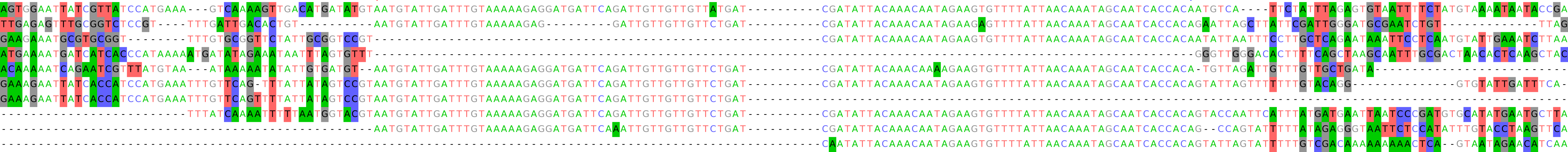


Start Crop Point

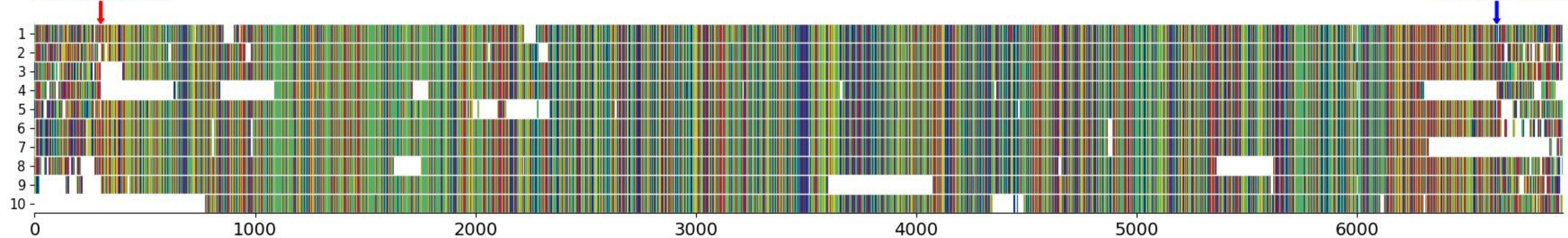
End Crop Point

MSA length = 6334



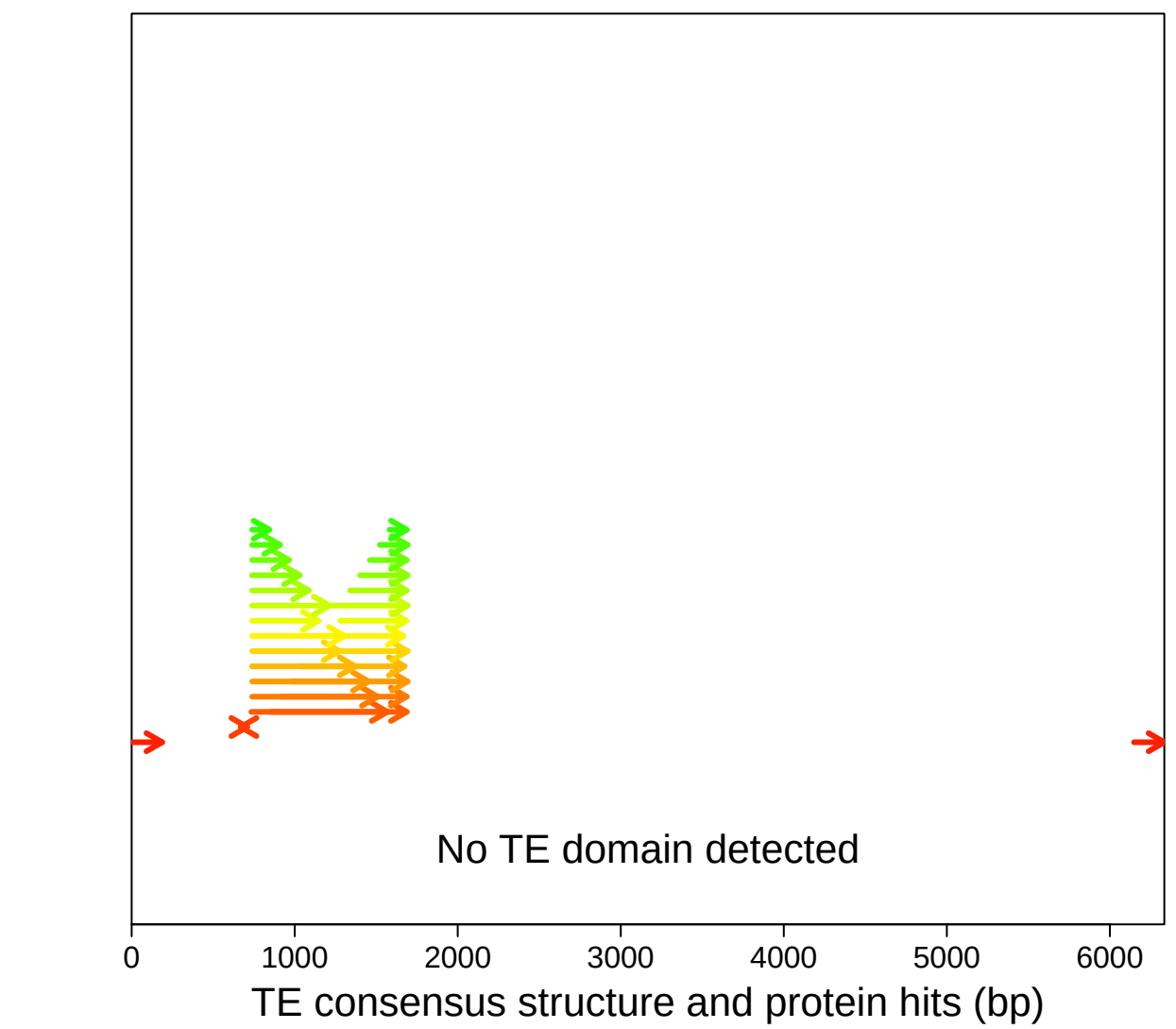
