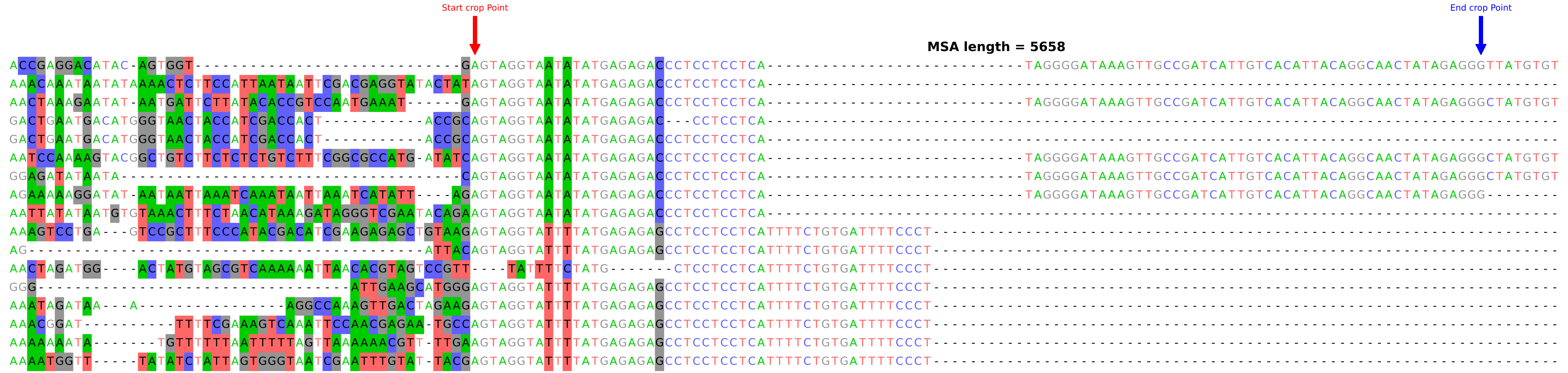
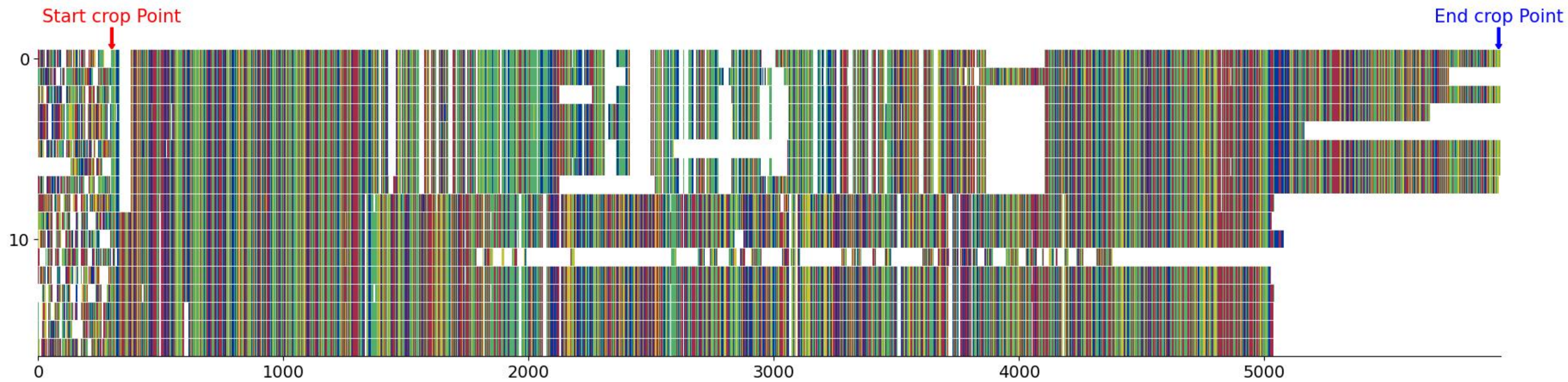




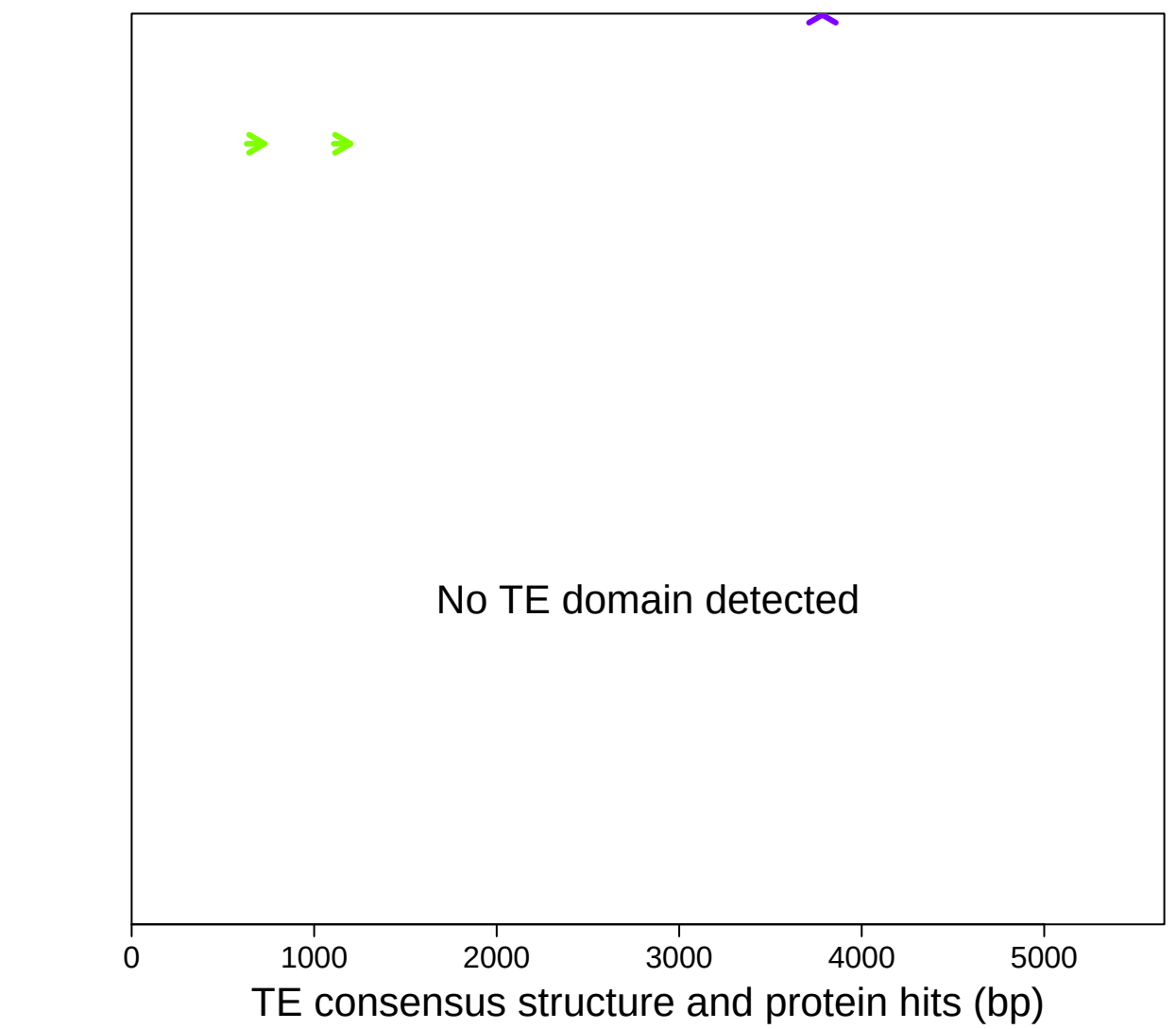
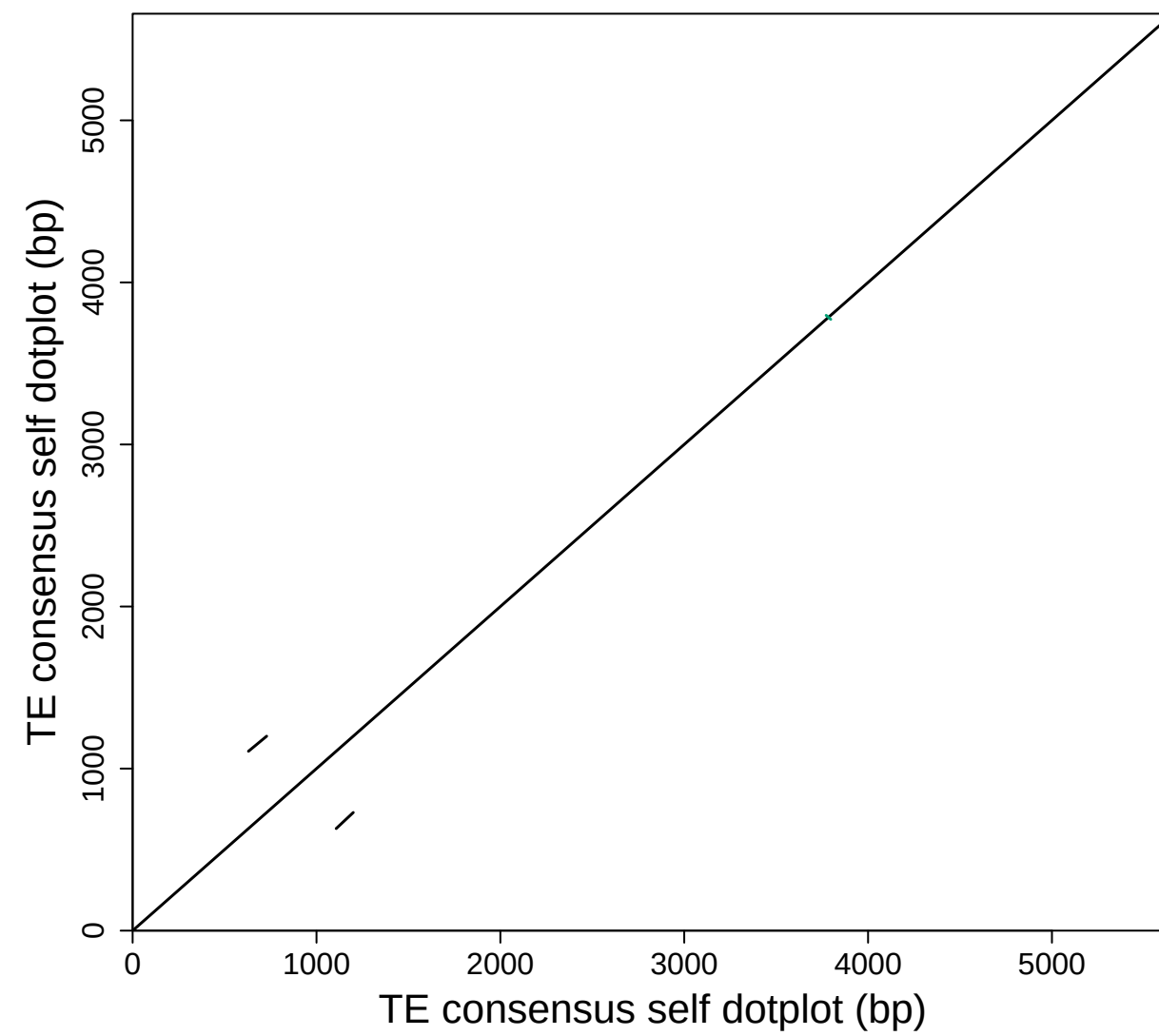
