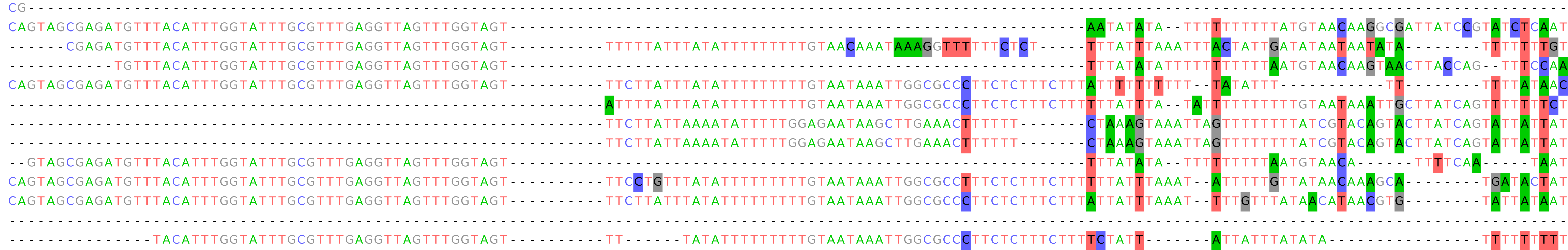
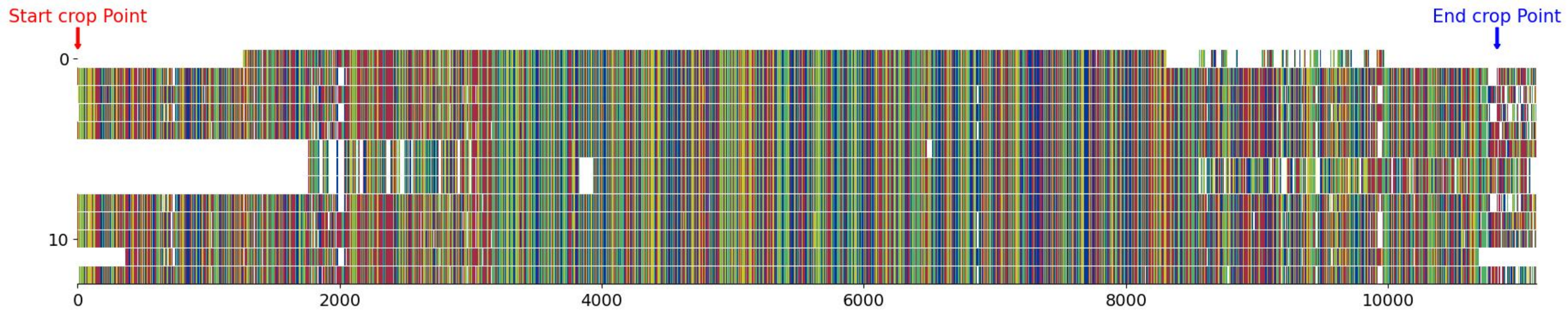


Start crop Point

End crop Point

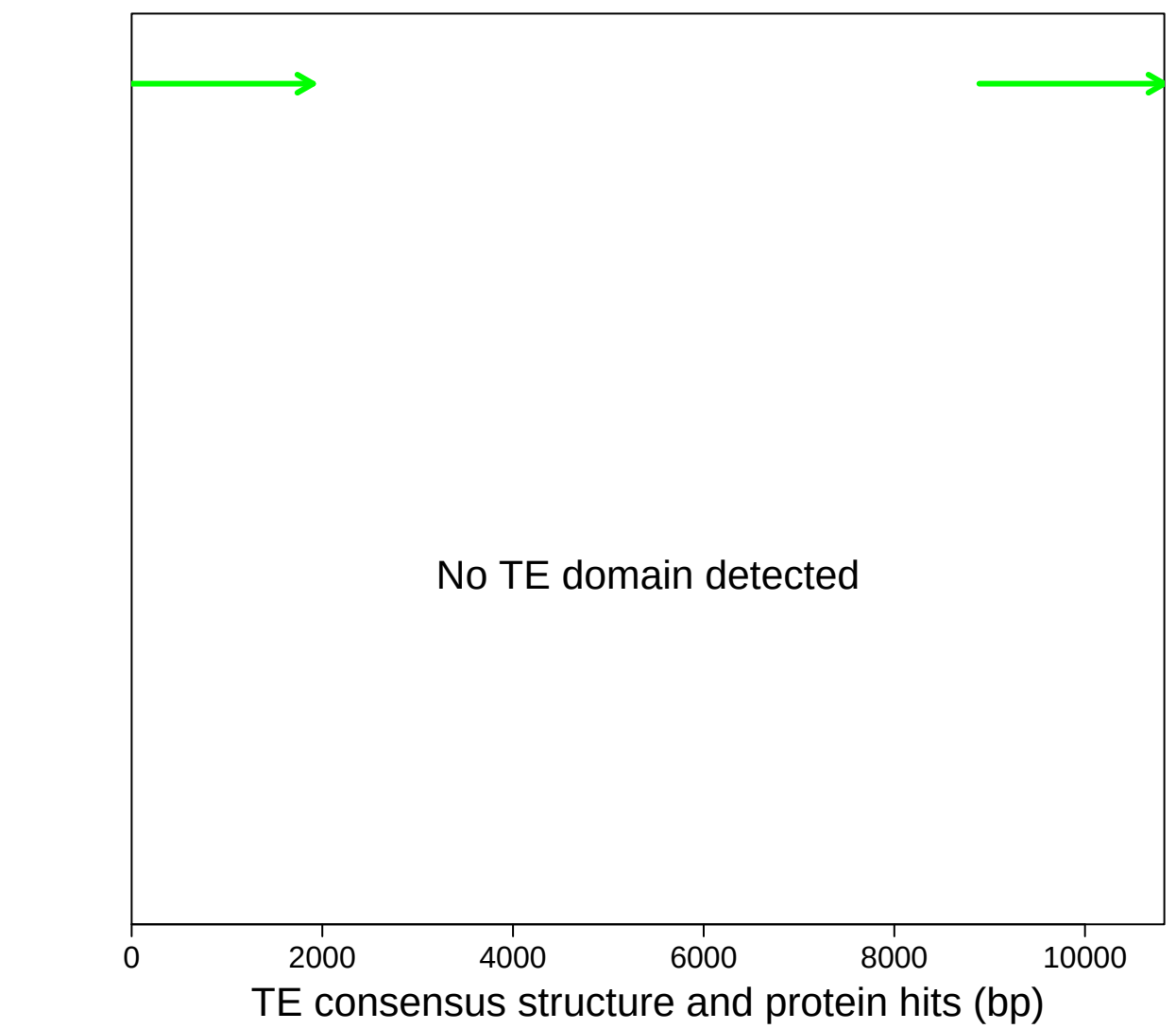
