

Start crop Point

End crop Point

MSA length = 1867



Start crop Point

End crop Point

0

10

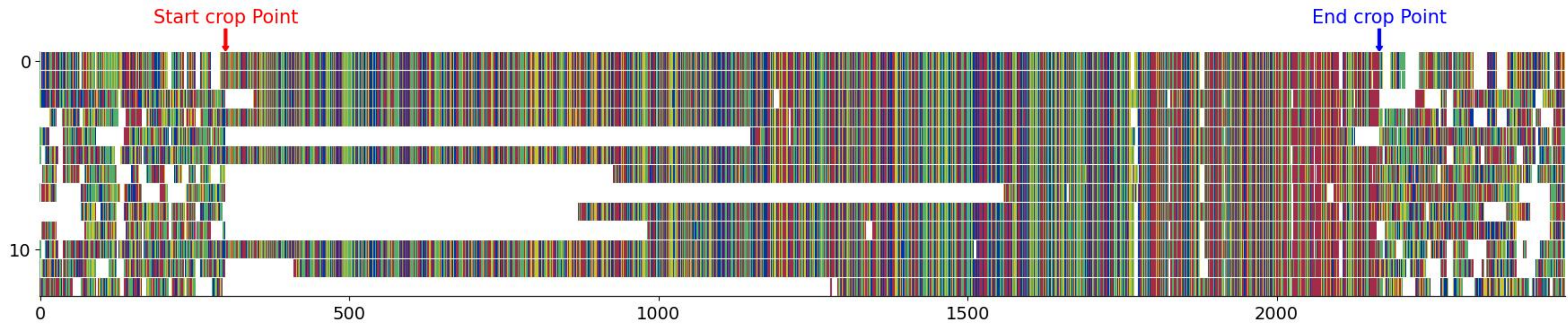
0

500

1000

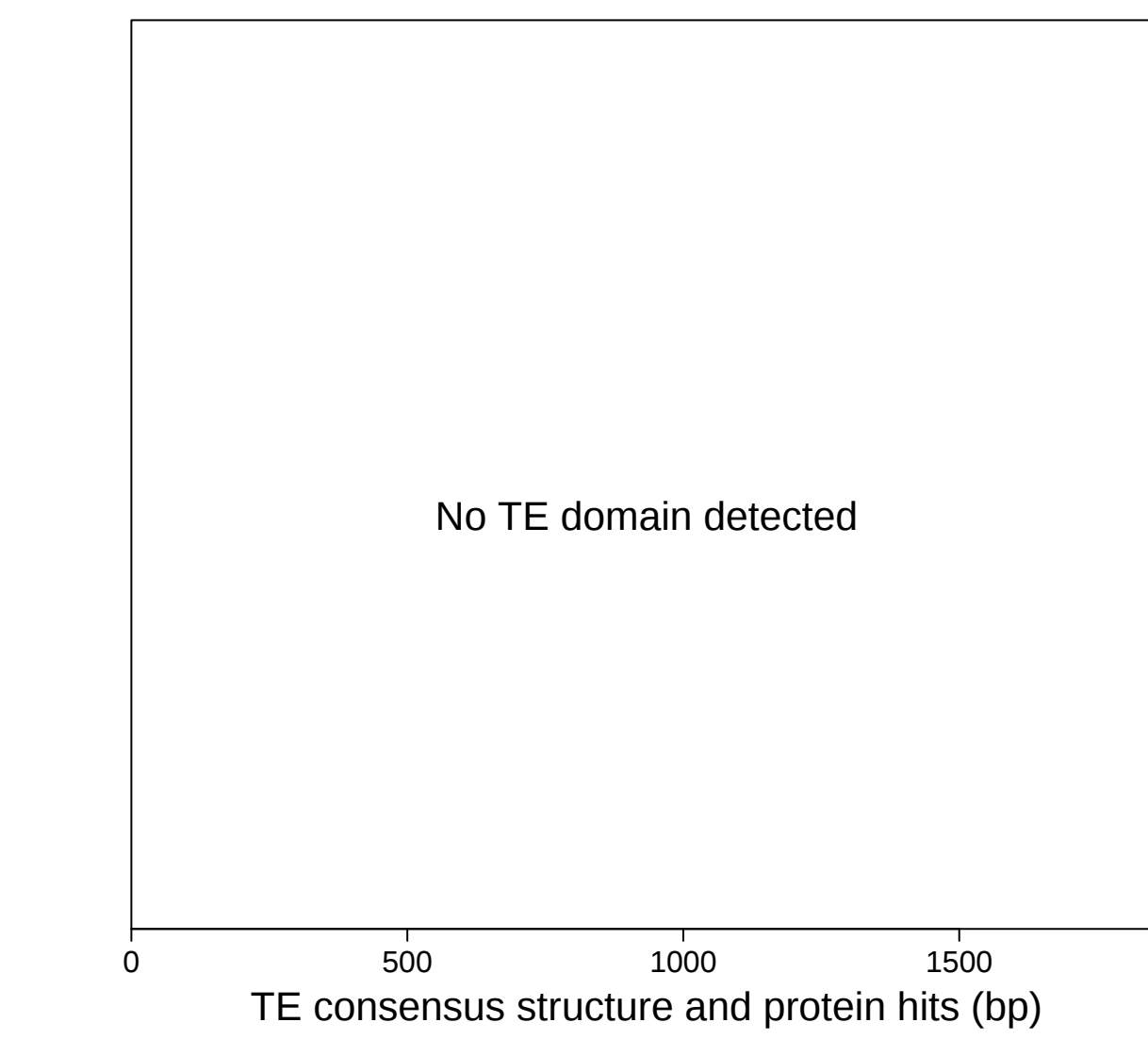
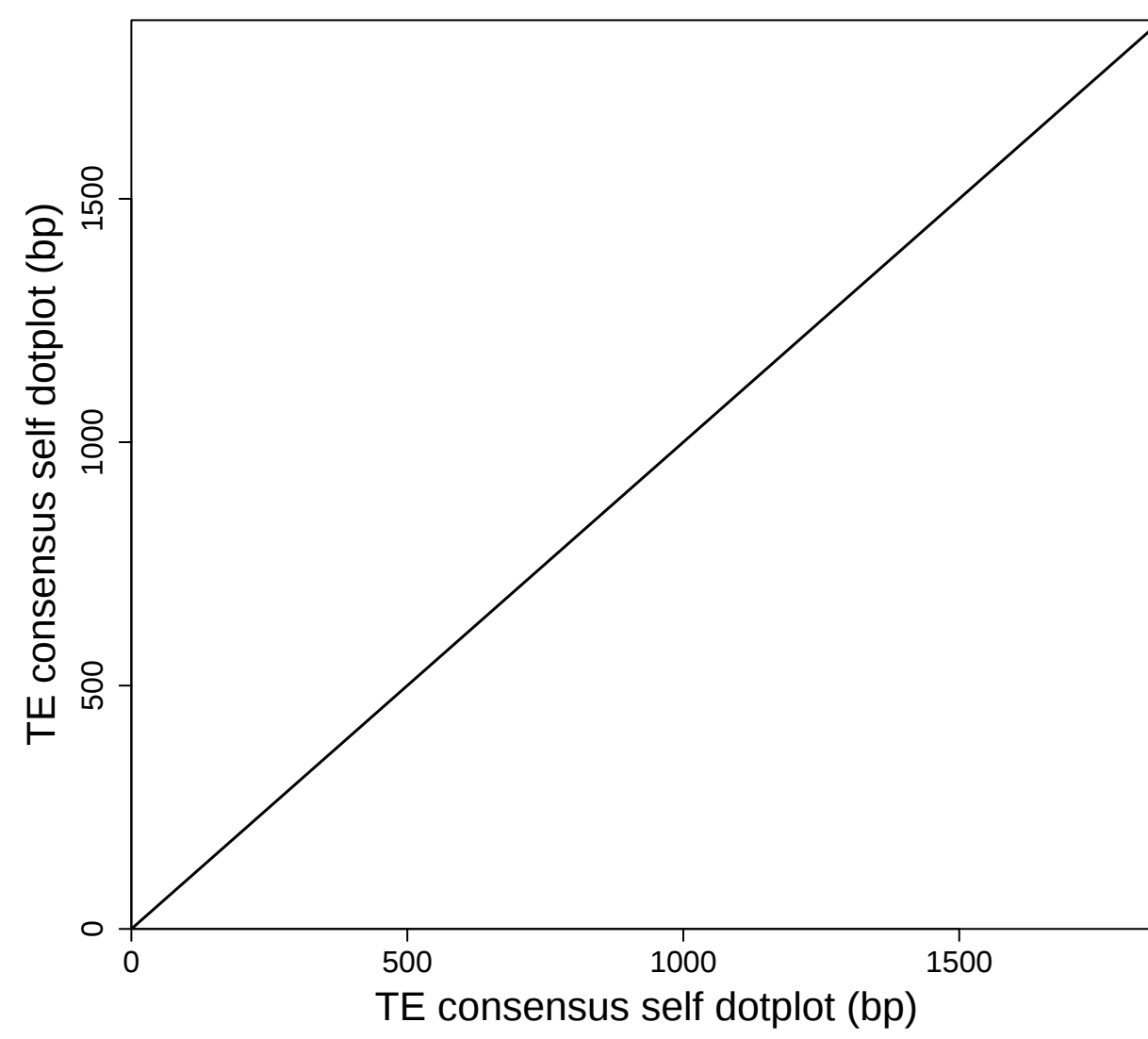
