

Start Crop Point

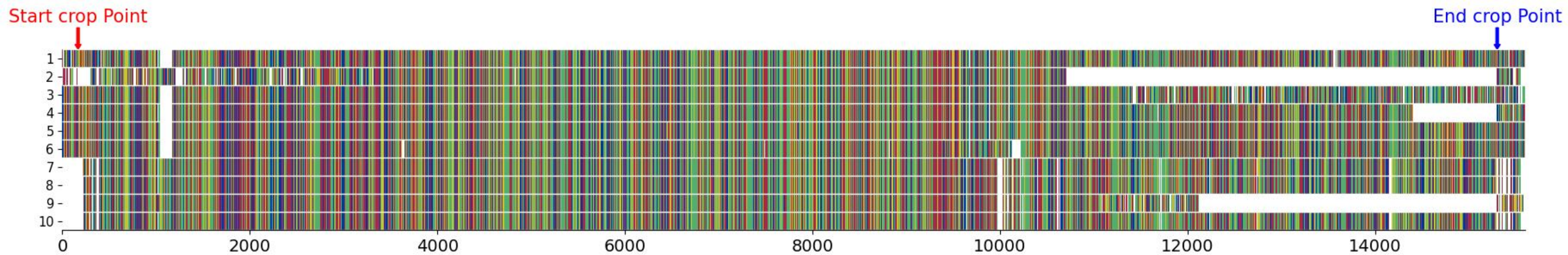


End Crop Point



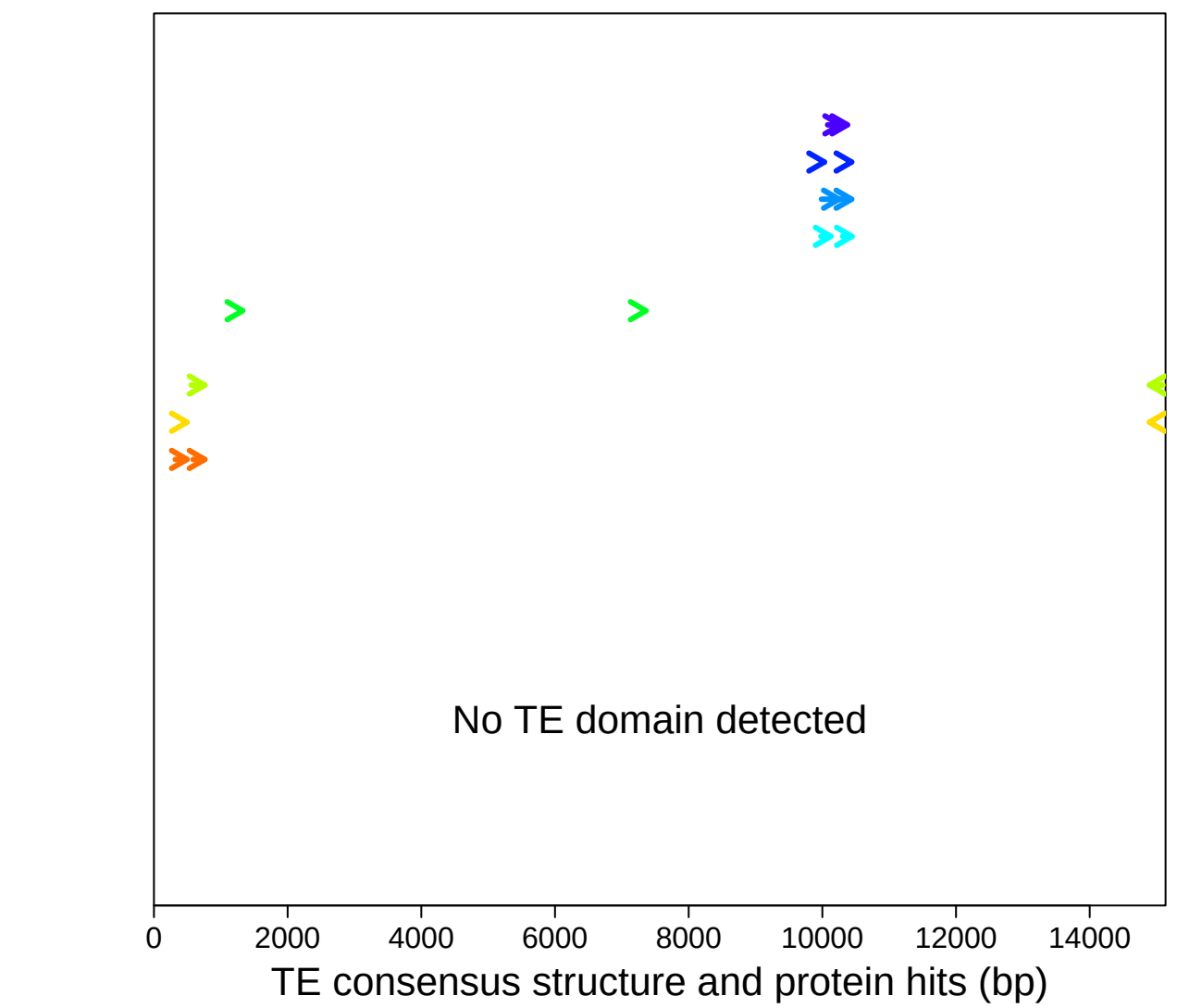
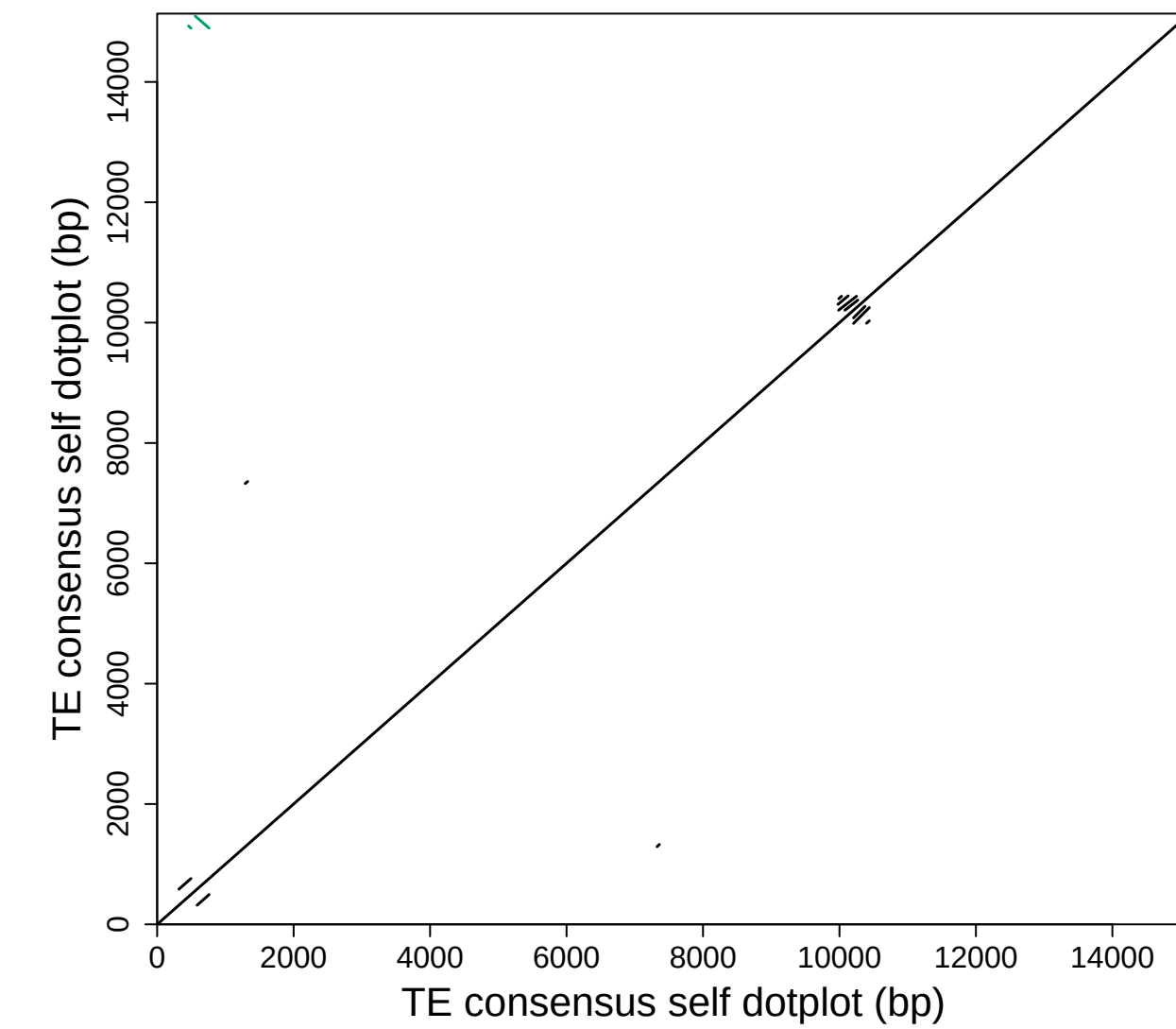
