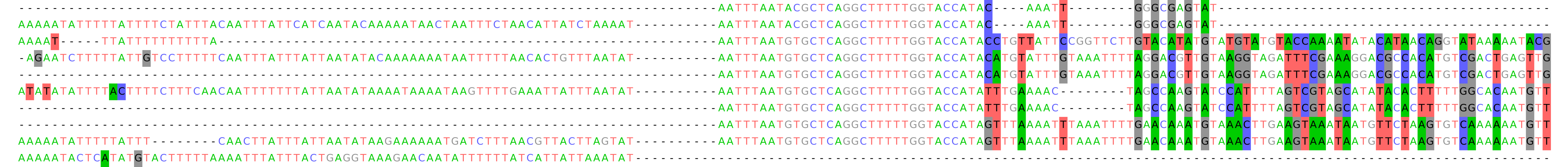


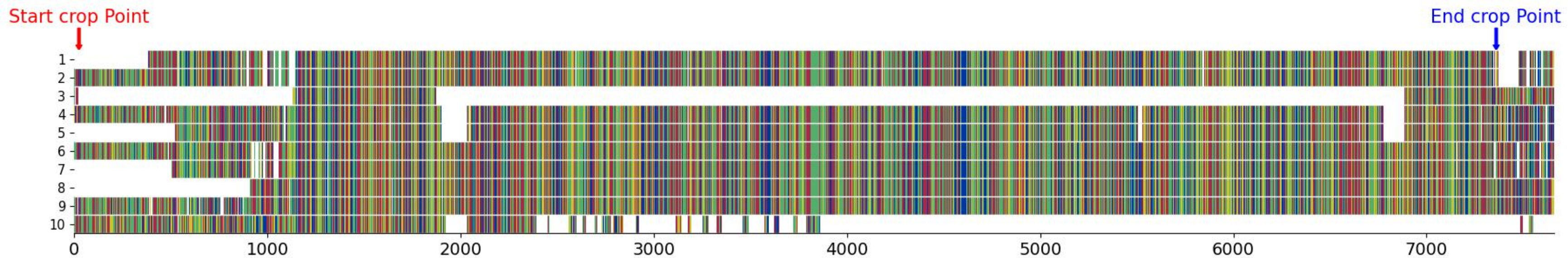
Start crop Point



MSA length = 7338

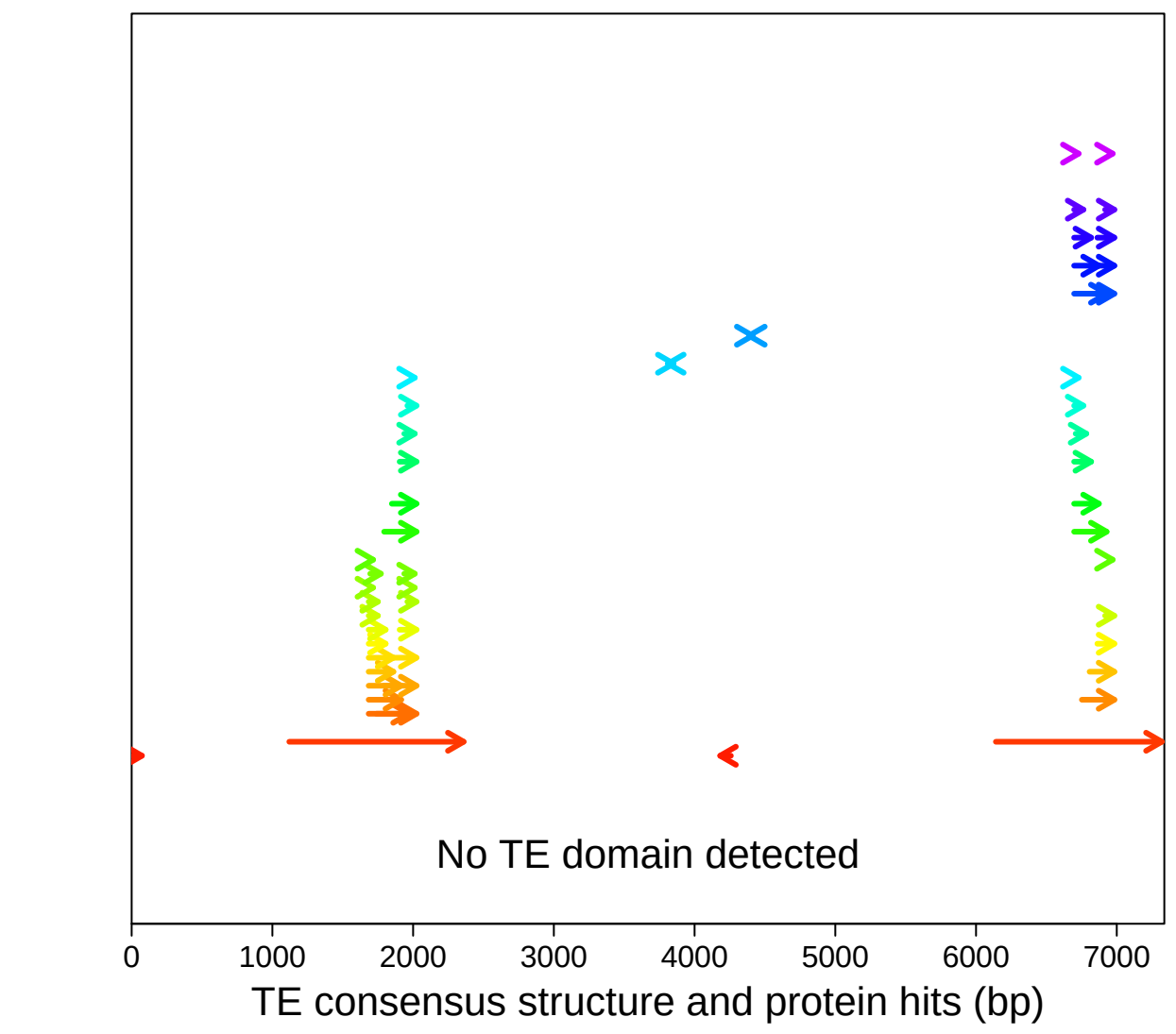
