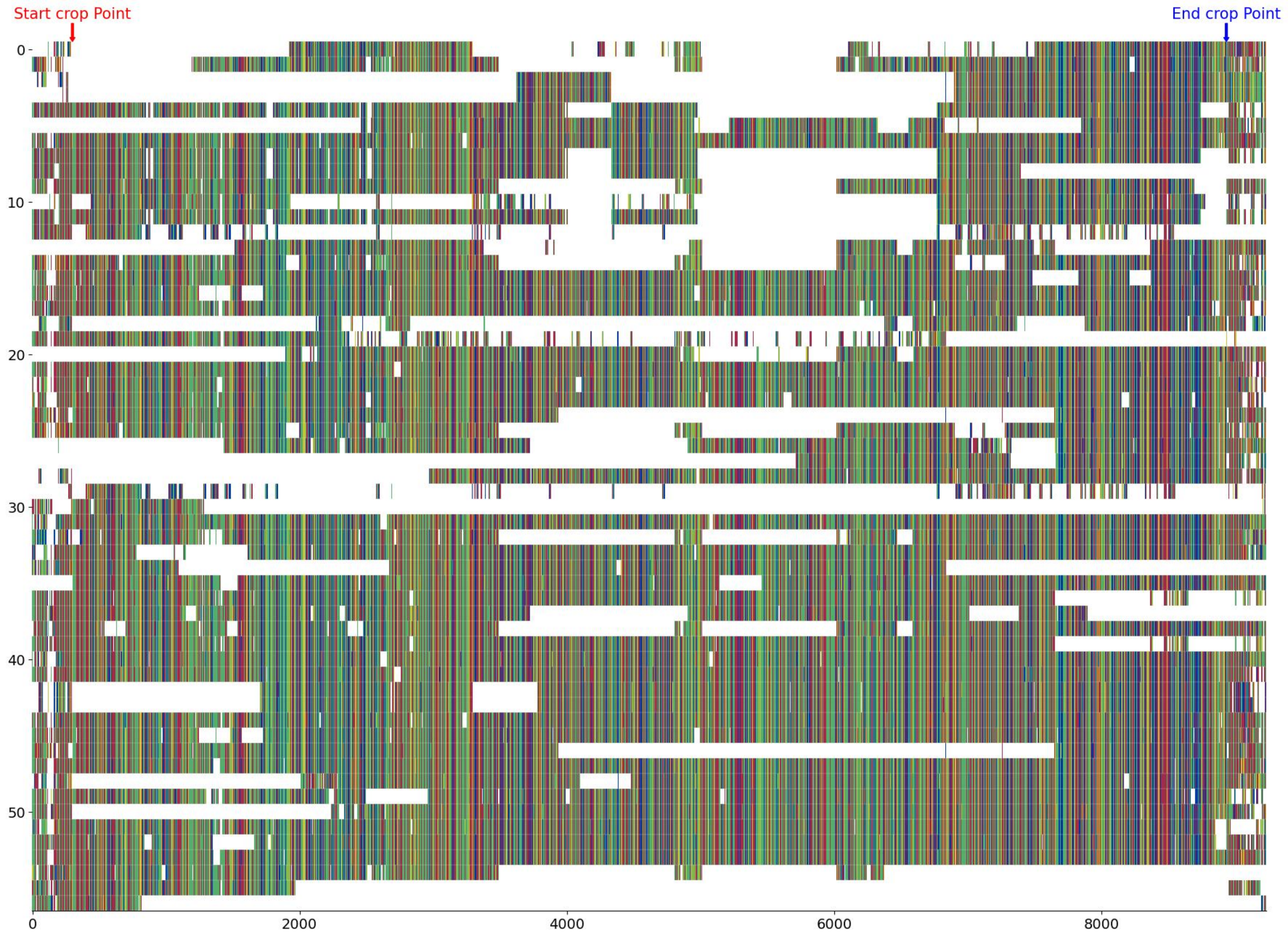




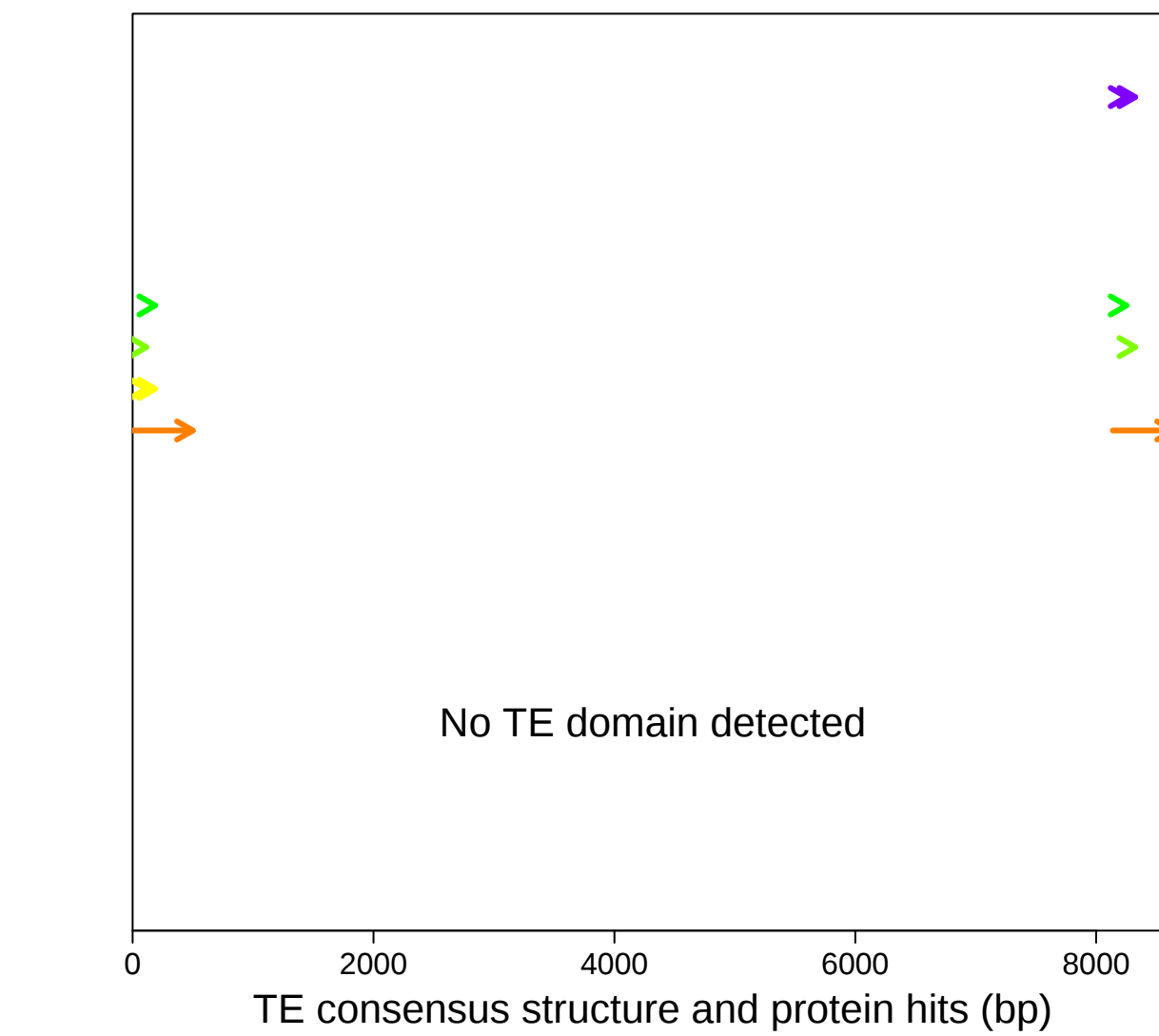
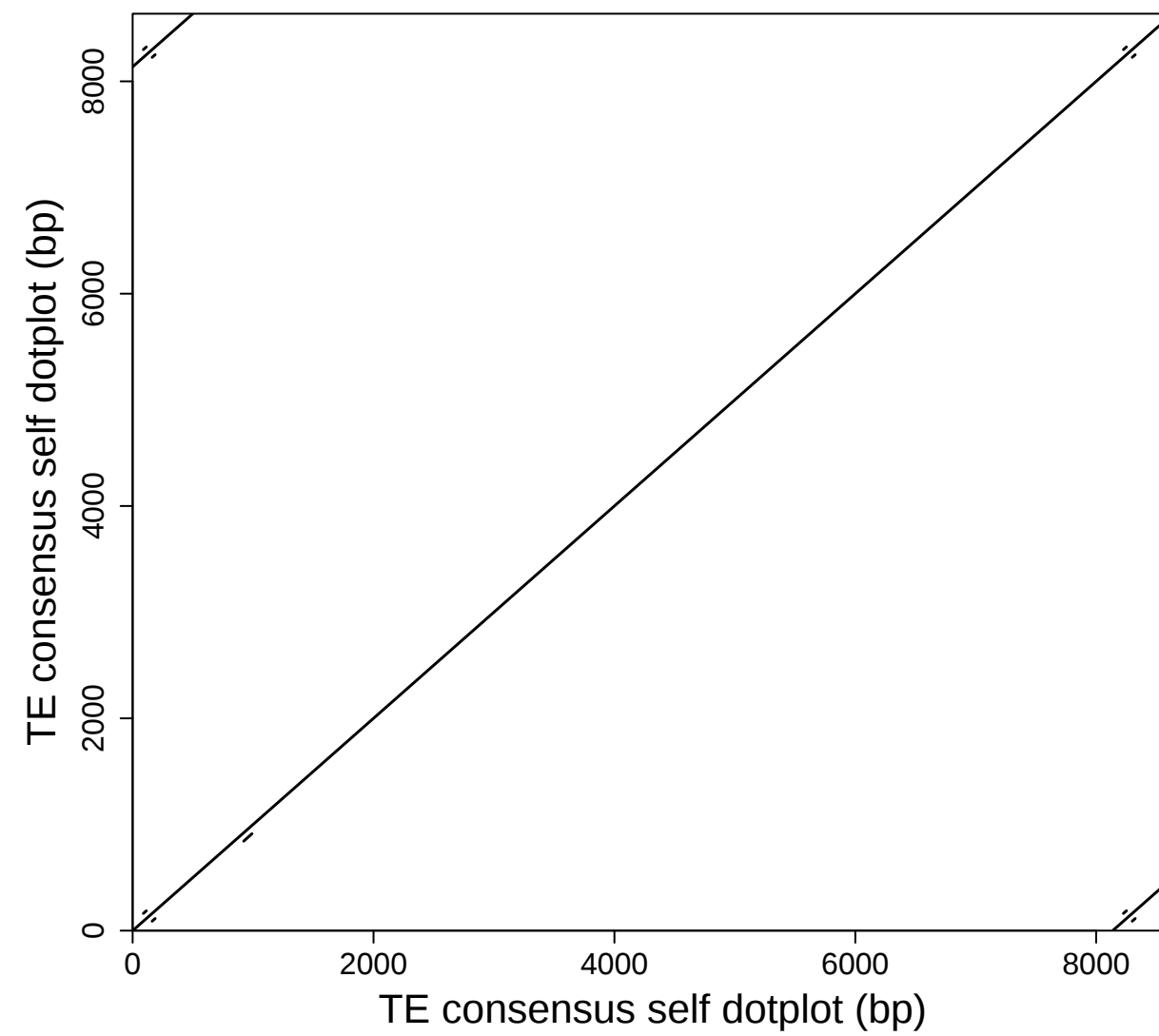
