

Start crop Point

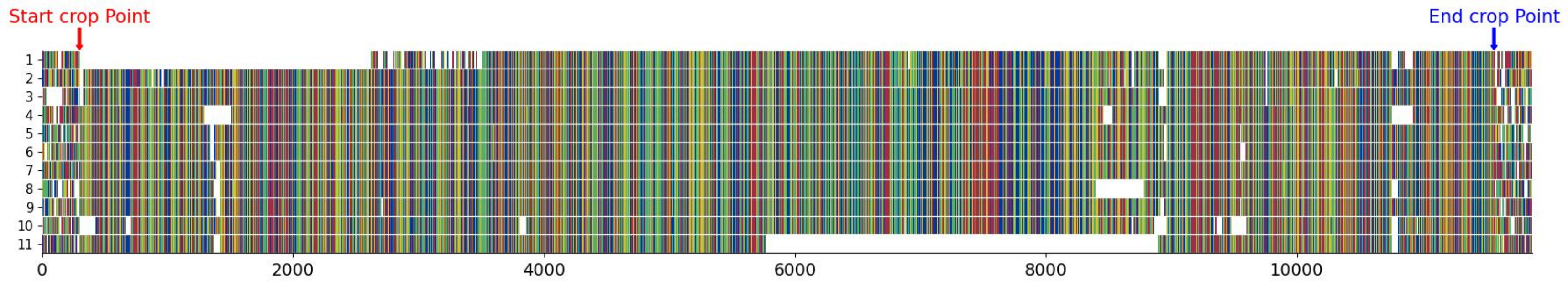


End crop Point



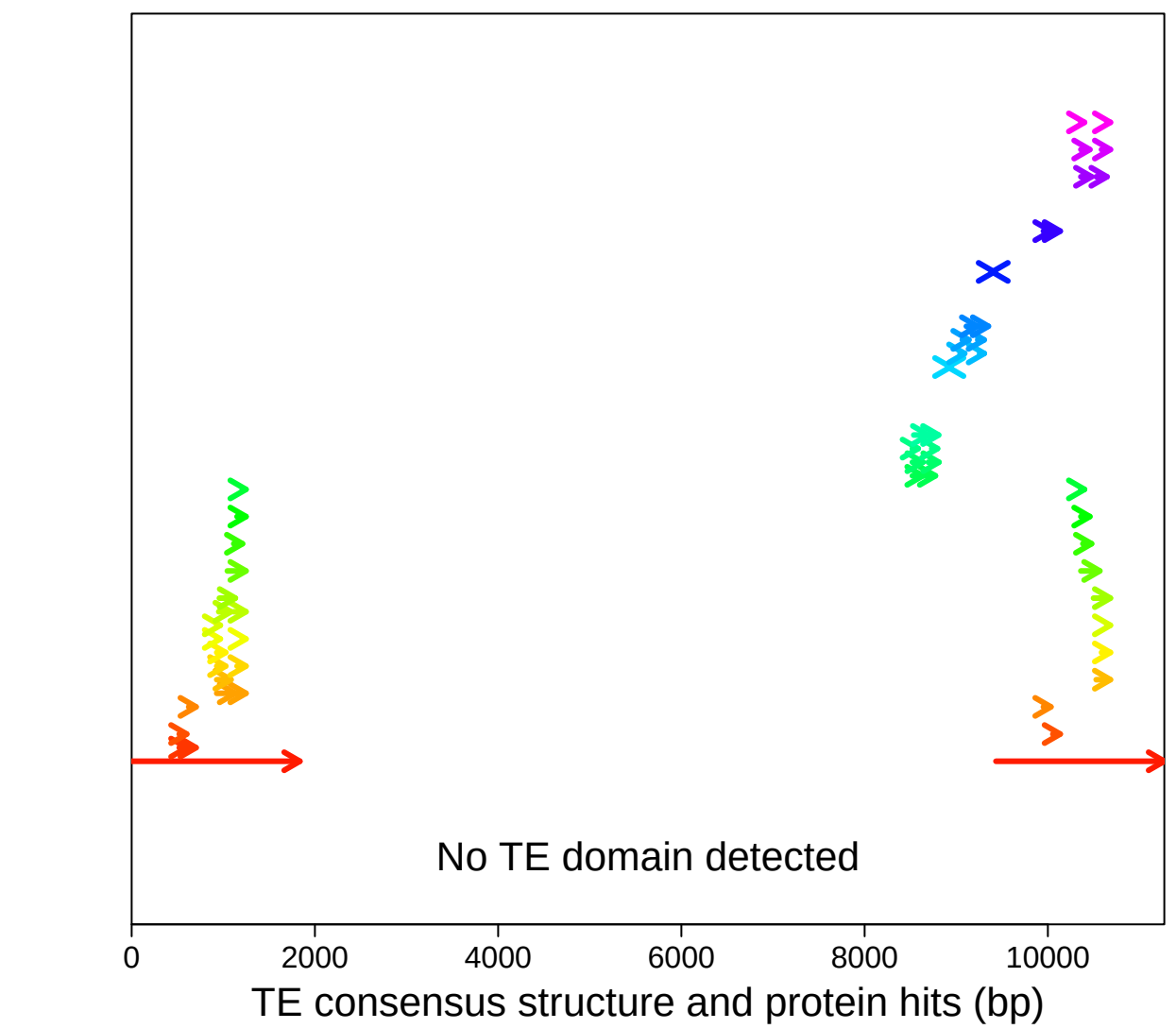
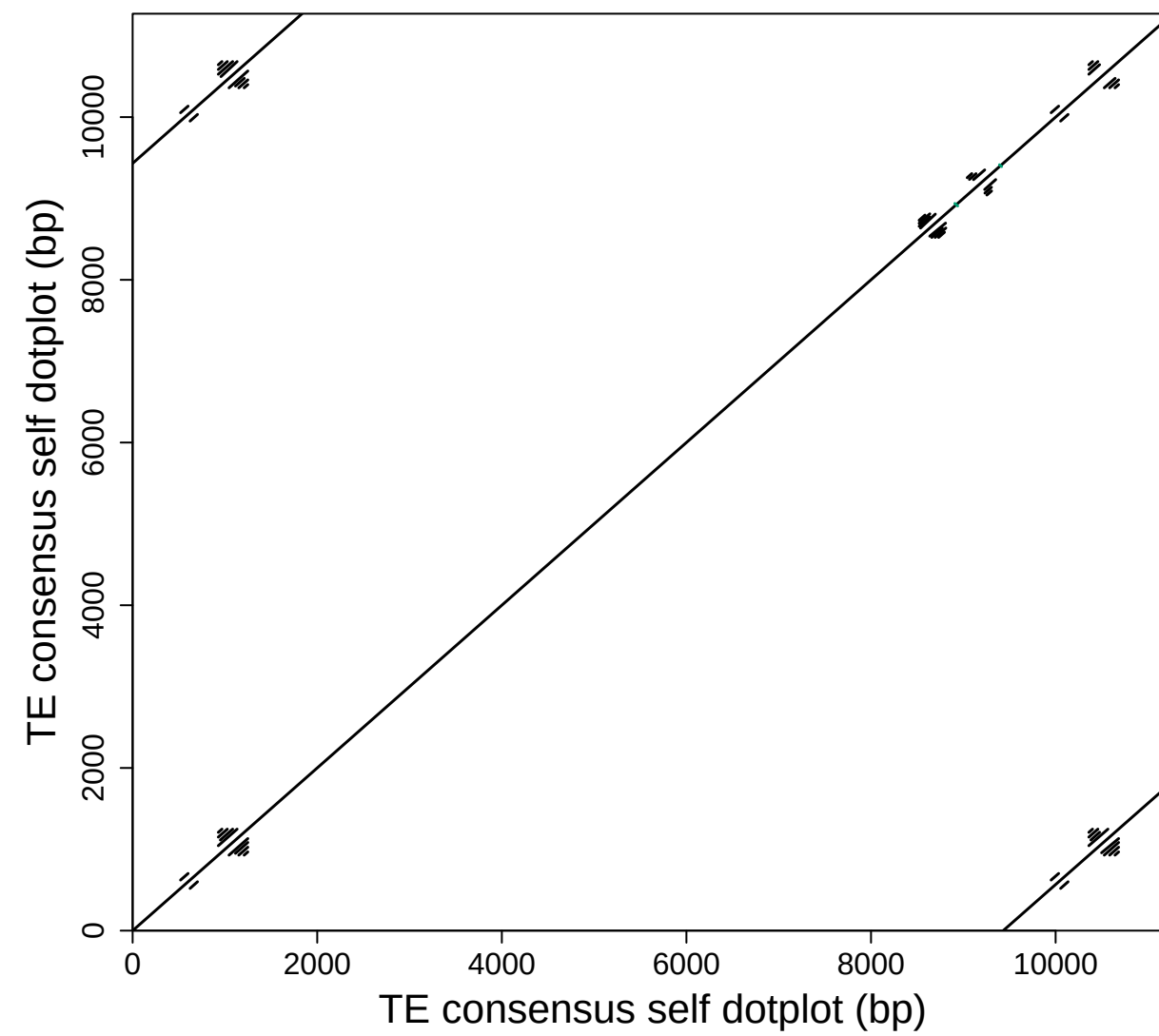
