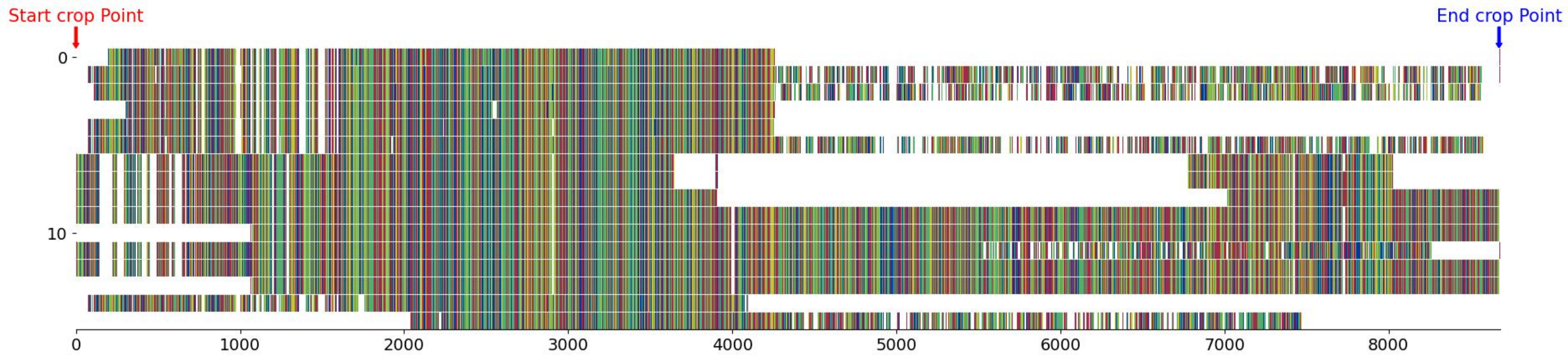


MSA length = 8675

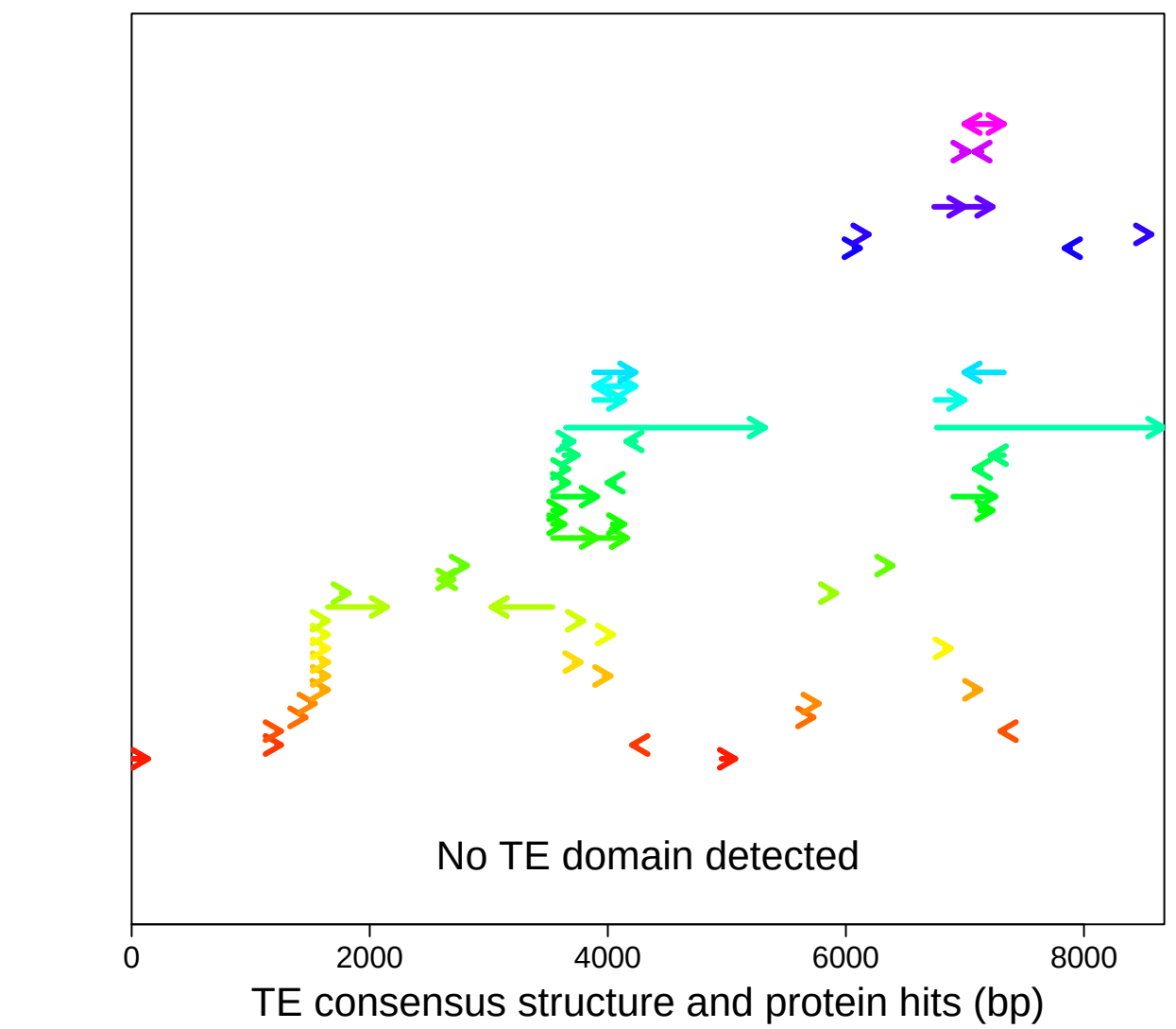
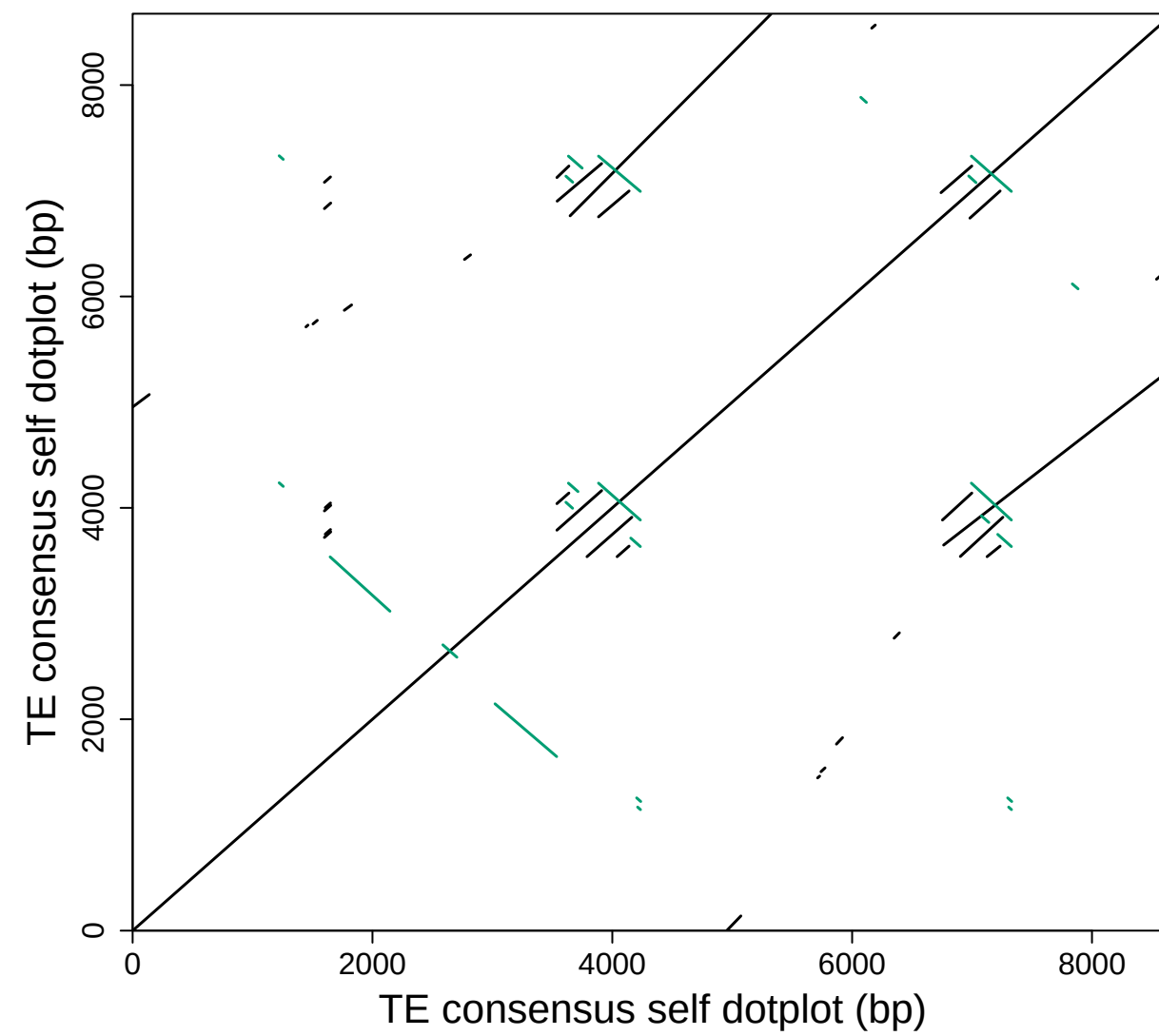
