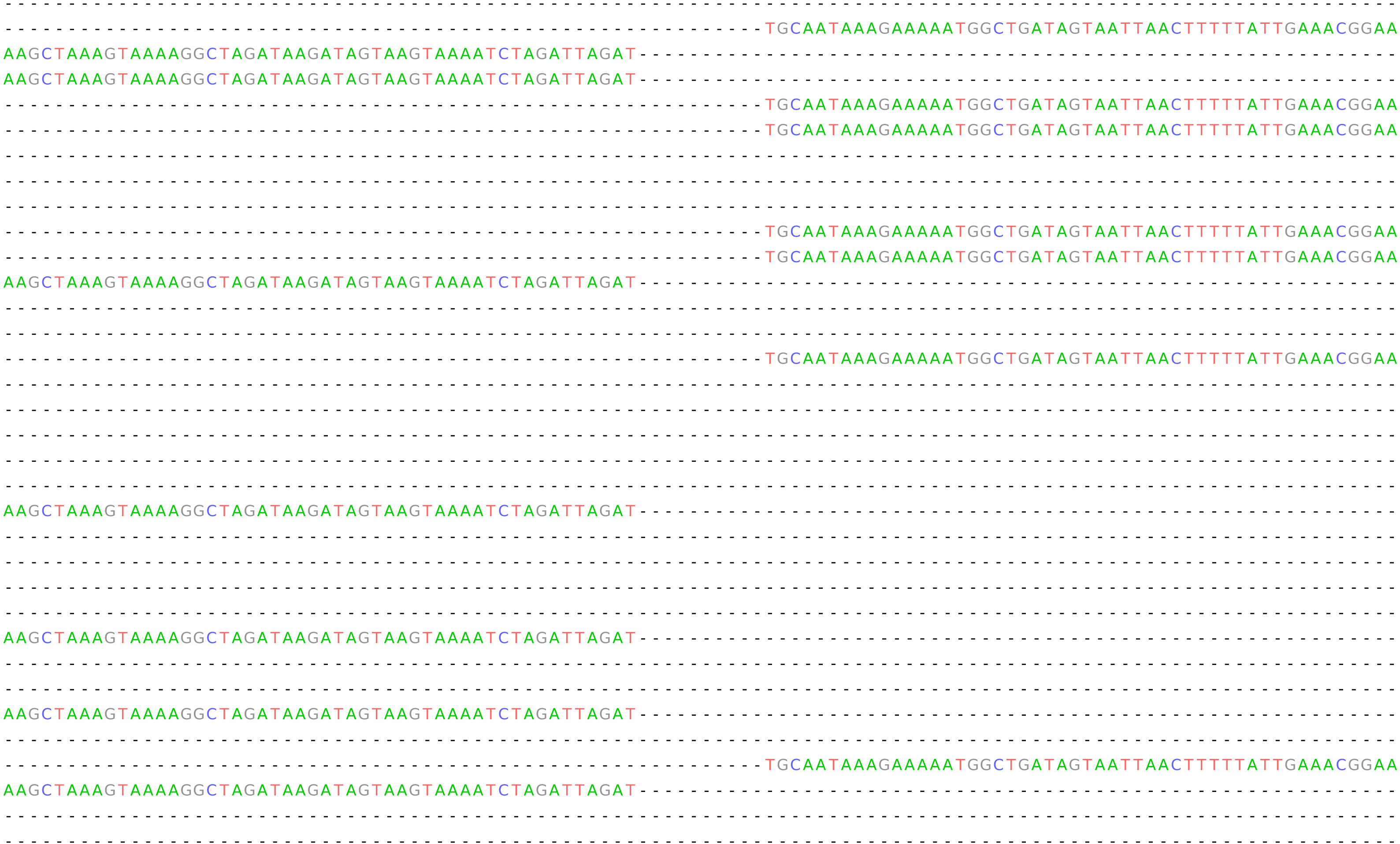
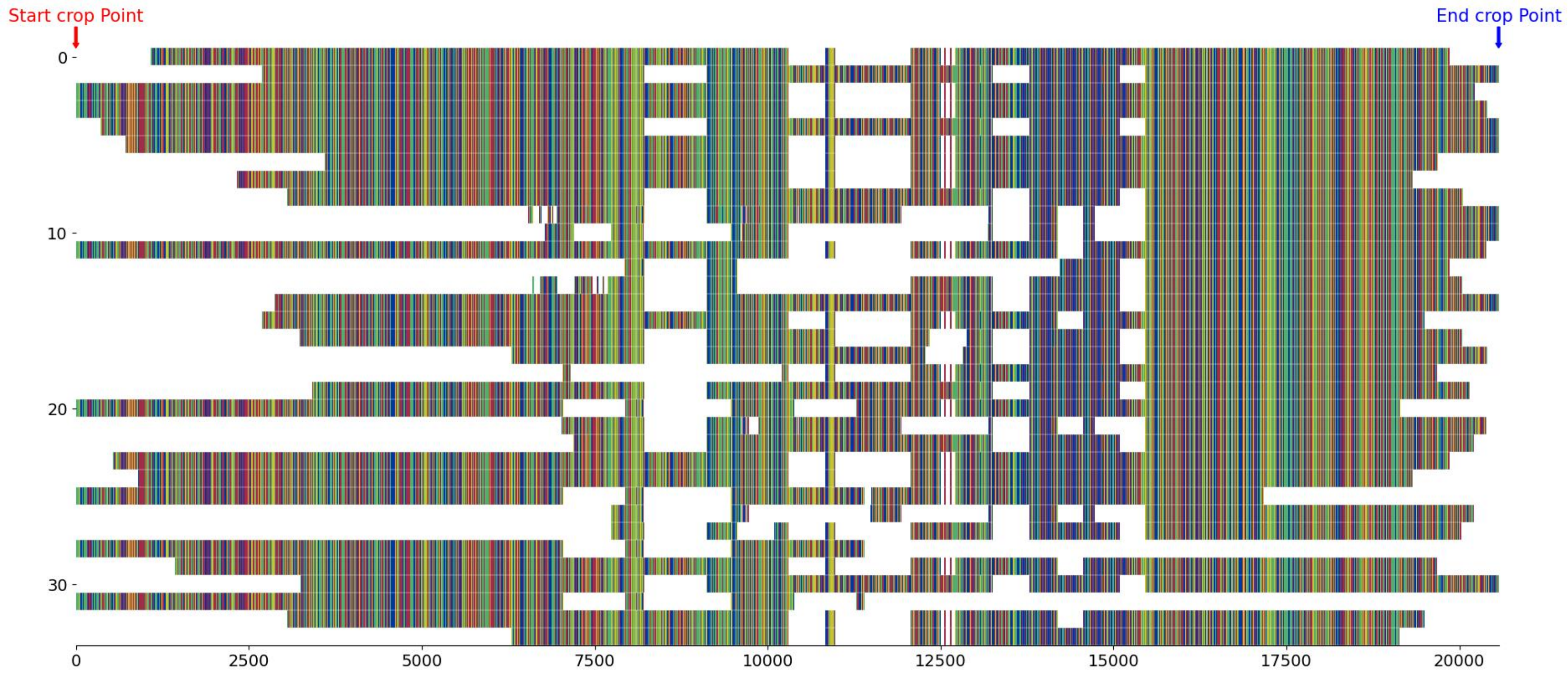


