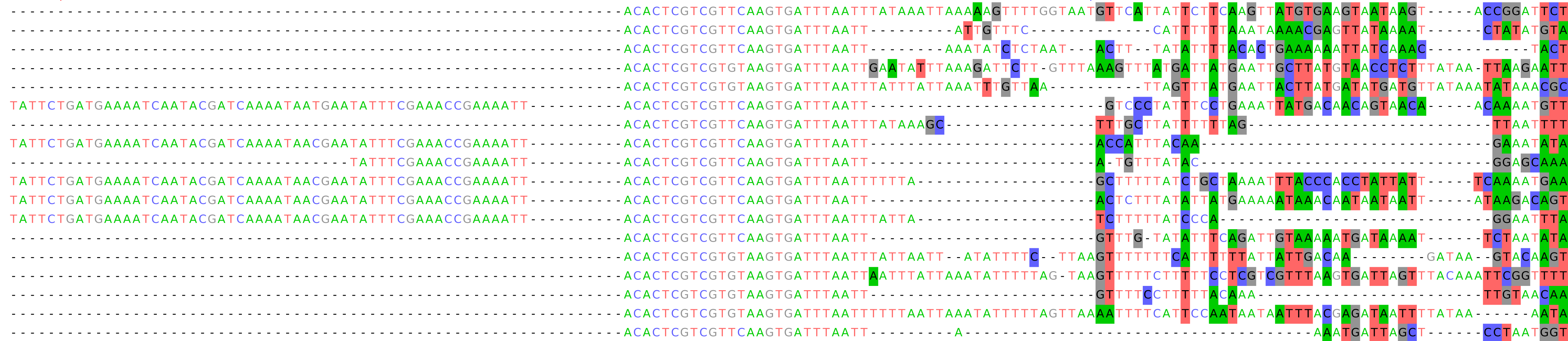
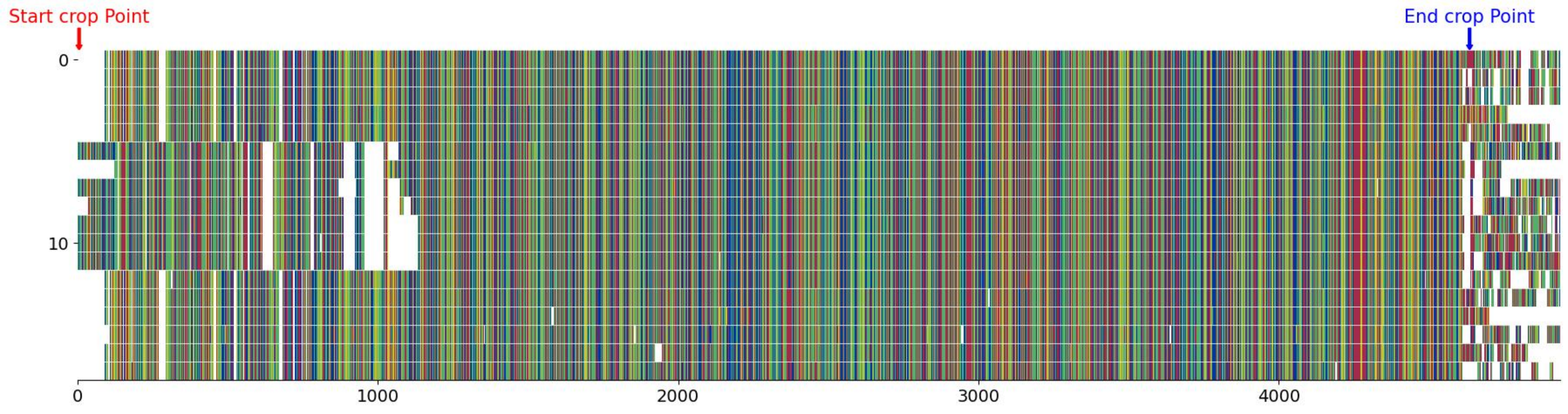


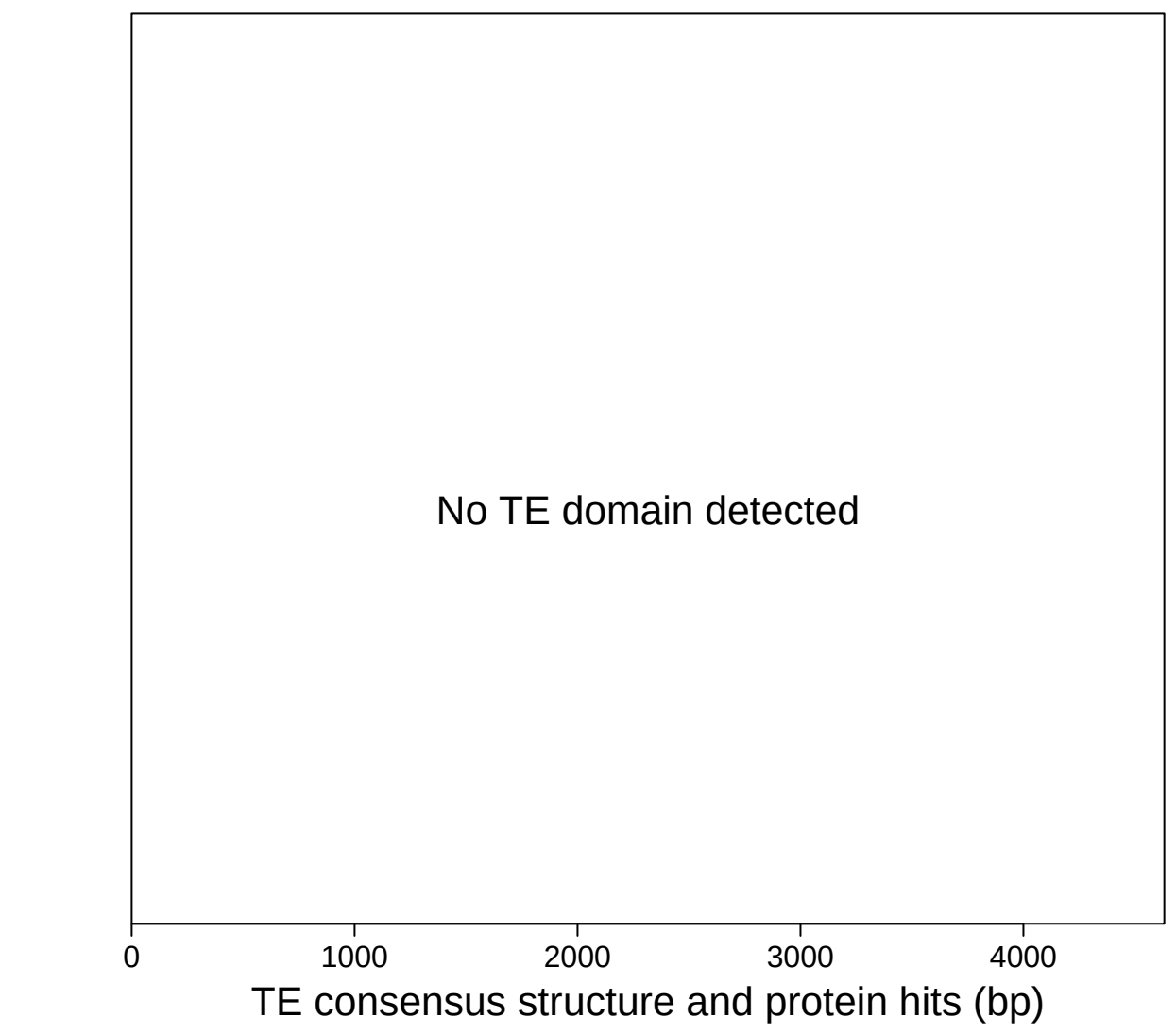