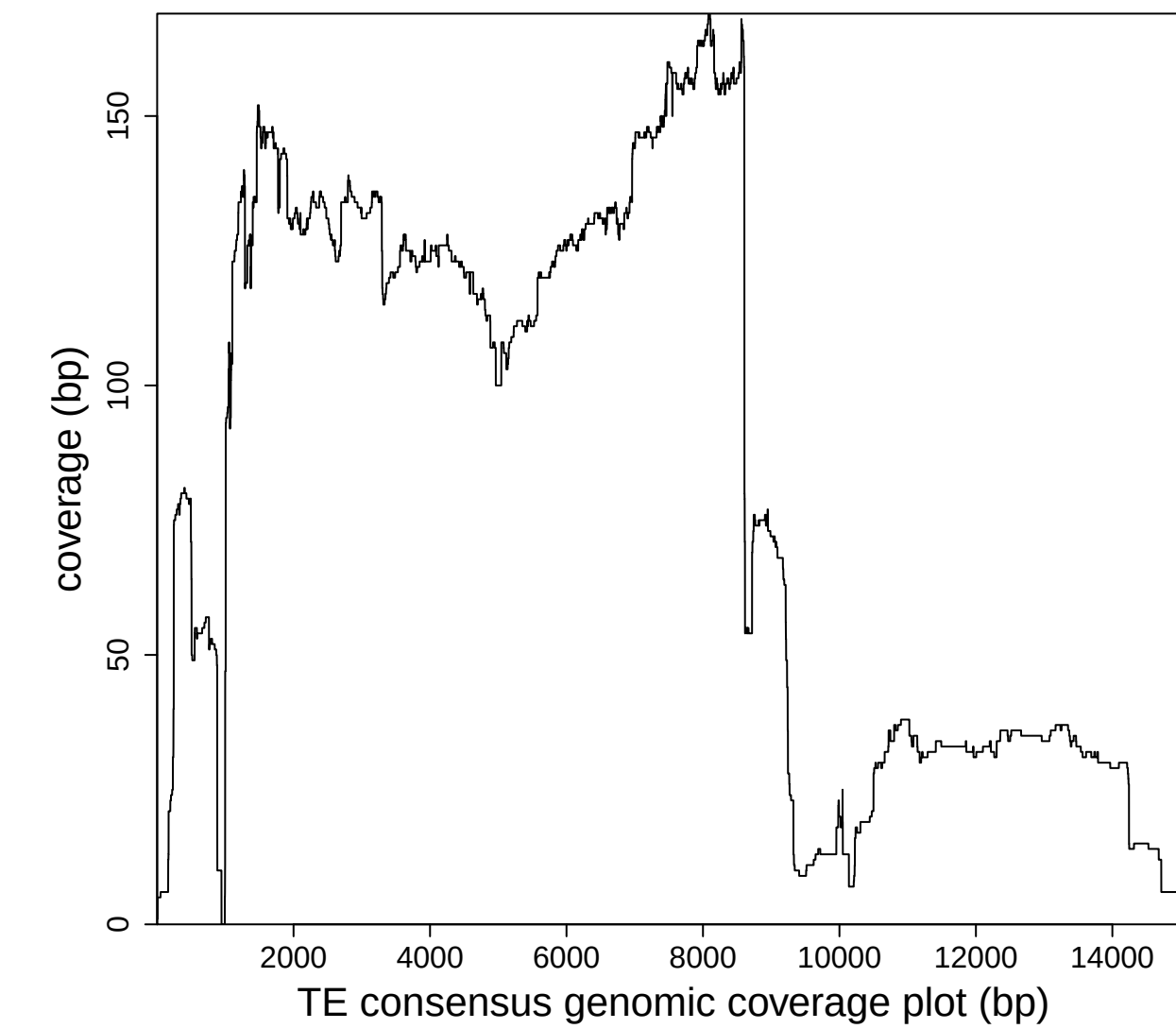
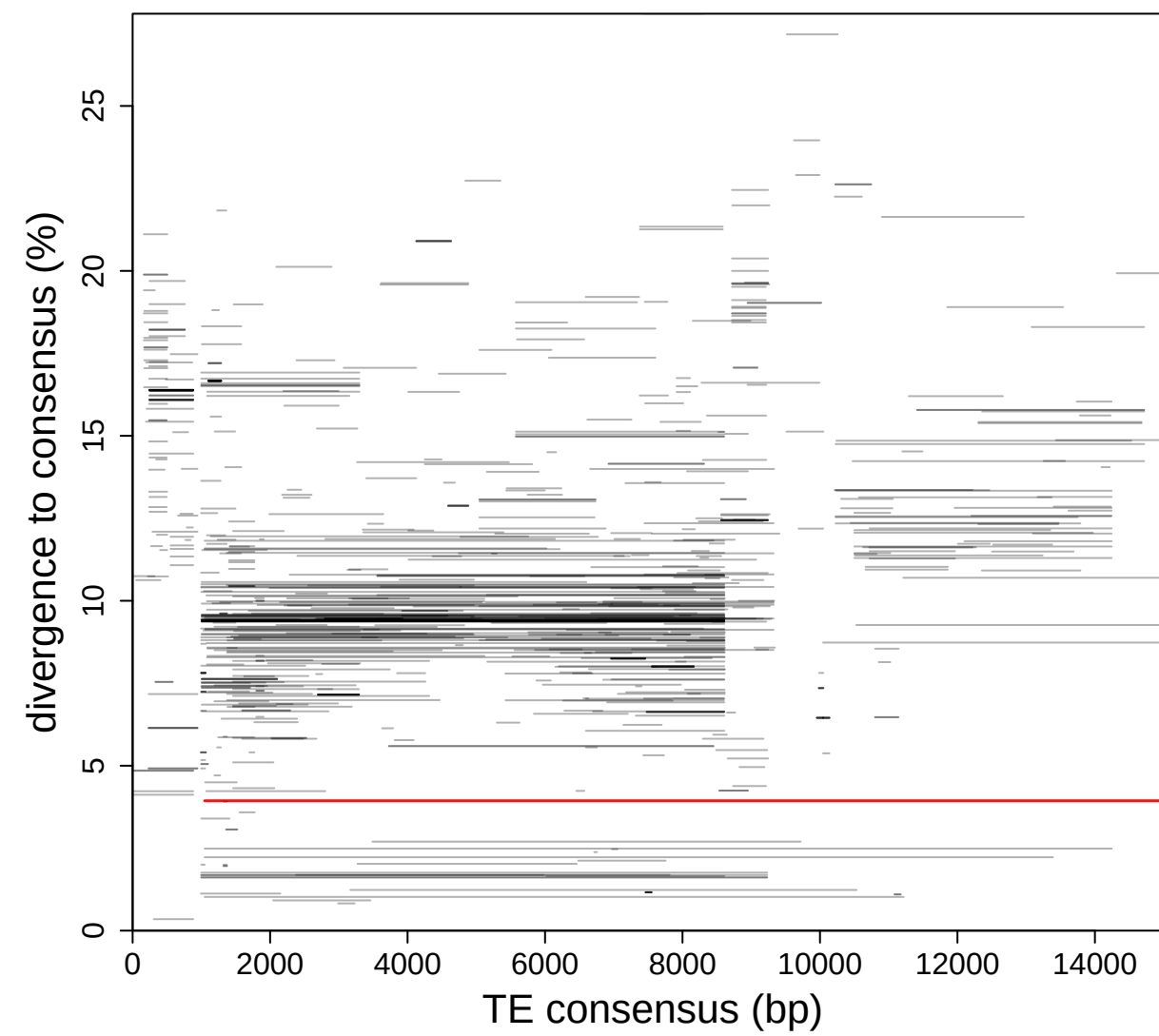
MSA length = 15135





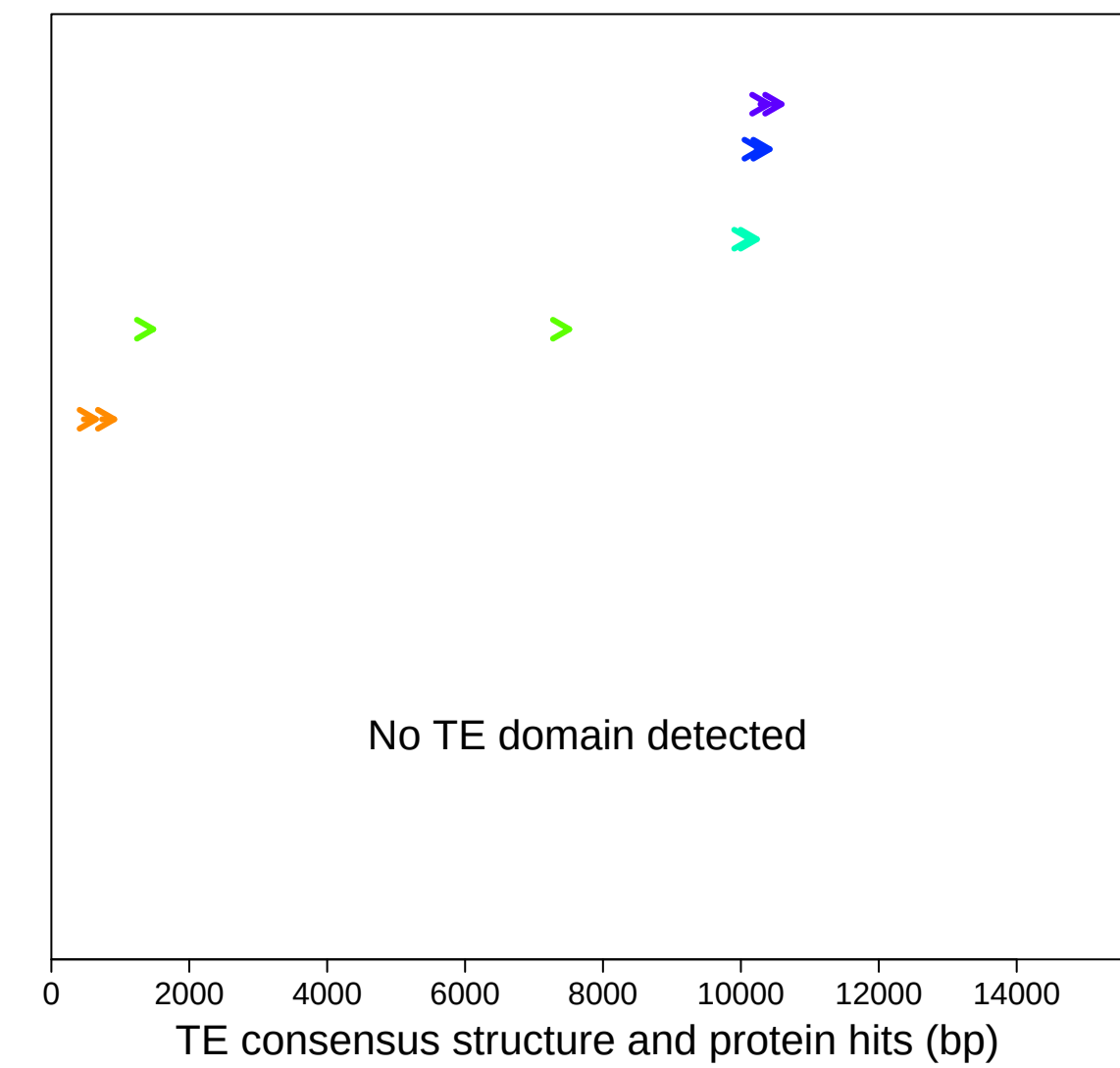
