

Start crop Point



End crop Point

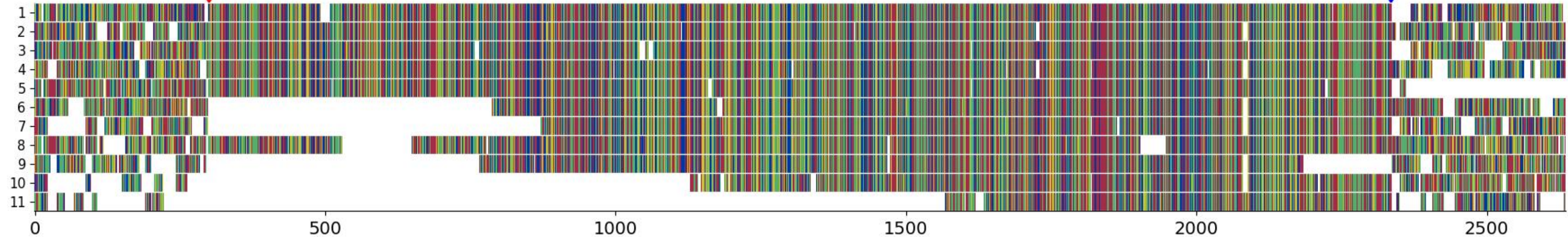


MSA length = 2036



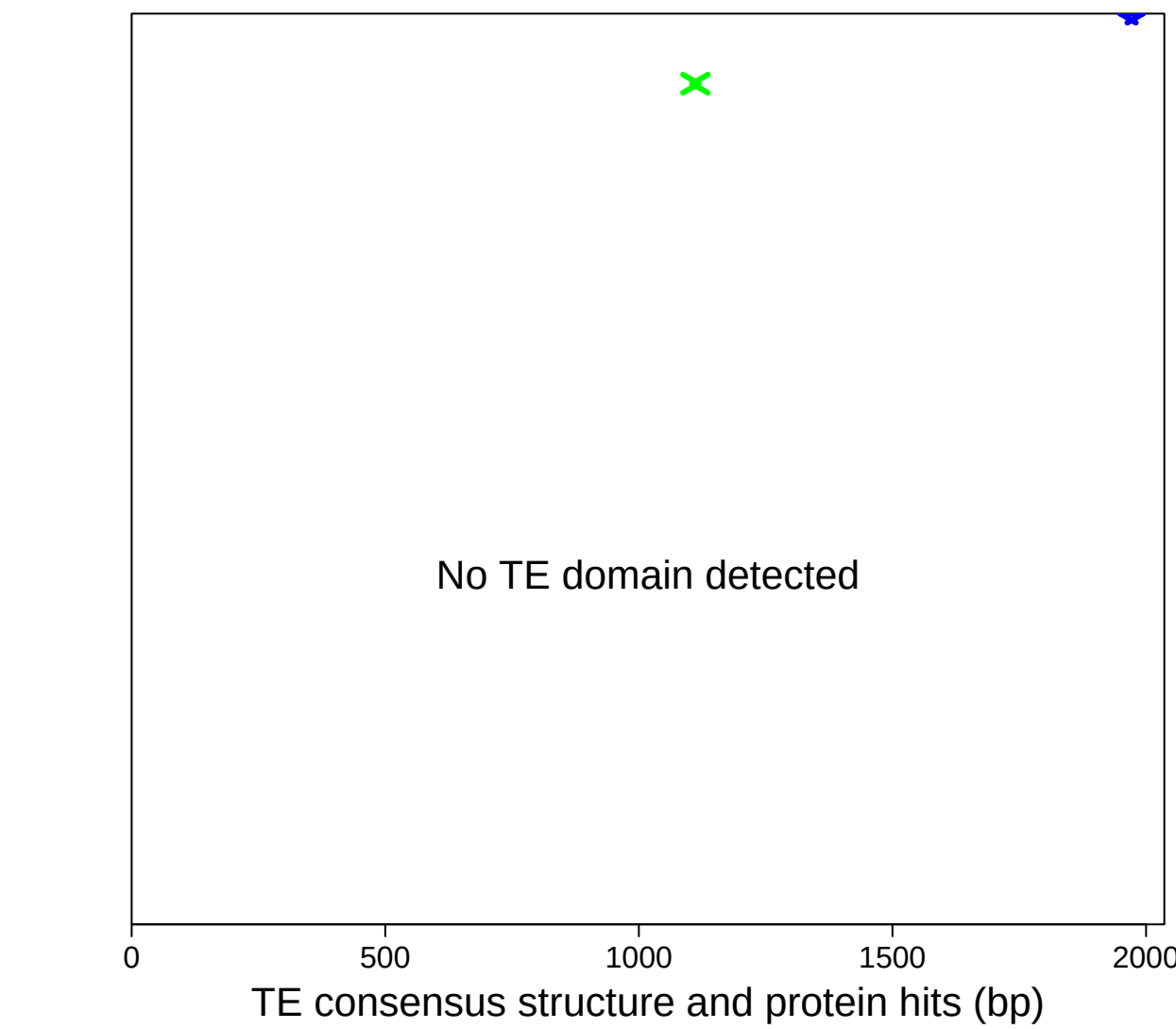
