

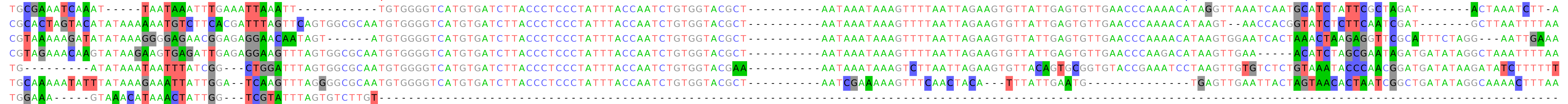
Start crop Point



End crop Point



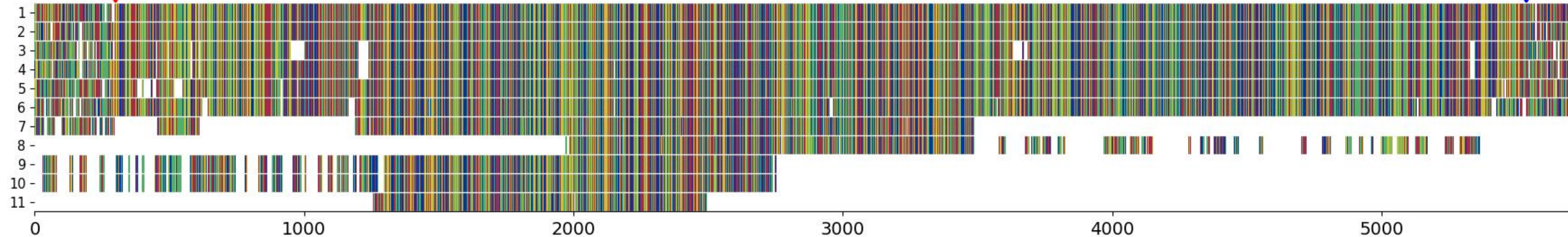
MSA length = 5239



TGTTCAAATAAAT - - - - - TAGACTTTTATATCCACACGGAAACCA - - - - - CG - - - - -
TGTTCAAATAAAT - - - - - TAGACTTTTATATCCACACGGAAACCA - - - - - CG - - - - -

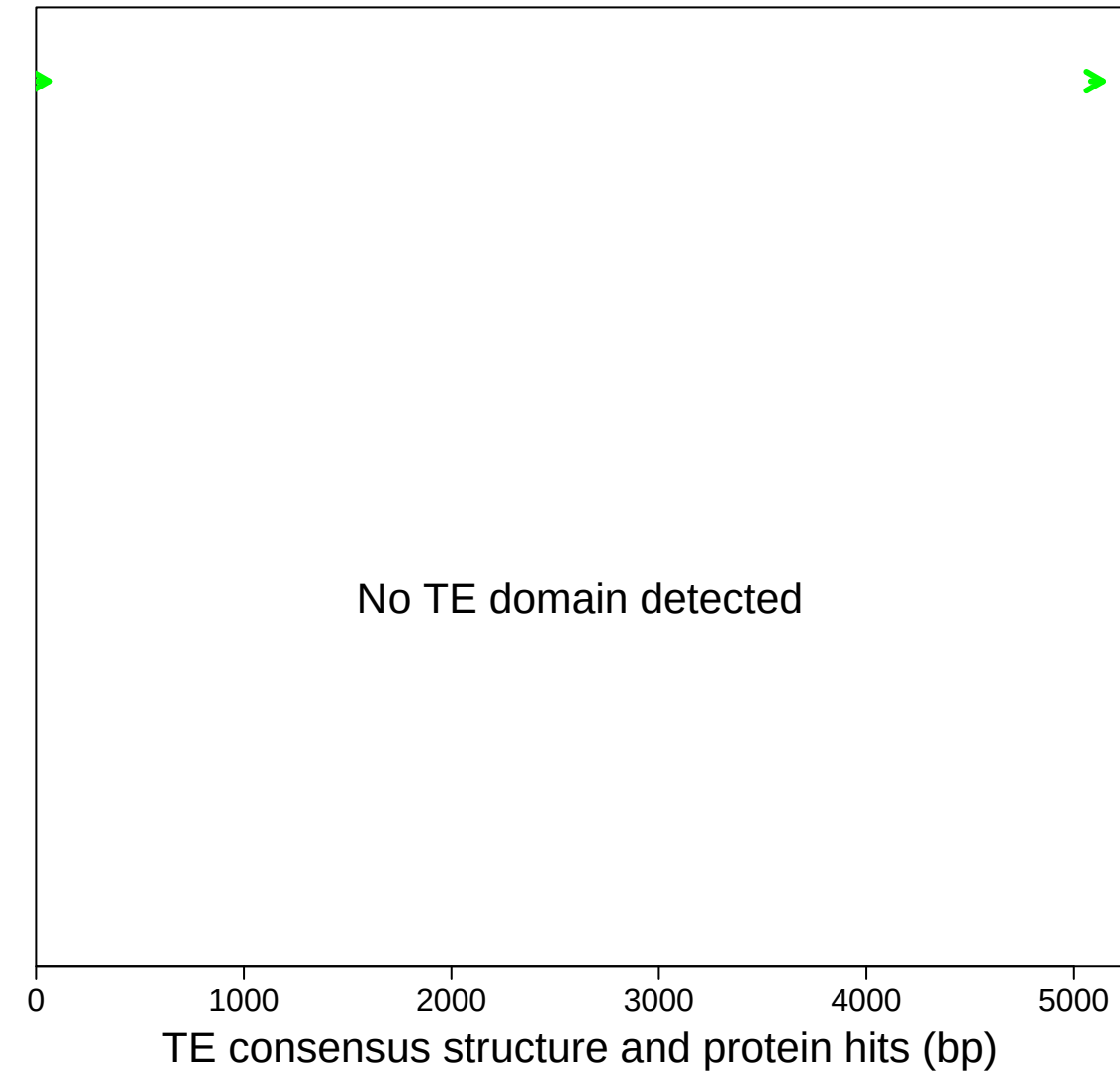
