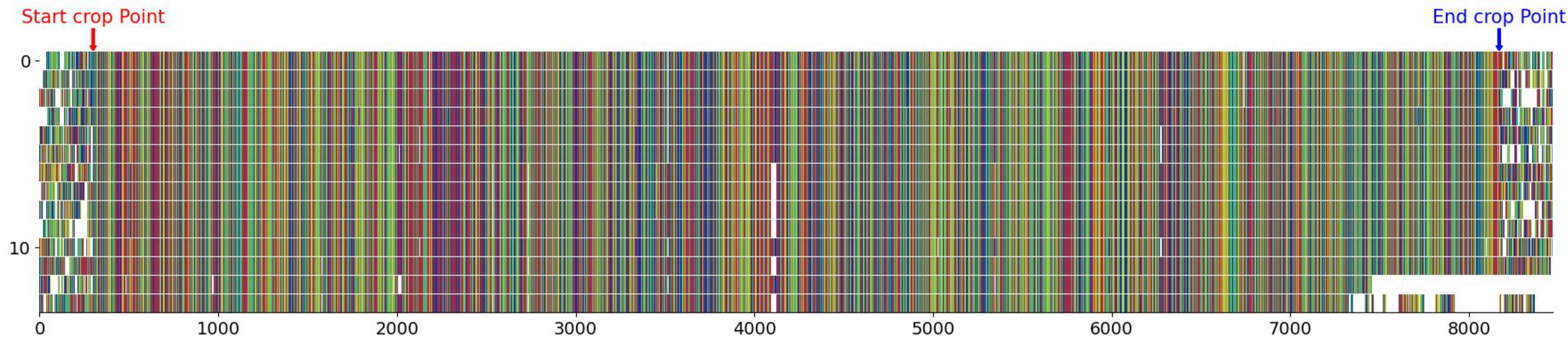


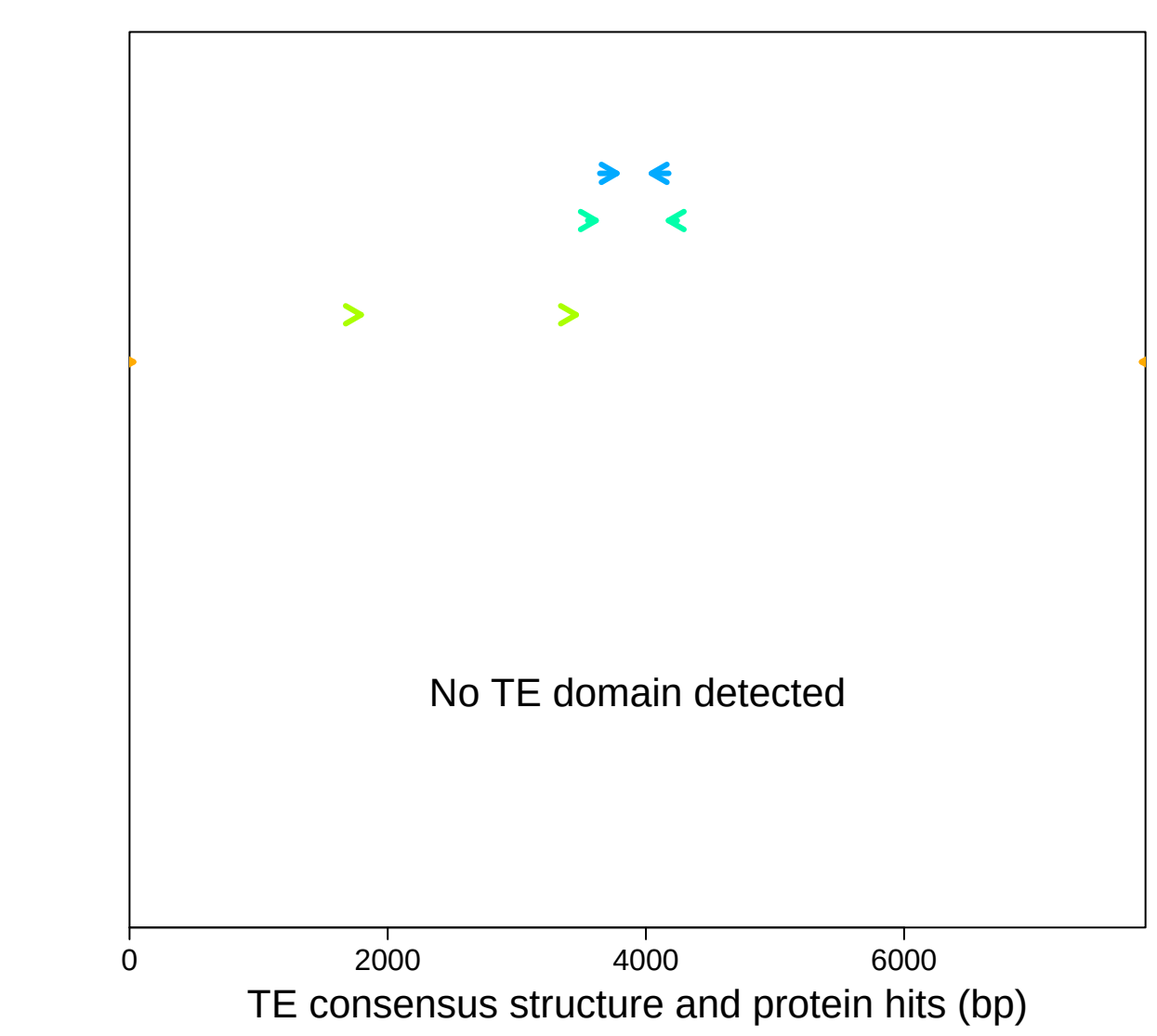
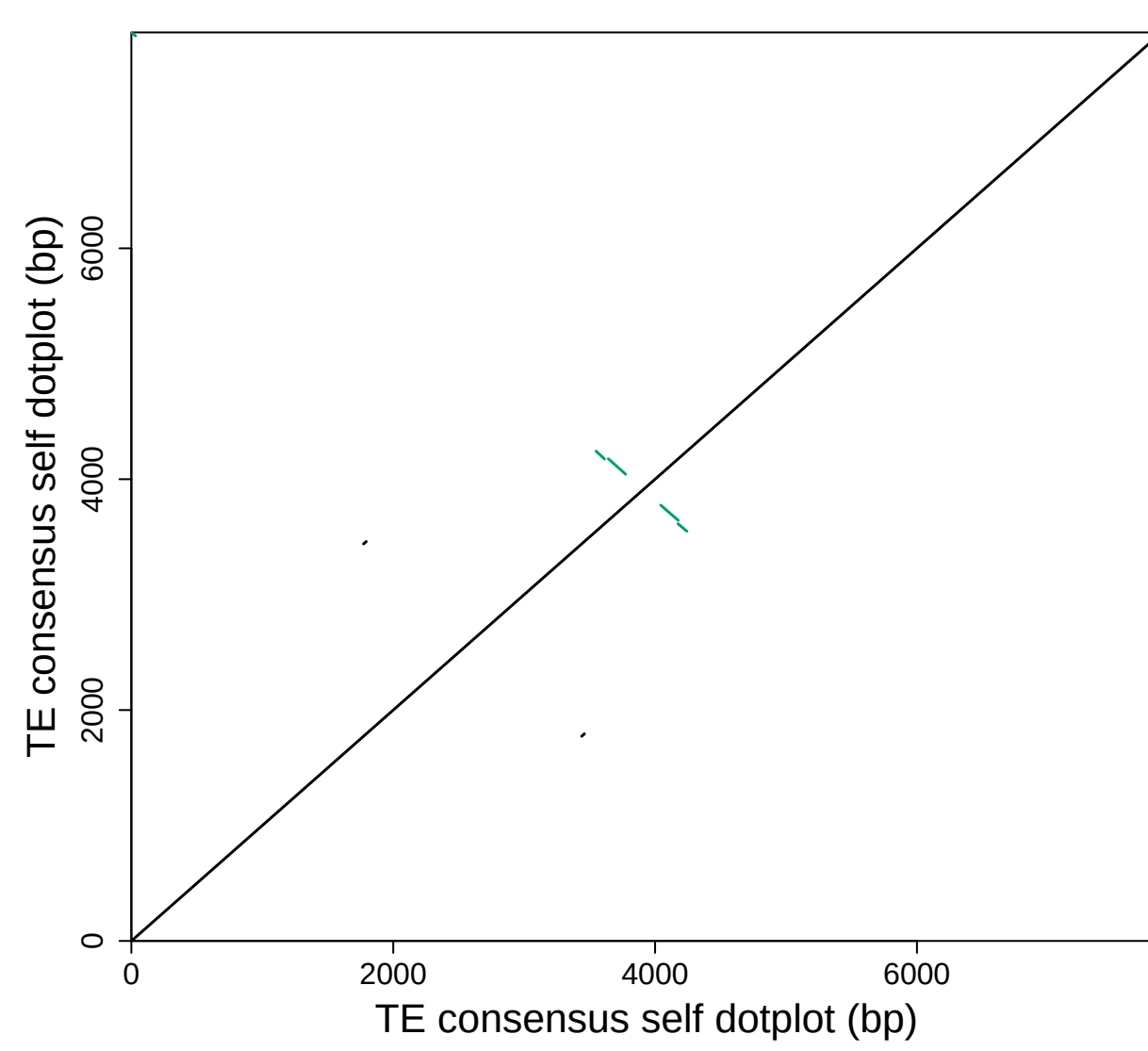
