

Start crop Point

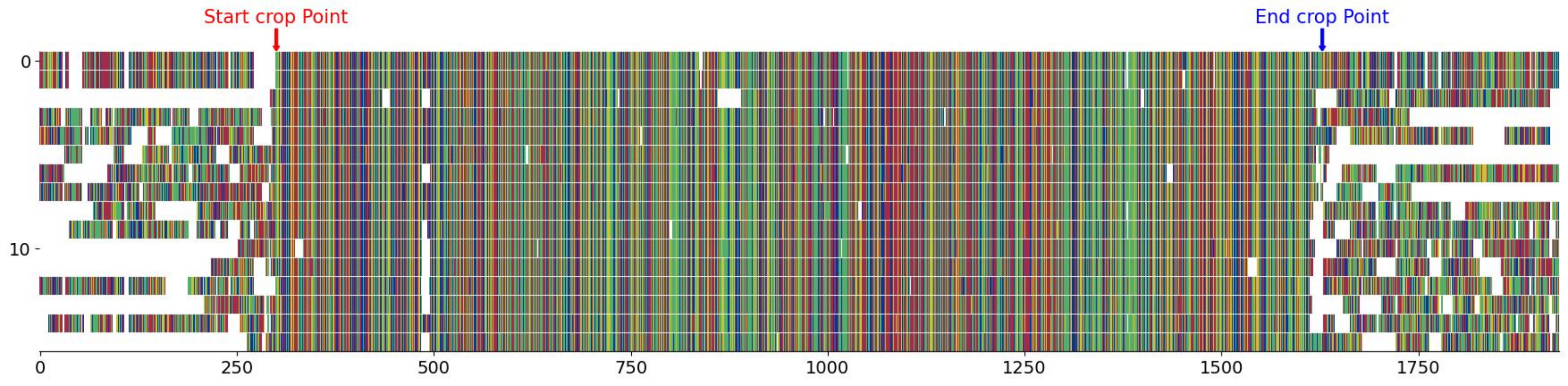
End crop Point

MSA length = 1329



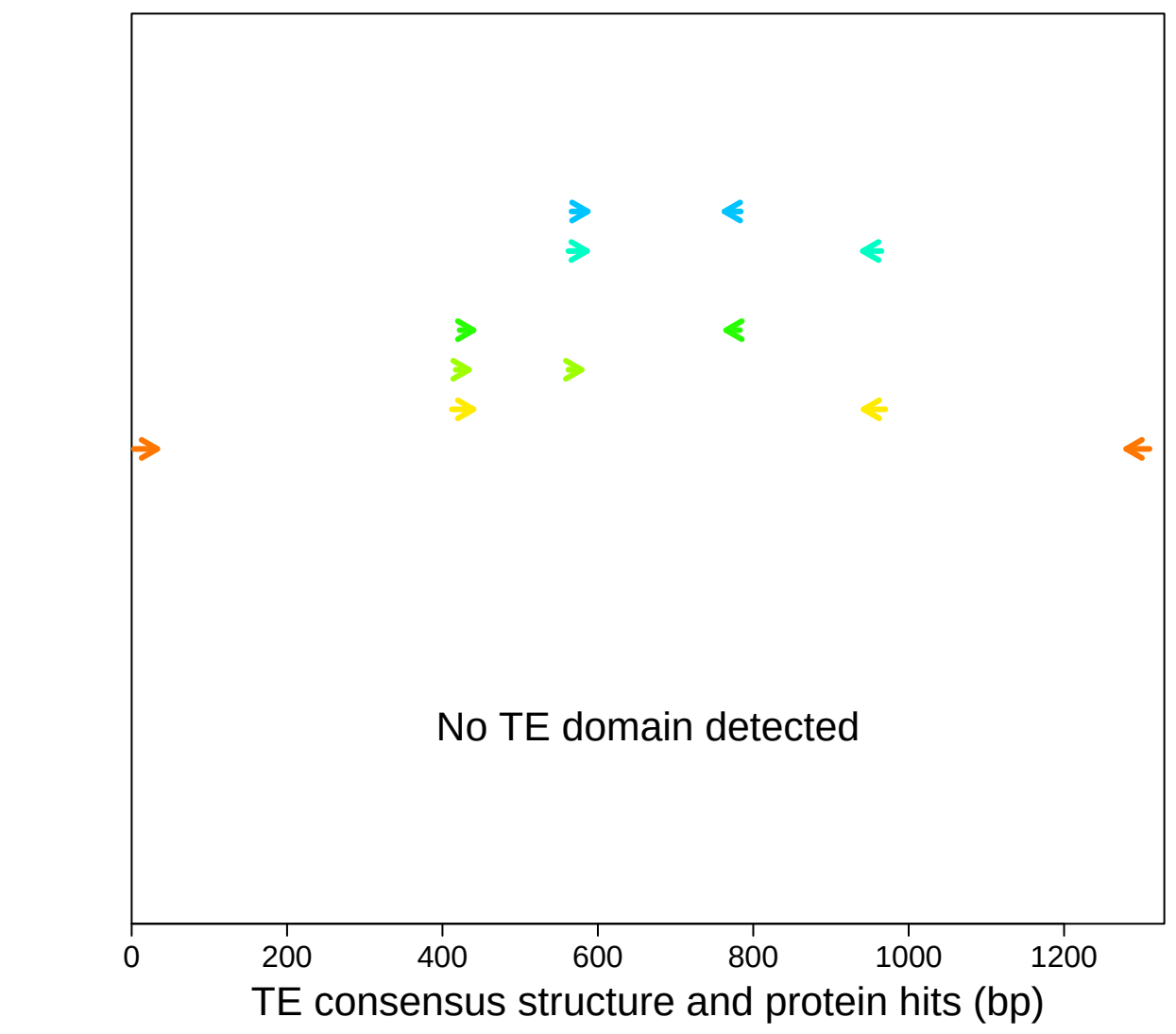
