

Start Crop Point

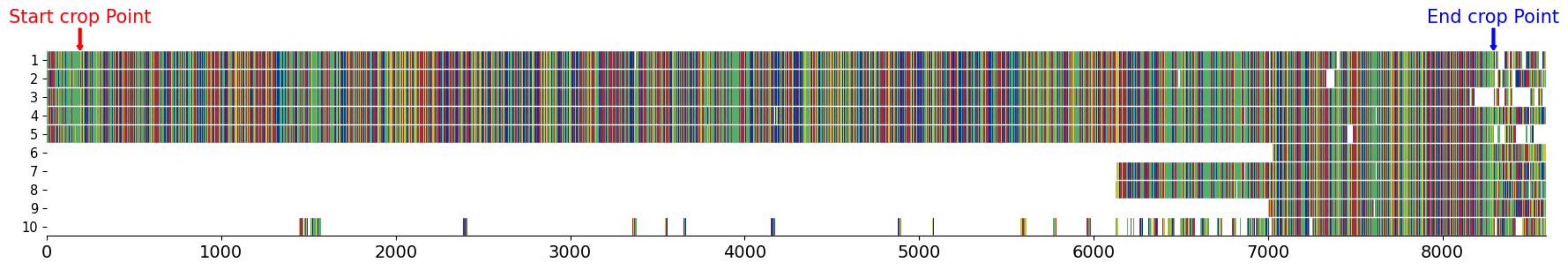


End Crop Point



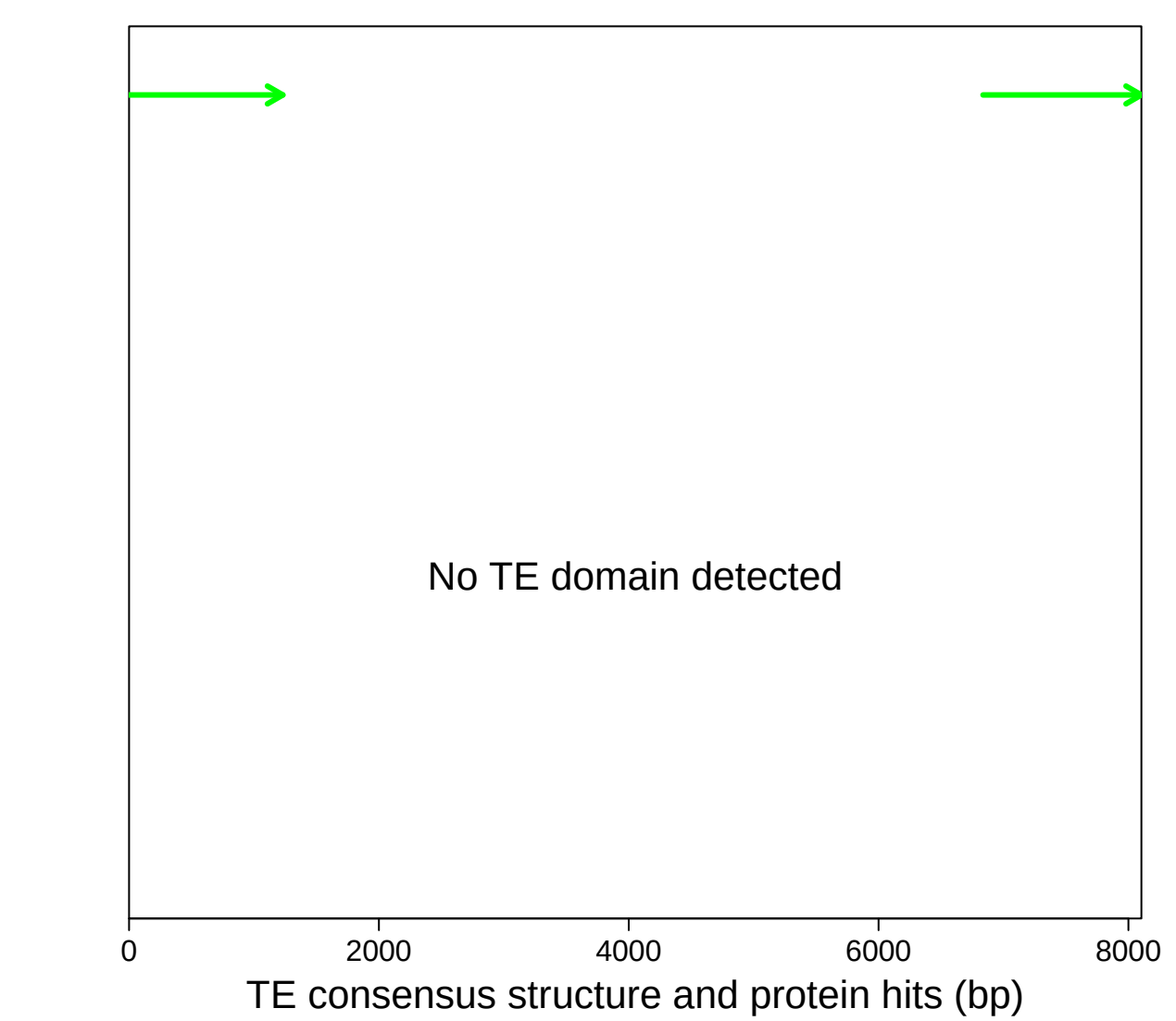
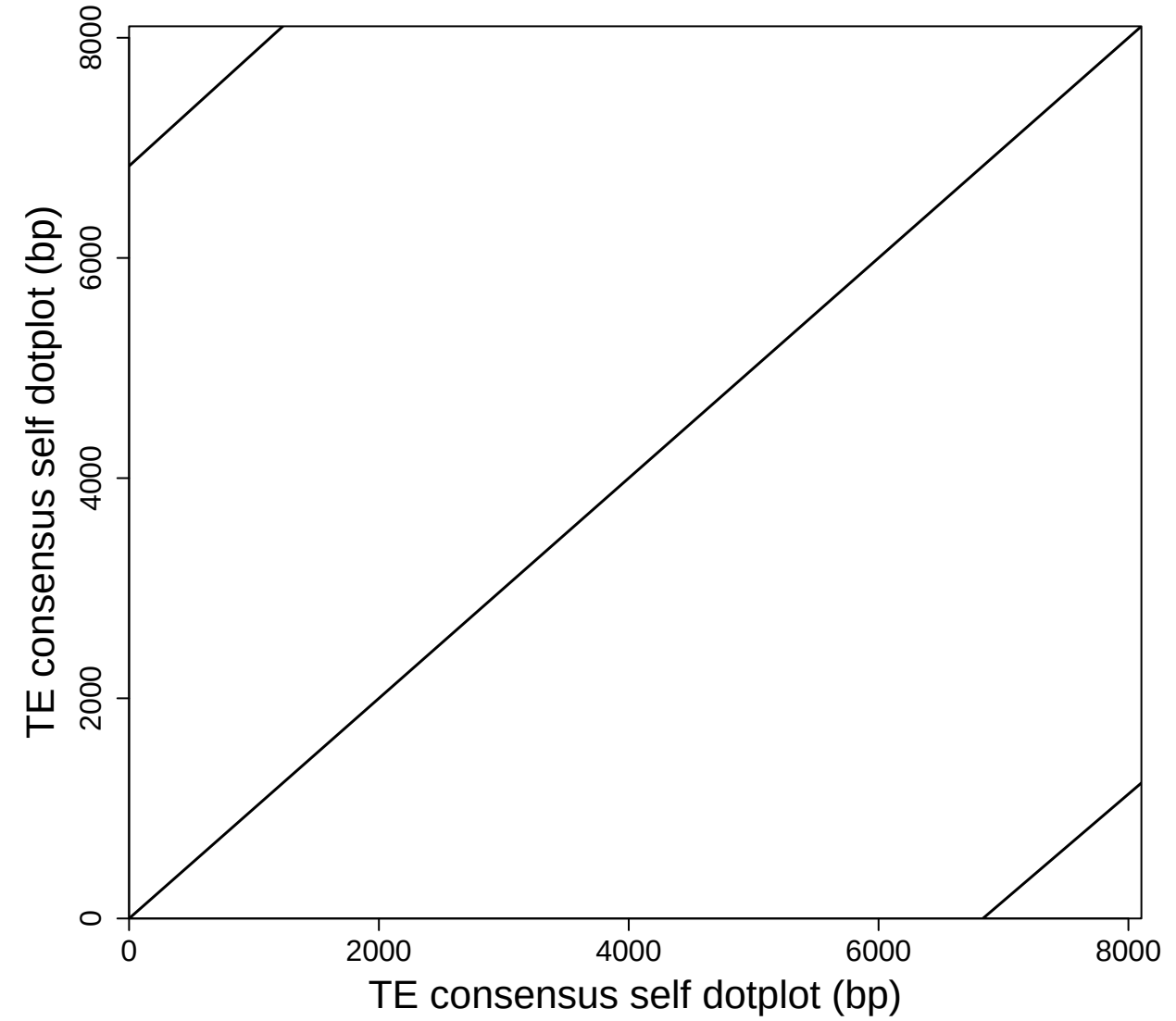
