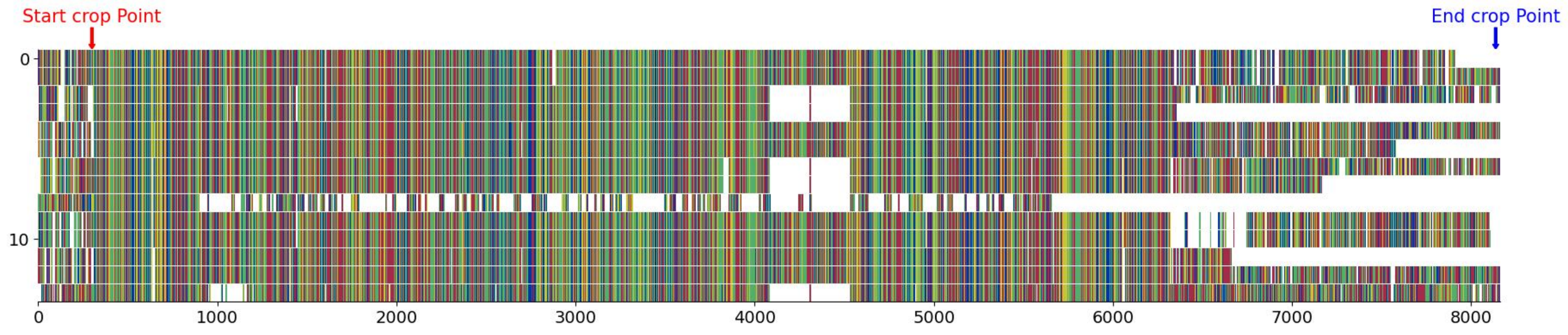
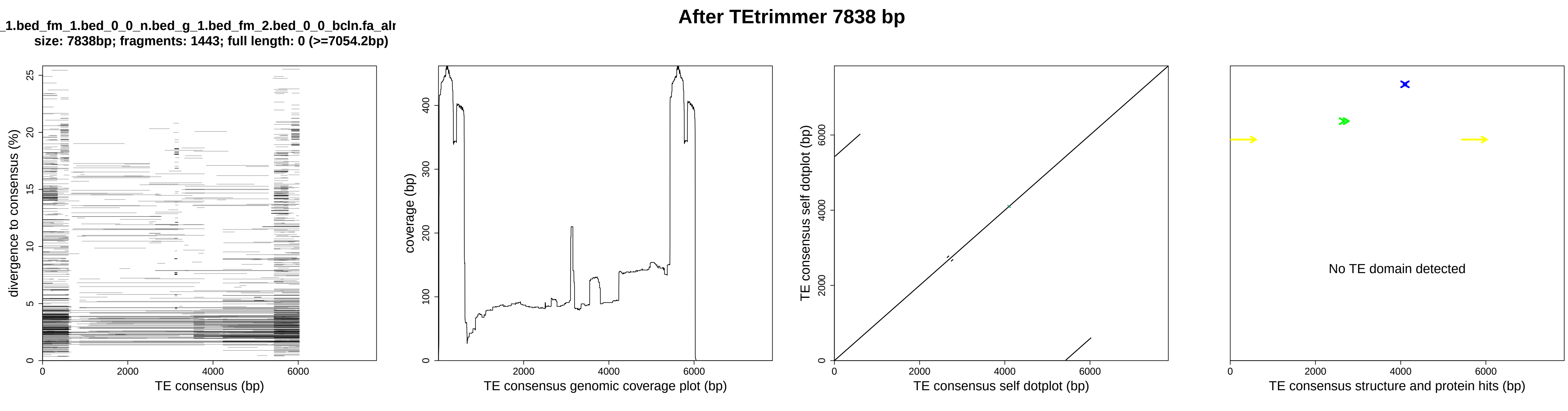




MSA length = 7838