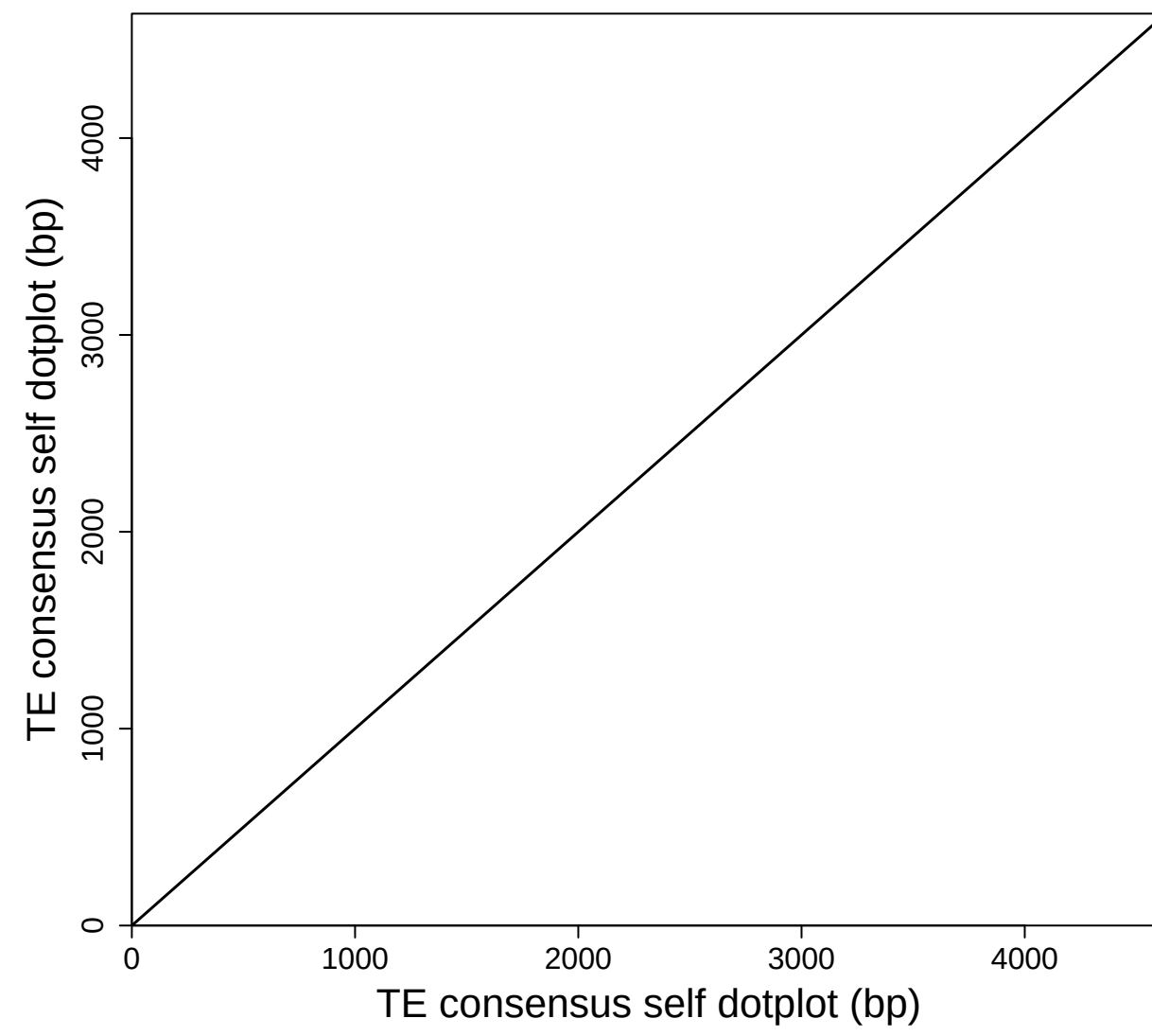
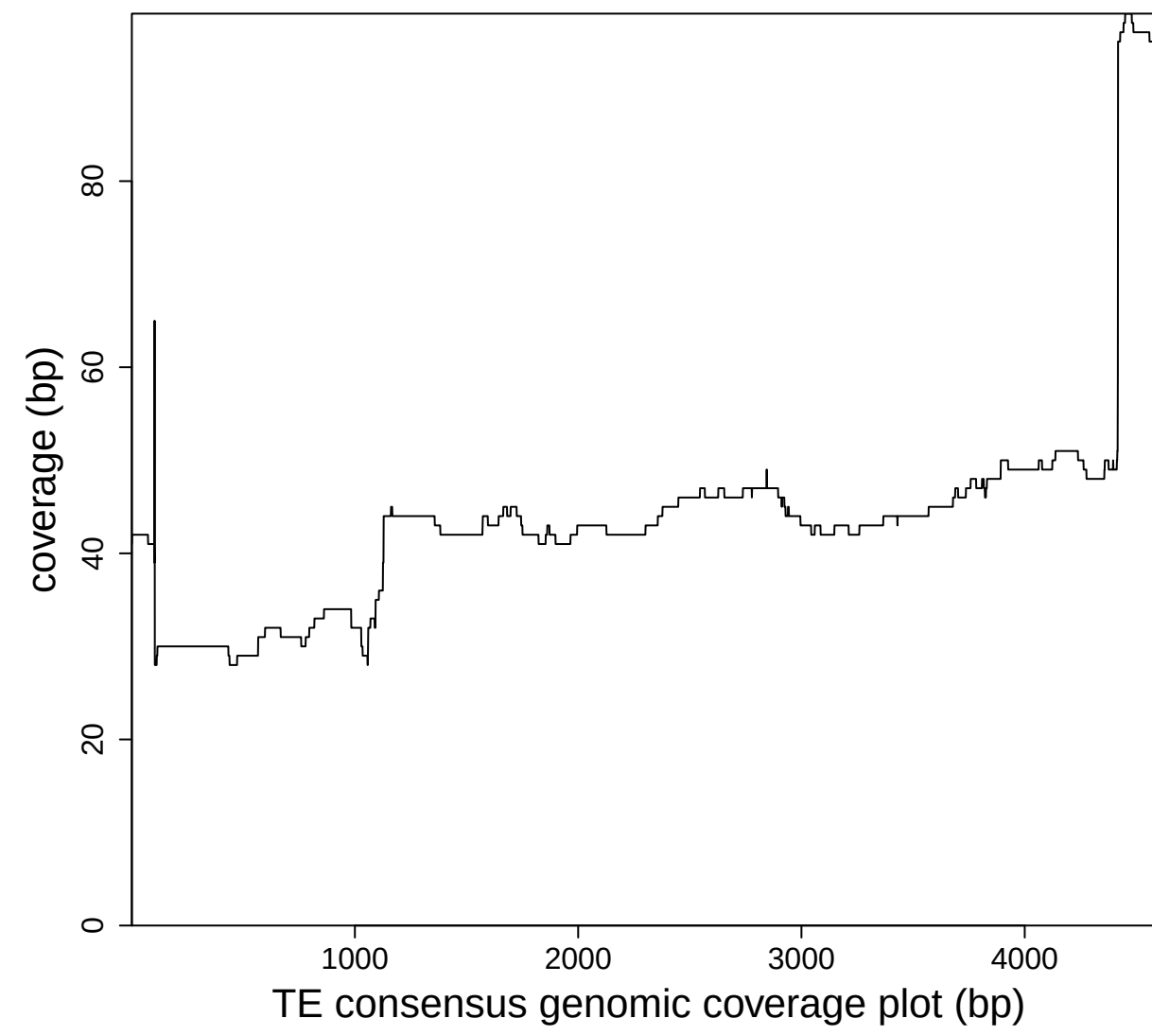
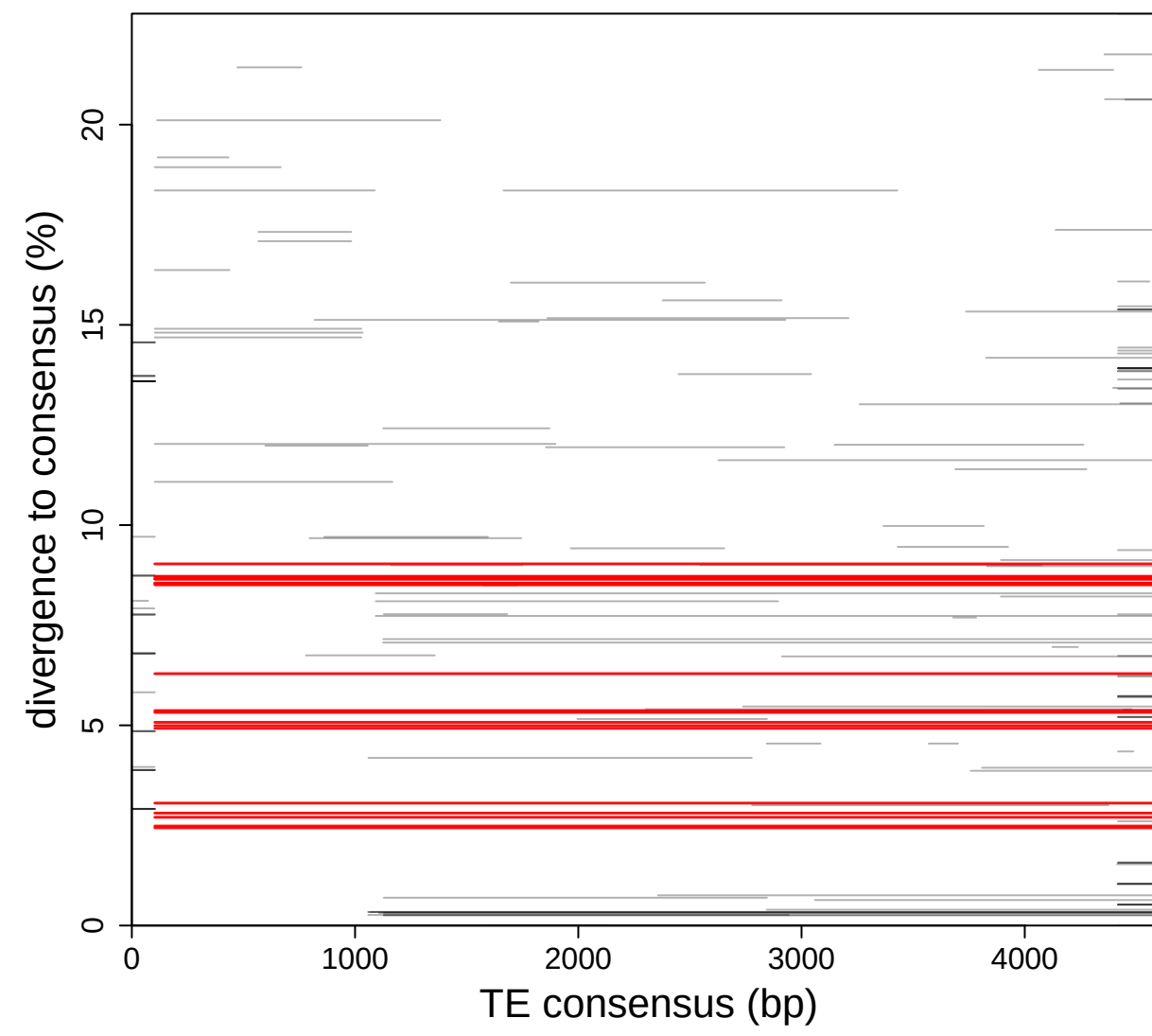
MSA length = 4632