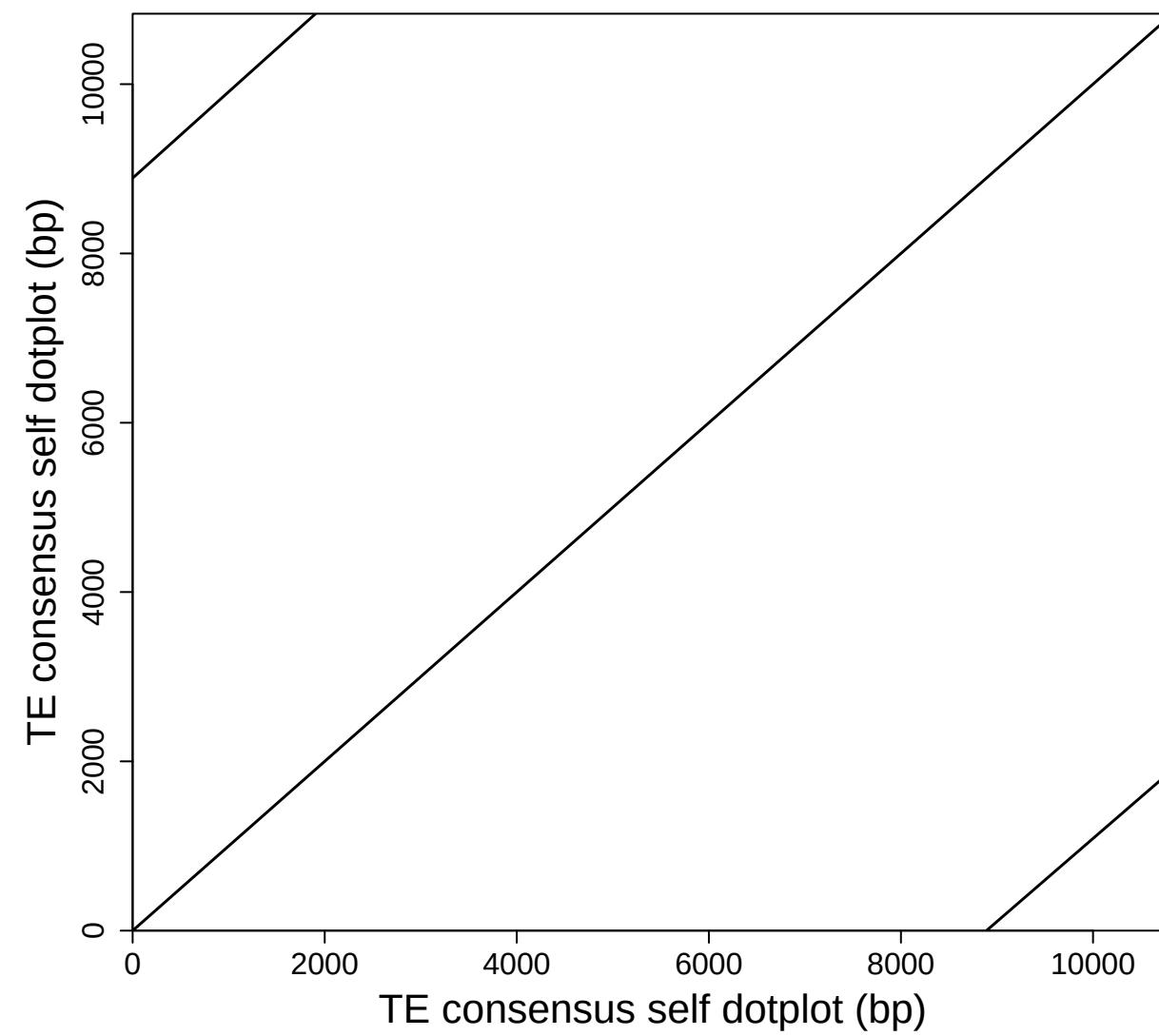
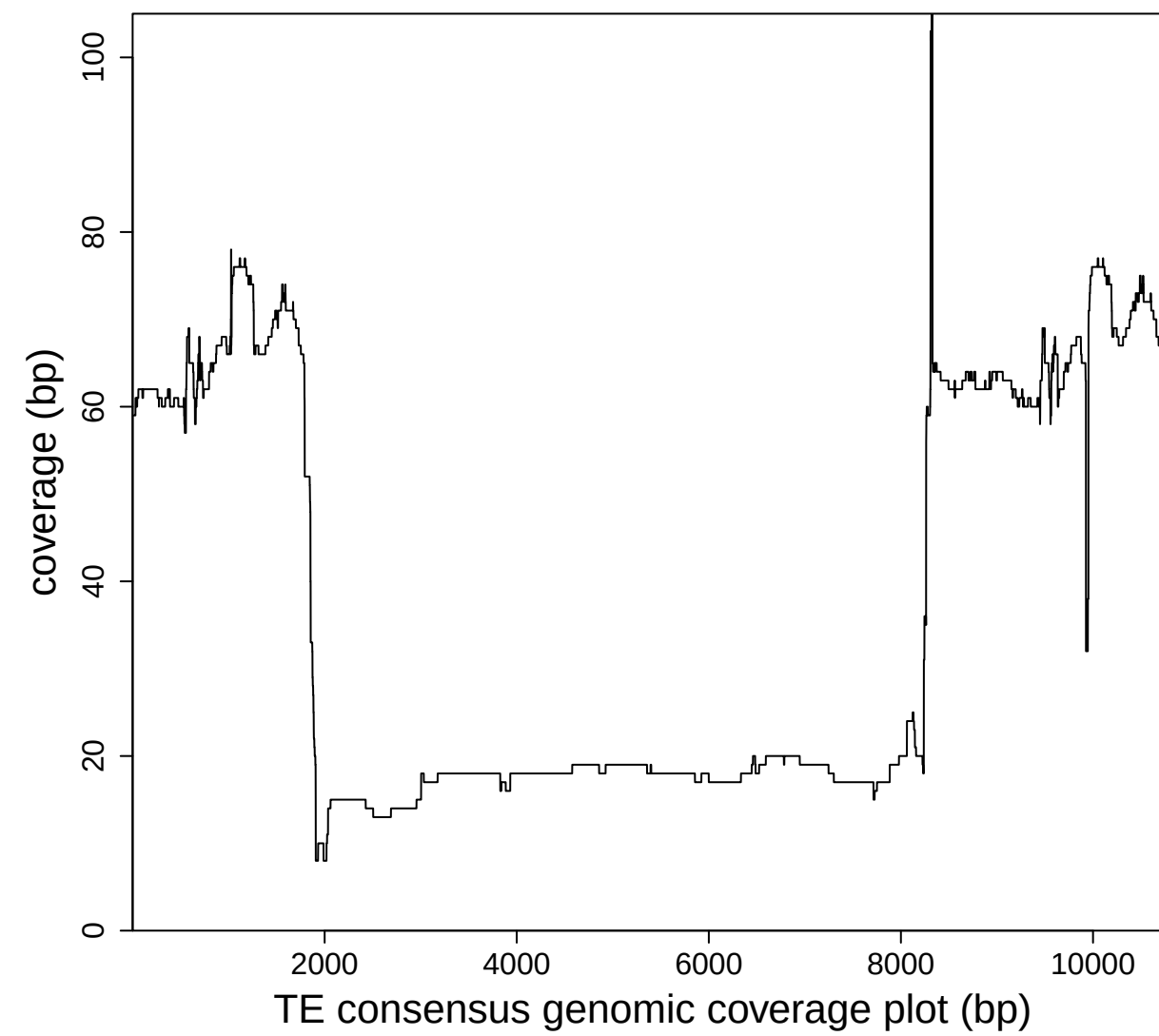
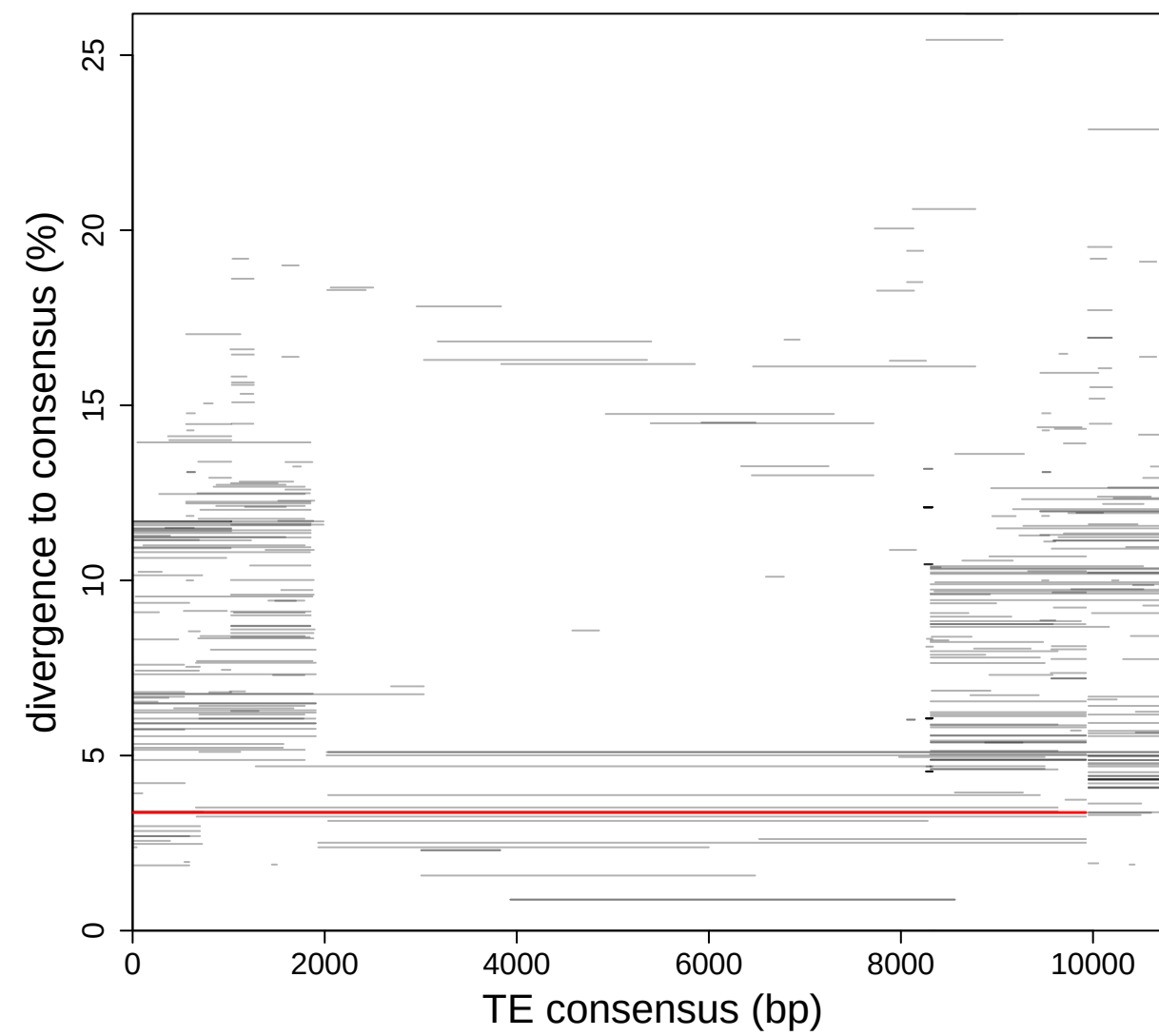
MSA length = 10832





