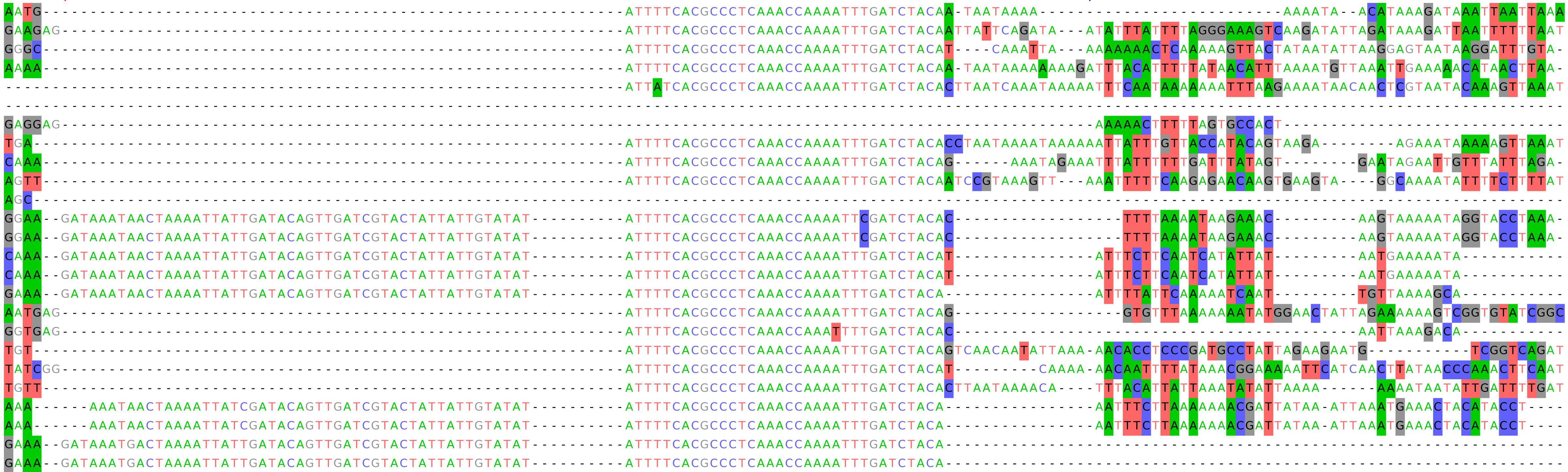
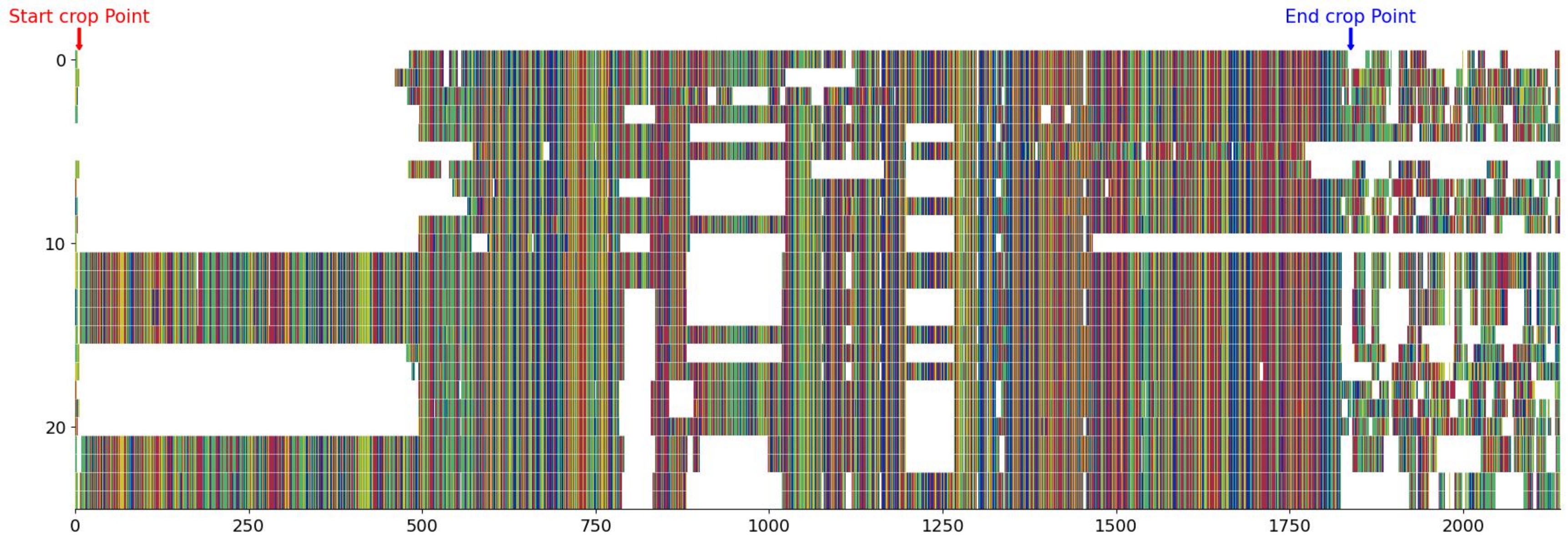


Start crop Point

End crop Point

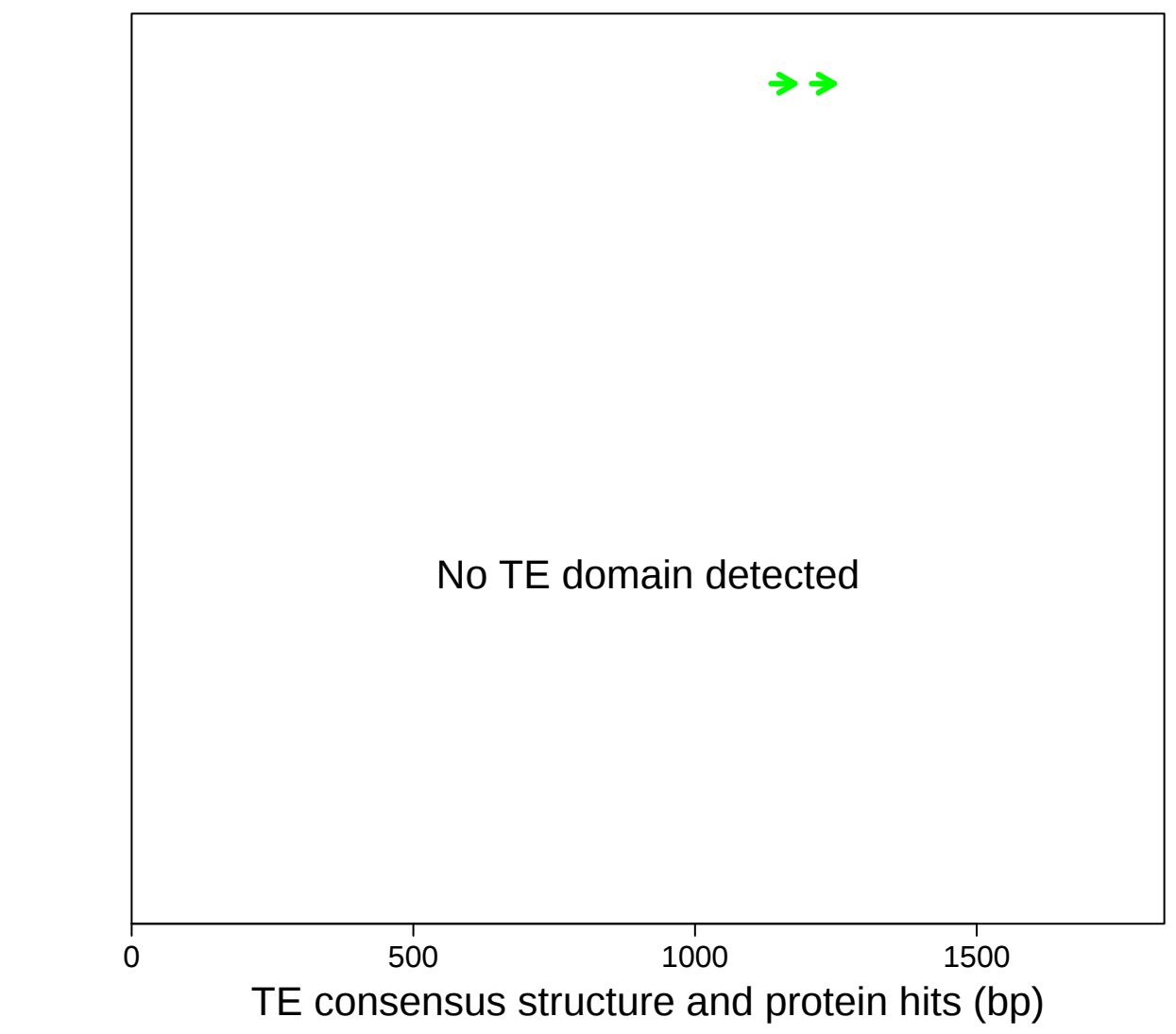