Start crop Point

End crop Point

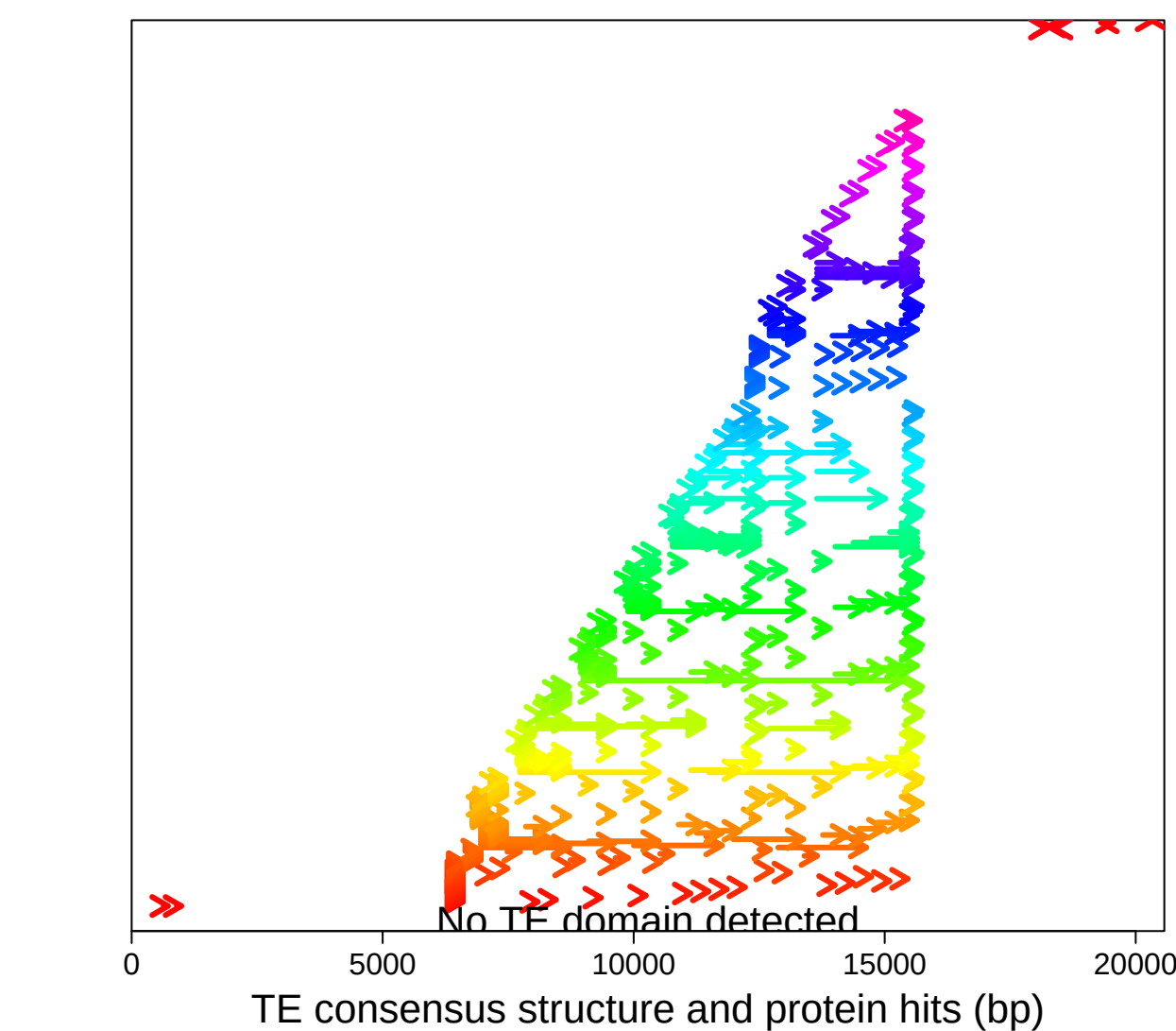
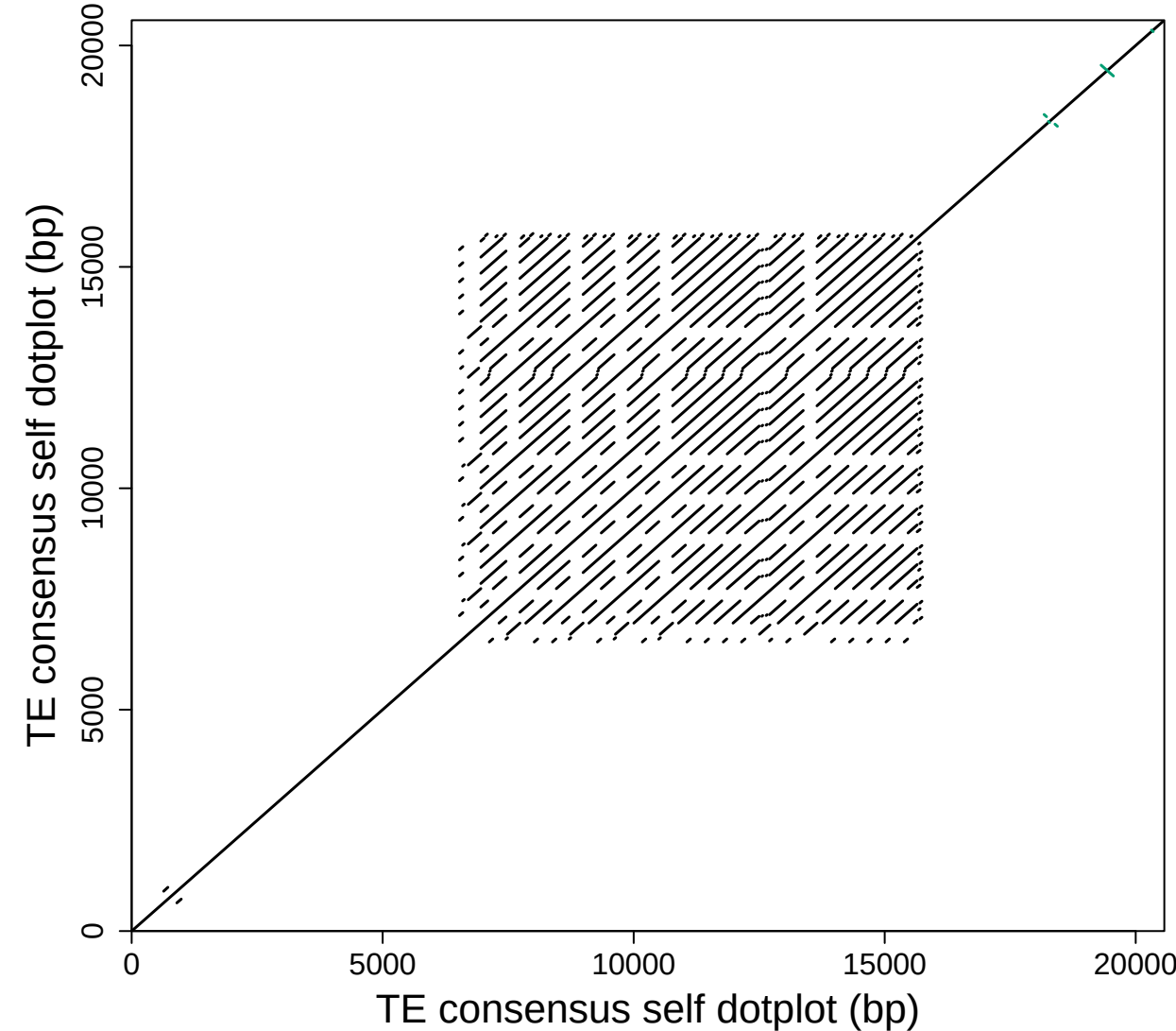
