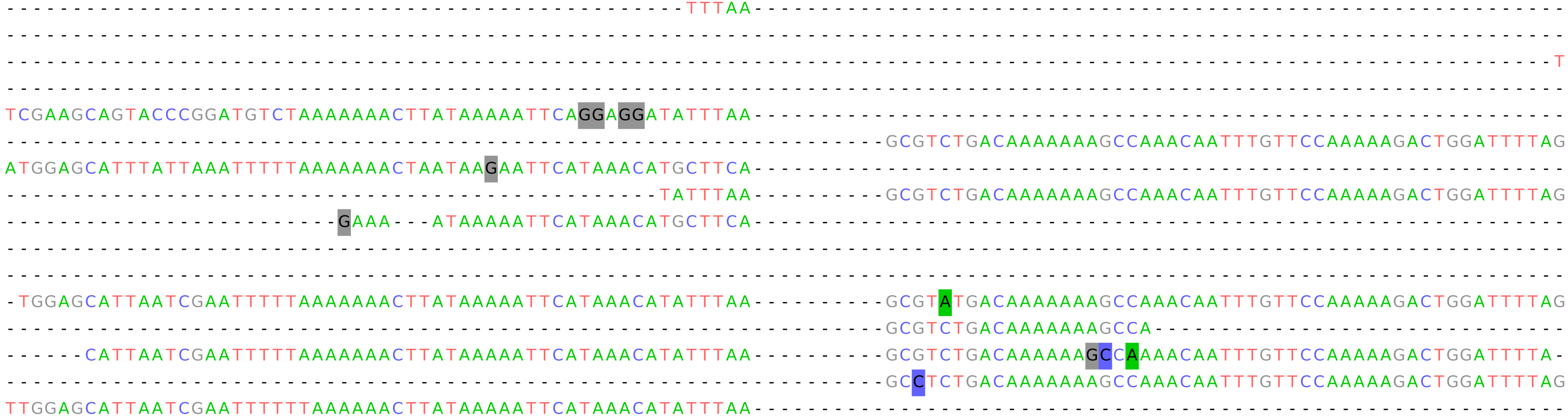


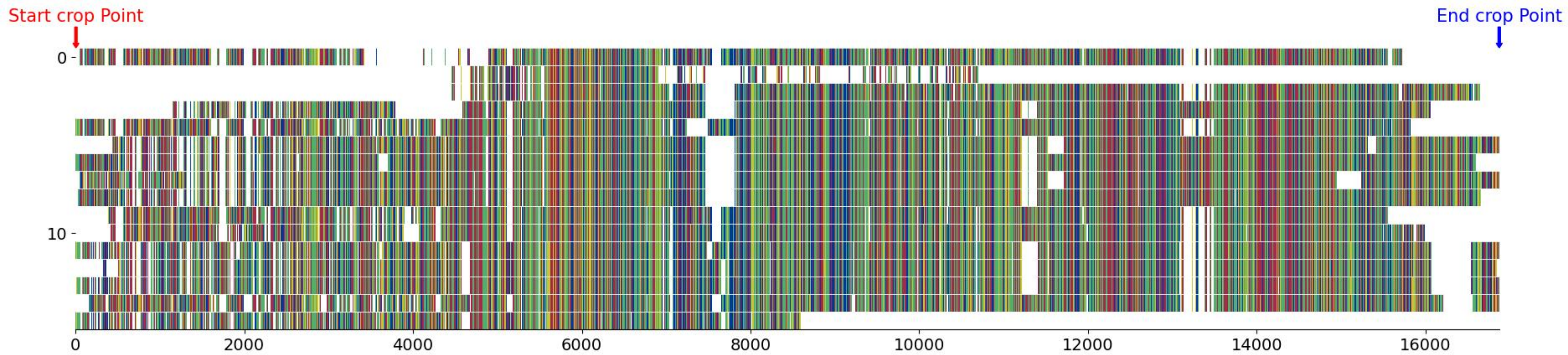
Start crop Point

End crop Point



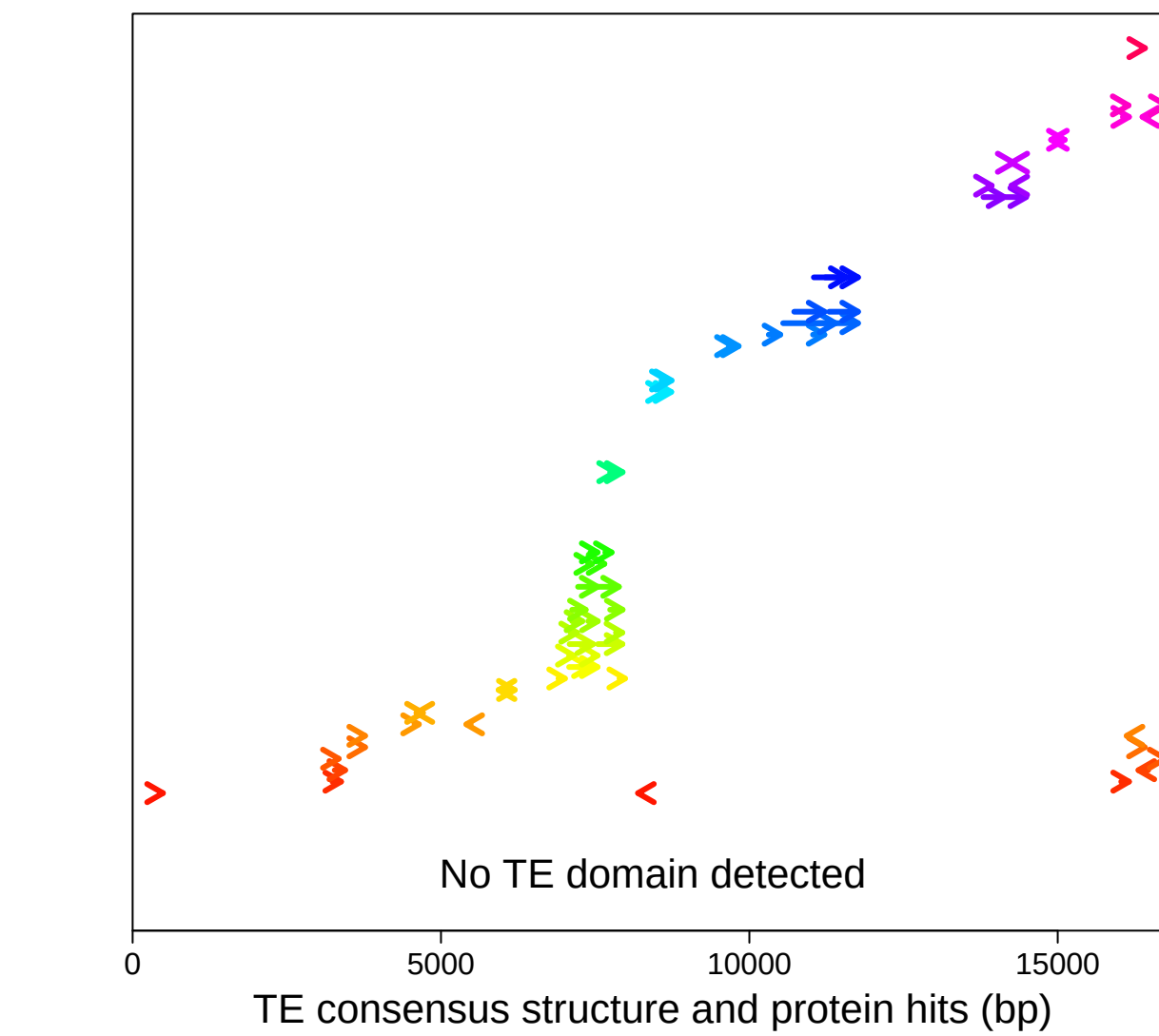
