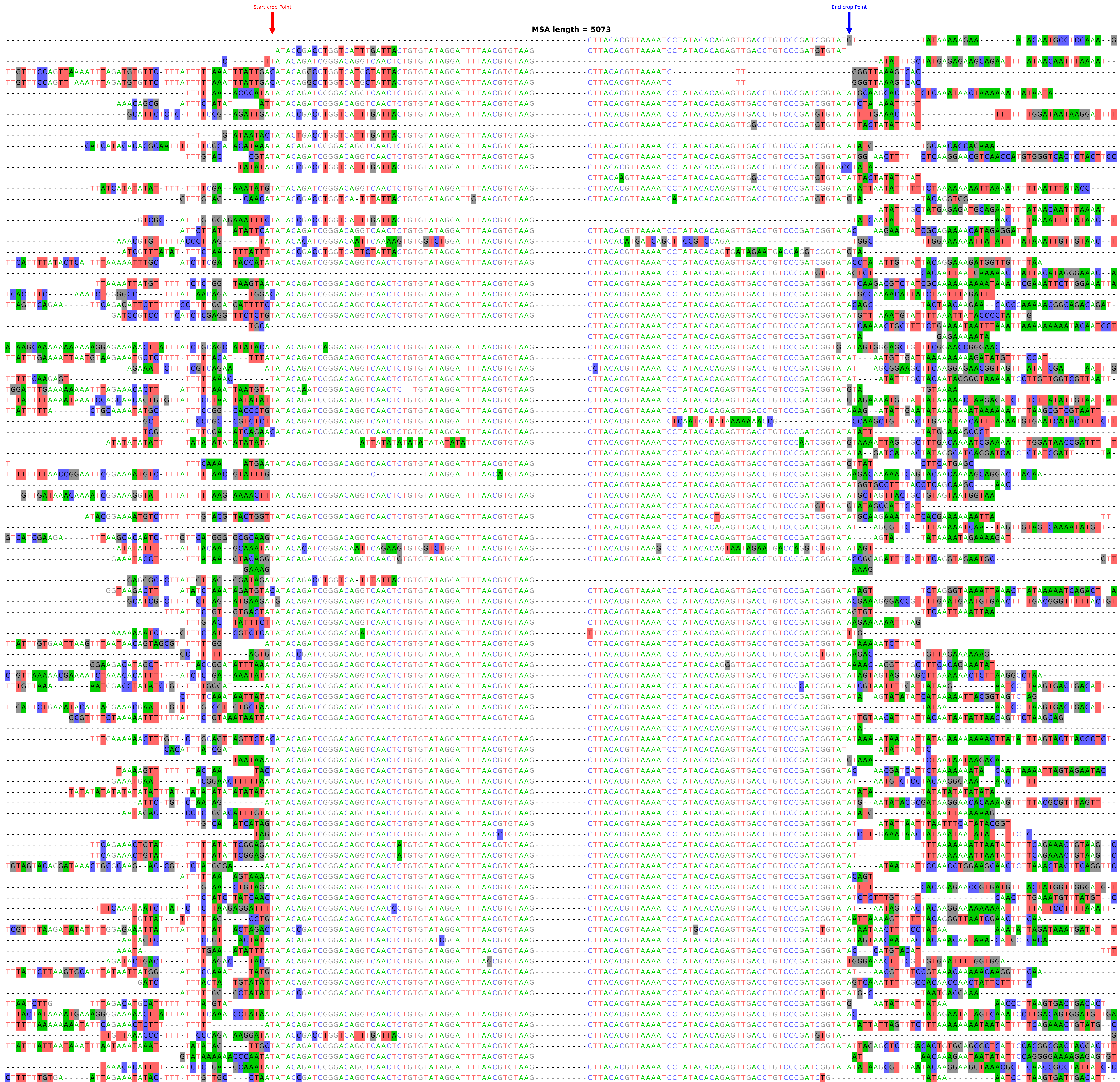
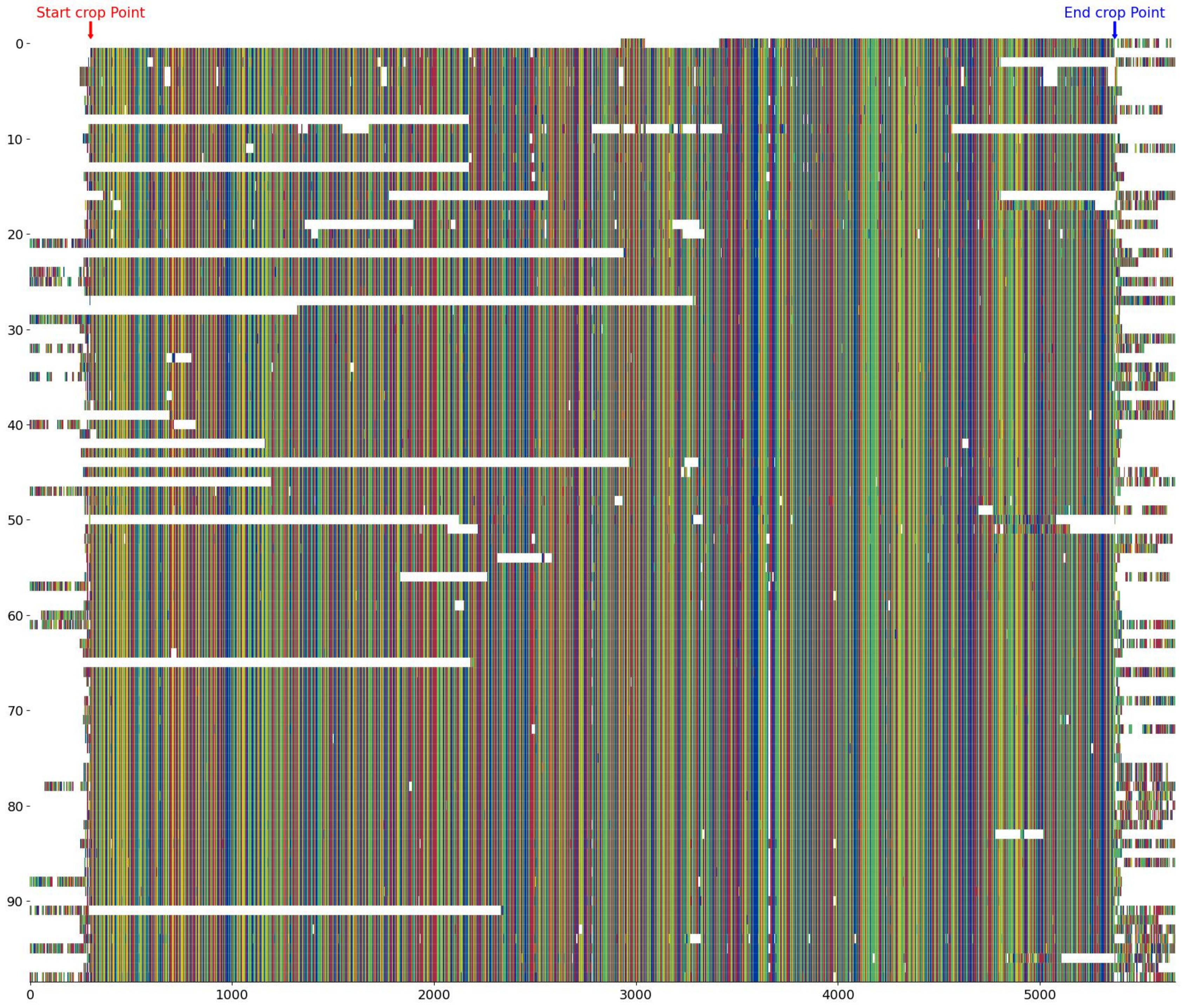
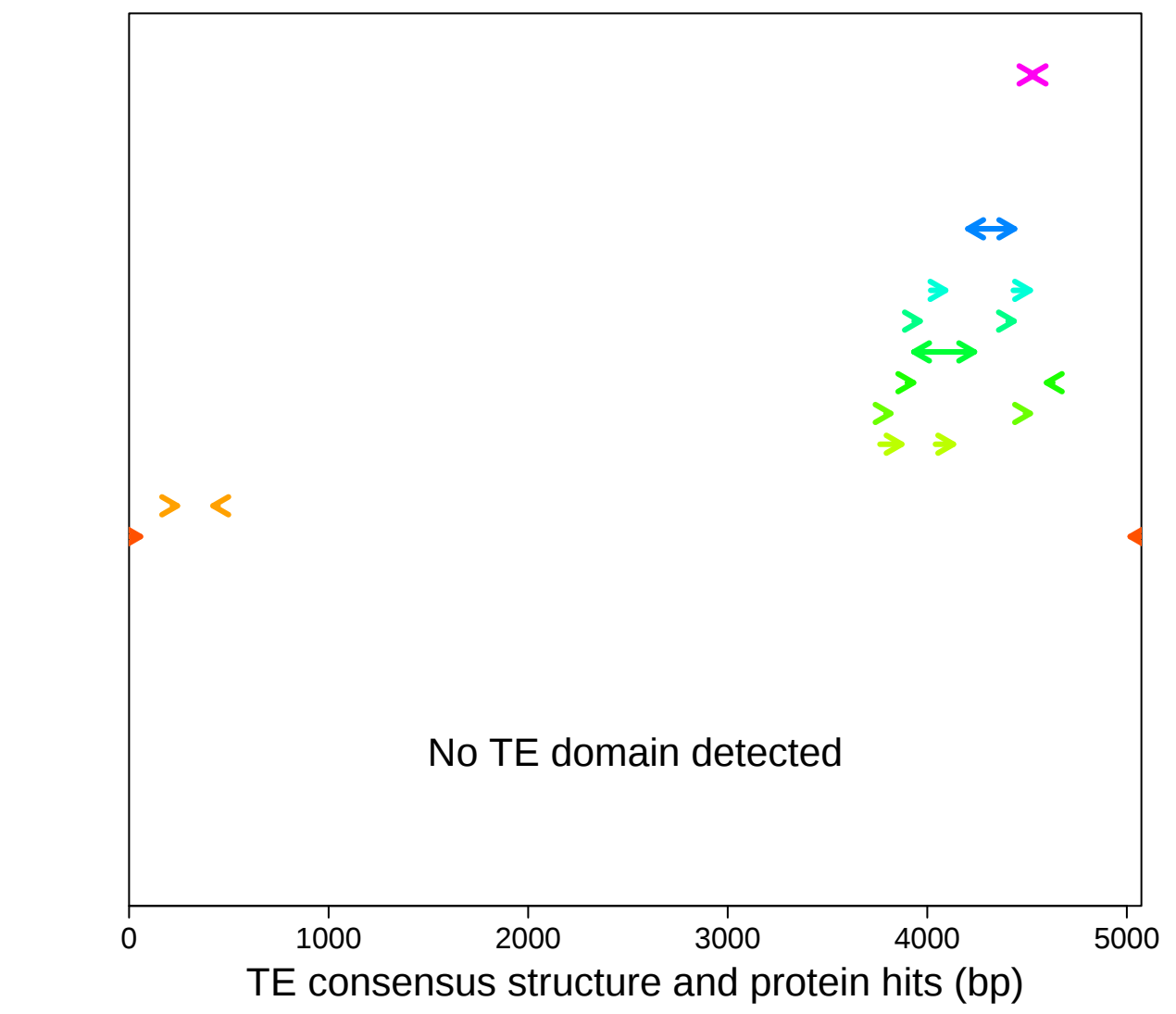
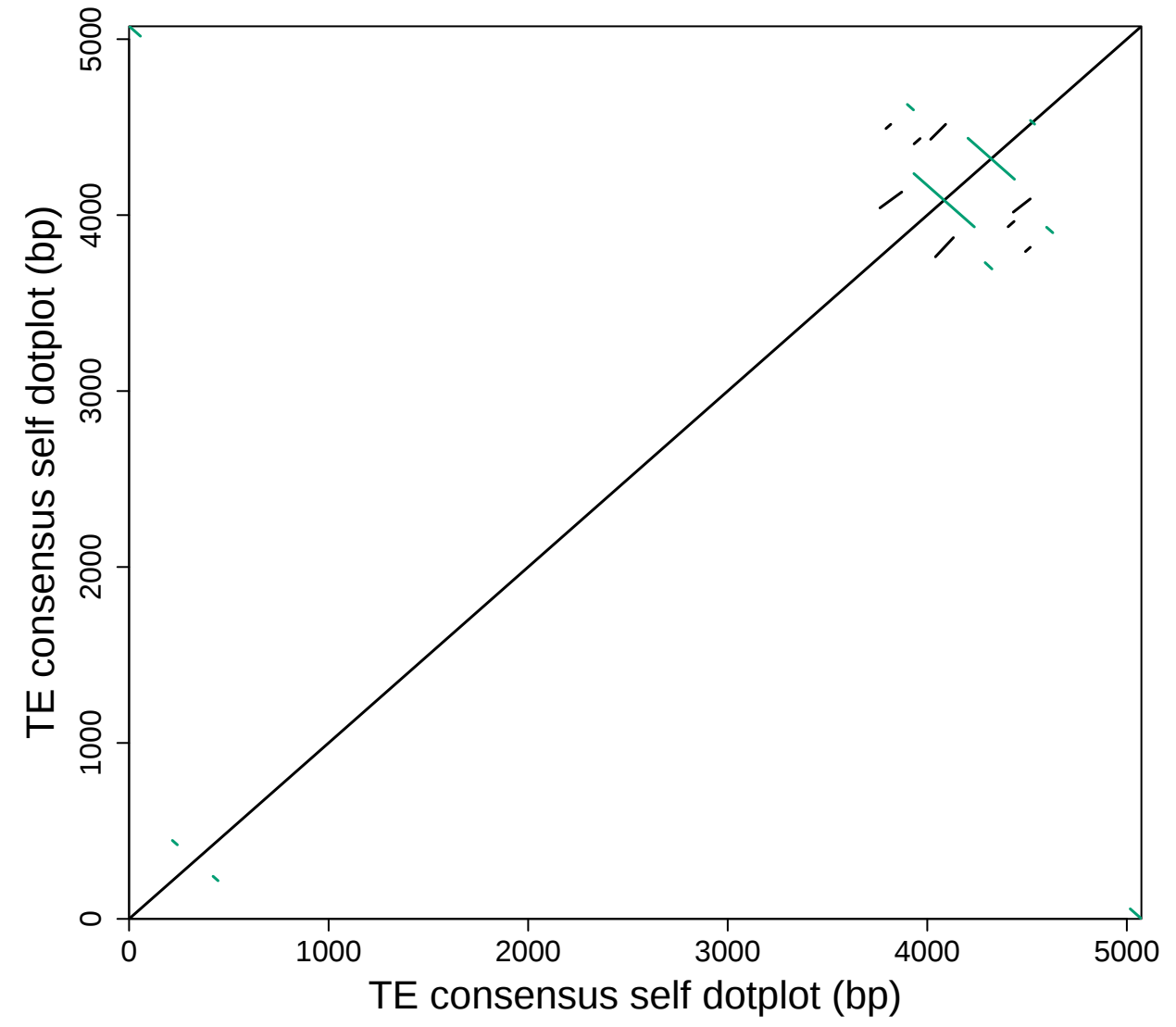
