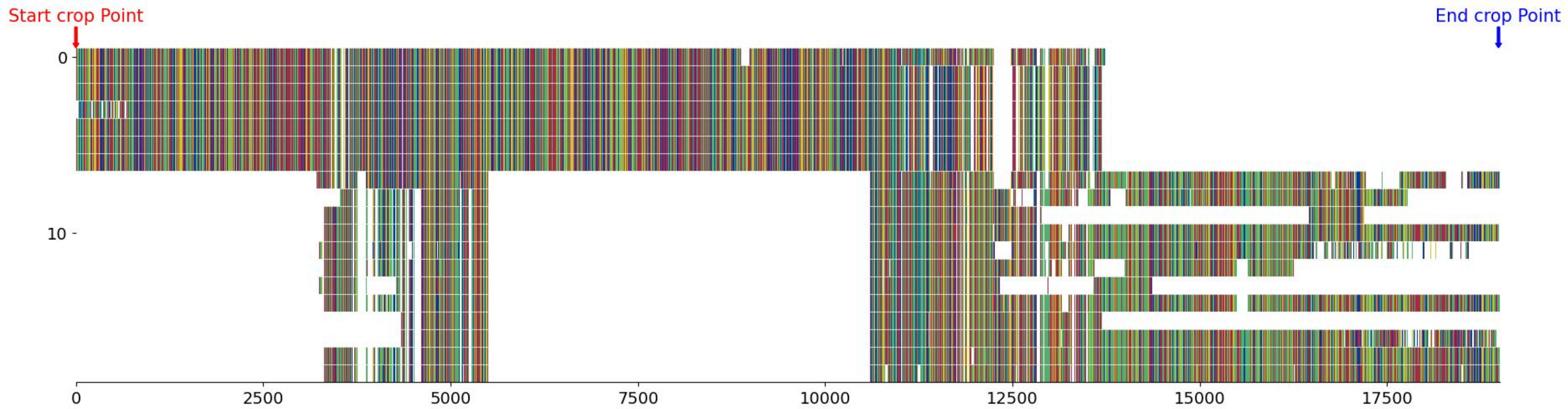


Start crop Point

End crop Point

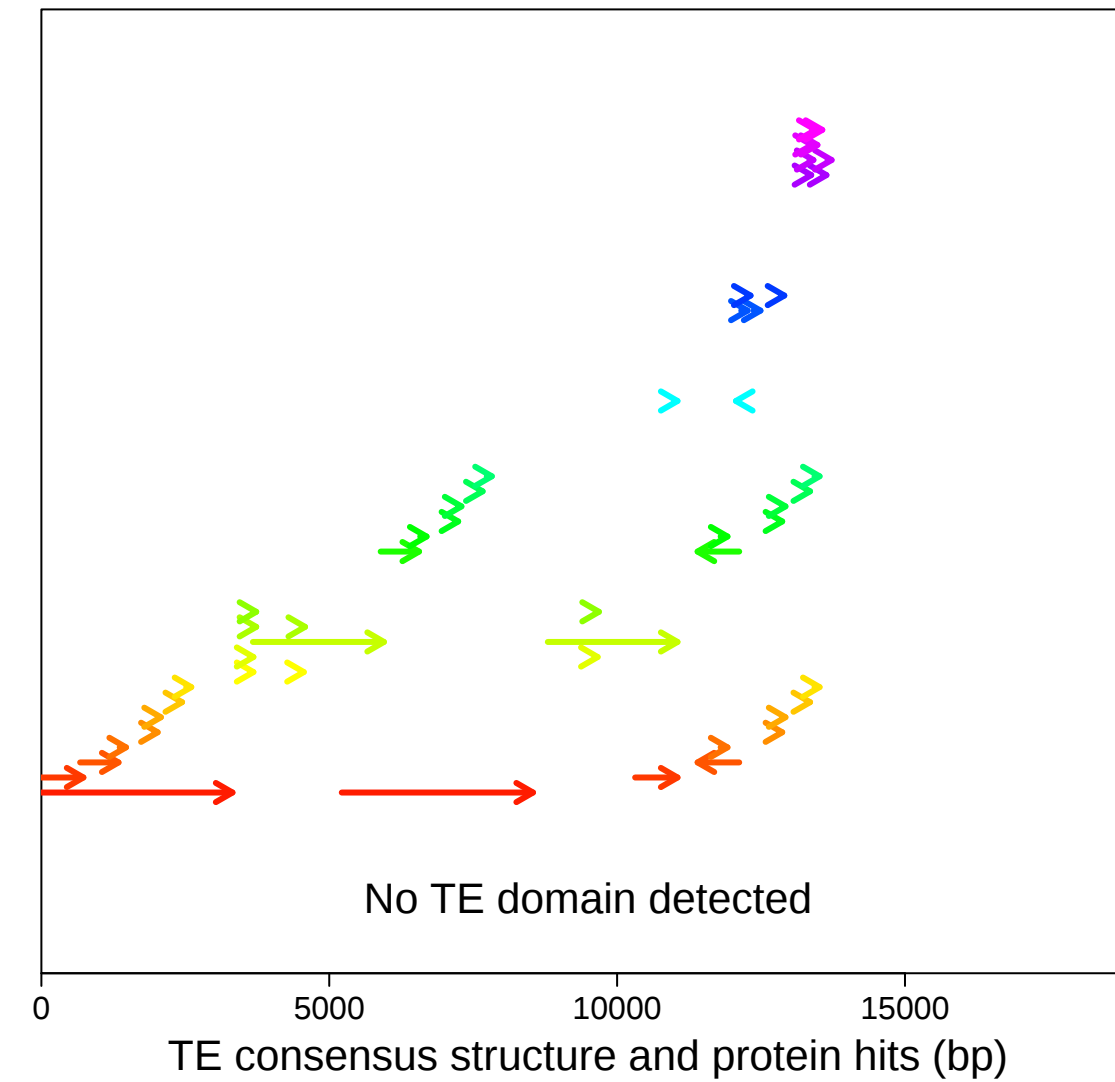
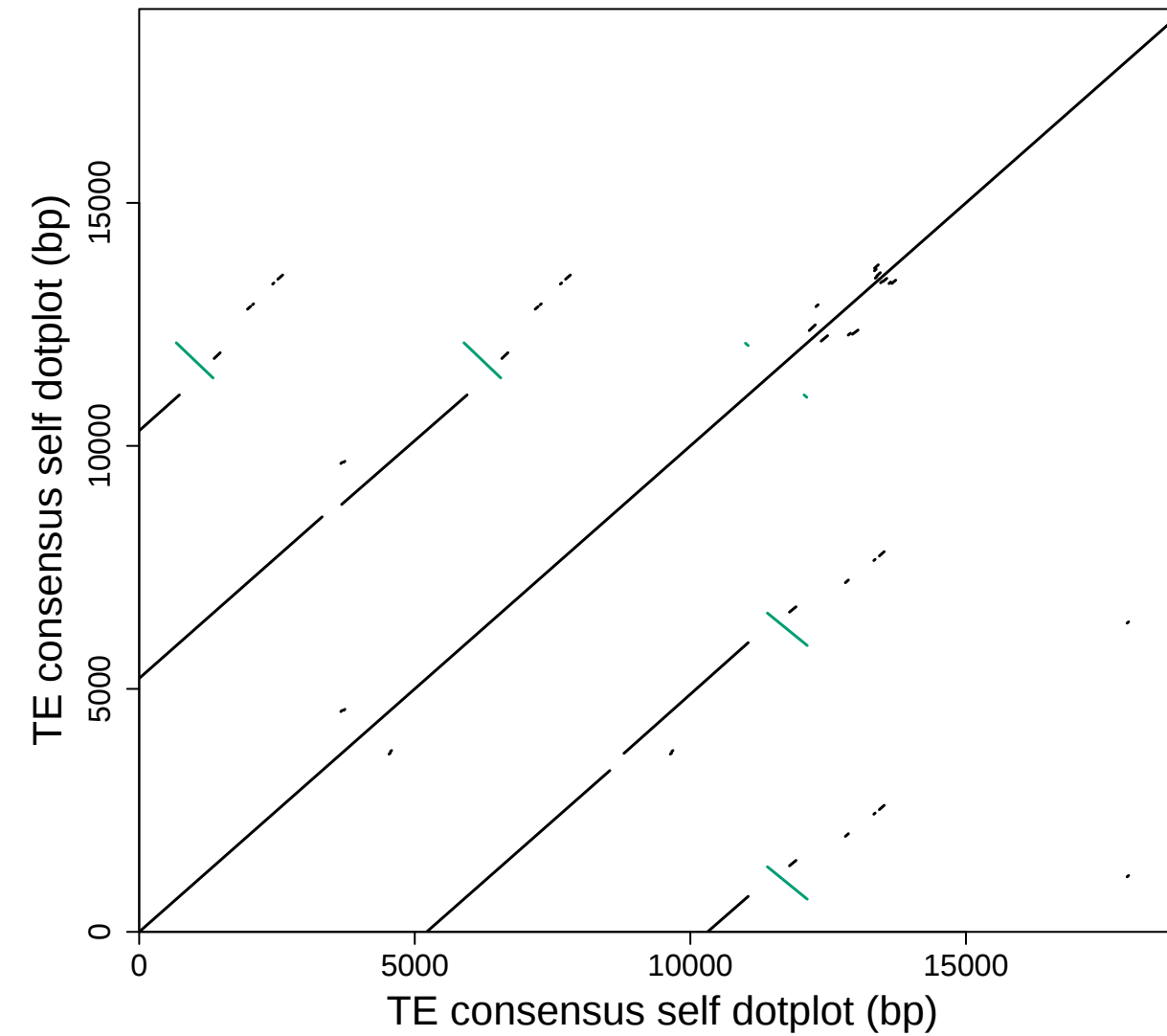
