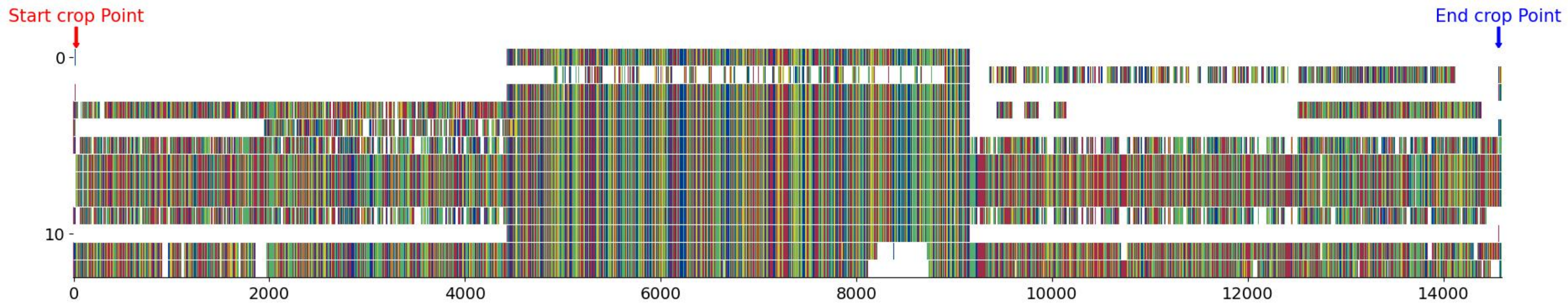


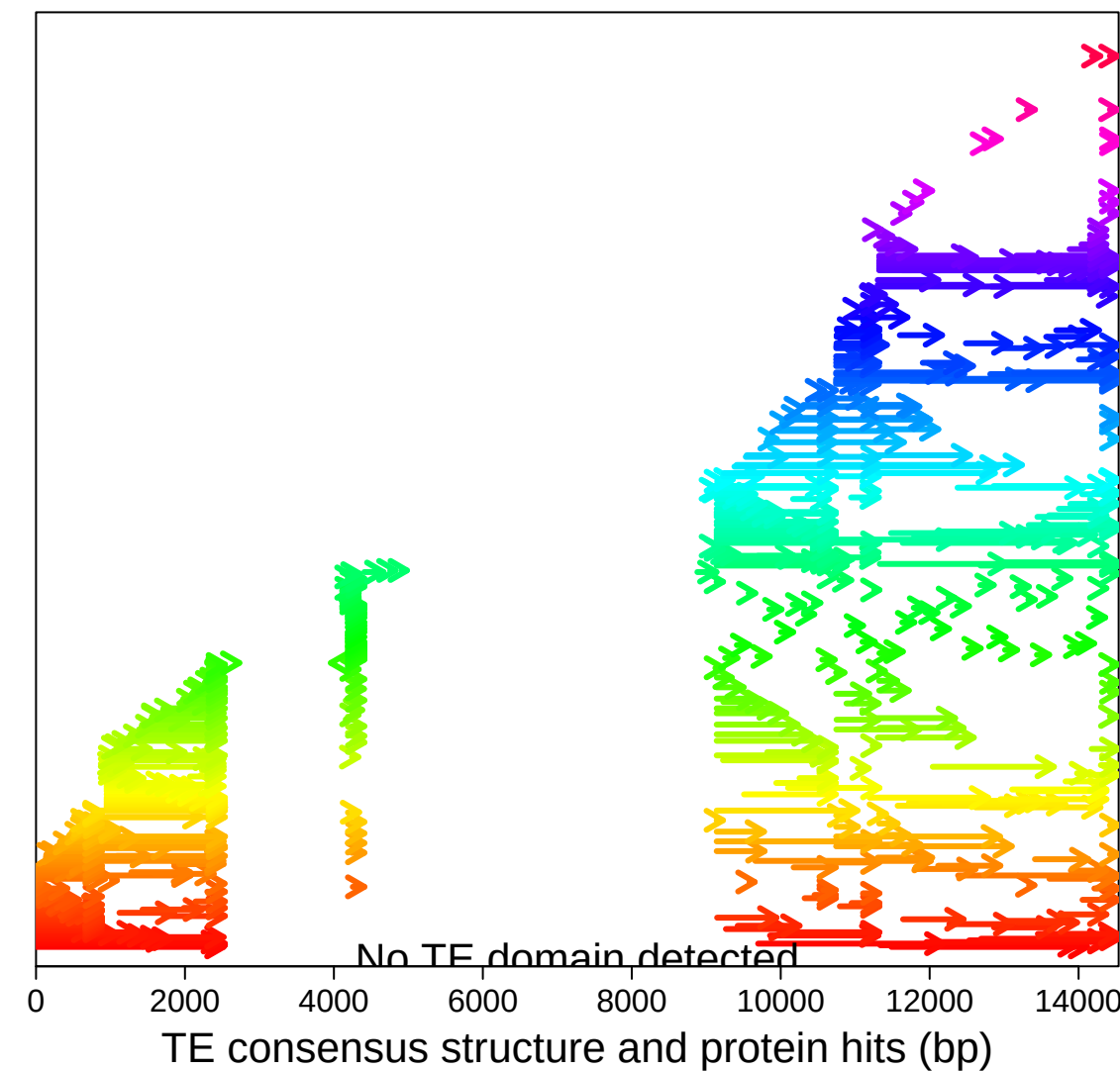
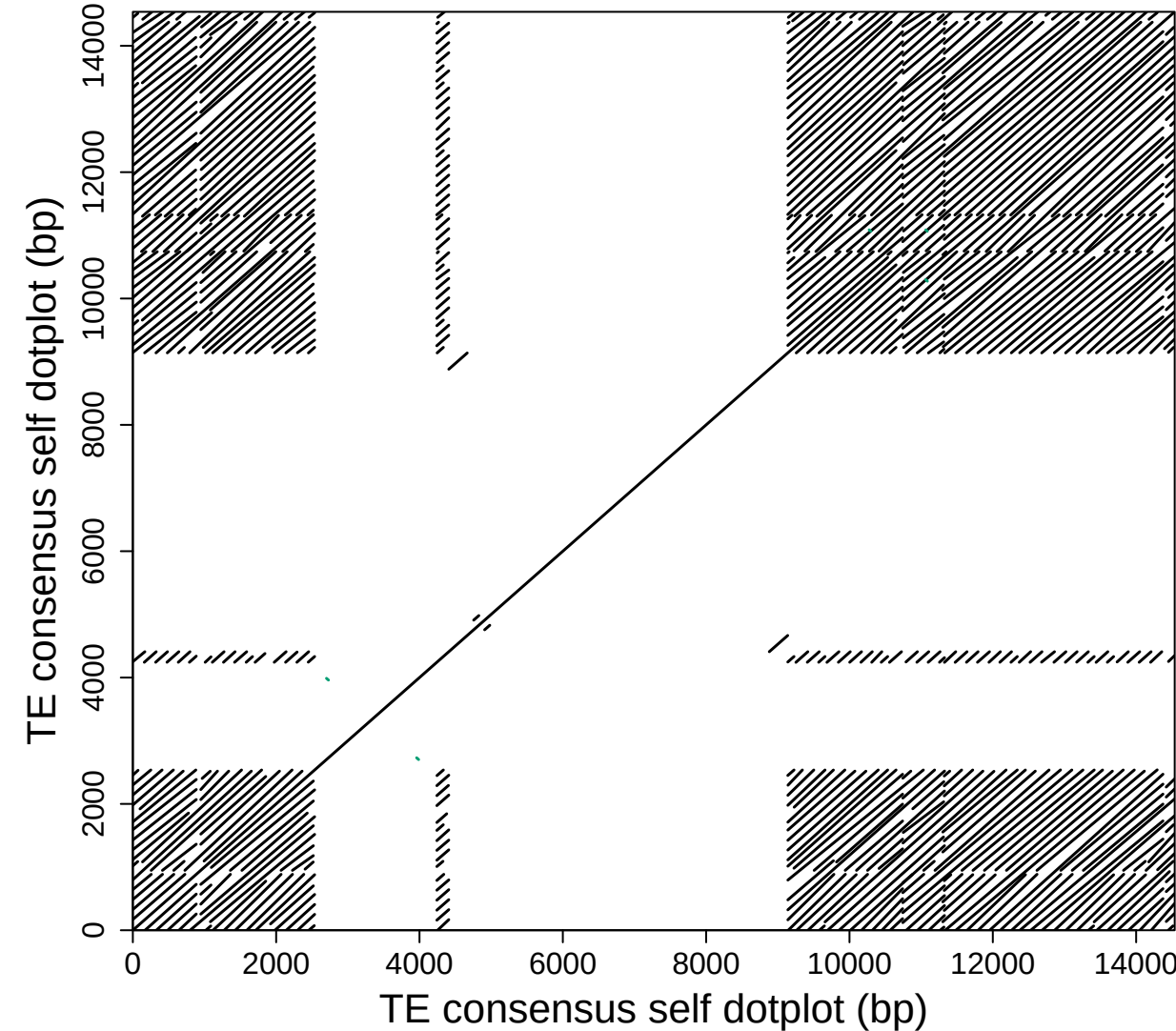
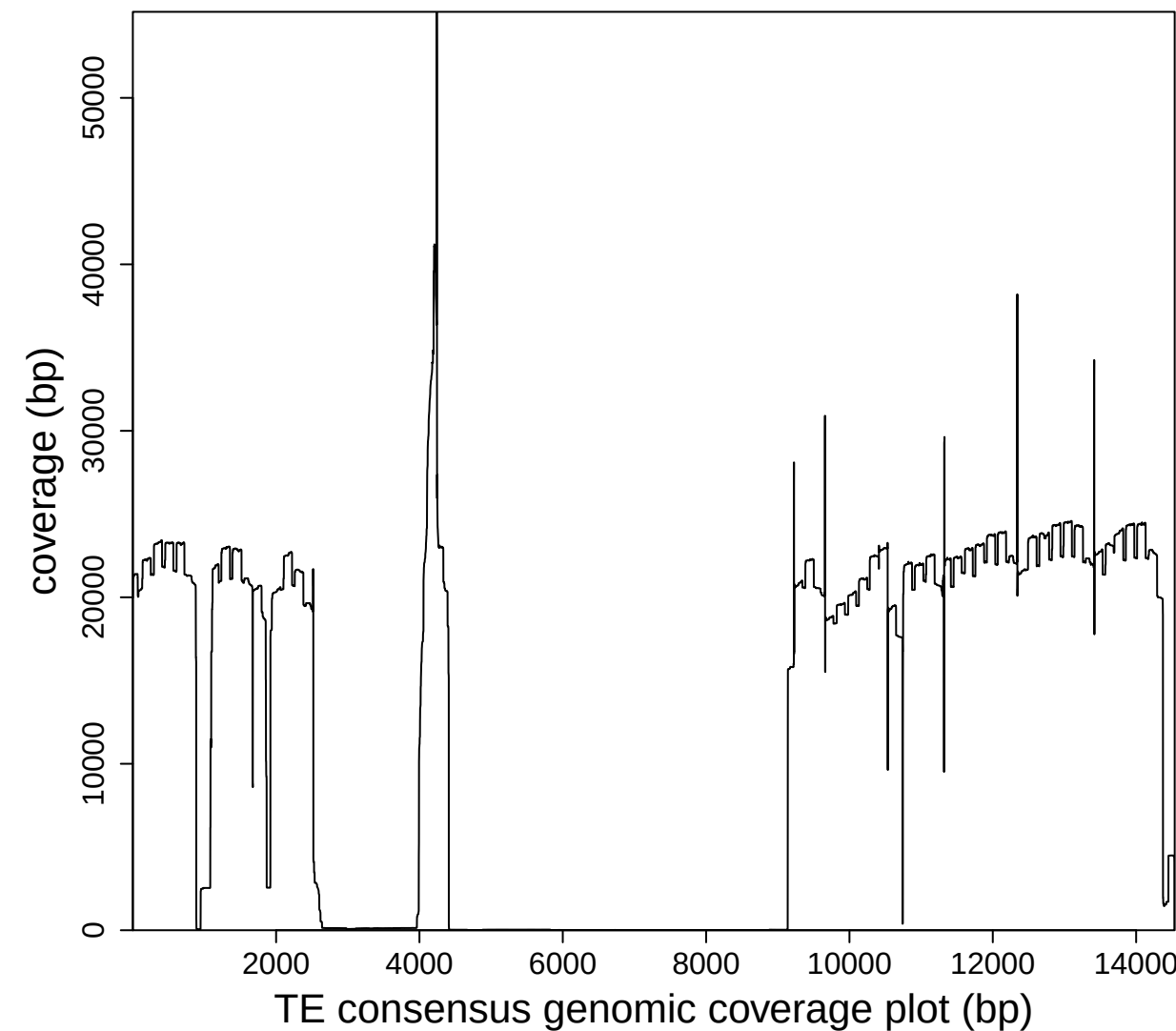
