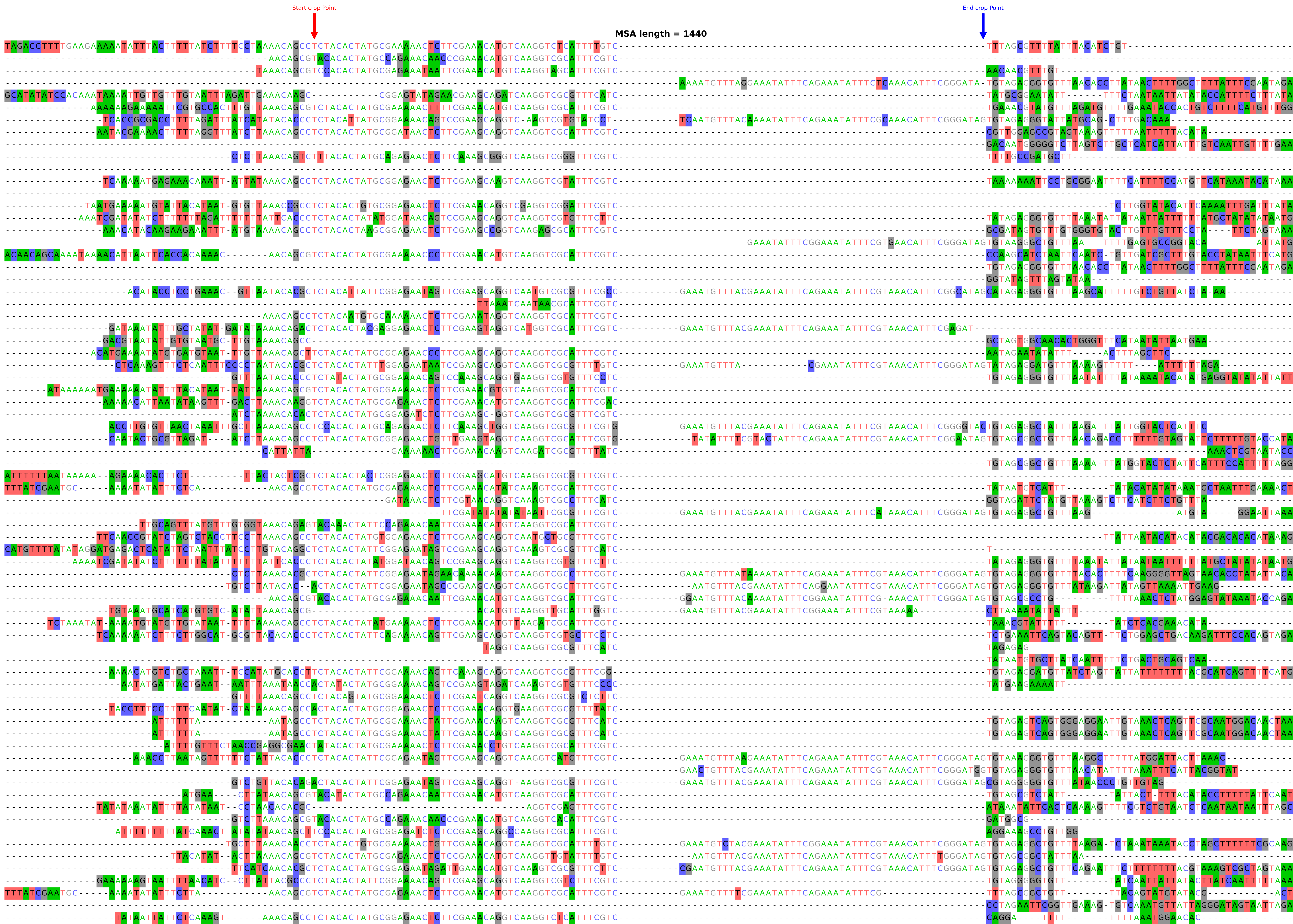
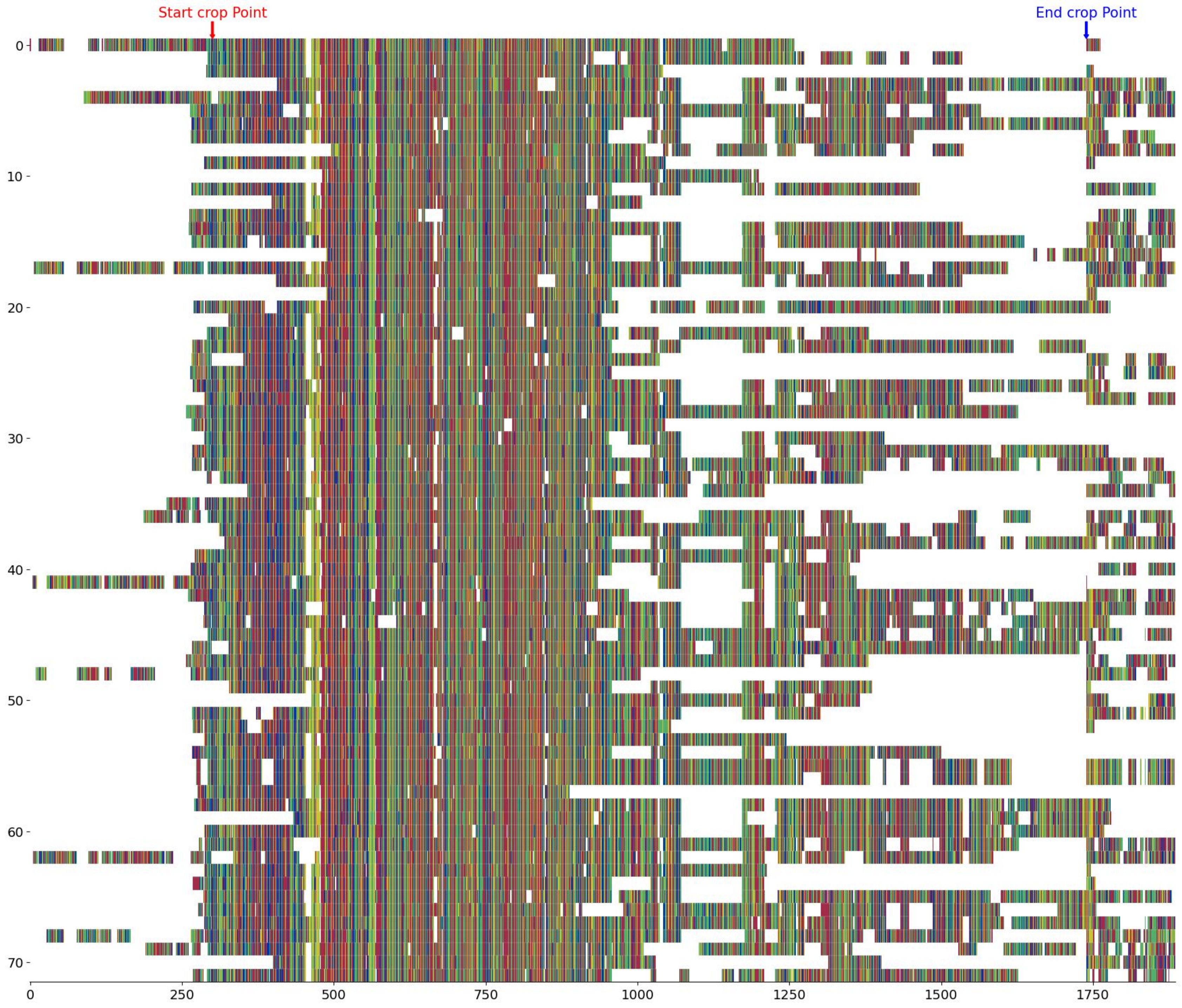
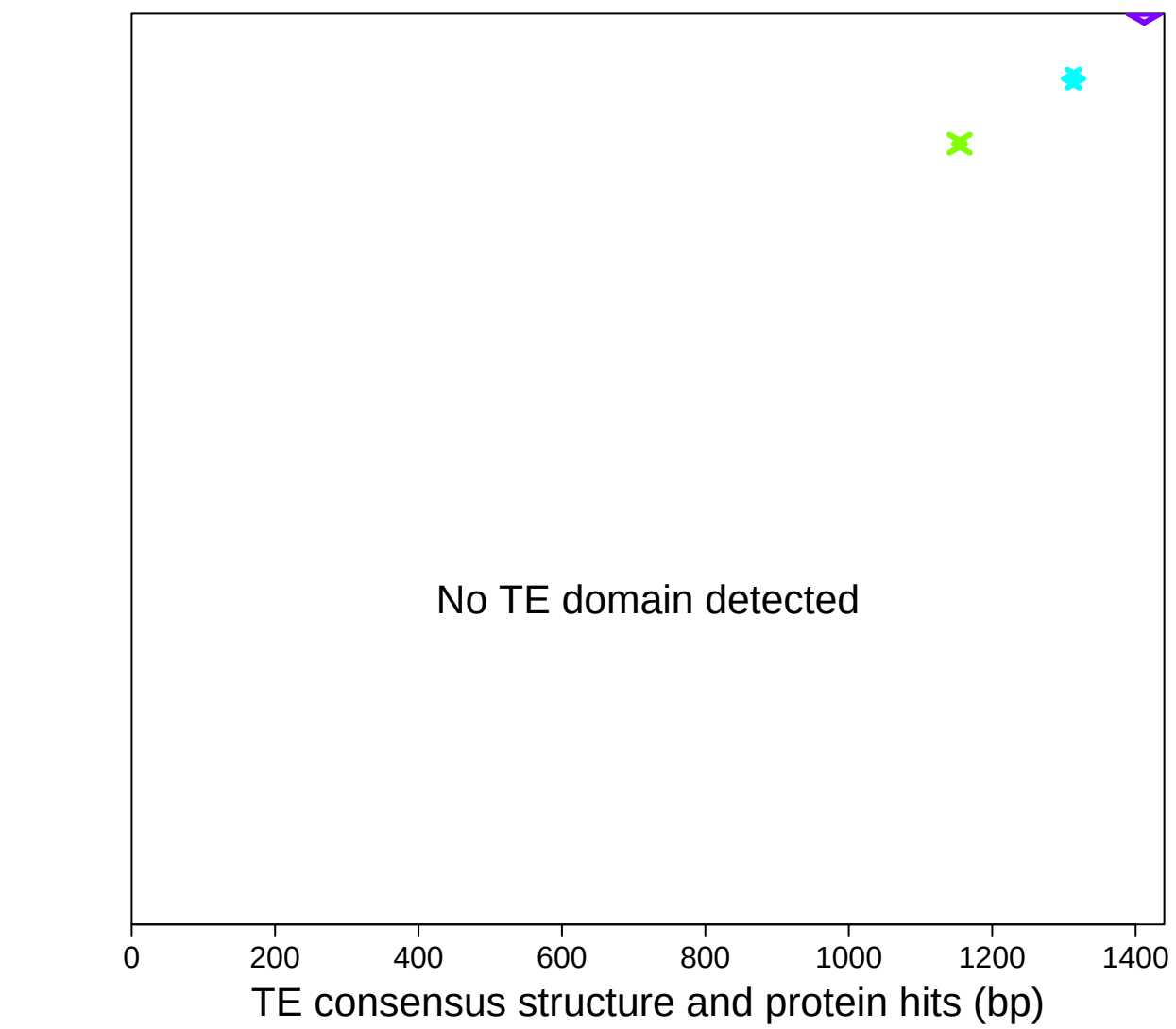