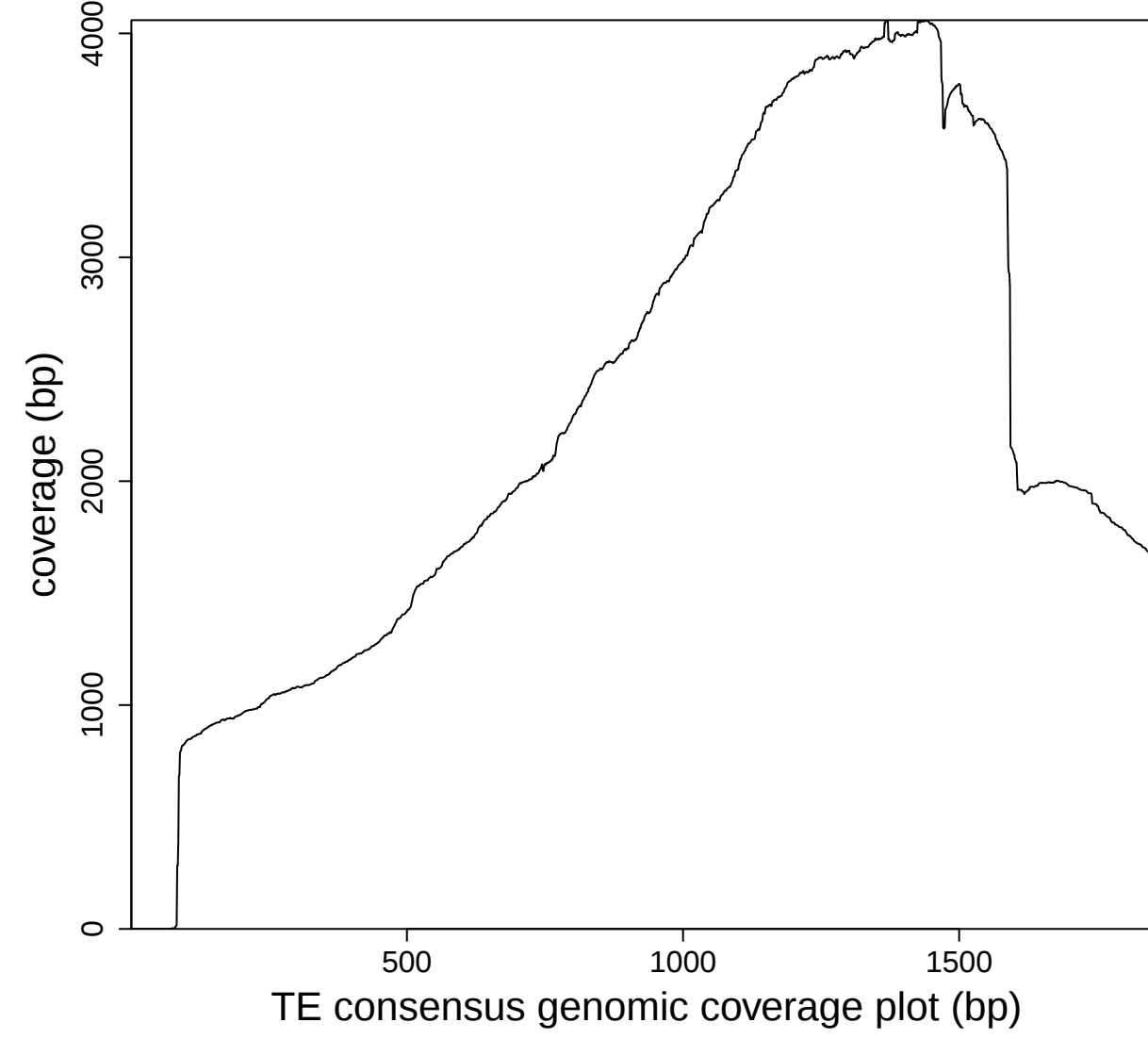
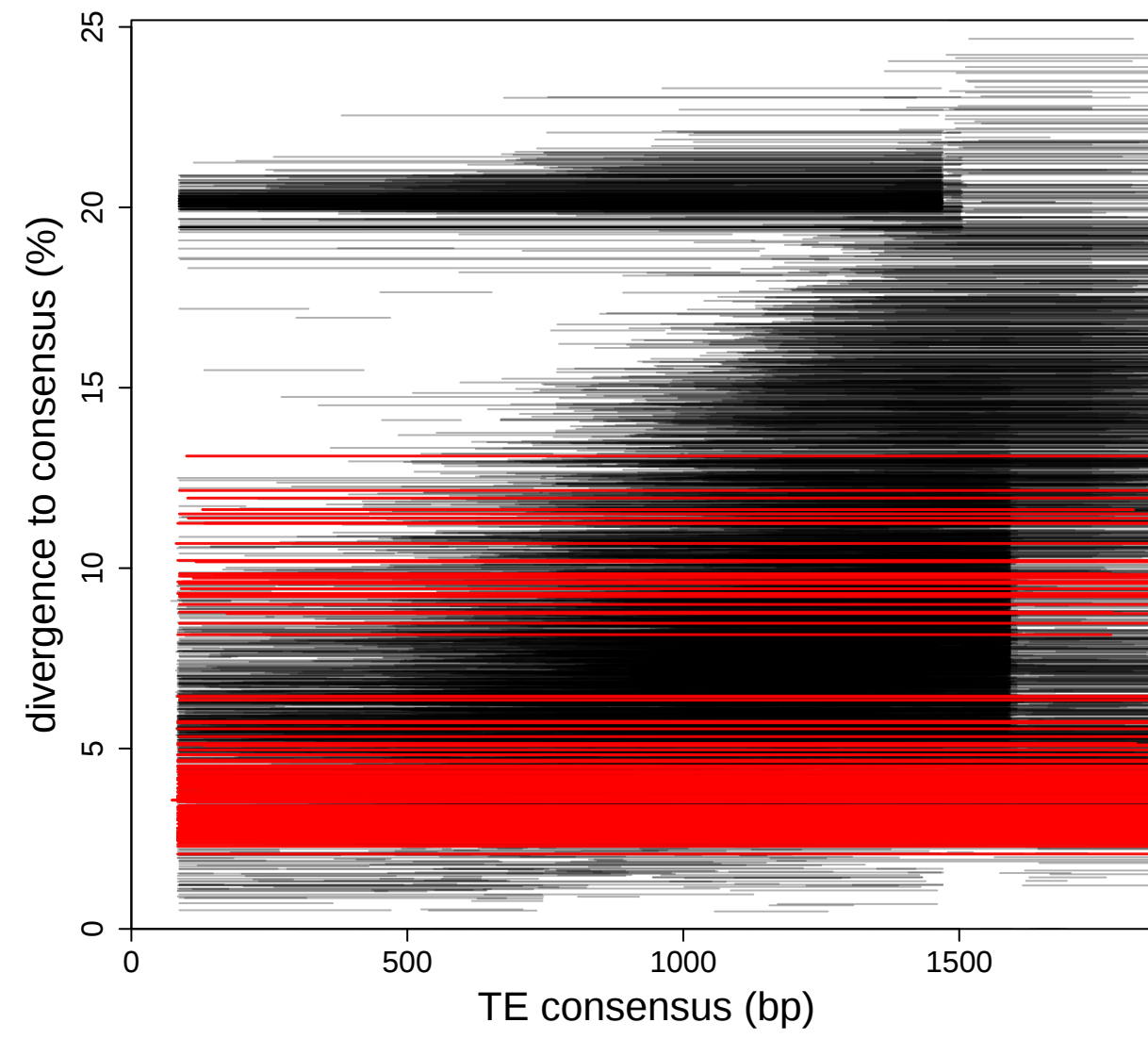
1500