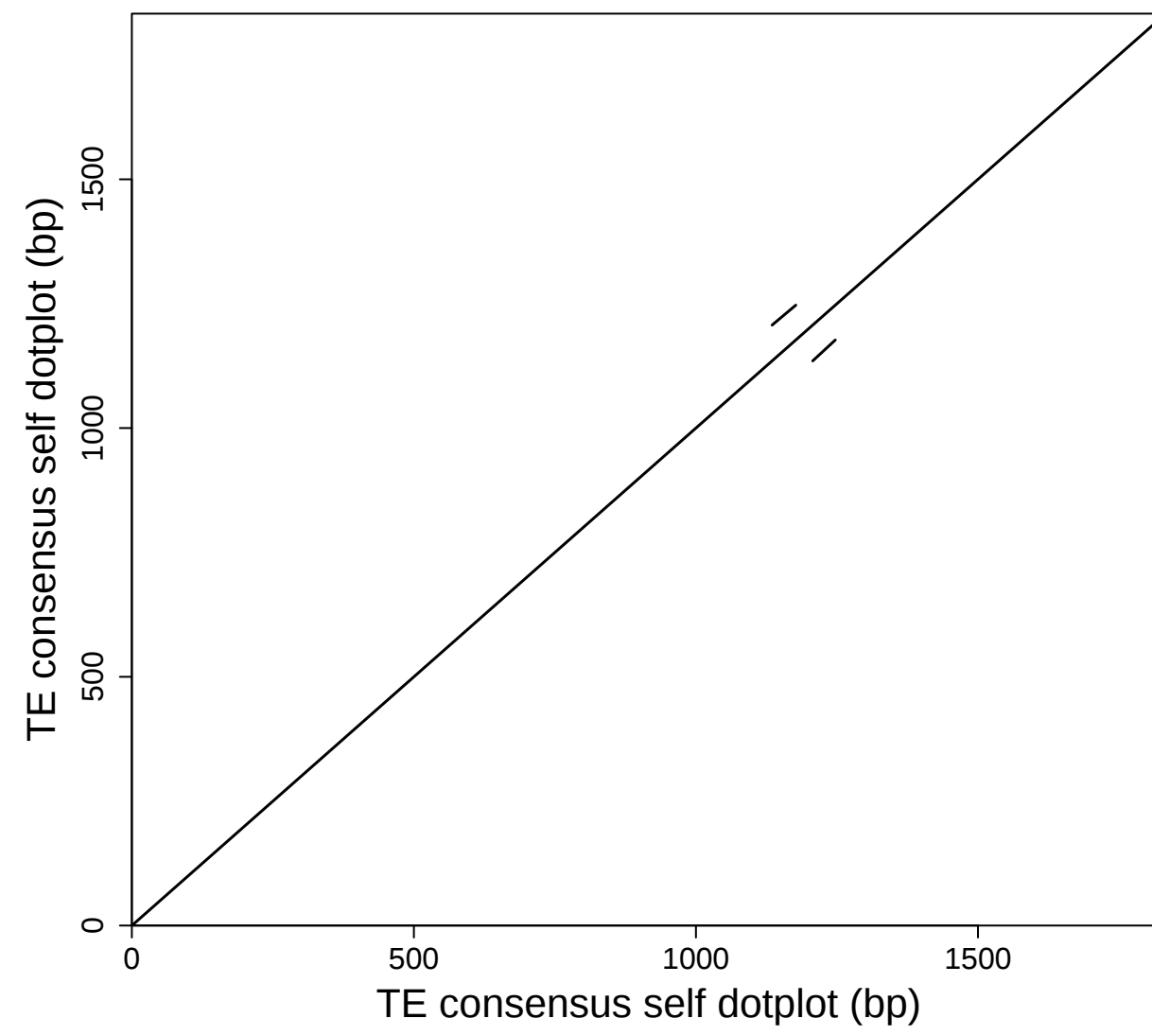
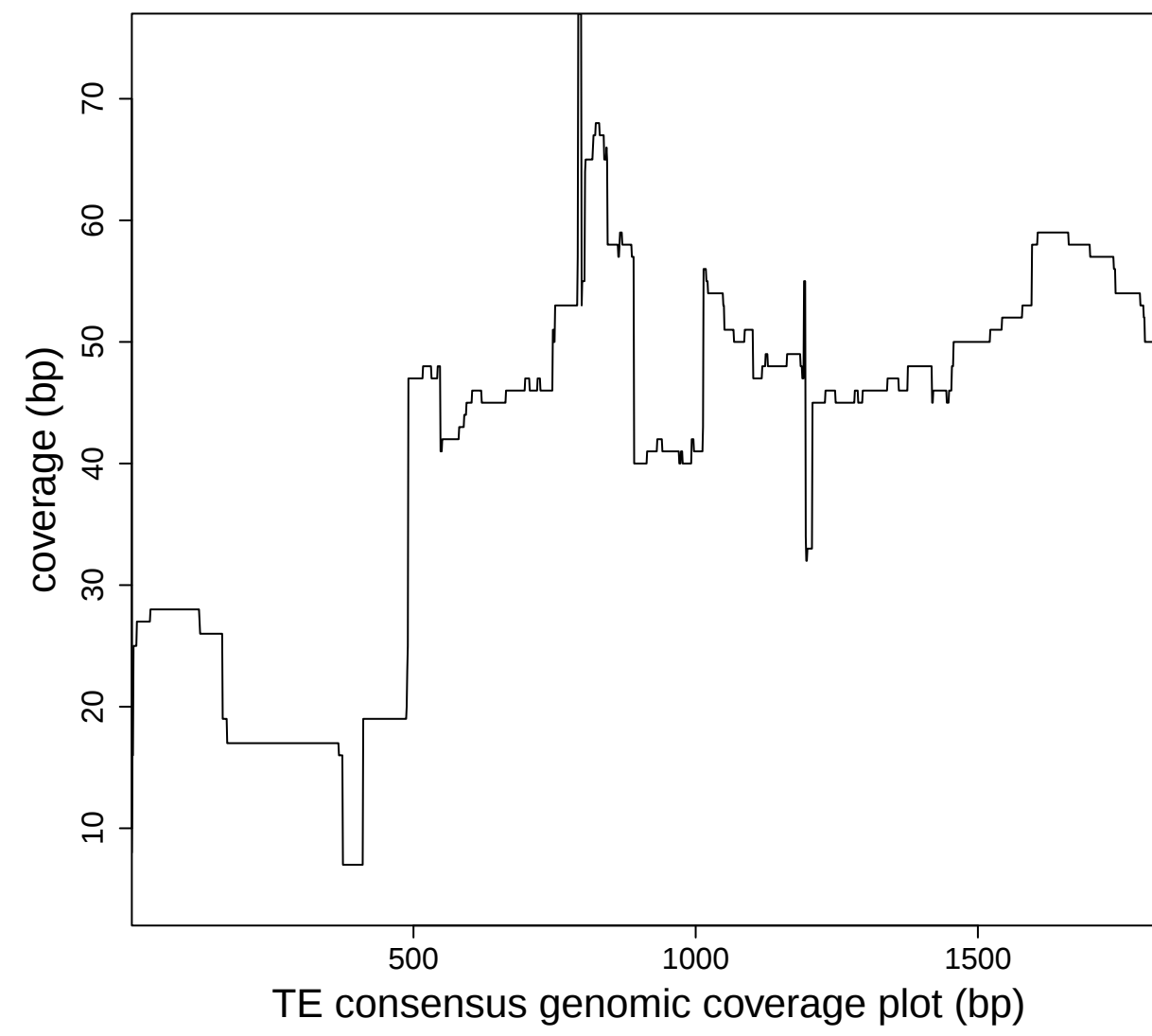
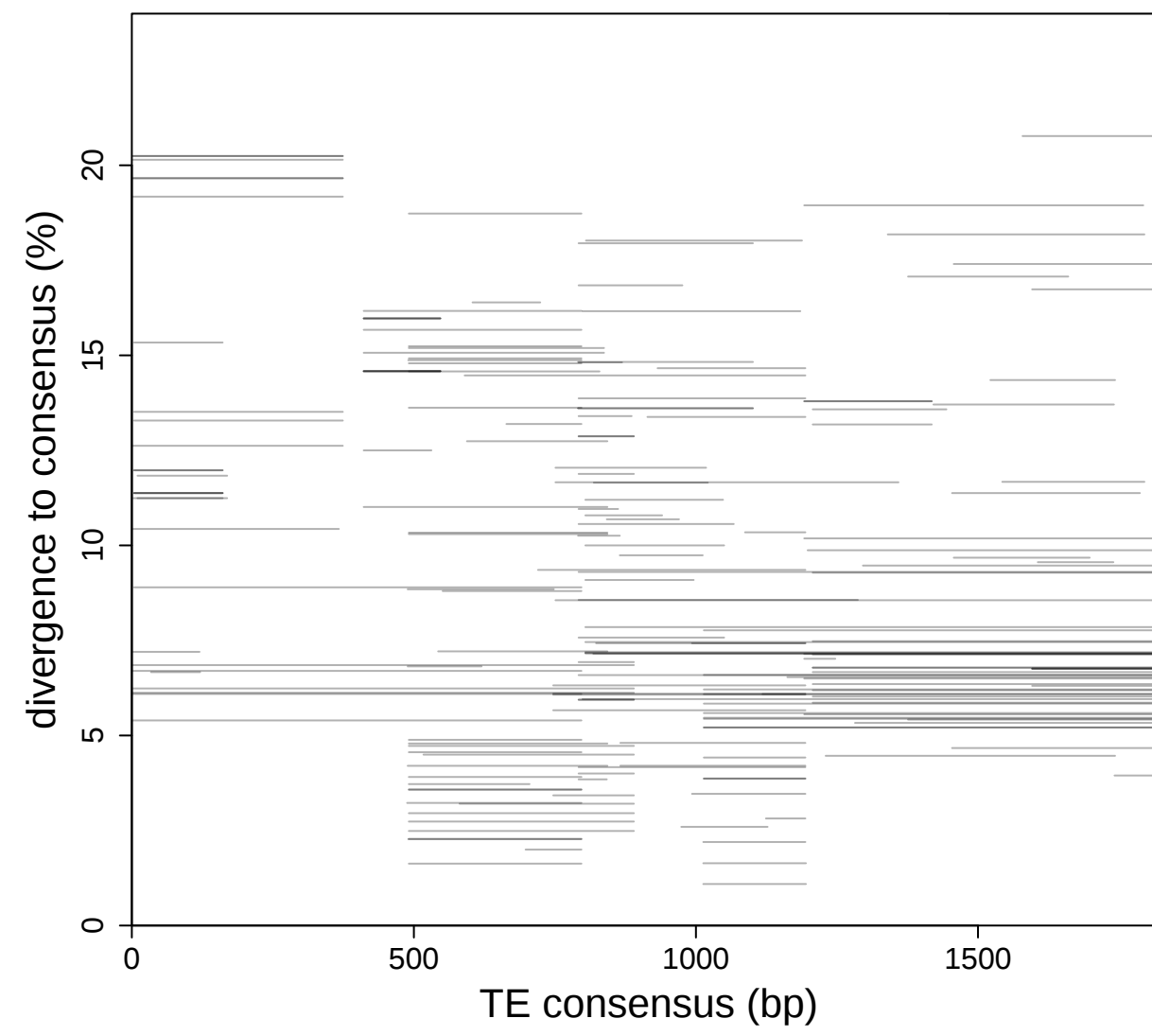
MSA length = 1833





