

MSA length = 4493

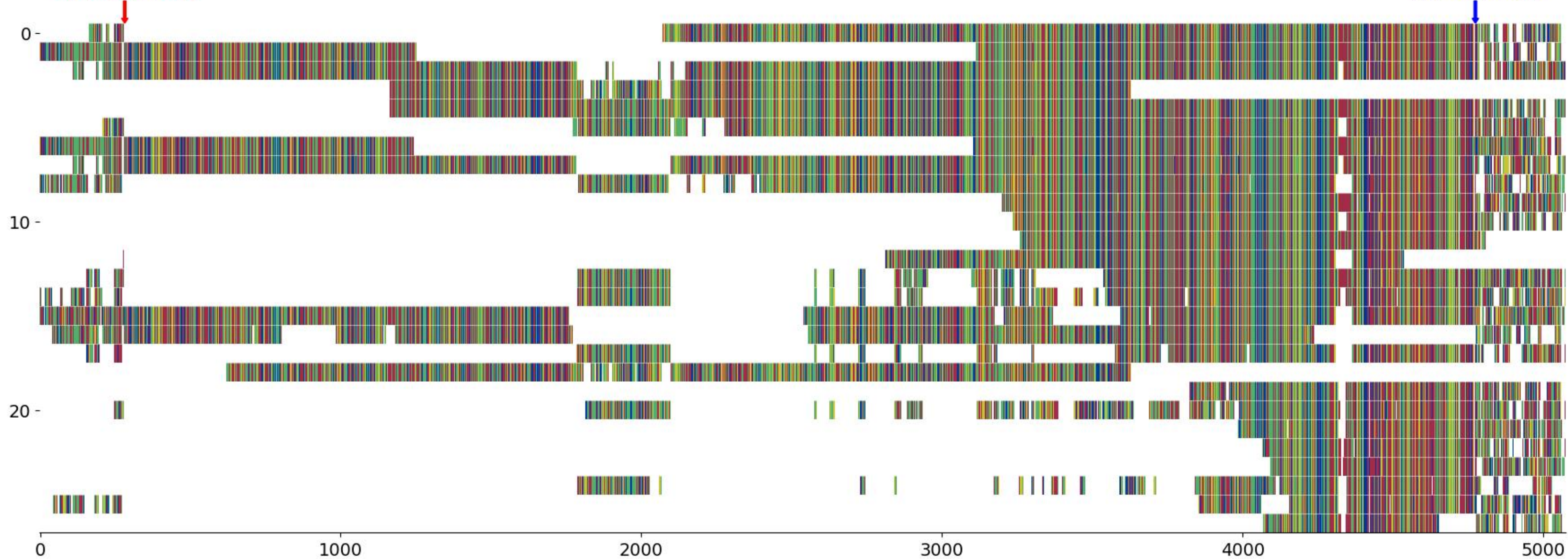
Start crop Point

End crop Point



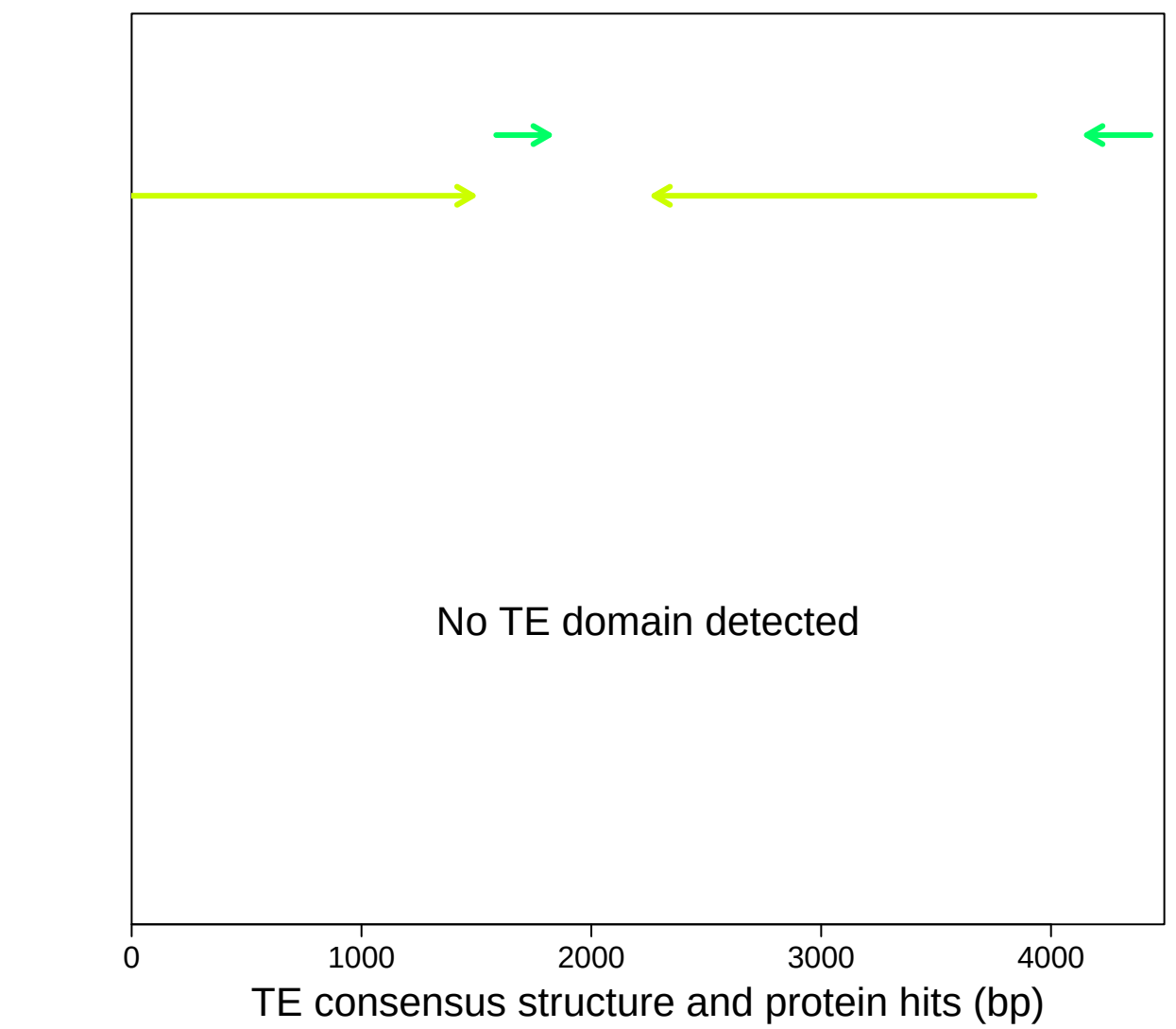
