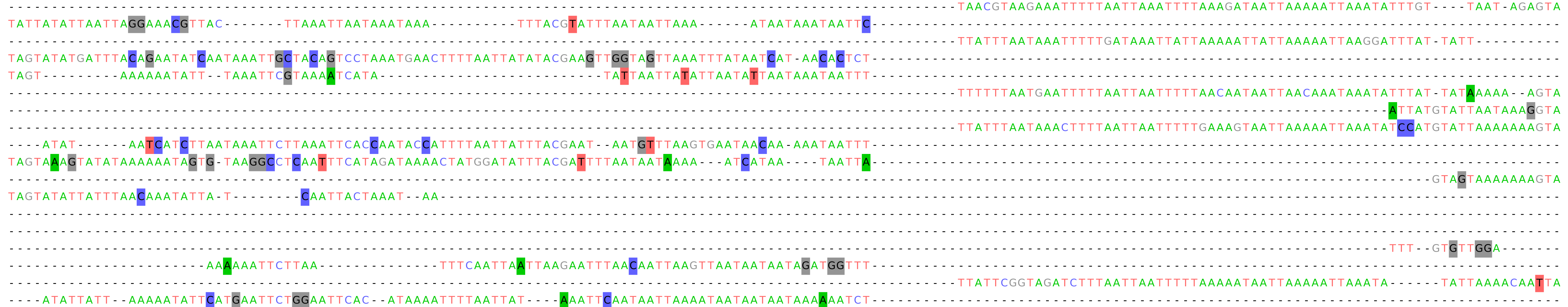
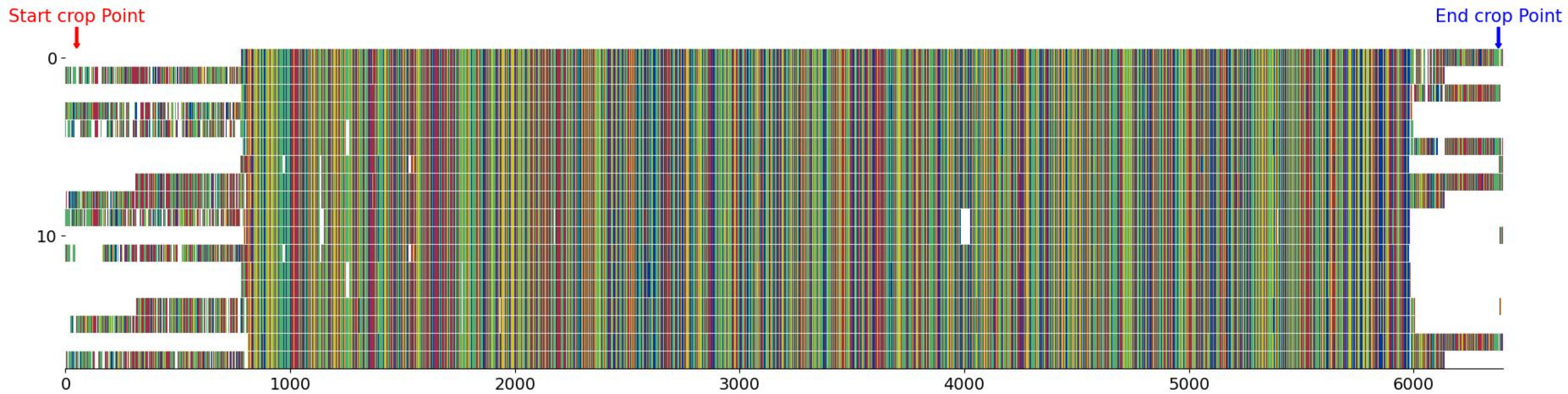


Start crop Point

End crop Point

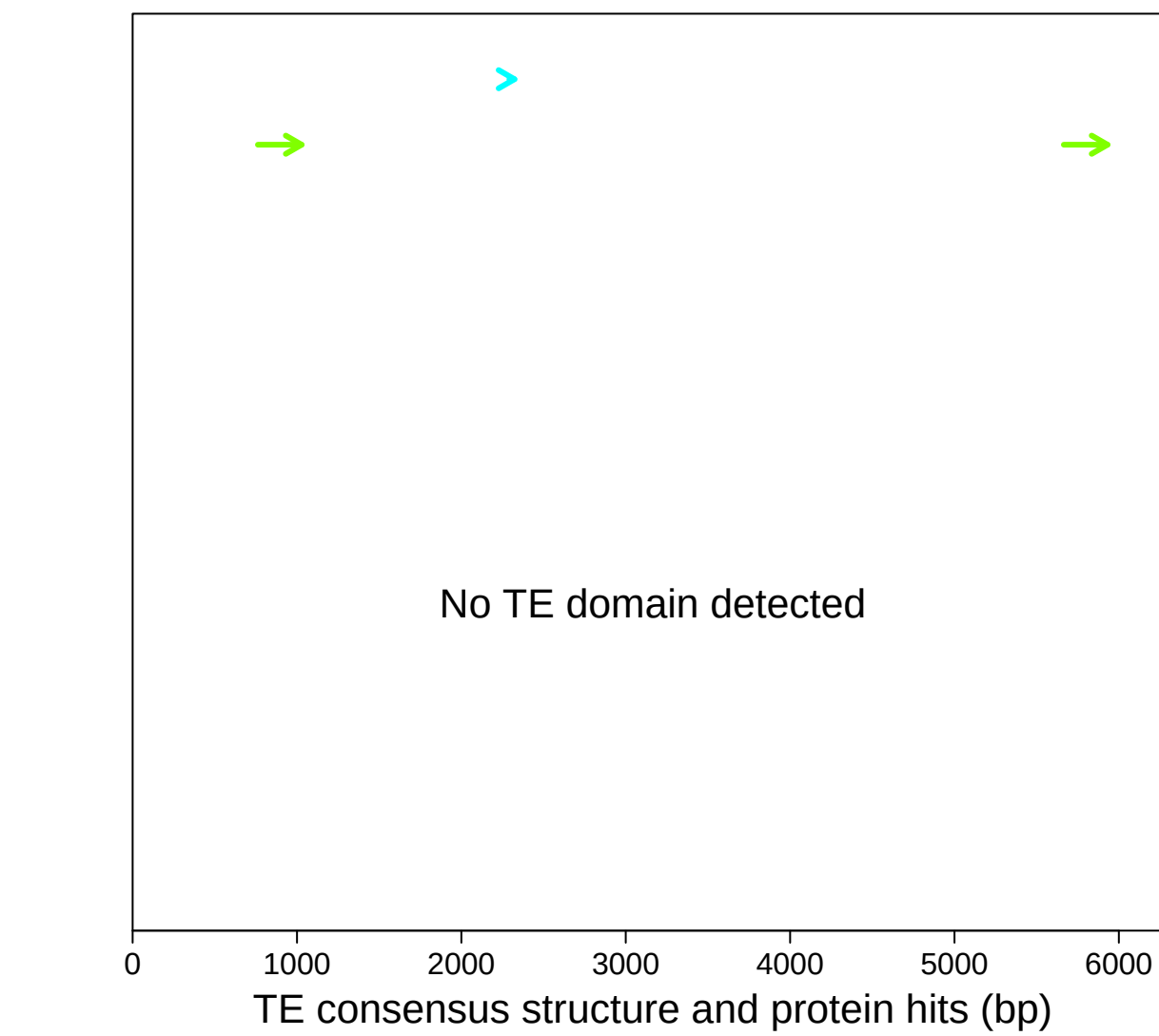
