

Start crop Point



End crop Point



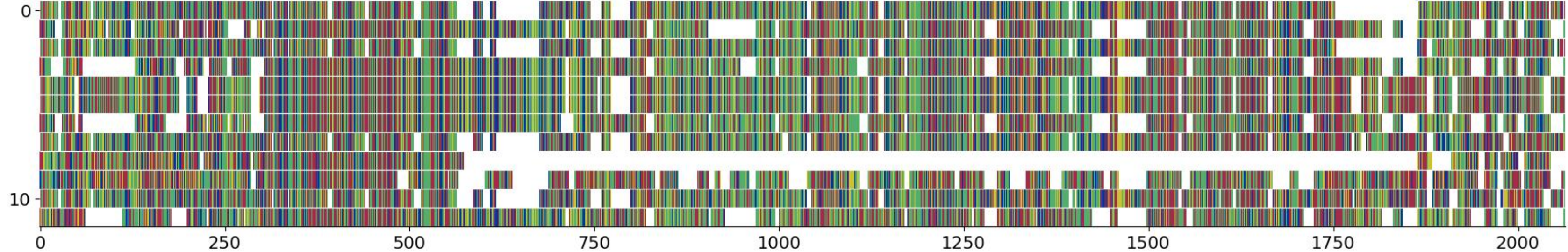
MSA length = 1578



Start crop Point

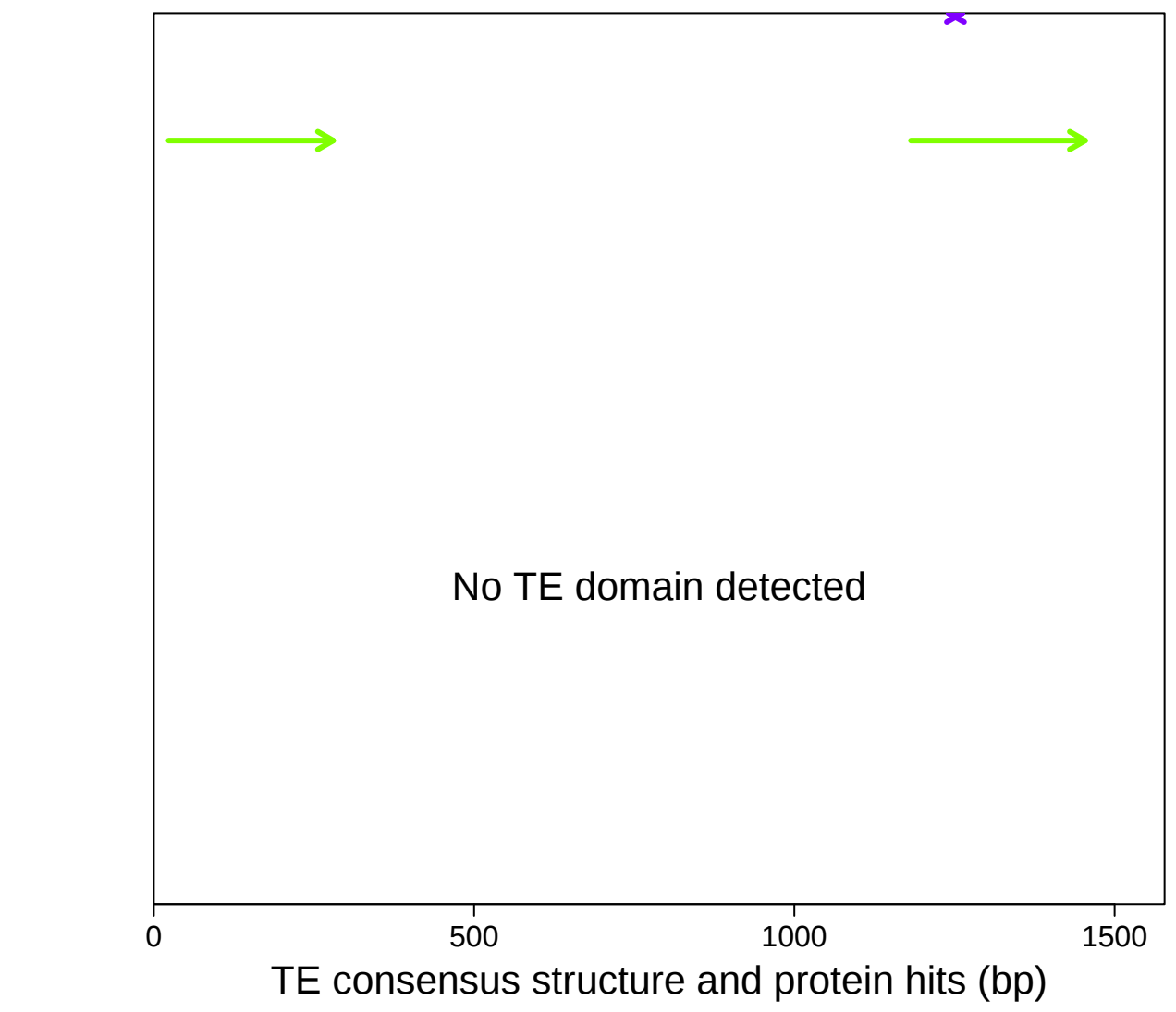
