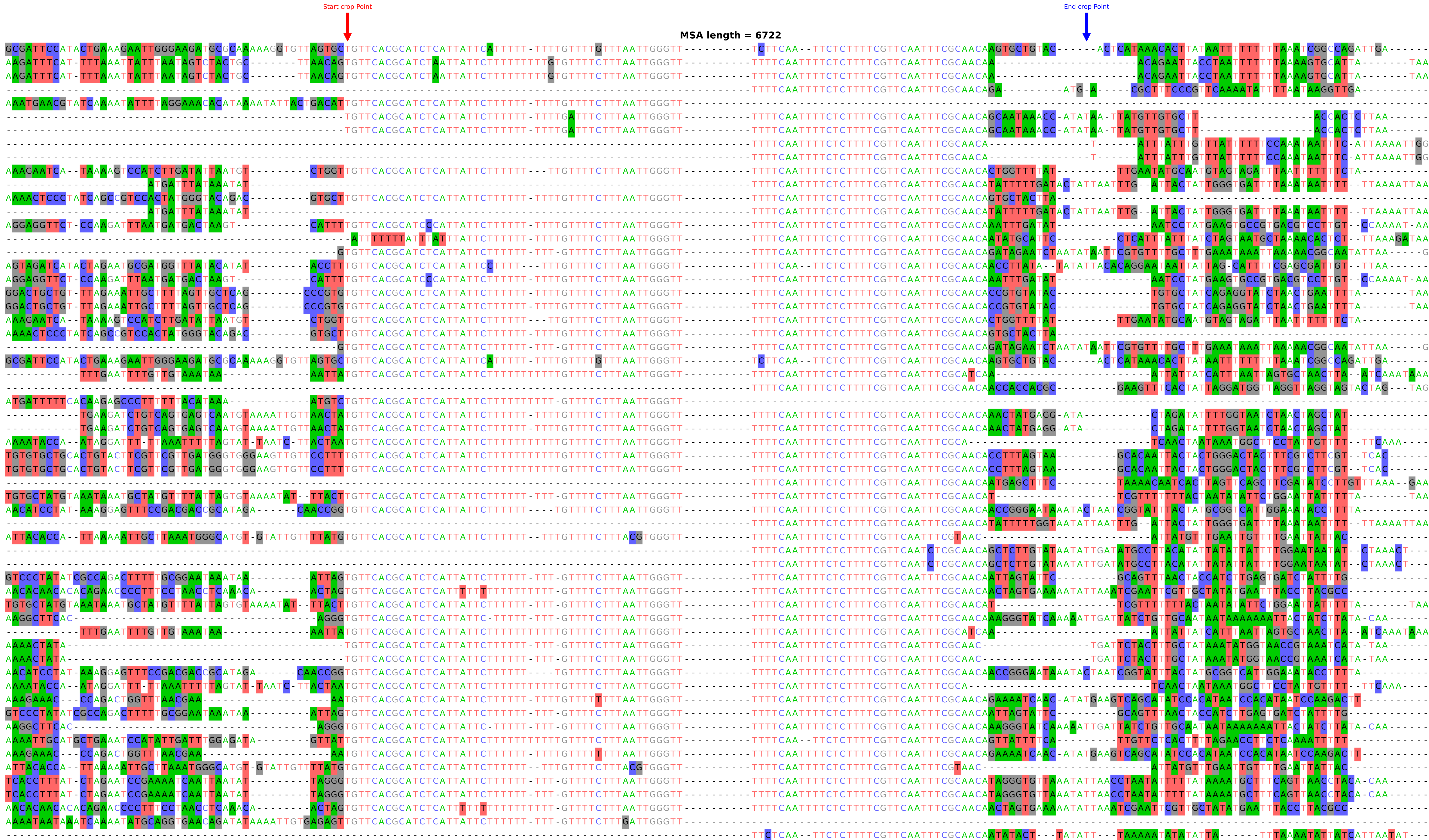
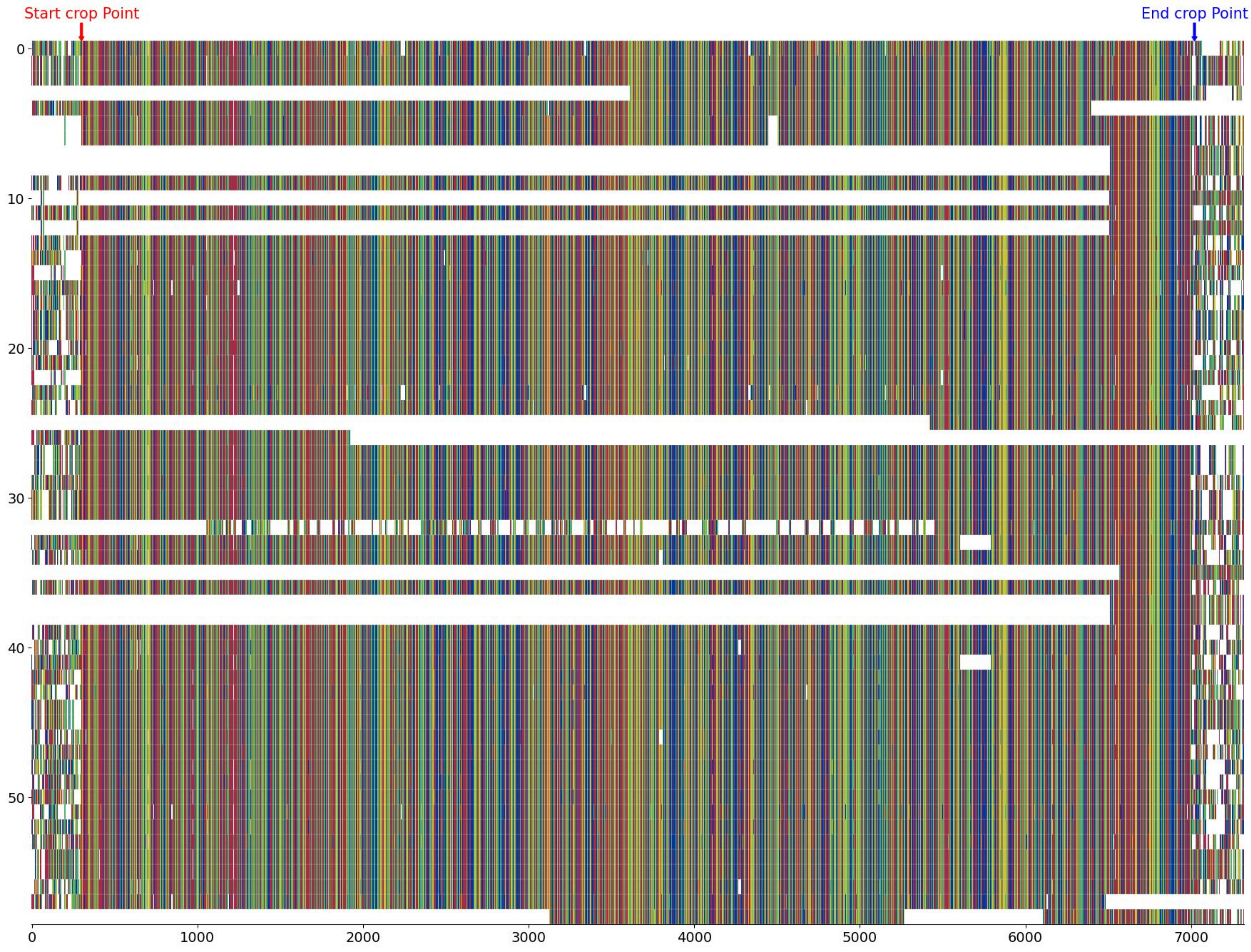




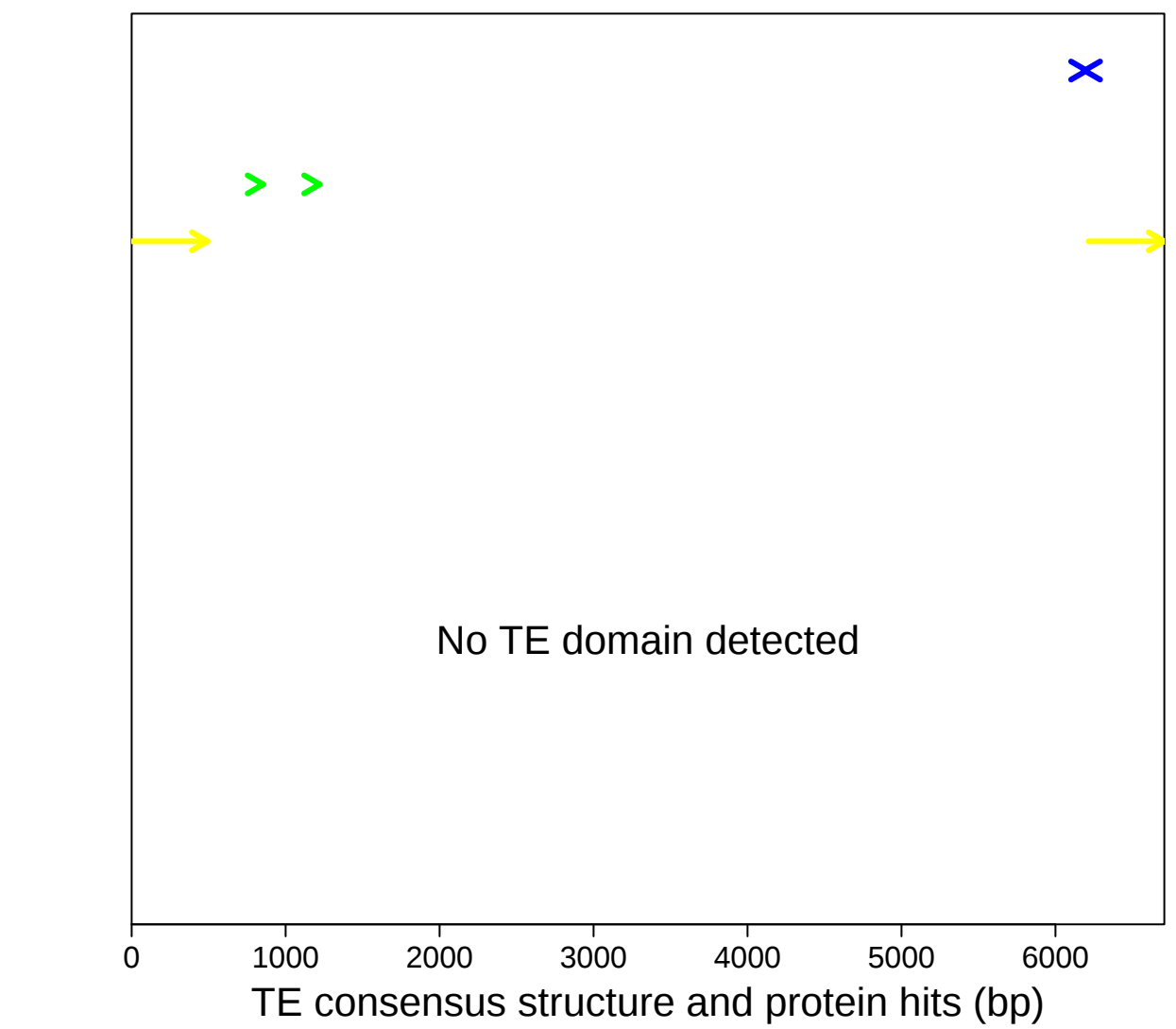