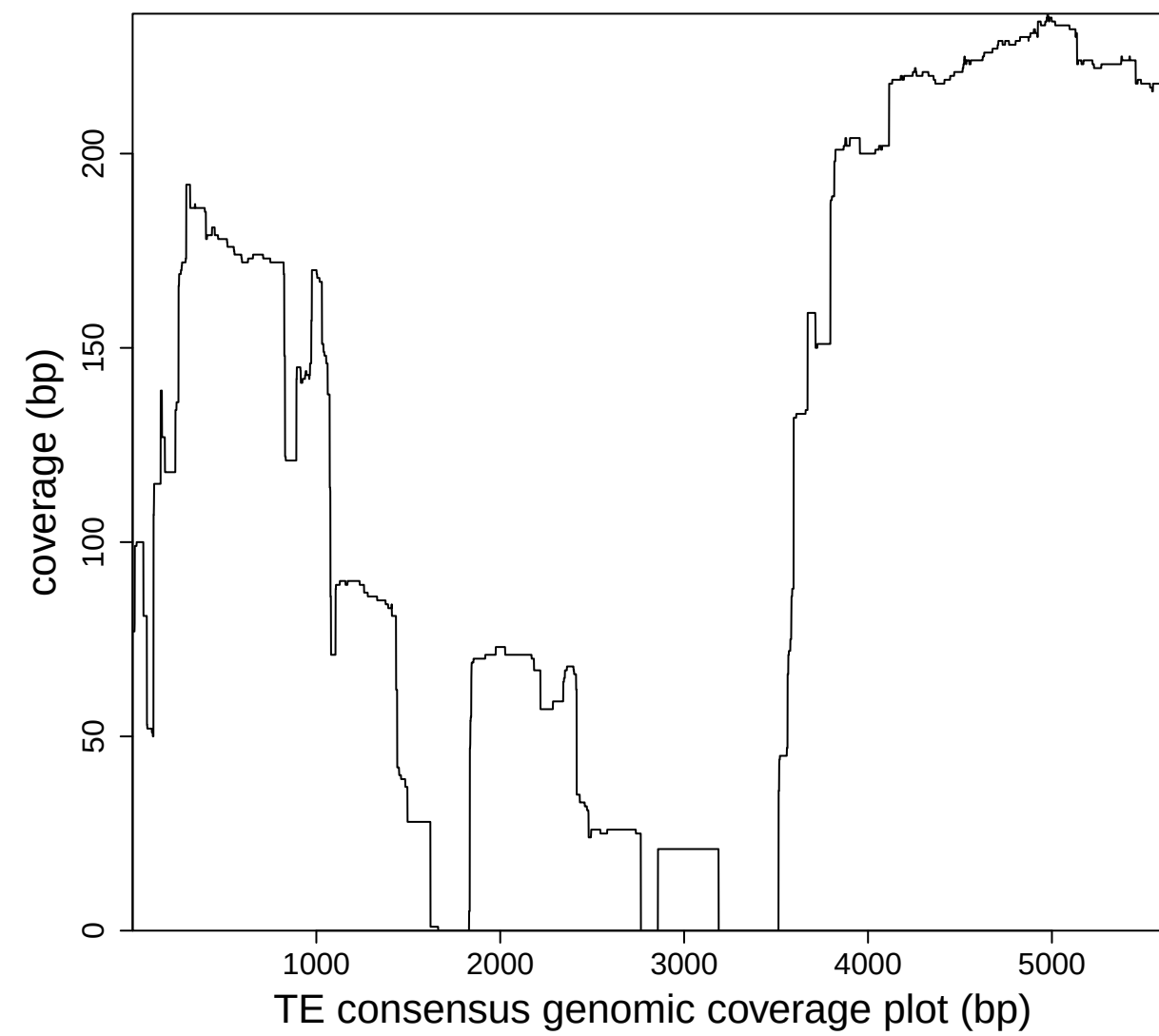
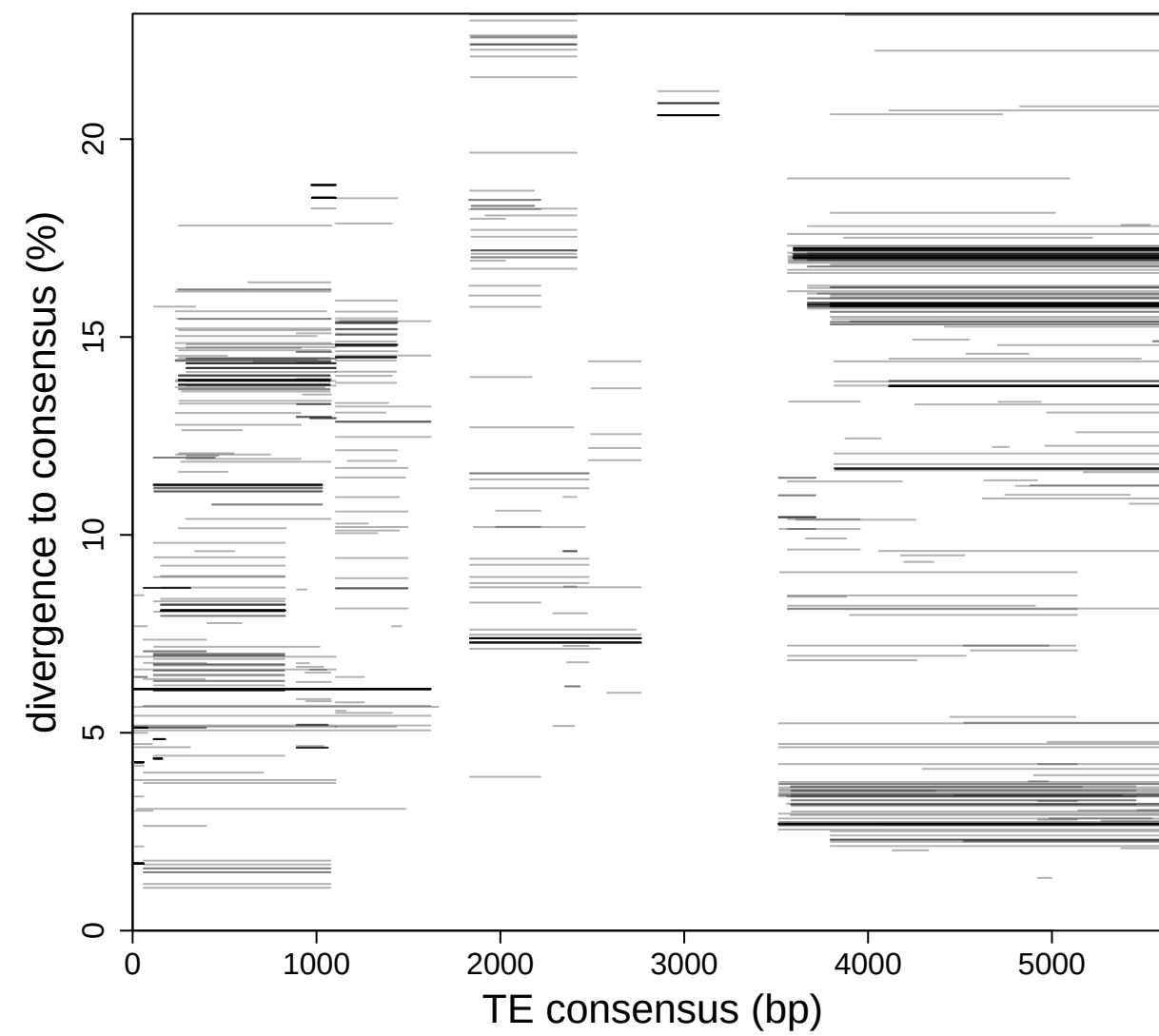
Start crop Point

End crop Point



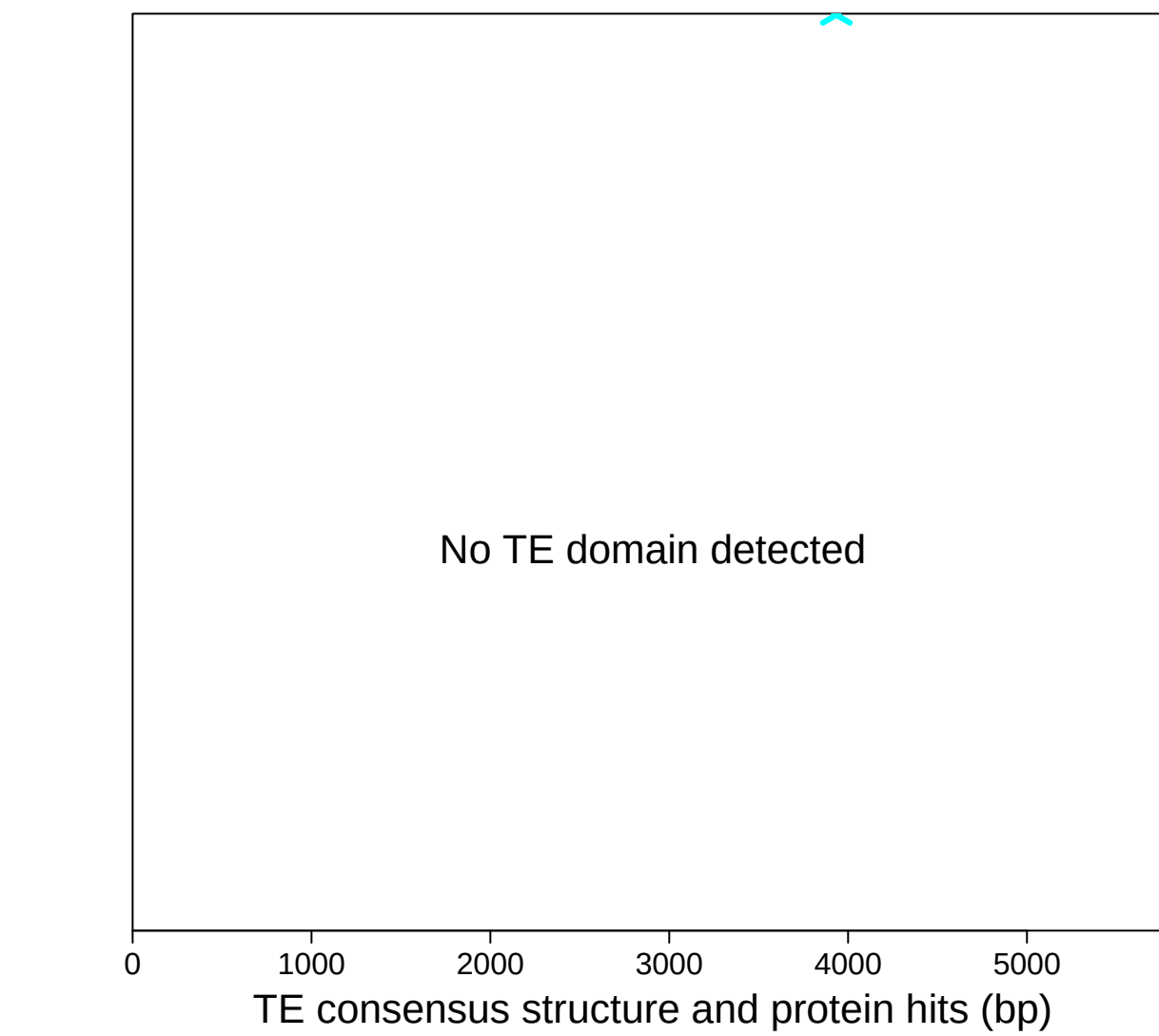
