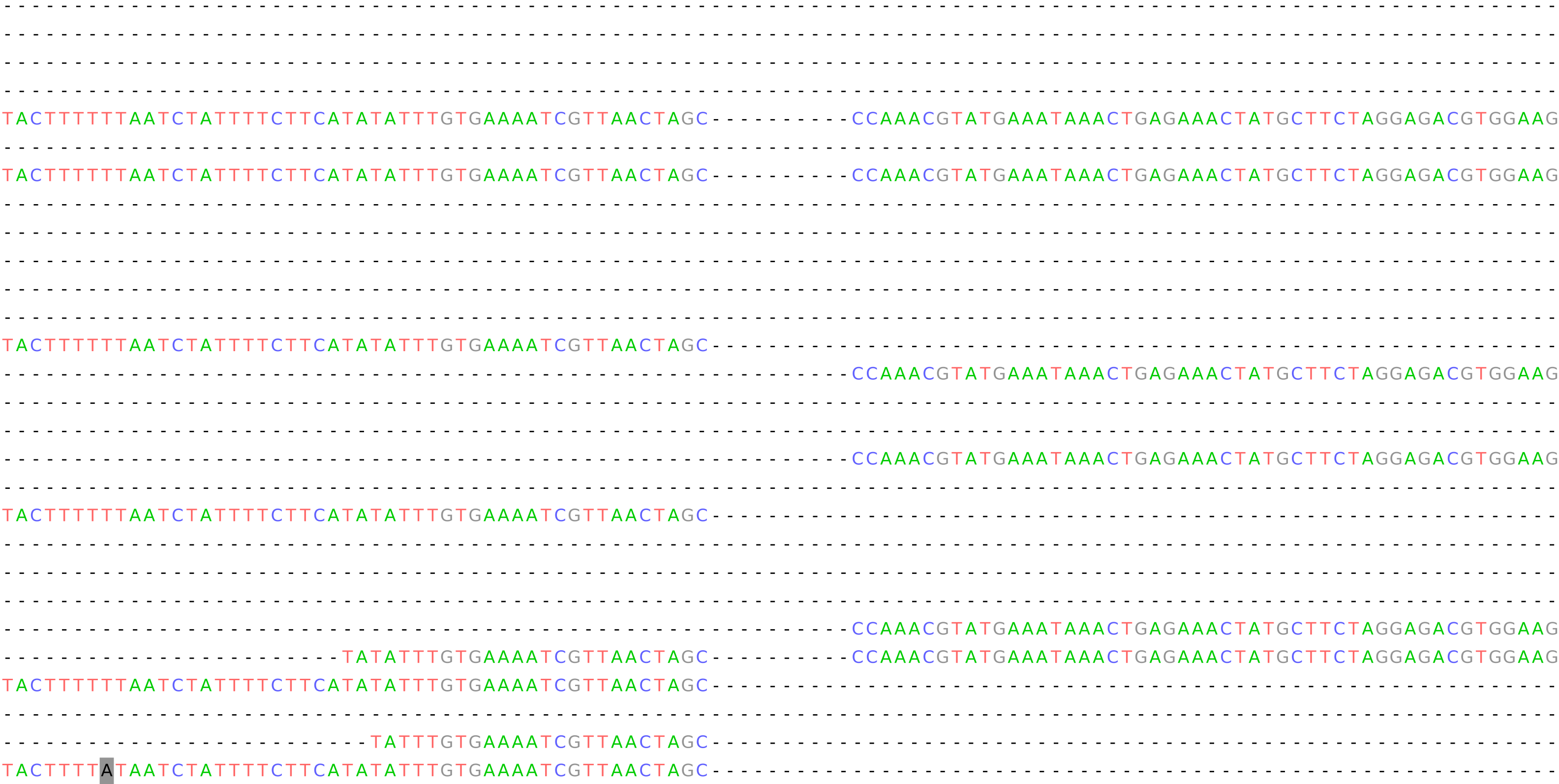
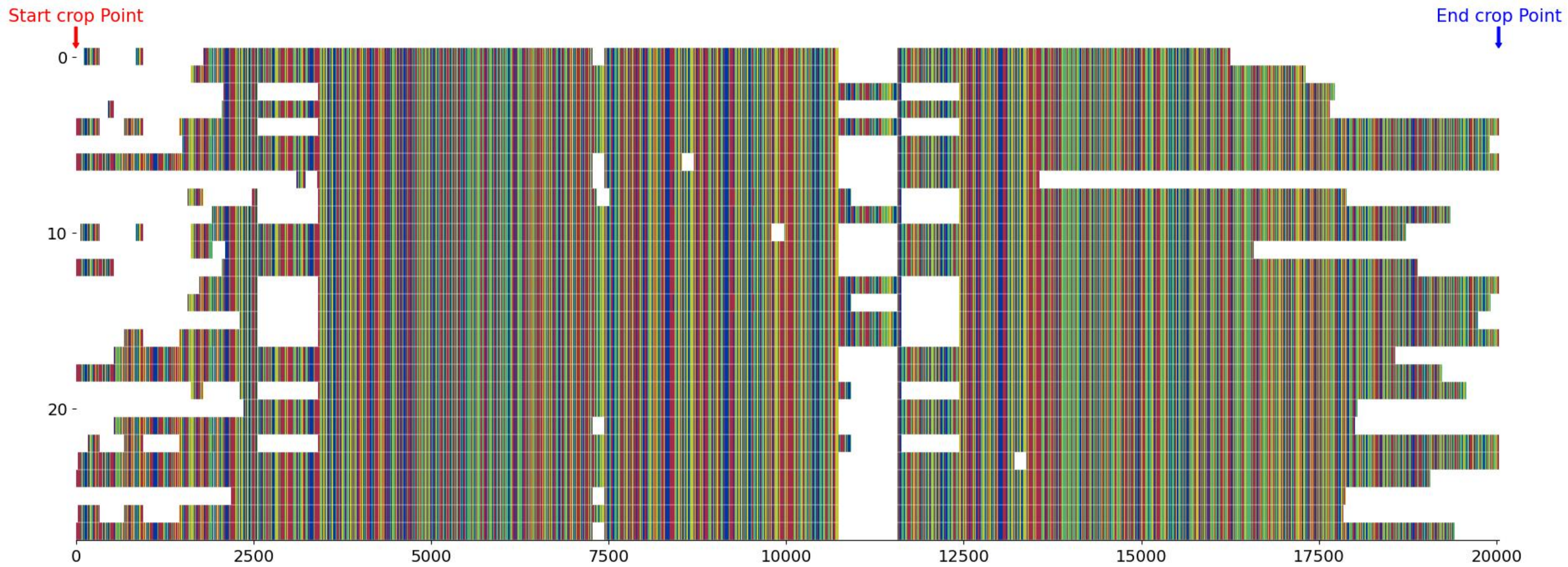


Start crop Point

End crop Point

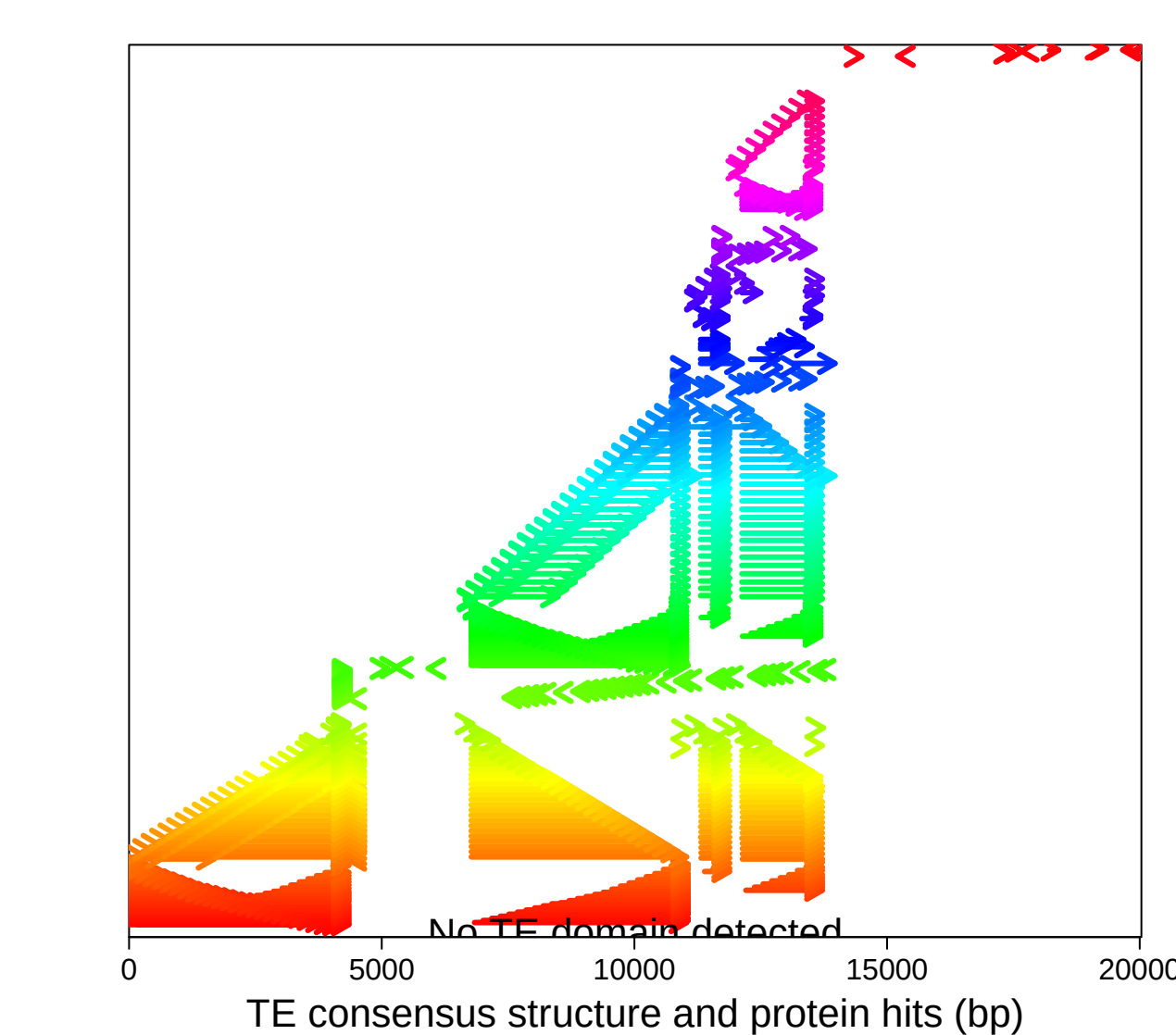
