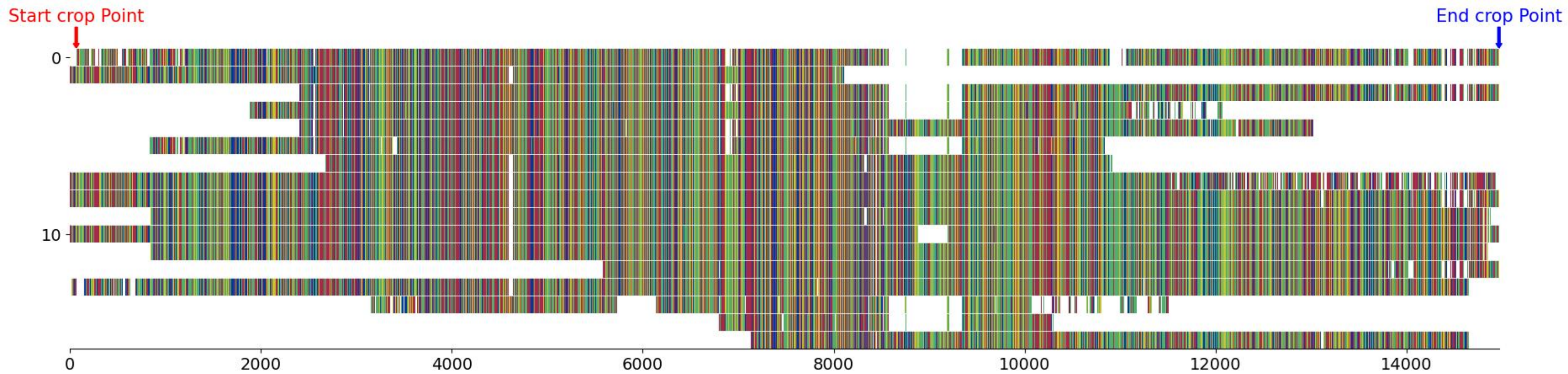


Start crop Point

End crop Point

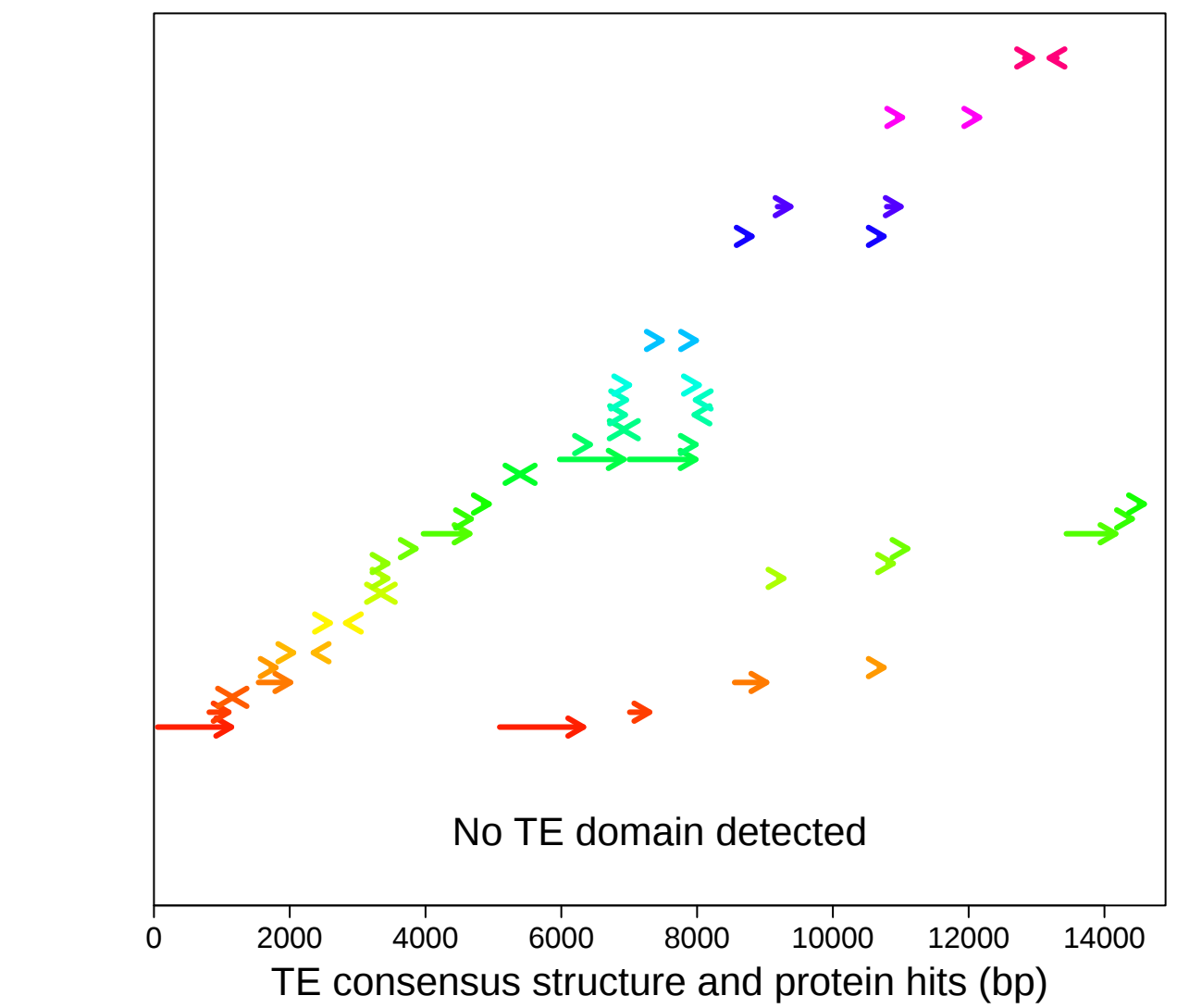
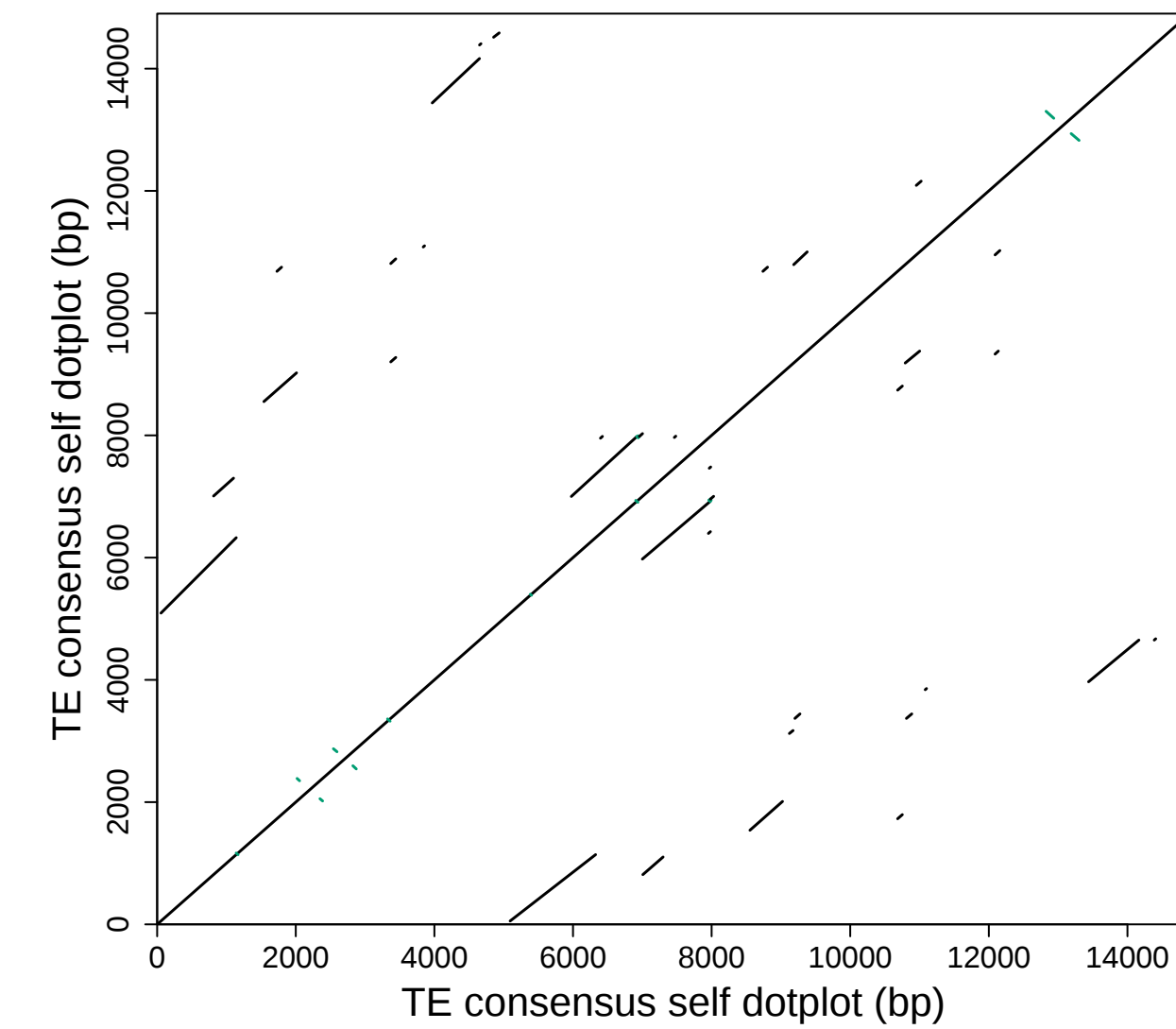
