

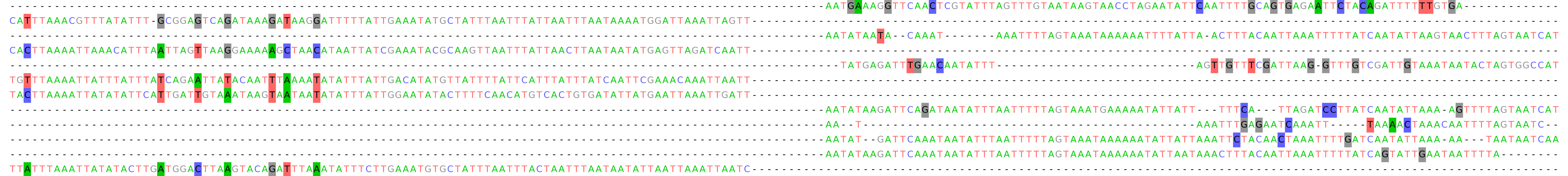
Start crop Point

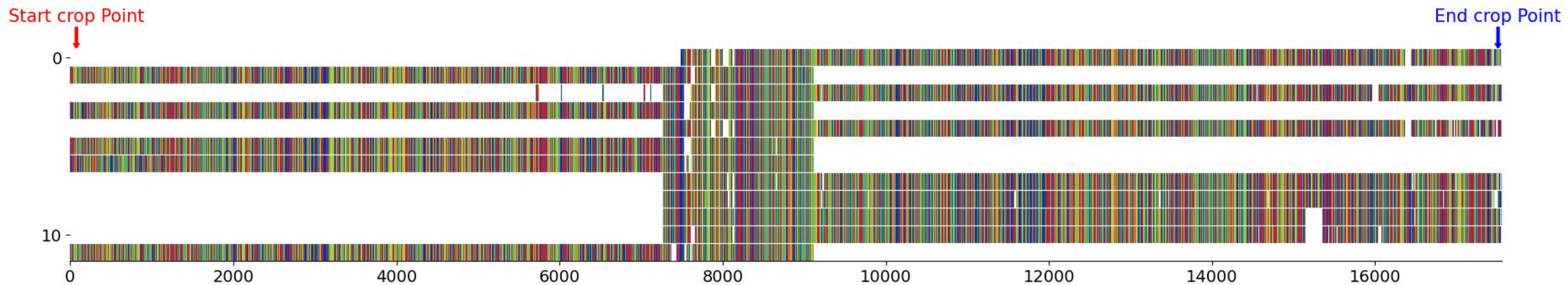


End crop Point



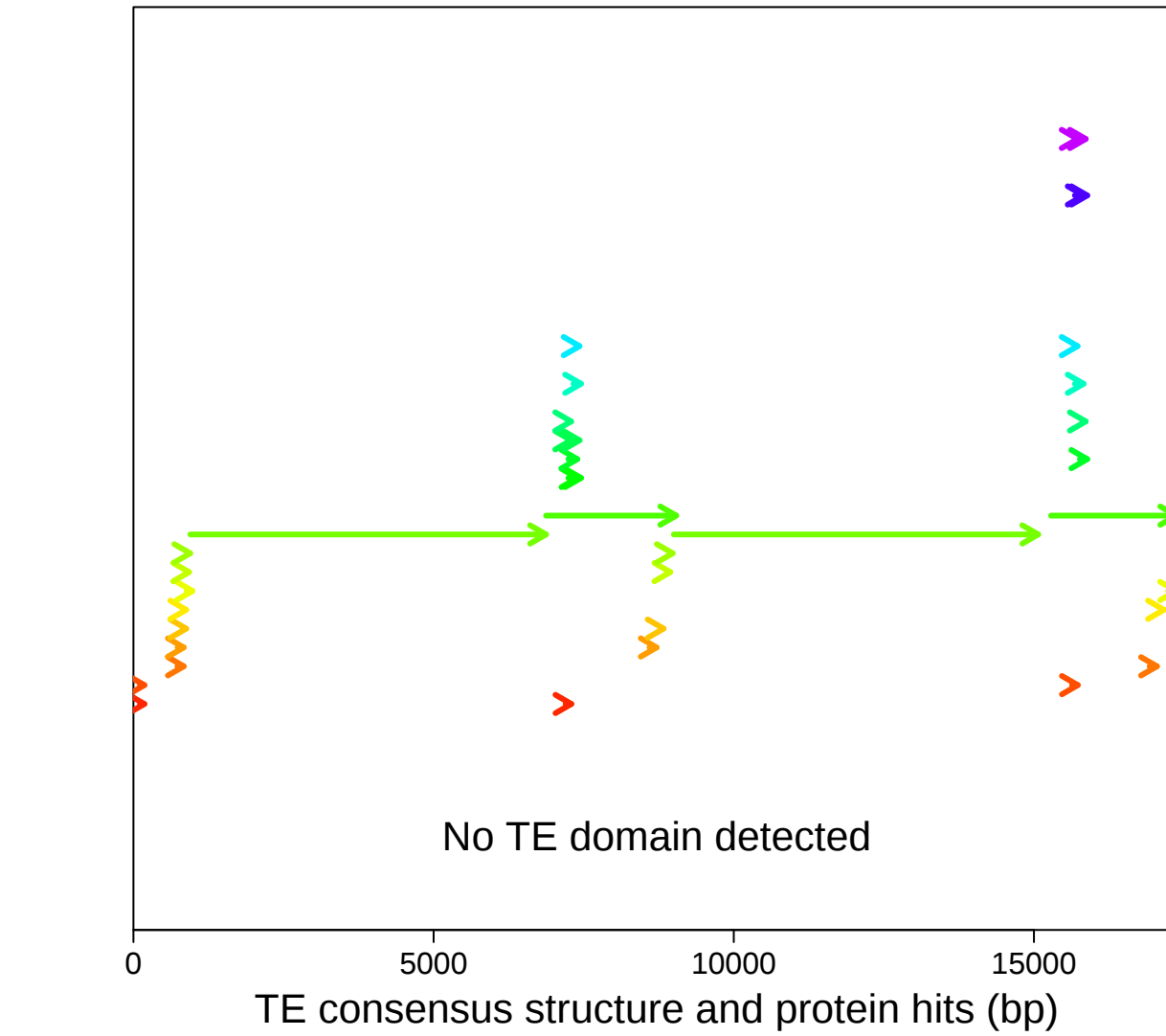
