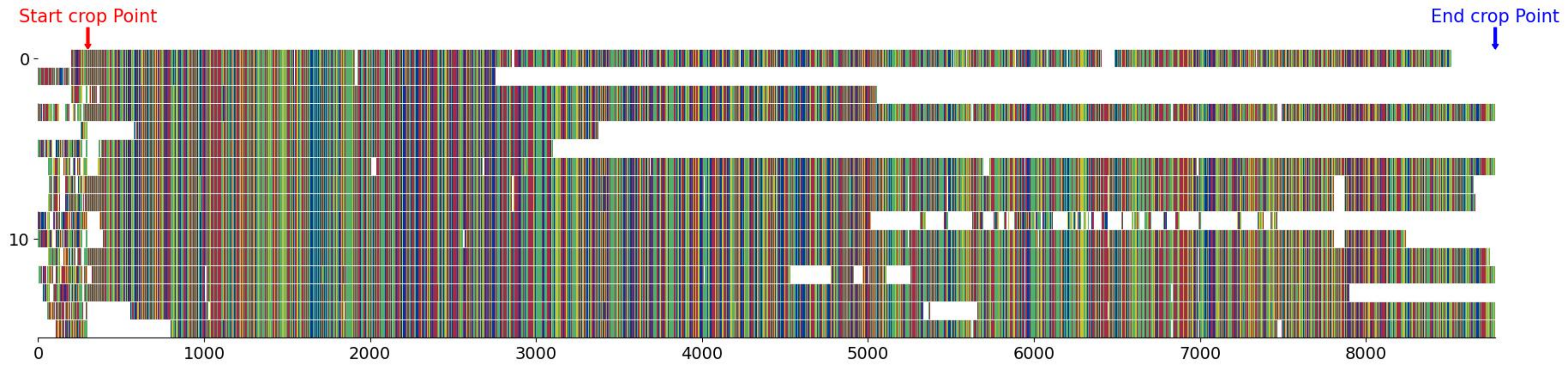




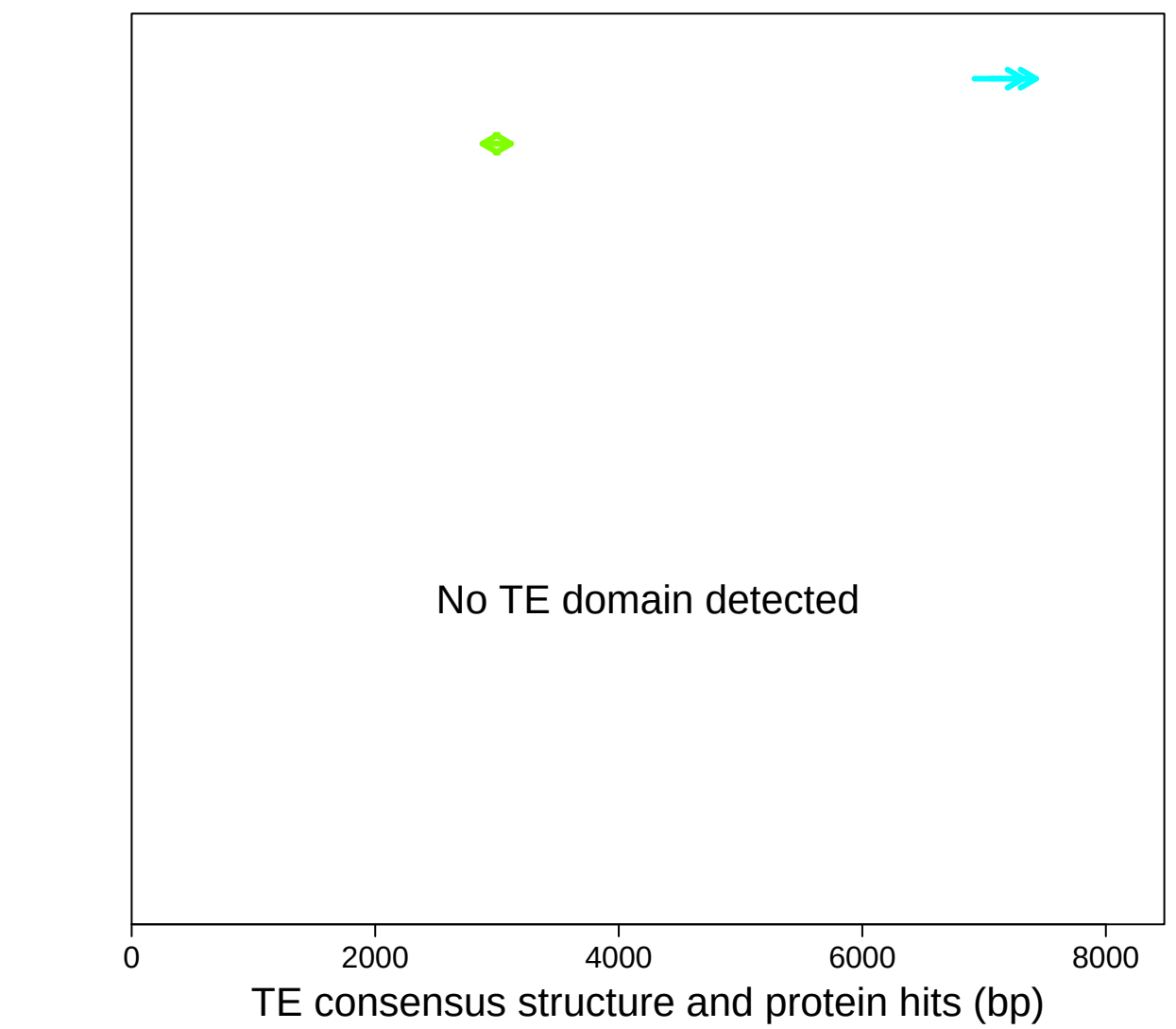
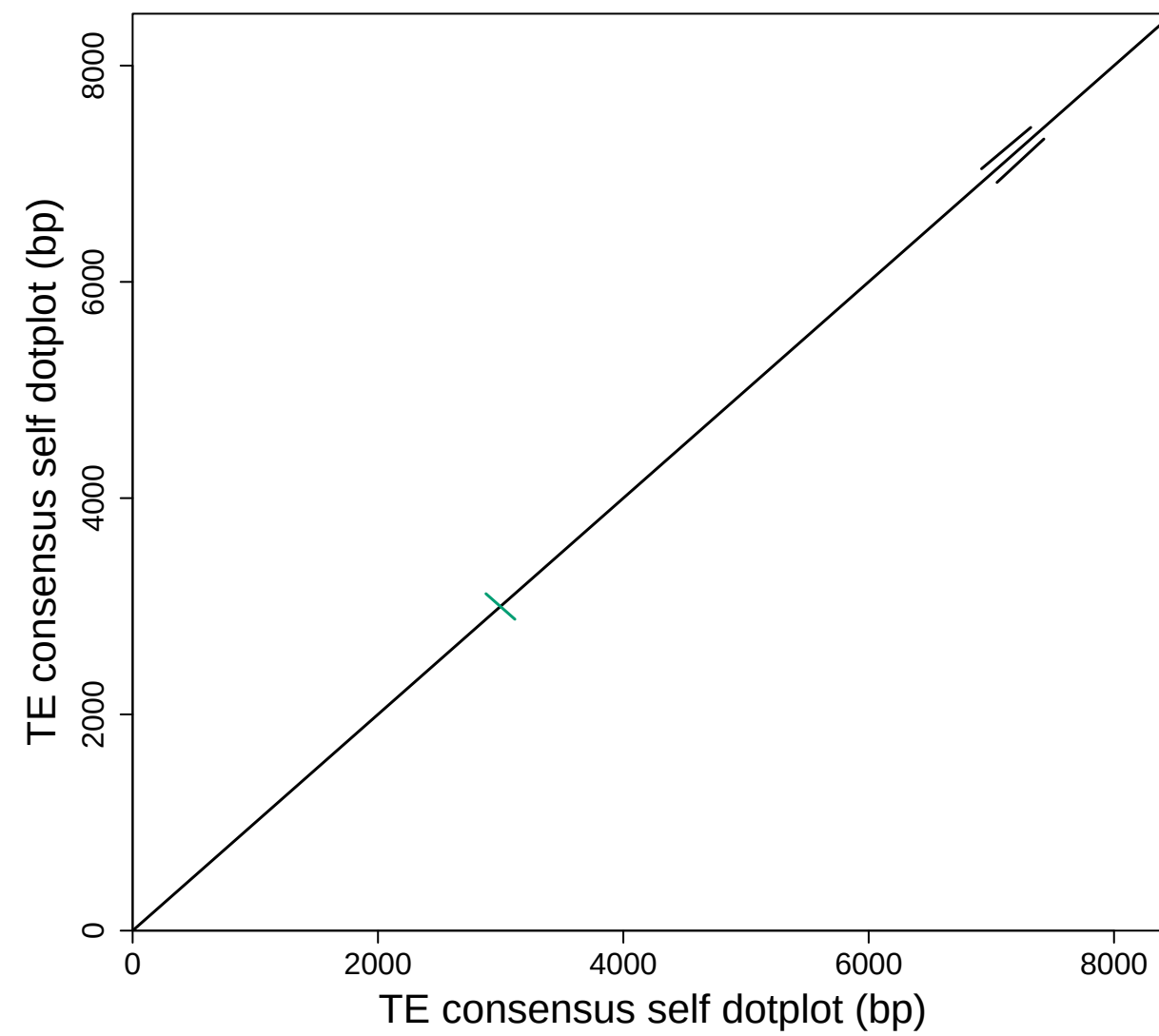
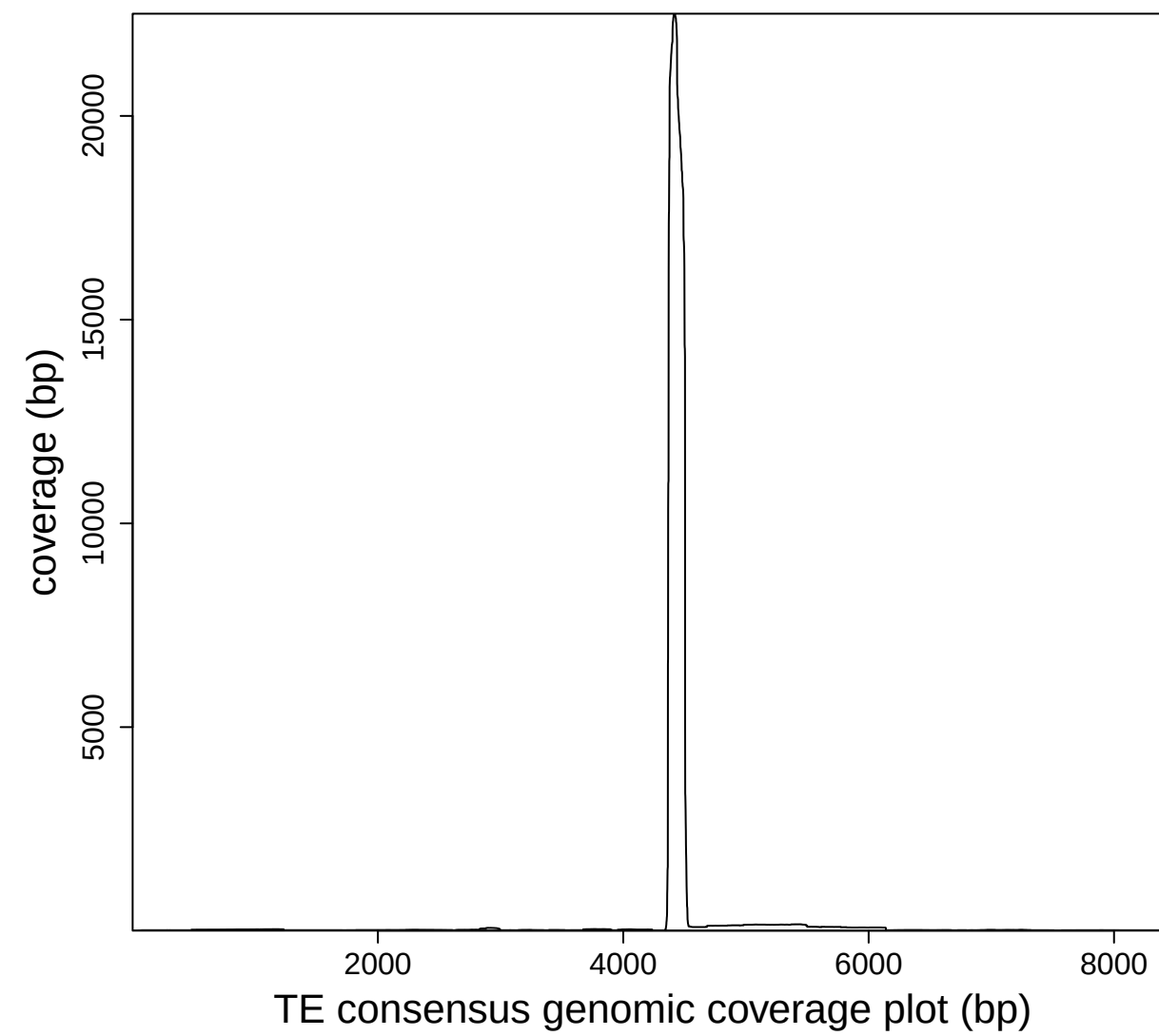
