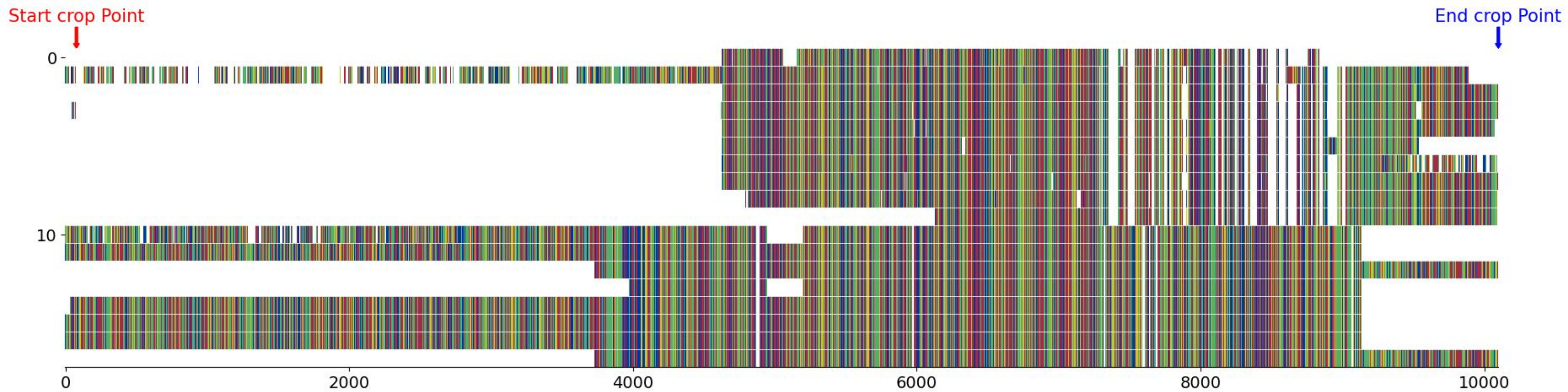


Start crop Point

End crop Point

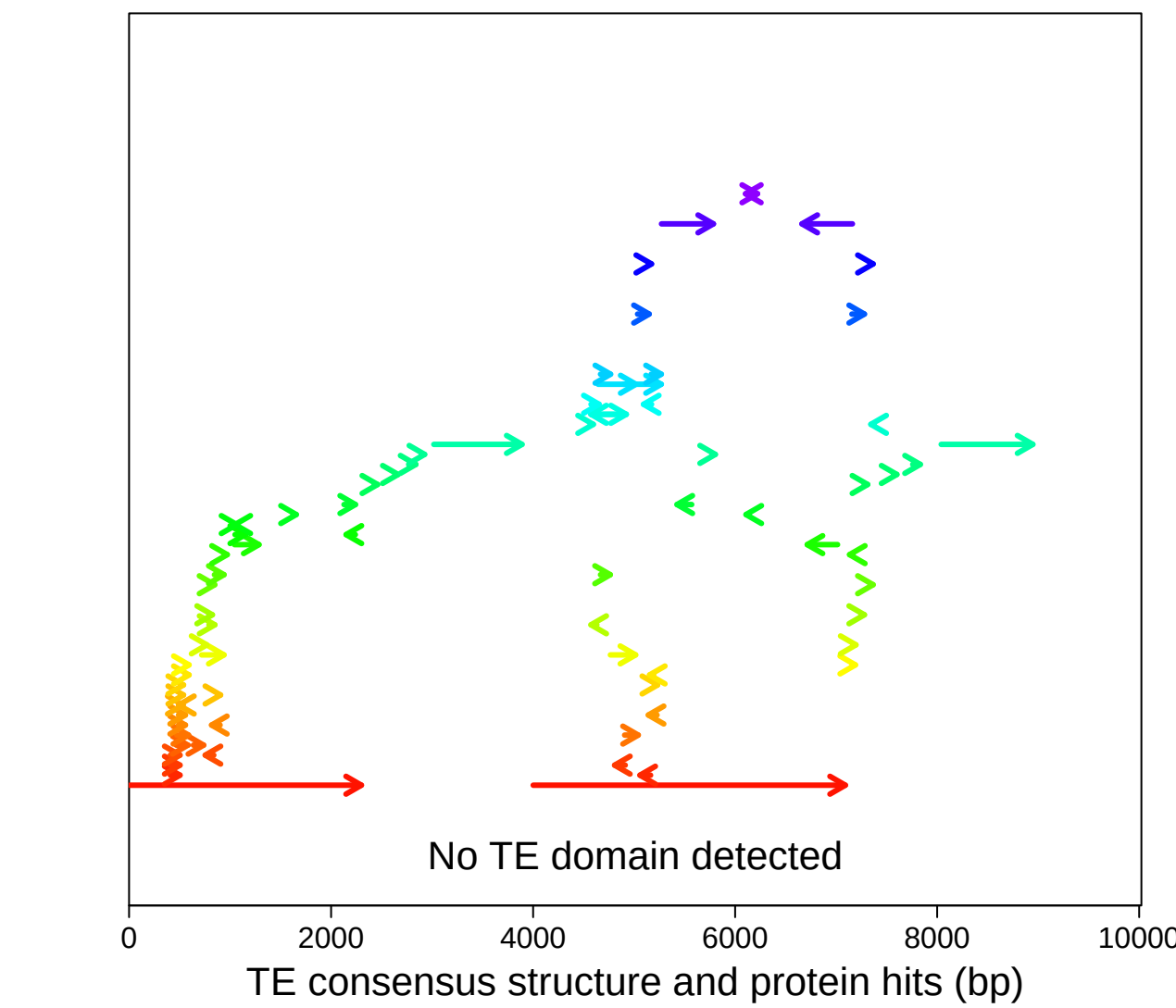
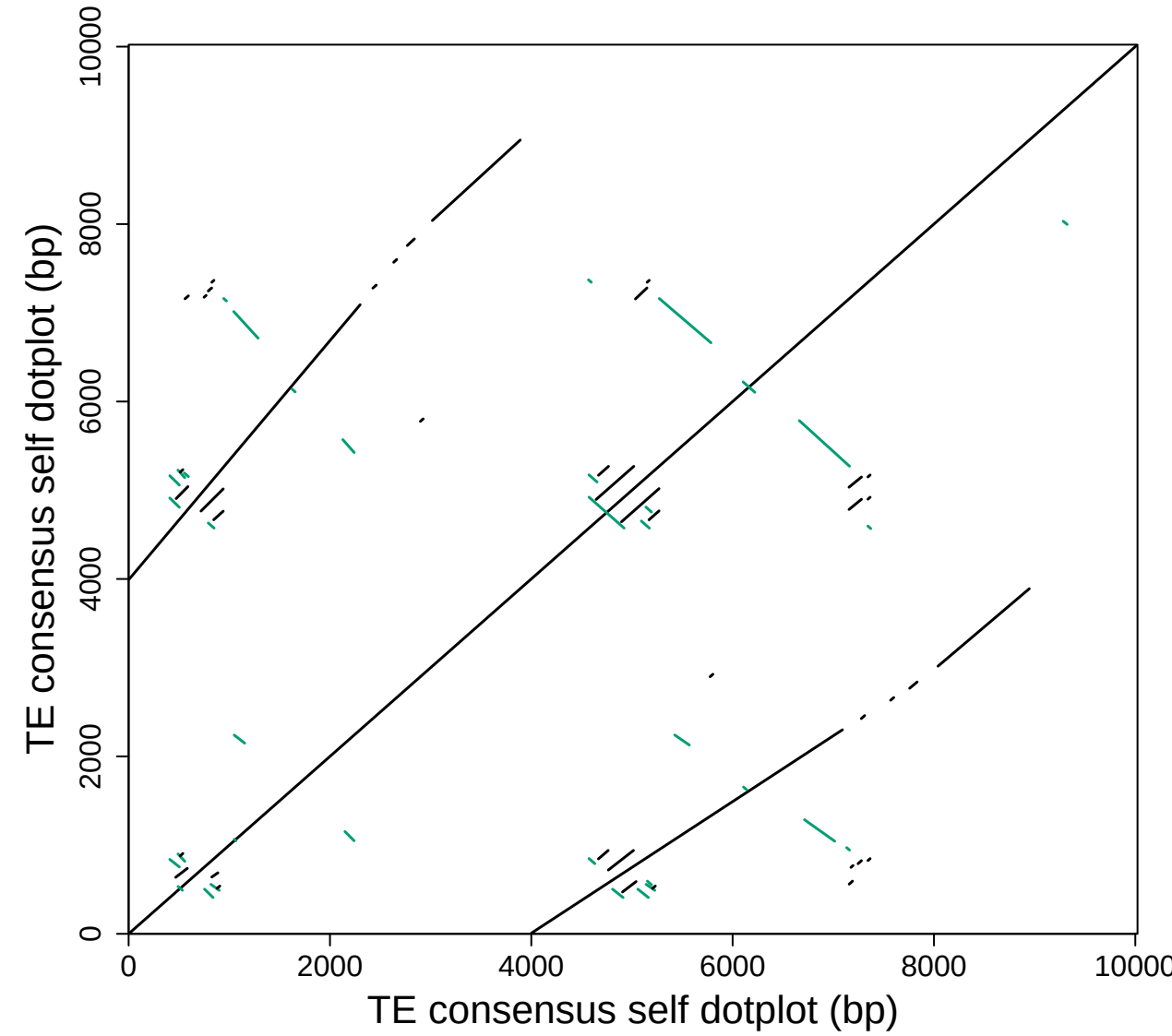
