

MSA length = 5029

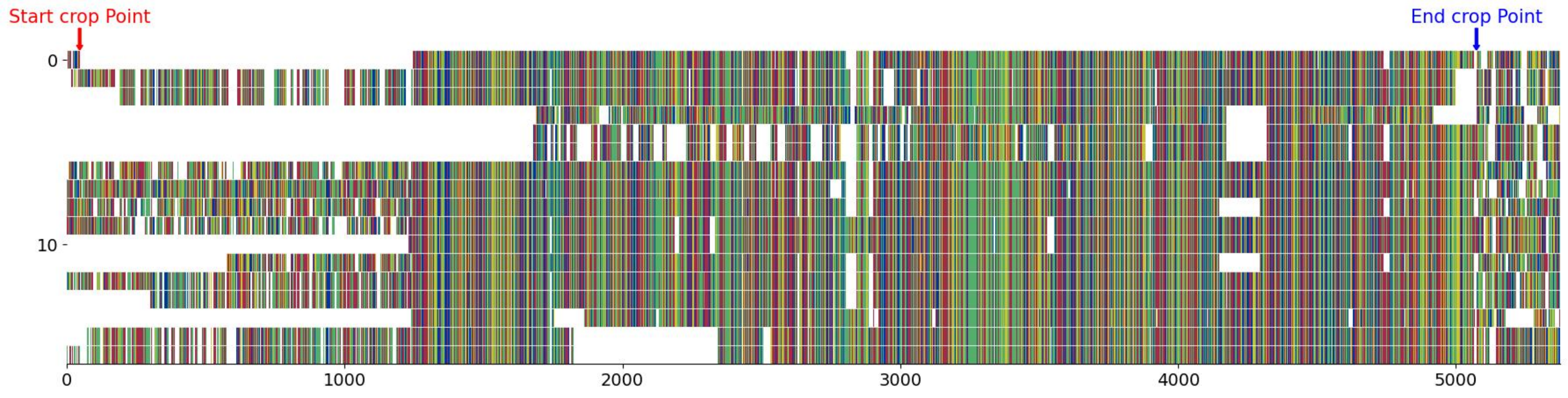
Start crop Point

End crop Point