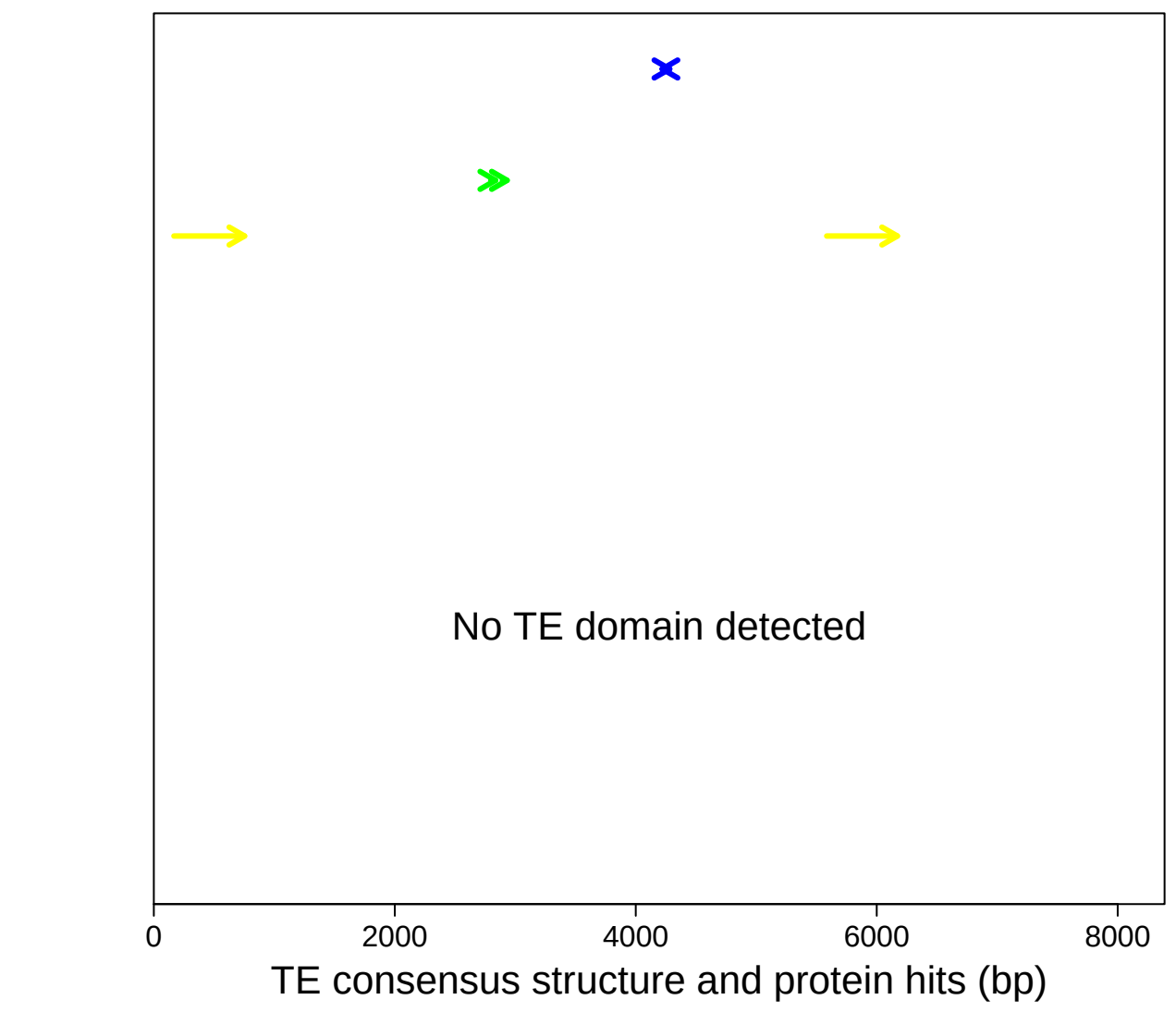
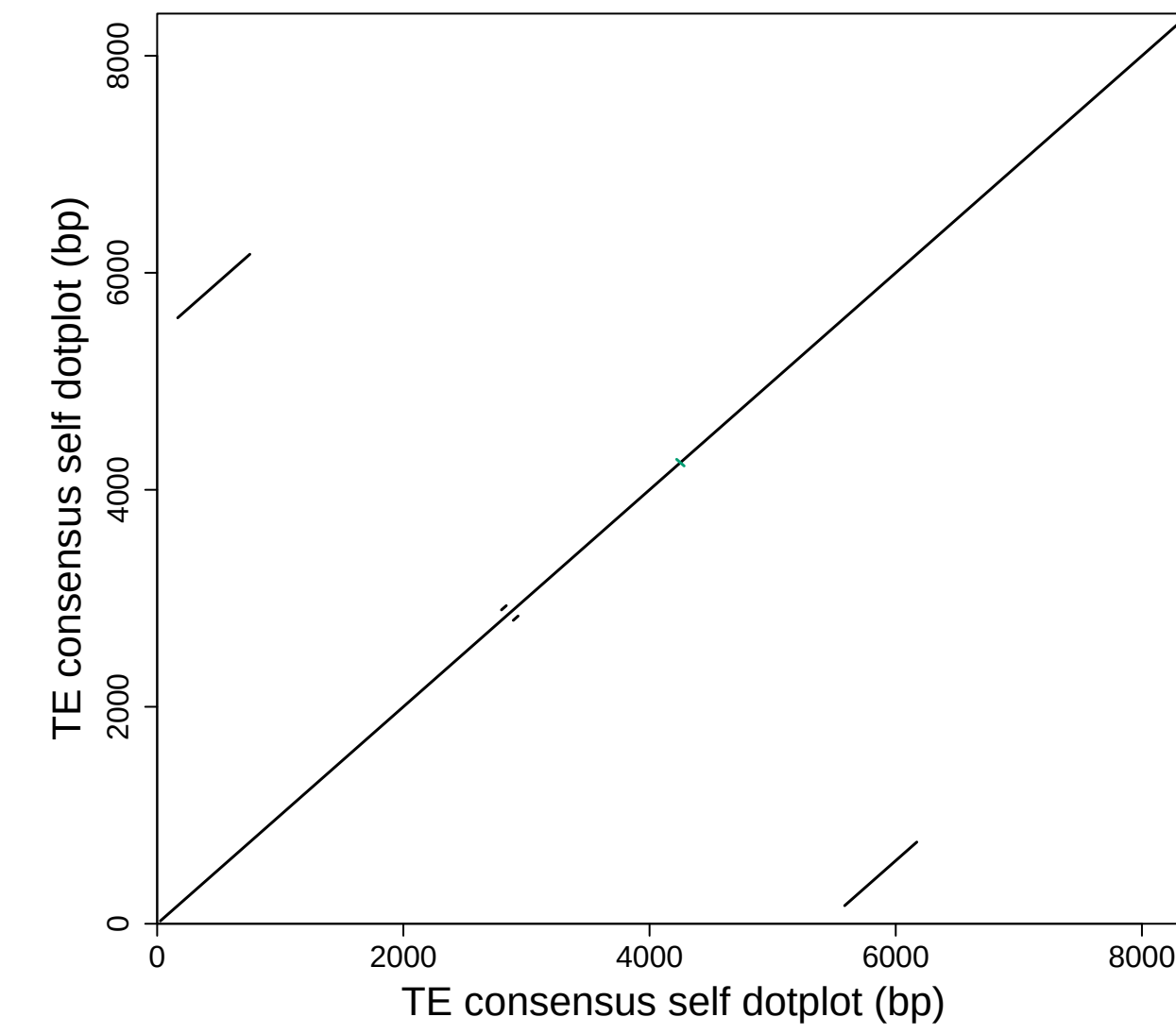
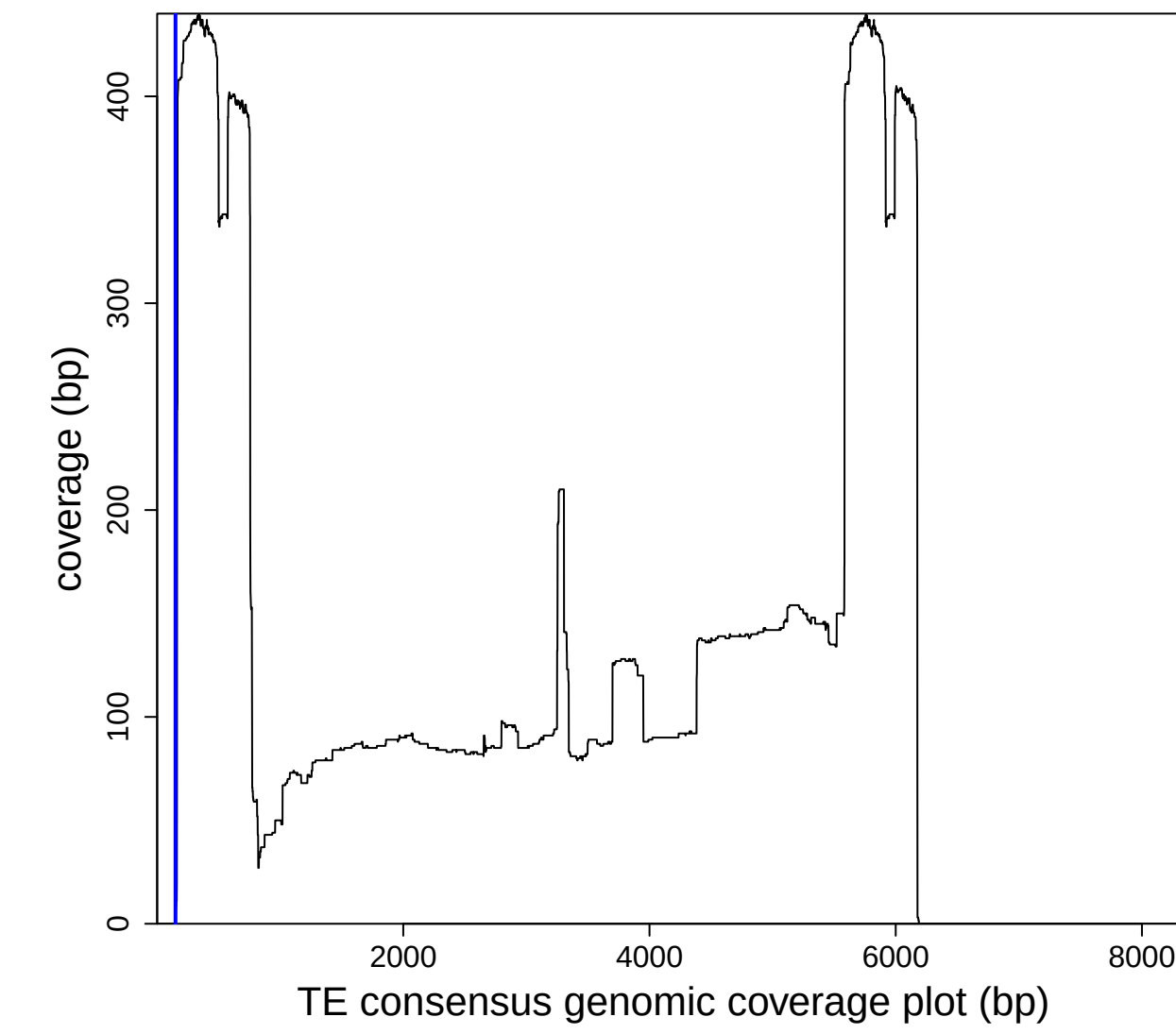
