

Start crop Point

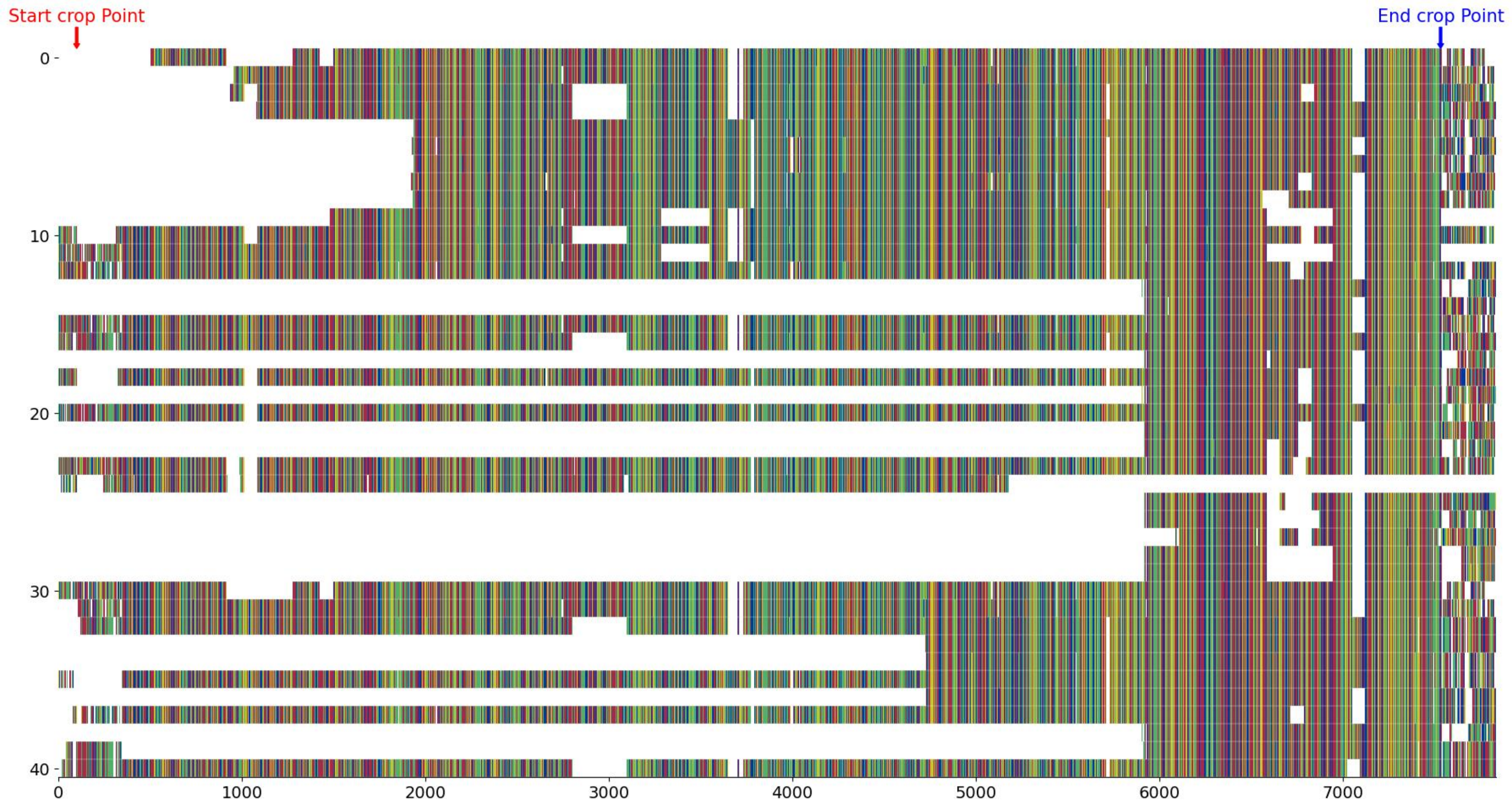


End crop Point



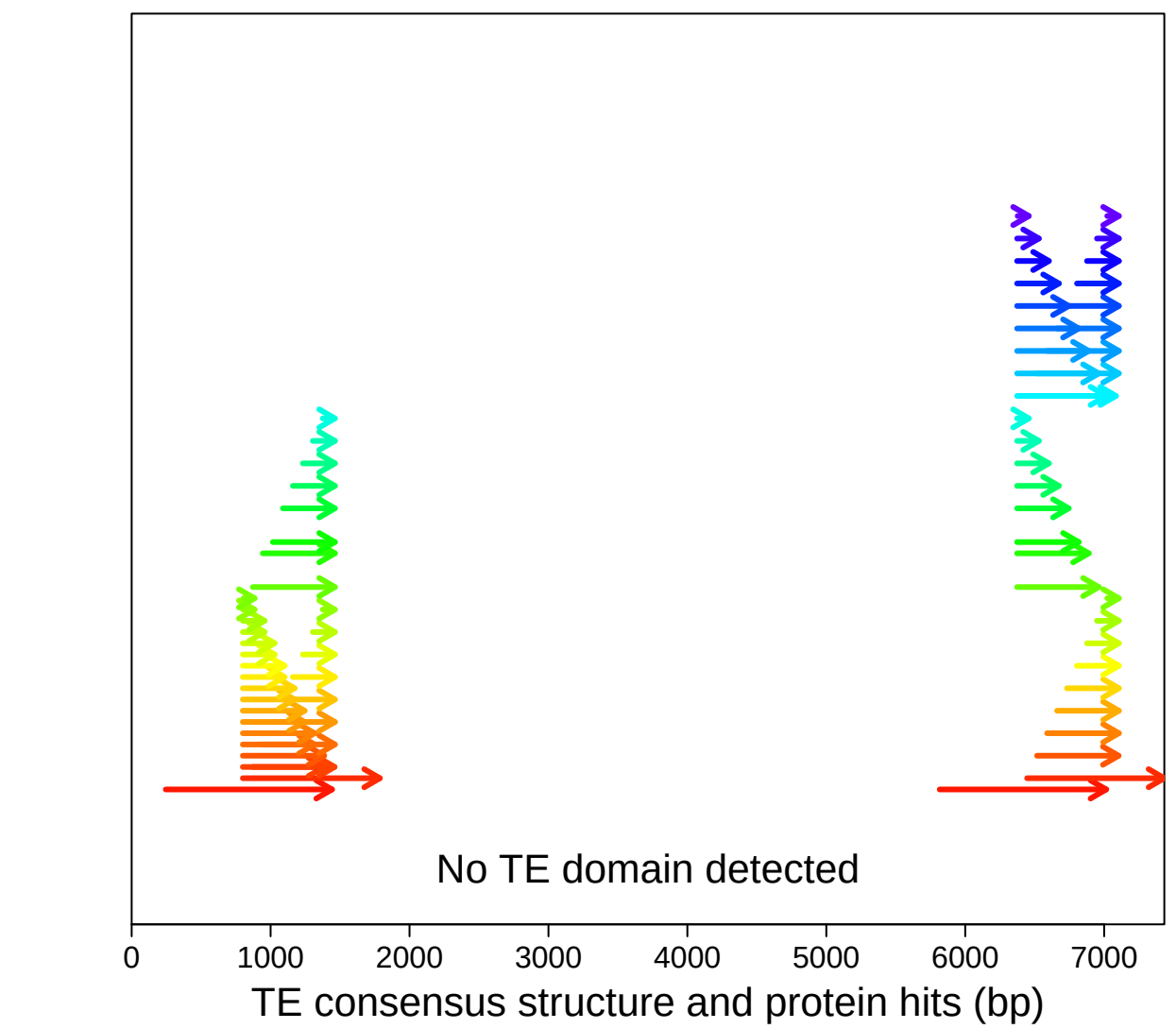
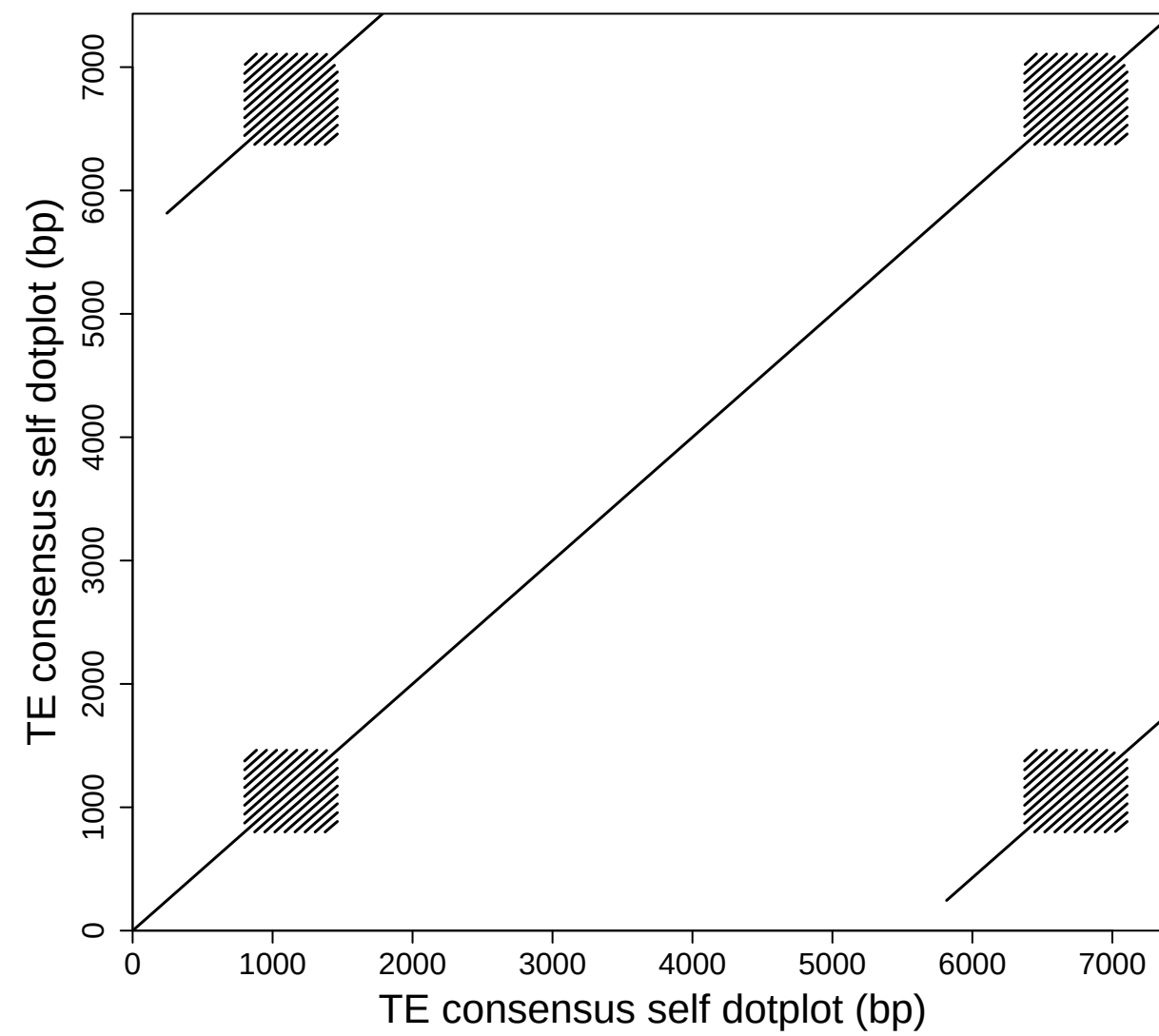
