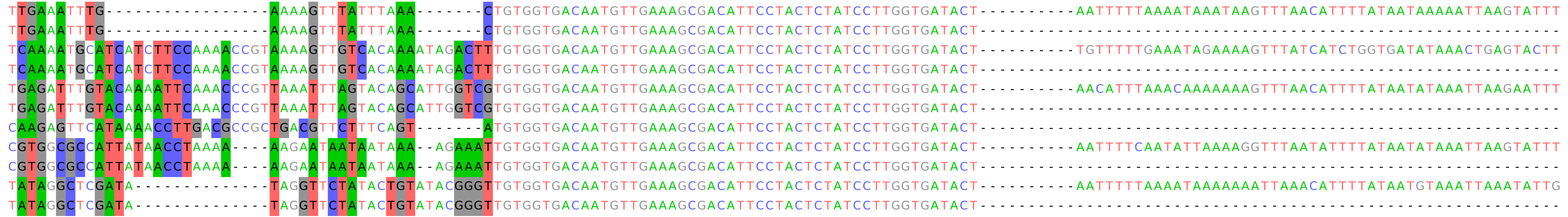


Start crop Point

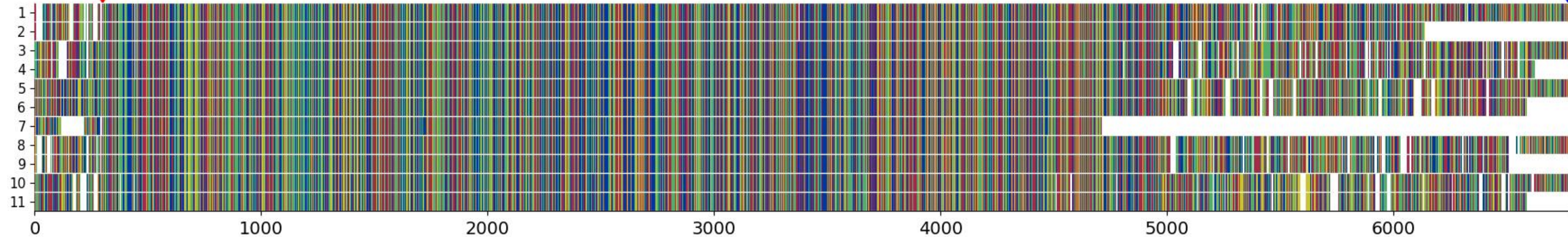
End crop Point

MSA length = 6474



