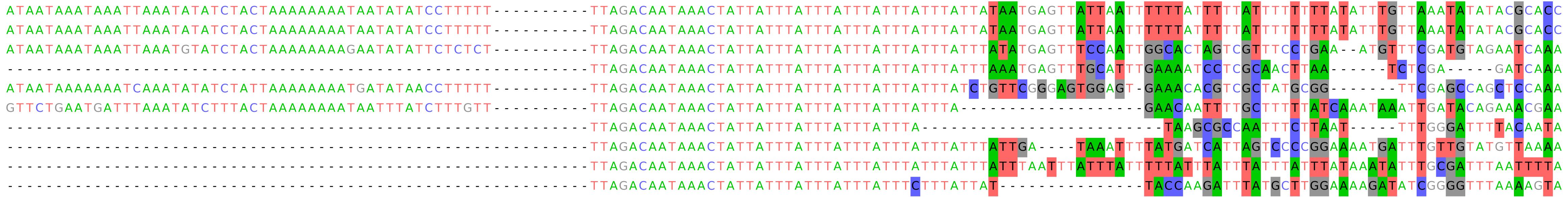
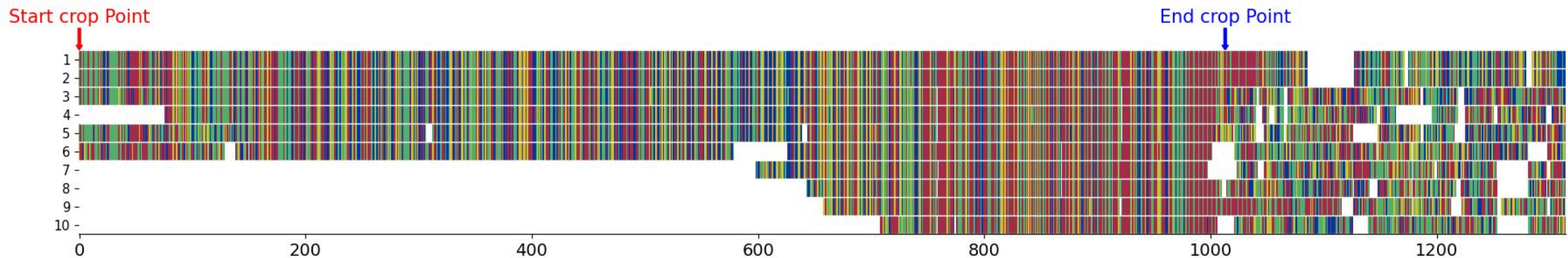


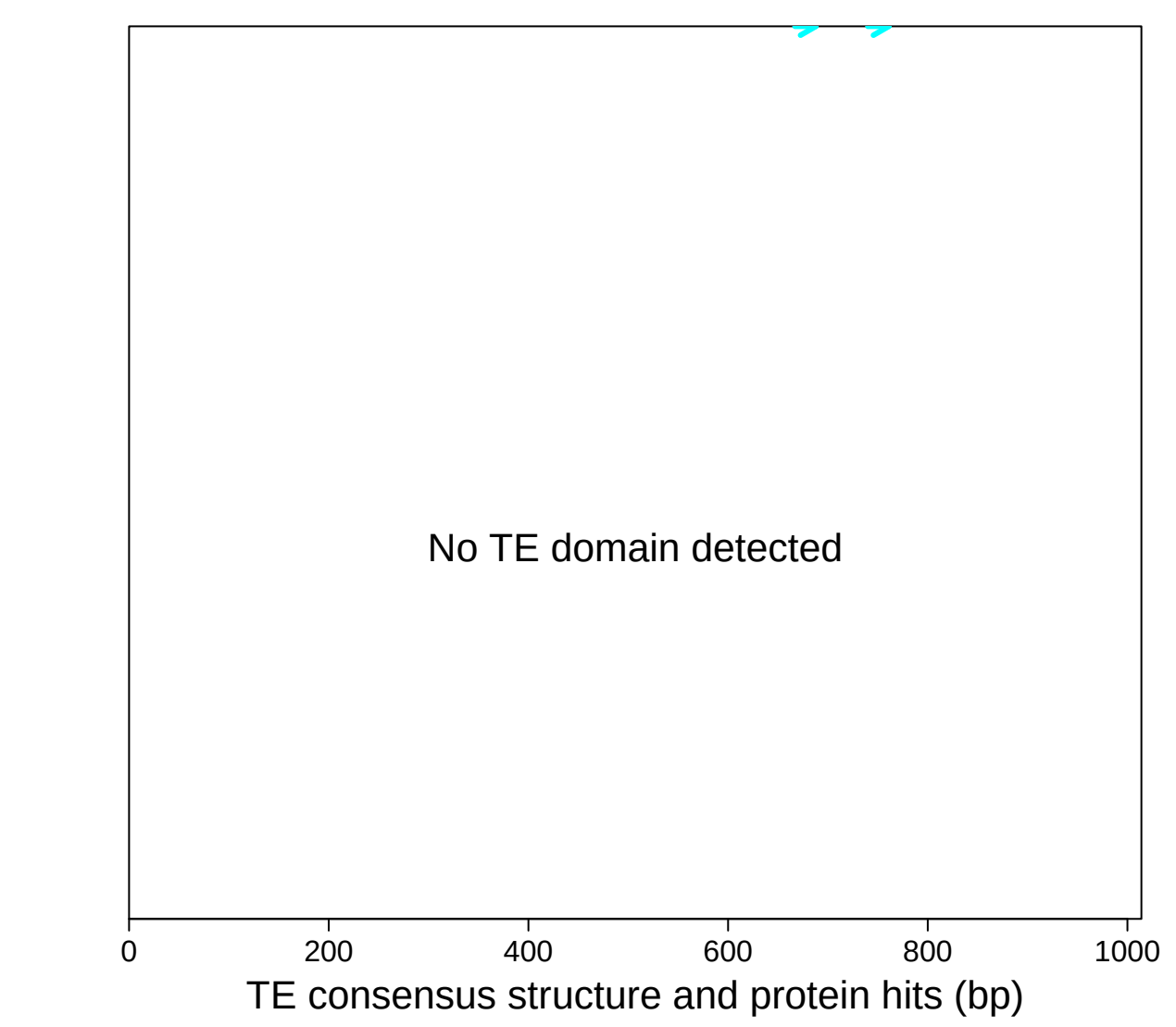
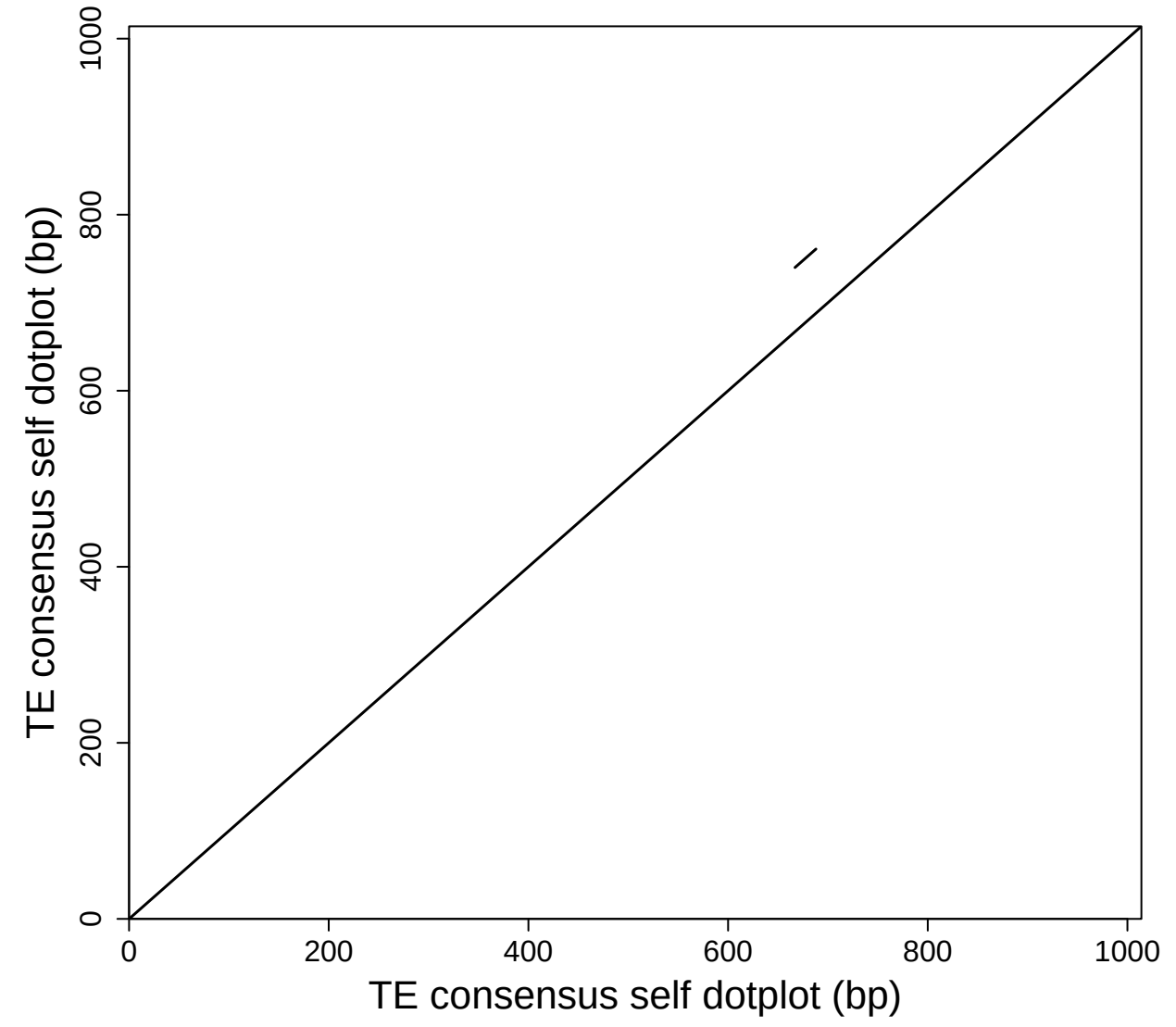
