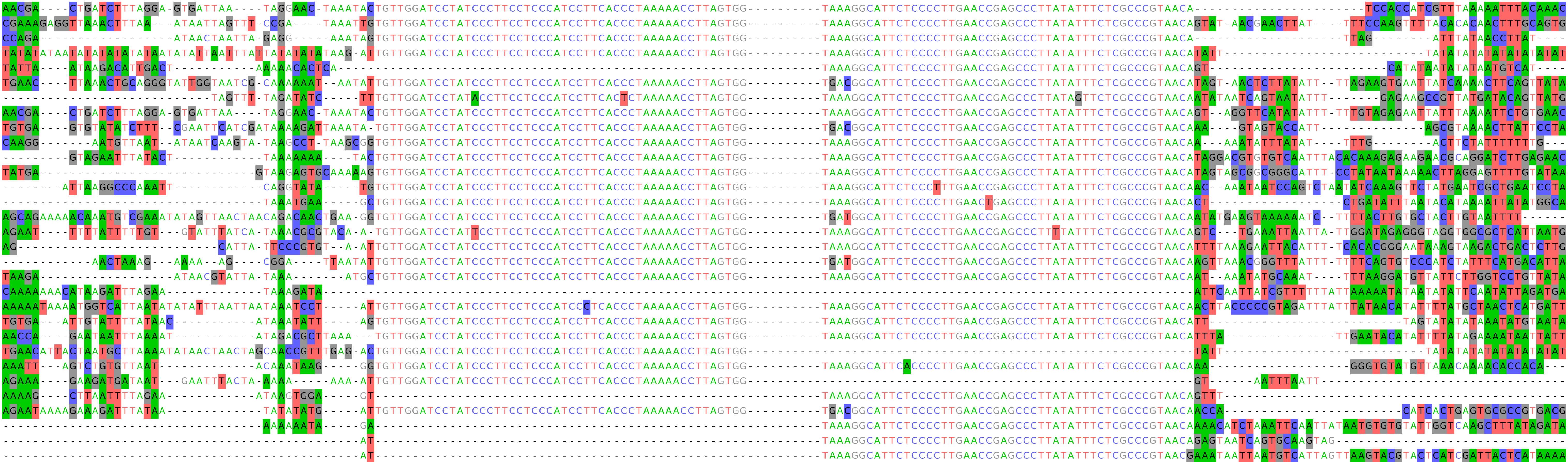


MSA length = 6788

Start crop Point

End crop Point



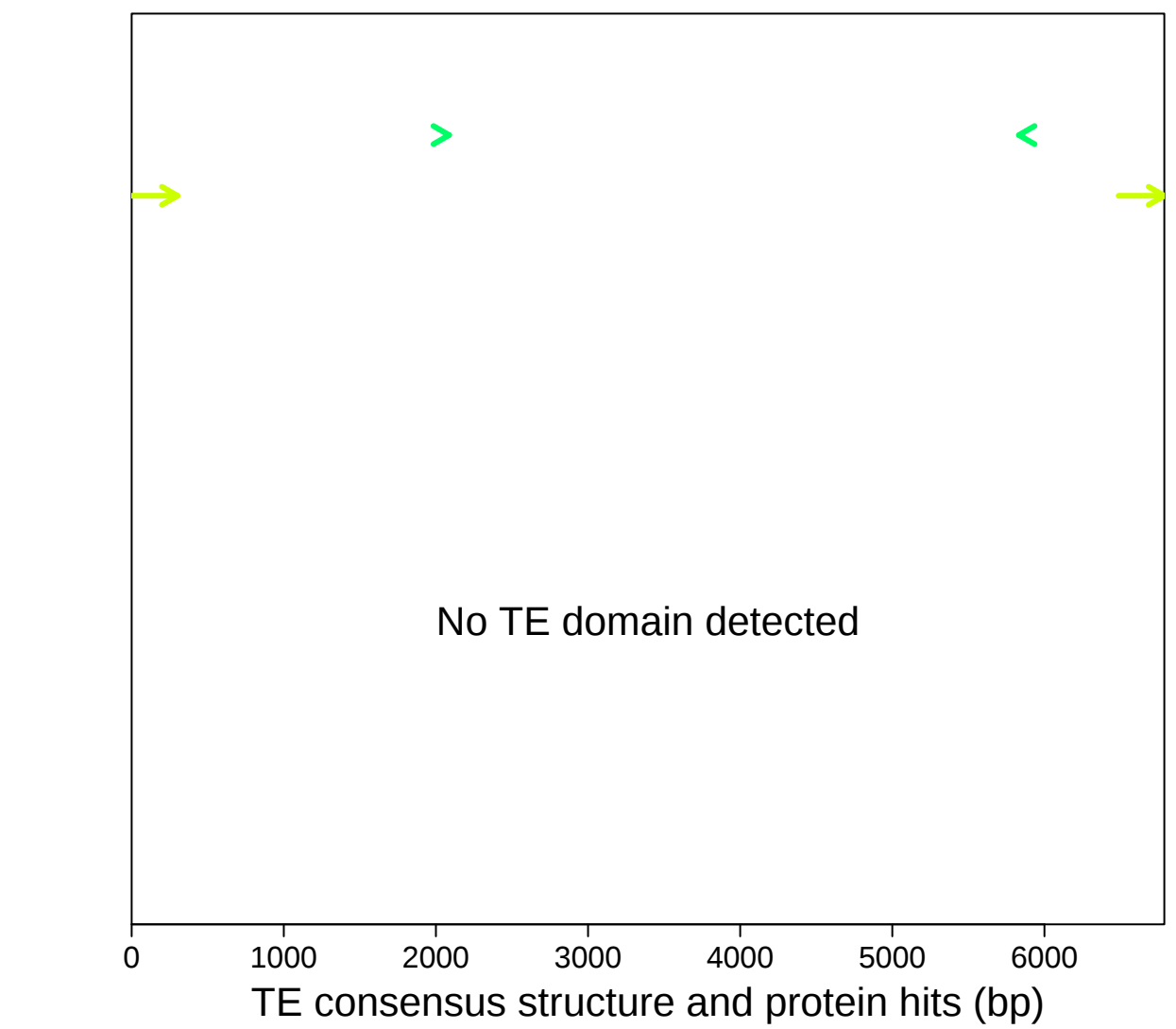