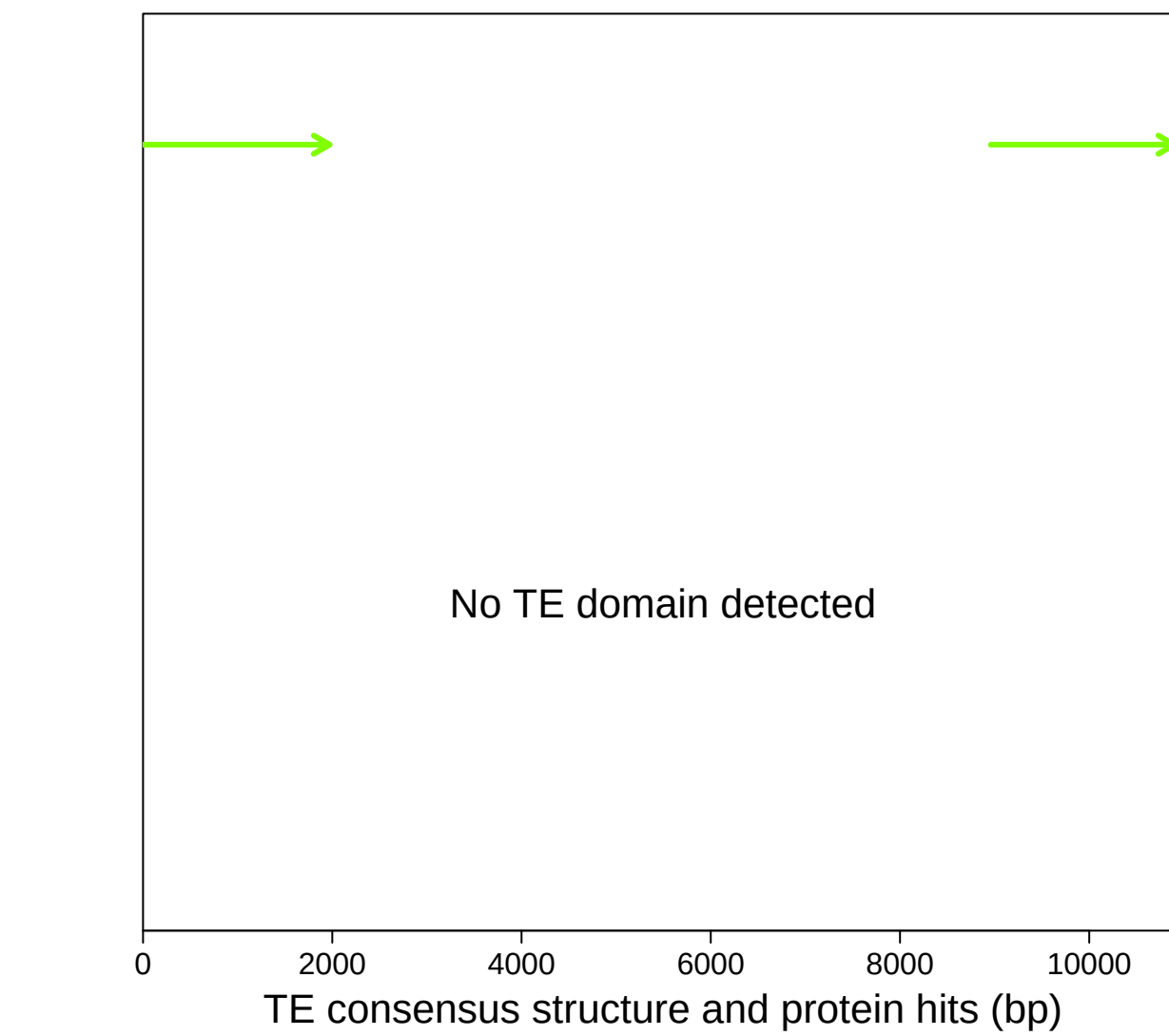
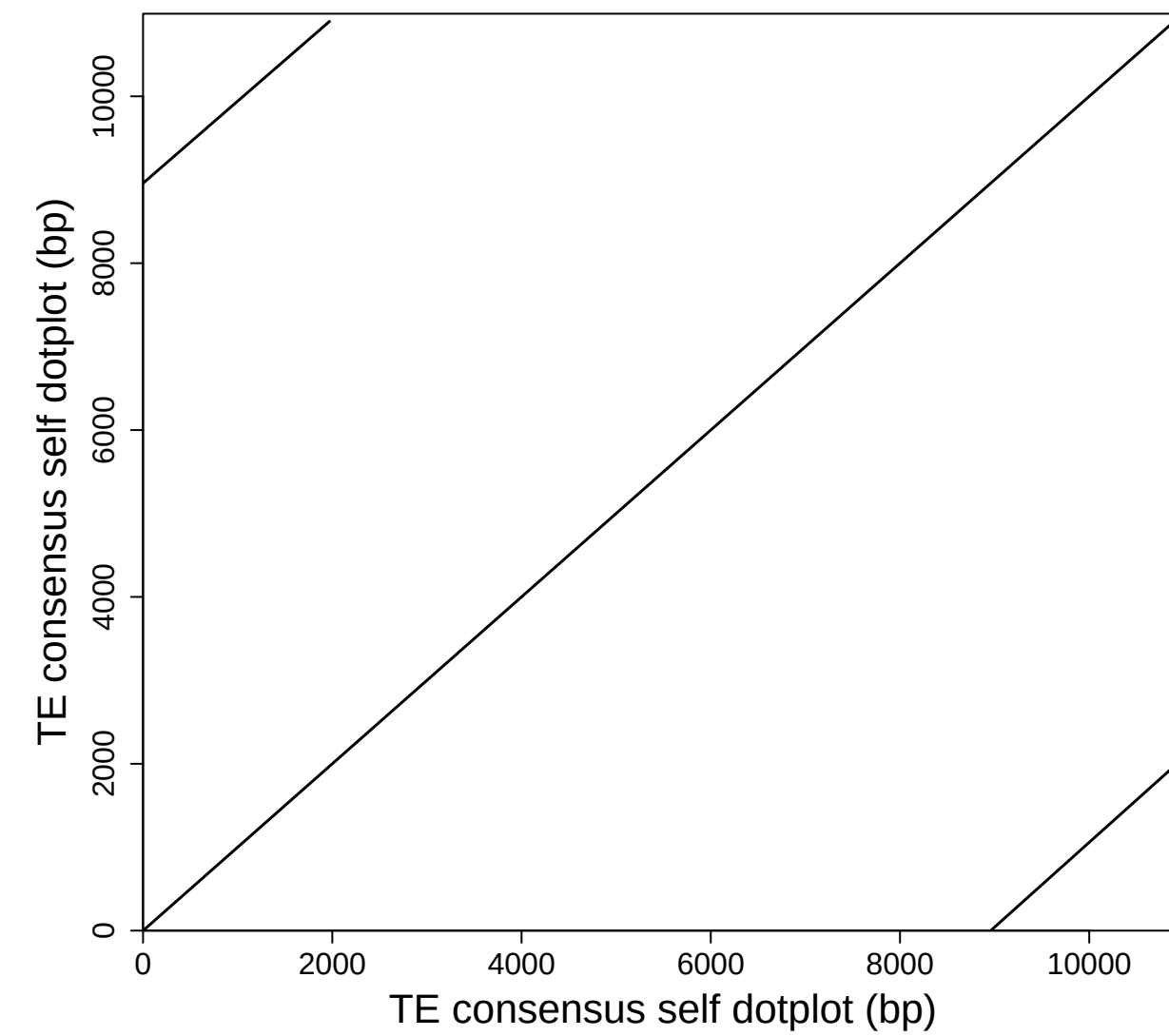
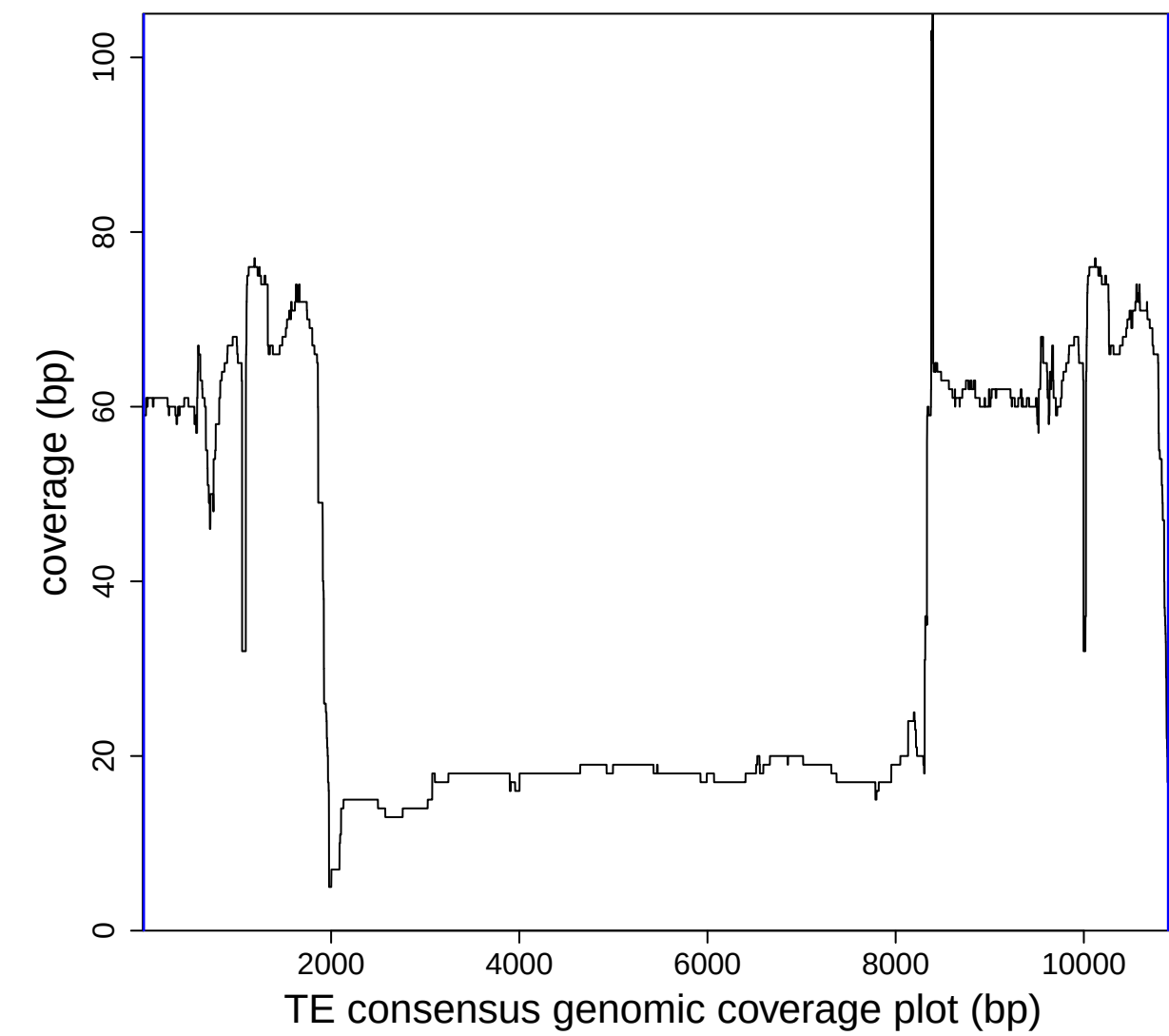
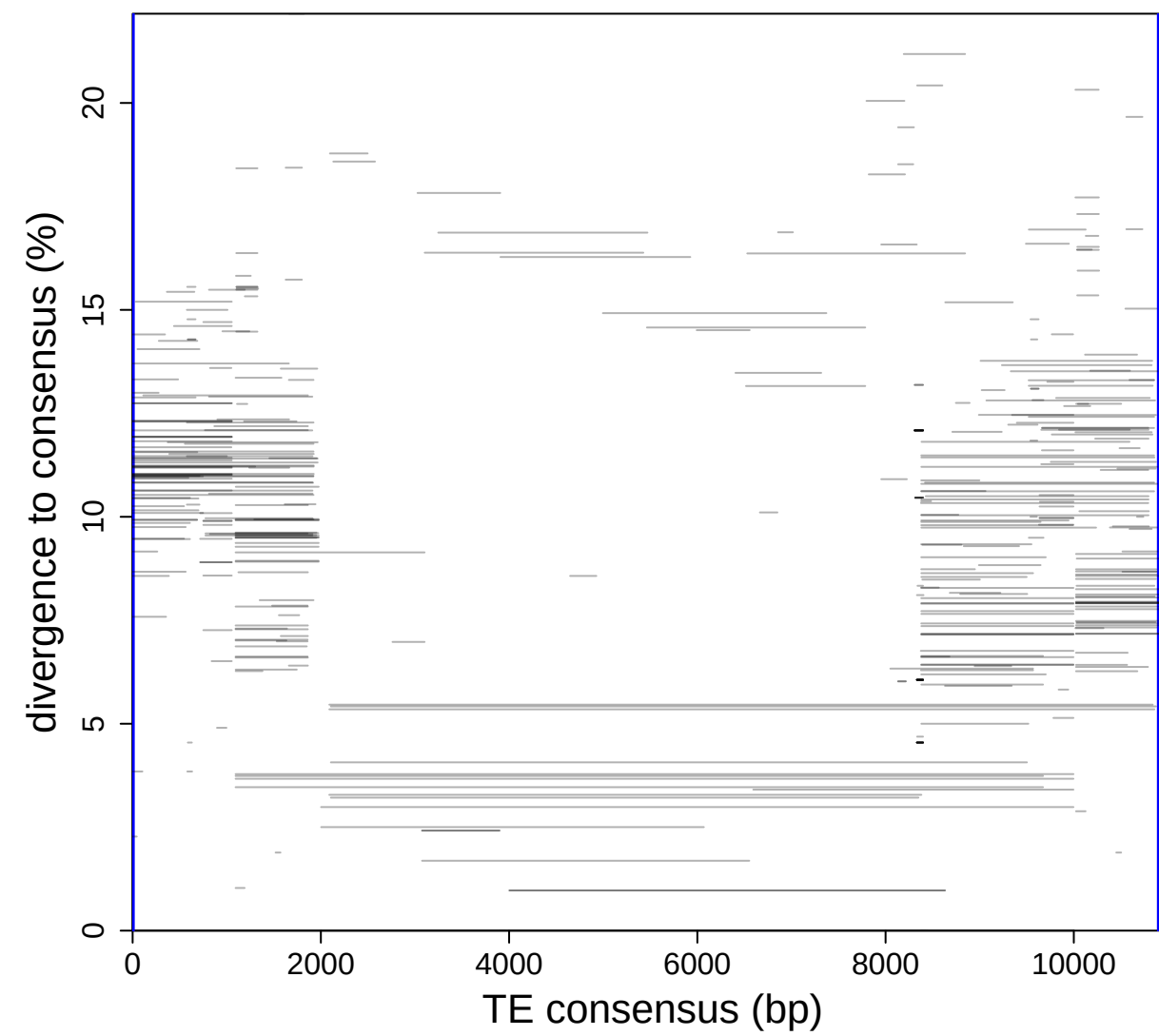