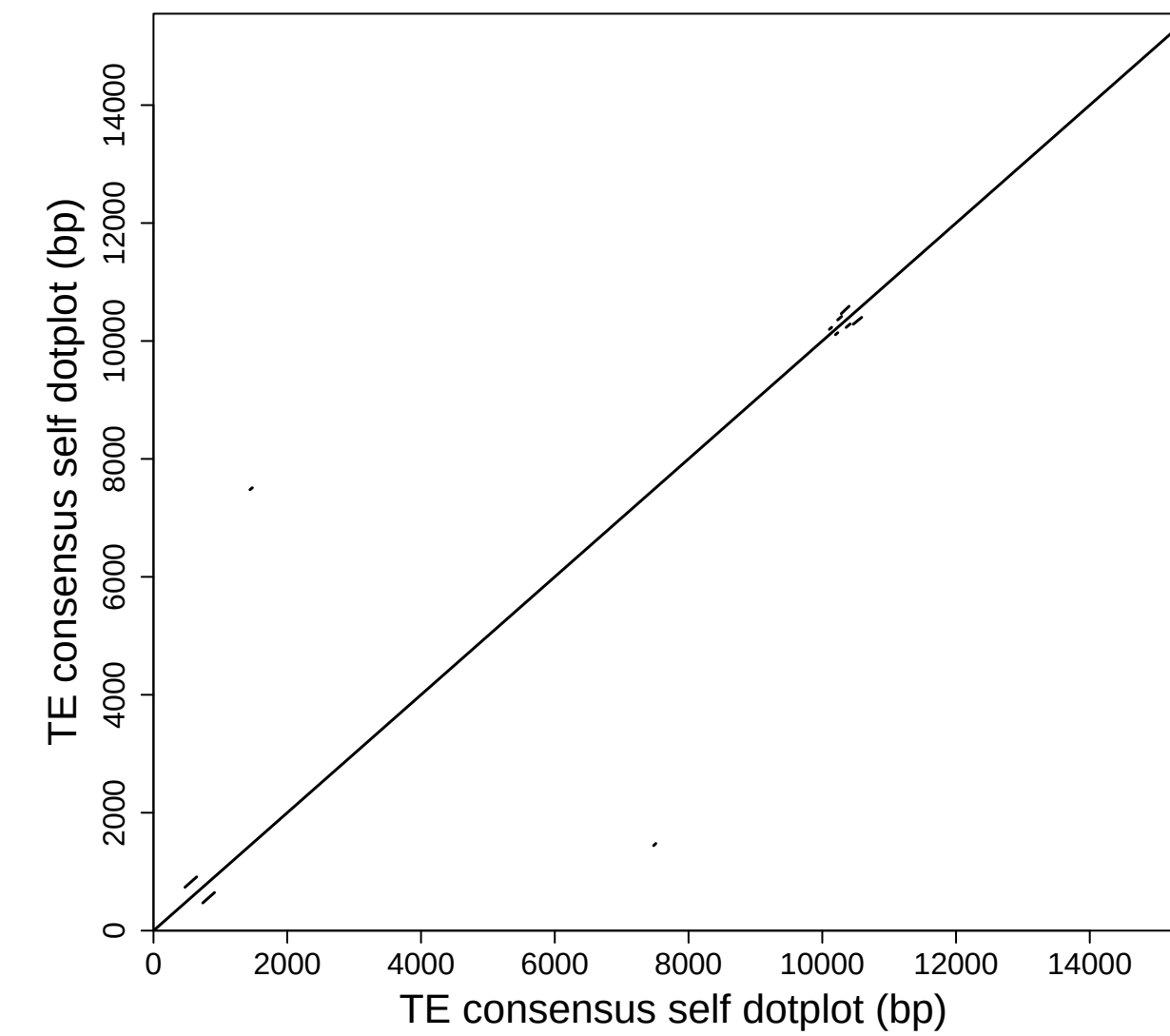
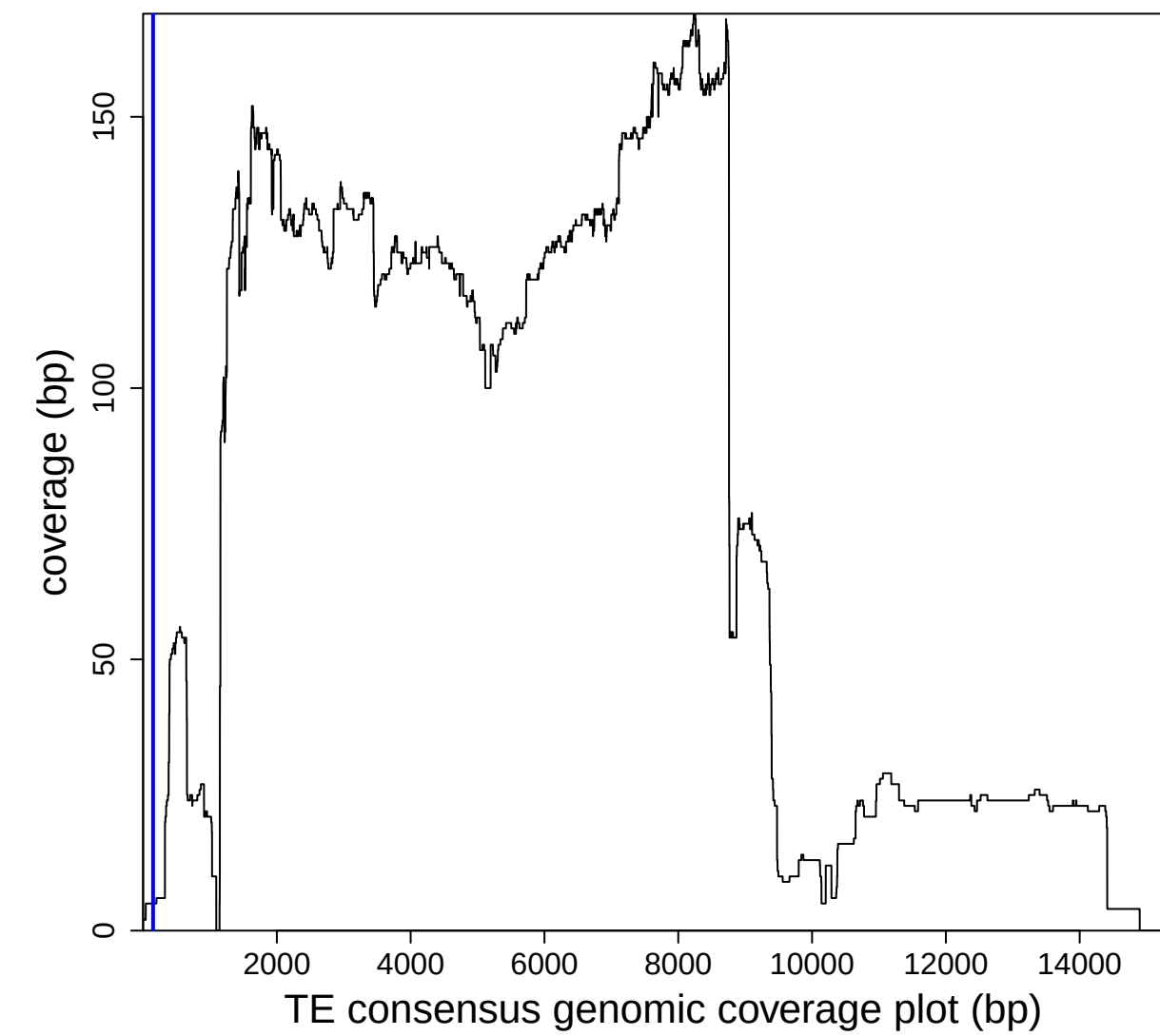
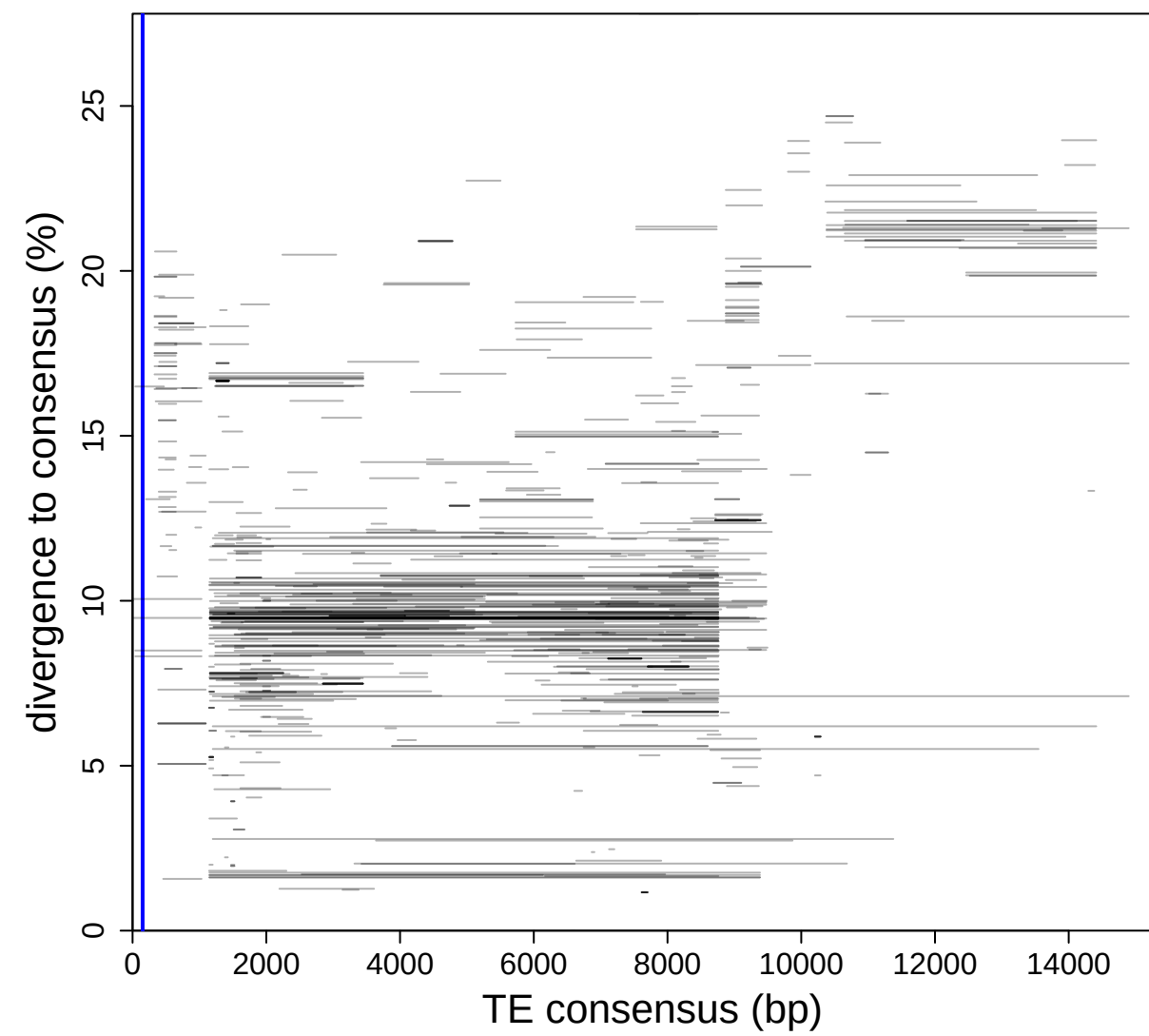
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 