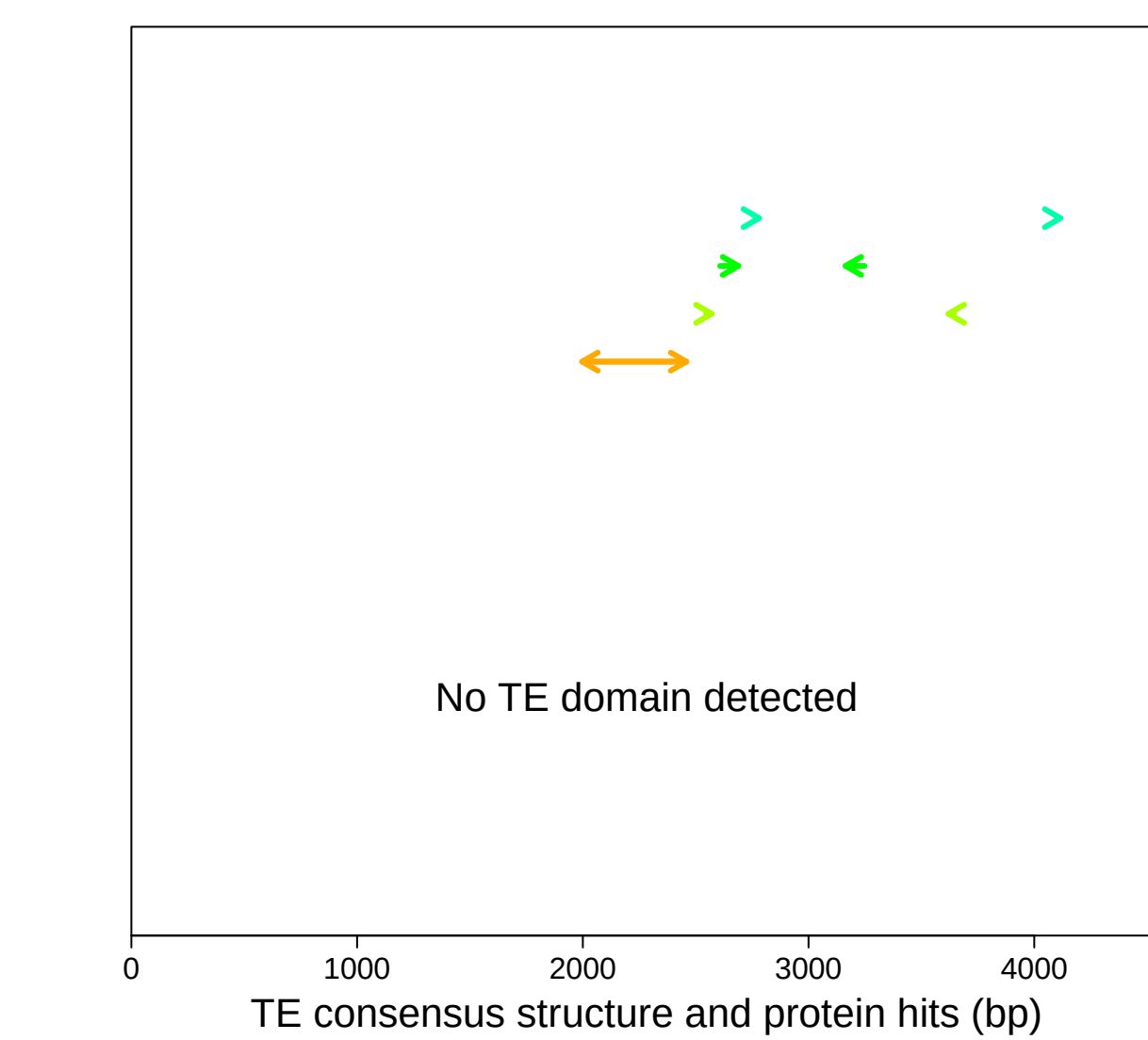
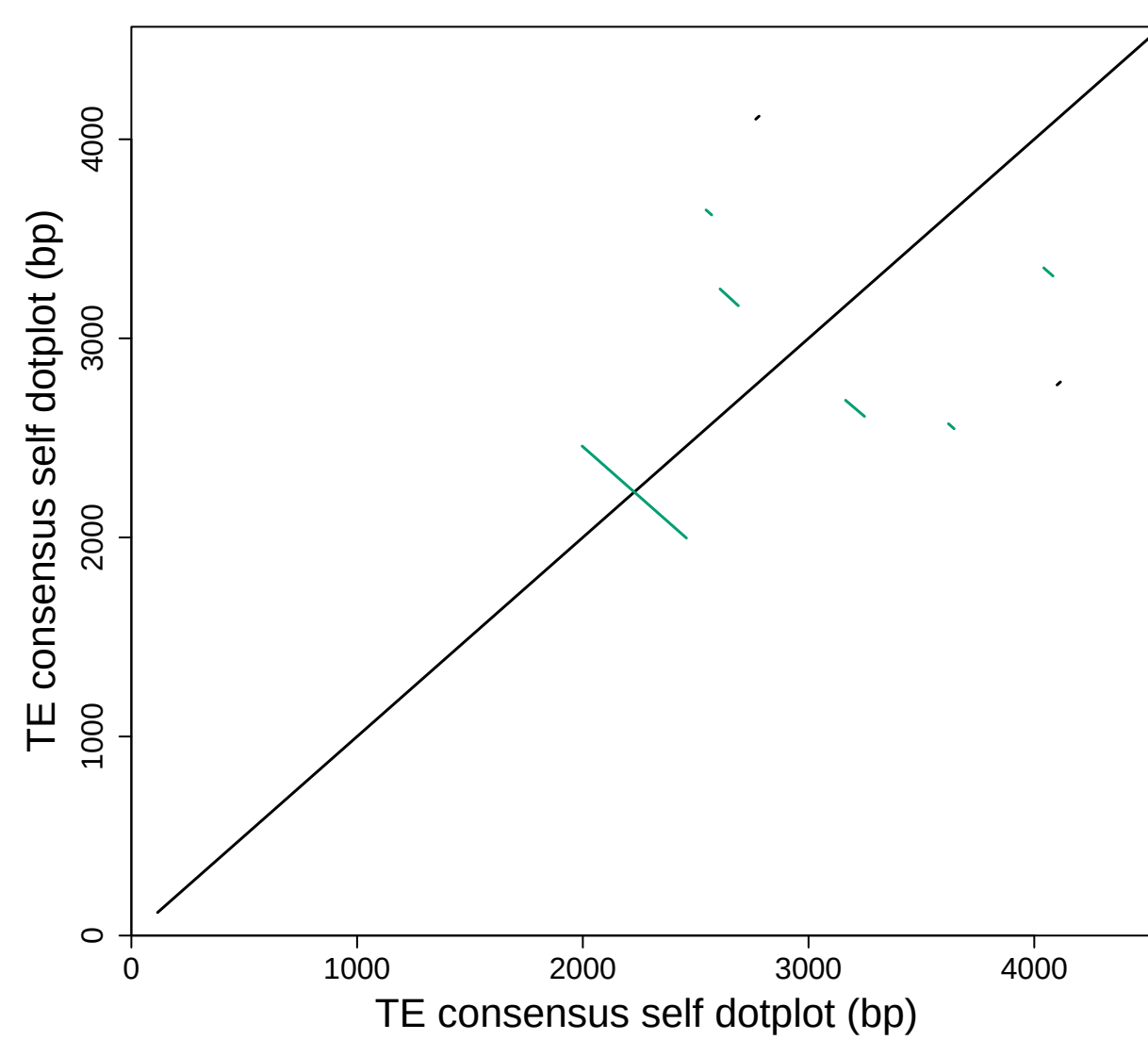
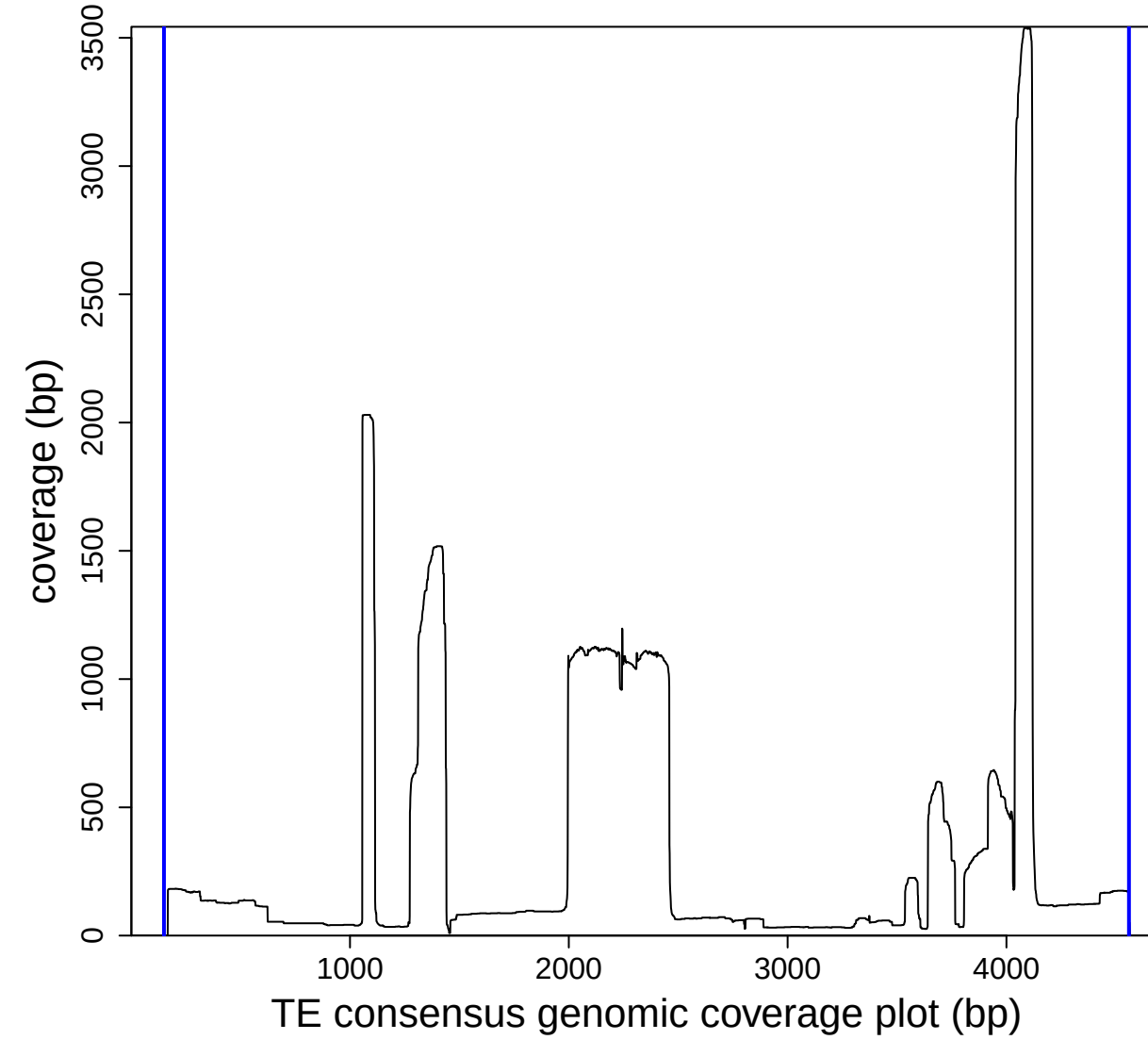
