

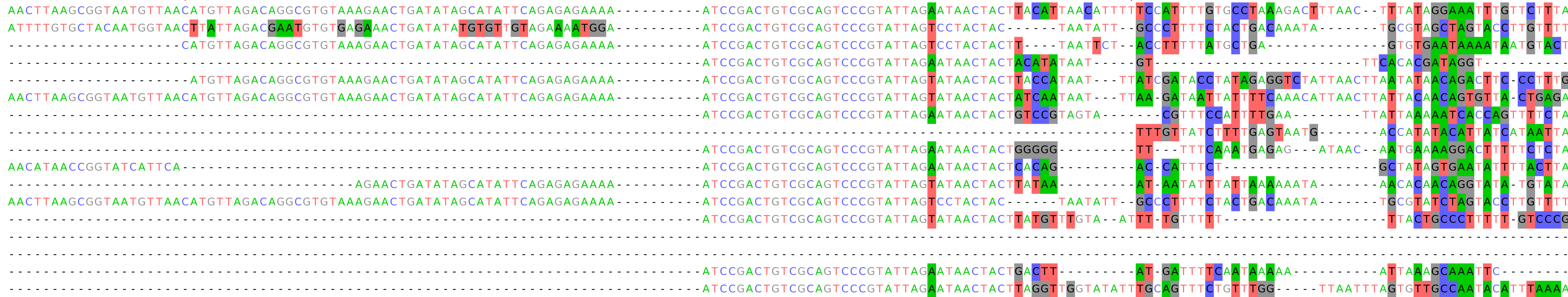
Start crop Point

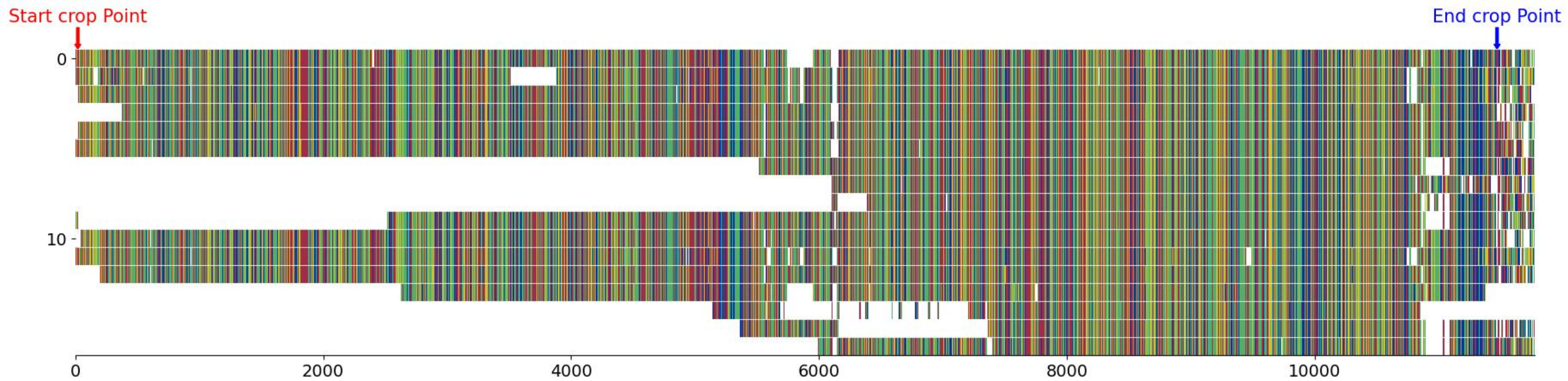


End crop Point



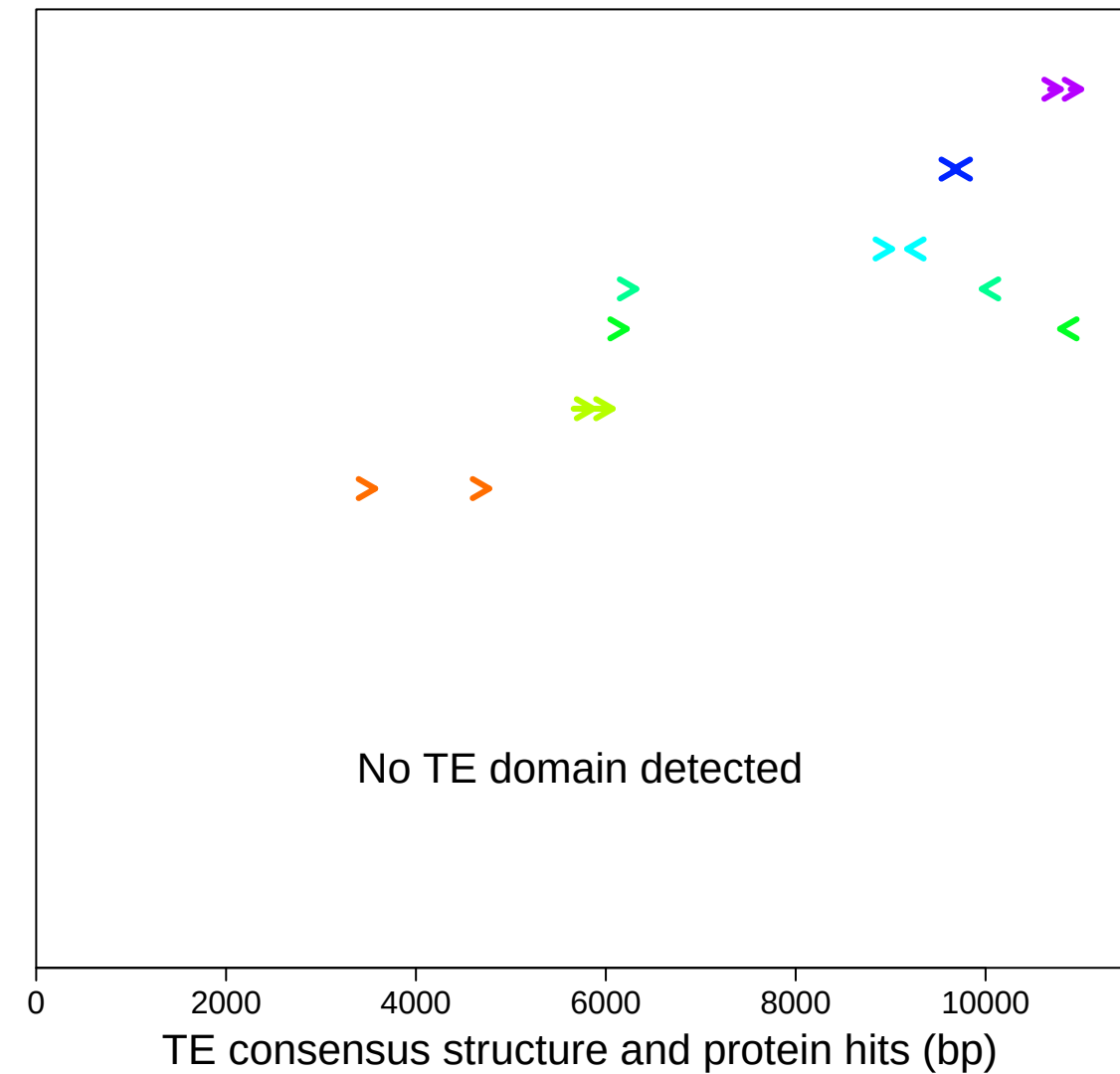