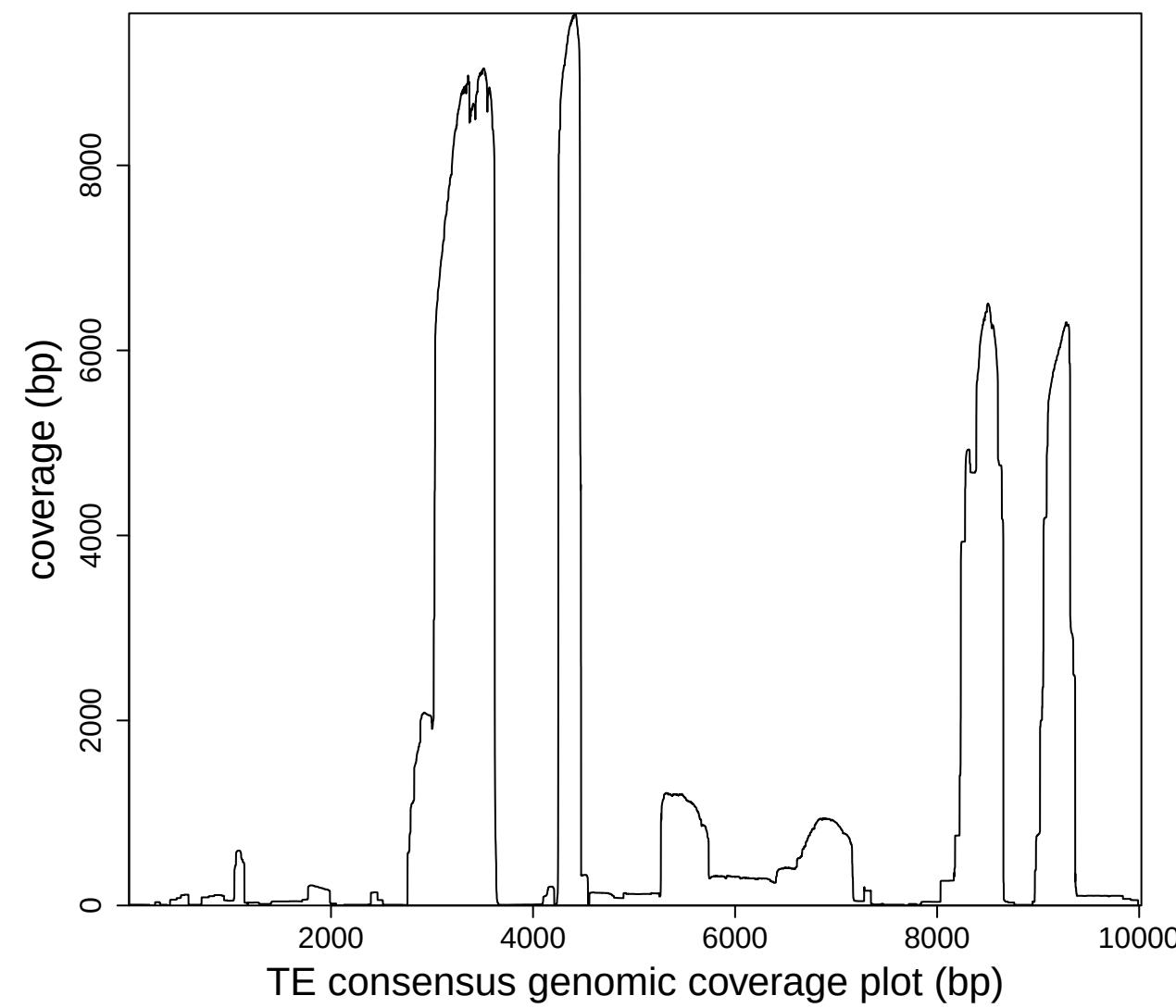
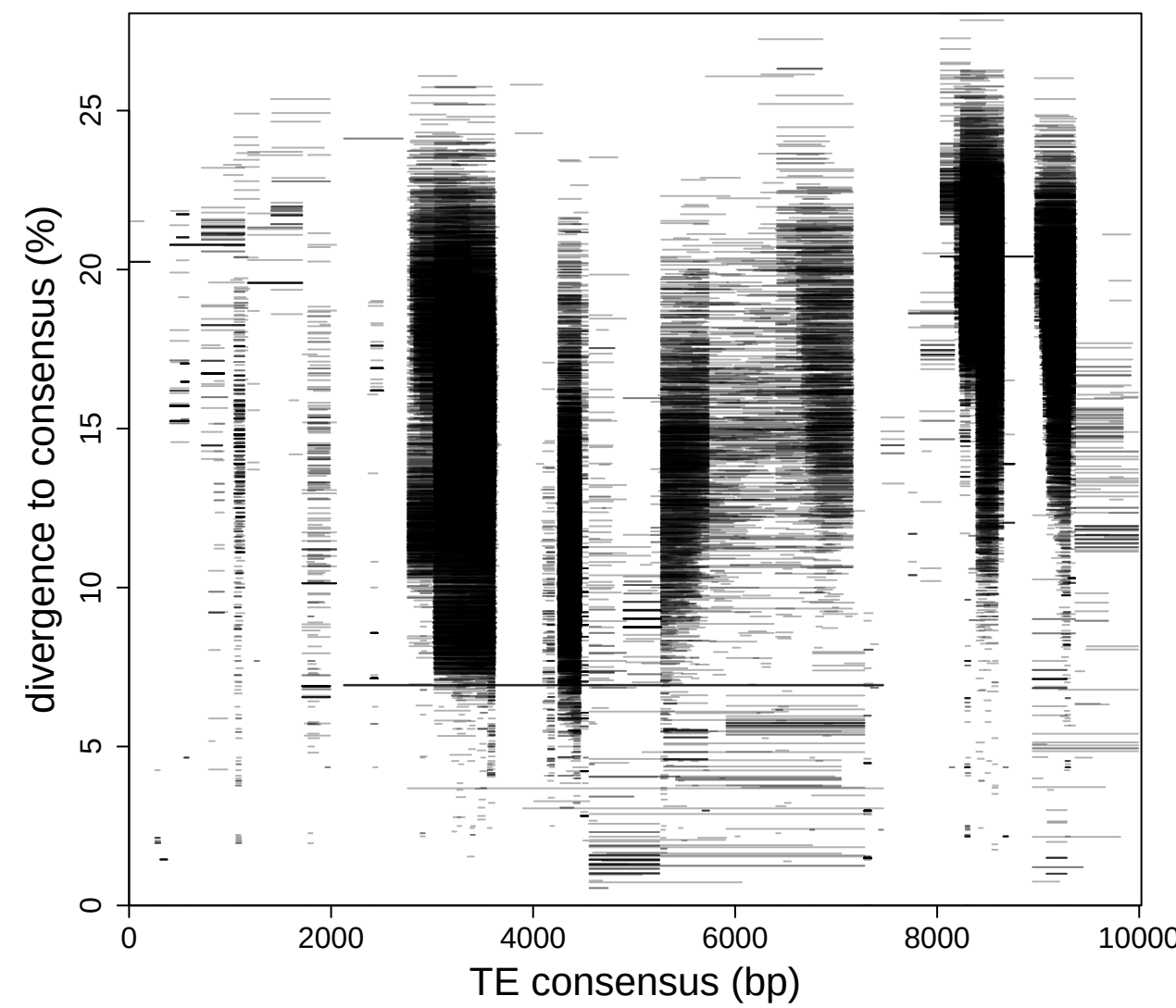
MSA length = 10022





