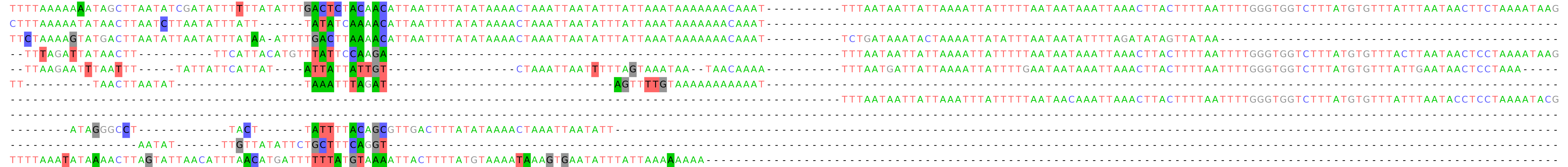


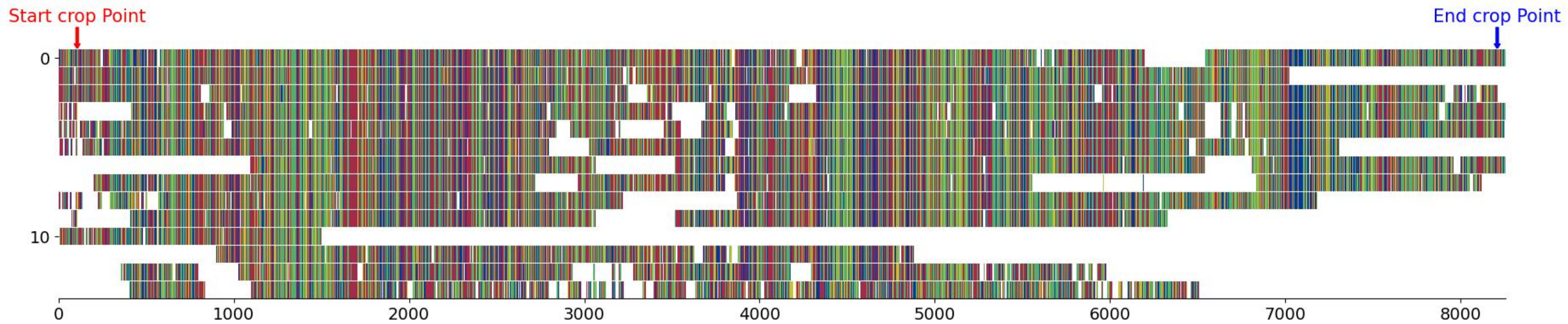
MSA length = 8105

Start crop Point



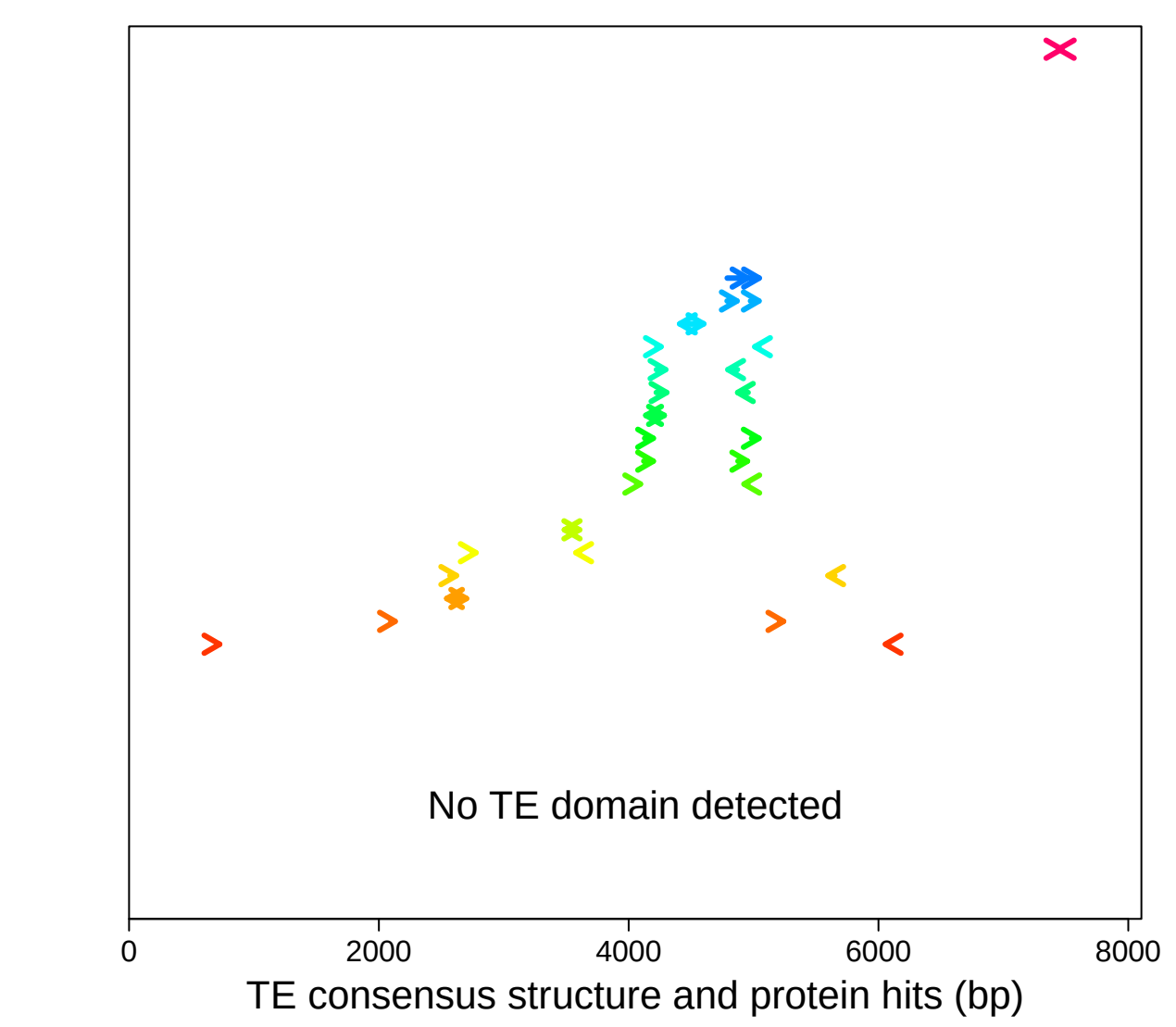