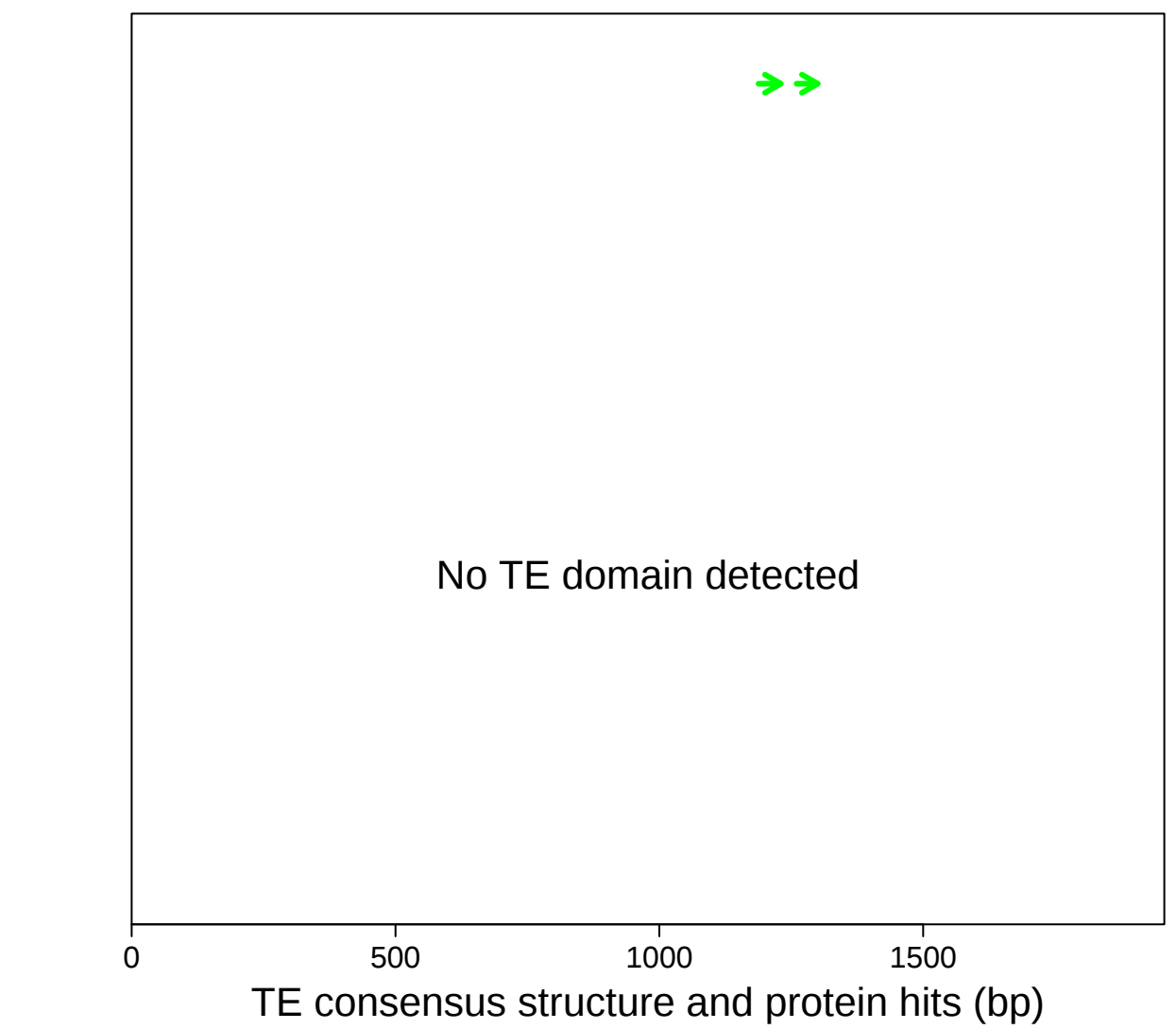
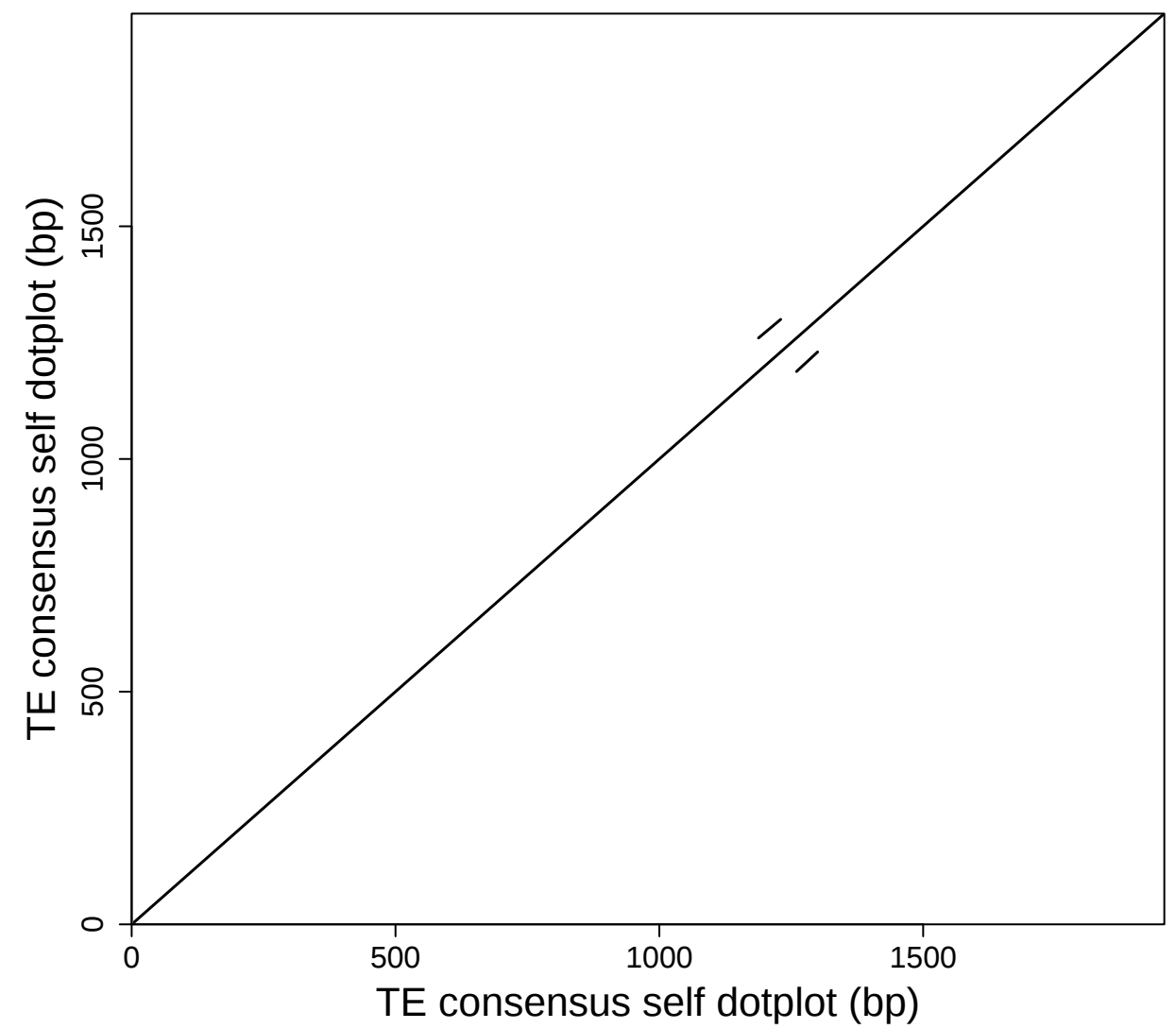
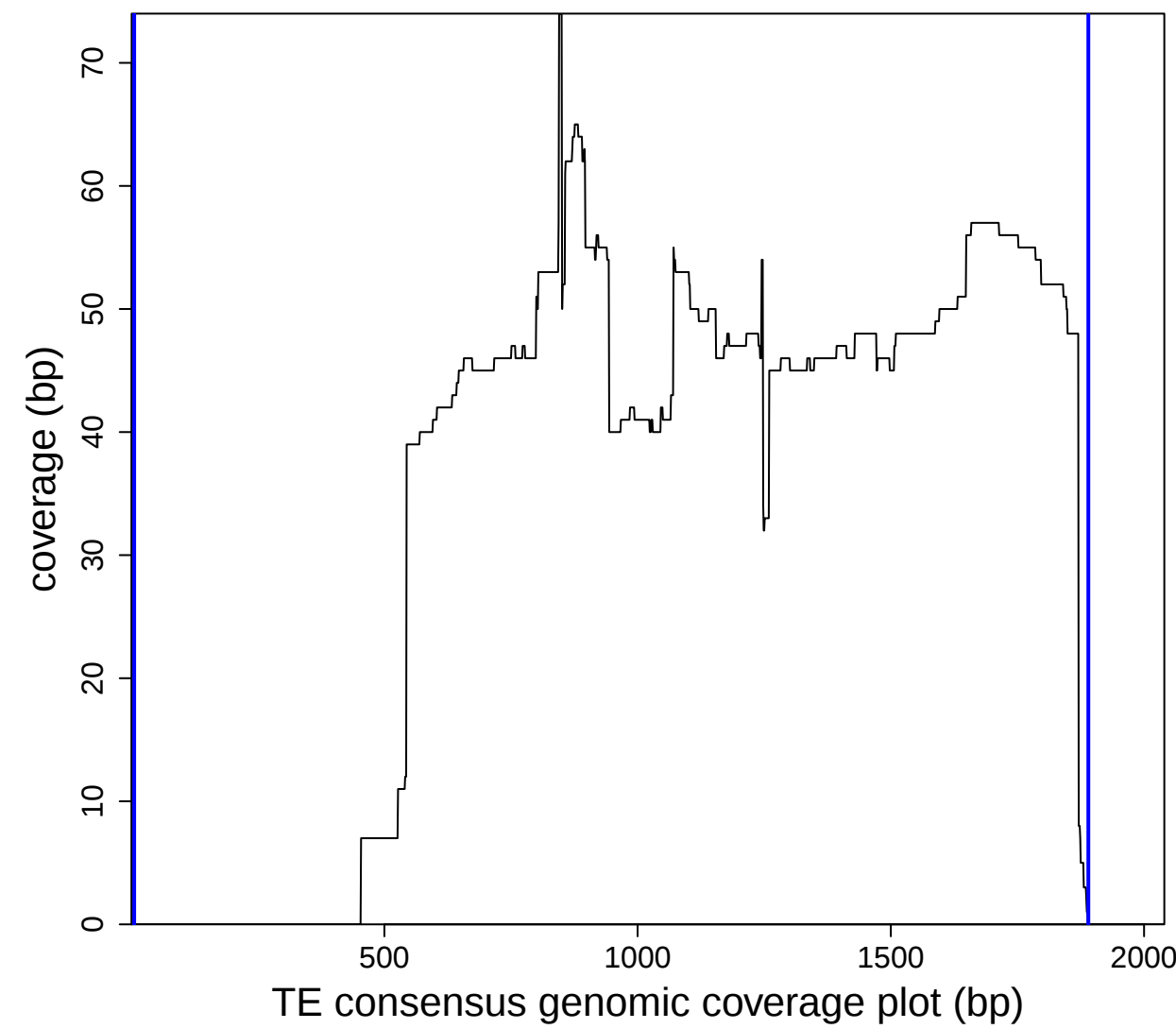
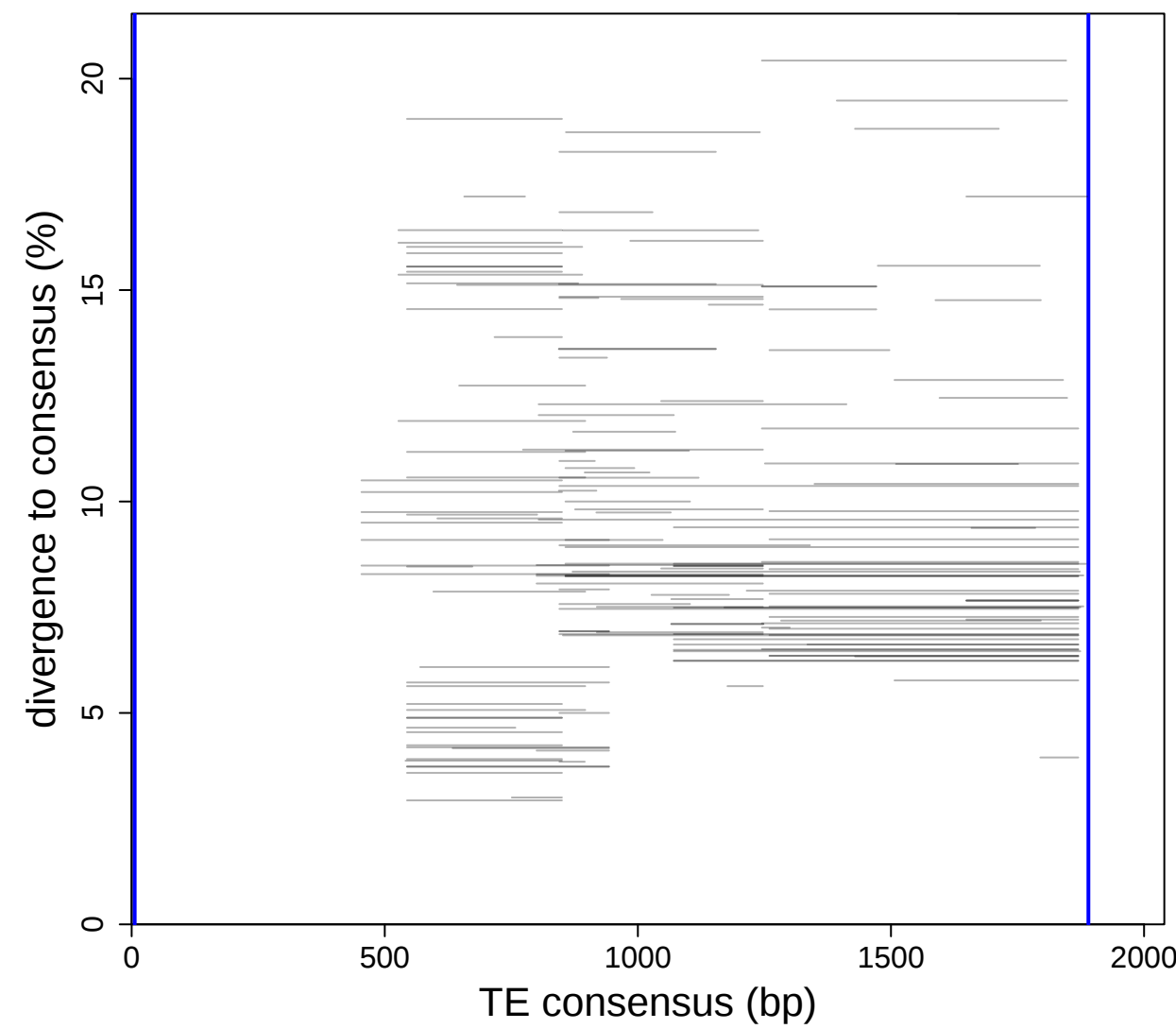
