

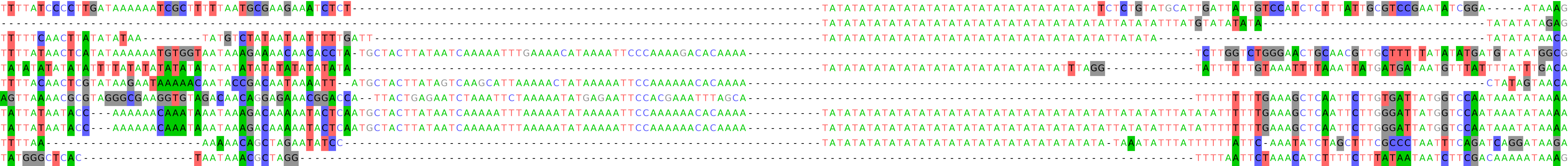
Start crop Point



End crop Point

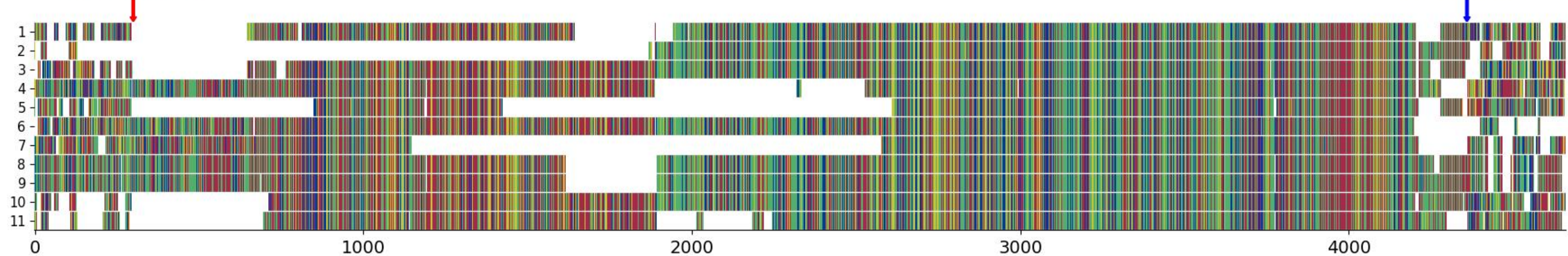


MSA length = 4061



