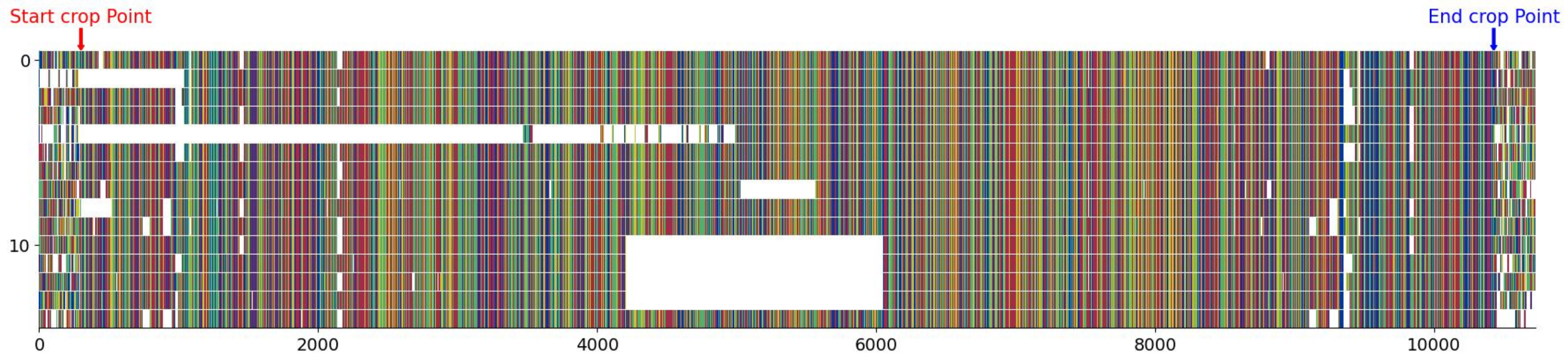
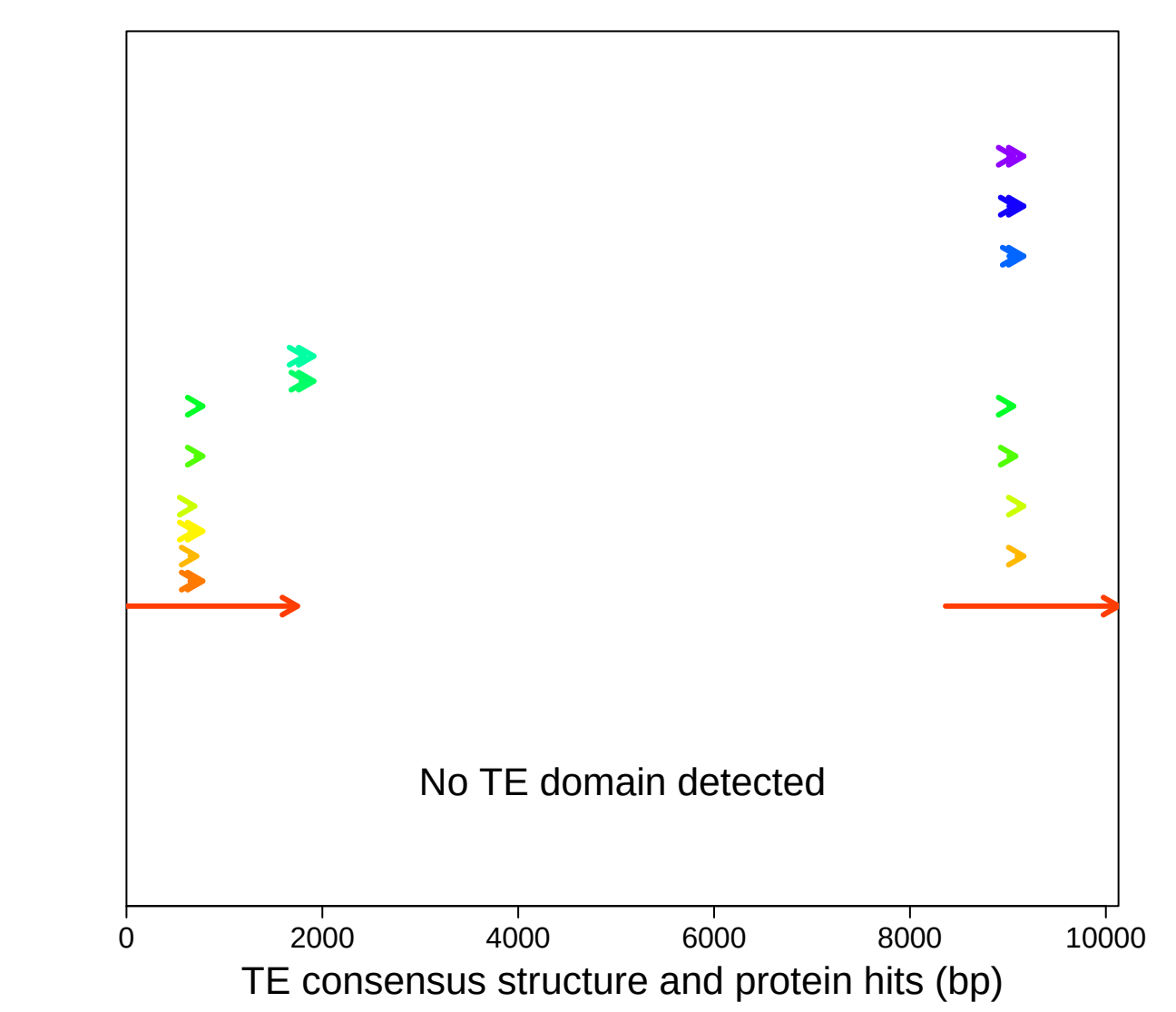
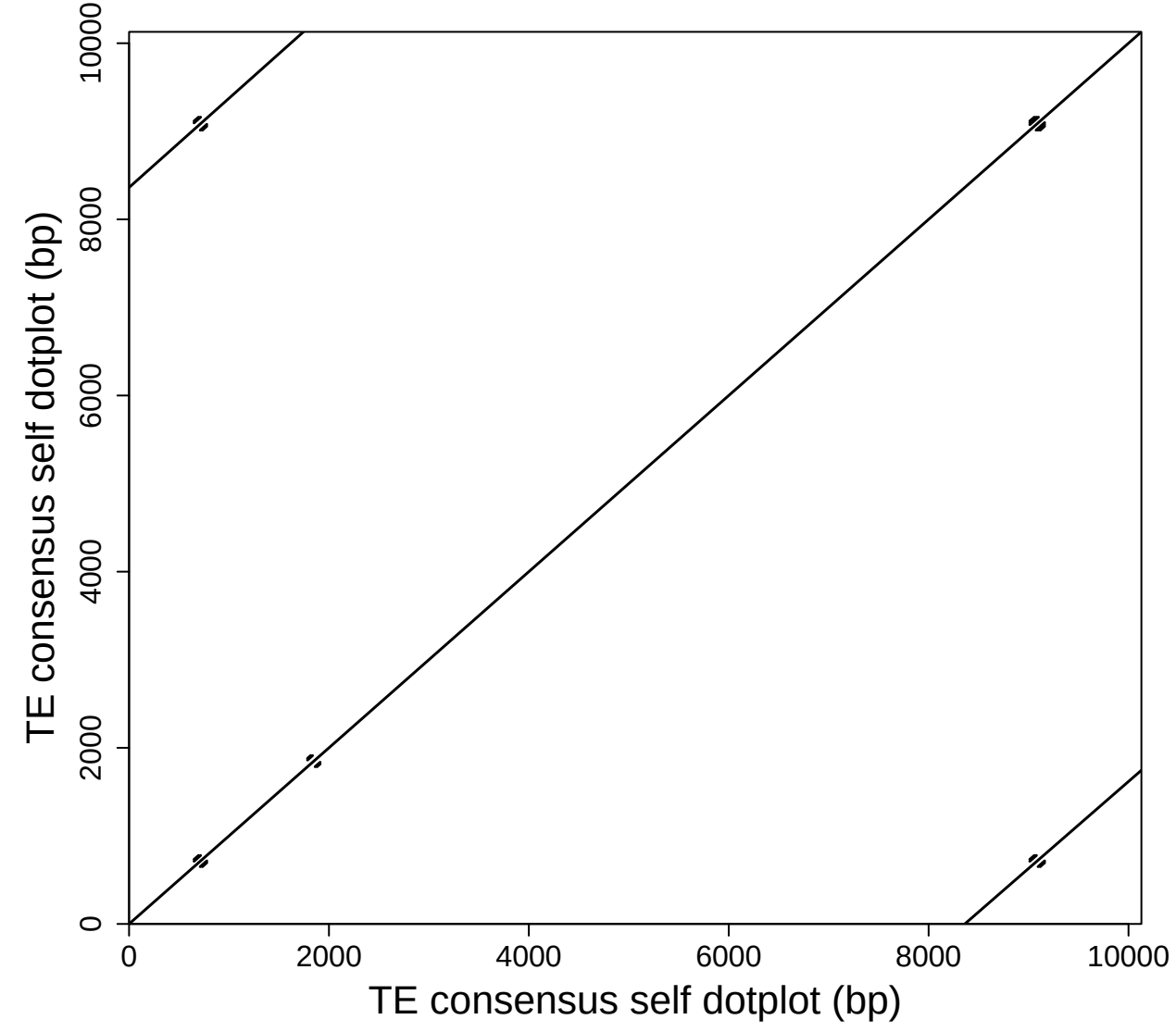
