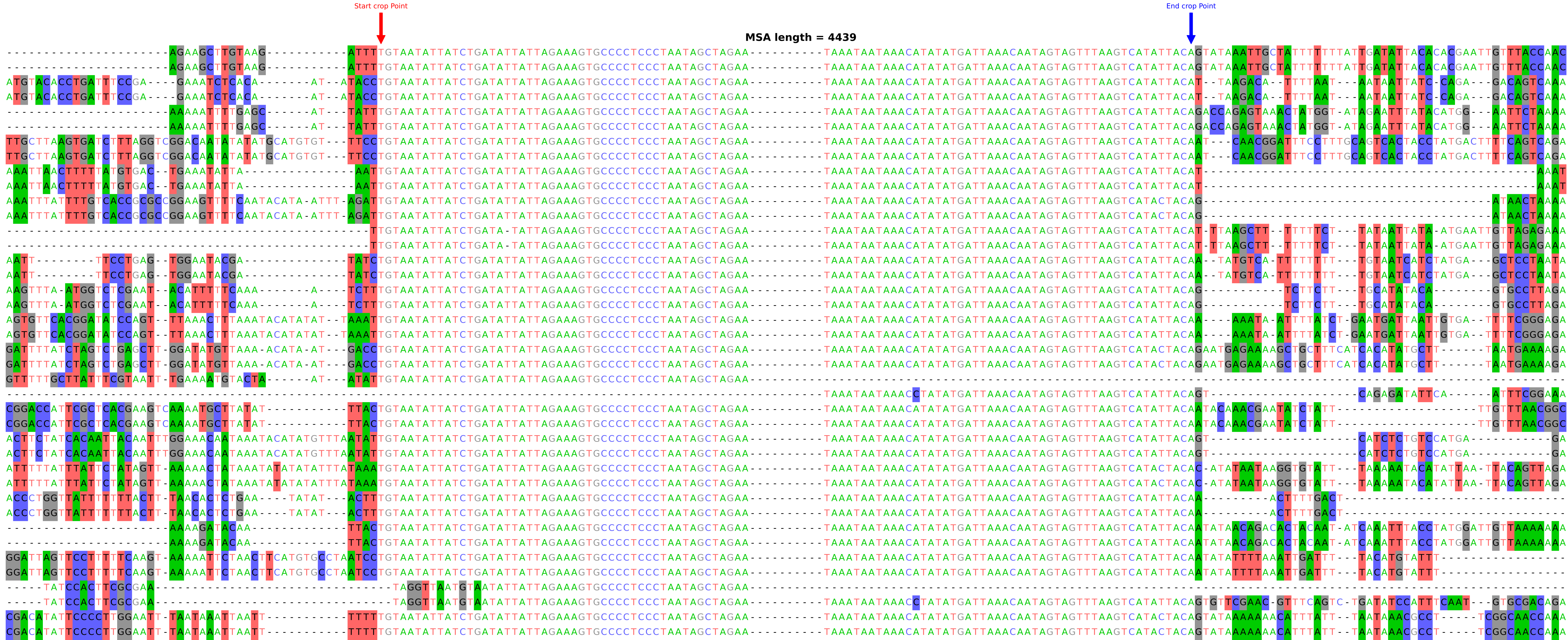
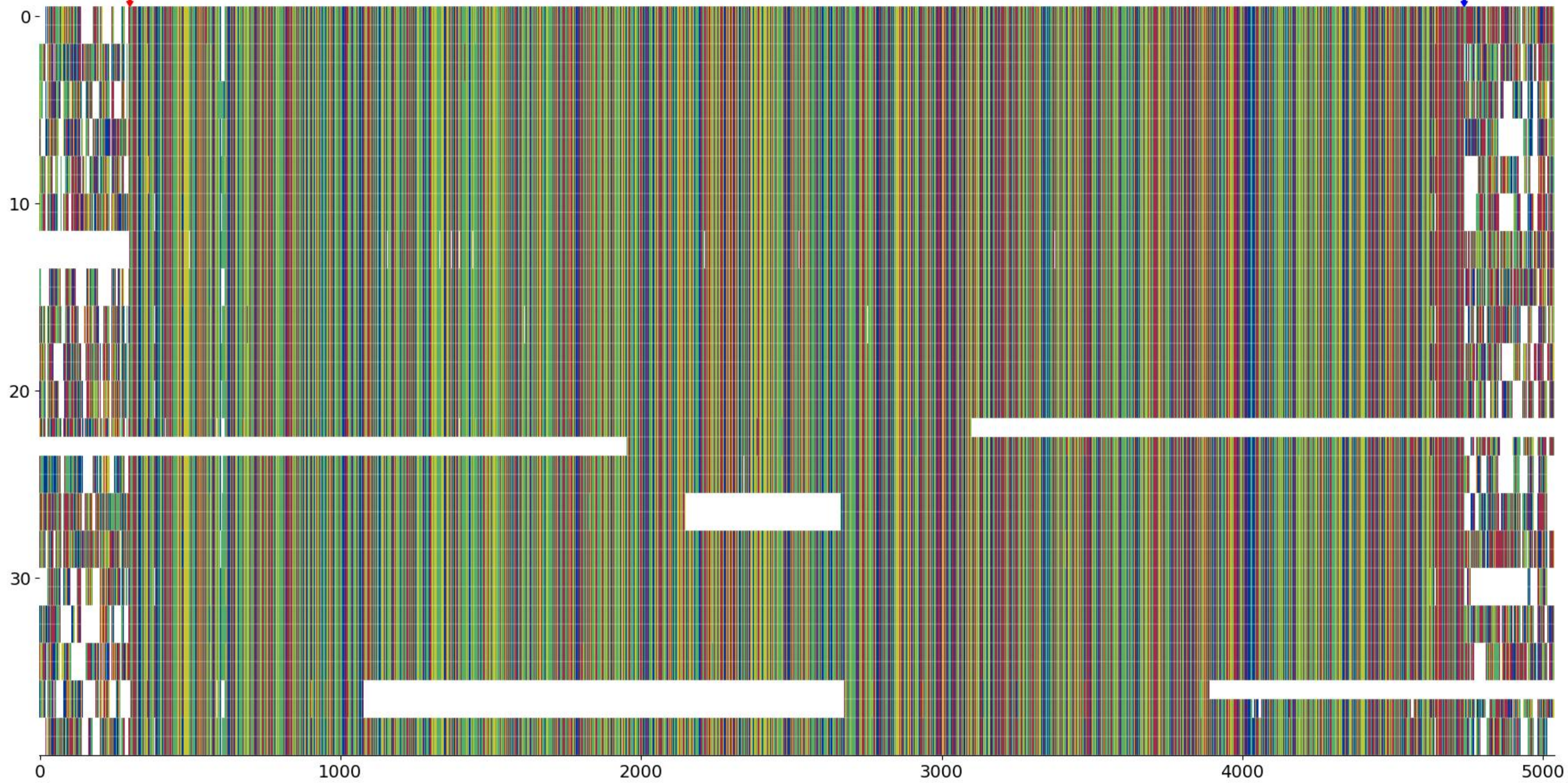




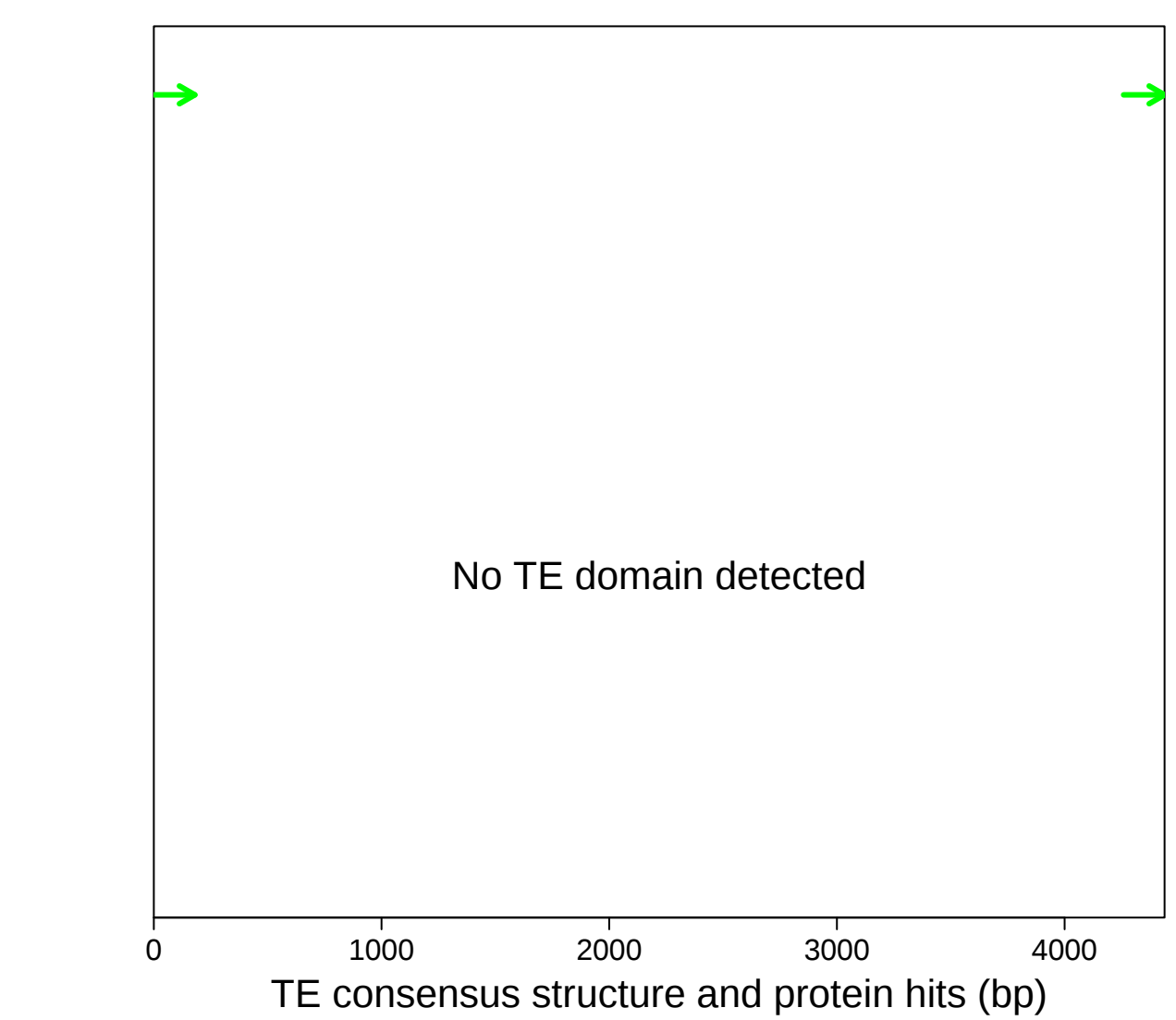
