

Start crop Point

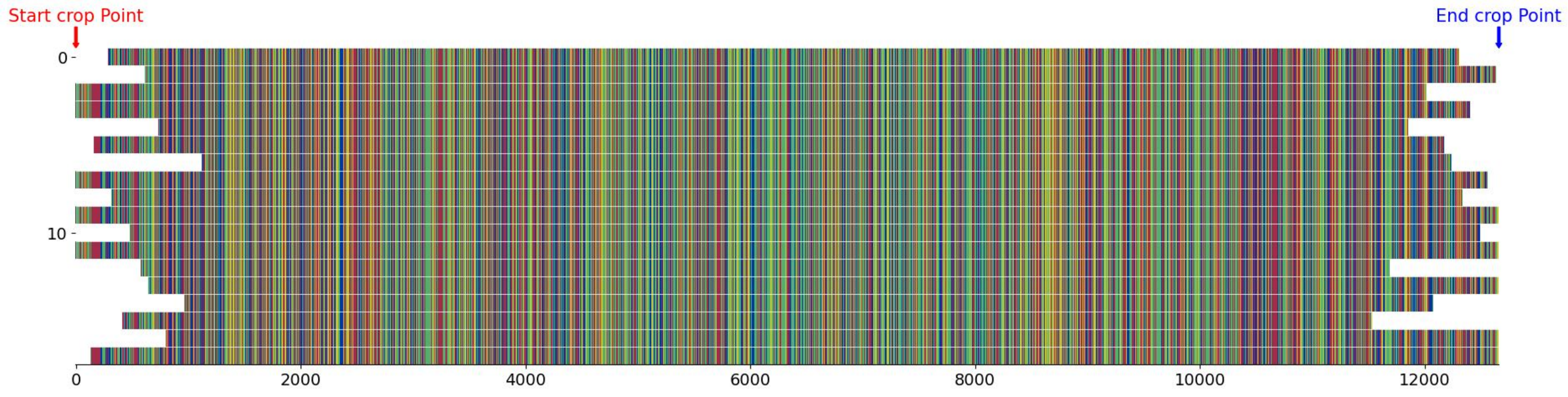


End crop Point



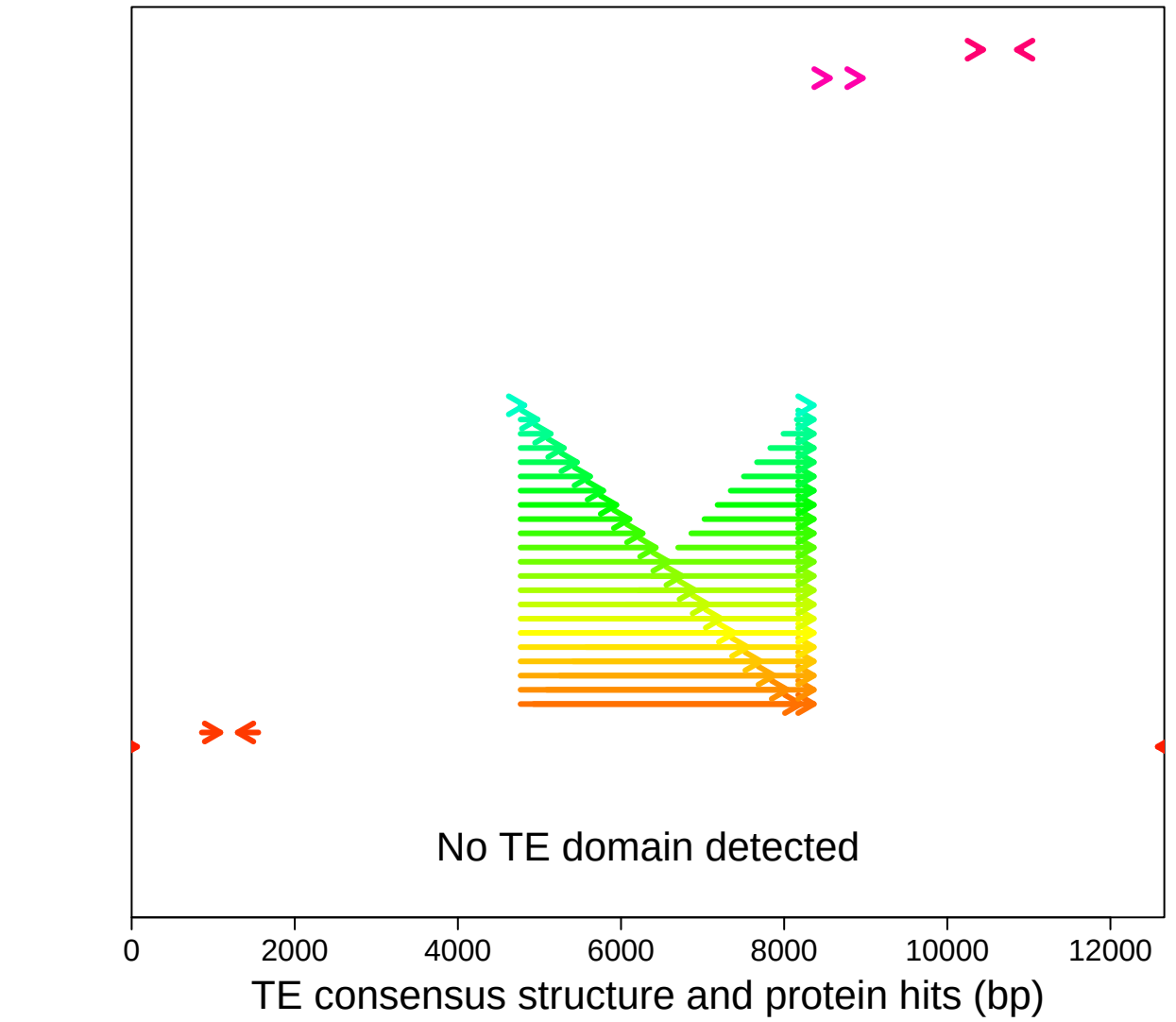
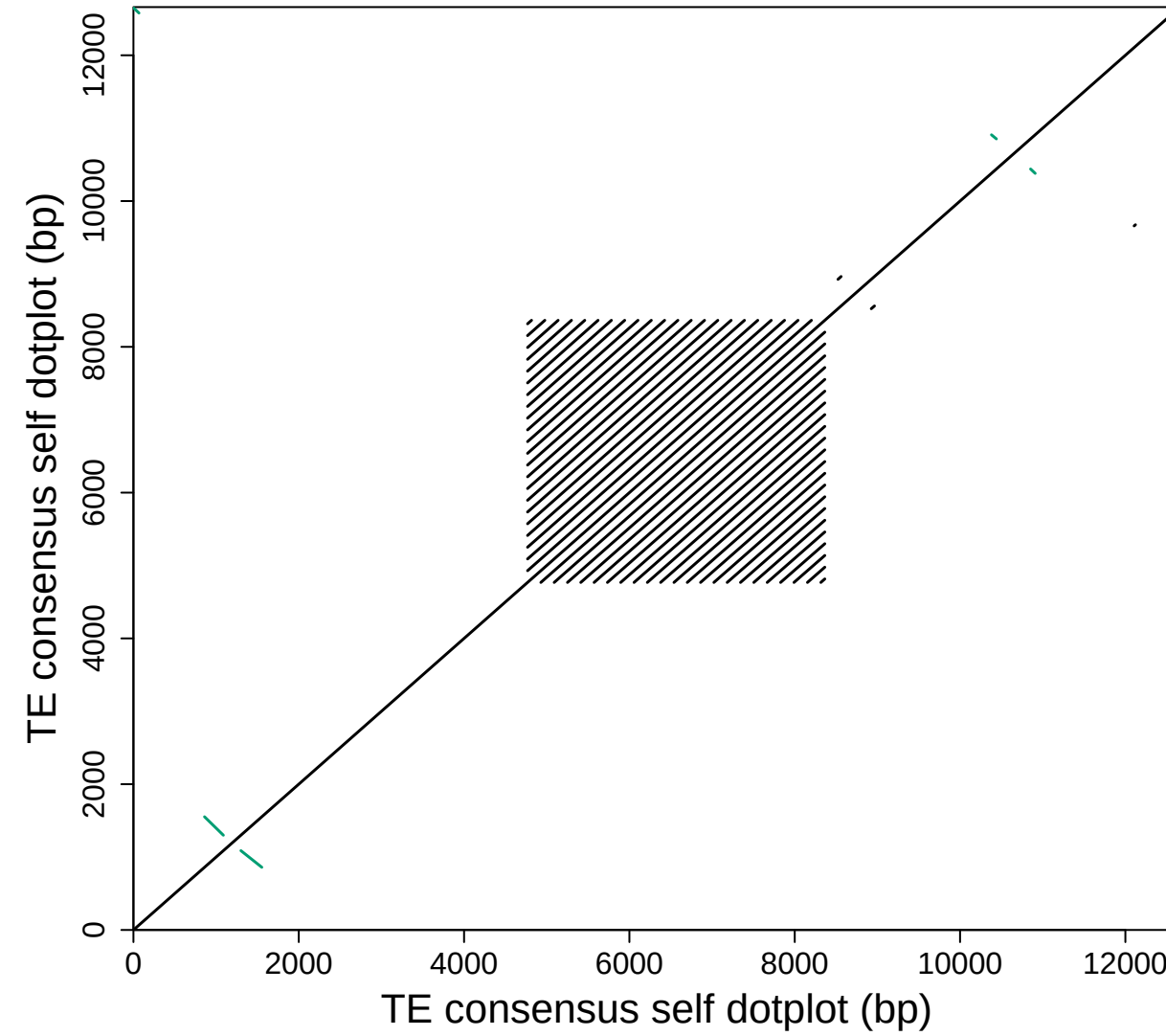