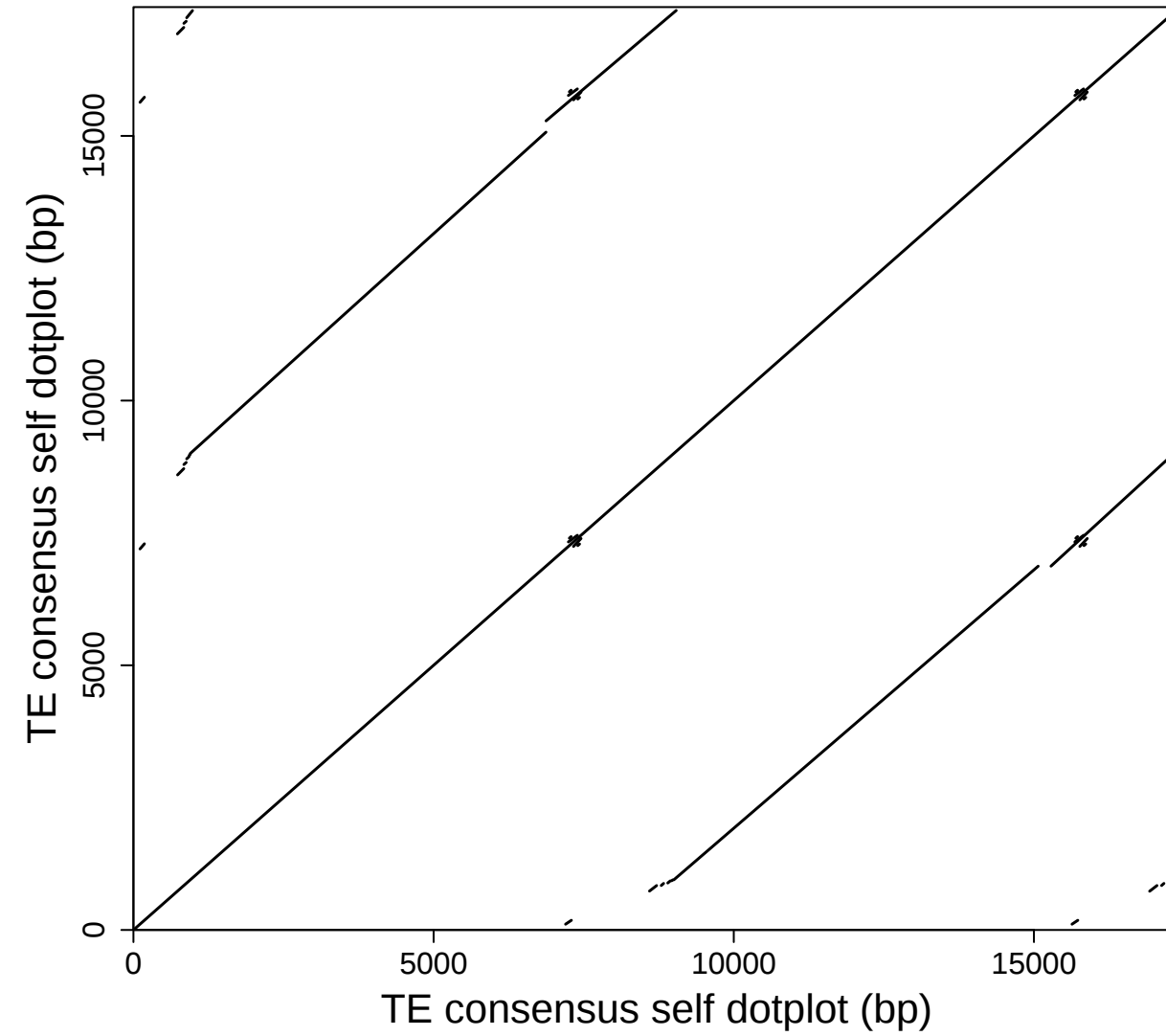
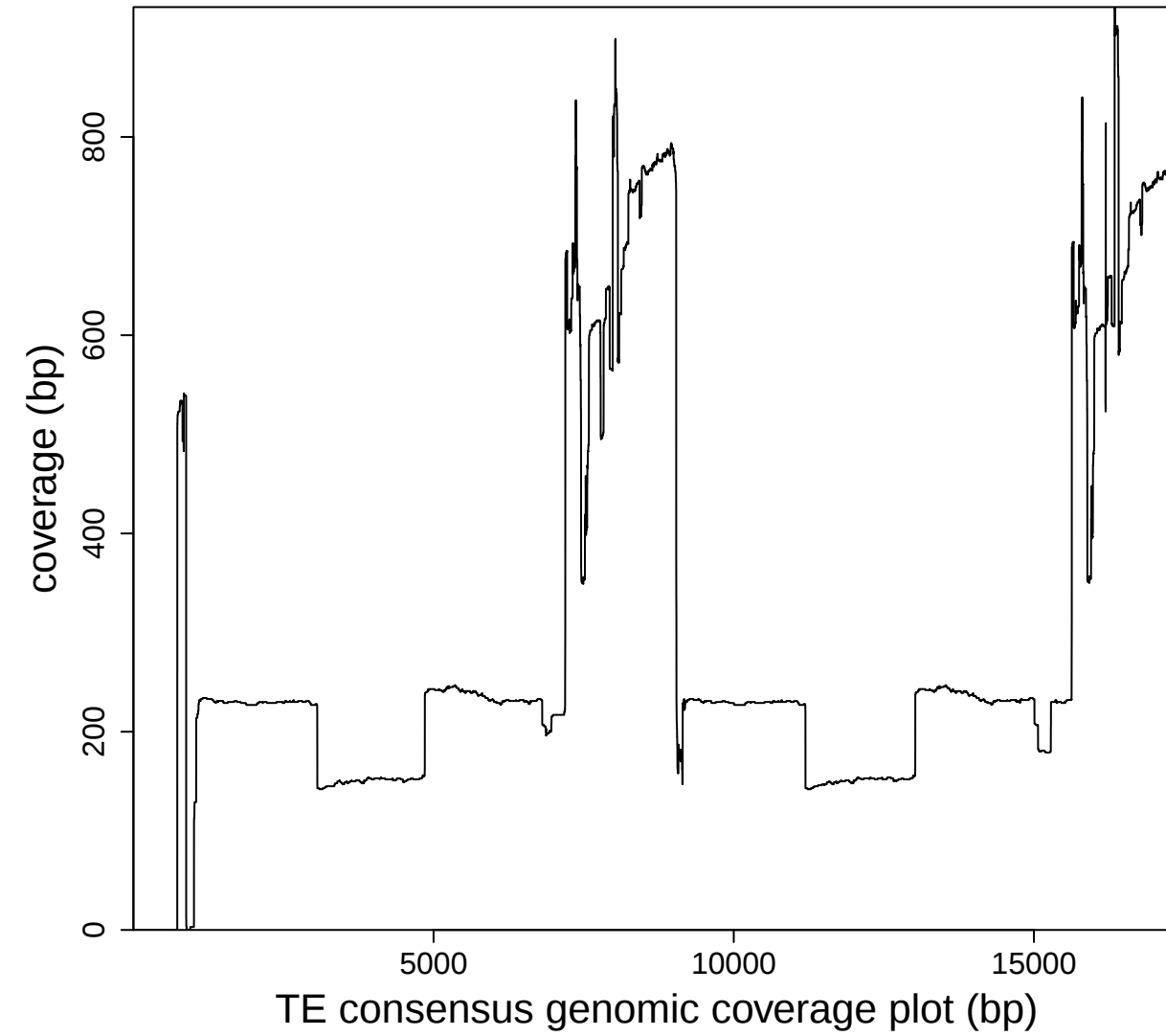
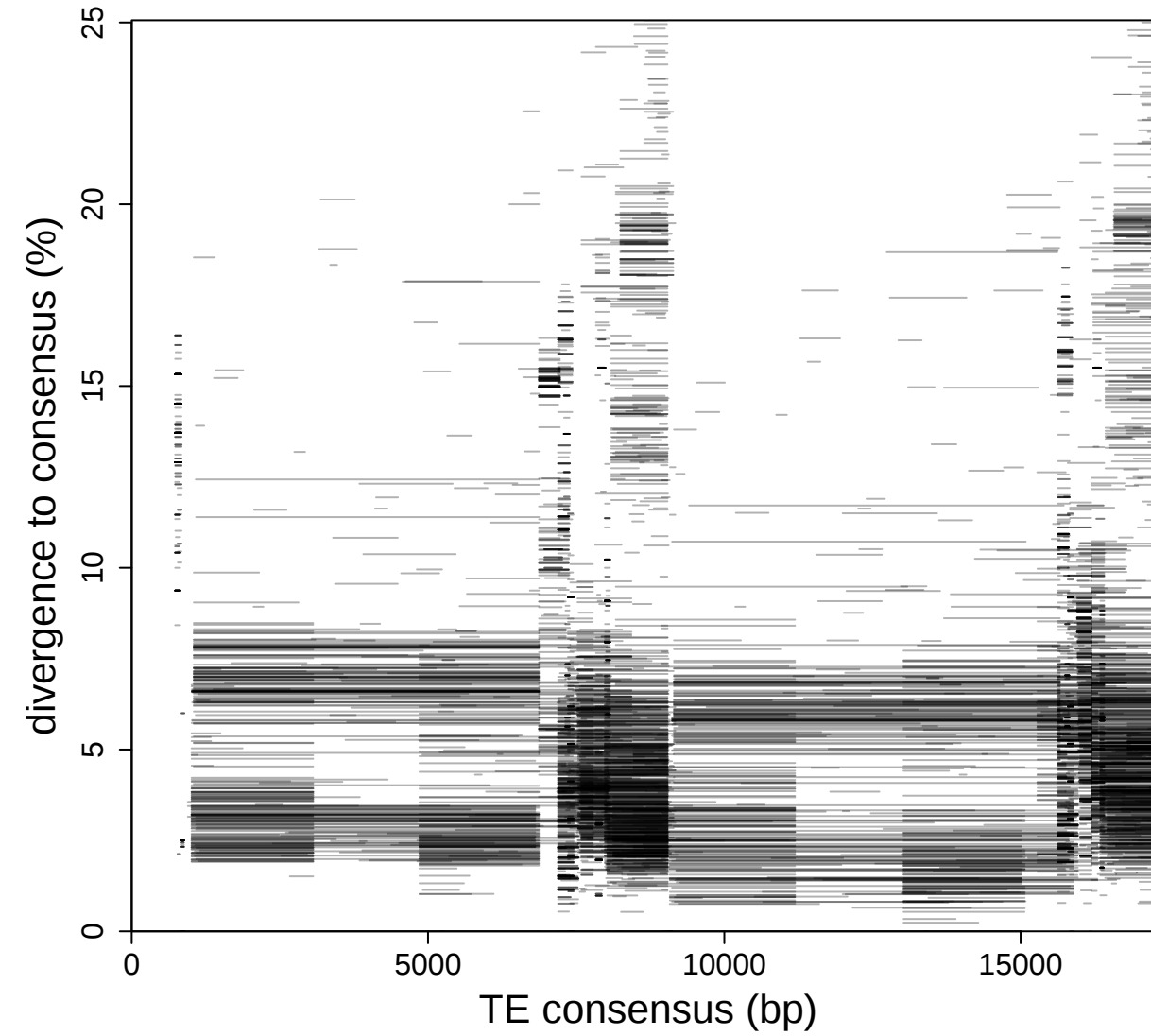
MSA length = 17433





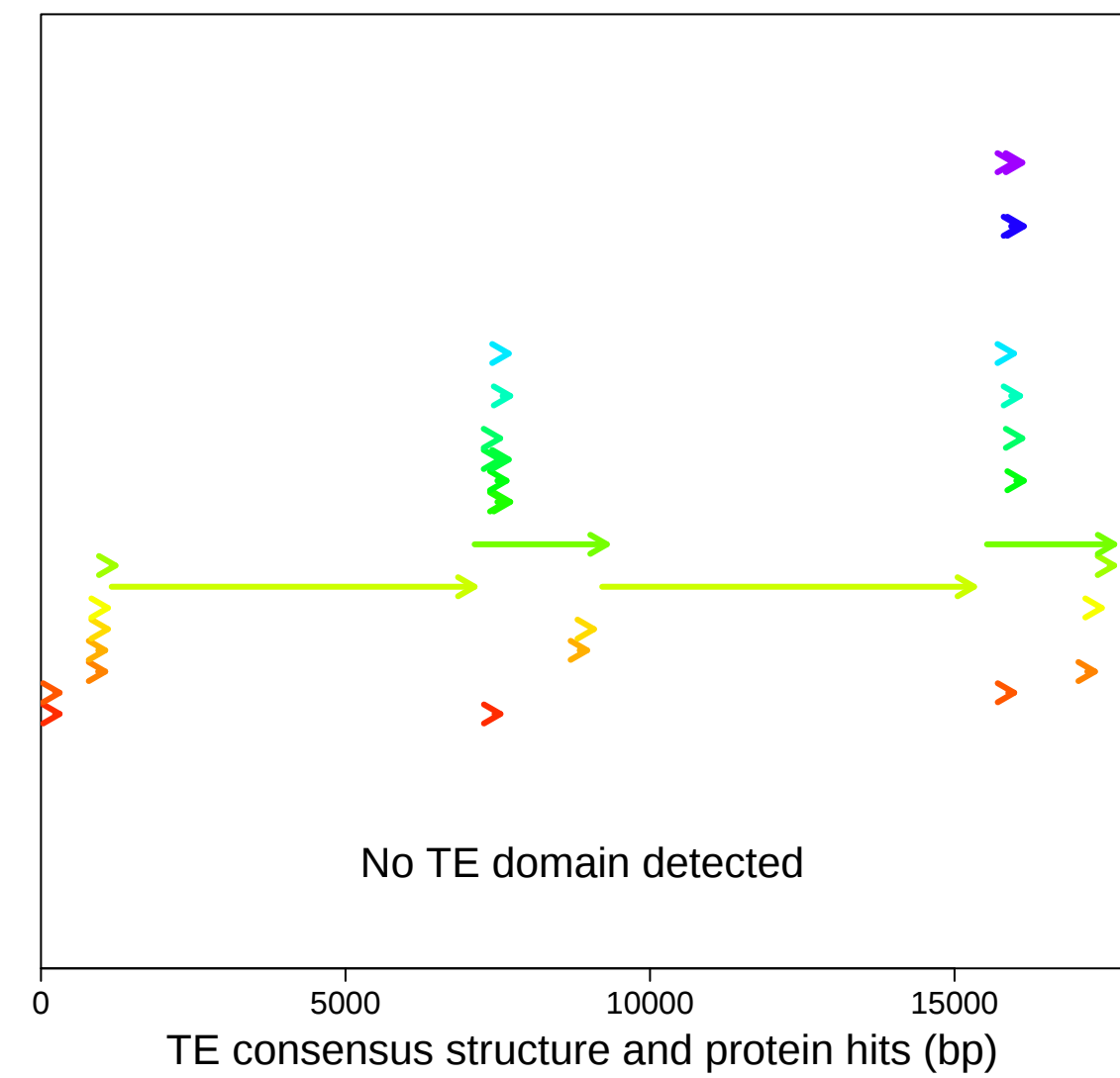
