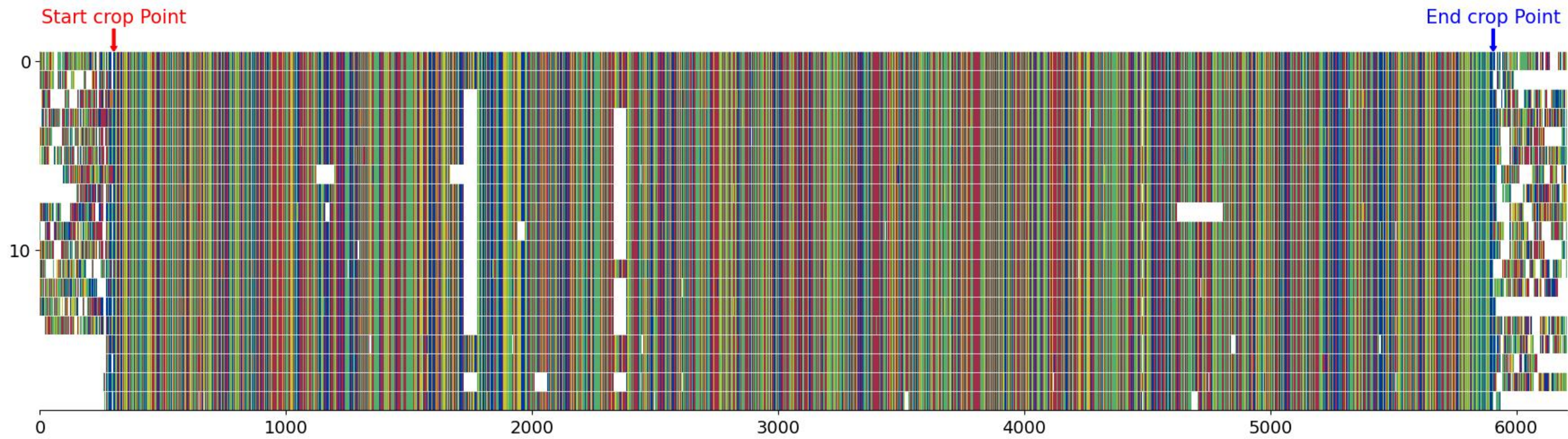




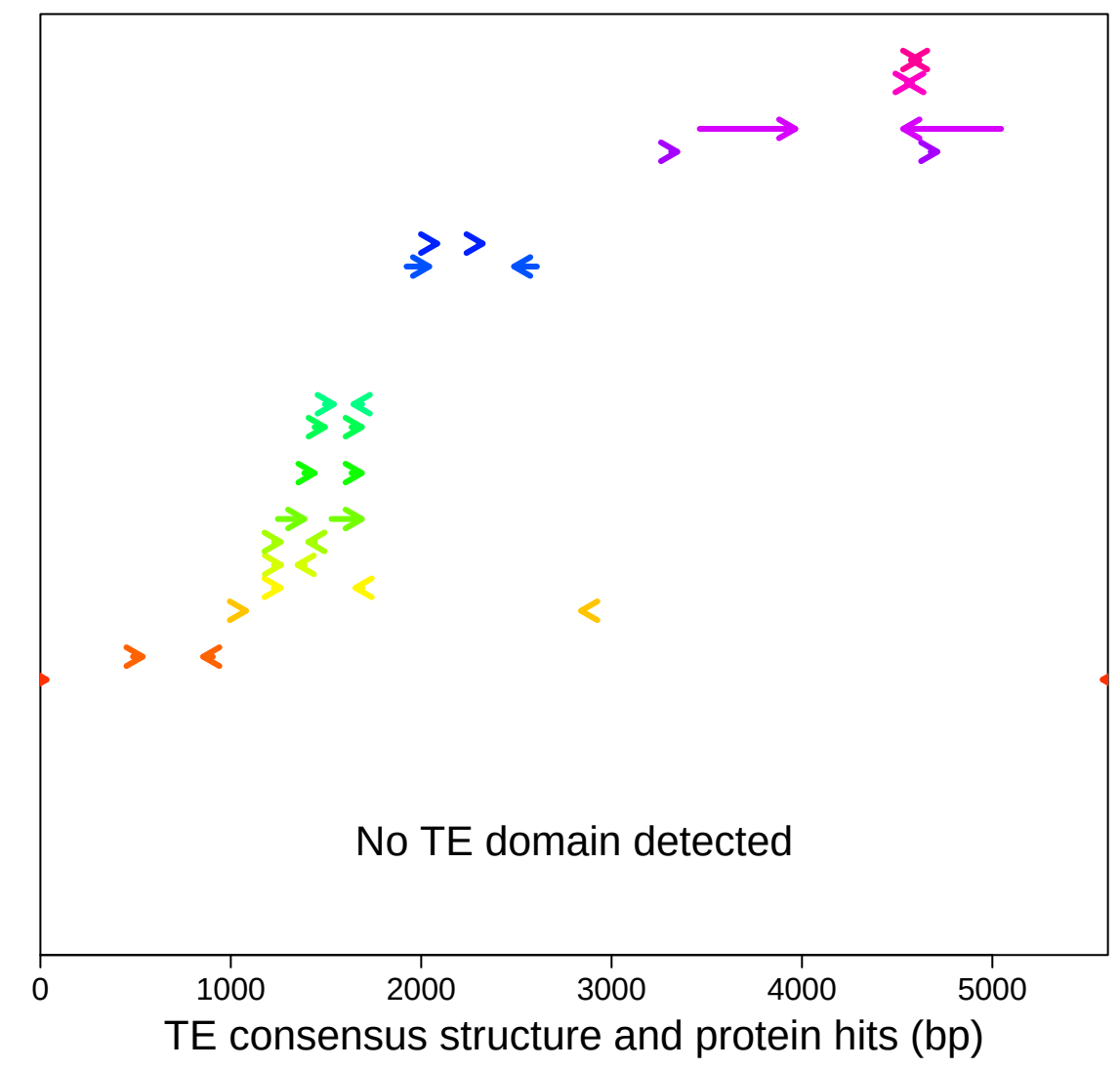
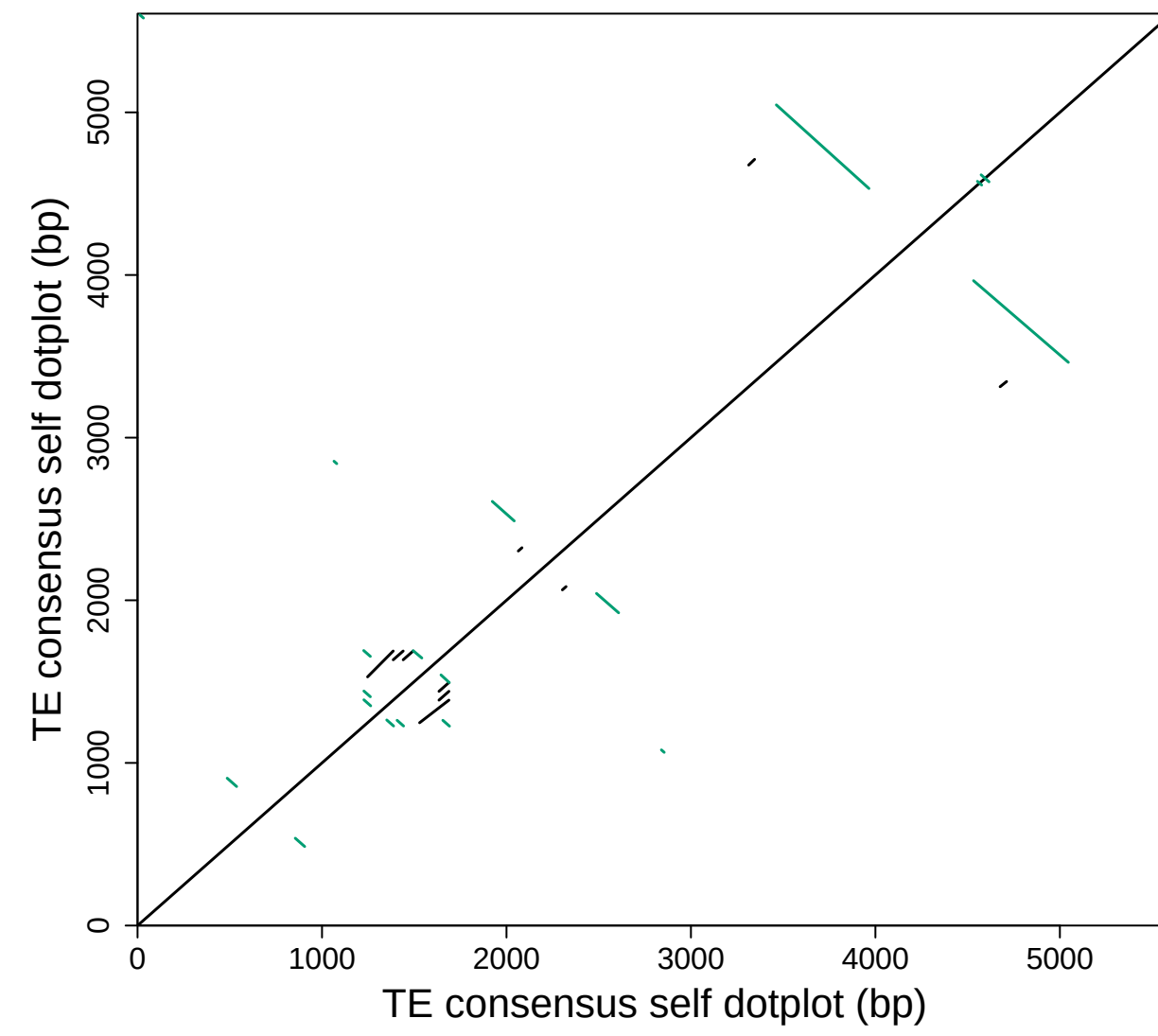
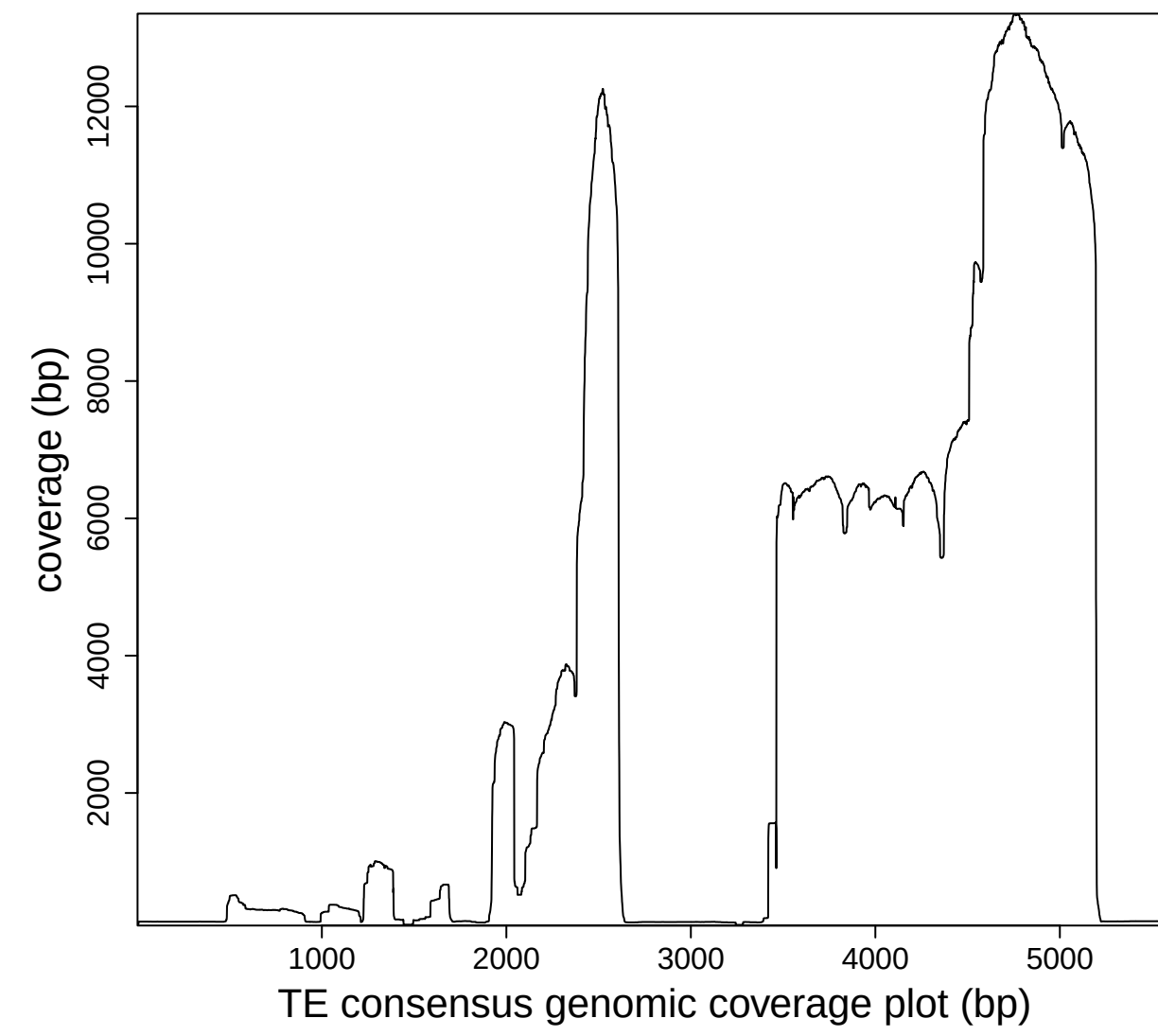
