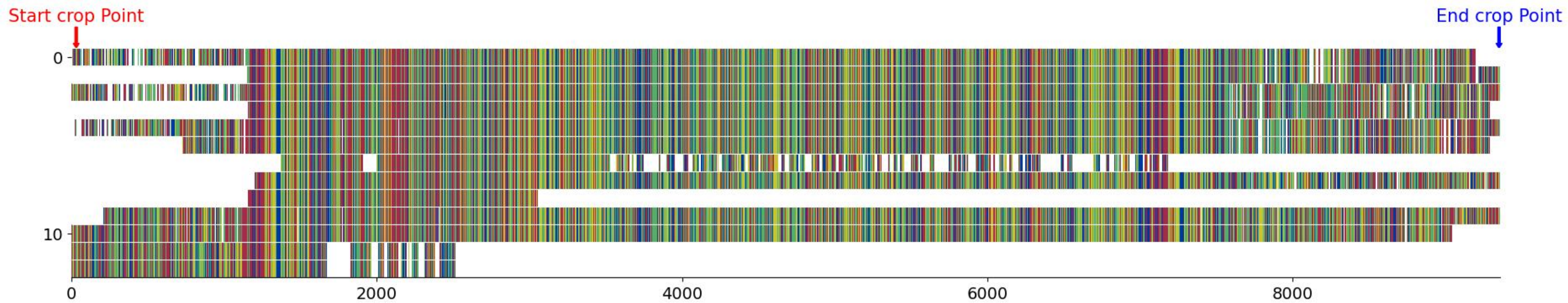
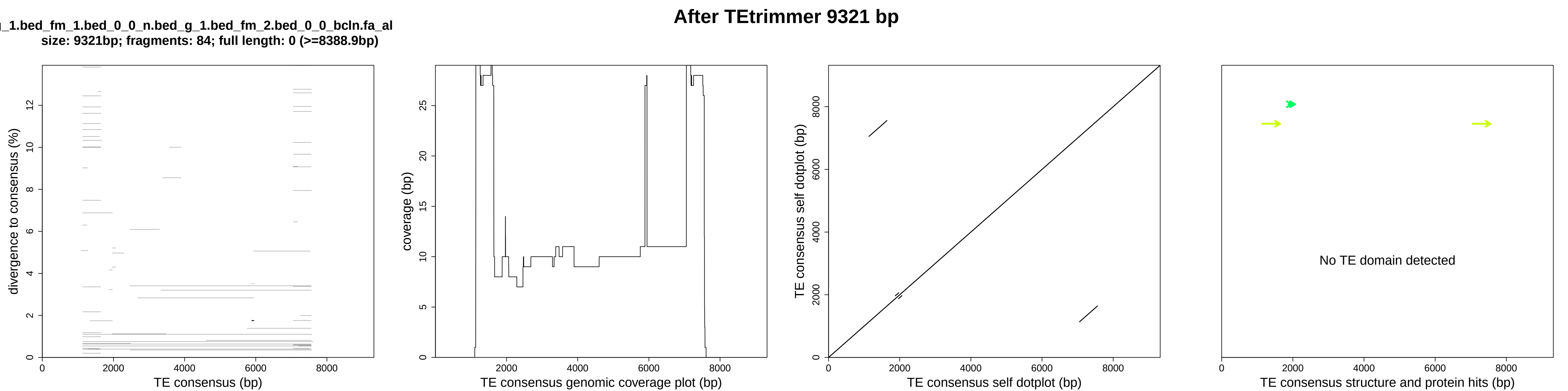


MSA length = 9321

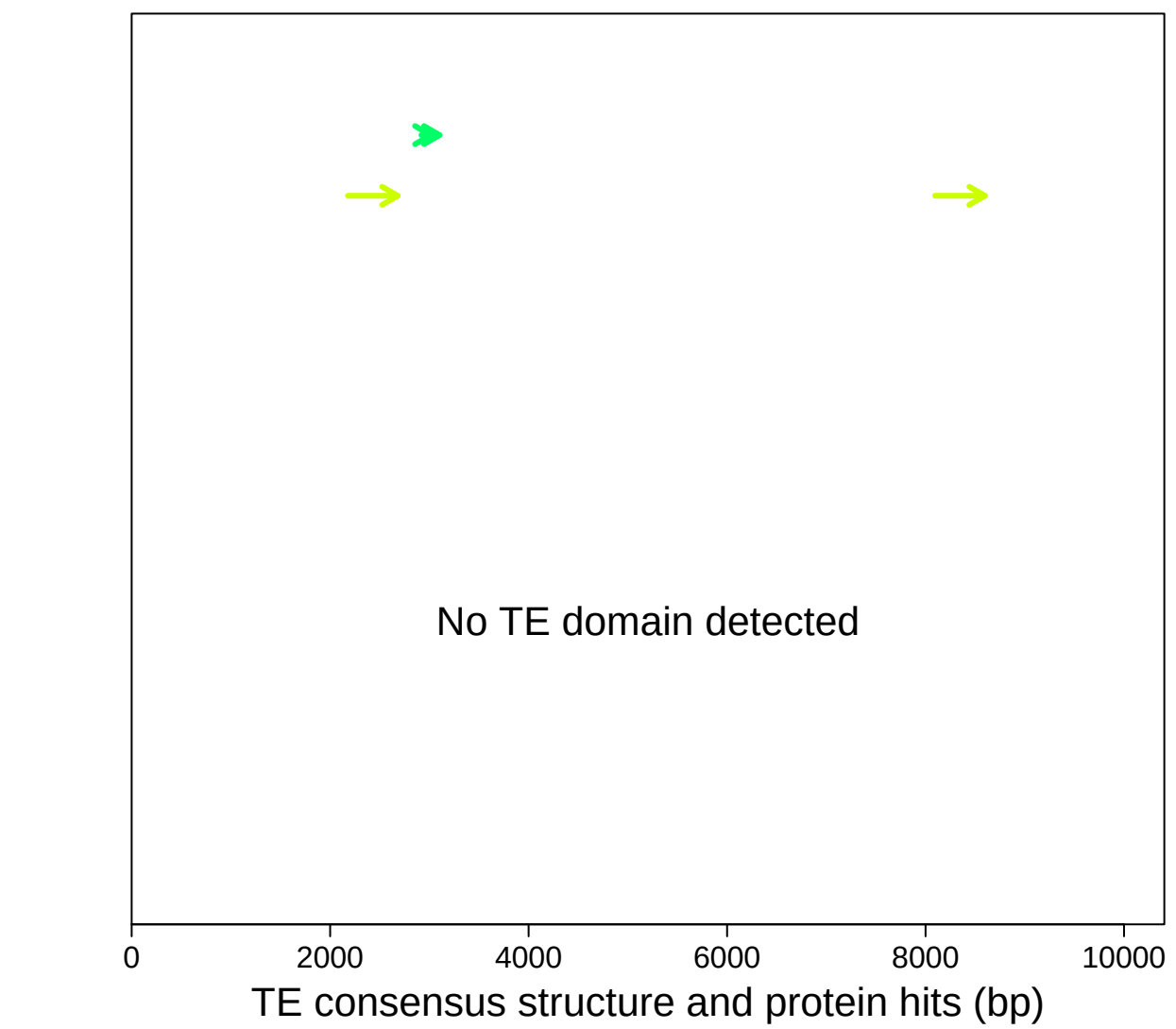
