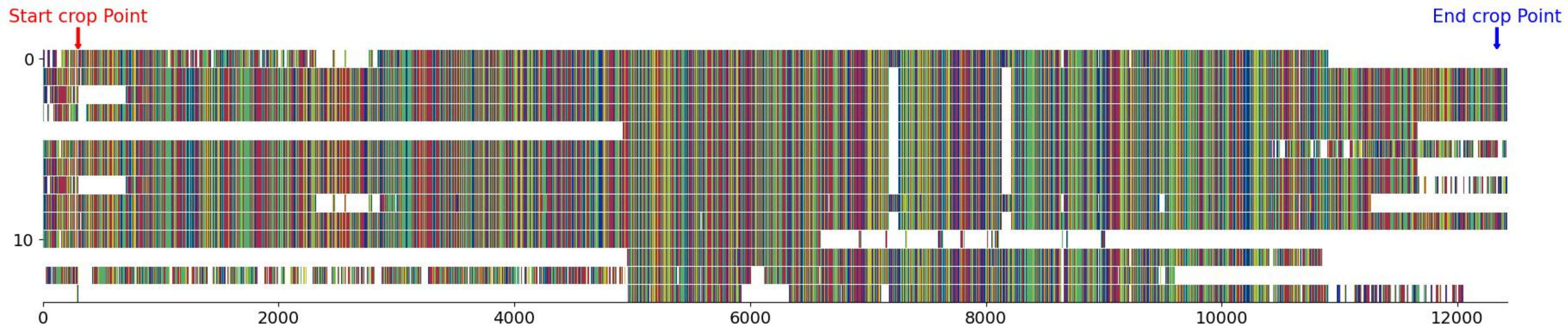
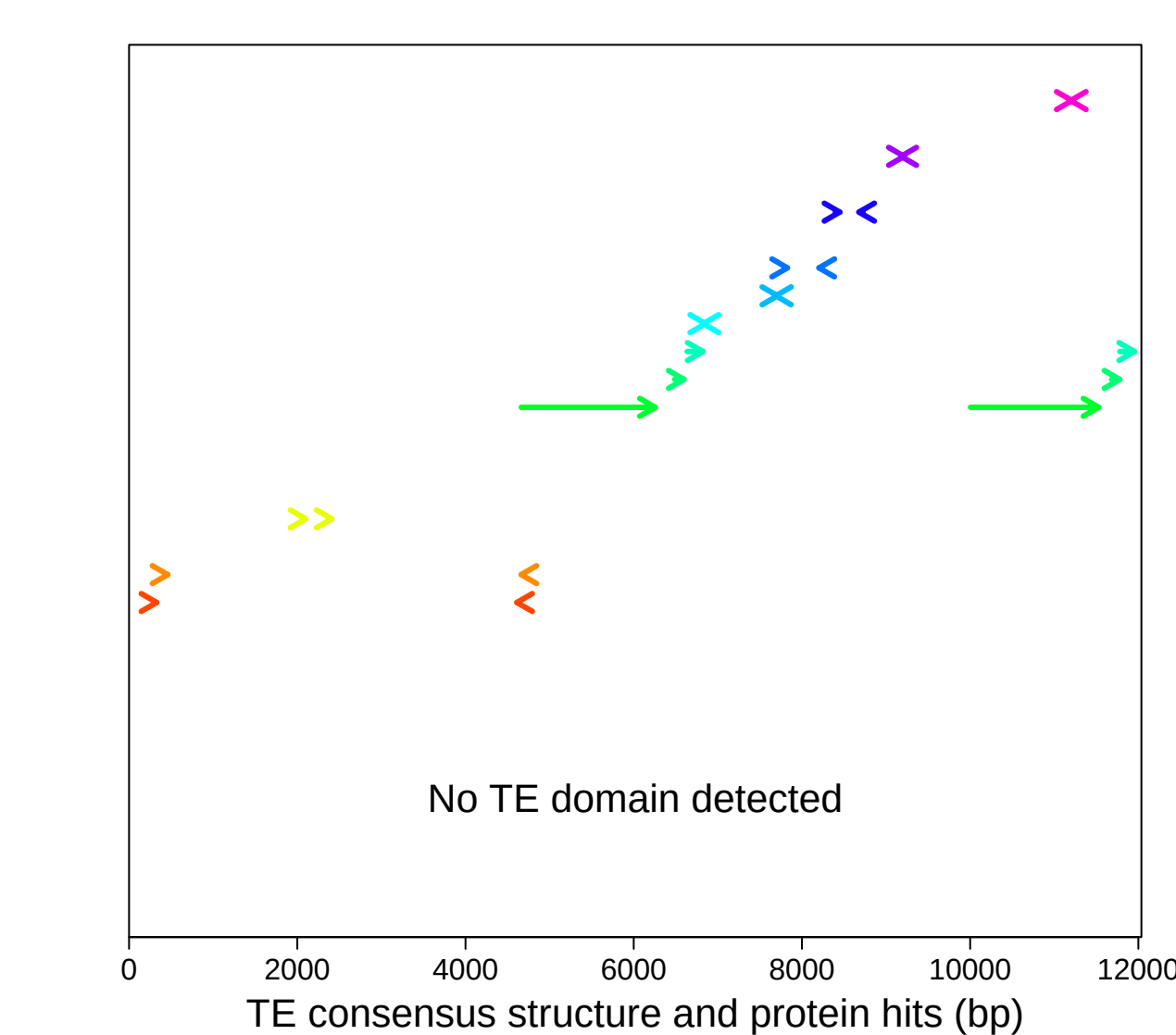
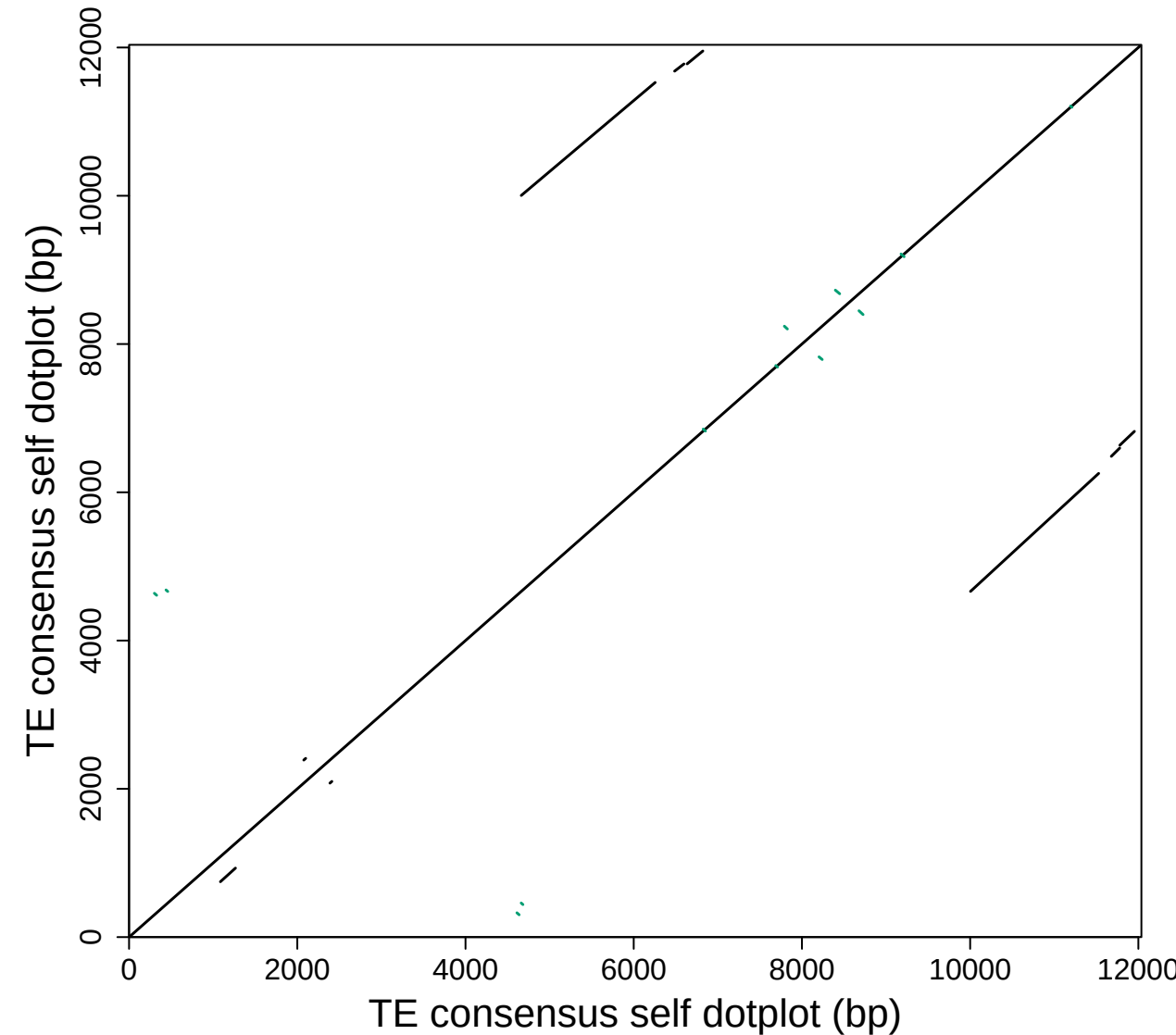
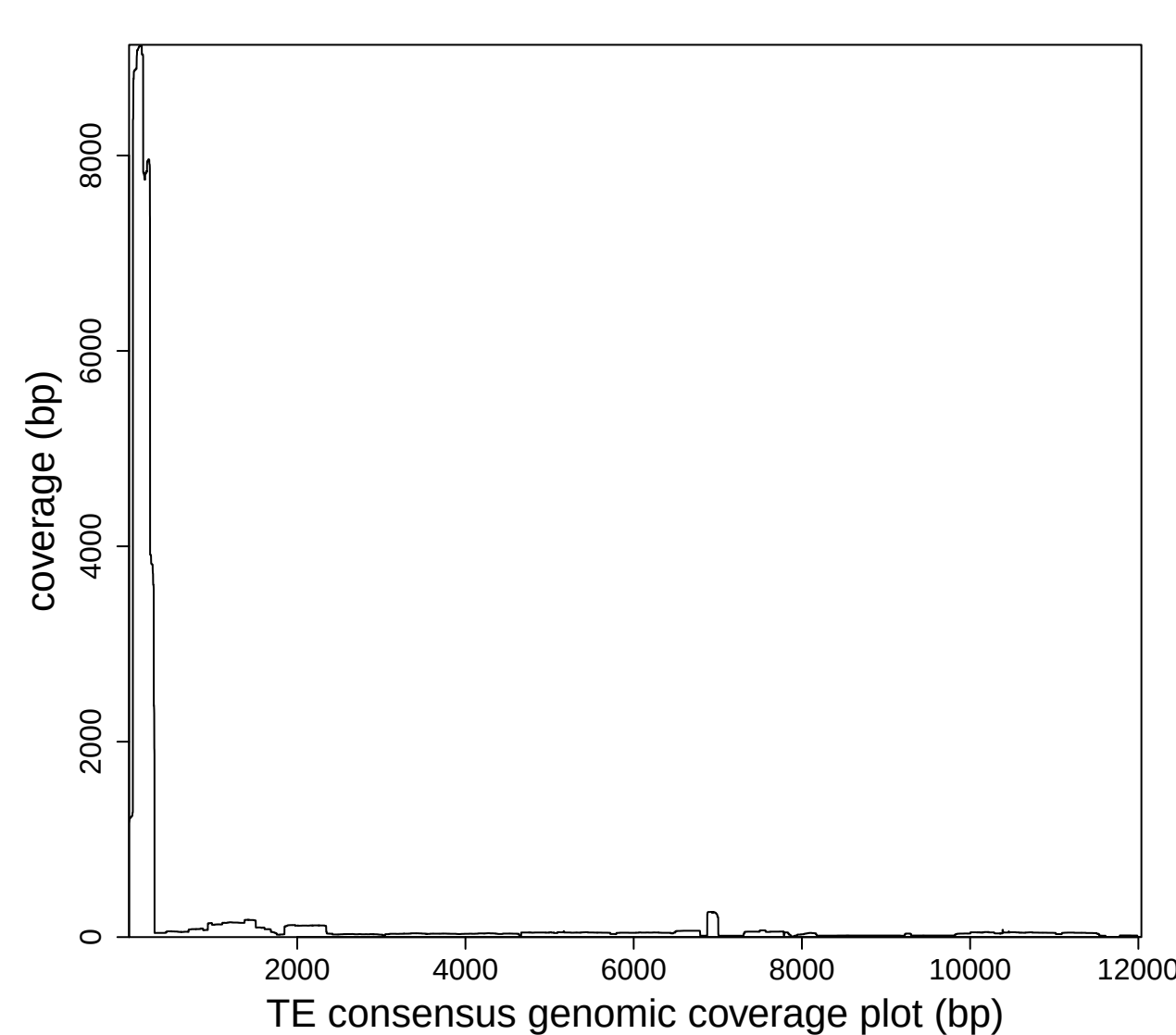
