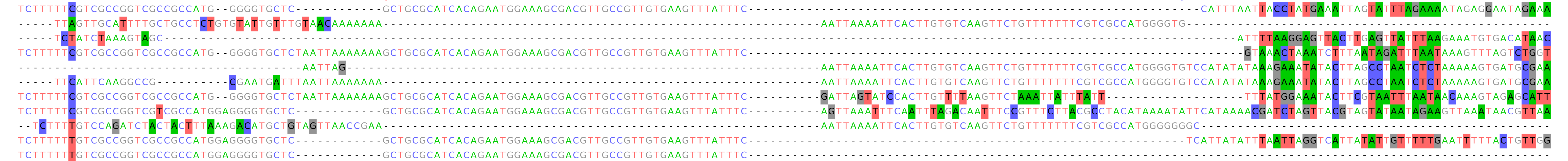


Start crop Point



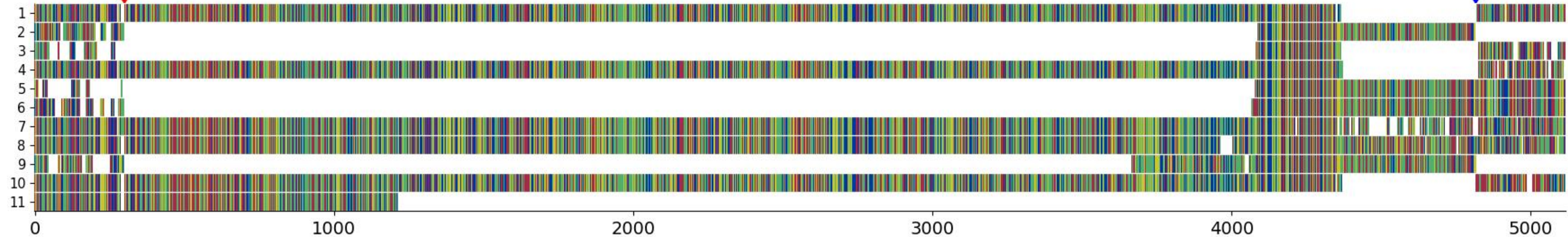
MSA length = 4517

