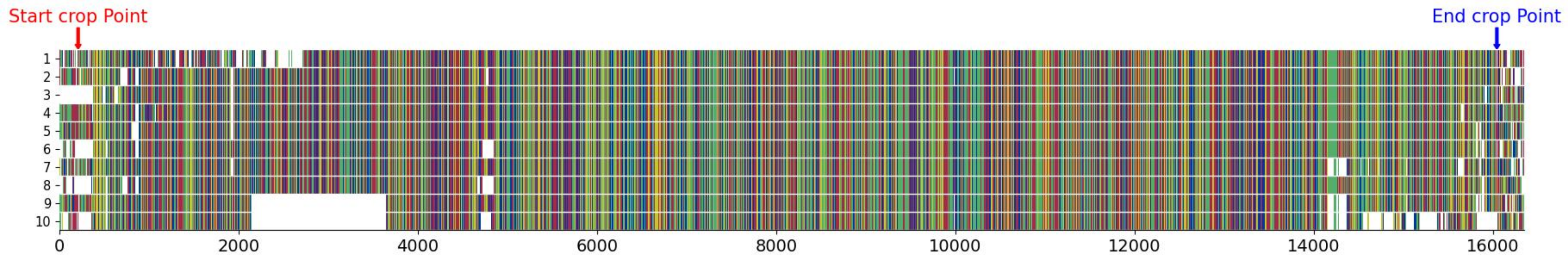


Start Crop Point

End Crop Point

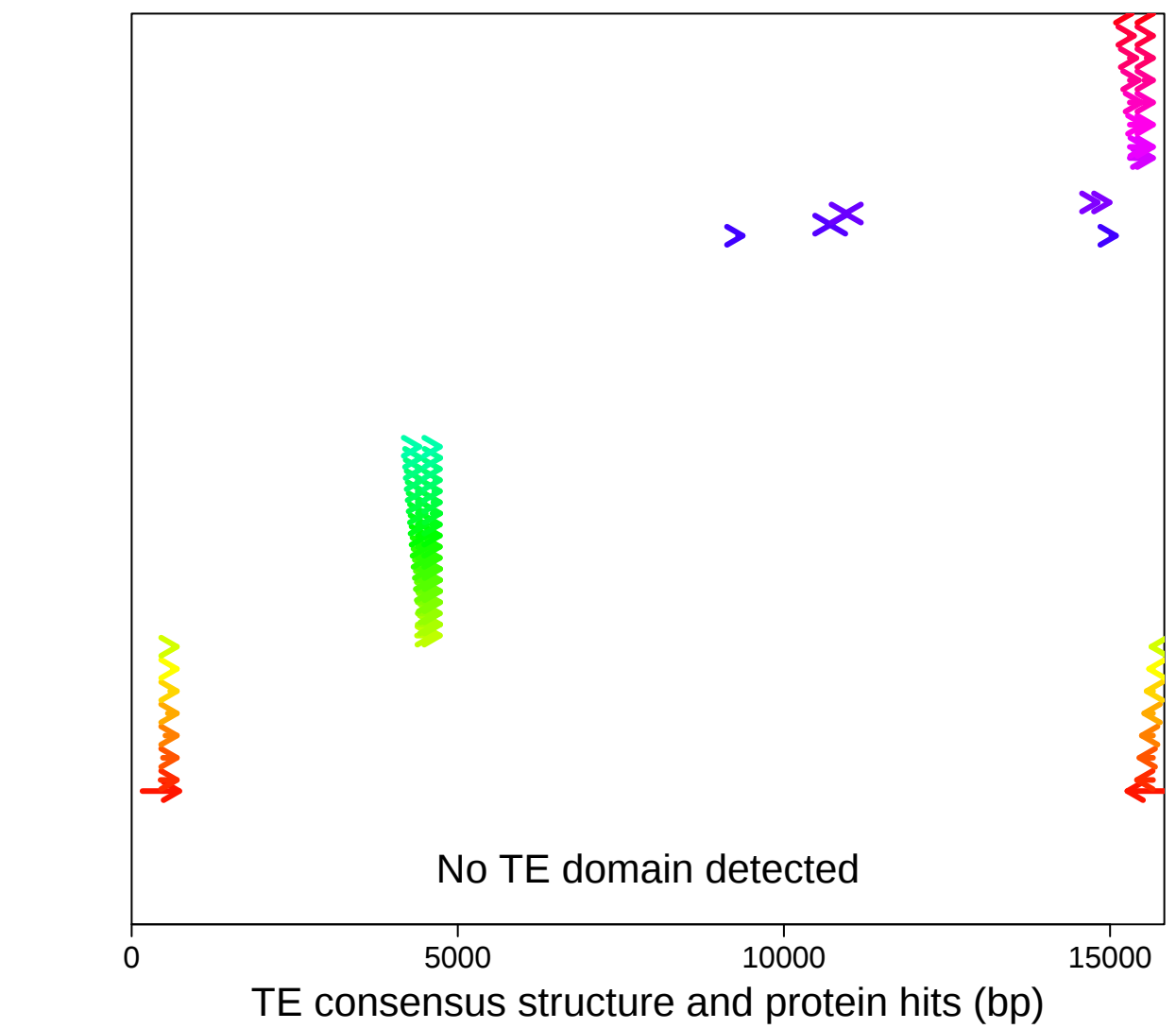
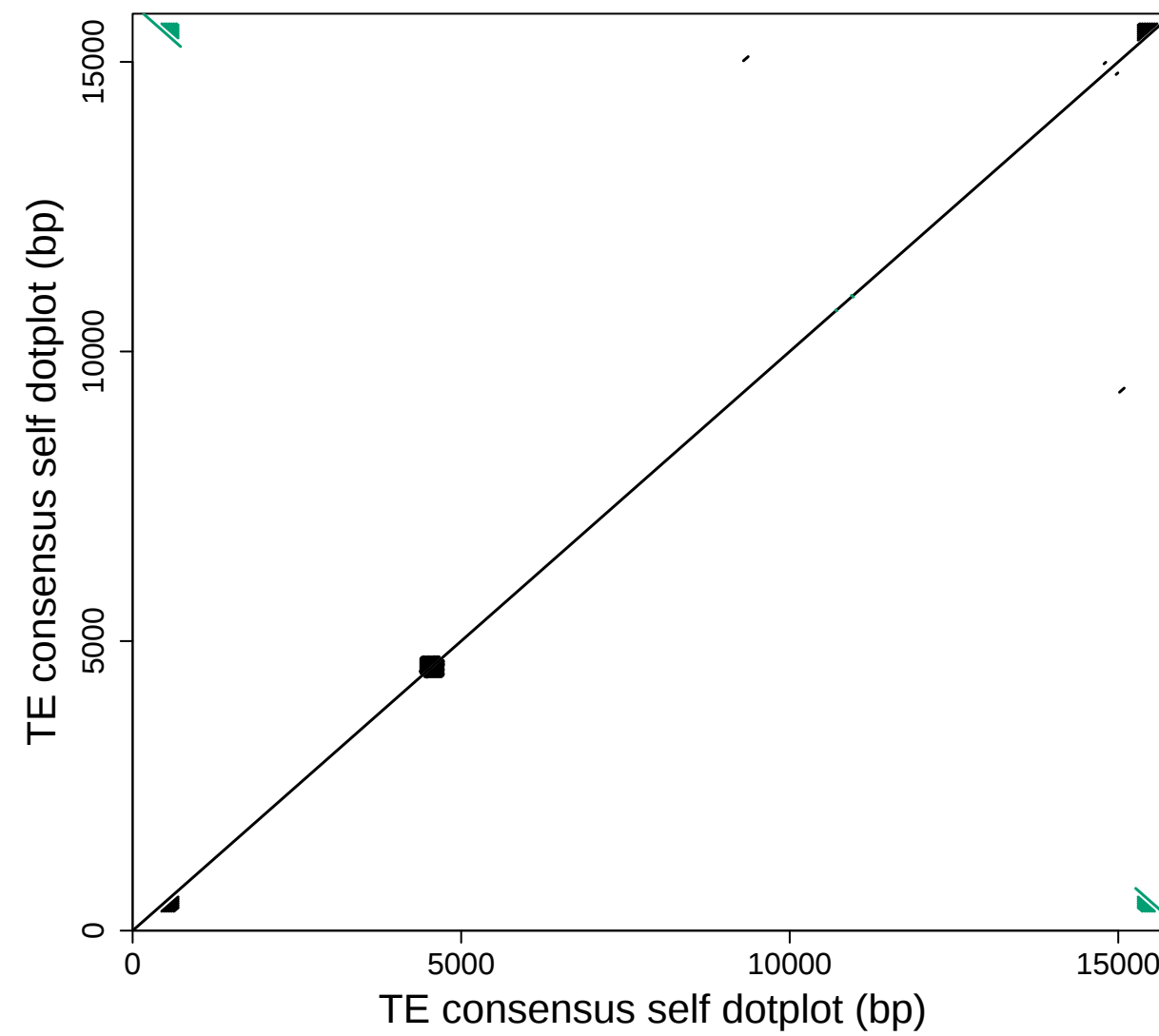