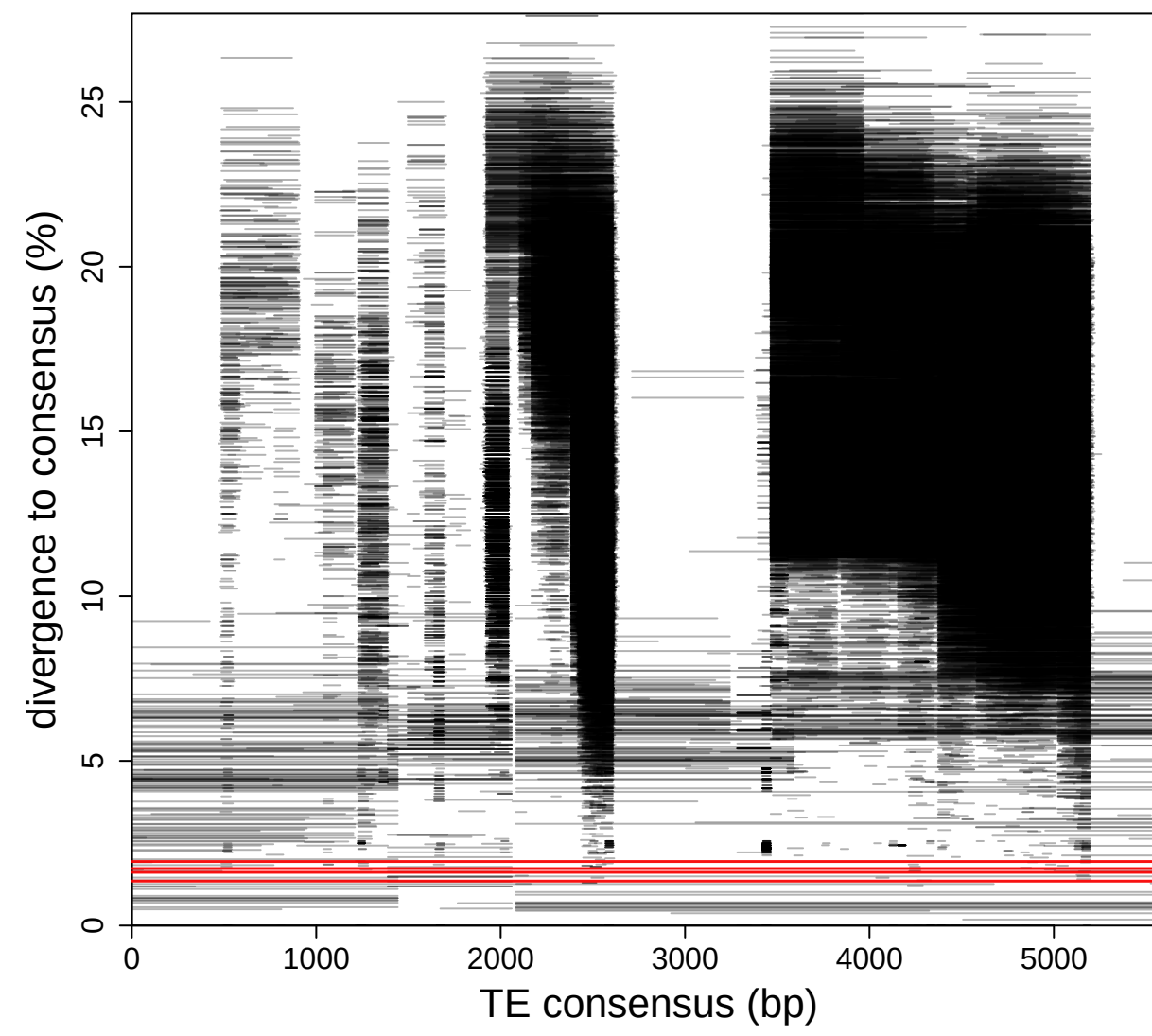
Start crop Point

End crop Point



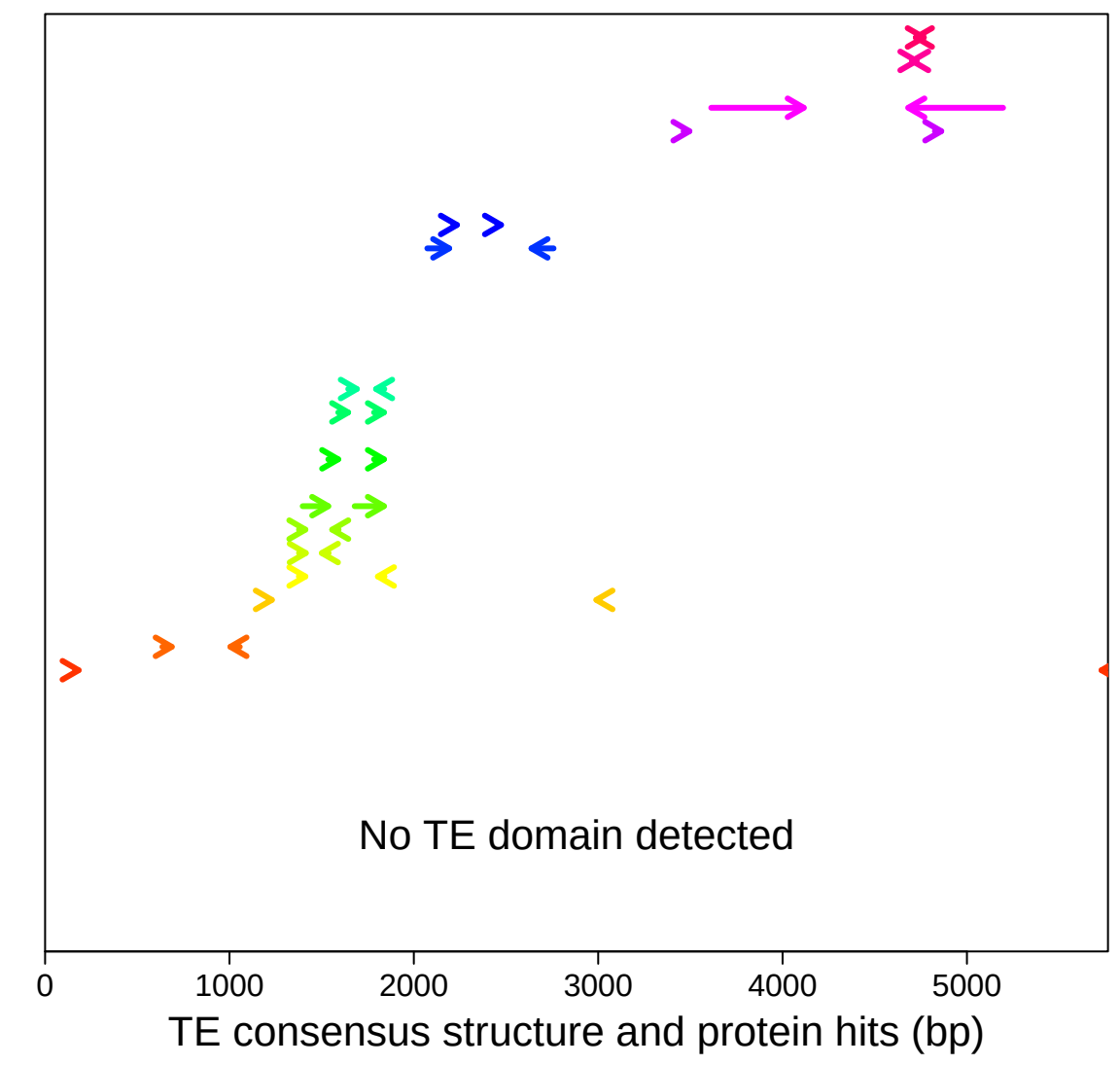
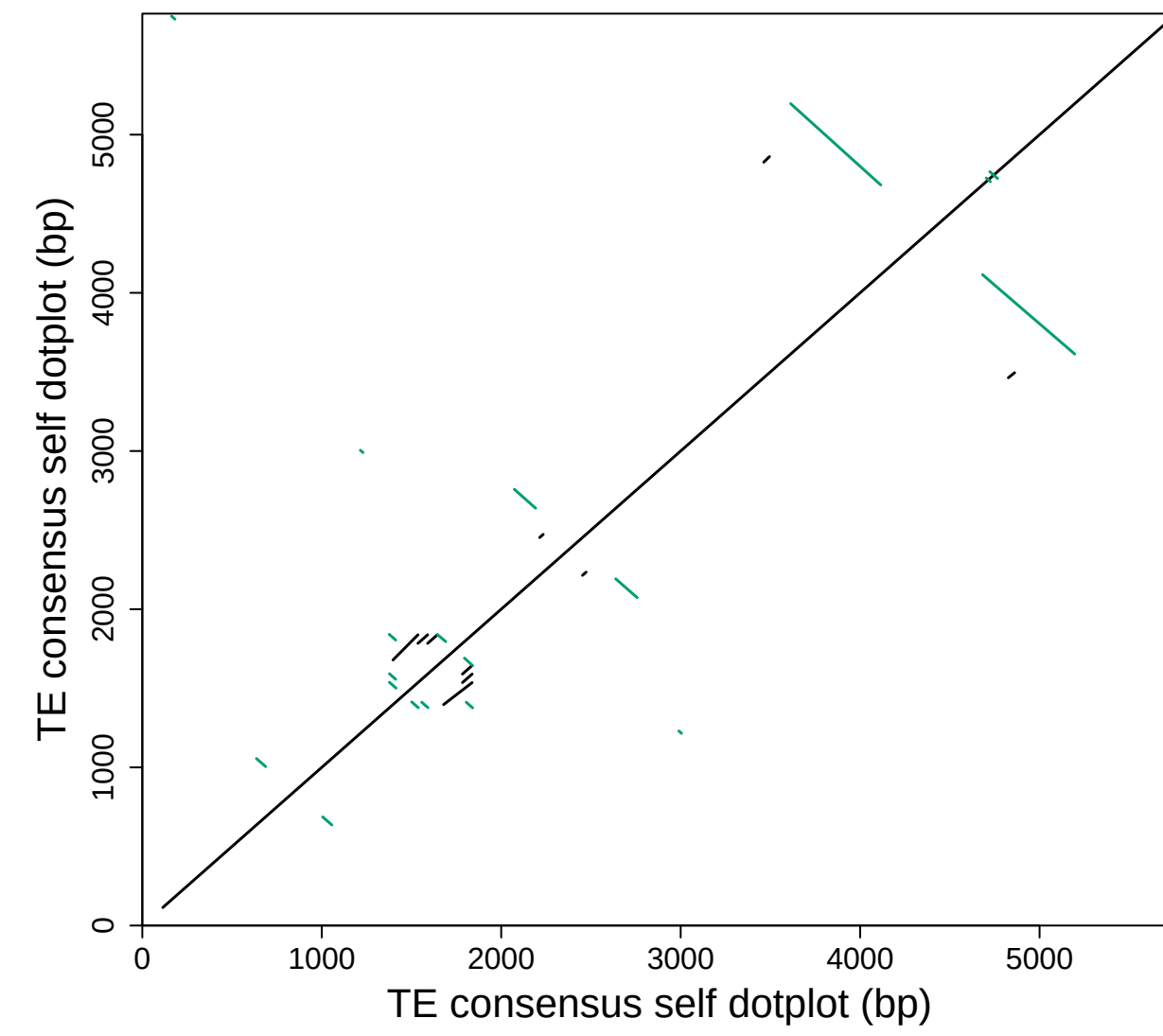
