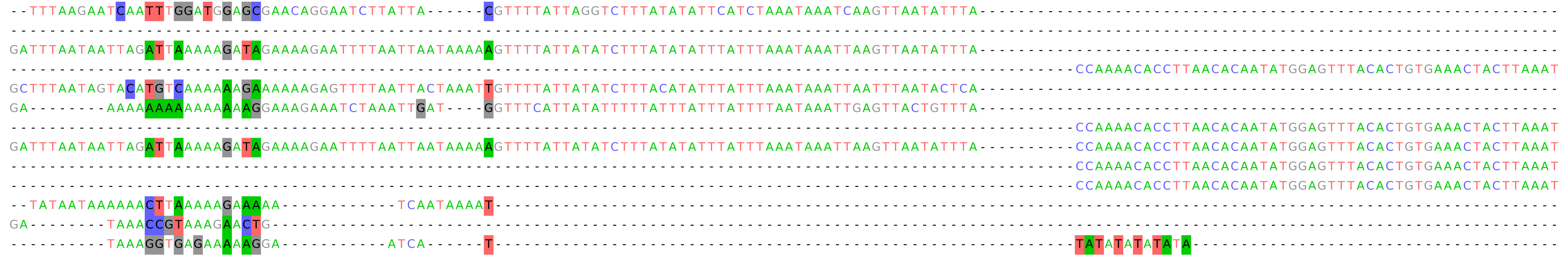
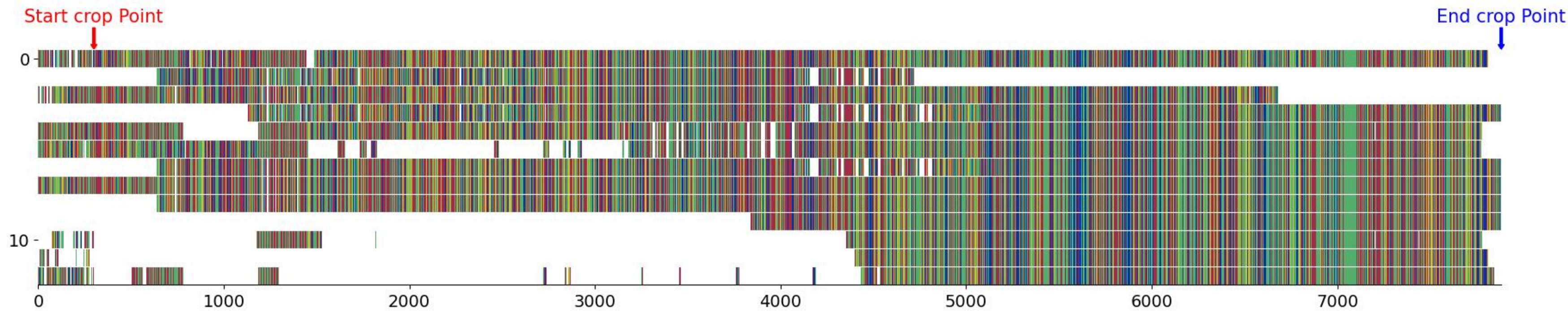


Start crop Point

End crop Point

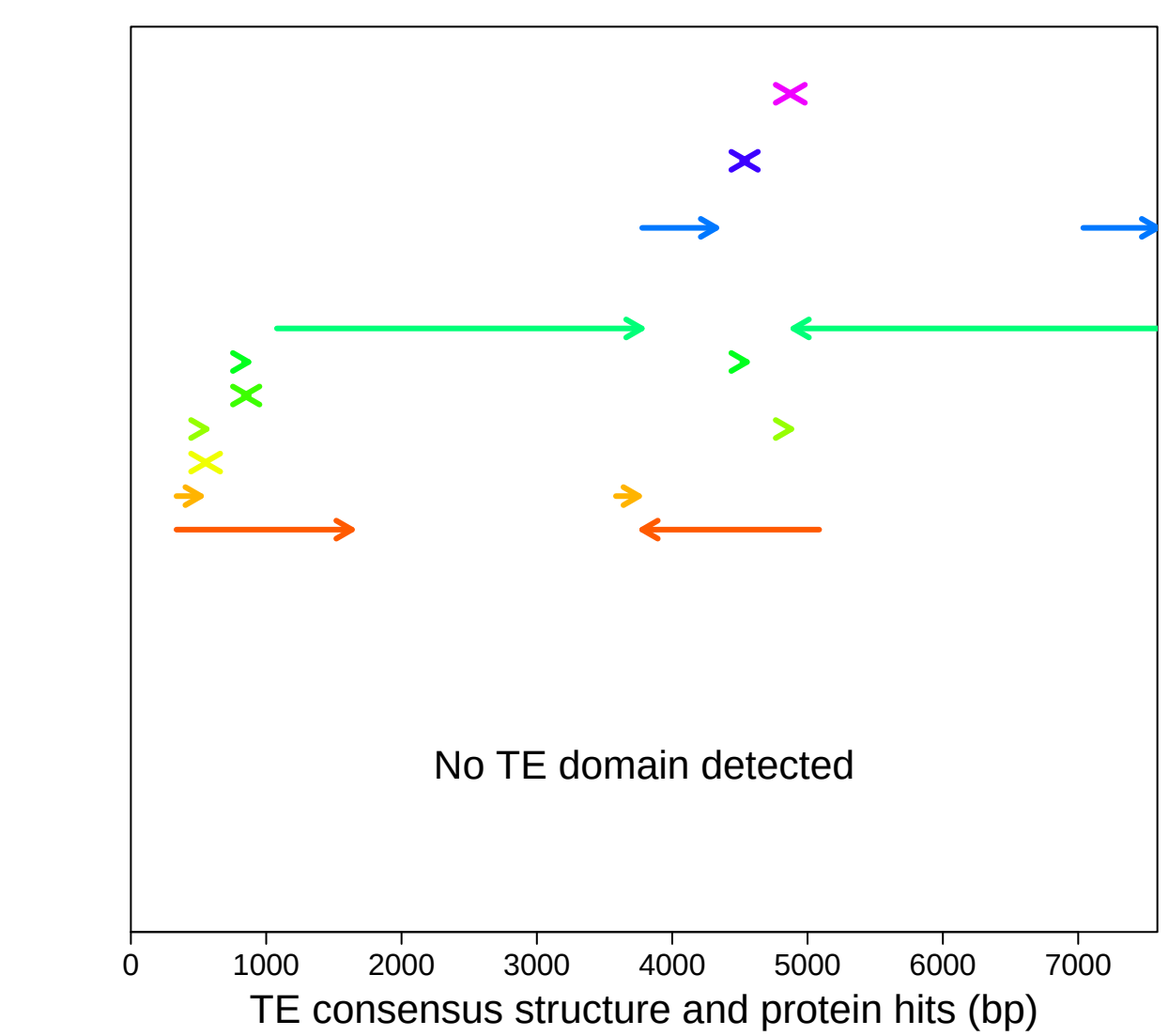
