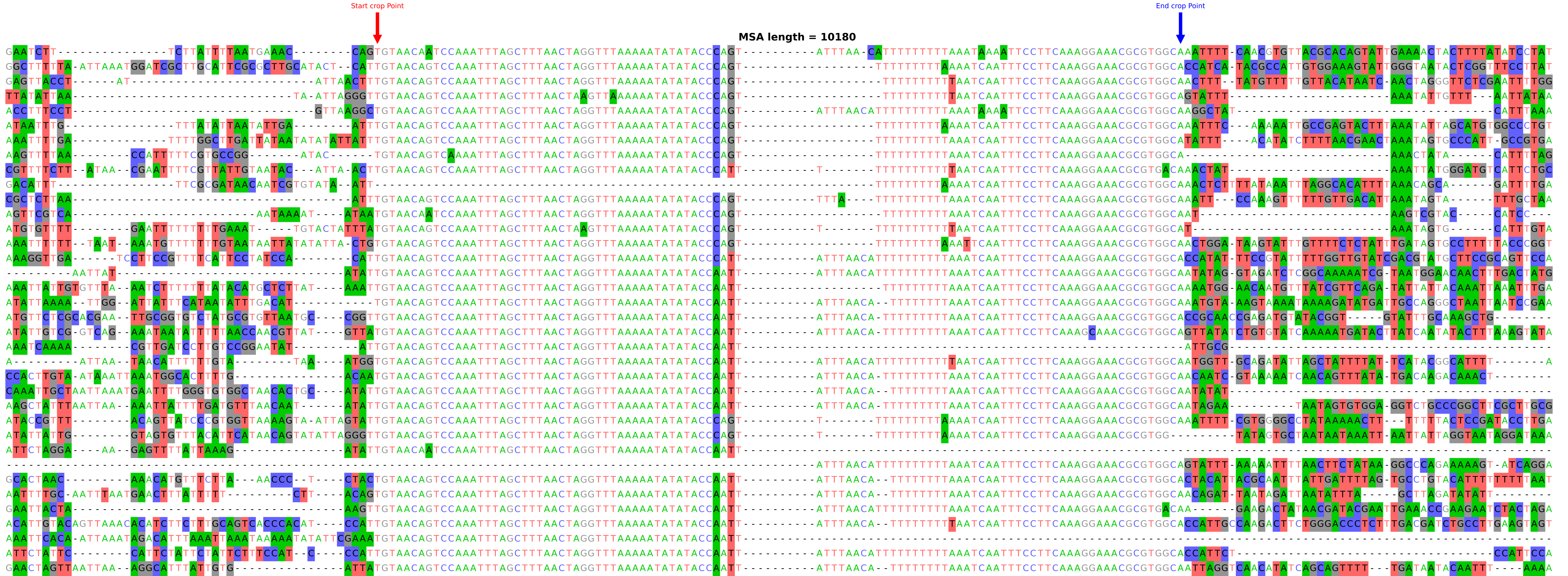
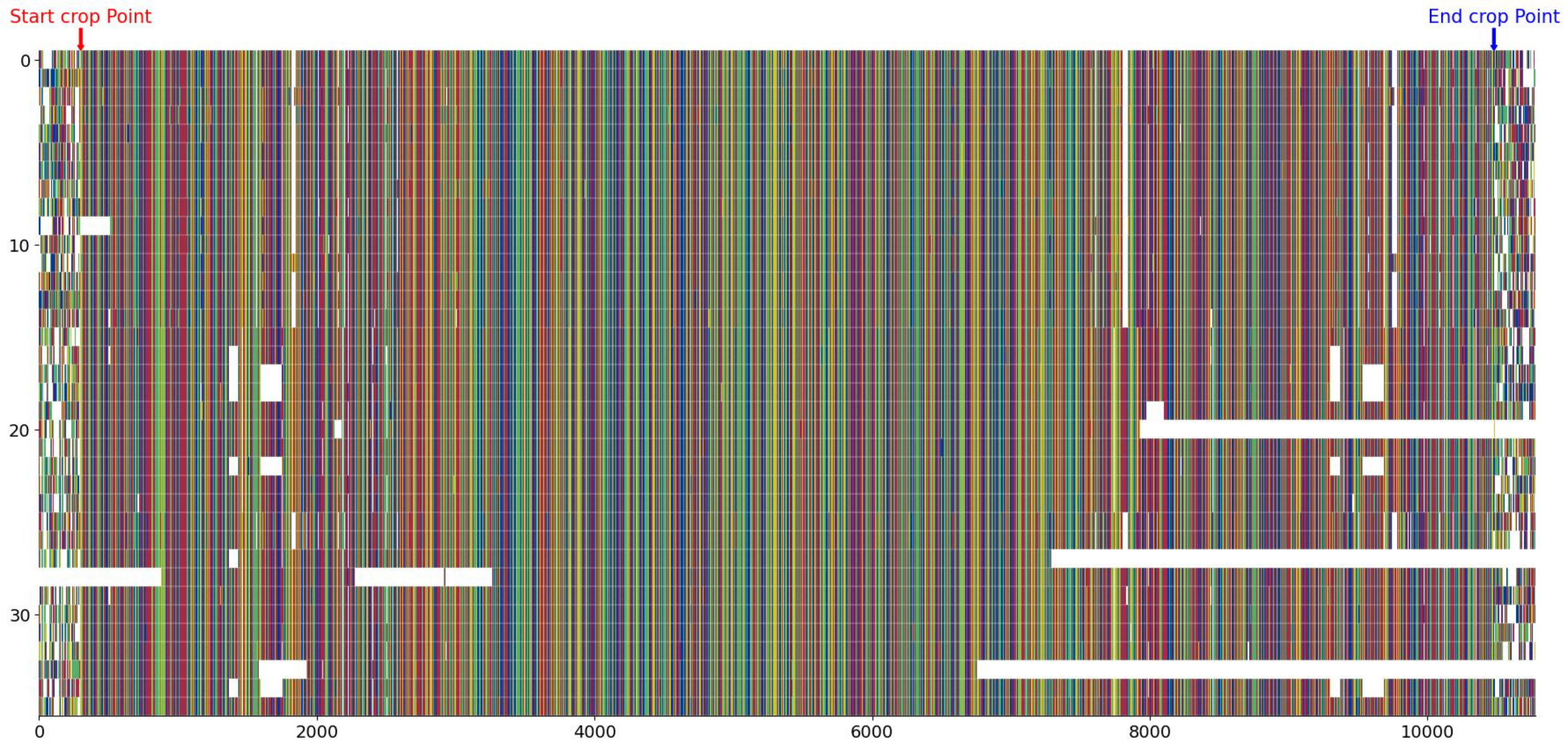
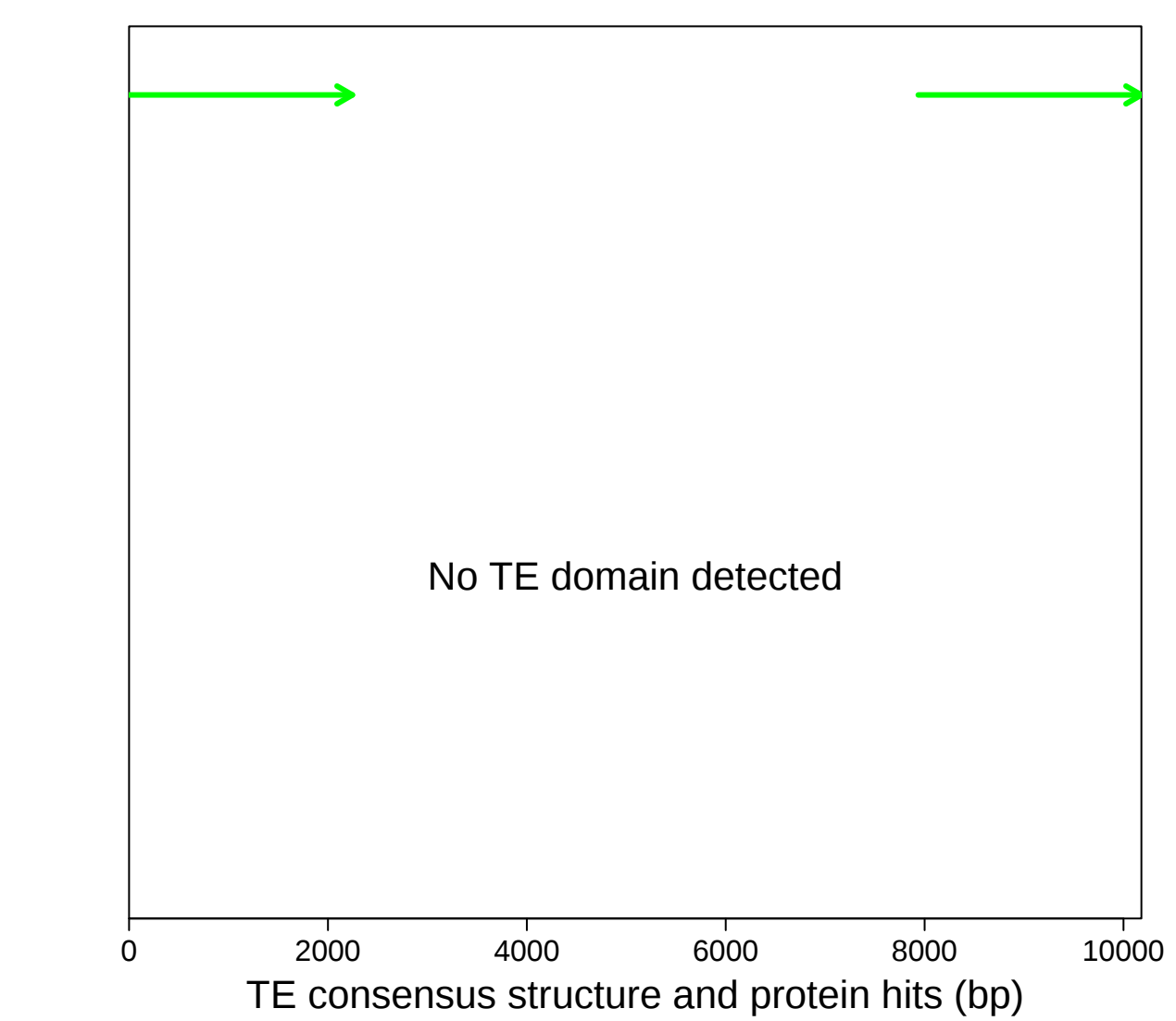
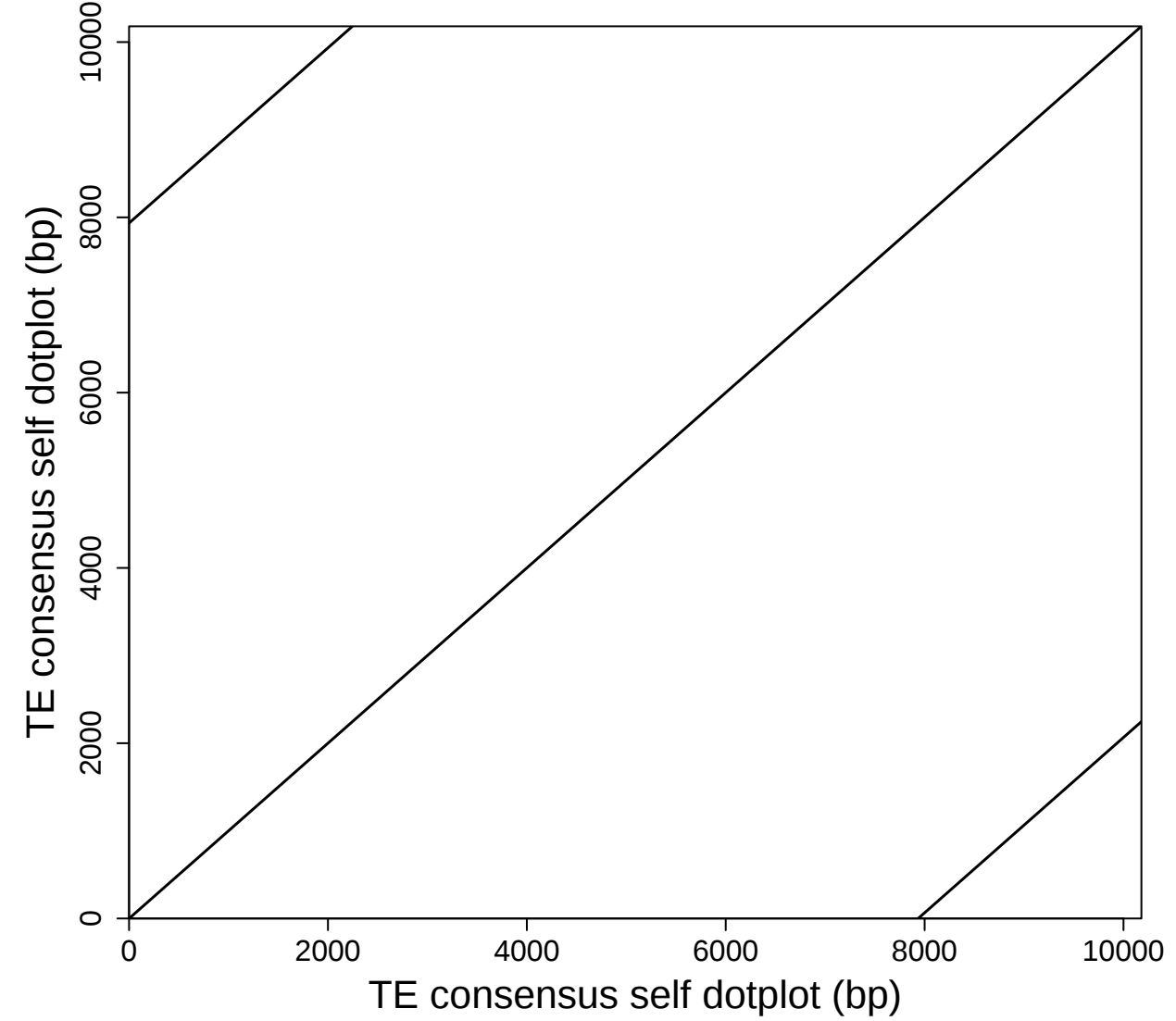