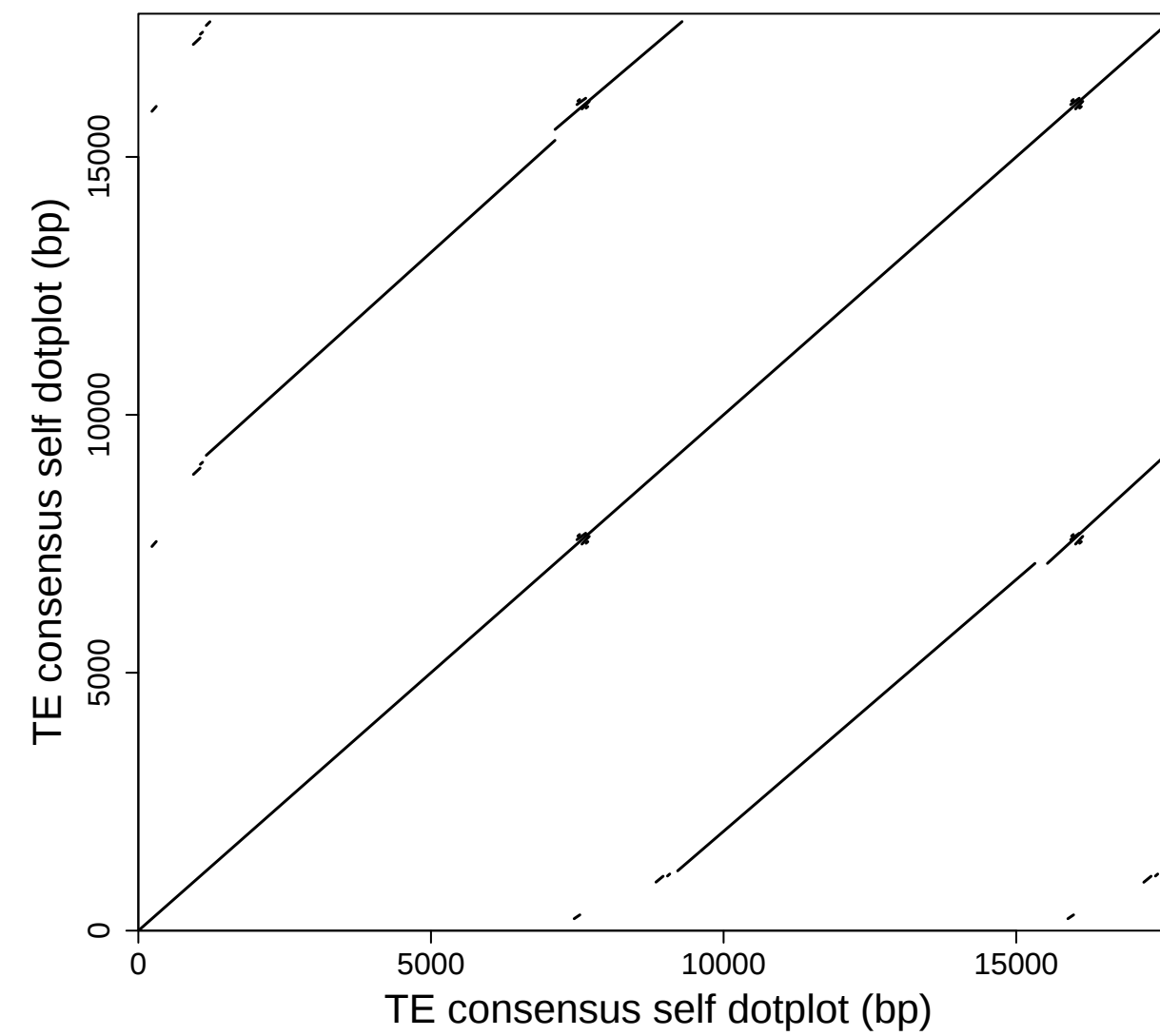
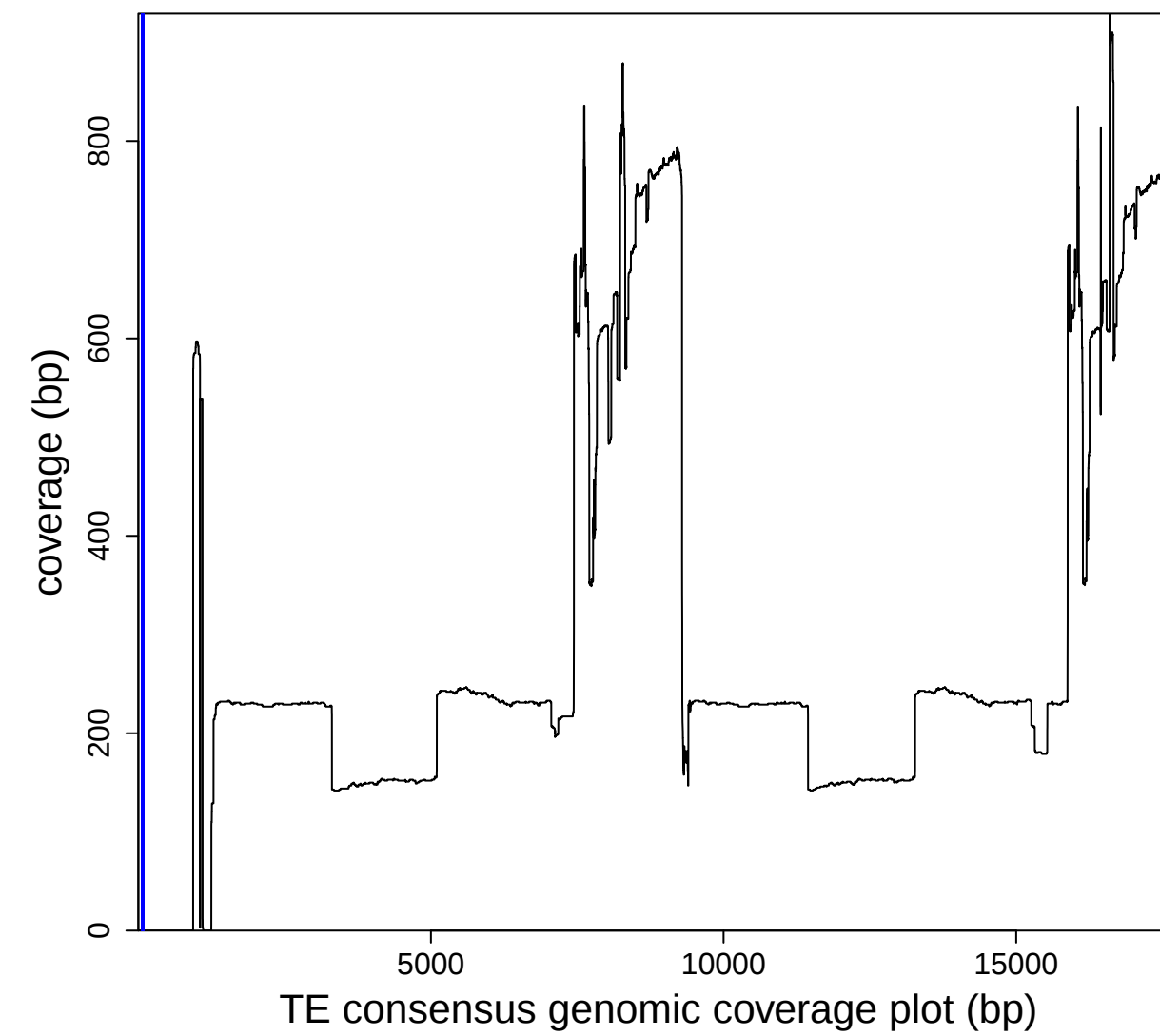
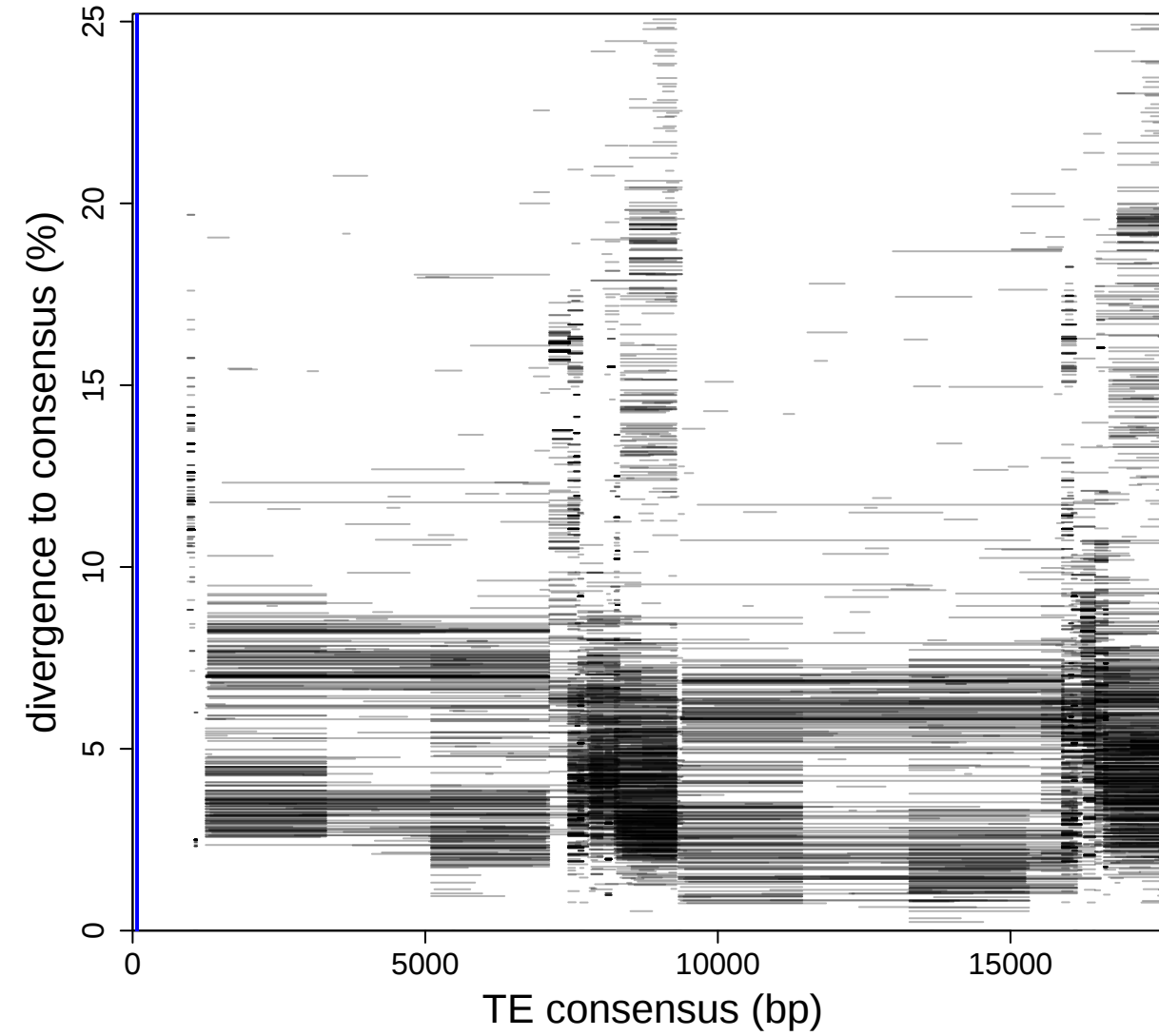
l.