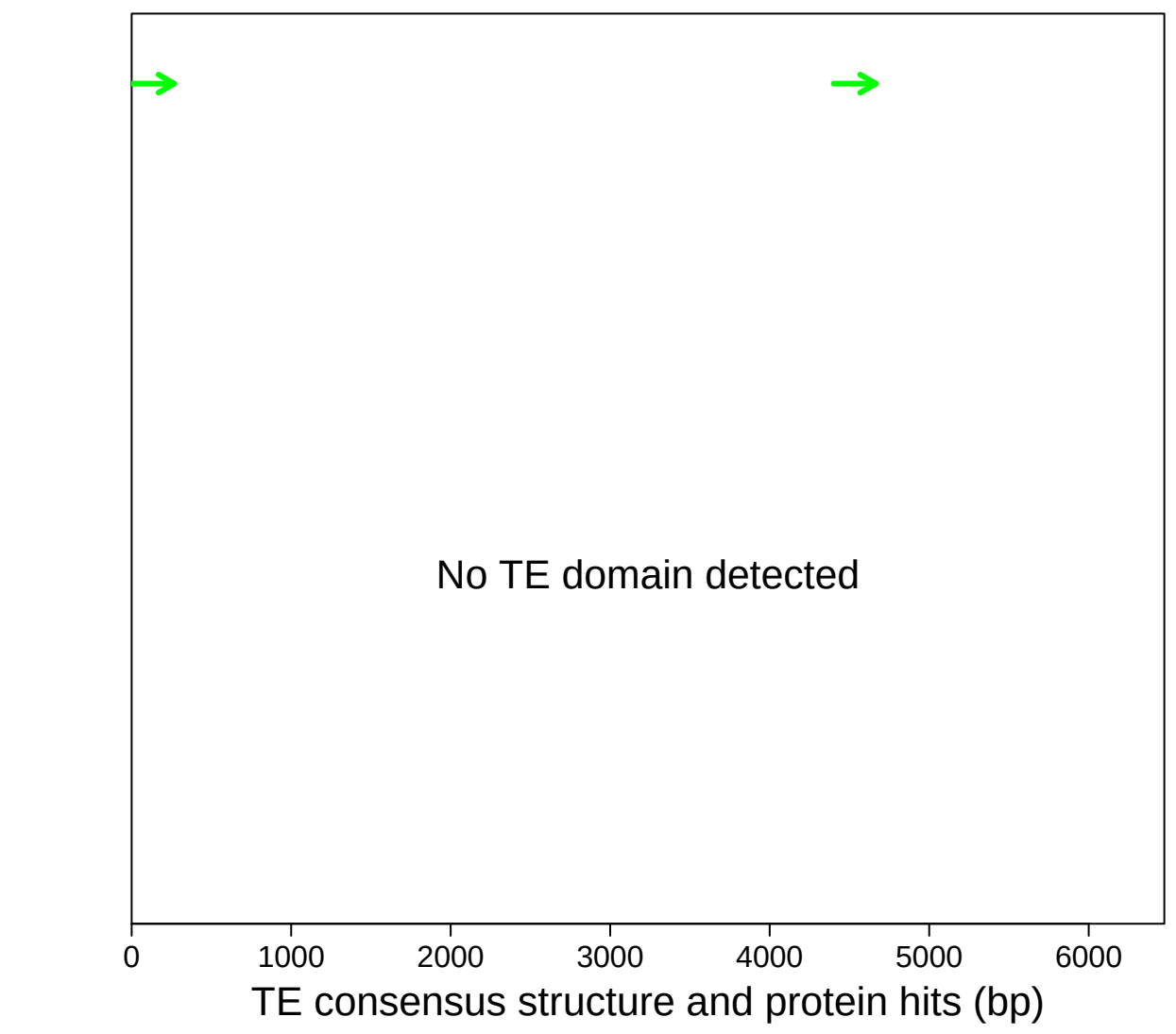
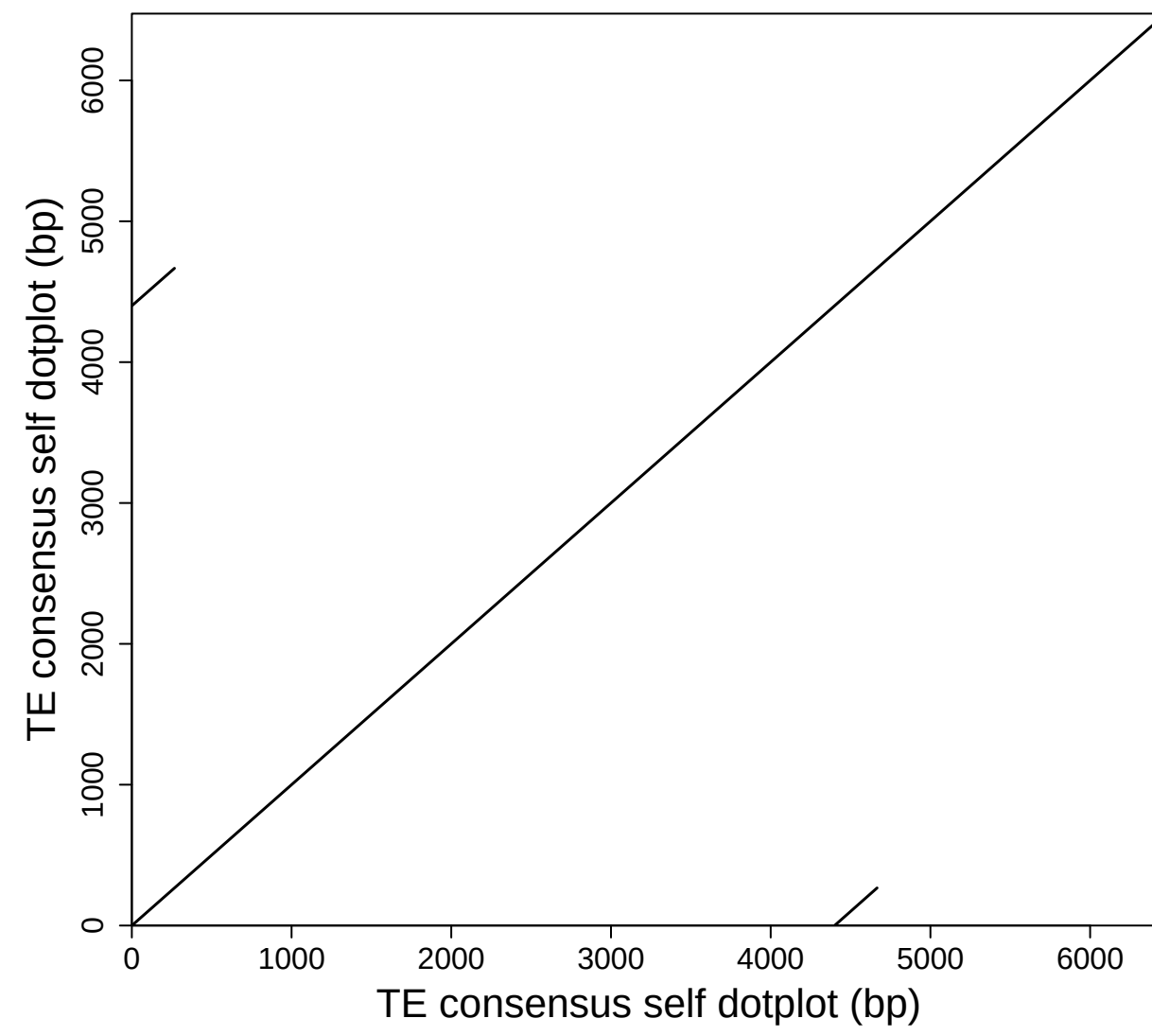
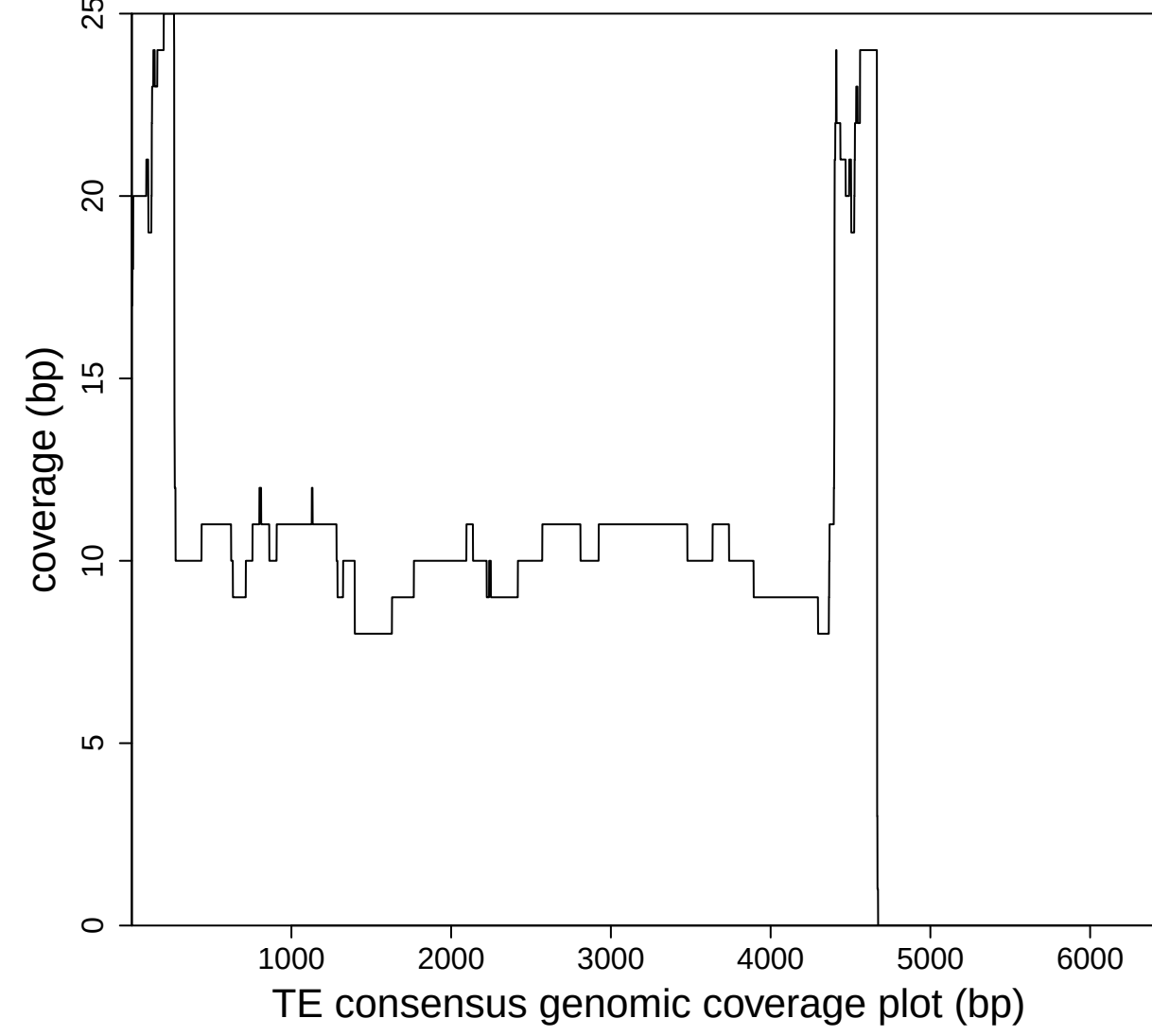
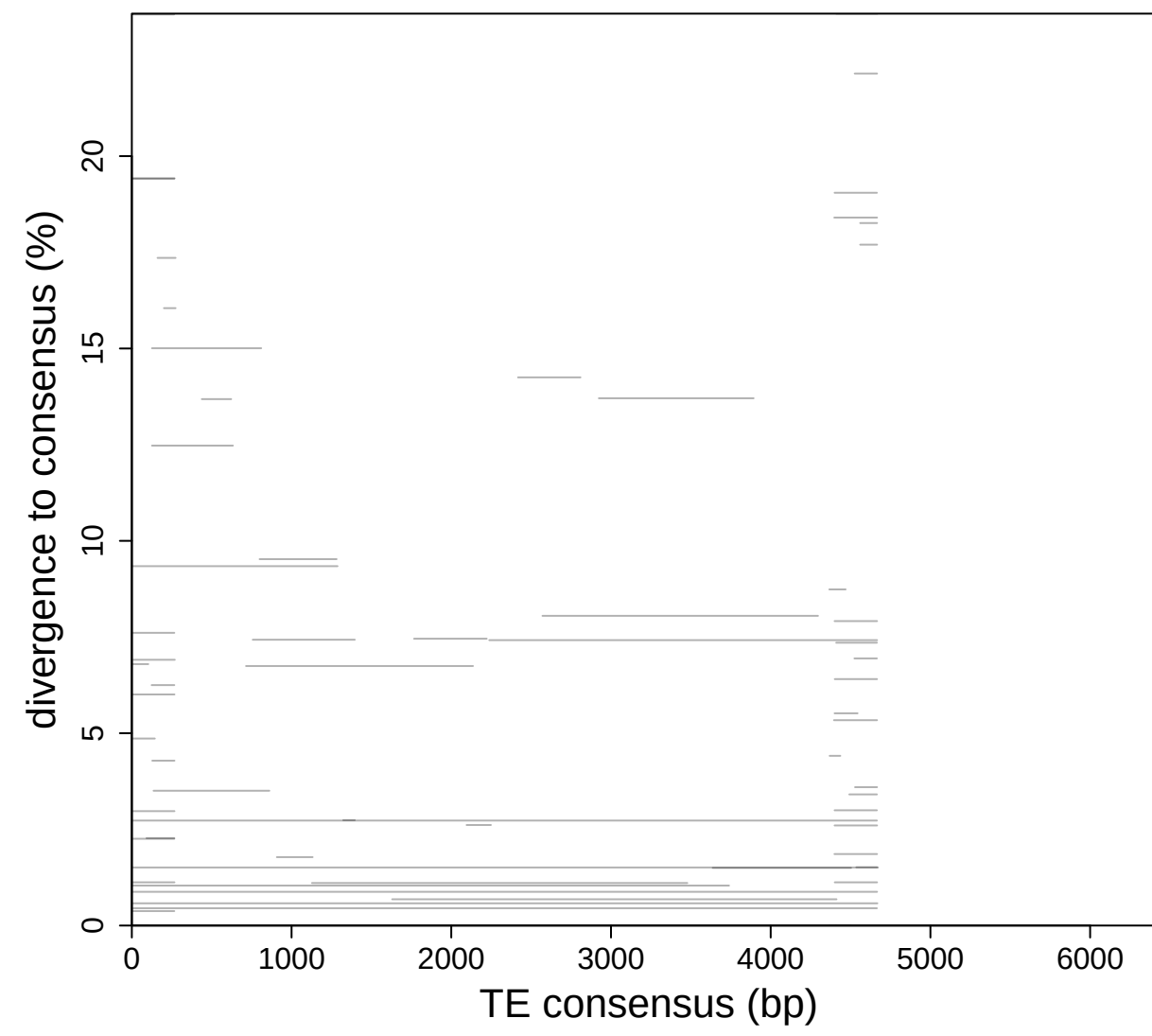