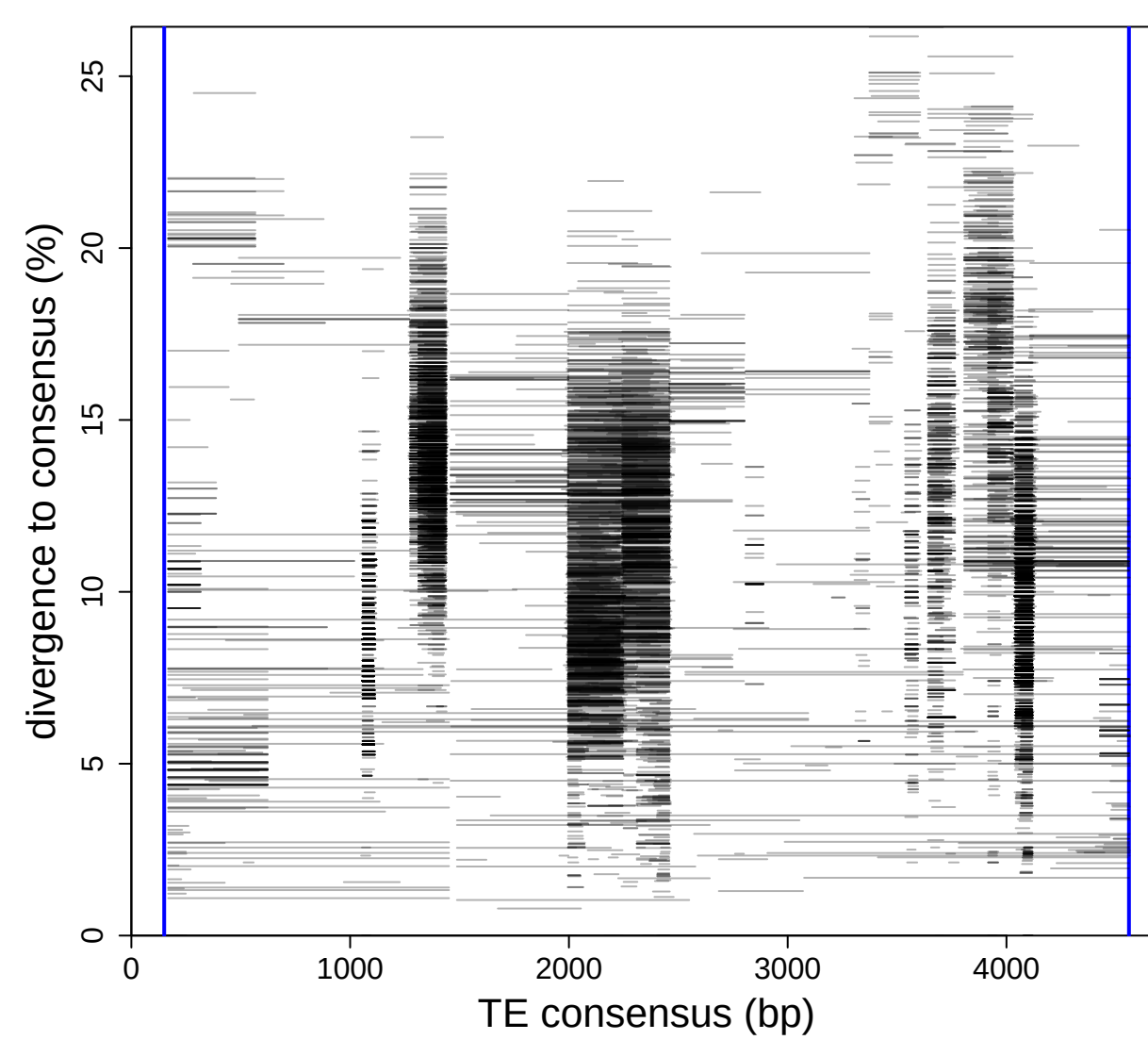
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 