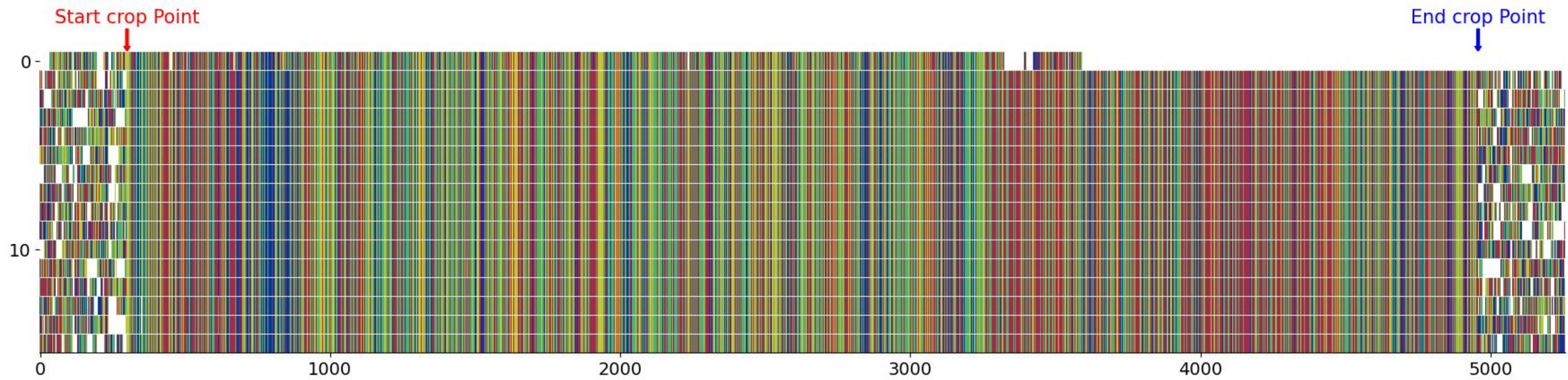




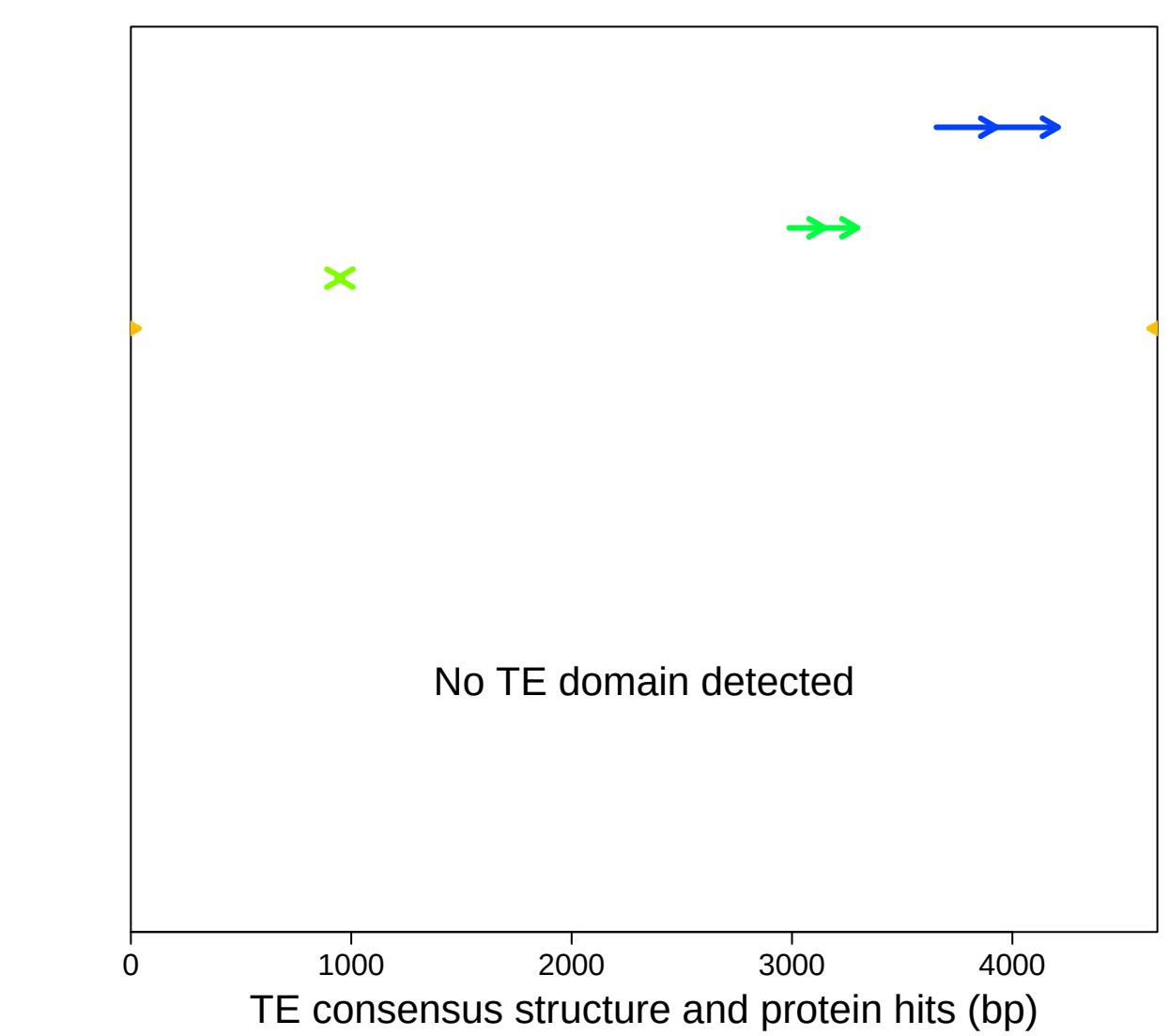
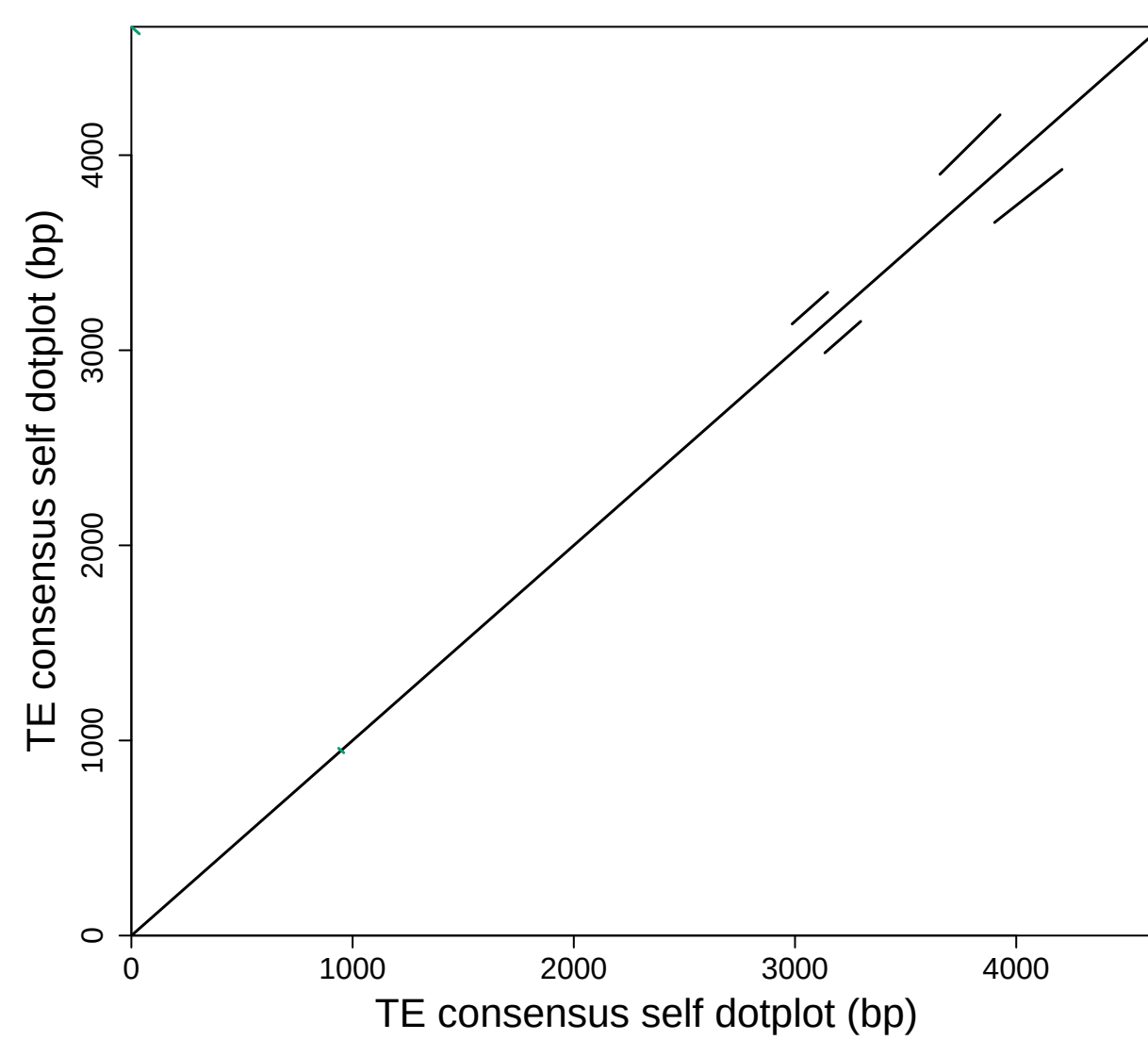
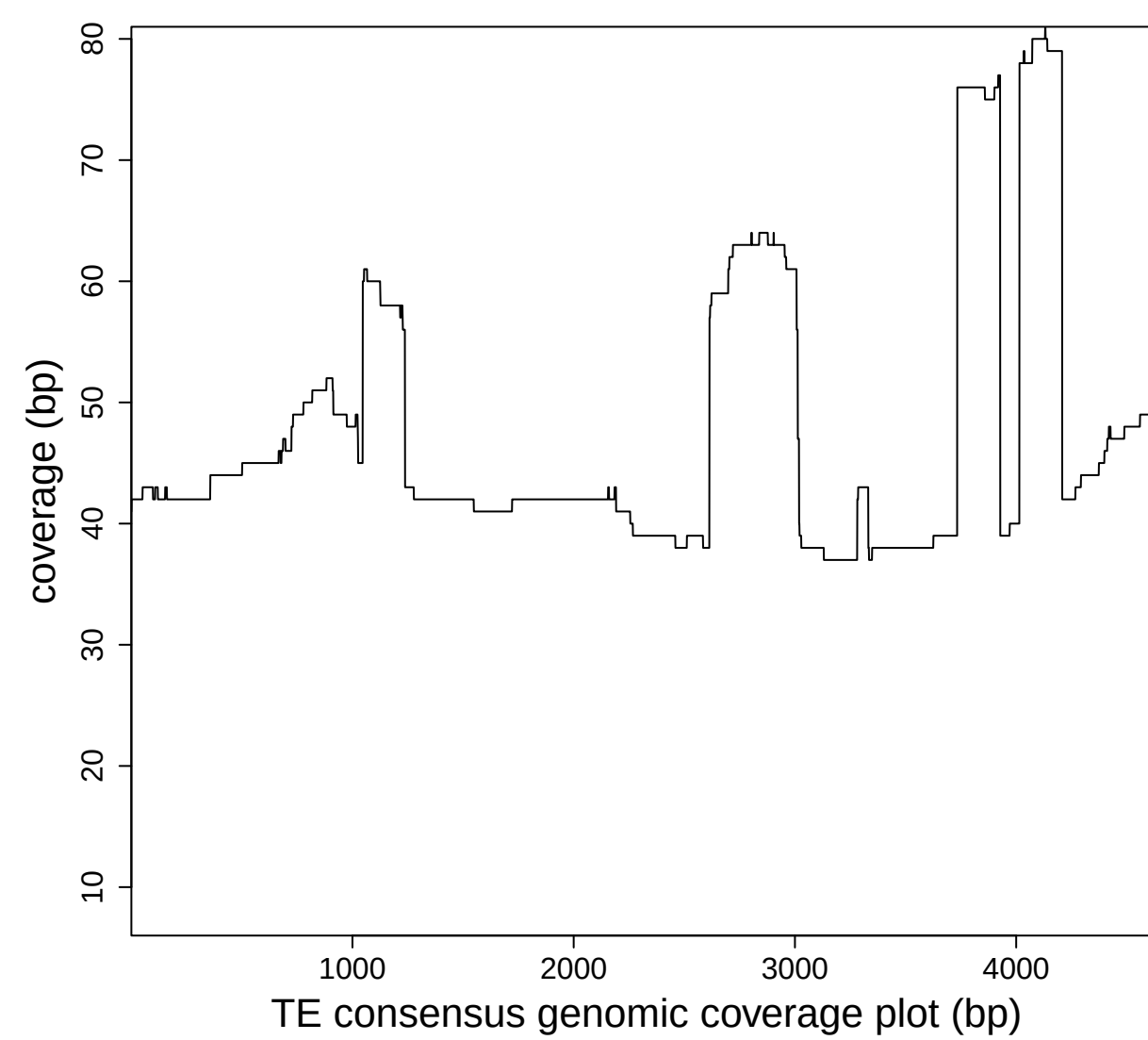