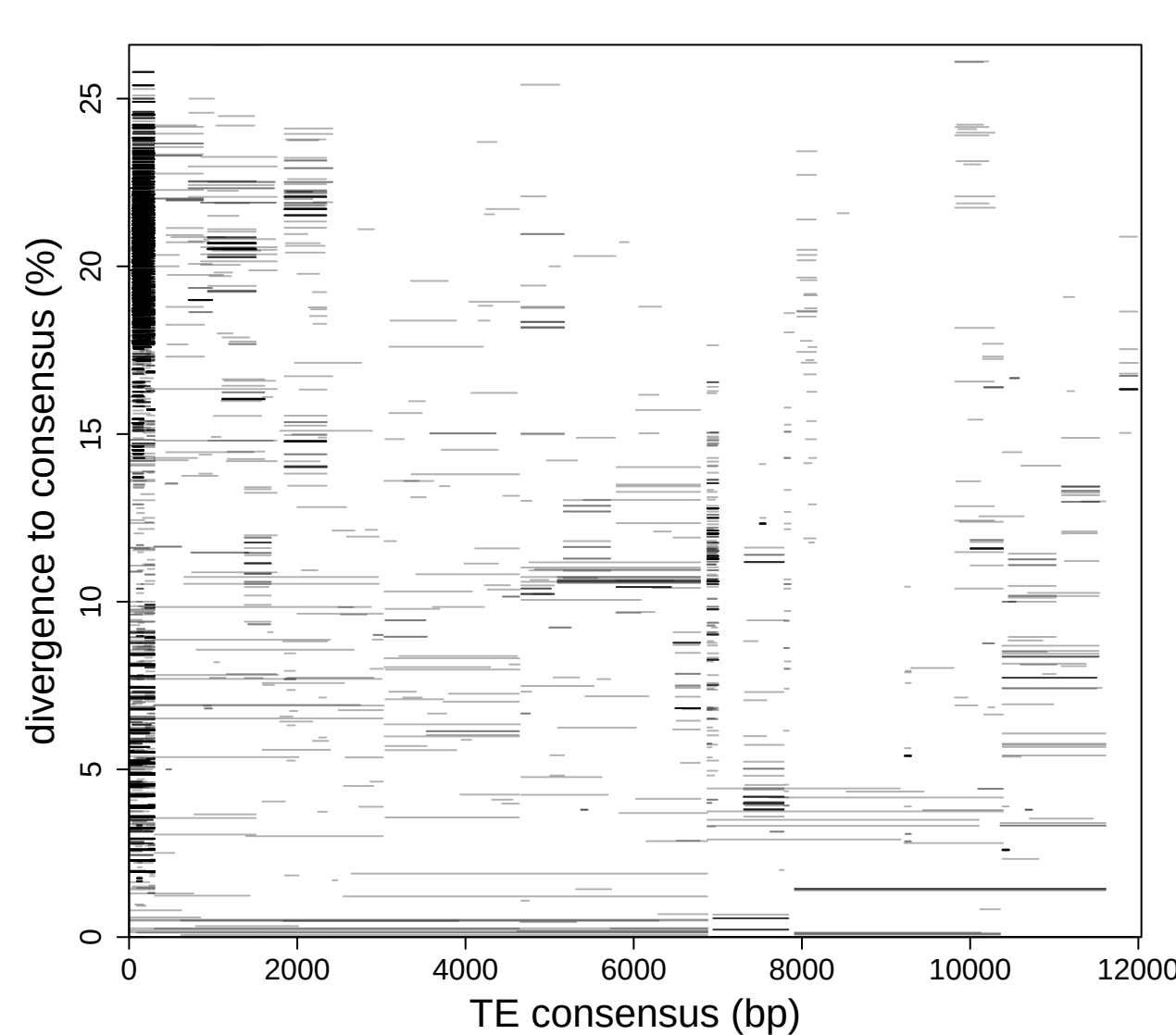




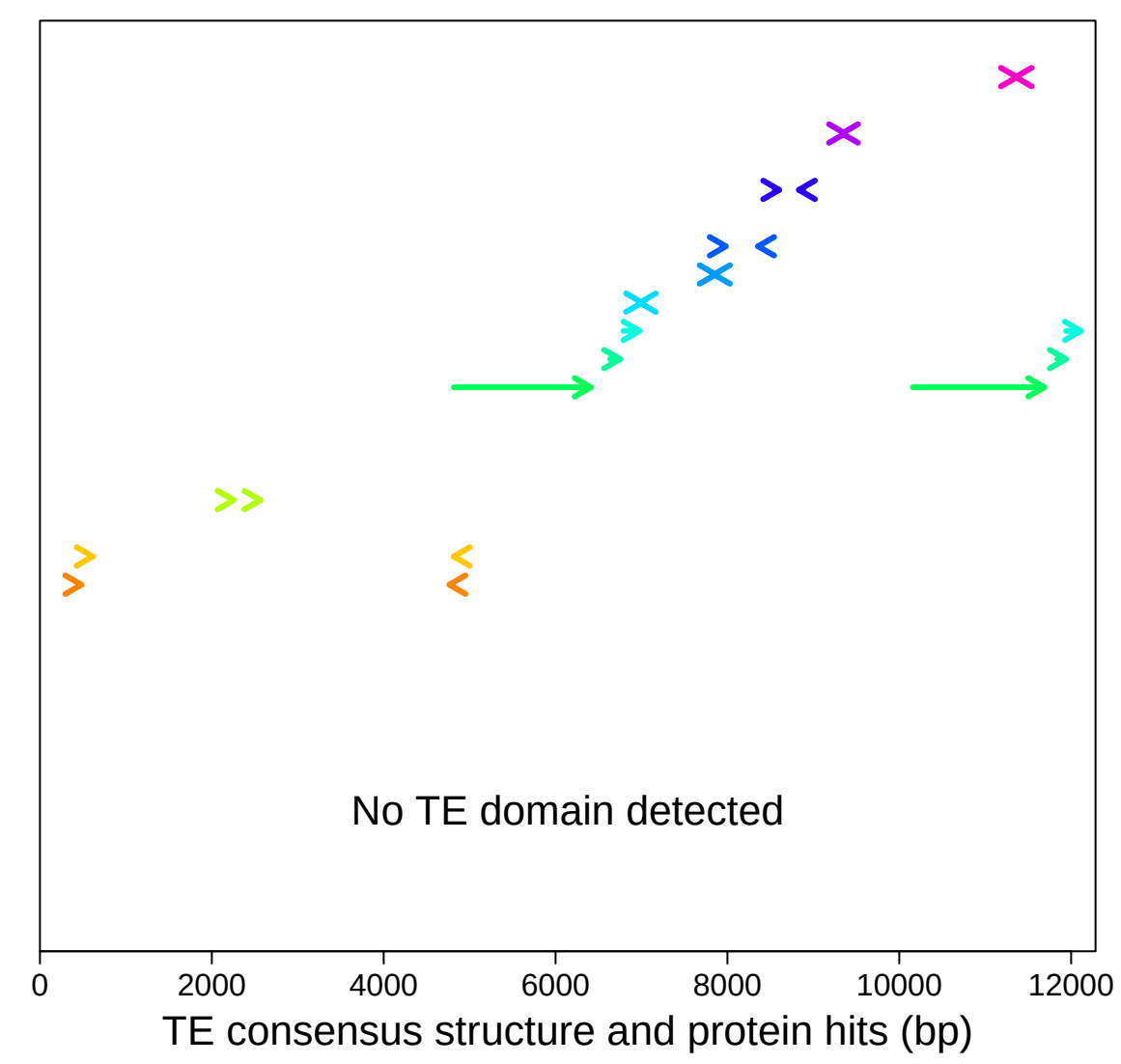
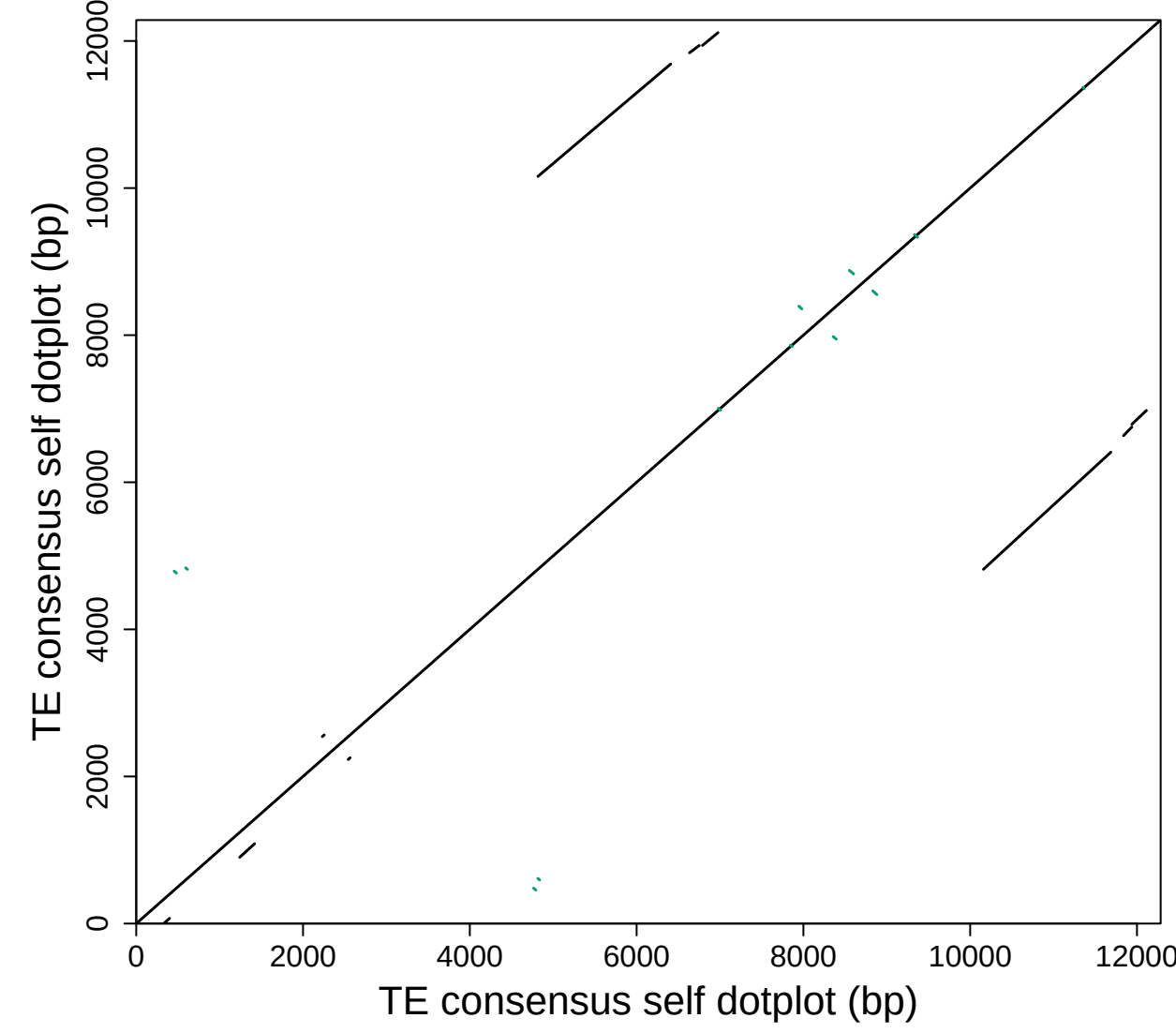
