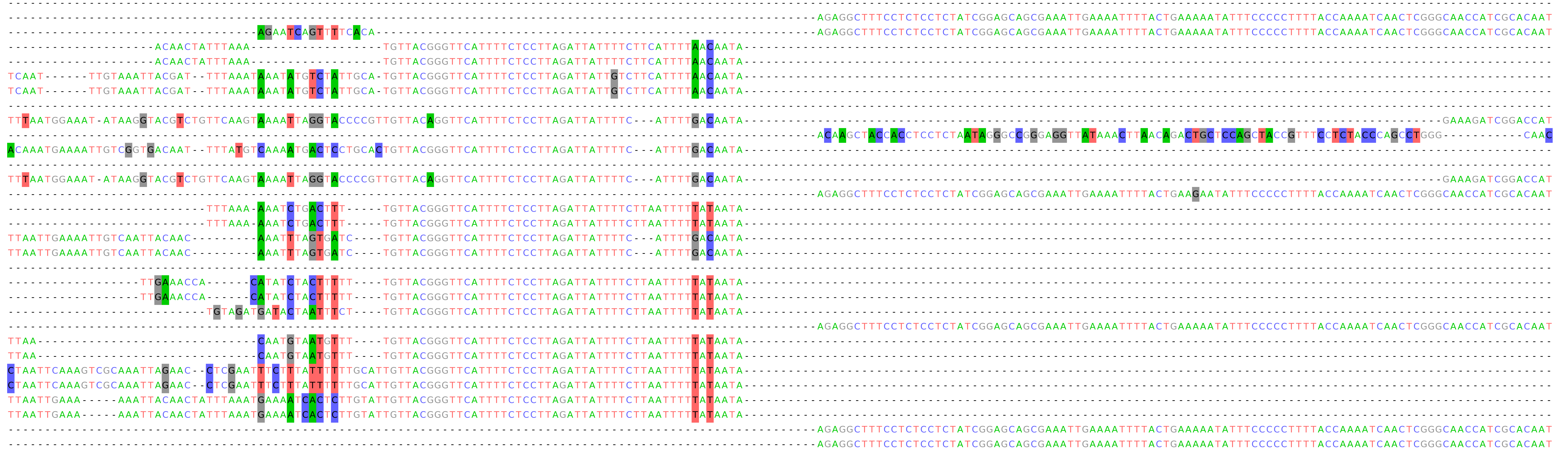
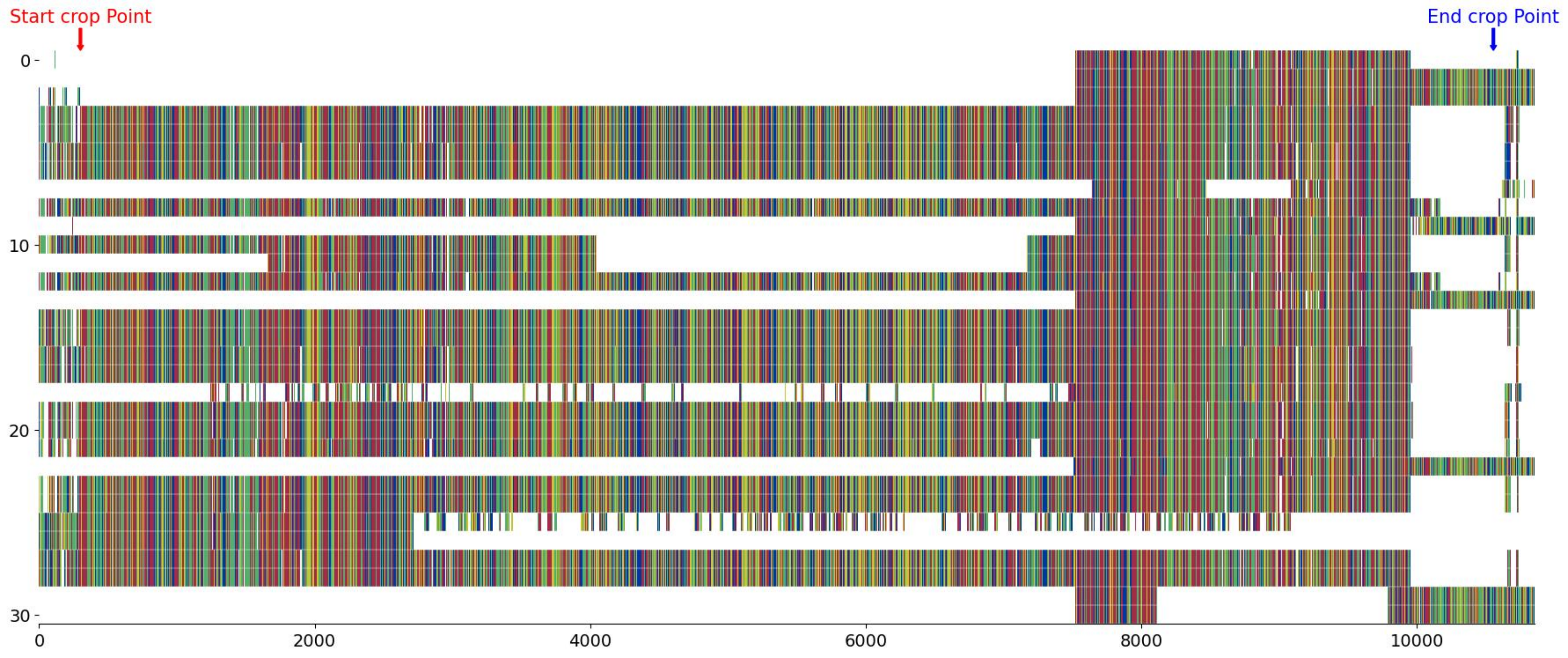


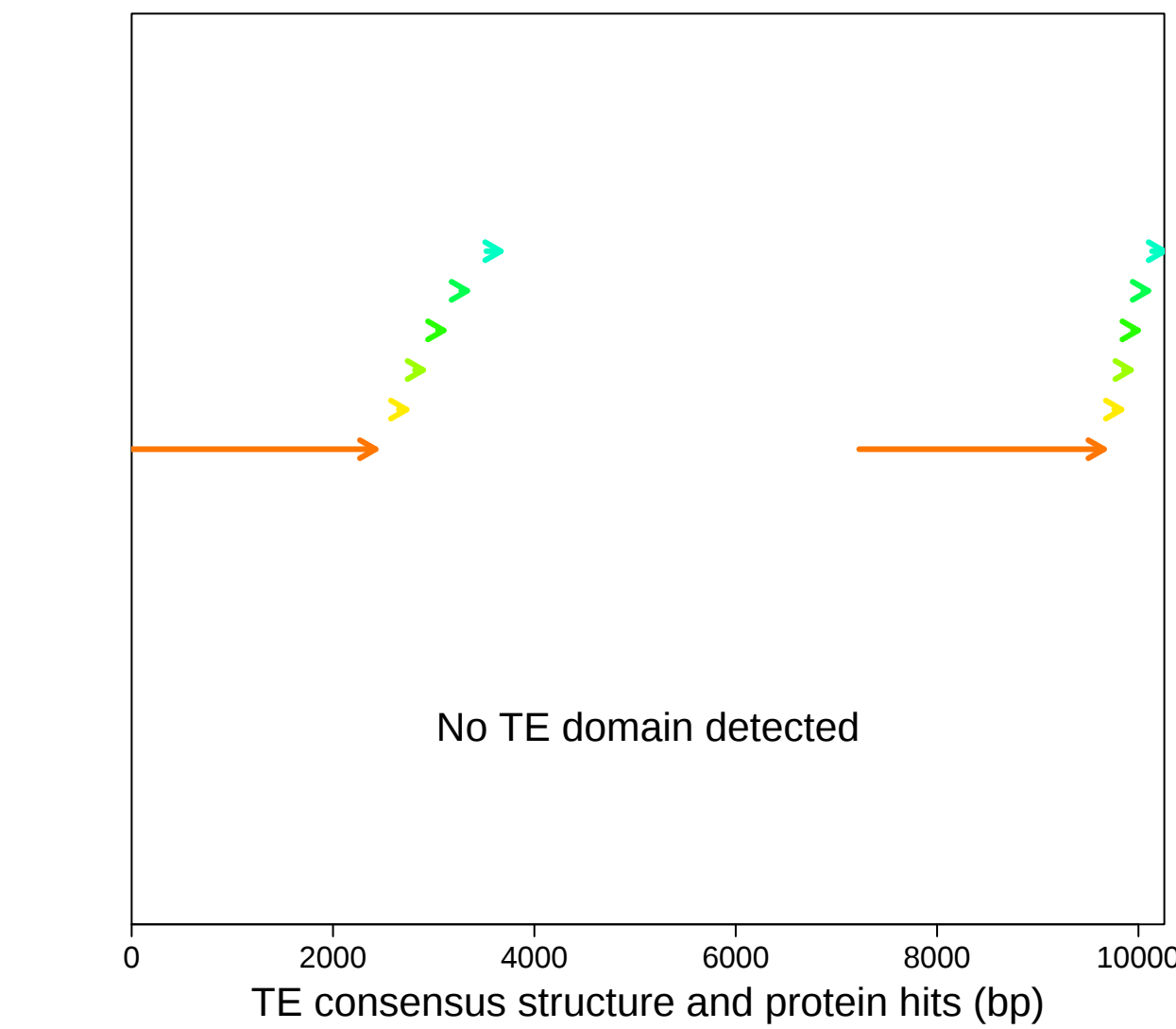
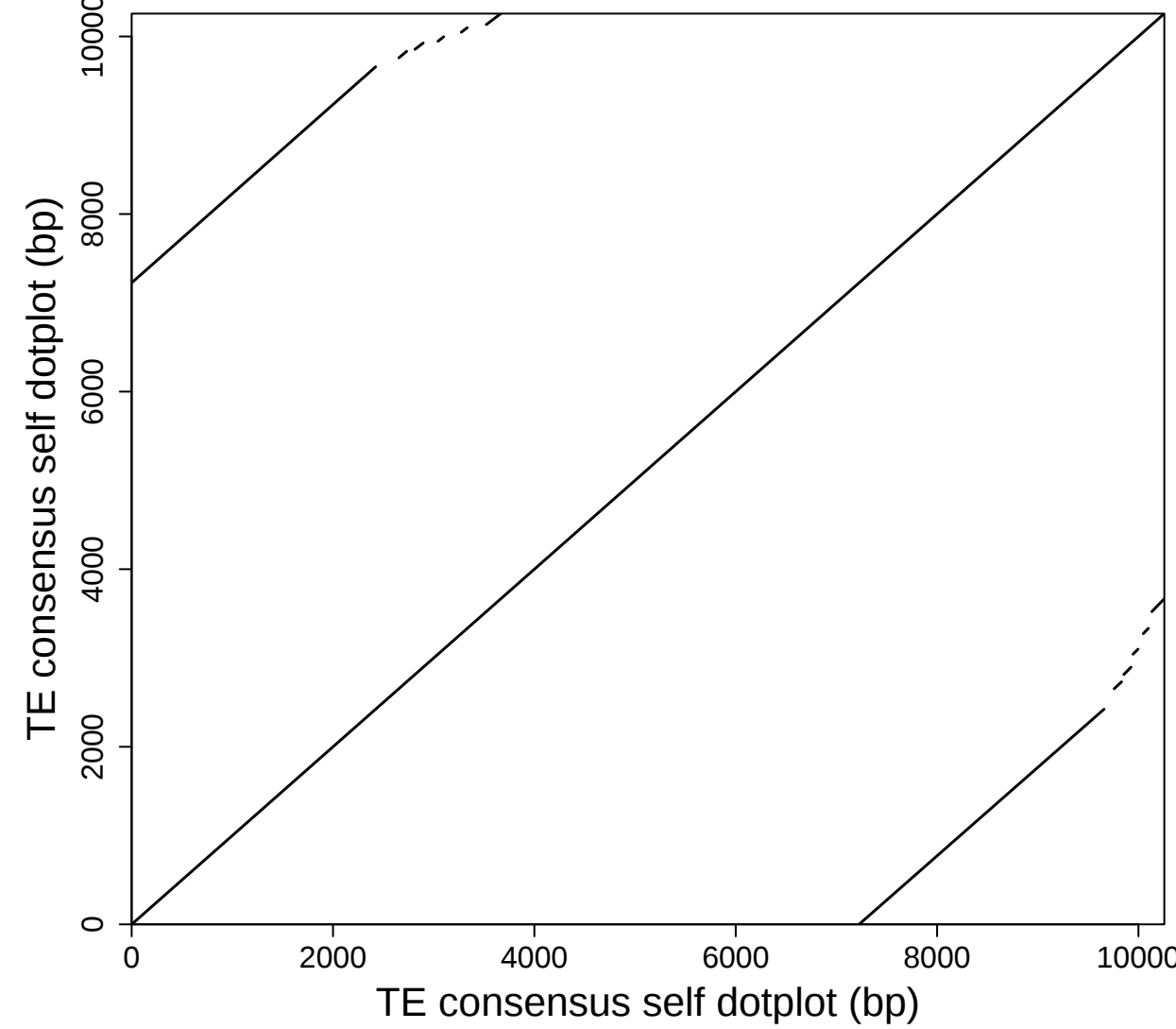
