

Start crop Point

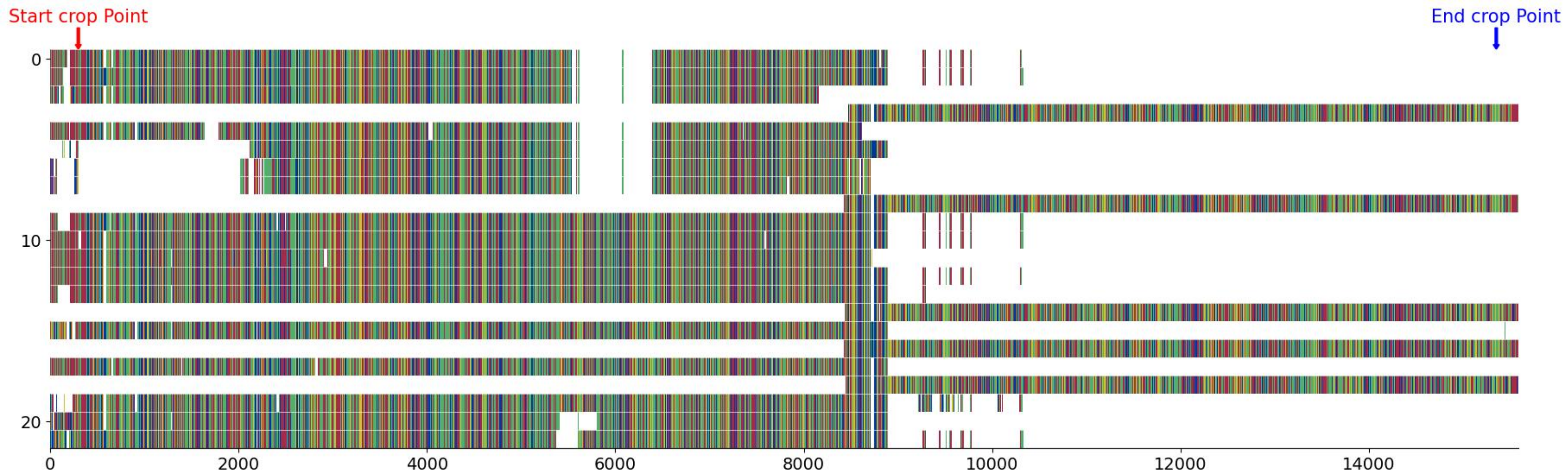


End crop Point



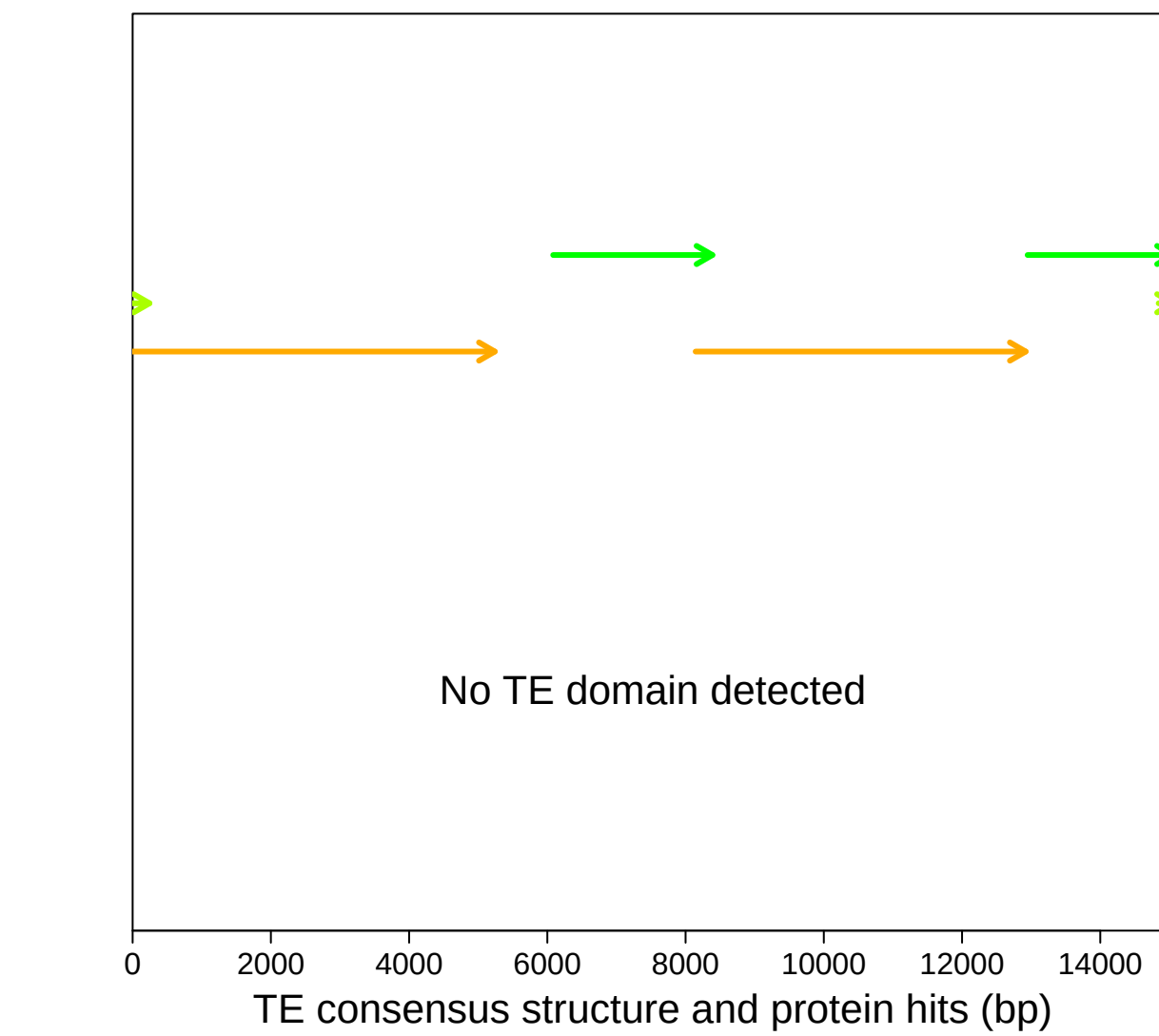
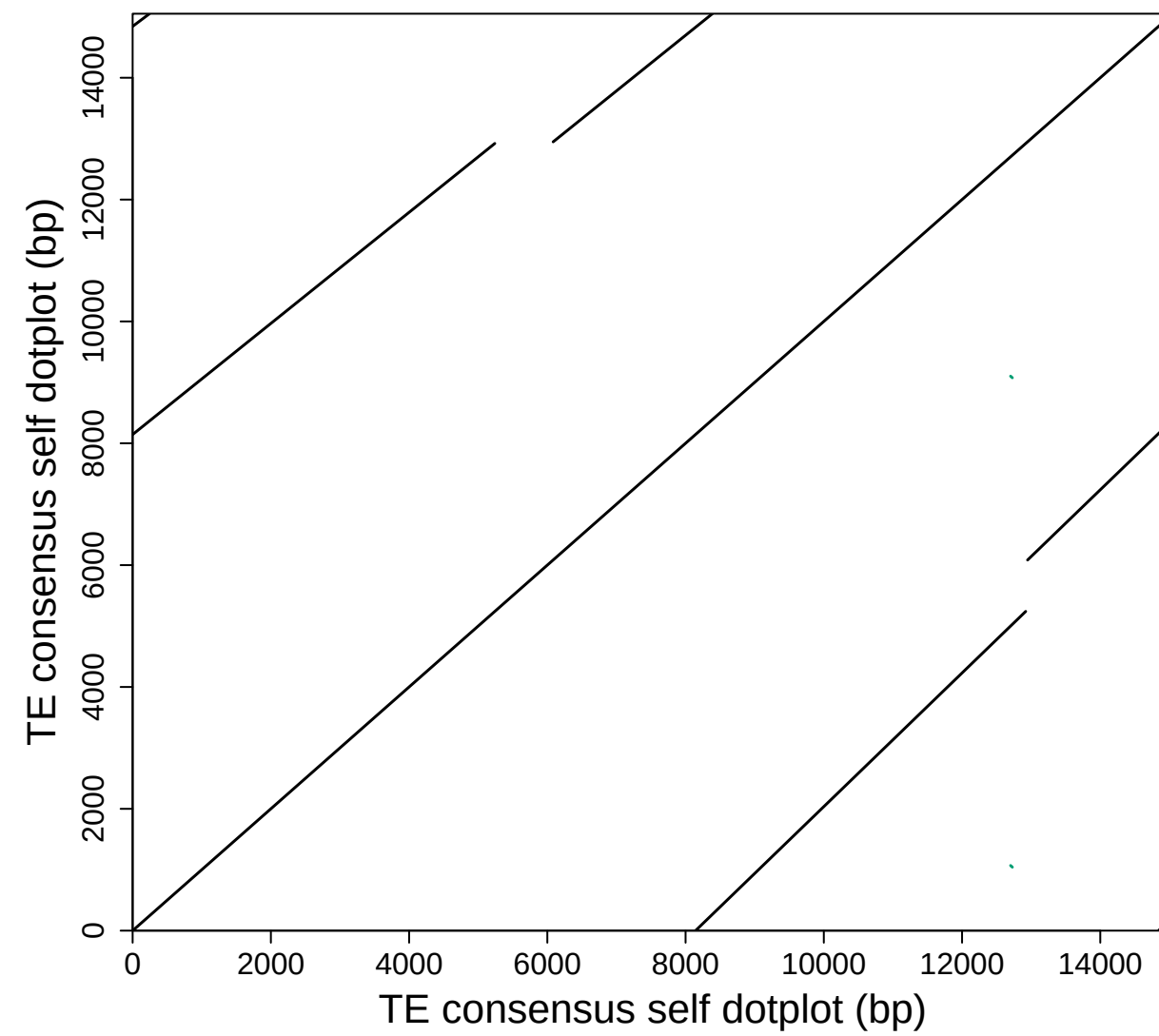
