

Start Crop Point

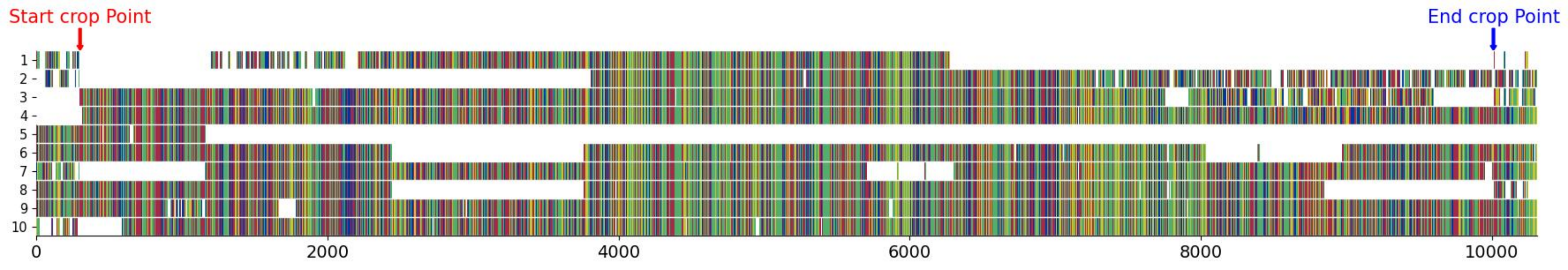


End Crop Point



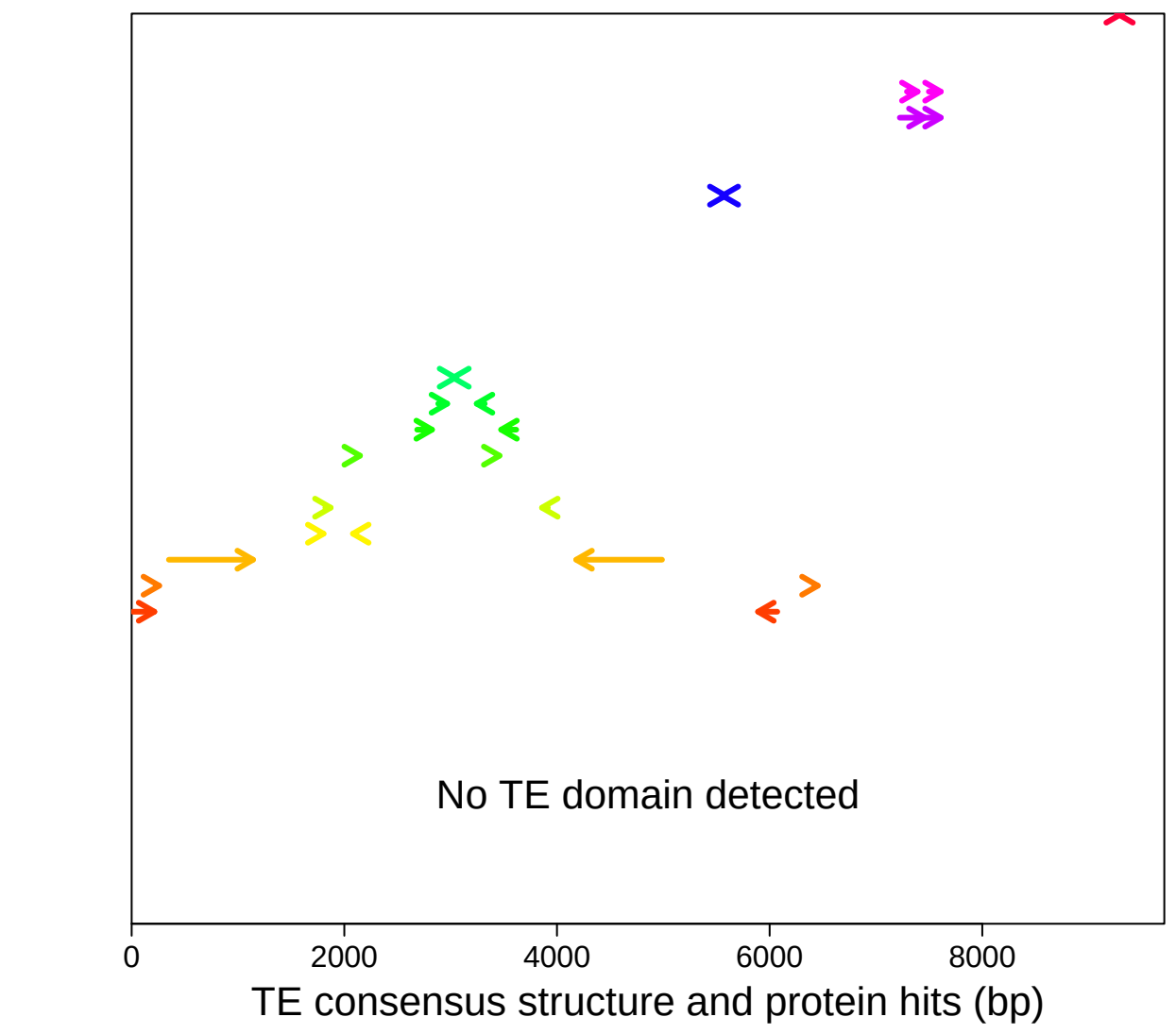
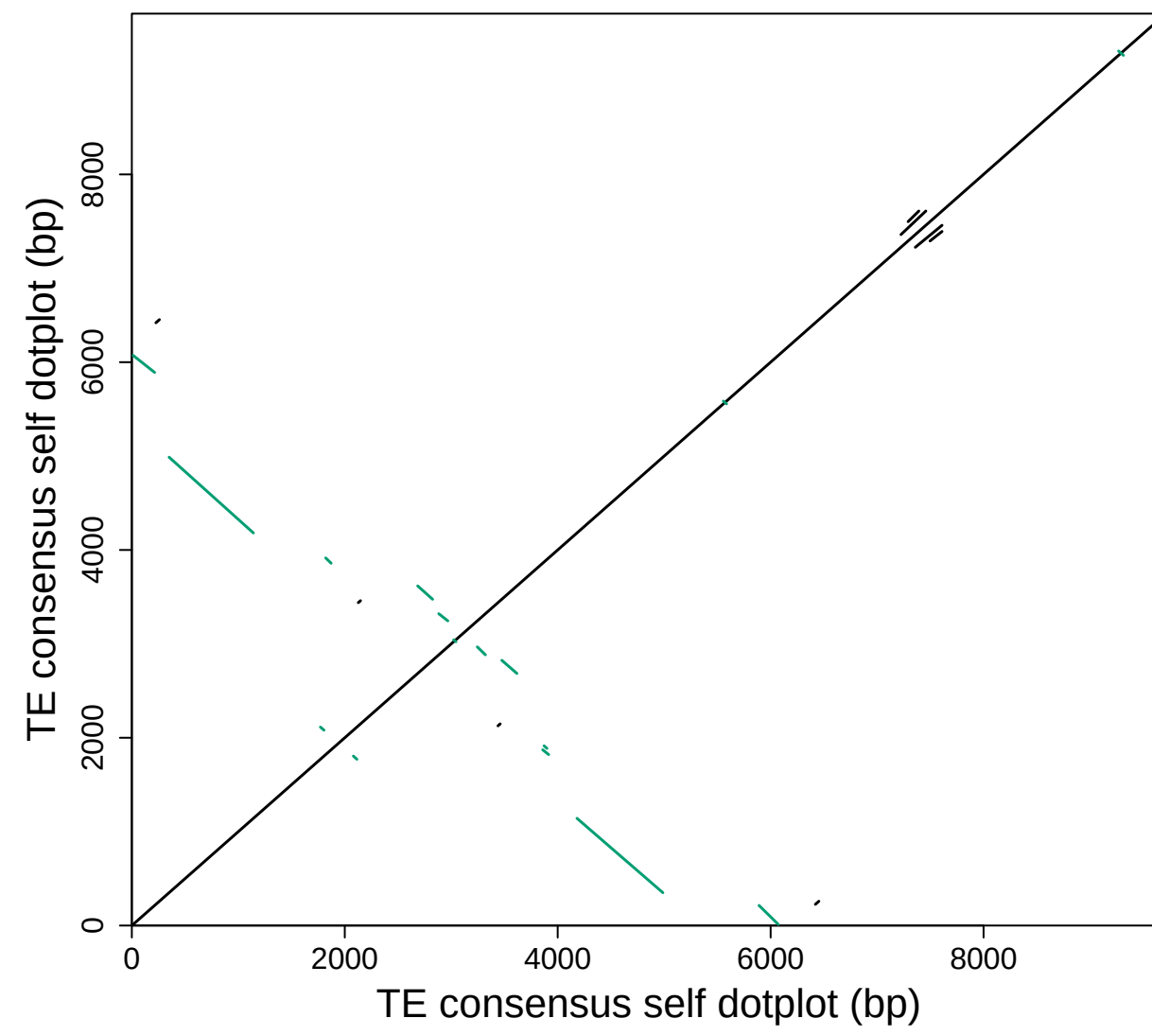