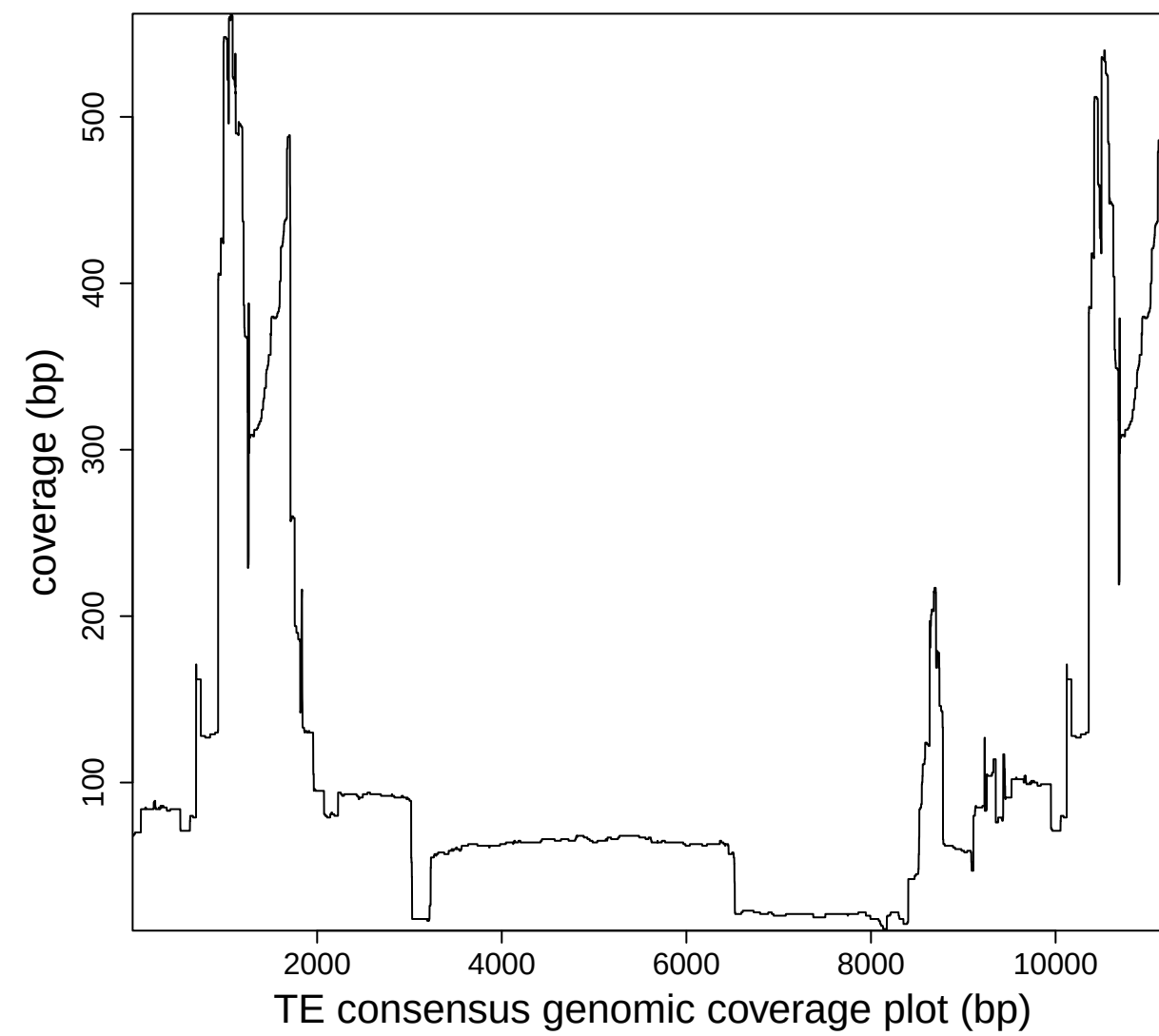
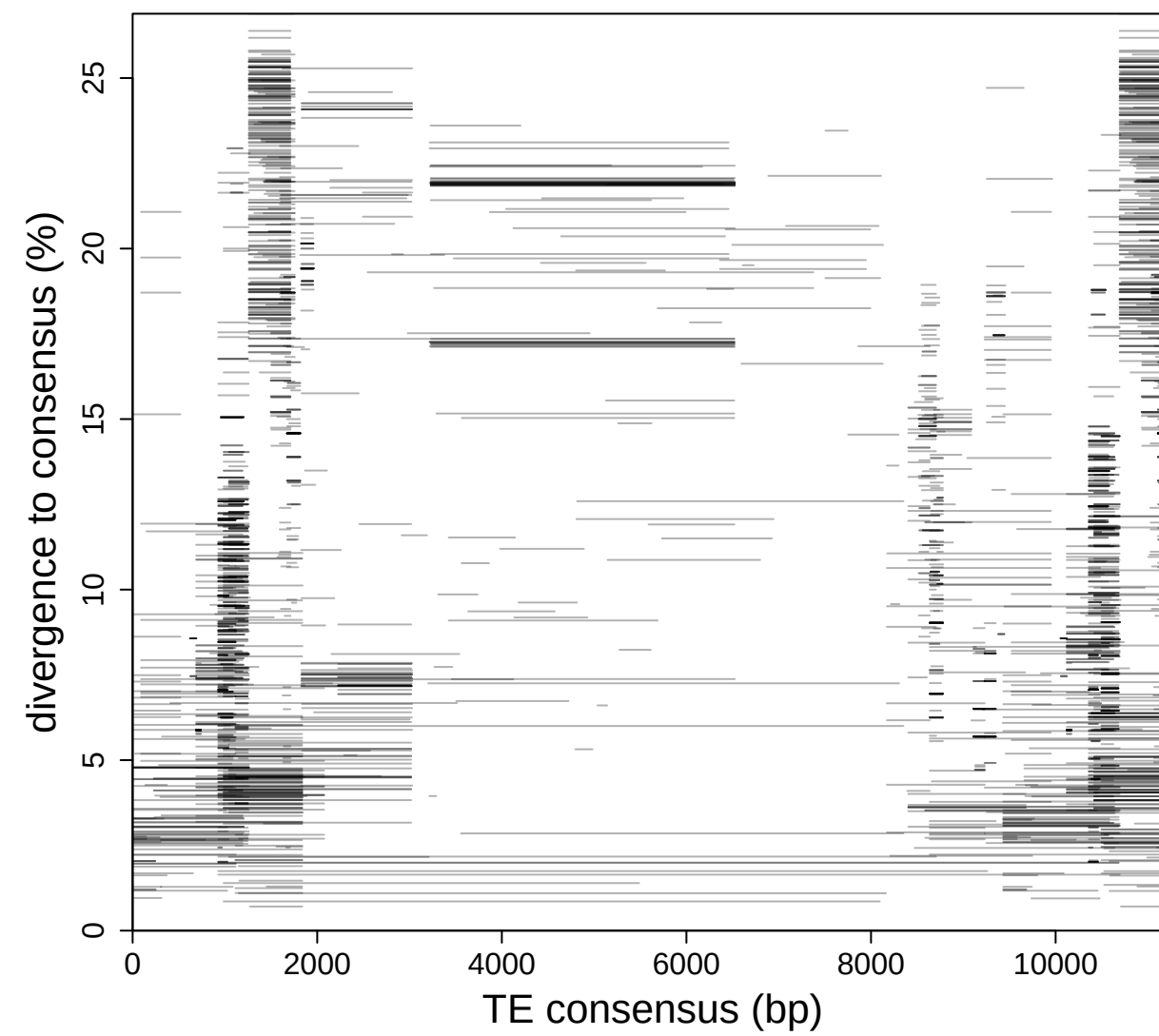
MSA length = 11271





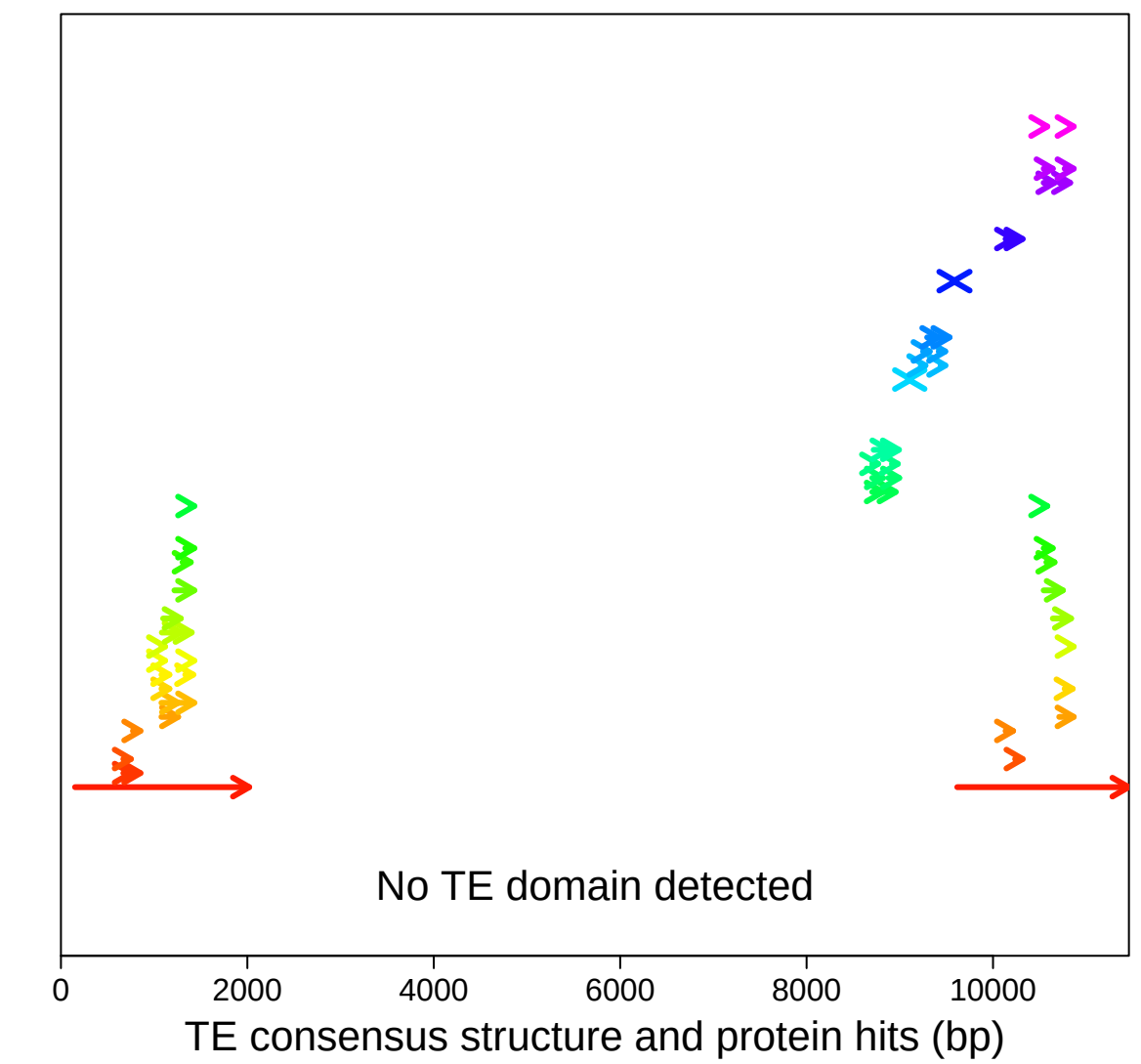