2000



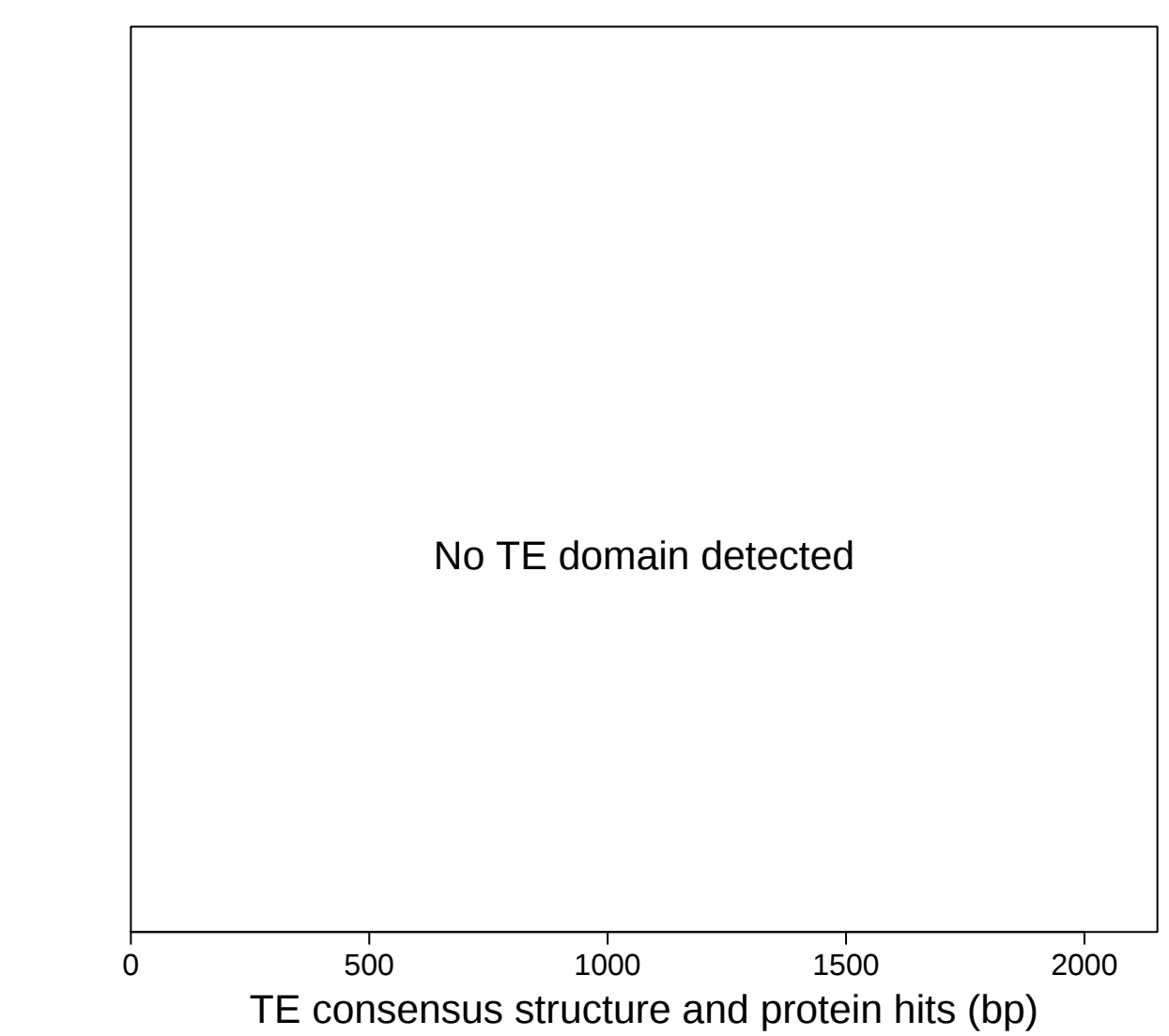
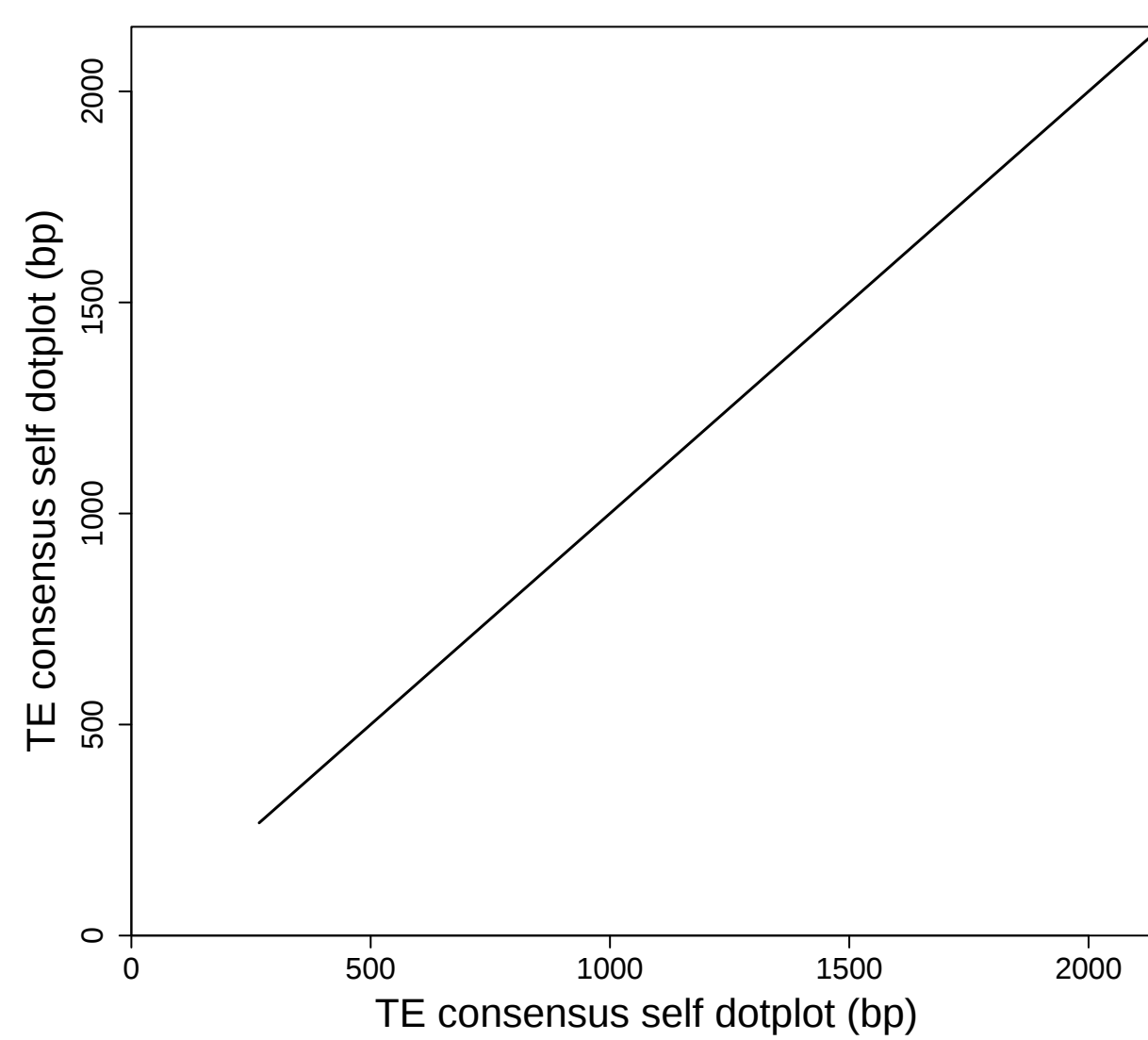
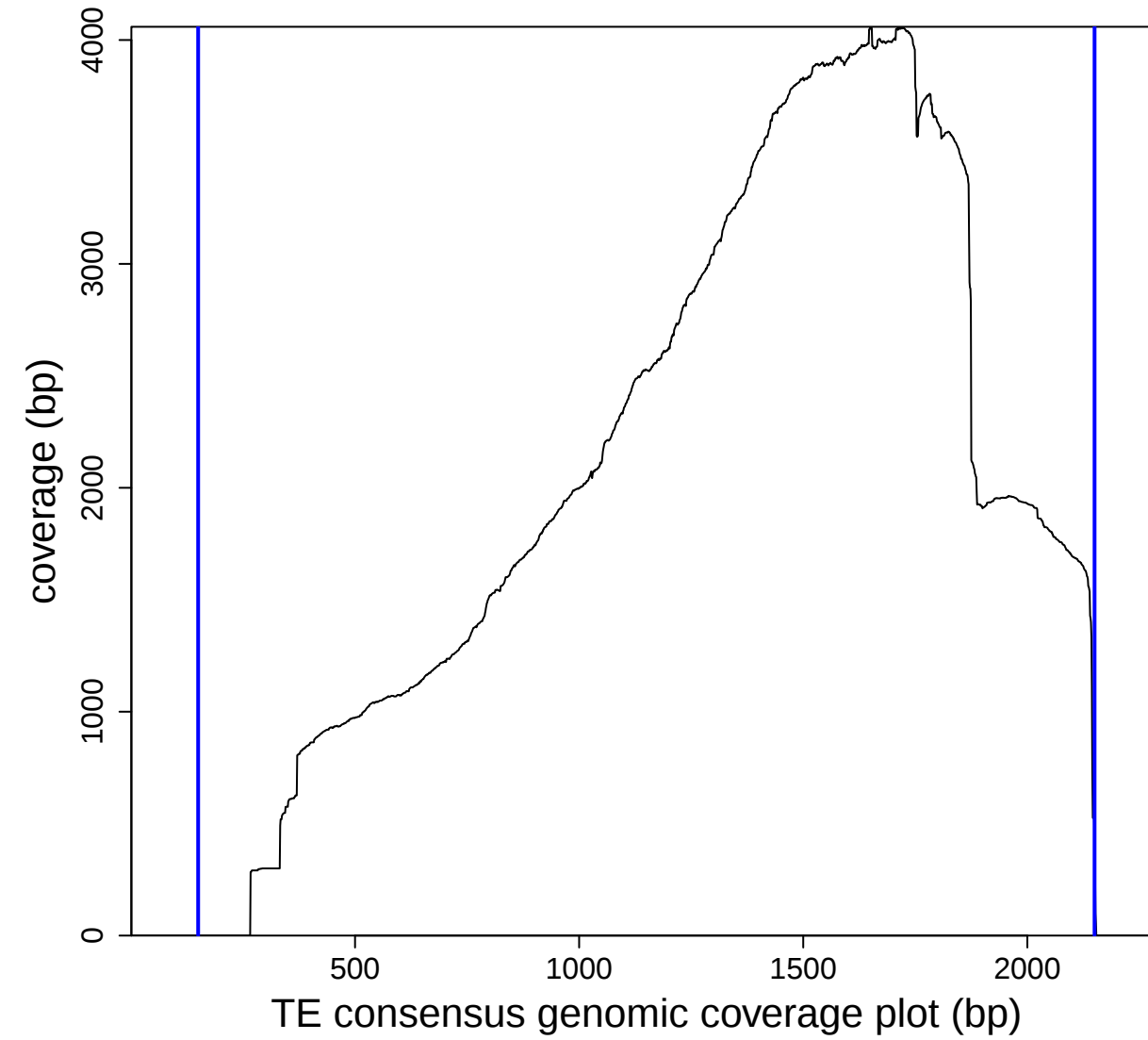
