

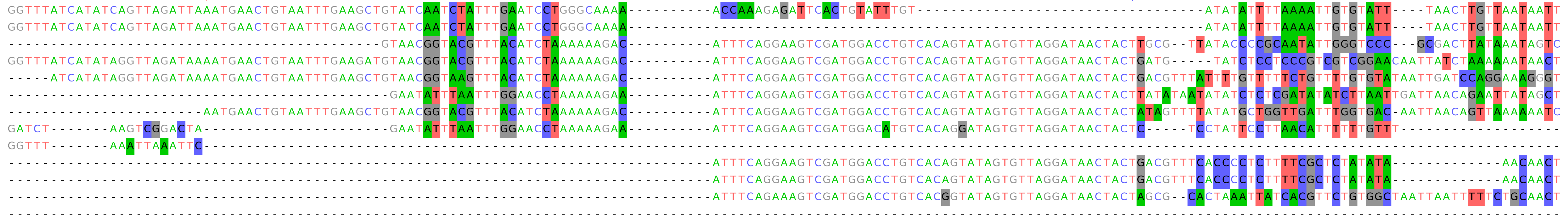
Start crop Point

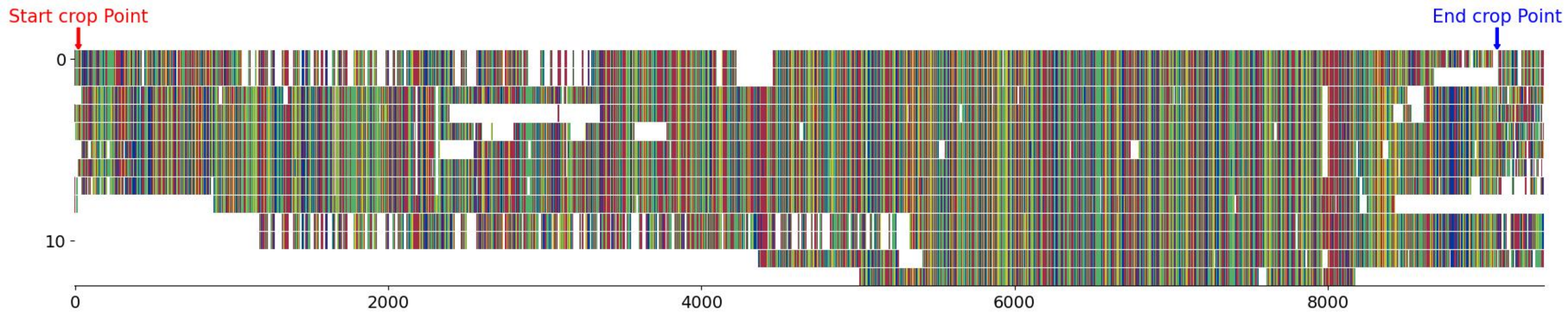


End crop Point



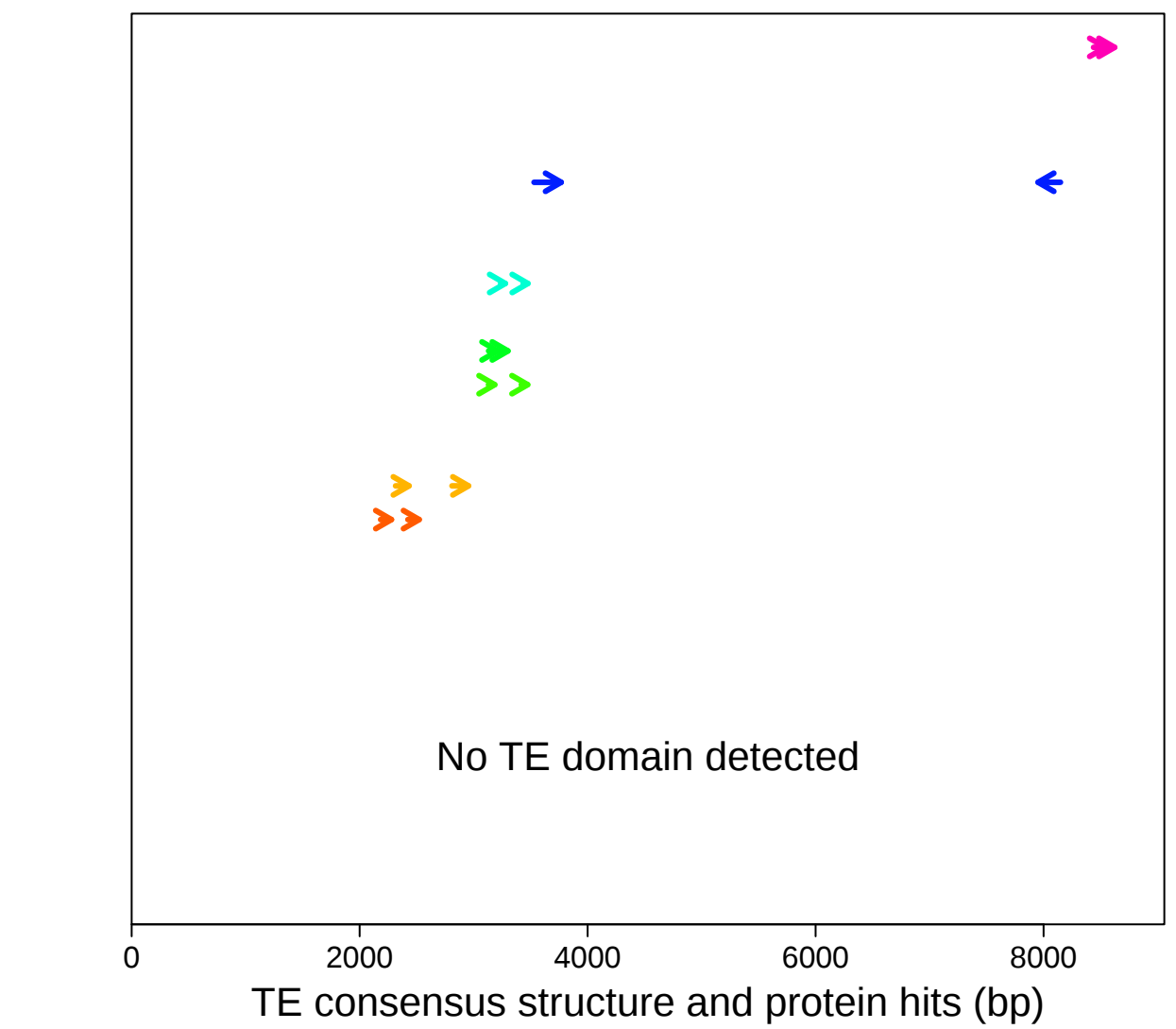