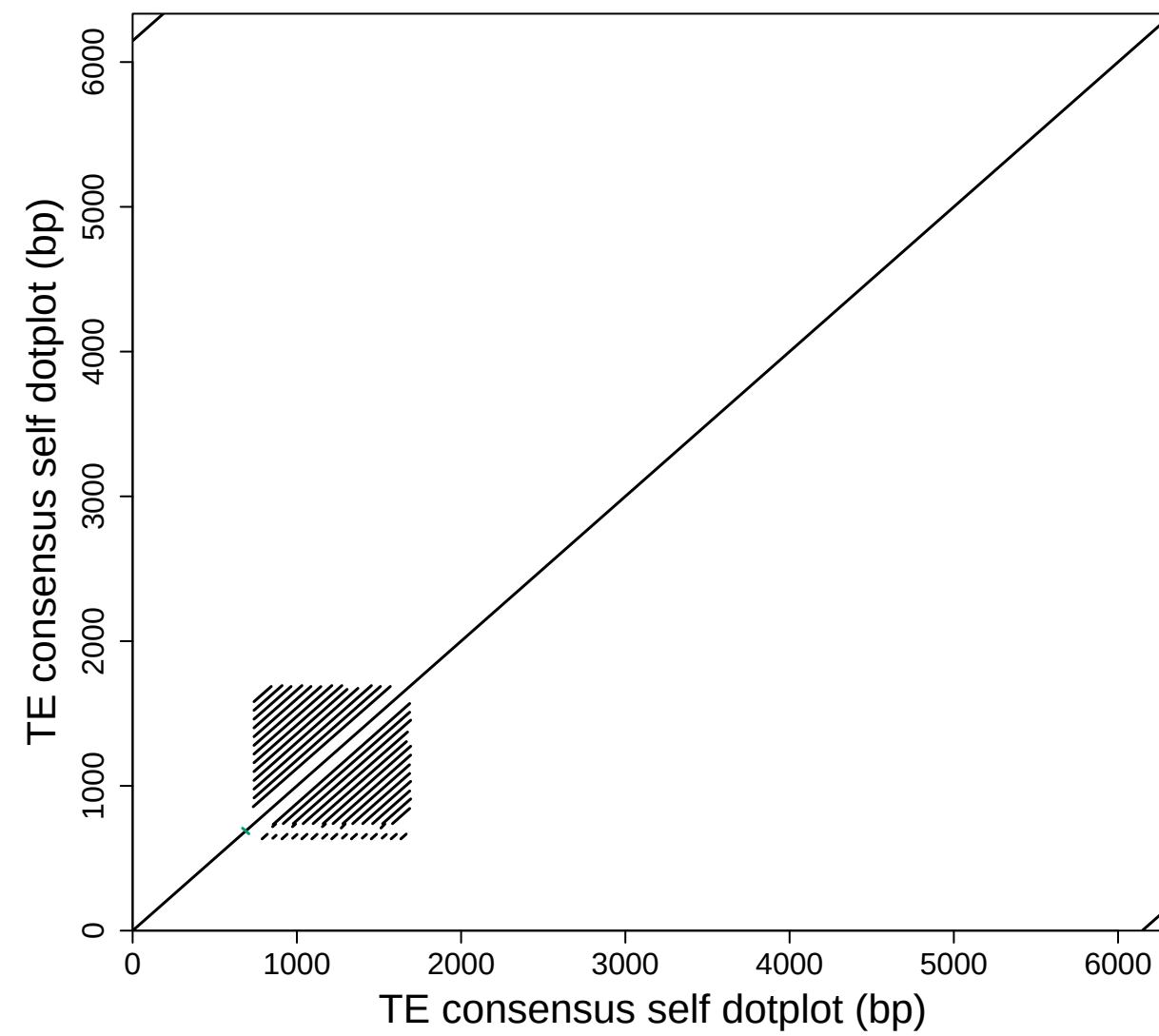
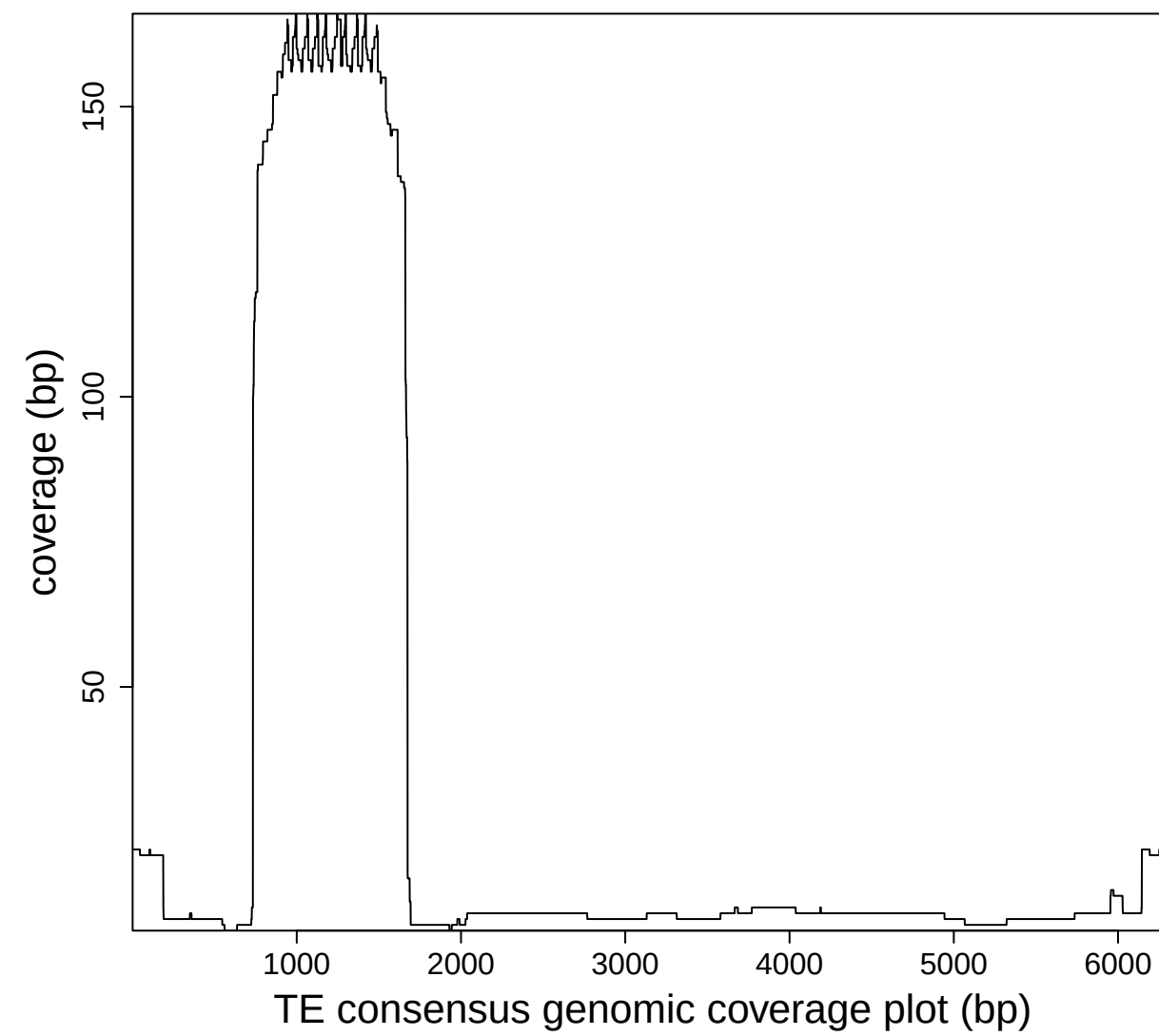
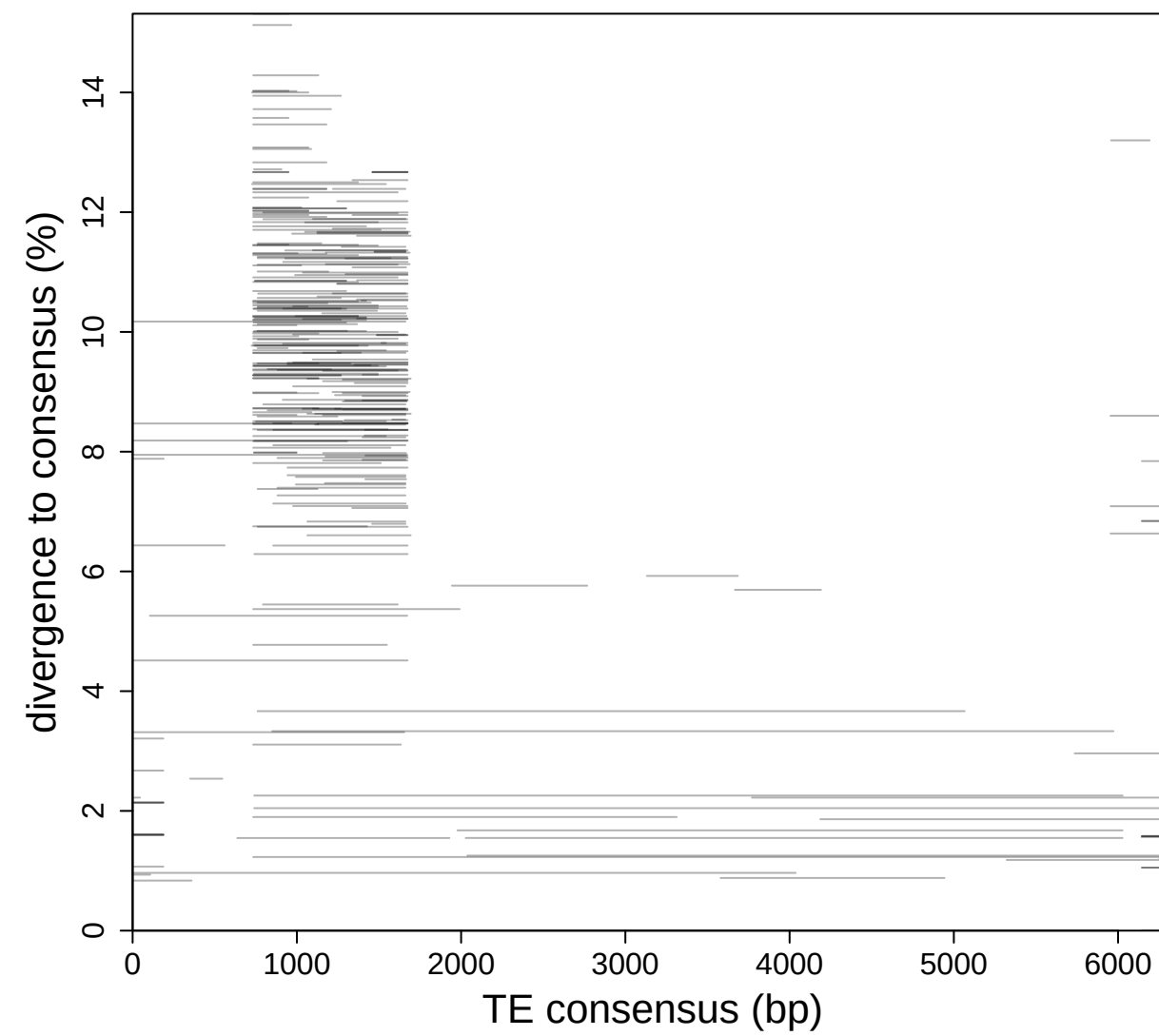
Start crop Point