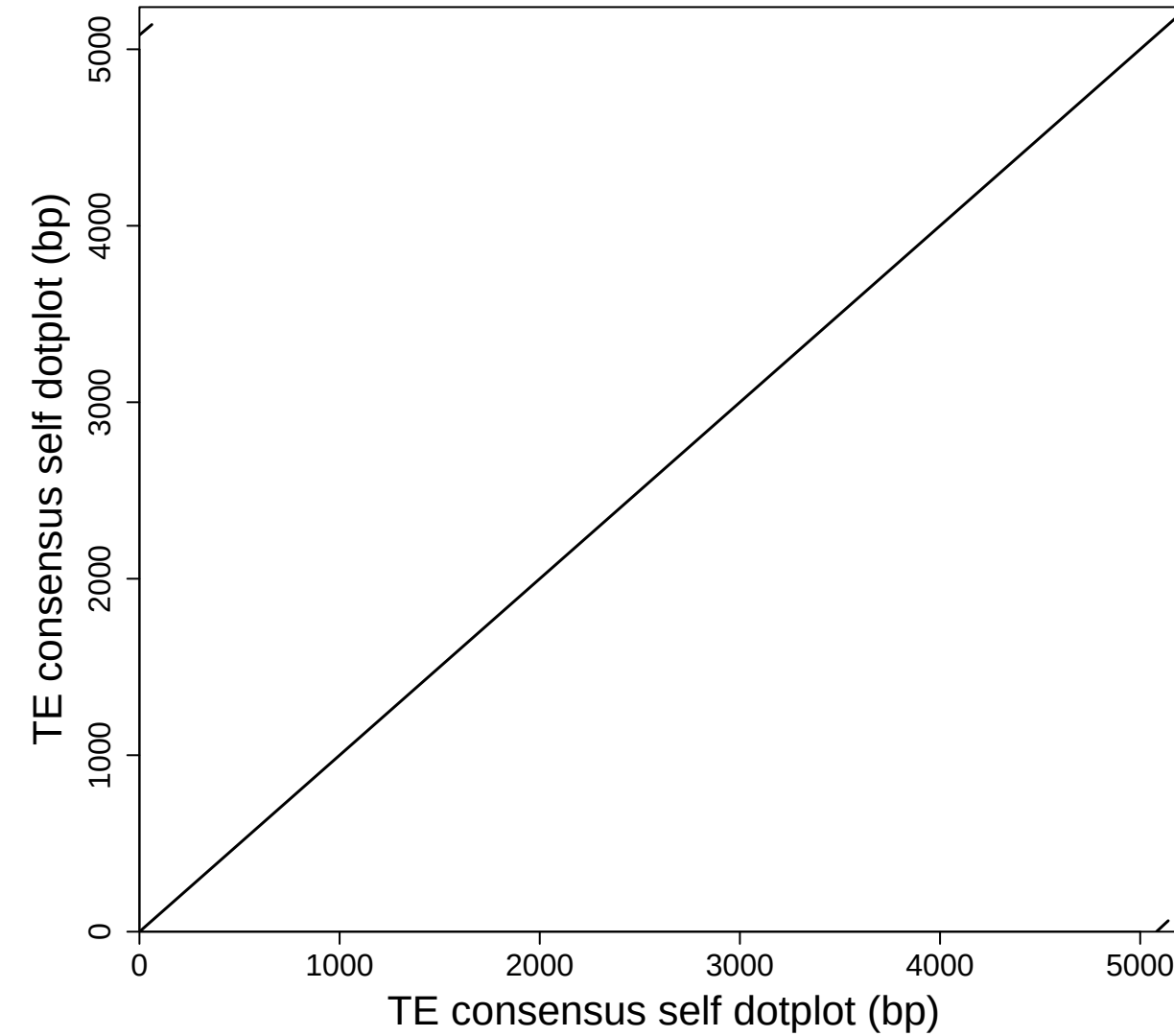
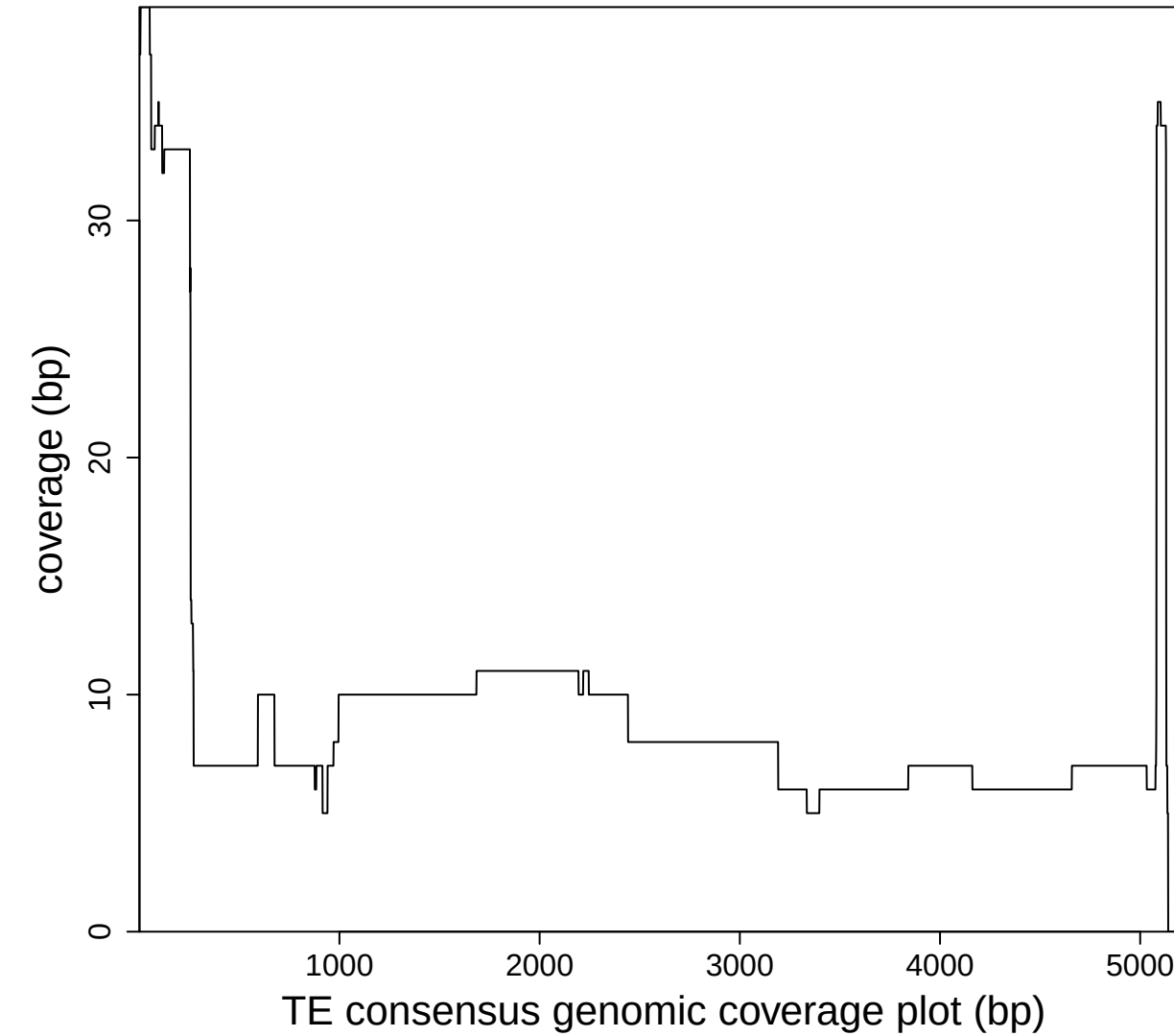
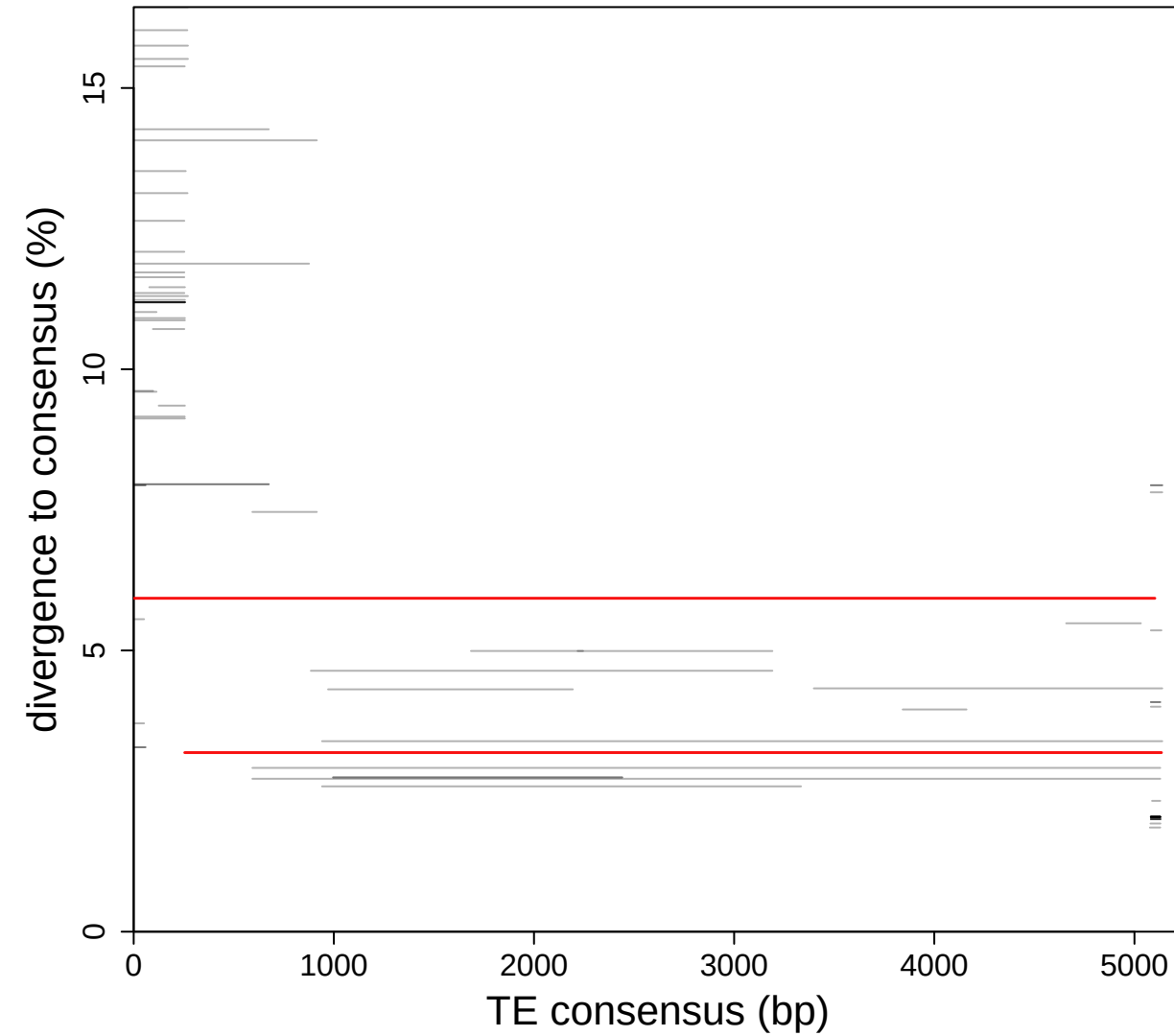
Start crop Point

End crop Point