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 17433bp; fragments: 6585; full length: 0 (>=15689.7bp)

After TEtrimmer 17433 bp



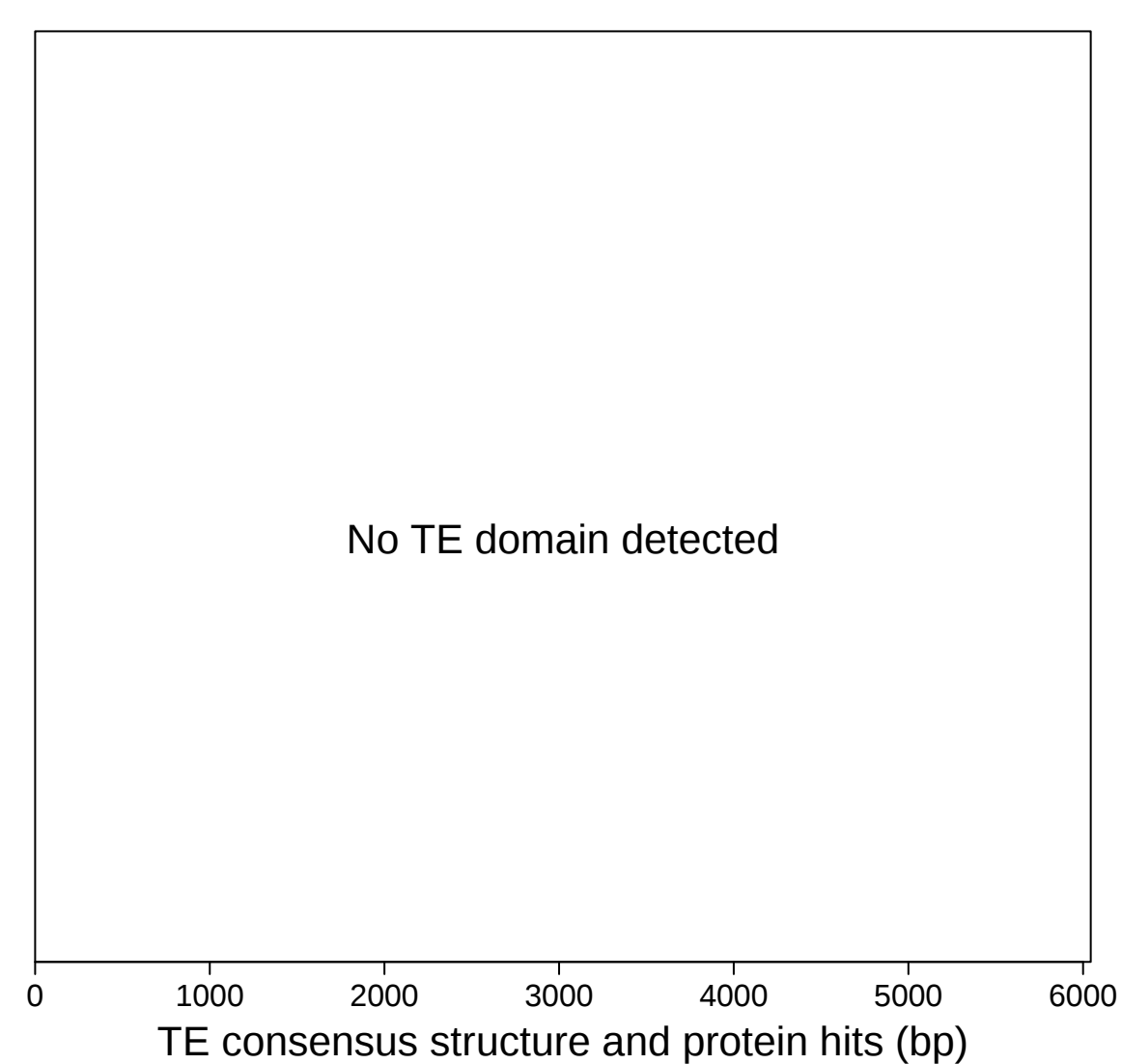