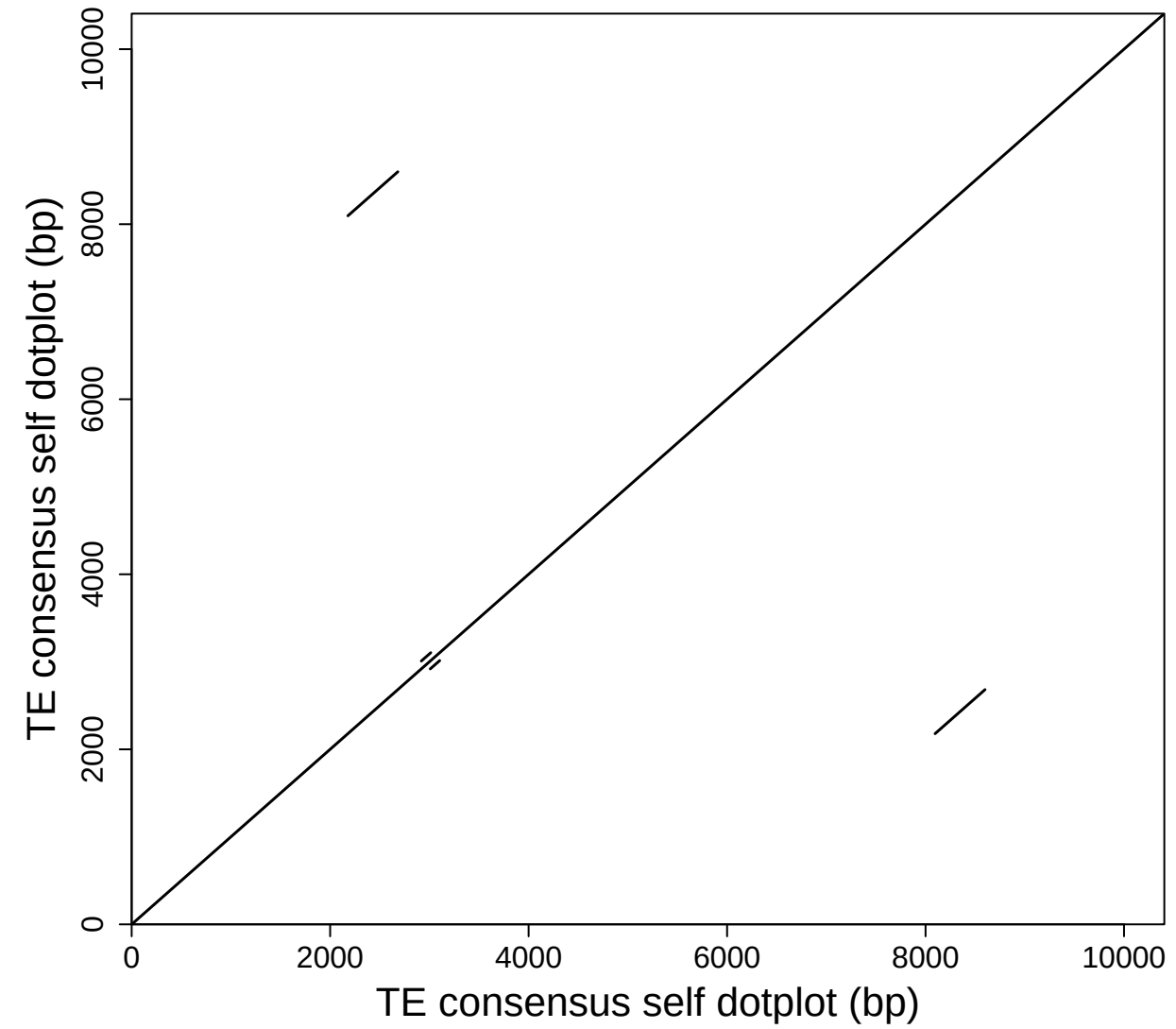
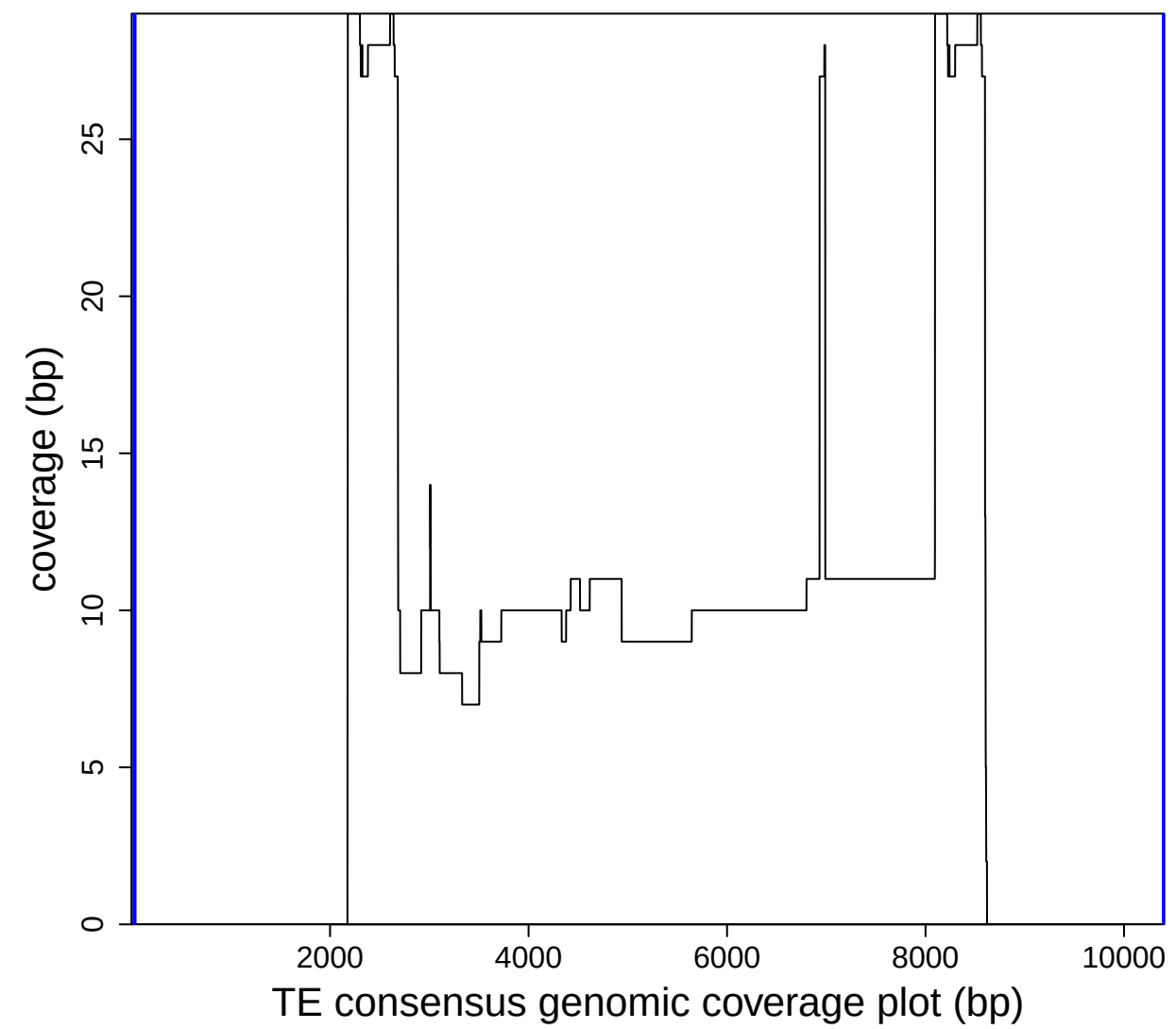
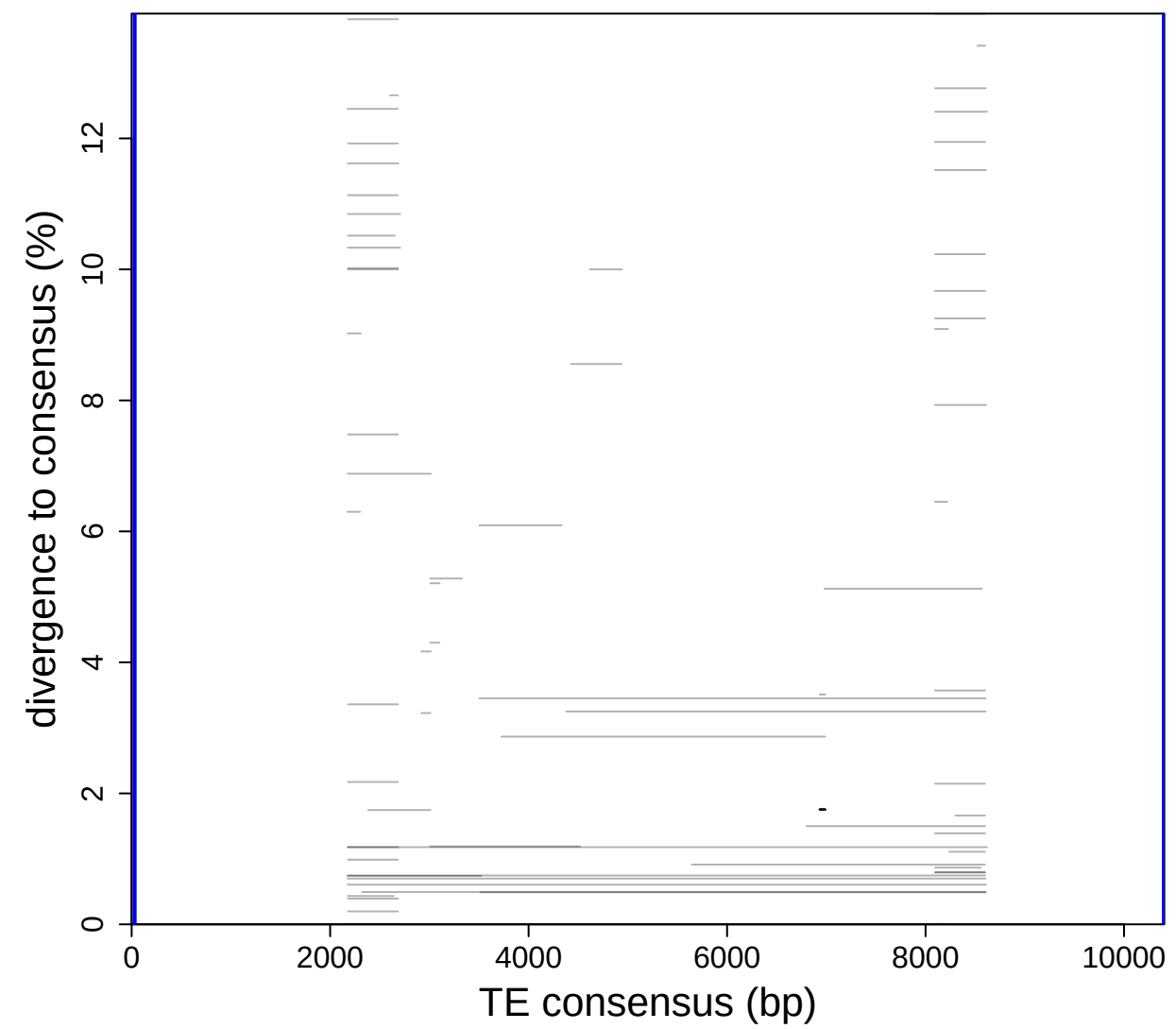