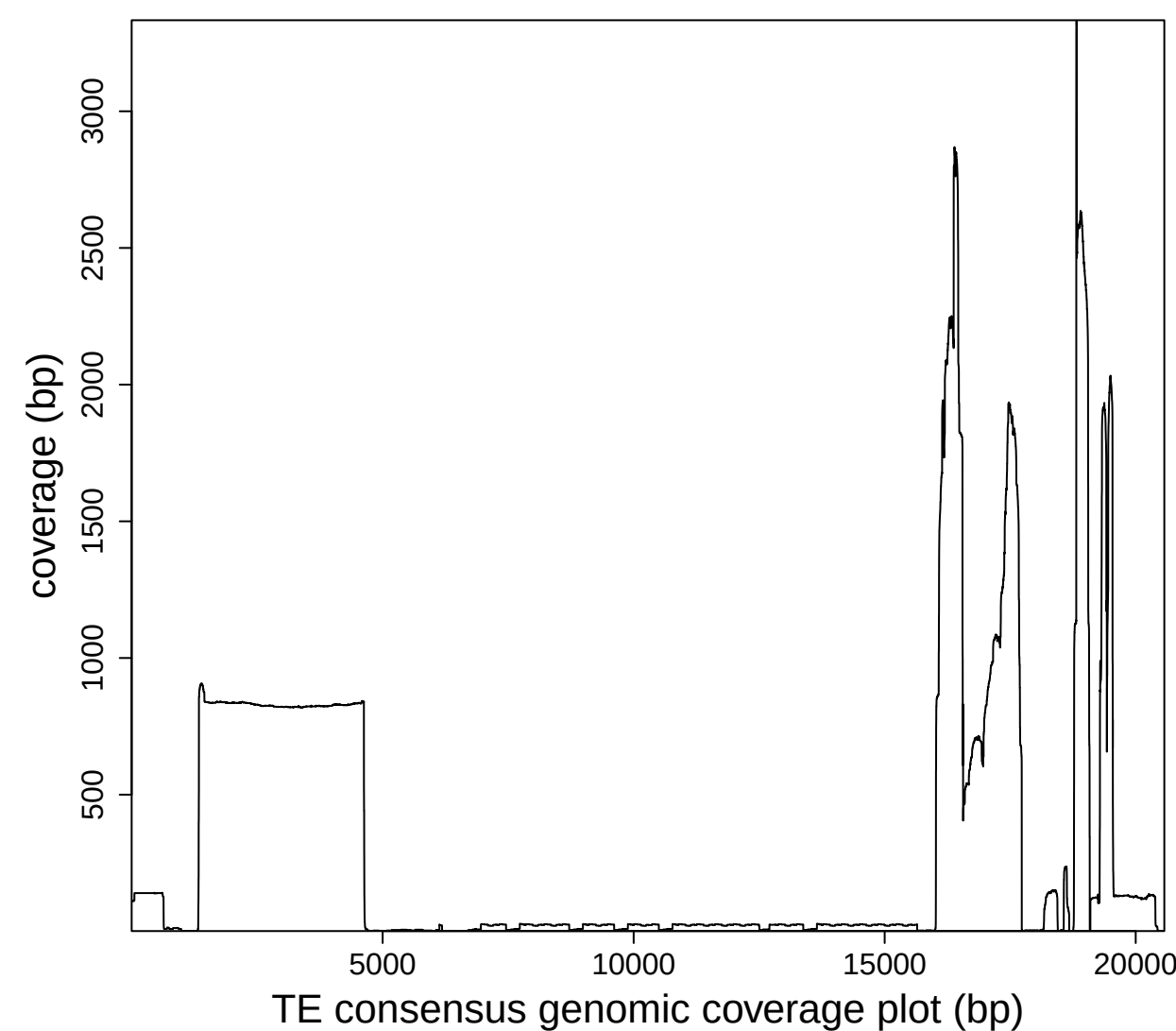
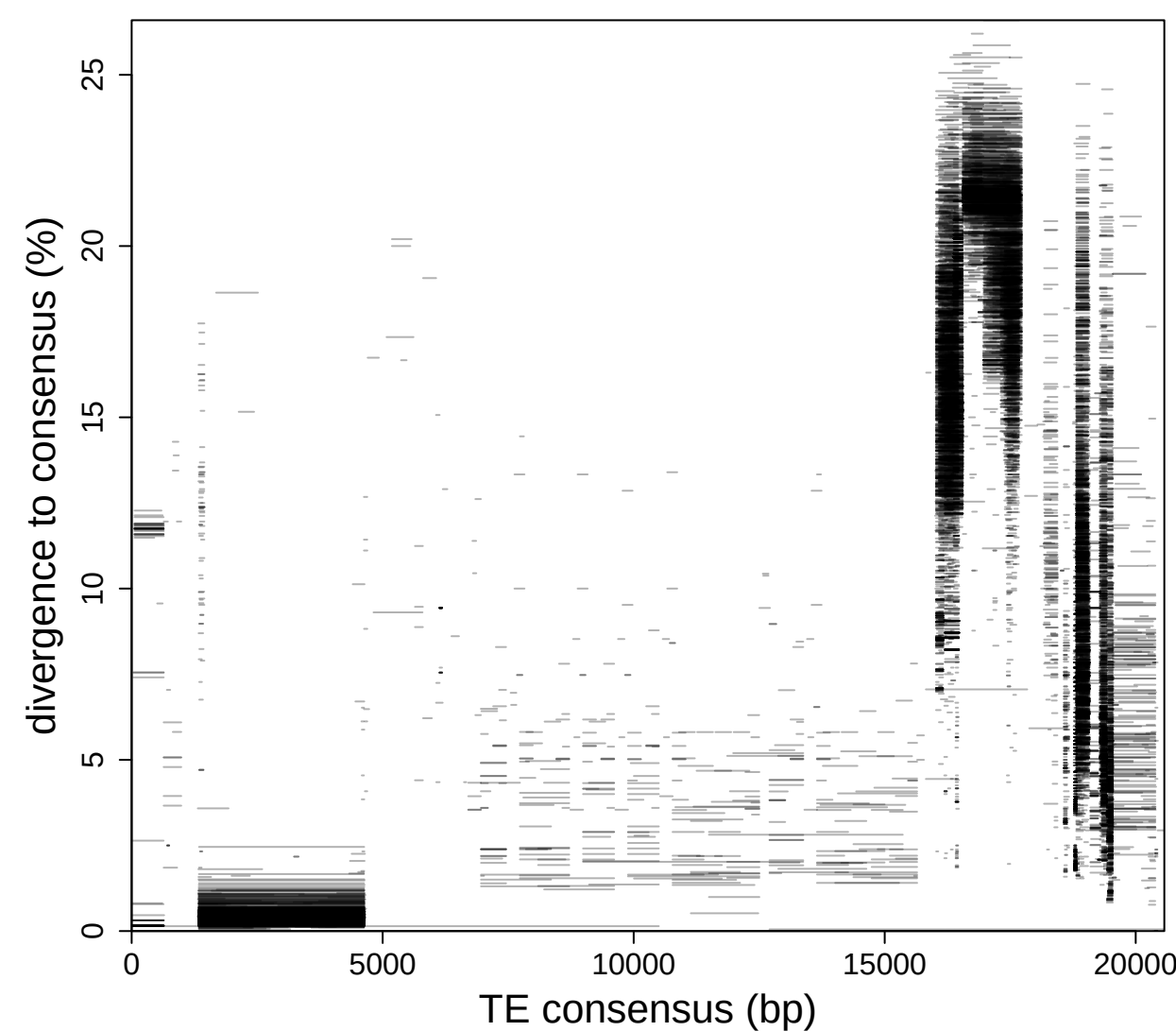
MSA length = 20570