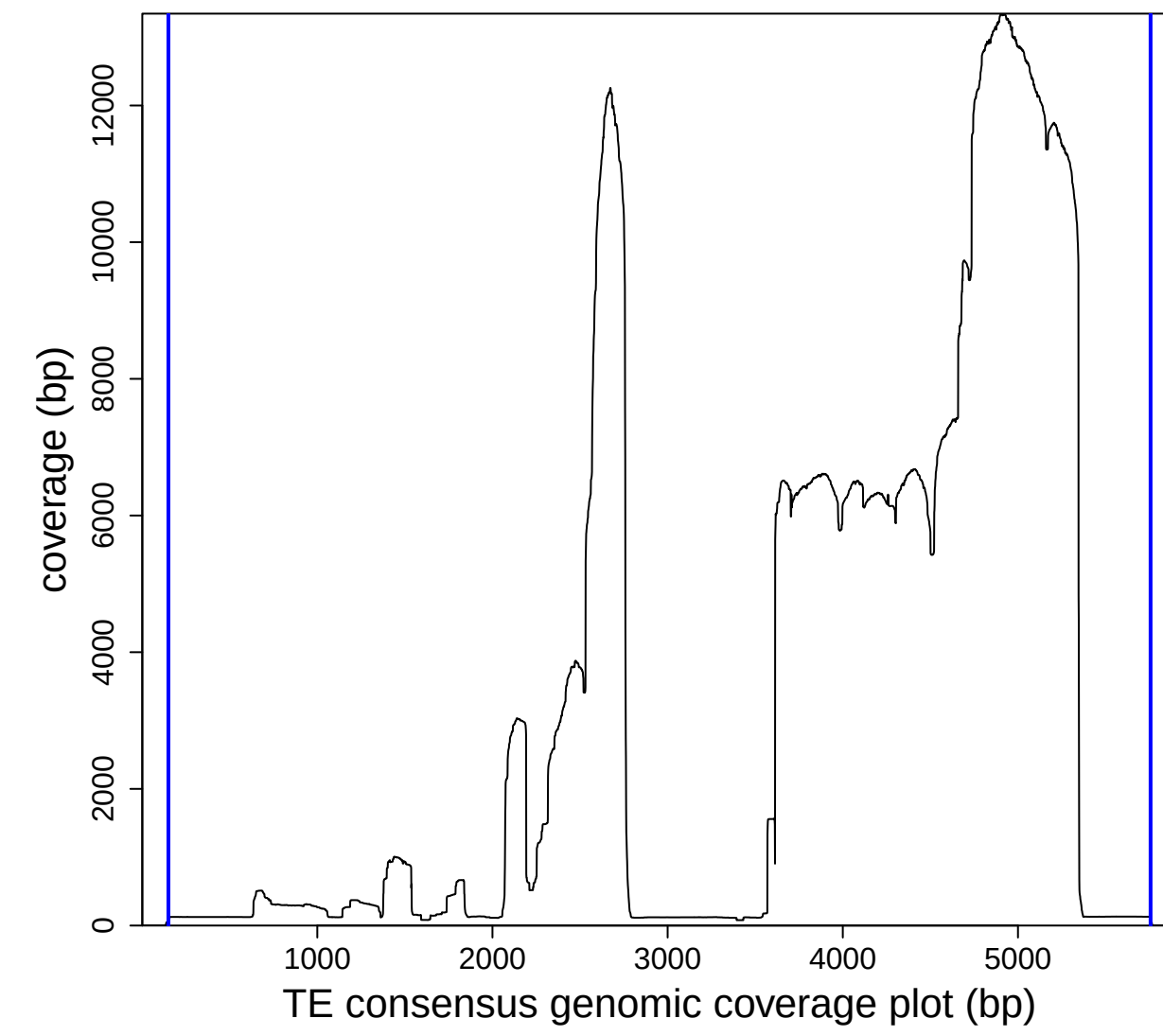
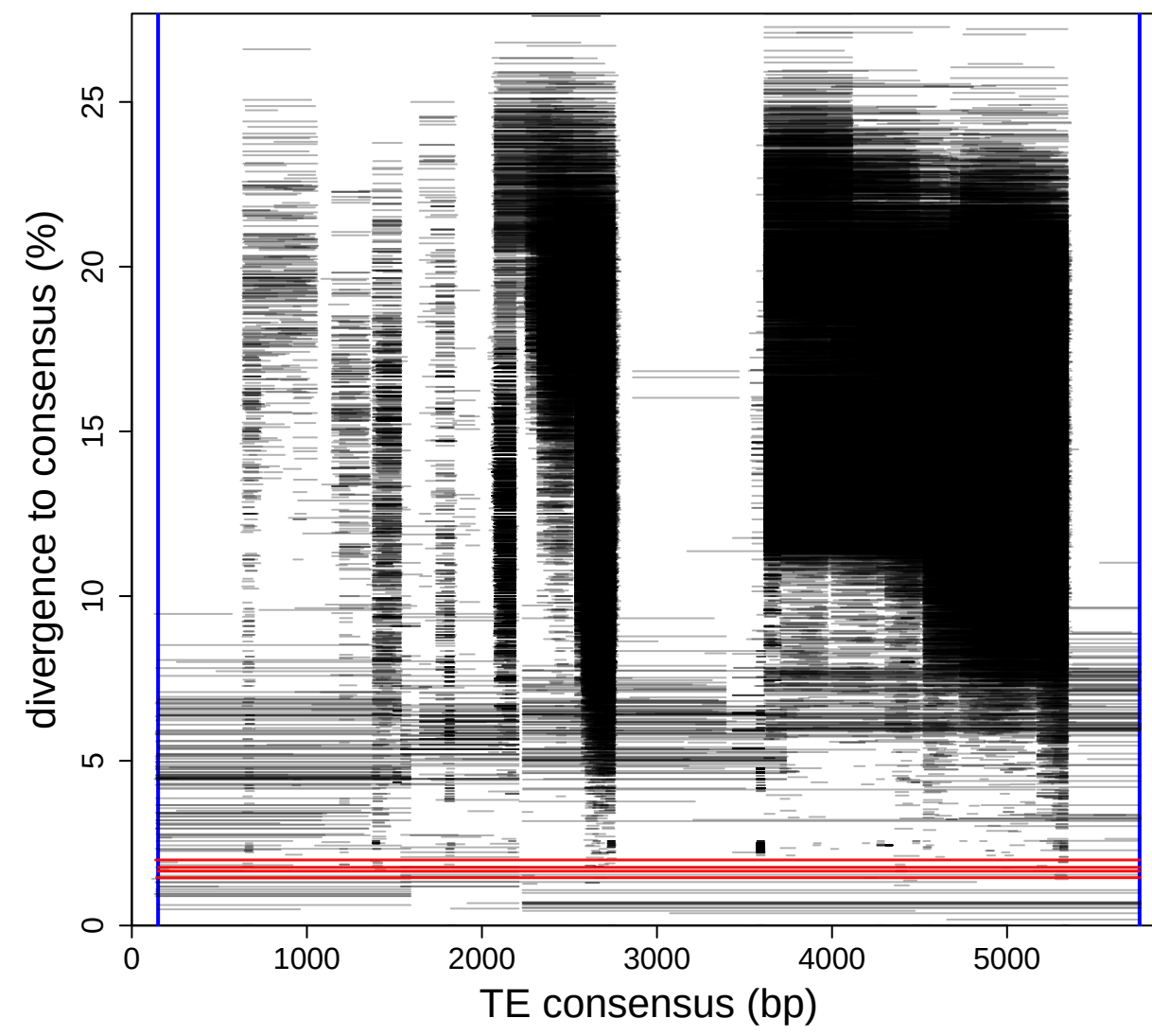
l.b.bed uf.bed g_3.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 