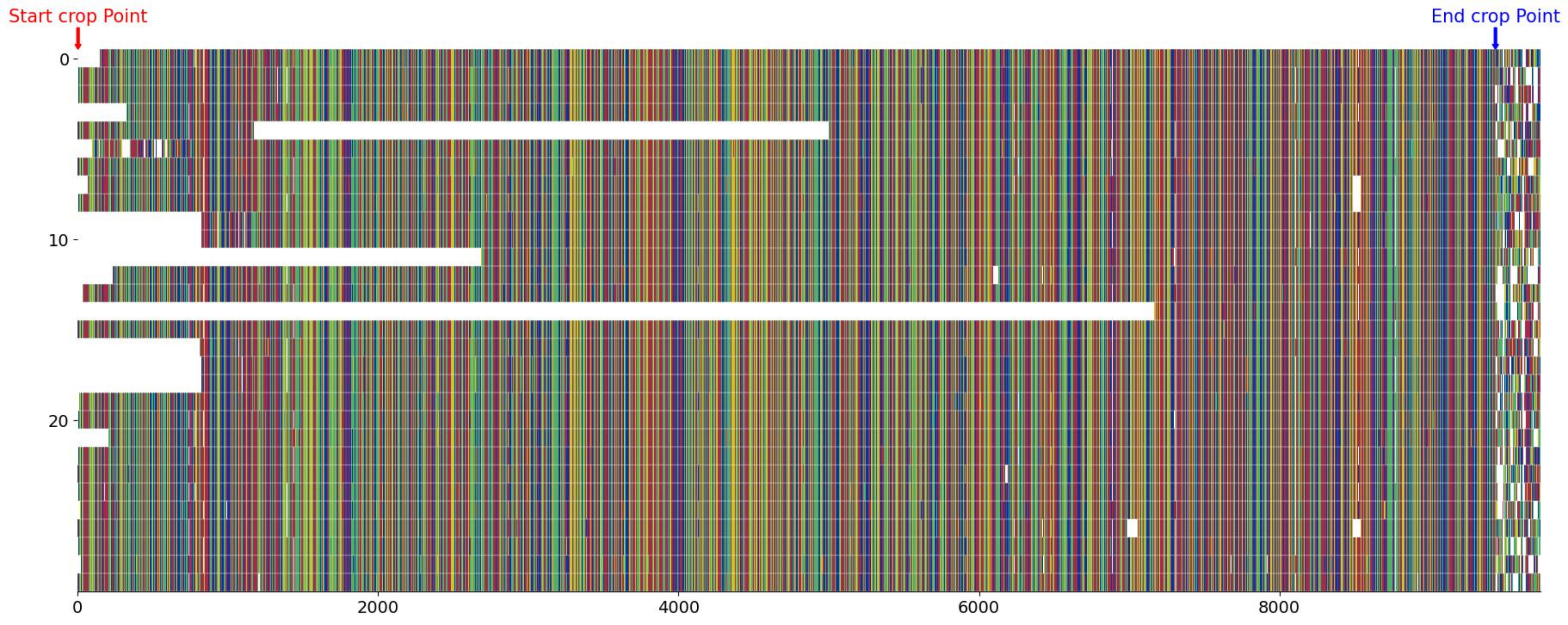


Start crop Point

End crop Point

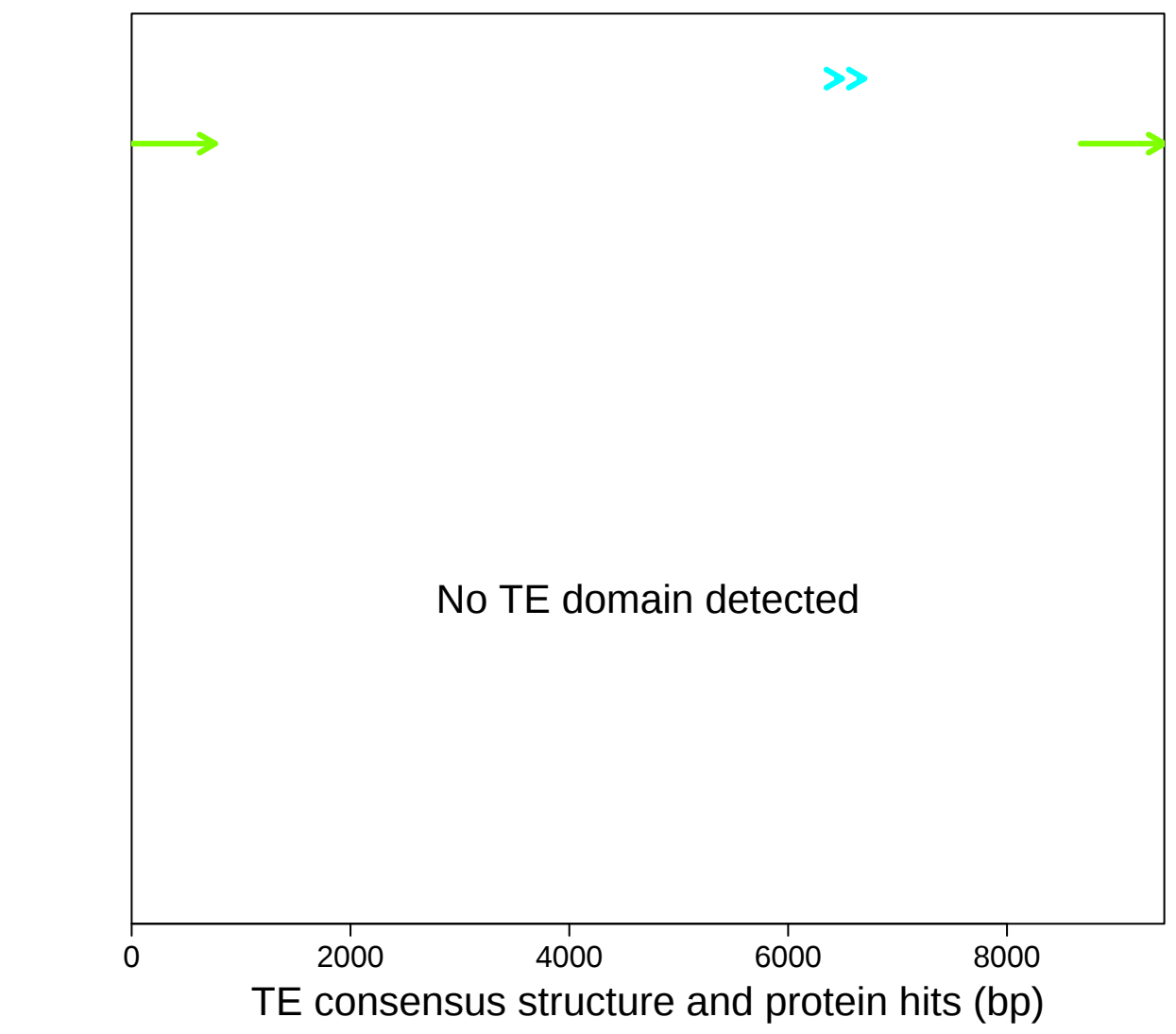
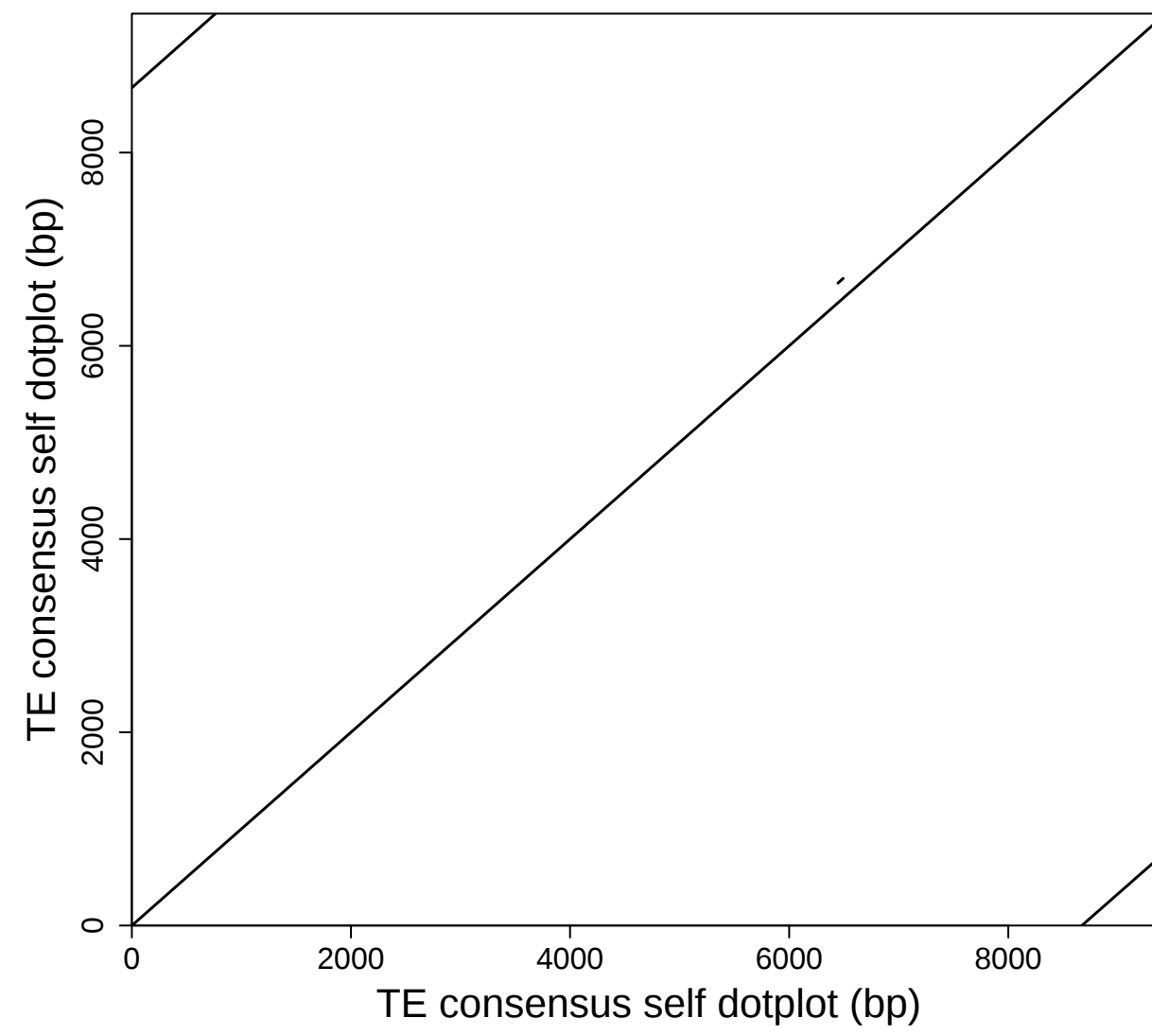