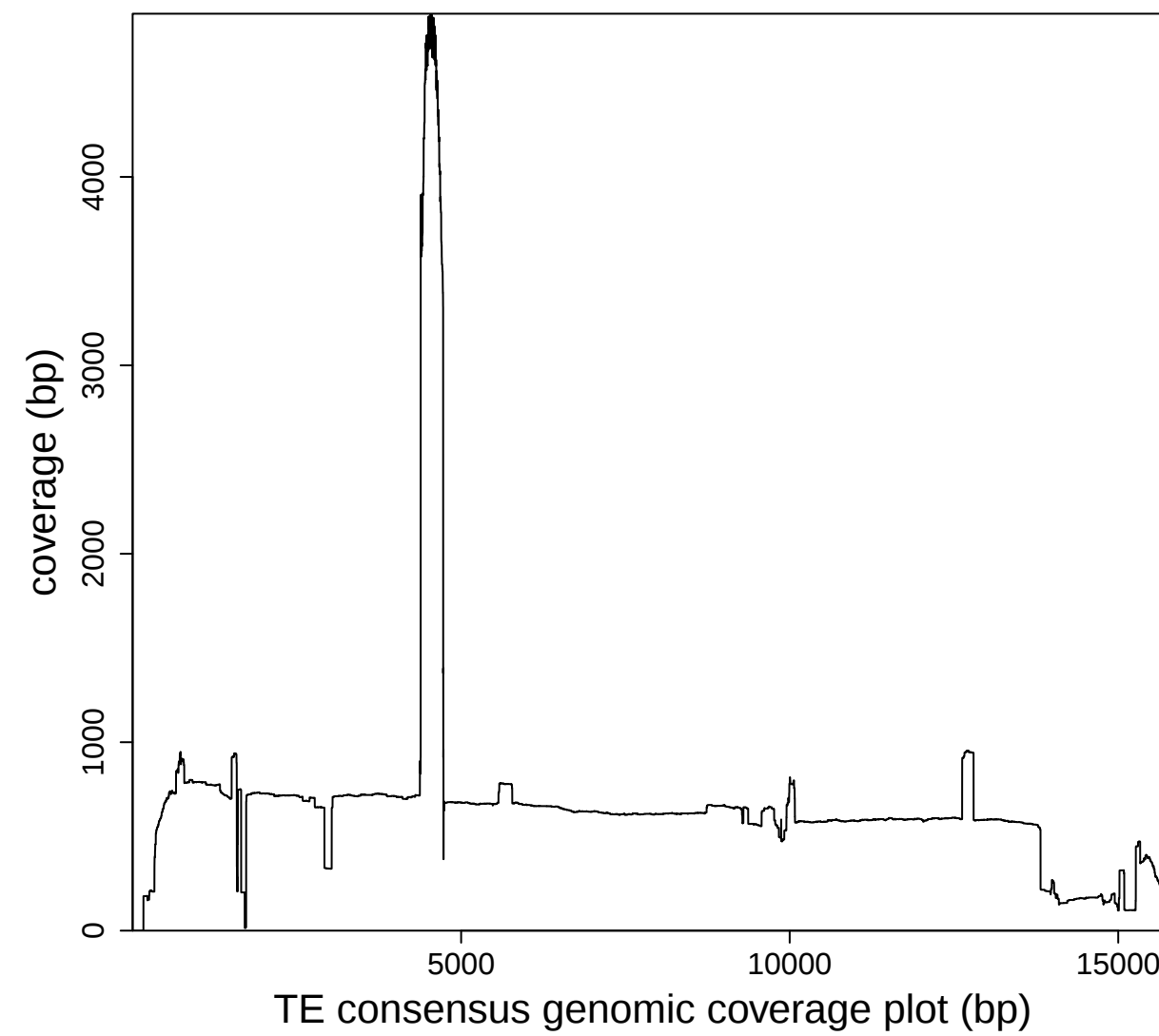
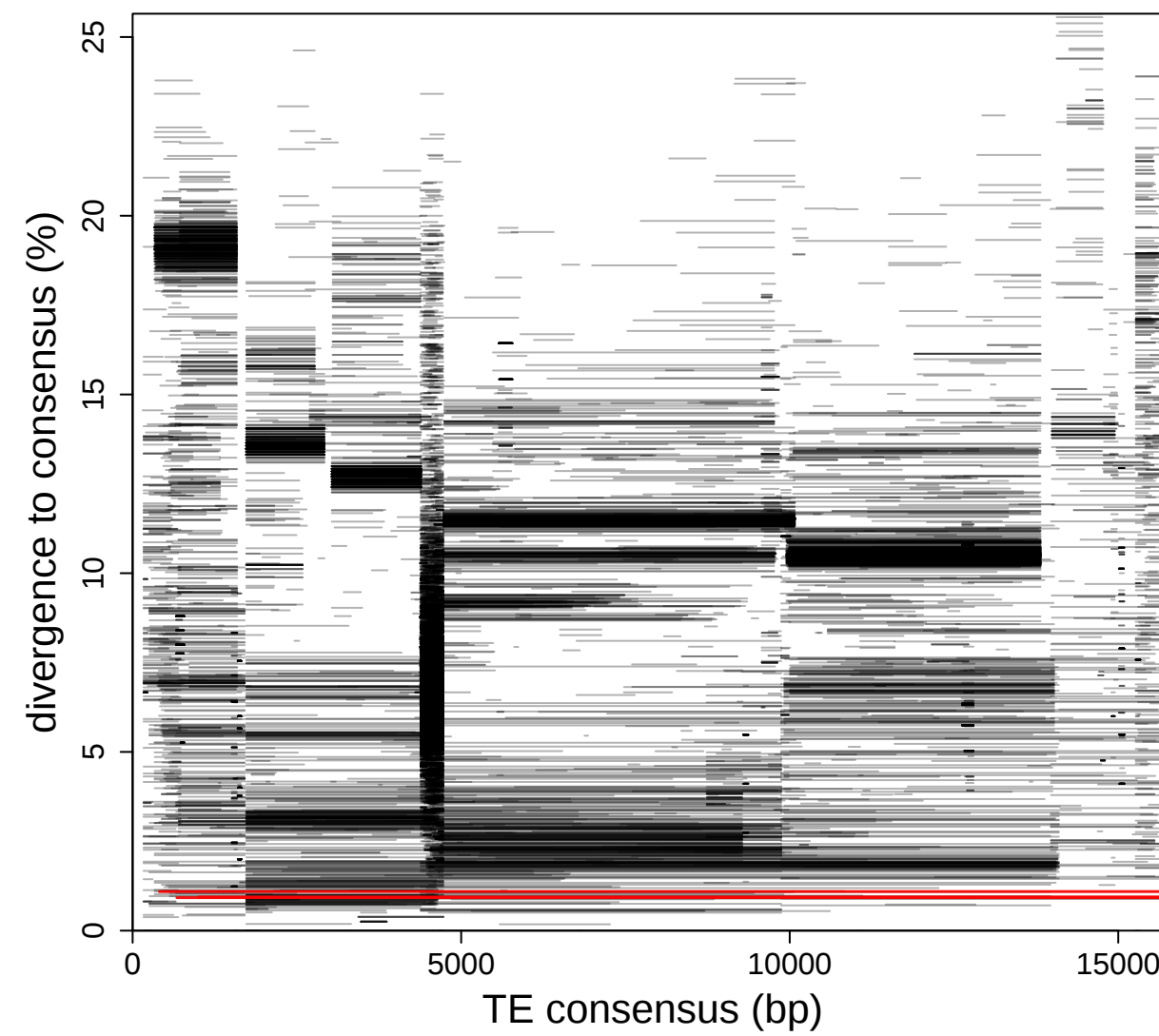
MSA length = 15832





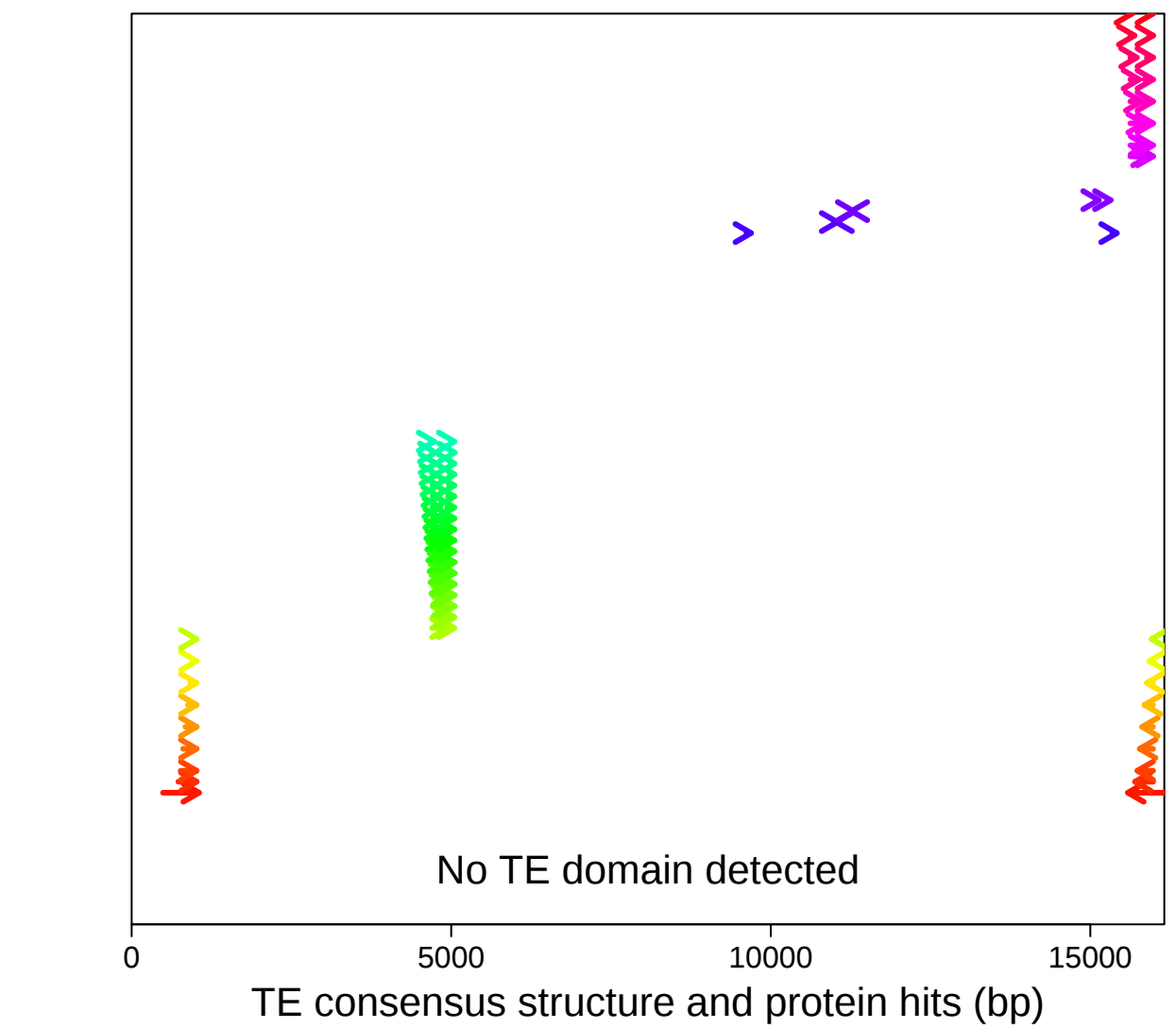