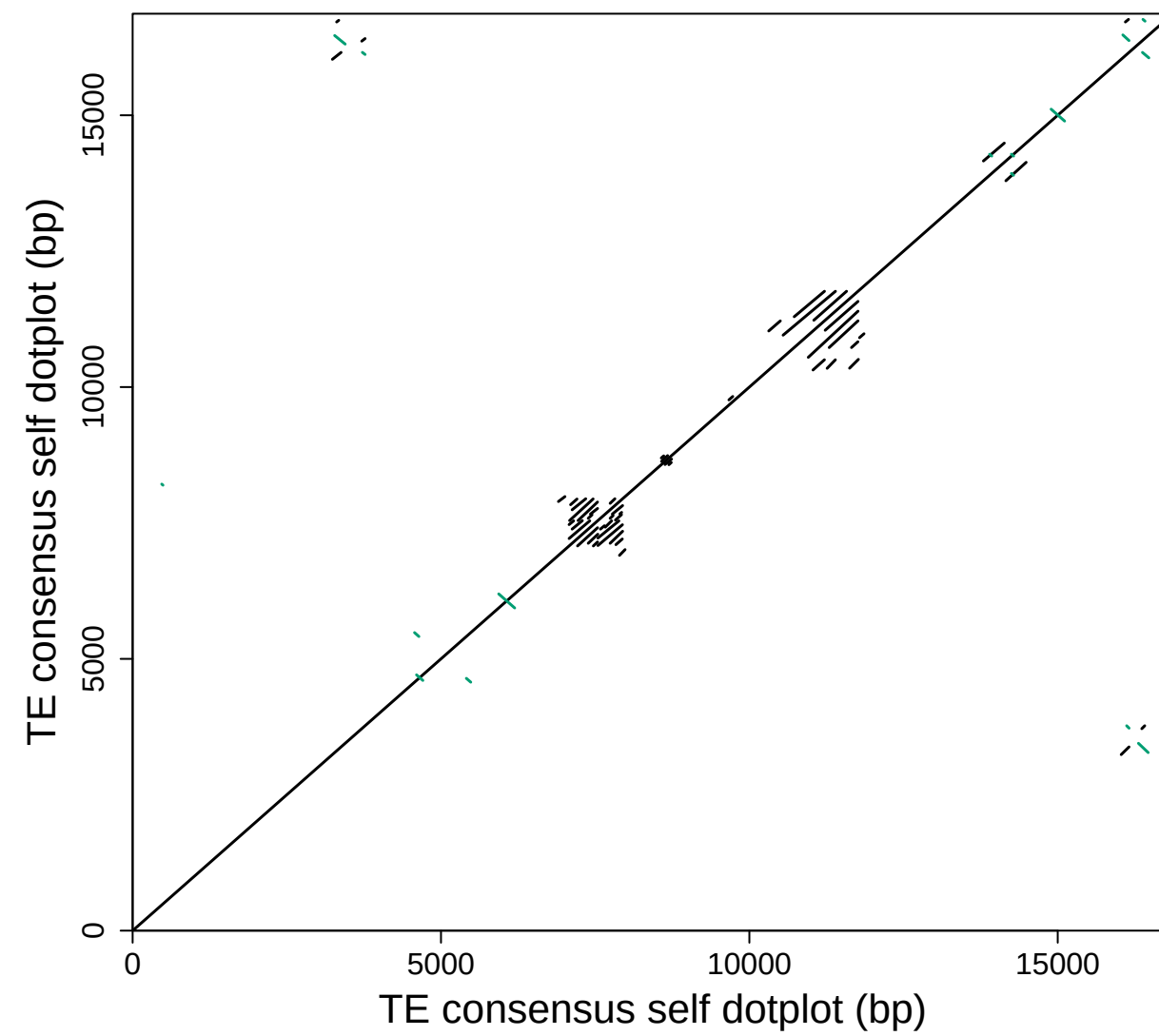
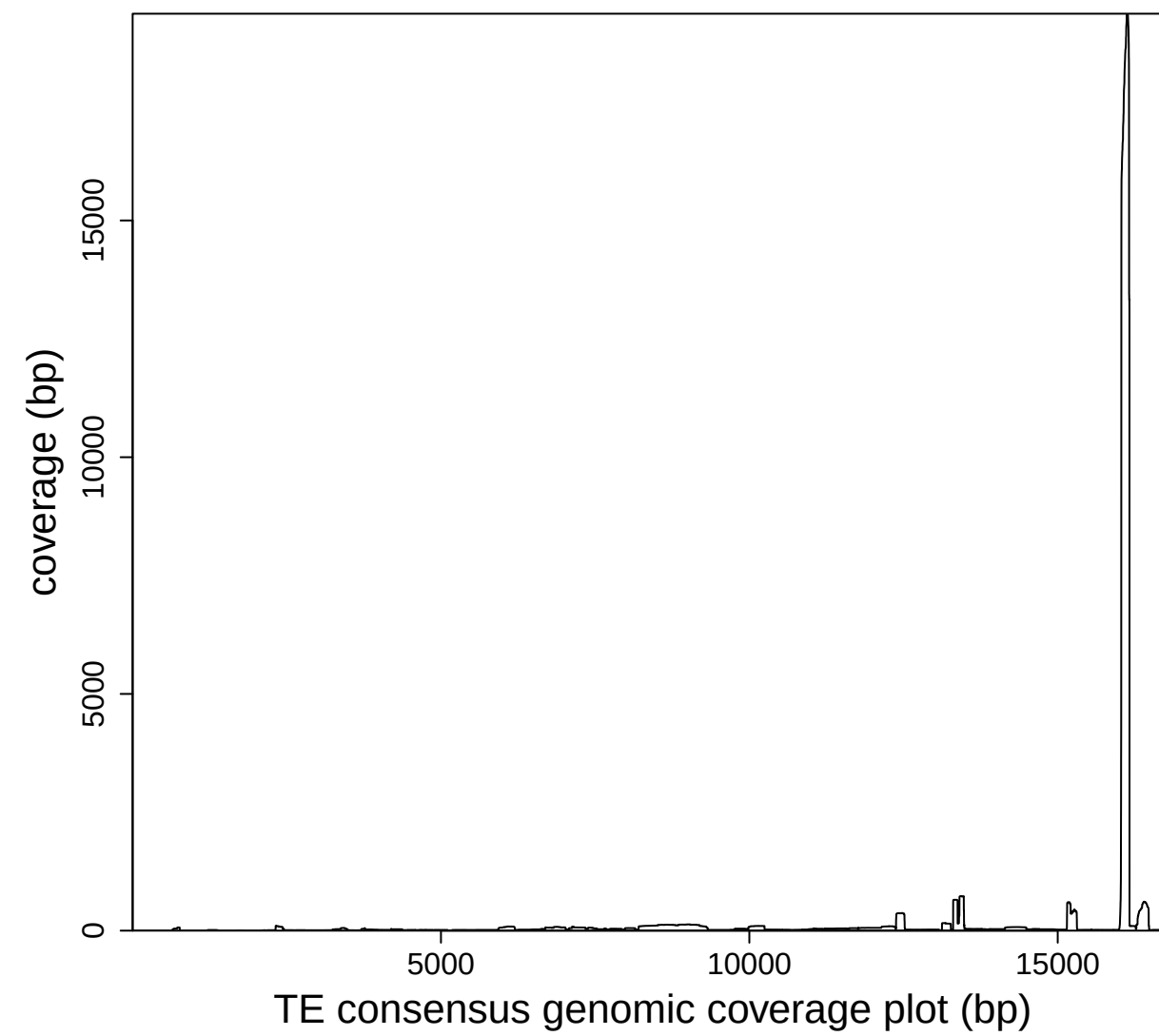
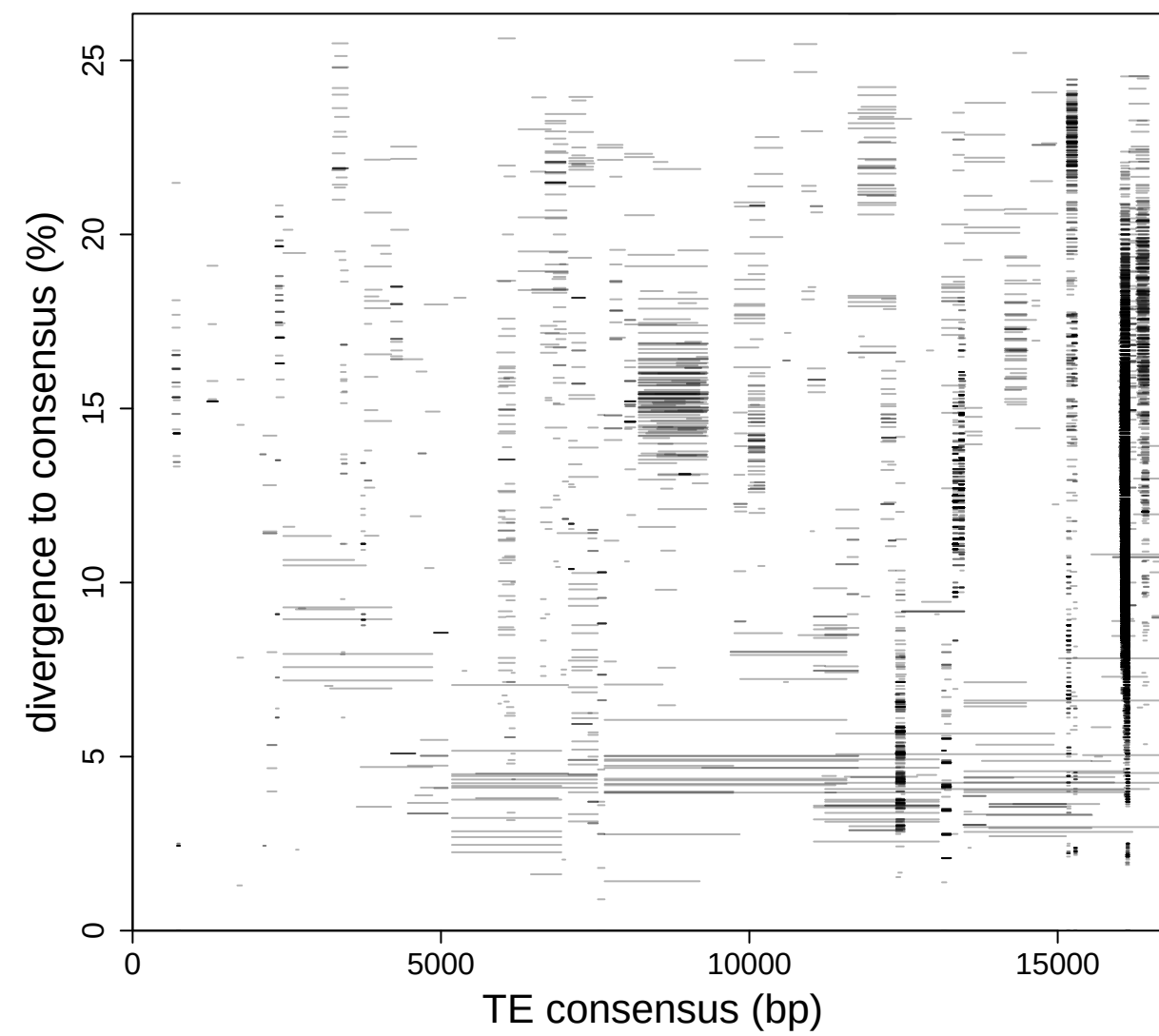
MSA length = 16868