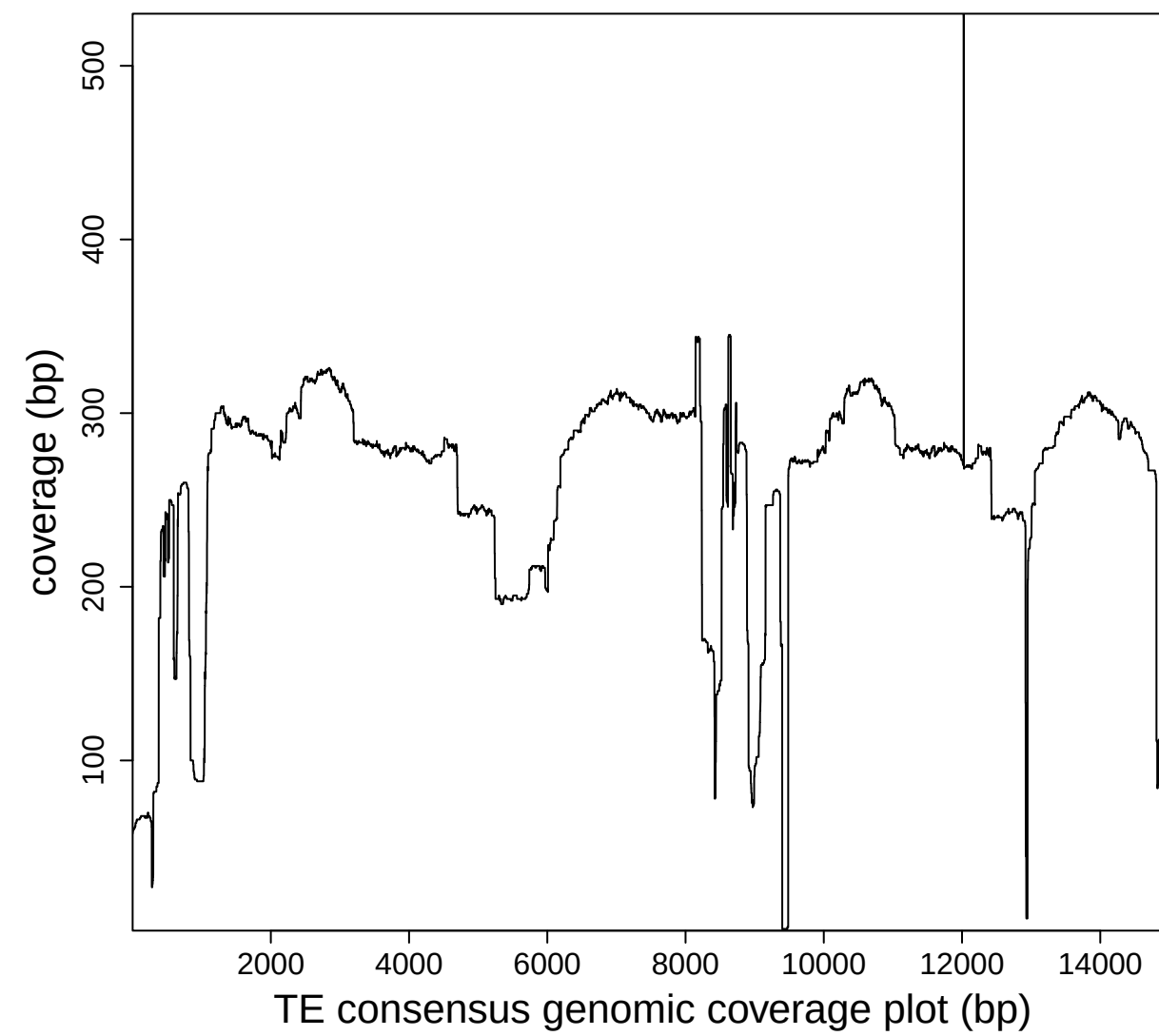
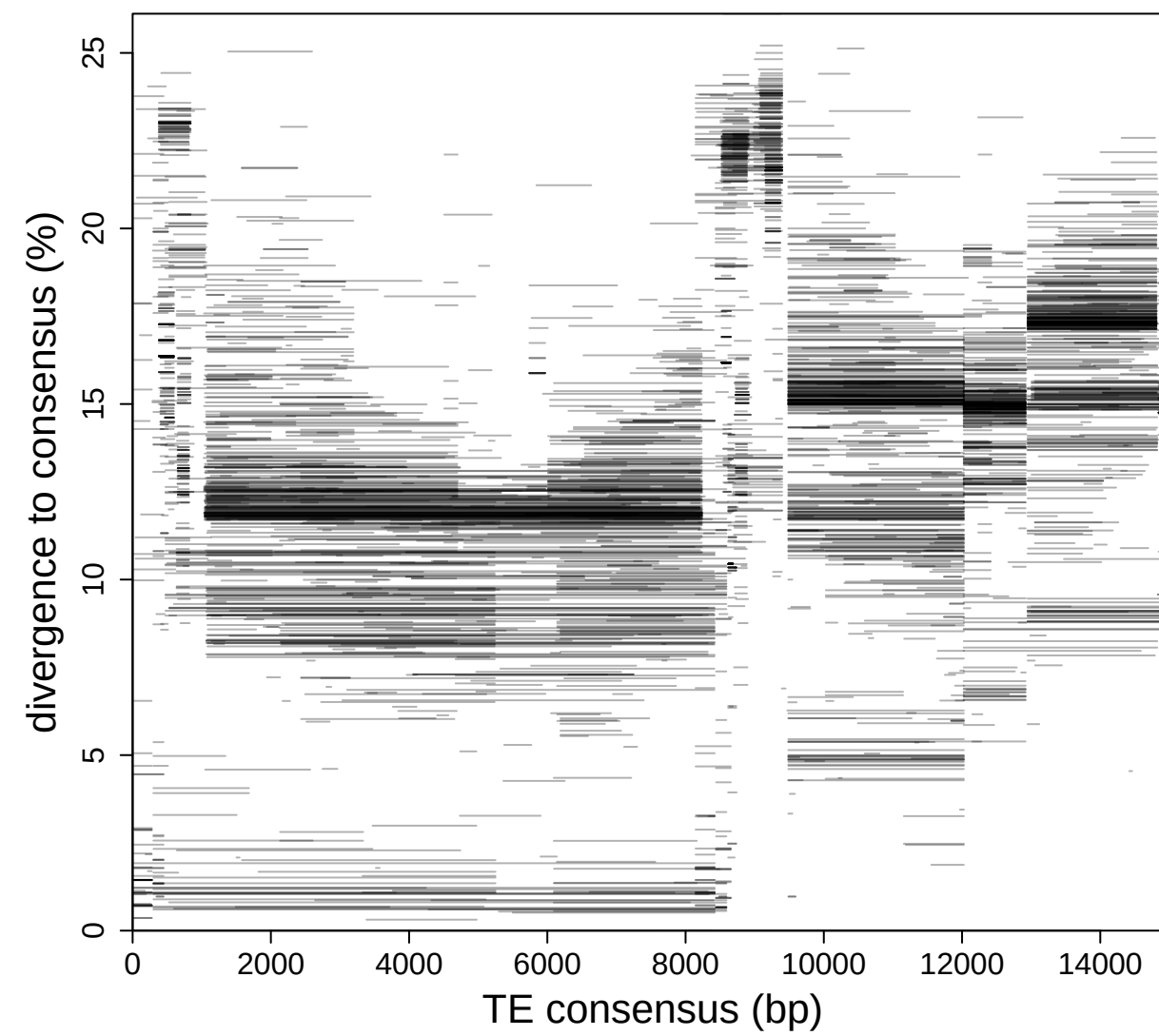
MSA length = 15051





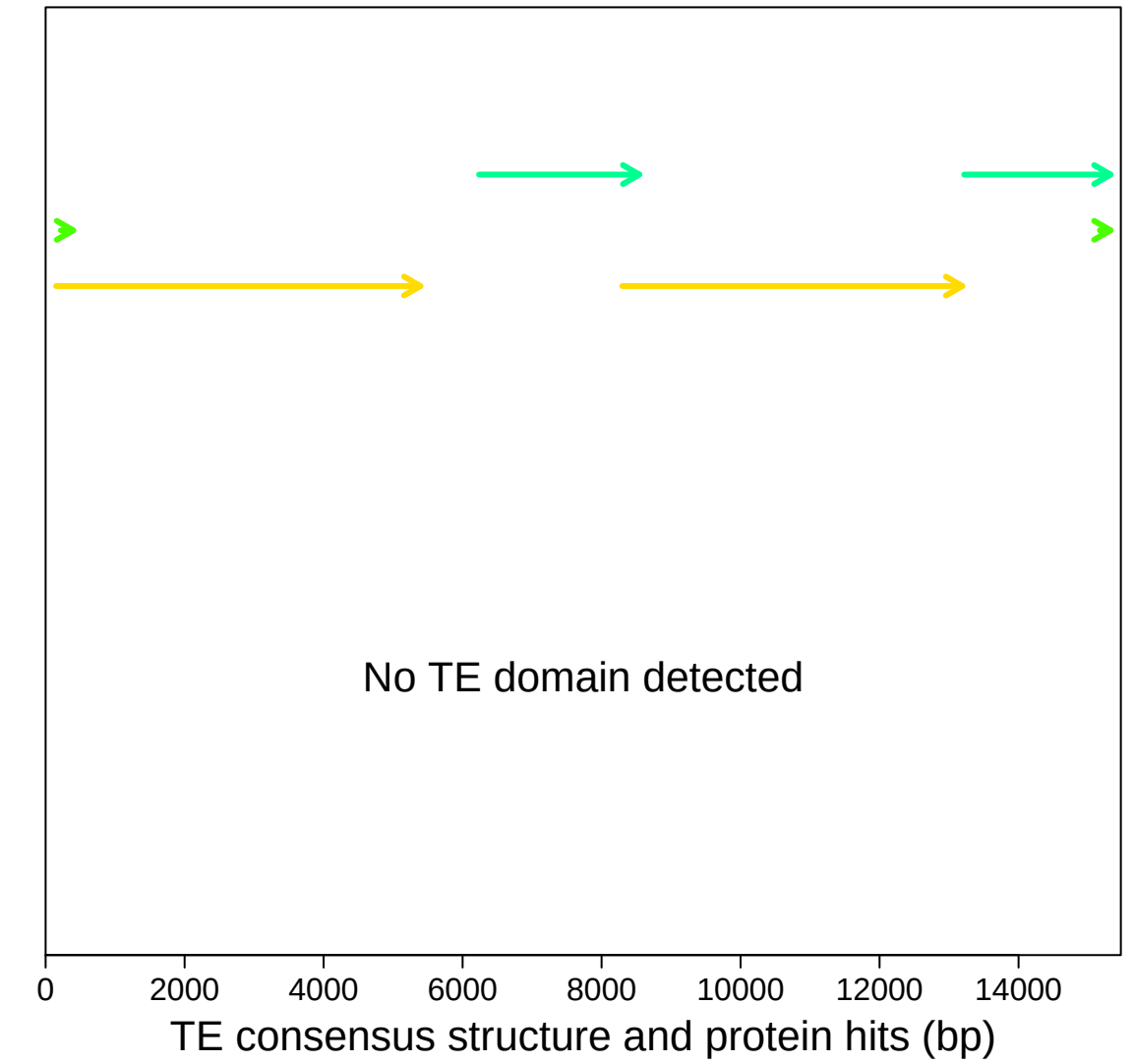