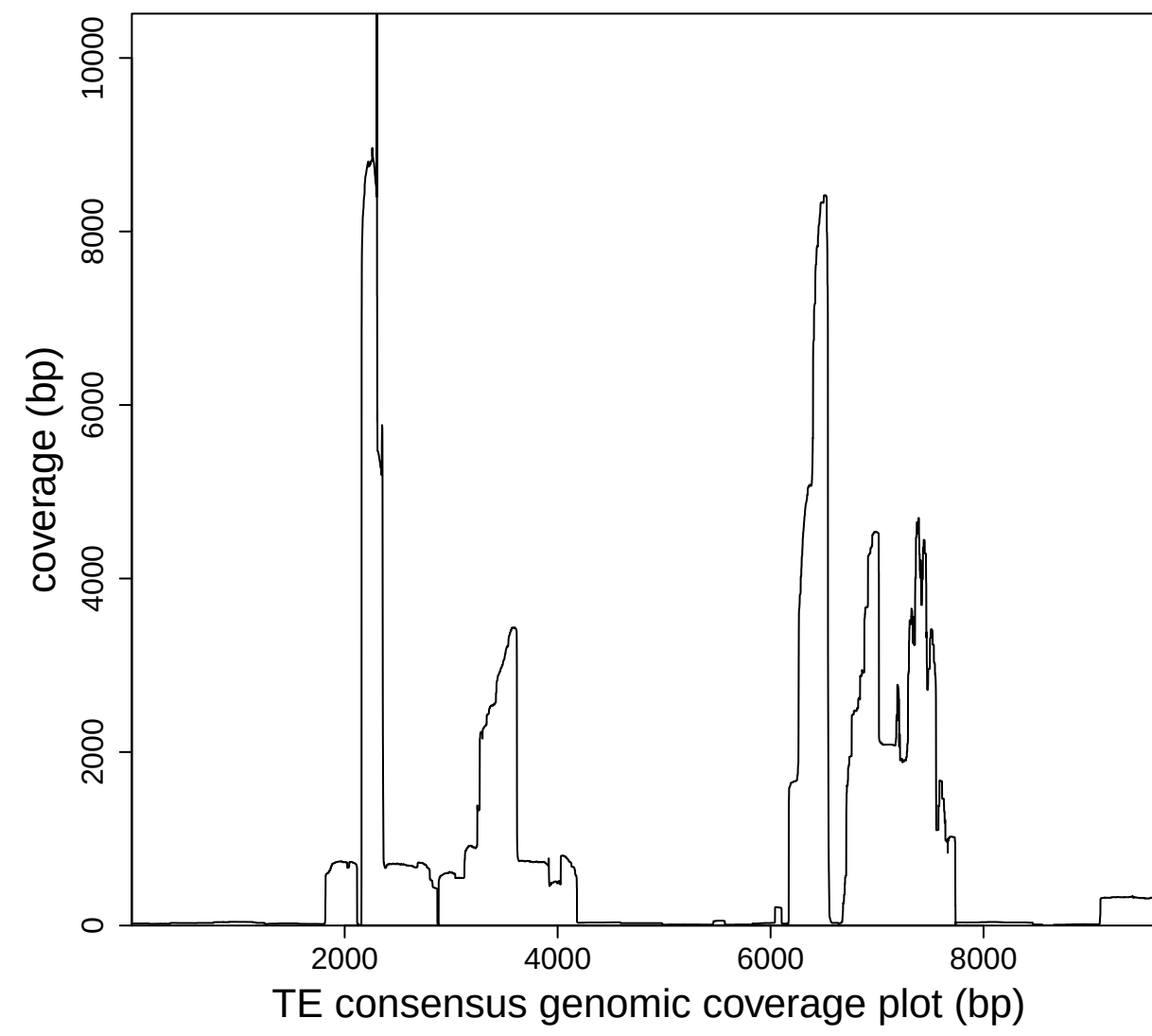
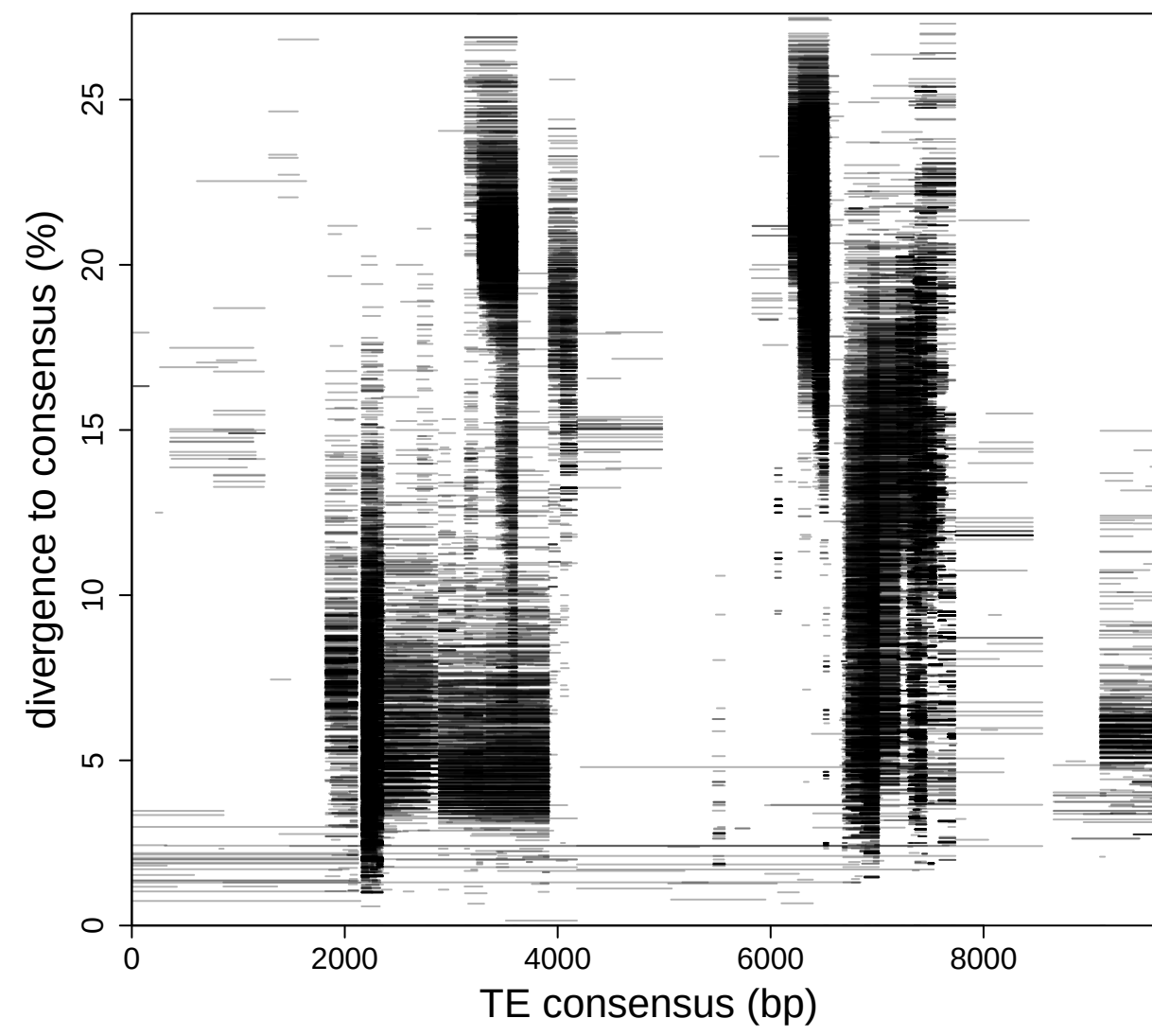
MSA length = 9712





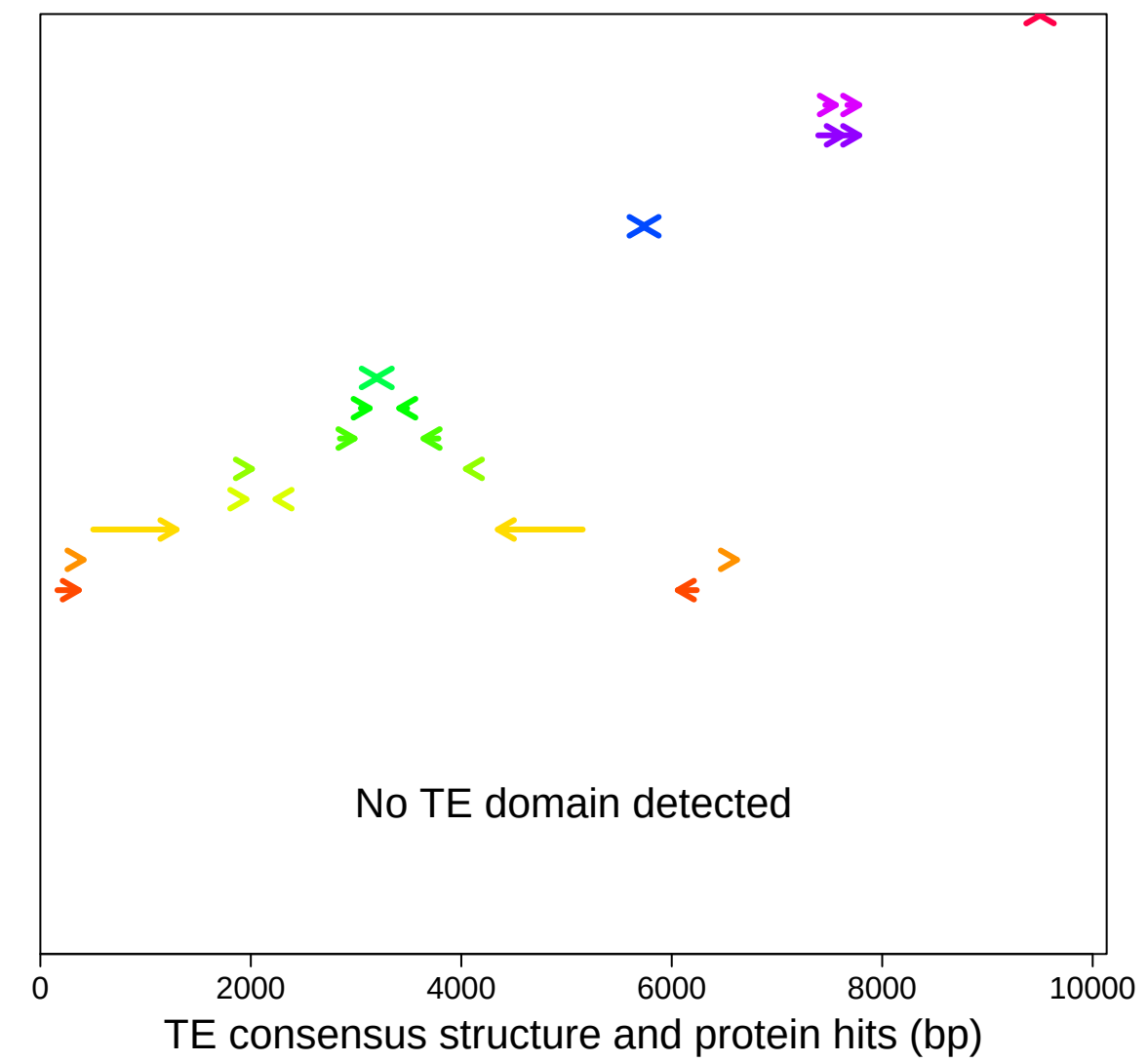