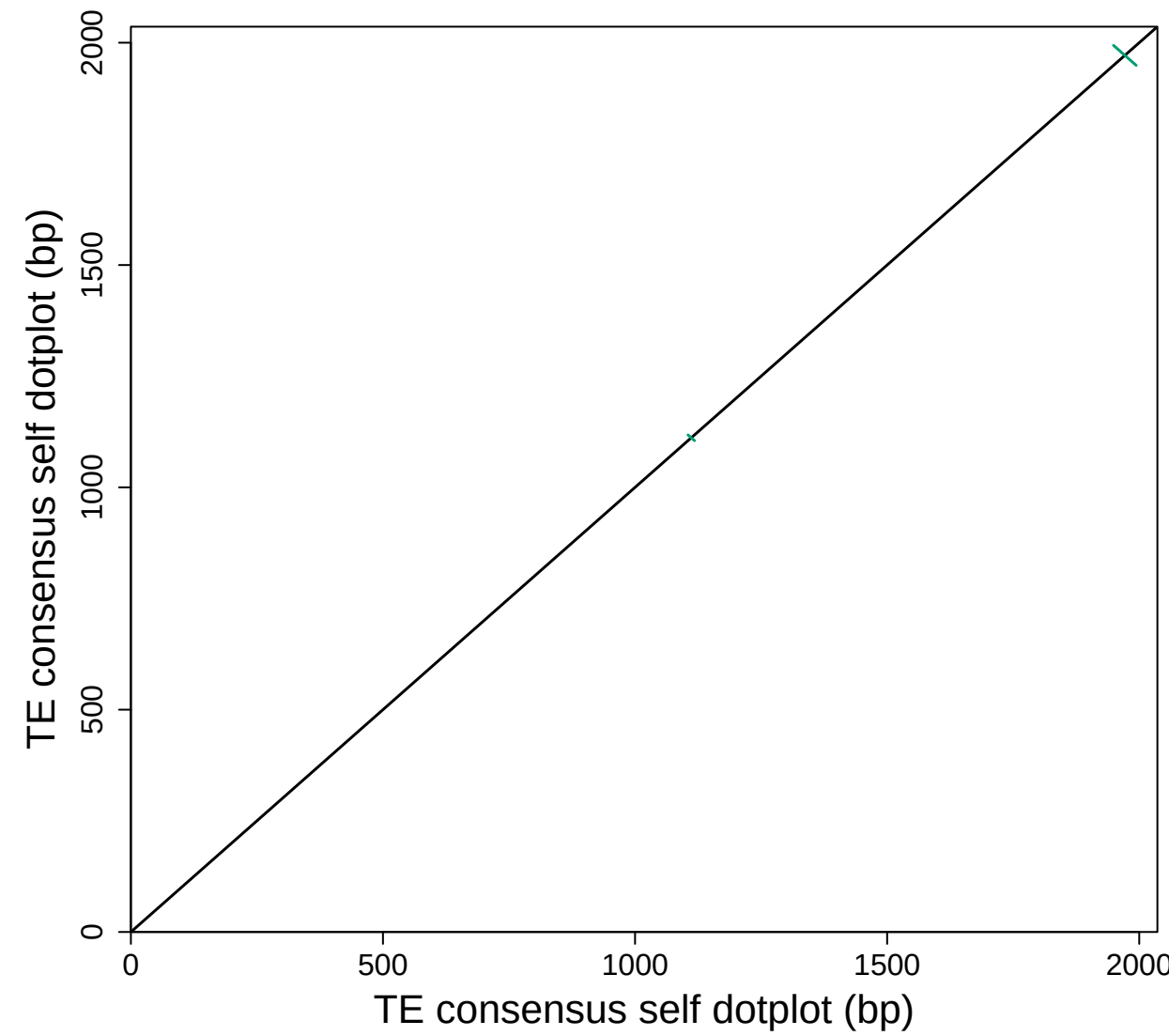
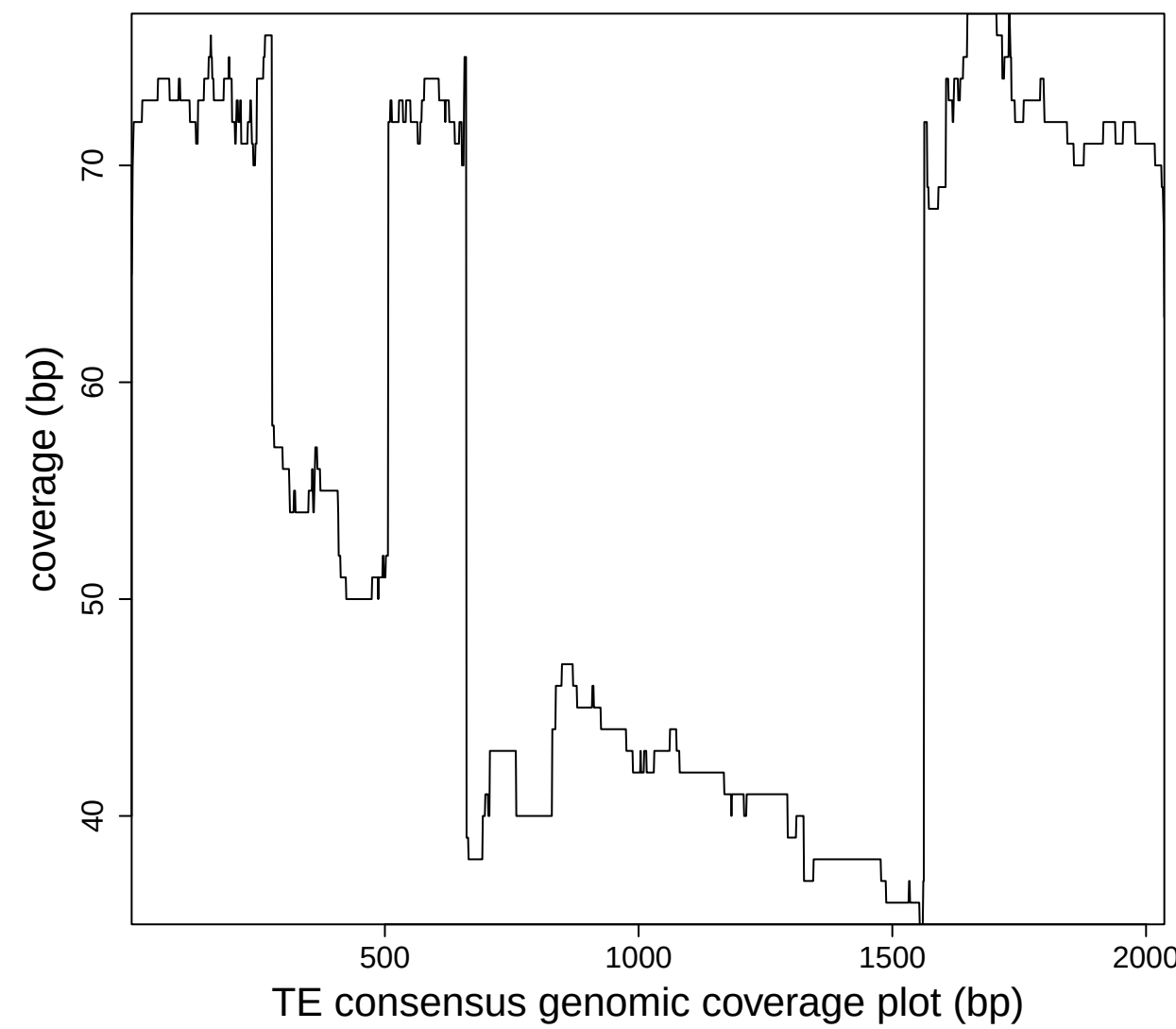
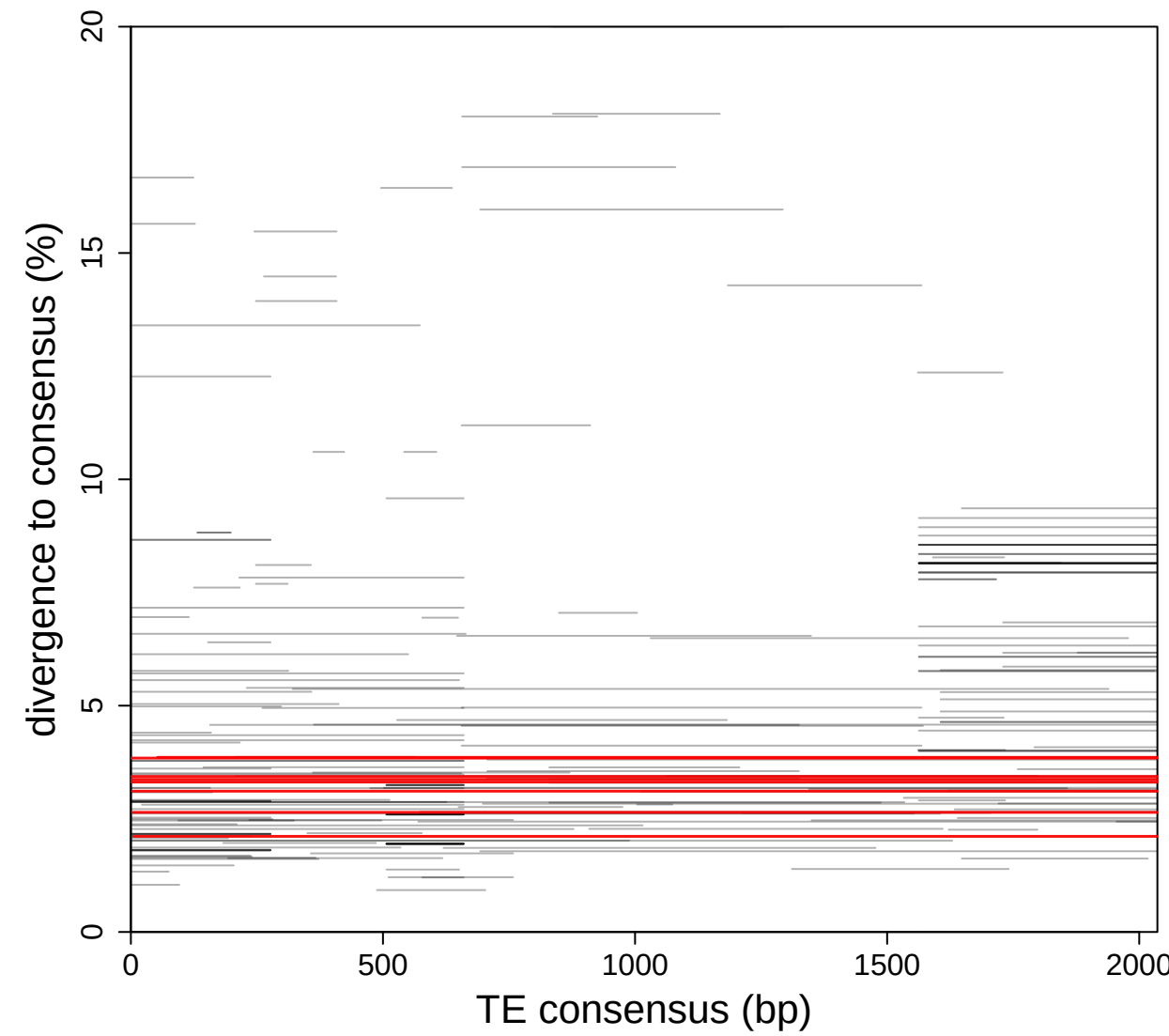
Start crop Point

End crop Point