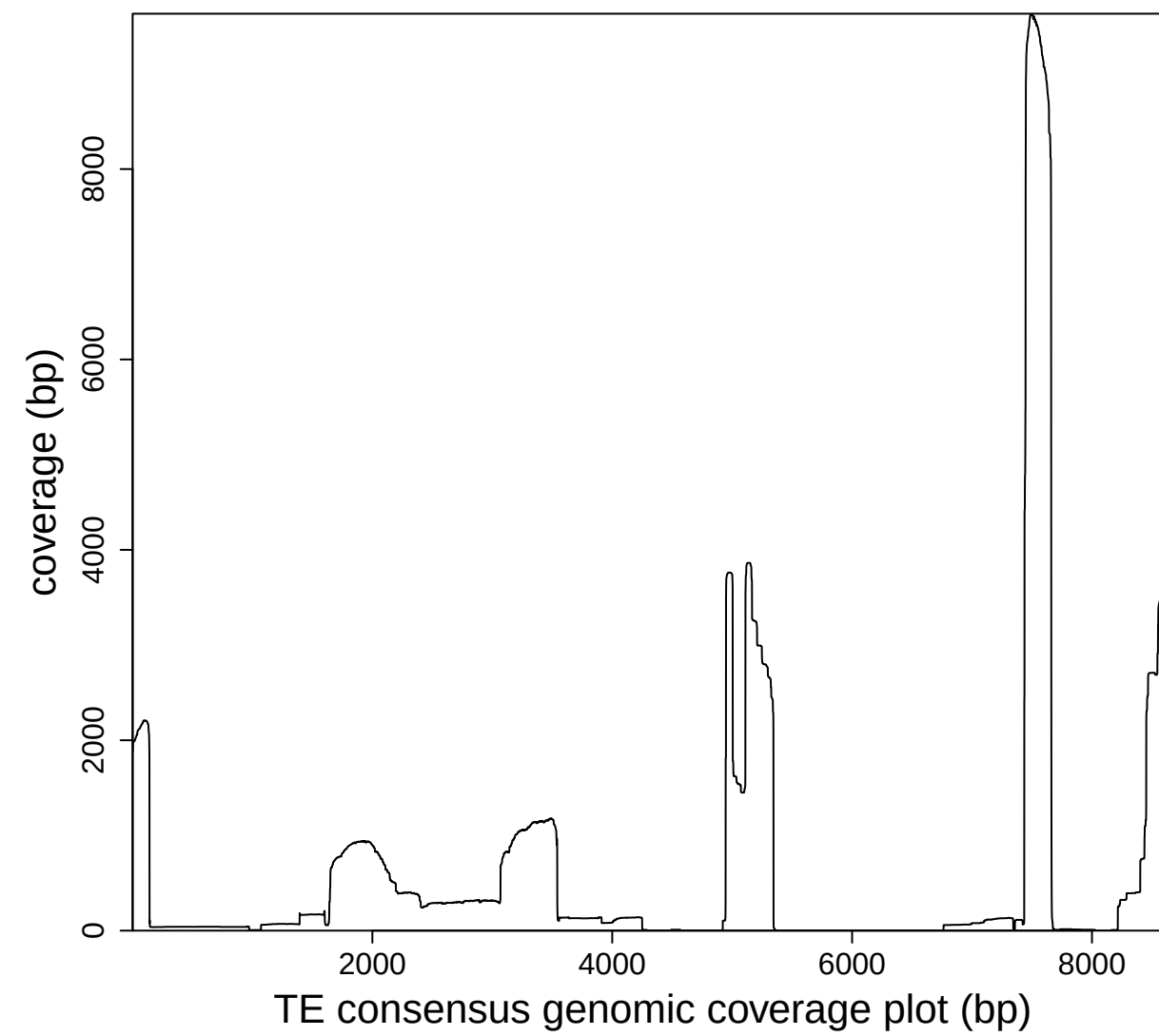
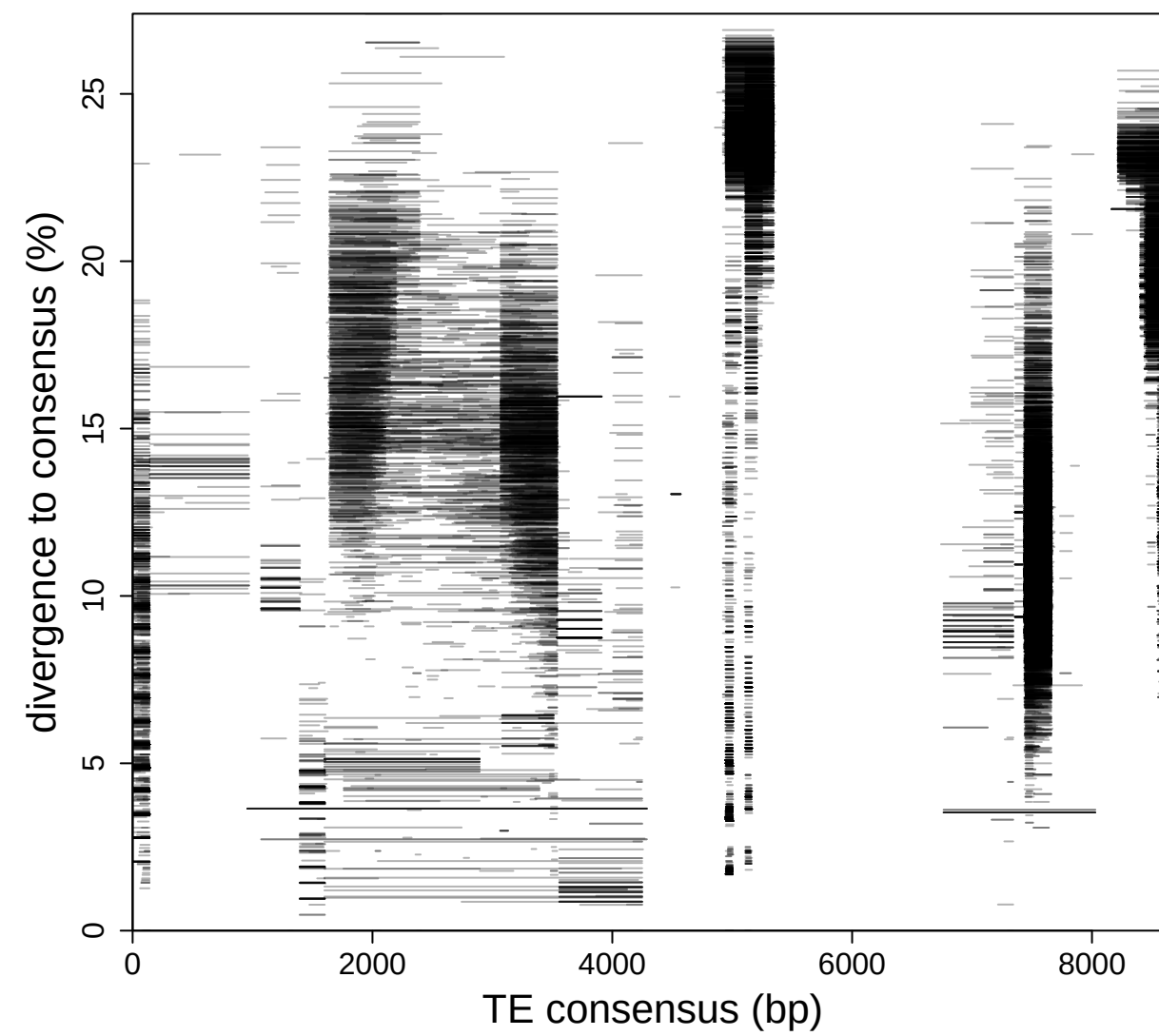
End crop Point

[illegible]



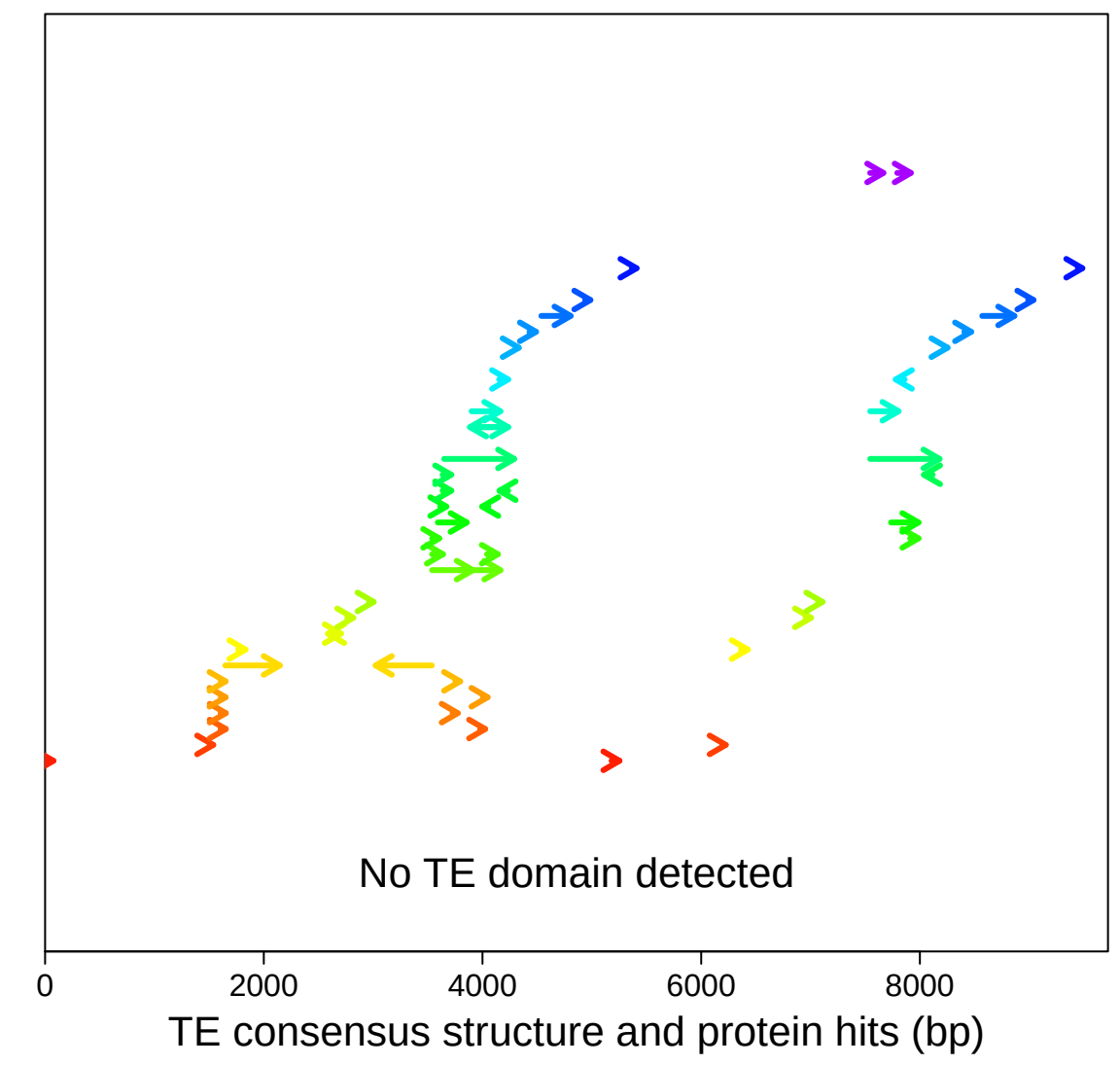