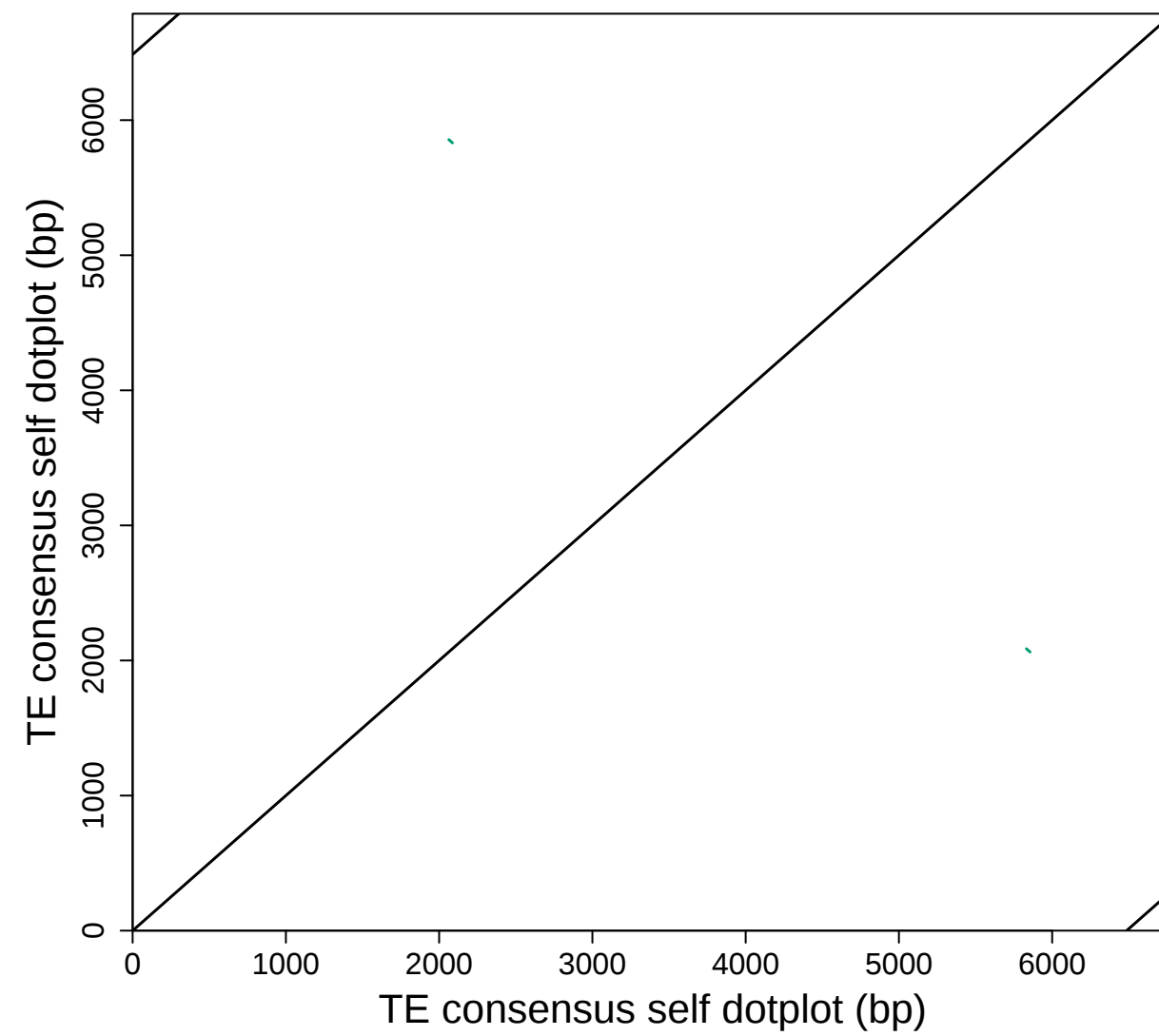
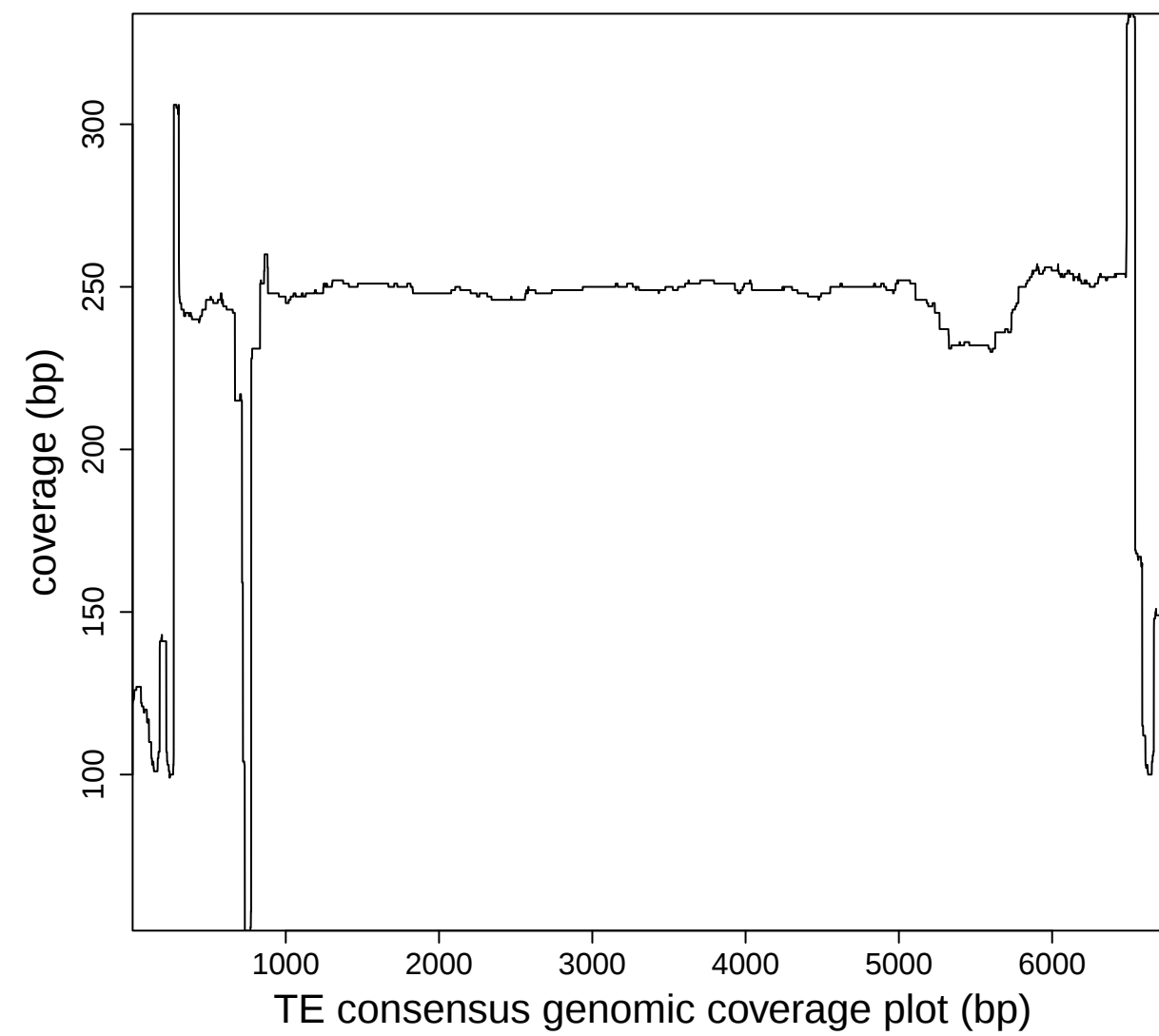
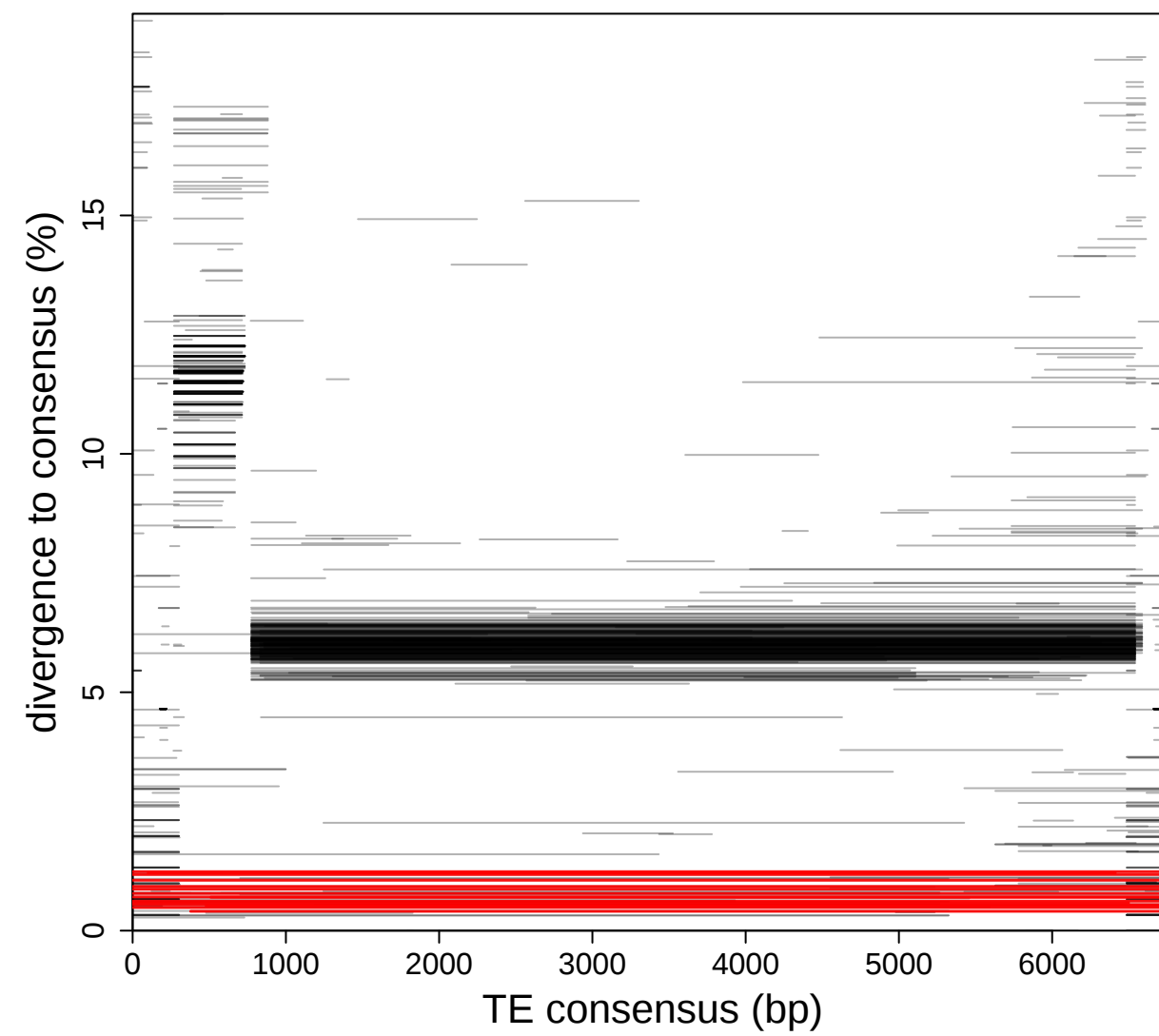
Start crop Point

End crop Point



