

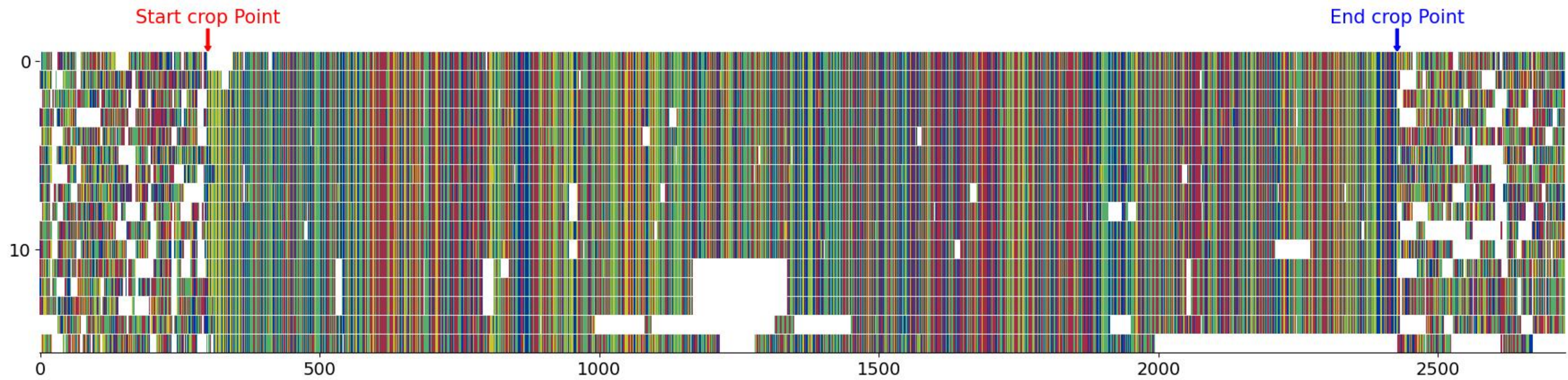
End crop Point

MSA length = 2128



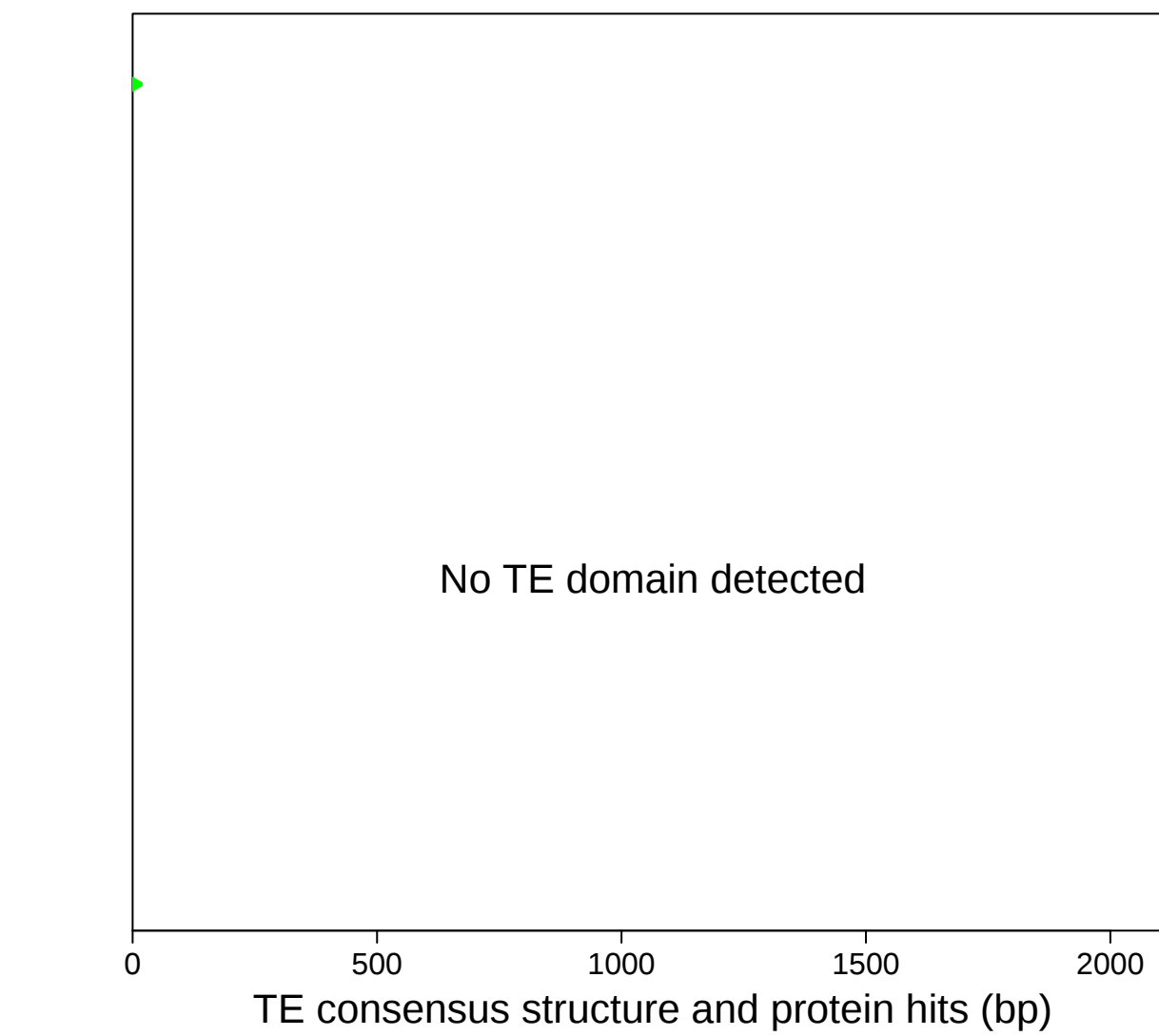
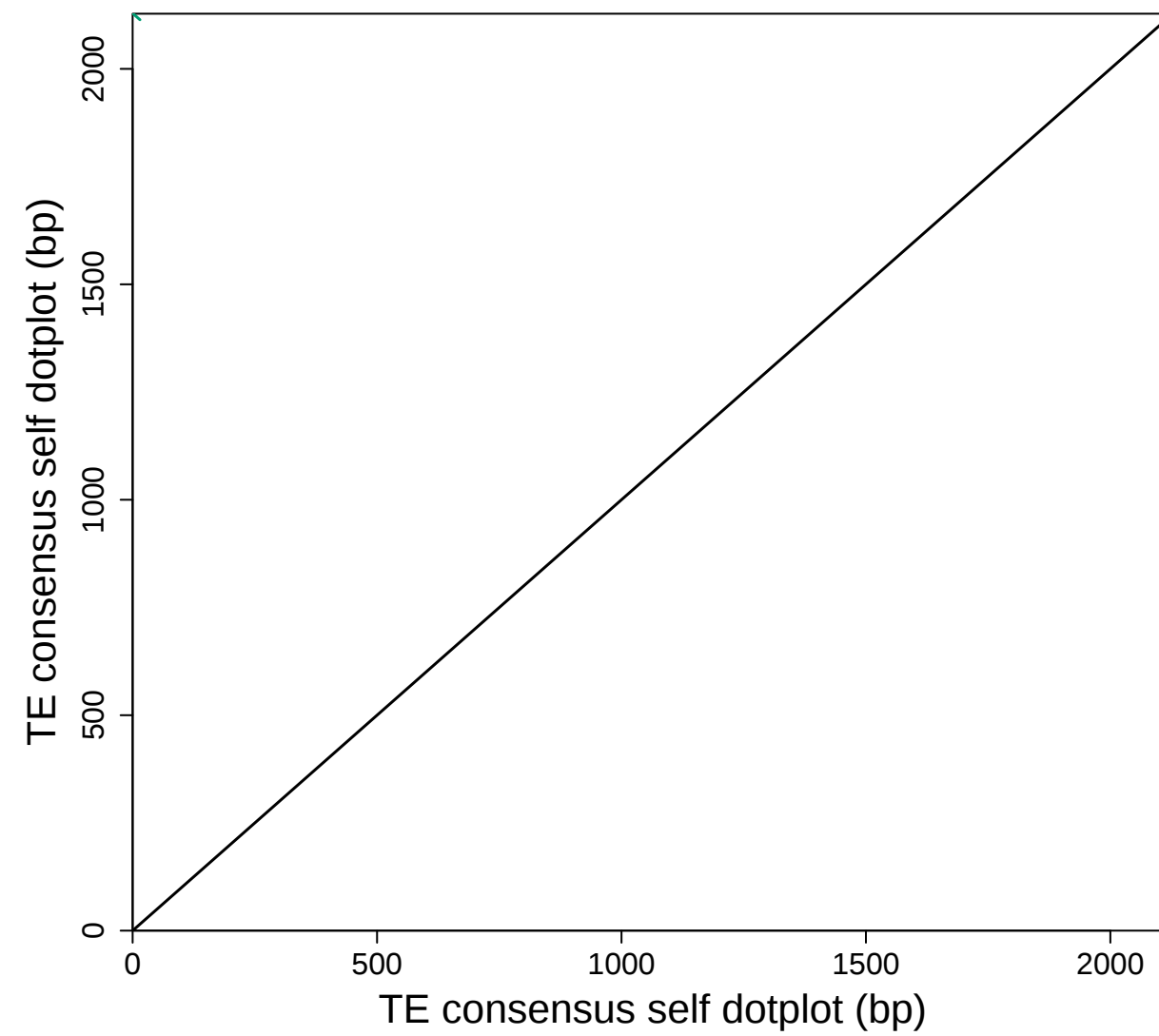