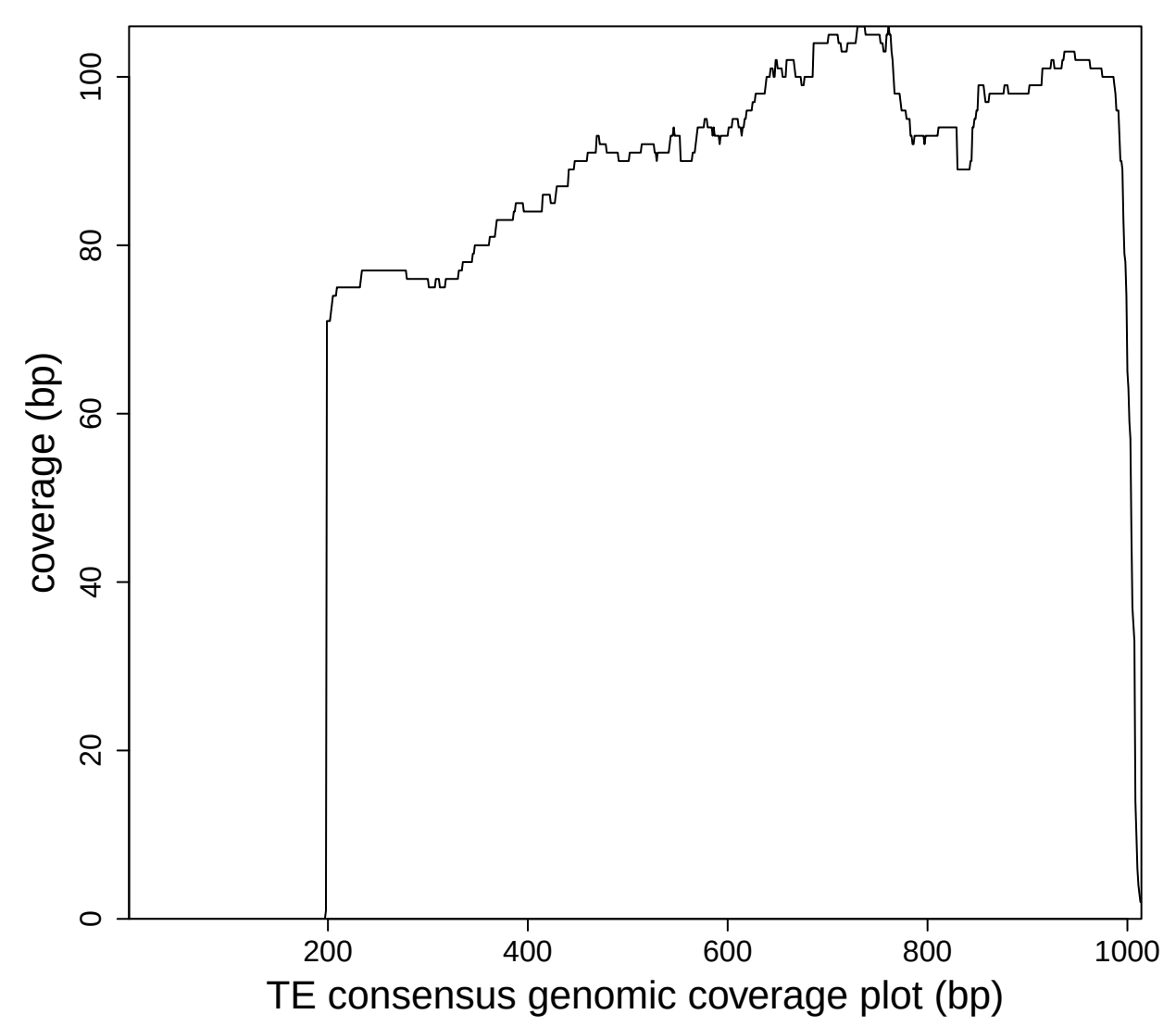
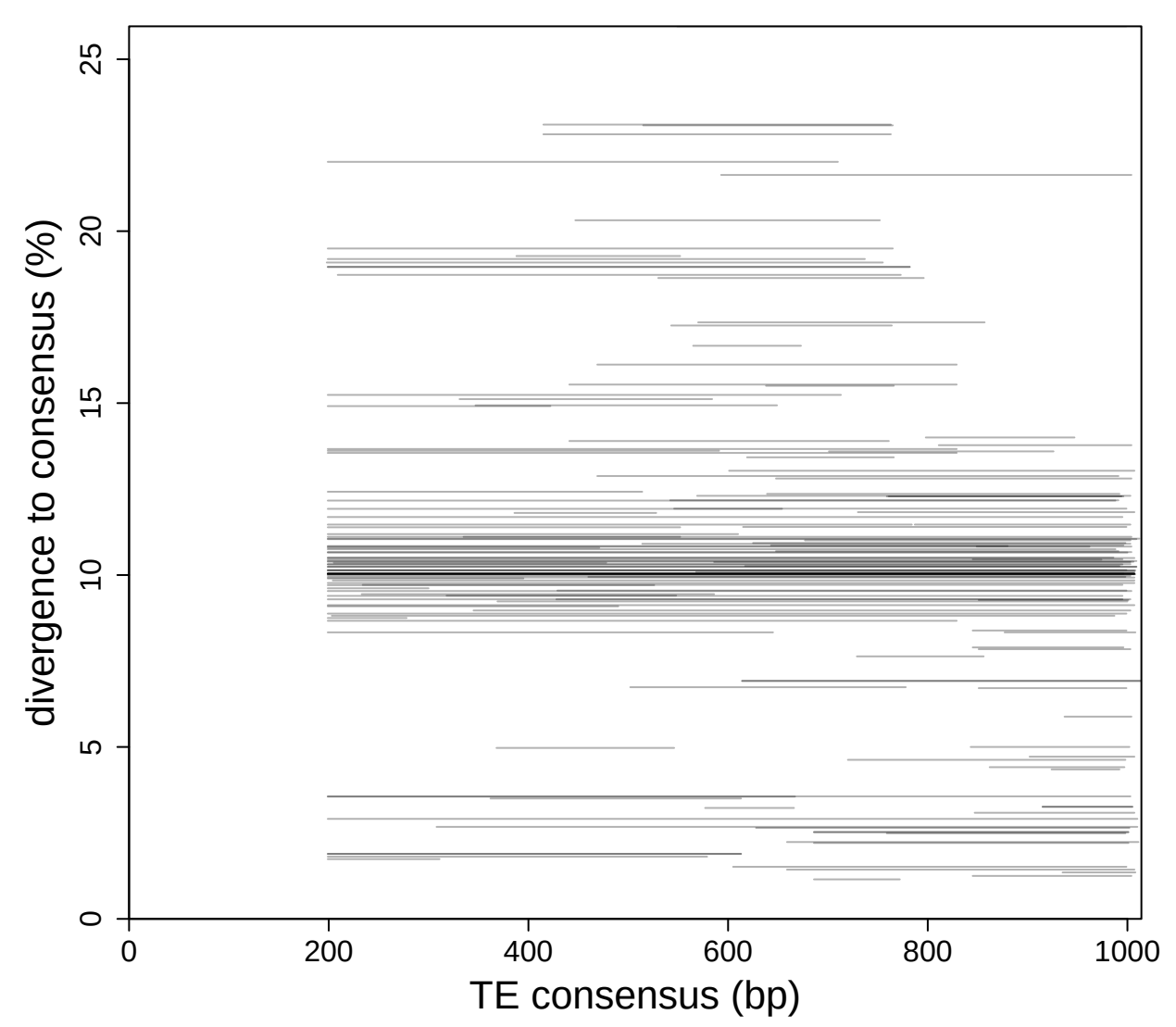
MSA length = 1014





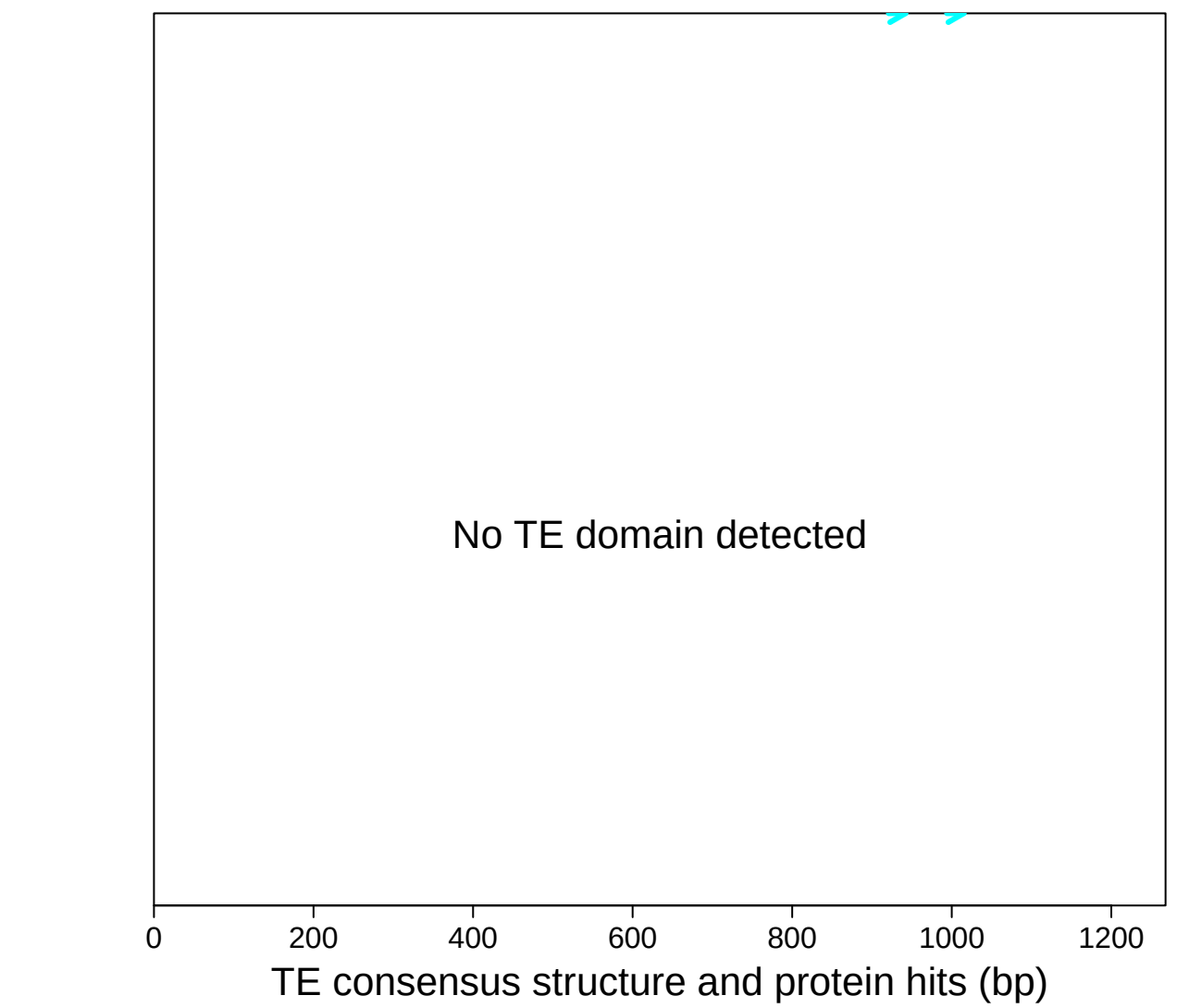
