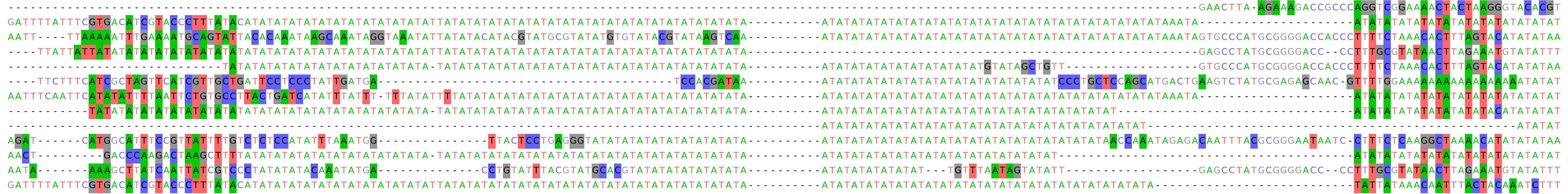
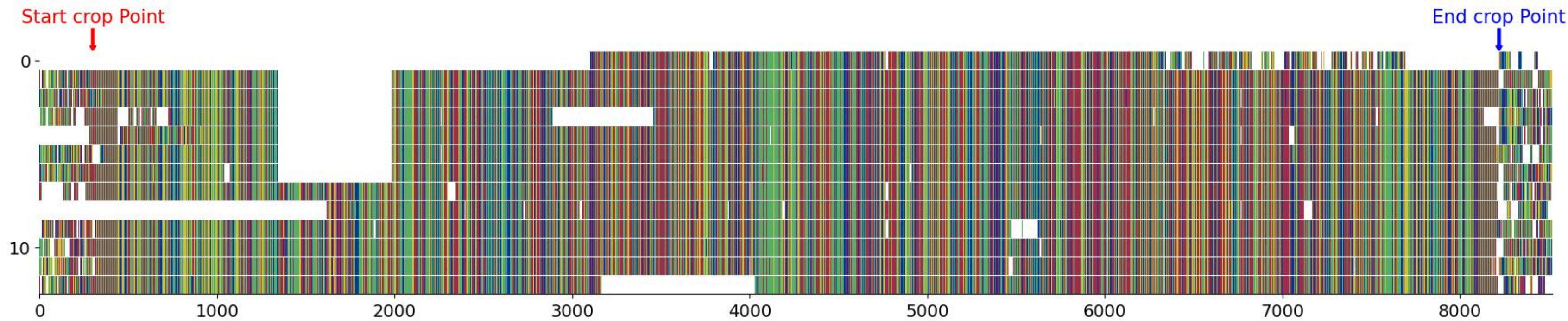


End crop Point

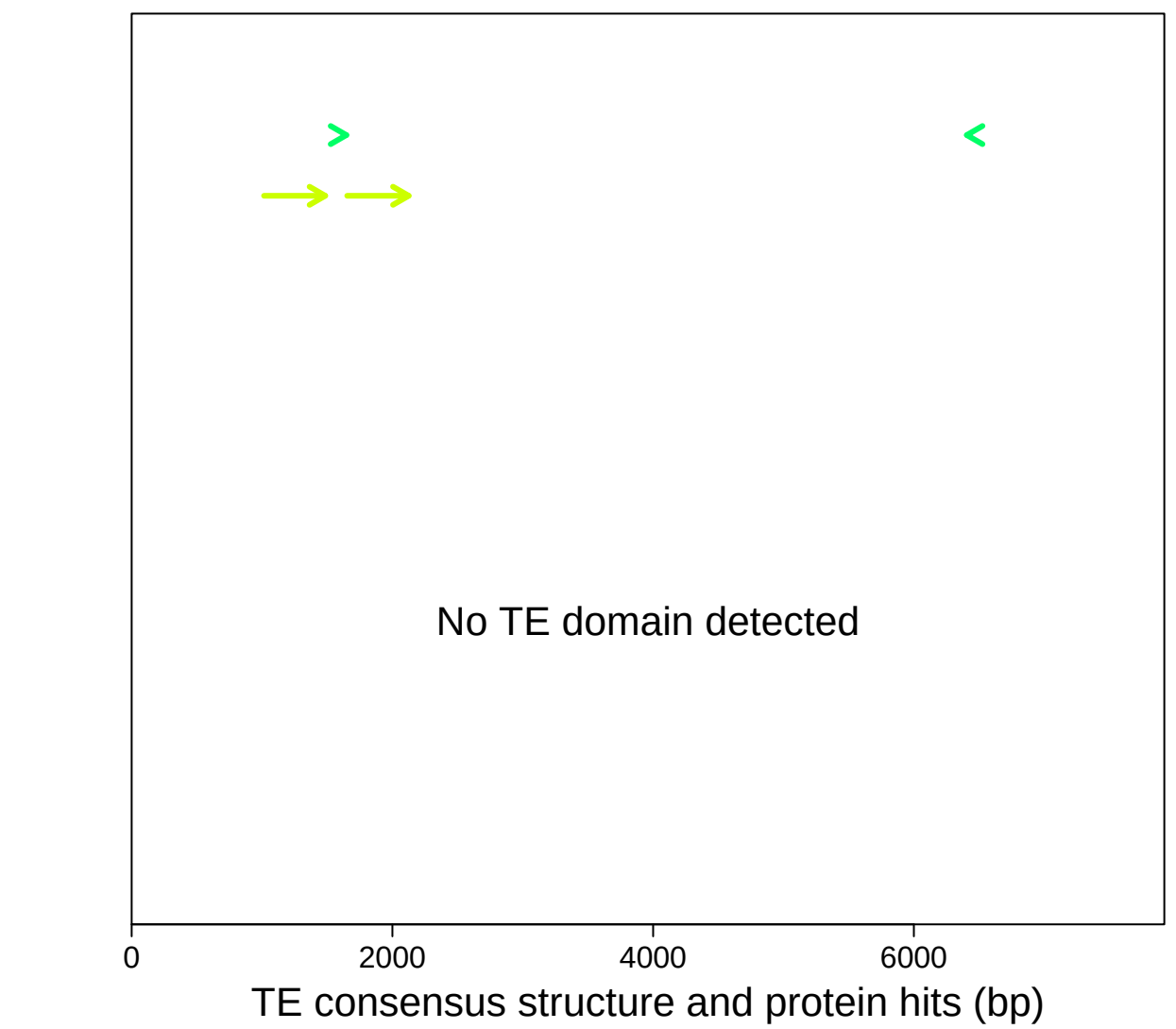