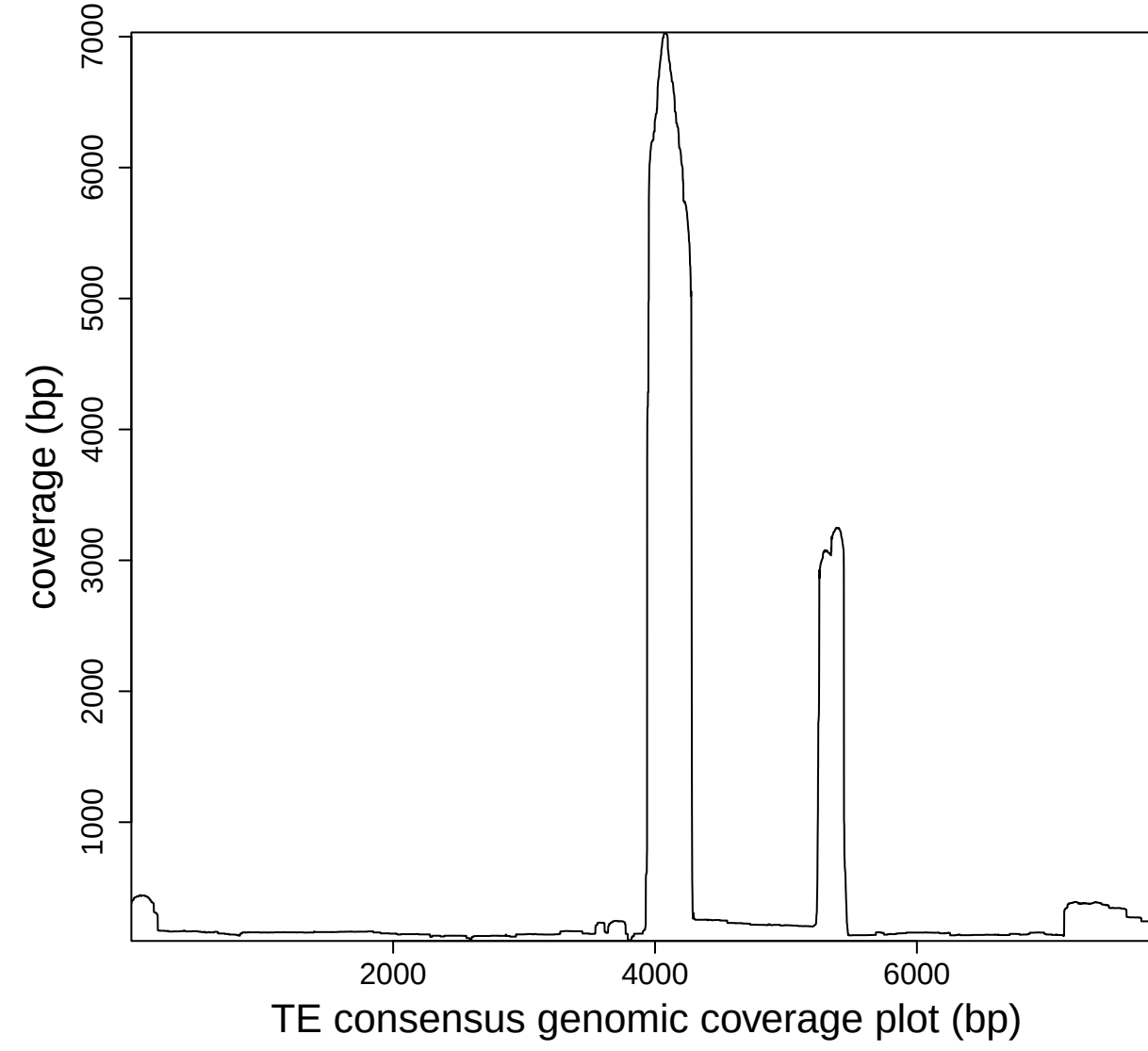
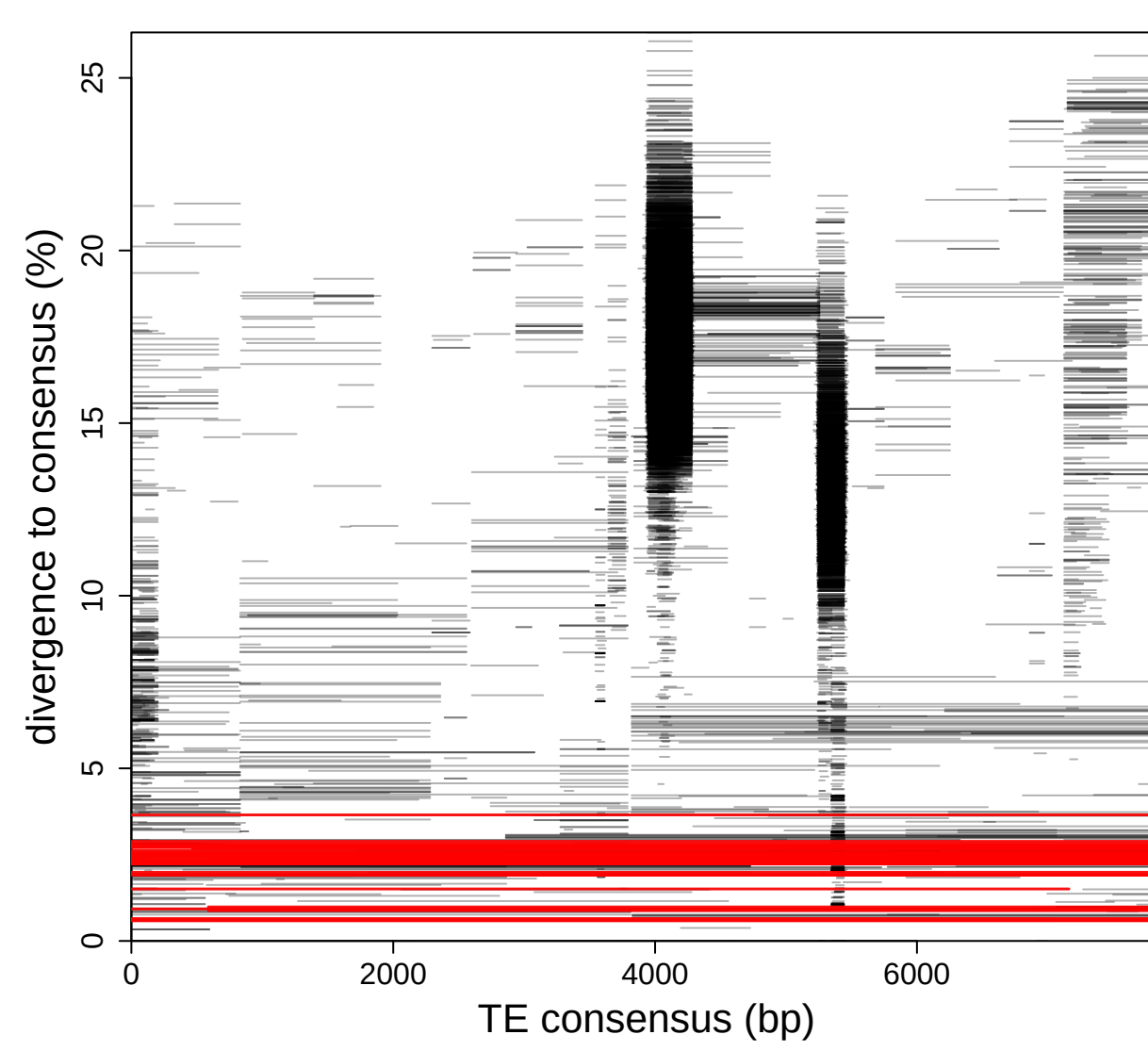
MSA length = 7870

[illegible]



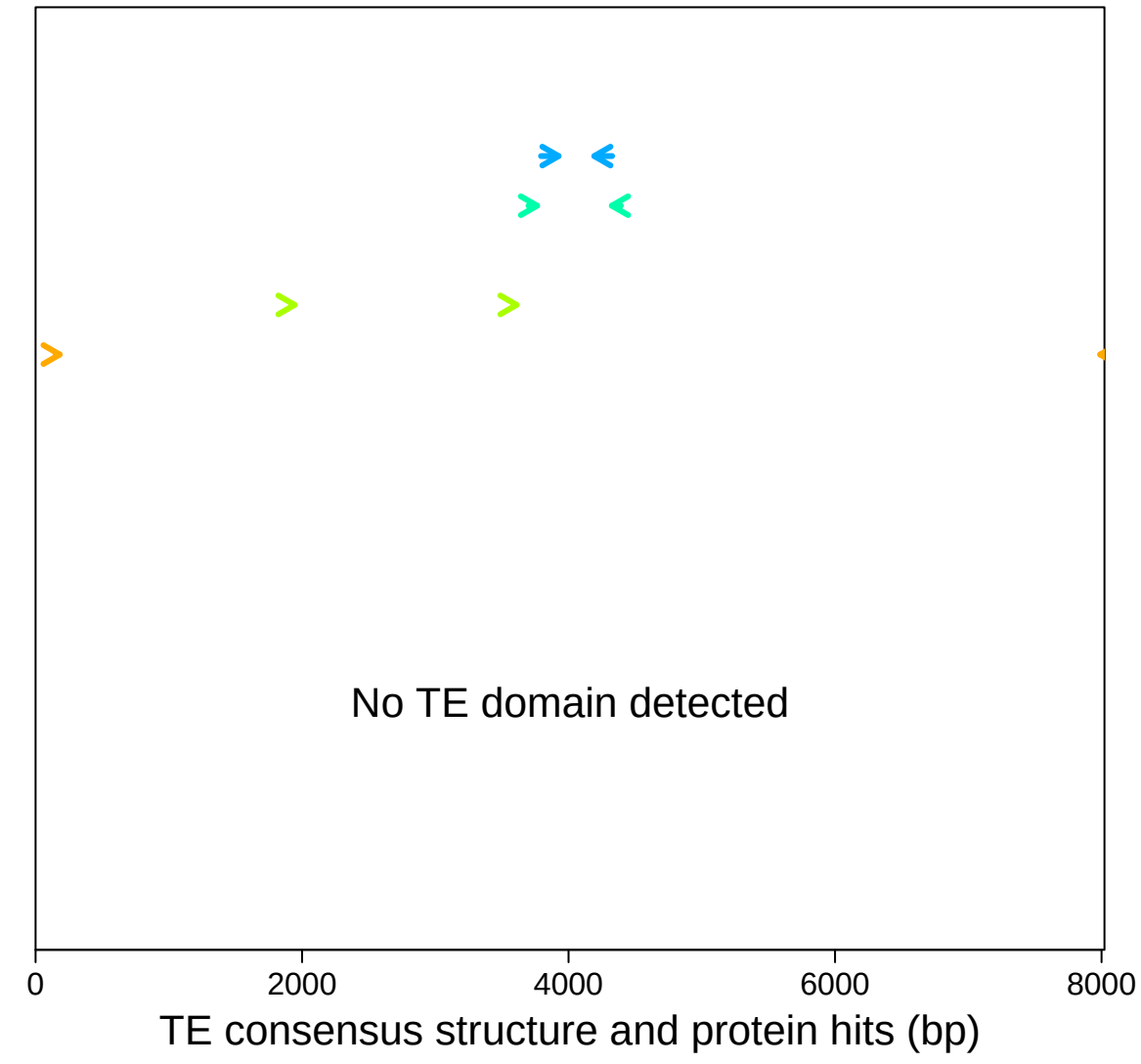