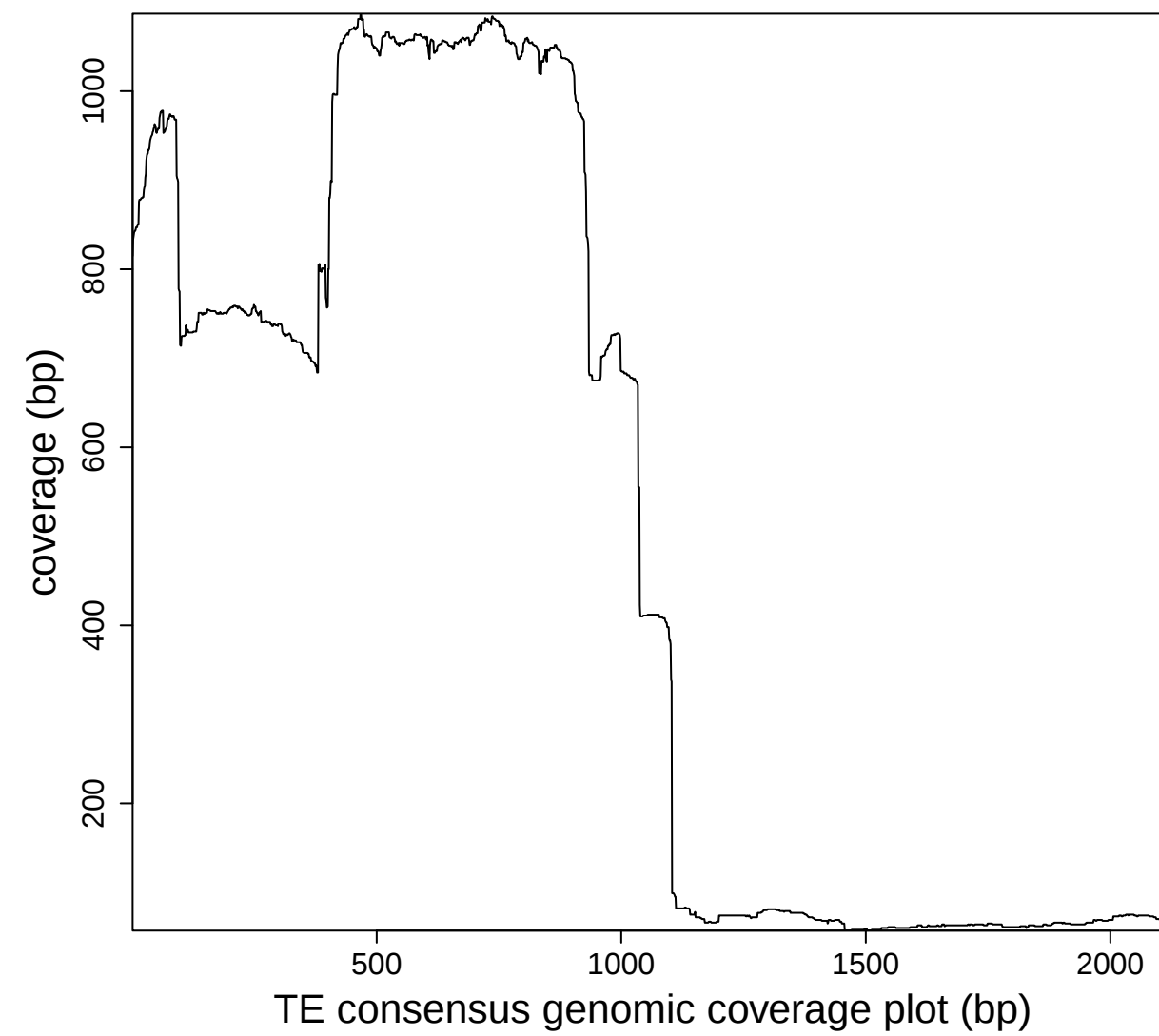
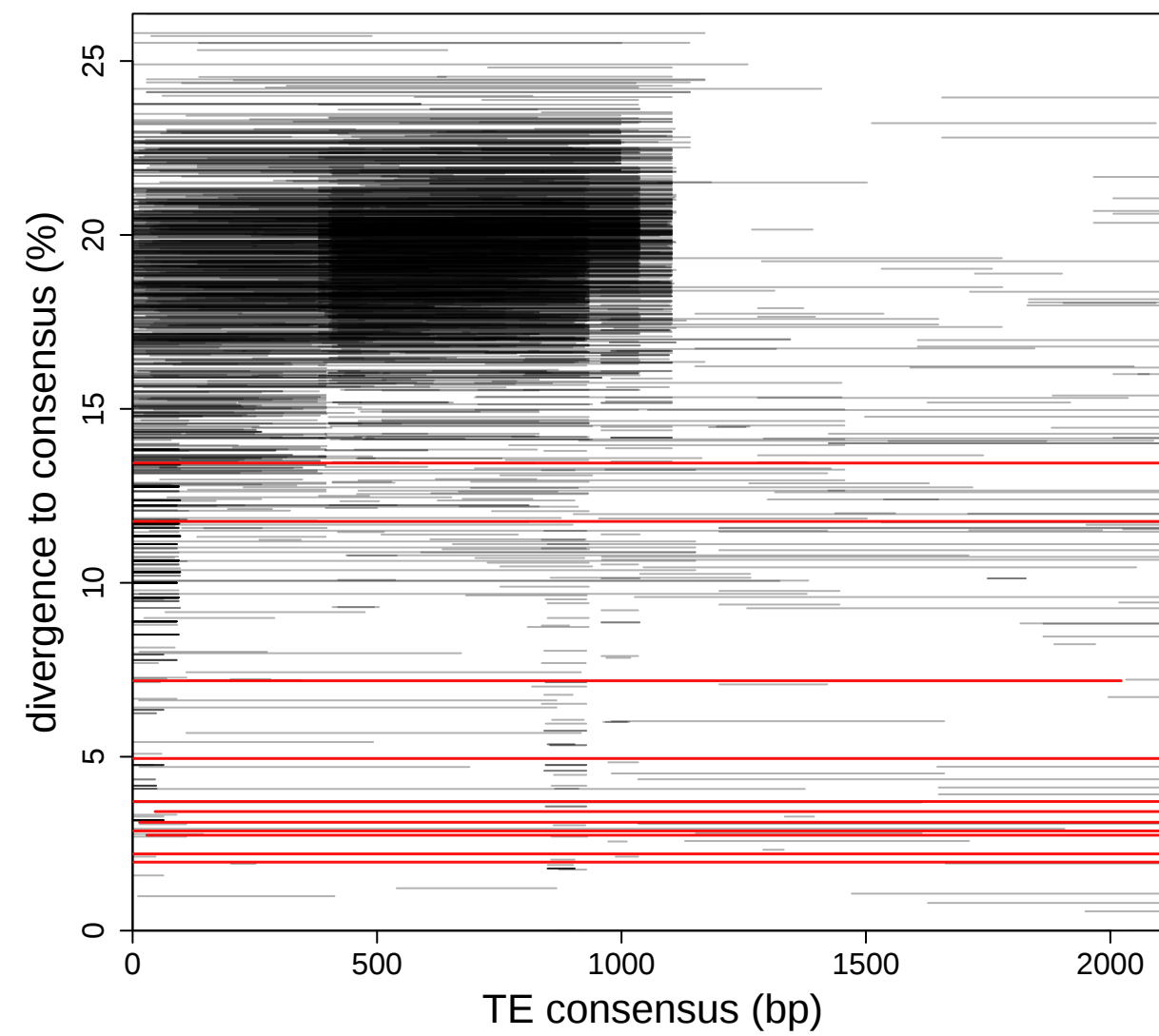
Start crop Point

End crop Point



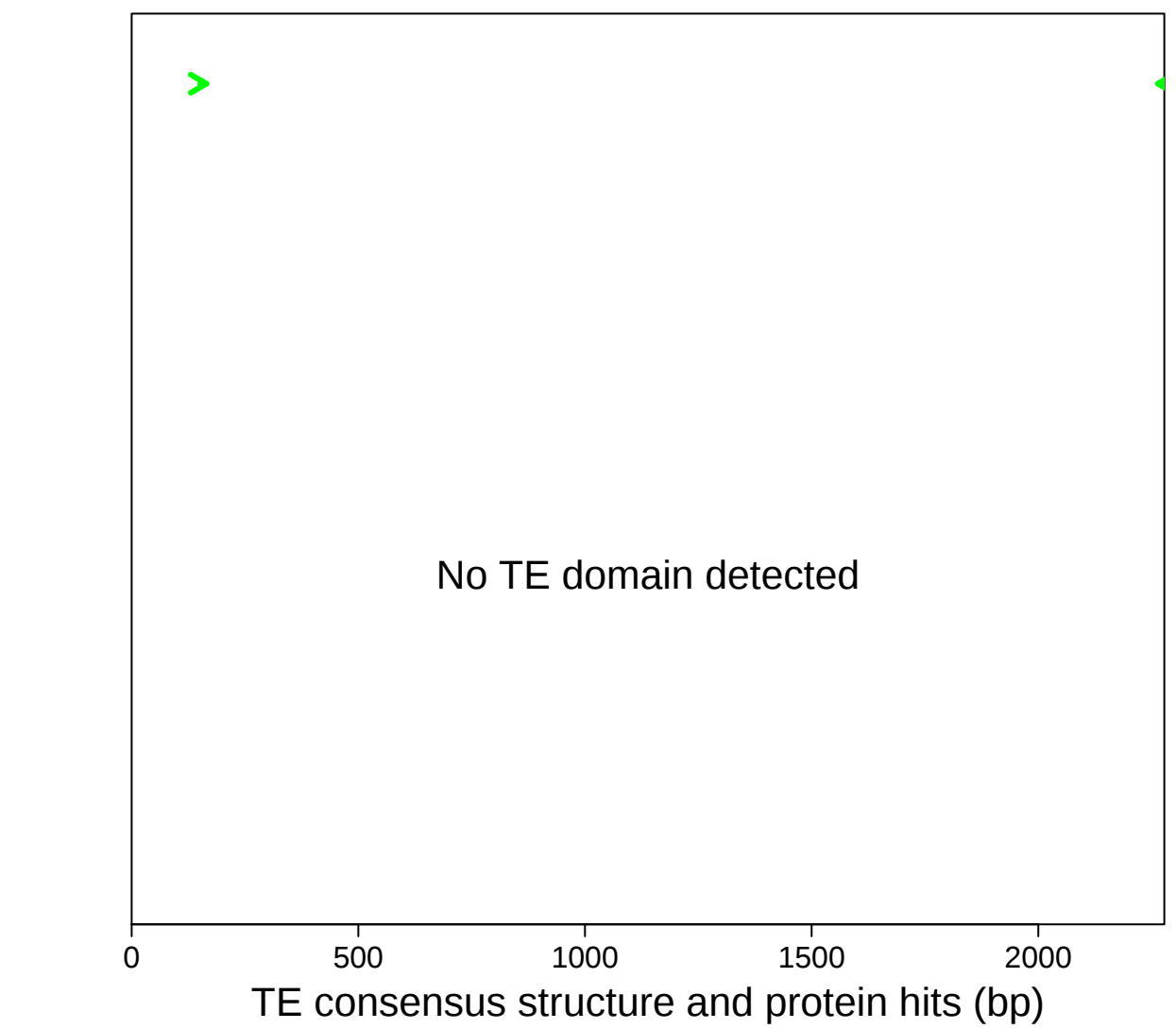