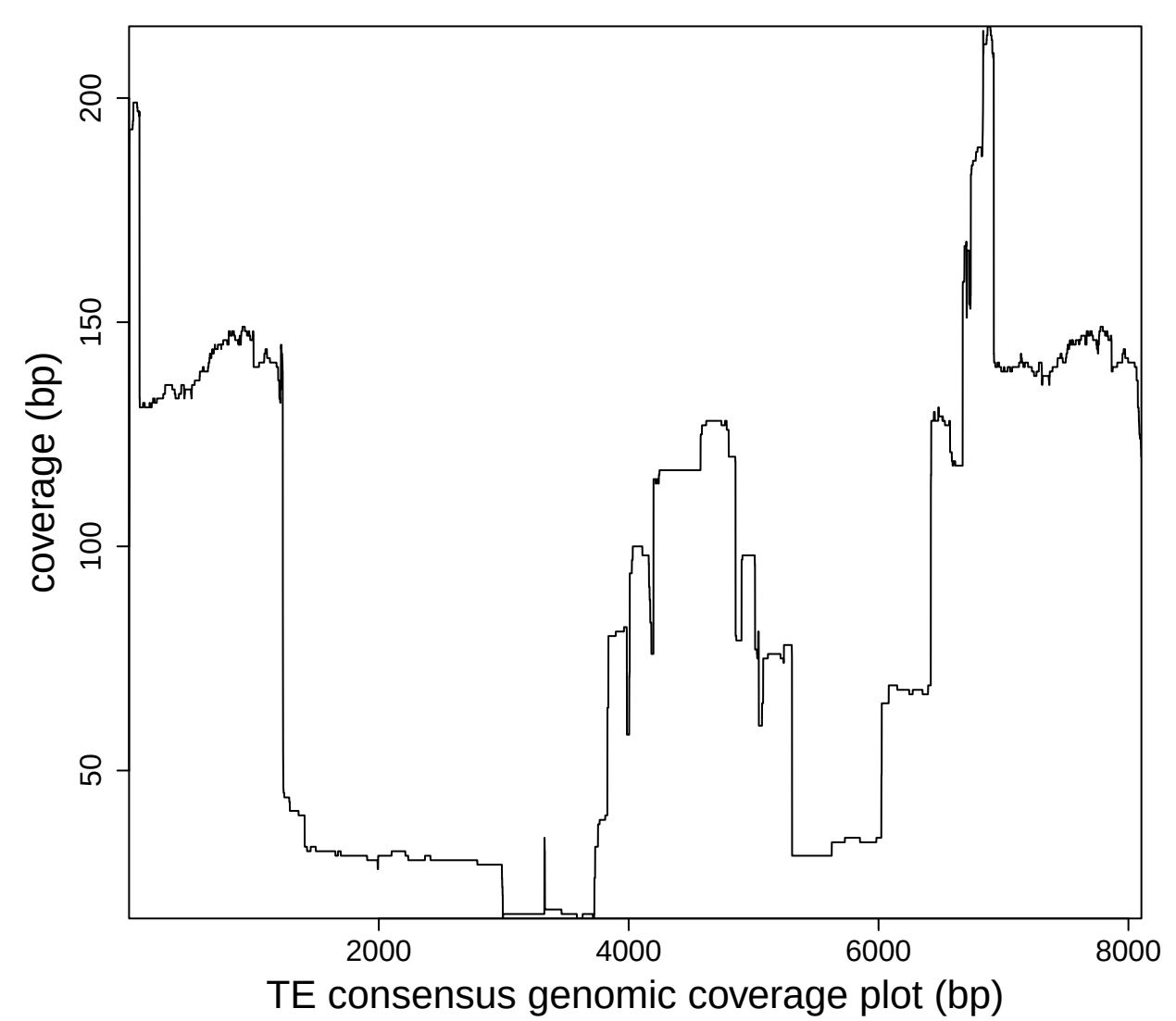
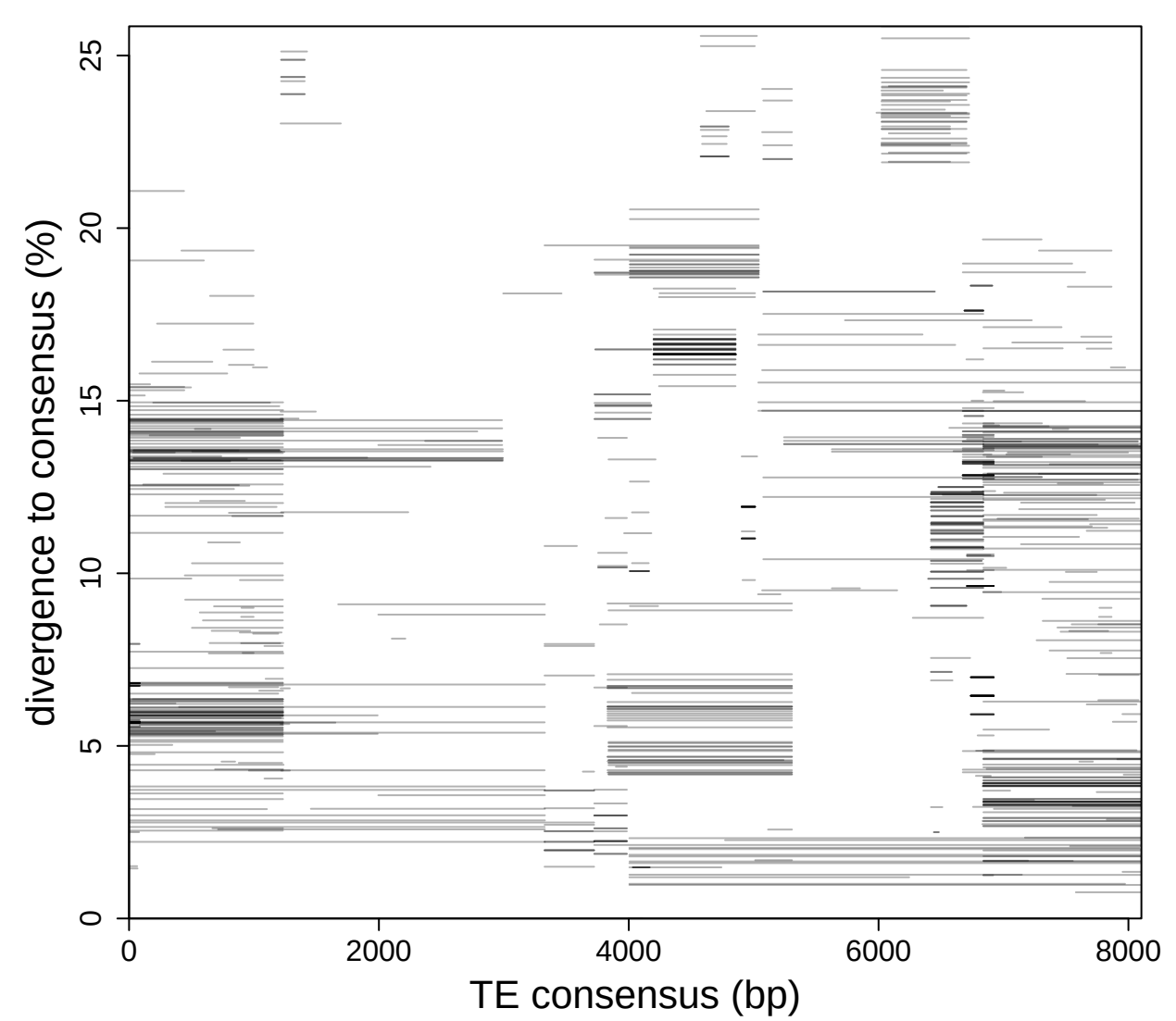
MSA length = 8104





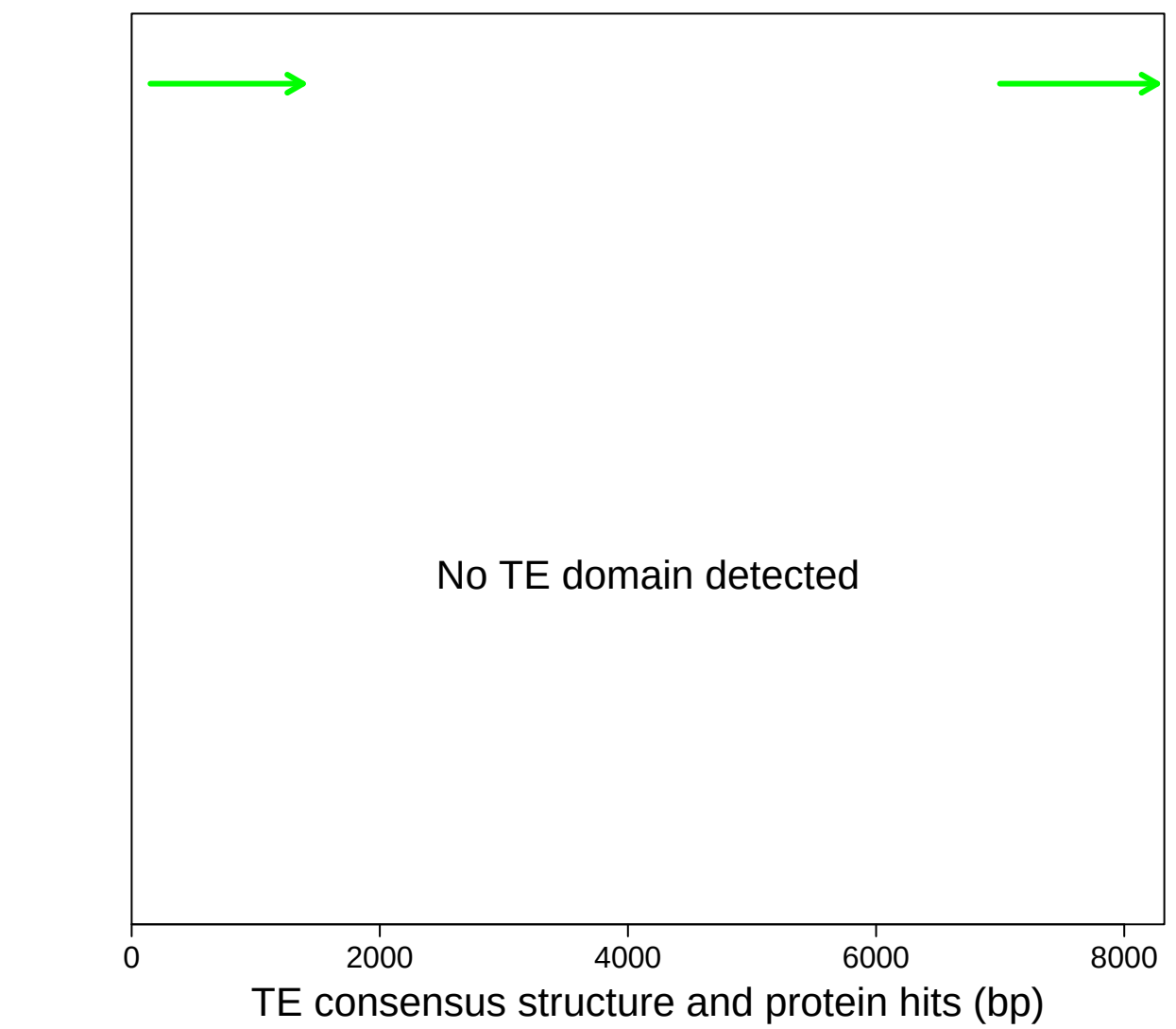