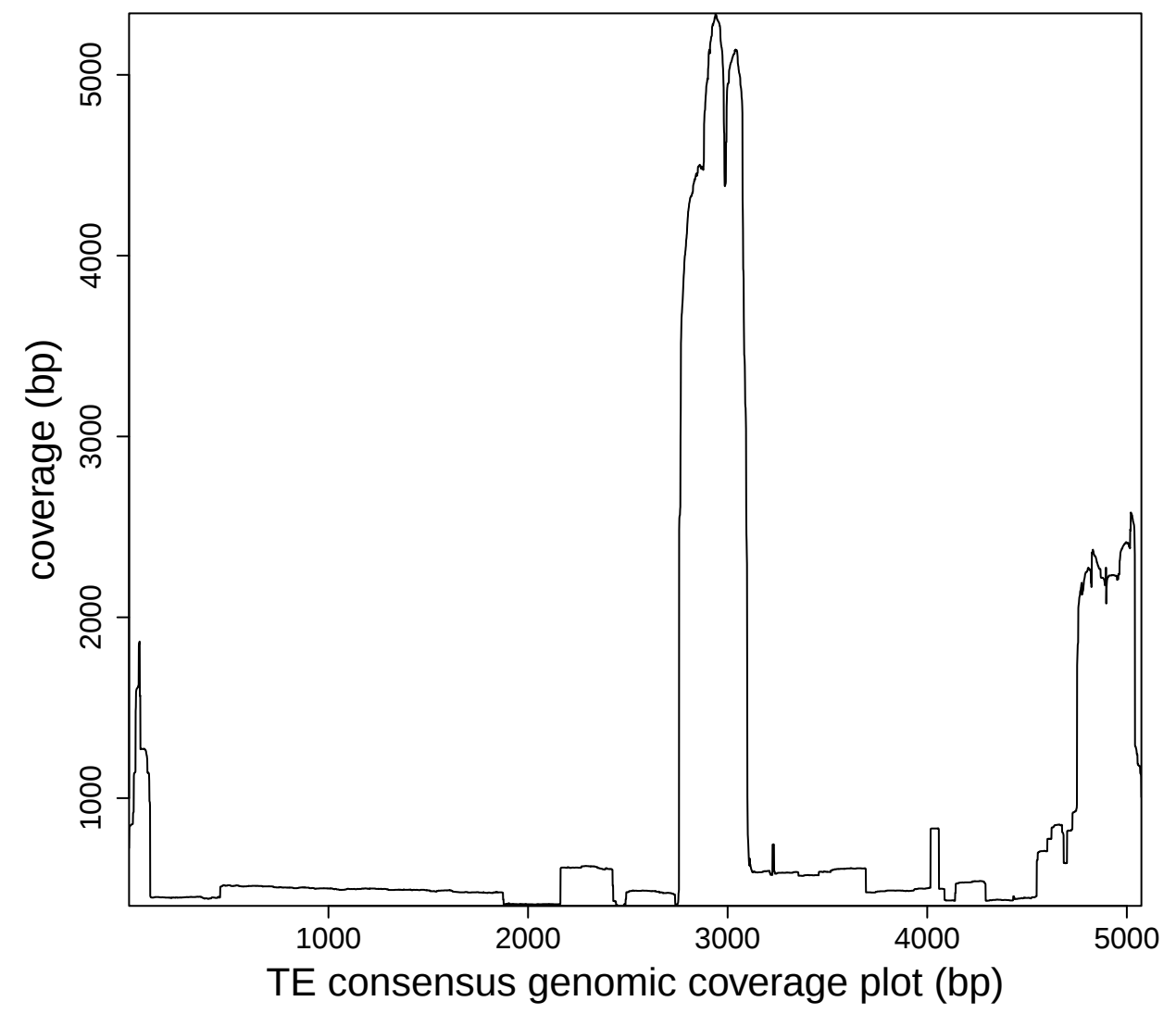
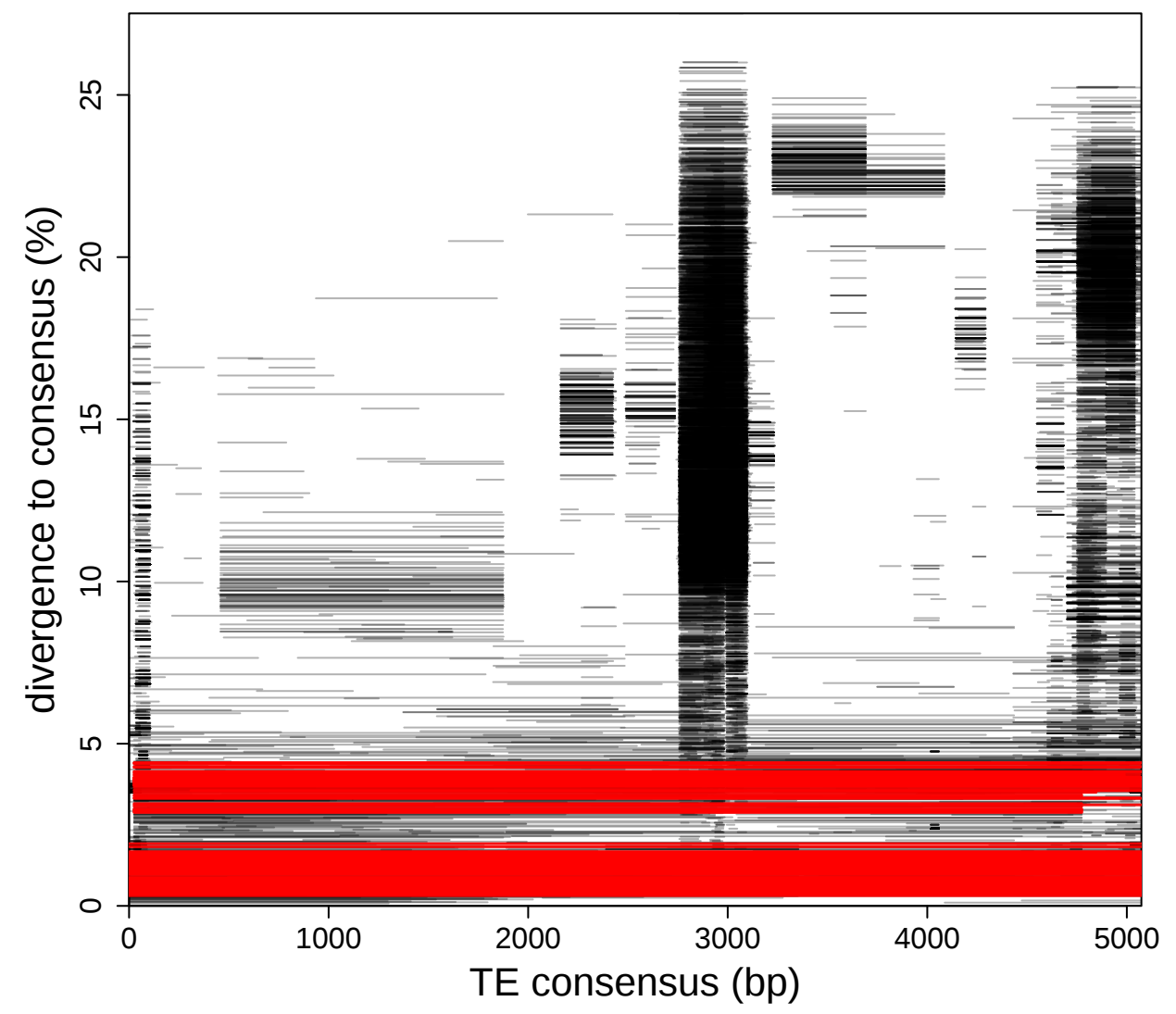






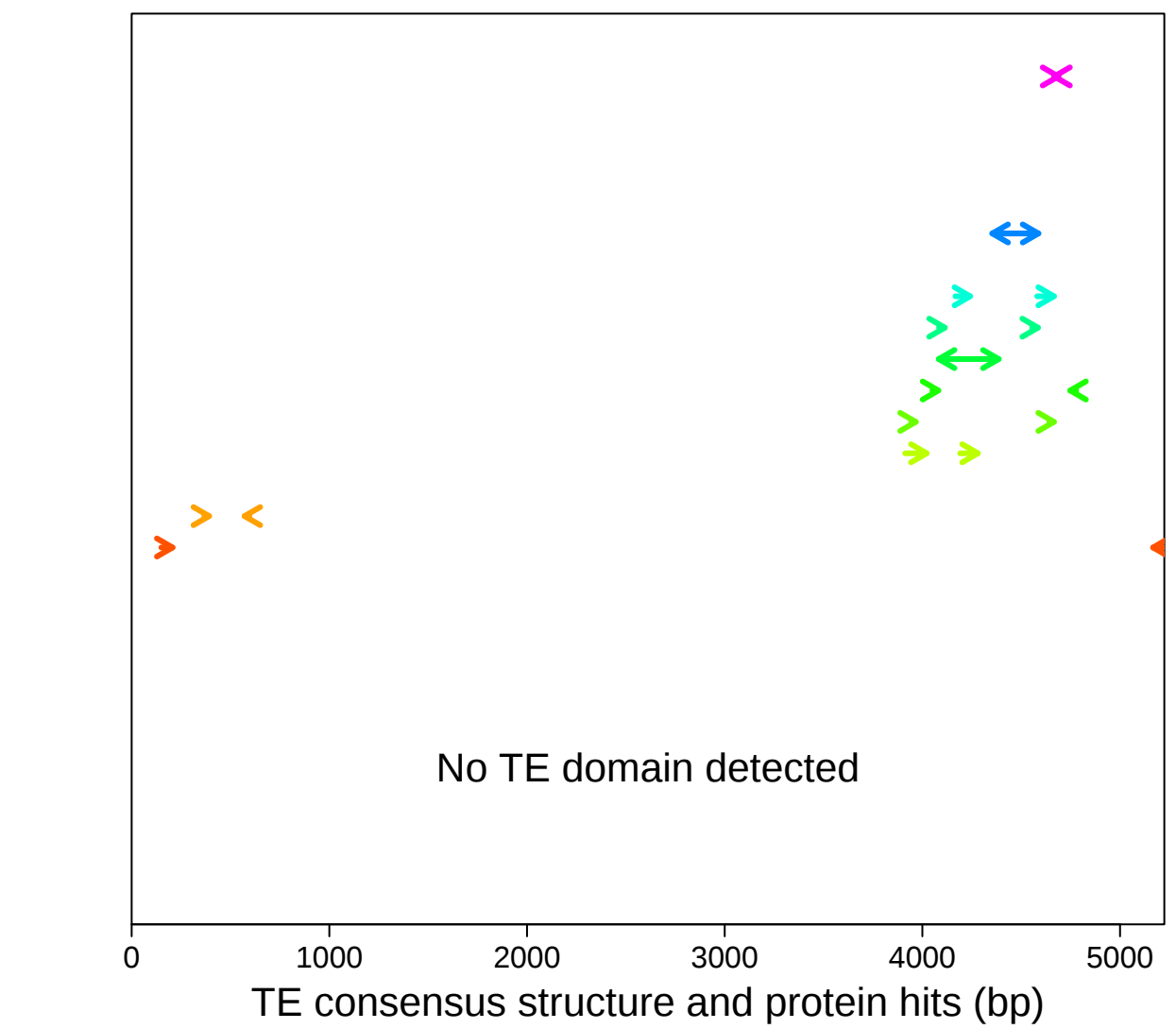