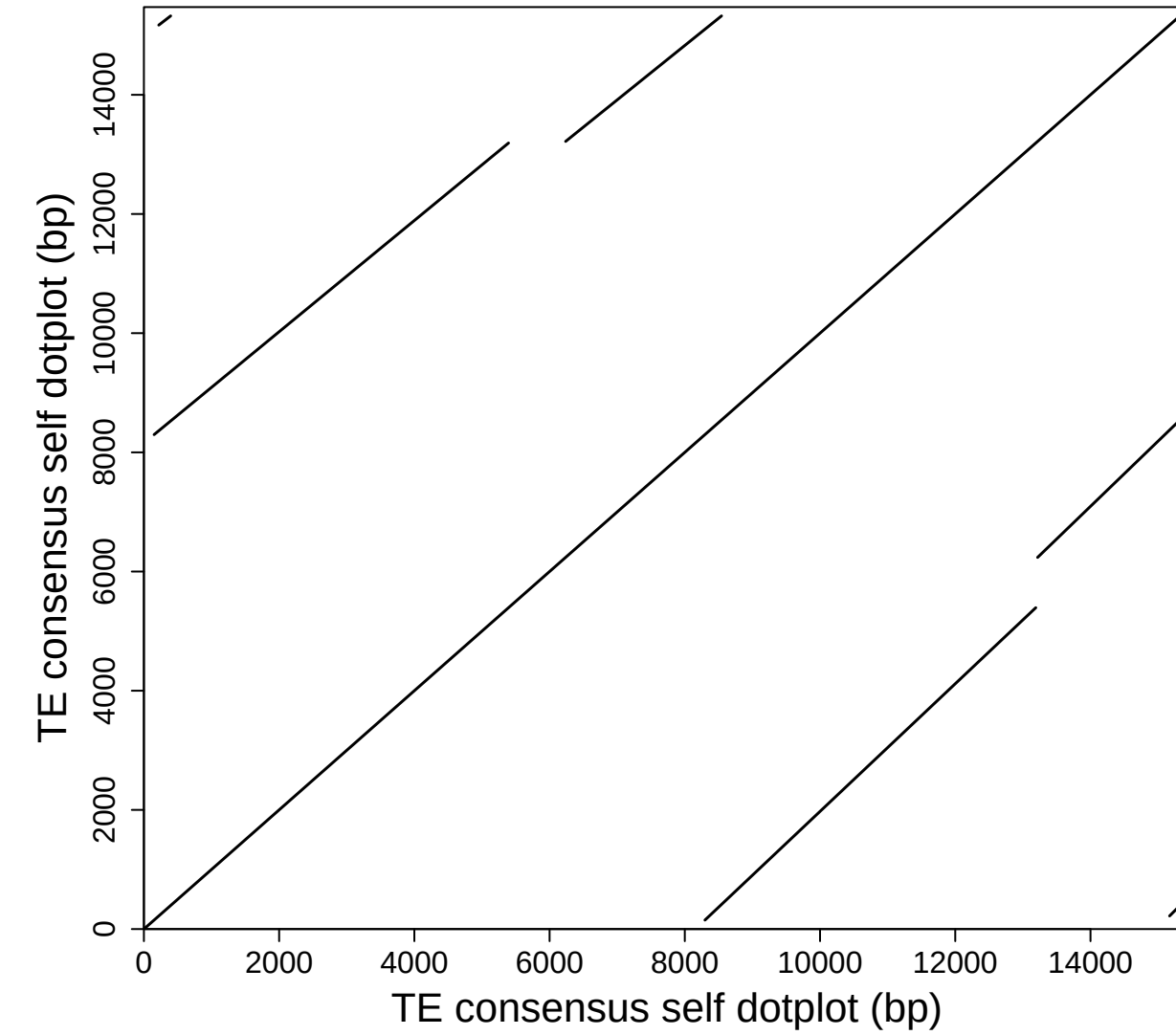
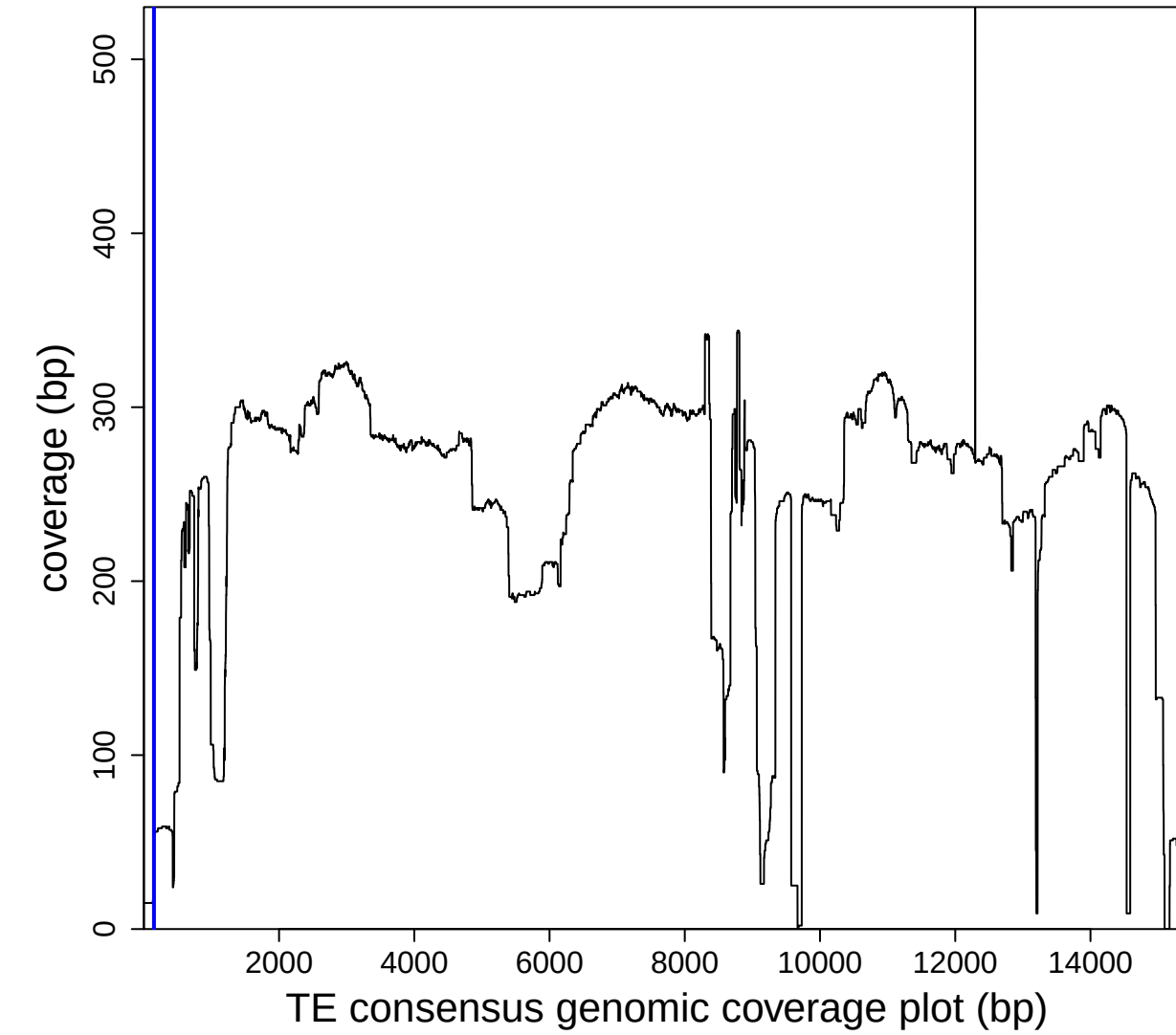
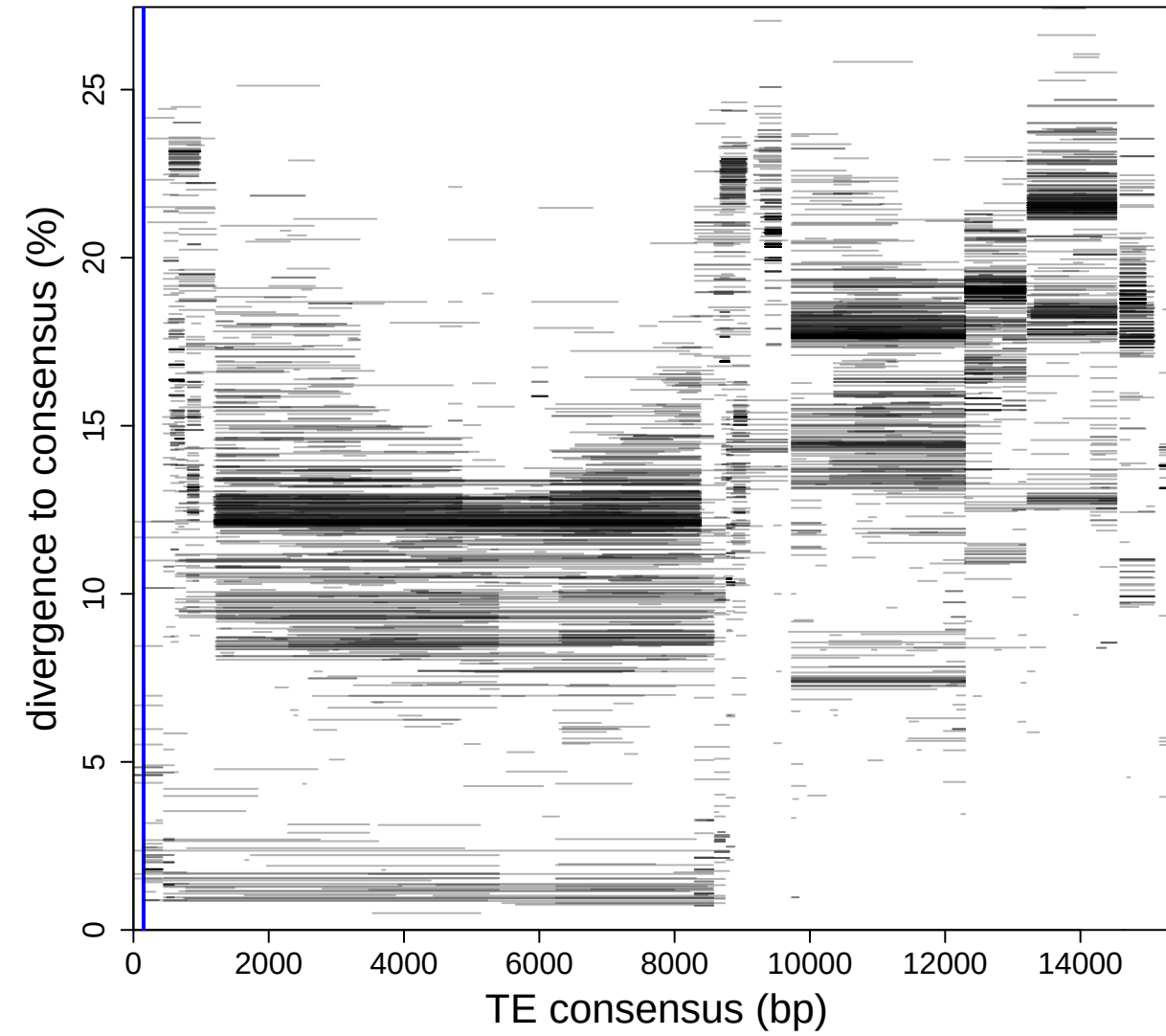
ed