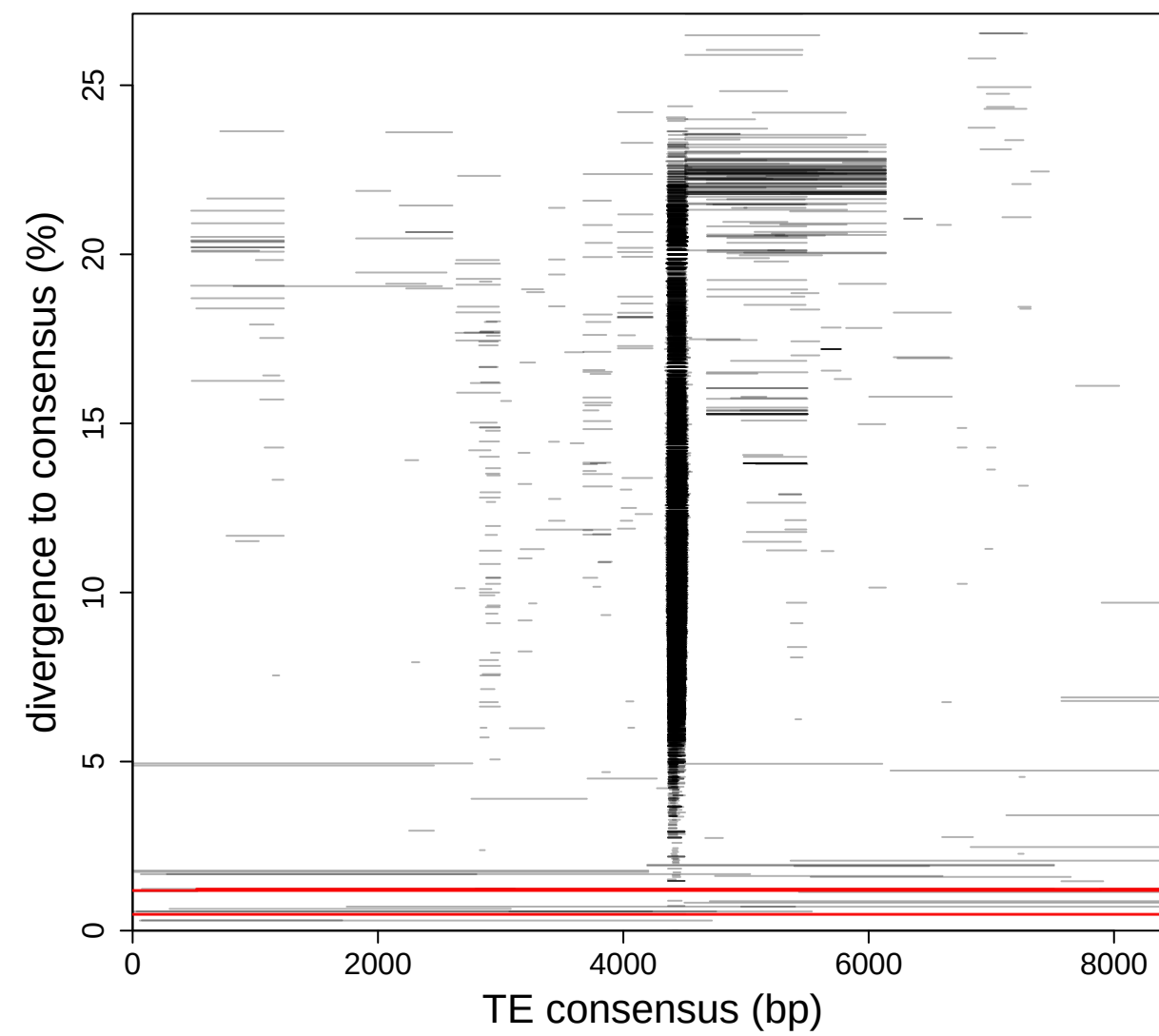
MSA length = 8480

[illegible]



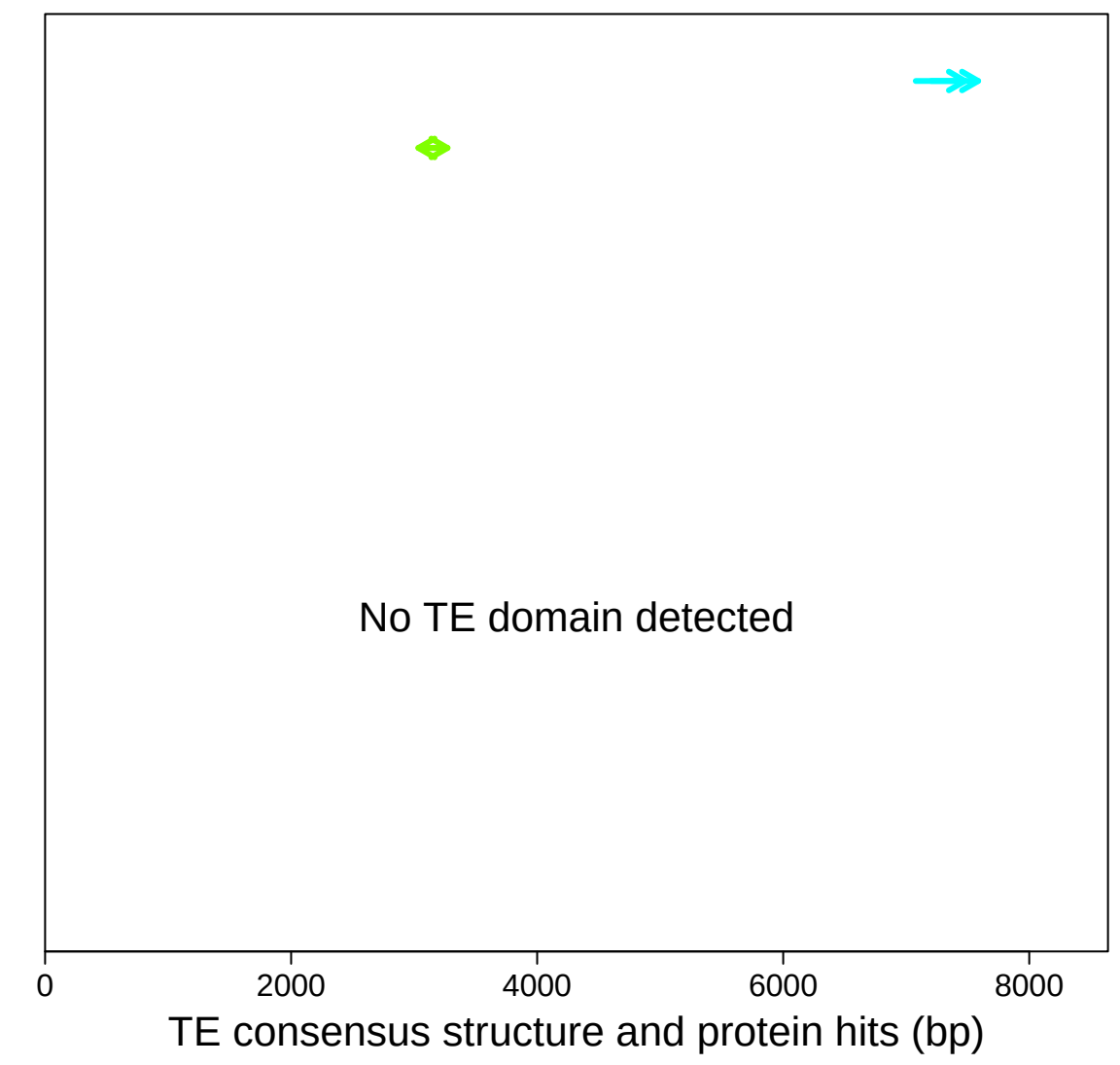