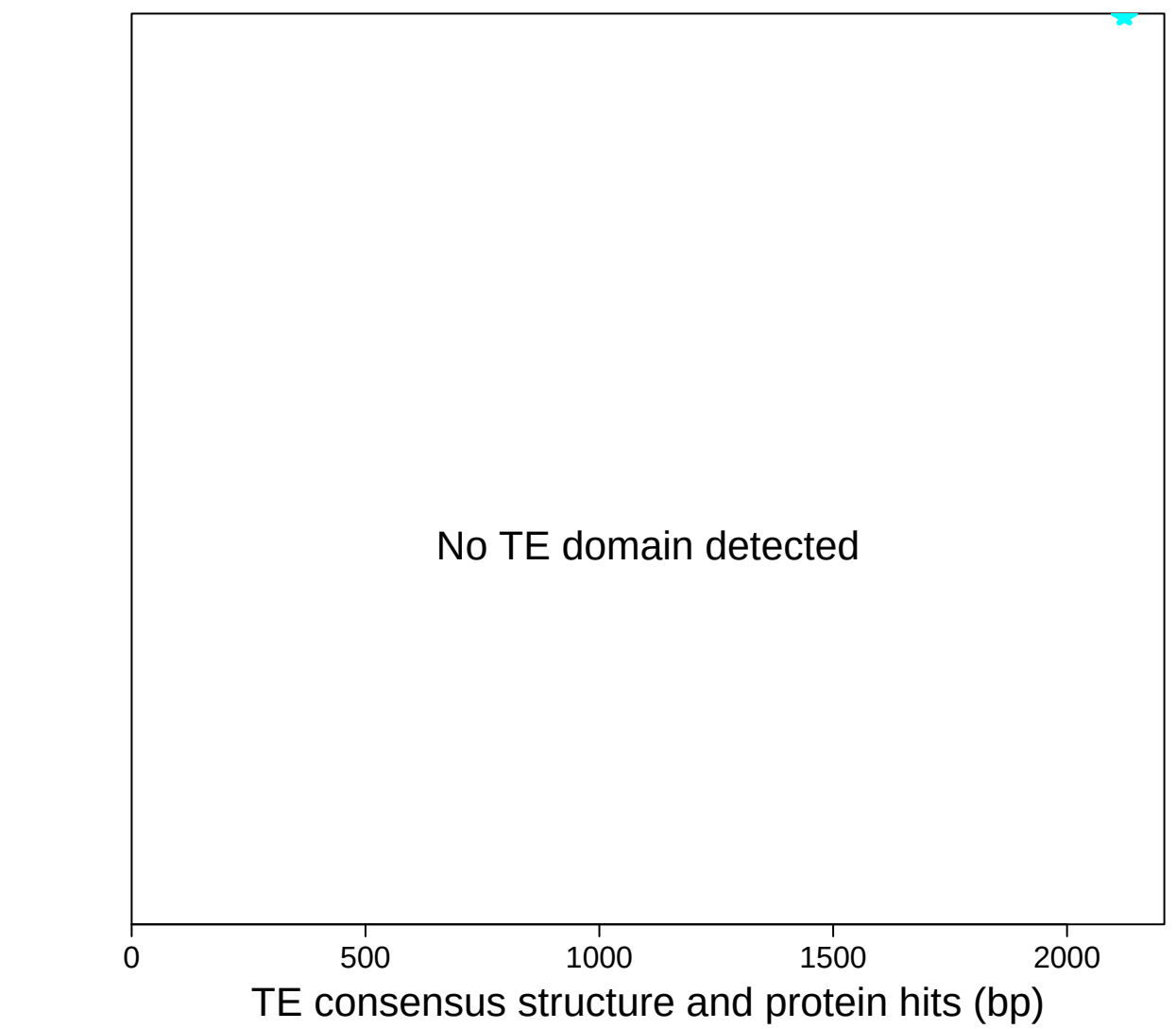
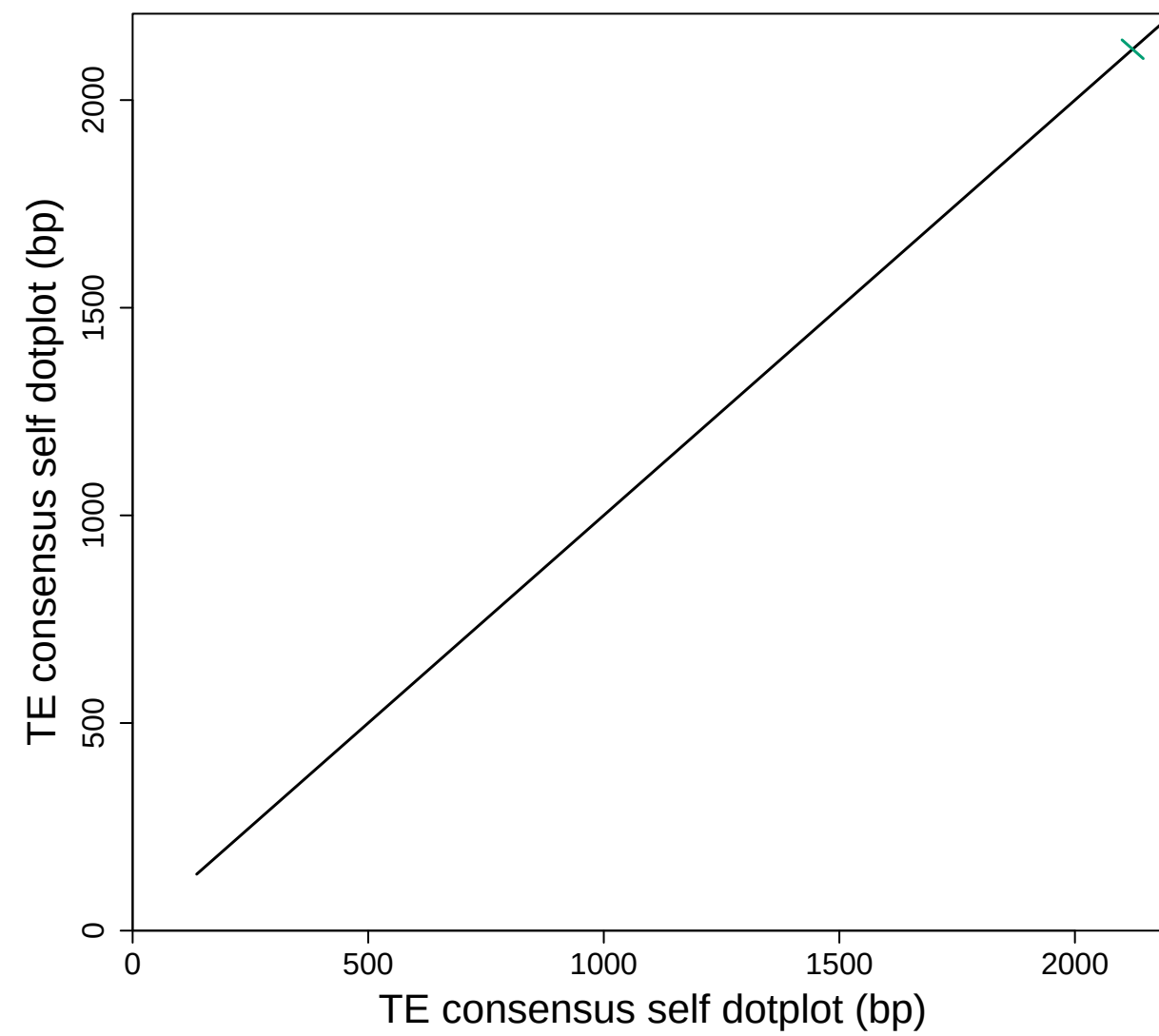
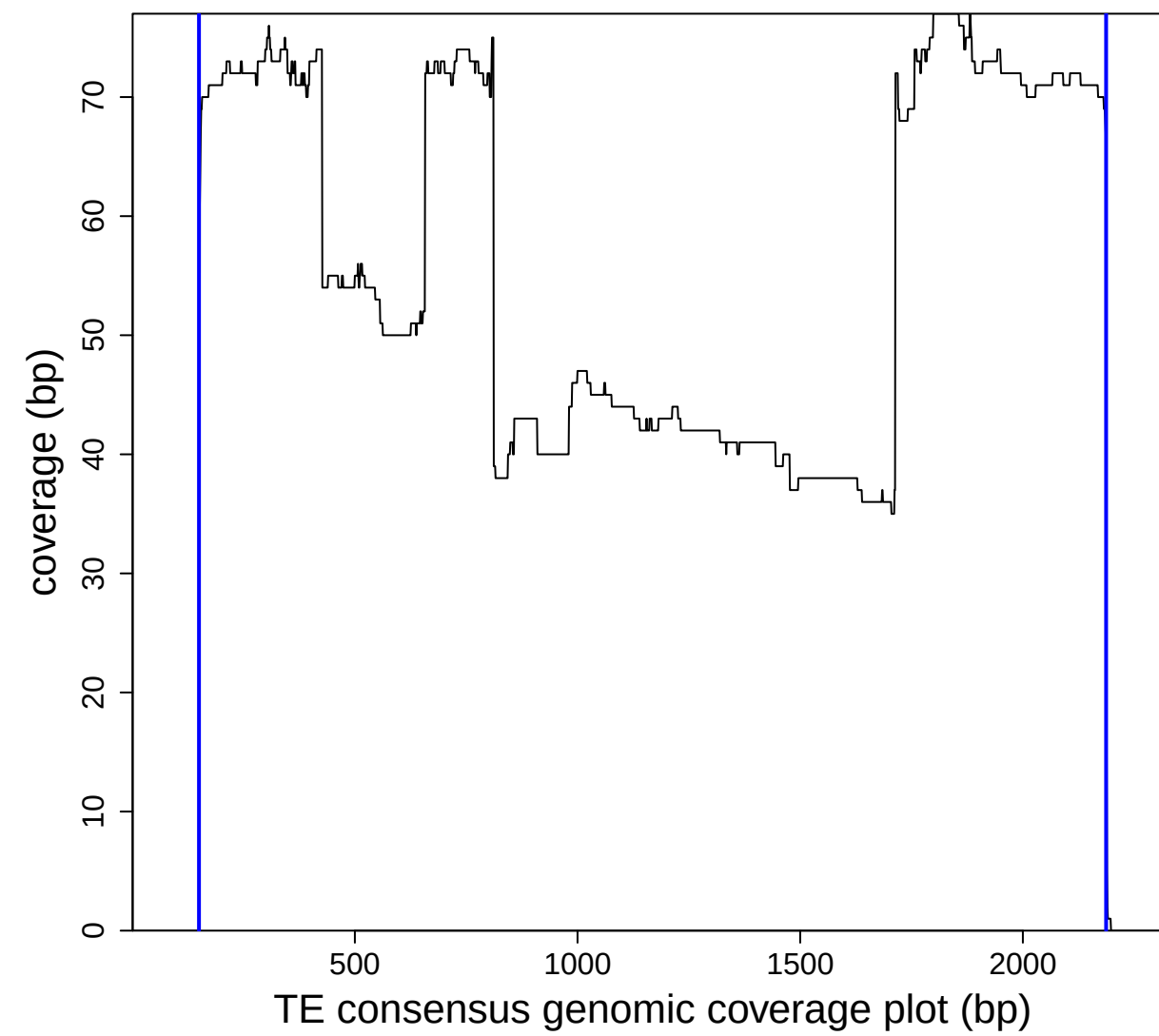