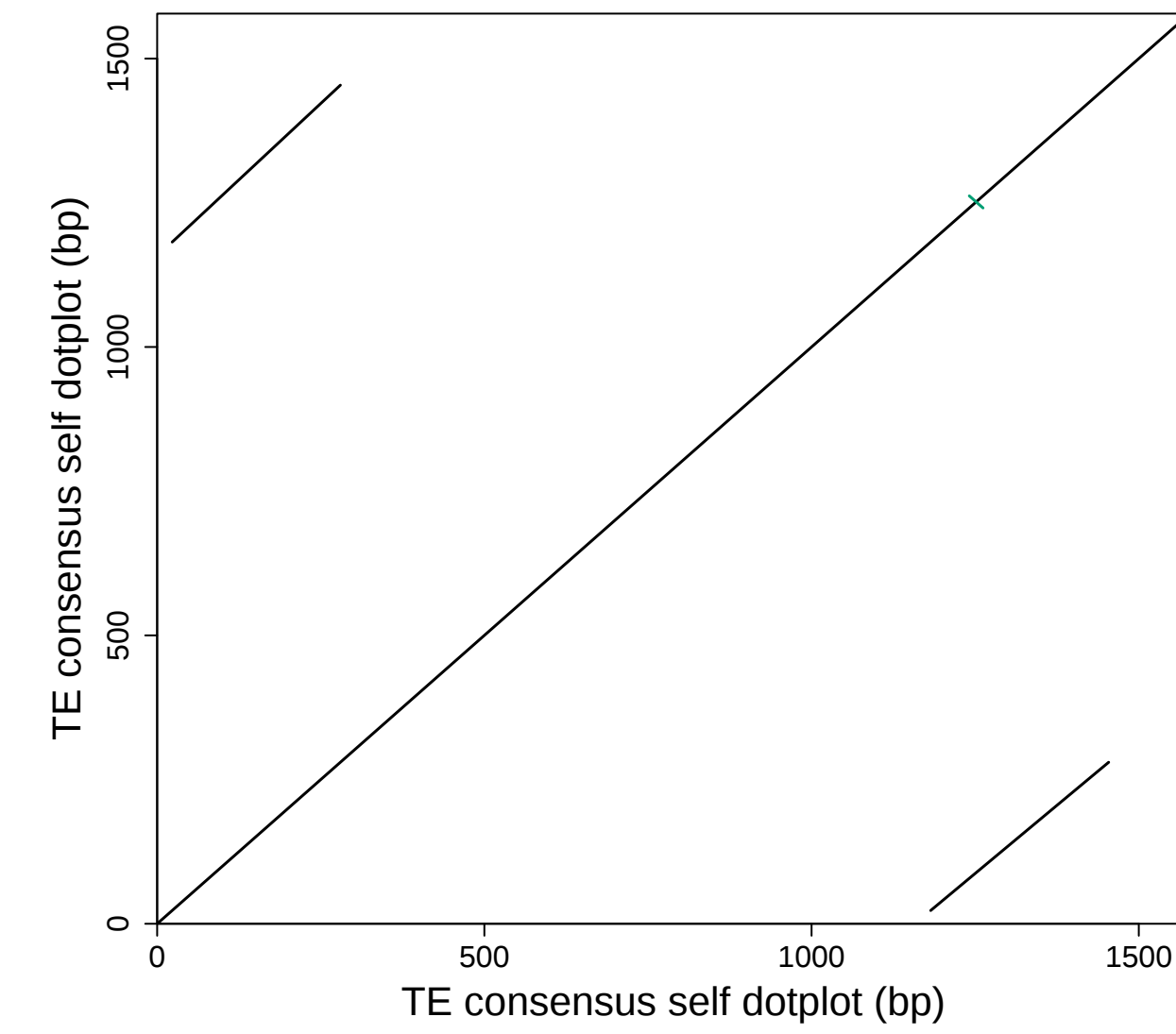
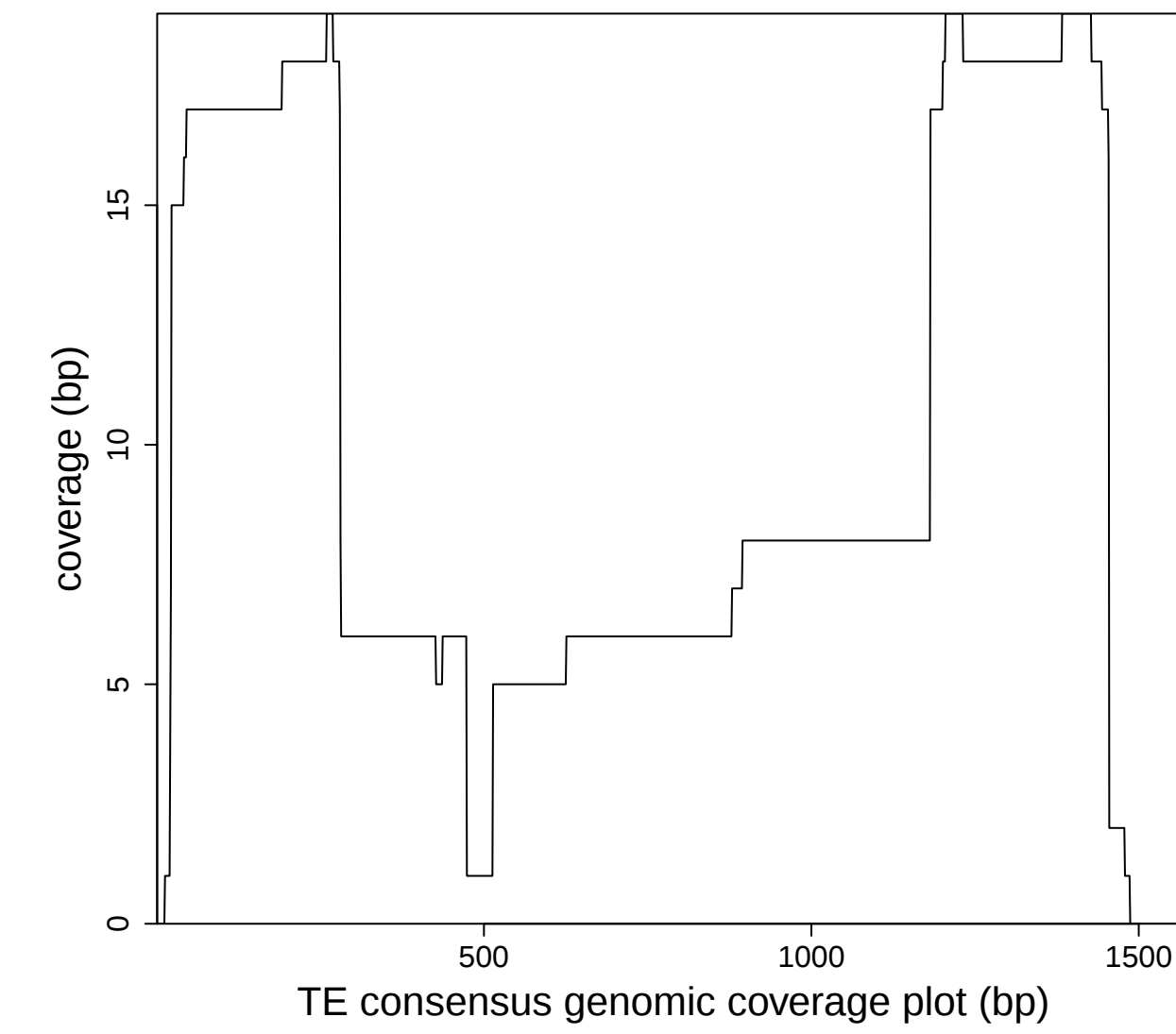
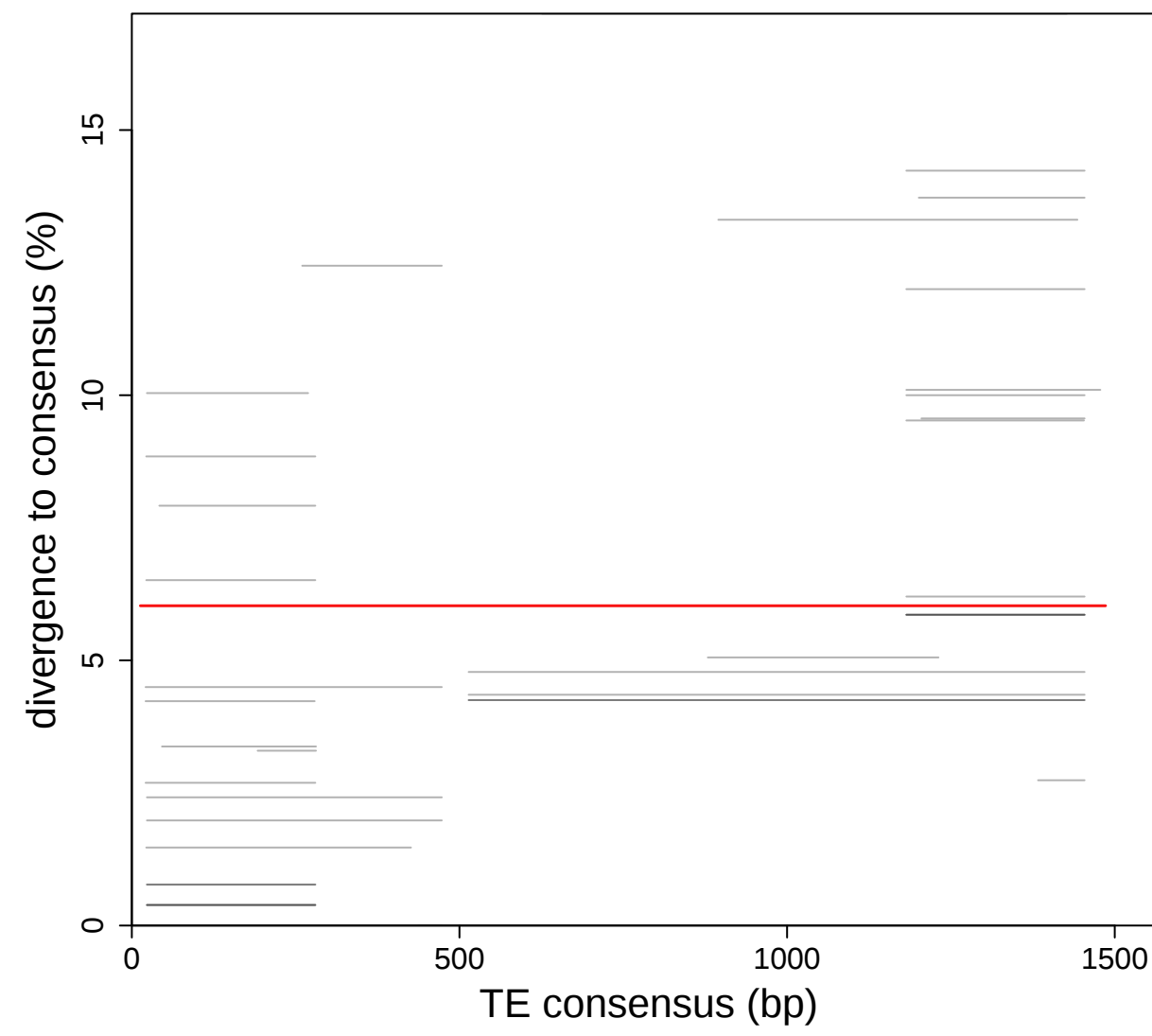


End crop Point