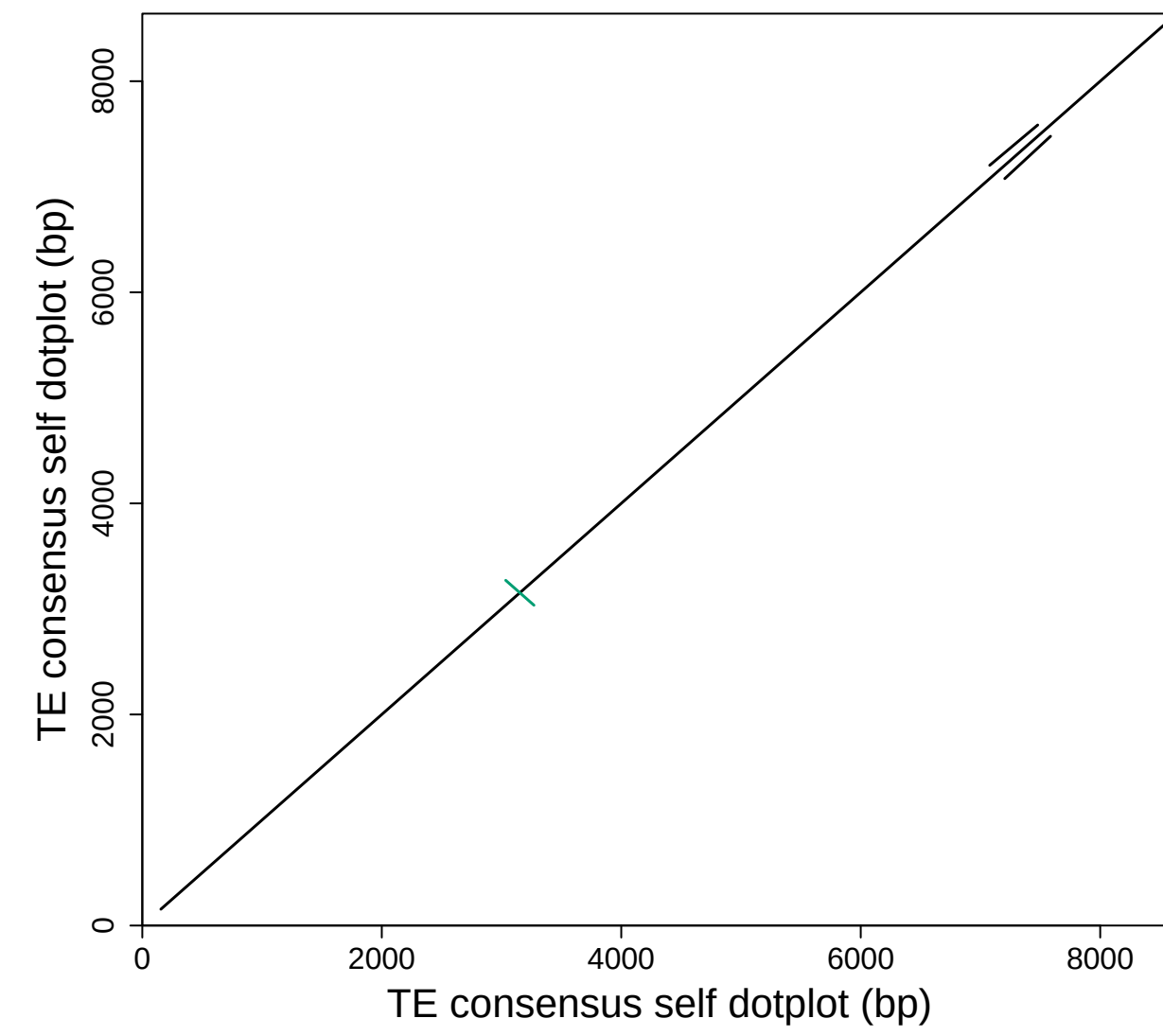
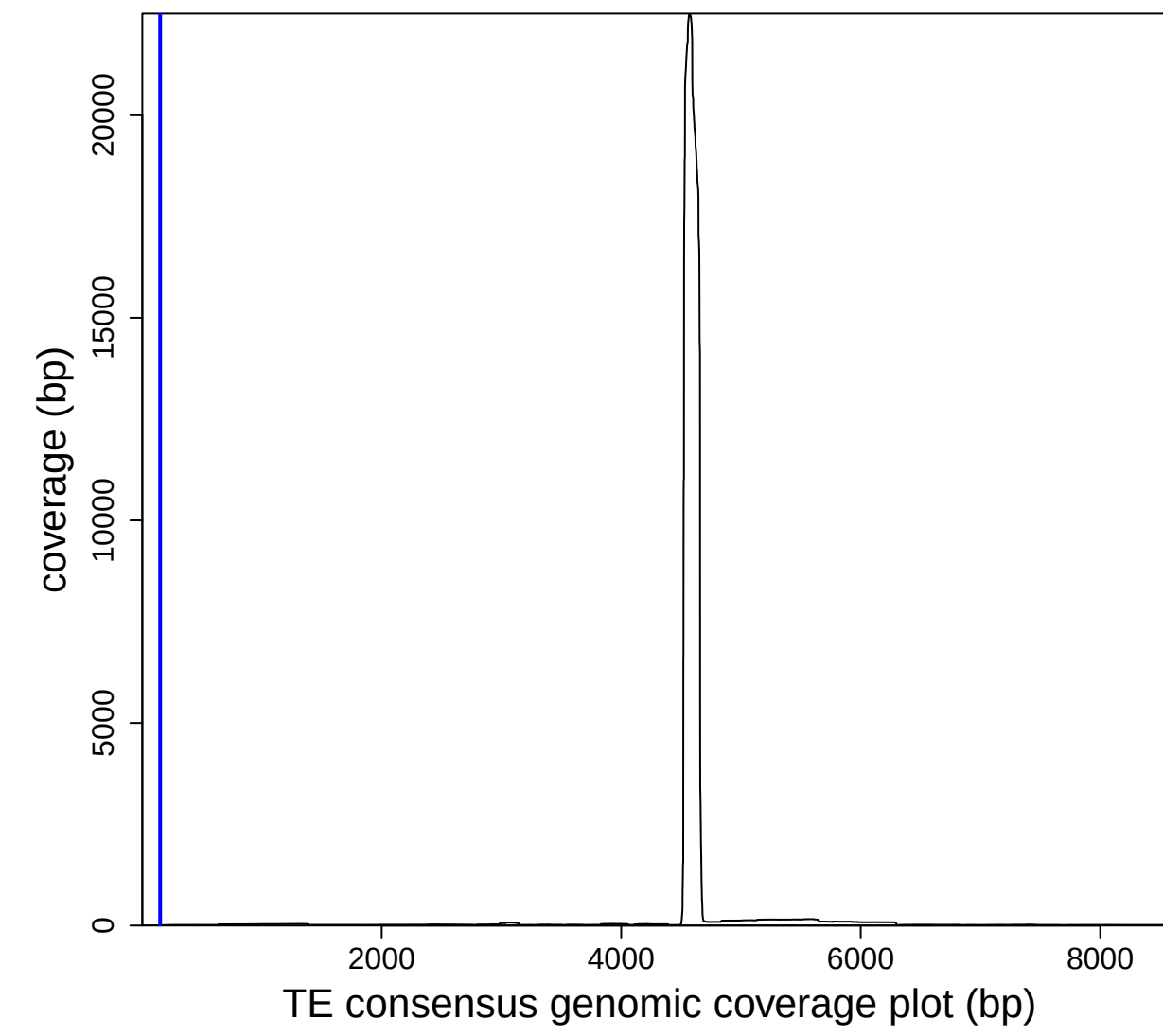
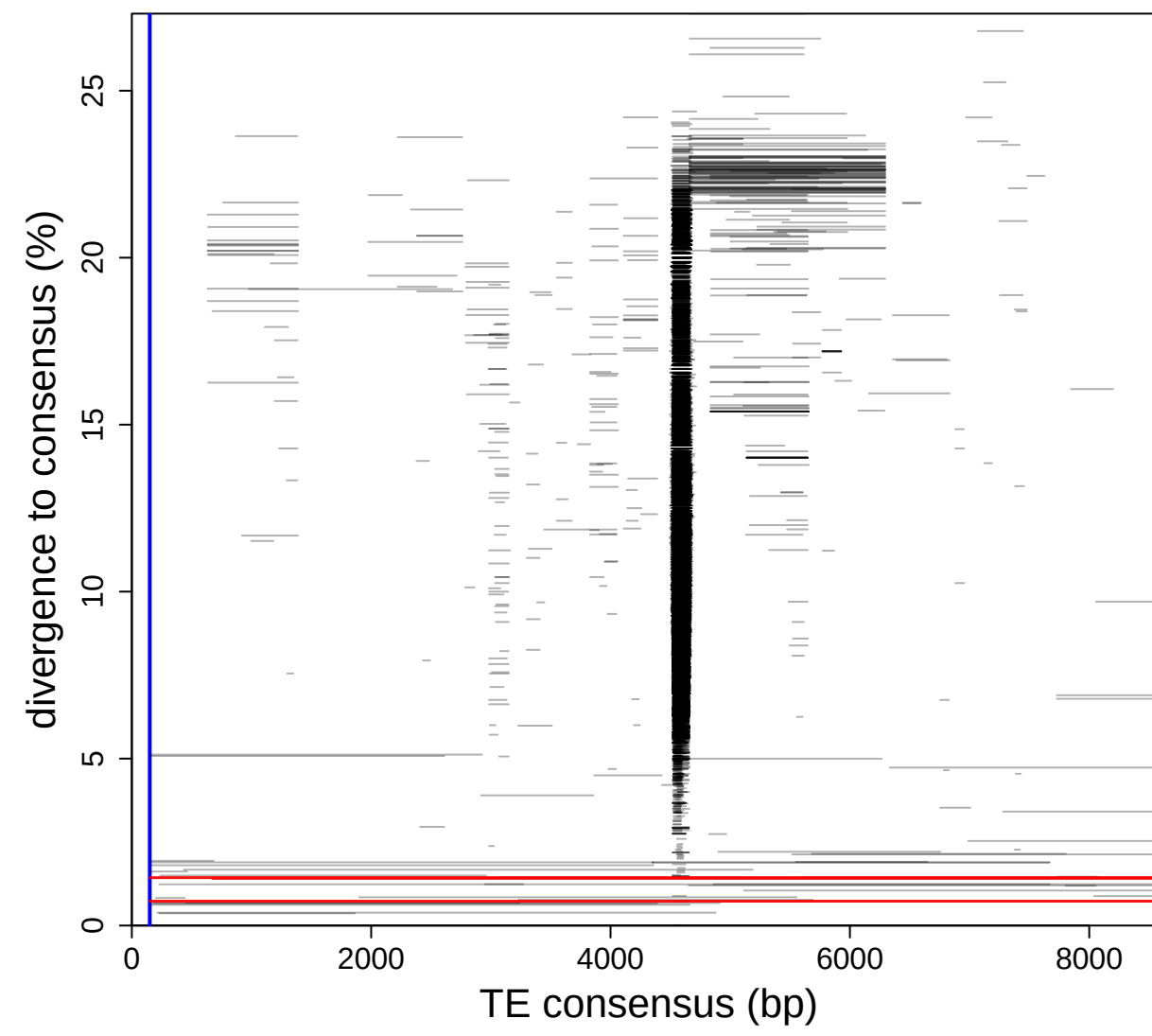
ta.b.bed uf.bed g_3.bed fm_1.bed 0_0 bcln.fa aln.fa cl.fa gs.f