g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 20570bp; fragments: 16160; full length: 0 (>=18513bp)

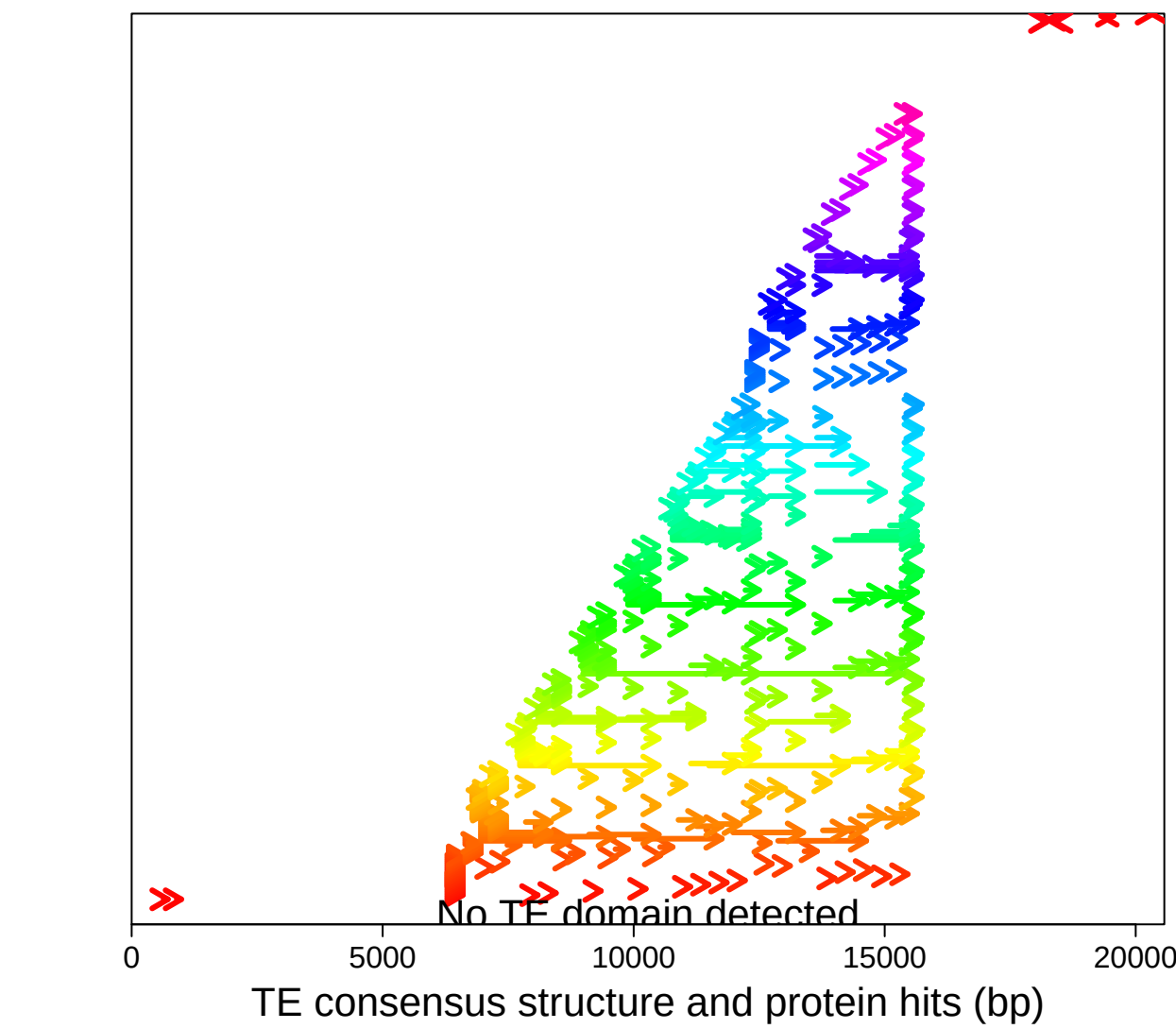