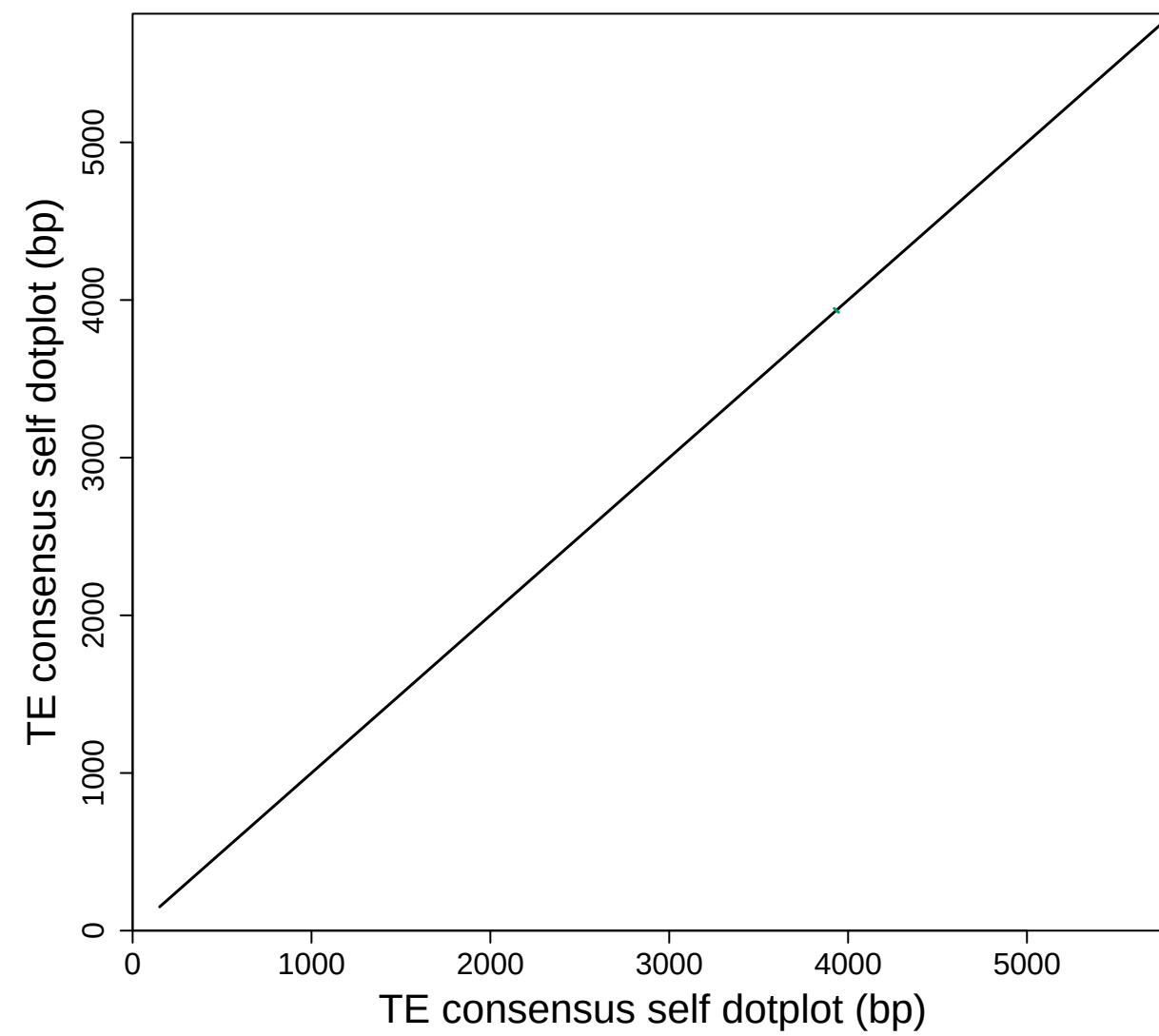
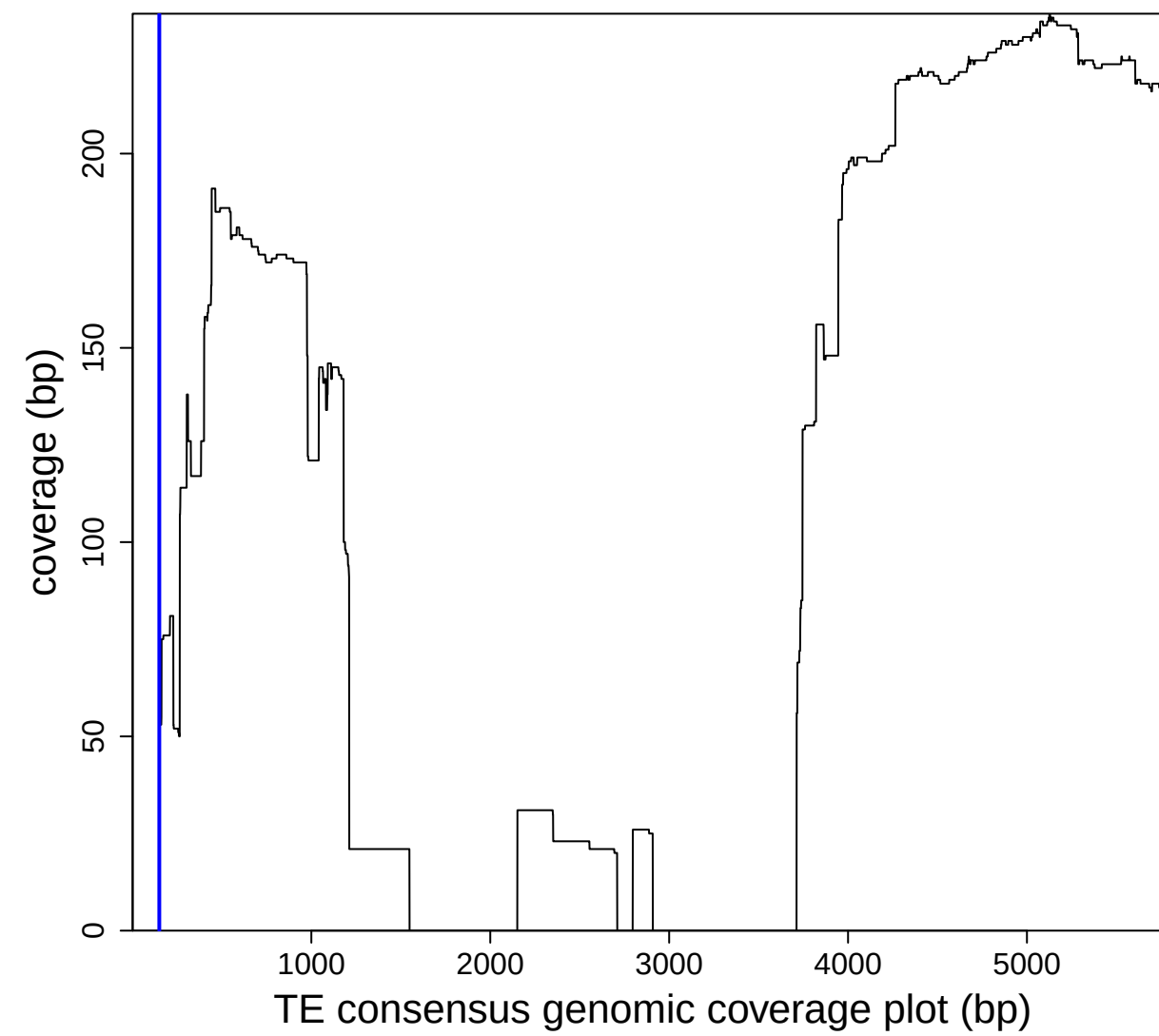
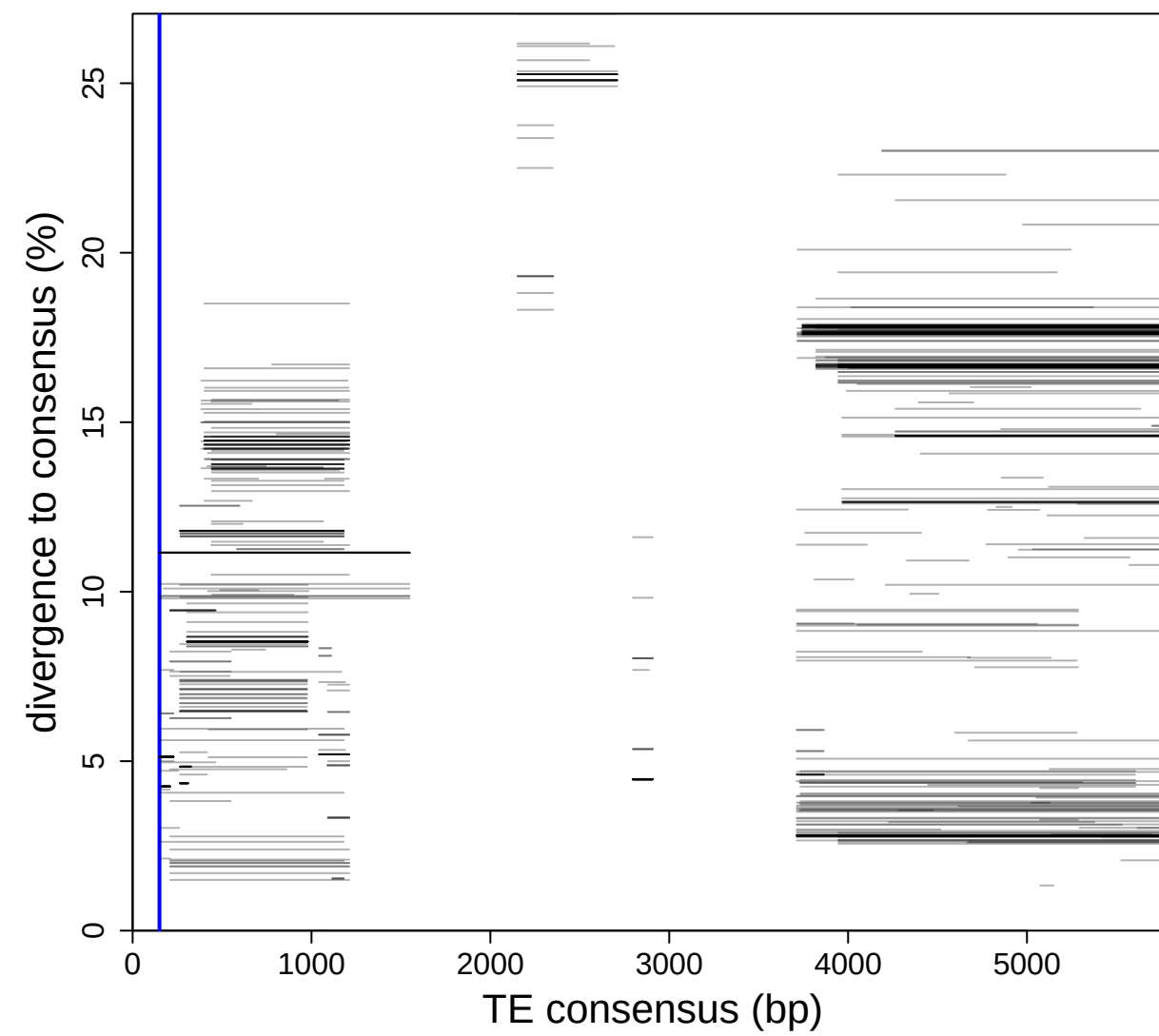
l.b.bed uf.bed g_4.