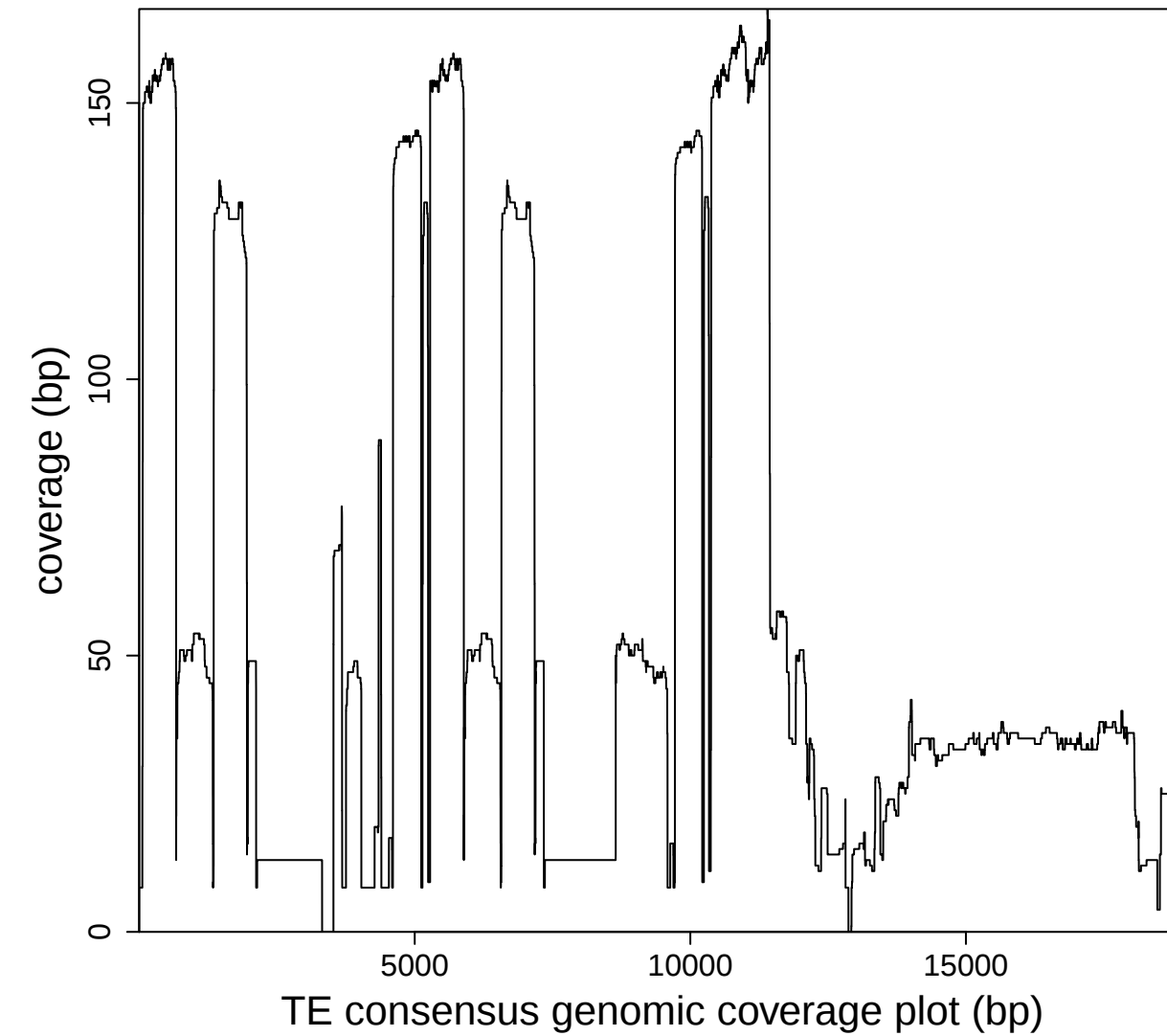
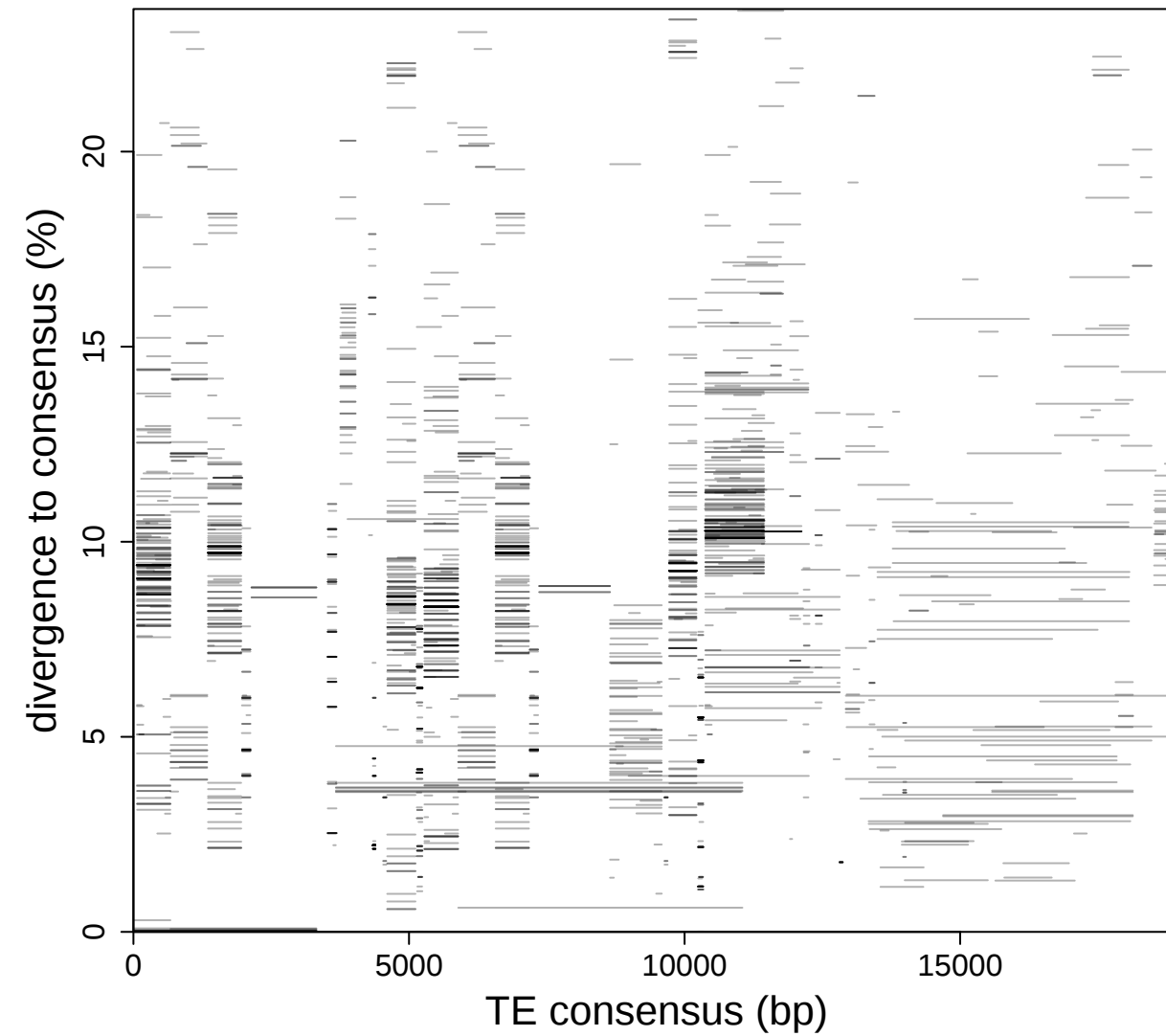
MSA length = 18989





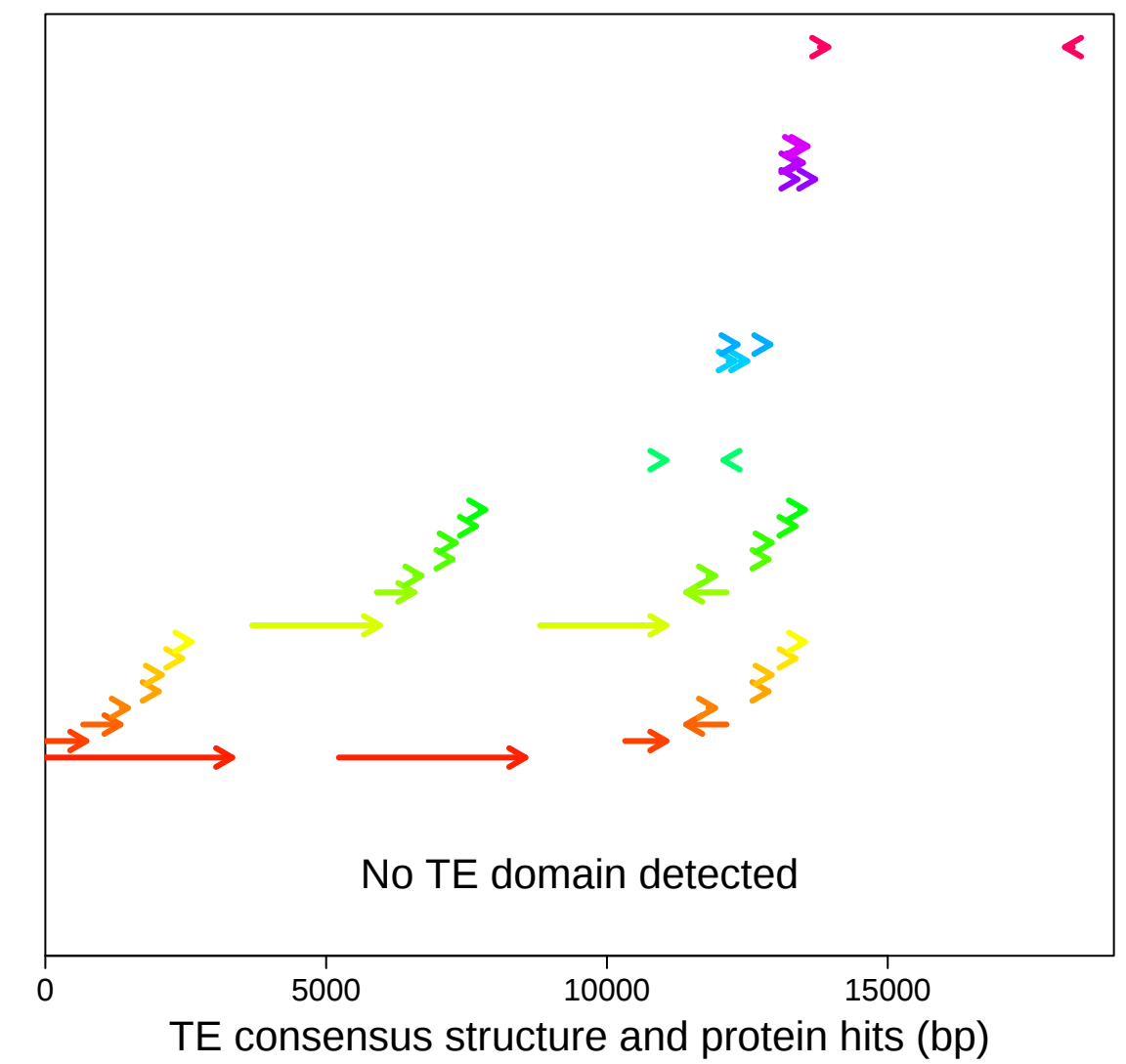