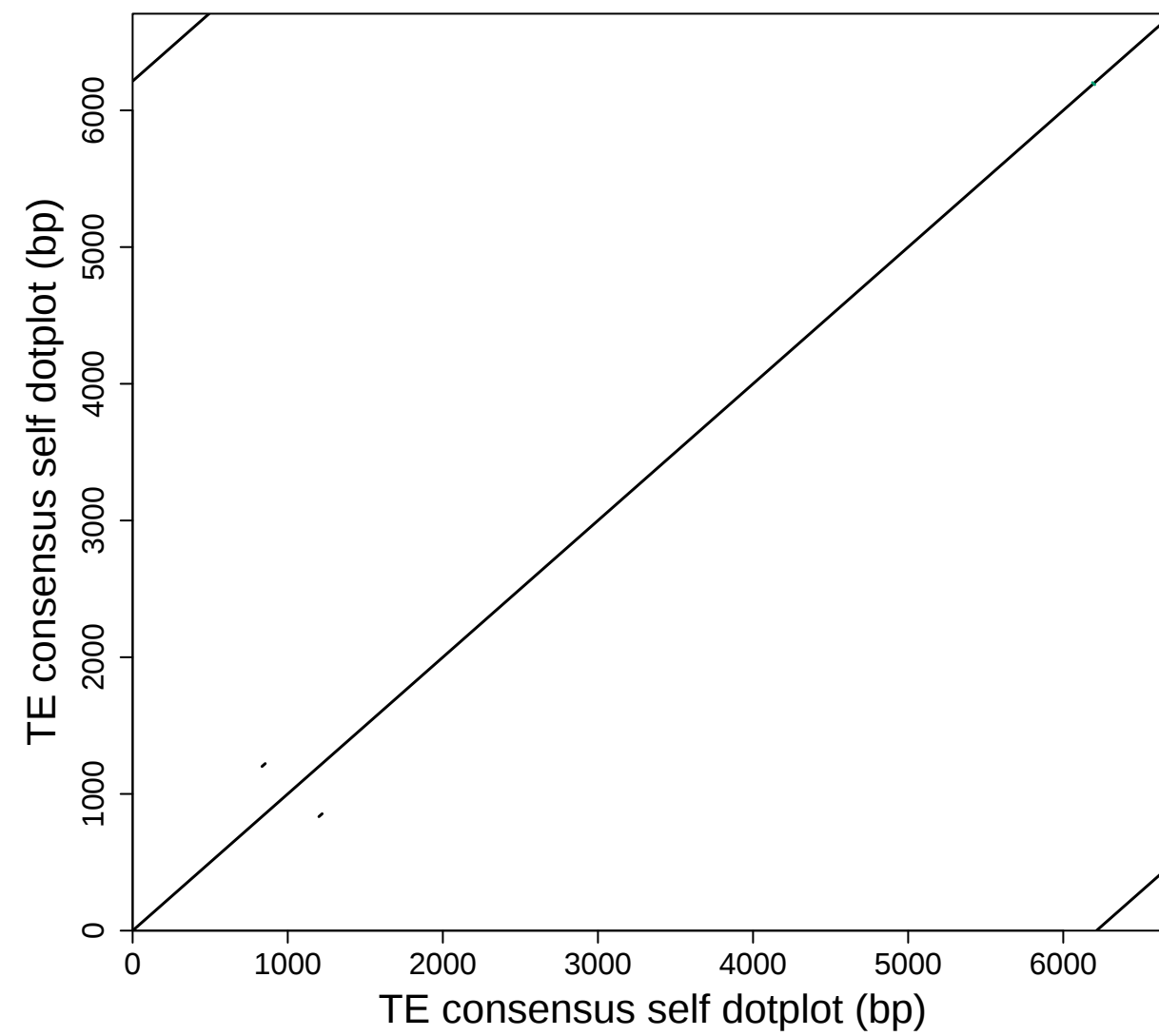
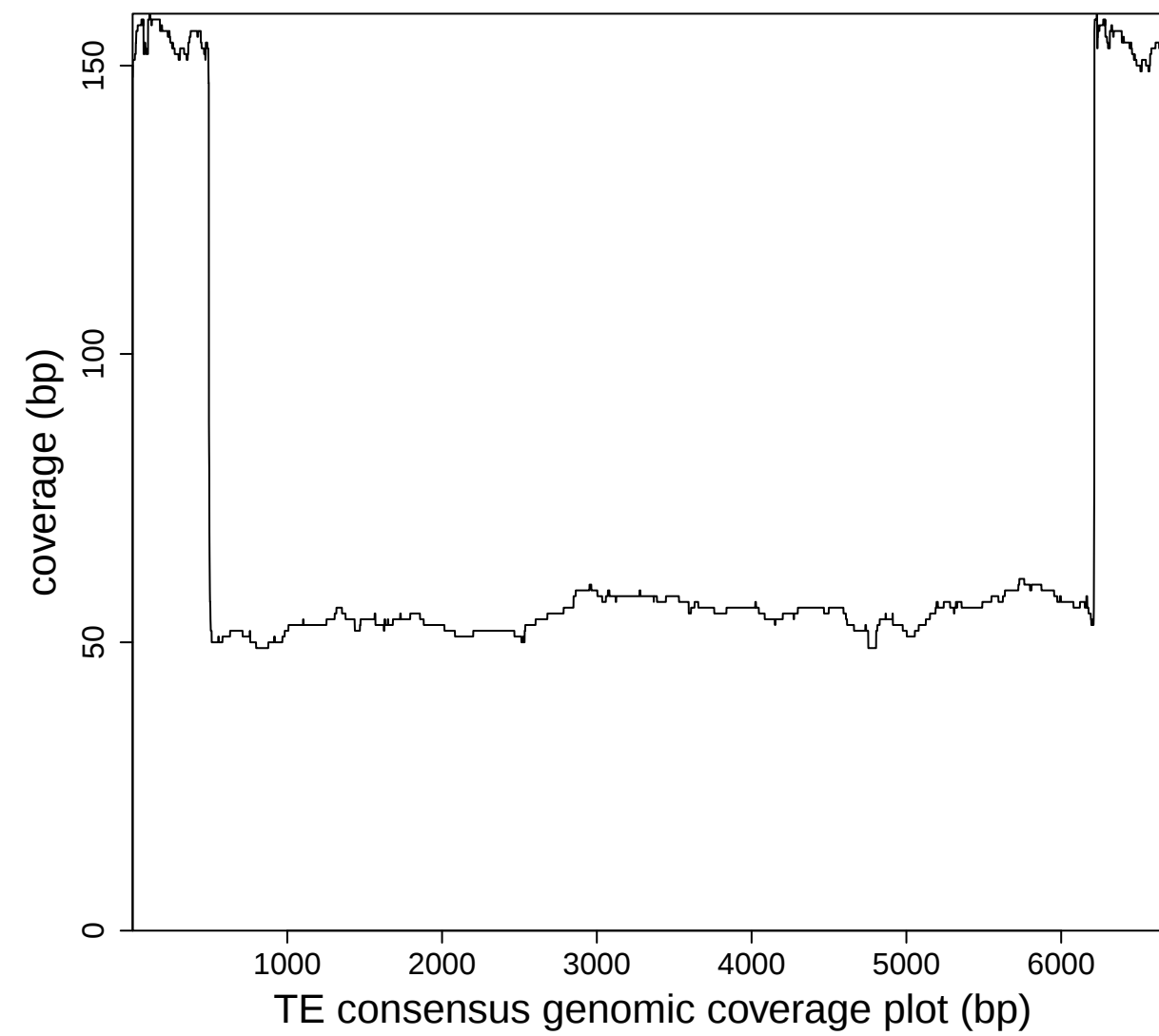
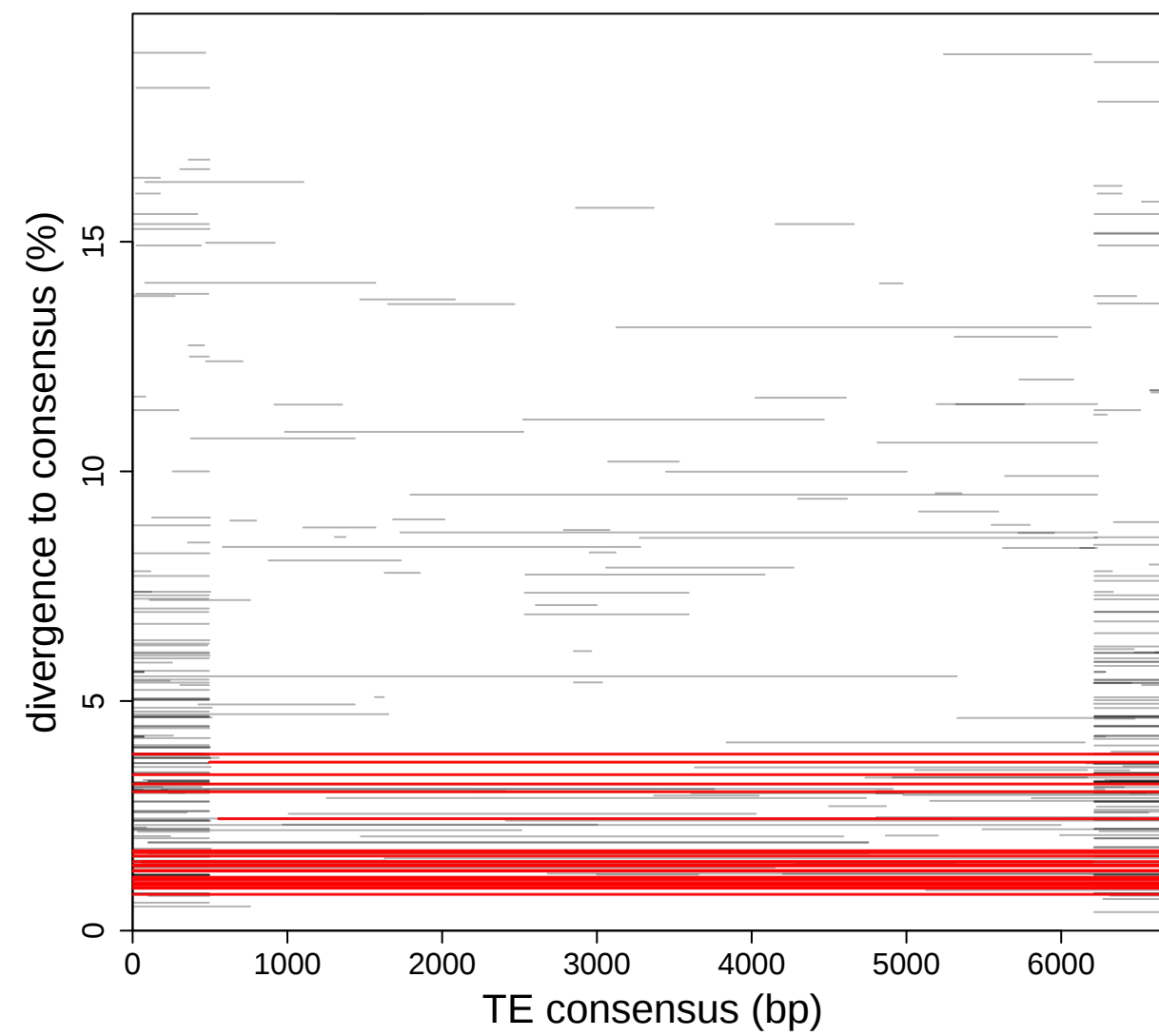
Start crop Point

End crop Point