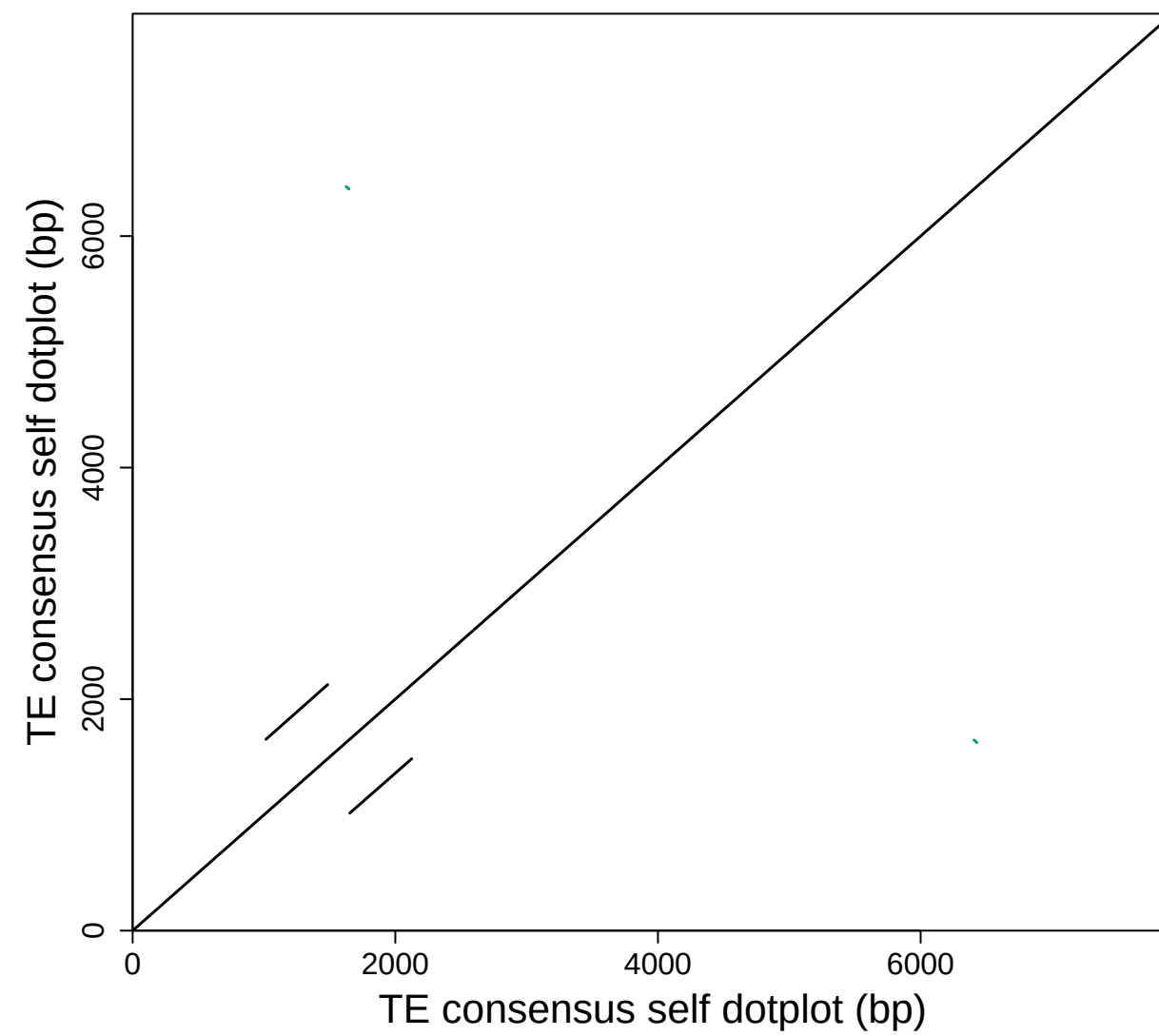
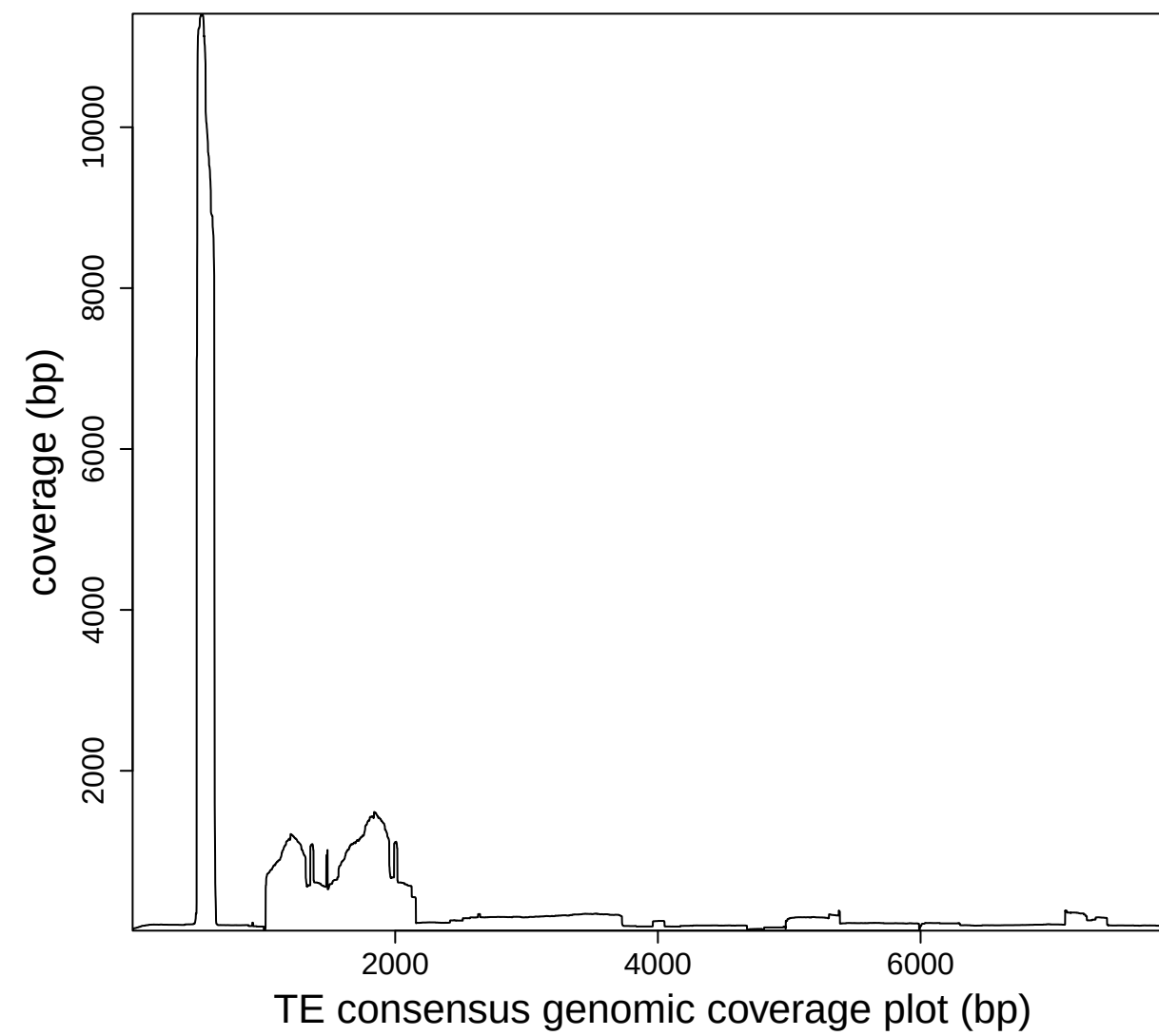
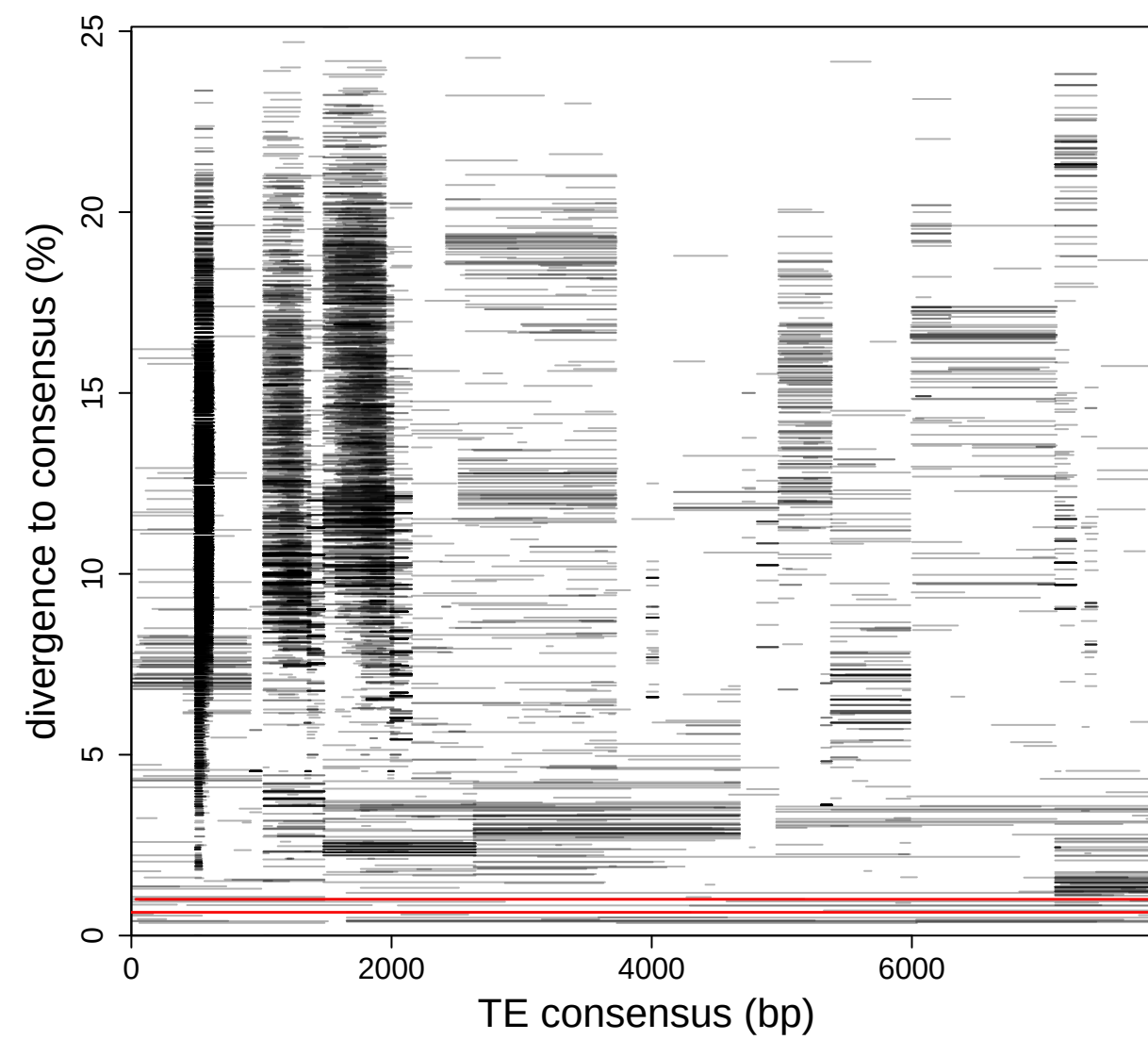
MSA length = 7921