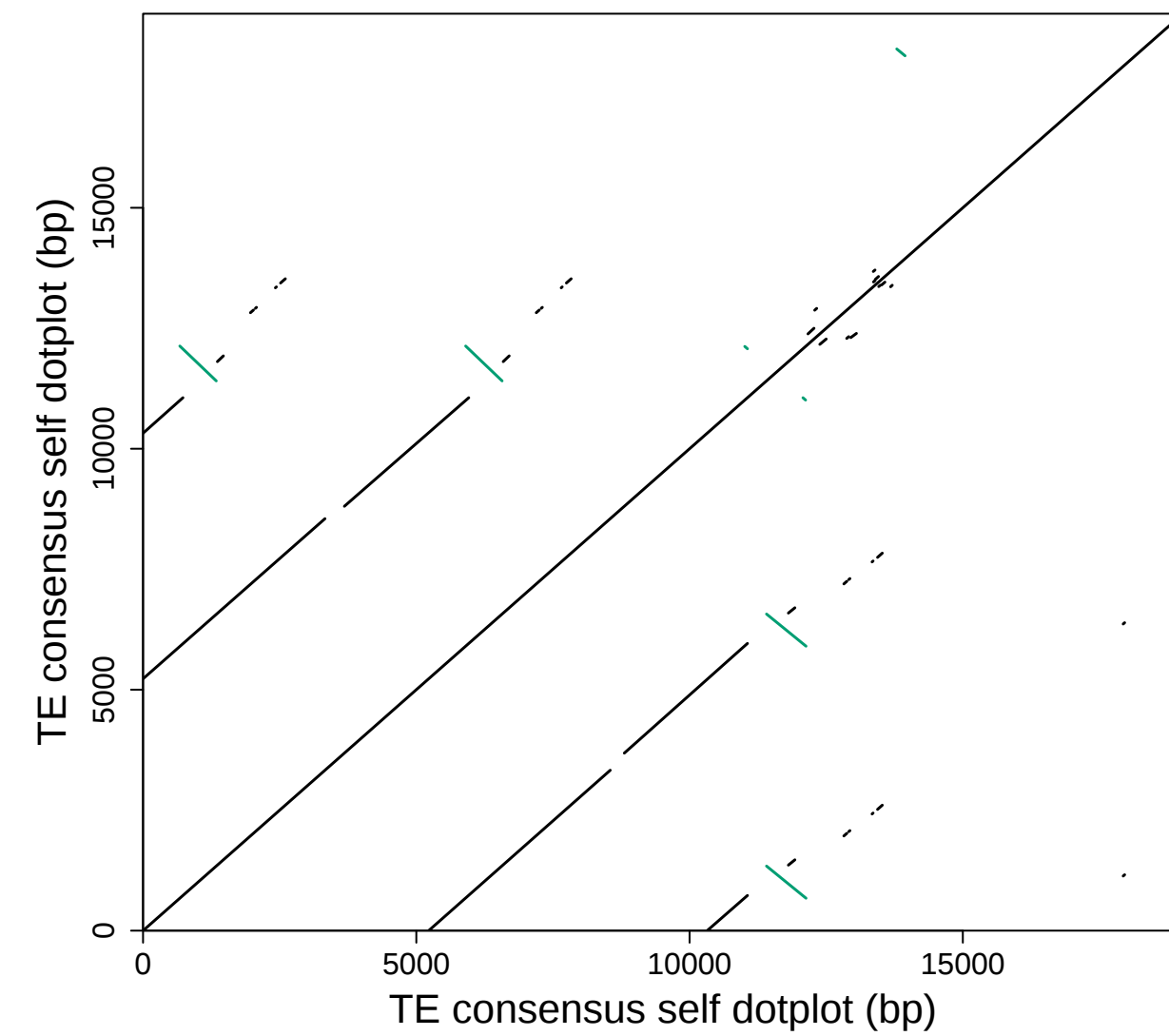
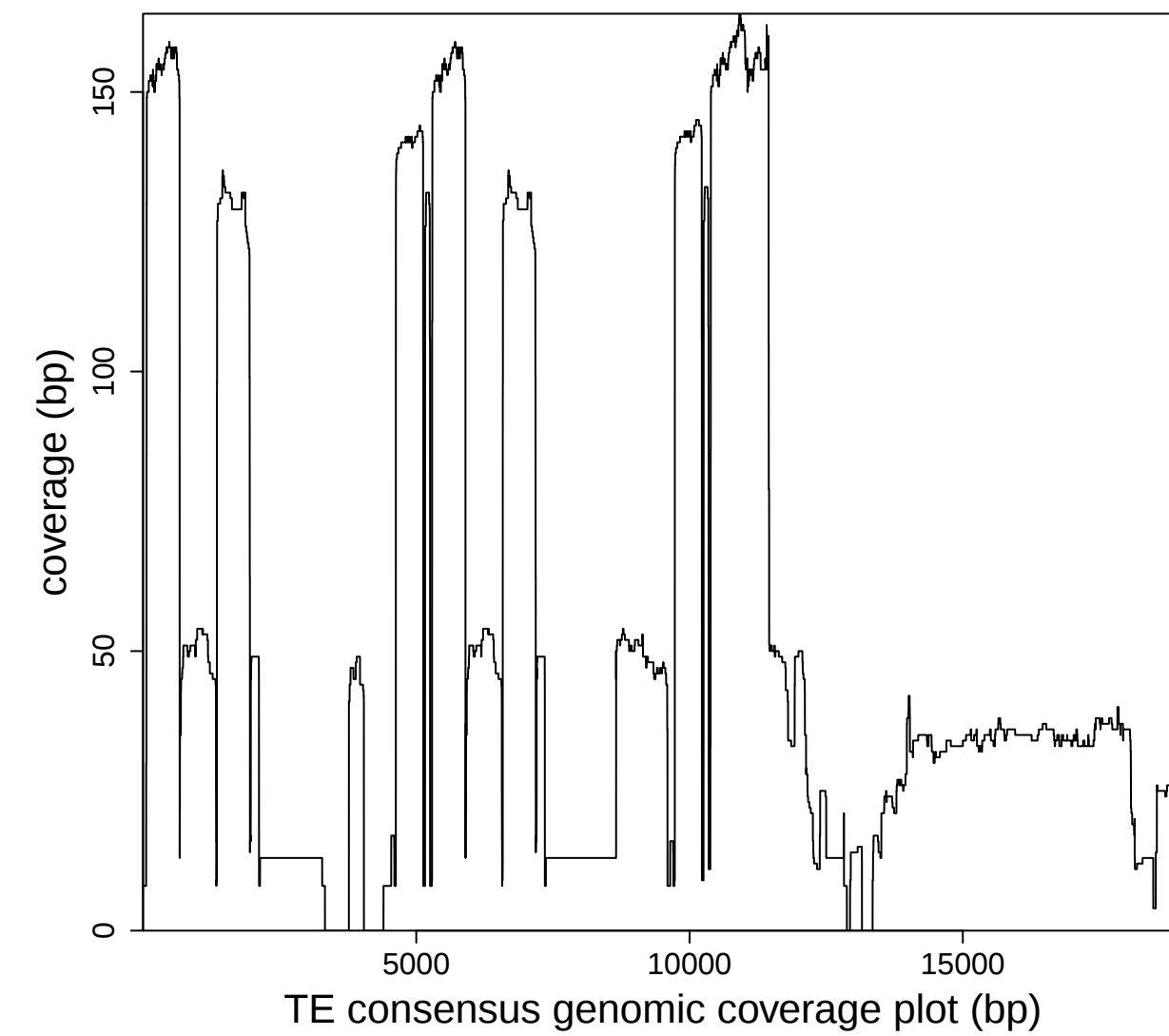
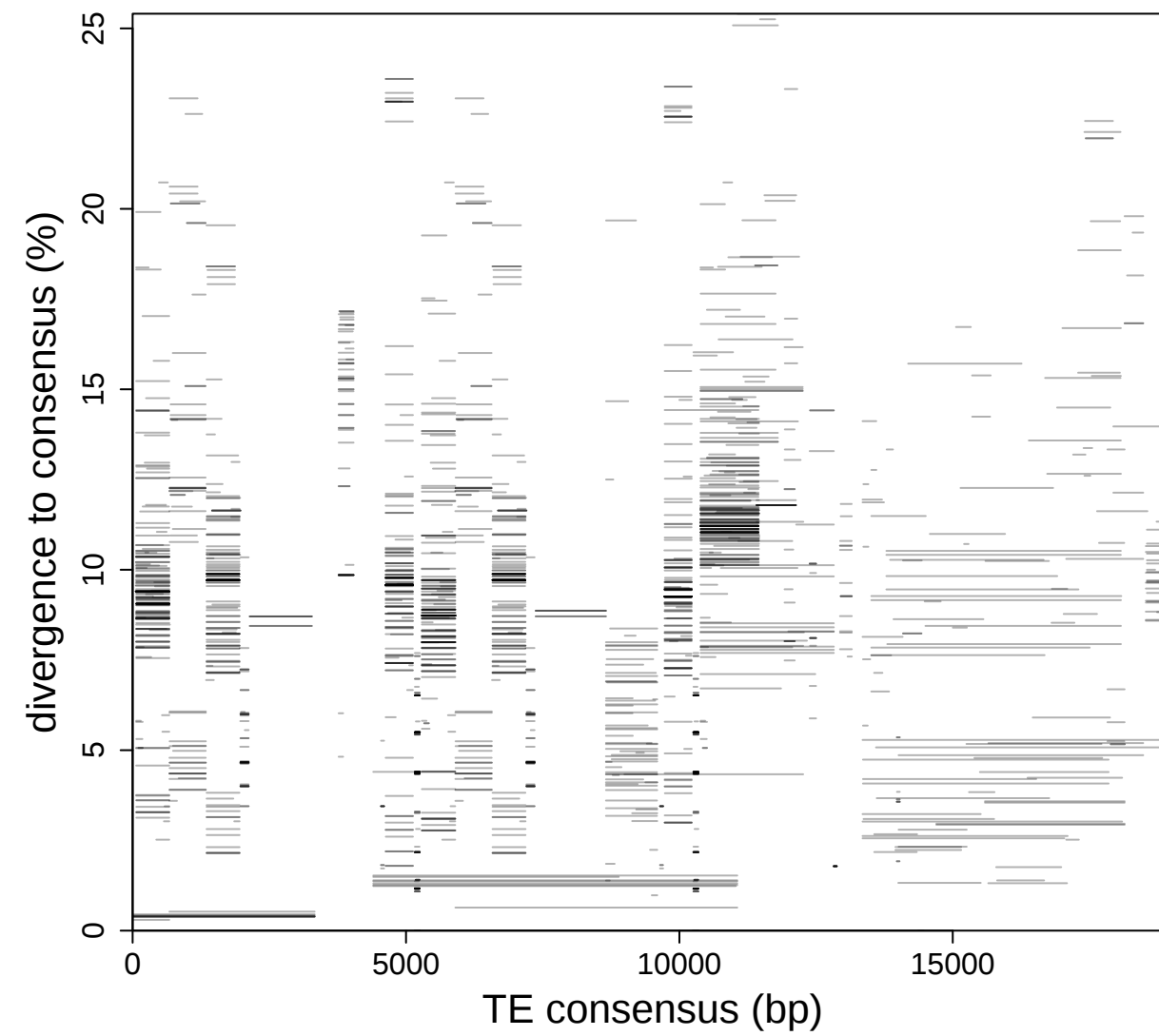
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