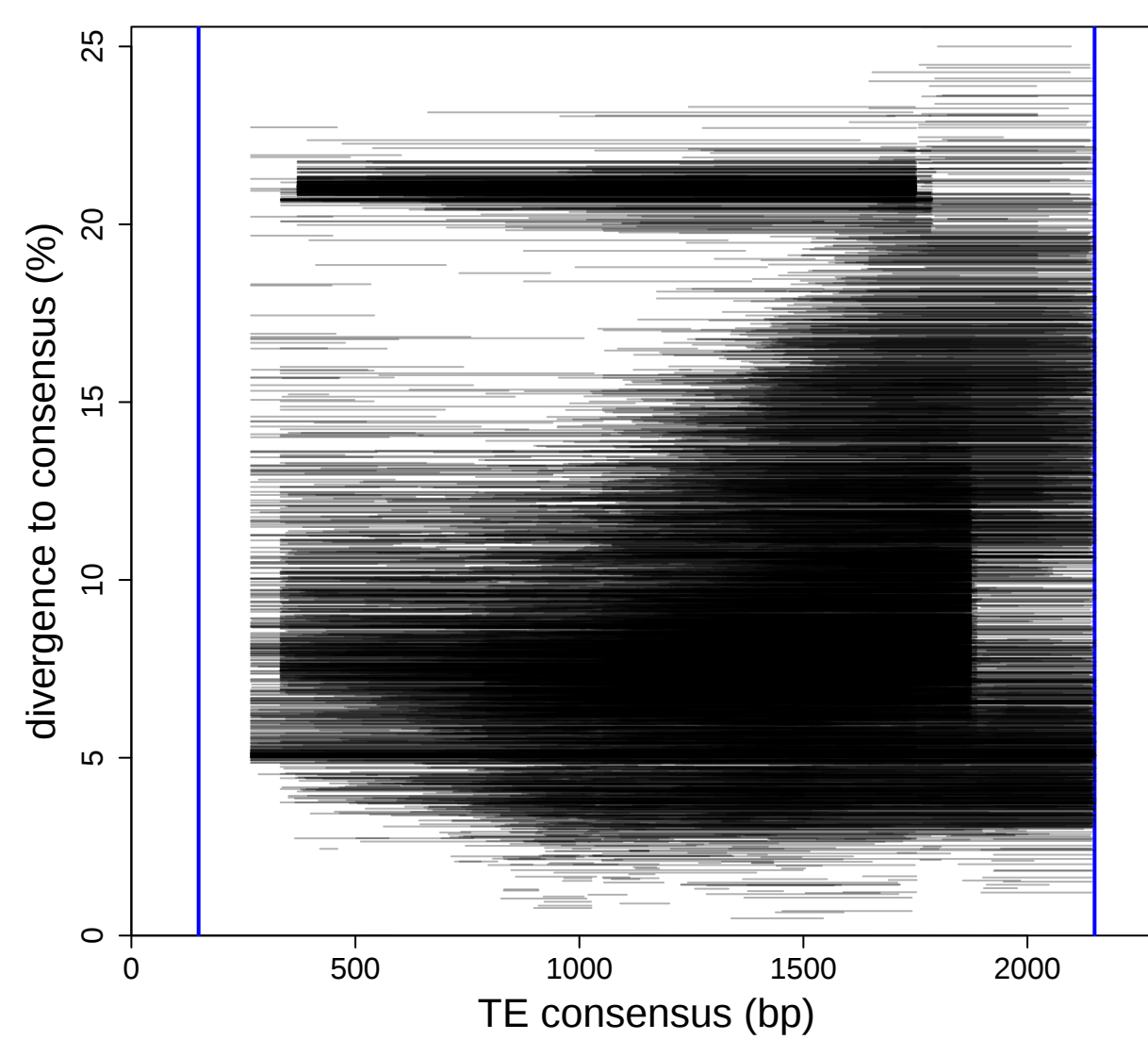
ed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.fa
size: 1867bp; fragments: 6310; full length: 196 (≥ 1680.3 bp)