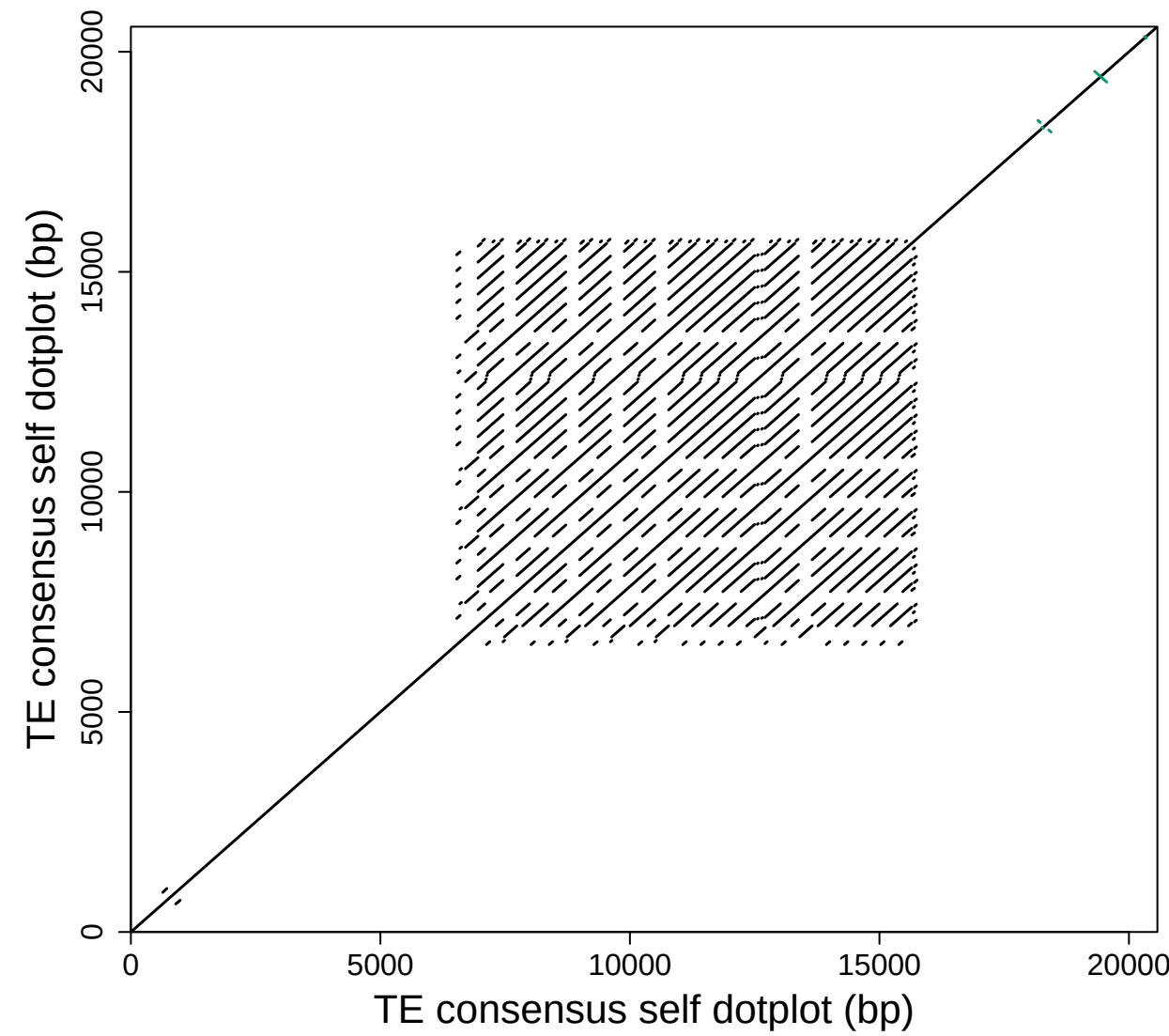
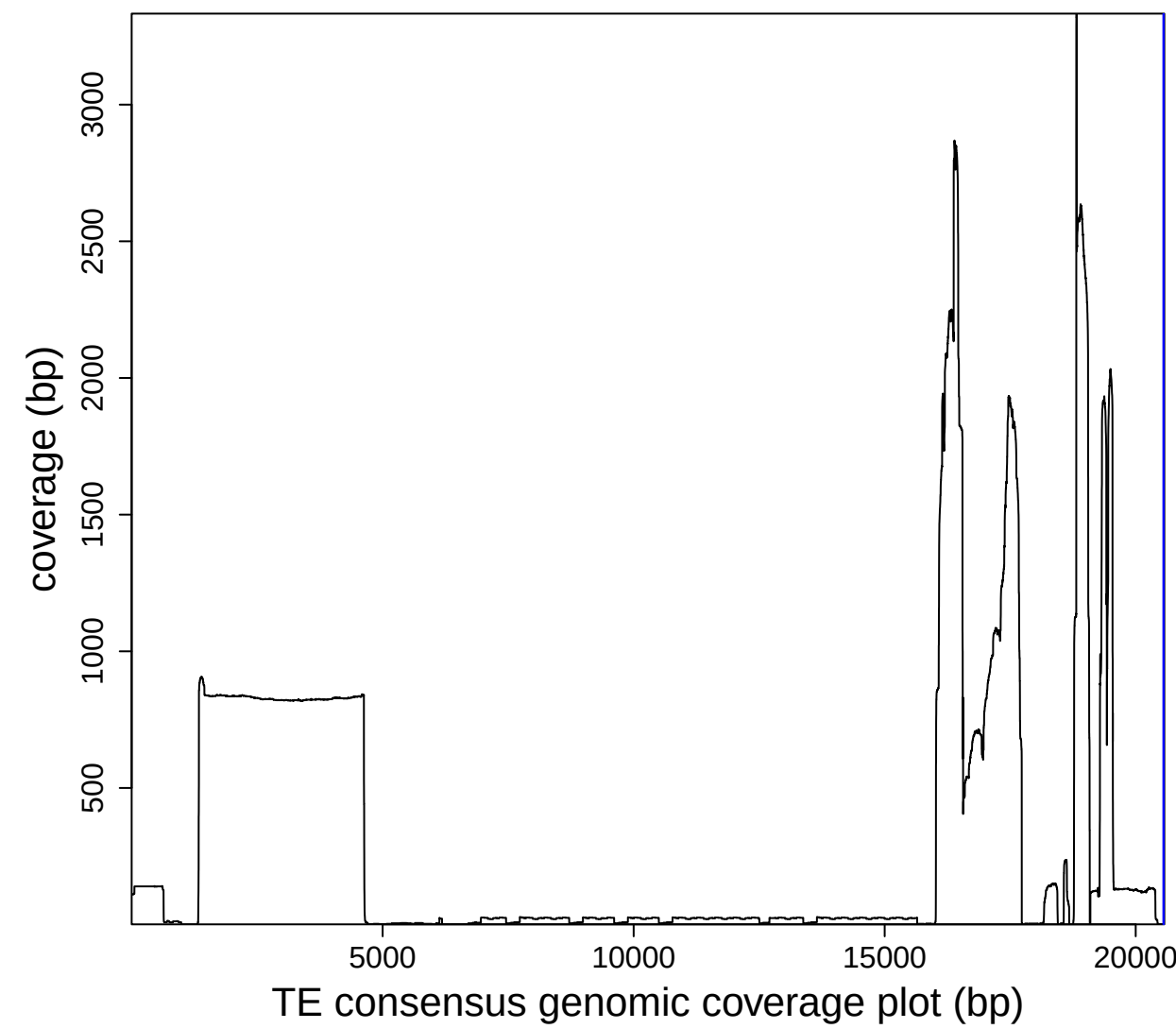
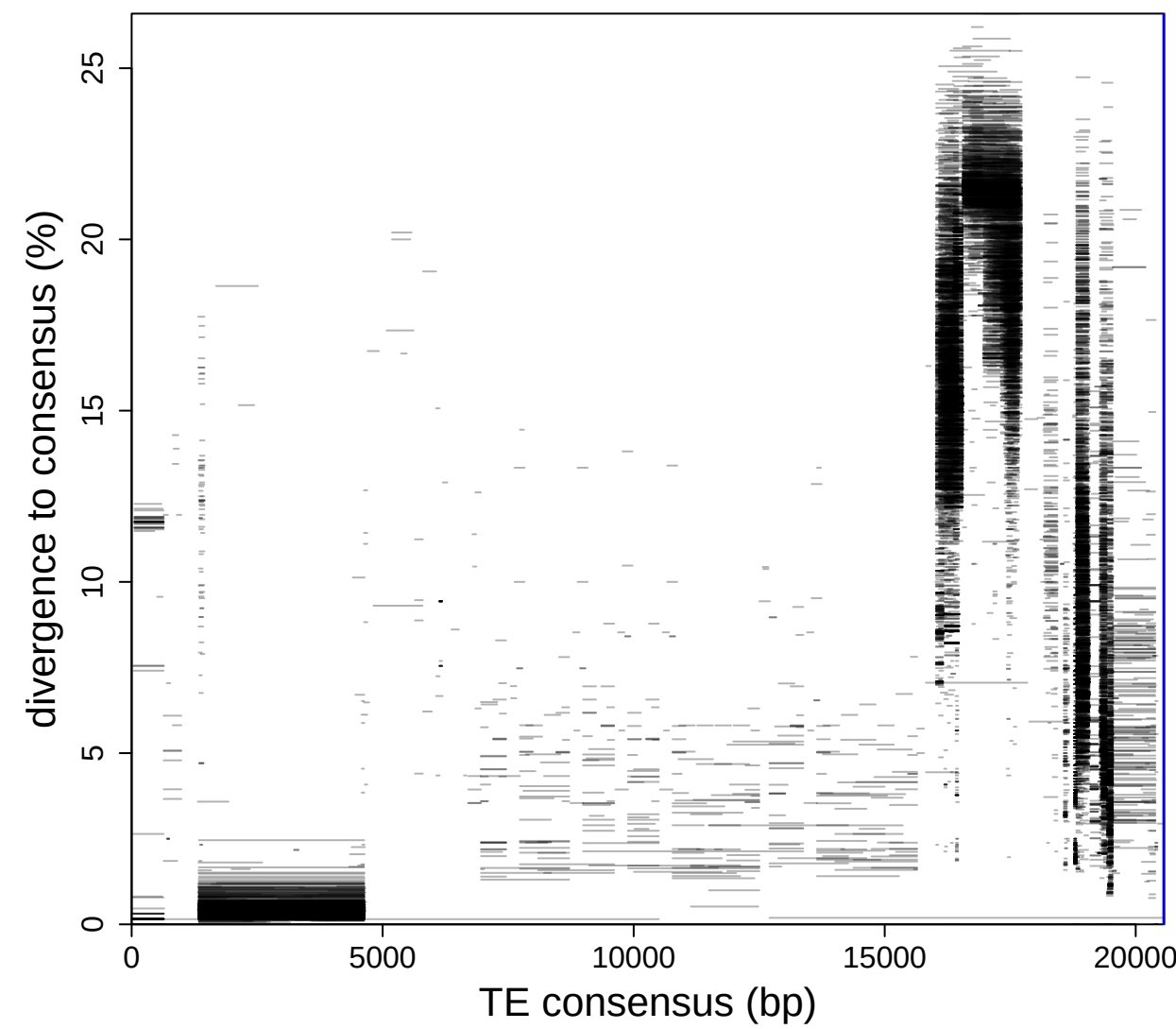
After TEtrimmer 20570 bp