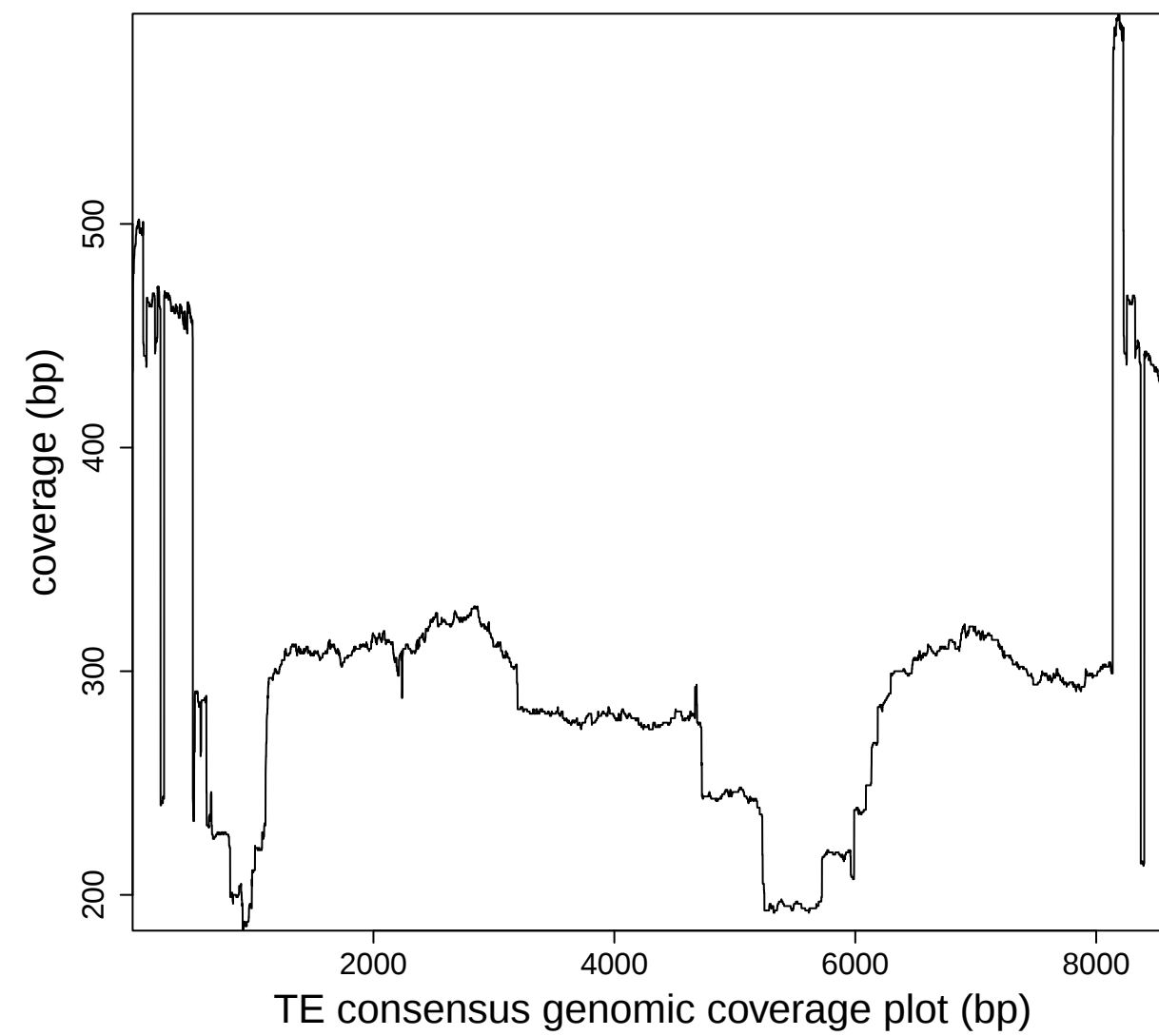
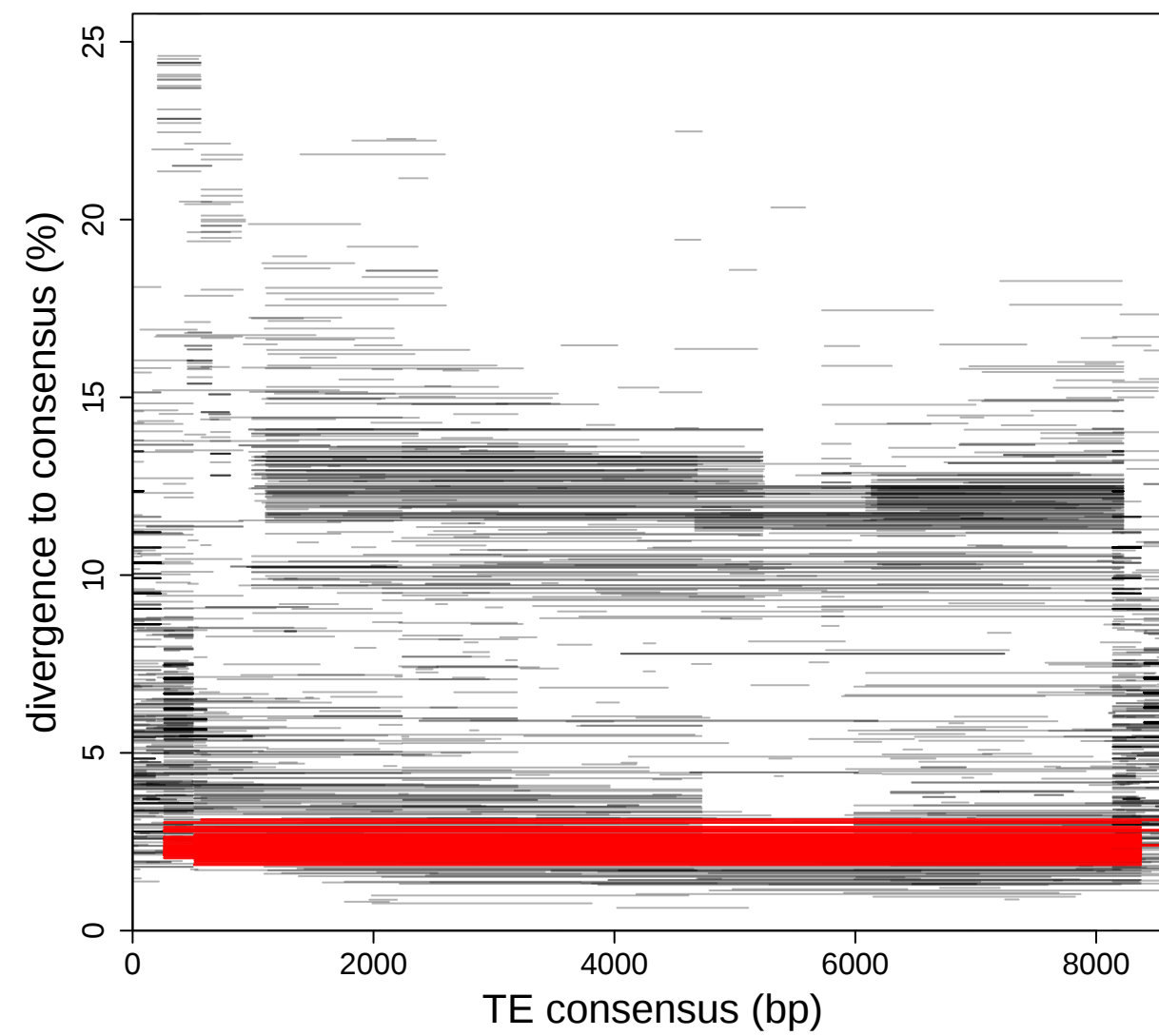
MSA length = 8637

[illegible]



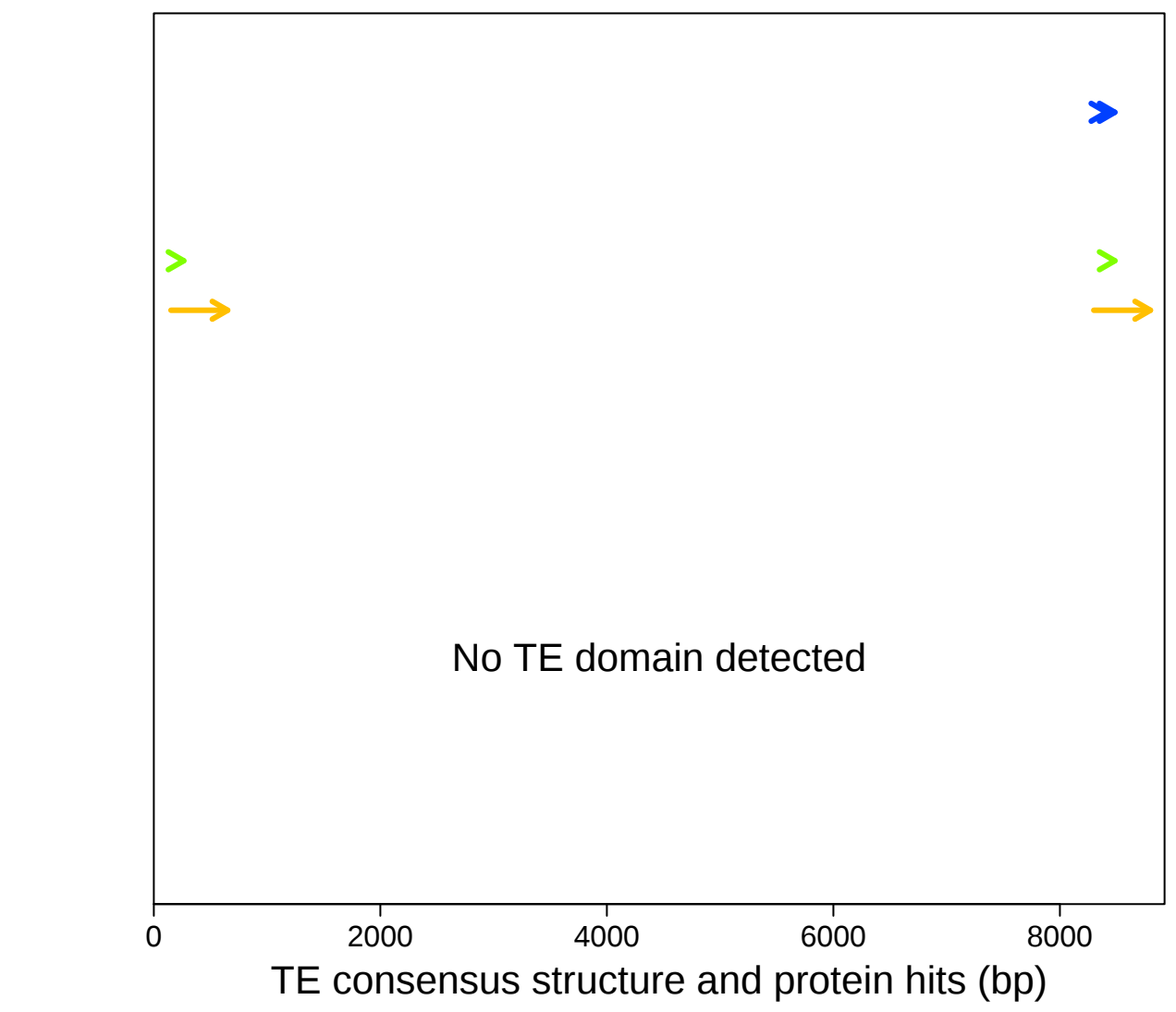