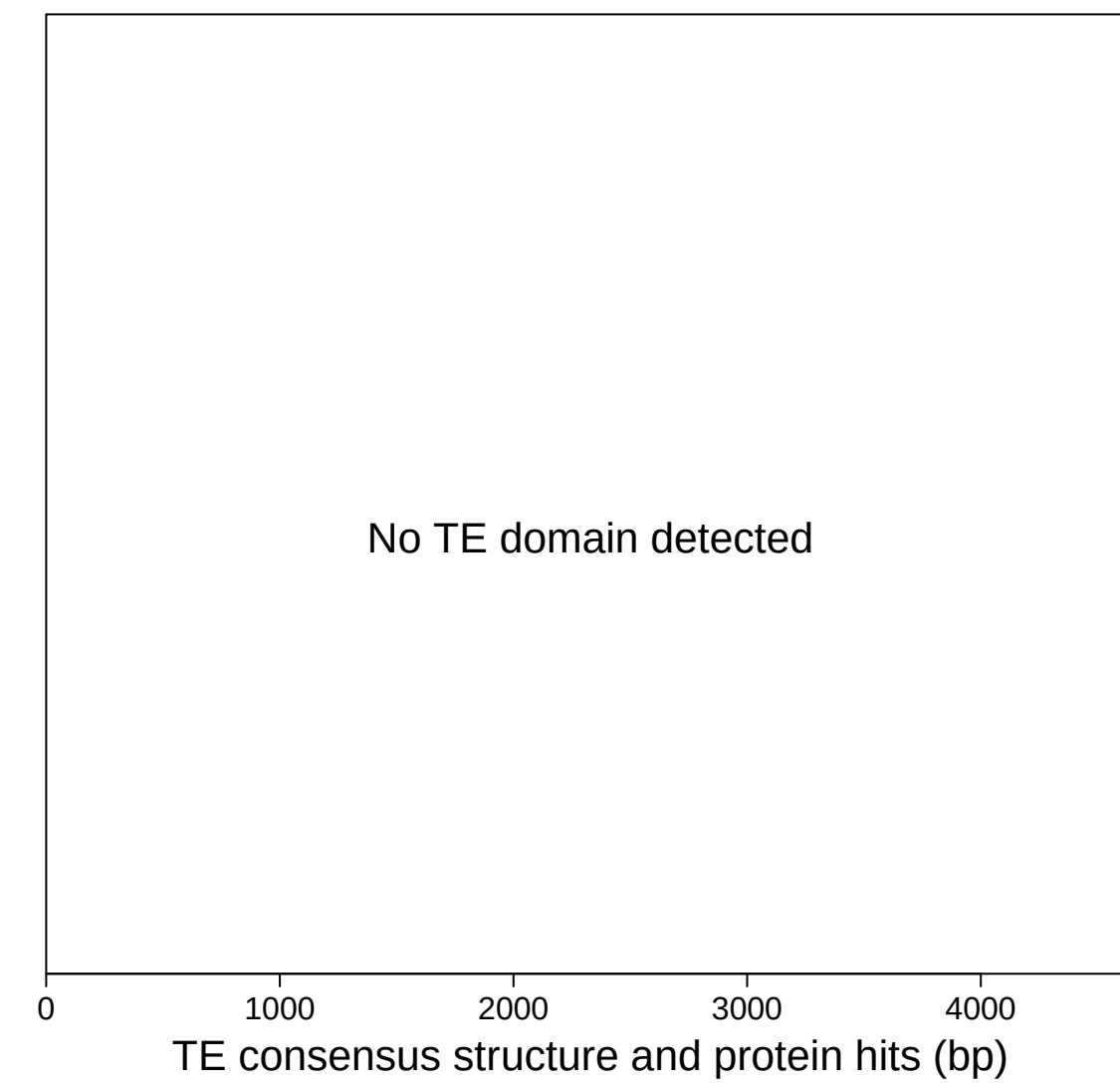
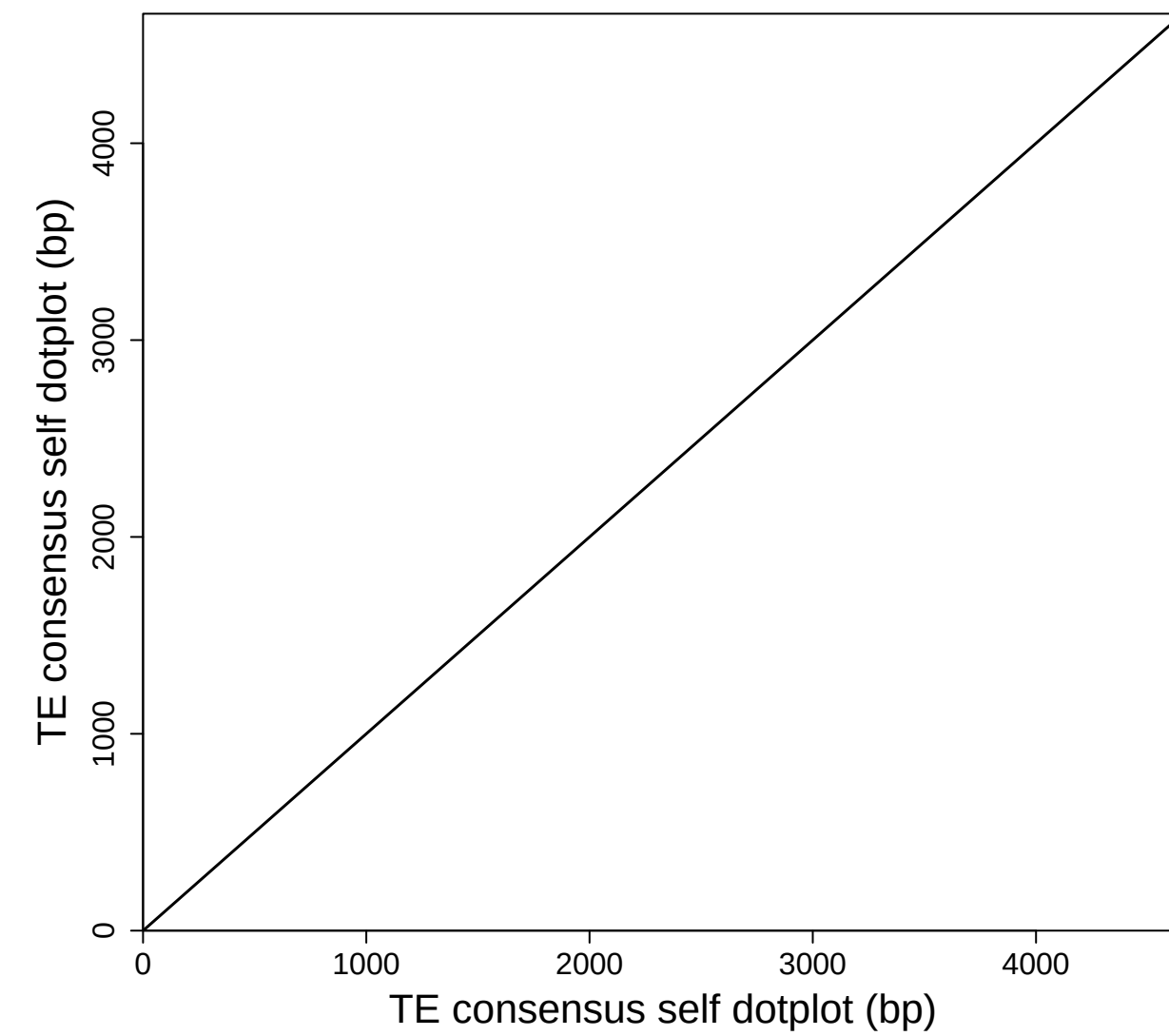
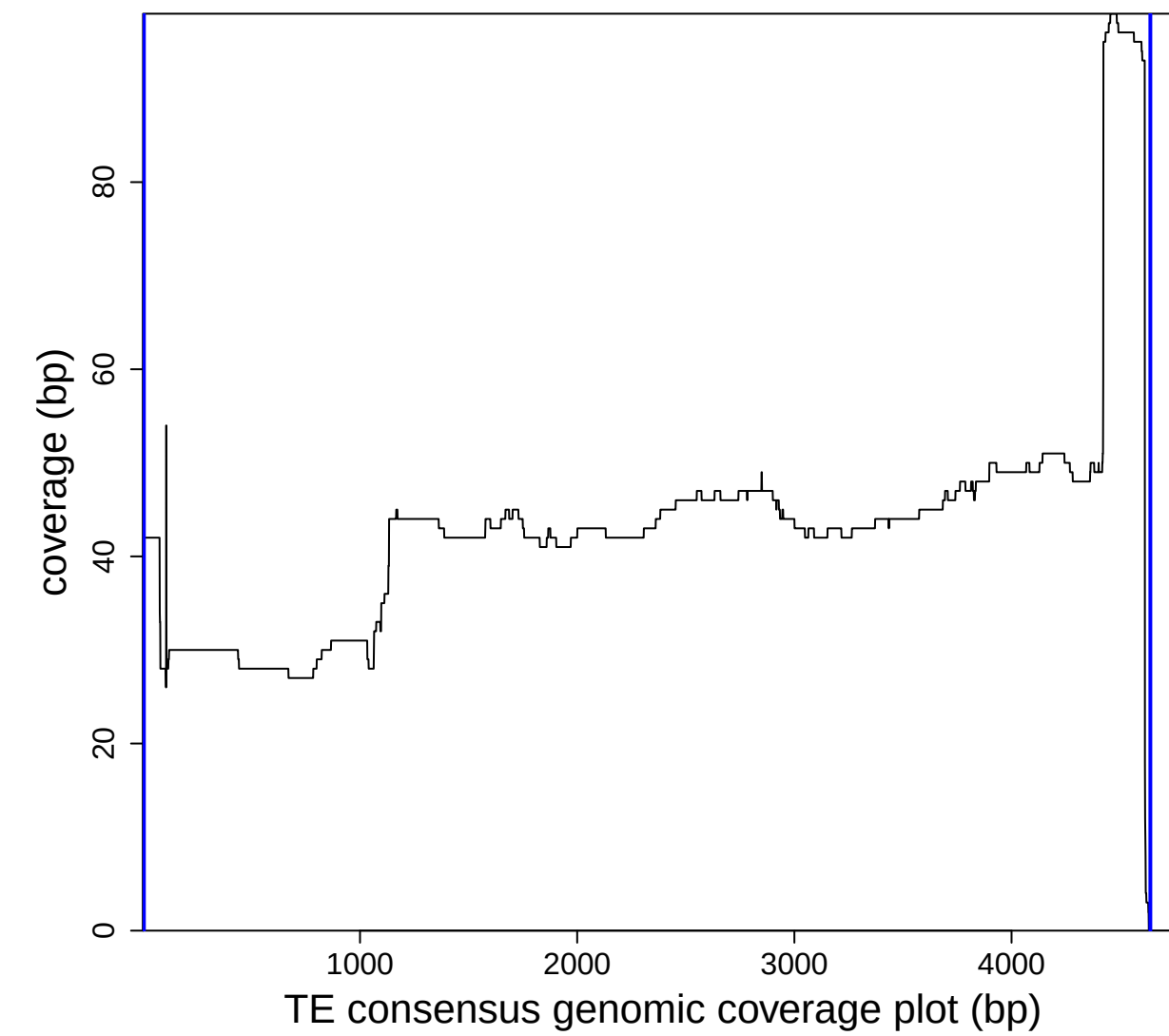
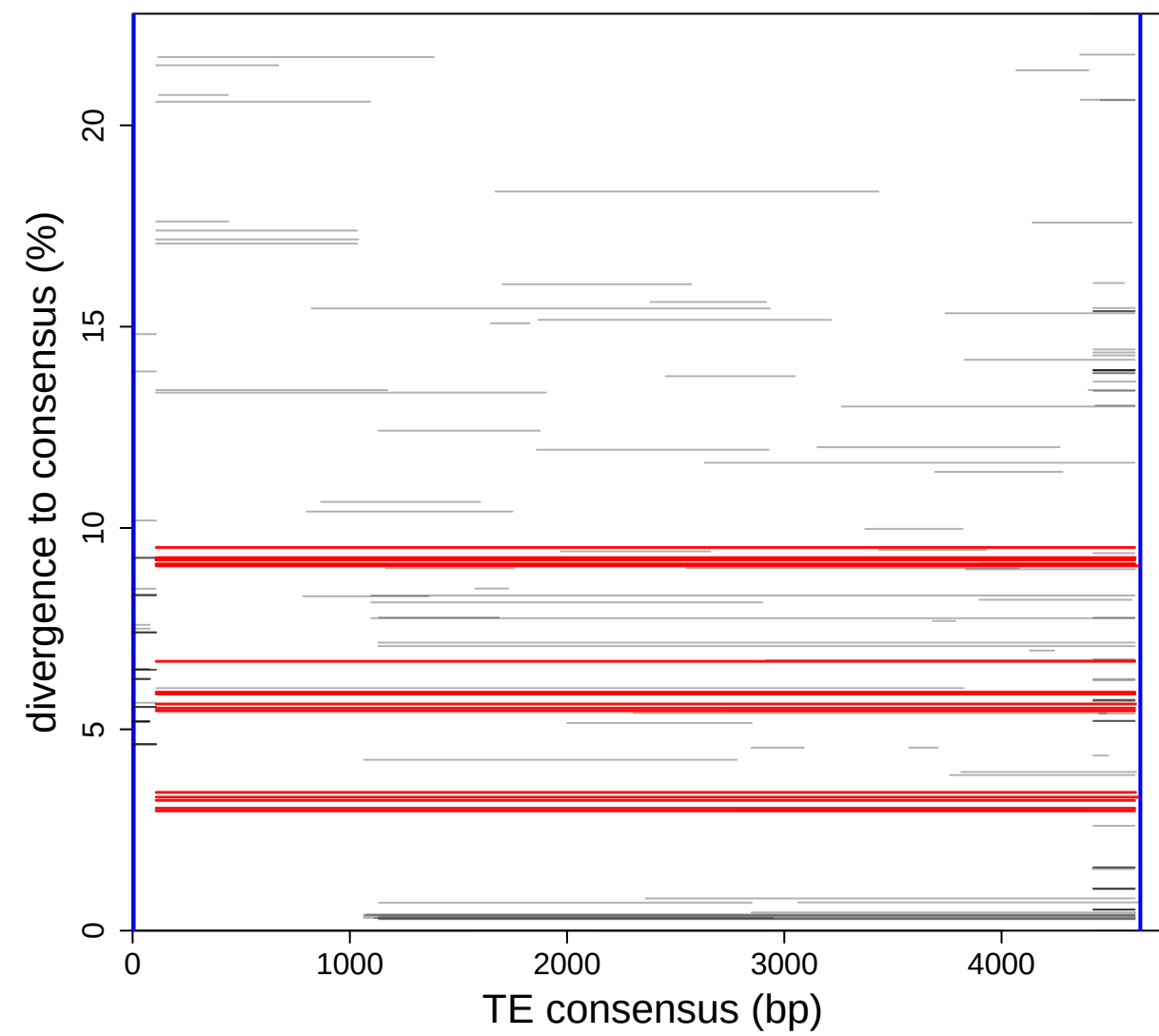
