

Start crop Point



End crop Point



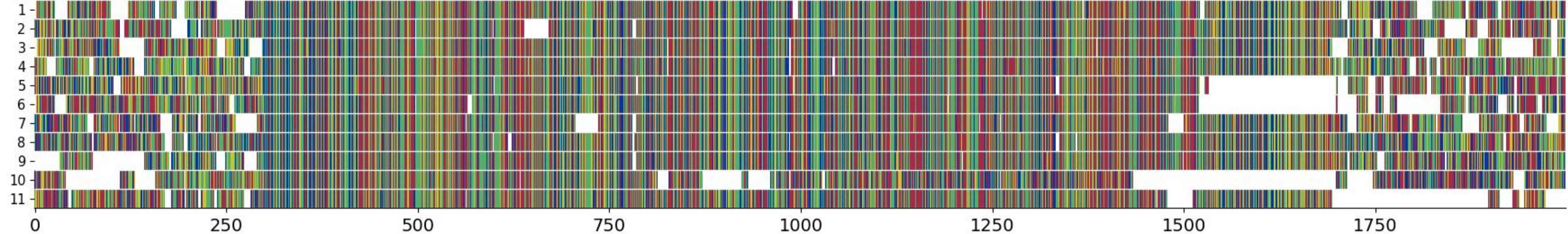
MSA length = 1399



Start crop Point

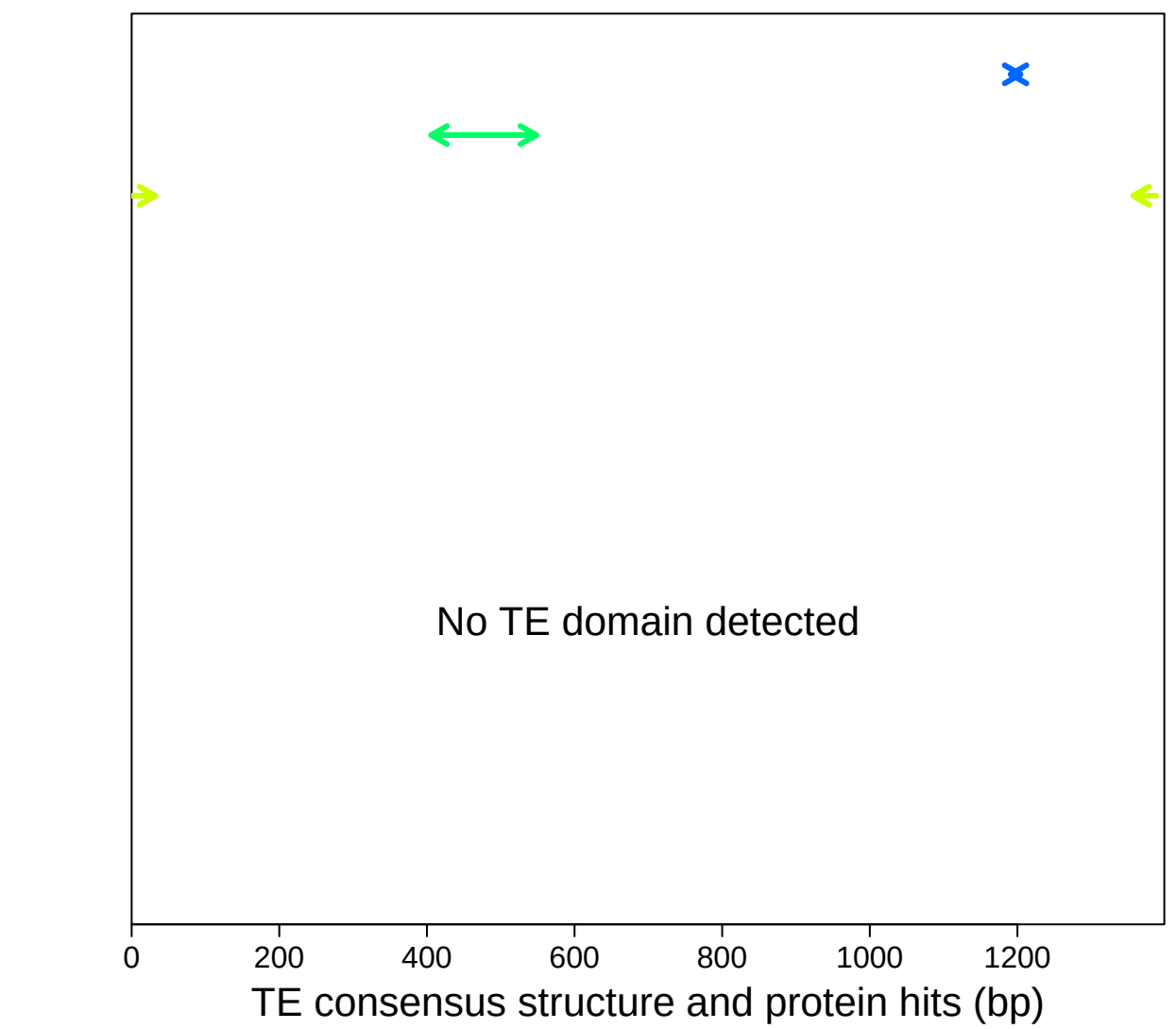