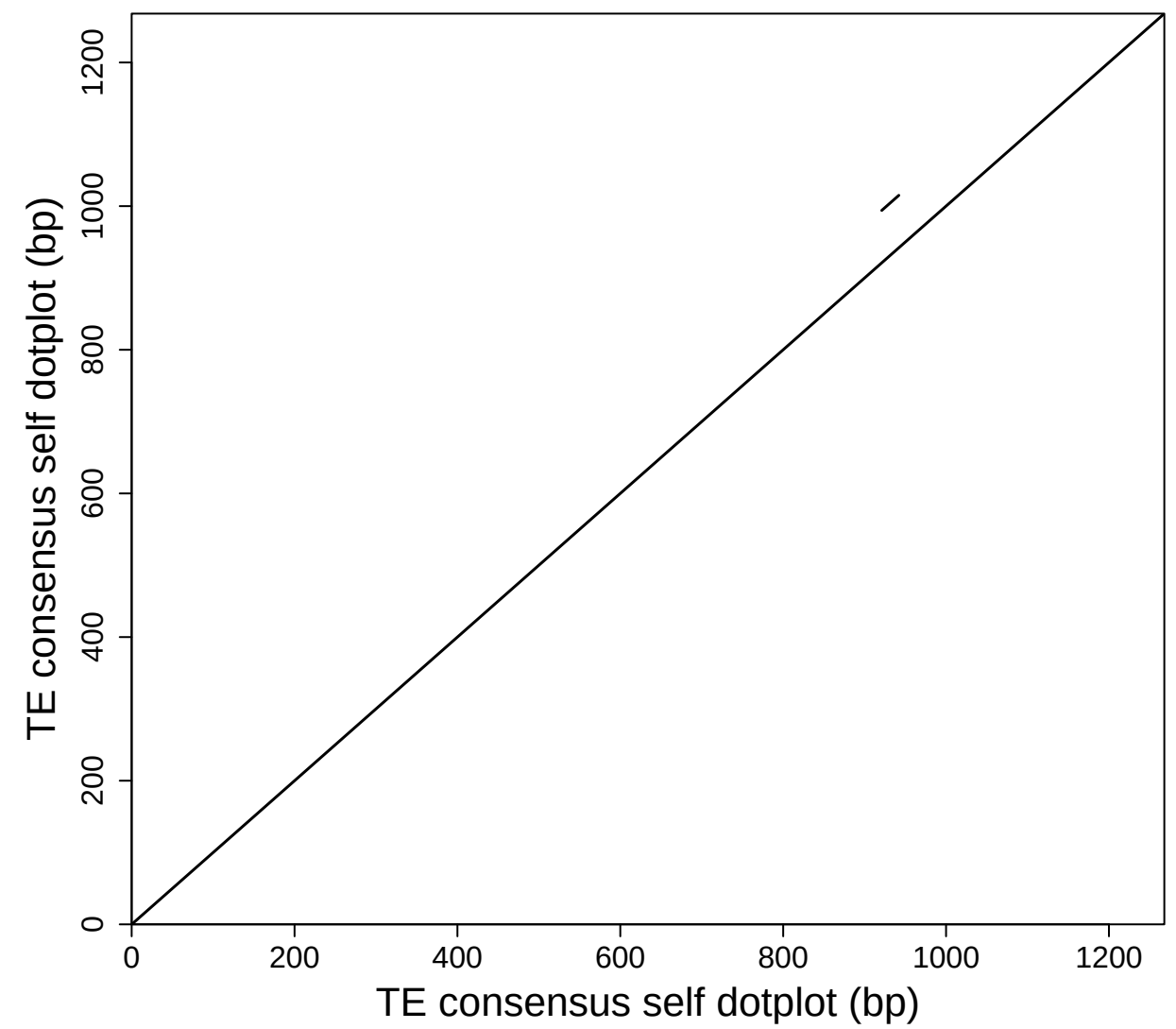
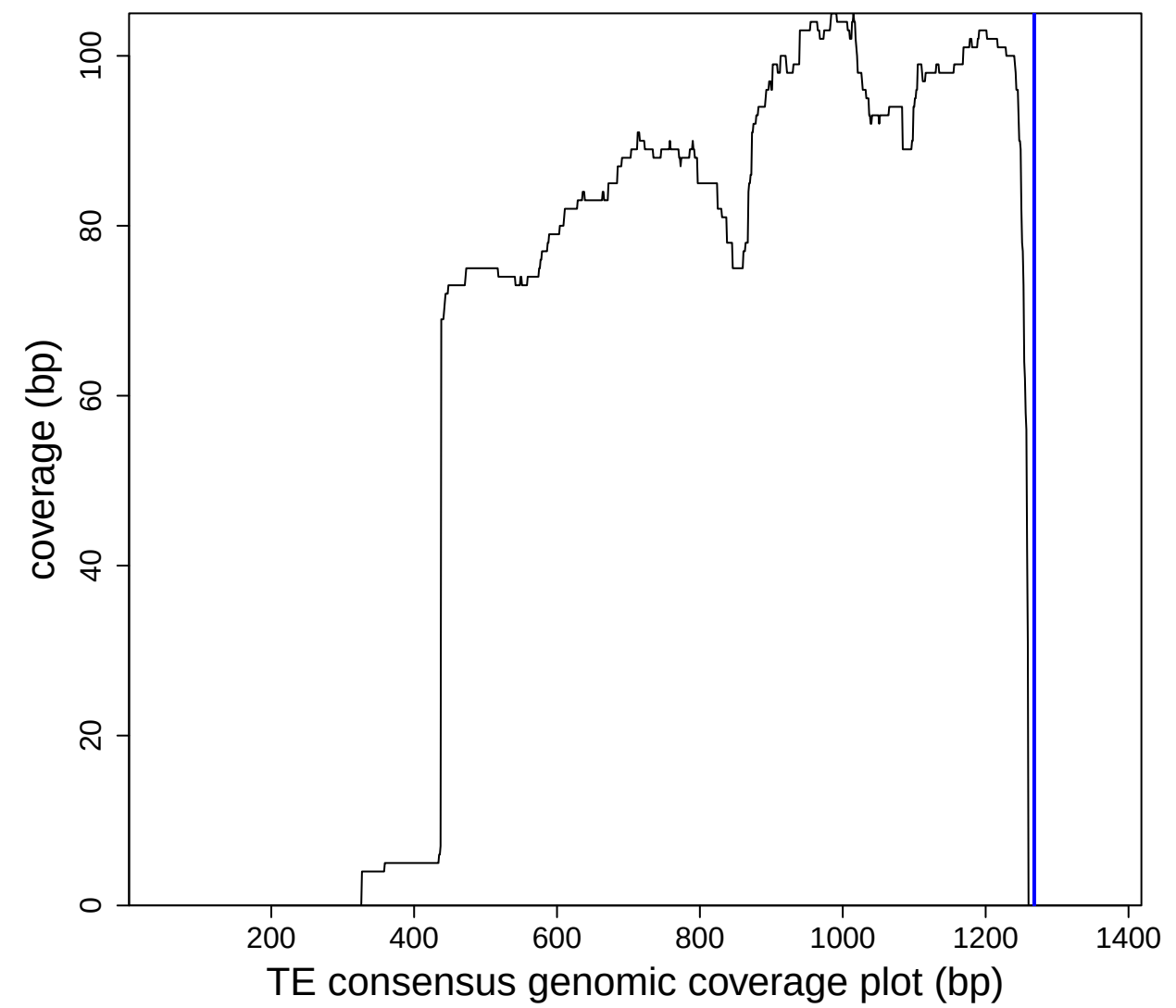
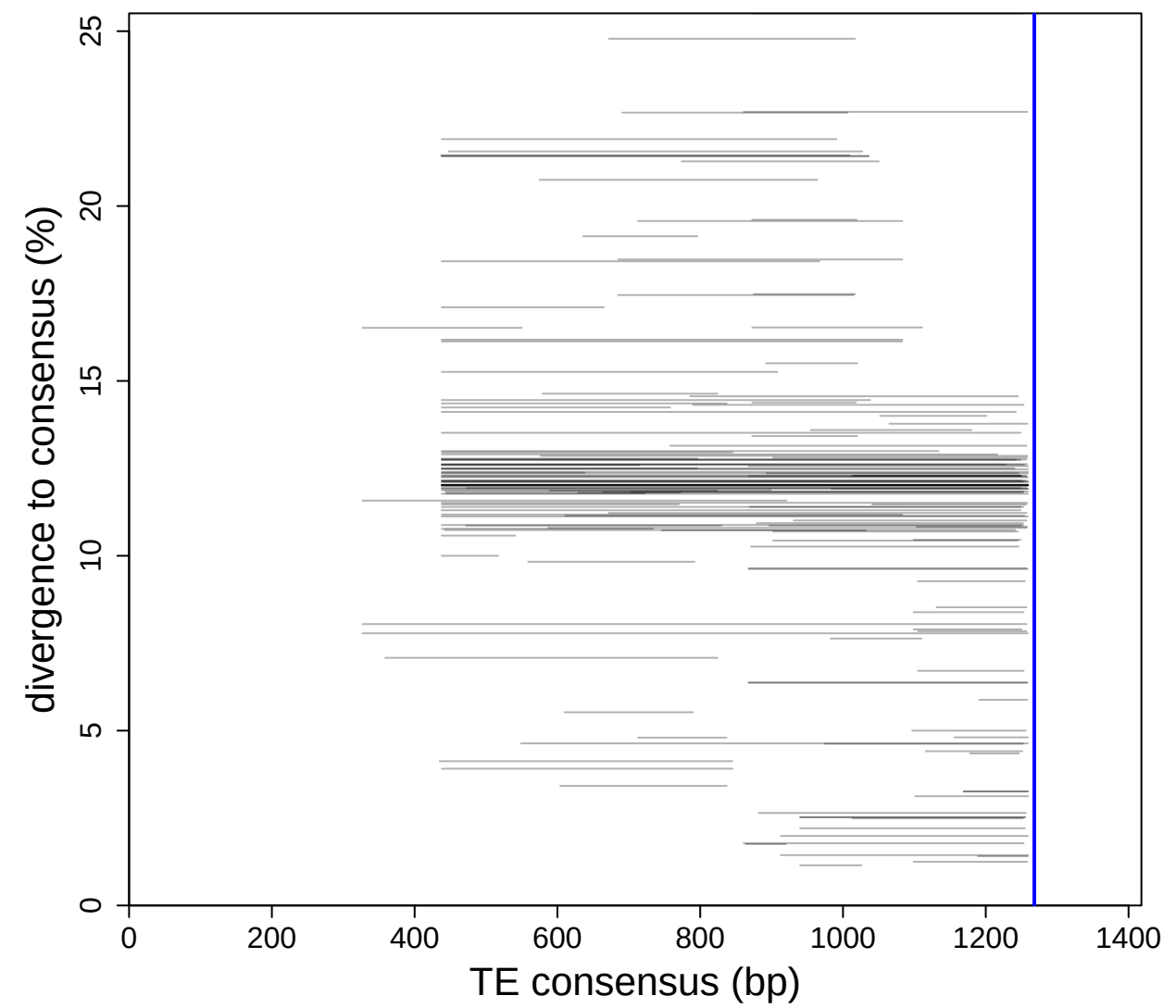
bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 1014bp; fragments: 162; full length: 0 (>=912.6bp)

After TEtrimmer 1014 bp

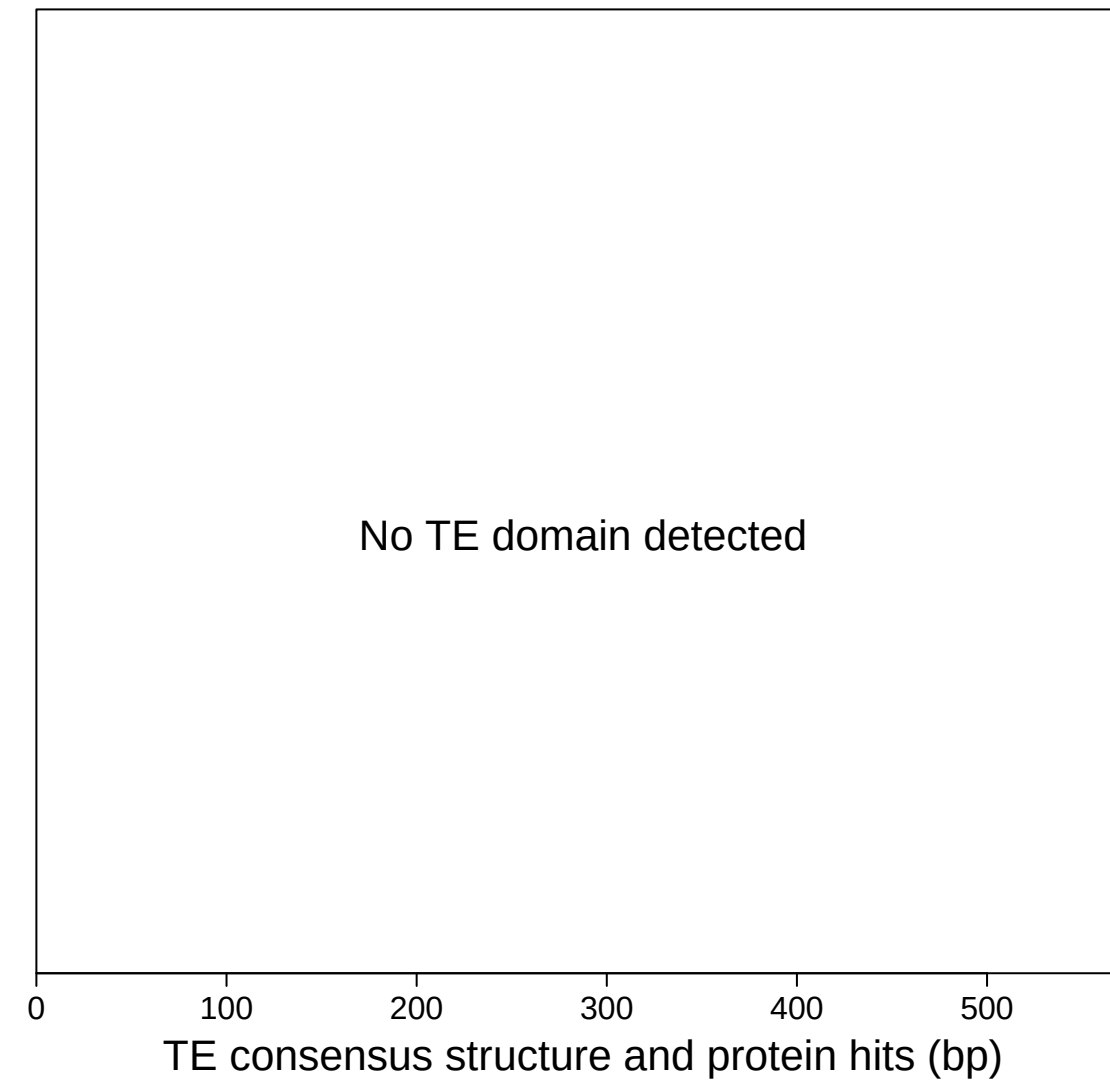