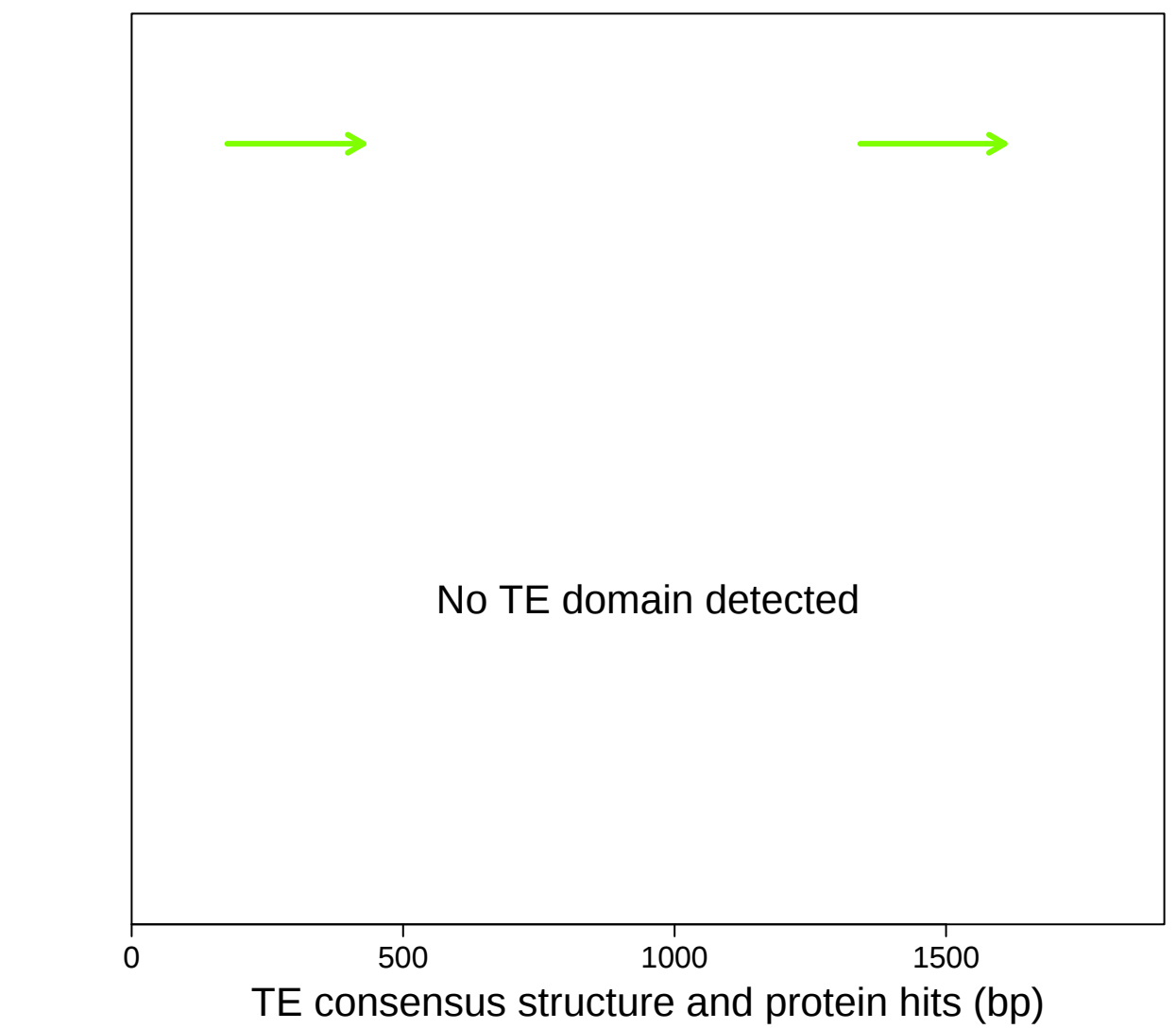
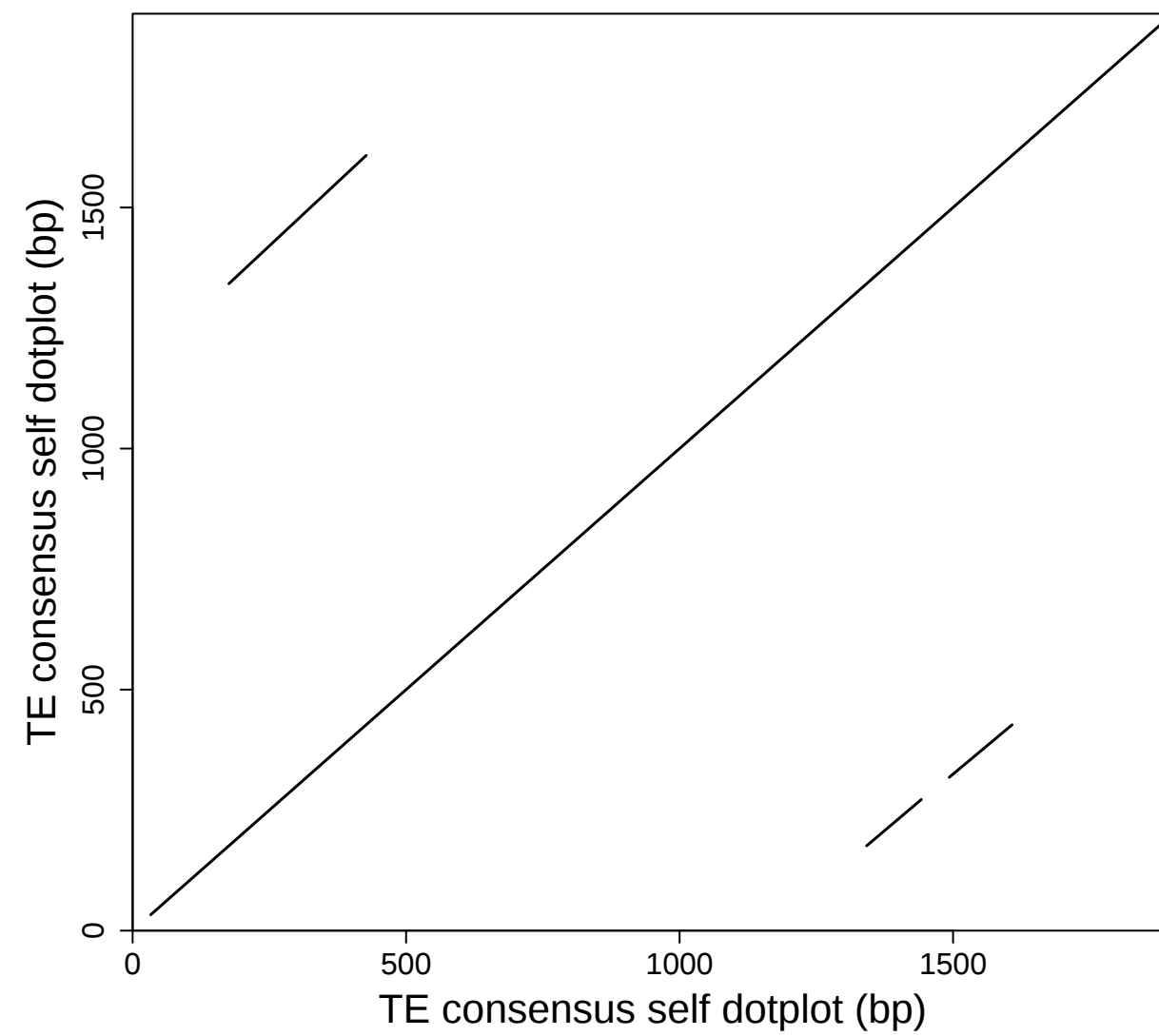
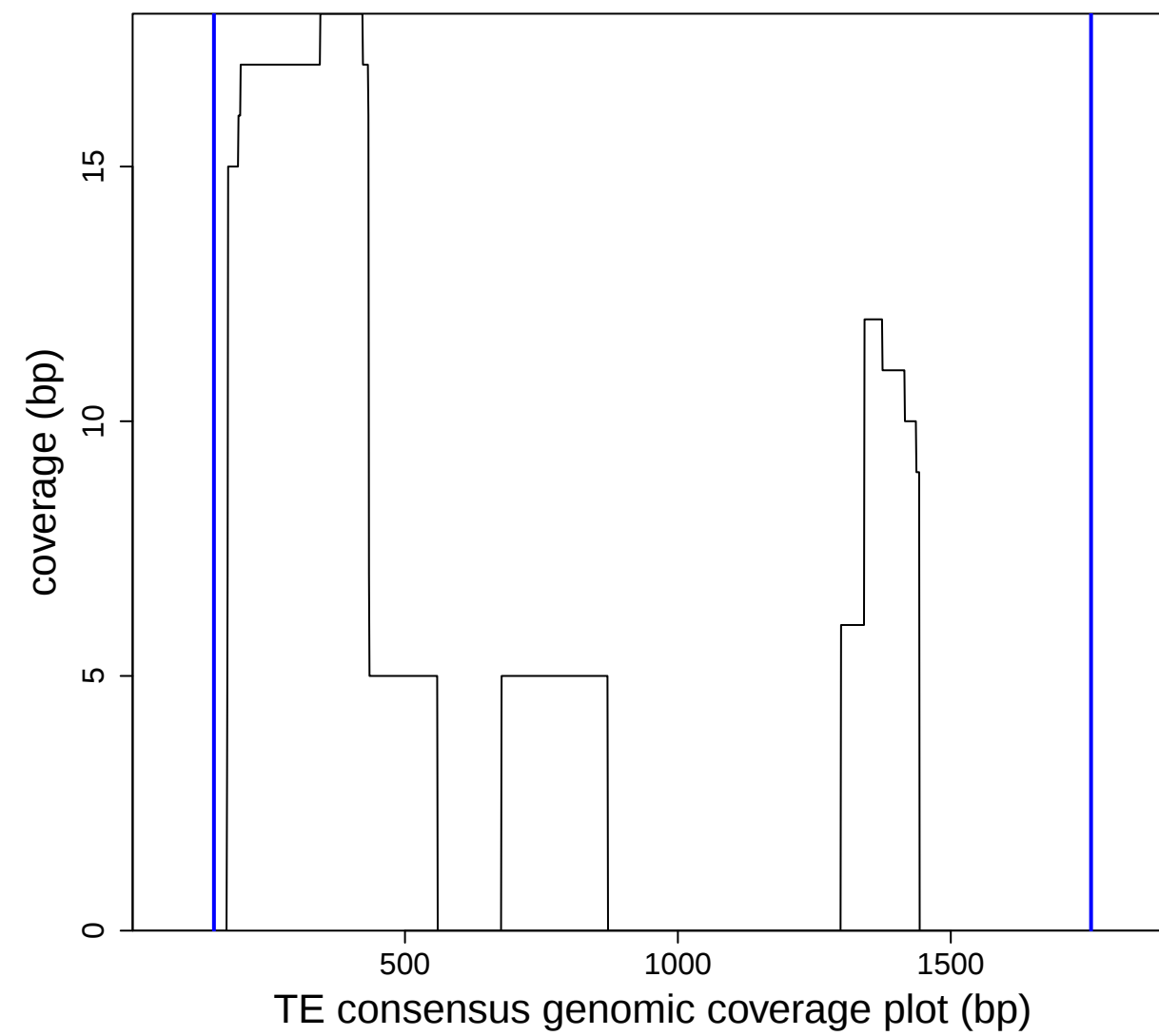