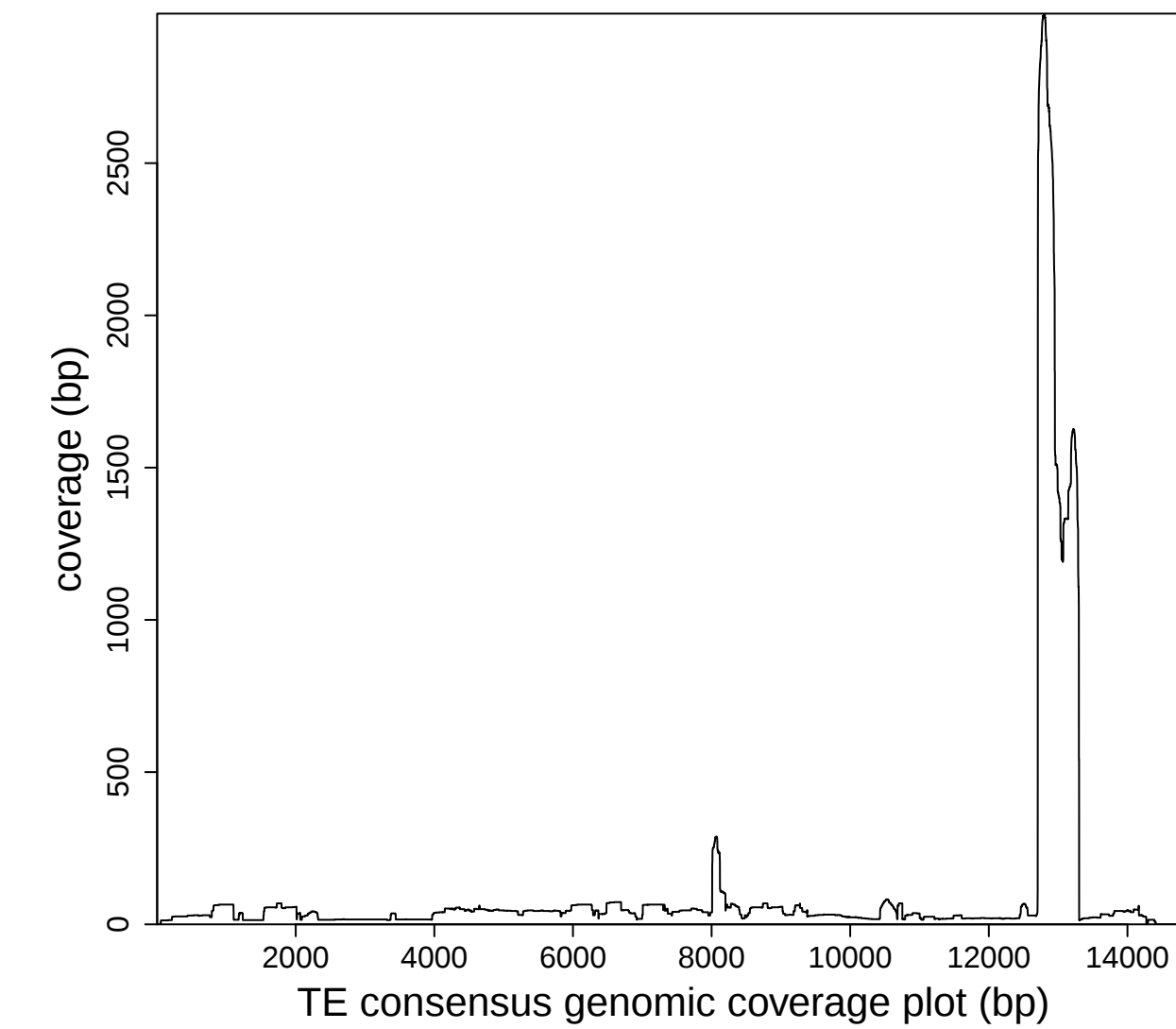
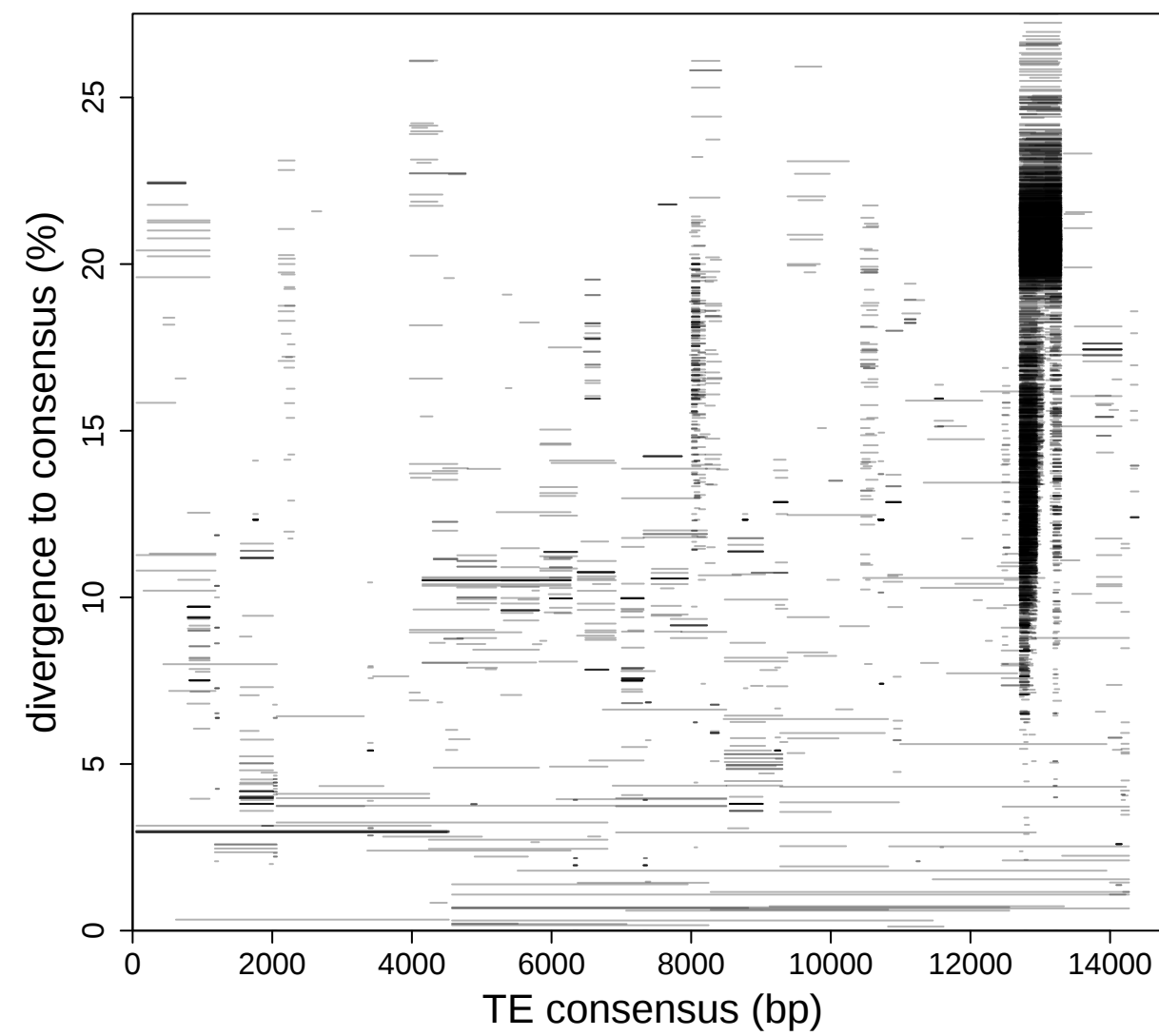
MSA length = 14900





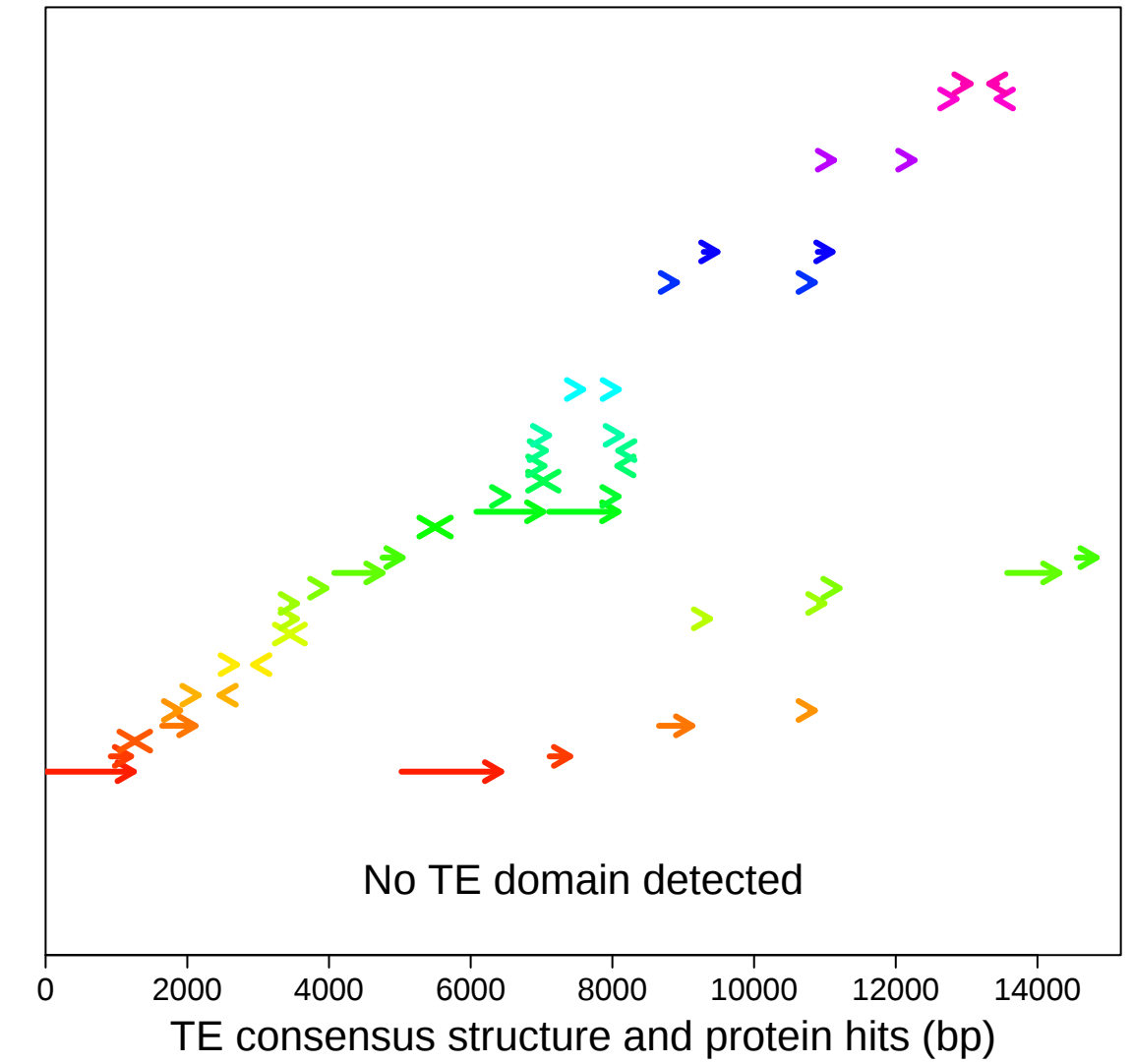