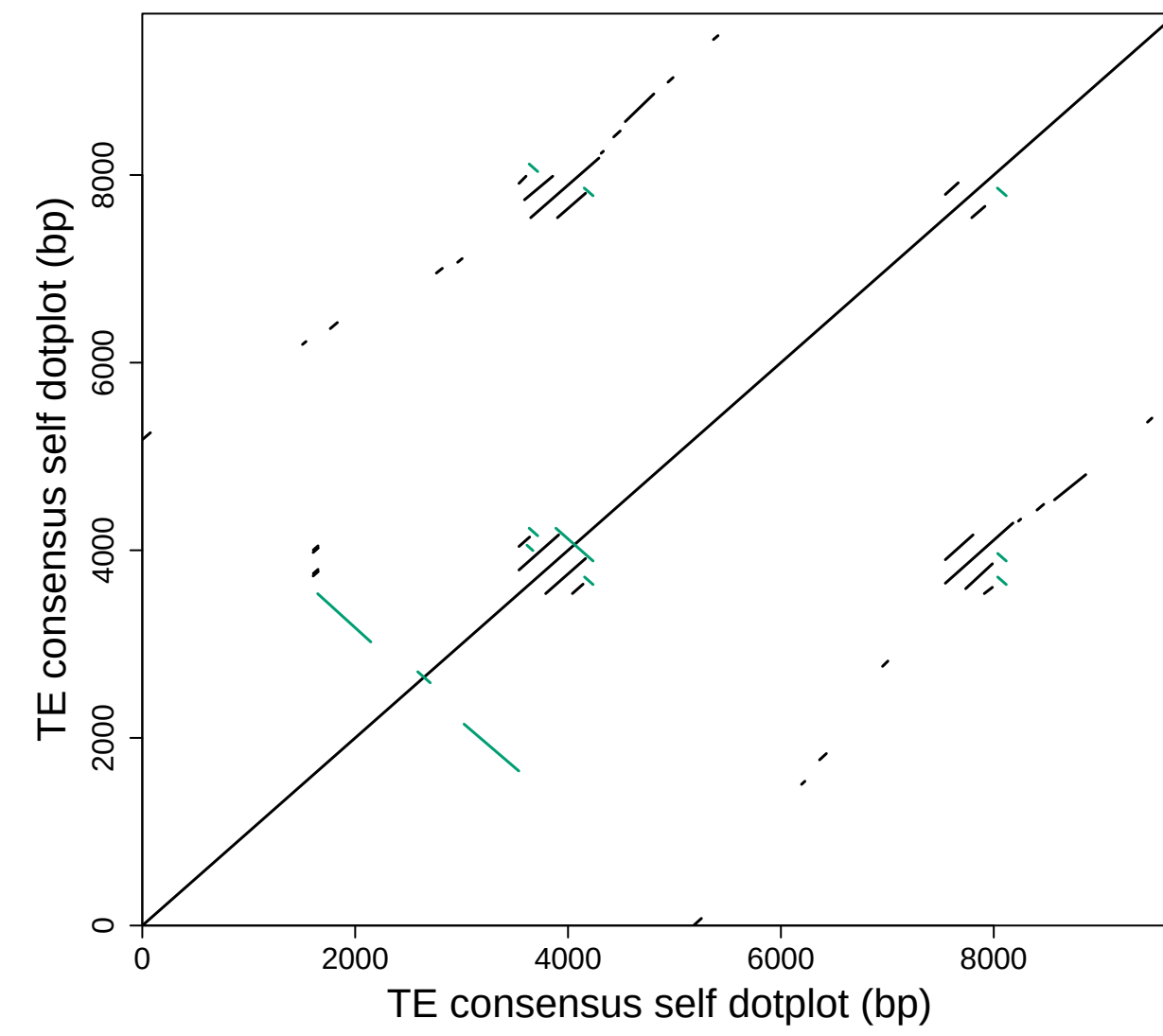
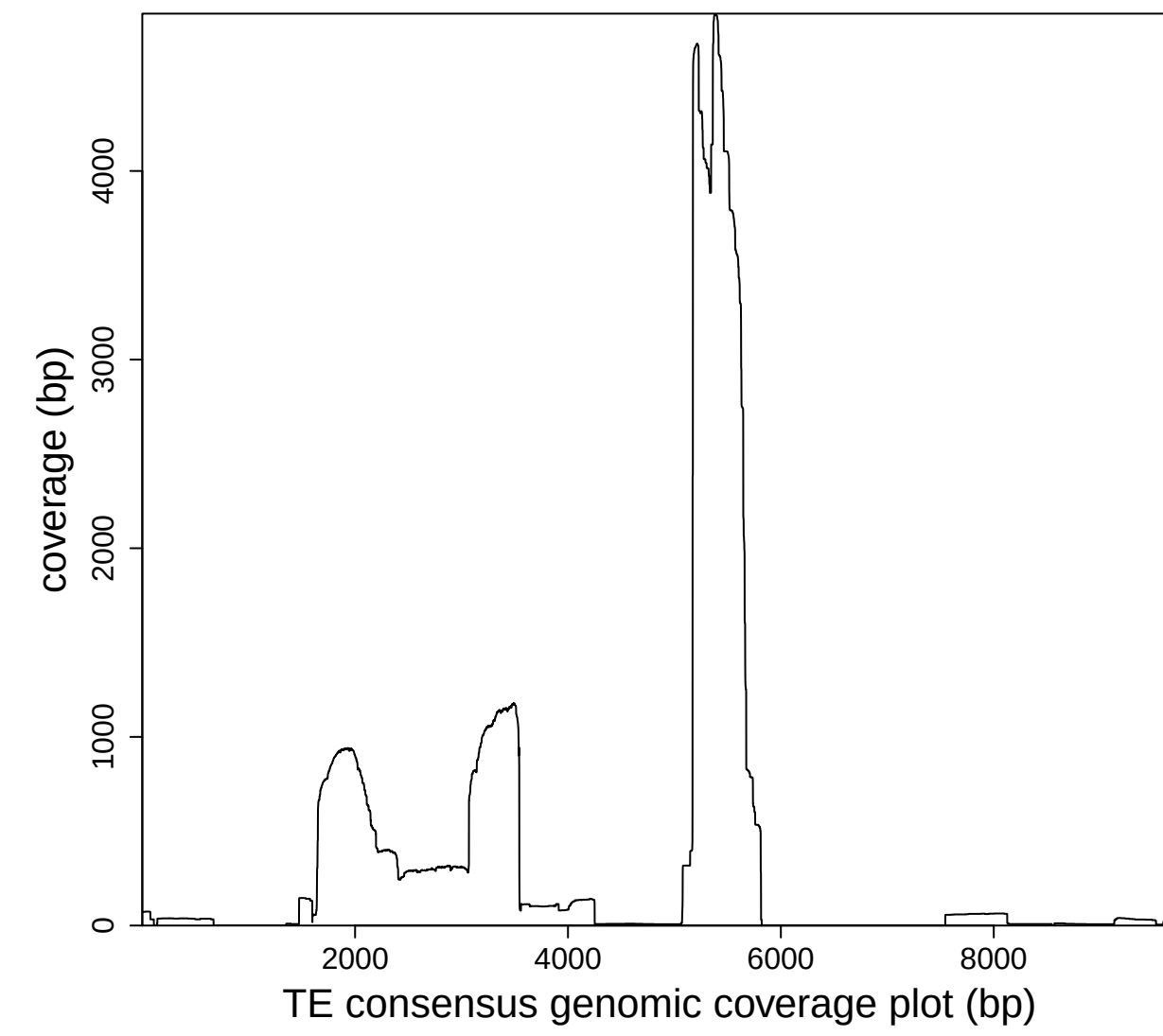
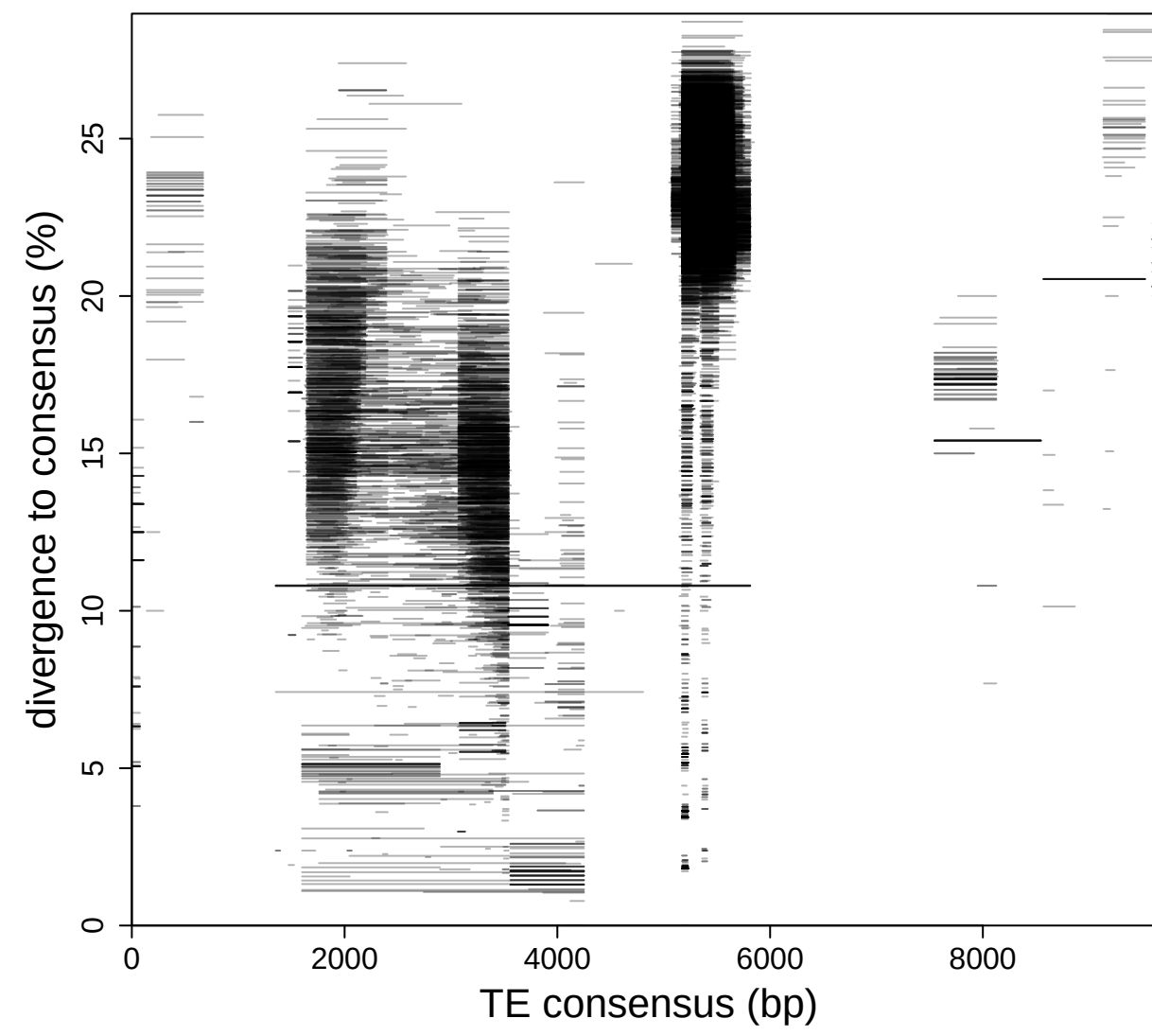
l.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 