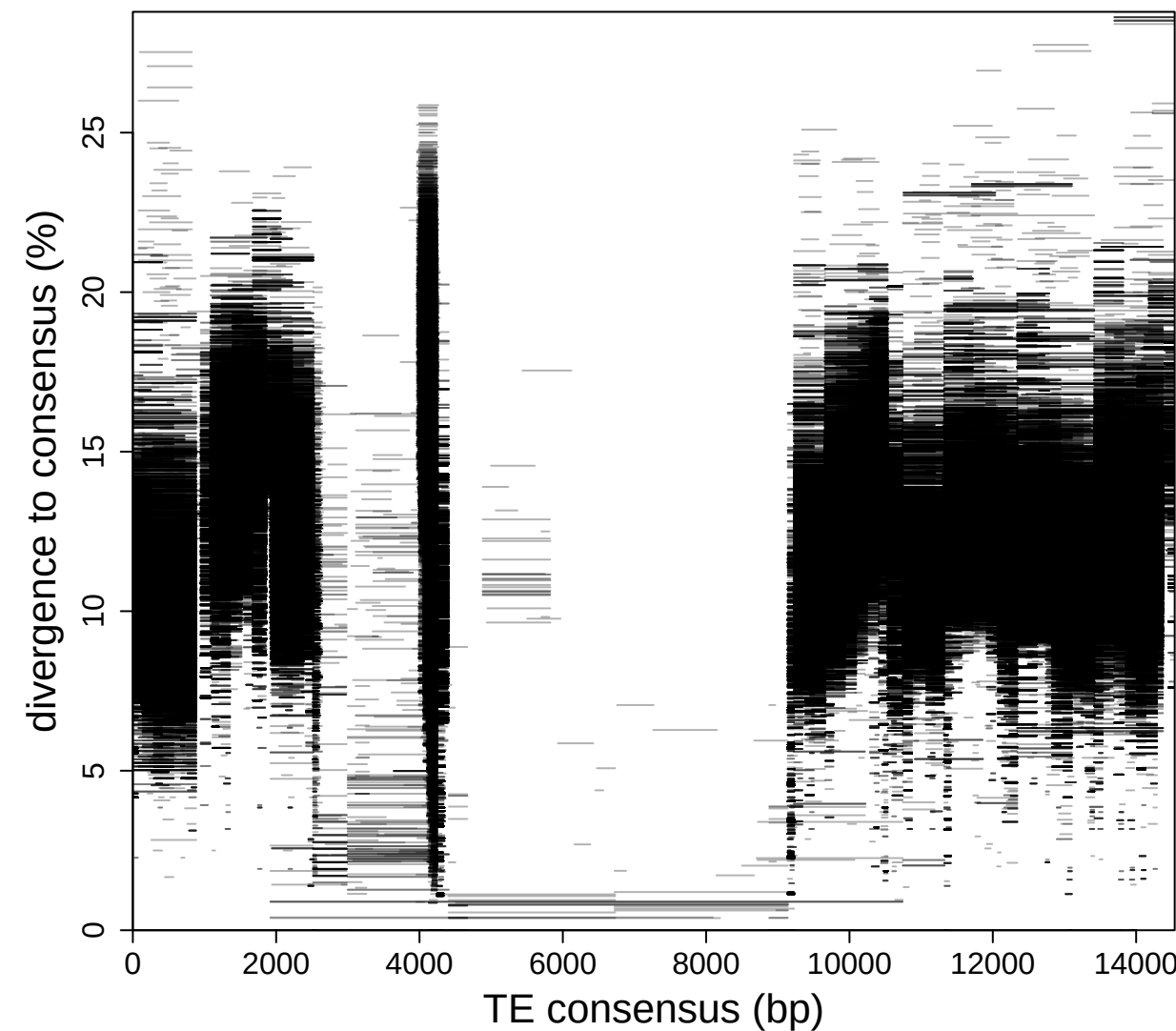
MSA length = 14538





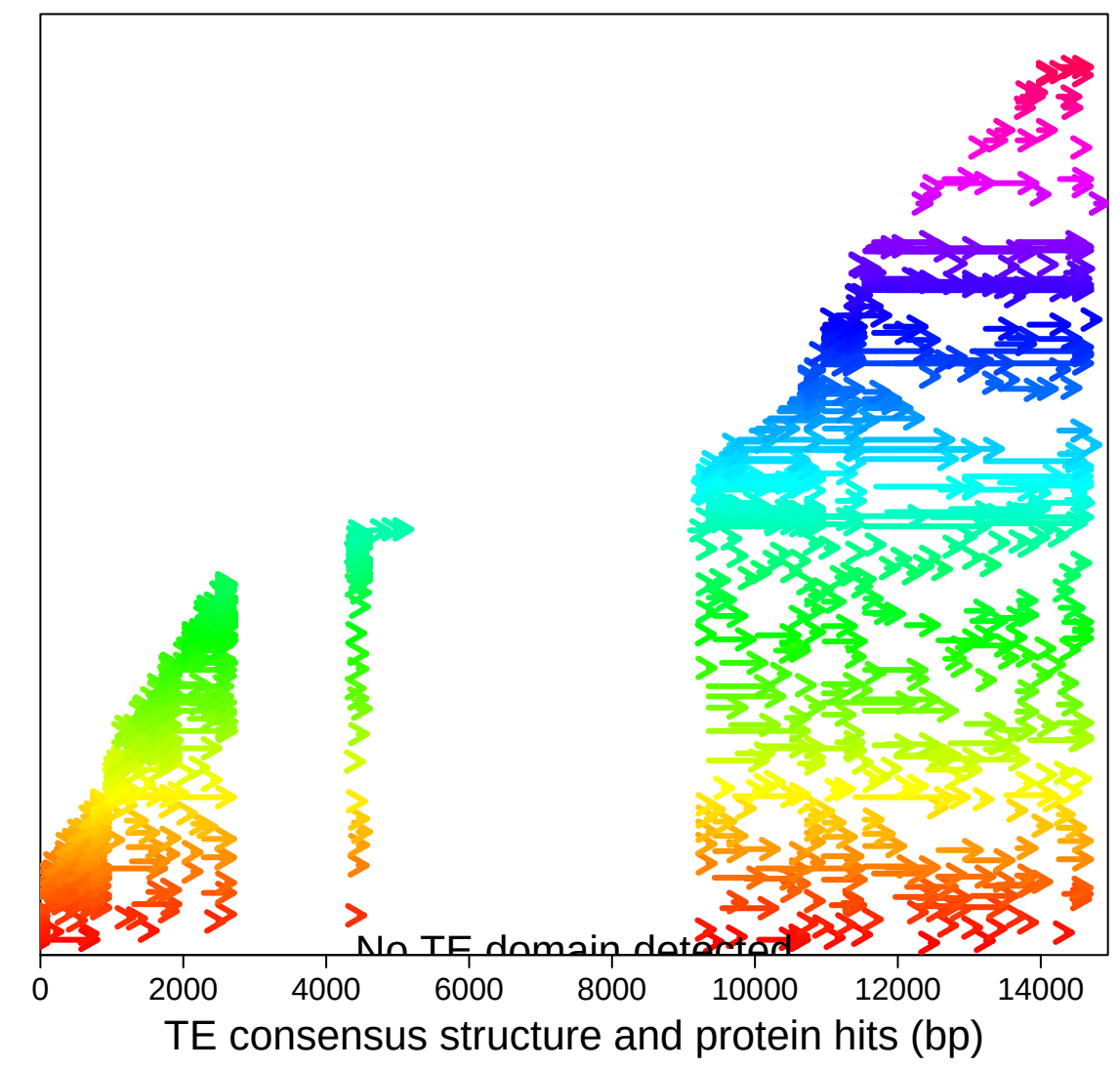
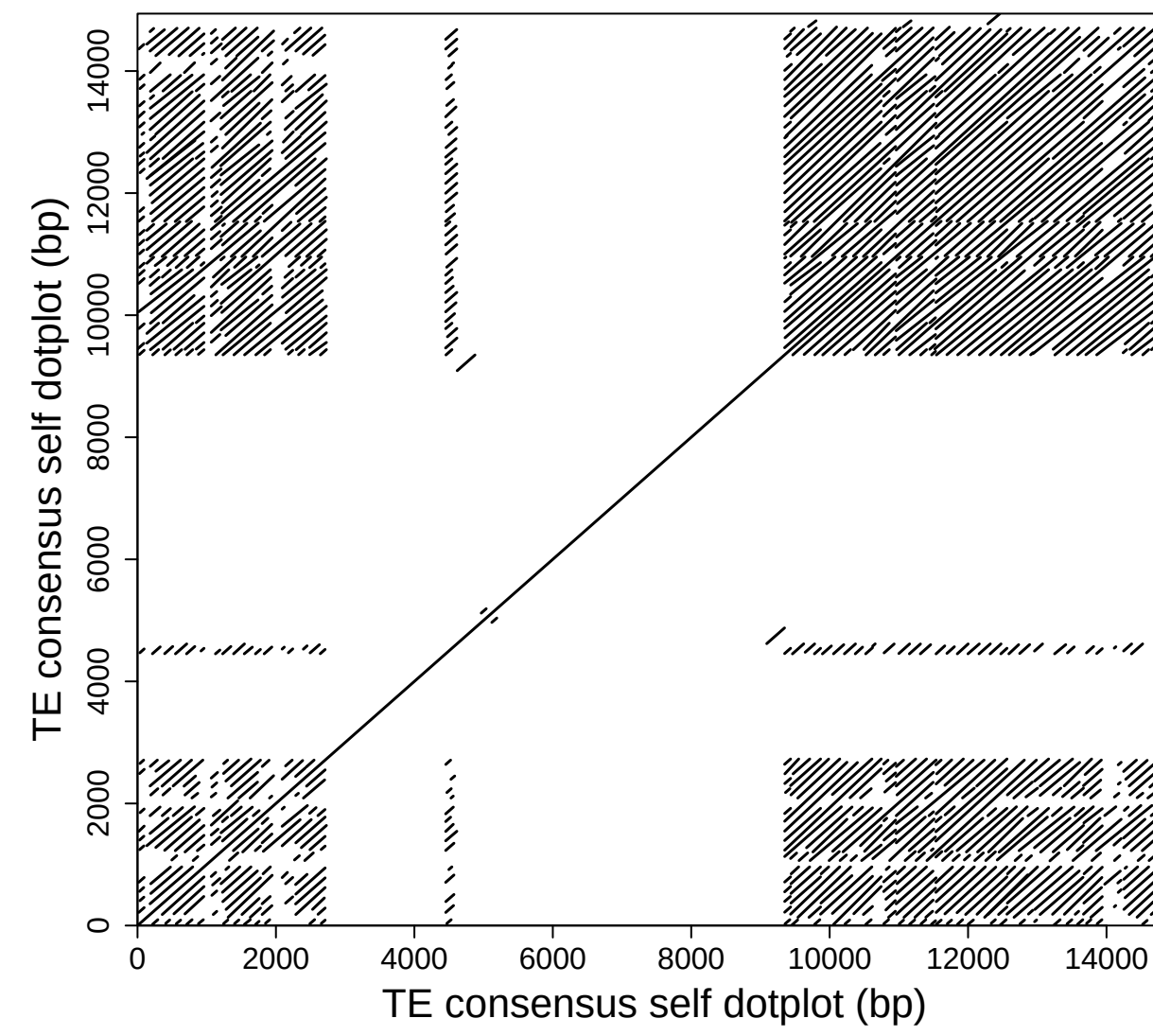