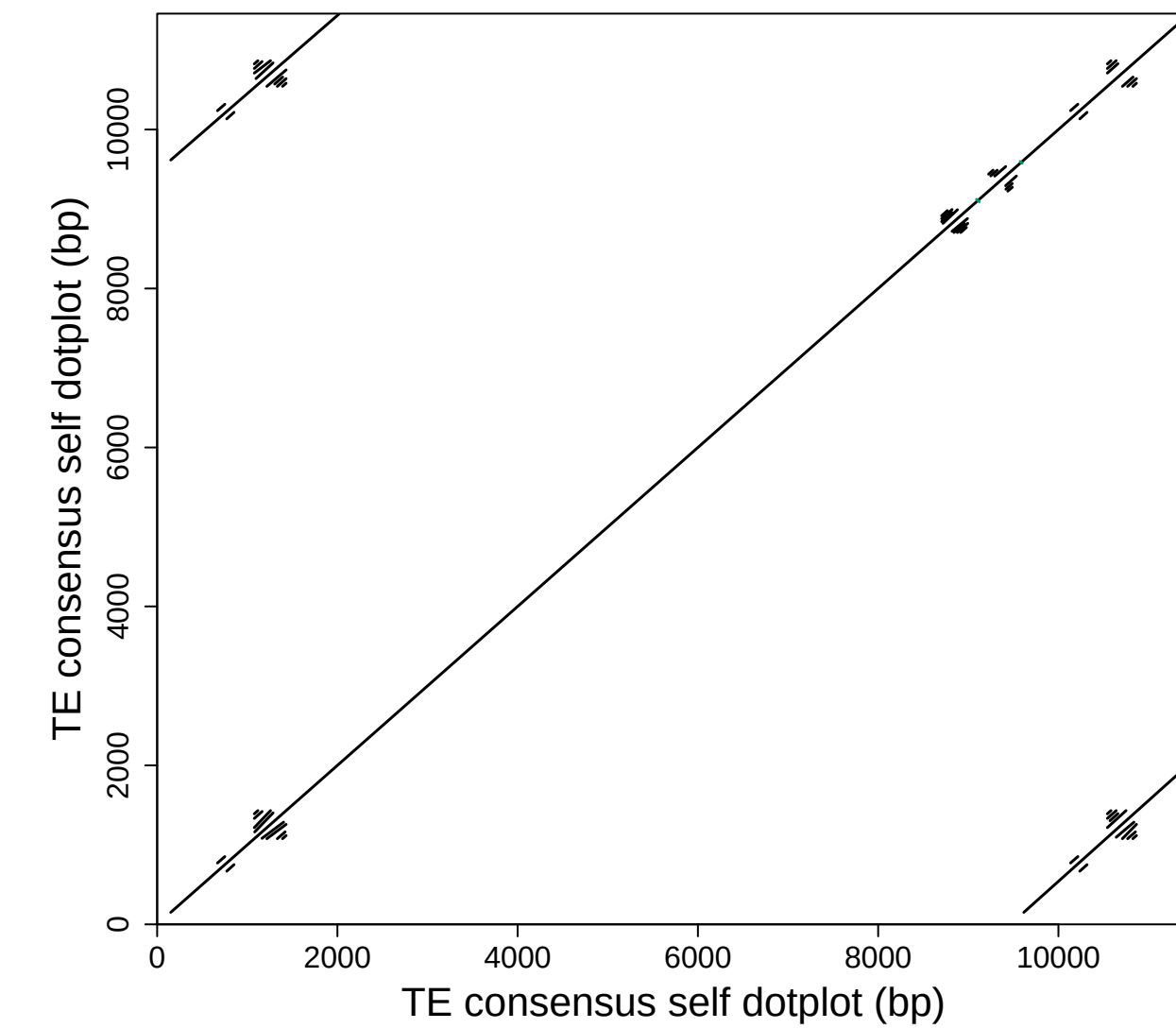
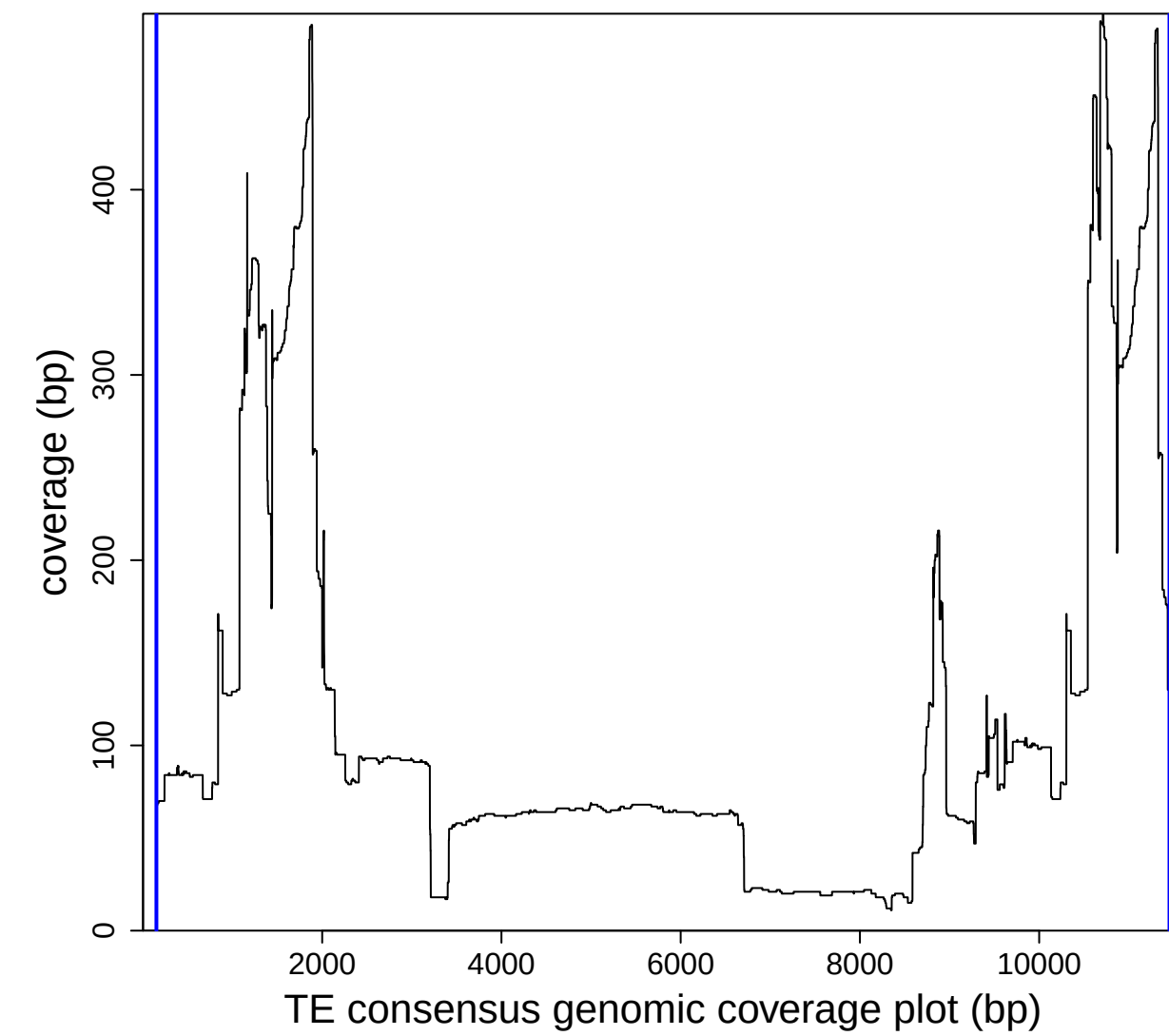
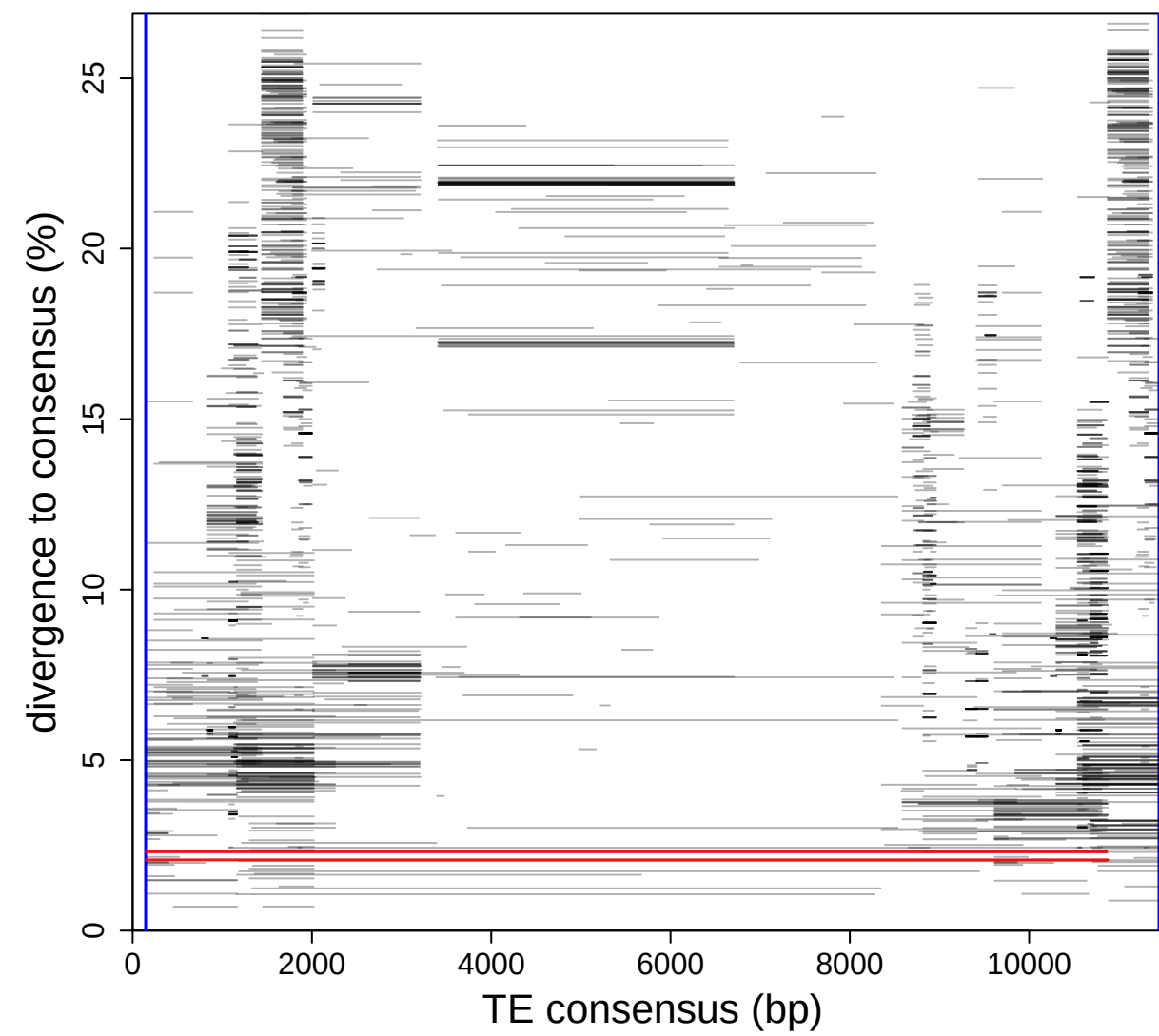
ed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