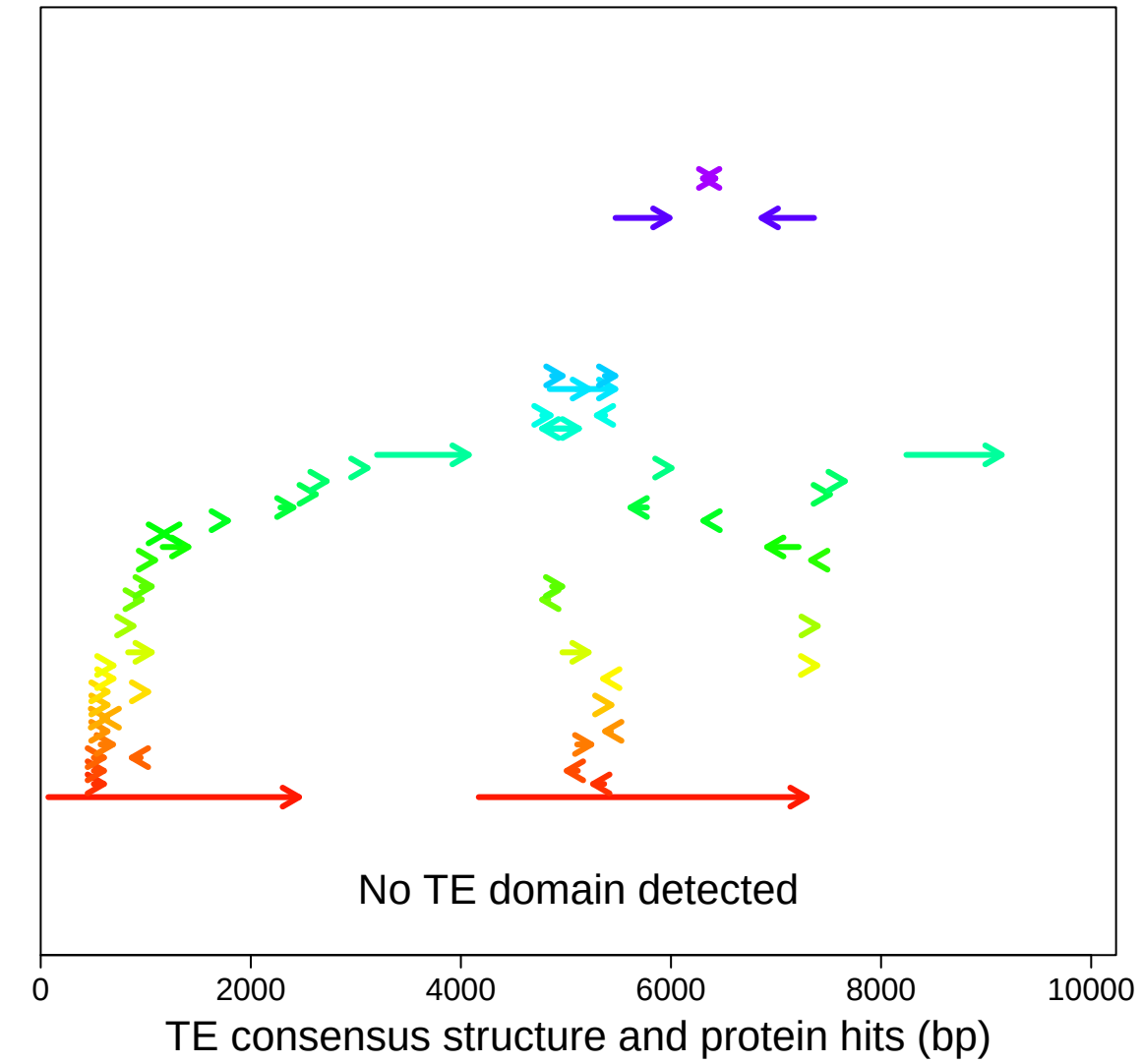
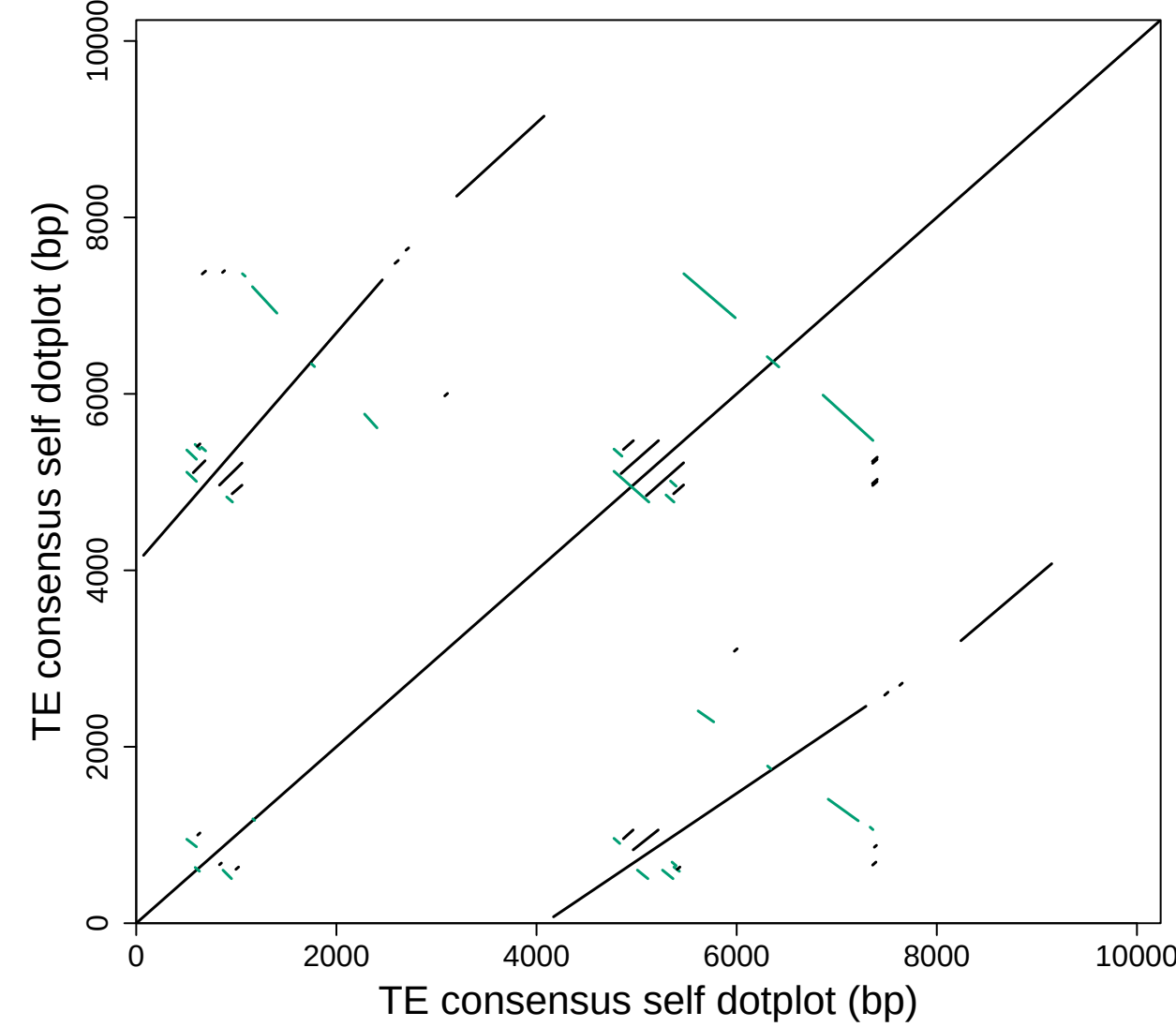
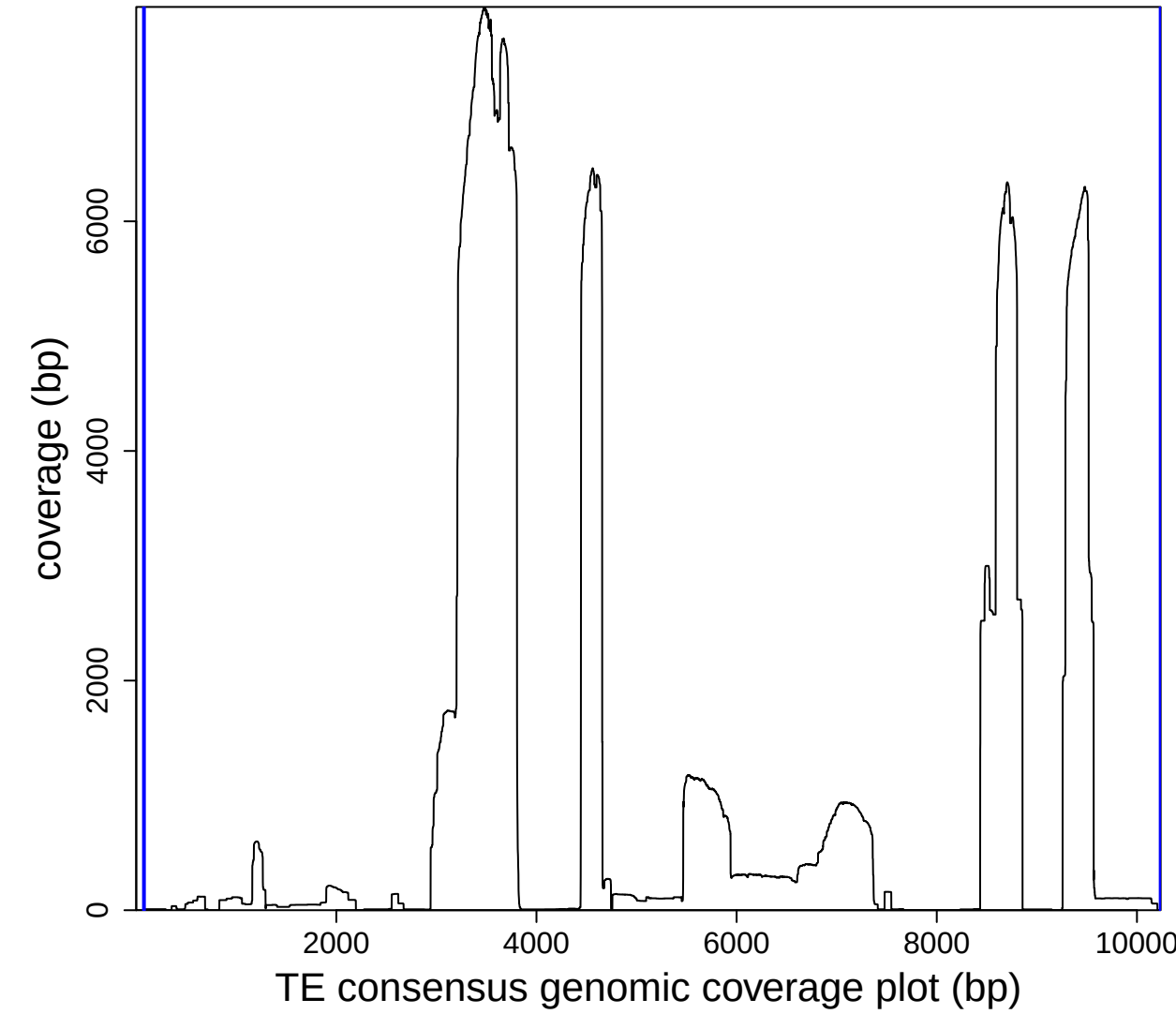
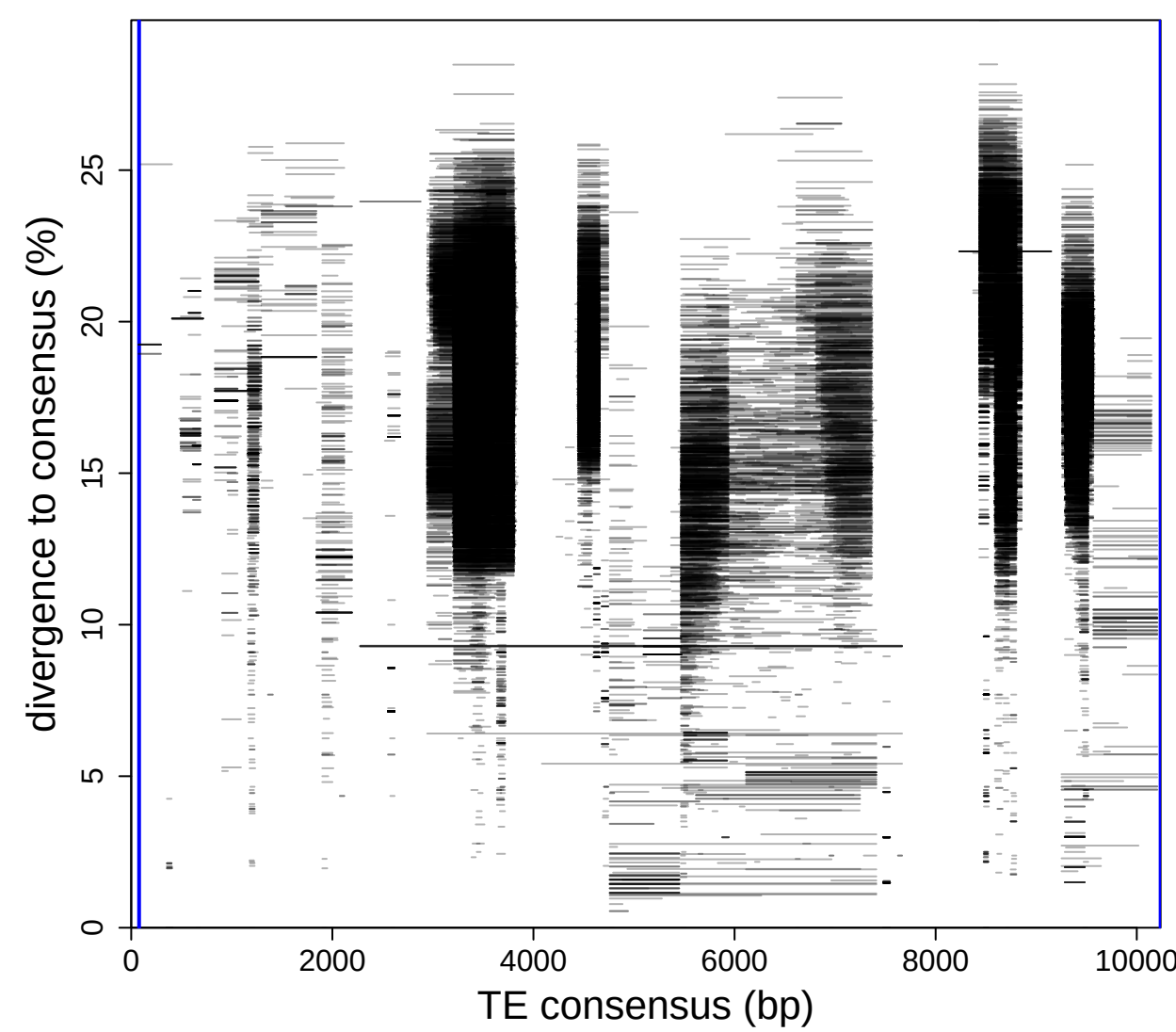
