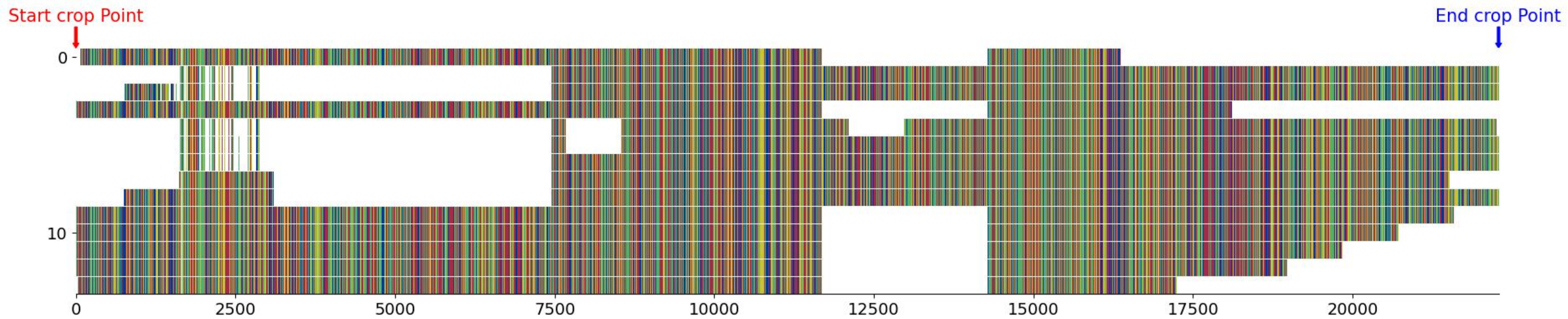


Start crop Point

End crop Point

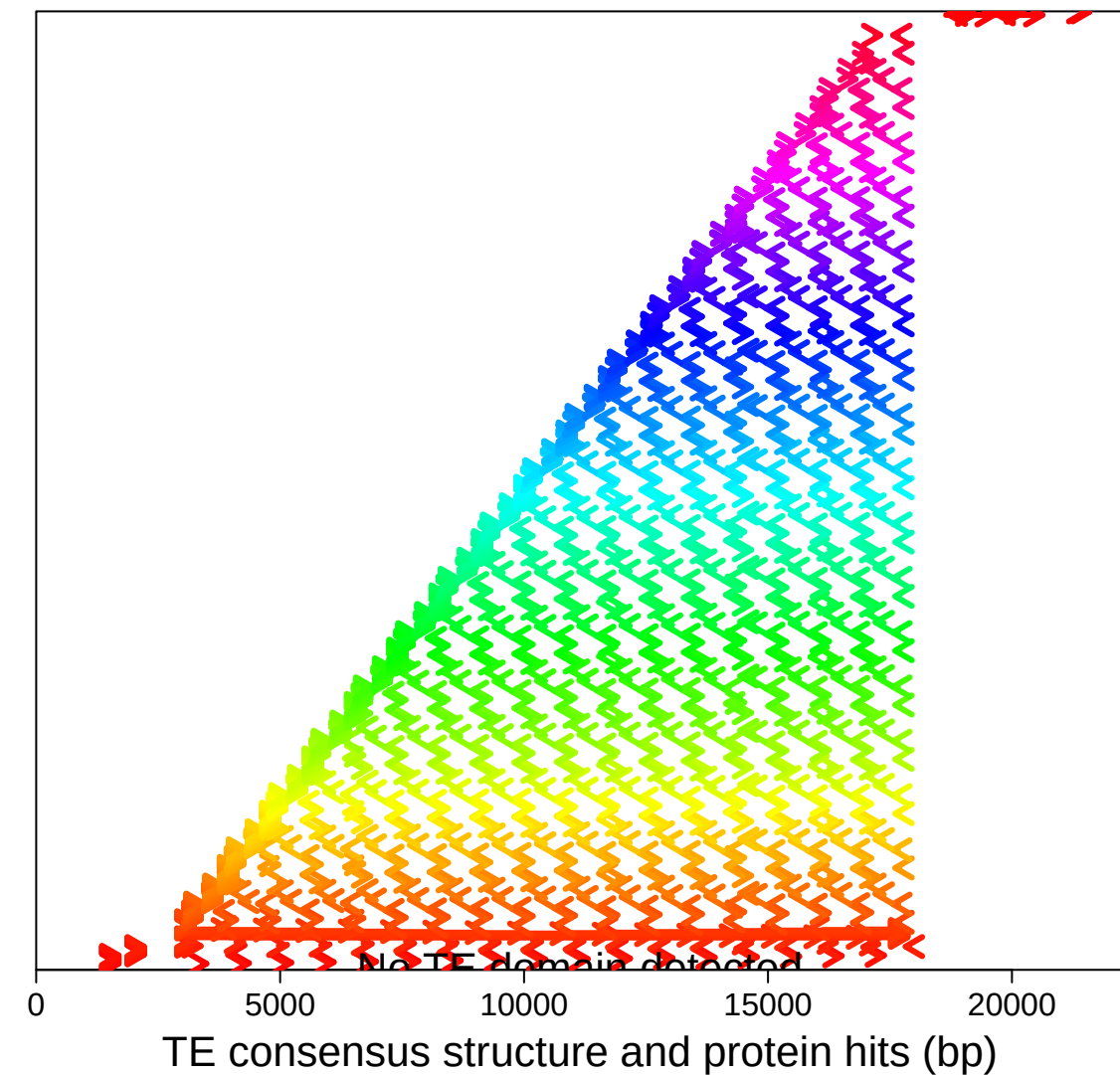
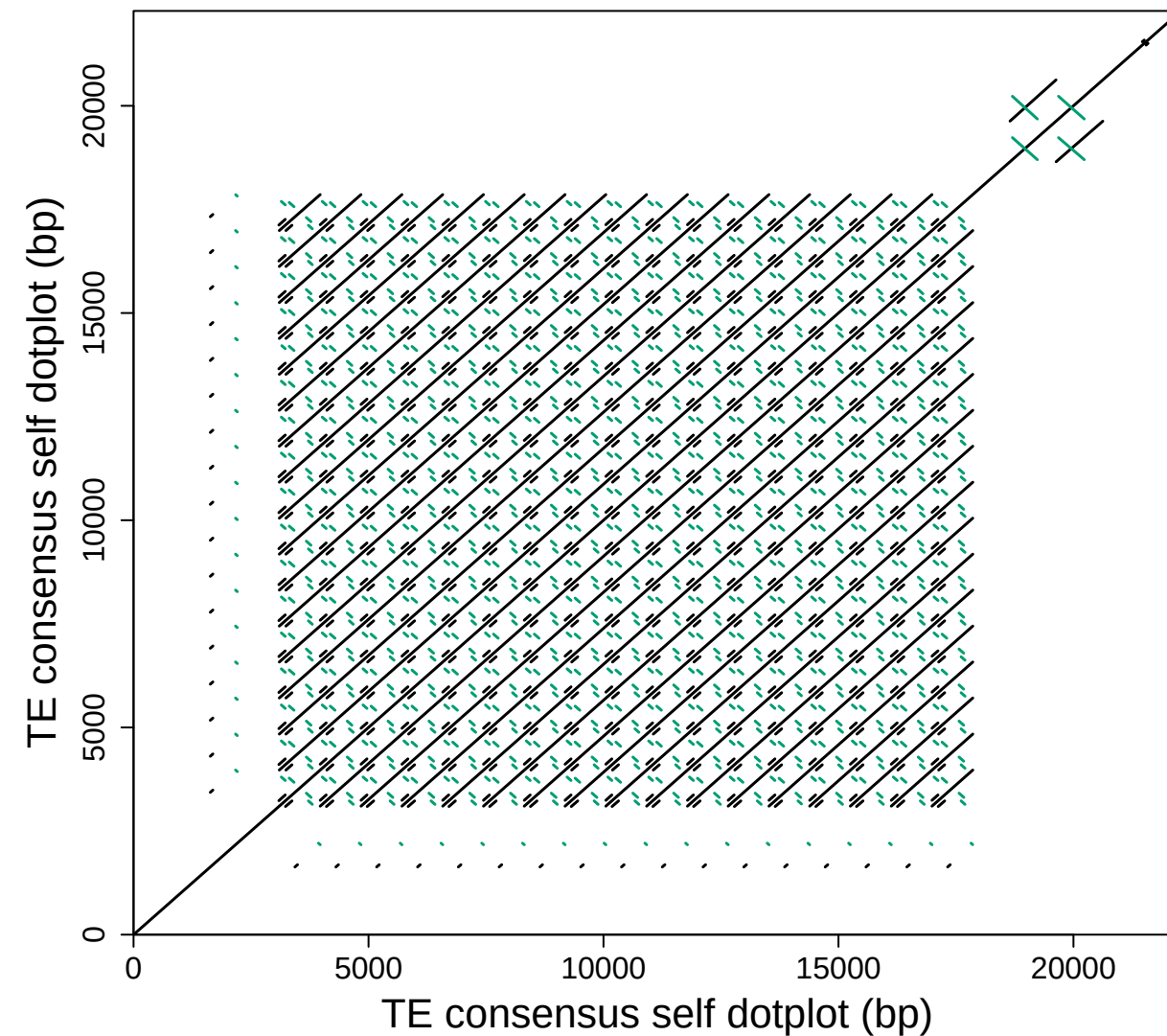
