

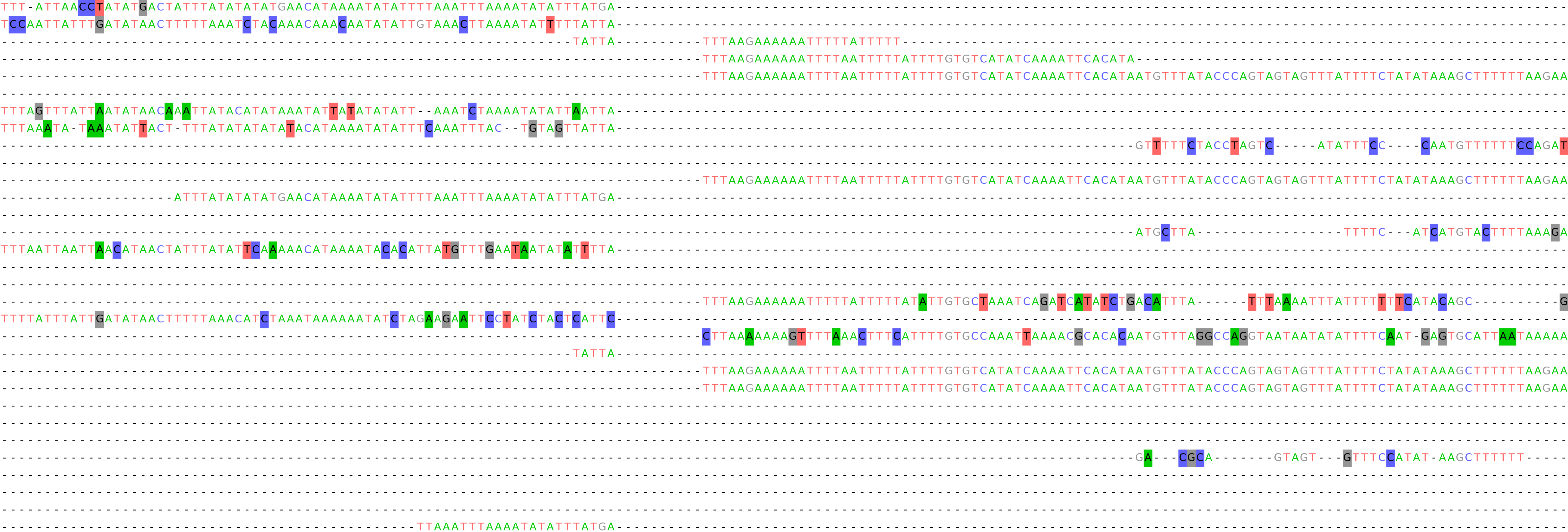
Start crop Point

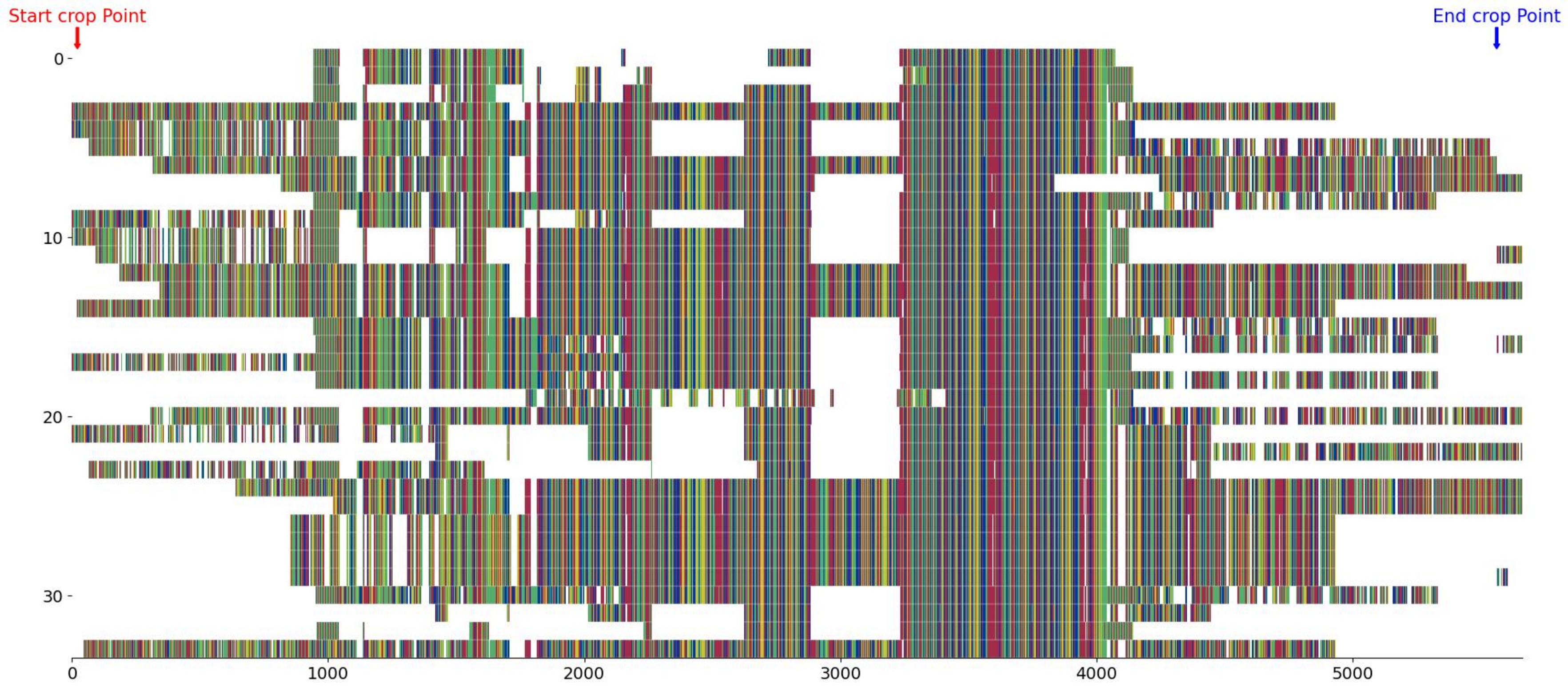