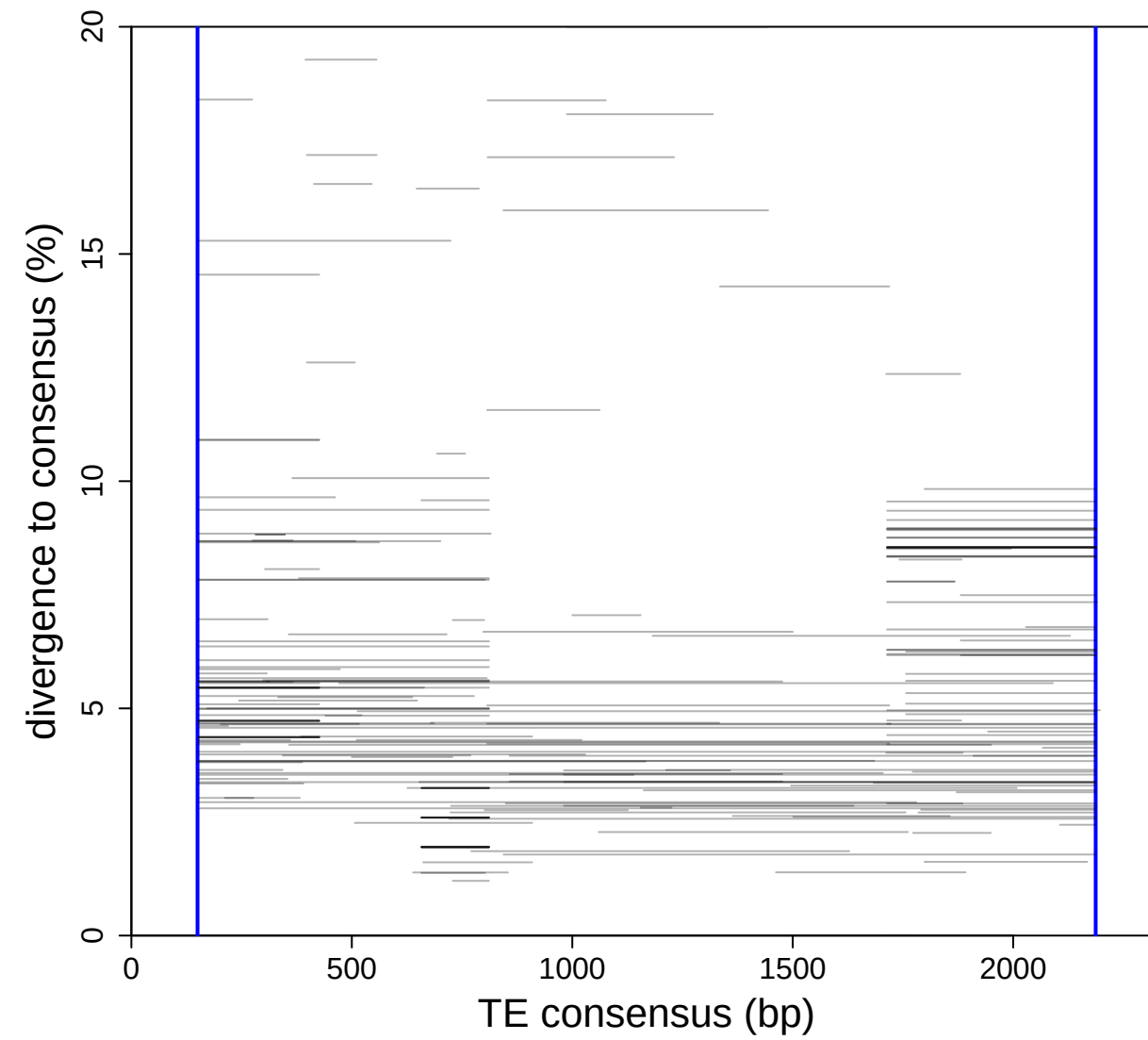
g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_ε
size: 2036bp; fragments: 224; full length: 8 (>=1832.4bp)