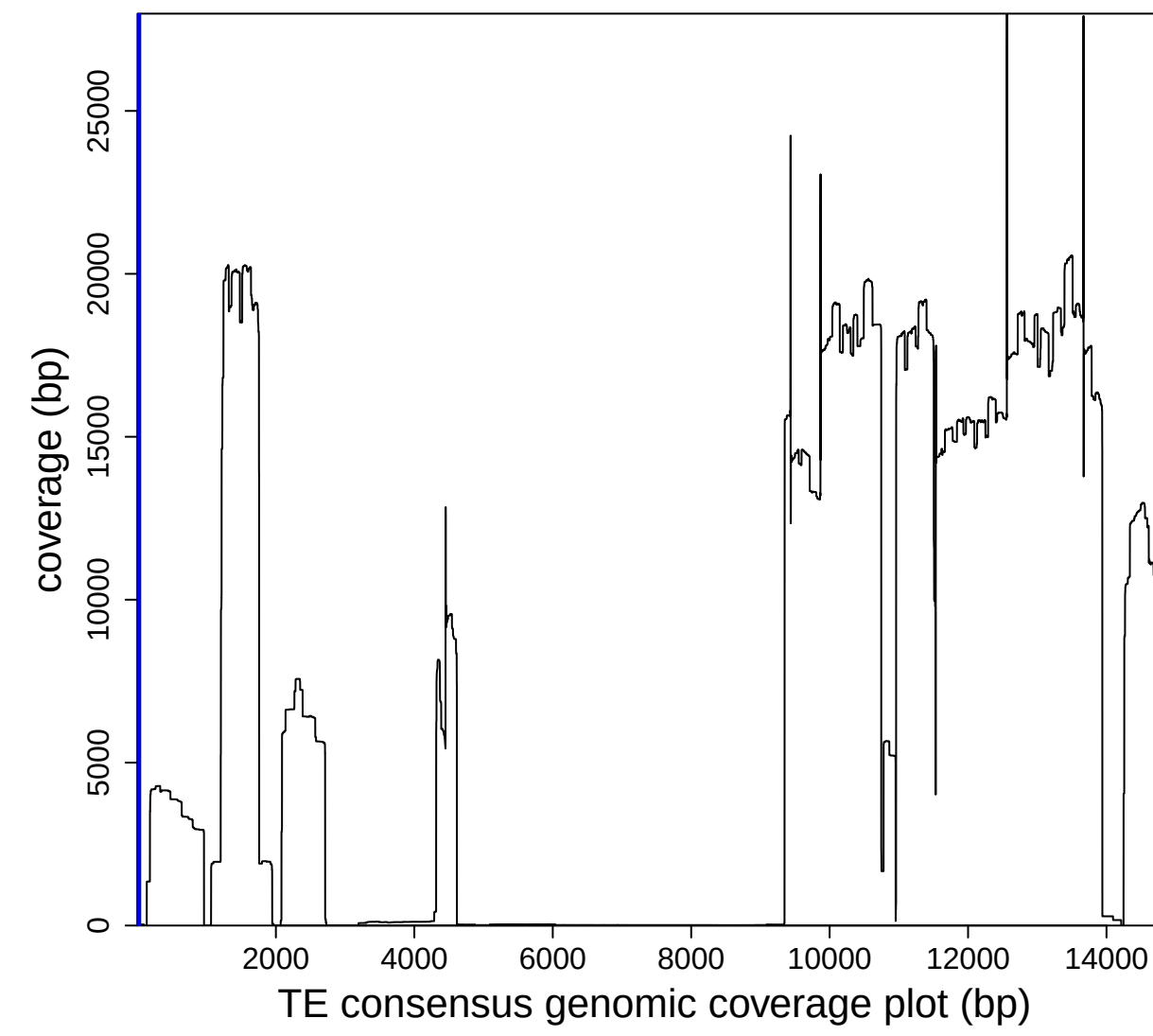
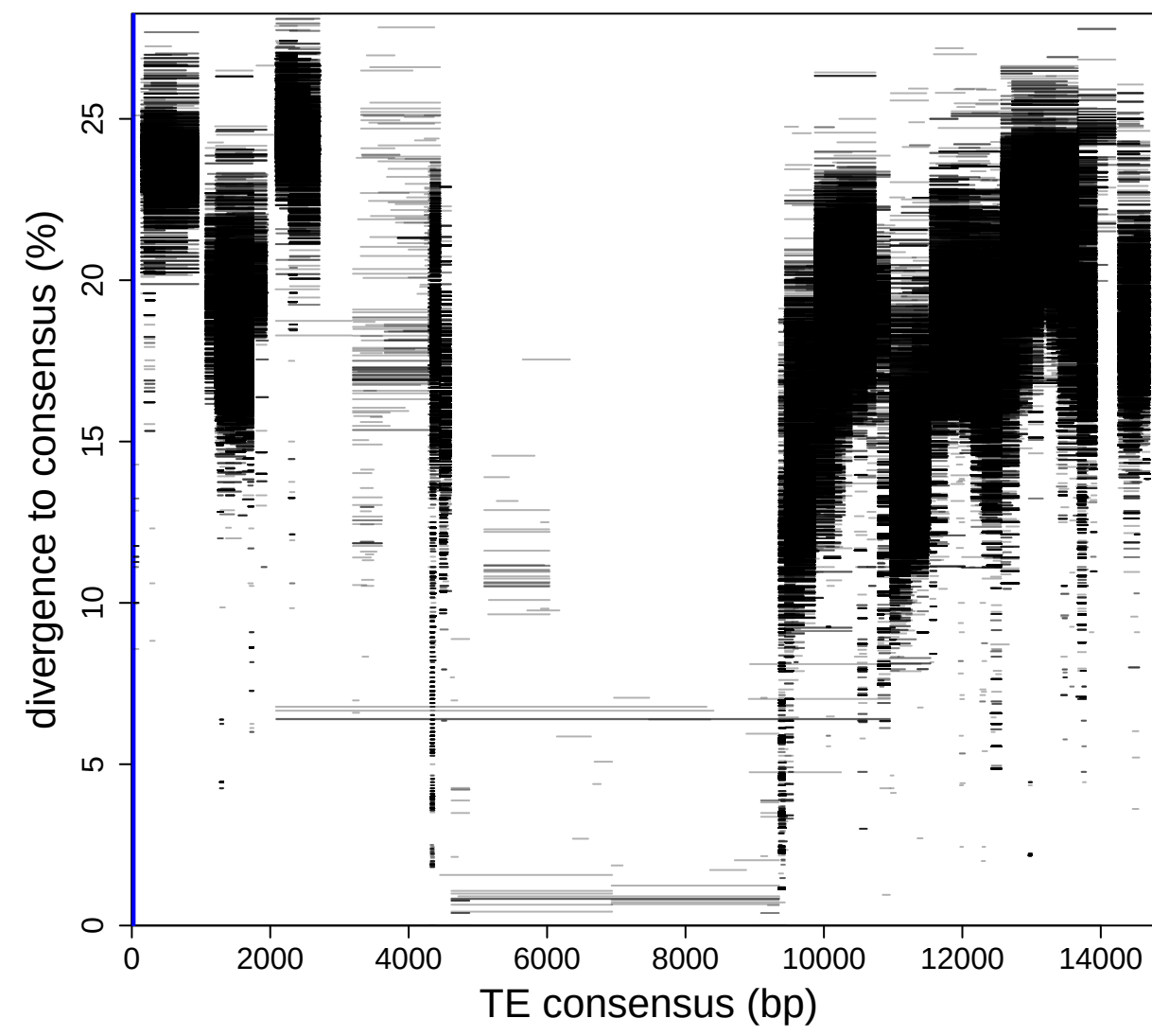
ed_uf.bed g_4.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa ce.