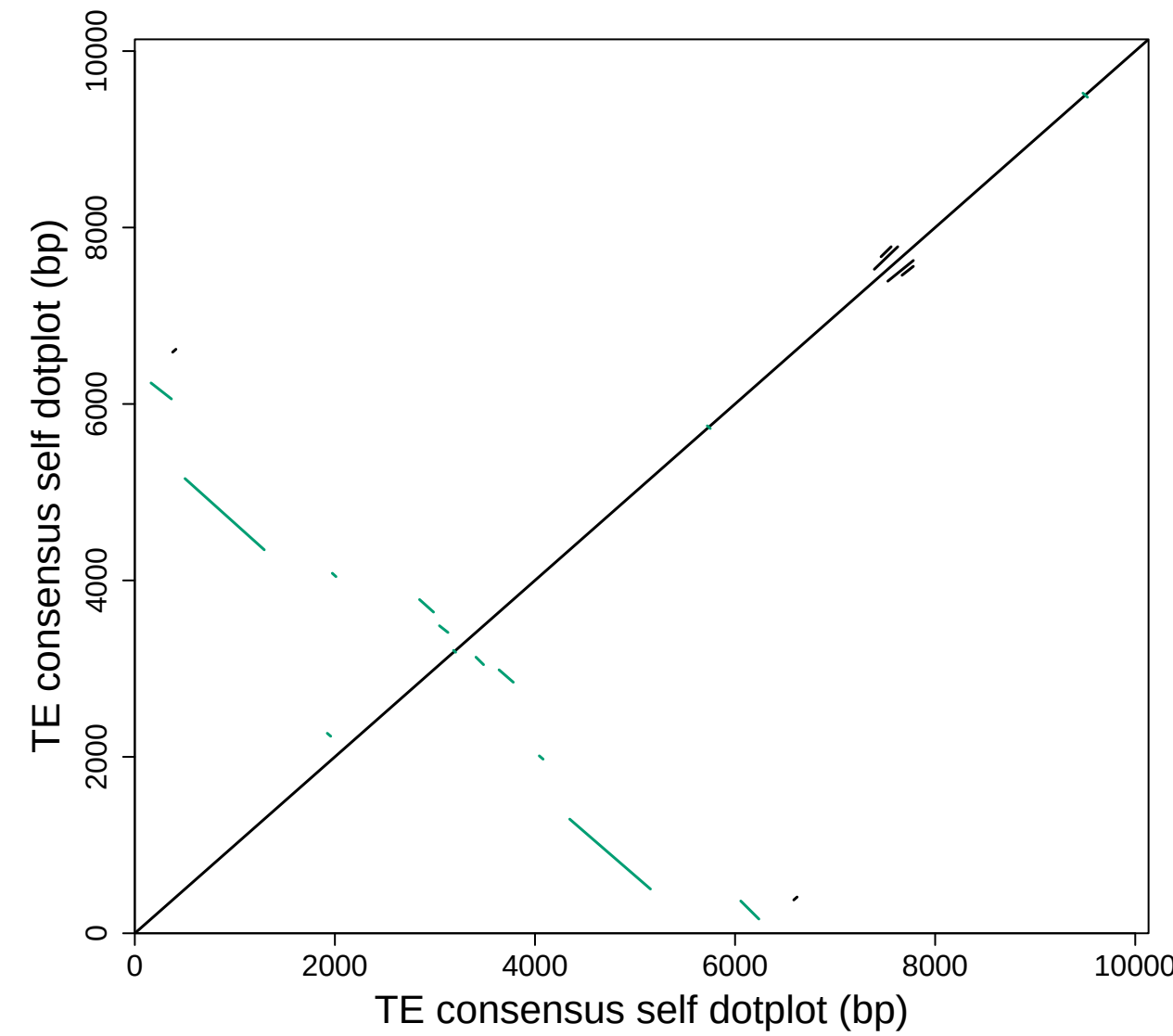
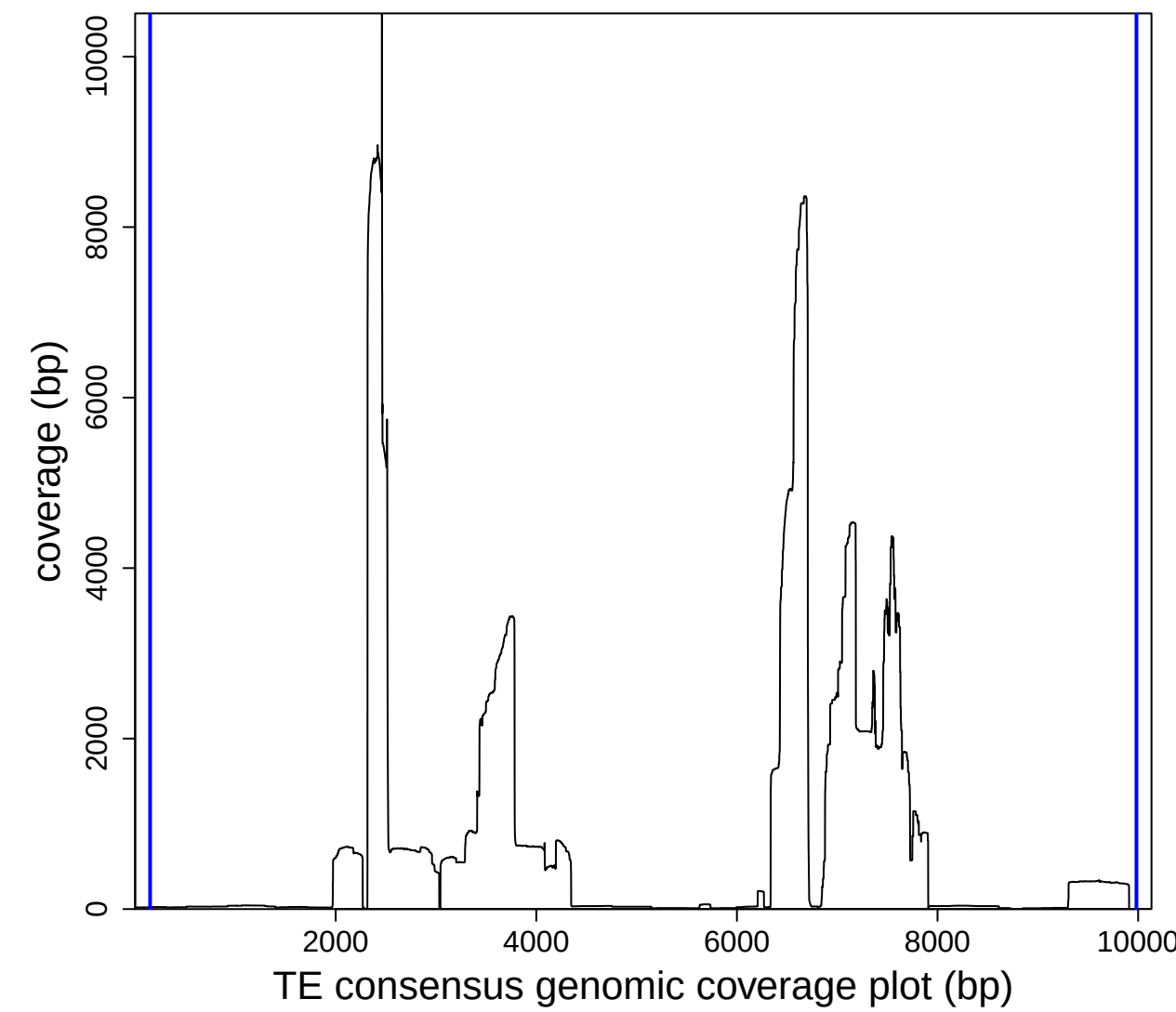
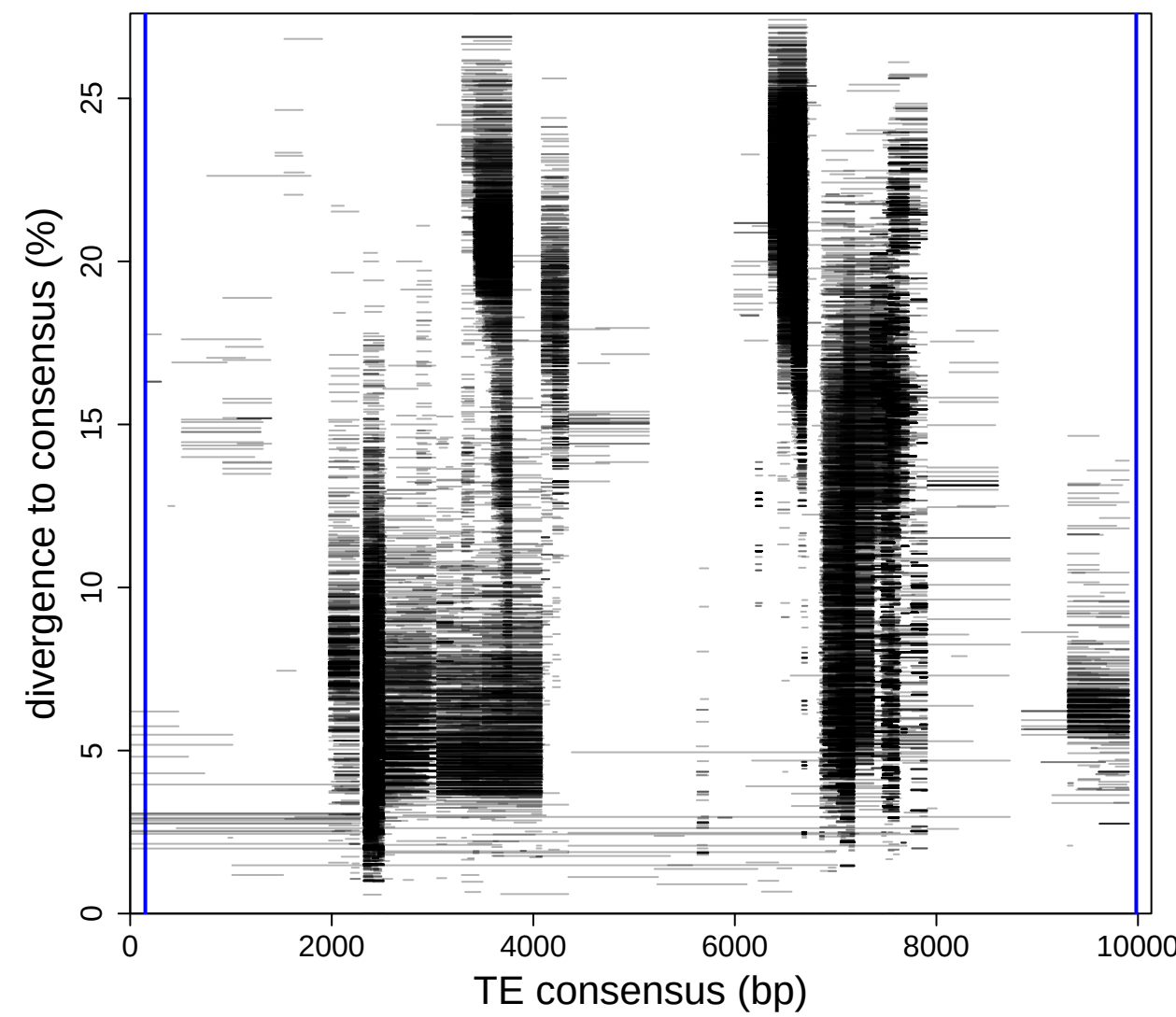
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 