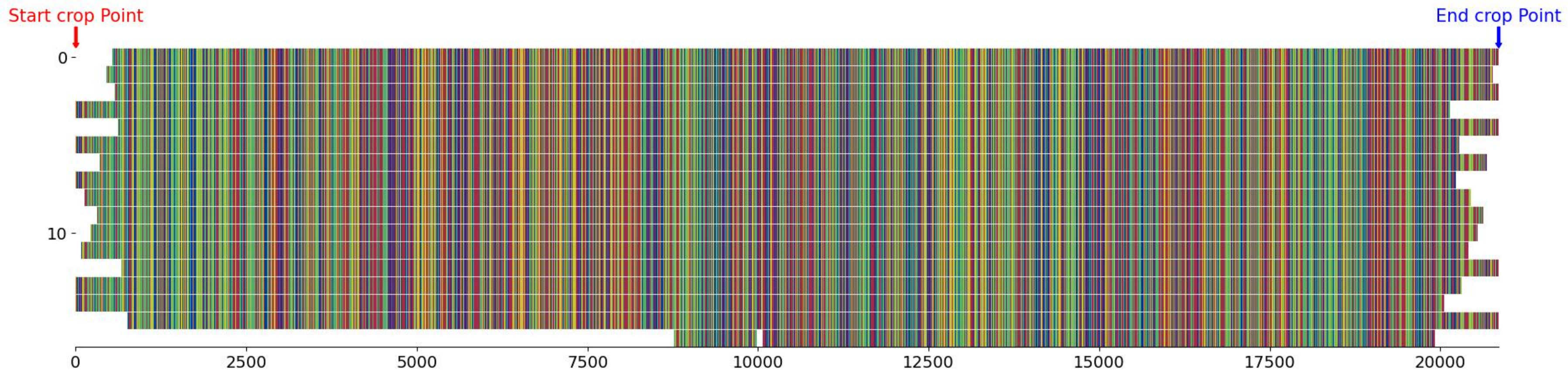


Start crop Point

End crop Point

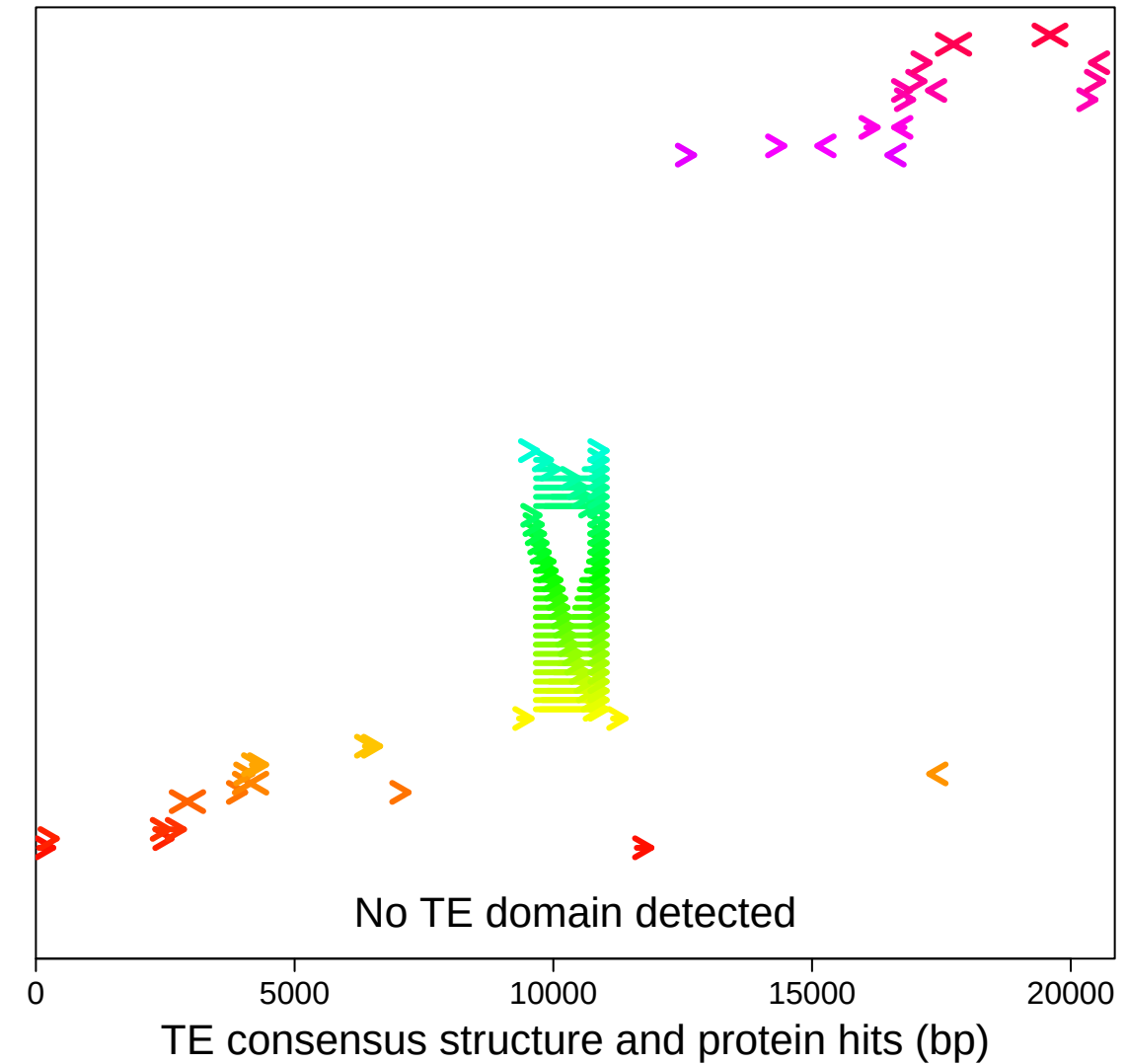
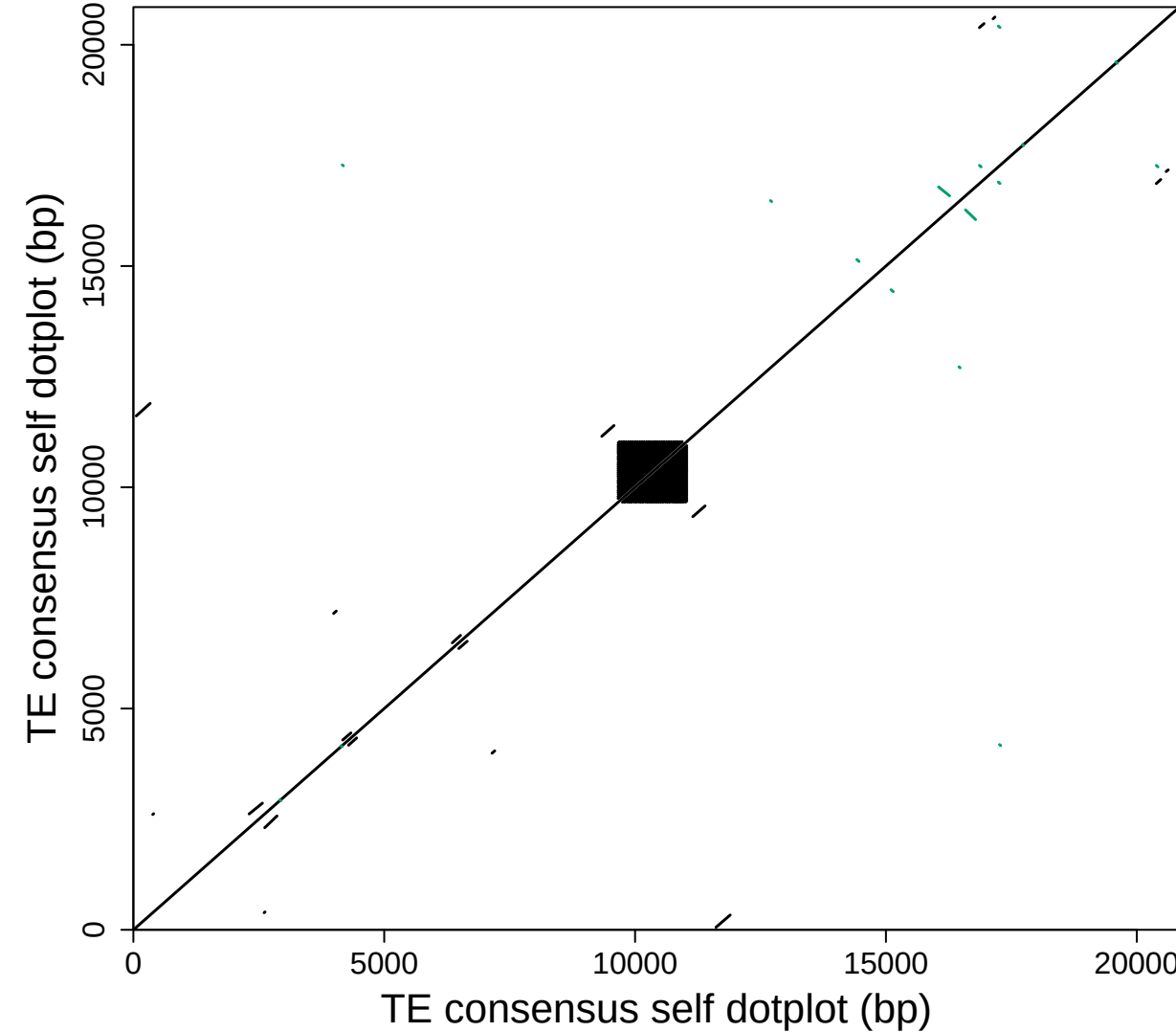