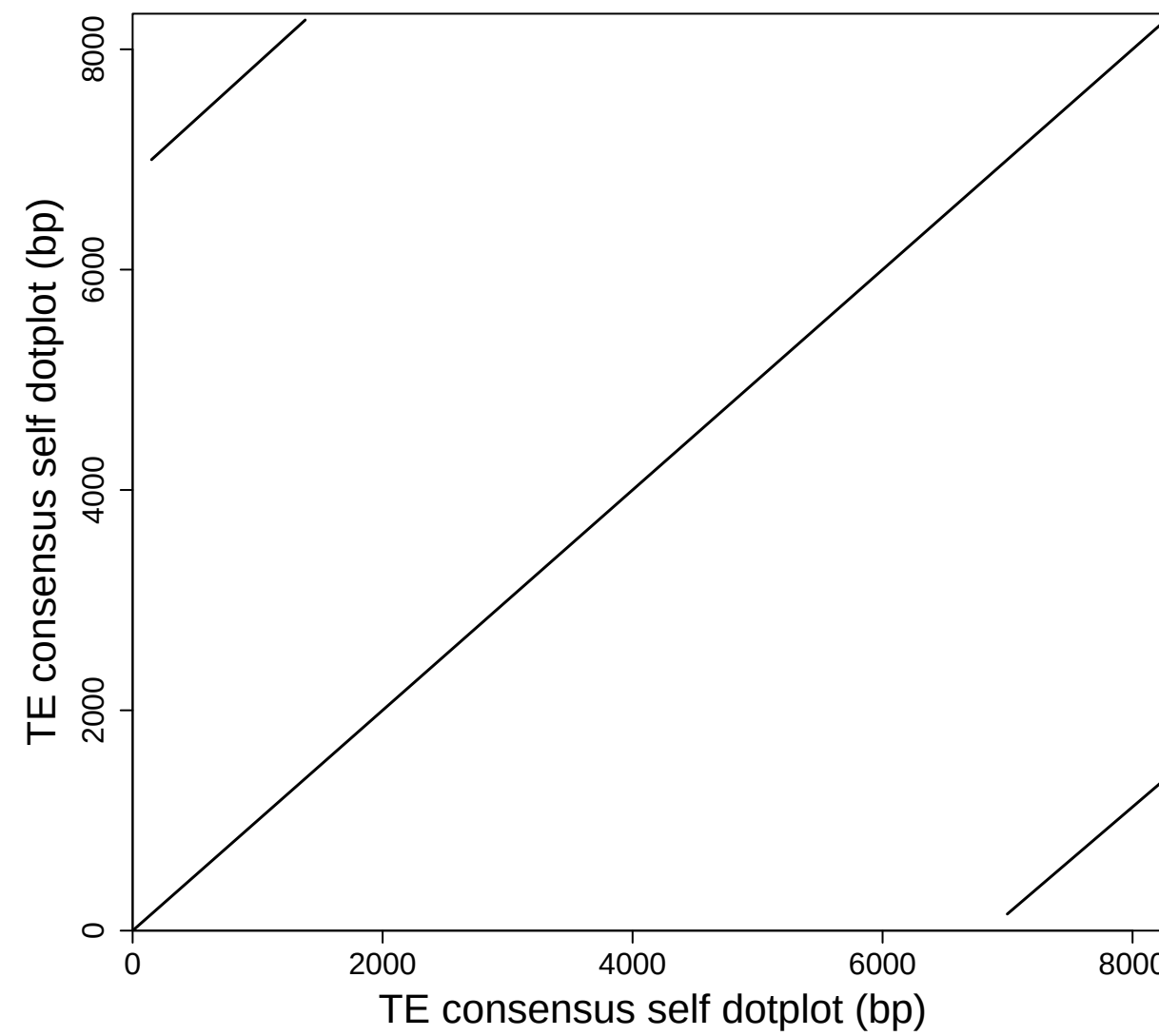
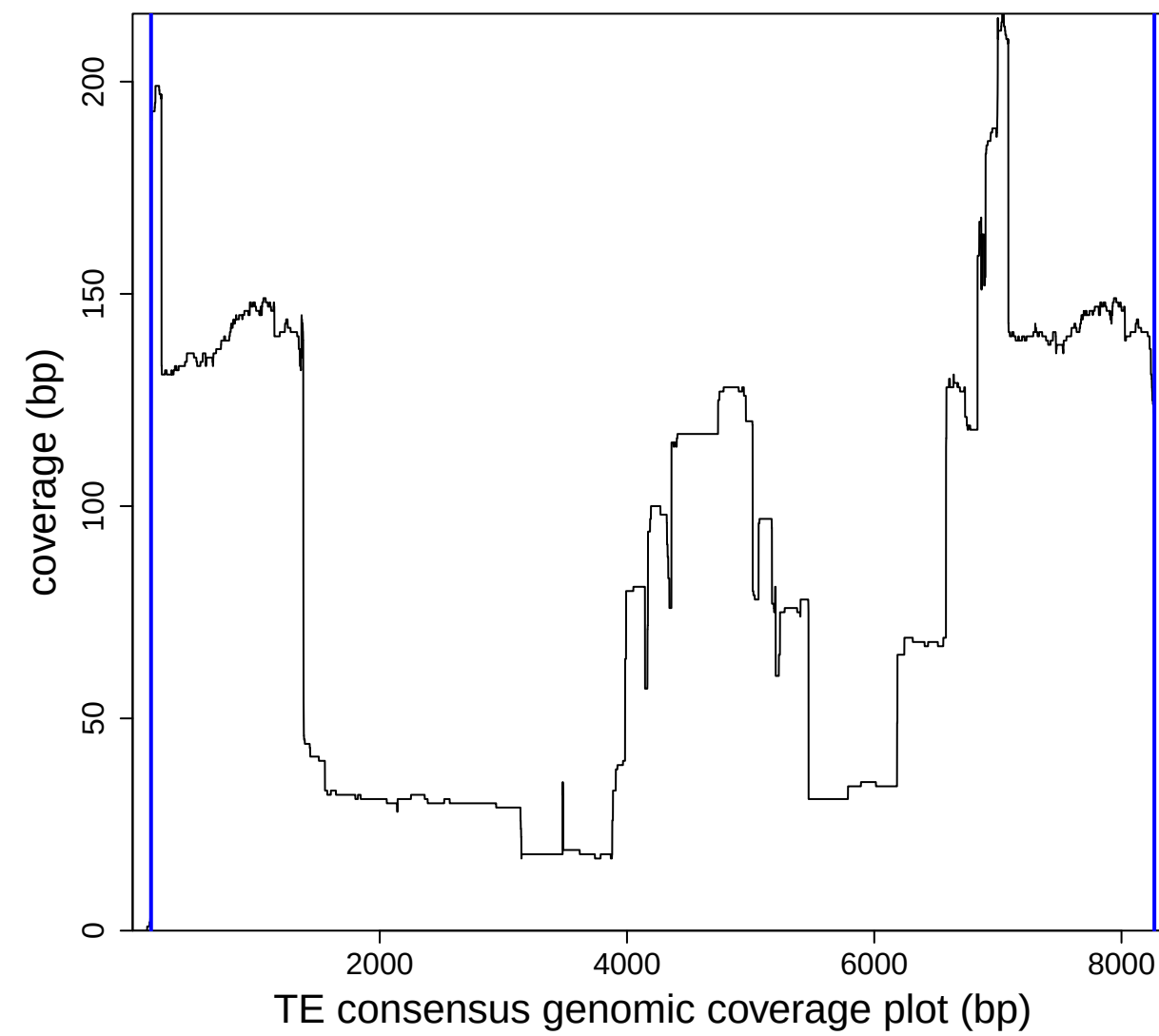
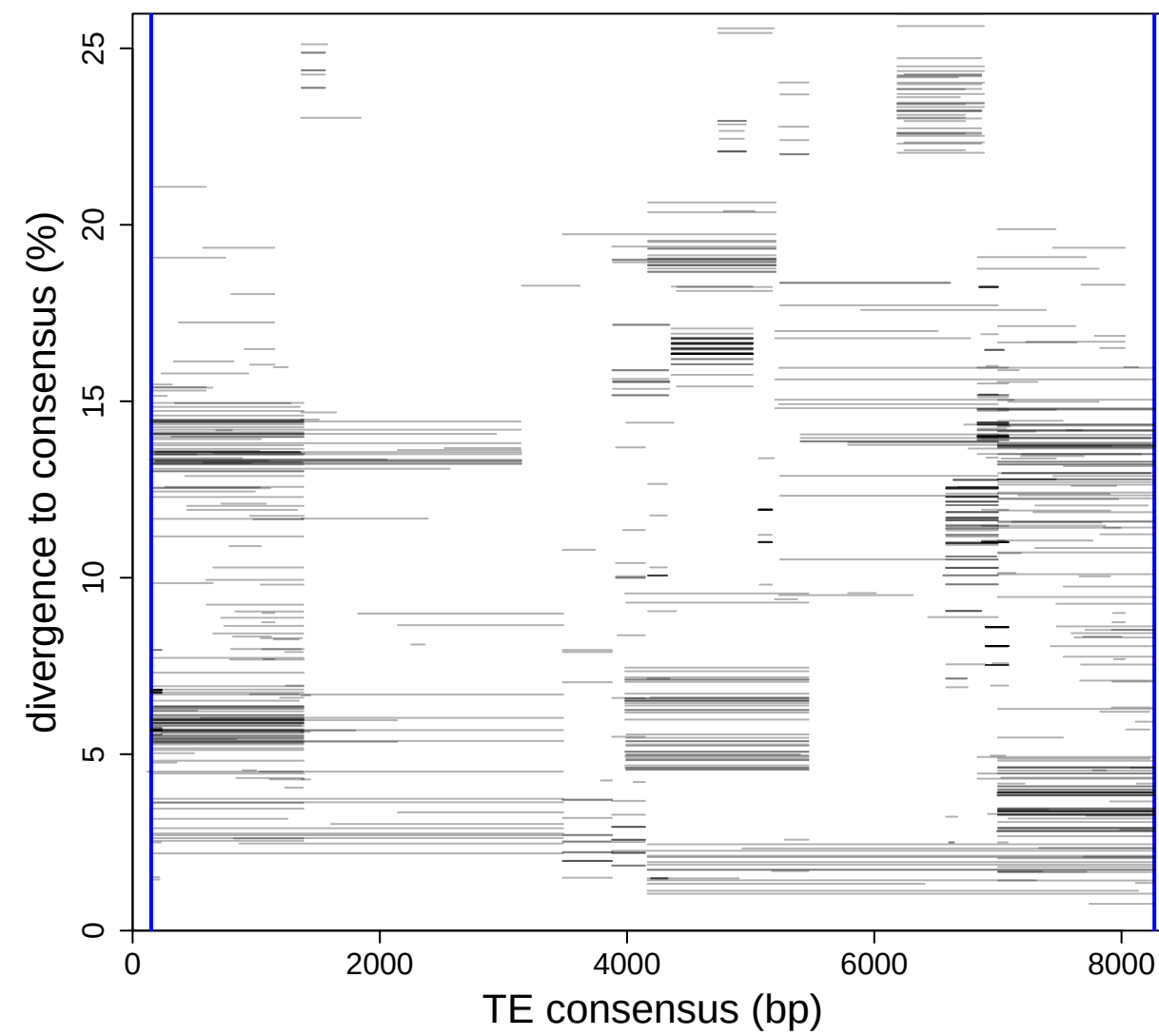
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