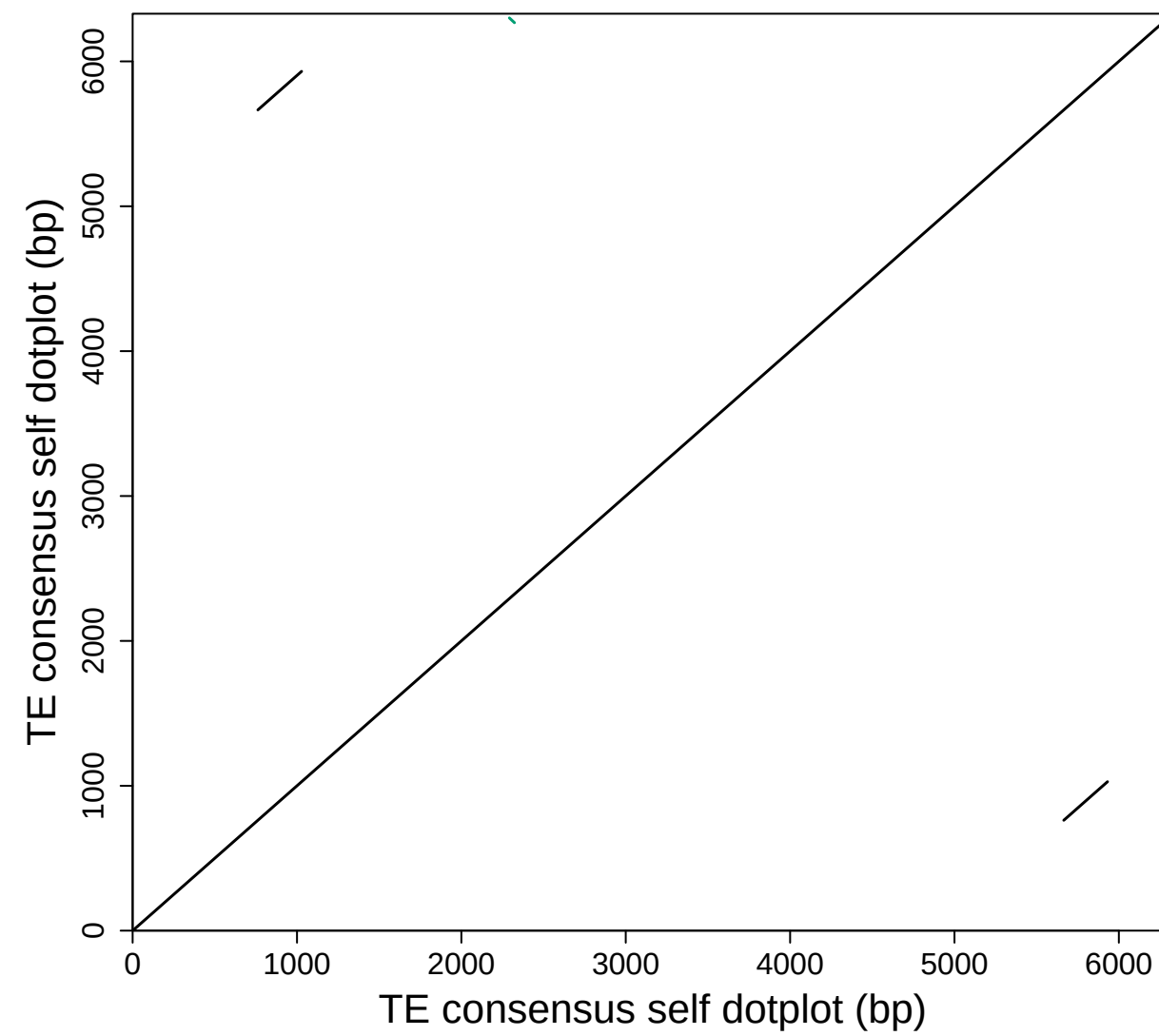
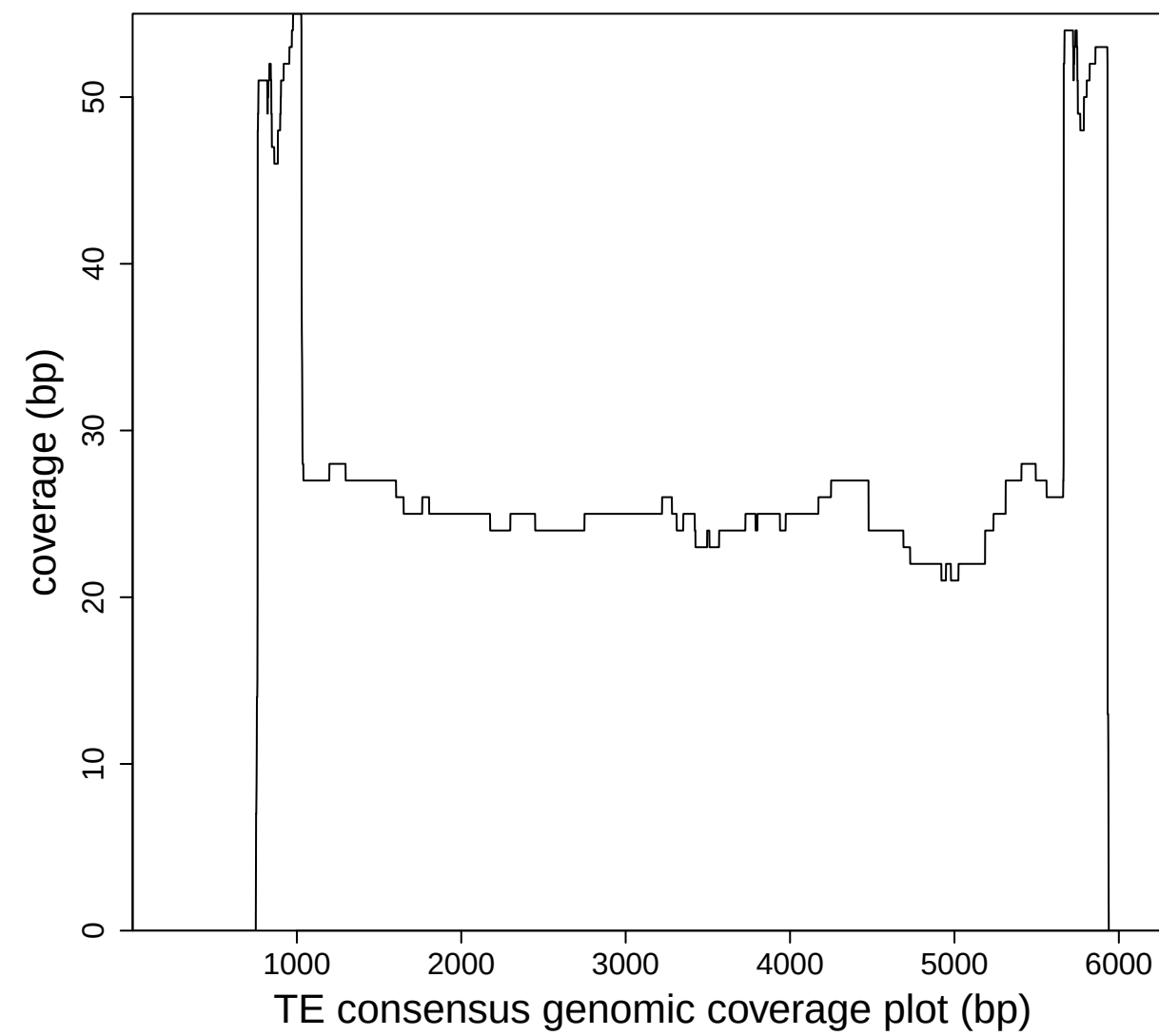
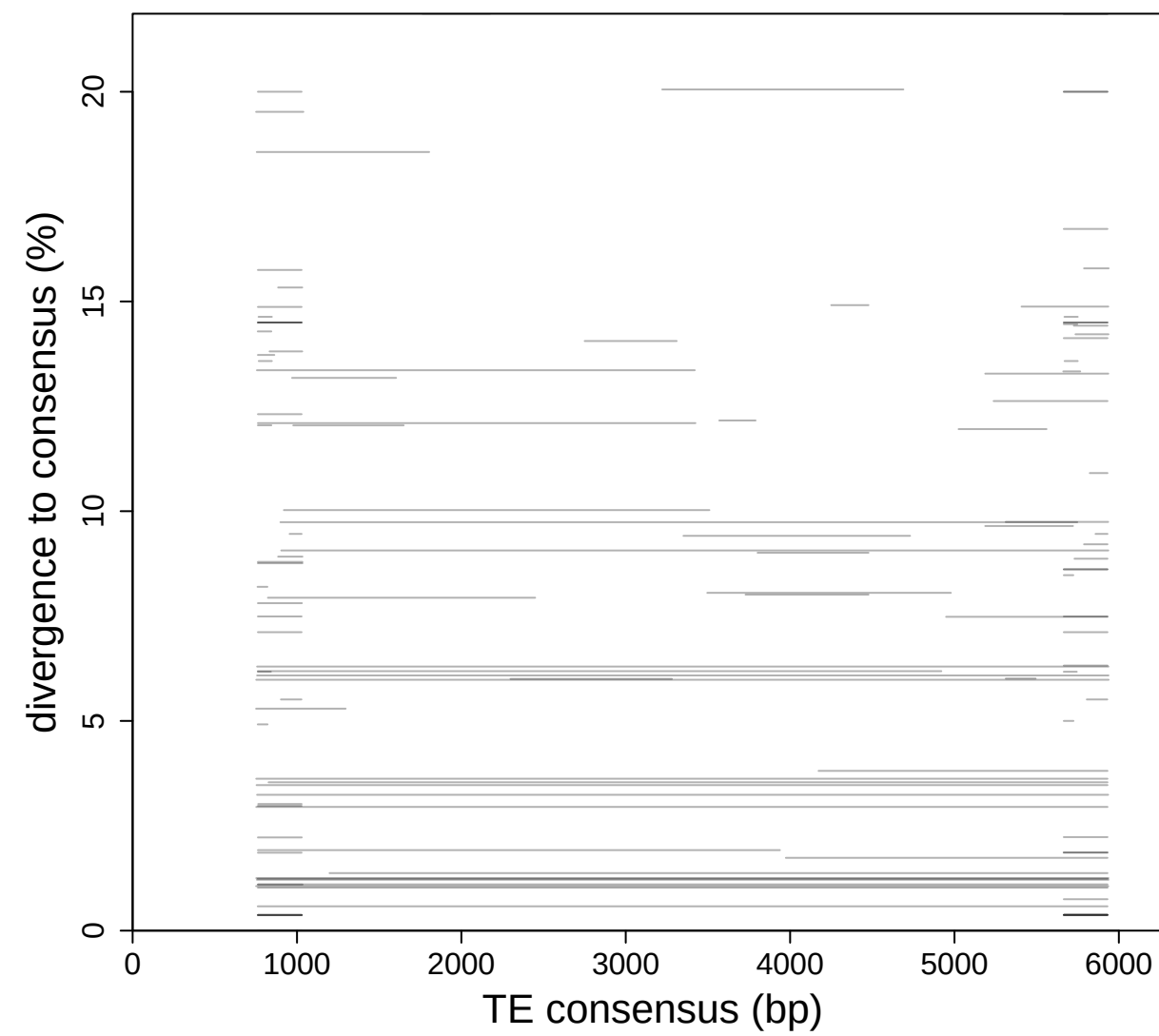