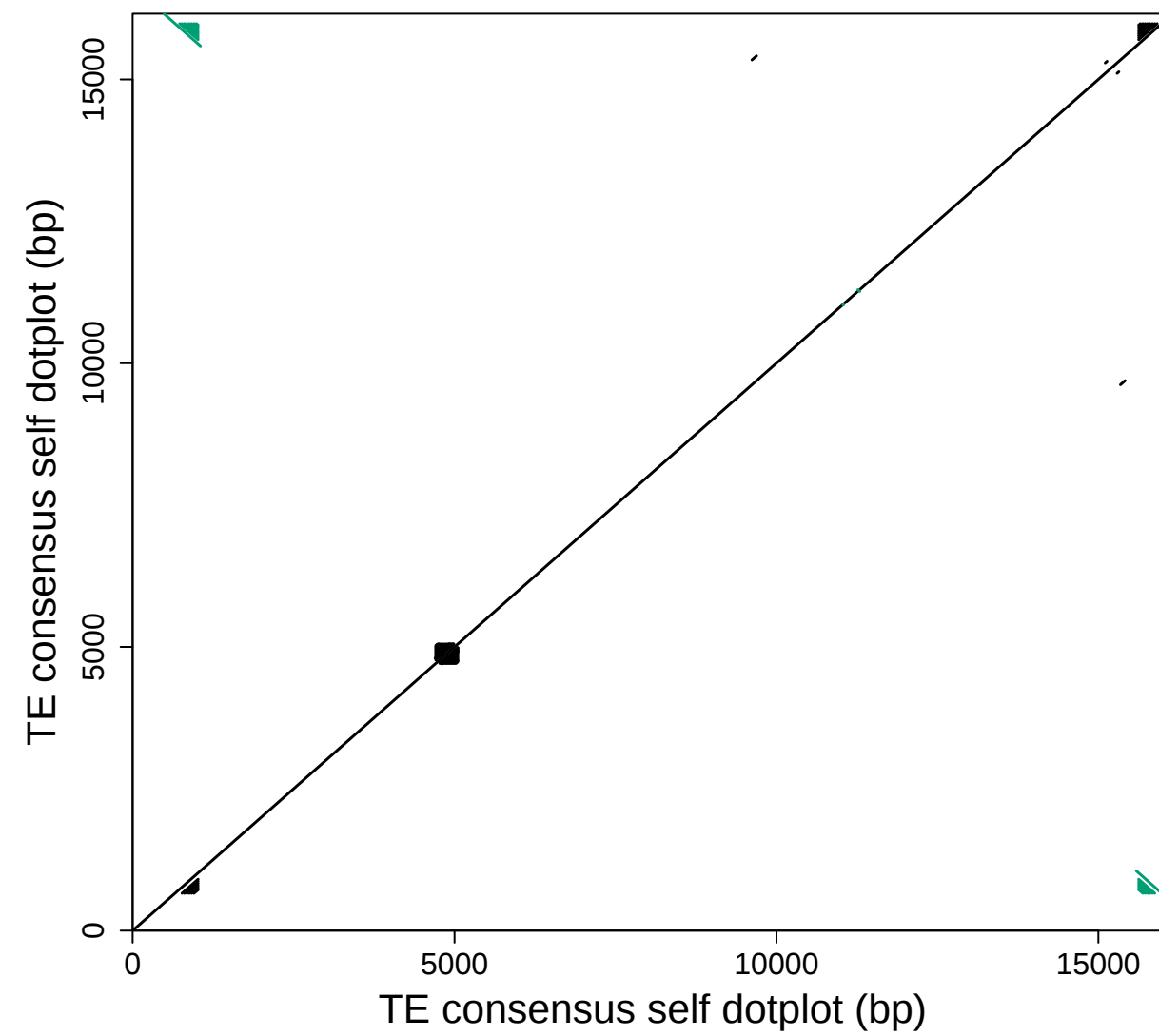
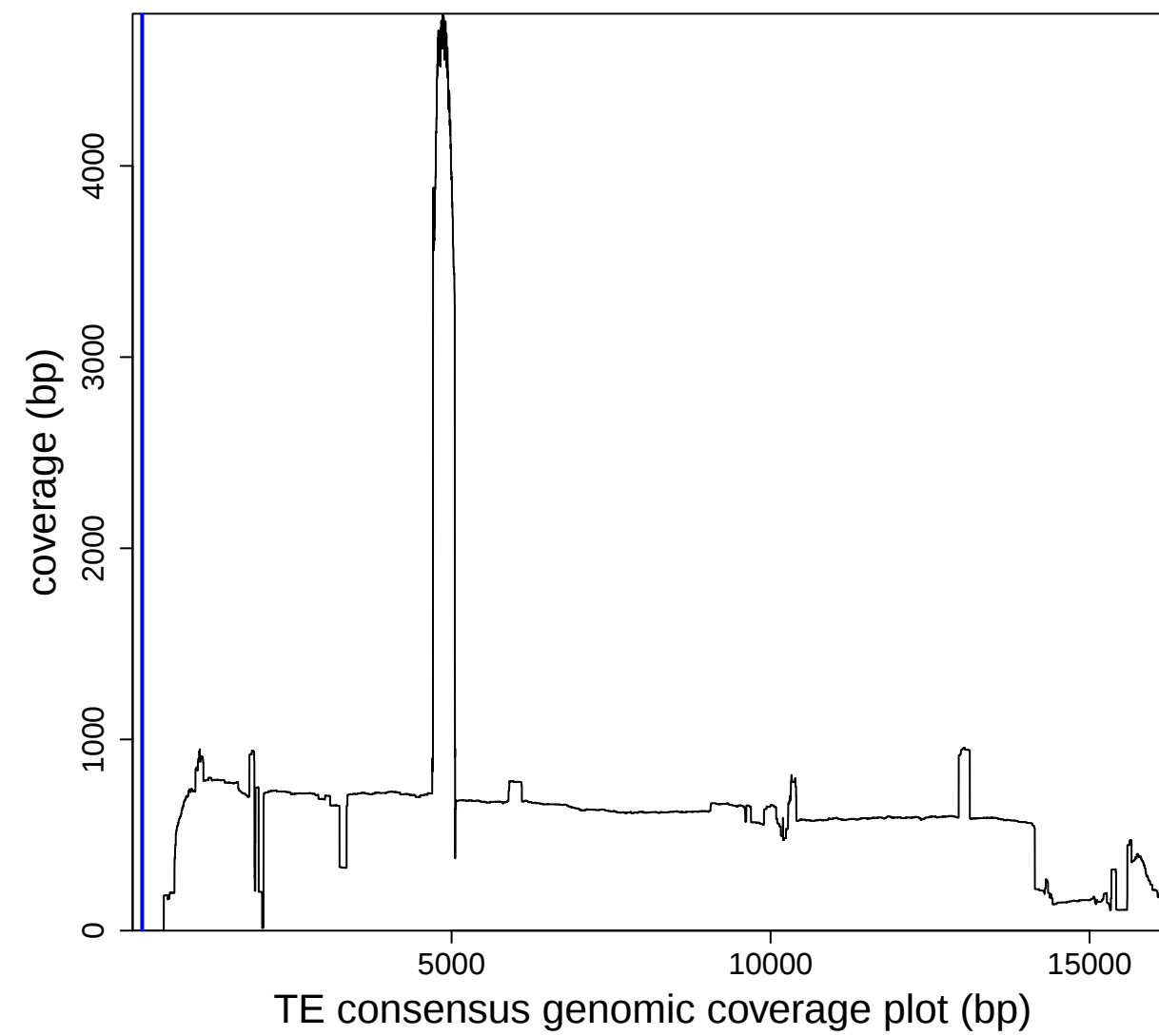
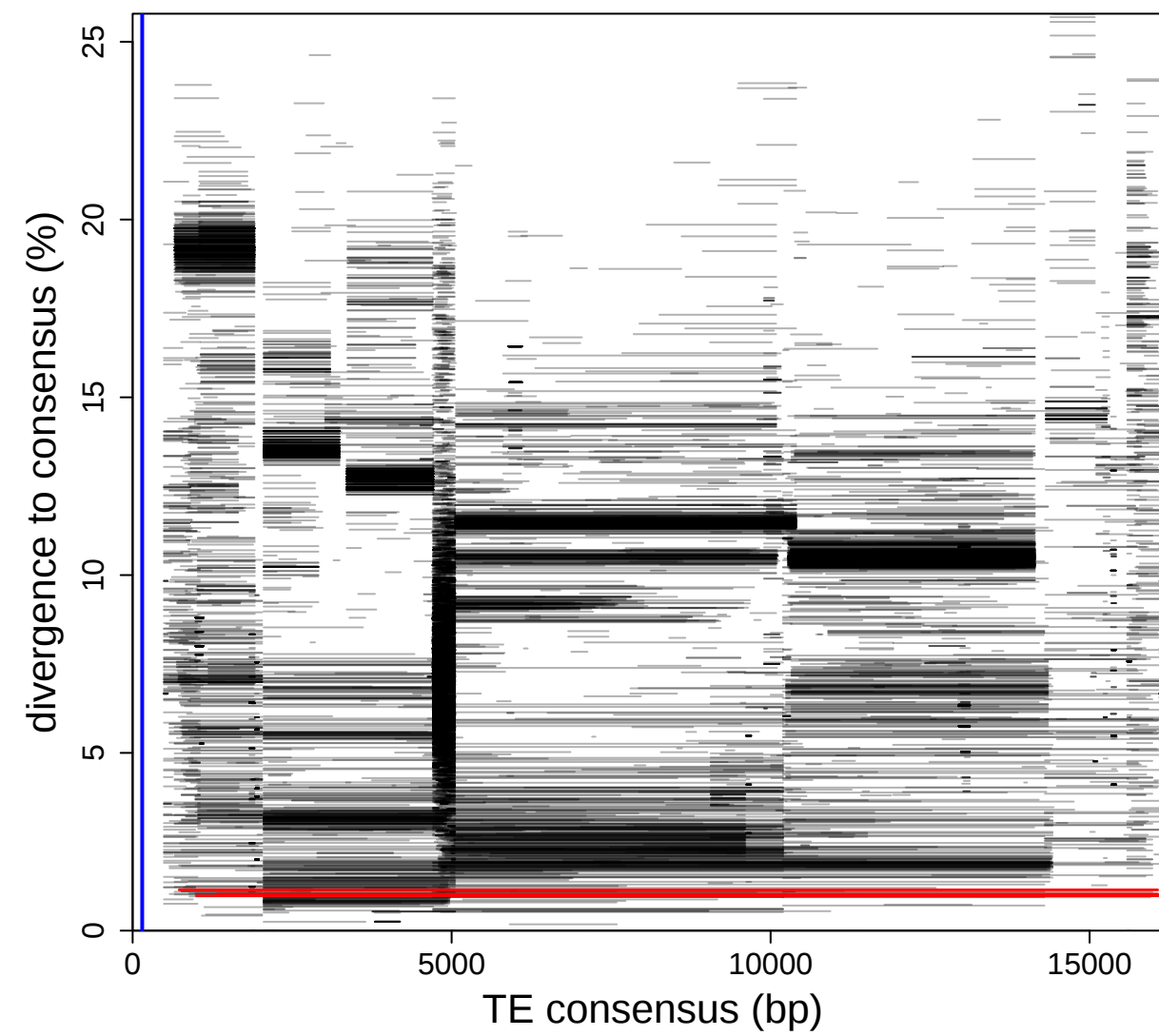
g_3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 