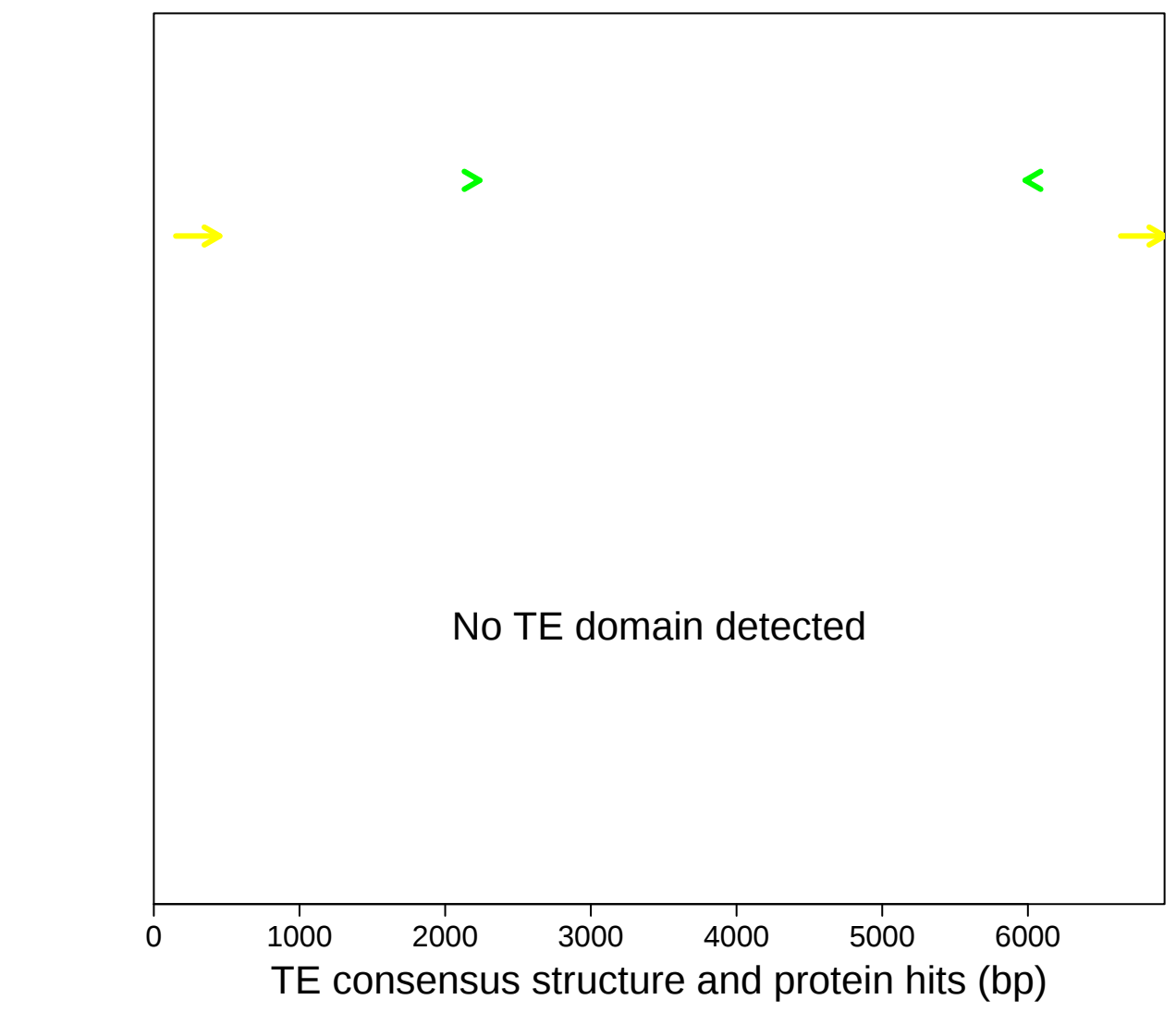
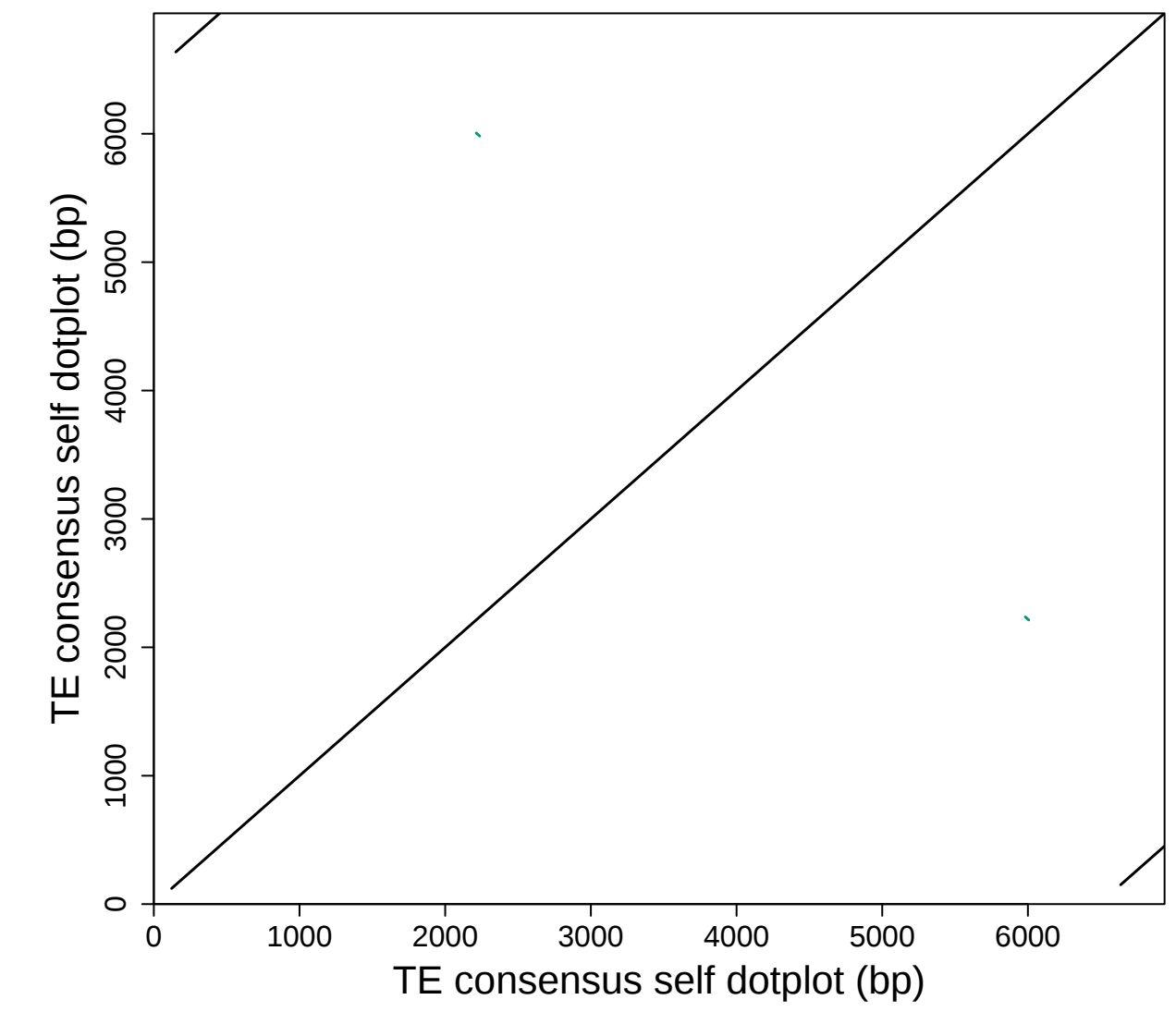
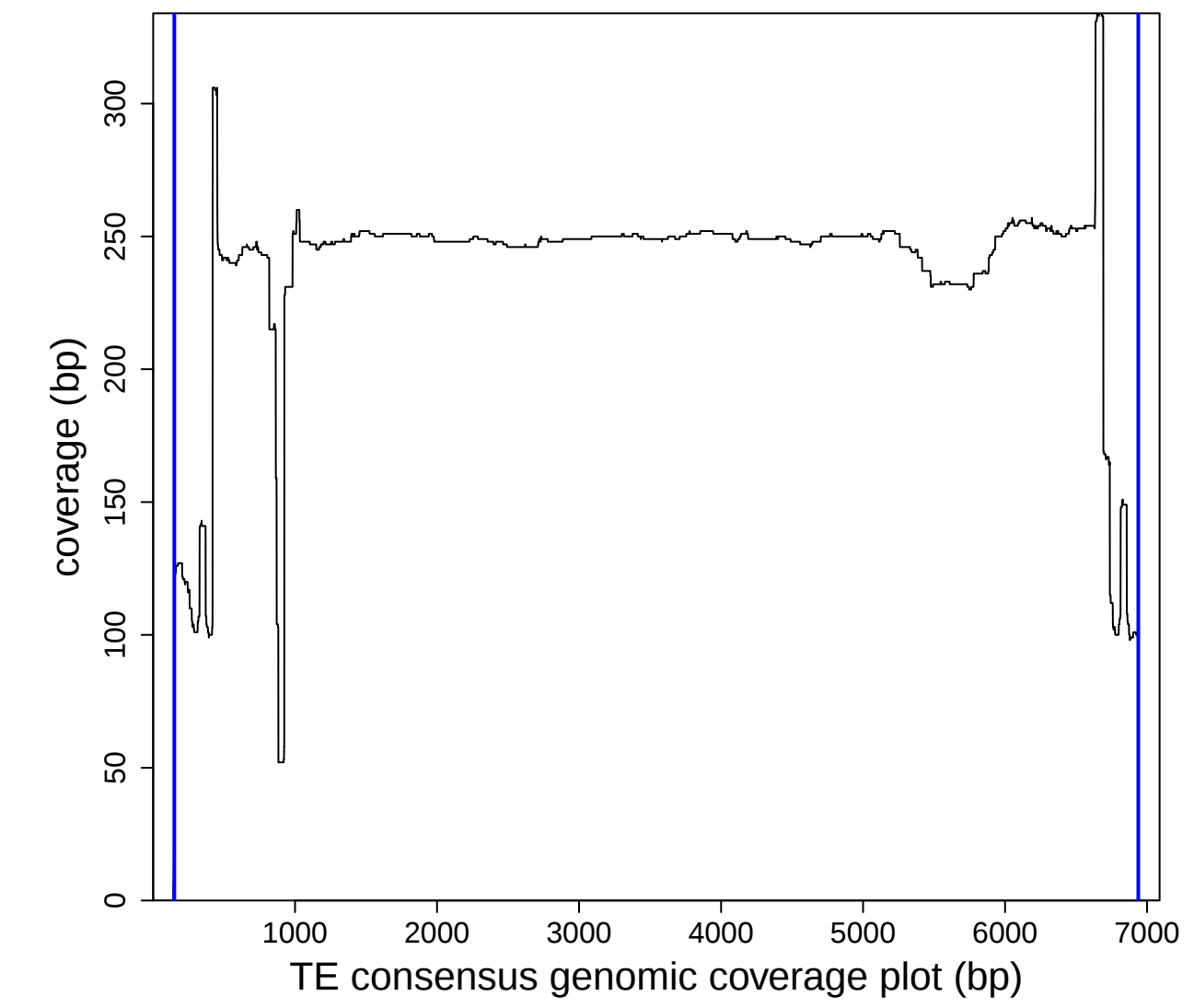
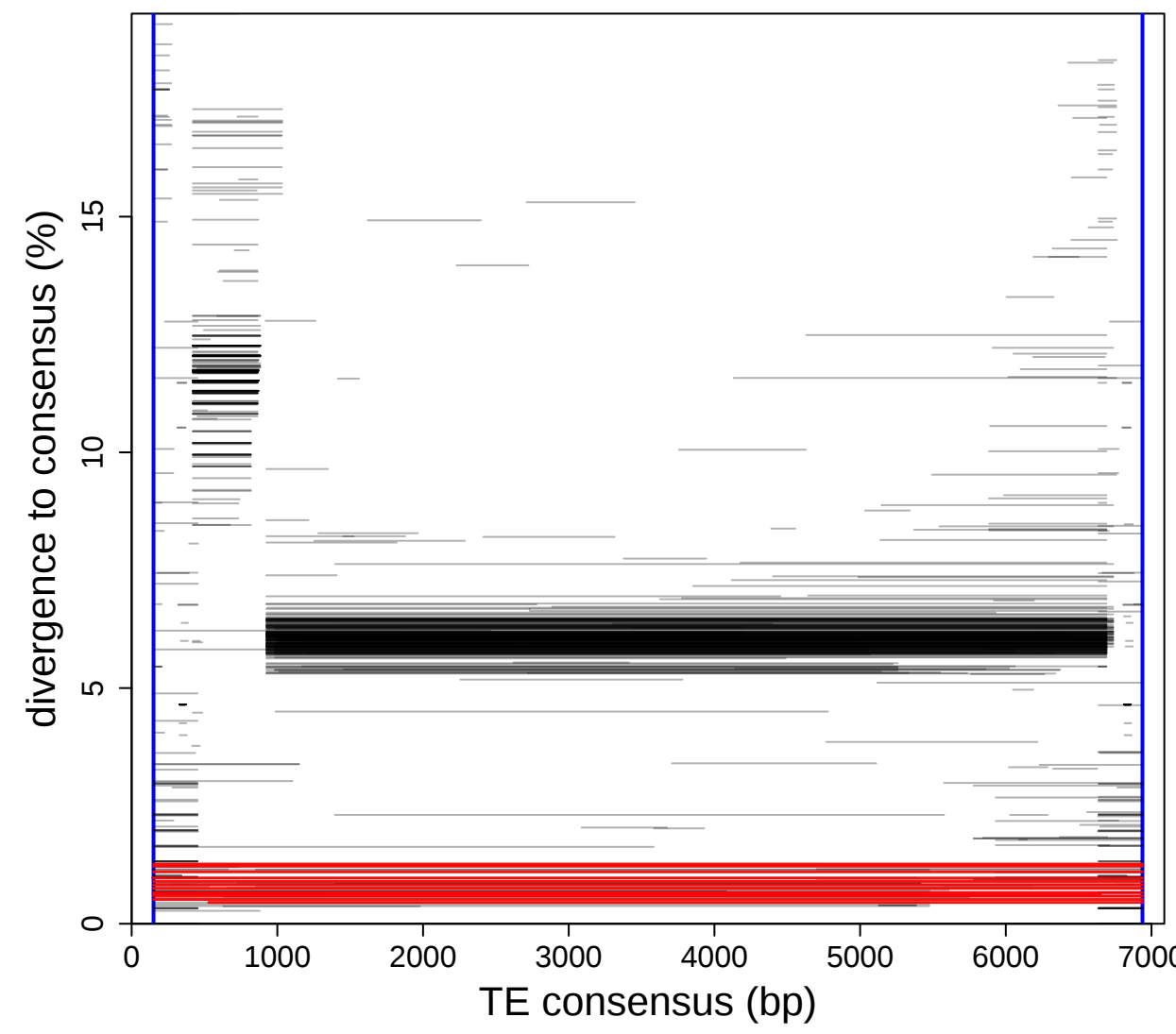
