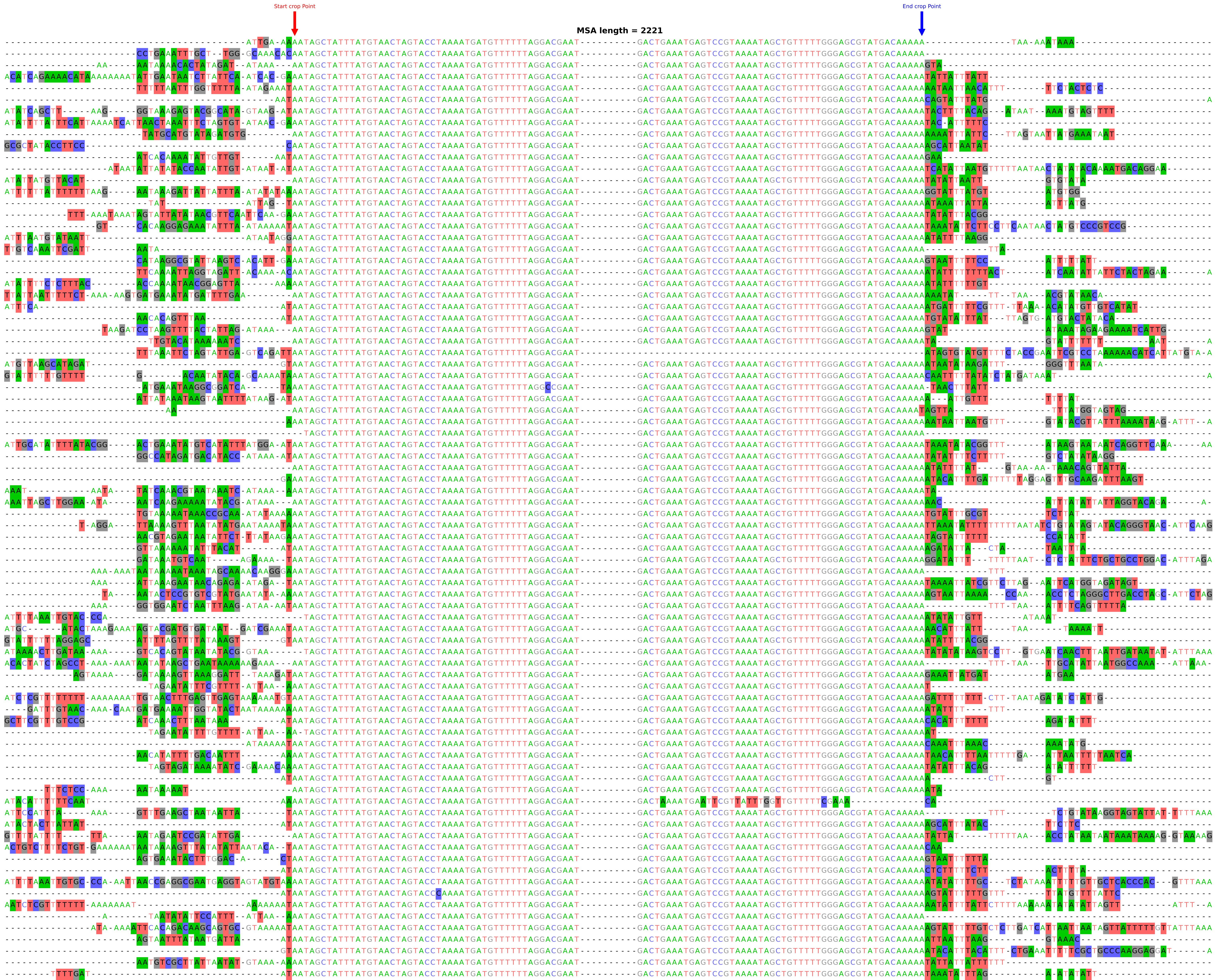
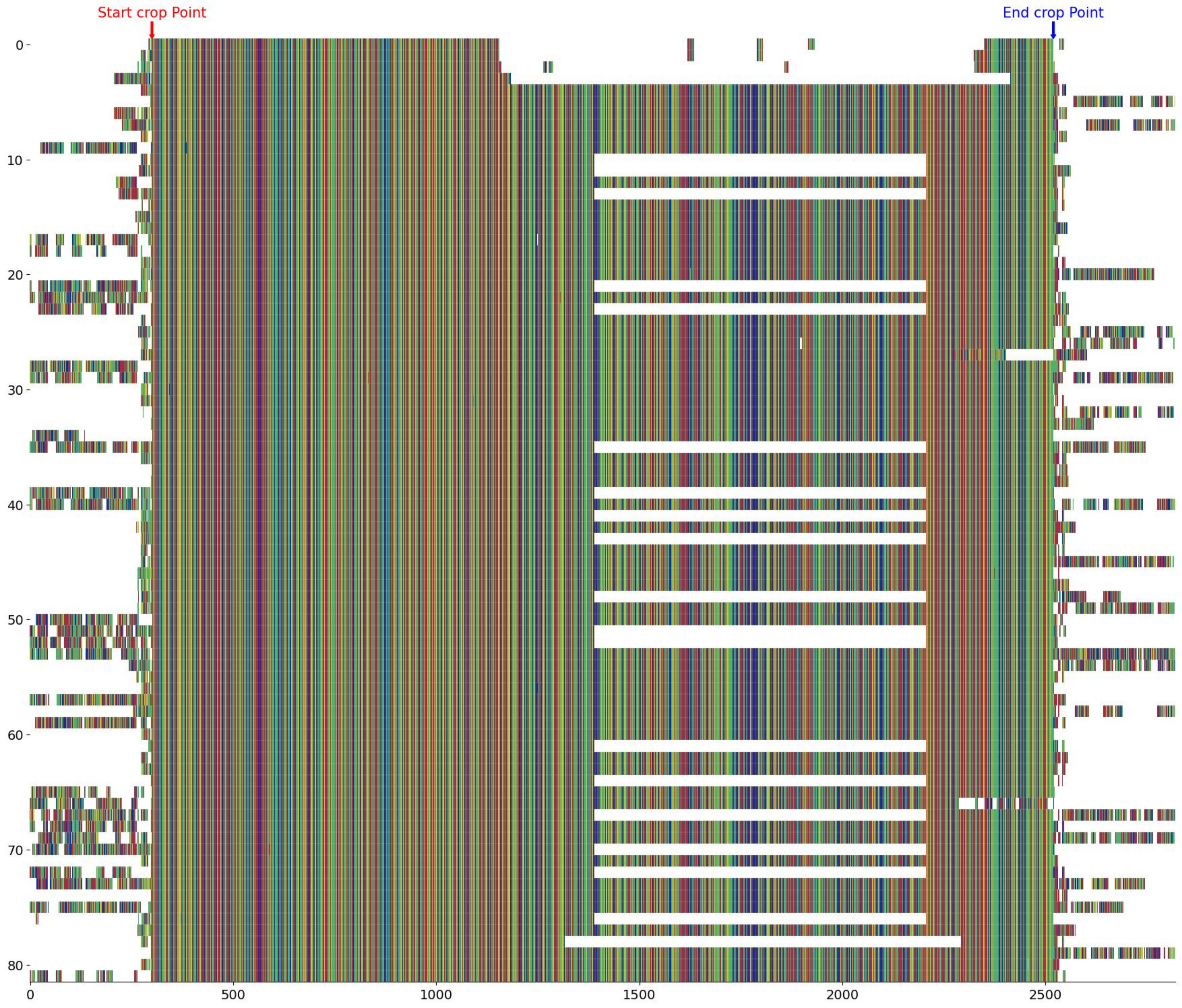
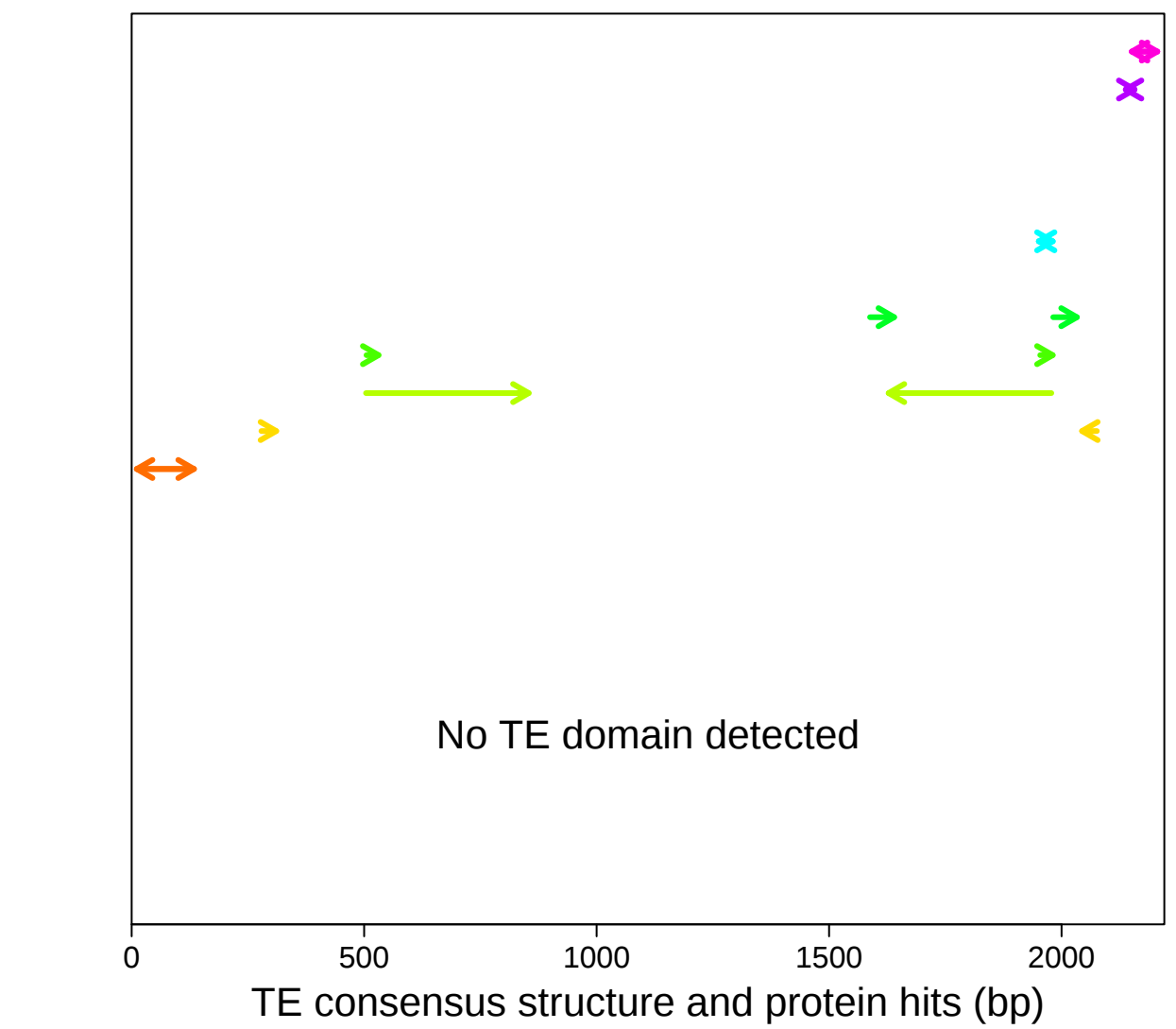
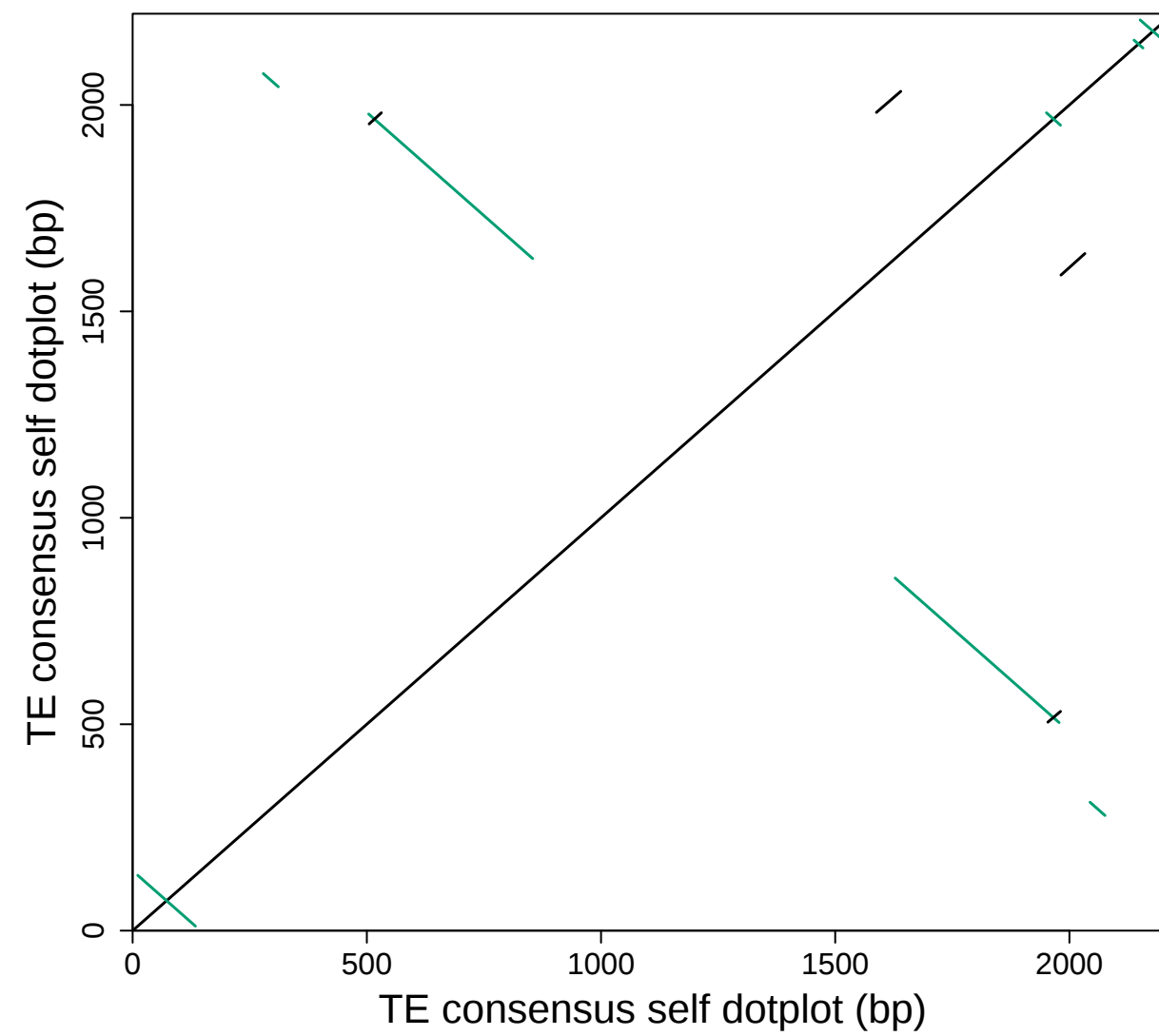