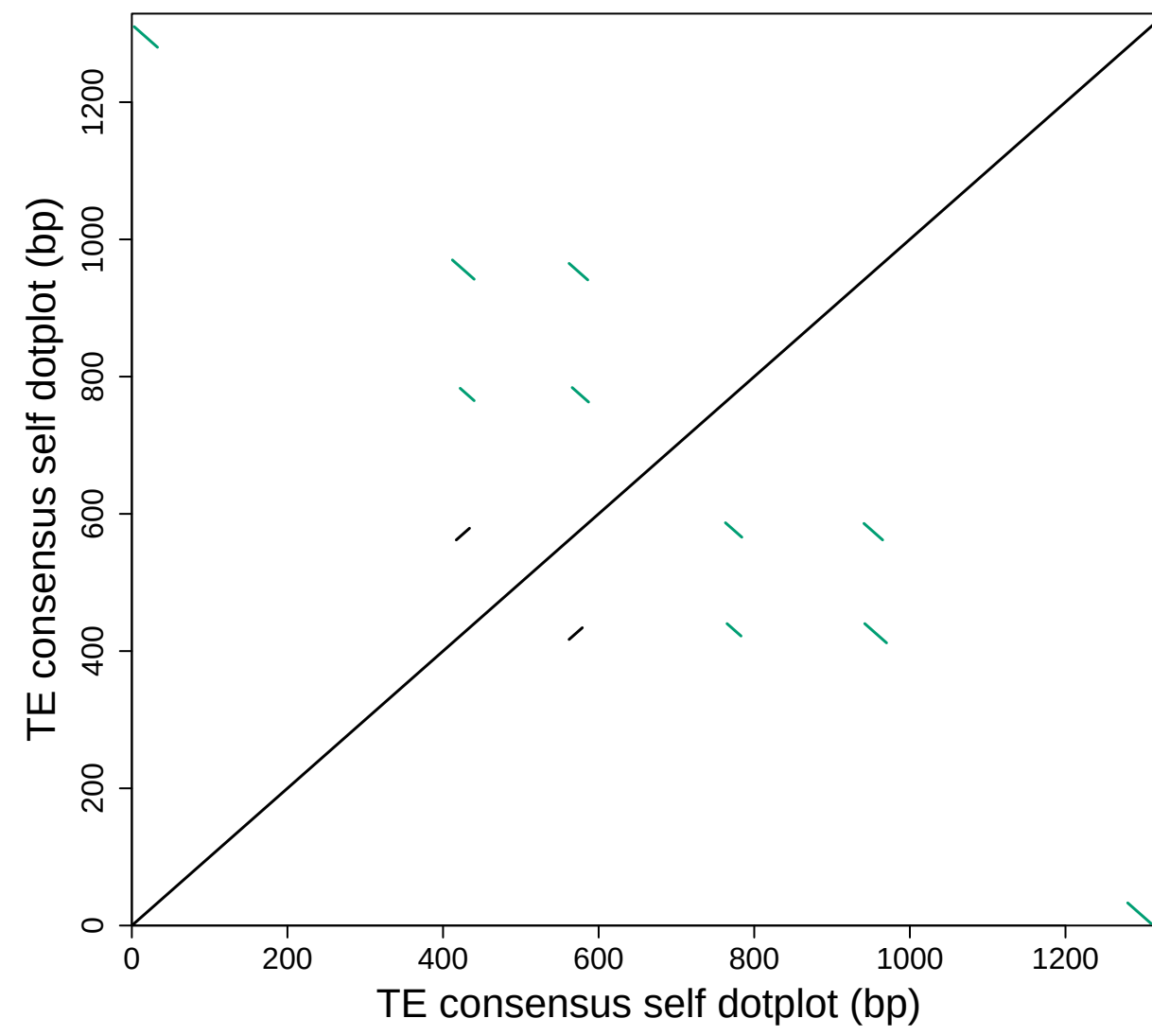
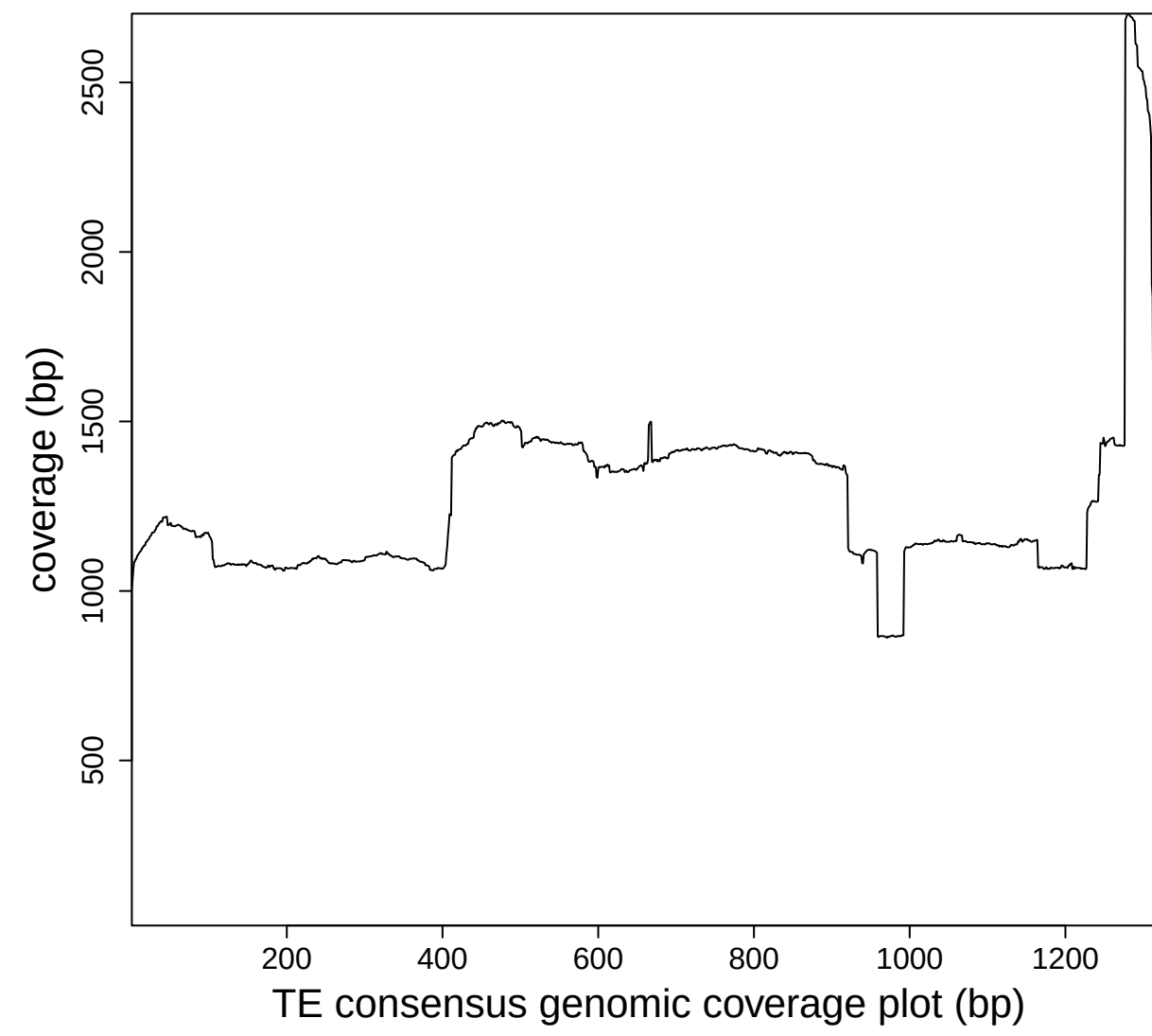
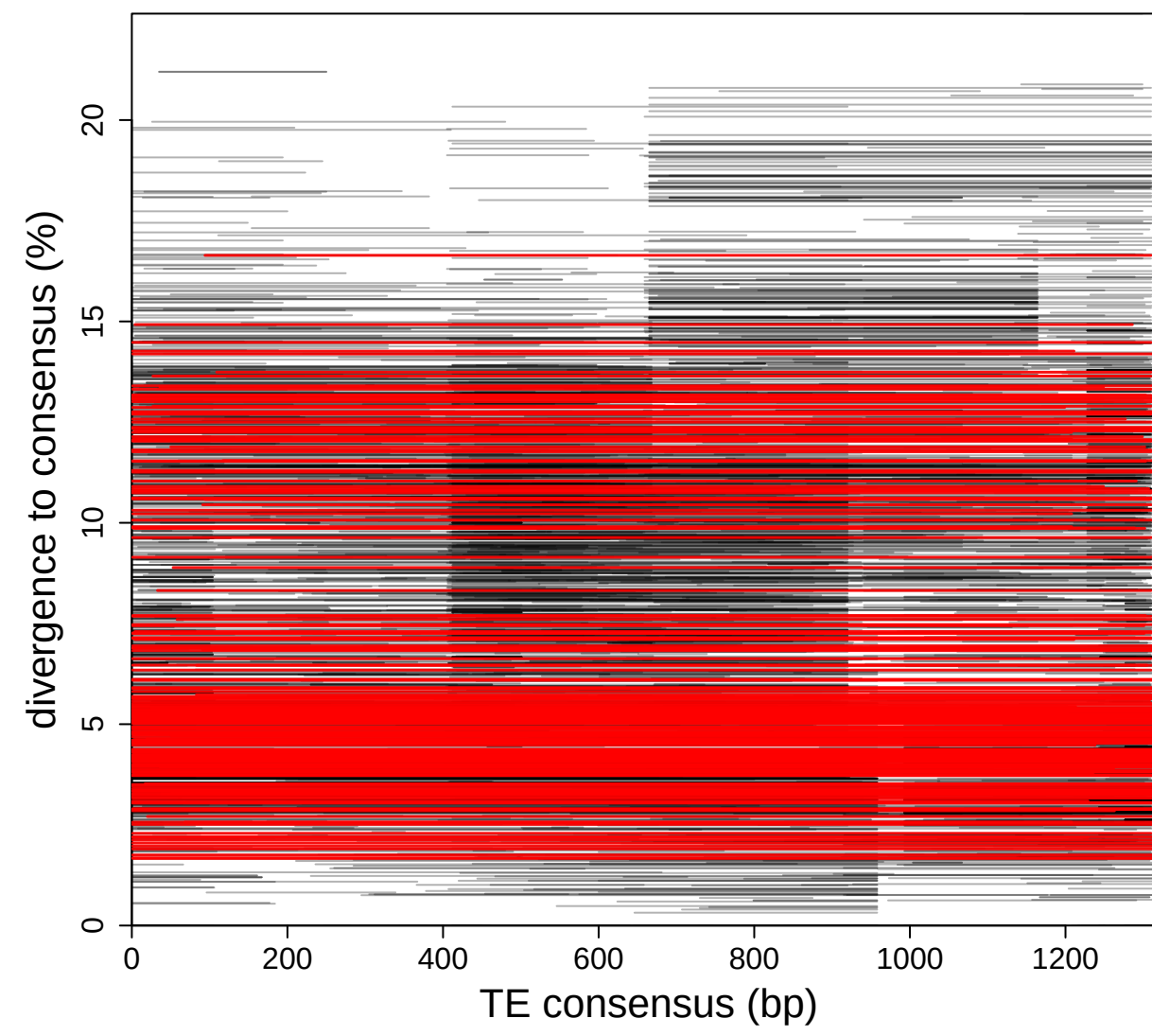
Start crop Point

End crop Point