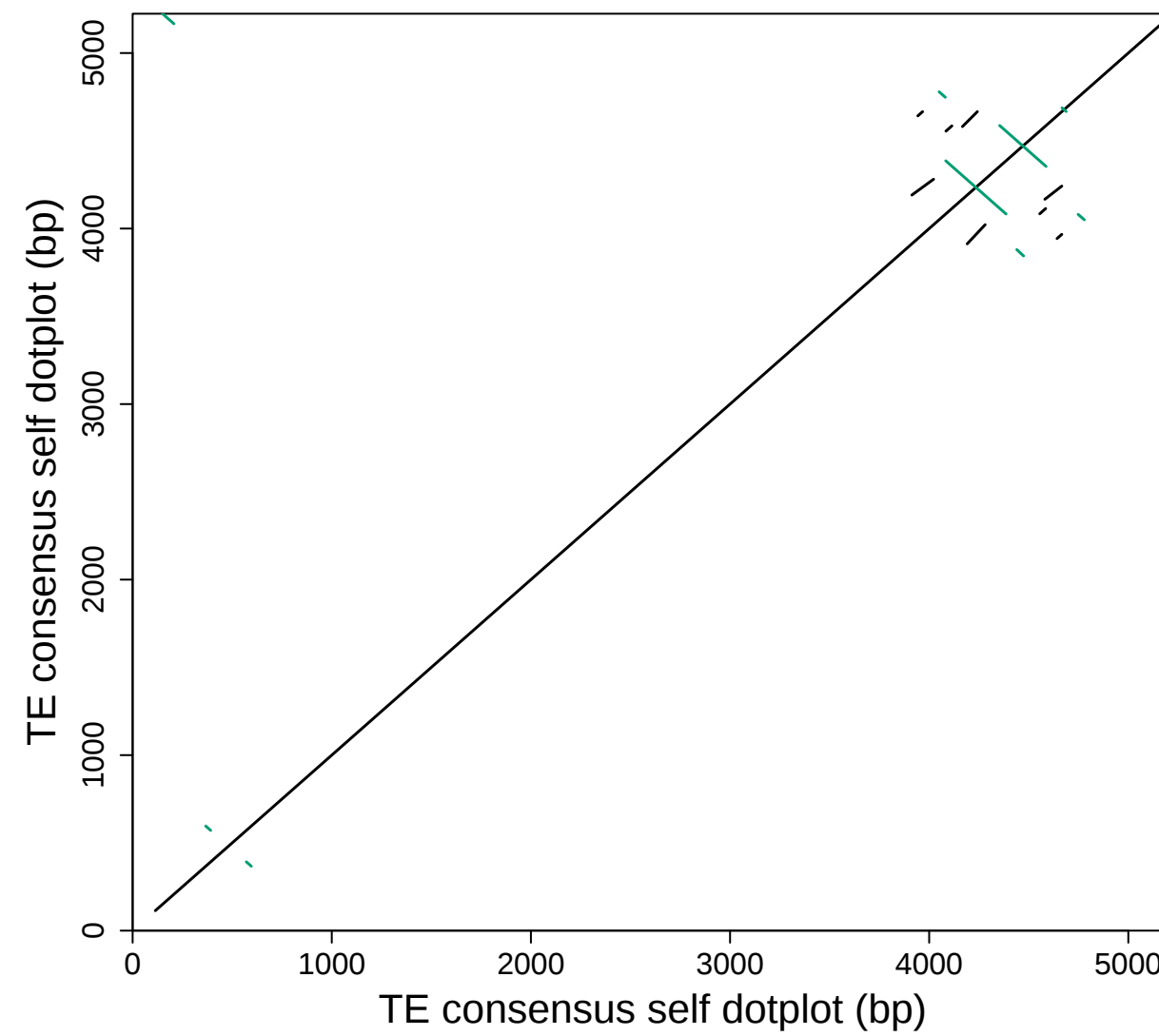
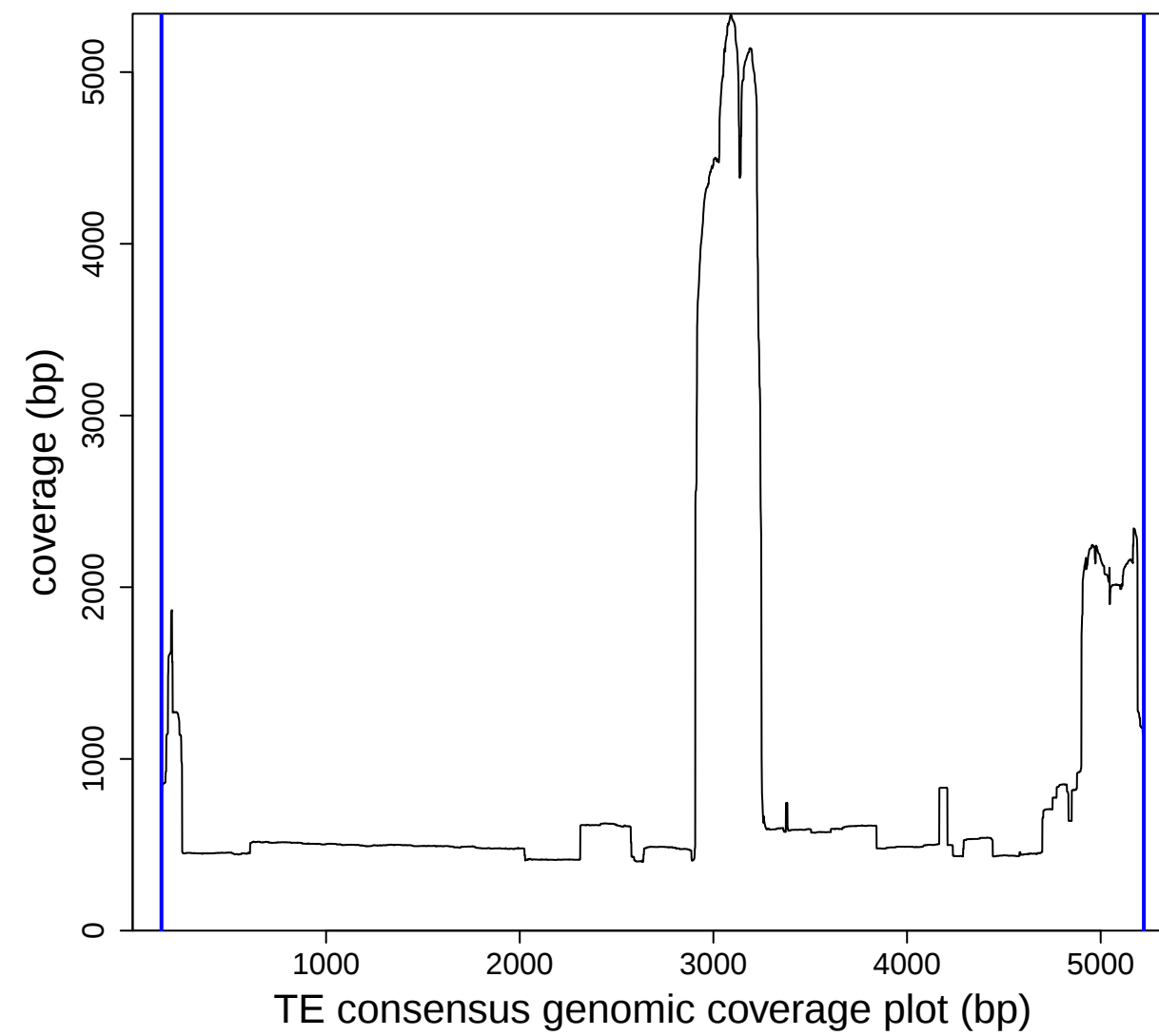
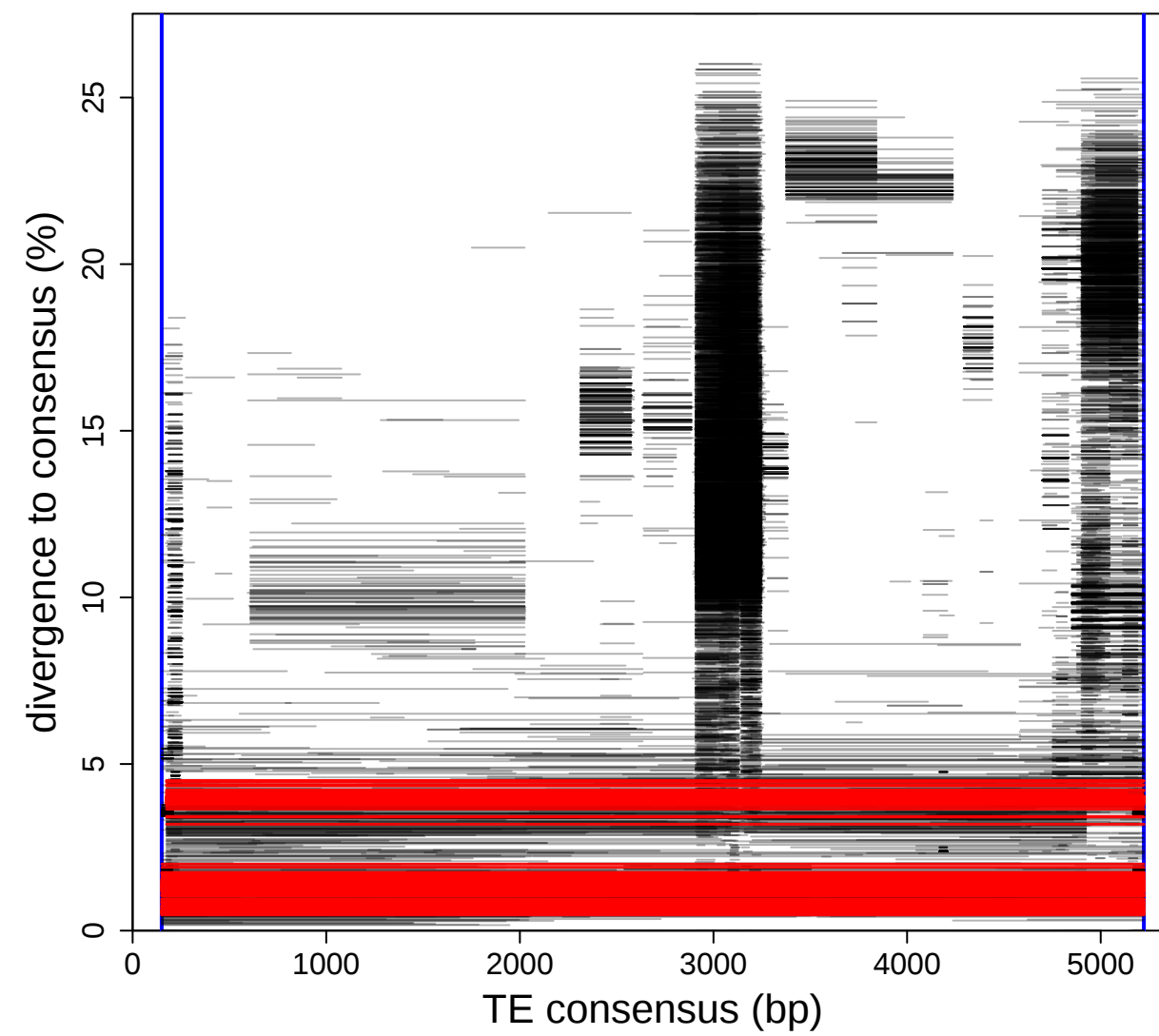
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 