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 5658bp; fragments: 815; full length: 0 (>=5092.2bp)

After TEtrimmer 5658 bp



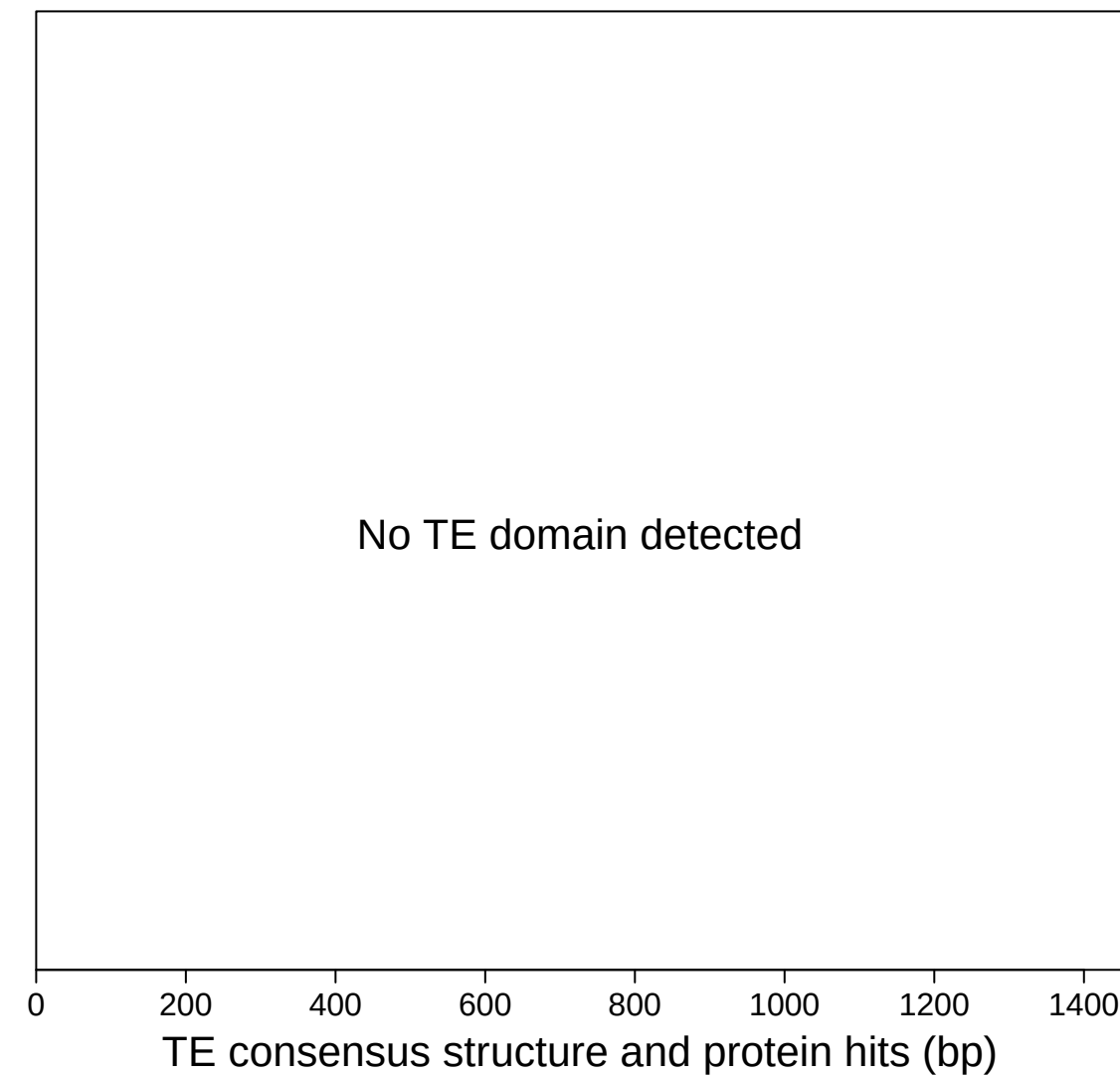