After TEtrimmer 1867 bp

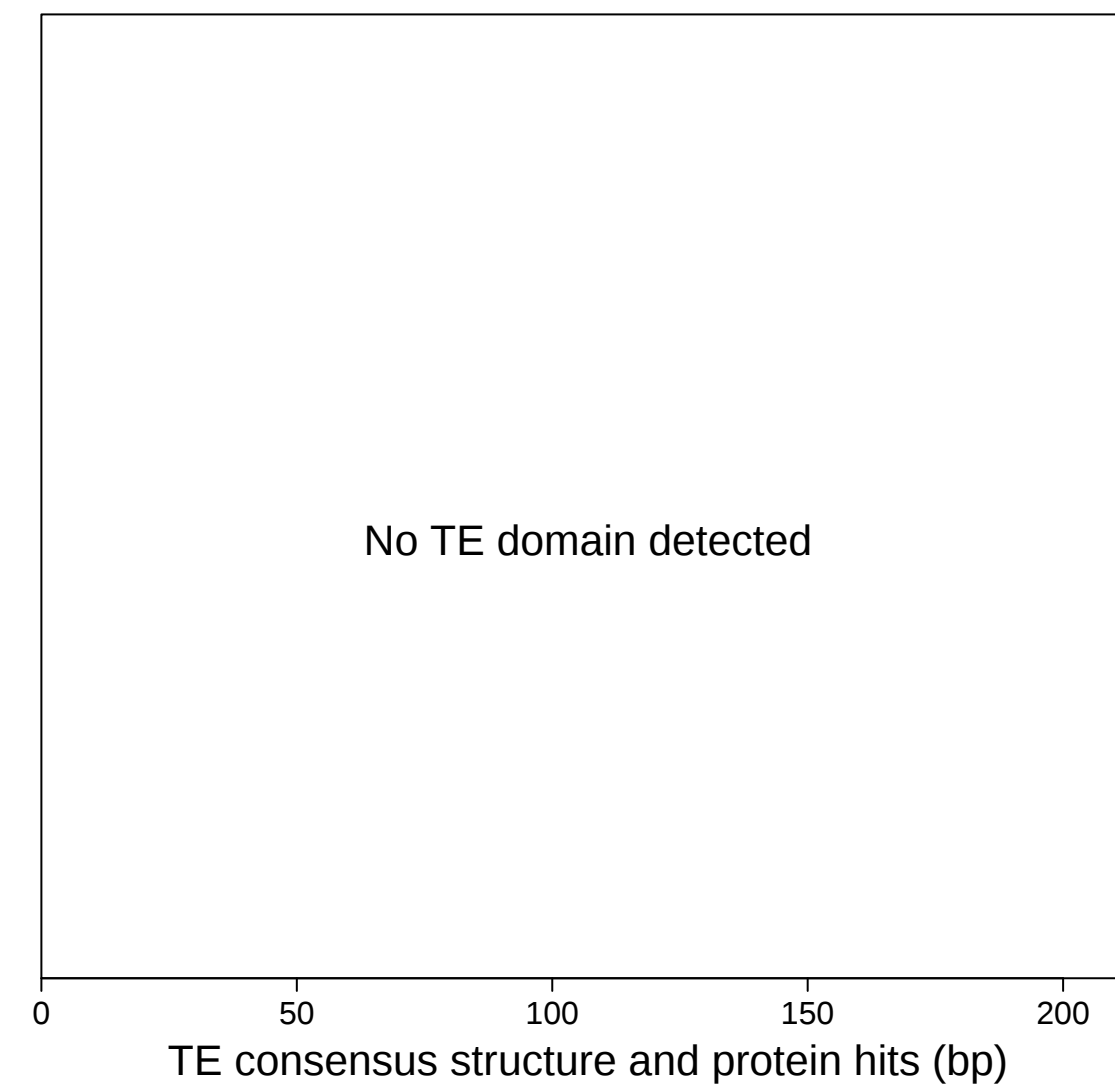
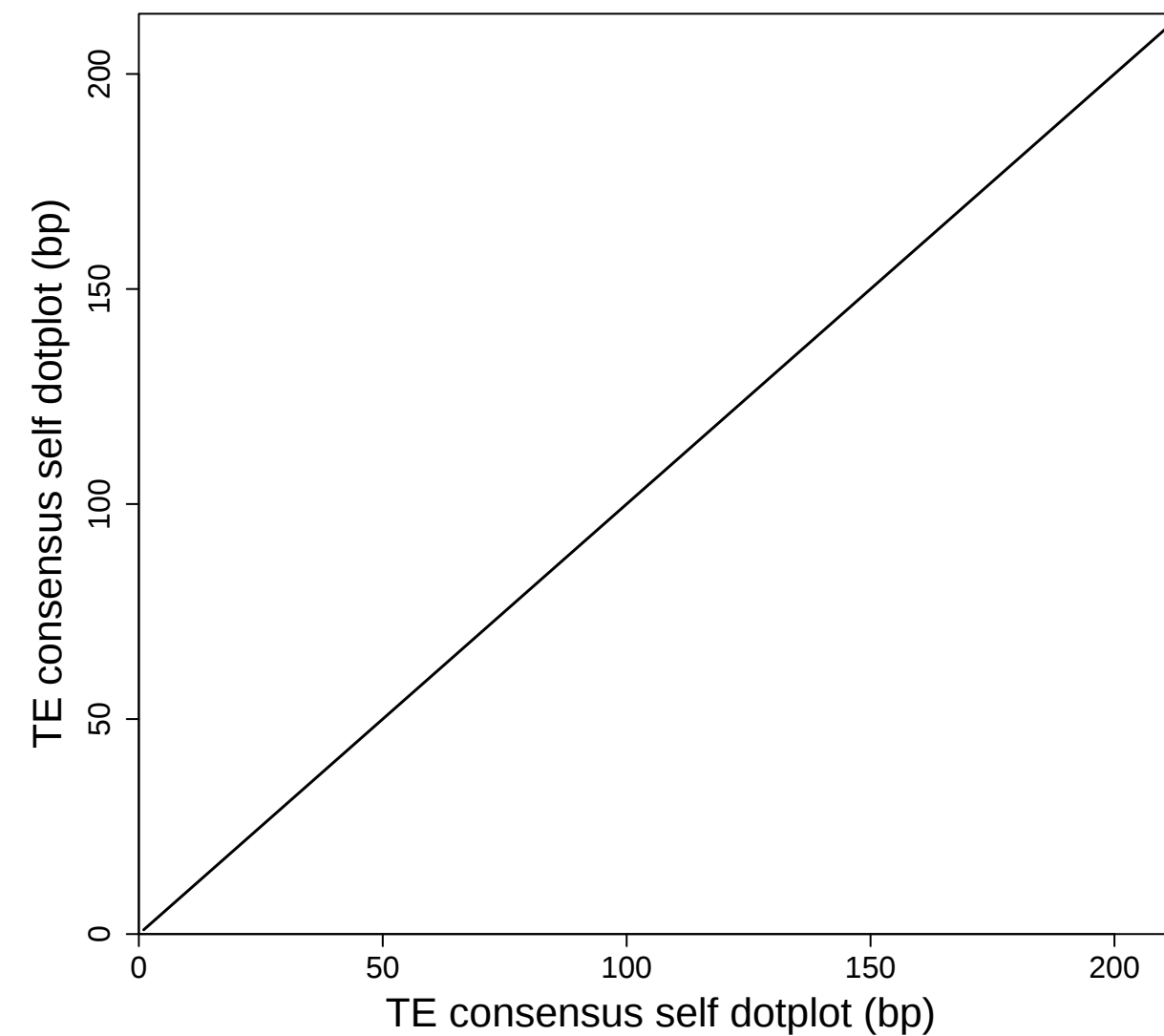