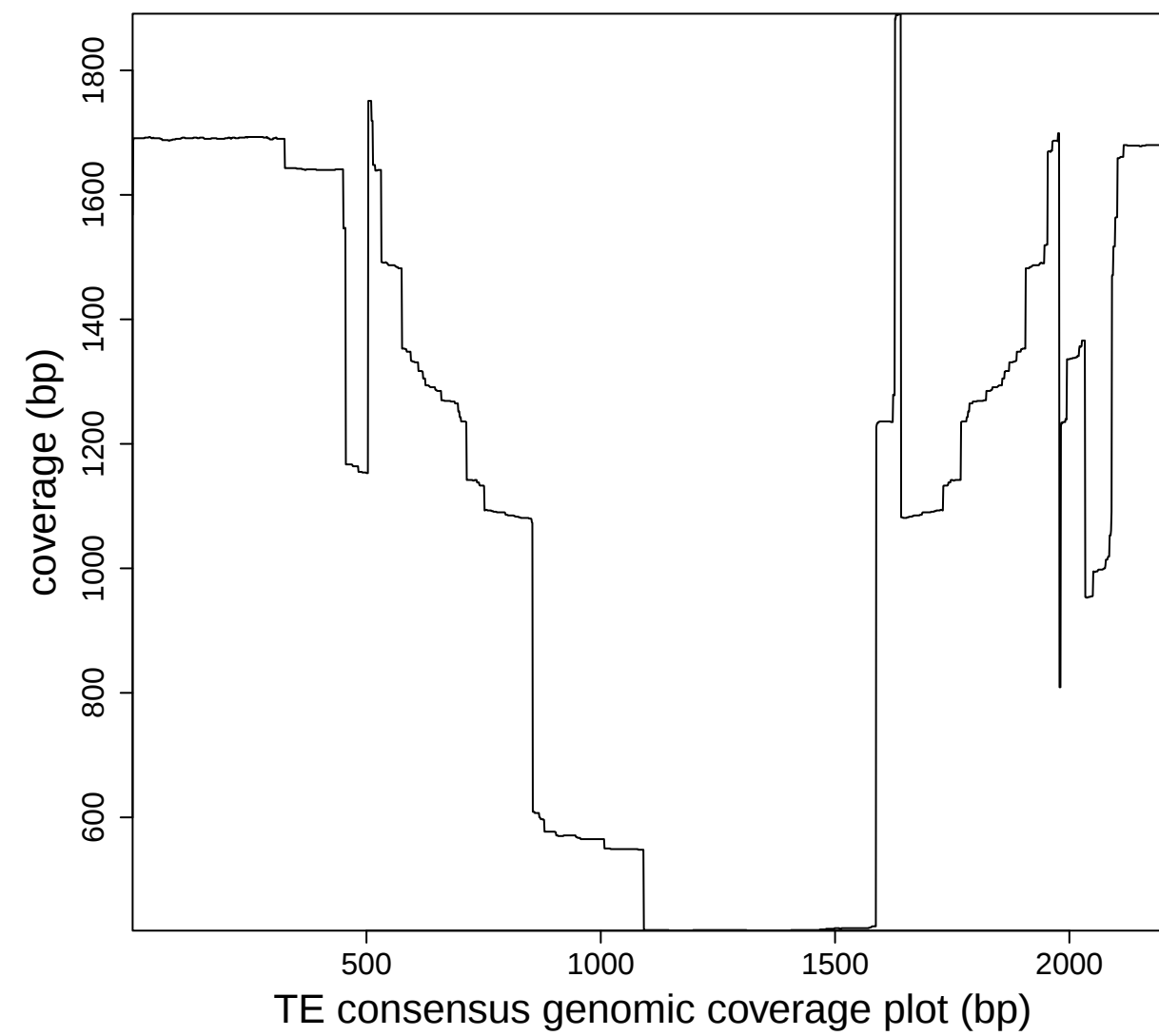
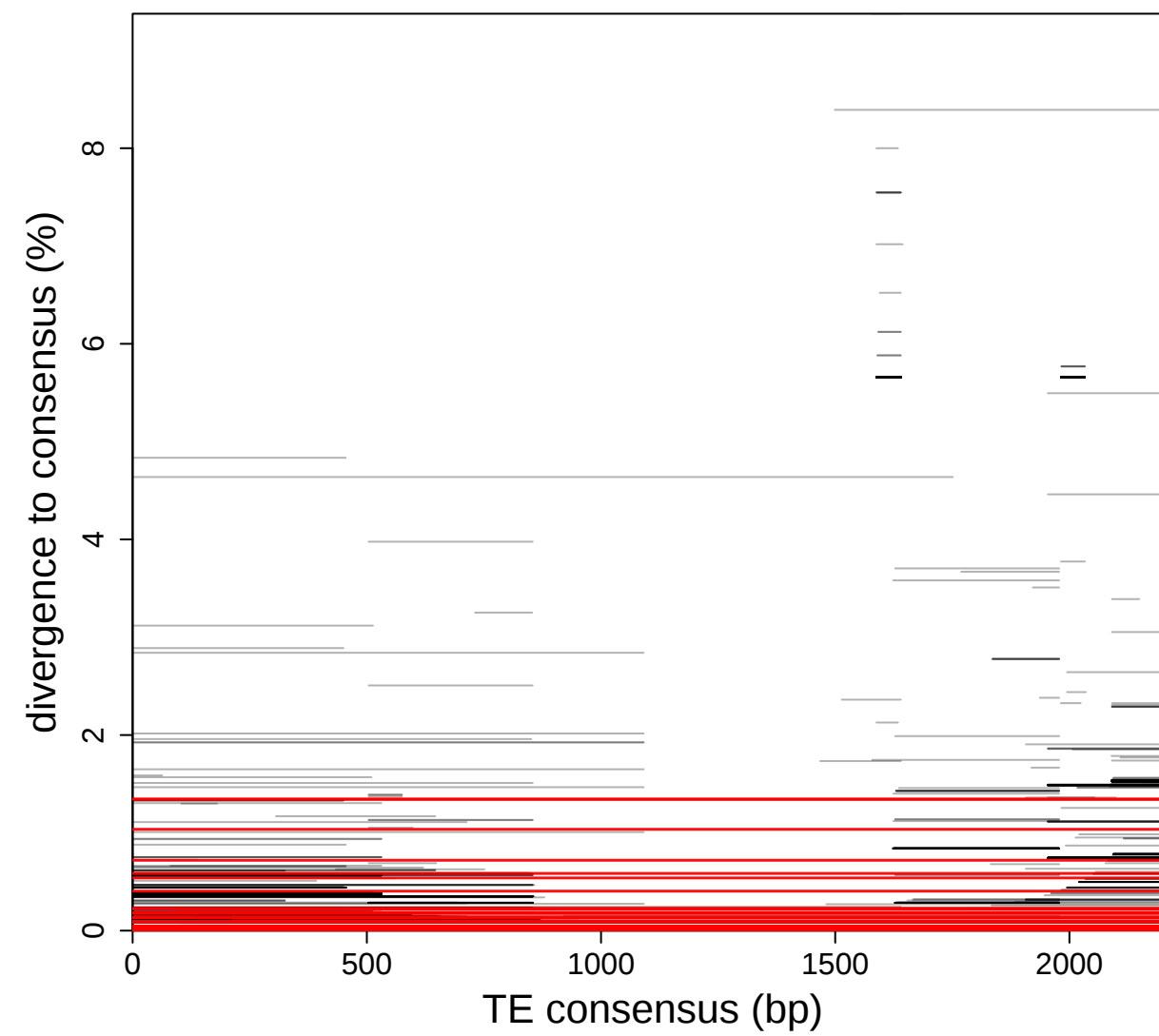






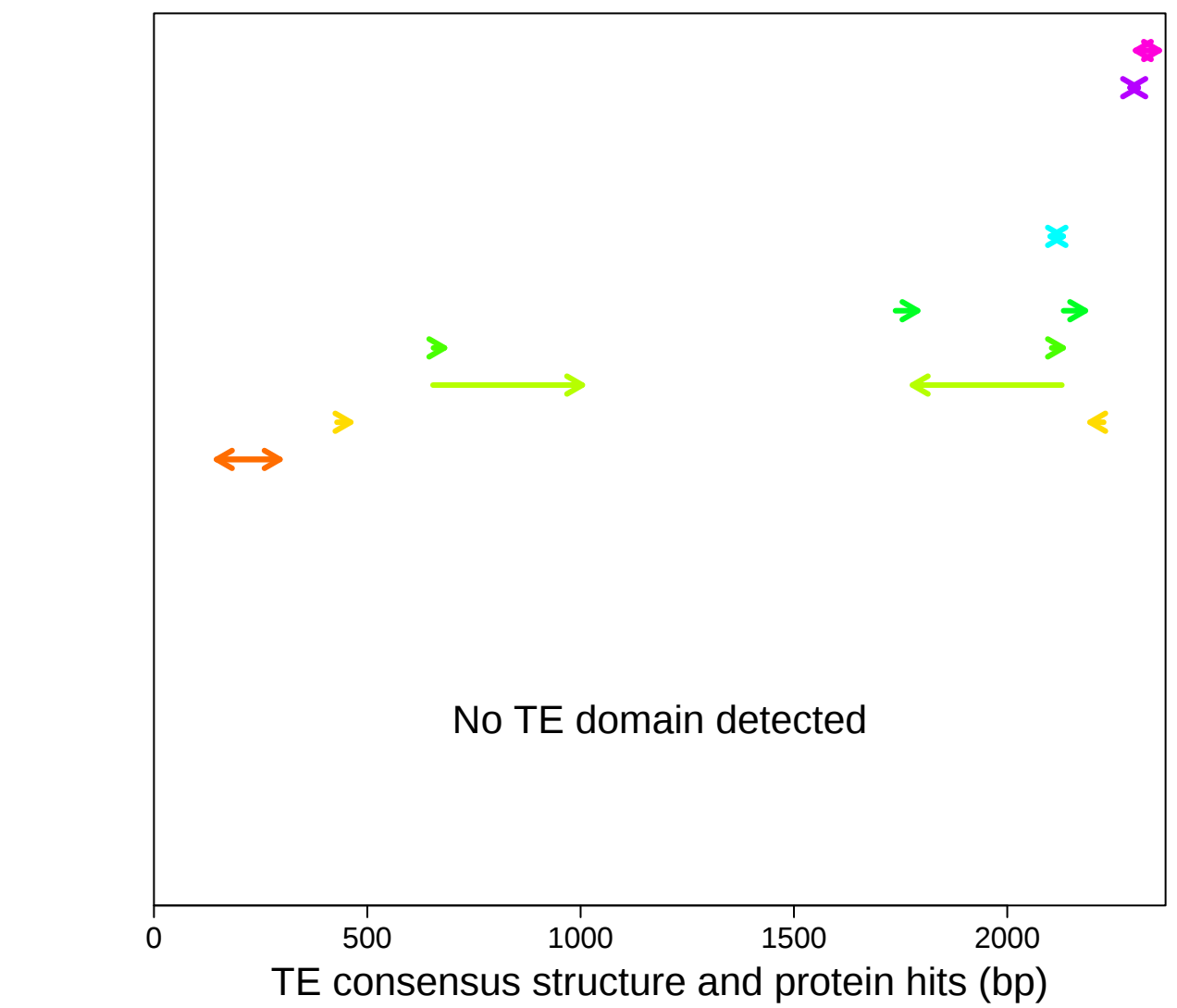