 8104bp; fragments: 883; full length: 0 (>=7293.6bp)

After TETrimmer 8104 bp



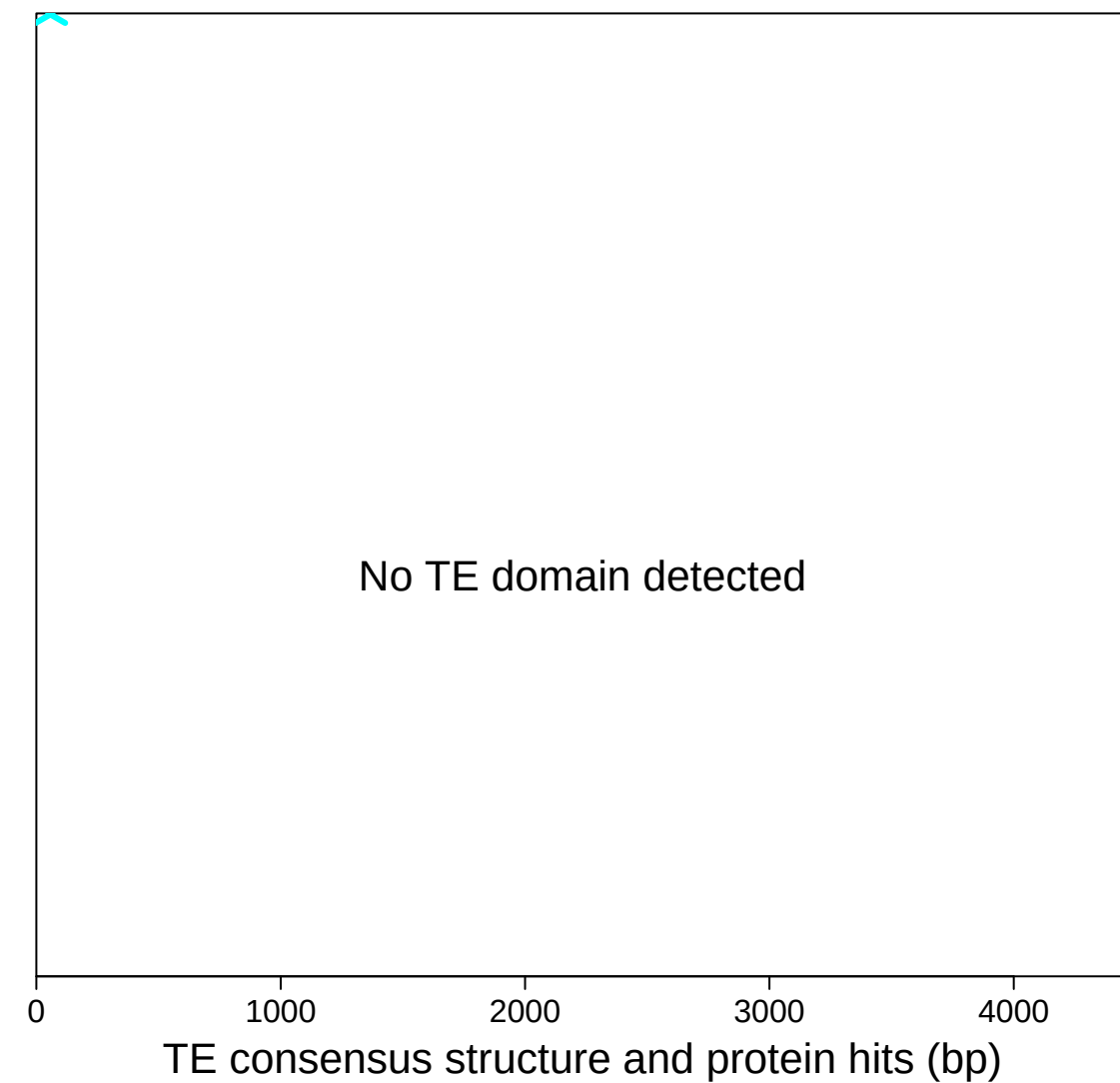