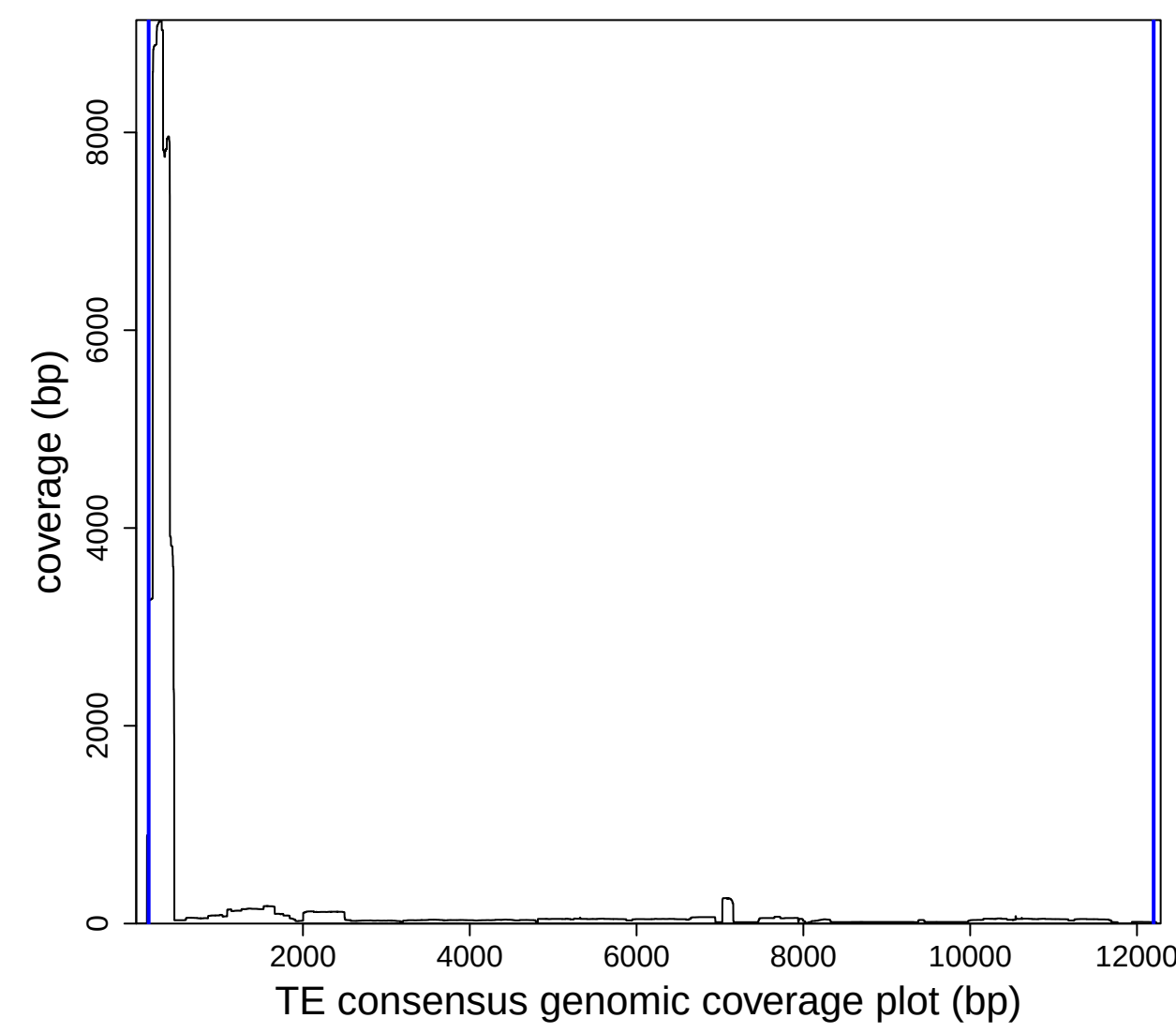
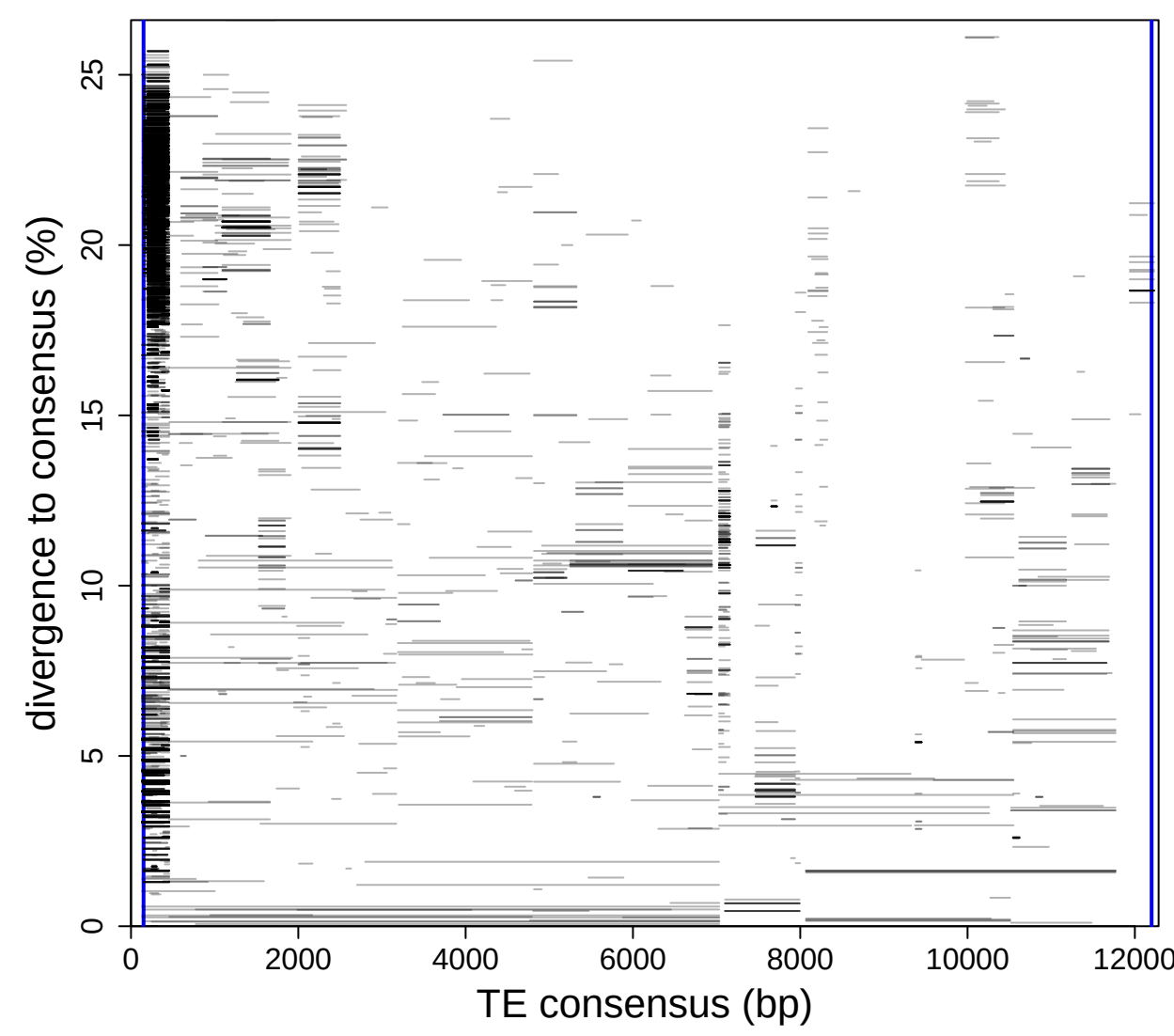
l.b.bed uf.bed g_3.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 