ed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 10832bp; fragments: 394; full length: 1 (>=9748.8bp)

After TEtrimmer 10832 bp



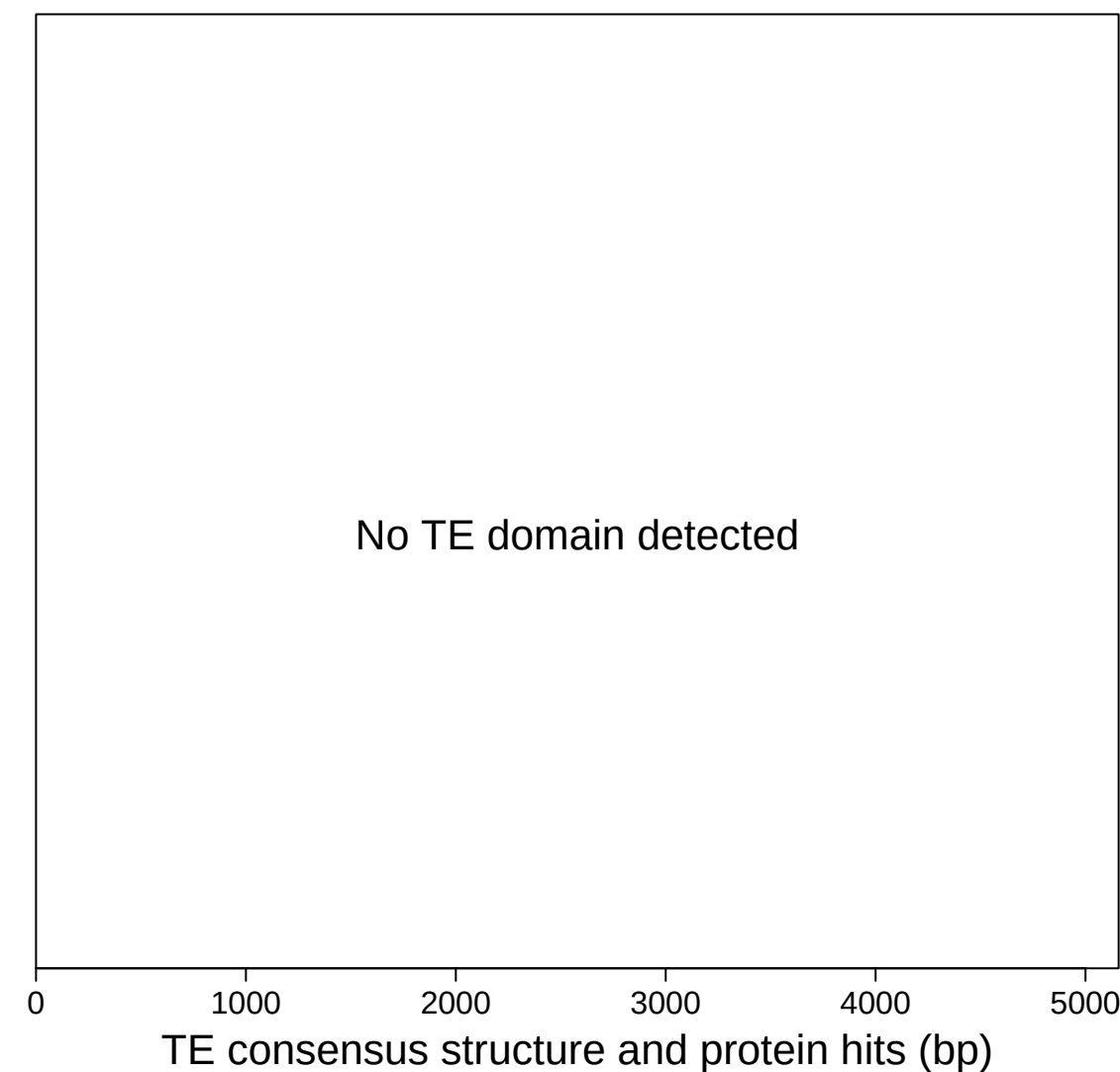