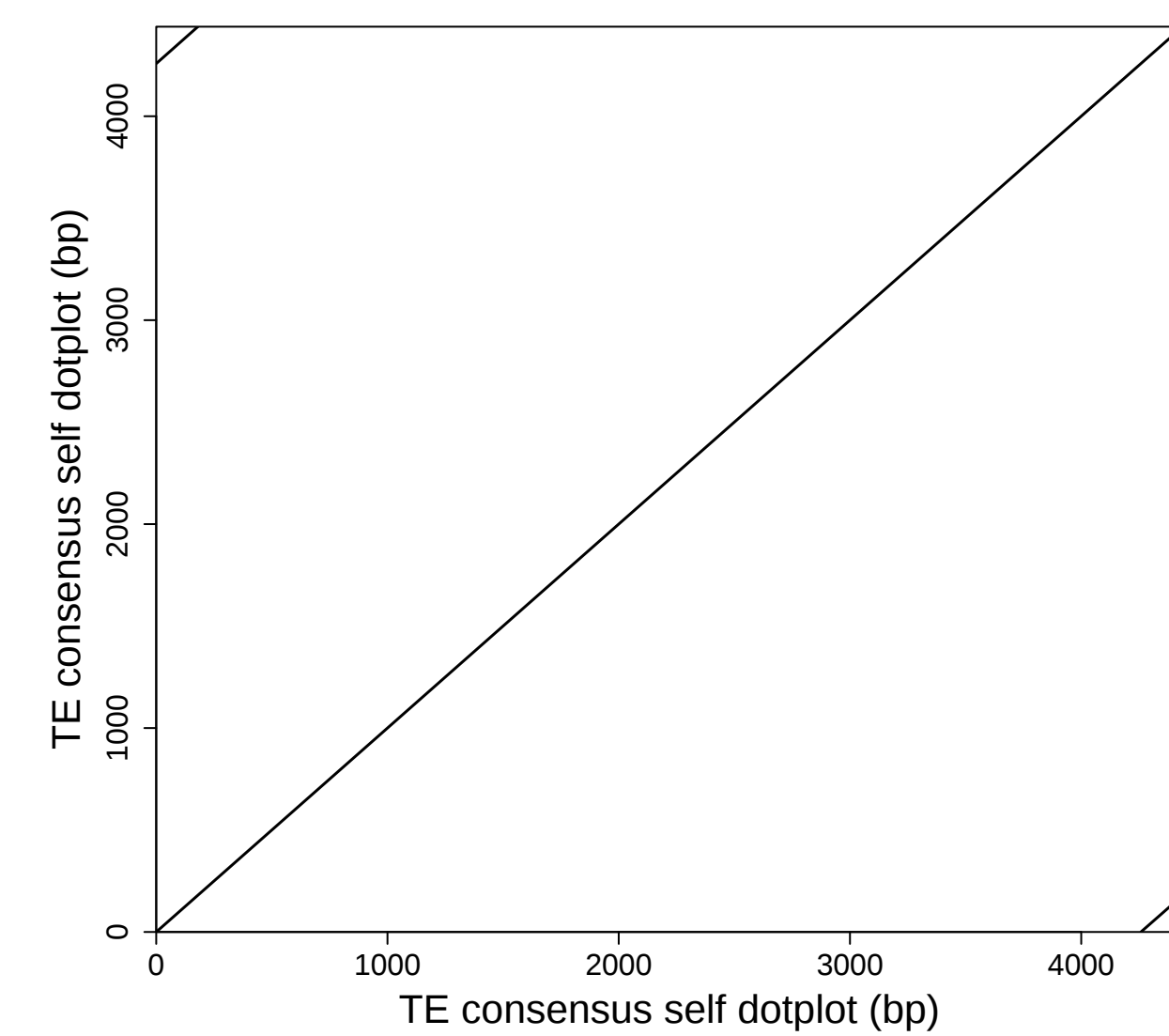
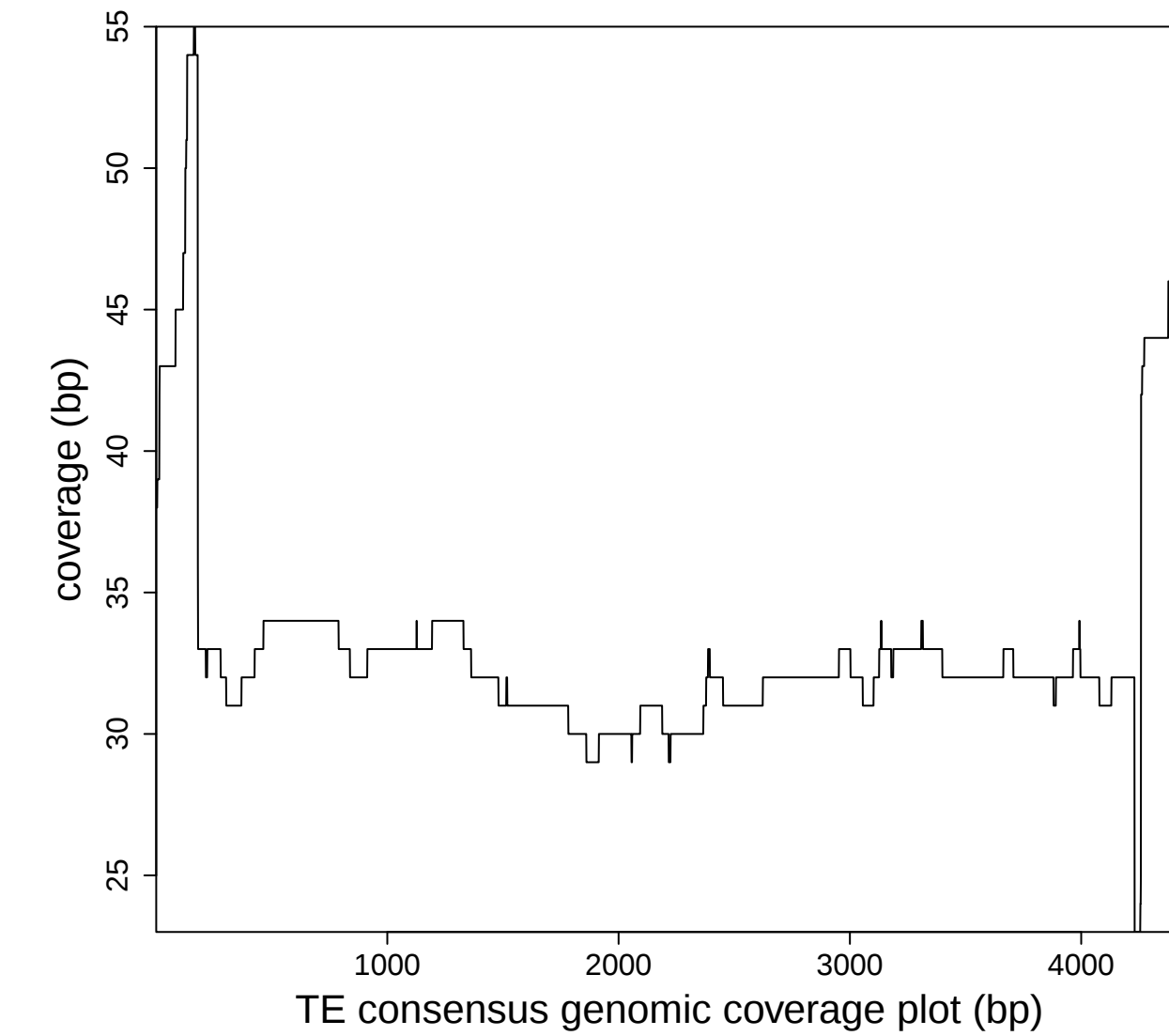
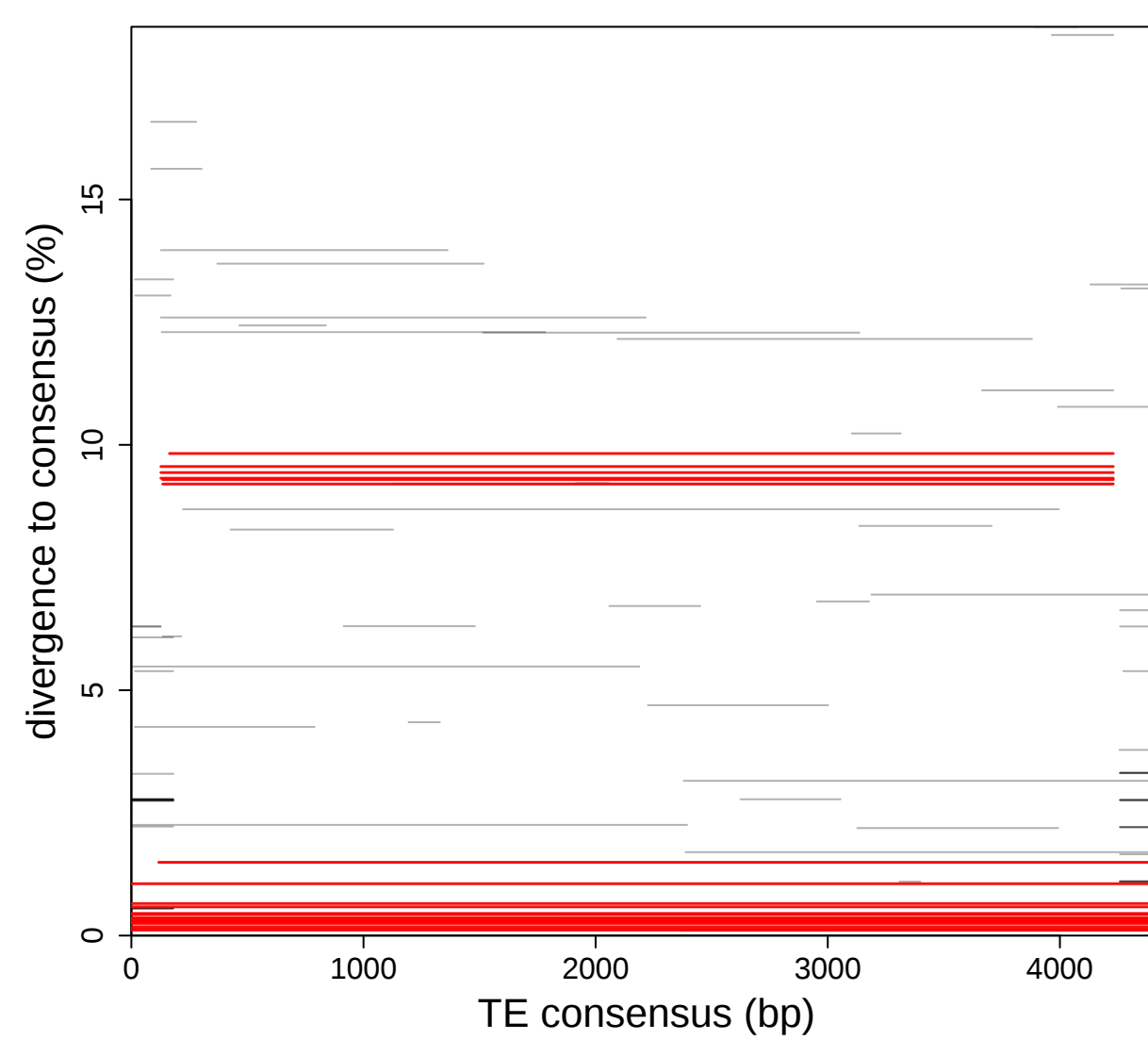
Start crop Point

End crop Point