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_al
size: 1833bp; fragments: 197; full length: 0 (>=1649.7bp)

After TEtrimmer 1833 bp

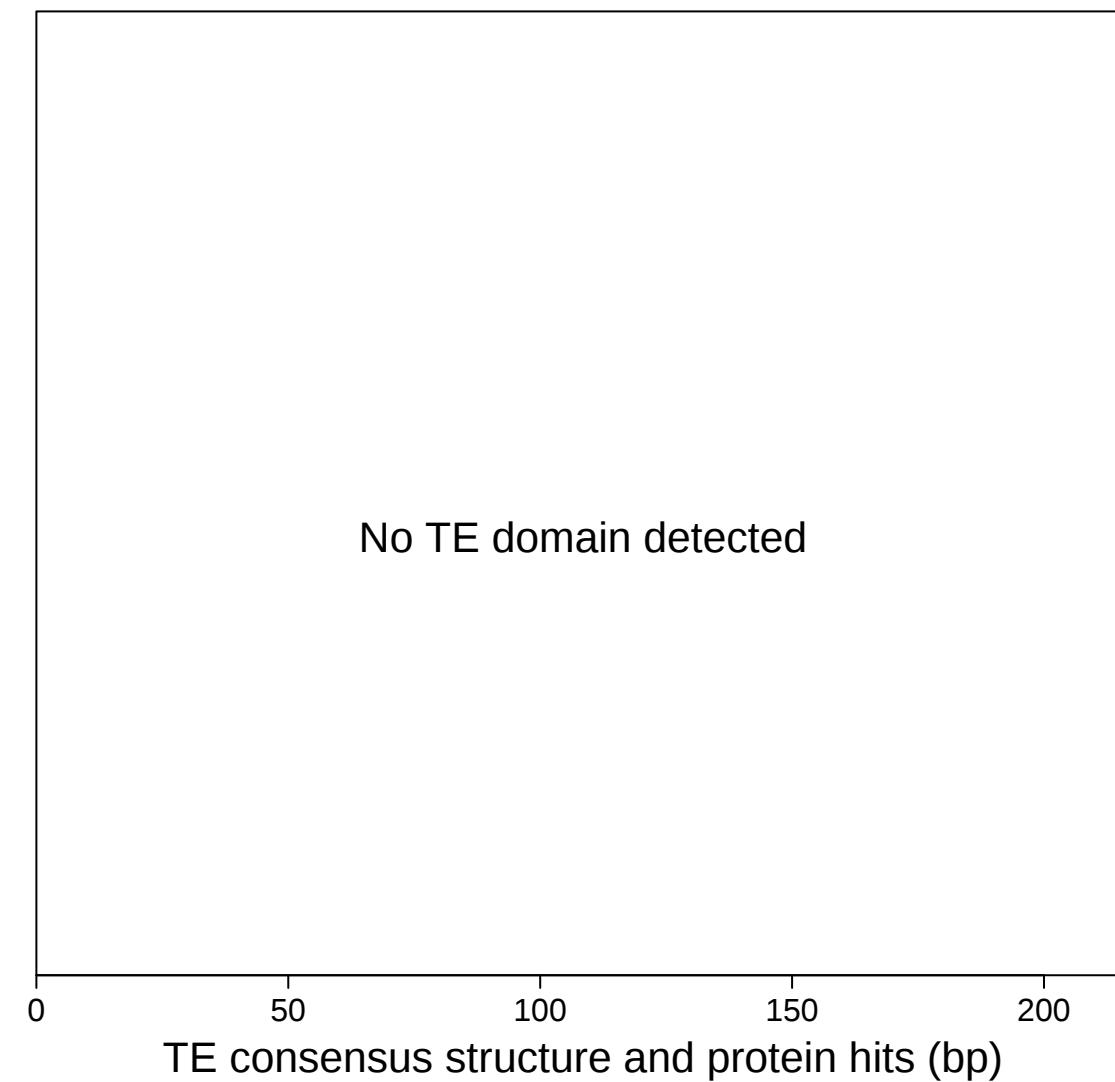
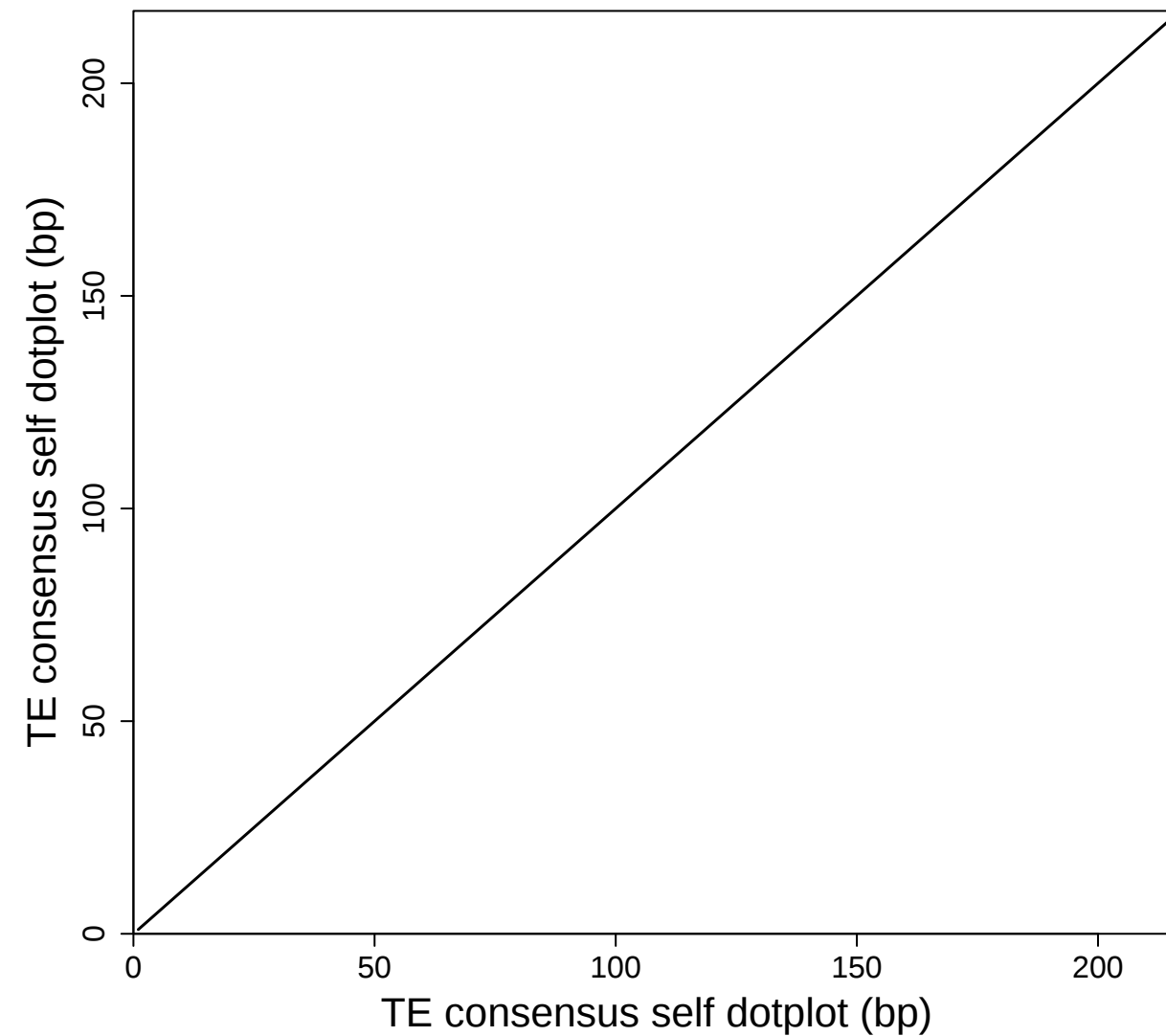
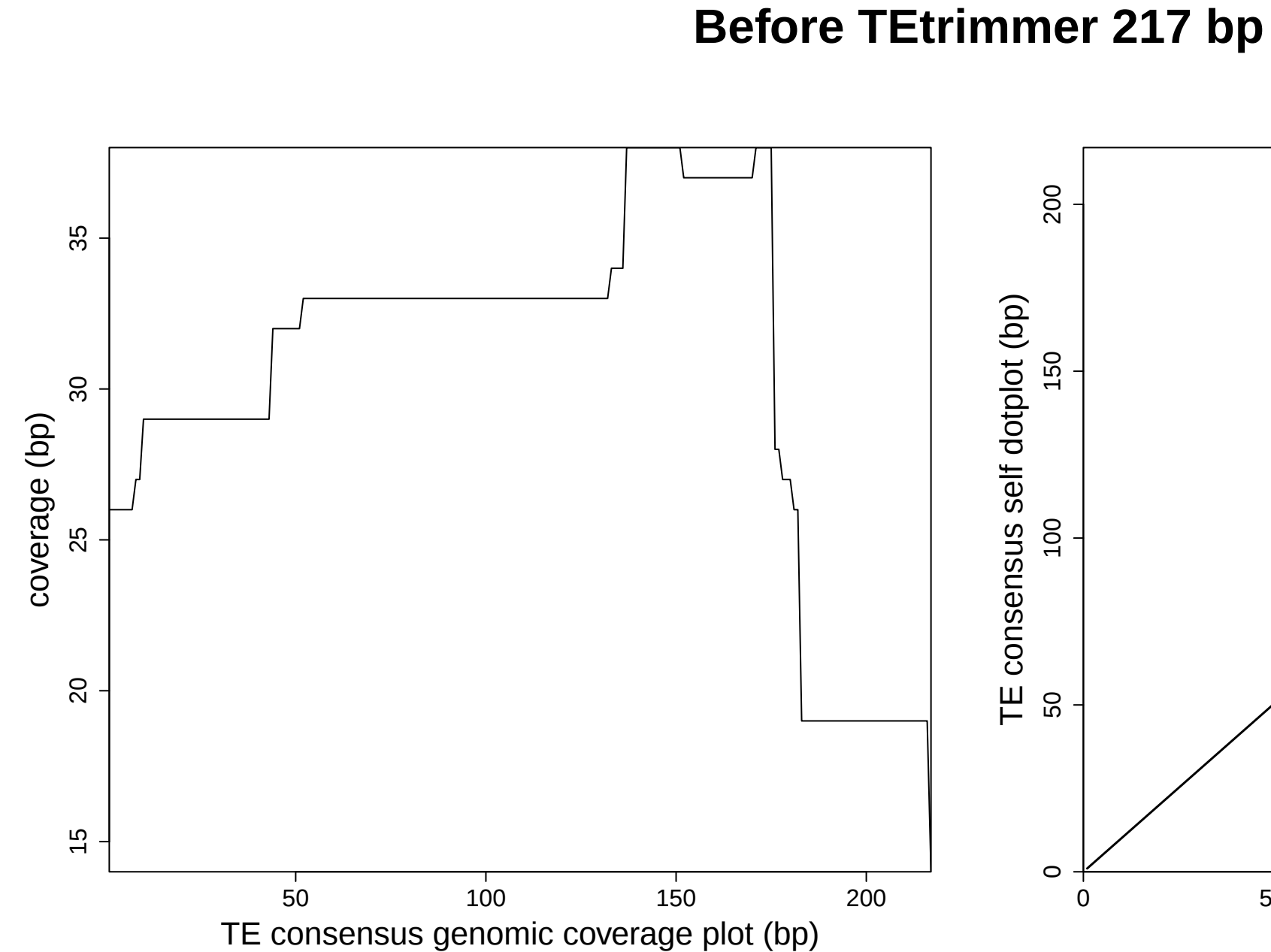
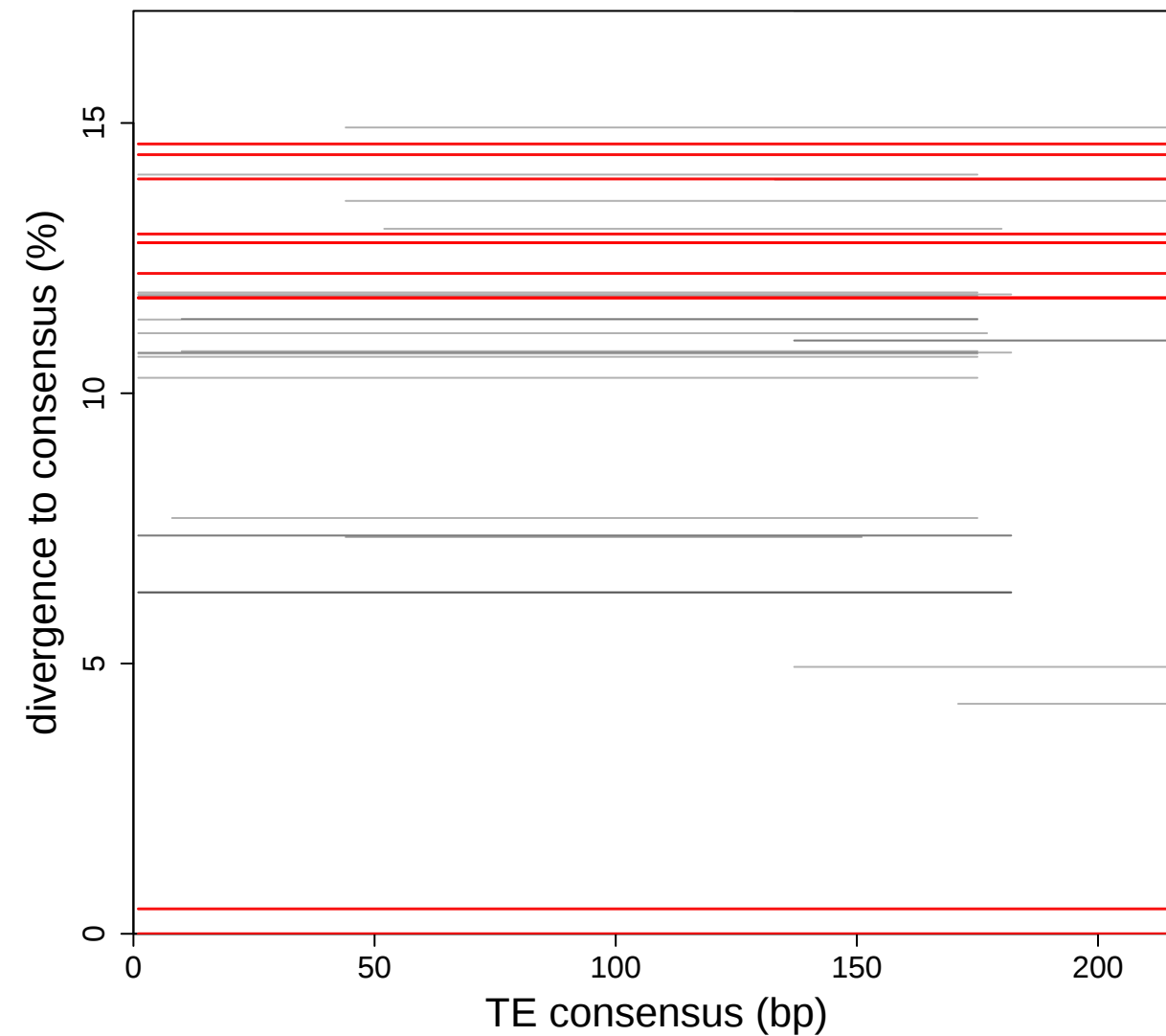


bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 2040bp; fragments: 162; full length: 0 (>=1836bp)

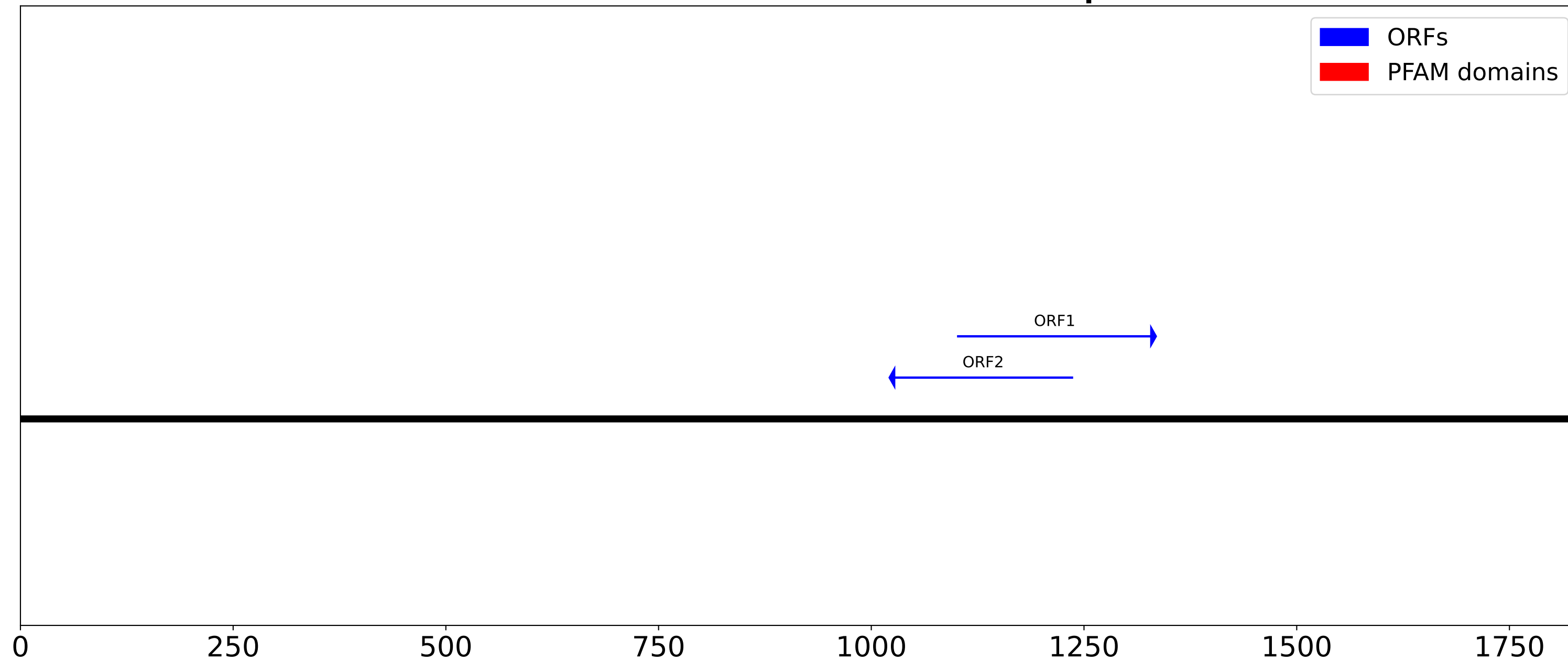