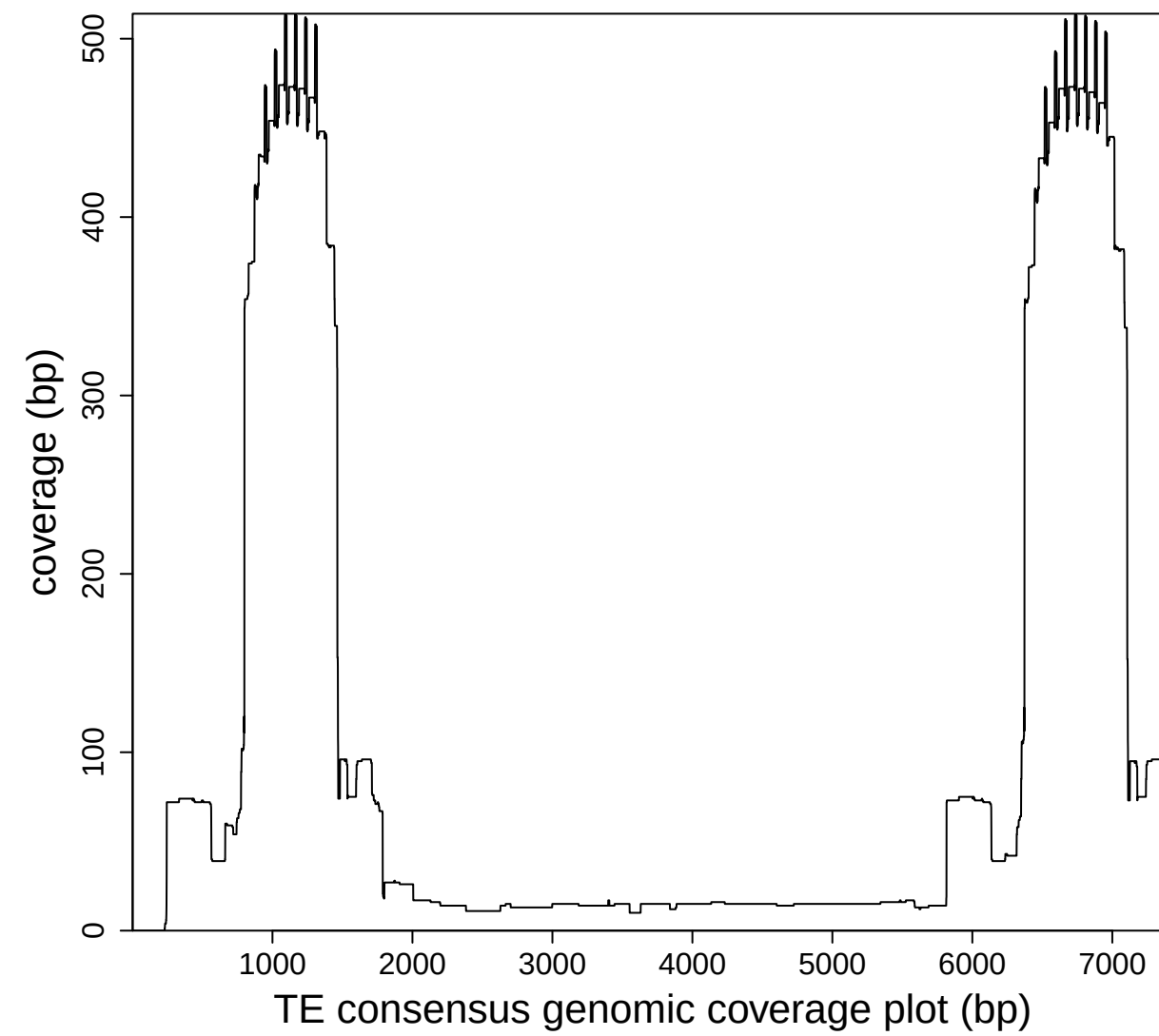
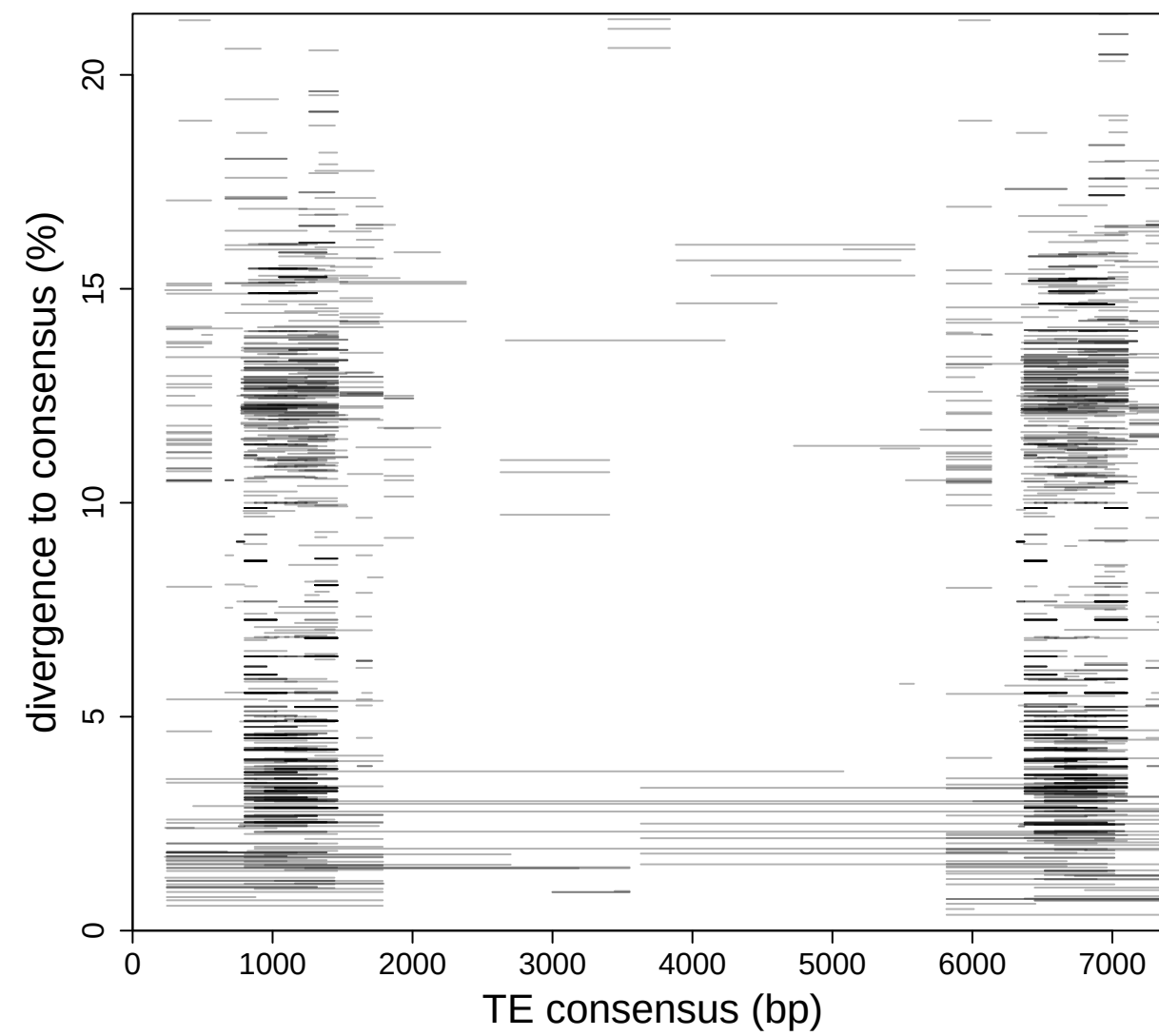
MSA length = 7433





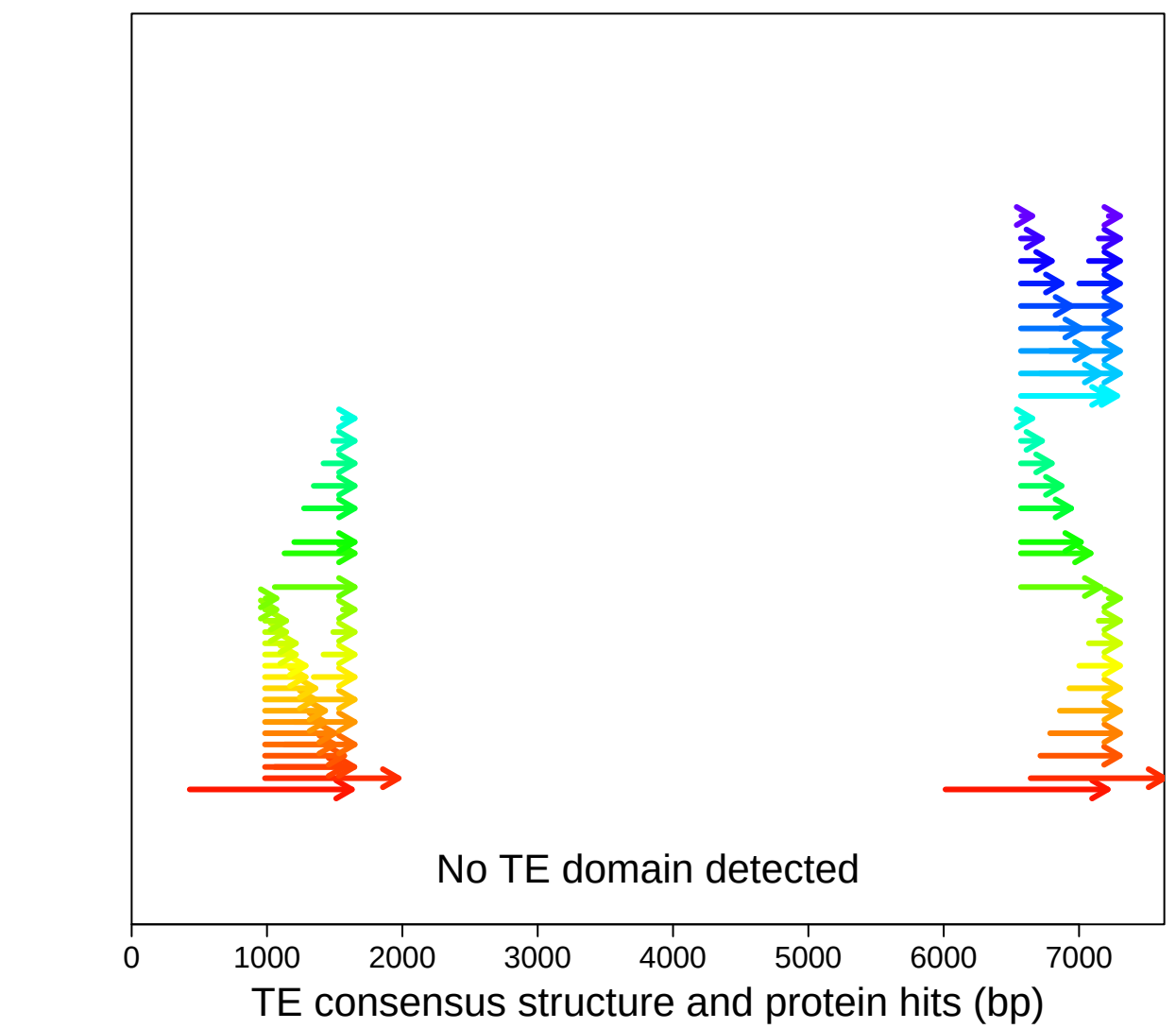