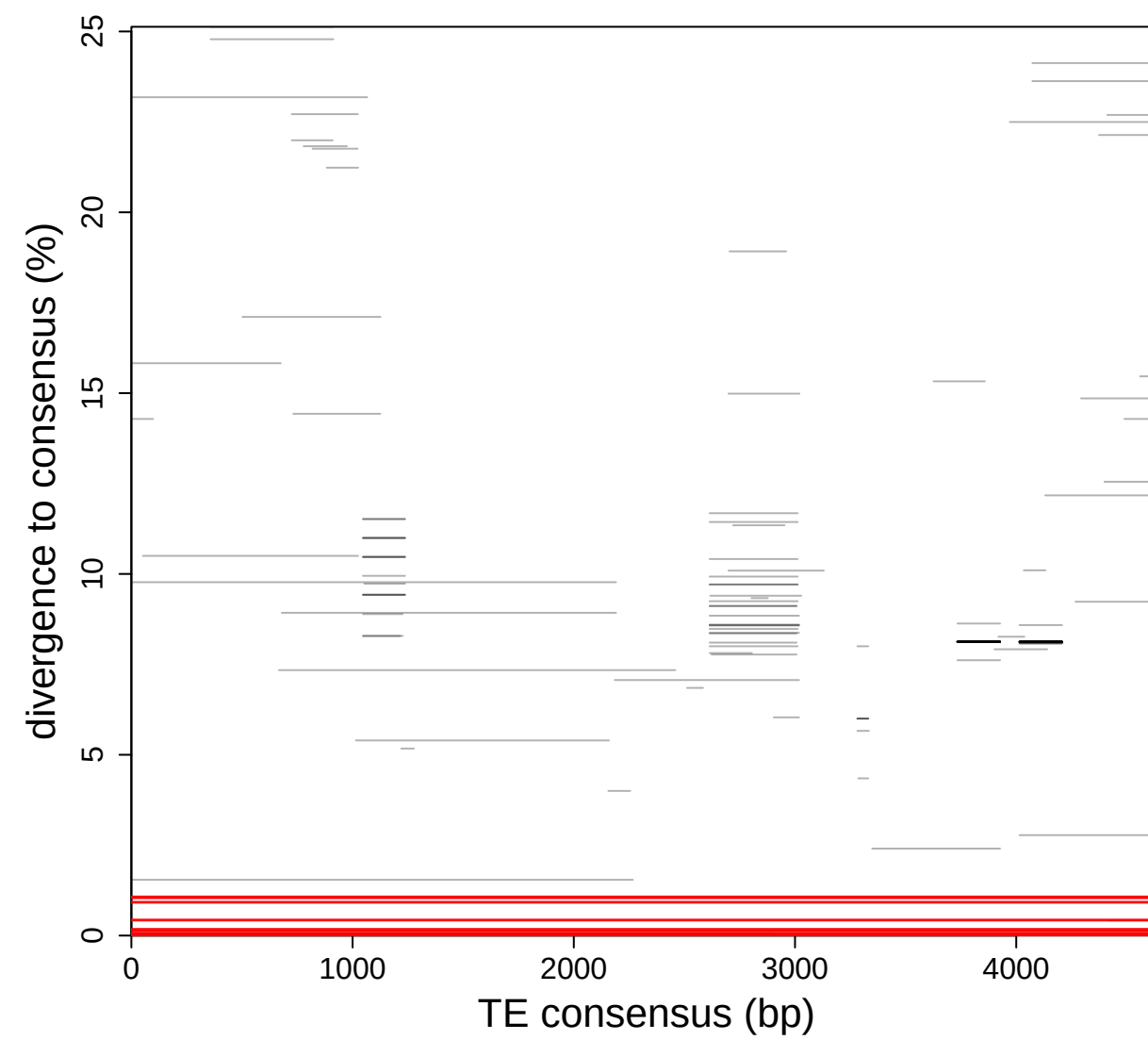
Start crop Point

End crop Point



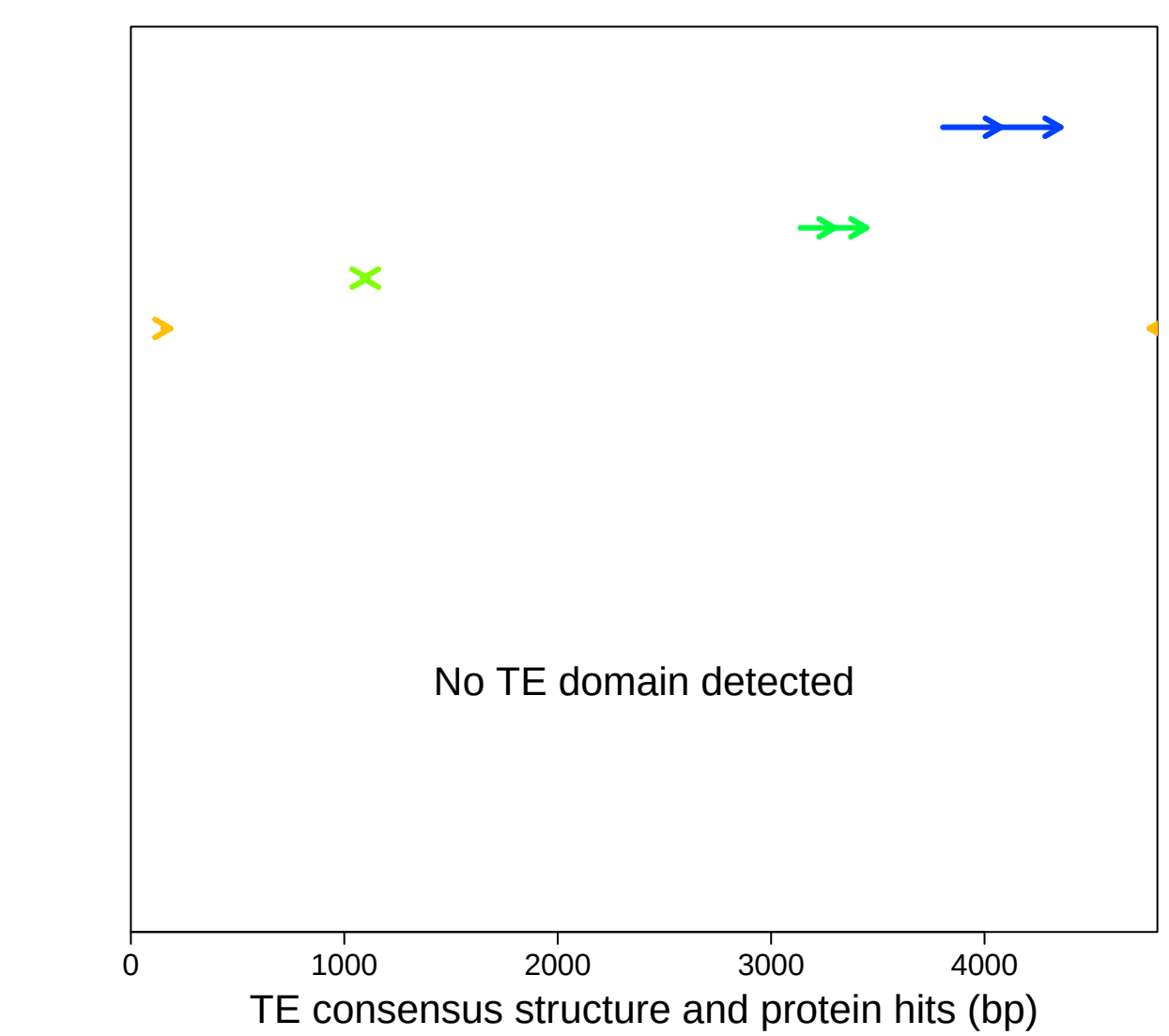