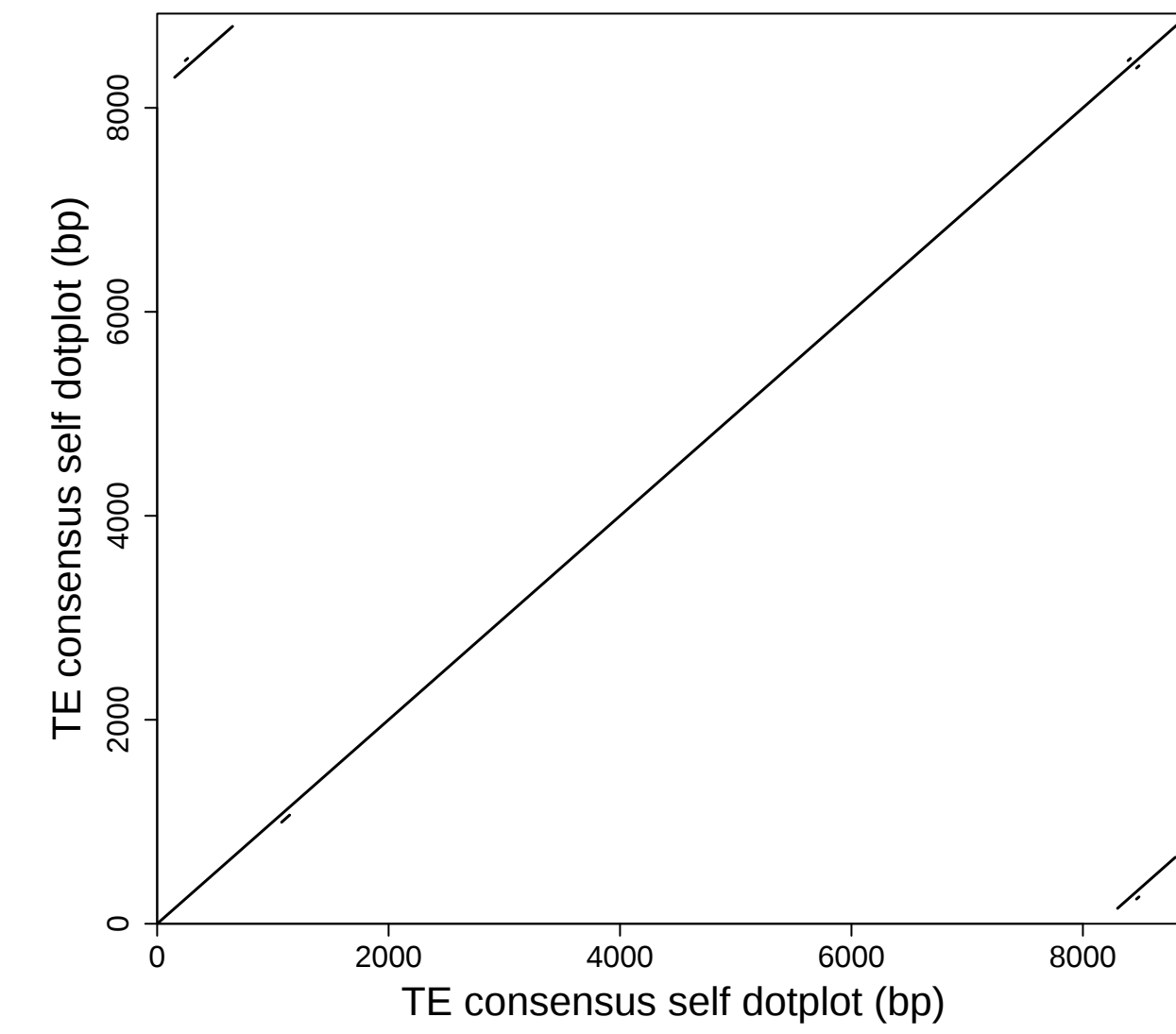
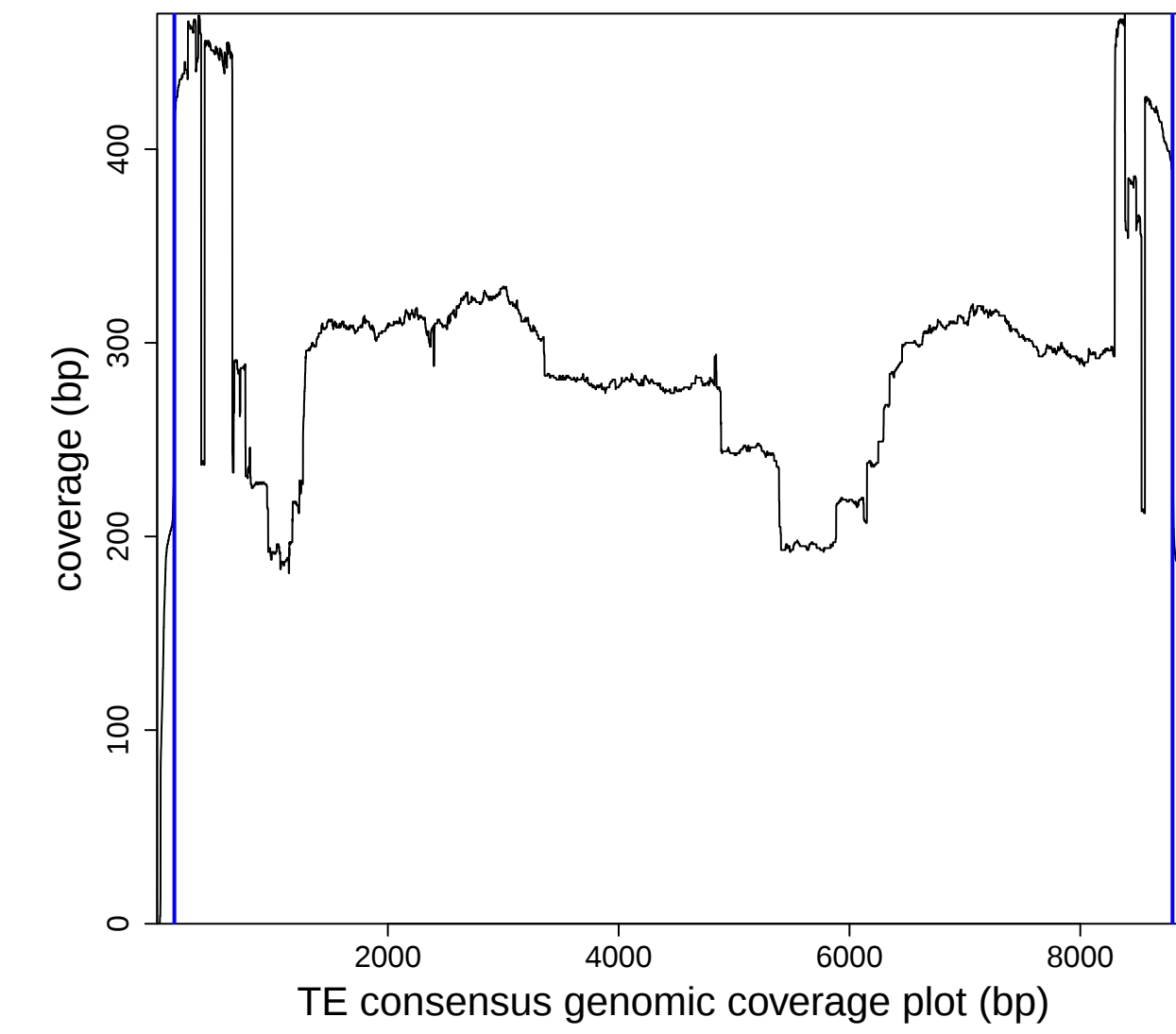
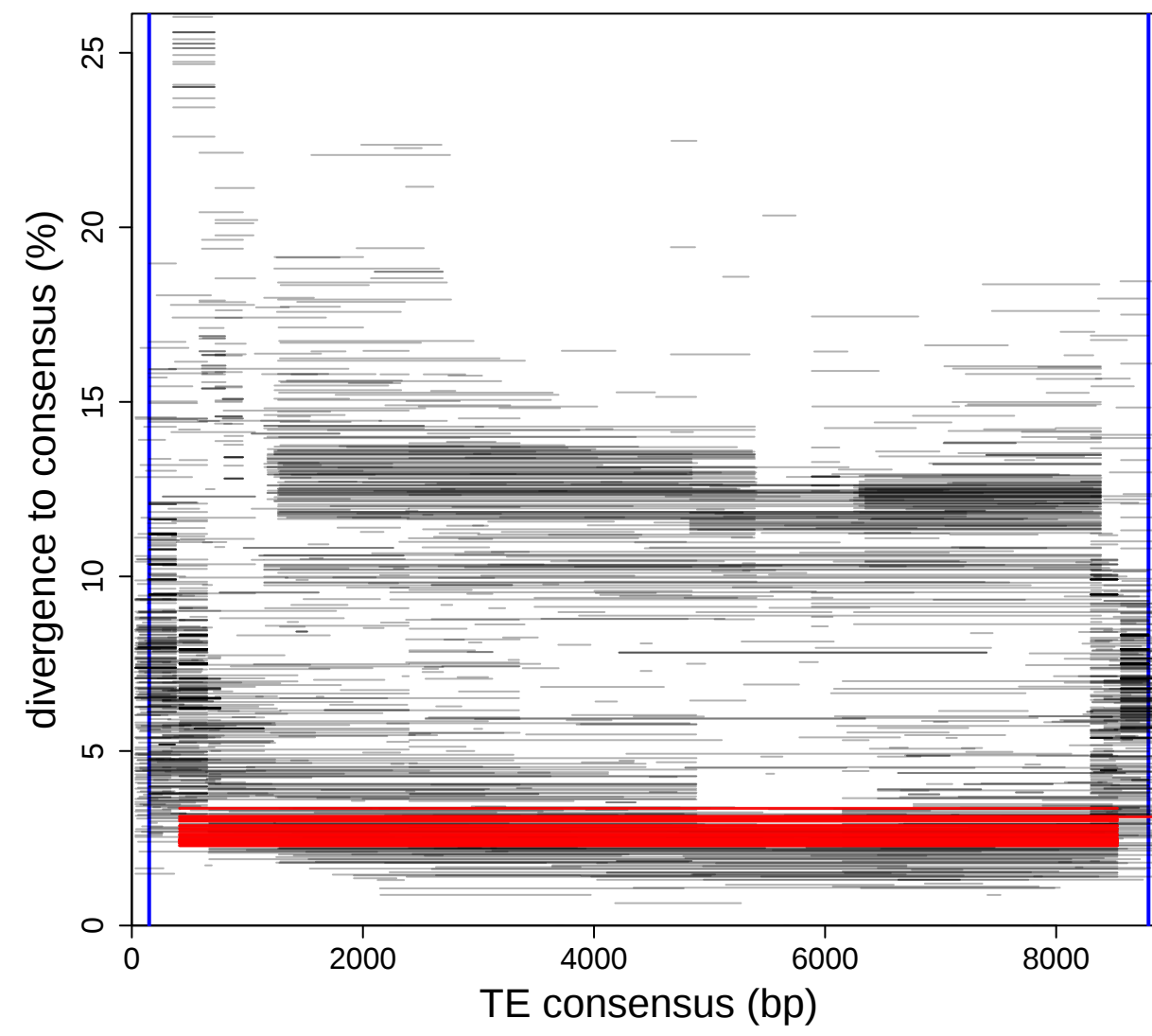
bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.faaln.fa
size: 8637bp; fragments: 2432; full length: 54 (>=7773.3bp)

After TEtrimmer 8637 bp

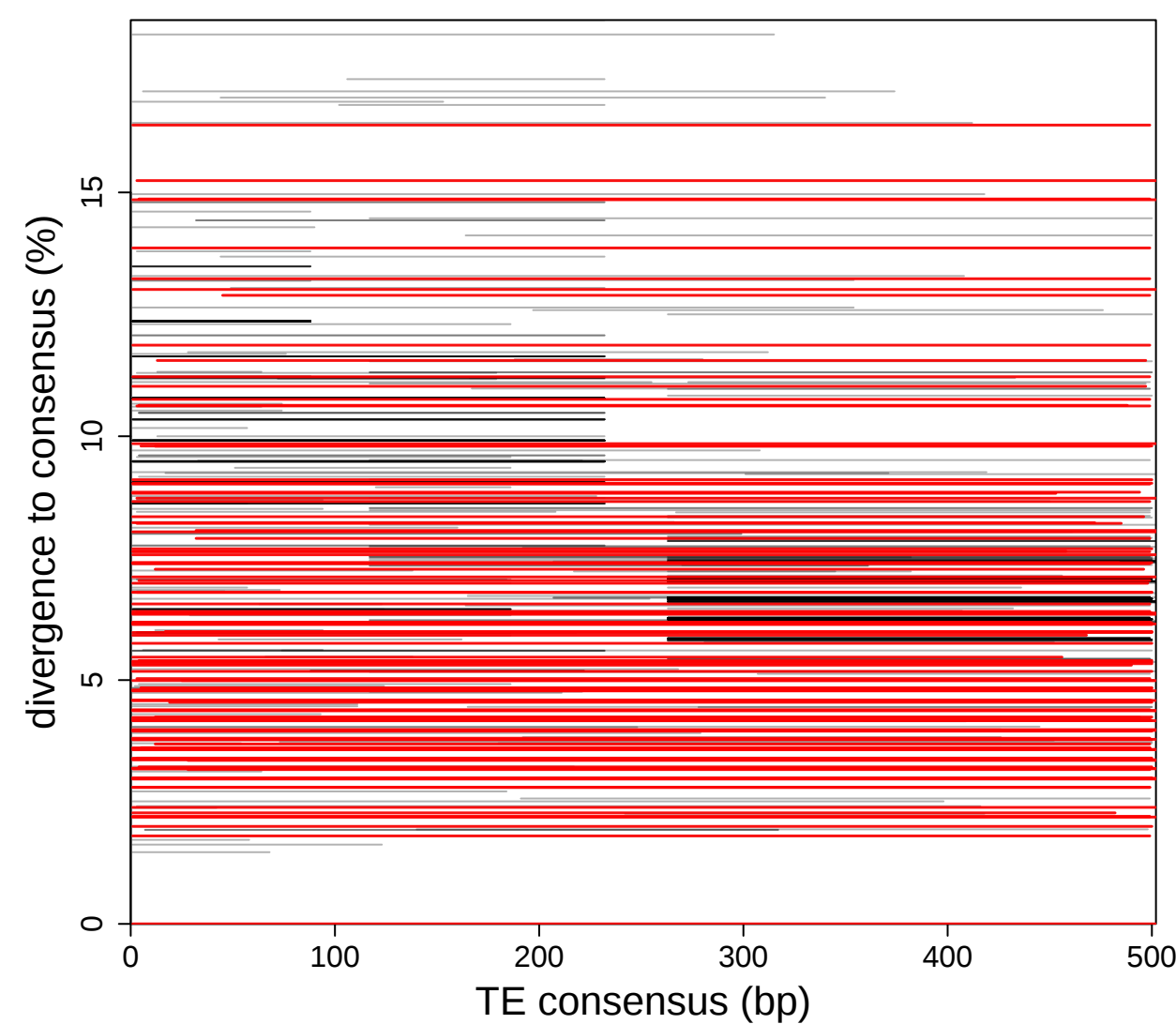


ed_uf.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_
size: 8949bp; fragments: 2208; full length: 28 (>=8054.1bp)