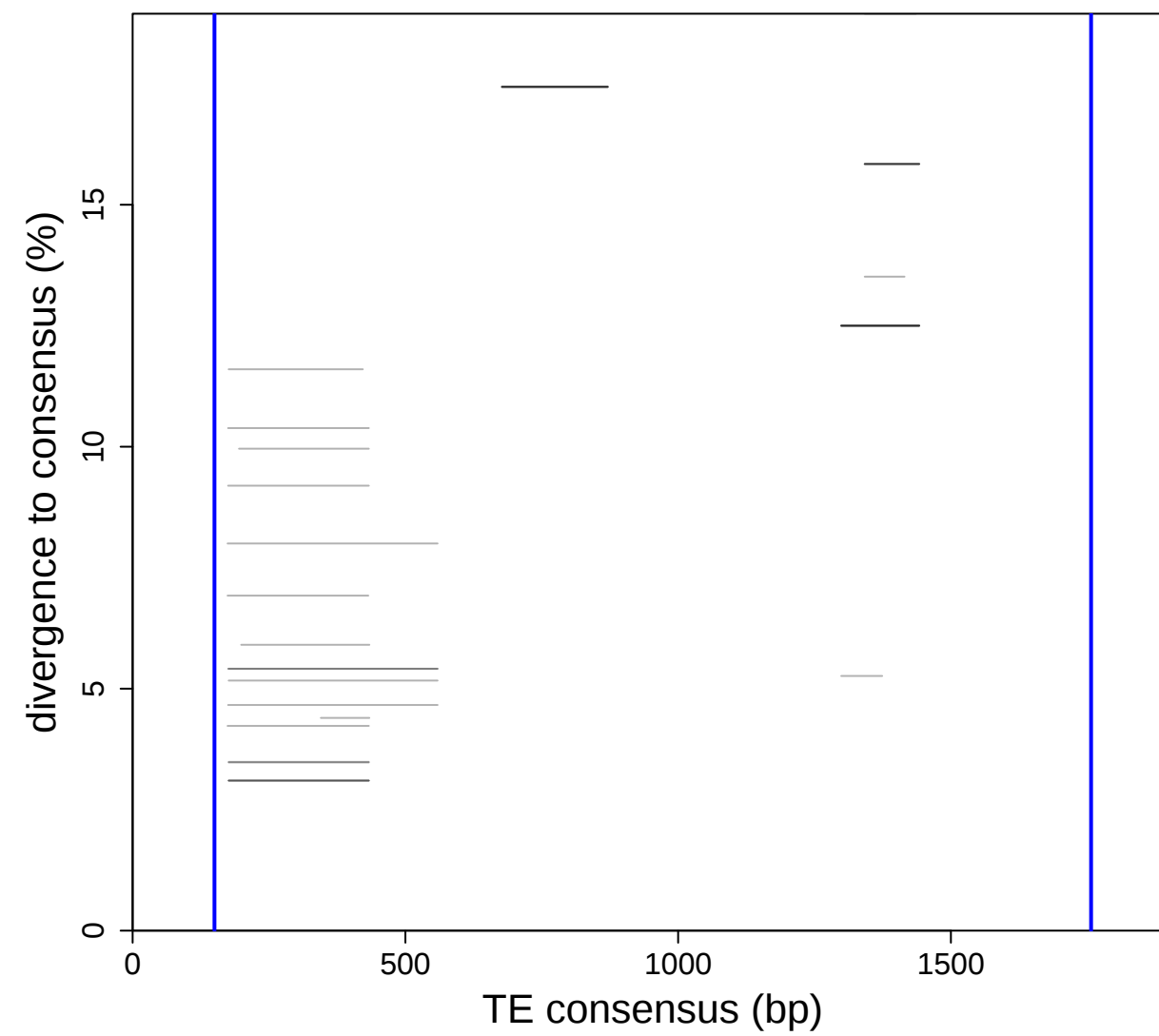
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_al
size: 1578bp; fragments: 39; full length: 1 (>=1420.2bp)