End crop Point



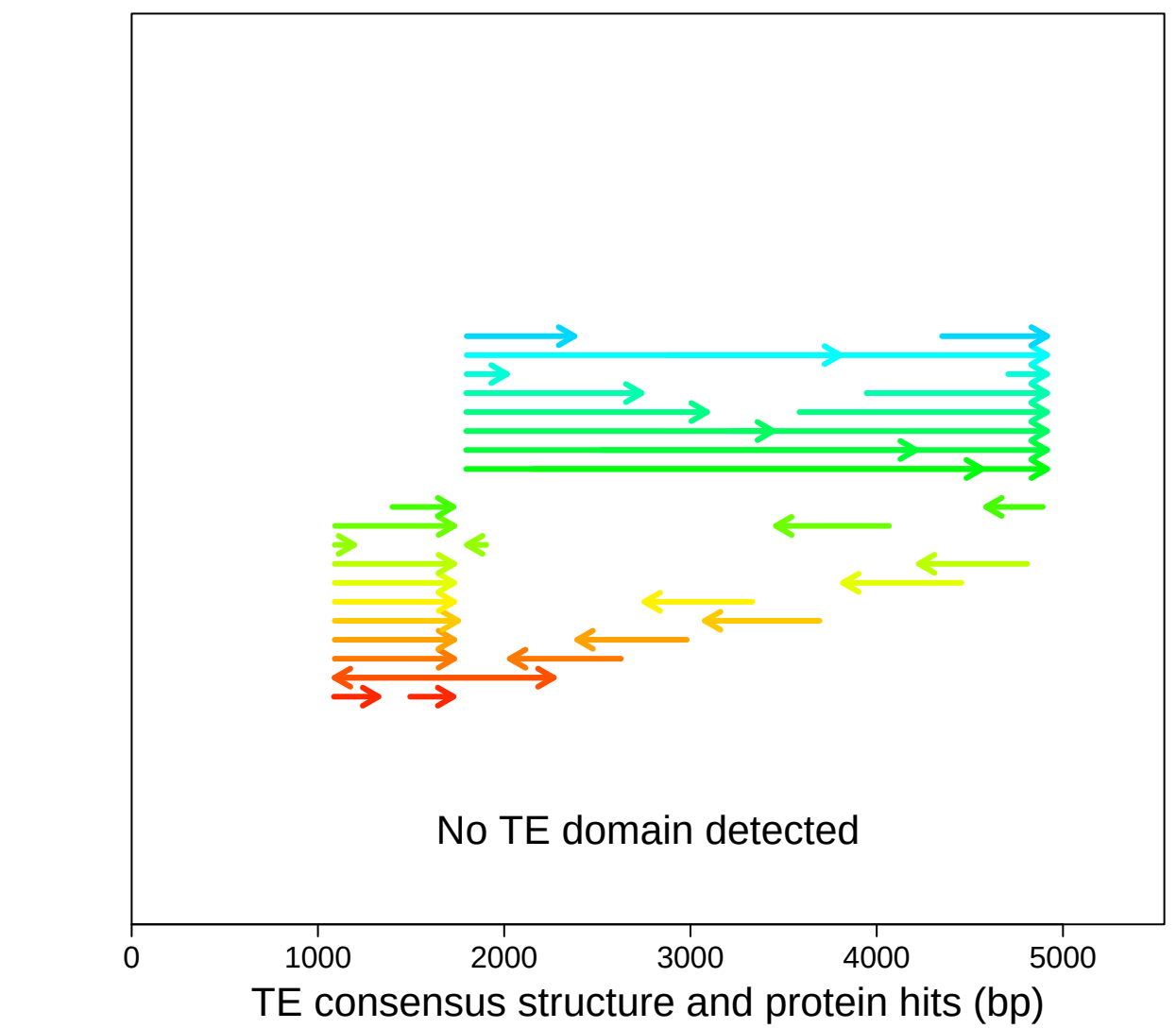
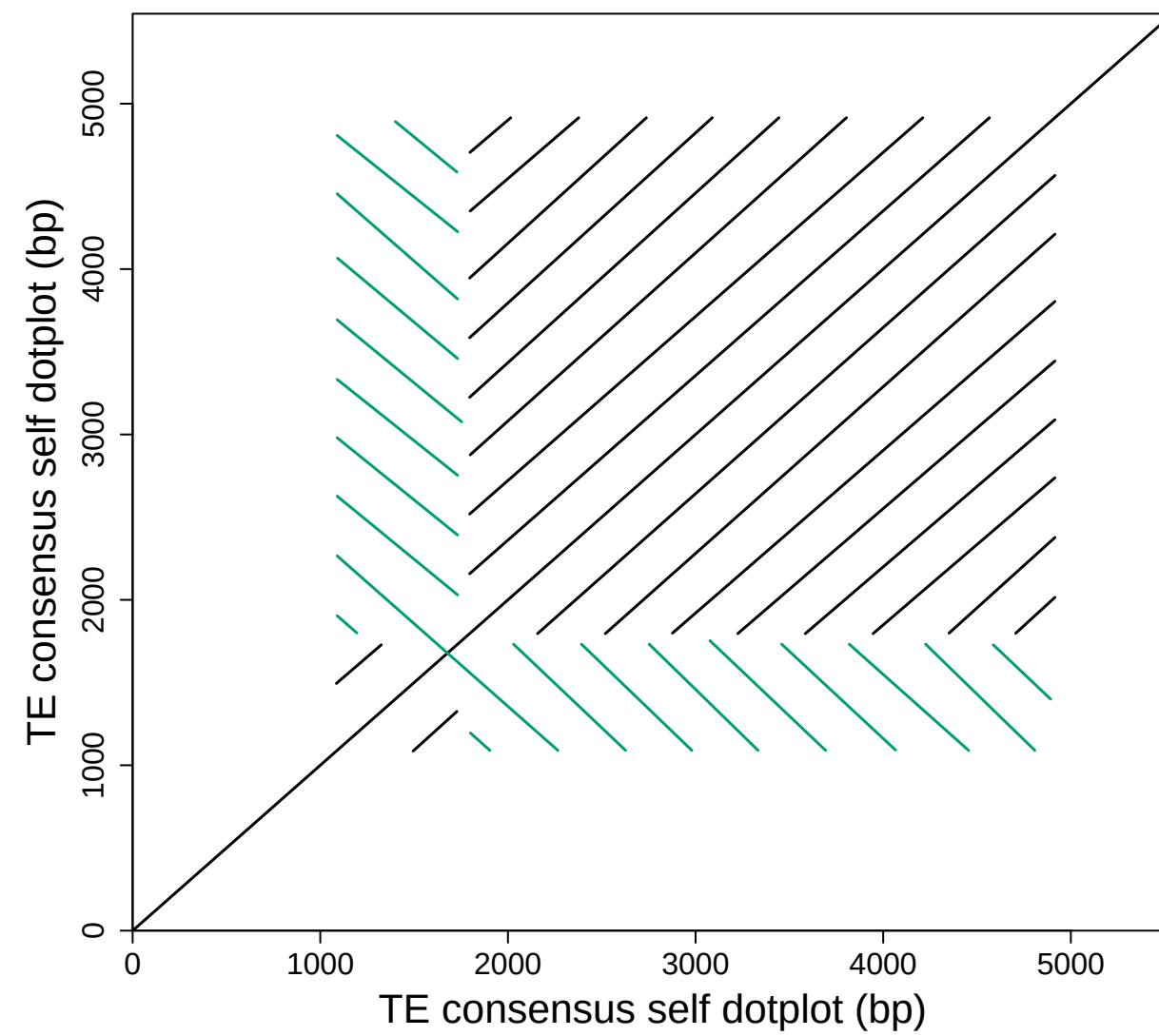