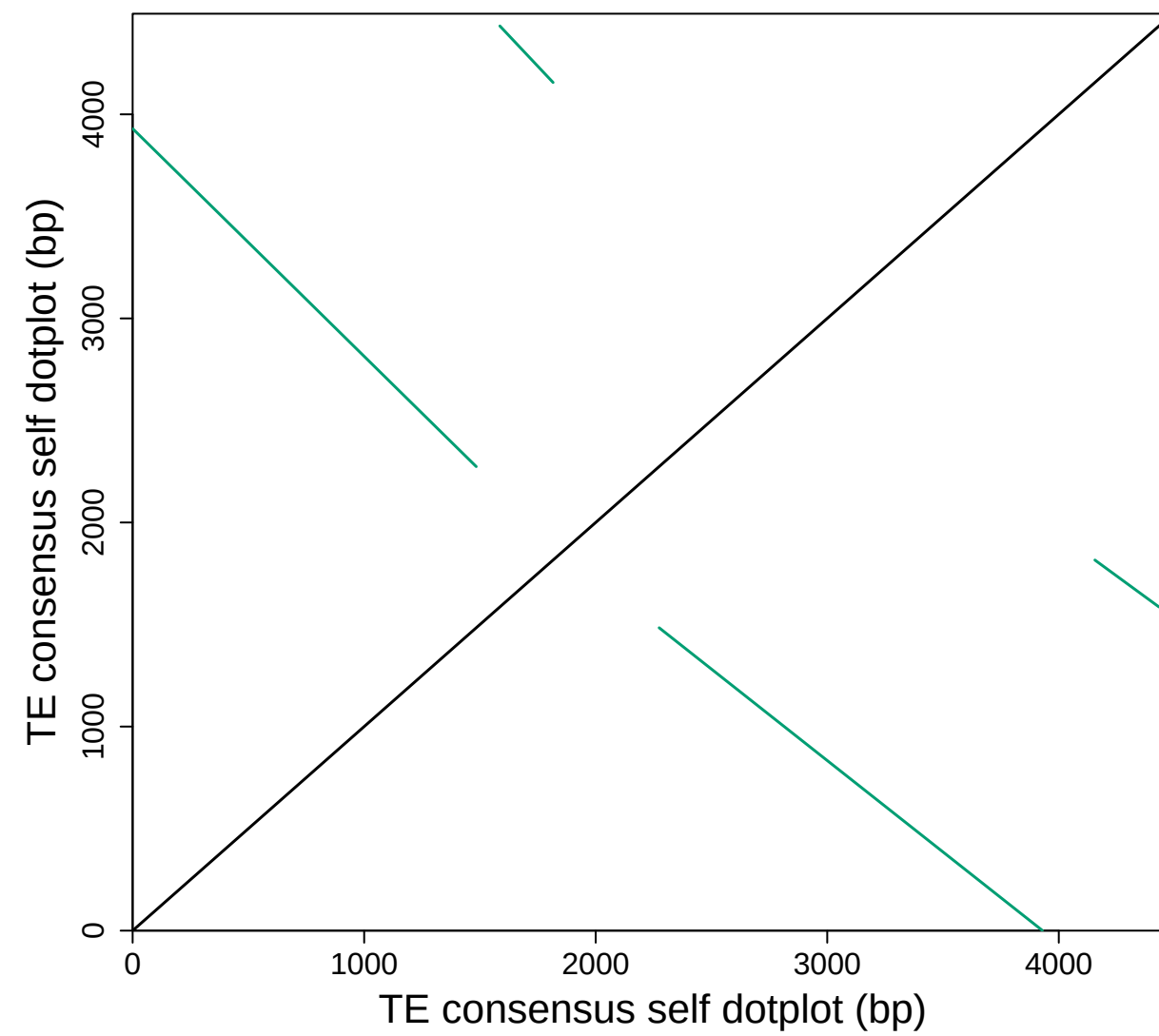
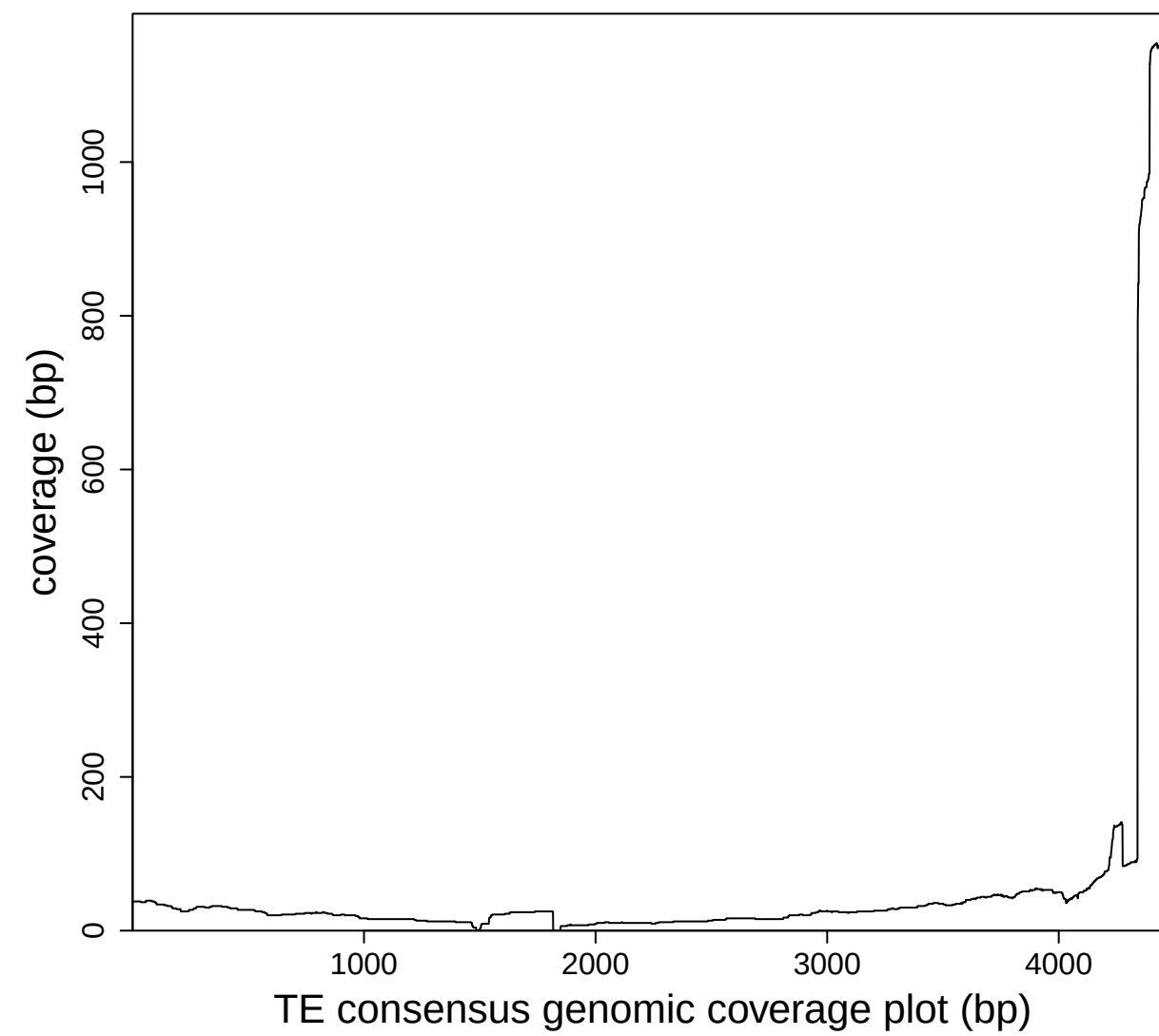
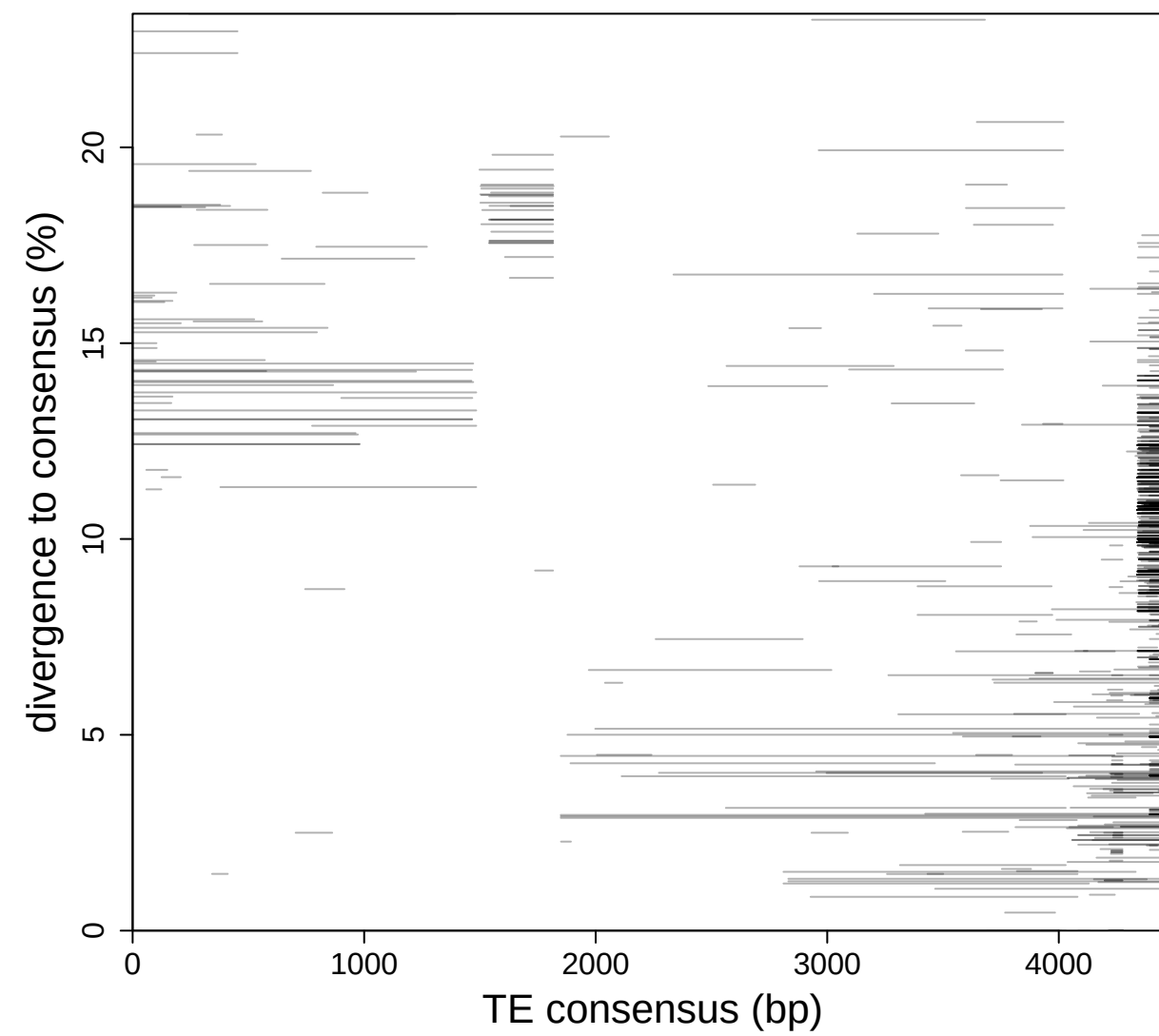
Start crop Point

End crop Point