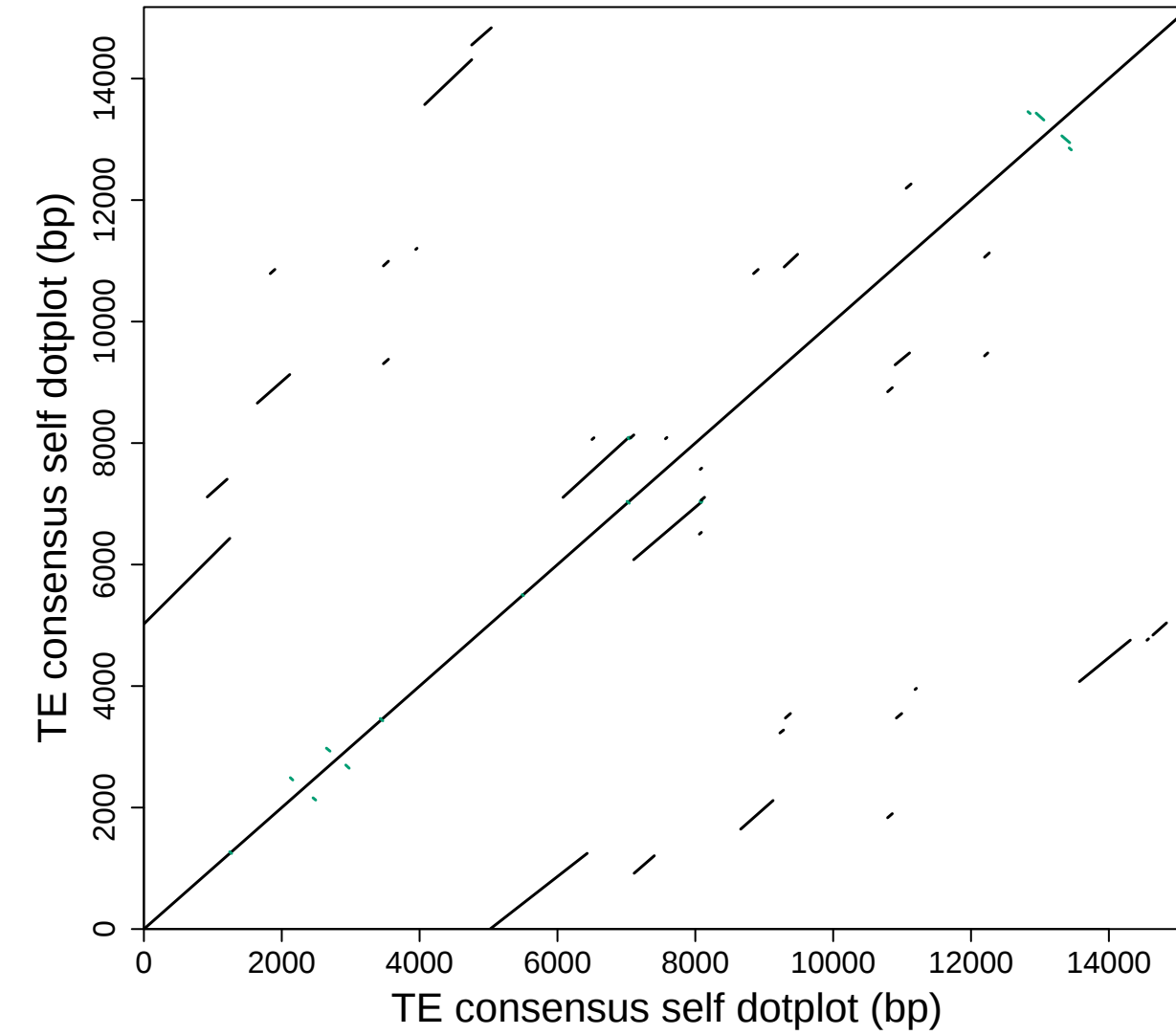
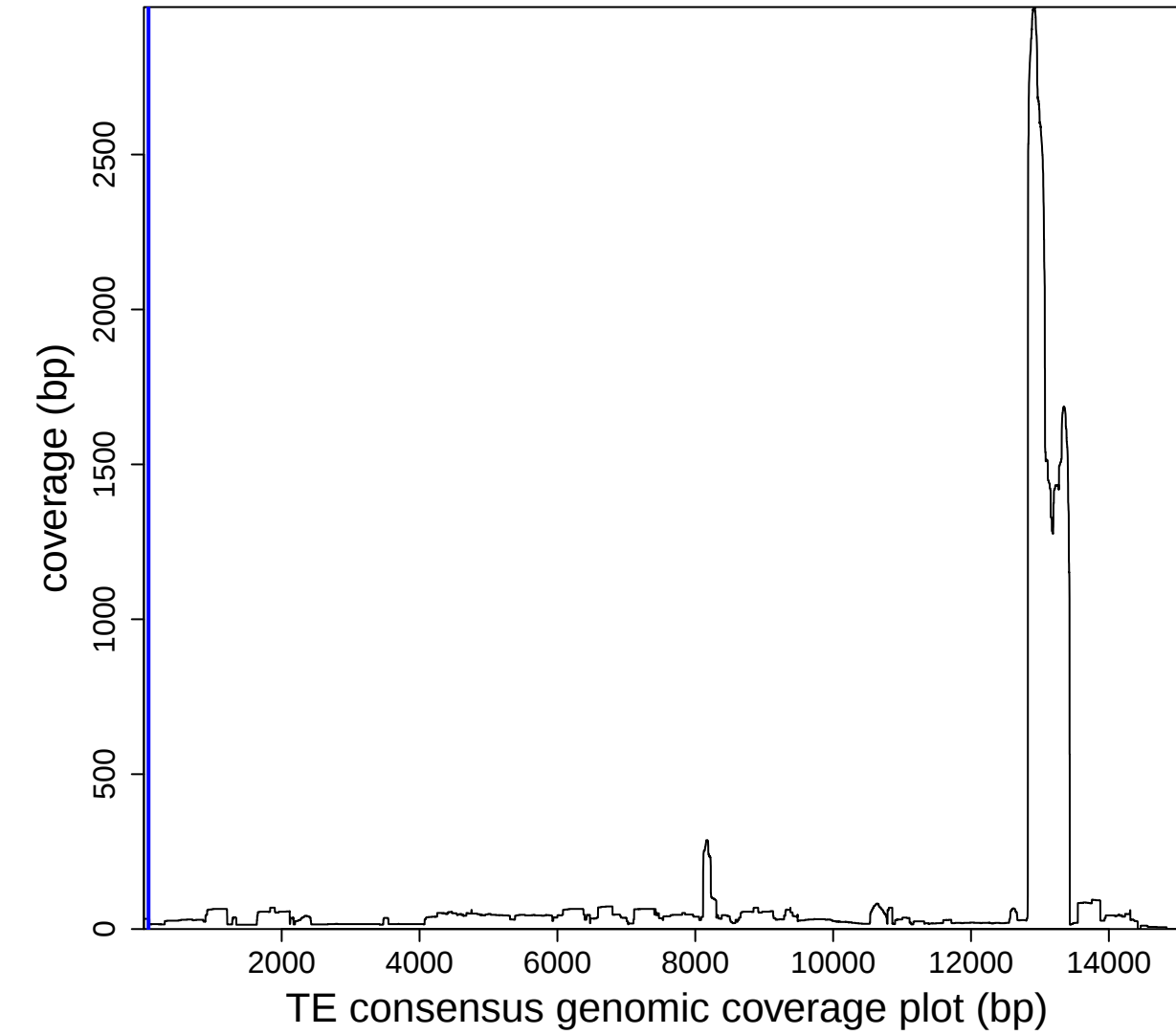
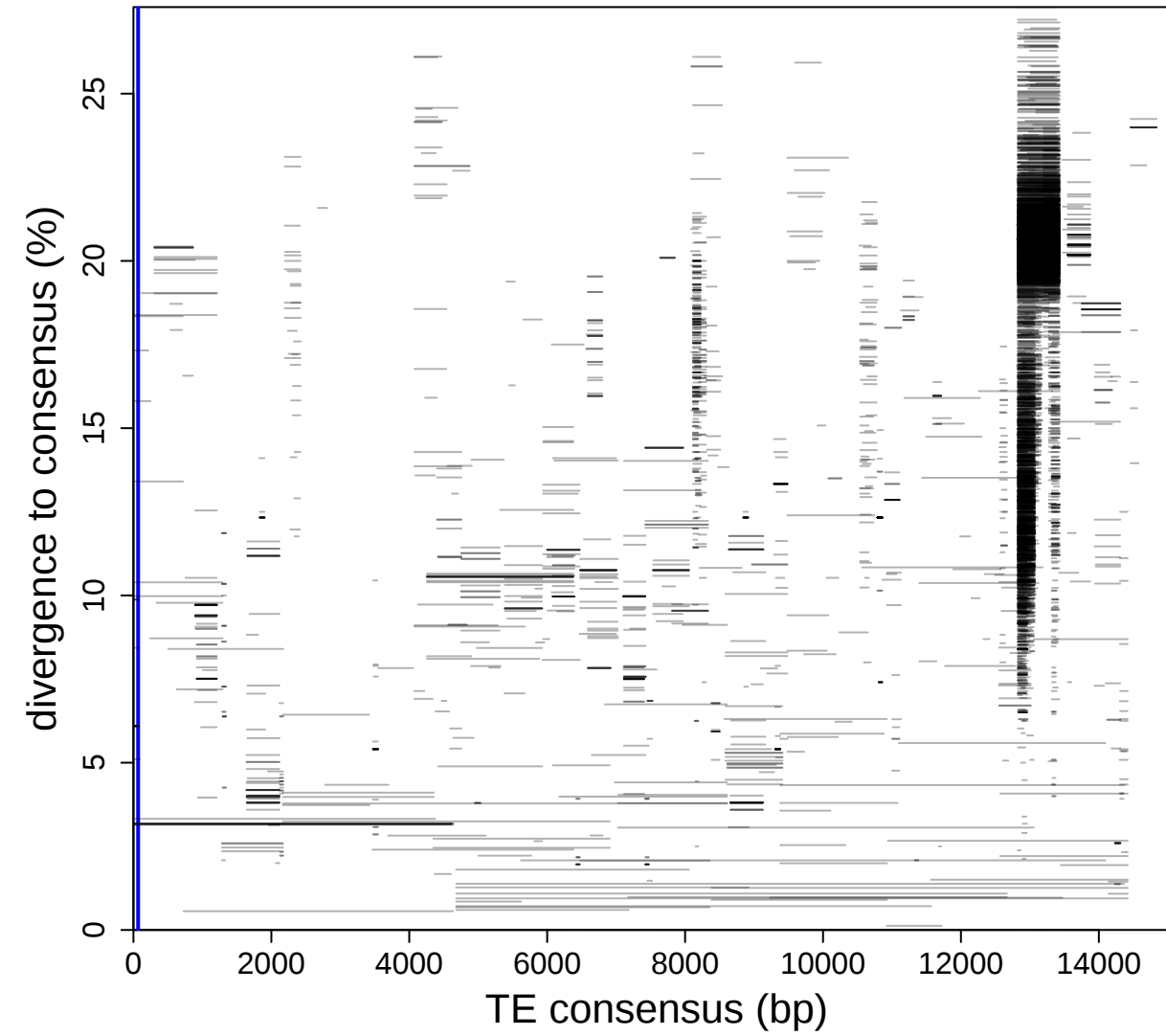
l.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 