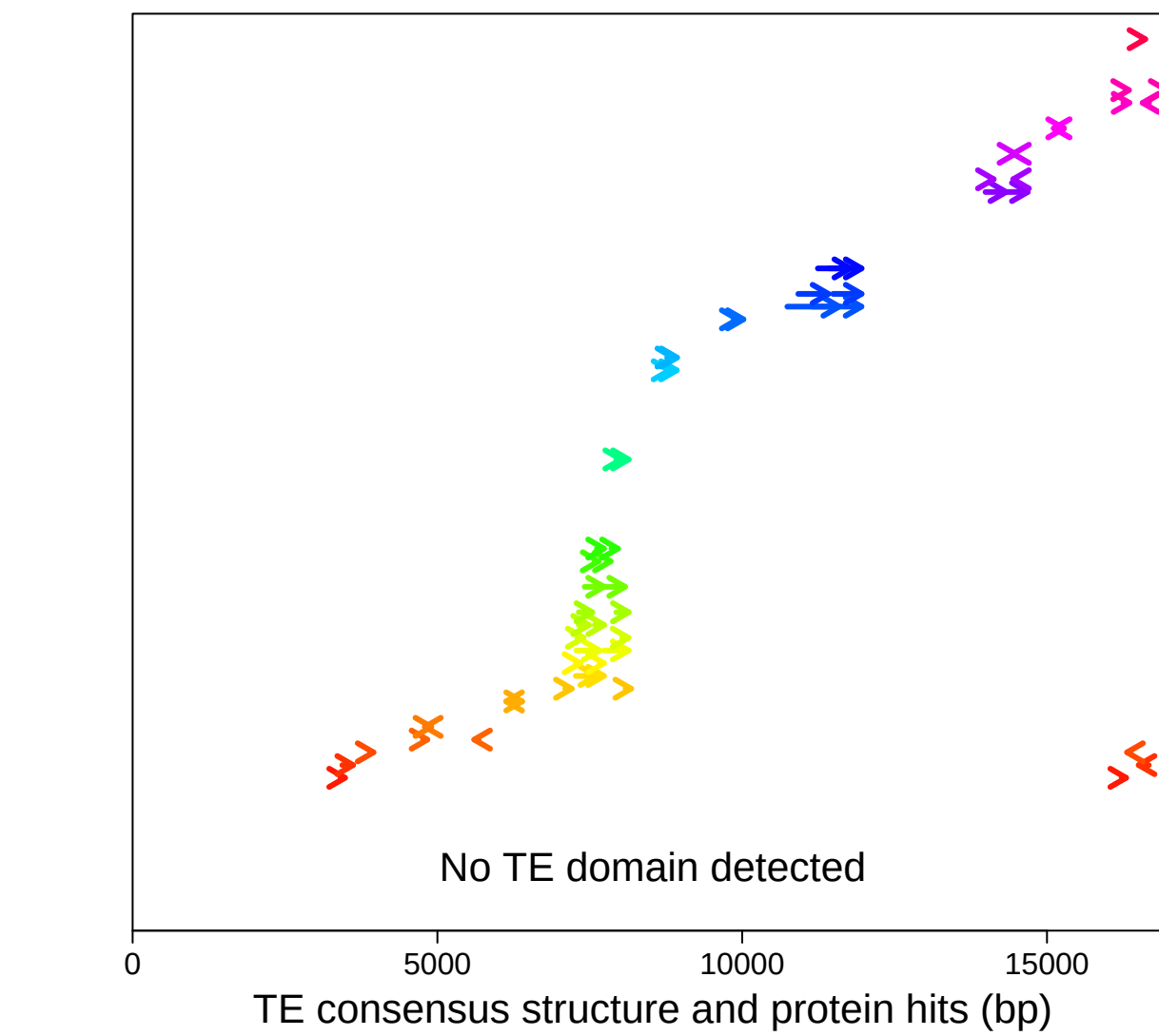
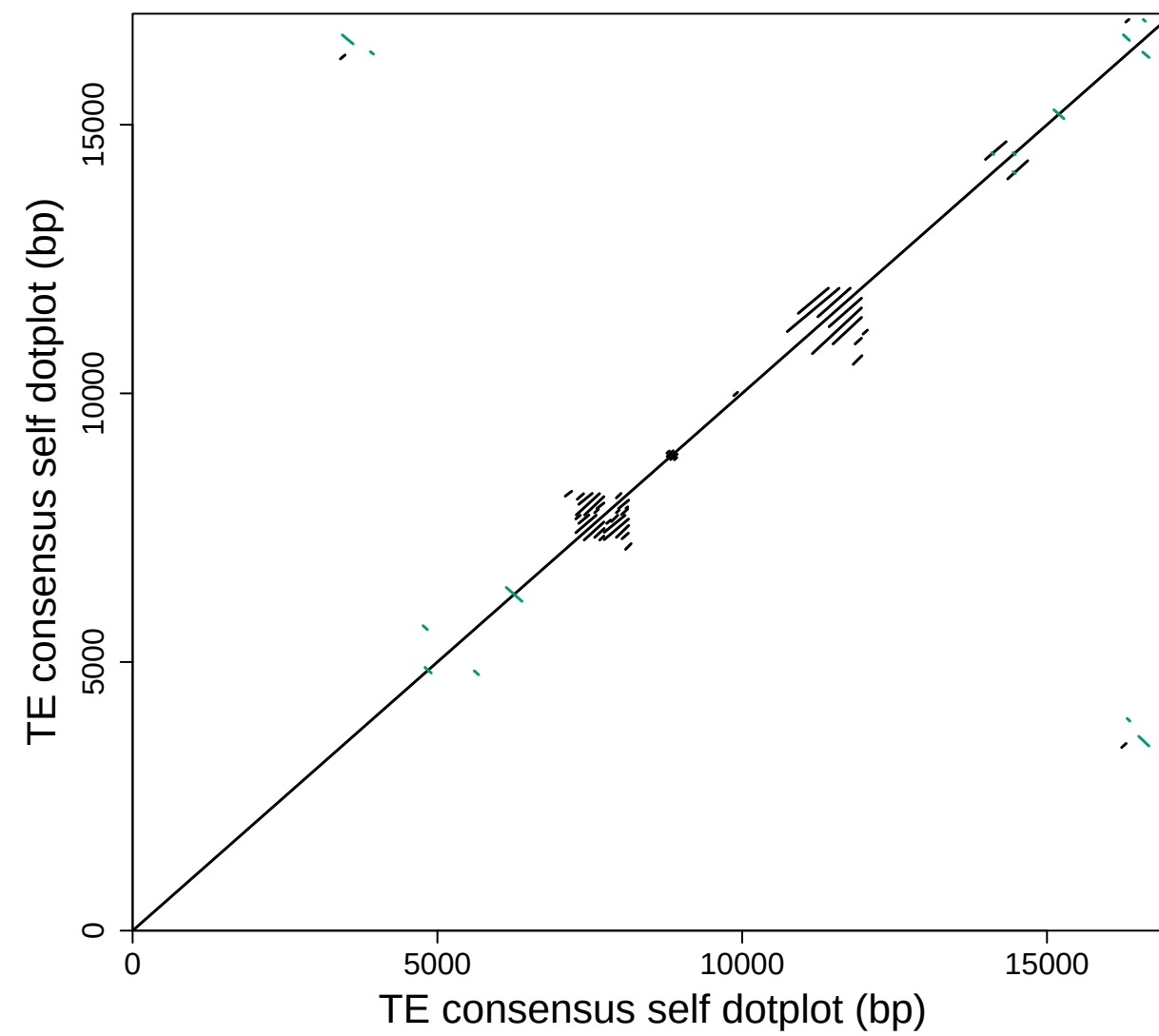
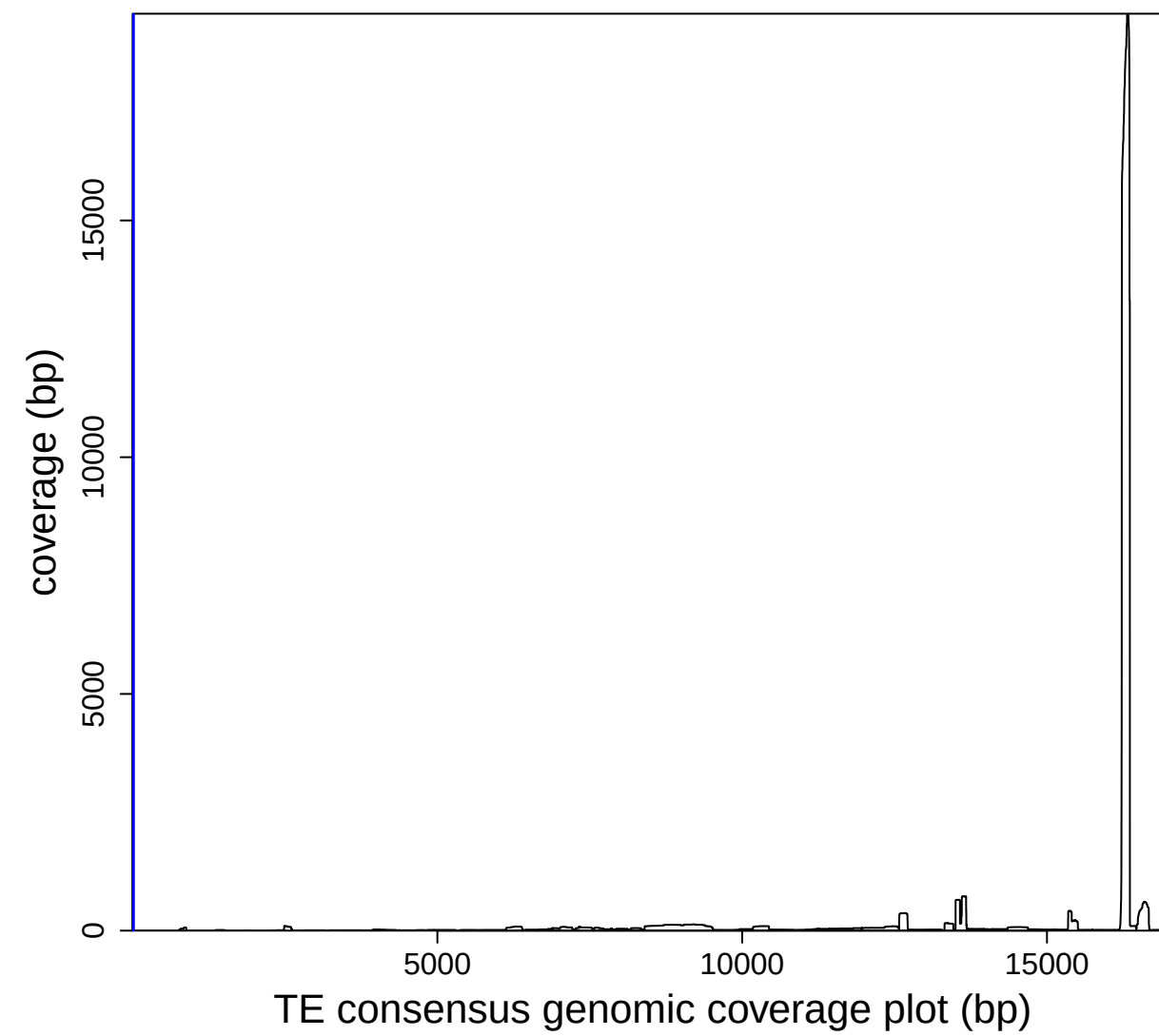