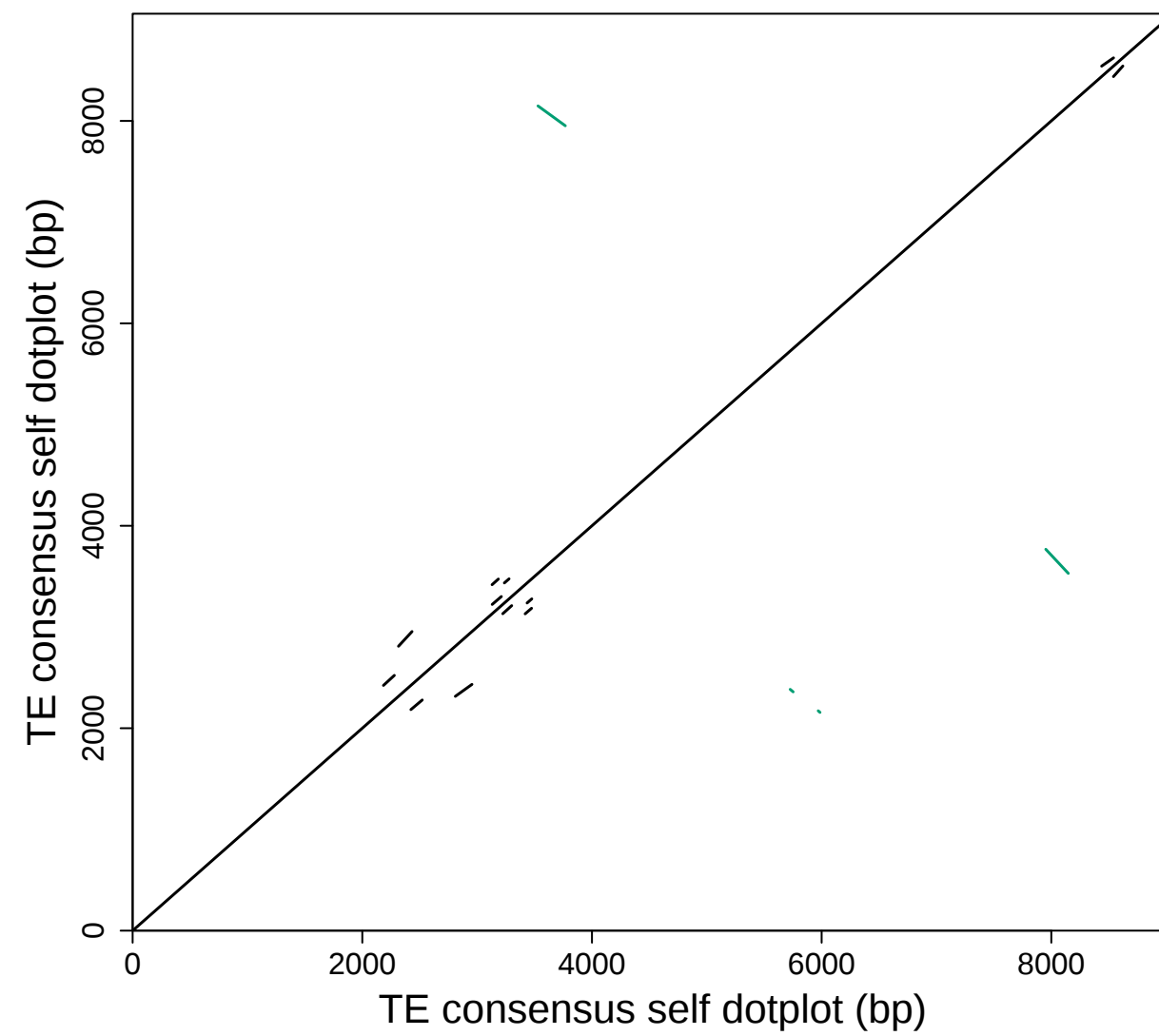
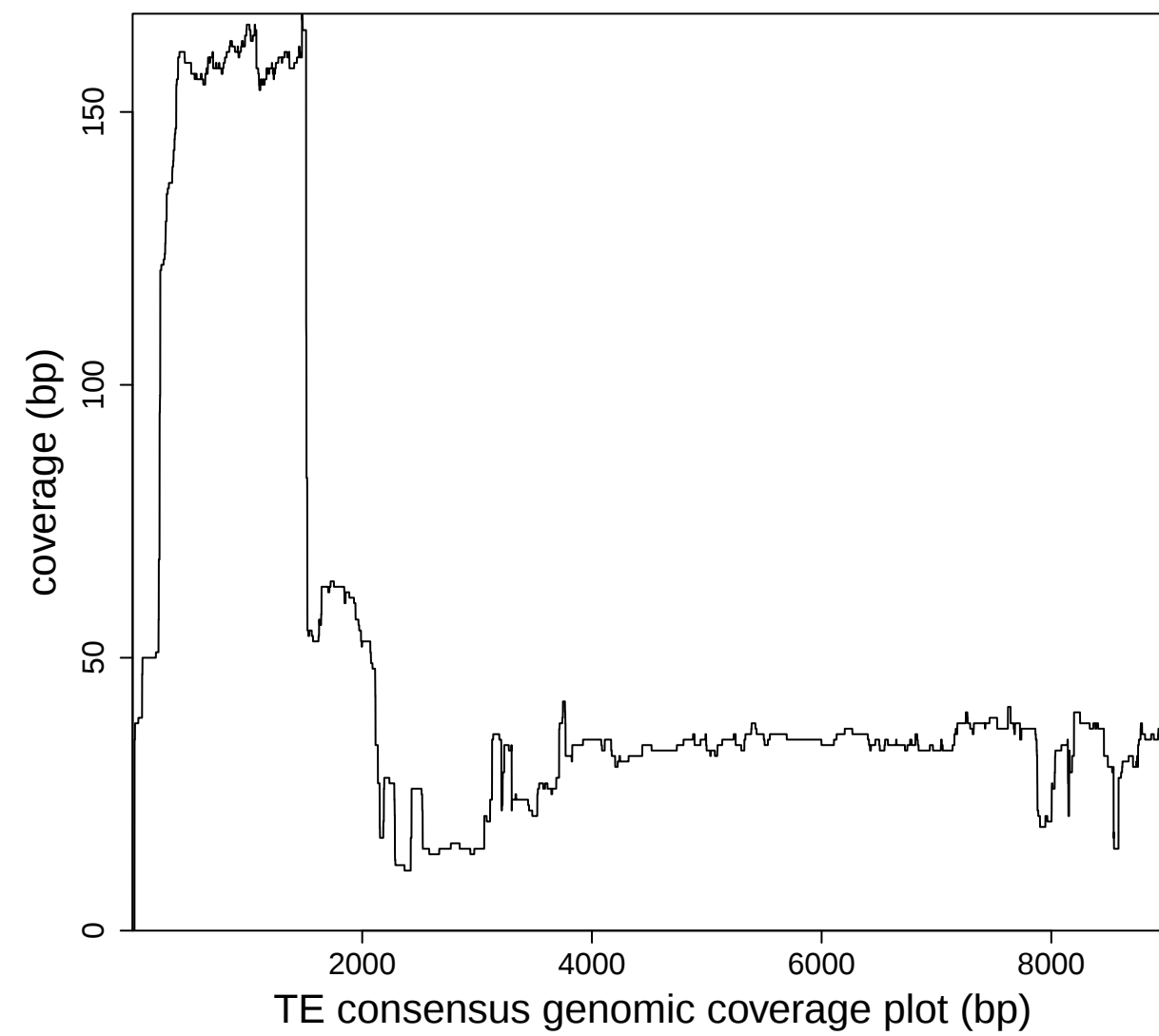
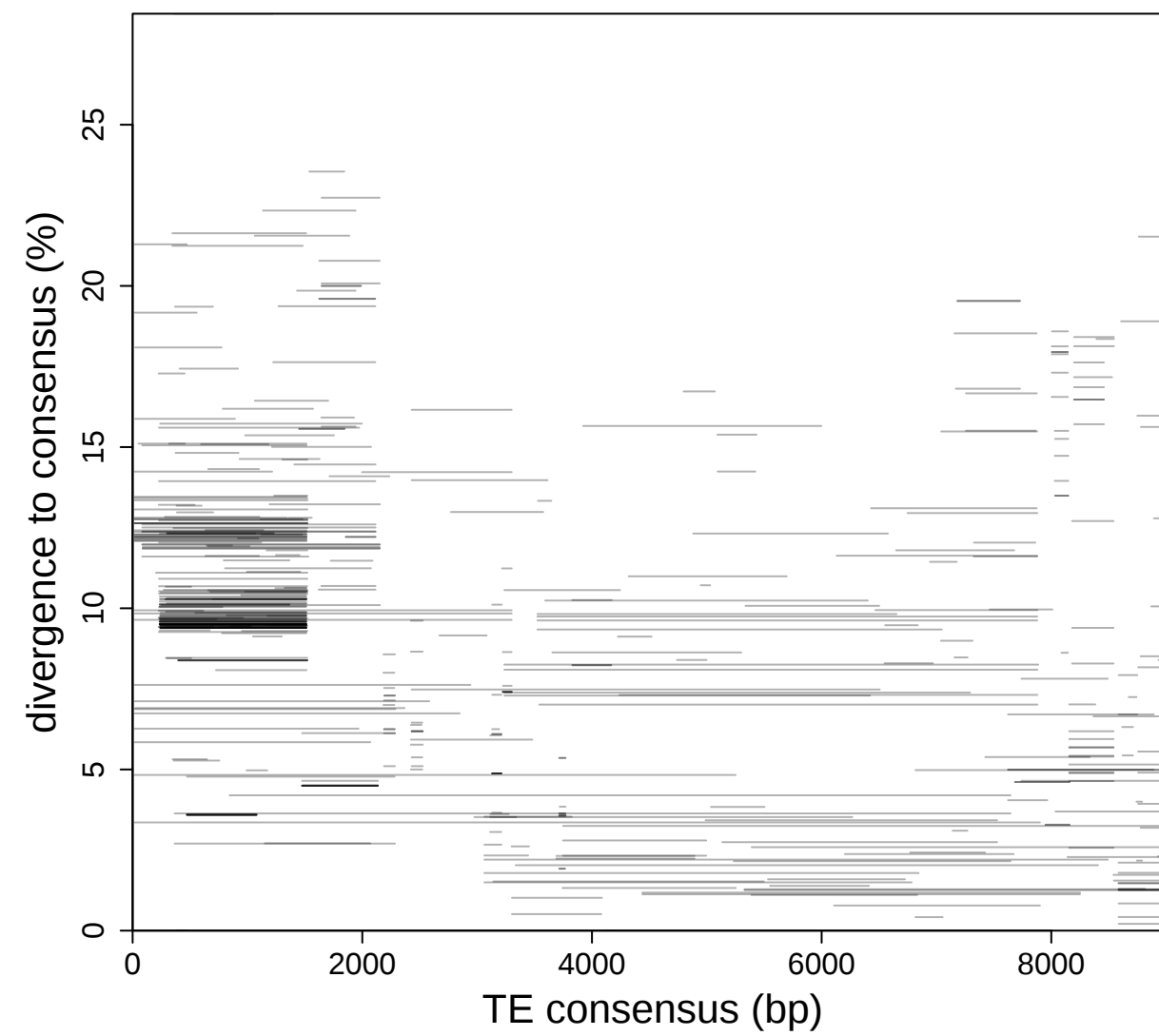
MSA length = 9059