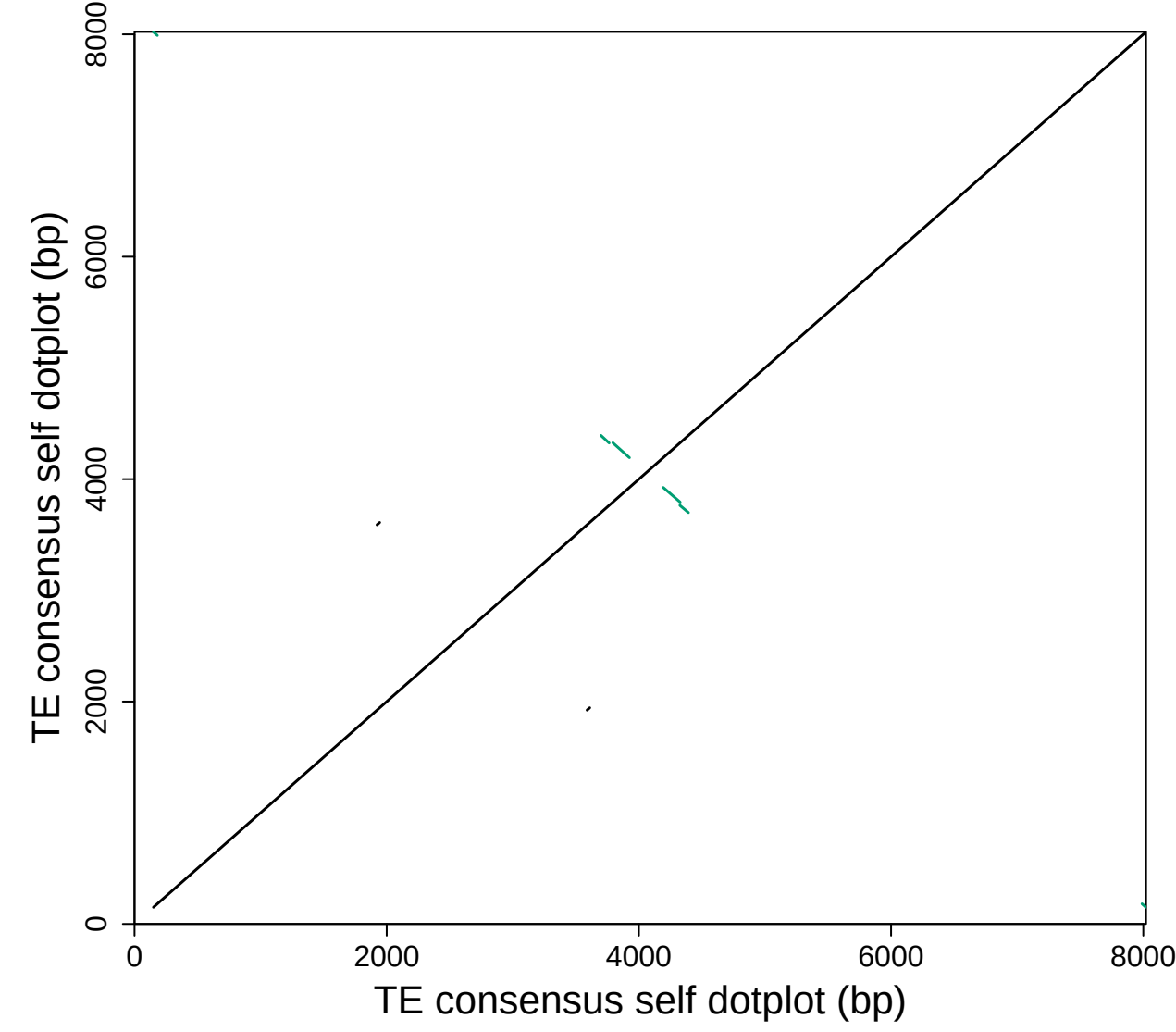
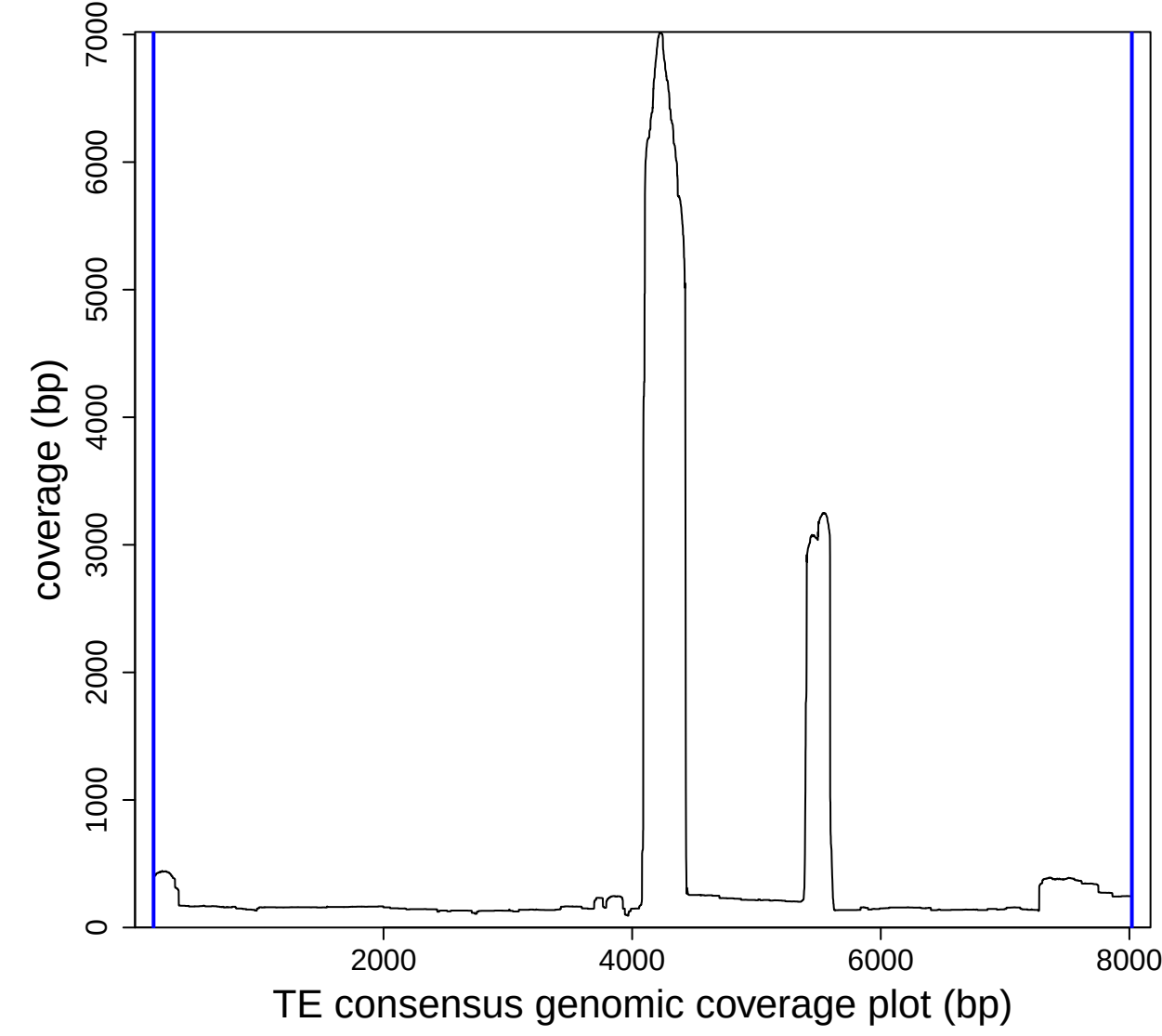
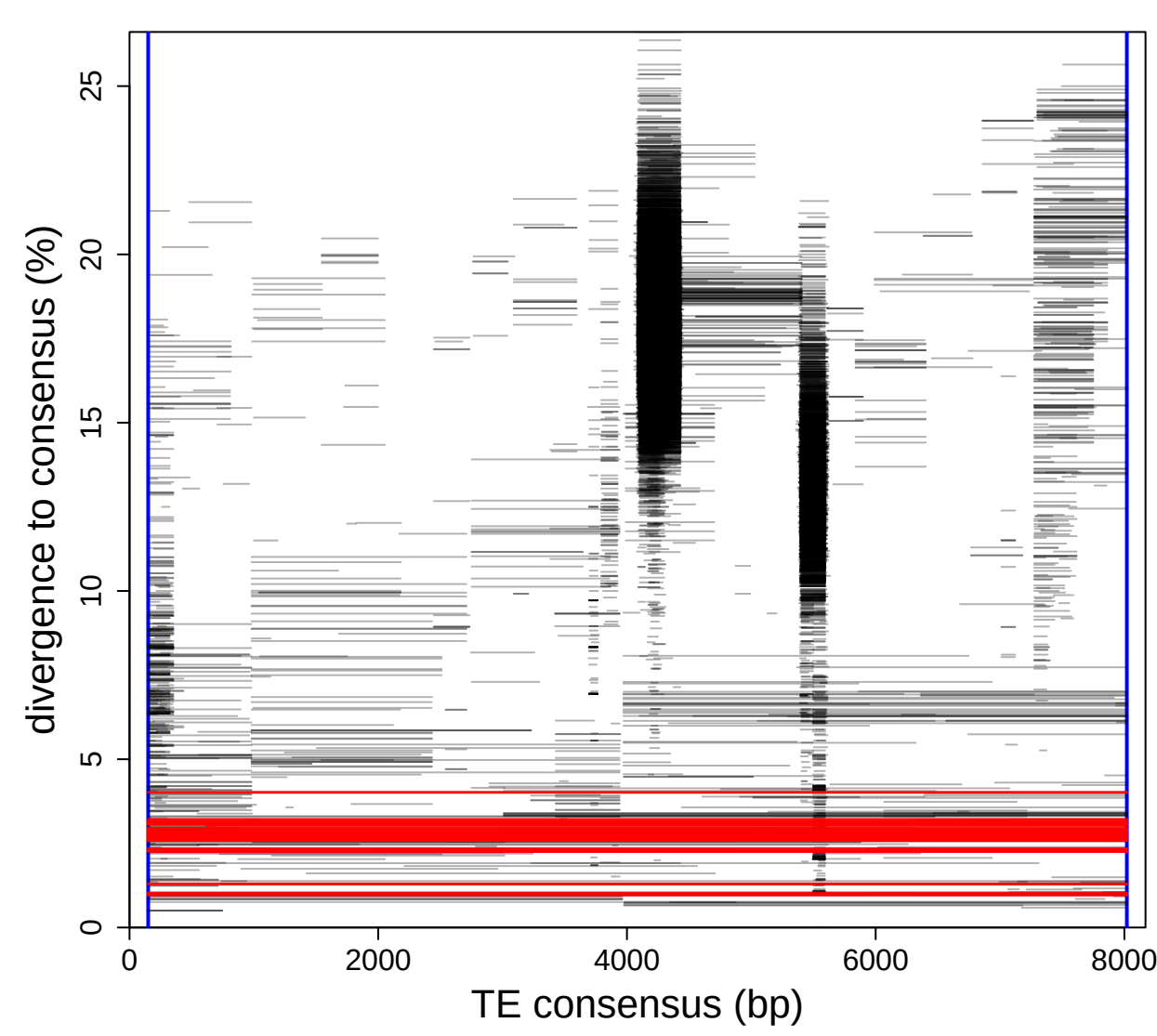