b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 4632bp; fragments: 189; full length: 19 (≥ 4168.8 bp)

After TETrimmer 4632 bp

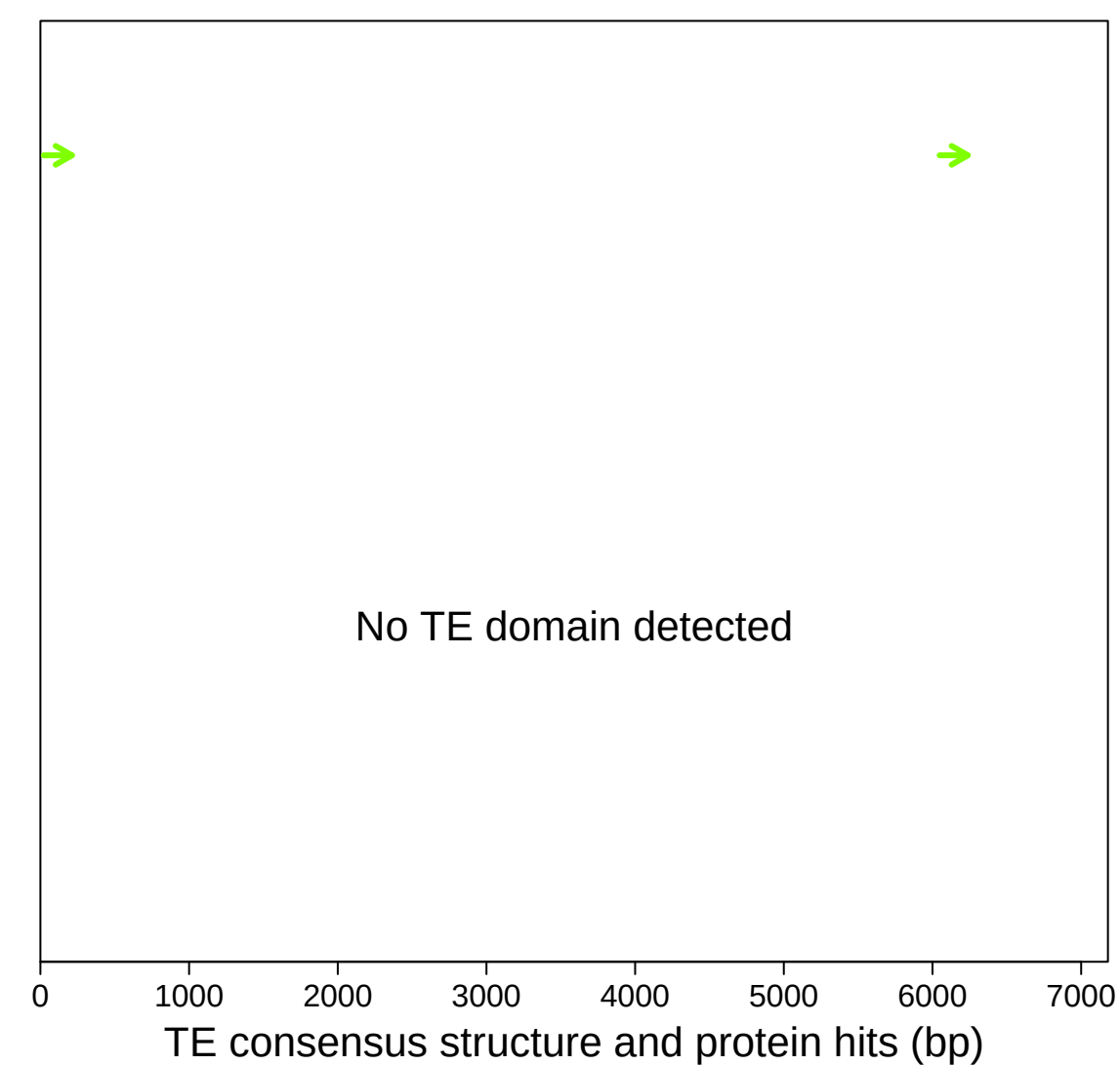
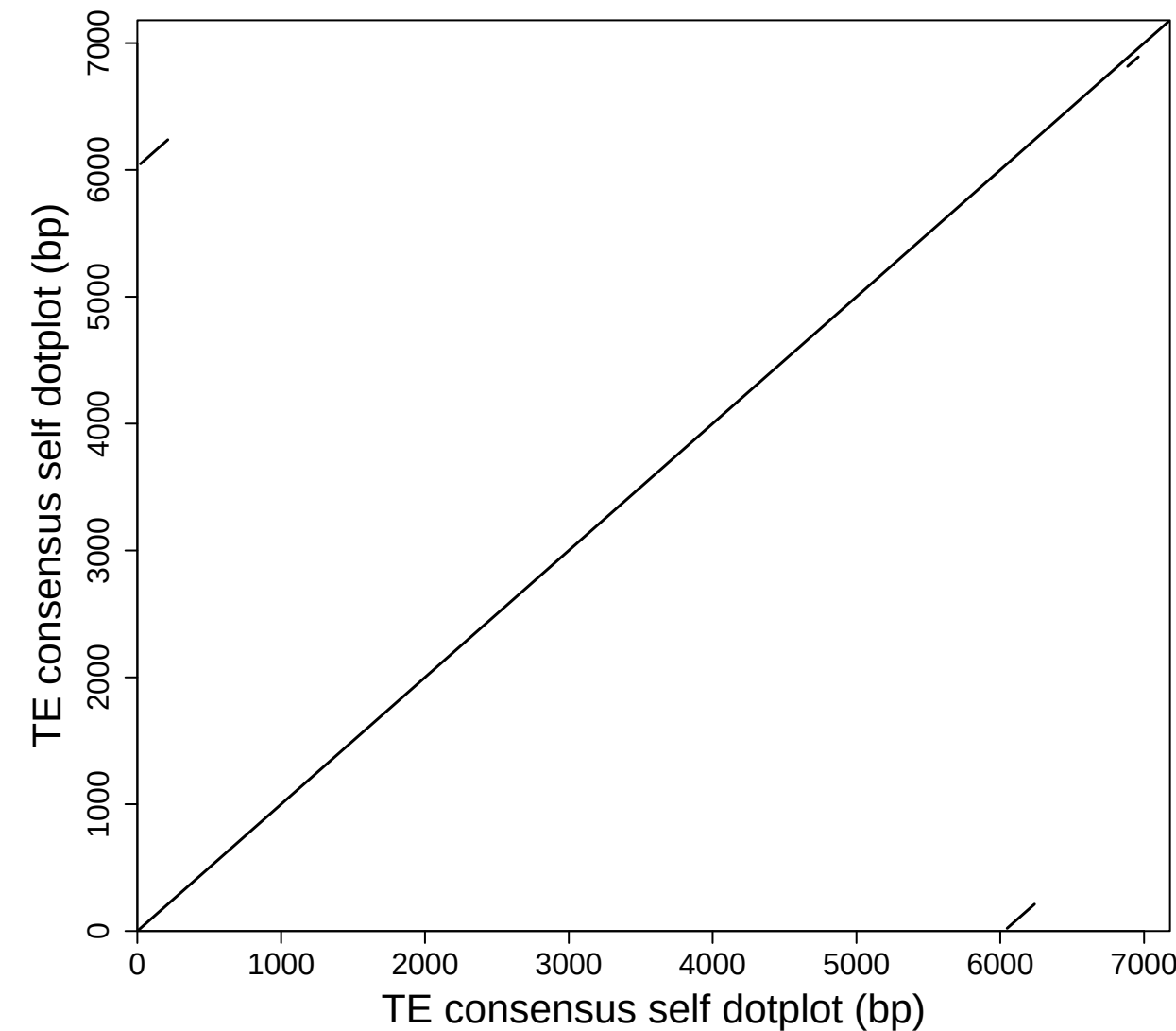
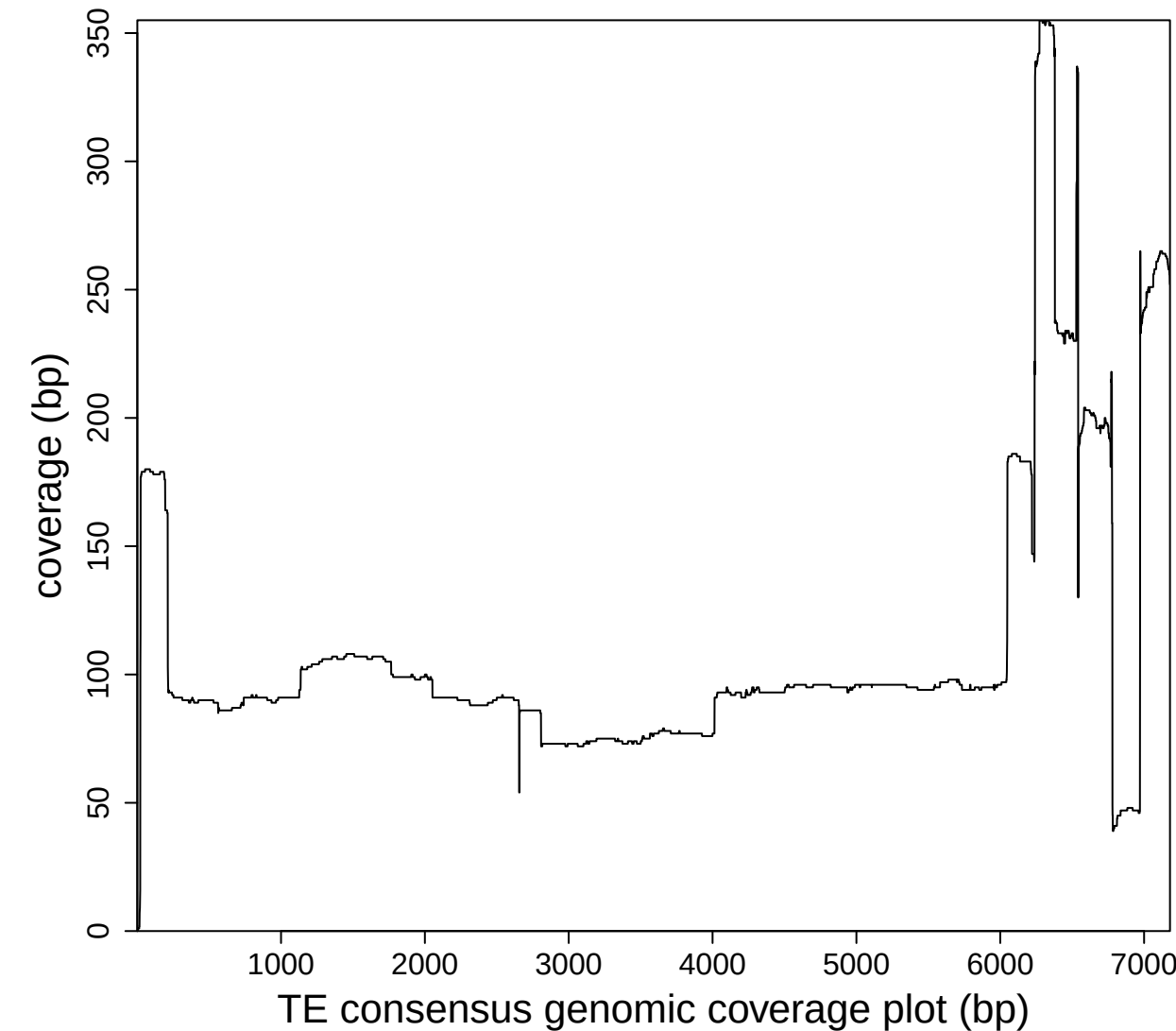