Start crop Point

End crop Point



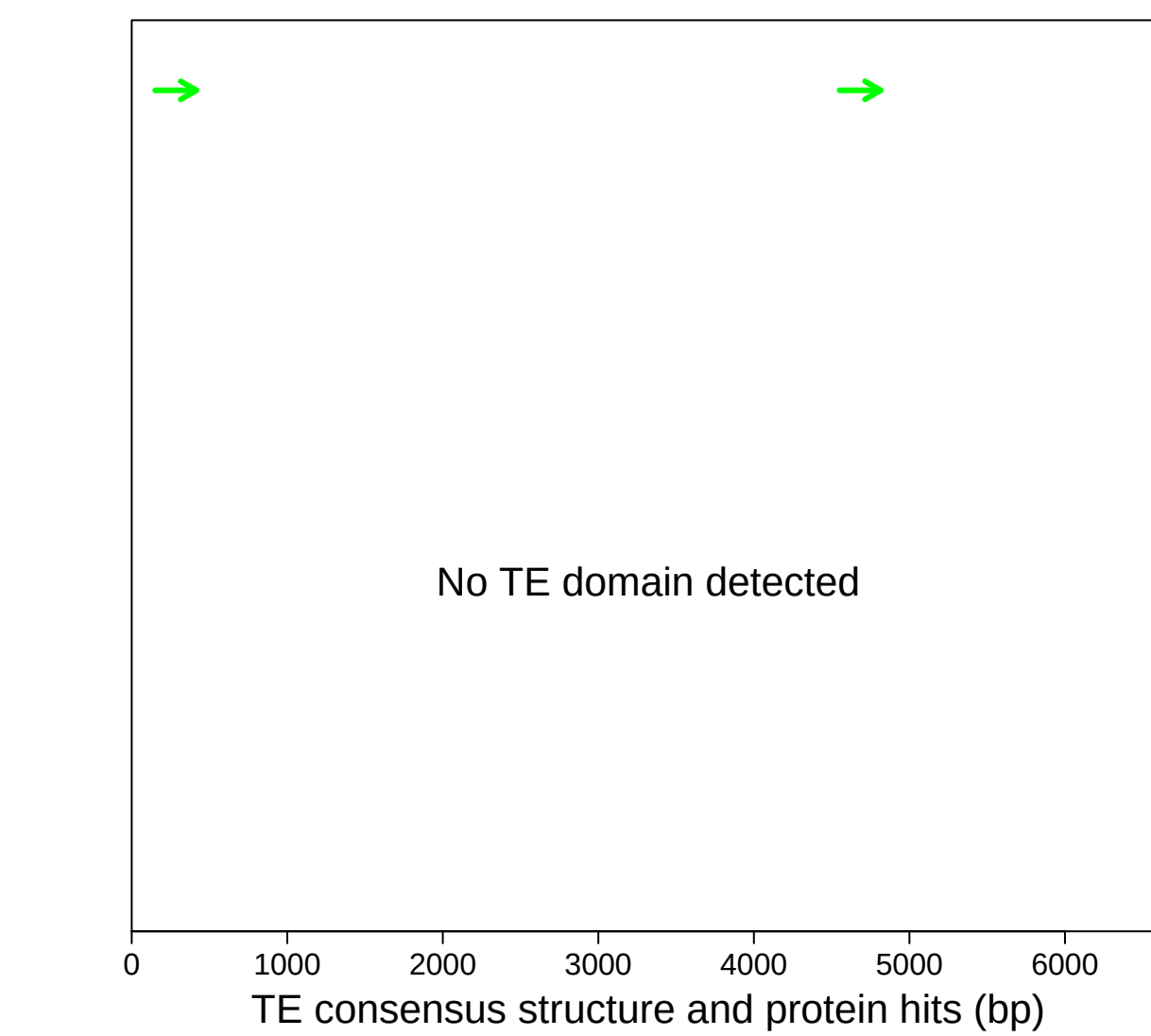
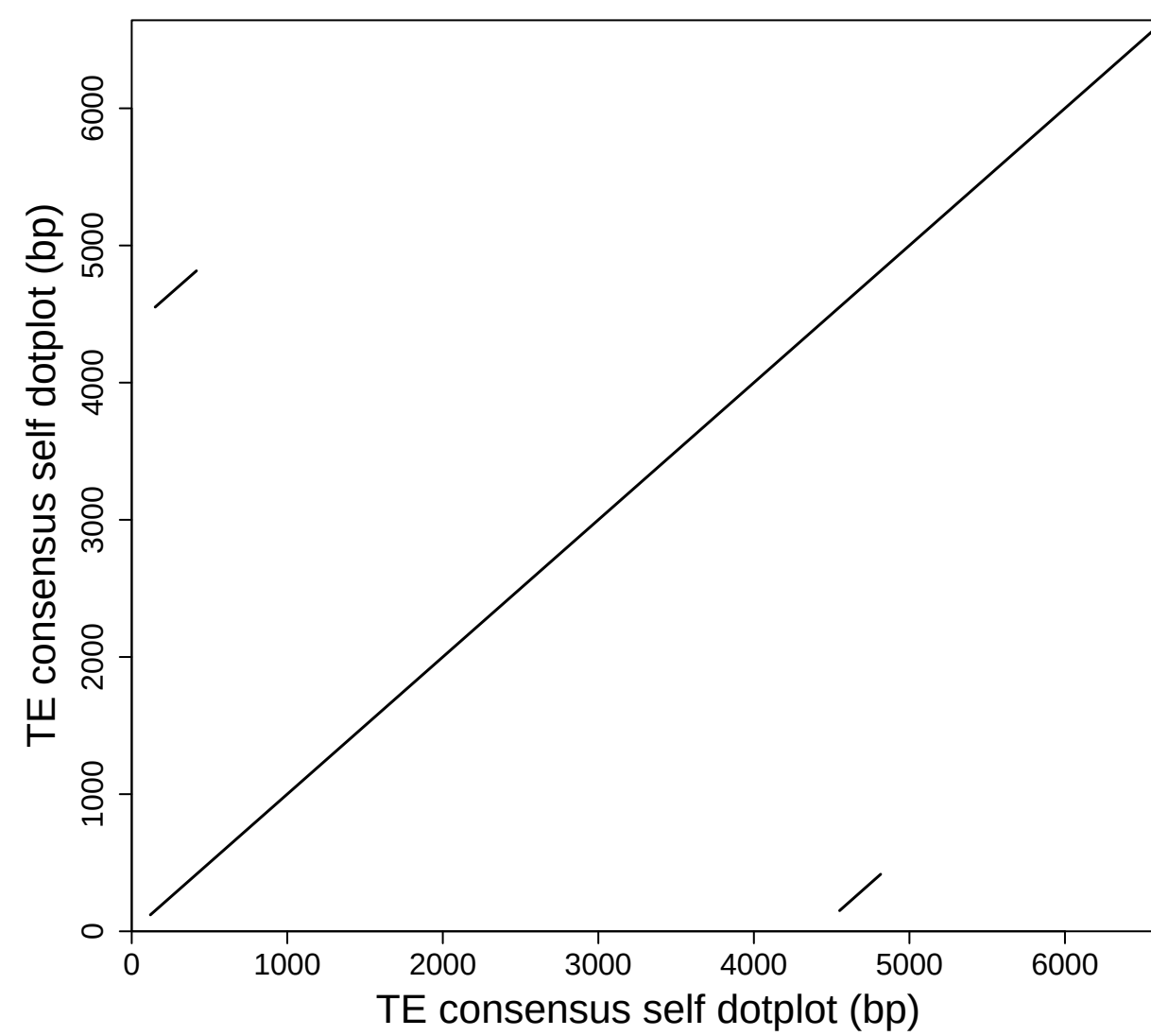
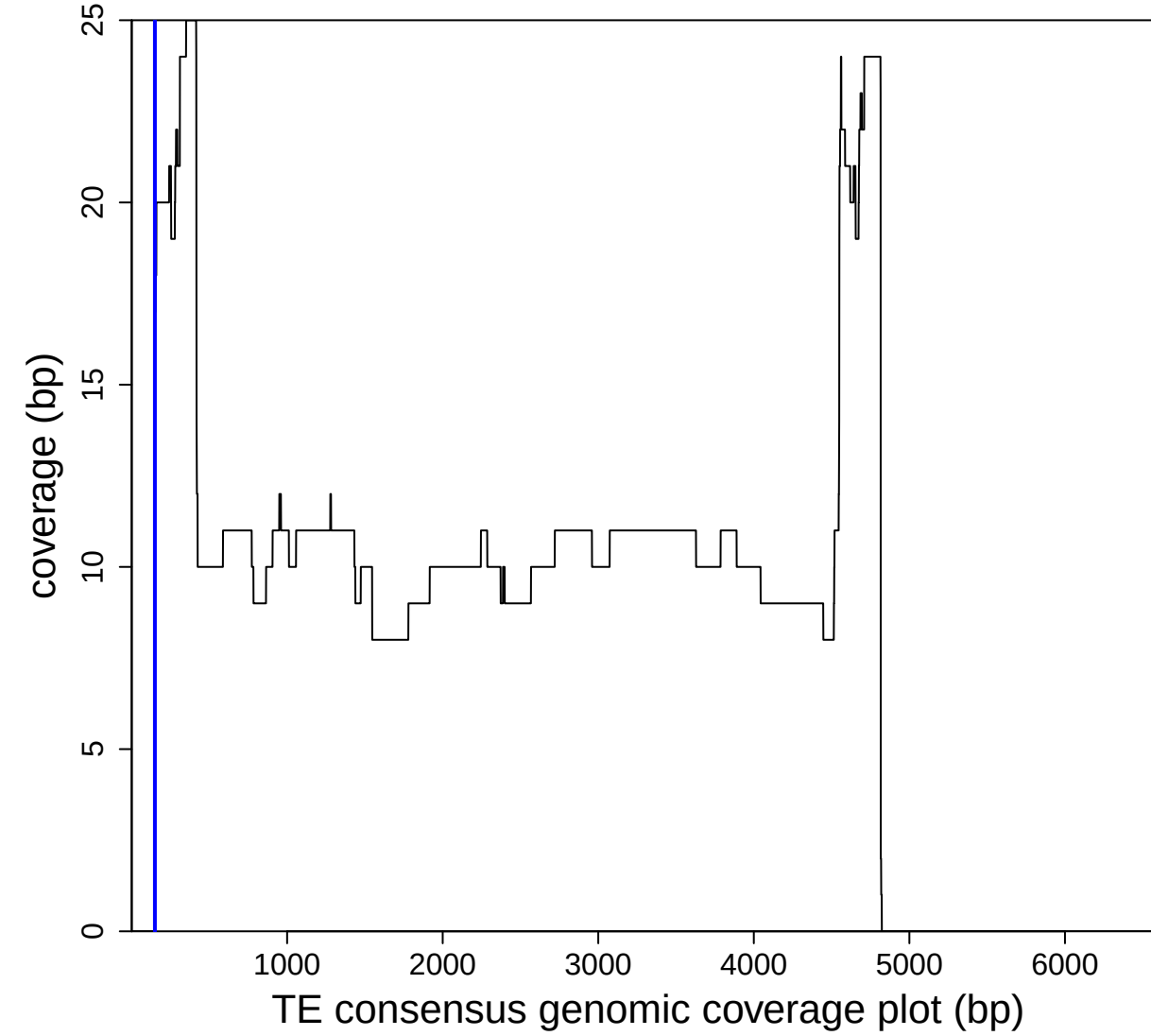