15832bp; fragments: 14047; full length: 3 (>=14248.8bp)

After TEtrimmer 15832 bp



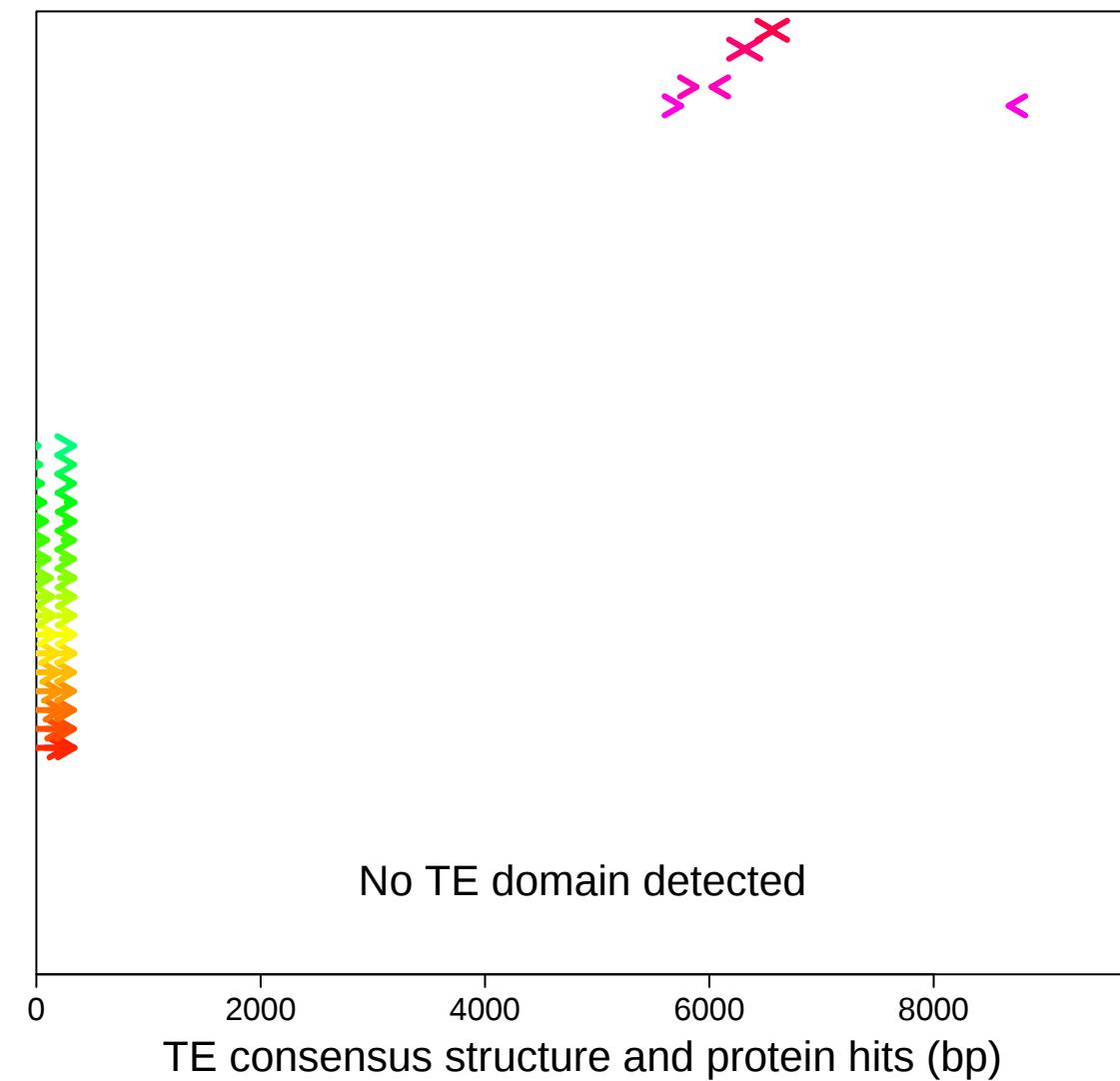