8675bp; fragments: 25404; full length: 0 (>=7807.5bp)

After TEtrimmer 8675 bp



9.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 9720bp; fragments: 8903; full length: 0 (>=8748bp)

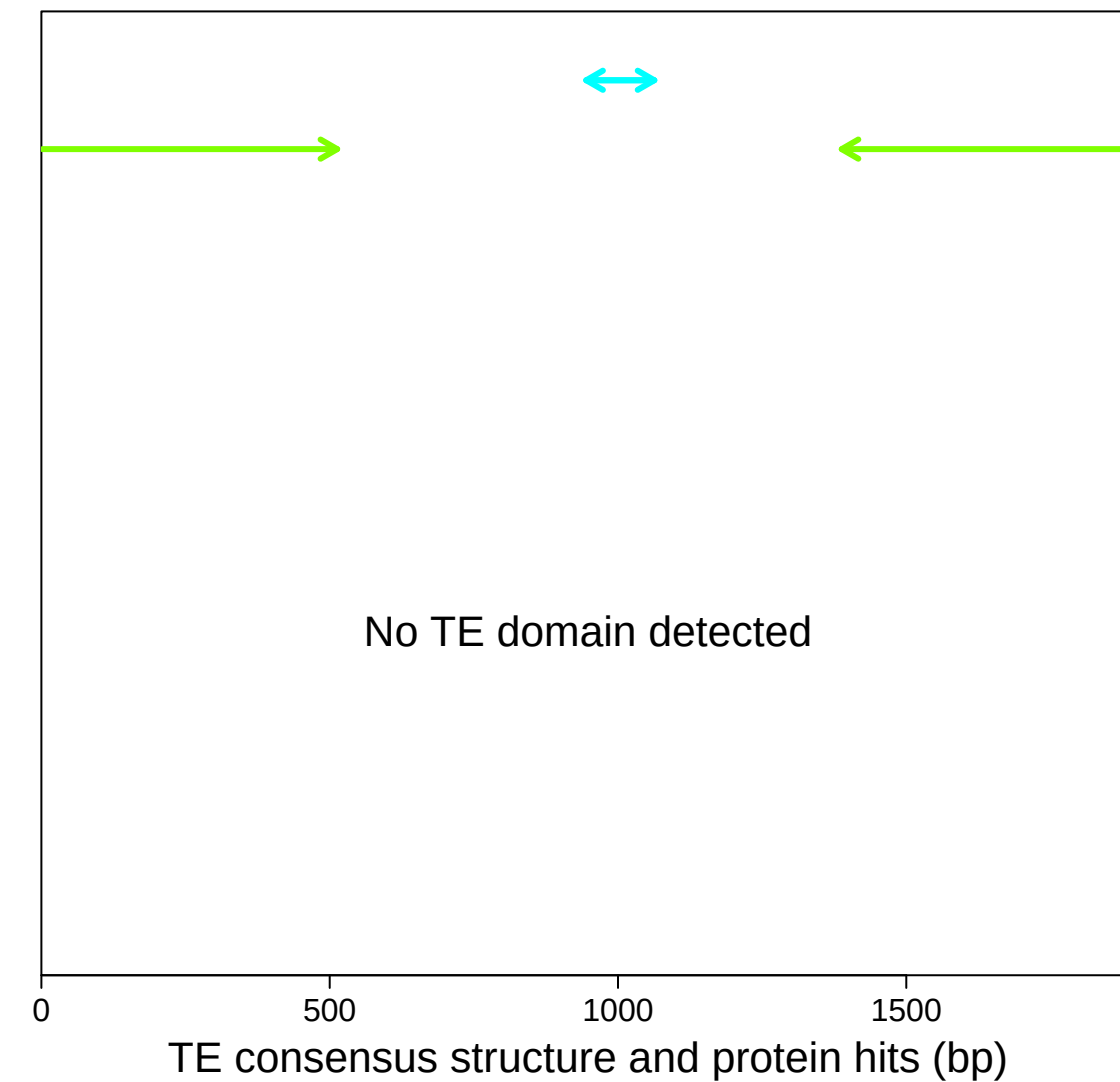
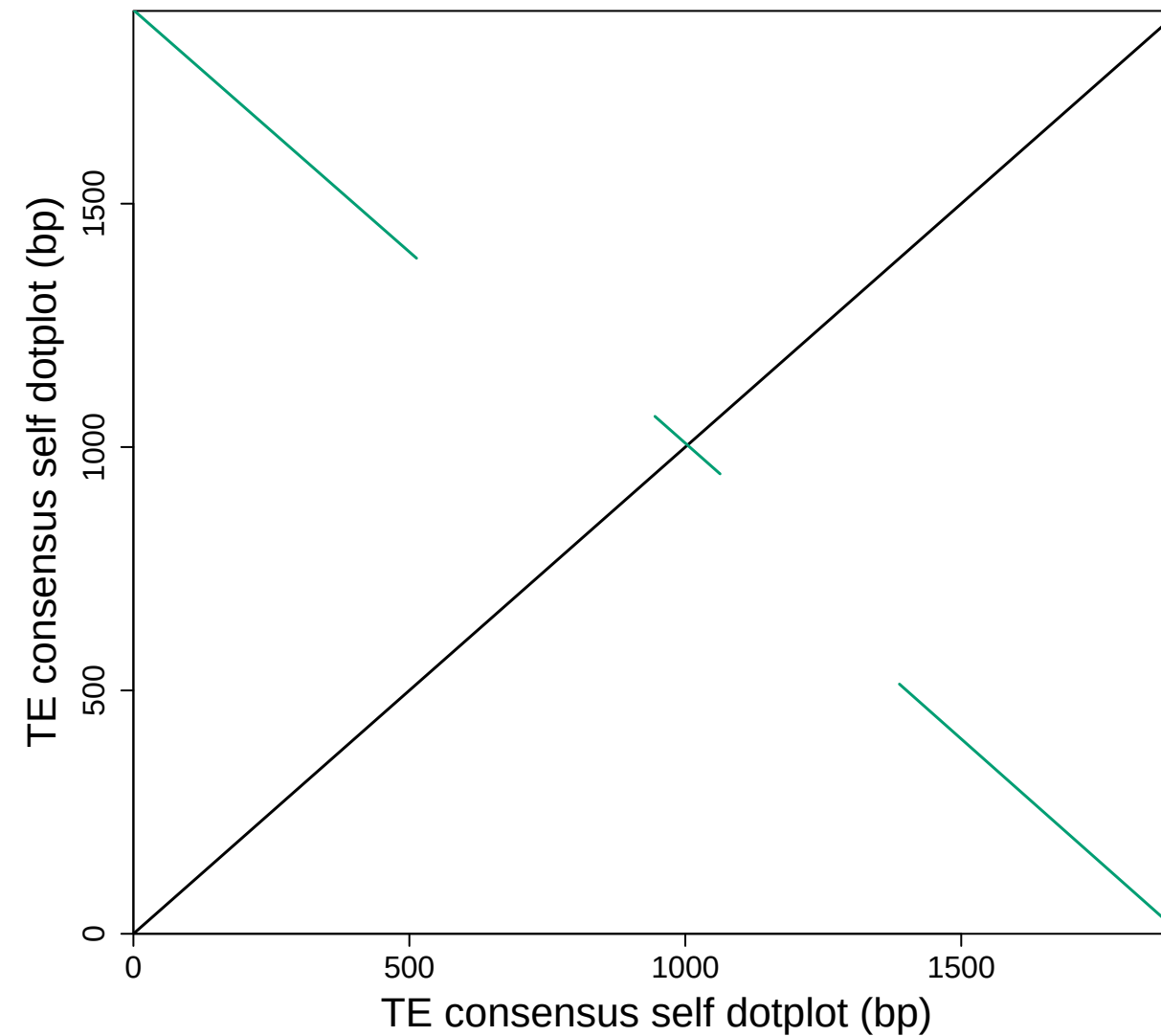