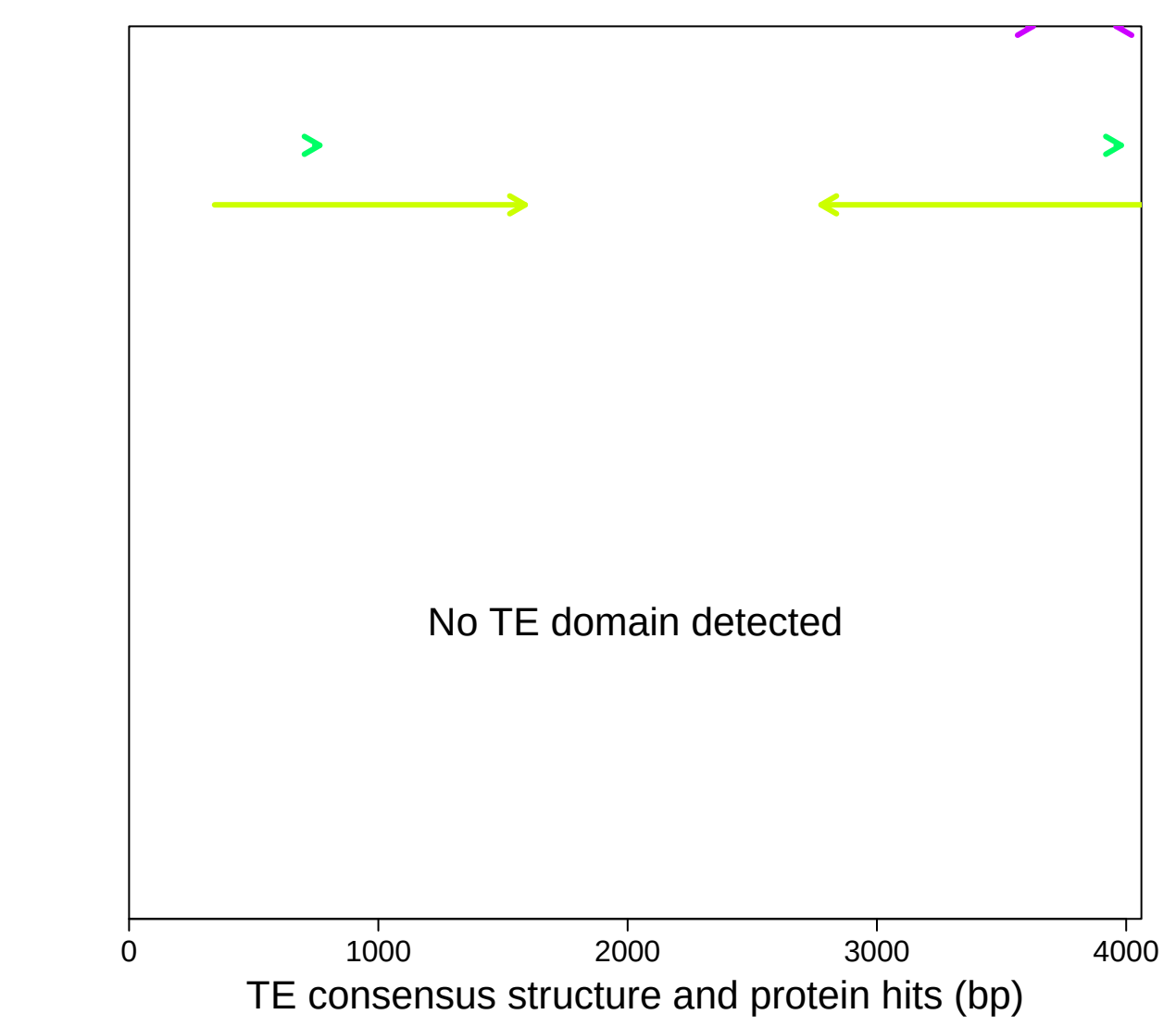
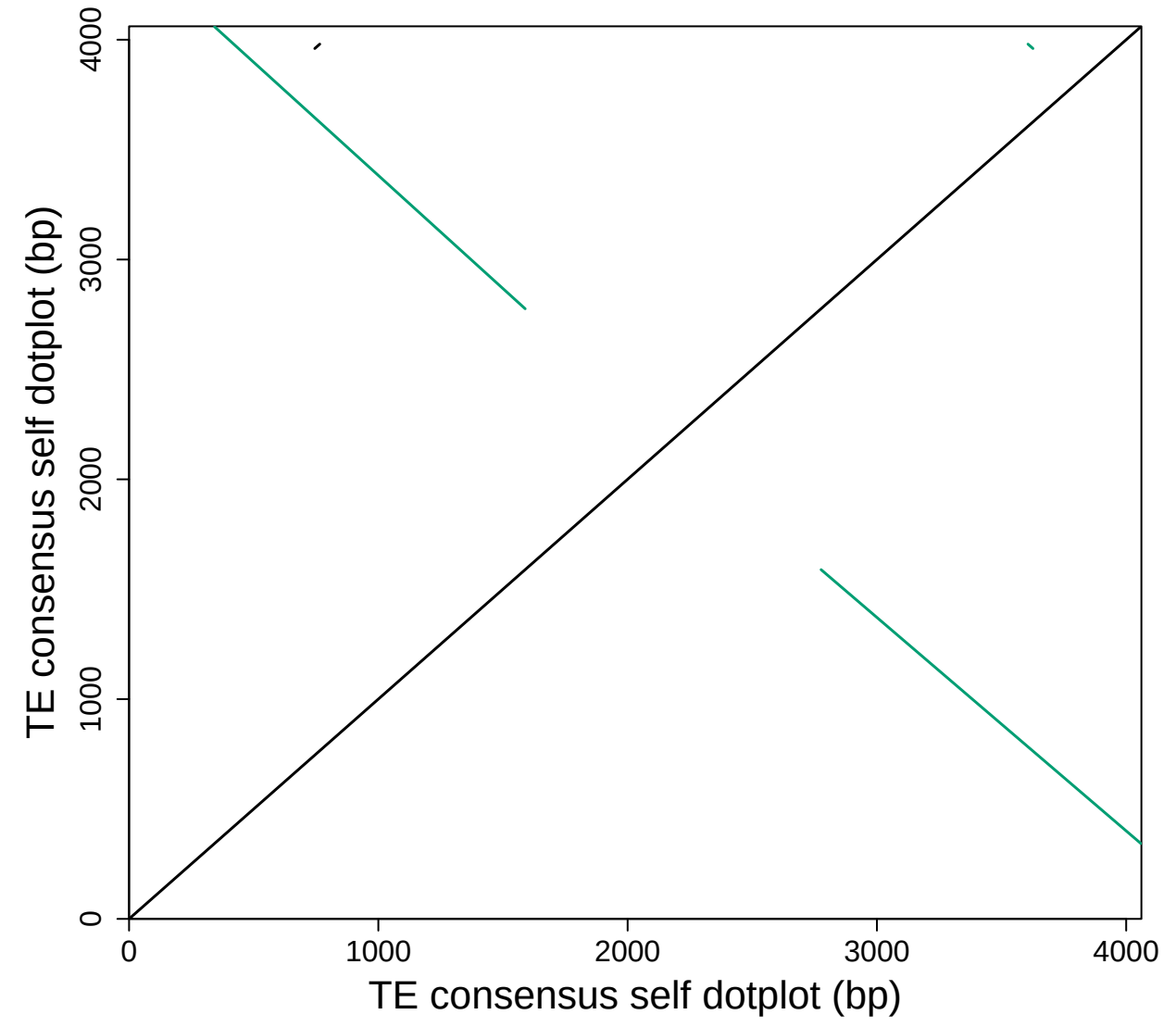
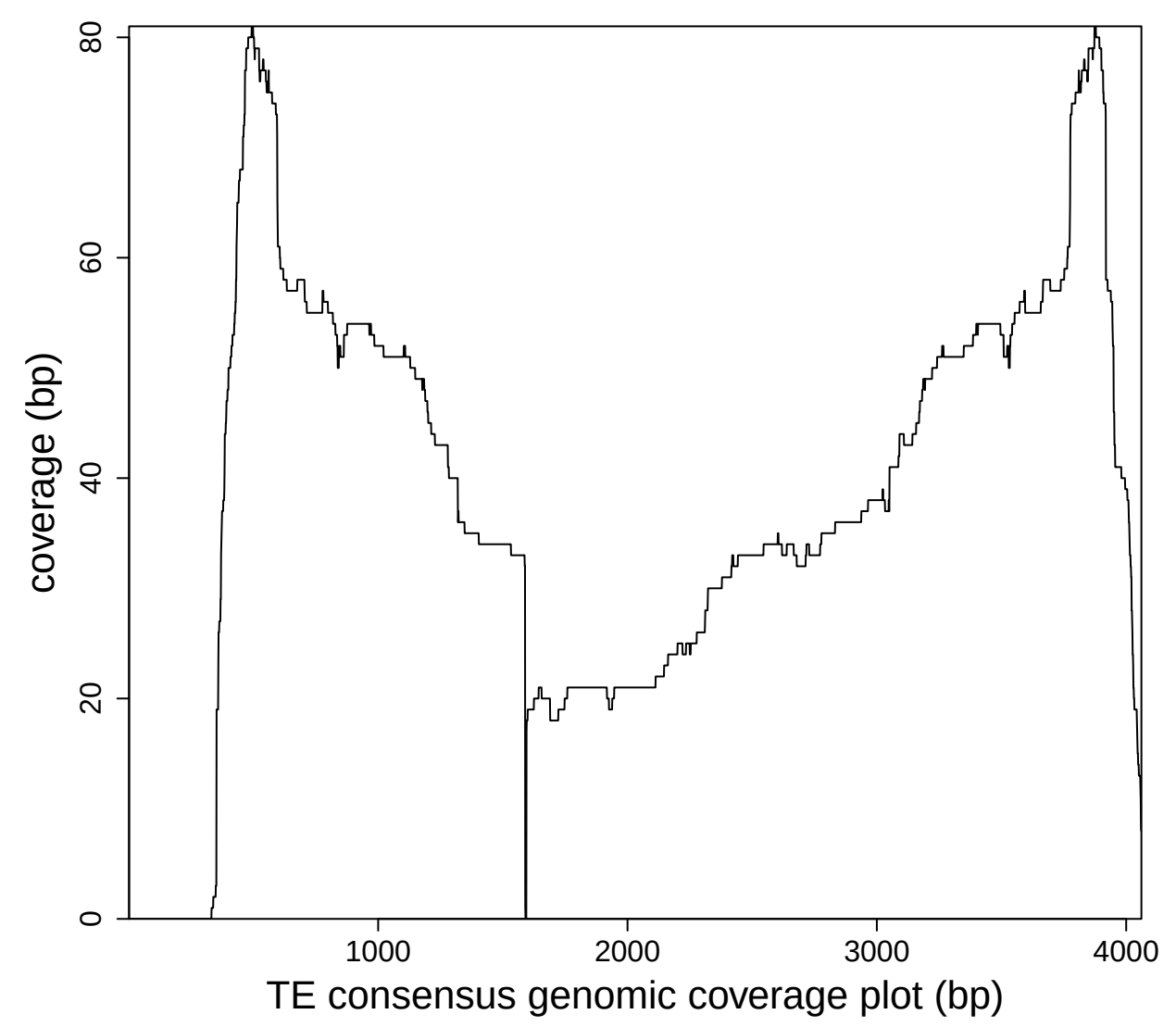
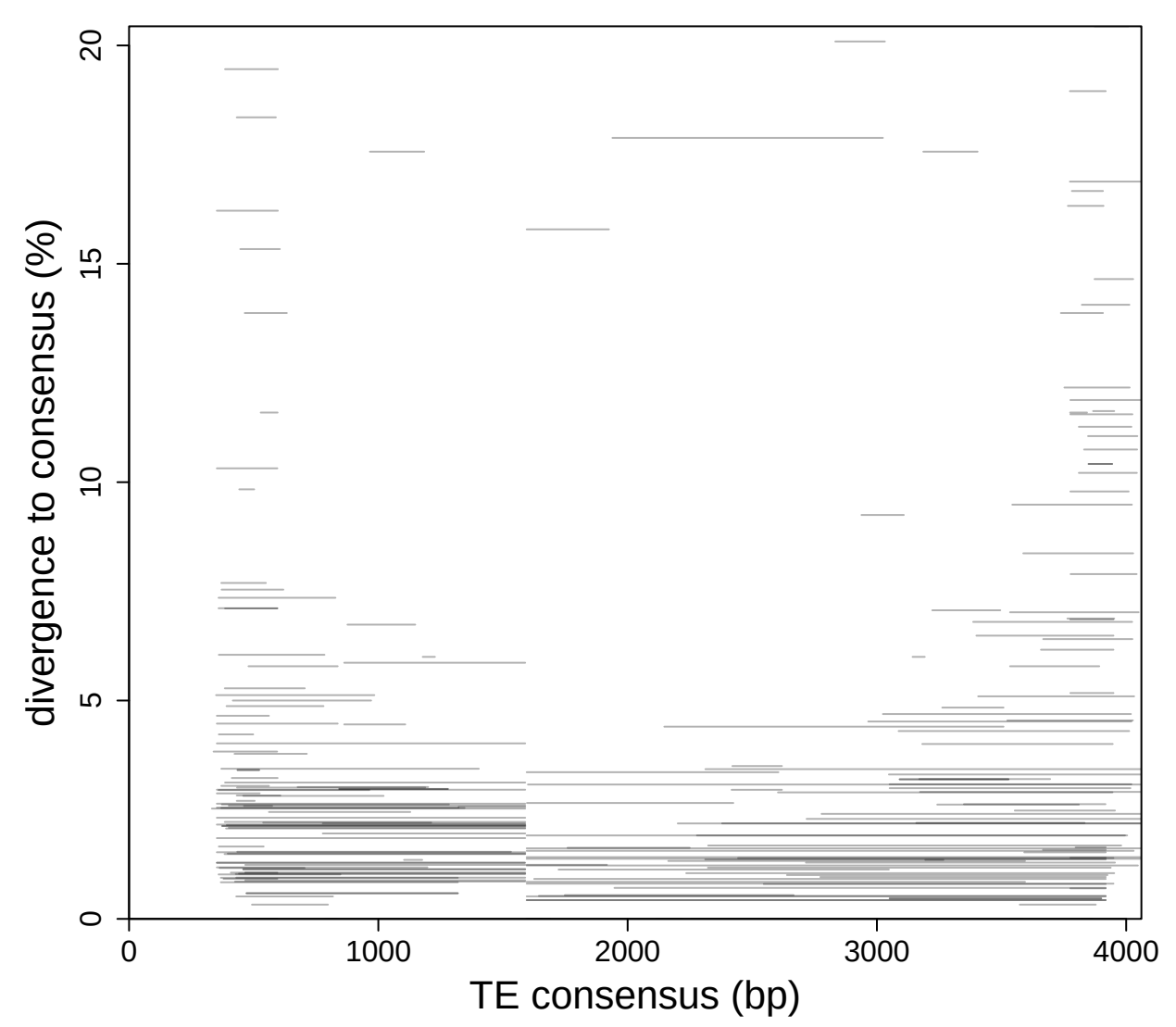