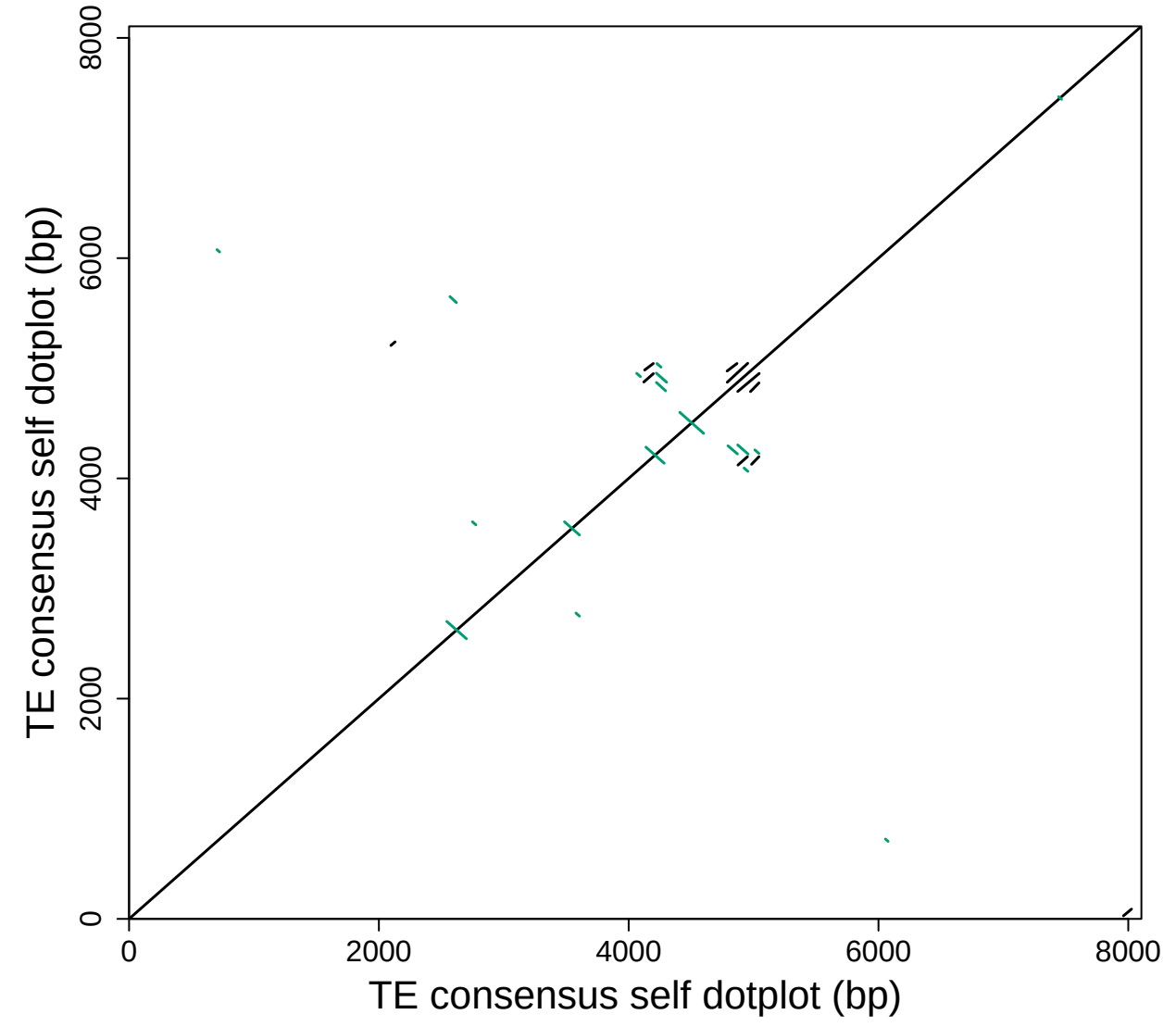
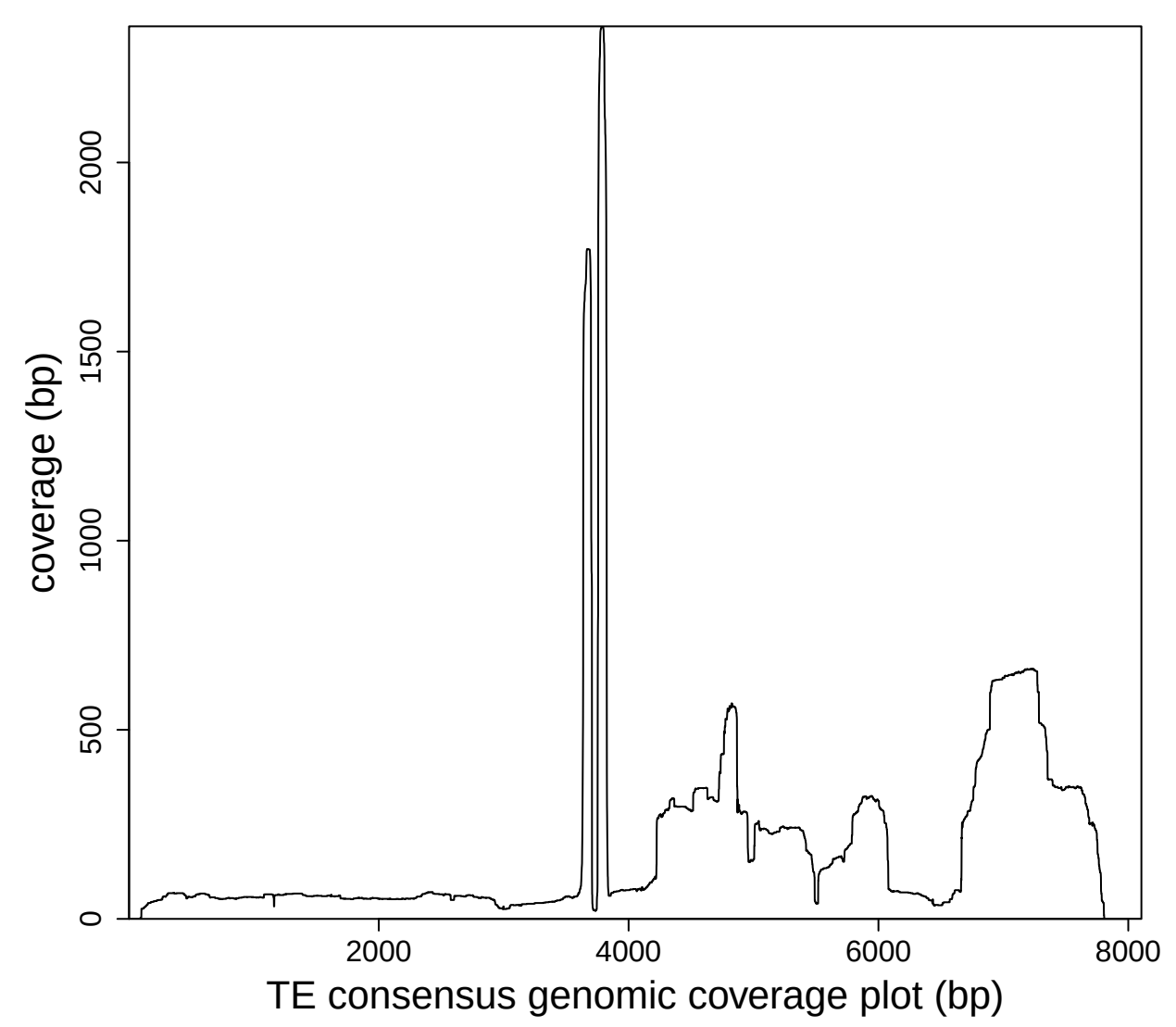
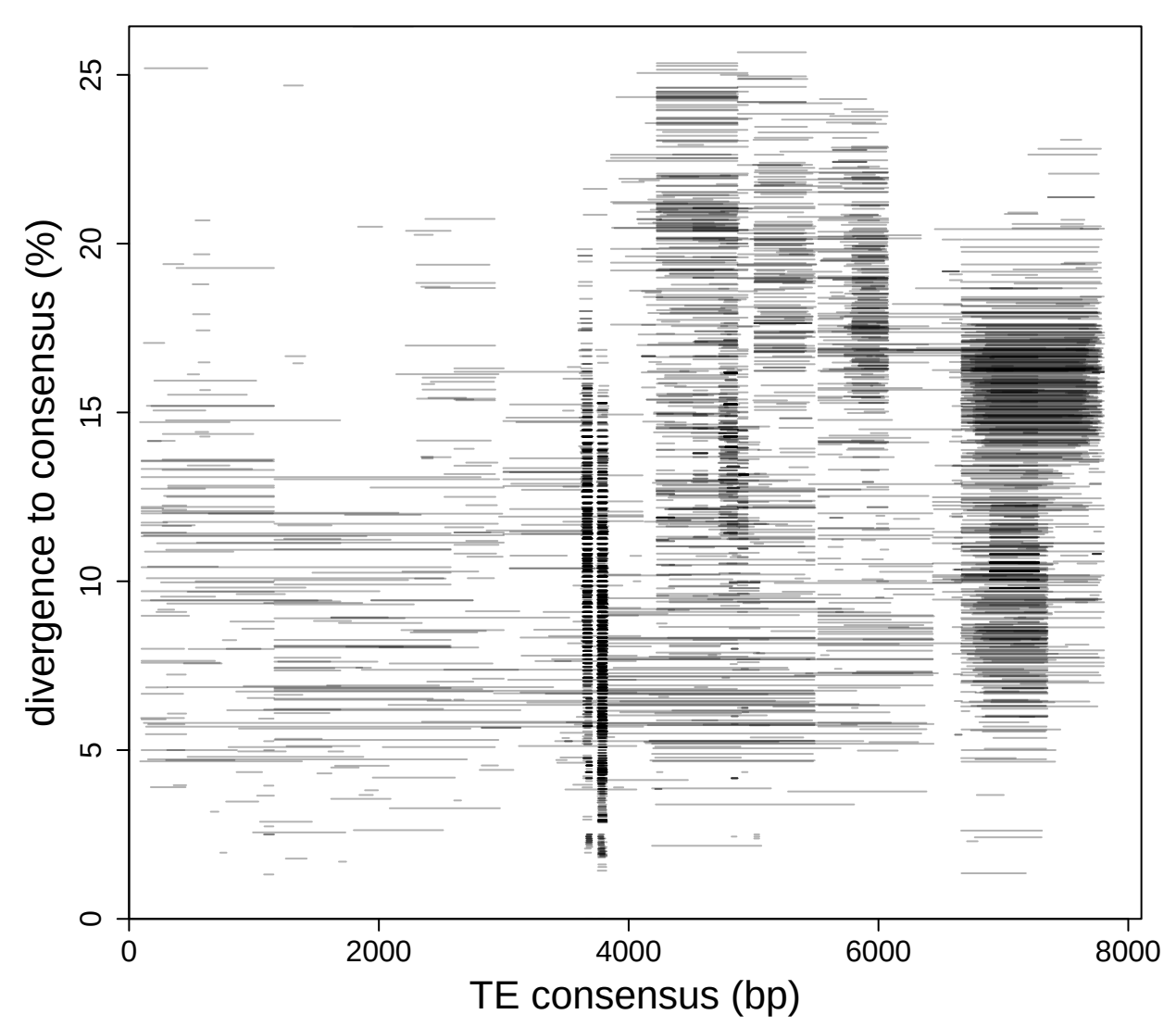
End crop Point