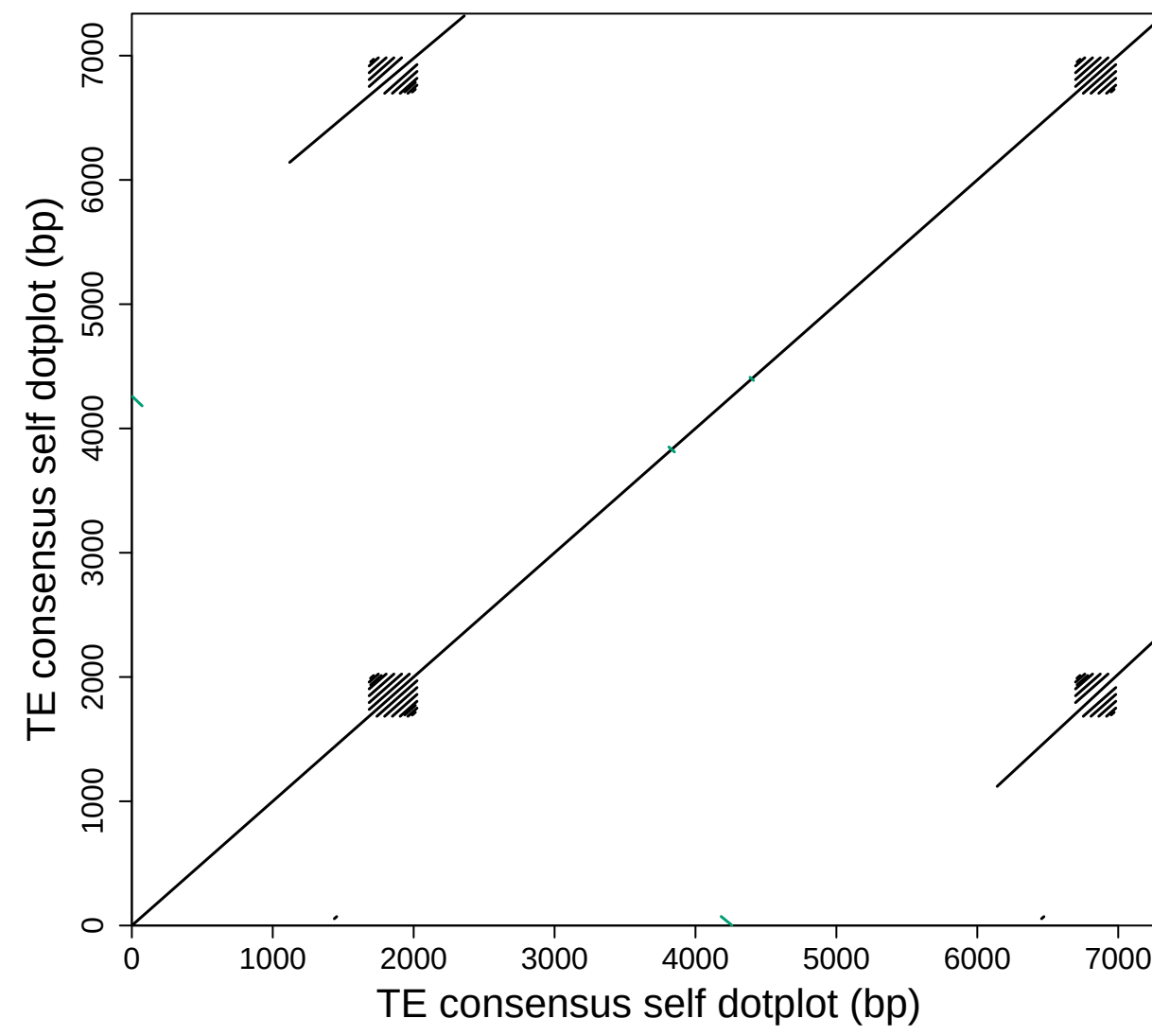
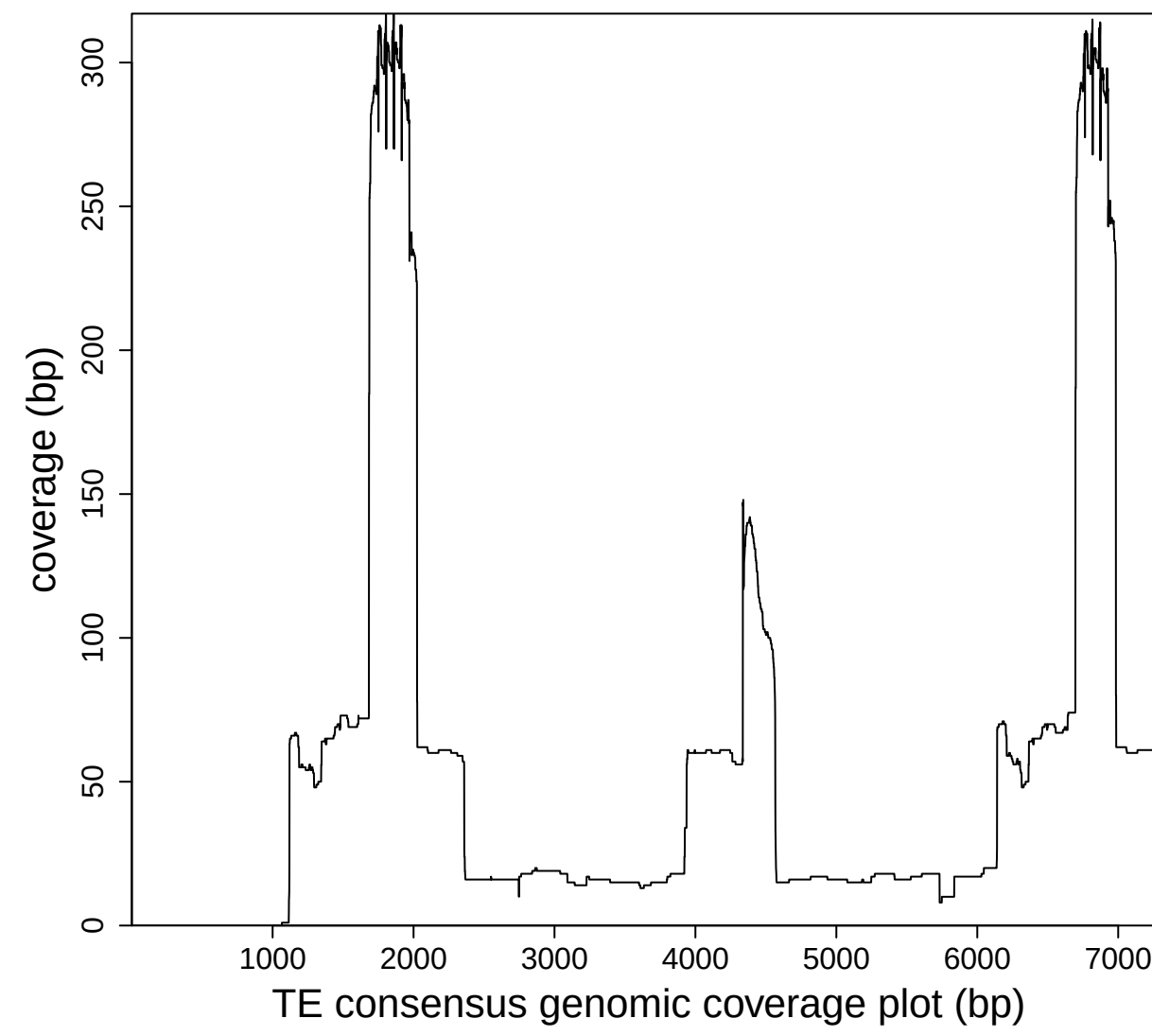
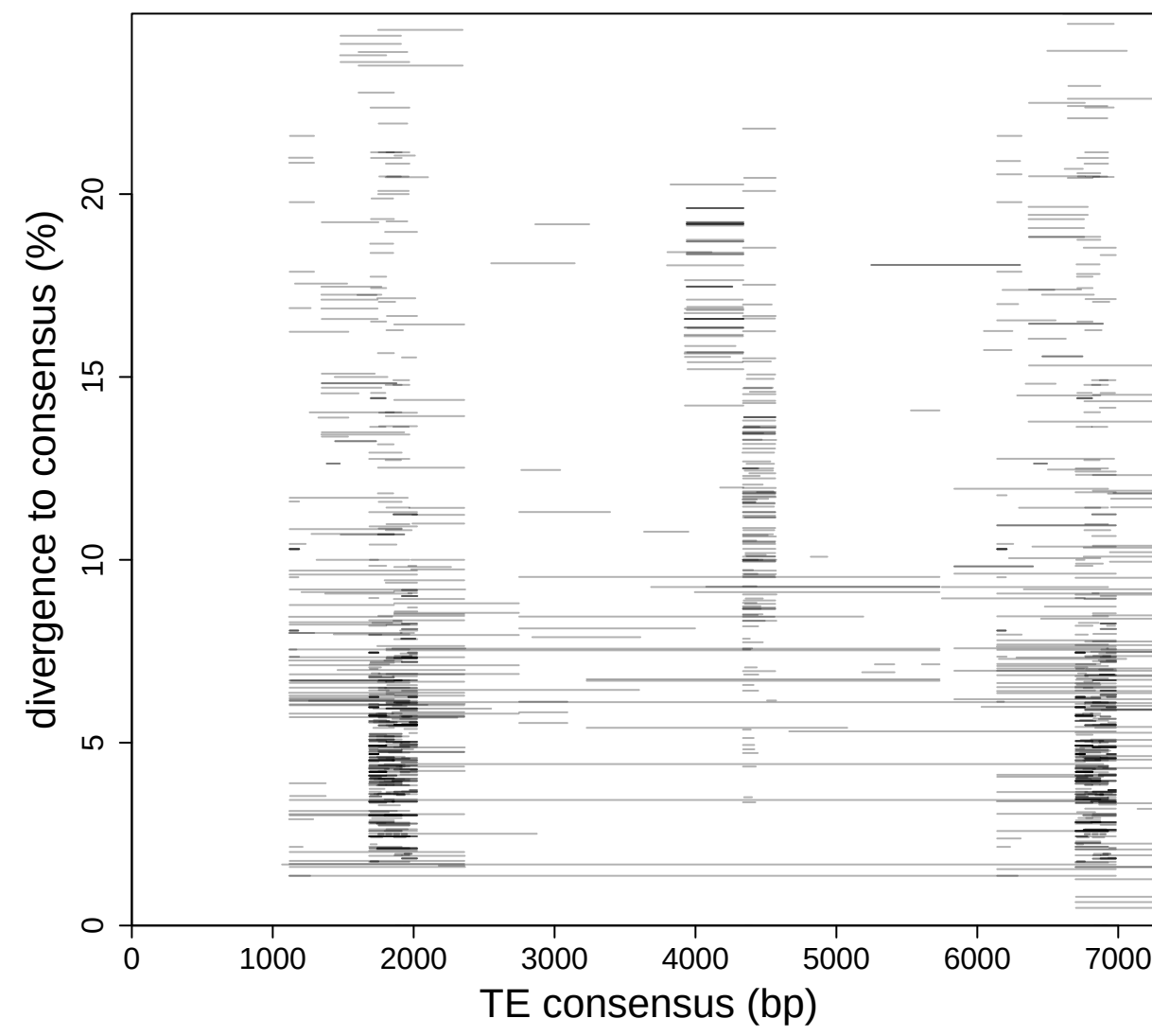
End crop Point