After TEtrimmer 1578 bp



bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 1907bp; fragments: 35; full length: 0 (>=1716.3bp)

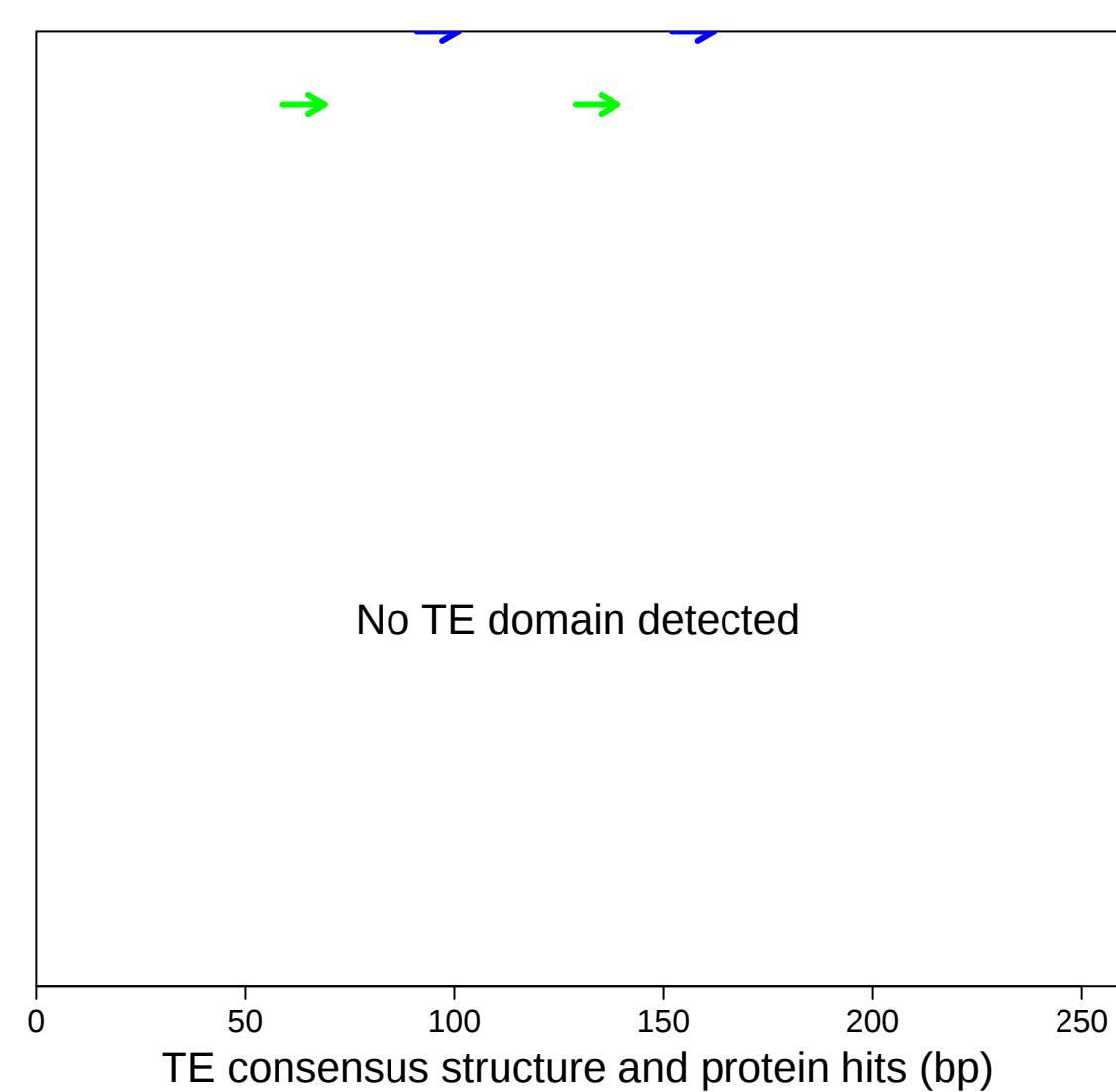