4361bp; fragments: 10978; full length: 0 (>=3924.9bp)

After TEtrimmer 4361 bp

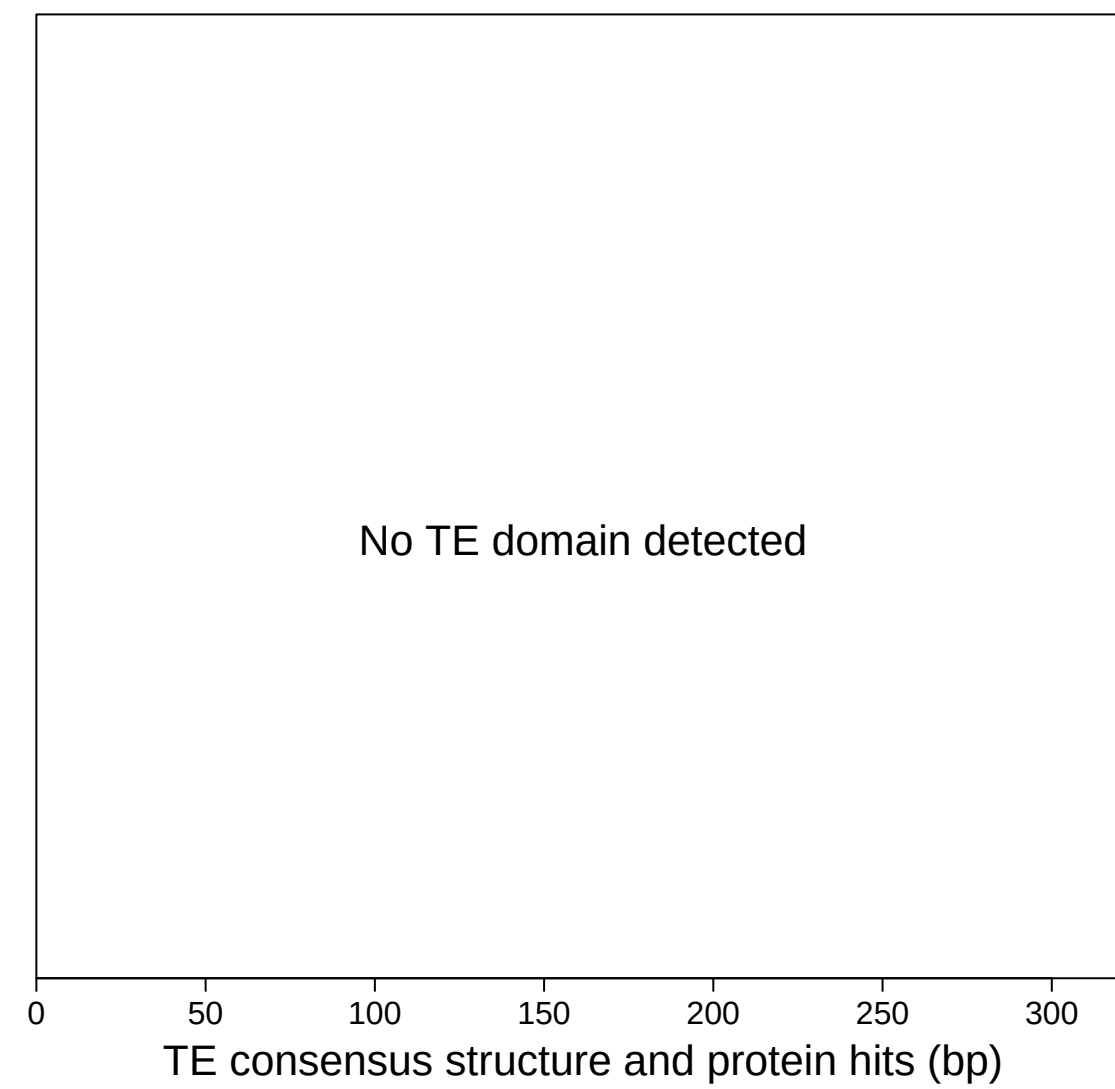