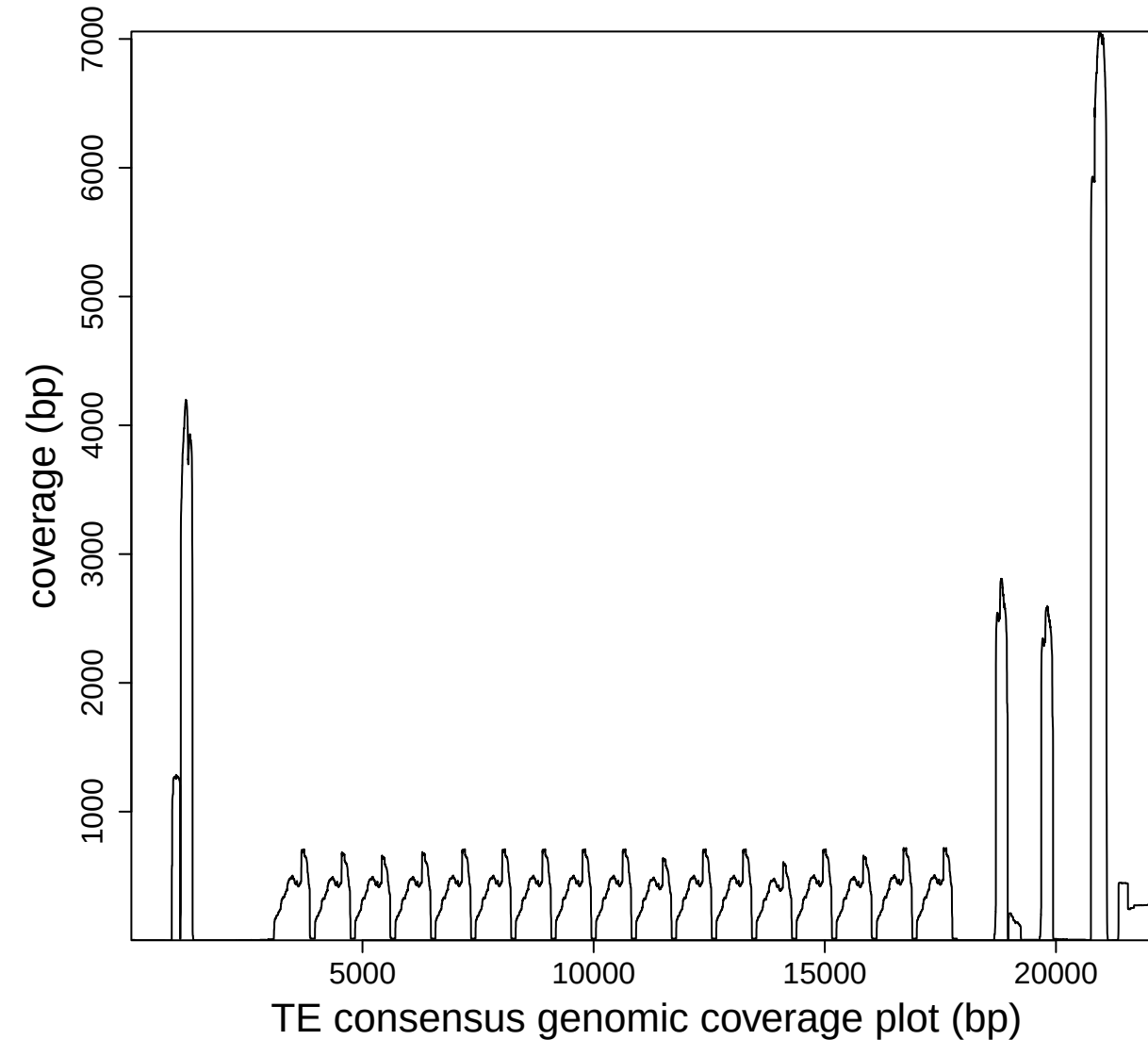
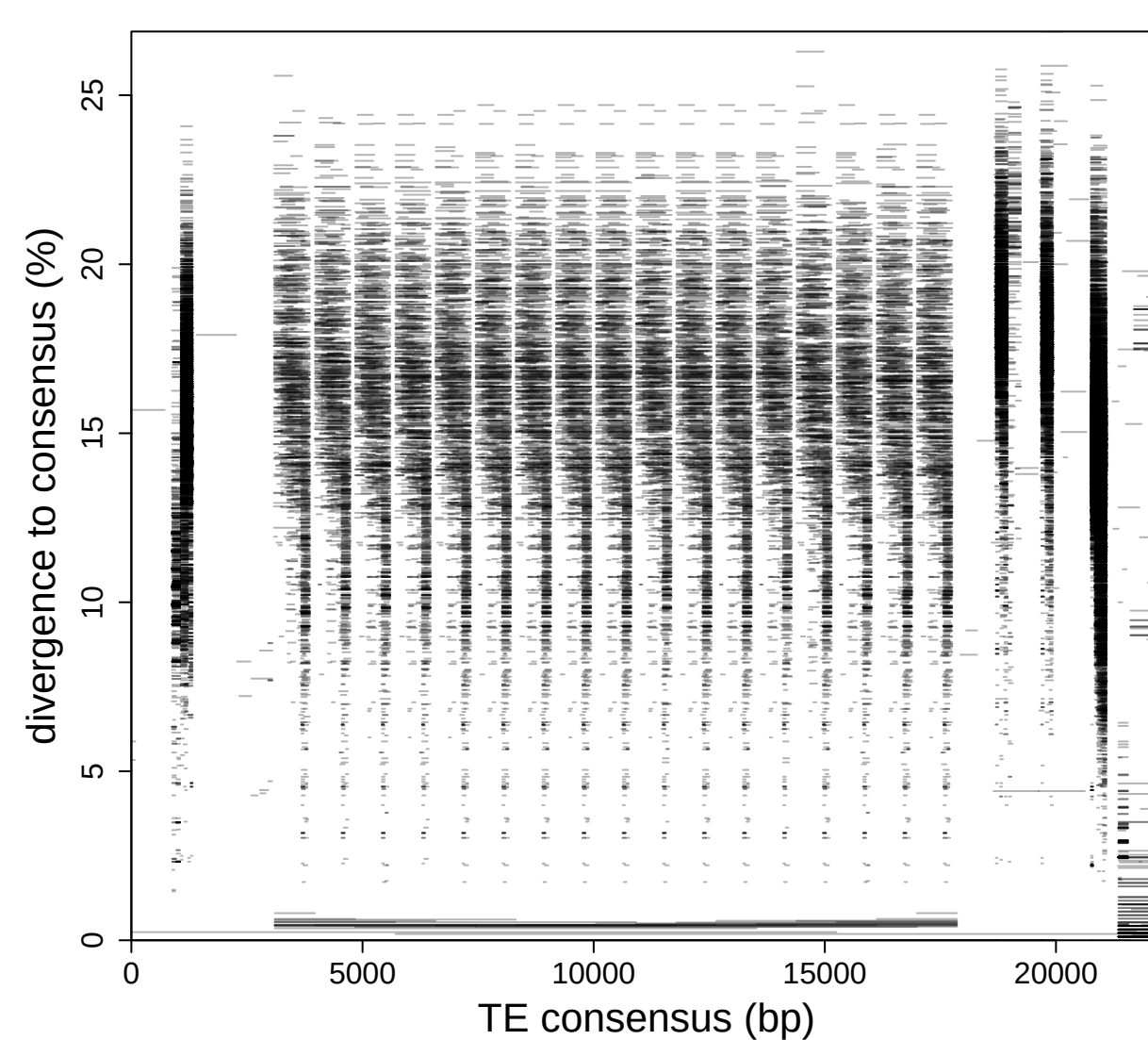
MSA length = 22289



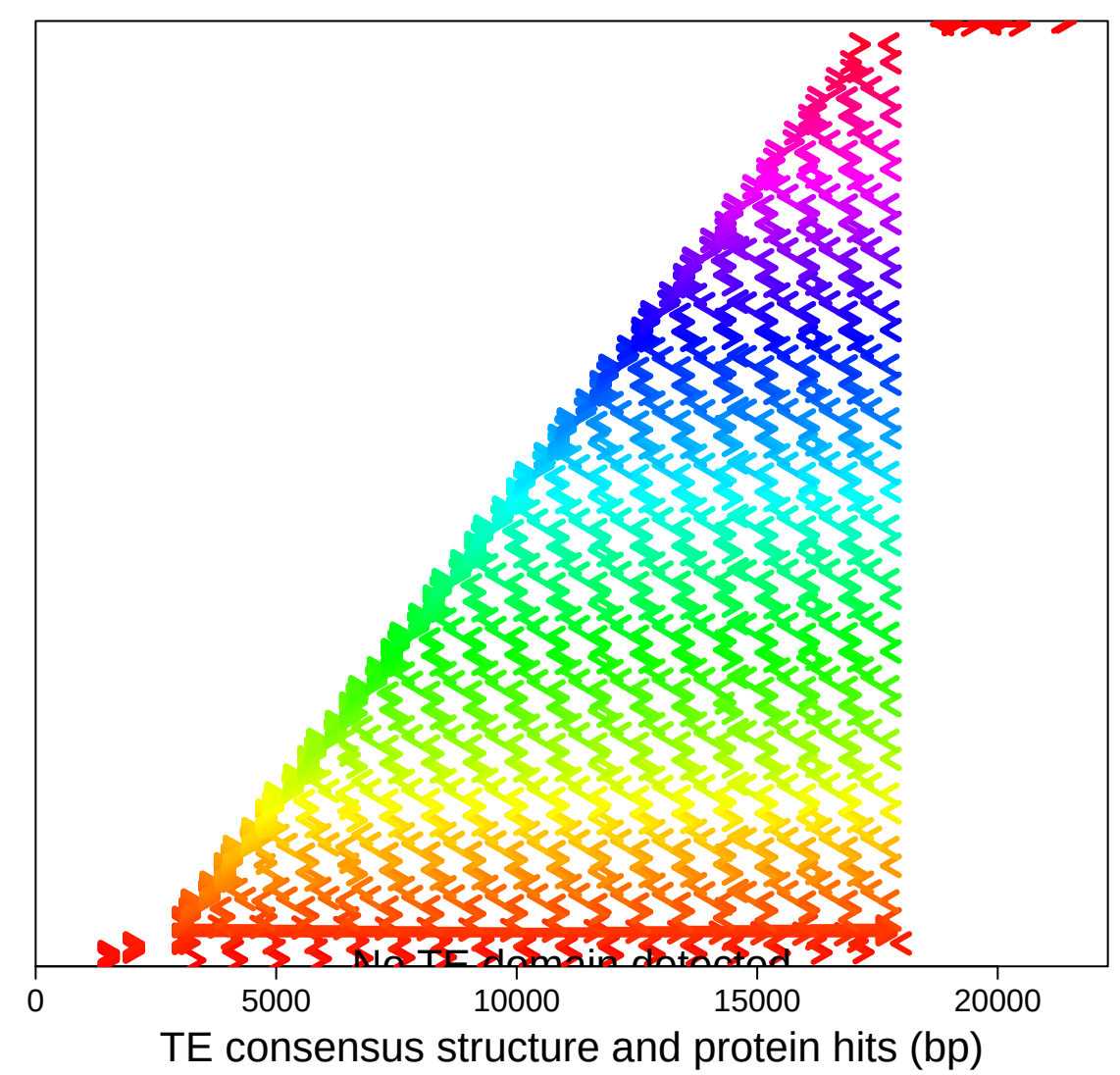