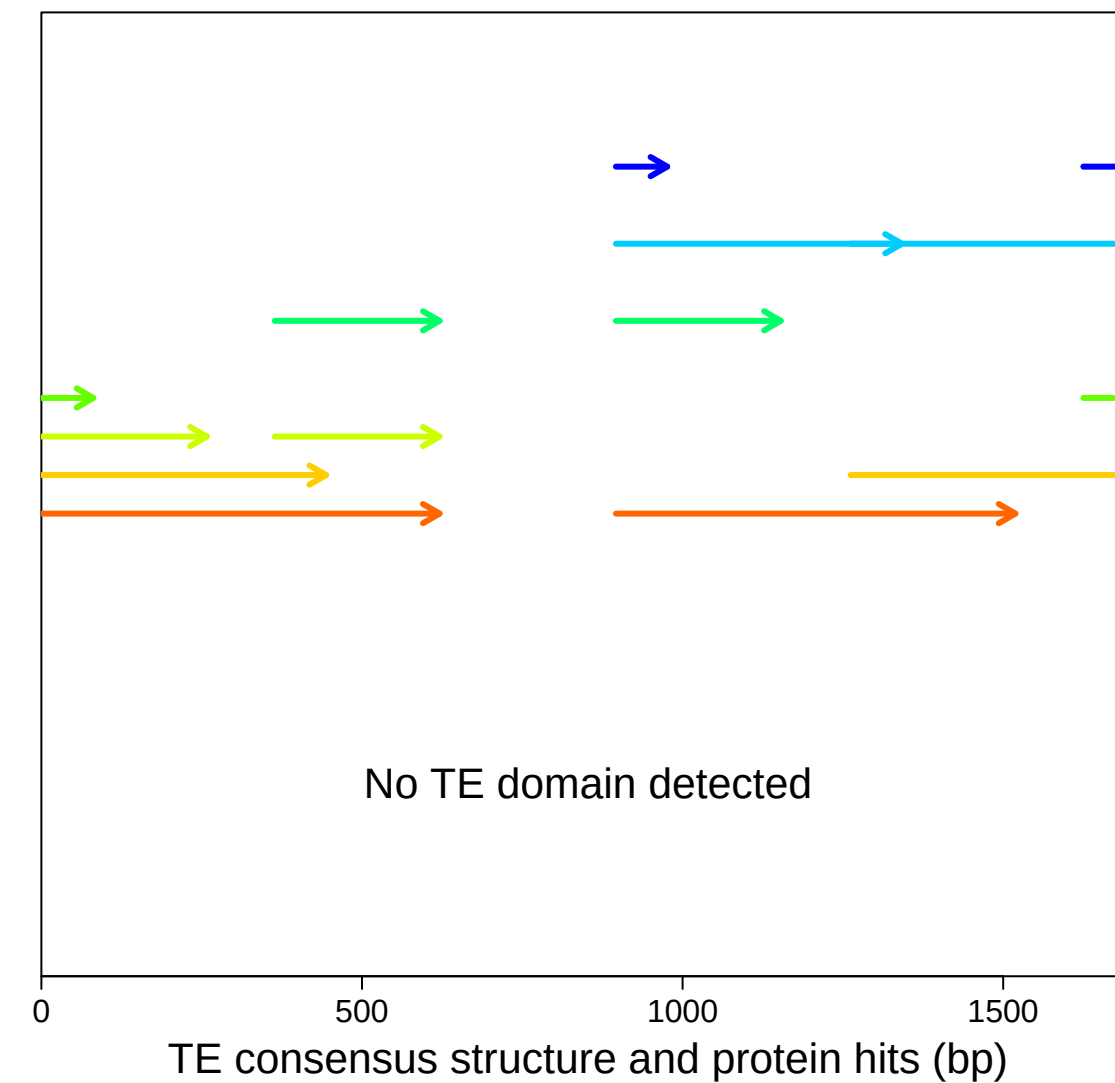
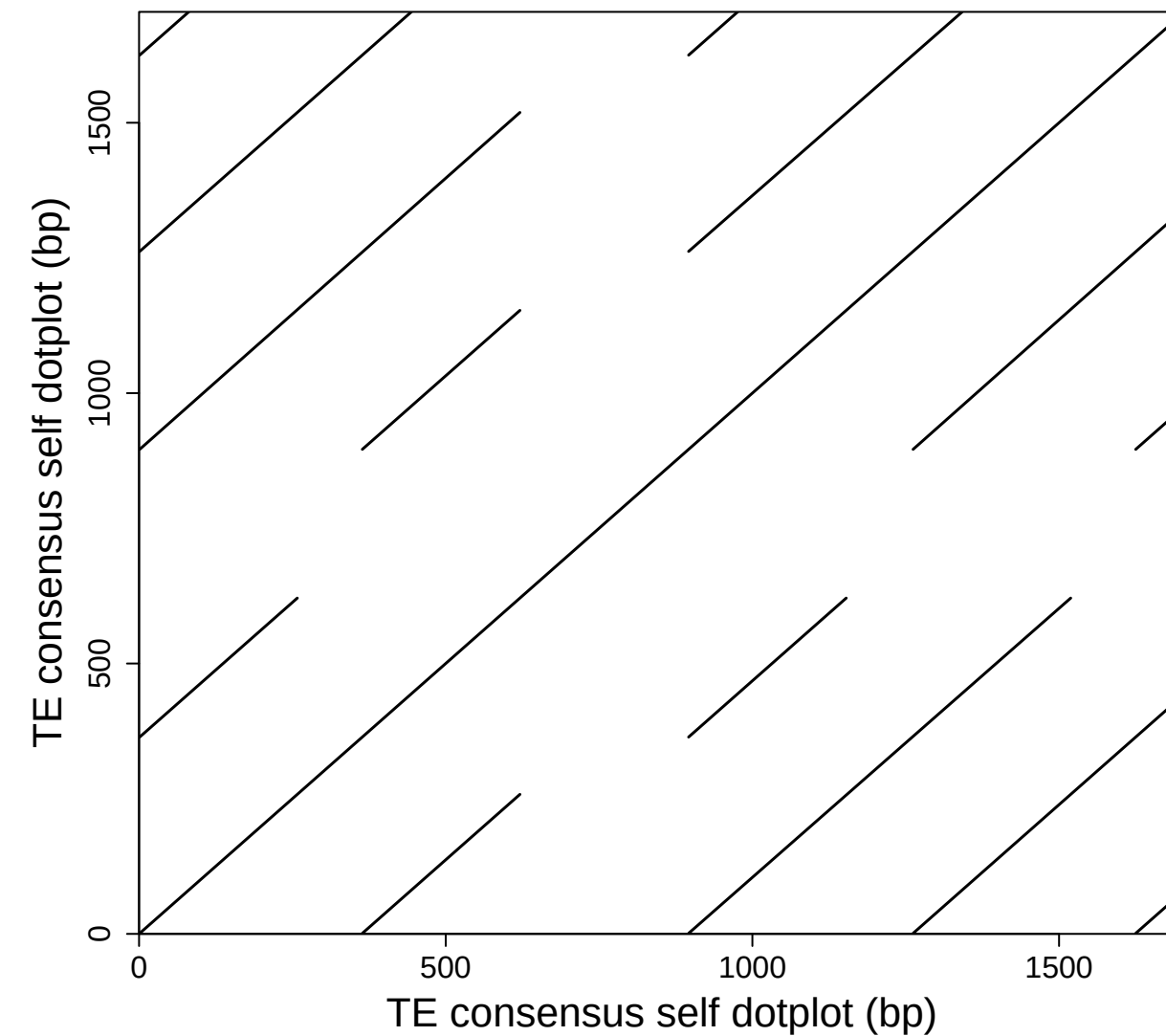
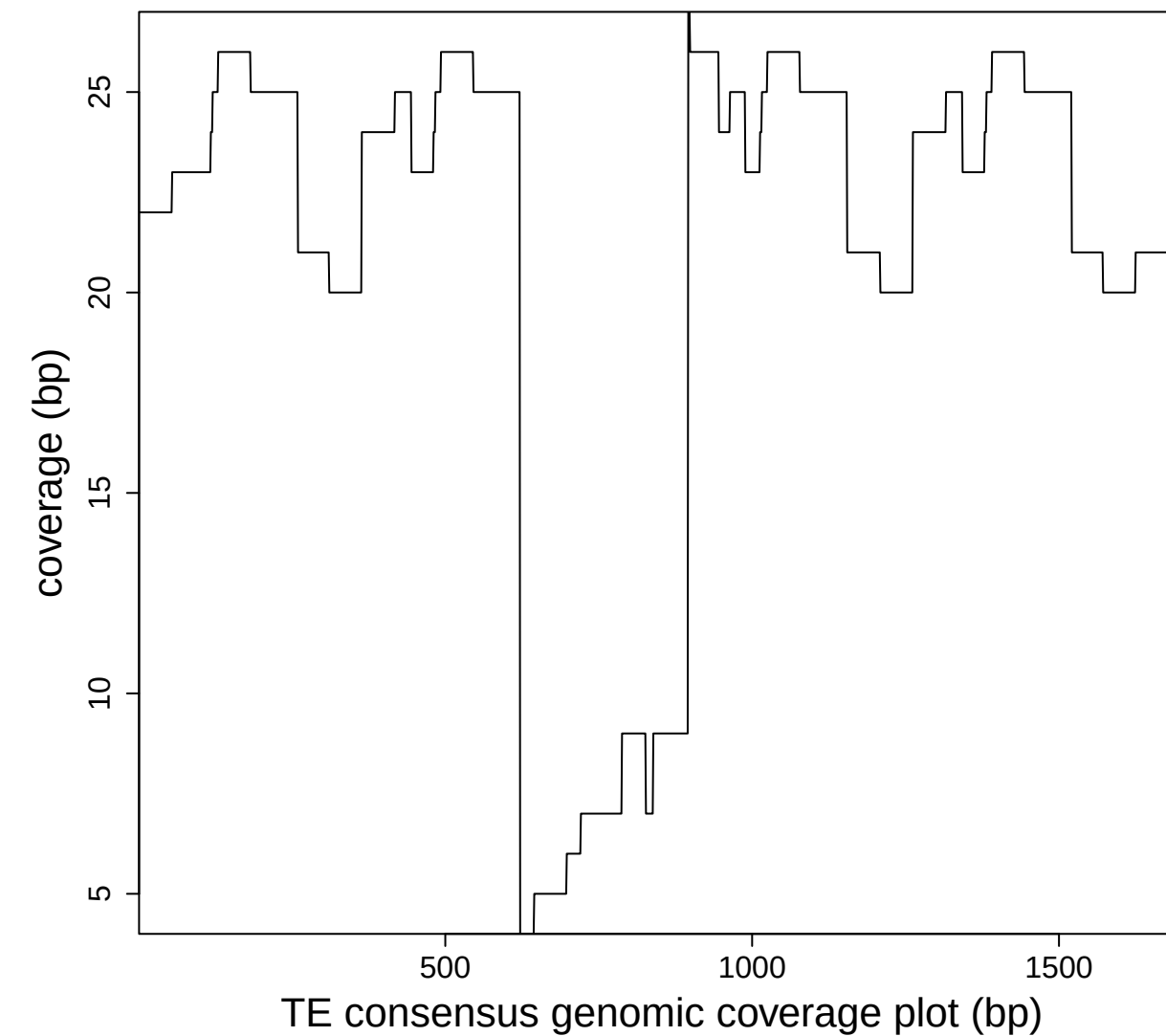
