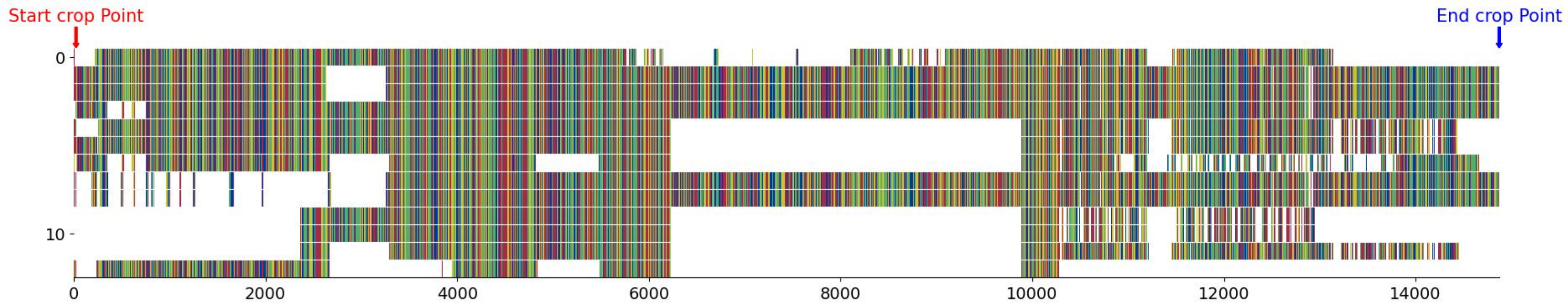


Start crop Point

End crop Point

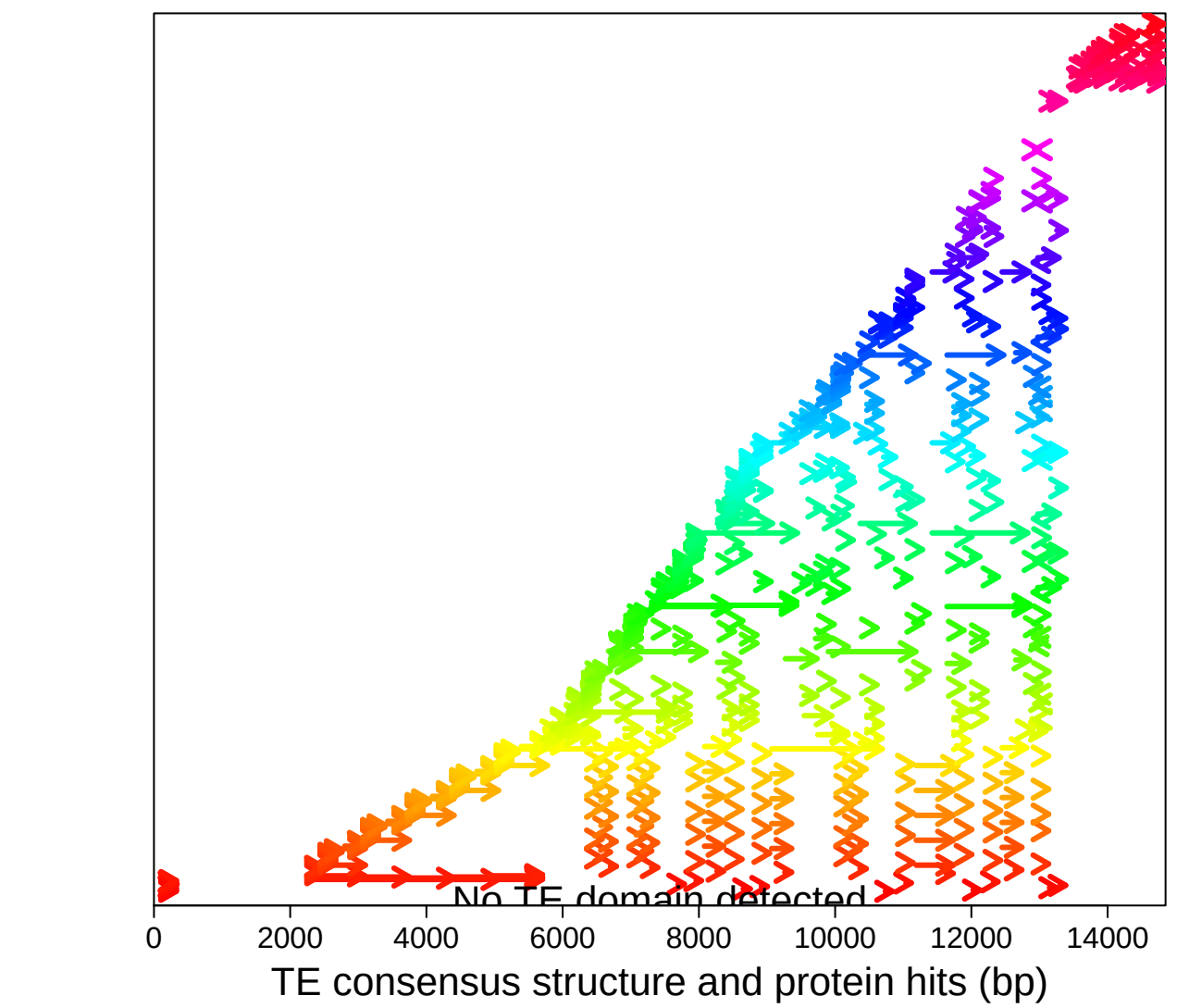
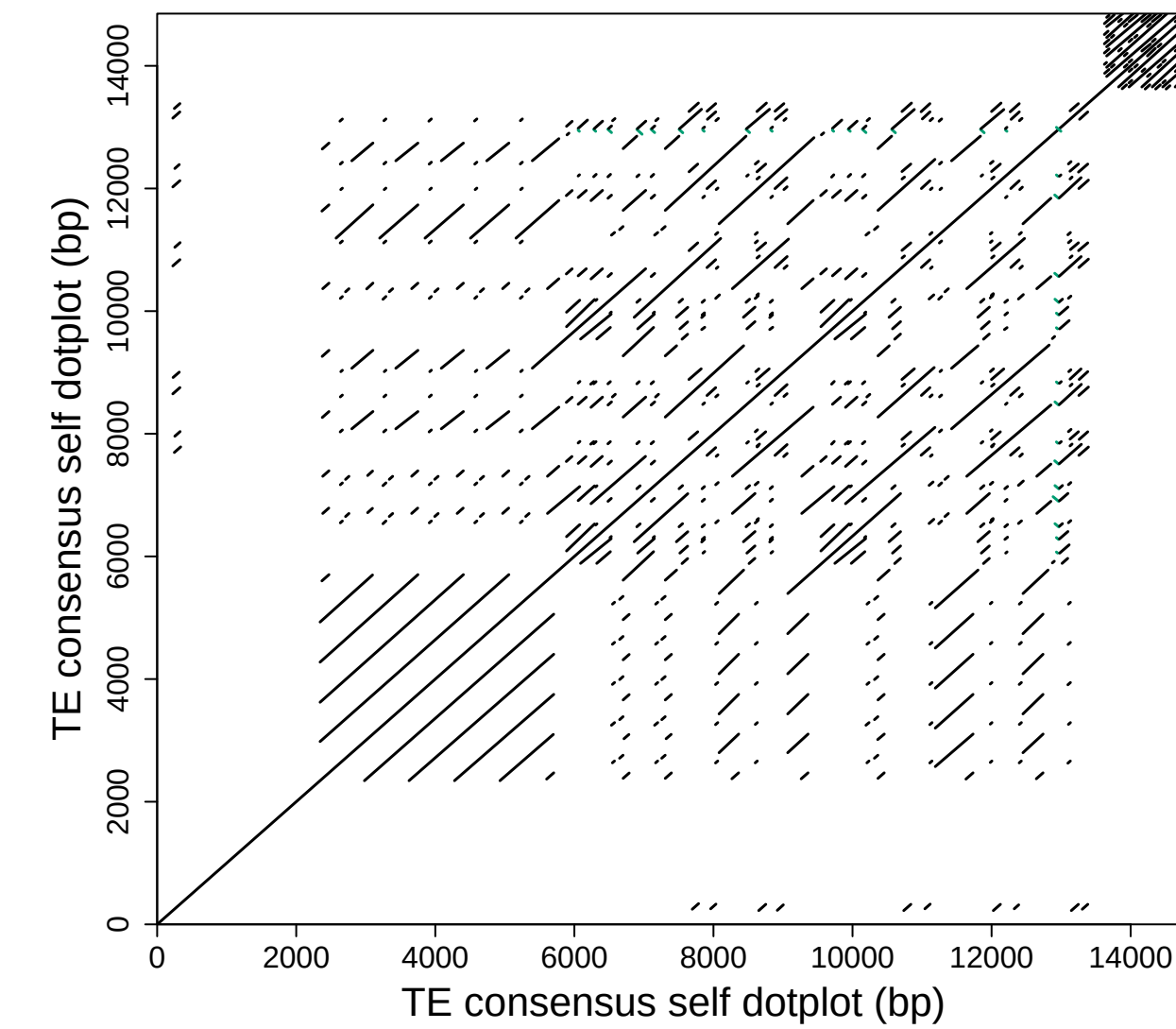