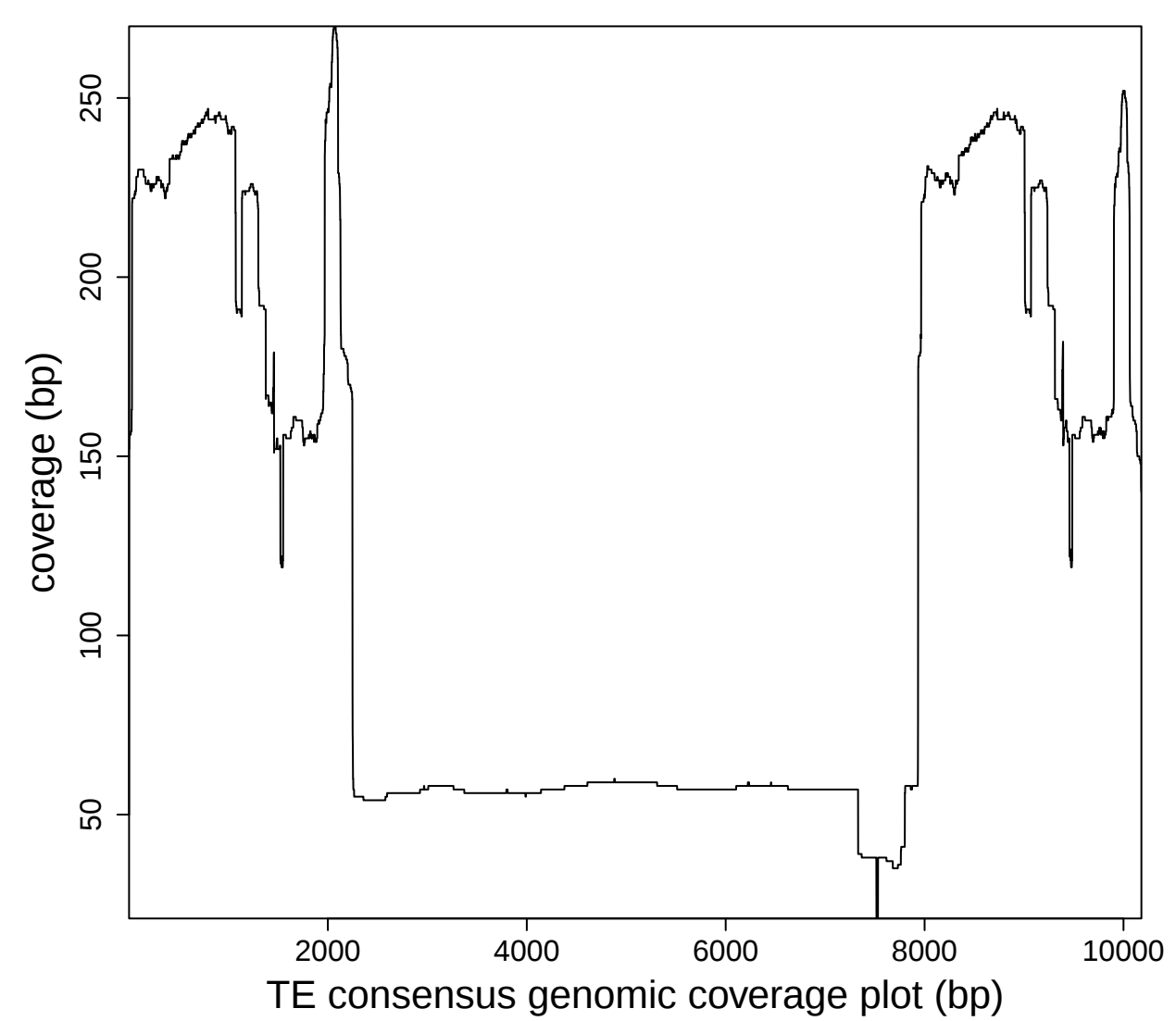
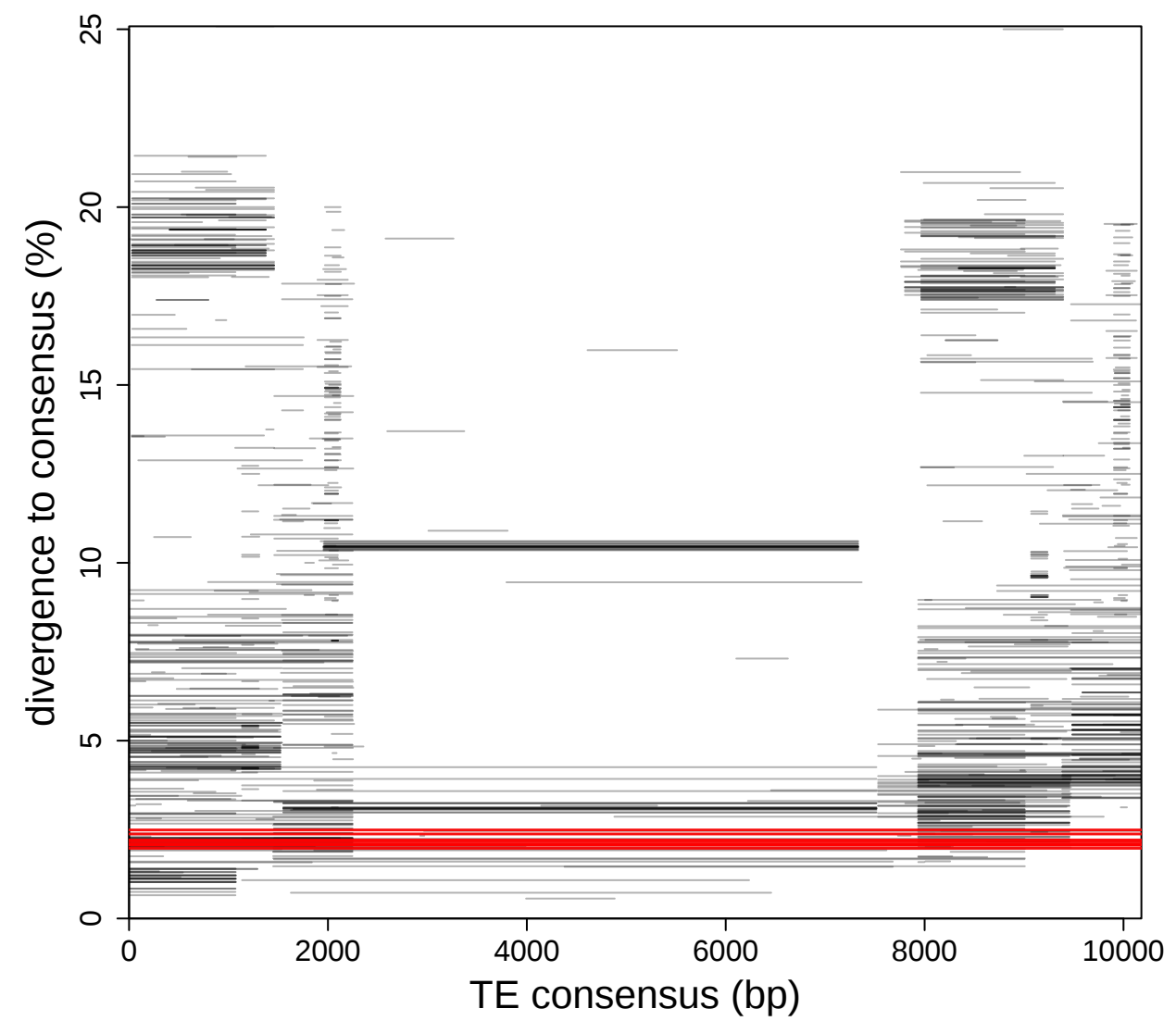






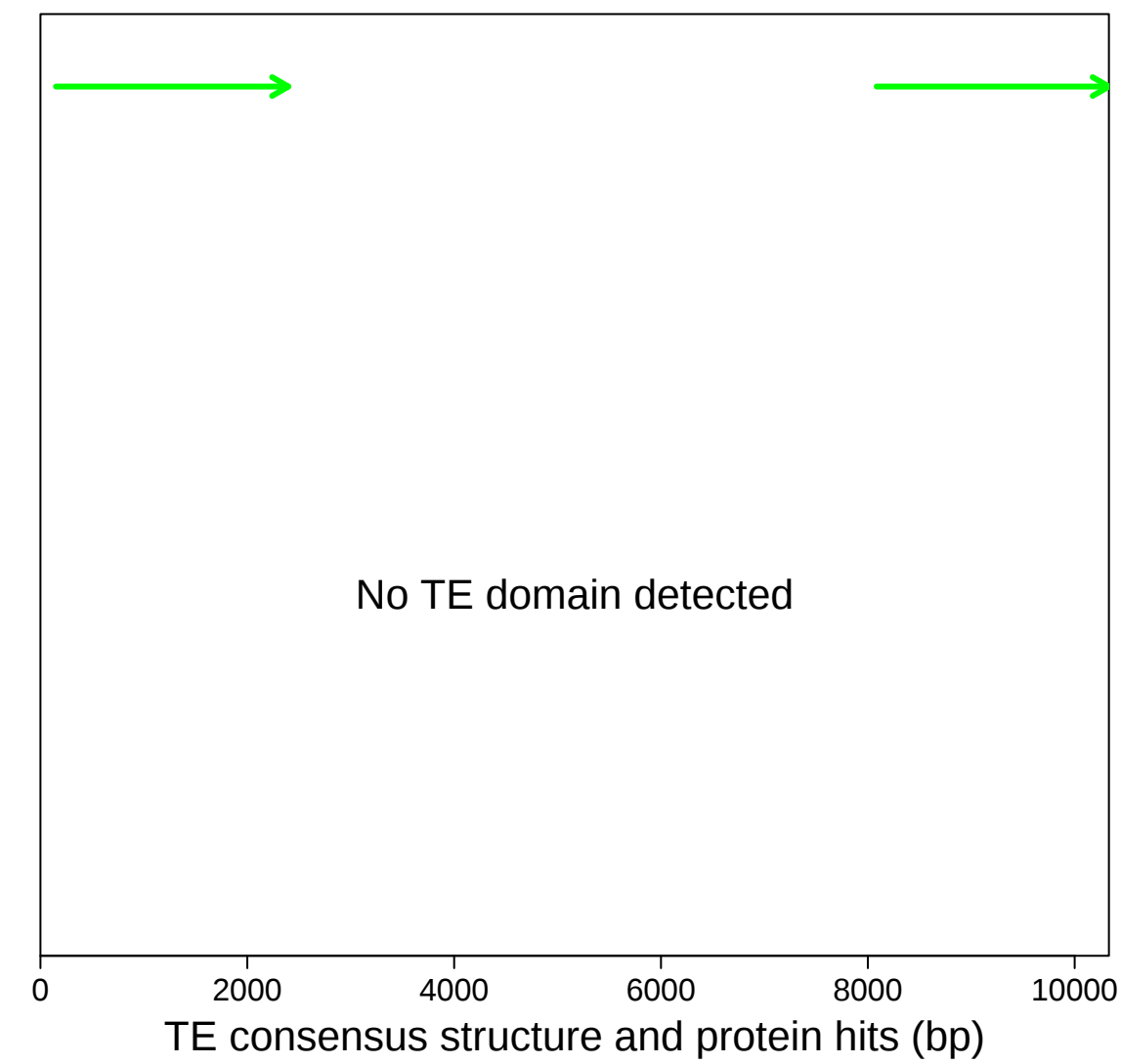