9712bp; fragments: 39729; full length: 0 (>=8740.8bp)

After TEtrimmer 9712 bp



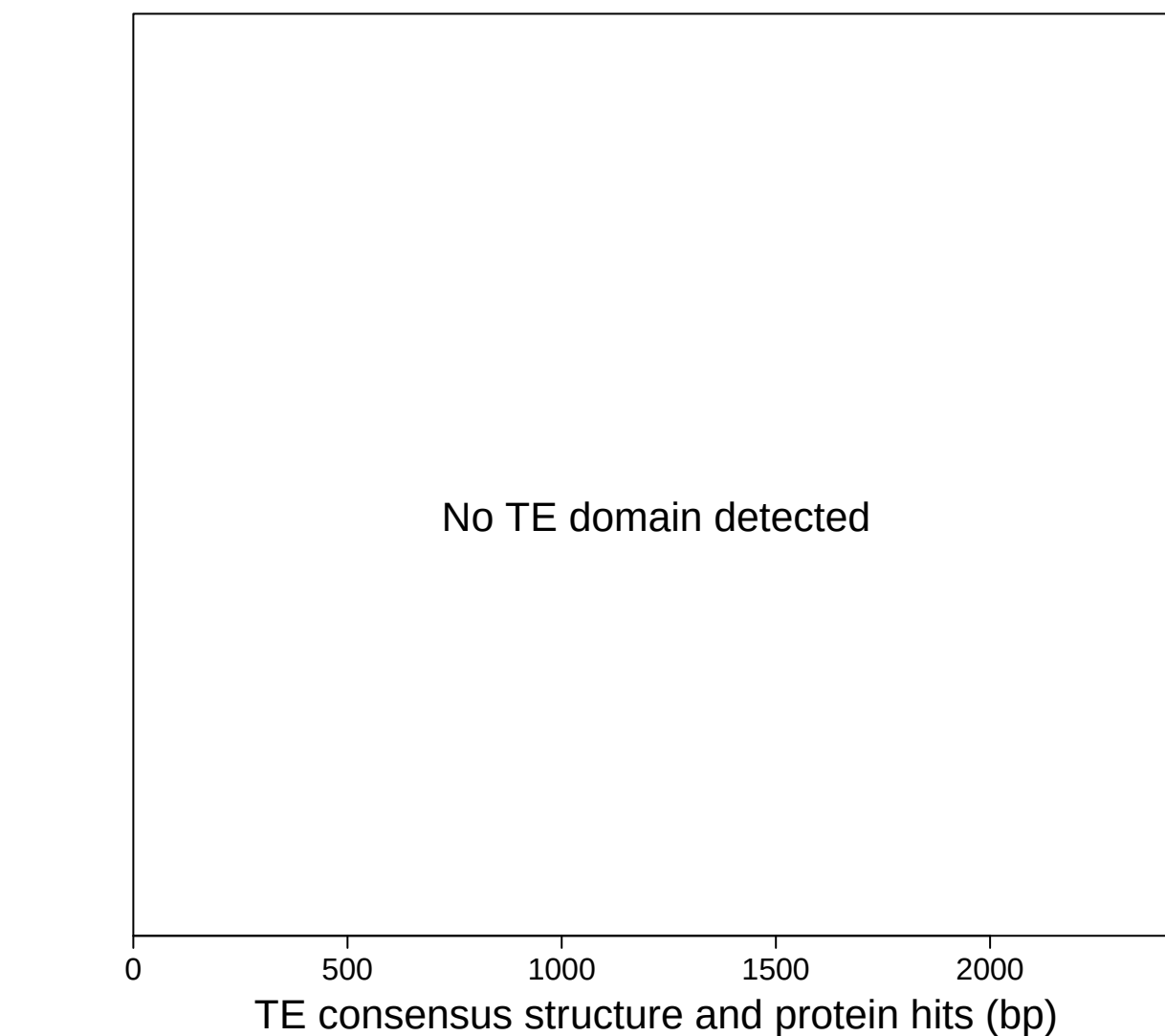