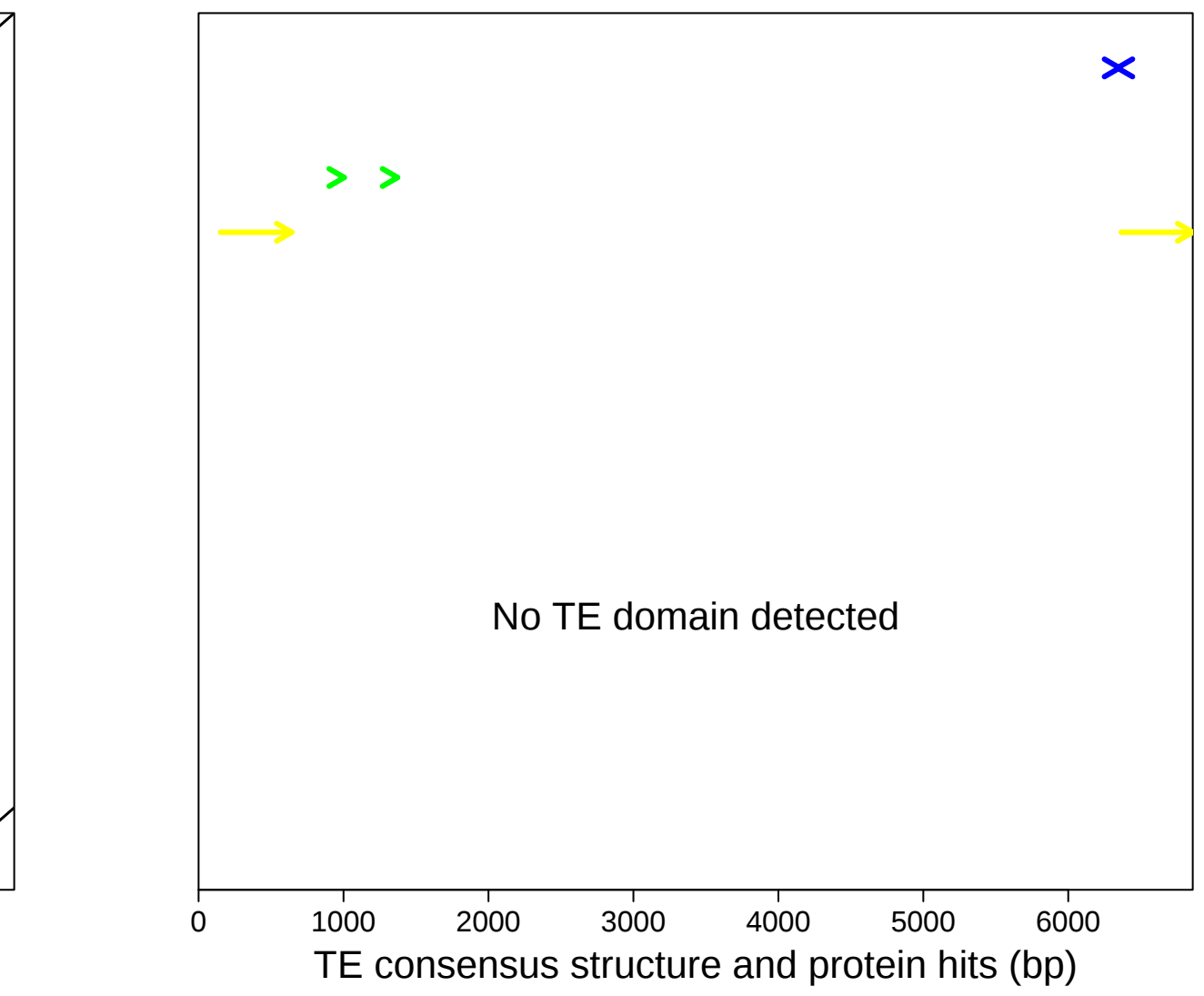
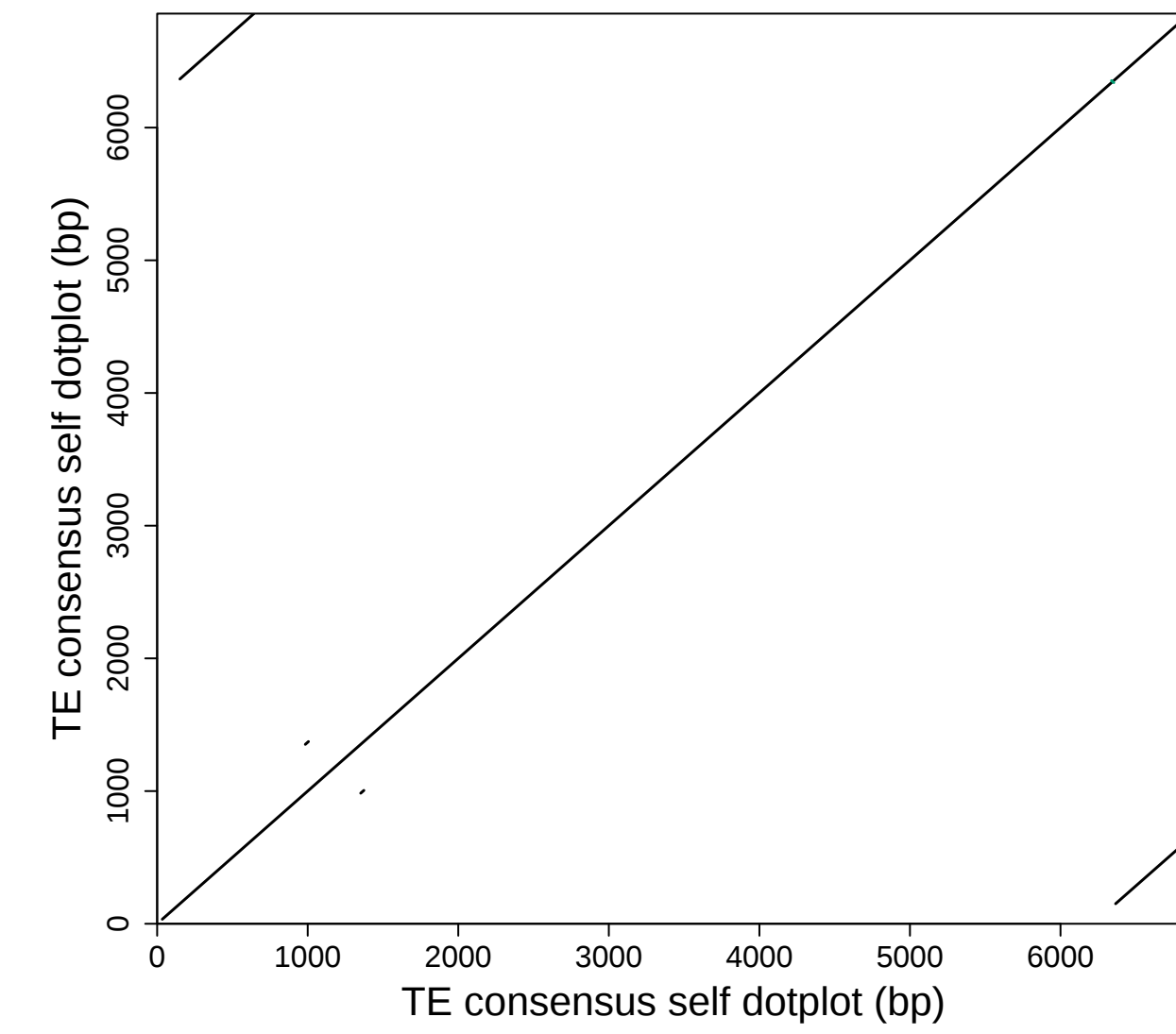
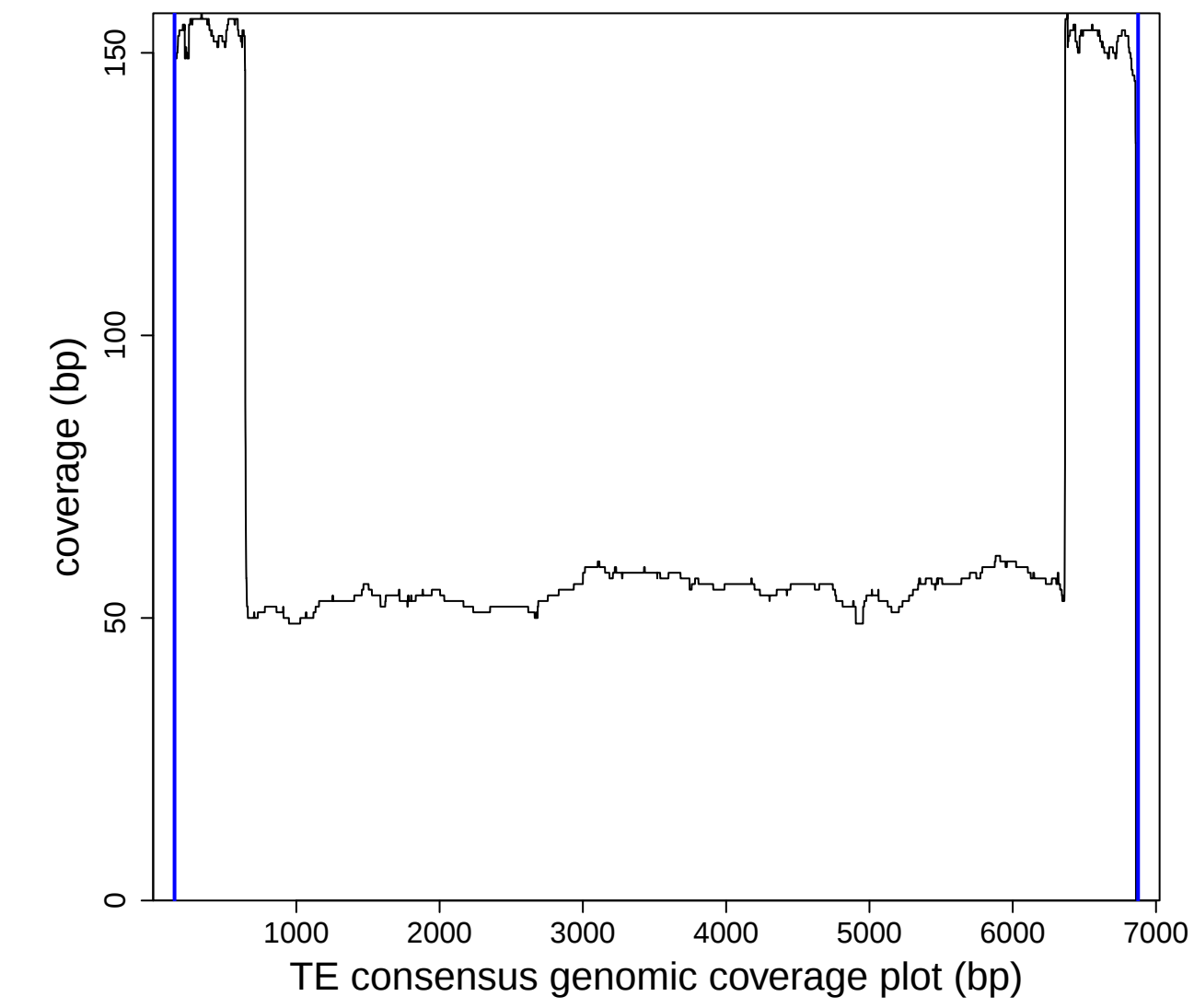
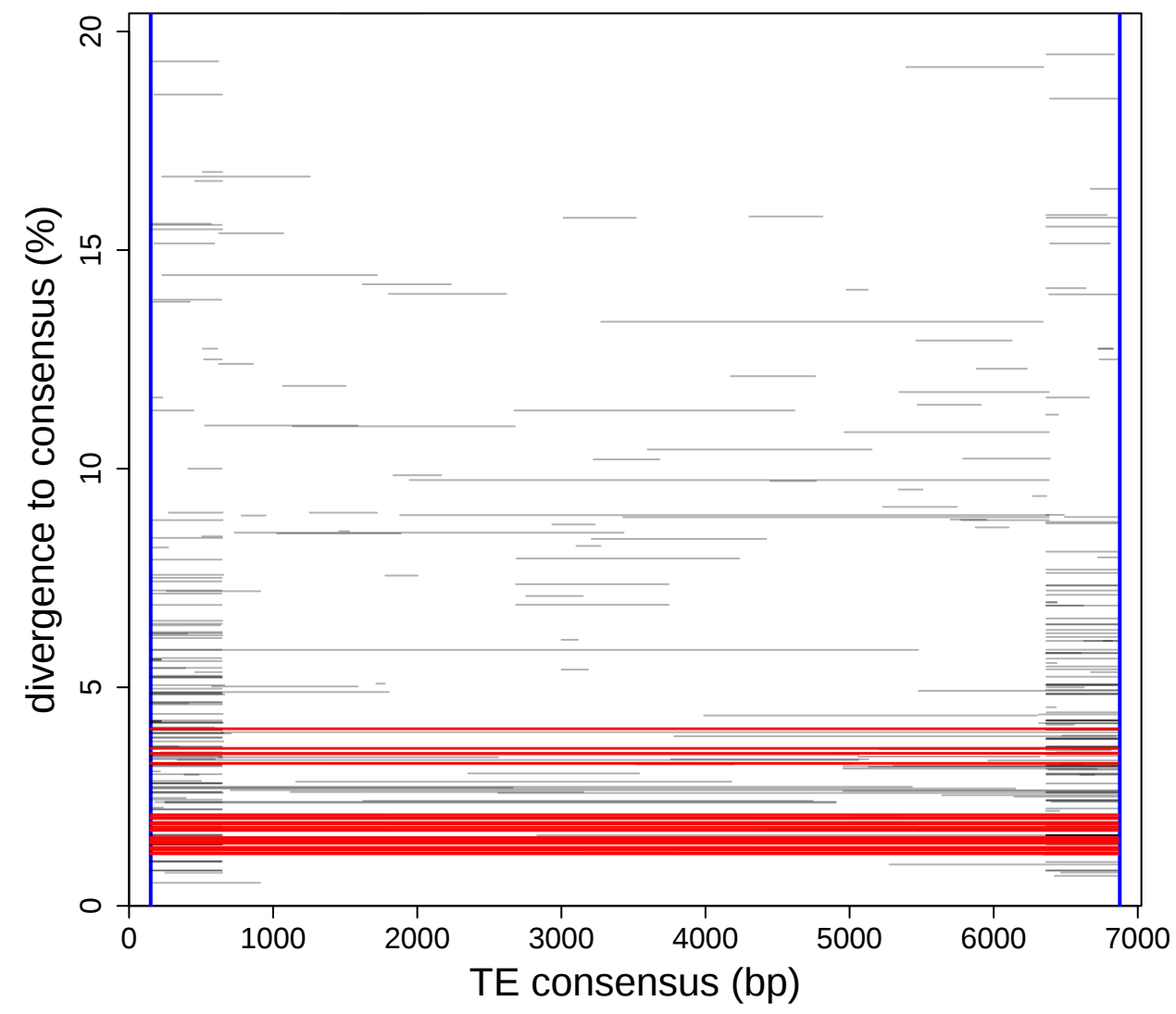