a.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 7870bp; fragments: 11713; full length: 53 (>=7083bp)

After TEtrimmer 7870 bp



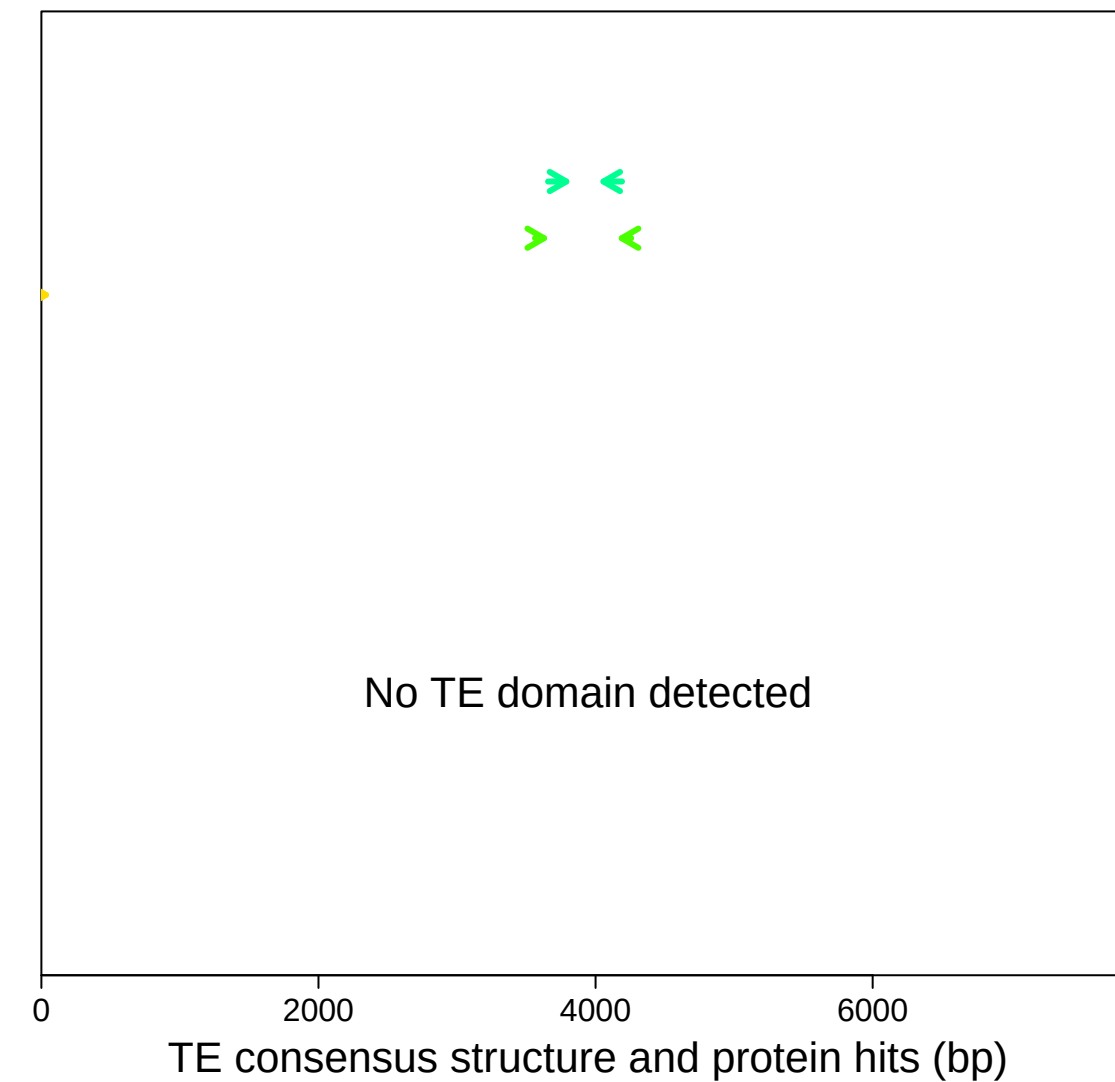