End crop Point



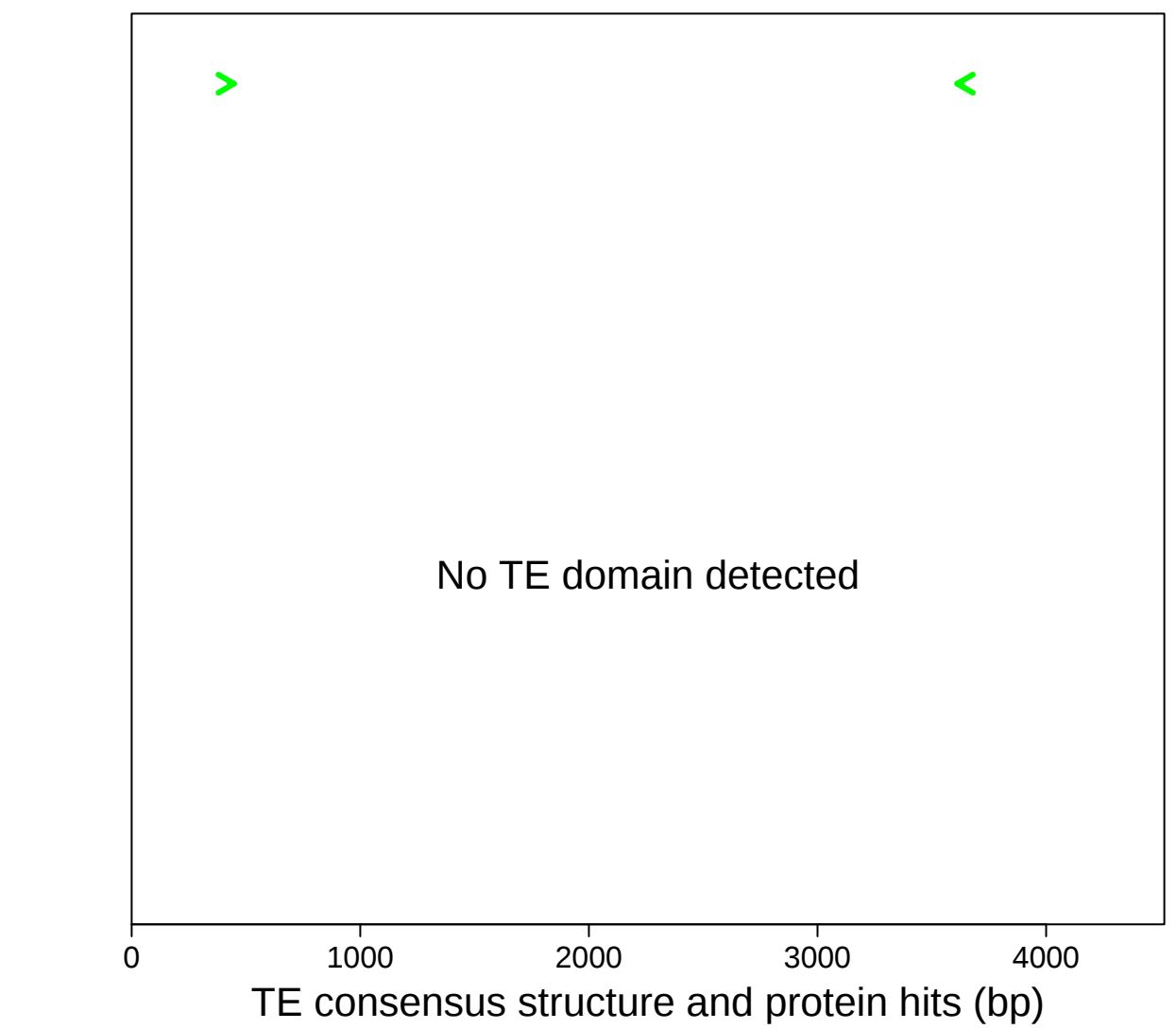
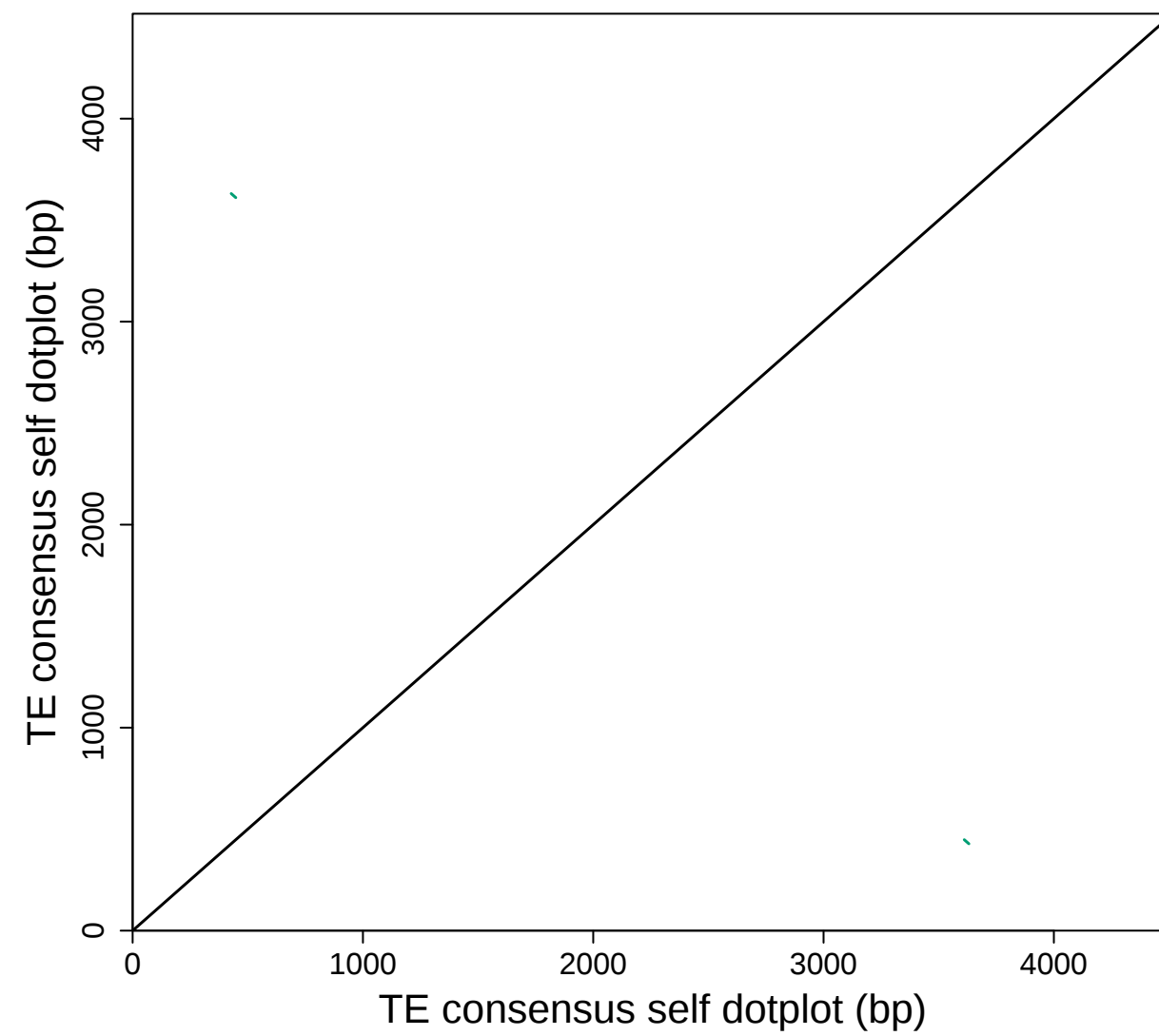
