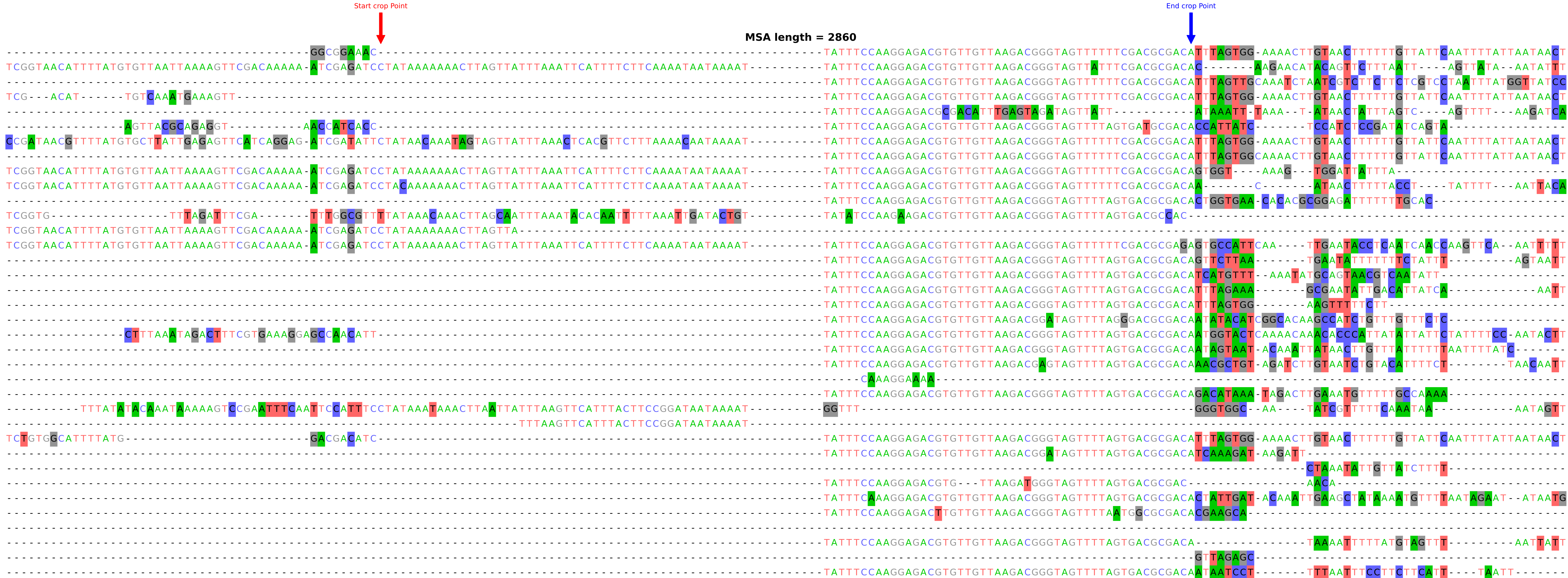
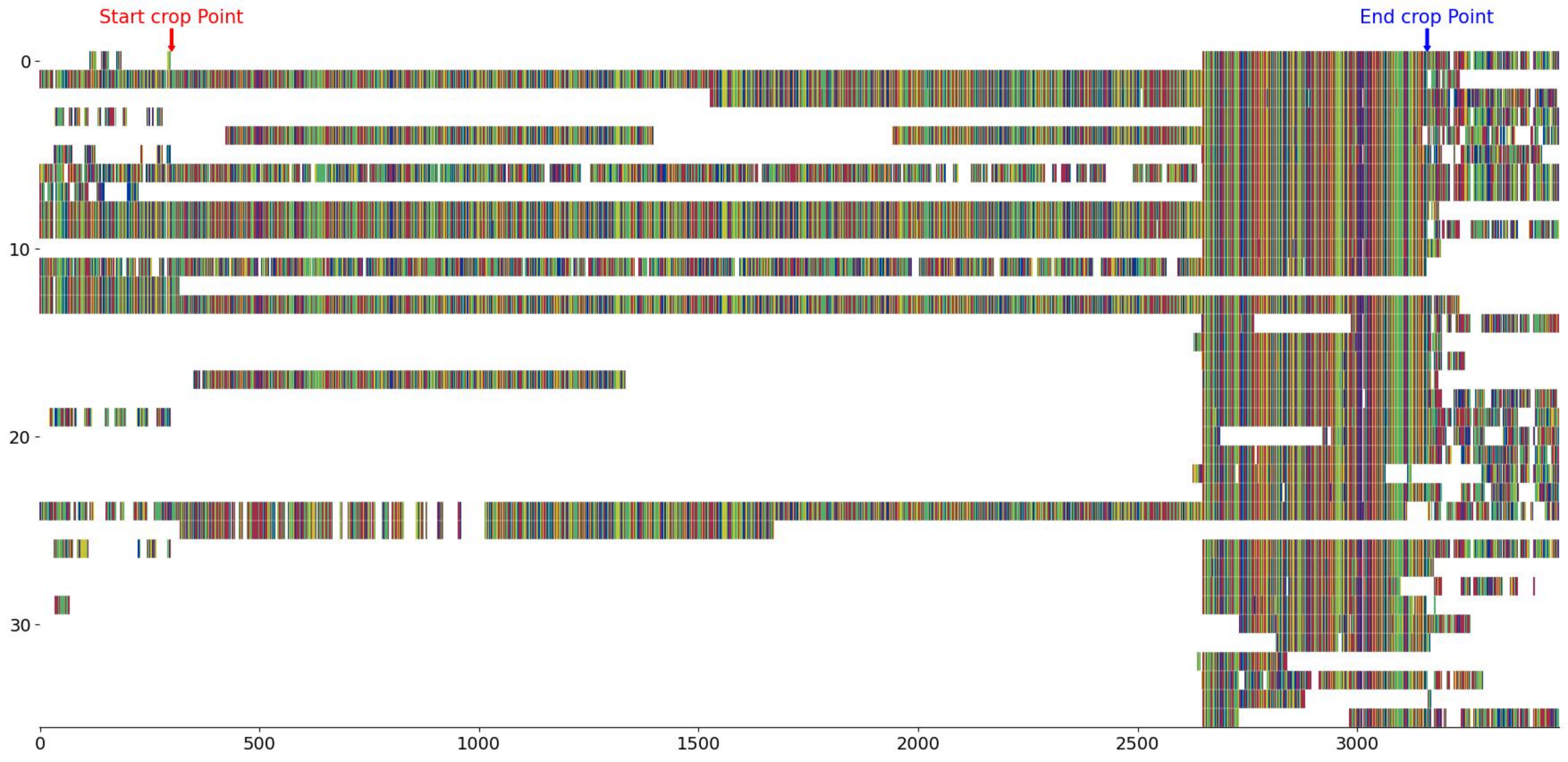
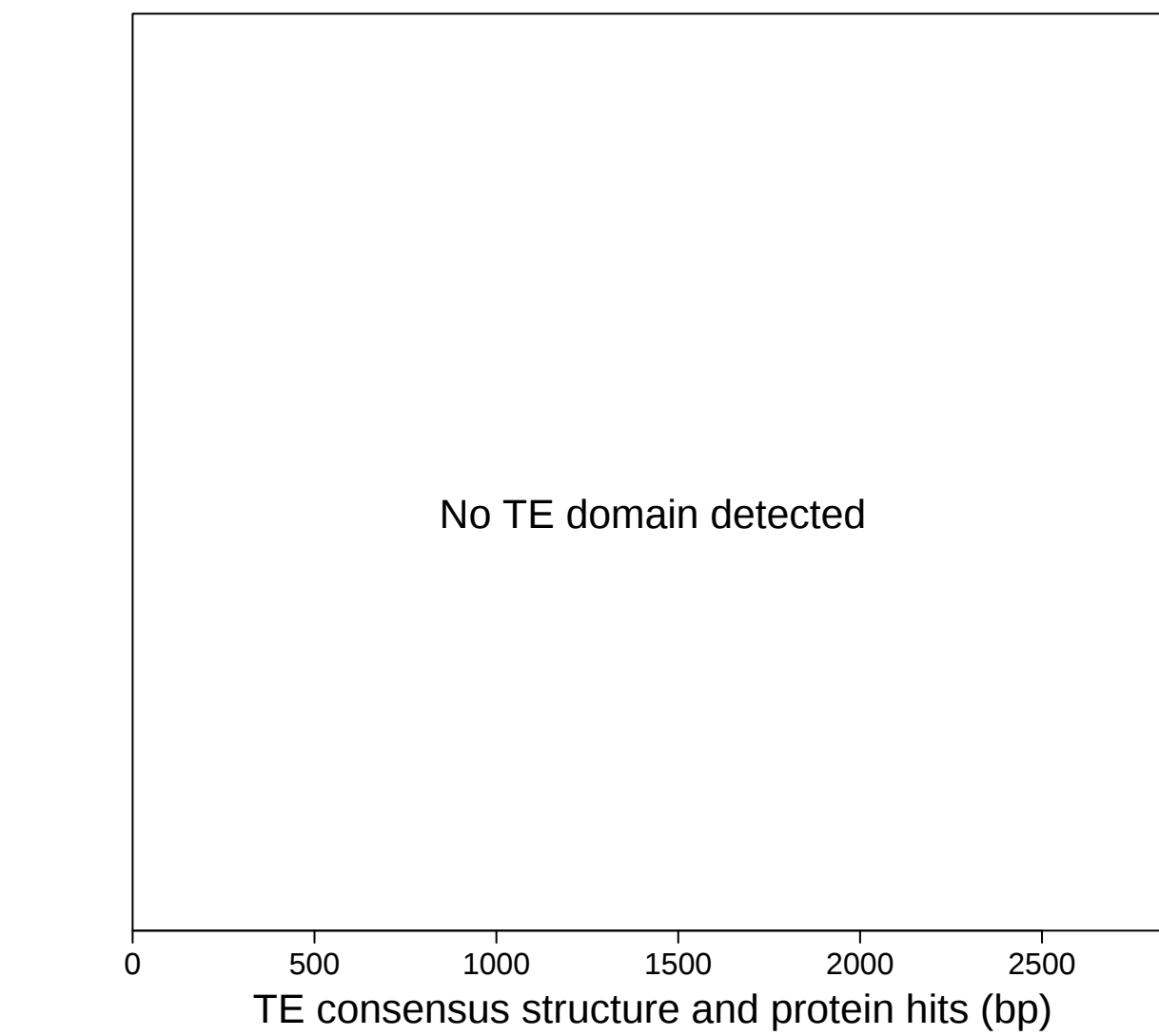