size: 8480bp; fragments: 22984; full length: 3 (>=7632bp)

After TEtrimmer 8480 bp

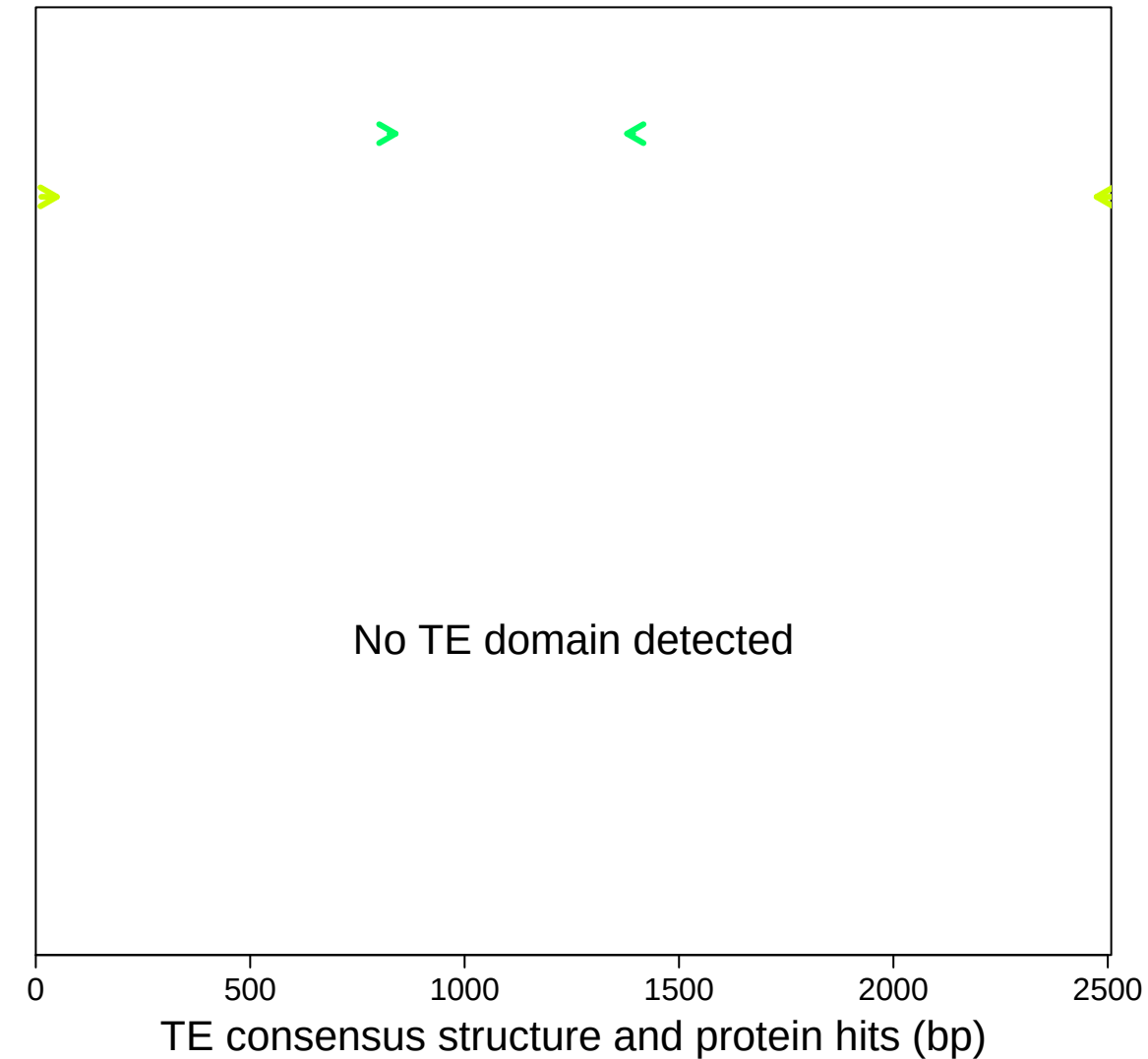