_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa al
size: 7921bp; fragments: 16819; full length: 2 (>=7128.9bp)

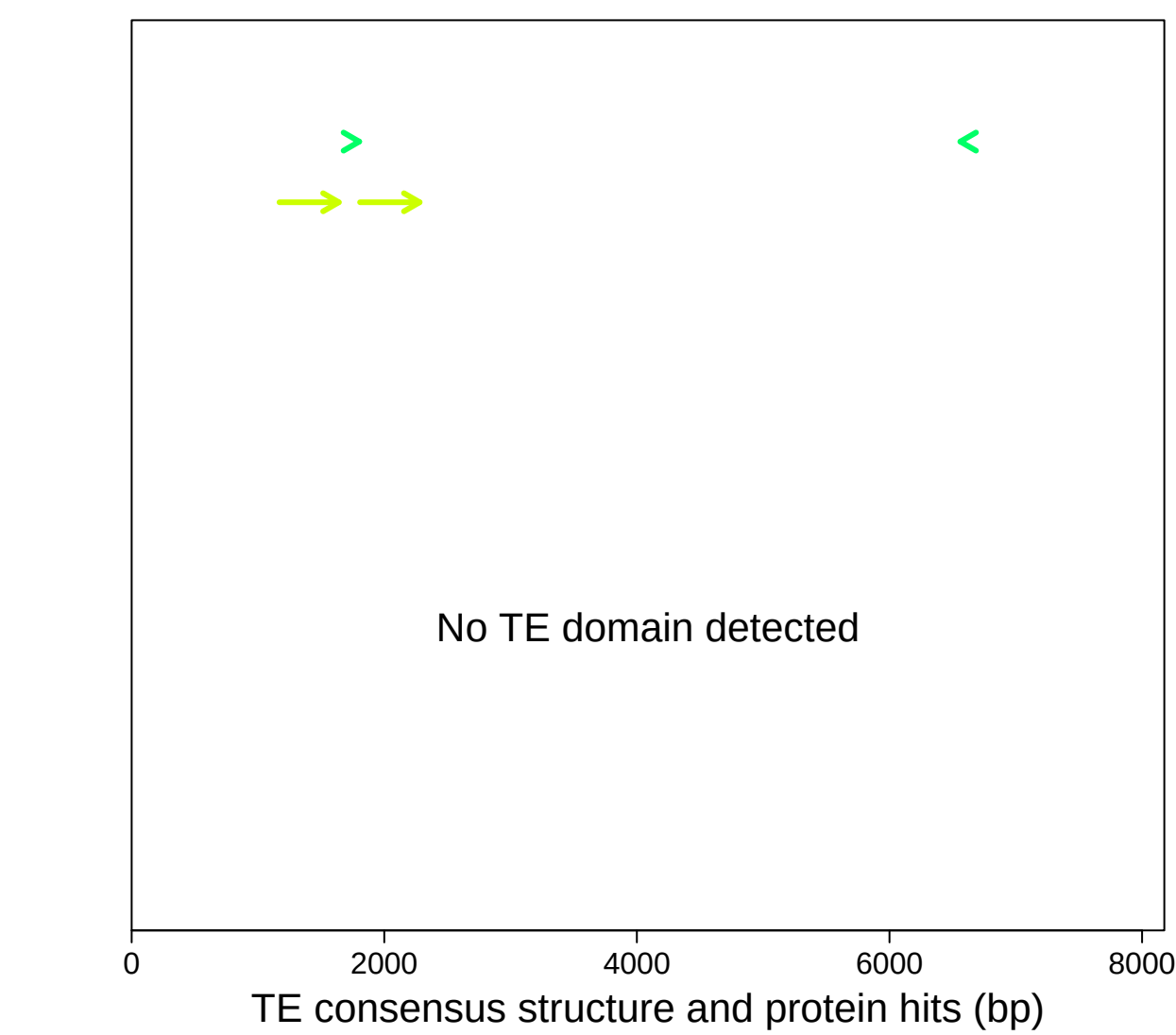