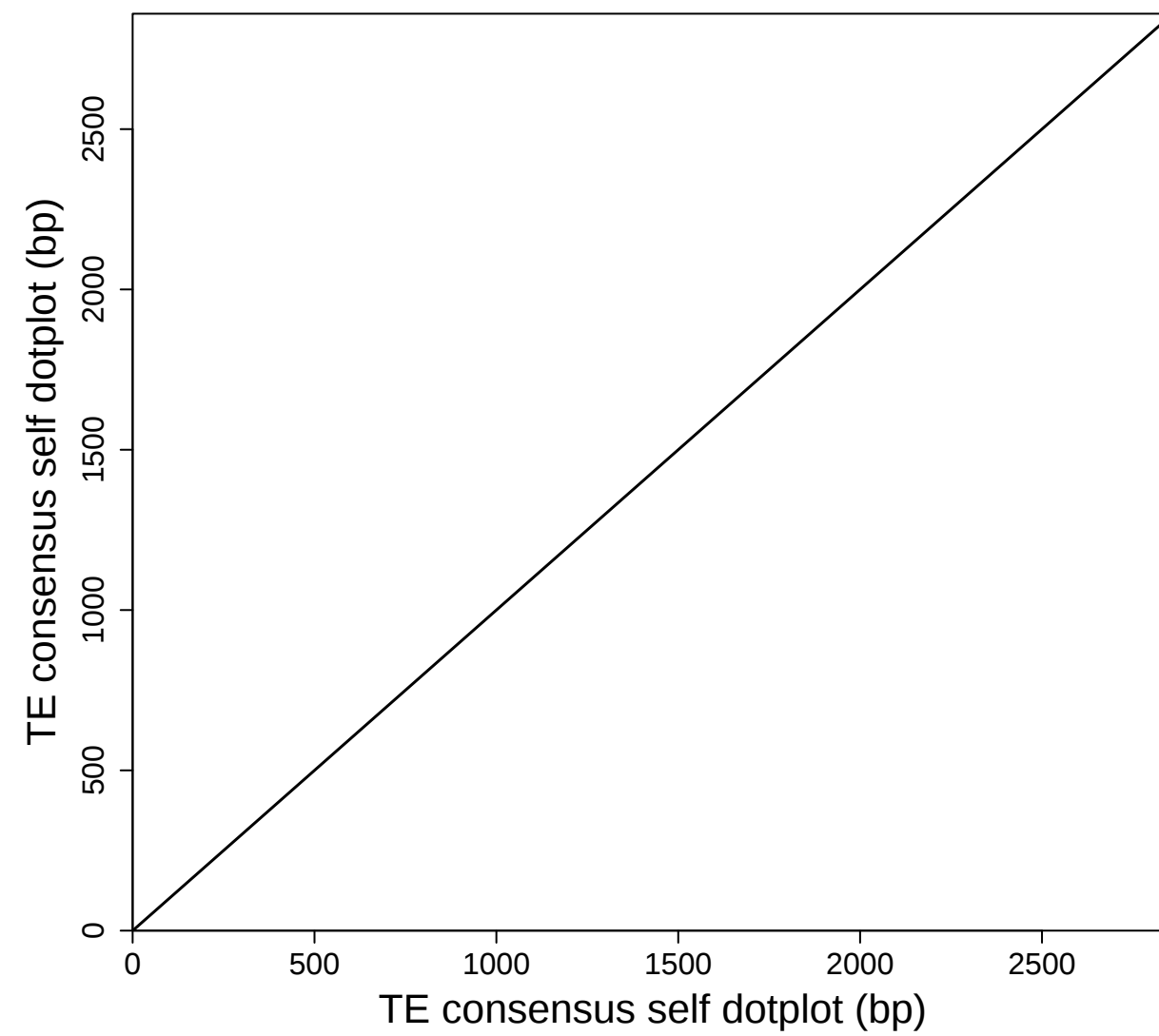
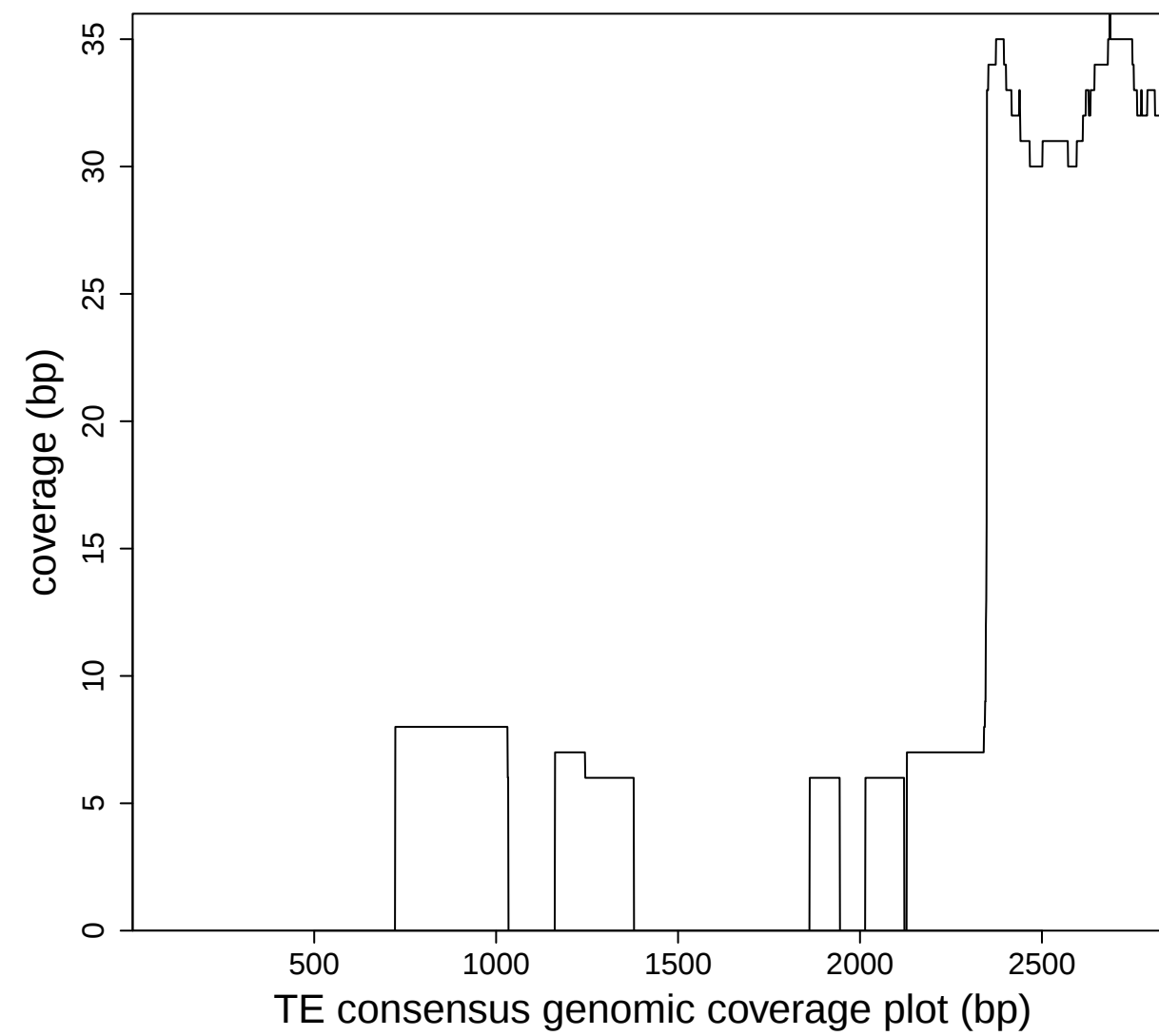
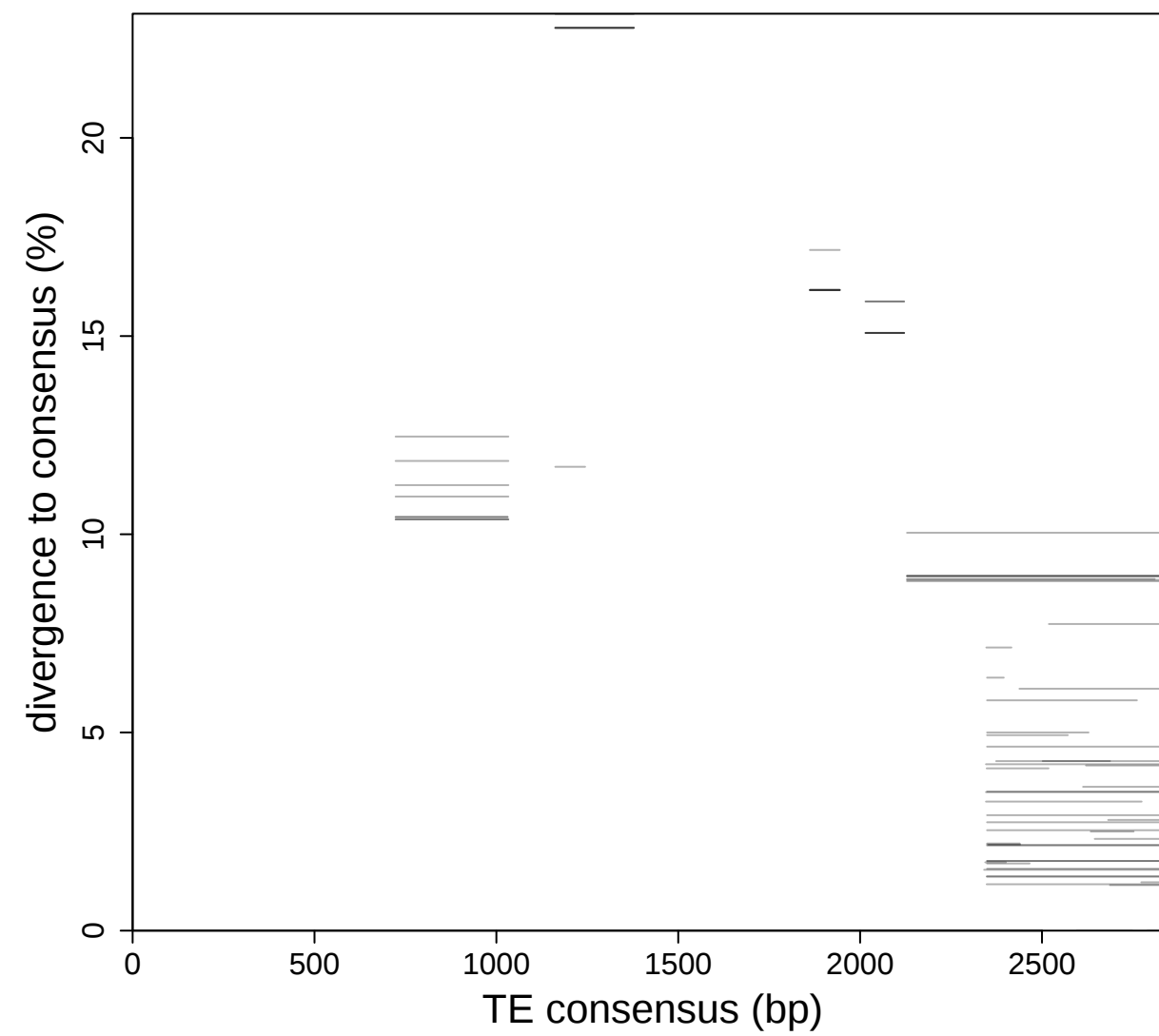






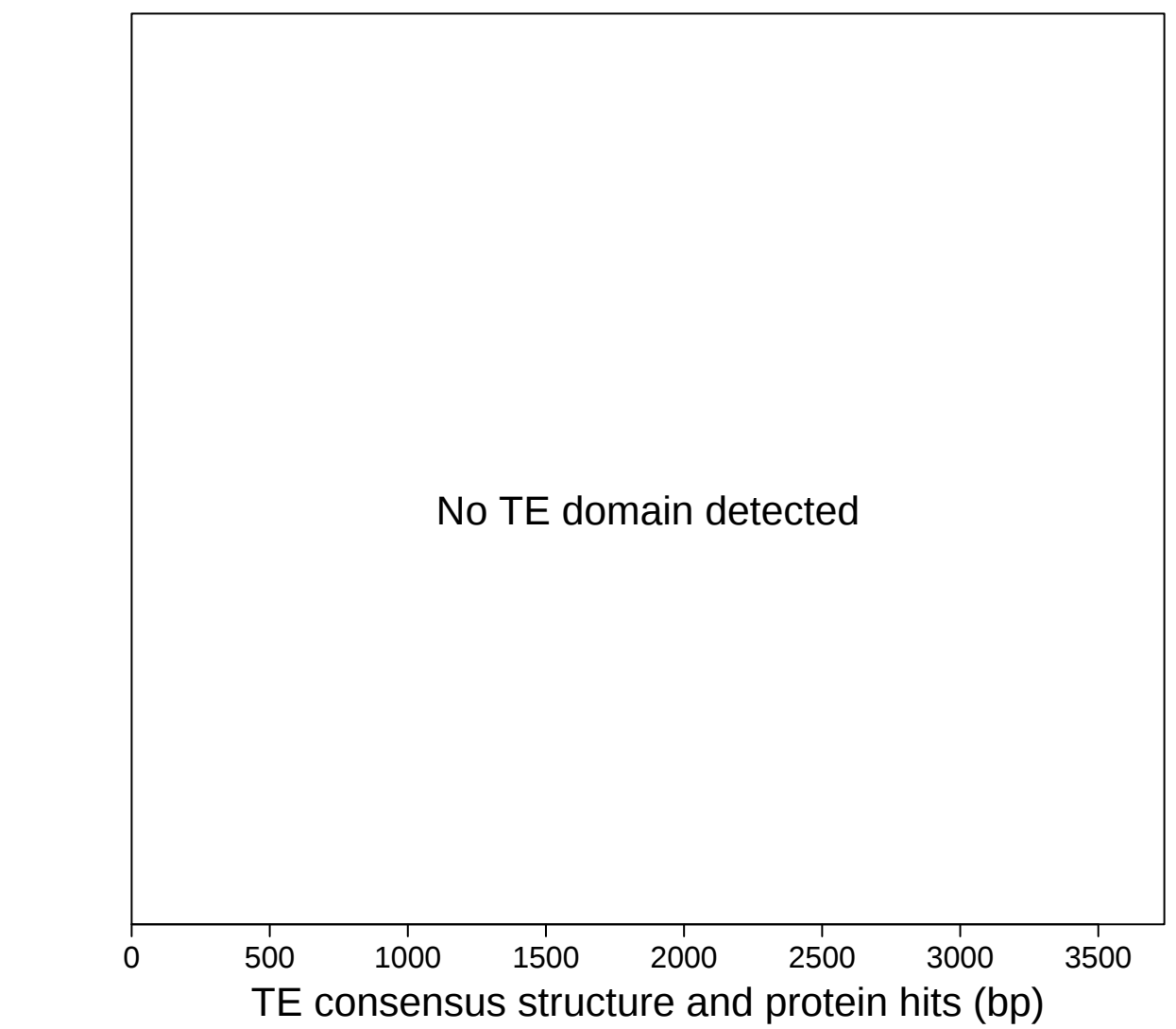