After TEtrimmer Extended plot Blue lines are boundaries



TE: ltr_1_family_337
size: 217bp; fragments: 39; full length: 11 (≥ 195.3 bp)



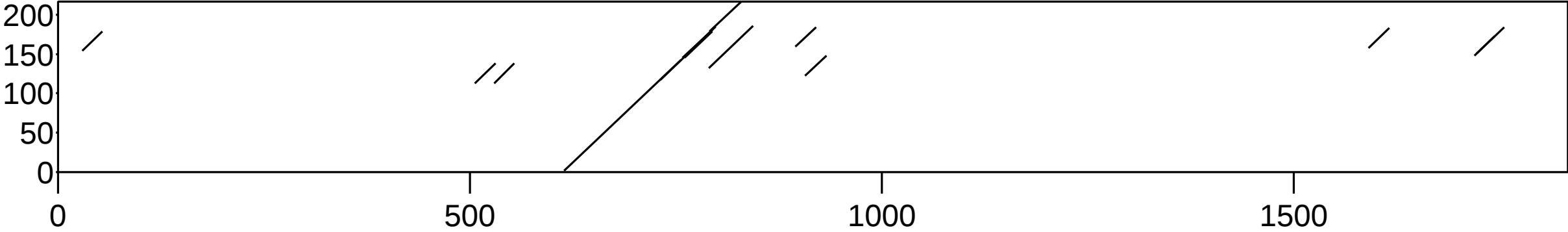
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

sensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)