MSA length = 5029

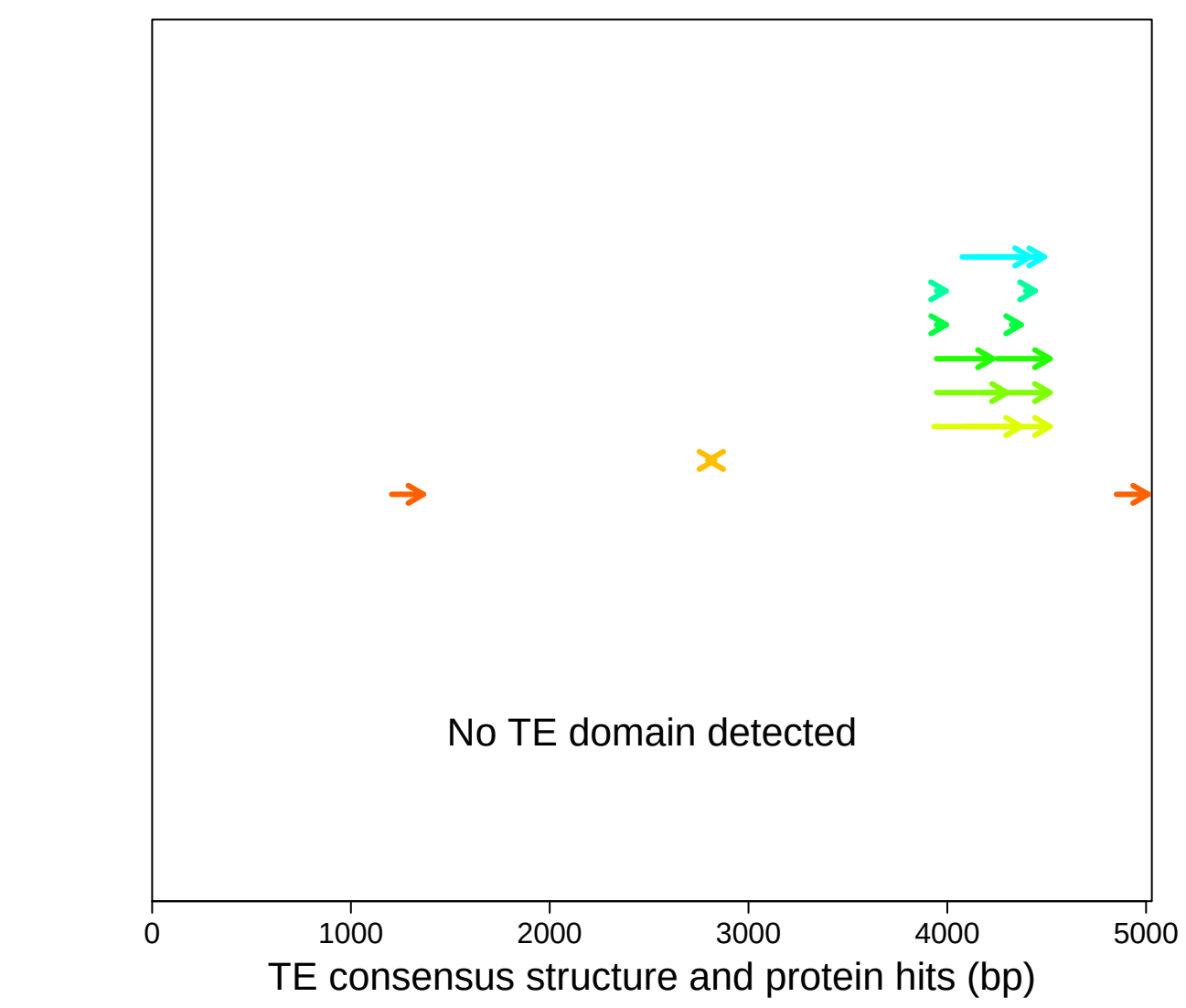
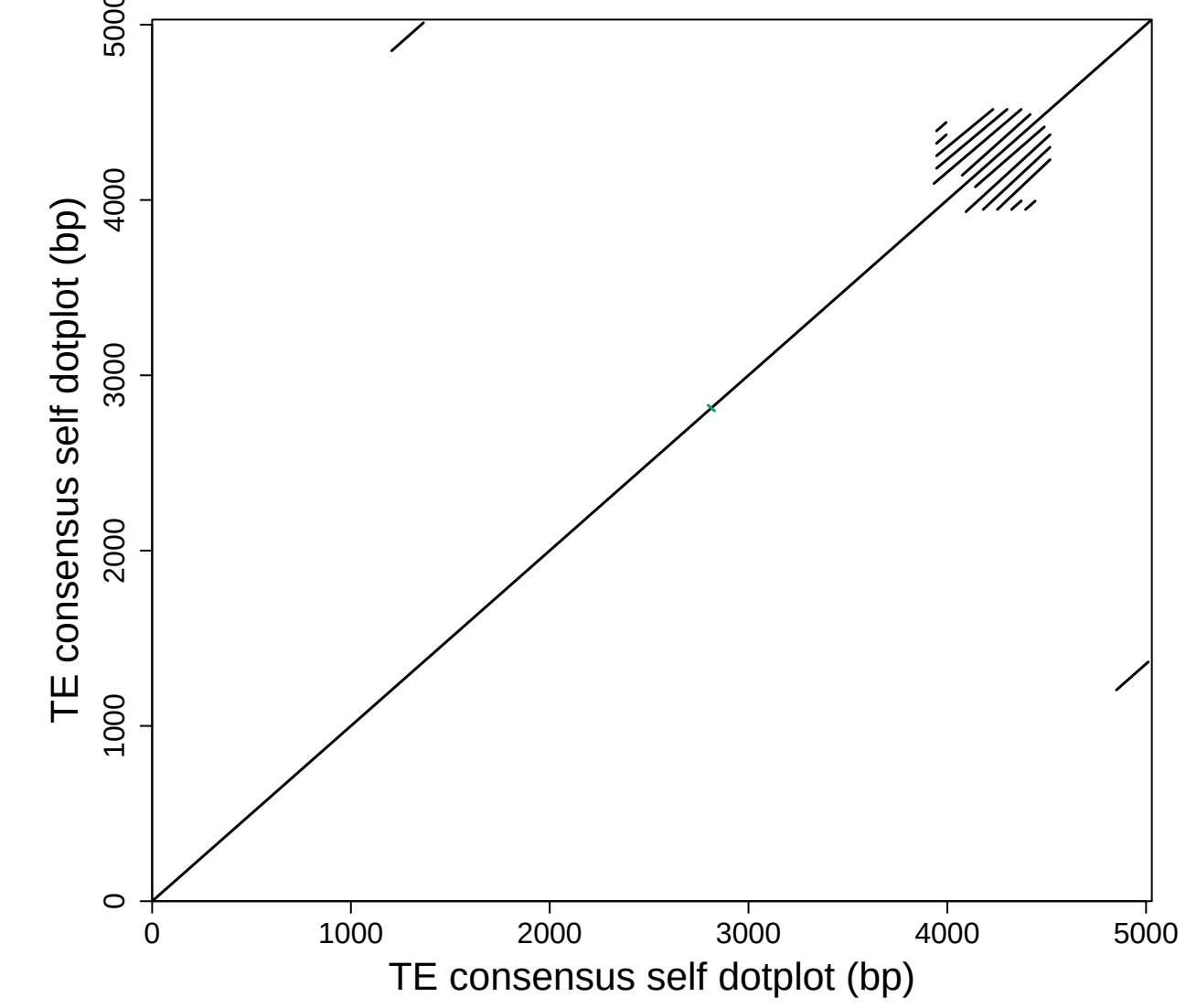