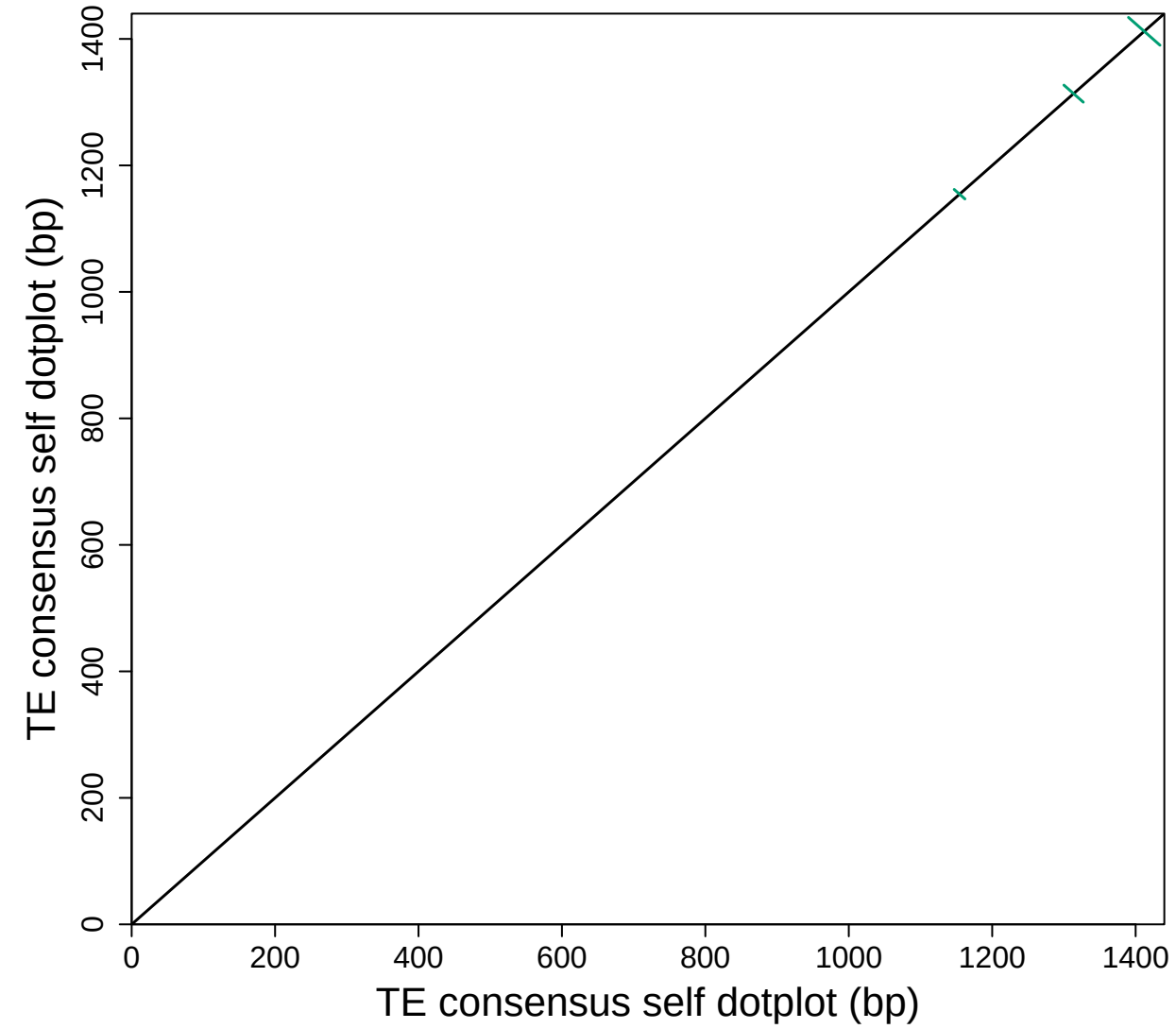
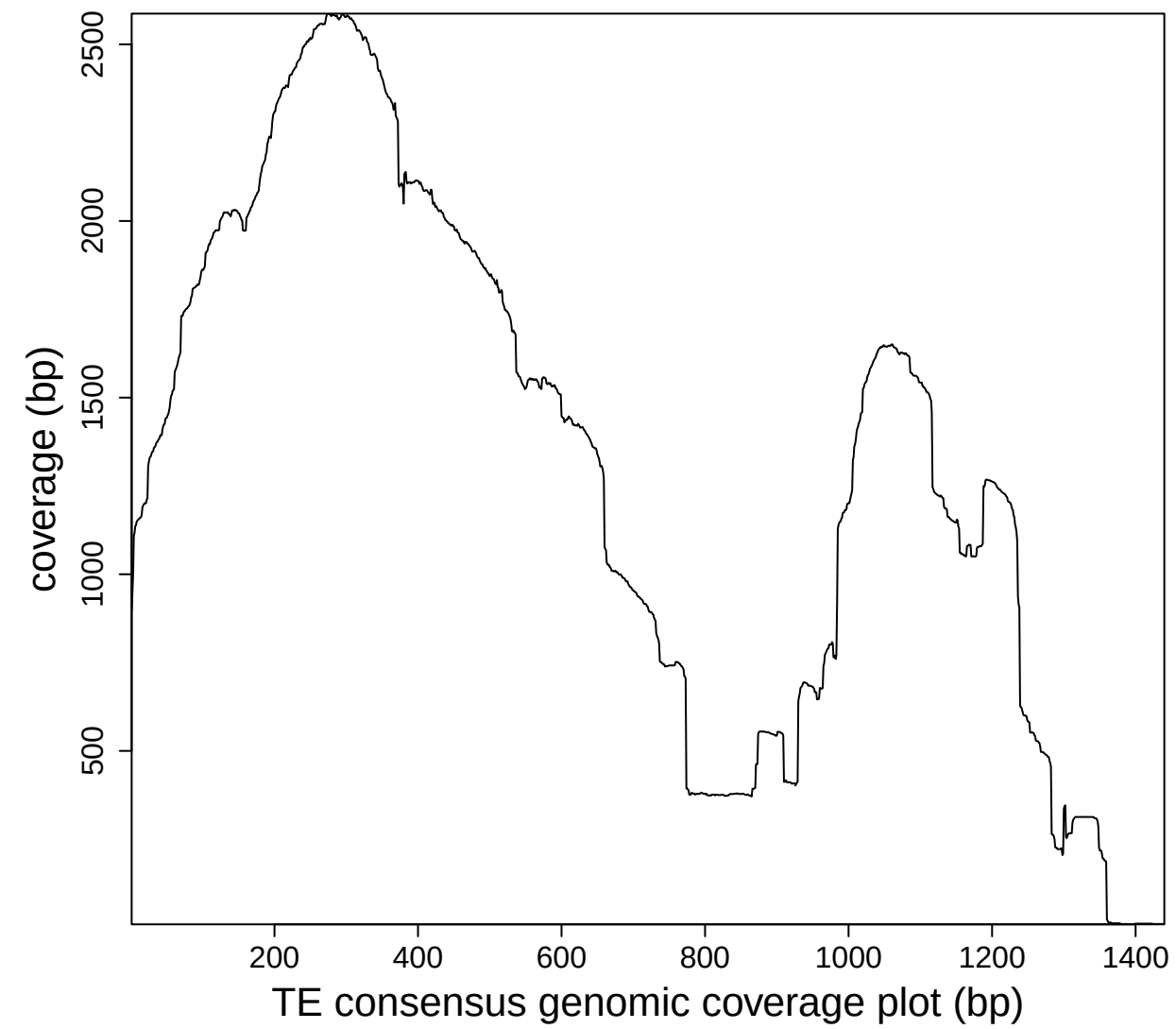
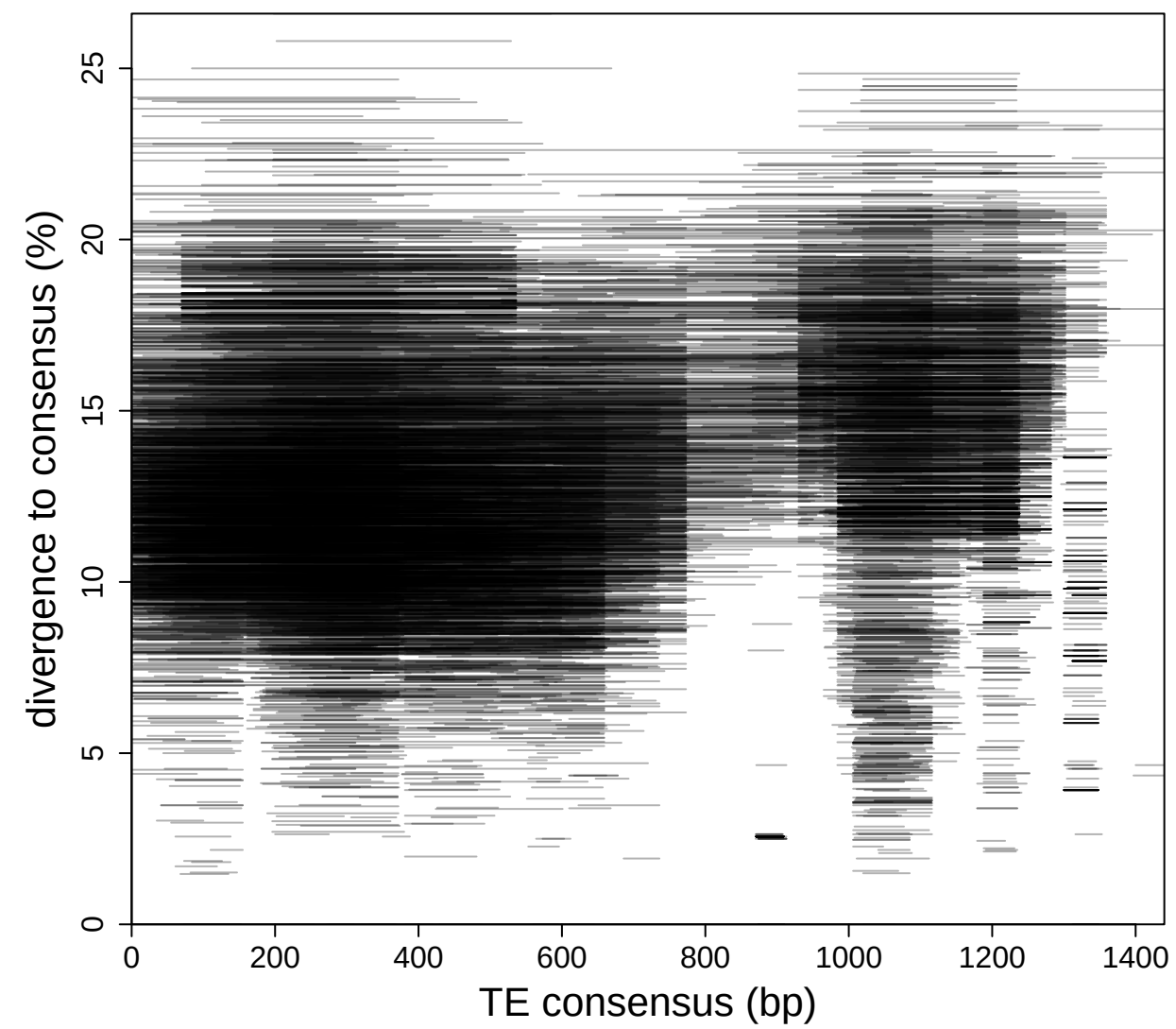






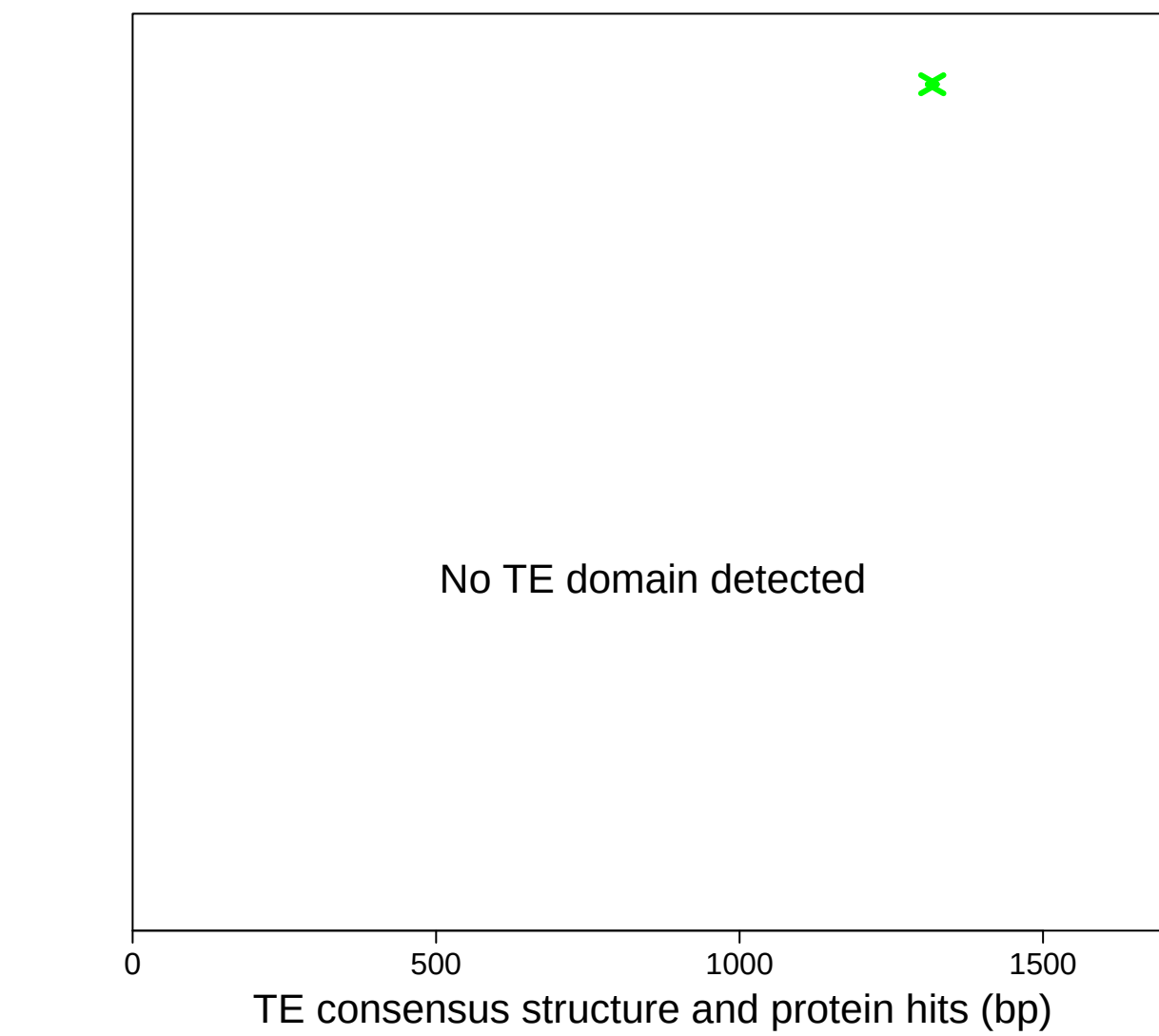