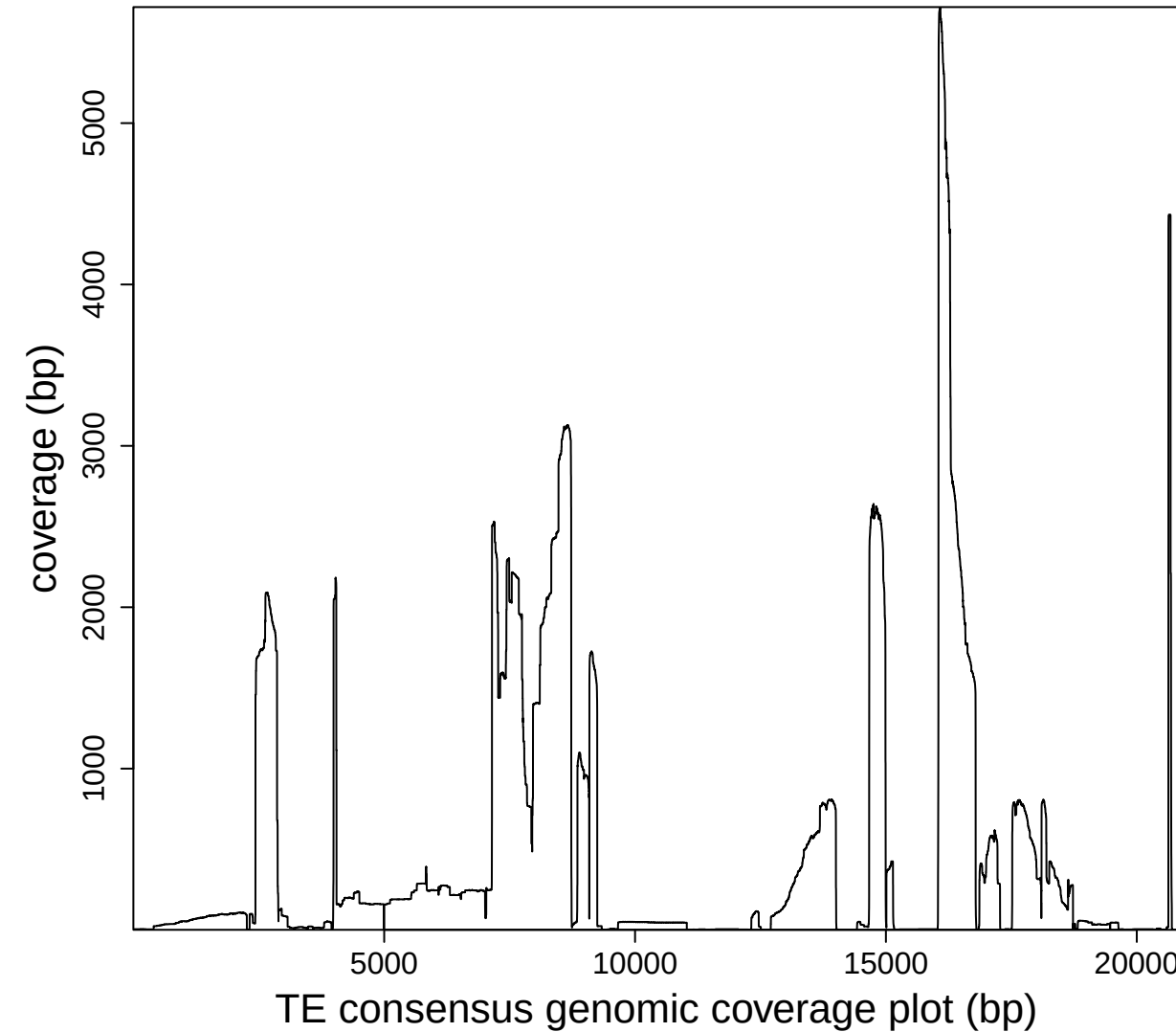
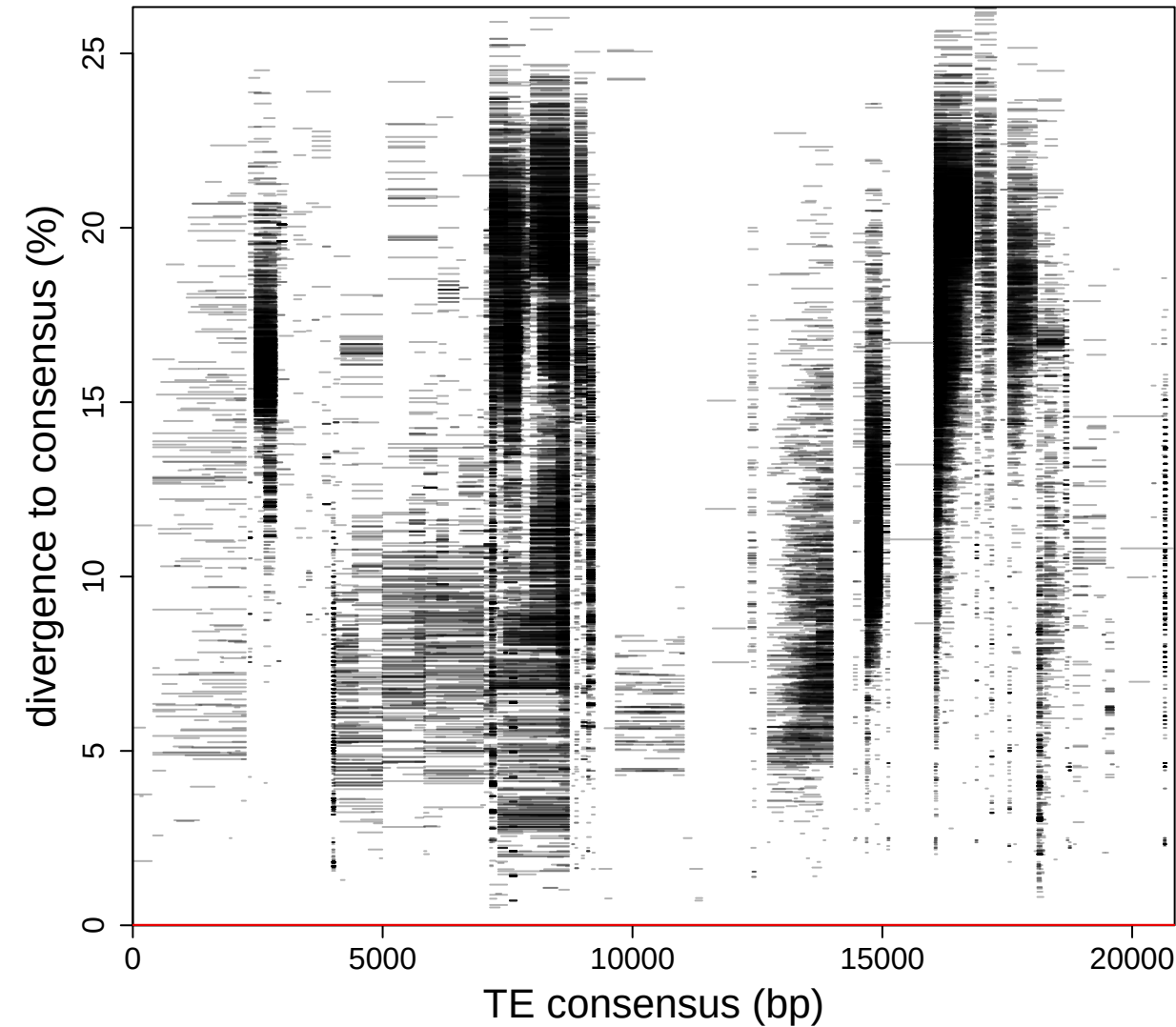
MSA length = 20851





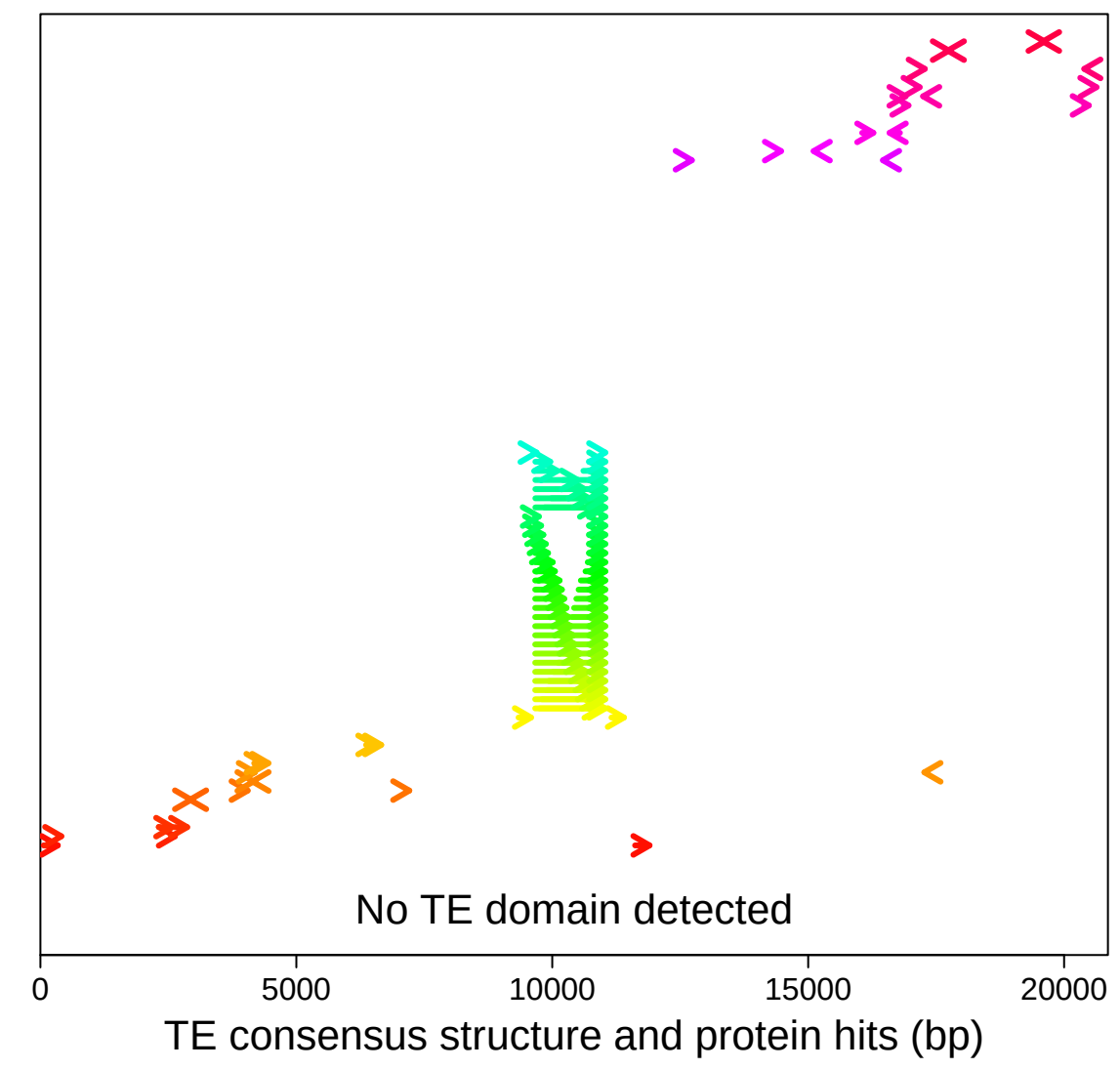