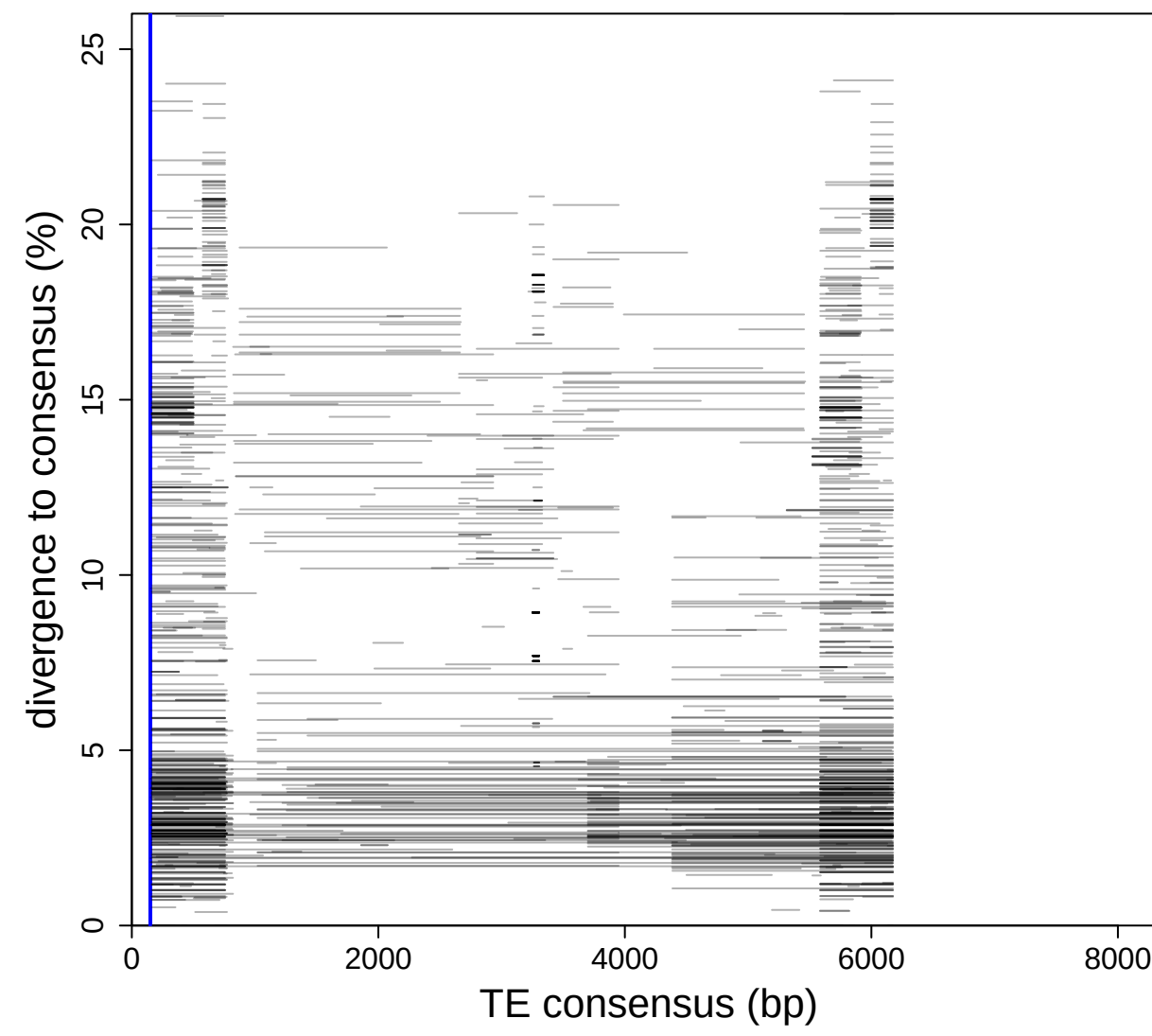
[illegible]



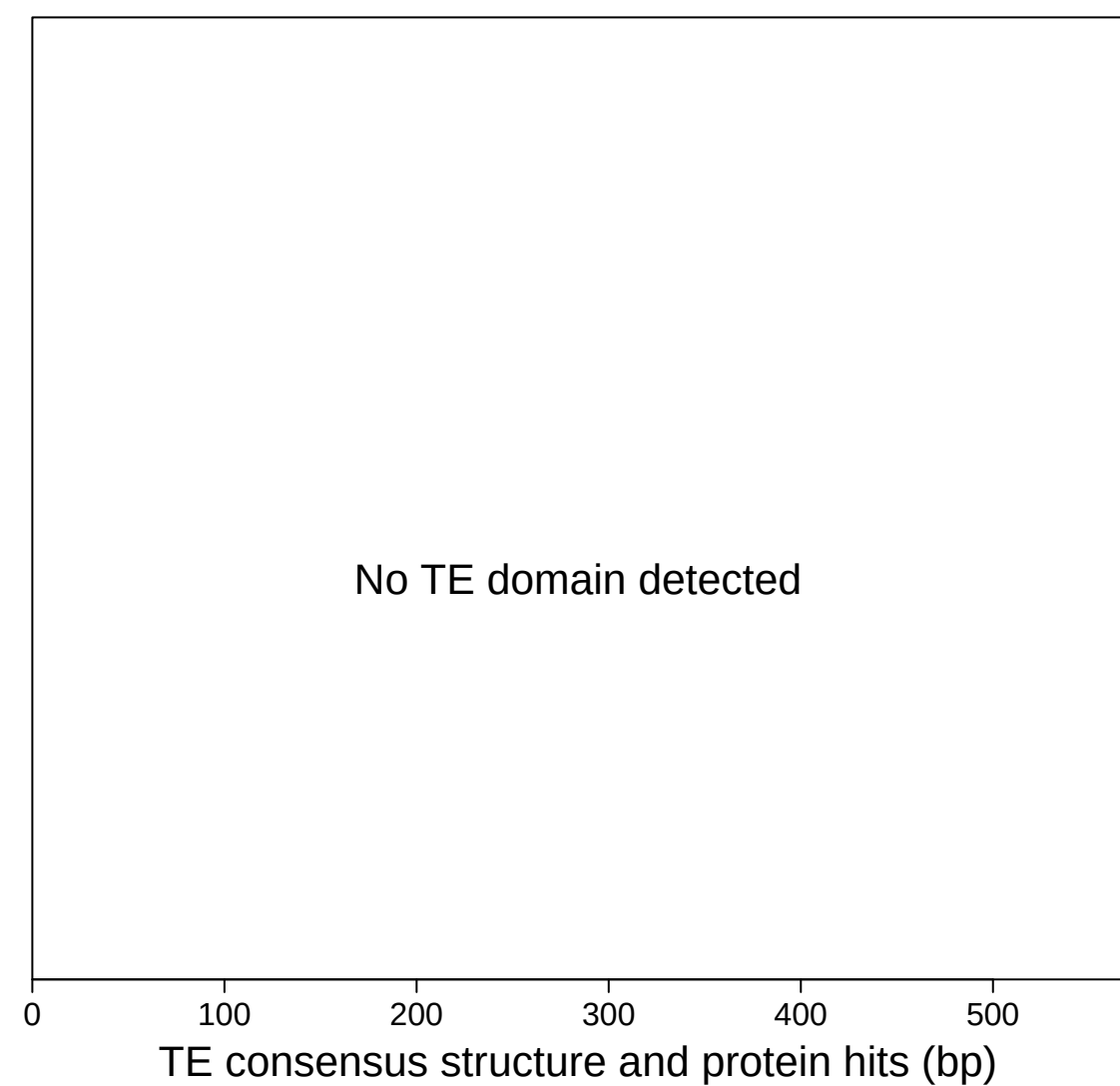
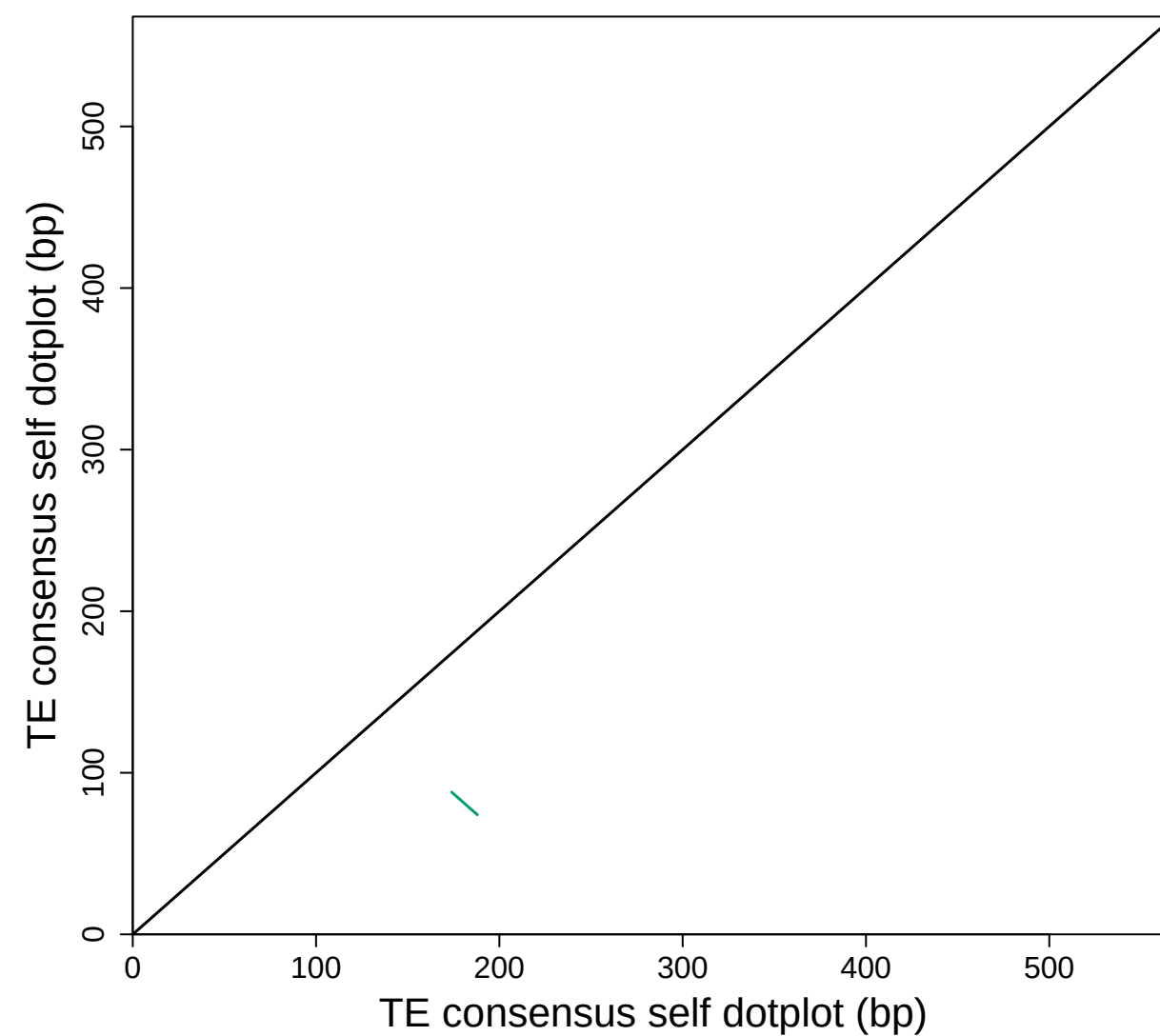