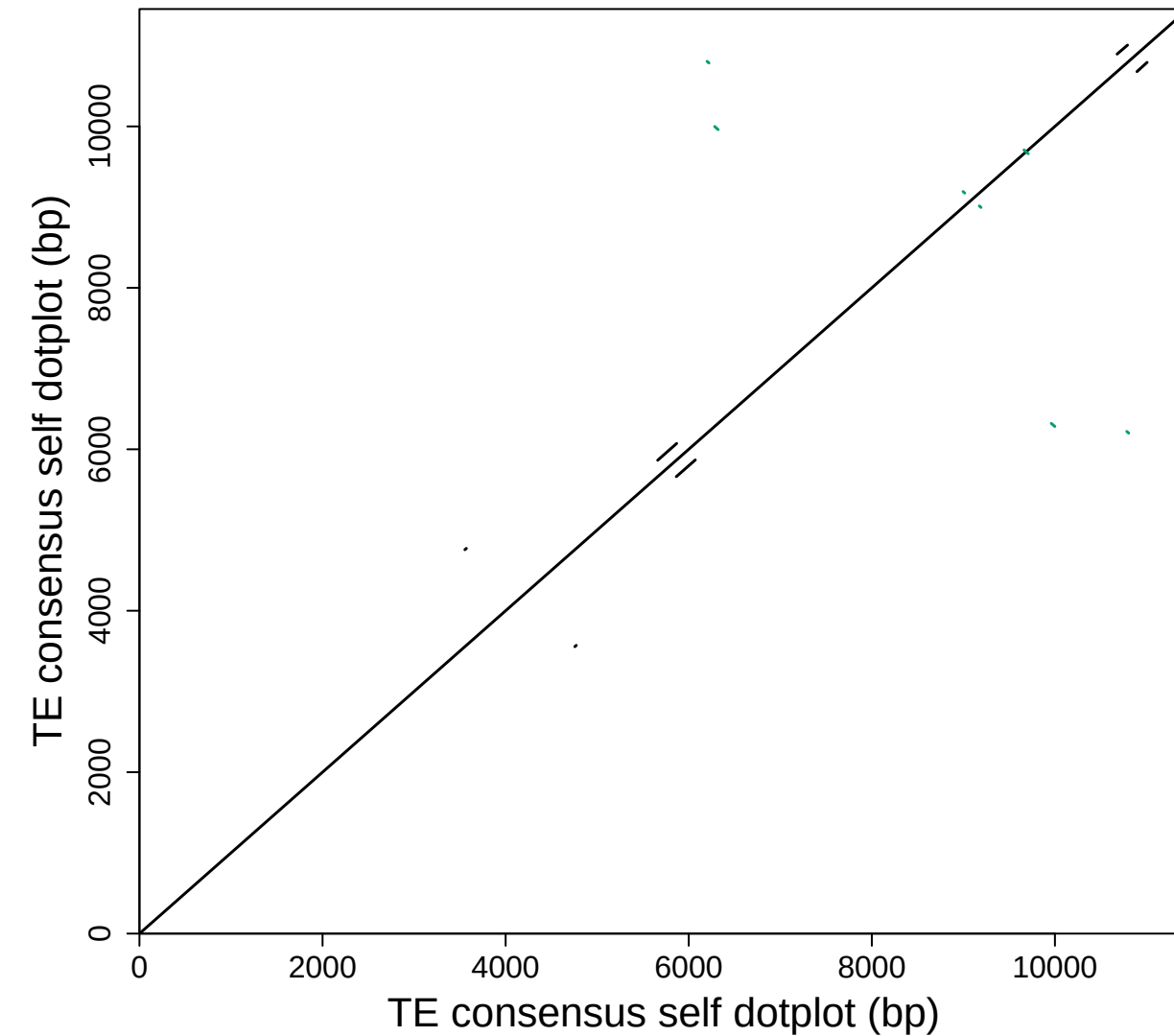
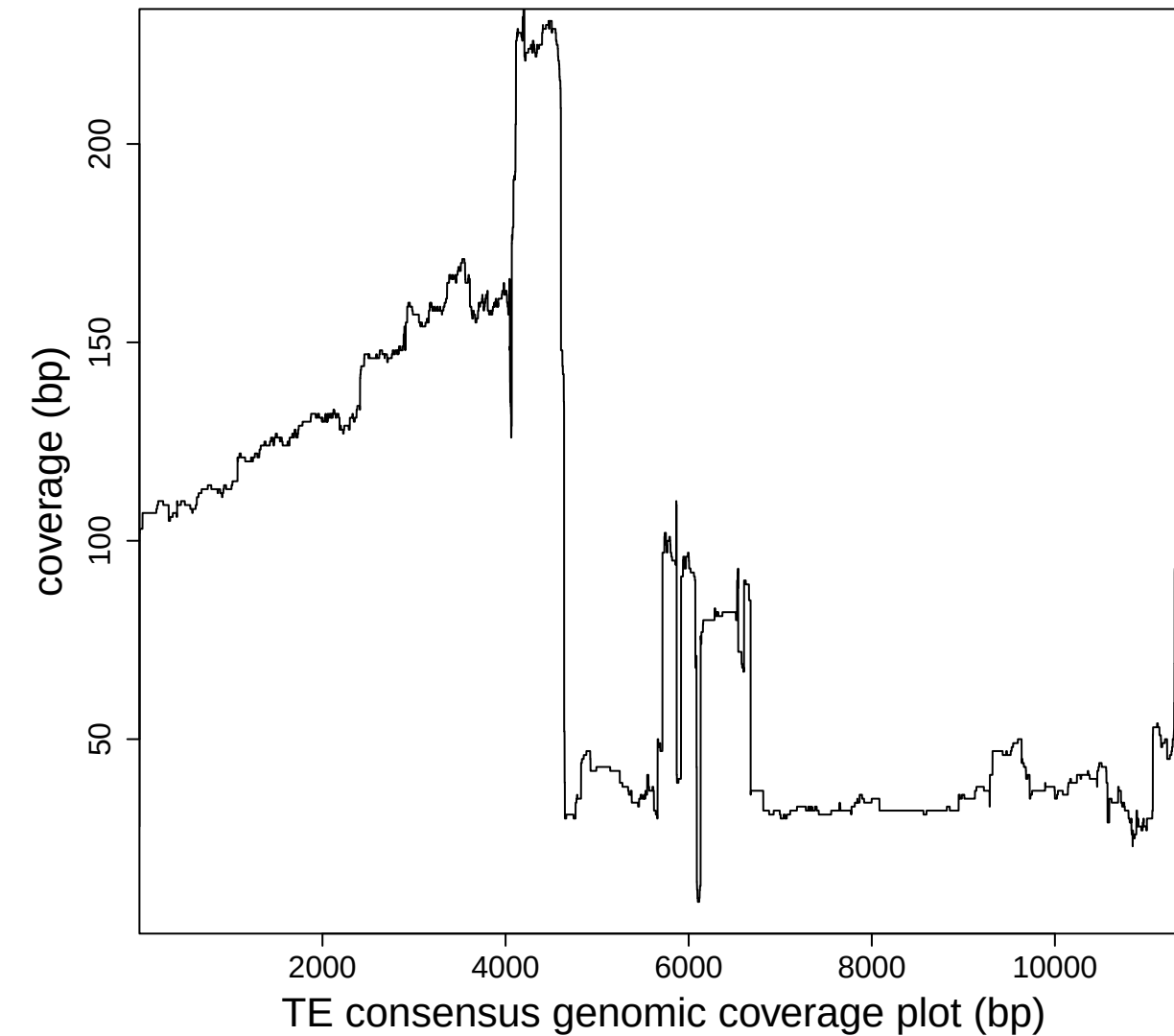
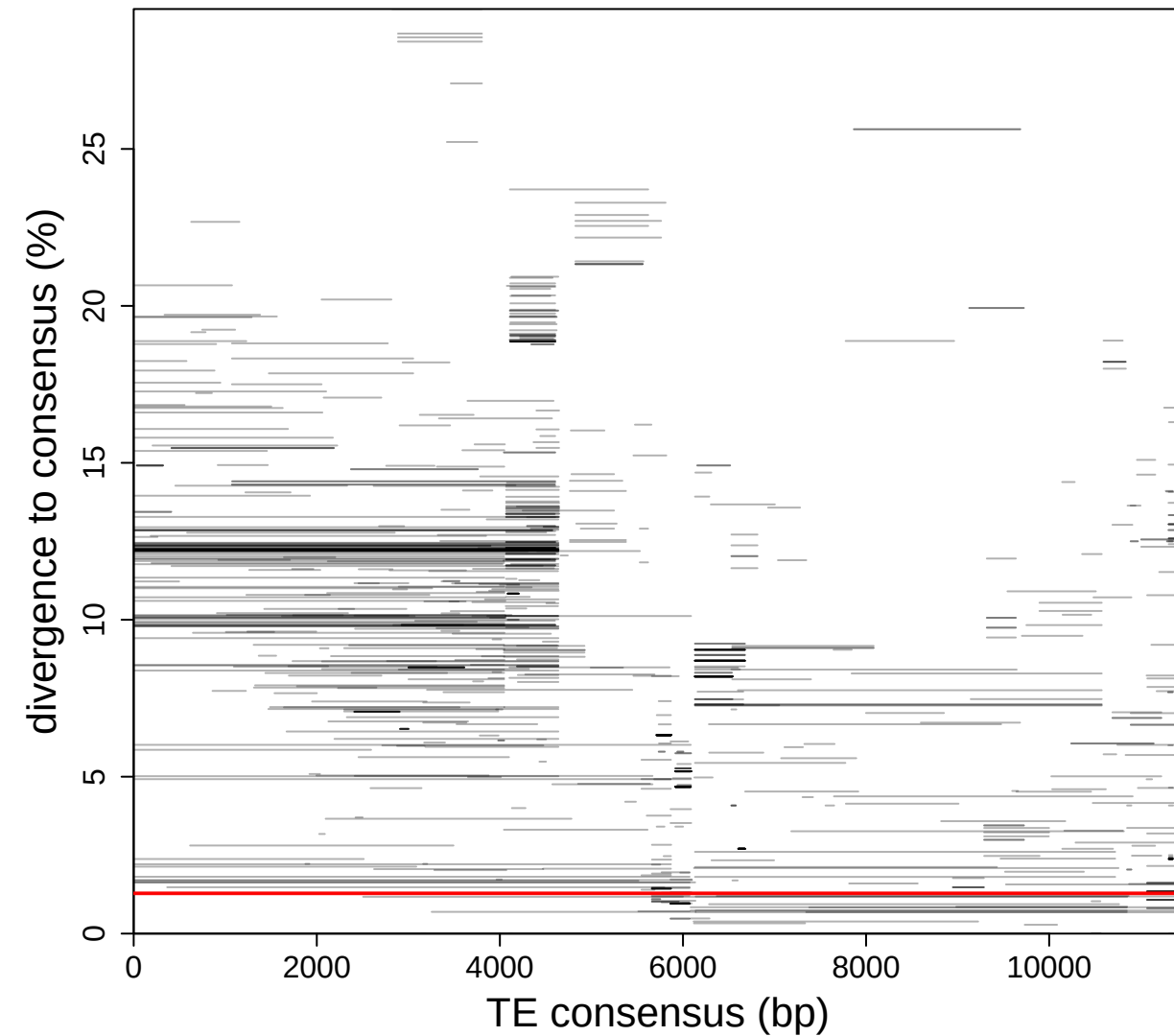
MSA length = 11454