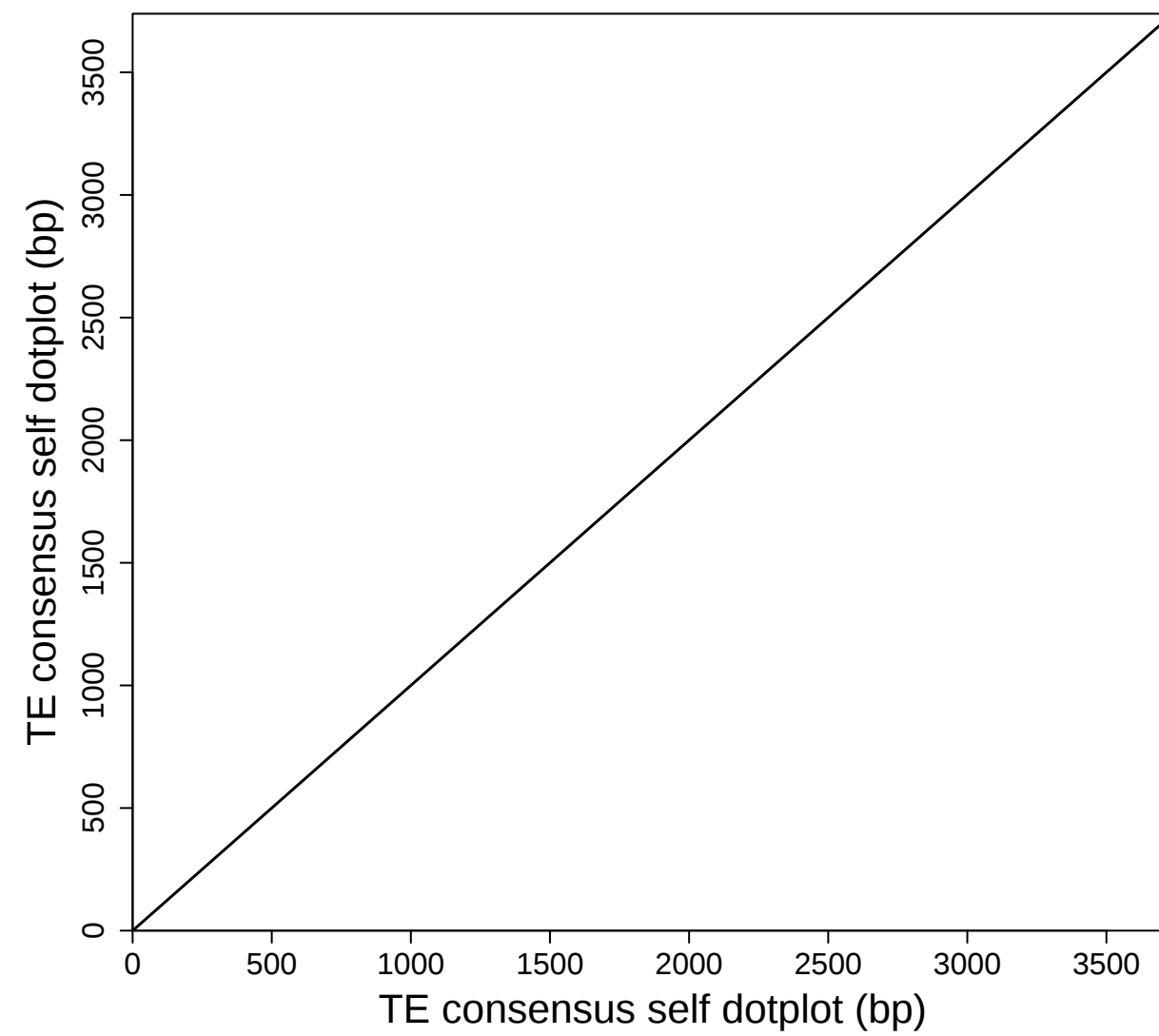
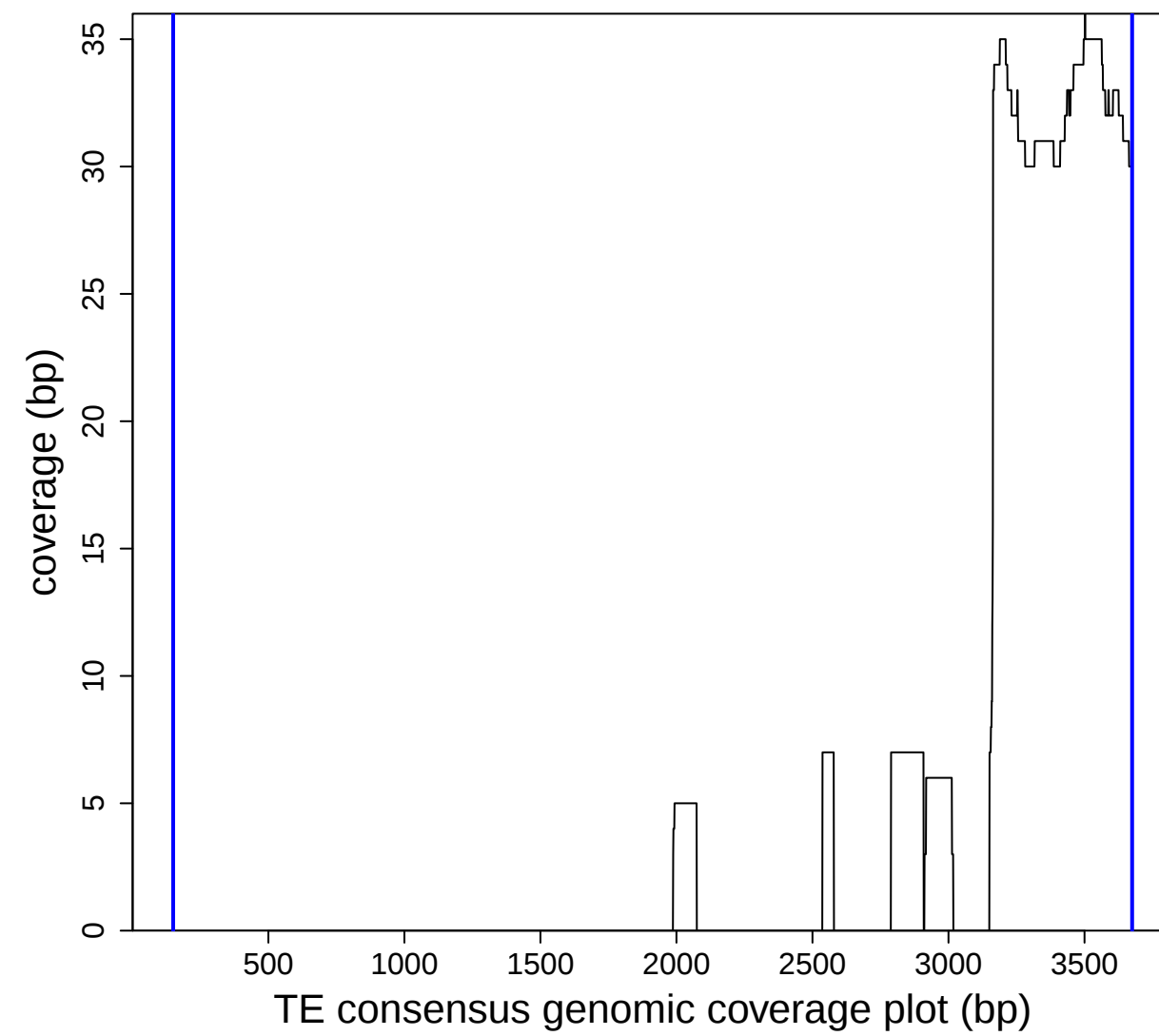
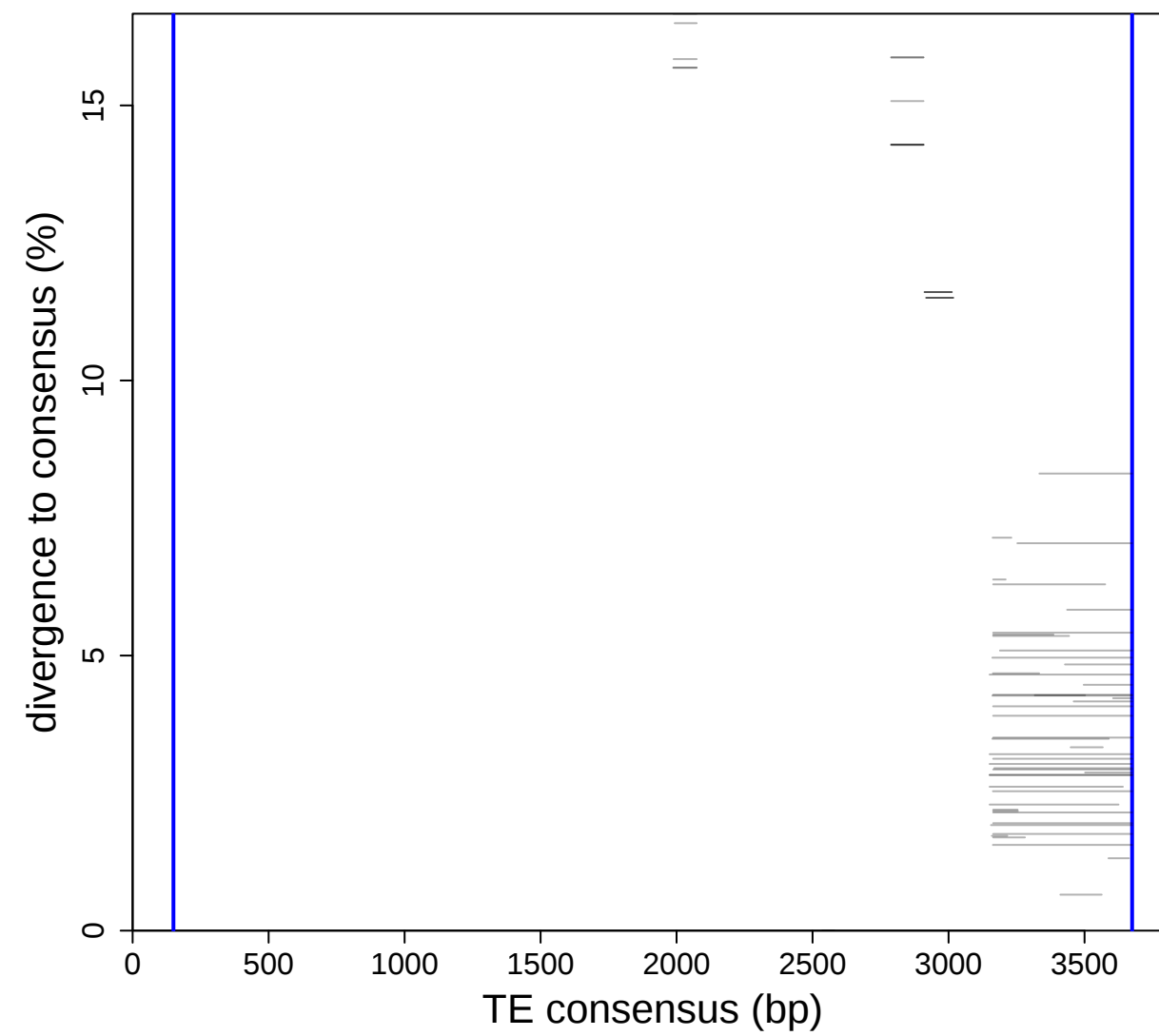