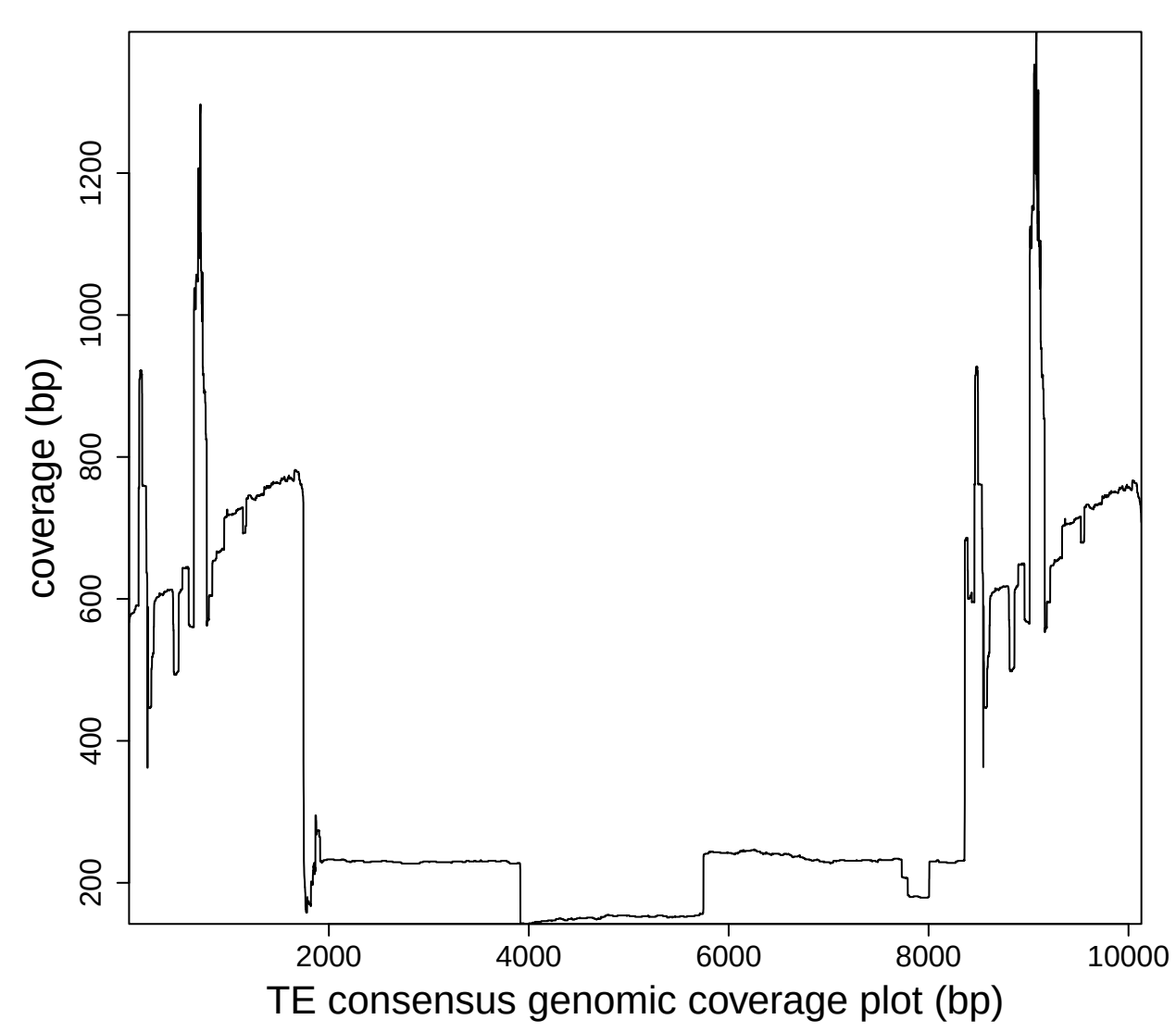
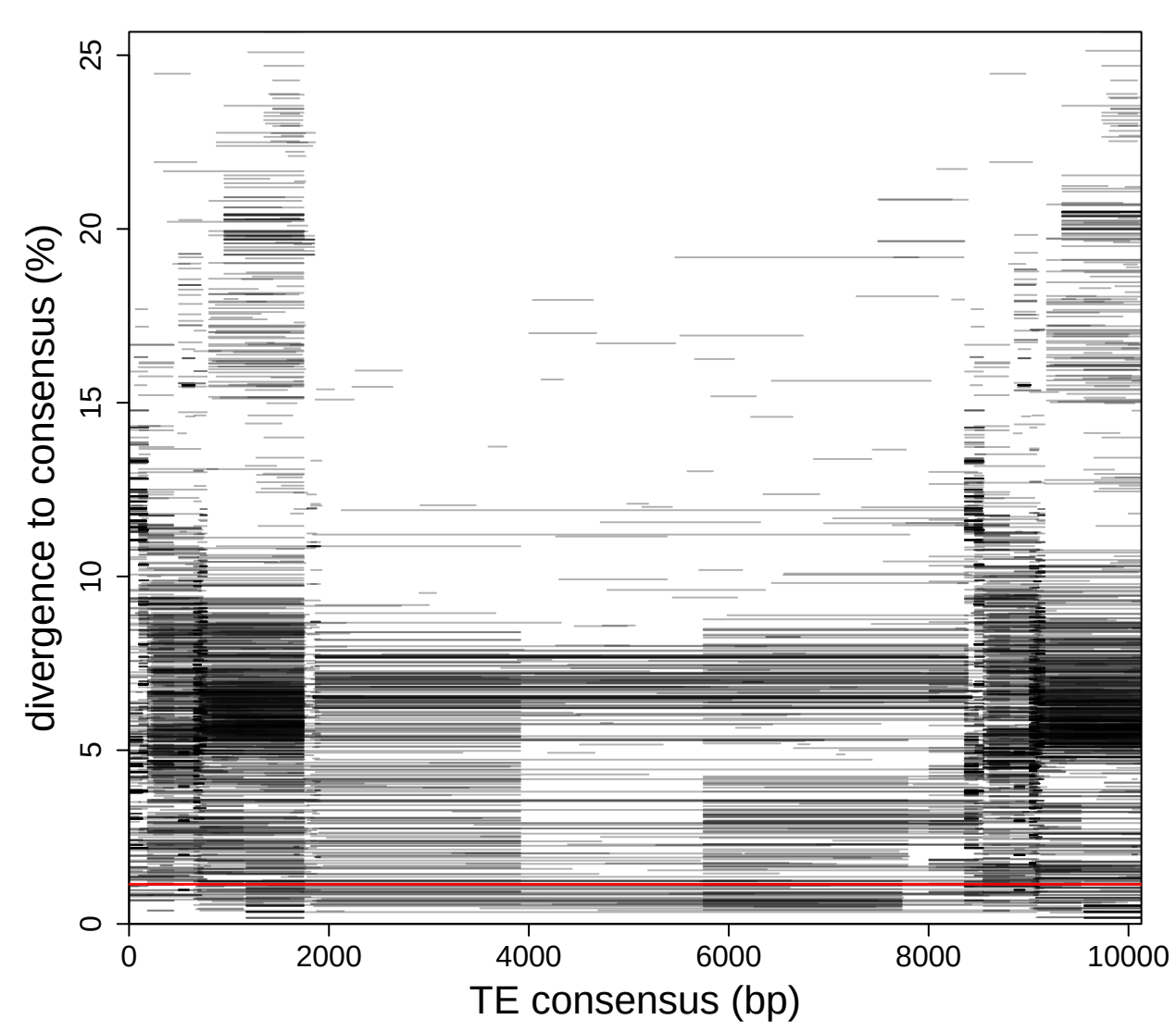
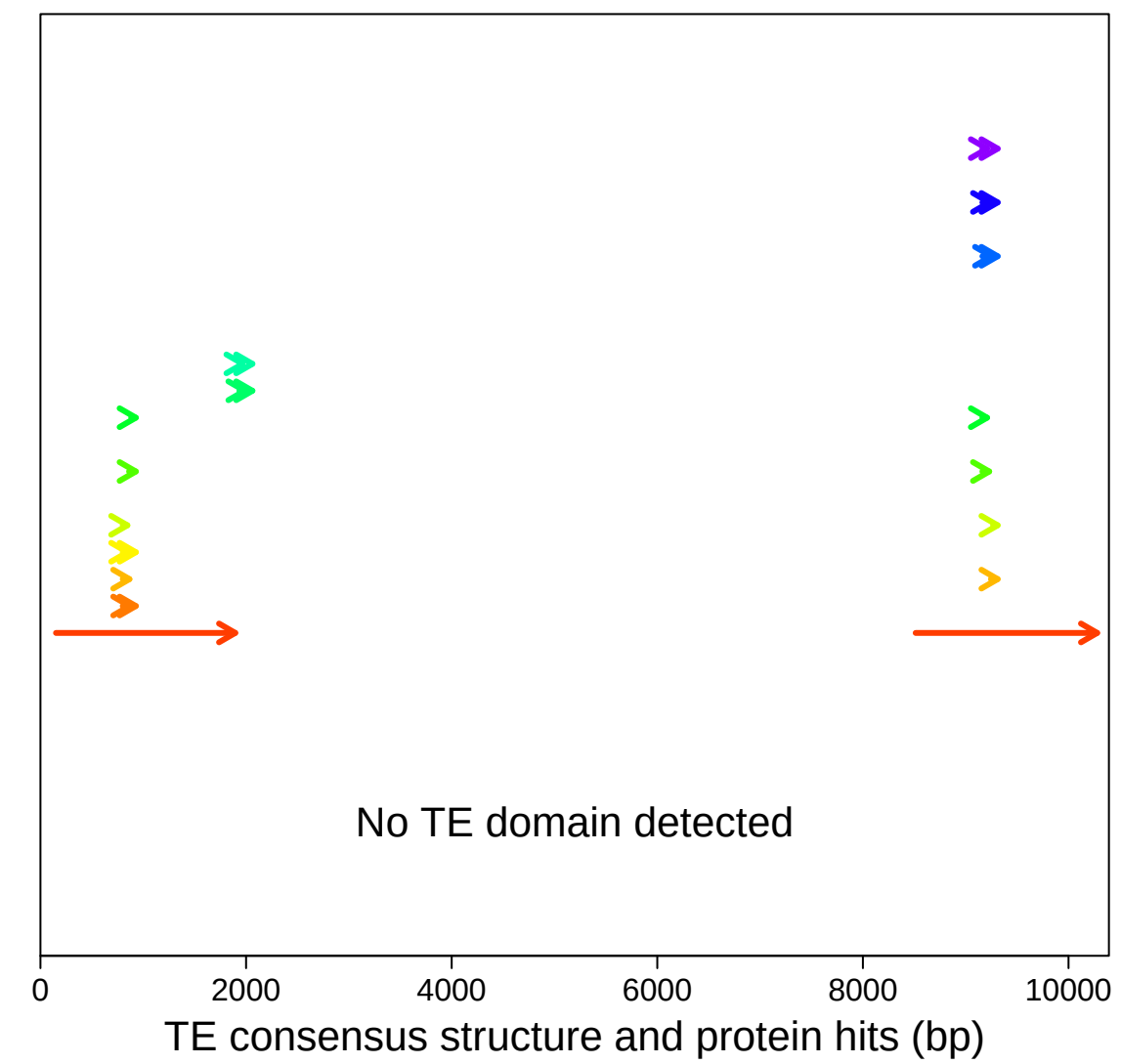