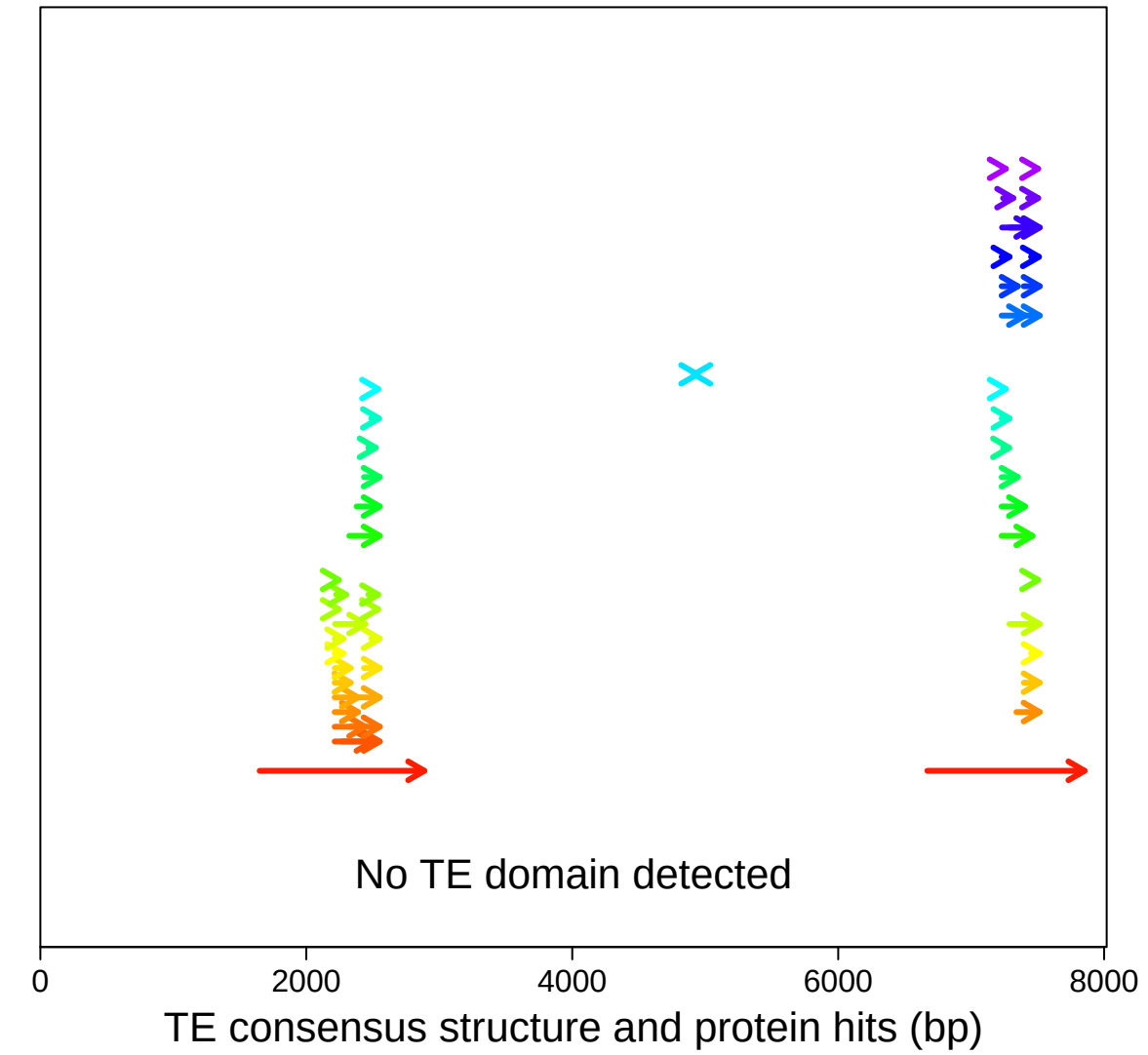
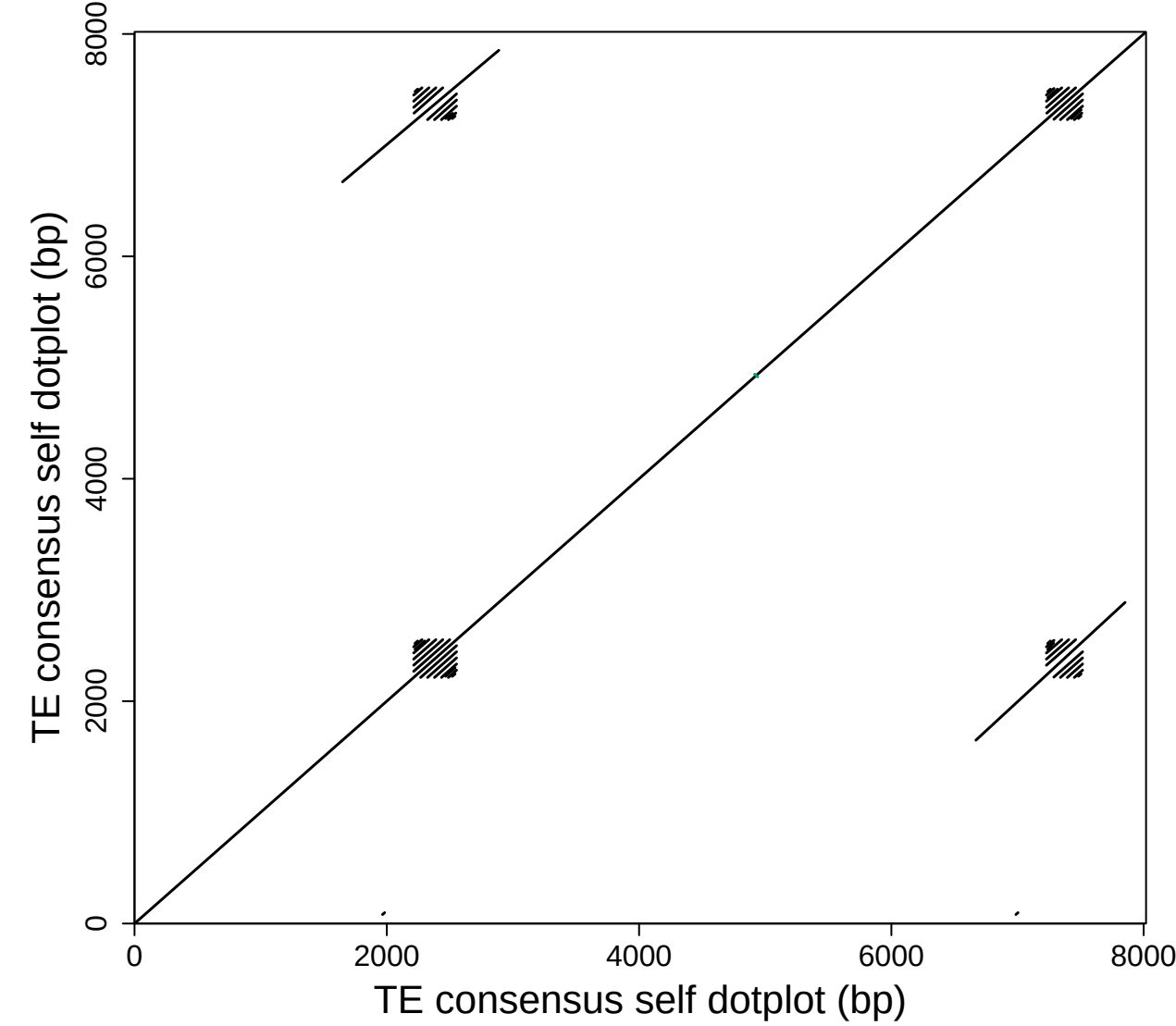
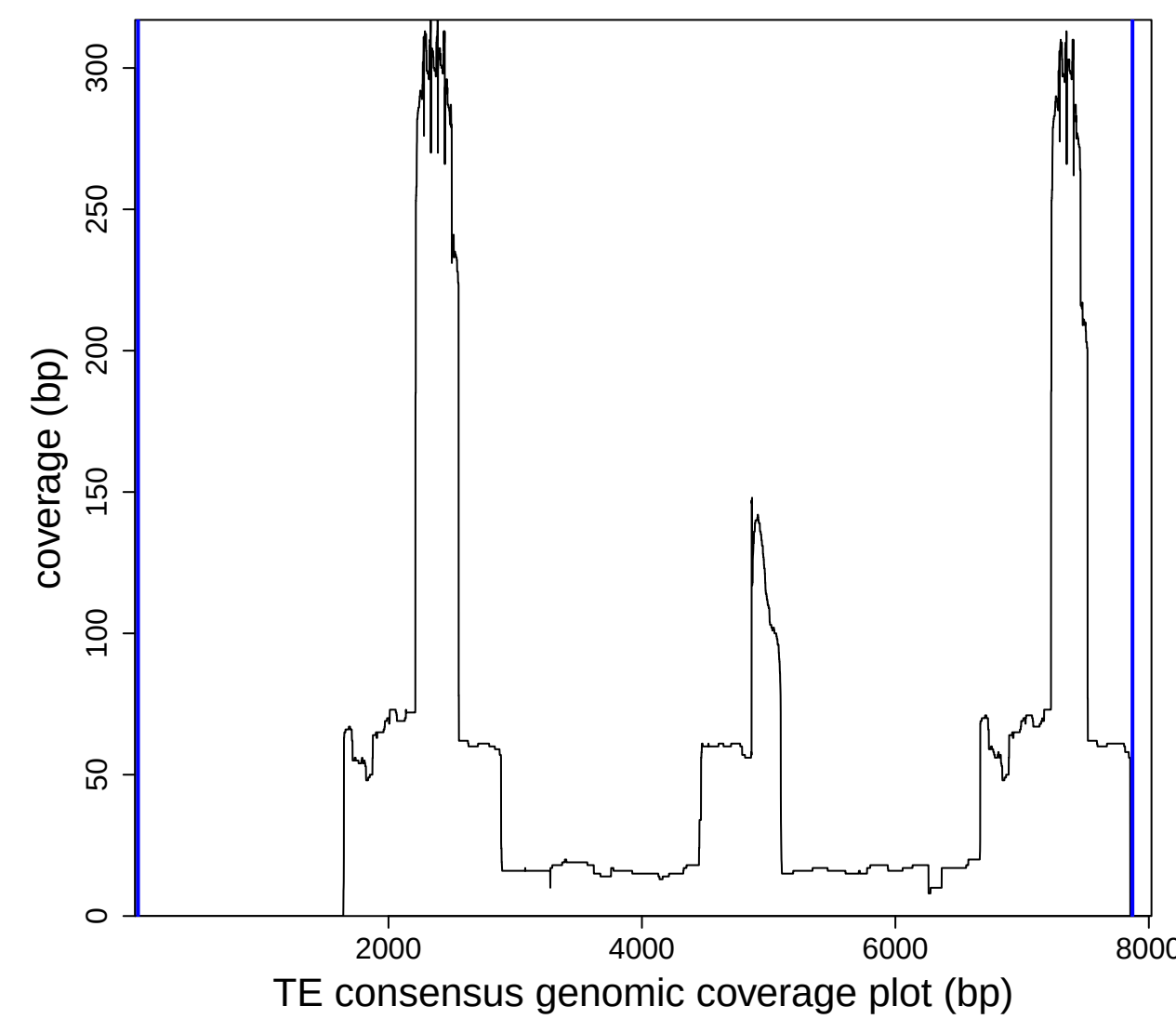
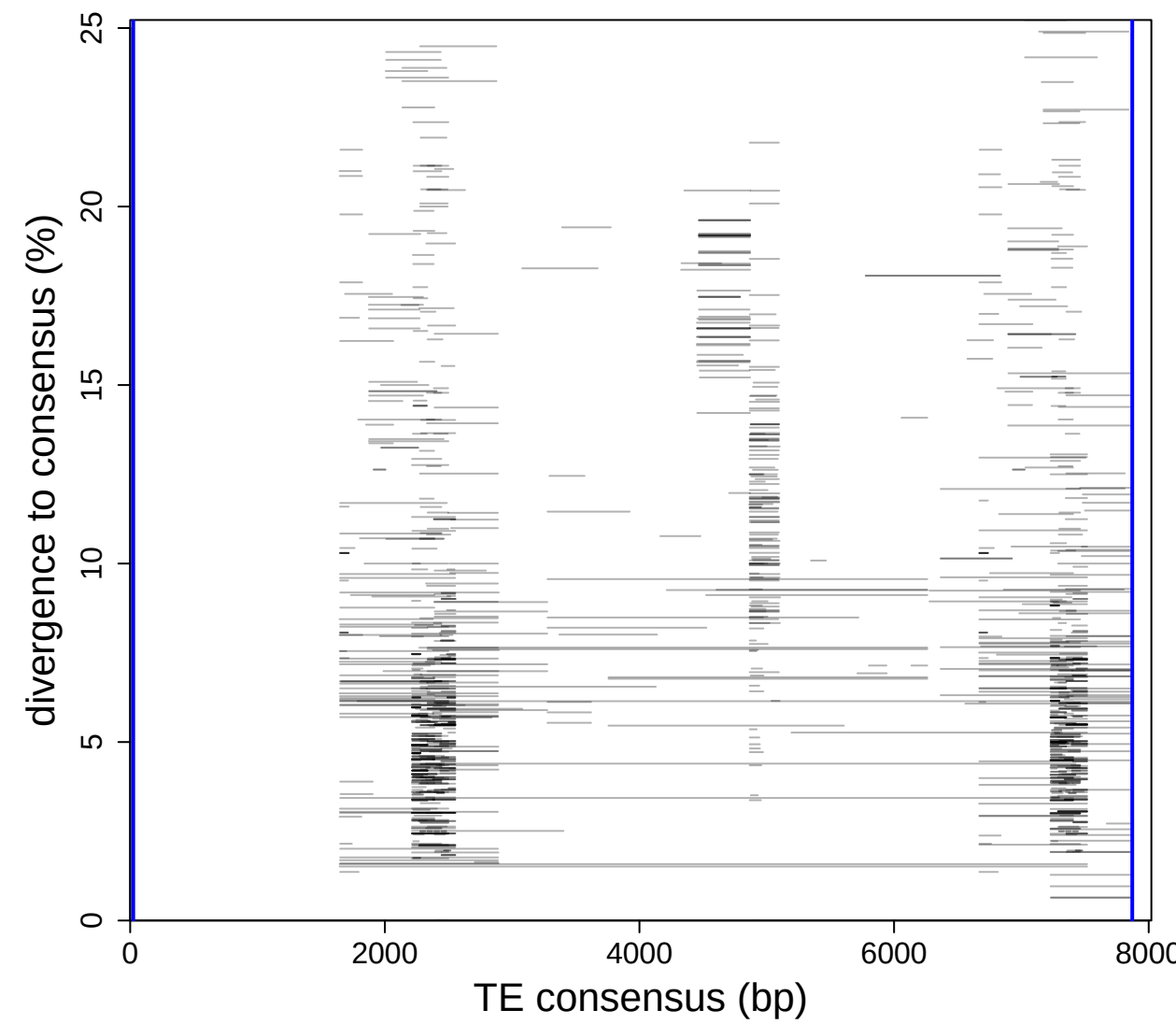