b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 2860bp; fragments: 74; full length: 0 (>=2574bp)

After TEtrimmer 2860 bp

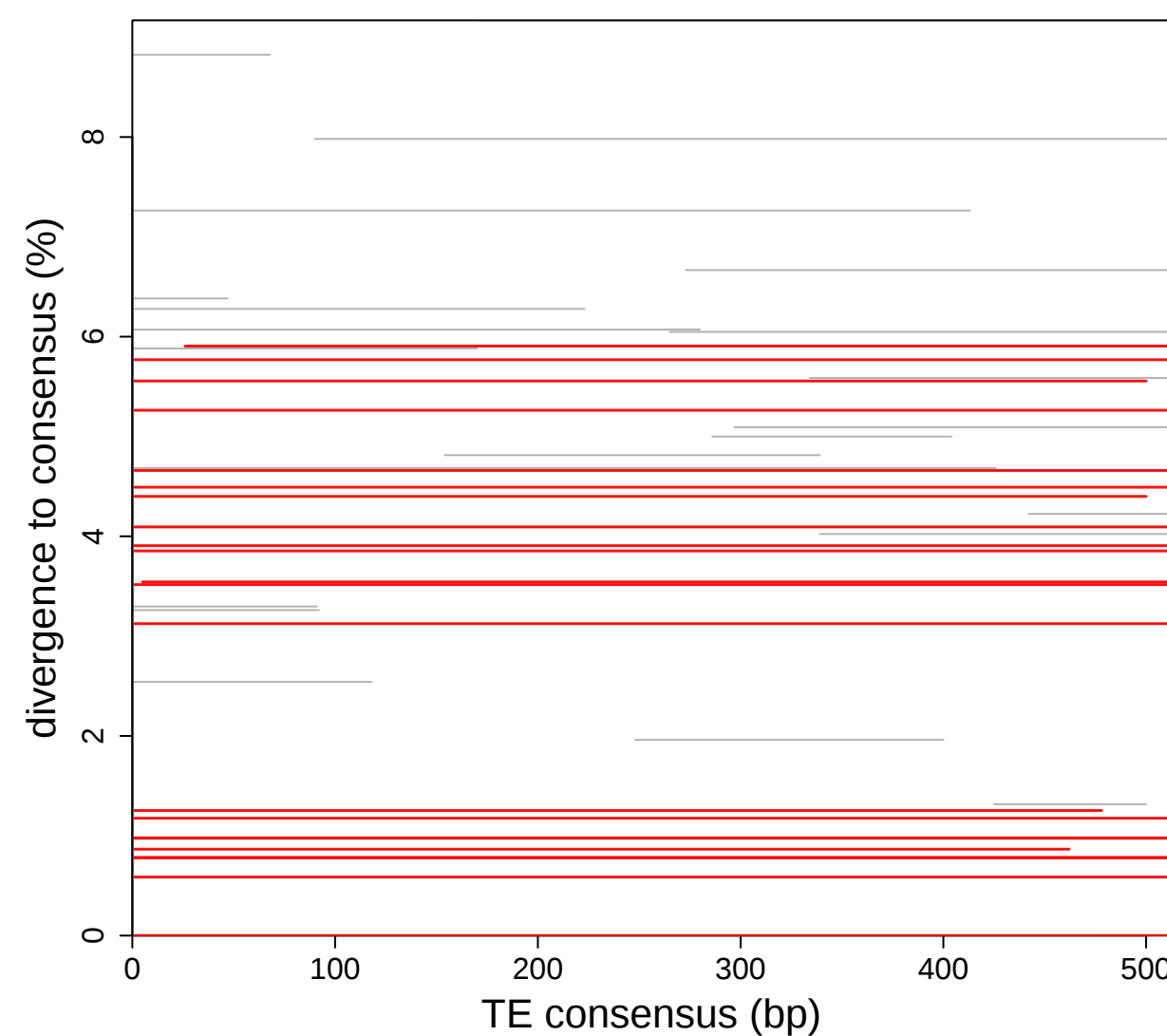


3.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 3825bp; fragments: 72; full length: 0 (>=3442.5bp)