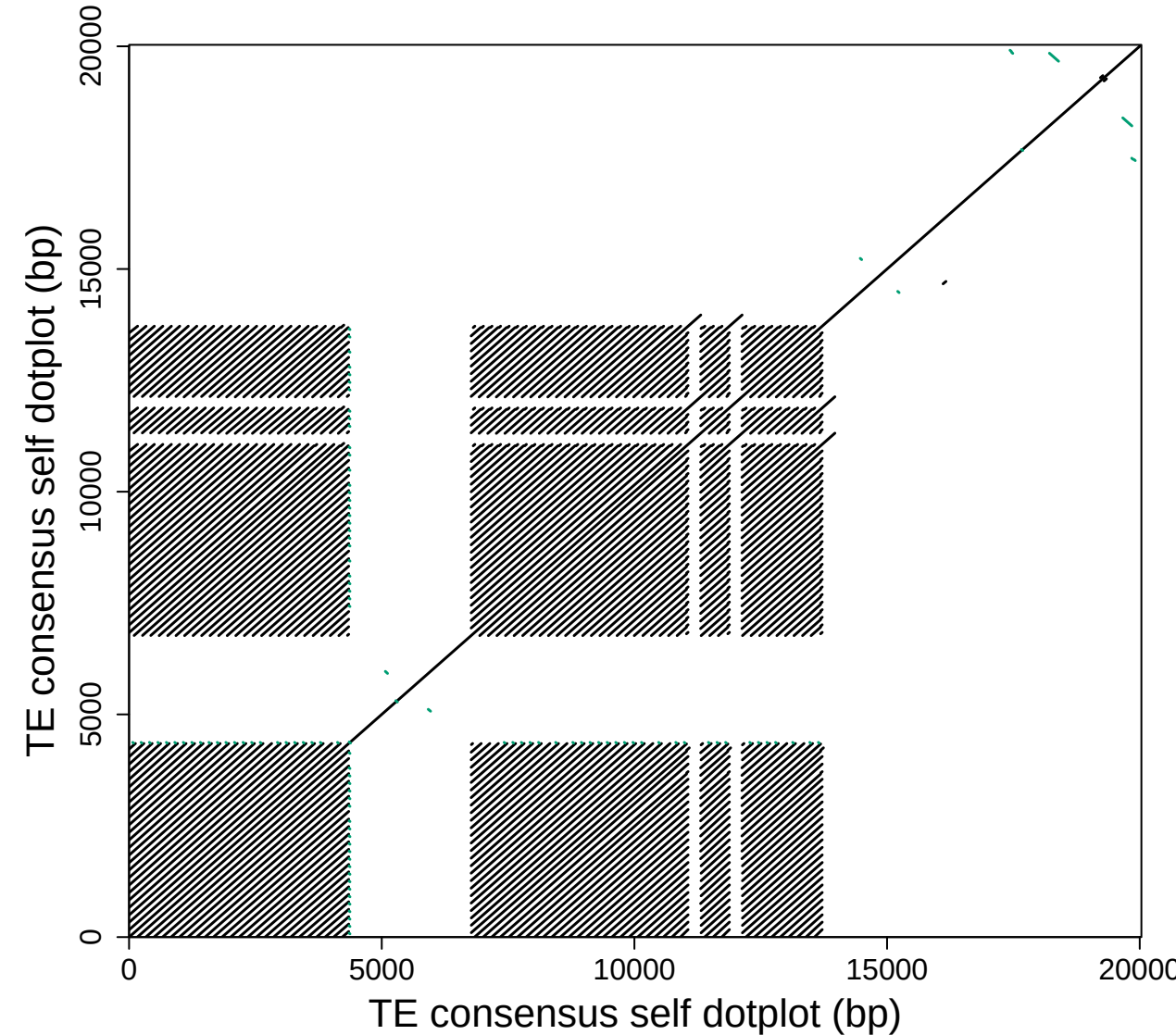
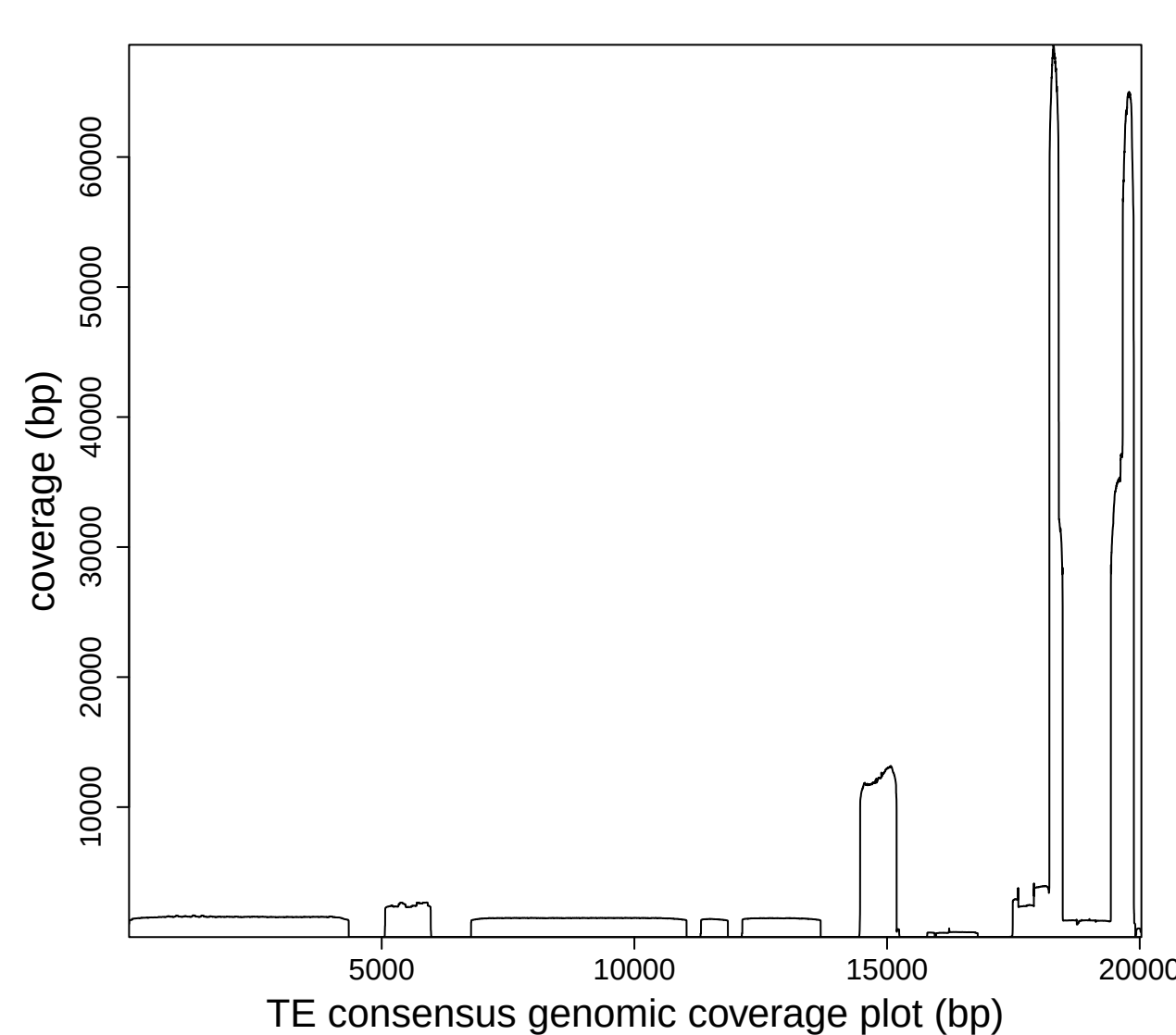
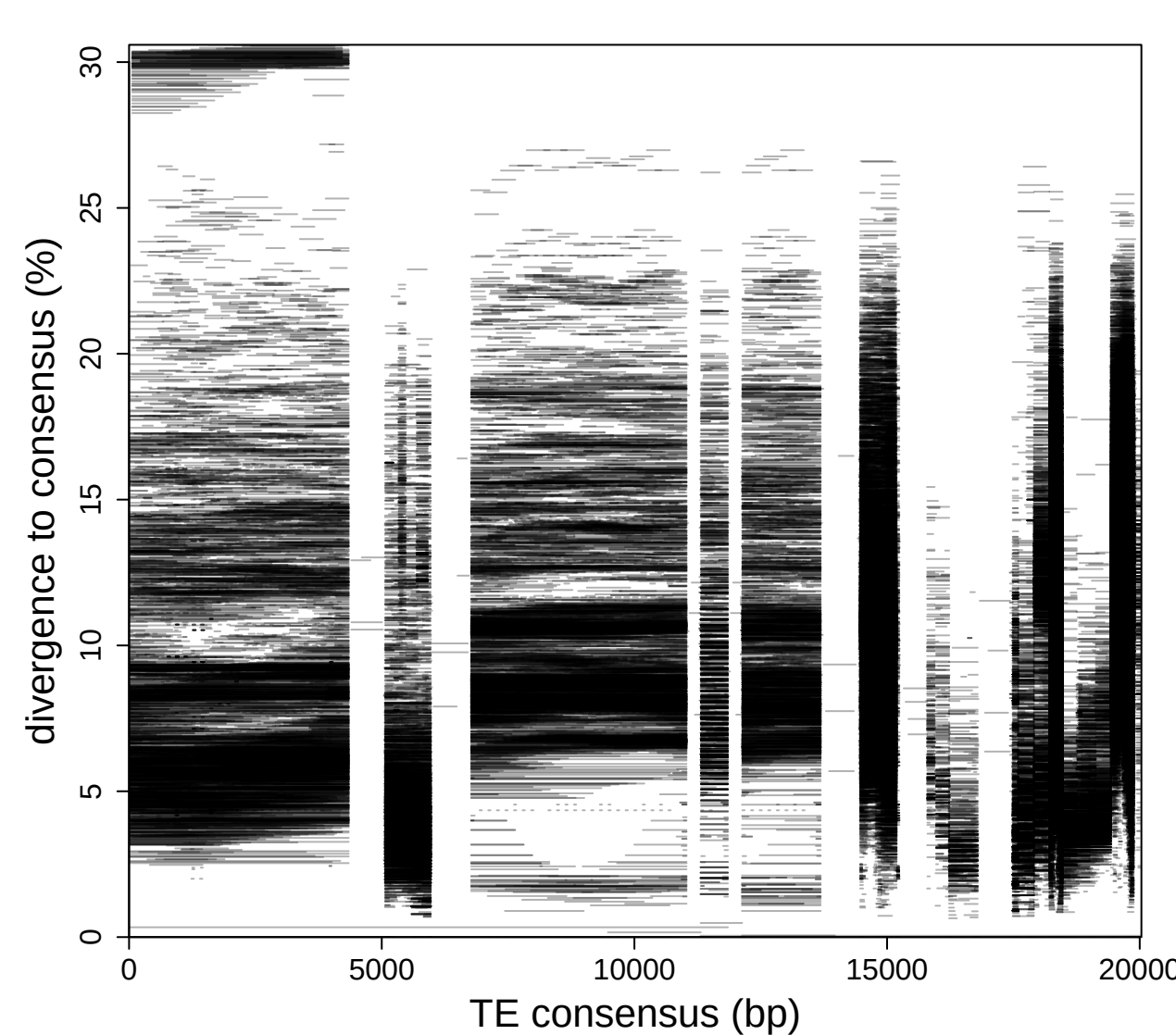
MSA length = 20033