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa al
size: 7338bp; fragments: 1374; full length: 0 (>=6604.2bp)

After TEtrimmer 7338 bp



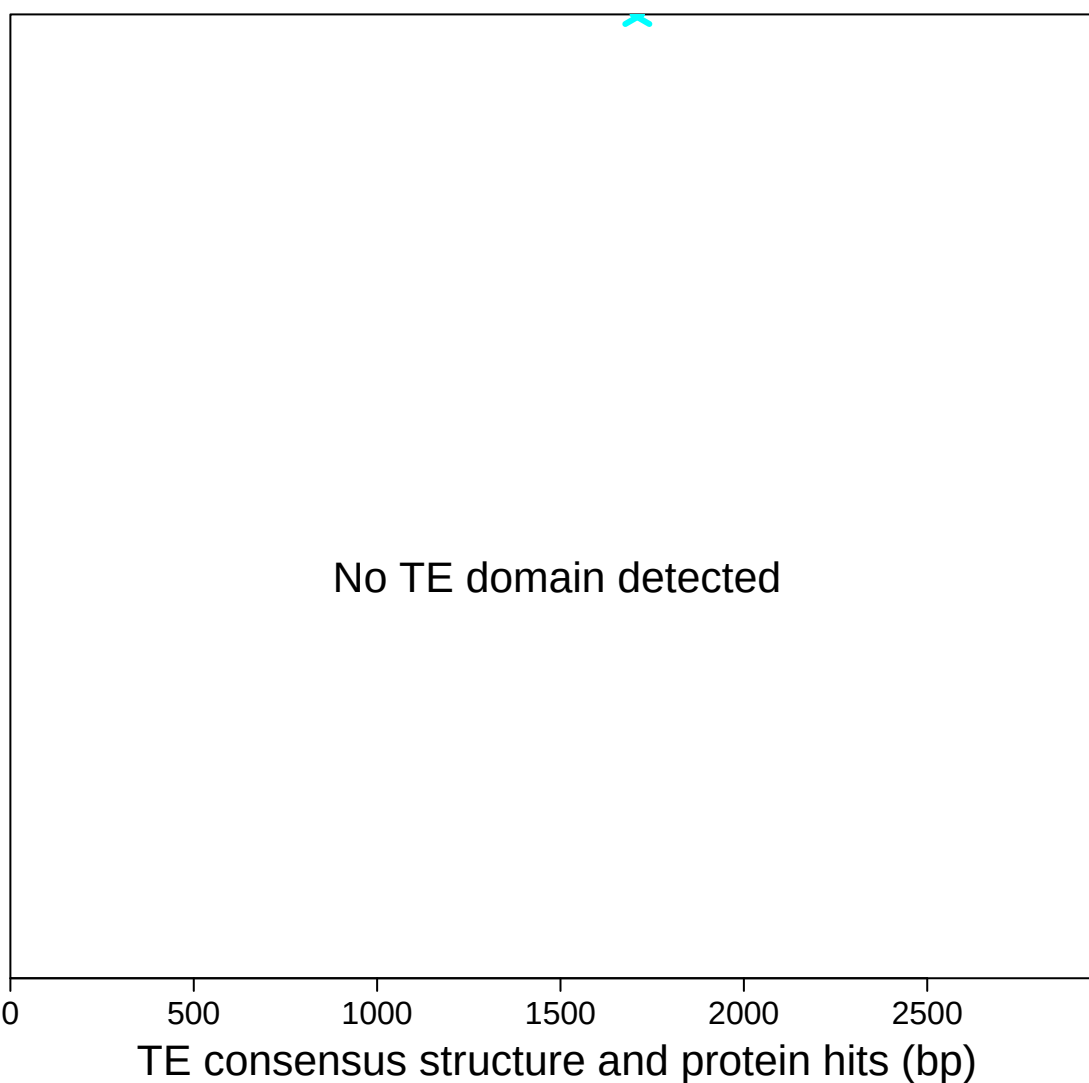