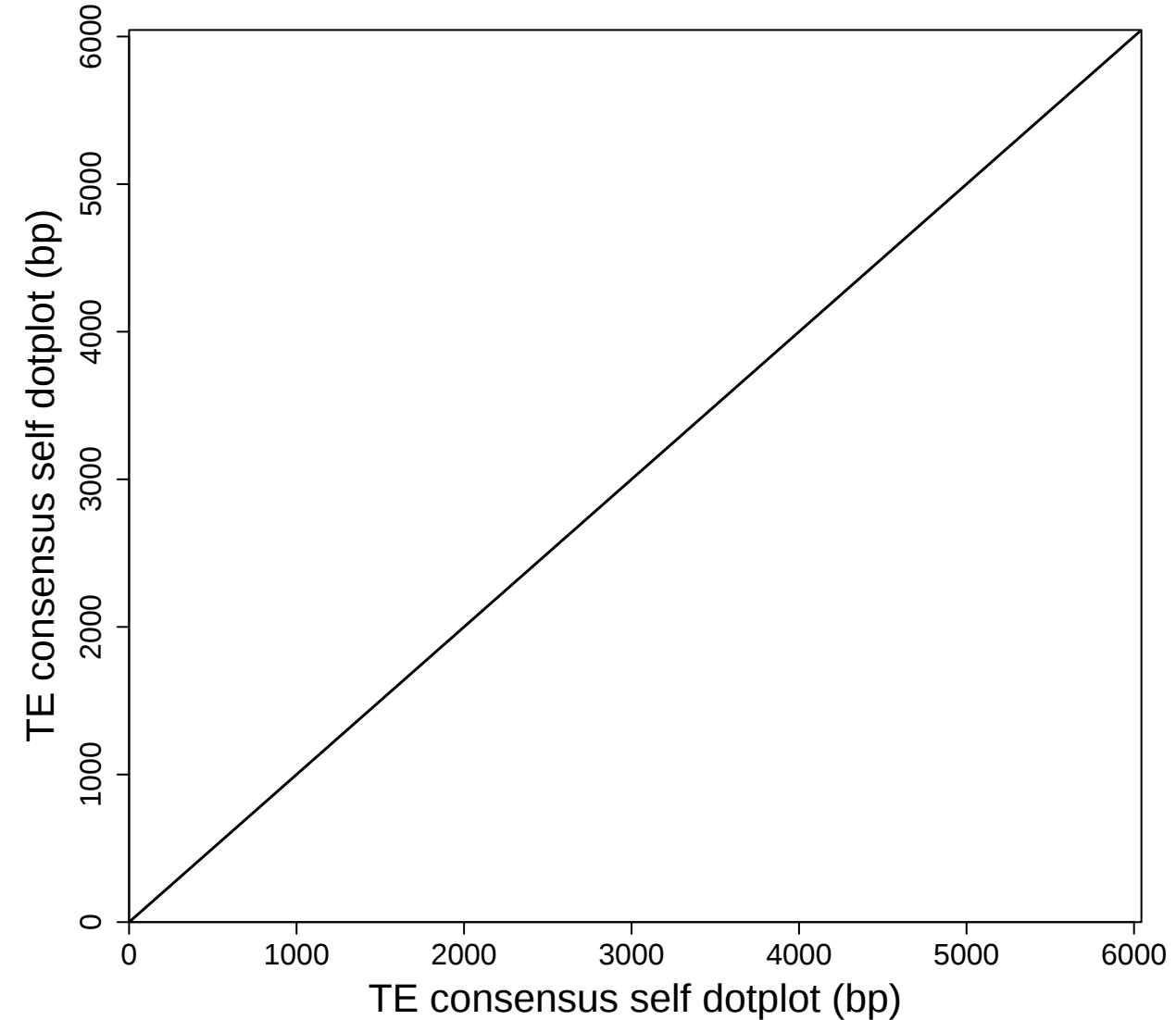
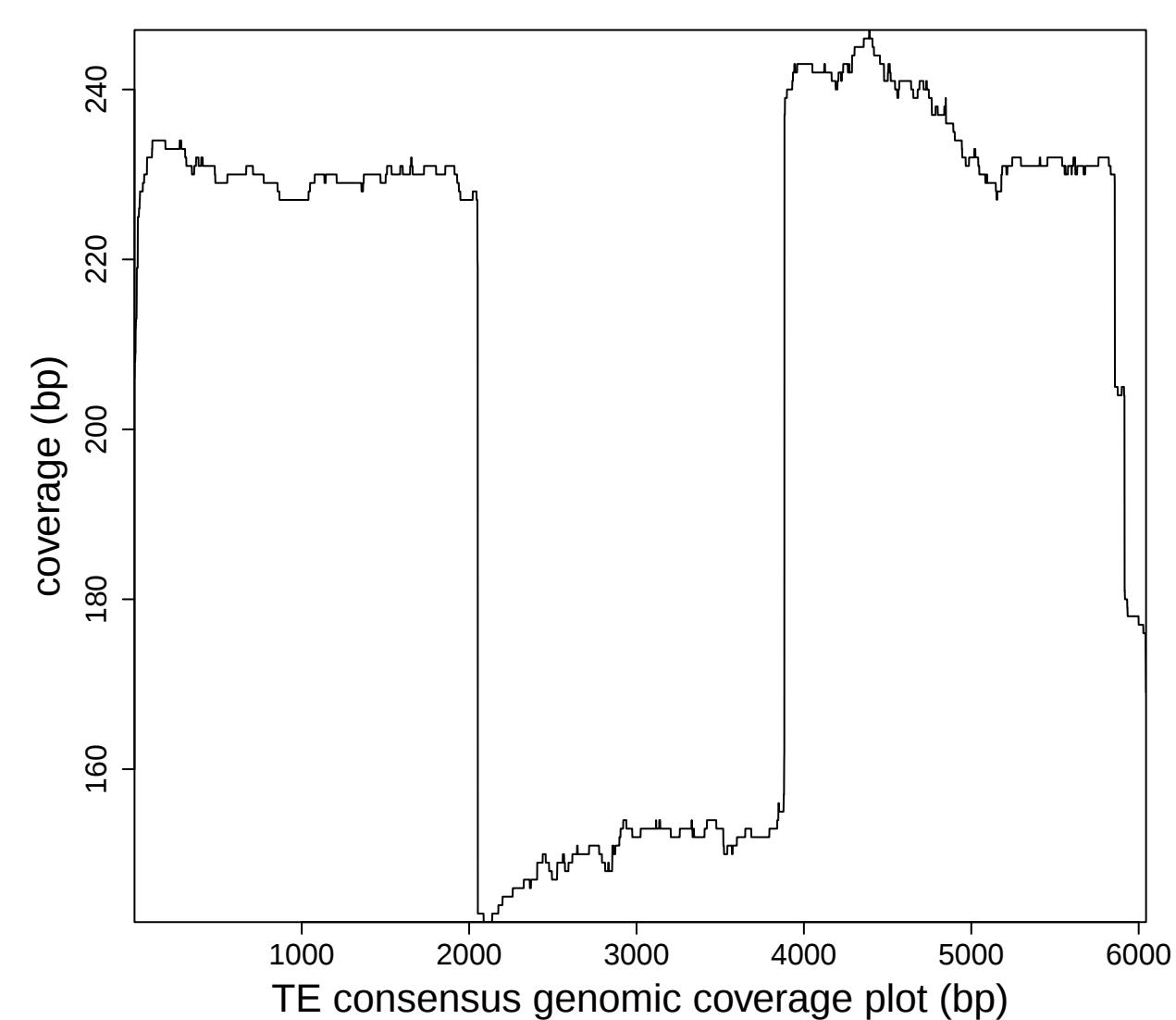
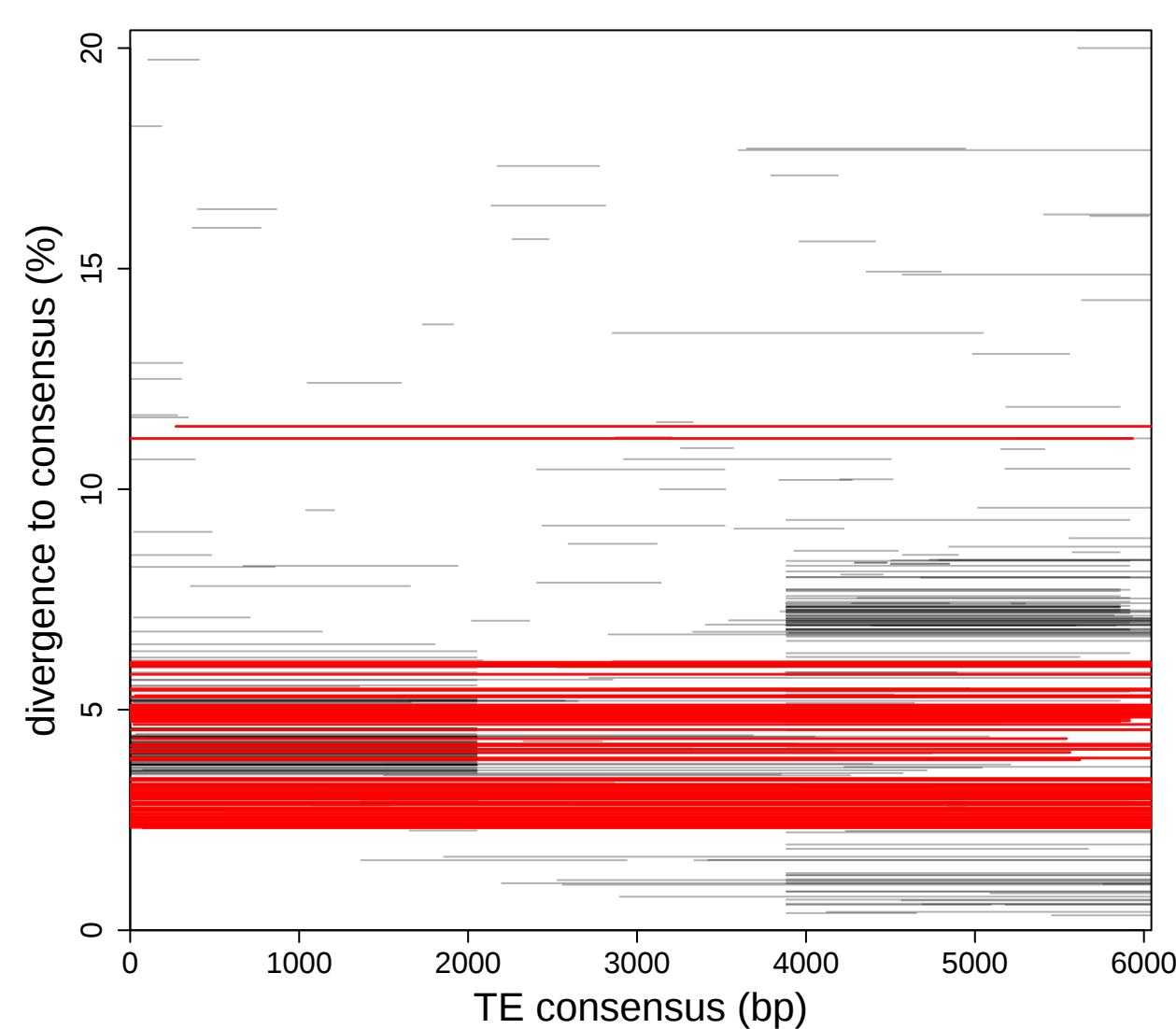
4.fasta.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 