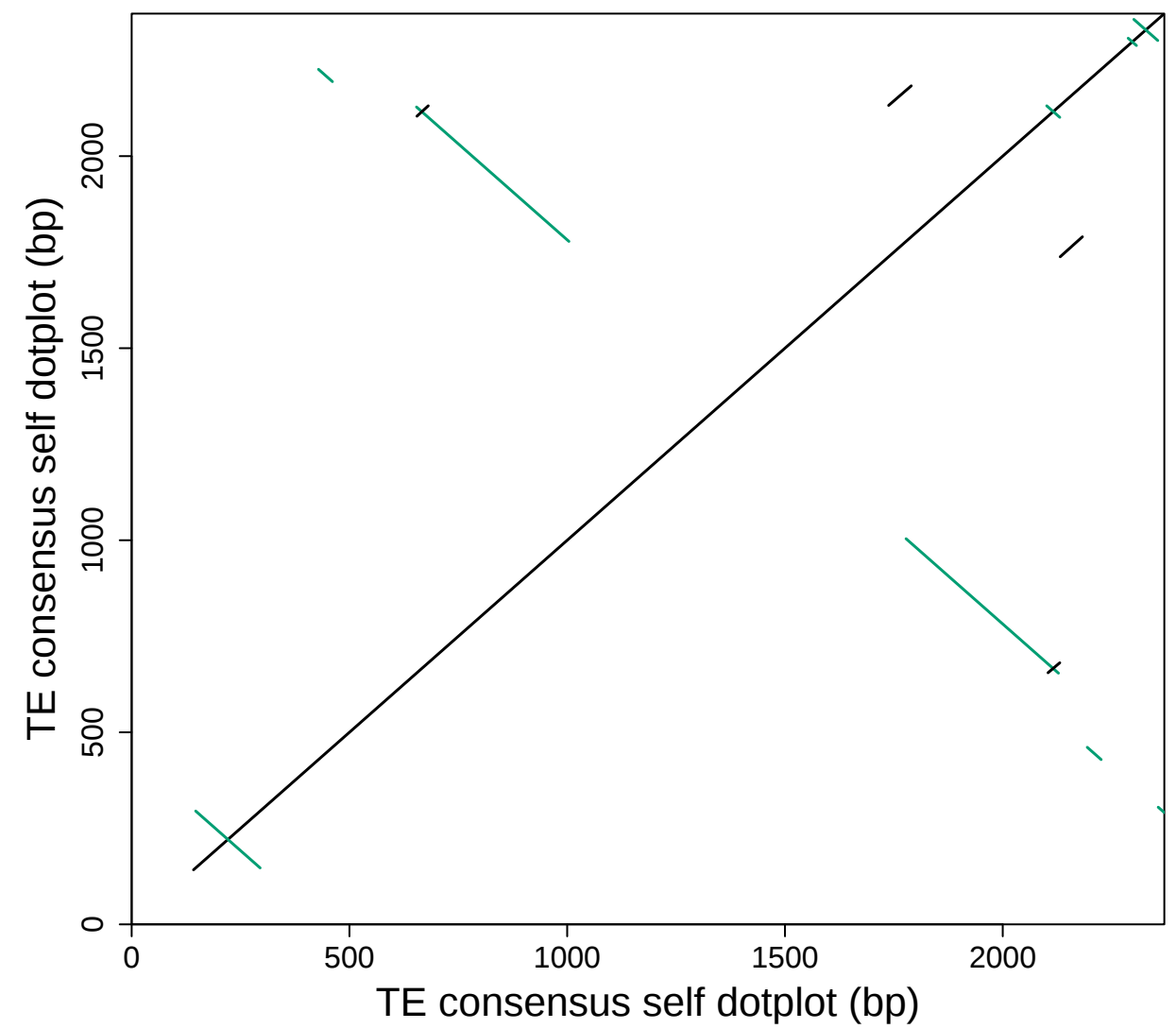
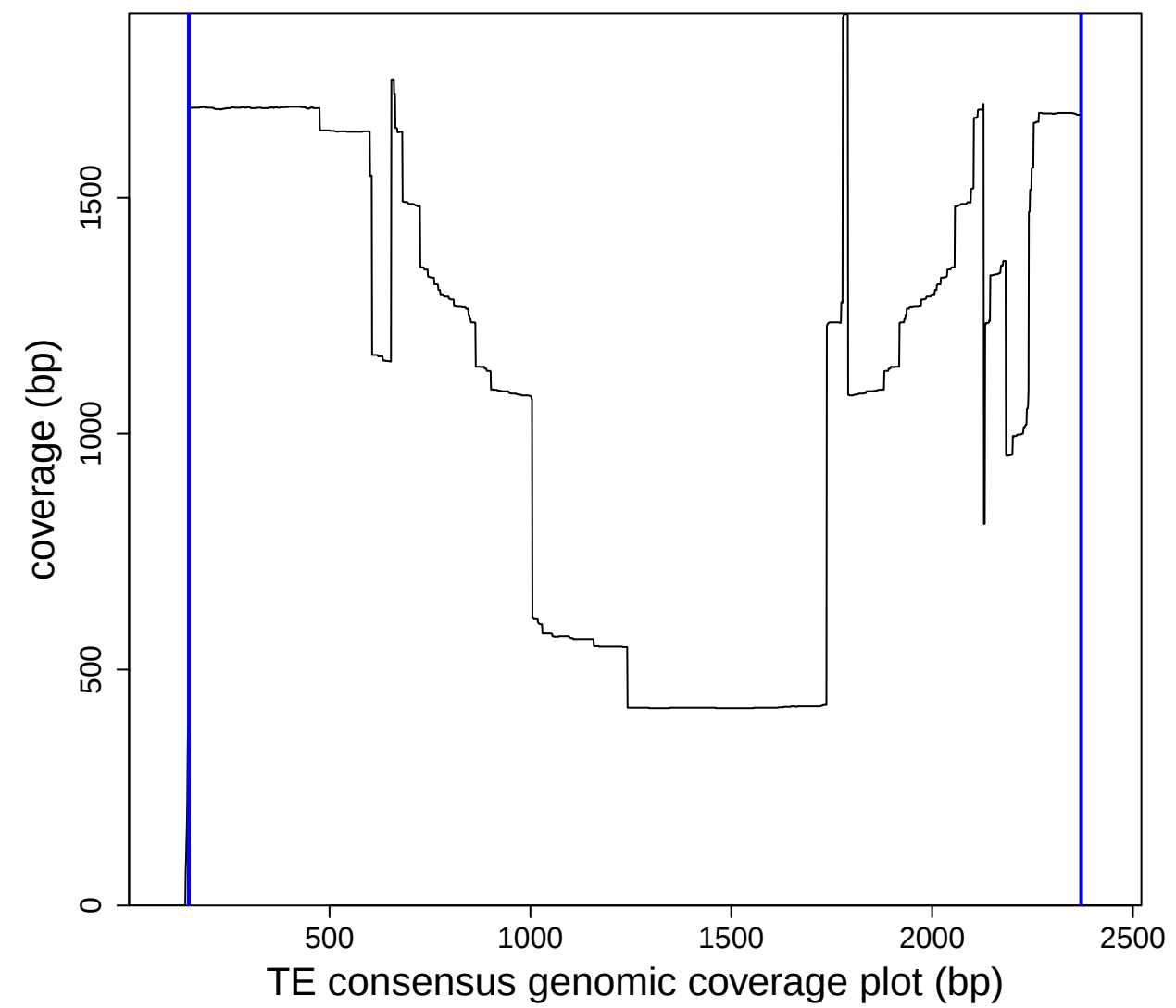
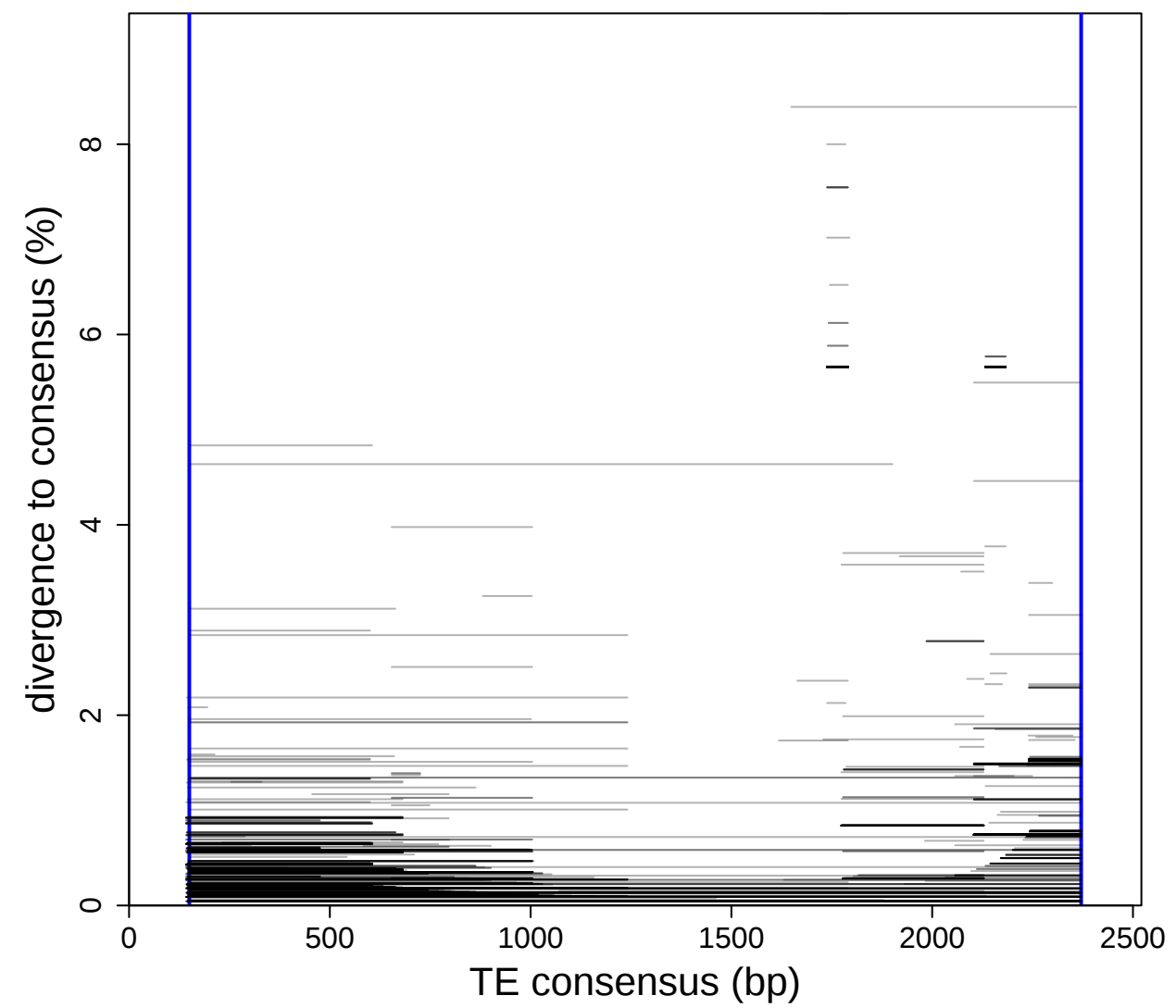
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 