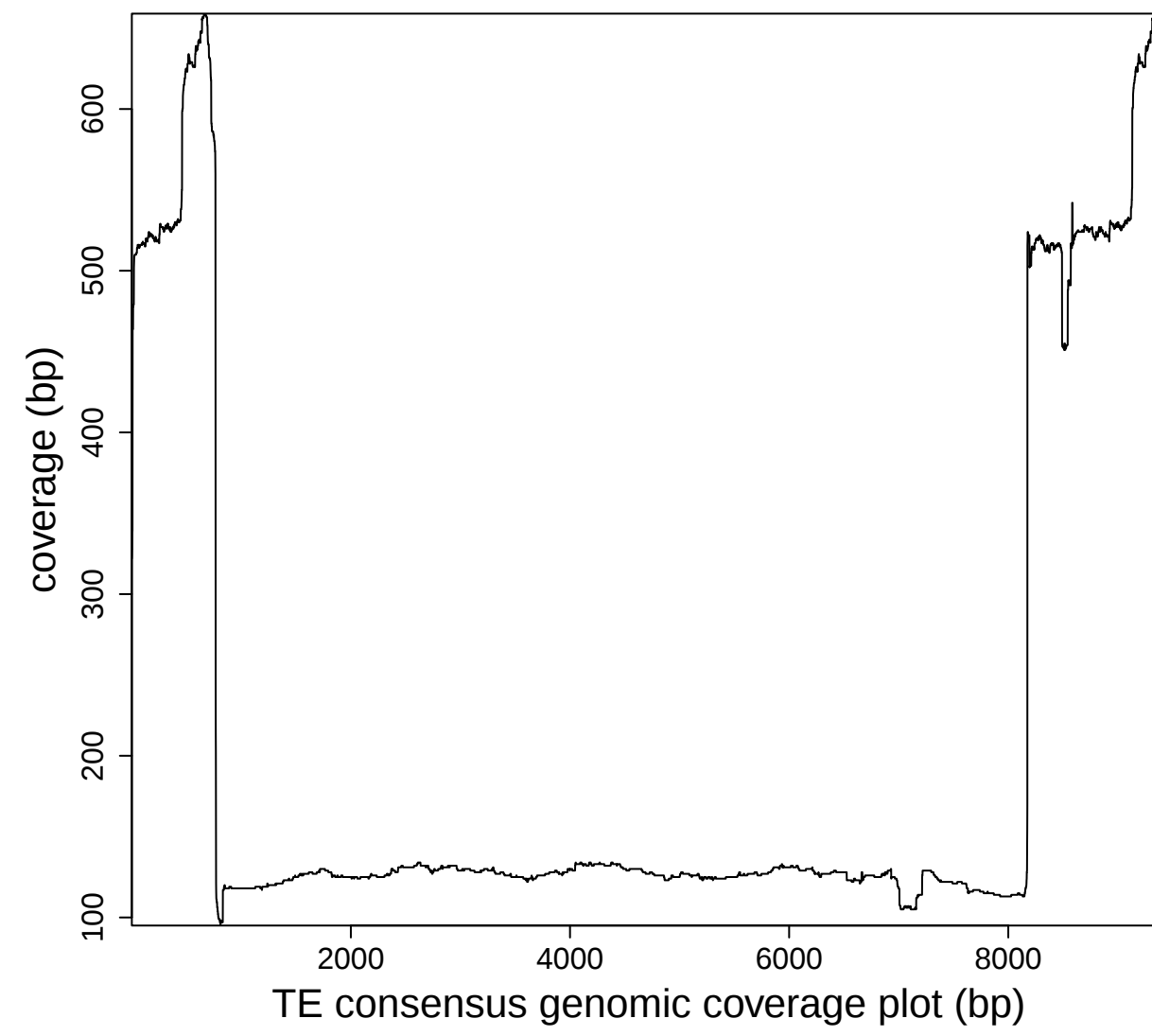
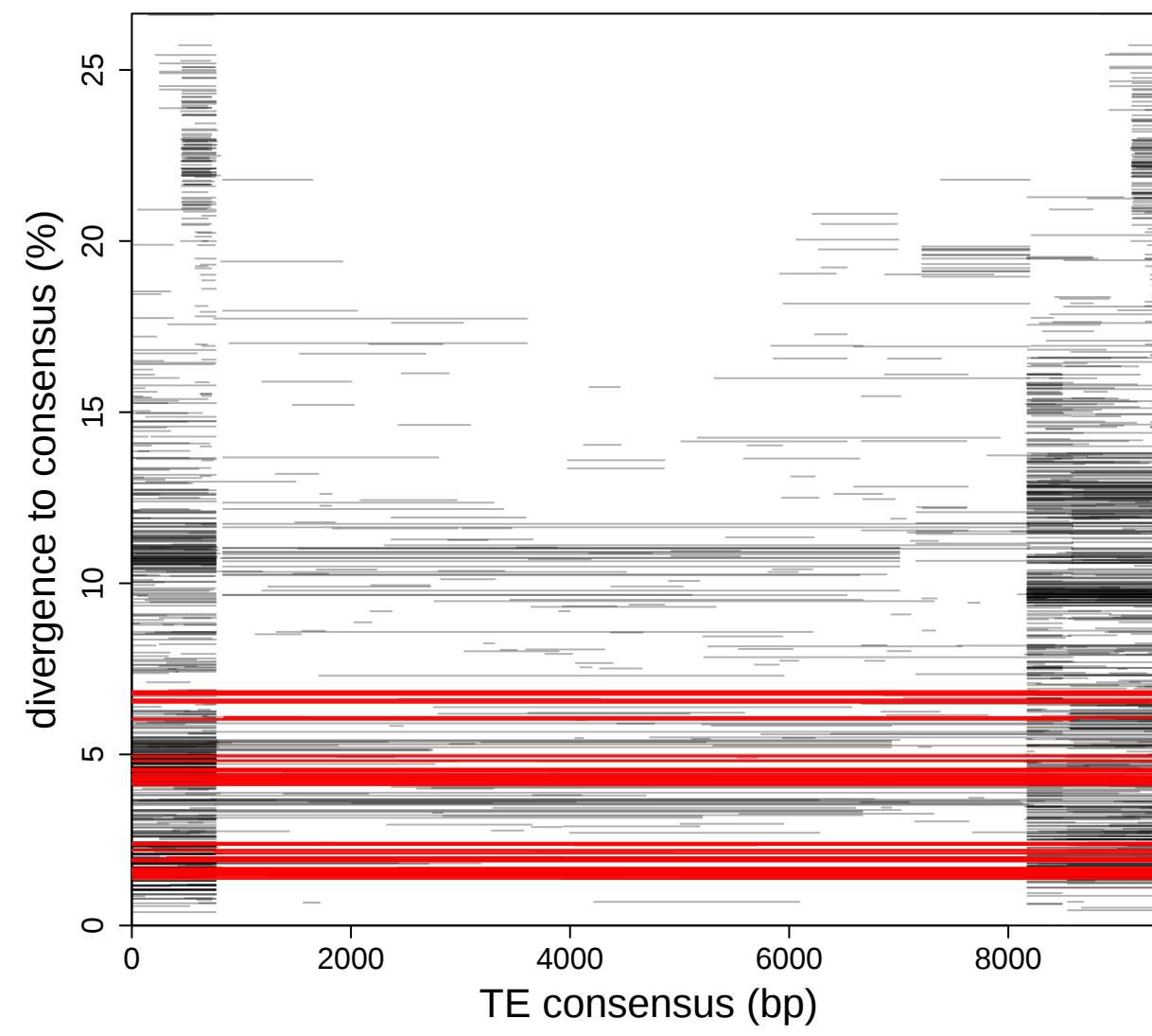
MSA length = 9438





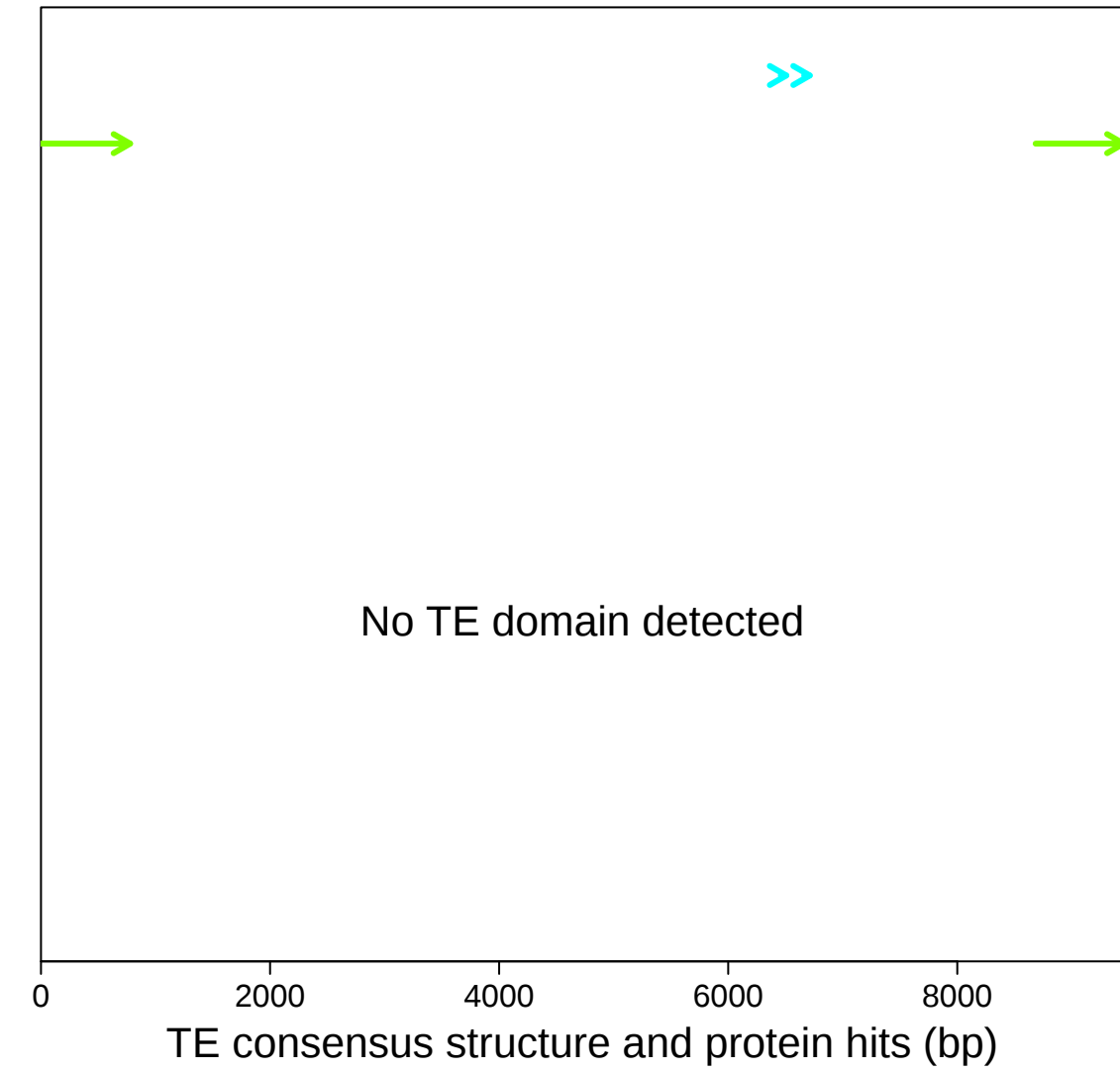