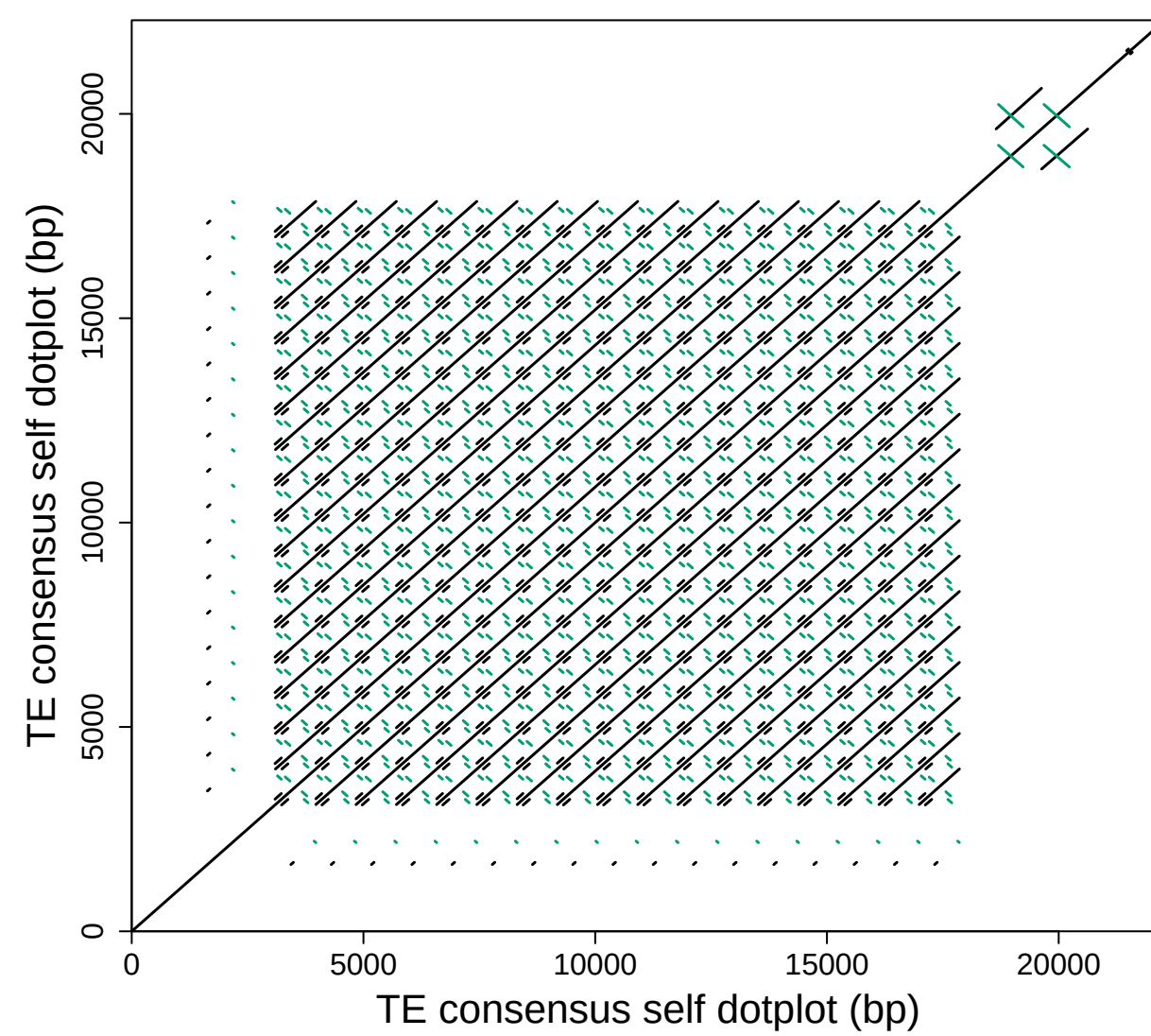
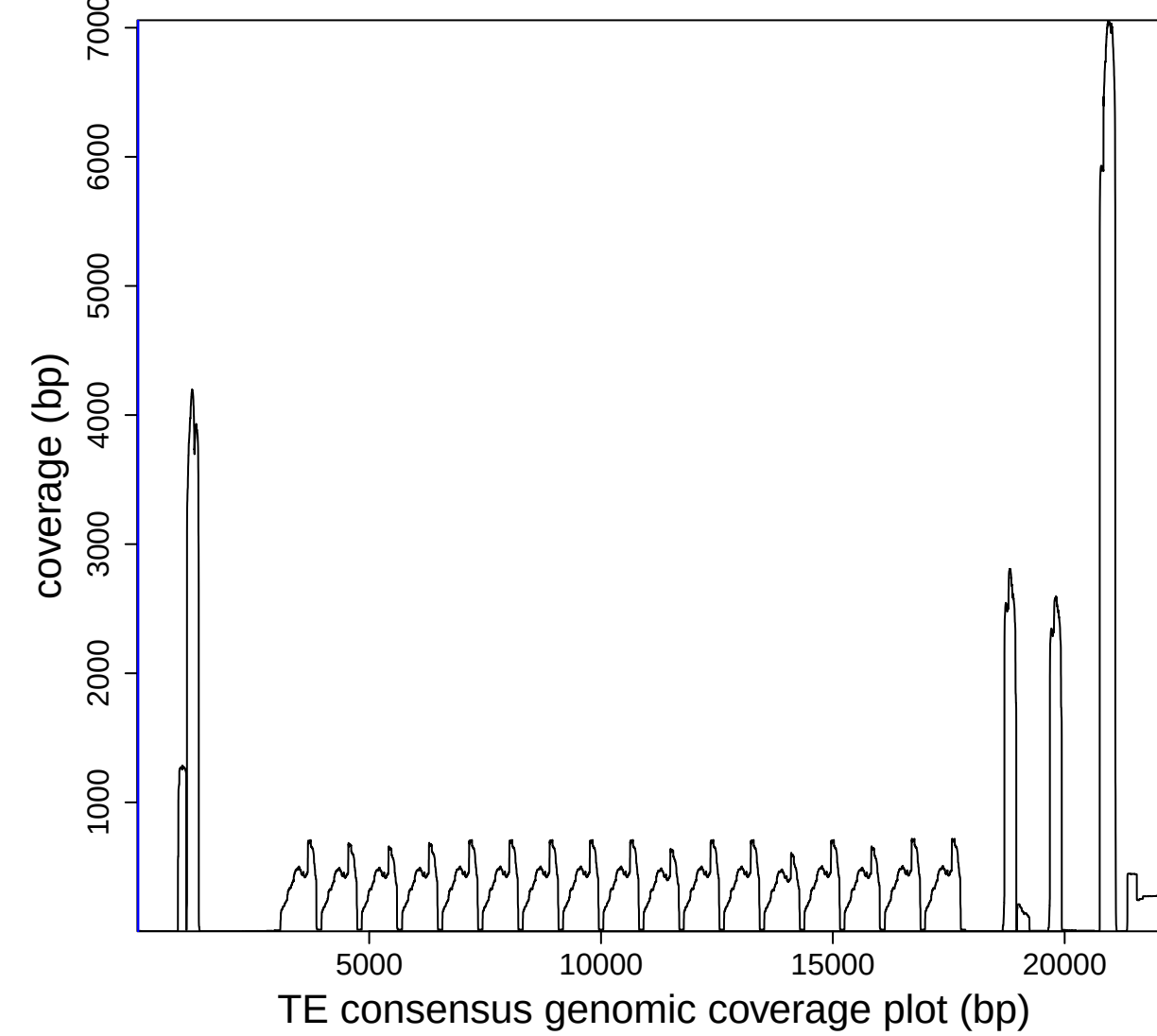
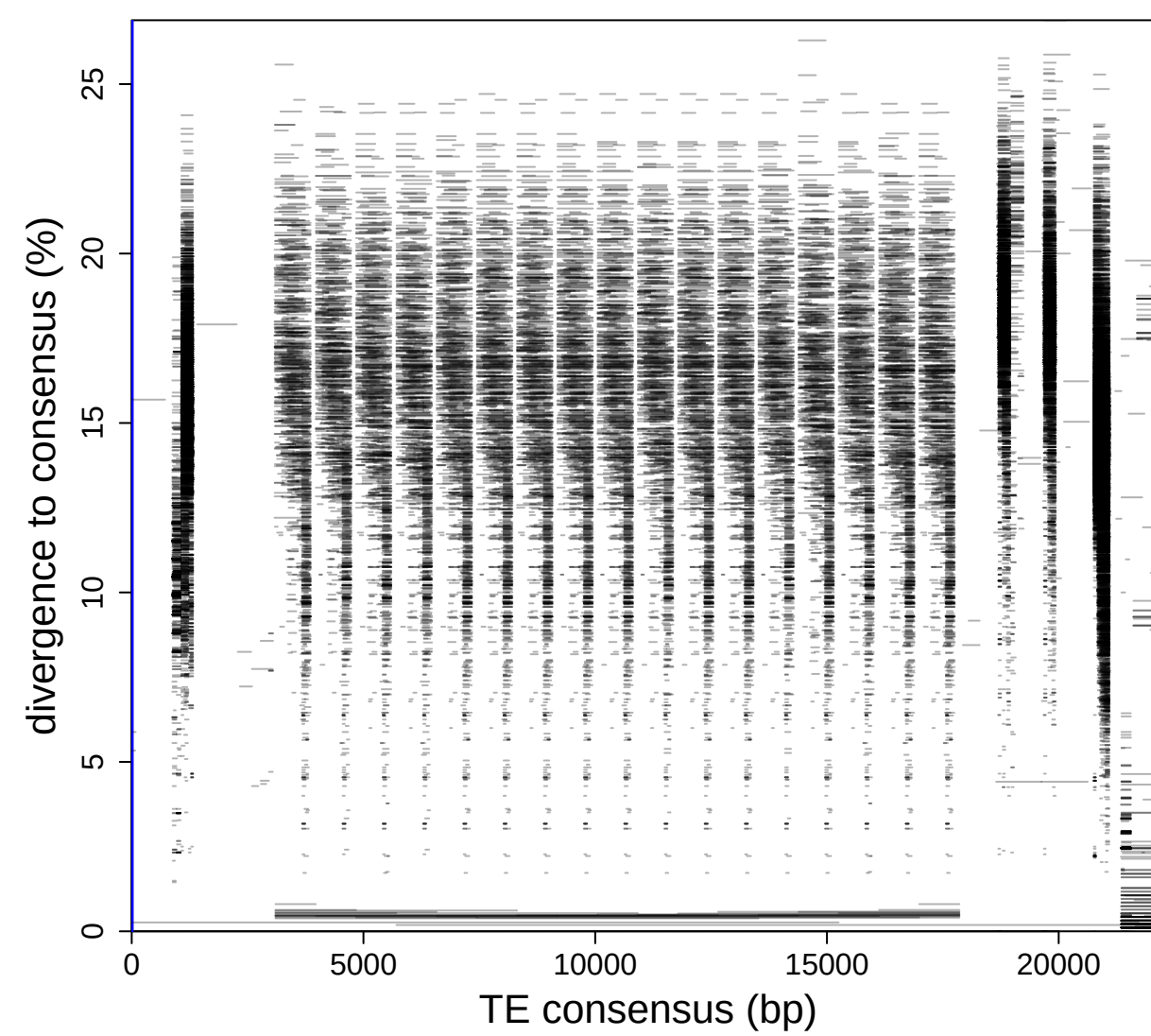


g_1.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcIn.fa a
size: 22289bp; fragments: 37403; full length: 0 (>=20060.1bp)

After TEtrimmer 22289 bp

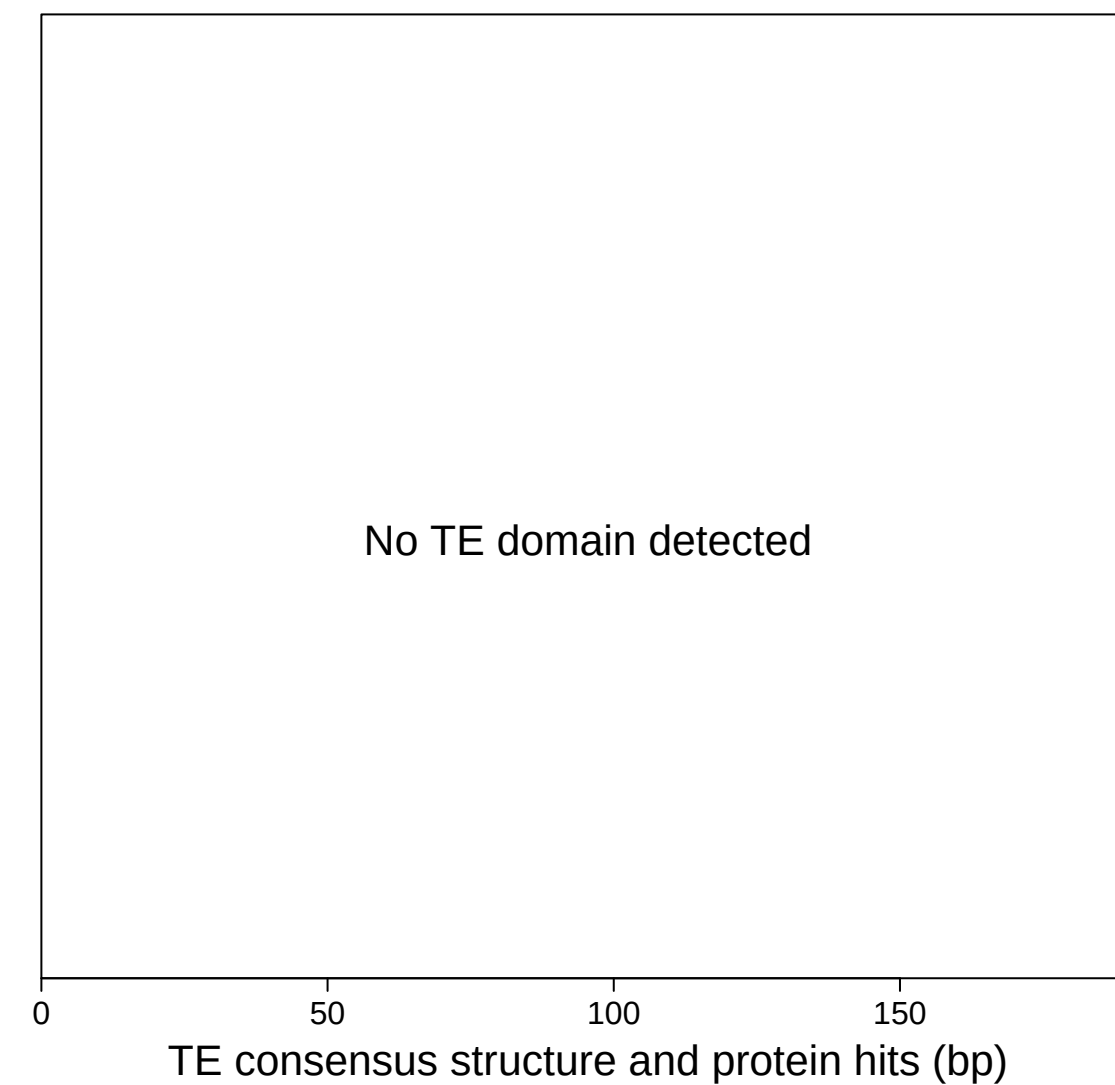