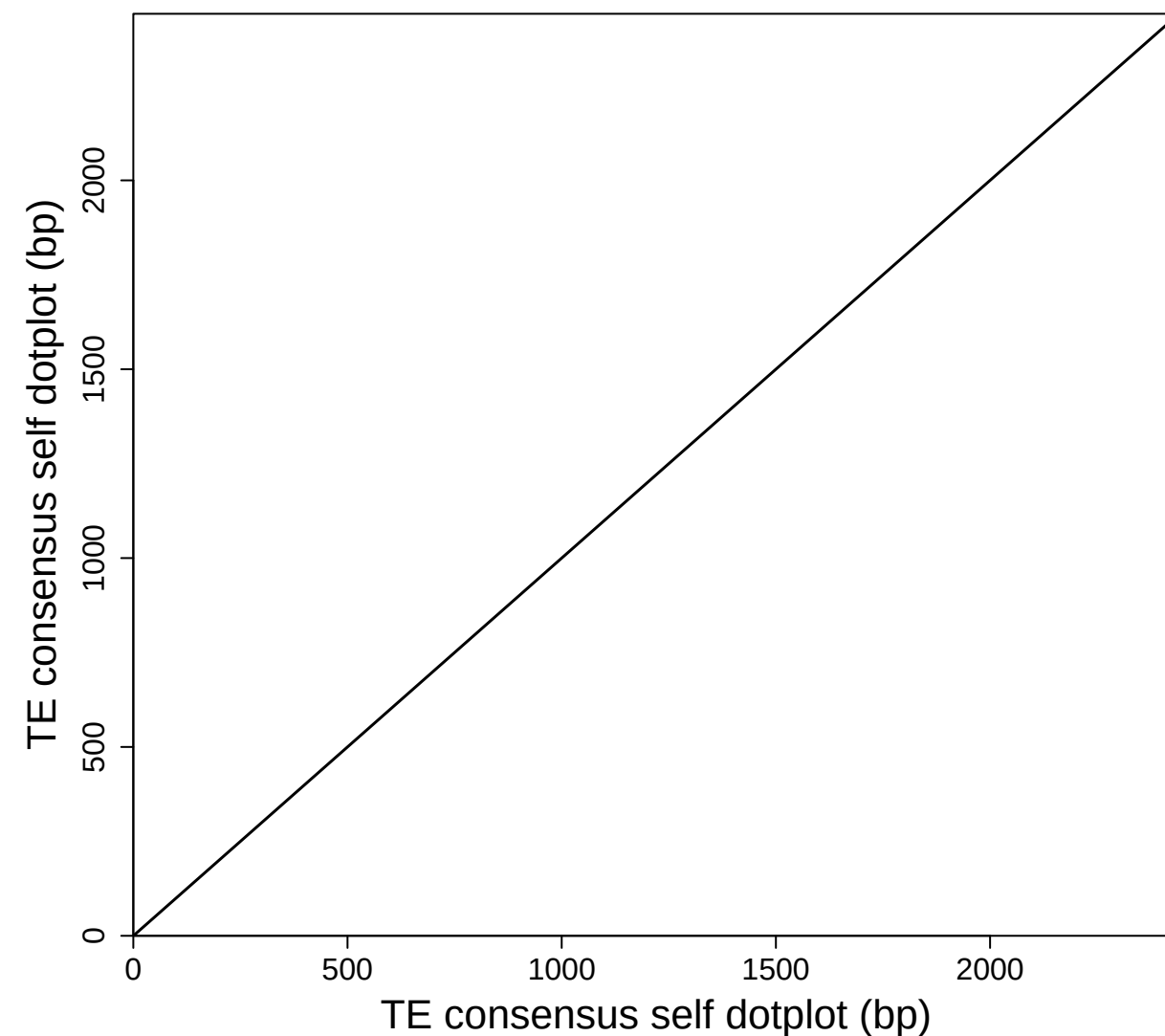
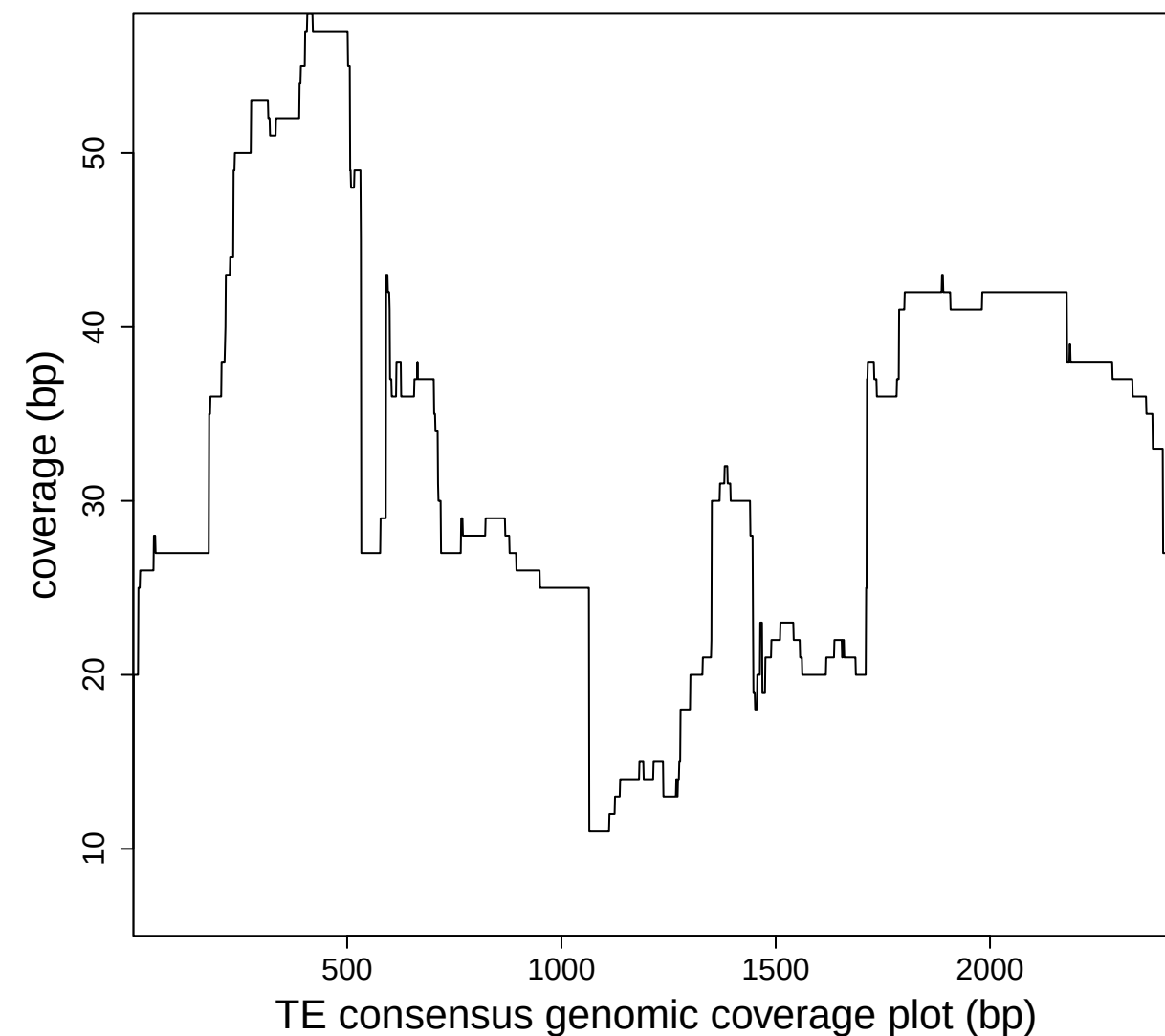
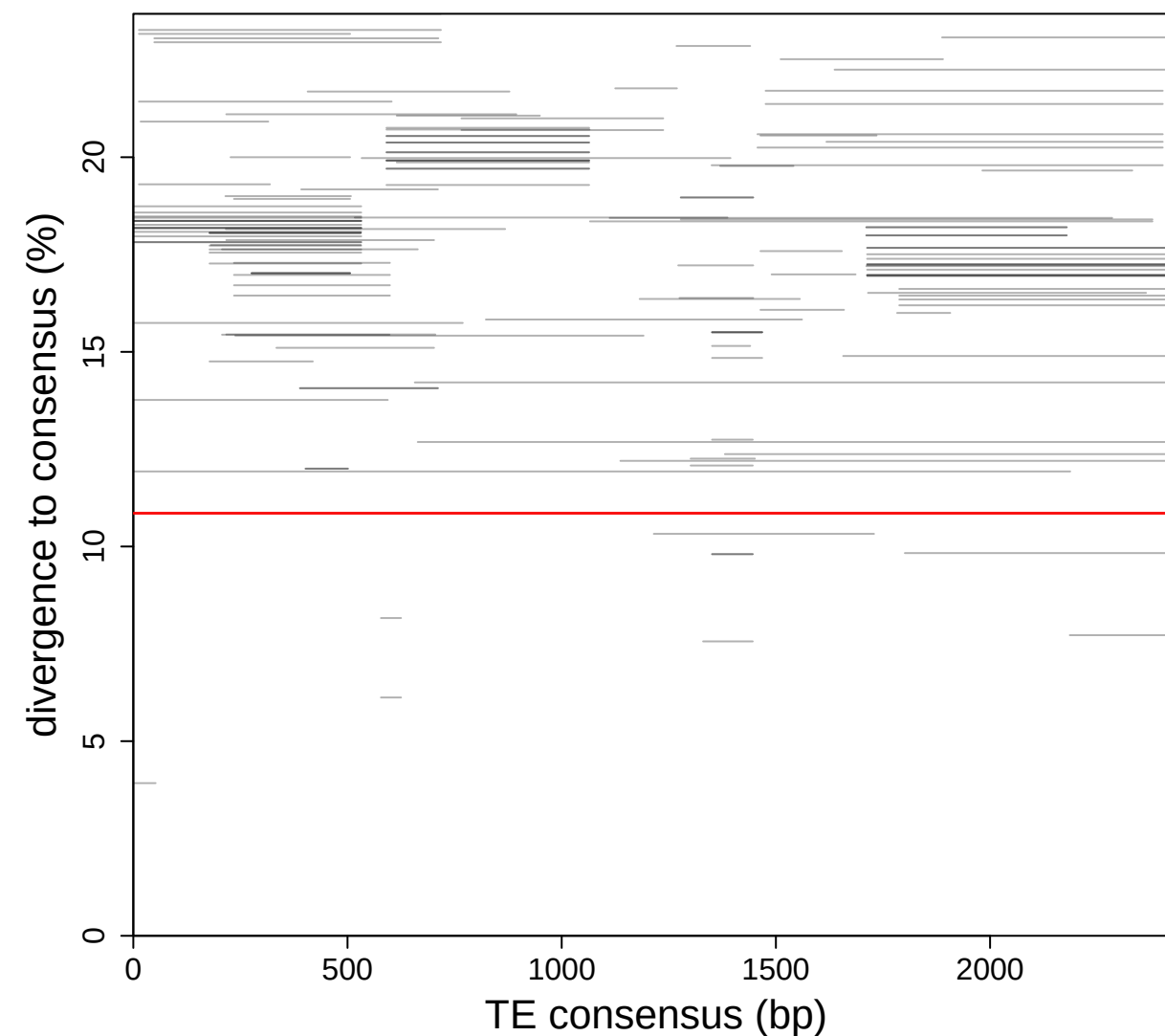
ly_9737.fasta.b.bed uf.bed g_3.bed fm_1.bed 0_0 bcln.fa aln.f