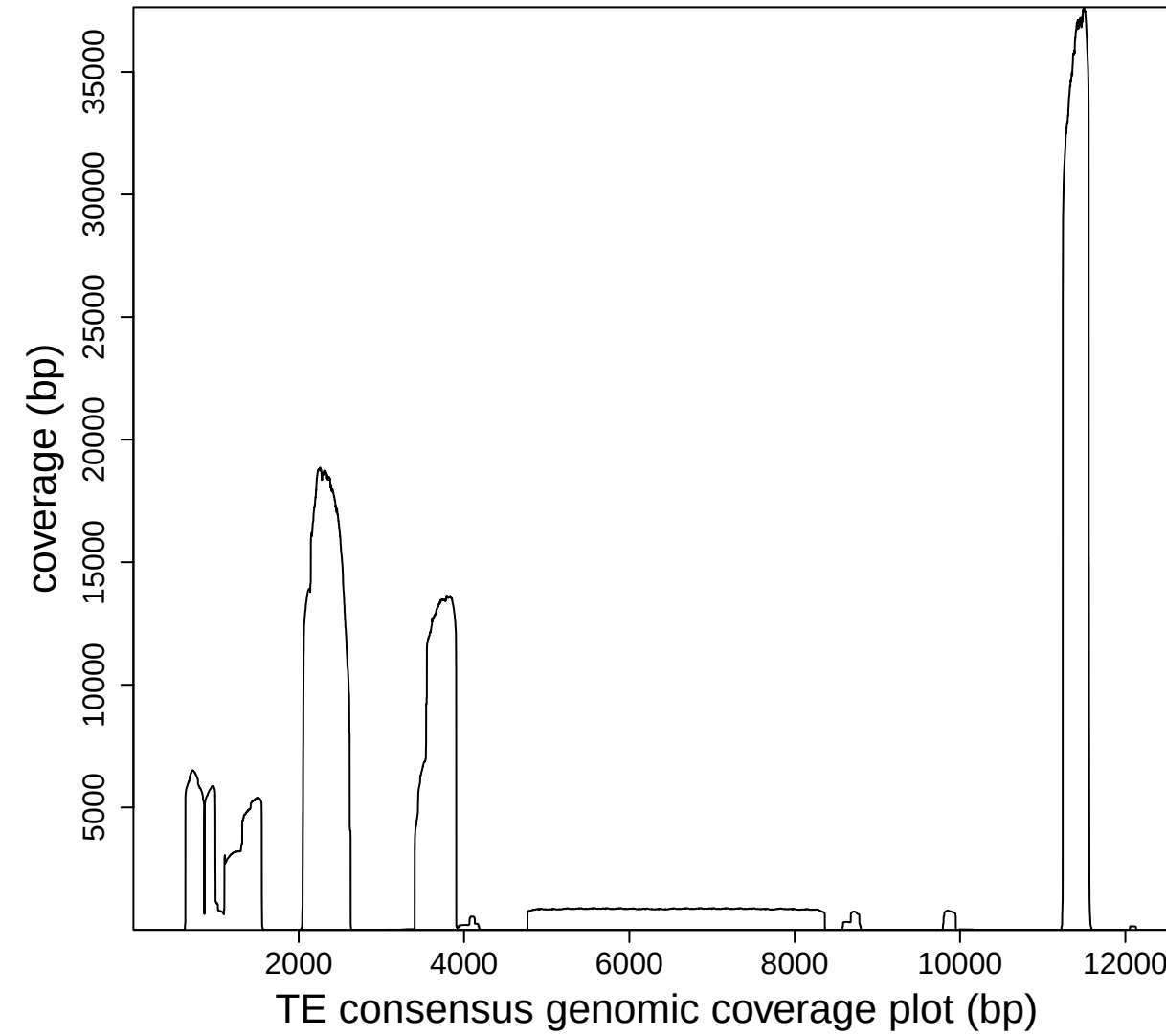
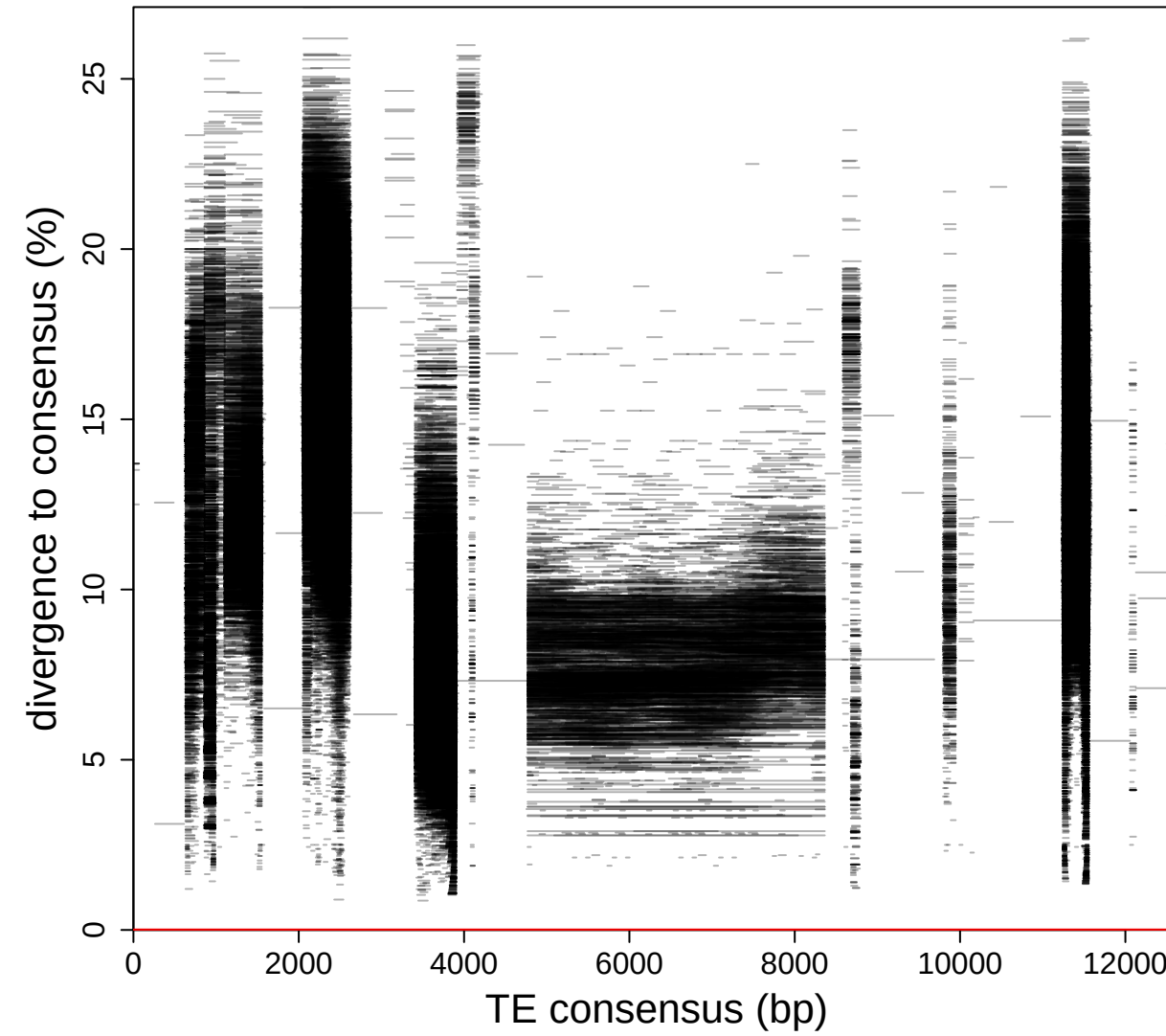
MSA length = 12661





_