5073bp; fragments: 12823; full length: 200 (≥ 4565.7 bp)

After TEtrimmer 5073 bp



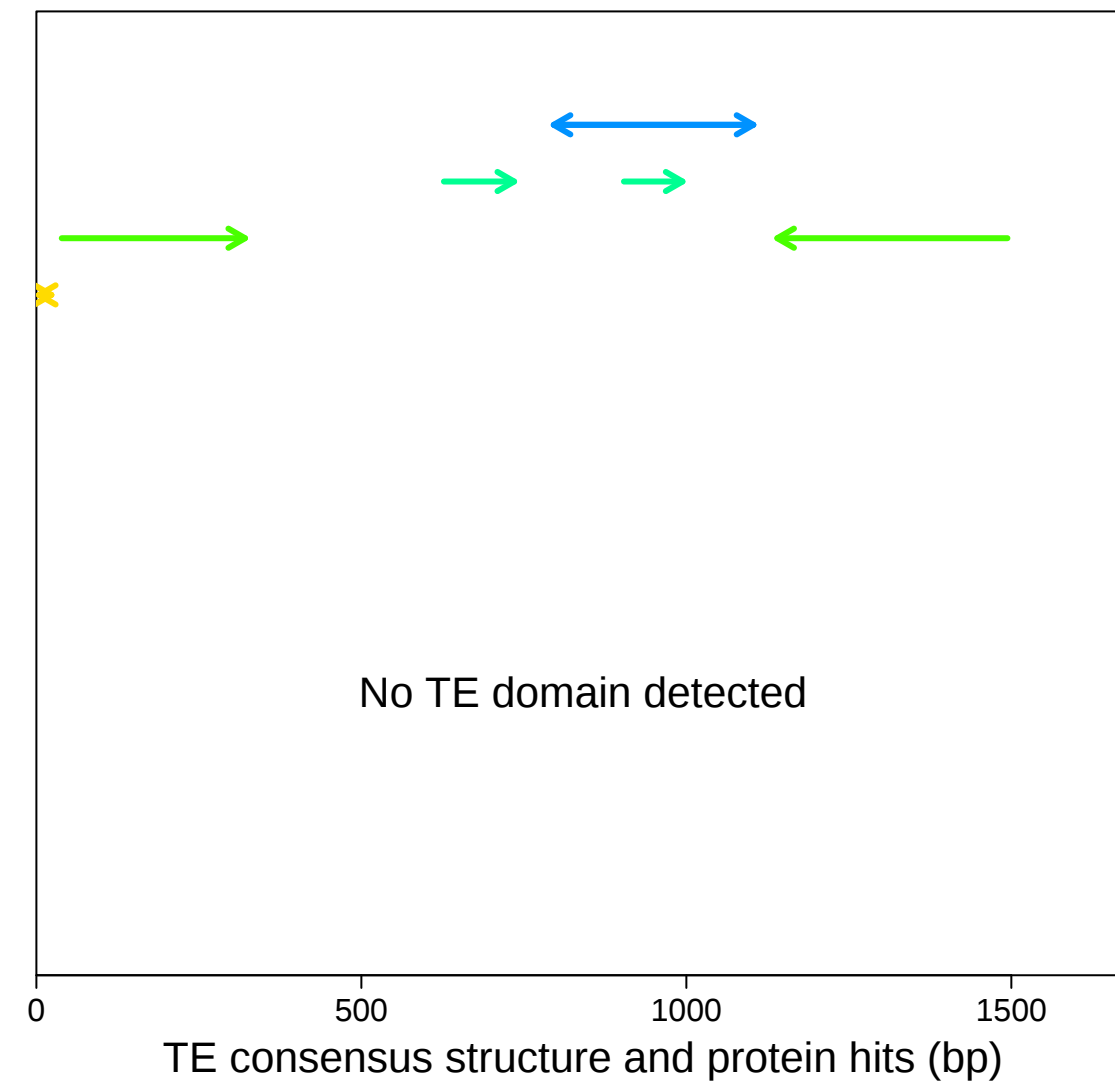