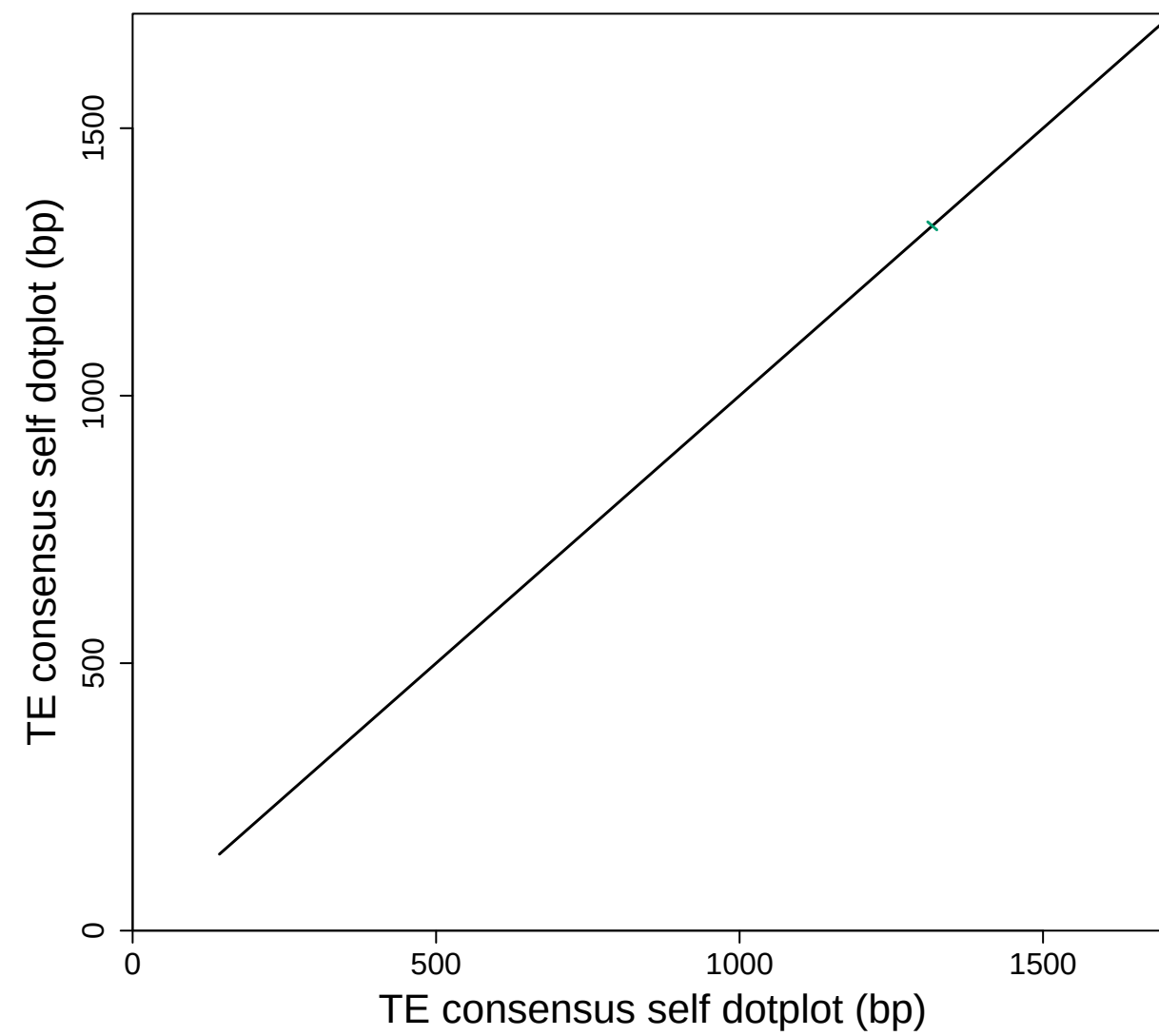
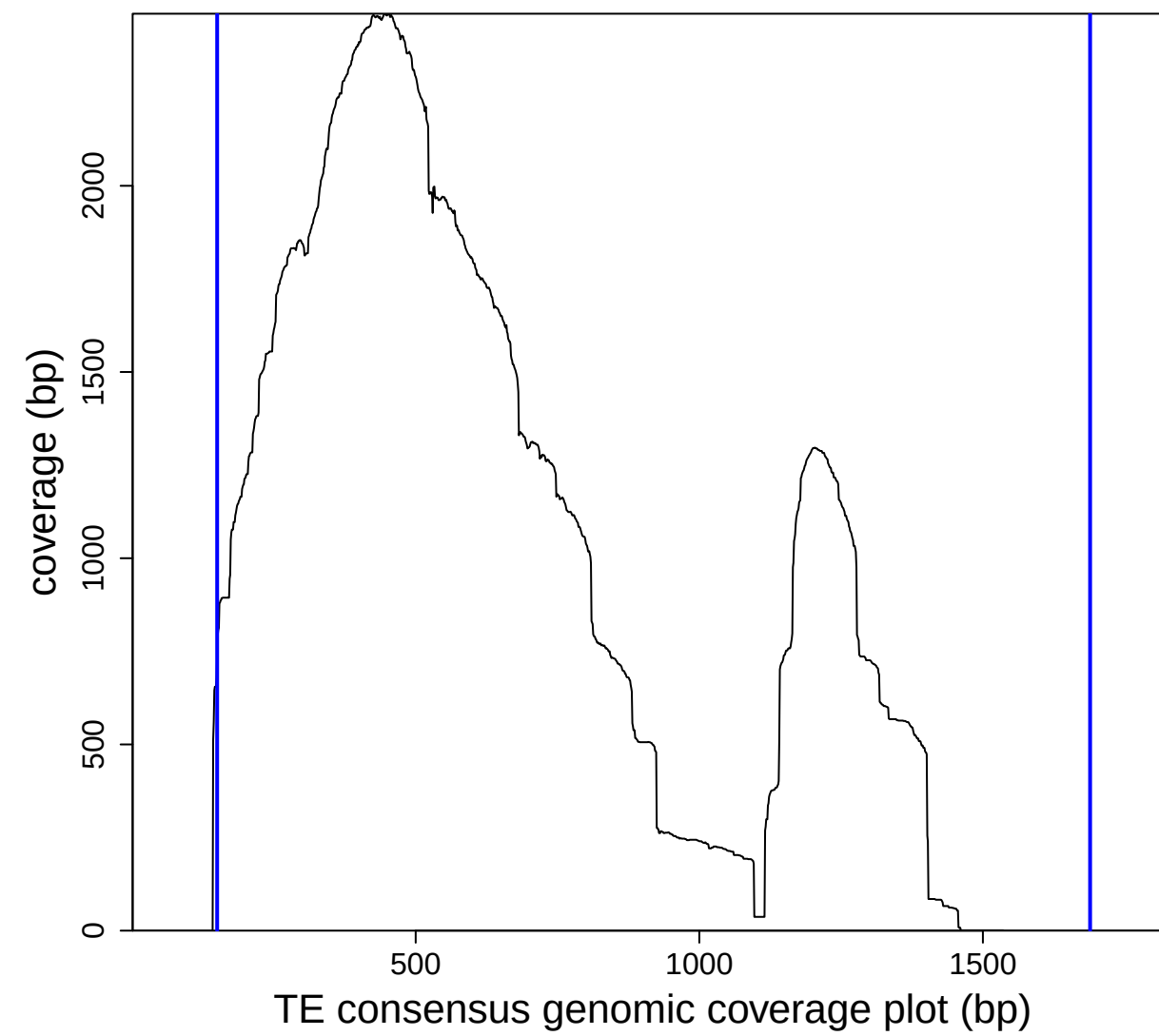
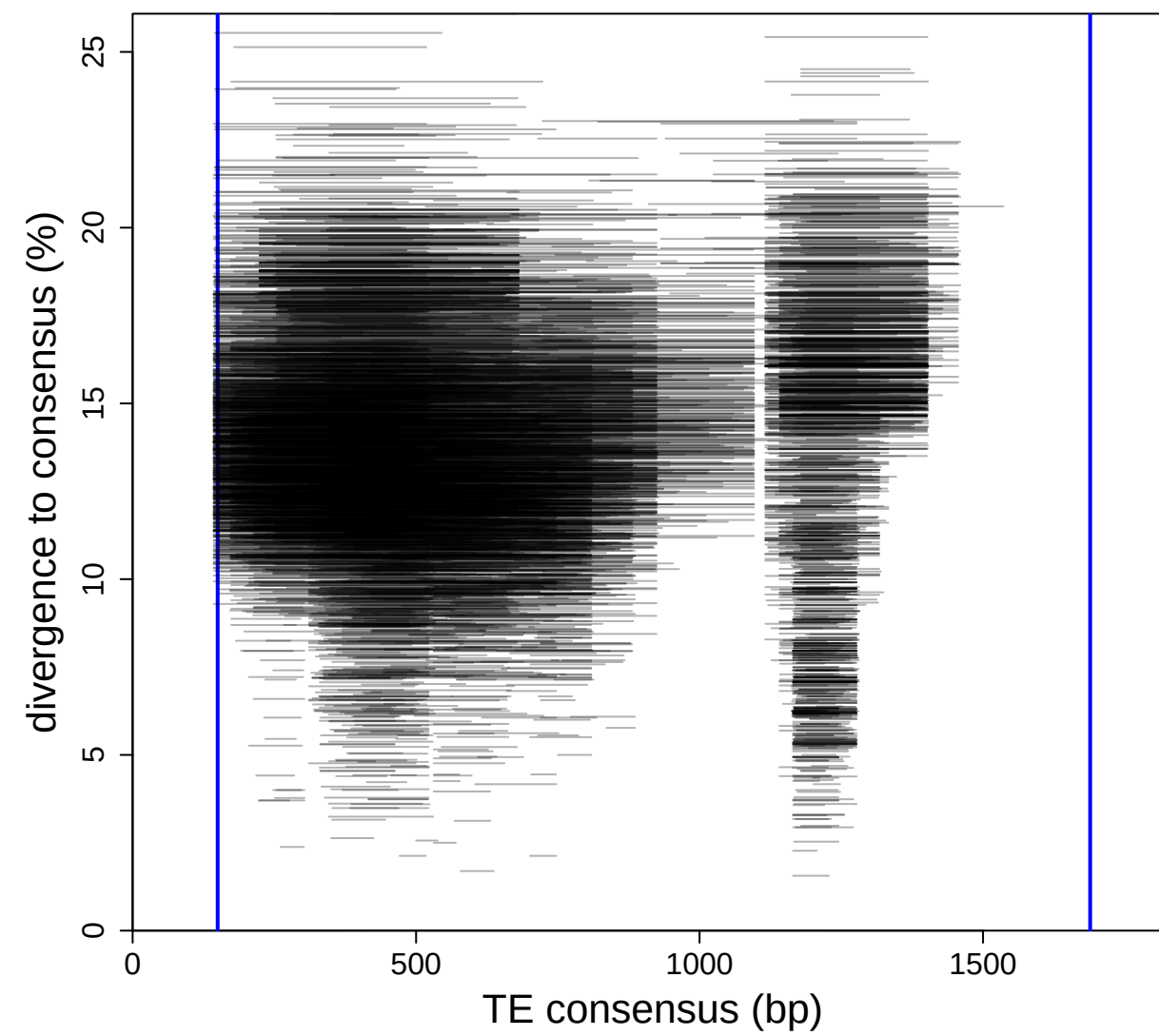