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 10022bp; fragments: 39767; full length: 0 (>=9019.8bp)

After TEtrimmer 10022 bp



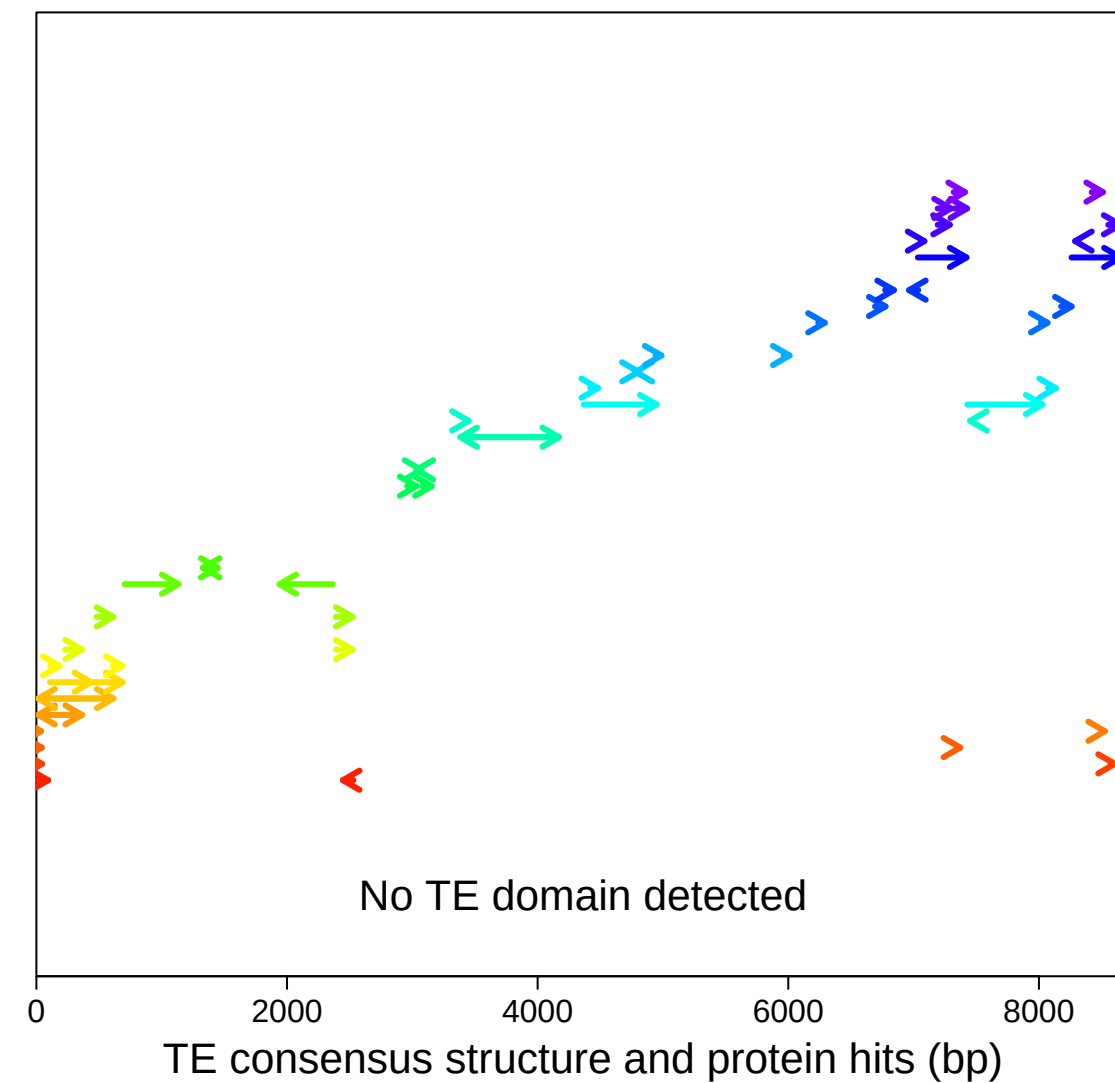
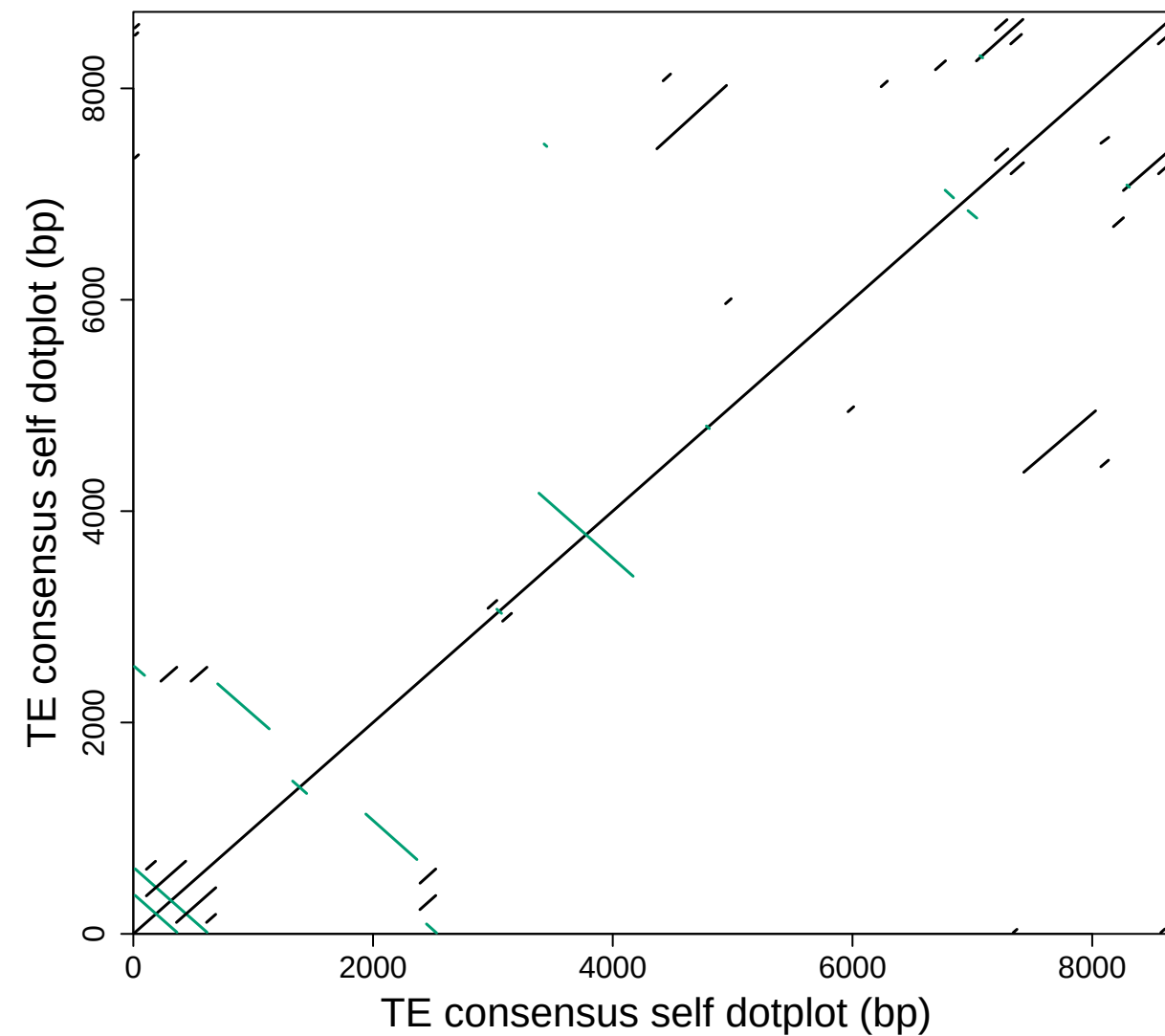
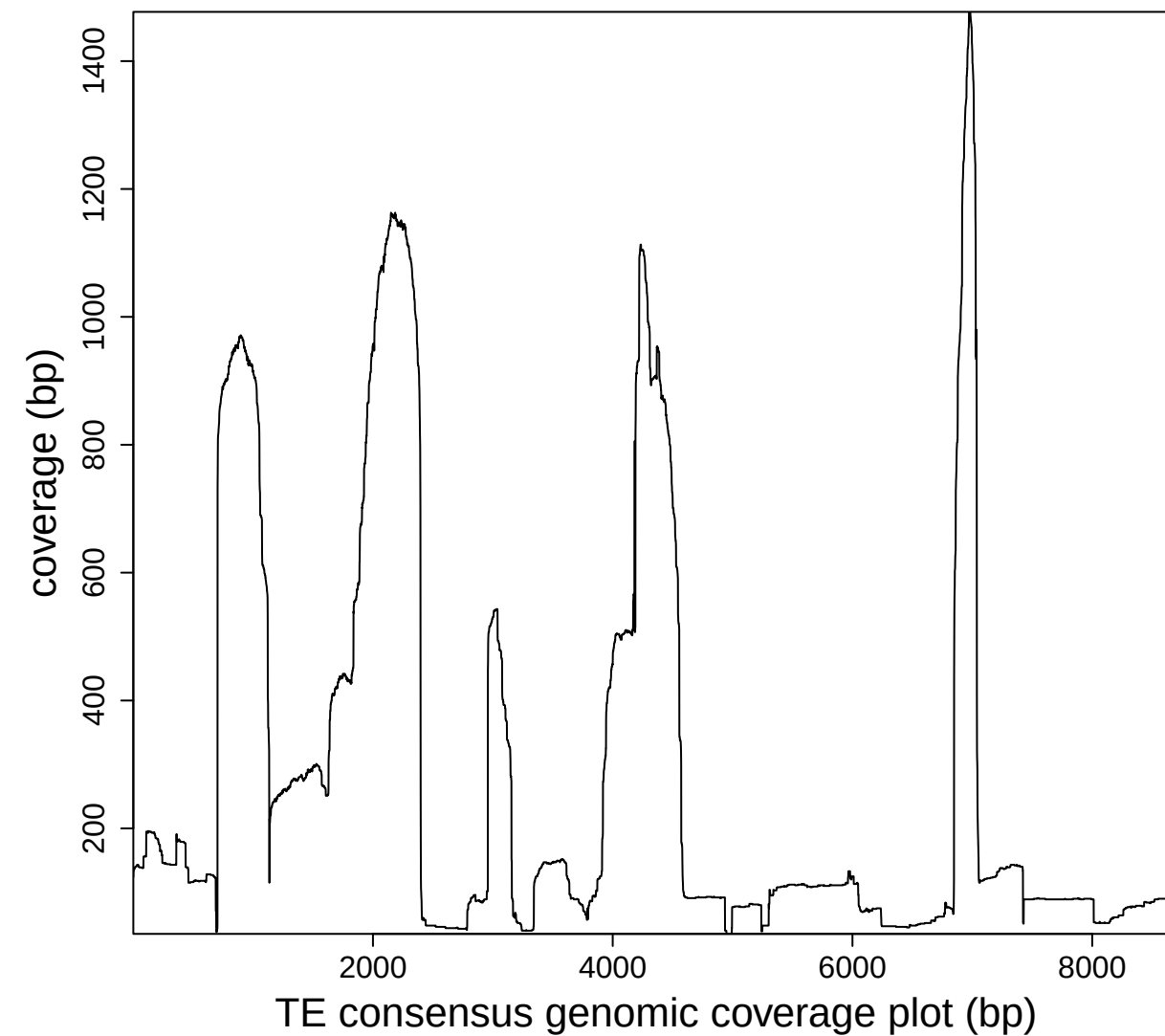
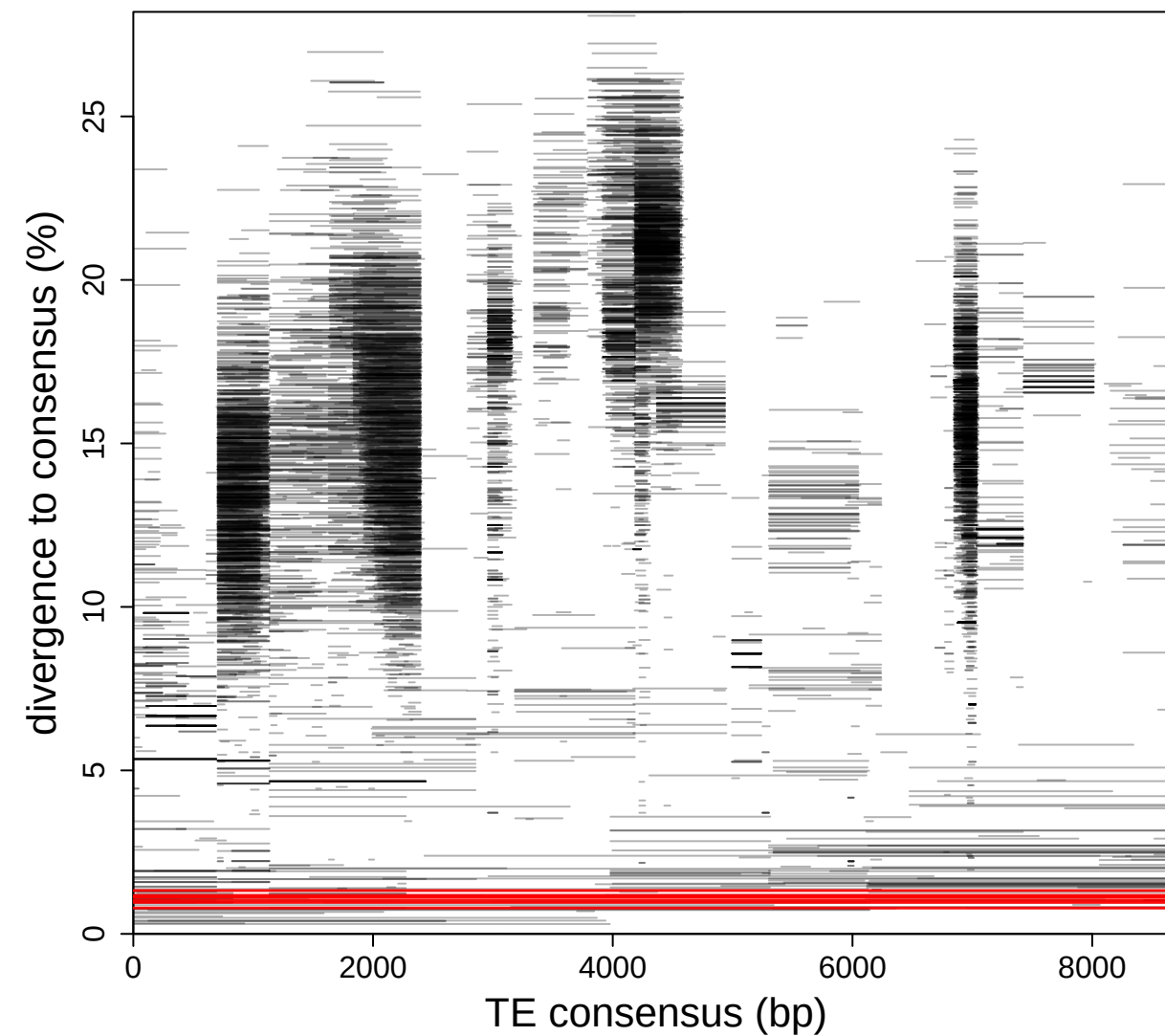