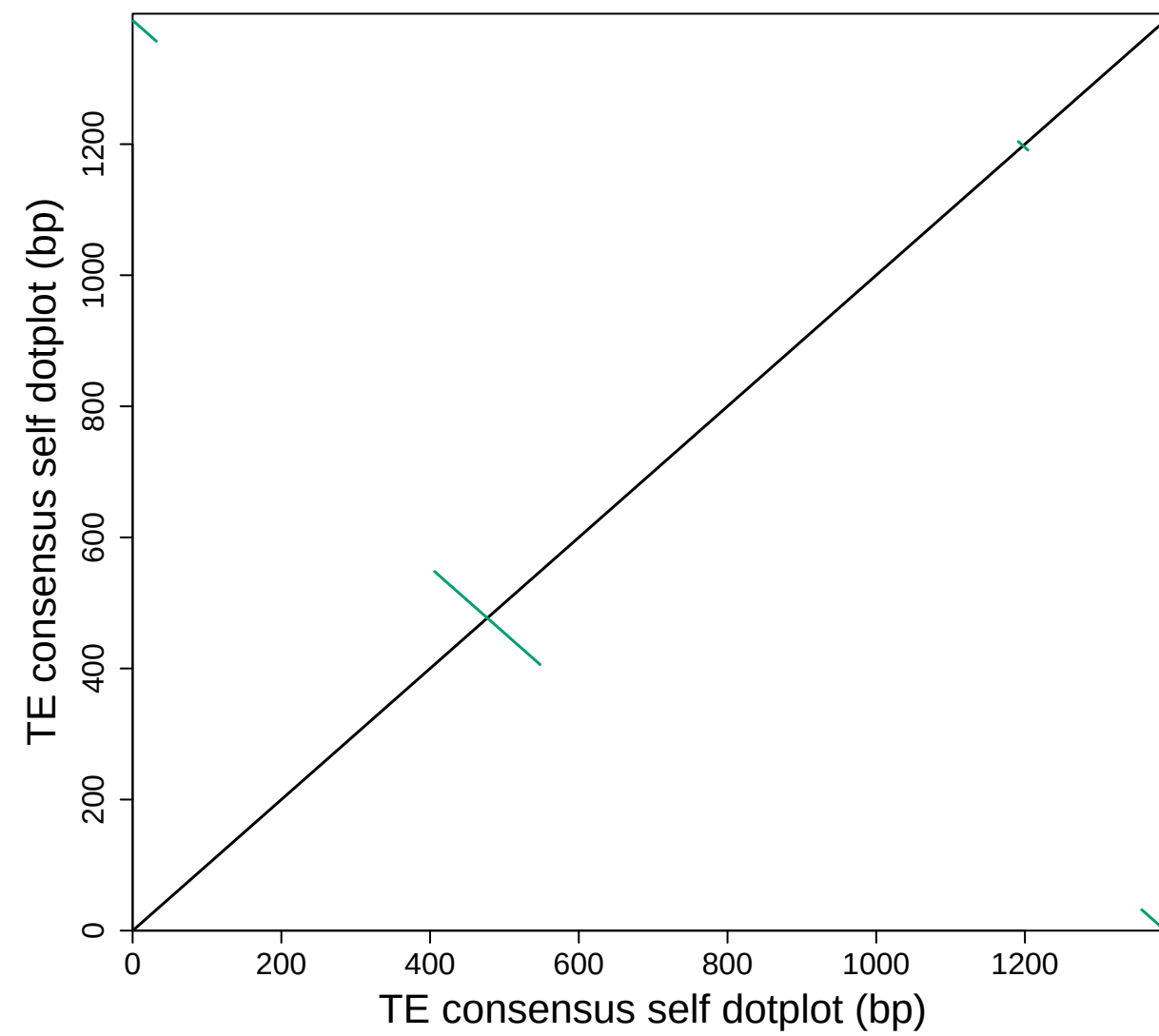
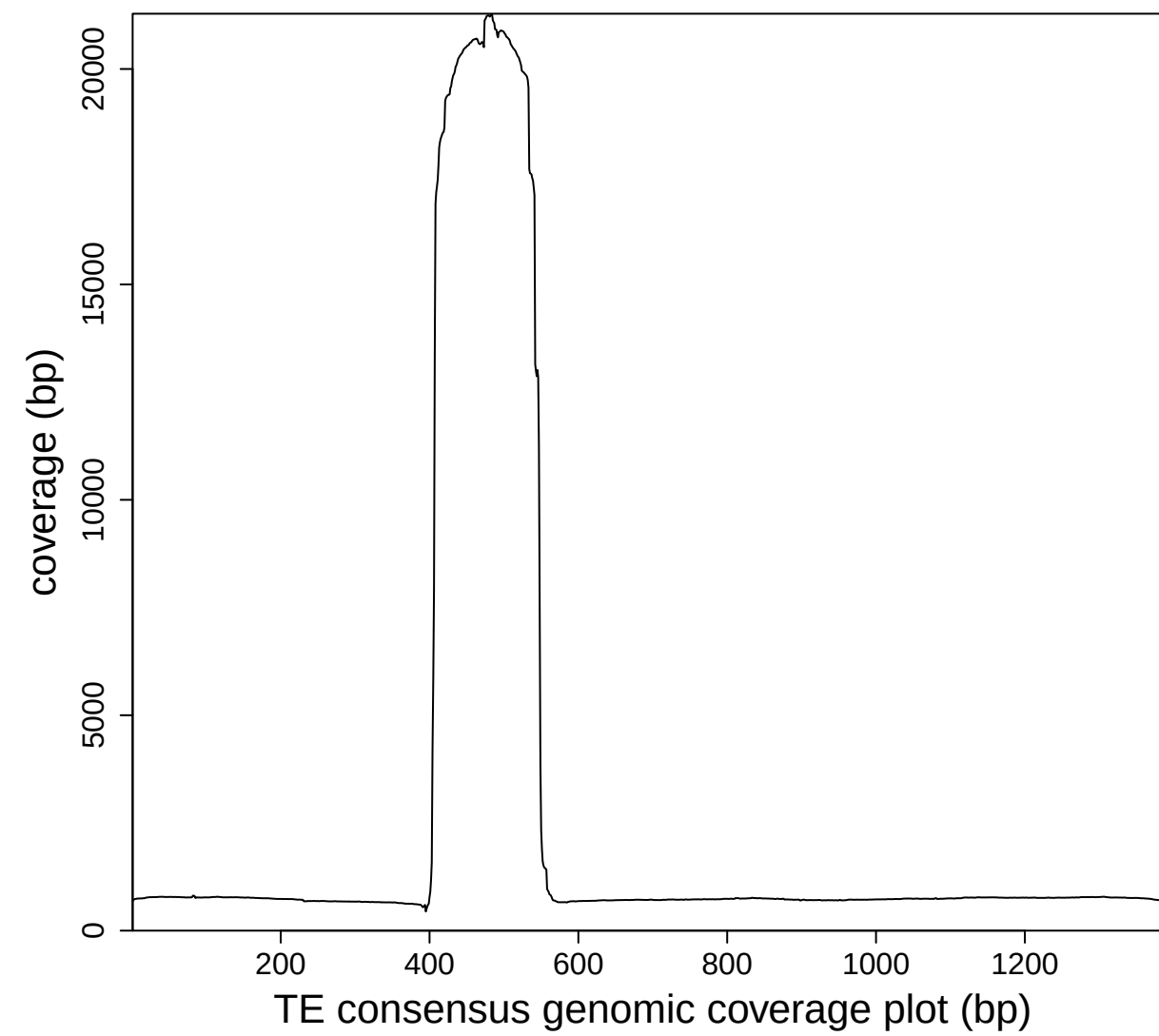
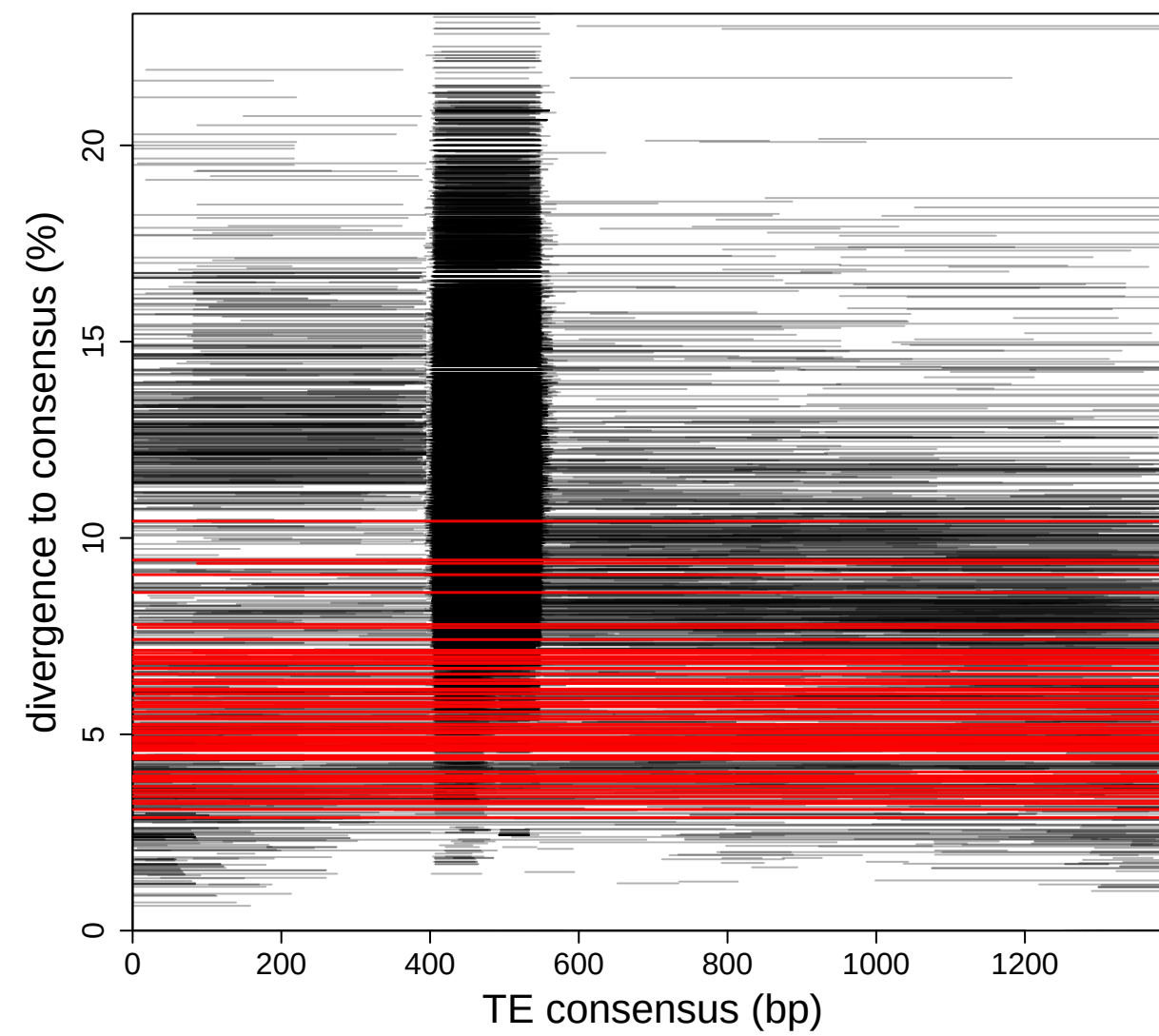


End crop Point