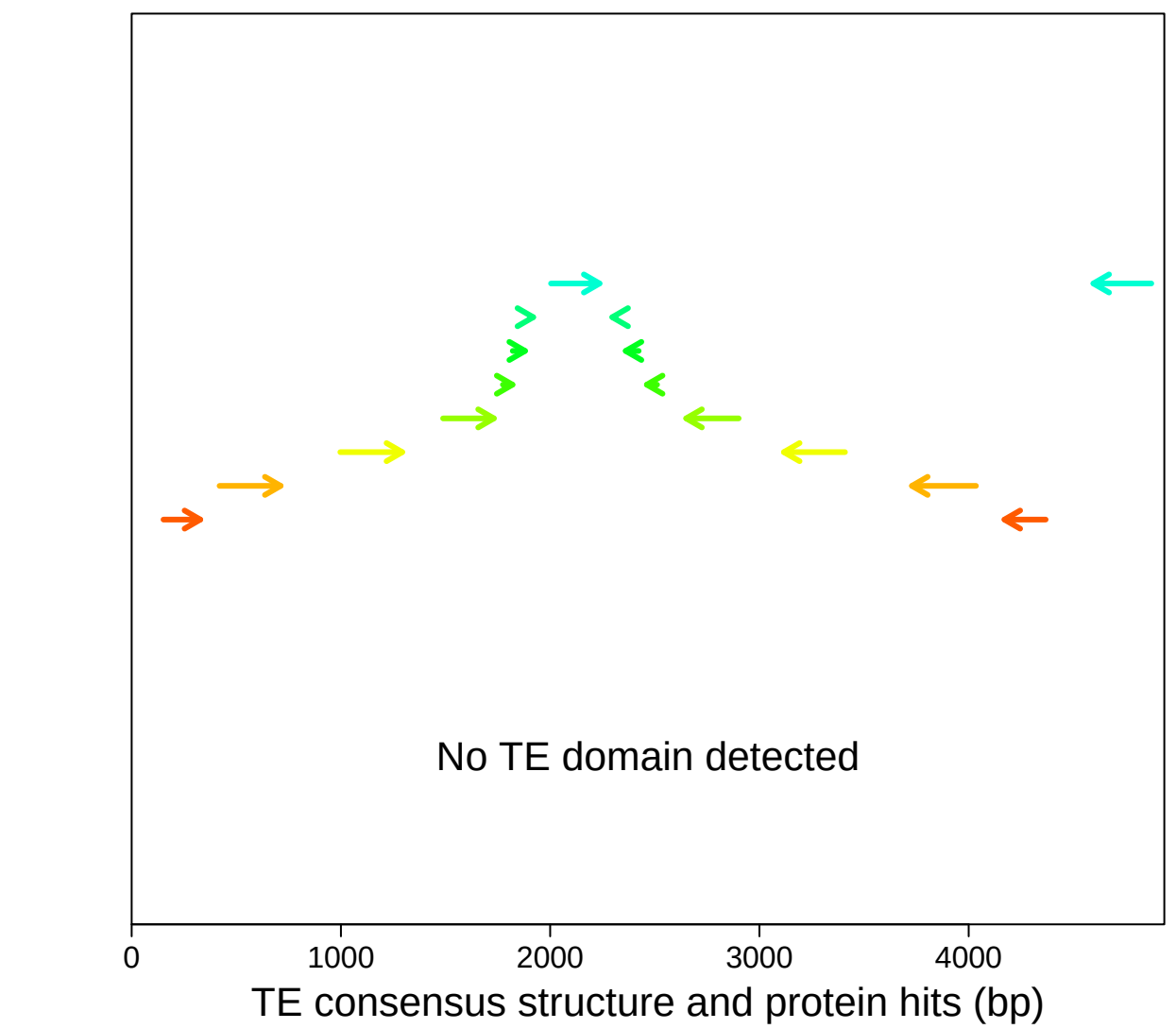
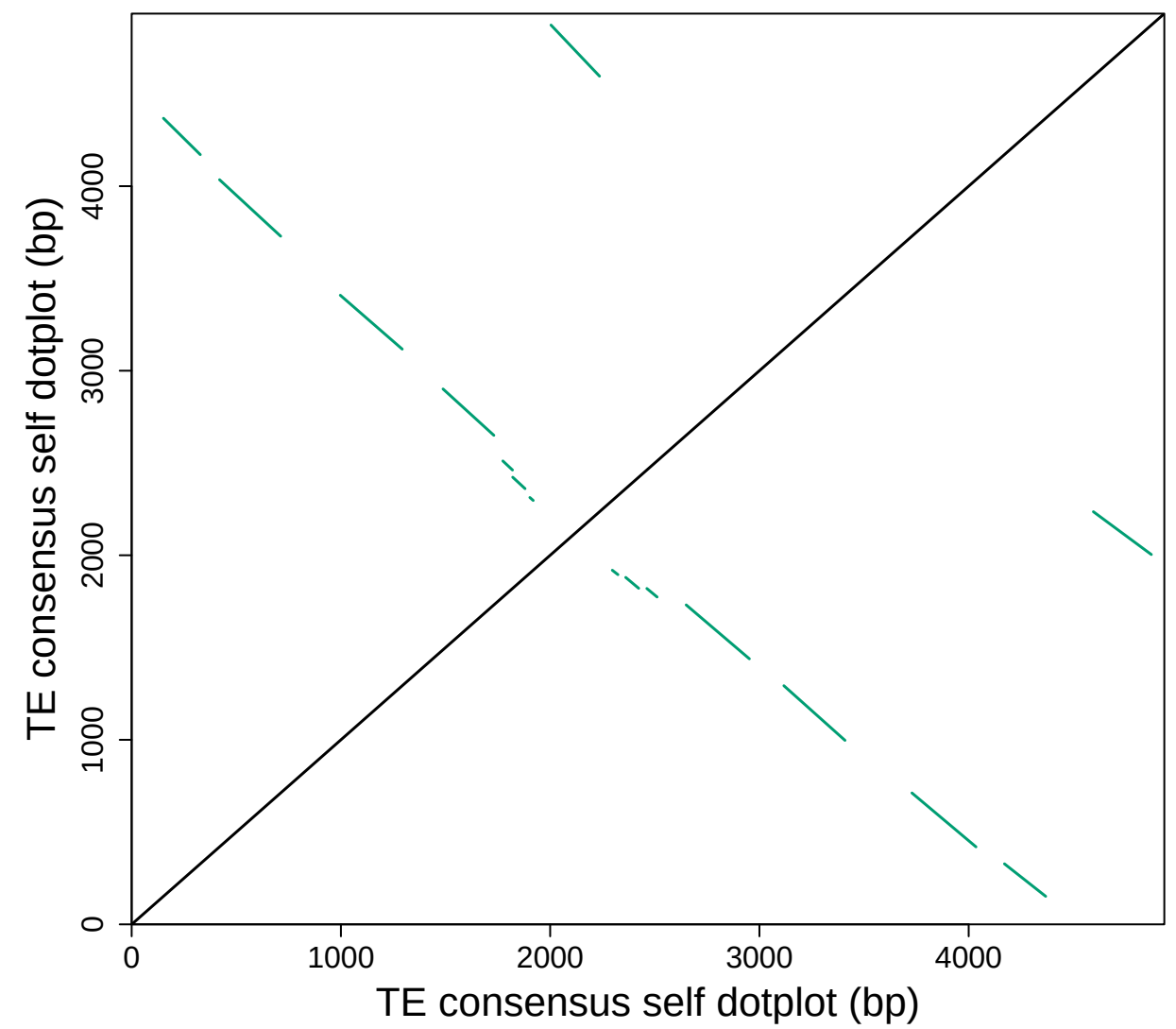
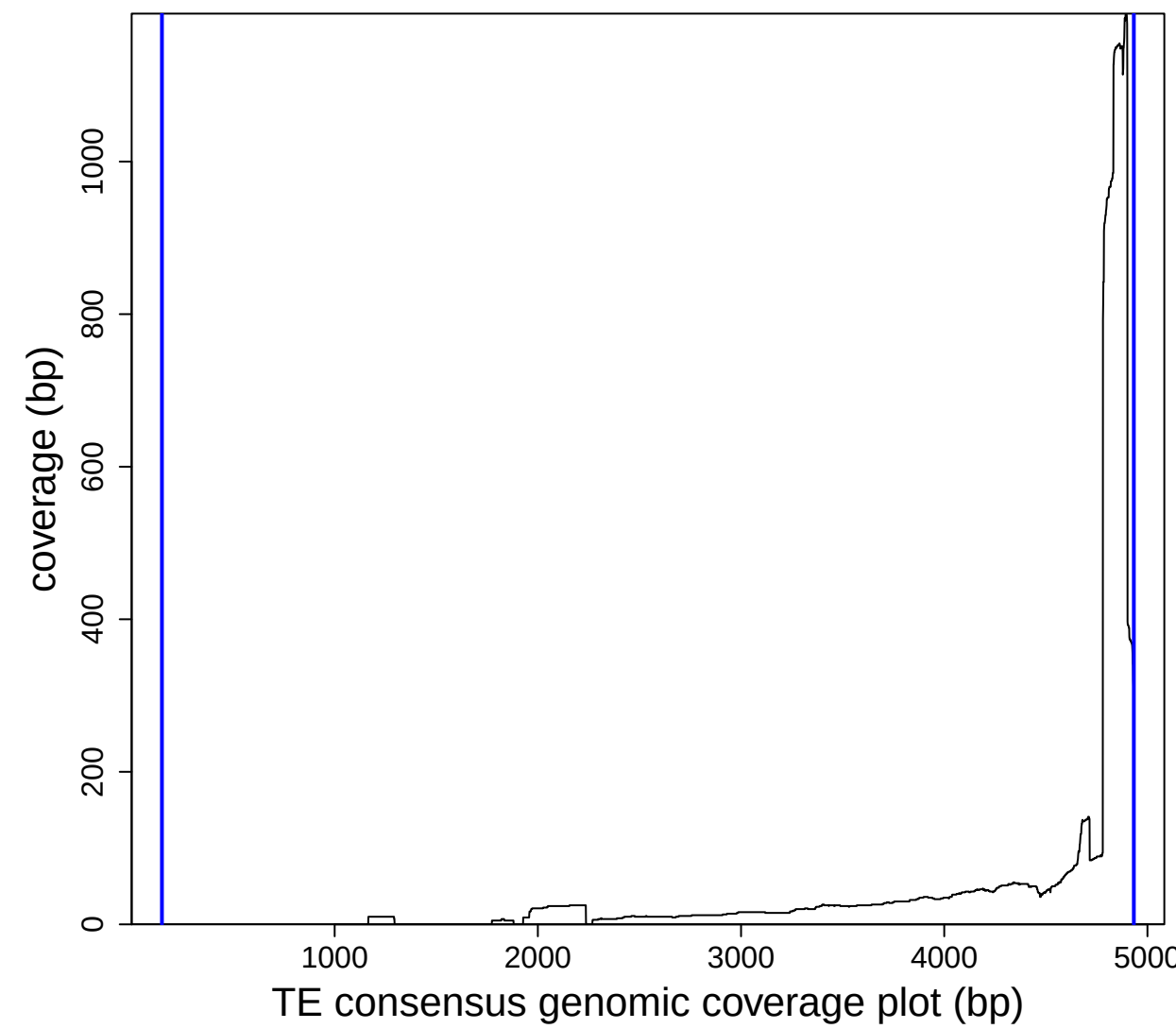
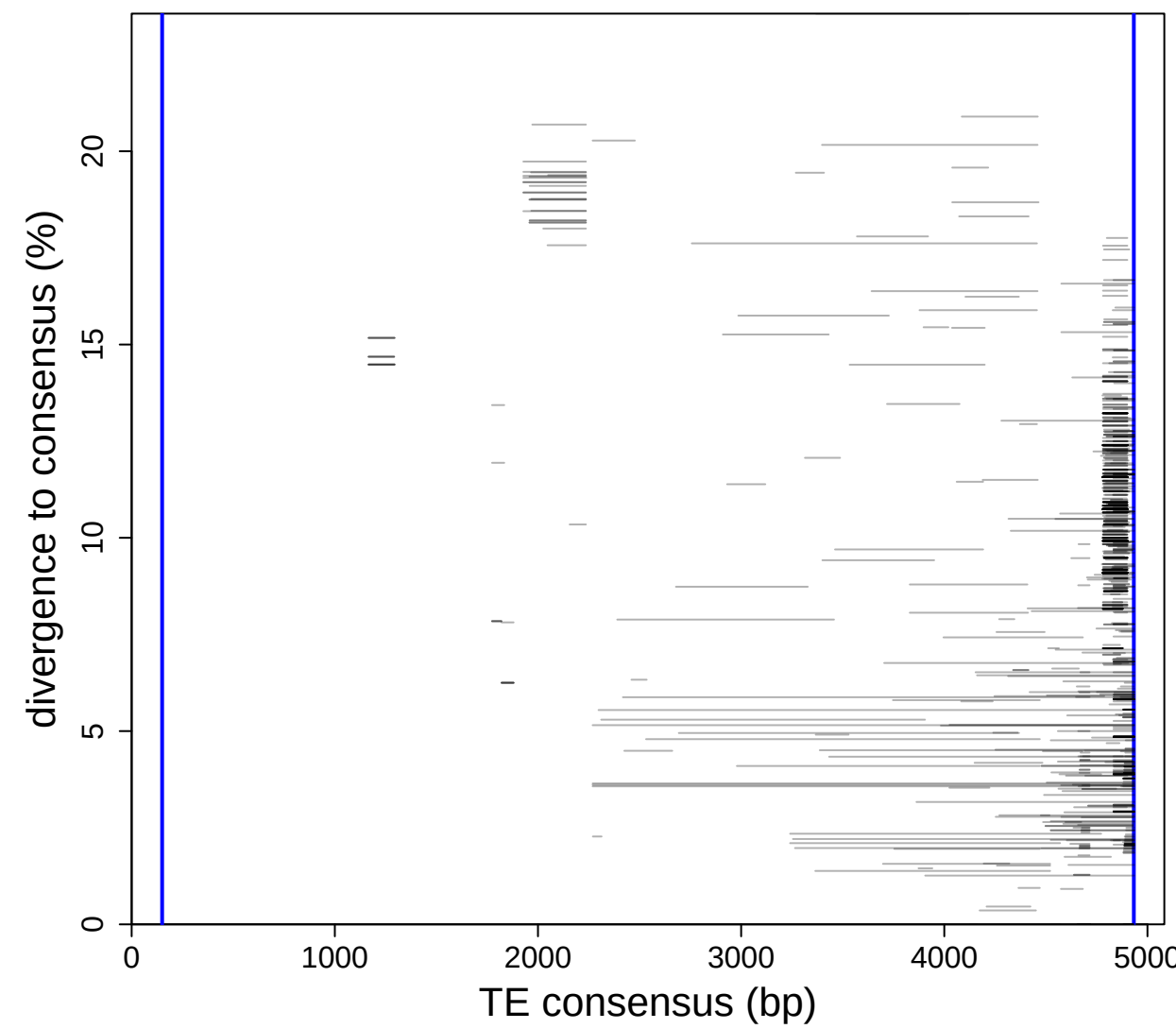
bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.fa
size: 4493bp; fragments: 1458; full length: 0 (>=4043.7bp)

After TETrimmer 4493 bp



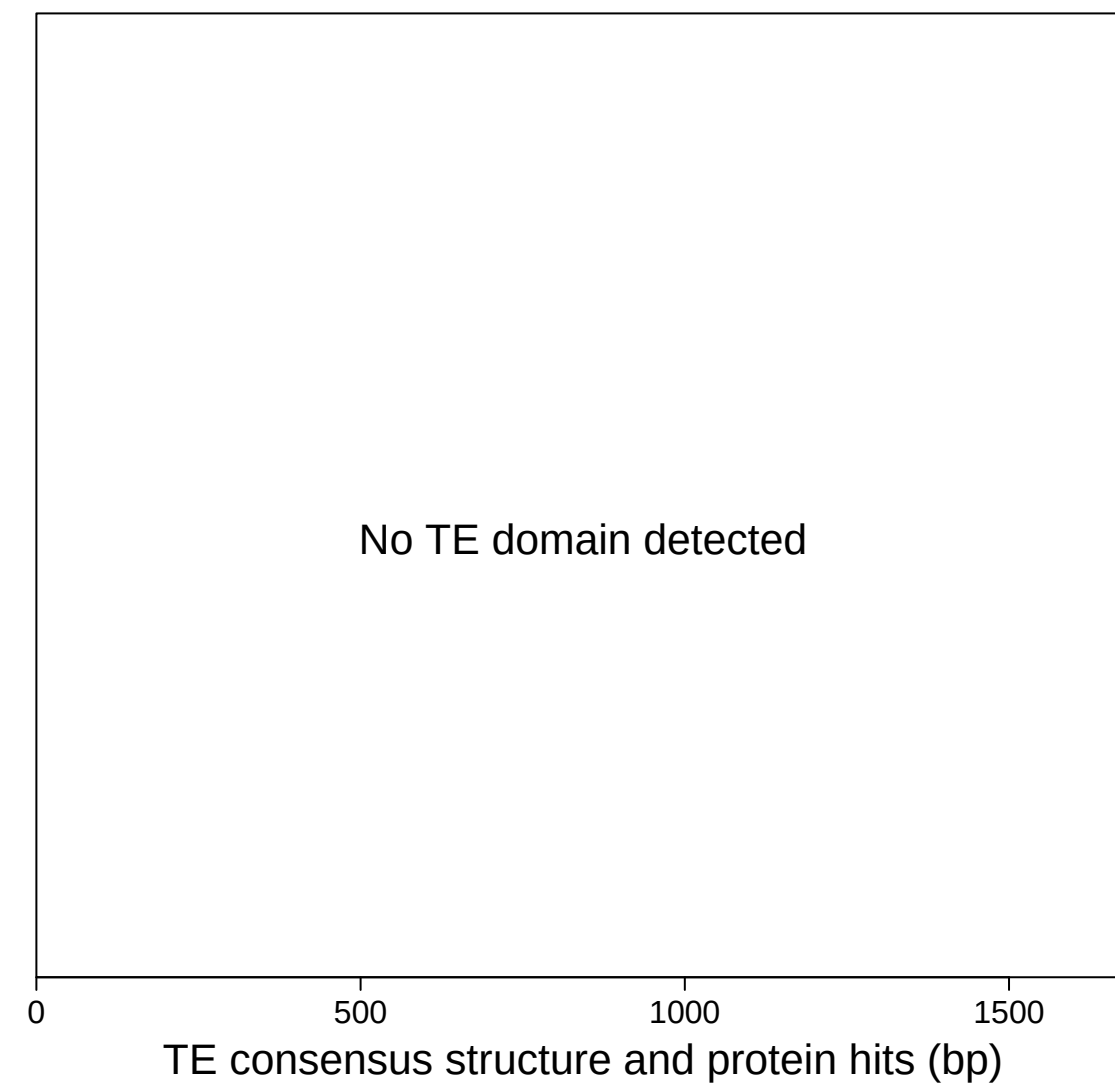
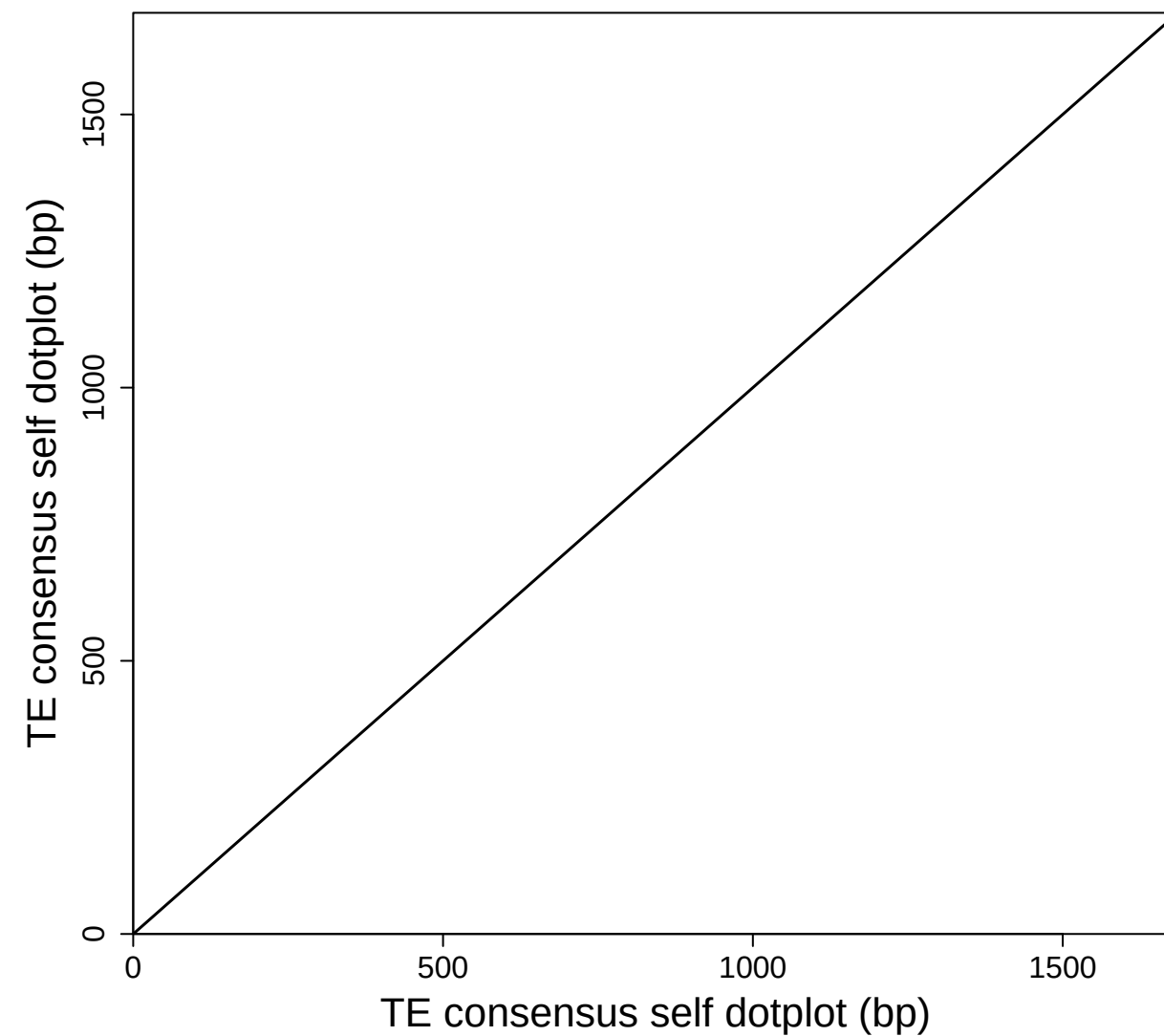
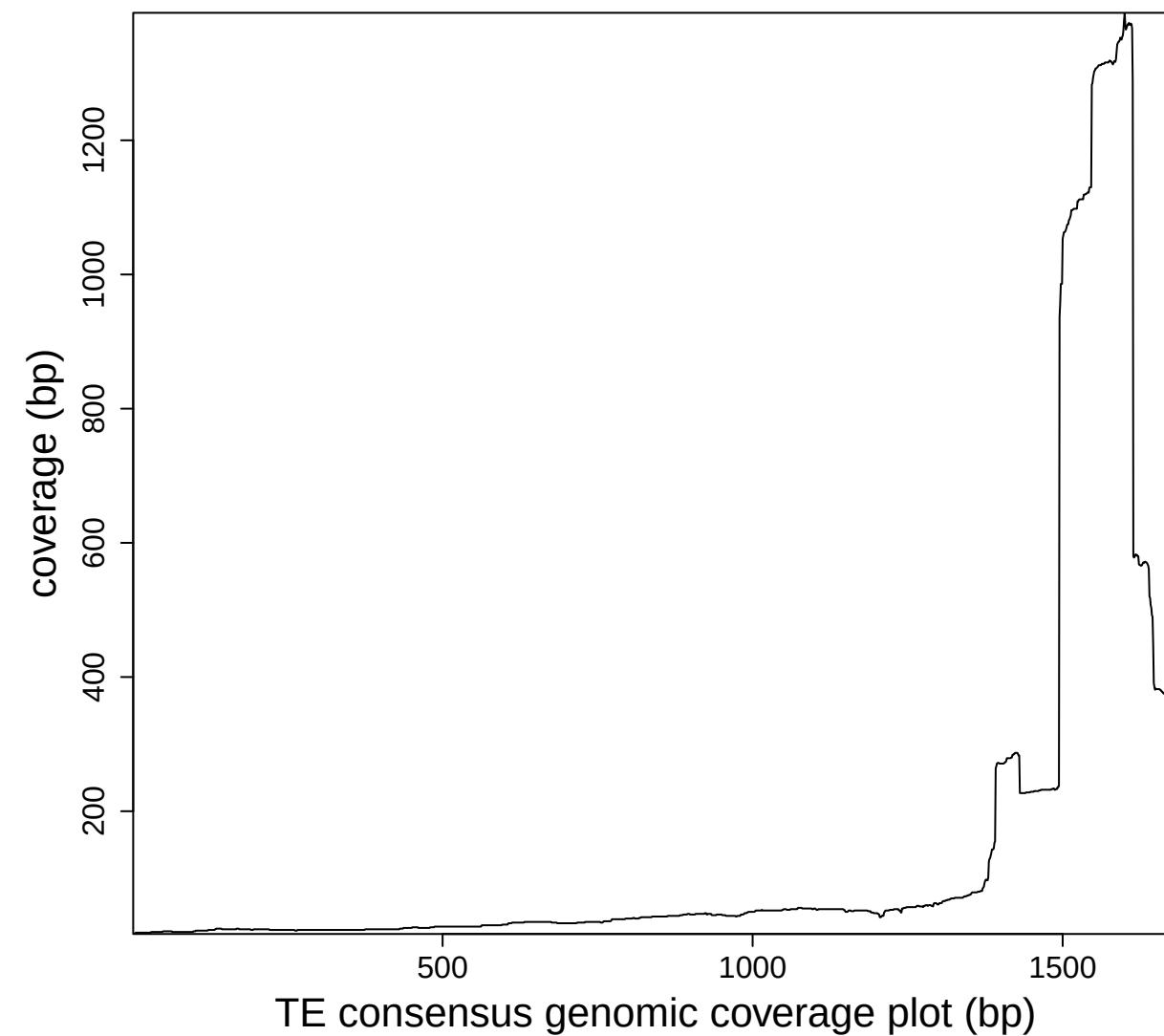