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-----TCTACTTTA-----ACTTTACCTAAACTC-----GTATCTTCTCAAATATTAGTTTAAATAAATTTAAGAGCTAATTAGTAAGCAAGAAT
-----GCATTTTCTTAAGTAACAGTCCAAATAATGGGAAGATCTCATTGGTAGACGAGAGT
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bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 10406bp; fragments: 85; full length: 0 (>=9365.4bp)

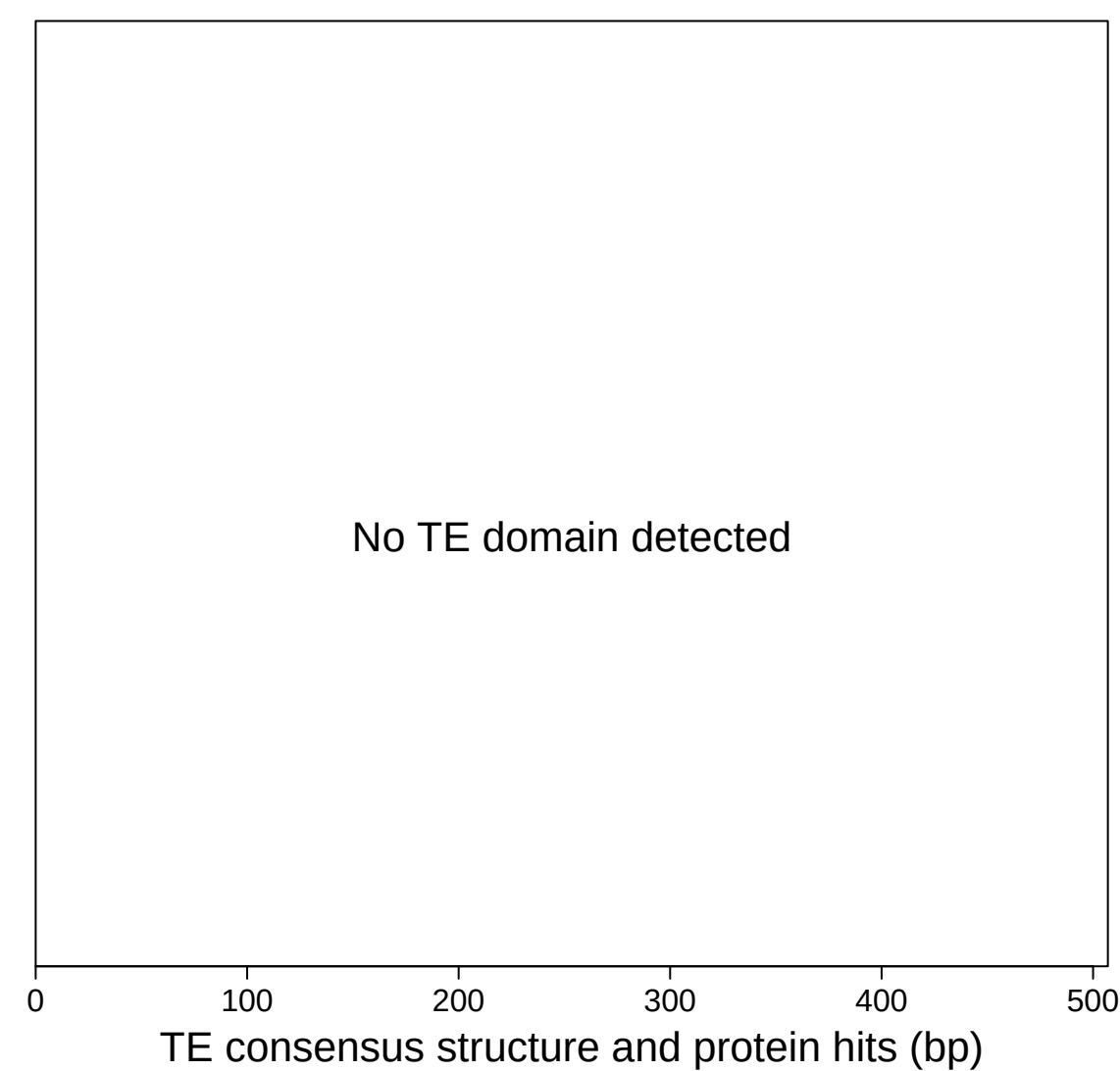
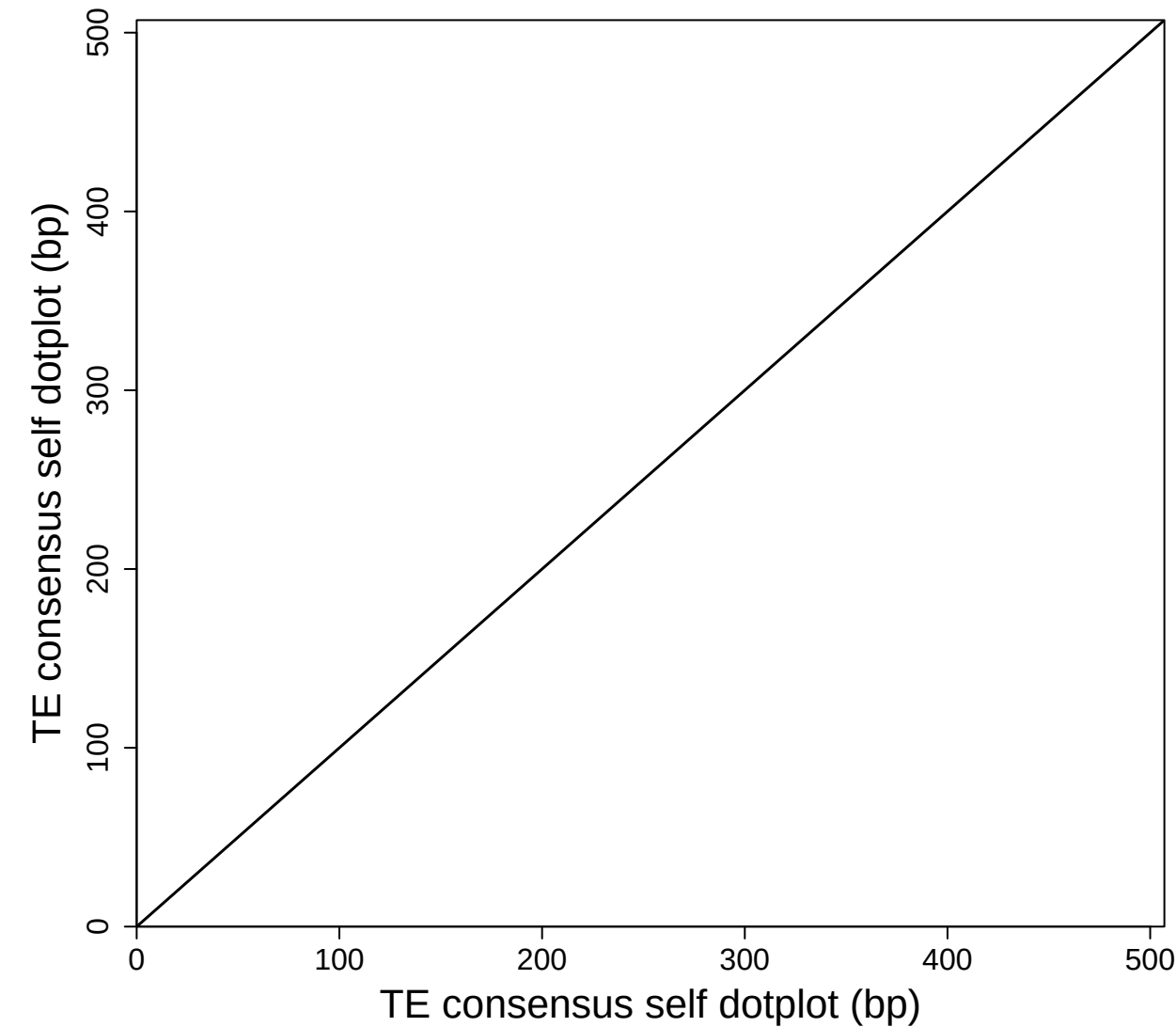