ed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 6788bp; fragments: 856; full length: 15 (≥ 6109.2 bp)

After TEtrimmer 6788 bp



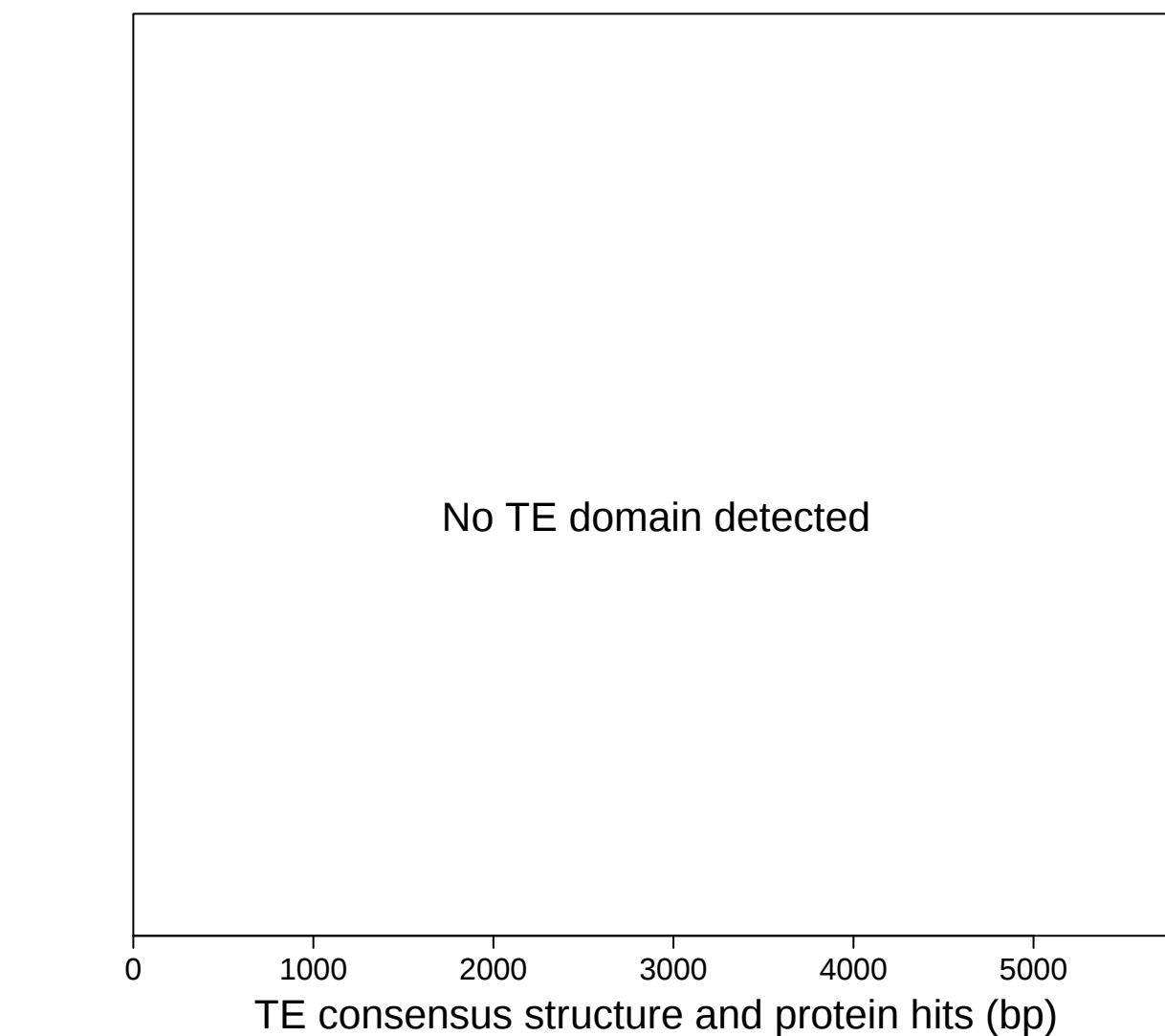
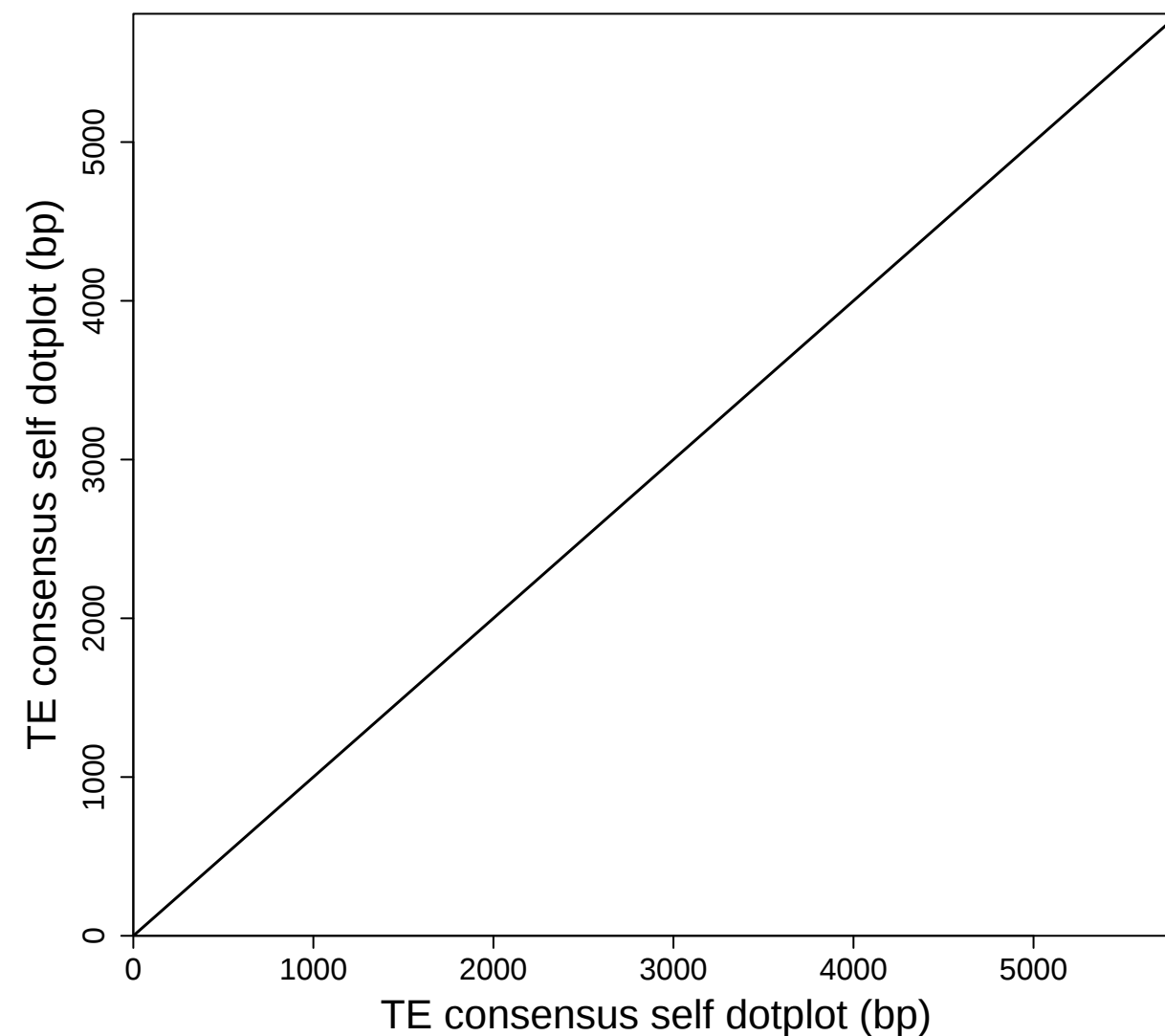
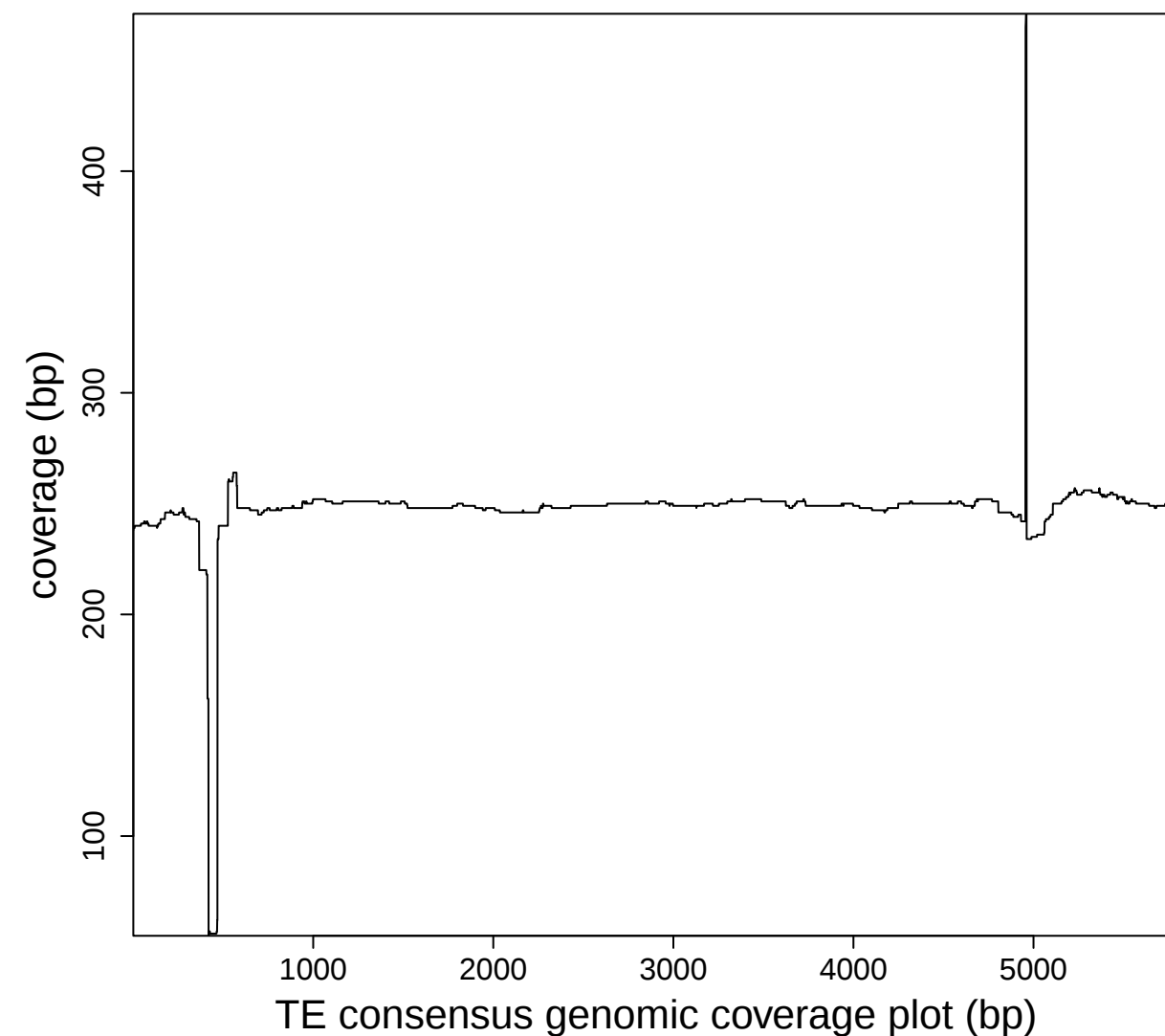