5607bp; fragments: 43368; full length: 4 (>=5046.3bp)

After TEtrimmer 5607 bp



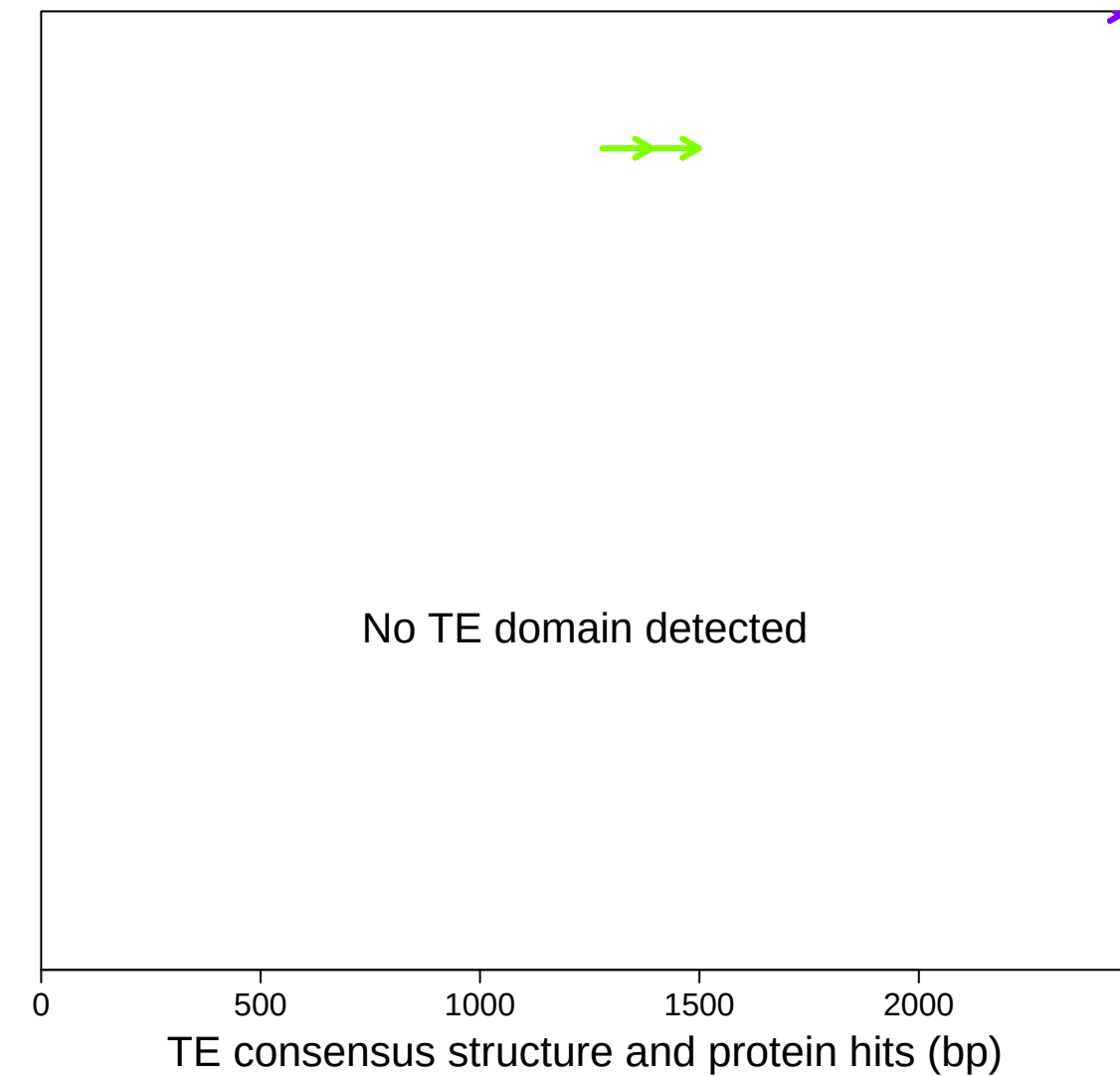