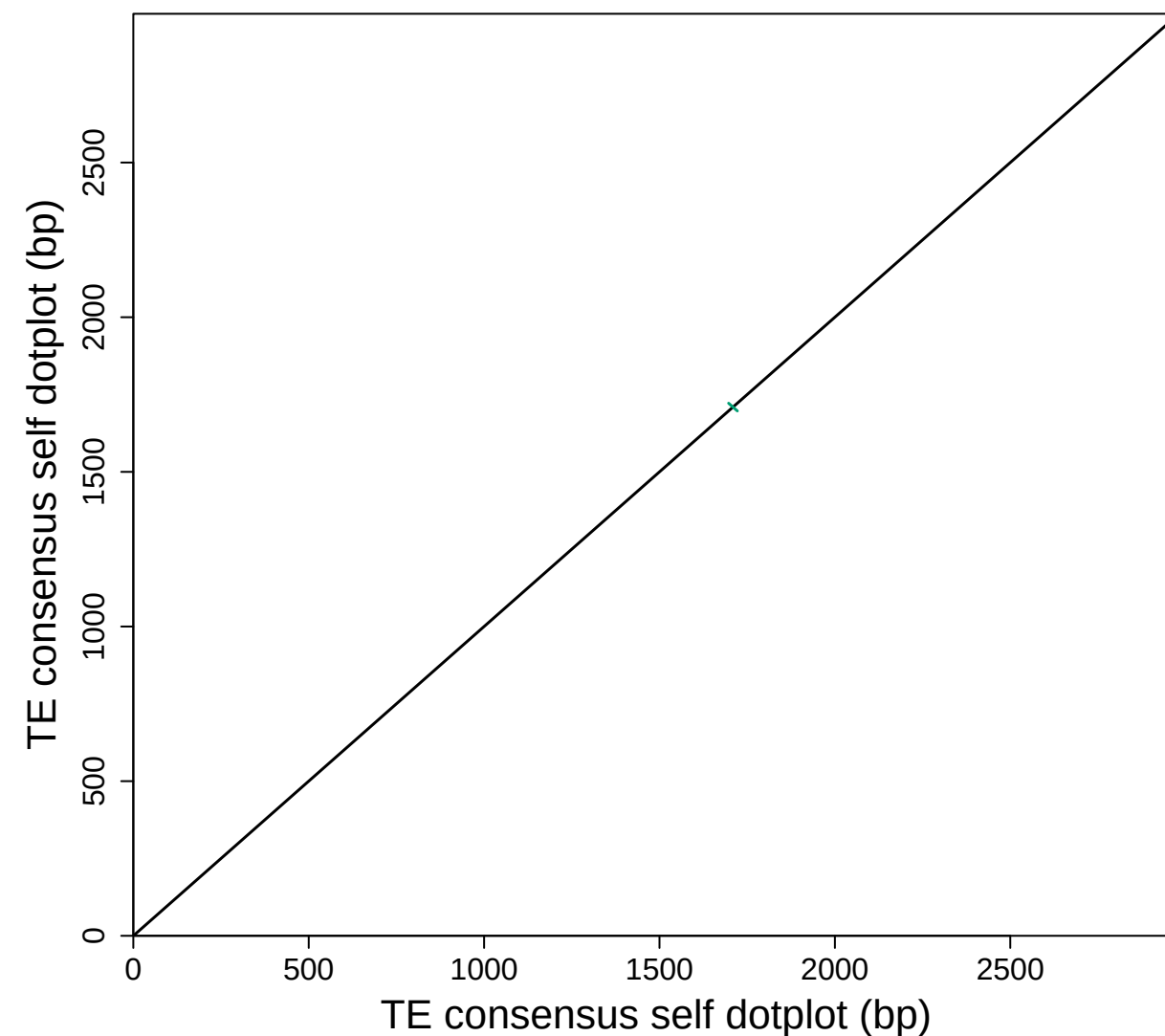
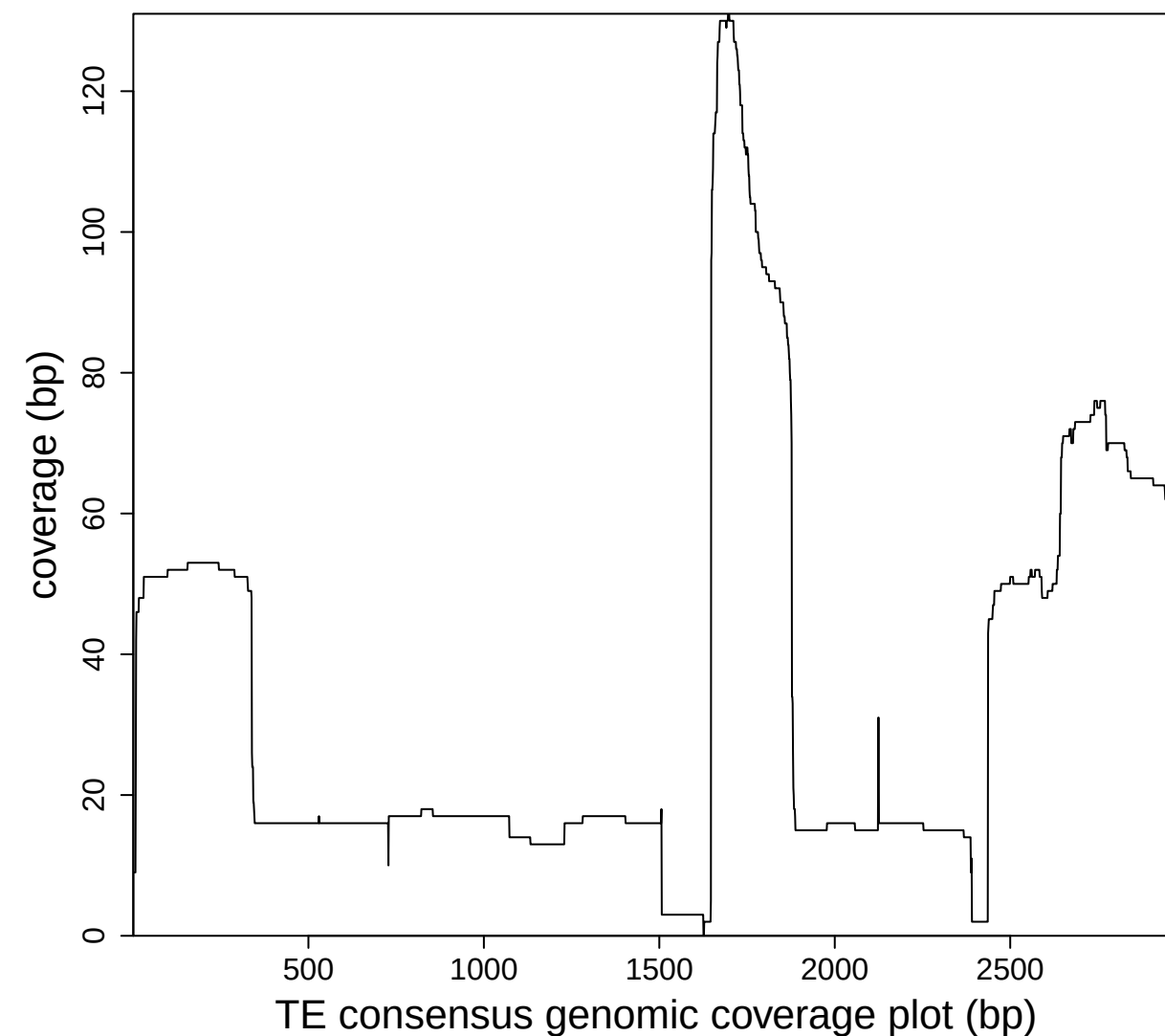
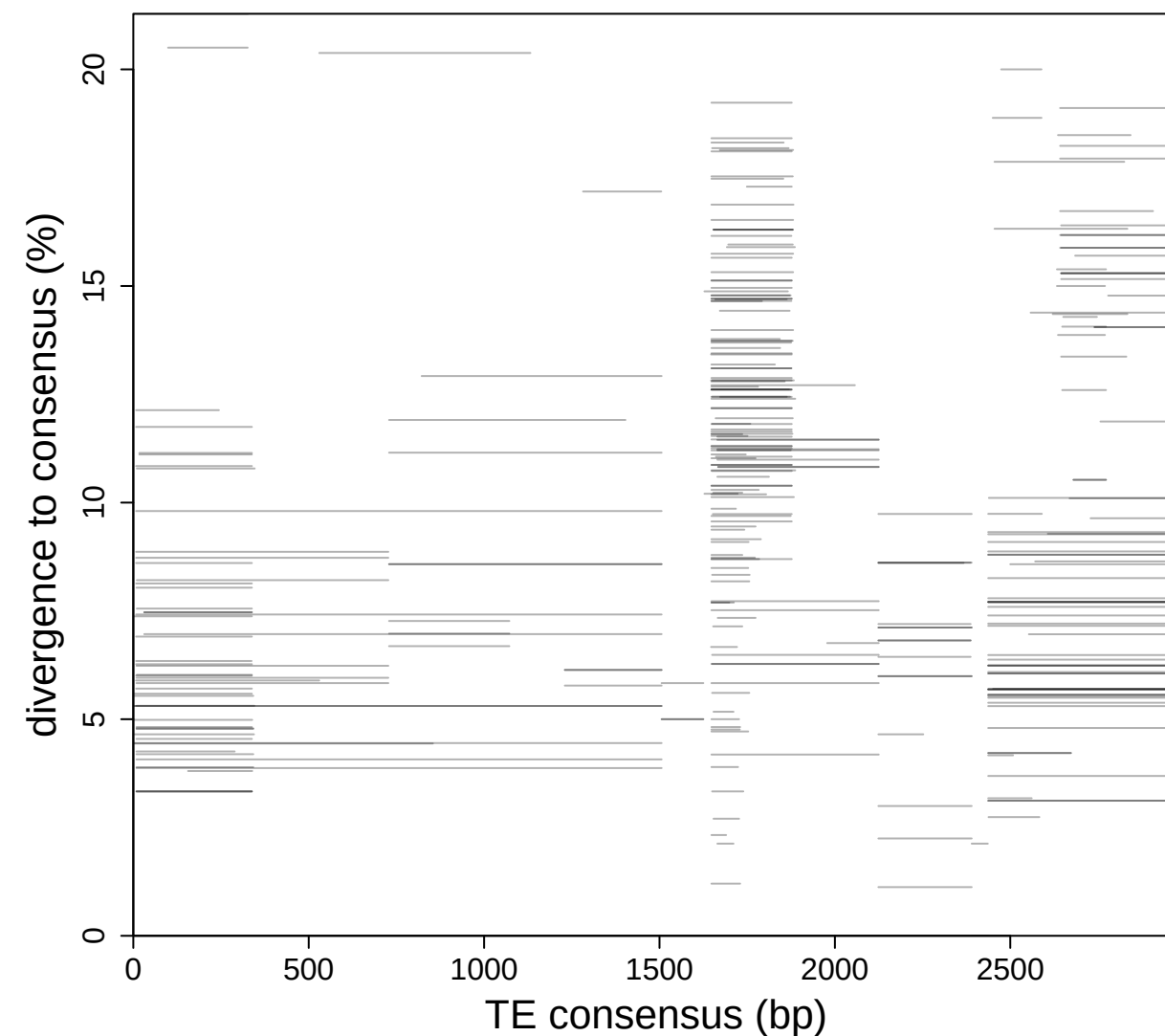