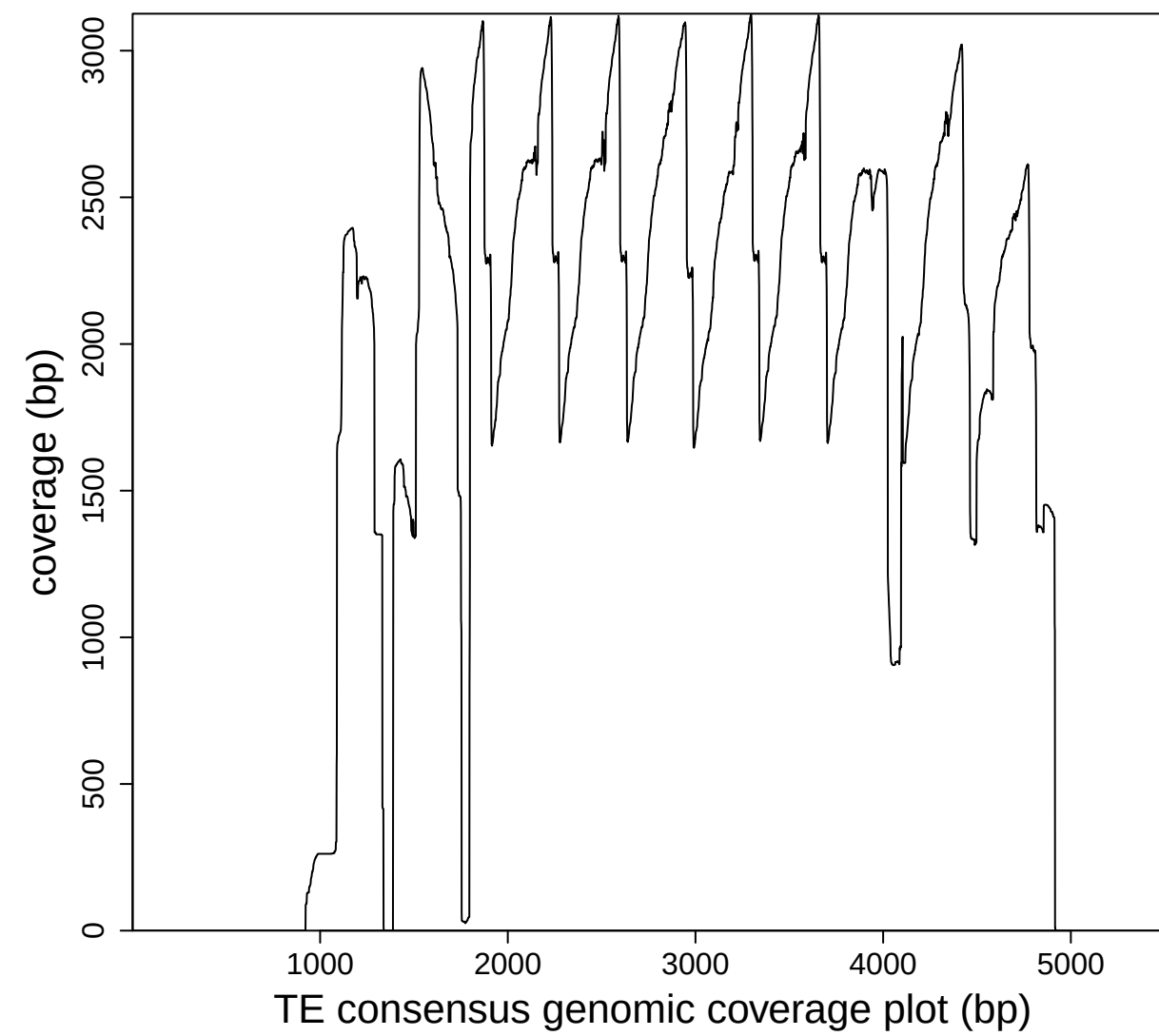
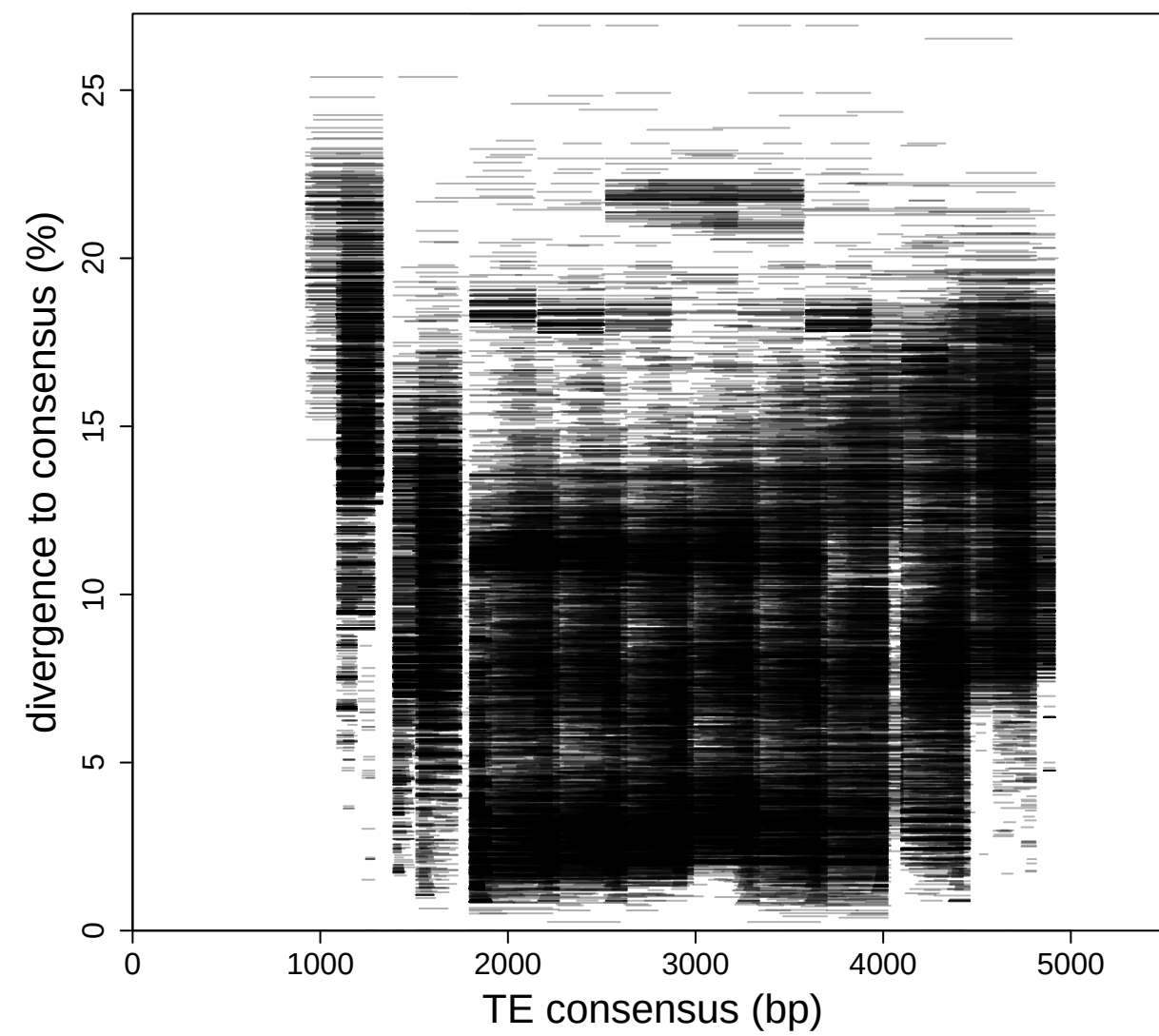
MSA length = 5544