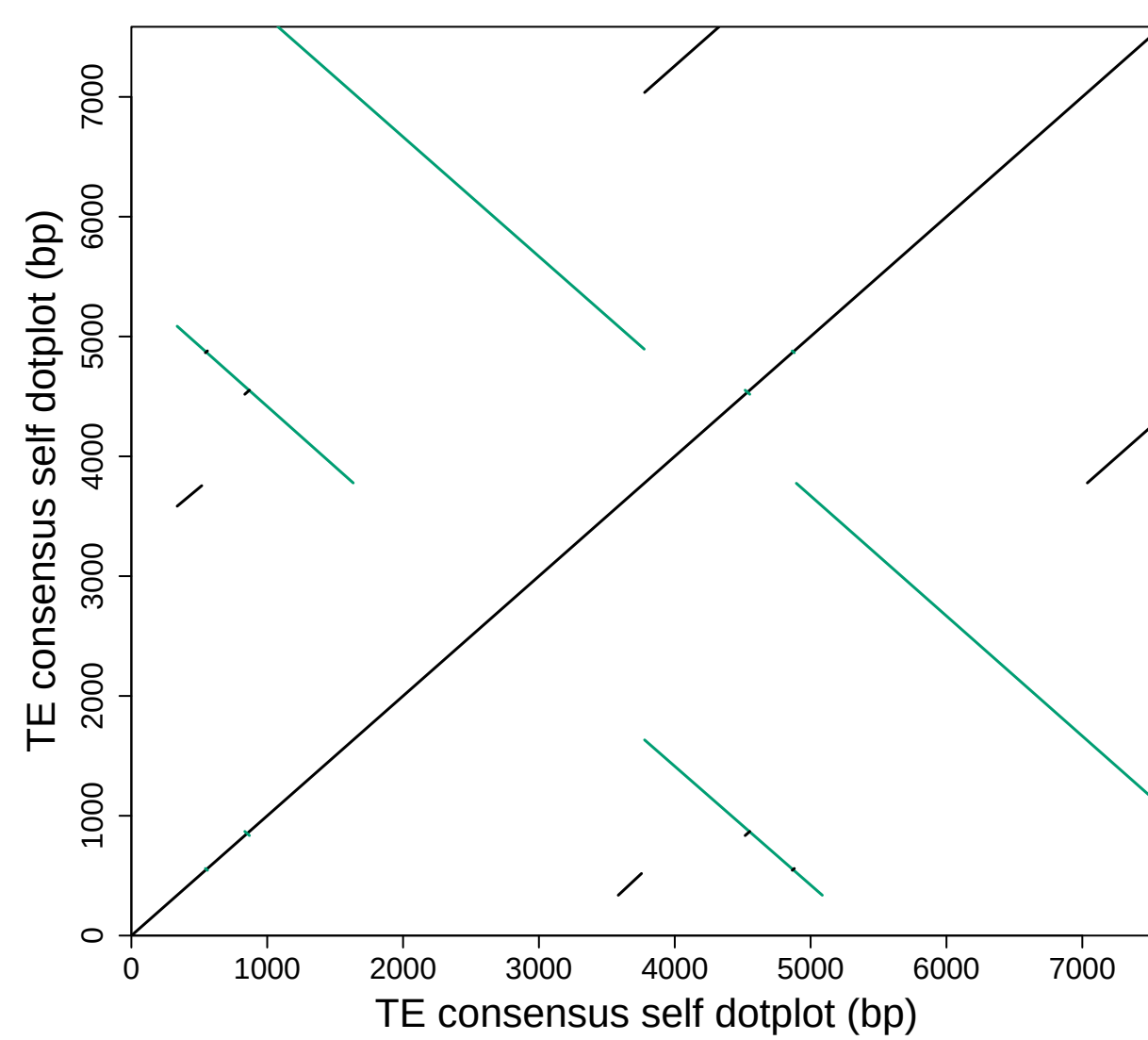
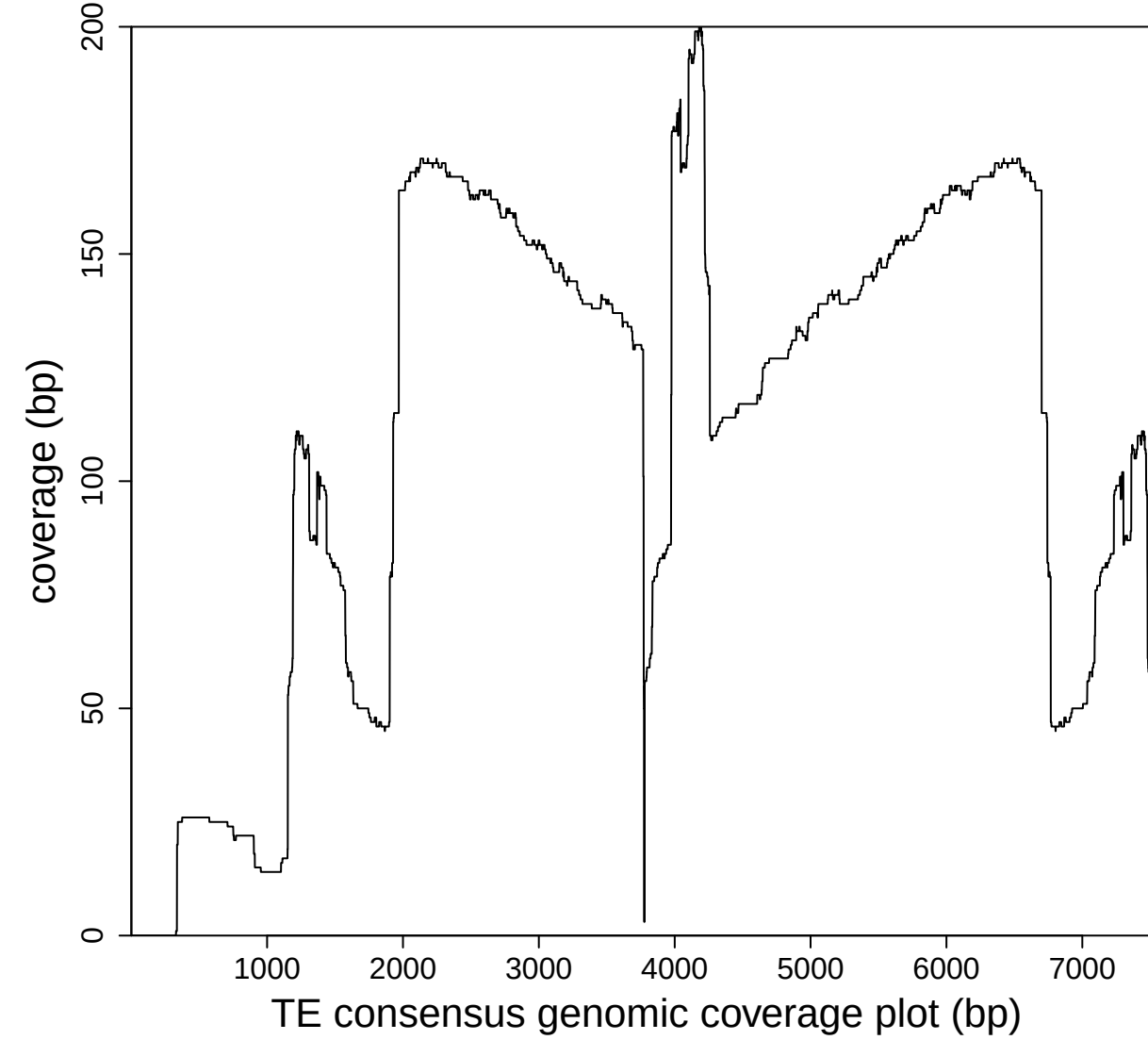
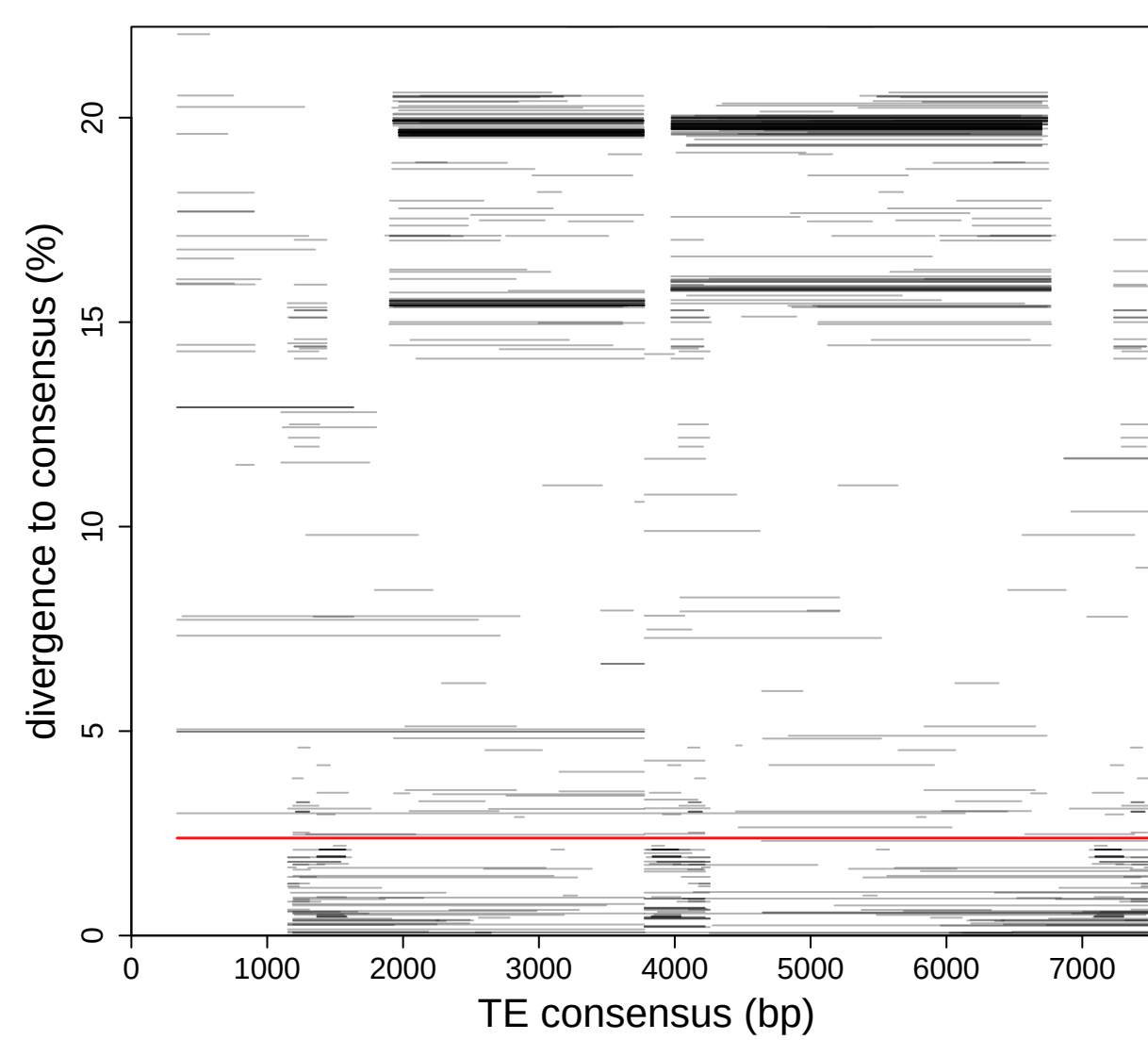
MSA length = 7585