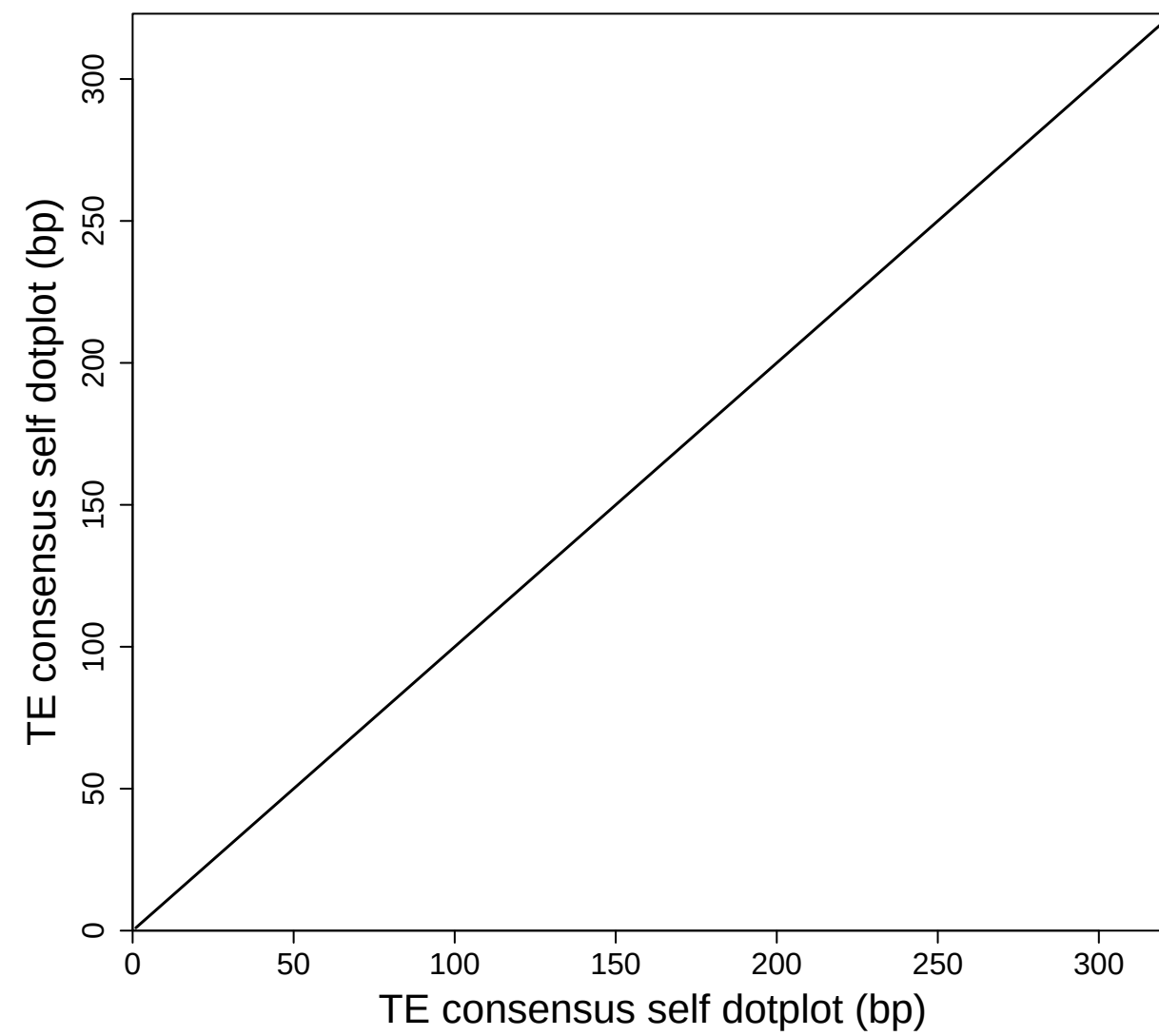
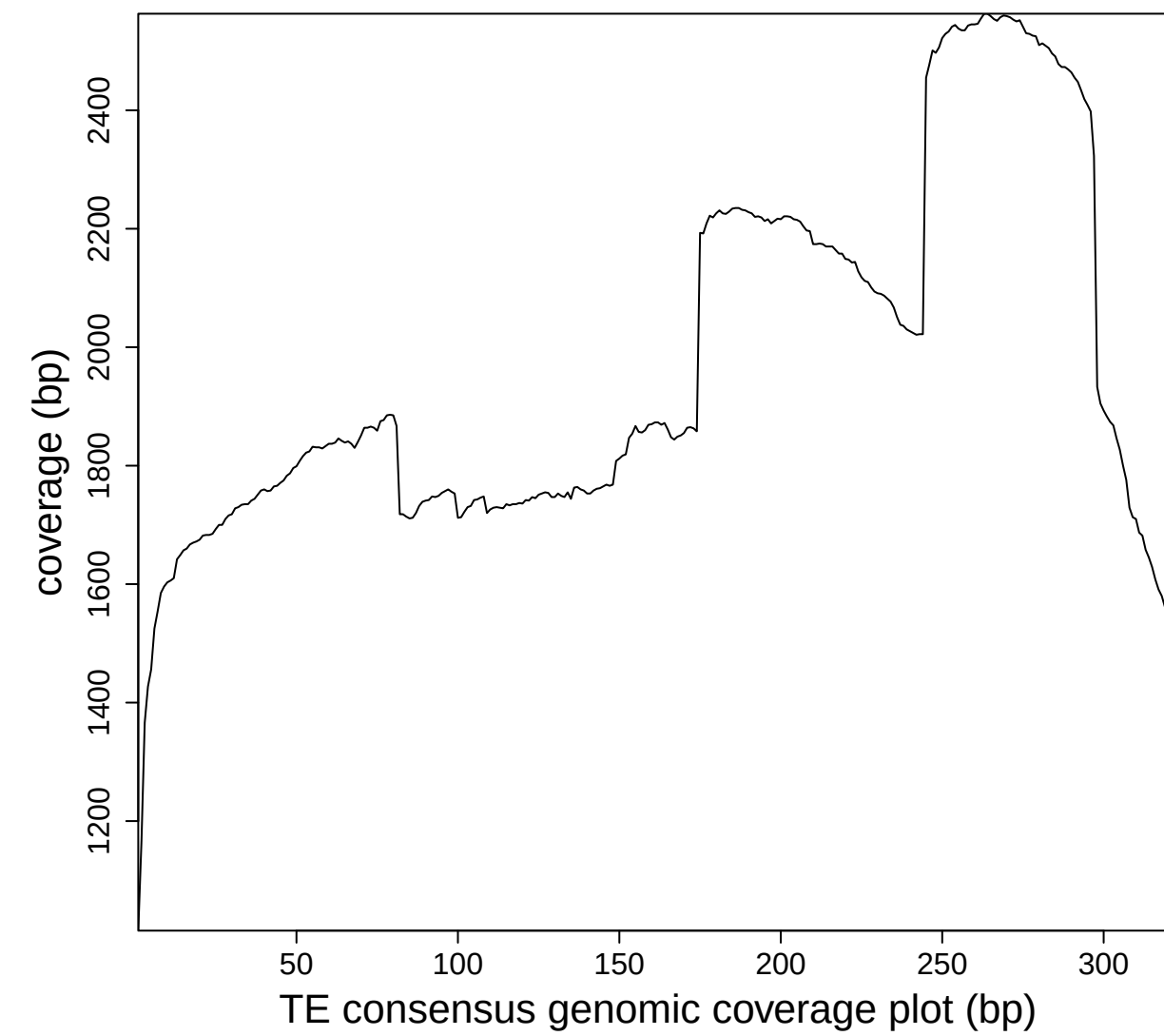