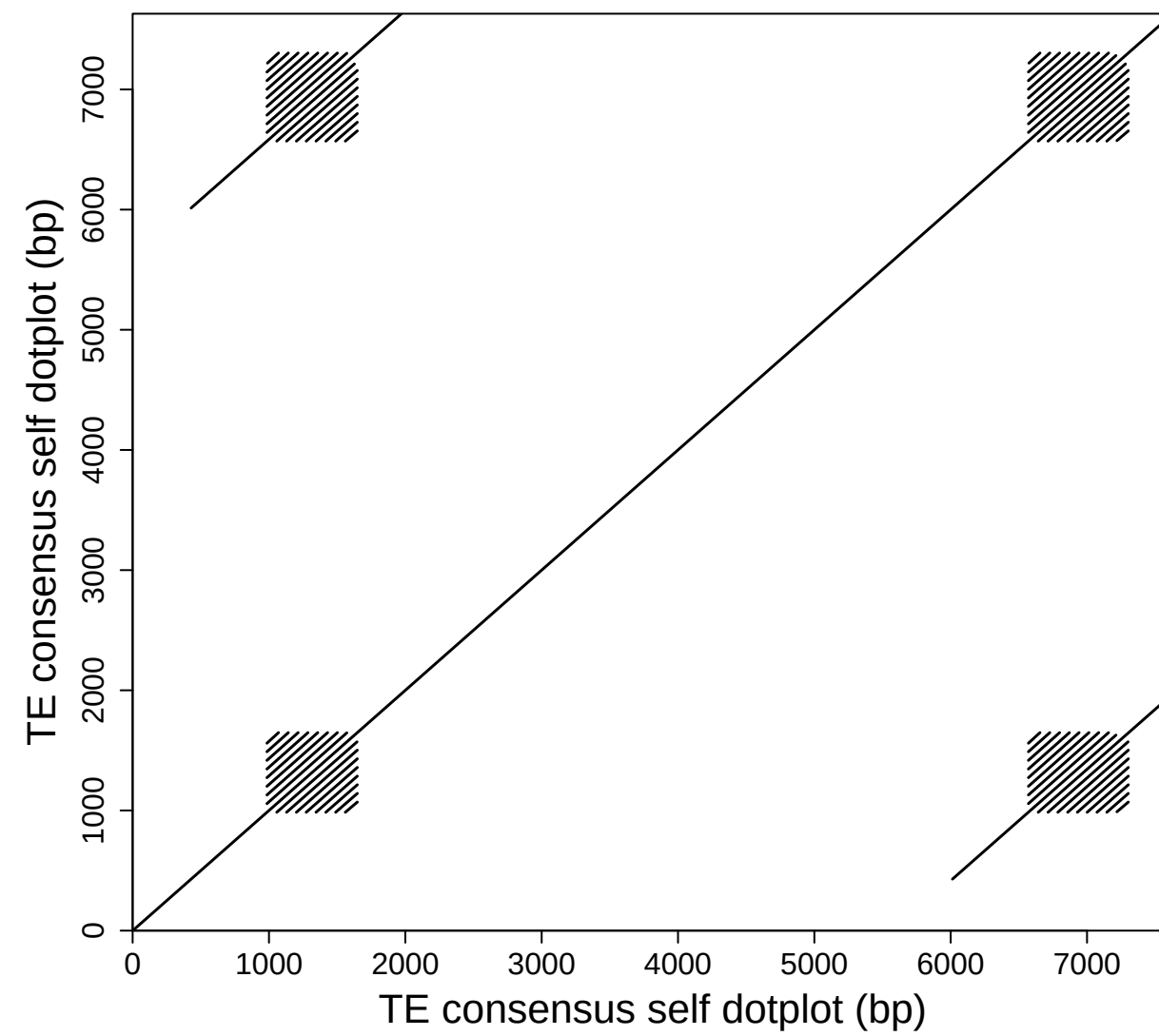
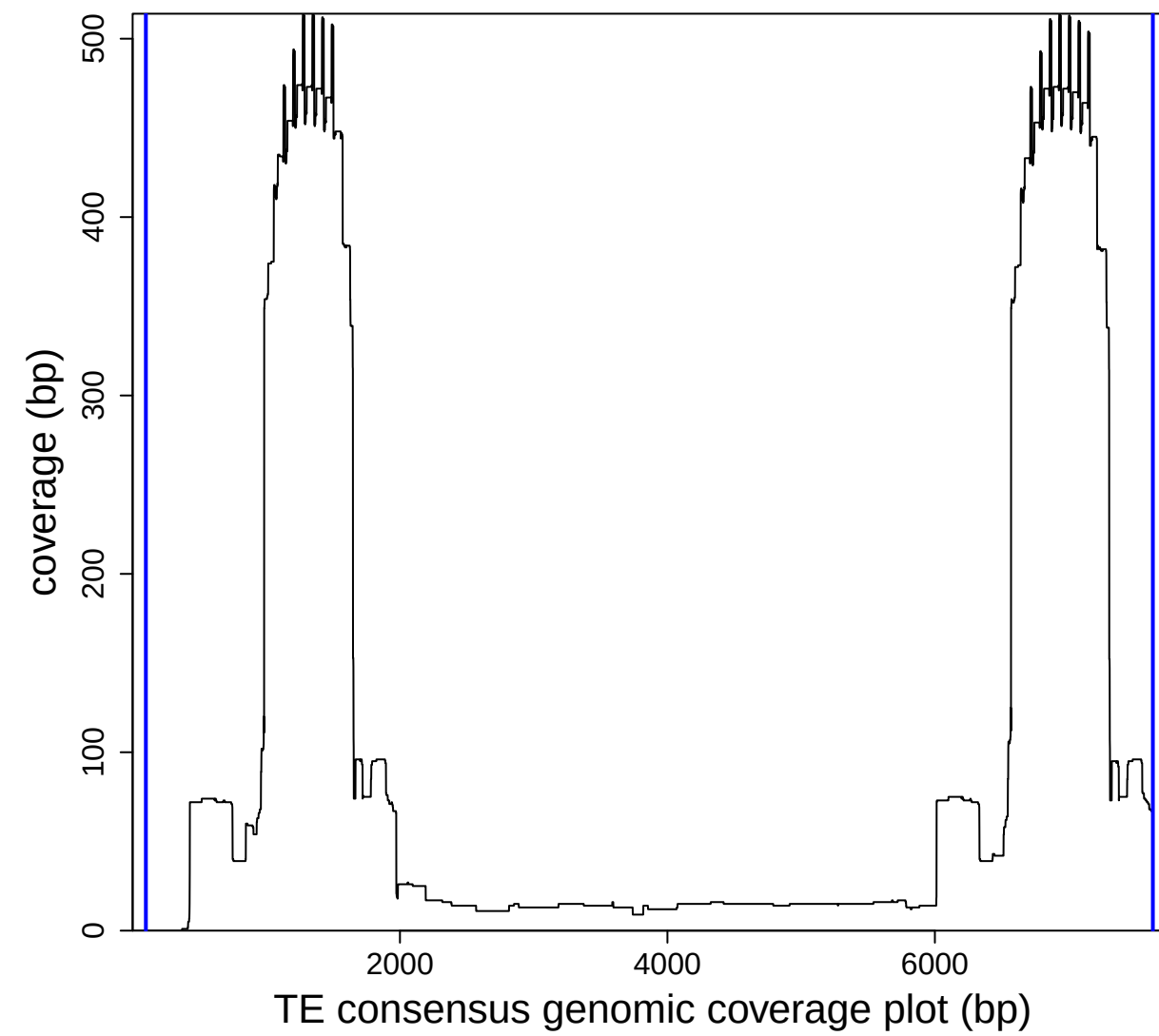
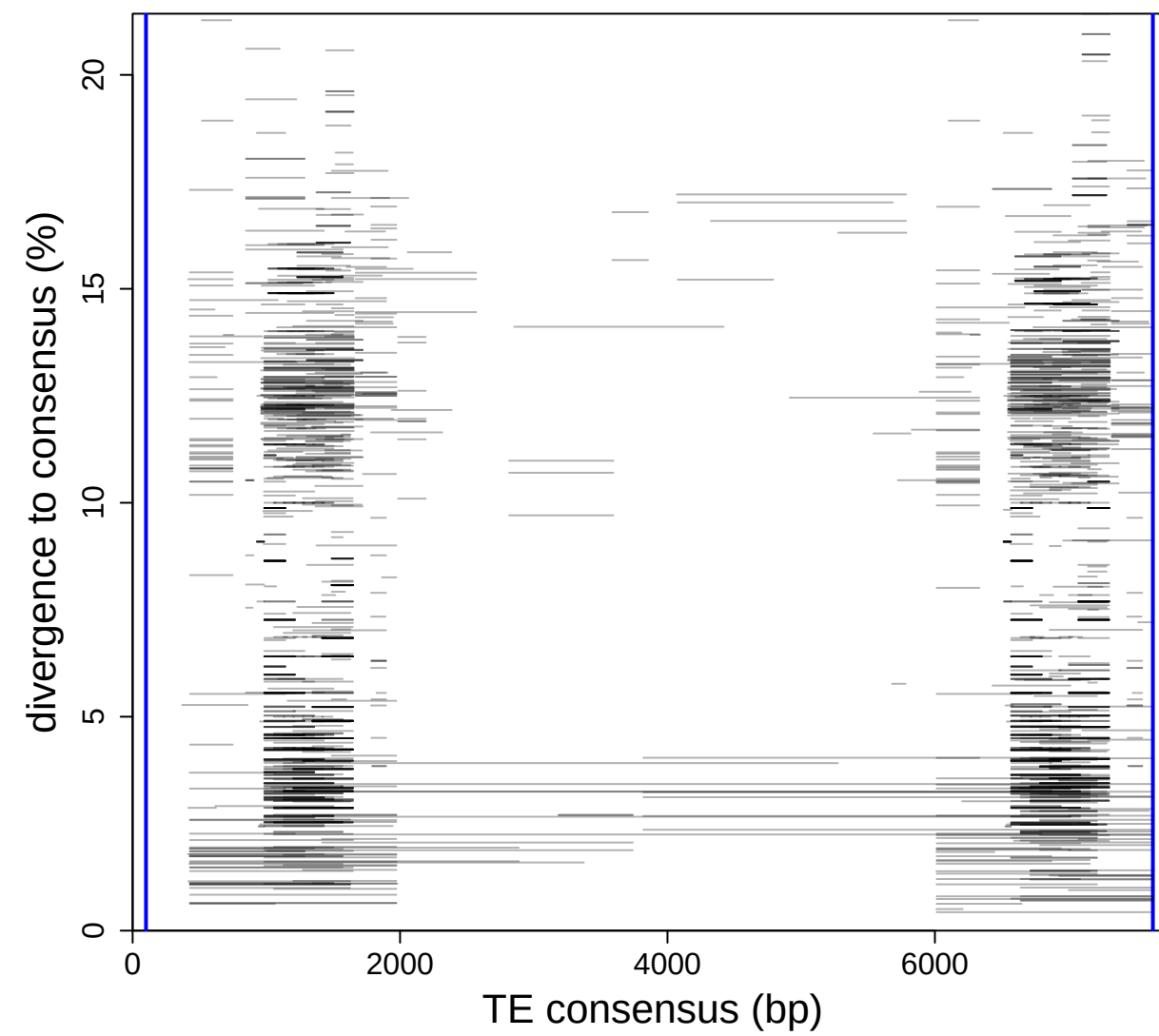
g_1.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln.fa ε
size: 7433bp; fragments: 1971; full length: 0 (>=6689.7bp)

After TETrimmer 7433 bp



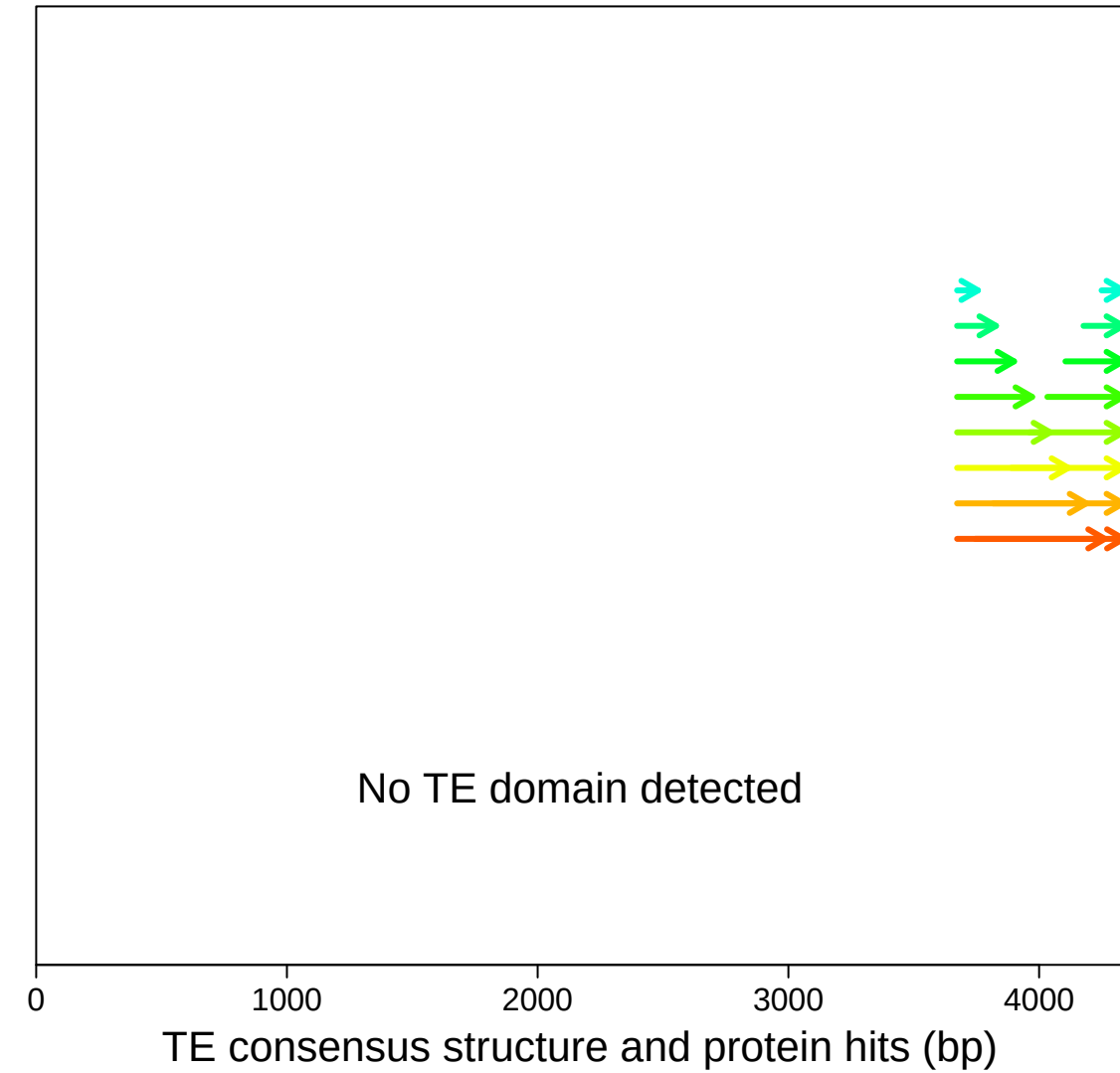