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa al
size: 12661bp; fragments: 111340; full length: 1 (>=11394.9bp)

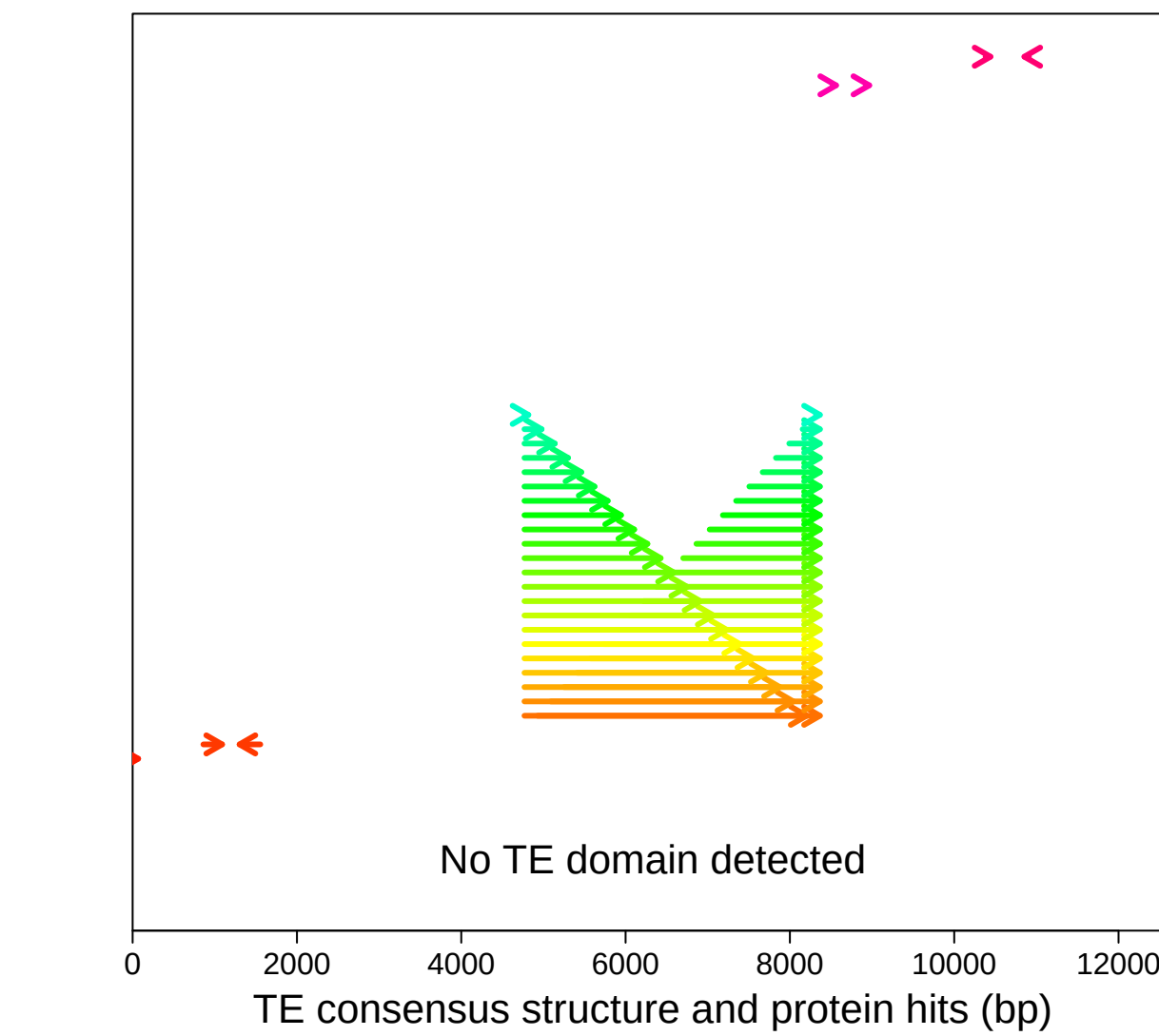