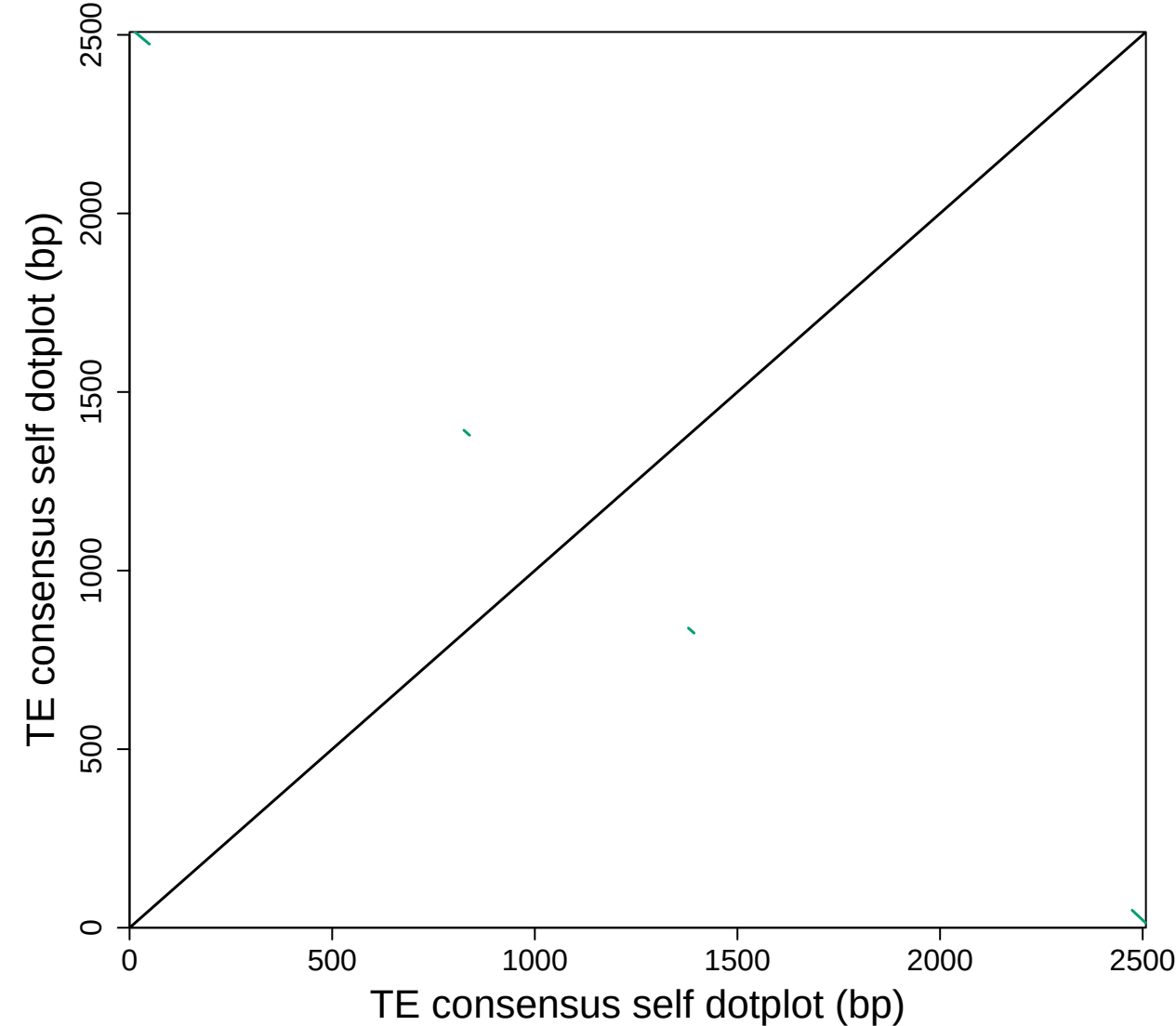
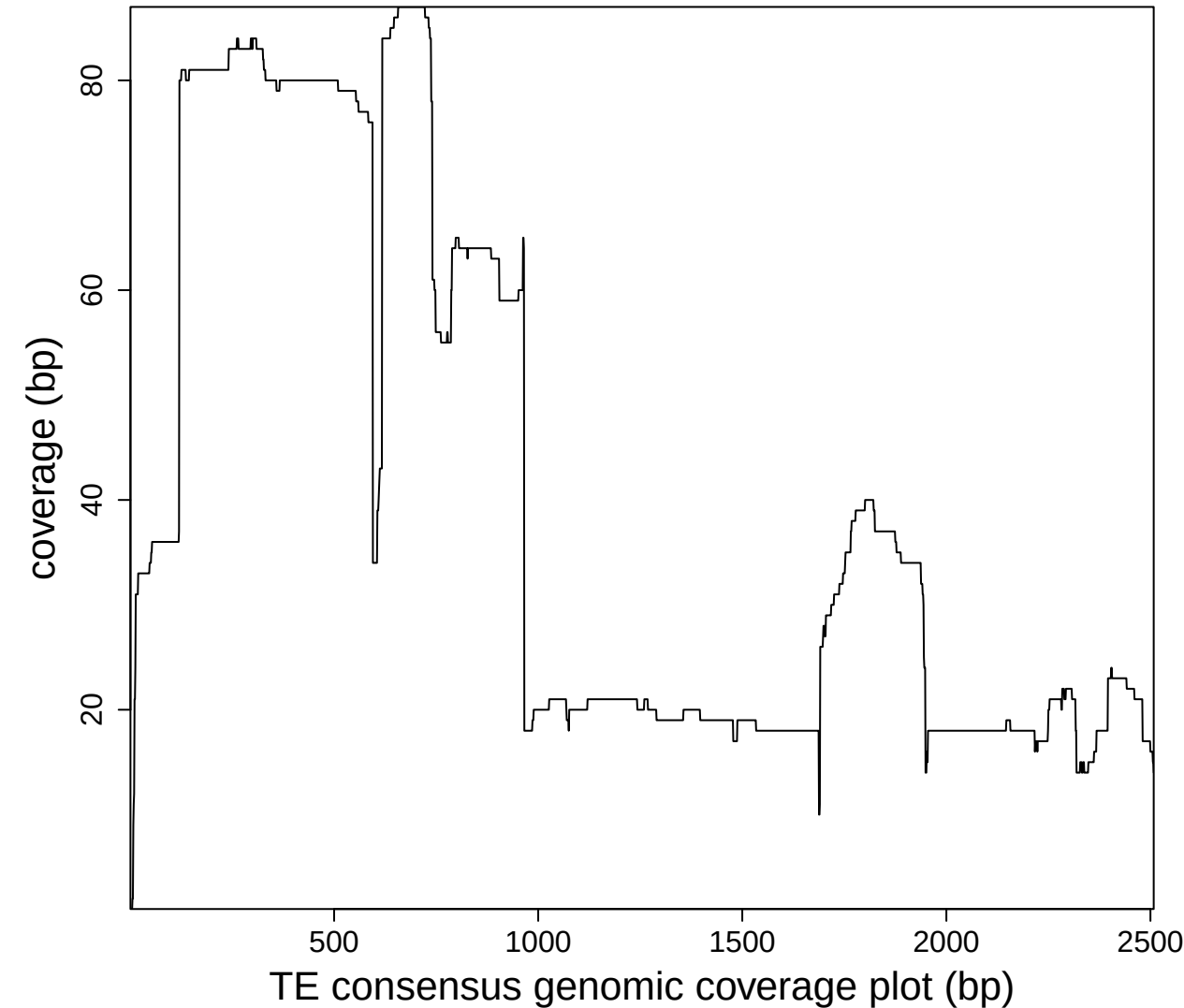
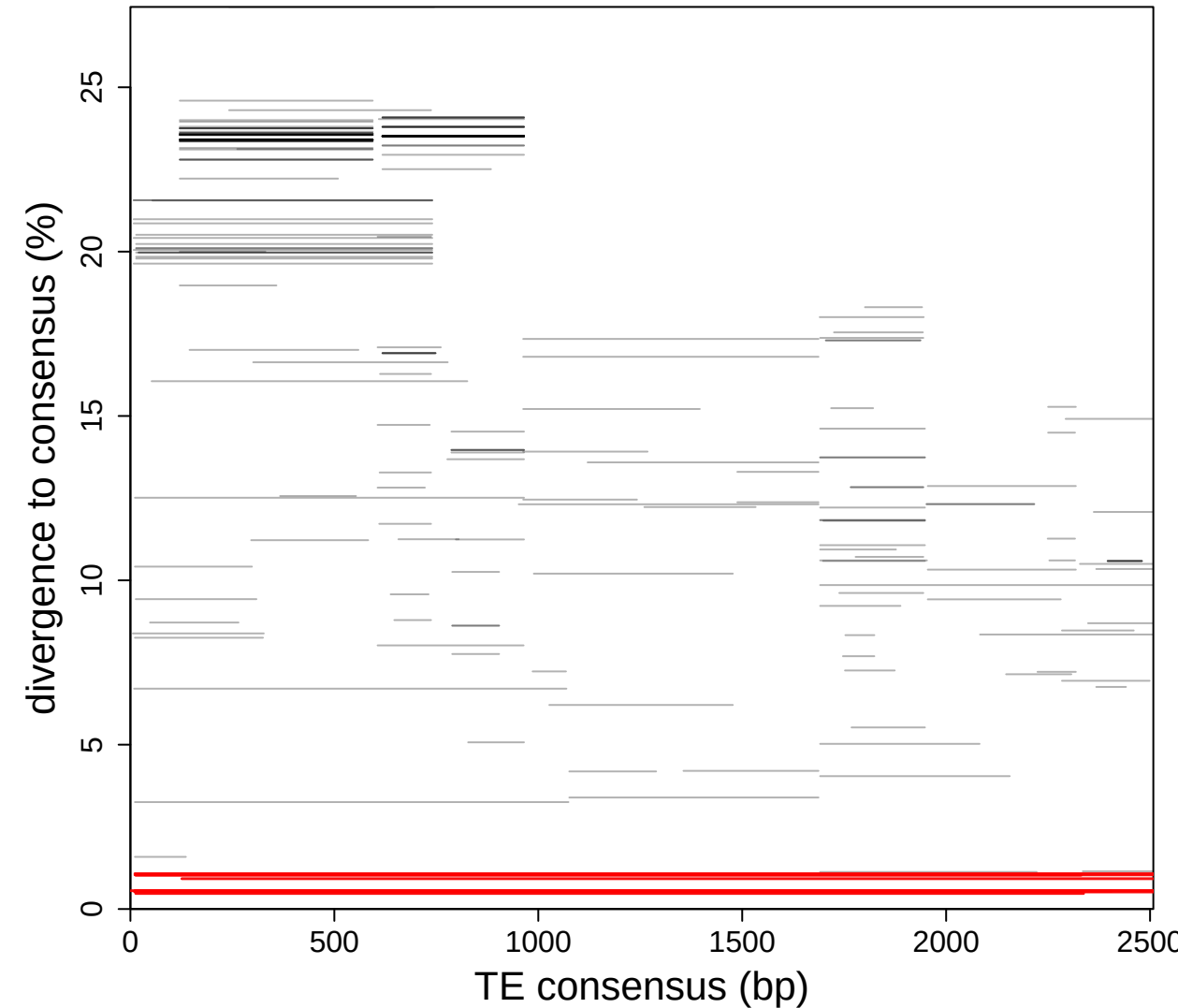


81.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