size: 10133bp; fragments: 38007; full length: 0 (>=9119.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

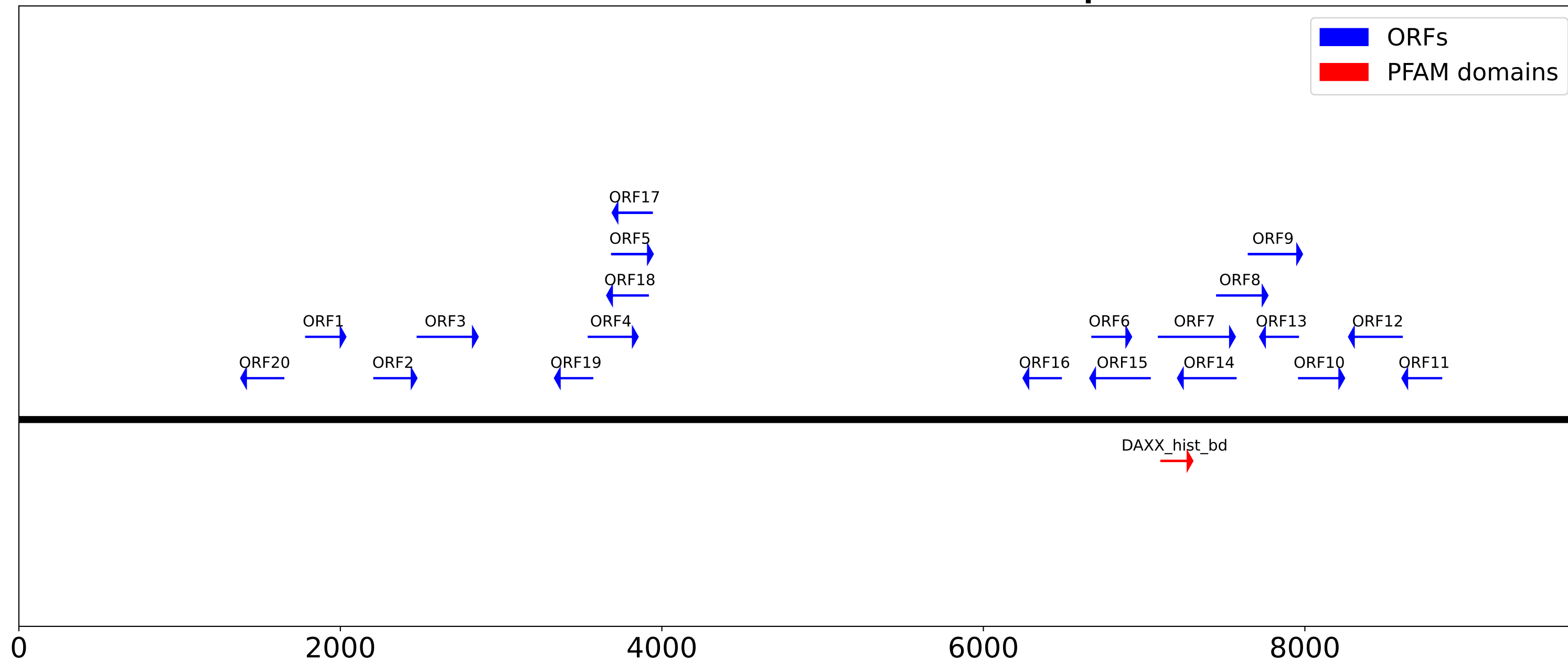