14900bp; fragments: 4876; full length: 0 (>=13410bp)

After TEtrimmer 14900 bp

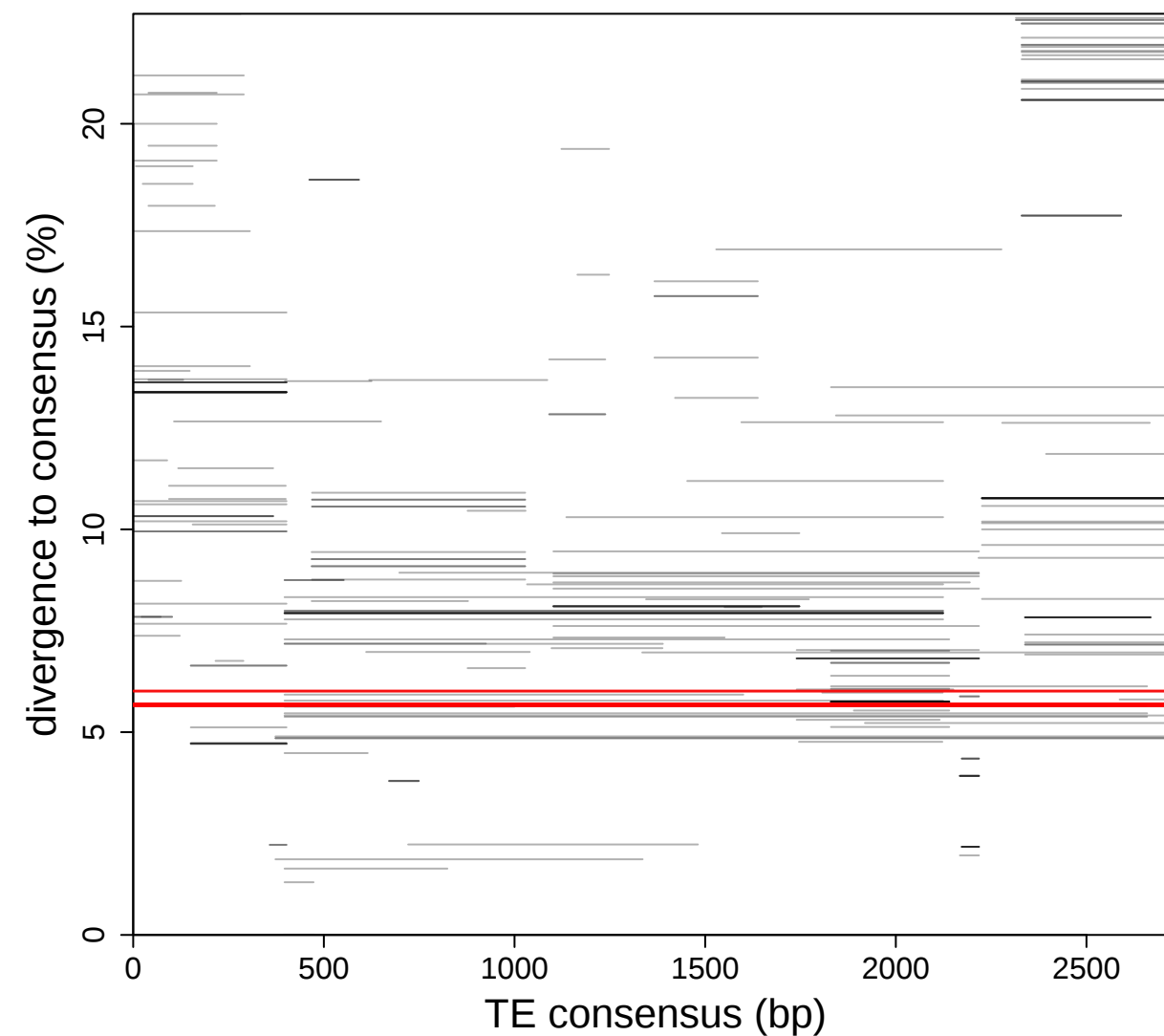