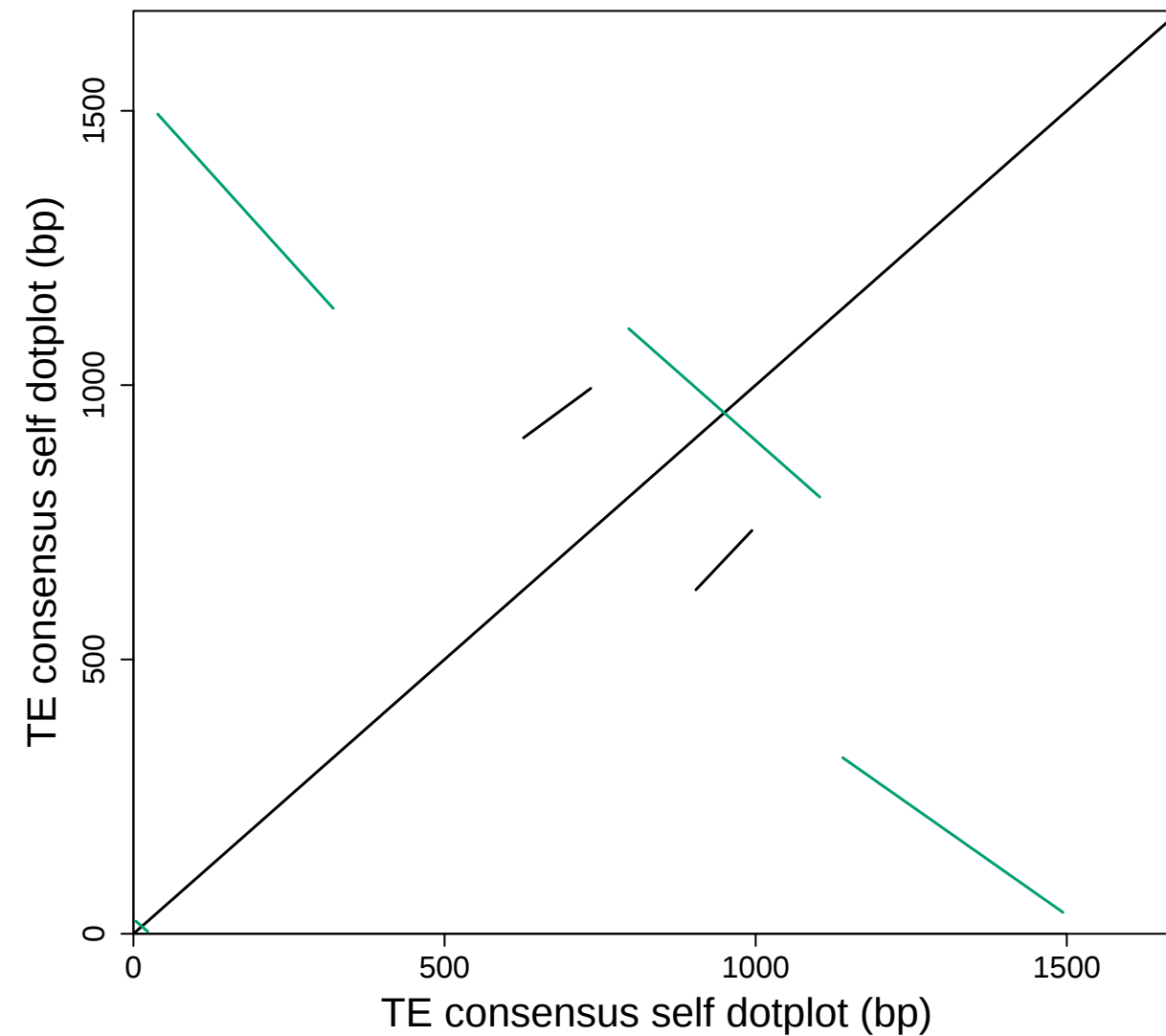
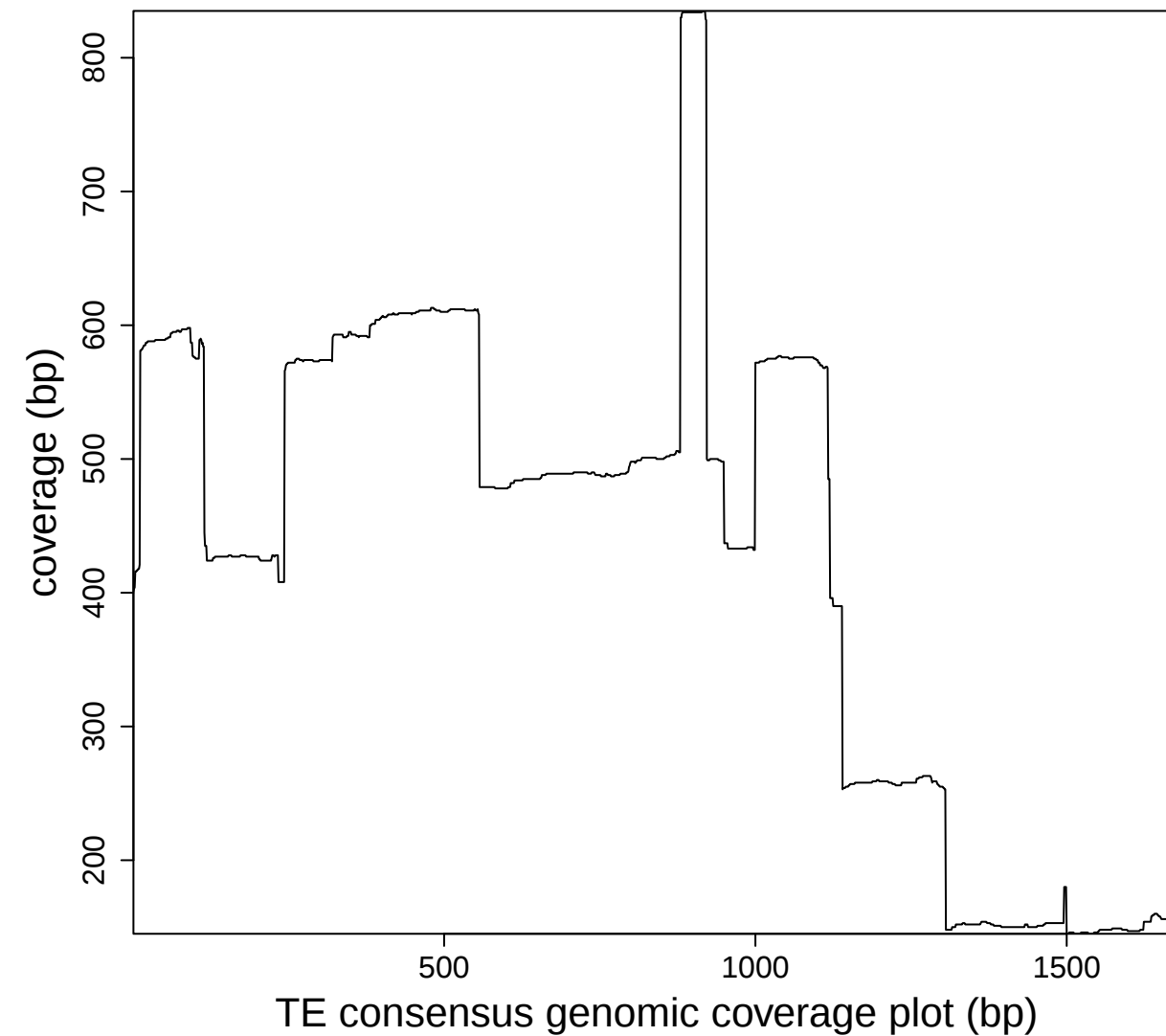
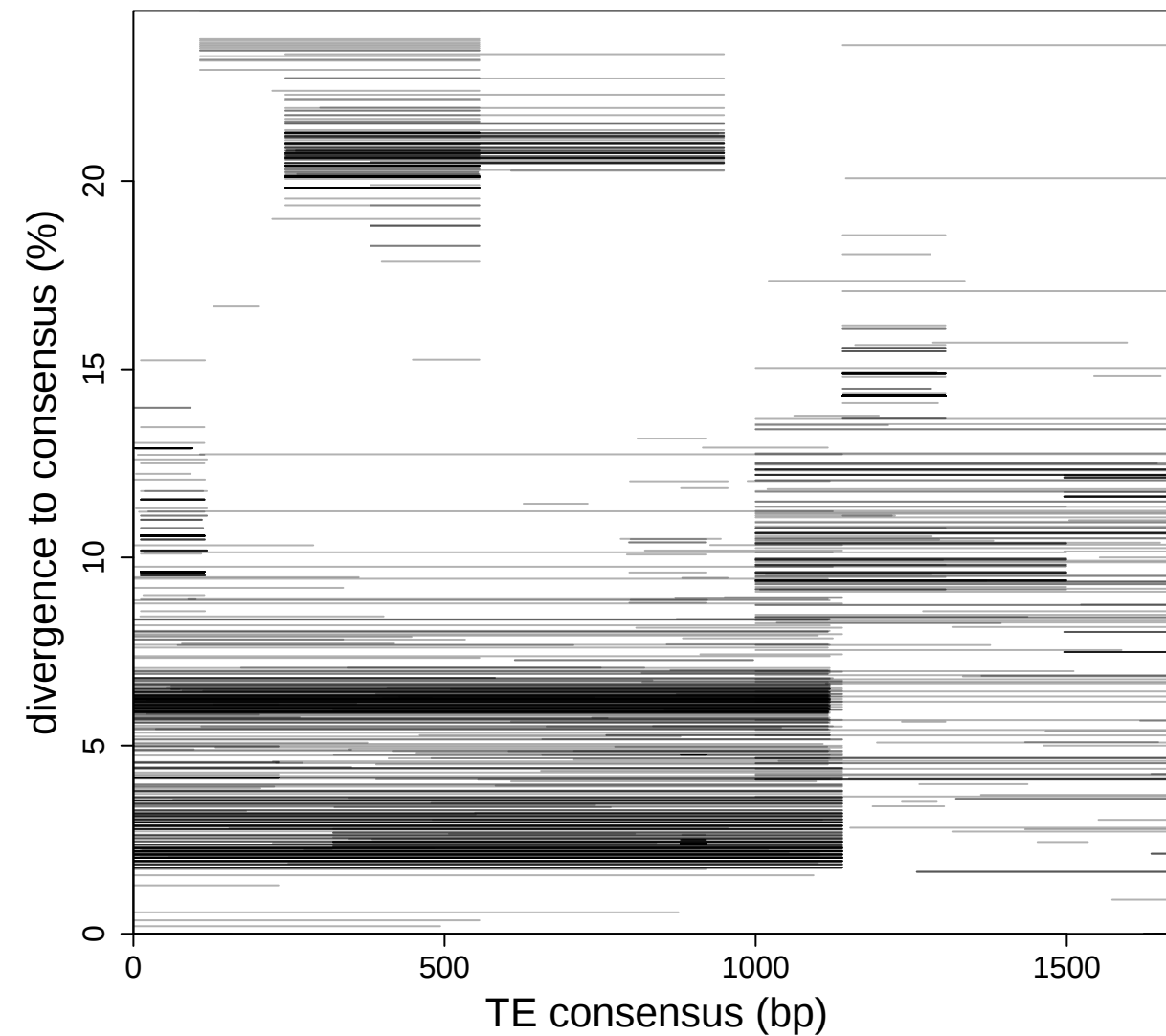
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 