.1
size: 8640bp; fragments: 22973; full length: 3 (>=7776bp)

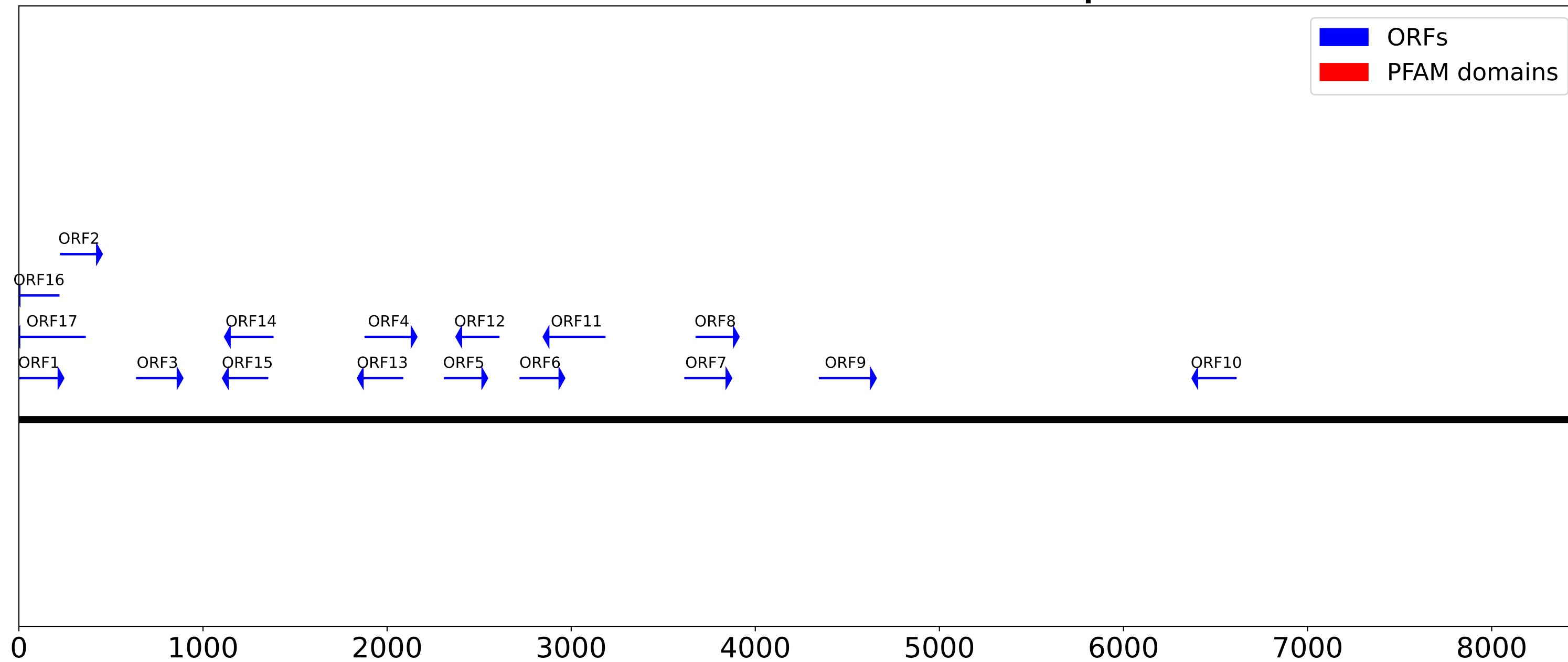