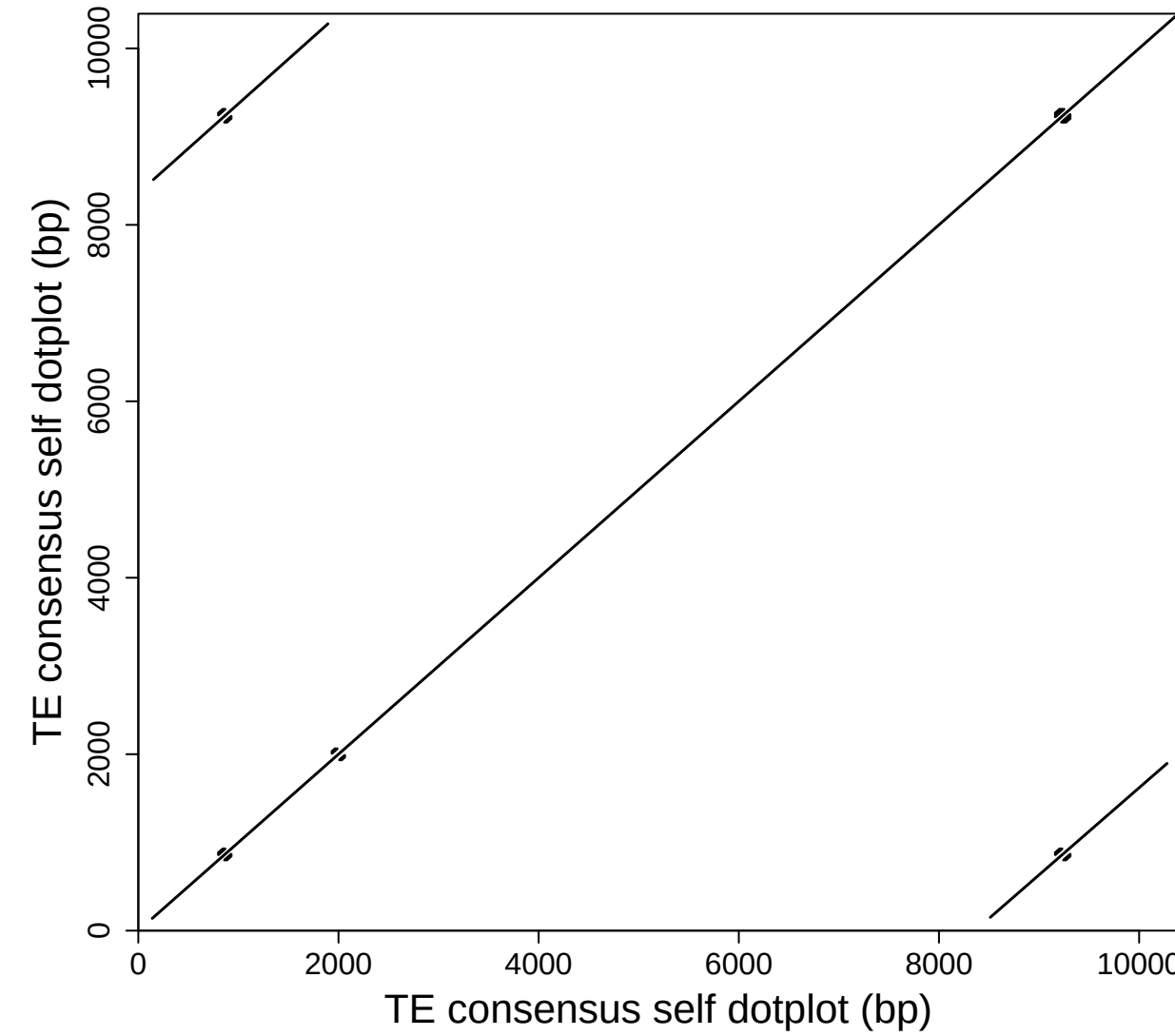
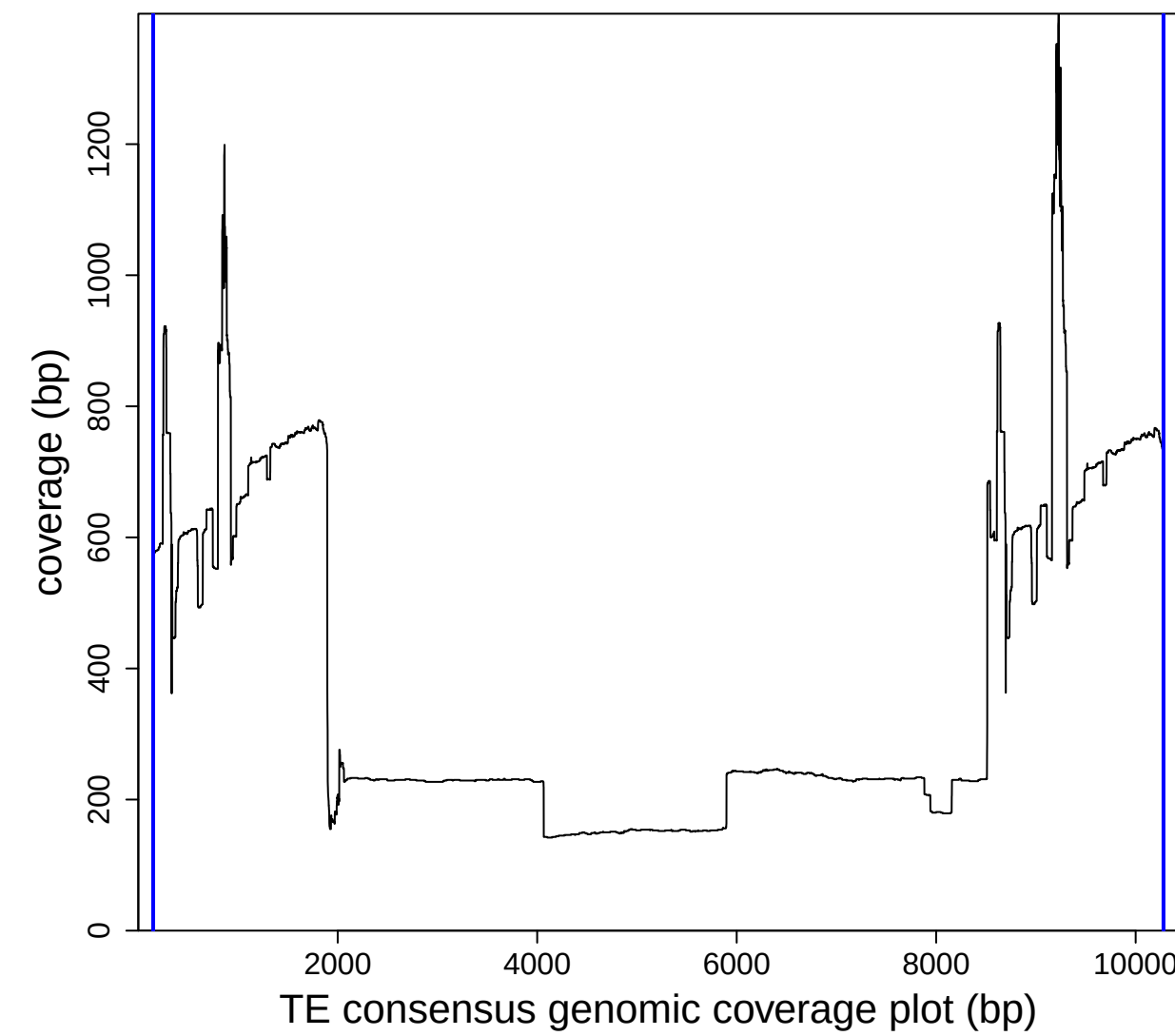
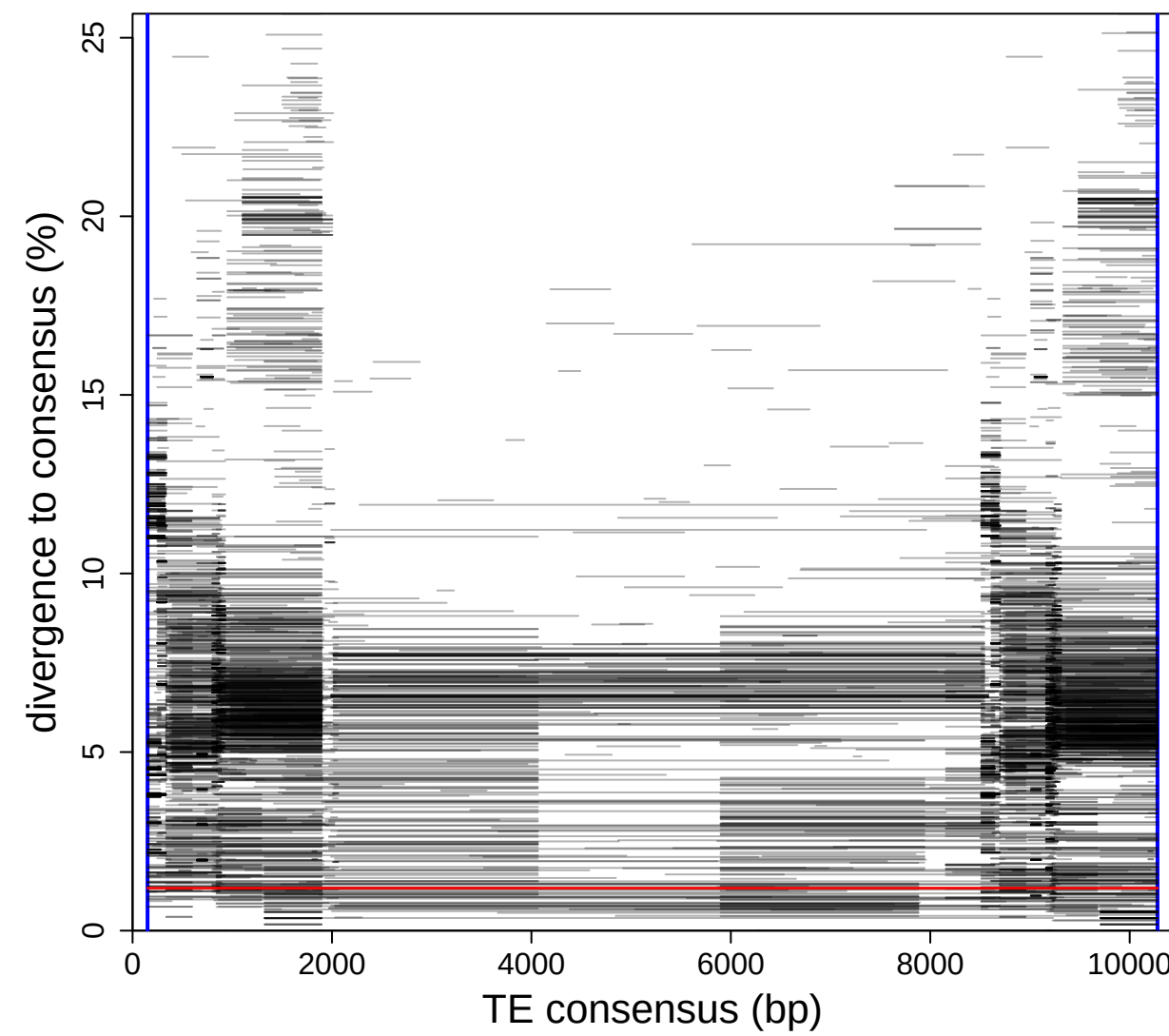