2221bp; fragments: 5726; full length: 399 (>=1998.9bp)

After TEtrimmer 2221 bp

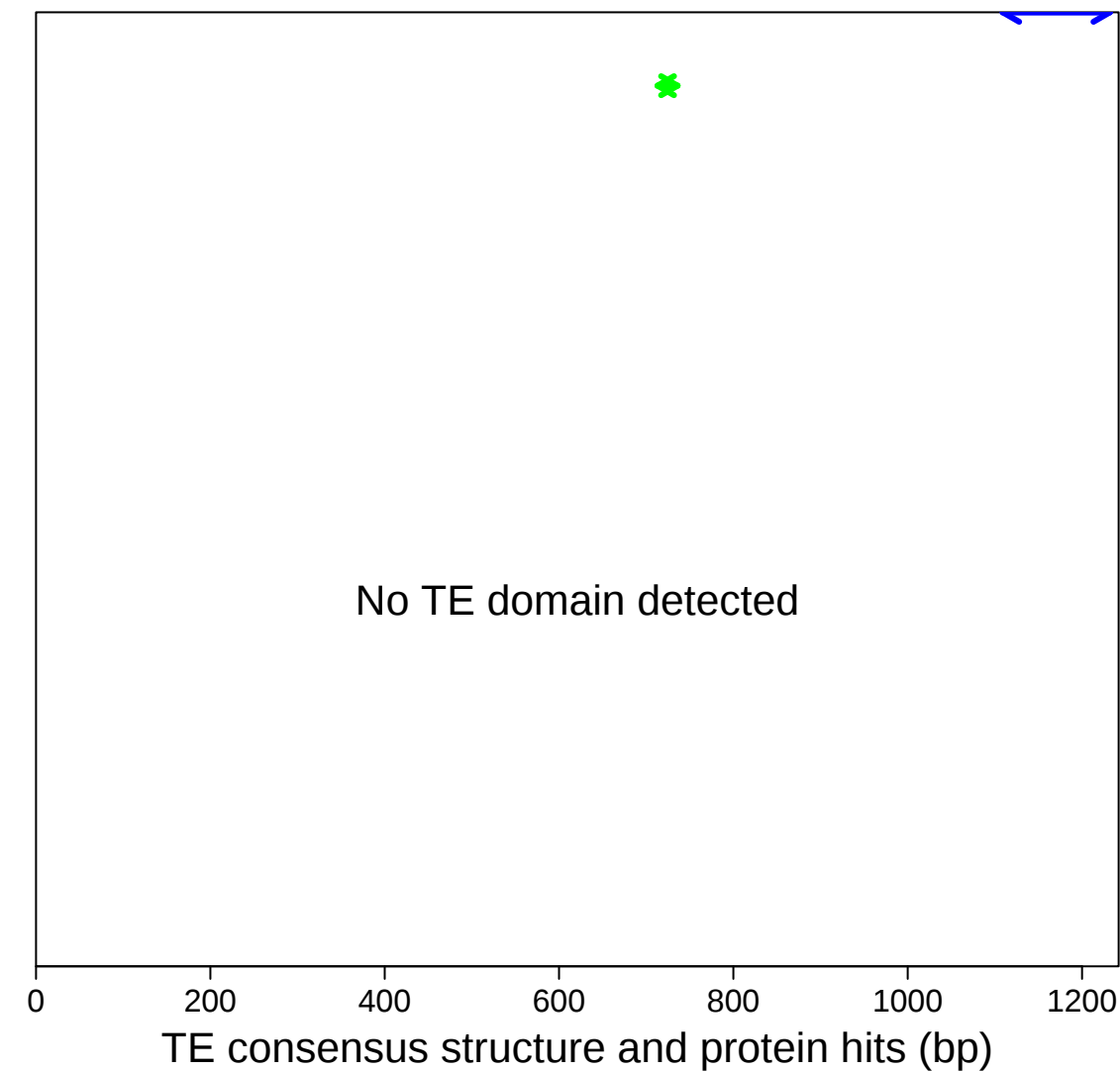