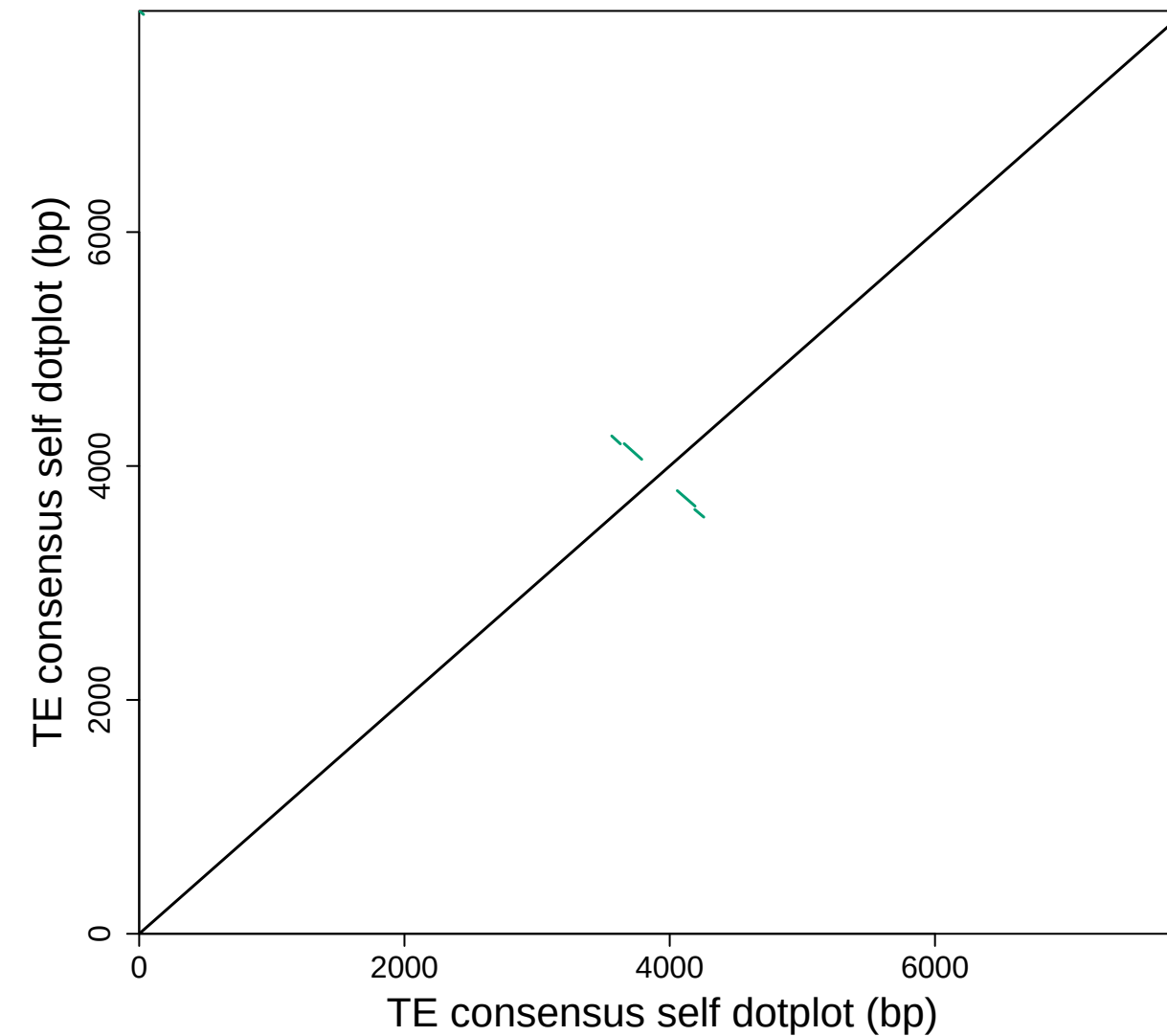
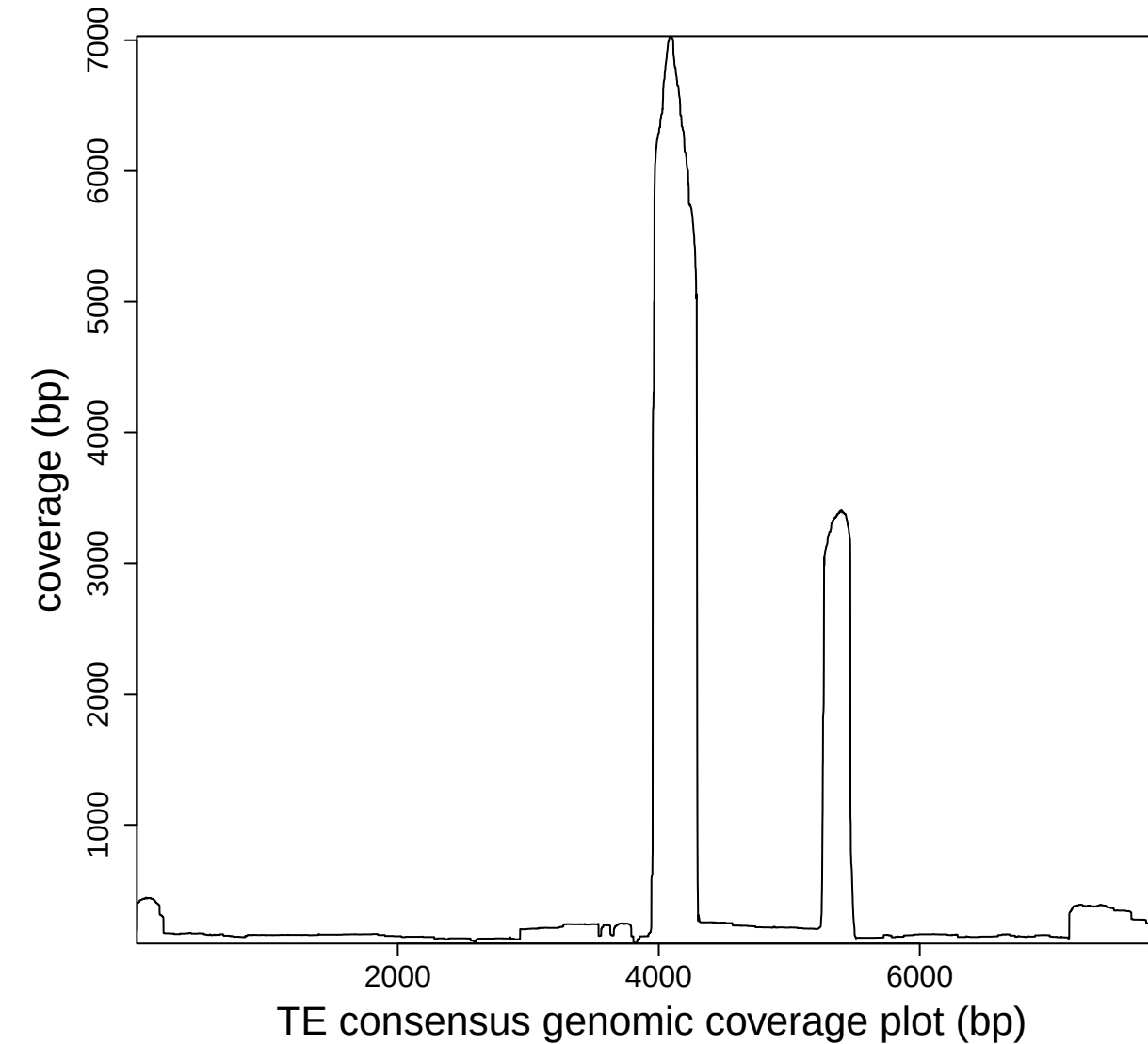
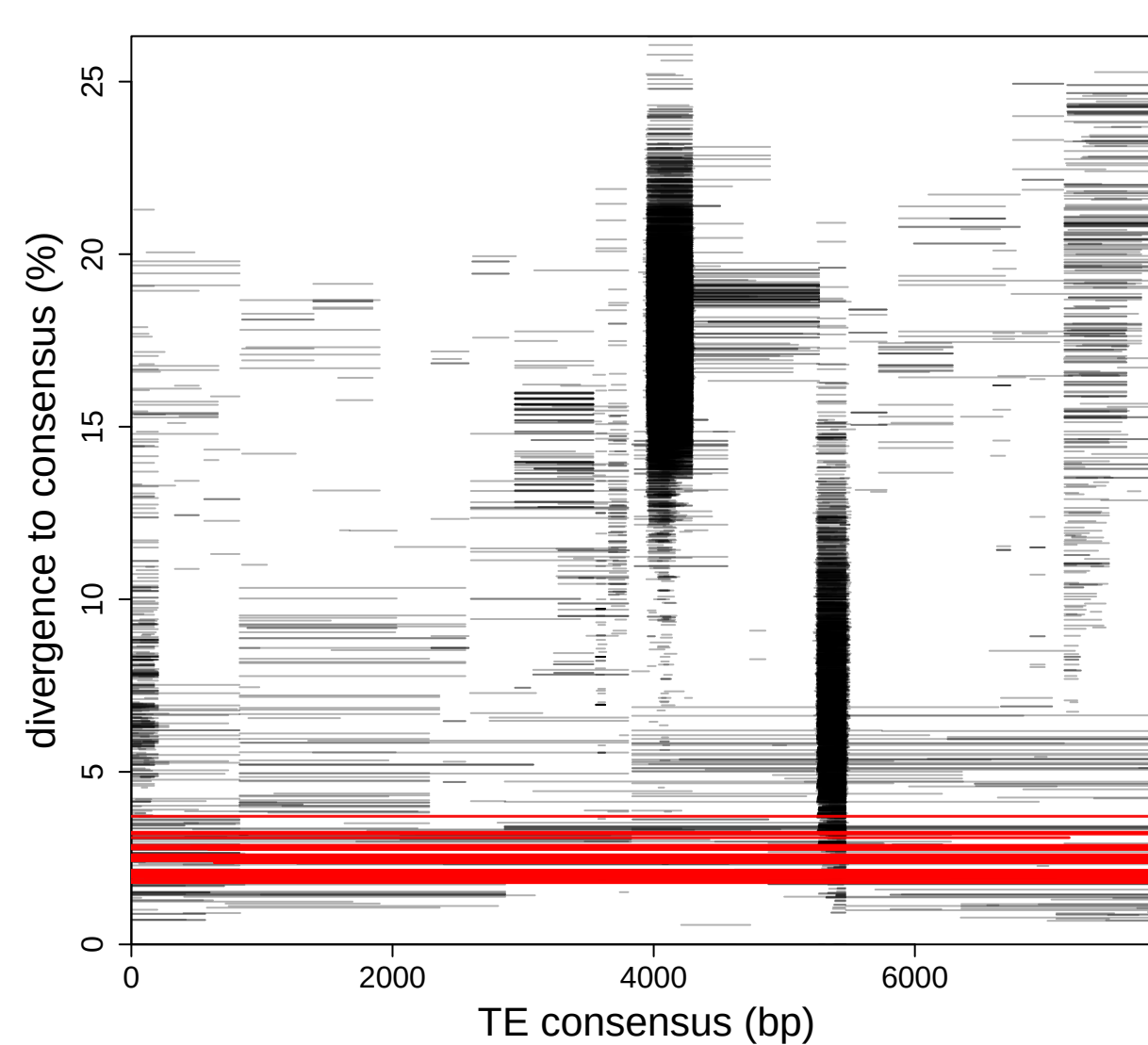