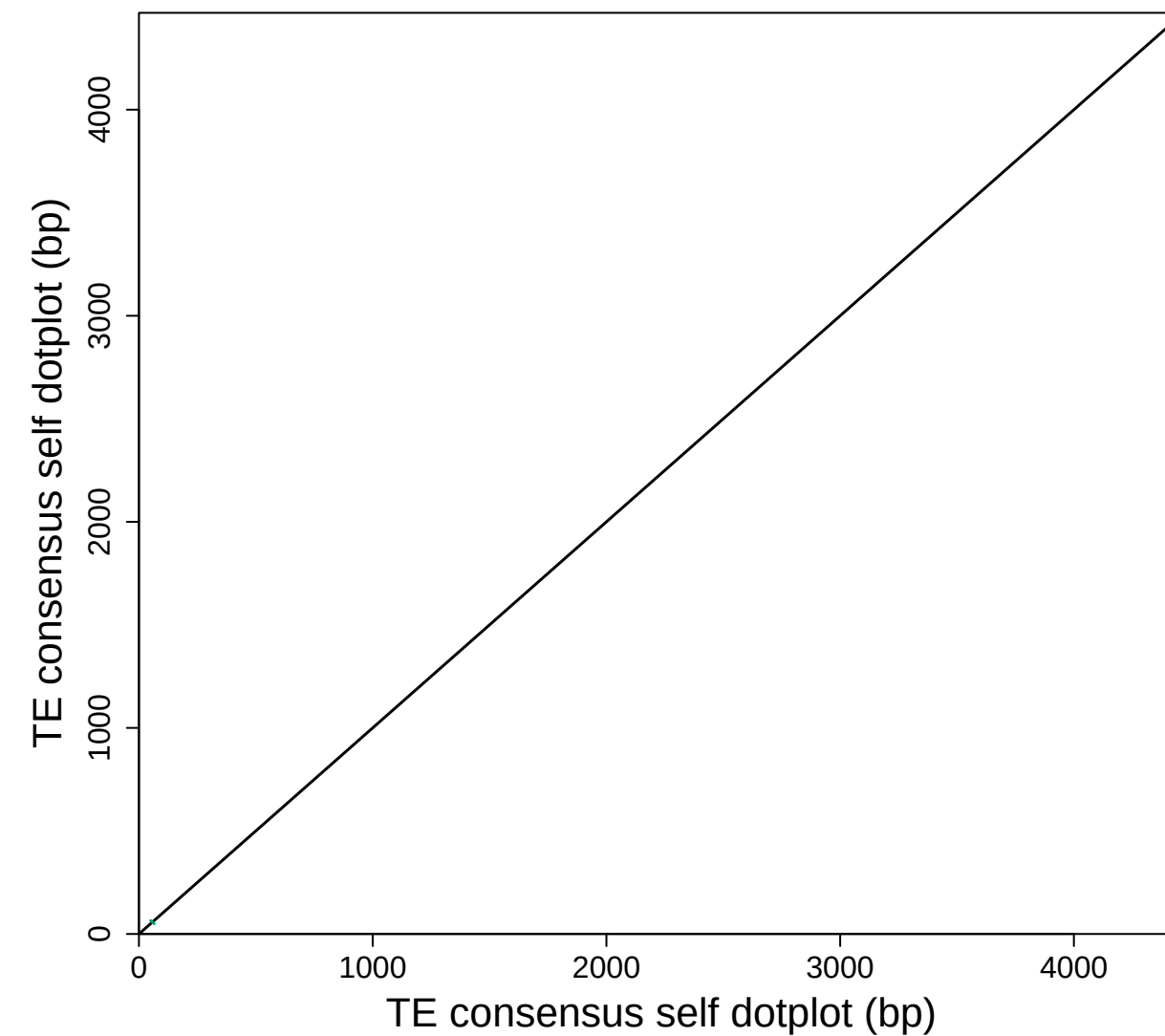
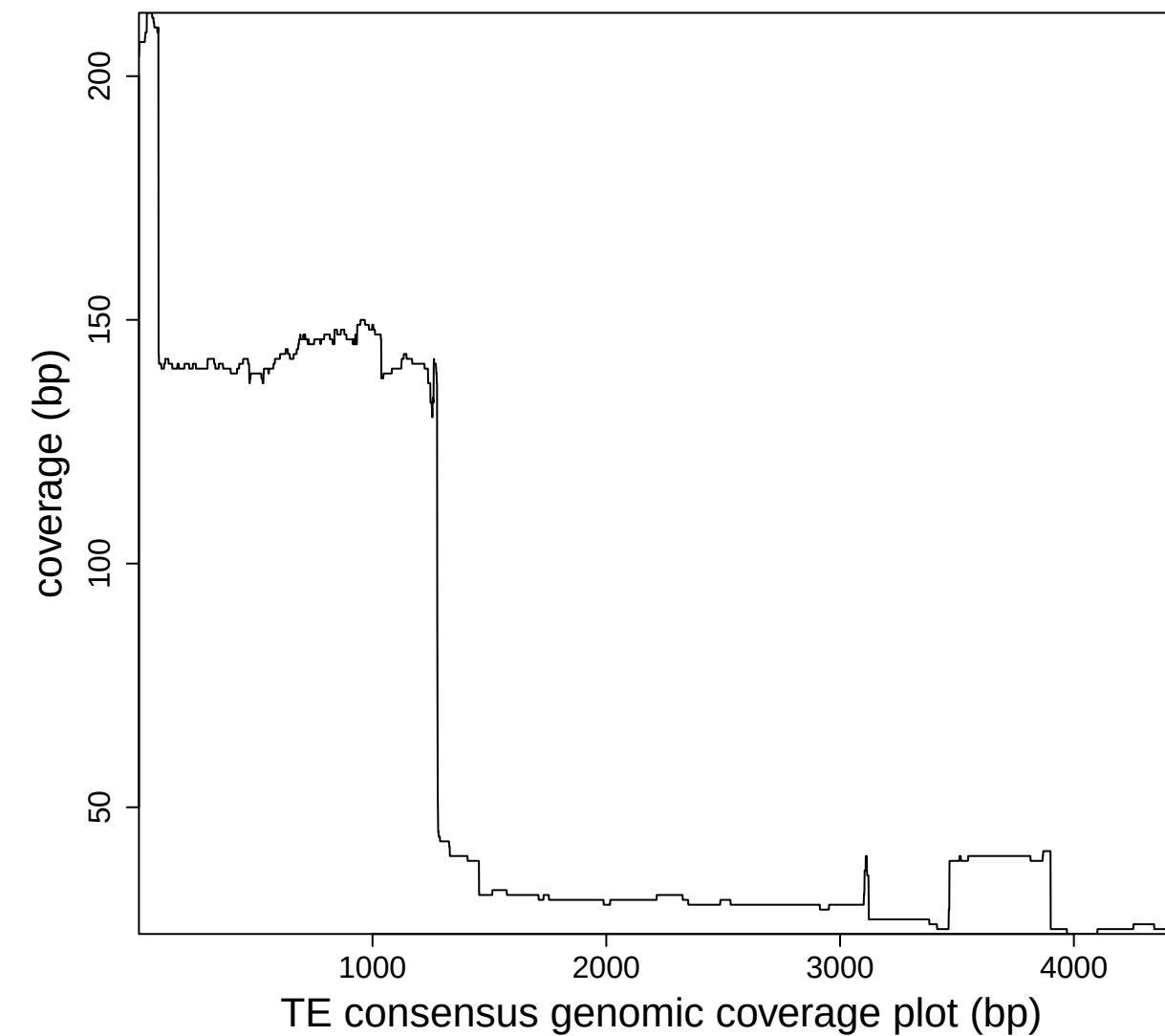
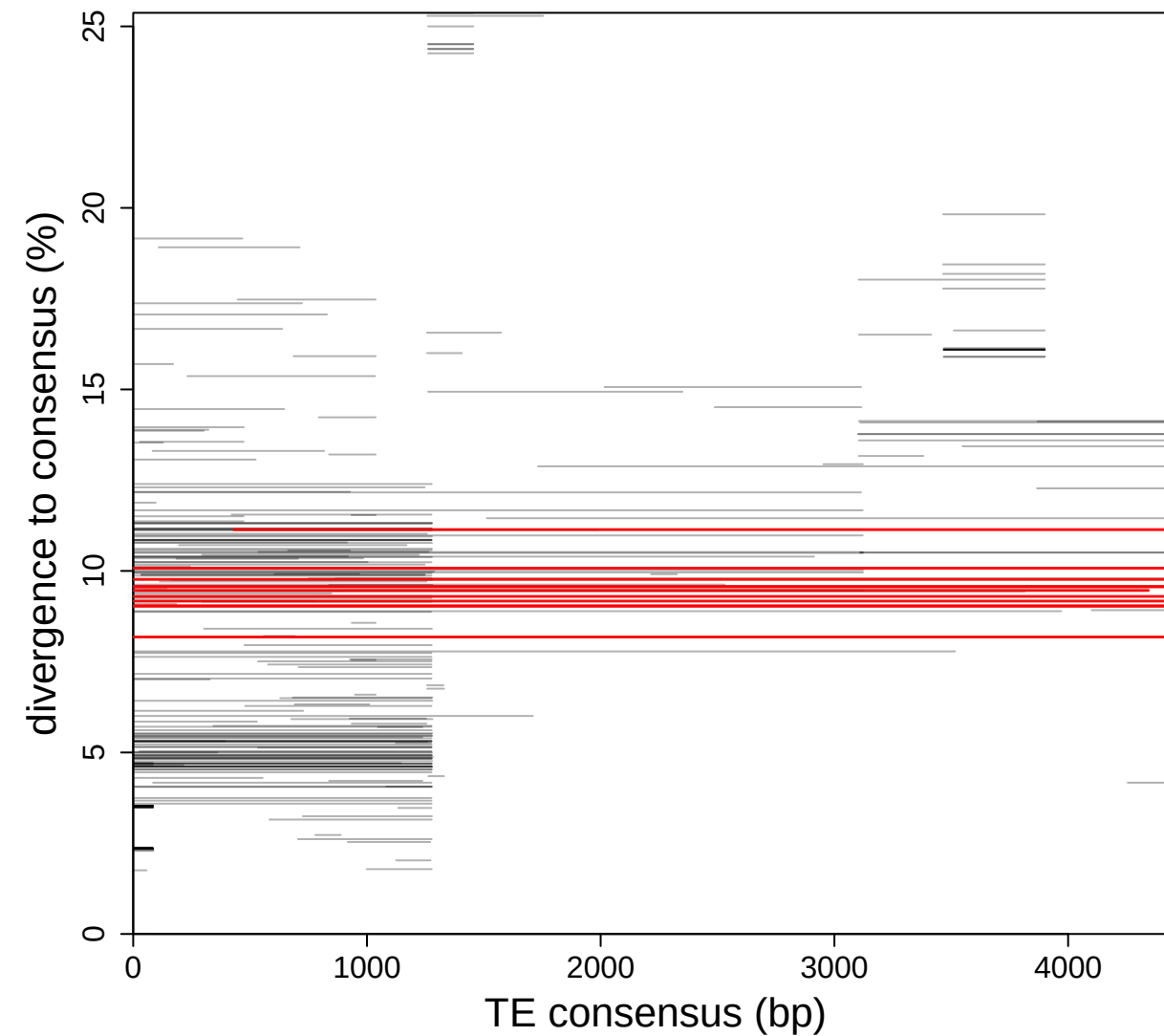
ly_2686.fasta.b.bed uf.bed g_3.bed fm_1.bed_0_0 bcln.fa aln.f