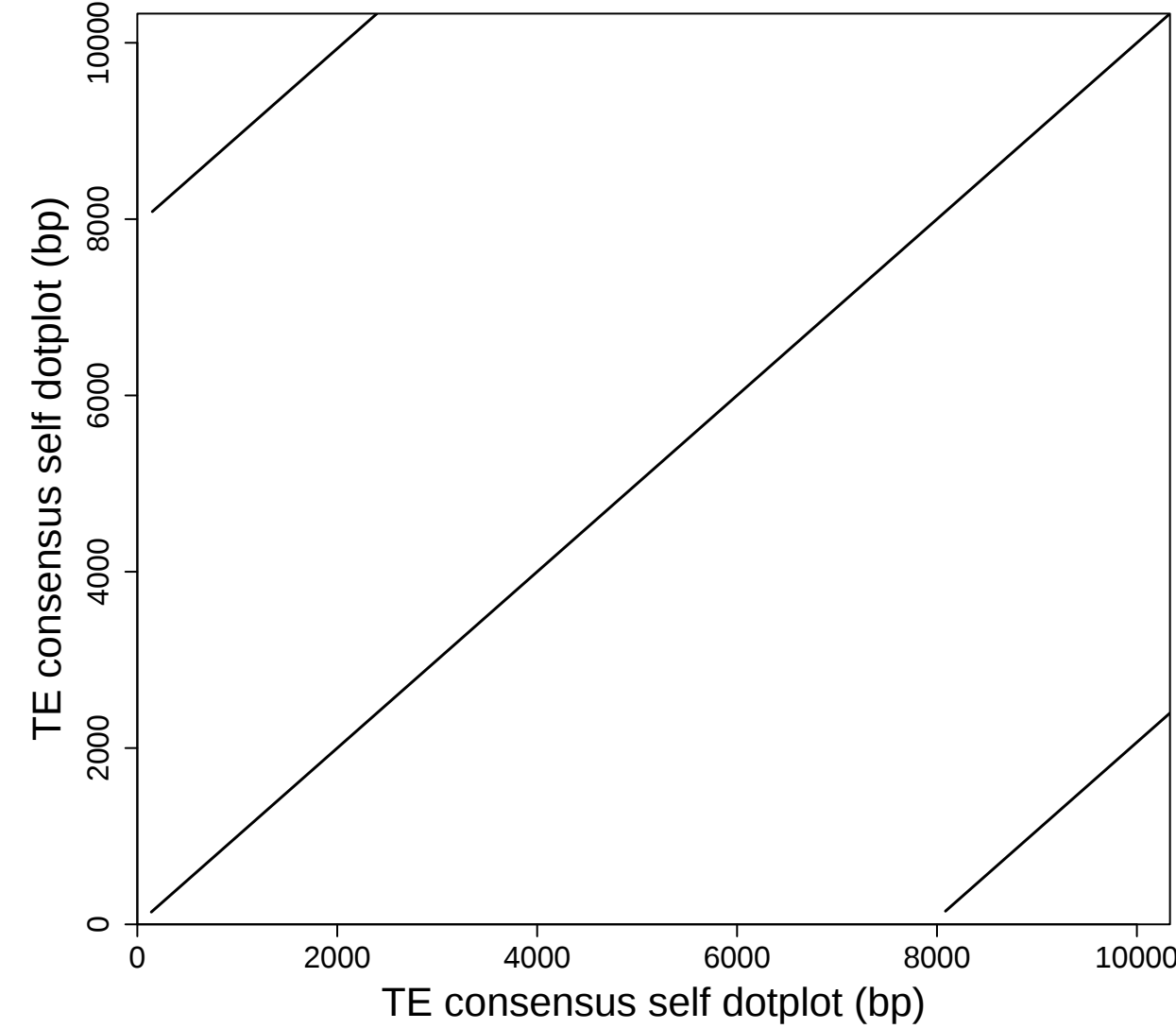
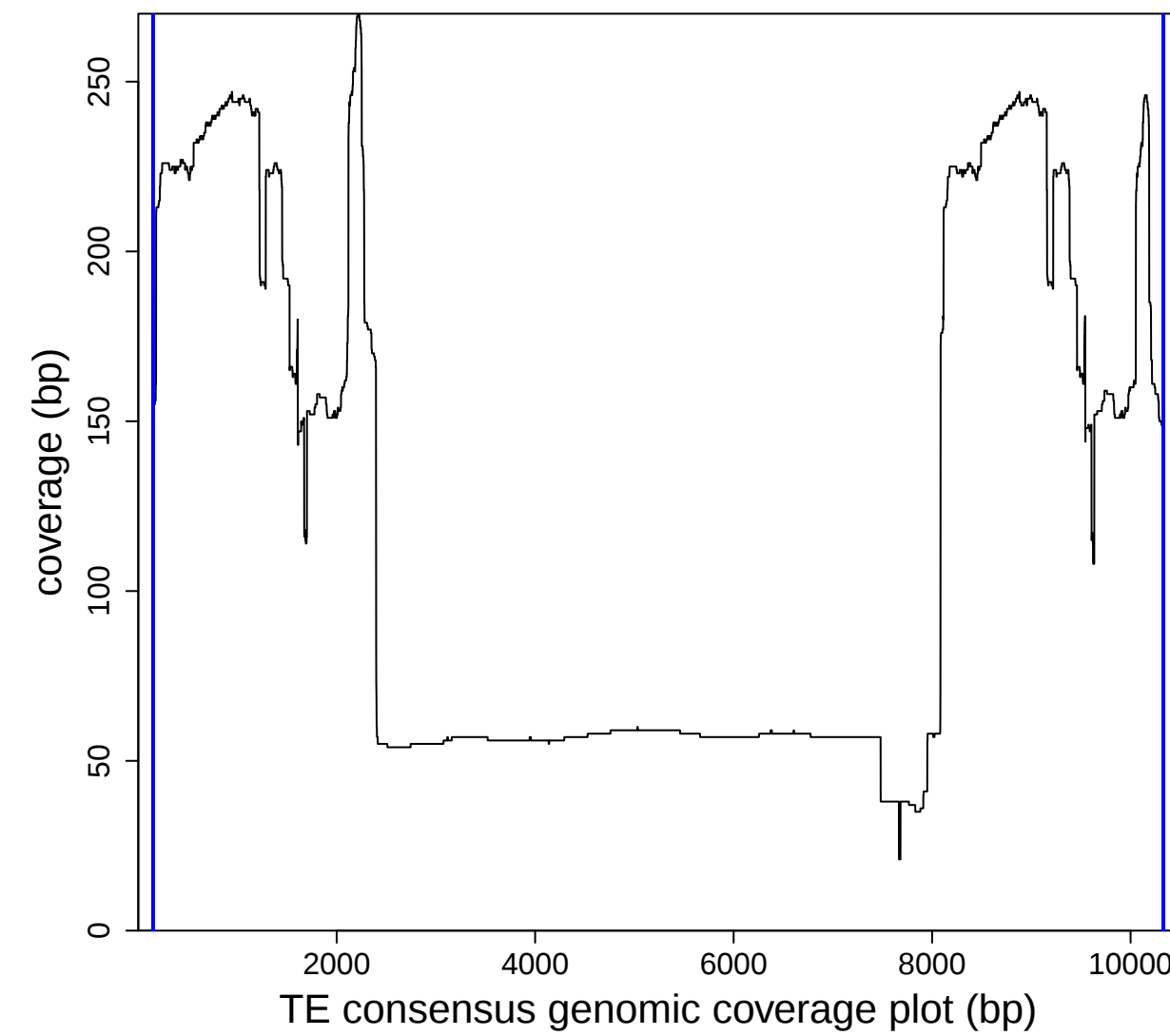
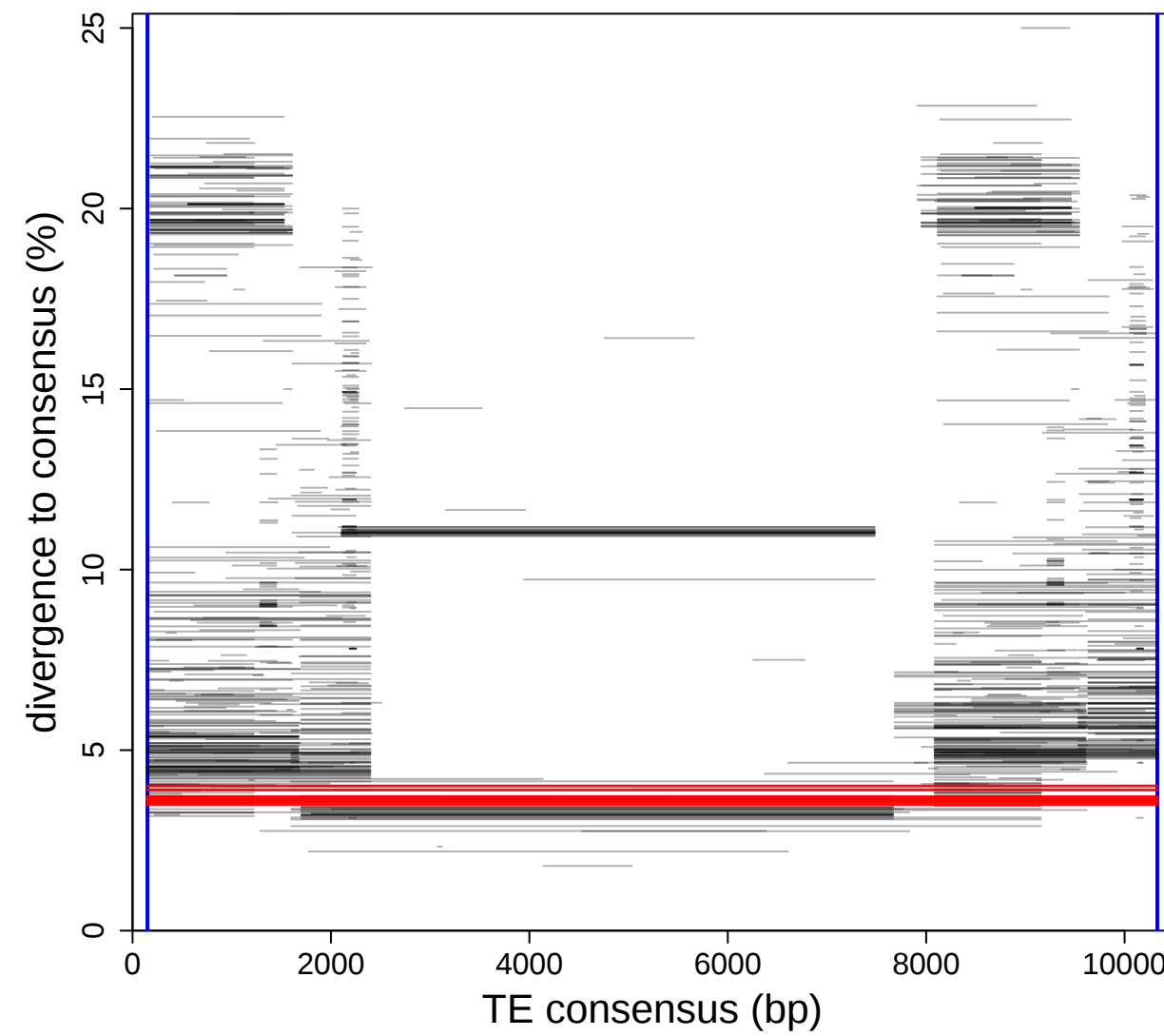
ed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 