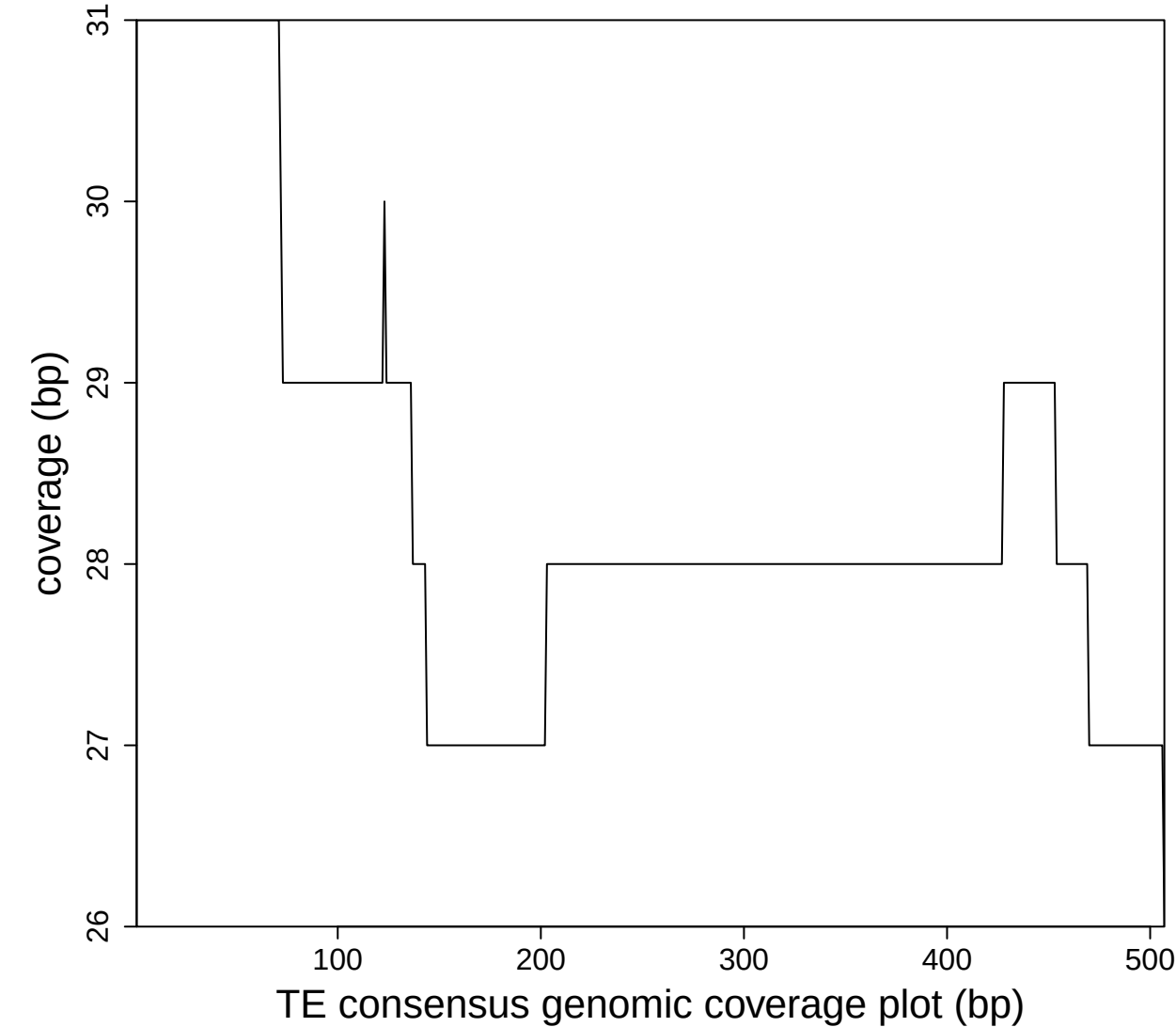
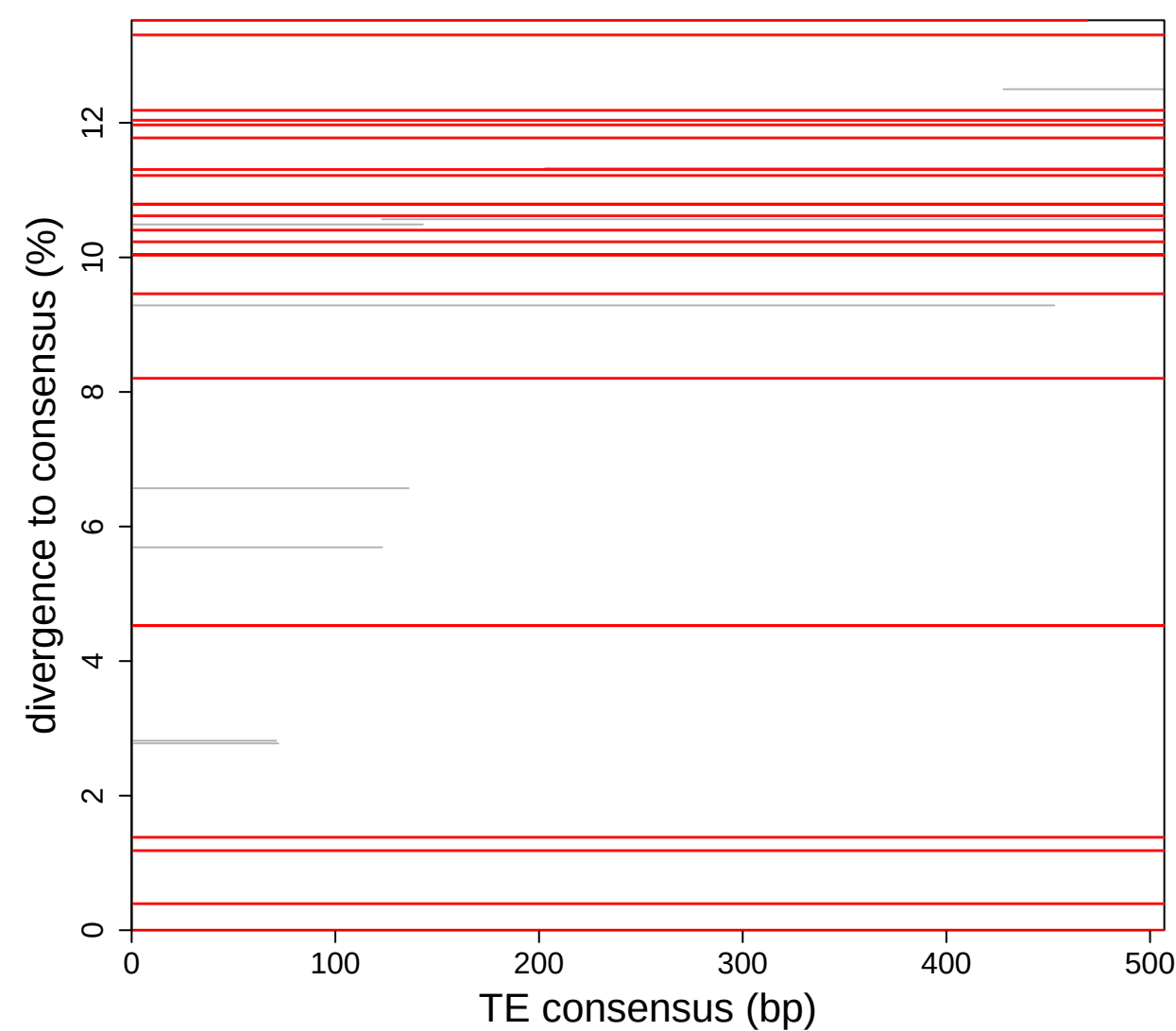
After TEtrimmer Extended plot Blue lines are boundaries