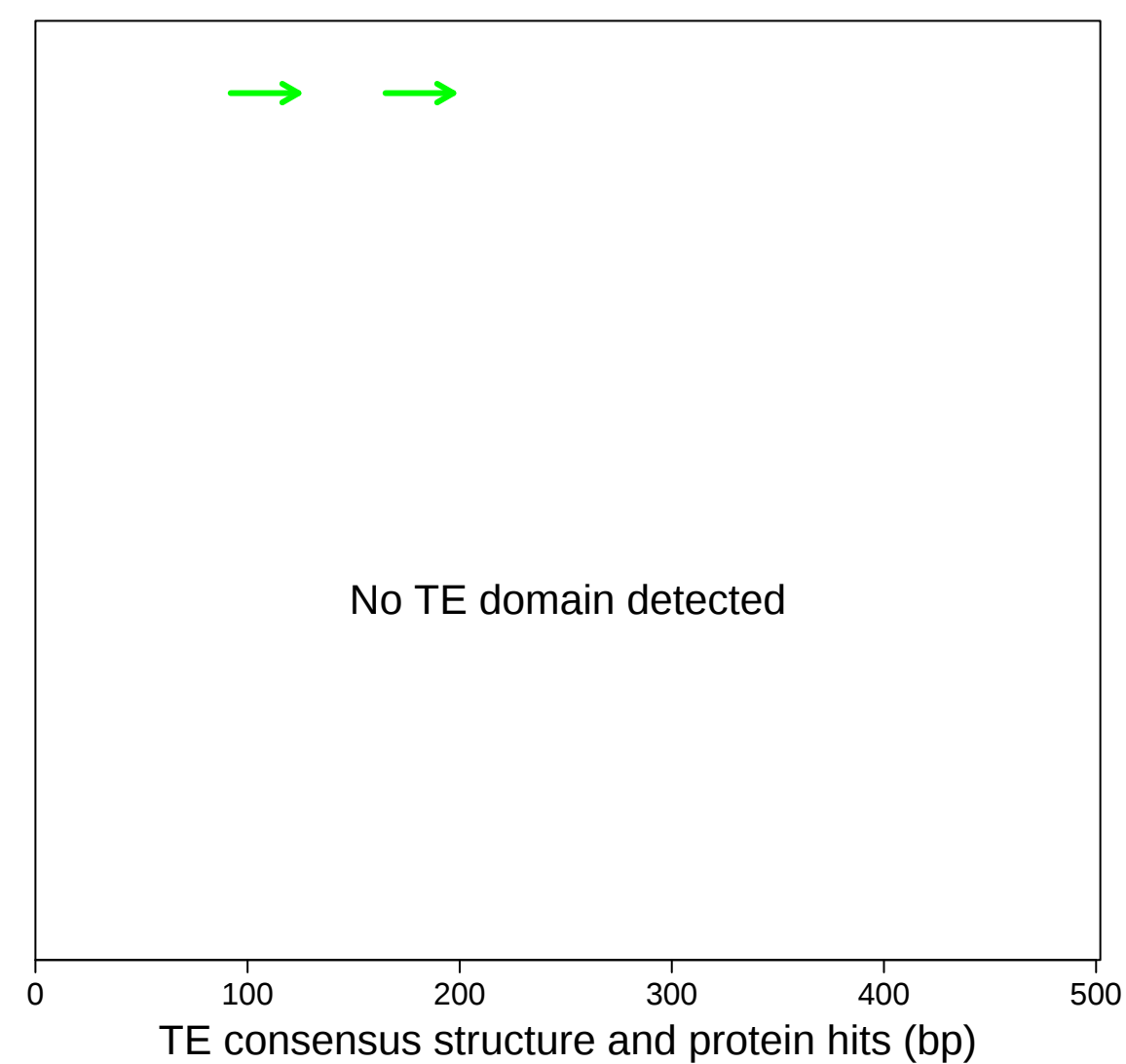
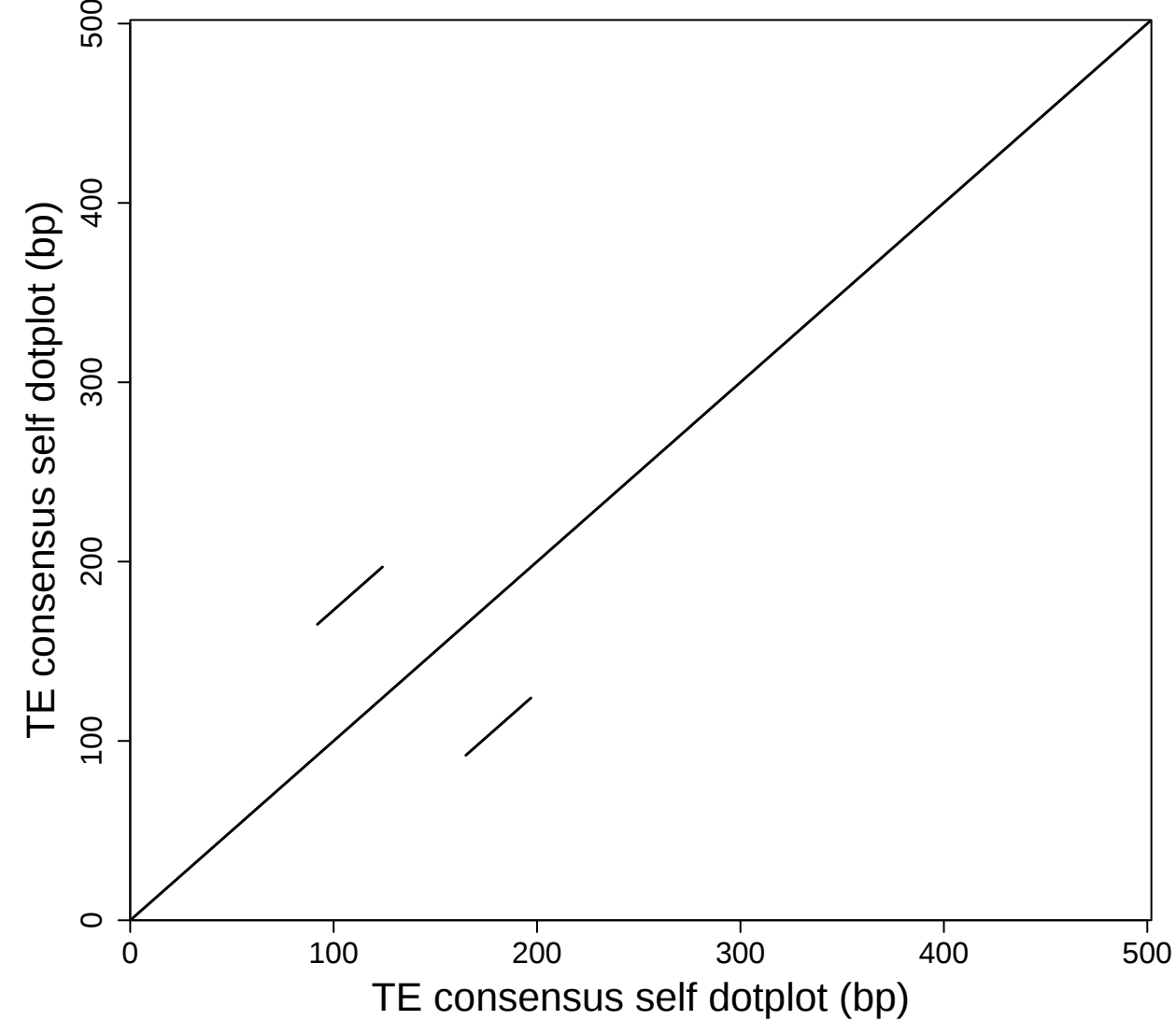
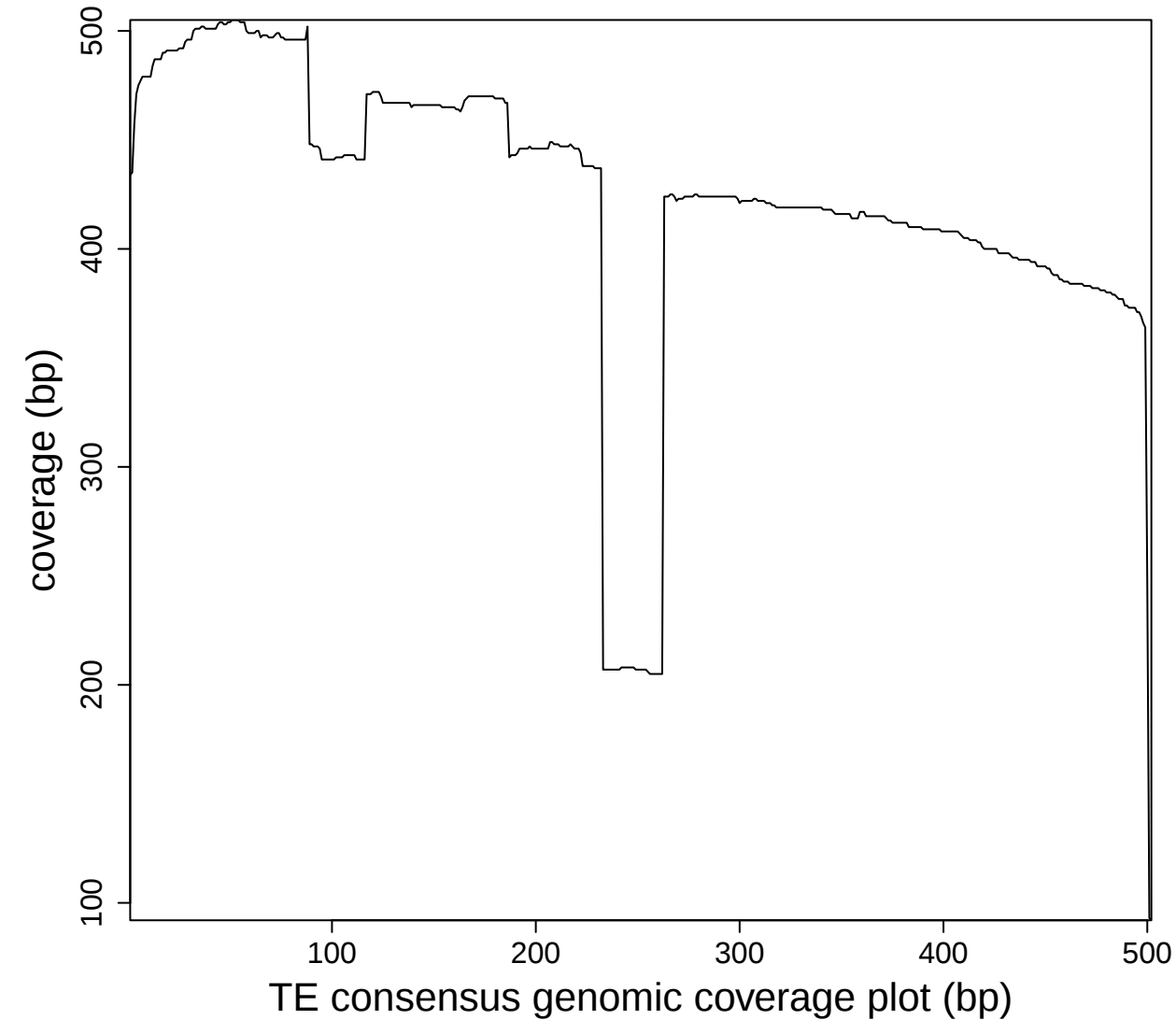