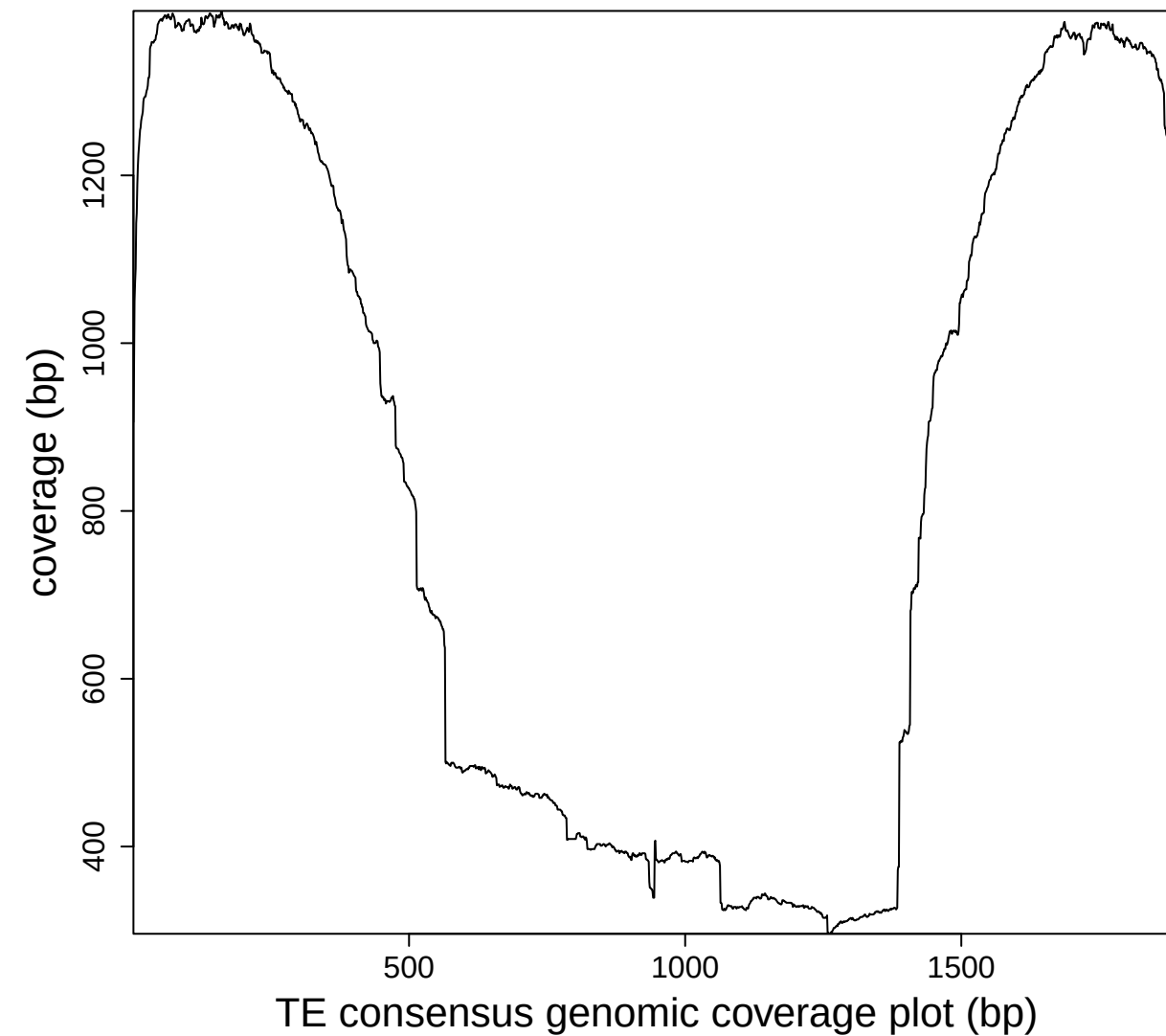
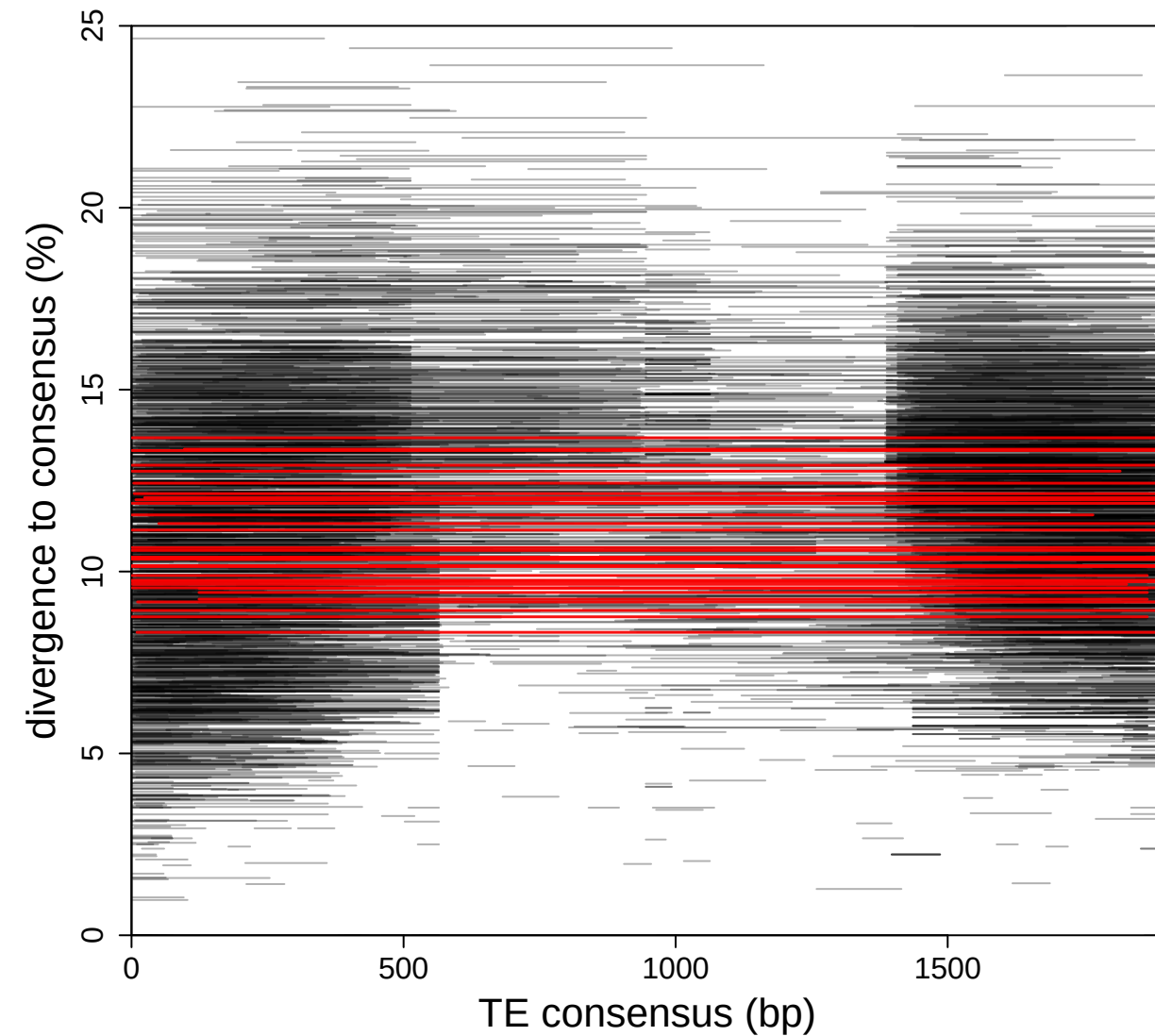
After TEtrimmer Extended plot Blue lines are boundaries



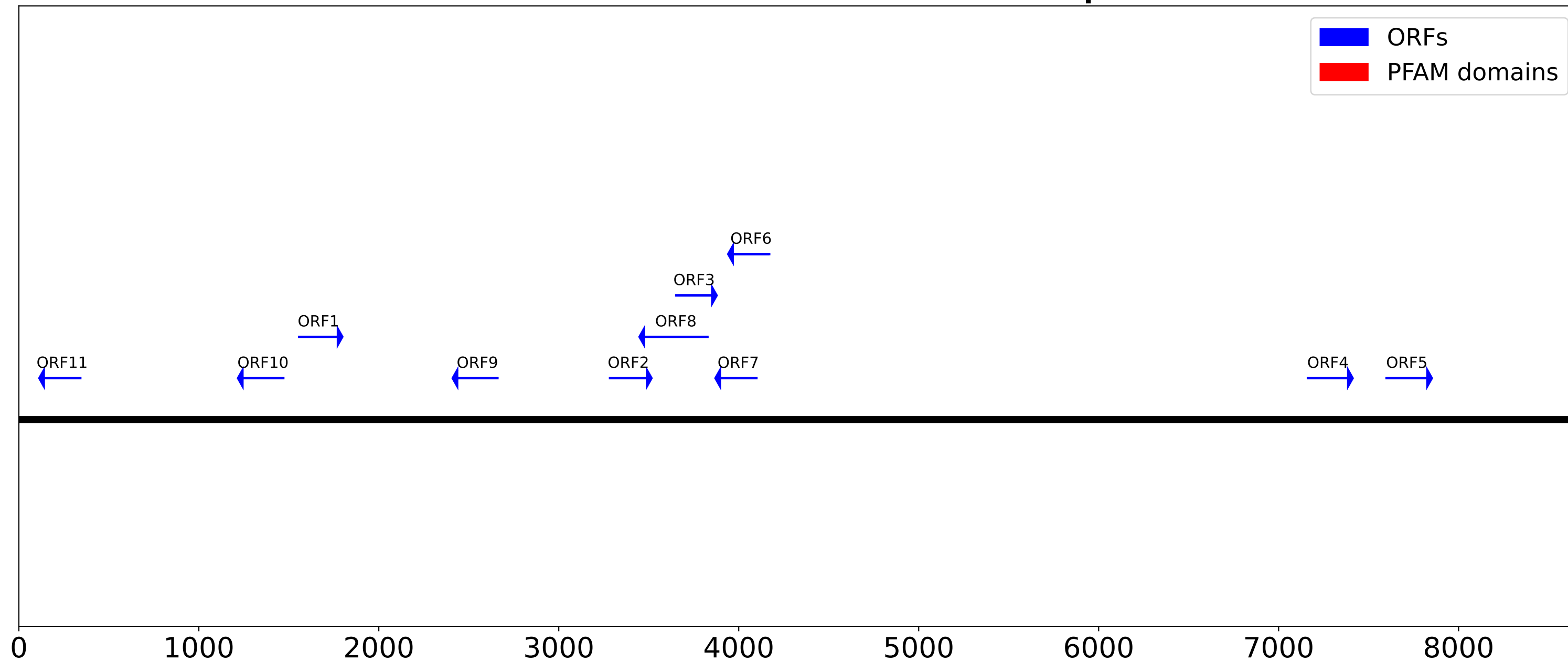