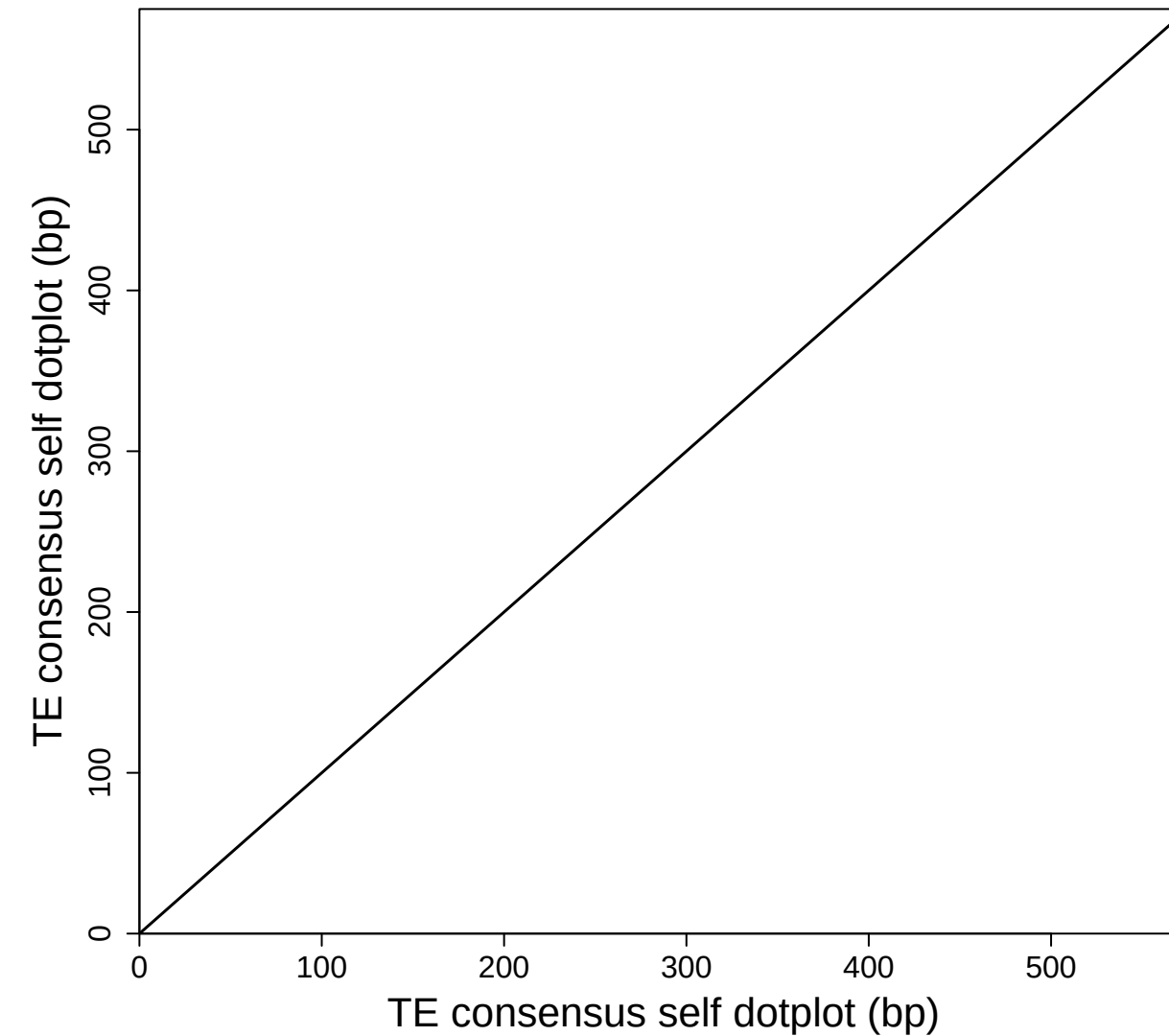
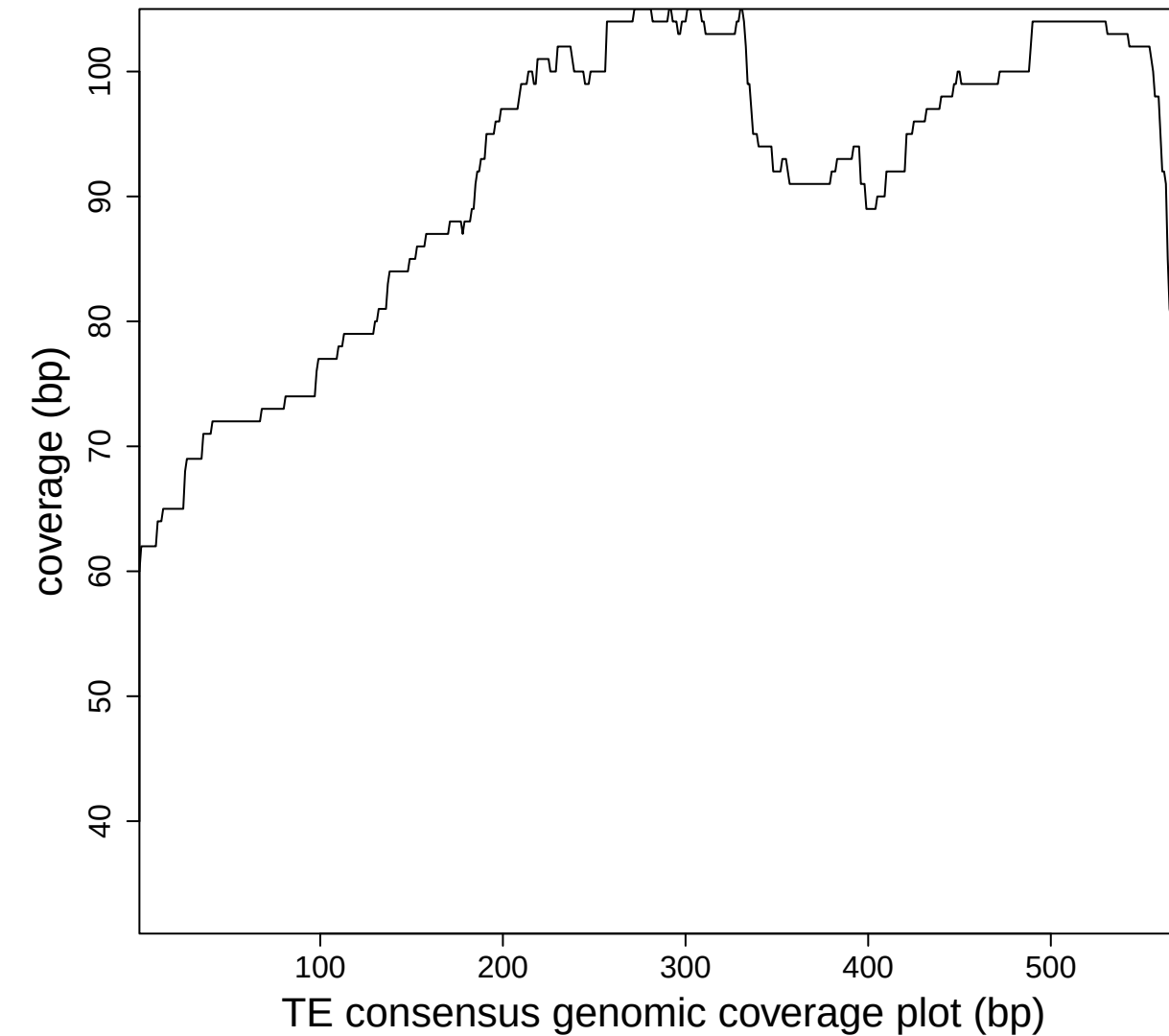
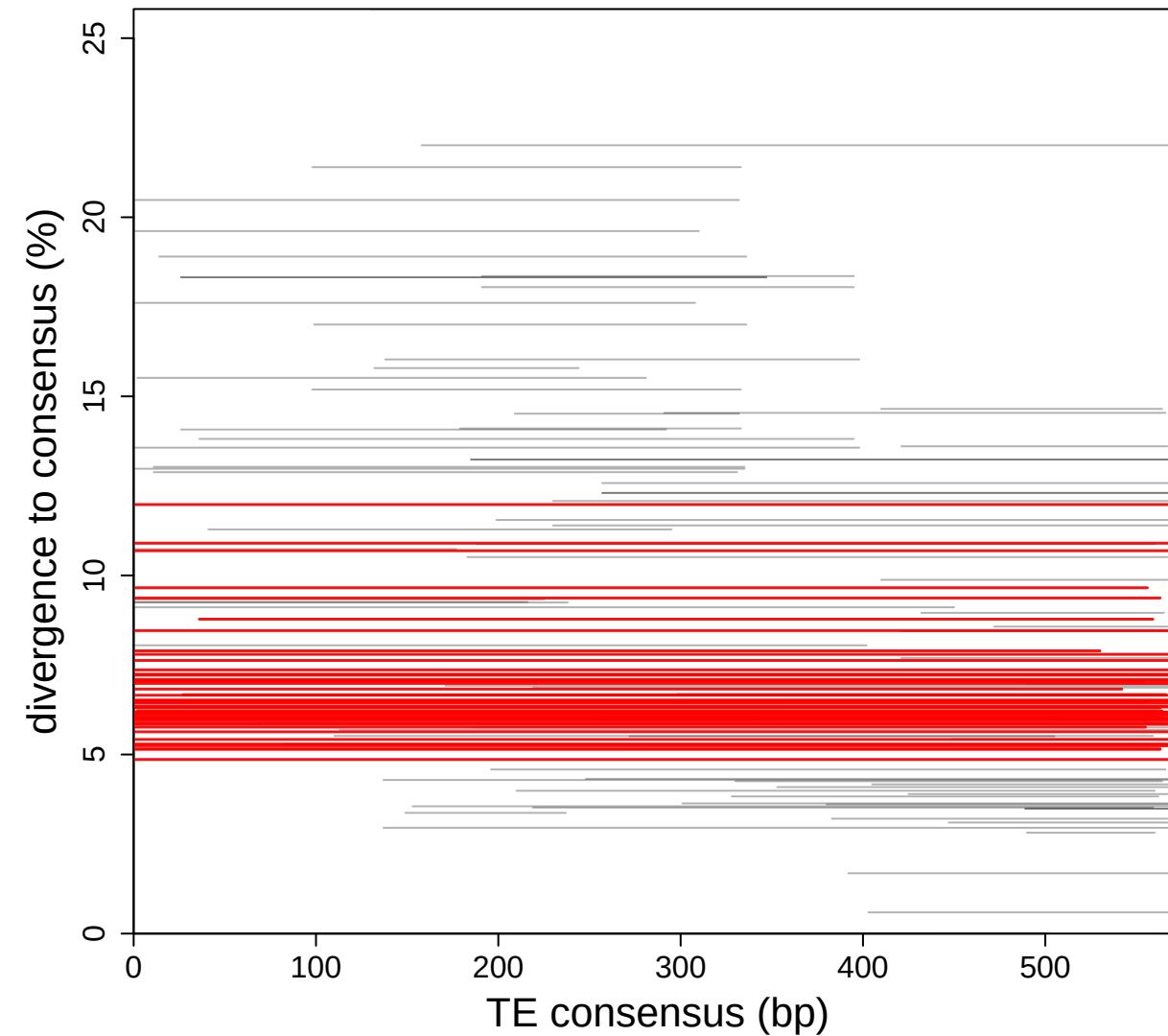


4.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