size: 14538bp; fragments: 364609; full length: 0 (≥ 13084.2 bp)

After TEtrimmer 14538 bp



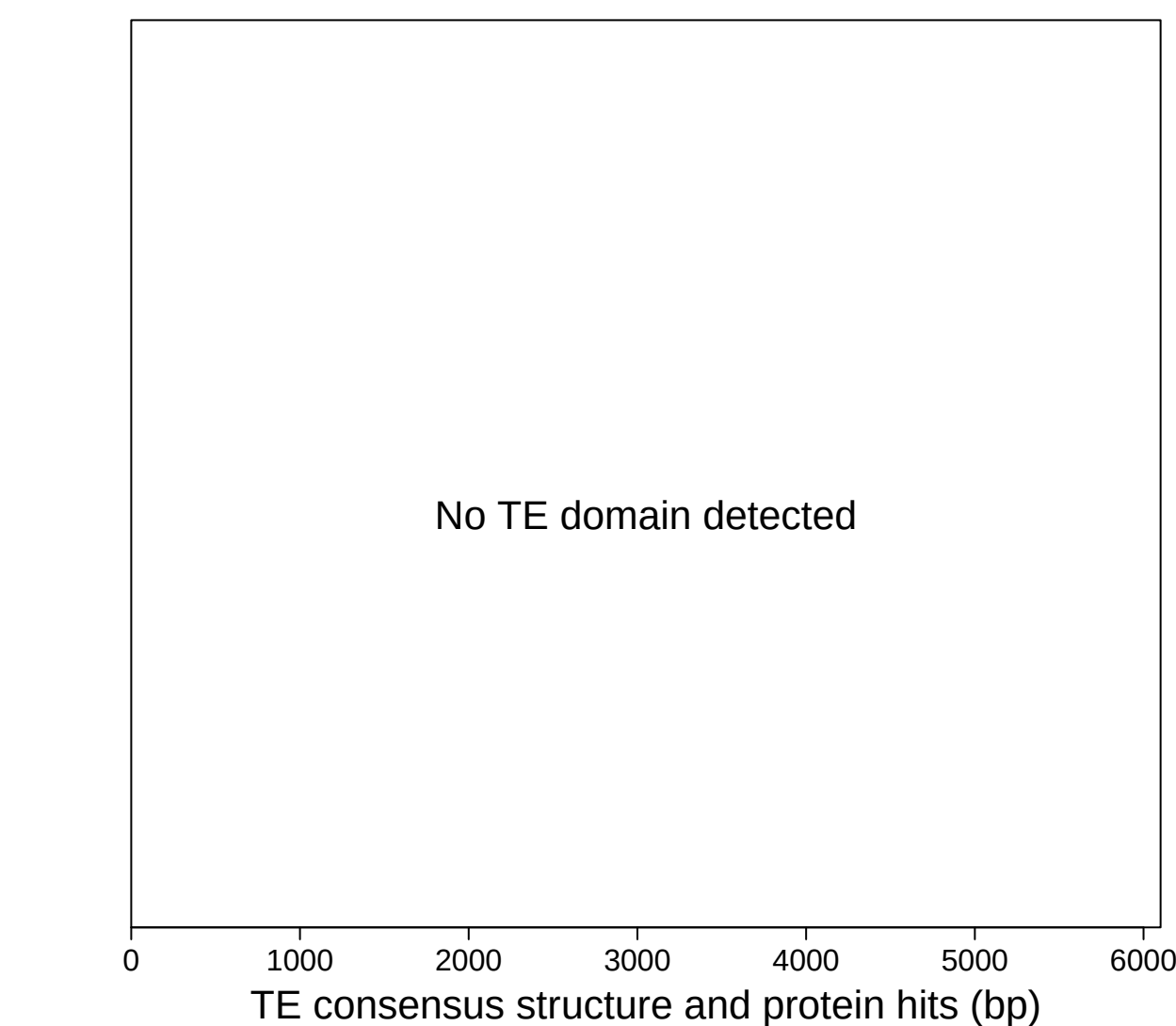
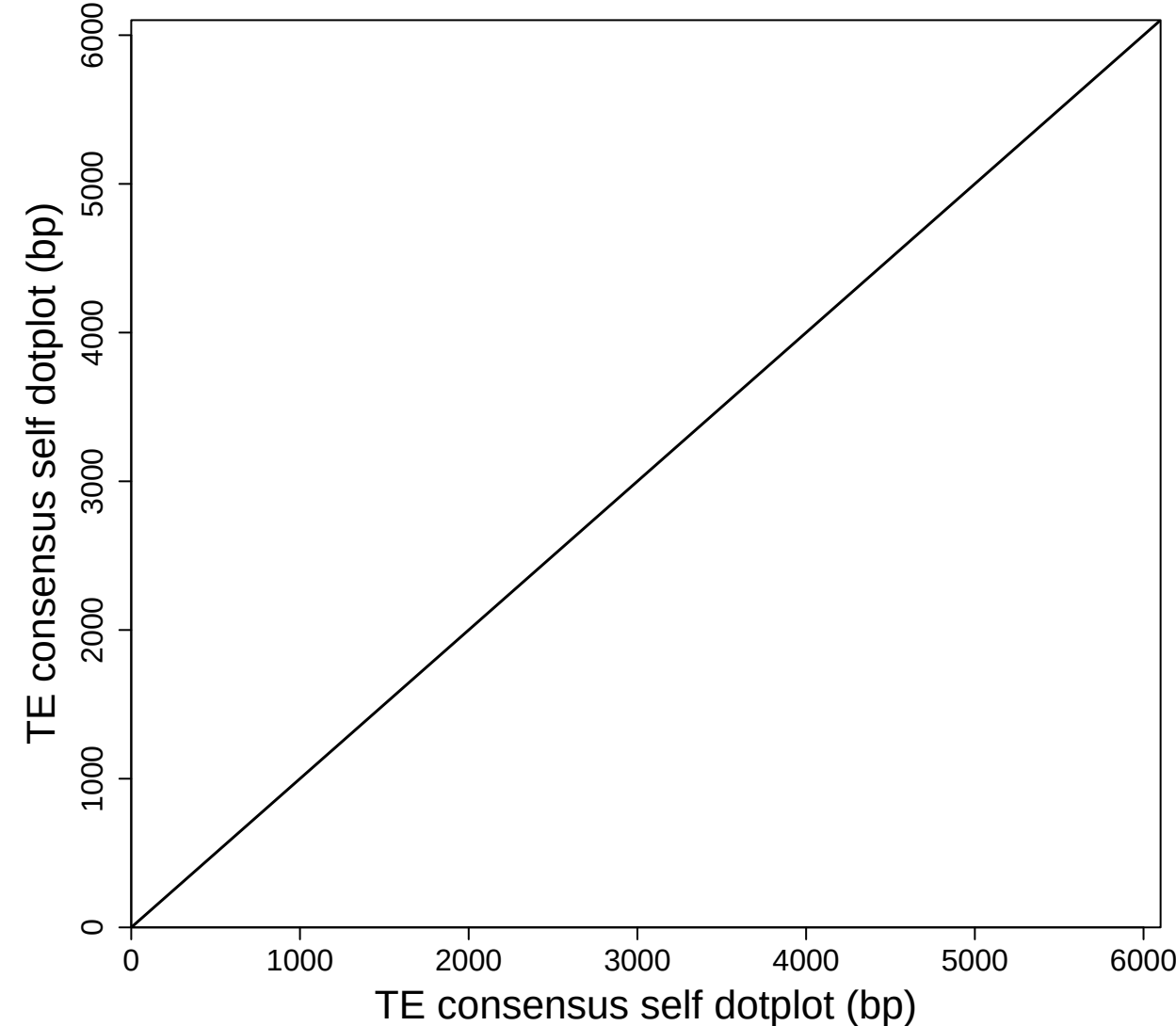