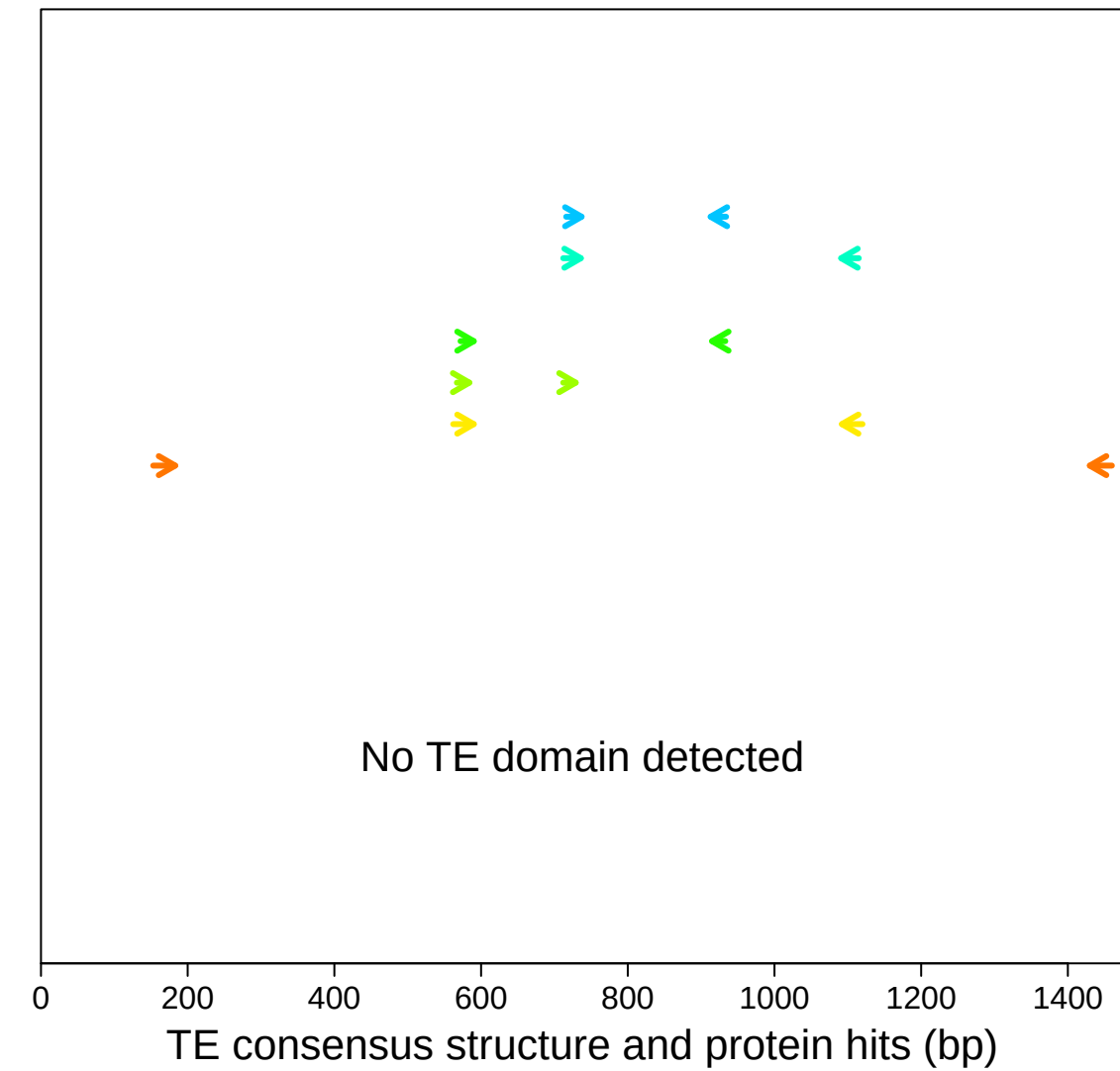
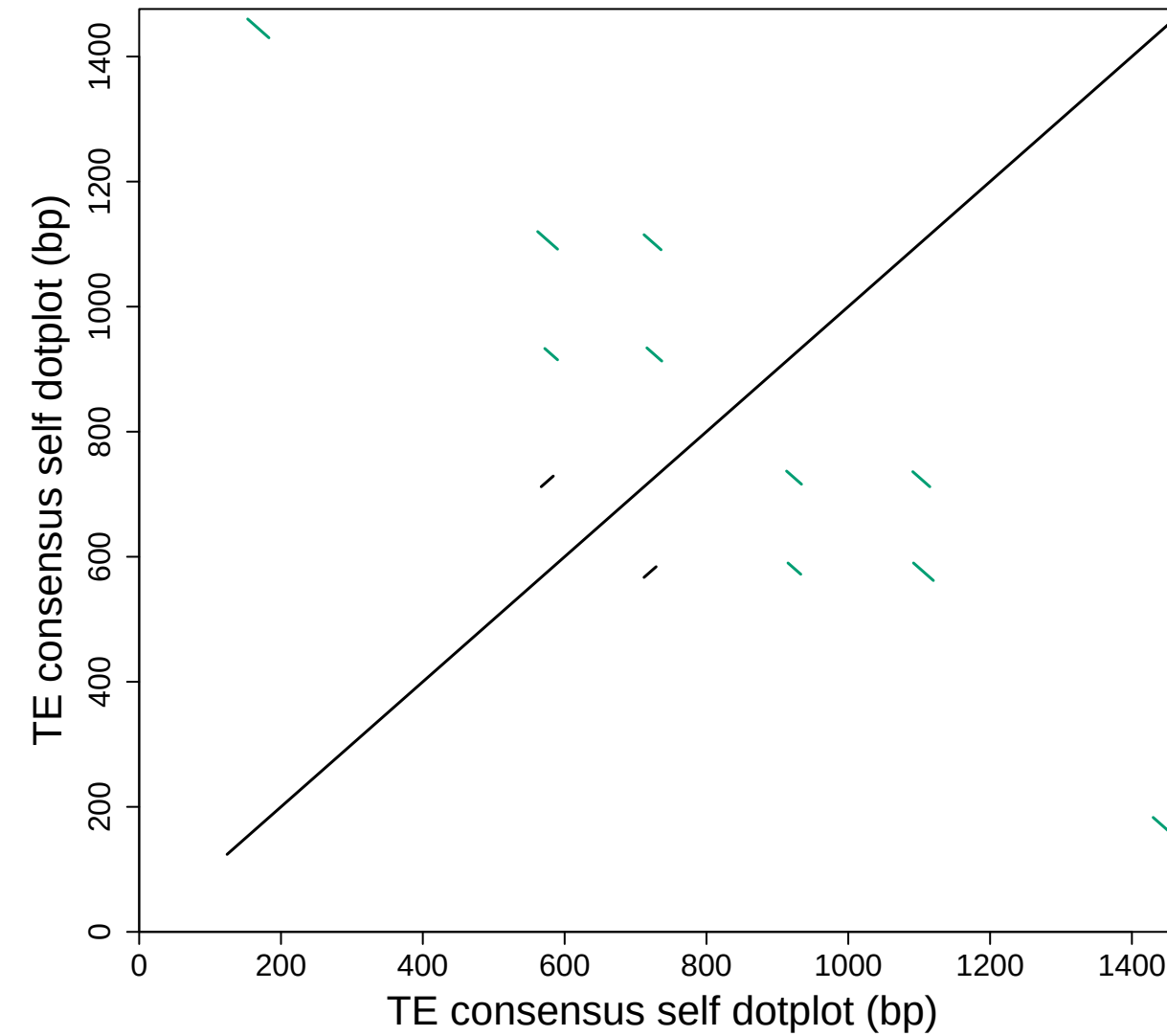
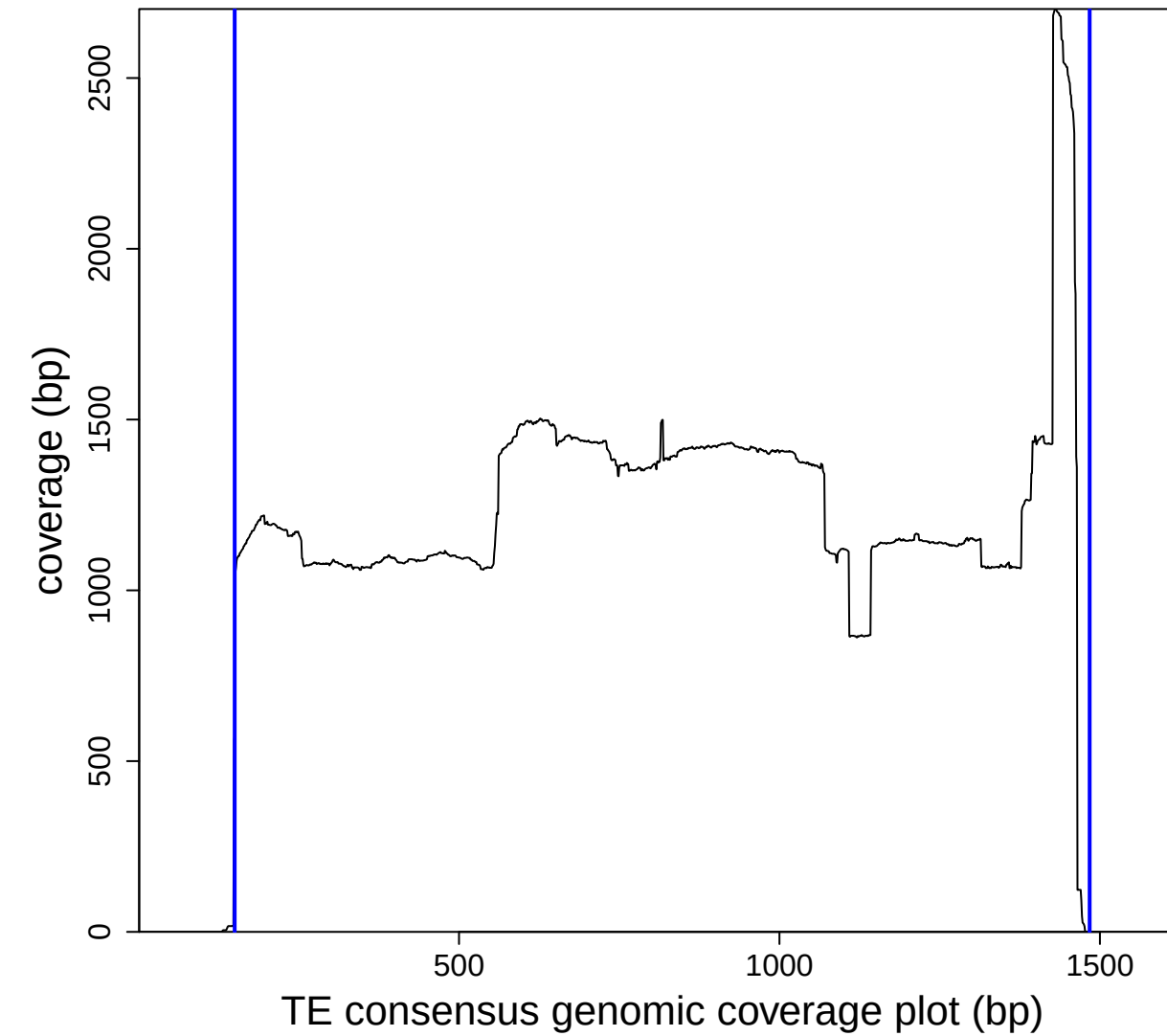