3.fasta.b.bed uf.bed g_4.bed fm_1.bed 0 0 bcIn.fa aln.fa cl.f
size: 8170bp; fragments: 11697; full length: 44 (>=7353bp)

After TEtrimmer Extended plot Blue lines are boundaries

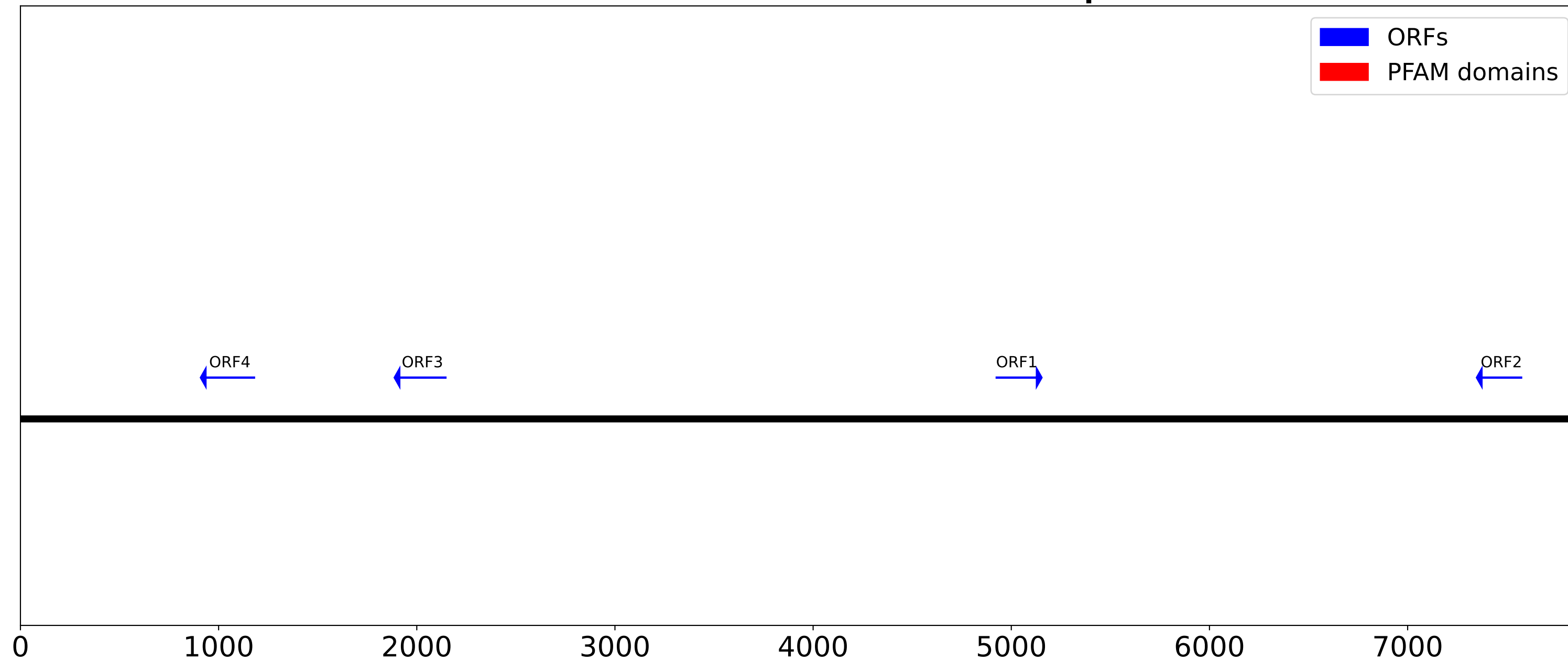


TE: rnd_5_family_3343
size: 7891bp; fragments: 11996; full