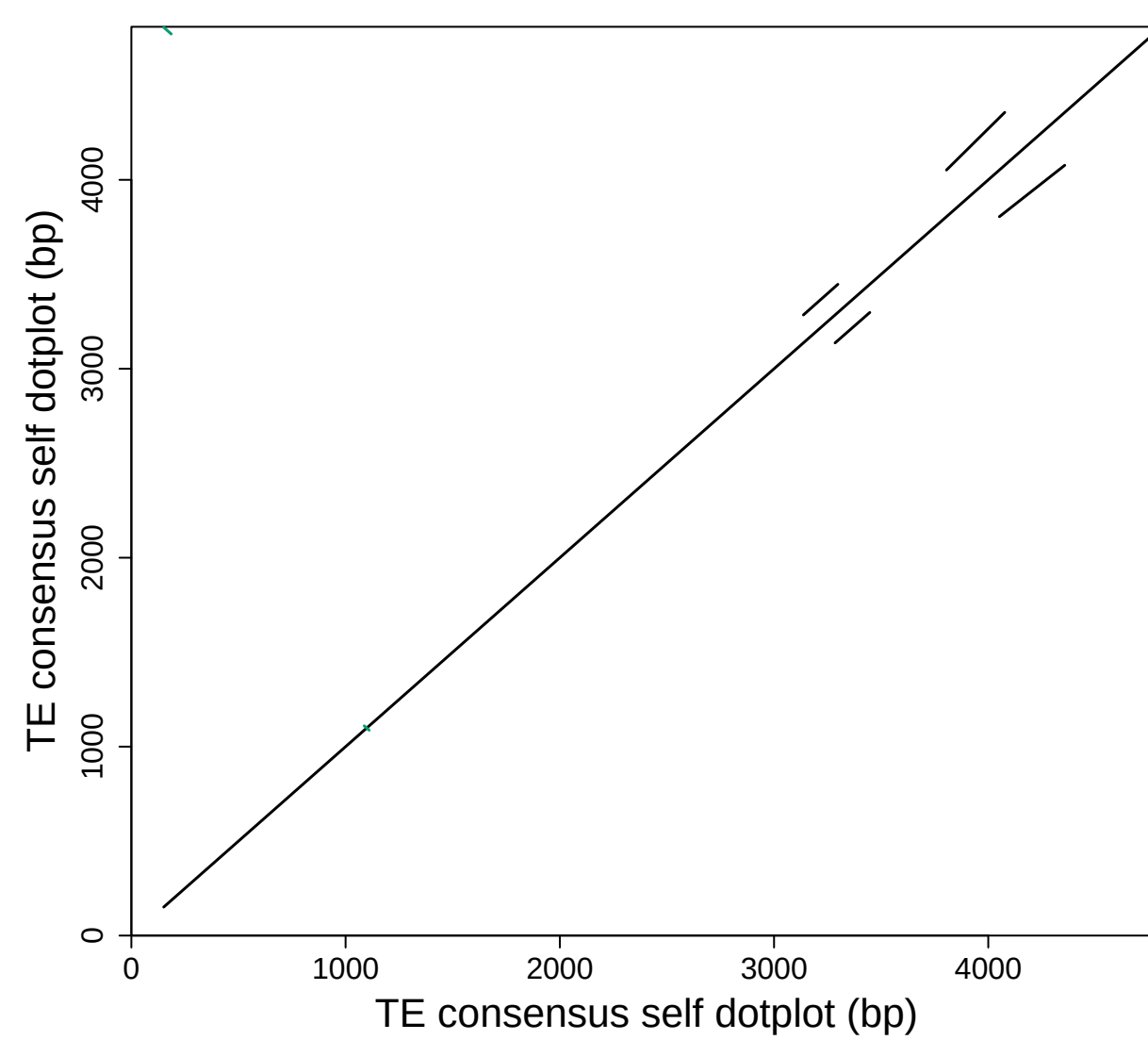
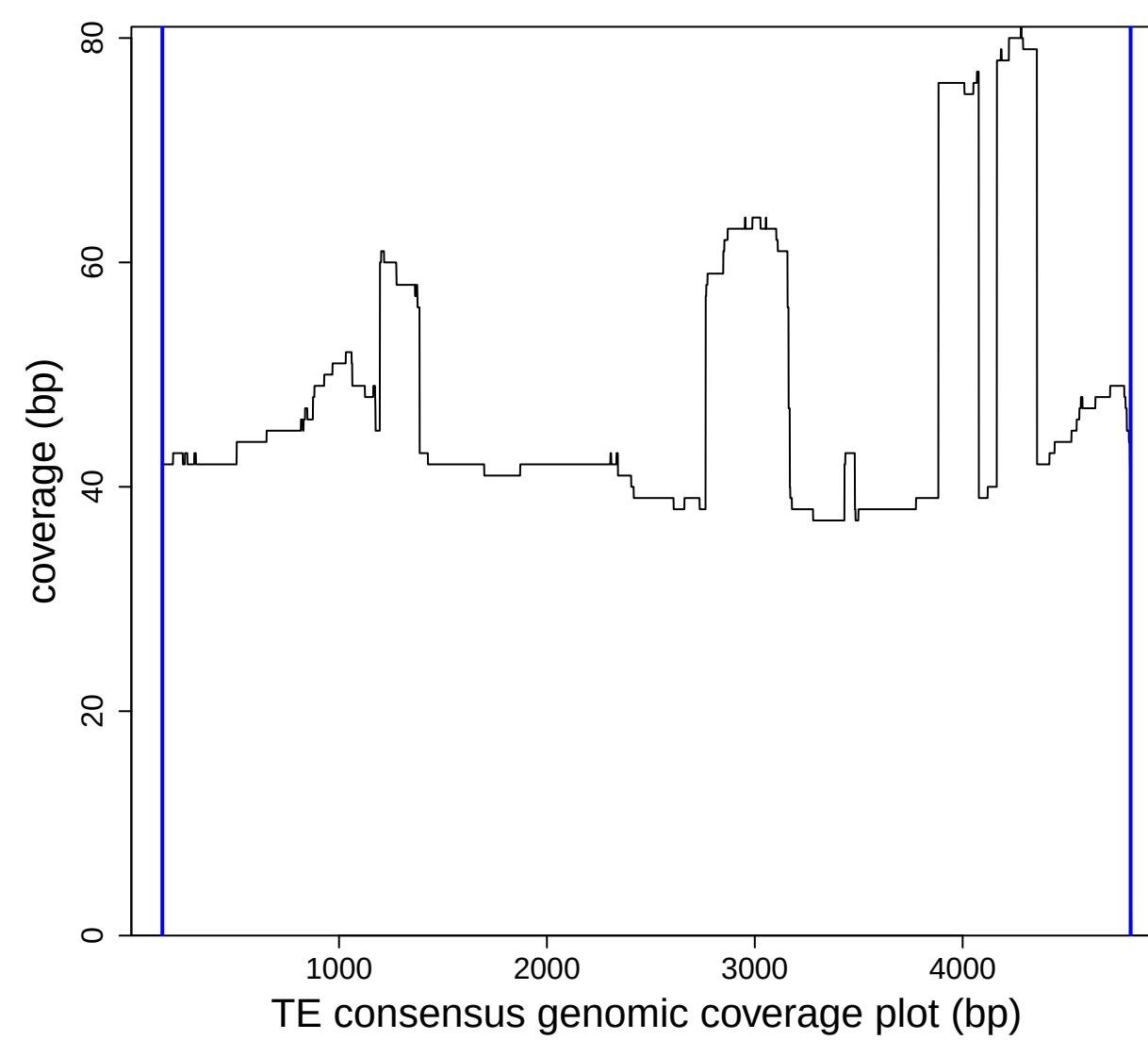
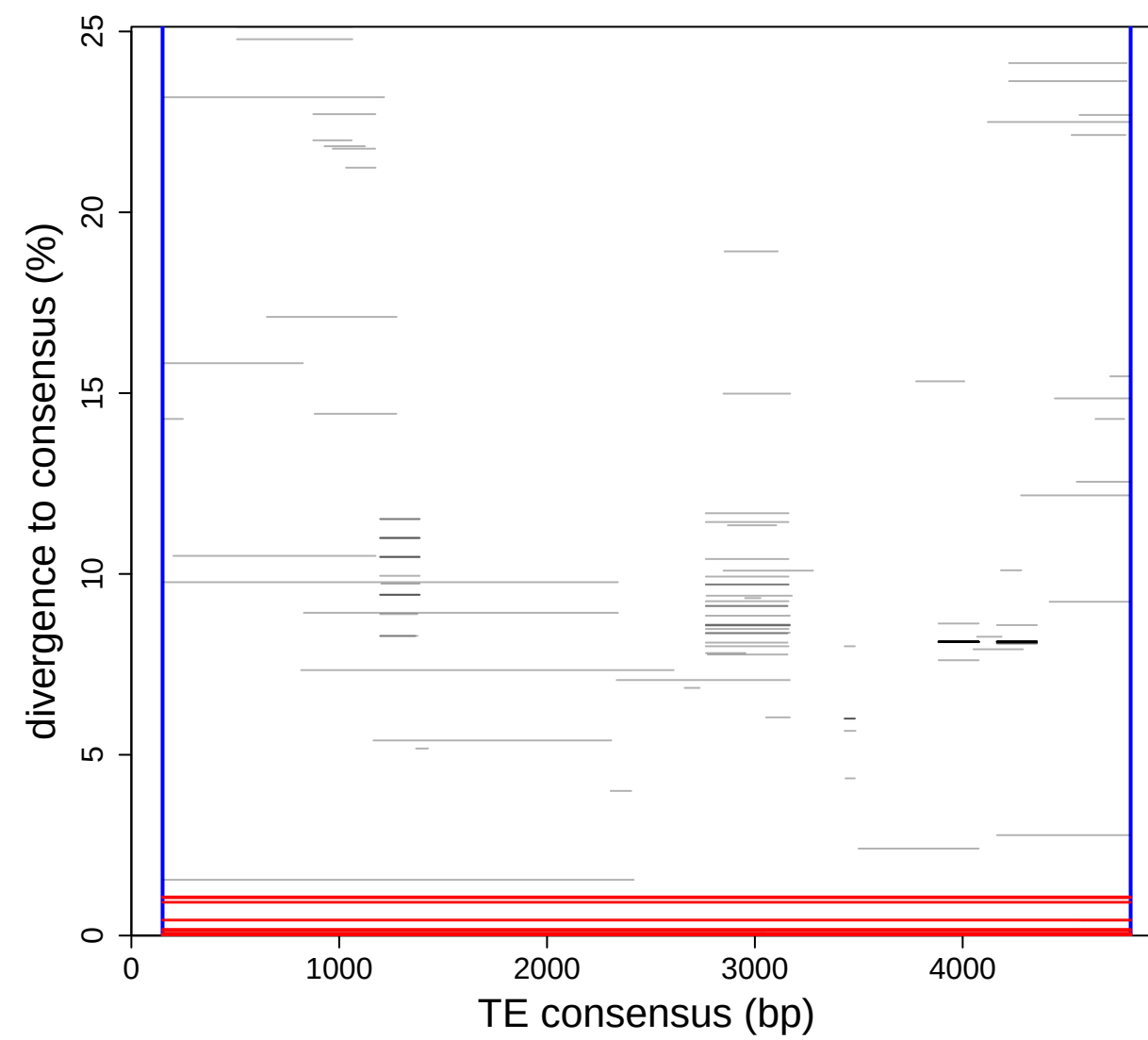
ta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.f