Before TEtrimmer 1896 bp

TE: rnd_4_family_379

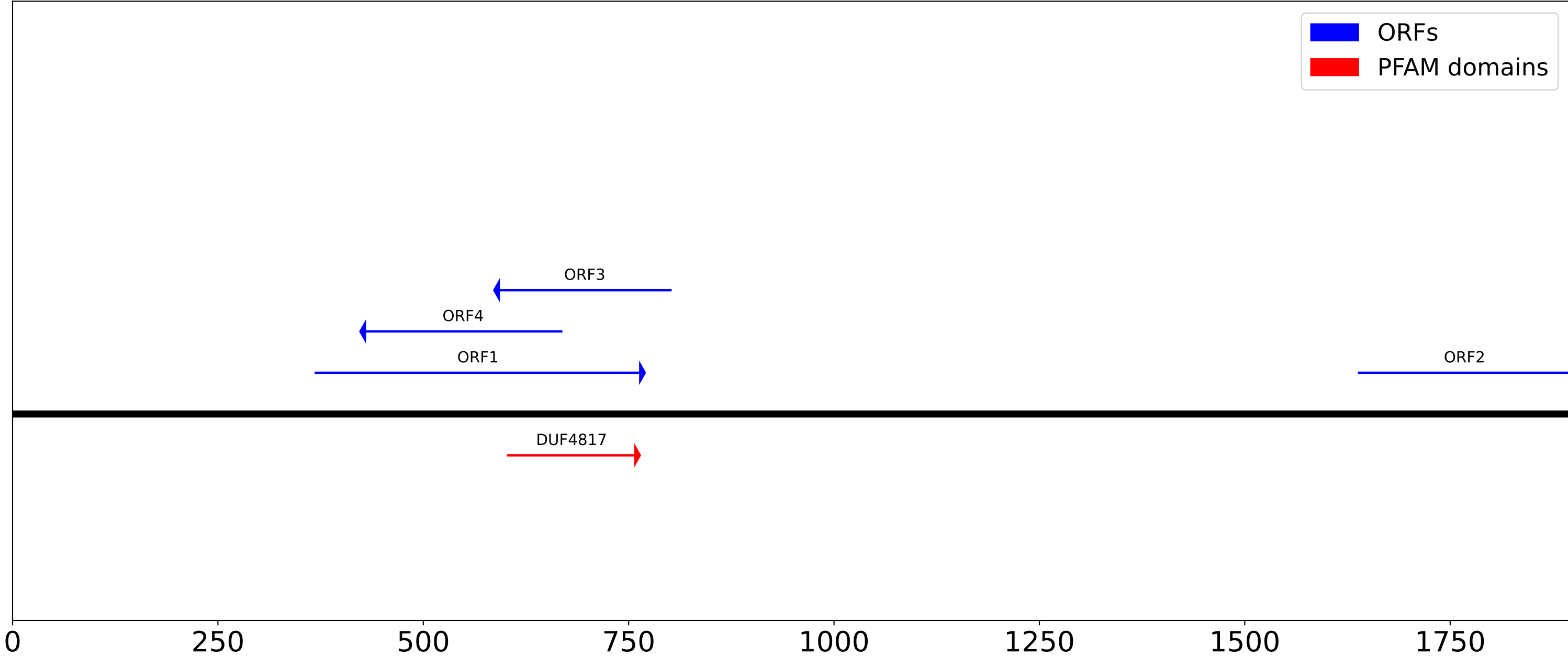
size: 1896bp; fragments: 3918; full length: 40 (≥ 1706.4 bp)