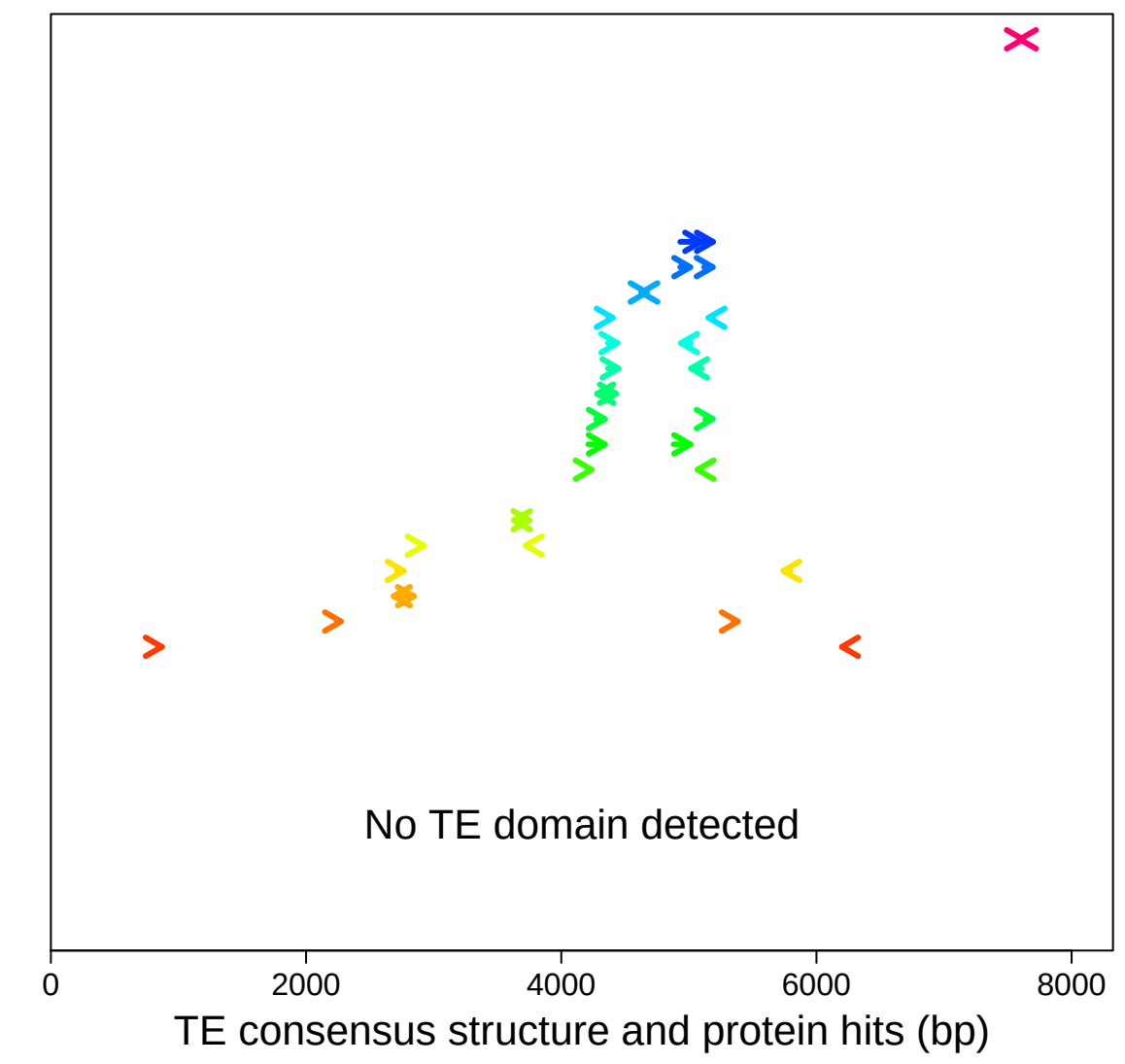
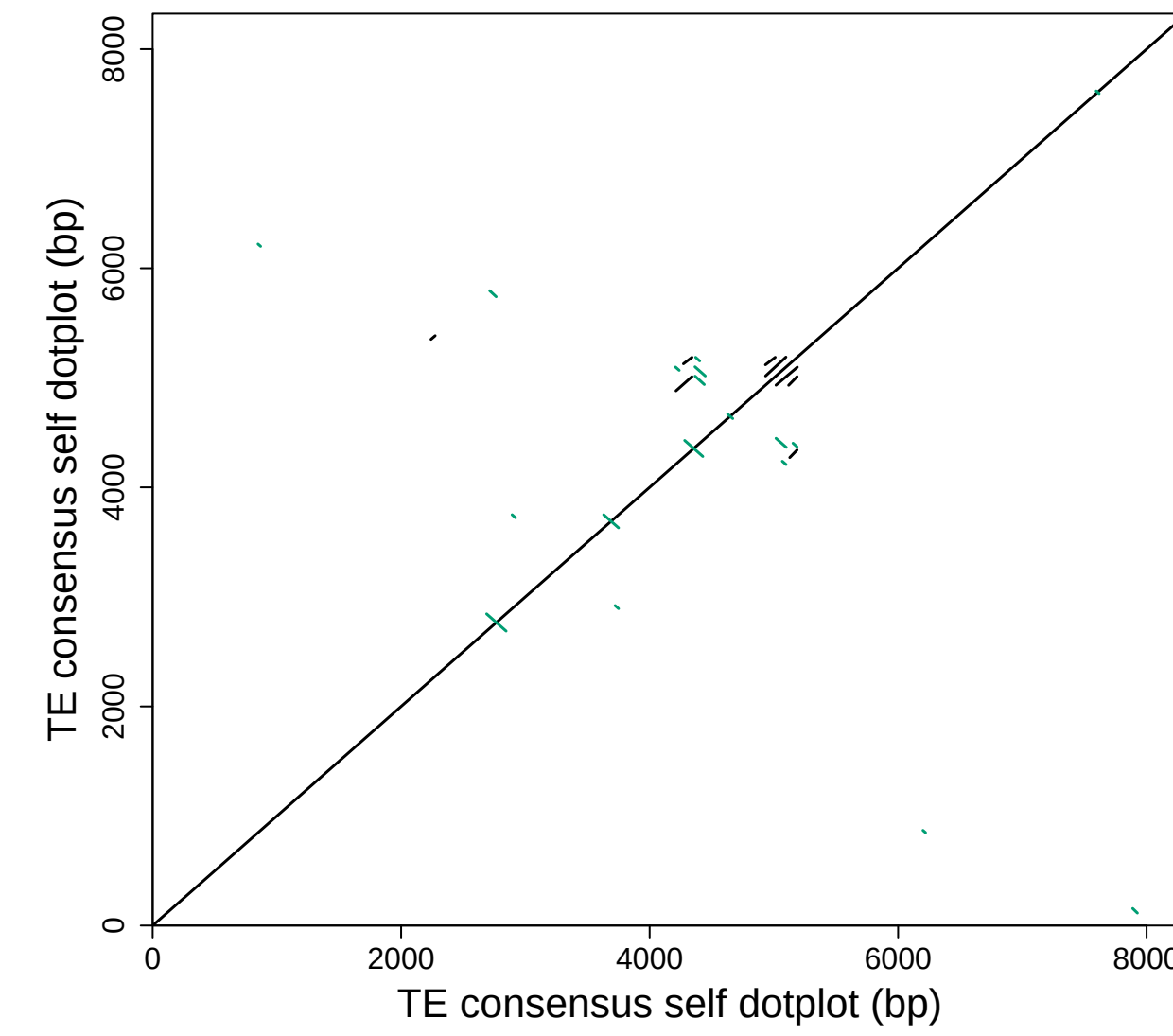
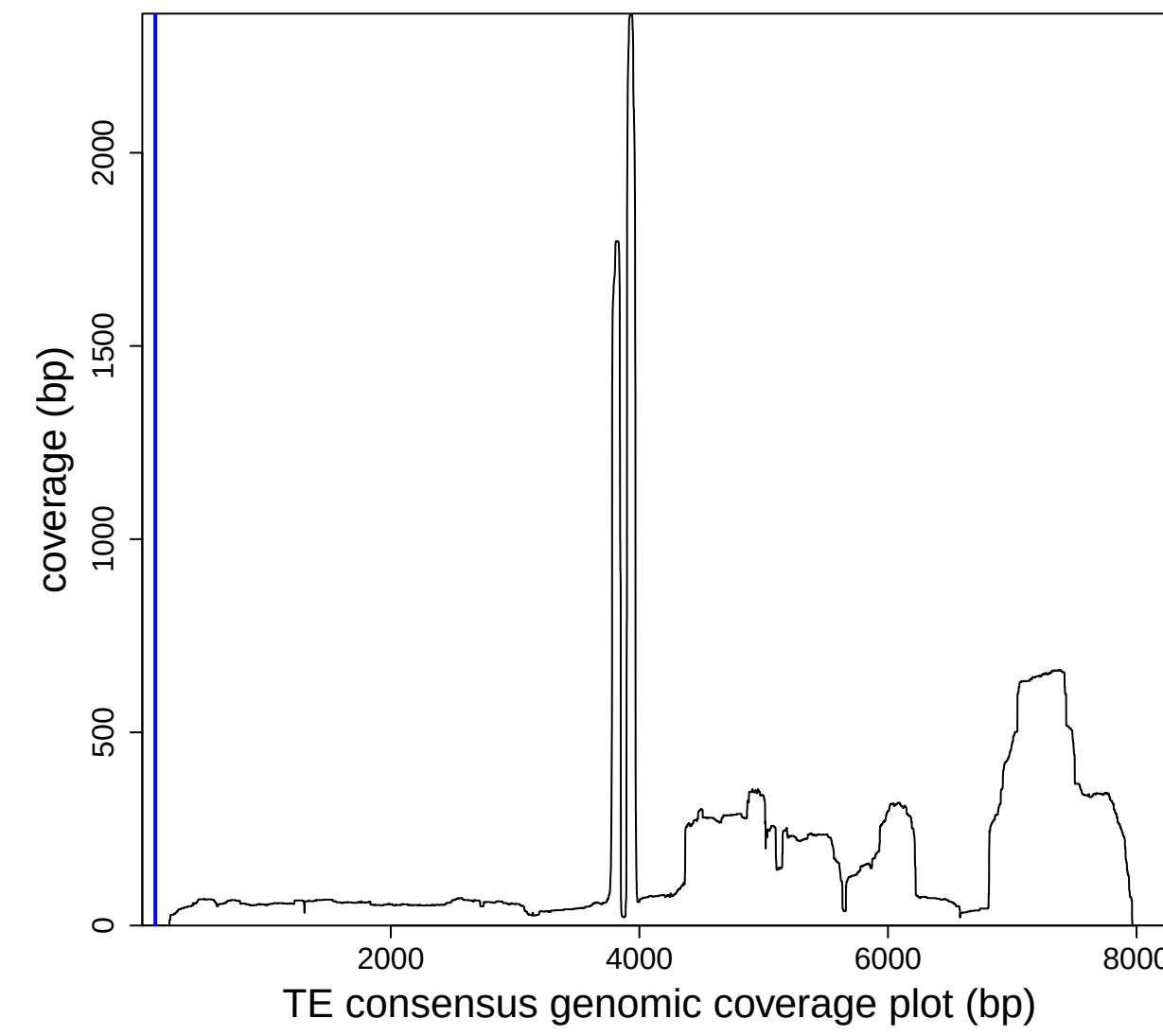
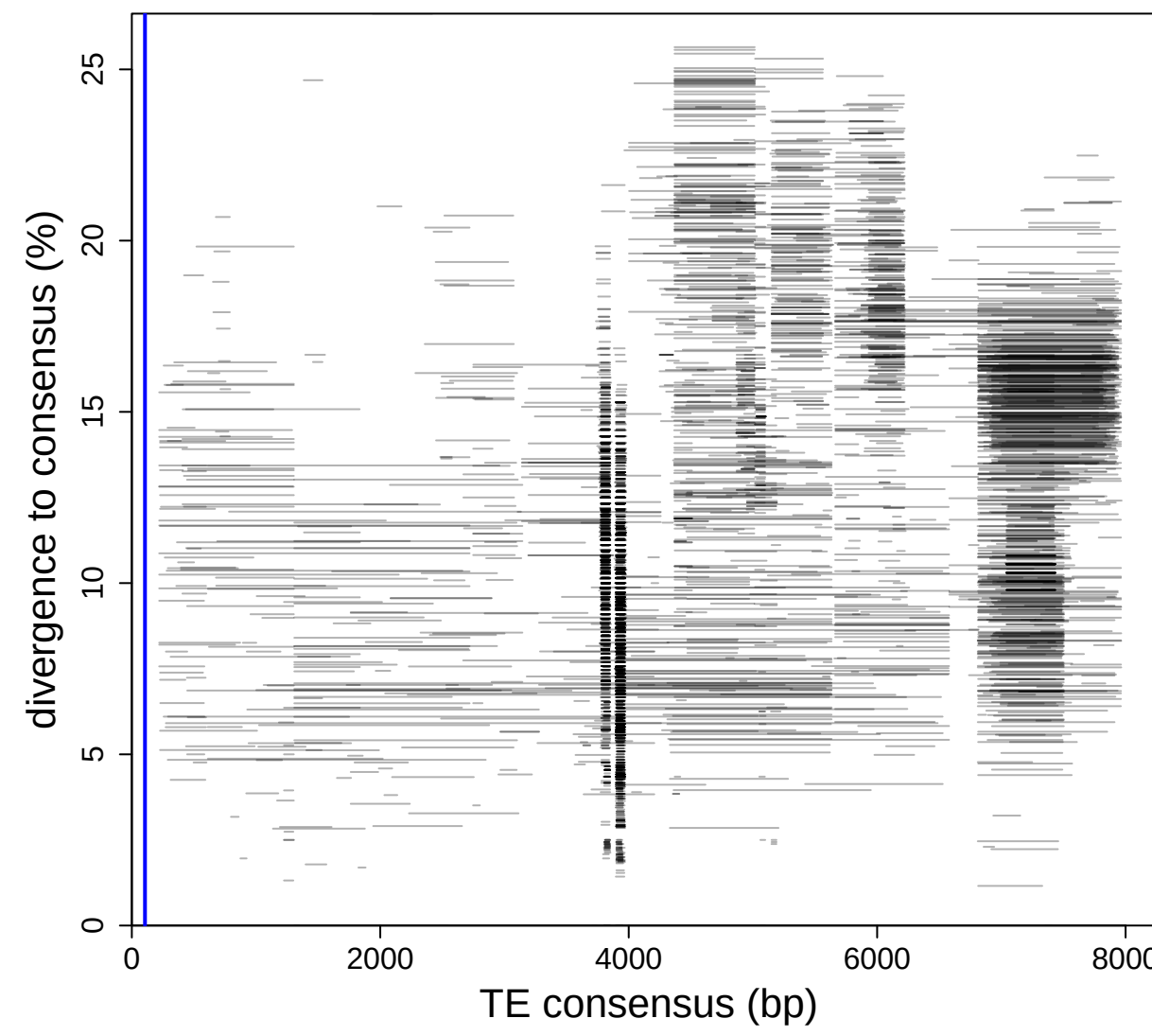