15135bp; fragments: 763; full length: 1 (>=13621.5bp)

After TEtrimmer 15135 bp



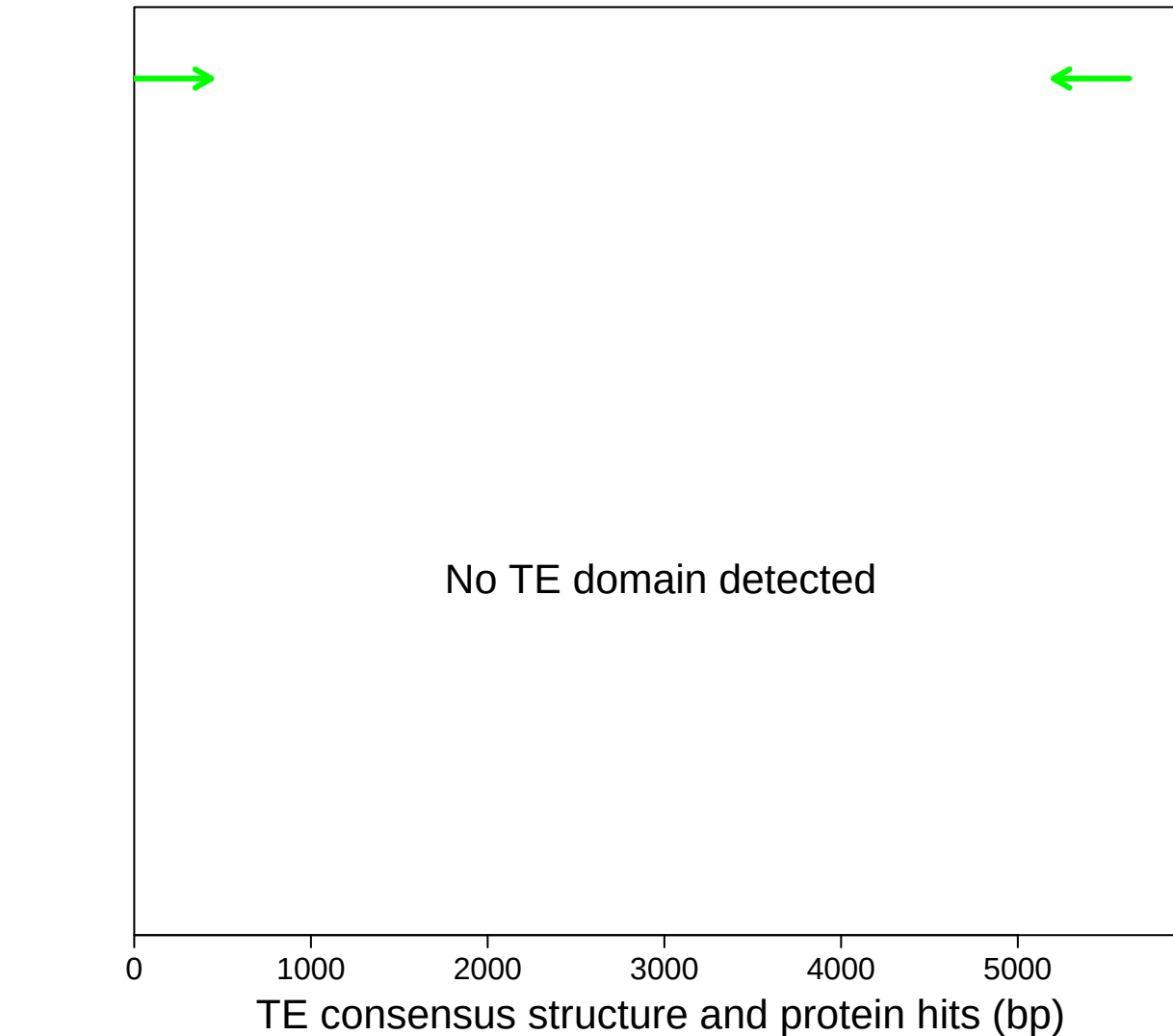