 1418bp; fragments: 161; full length: 0 (>=1276.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

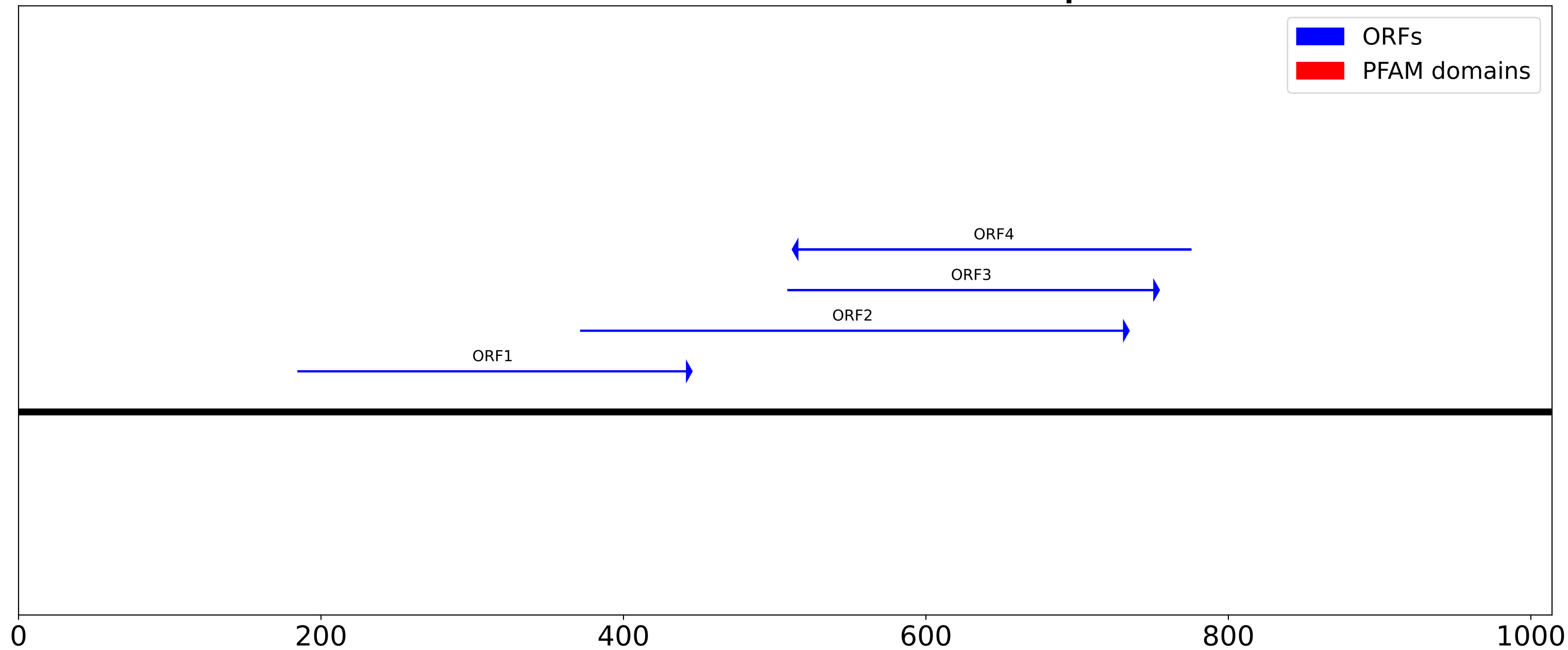