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 15051bp; fragments: 3156; full length: 0 (>=13545.9bp)

After TEtrimmer 15051 bp

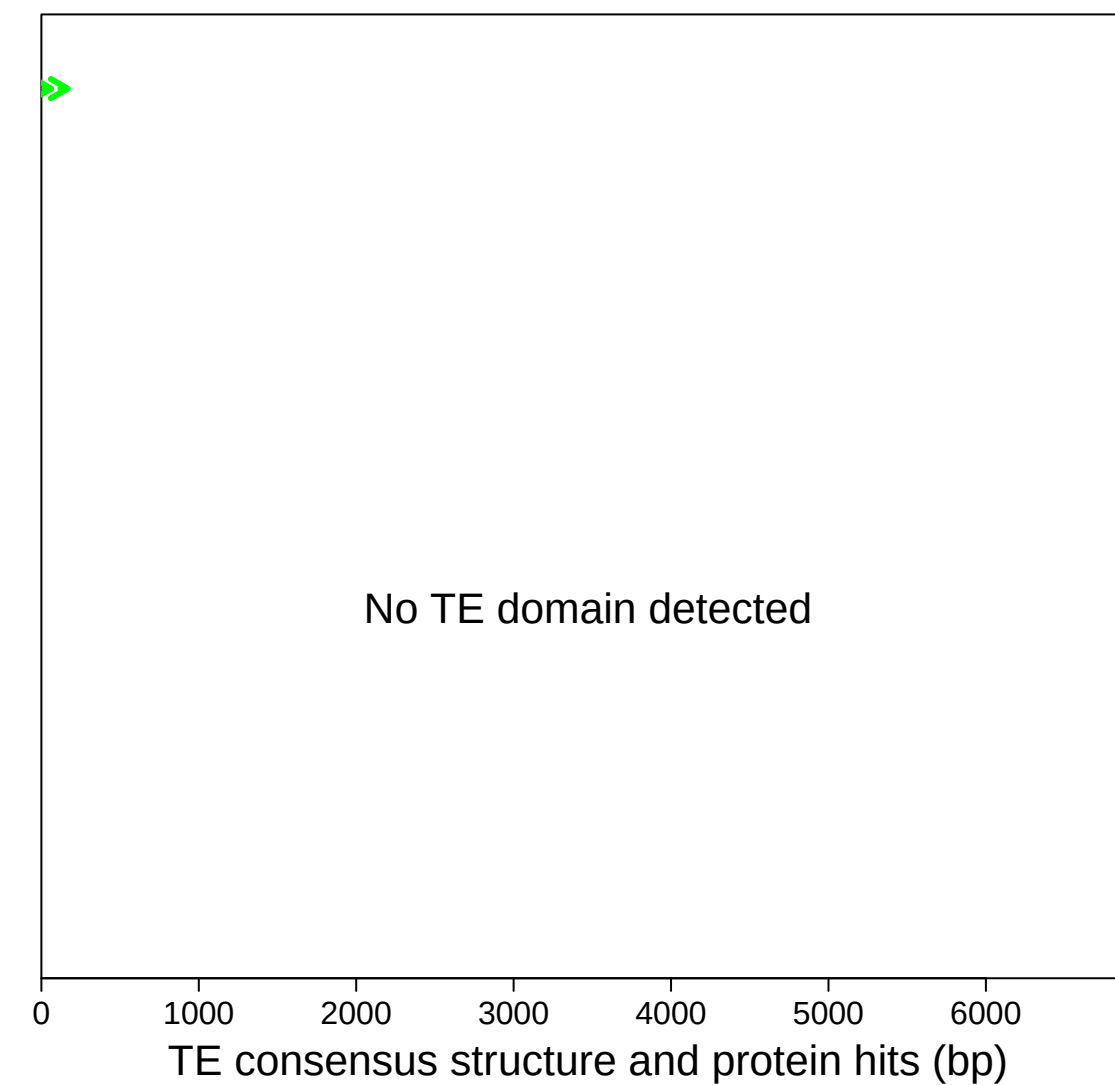