5373bp; fragments: 12586; full length: 178 (>=4835.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

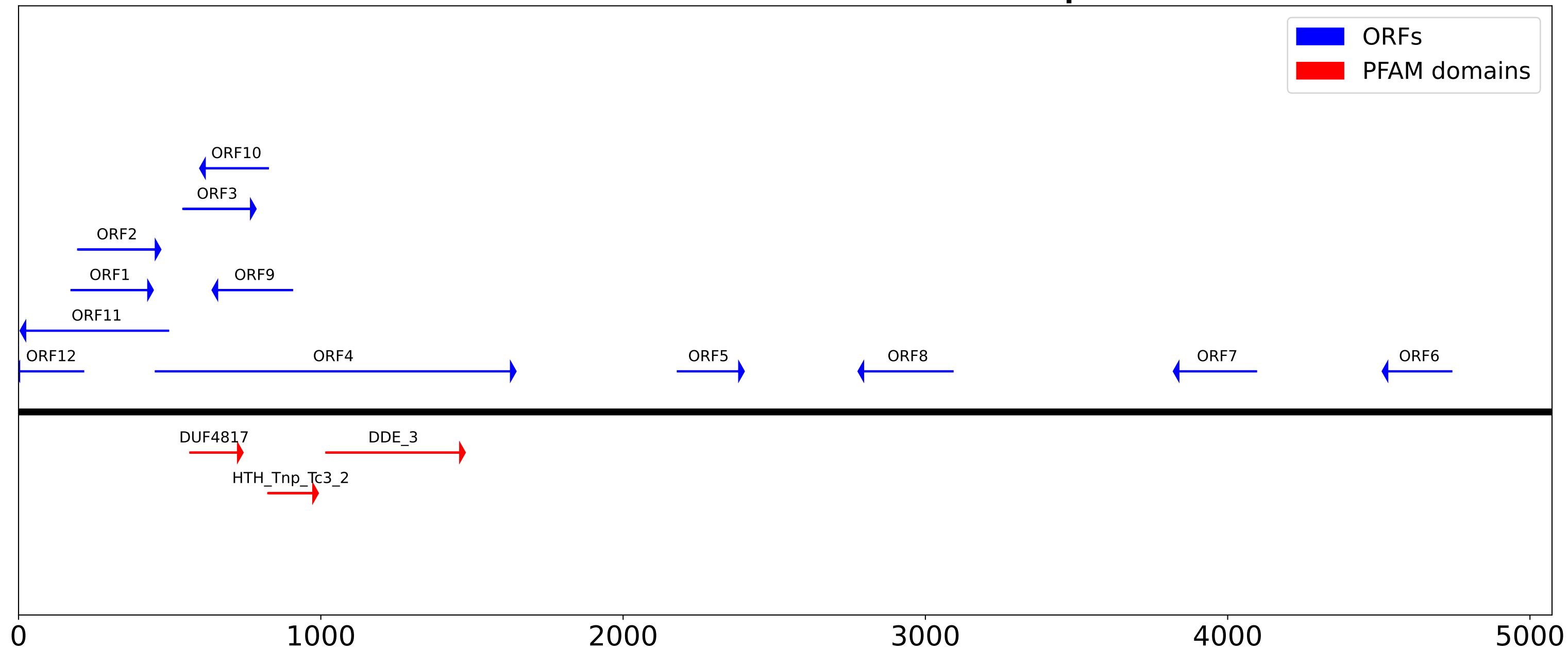