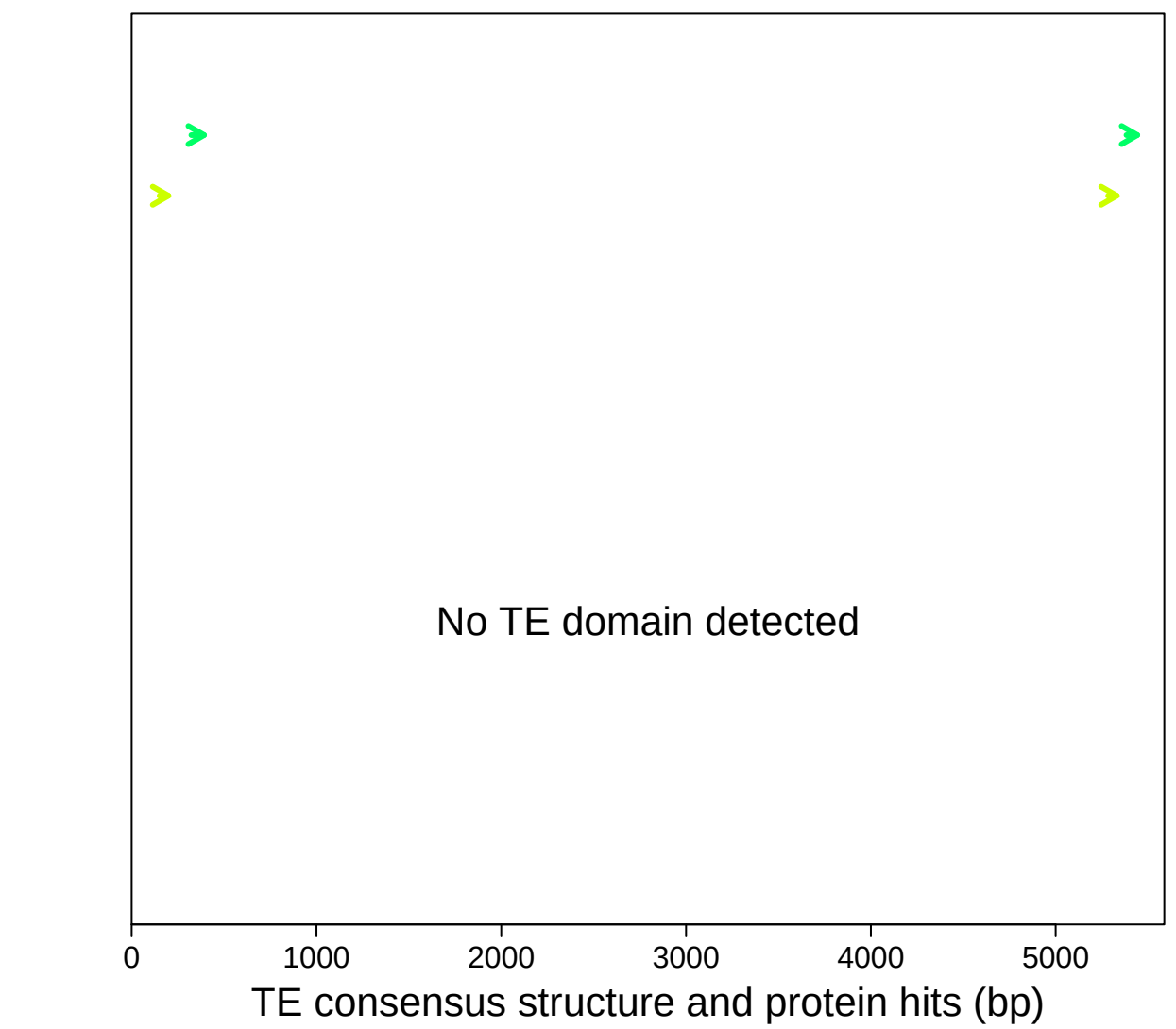
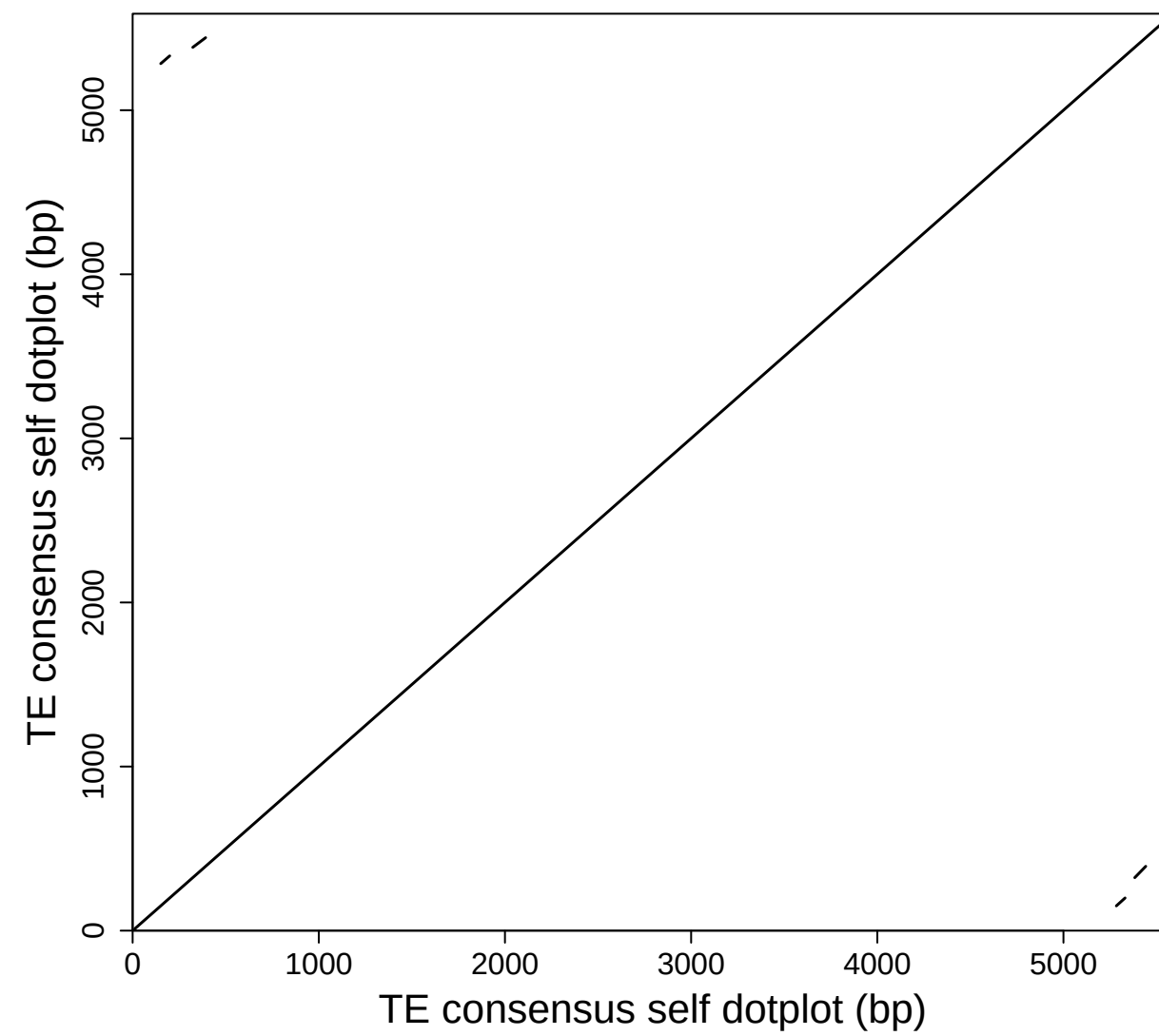
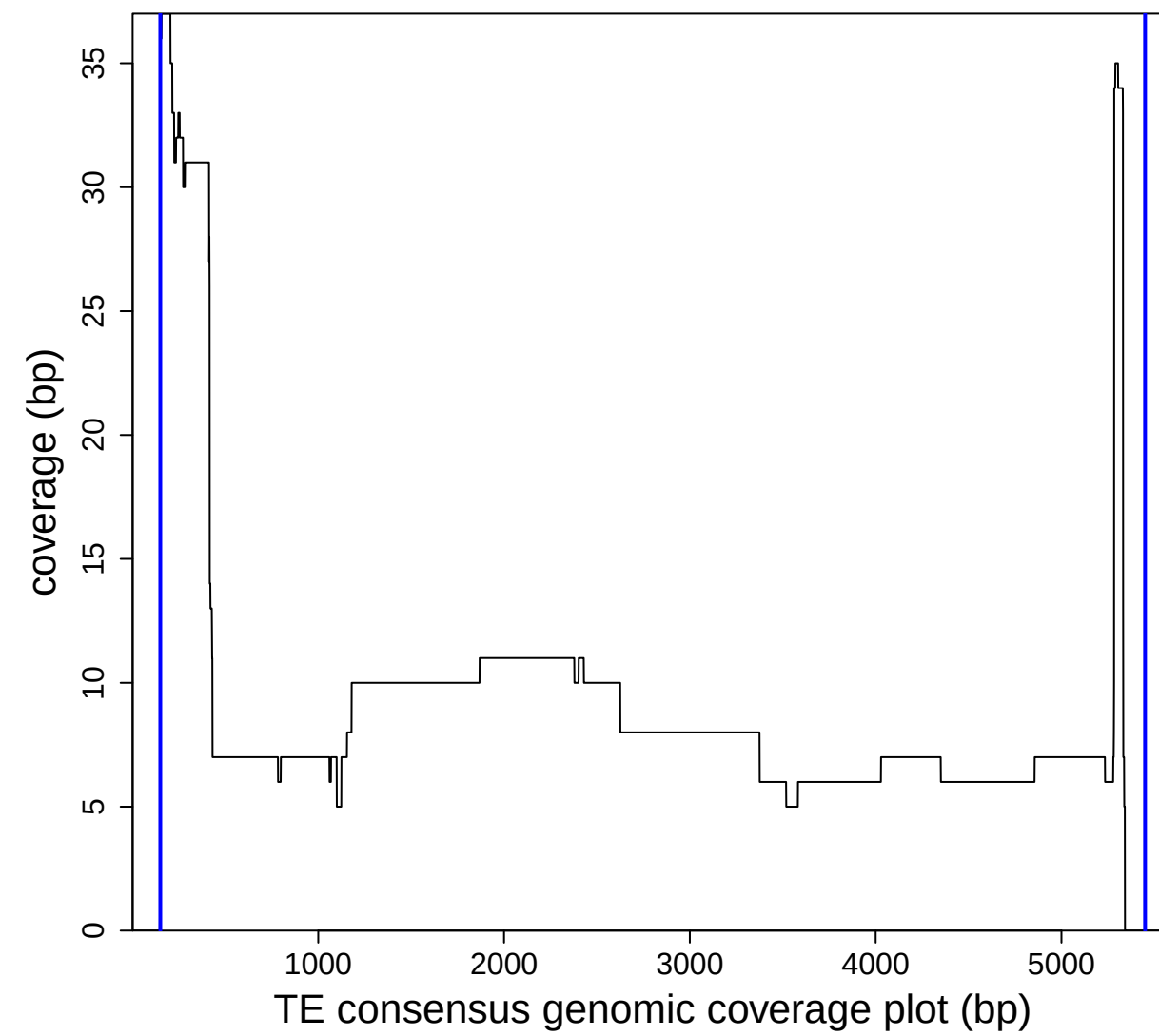
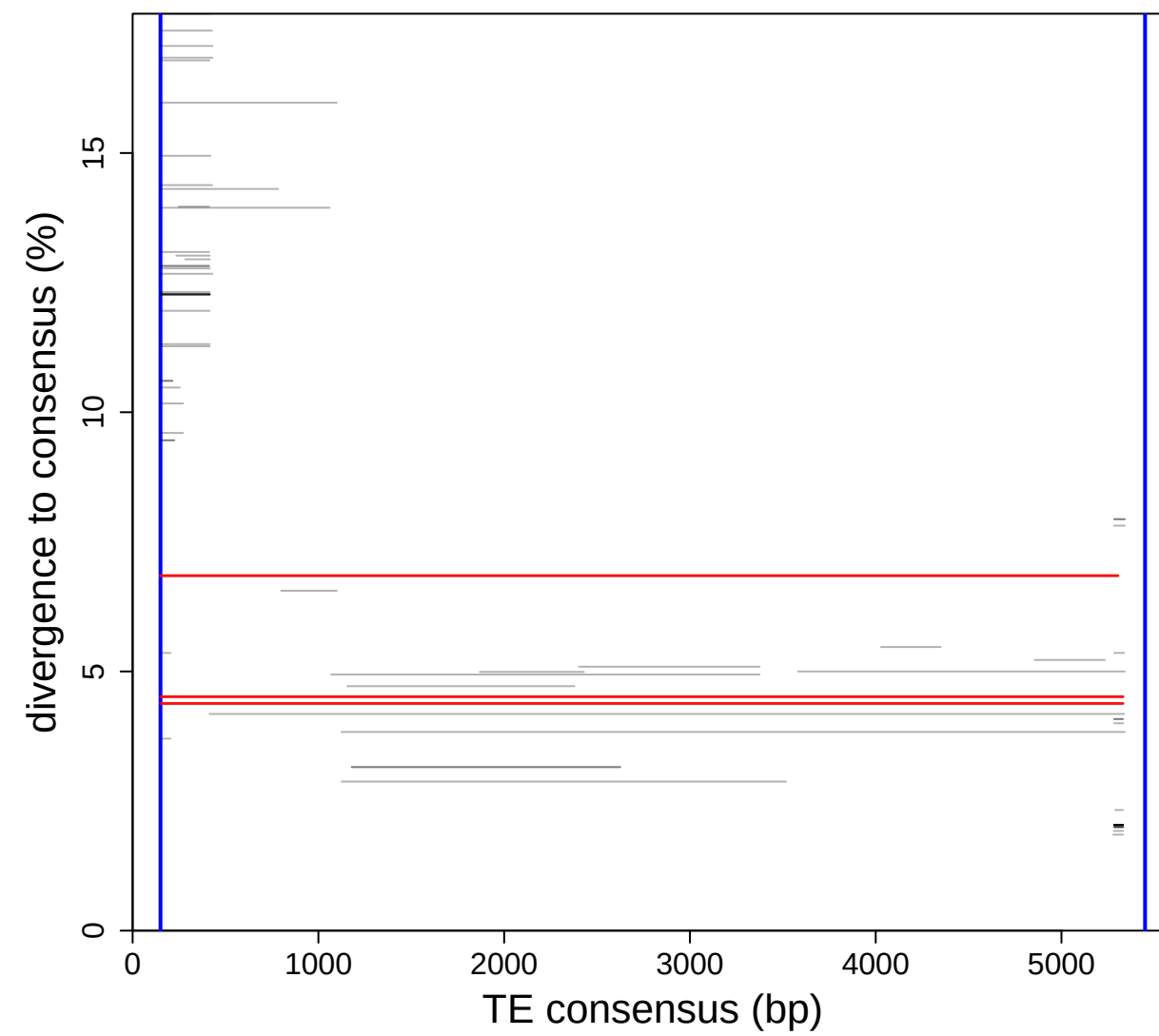