size: 8416bp; fragments: 883; full length: 0 (>=7574.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

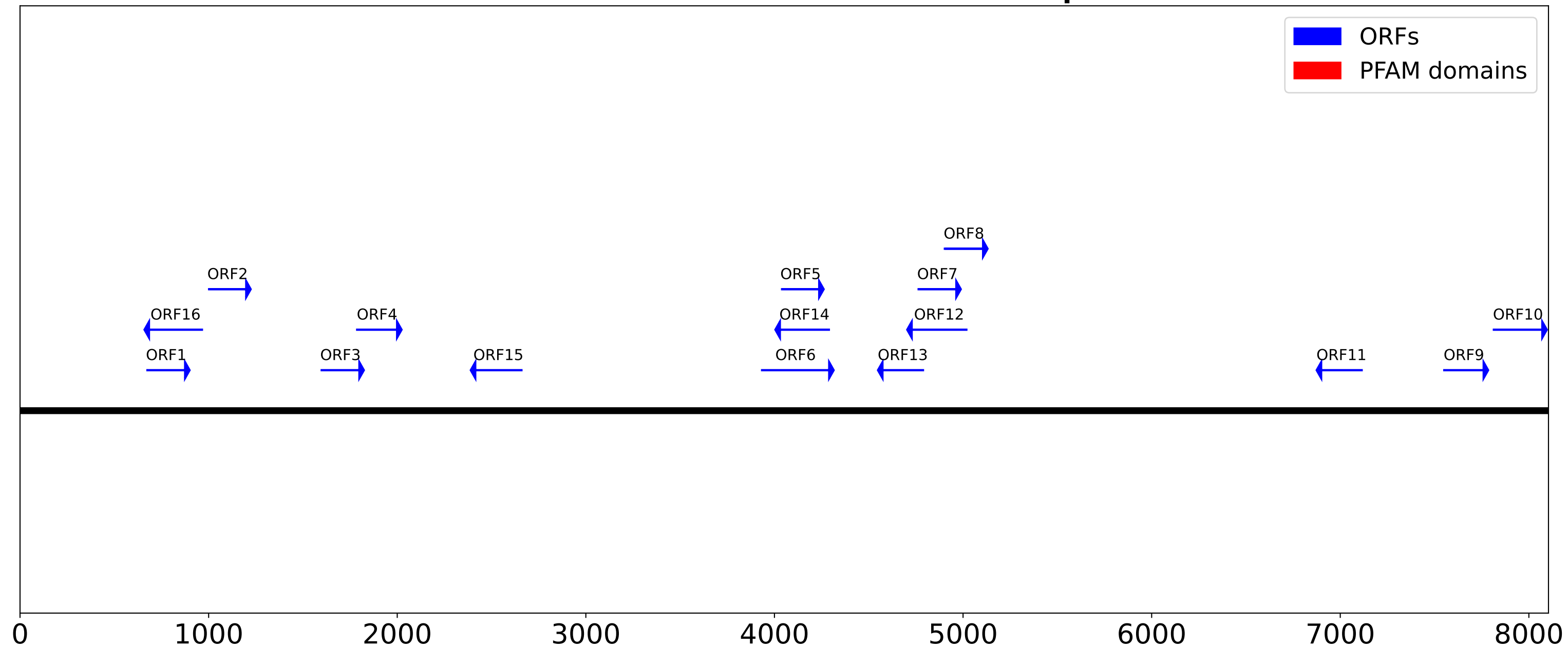