17776bp; fragments: 6636; full length: 0 (>=15998.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

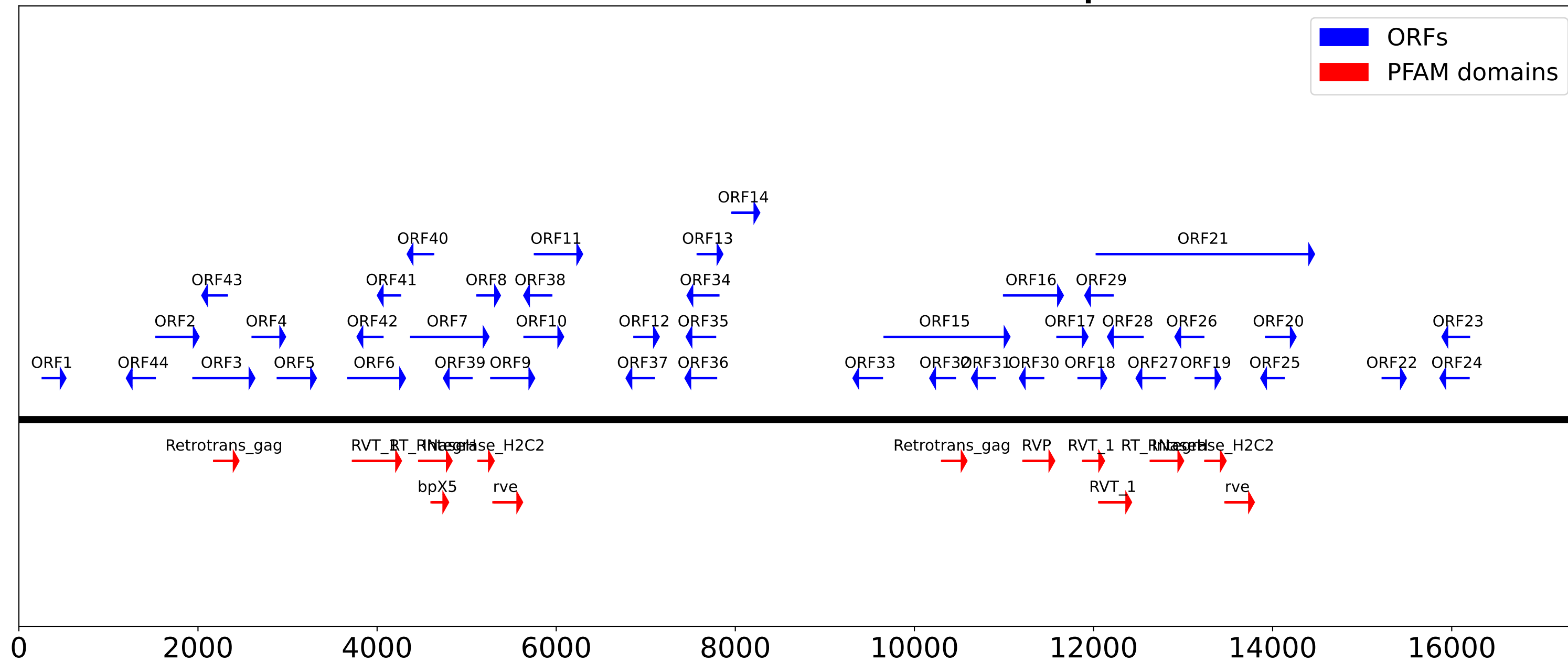


TE: rnd_1_family_284