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 1440bp; fragments: 6028; full length: 0 (>=1296bp)

After TEtrimmer 1440 bp



f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 1835bp; fragments: 4455; full length: 0 (>=1651.5bp)

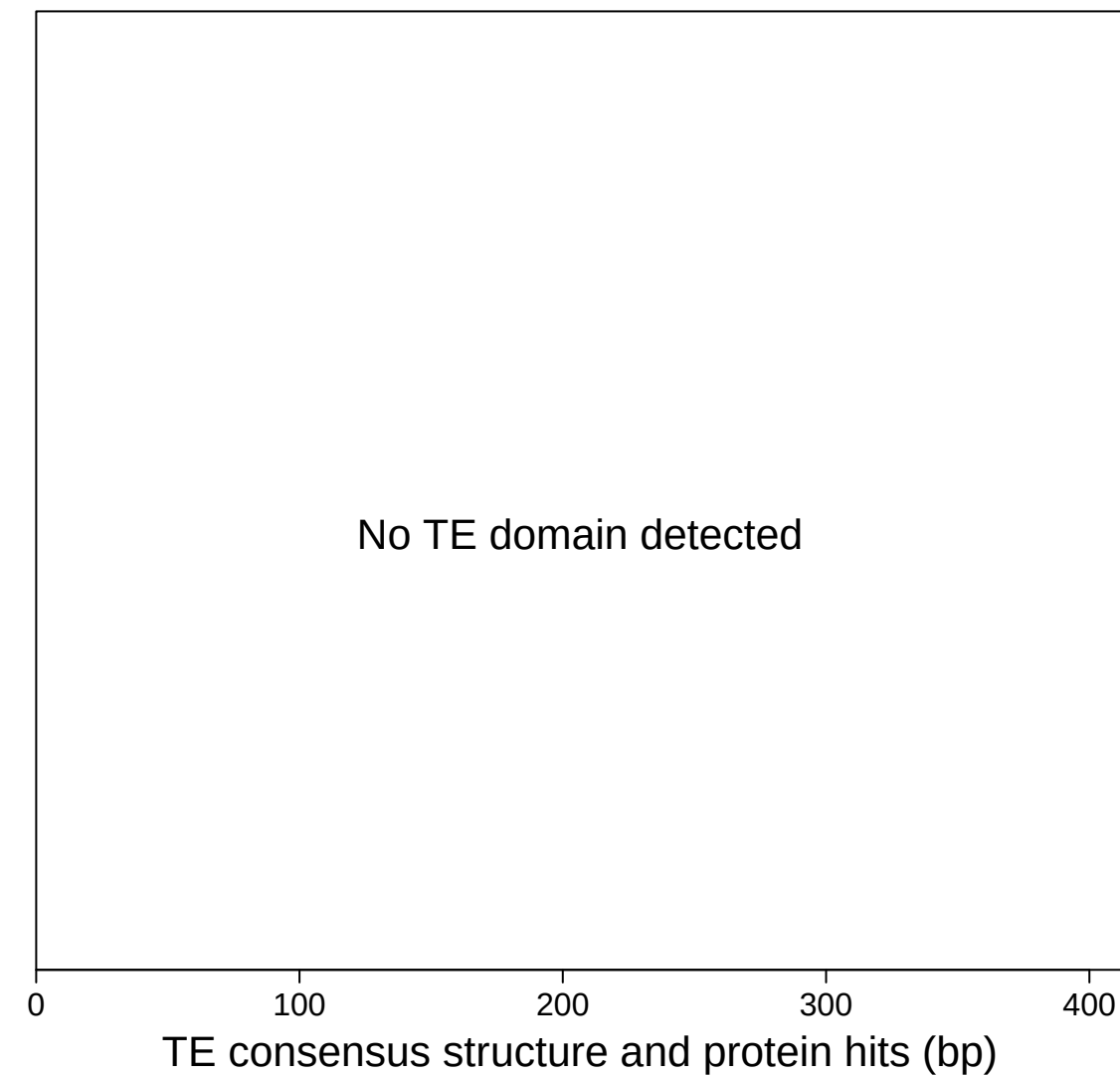