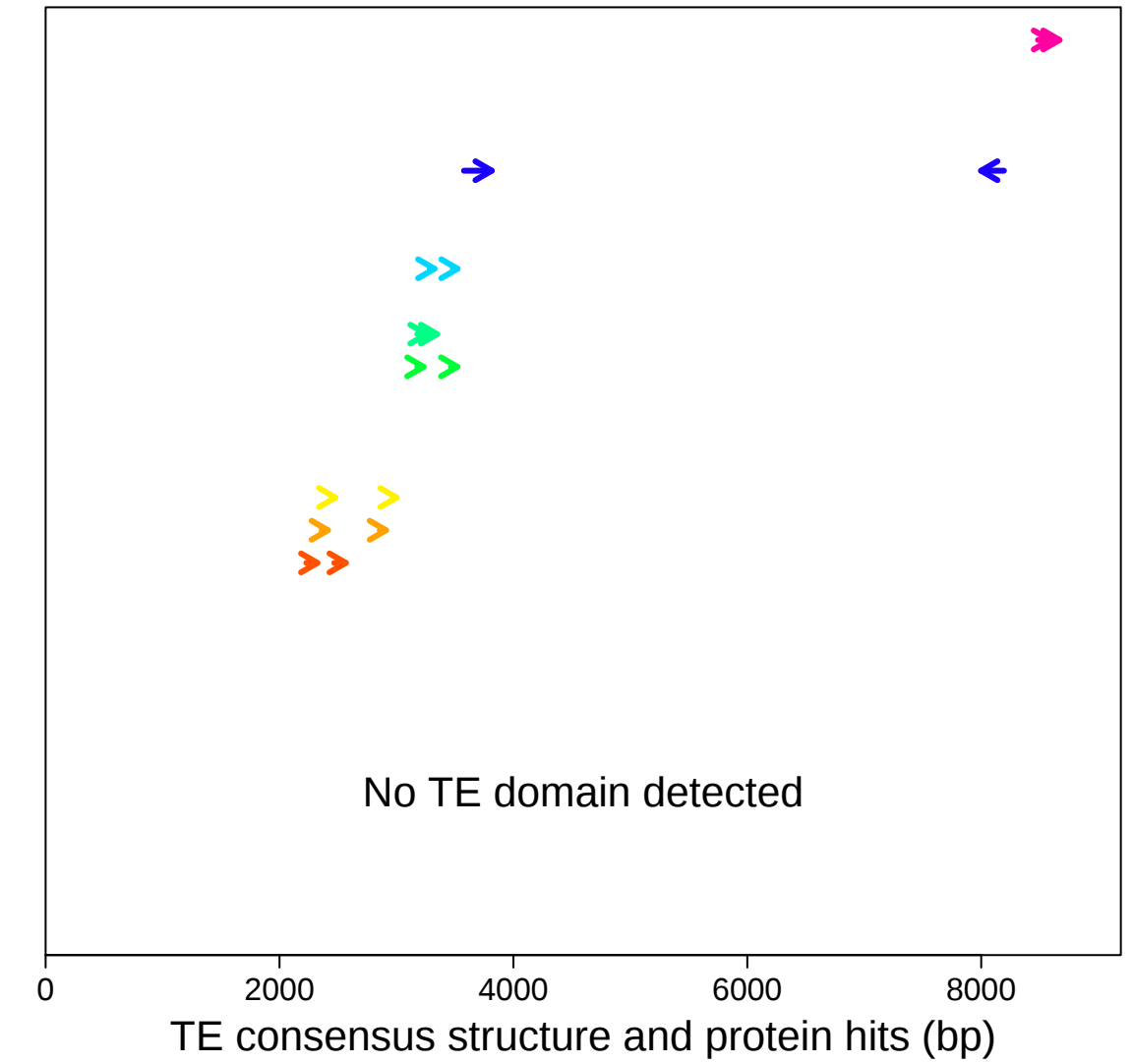
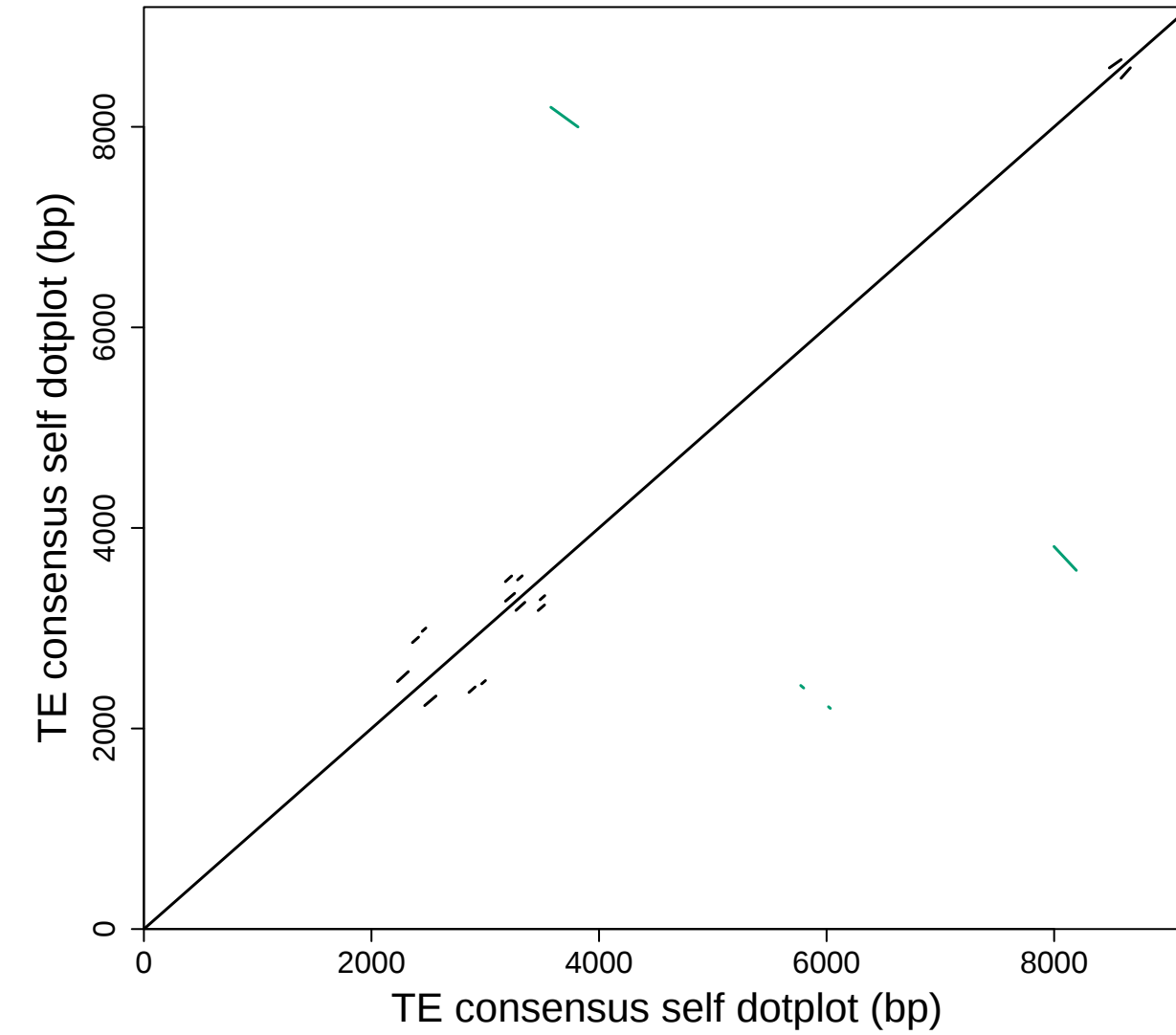
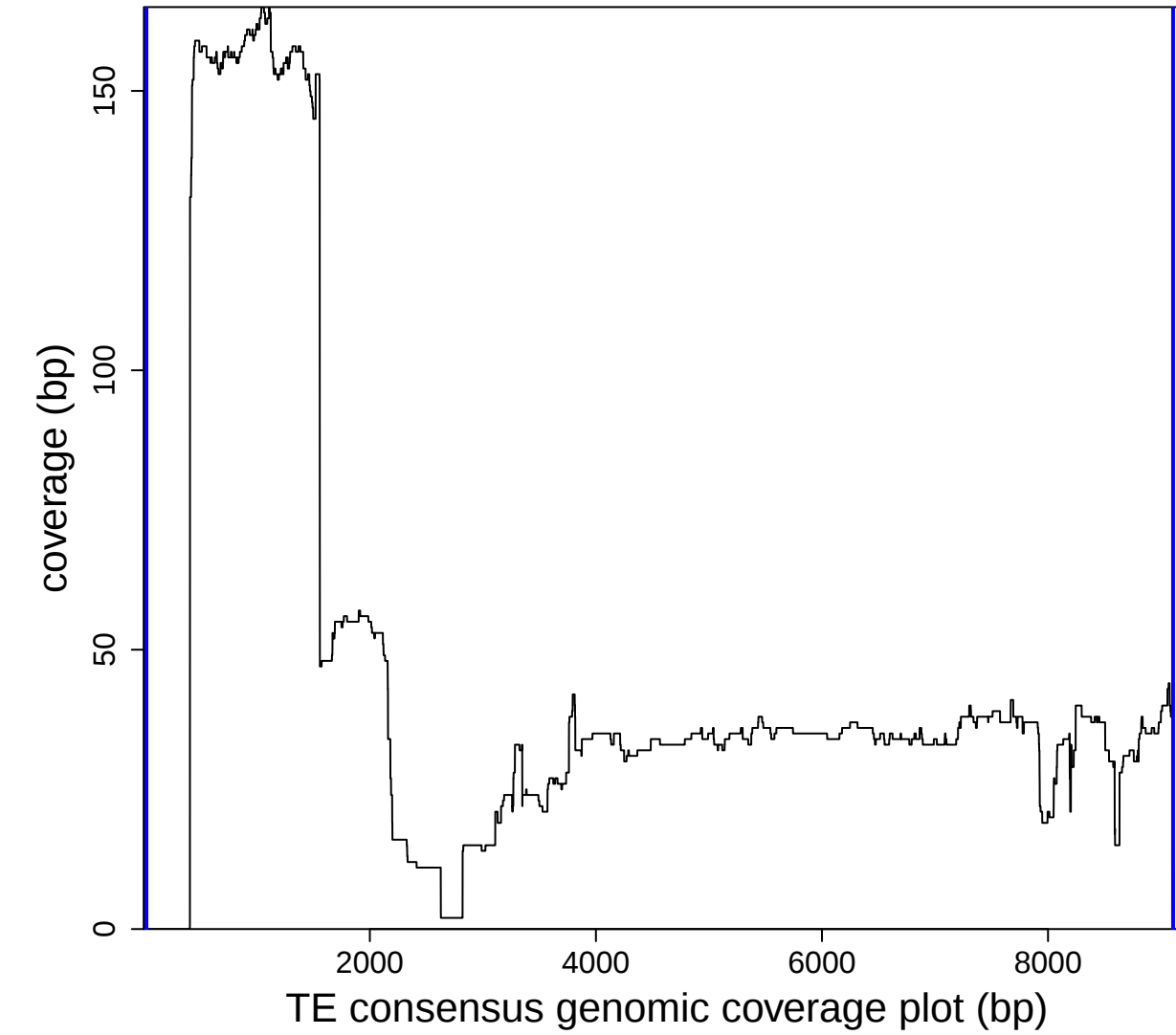
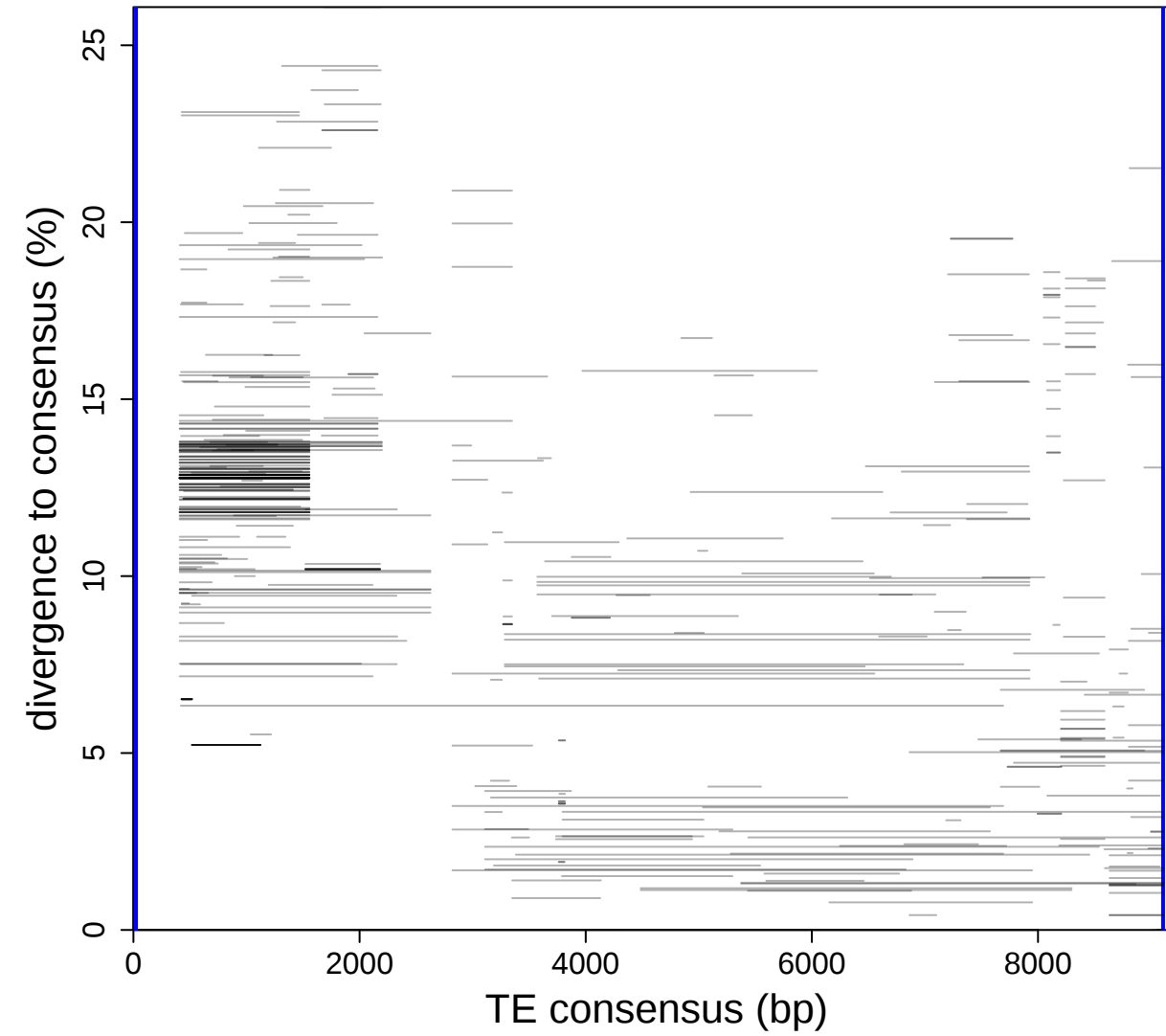