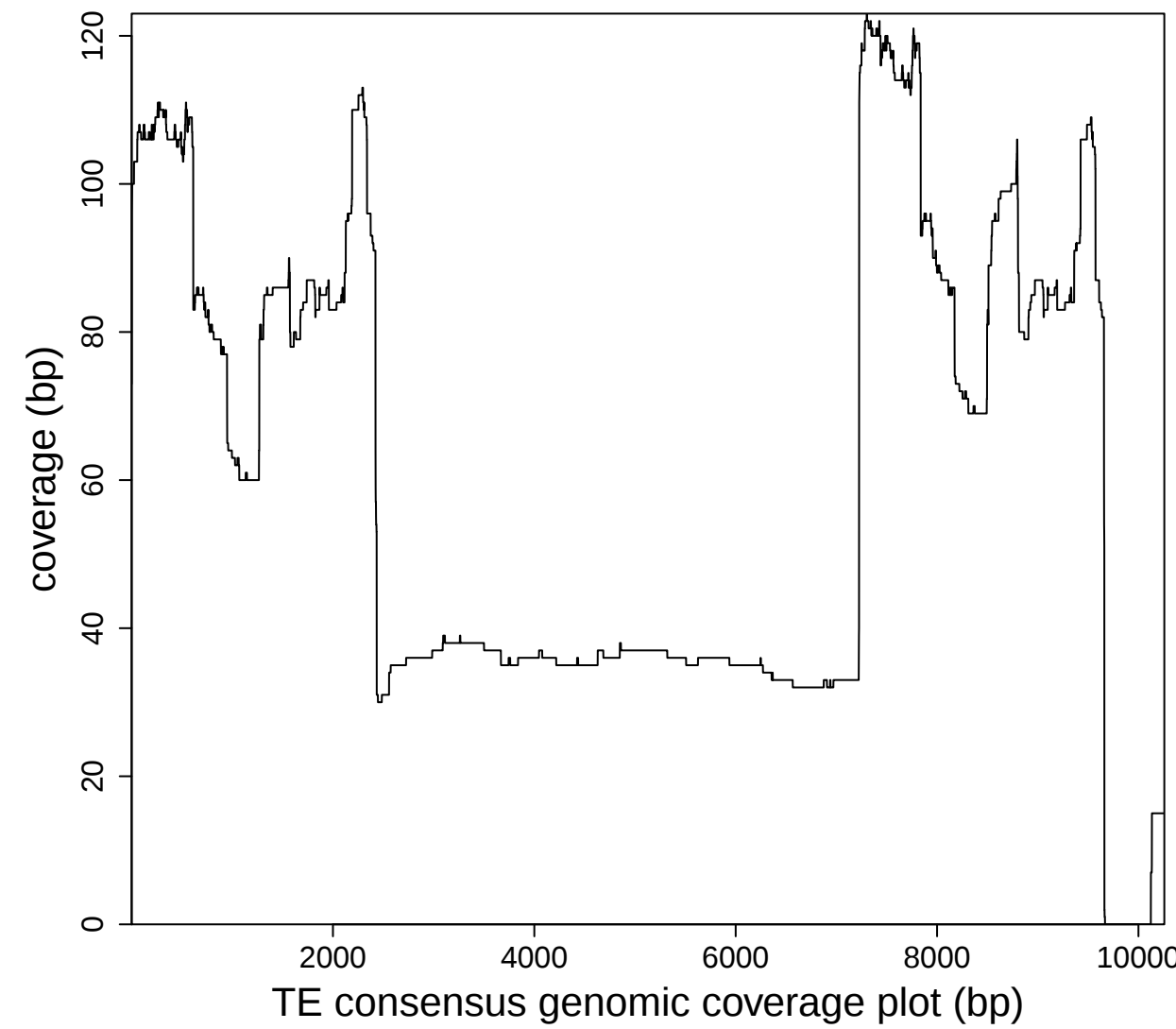
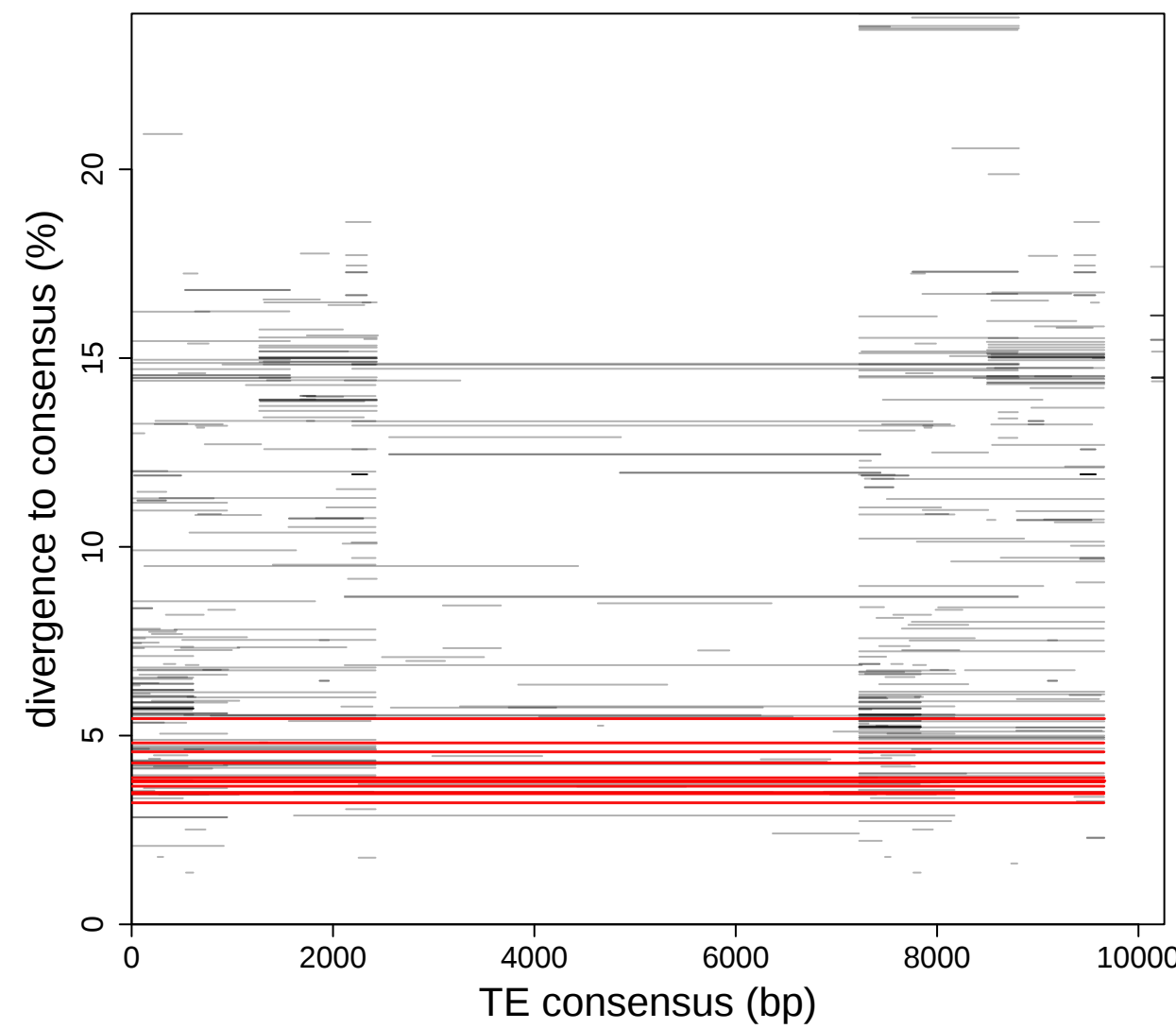
MSA length = 10257





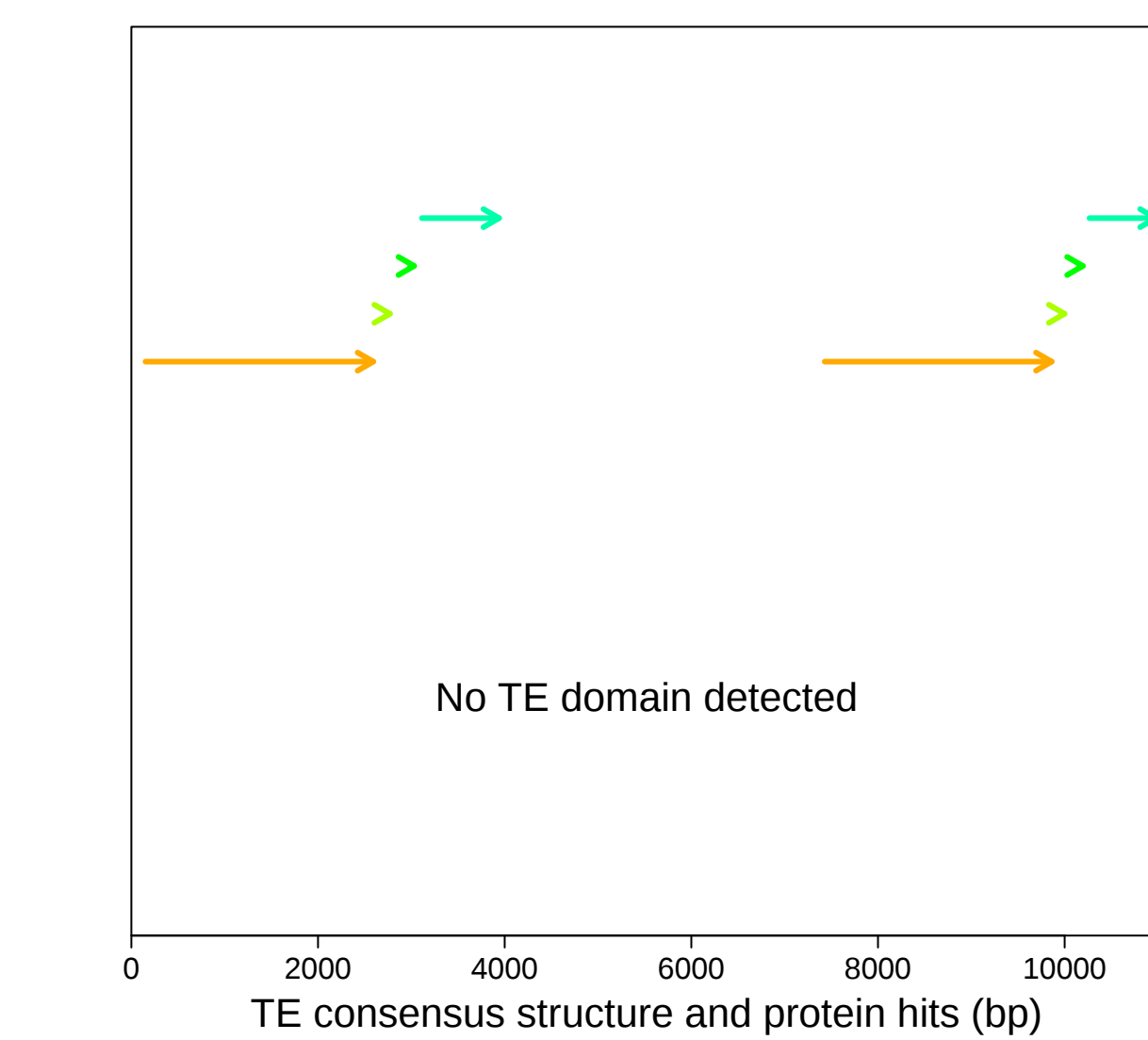