12036bp; fragments: 10564; full length: 0 (>=10832.4bp)

After TEtrimmer 12036 bp

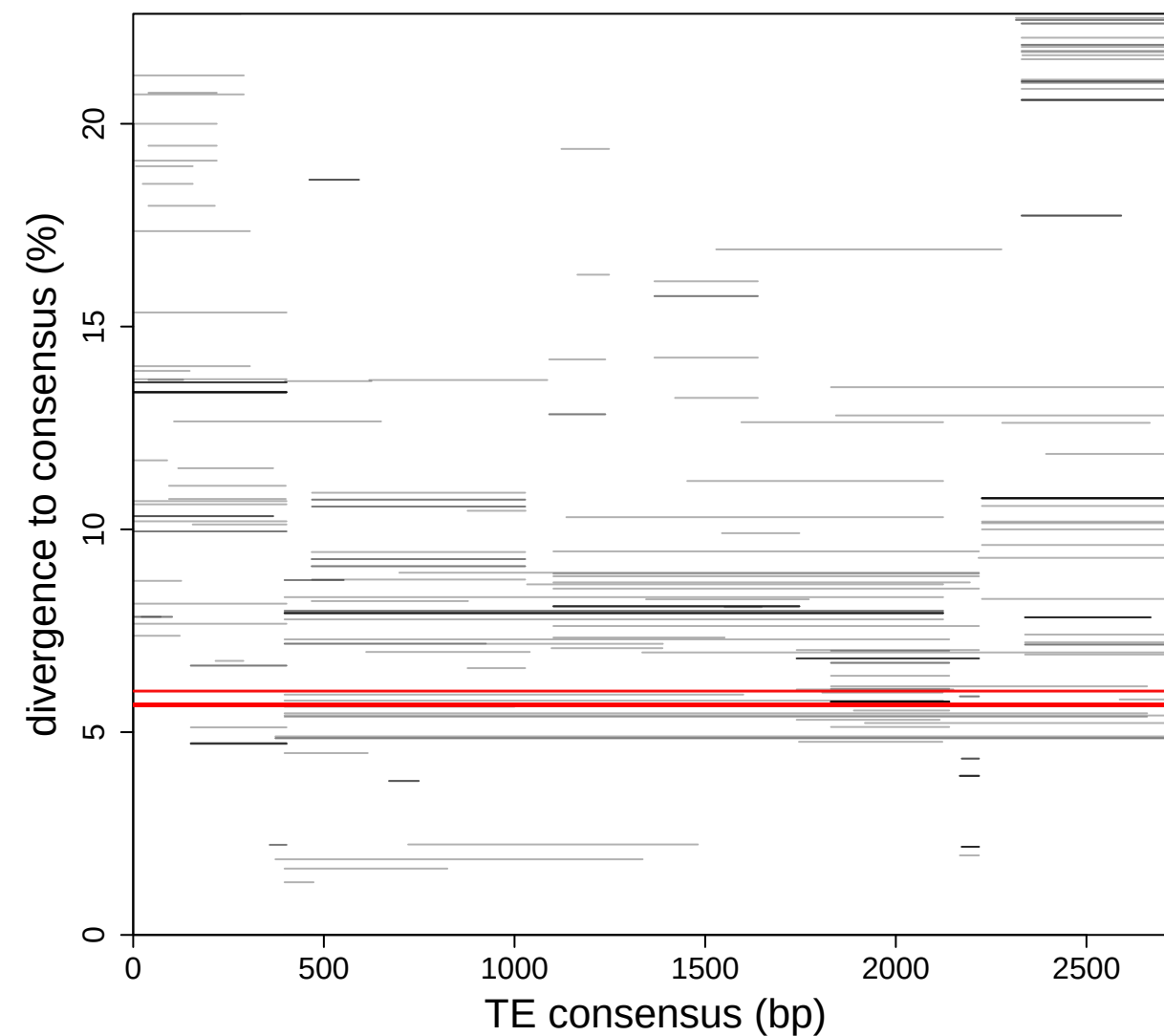


2.fasta.b.bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 12284bp; fragments: 10576; full length: 0 (>=11055.6bp)

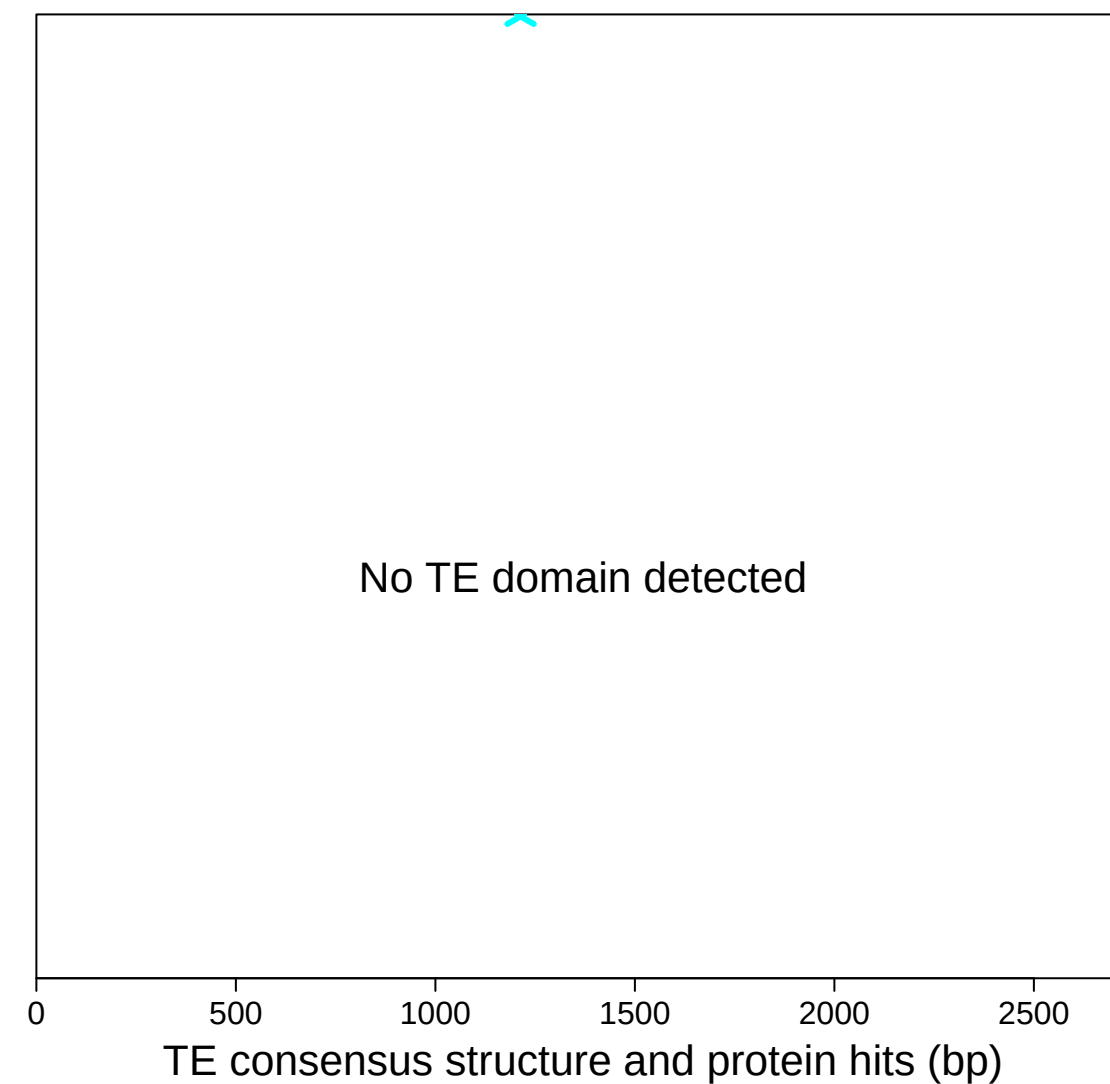