a.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 9059bp; fragments: 483; full length: 0 (>=8153.1bp)

After TEtrimmer 9059 bp

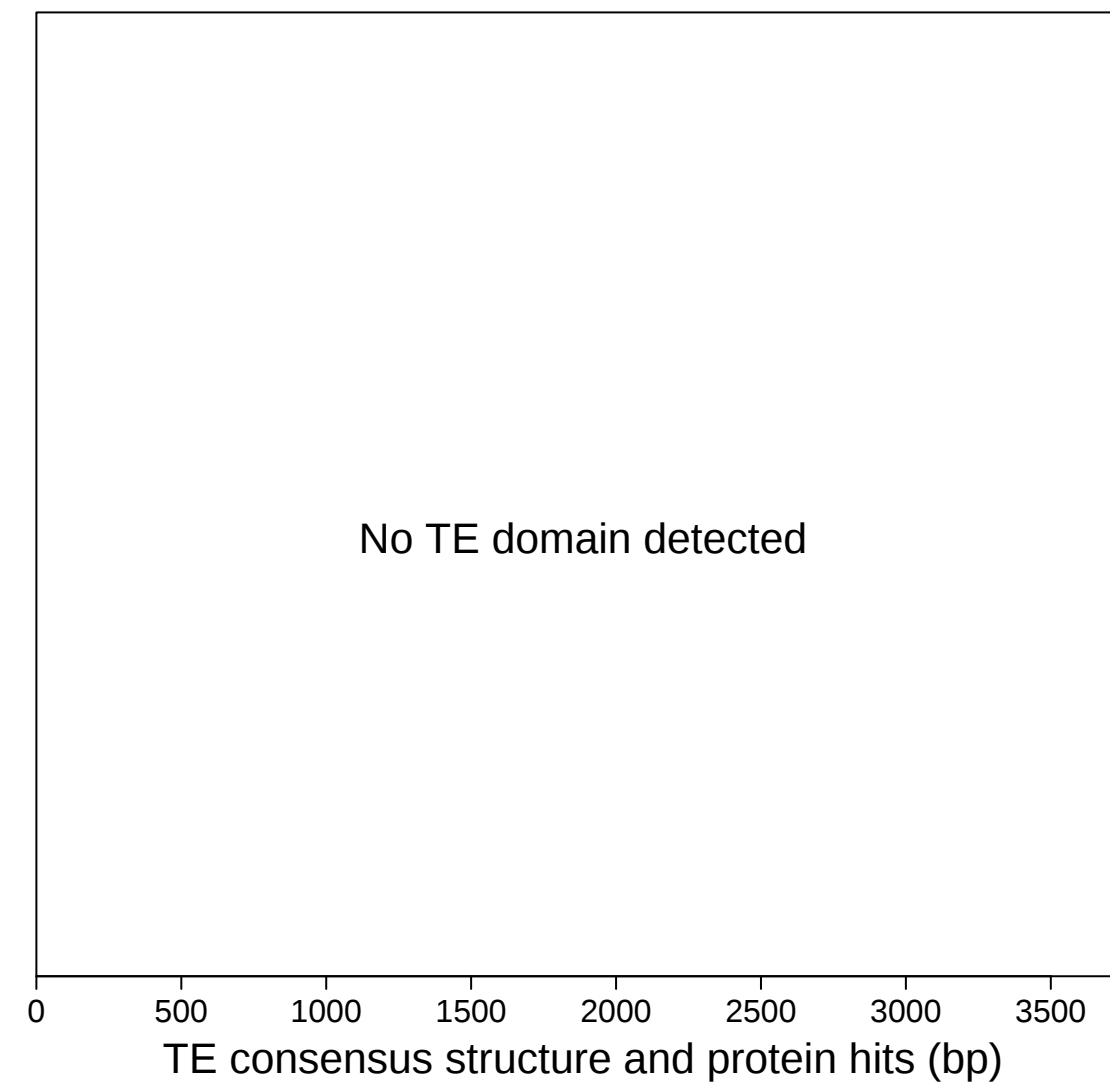
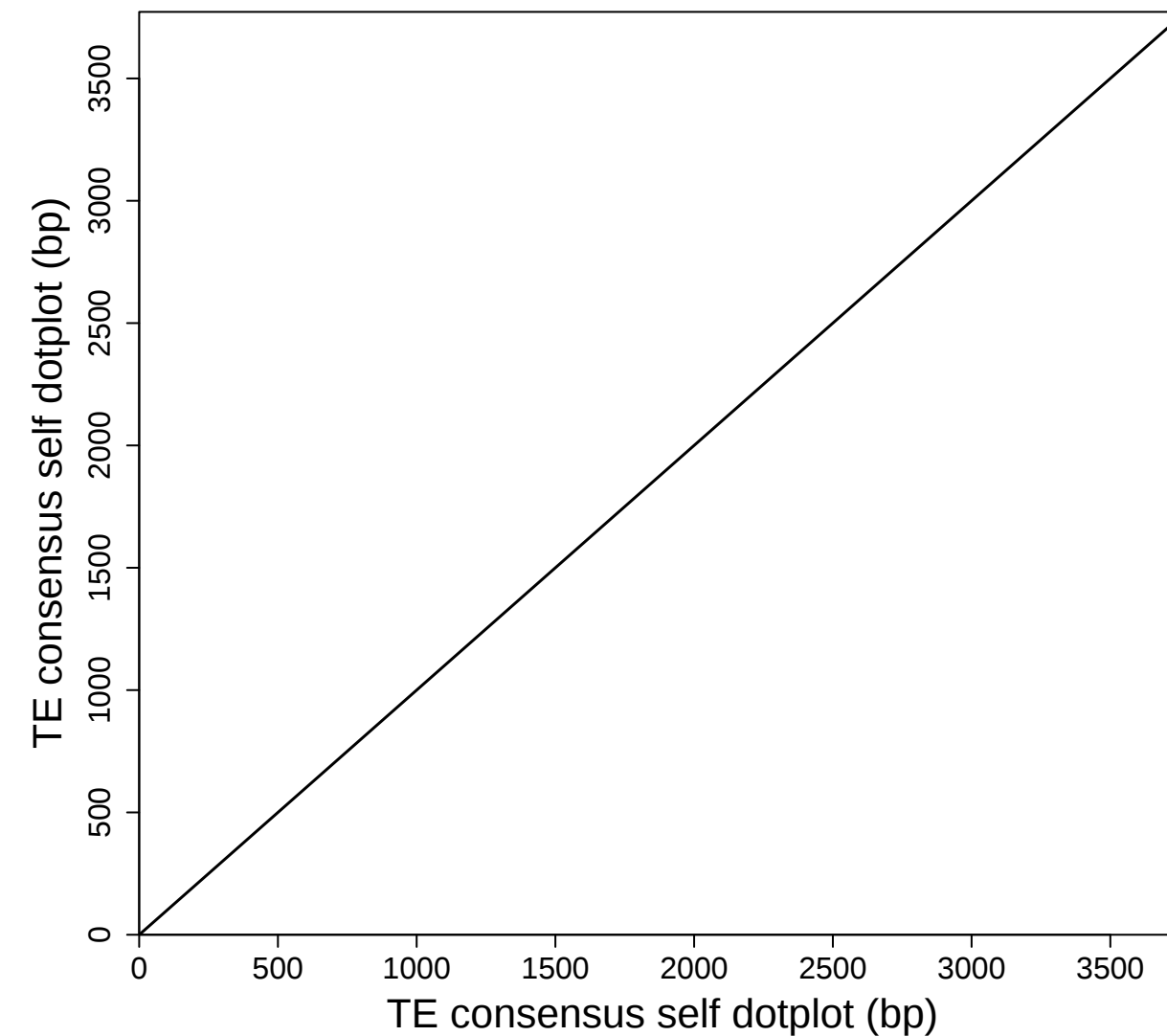
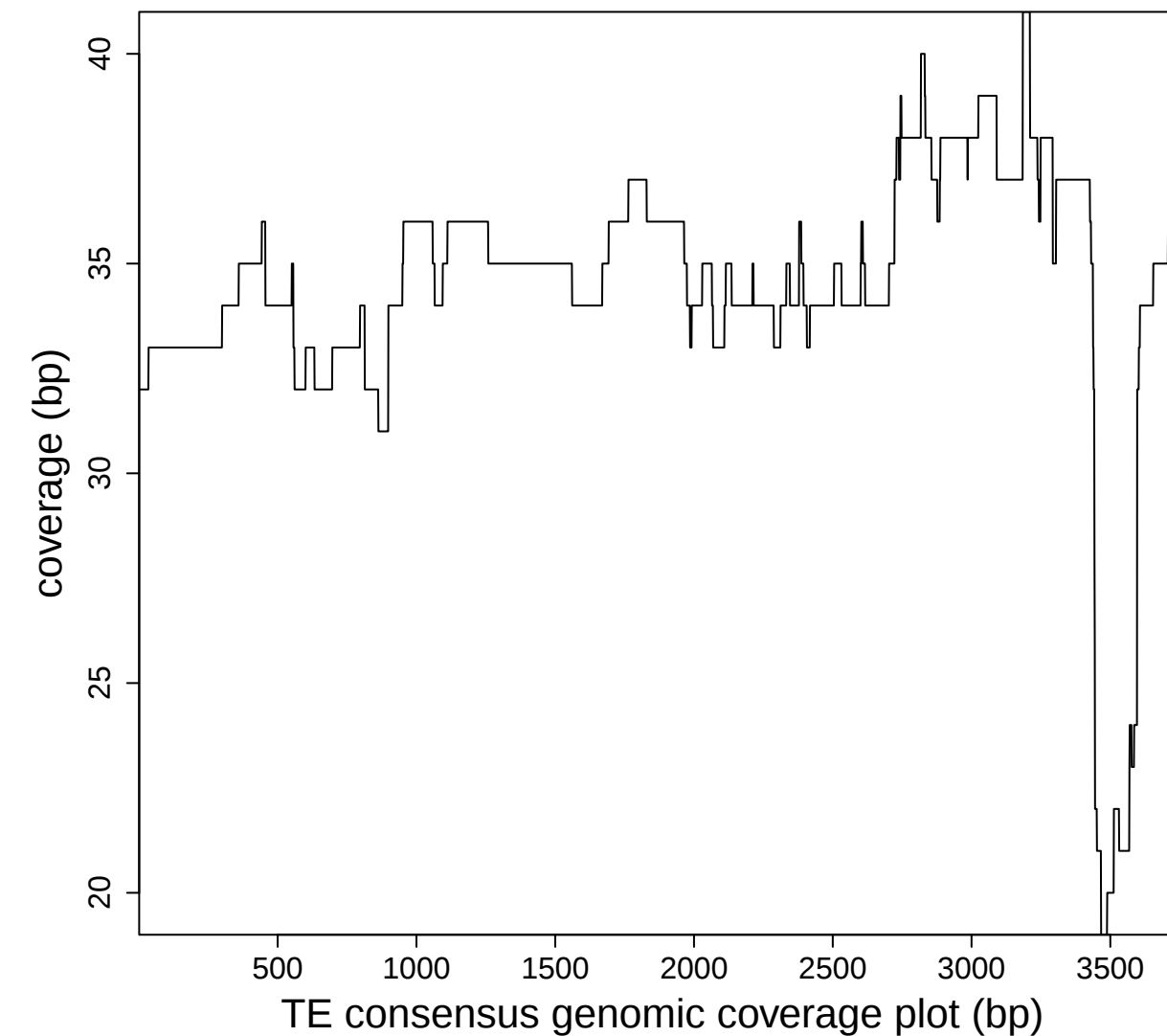
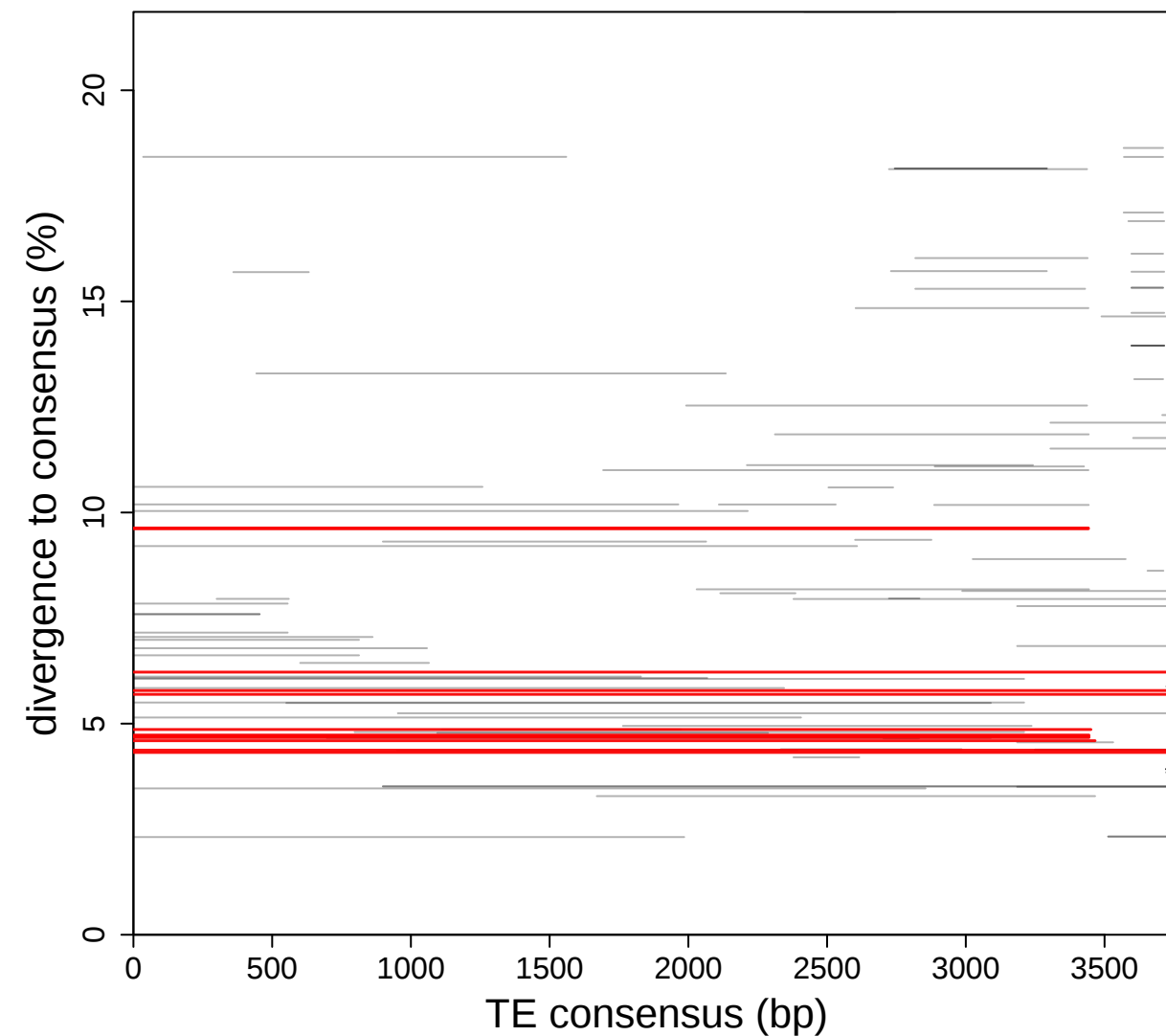