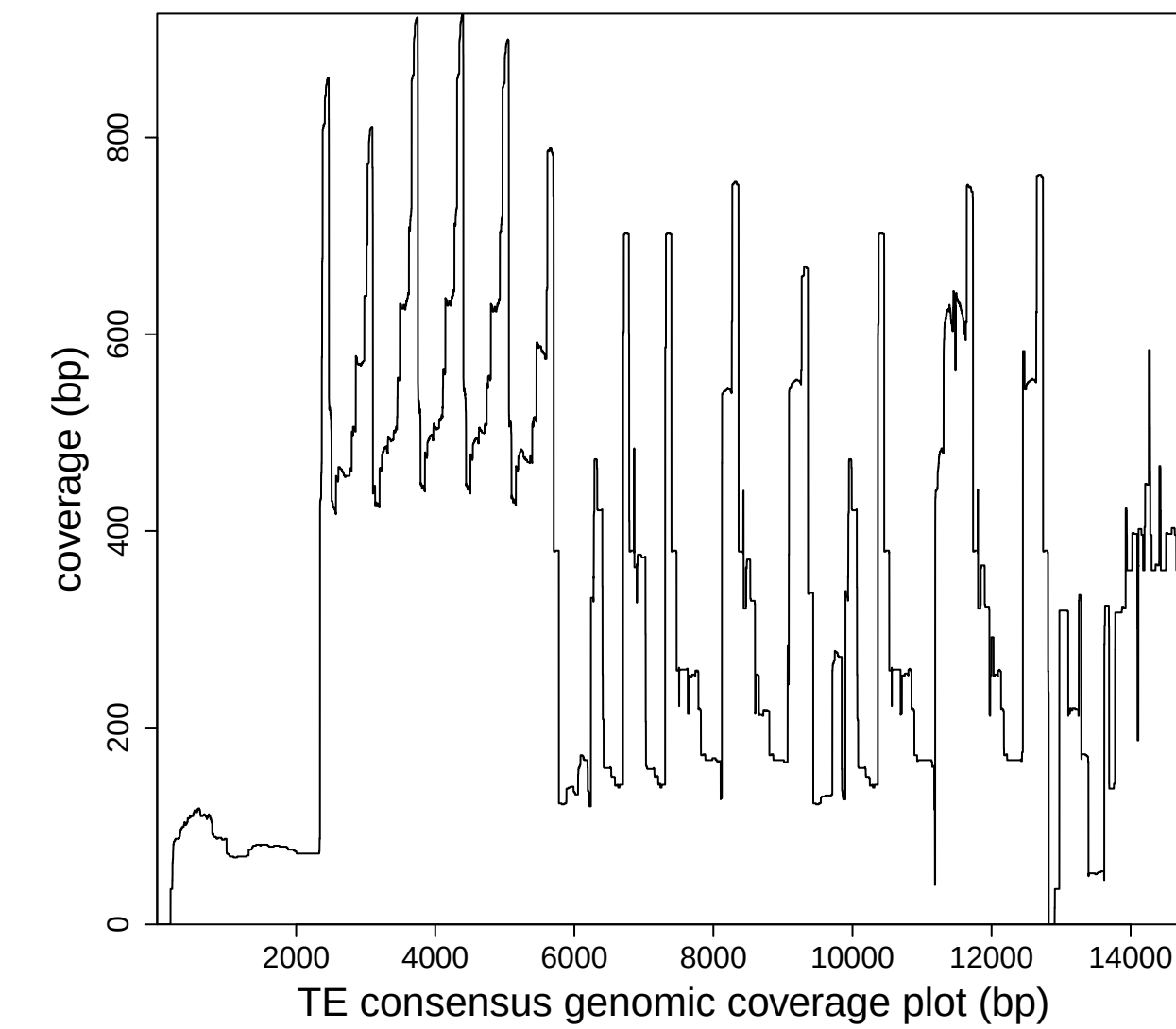
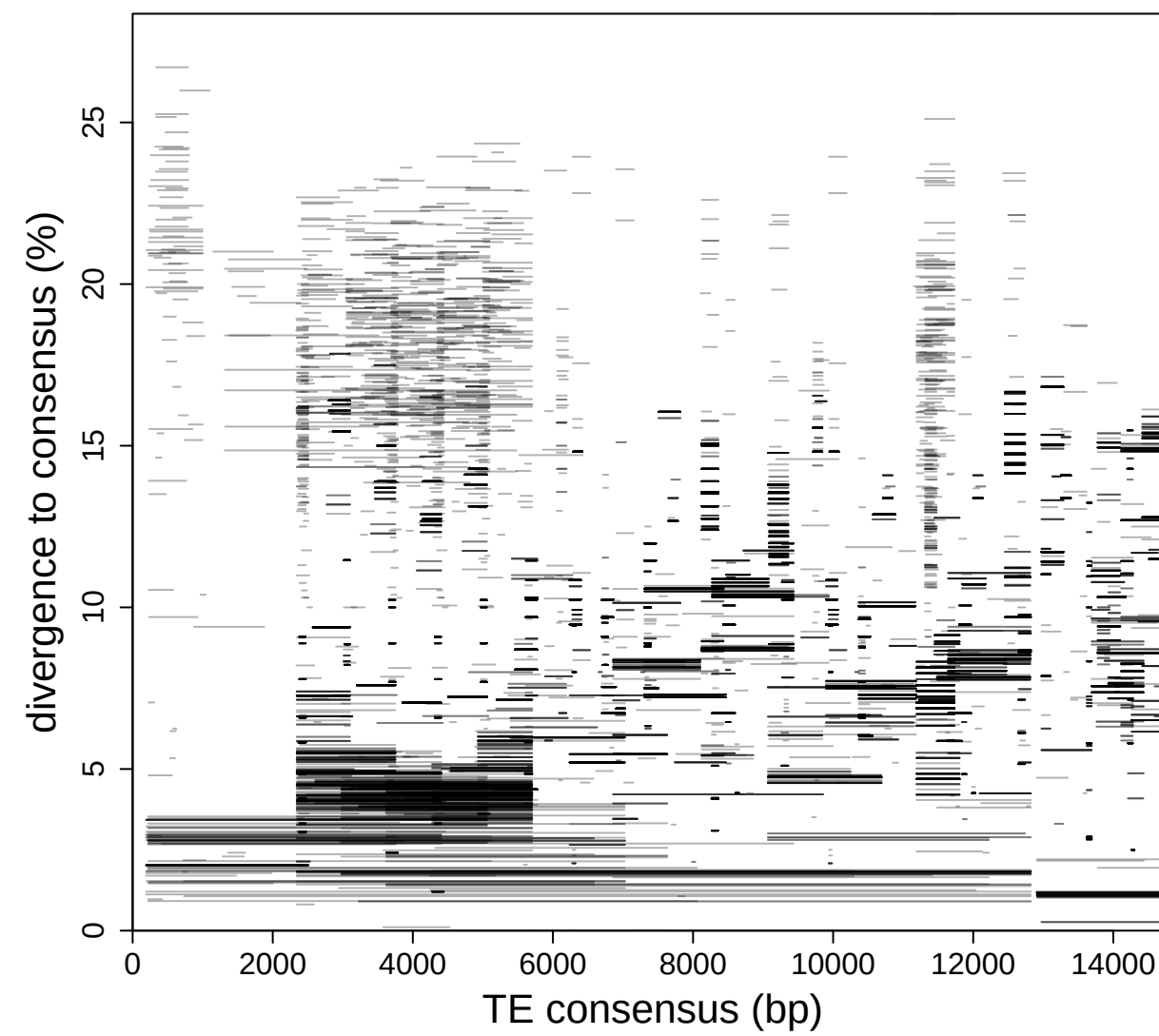
MSA length = 14851





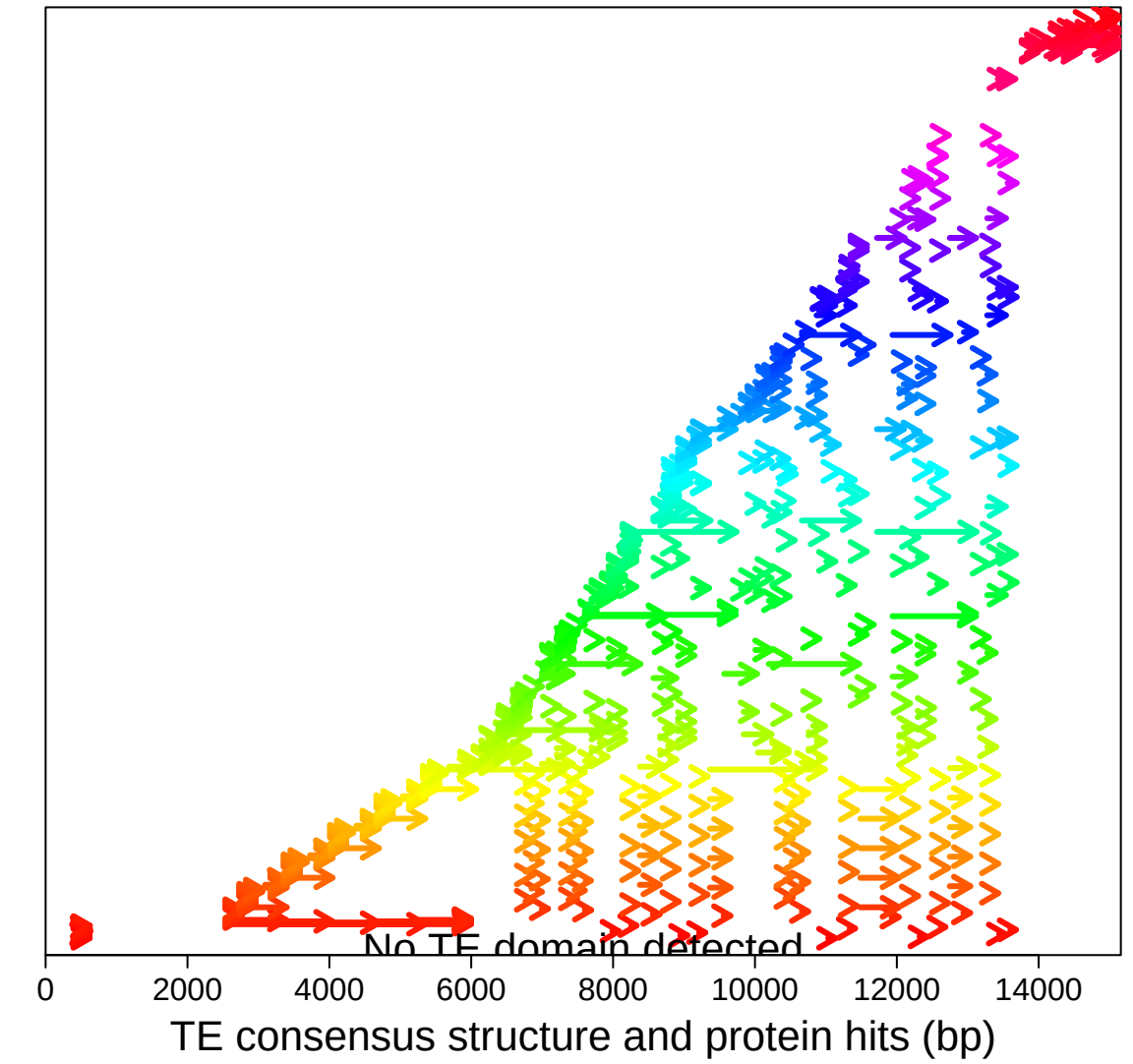