g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 5544bp; fragments: 27609; full length: 0 (>=4989.6bp)

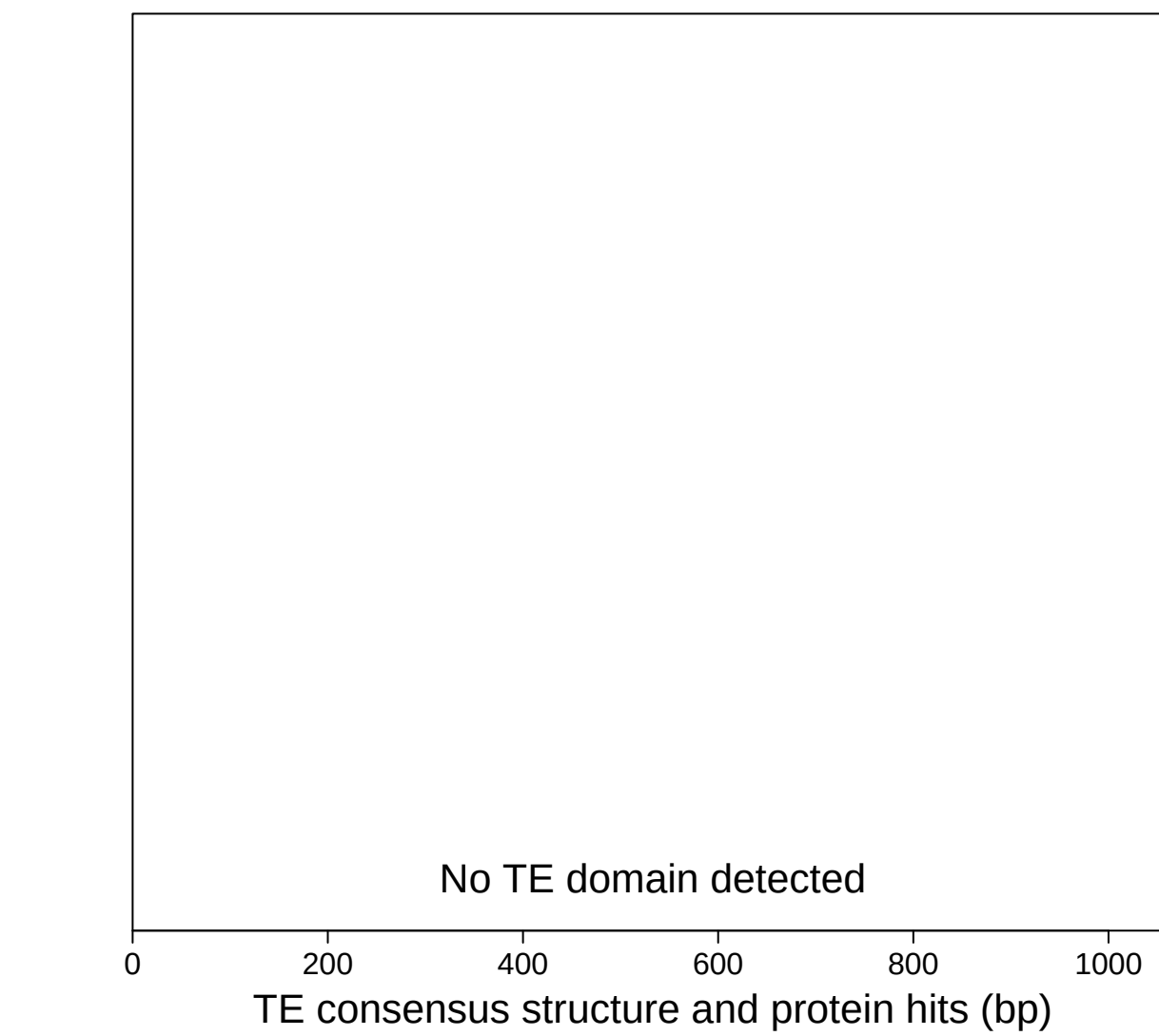