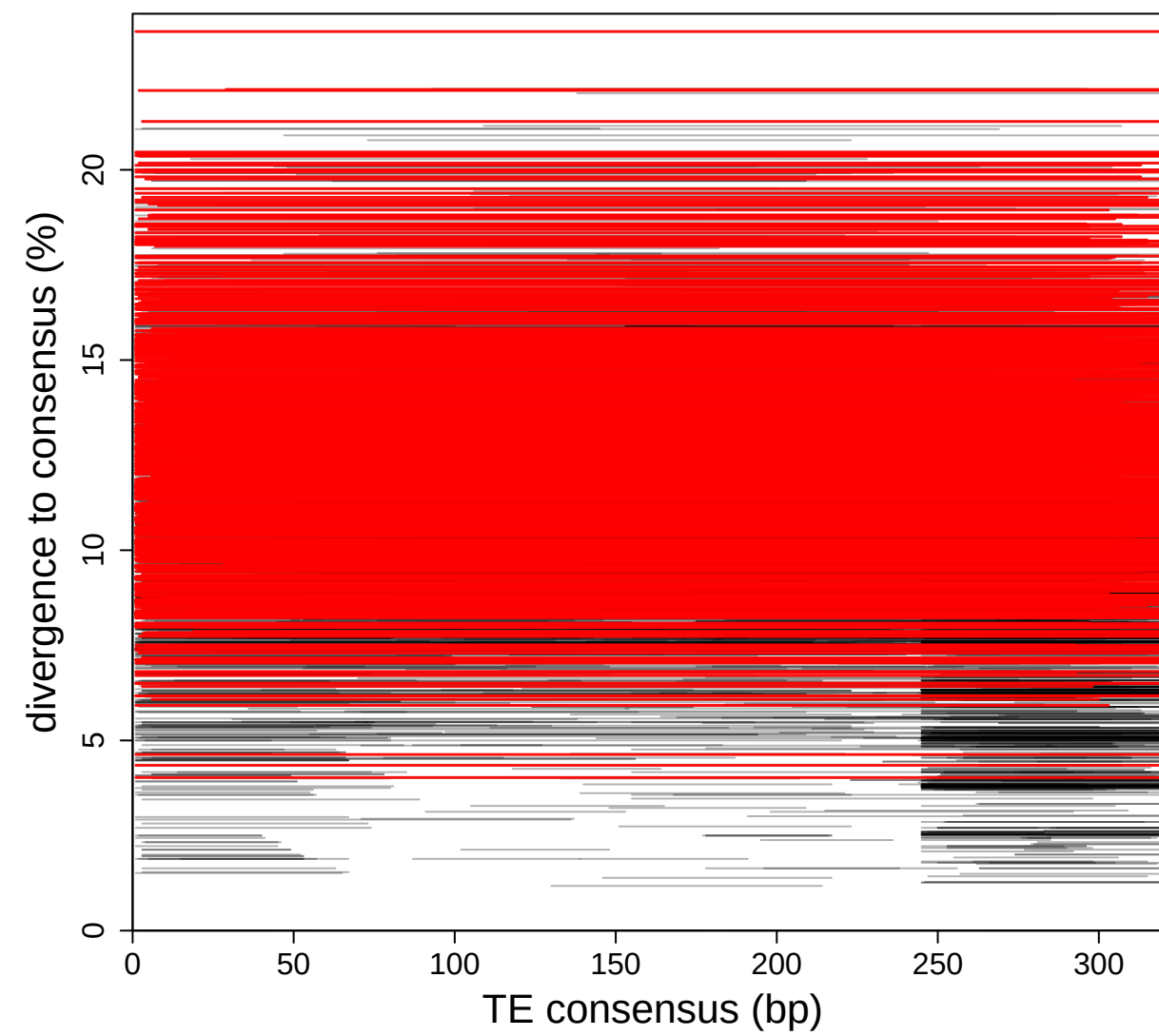


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 4710bp; fragments: 10875; full