size: 4658bp; fragments: 205; full length: 34 (≥ 4192.2 bp)

After TEtrimmer 4658 bp

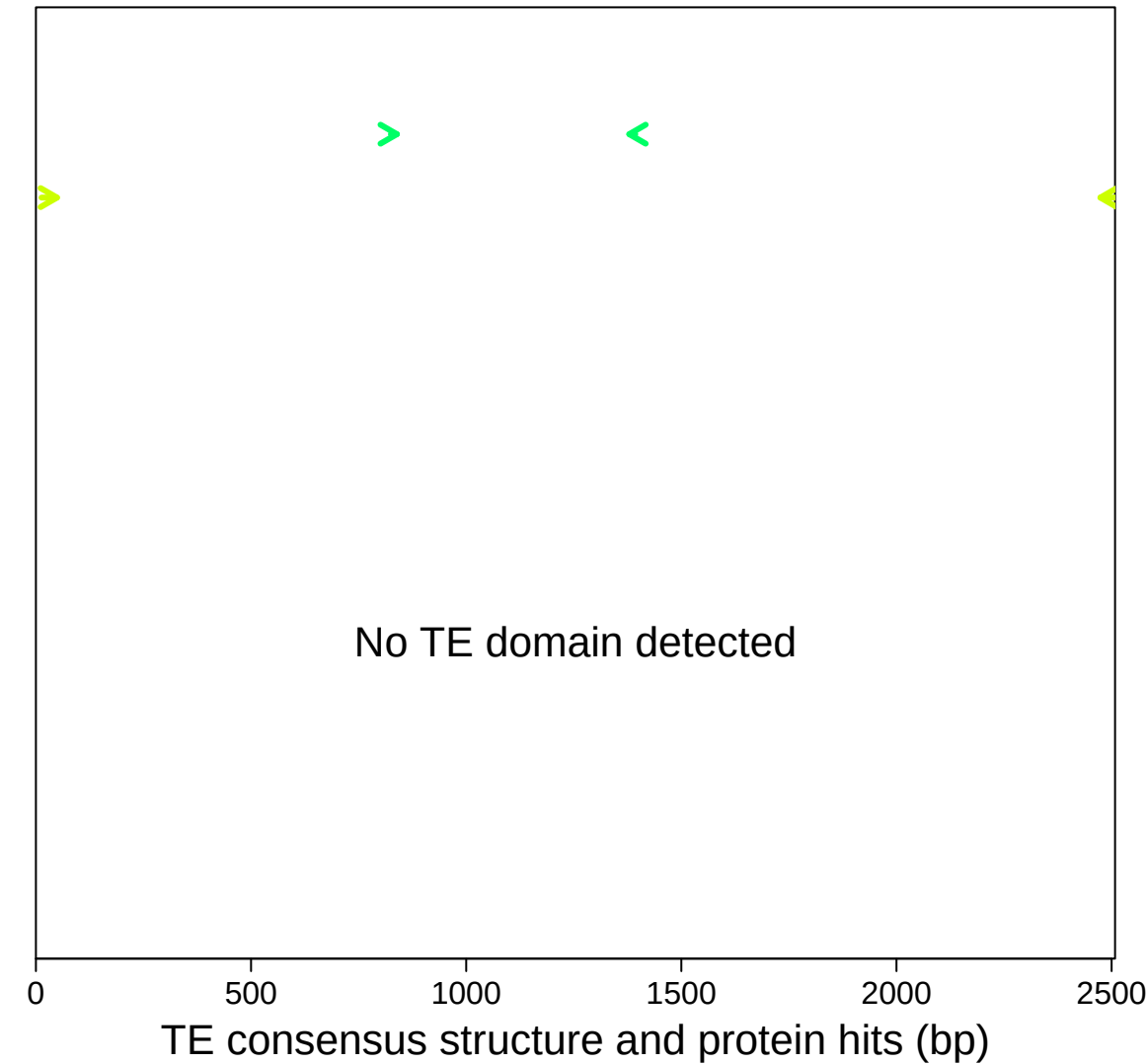