2.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 15177bp; fragments: 4906; full length: 0 (>=13659.3bp)

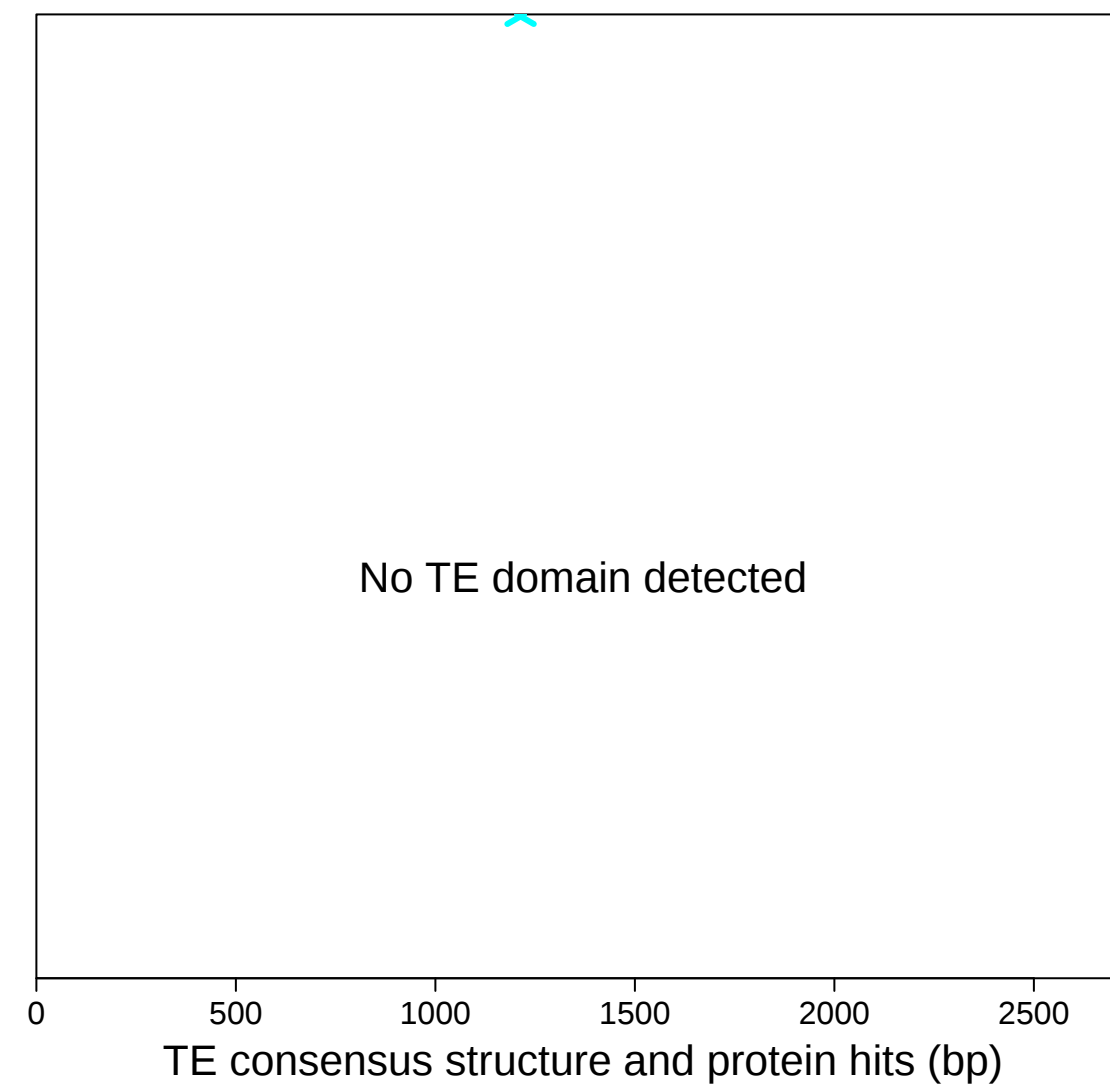