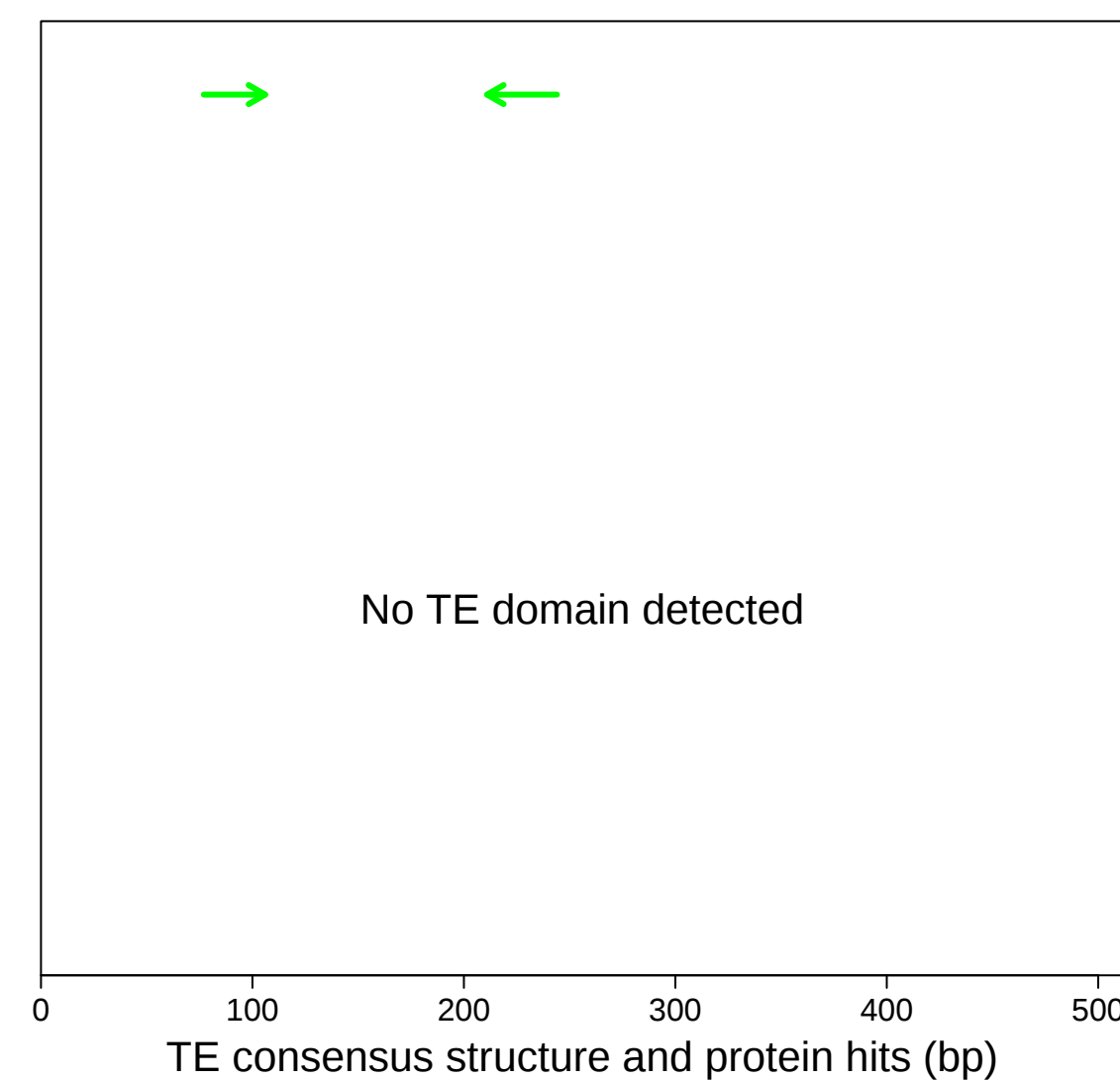
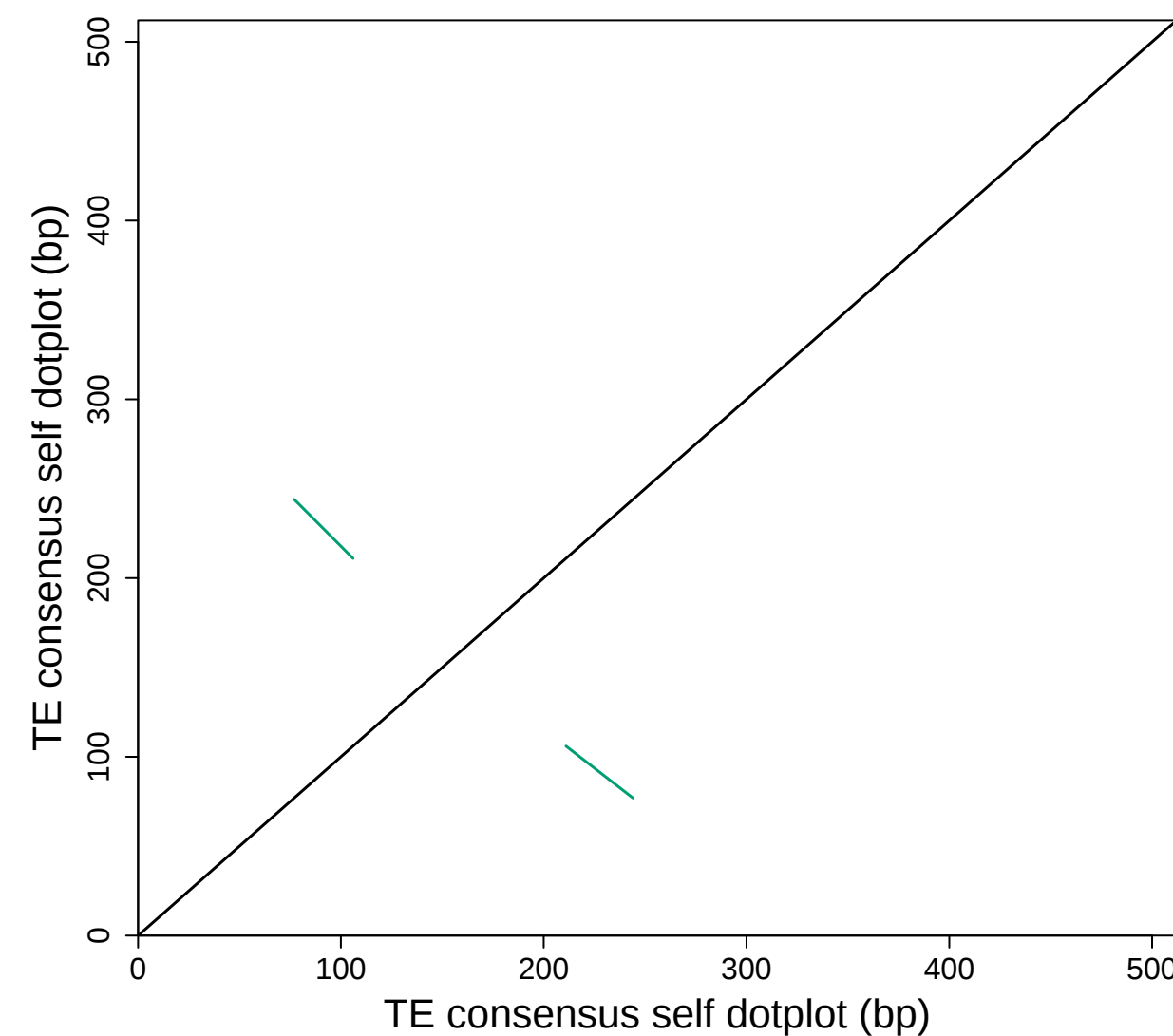
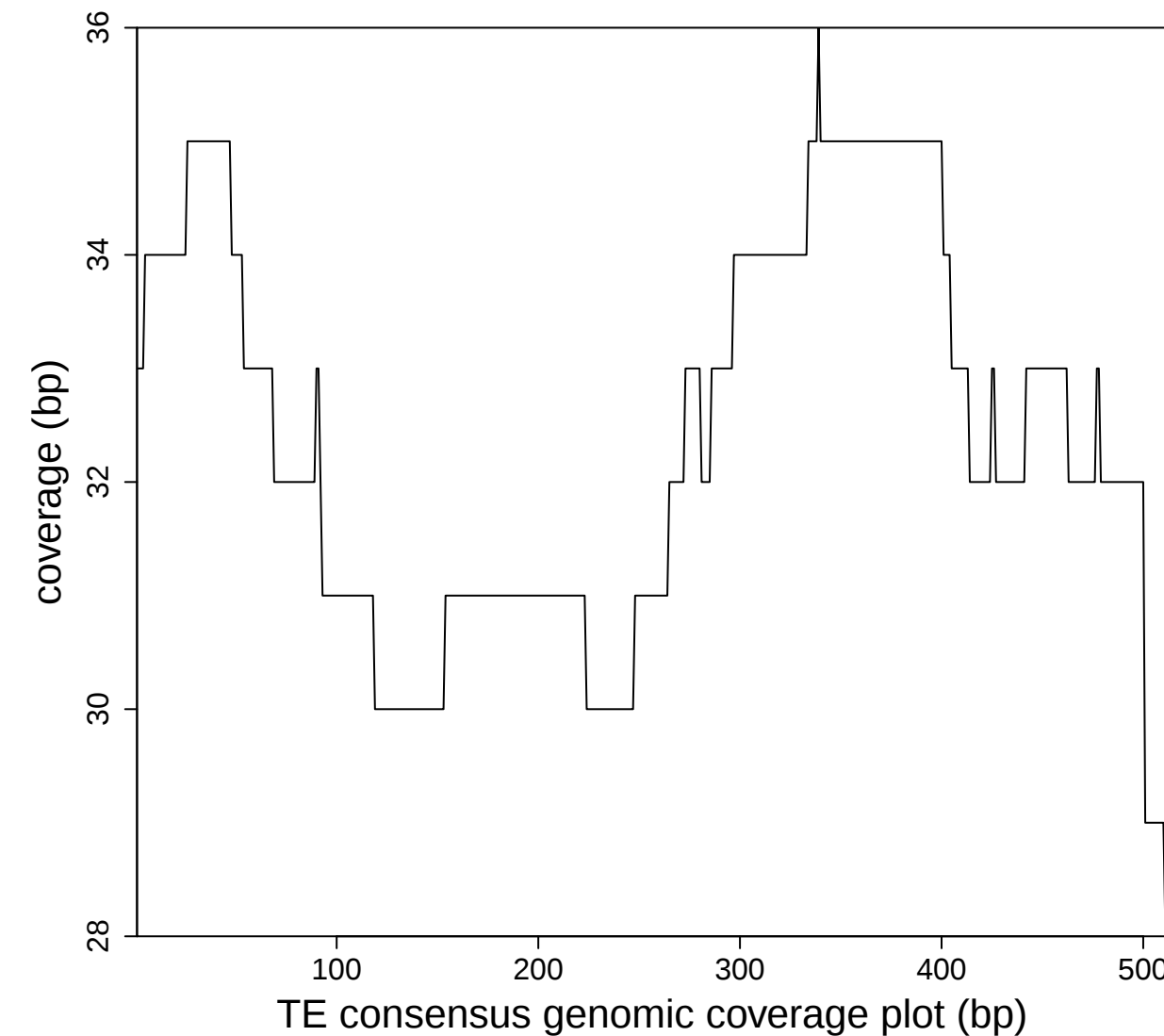
After TEtrimmer Extended