End crop Point



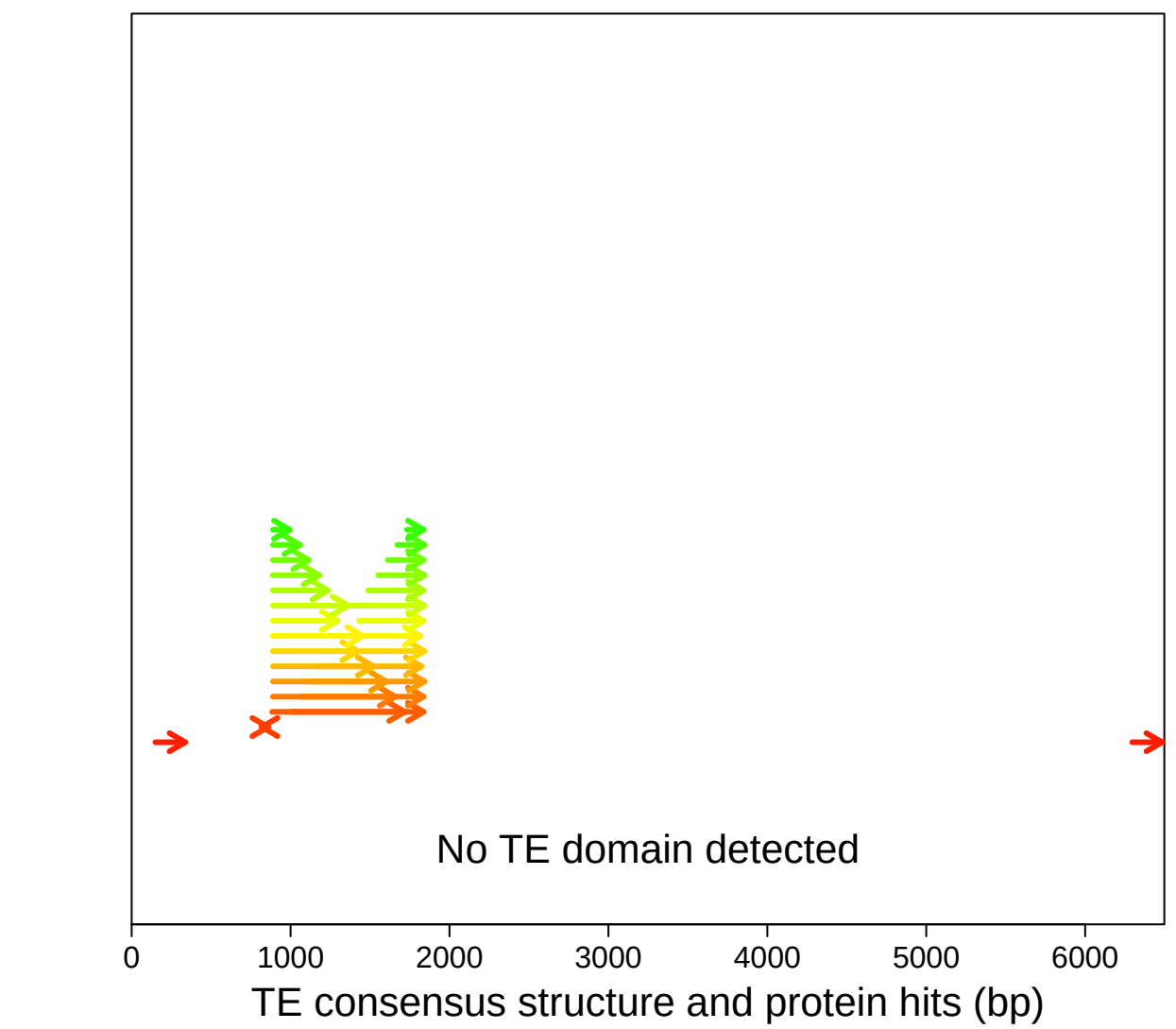
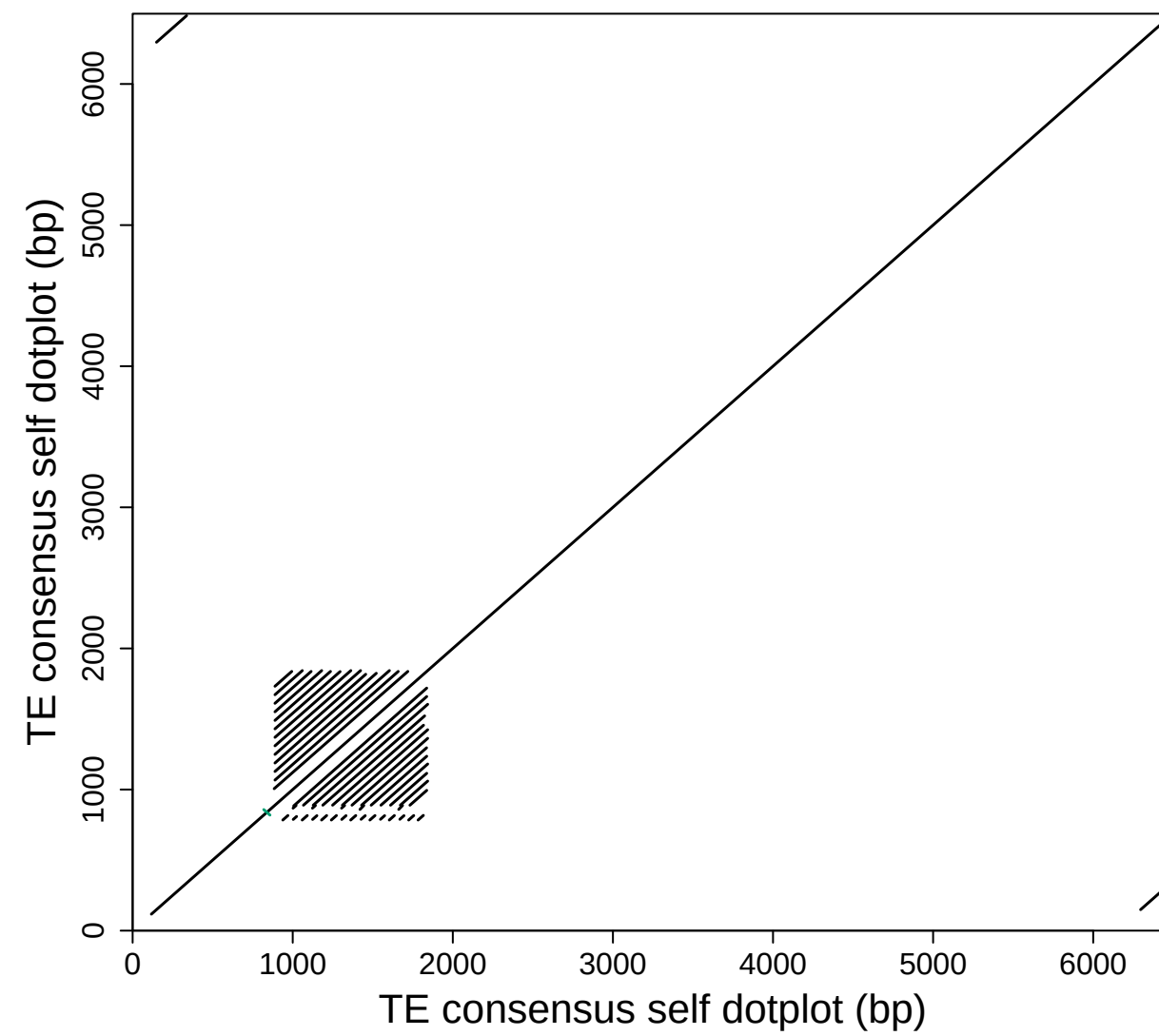
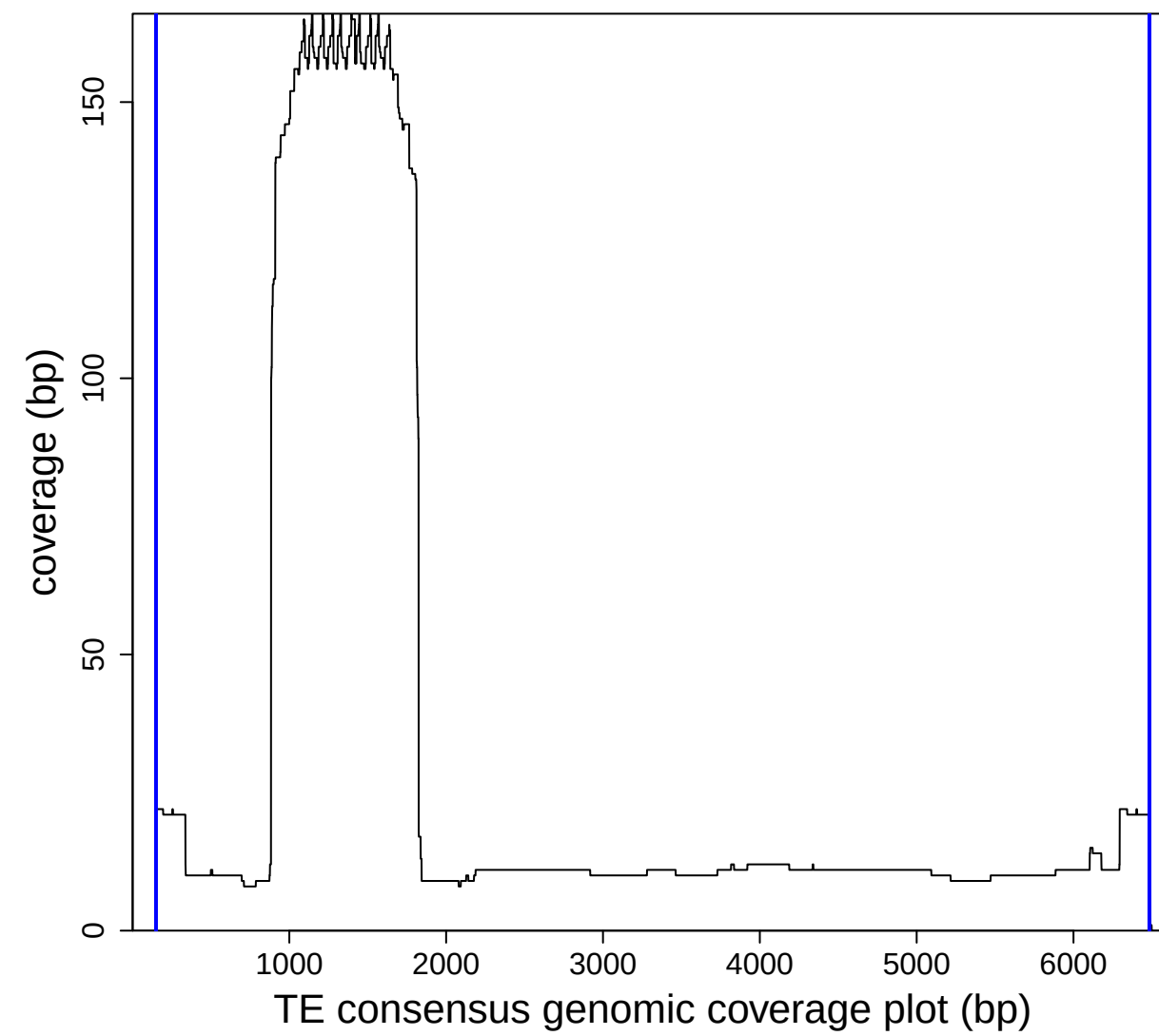