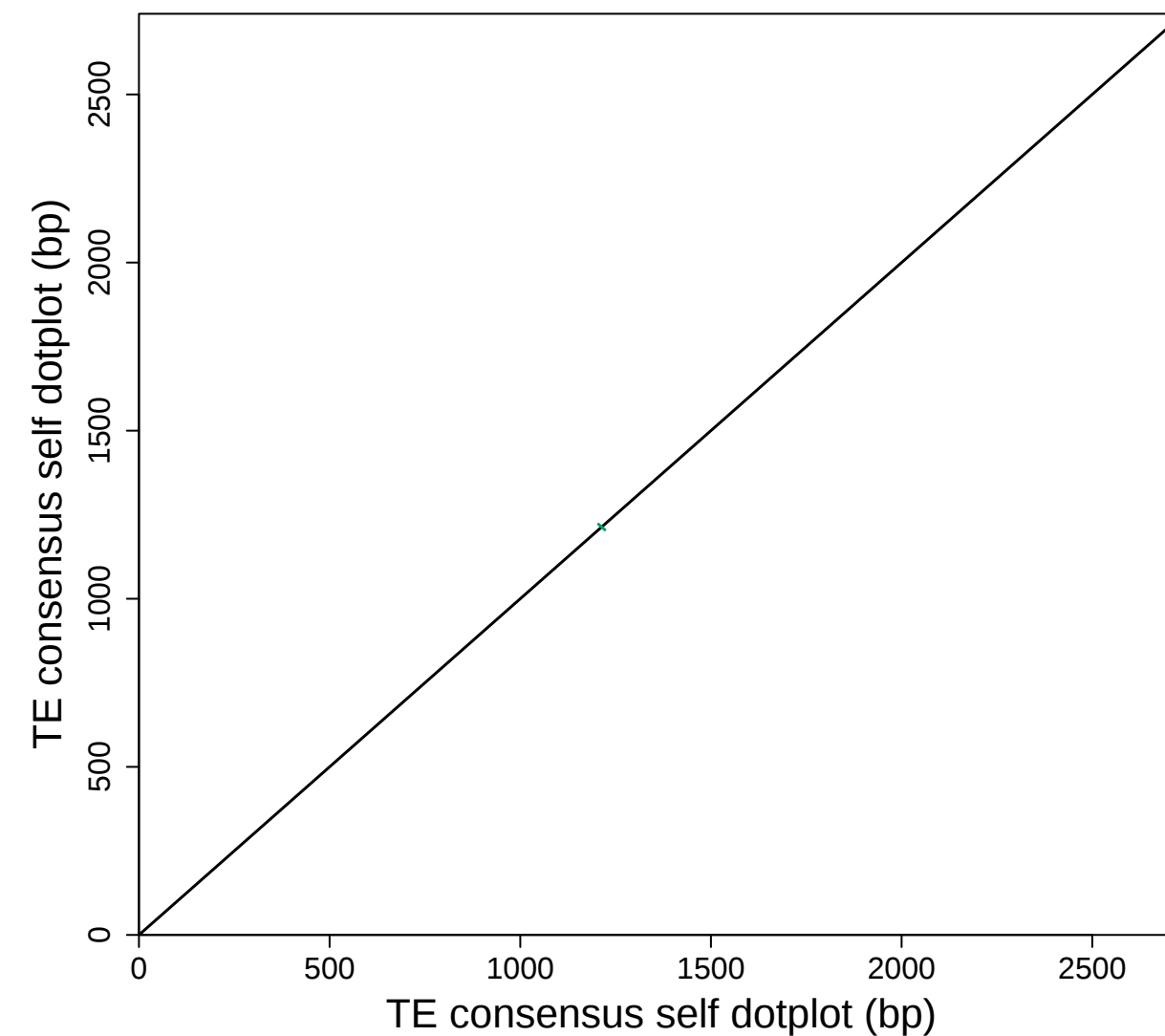
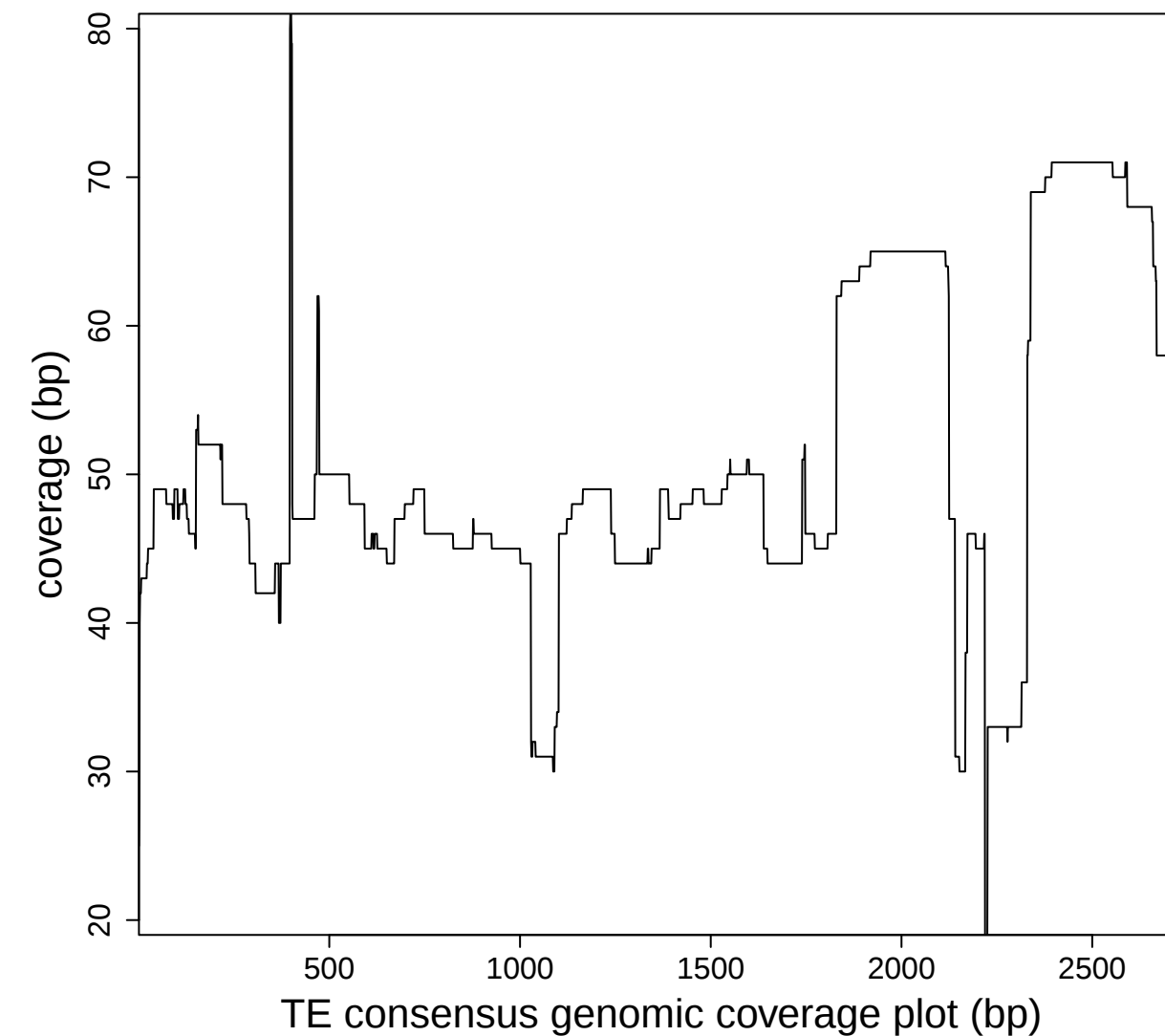
After TEtrimmer Extended plot Blue lines are boundaries



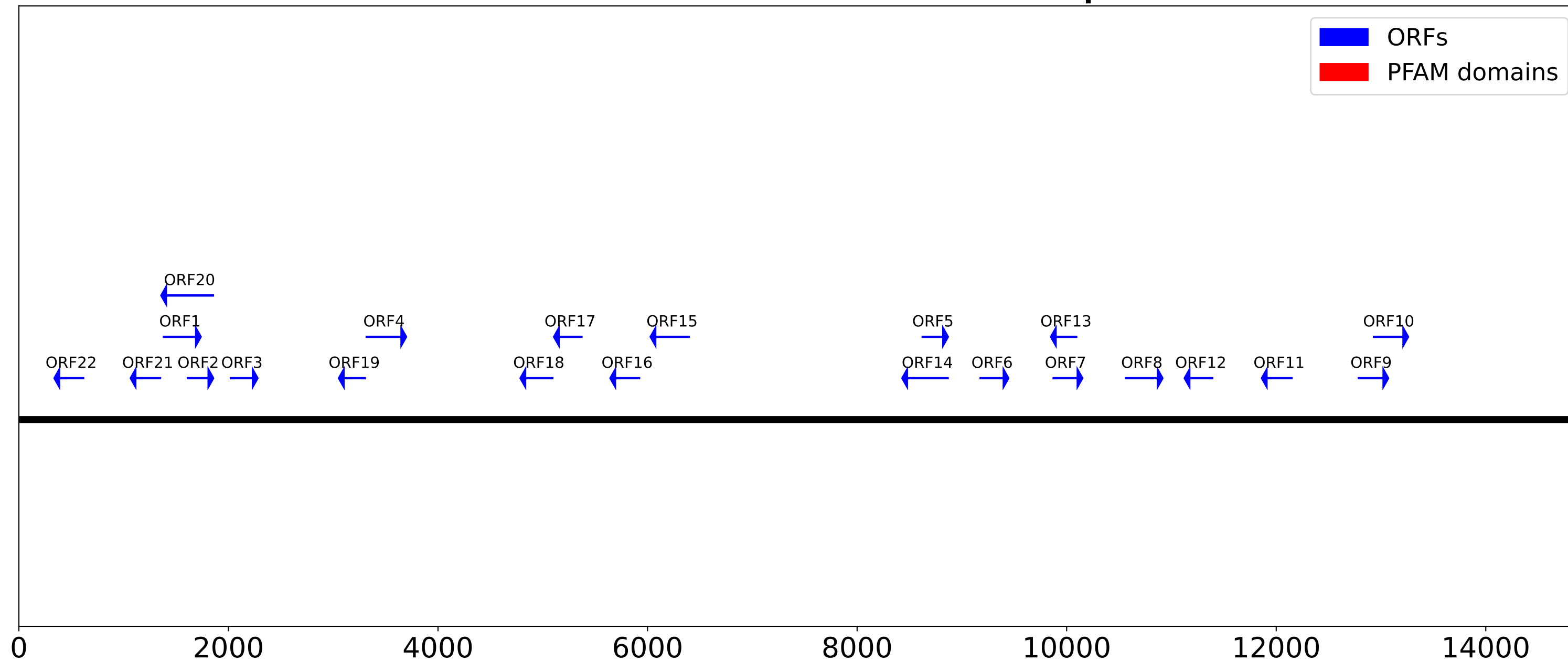