10180bp; fragments: 1071; full length: 8 (>=9162bp)

After TEtrimmer 10180 bp

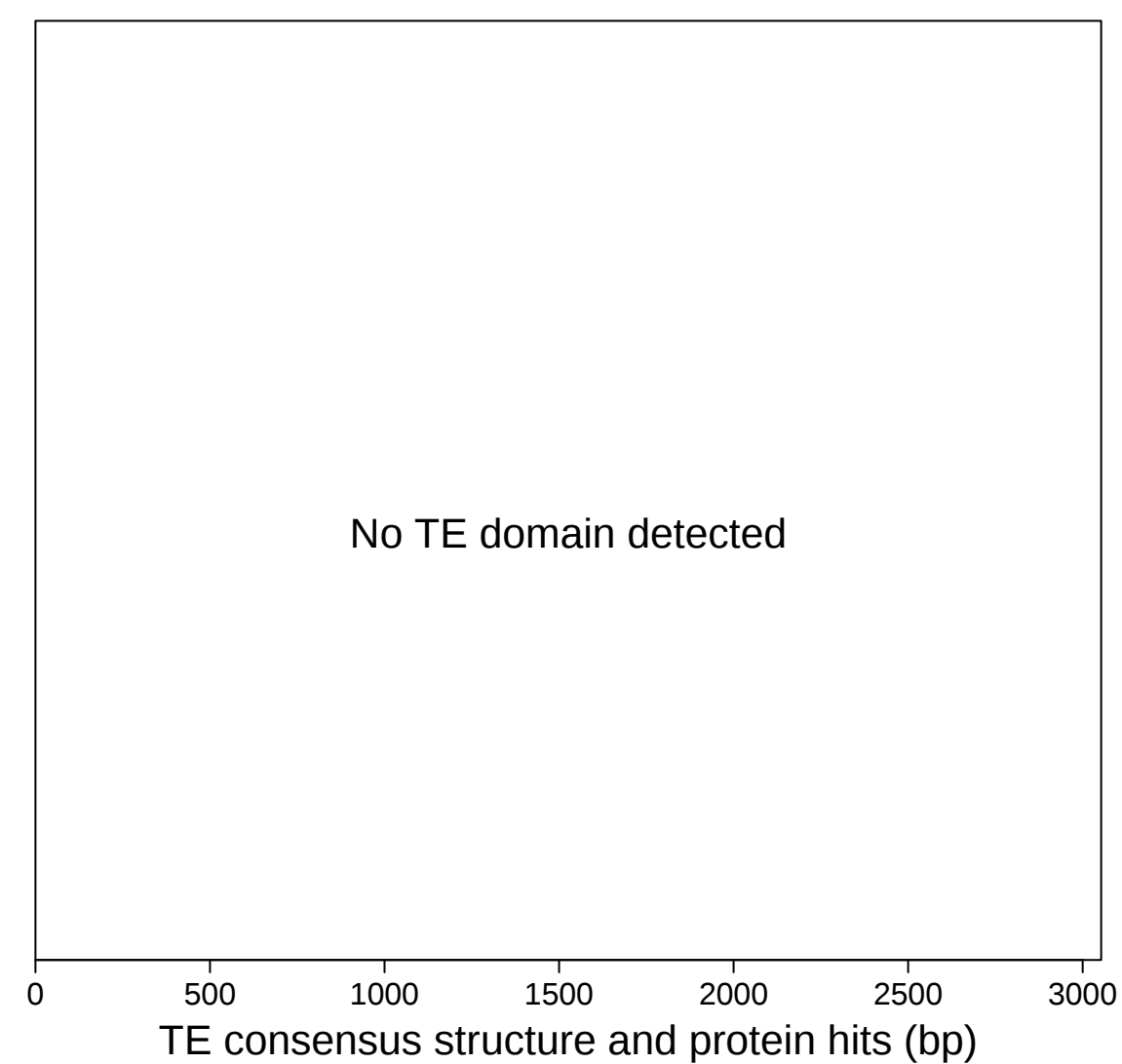