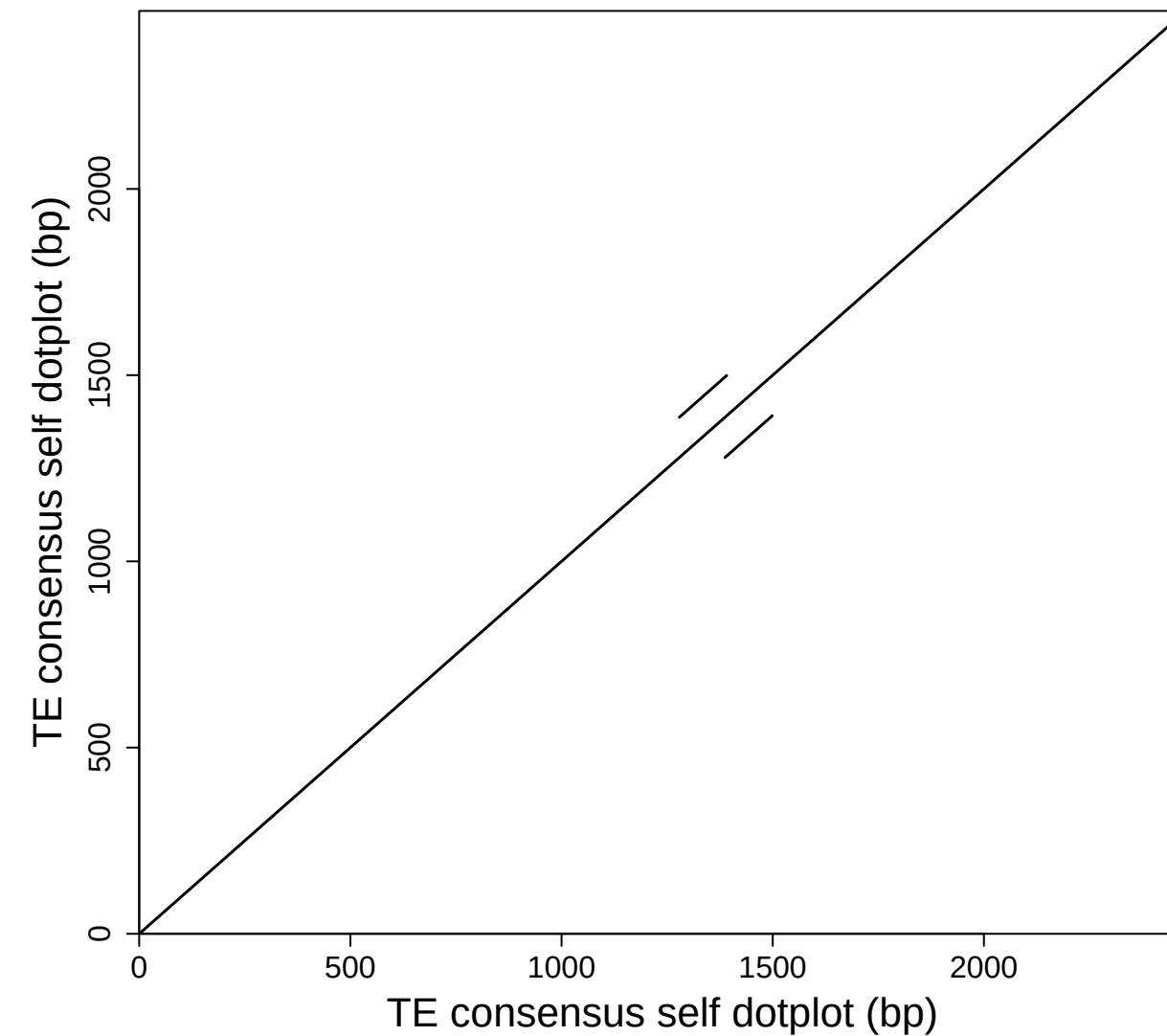
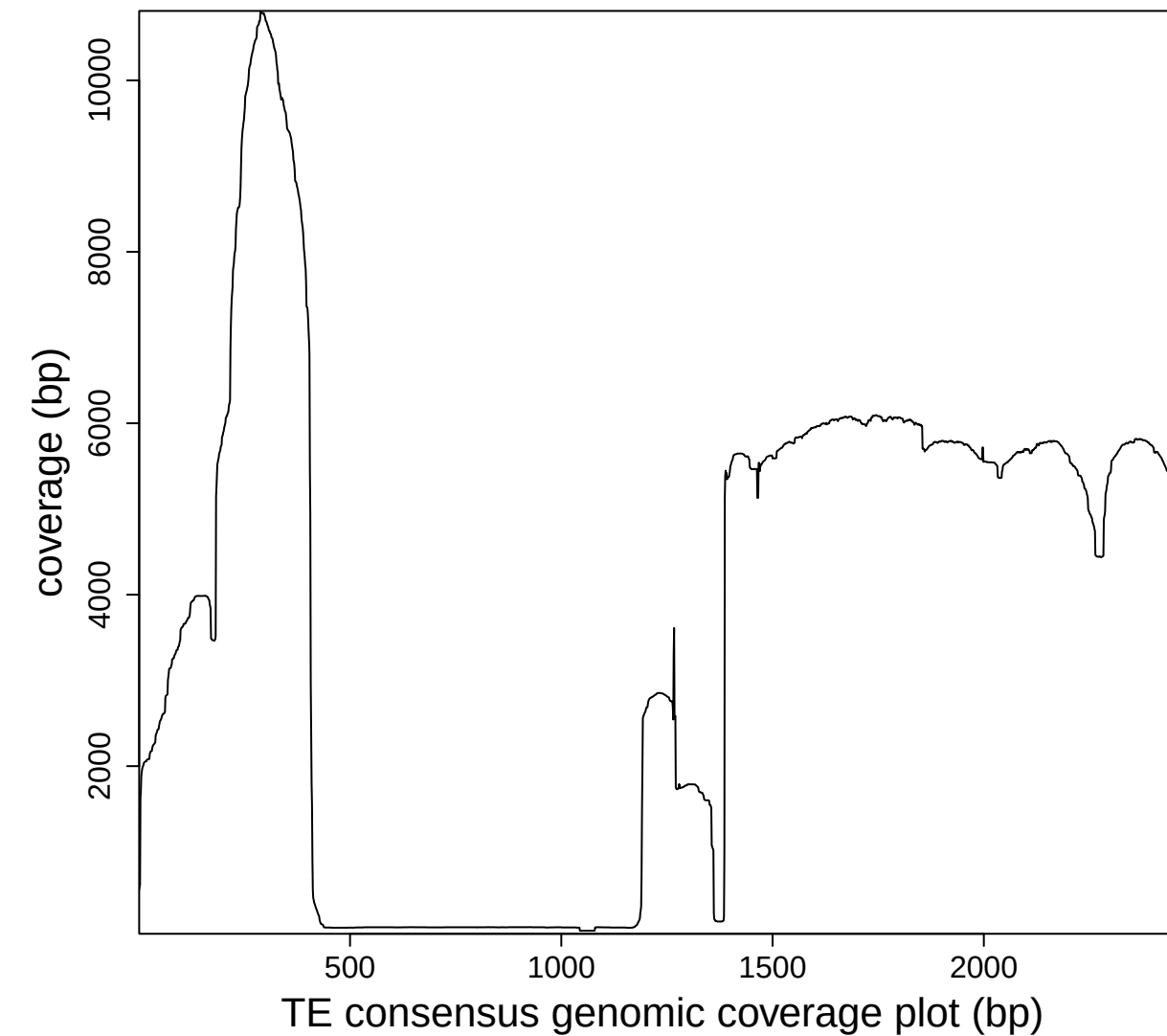
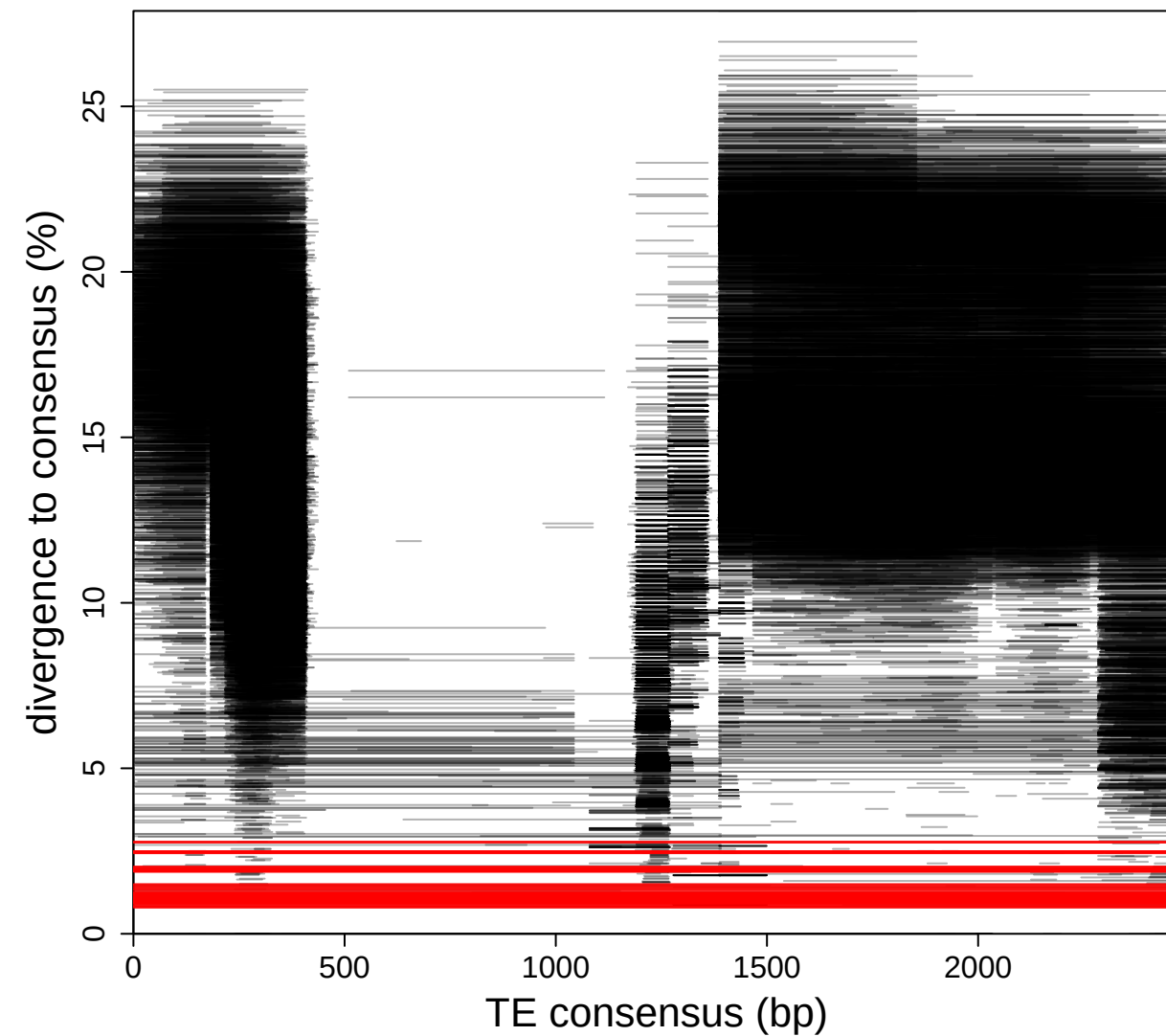
9.fasta.b.bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 