Start crop Point

End crop Point



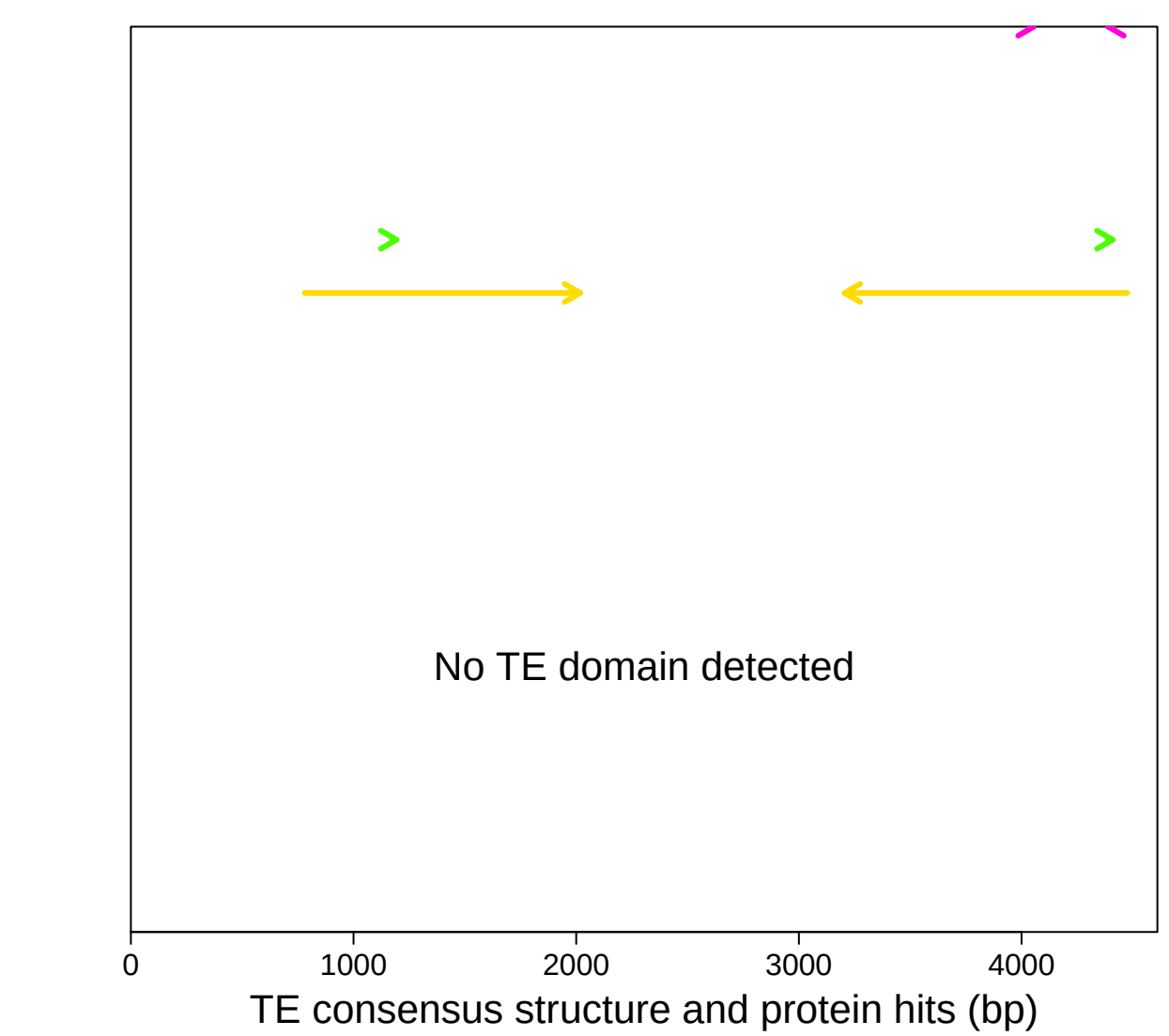
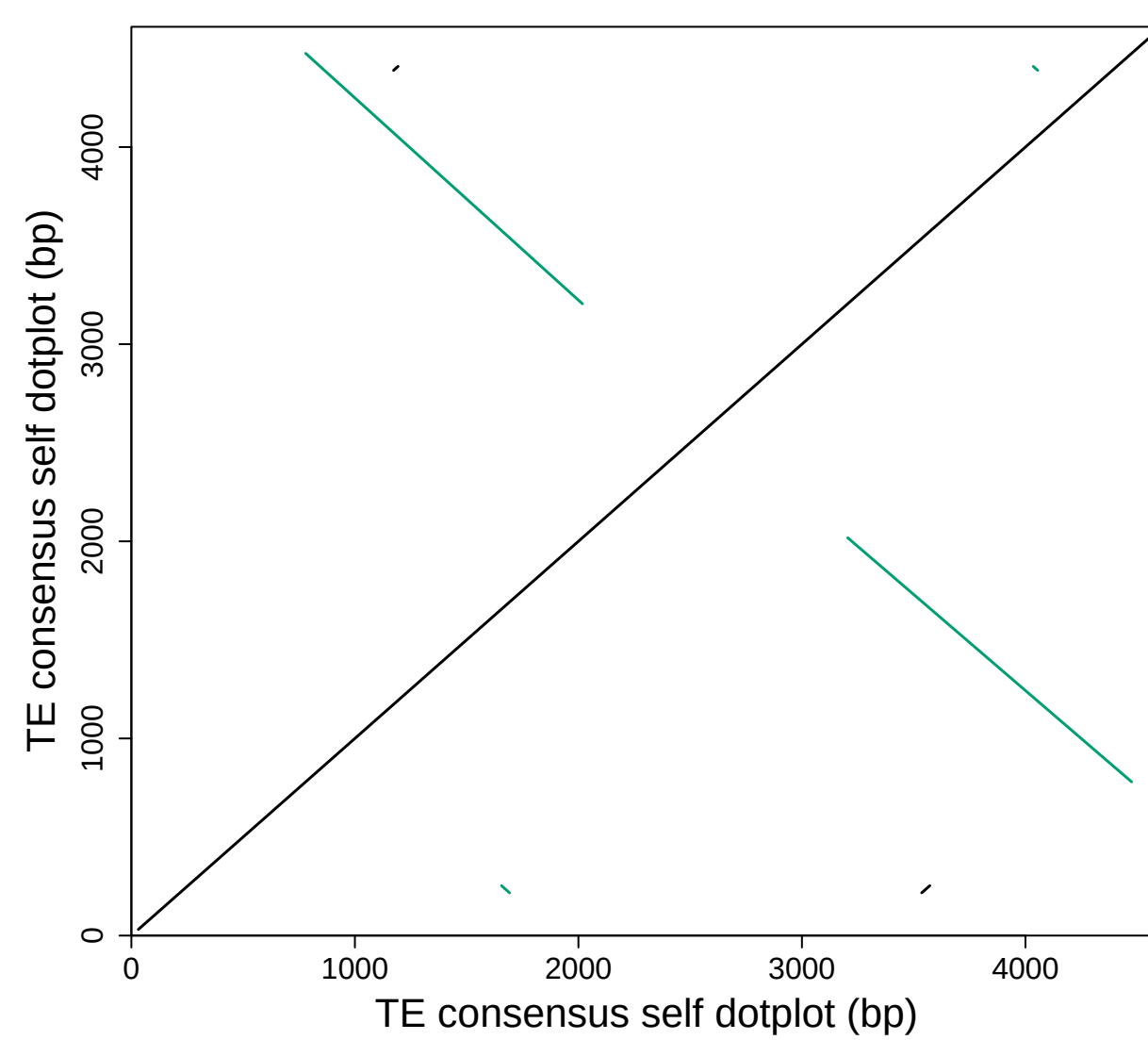
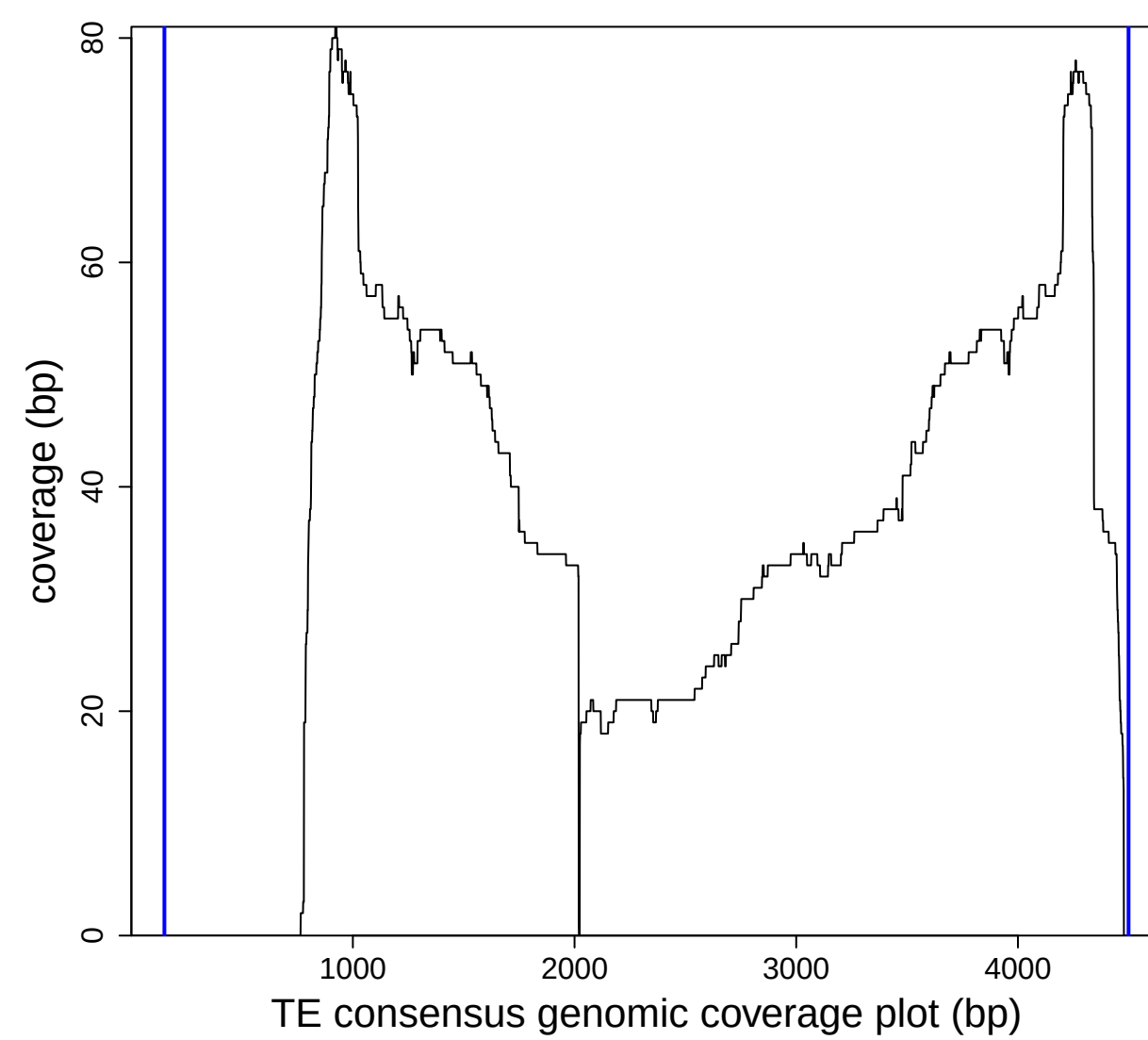