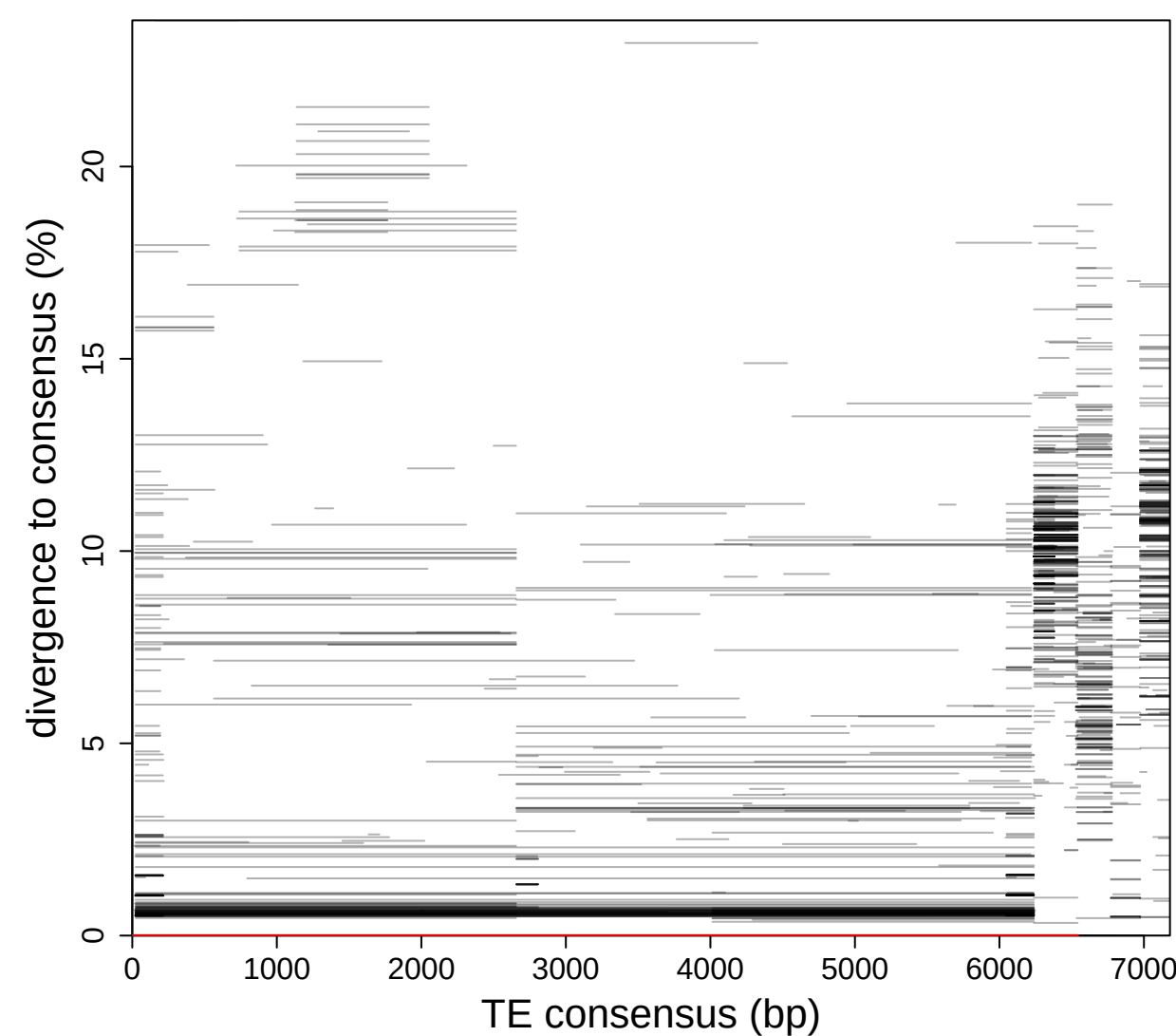


.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.fa
size: 4789bp; fragments: 185; full length: 19 (>=4310.1bp)