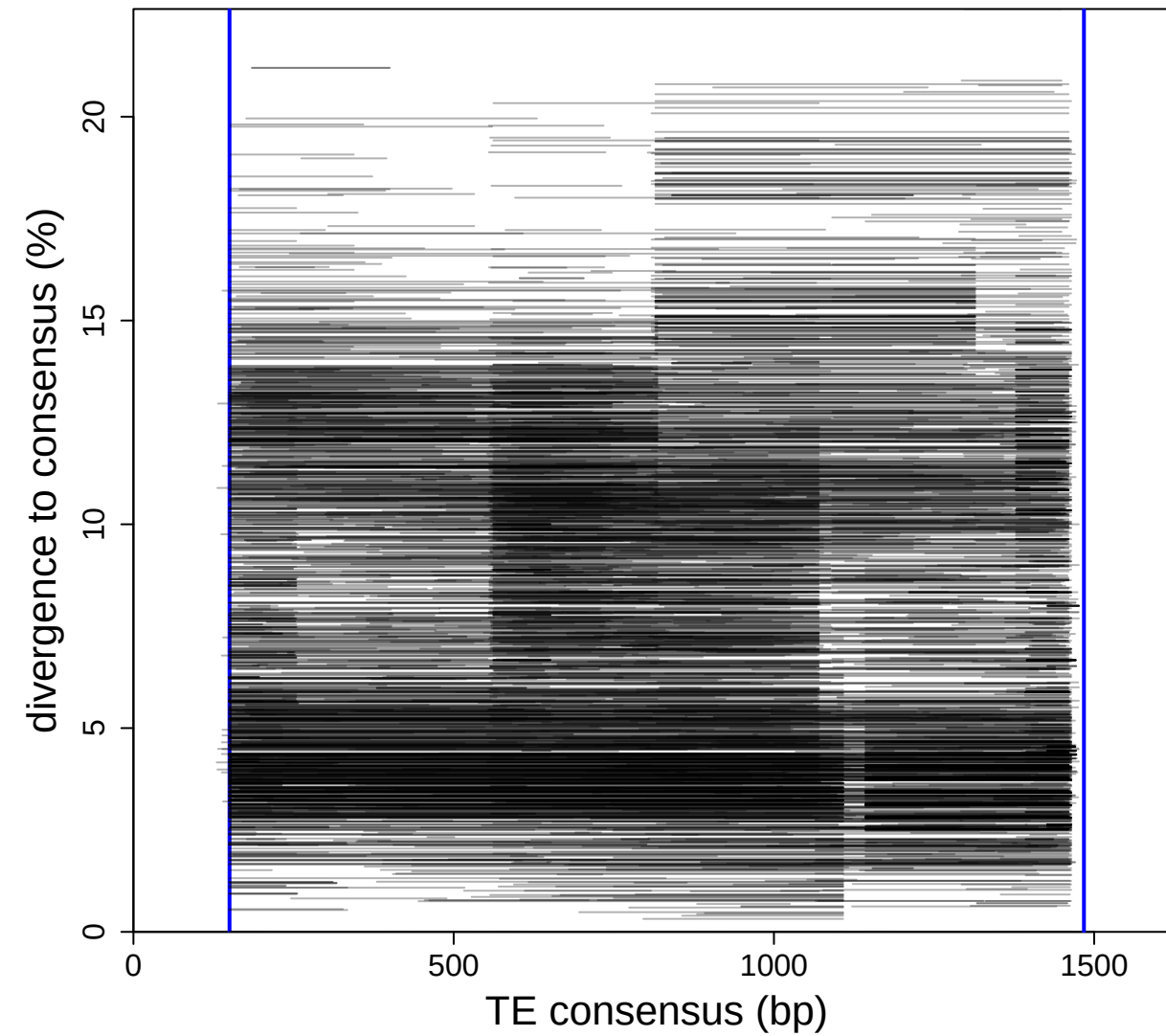
l.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 1329bp; fragments: 5148; full length: 235 (≥ 1196.1 bp)