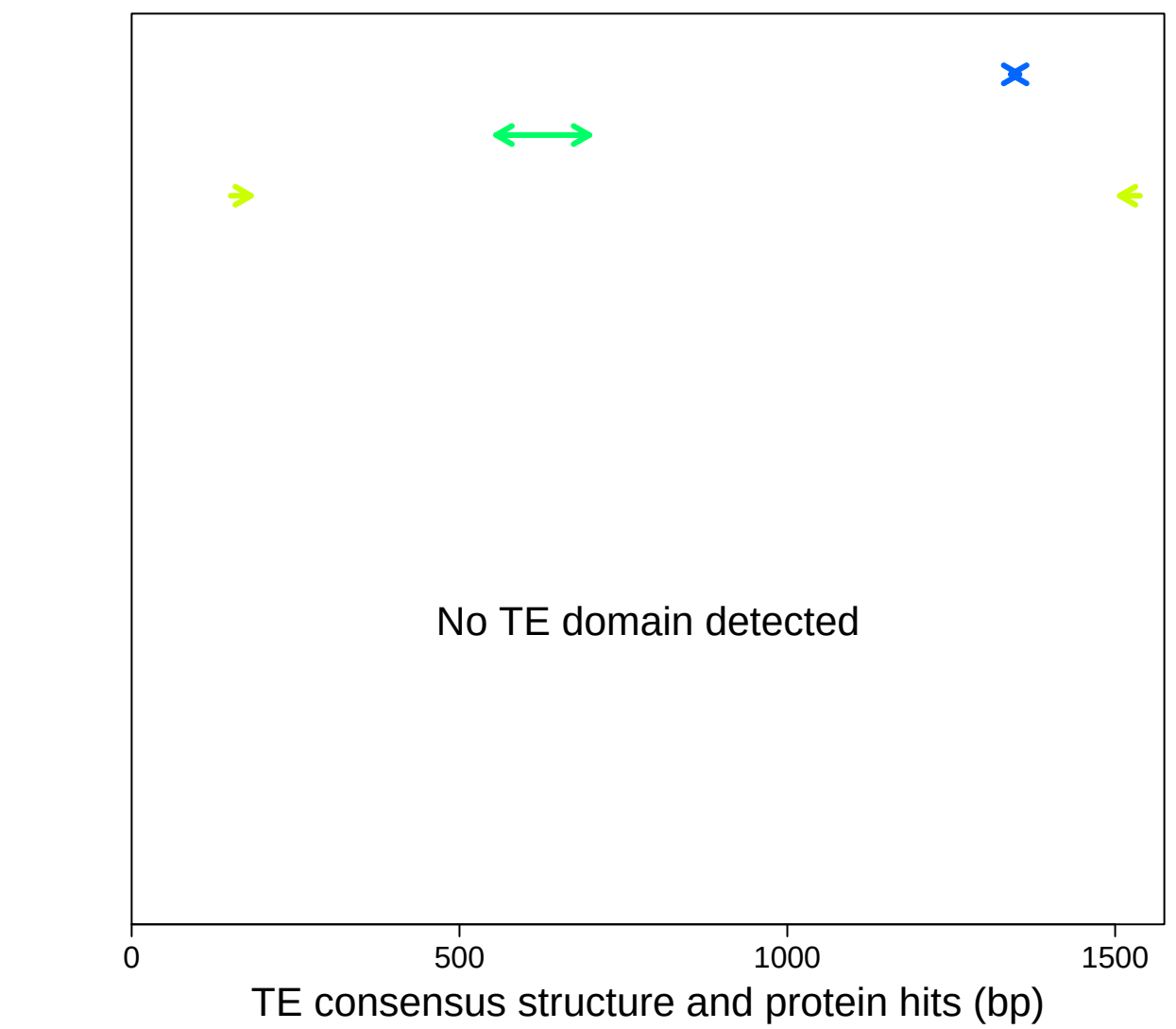
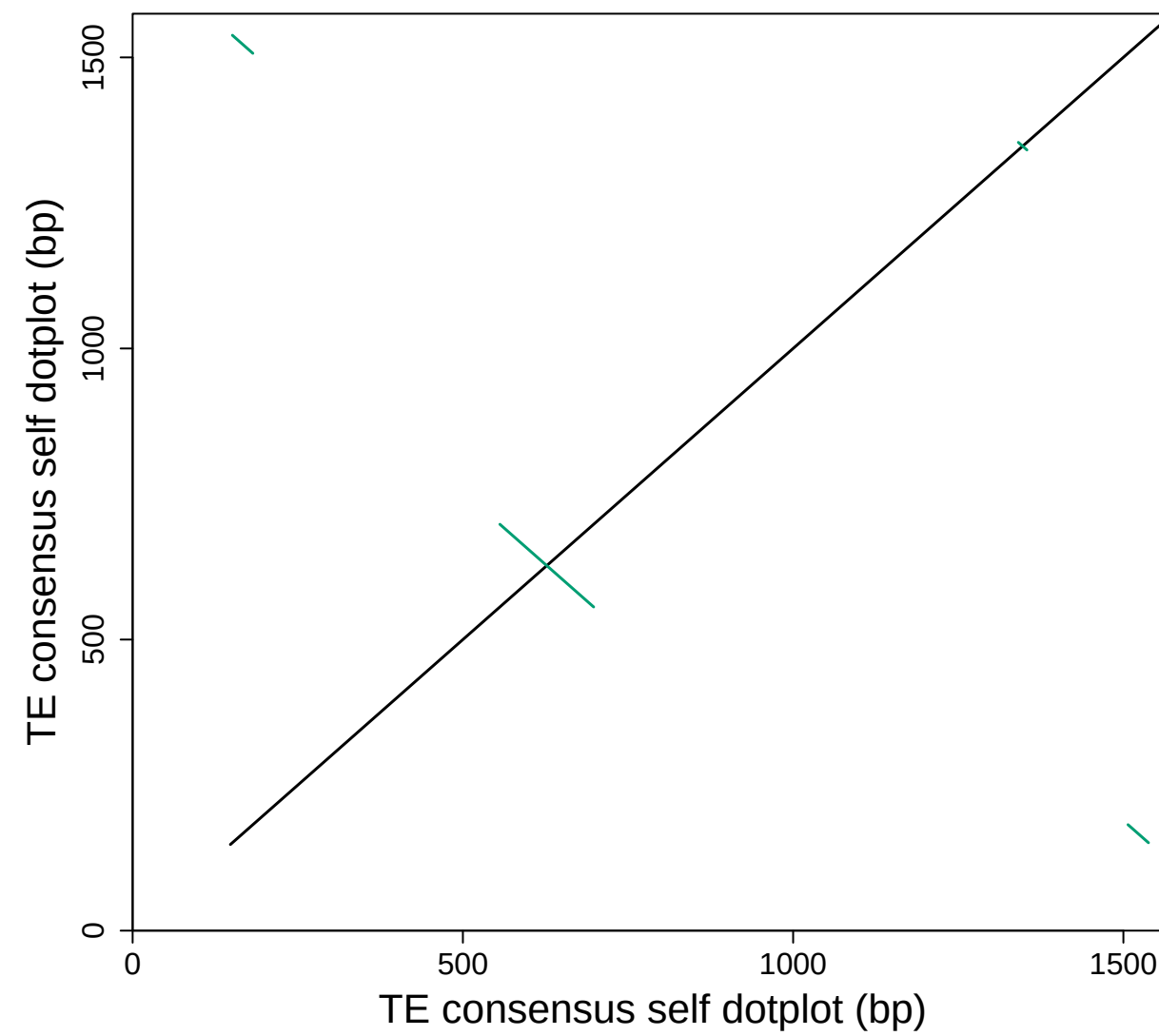
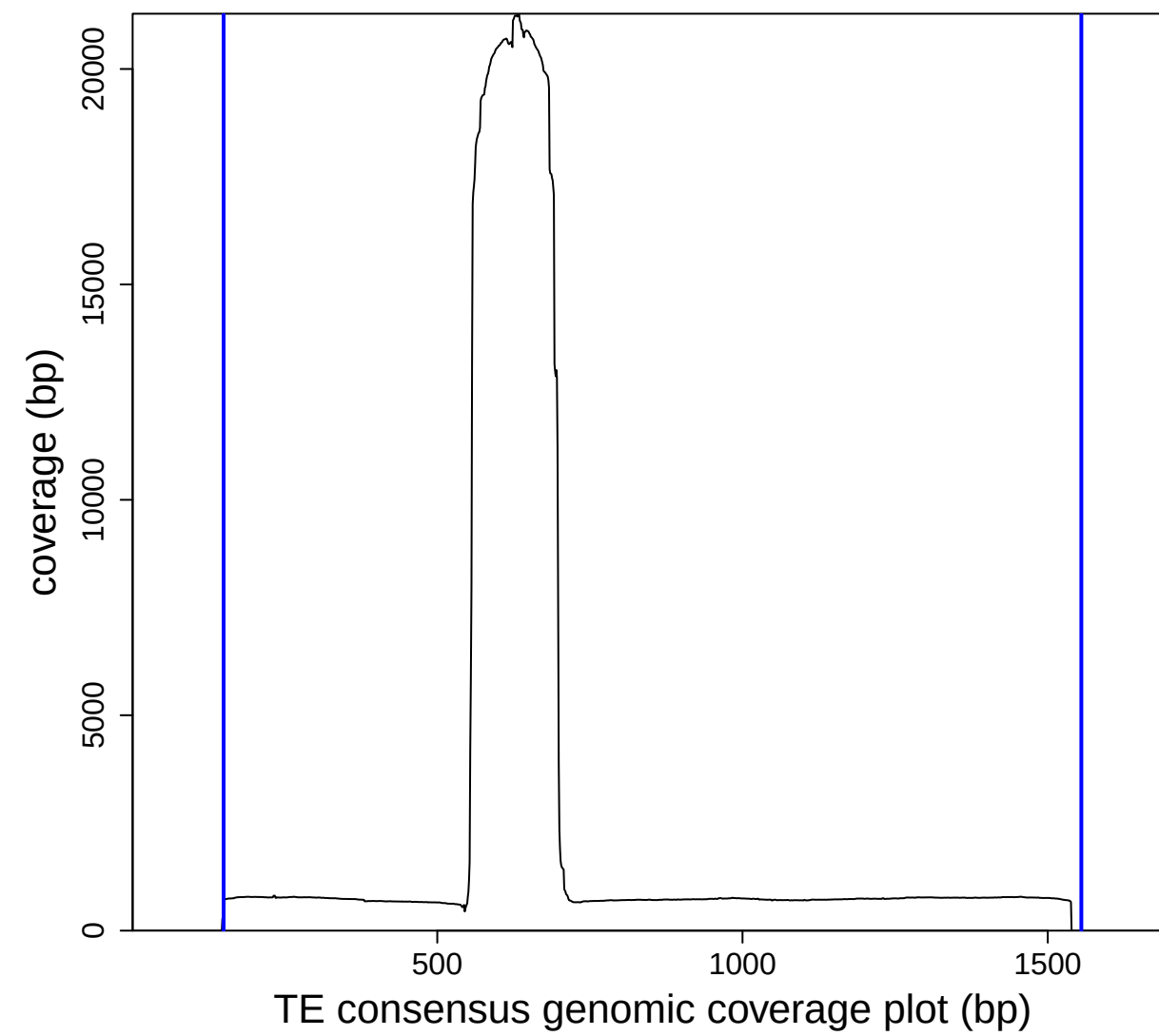