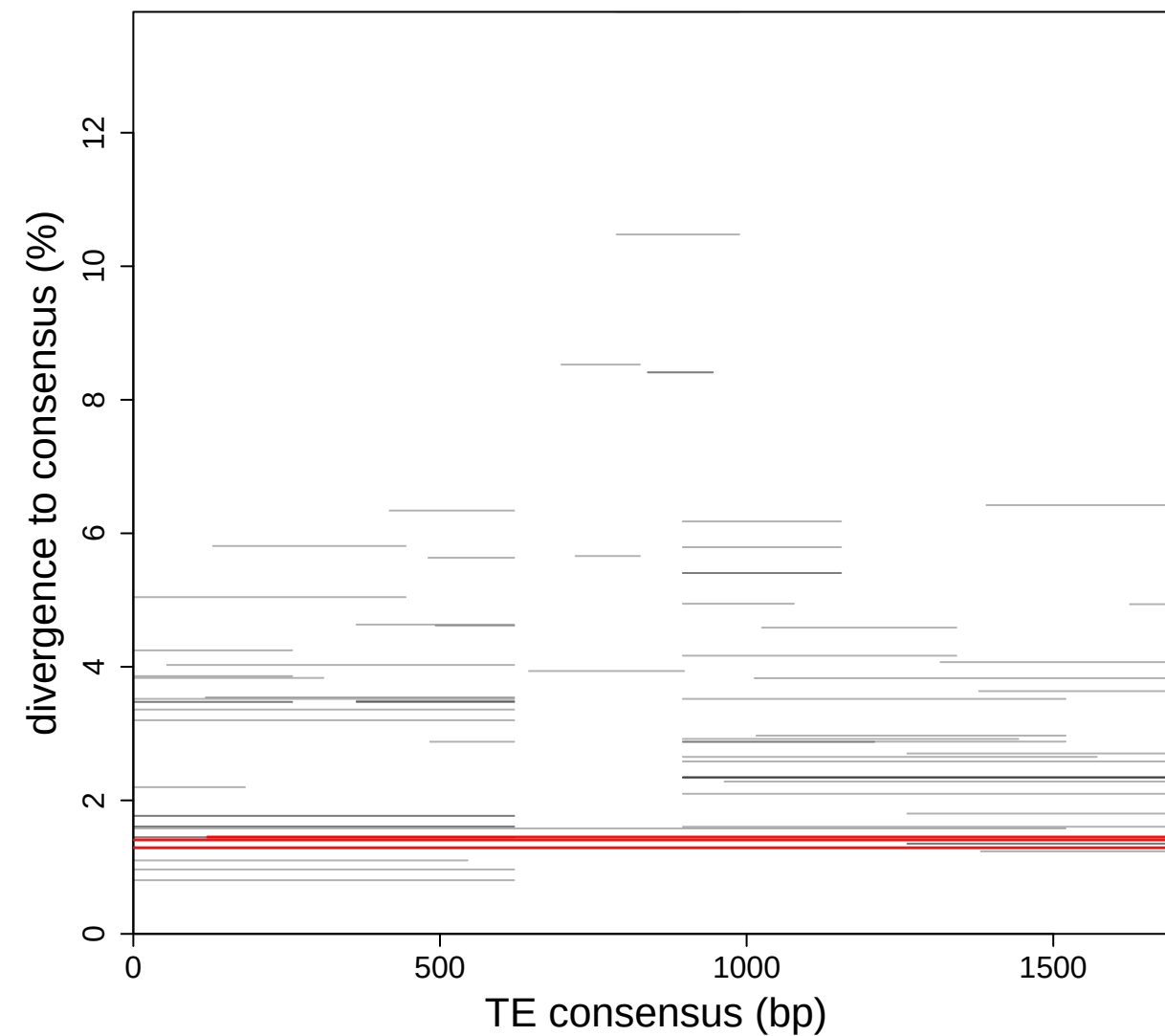
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 20570bp; fragments: 16160; full length: 0 (>=18513bp)