TE: rnd_5_family_9737
size: 2441bp; fragments: 151; full length: 1 (≥ 2196.9 bp)

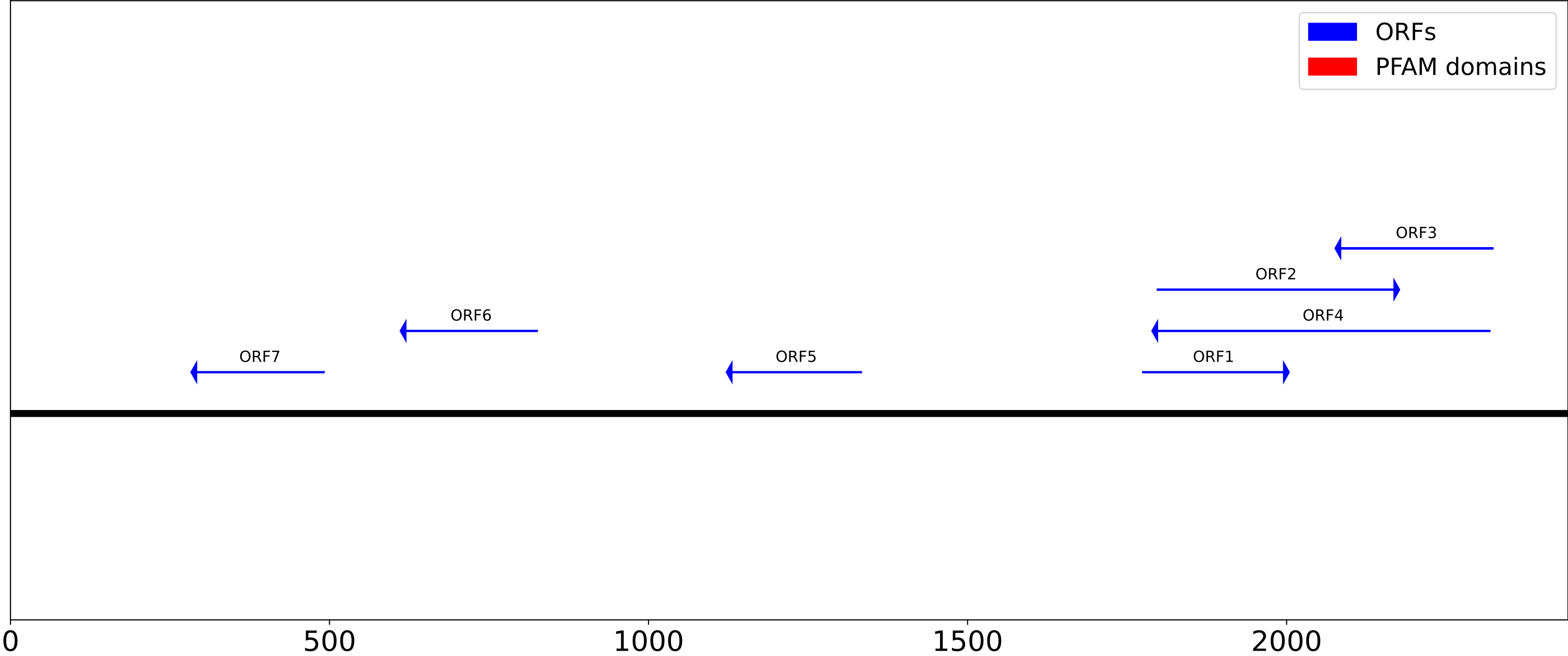