2.fasta.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 10237bp; fragments: 33888; full length: 0 (>=9213.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

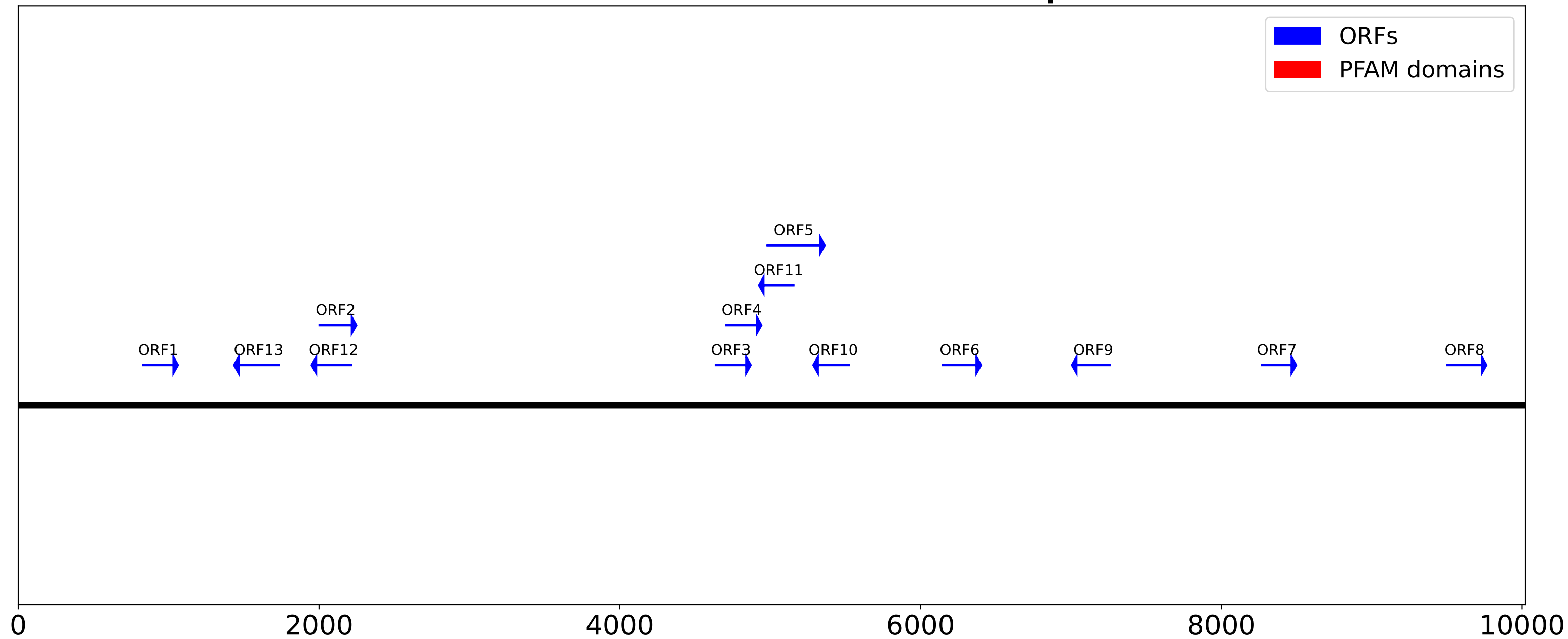


TE: rnd_5_family_2772
size: 8724bp; fragments: 7111; full