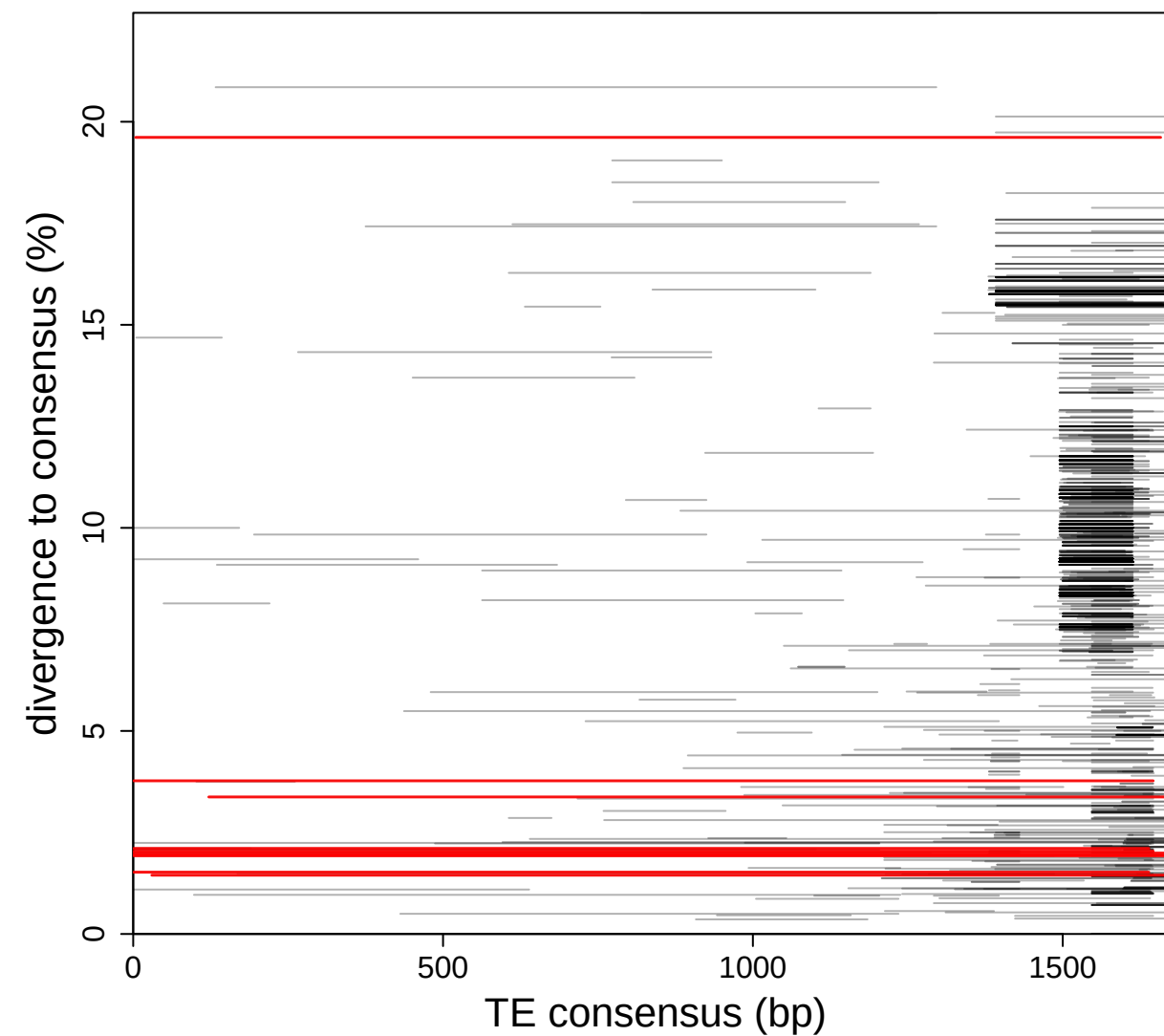
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 5083bp; fragments: 1422; full length: 0 (>=4574.