plot Blue lines are boundaries



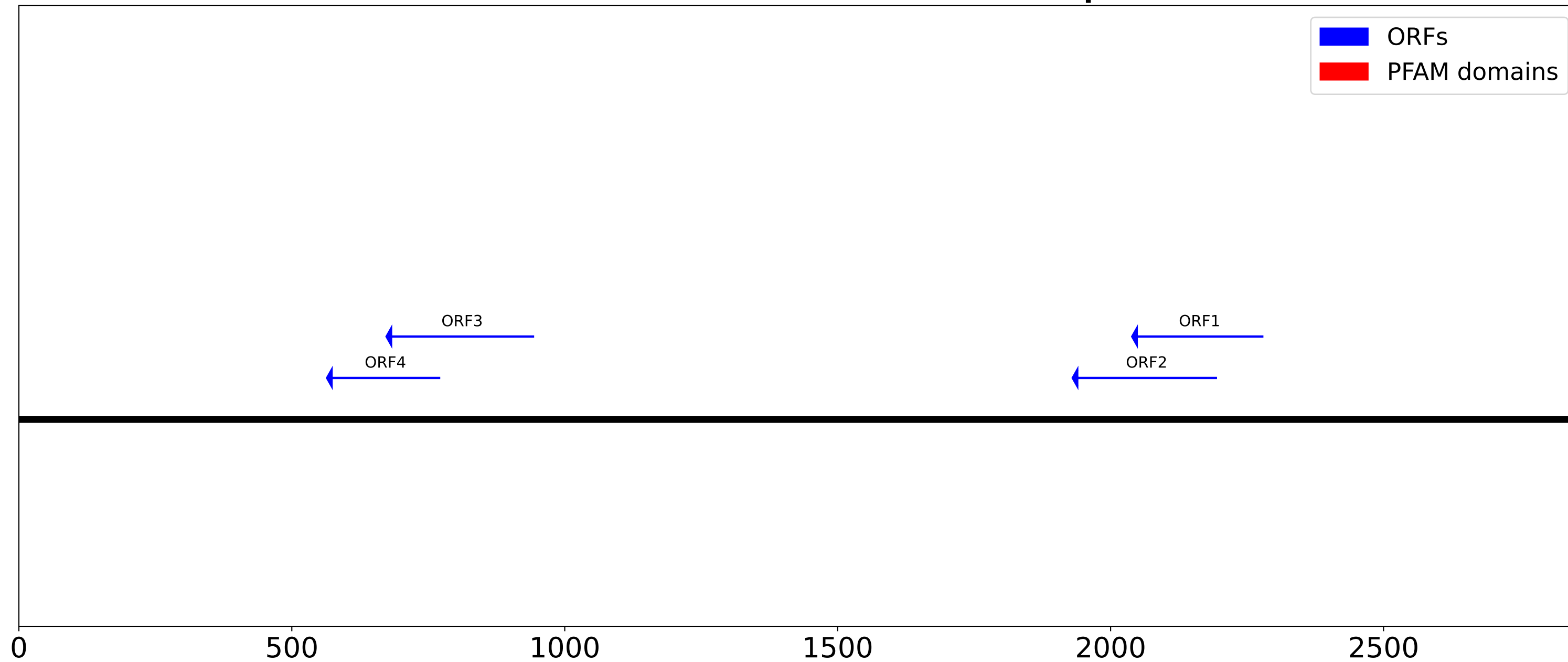
TE: ltr_1_family_253
size: 512bp; fragments: 48; full length: 24 (≥ 460.8 bp)