length: 7 (≥ 7851.6 bp)

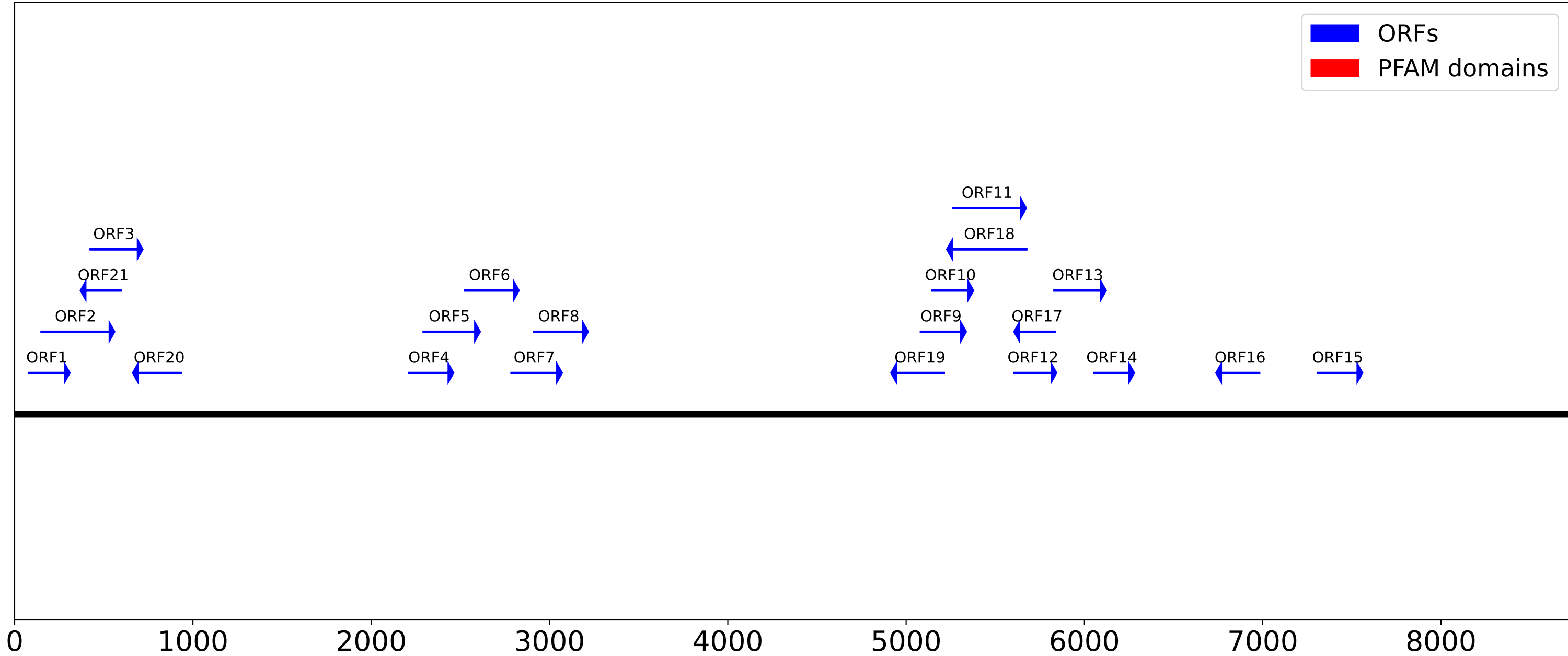
Before TEtrimmer 8724 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