MSA length = 6329





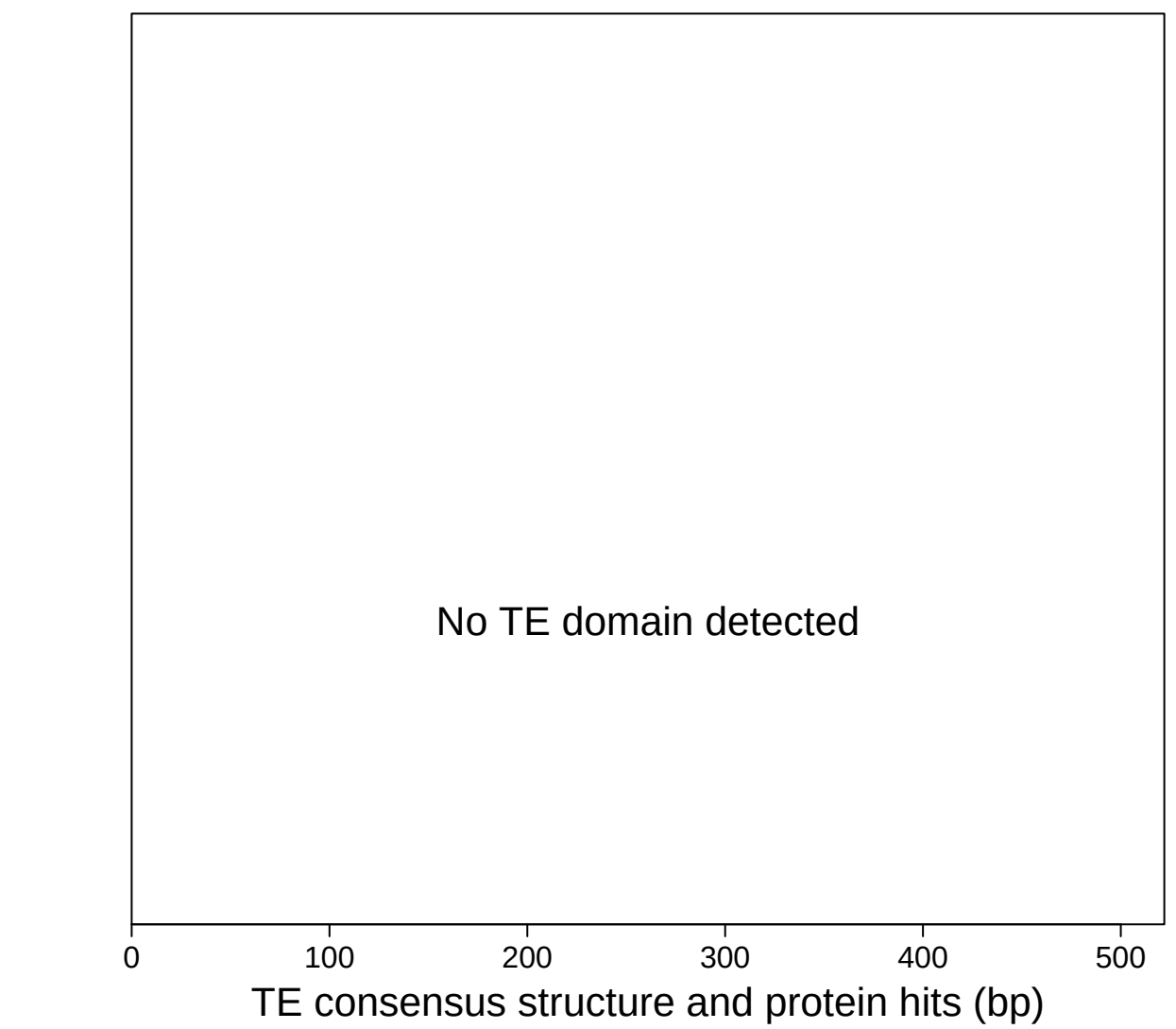
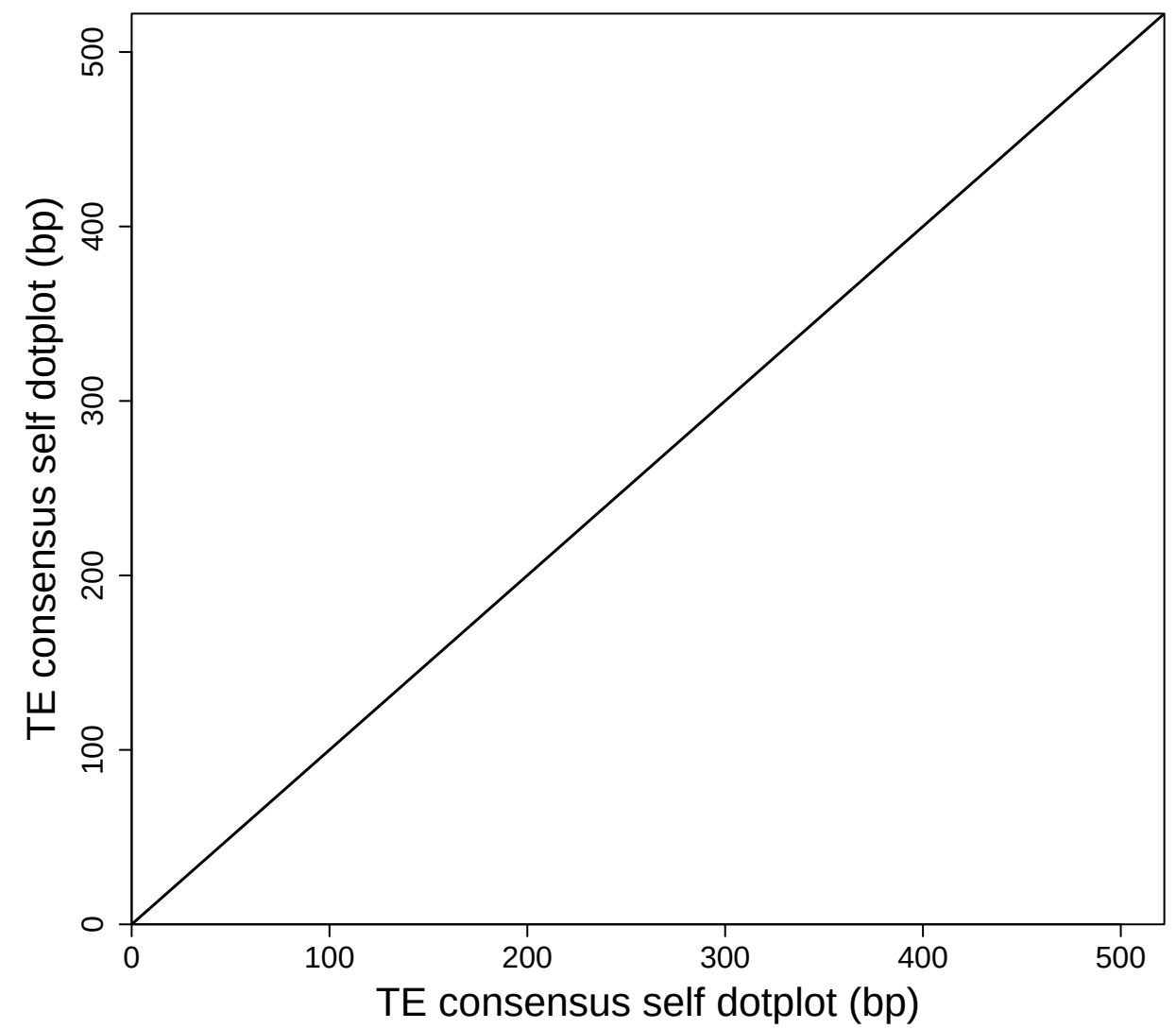
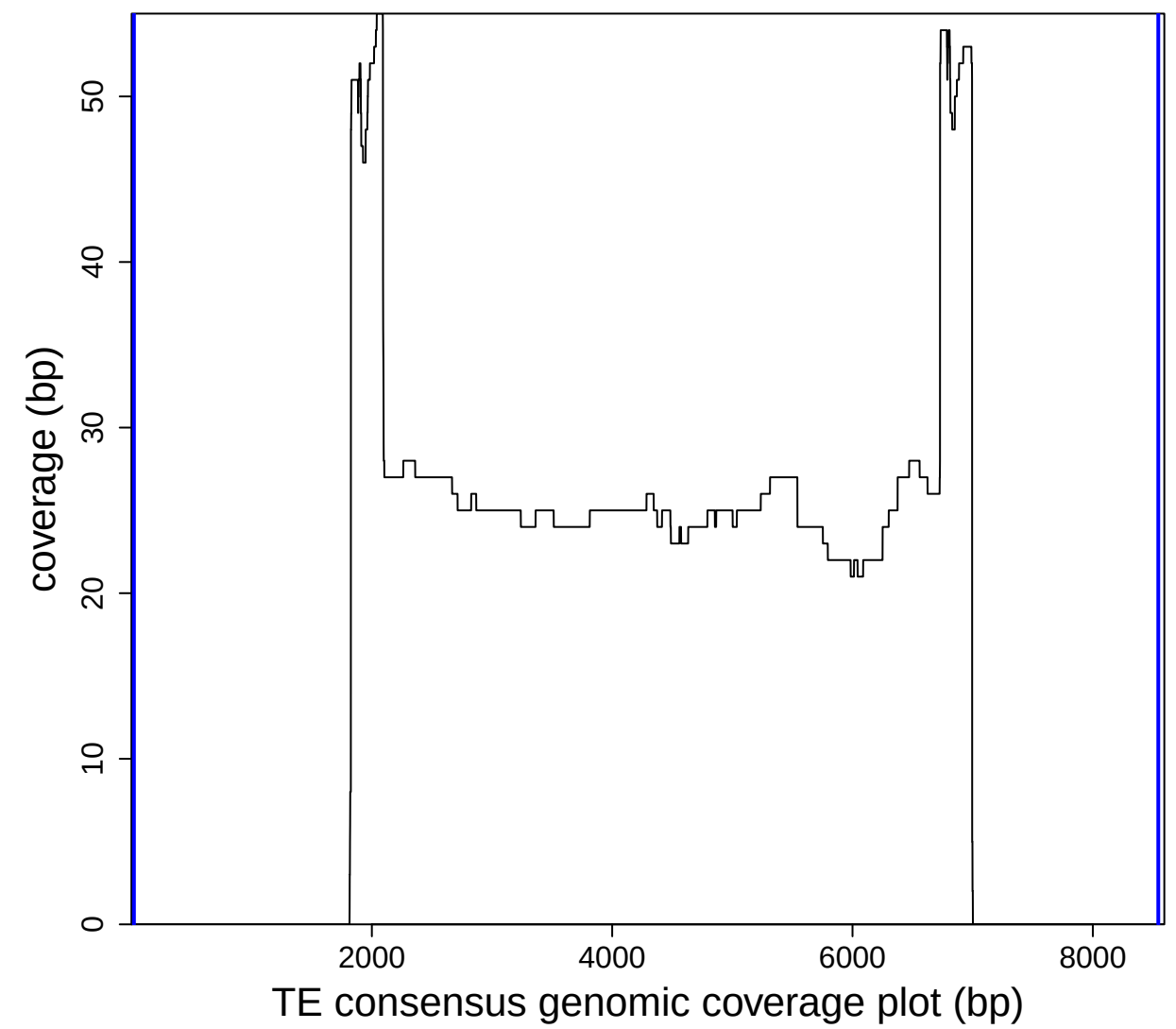