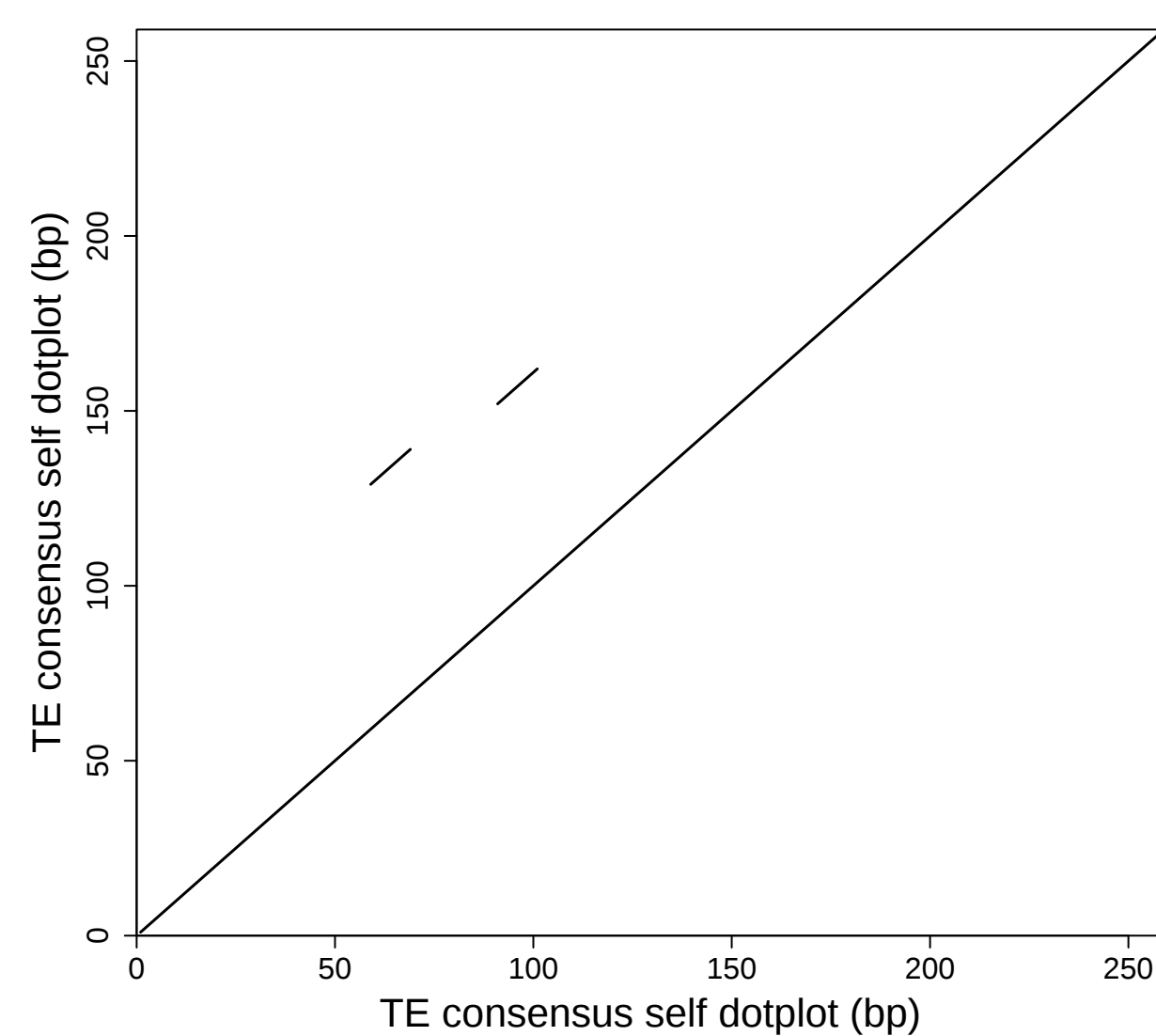
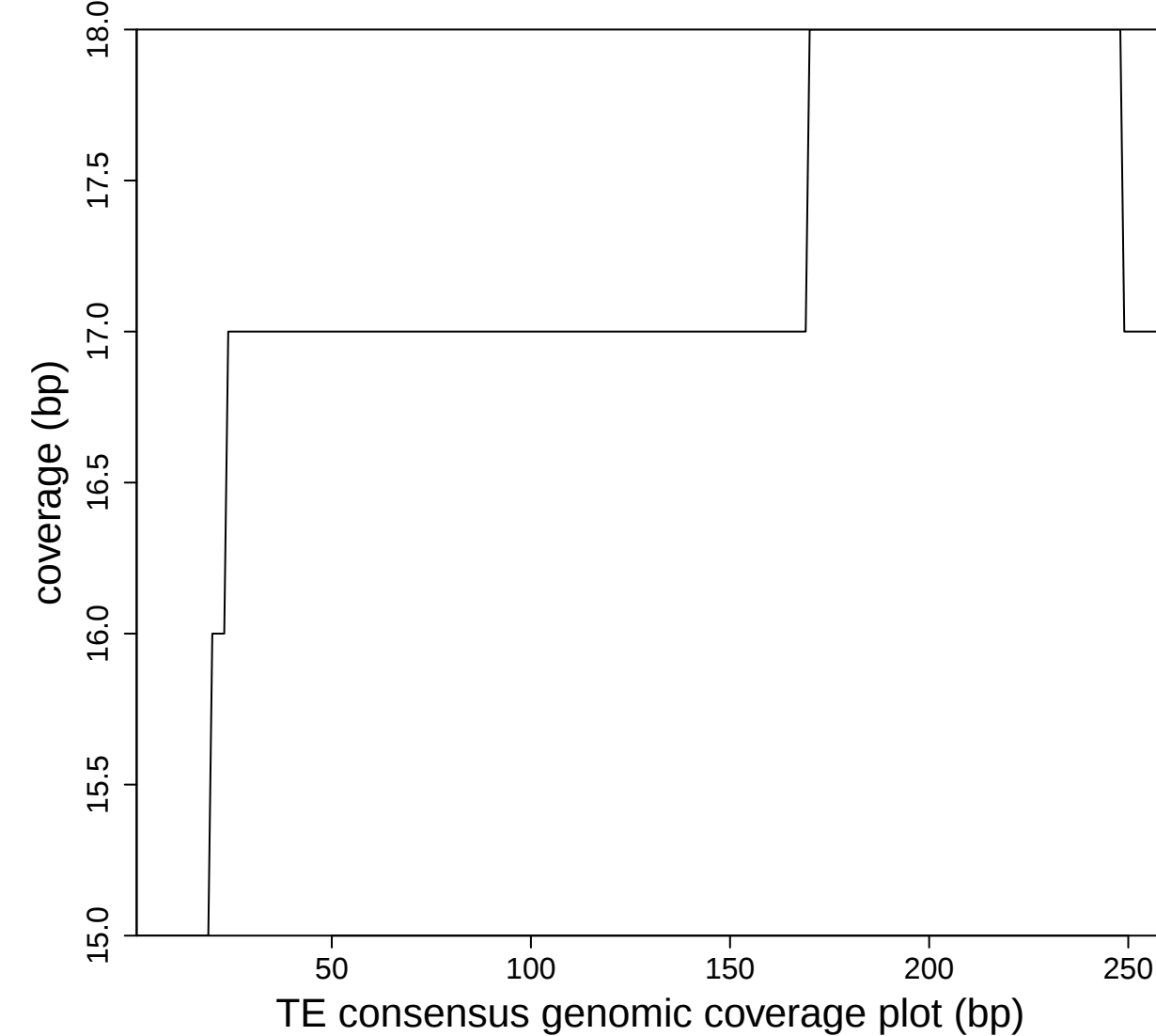
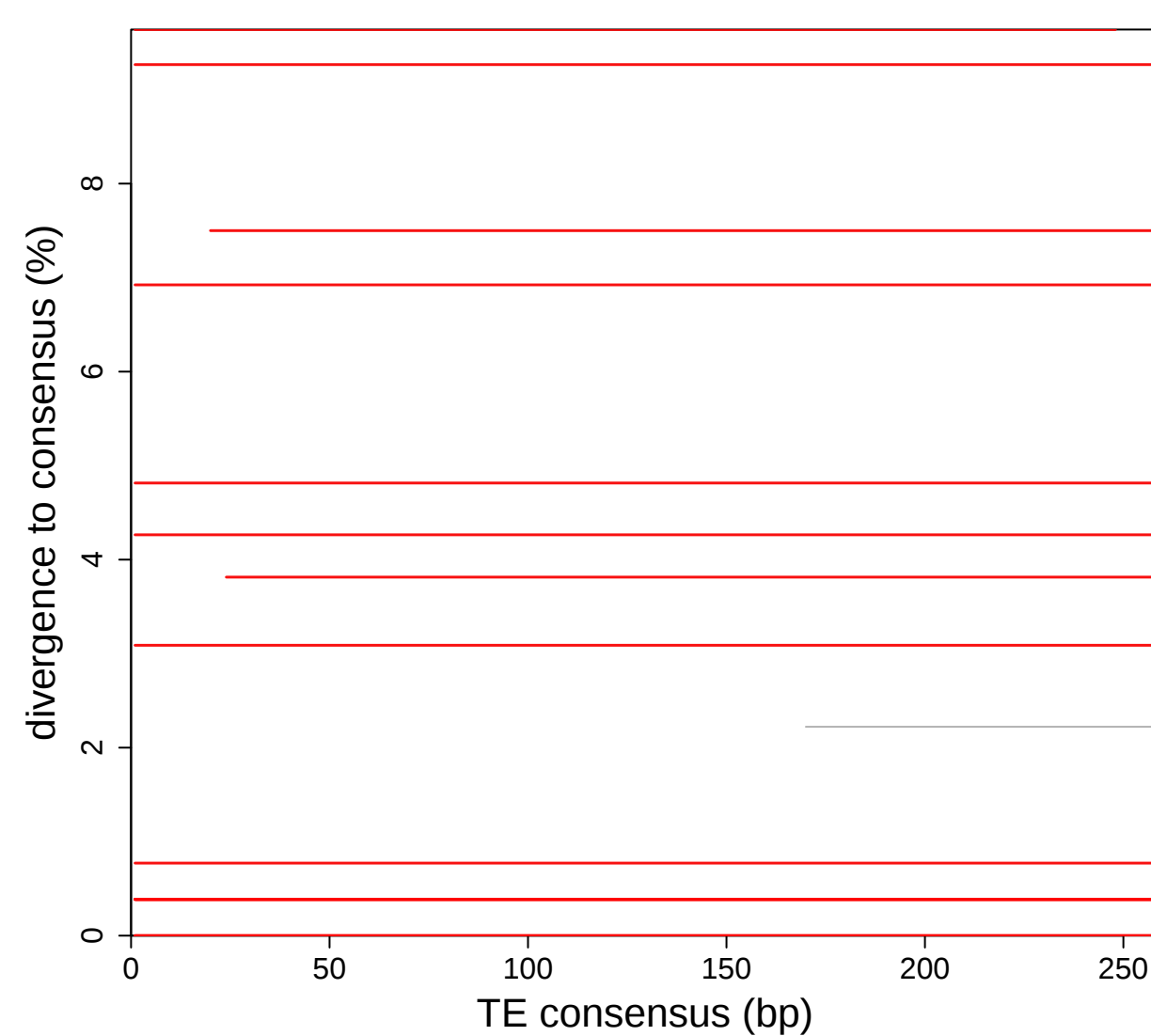