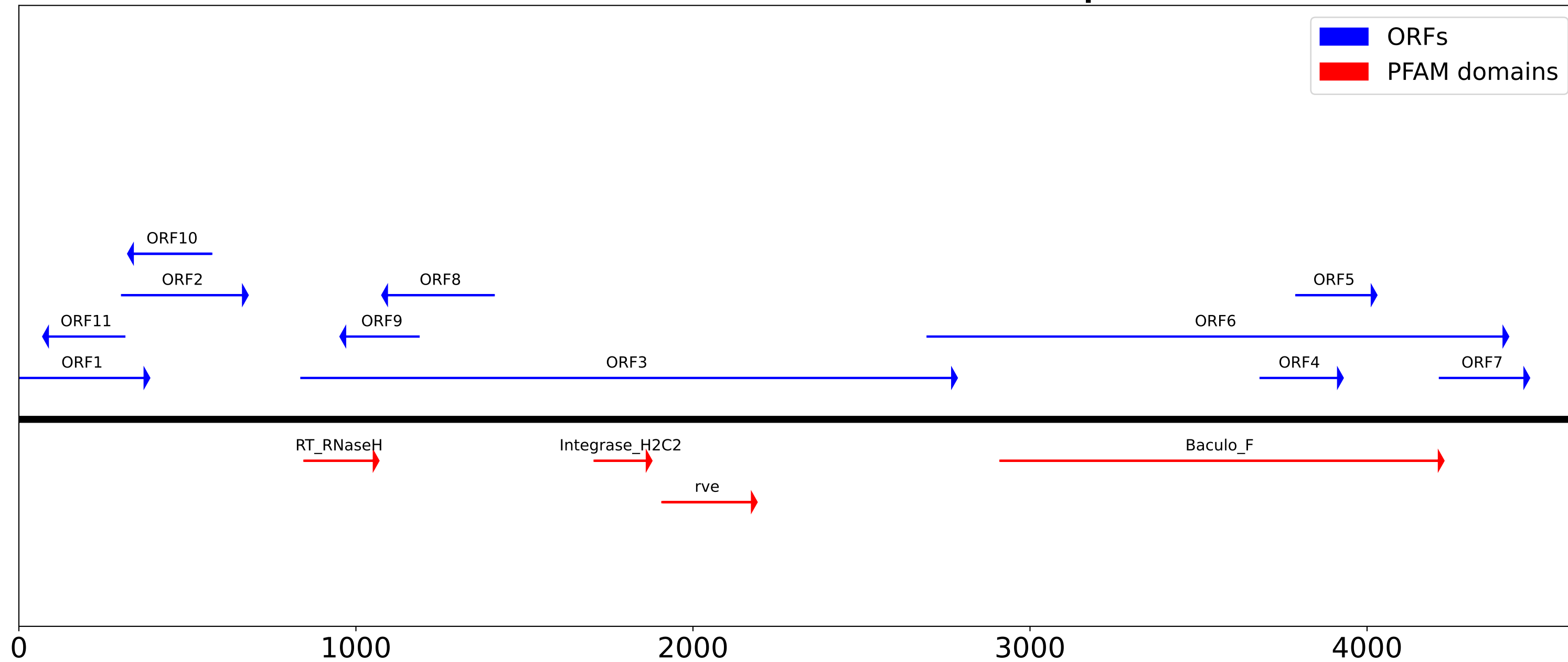
After TEtrimmer Extended plot Blue lines are boundaries



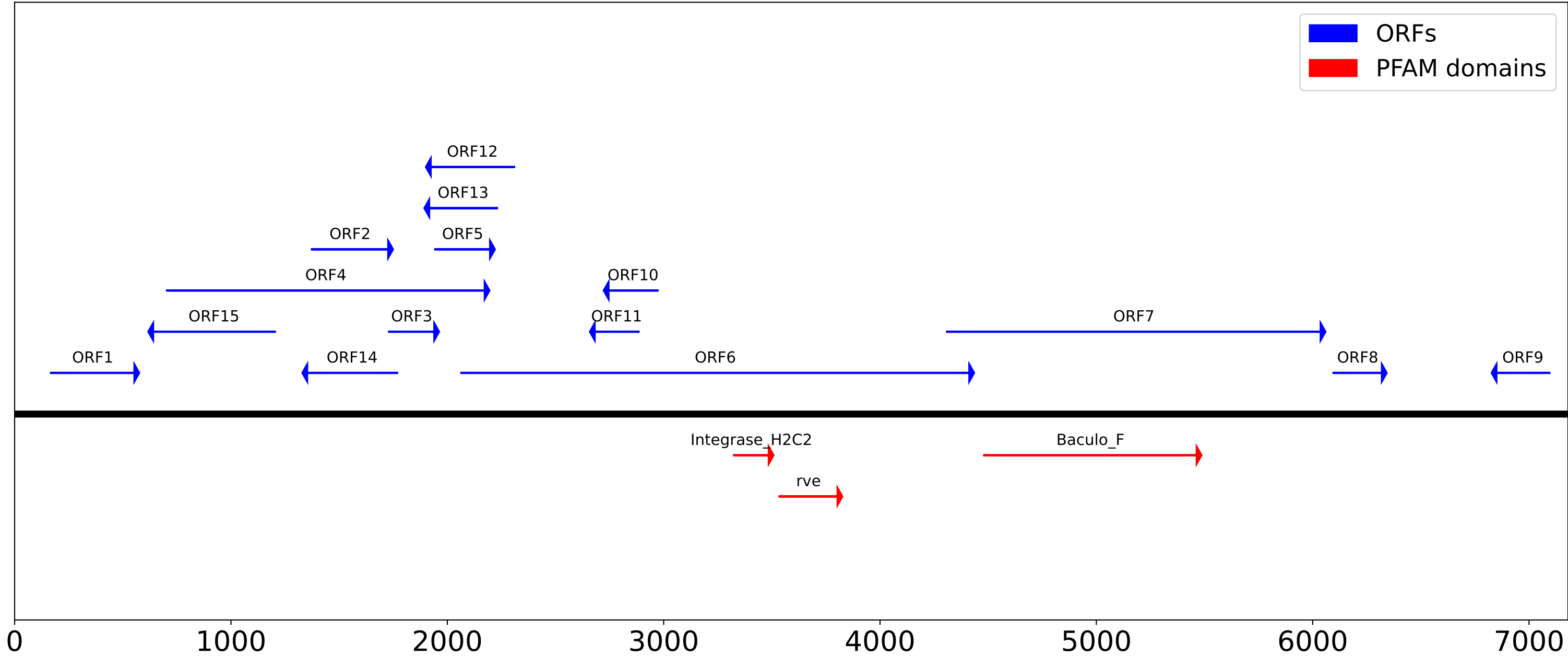
TE: ltr_1_family_994
size: 7180bp; fragments: 1322; full length: 1 (>=6462bp)



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