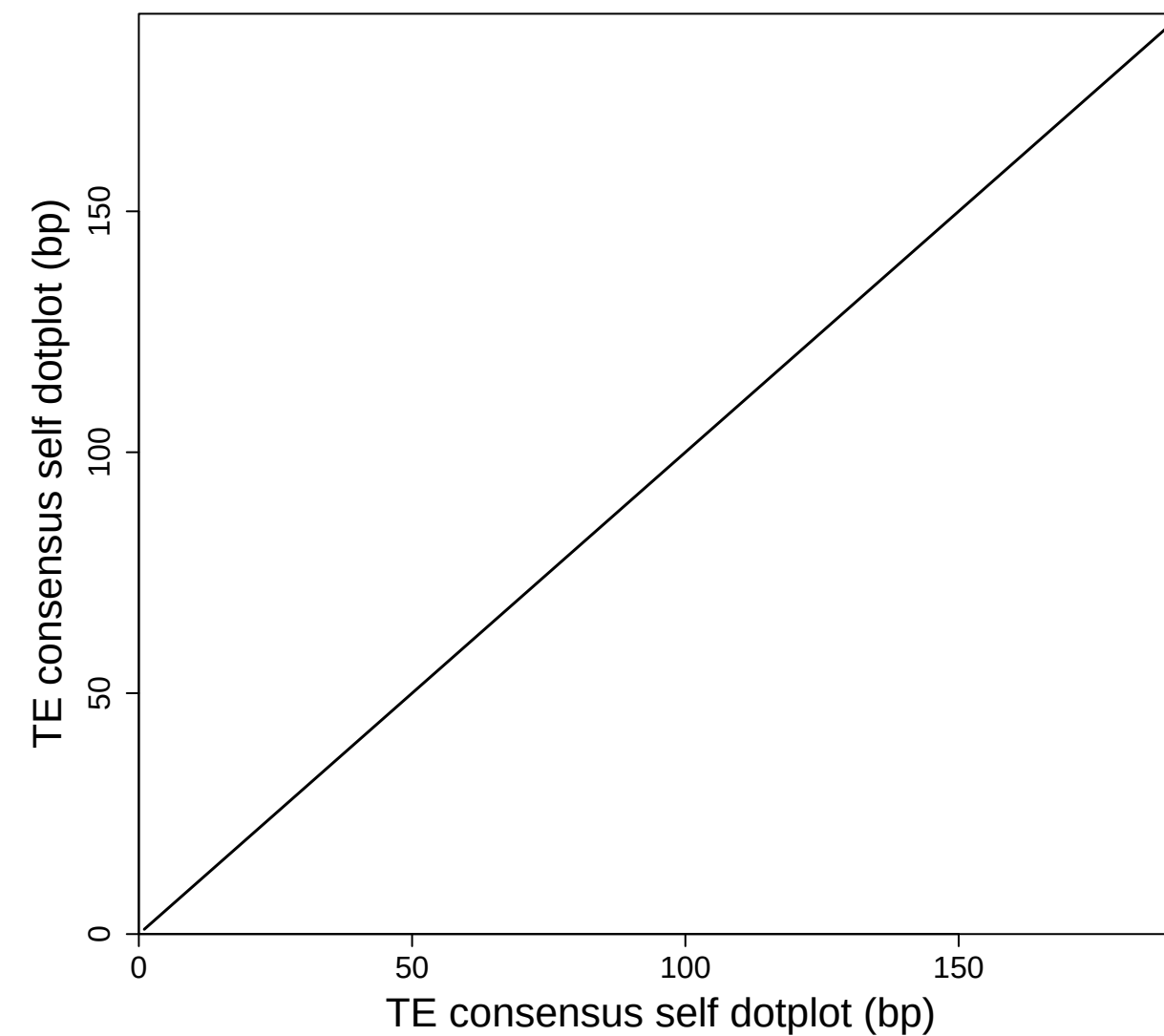
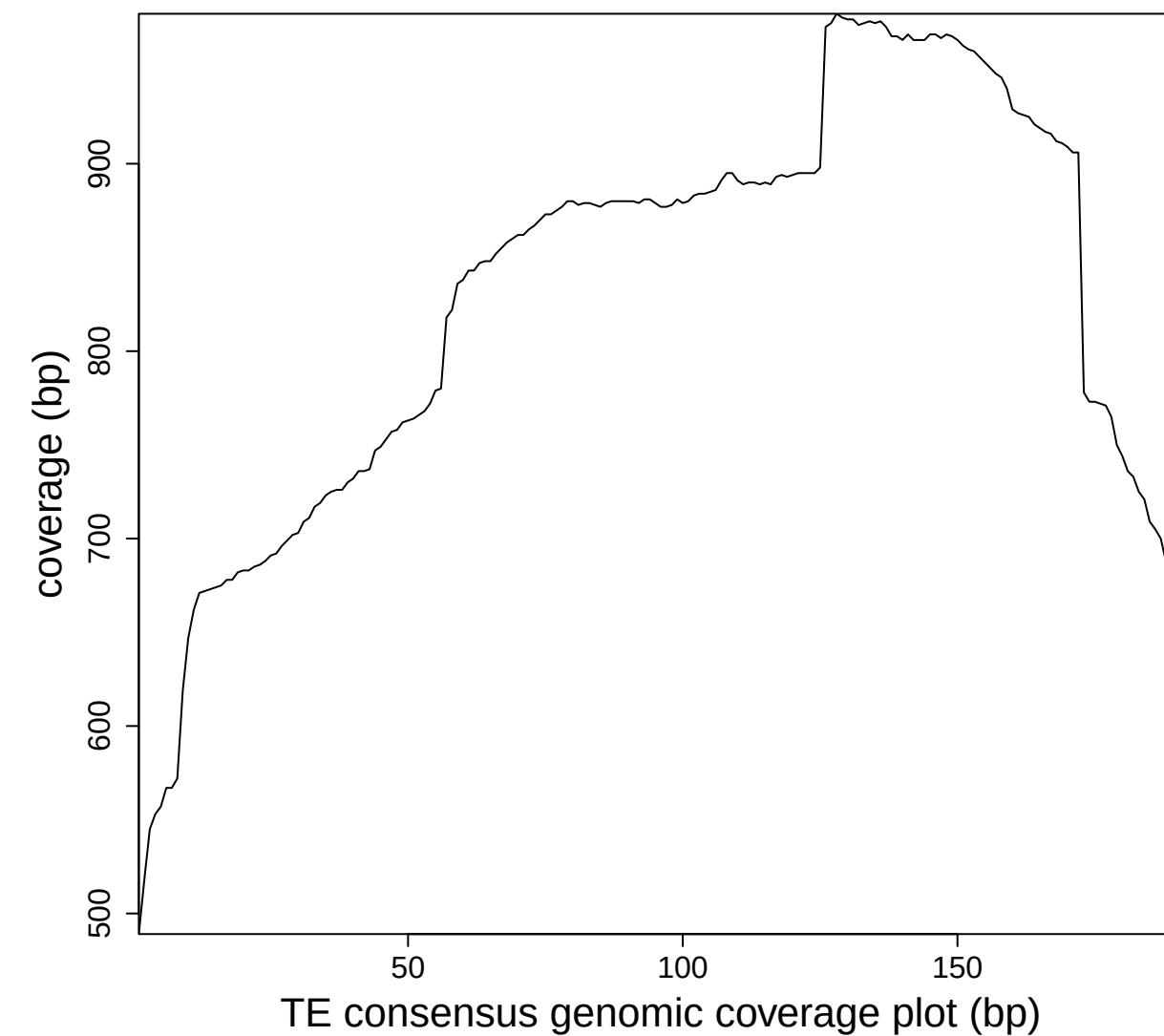
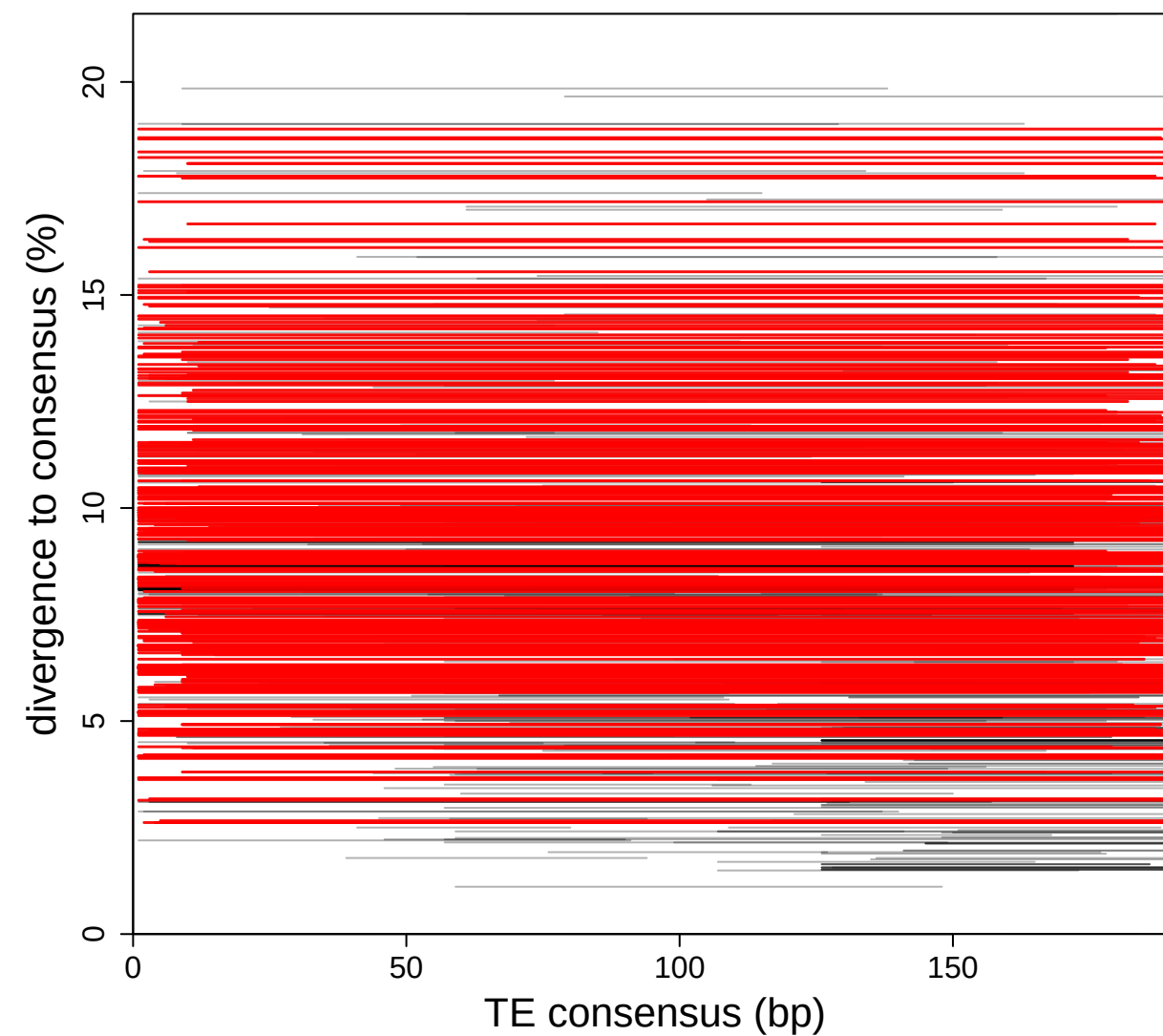