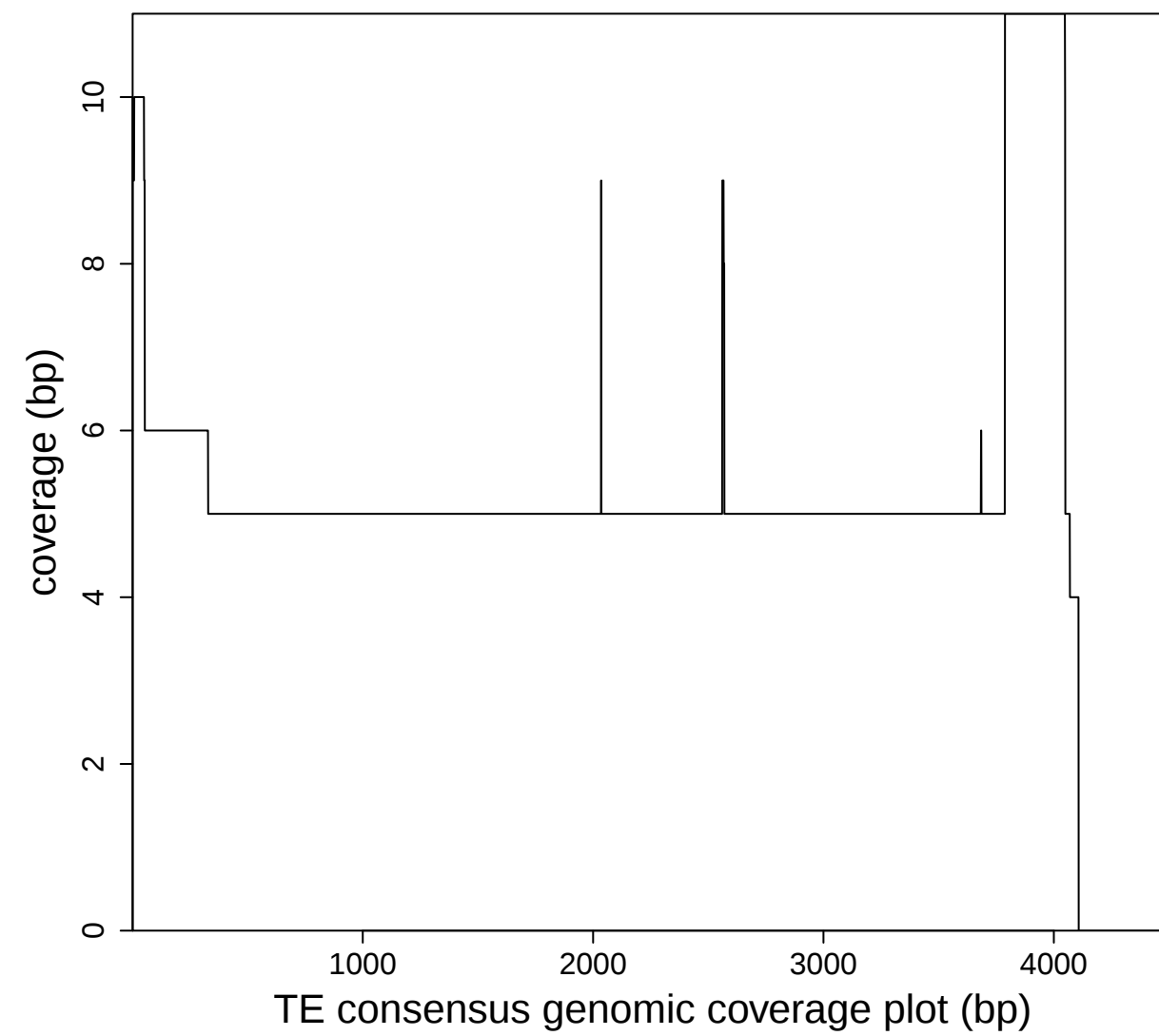
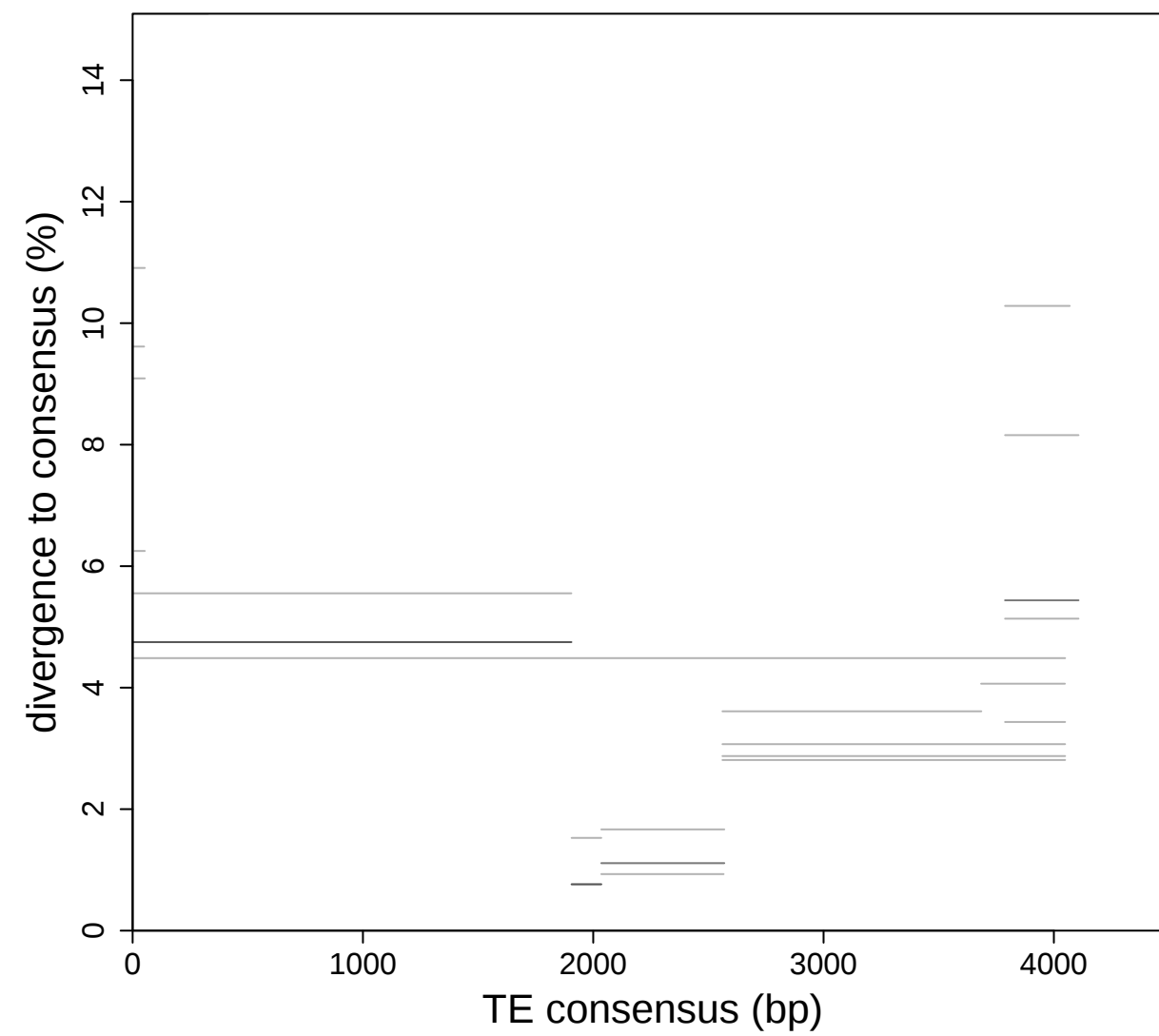
Start crop Point

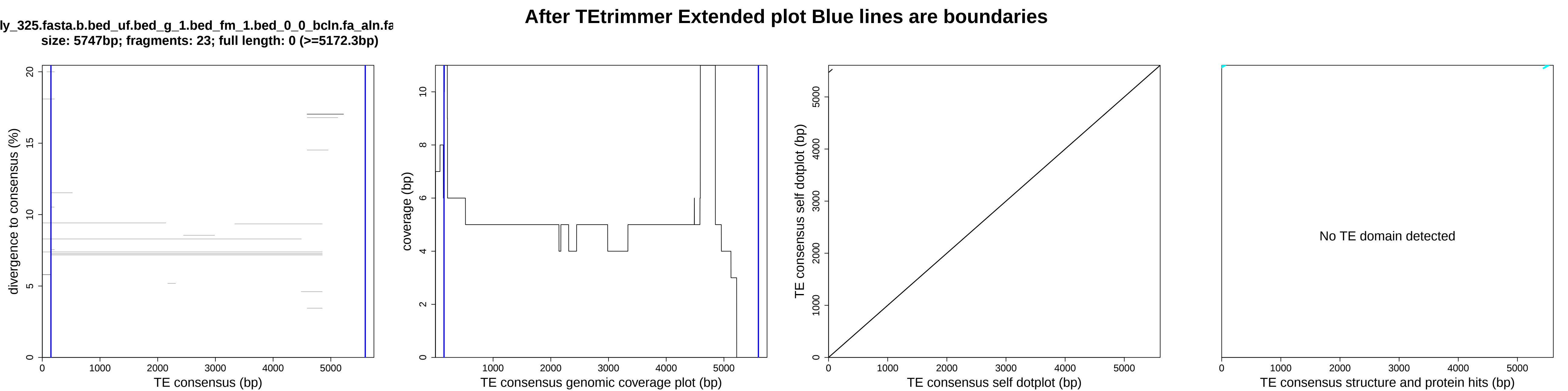