After TETrimmer 1329 bp



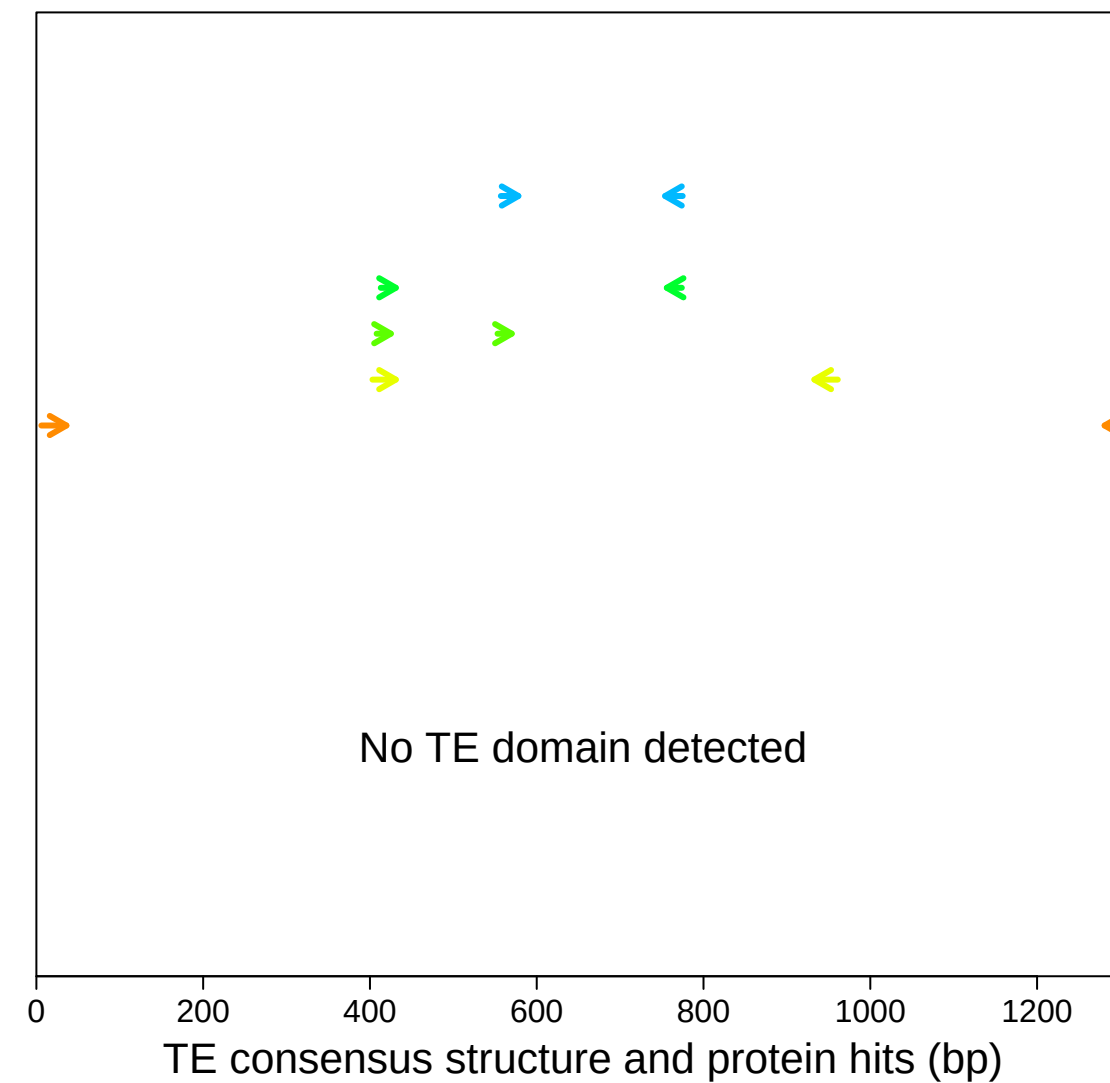
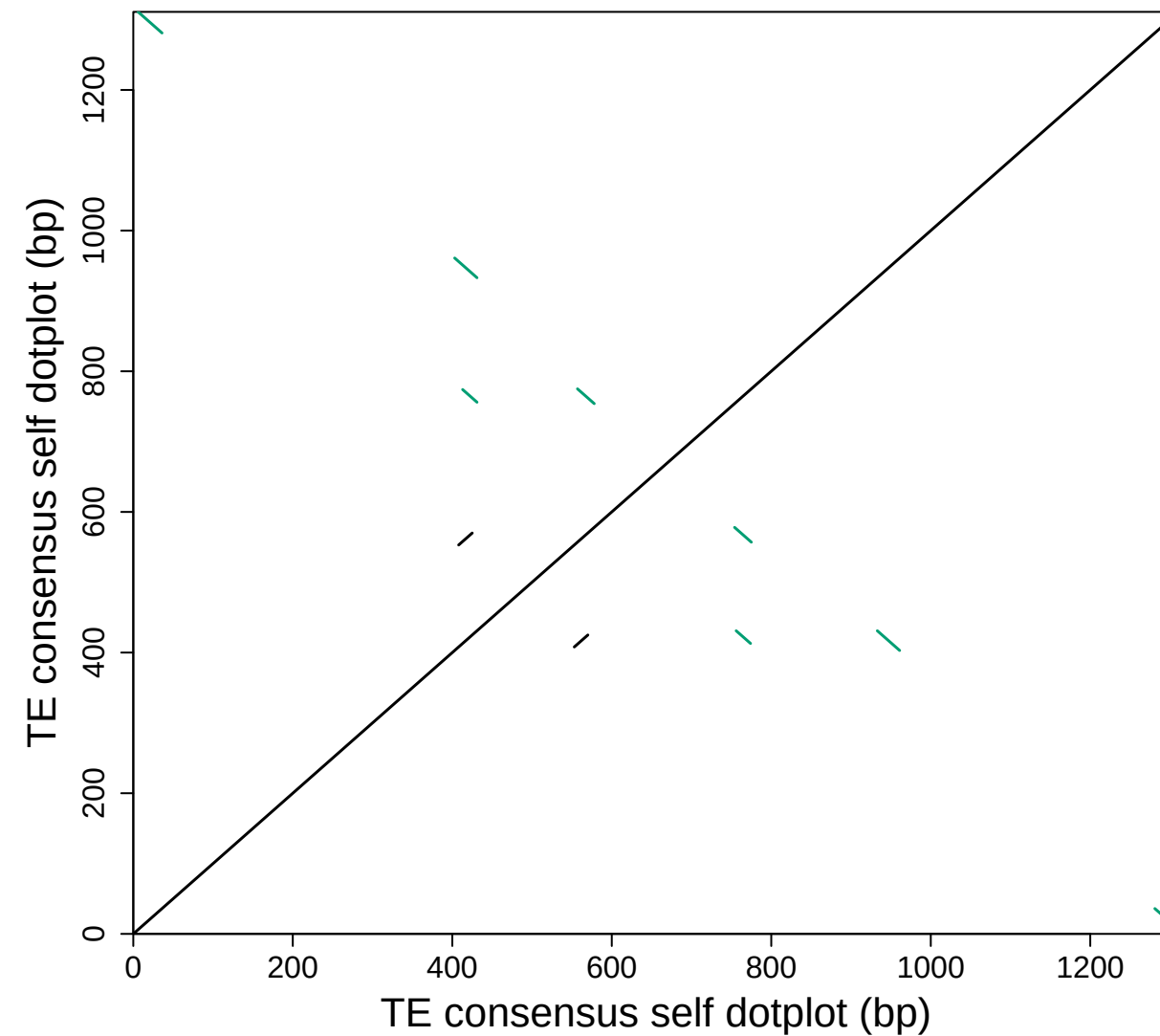
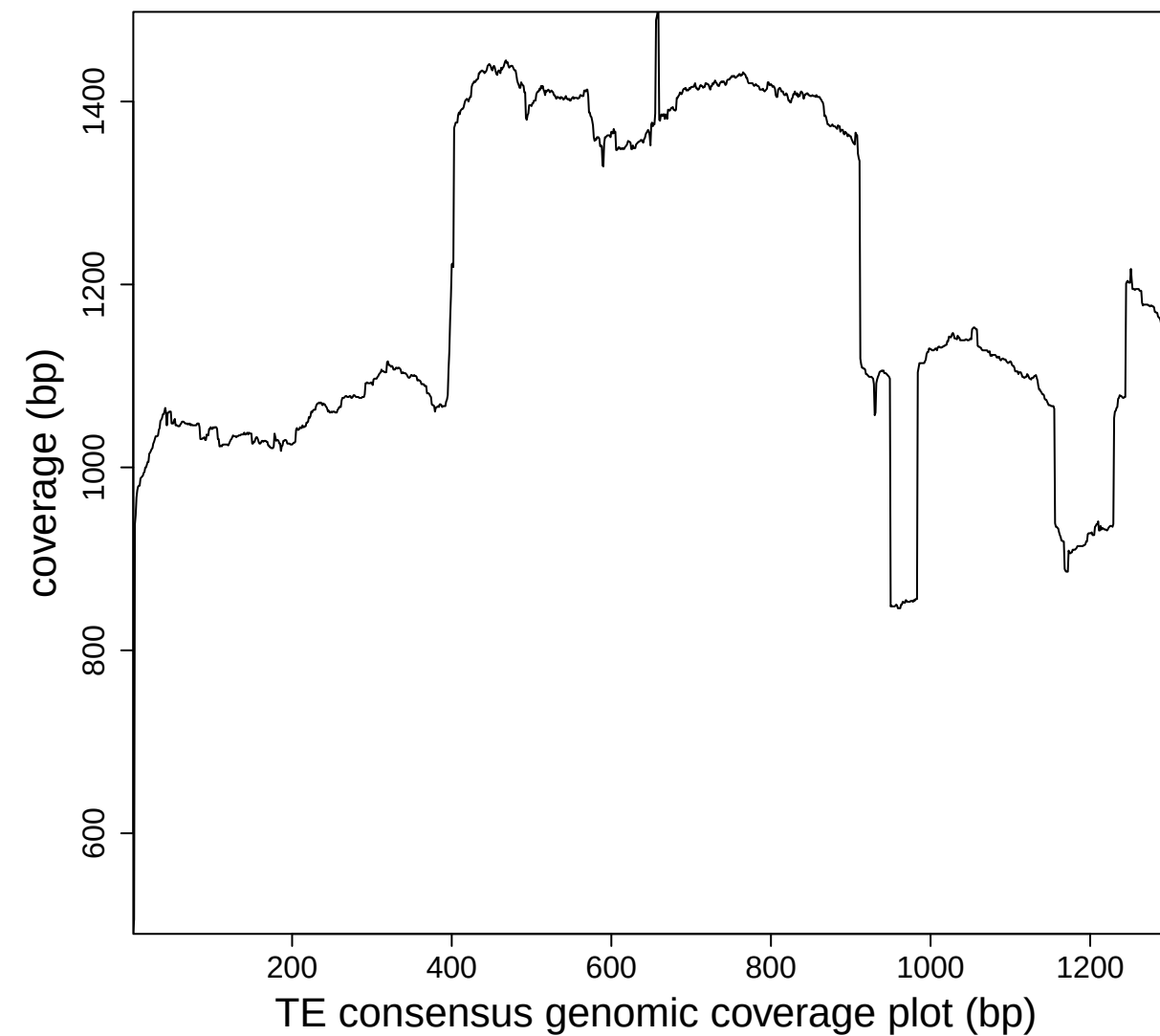