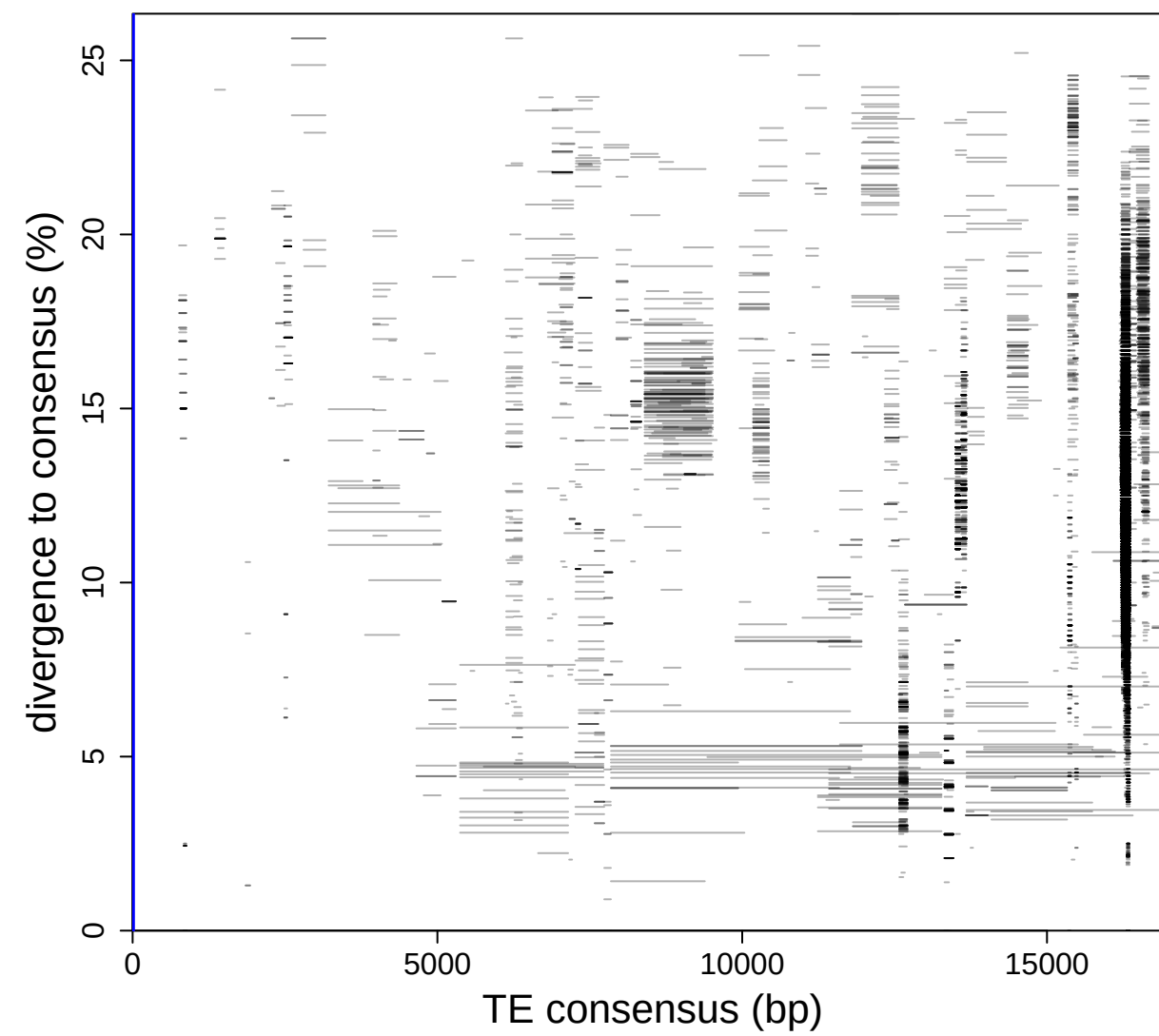
g_1.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln.fa a
size: 16868bp; fragments: 23732; full length: 0 (>=15181.2bp)