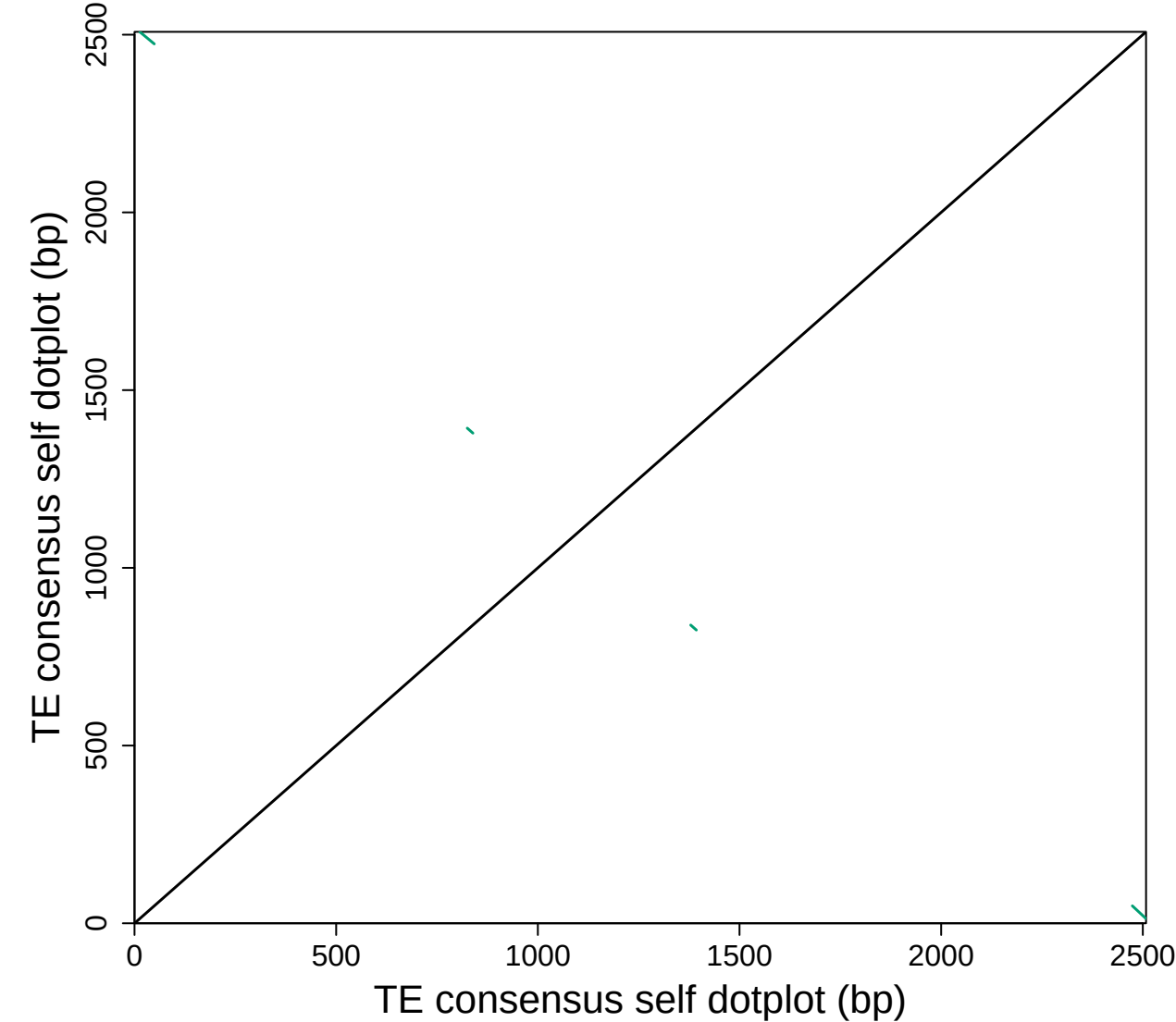
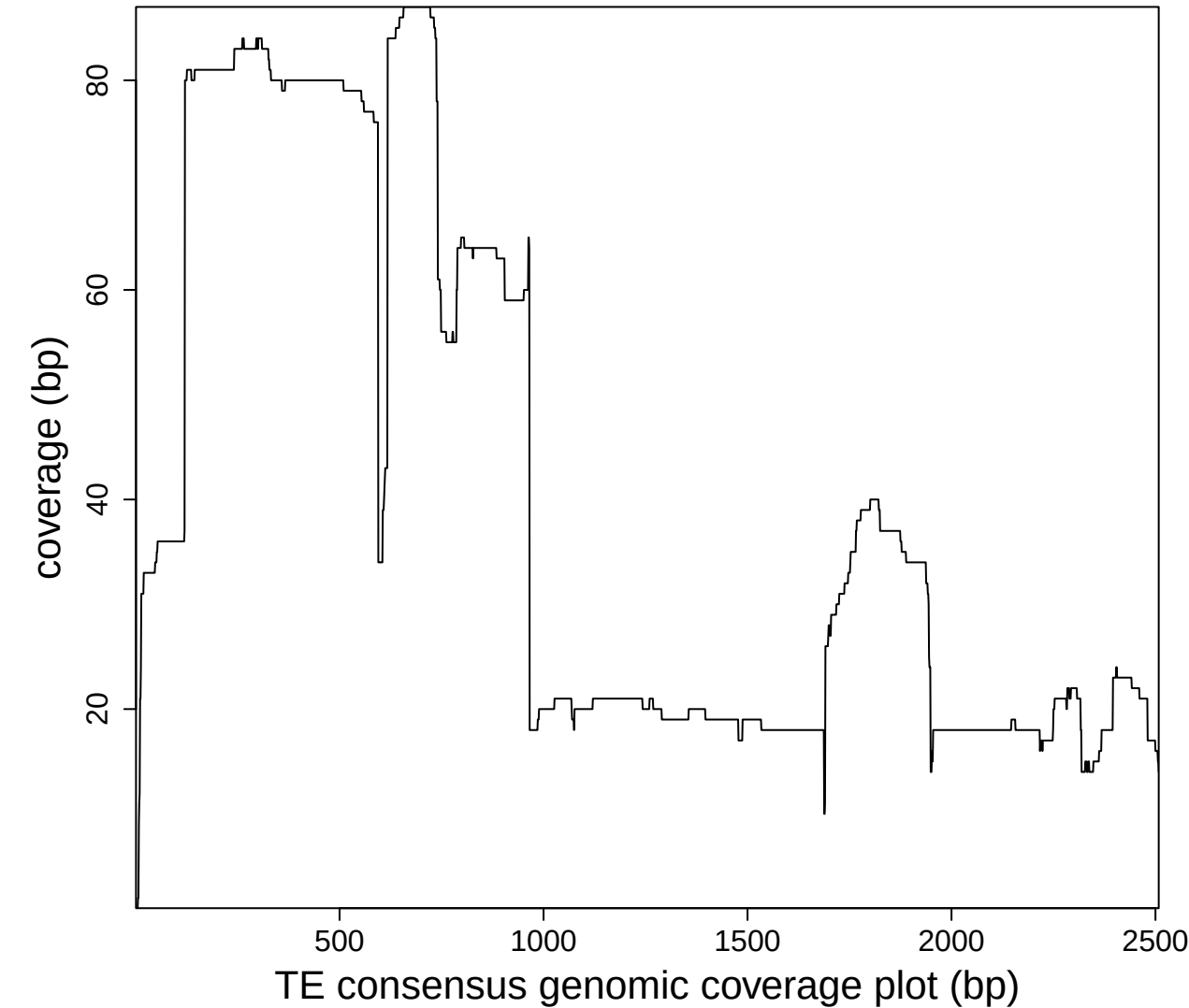
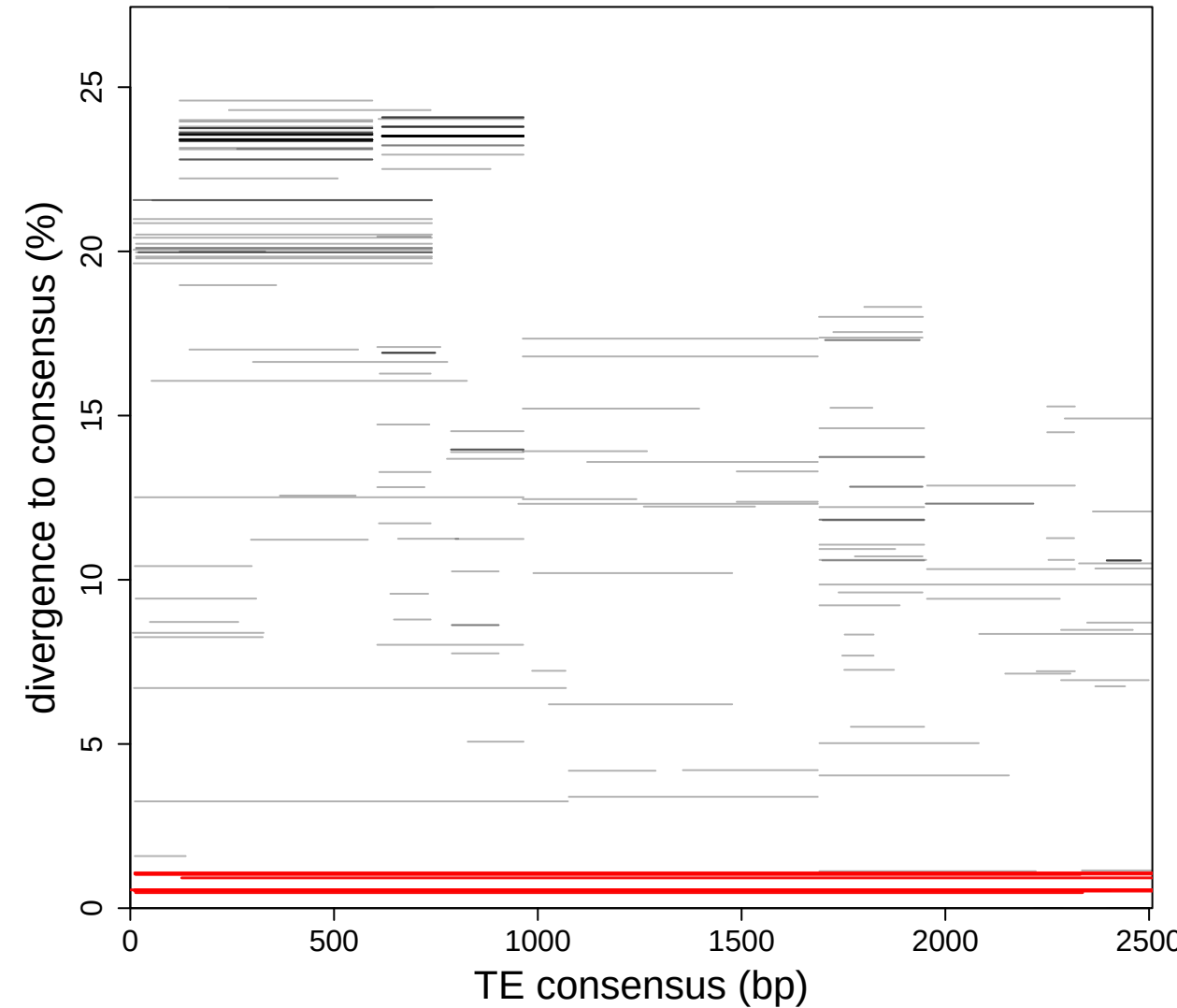


fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_