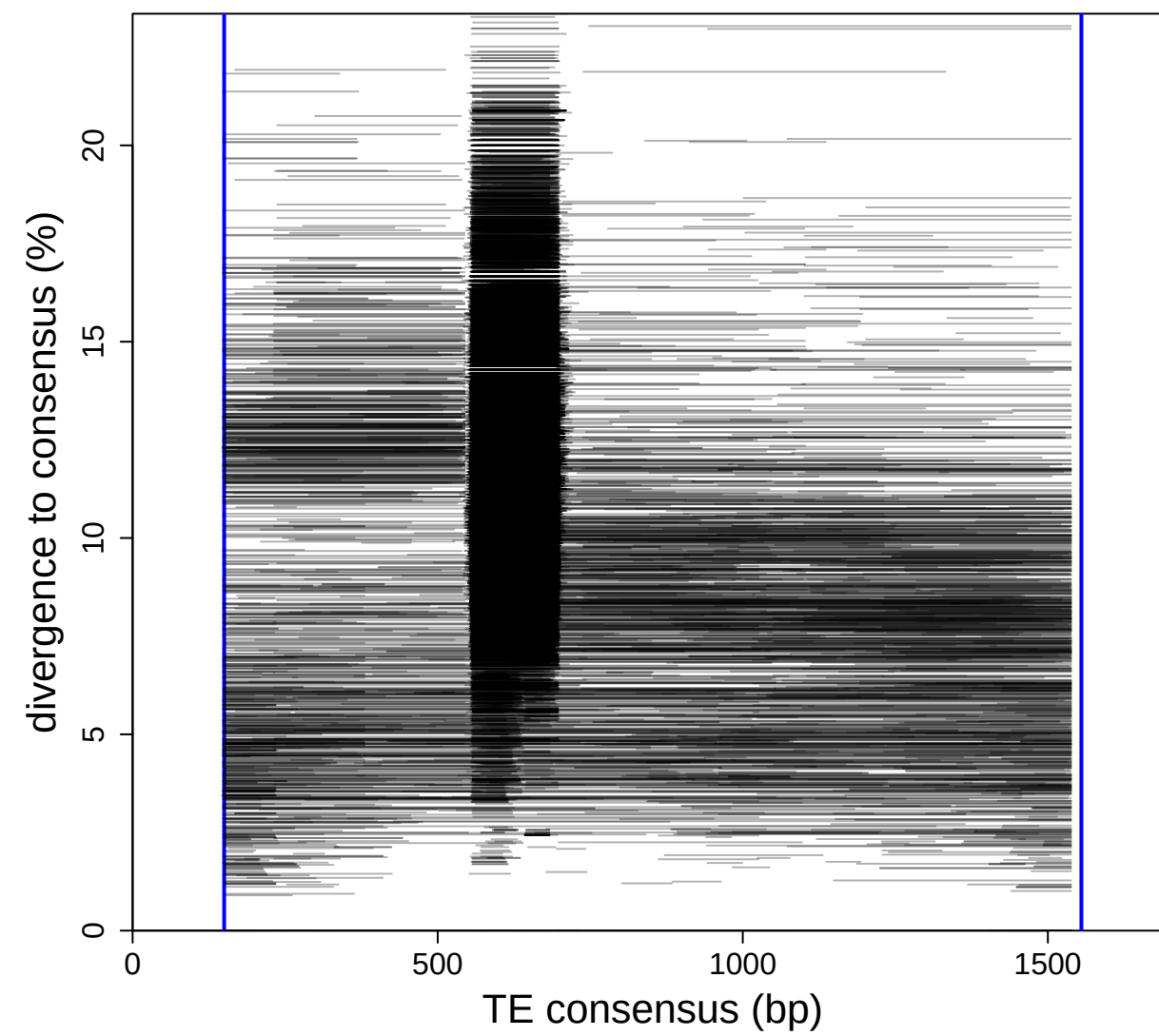
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 1399bp; fragments: 24227; full length: 74 (>=1259.1bp)