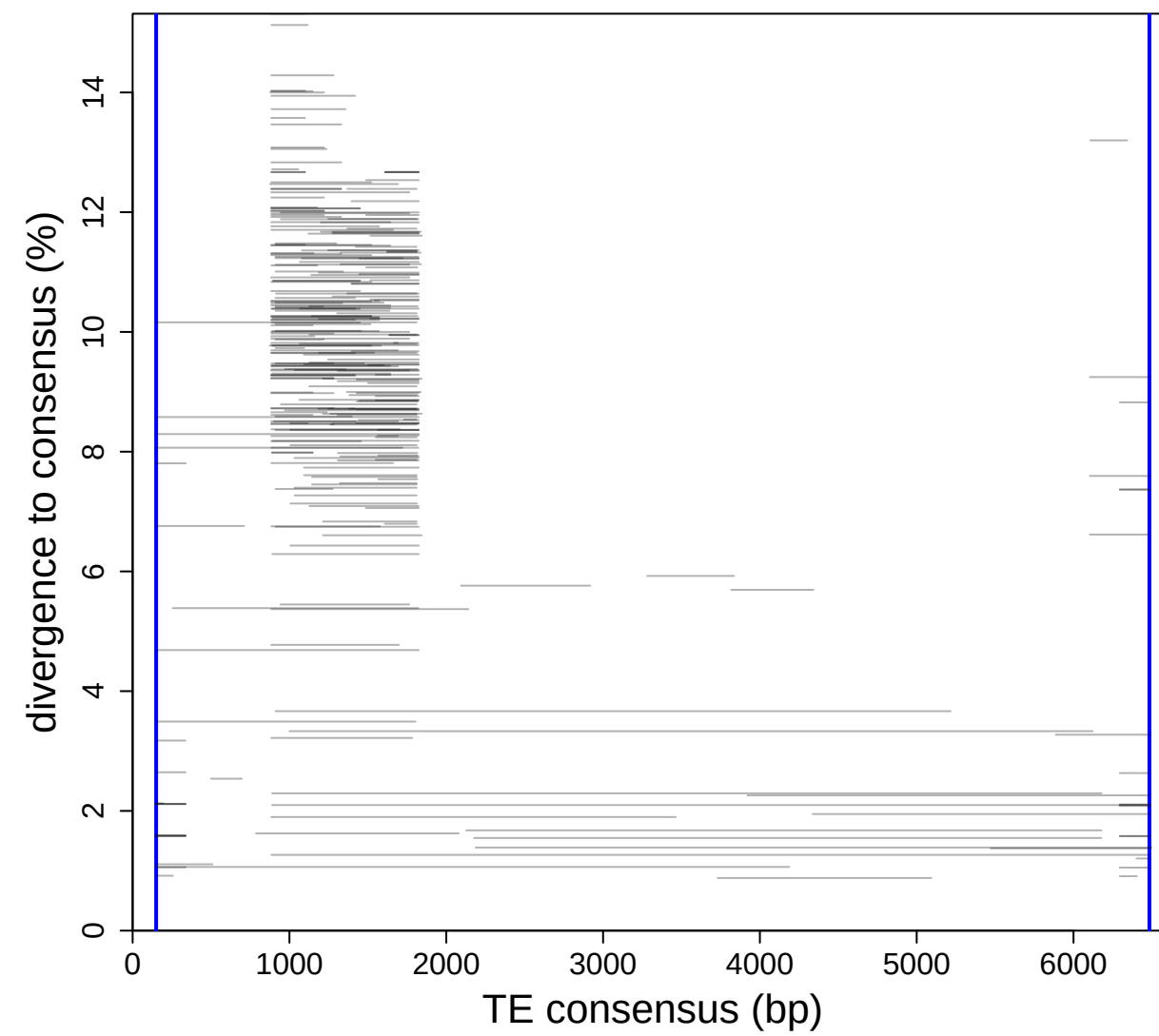
ed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 6334bp; fragments: 299; full length: 0 (>=5700.6bp)