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 8105bp; fragments: 6443; full length: 0 (>=7294.5bp)

After TEtrimmer 8105 bp



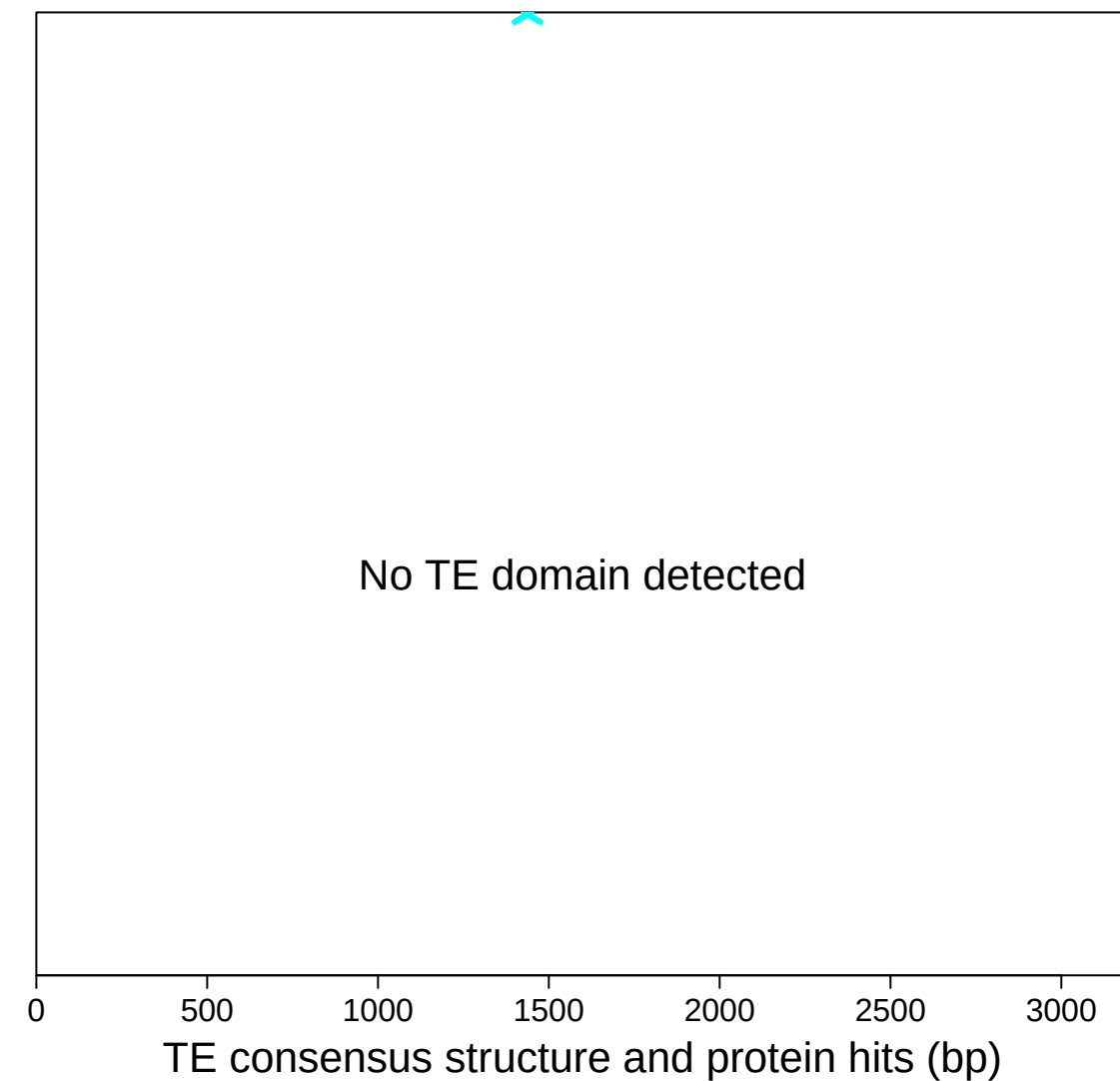