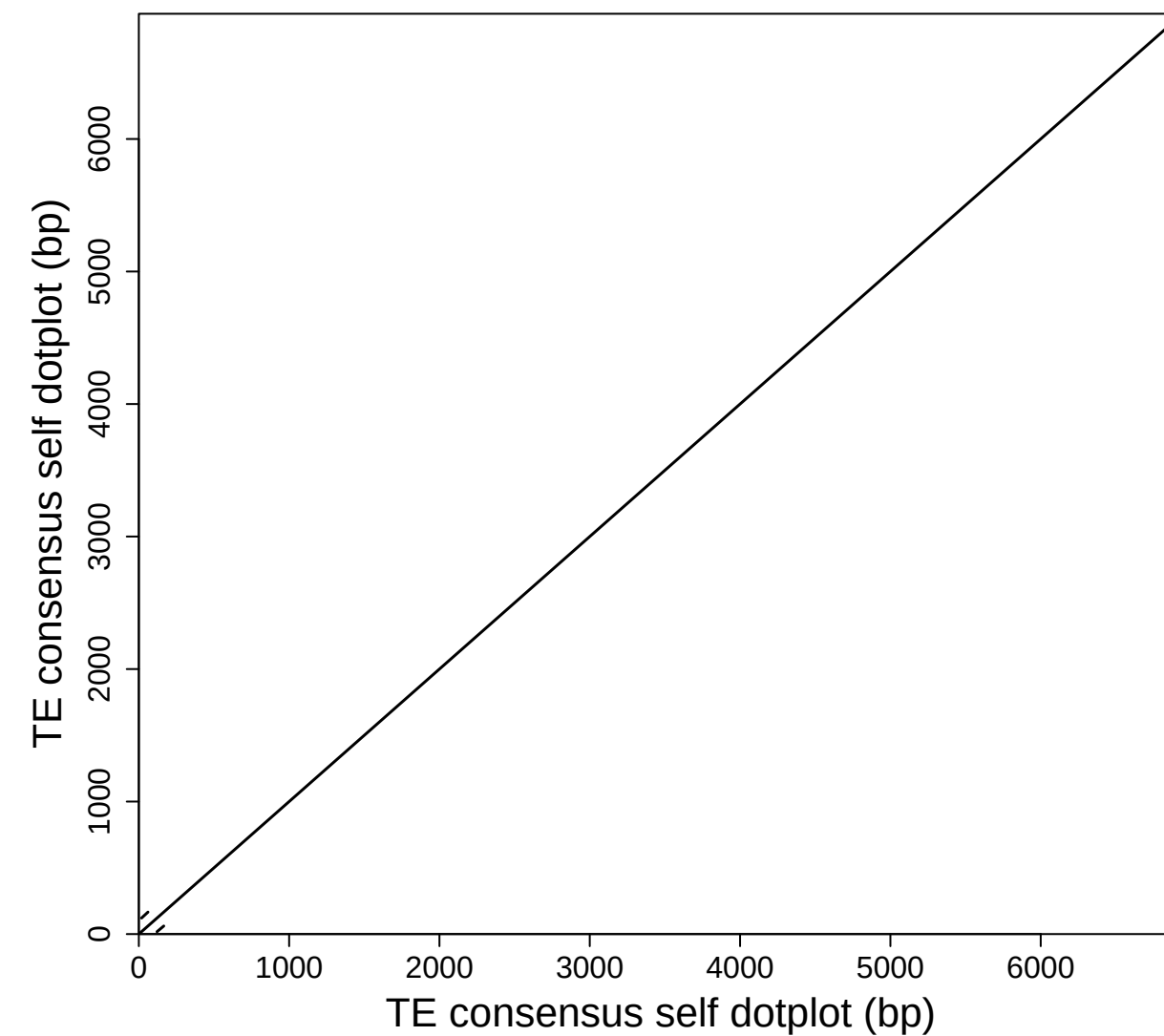
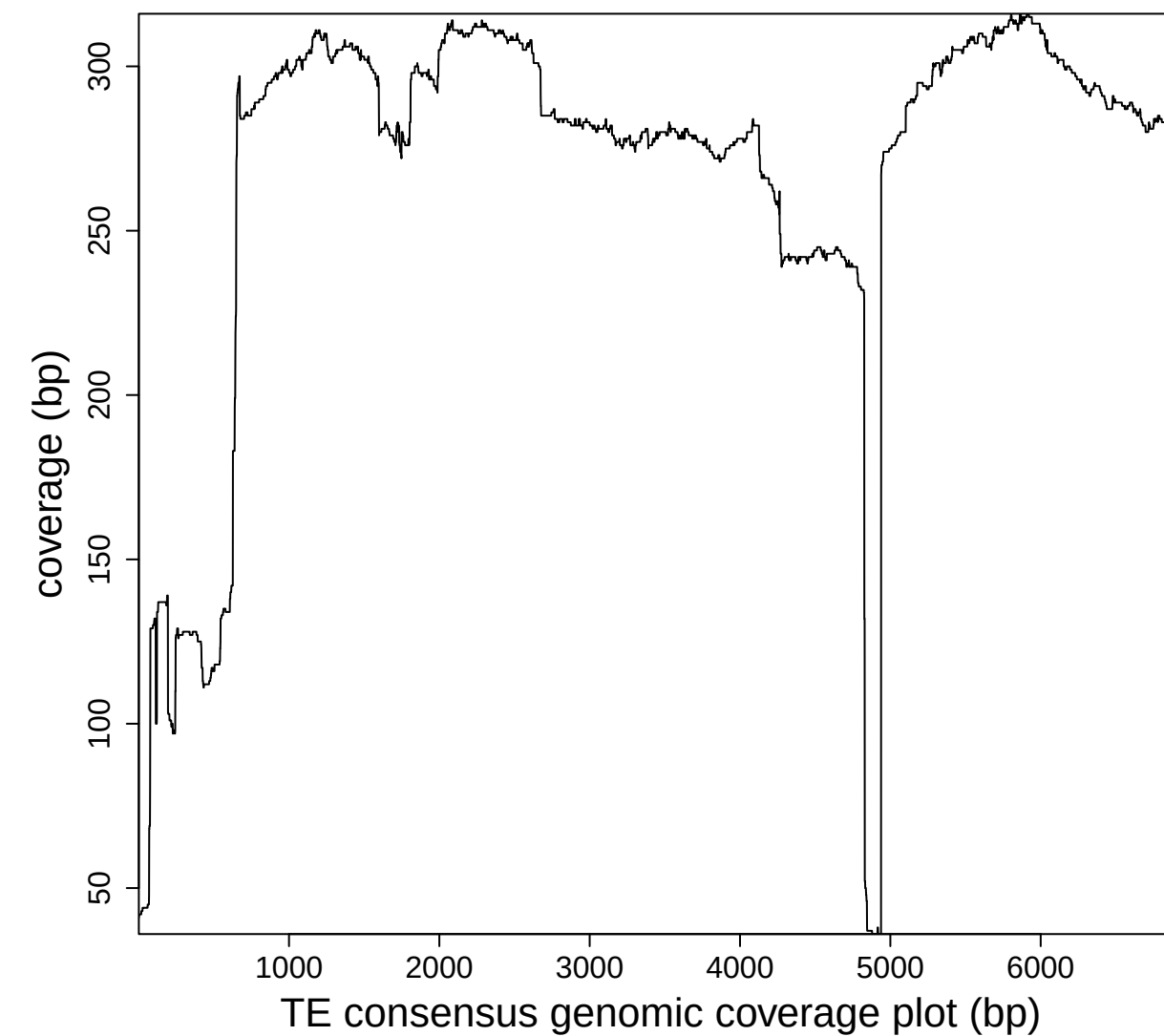
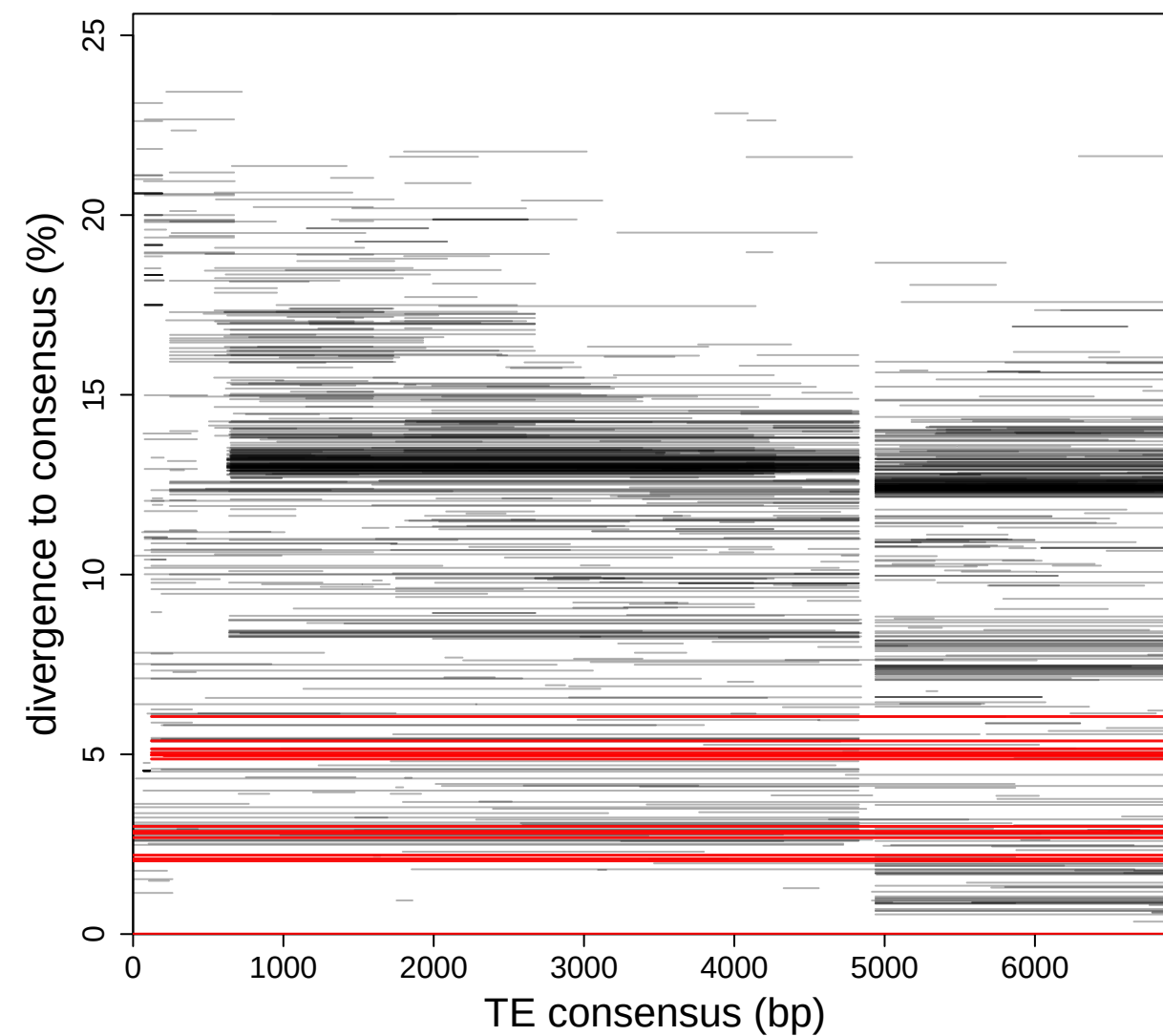


ly_678.