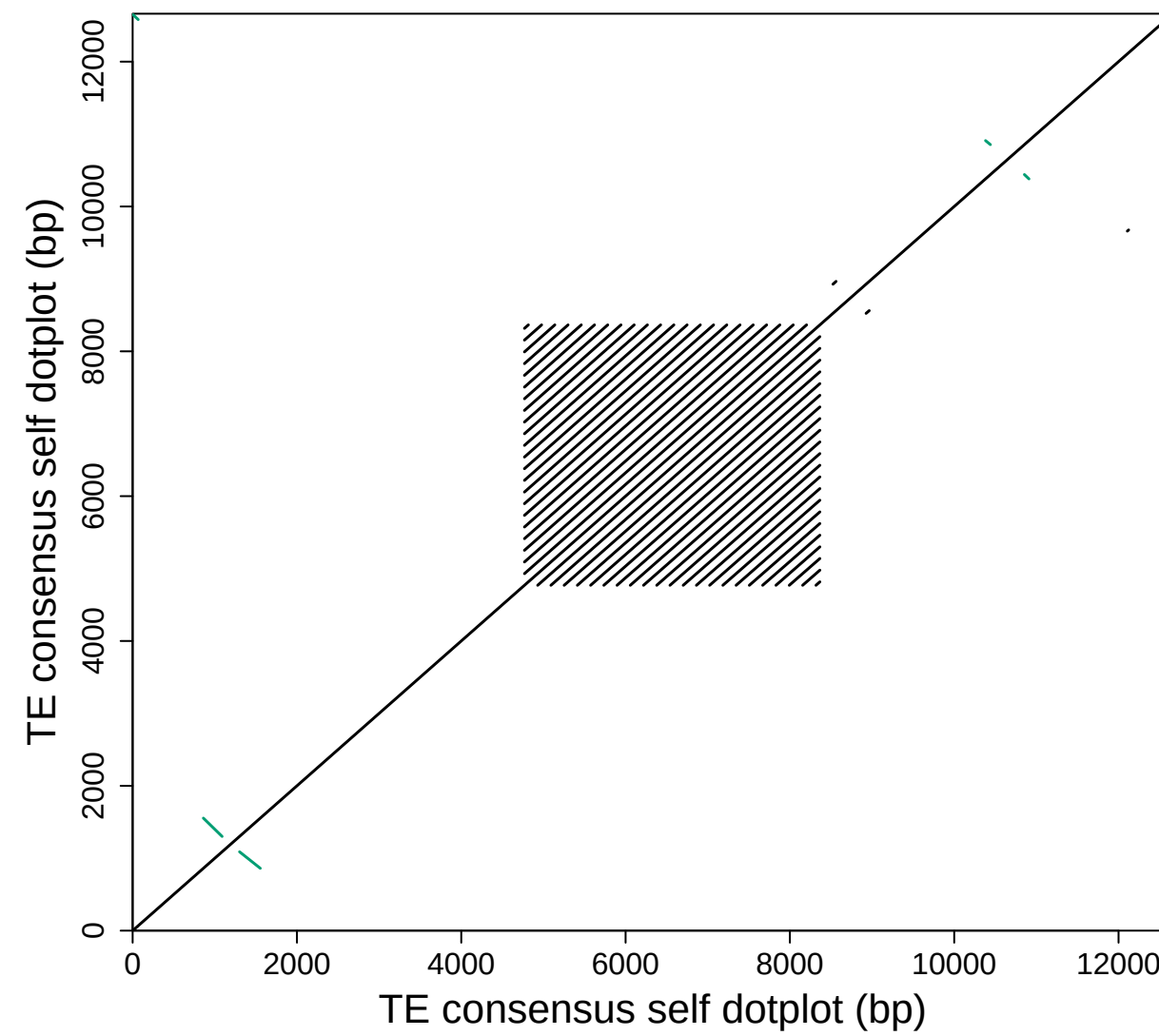
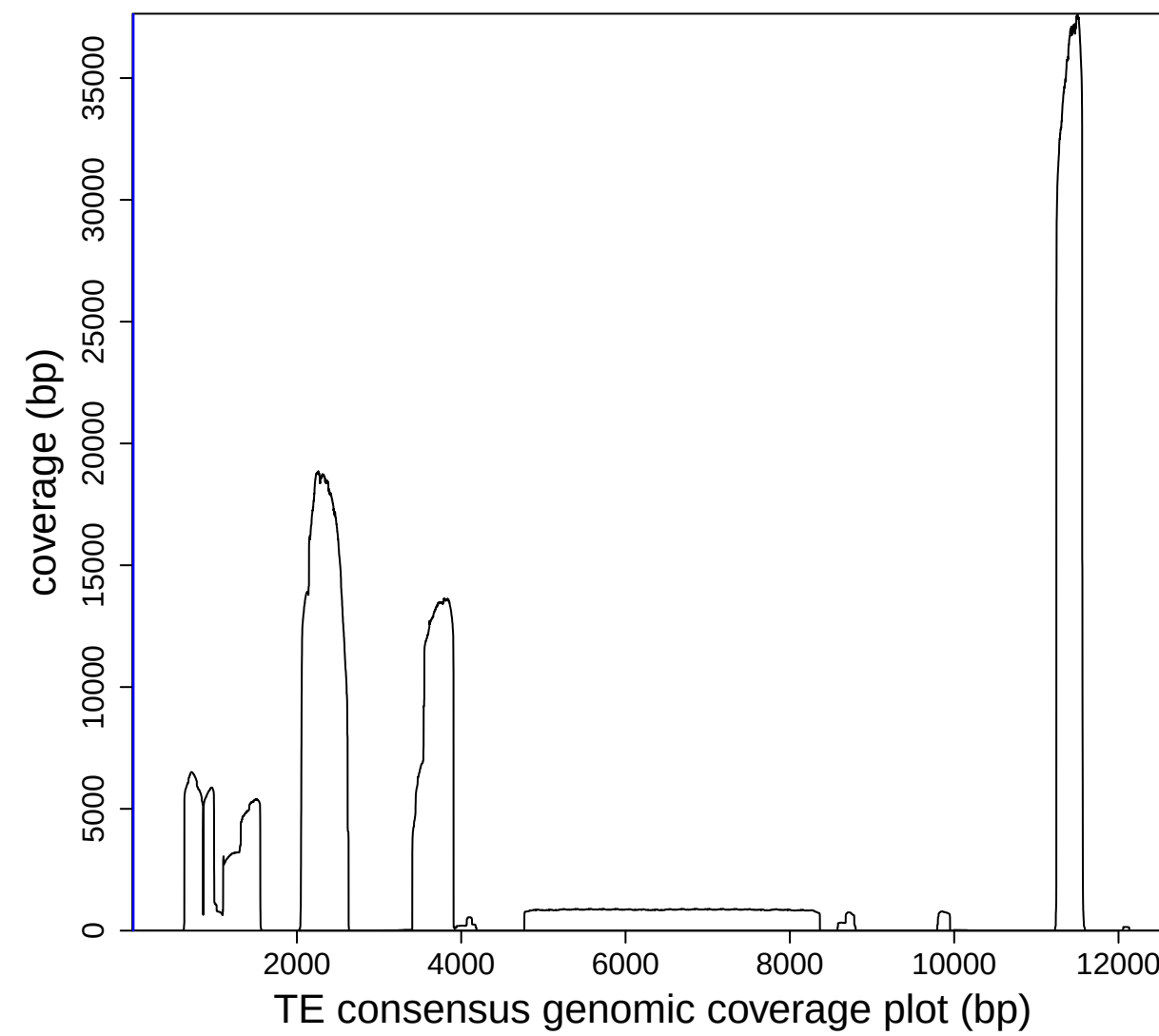
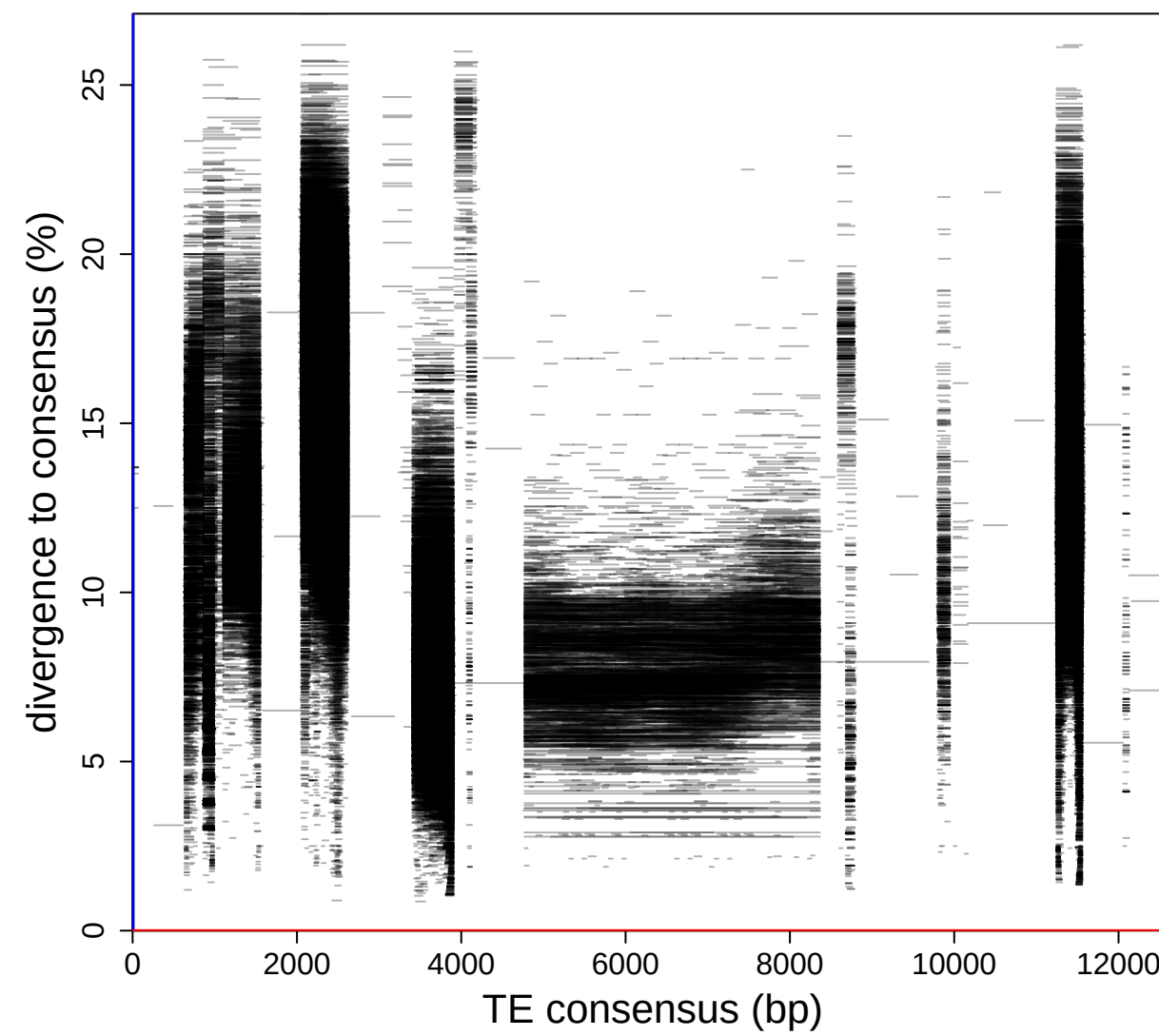
After TEtrimmer 12661 bp