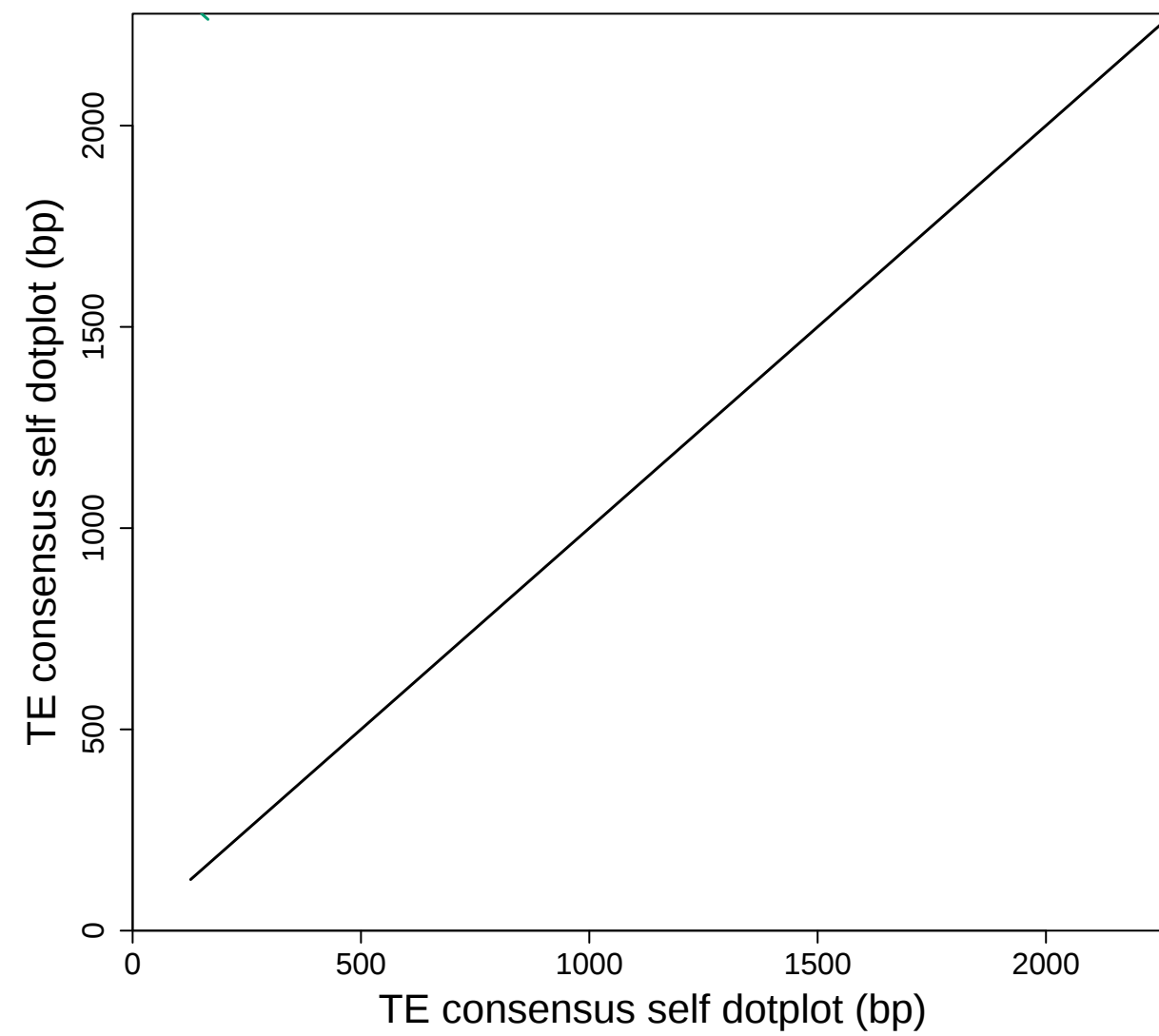
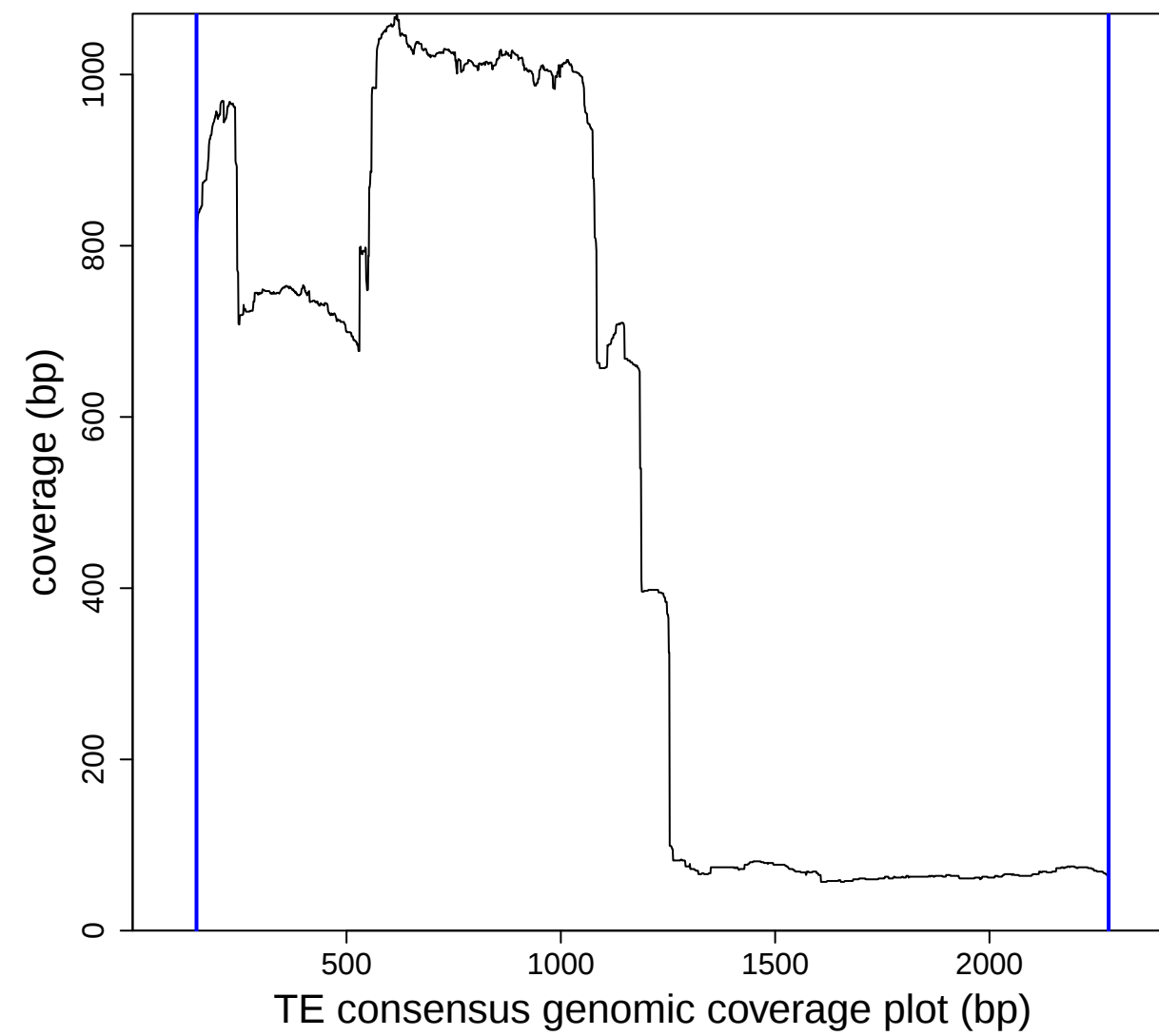
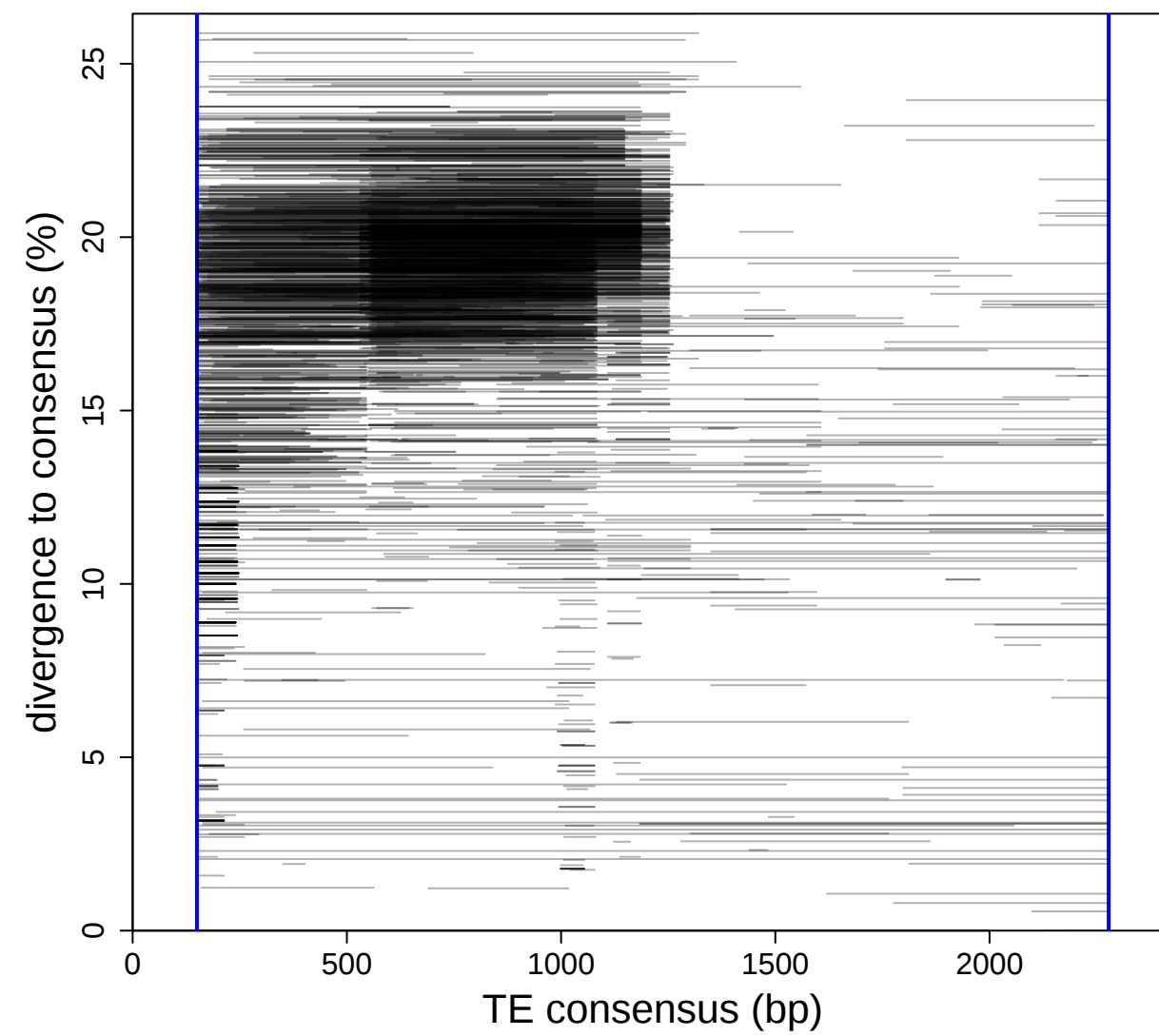