b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 5239bp; fragments: 86; full length: 2 (≥ 4715.1 bp)

After TETrimmer 5239 bp



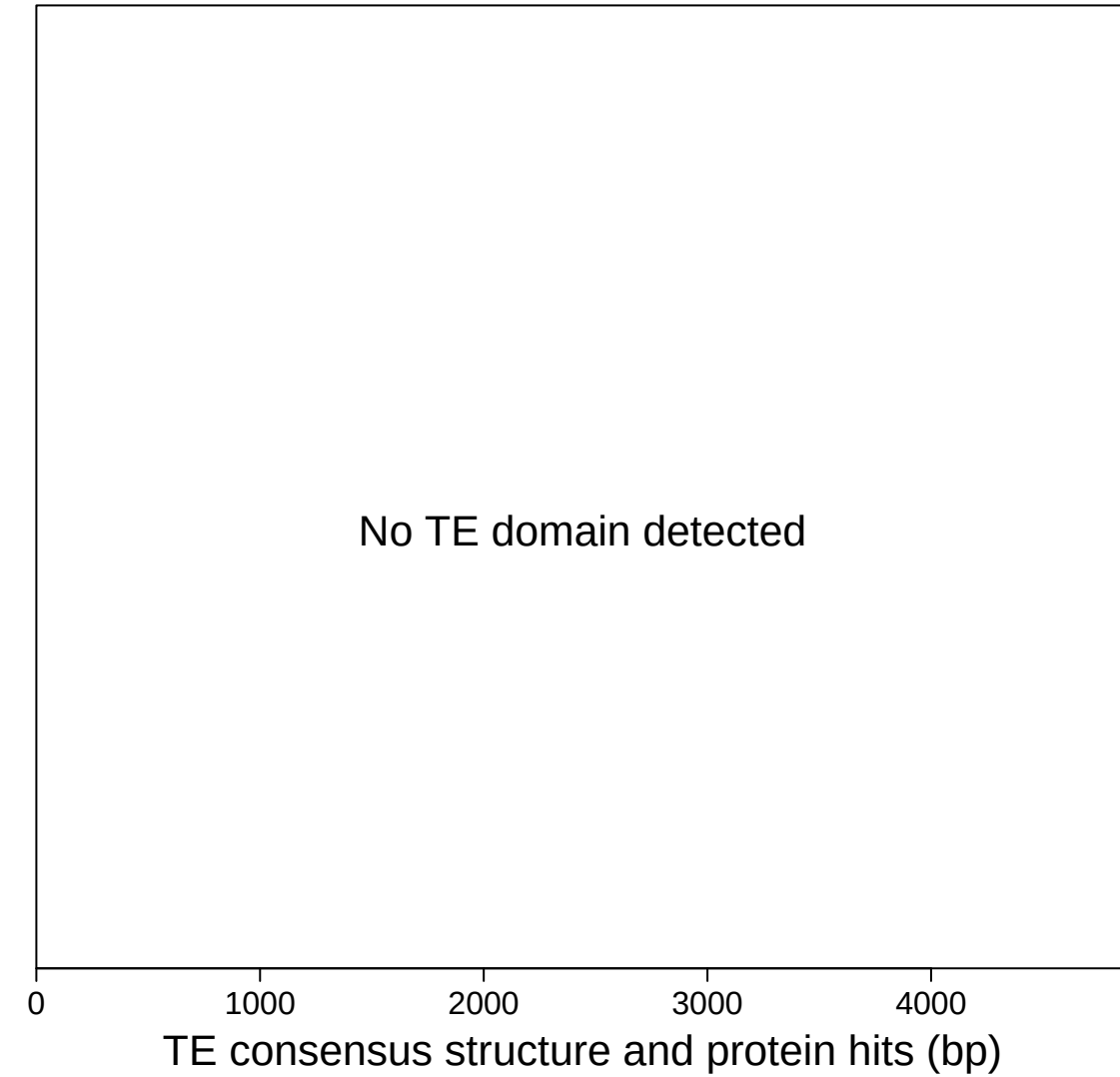