End crop Point



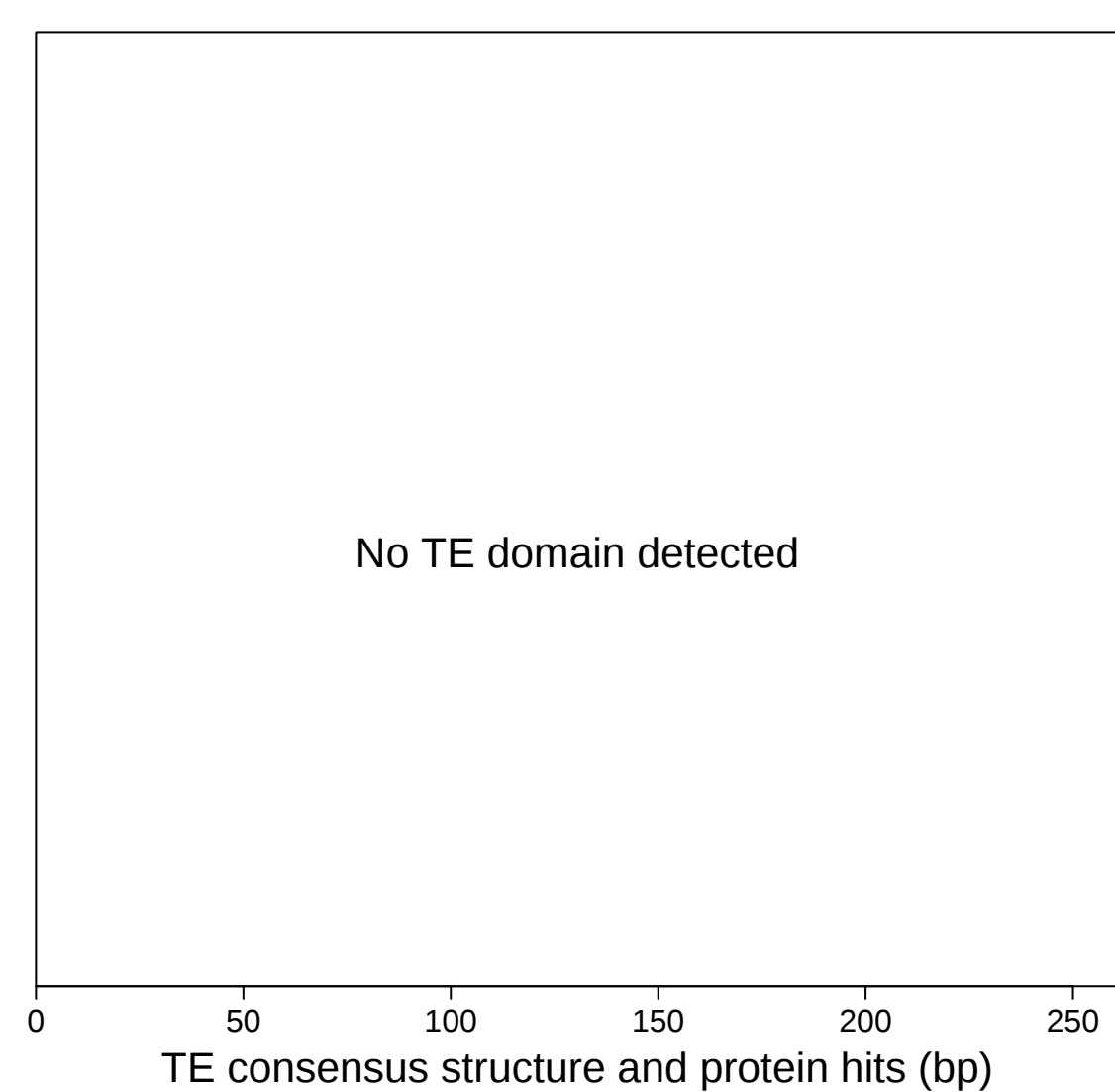
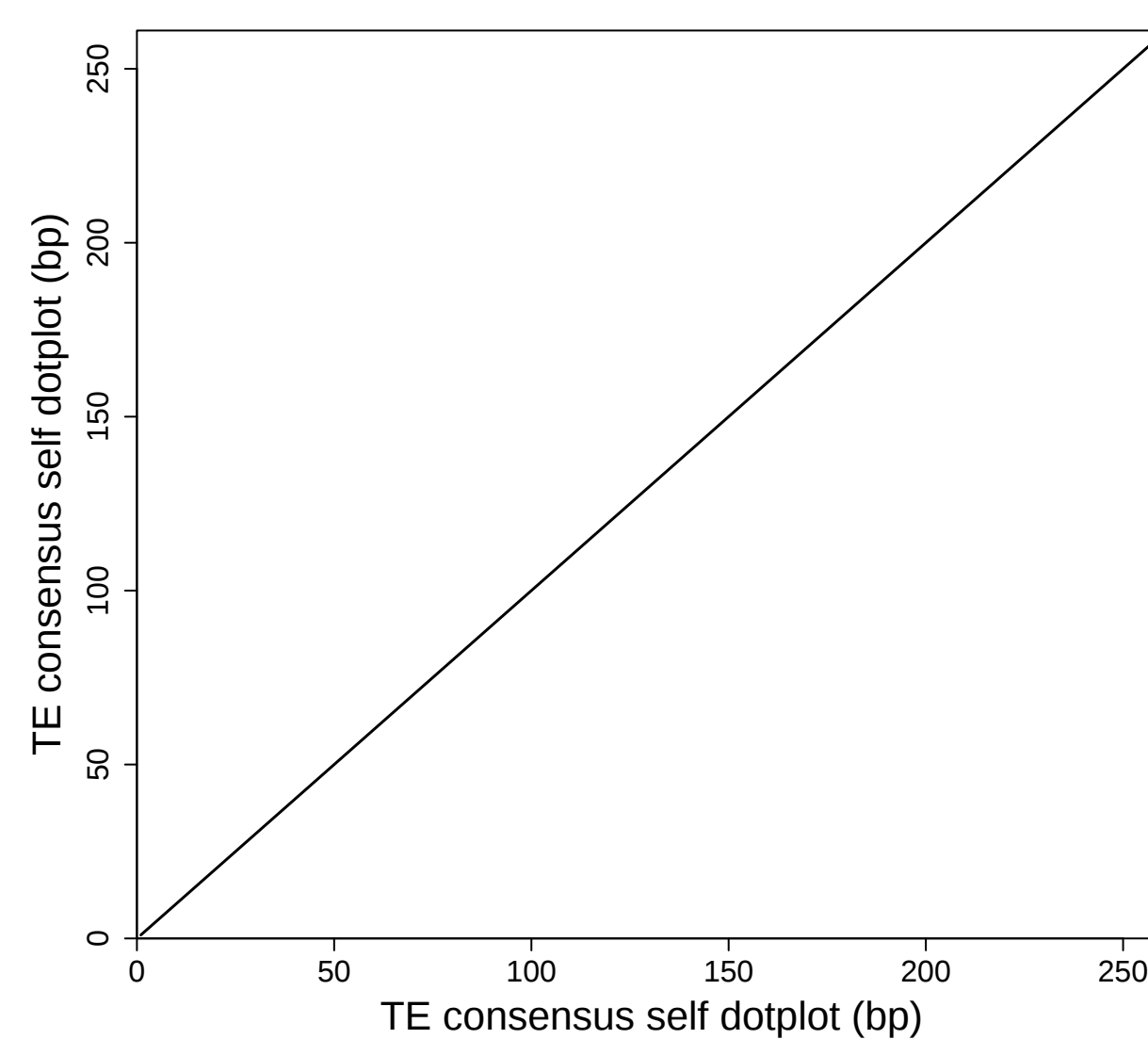