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 2128bp; fragments: 2080; full length: 11 (>=1915.2bp)

After TEtrimmer 2128 bp



6.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 2428bp; fragments: 2024; full length: 0 (>=2185.2bp)

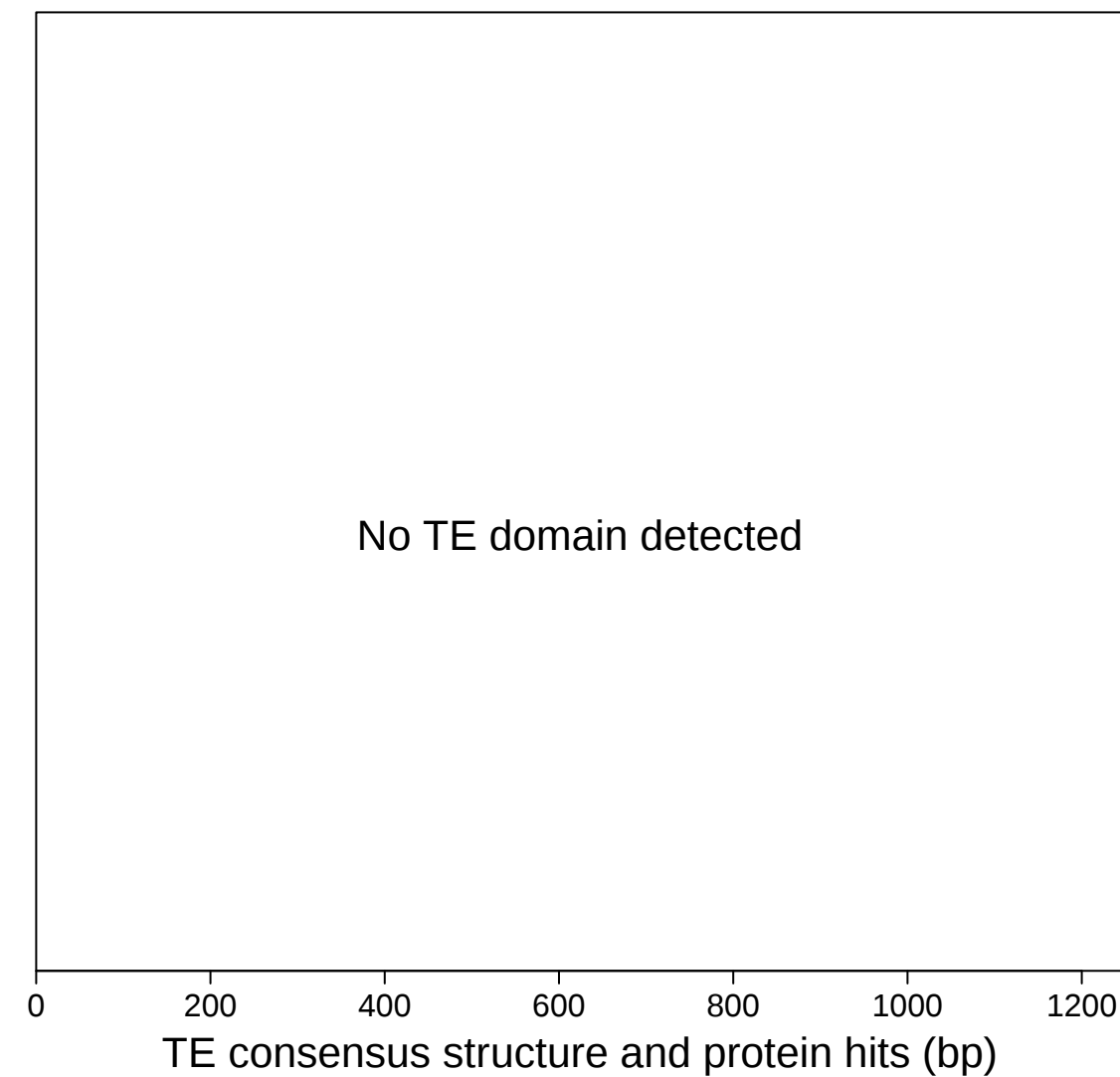
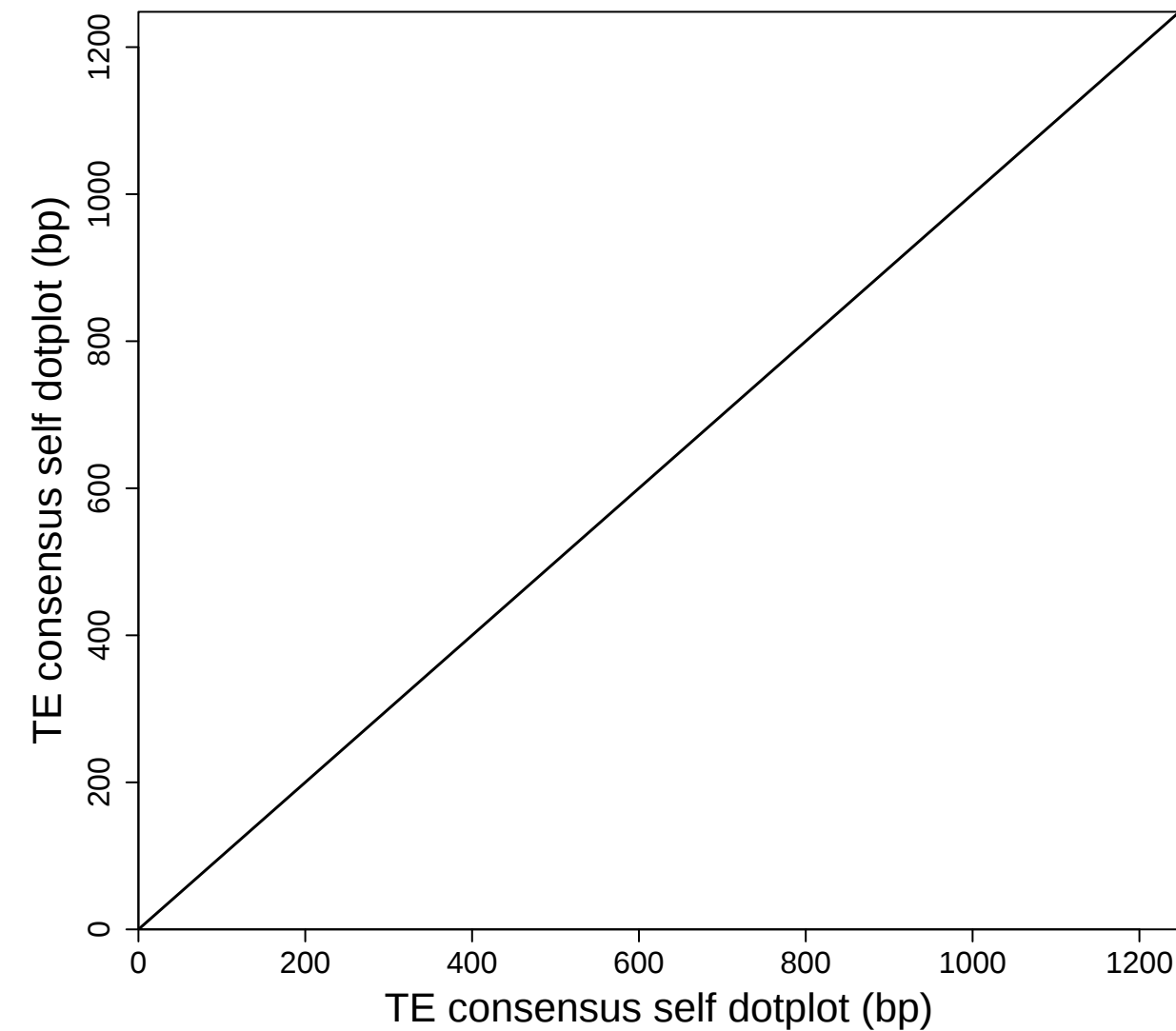