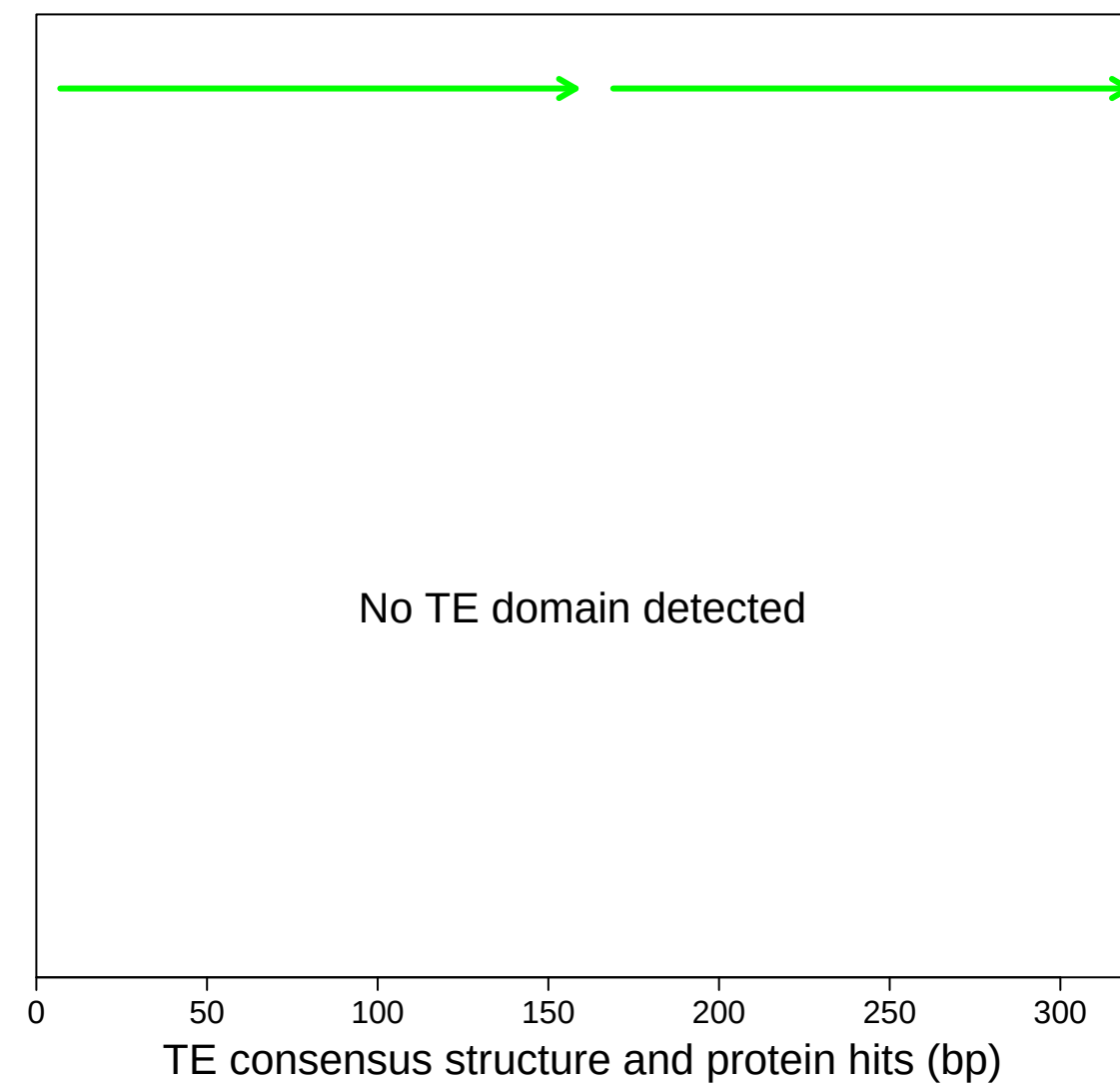
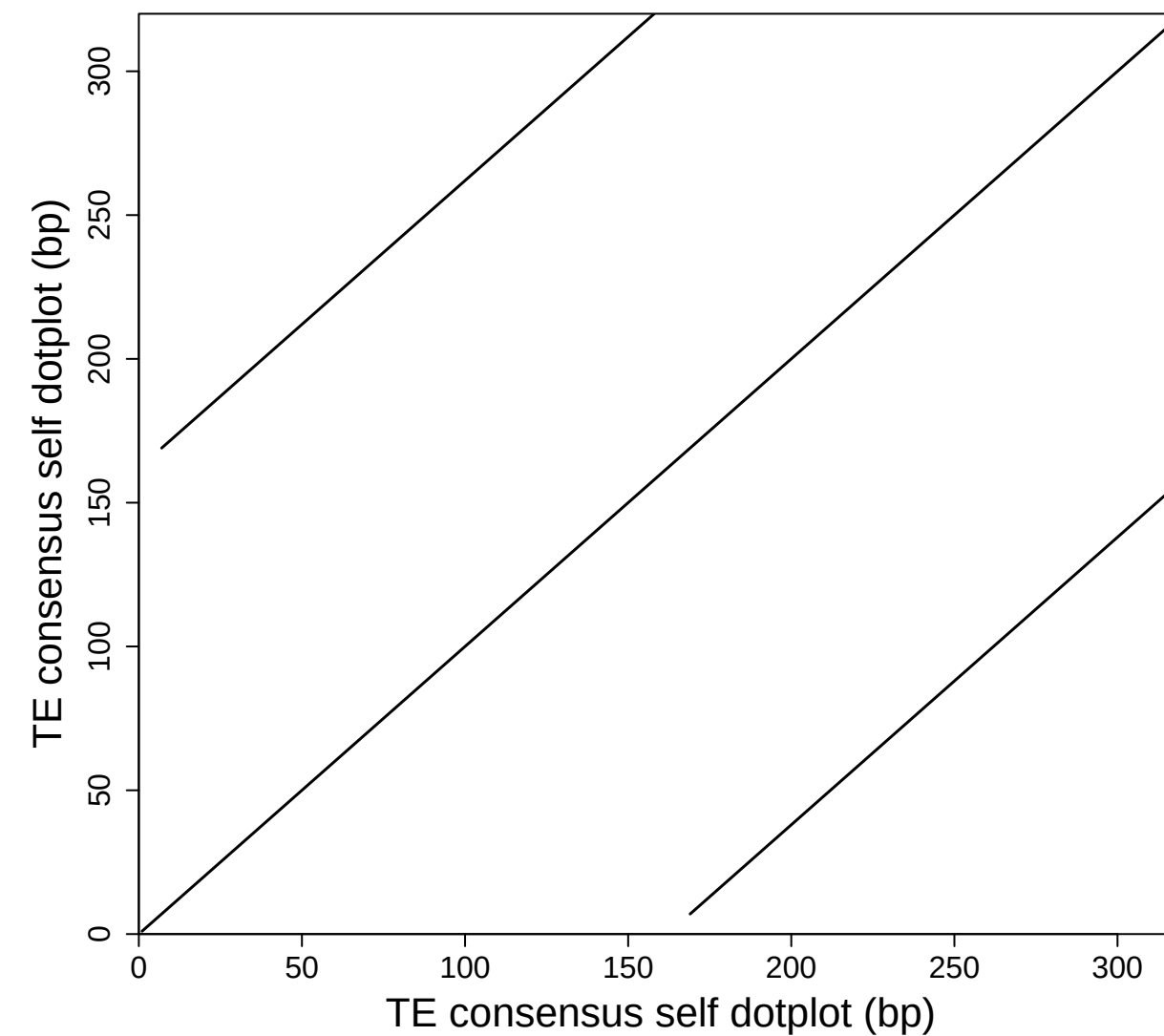
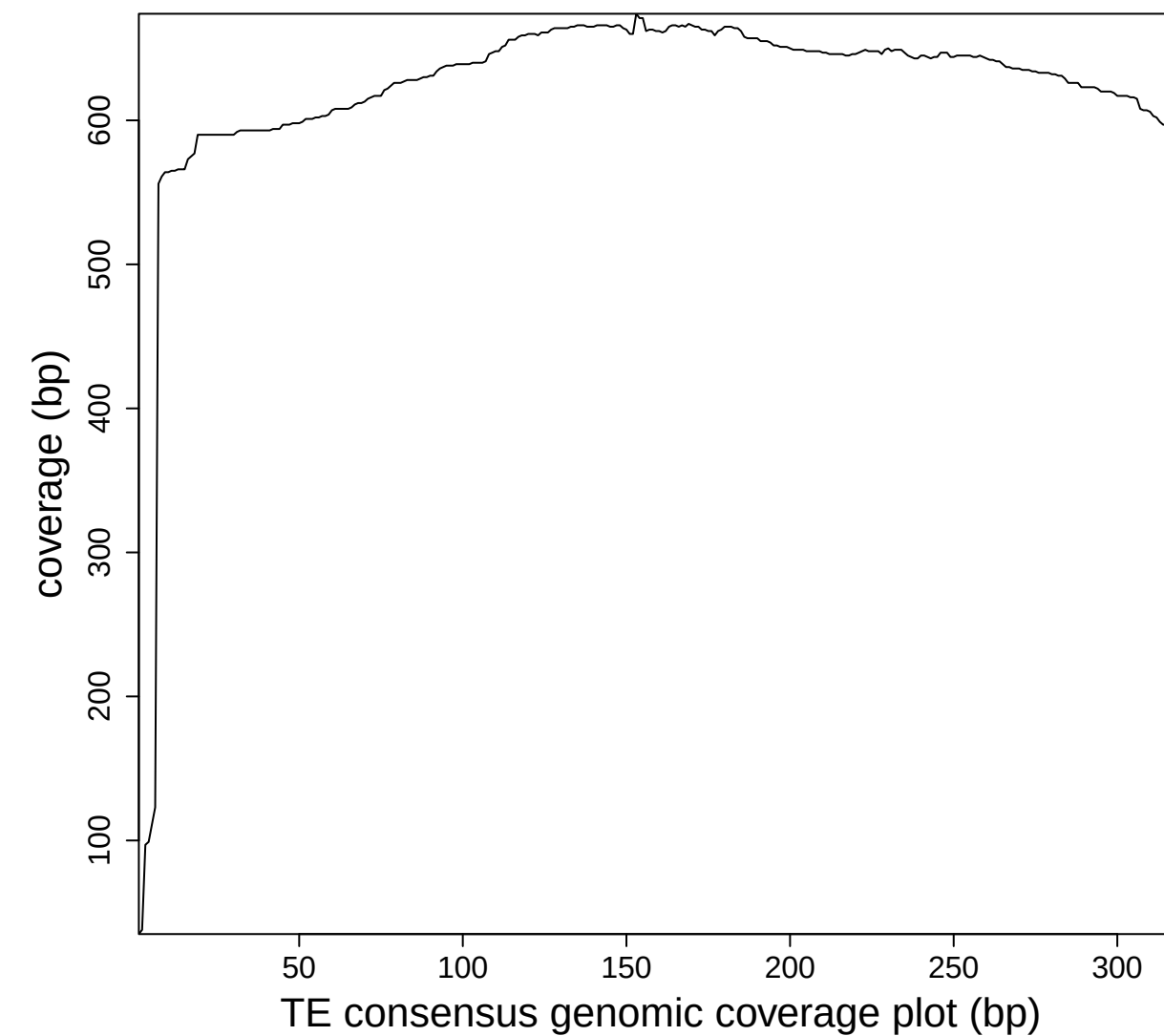
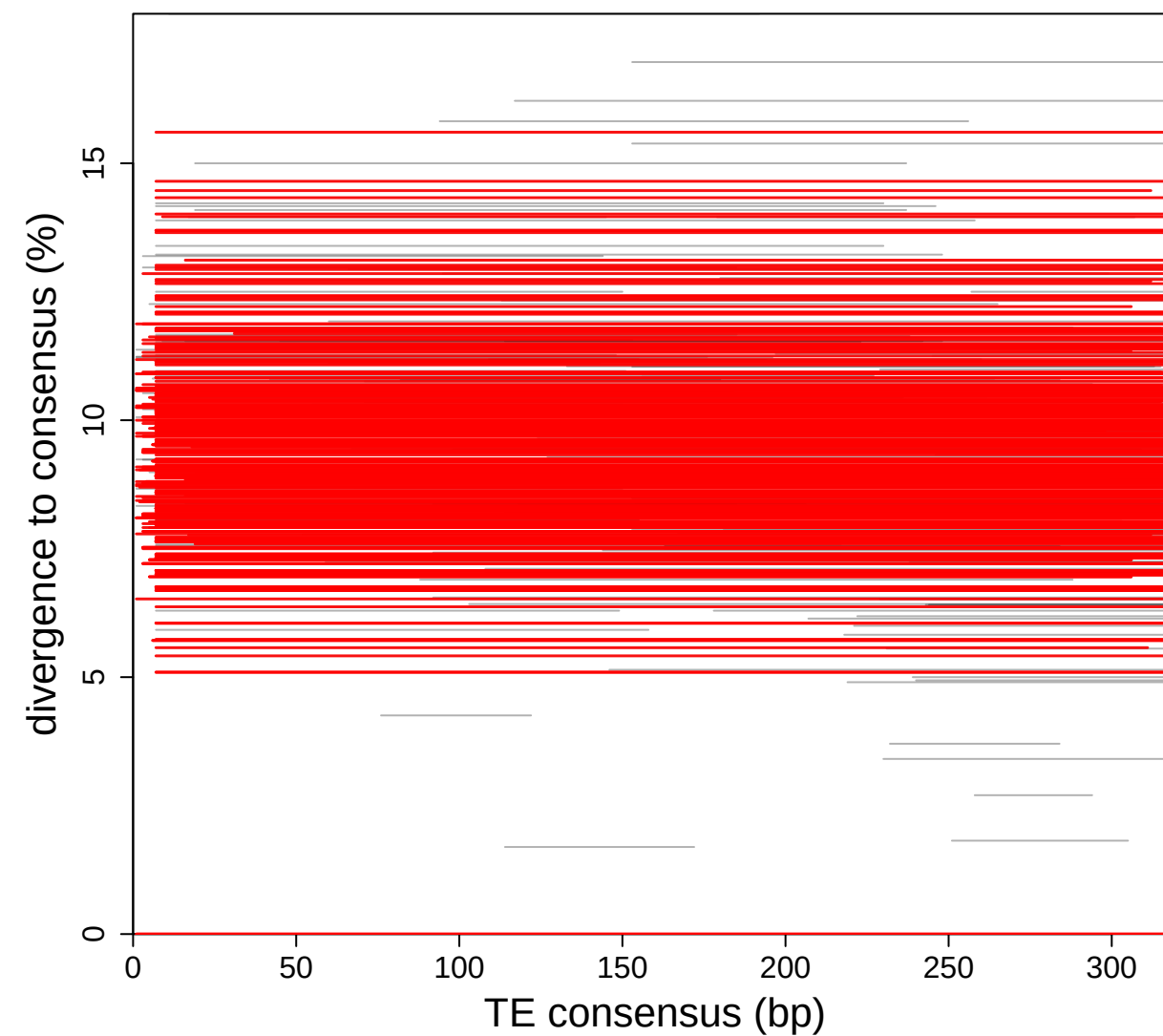