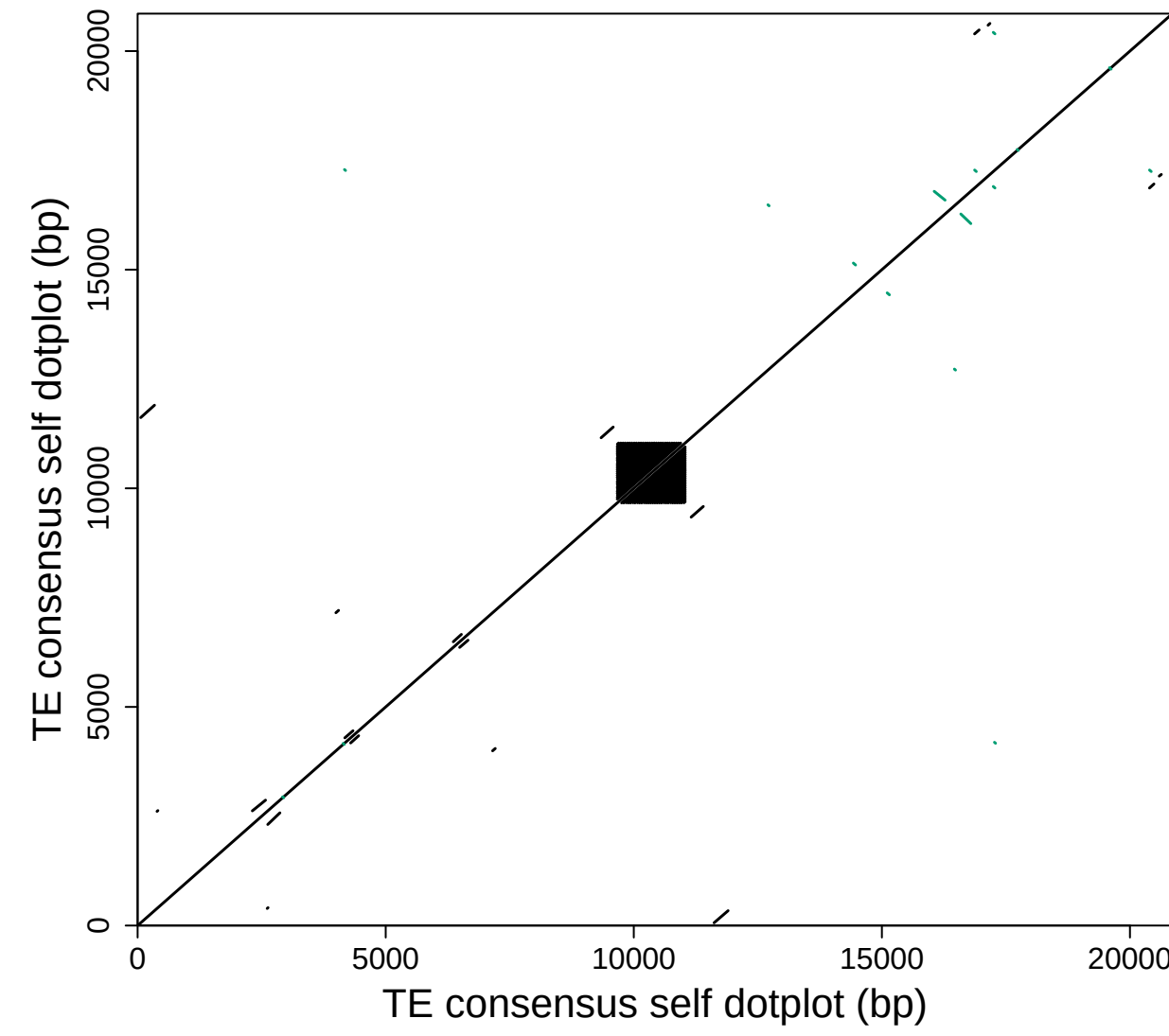
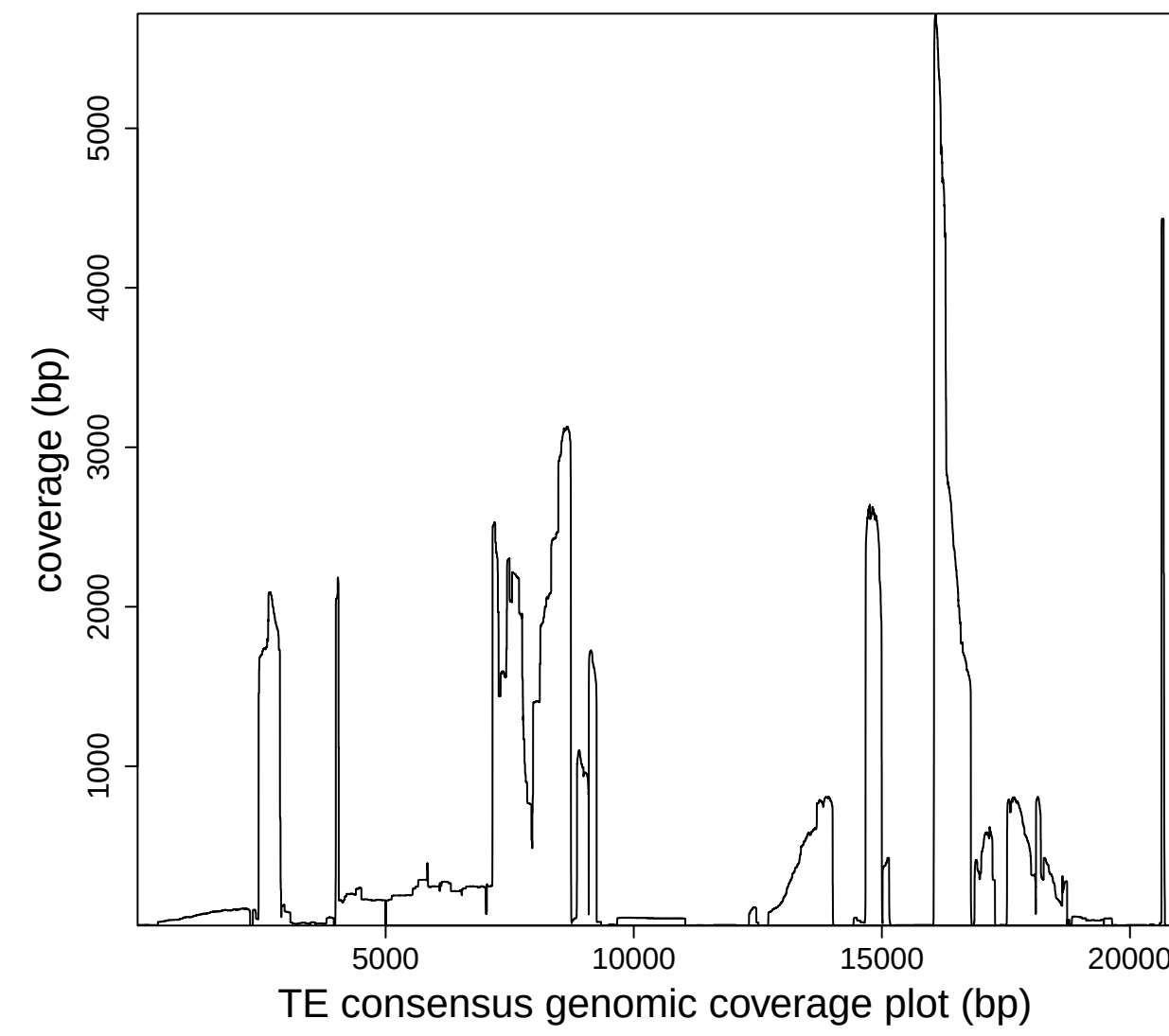
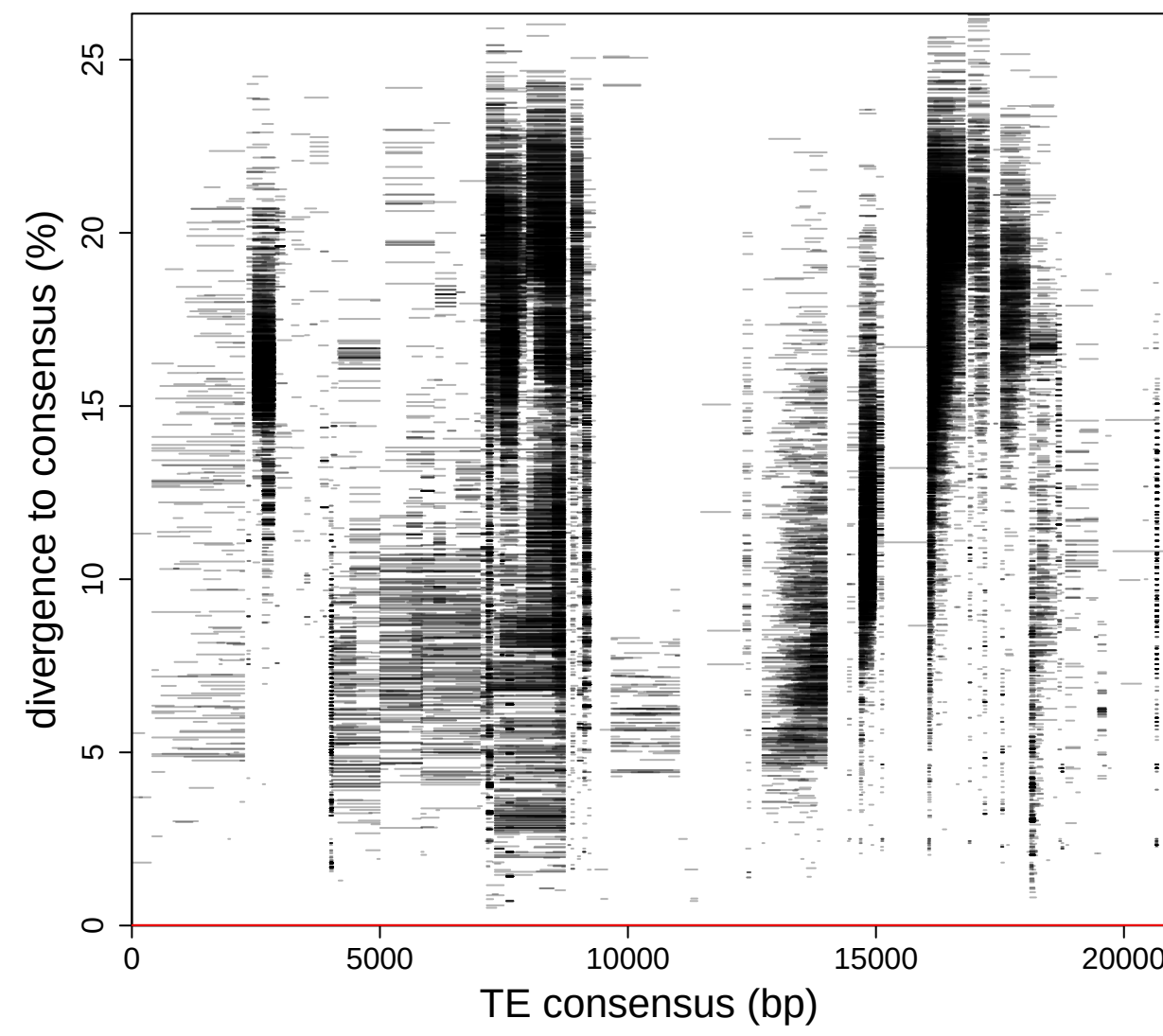
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa al
size: 20851bp; fragments: 34090; full length: 1 (>=18765.9bp)

After TEtrimmer 20851 bp

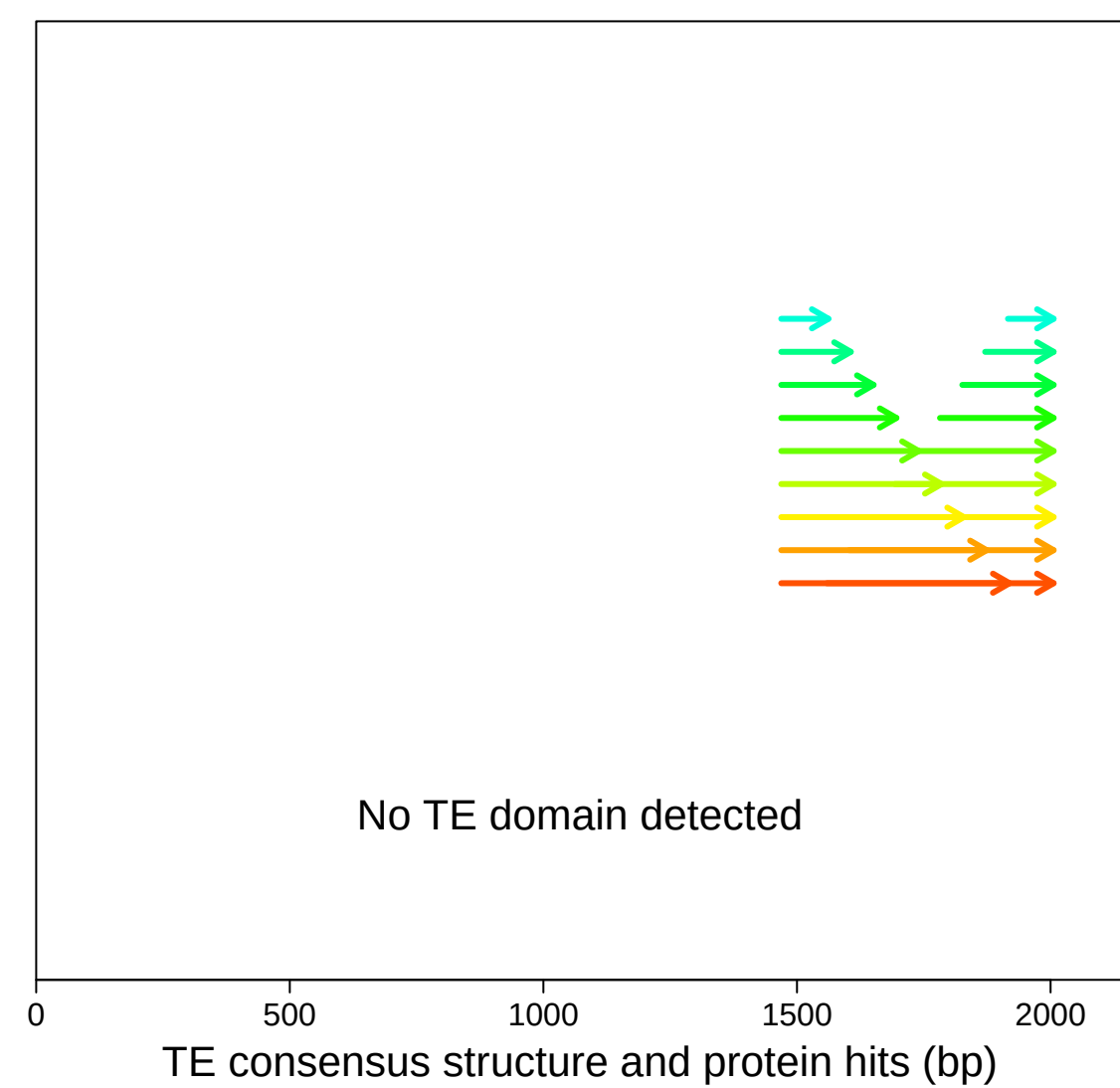