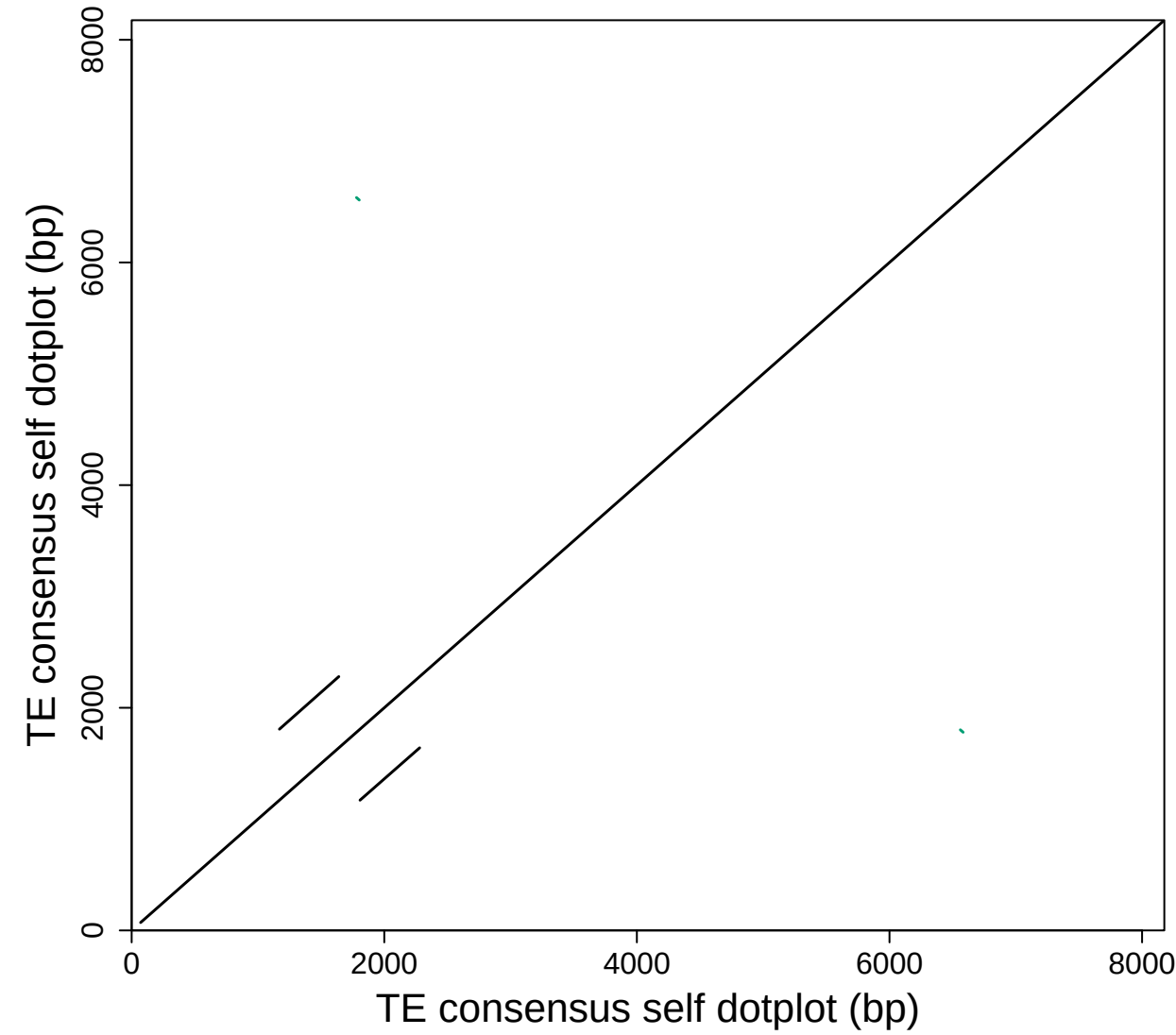
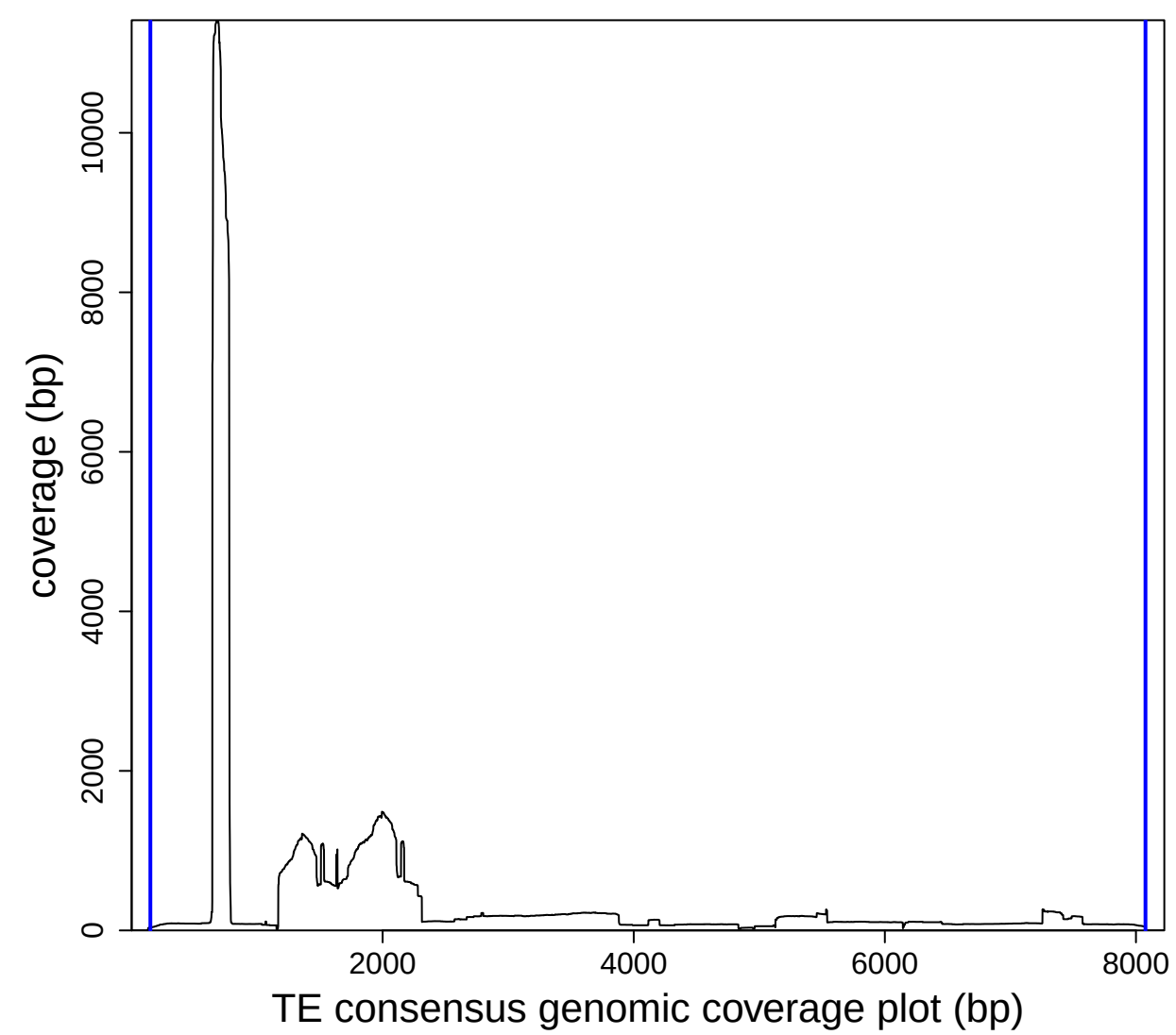
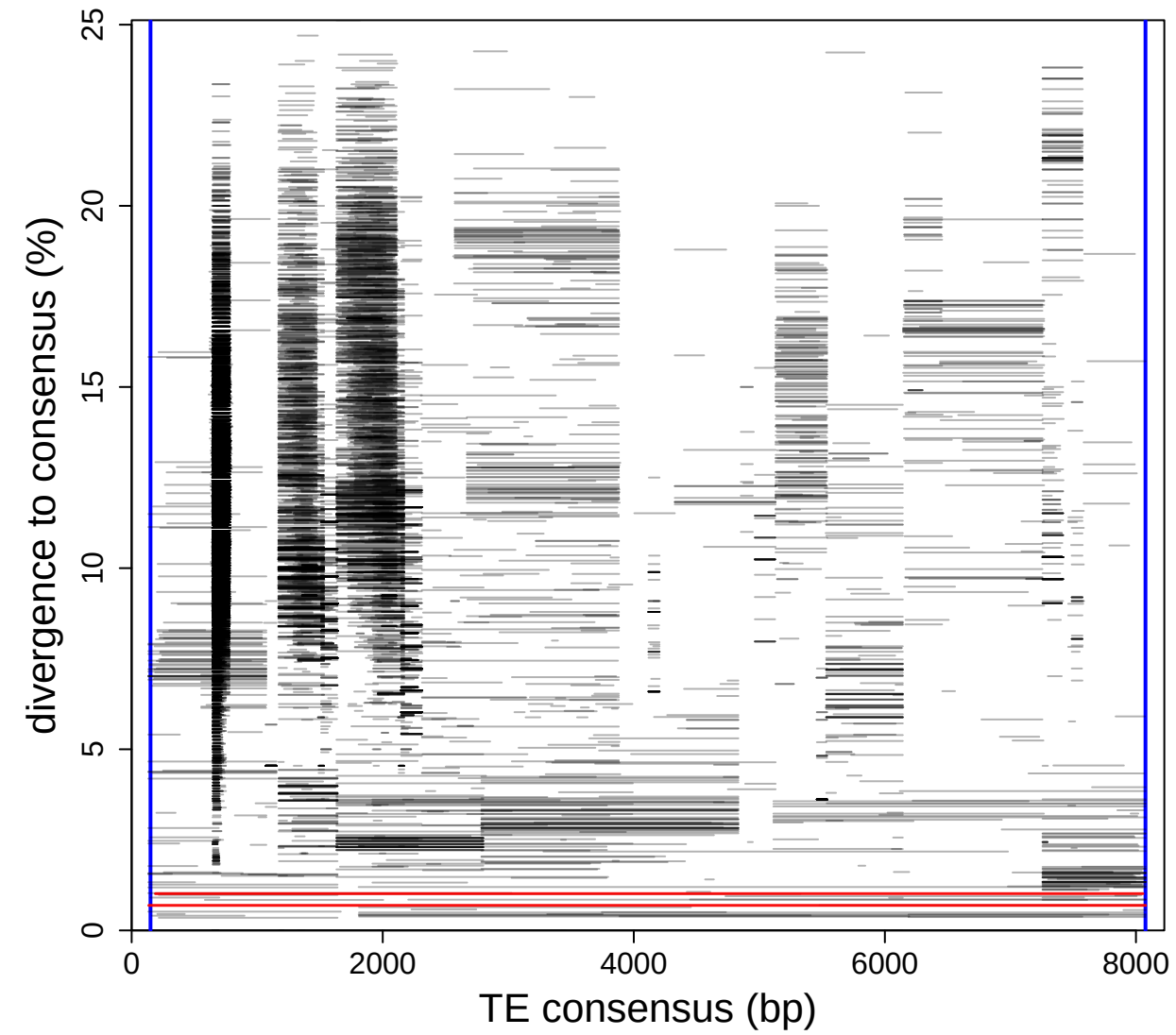
After TEtrimmer 7921 bp