After TEtrimmer 2036 bp



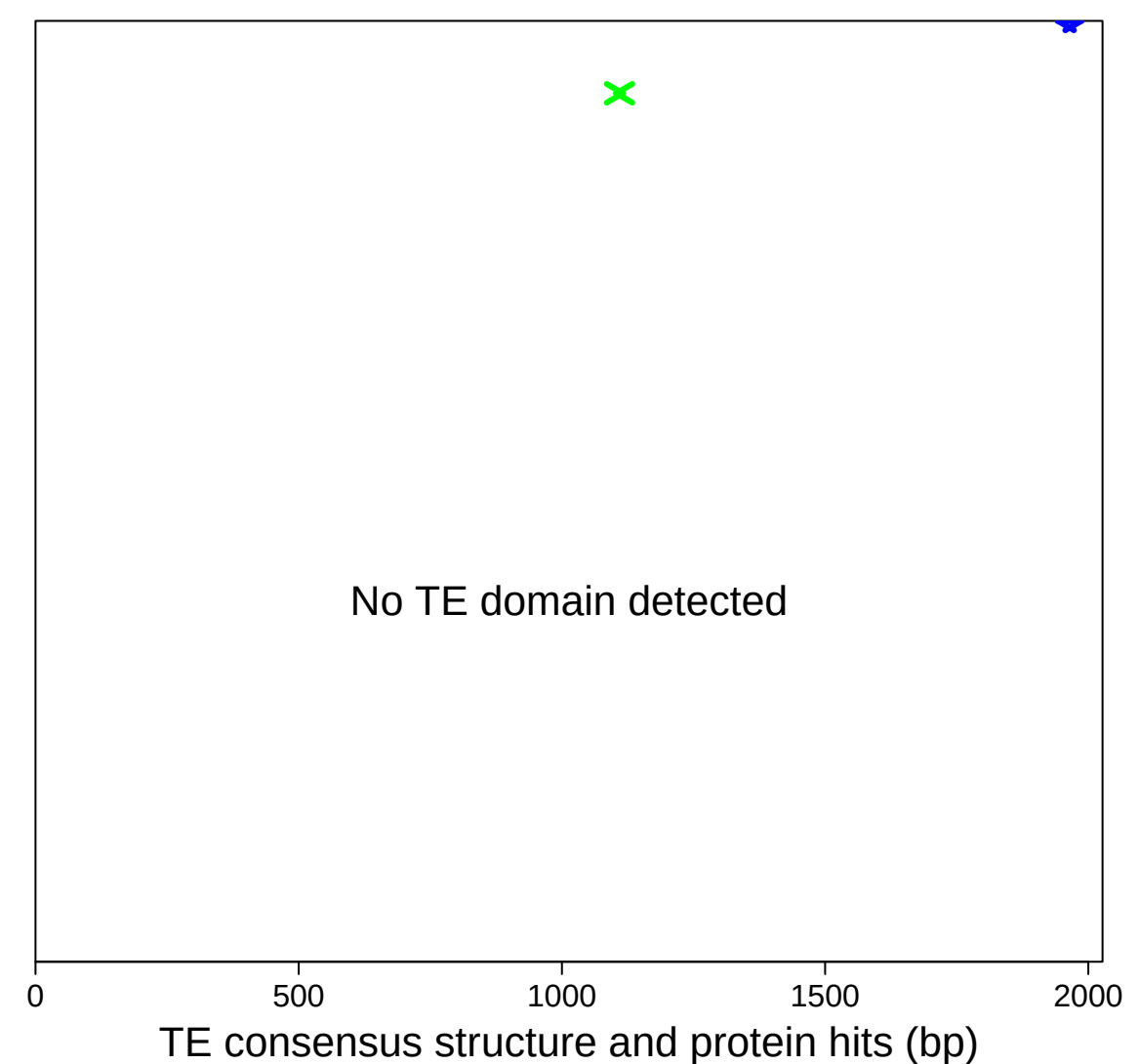
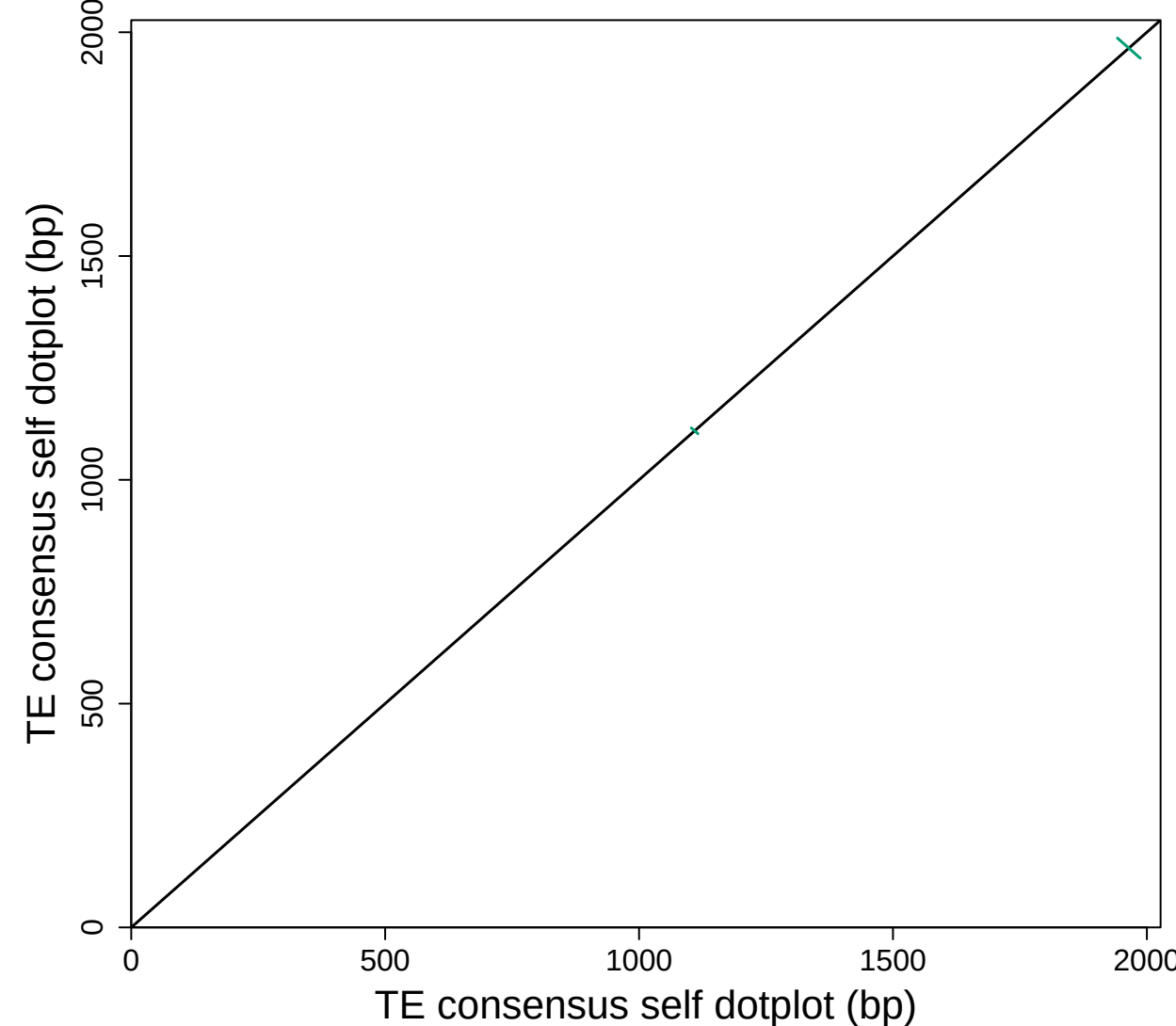
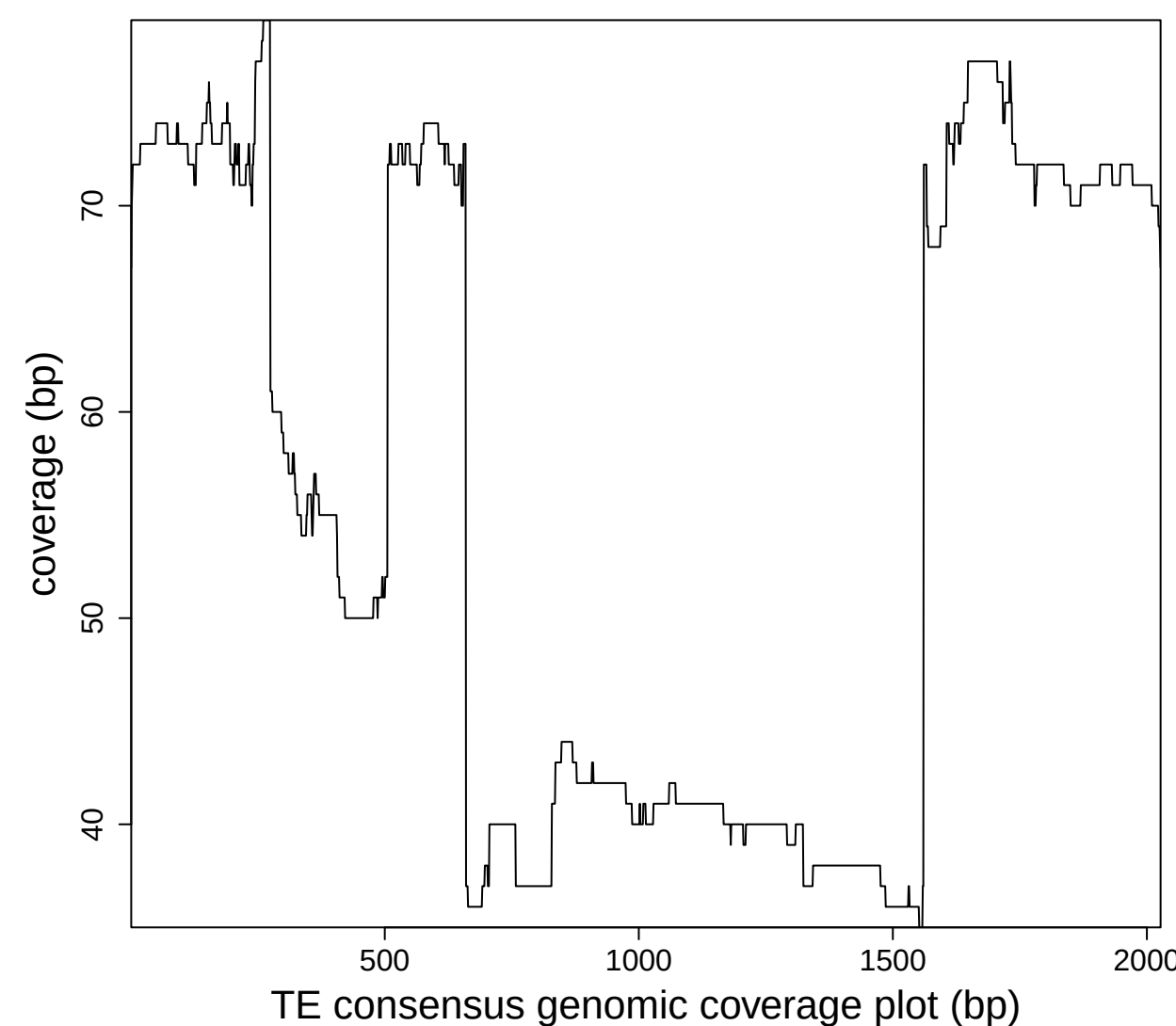