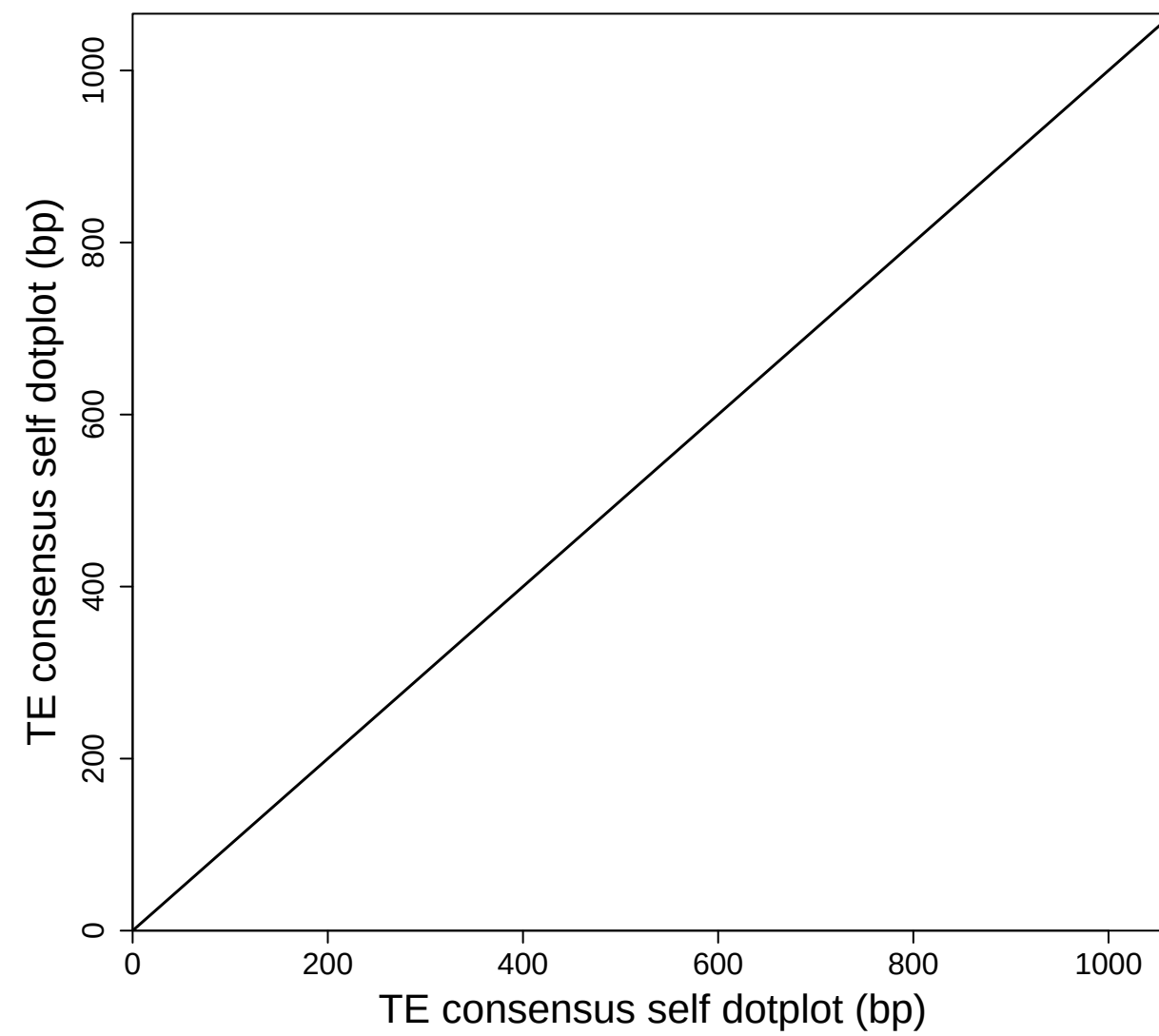
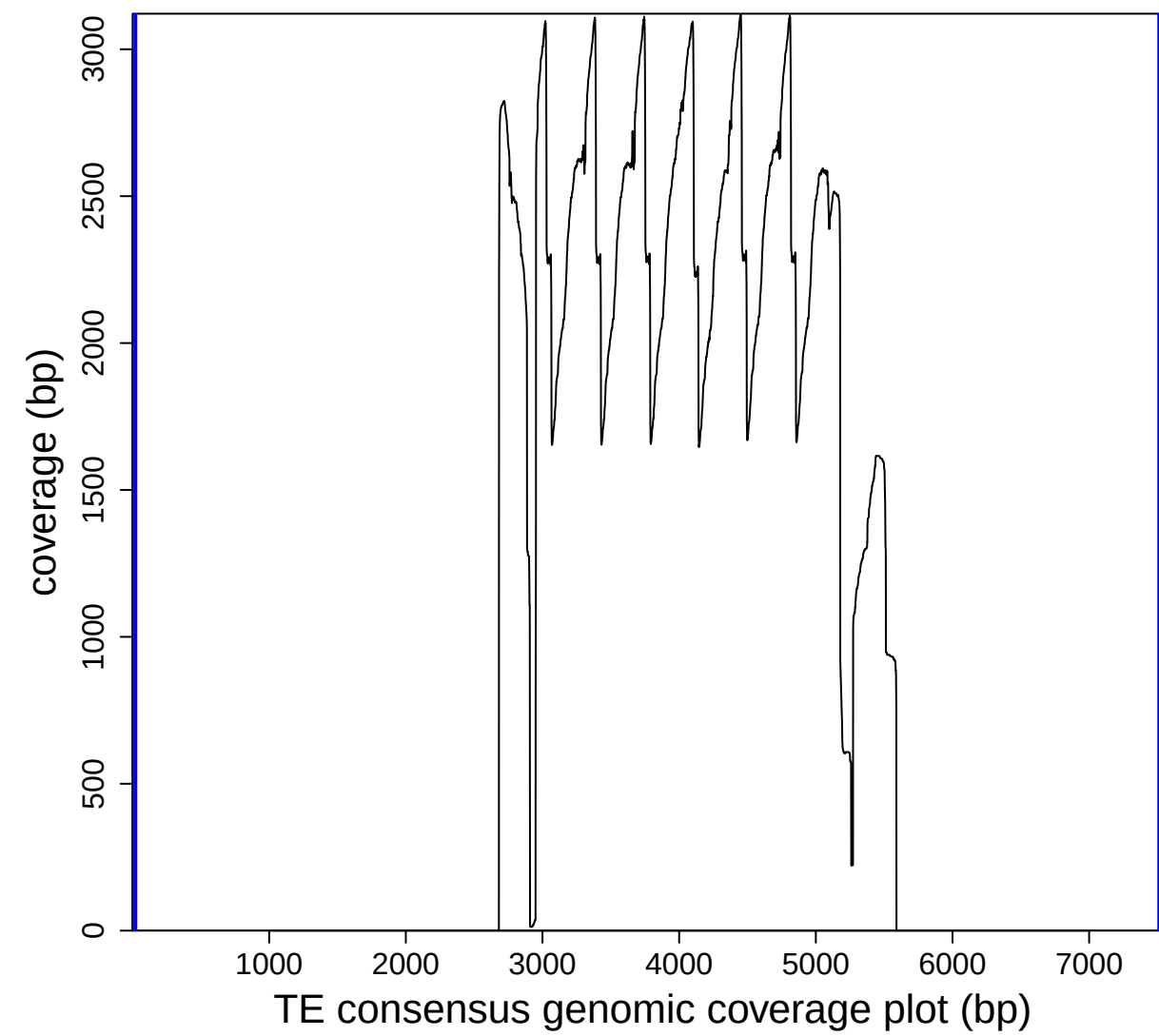
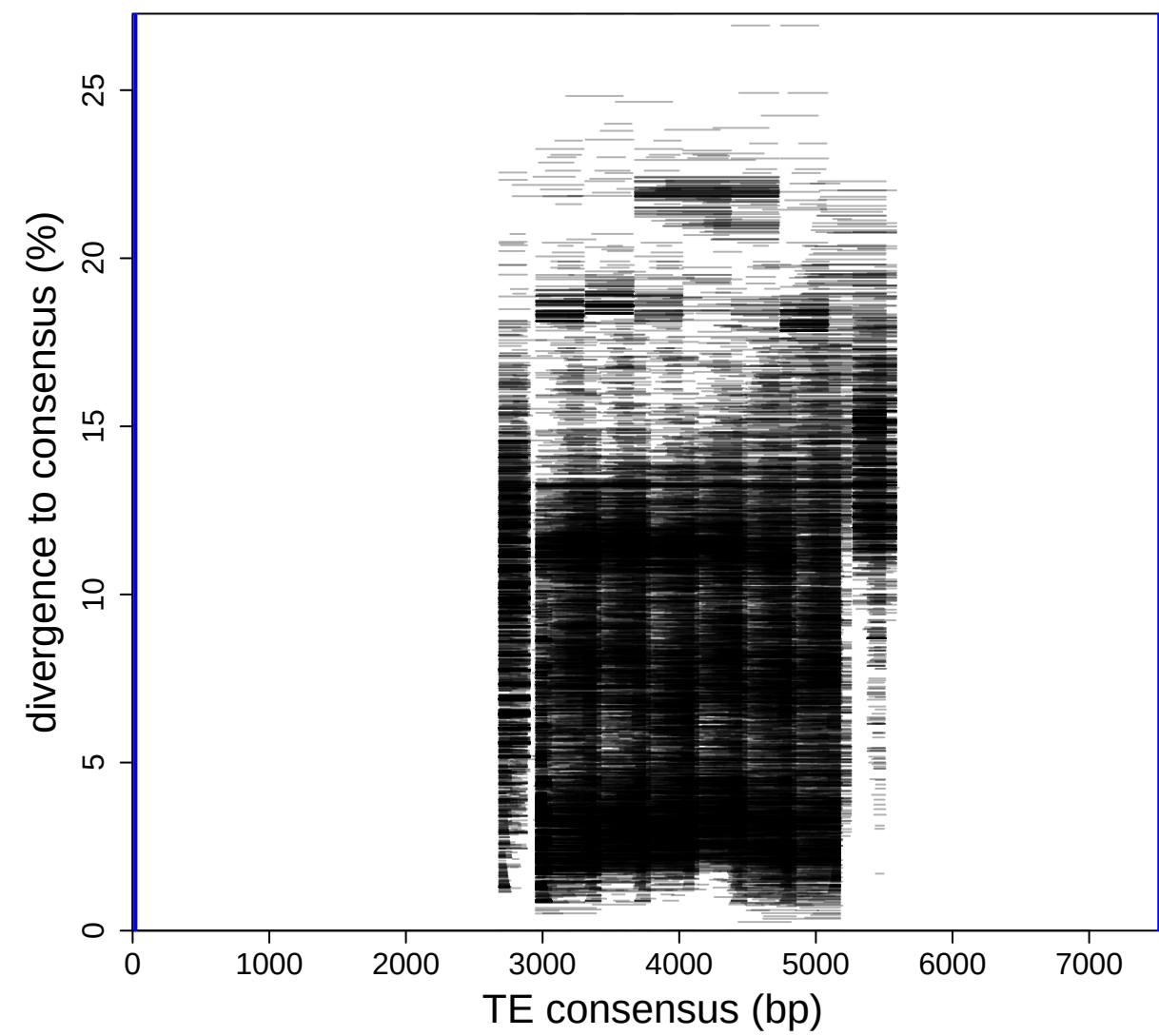
After TEtrimmer 5544 bp