 11271bp; fragments: 2732; full length: 0 (>=10143.9bp)

After TETrimmer 11271 bp



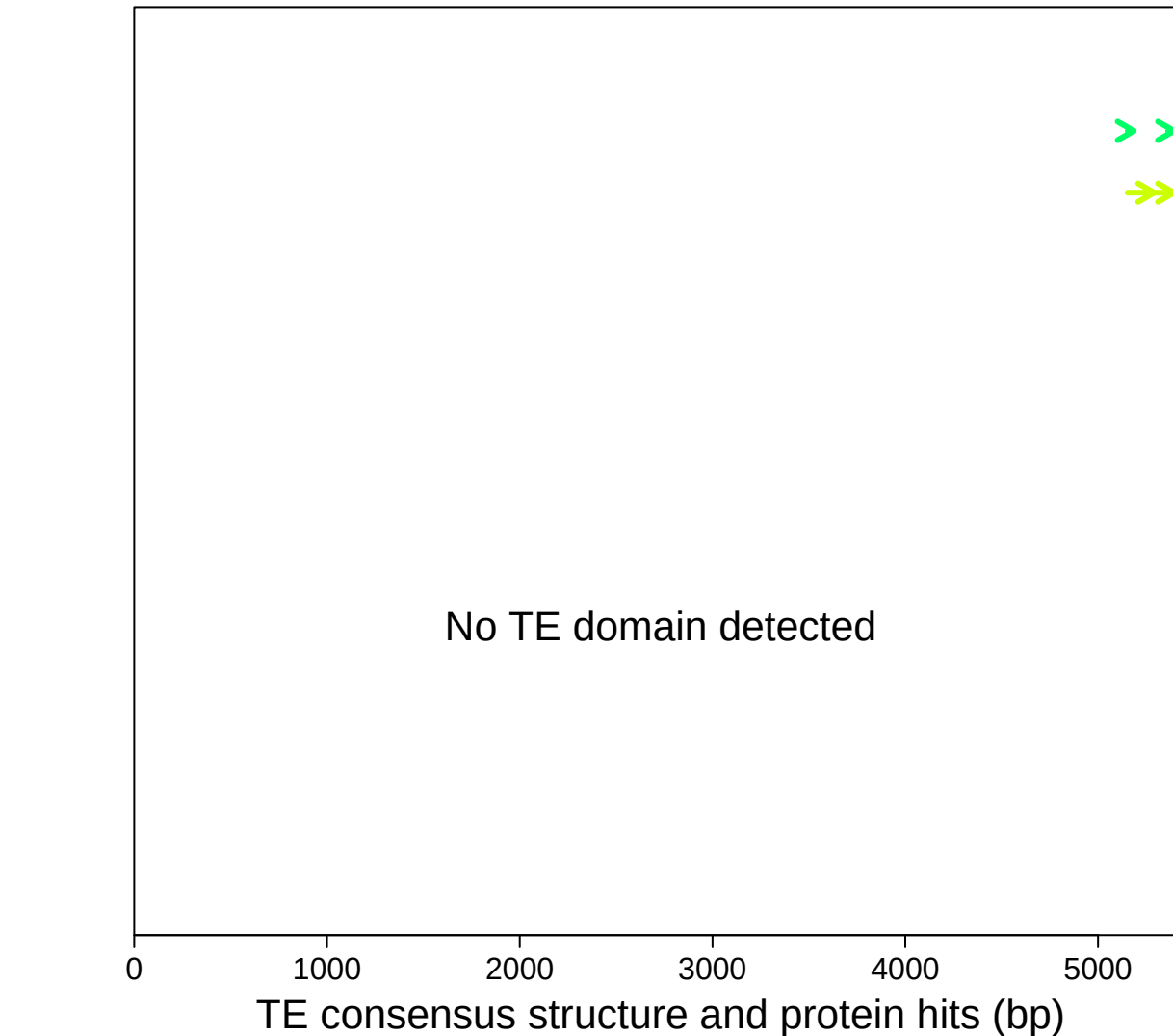