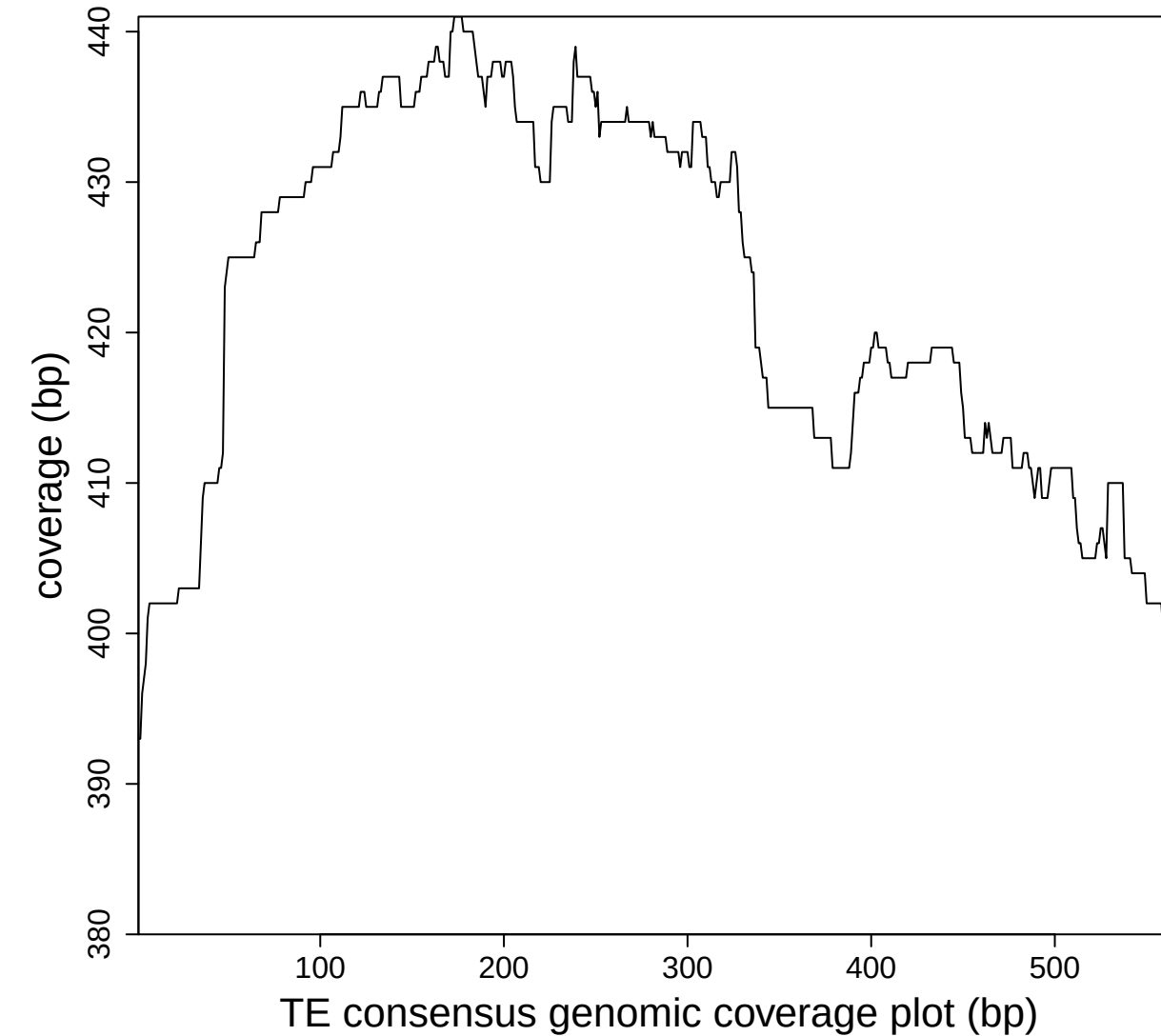
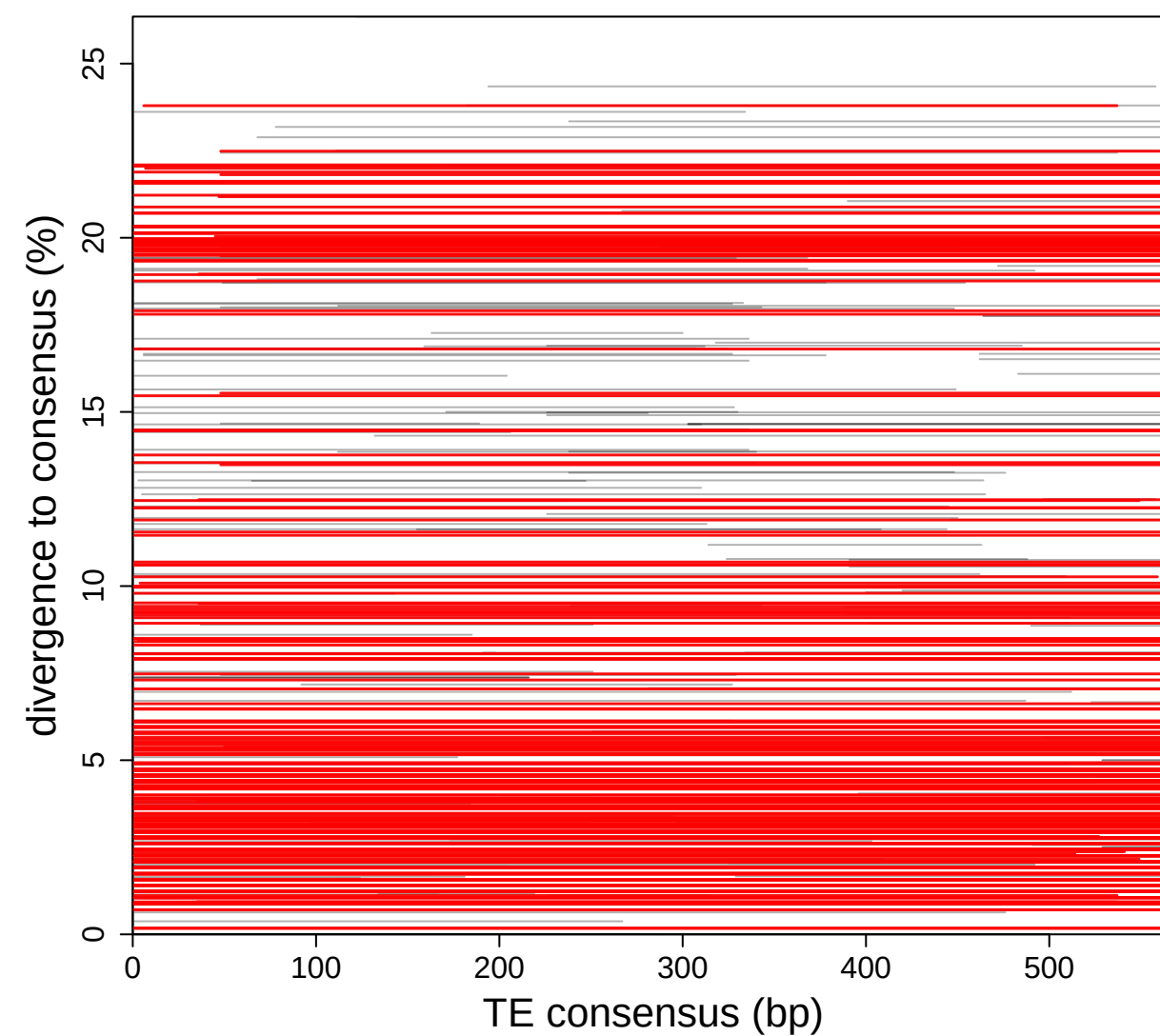


bed_g_1.