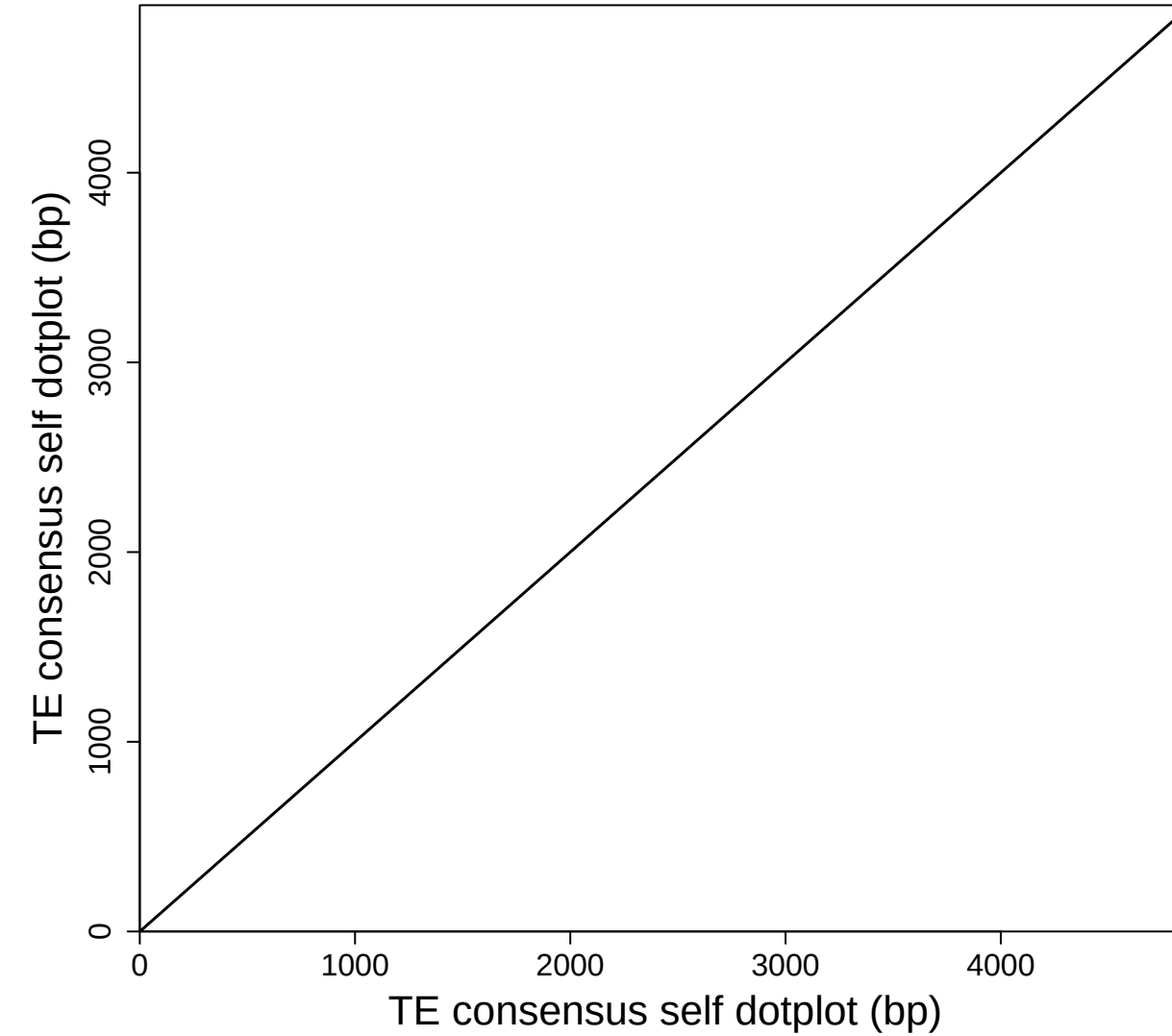
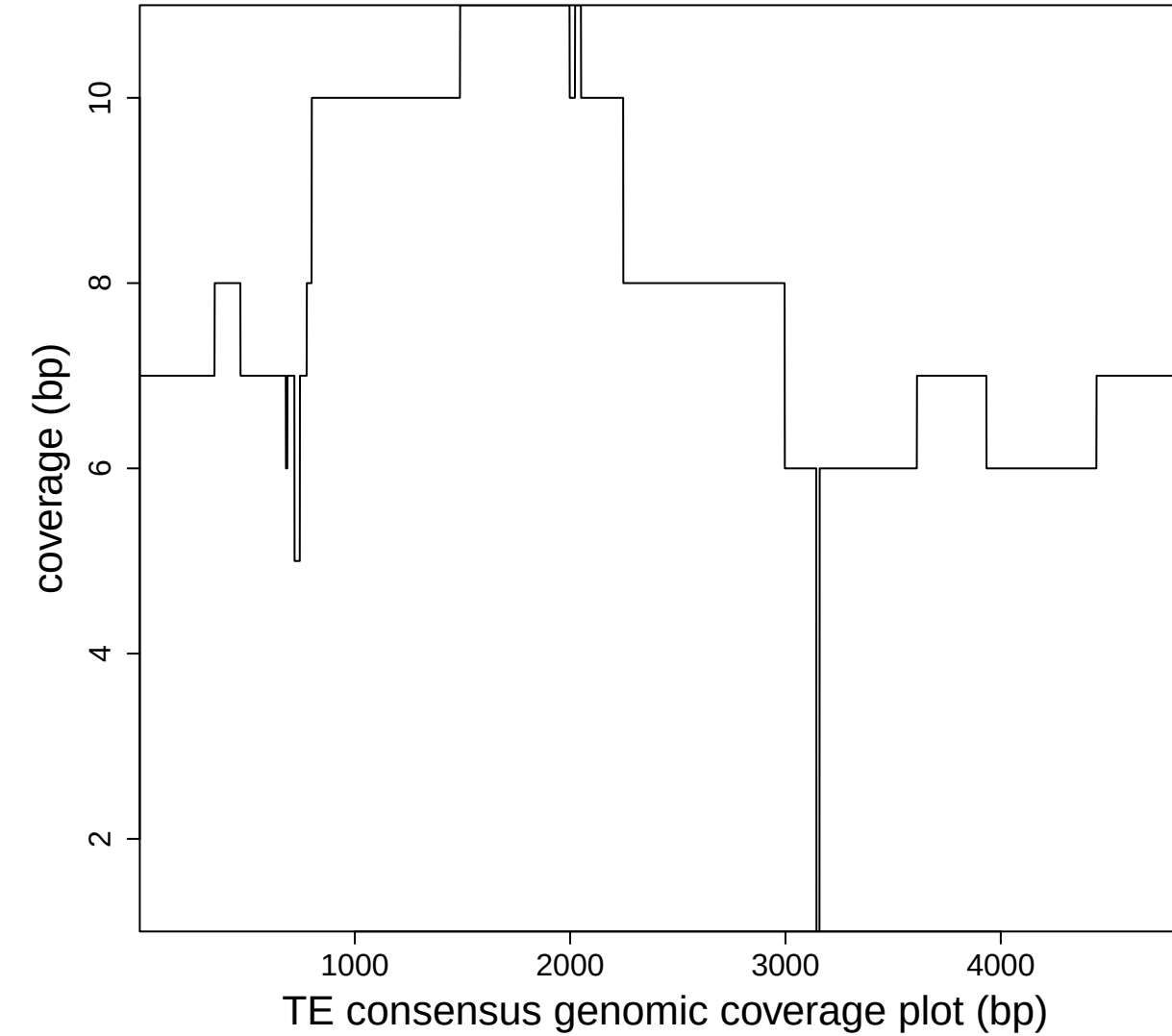
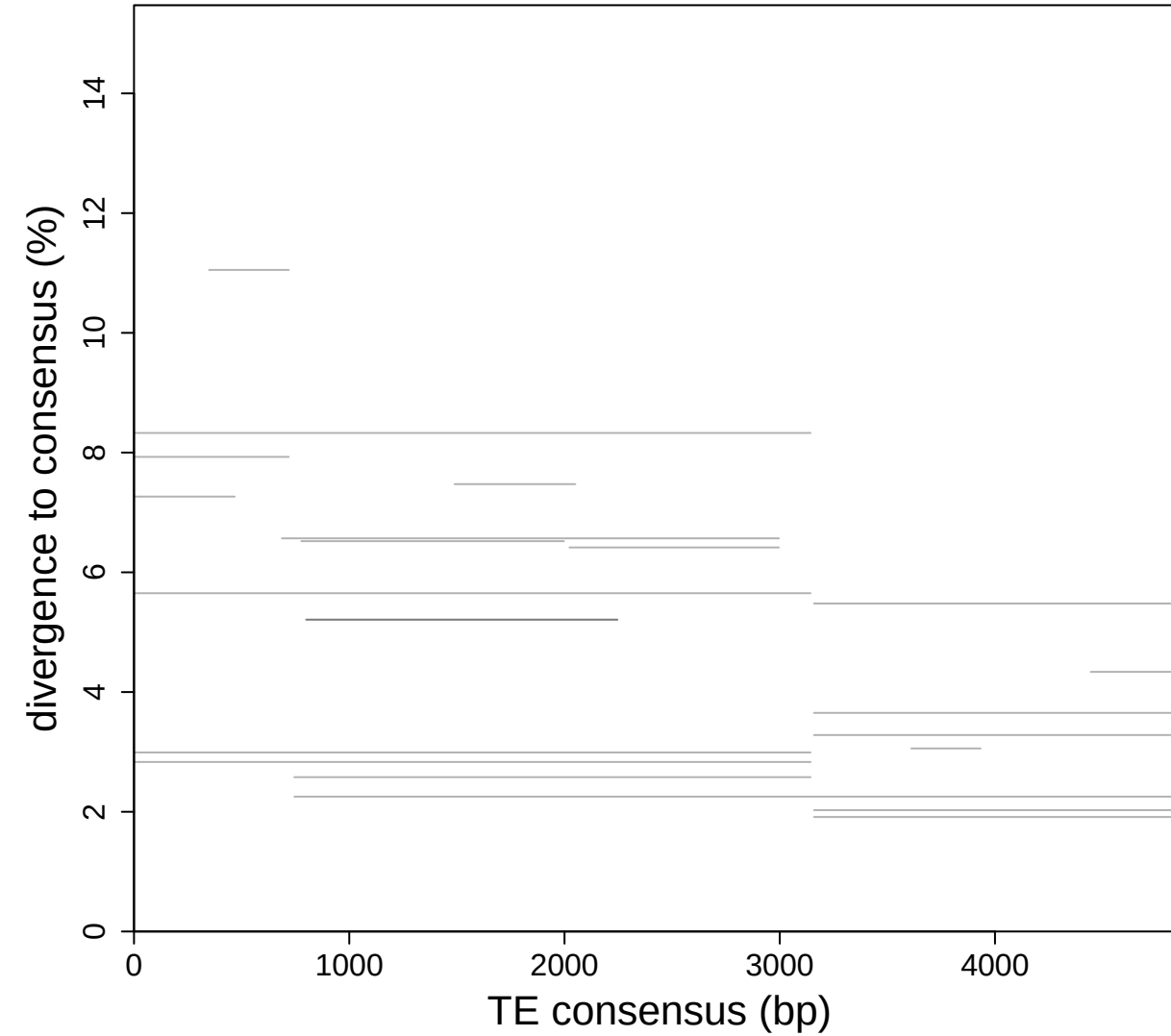
.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 