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

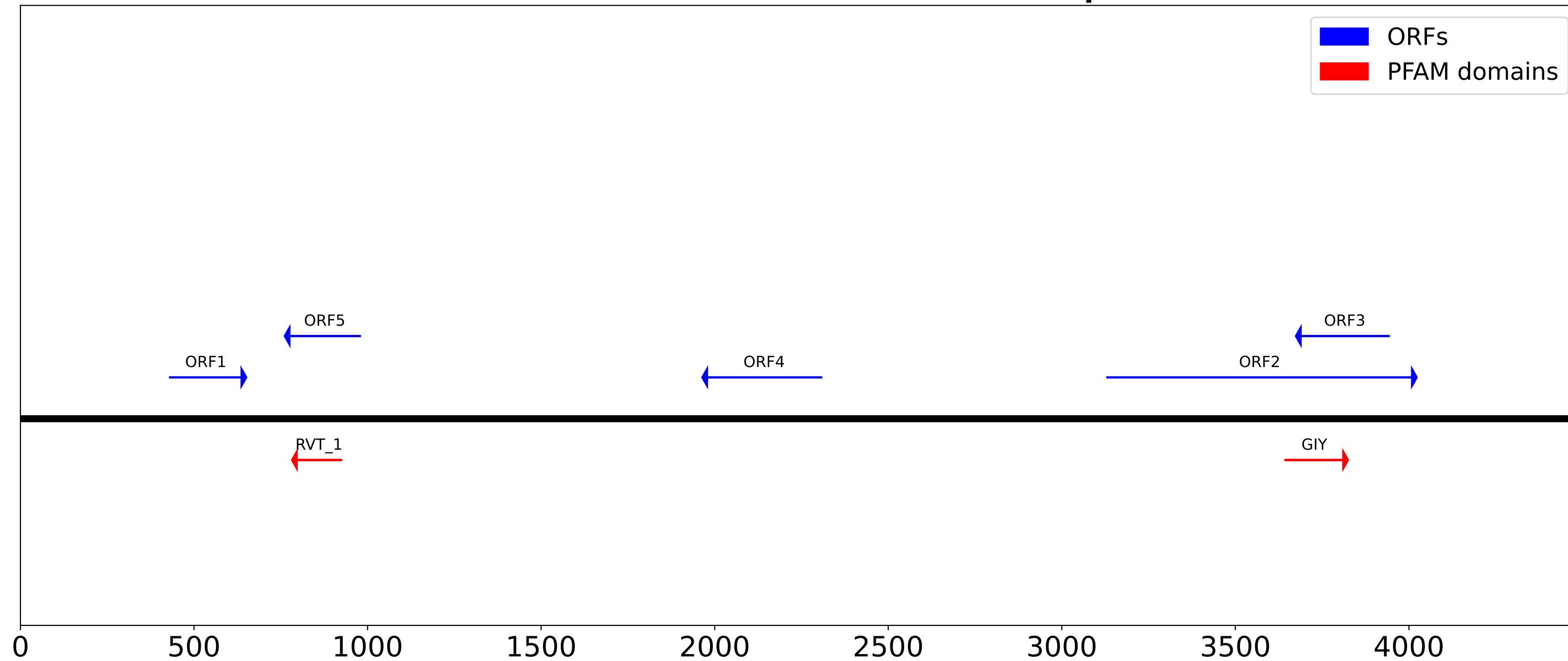


TE: rnd_4_family_637
size: 1686bp; fragments: 1588; full