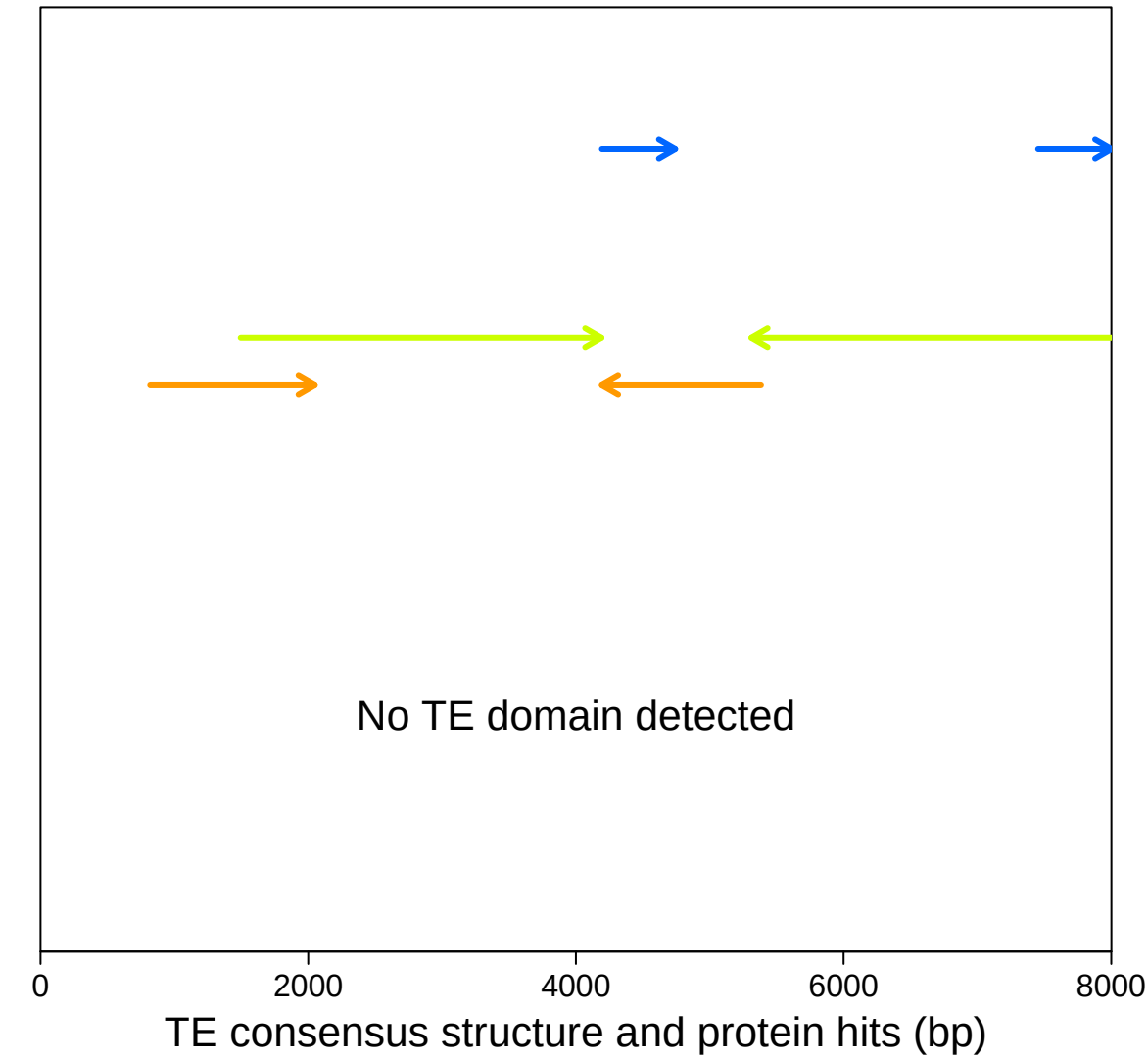
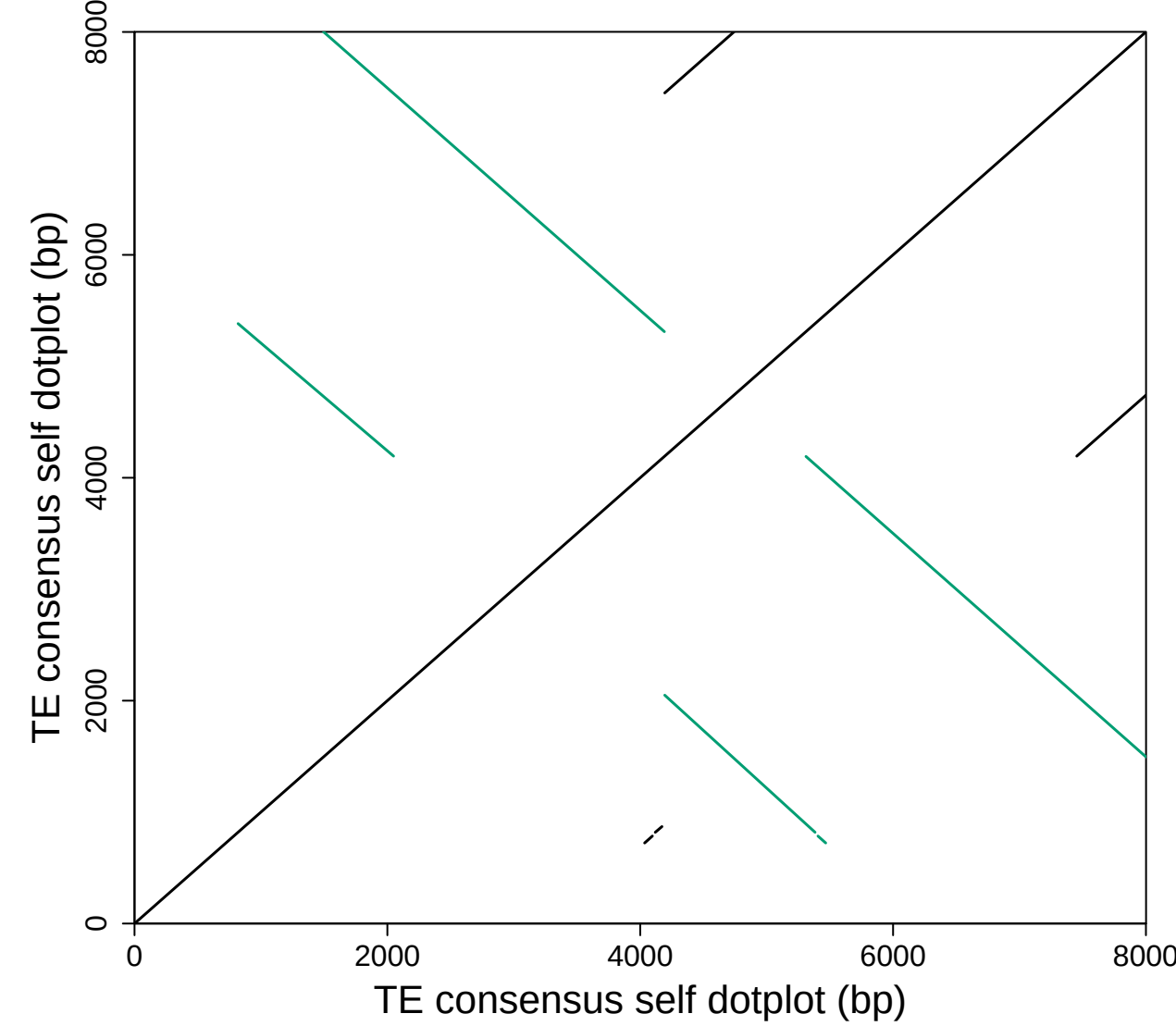
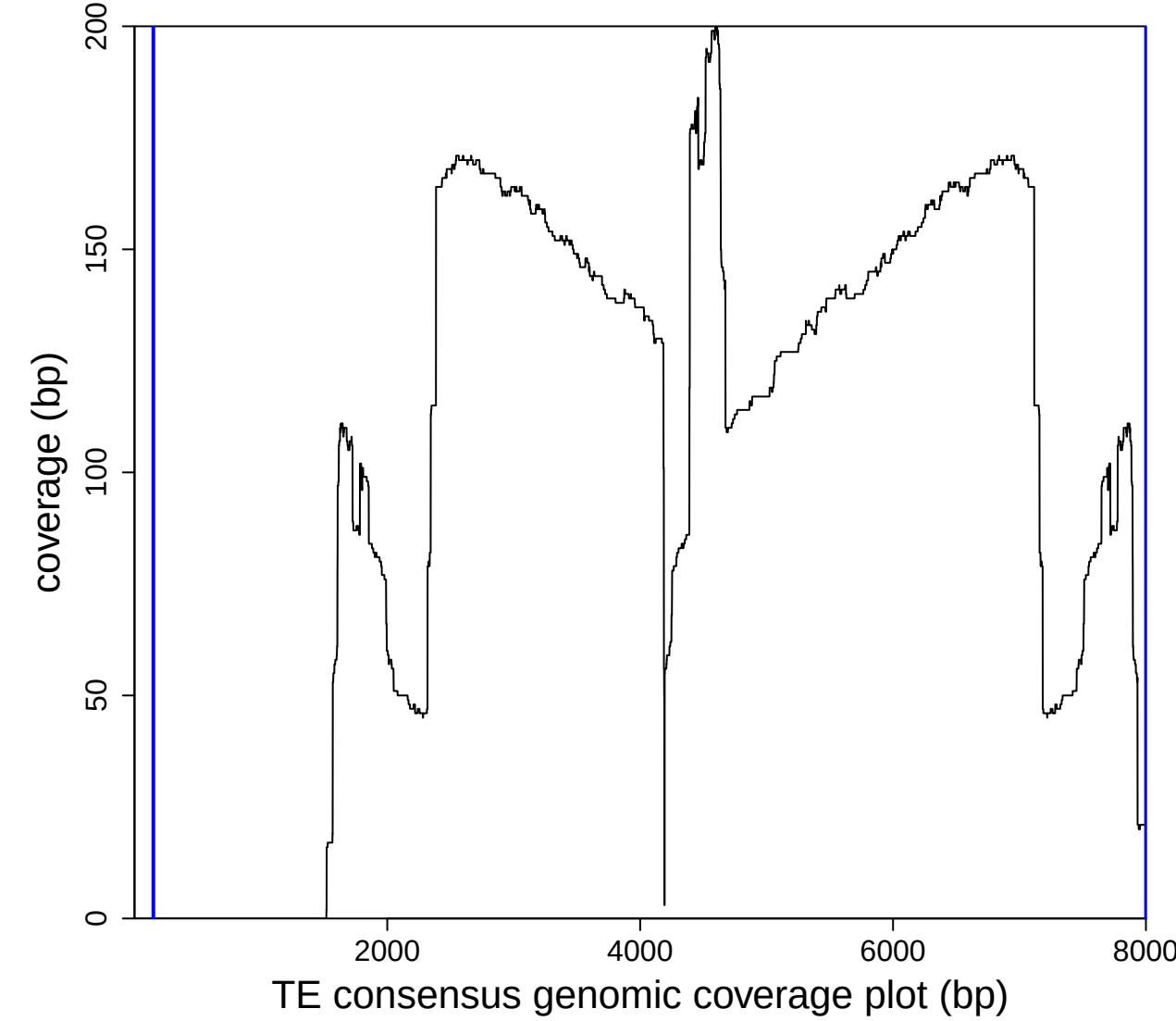
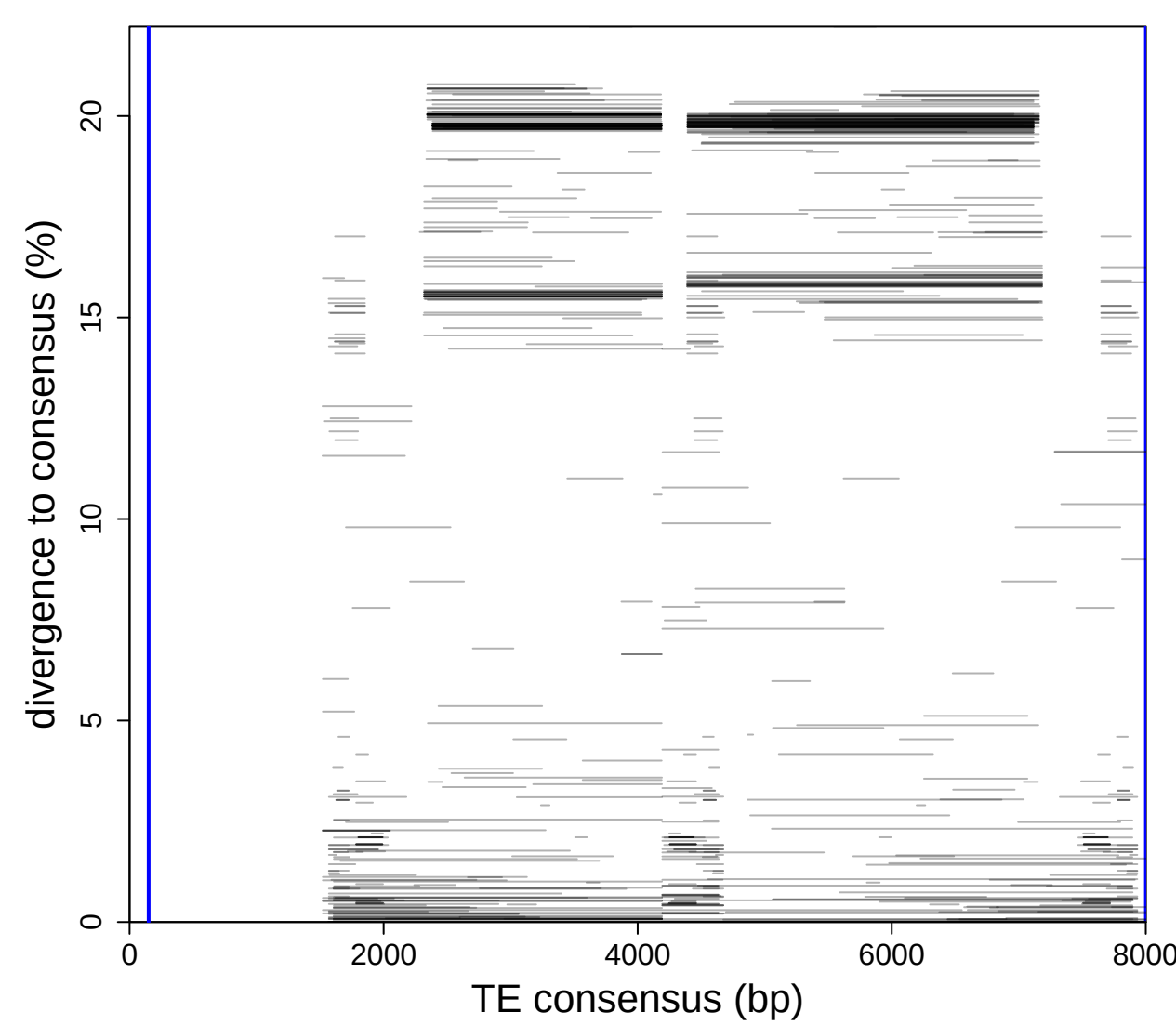