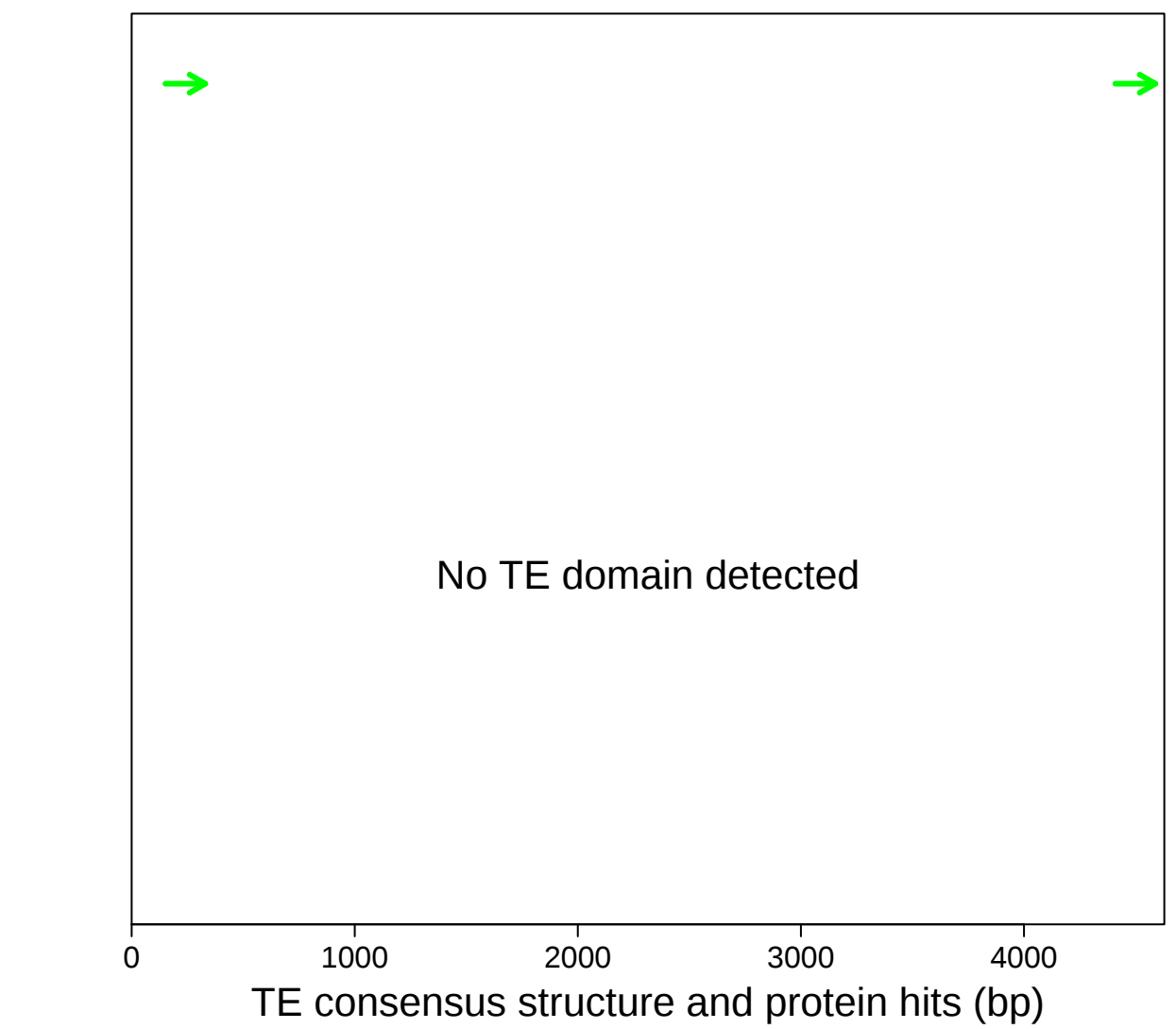
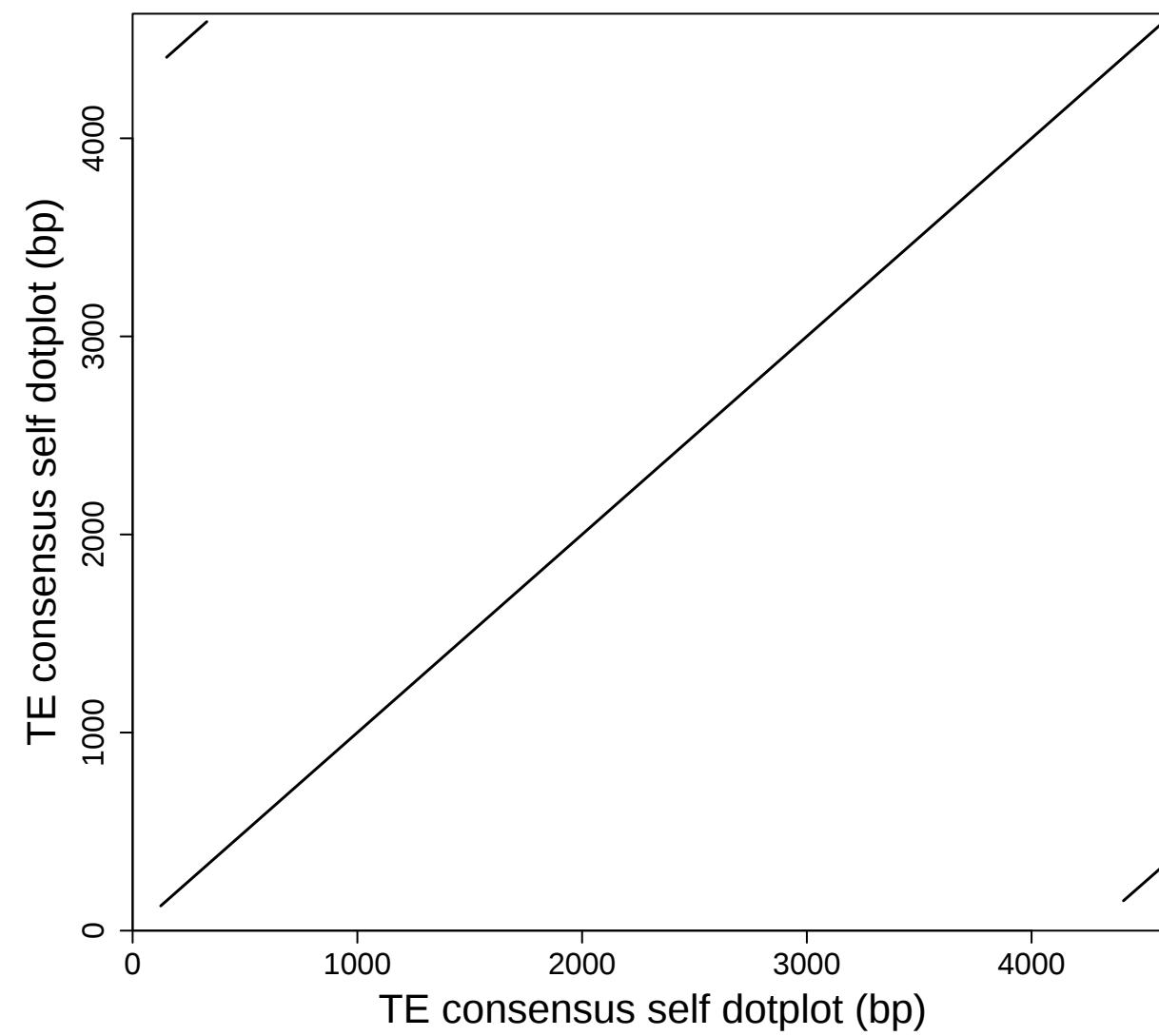
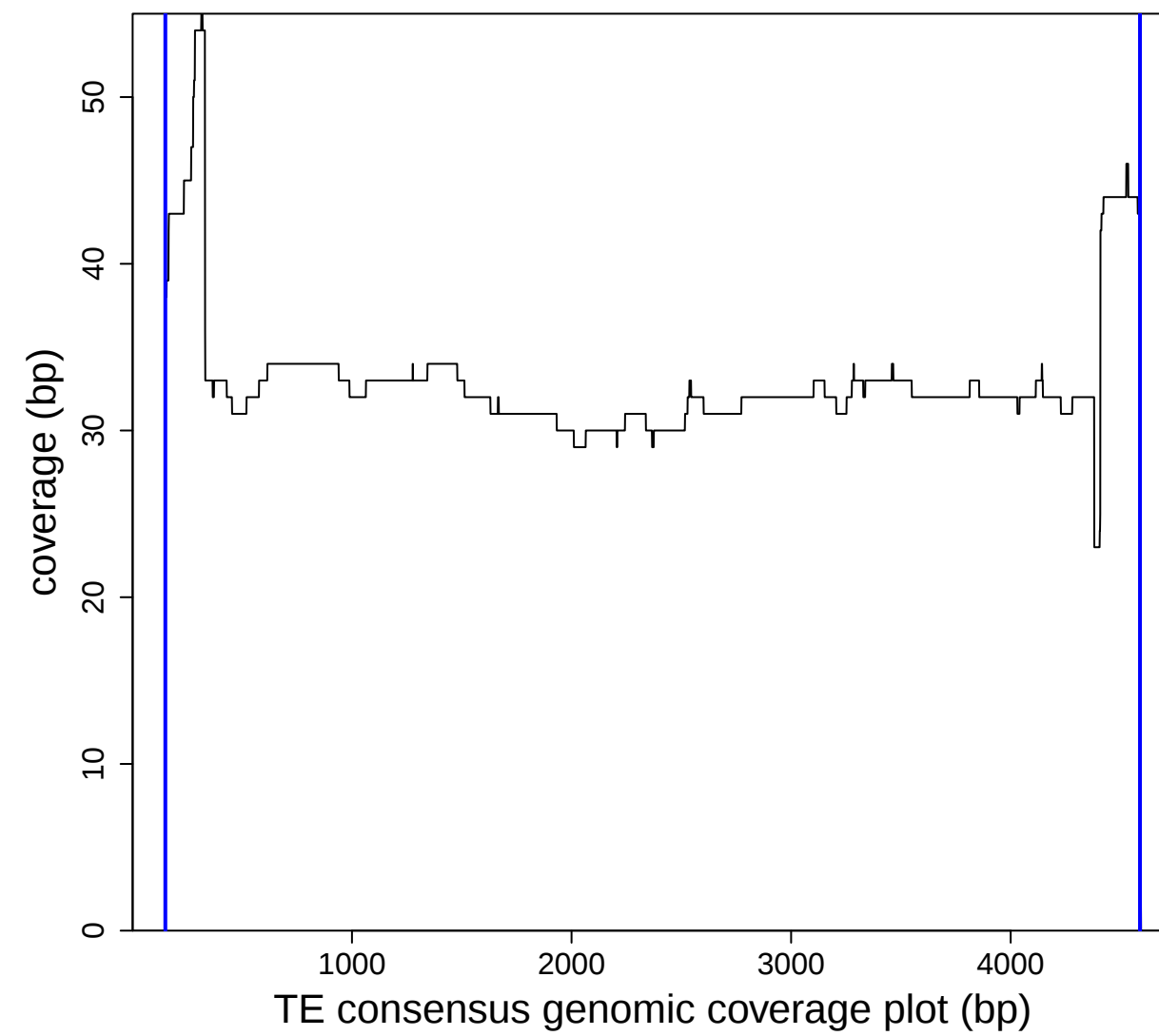
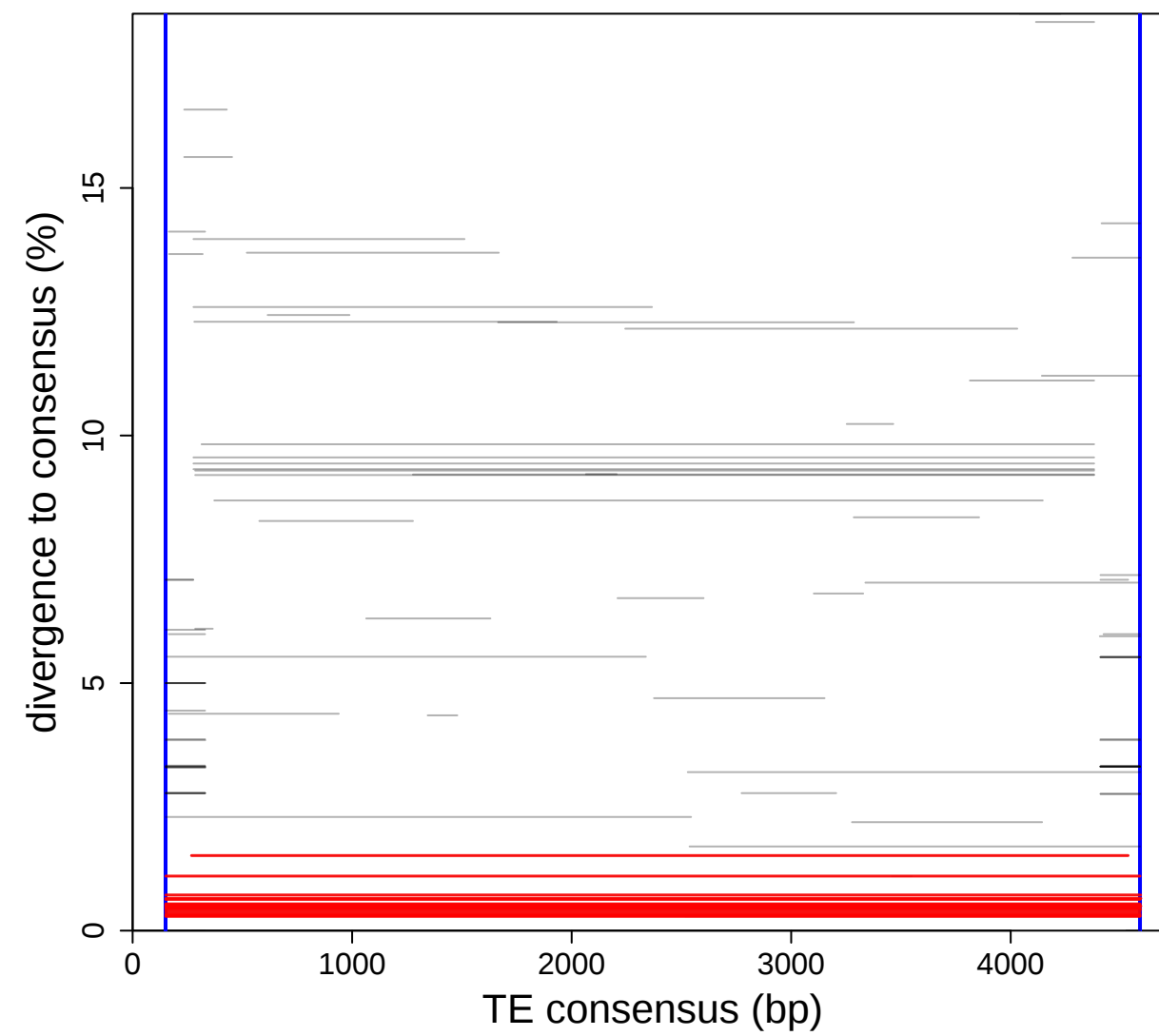