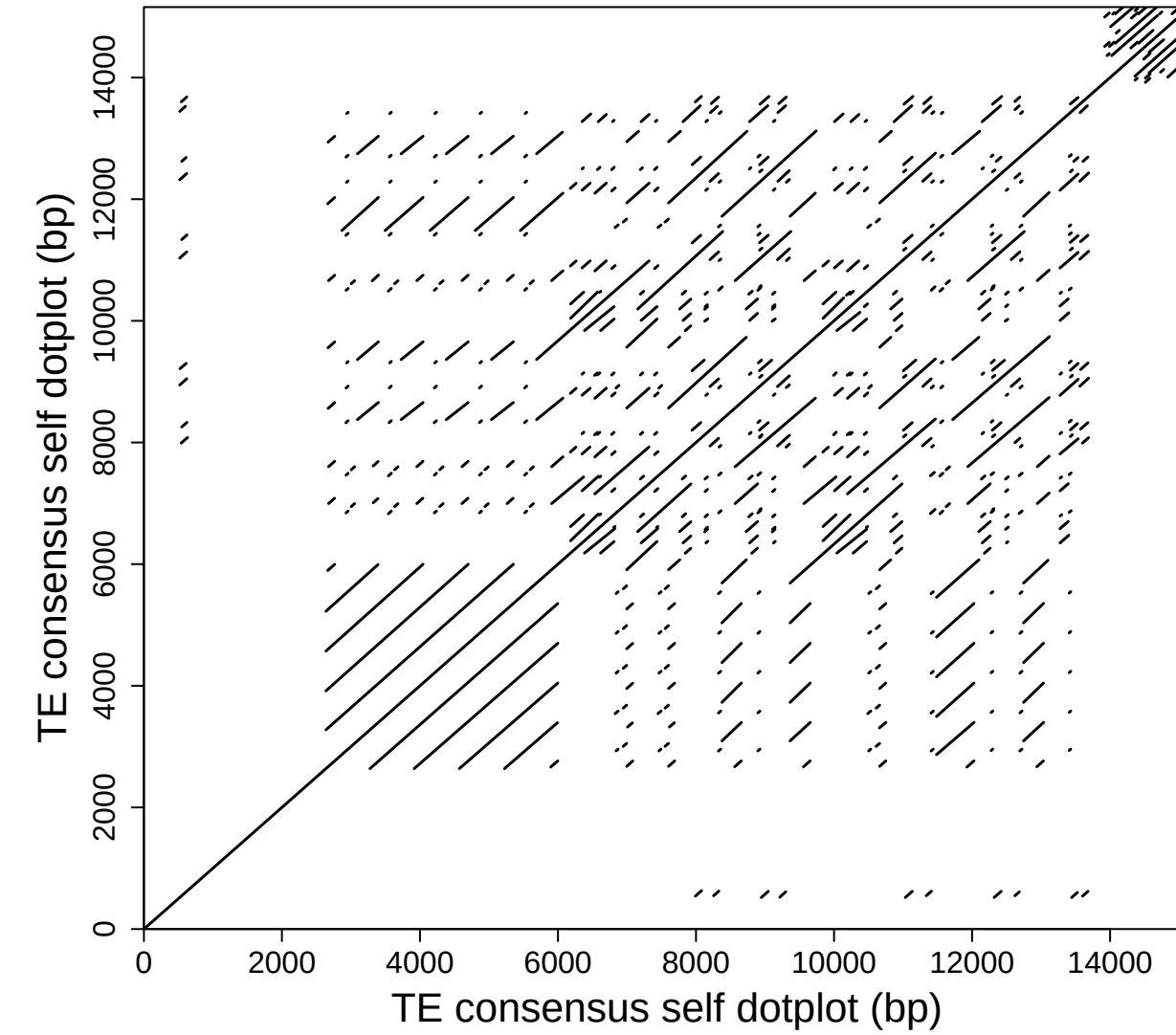
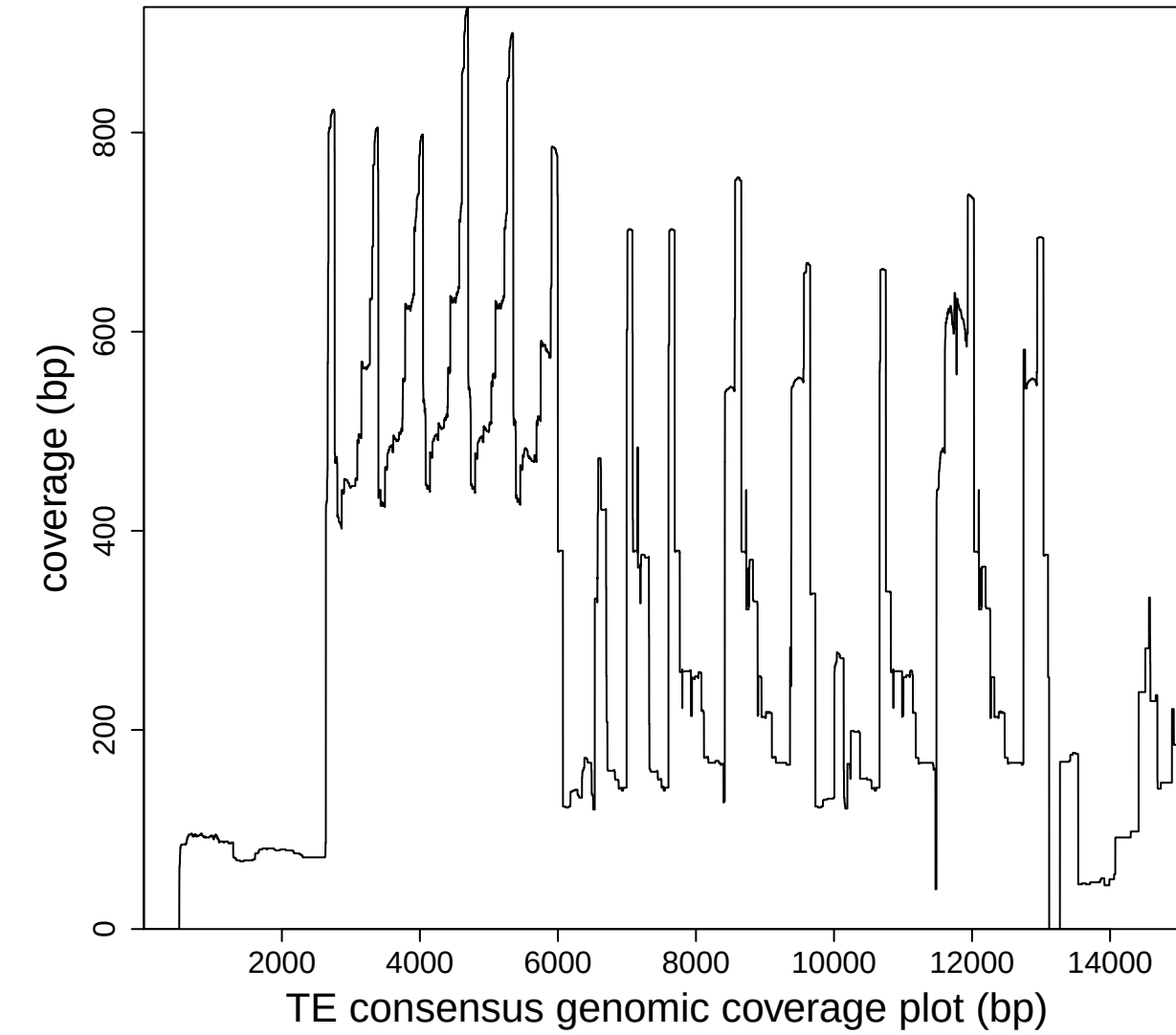
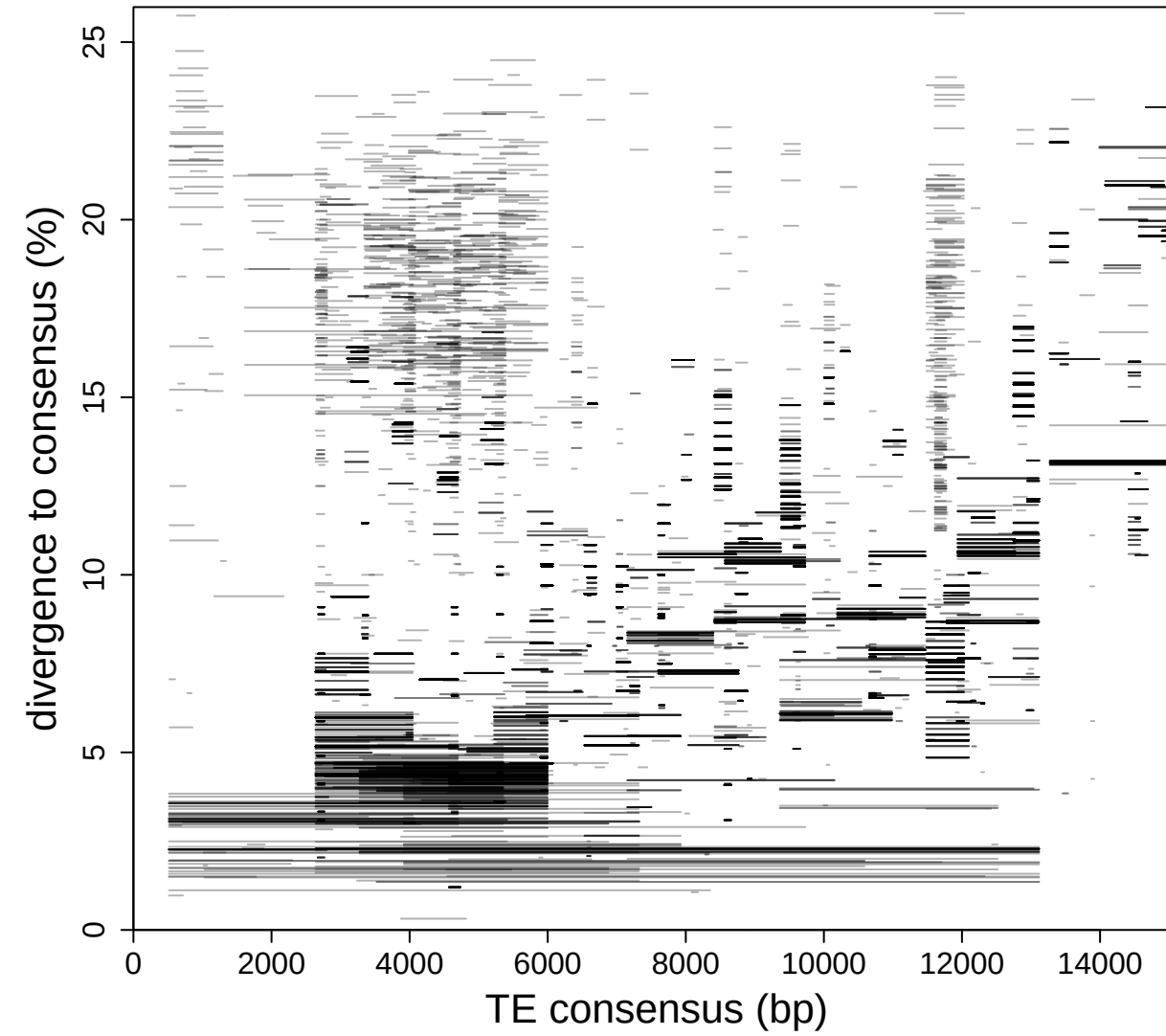
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 