5600bp; fragments: 82; full length: 3 (>=5040bp)

After TEtrimmer Extended plot Blue lines are boundaries

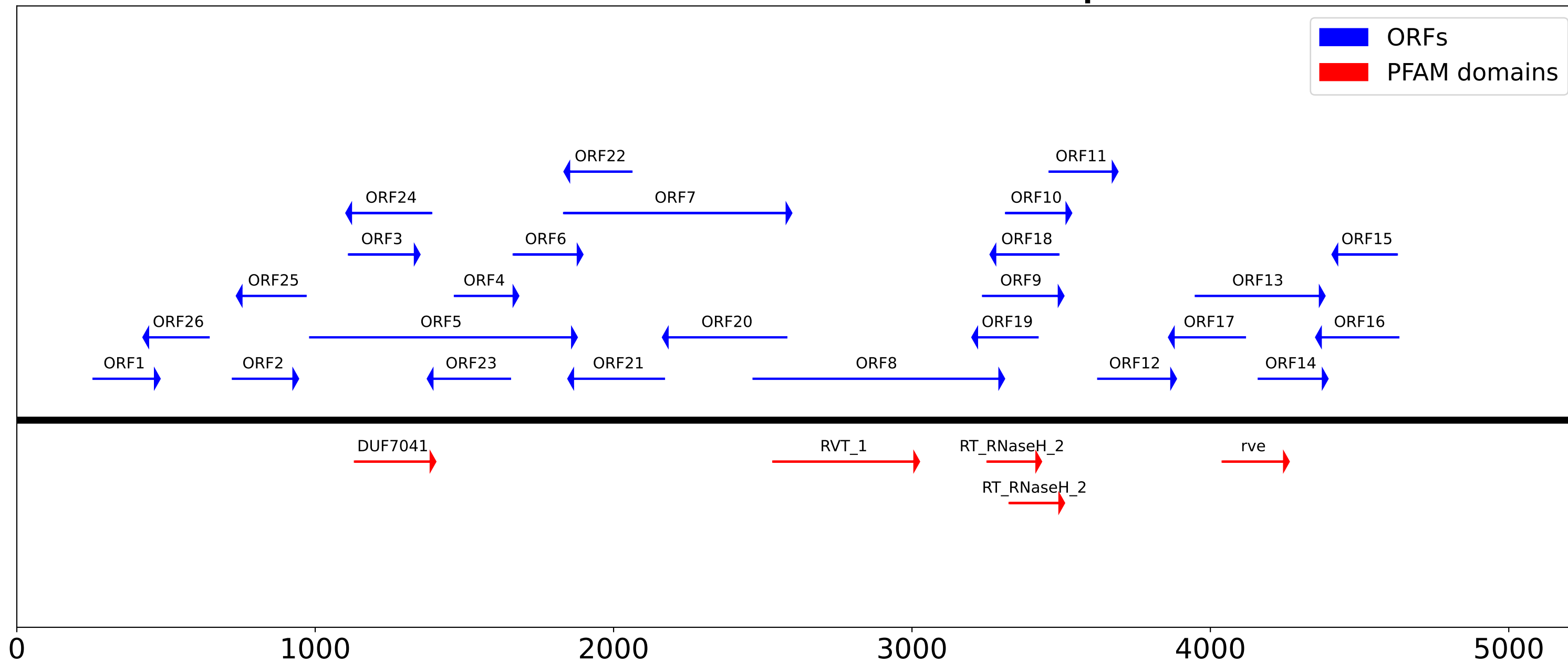


TE: ltr_1_family_631