TE: rnd_5_family_752
size: 2740bp; fragments: 261; full length: 5 (≥ 2466 bp)



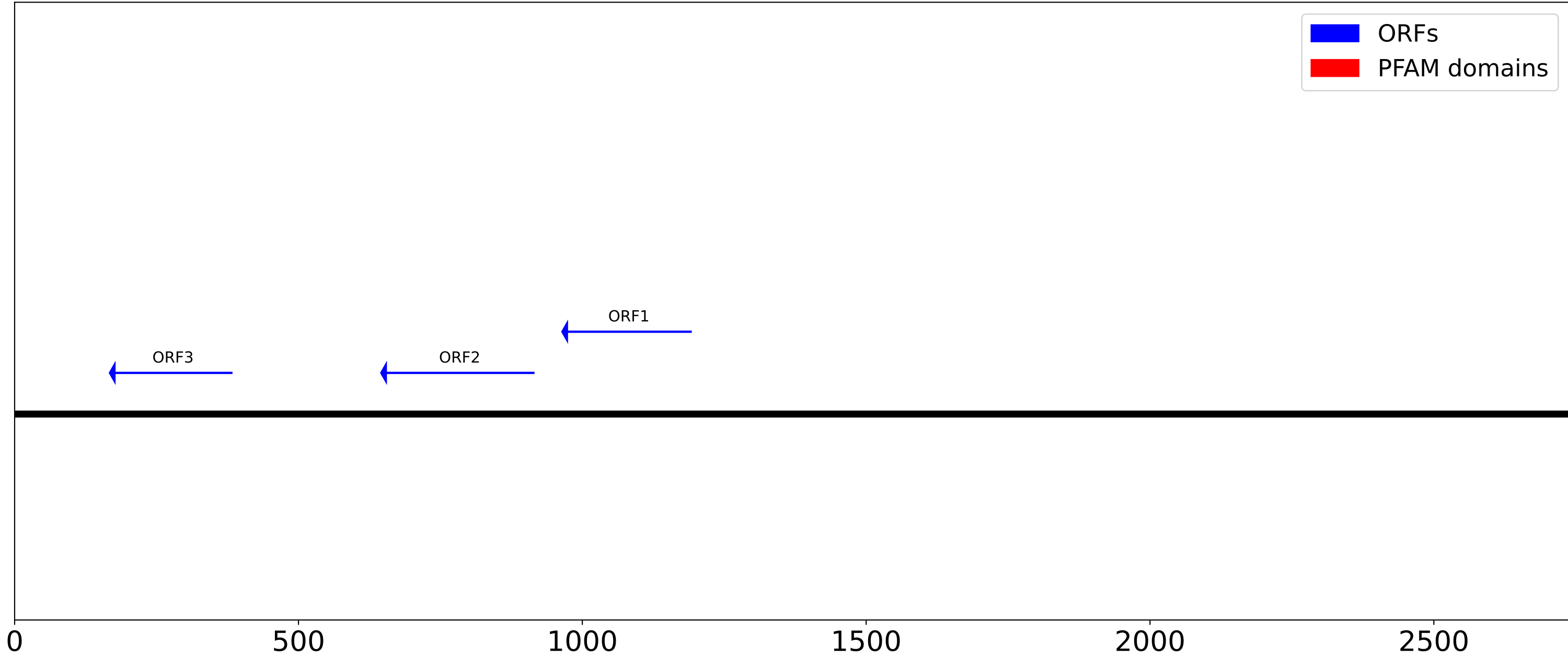