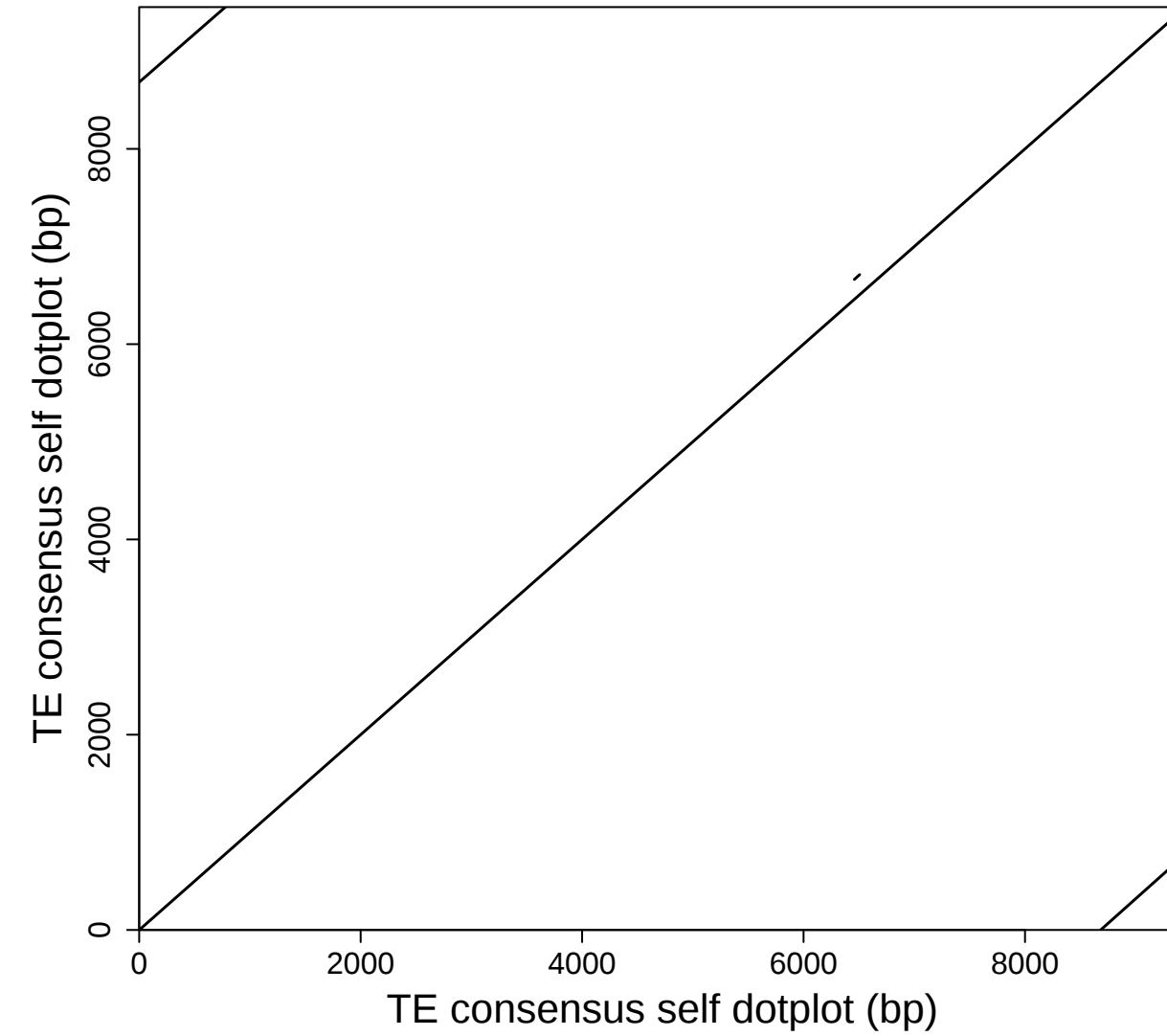
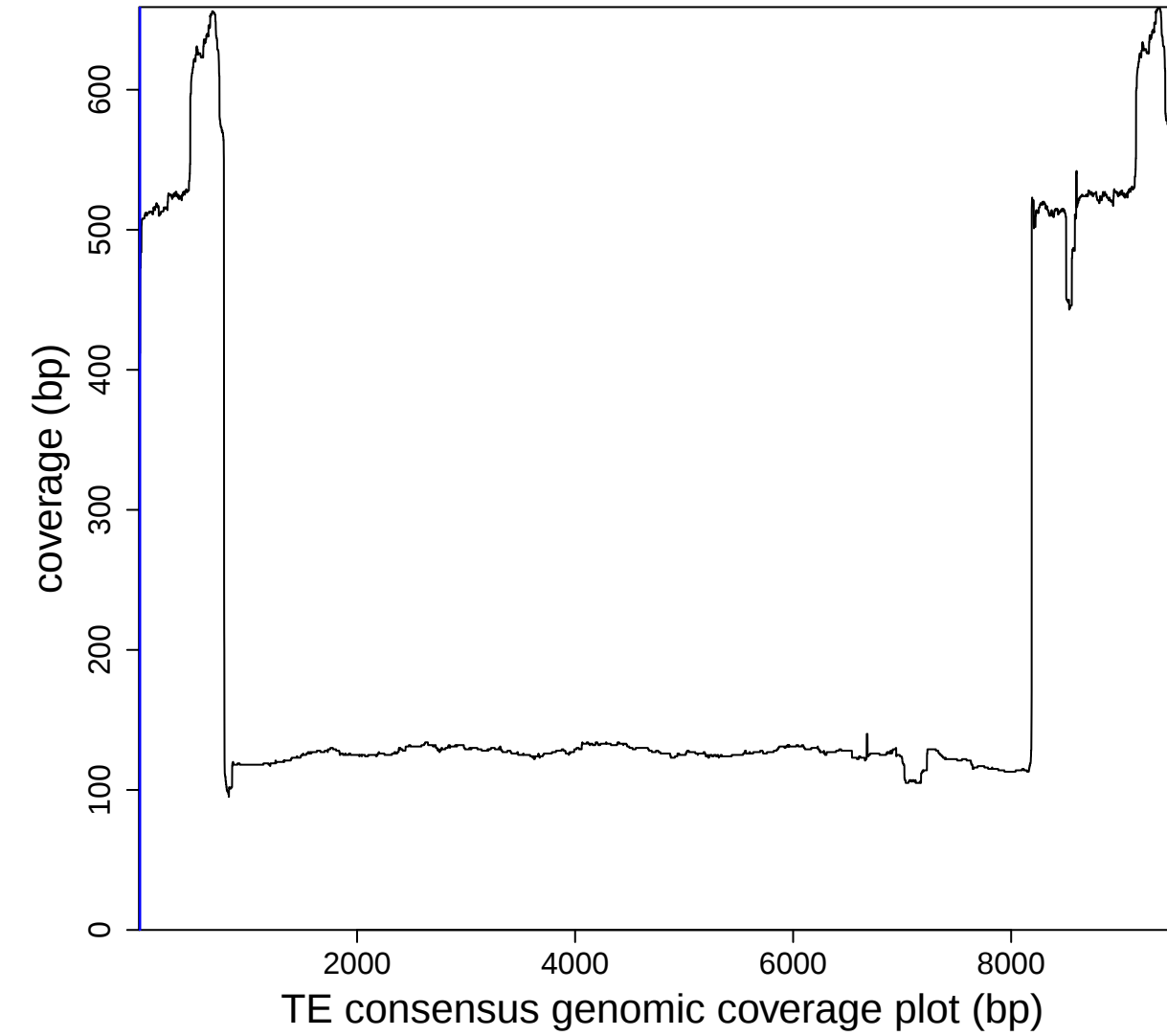
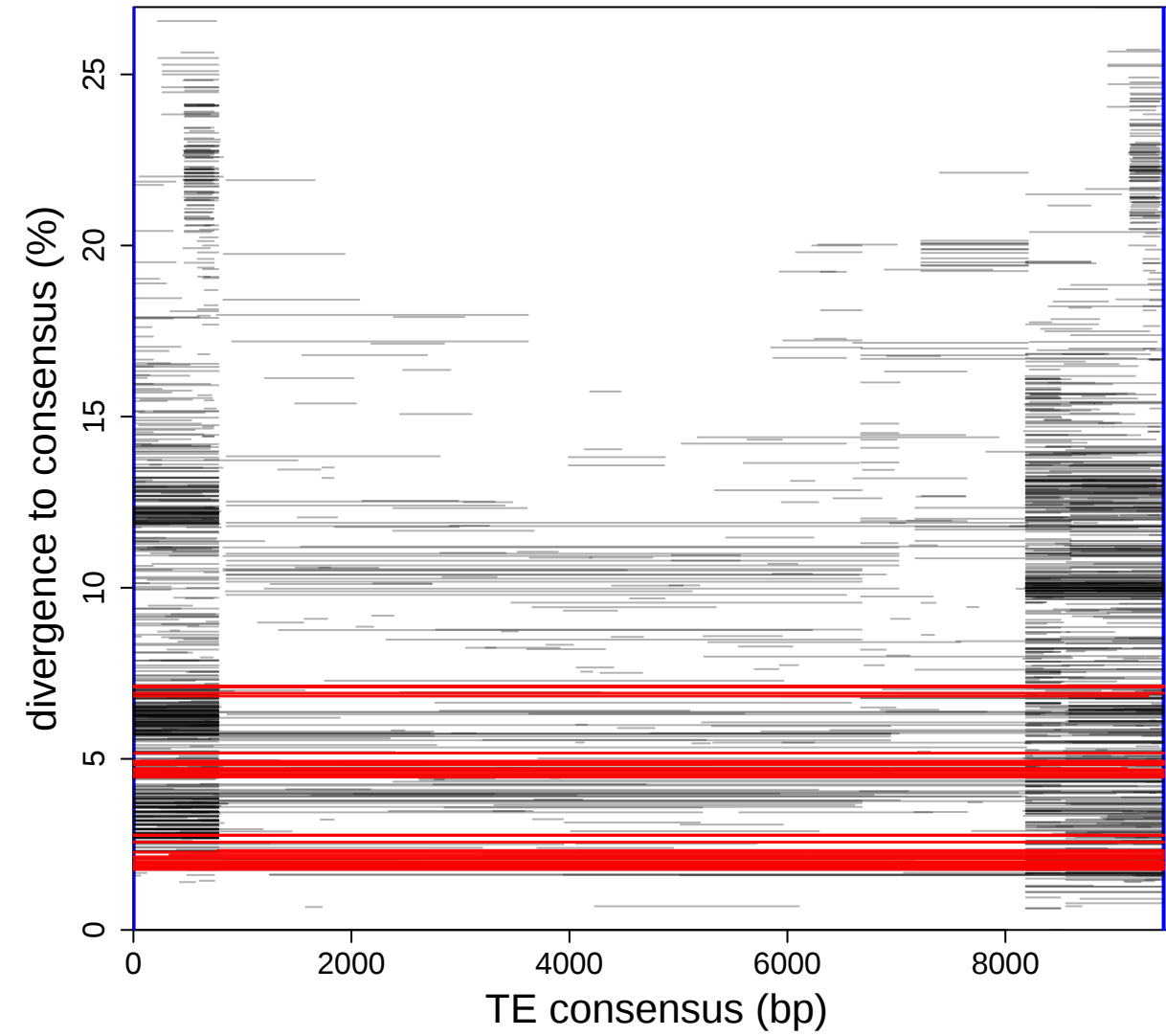
ed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 