size: 6044bp; fragments: 426; full length: 111 (≥ 5439.6 bp)

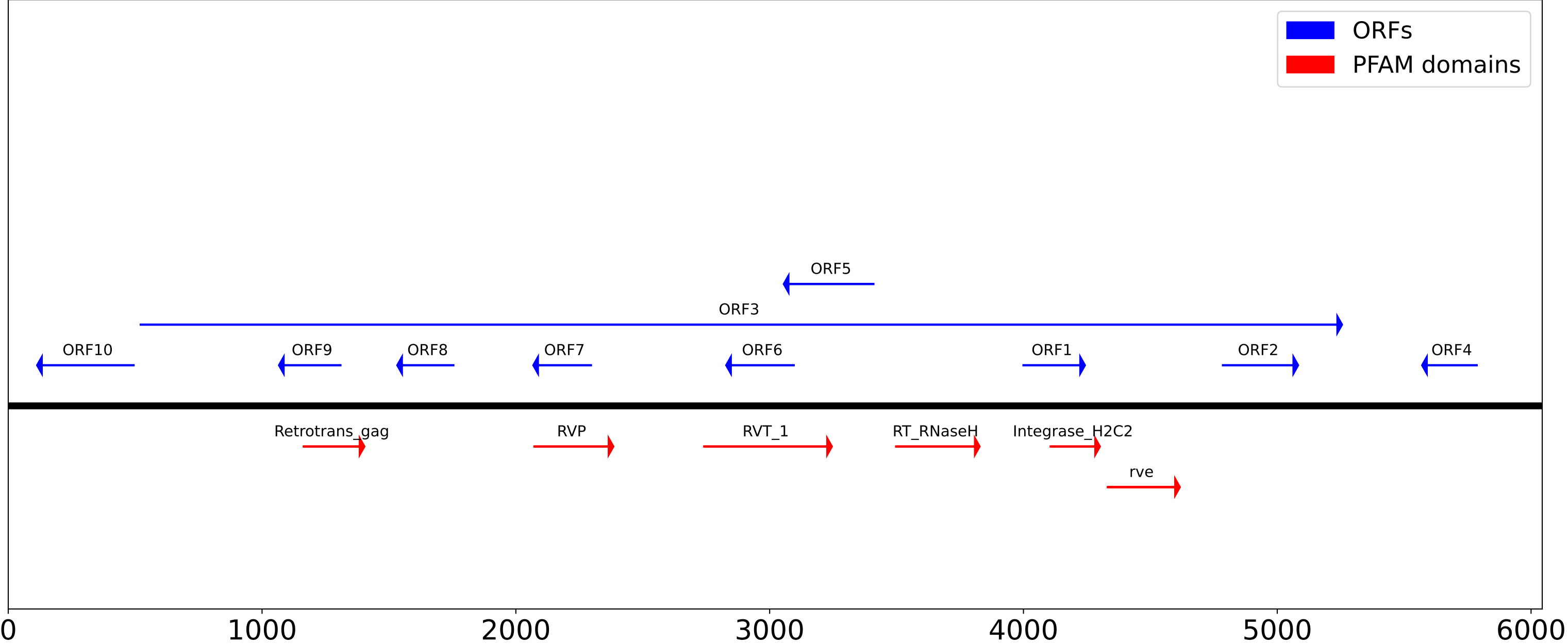
Before TEtrimmer 6044 bp



After TETrimmer ORF and PFAM domain plot