bed_g_1.bed fm_1.bed 0 0 n.bed_g_1.bed fm_2.bed 0 0 bclr
size: 8021bp; fragments: 1341; full length: 0 (>=7218.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

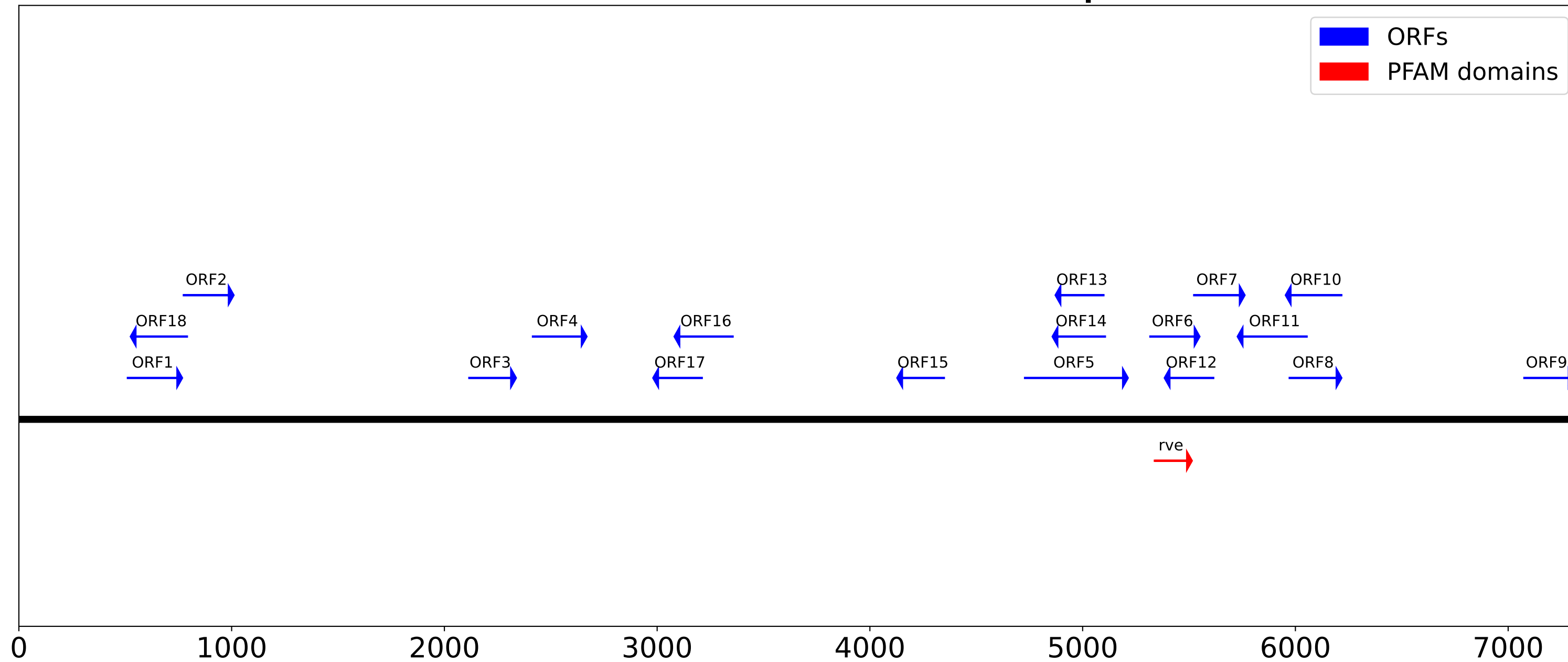


TE: ltr_1_family_653
size: 2981bp; fragments: 307; full length: 0 (>=2682.9bp)

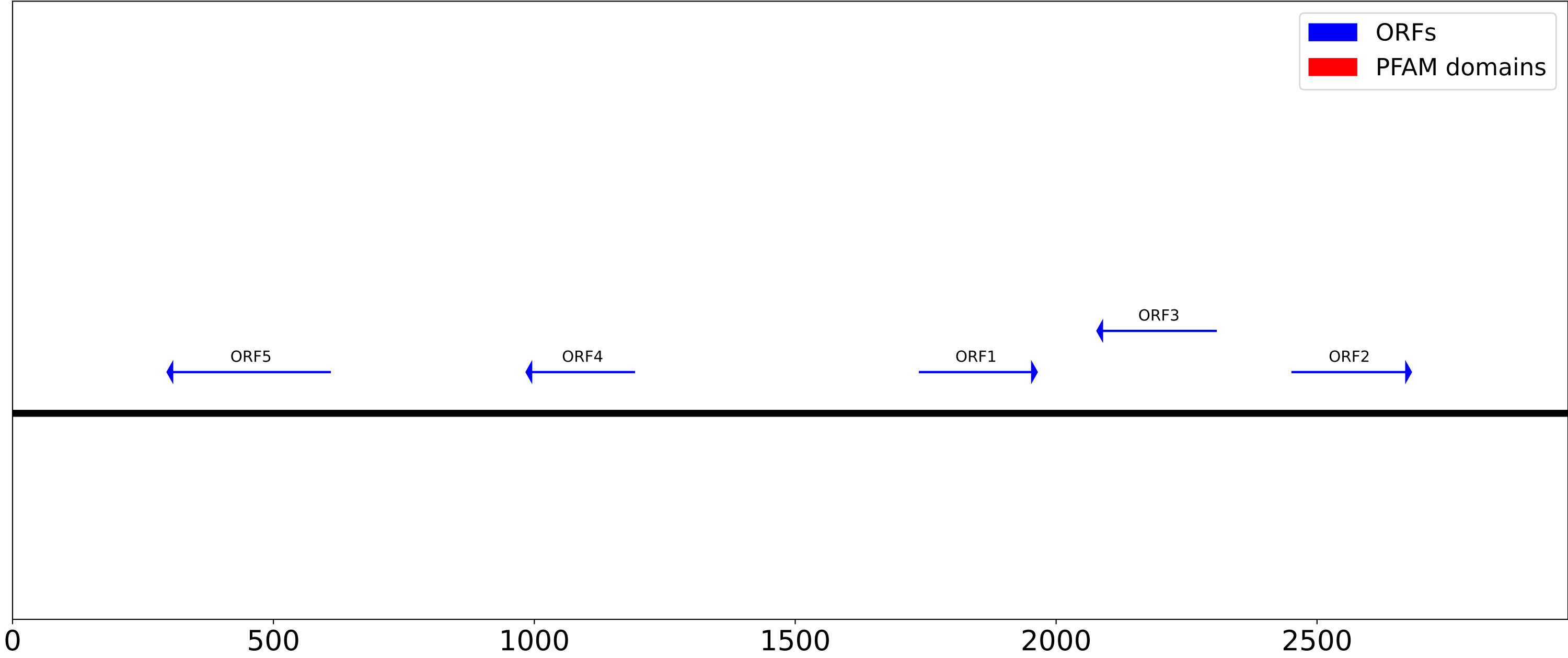