9438bp; fragments: 1804; full length: 43 (>=8494.2bp)

After TEtrimmer 9438 bp



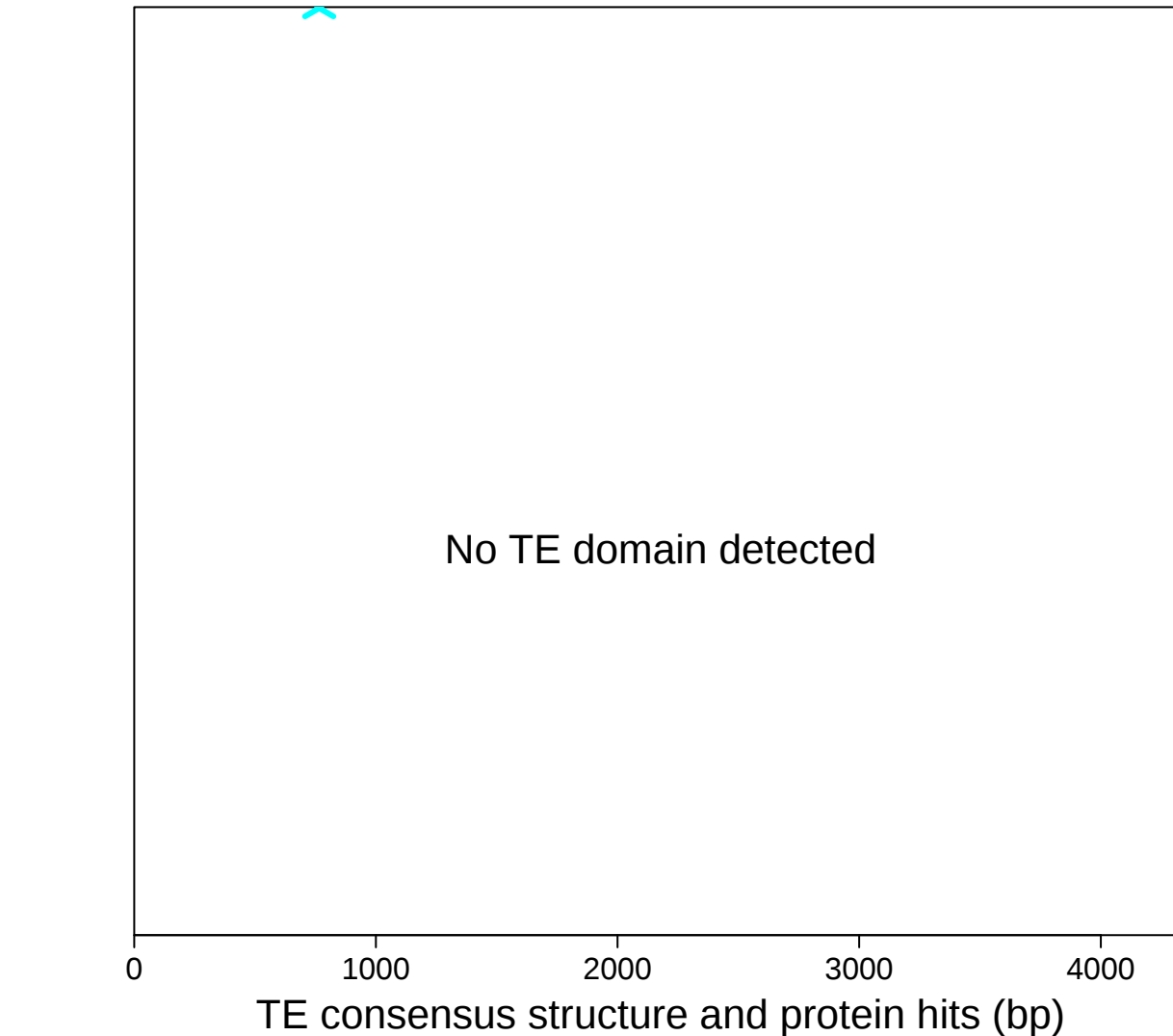