14851bp; fragments: 11640; full length: 0 (>=13365.9bp)

After TEtrimmer 14851 bp



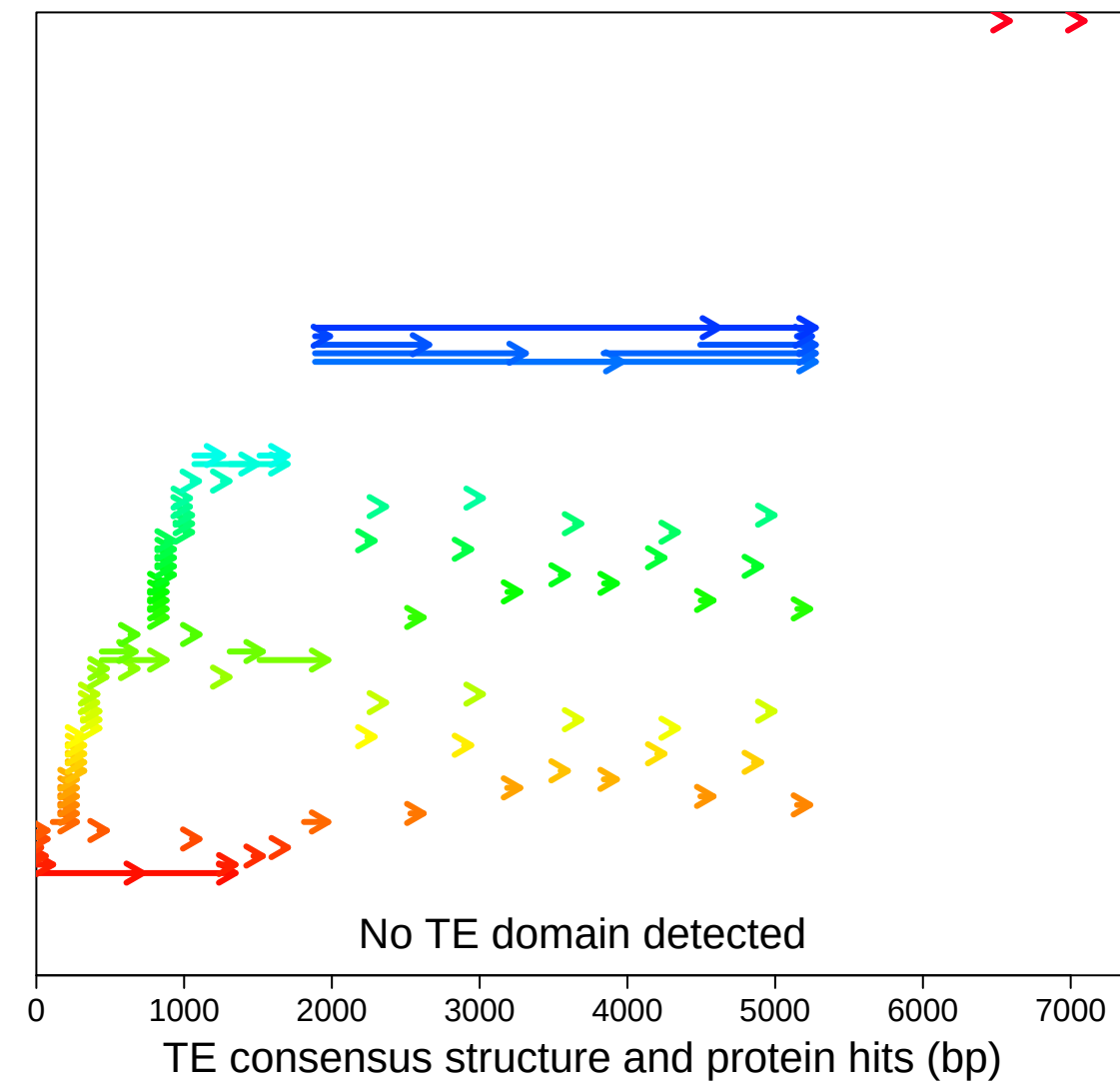