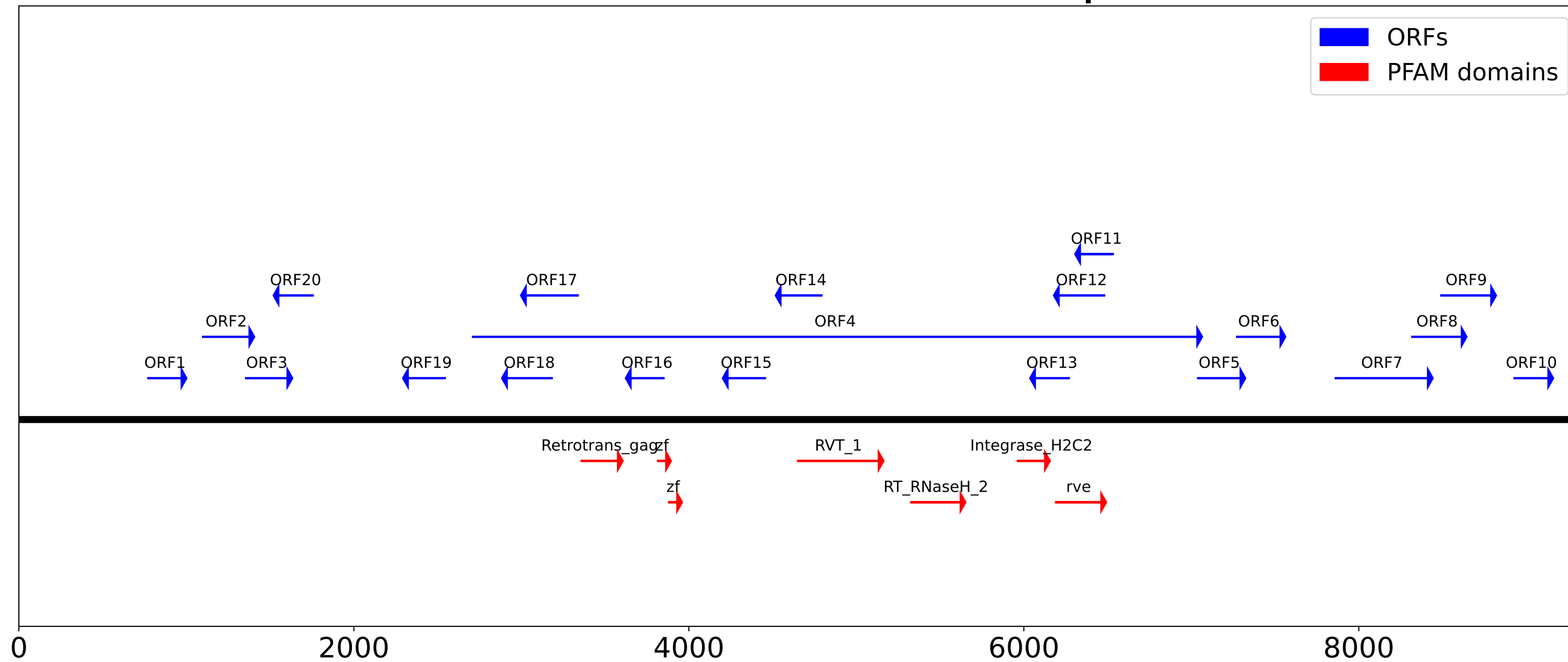
Before TEtrimmer 507 bp

TE: ltr_1_family_377

size: 507bp; fragments: 34; full length: 25 (≥ 456.3 bp)



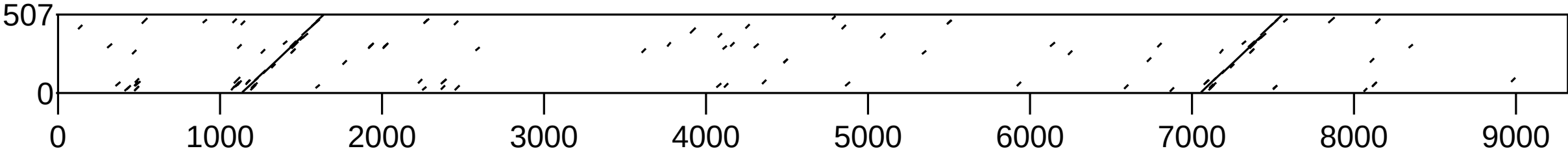
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

Consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)