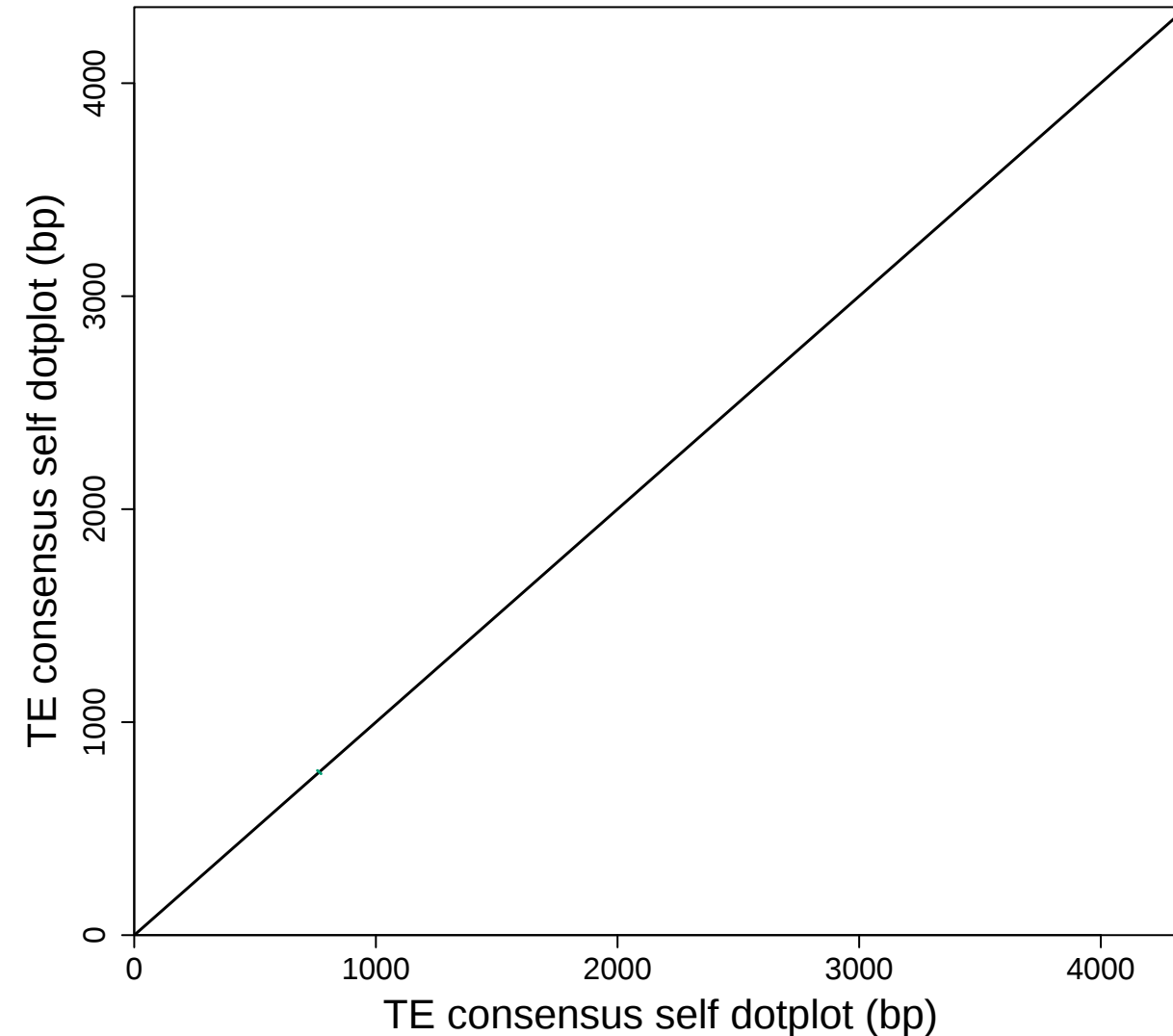
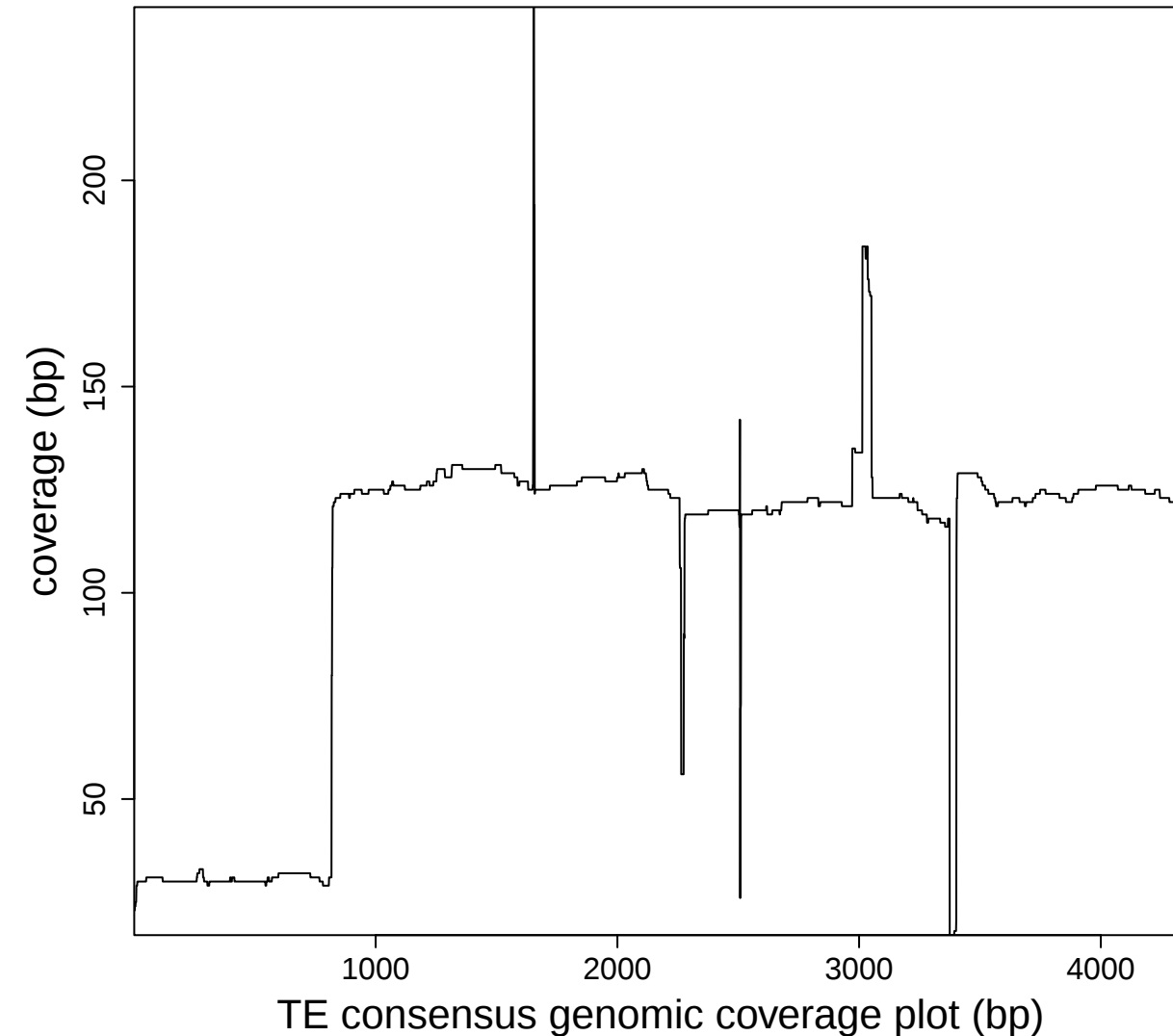
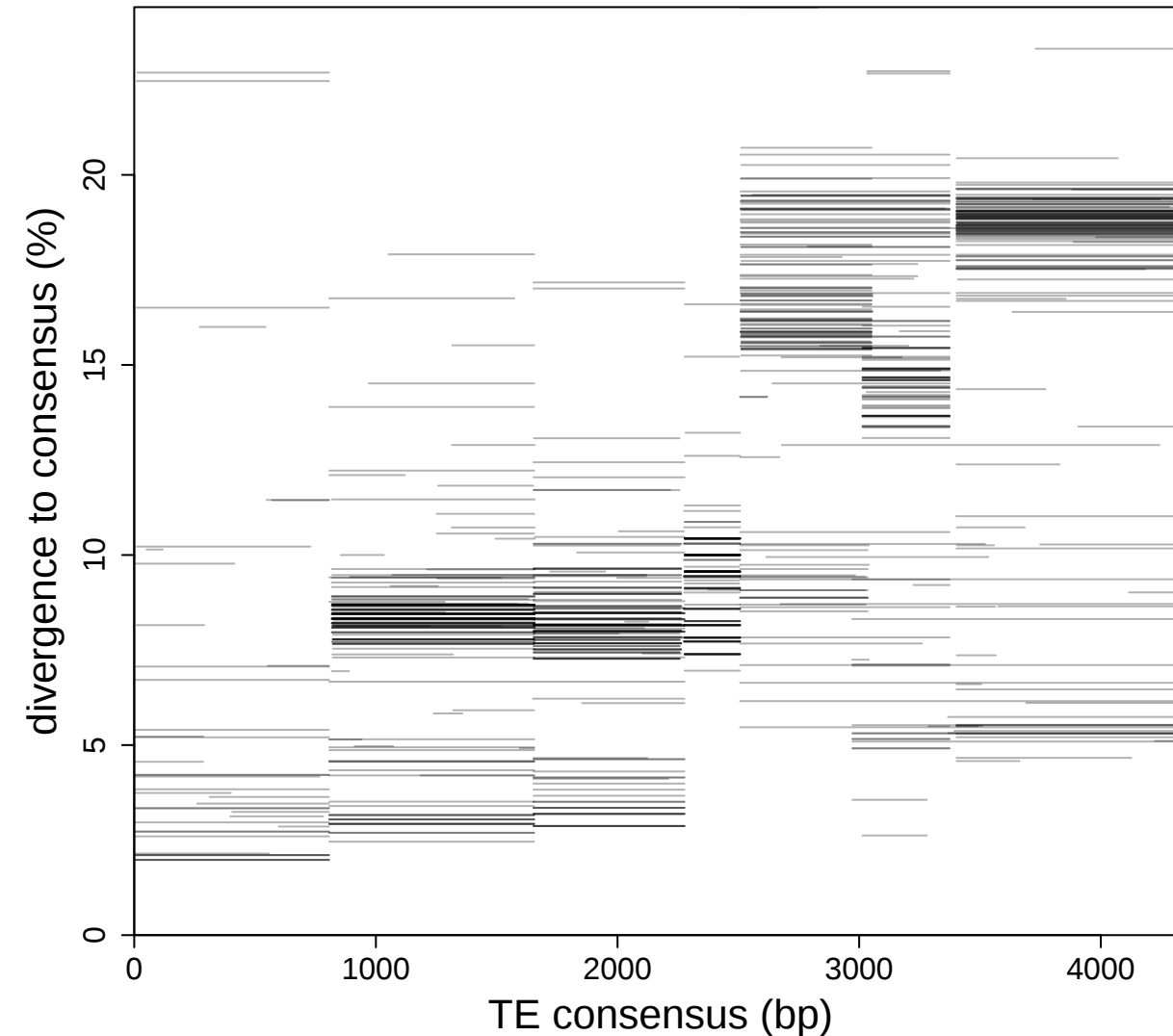
ly_822.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcIn.fa_aln.fa
size: 9602bp; fragments: 1813; full length: 37 (>=8641.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

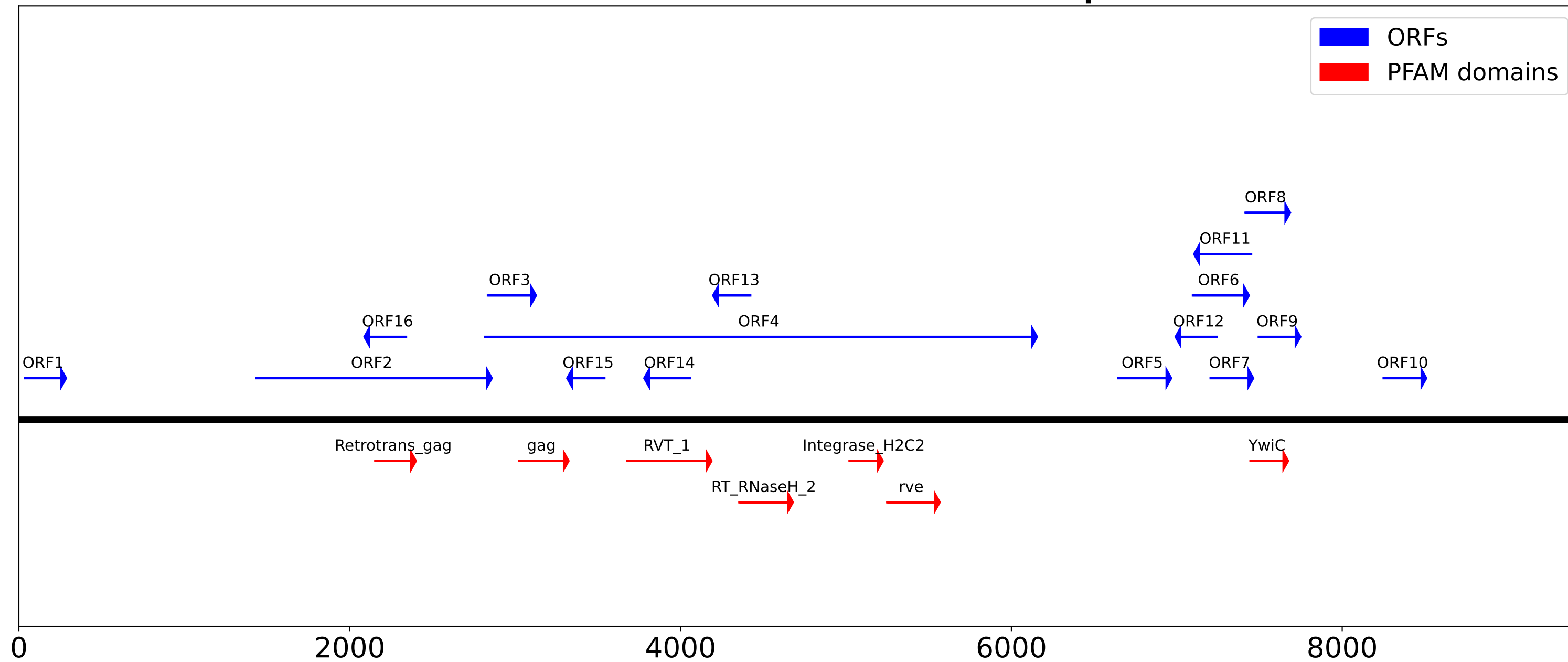


TE: ltr_1_family_822
size: 4357bp; fragments: 757; full length: 0 (>=3921.3bp)

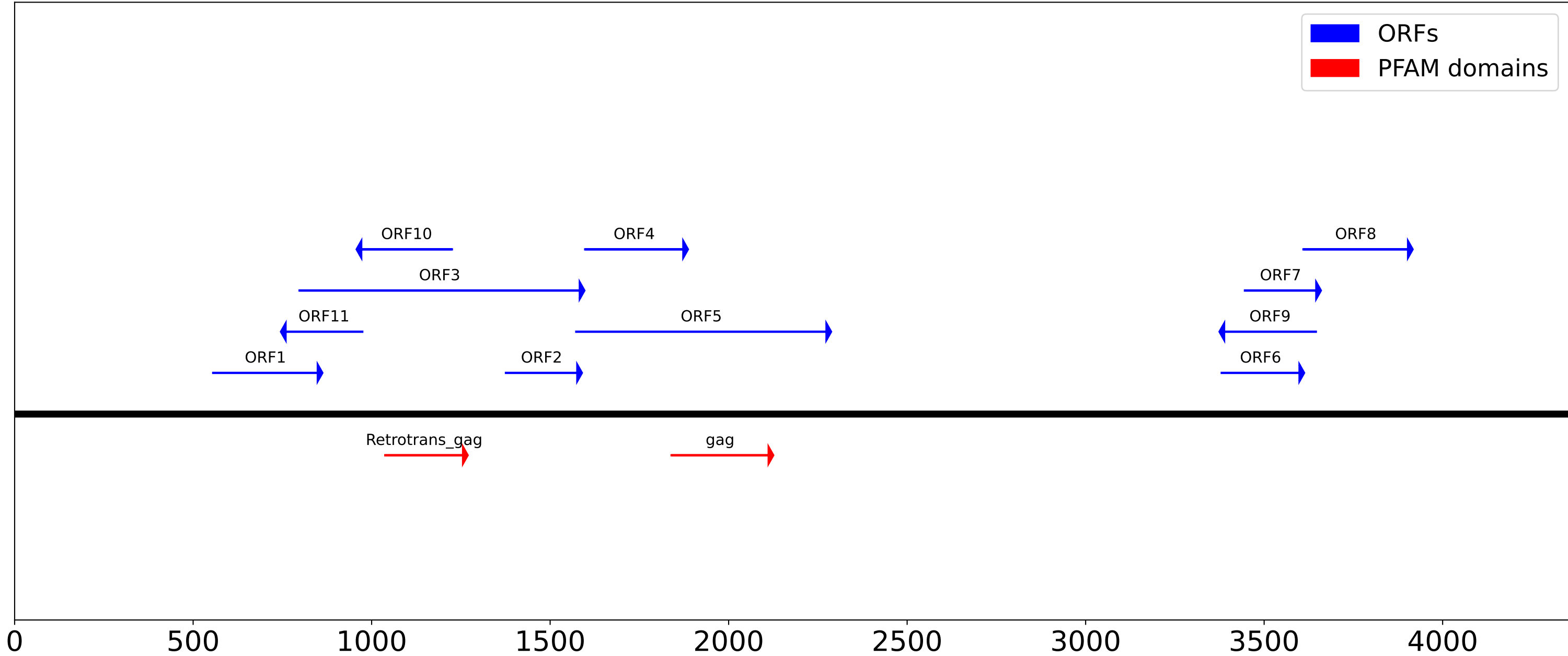