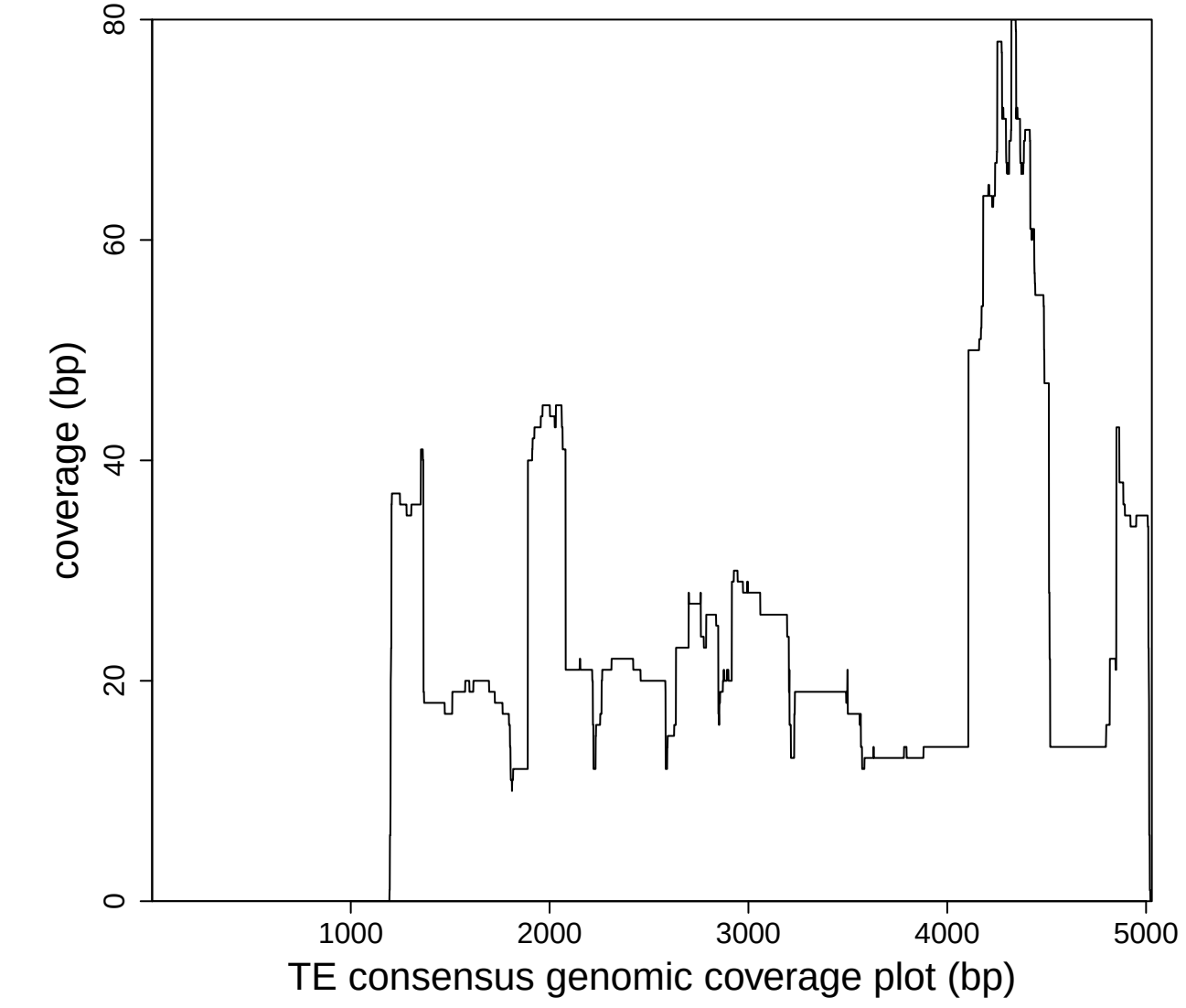
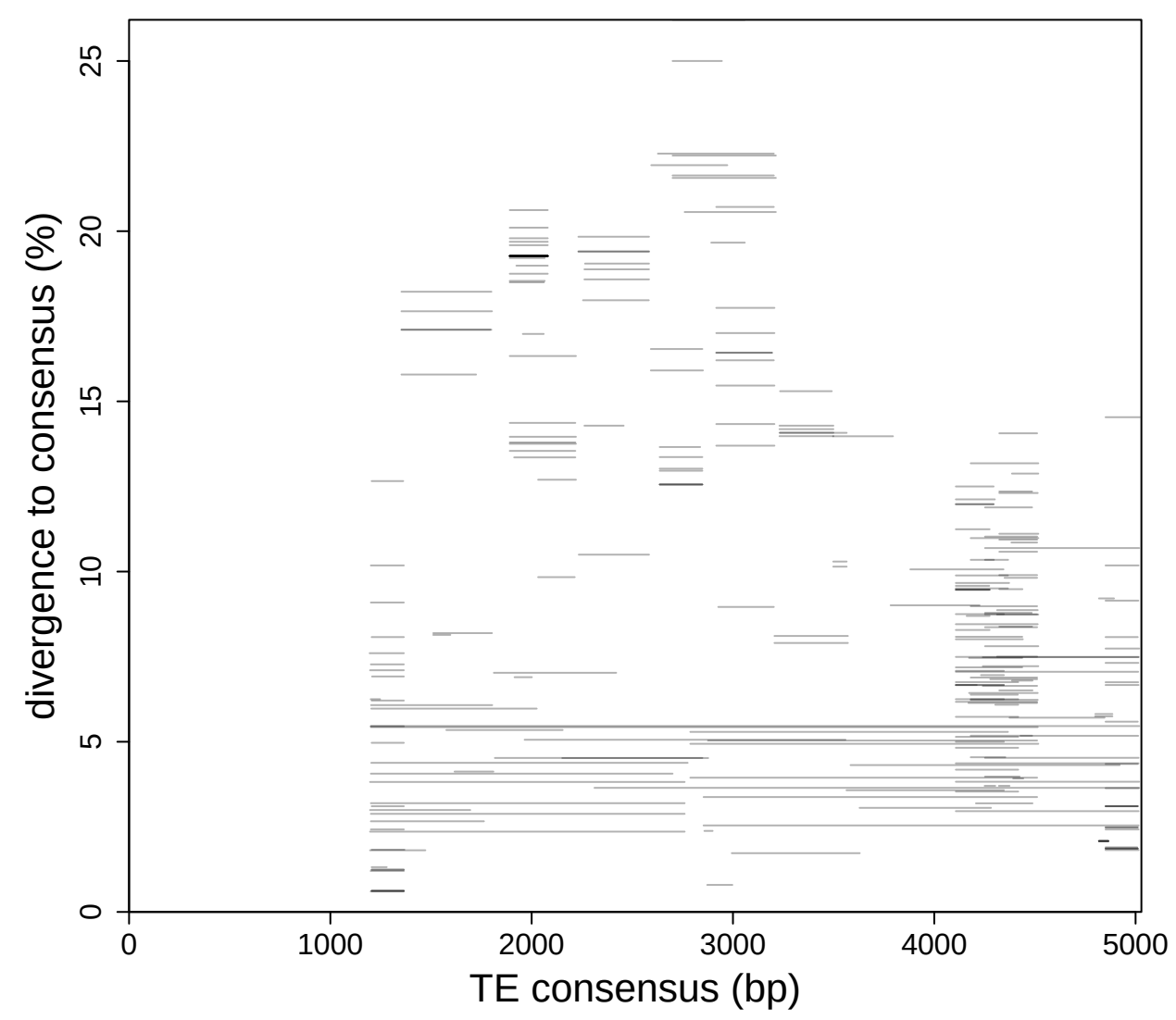