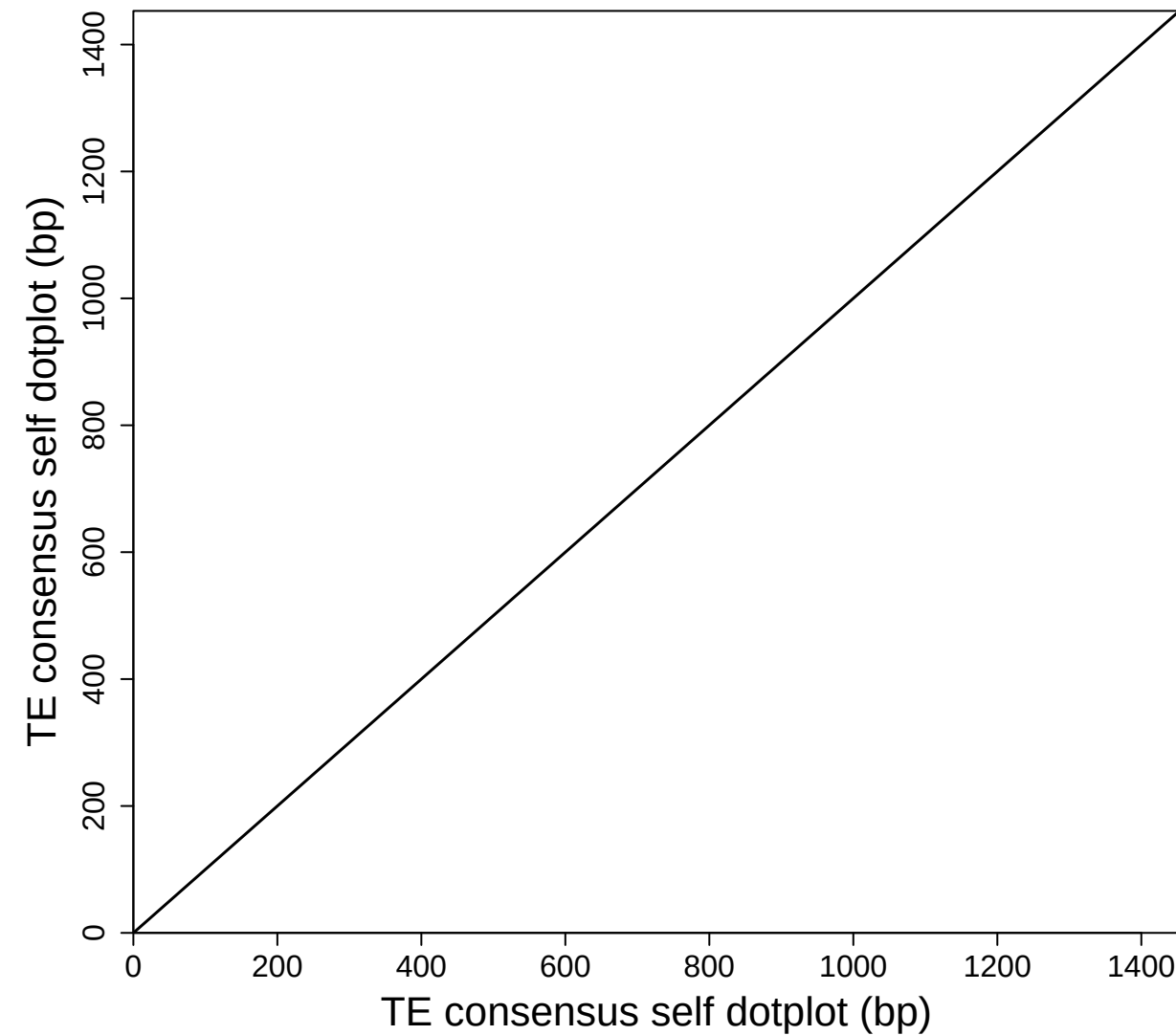
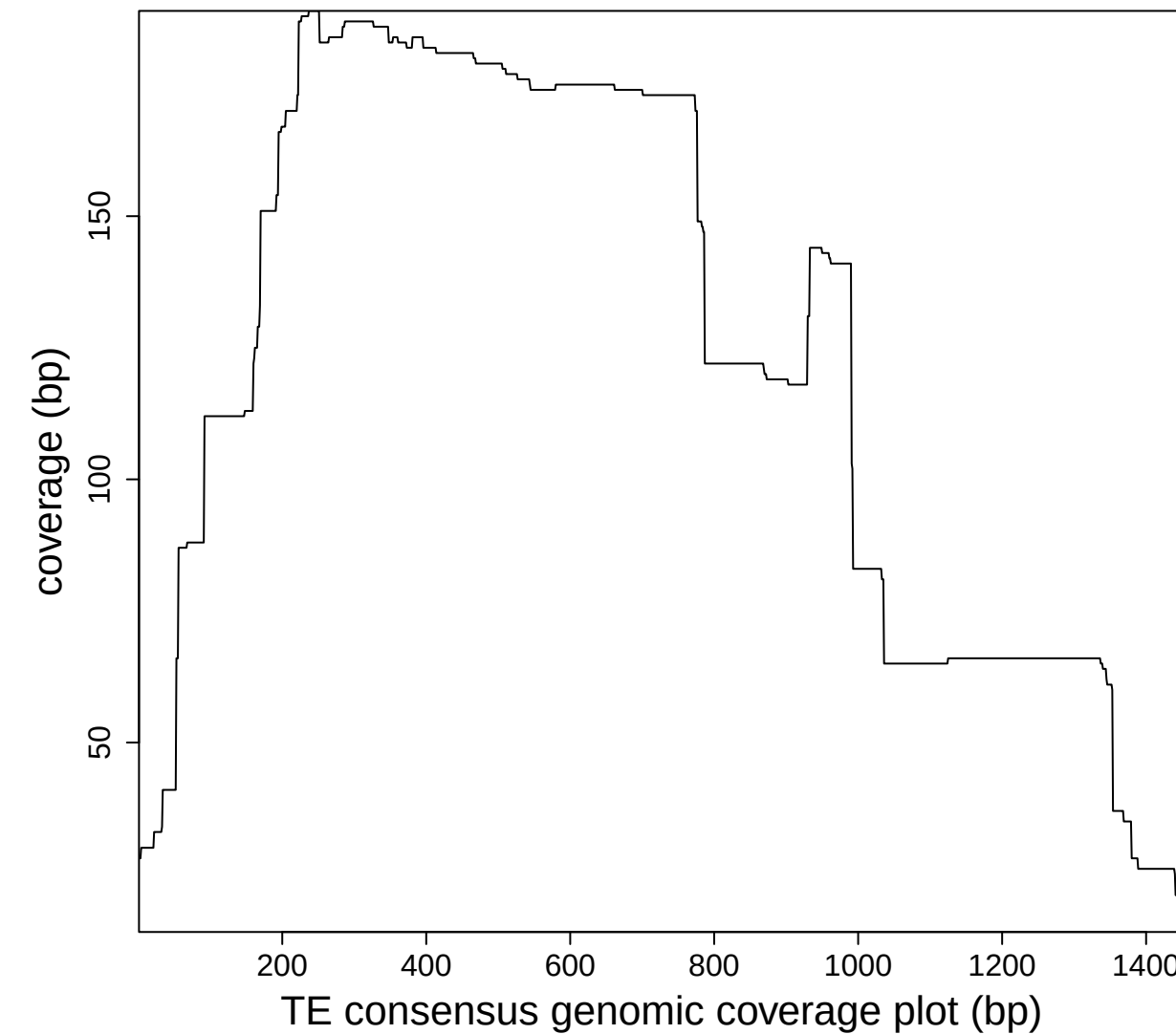
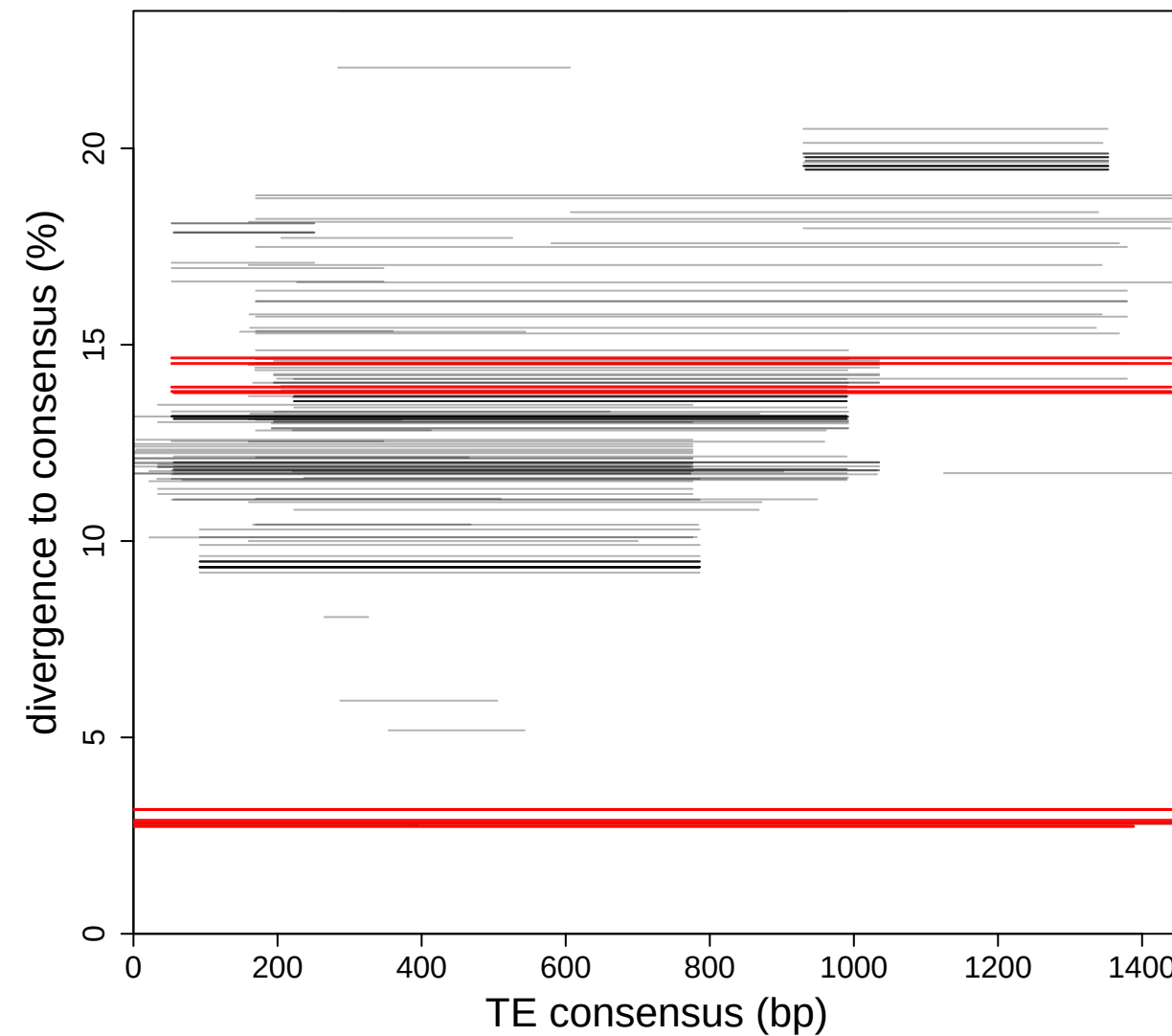
8.fasta.b.bed uf.bed g 4.