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 15472bp; fragments: 3461; full length: 0 (>=13924.8bp)

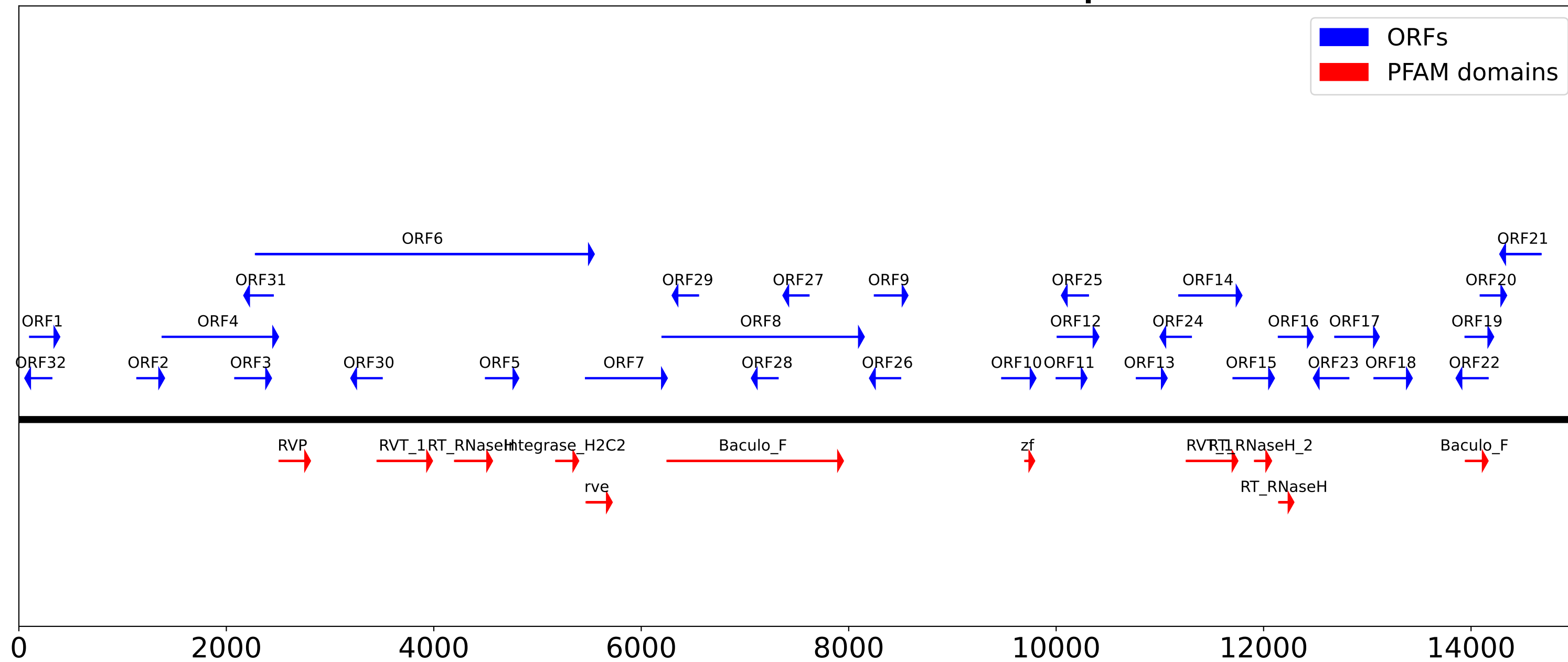