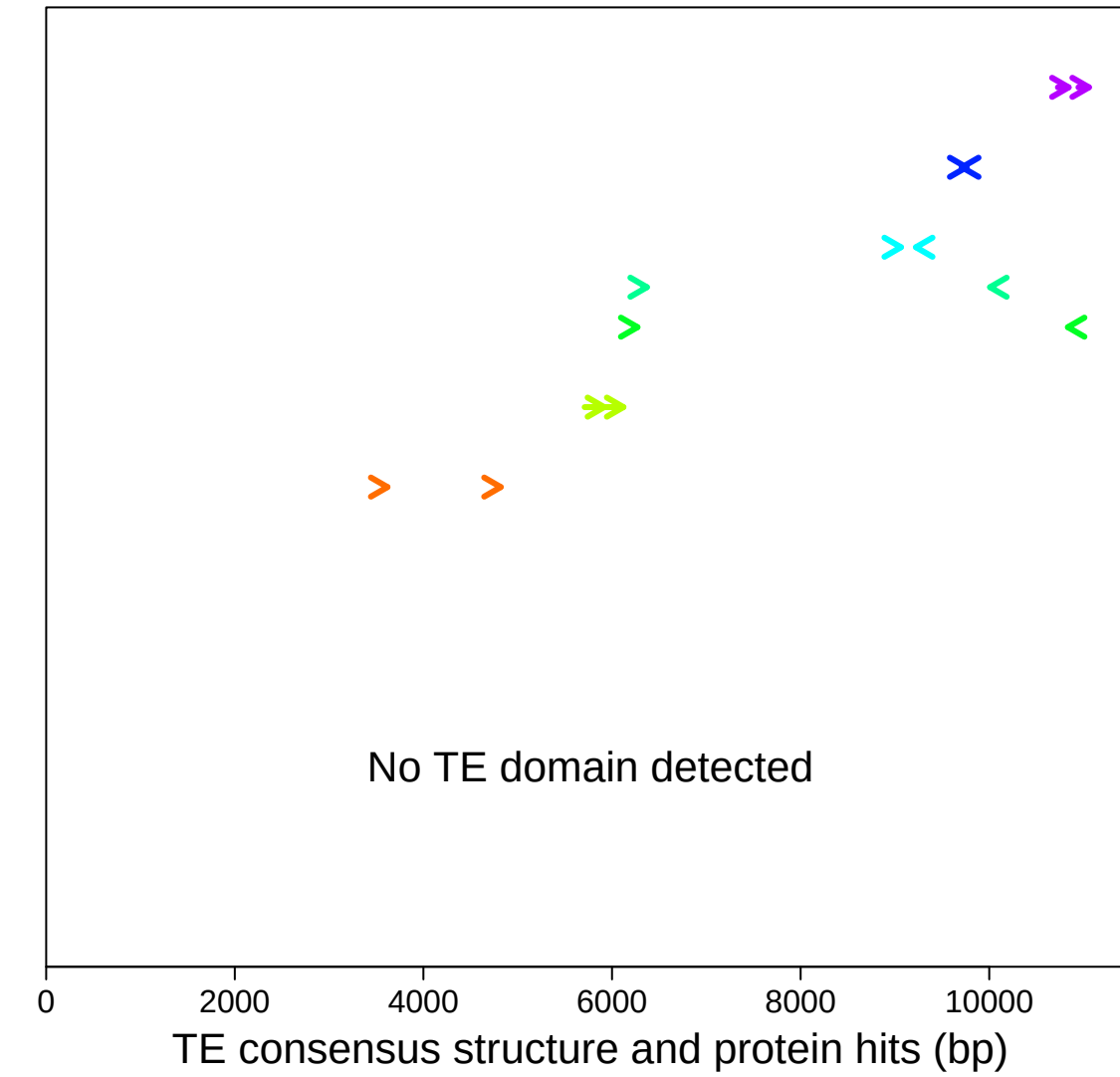
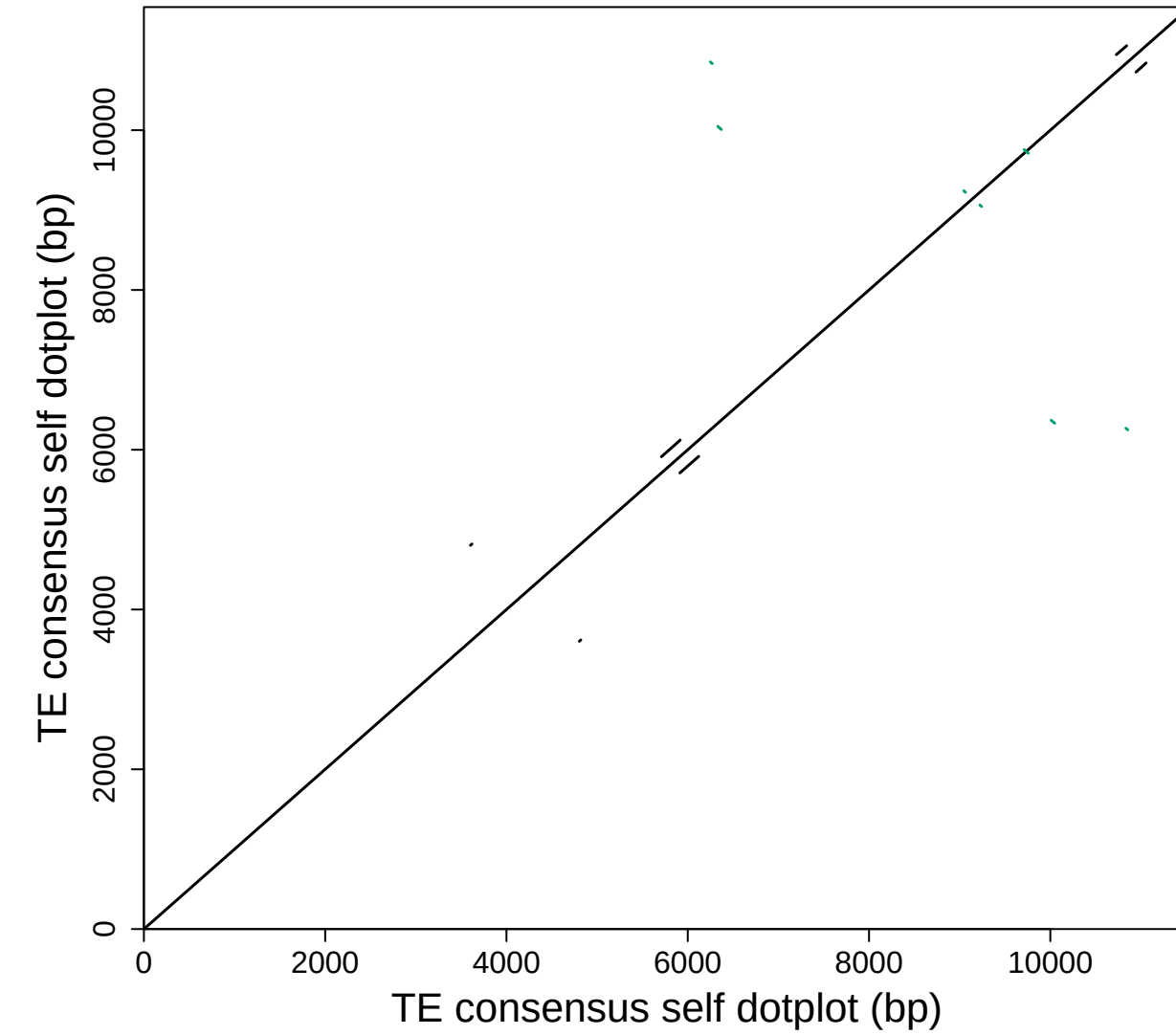
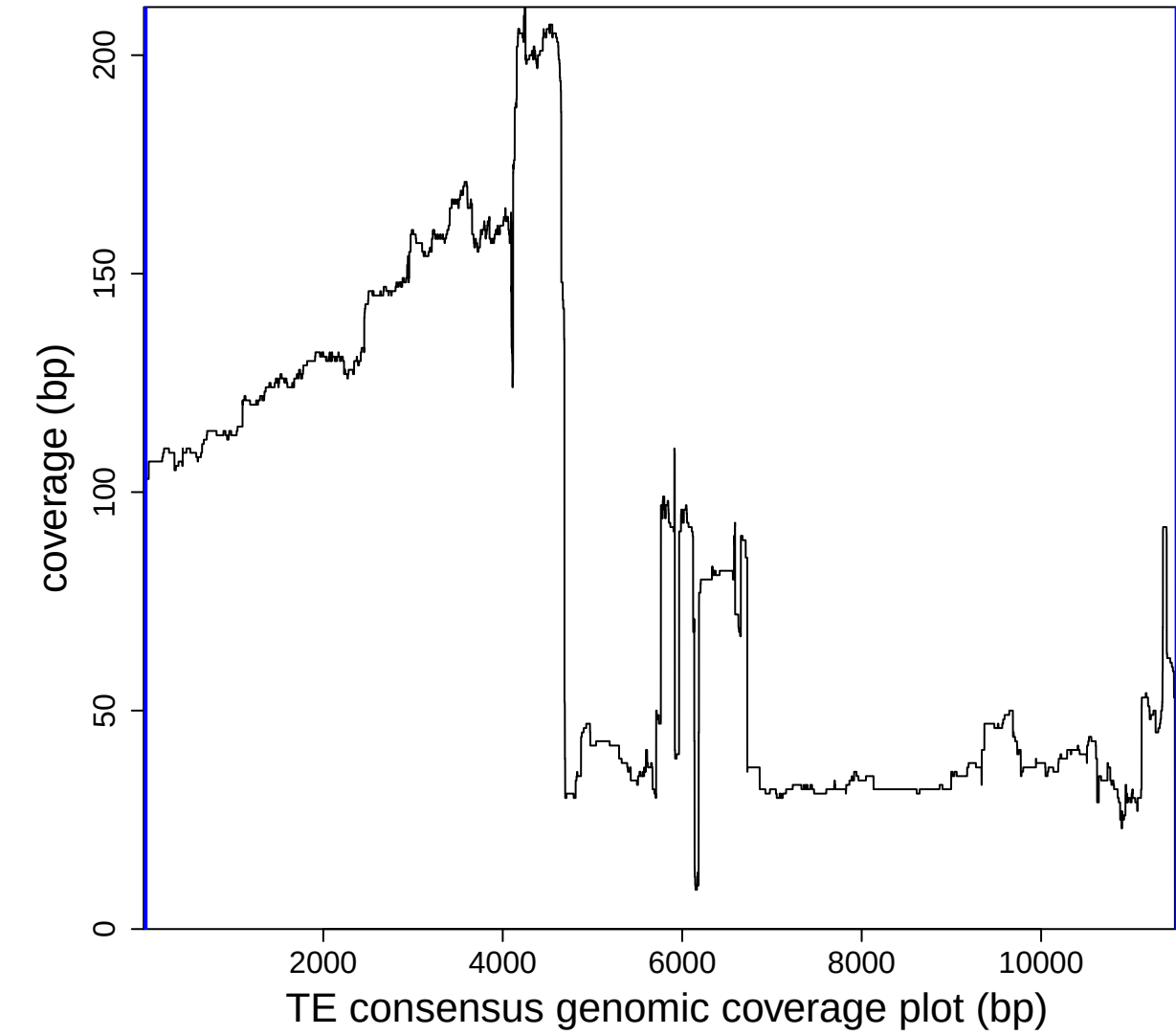
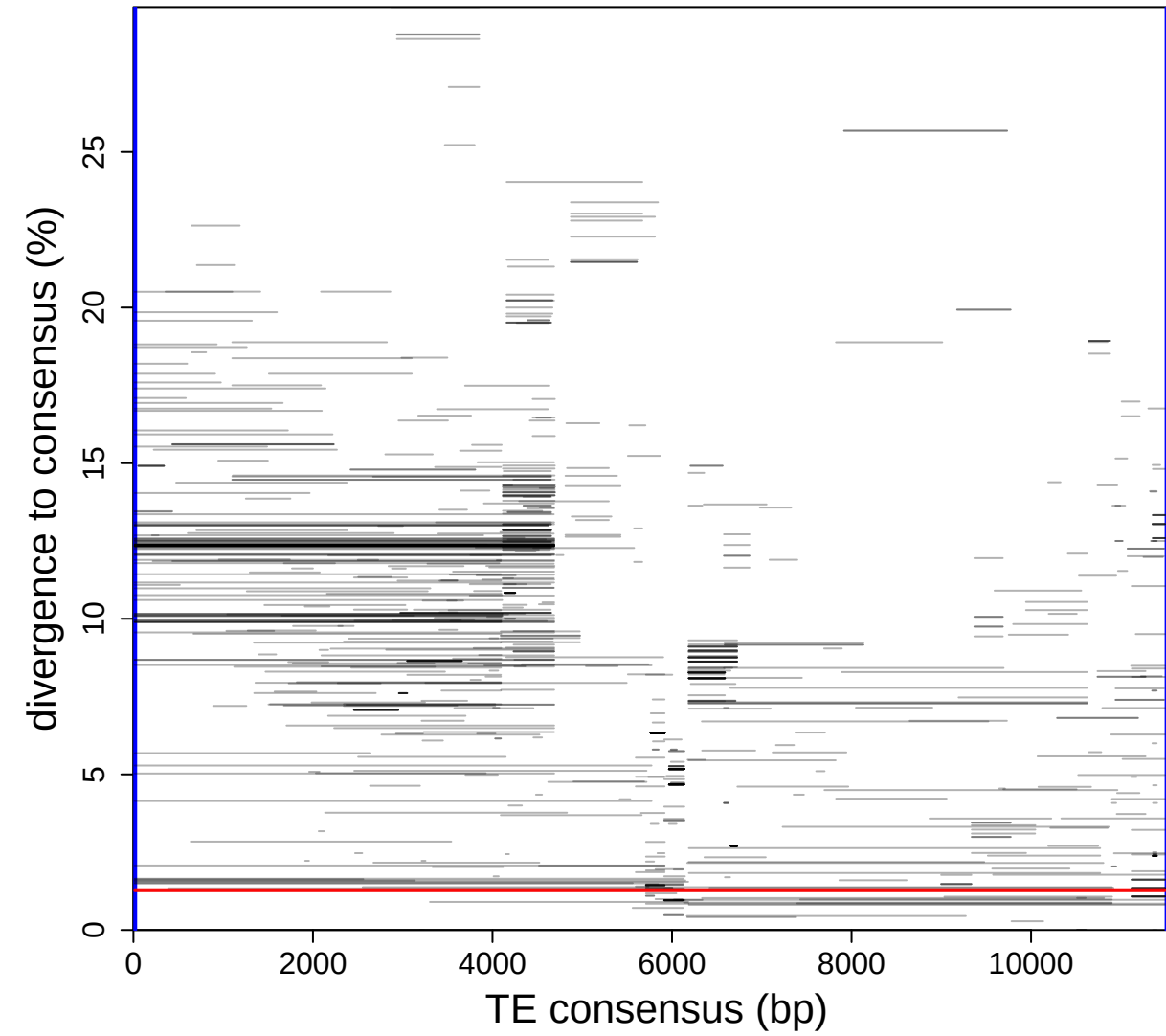