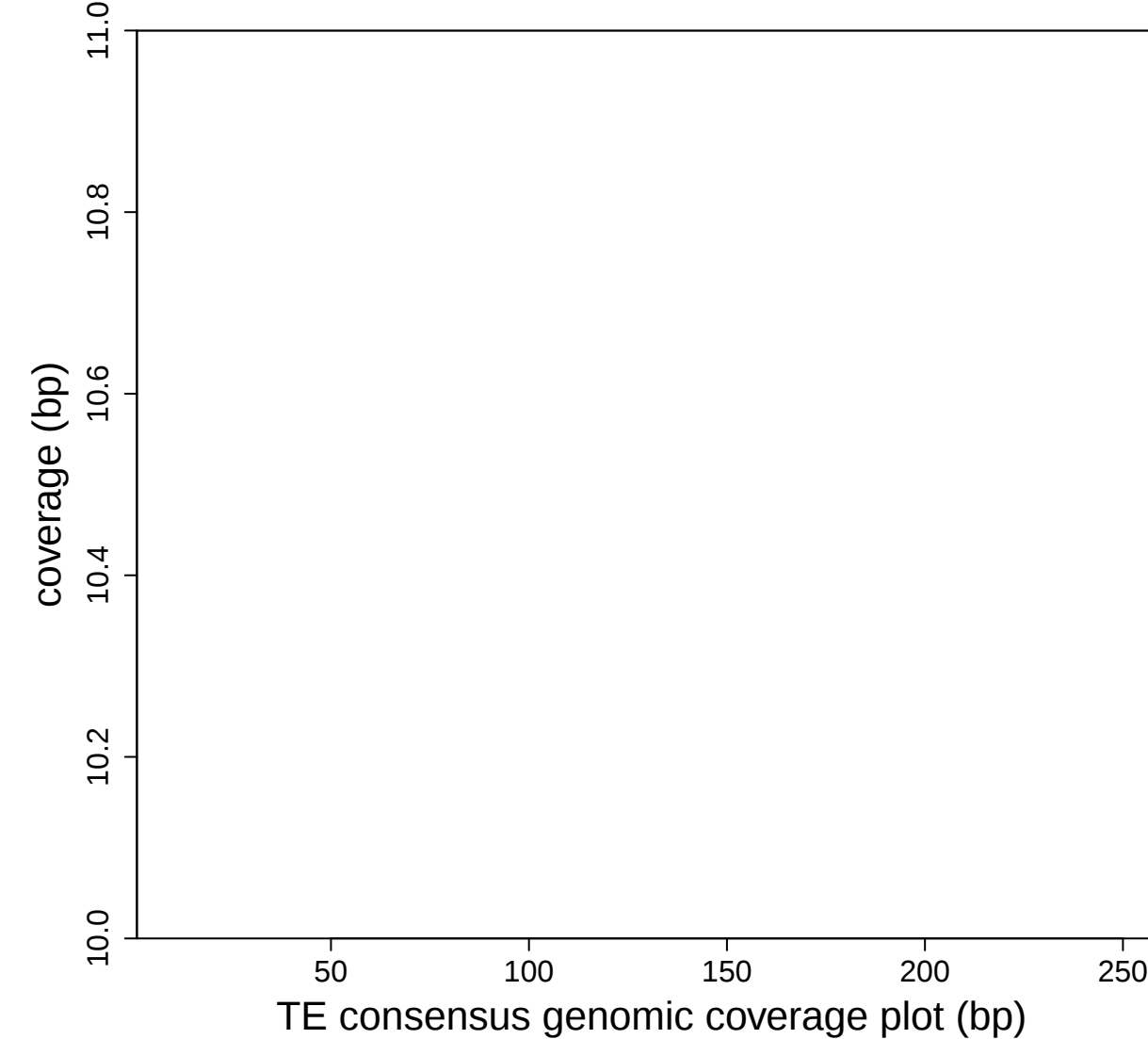
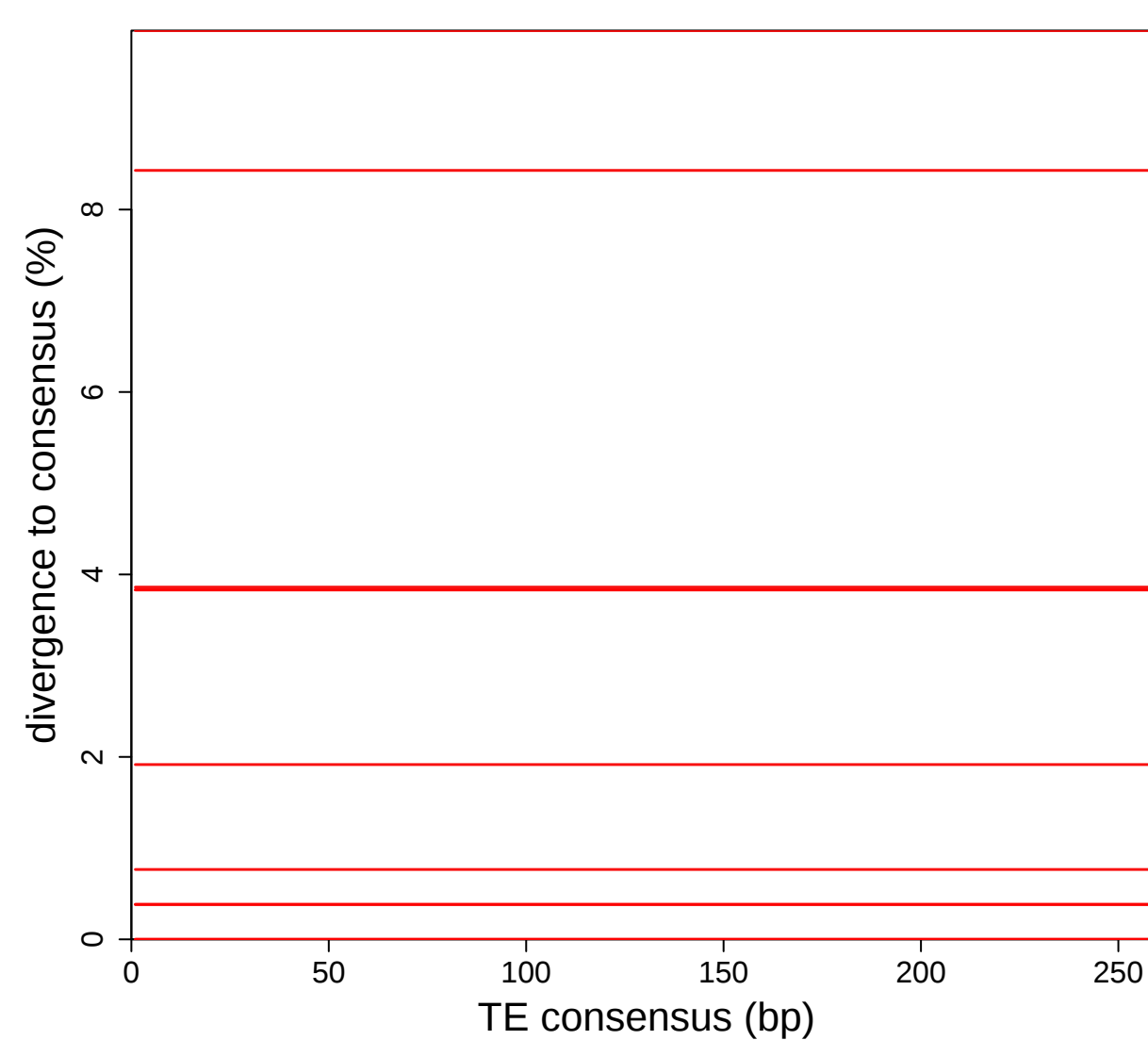
ed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