After TEtrimmer 1399 bp

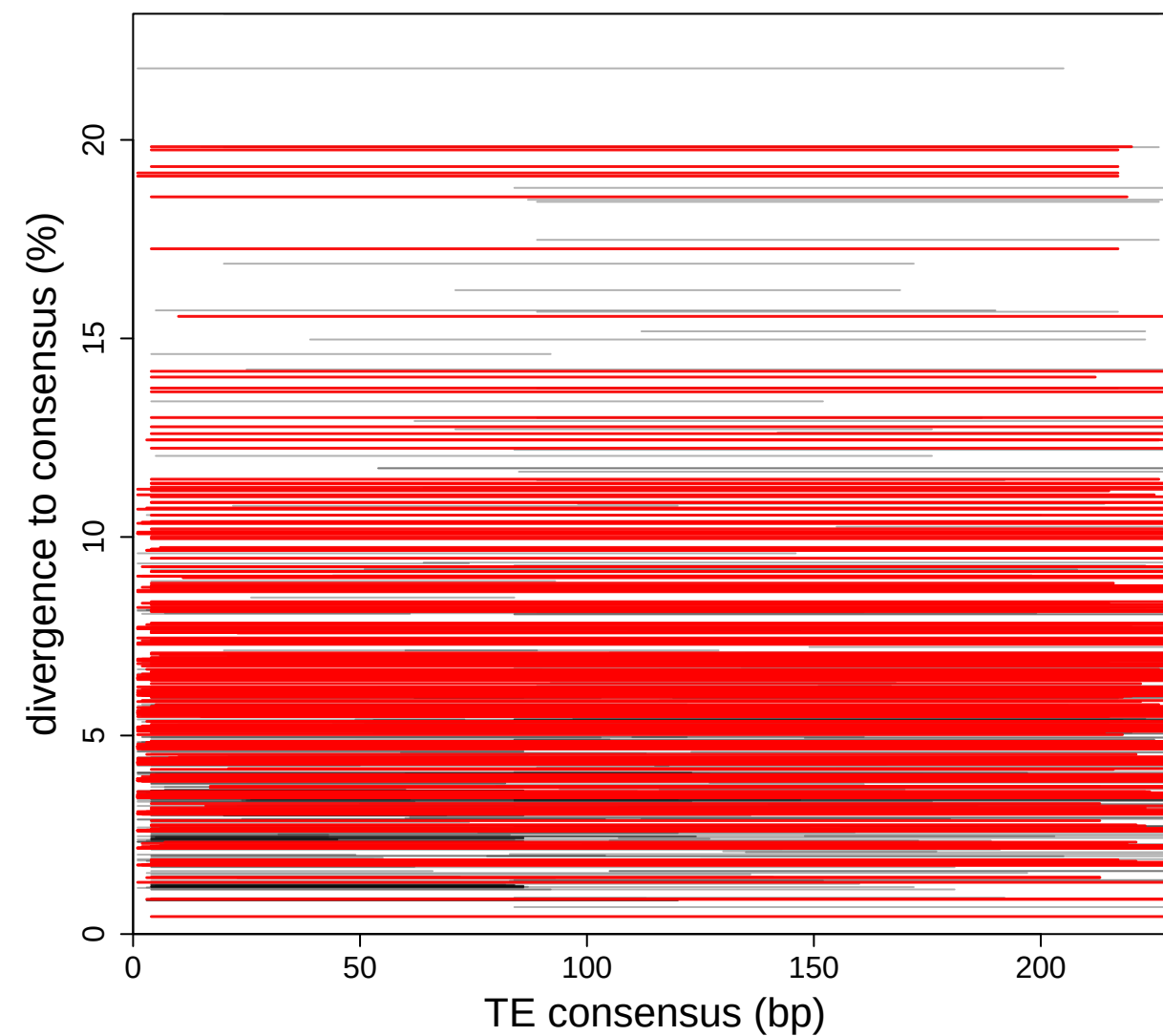


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 1705bp; fragments: 24227; full length: 0 (>=1534.5bp)