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 11454bp; fragments: 943; full length: 2 (>=10308.6bp)

After TEtrimmer 11454 bp

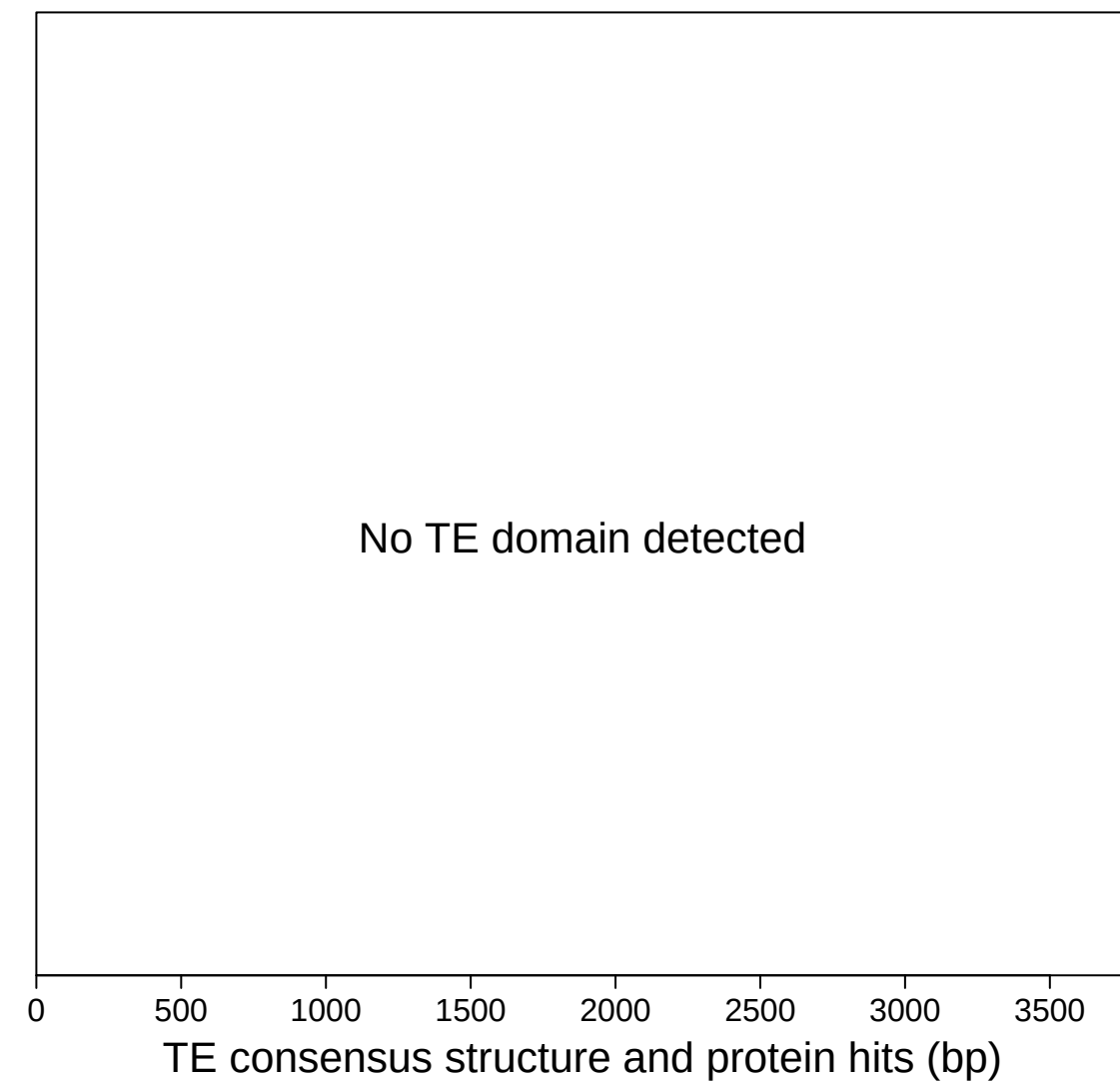