After TEtrimmer Extended plot Blue lines are boundaries



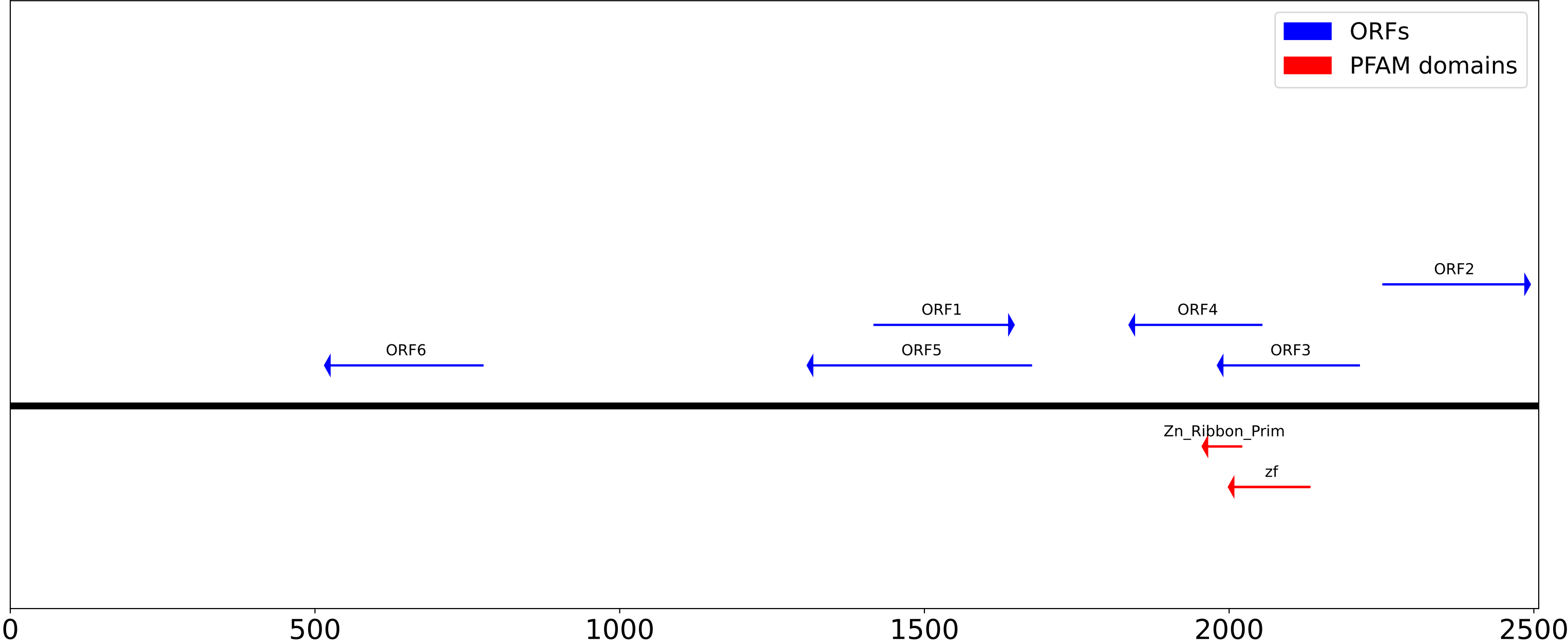
TE: rnd_5_family_13281
size: 2508bp; fragments: 228; full length: 9 (>=2257.2bp)