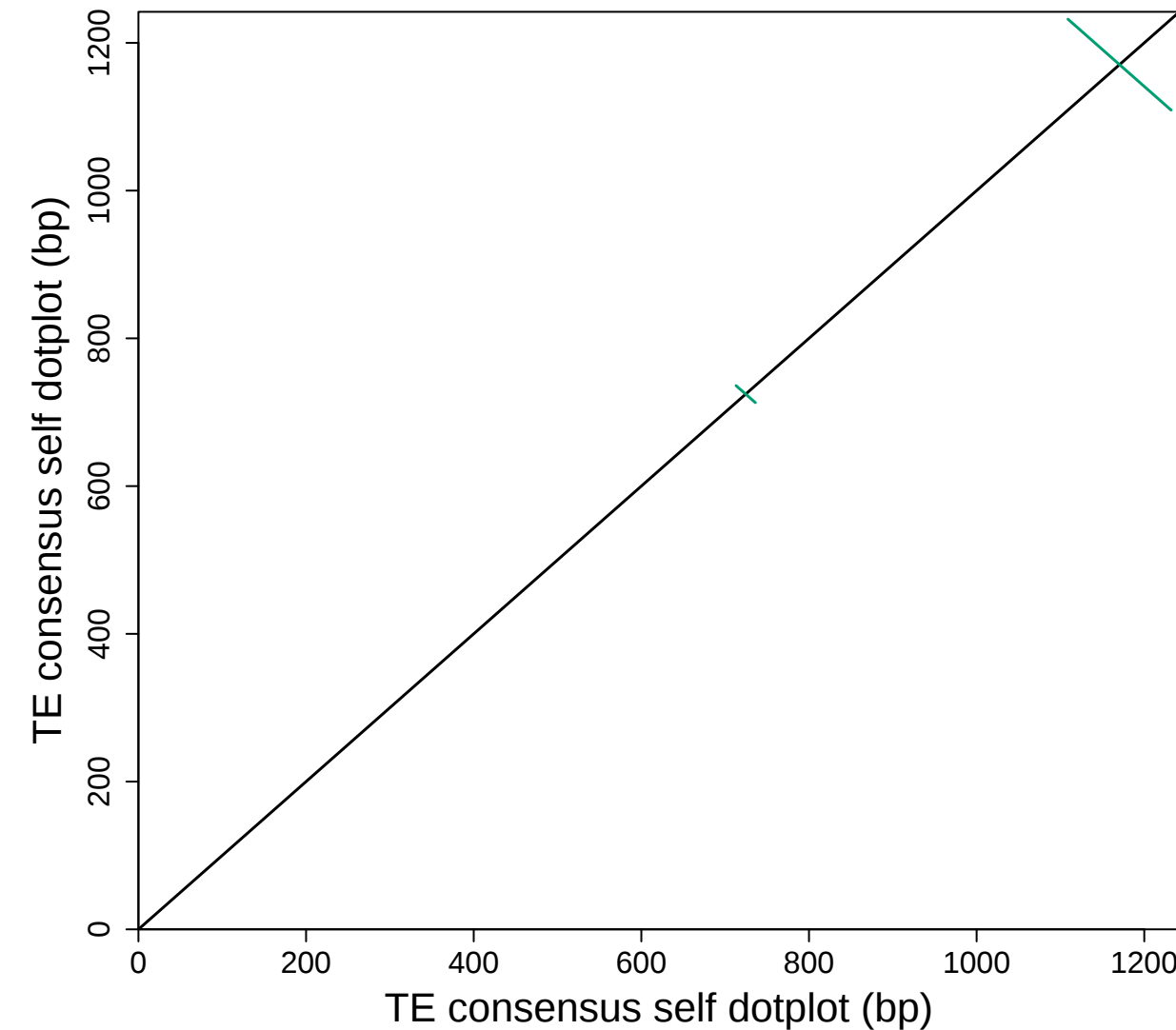
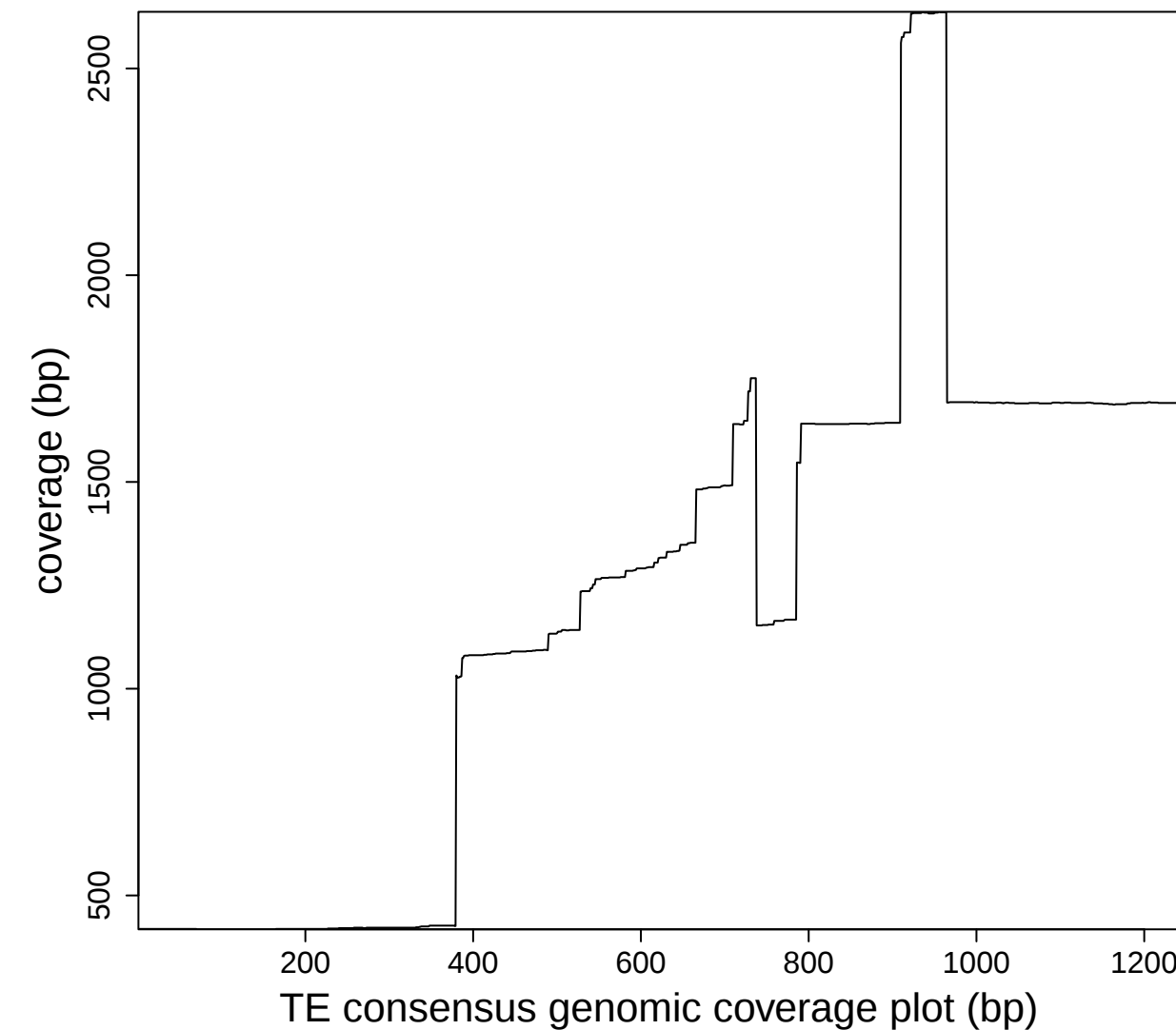
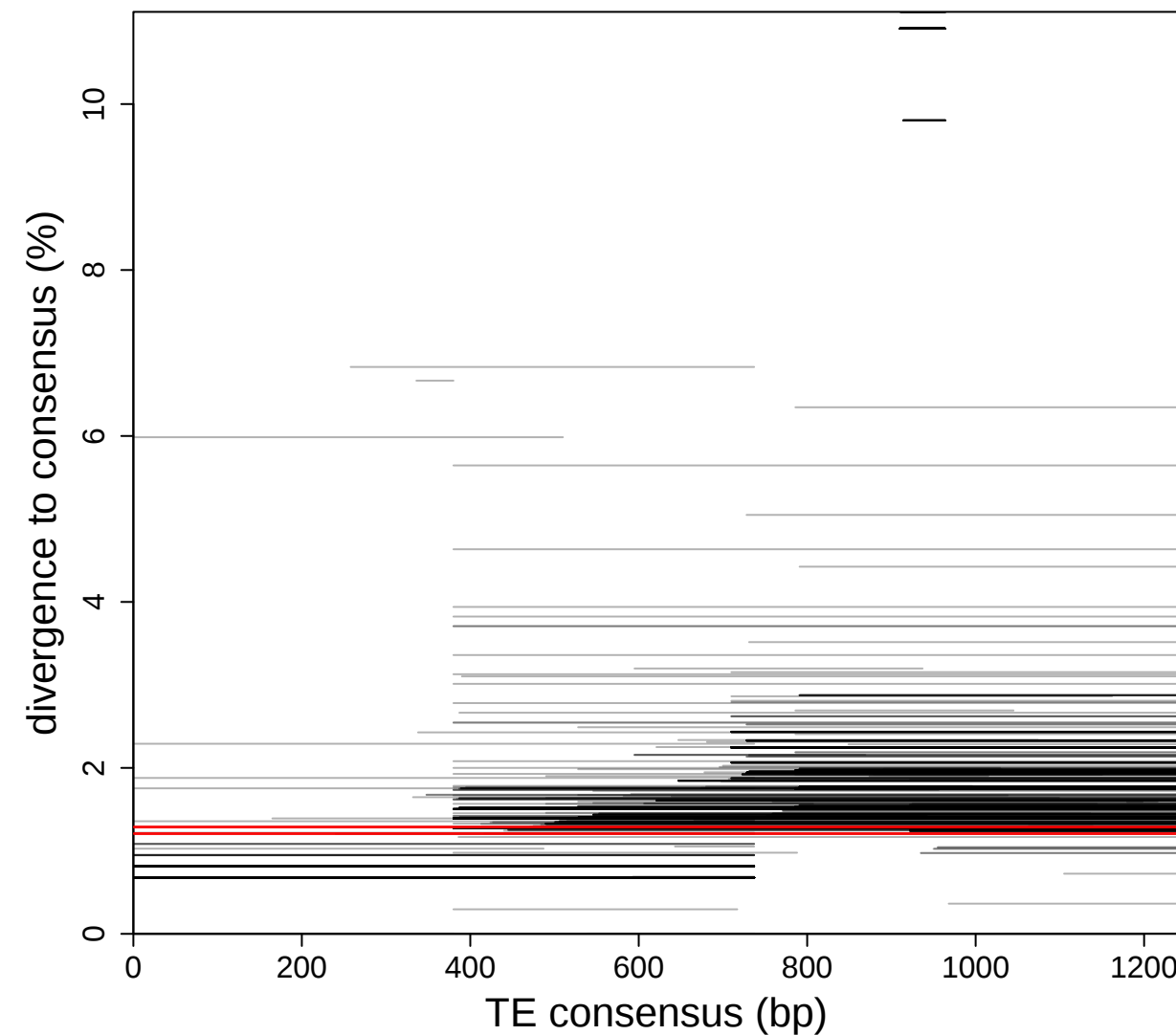


ed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.f