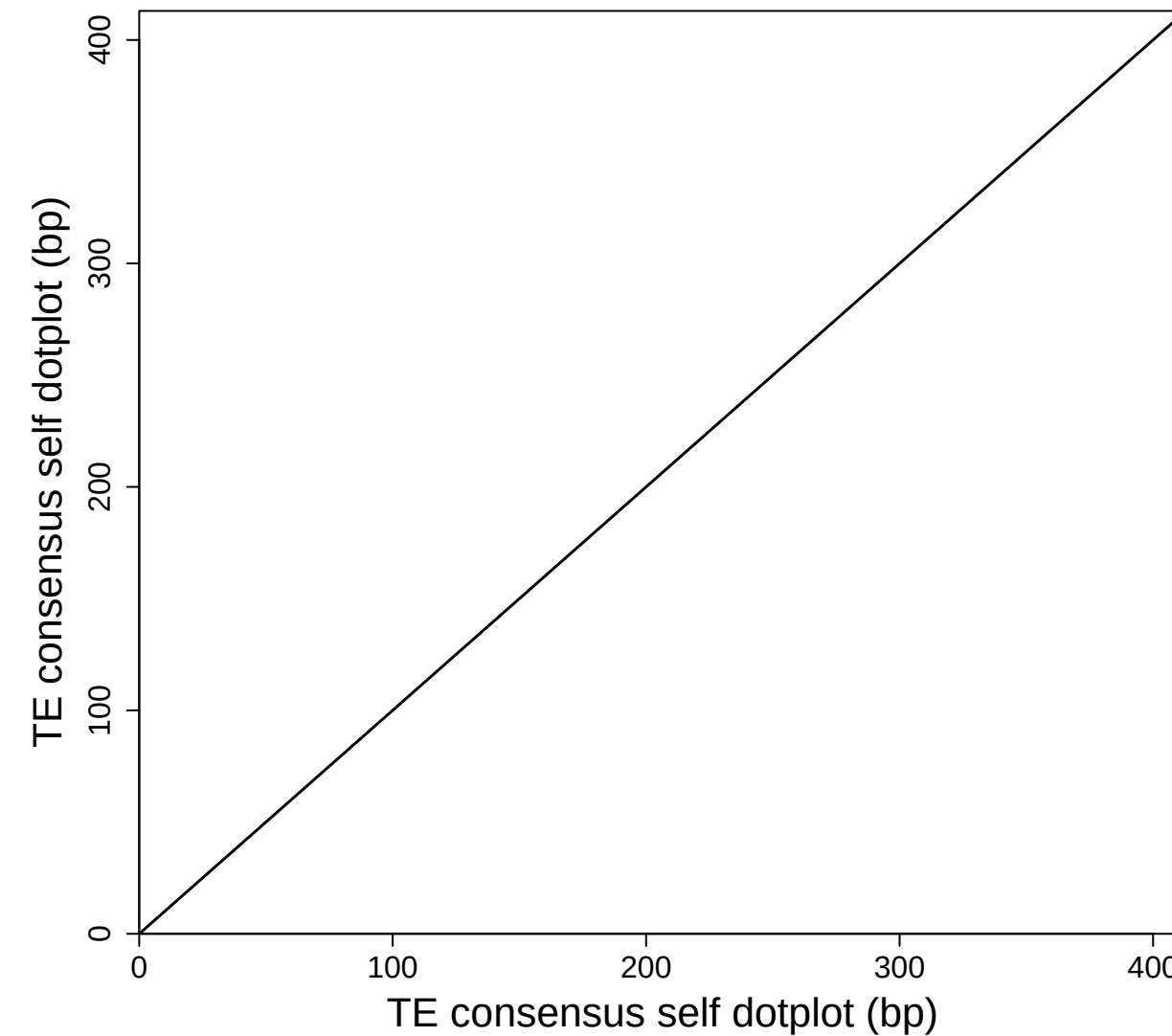
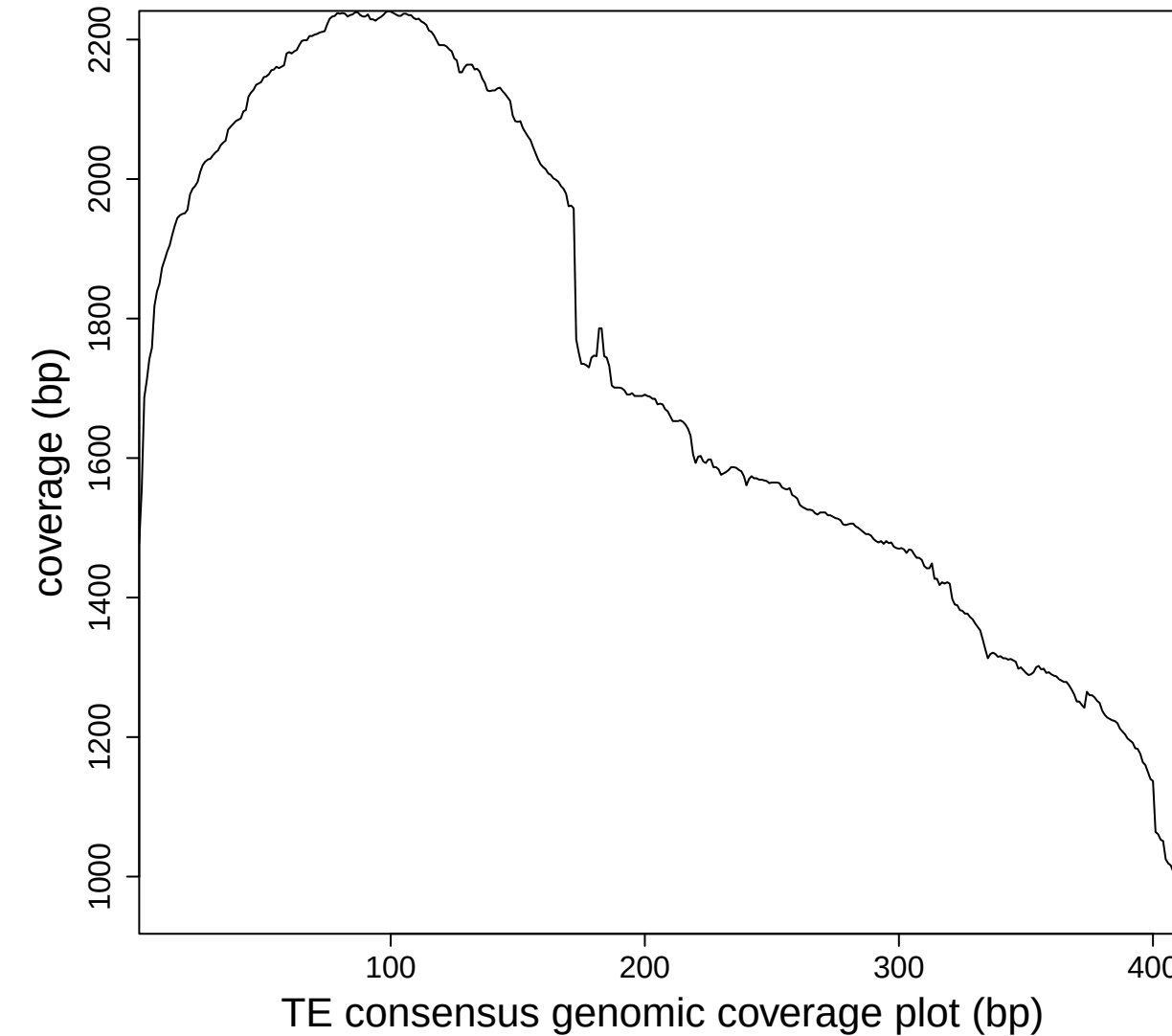
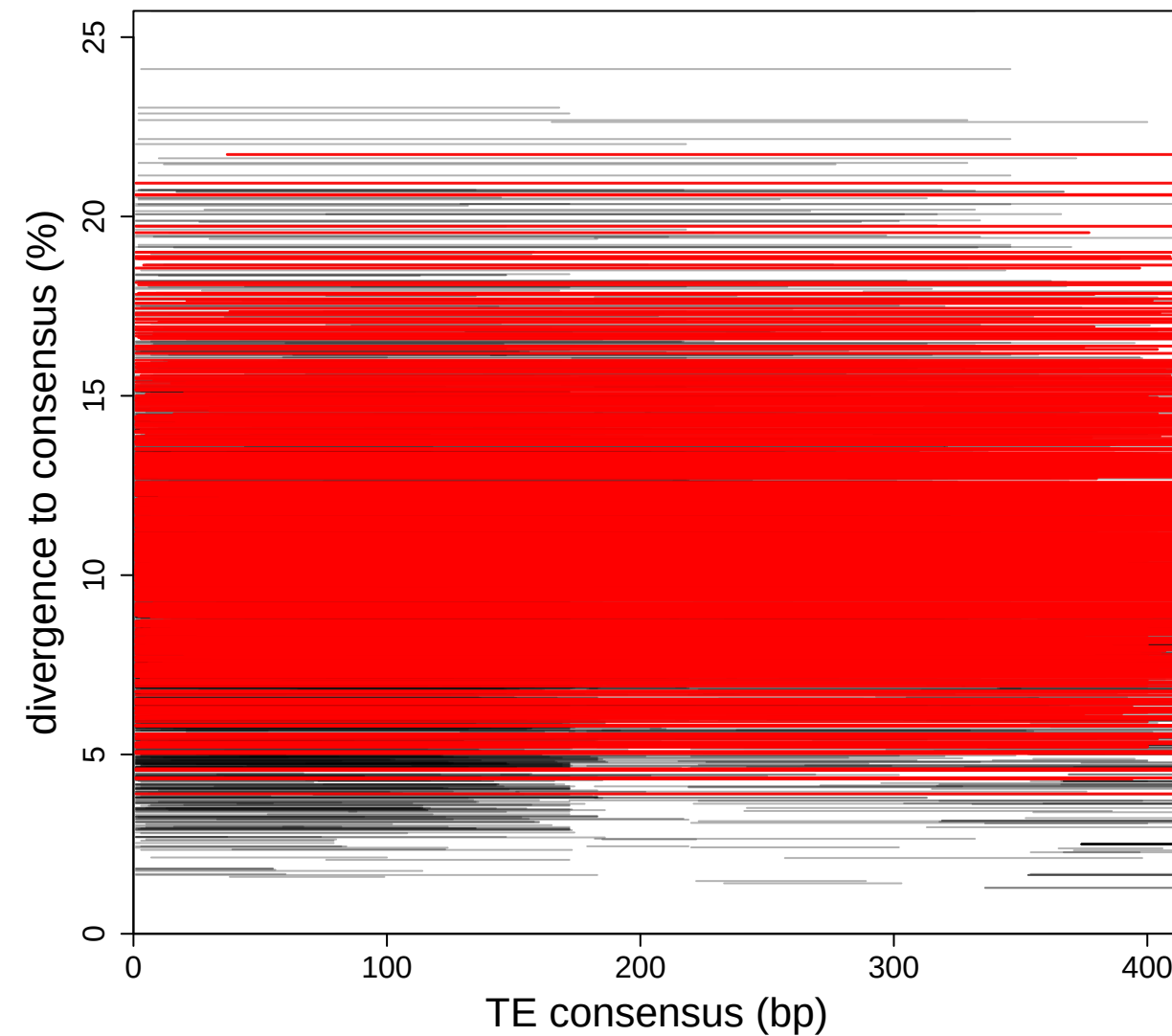
After TEtrimmer Extended