Before TEtrimmer 512 bp



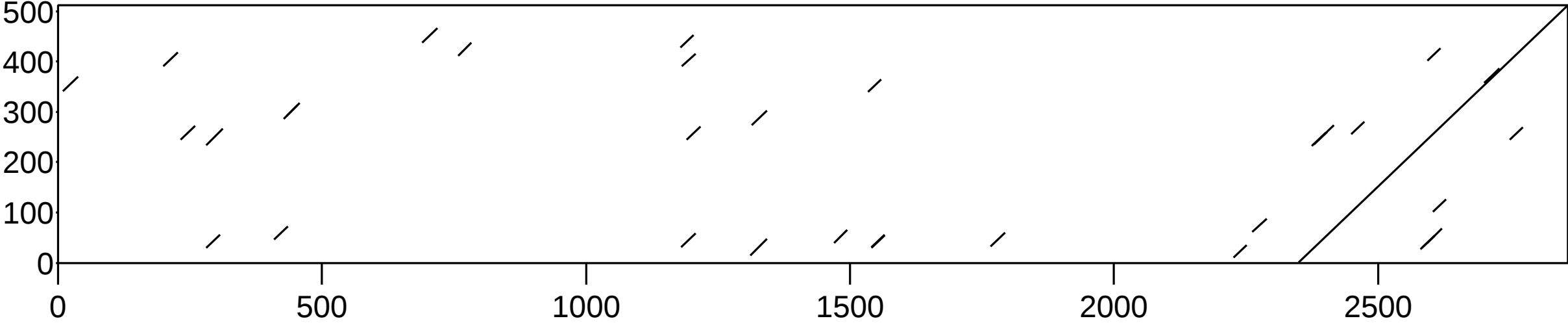
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)