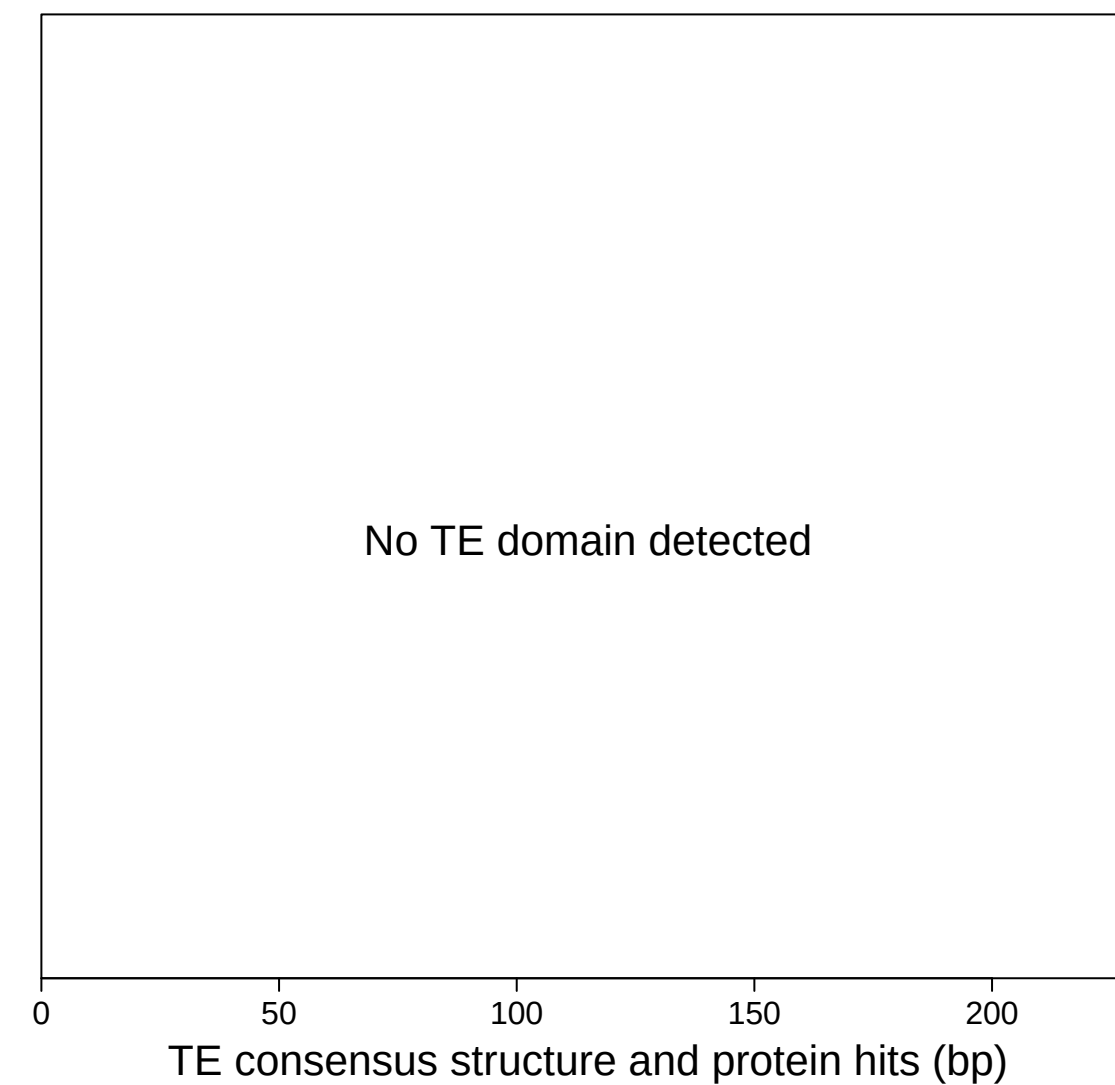
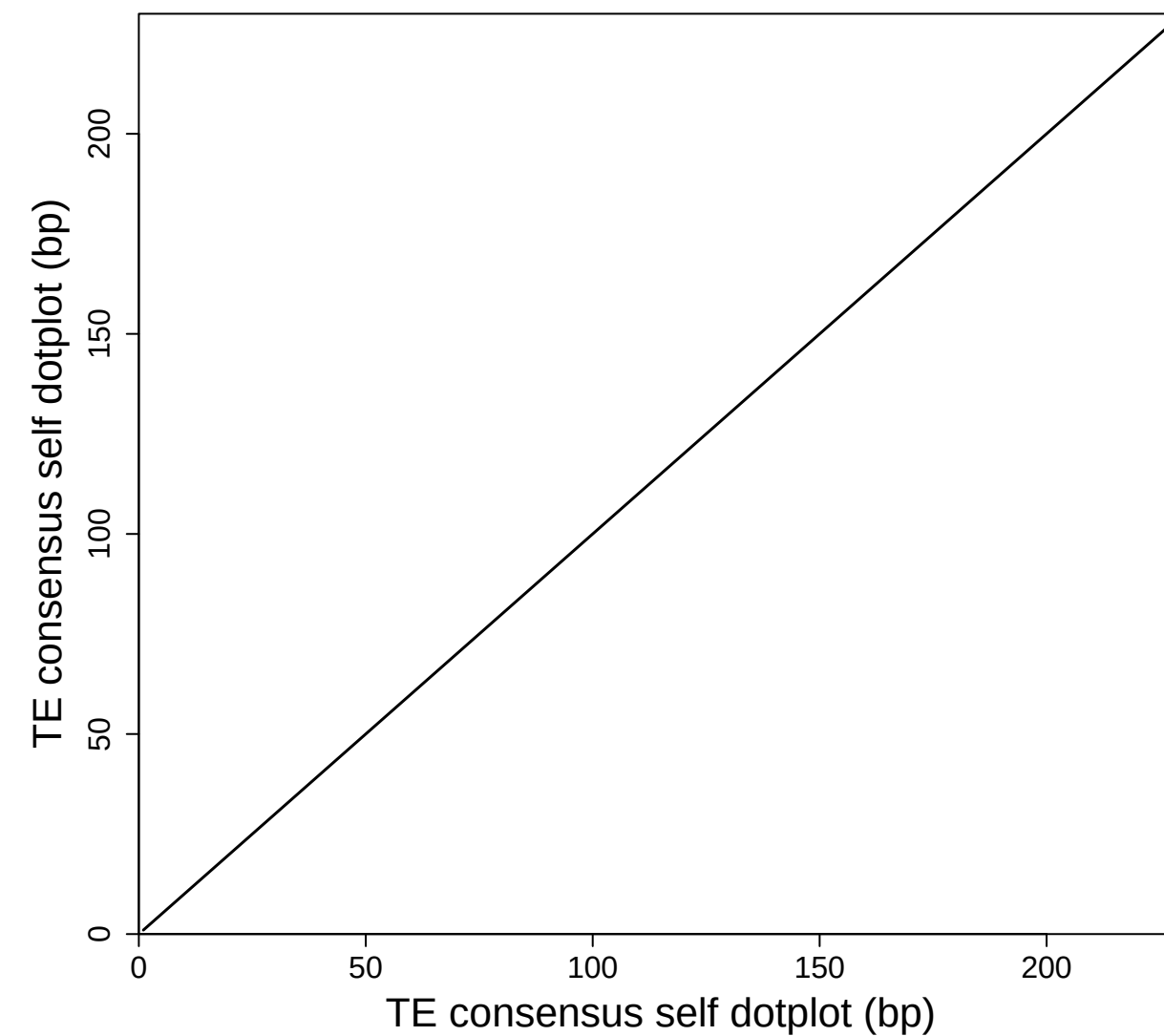
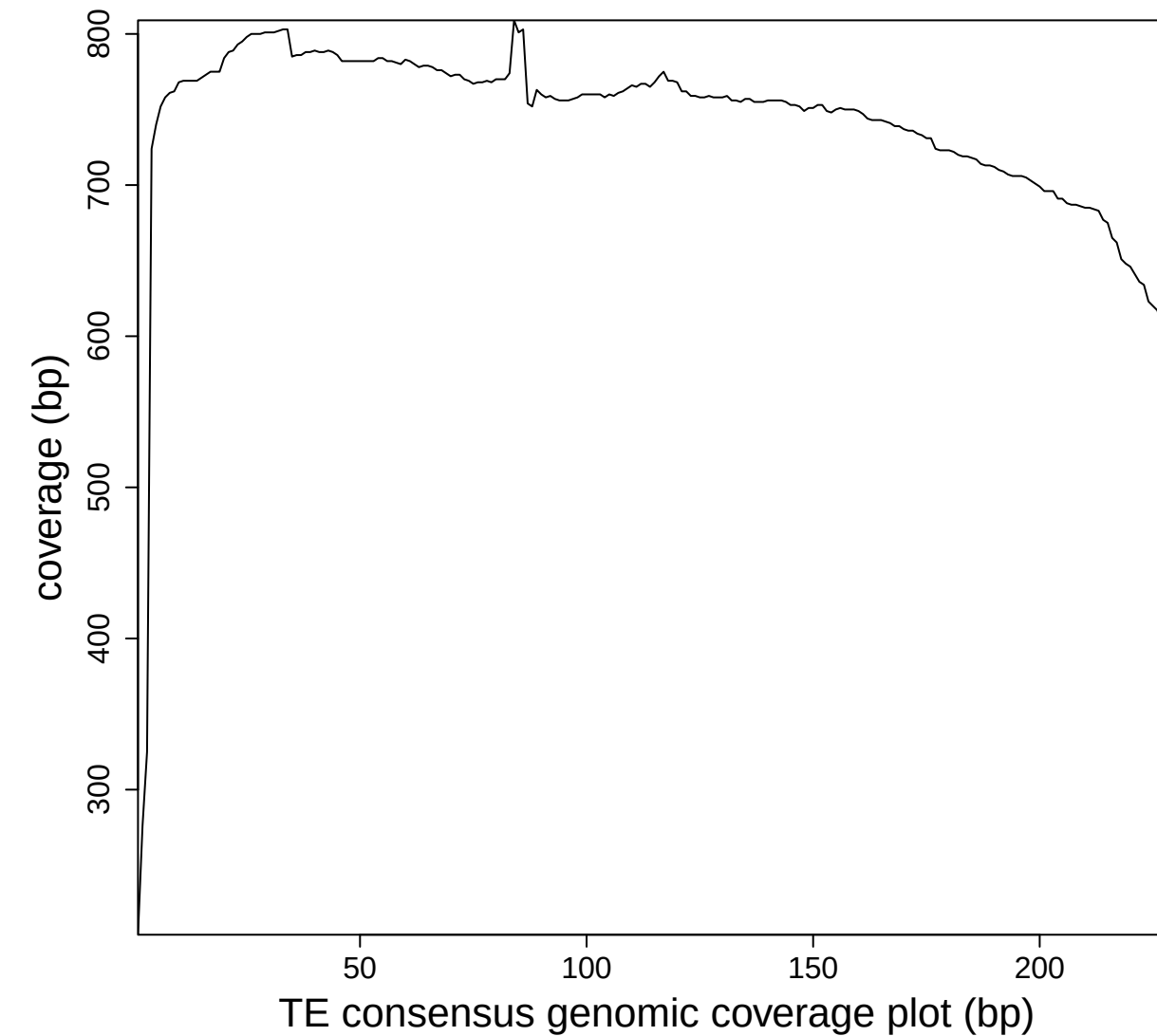