size: 2521bp; fragments: 5726; full length: 0 (>=2268.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

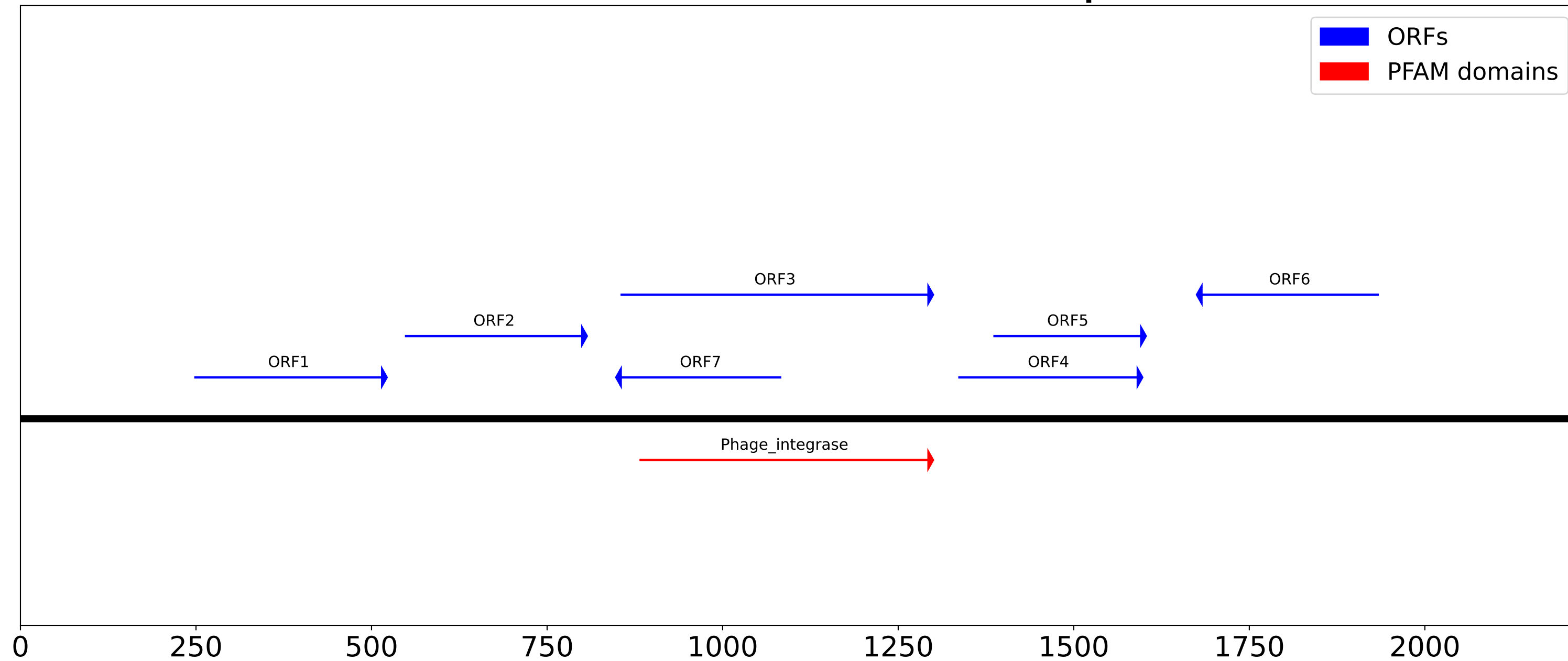