After TEtrimmer ORF and PFAM domain plot



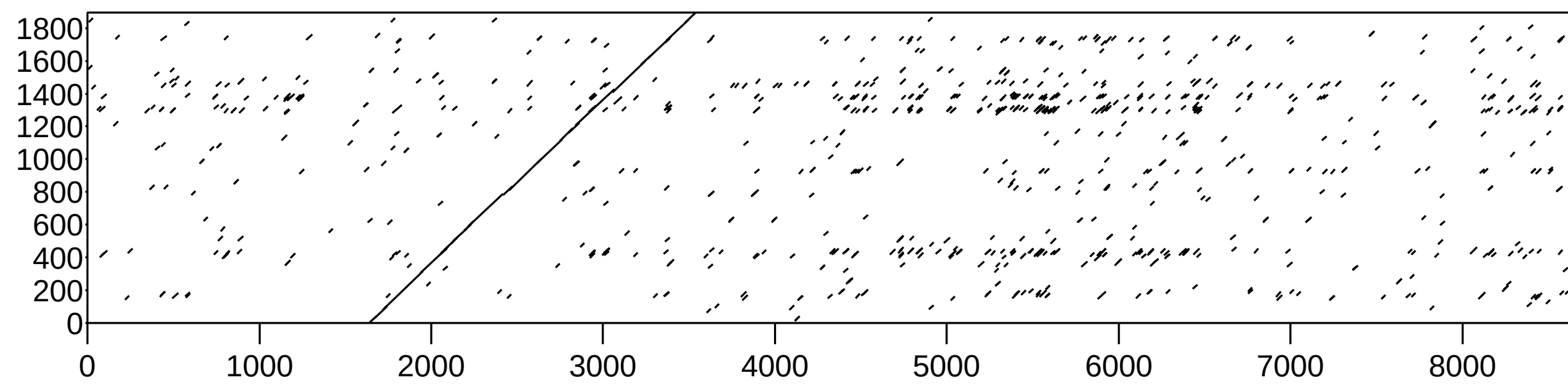
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)