bed_uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa c
size: 7585bp; fragments: 763; full length: 1 (>=6826.5bp)

After TEtrimmer 7585 bp

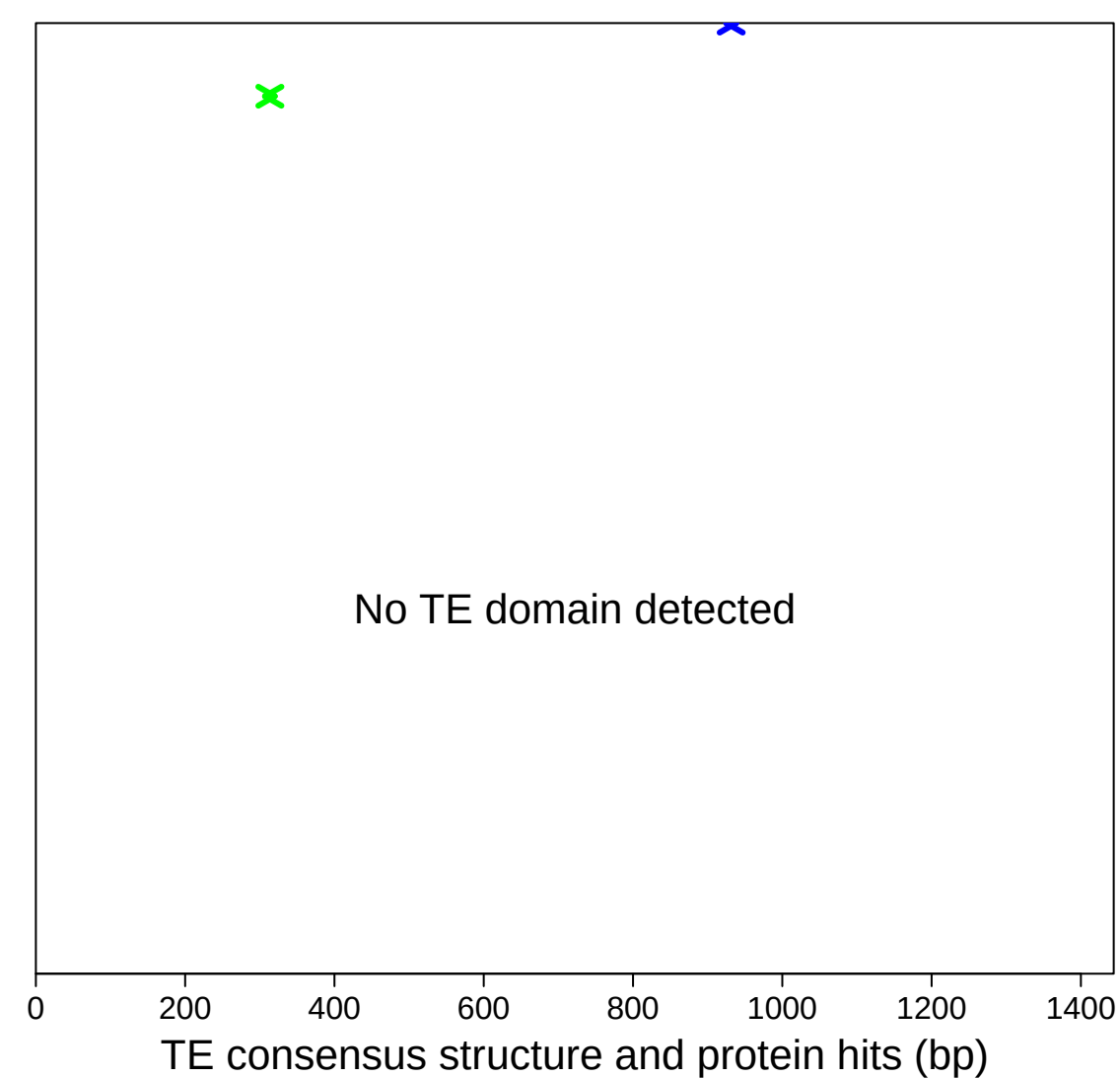
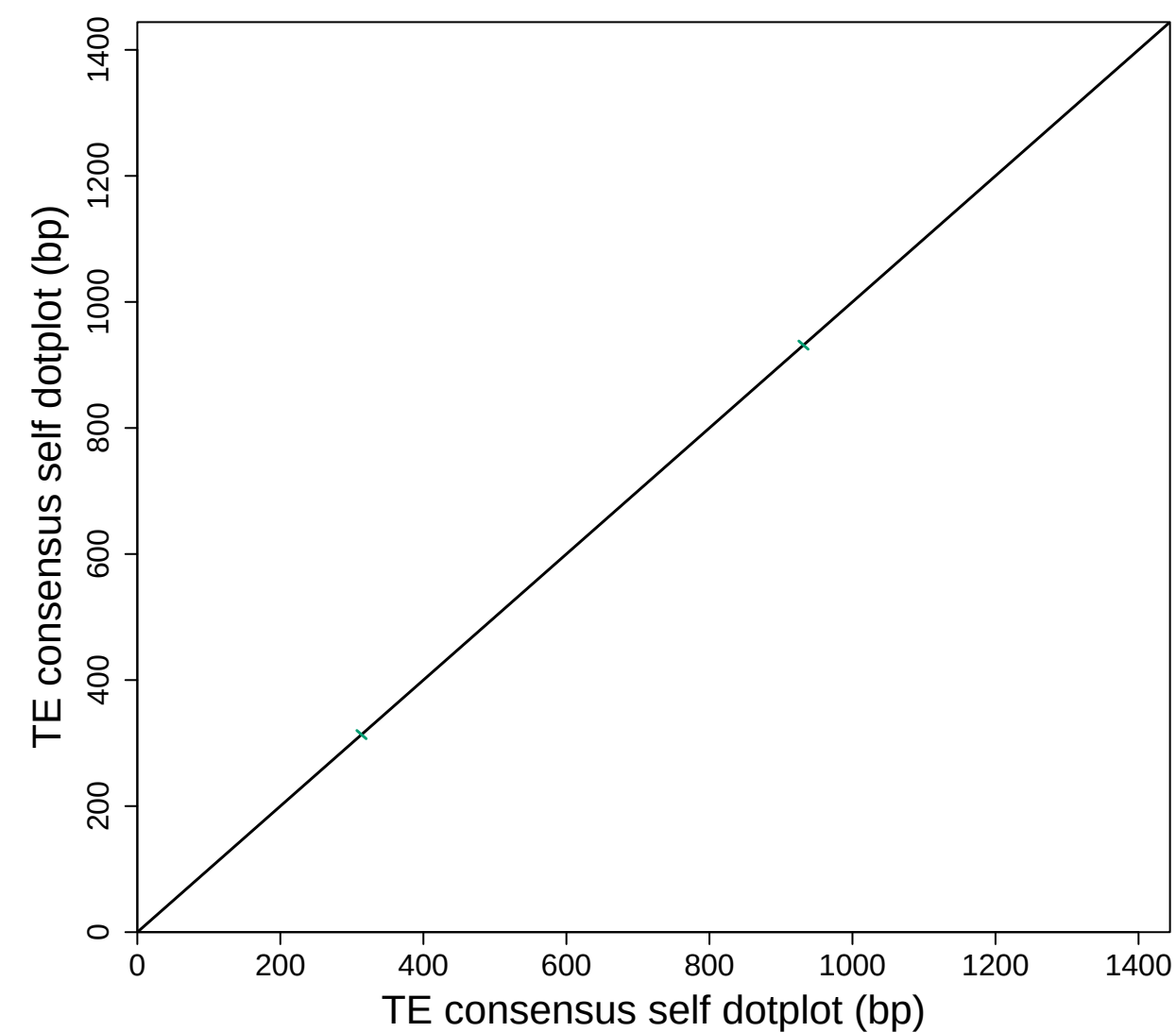
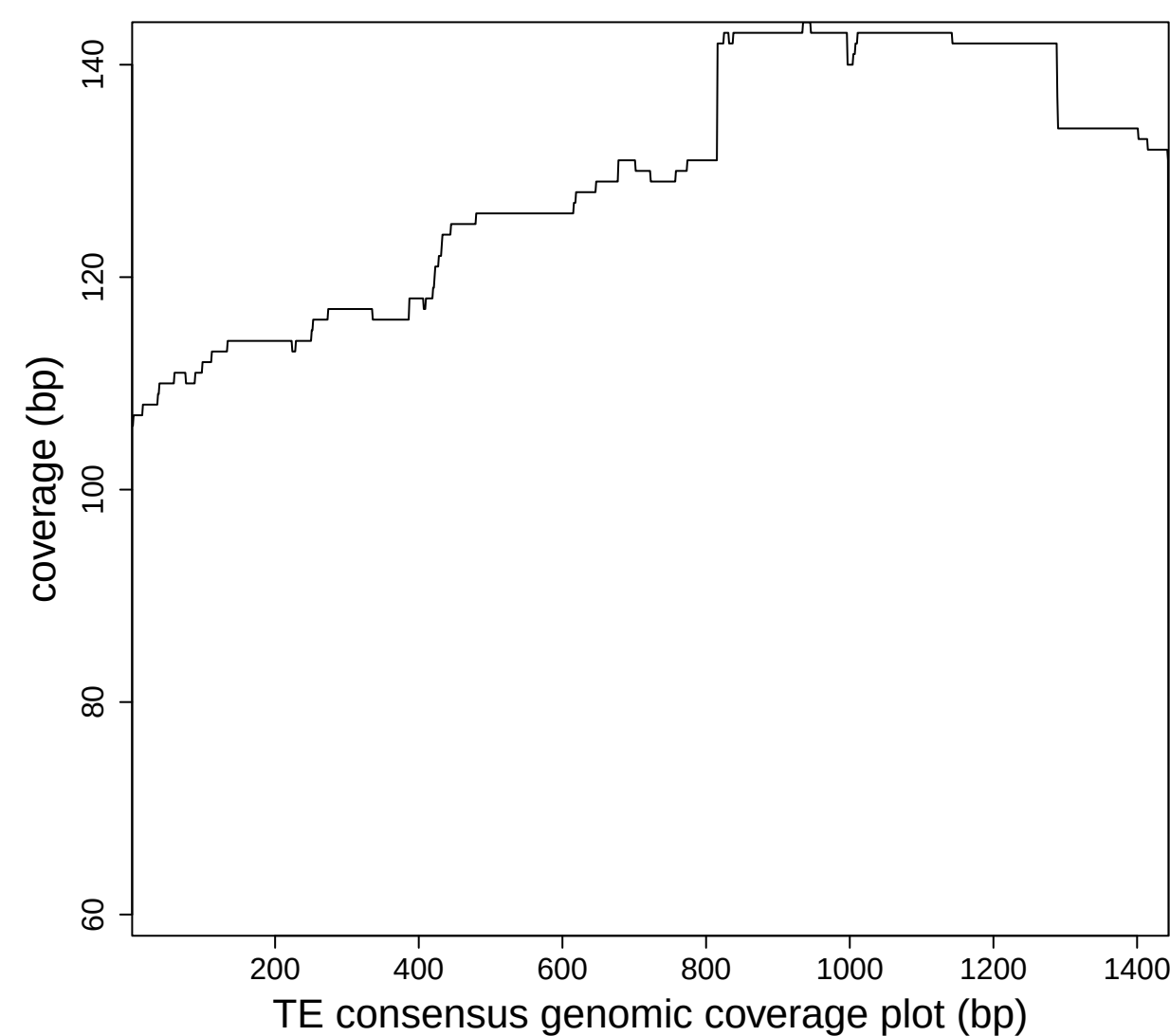
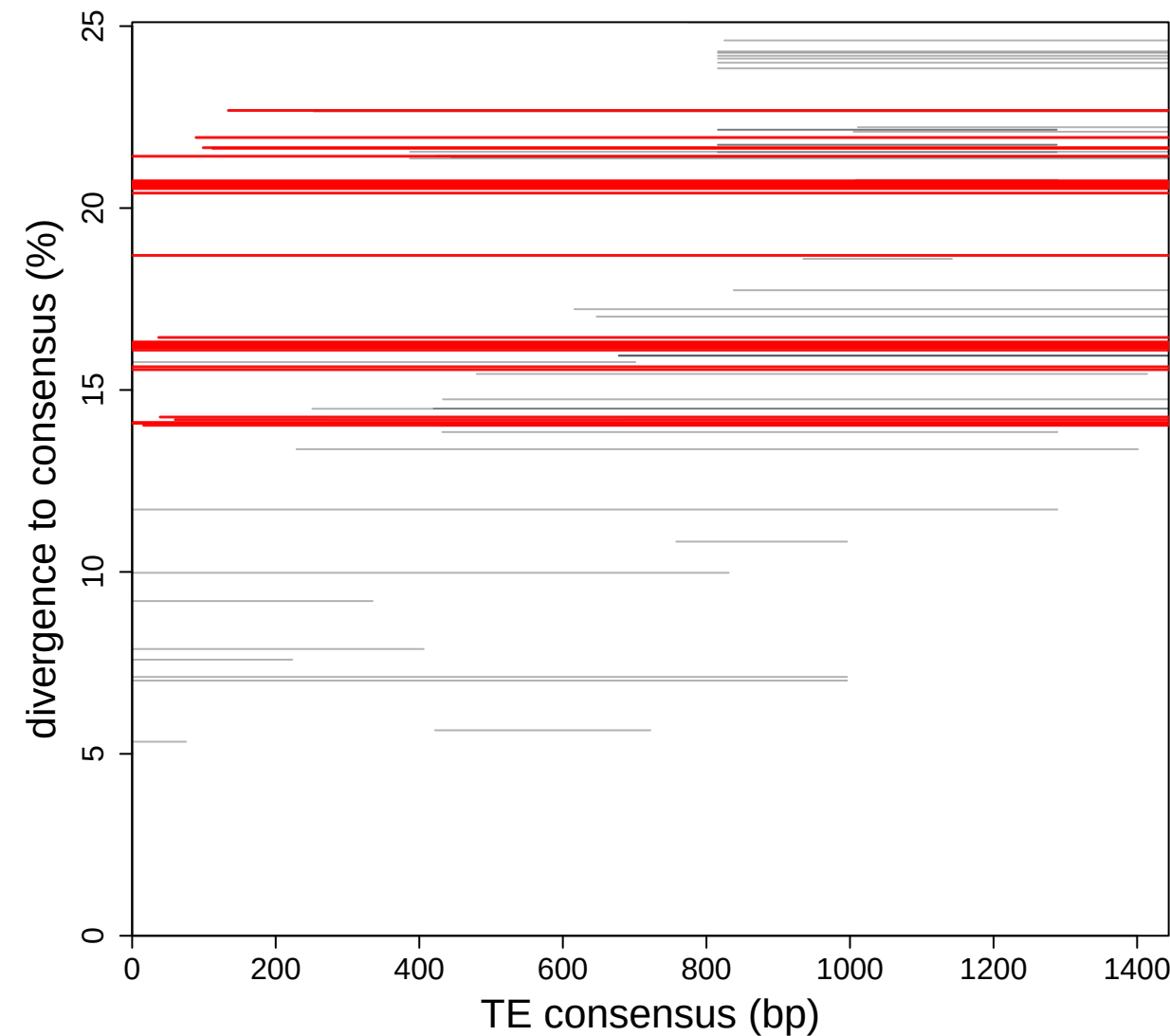