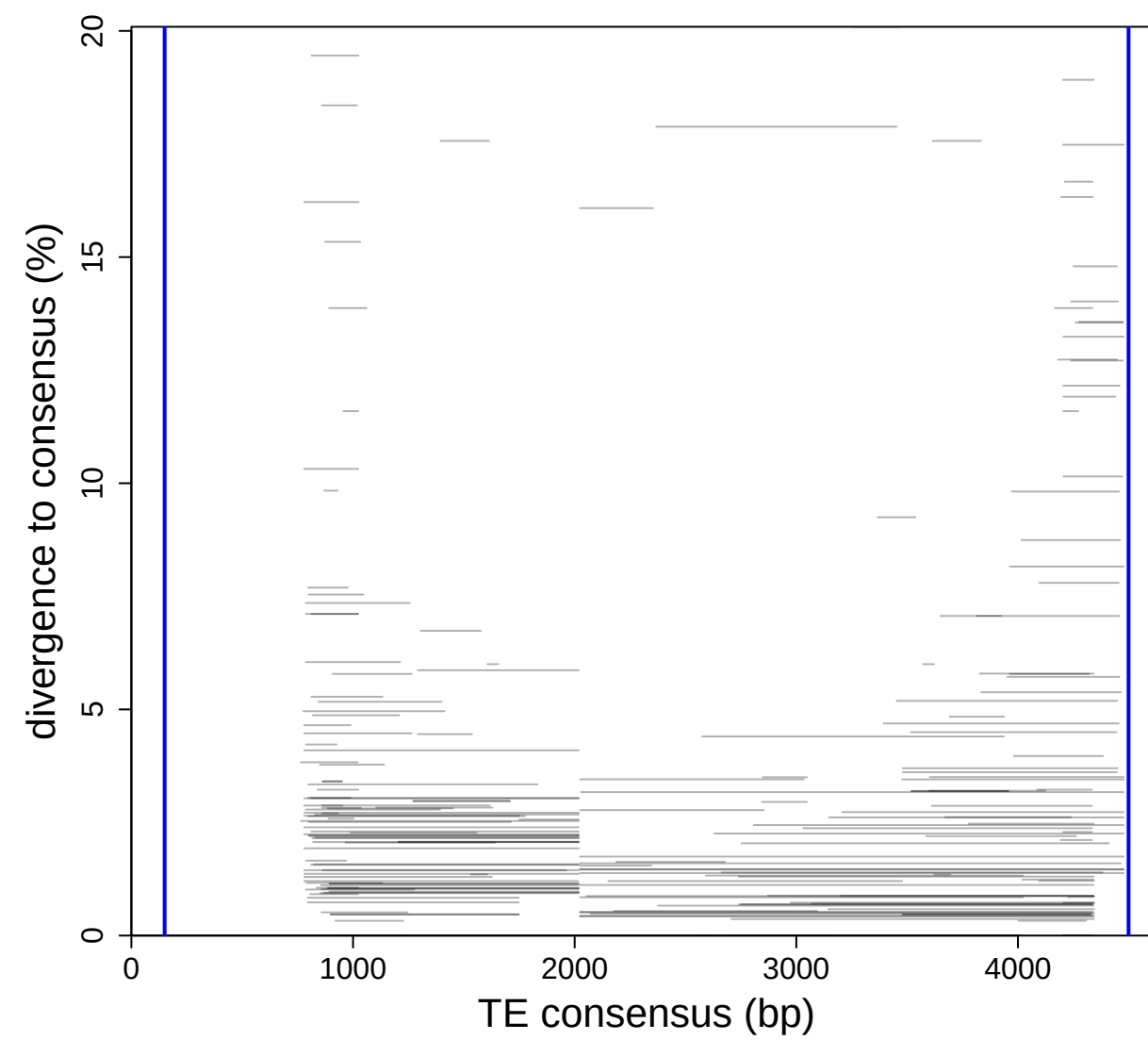
bed_fm_1.bed 0 0 n.bed g 1.bed fm_2.bed 0 0 bcln.fa aln.fa
size: 4061bp; fragments: 215; full length: 0 (>=3654.9bp)