g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 20033bp; fragments: 203489; full length: 0 (>=18029.7bp)

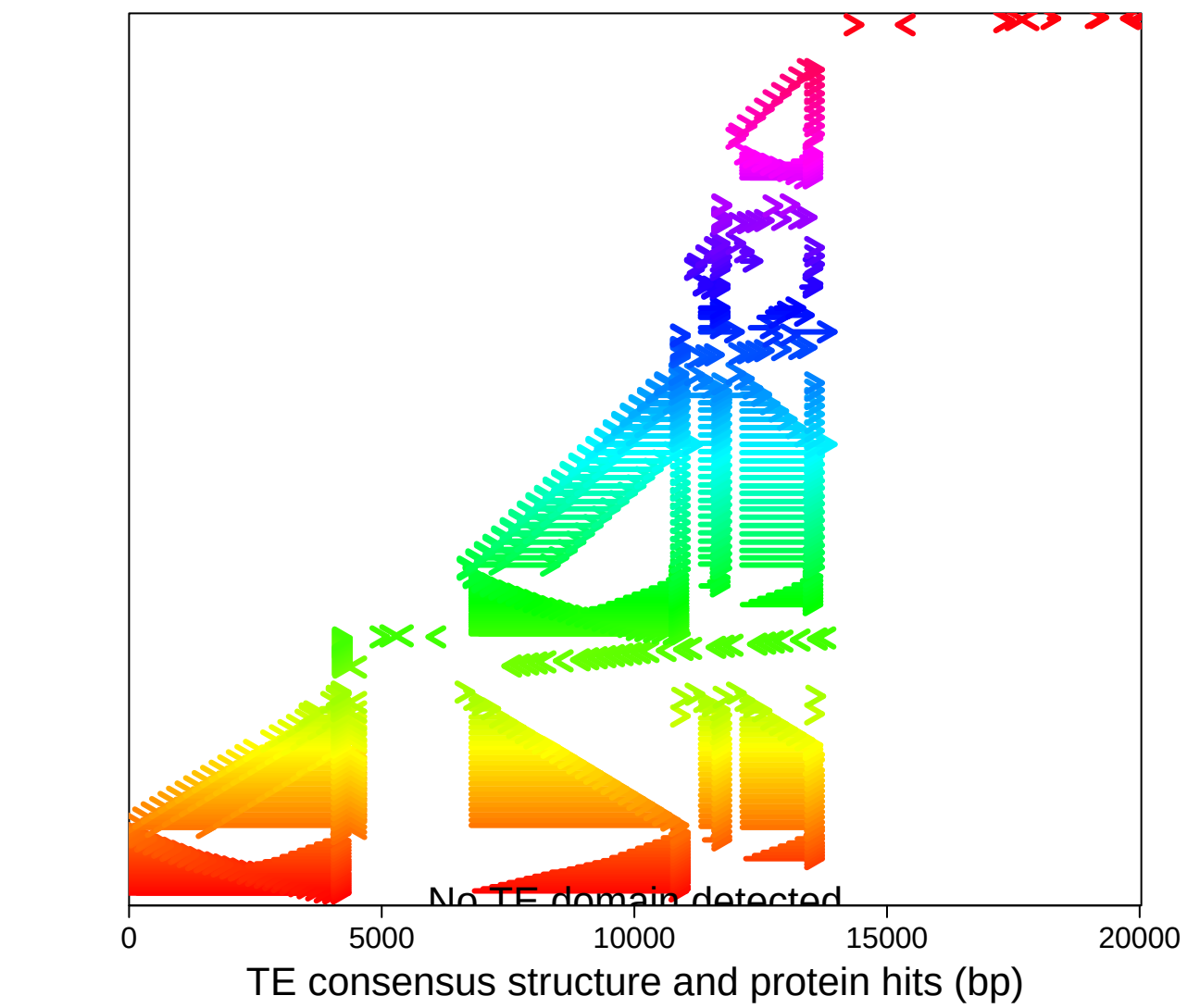