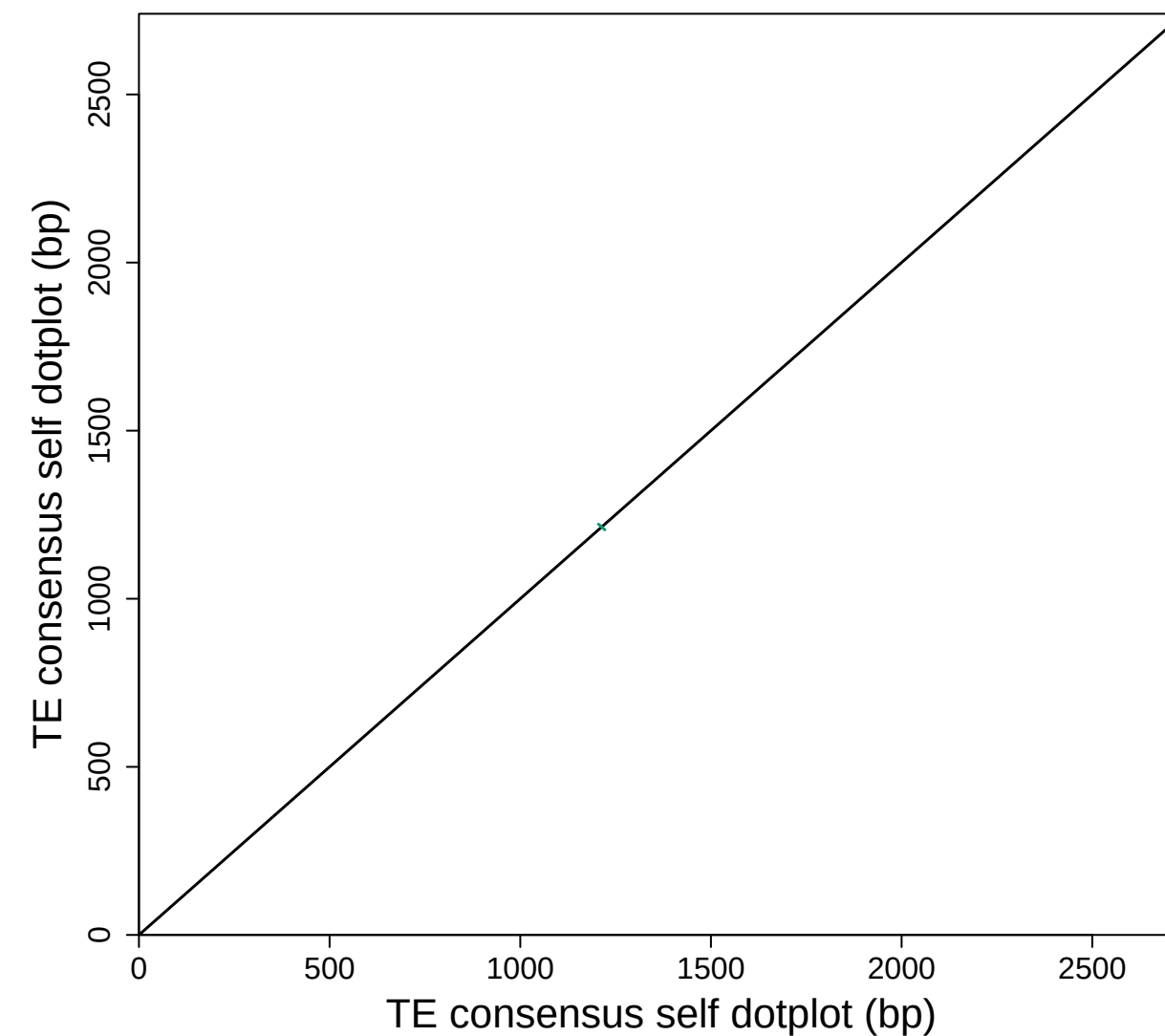
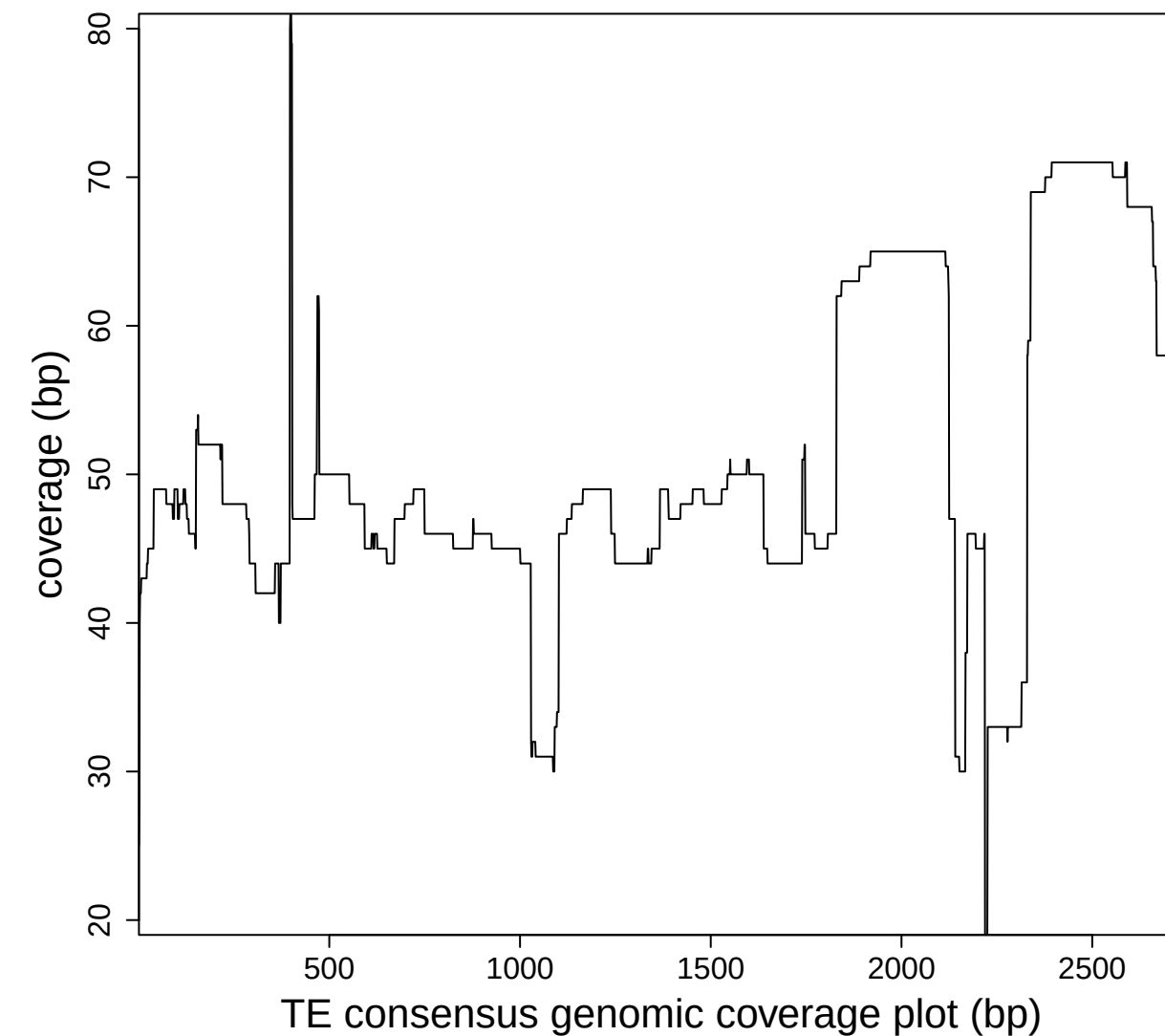
After TEtrimmer Extended plot Blue lines are boundaries



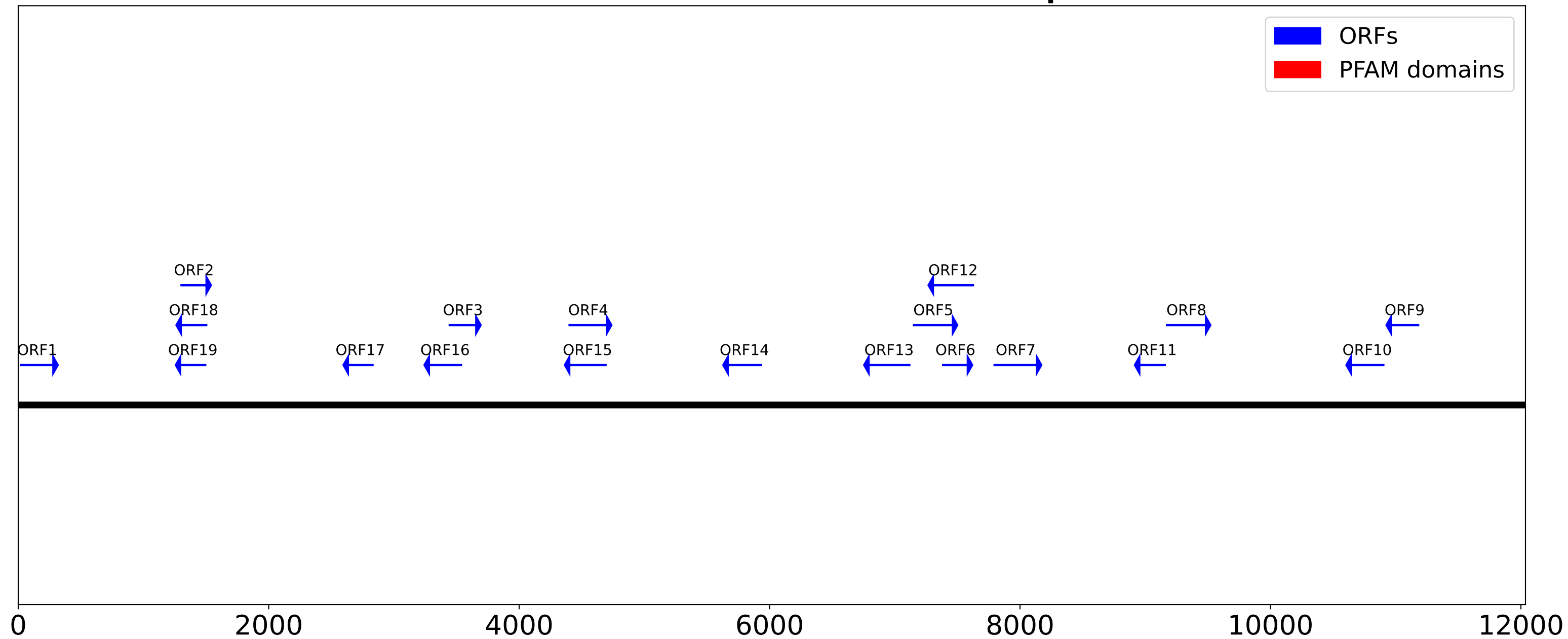