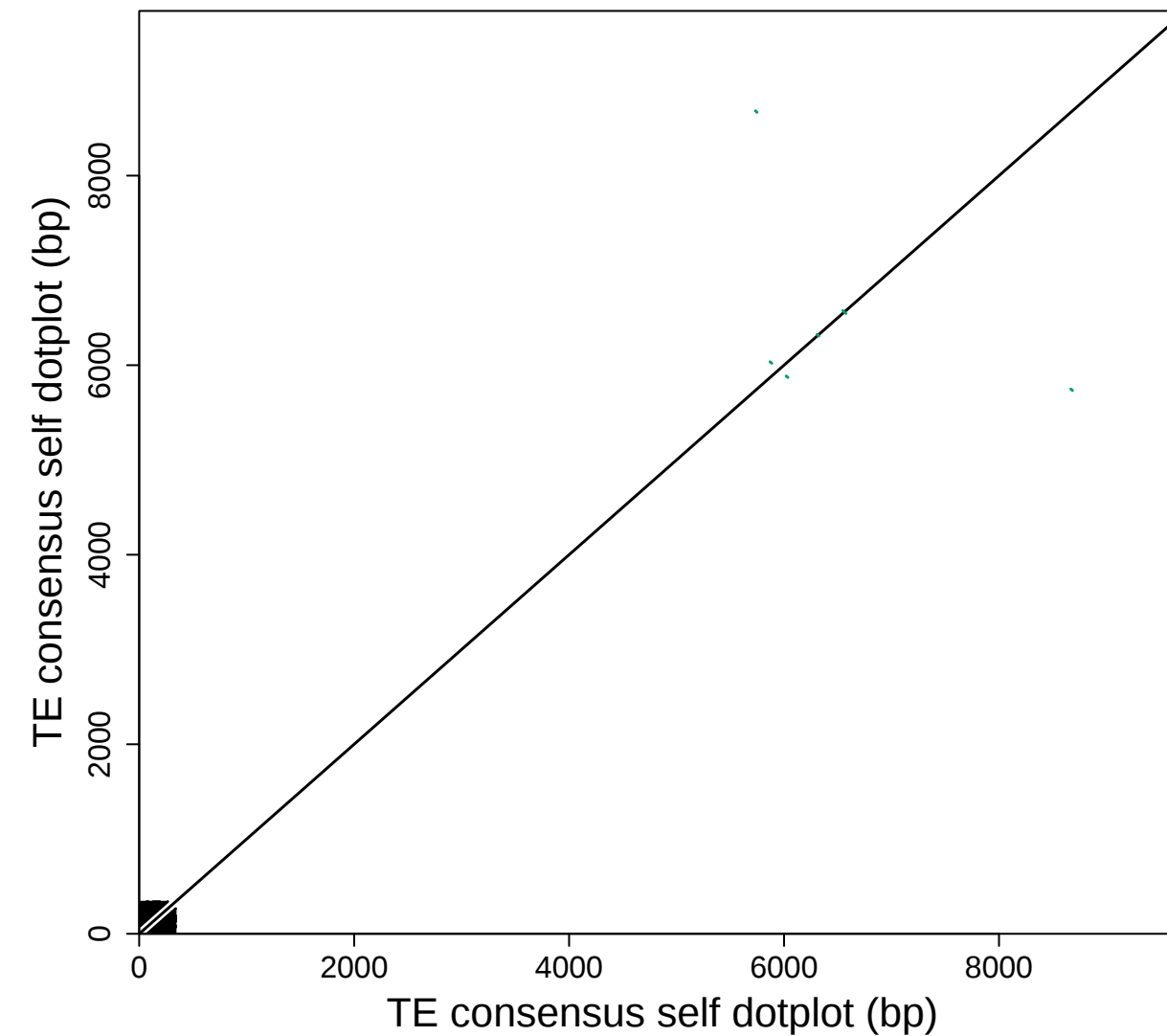
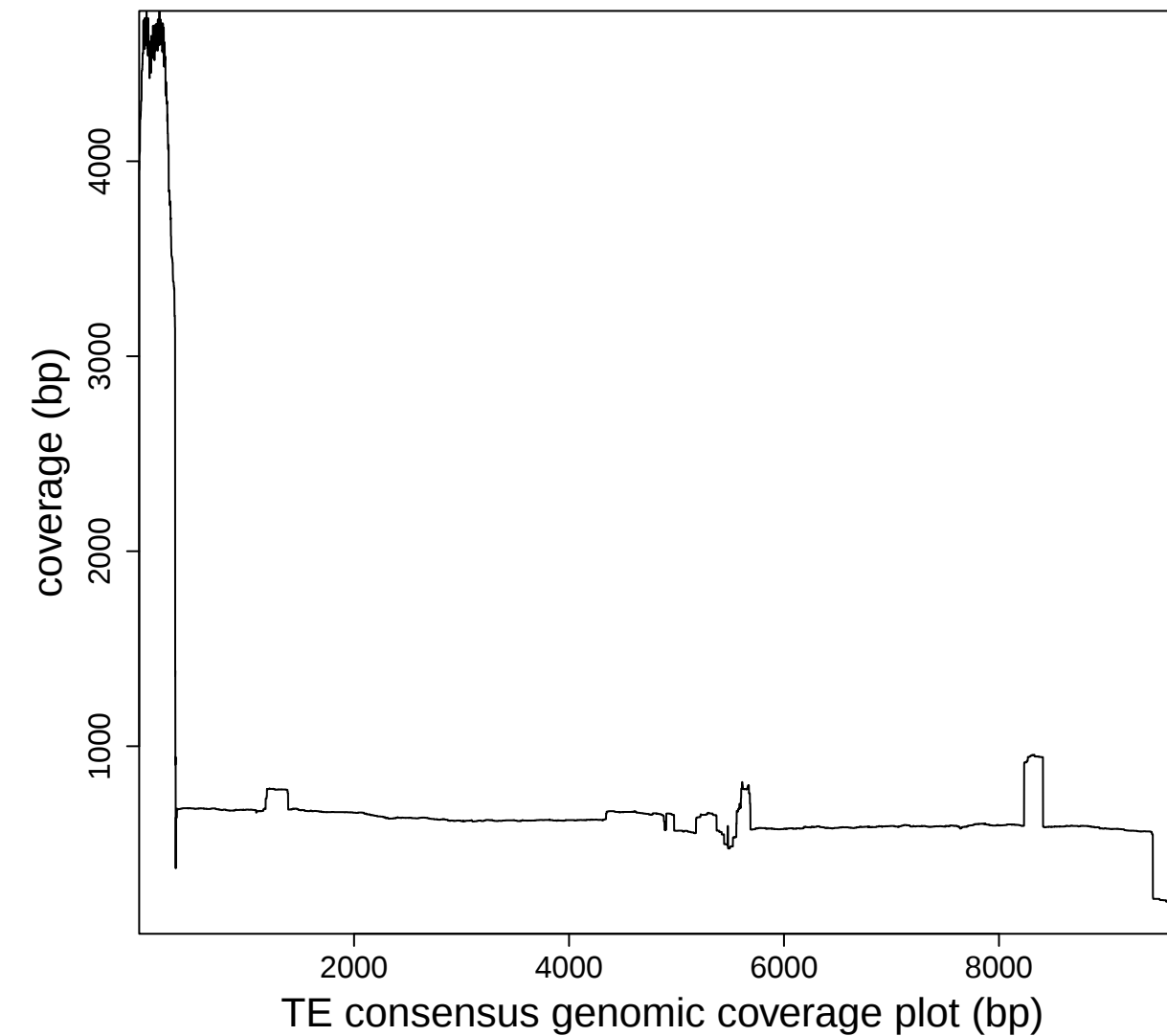
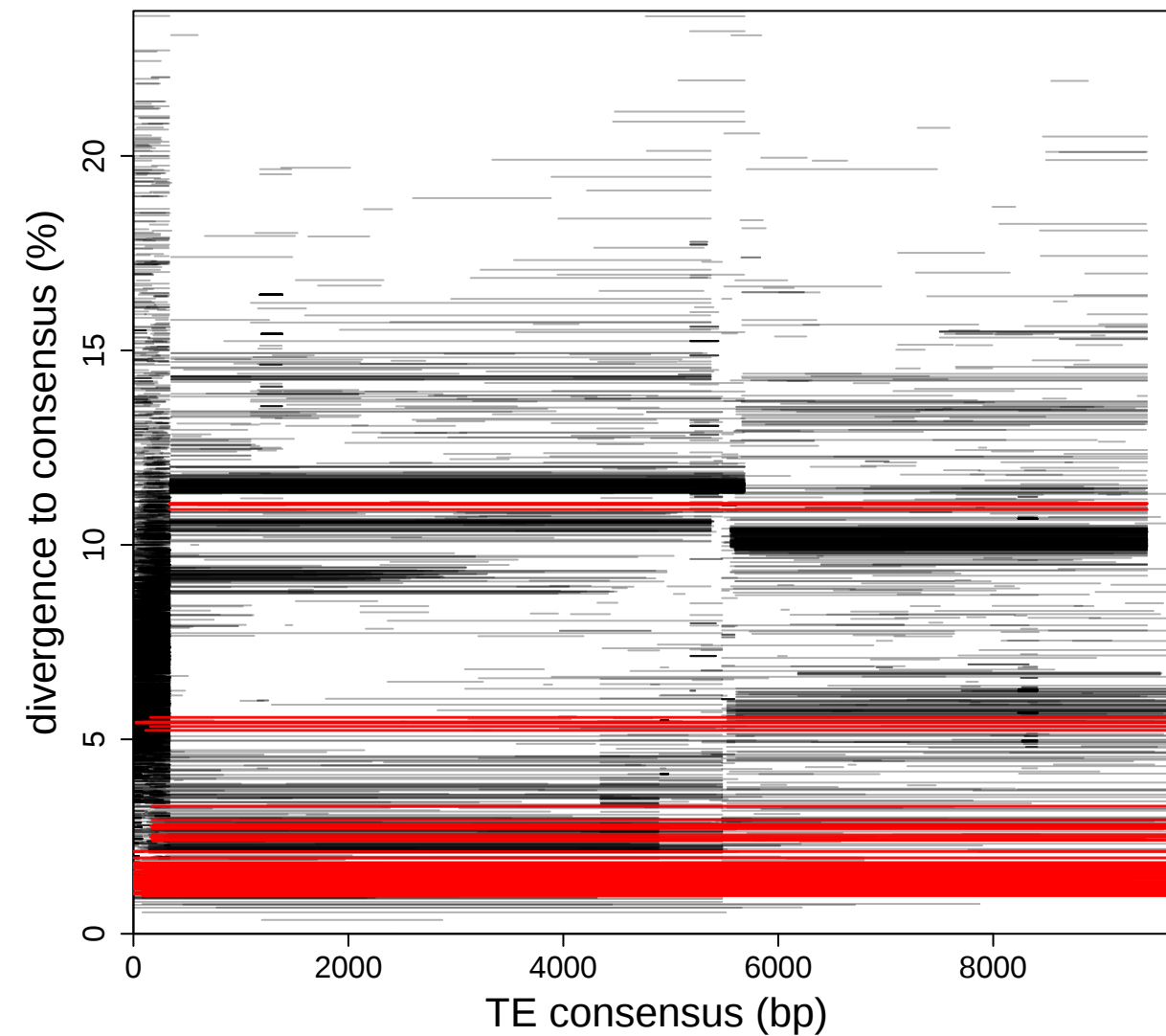
f.bed_g_3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