After TETrimmer 16868 bp



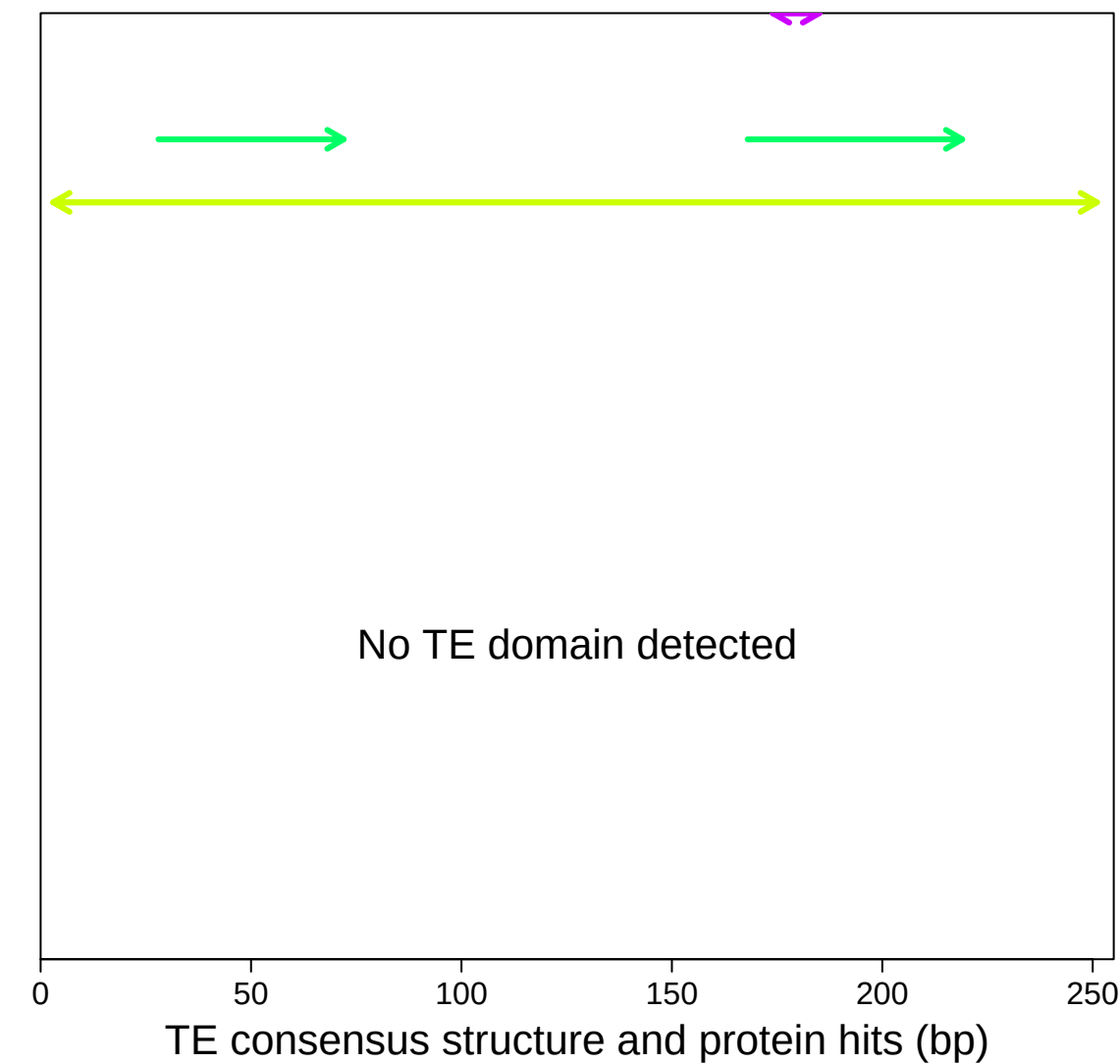
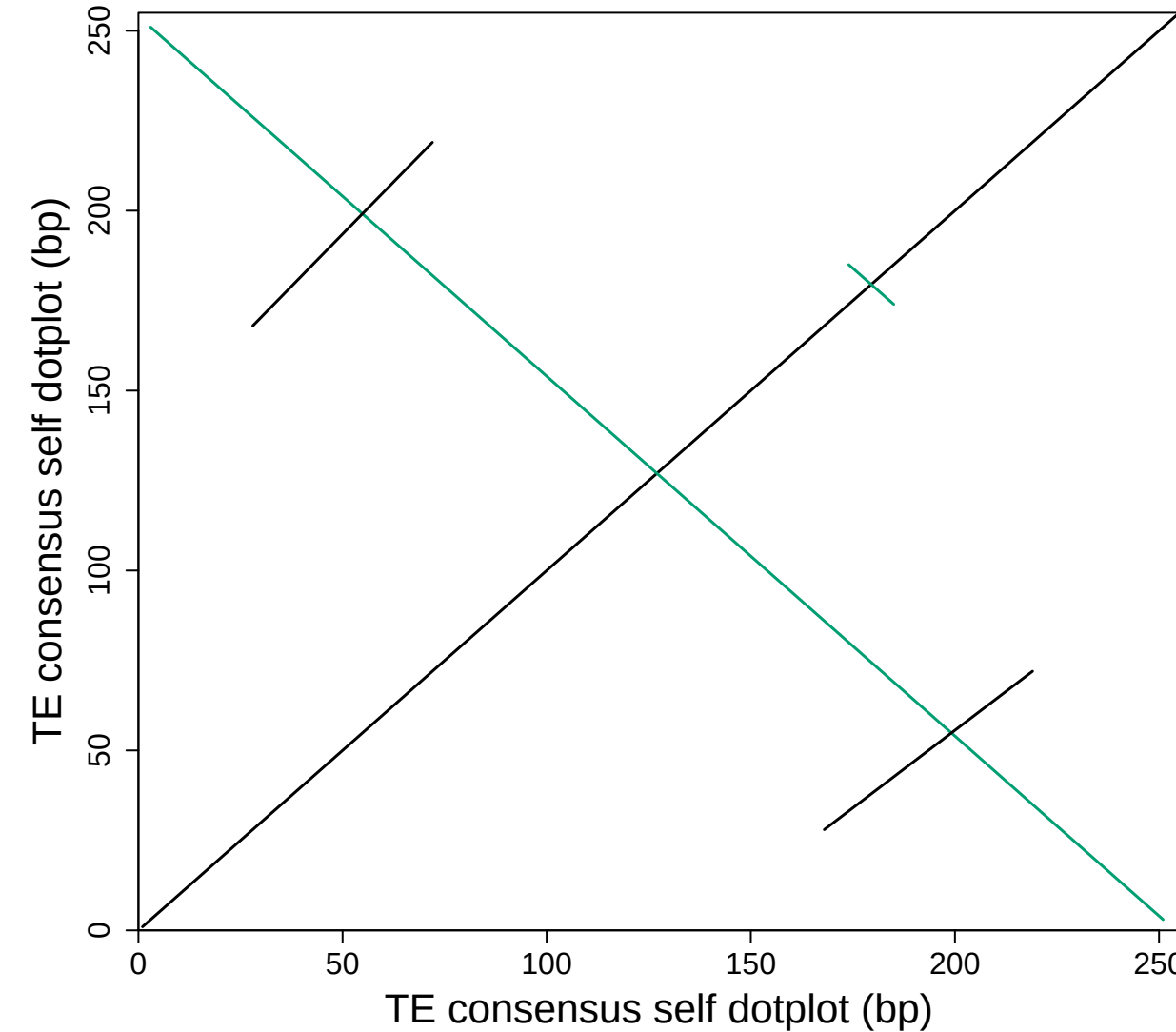
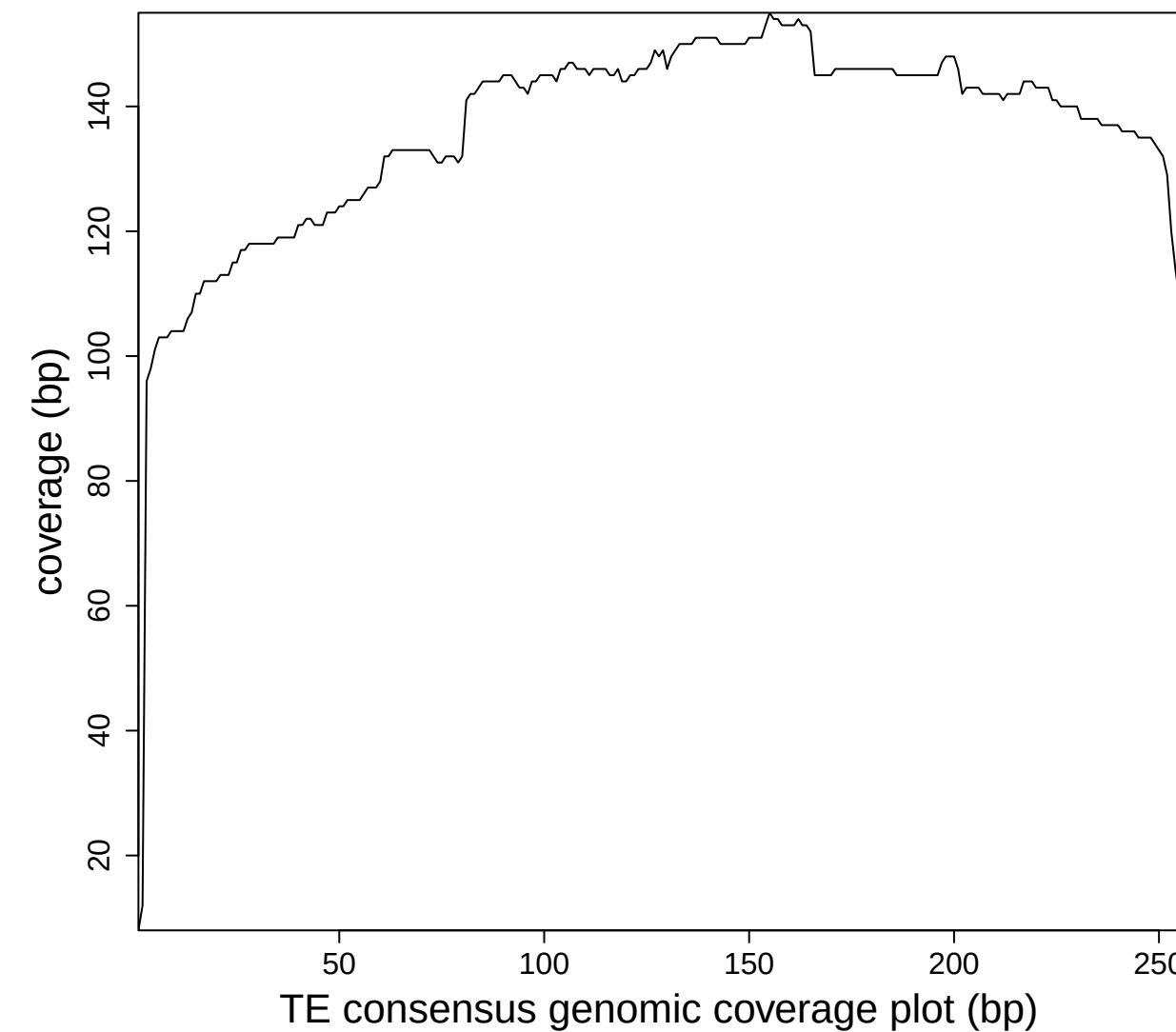