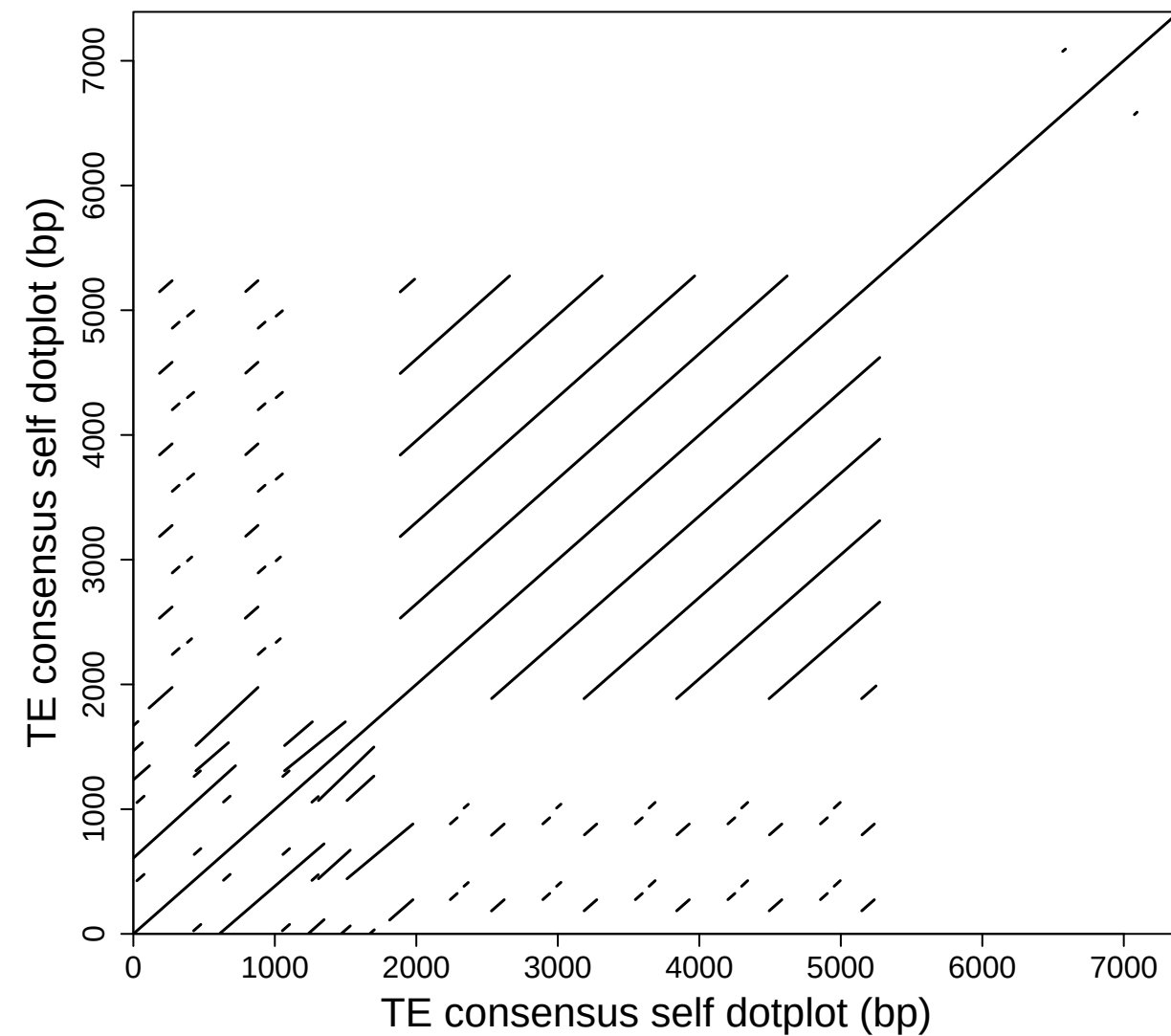
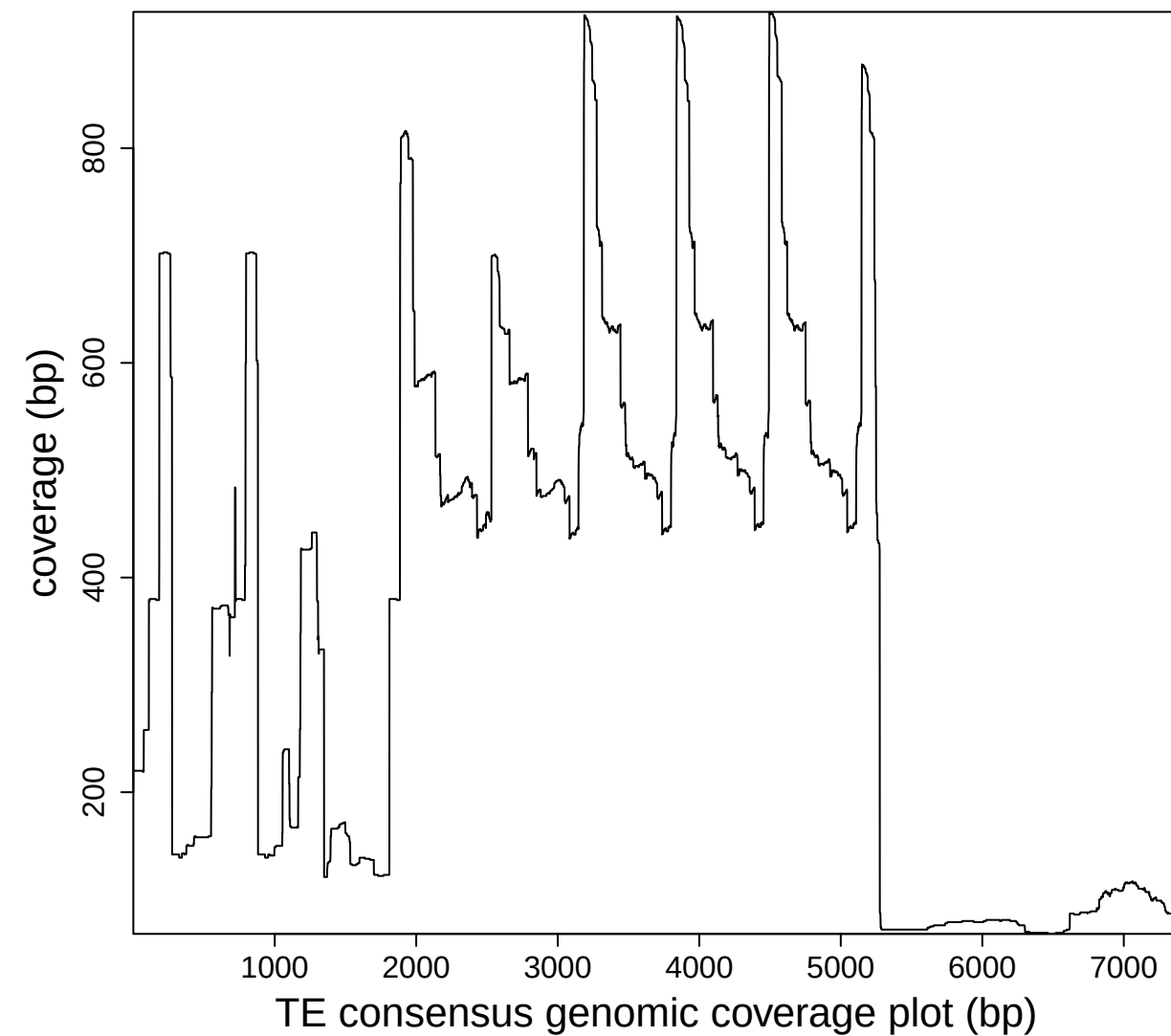
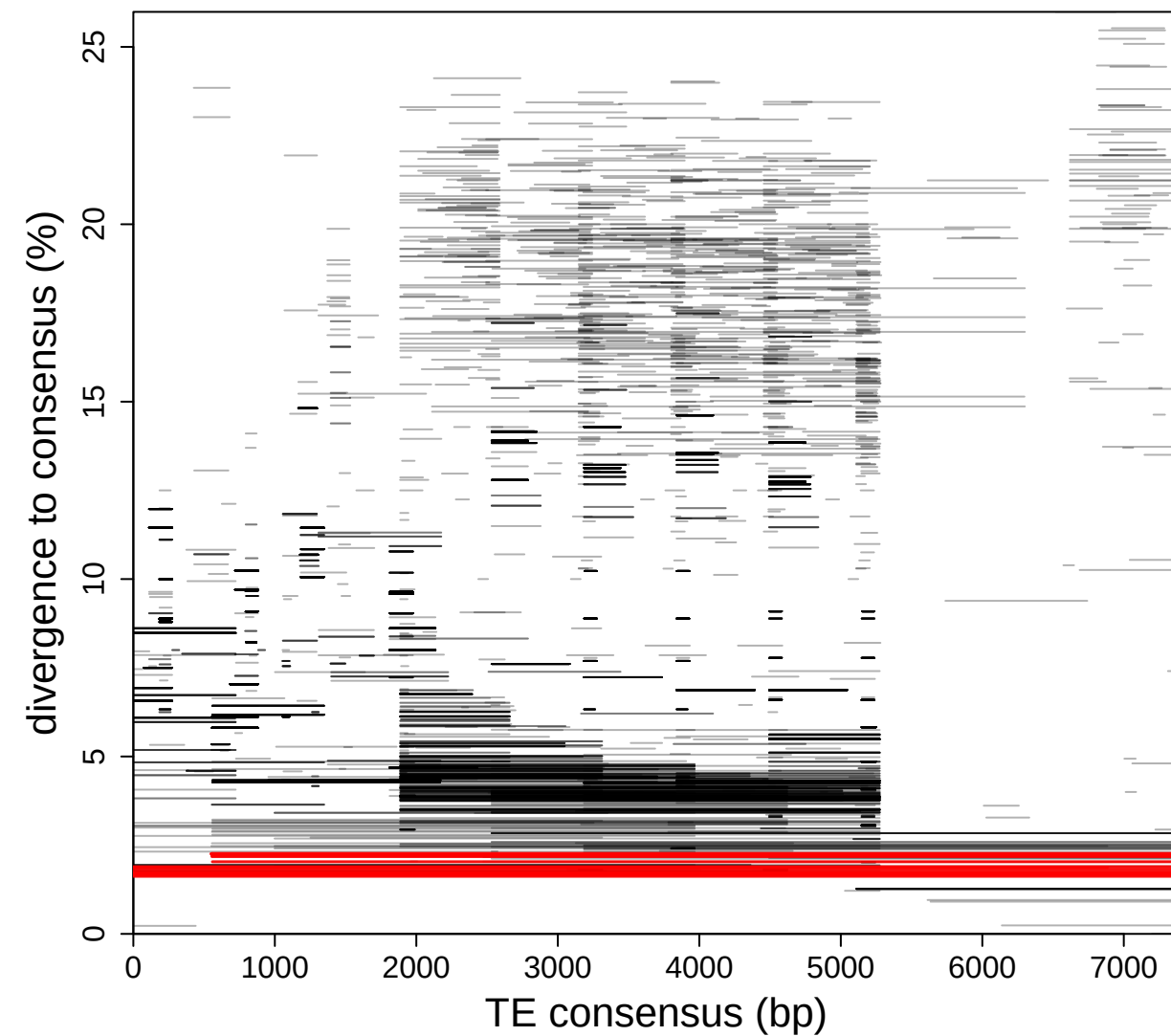
asta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_c