TE: rnd_5_family_752
size: 2740bp; fragments: 261; full length: 5 (≥ 2466 bp)



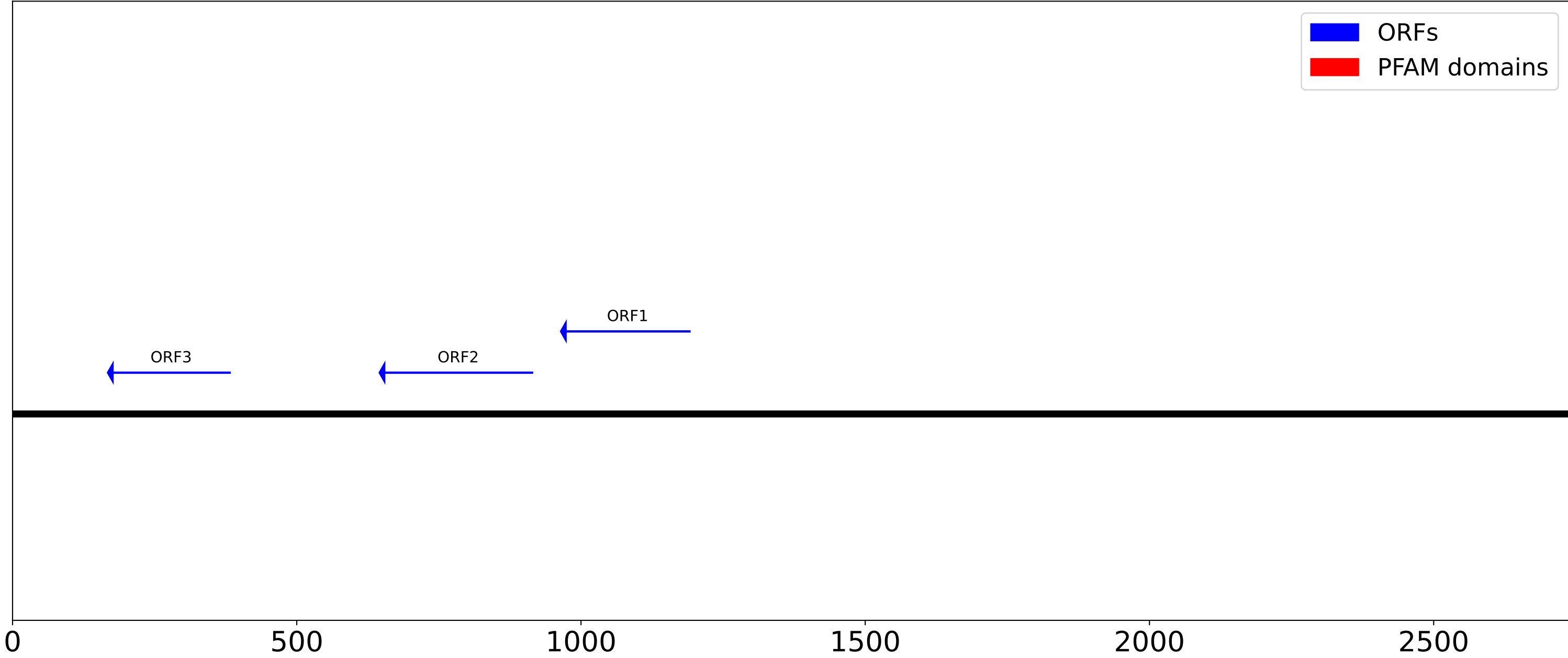
Before TEtrimmer 2740 bp