After TEtrimmer 4061 bp



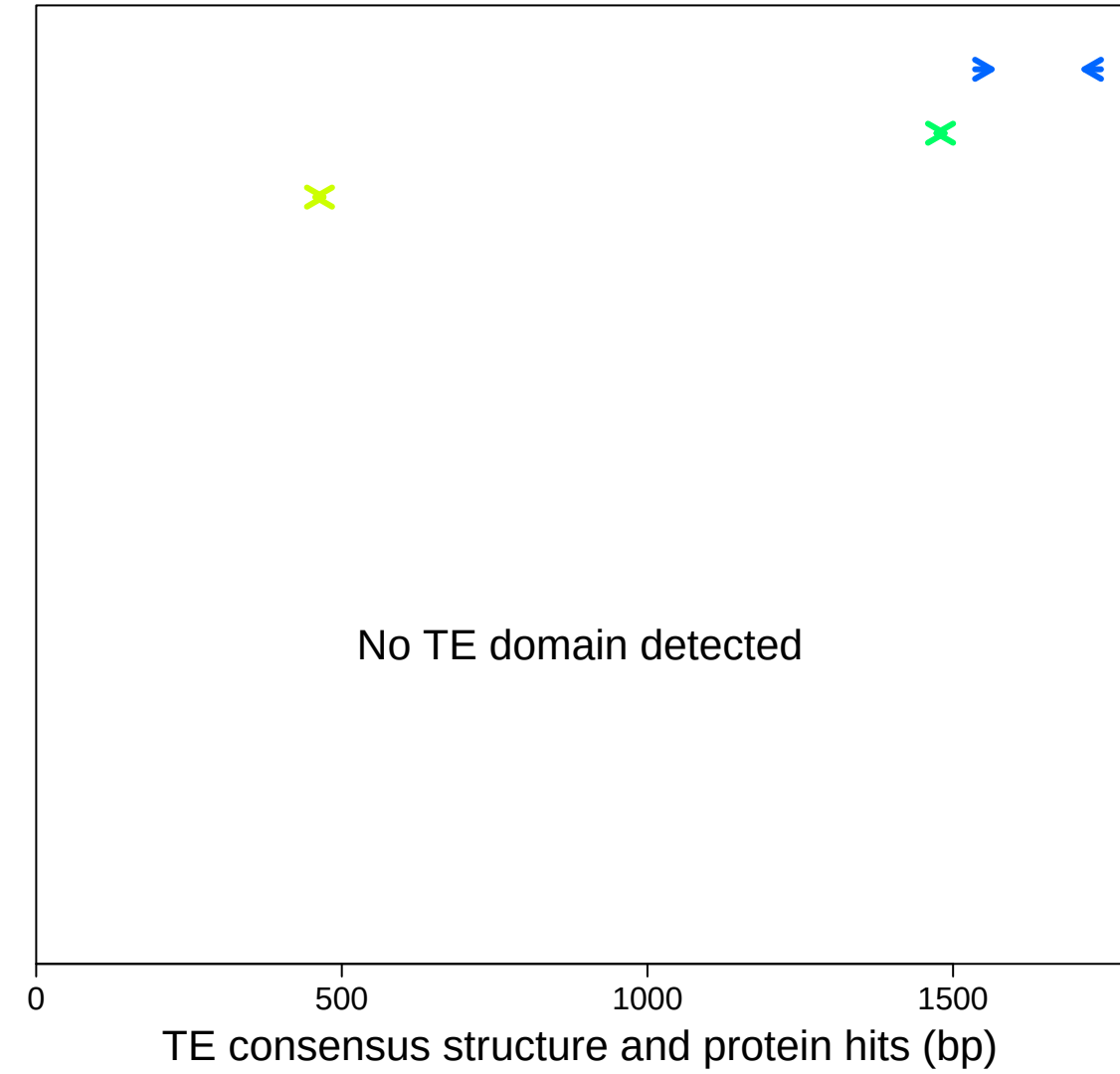
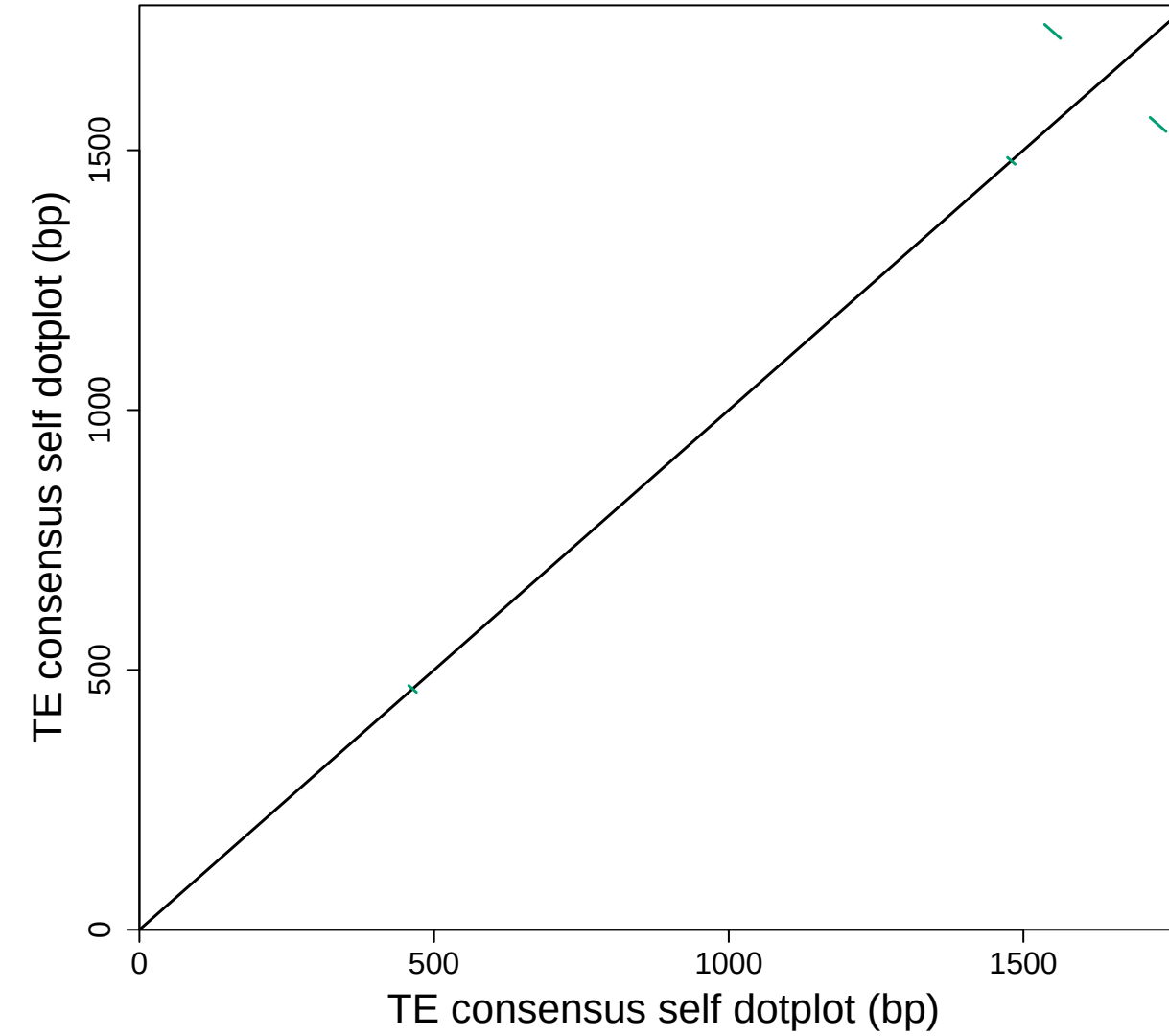
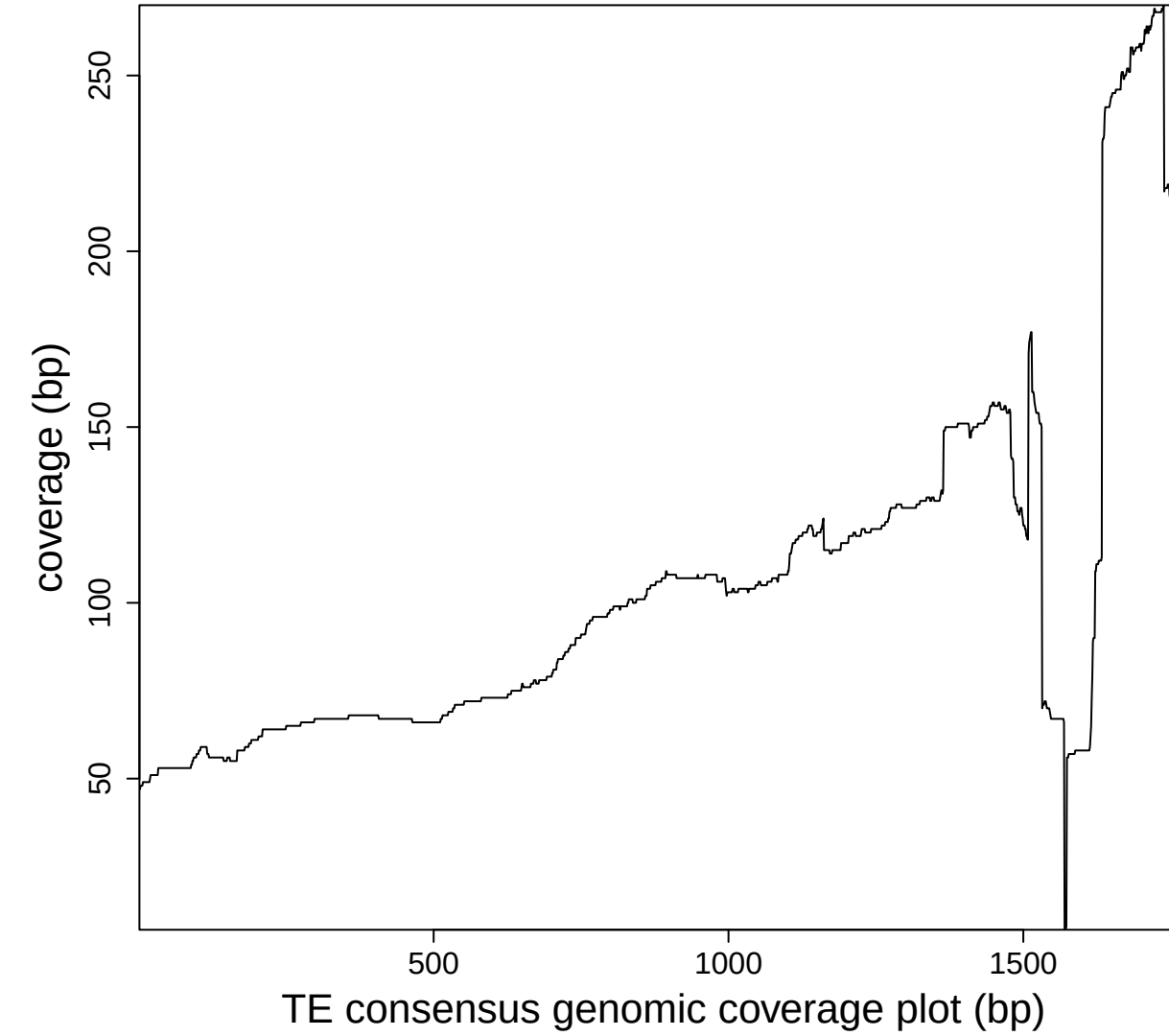