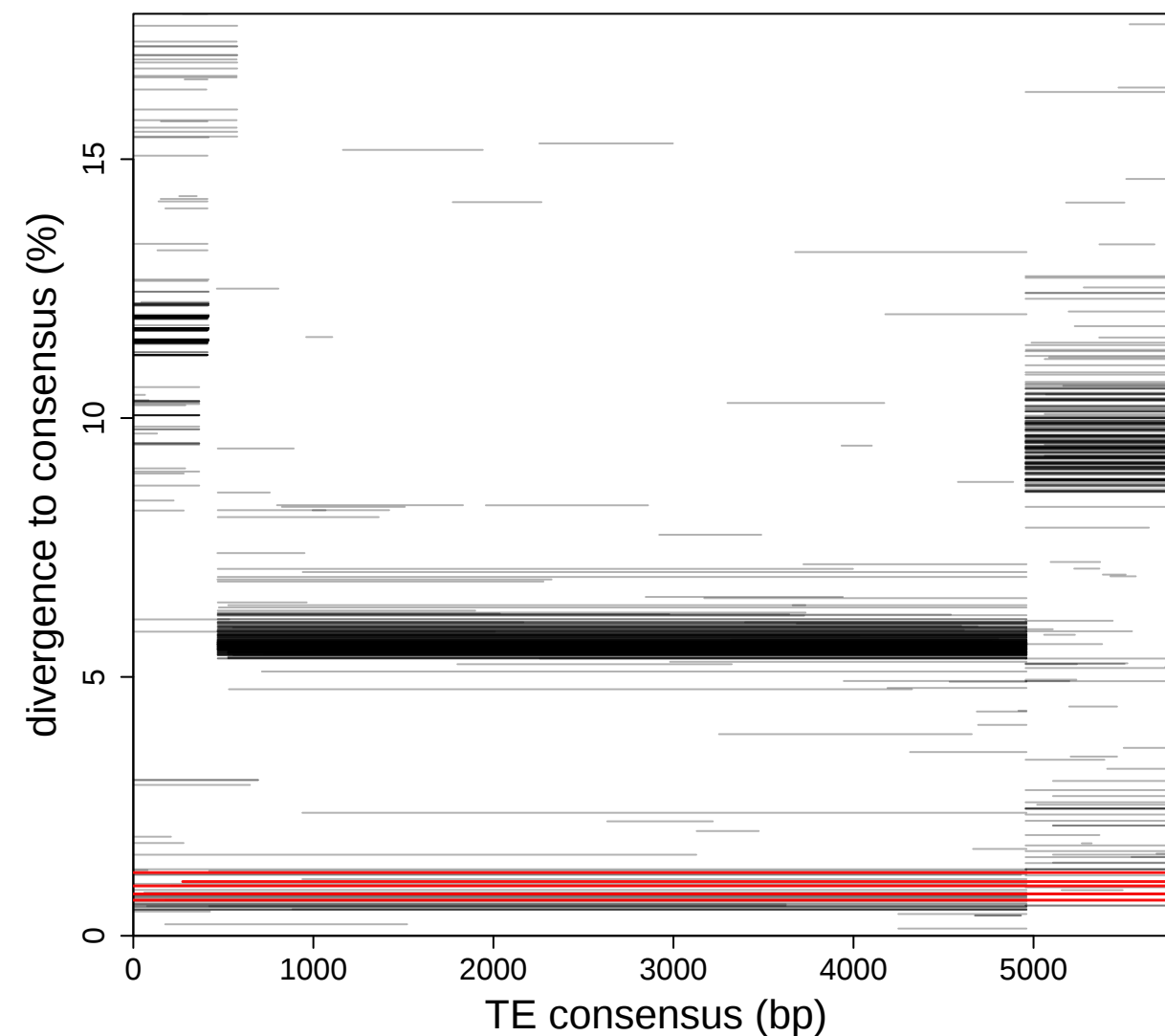
ly_756.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcIn.fa_aln.fa
size: 7088bp; fragments: 856; full length: 12 (>=6379.