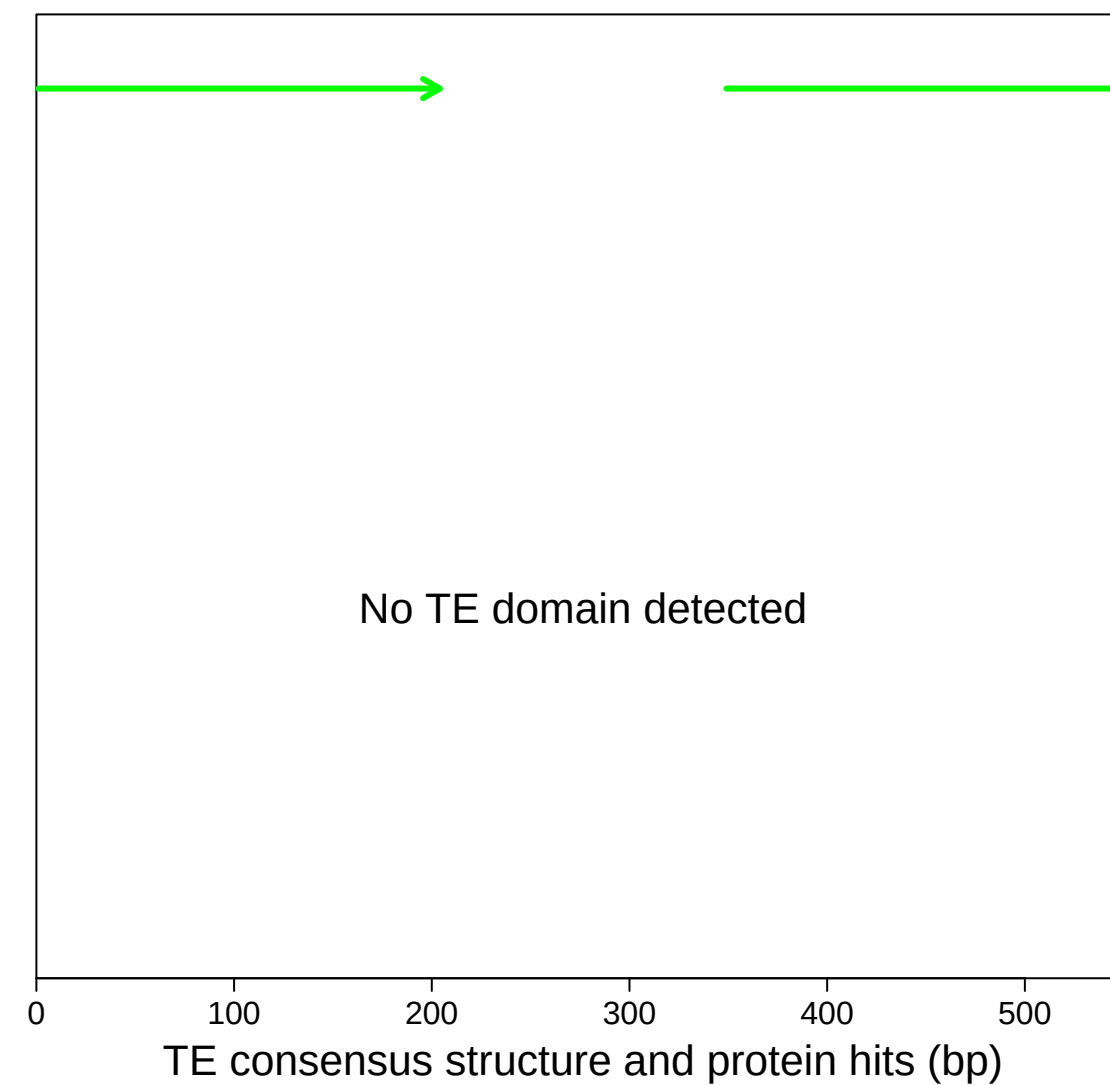
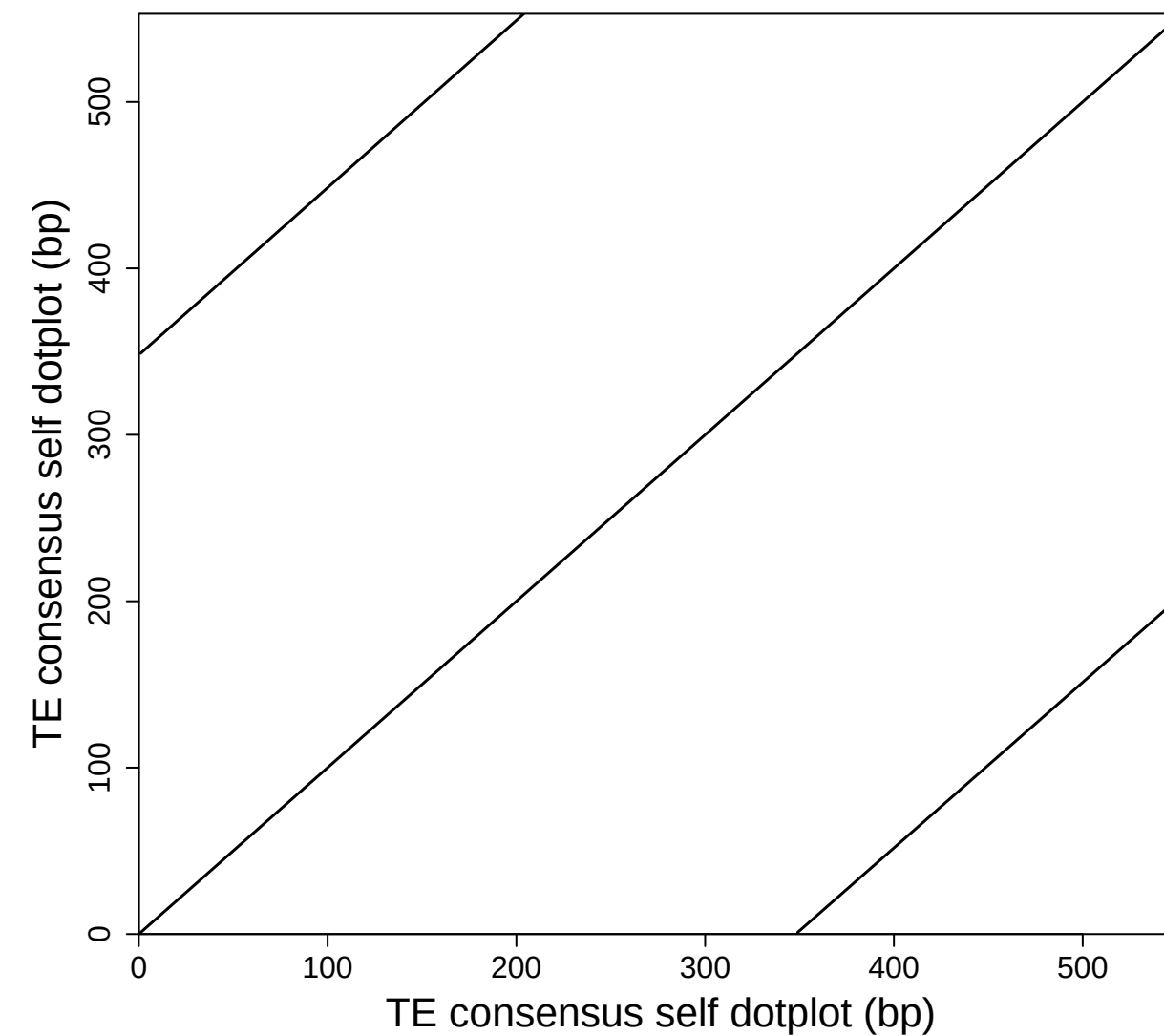
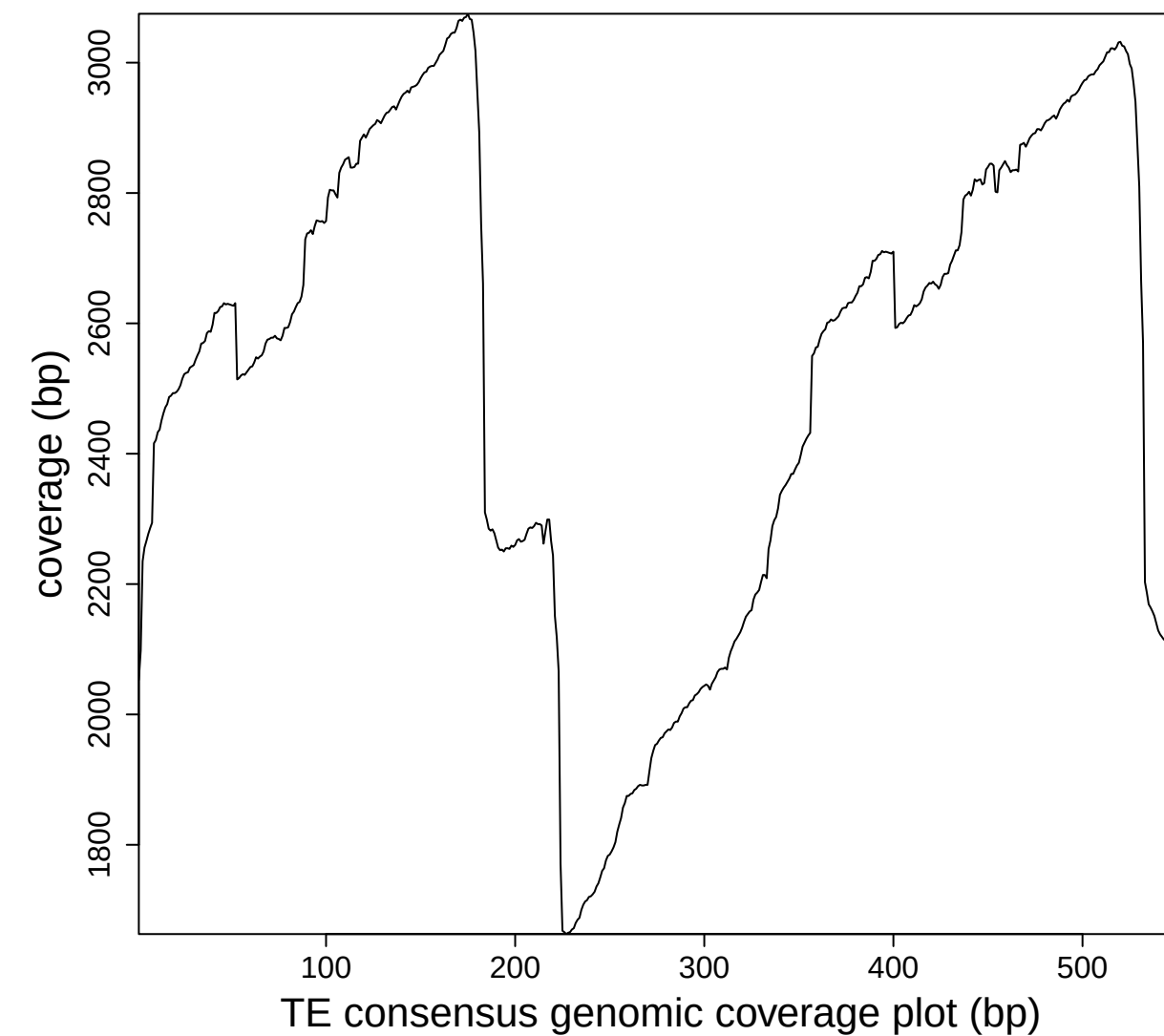