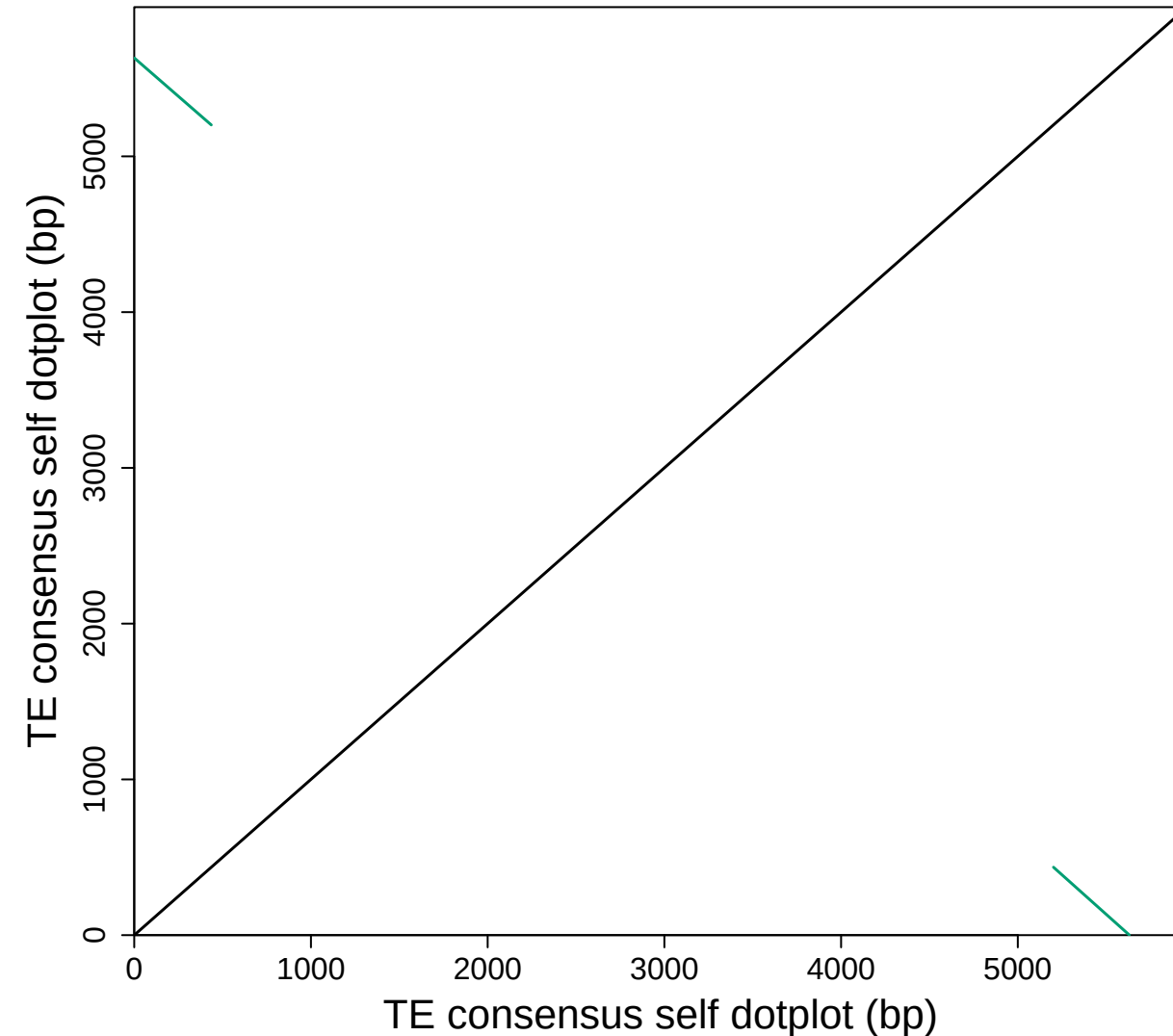
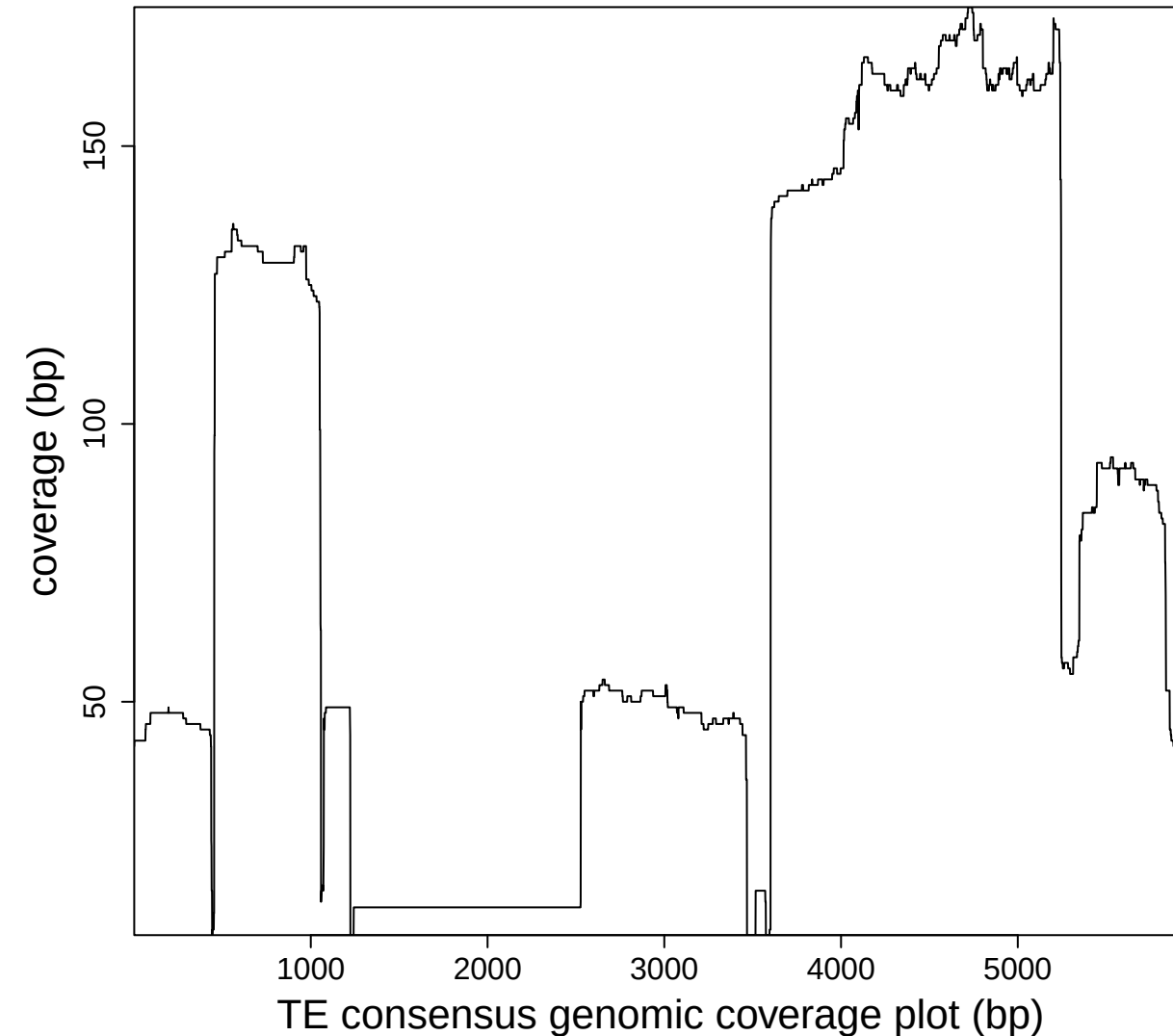
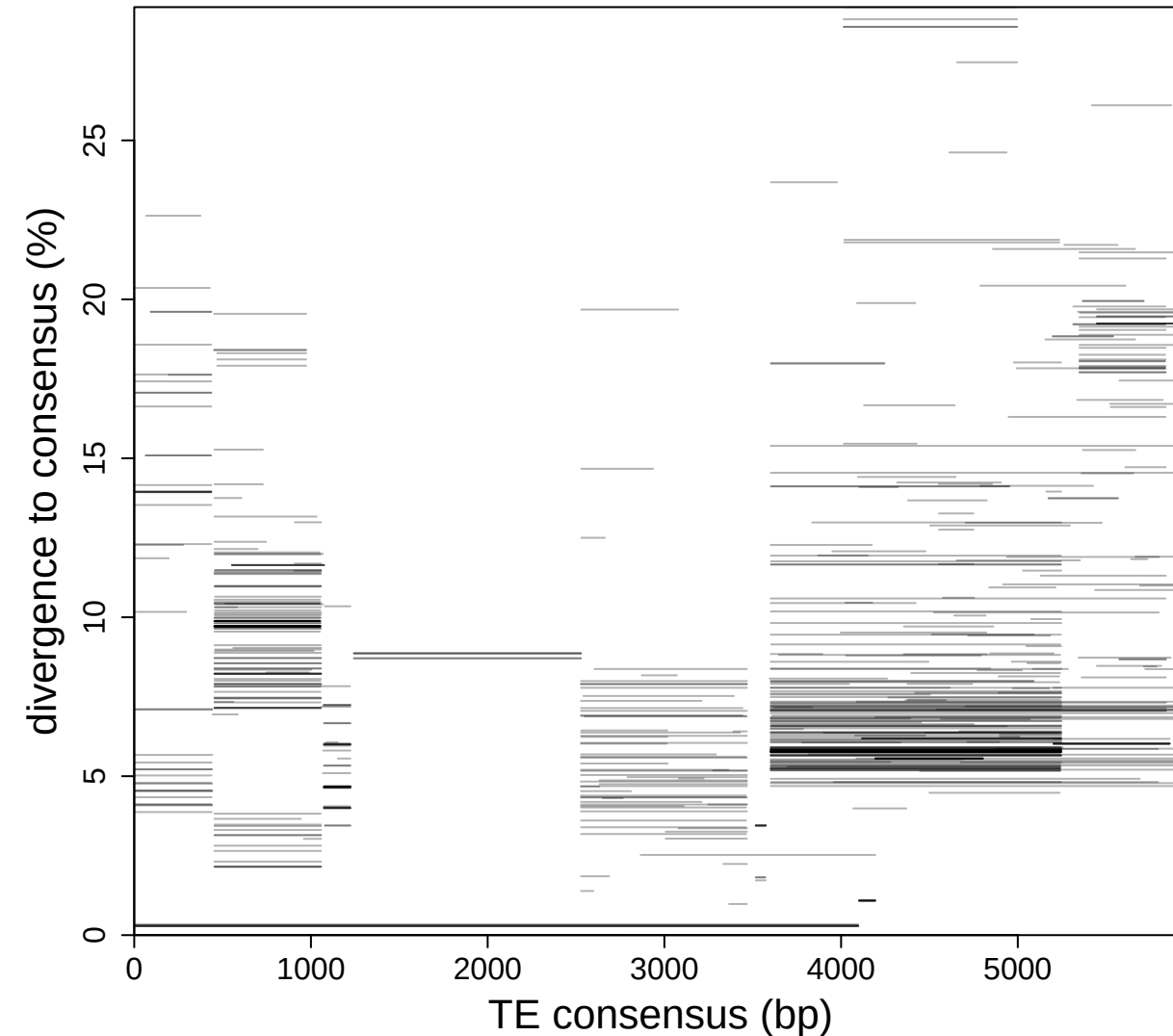
3.fasta.b.bed uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 15551bp; fragments: 691; full length: 0 (>=13995.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

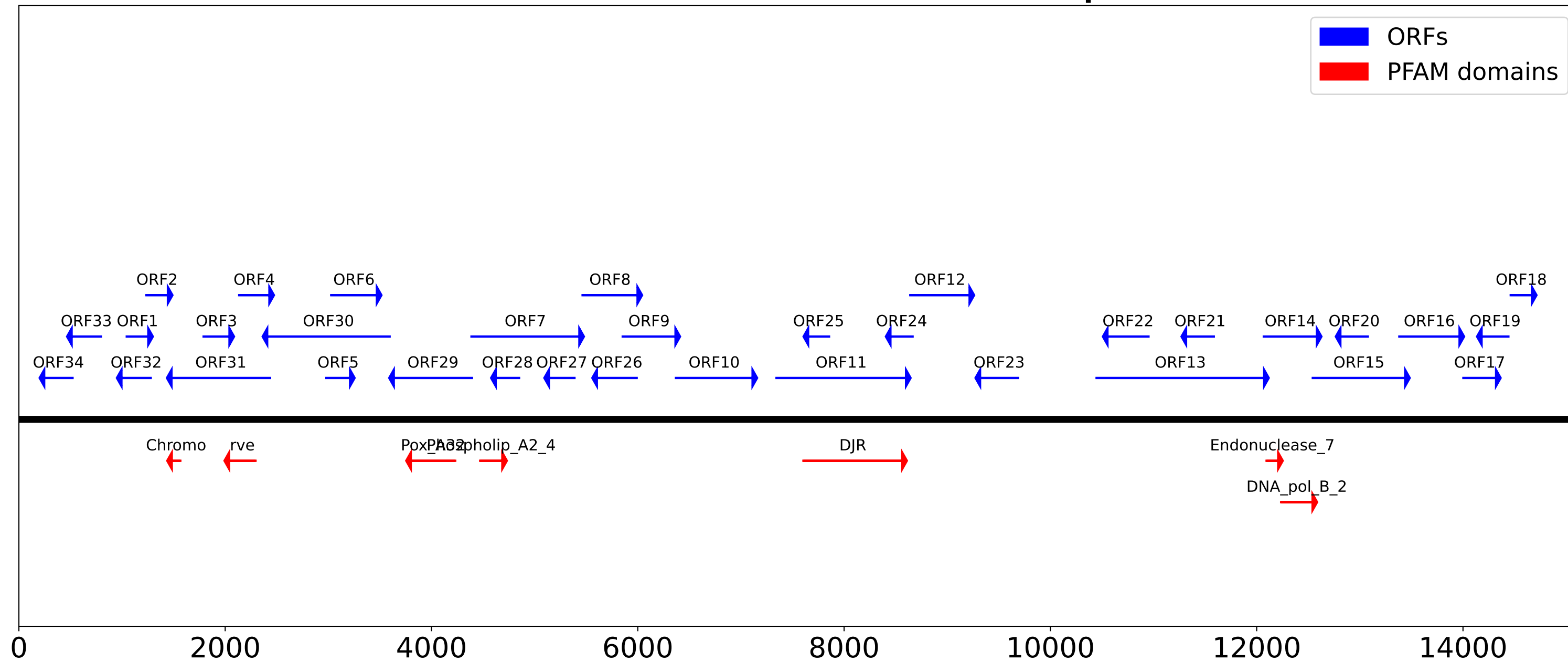