Before TEtrimmer 2981 bp



After TEtrimmer ORF and PFAM domain plot



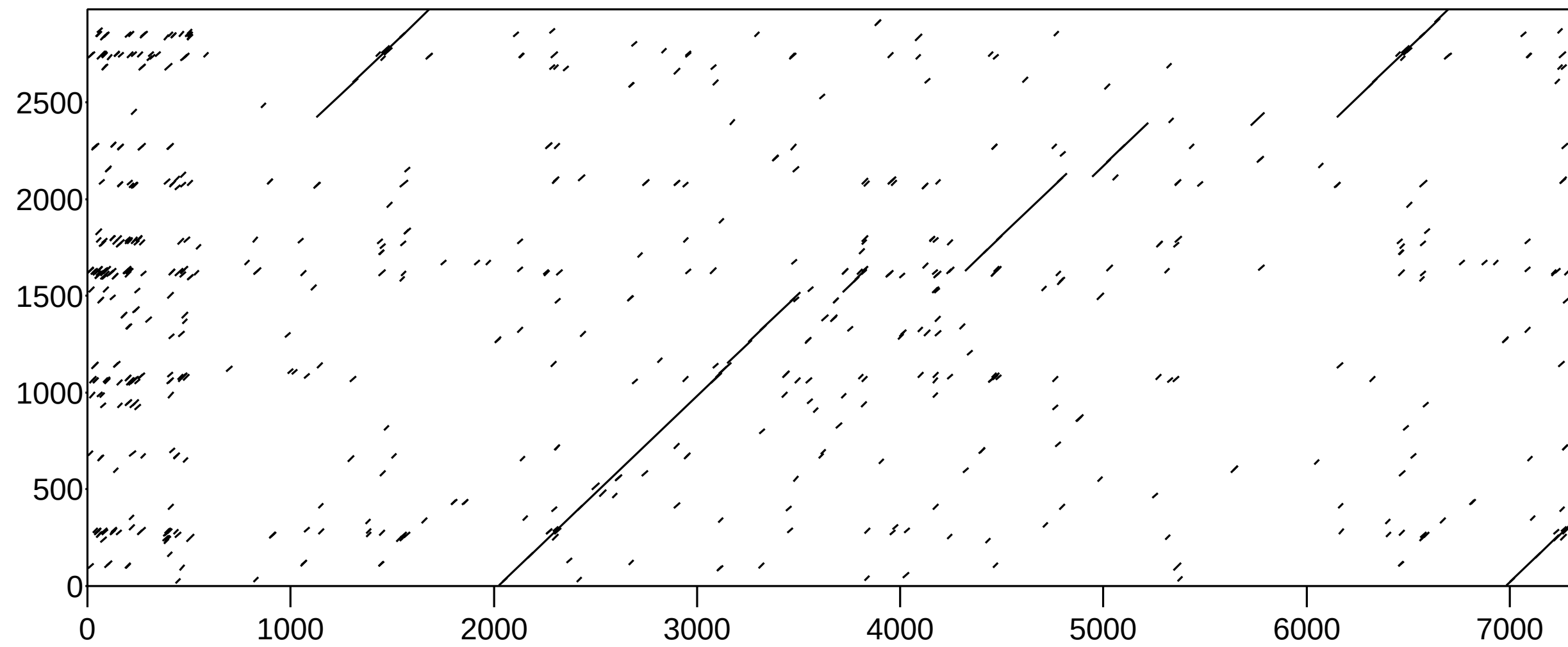
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)