After TEtrimmer ORF and PFAM domain plot



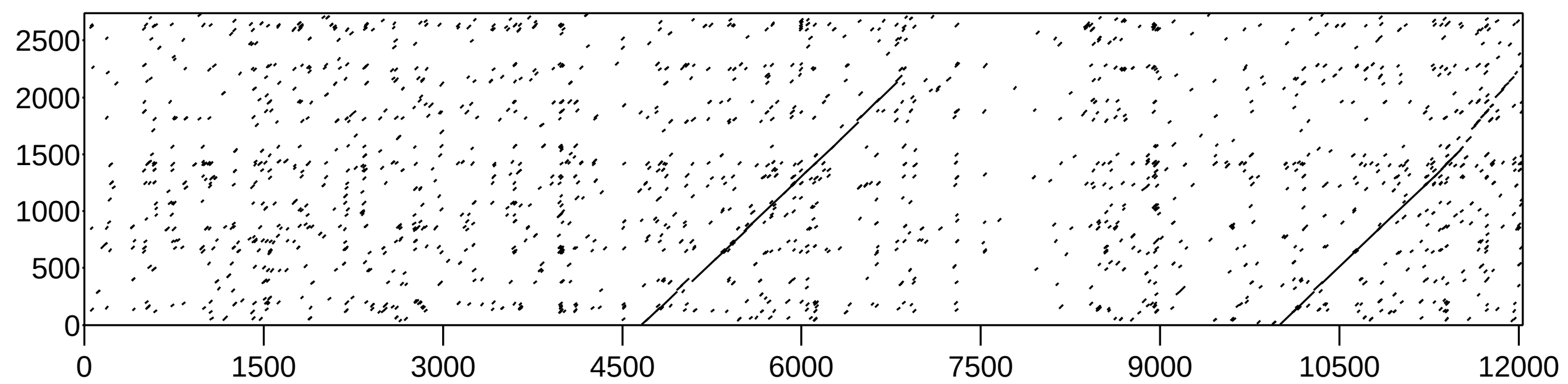
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)