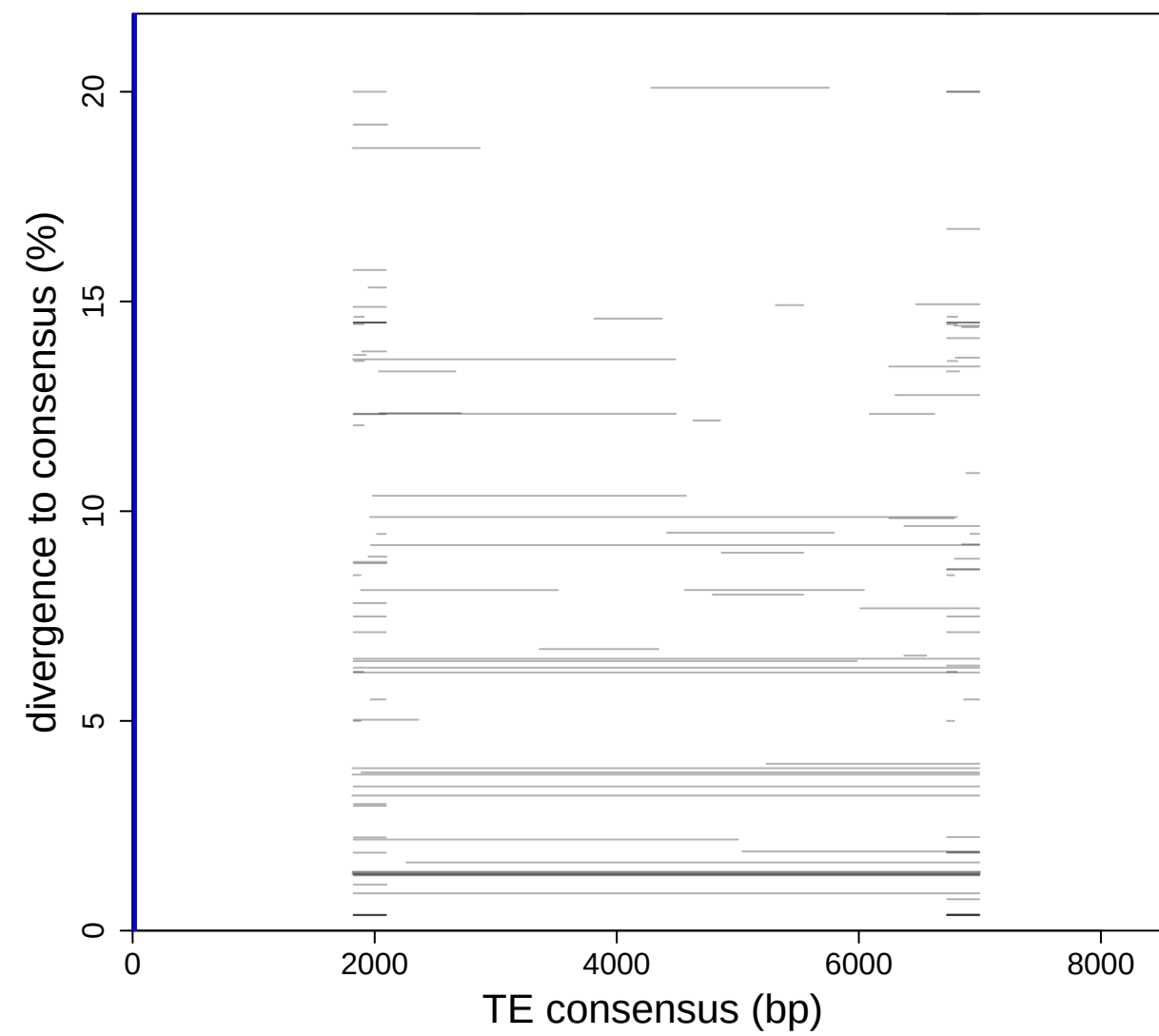
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_al
size: 6329bp; fragments: 120; full length: 0 (>=5696.1bp)