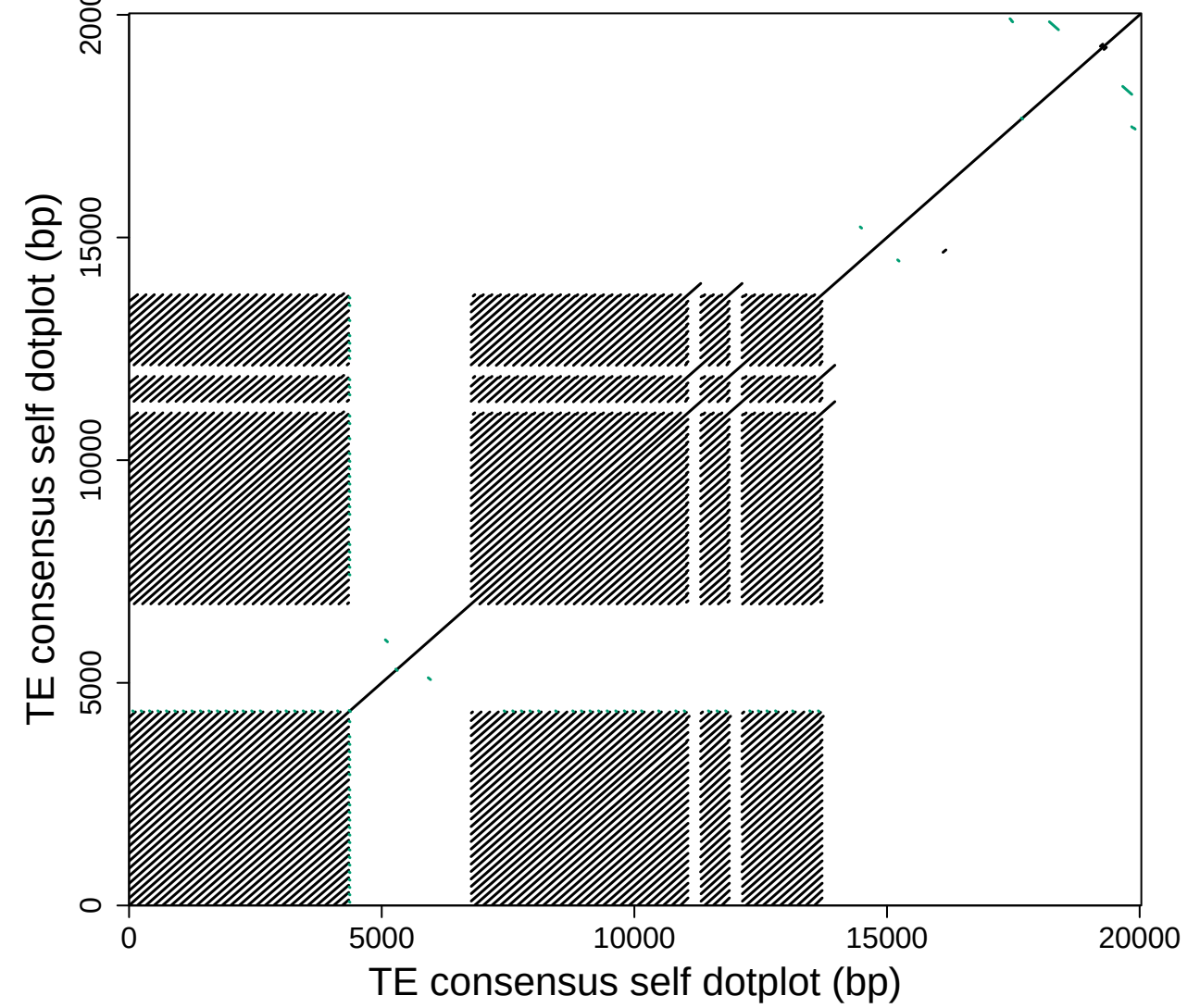
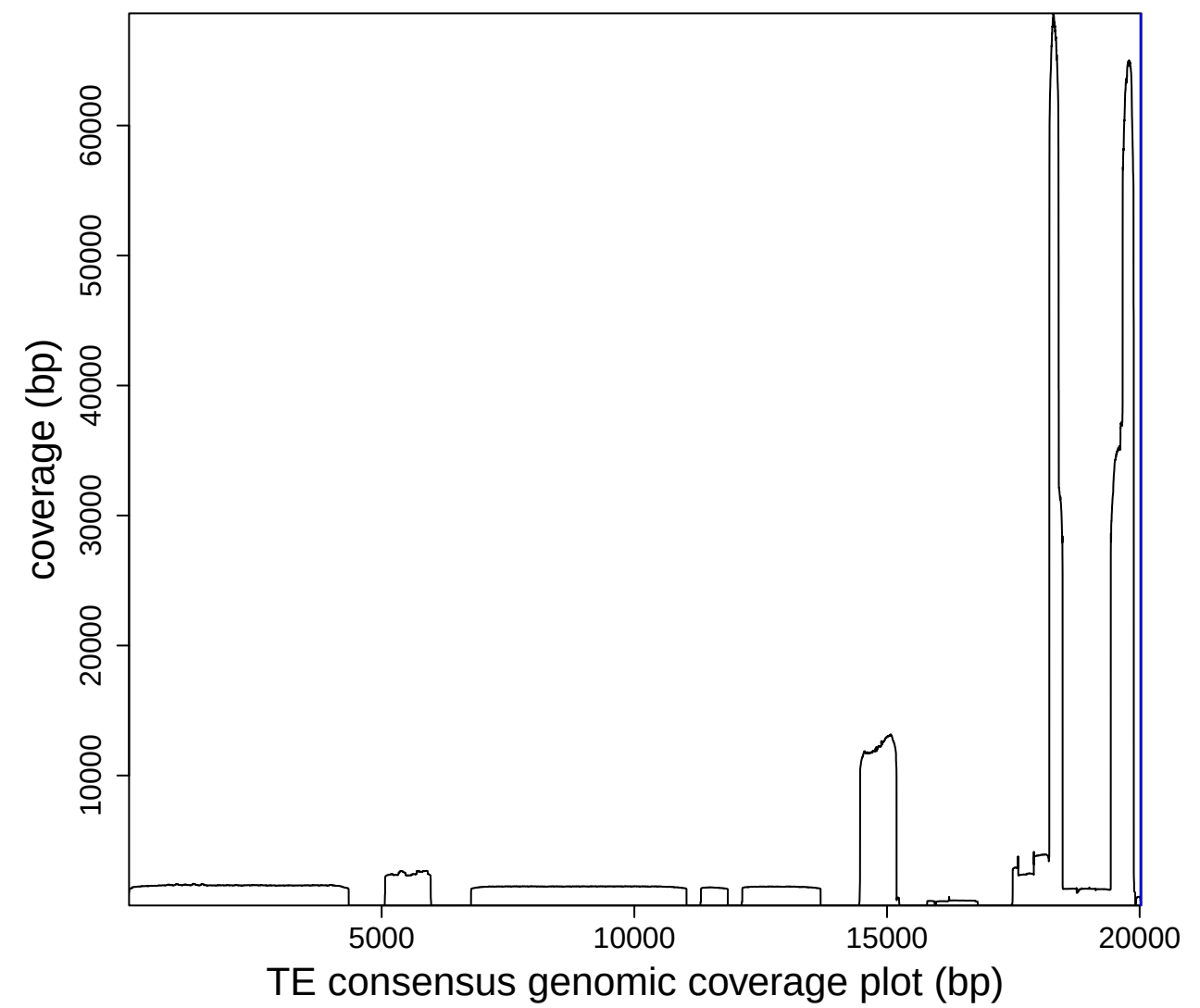
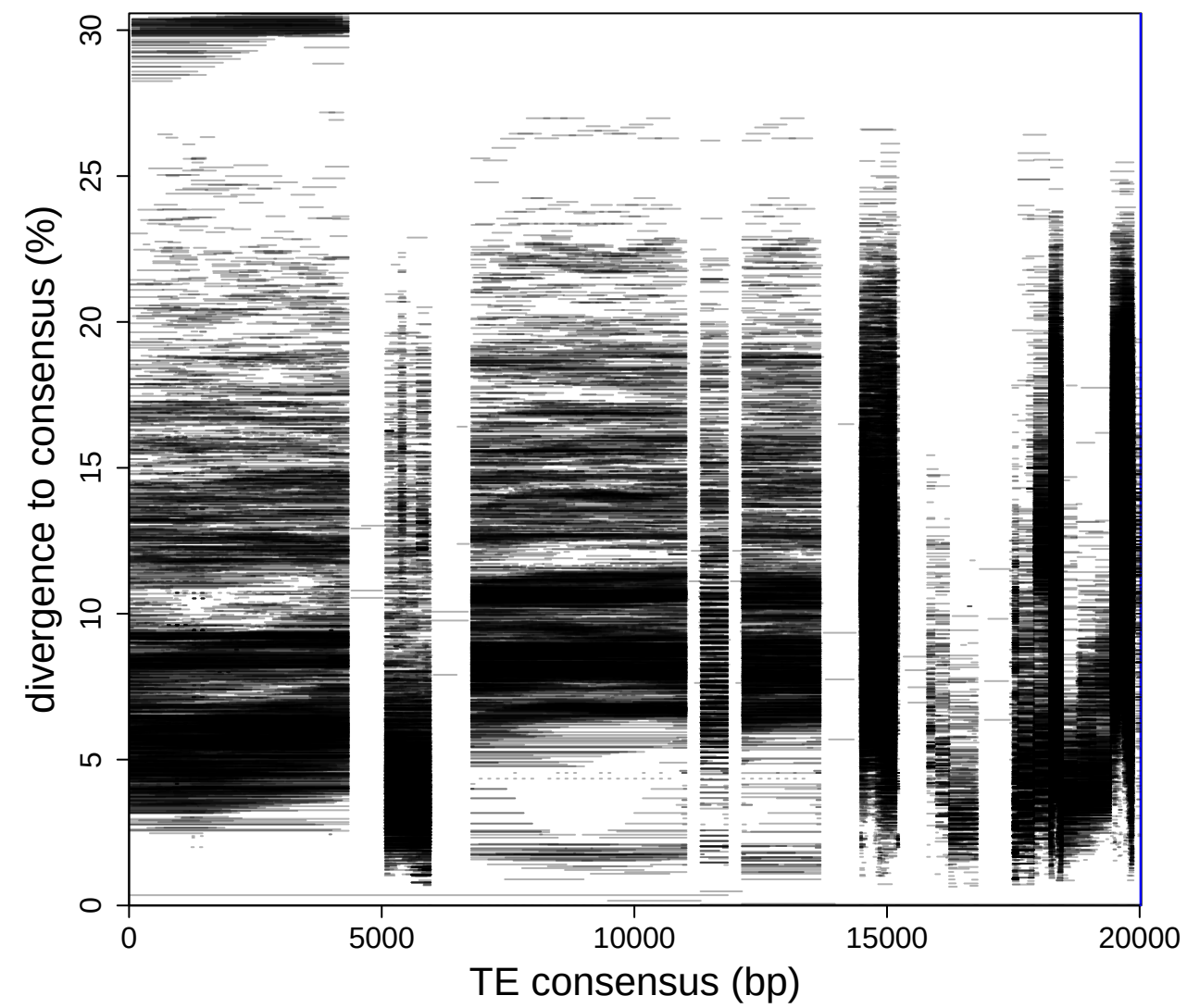
After TEtrimmer 20033 bp



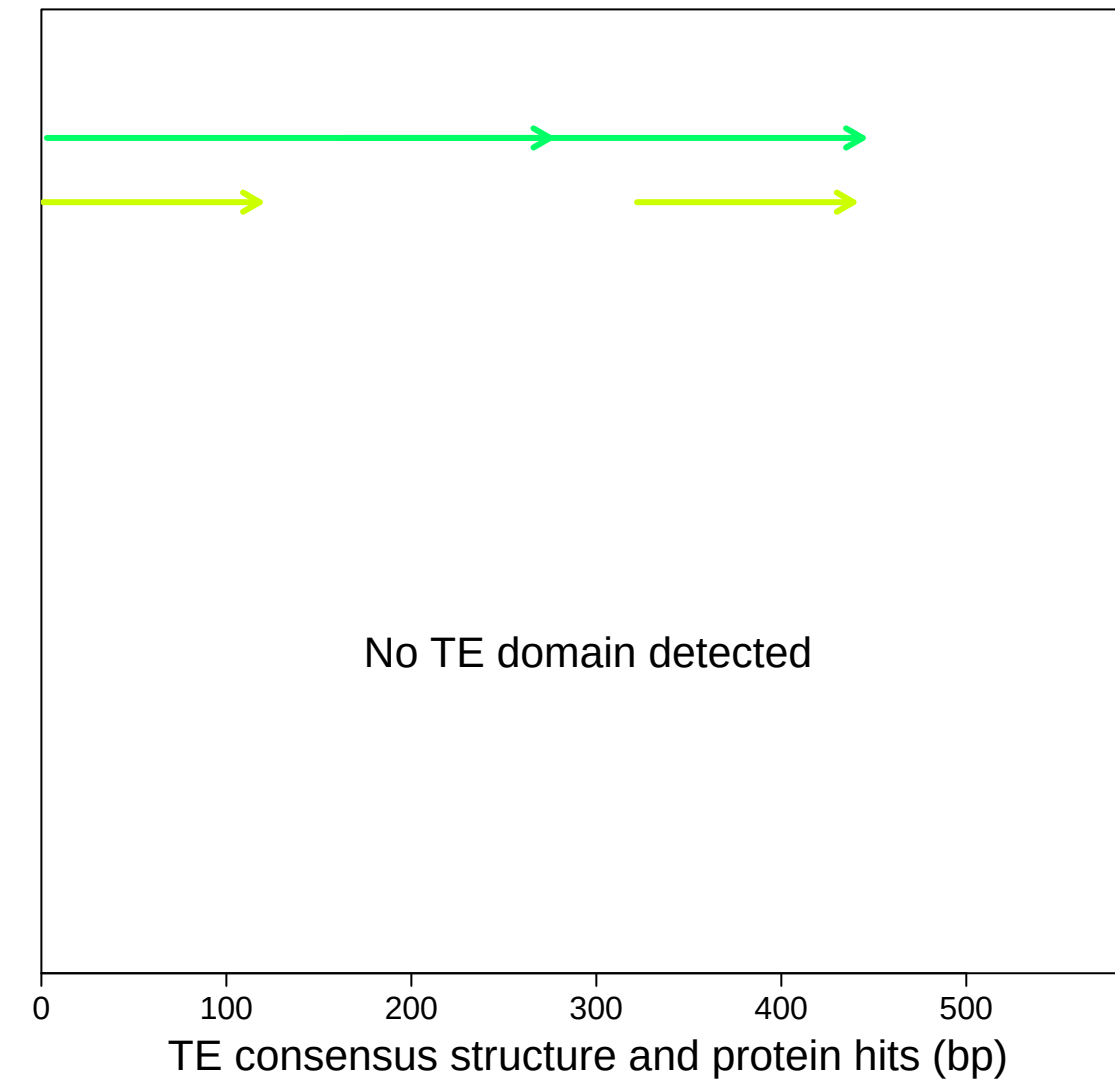