ed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.f
size: 4439bp; fragments: 107; full length: 23 (≥ 3995.1 bp)

After TETrimmer 4439 bp

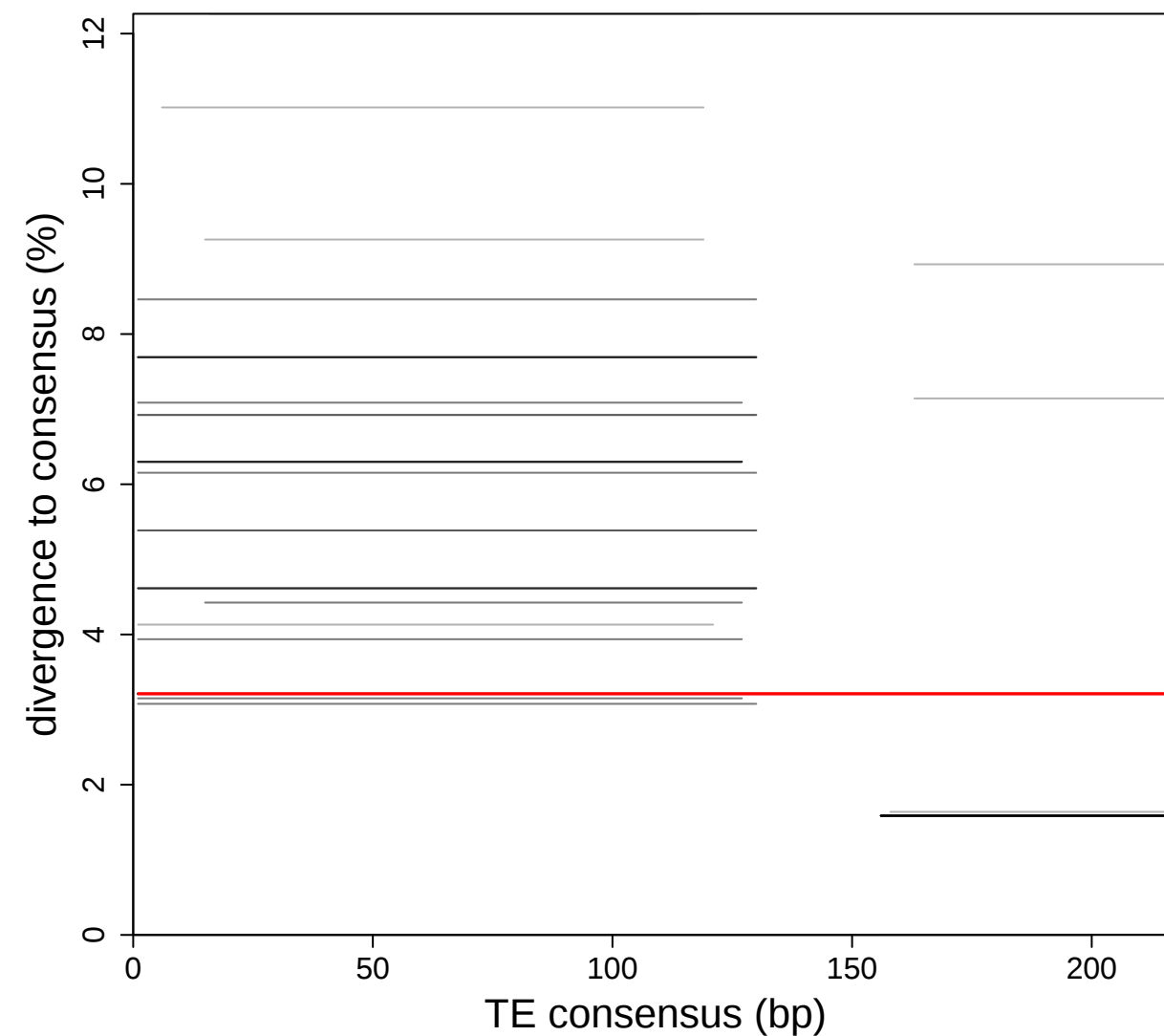


ily_38.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 4739bp; fragments: 107; full length: 17 (>=4265.1bp)