After TEtrimmer Extended plot

Blue lines are boundaries

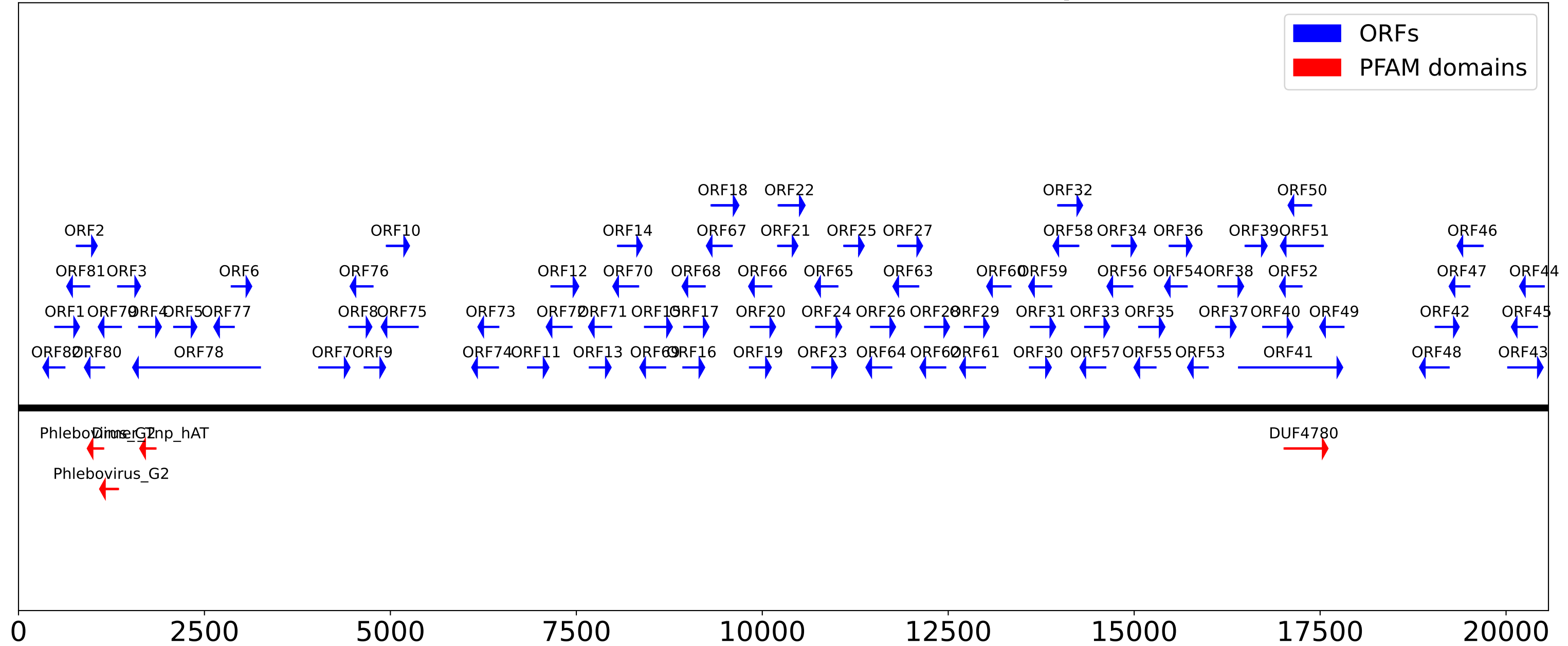


TE: rnd_1_family_618
size: 1705bp; fragments: 72; full length: 3 (≥ 1534.5 bp)

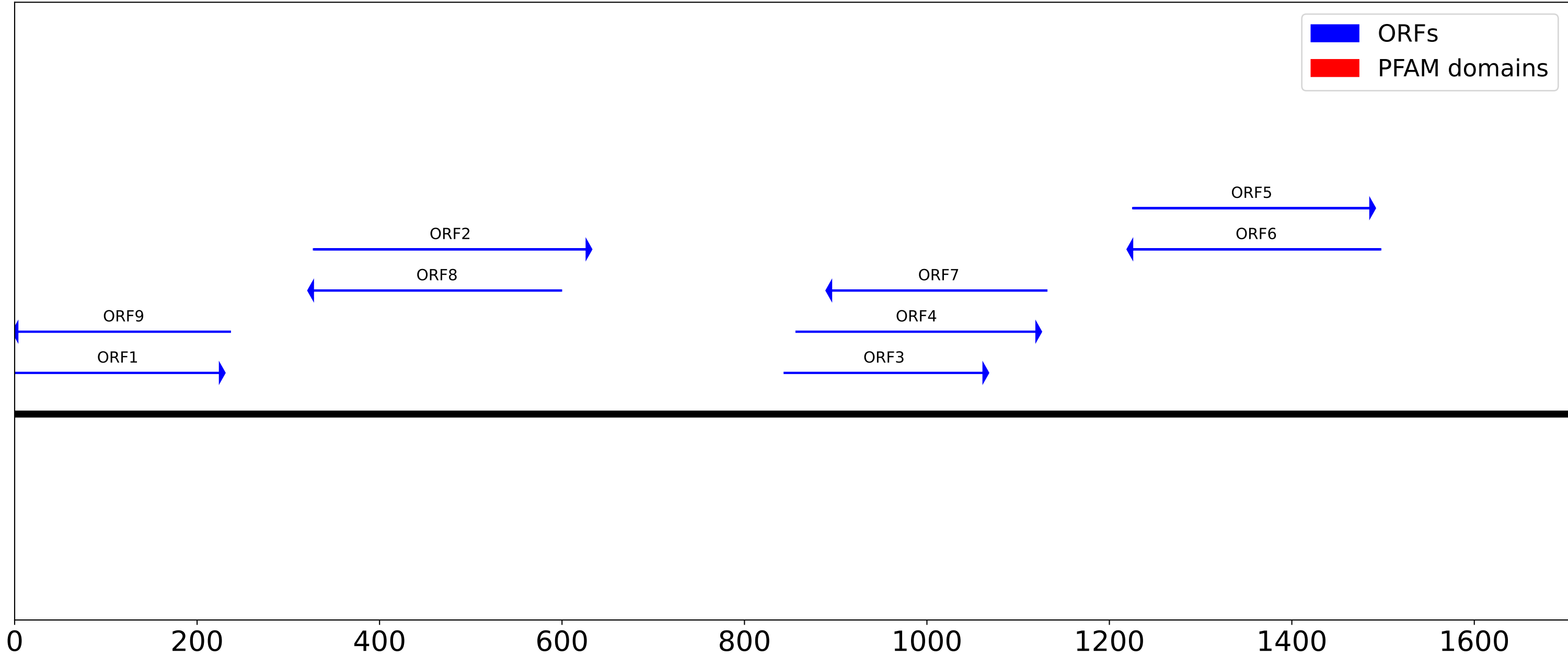


Before TETrimmer 1705 bp

After TETrimmer ORF and PFAM domain plot



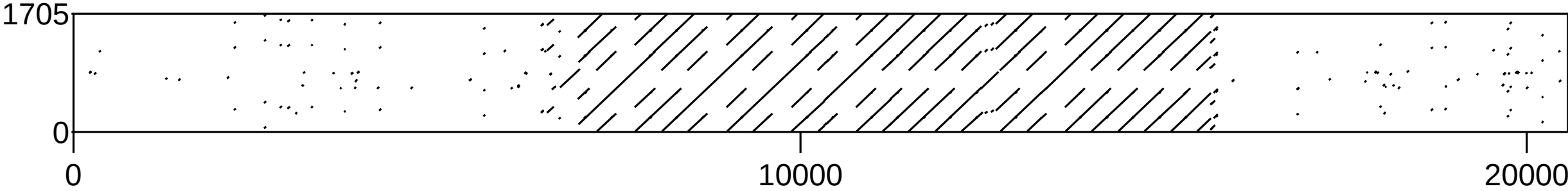
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

ensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)