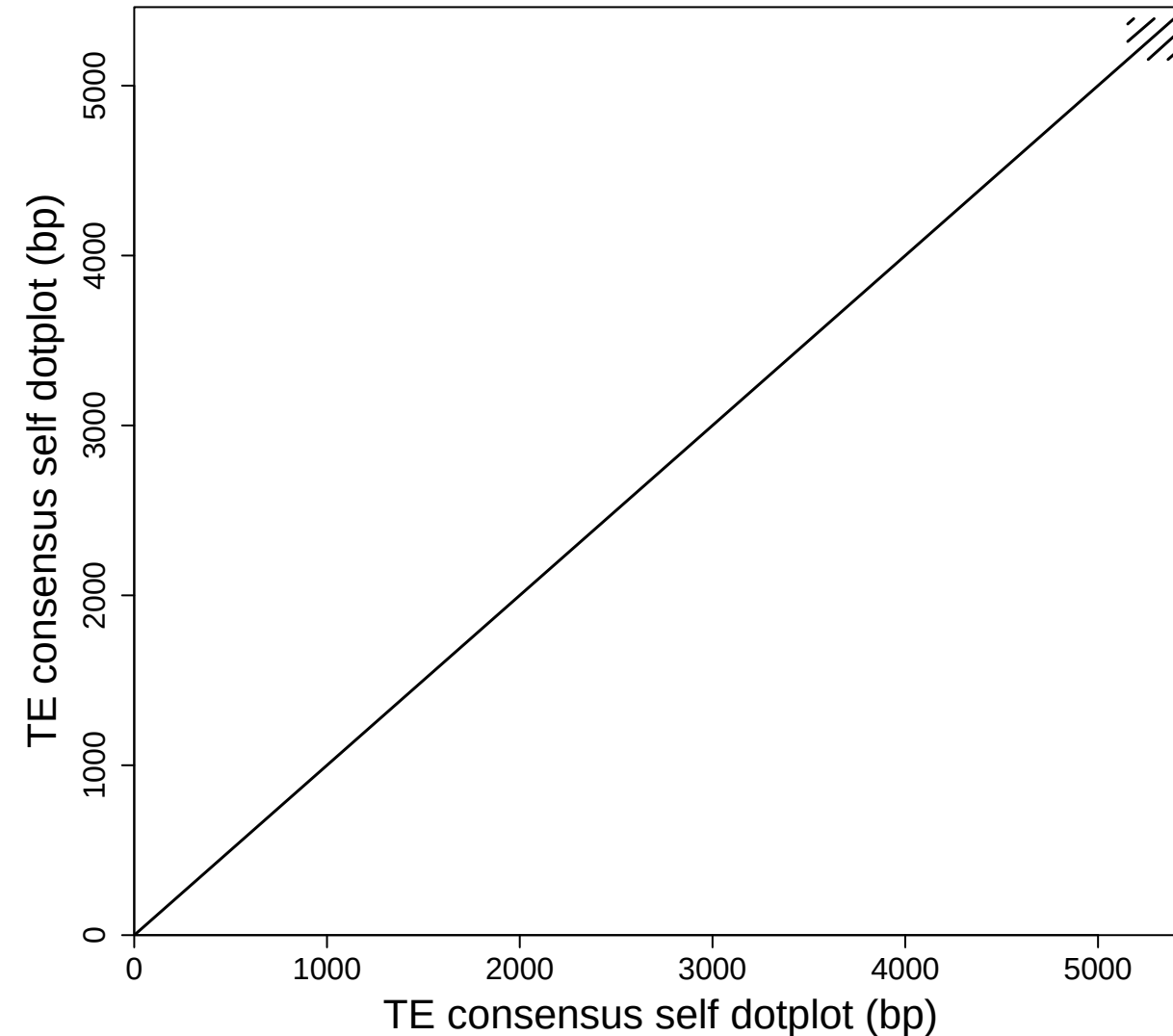
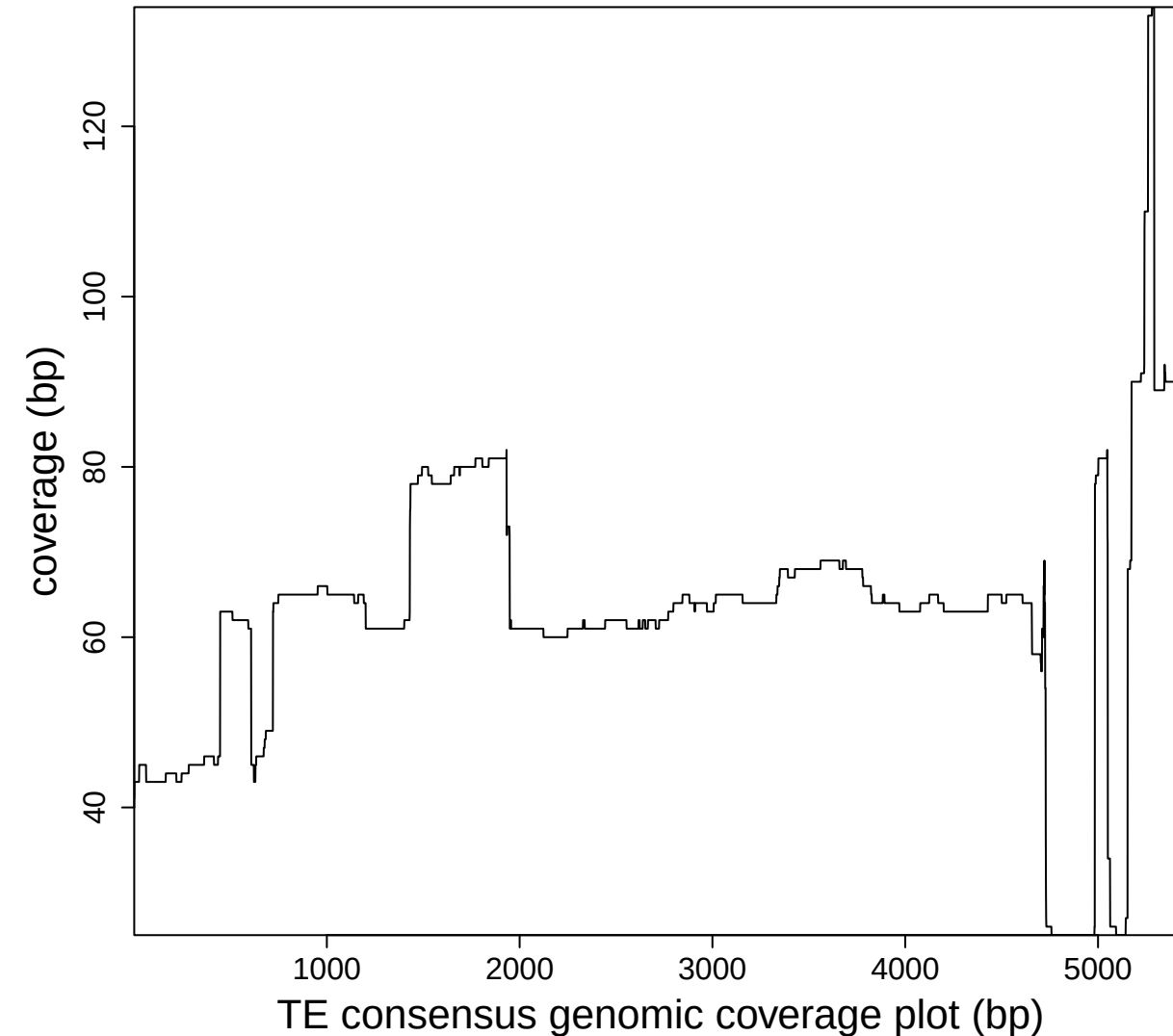
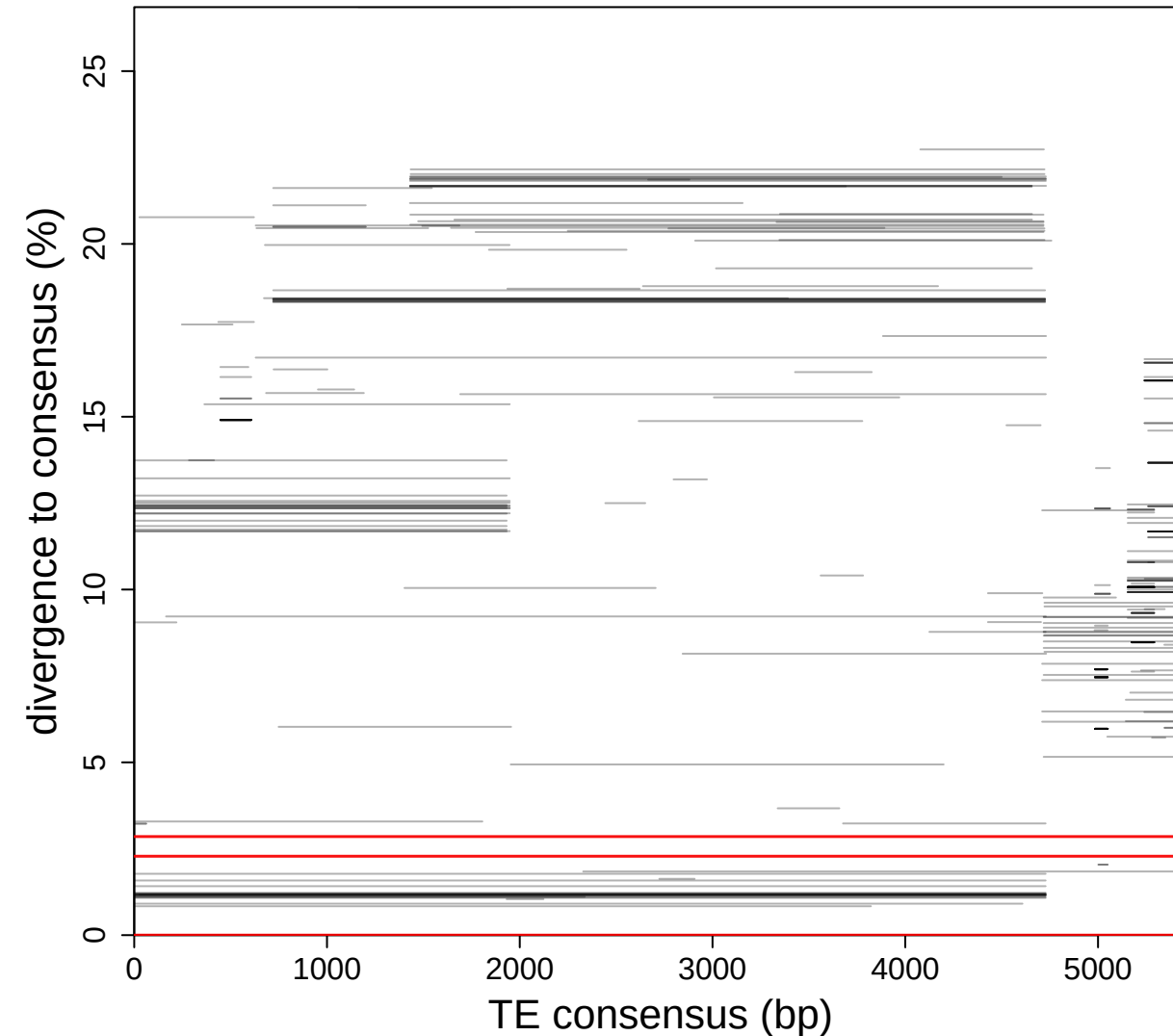
ly_639.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