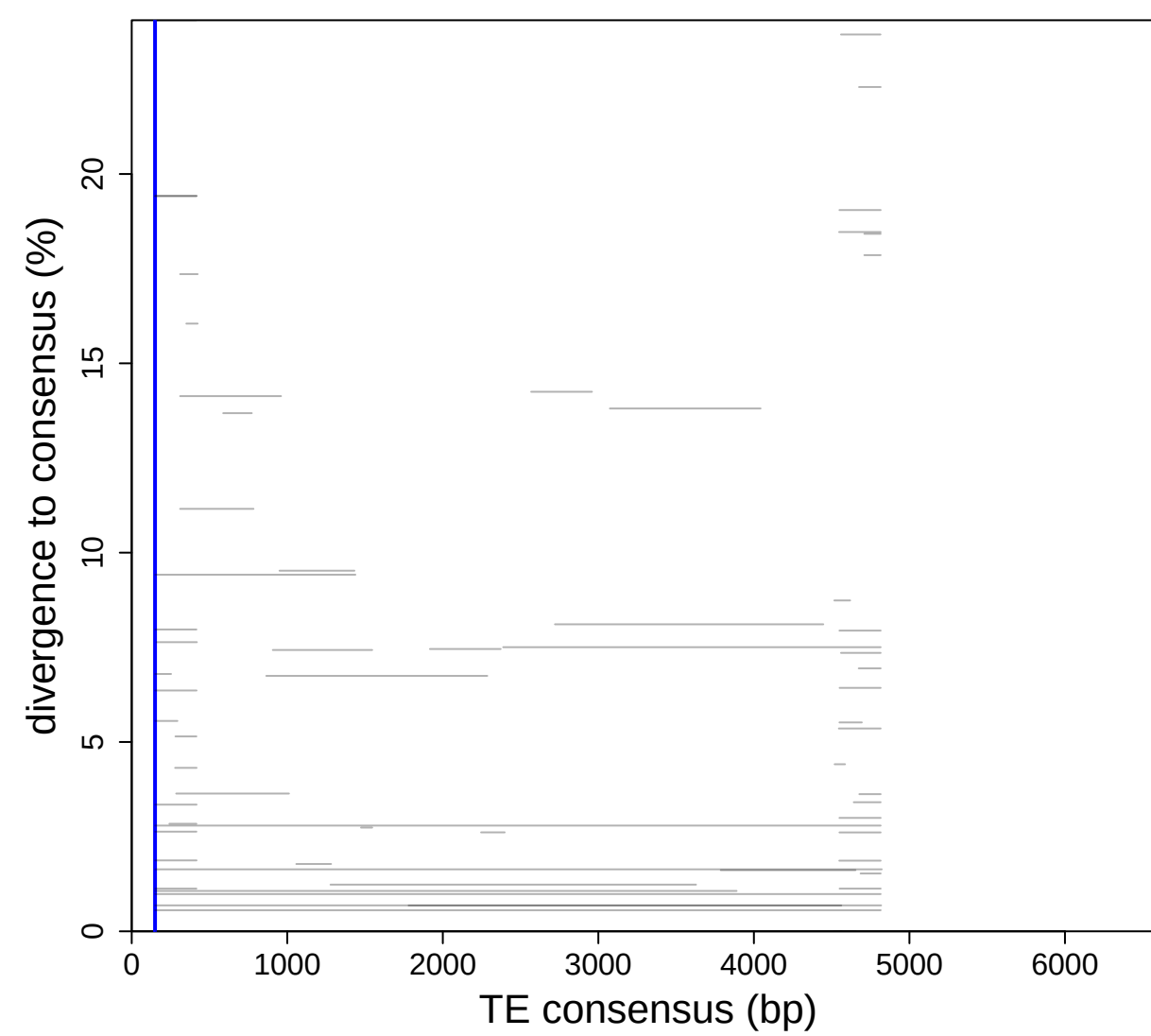
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_al
size: 6474bp; fragments: 66; full length: 0 (>=5826.6bp)