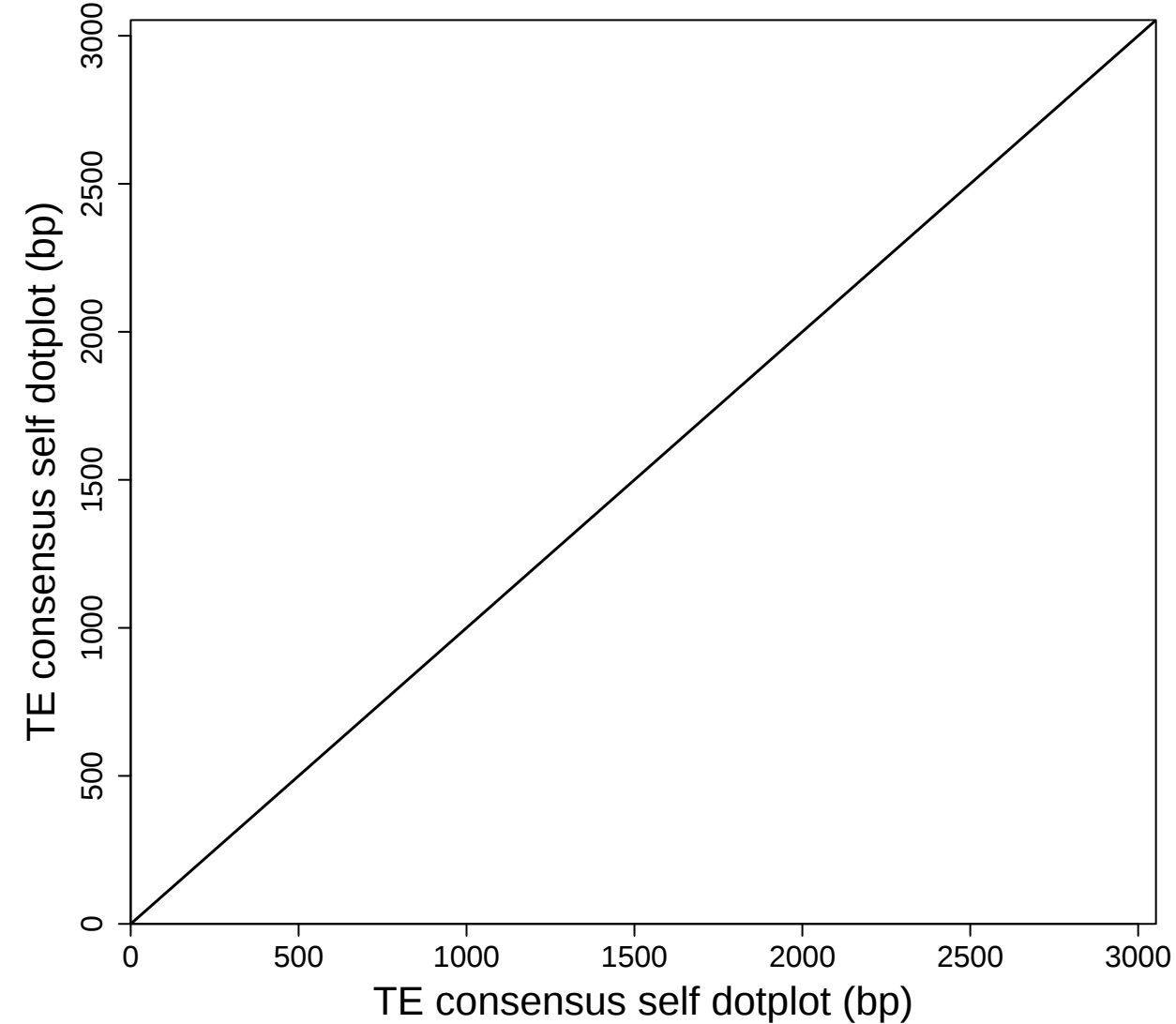
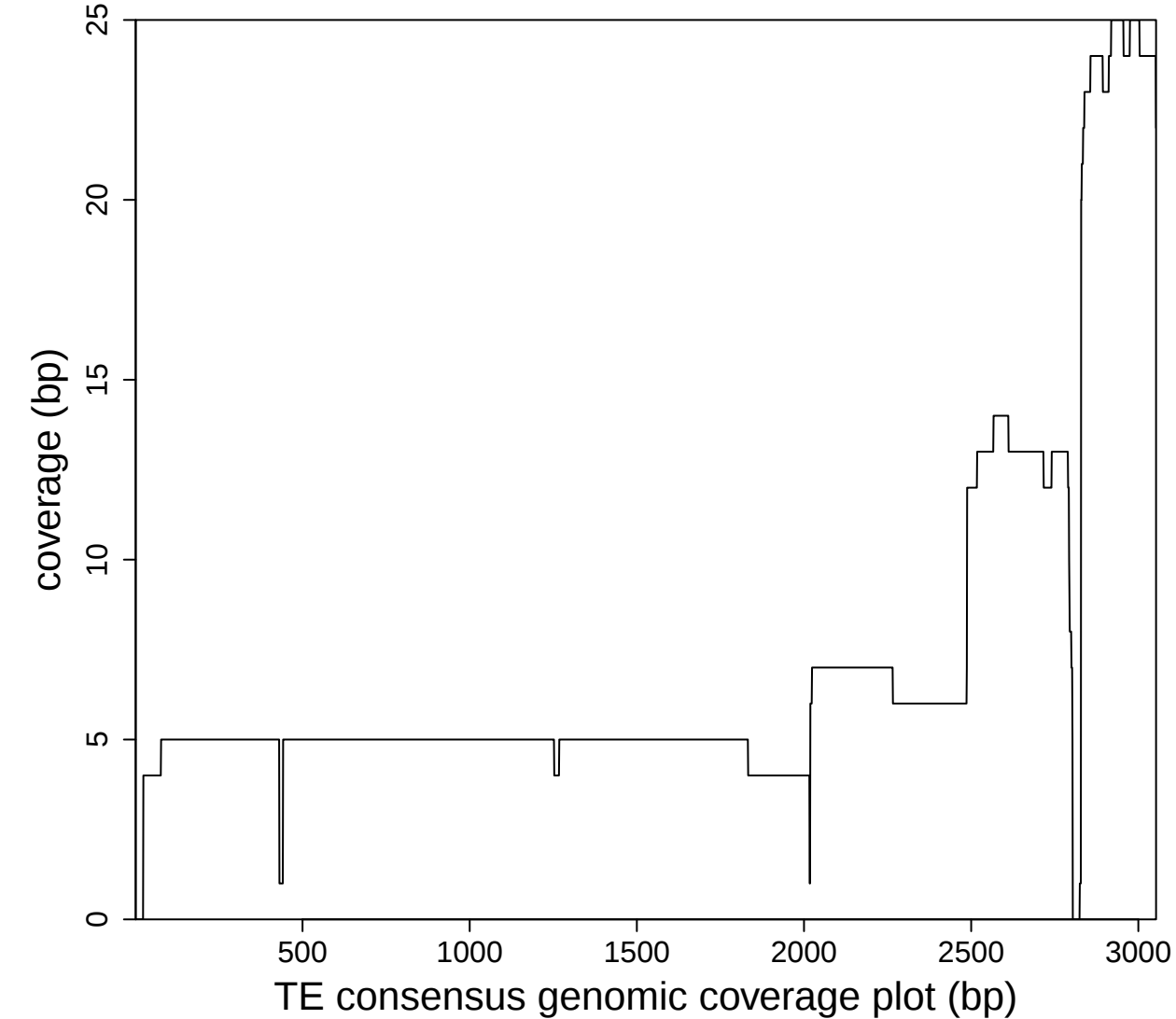
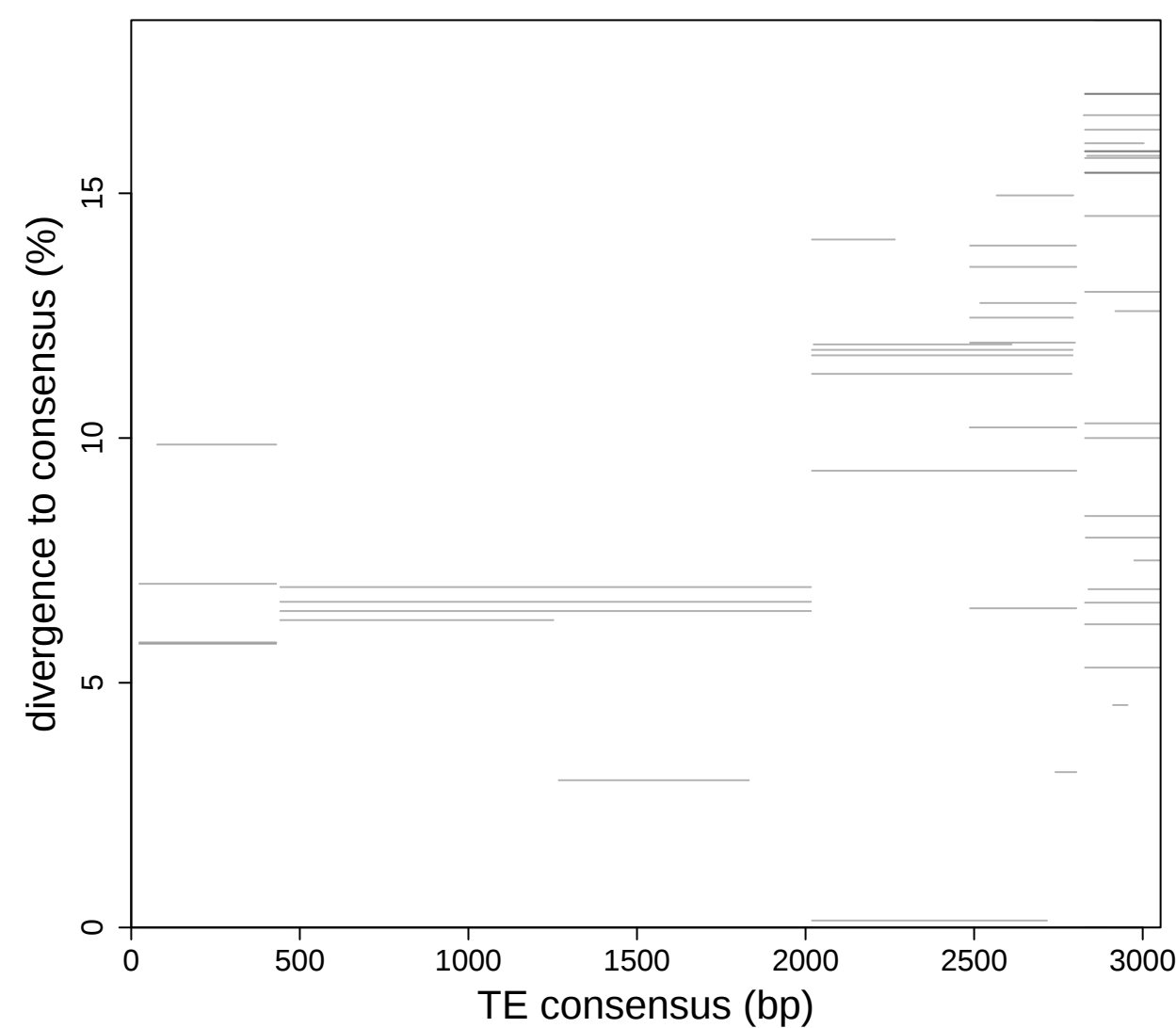


ly_869.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 