After TEtrimmer Extended plot Blue lines are boundaries



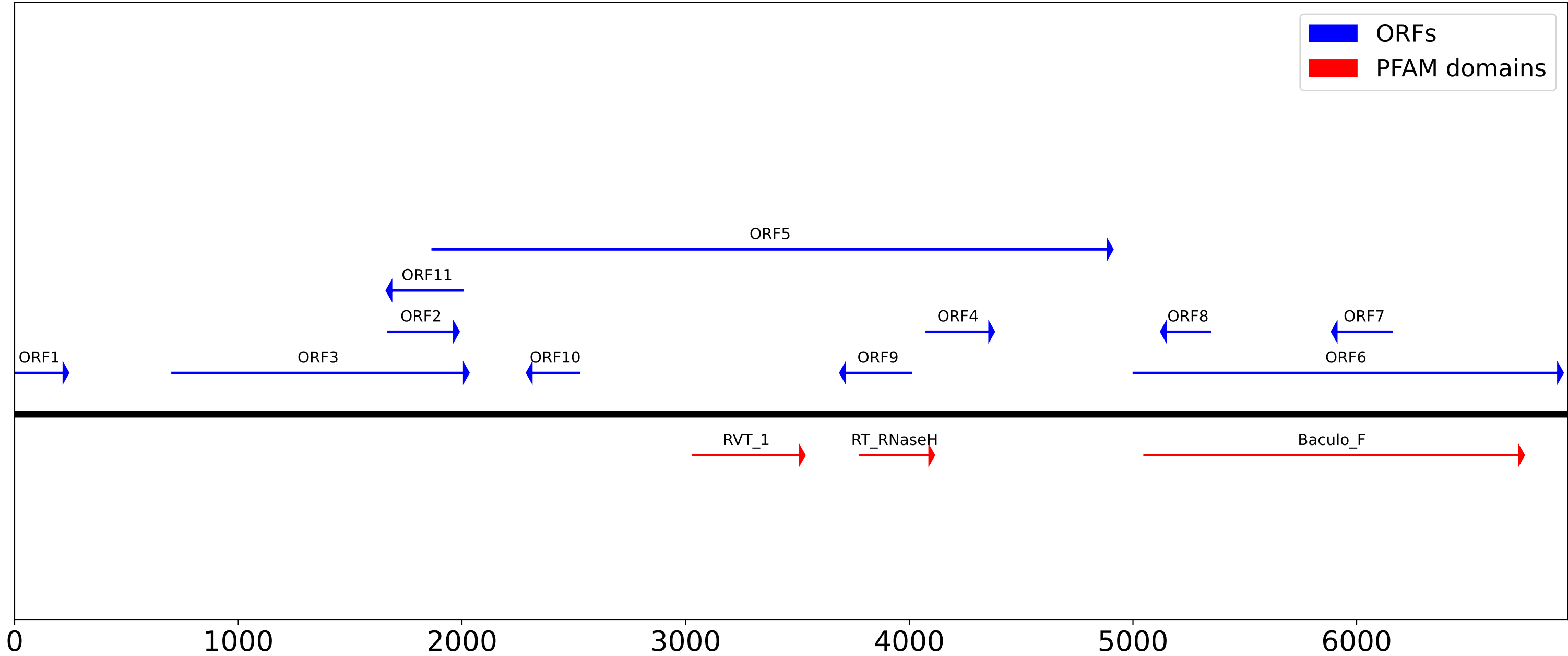
TE: ltr_1_family_678
size: 6945bp; fragments: 1061; full length: 16 (≥ 6250.5 bp)