5907bp; fragments: 43327; full length: 4 (>=5316.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

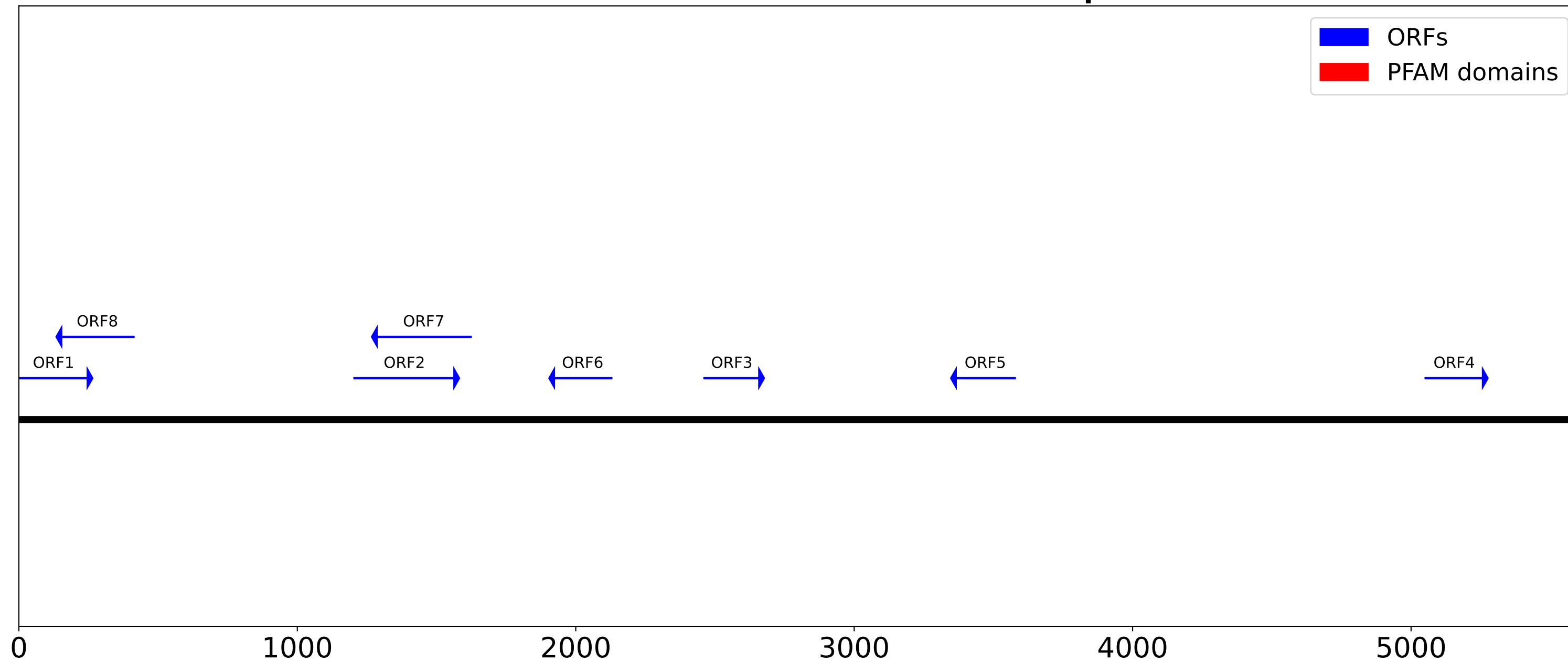