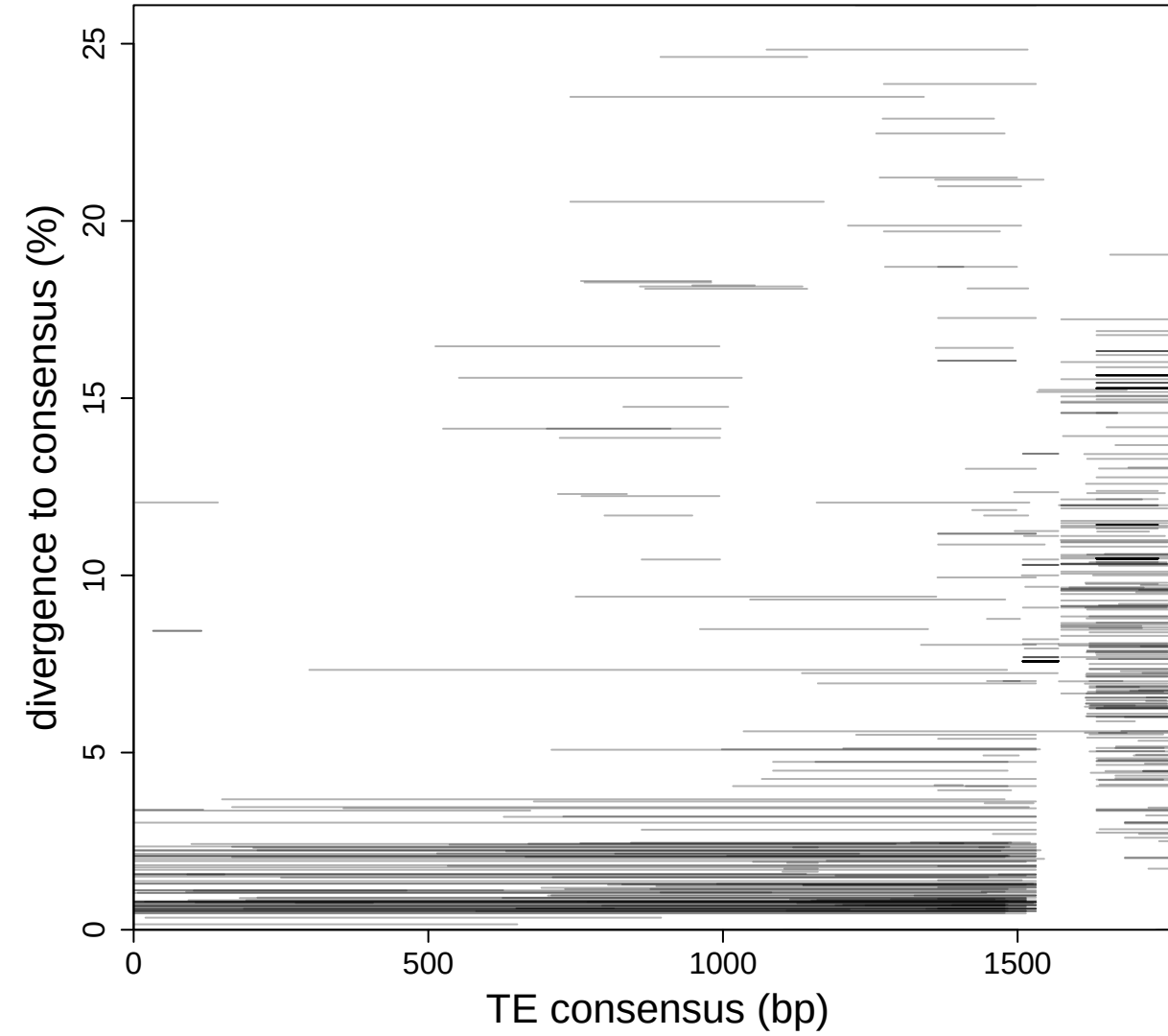
f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 4649bp; fragments: 210; full length: 0 (>=4184.