After TEtrimmer Extended plot Blue lines are boundaries



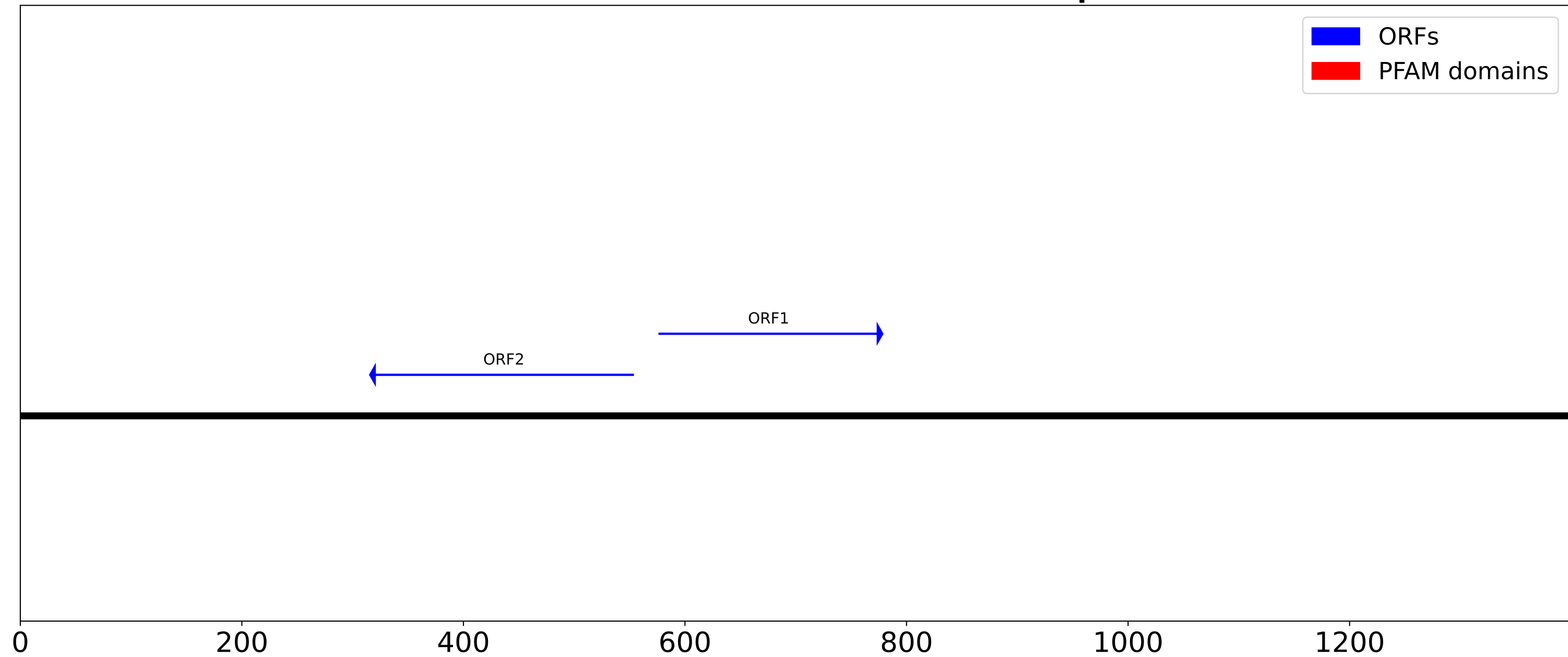
TE: rnd_1_family_567
size: 230bp; fragments: 969; full