After TEtrimmer 6329 bp

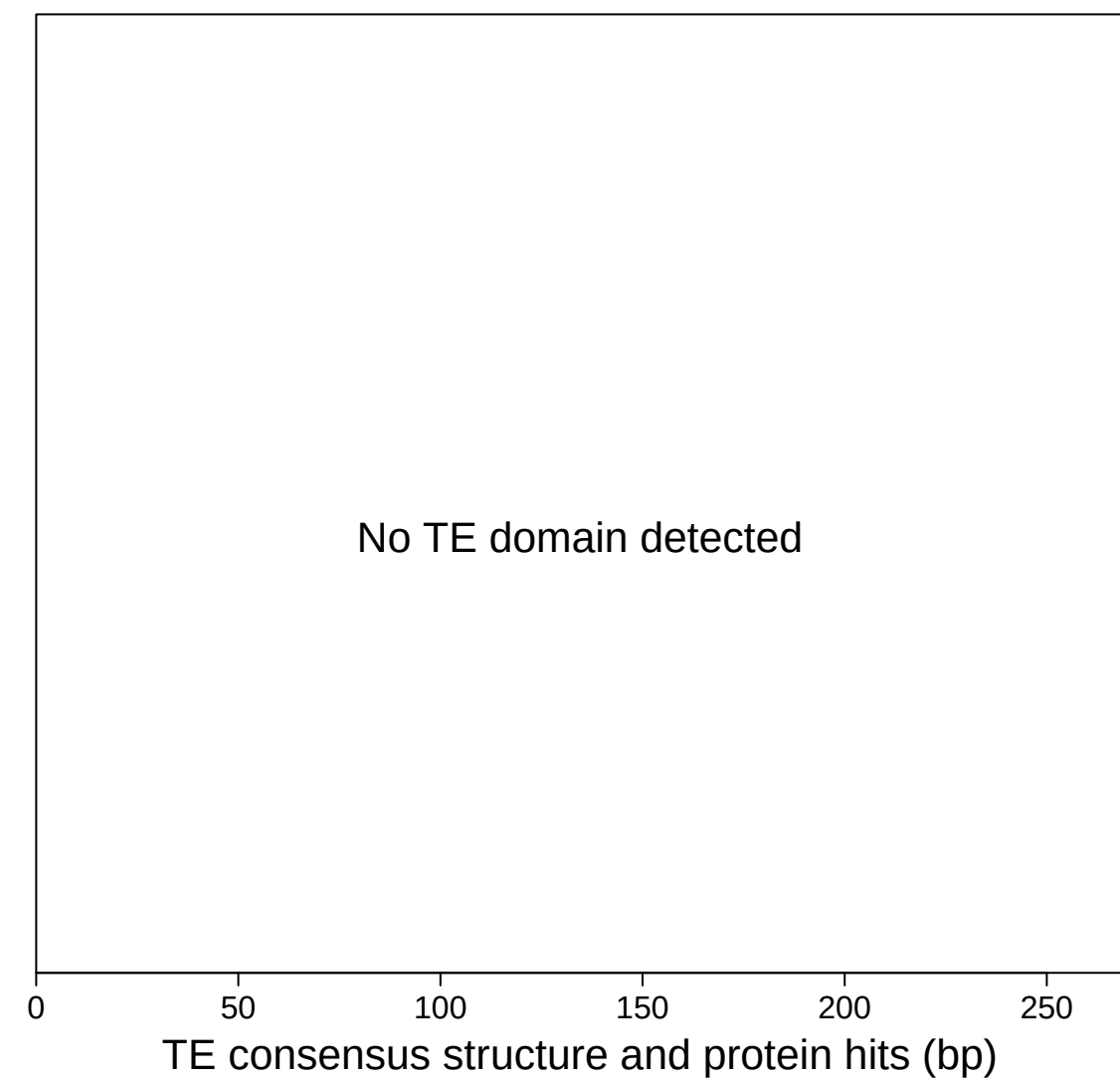
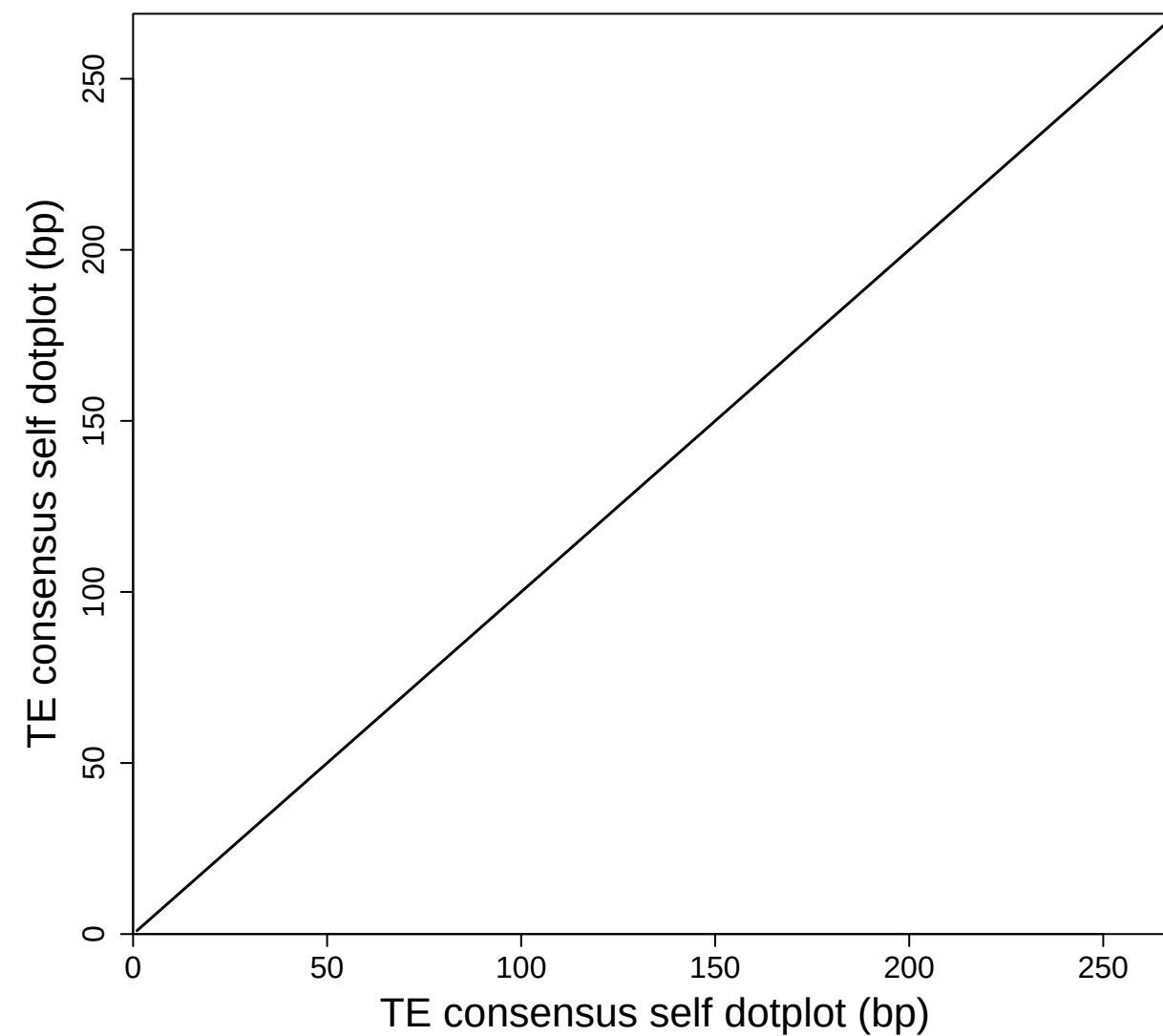
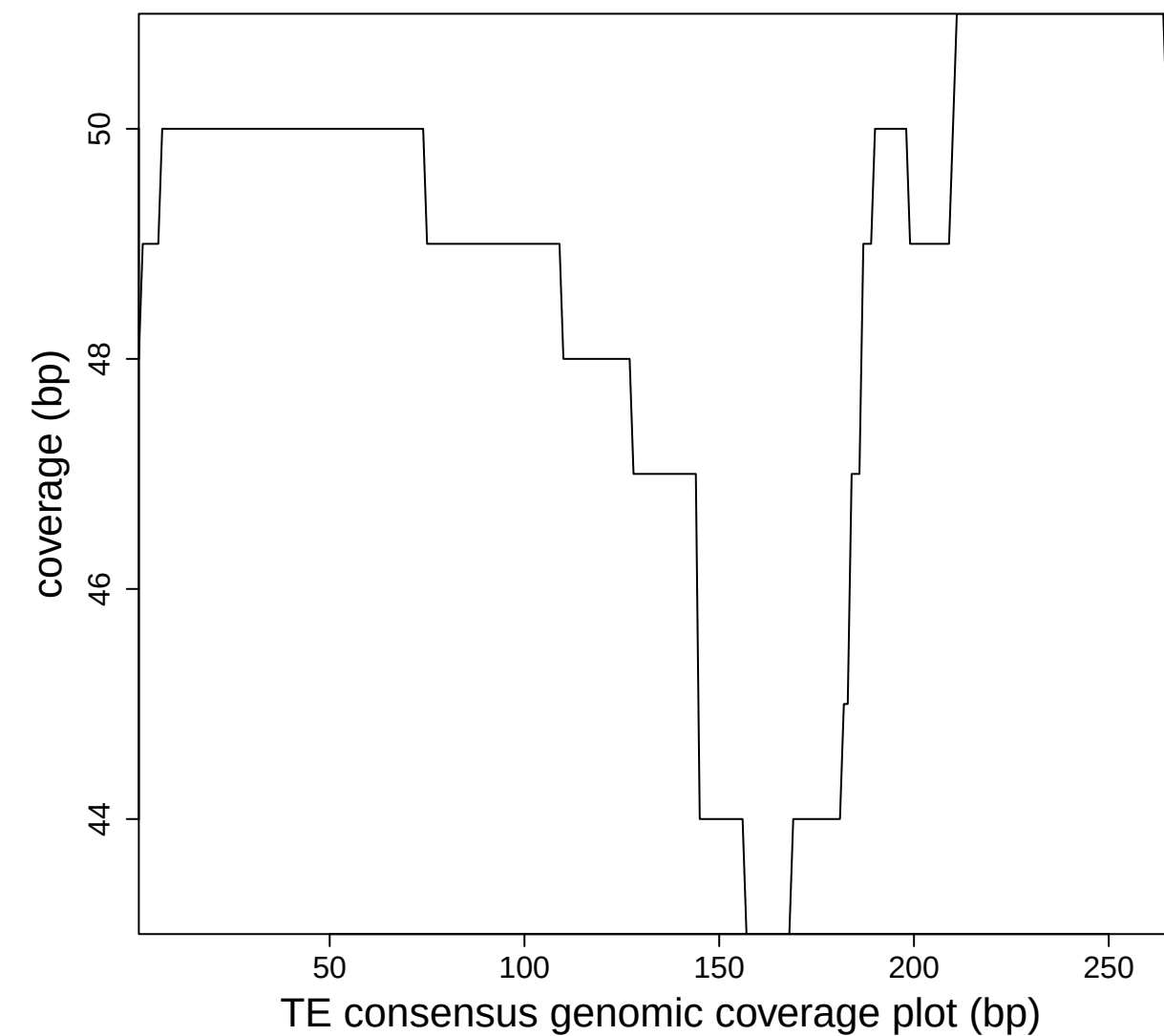