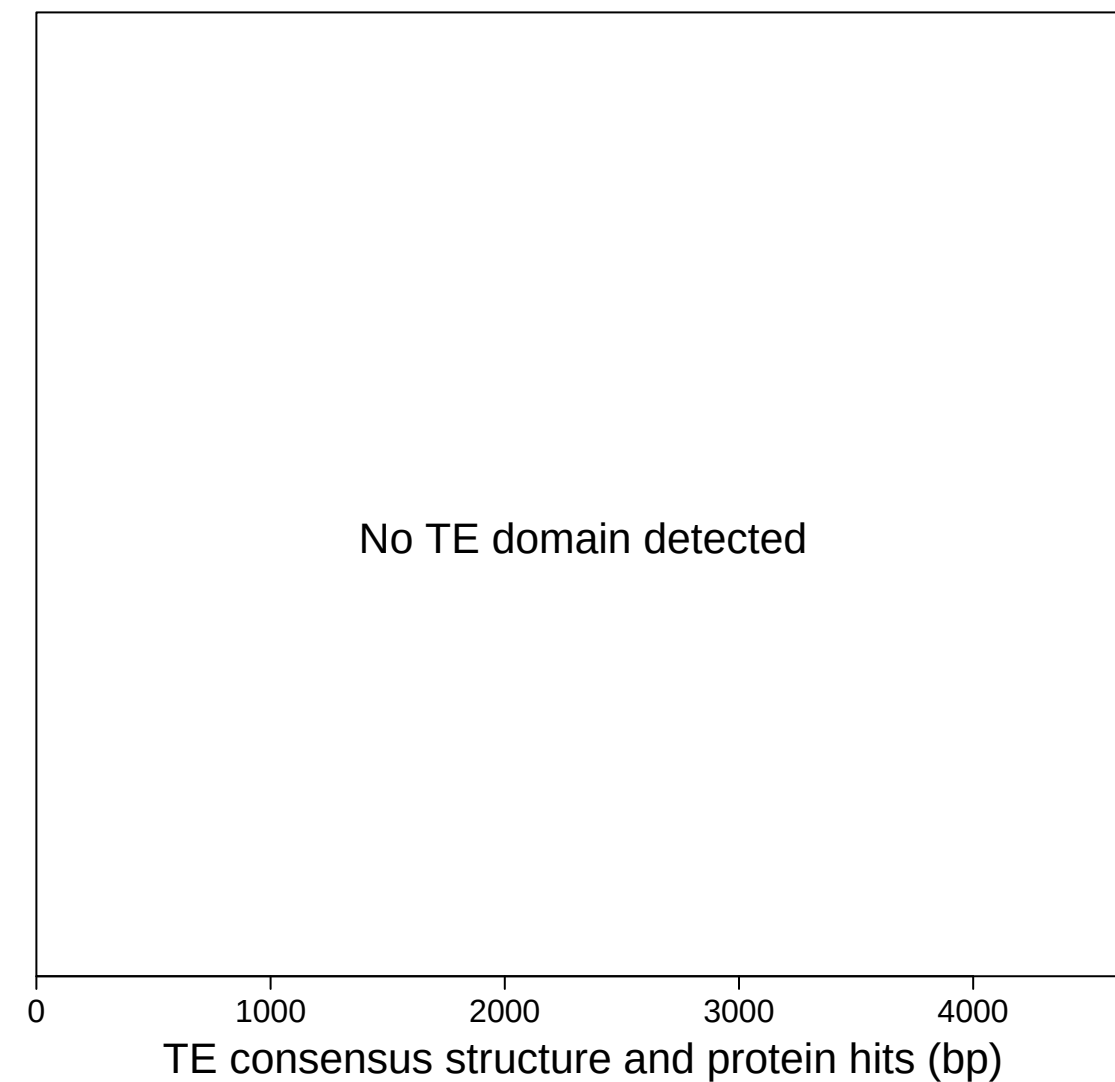
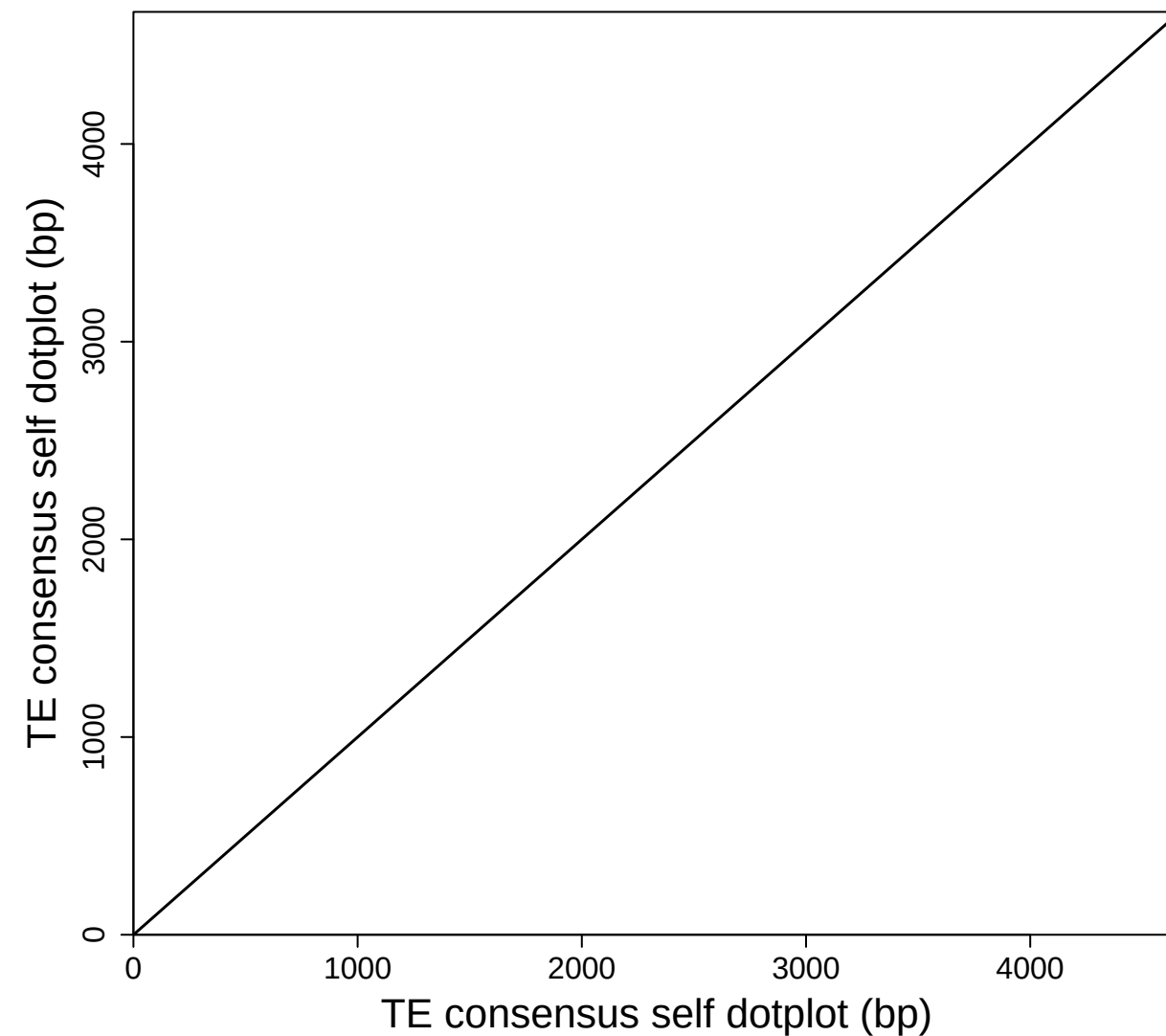
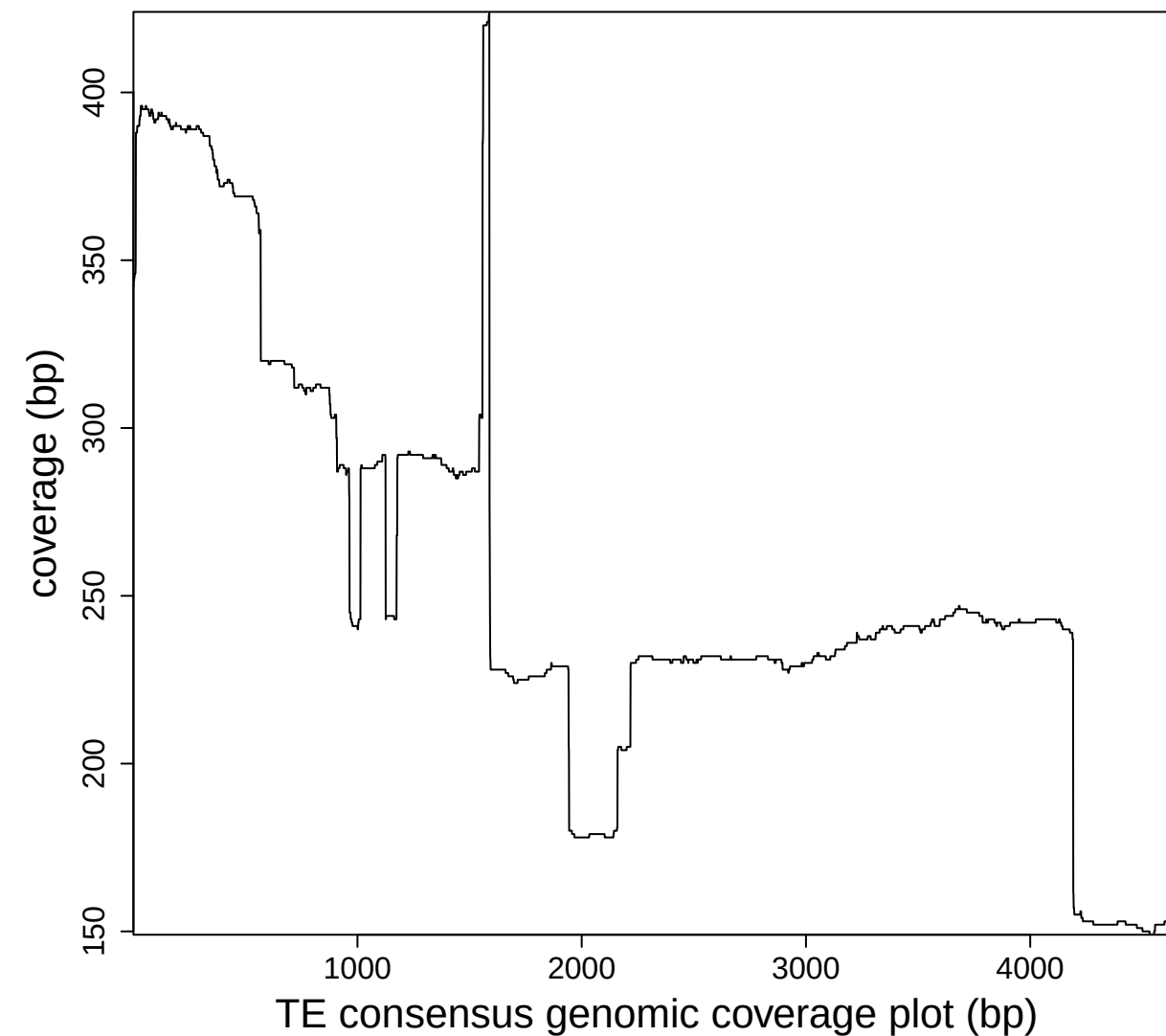
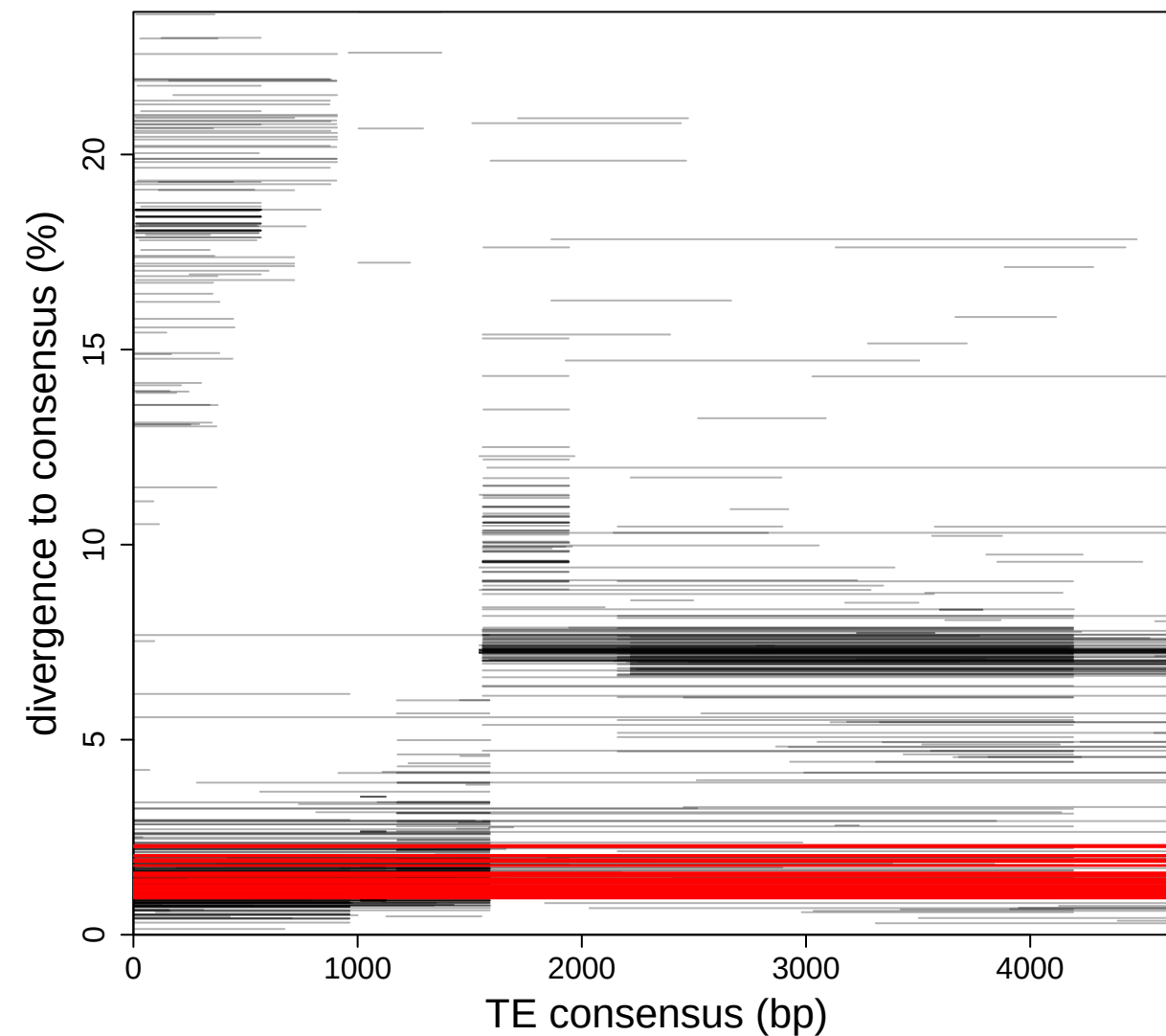