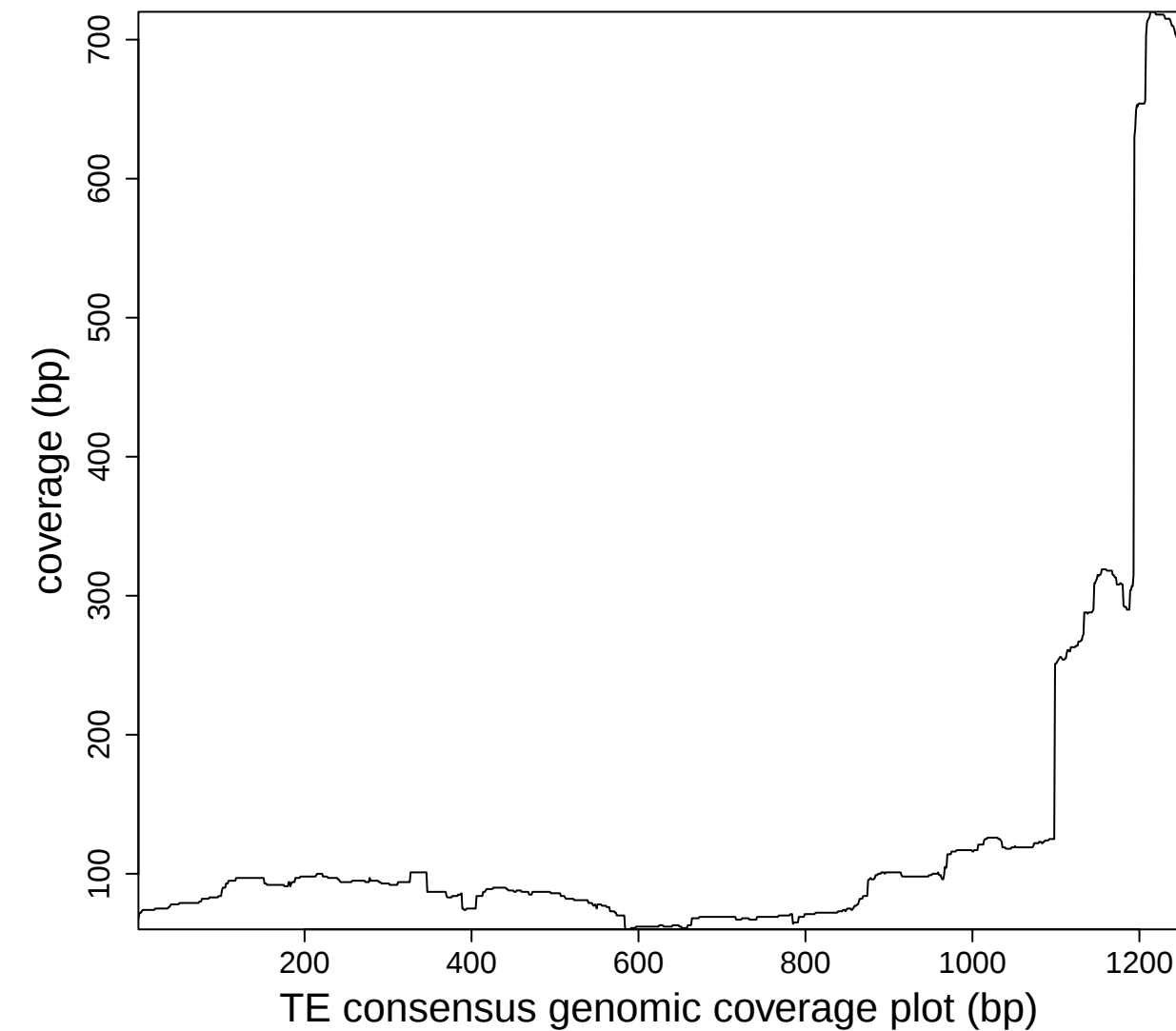
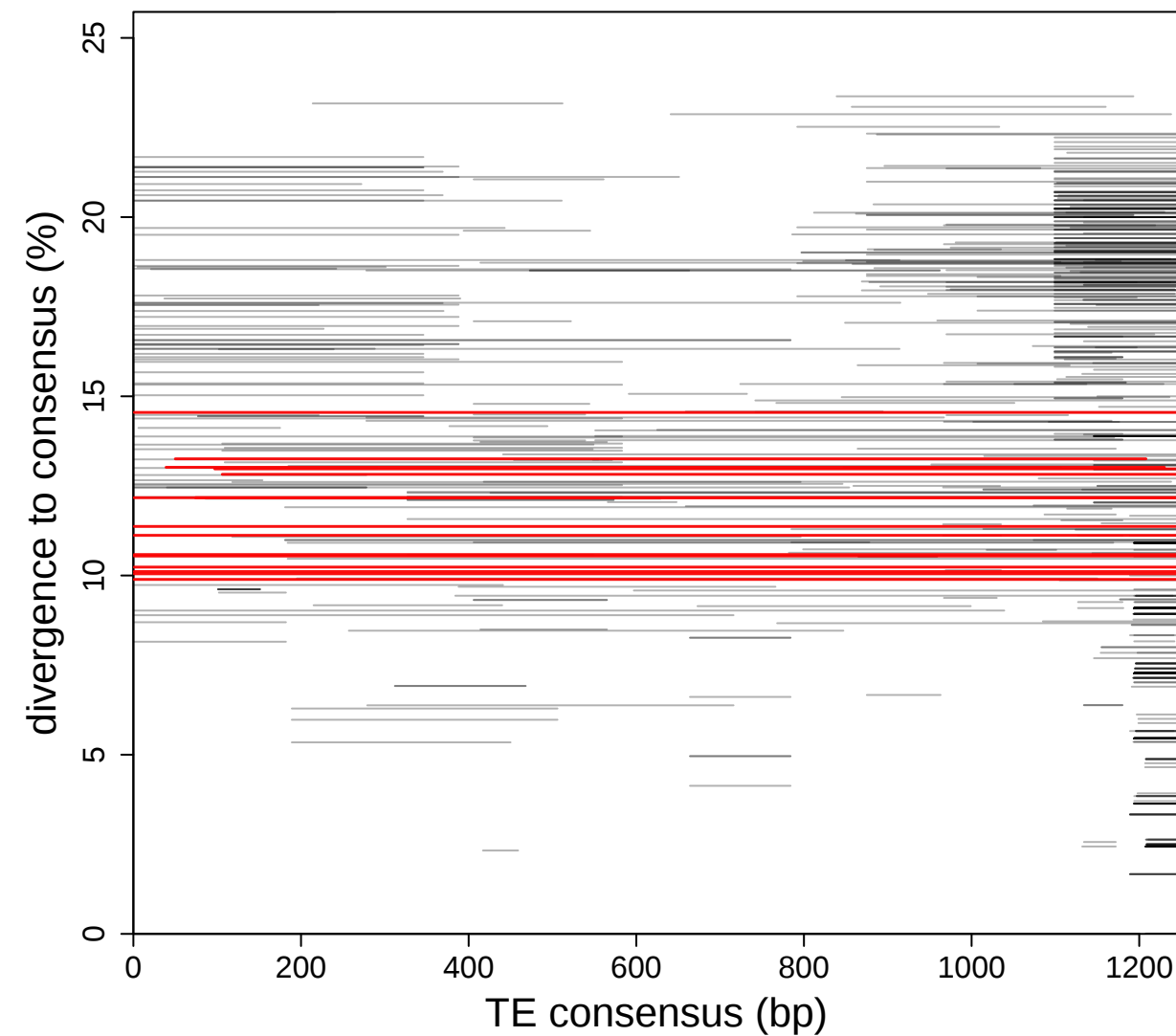
After TEtrimmer Extended plot Blue lines are boundaries



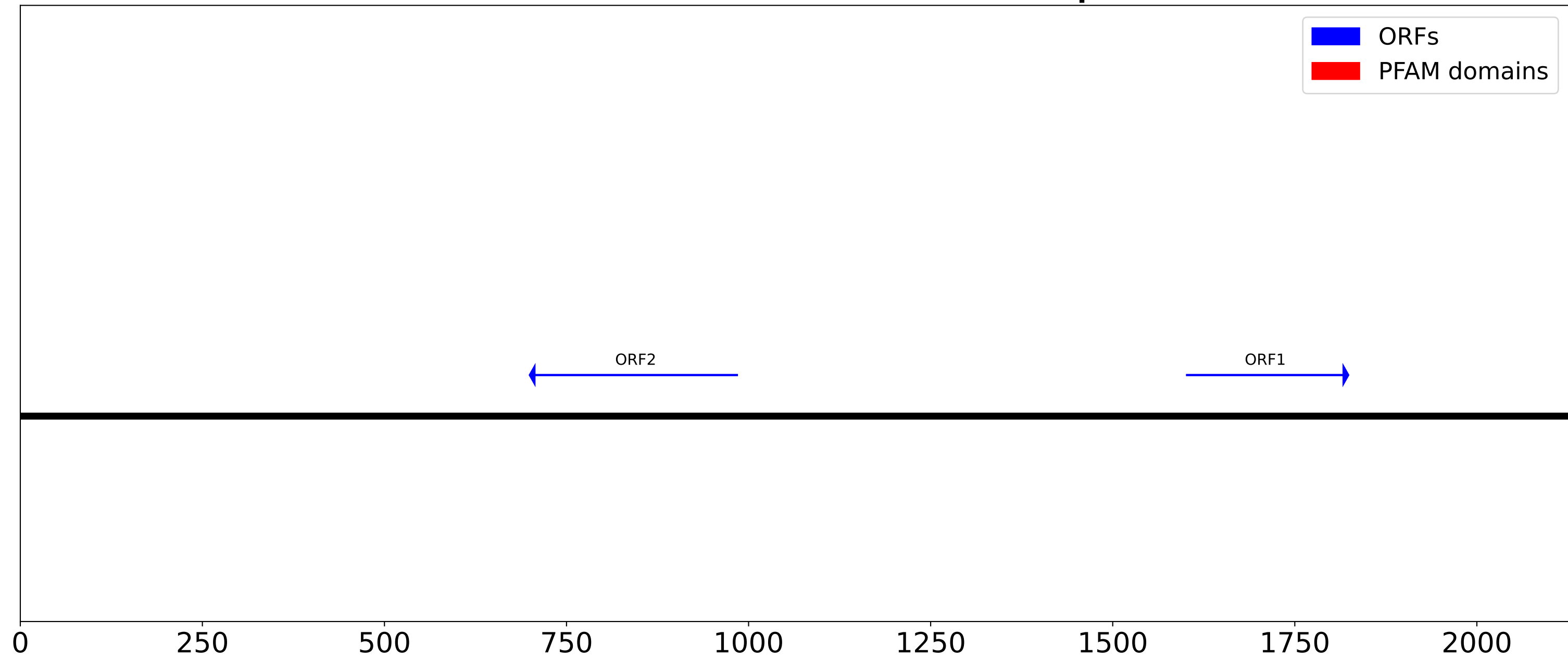