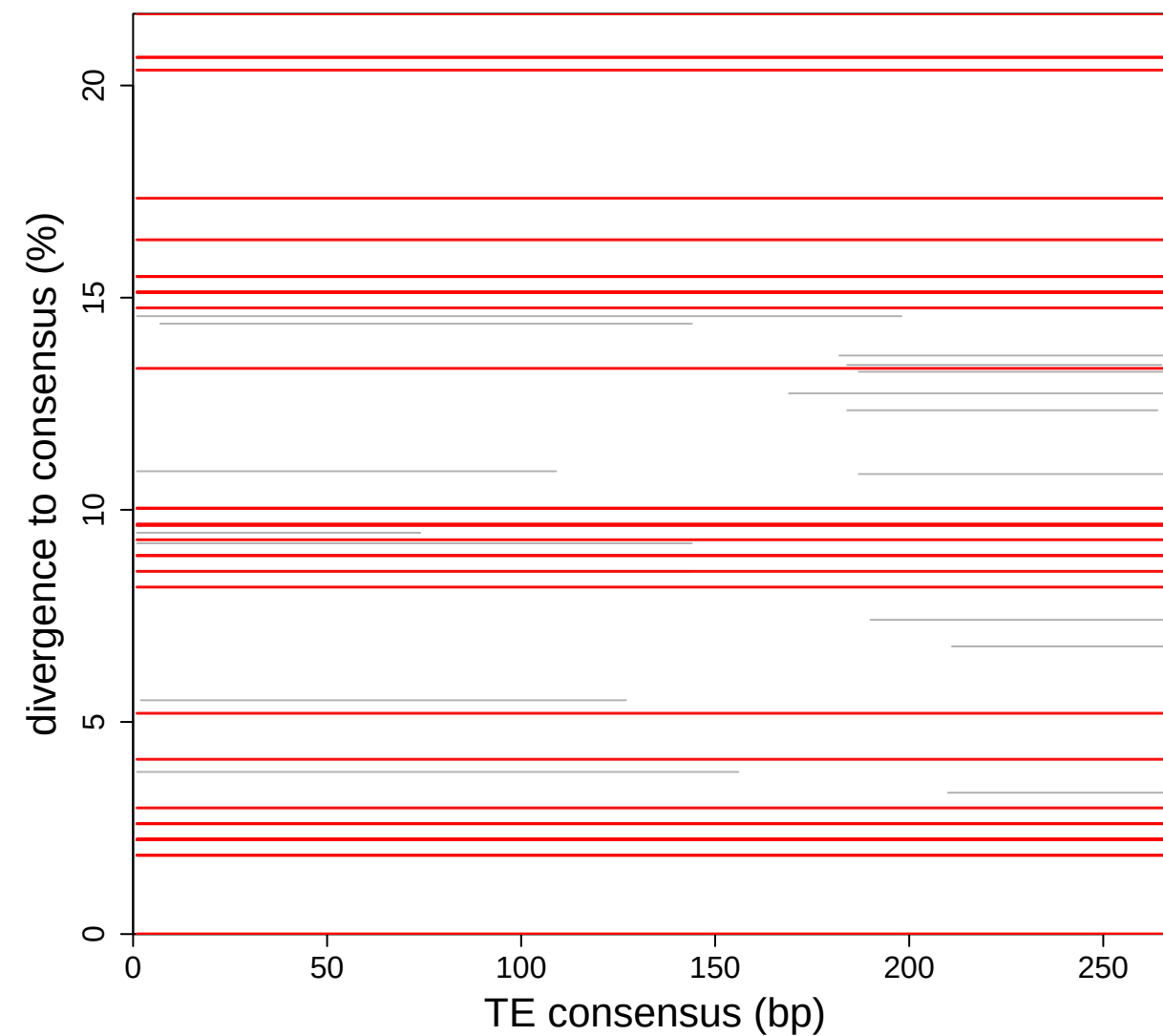


d_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa
size: 8595bp; fragments: 120; full length: 0 (>=7735.5bp)