617.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa c
size: 10429bp; fragments: 6390; full length: 1 (>=9386.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

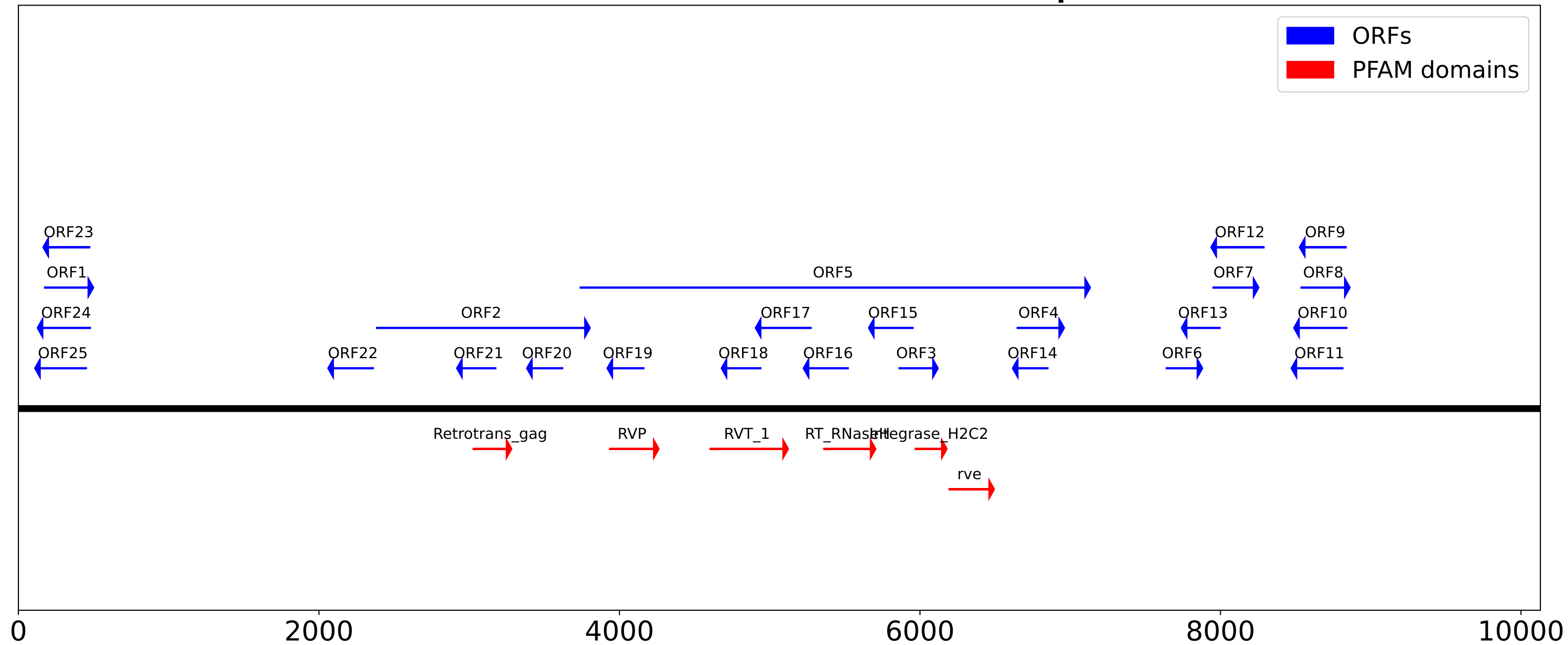


TE: rnd_1_family_617