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_0
size: 6722bp; fragments: 395; full length: 26 (>=6049.8bp)

After TETrimmer 6722 bp

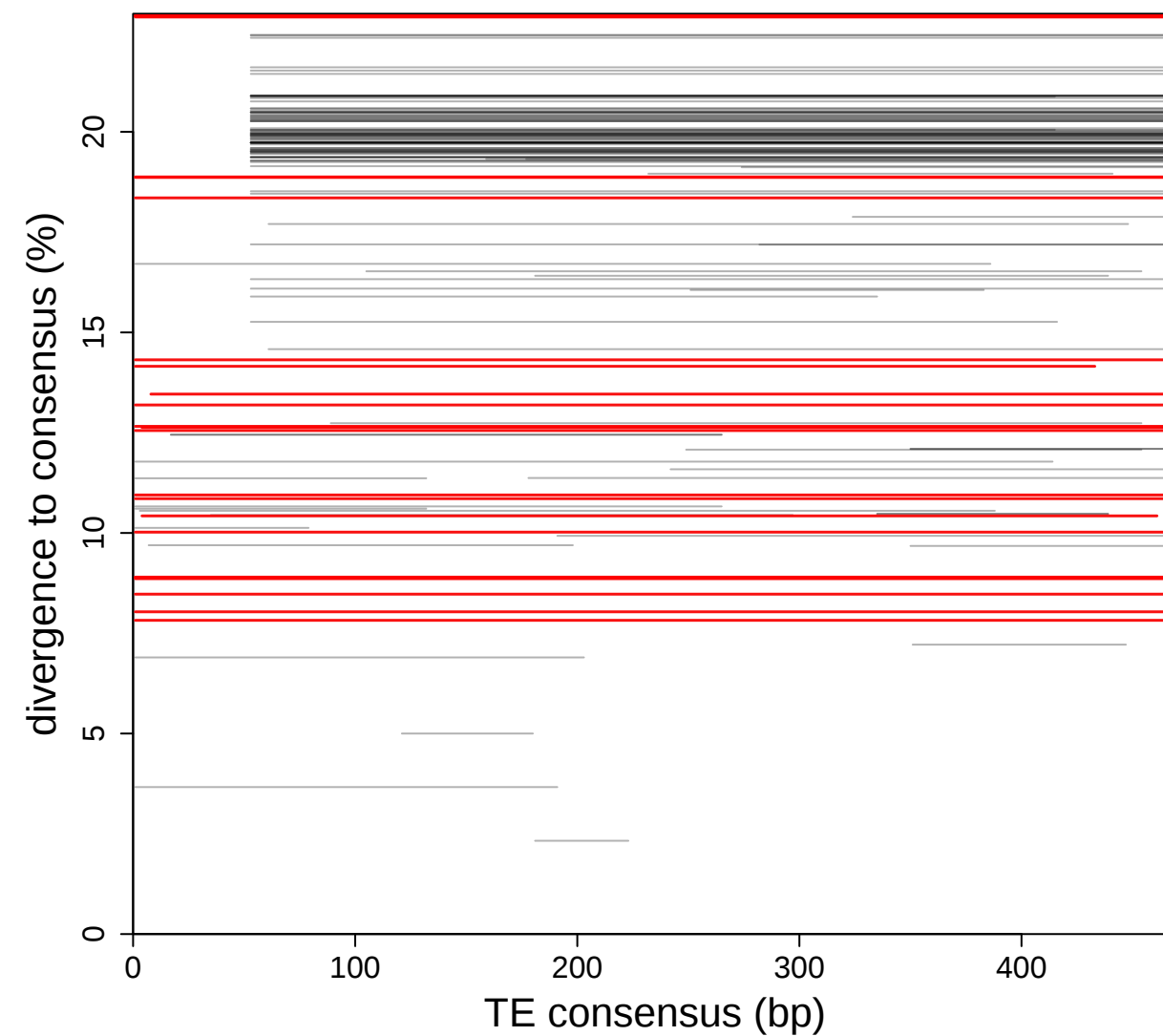


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 7025bp; fragments: 389; full length: 23 (≥ 6322.5 bp)