After TEtrimmer 6474 bp

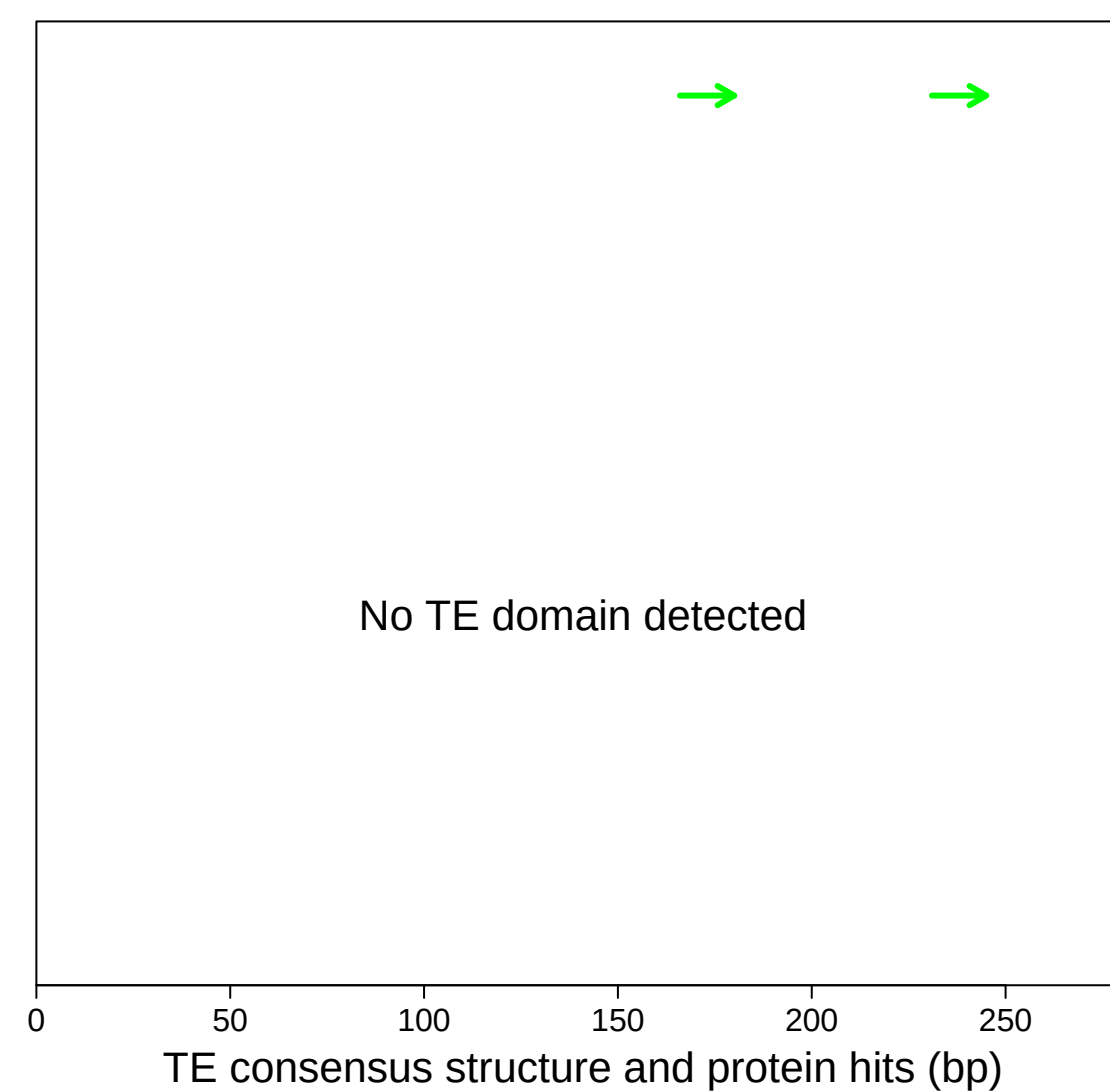
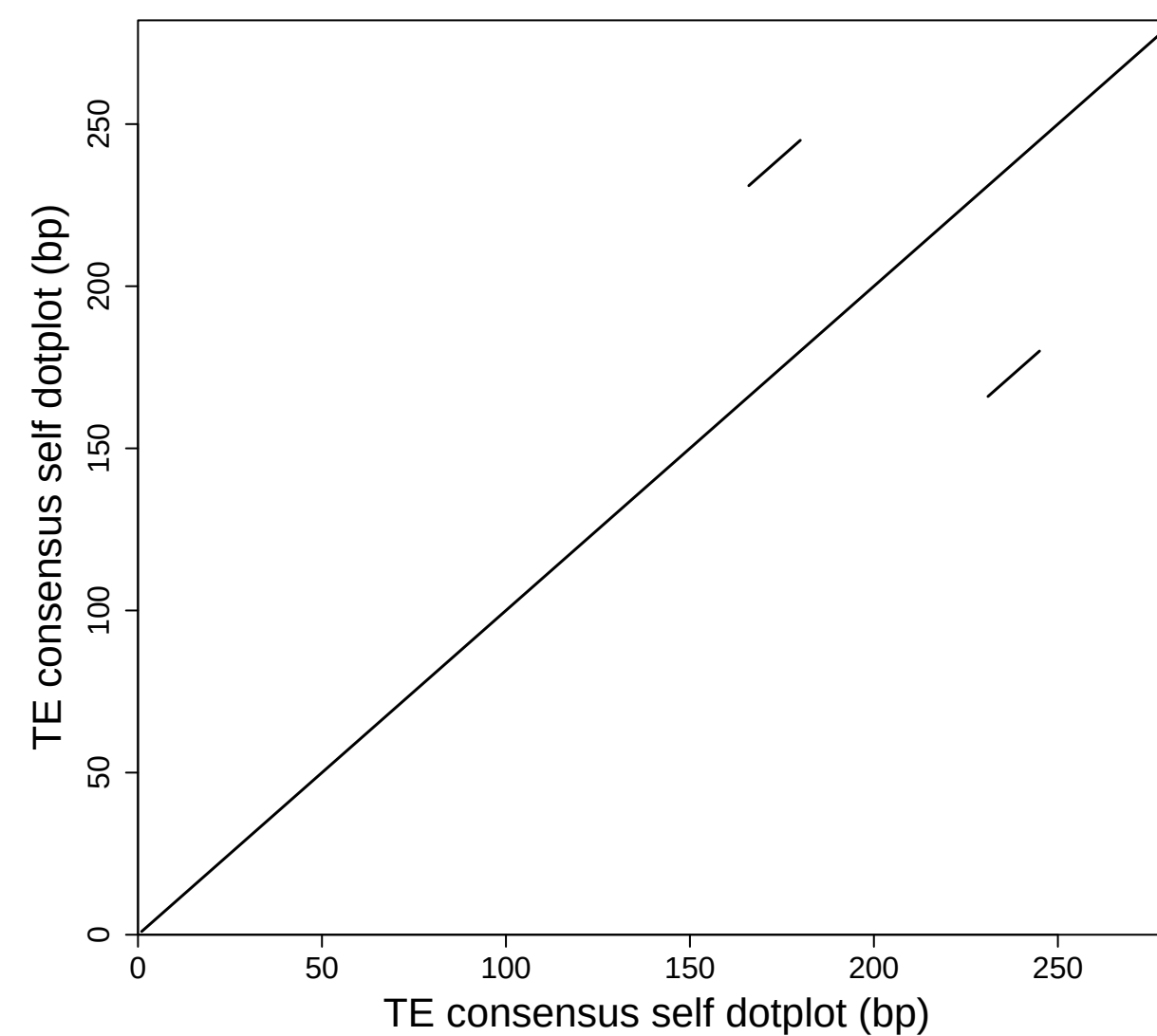
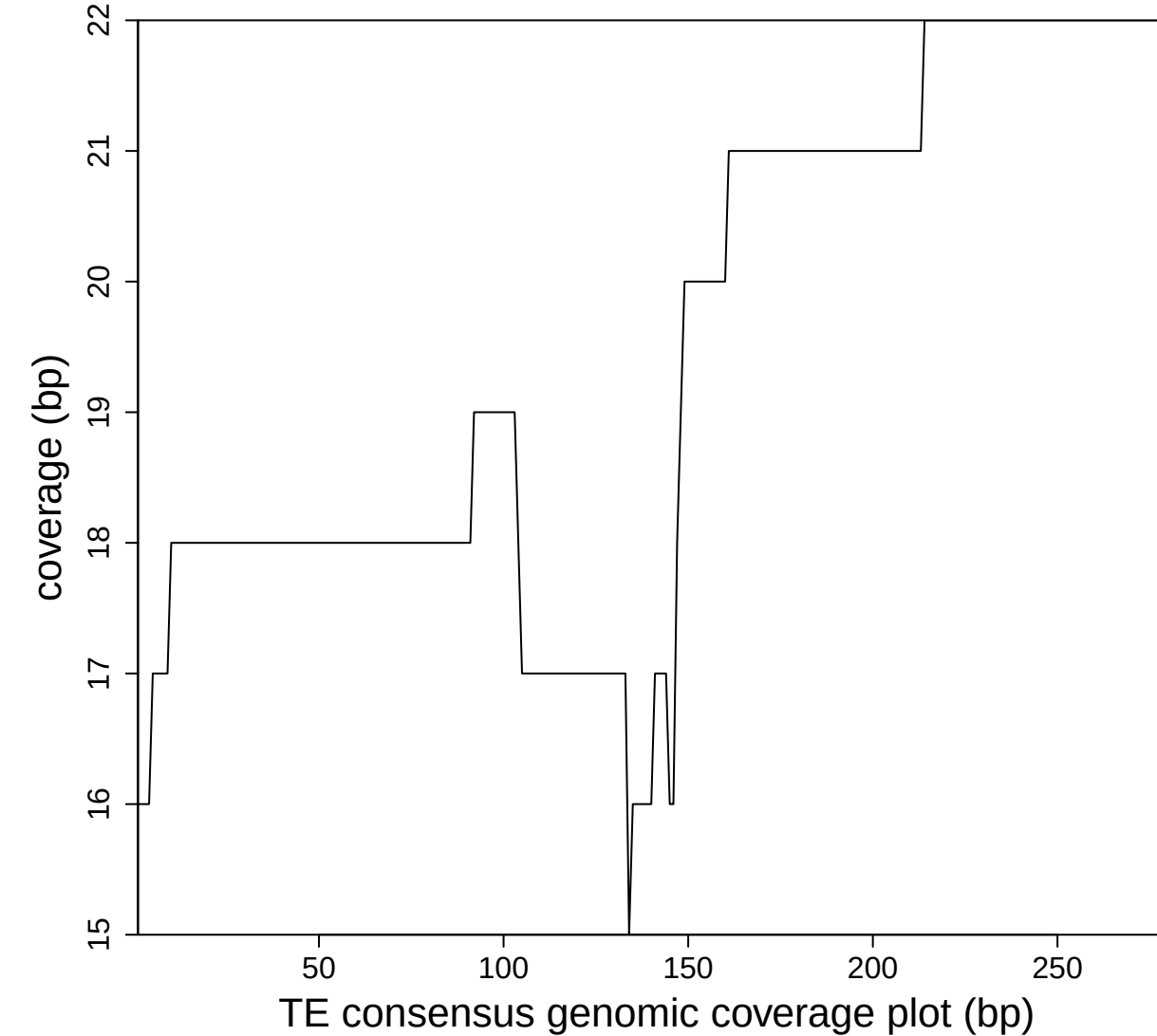