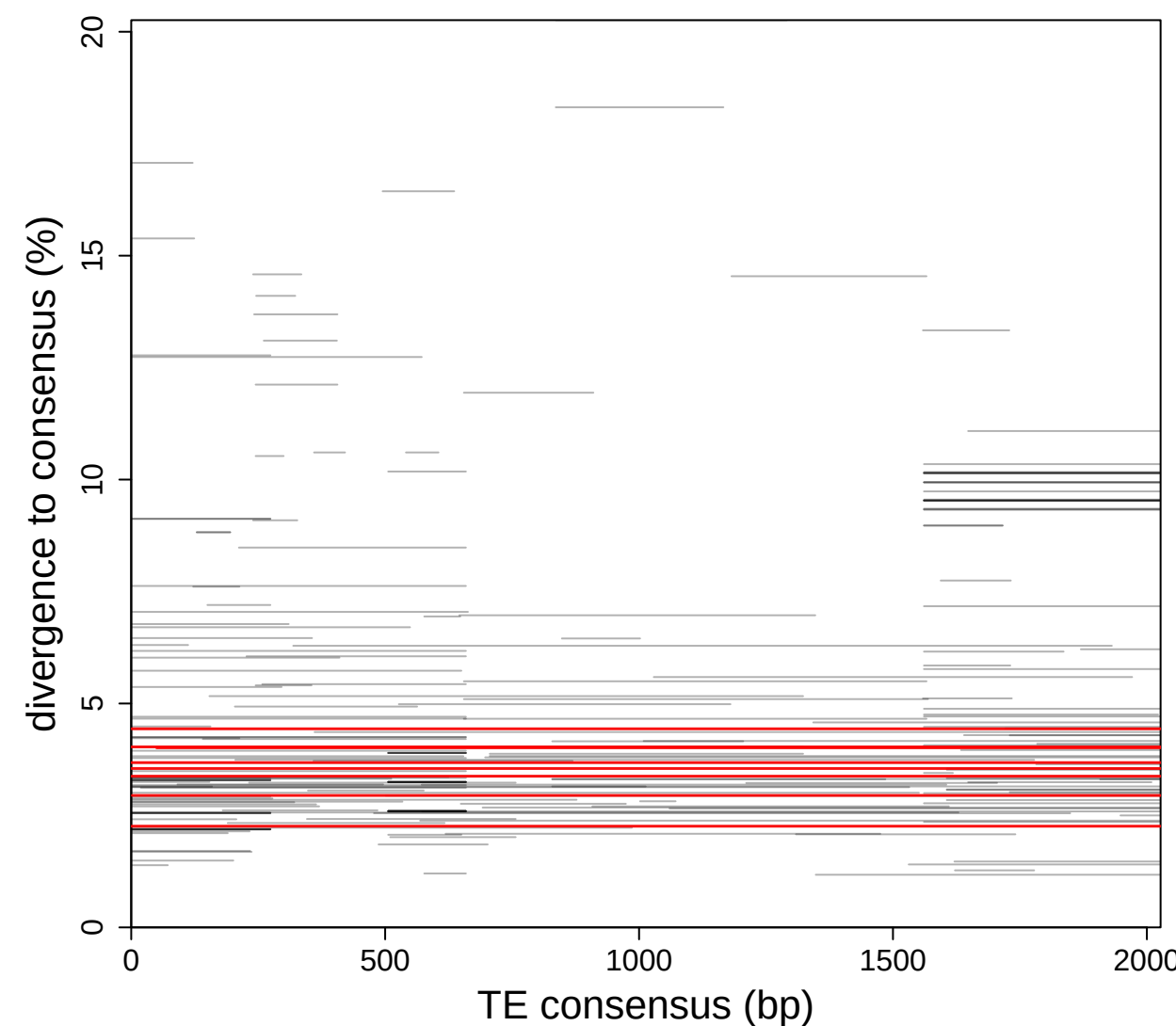
f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 2337bp; fragments: 221; full length: 0 (>=2103.