Before TEtrimmer 4357 bp



After TEtrimmer ORF and PFAM domain plot



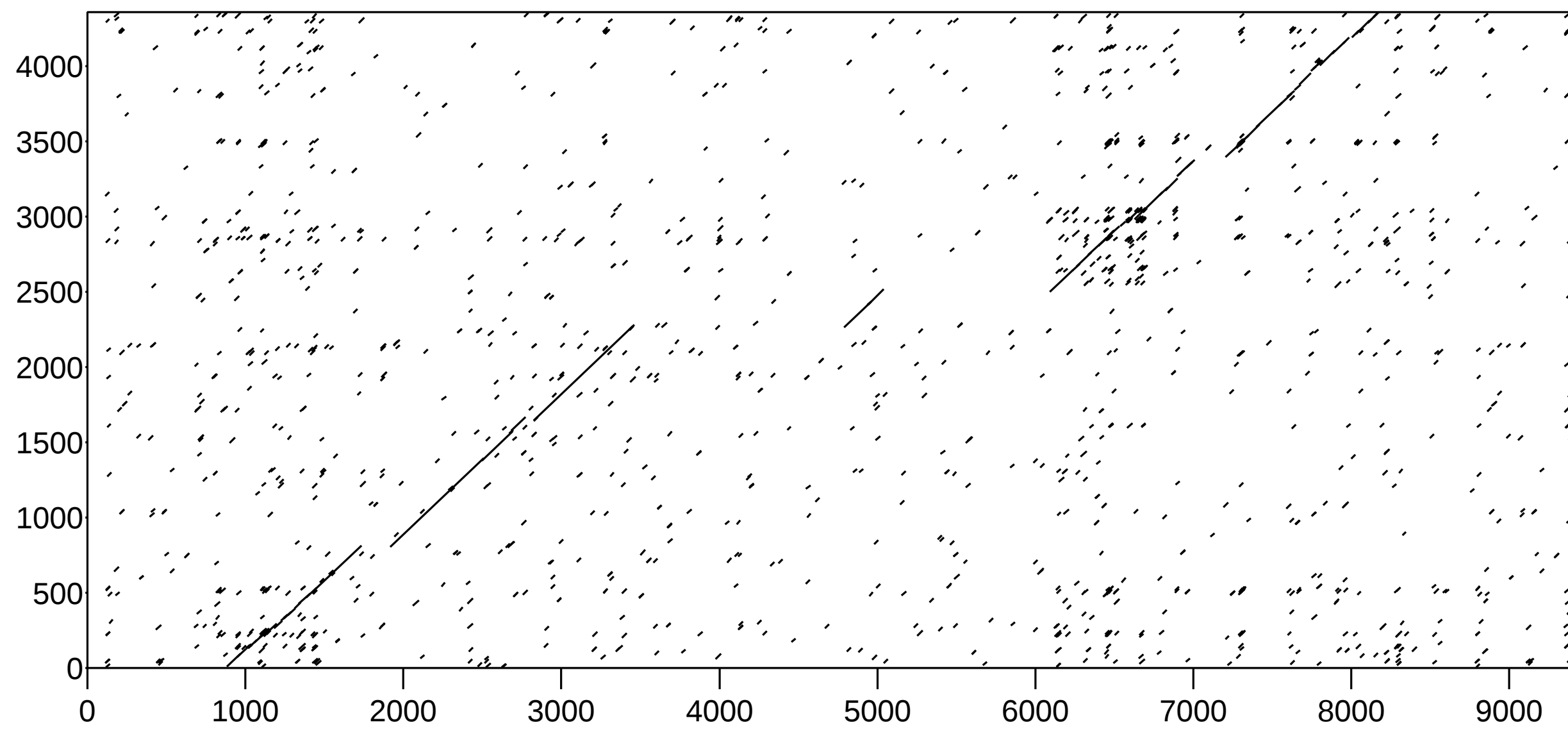
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)