 18989bp; fragments: 2051; full length: 0 (>=17090.1bp)

After TEtrimmer 18989 bp



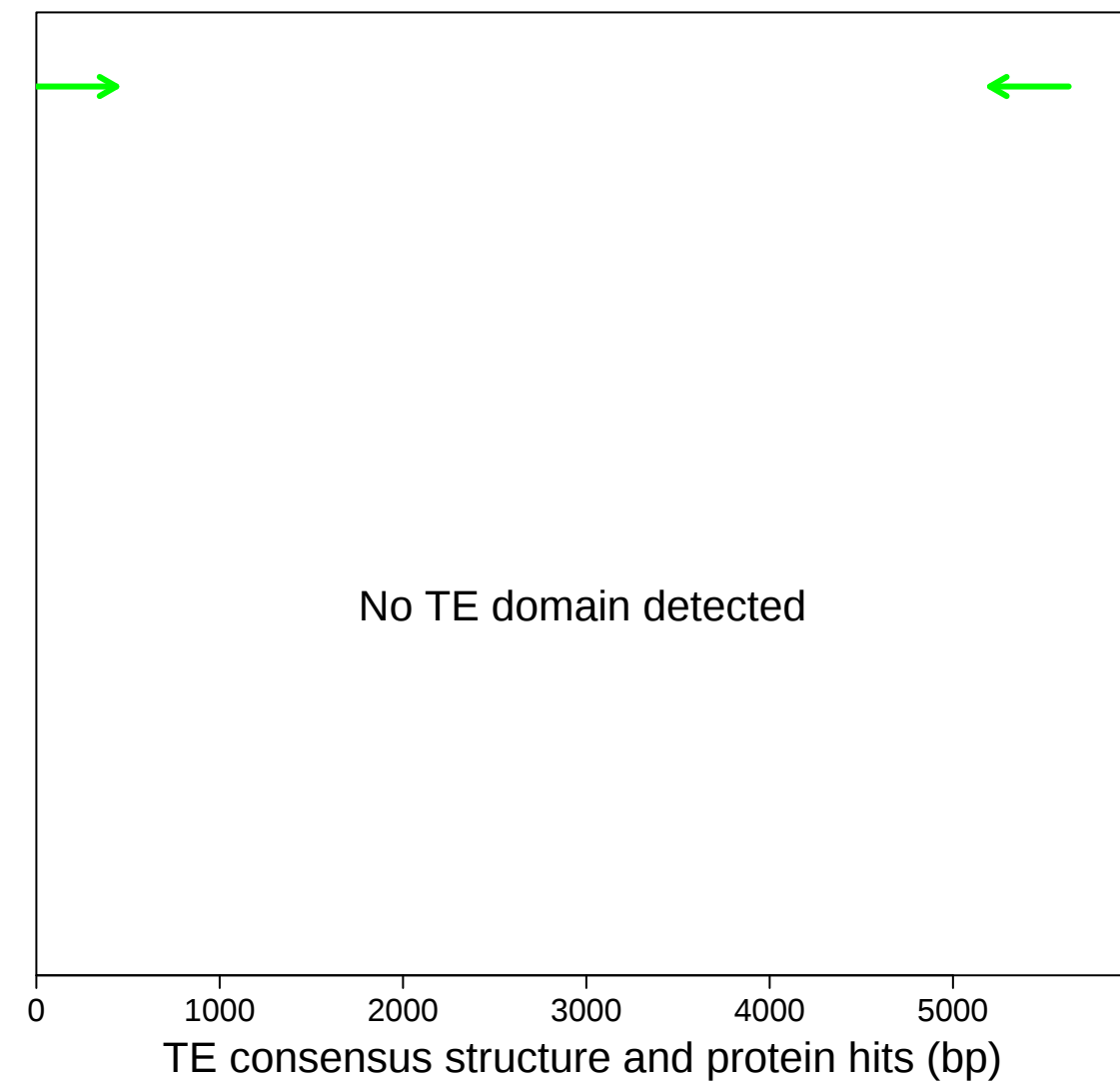