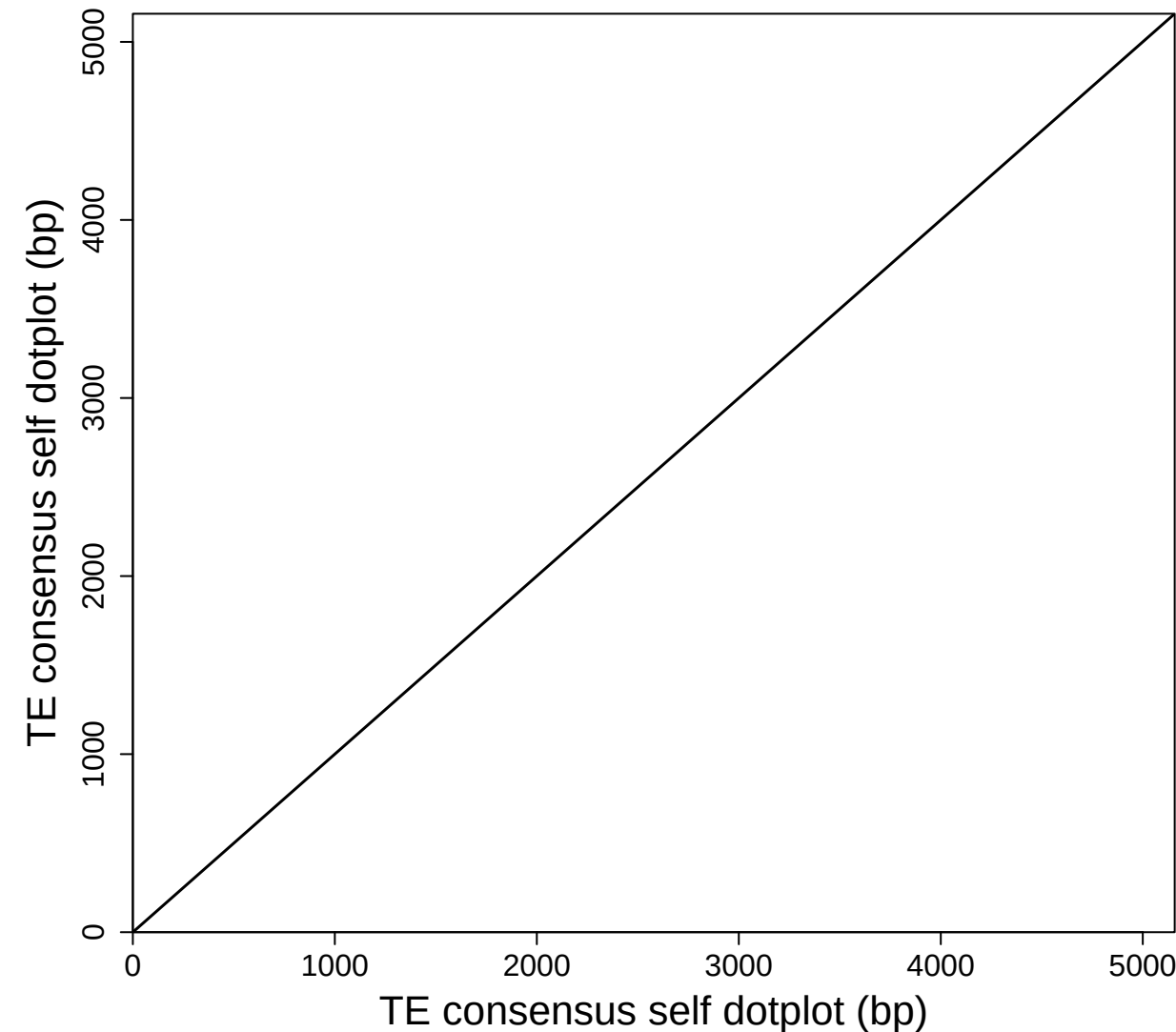
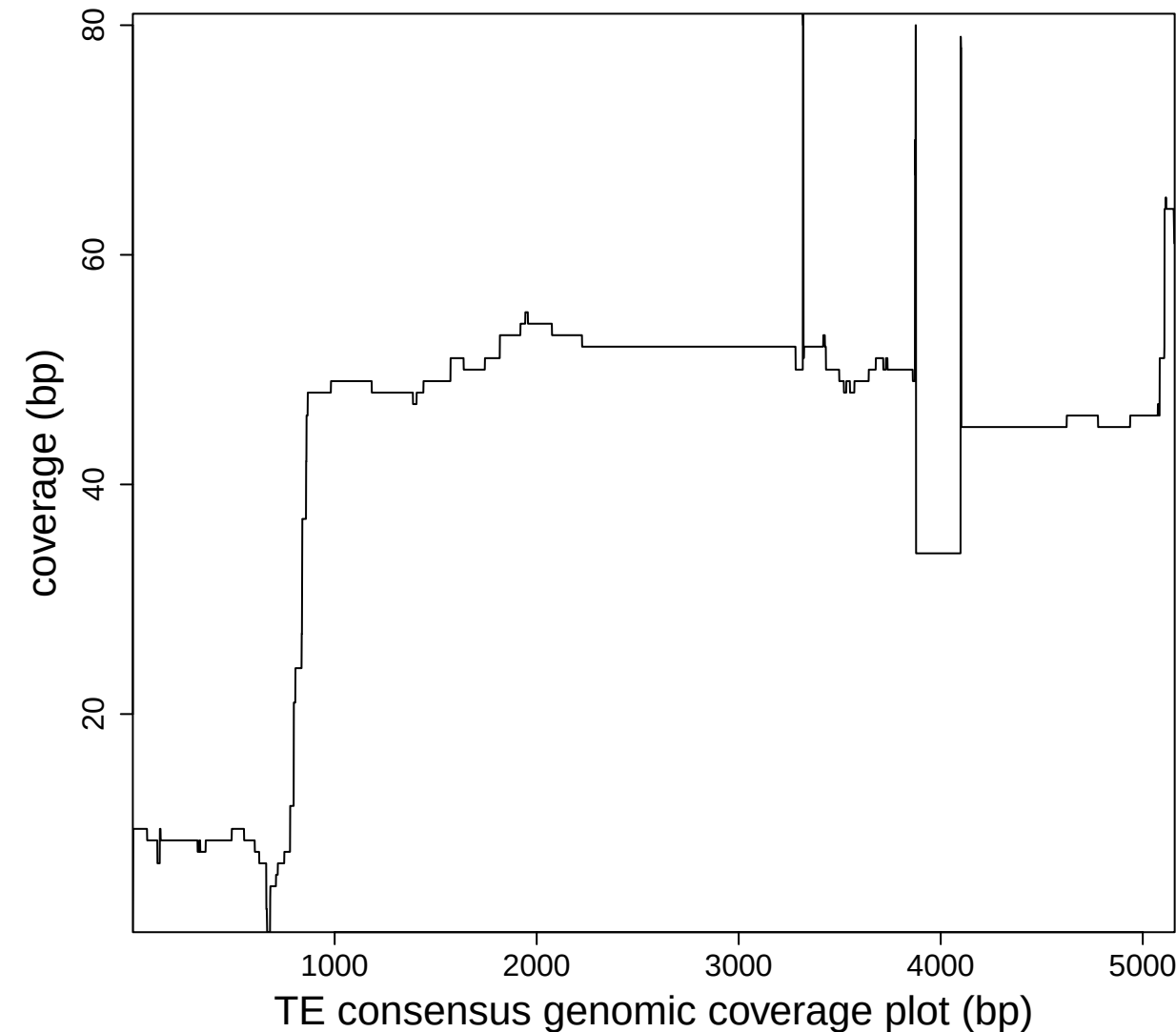
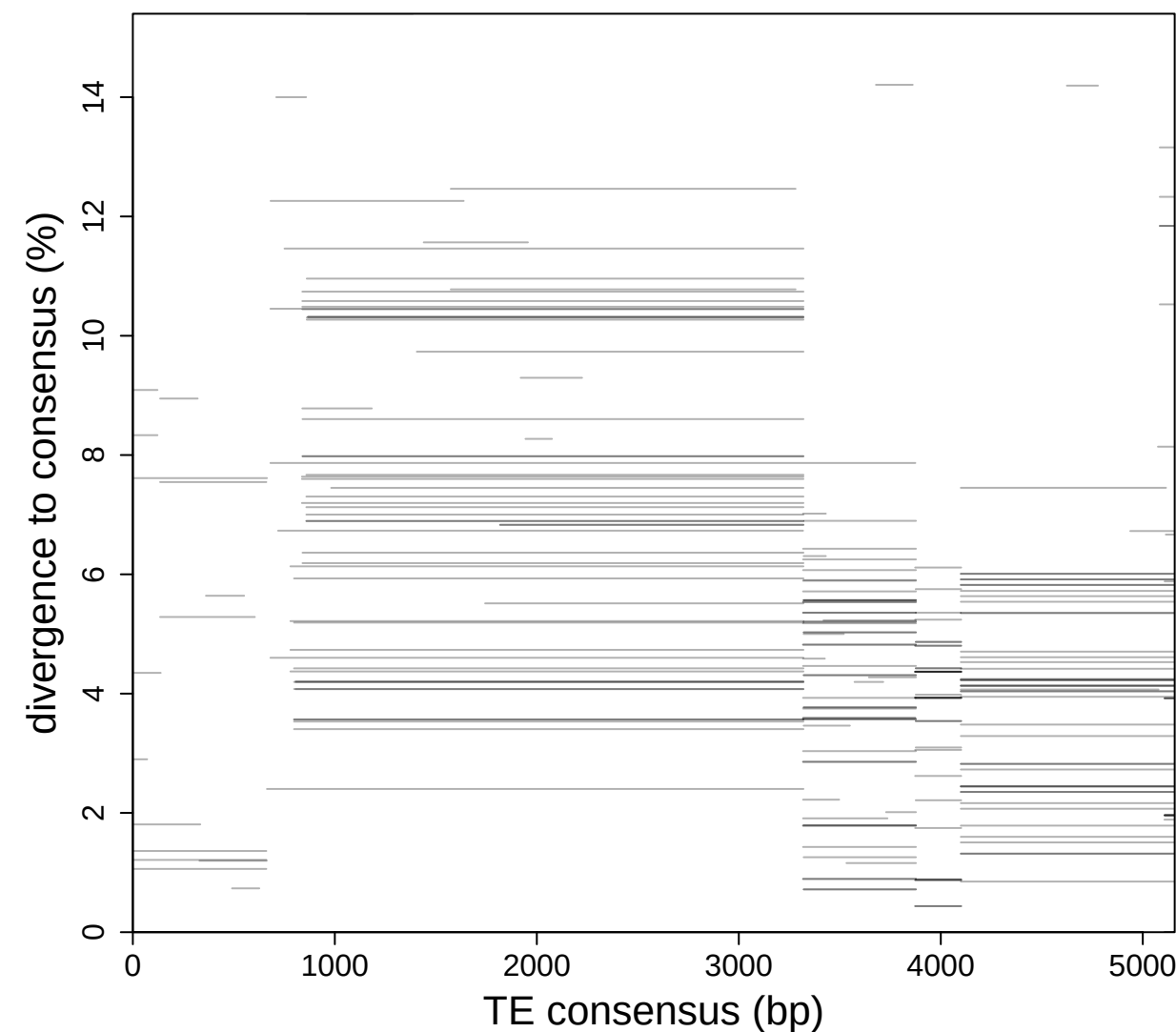