TE: rnd_5_family_2686
size: 4470bp; fragments: 317; full length: 11 (≥ 4023 bp)

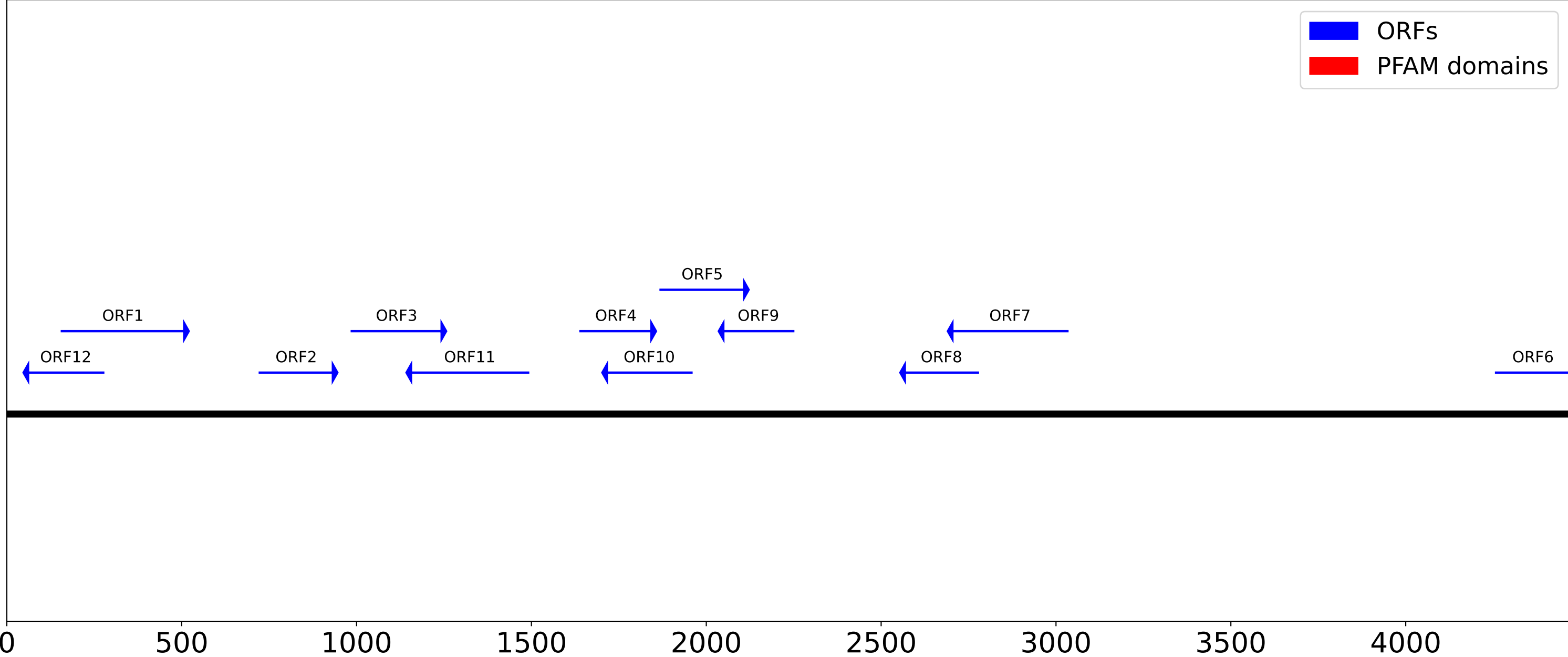