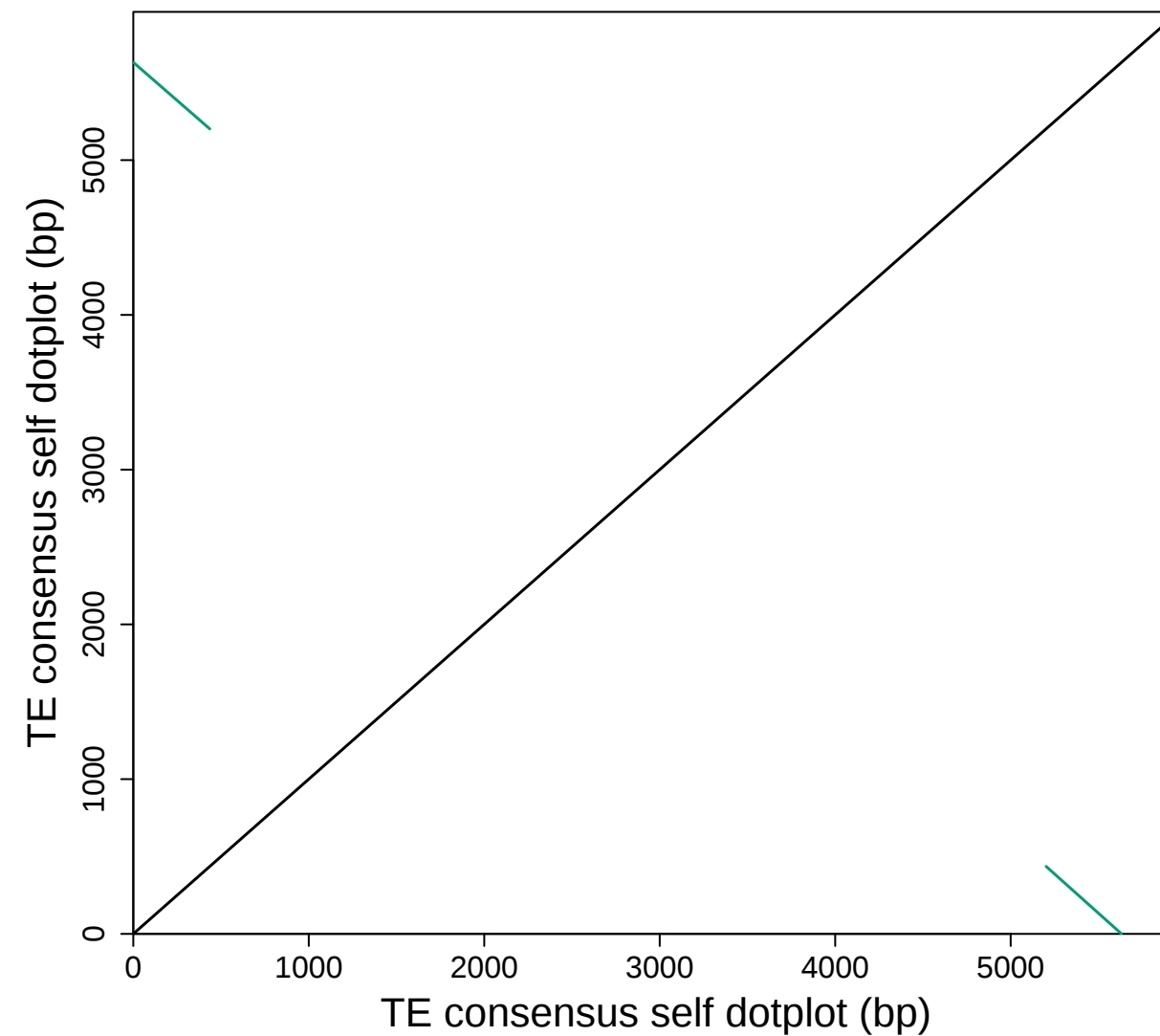
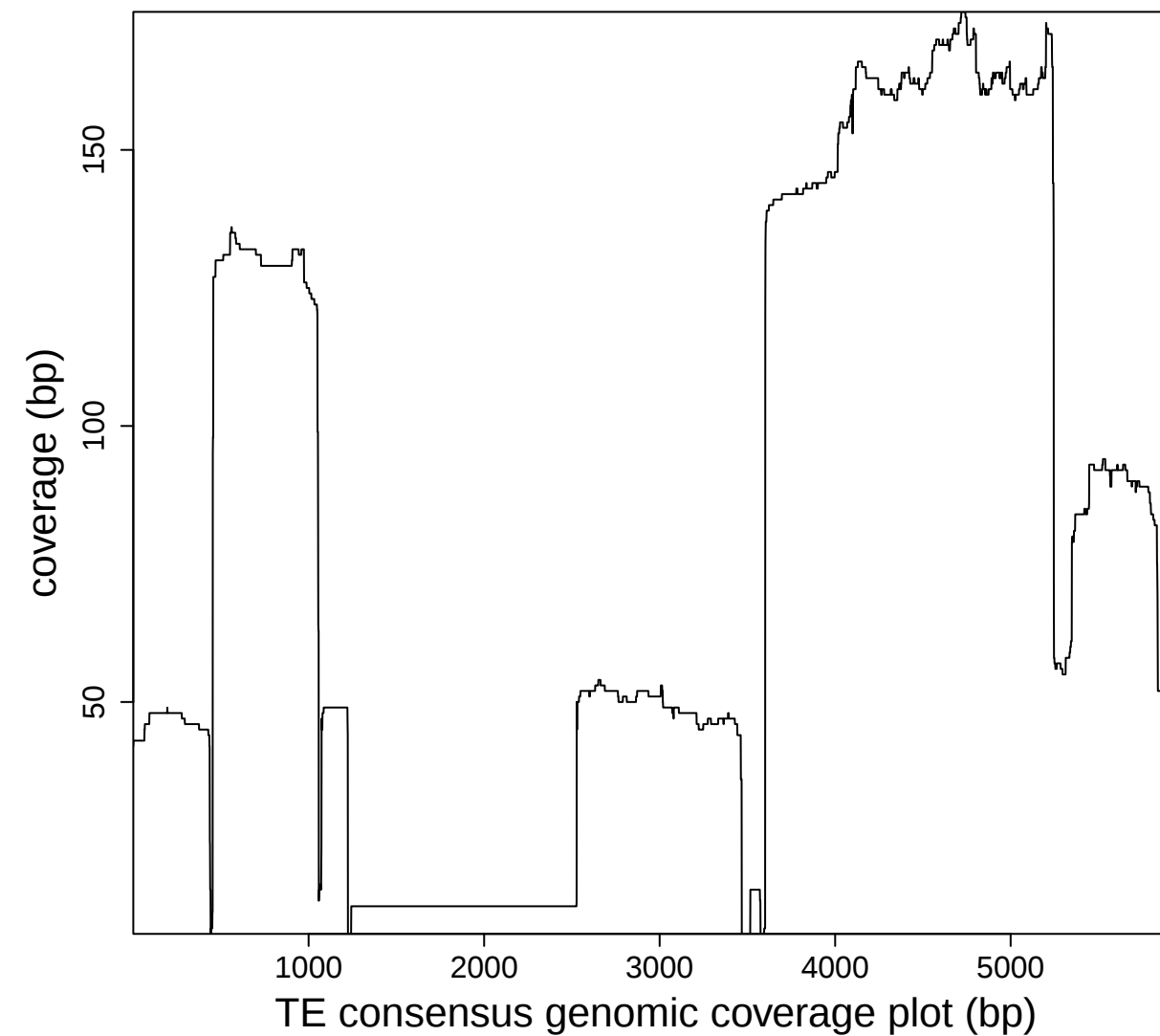
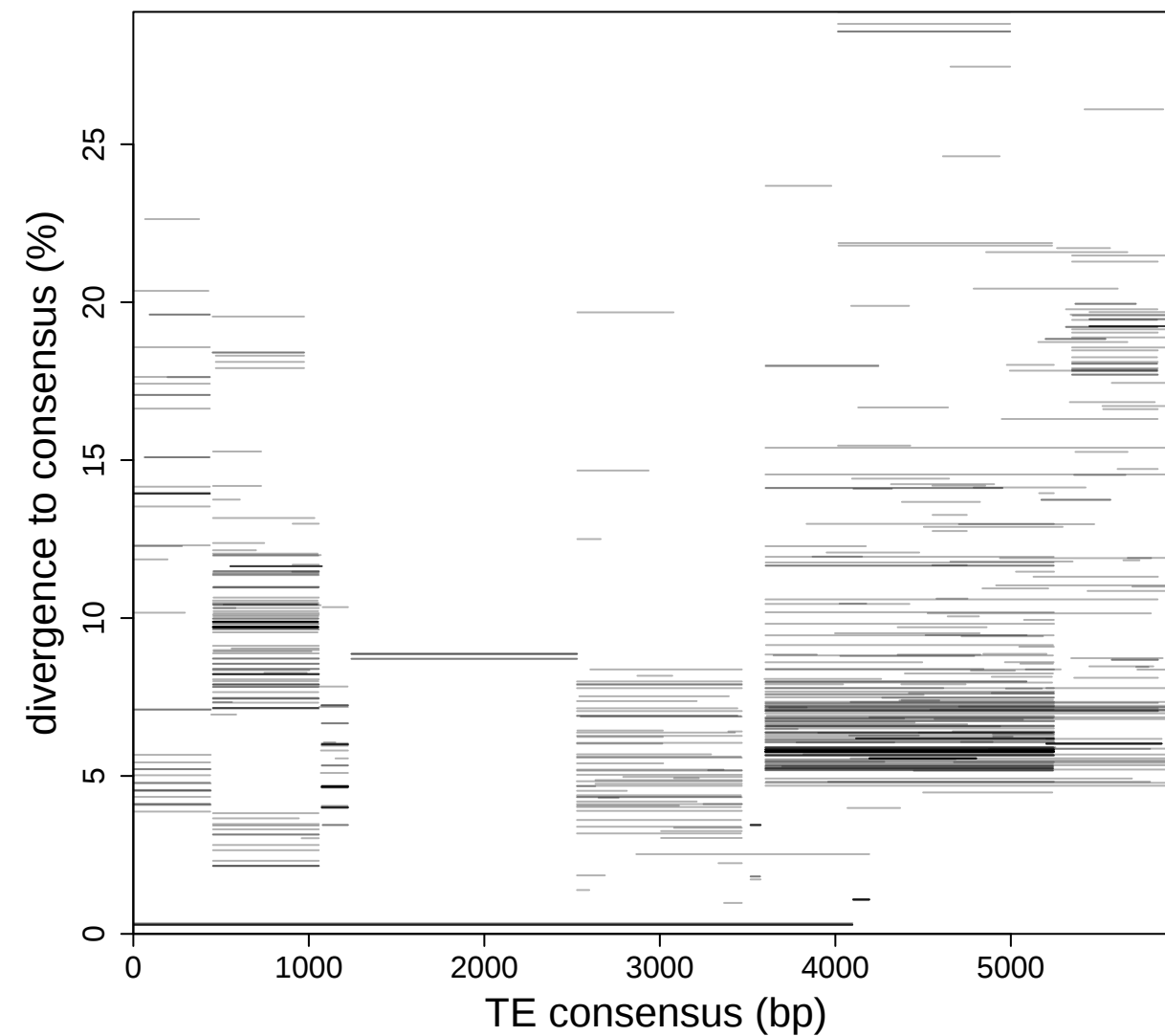
3.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 