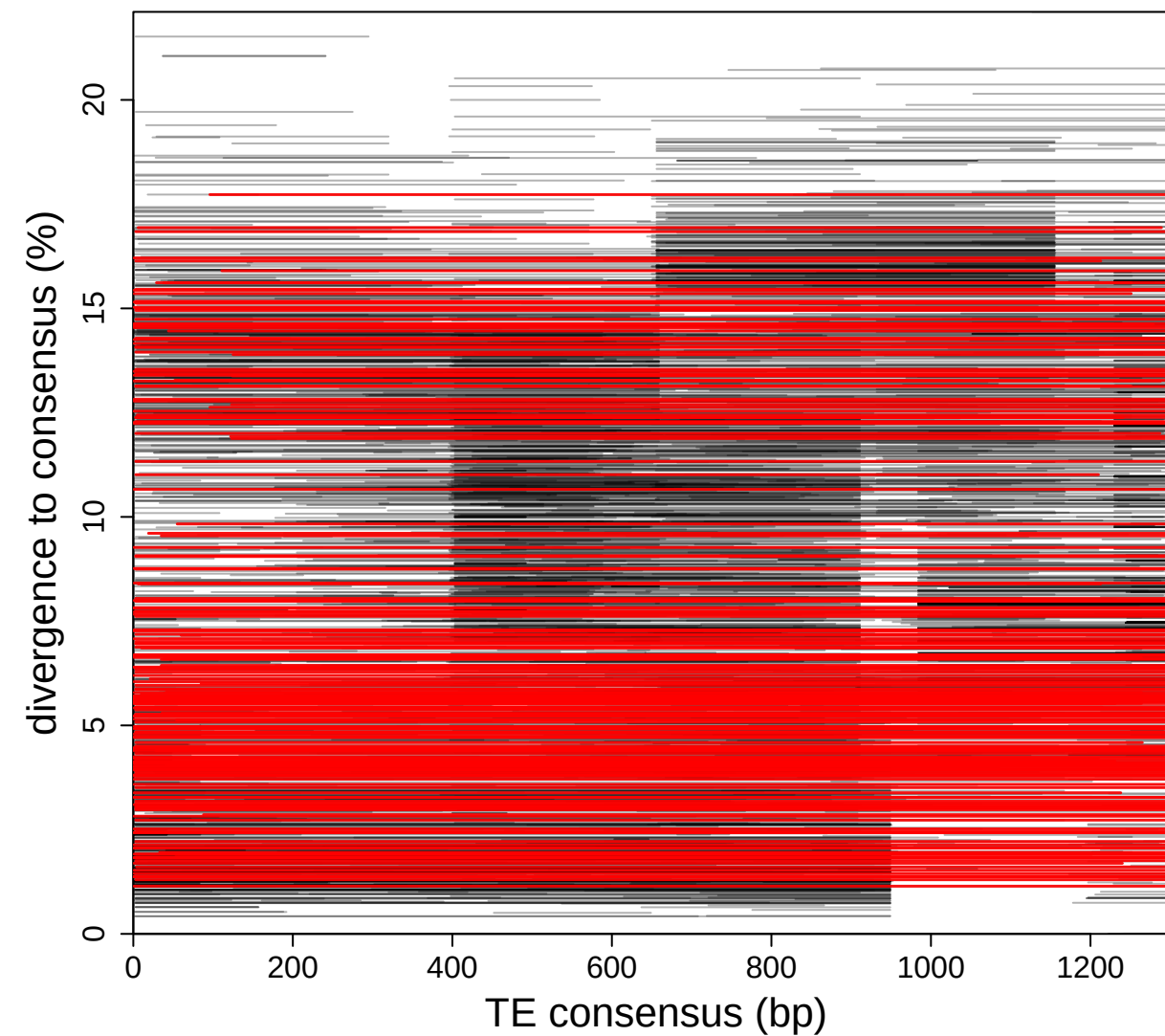
9.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 1634bp; fragments: 5147; full length: 0 (>=1470.