TE: rnd_5_family_3973
size: 5958bp; fragments: 598; full length: 0 (>=5362.2bp)

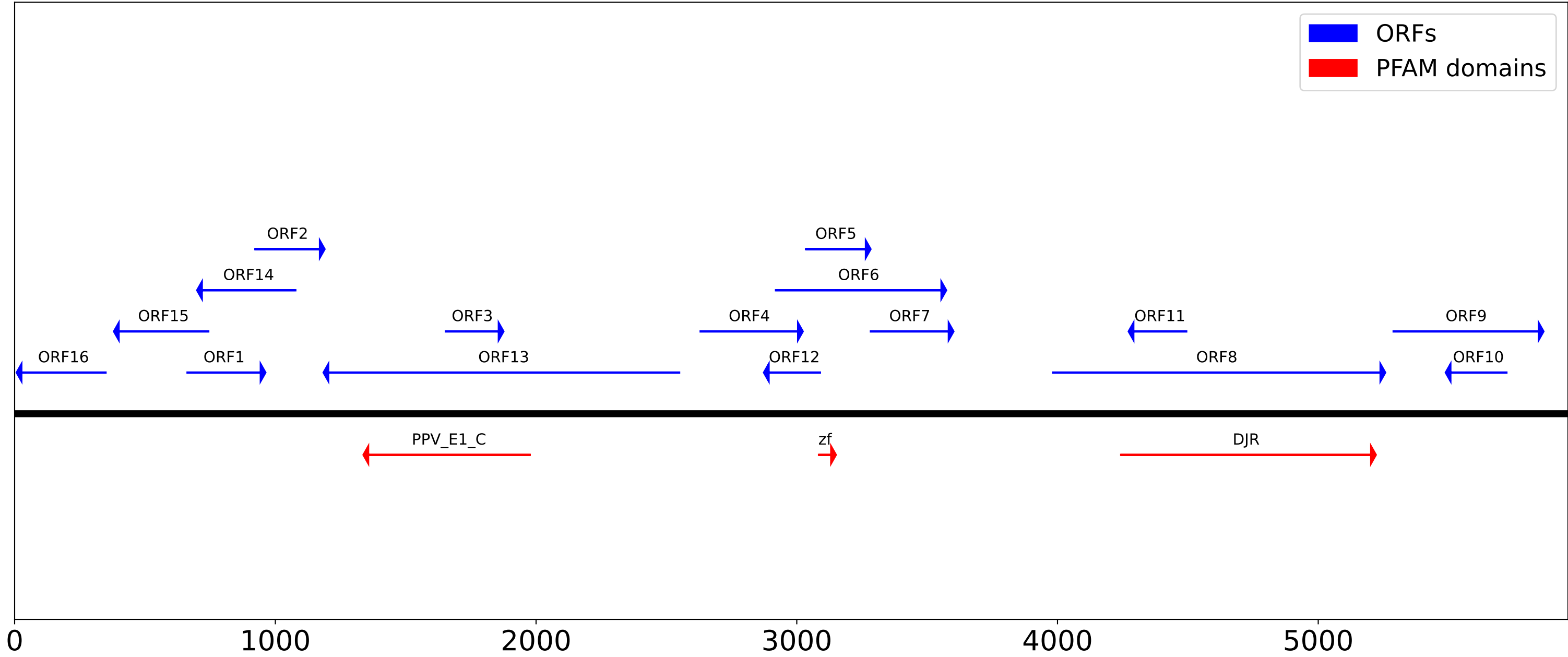