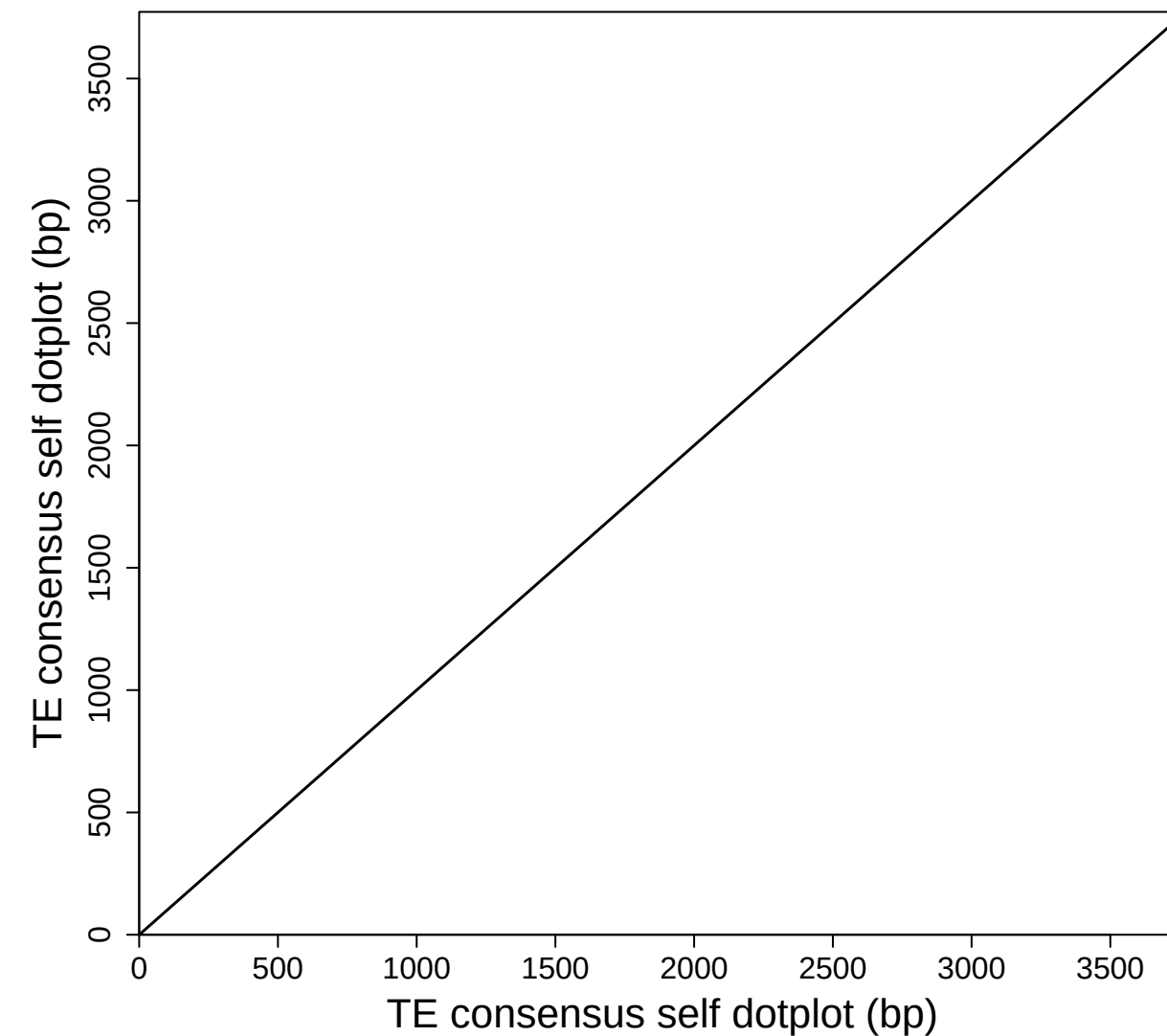
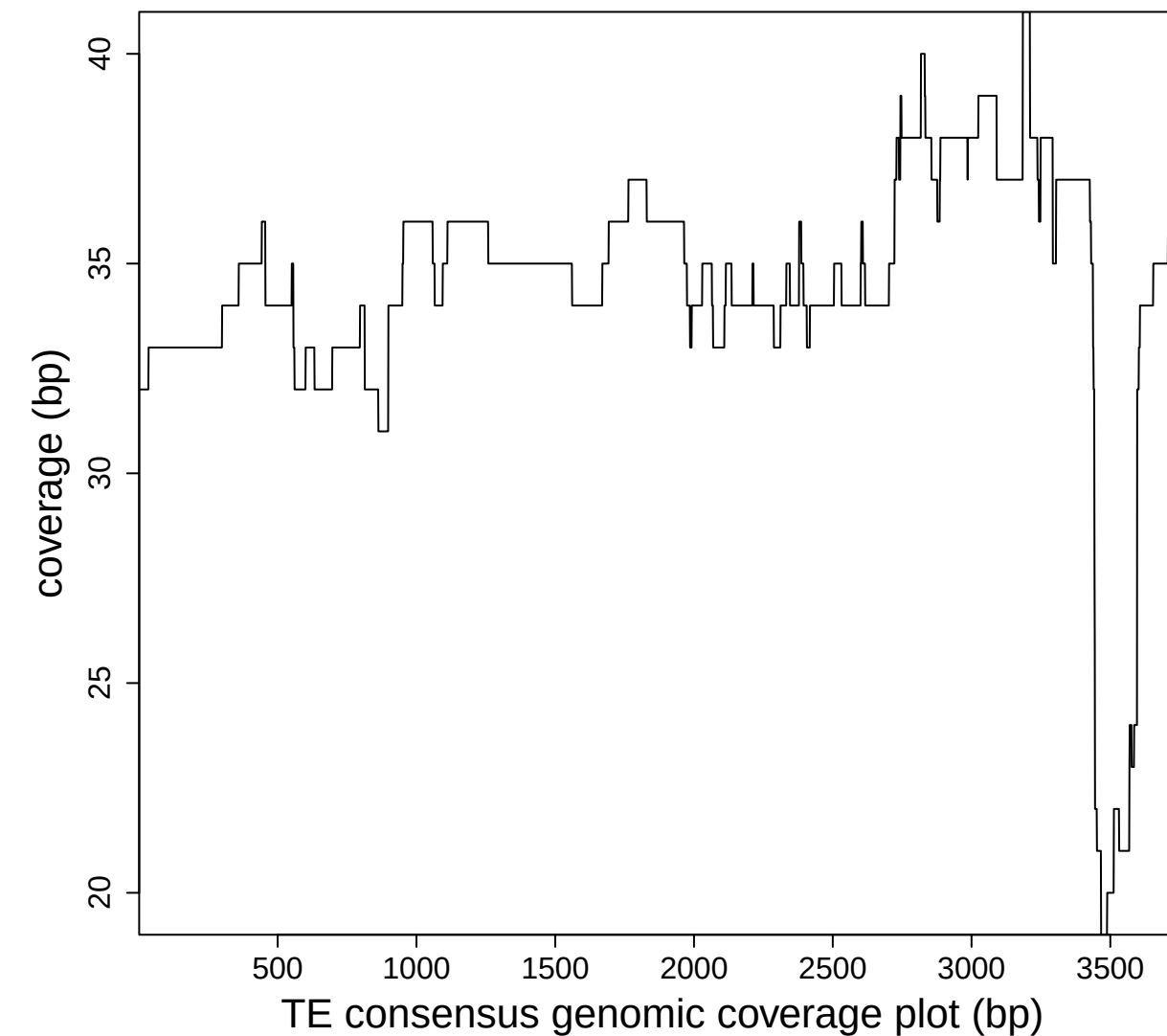
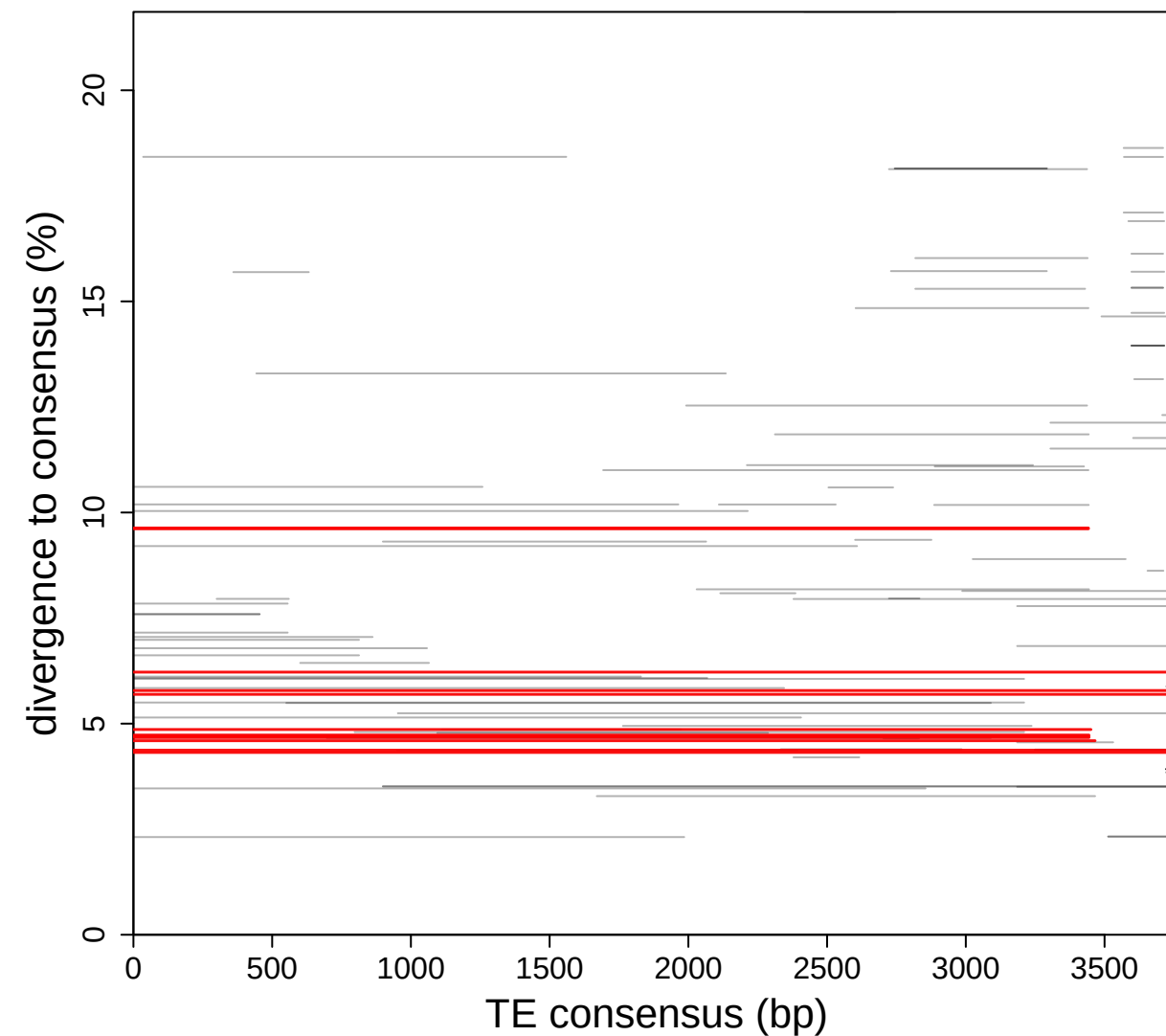