 4517bp; fragments: 29; full length: 0 (>=4065.3bp)

After TETrimmer 4517 bp



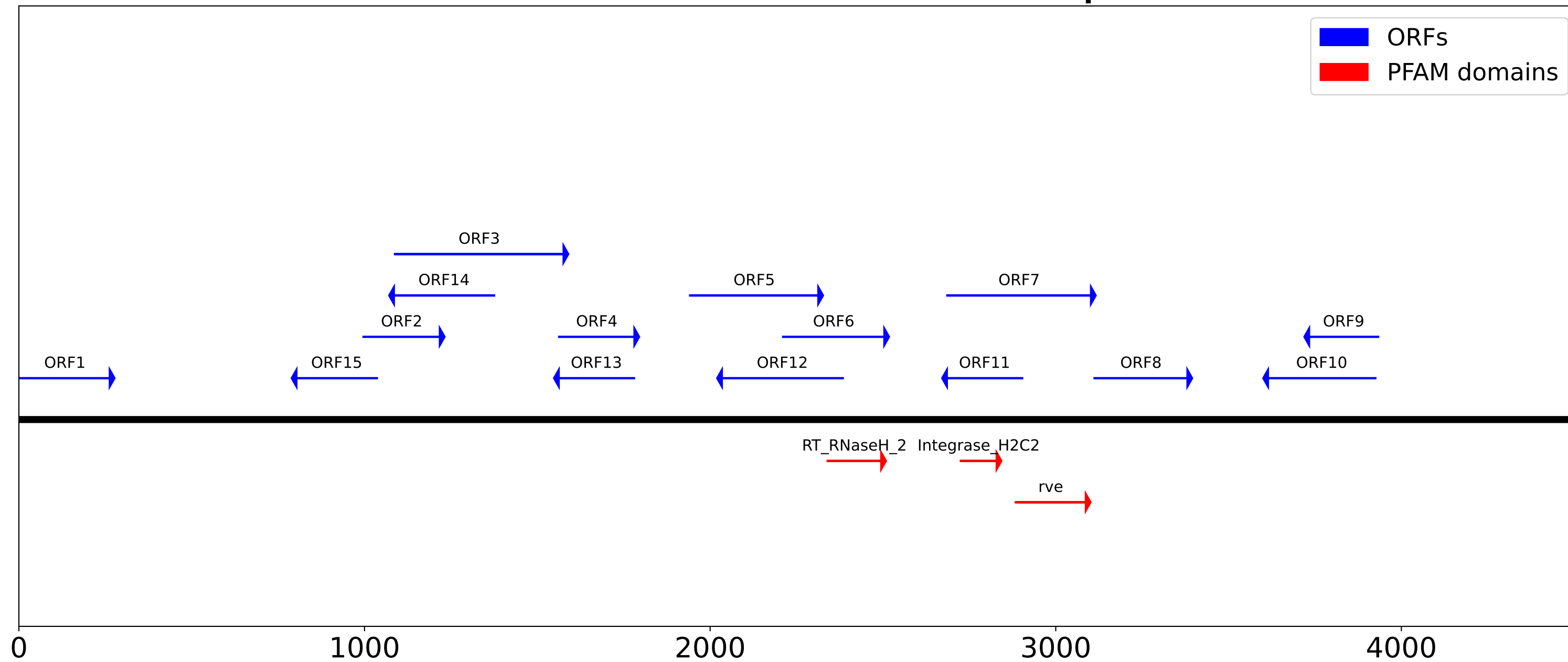


TE: ltr_1_family_325
size: 261bp; fragments: 11; full length: 11 (≥ 234.9 bp)



Before TEtrimmer 261 bp

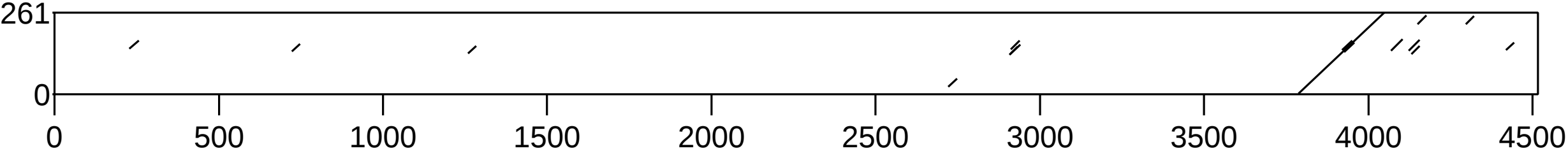
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)