5.fasta.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 9256bp; fragments: 446; full length: 0 (>=8330.4bp)

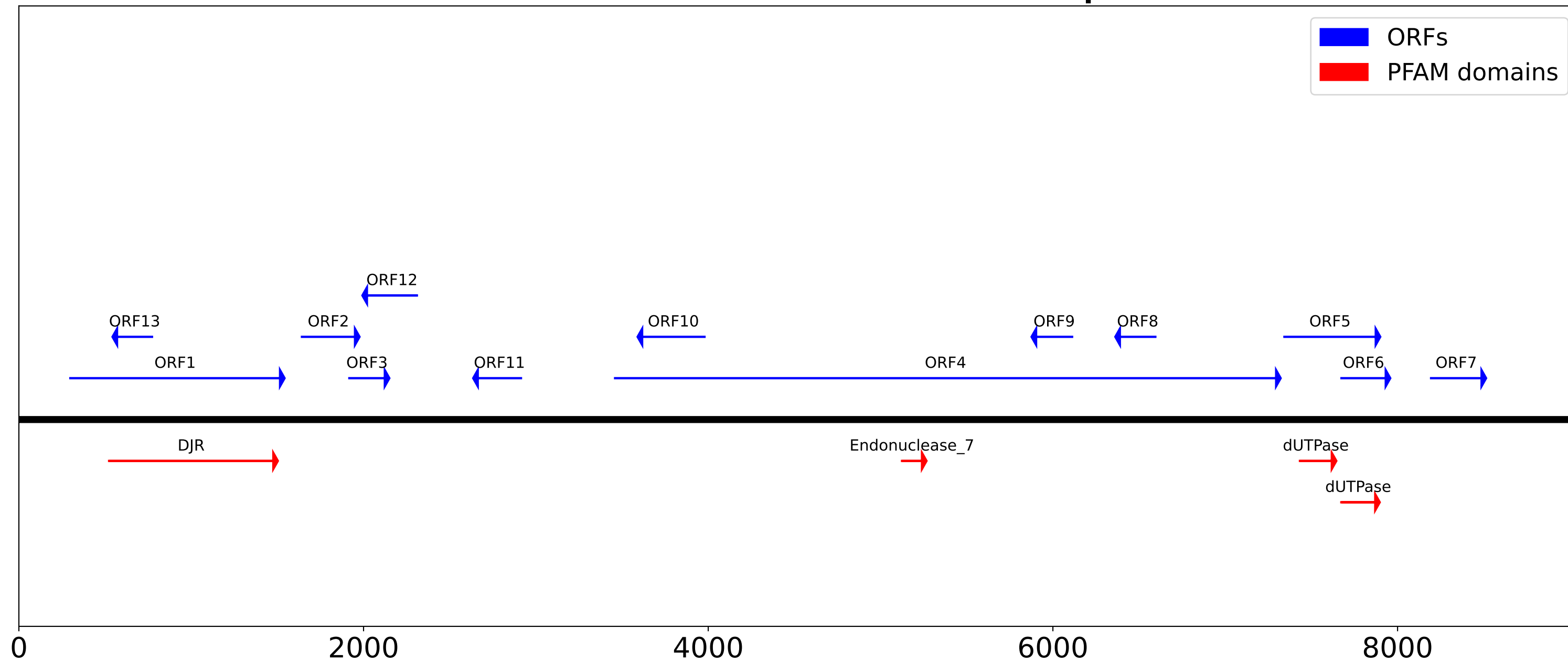