TE: rnd_5_family_9444
size: 575bp; fragments: 138; full length: 51 (≥ 517.5 bp)

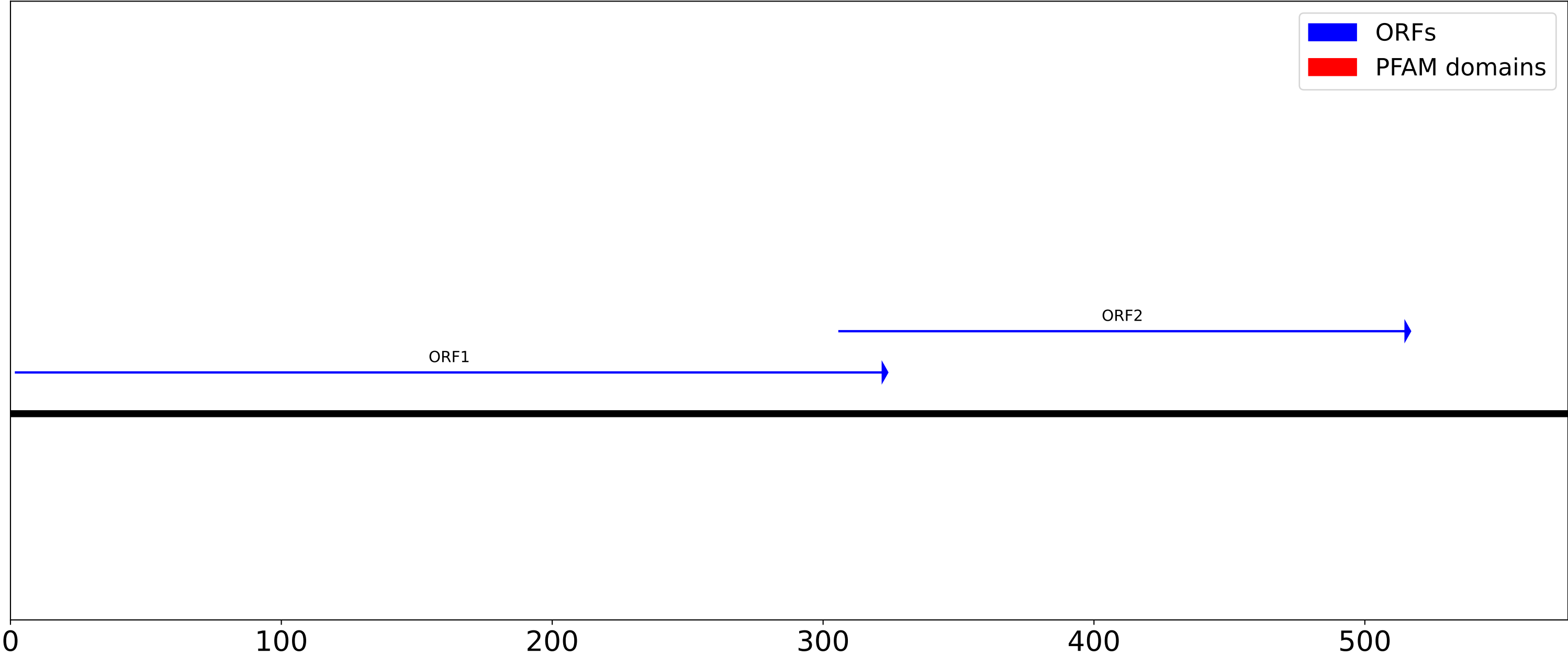