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln
size: 8389bp; fragments: 1395; full length: 0 (>=7550.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

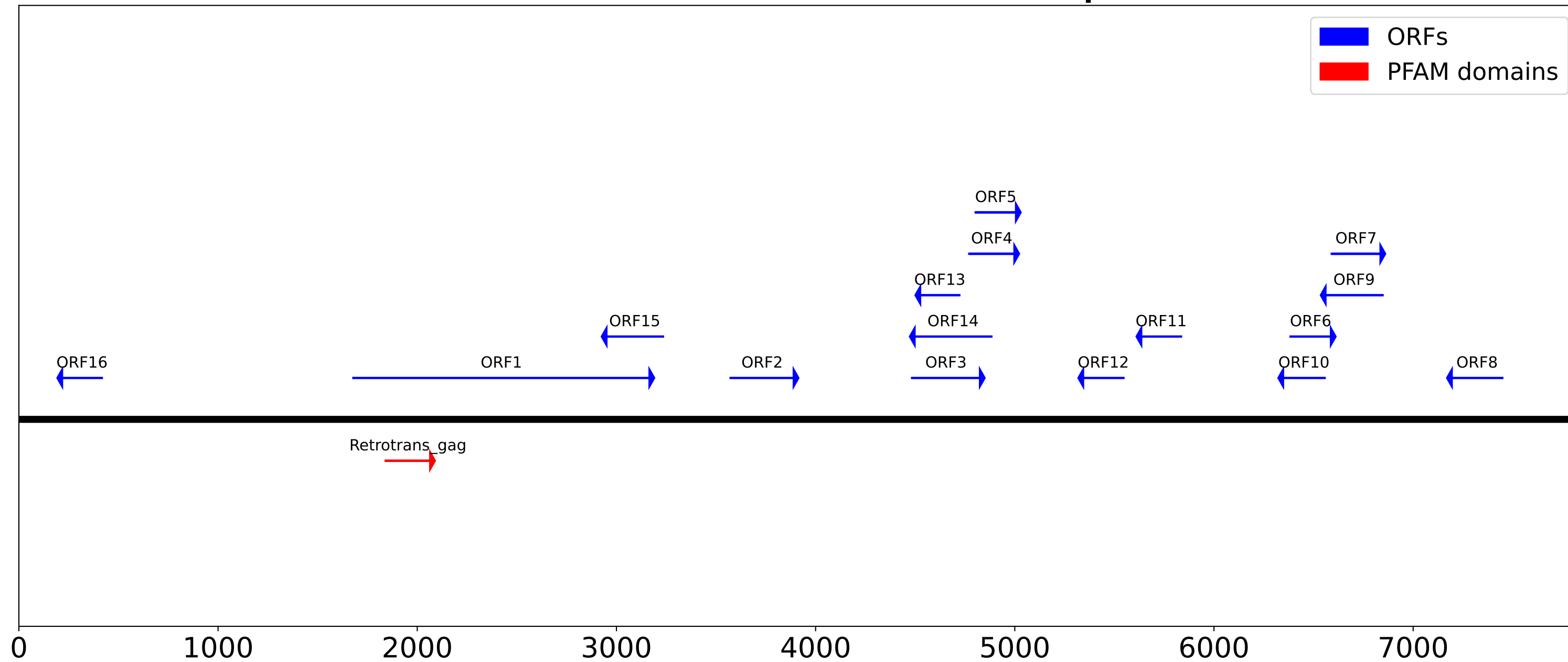


TE: ltr_1_family_81
size: 568bp; fragments: 