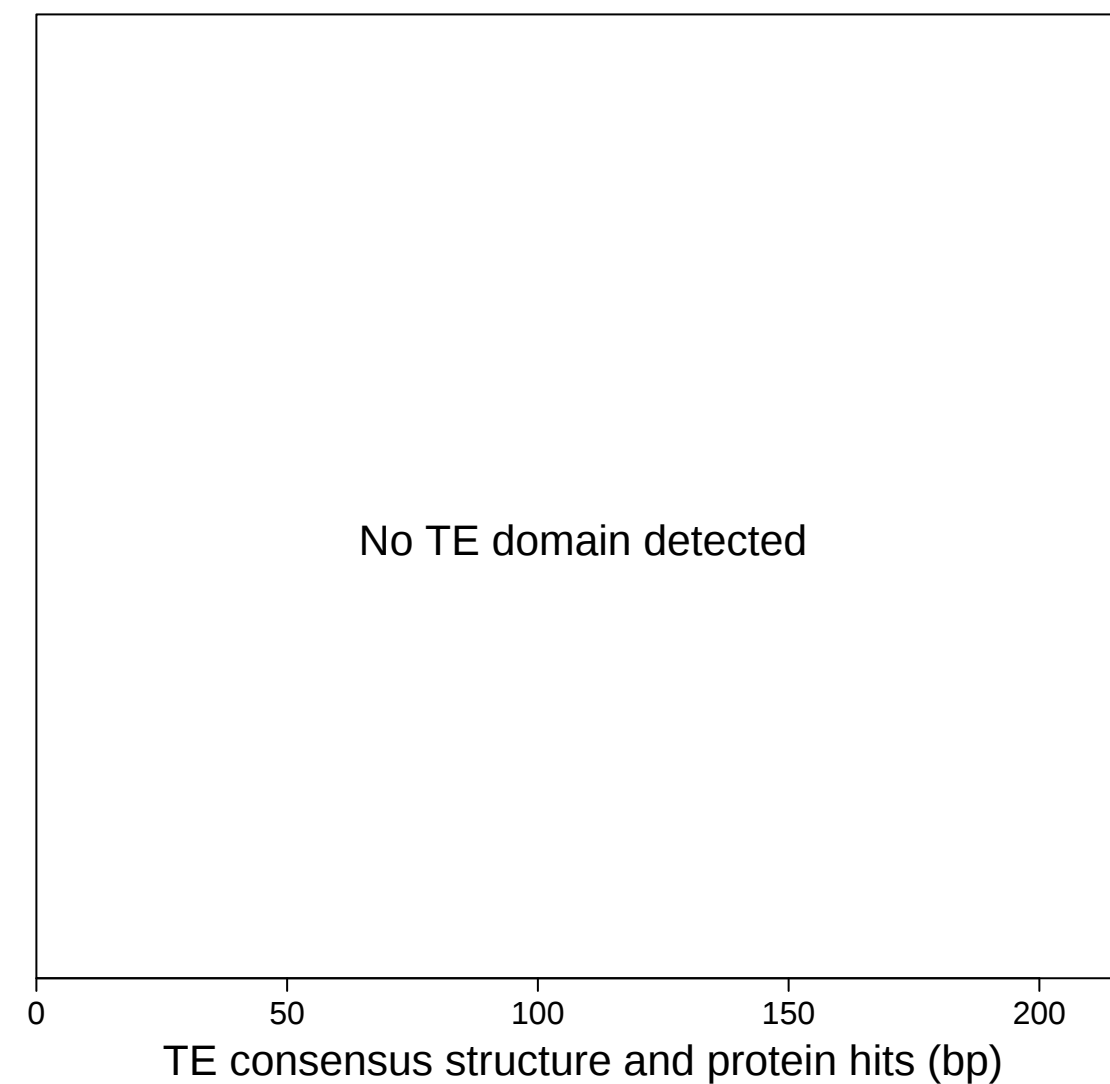
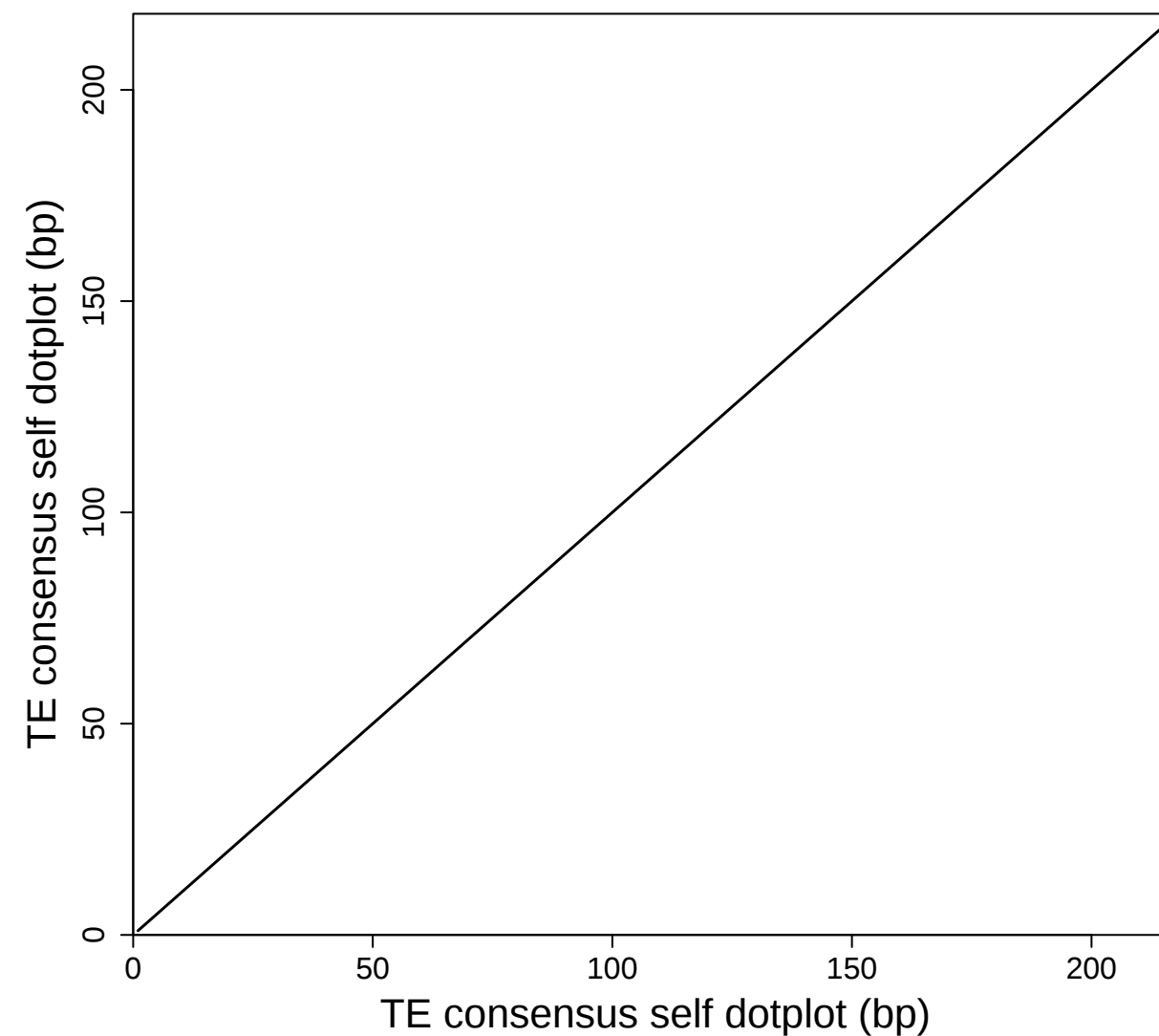
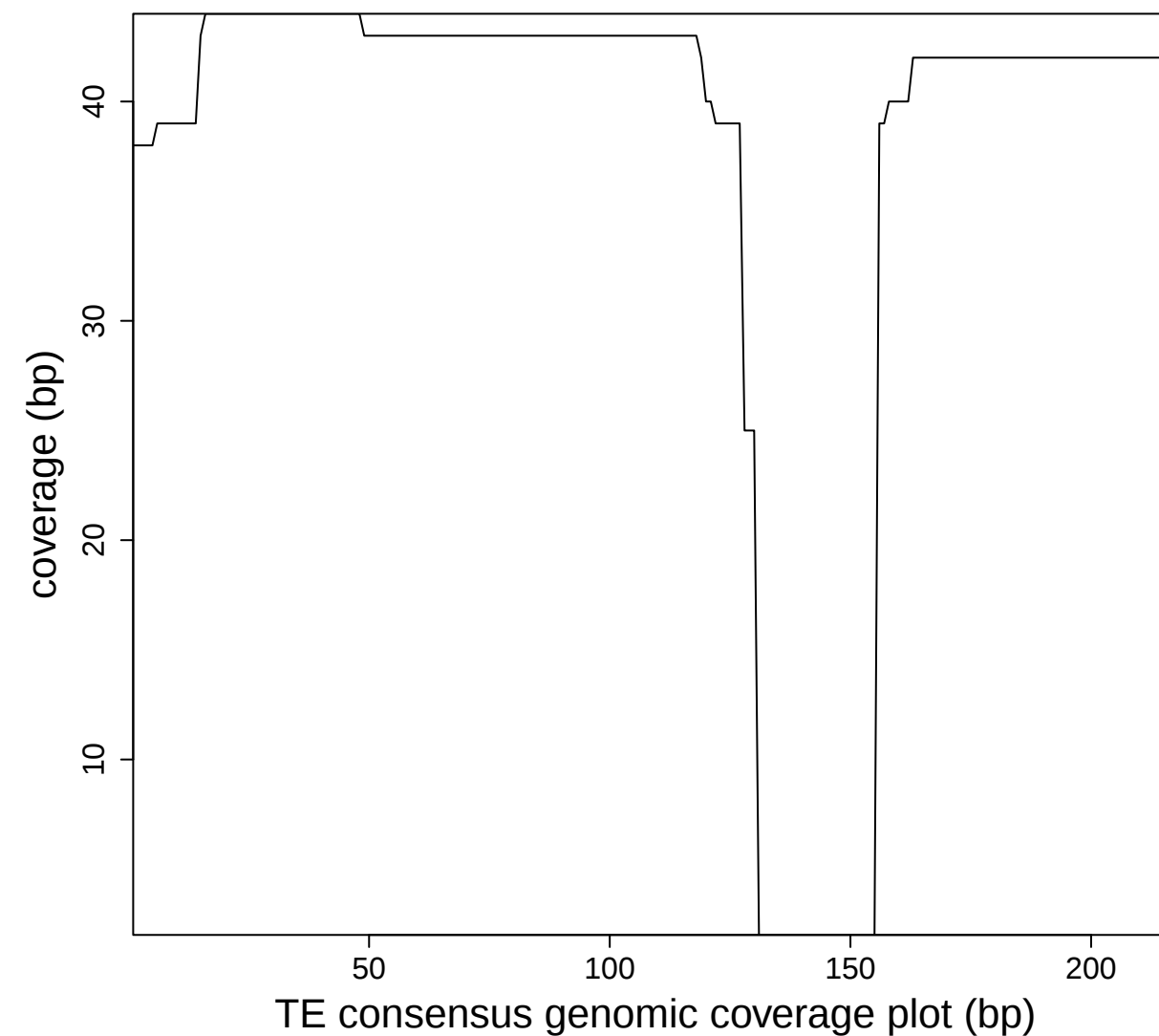