ly_921.fasta.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 11053bp; fragments: 412; full length: 0 (>=9947.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

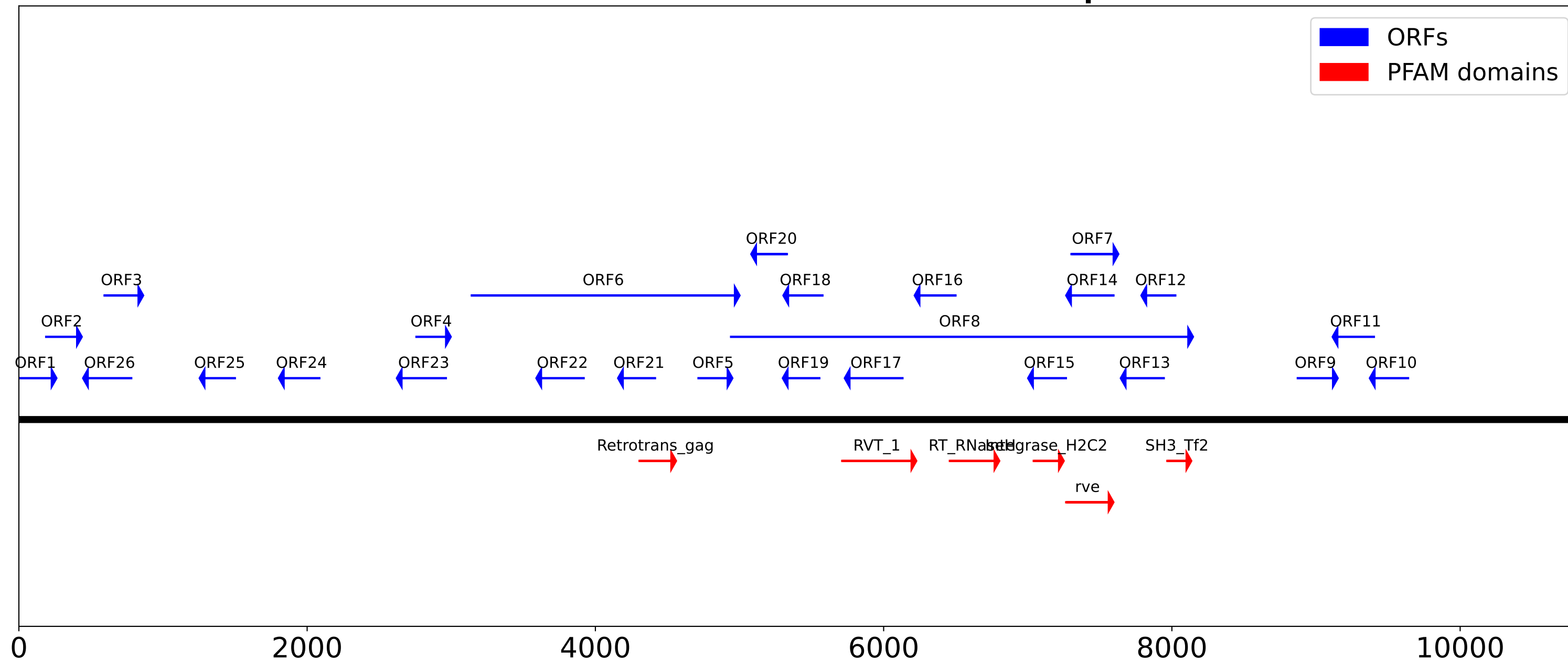


TE: ltr_1_family_921
size: 5158bp; fragments: 233; full length: 0 (>=4642.2bp)

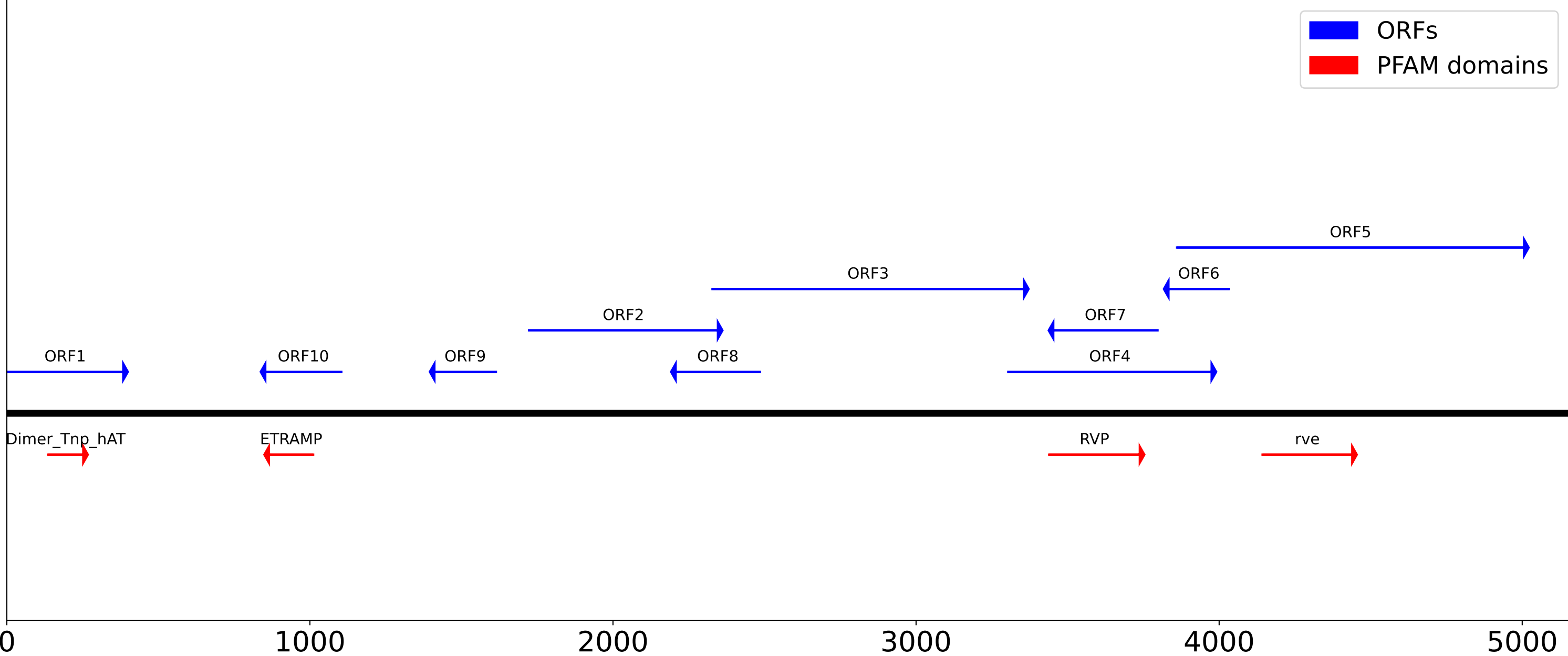