size: 15158bp; fragments: 9772; full length: 0 (>=13642.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

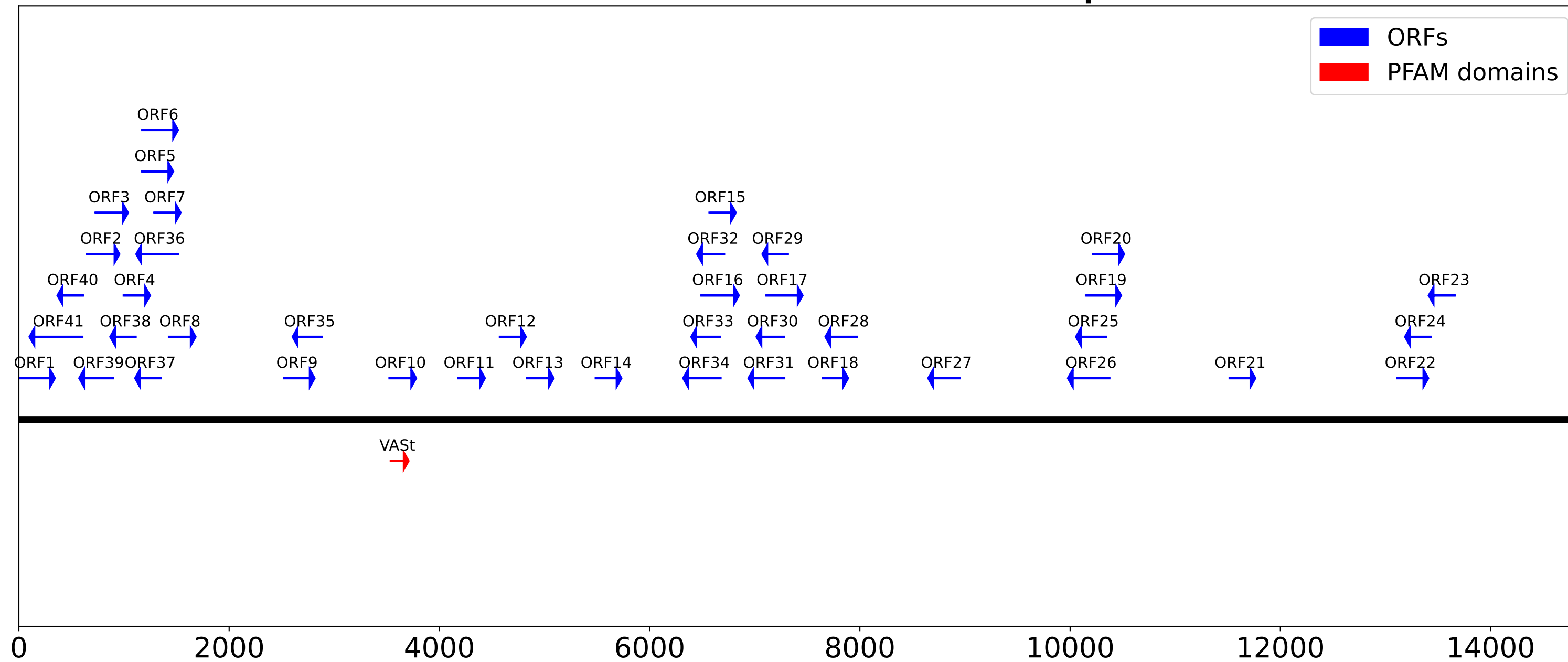