After TEtrimmer 6334 bp

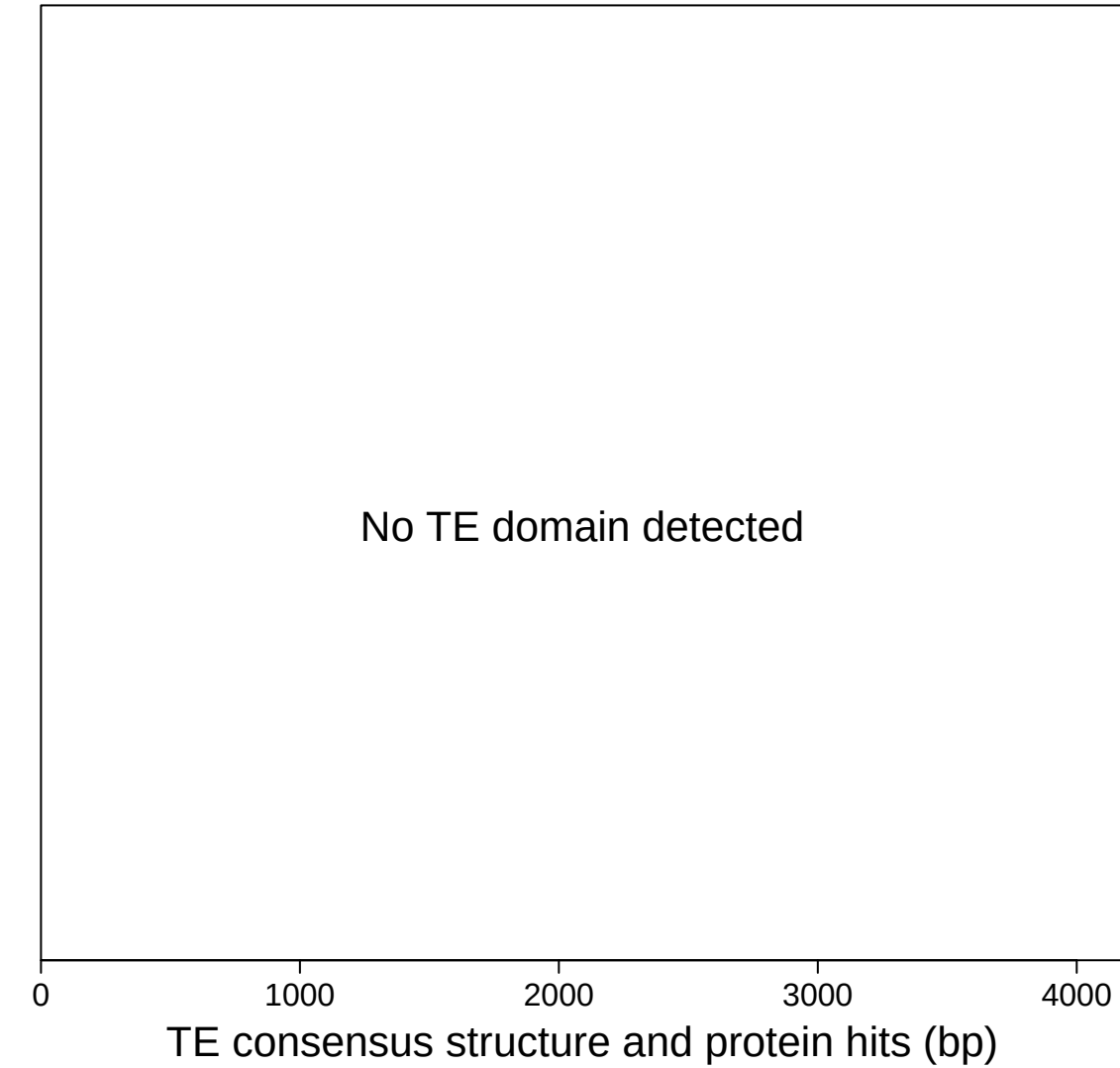
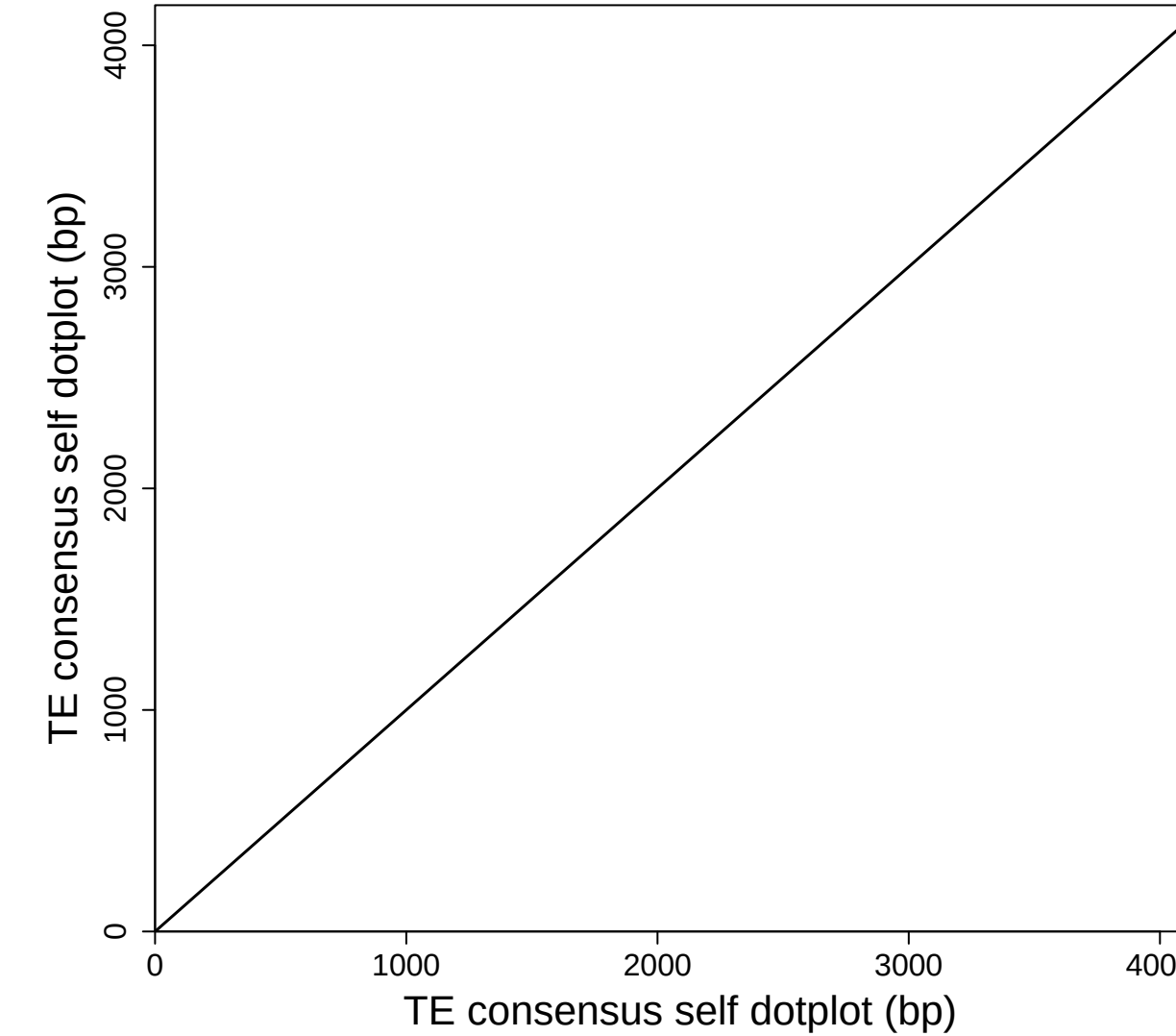