Start crop Point

End crop Point



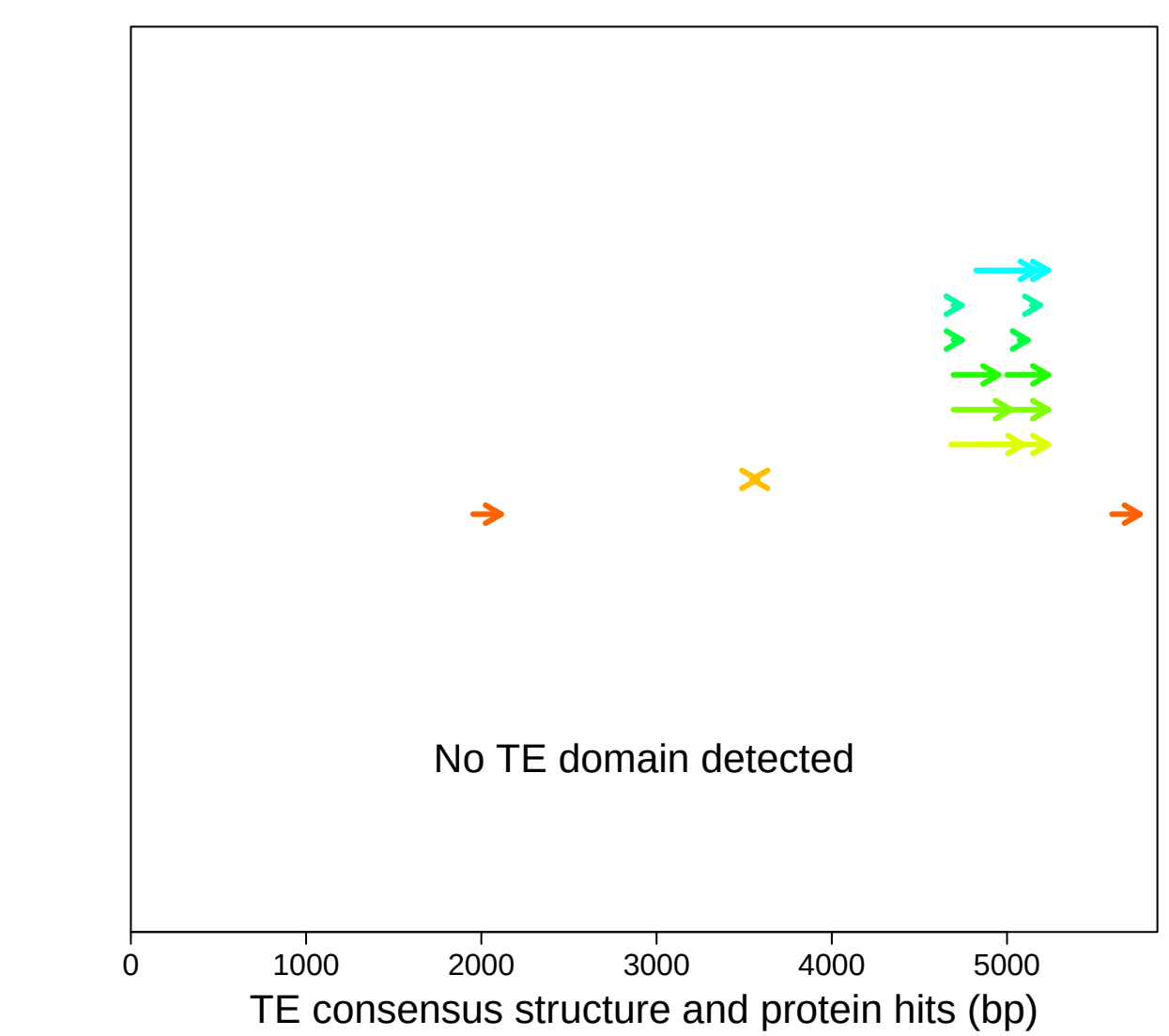