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

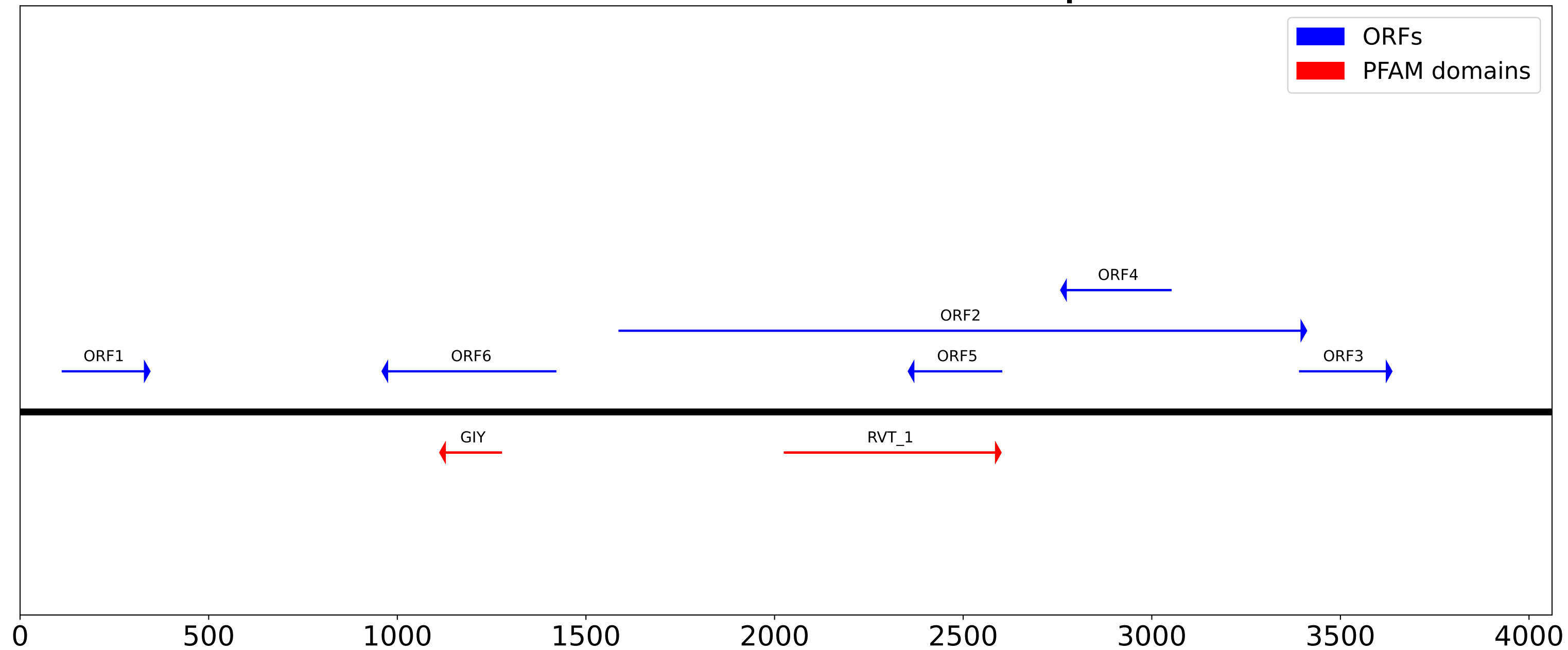


TE: rnd_5_family_983
size: 1779bp; fragments: 558; full