5.fasta.b.bed uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 11659bp; fragments: 921; full length: 2 (>=10493.1bp)

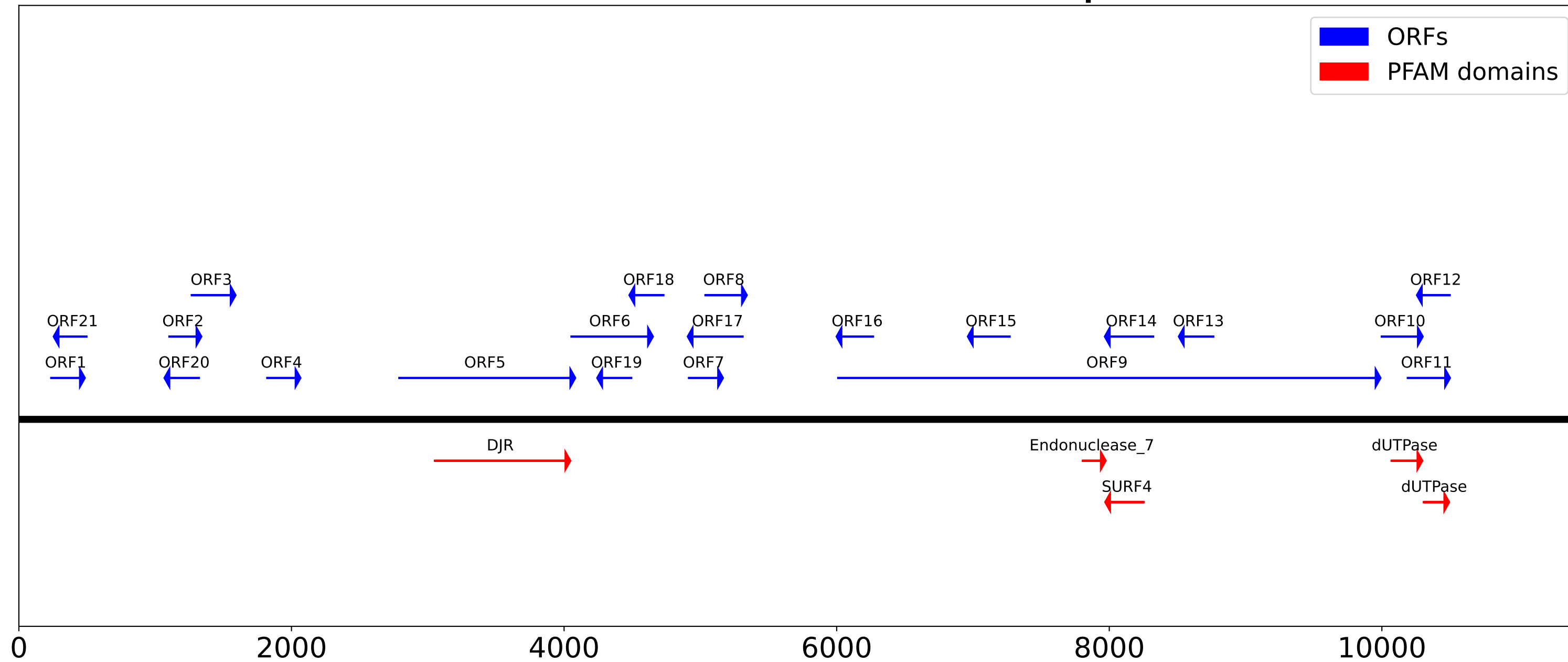