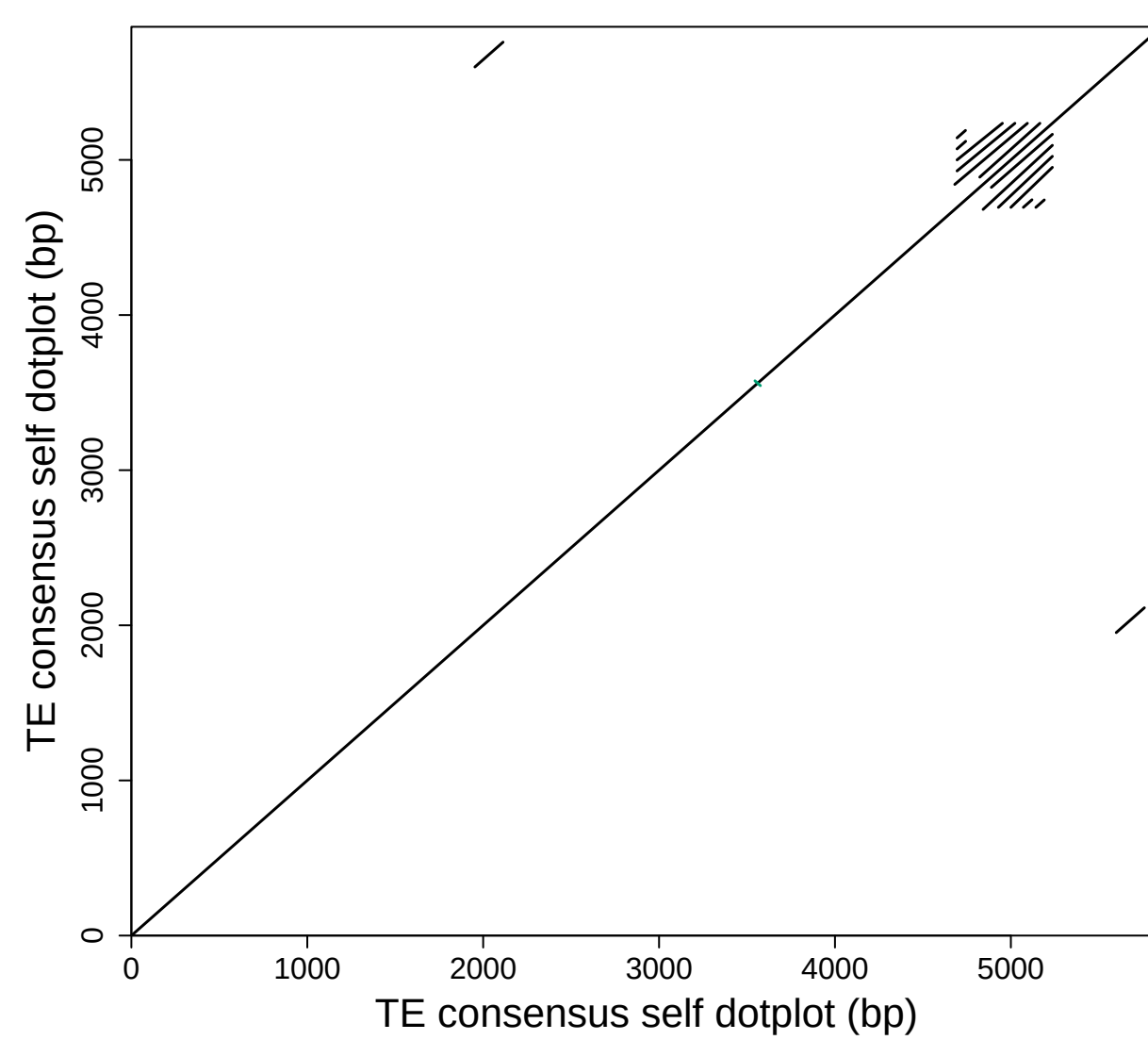
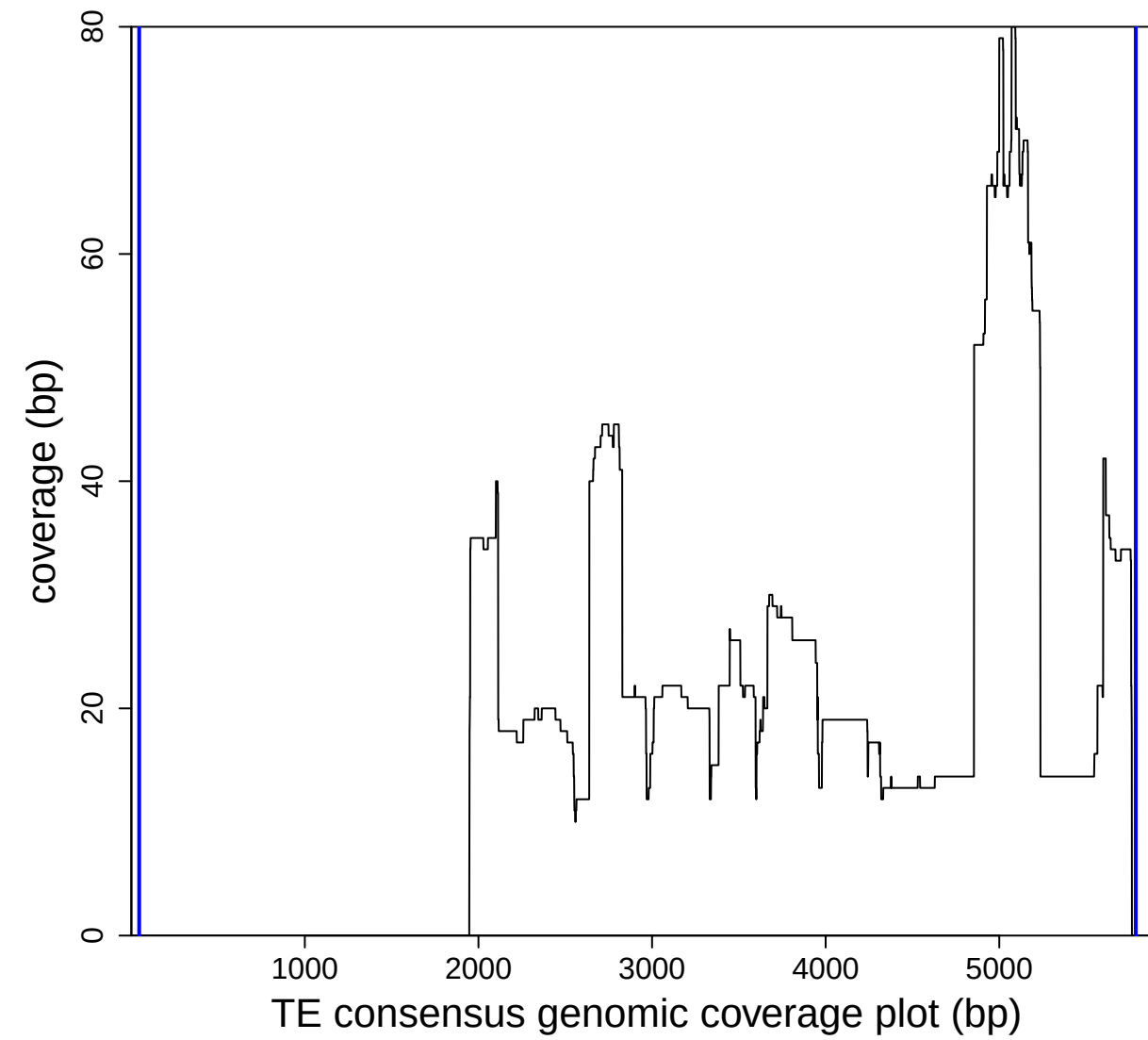
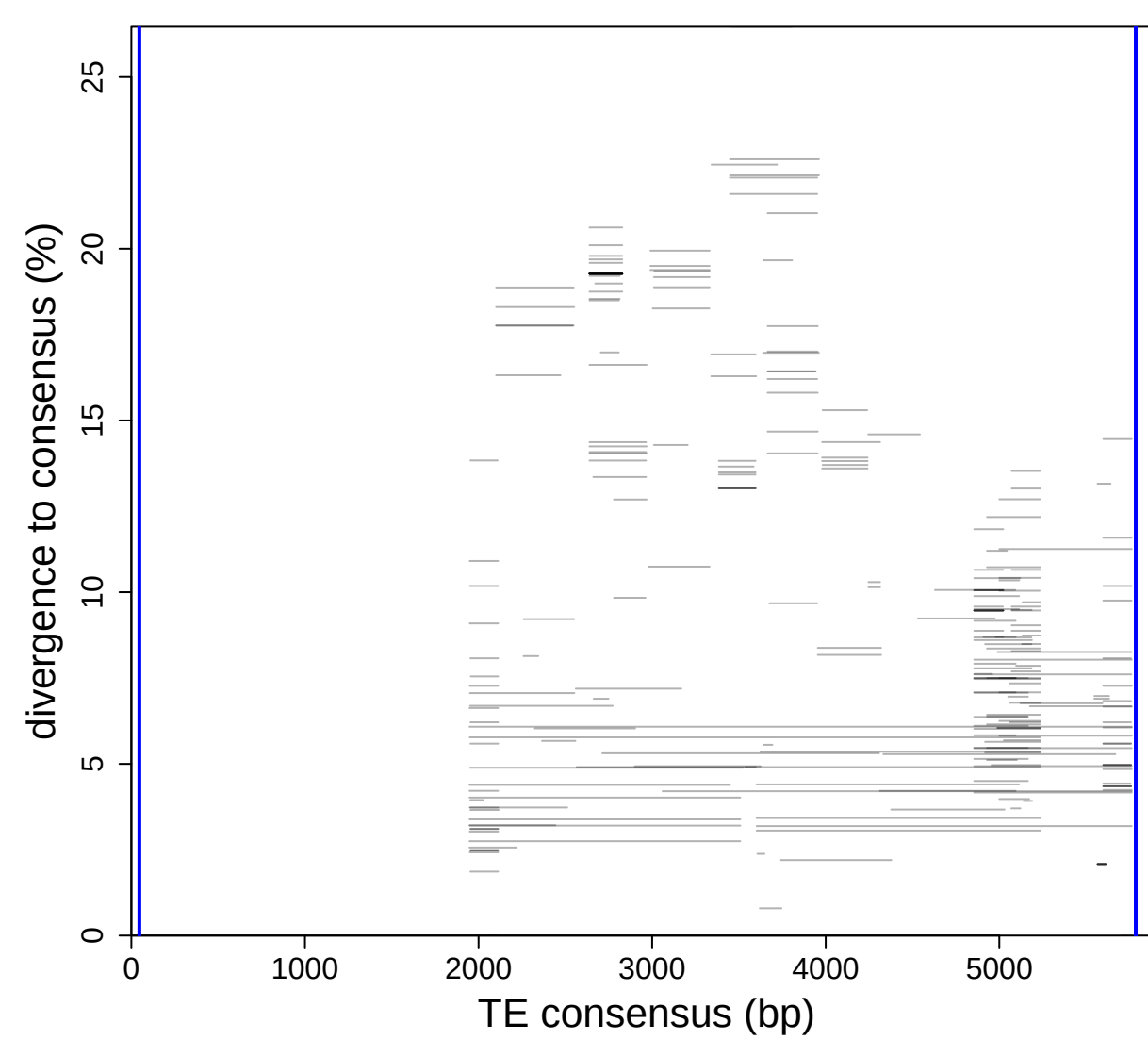