TE: rnd_1_family_989
size: 2478bp; fragments: 27422; full length: 31 (≥ 2230.2 bp)

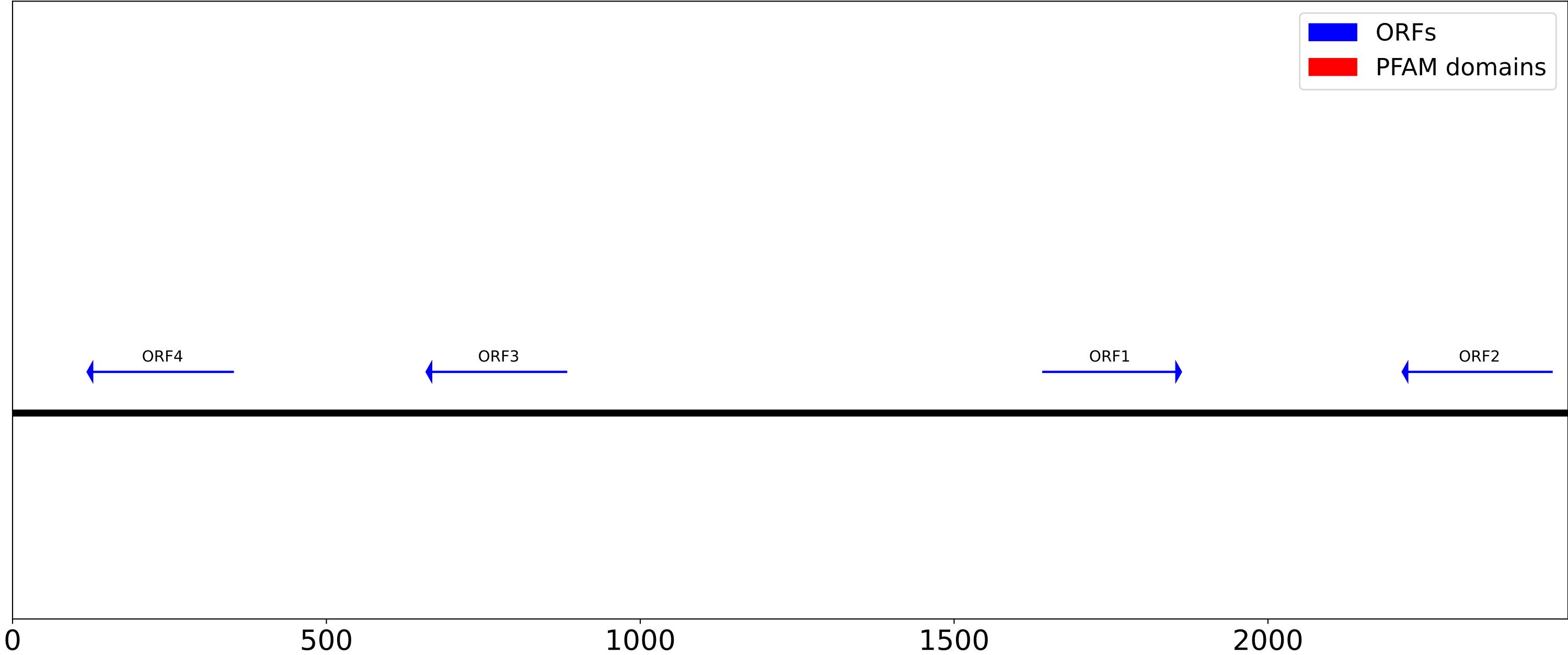