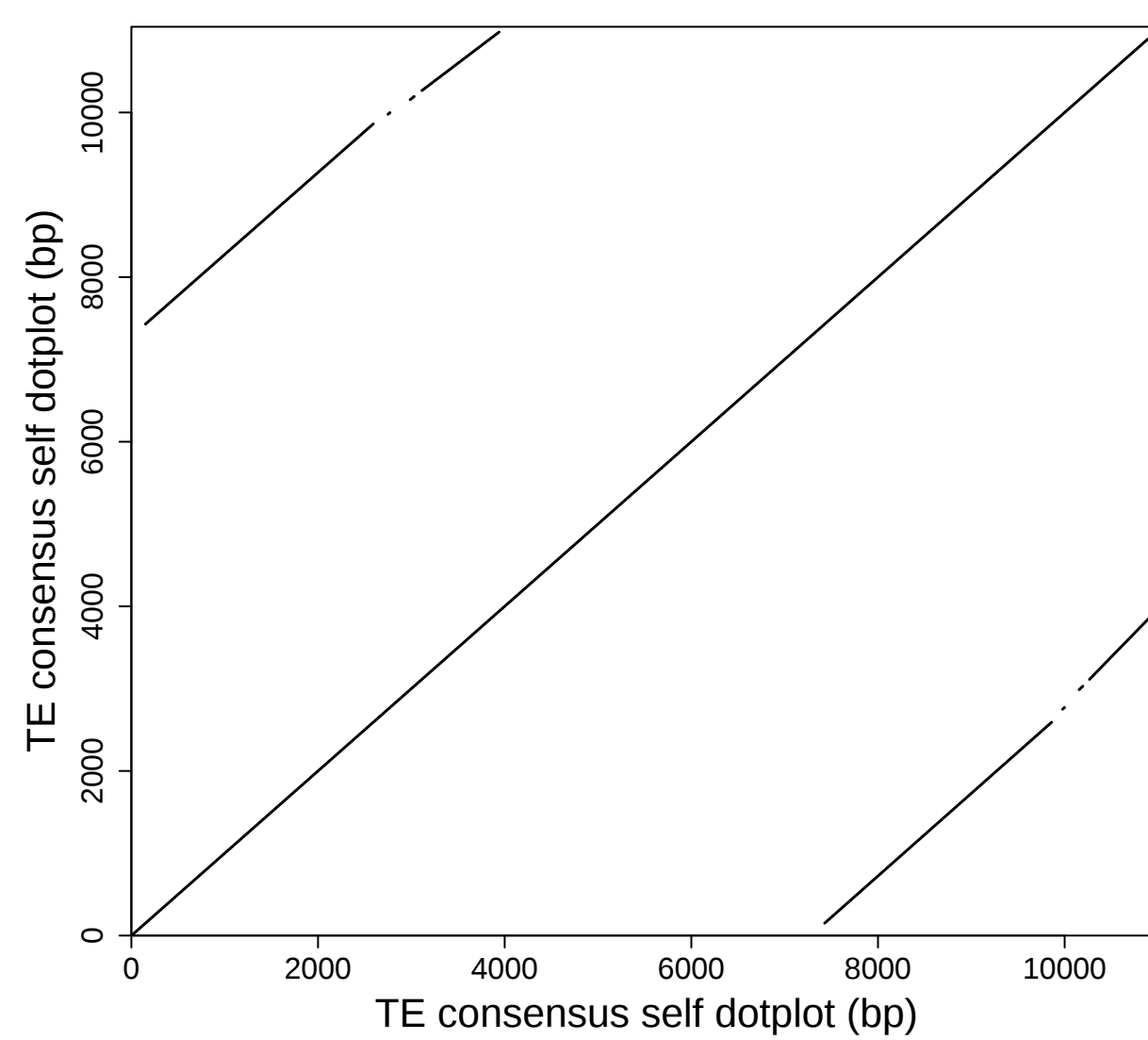
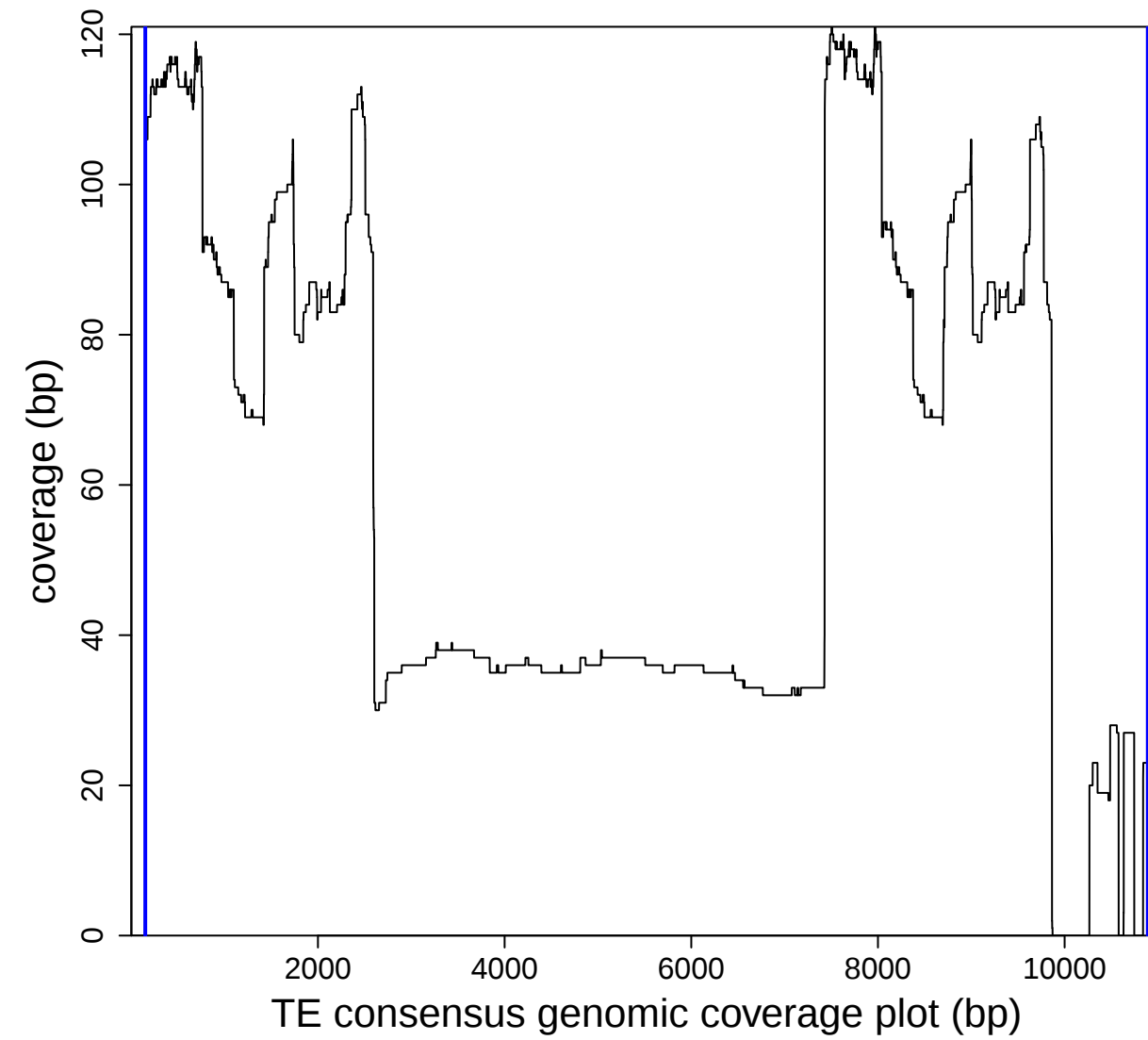
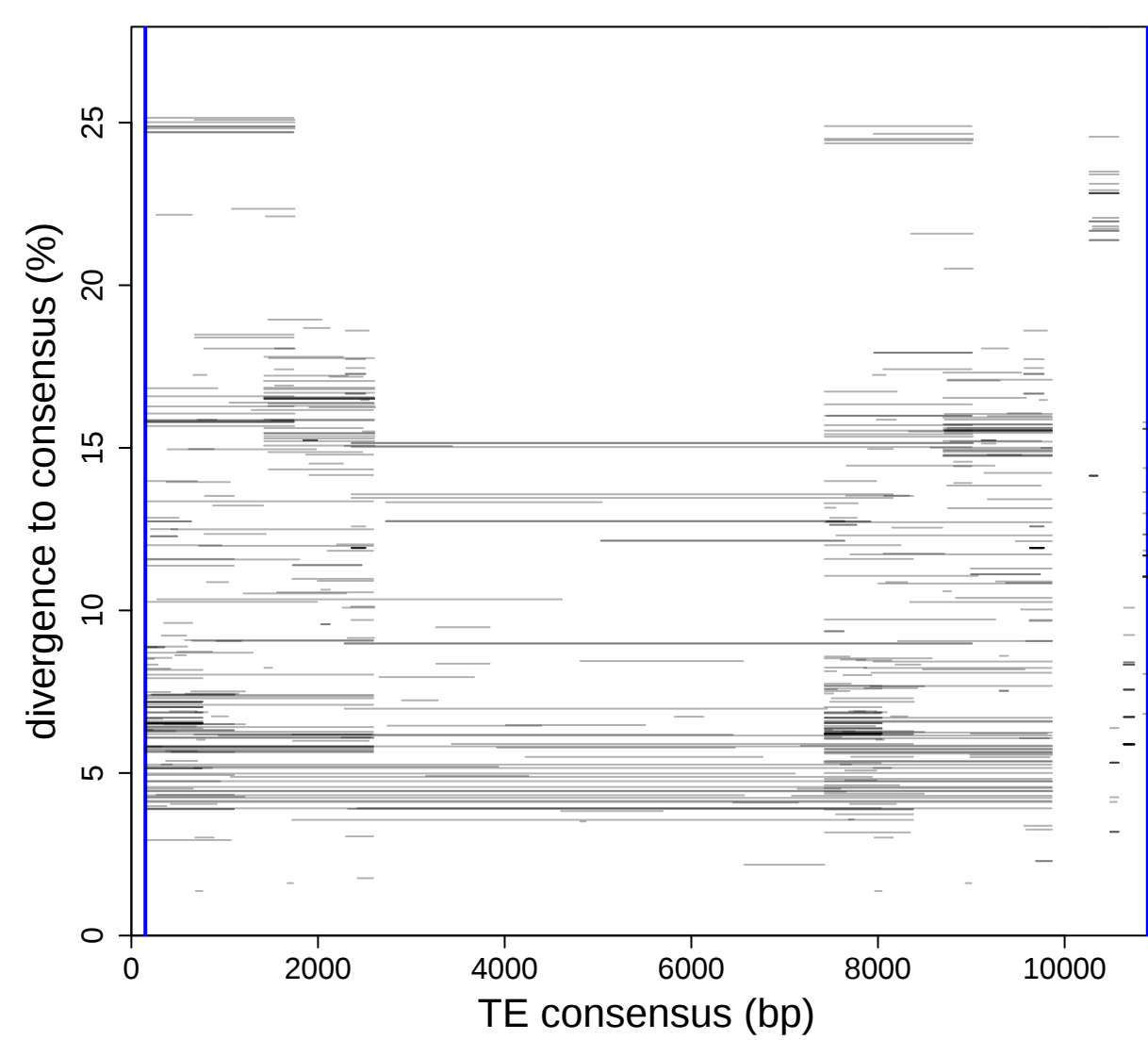
bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.fa
size: 10257bp; fragments: 503; full length: 11 (>=9231.3bp)

After TEtrimmer 10257 bp

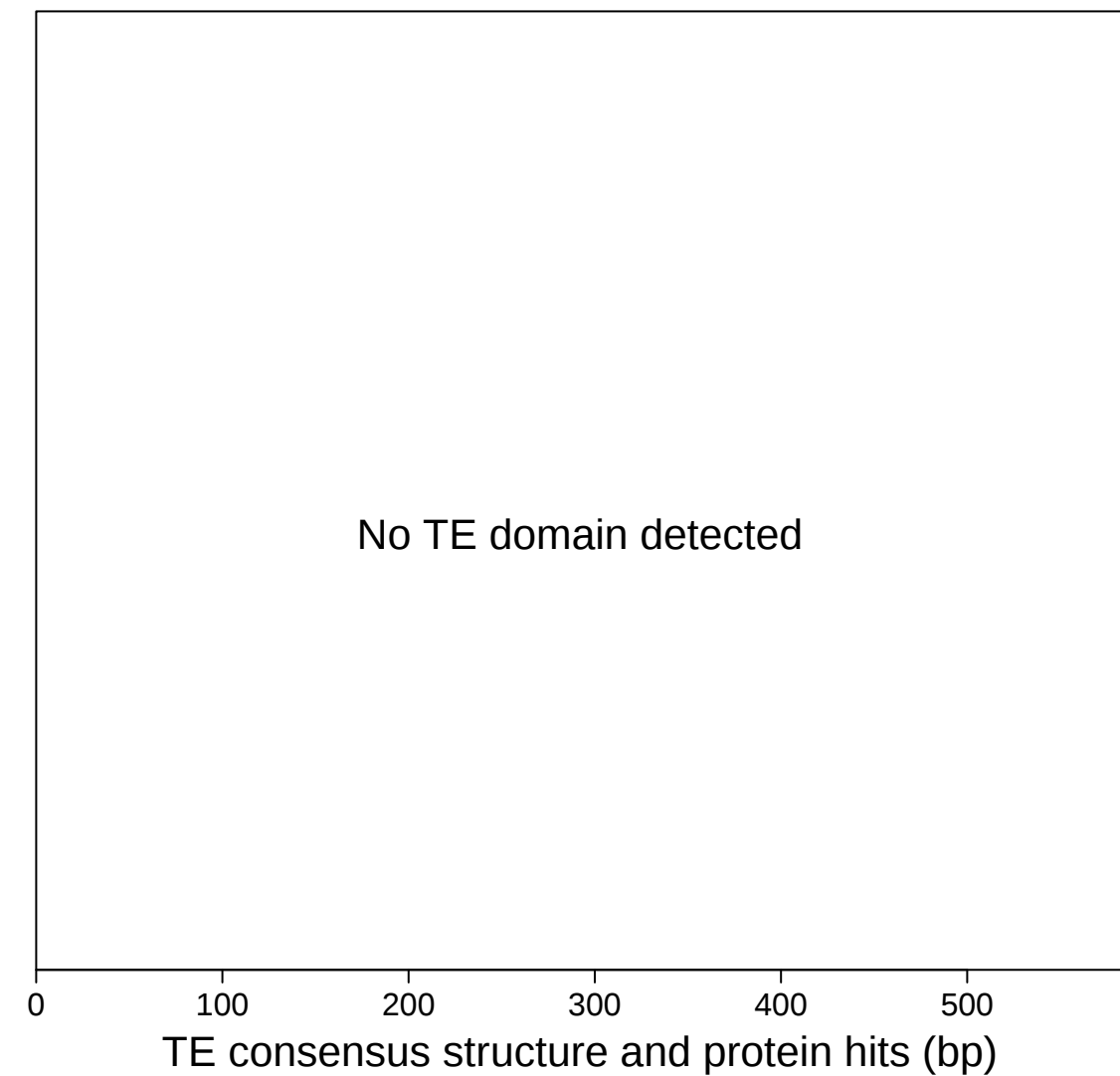
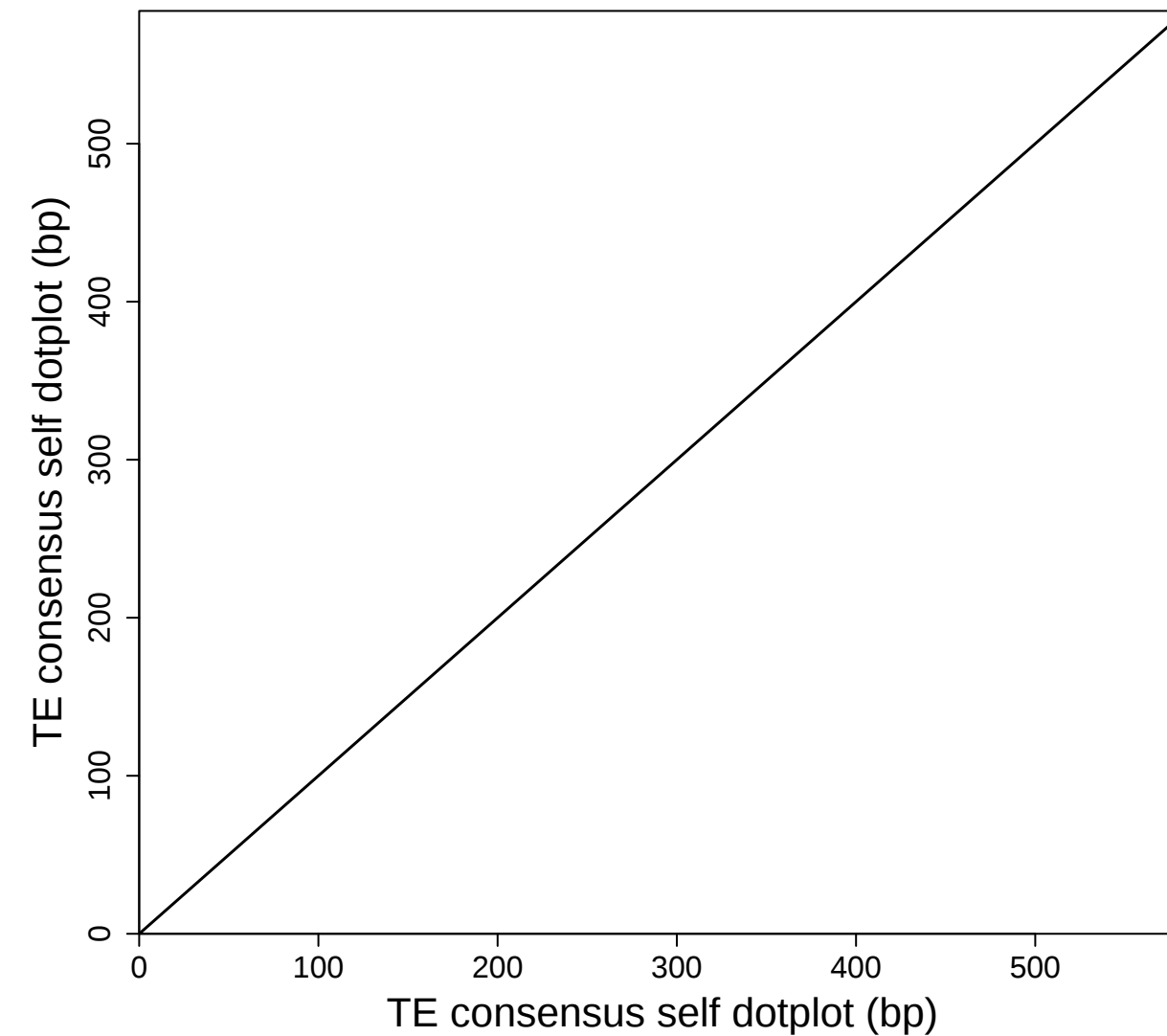
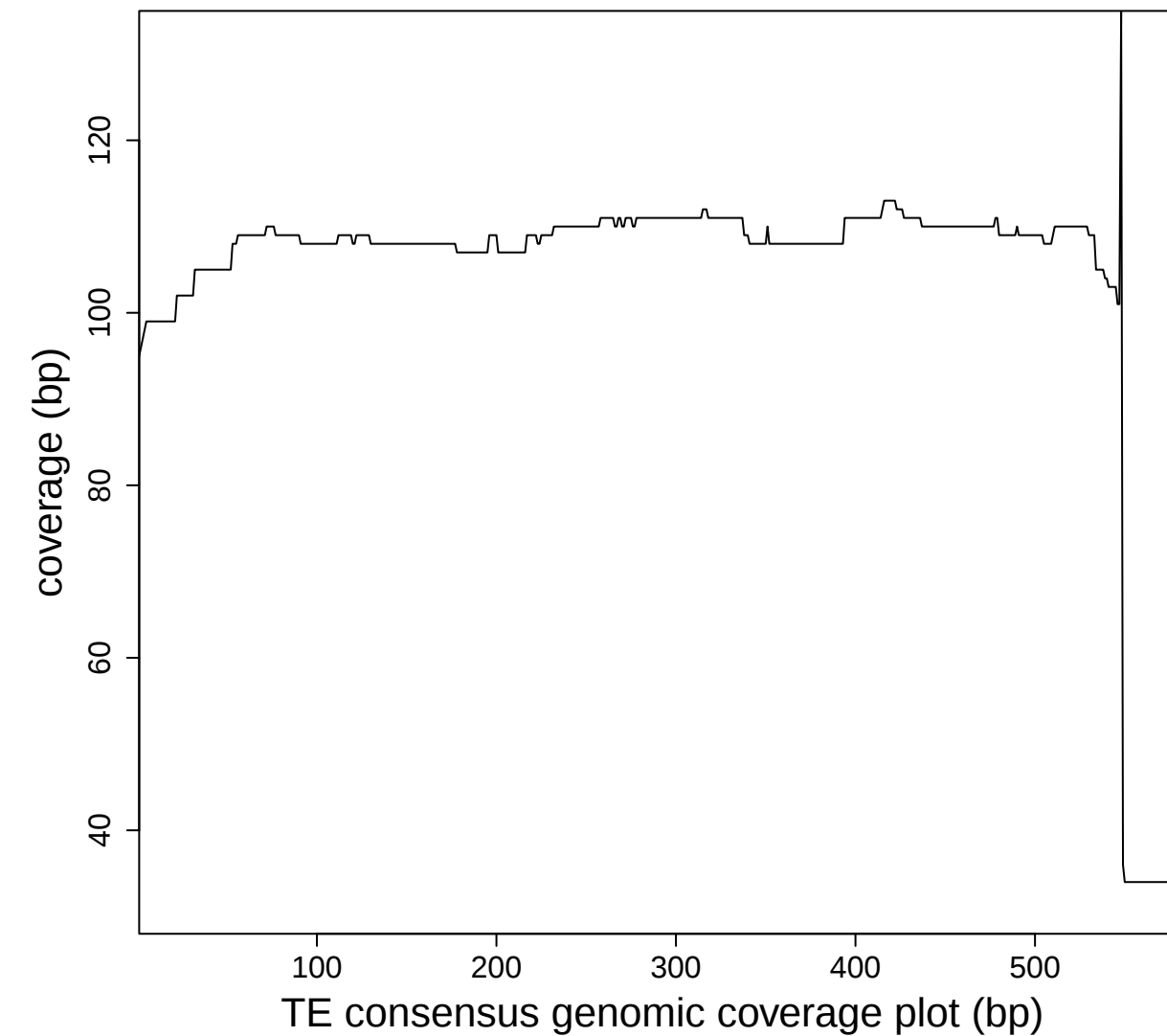
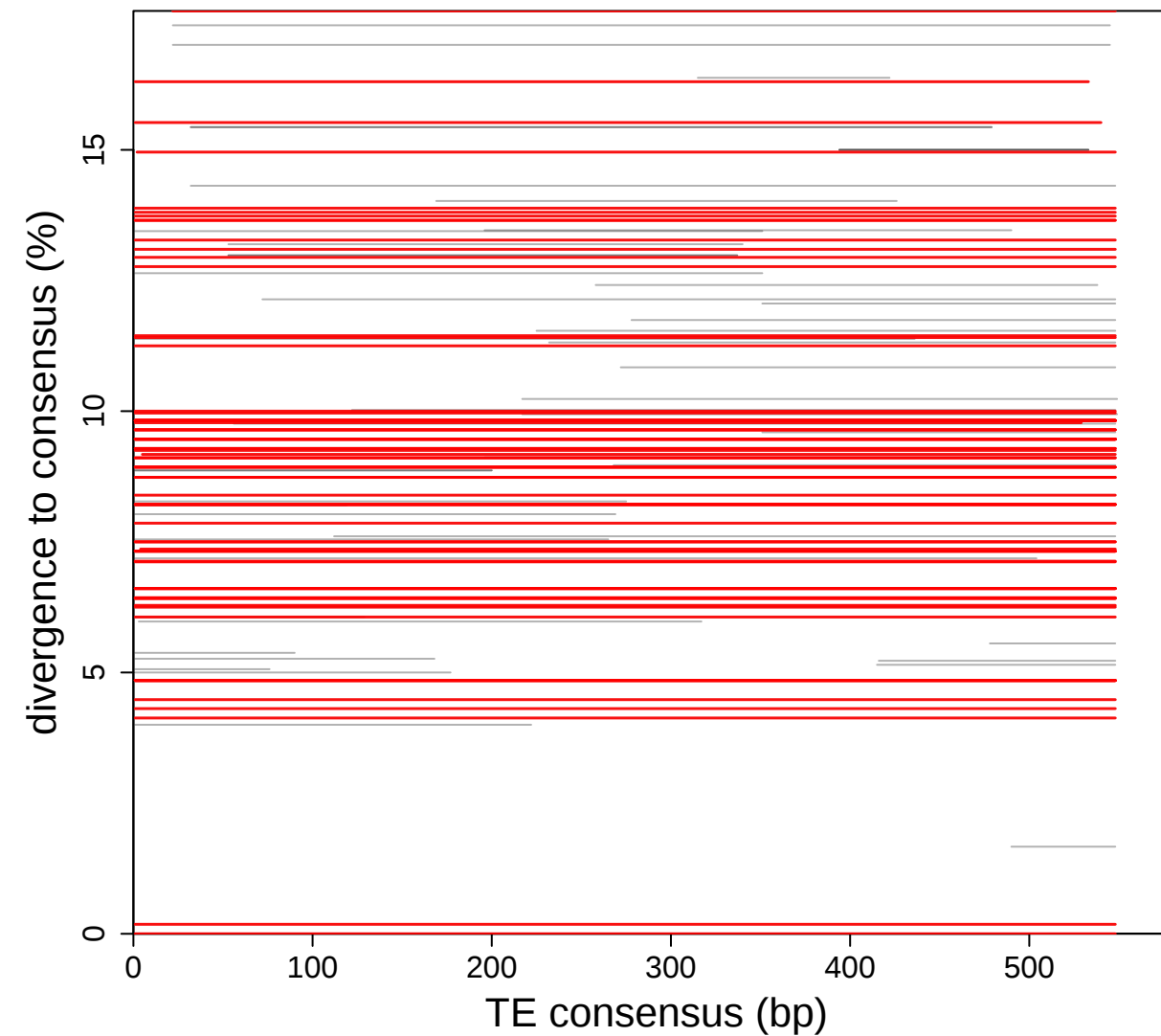