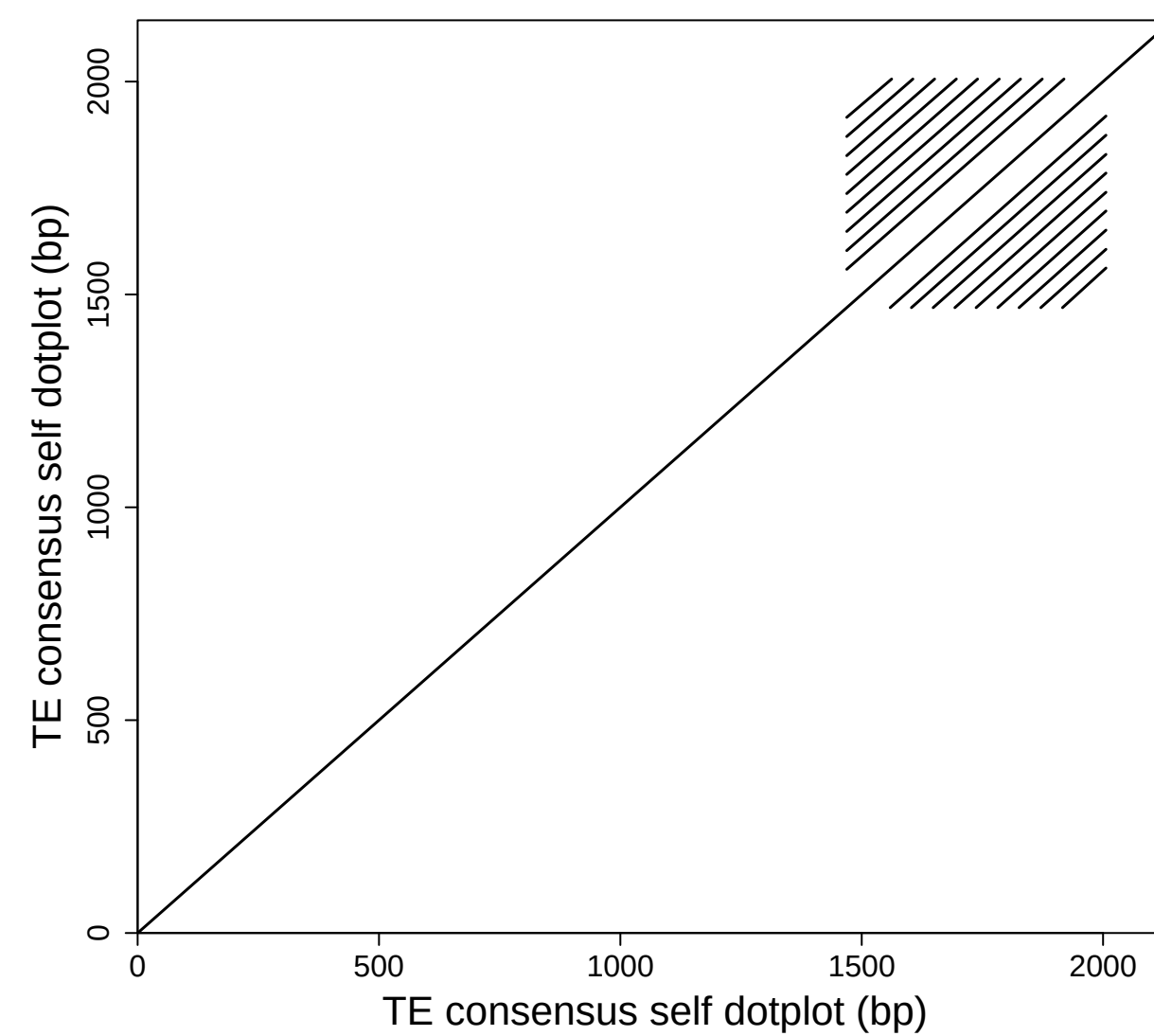
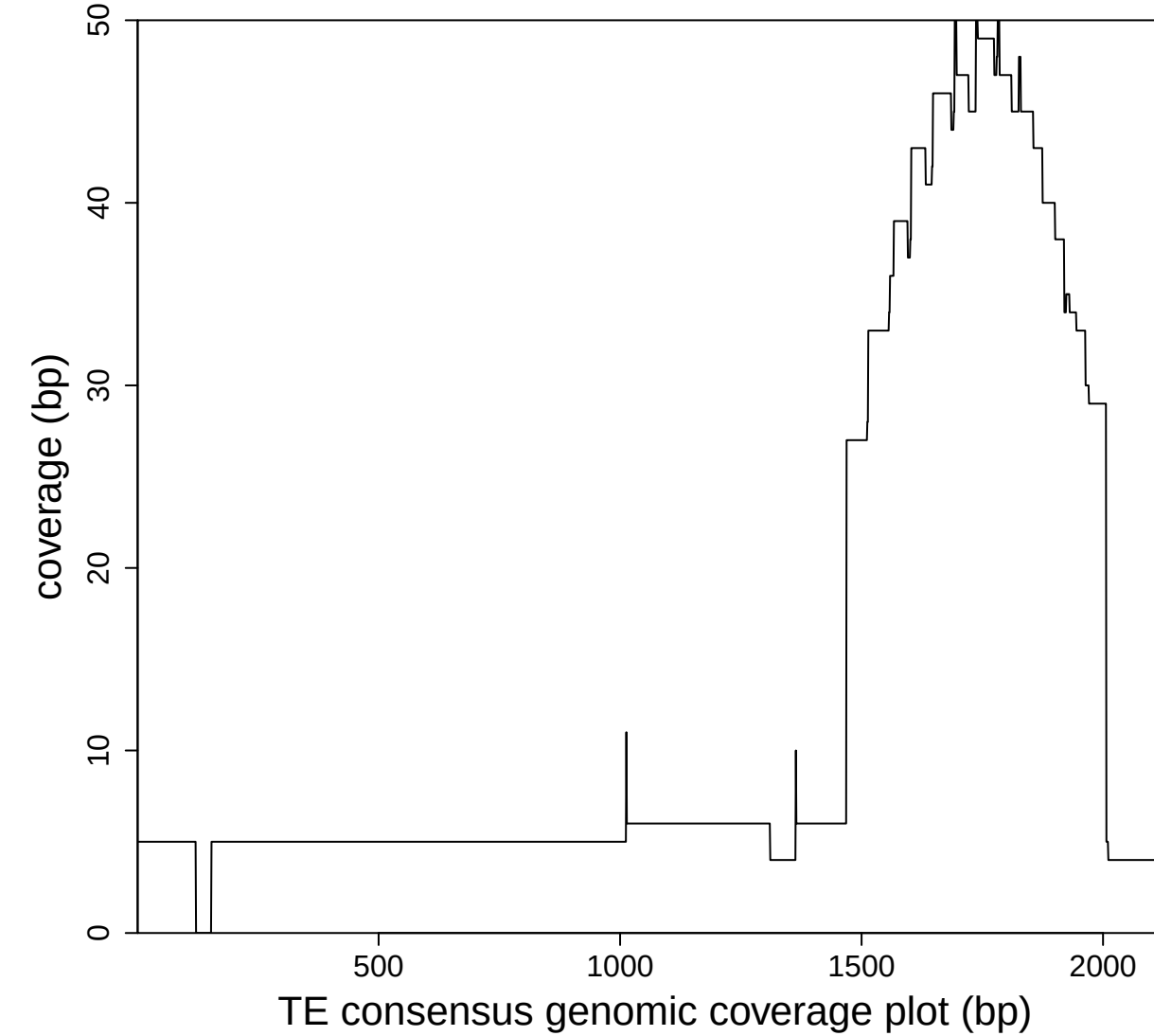
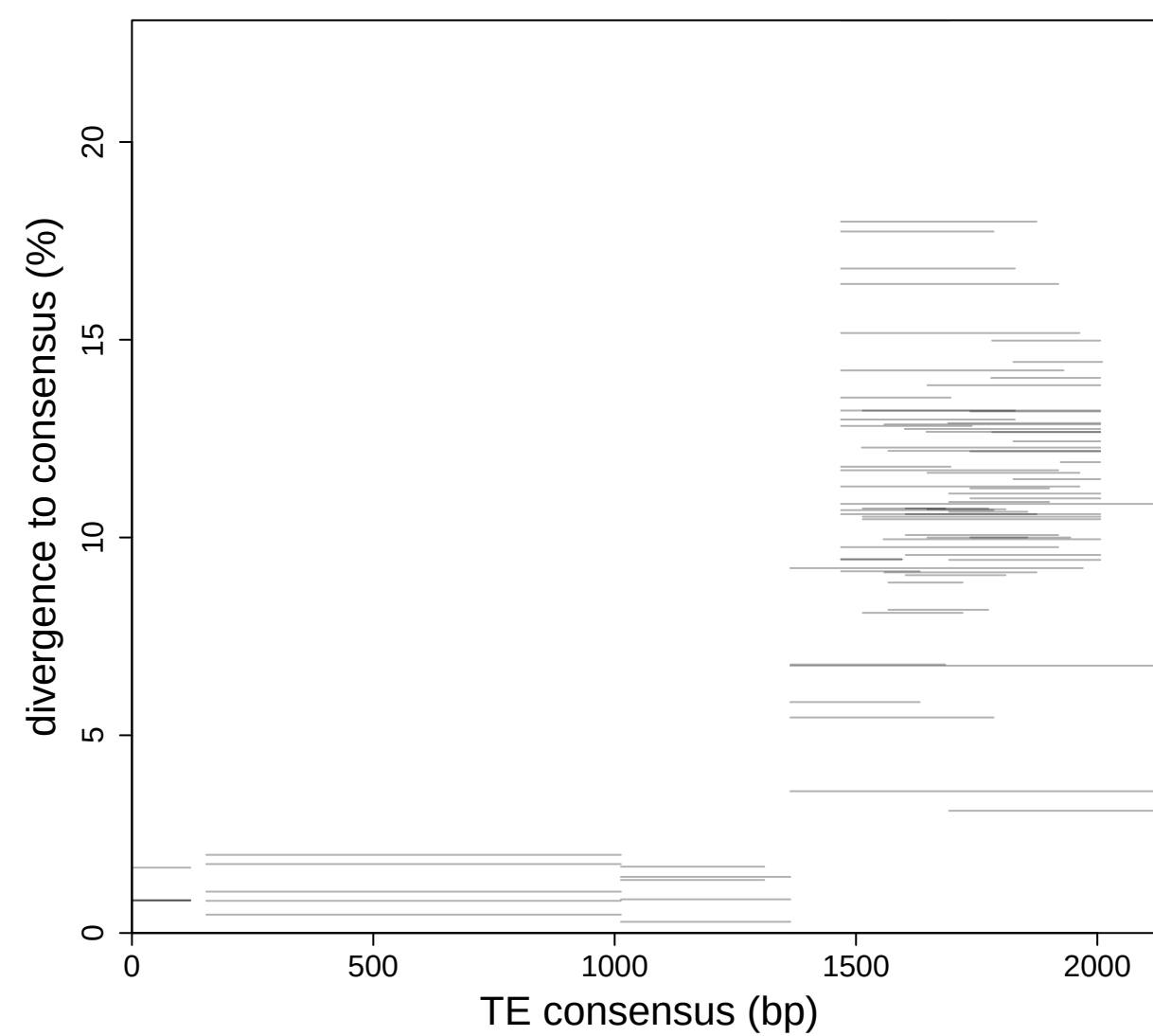