Before TEtrimmer 2478 bp



After TEtrimmer ORF and PFAM domain plot



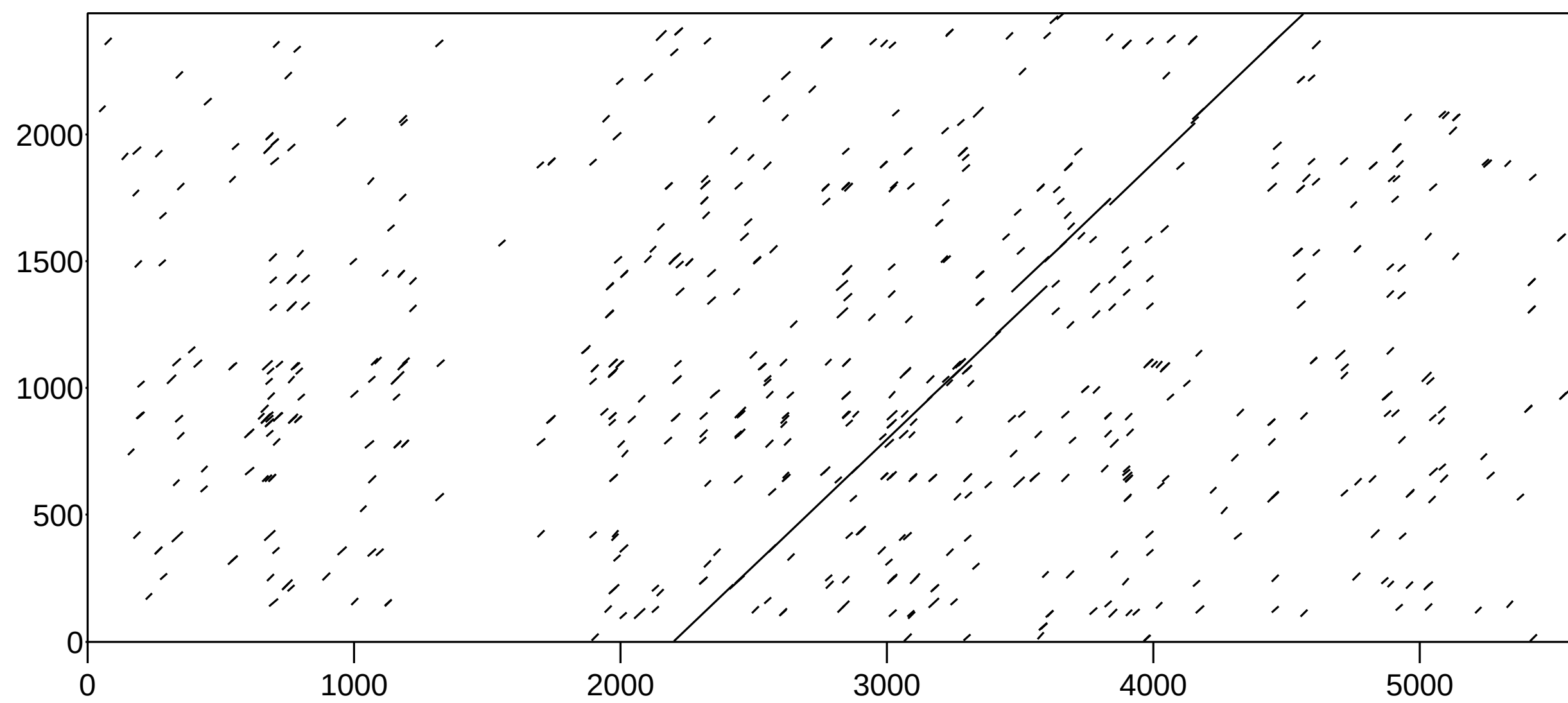
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)