length: 0 (>=1601.1bp)

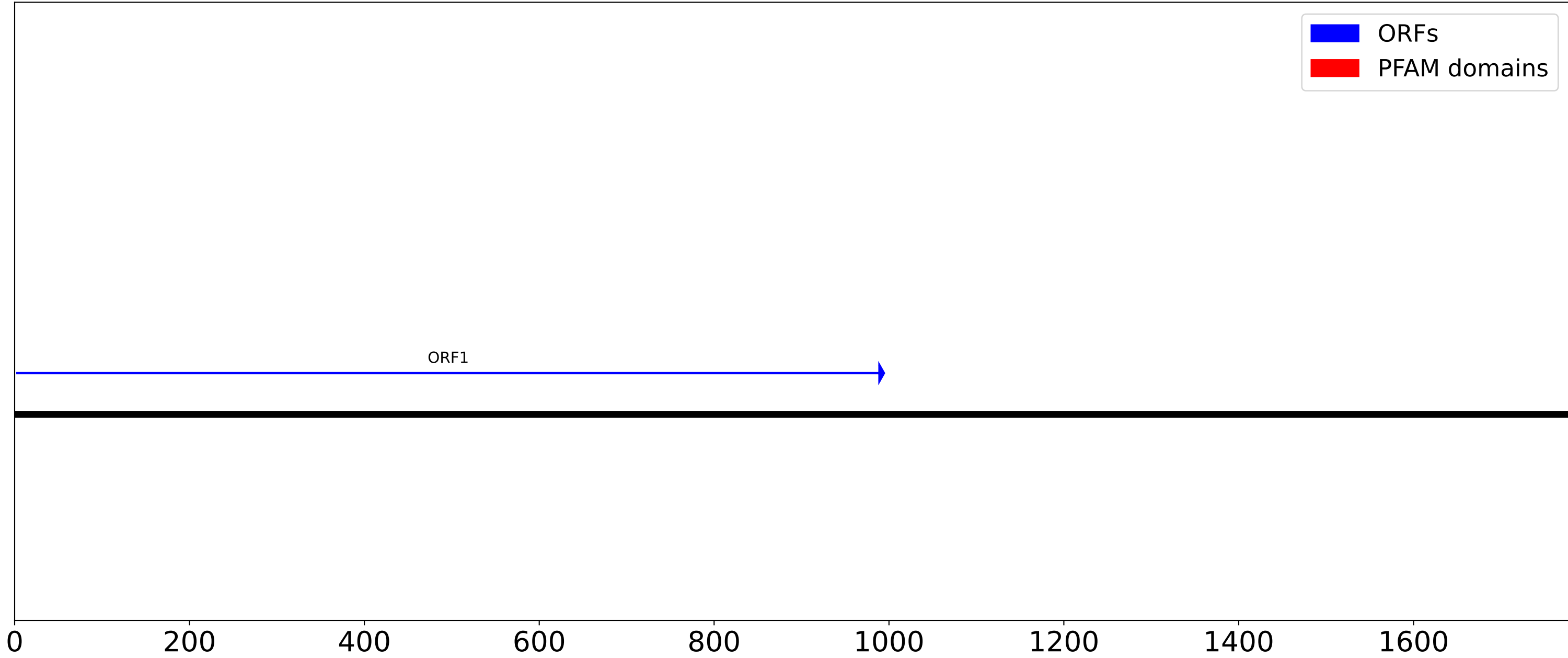
Before TEtrimmer 1779 bp



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)