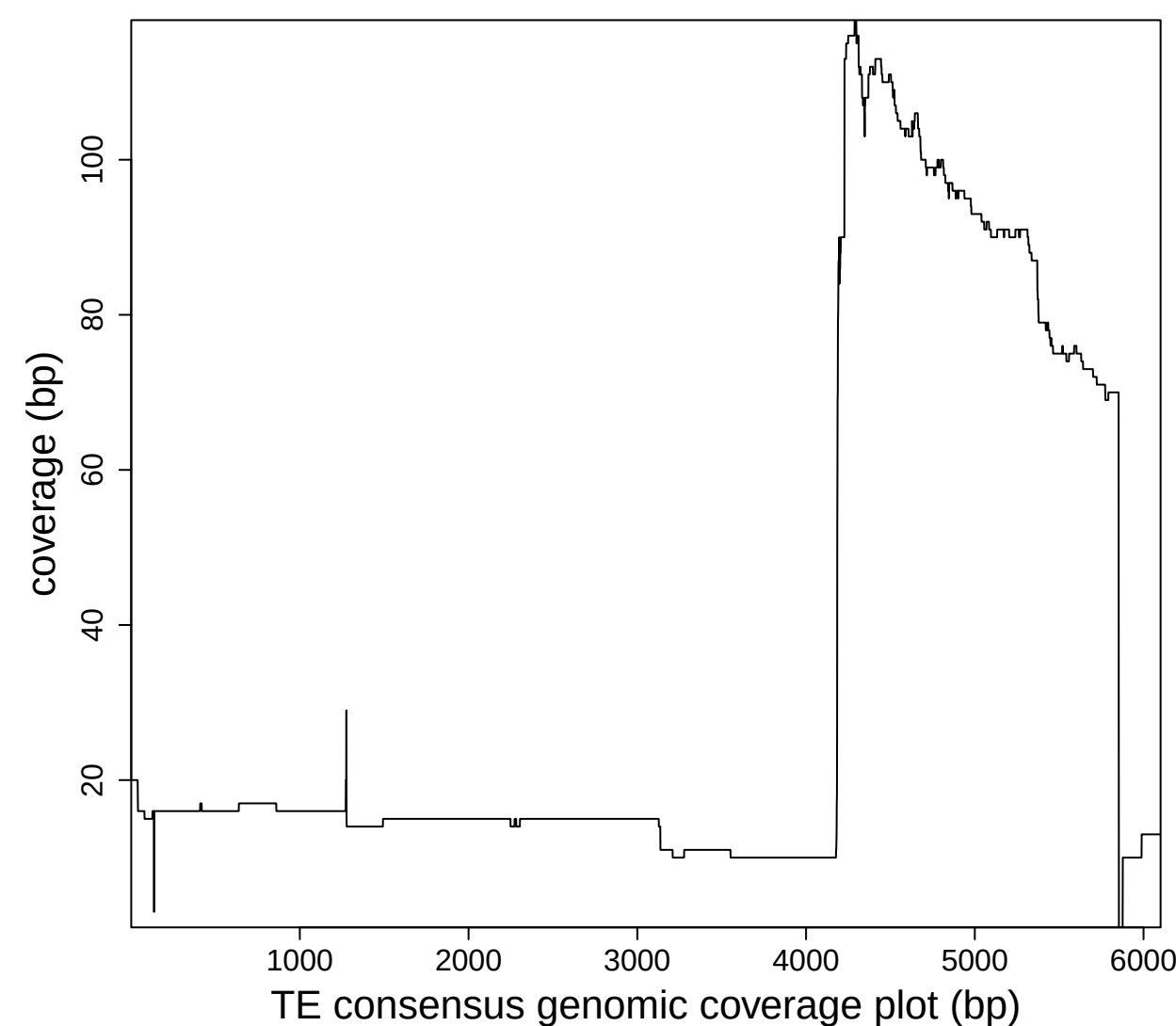
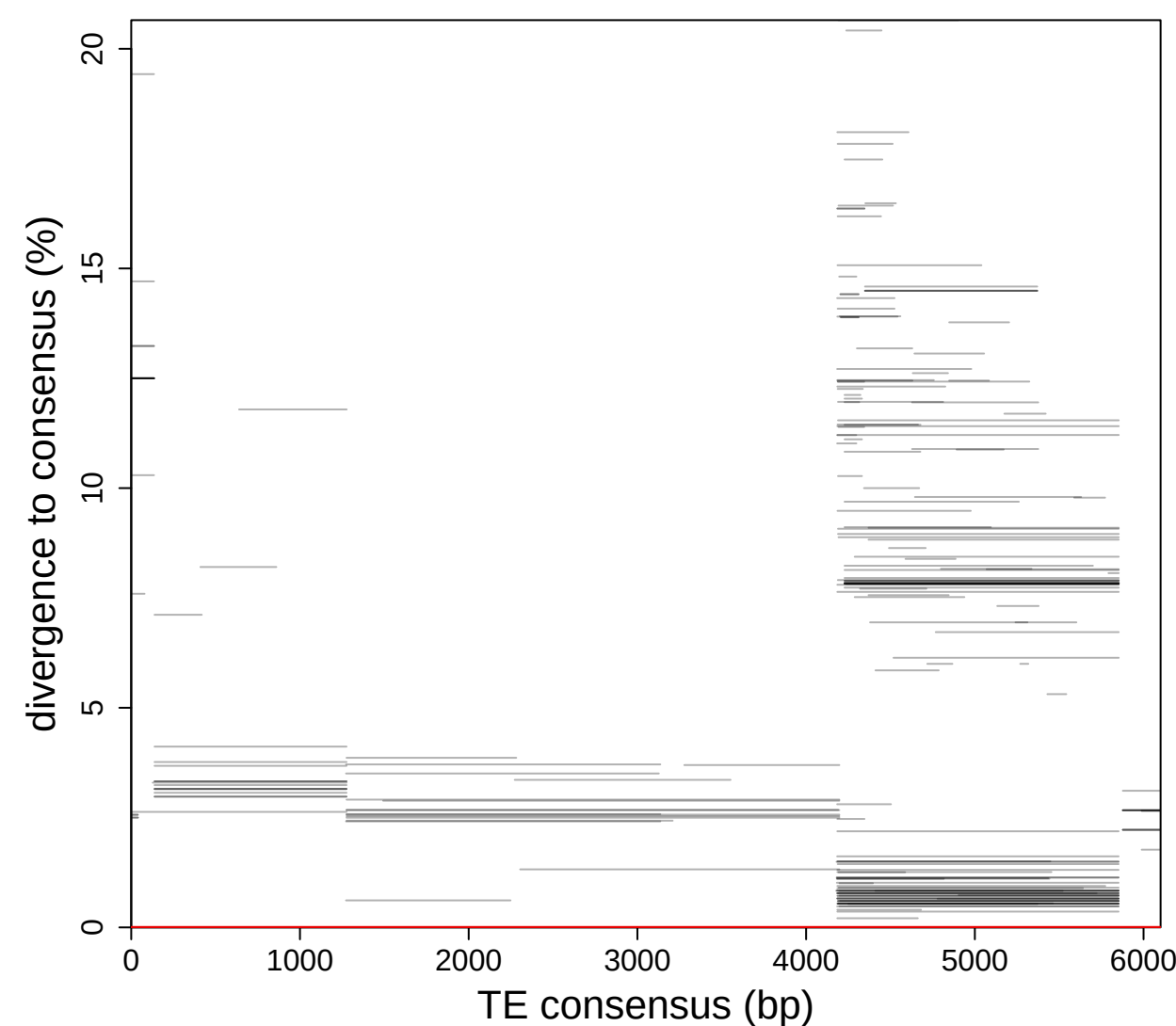
ly_606.fasta.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcIn.fa_aln.fa