Before TEtrimmer 4470 bp



After TEtrimmer ORF and PFAM domain plot



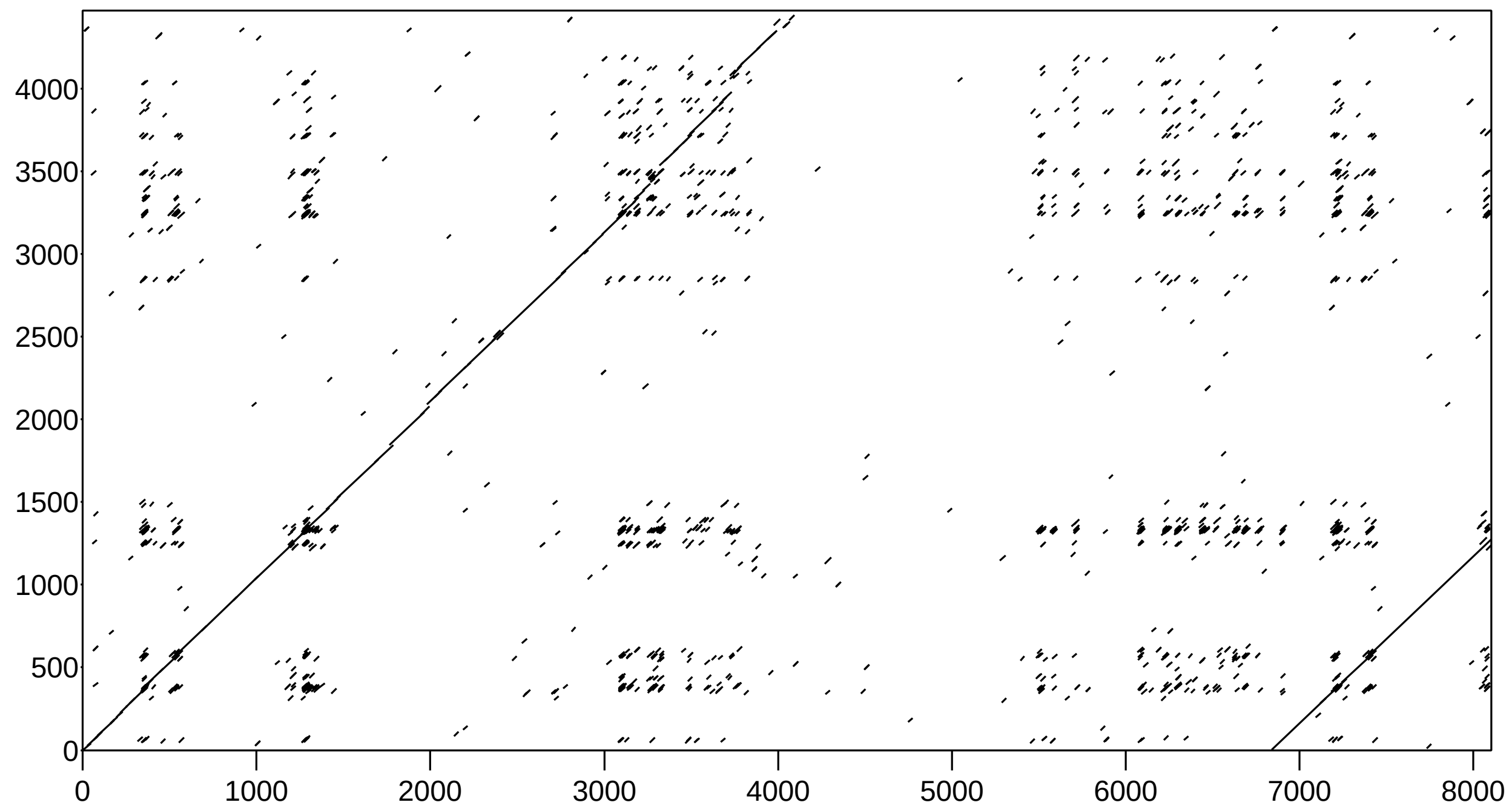
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)