10480bp; fragments: 1055; full length: 8 (>=9432bp)

After TEtrimmer Extended plot Blue lines are boundaries

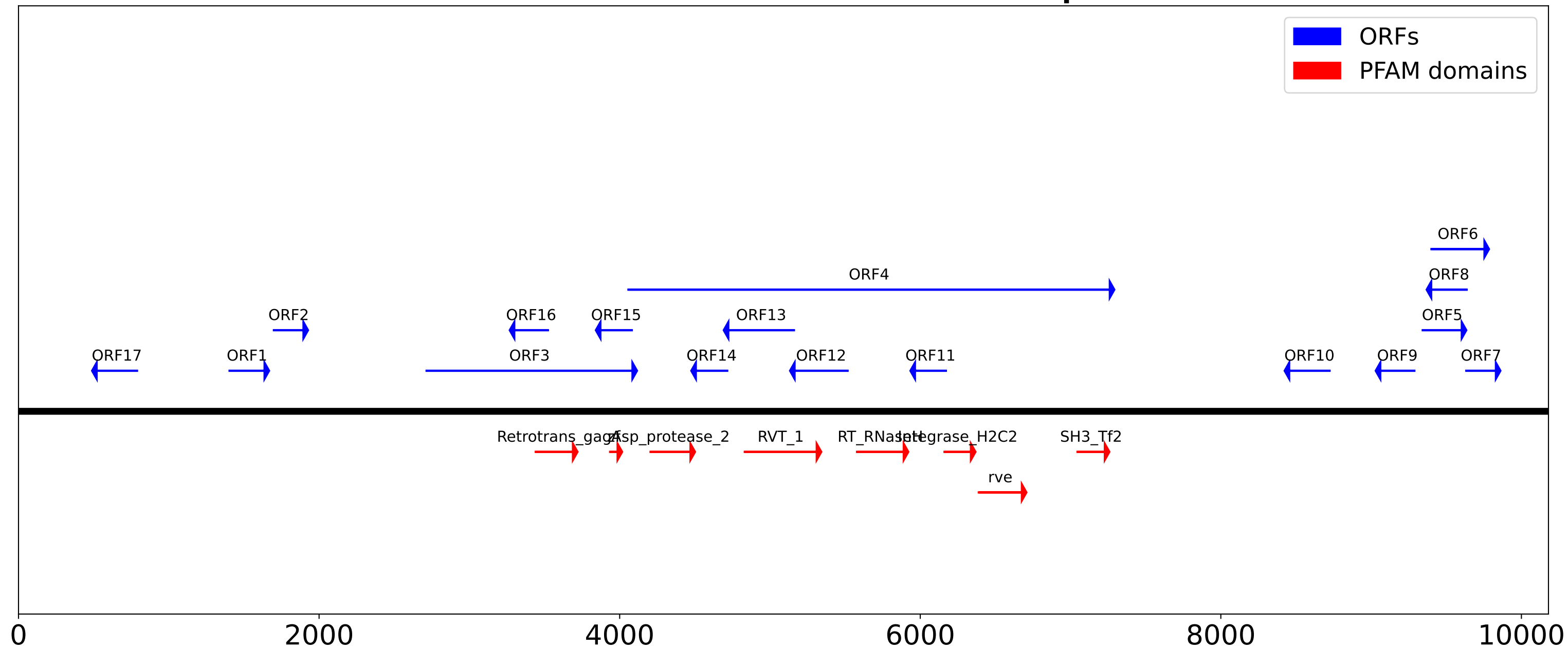