Before TEtrimmer 5158 bp



After TEtrimmer ORF and PFAM domain plot



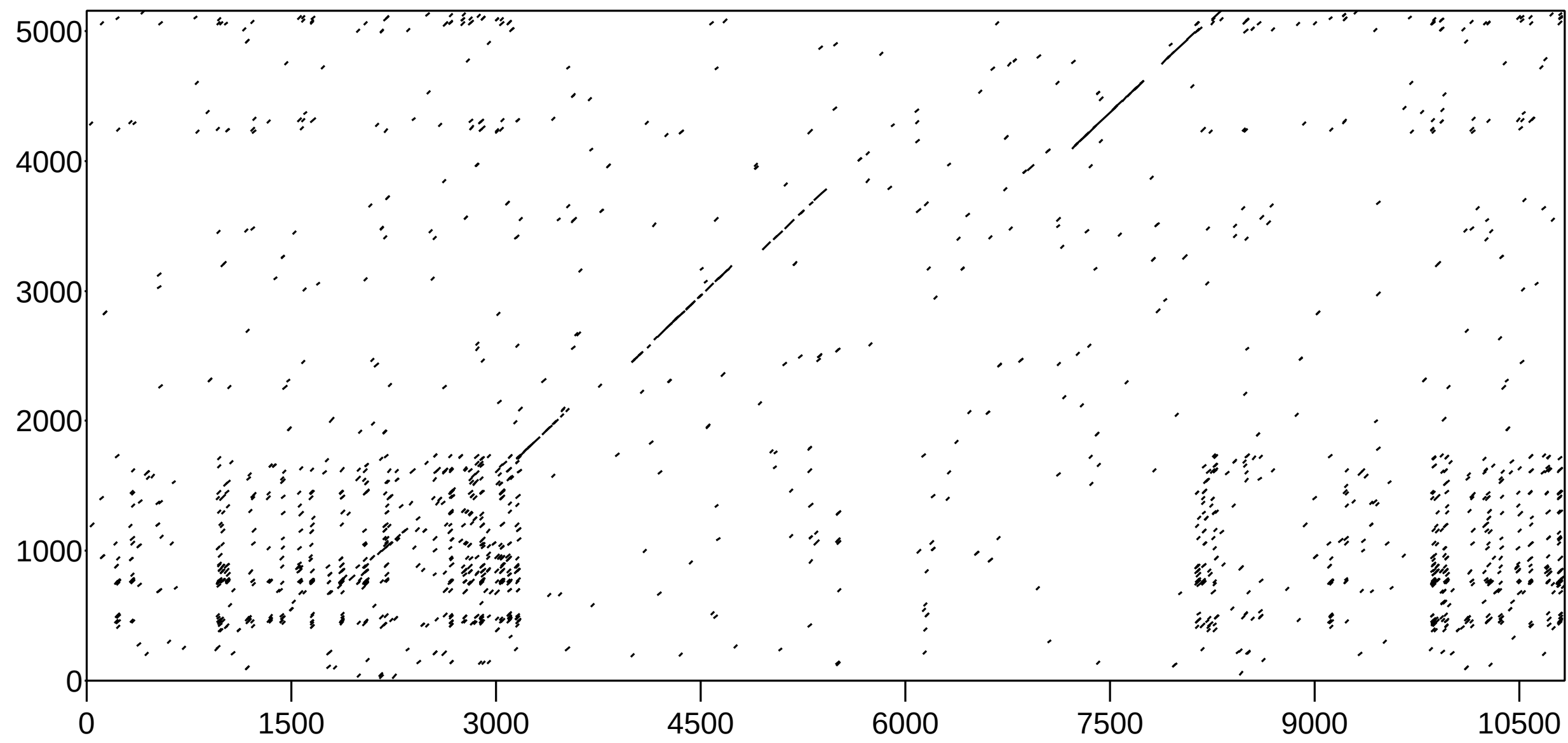
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)