length: 52 (≥ 7101.9 bp)

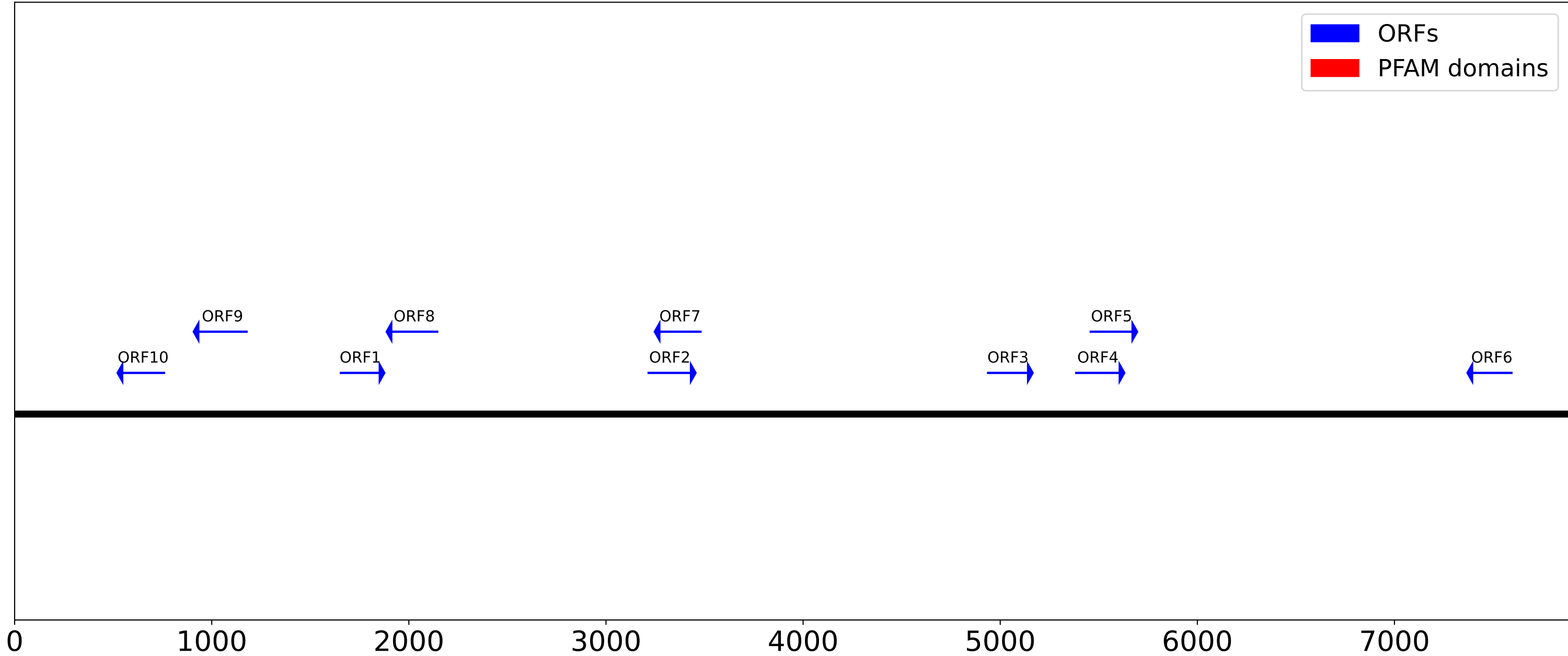
Before TEtrimmer 7891 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