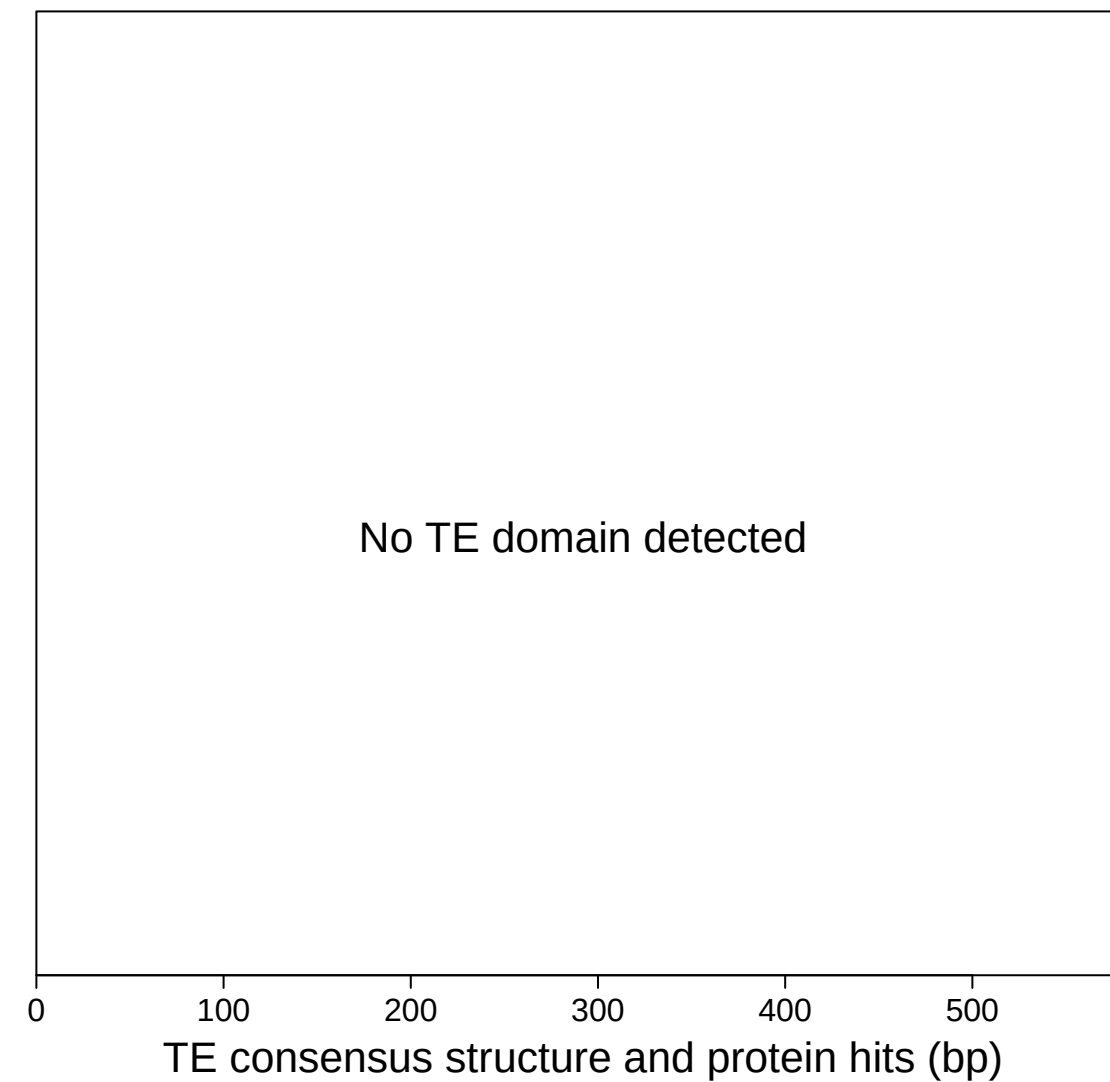
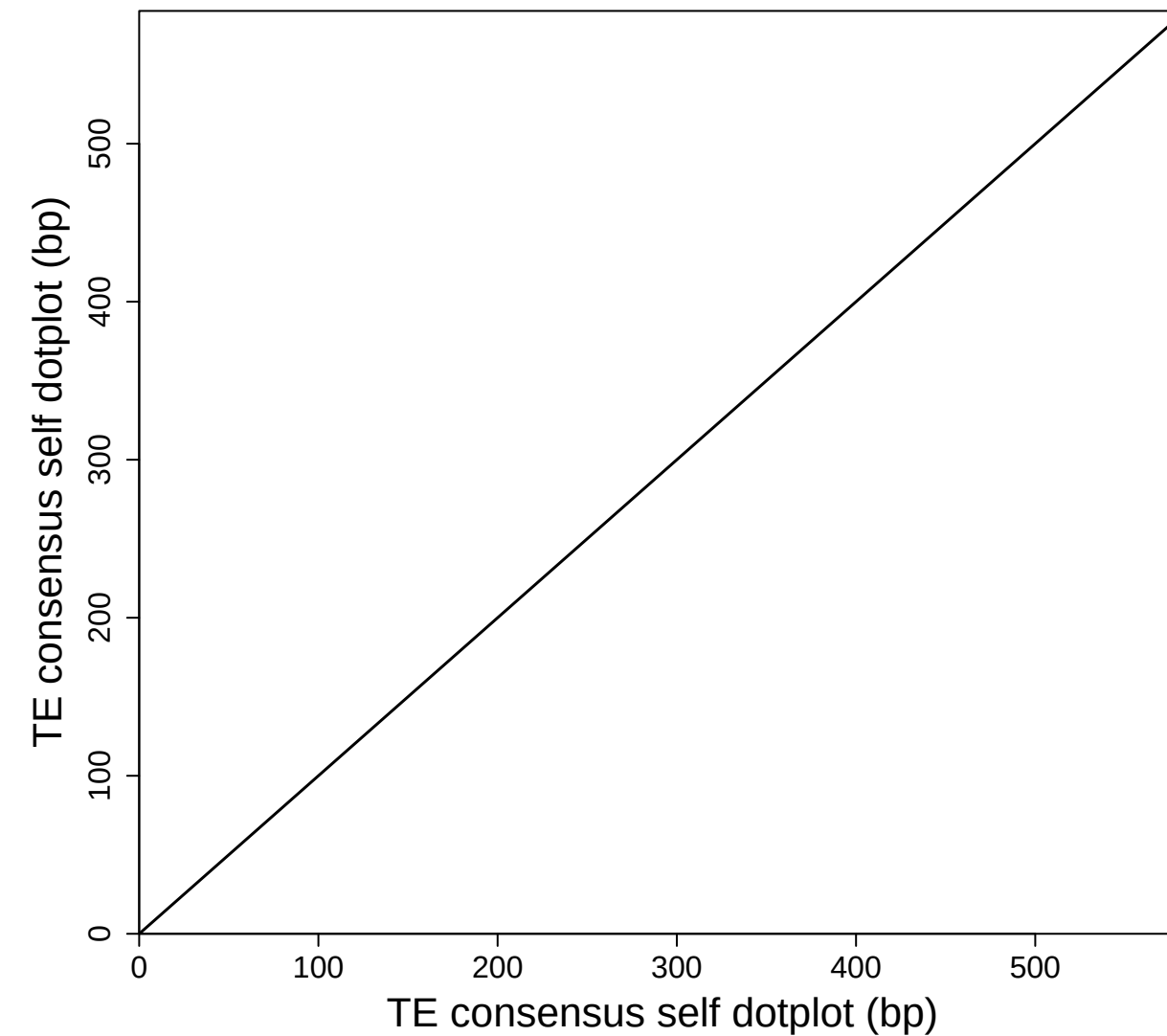
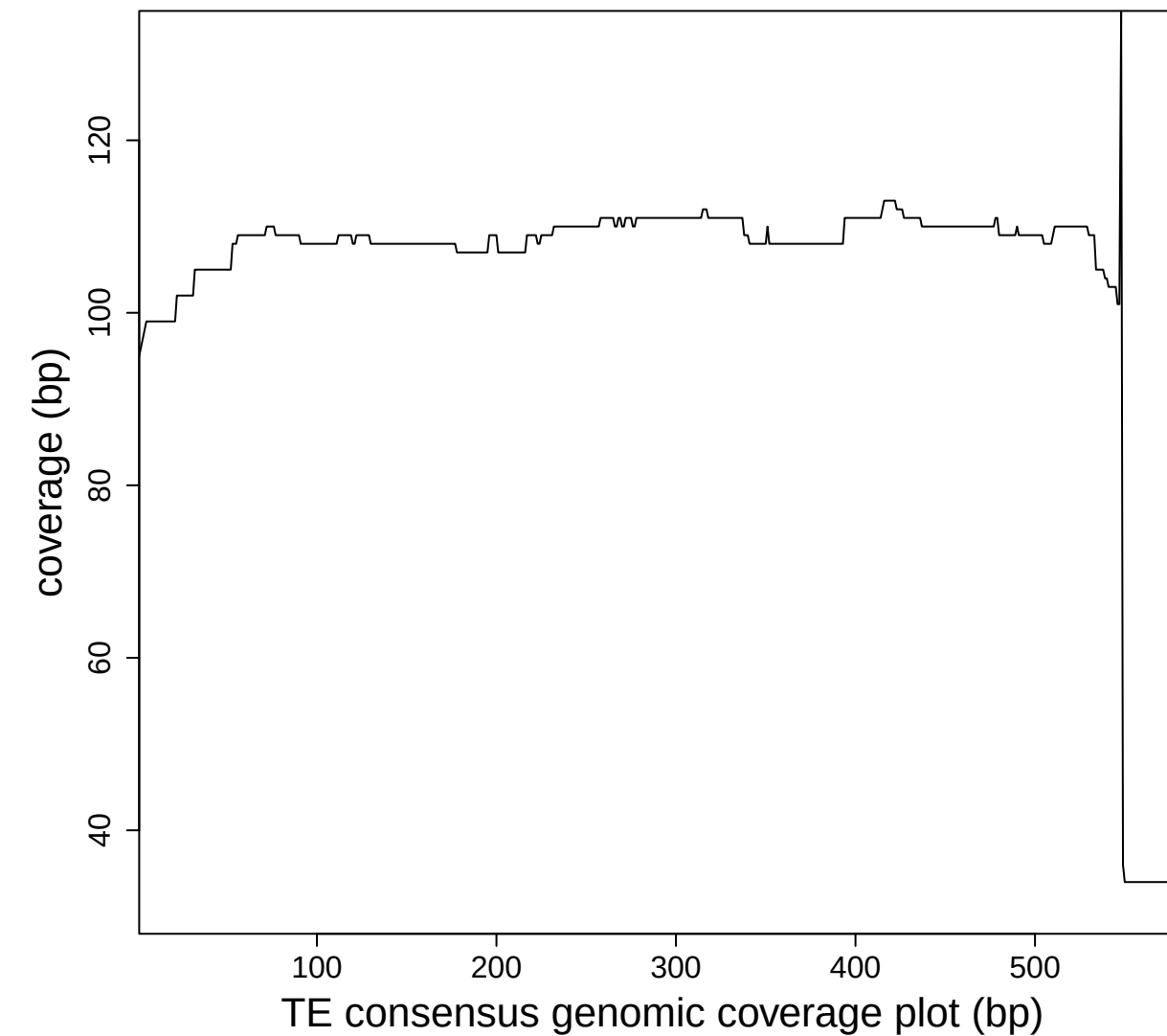
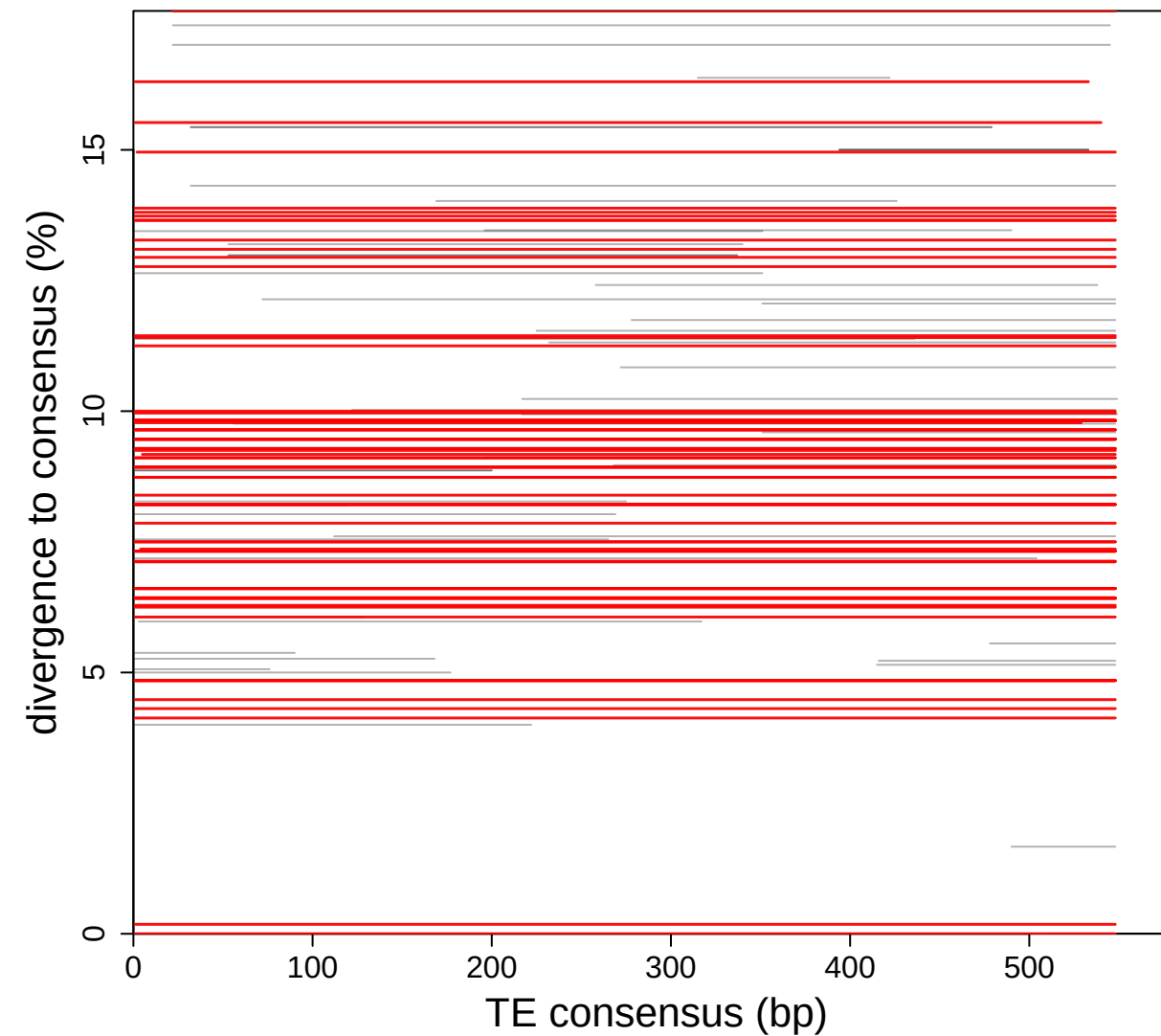