After TEtrimmer ORF and PFAM domain plot



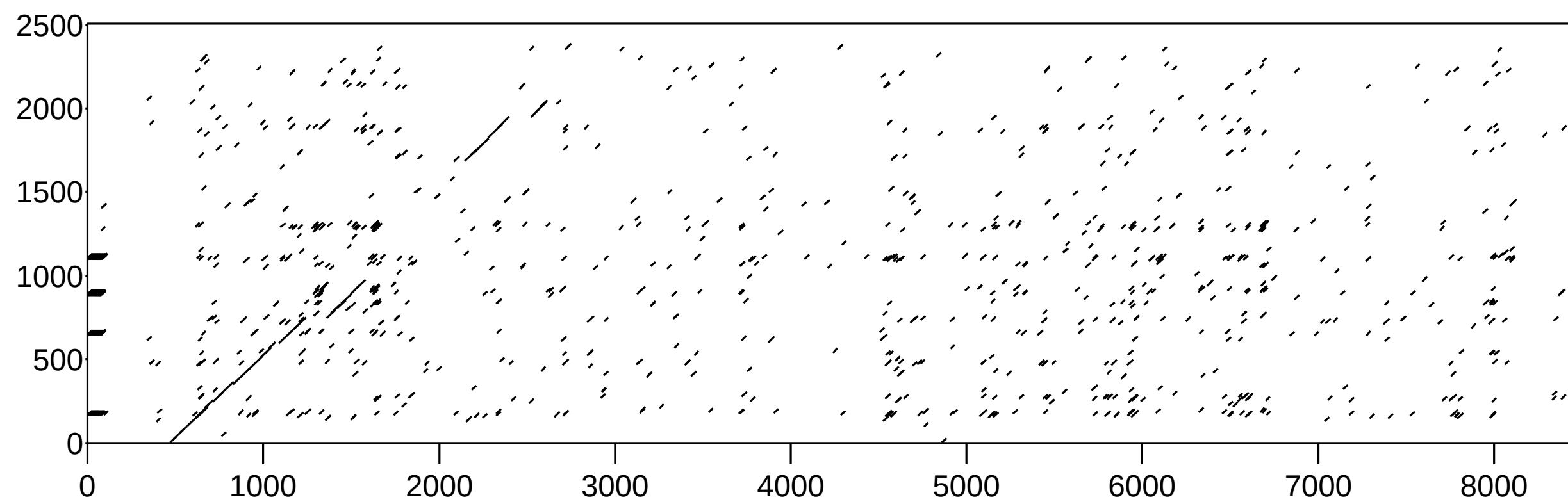
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)