 11604bp; fragments: 2481; full length: 2 (>=10443.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

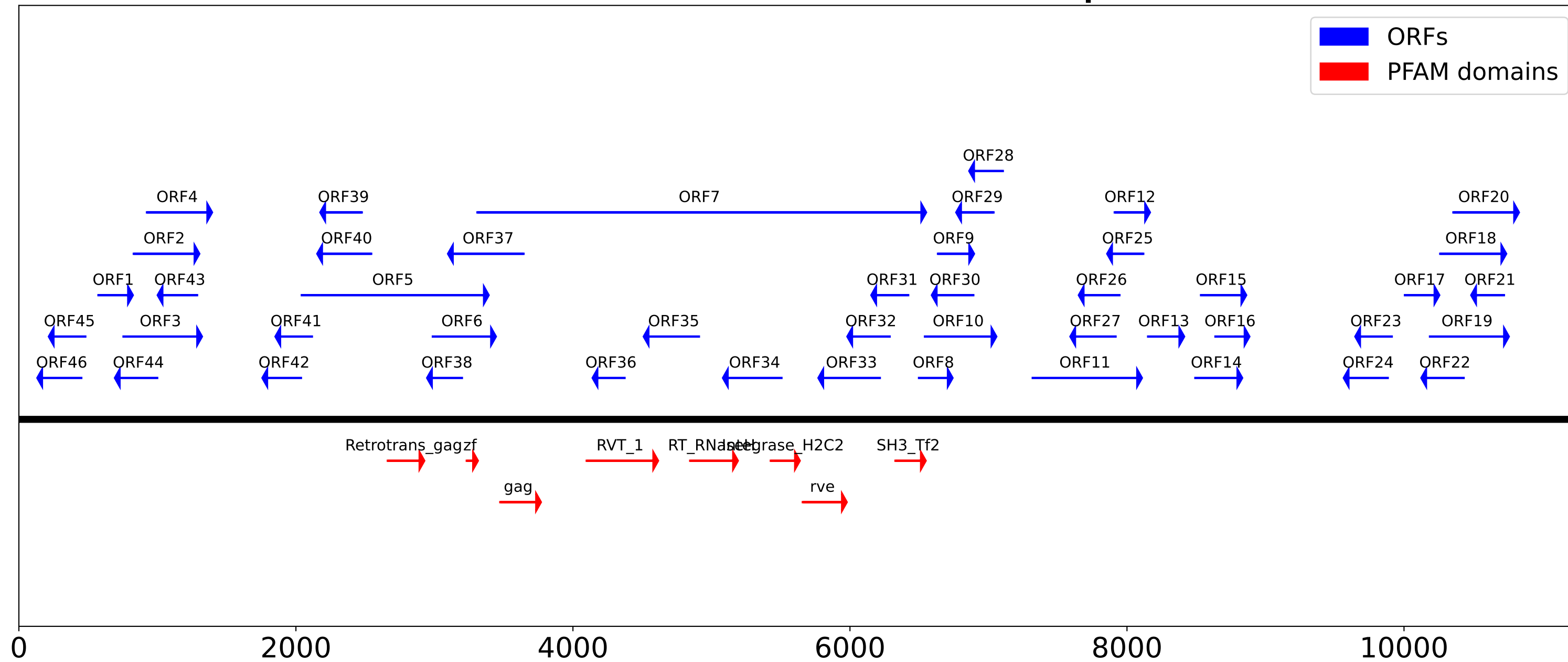