length: 10 (≥ 1517.4 bp)

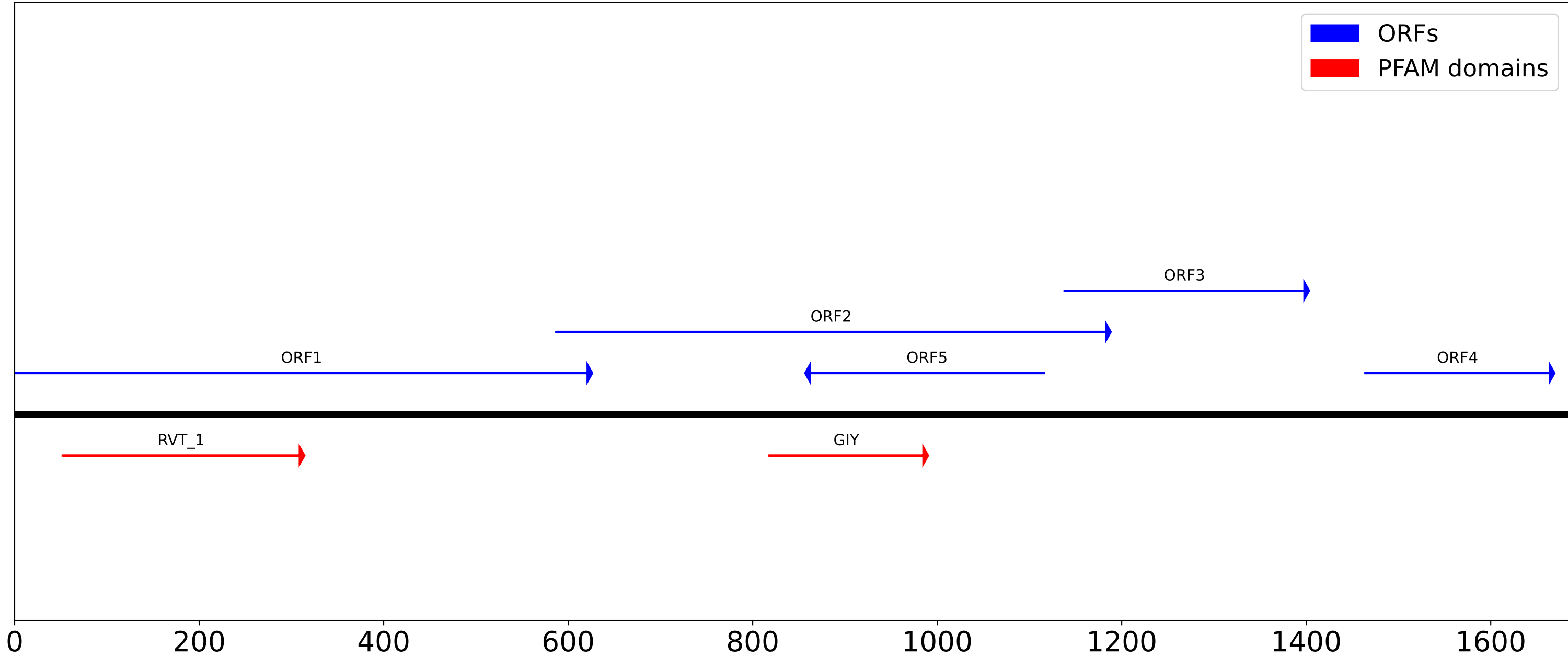
Before TEtrimmer 1686 bp



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)