plot Blue lines are boundaries



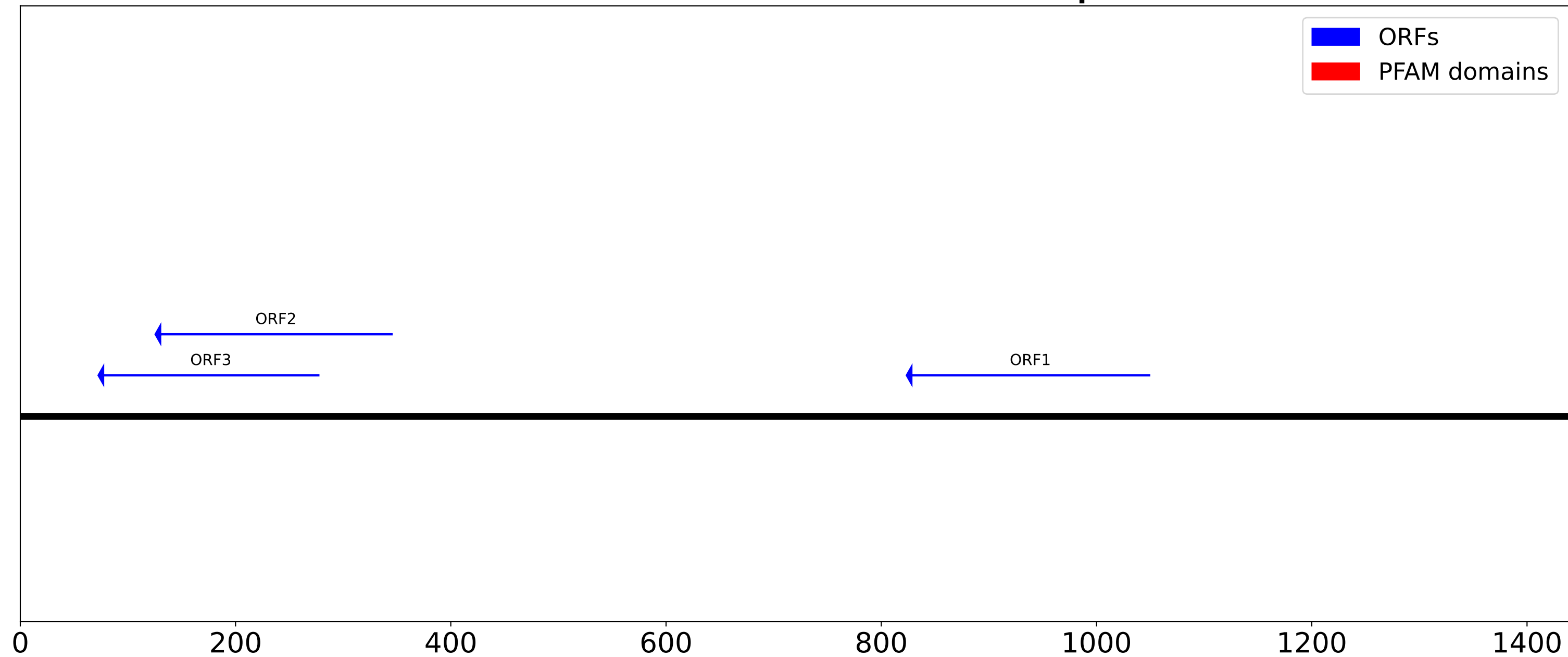
Before TEtrimmer 413 bp

TE: rnd_1_family_746