 16306bp; fragments: 13883; full length: 3 (>=14675.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

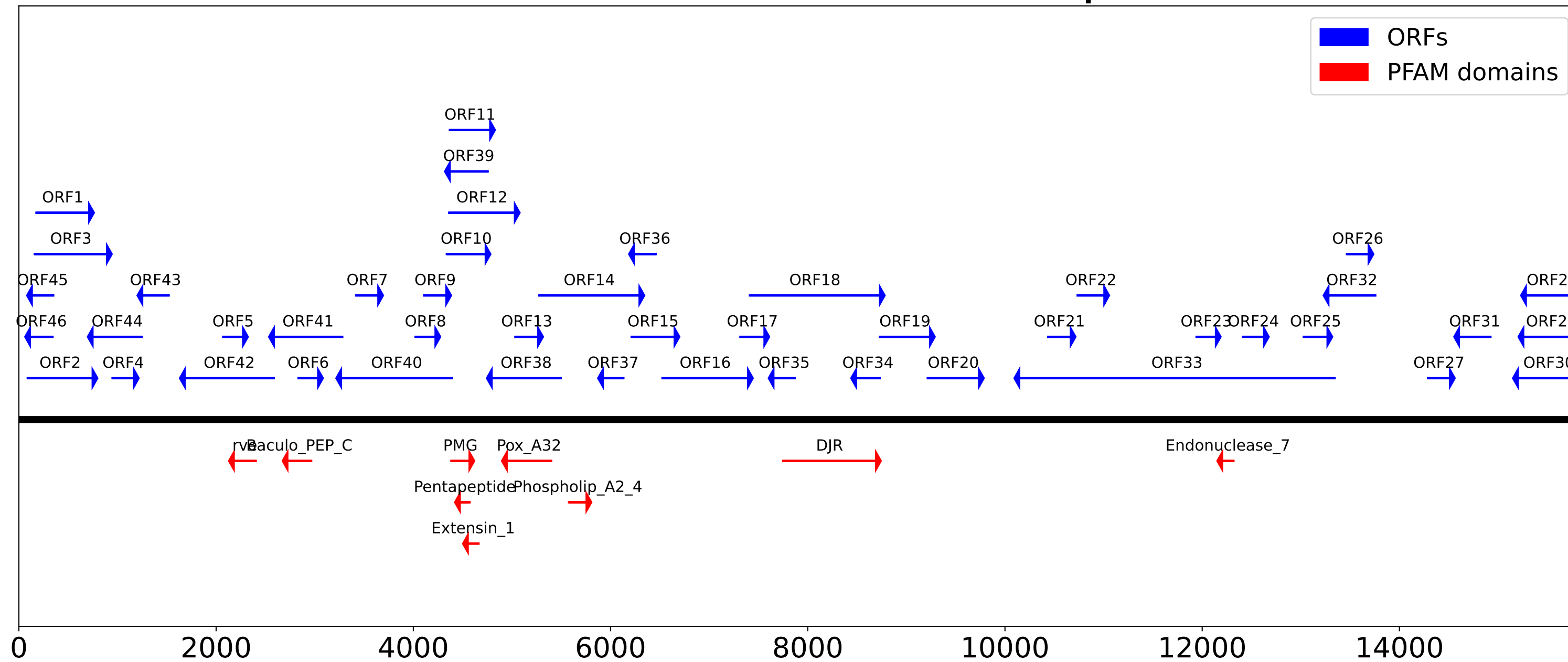