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

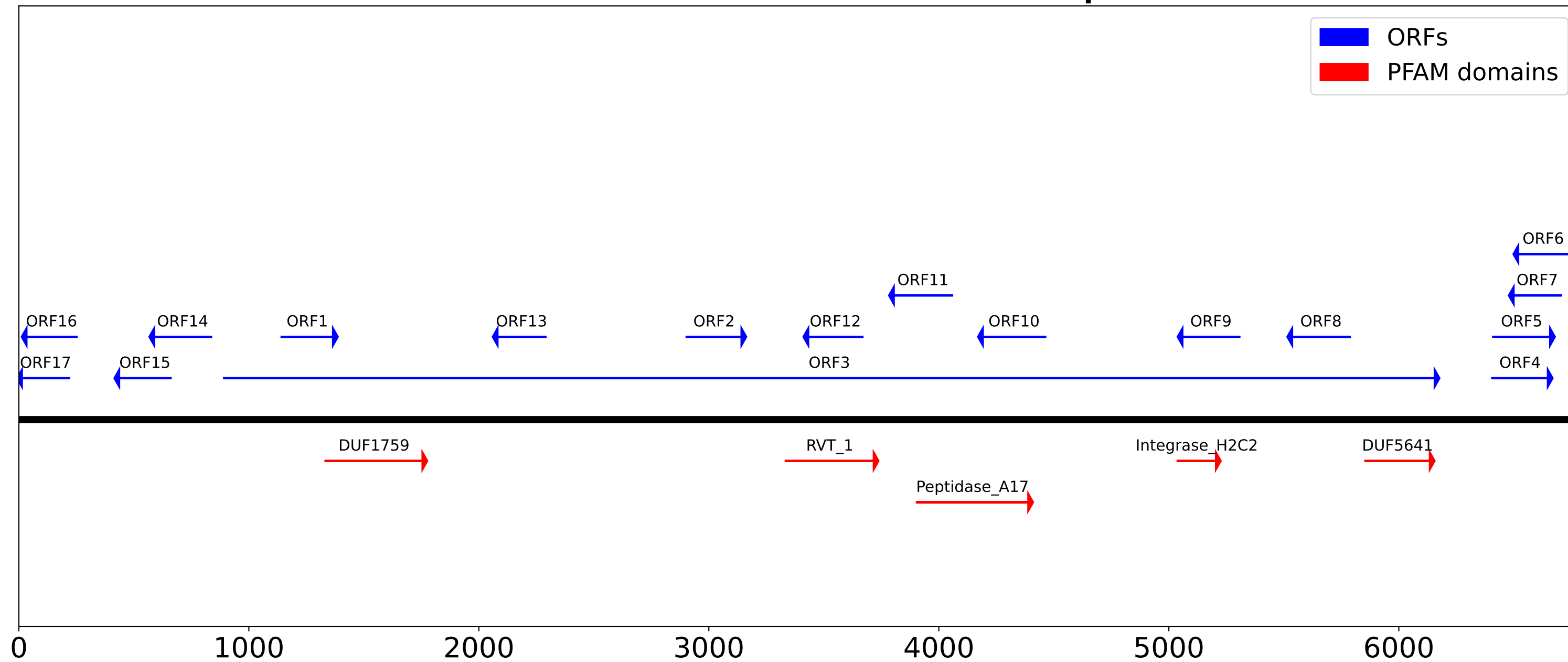


TE: ltr_1_family_756
size: 5808bp; fragments: 