After TEtrimmer Extended plot Blue lines are boundaries



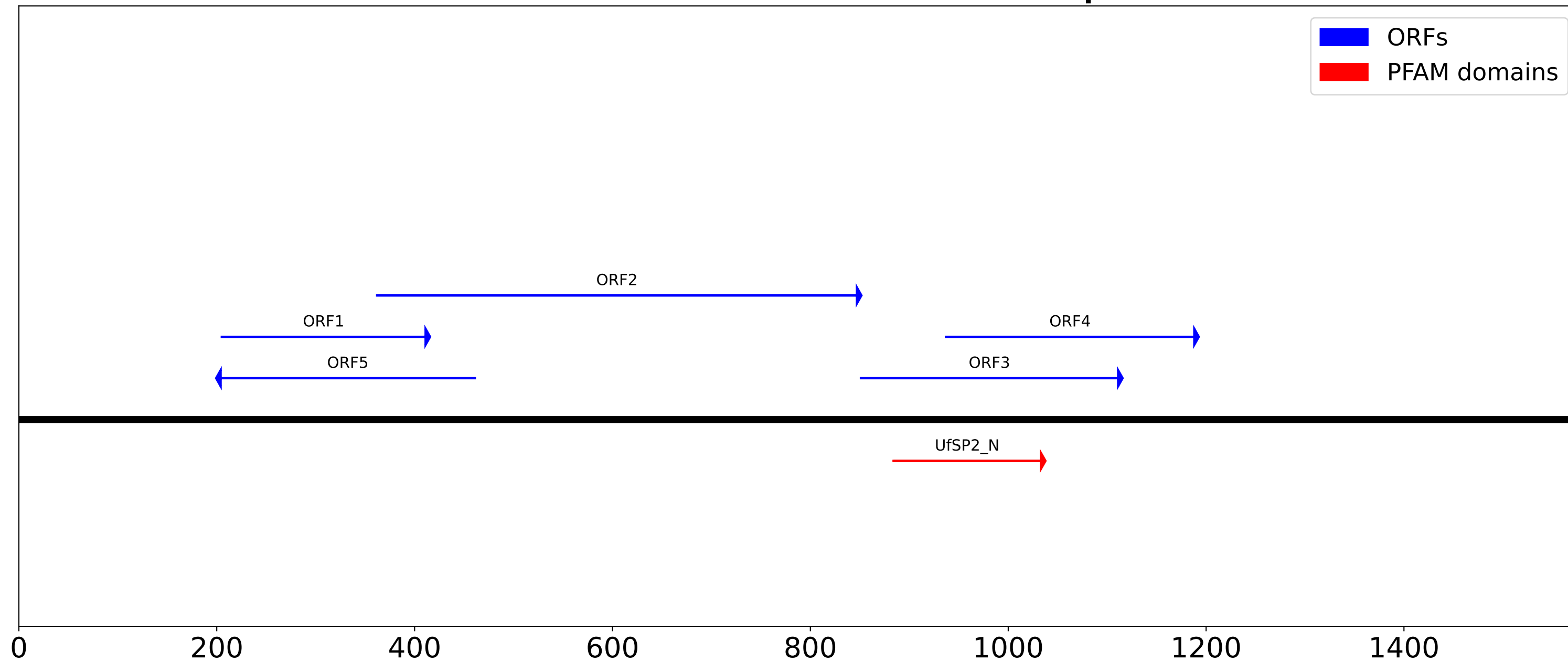
Before TEtrimmer 259 bp

TE: ltr_1_family_169