Before TEtrimmer 5958 bp



After TETrimmer ORF and PFAM domain plot



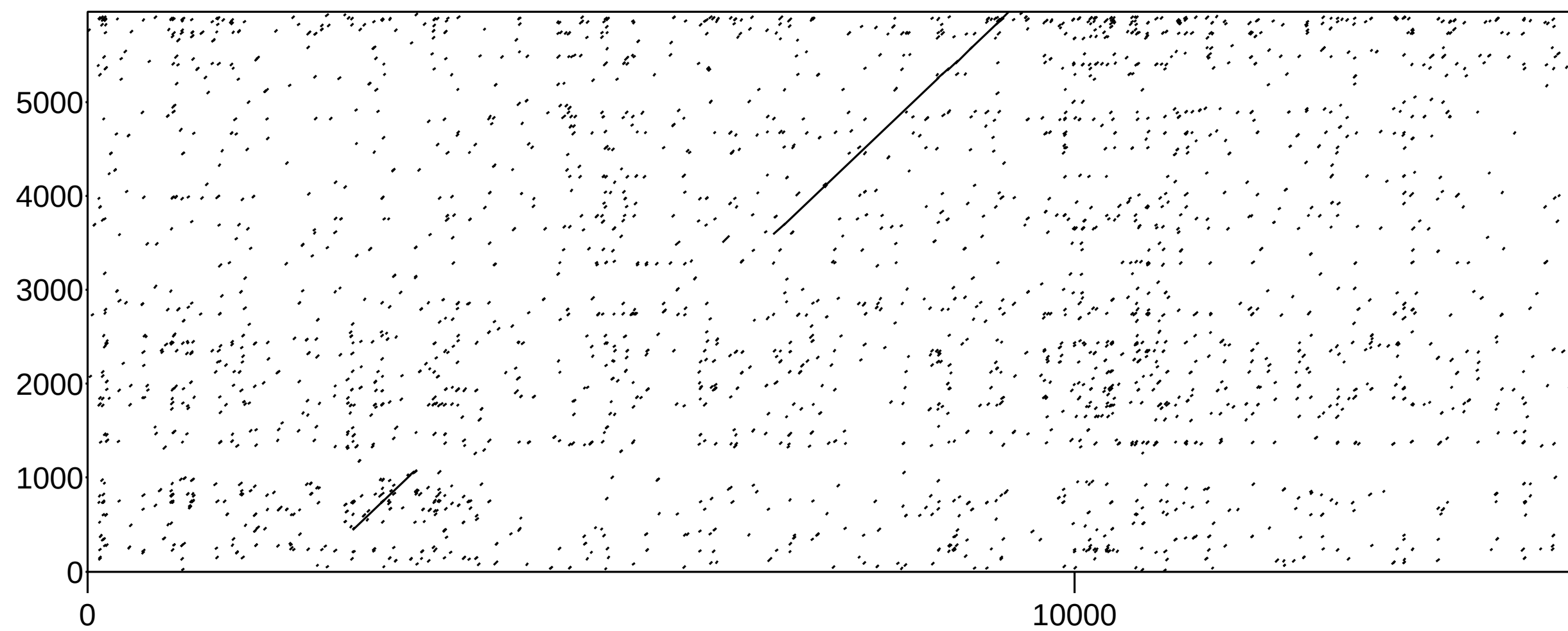
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)