After TEtrimmer Extended plot Blue lines are boundaries



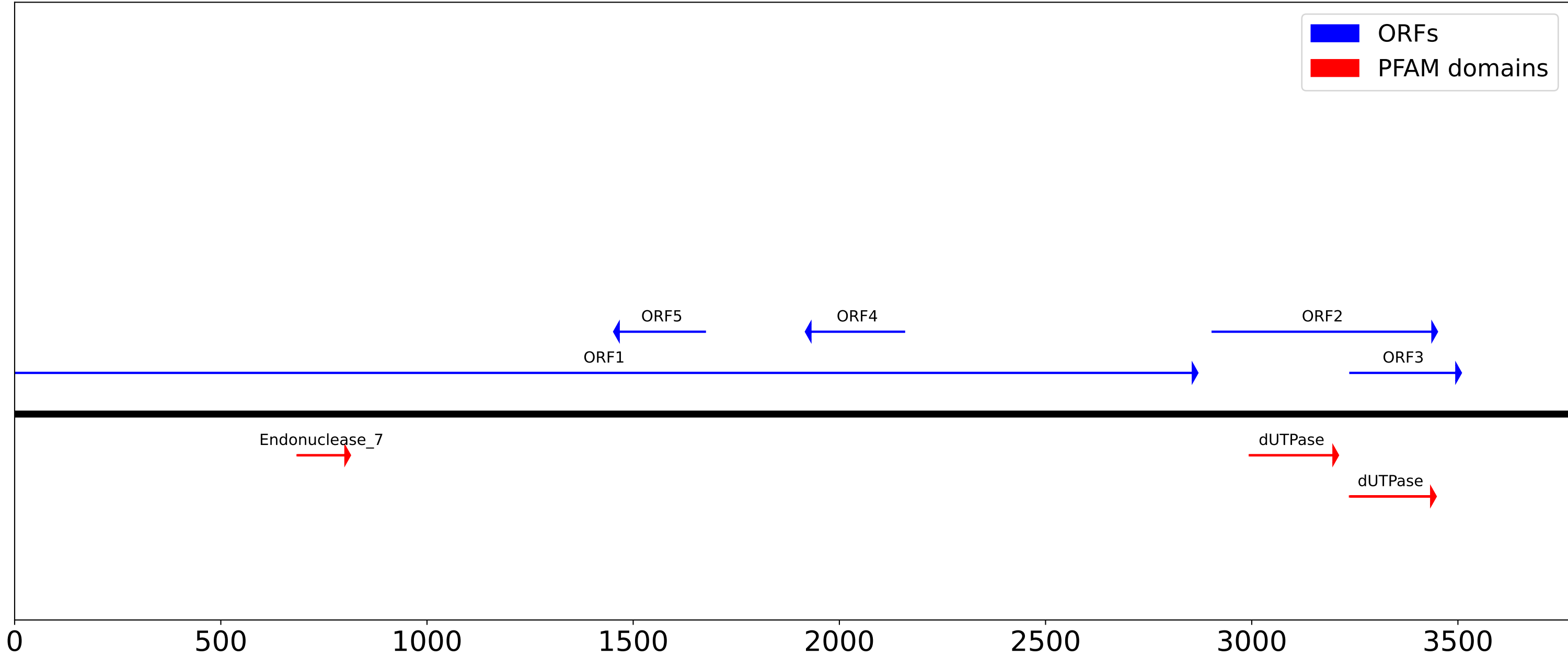
TE: rnd_5_family_7365
size: 3772bp; fragments: 110; full length: 12 (≥ 3394.8 bp)