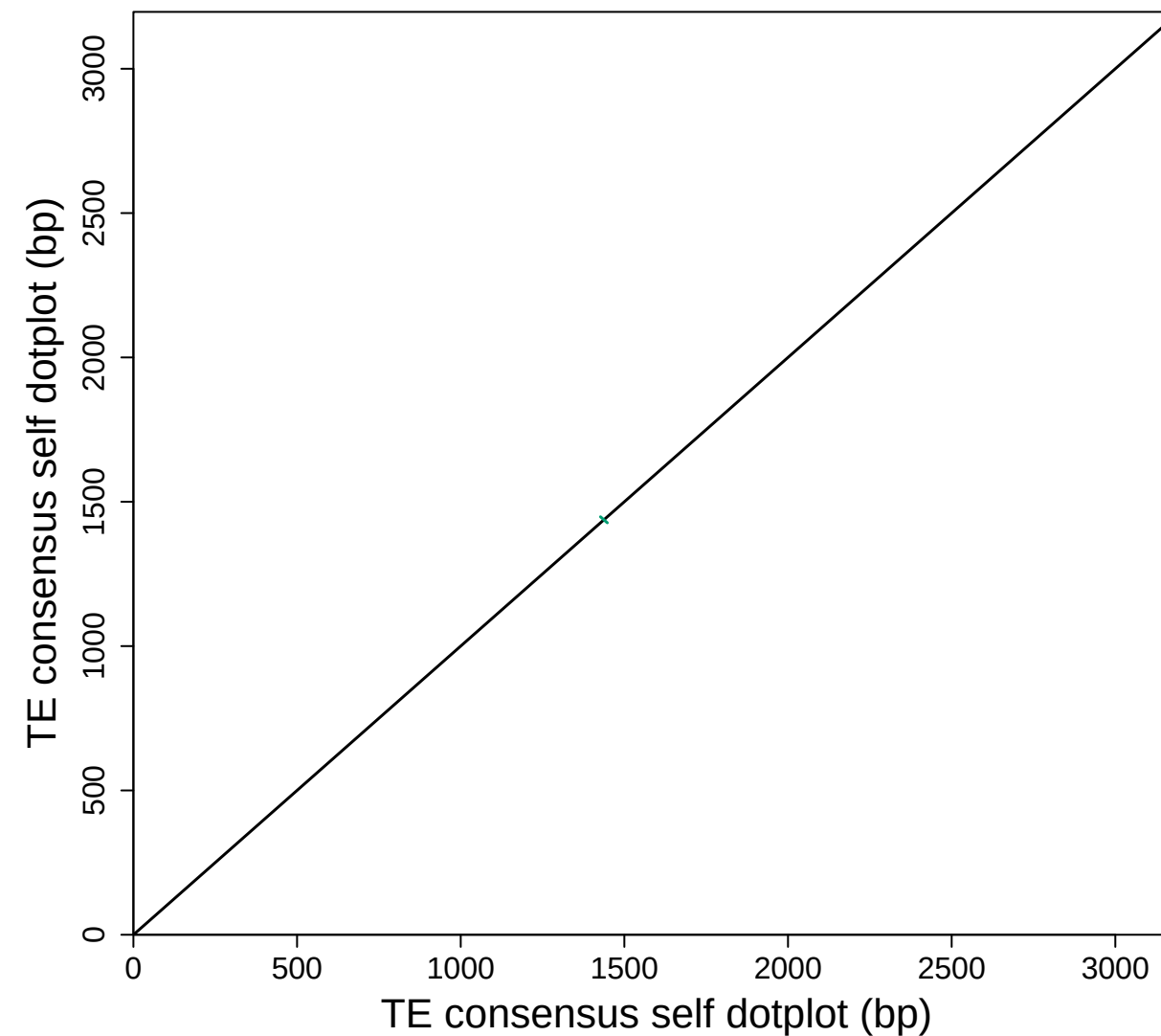
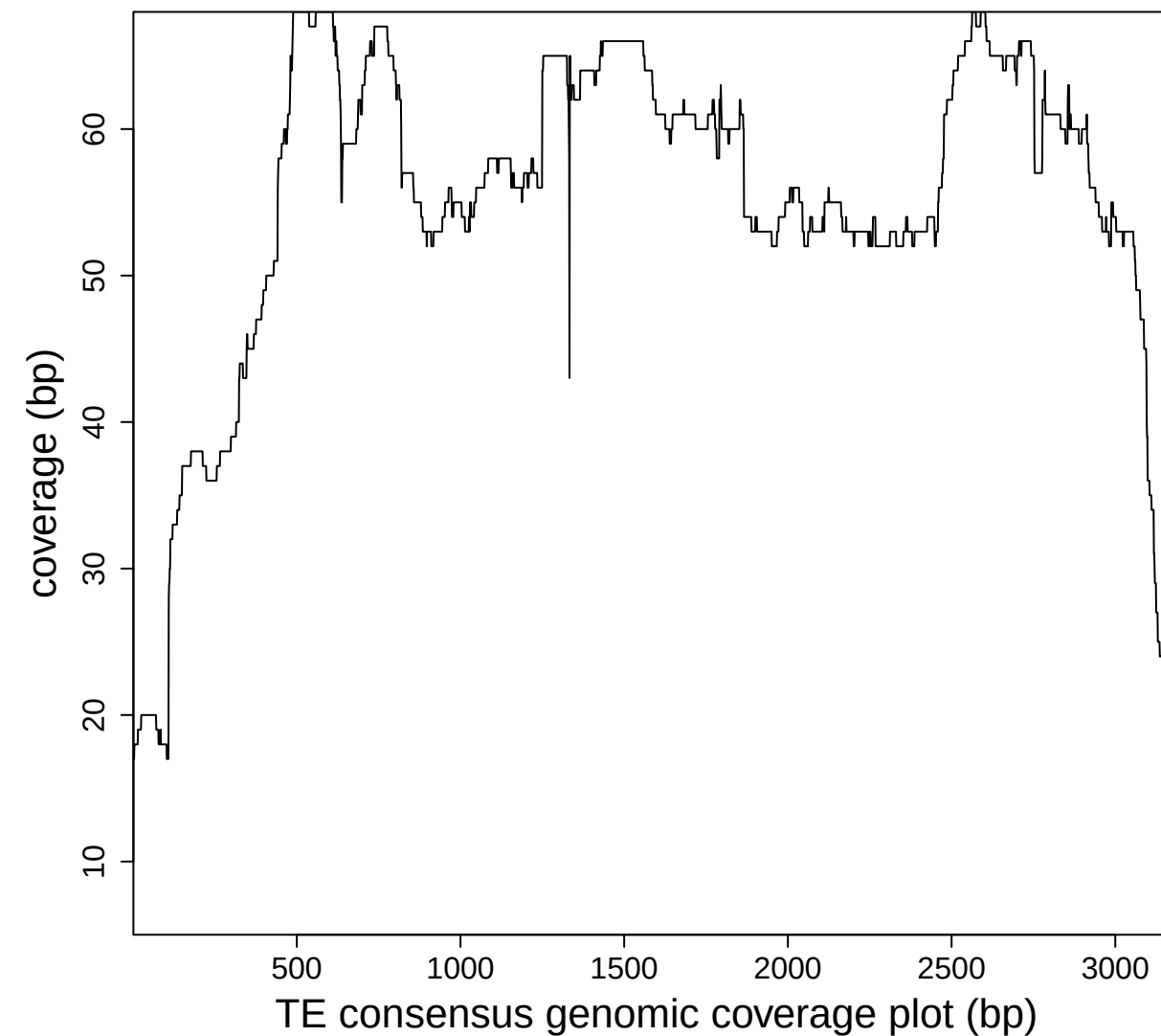
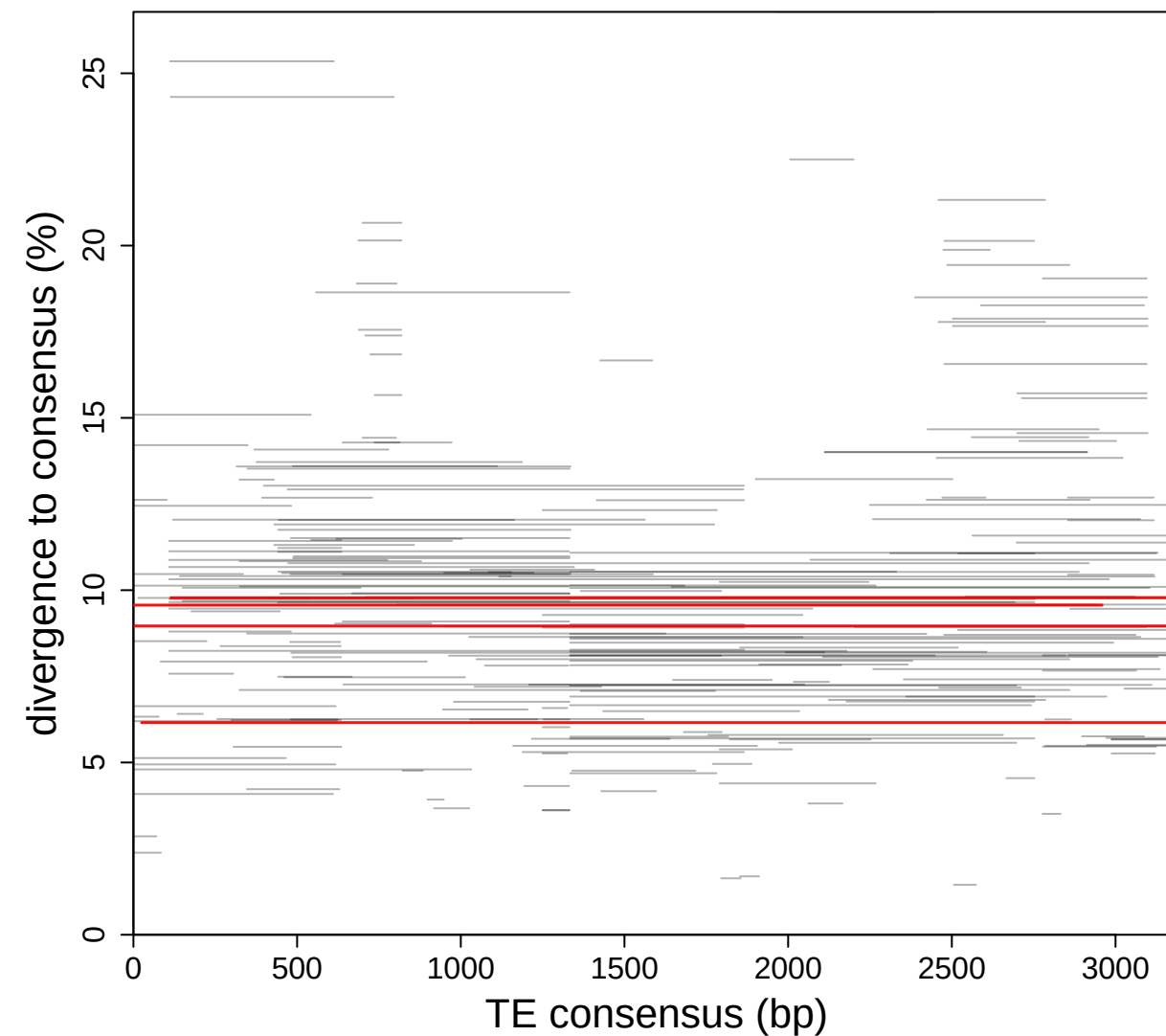