ed_uf.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0
size: 11041bp; fragments: 579; full length: 0 (>=9936.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

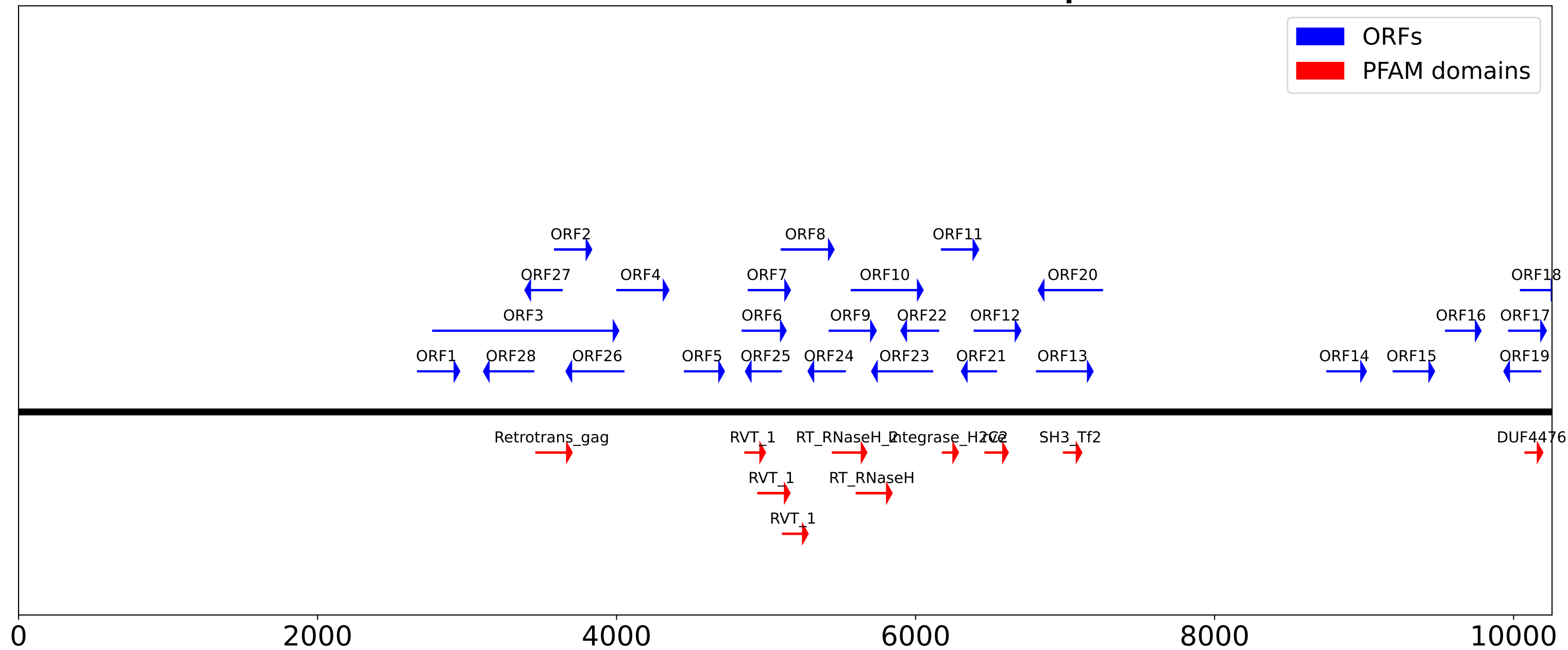


TE: ltr_1_family_354
size: 584bp; fragments: 169; full length: 83 (≥ 525.6 bp)



Before TEtrimmer 584 bp

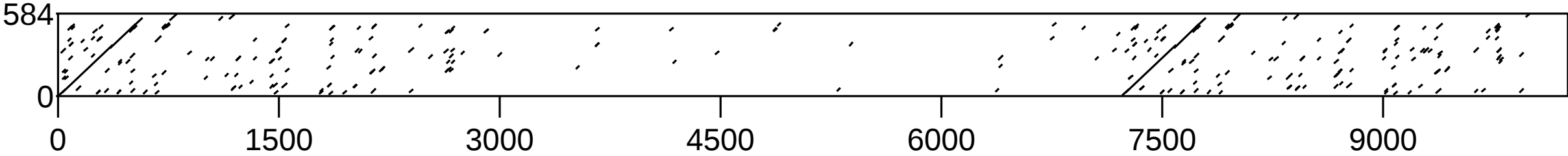
After TETrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)