_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_al
size: 5029bp; fragments: 268; full length: 0 (>=4526.1bp)

After TEtrimmer 5029 bp



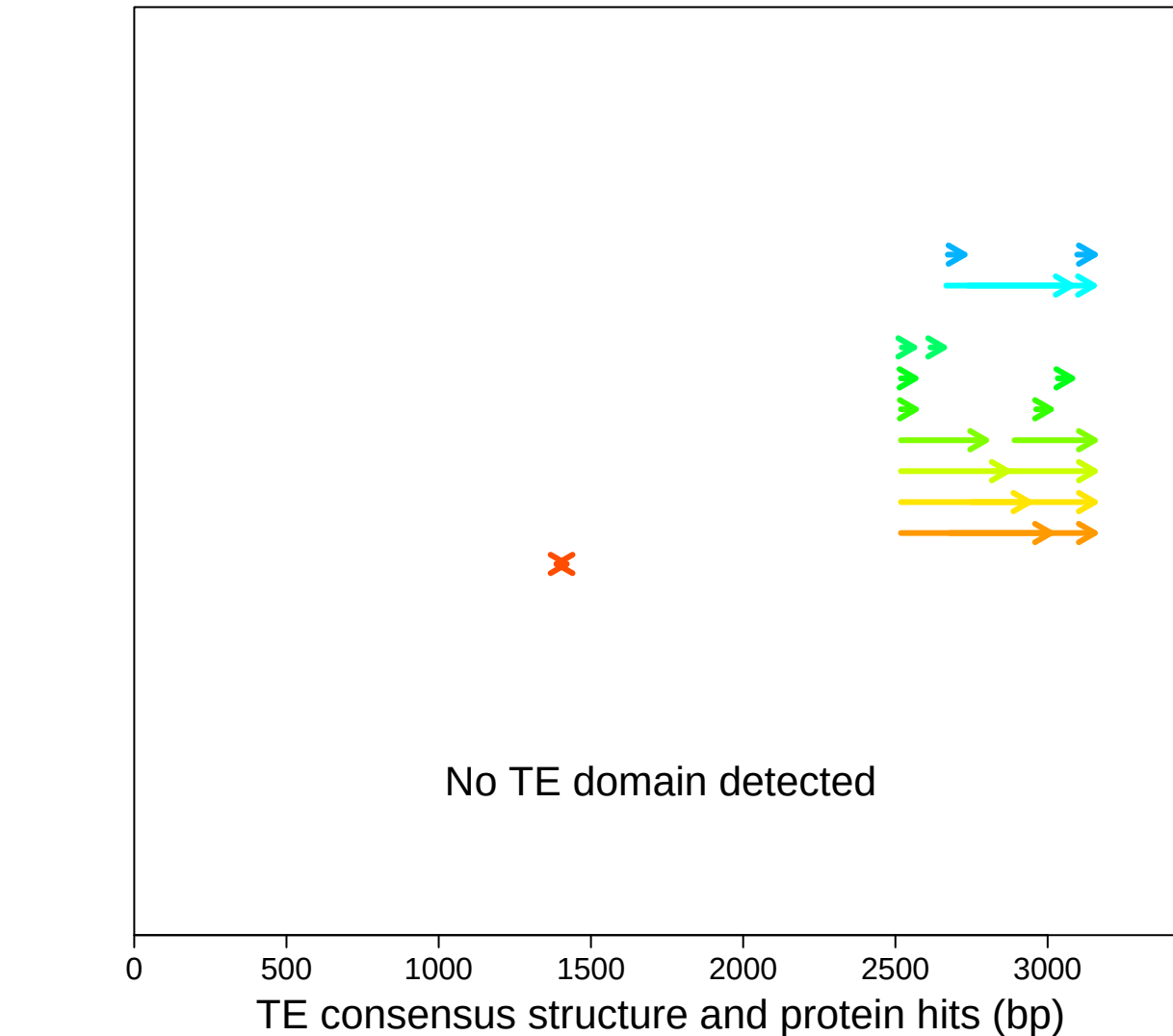