6.fasta.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 8324bp; fragments: 6138; full length: 0 (>=7491.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

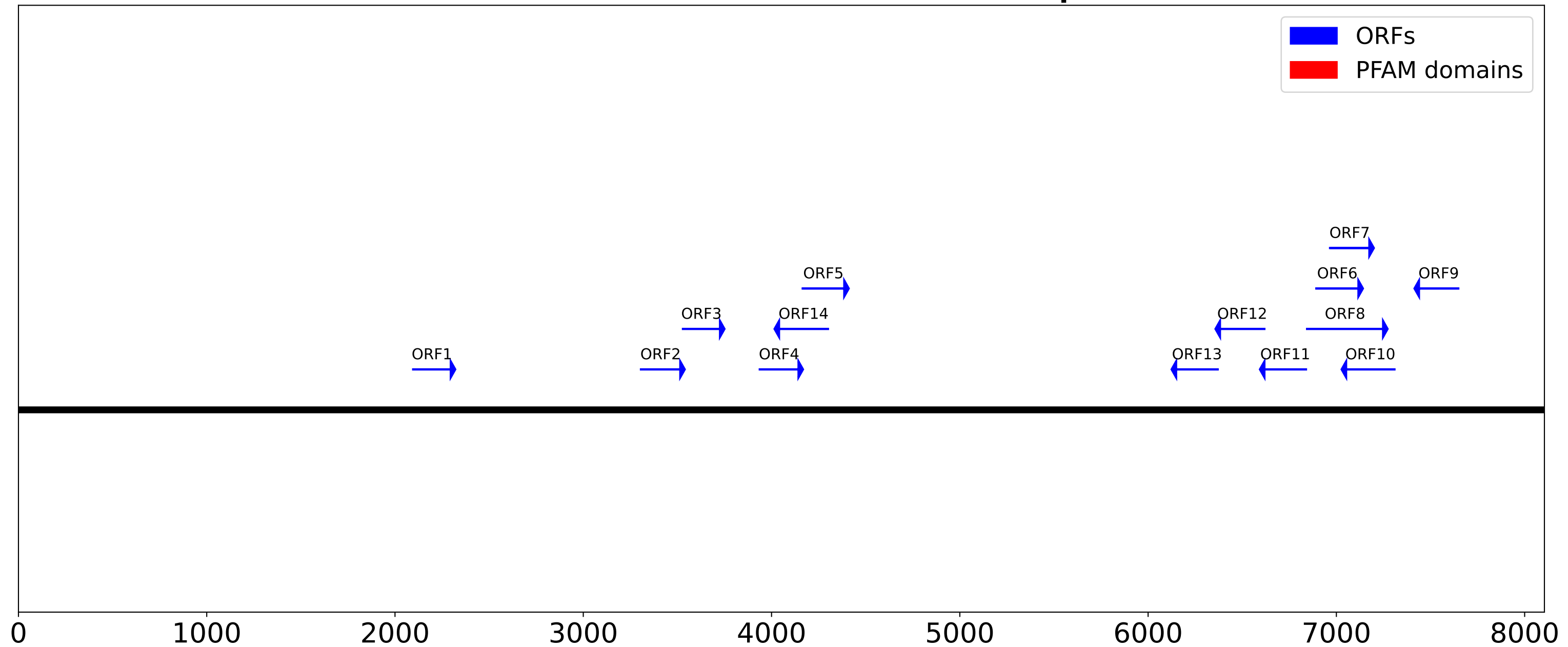


TE: rnd_5_family_8356
size: 3197bp; fragments: 242; full length: 4 (≥ 2877.3 bp)

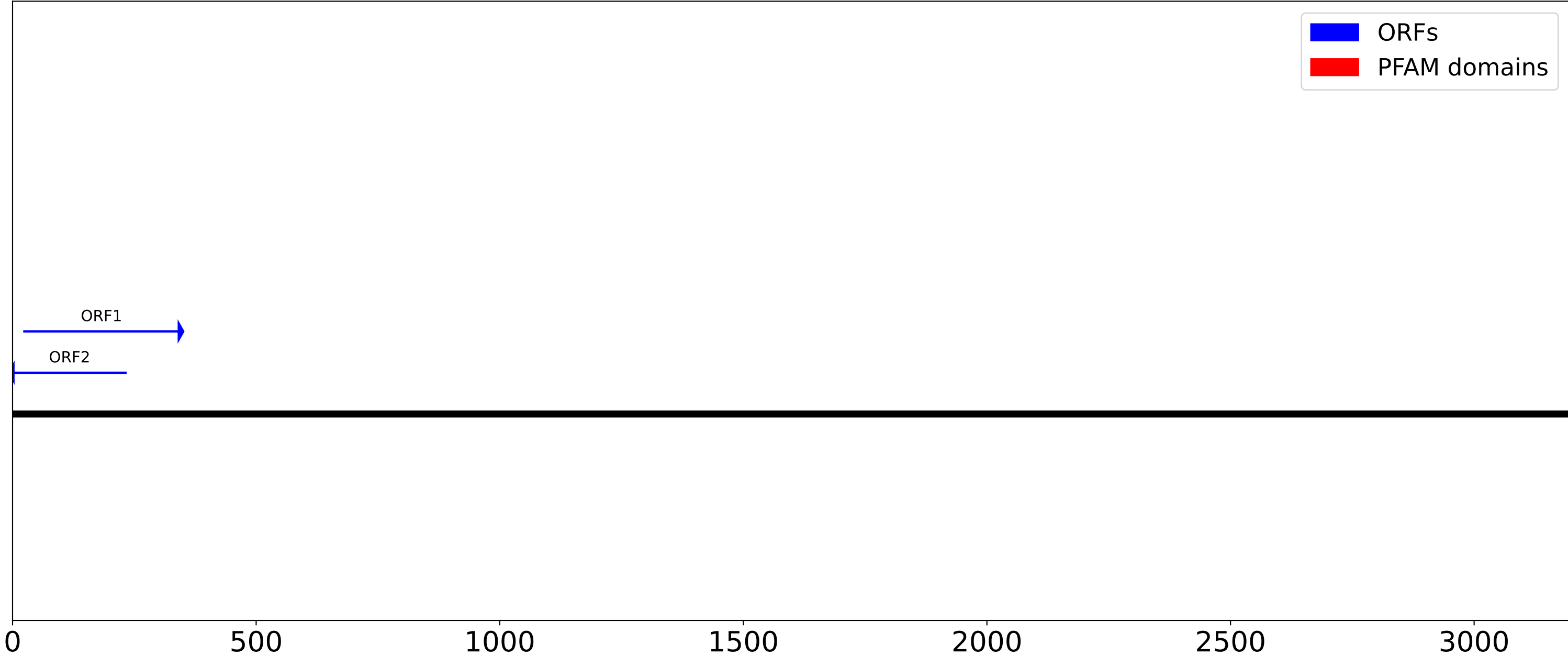