After TEtrimmer Extended plot Blue lines are boundaries



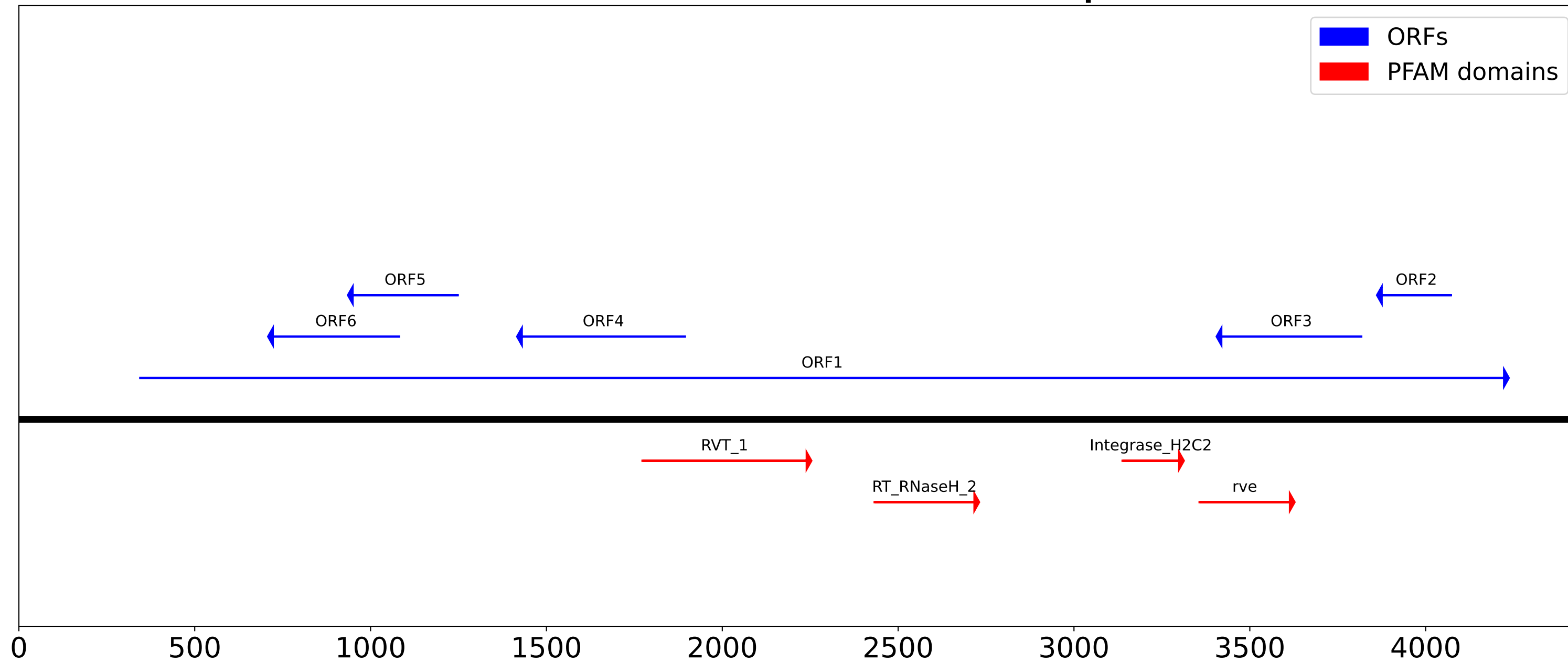
TE: ltr_1_family_38
size: 218bp; fragments: 84; full length: 2 (≥ 196.2 bp)



Before TEtrimmer 218 bp



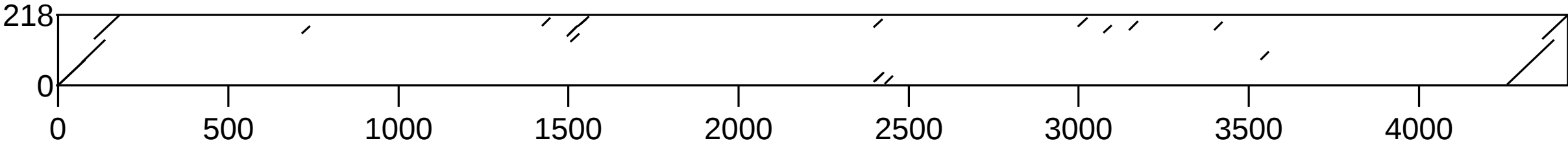
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)