TE: ltr_1_family_639
size: 5462bp; fragments: 334; full length: 3 (>=4915.8bp)

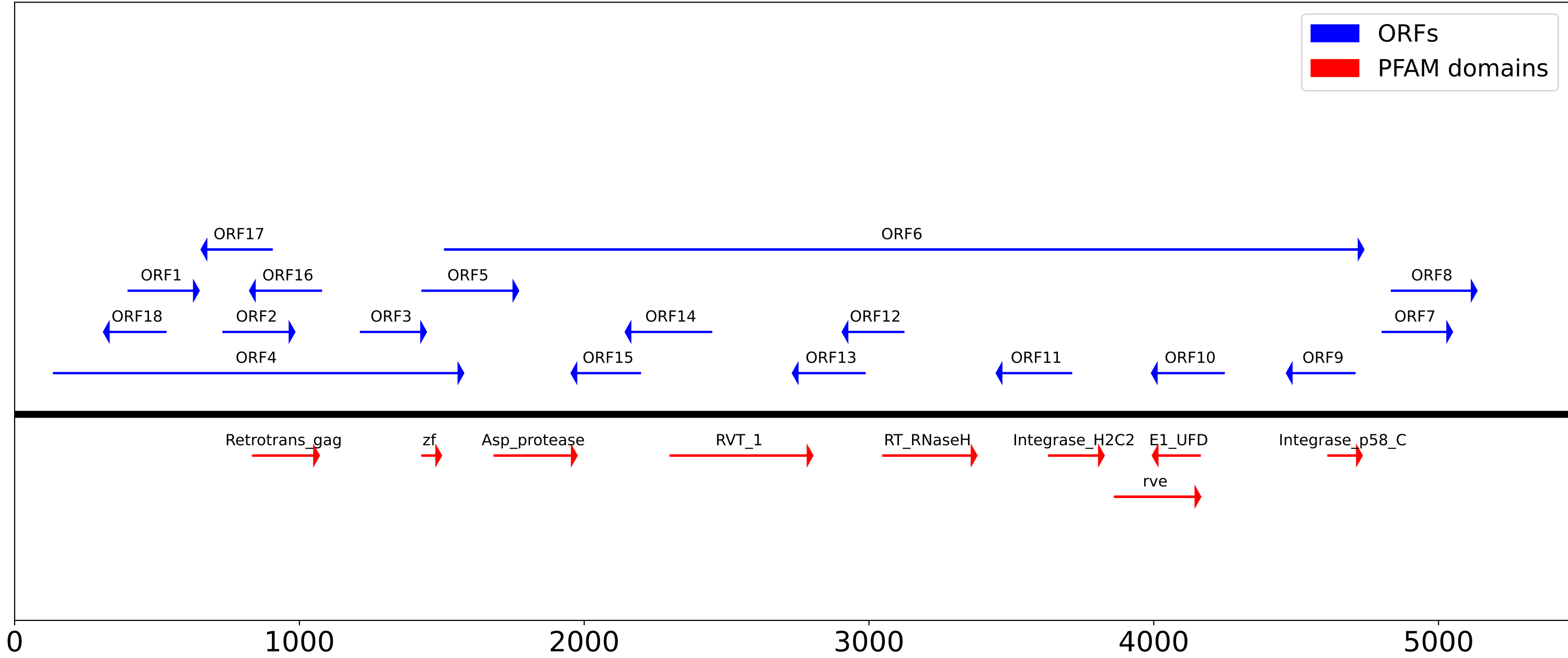