5.fasta.b.bed uf.bed g_3.bed fm_1.bed 0 0 bcIn.fa aln.fa cl.f
size: 8001bp; fragments: 750; full length: 0 (>=7200.9bp)

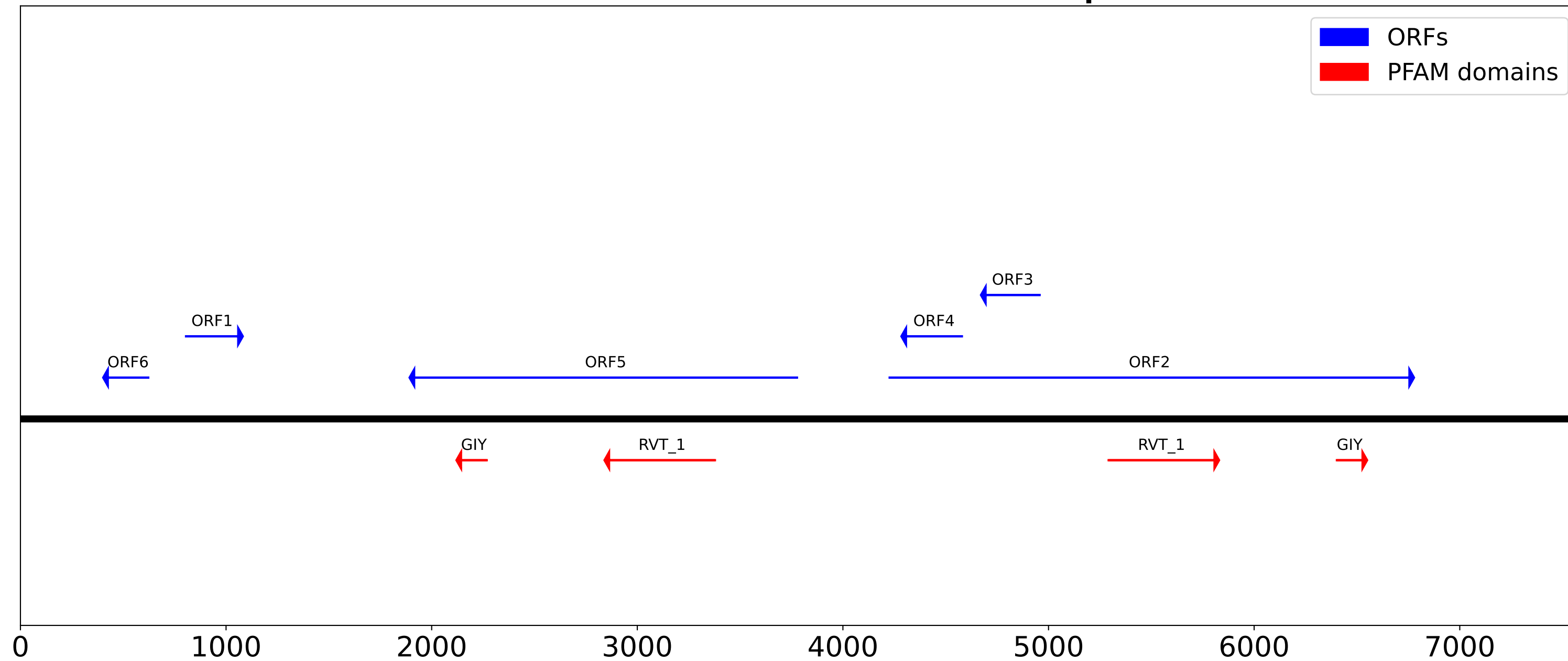