TE: ltr_1_family_869
size: 3053bp; fragments: 54; full length: 0 (≥ 2747.7 bp)

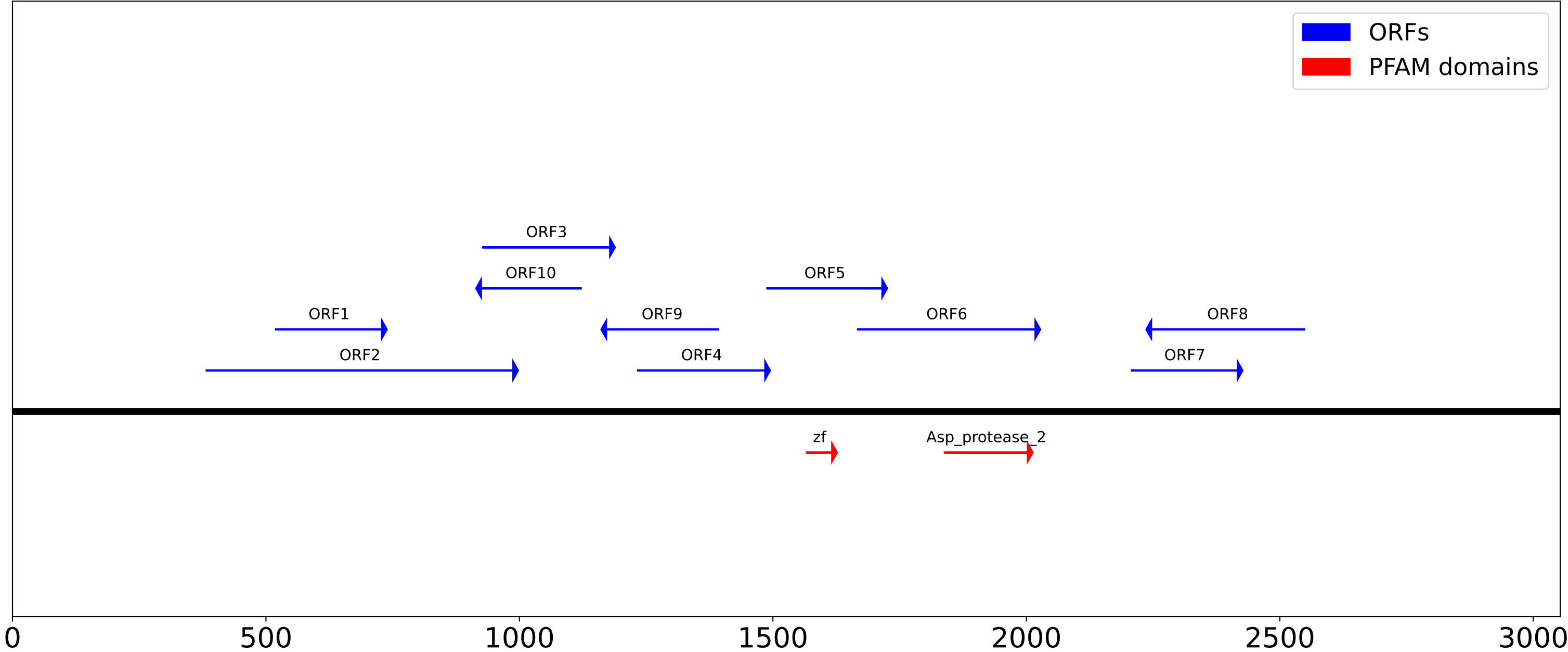