After TEtrimmer ORF and PFAM domain plot



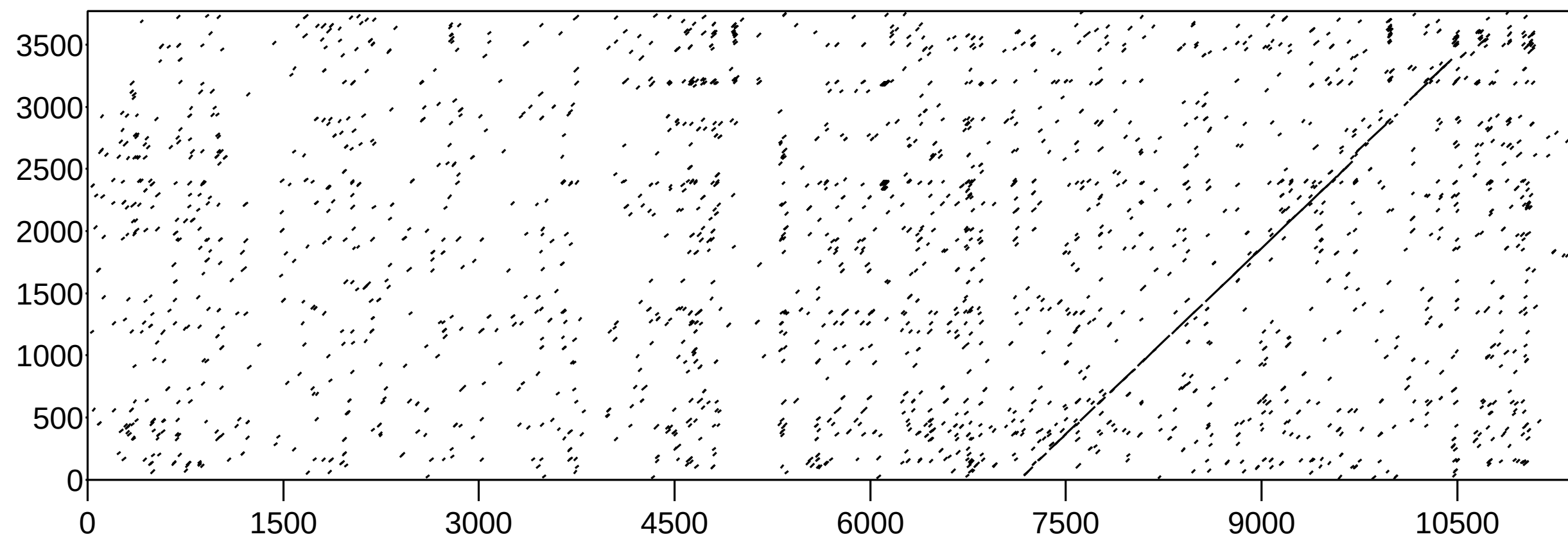
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)