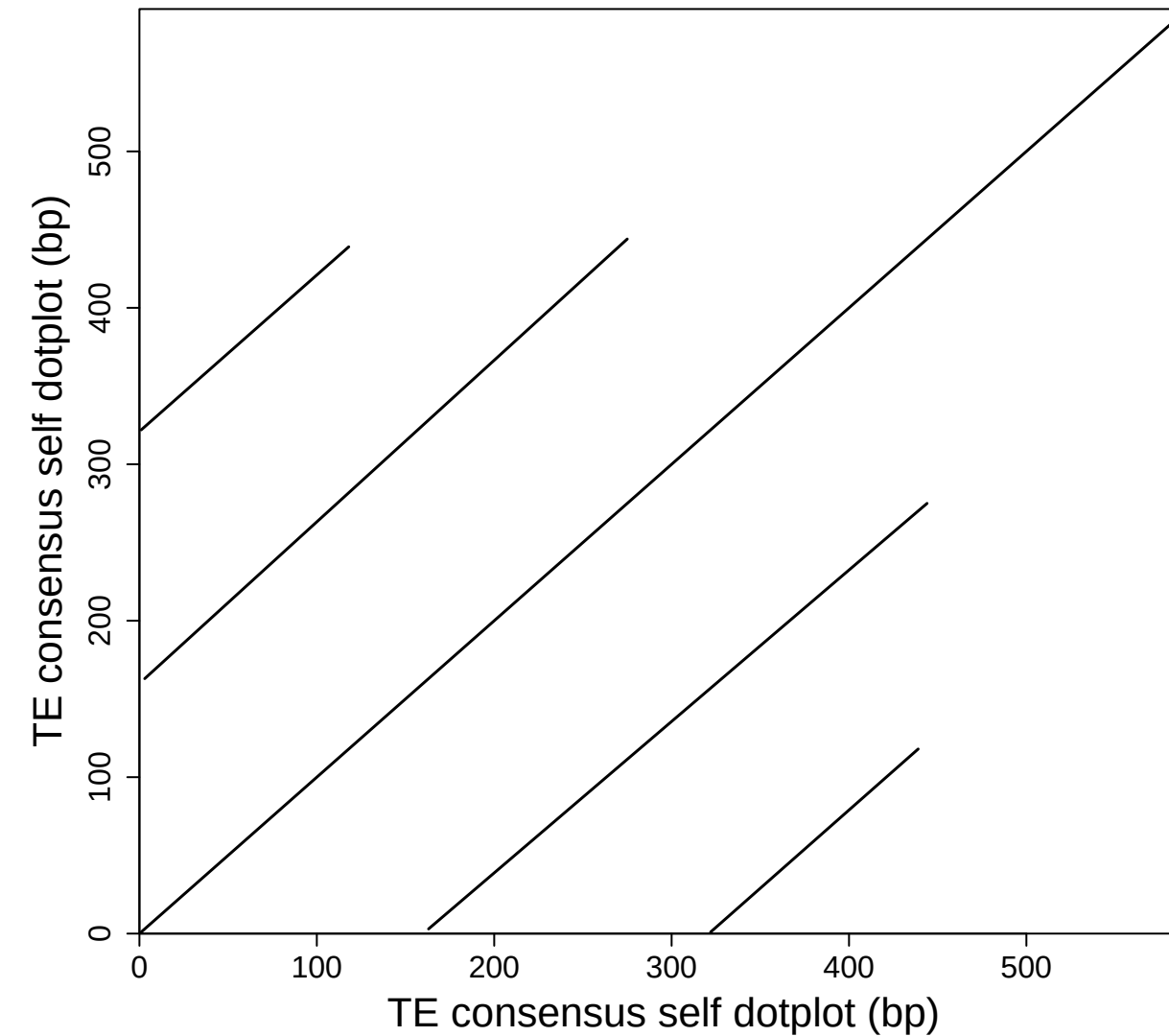
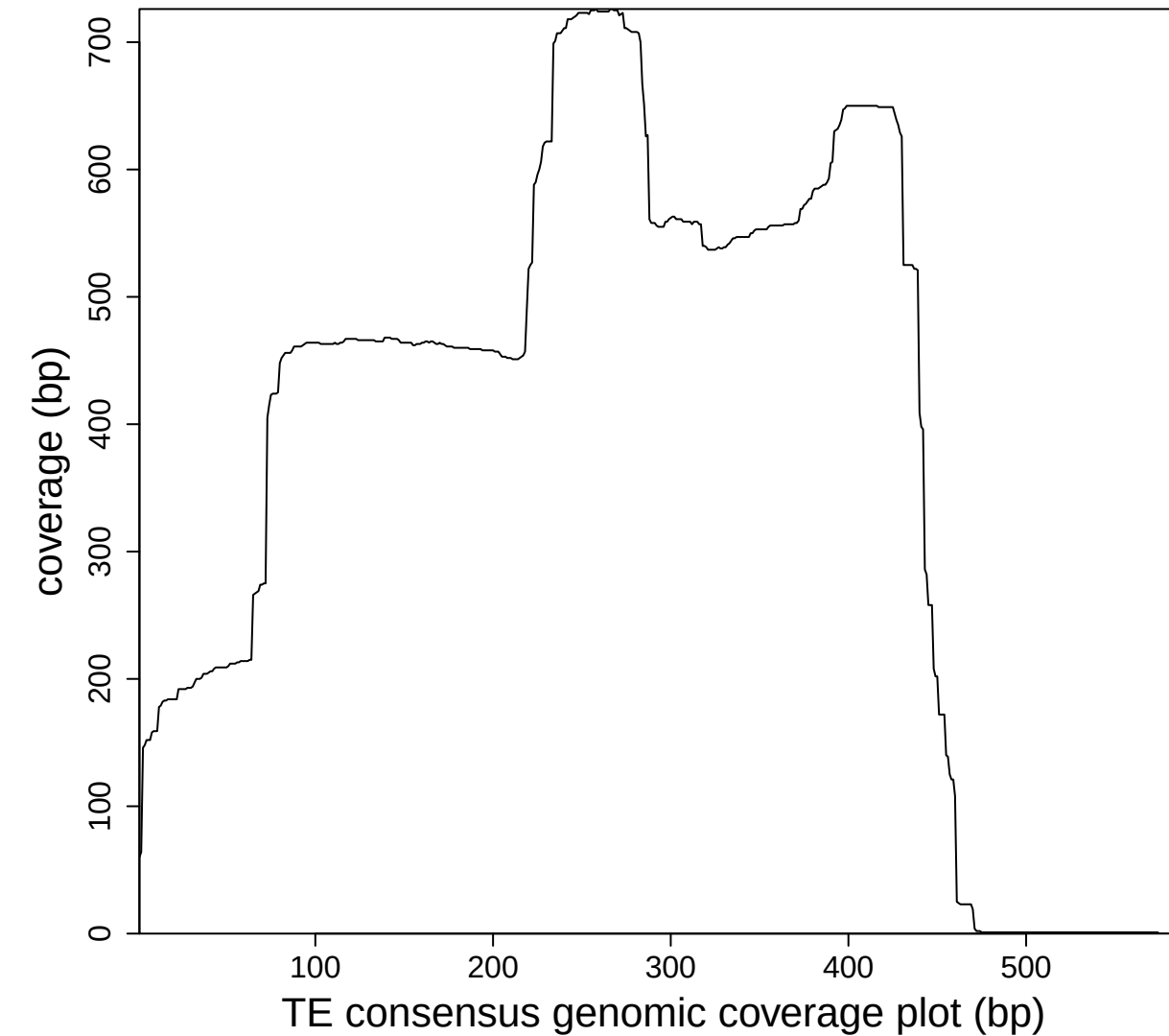
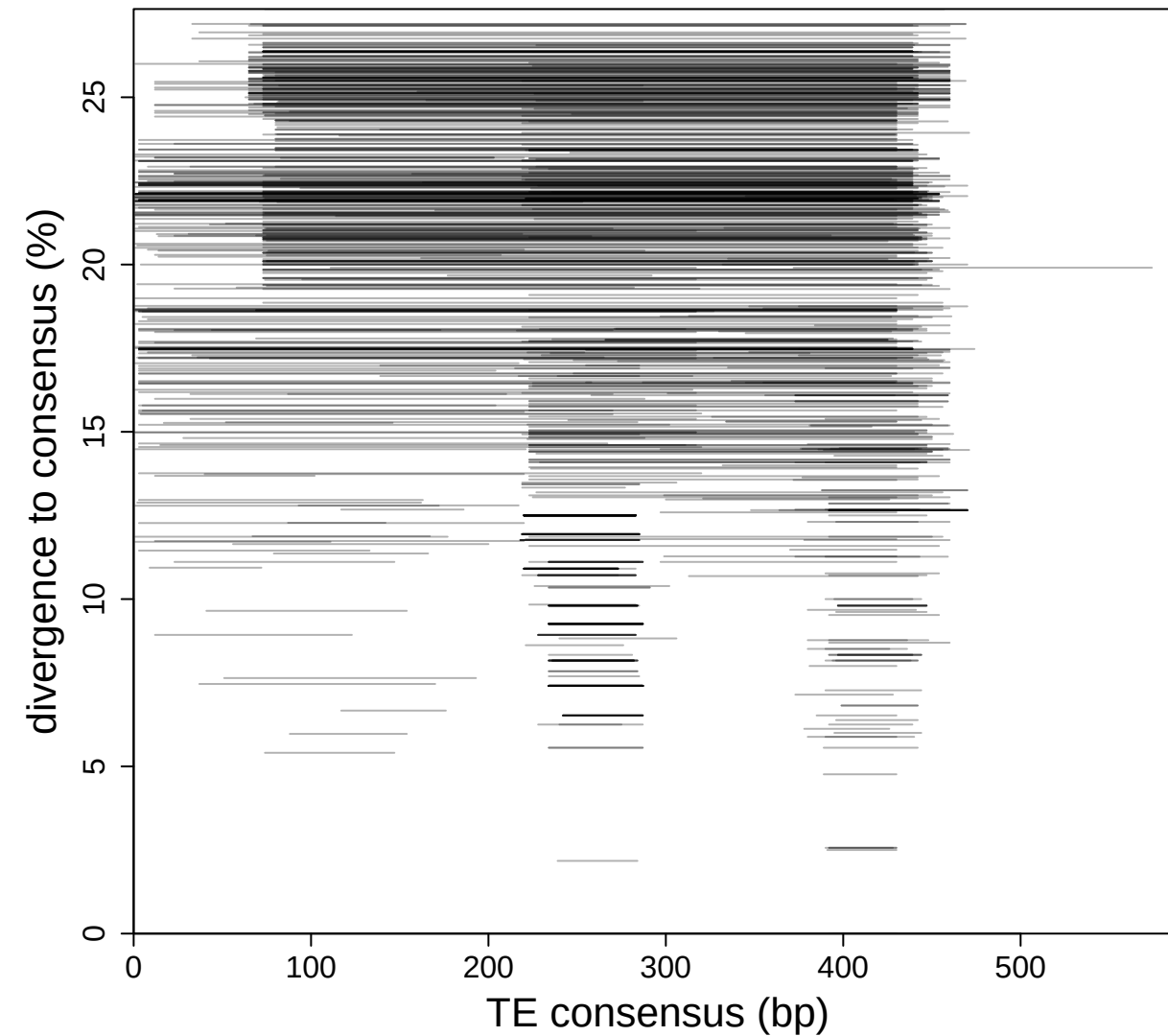