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 5816bp; fragments: 649; full length: 0 (>=5234.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

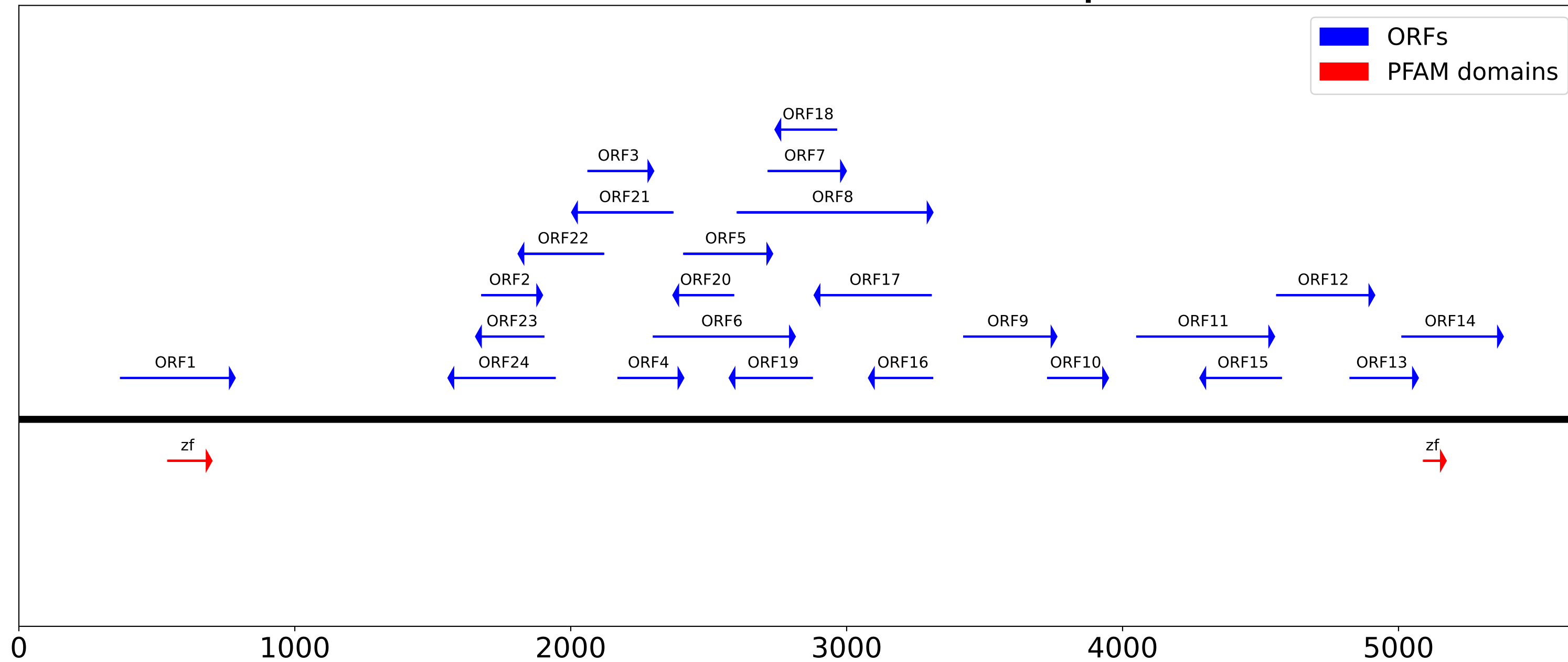


TE: rnd_5_family_588
size: 1453bp; fragments: 225; full length: 19 (≥ 1307.7 bp)

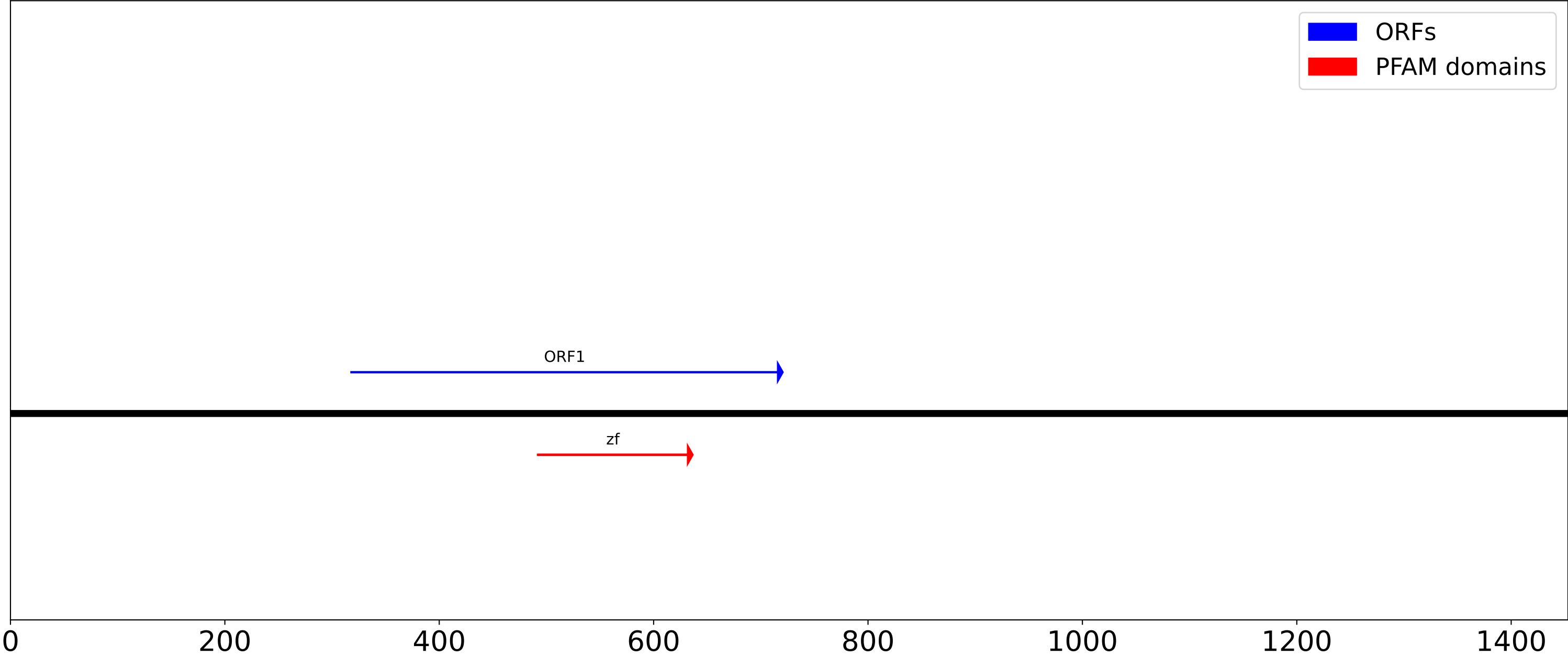