Before TEtrimmer 2441 bp



After TETrimmer ORF and PFAM domain plot



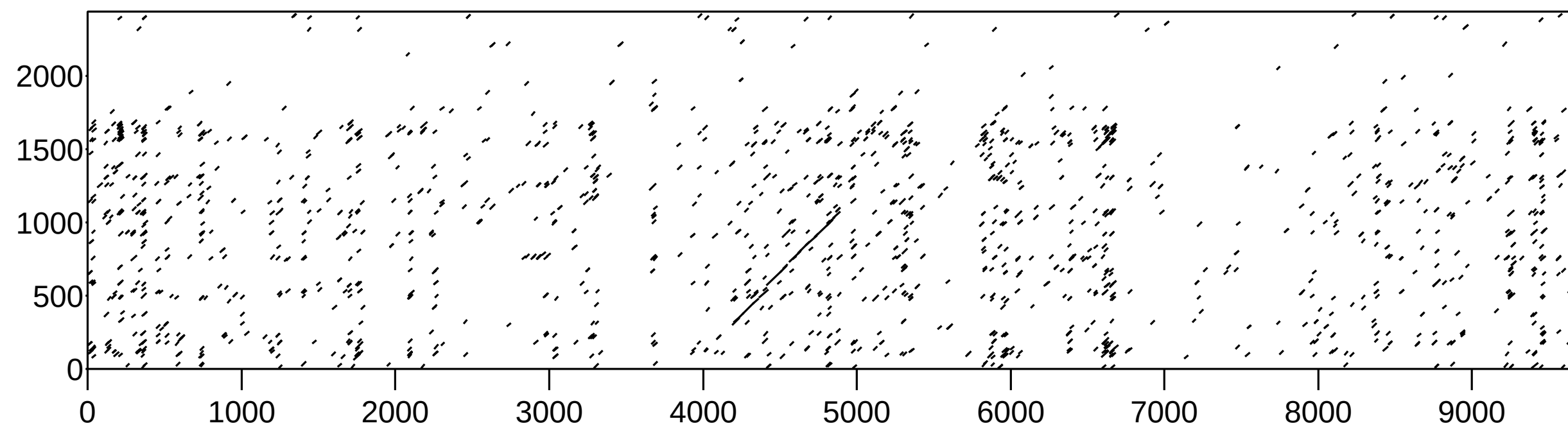
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)