bed_g_2.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bclr
size: 8226bp; fragments: 16819; full length: 2 (>=7403.4bp)

After TEtrimmer Extended plot

Blue lines are boundaries

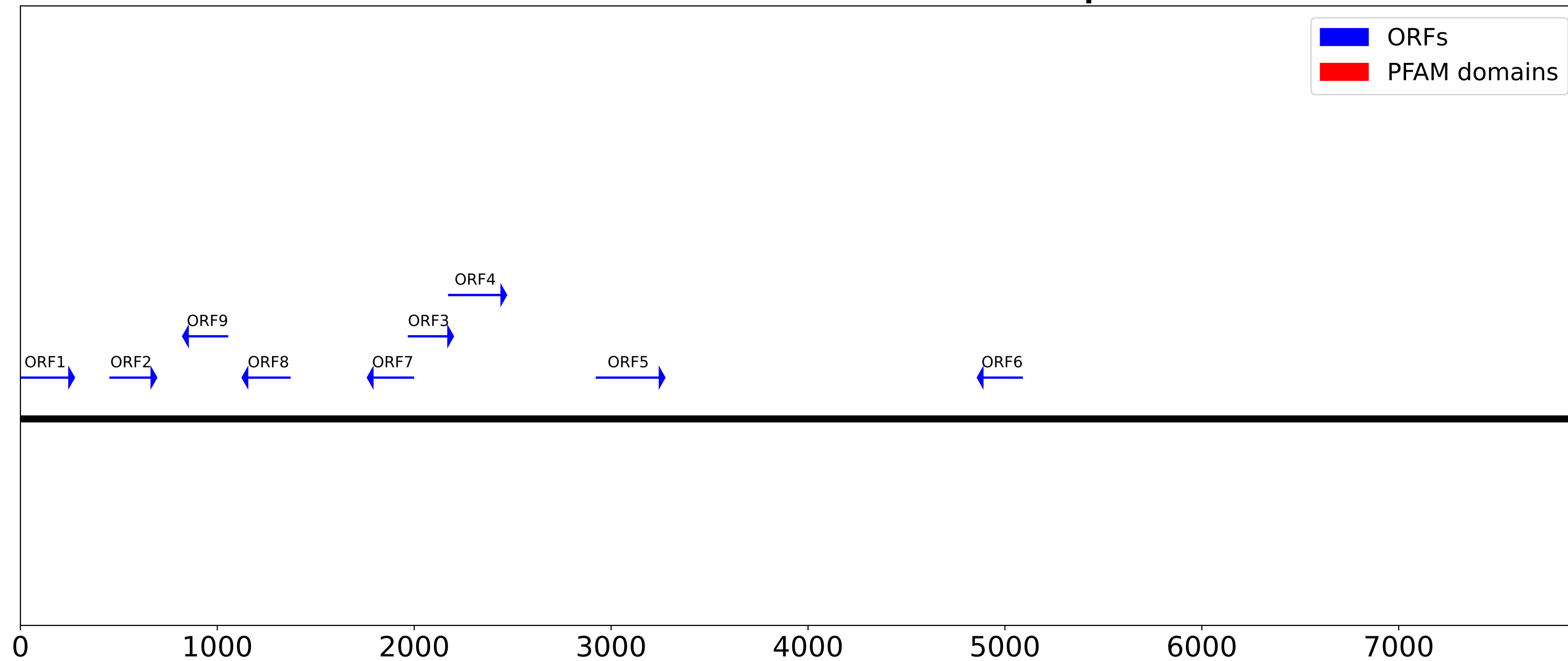


TE: ltr_1_family_354
size: 584bp; fragments: 169; full length: 83 (≥ 525.6 bp)



Before TEtrimmer 584 bp

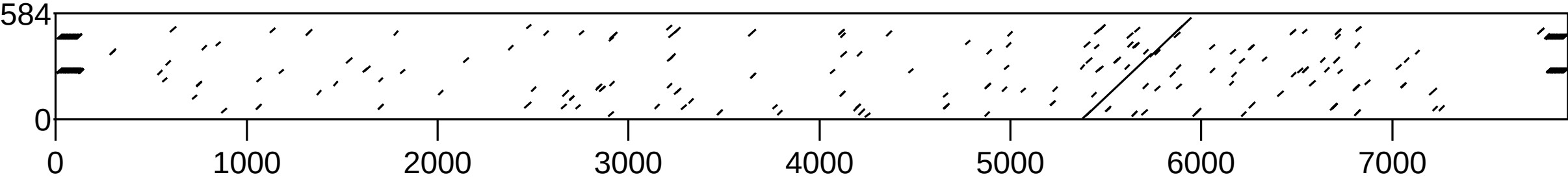
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)