bed_g_1.bed fm_1.bed 0 0 n.bed_g_1.bed fm_2.bed 0 0 bclr
size: 12662bp; fragments: 111340; full length: 1 (>=11395.8bp)

After TEtrimmer Extended plot

Blue lines are boundaries

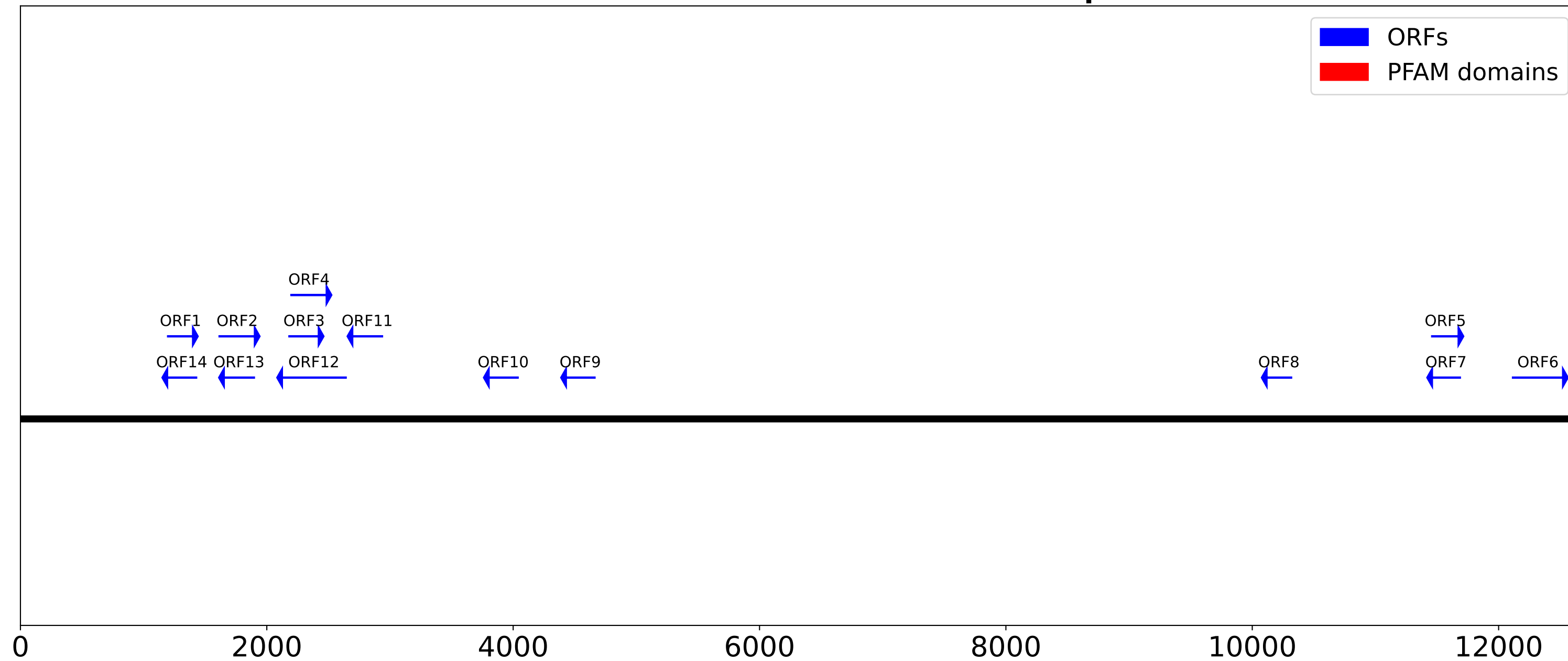


TE: ltr_1_family_153
size: 320bp; fragments: 735; full length: 499 (≥ 288 bp)



Before TEtrimmer 320 bp

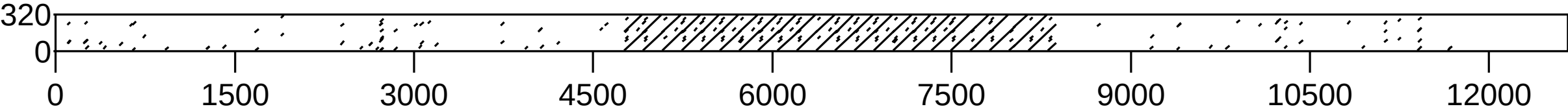
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)