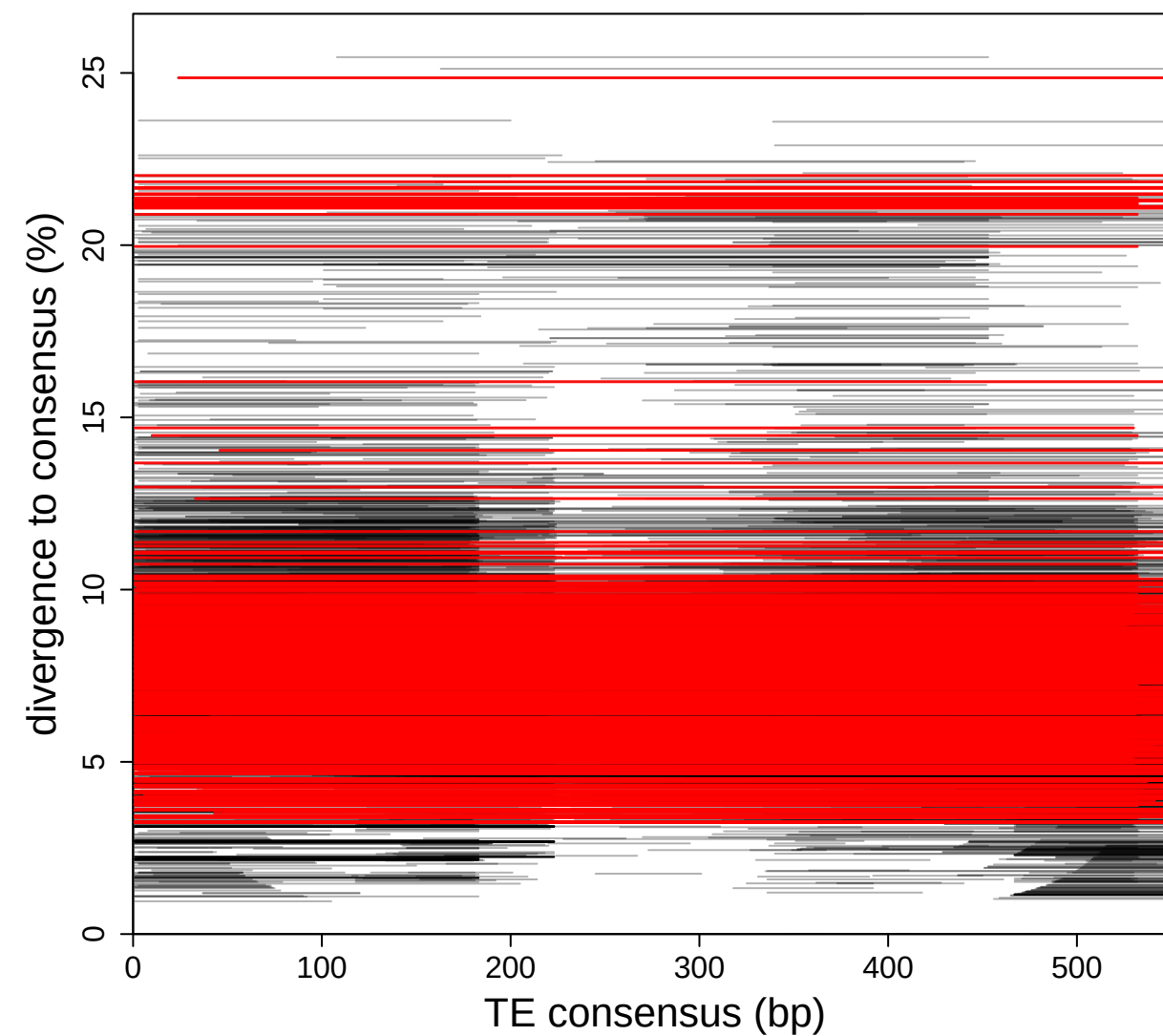
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 7613bp; fragments: 20740; full length: 0 (>=6851.7bp)