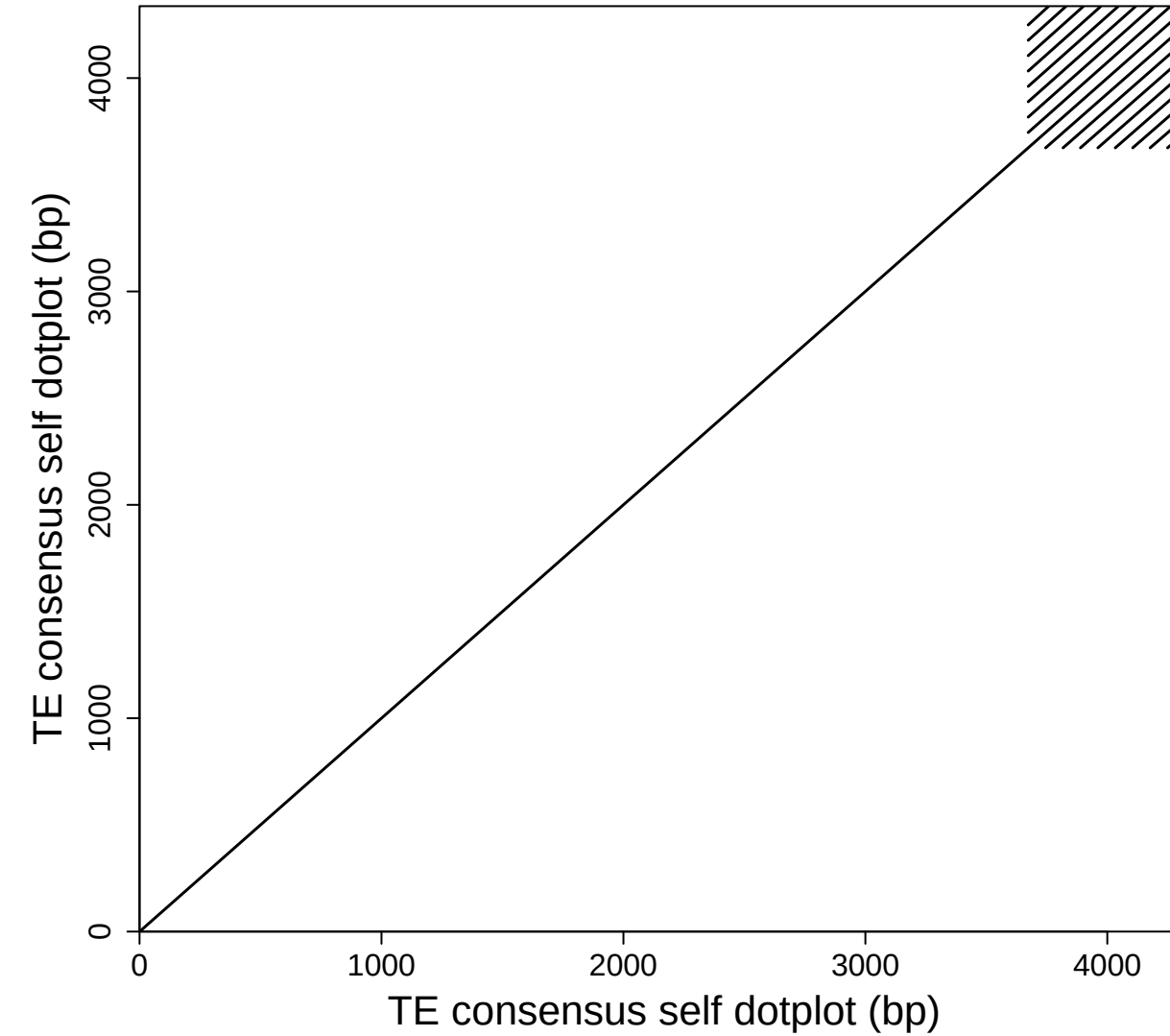
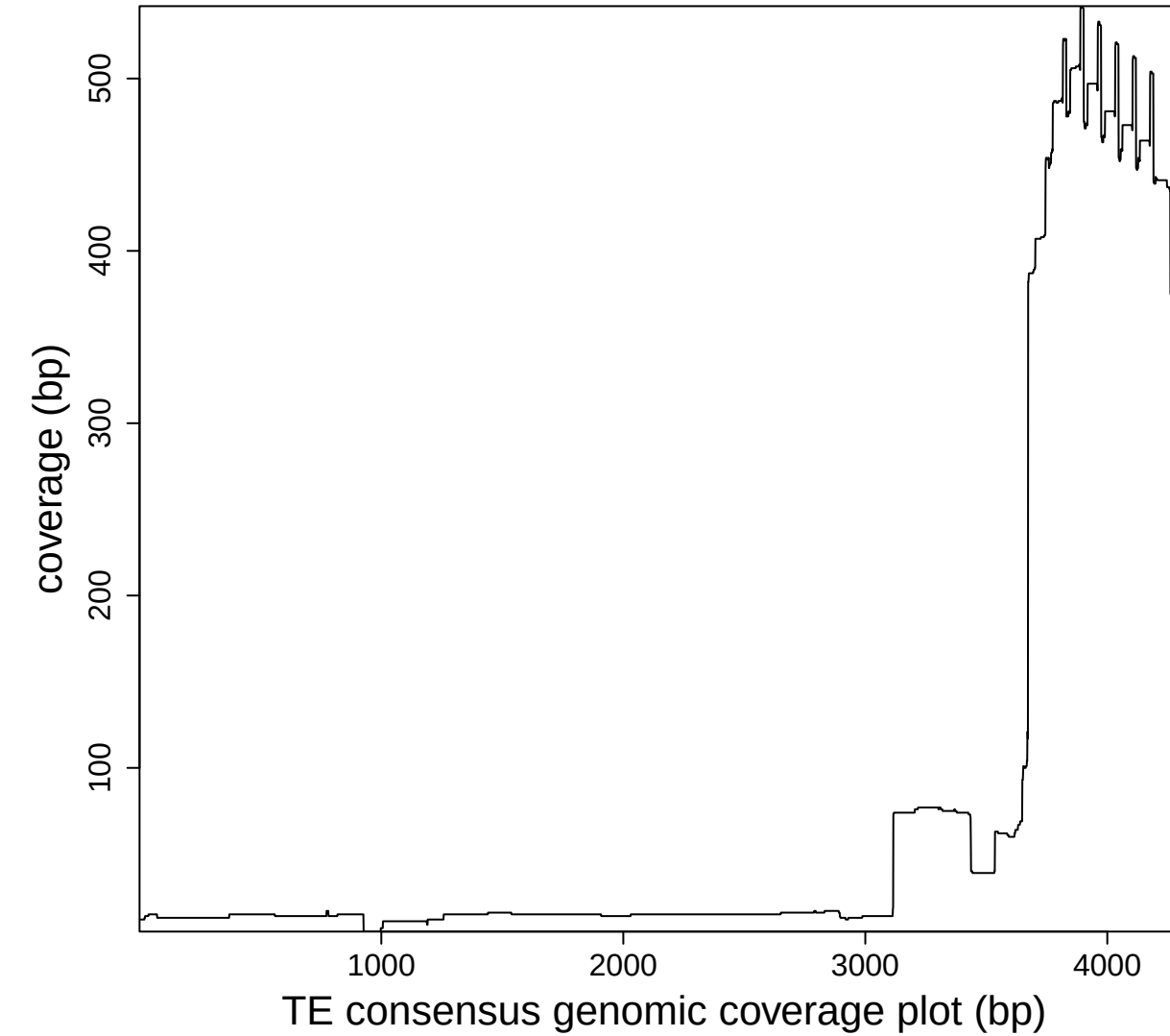
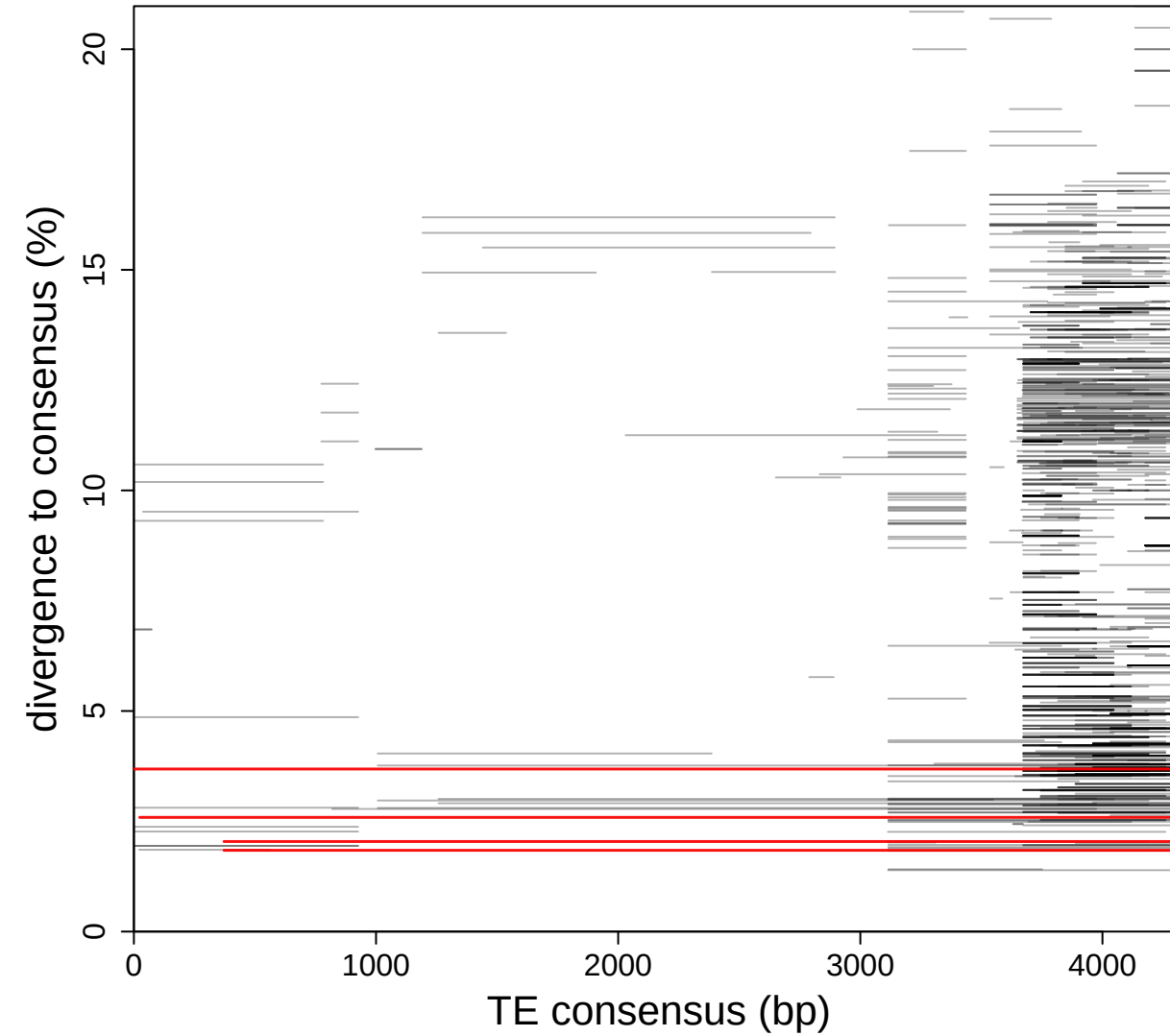
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