After TEtrimmer ORF and PFAM domain plot



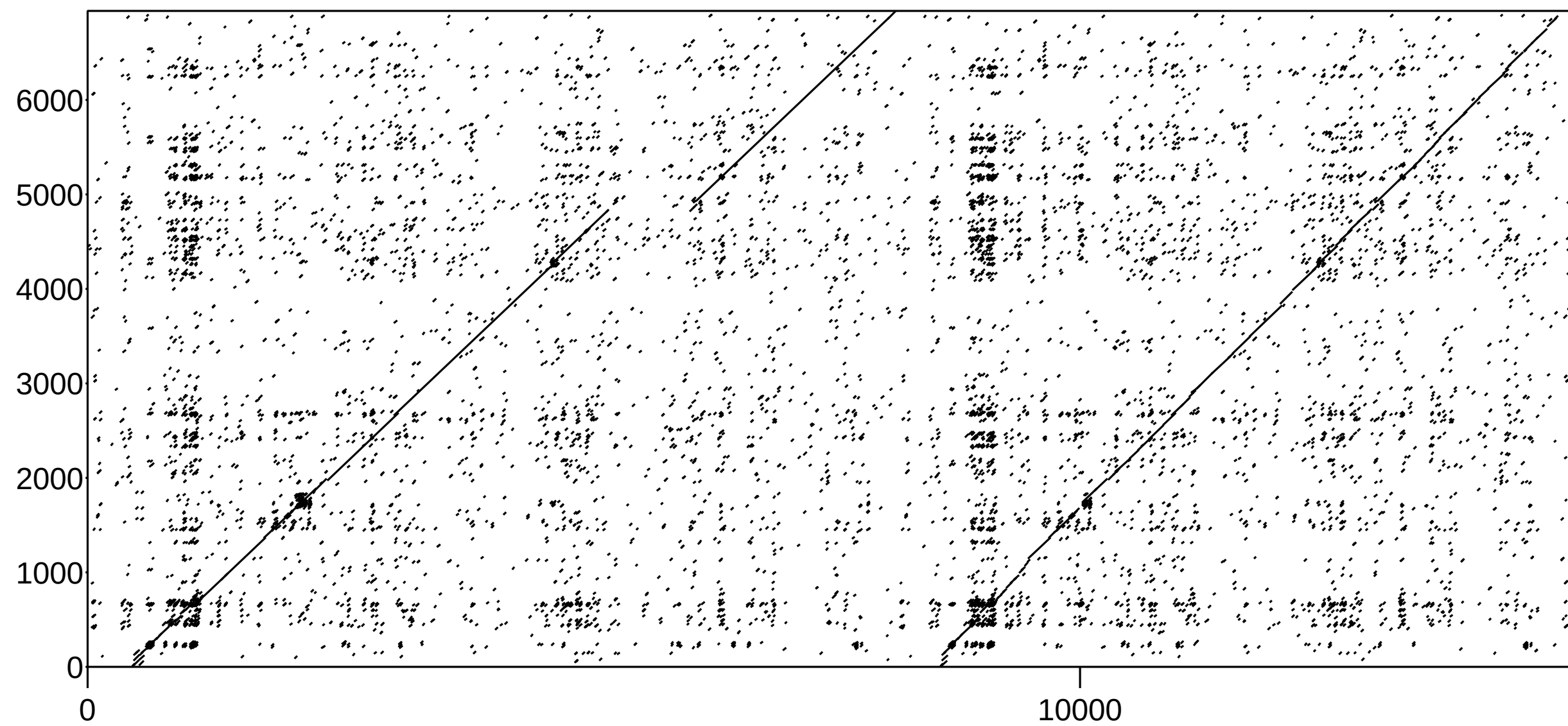
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)