Before TEtrimmer 2740 bp



After TEtrimmer ORF and PFAM domain plot



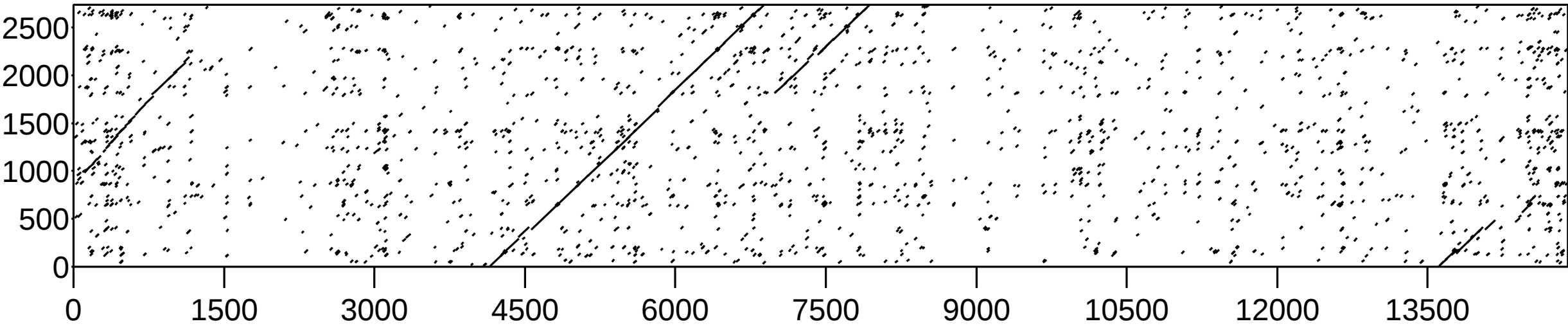
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)