 7780bp; fragments: 1968; full length: 0 (>=7002bp)

After TEtrimmer Extended plot Blue lines are boundaries

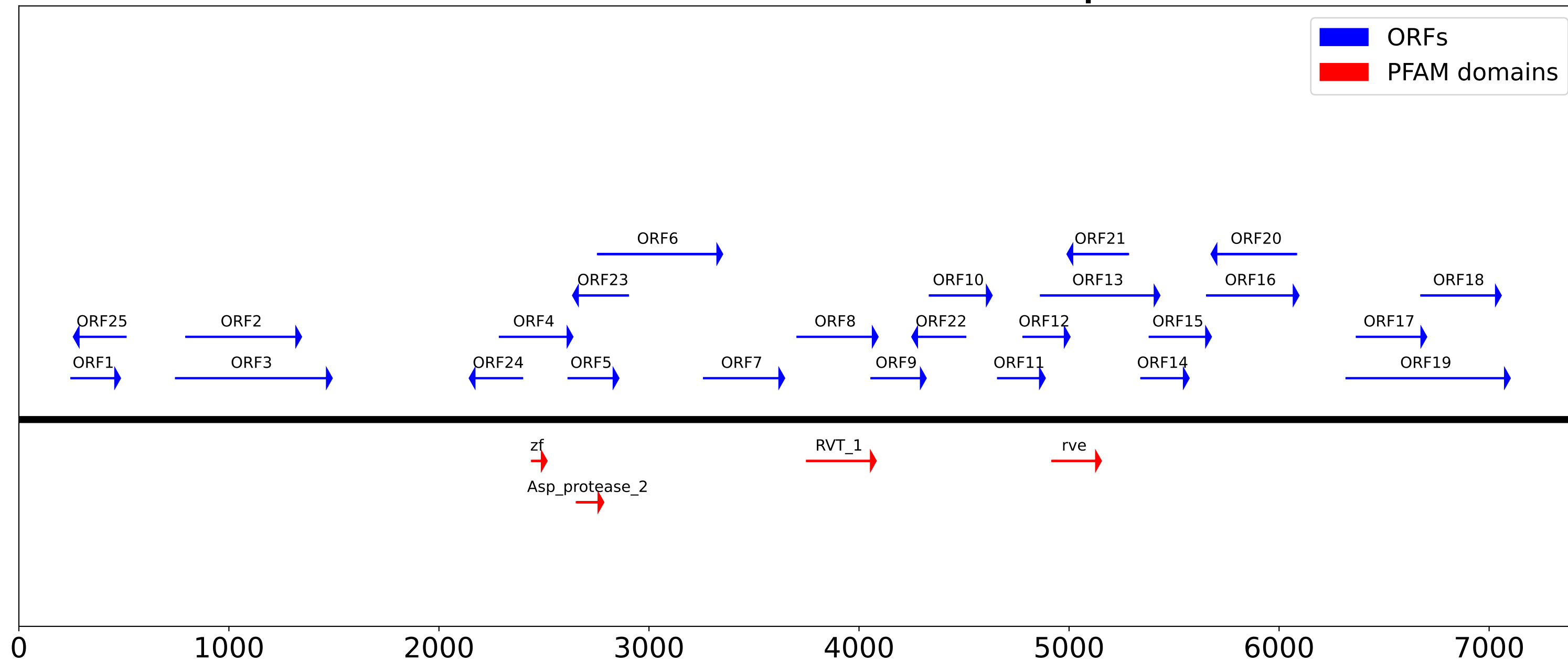


TE: rnd_5_family_2485
size: 4337bp; fragments: 952; full length: 4 (>=3903.3bp)

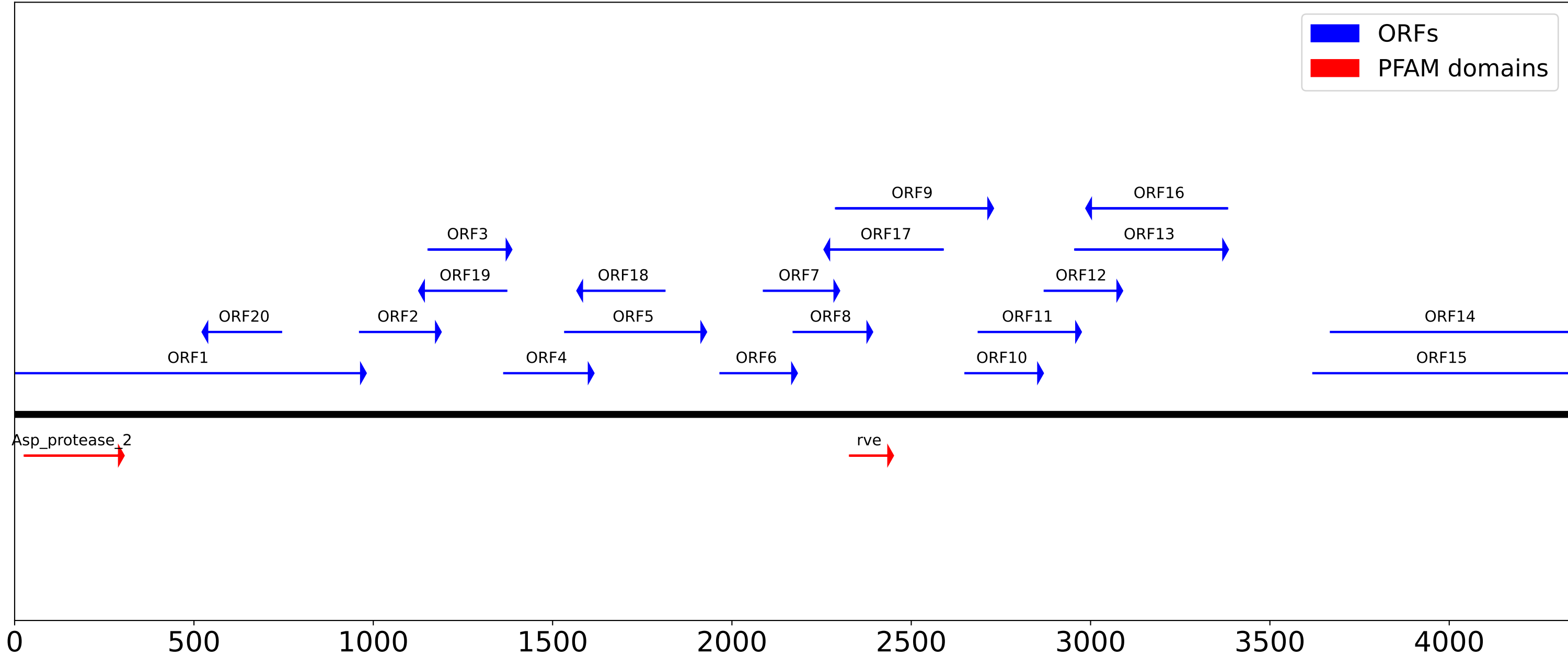