size: 4668bp; fragments: 801; full length: 39 (≥ 4201.2 bp)

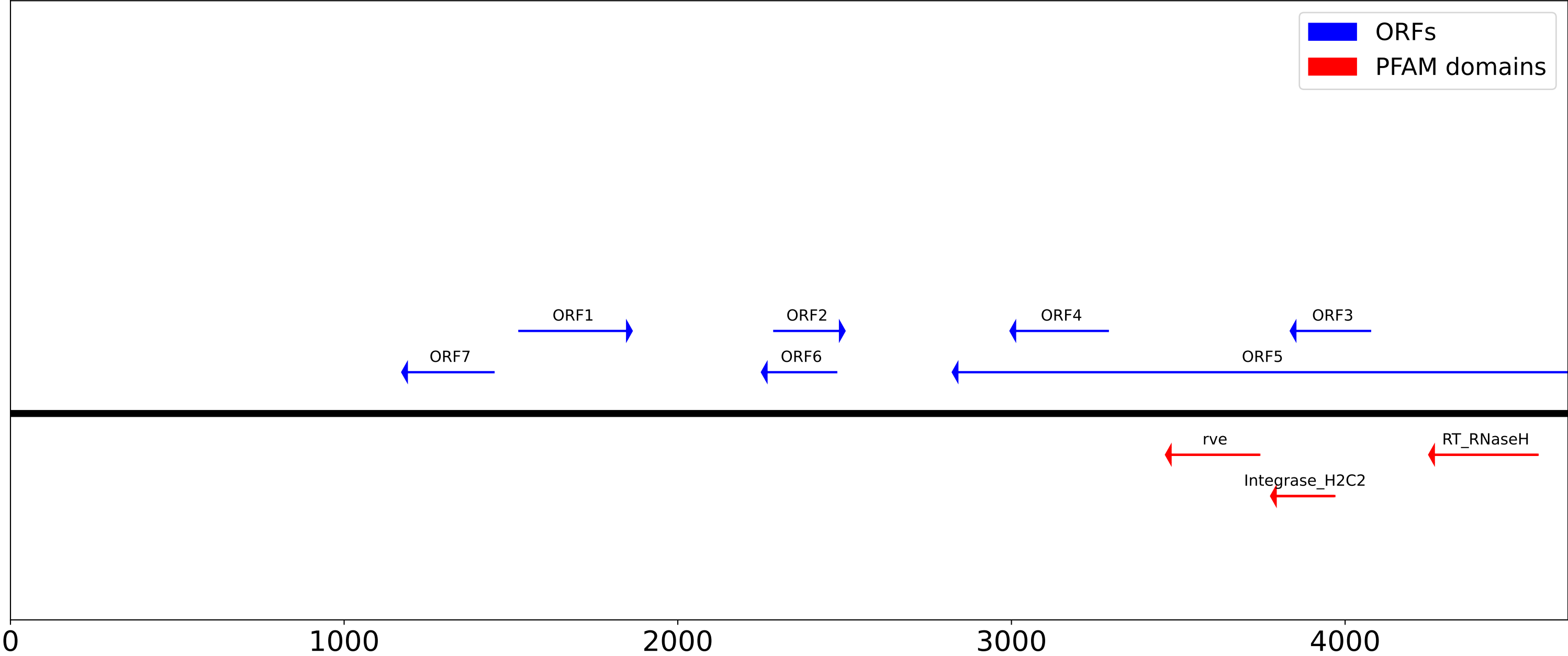
Before TEtrimmer 4668 bp