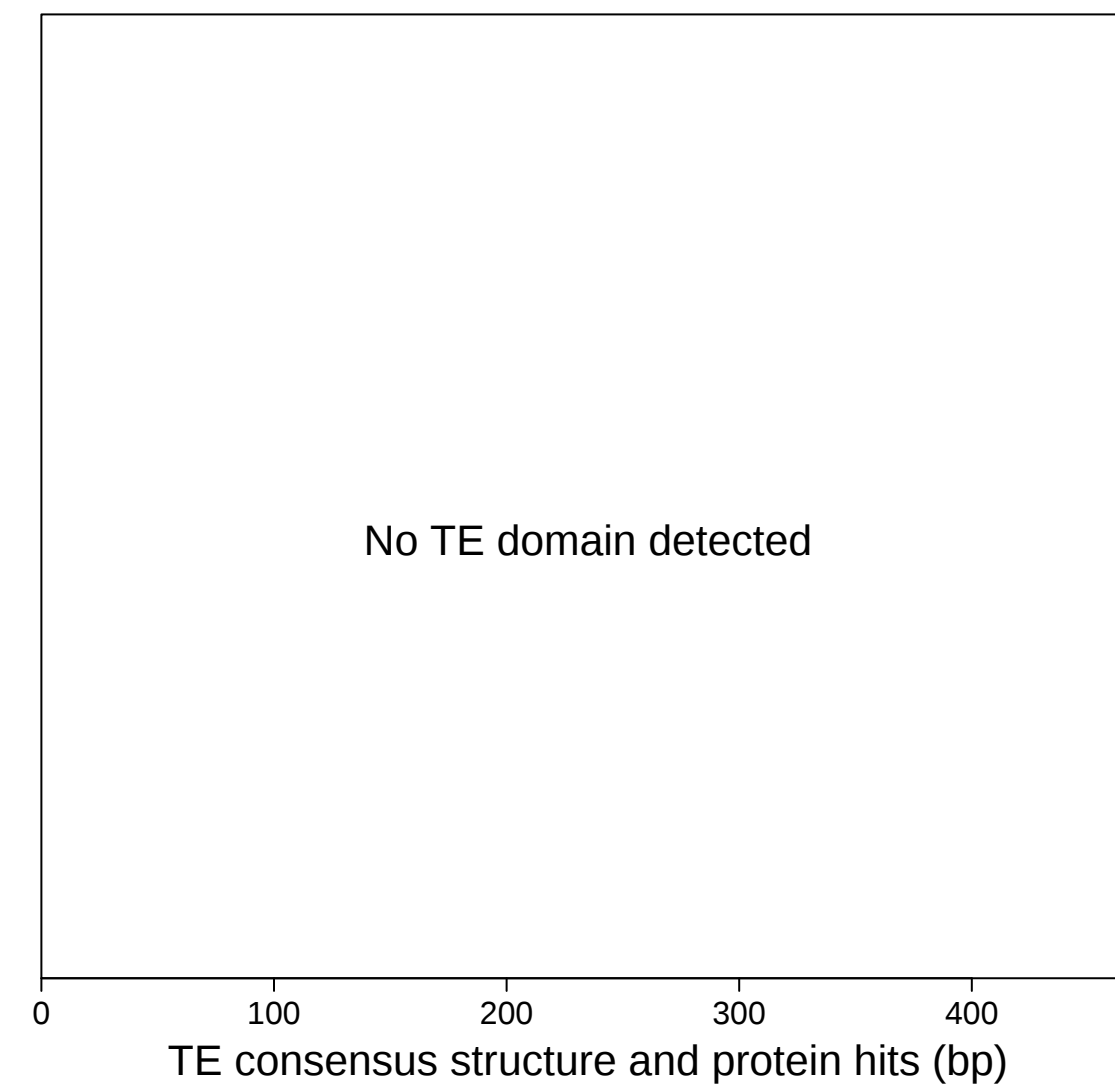
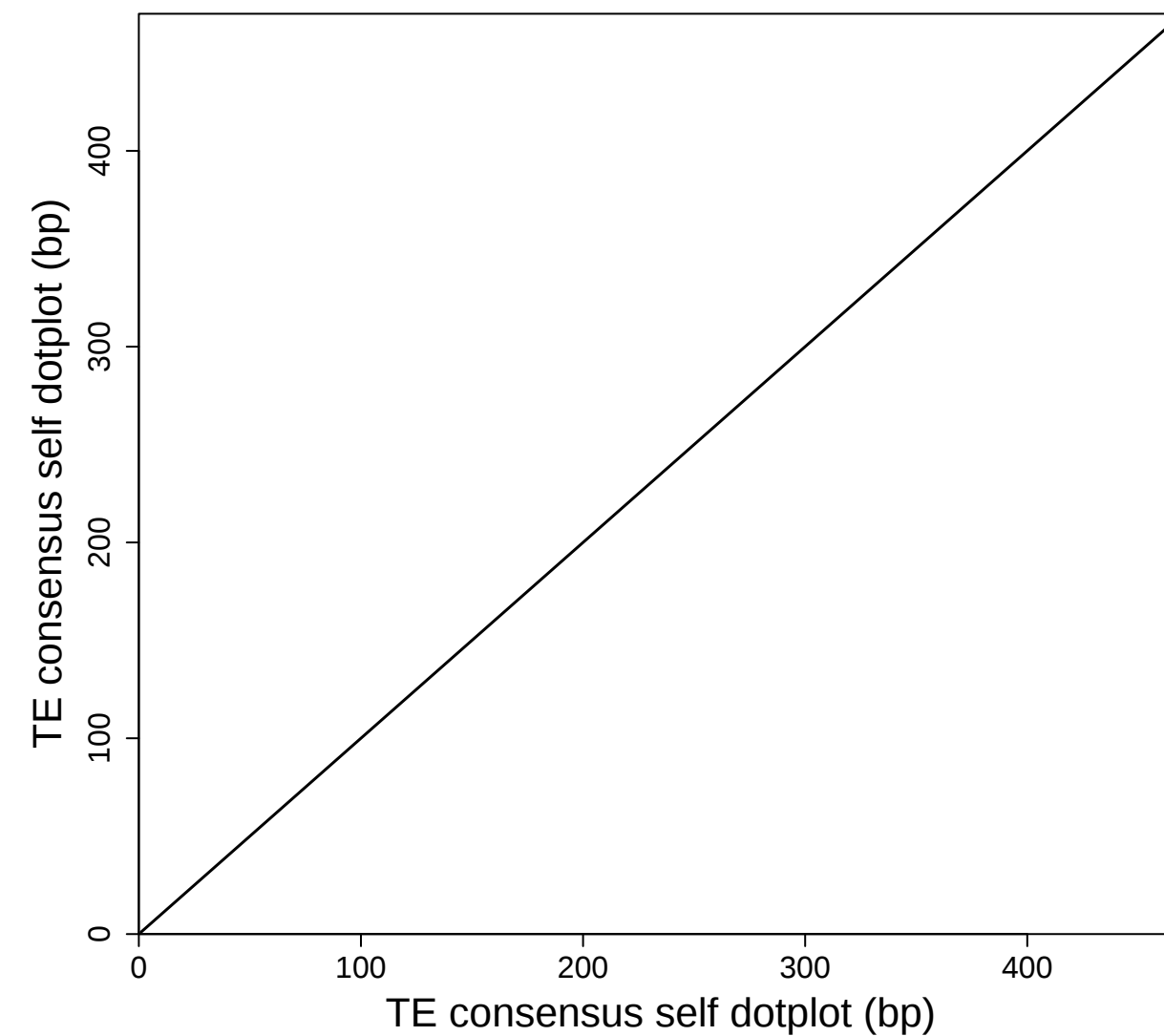
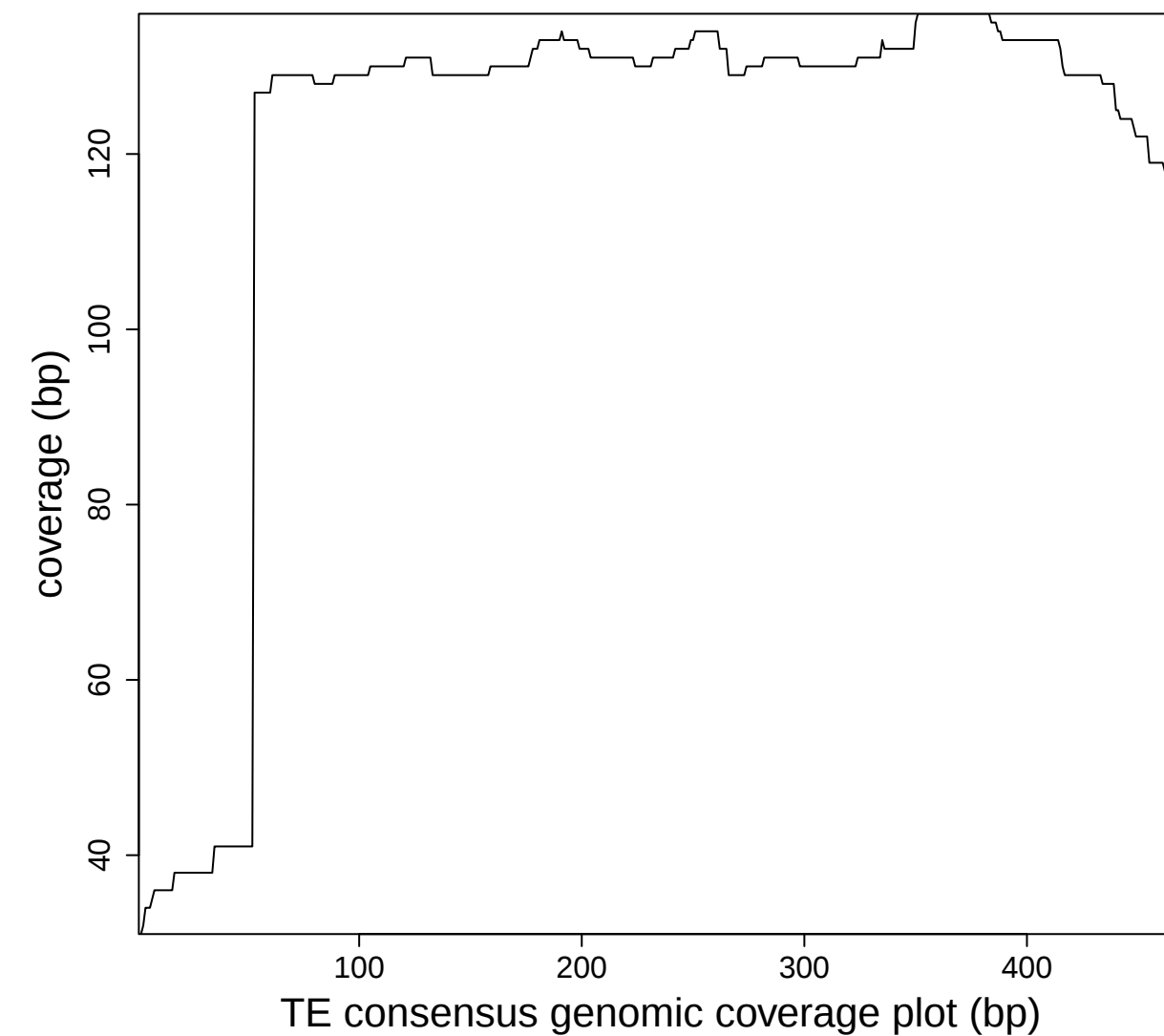