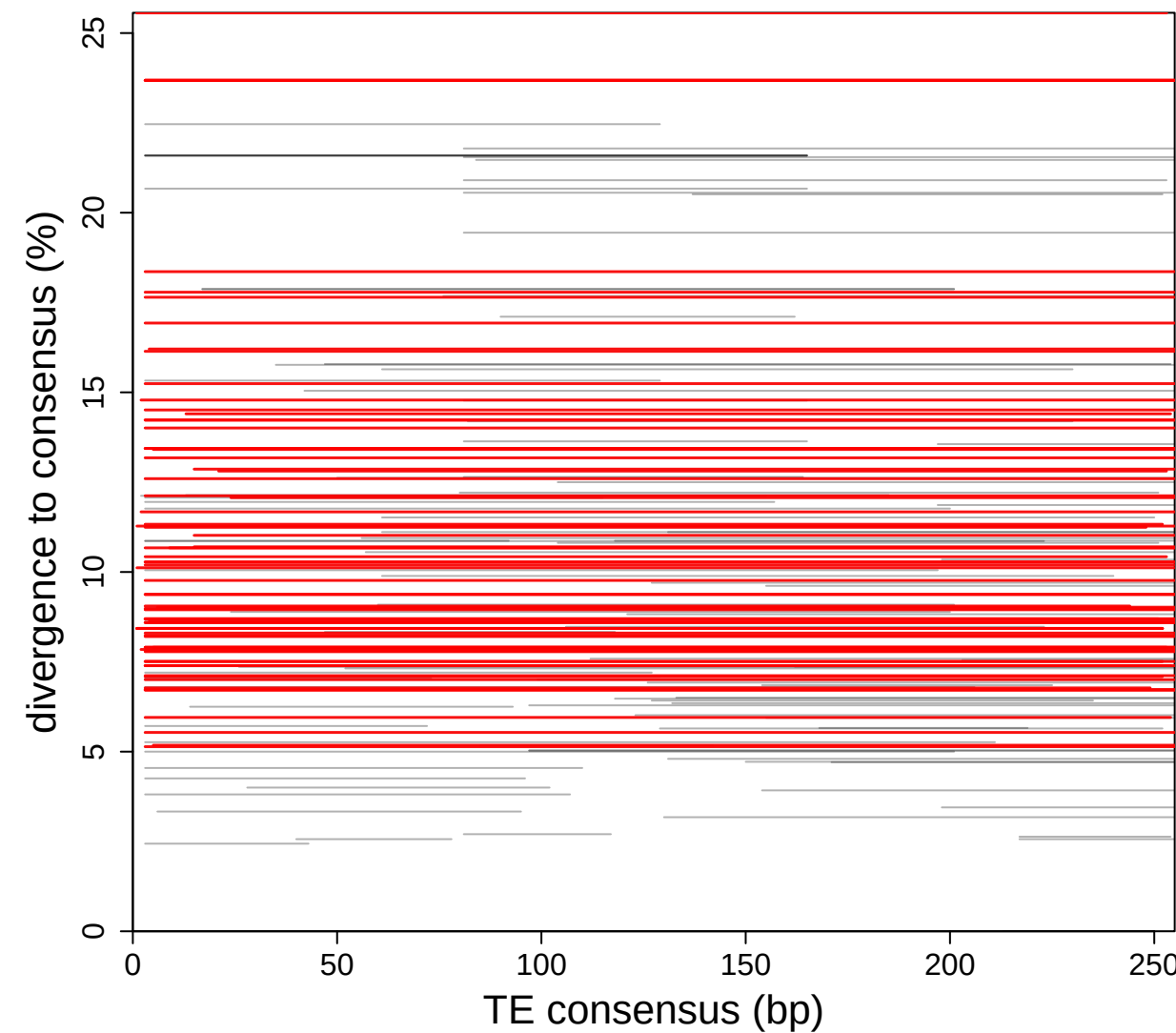
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 17066bp; fragments: 23370; full length: 0 (>=15359.