d_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa
size: 20856bp; fragments: 34090; full length: 1 (>=18770.4bp)

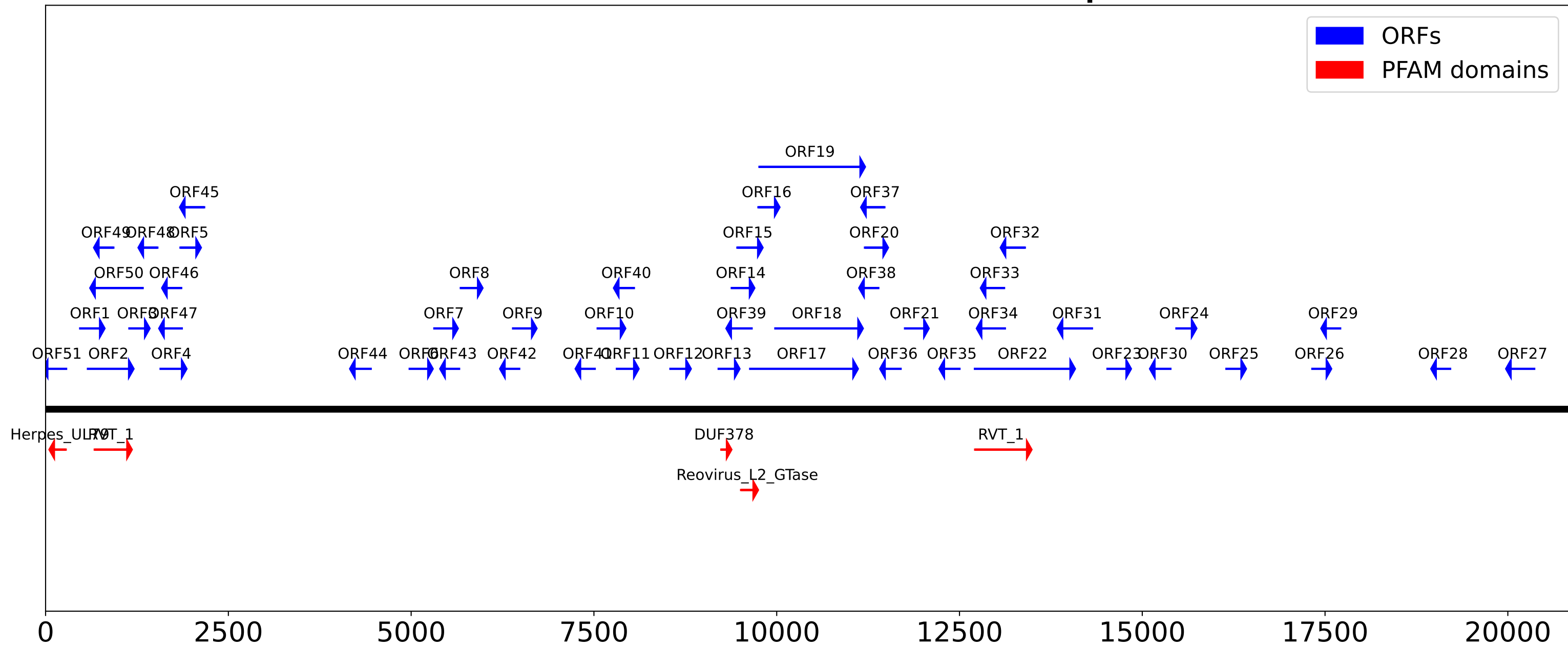