Before TEtrimmer 1248 bp

TE: rnd_5_family_6466

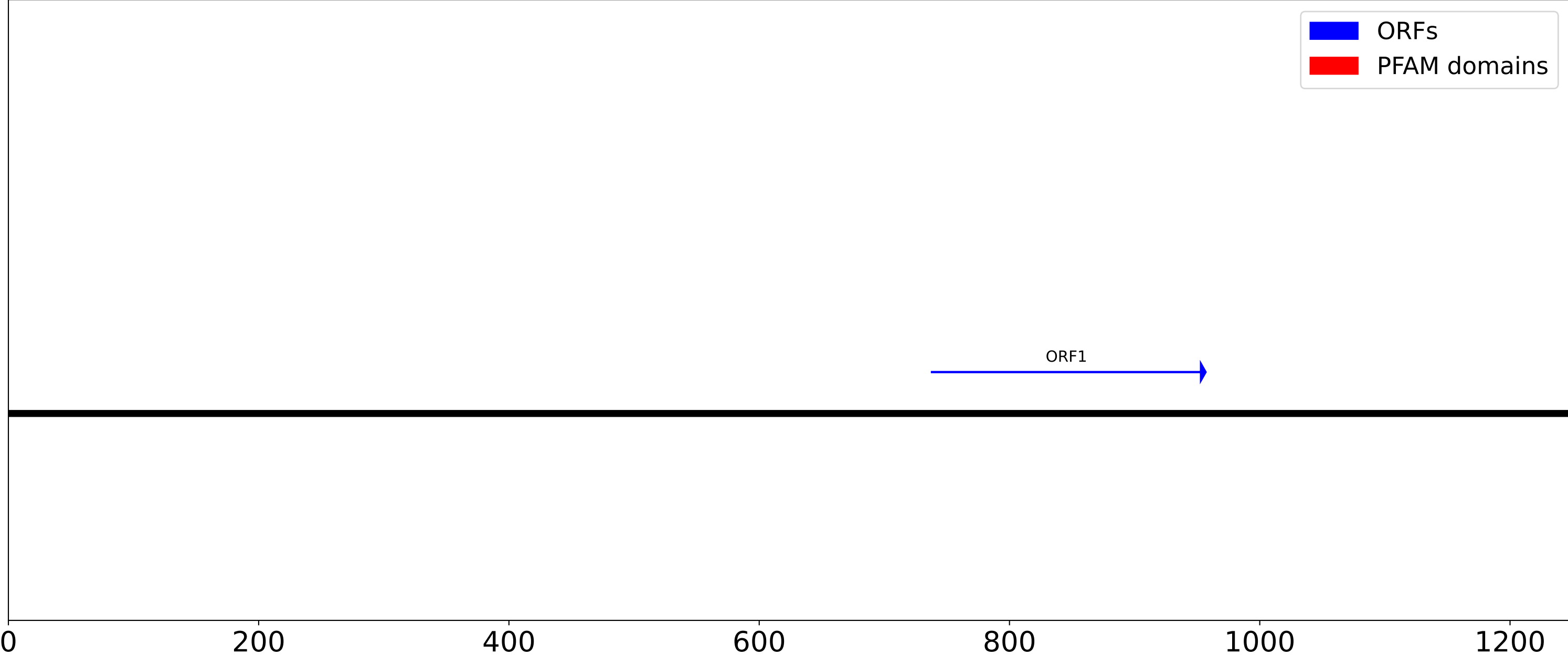
size: 1248bp; fragments: 901; full length: 15 (≥ 1123.2 bp)