length: 536 (≥ 207 bp)



Before TEtrimmer 230 bp



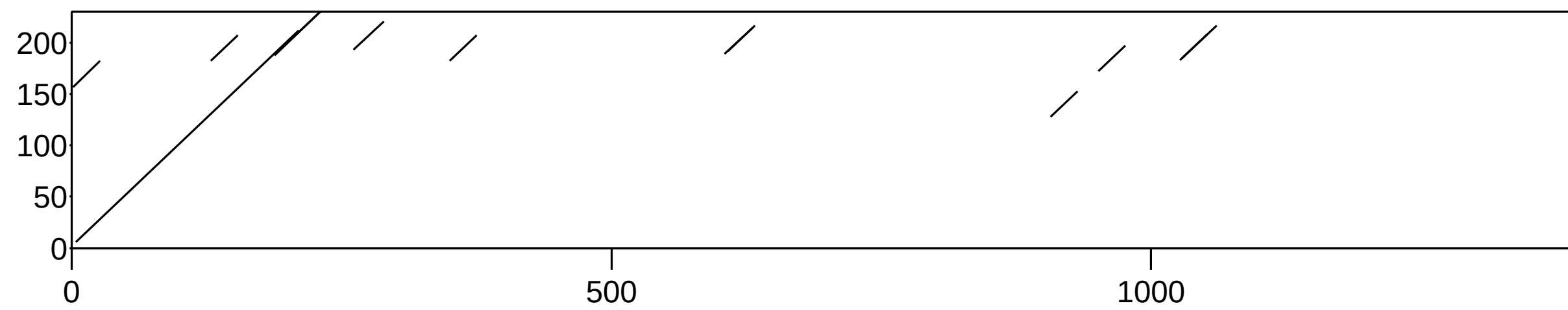
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)