TE: rnd_1_family_102
size: 1242bp; fragments: 3272; full length: 2 (≥ 1117.8 bp)

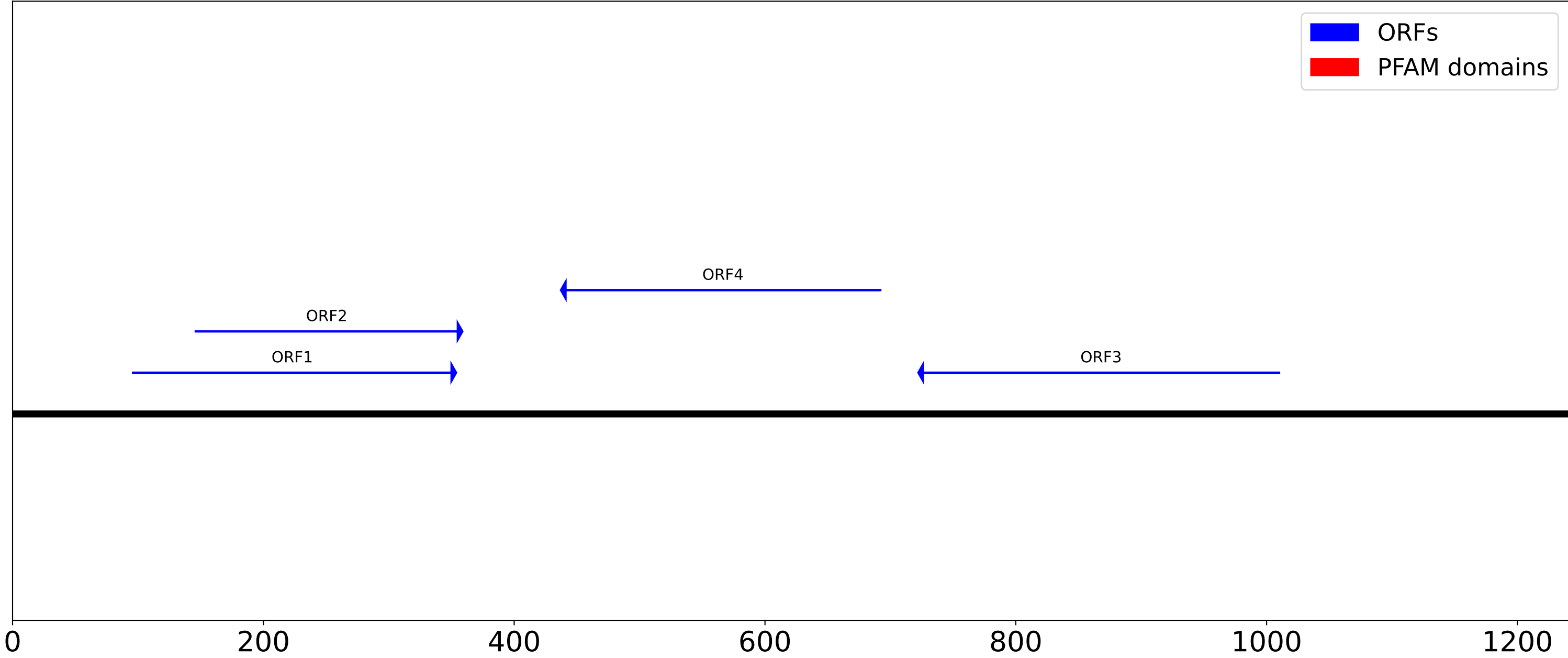