After TEtrimmer Extended plot Blue lines are boundaries



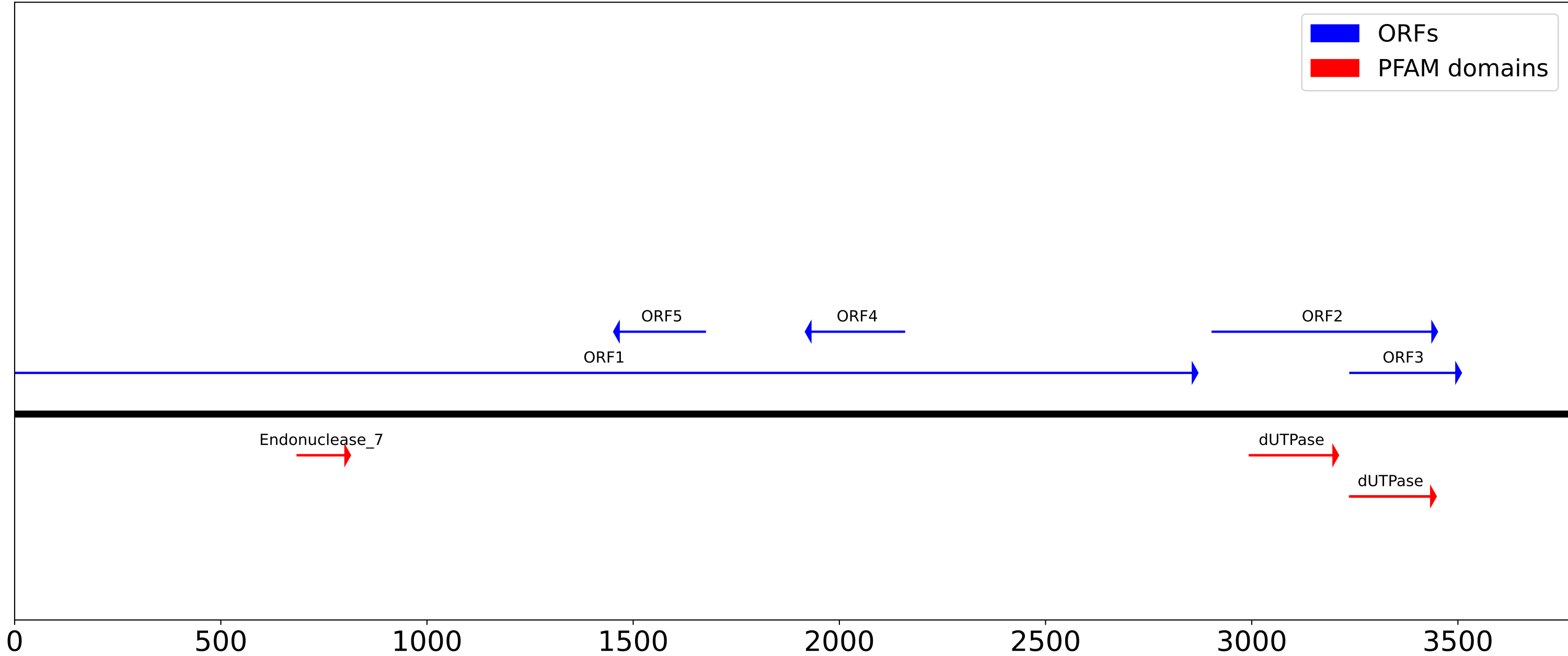
TE: rnd_5_family_7365
size: 3772bp; fragments: 110; full length: 12 (≥ 3394.8 bp)