TE: rnd_5_family_757
size: 9738bp; fragments: 10307; full length: 84 (≥ 8764.2 bp)

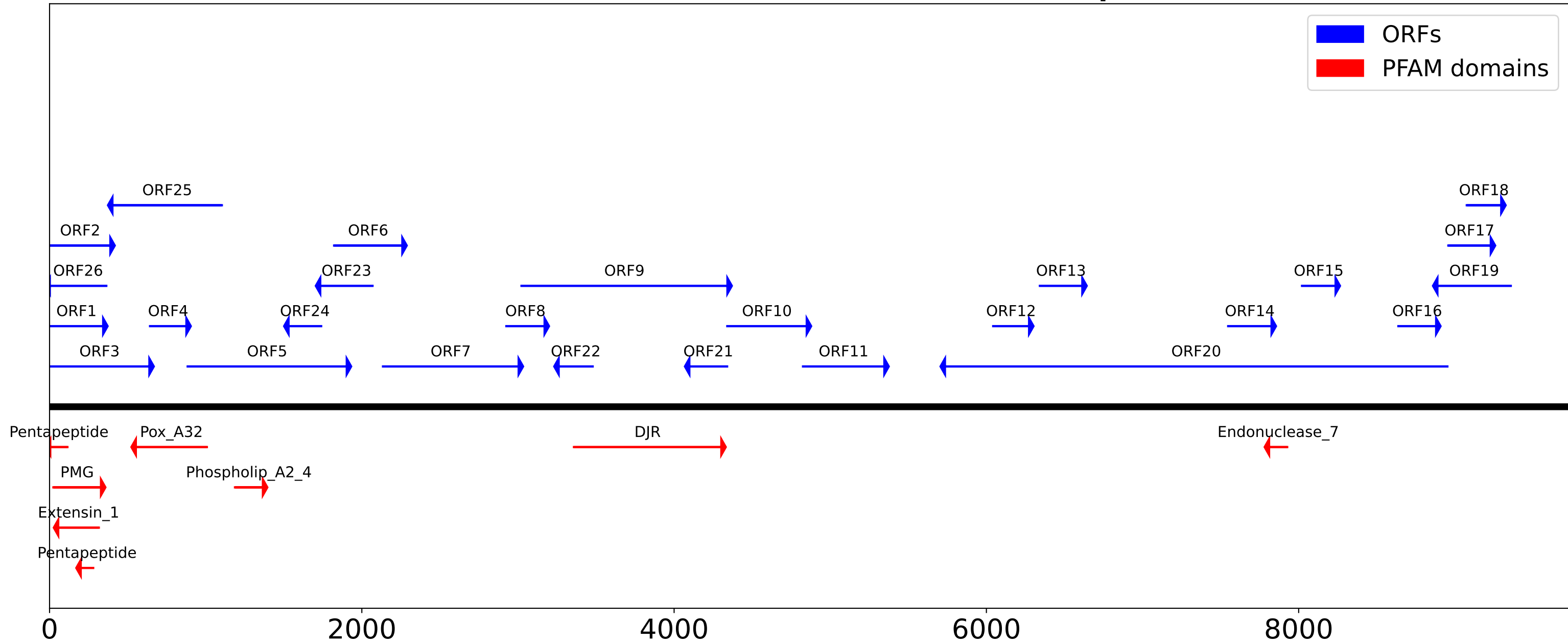