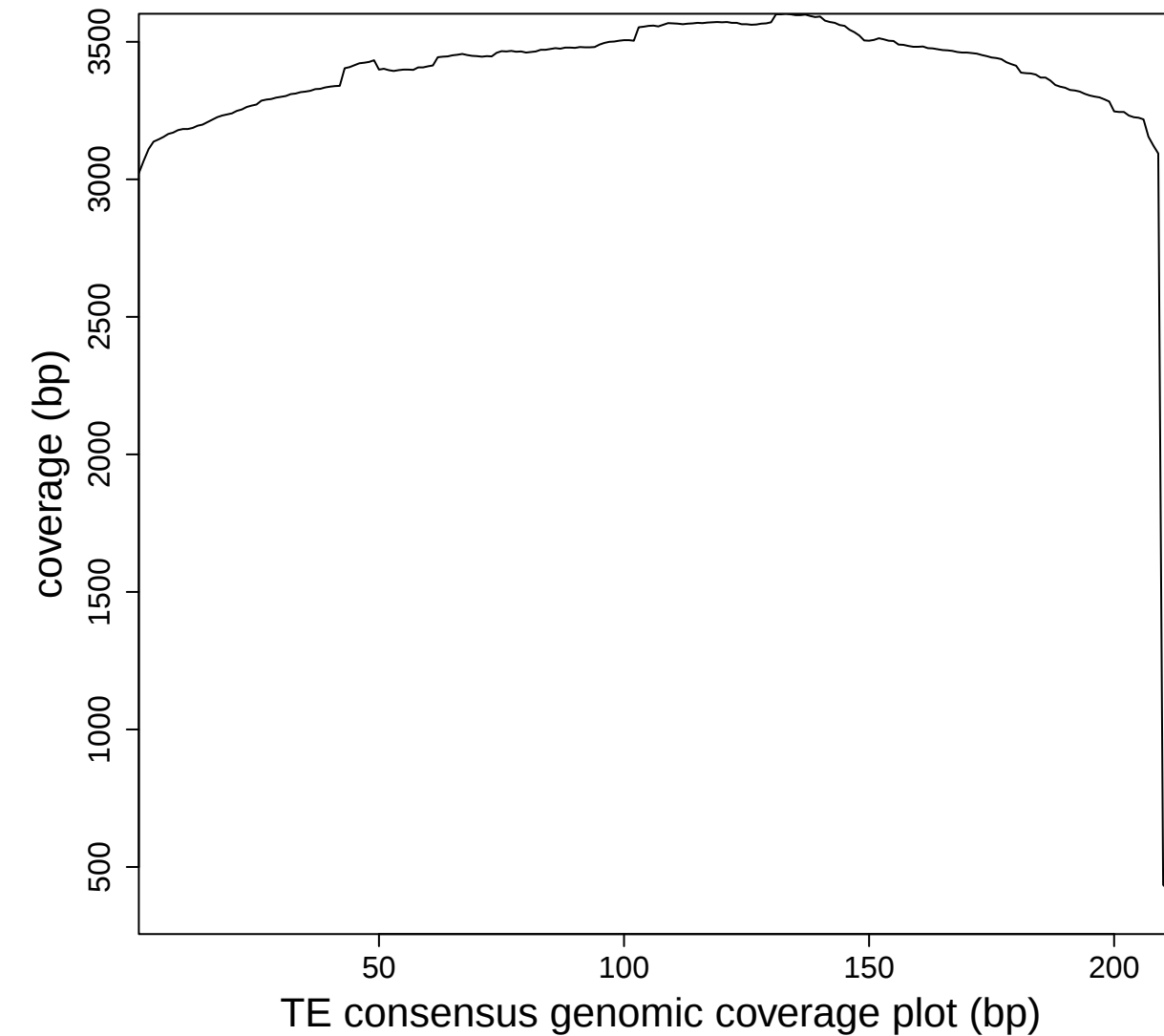
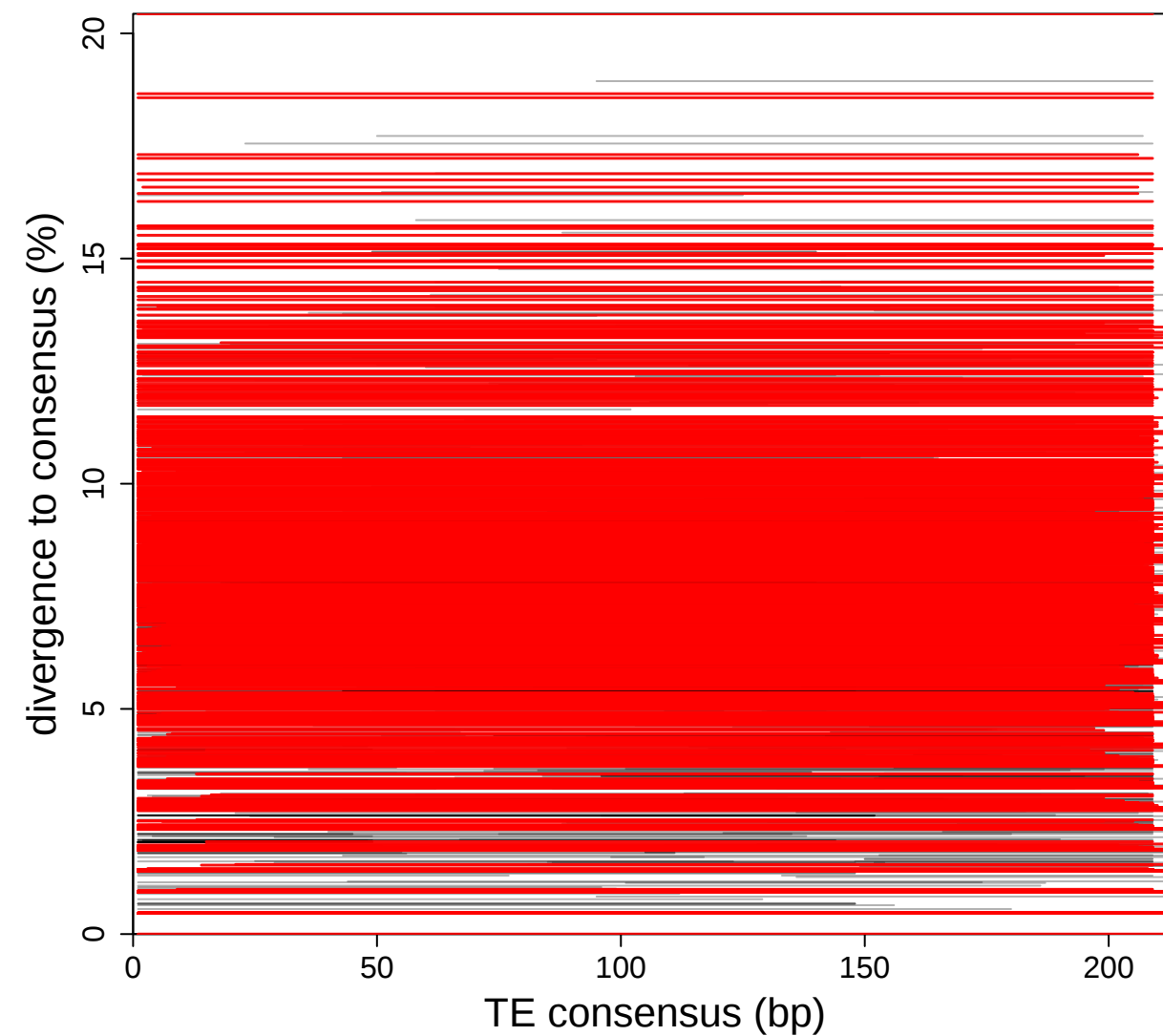