size: 259bp; fragments: 18; full length: 17 (≥ 233.1 bp)



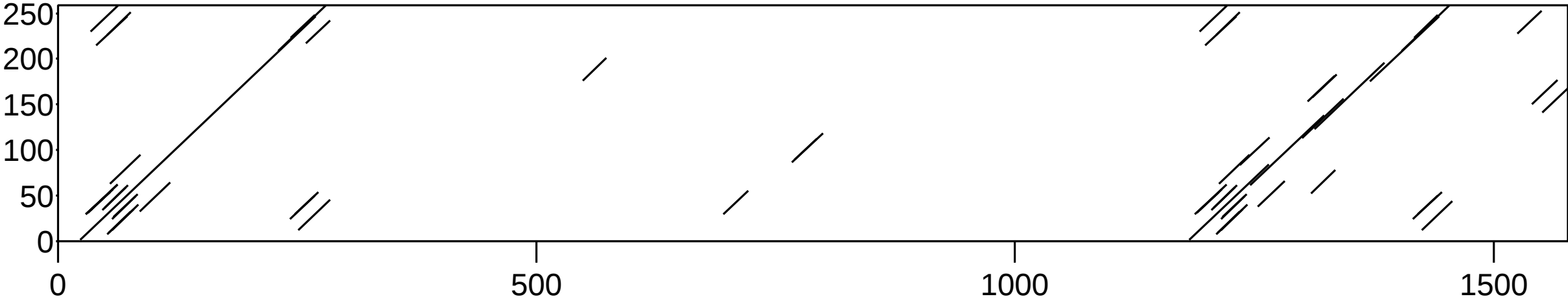
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)