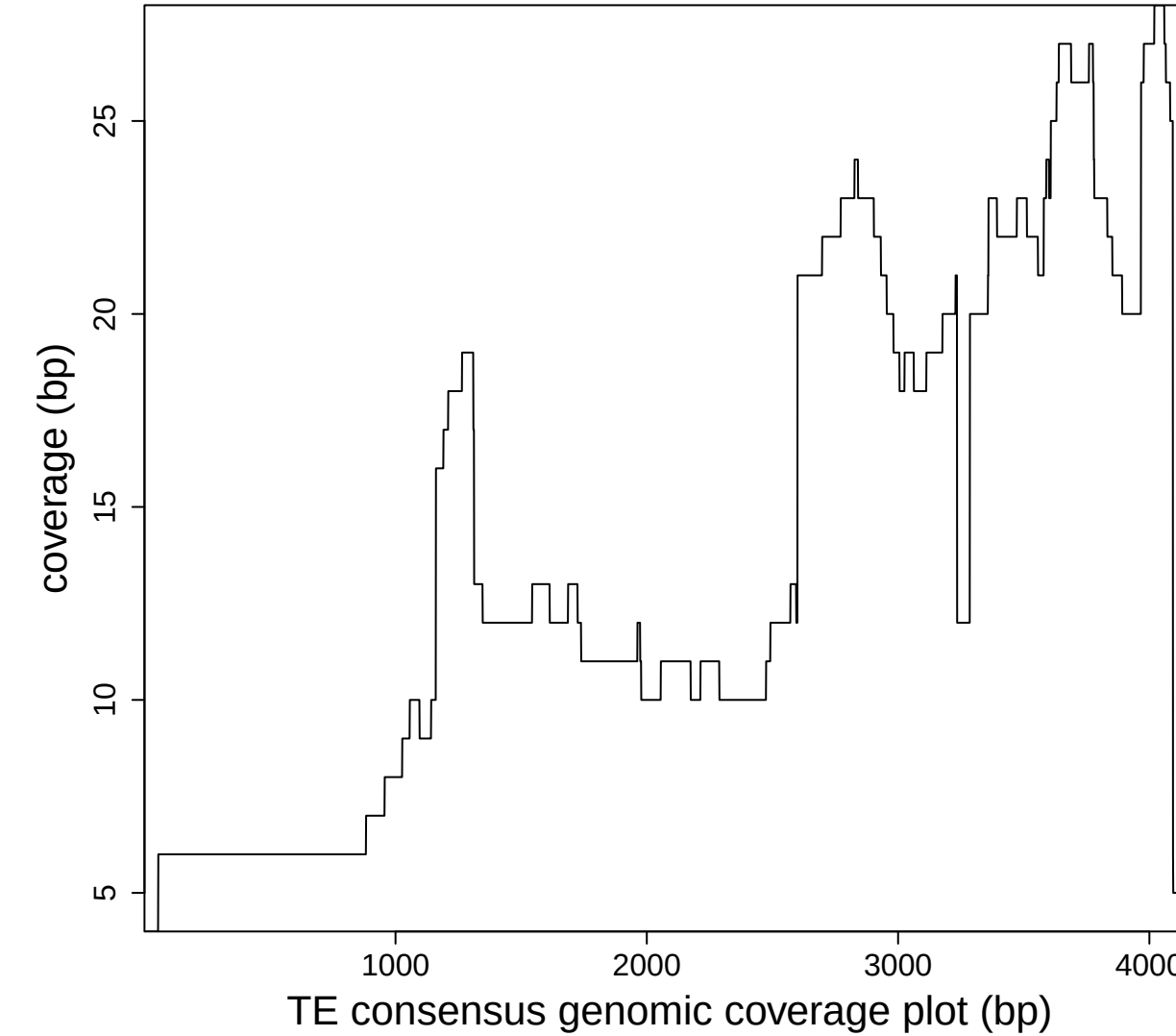
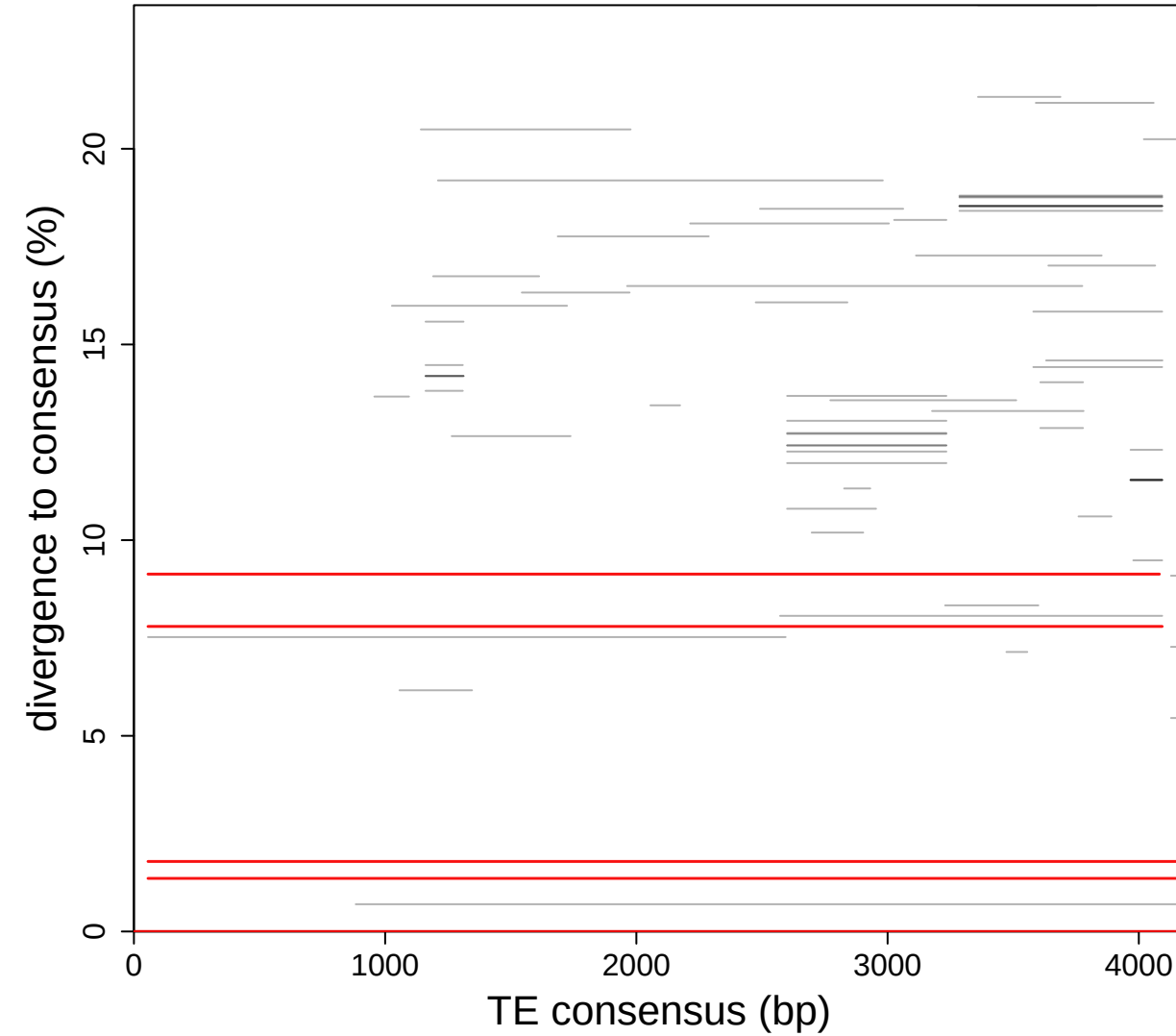