19027bp; fragments: 1875; full length: 0 (>=17124.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

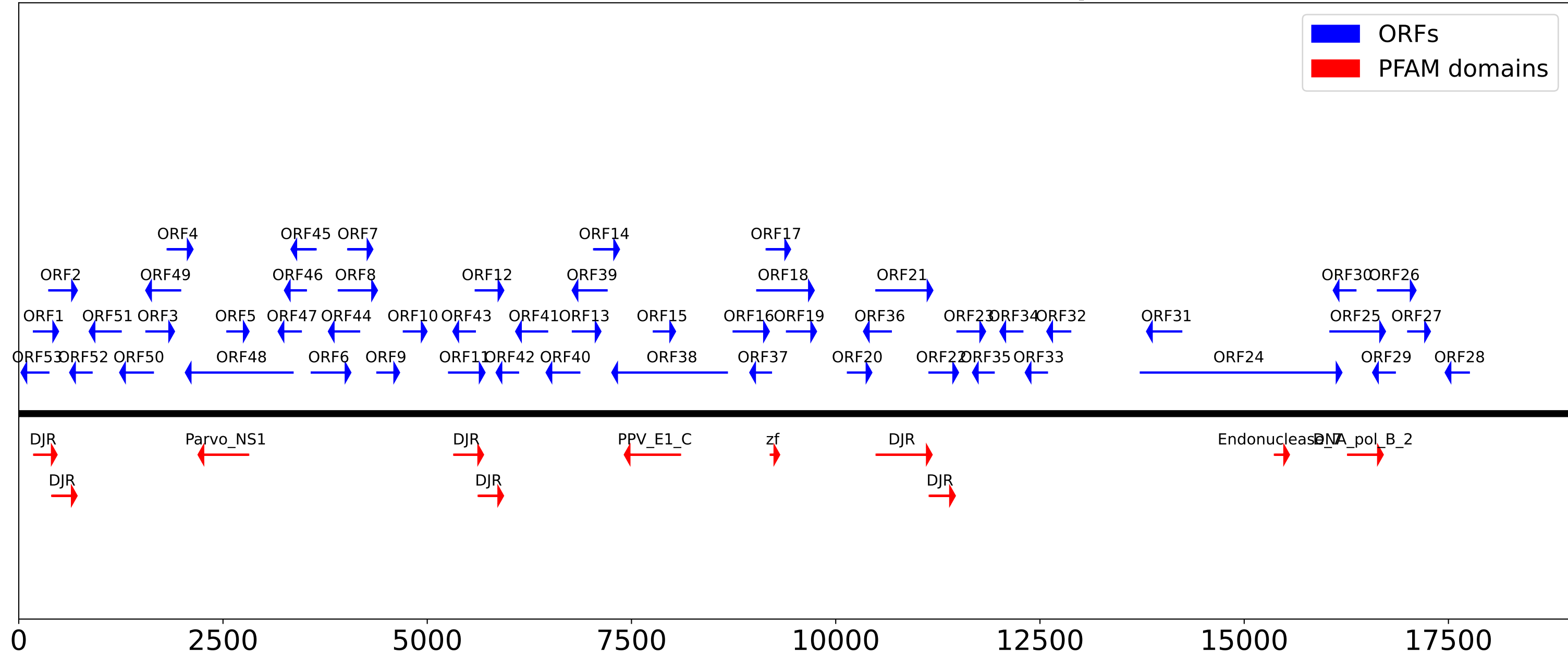