Before TEtrimmer 1453 bp



After TETrimmer ORF and PFAM domain plot



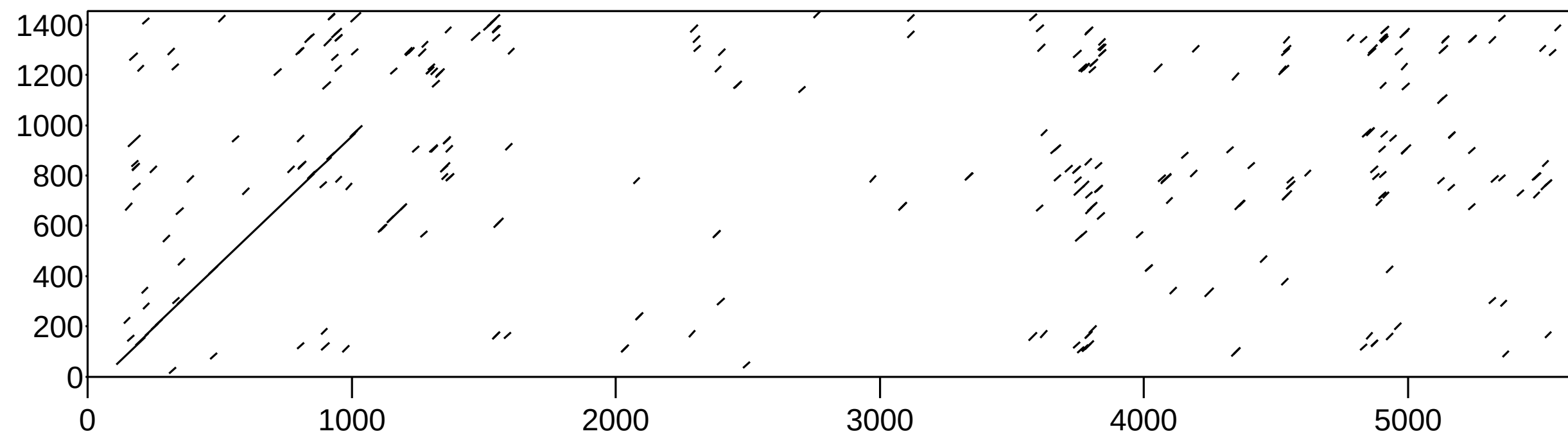
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)