ed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.
size: 2300bp; fragments: 6224; full length: 0 (>=2070bp)

After TEtrimmer Extended plot Blue lines are boundaries

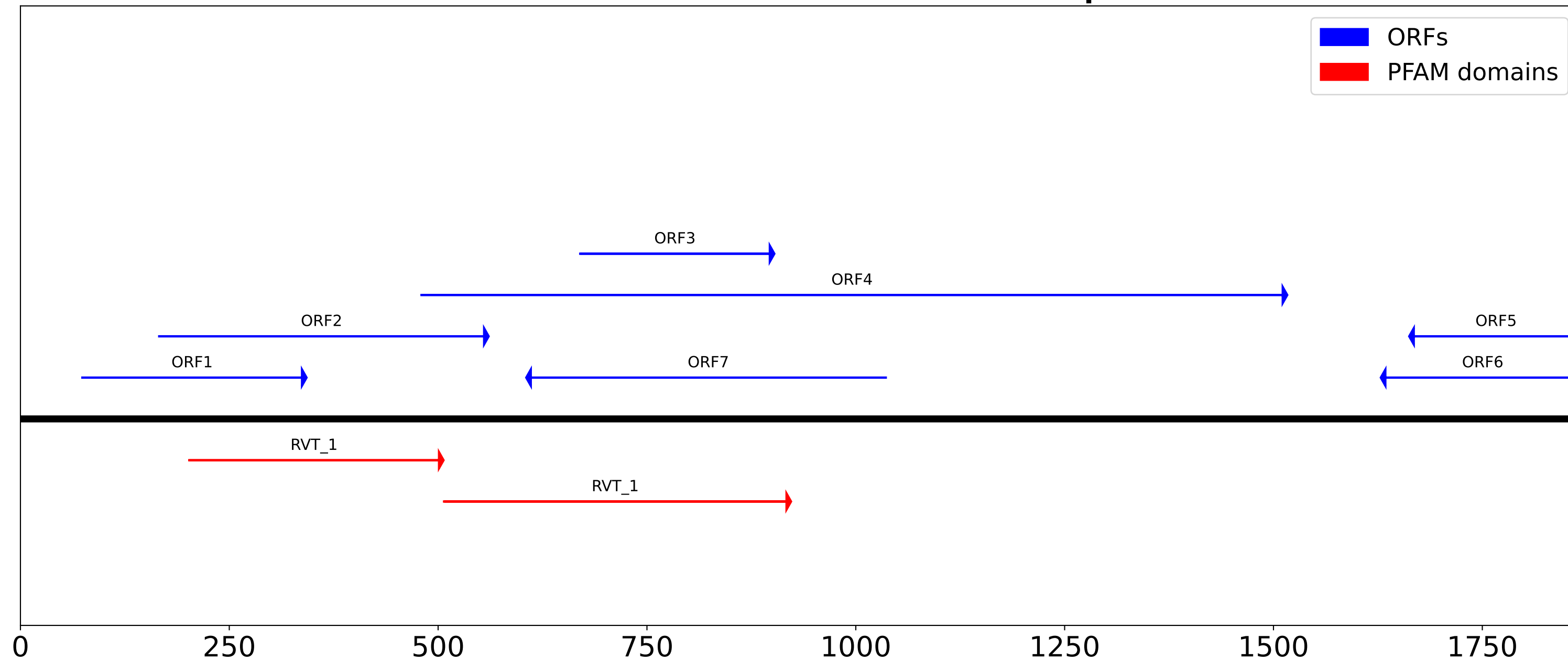


TE: ltr_1_family_5
size: 214bp; fragments: 3892; full length: 2714 (≥ 192.6 bp)



Before TEtrimmer 214 bp

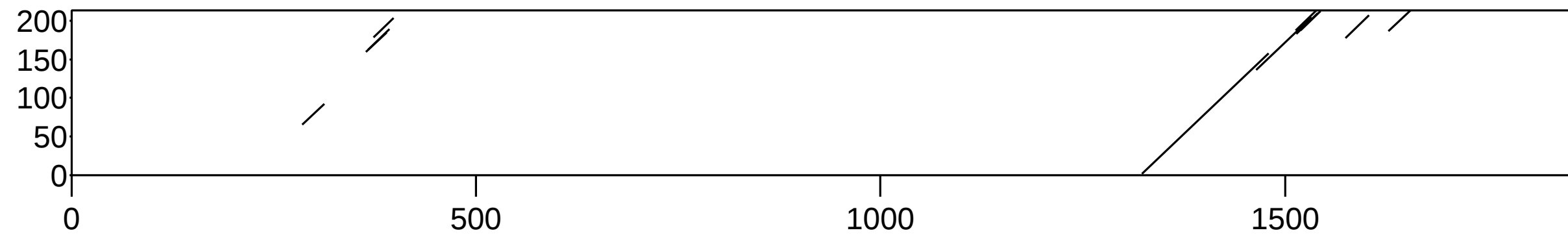
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

sensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)