TE: rnd_5_family_3973
size: 5958bp; fragments: 598; full length: 0 (>=5362.2bp)

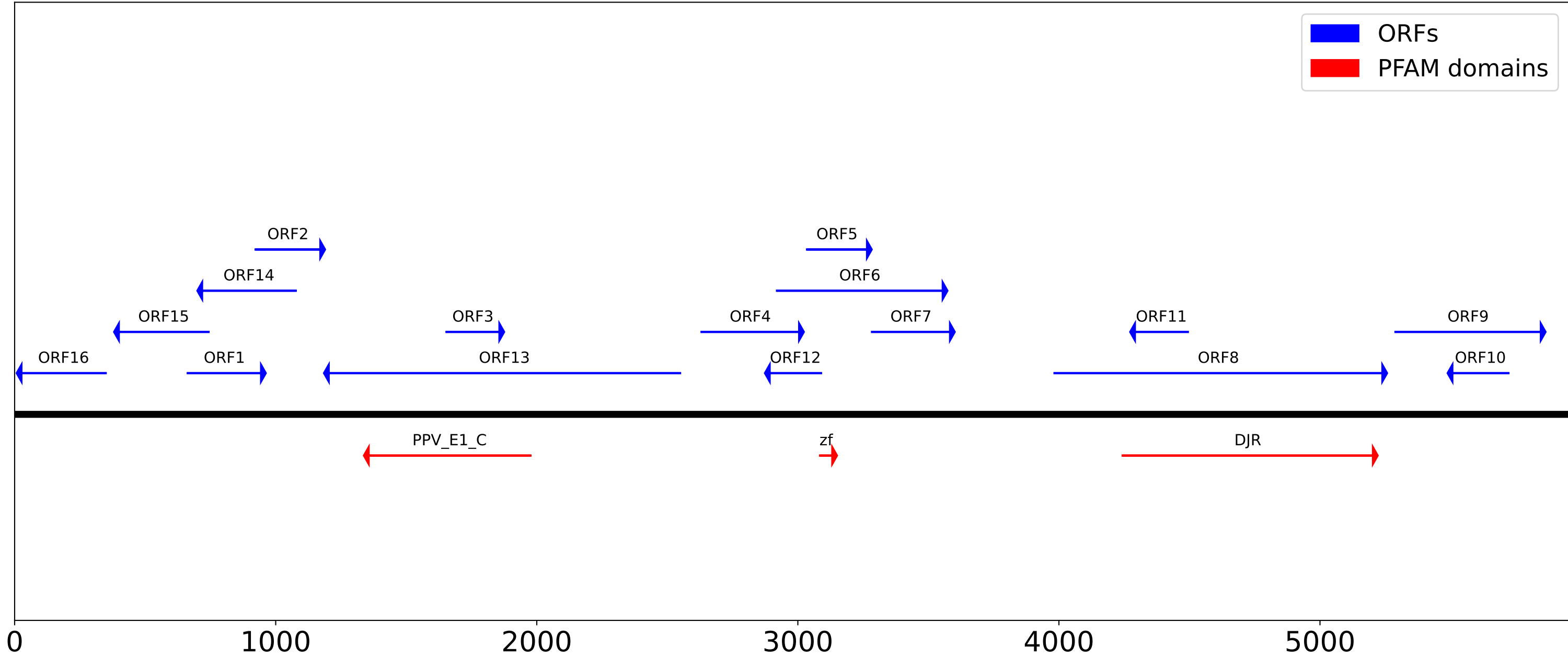