After TEtrimmer Extended plot Blue lines are boundaries



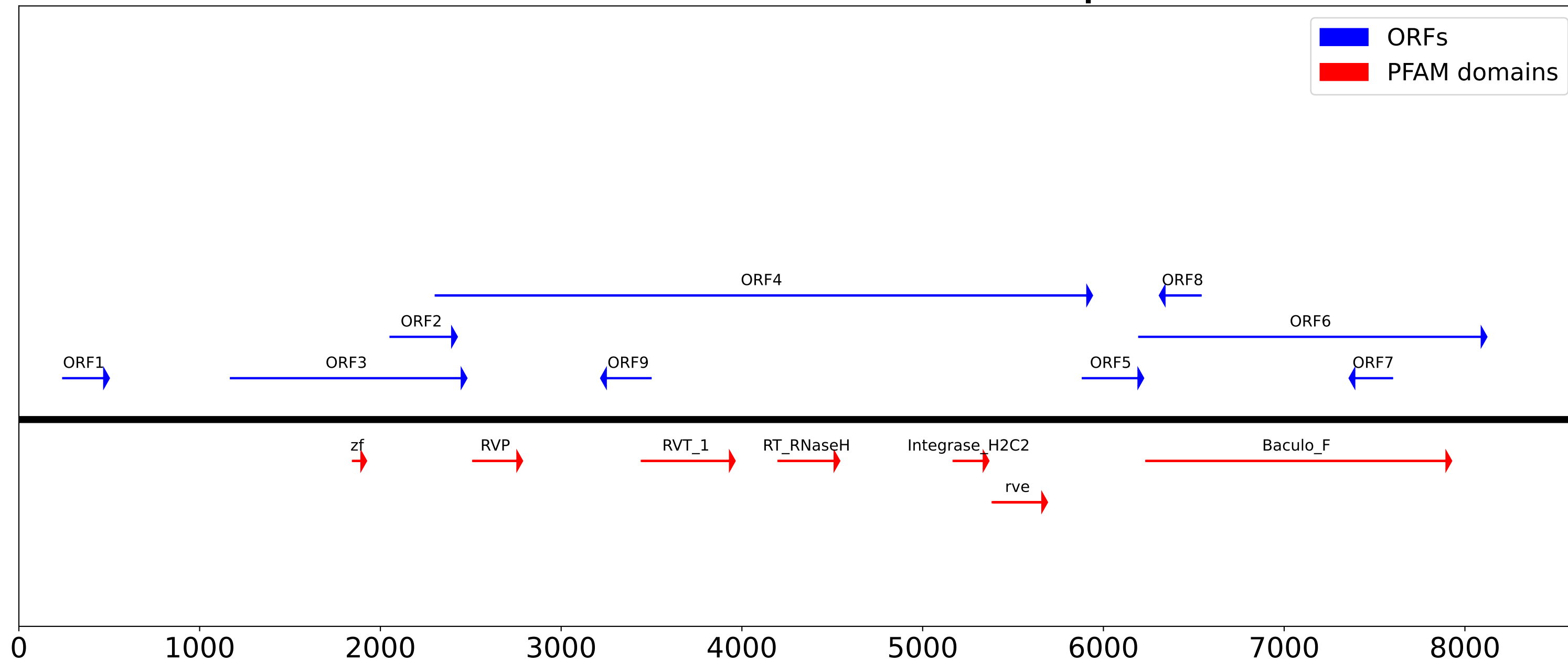
TE: ltr_1_family_39
size: 502bp; fragments: 802; full length: 125 (≥ 451.8 bp)



Before TEtrimmer 502 bp



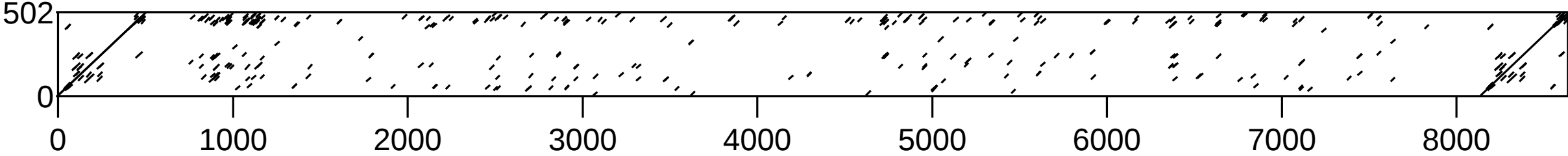
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)