TE: rnd_1_family_775
size: 1682bp; fragments: 1550; full length: 0 (≥ 1513.8 bp)

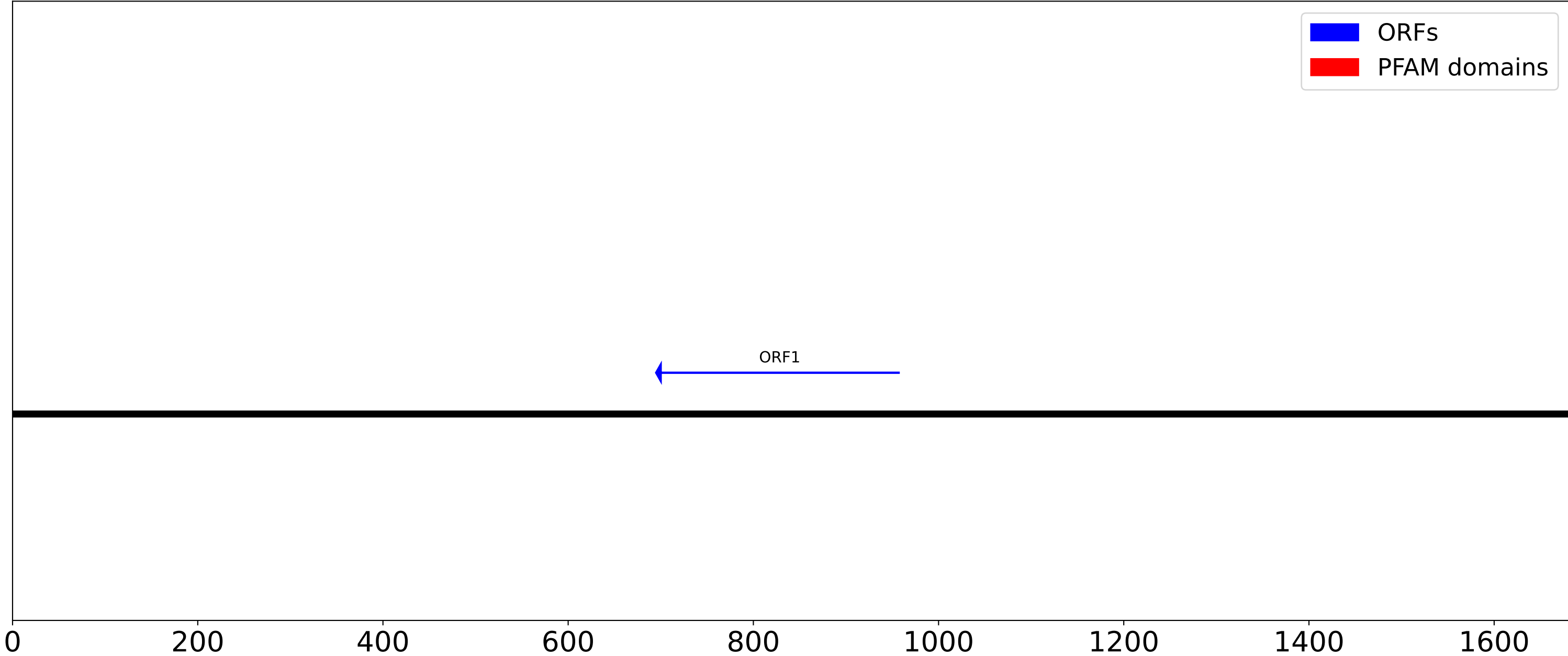
Before TEtrimmer 1682 bp