Before TEtrimmer 1242 bp

After TEtrimmer ORF and PFAM domain plot



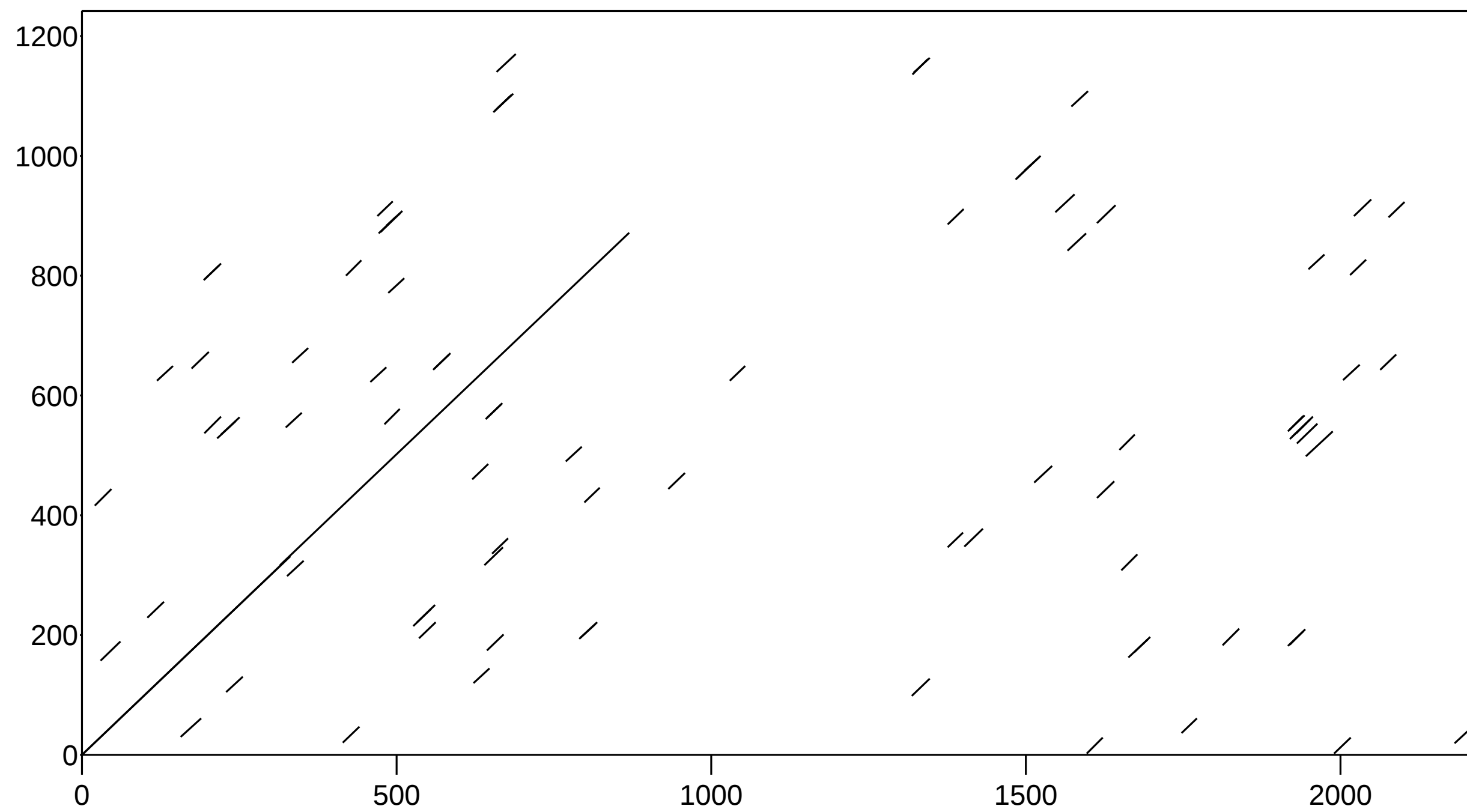
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)