size: 4883bp; fragments: 23; full length: 0 (≥ 4394.7 bp)

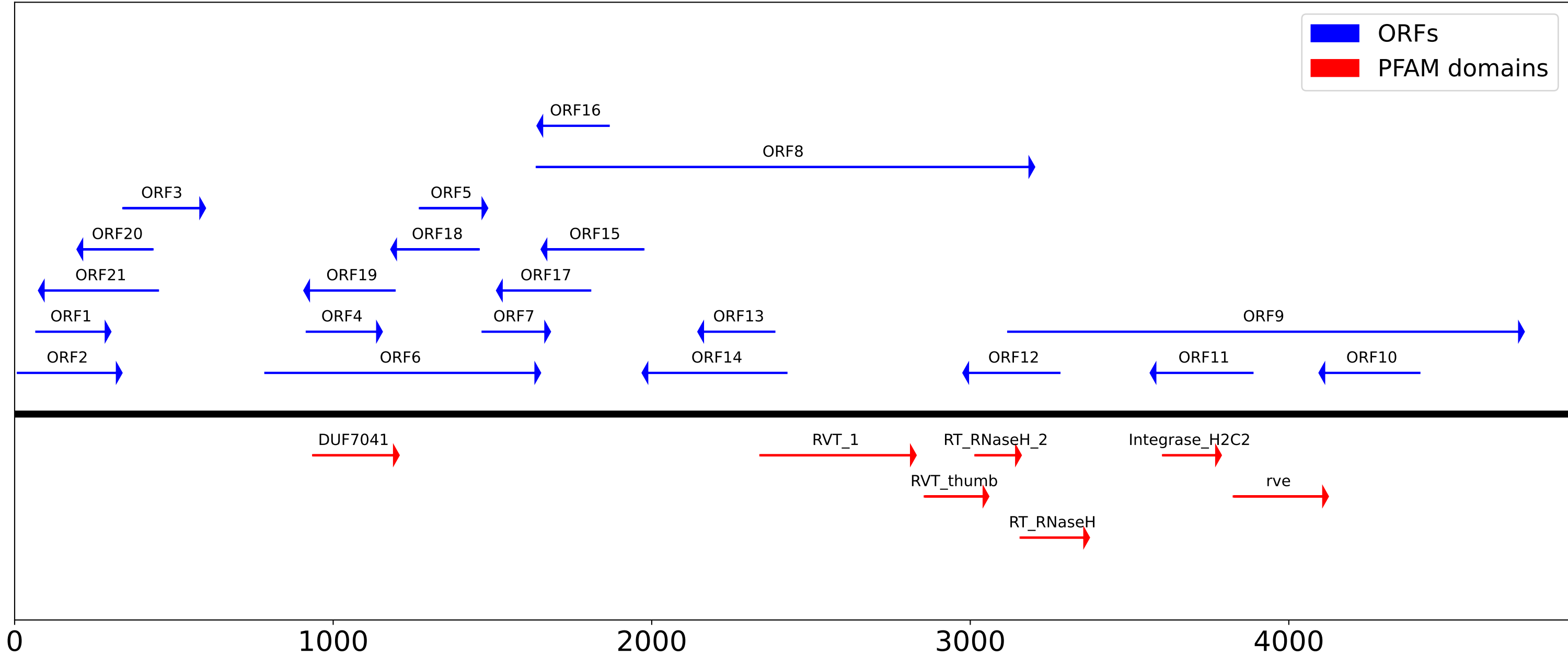
Before TEtrimmer 4883 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