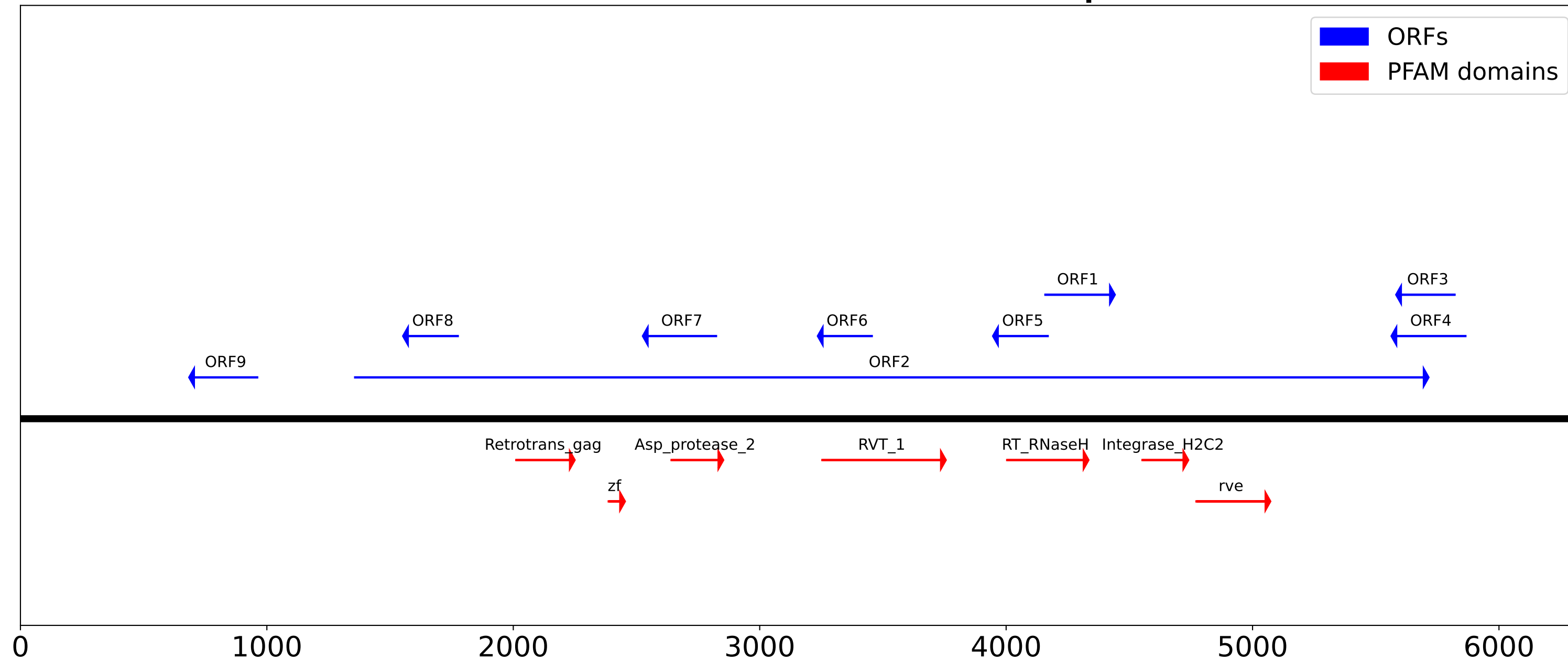
After TEtrimmer Extended plot Blue lines are boundaries



TE: ltr_1_family_352
size: 269bp; fragments: 59; full length: 42 (≥ 242.1 bp)



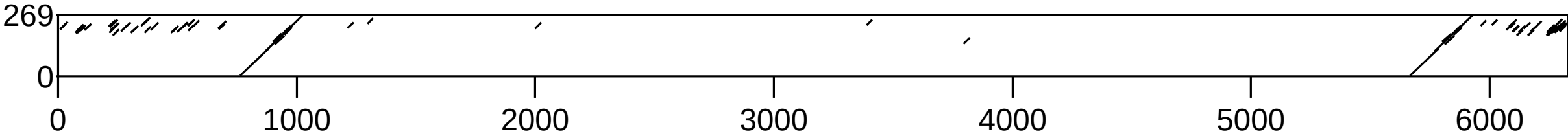
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)