size: 4958bp; fragments: 205; full length: 33 (>=4462.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

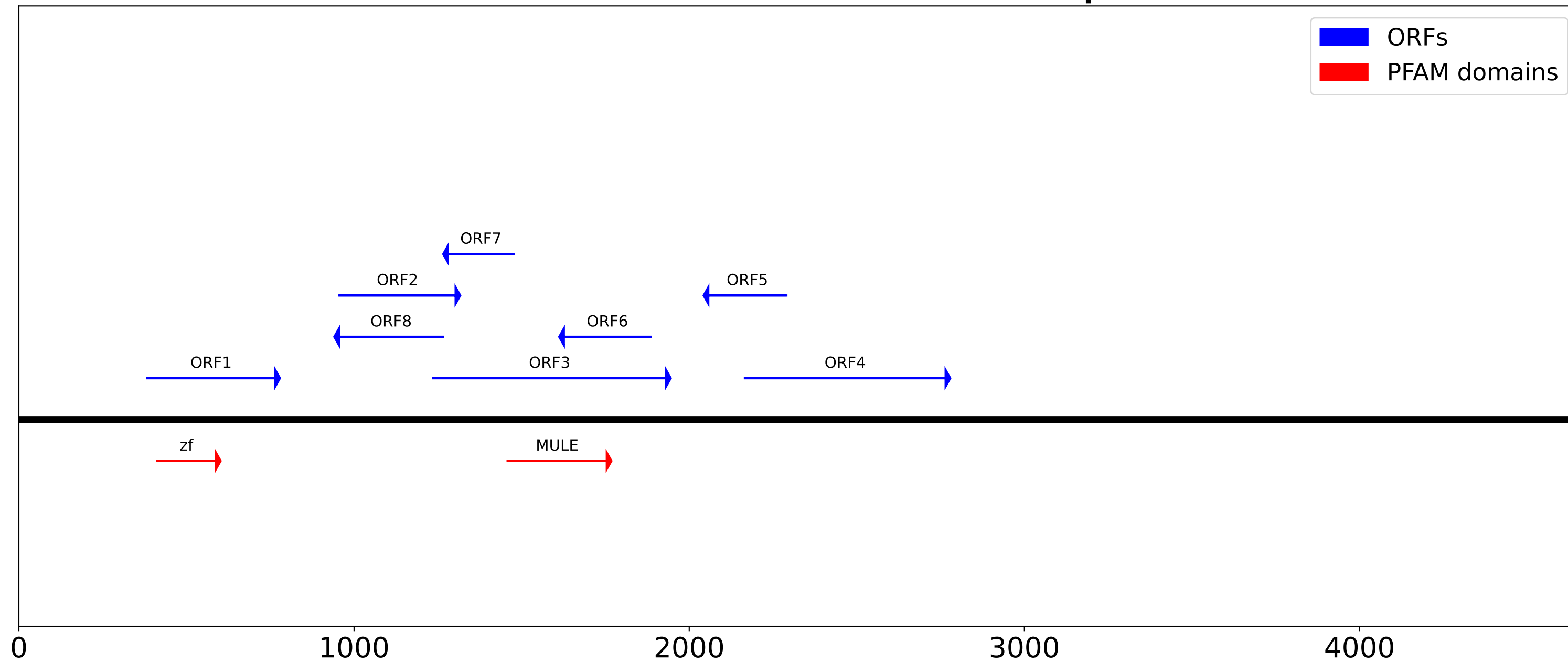


TE: rnd_5_family_13281
size: 2508bp; fragments: 228; full length: 9 (>=2257.2bp)

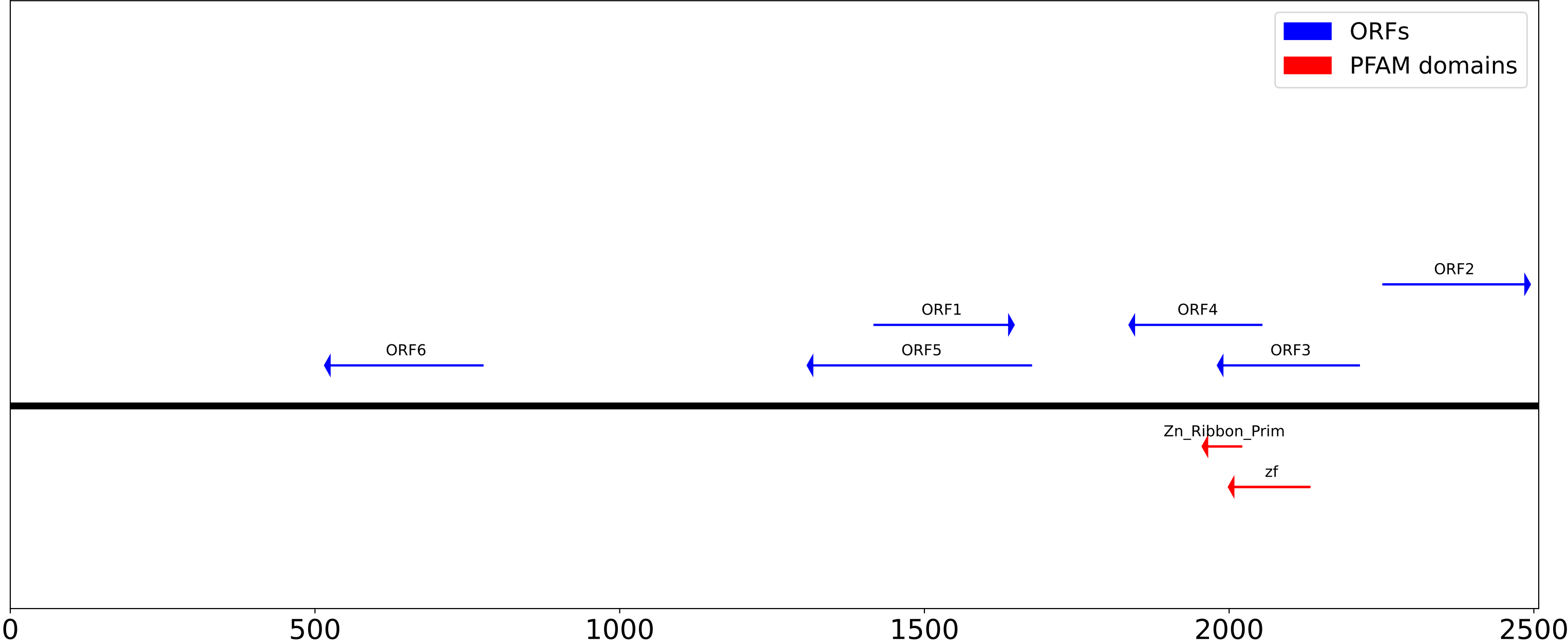