789; full length: 5 (≥ 5227.2 bp)

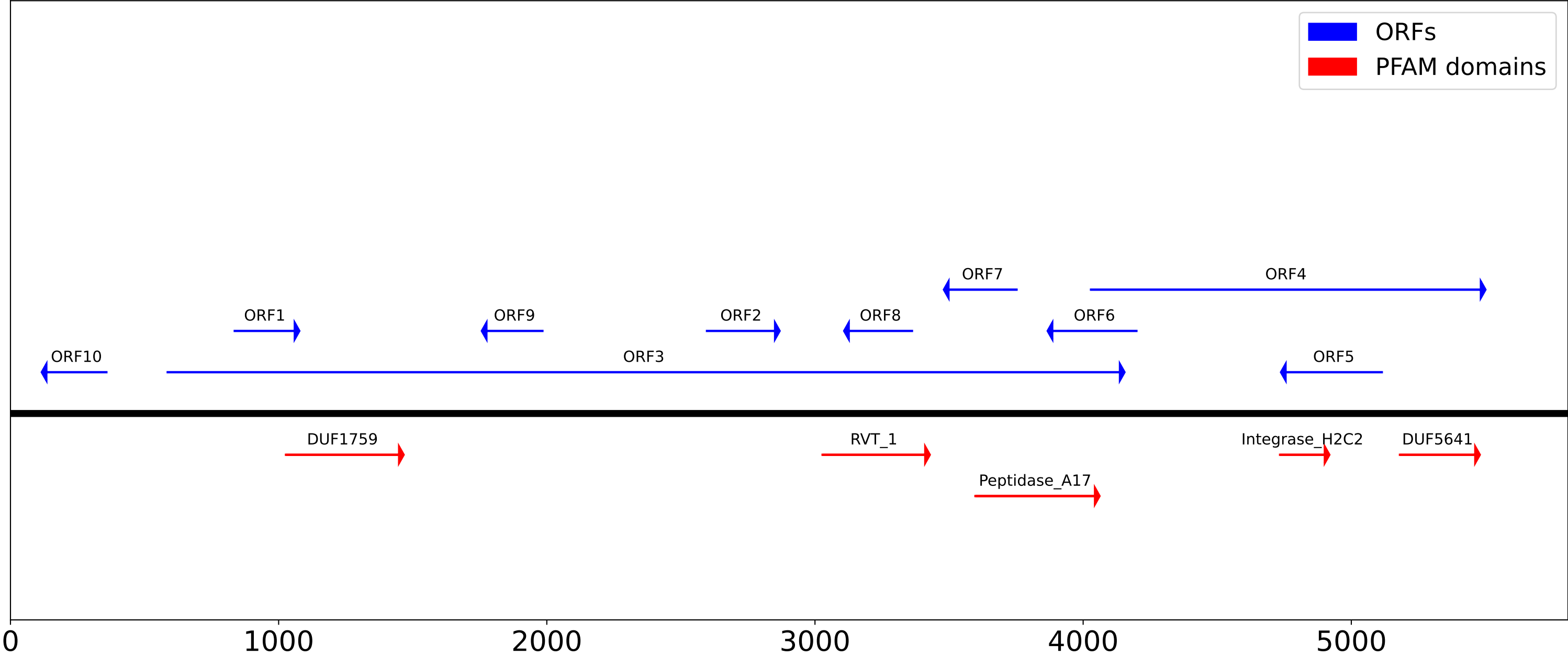
Before TEtrimmer 5808 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