After TEtrimmer Extended plot Blue lines are boundaries



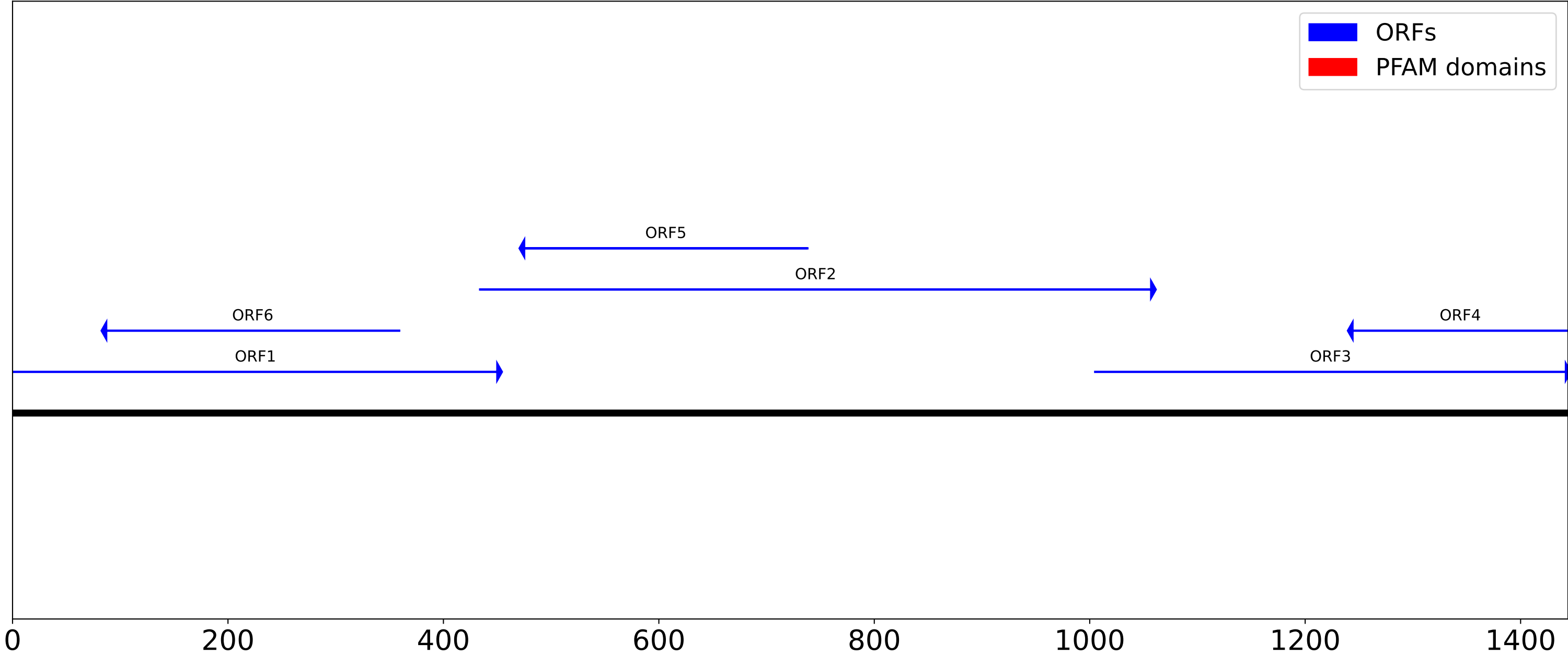
TE: rnd_5_family_3825
size: 1444bp; fragments: 155; full length: 106 (≥ 1299.6 bp)