TE: rnd_5_family_4748
size: 7392bp; fragments: 5293; full length: 18 (≥ 6652.8 bp)

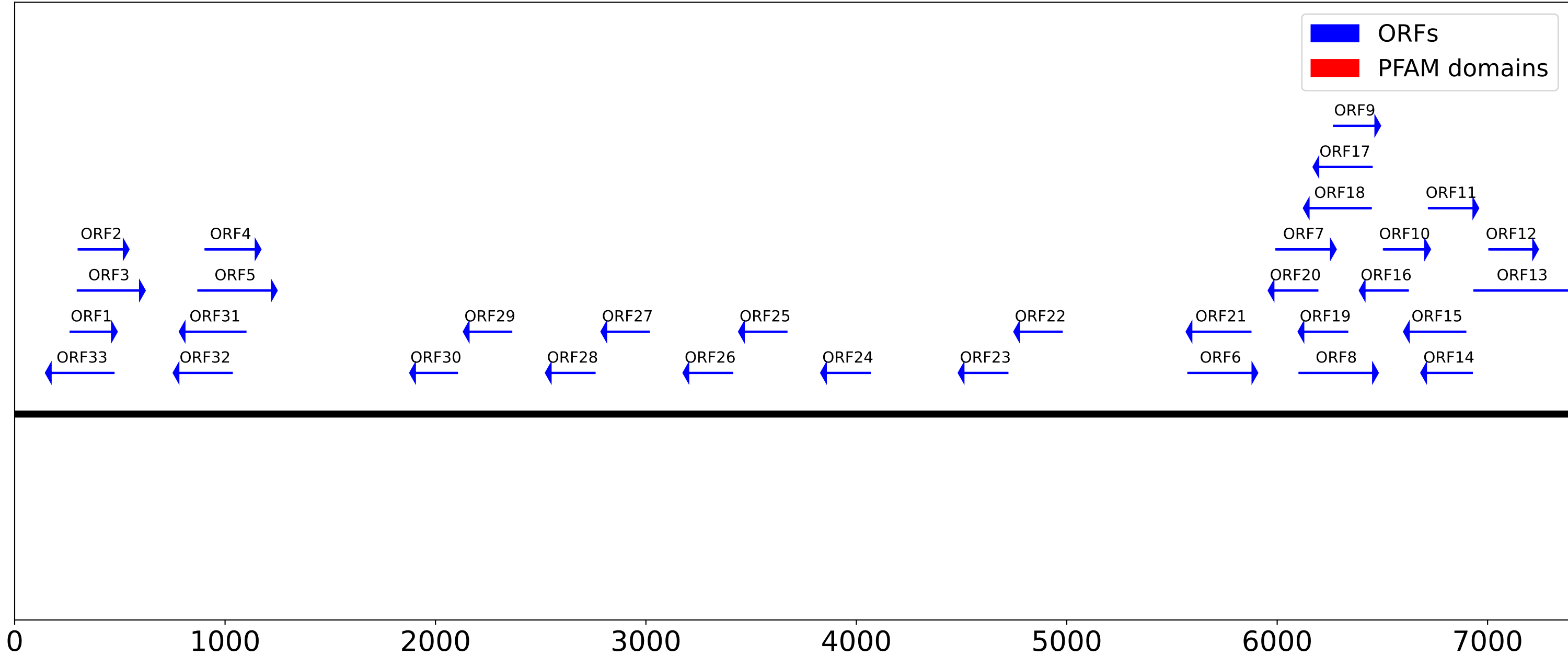