Before TEtrimmer 9738 bp



After TETrimmer ORF and PFAM domain plot



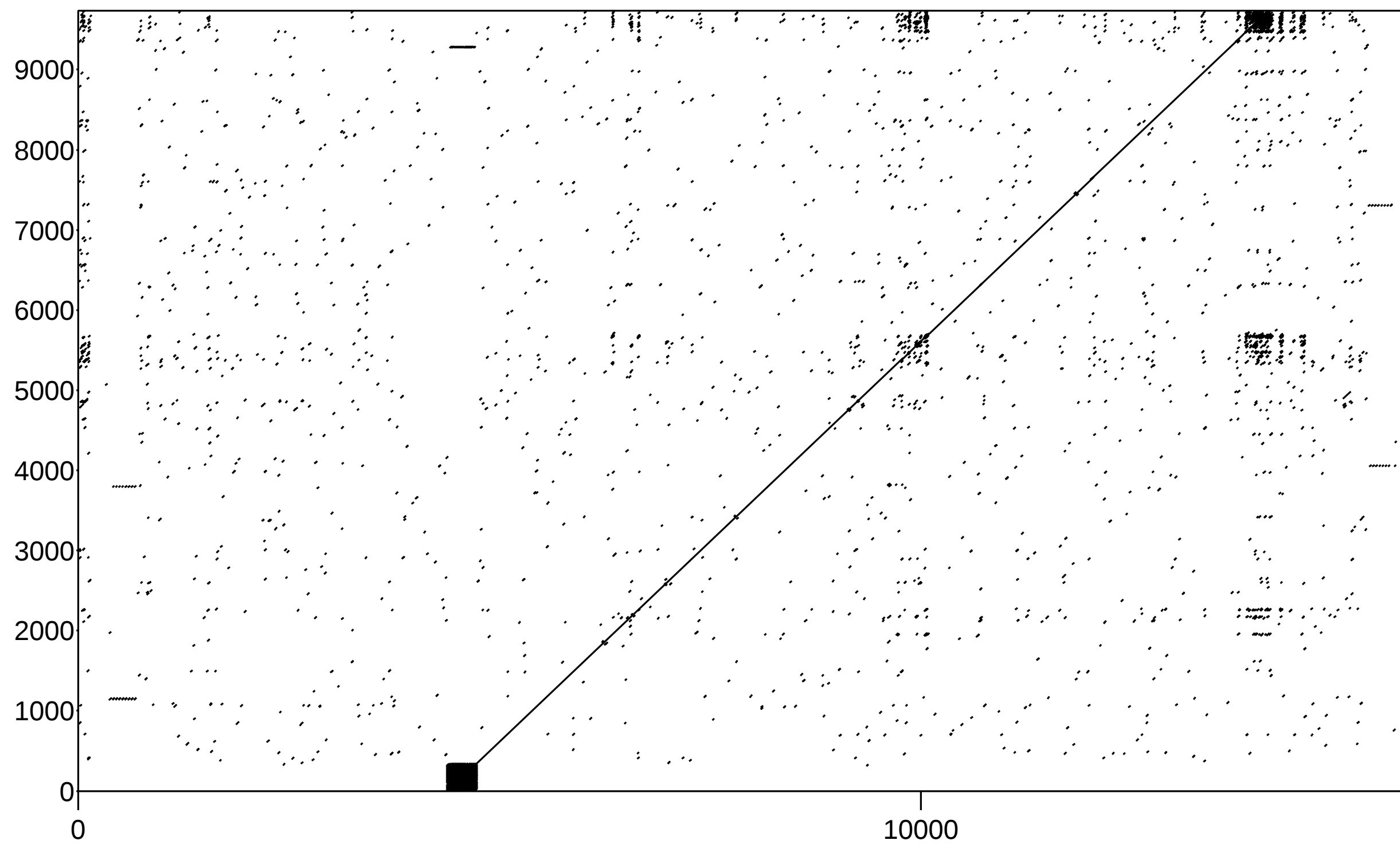
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)