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 20033bp; fragments: 203447; full length: 0 (>=18029.7bp)

After TEtrimmer Extended plot

Blue lines are boundaries

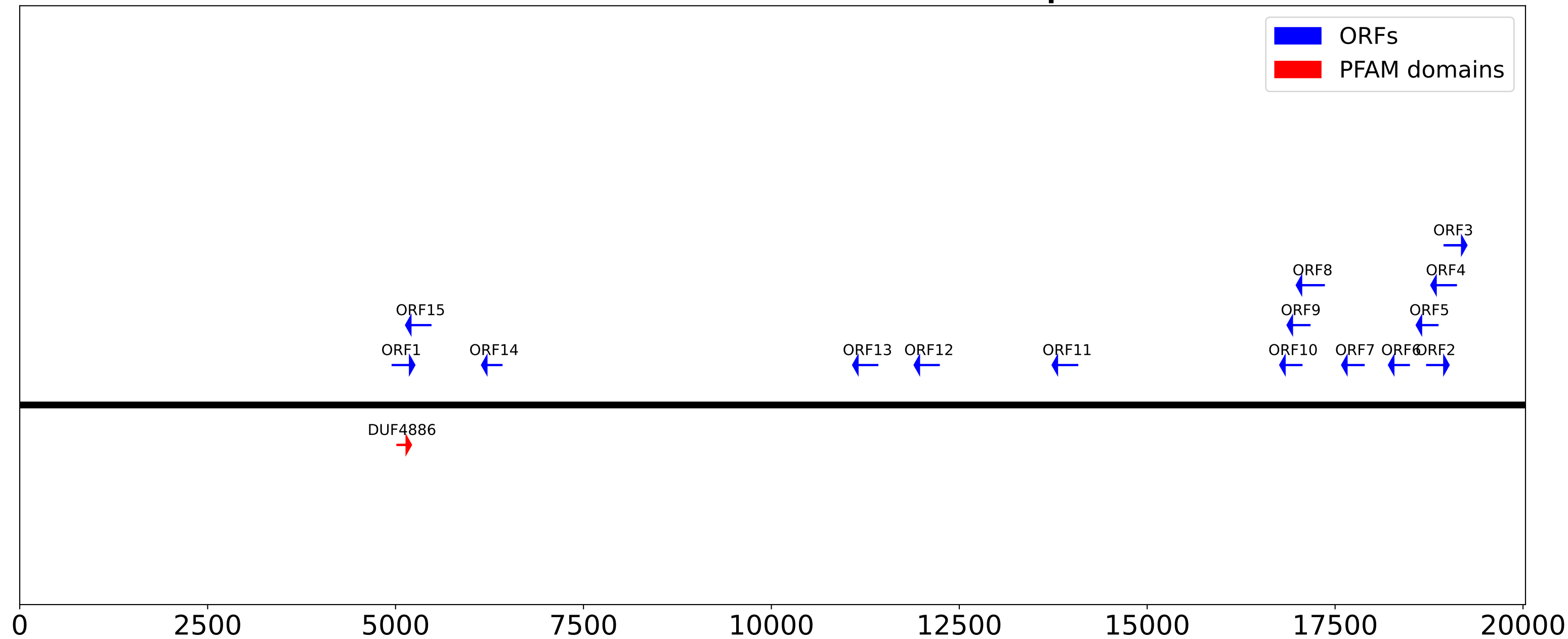


TE: rnd_4_family_157
size: 591bp; fragments: 901; full length: 0 (≥ 531.9 bp)

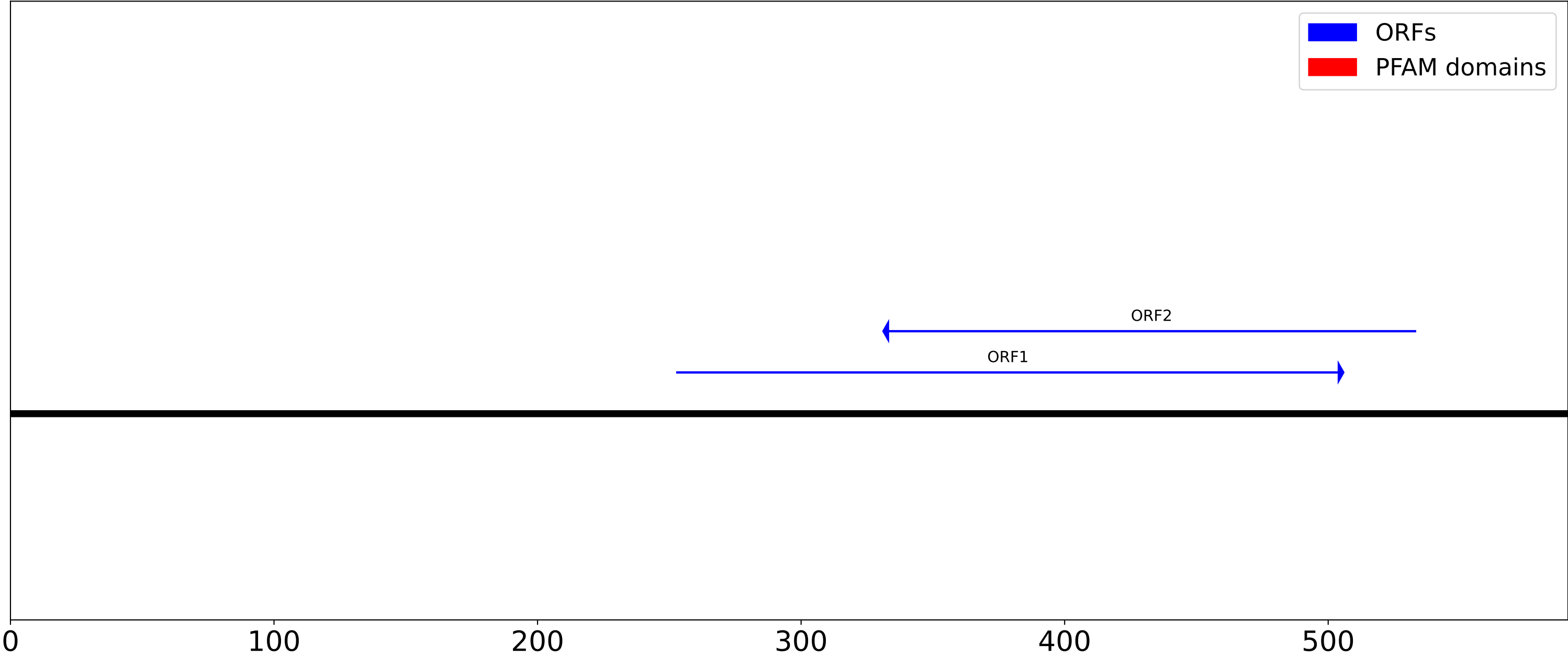


Before TEtrimmer 591 bp

After TEtrimmer ORF and PFAM domain plot



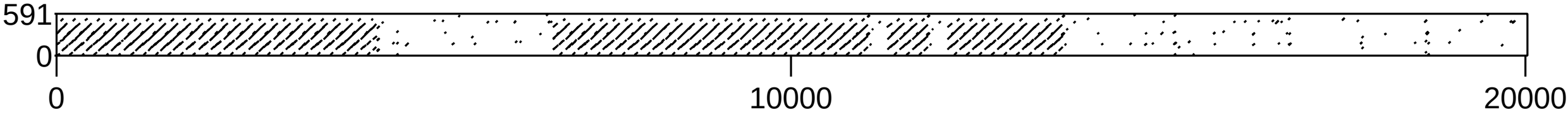
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)