ly_966.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 6634bp; fragments: 299; full length: 0 (>=5970.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

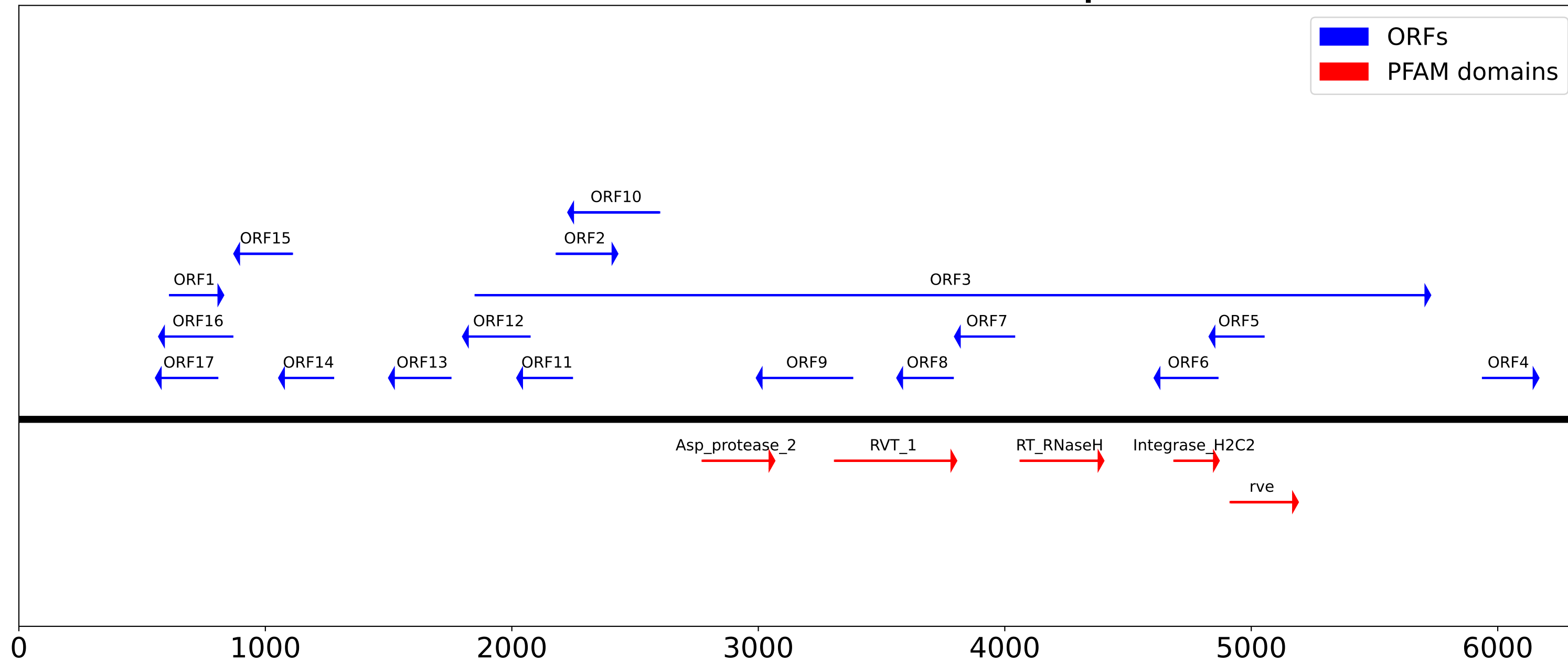


TE: ltr_1_family_966
size: 4181bp; fragments: 78; full length: 5 (≥ 3762.9 bp)