After TEtrimmer ORF and PFAM domain plot



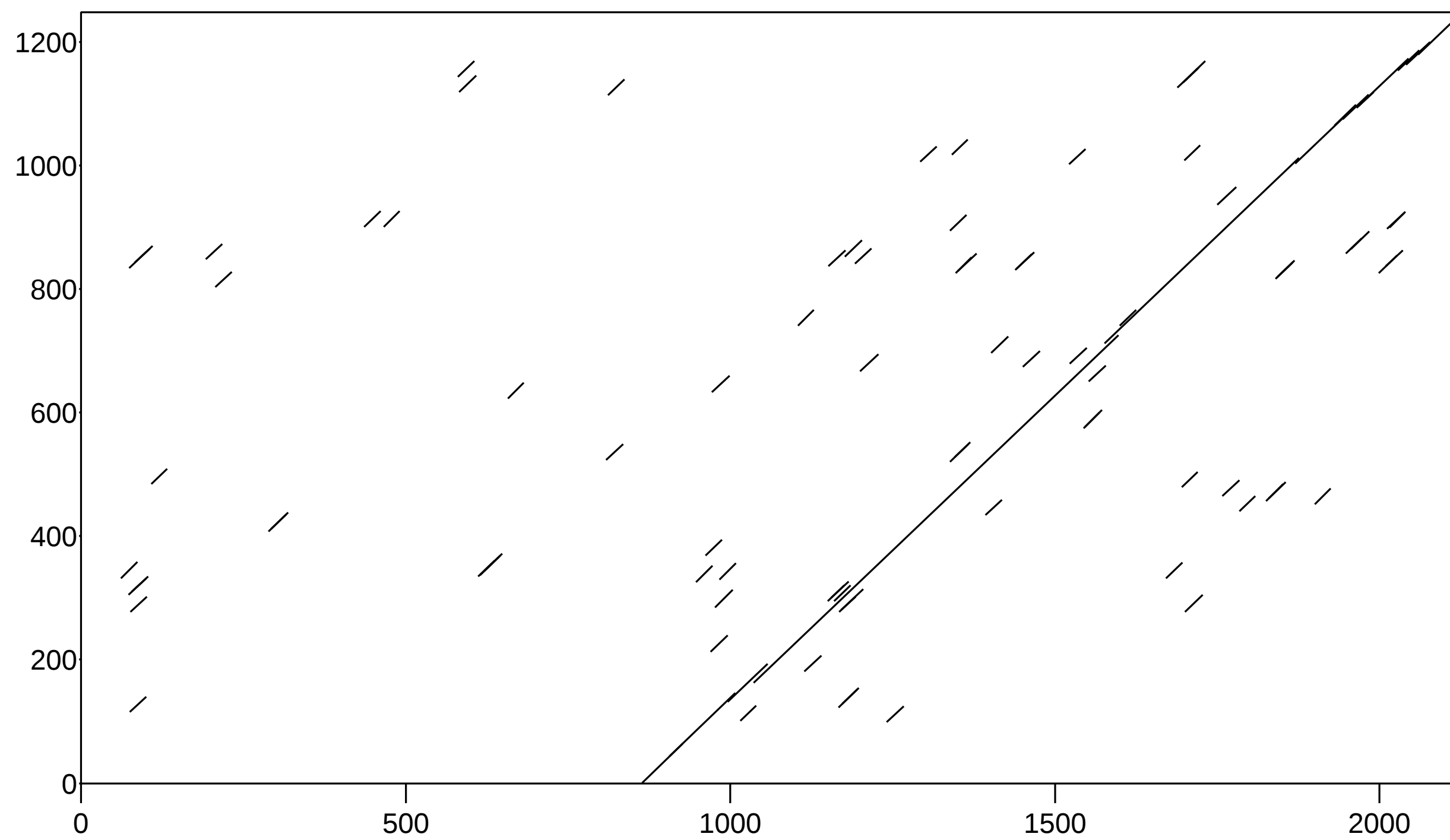
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)