After TEtrimmer Extended plot Blue lines are boundaries



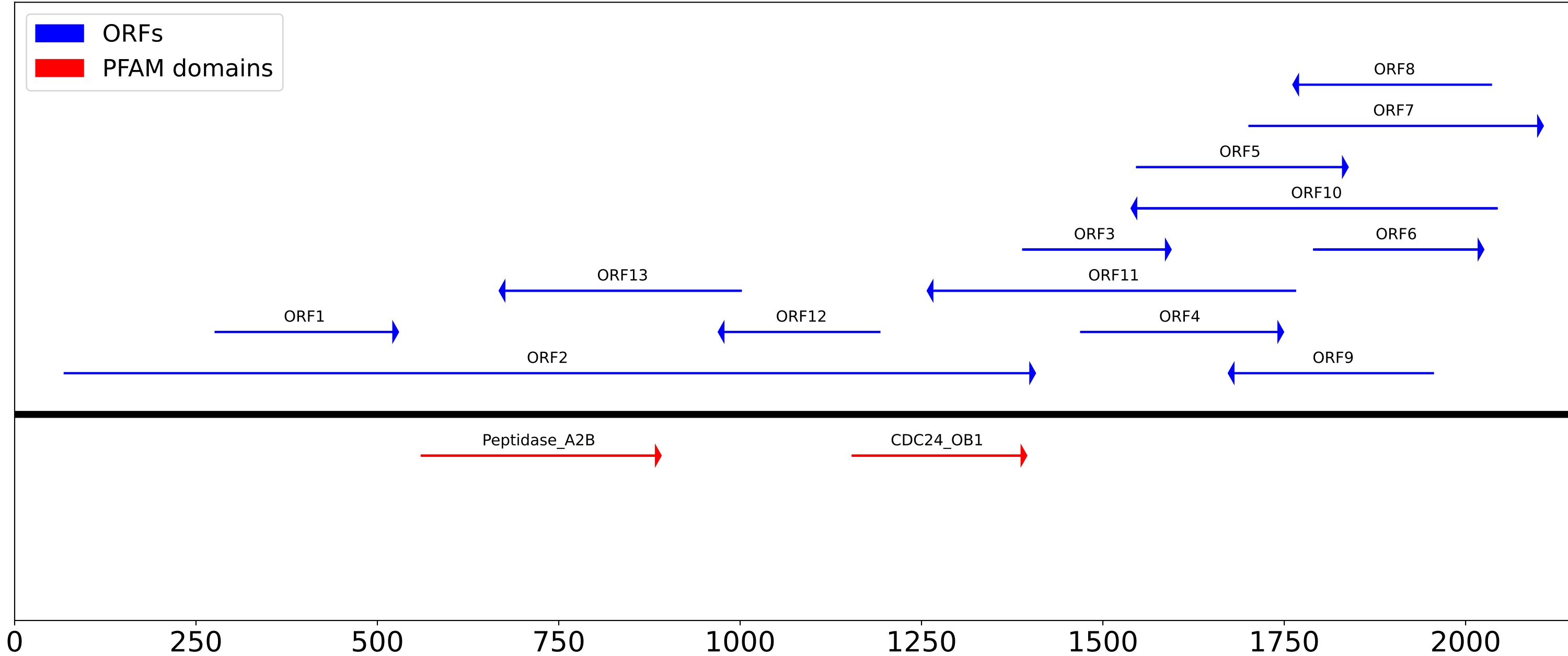
TE: ltr_1_family_690
size: 2144bp; fragments: 84; full length: 0 (>=1929.6bp)



After TETrimmer ORF and PFAM domain plot



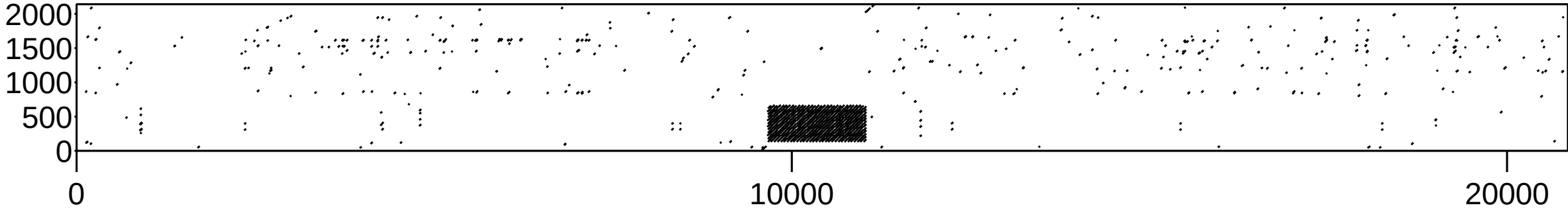
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)