506; full length: 351 (≥ 511.2 bp)

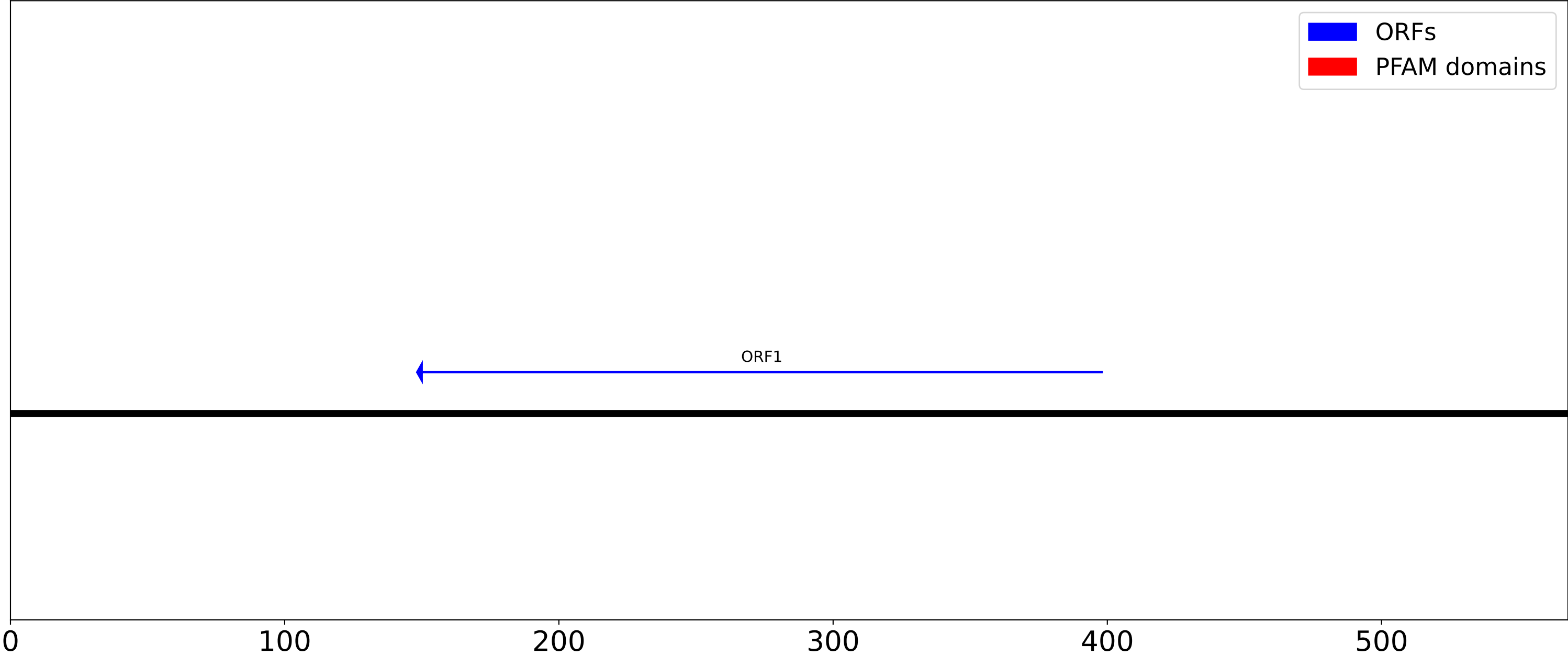


Before TEtrimmer 568 bp

After TEtrimmer ORF and PFAM domain plot



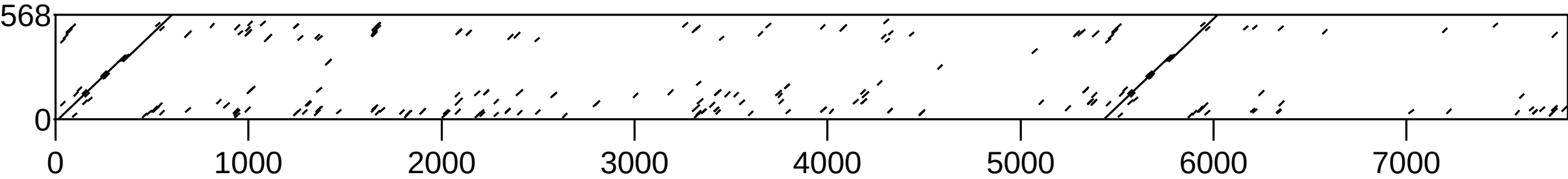
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)