Before TEtrimmer 5958 bp



After TETrimmer ORF and PFAM domain plot



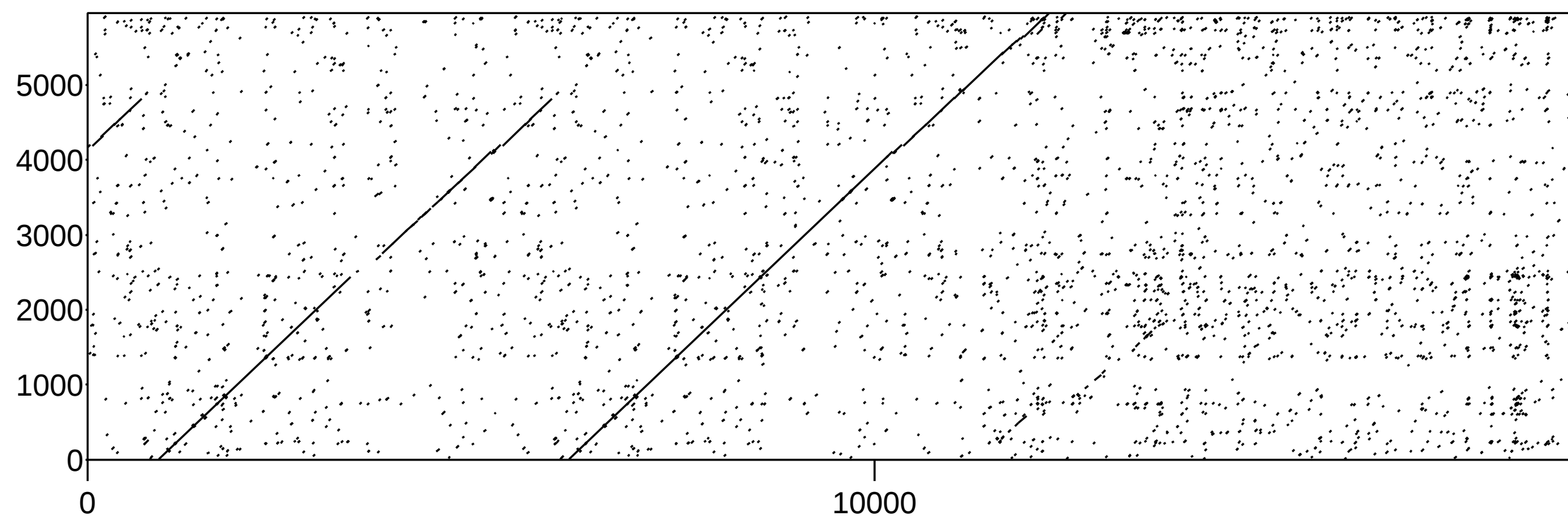
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)