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

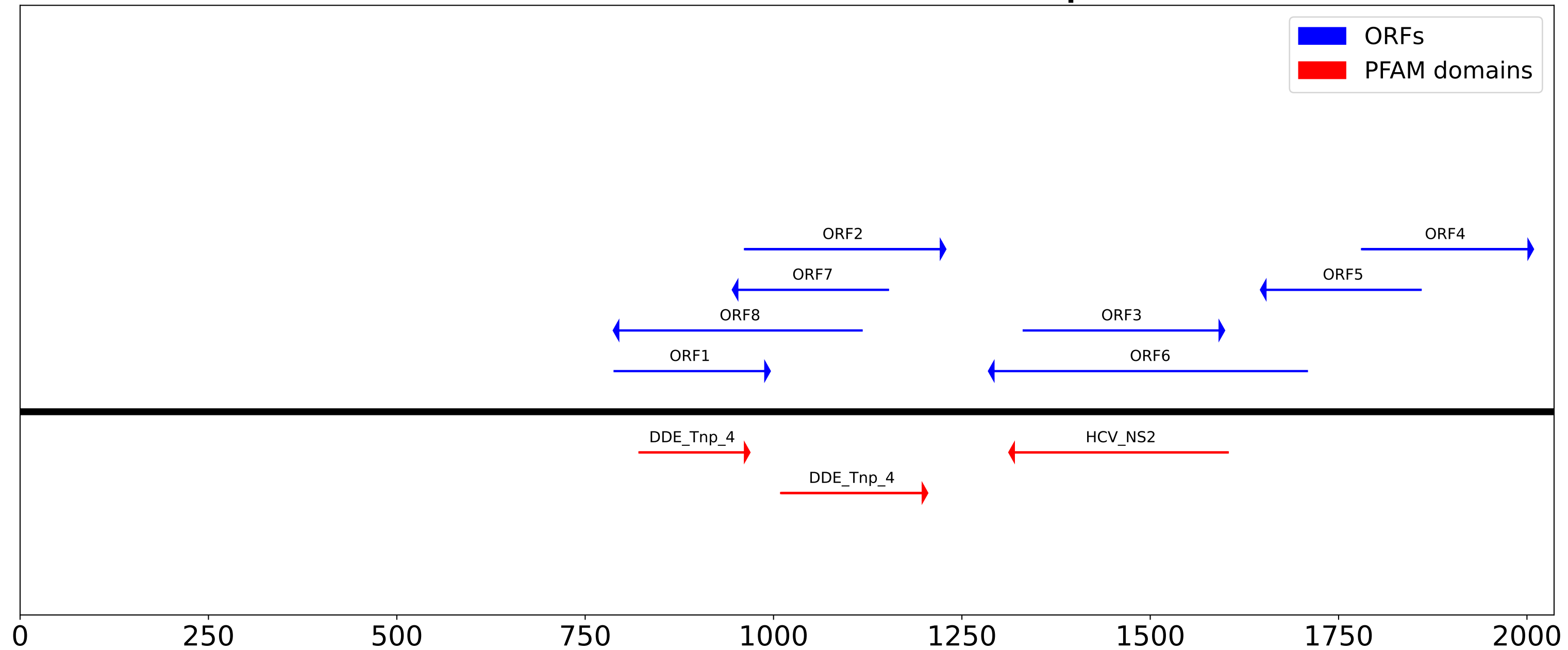


TE: rnd_5_family_1050
size: 2027bp; fragments: 224; full