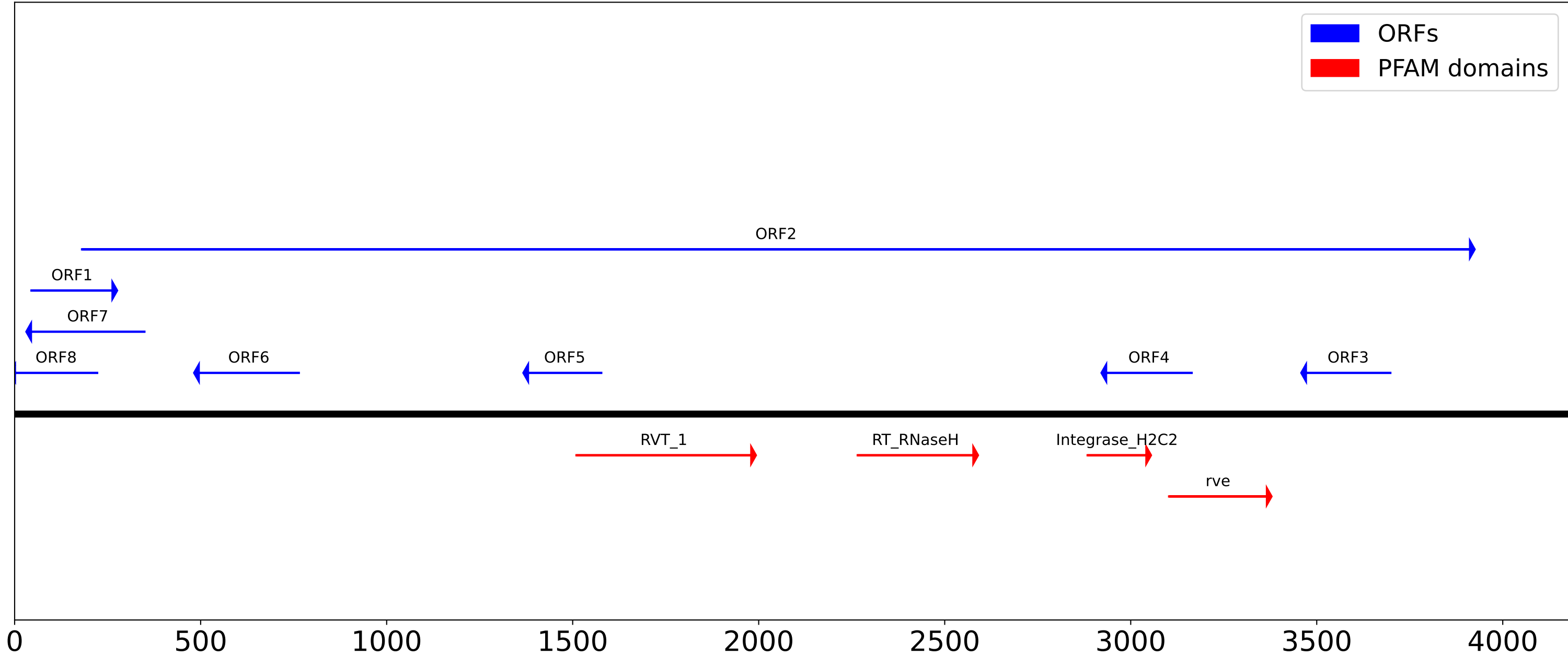


Before TEtrimmer 4181 bp

After TETrimmer ORF and PFAM domain plot



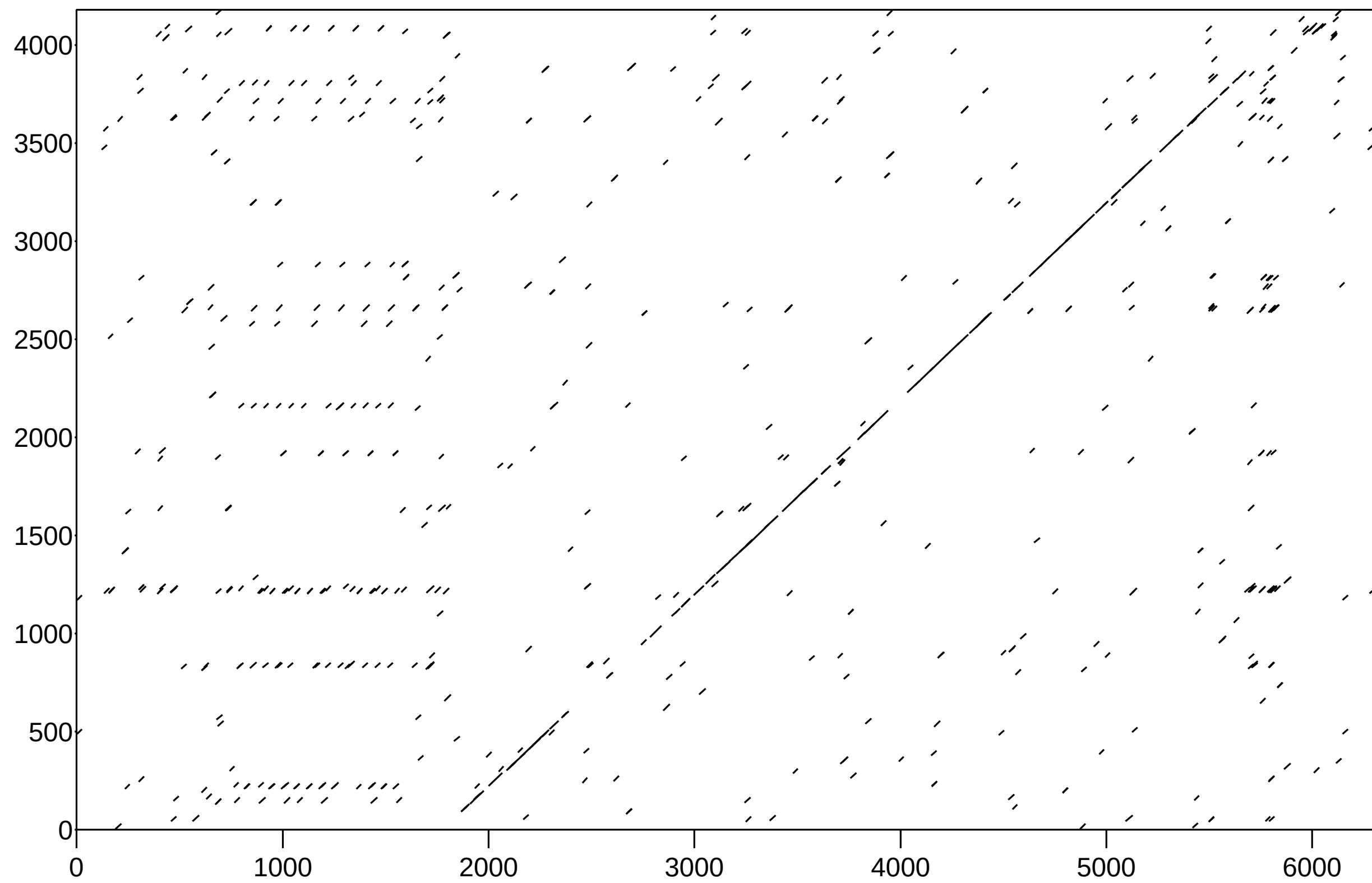
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)