After TEtrimmer Extended plot

Blue lines are boundaries

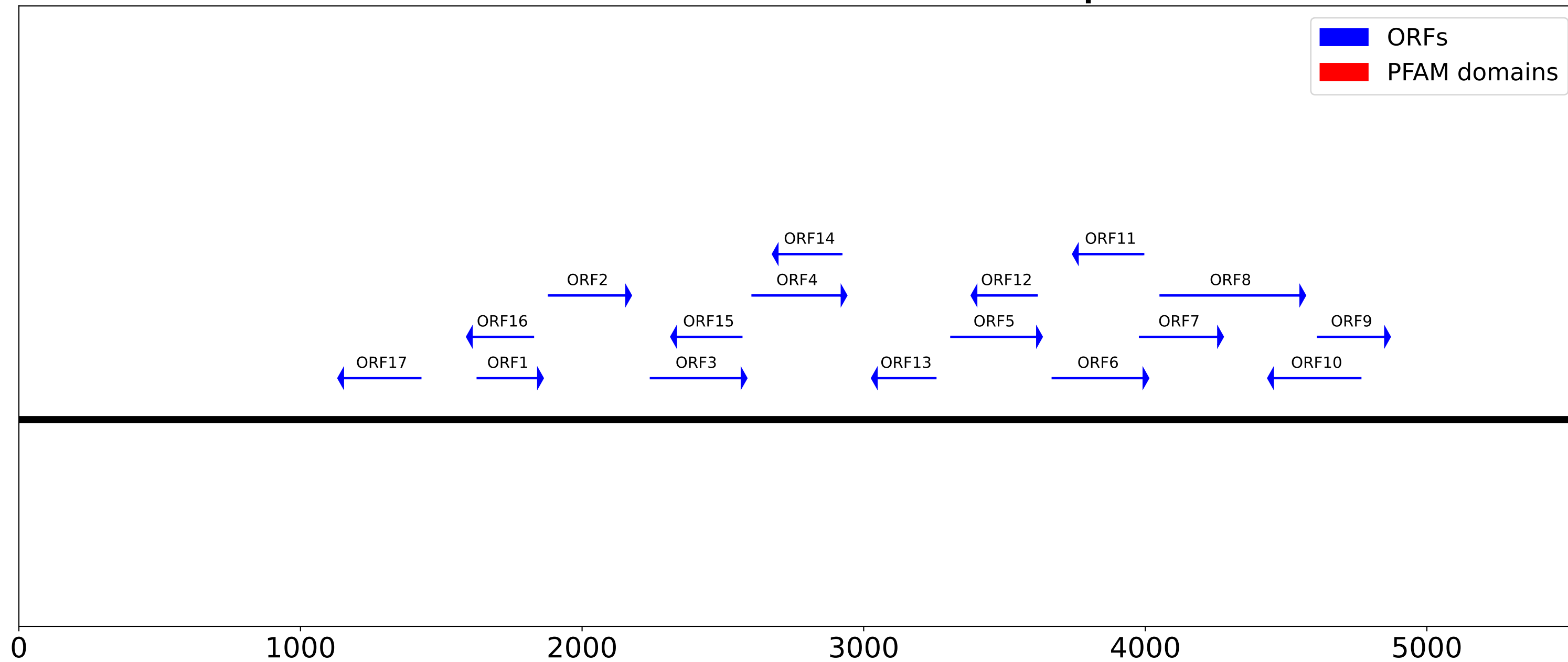


TE: rnd_1_family_747
size: 553bp; fragments: 5657; full length: 841 (≥ 497.7 bp)

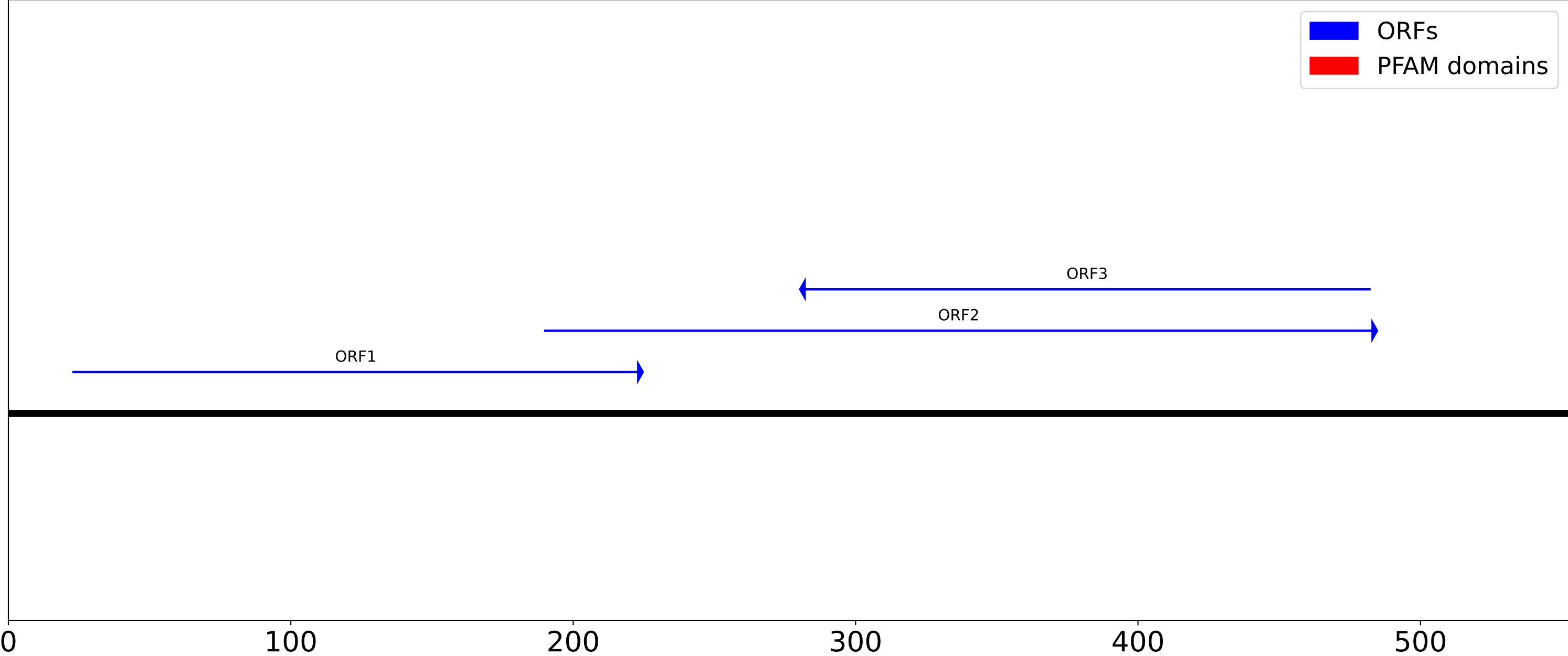


Before TEtrimmer 553 bp

After TEtrimmer ORF and PFAM domain plot



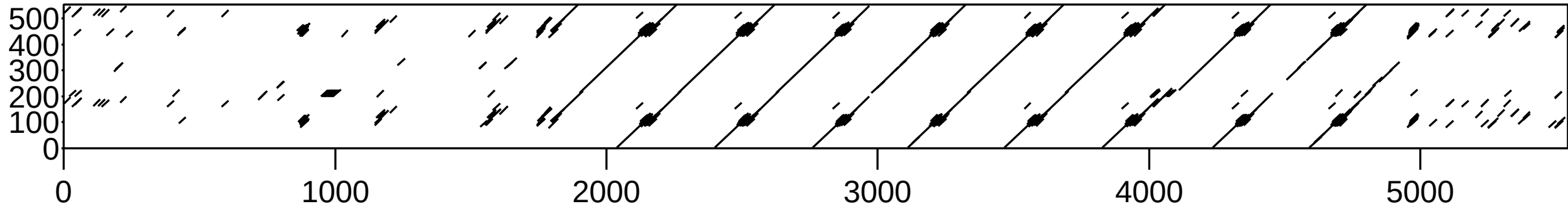
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)