Before TEtrimmer 3197 bp



After TEtrimmer ORF and PFAM domain plot



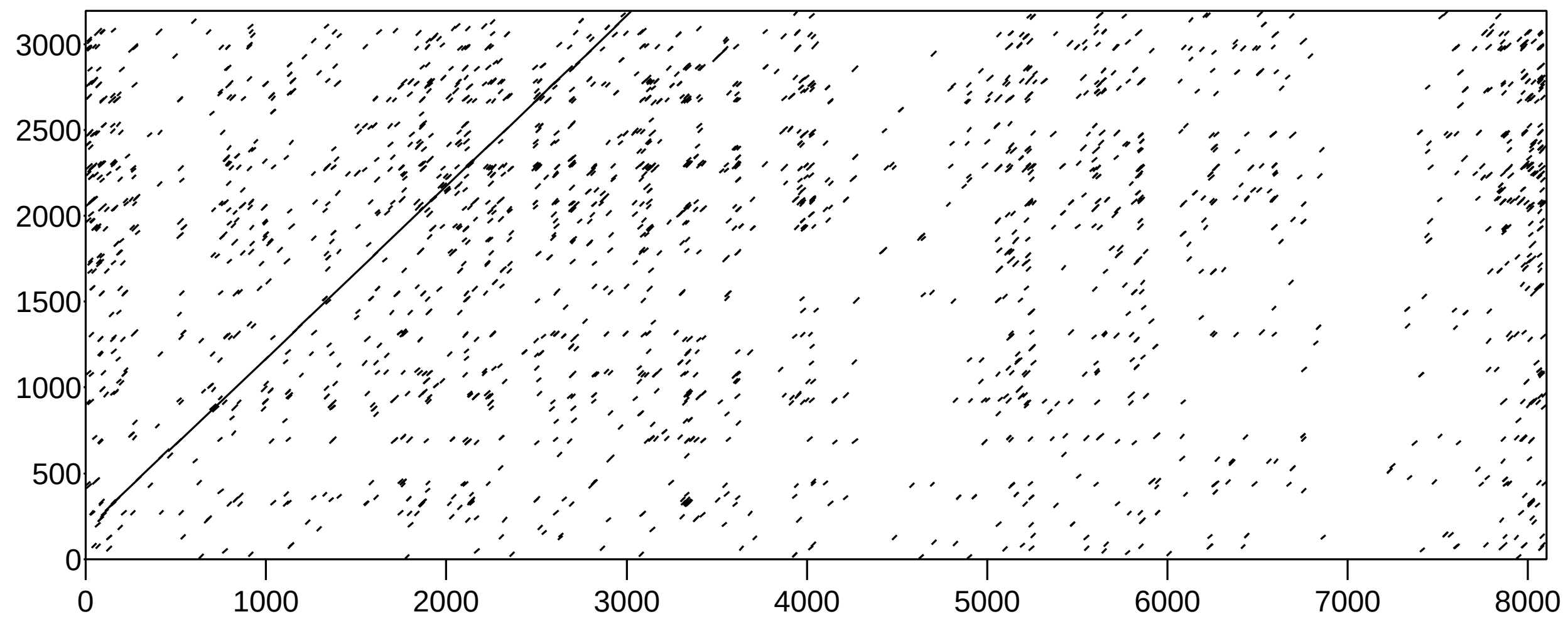
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)