Before TEtrimmer 3053 bp

After TEtrimmer ORF and PFAM domain plot



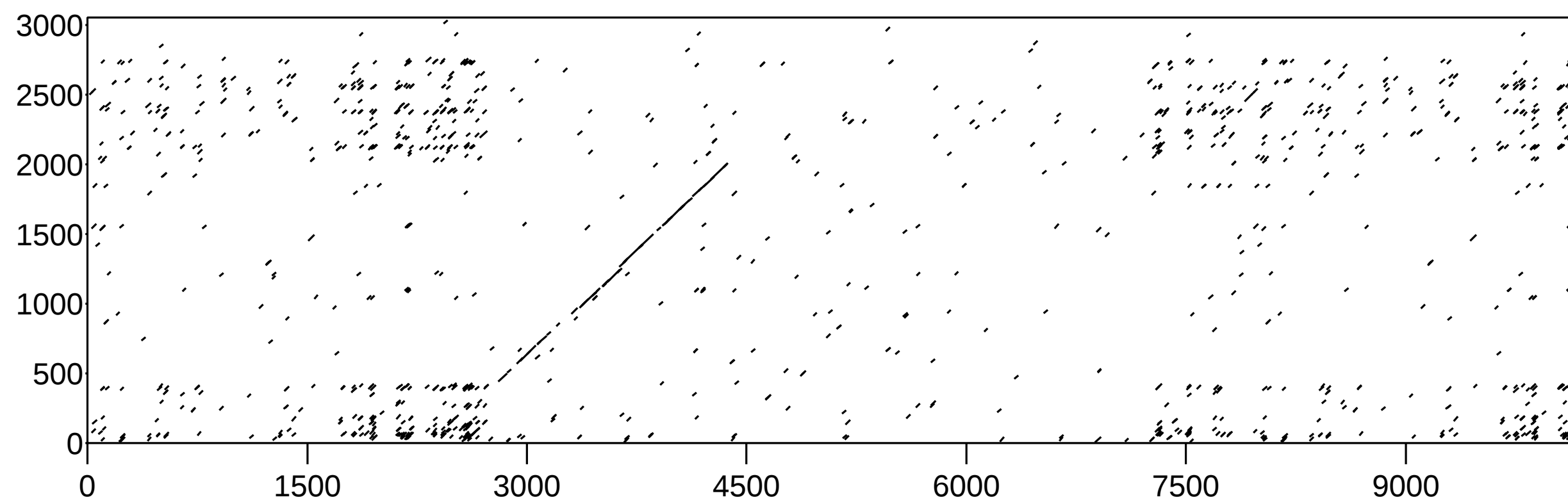
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)