After TEtrimmer ORF and PFAM domain plot



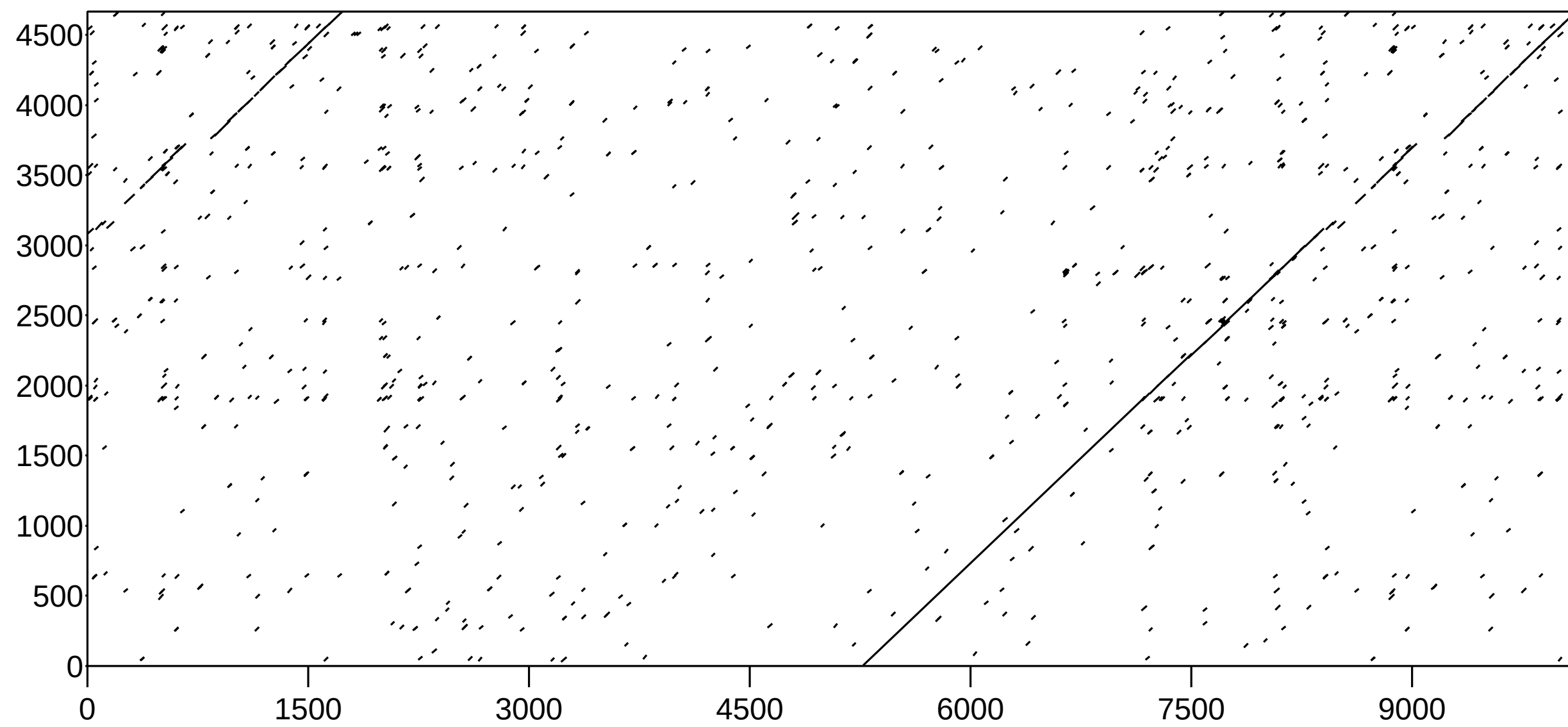
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)