length: 8 (≥ 1824.3 bp)

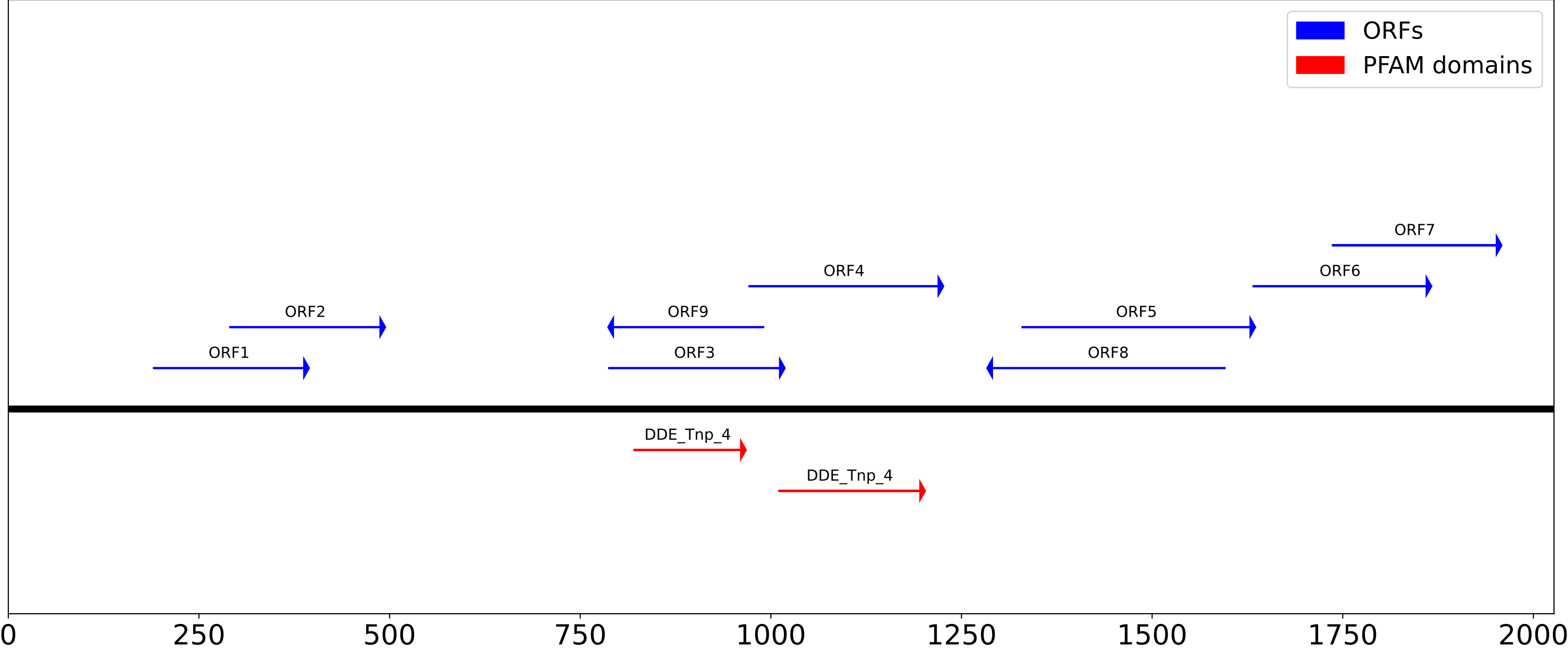
Before TEtrimmer 2027 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