Before TEtrimmer 4337 bp



After TETrimmer ORF and PFAM domain plot



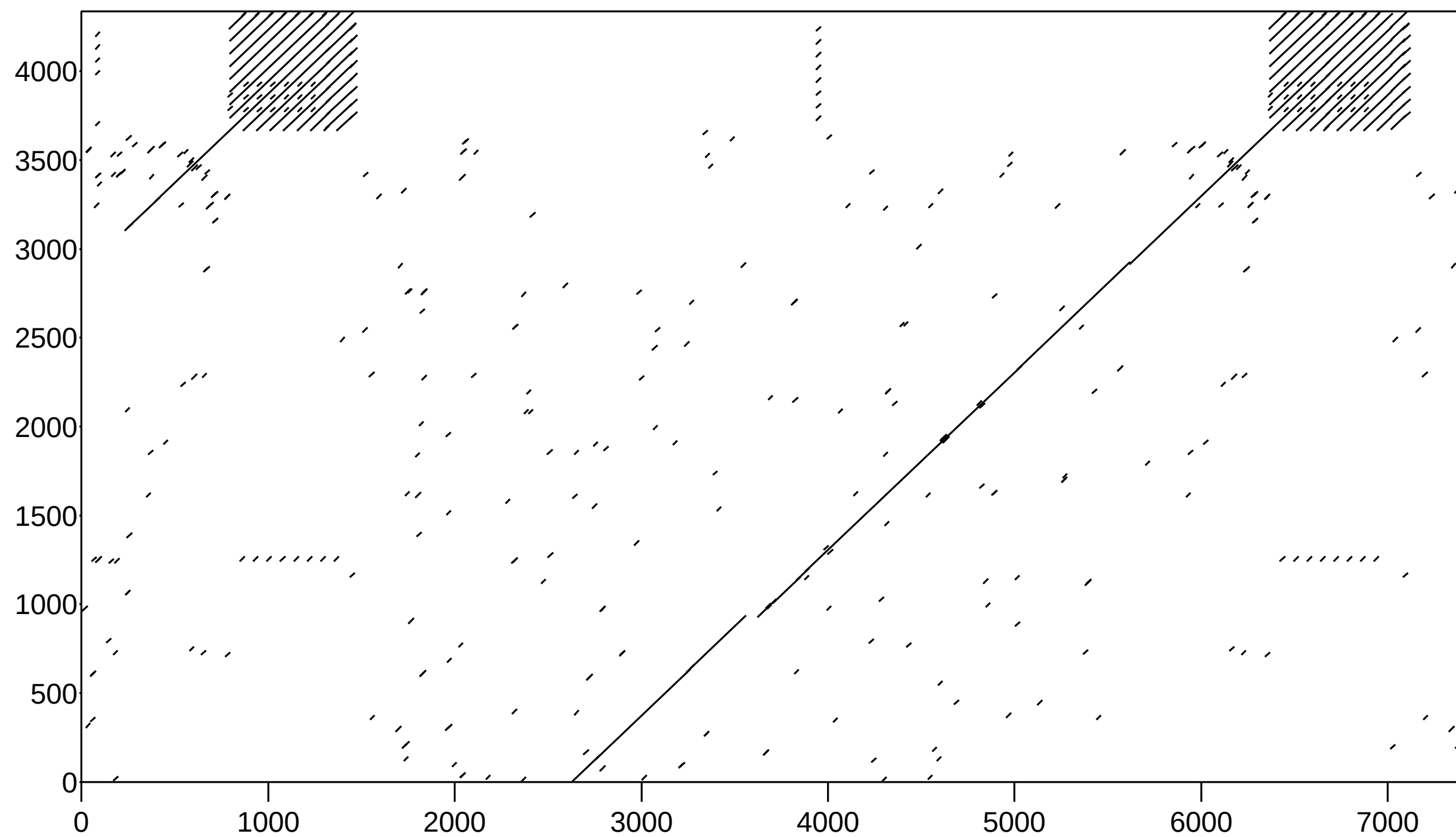
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)