Before TEtrimmer 7392 bp



After TEtrimmer ORF and PFAM domain plot



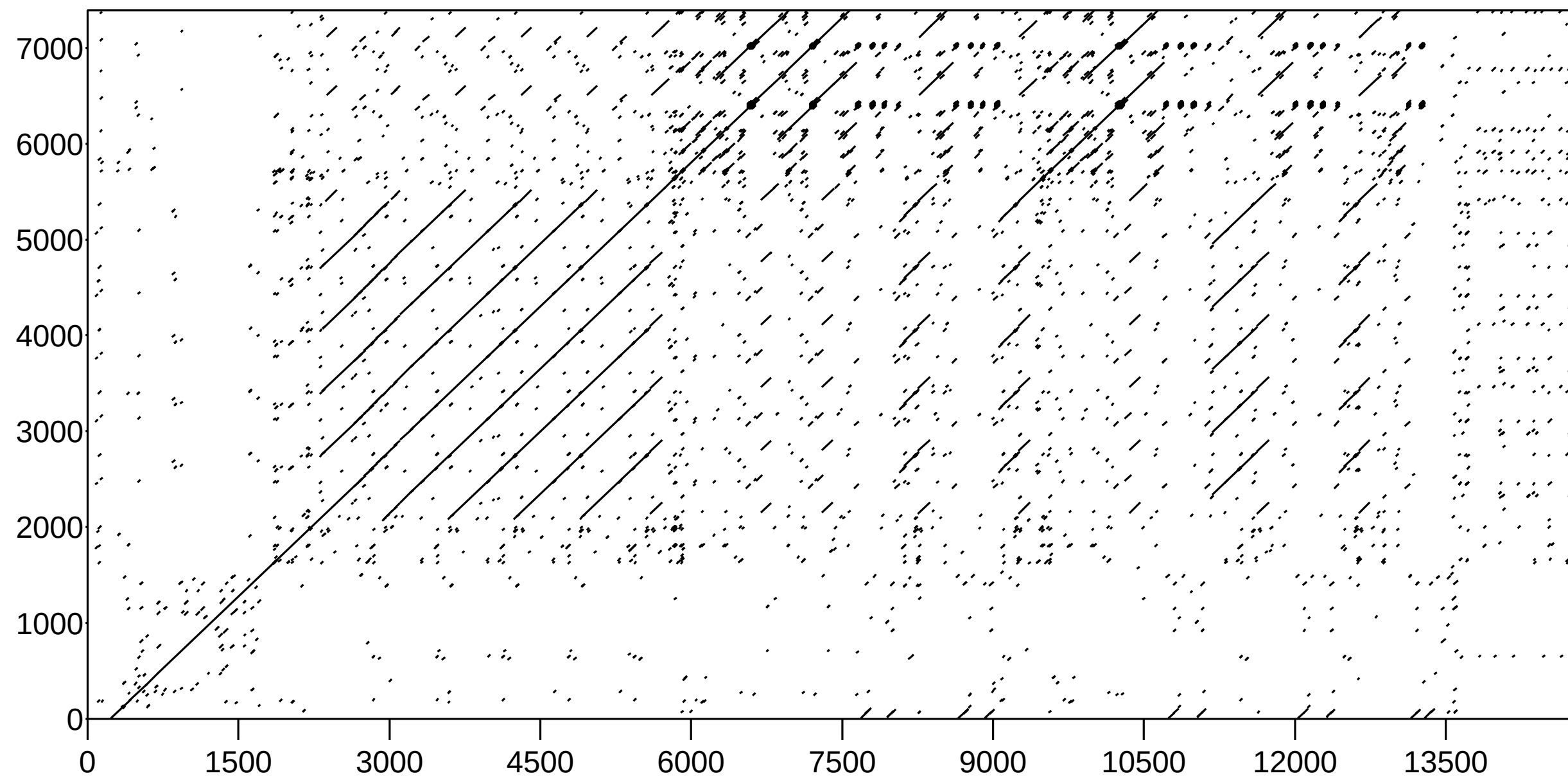
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)