Before TEtrimmer 5462 bp



After TEtrimmer ORF and PFAM domain plot



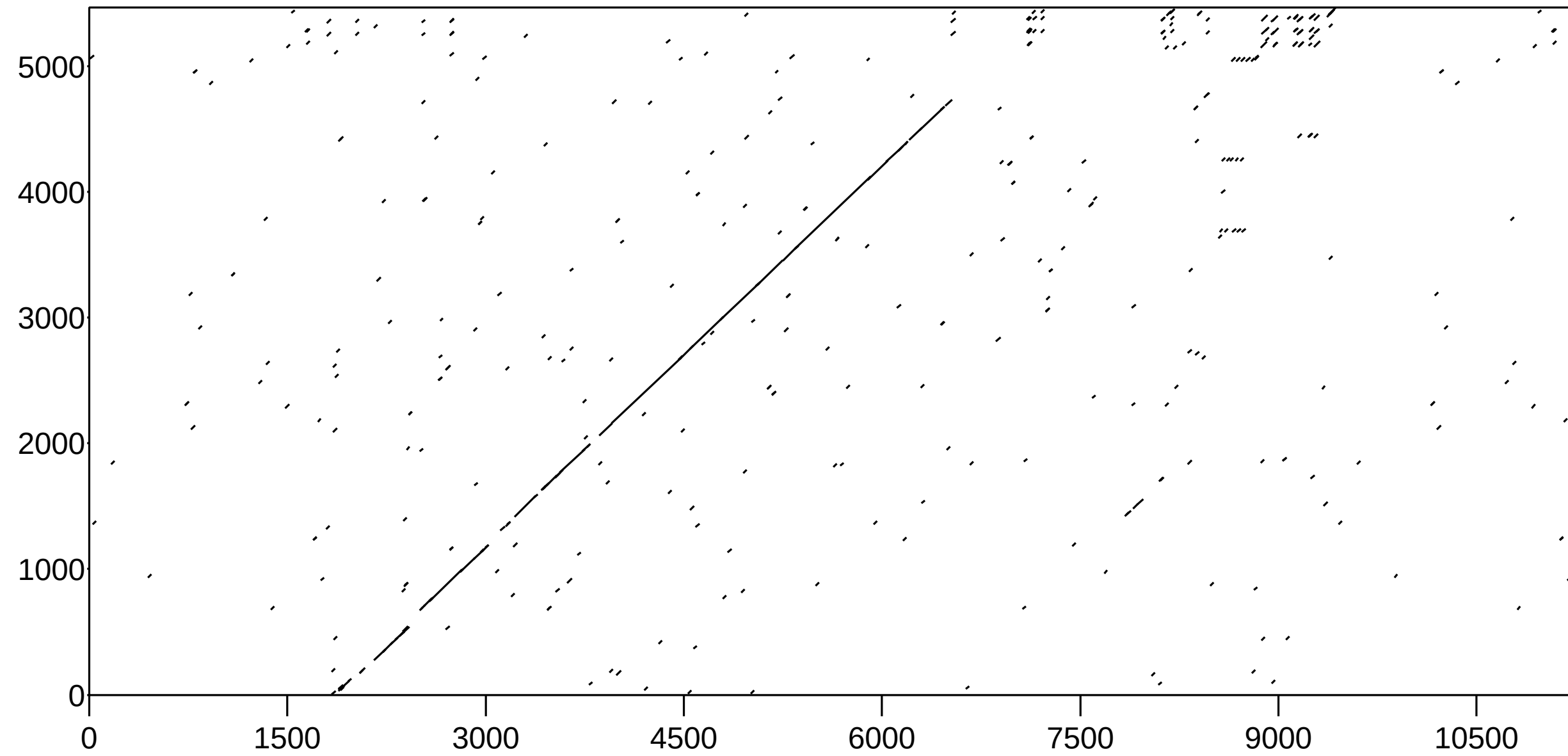
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)