size: 14940bp; fragments: 203713; full length: 0 (>=13446bp)

After TEtrimmer Extended plot Blue lines are boundaries

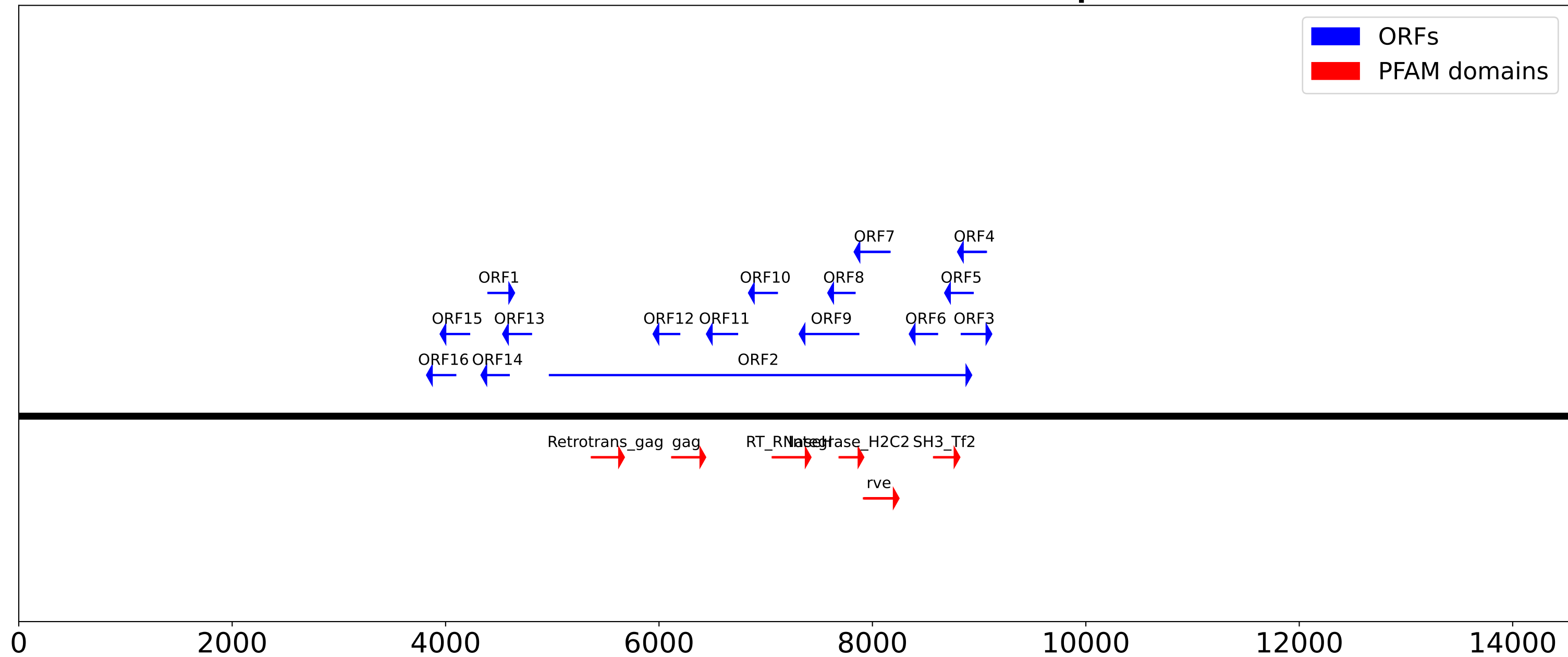


TE: ltr_1_family_606
size: 6101bp; fragments: 223; full length: 1 (>=5490.9bp)

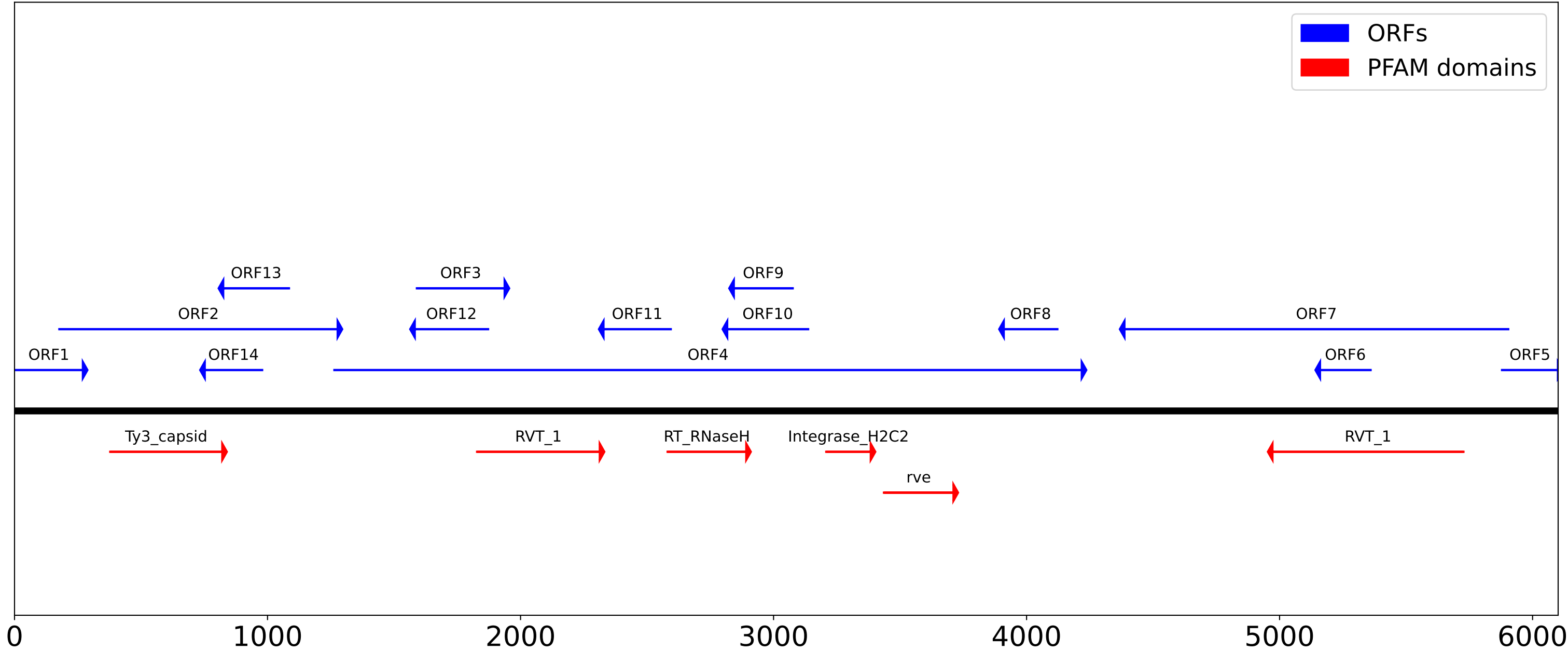