Before TEtrimmer 575 bp

After TEtrimmer ORF and PFAM domain plot



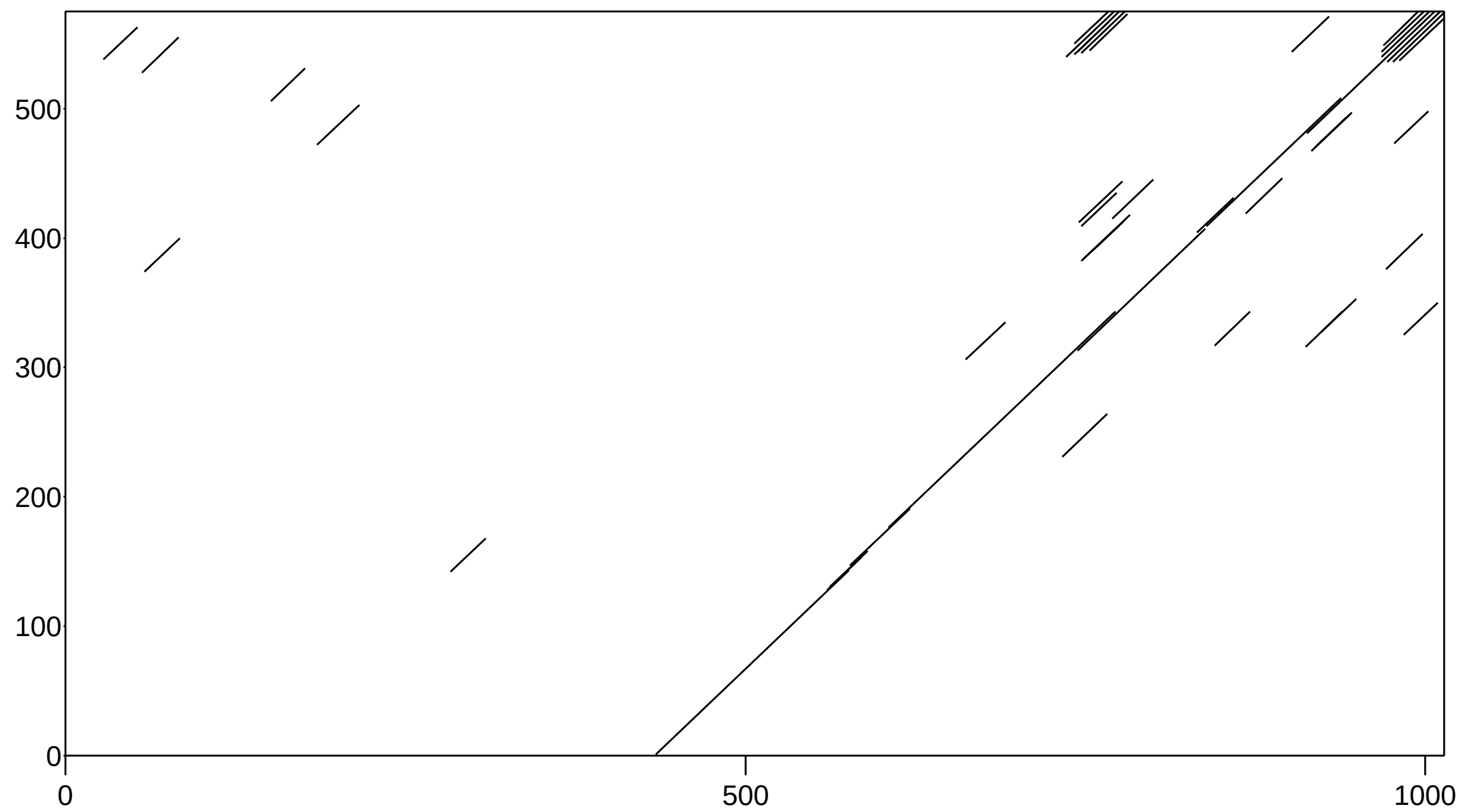
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)