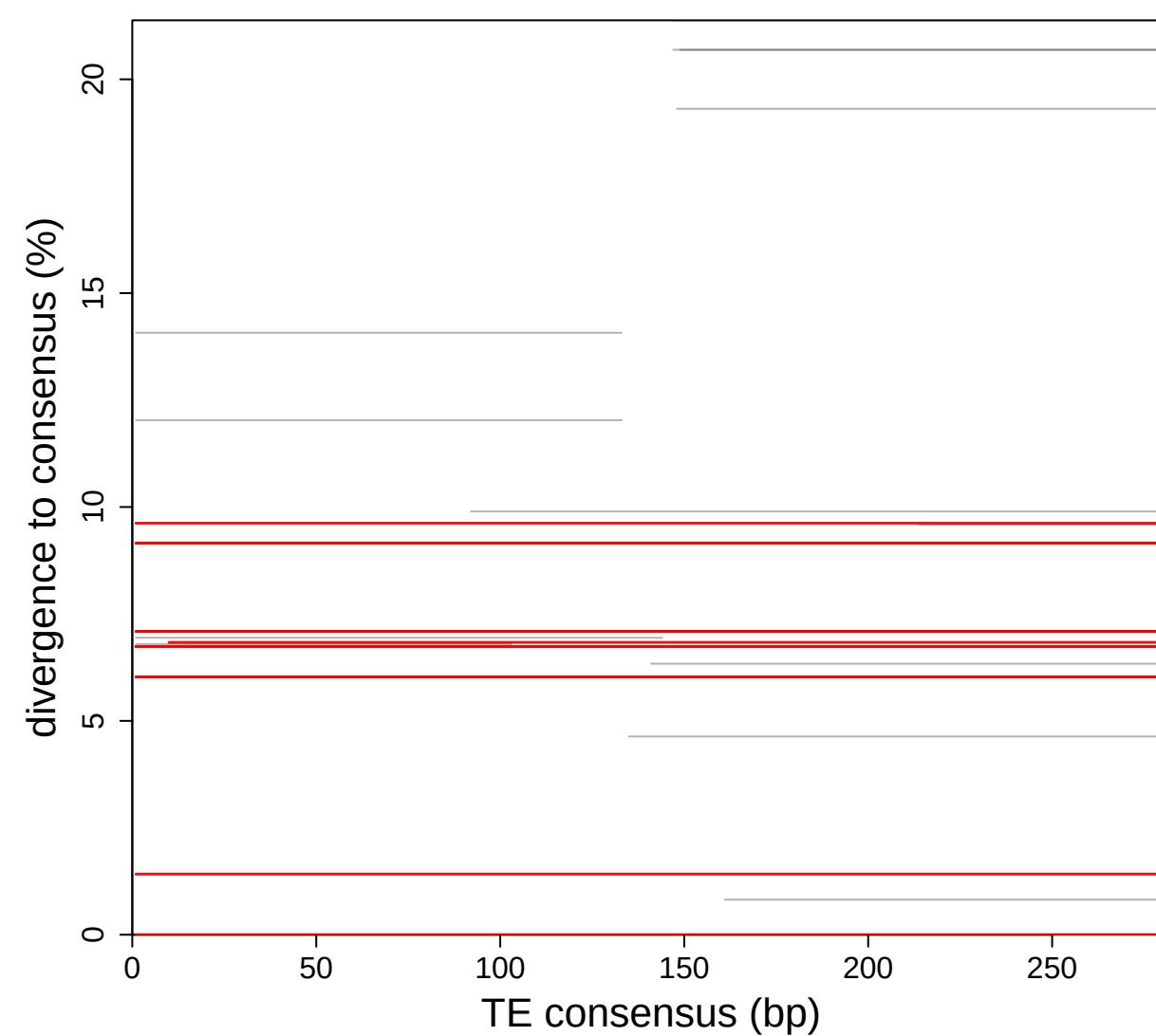


bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 6642bp; fragments: 66; full length: 0 (>=5977.