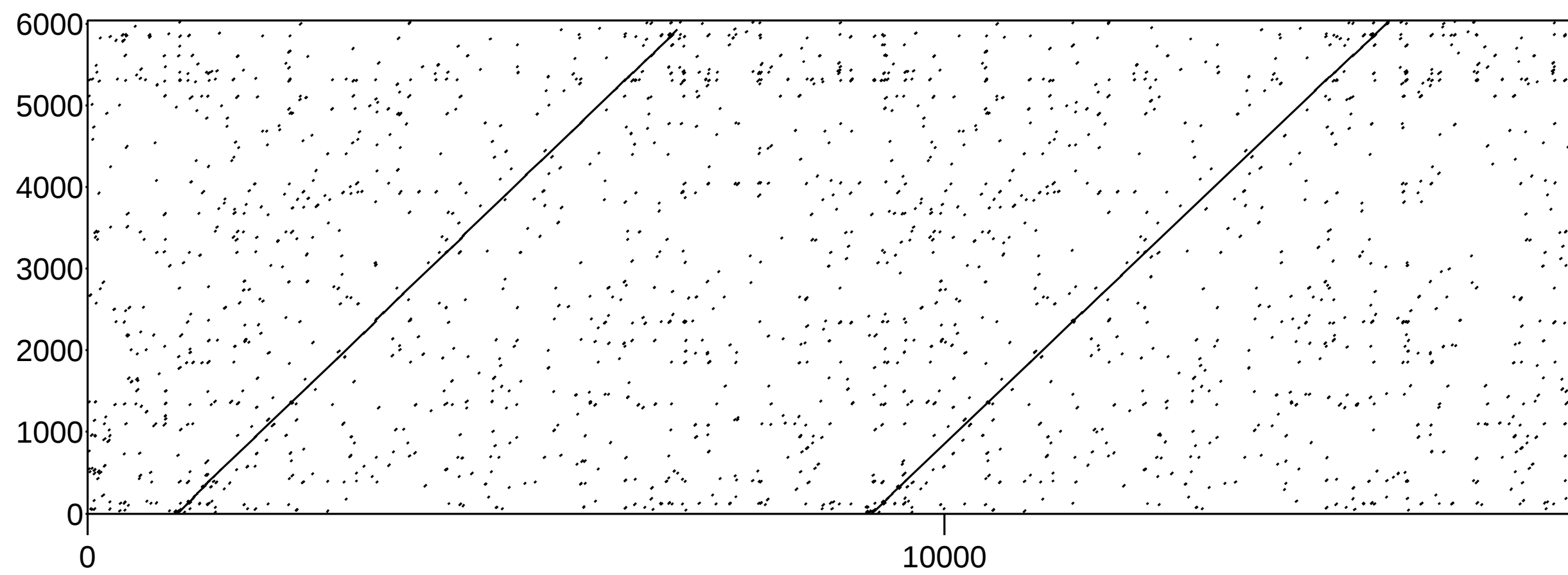
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

E consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)