Before TEtrimmer 2508 bp

After TEtrimmer ORF and PFAM domain plot



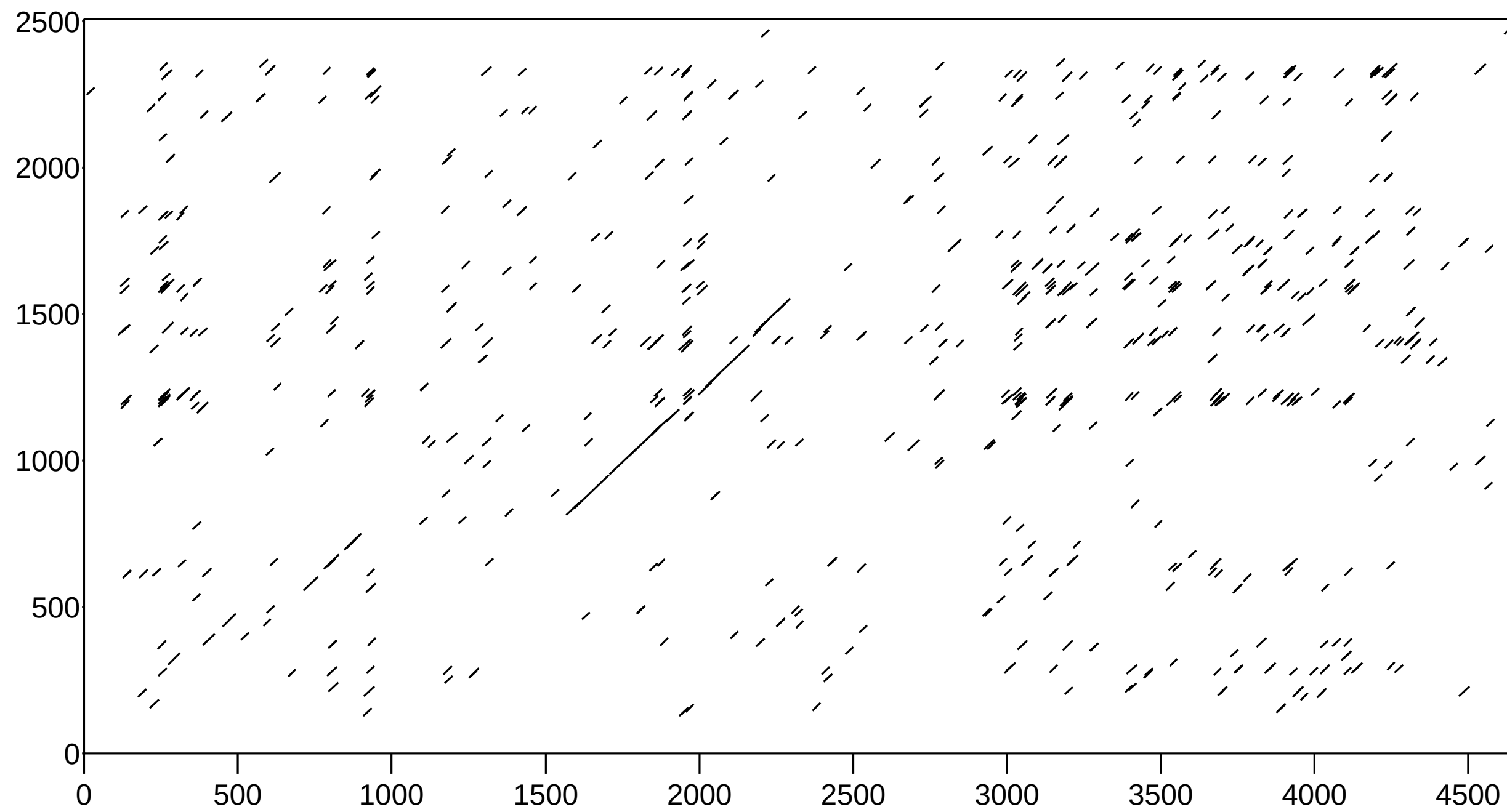
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)