After TEtrimmer ORF and PFAM domain plot



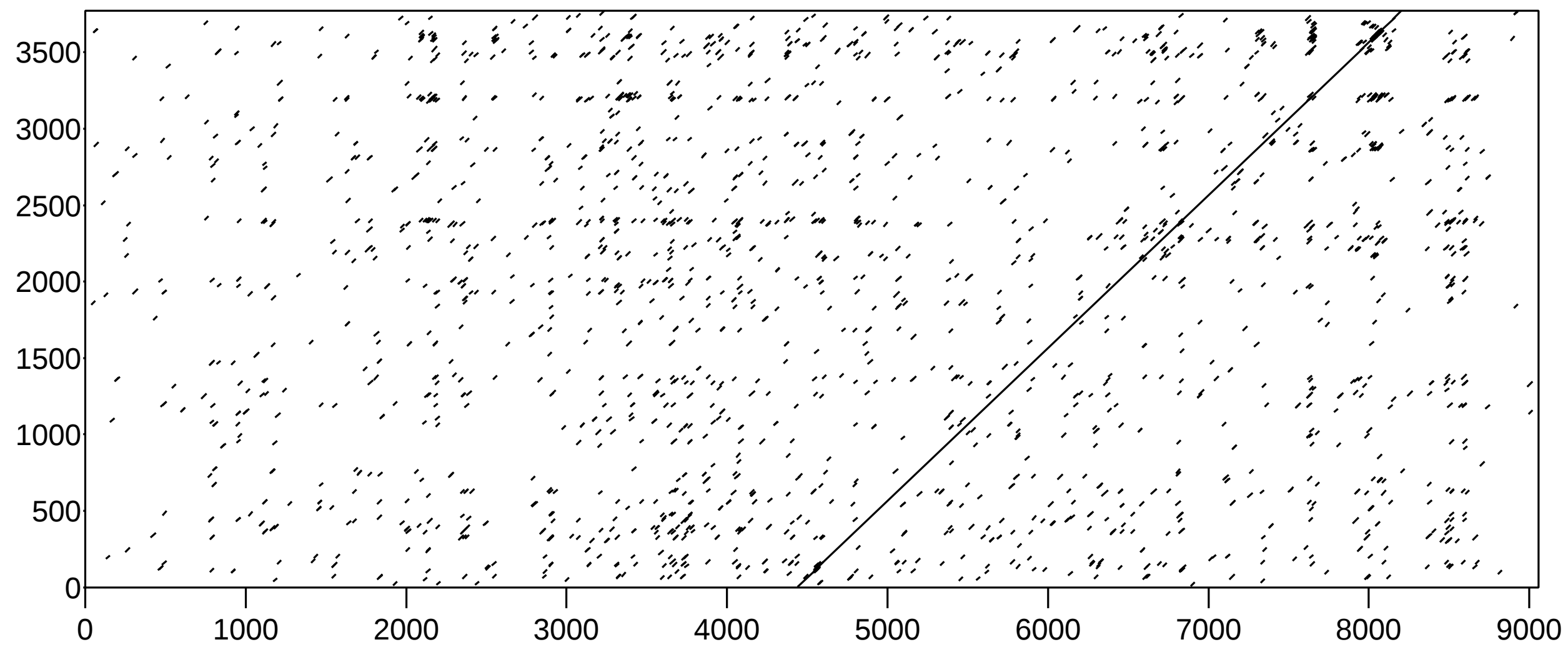
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)