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

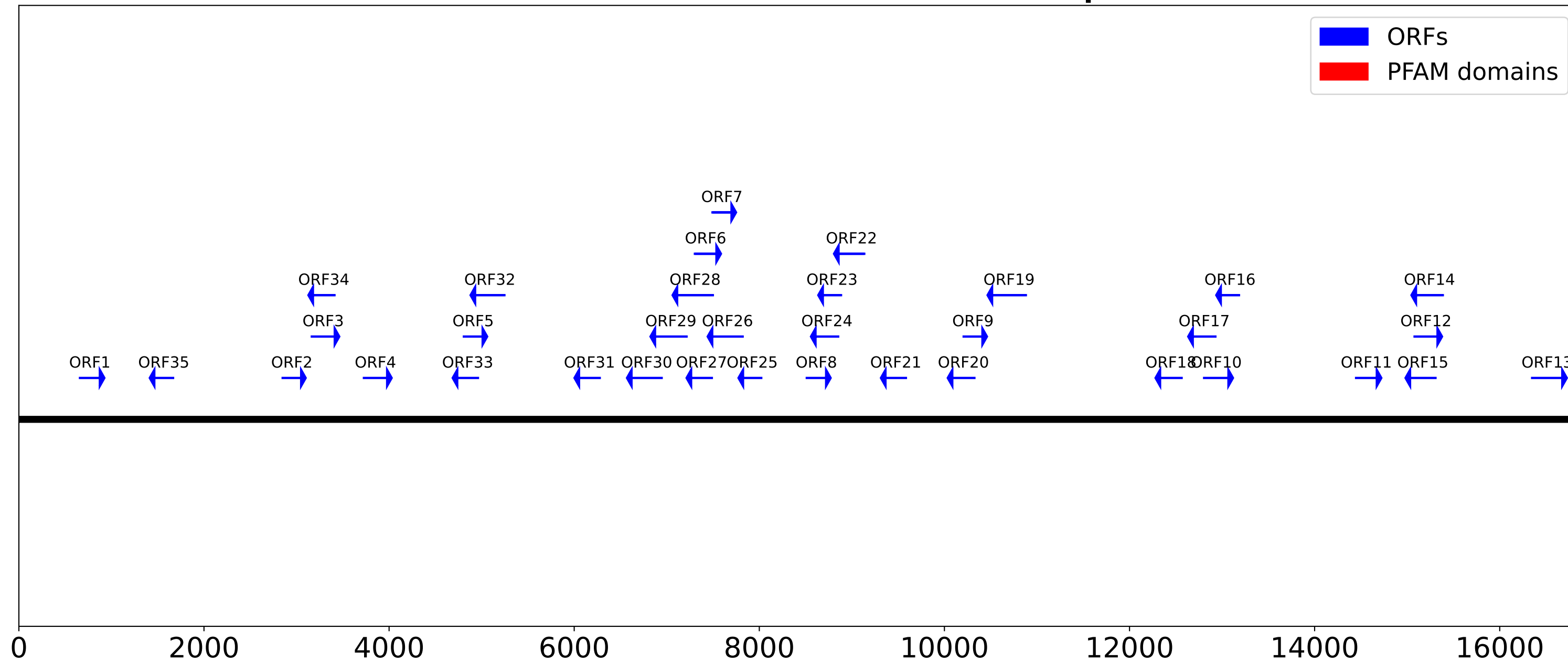


Before TEtrimmer 255 bp

TE: rnd_5_family_4610
size: 255bp; fragments: 187; full length: 81 (≥ 229.5 bp)



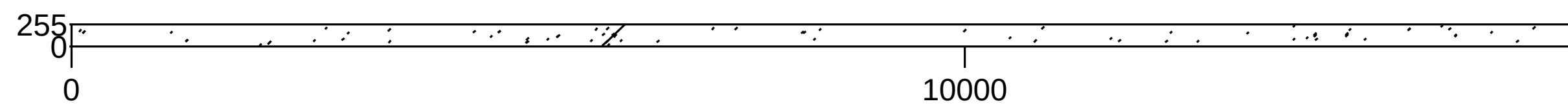
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)