size: 413bp; fragments: 2811; full length: 737 (≥ 371.7 bp)



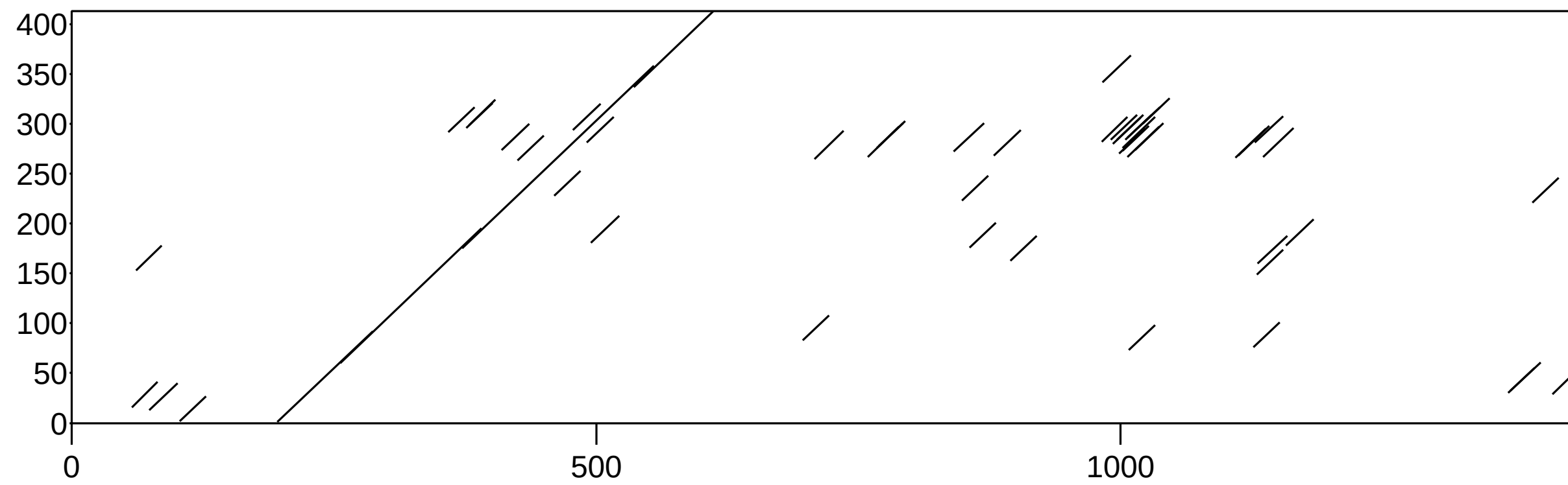
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)