After TEtrimmer Extended plot Blue lines are boundaries



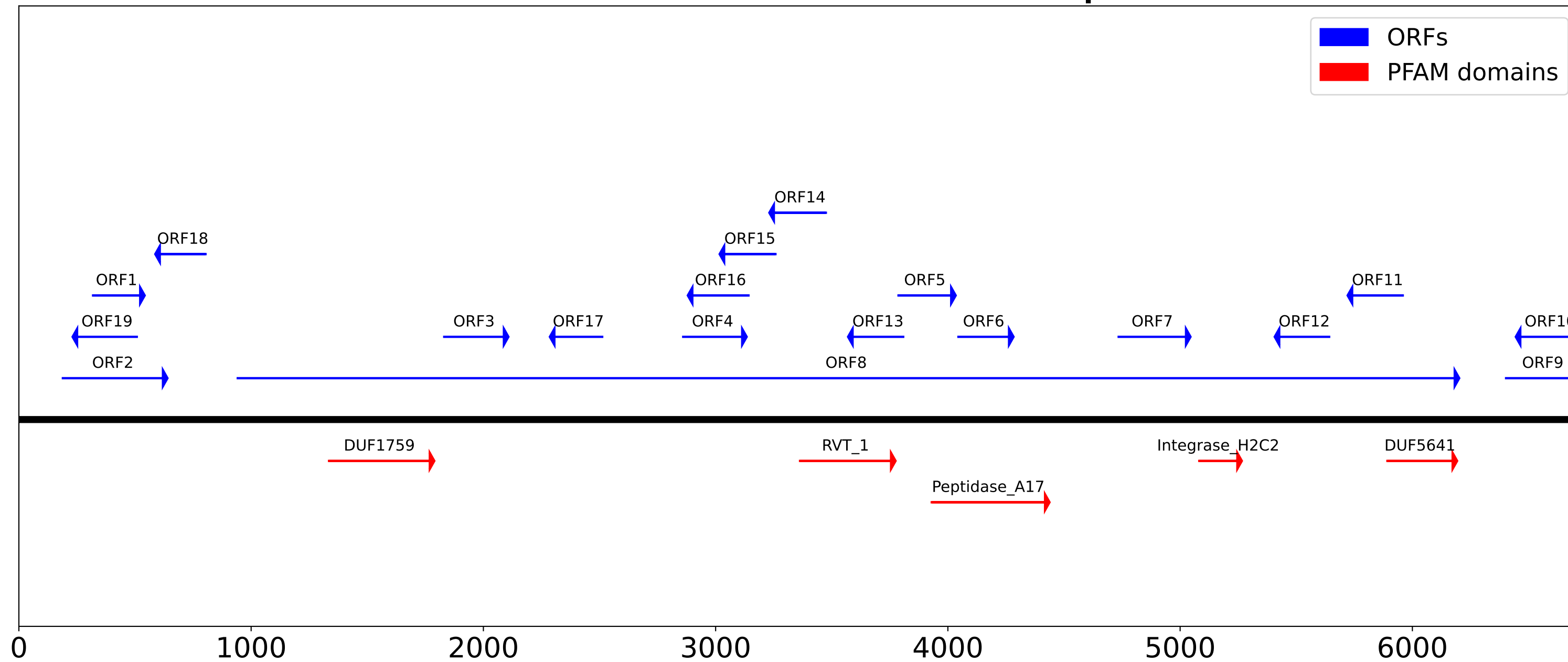
TE: rnd_5_family_2690
size: 470bp; fragments: 151; full length: 26 (≥ 423 bp)



Before TEtrimmer 470 bp



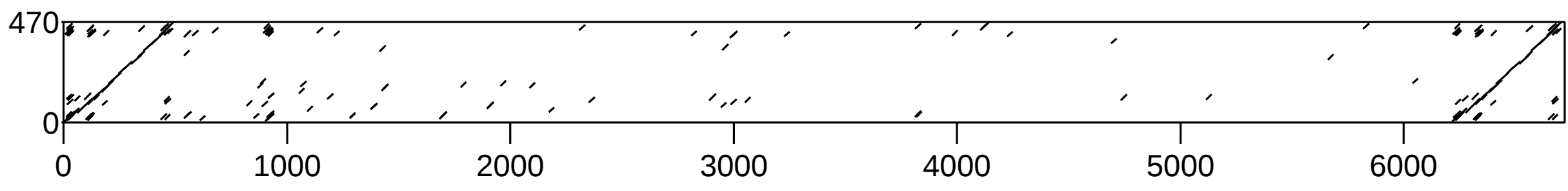
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)