length: 0 (>=4239bp)

After TEtrimmer Extended plot Blue lines are boundaries

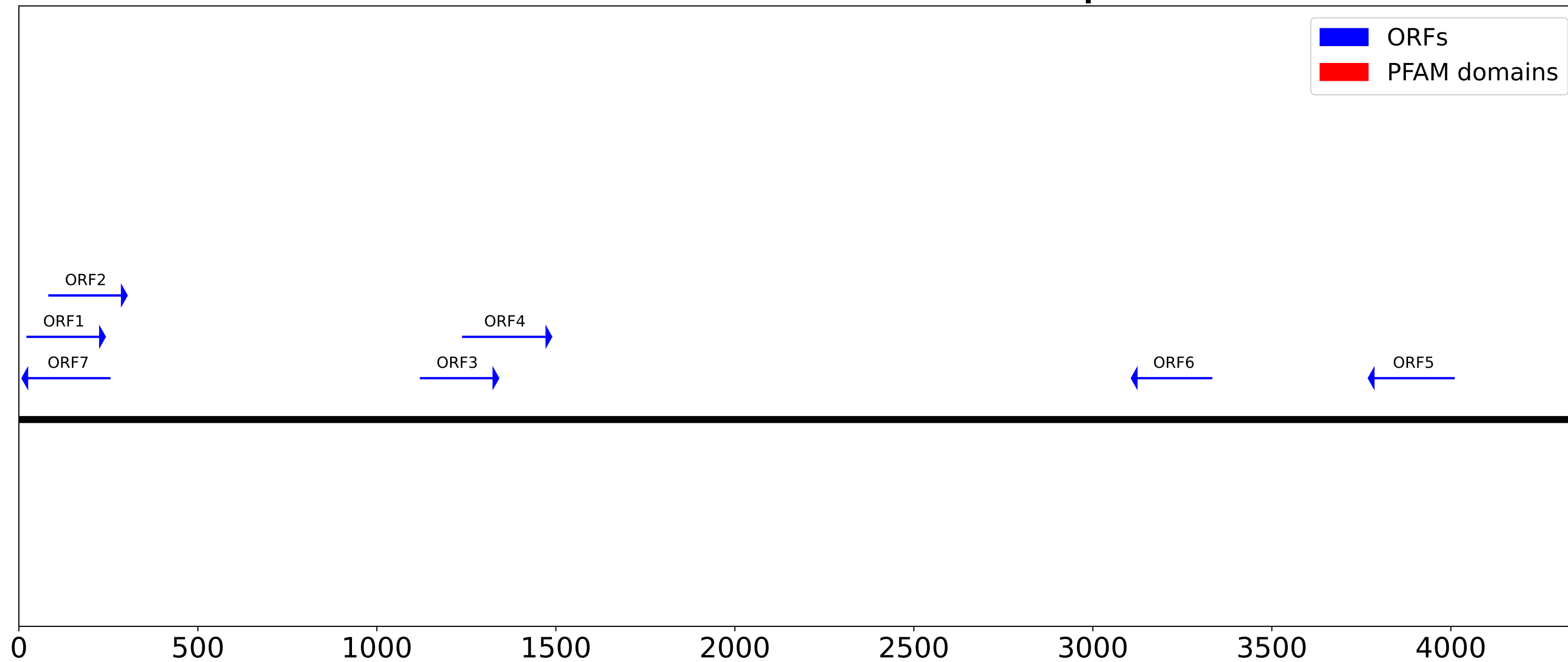


TE: rnd_1_family_706
size: 323bp; fragments: 3763; full length: 713 (≥ 290.7 bp)



Before TEtrimmer 323 bp

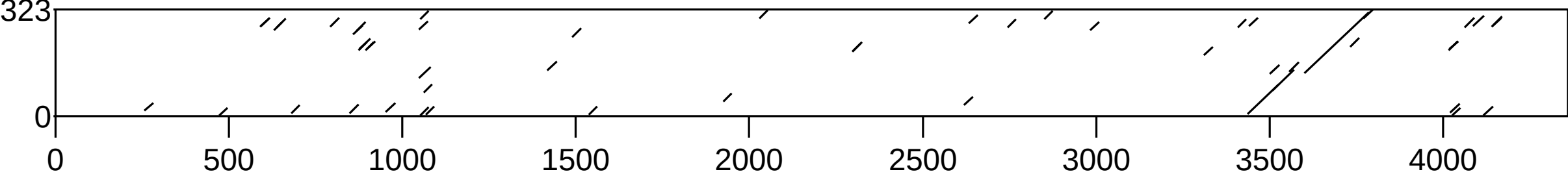
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)