After TEtrimmer ORF and PFAM domain plot



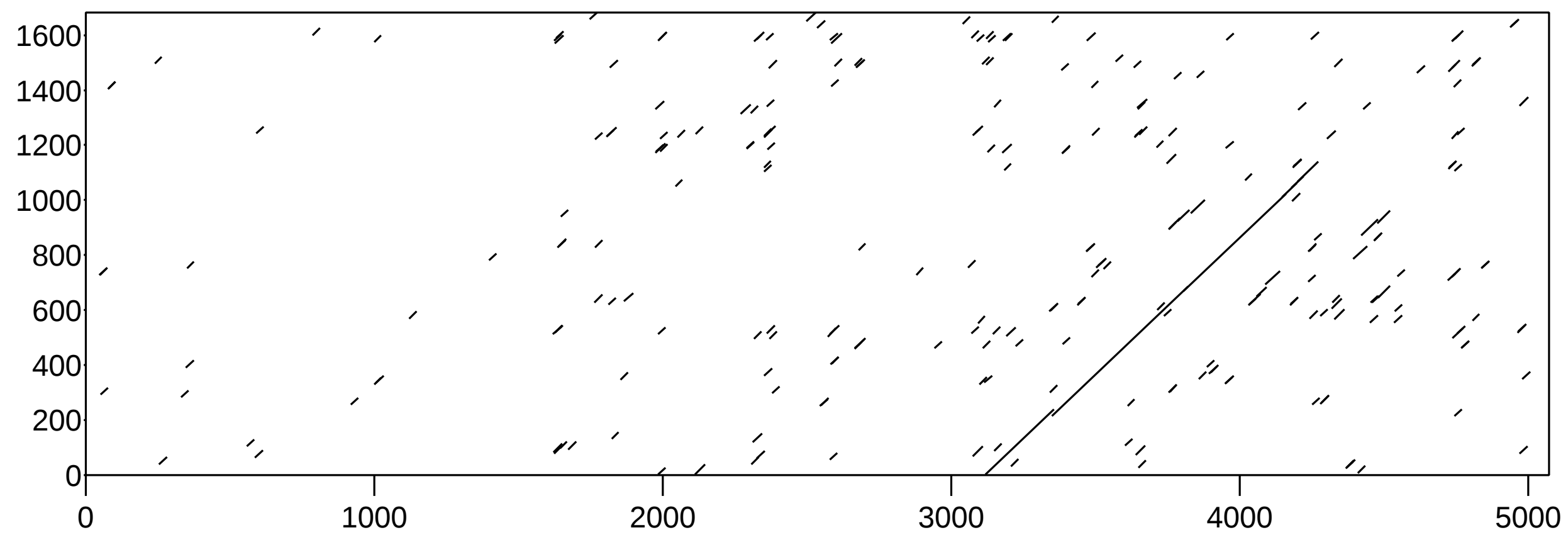
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)