After TEtrimmer ORF and PFAM domain plot



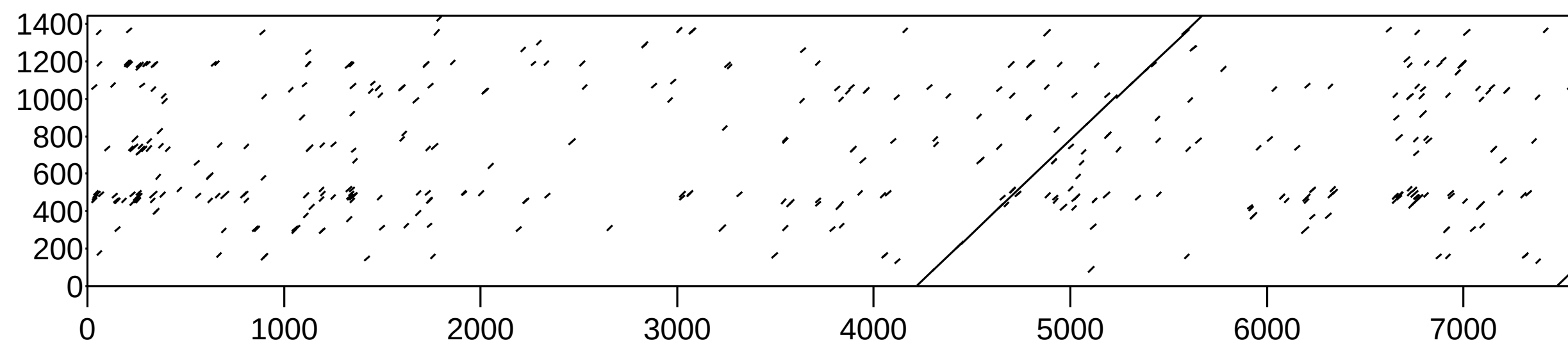
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)