ed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.f
size: 22290bp; fragments: 37405; full length: 0 (>=20061bp)



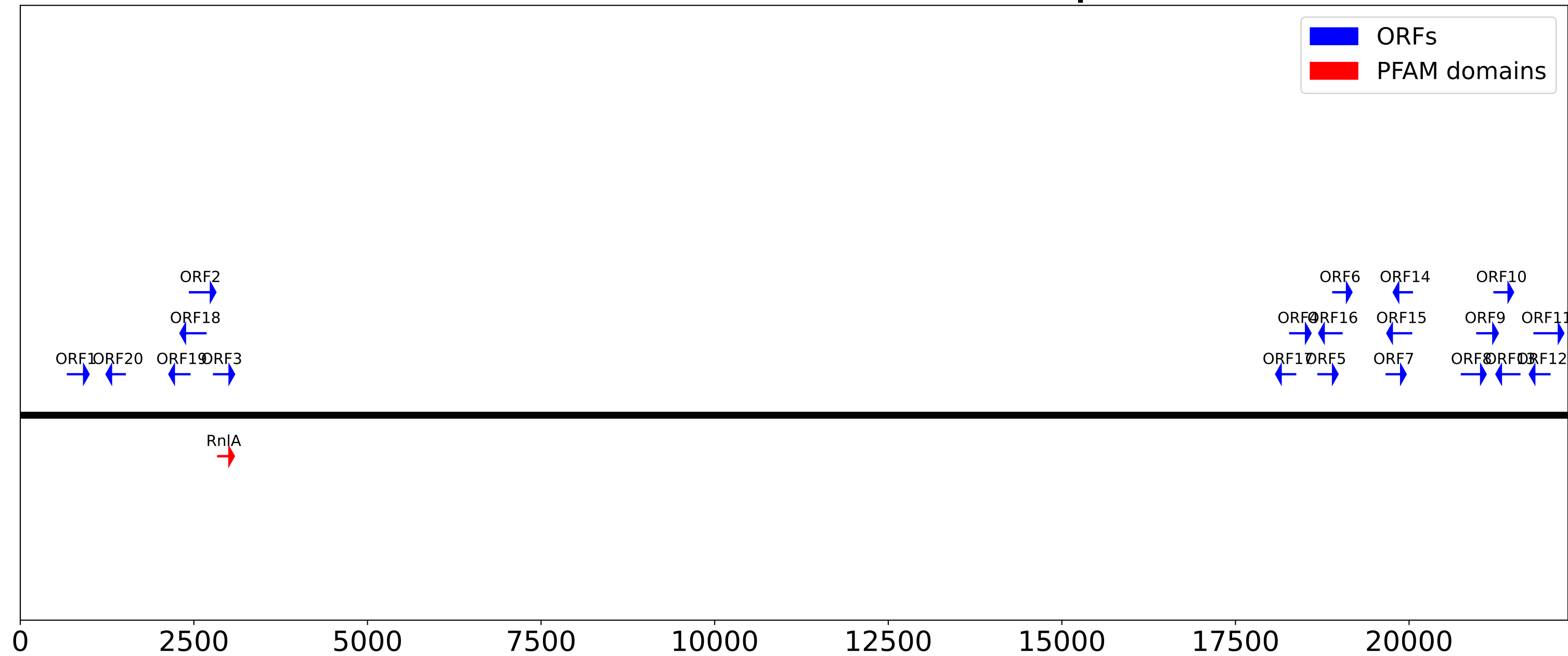
After TEtrimmer Extended plot Blue lines are boundaries

TE: rnd_4_family_1397
size: 191bp; fragments: 1049; full length: 461 (≥ 171.9 bp)



Before TEtrimmer 191 bp

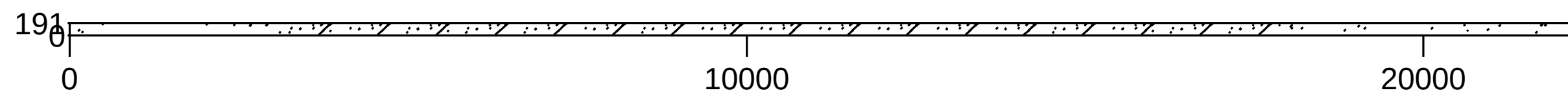
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

isus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)