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

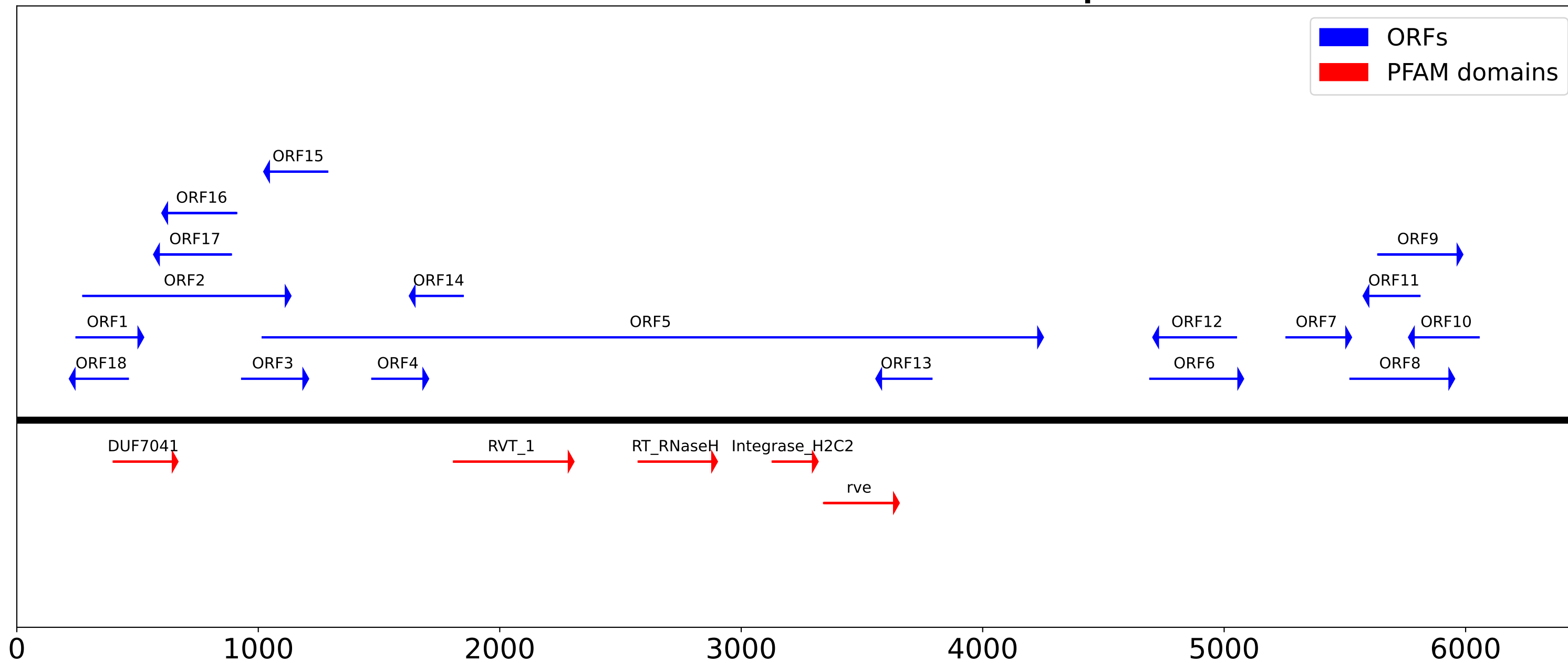


TE: ltr_1_family_267
size: 282bp; fragments: 27; full length: 13 (≥ 253.8 bp)



Before TEtrimmer 282 bp

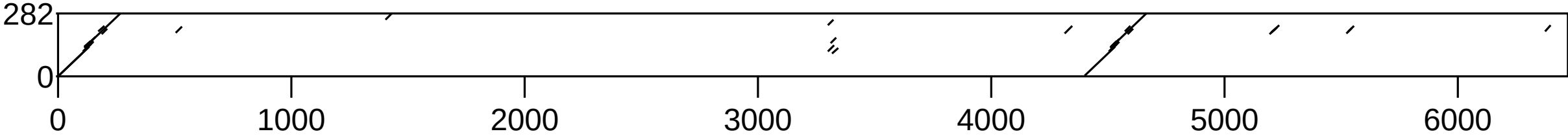
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)