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

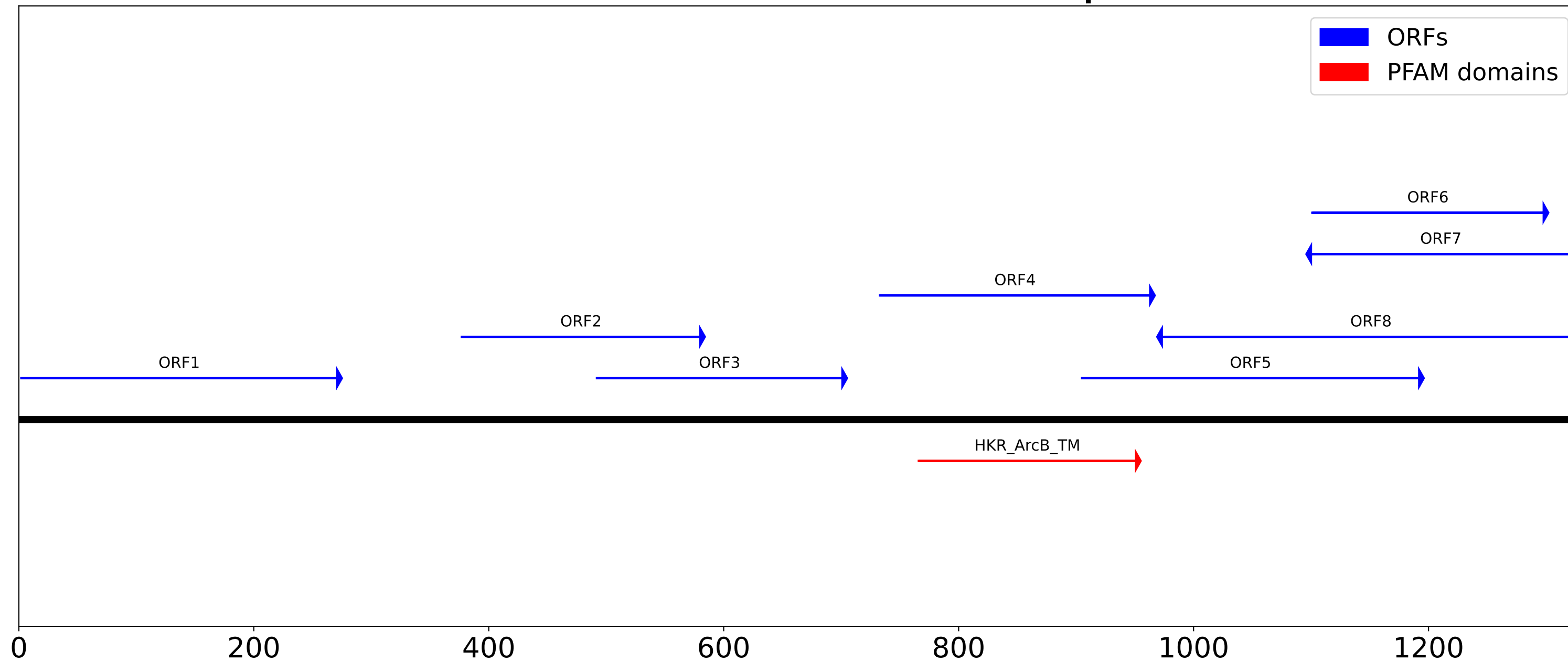


TE: rnd_1_family_259
size: 1311bp; fragments: 3419; full