Before TEtrimmer 6101 bp



After TEtrimmer ORF and PFAM domain plot



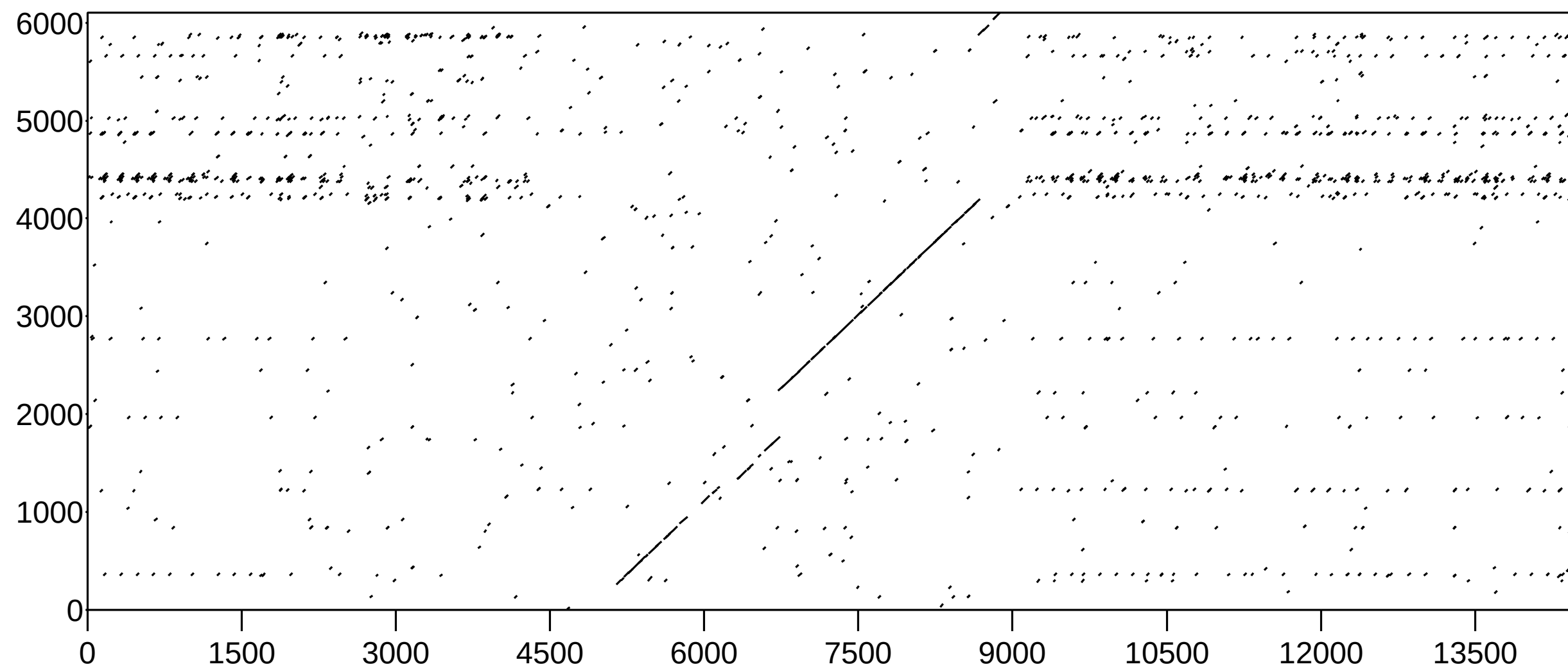
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)