length: 231 (≥ 1179.9 bp)

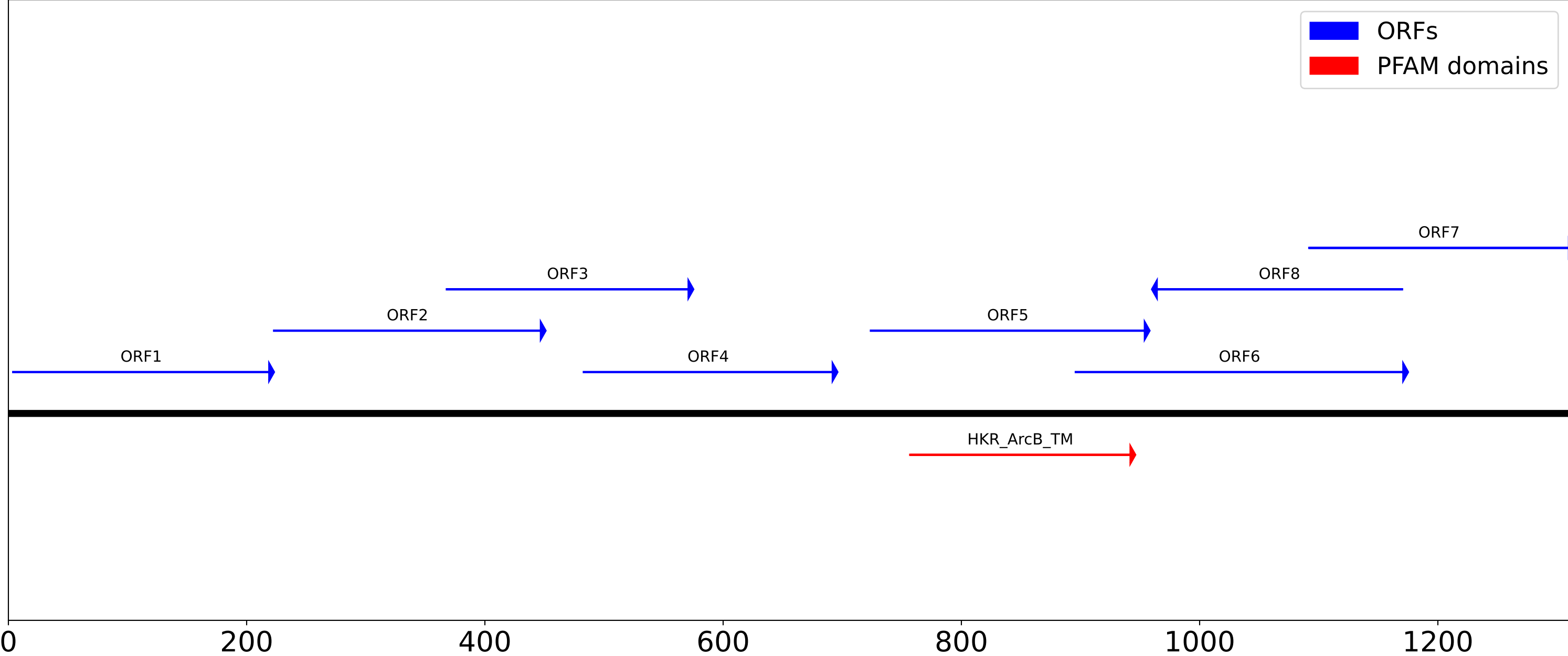
Before TEtrimmer 1311 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