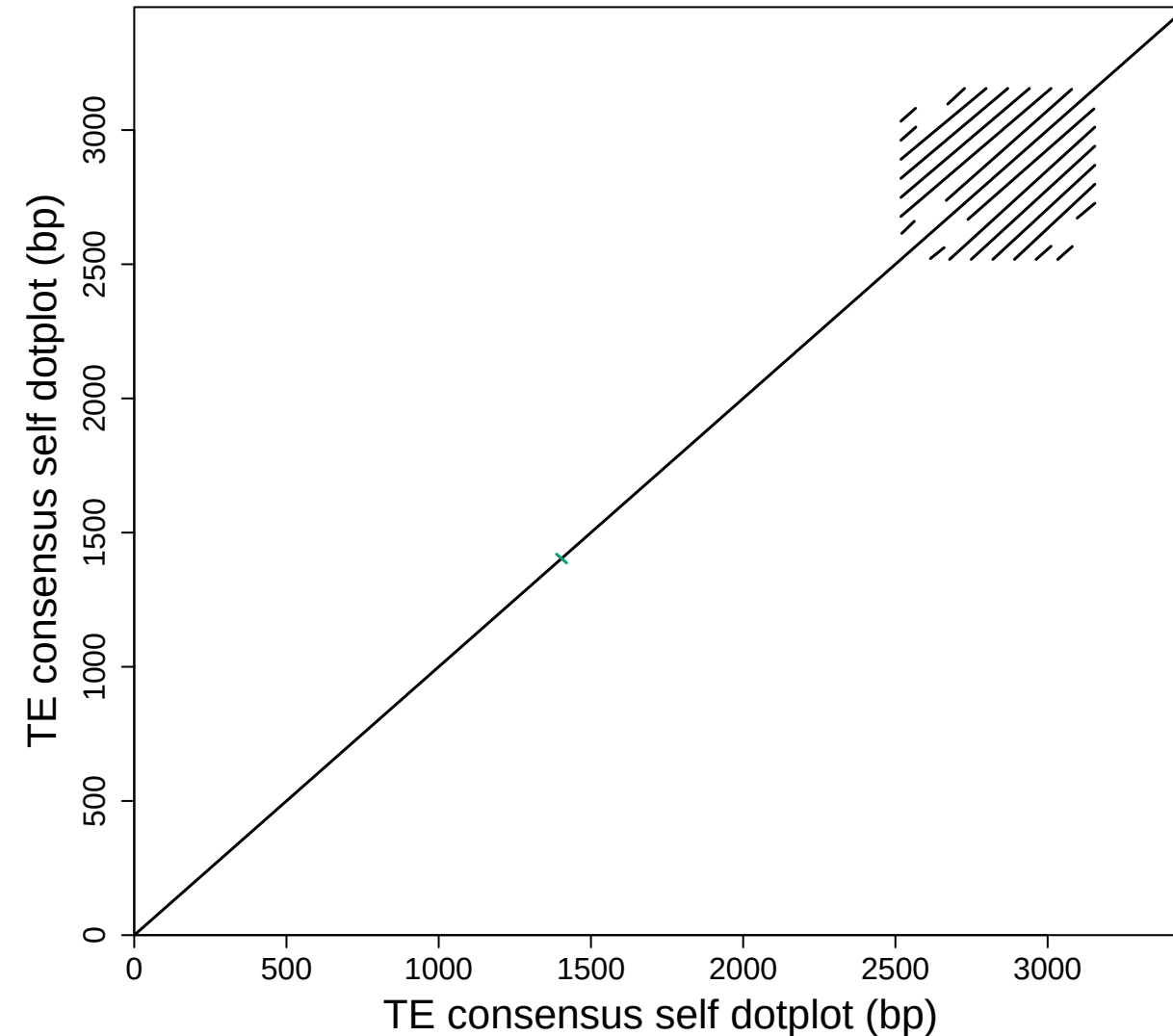
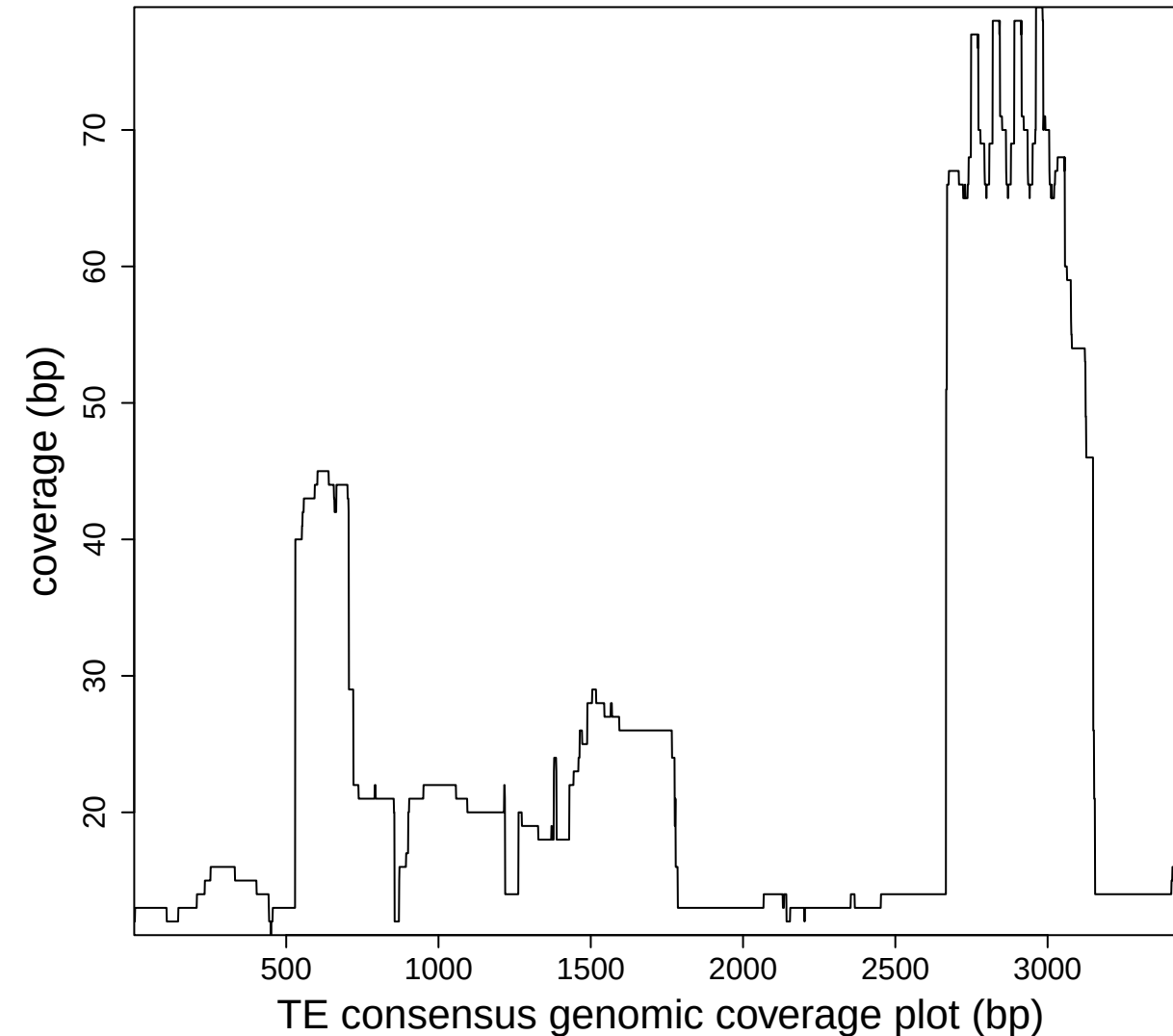
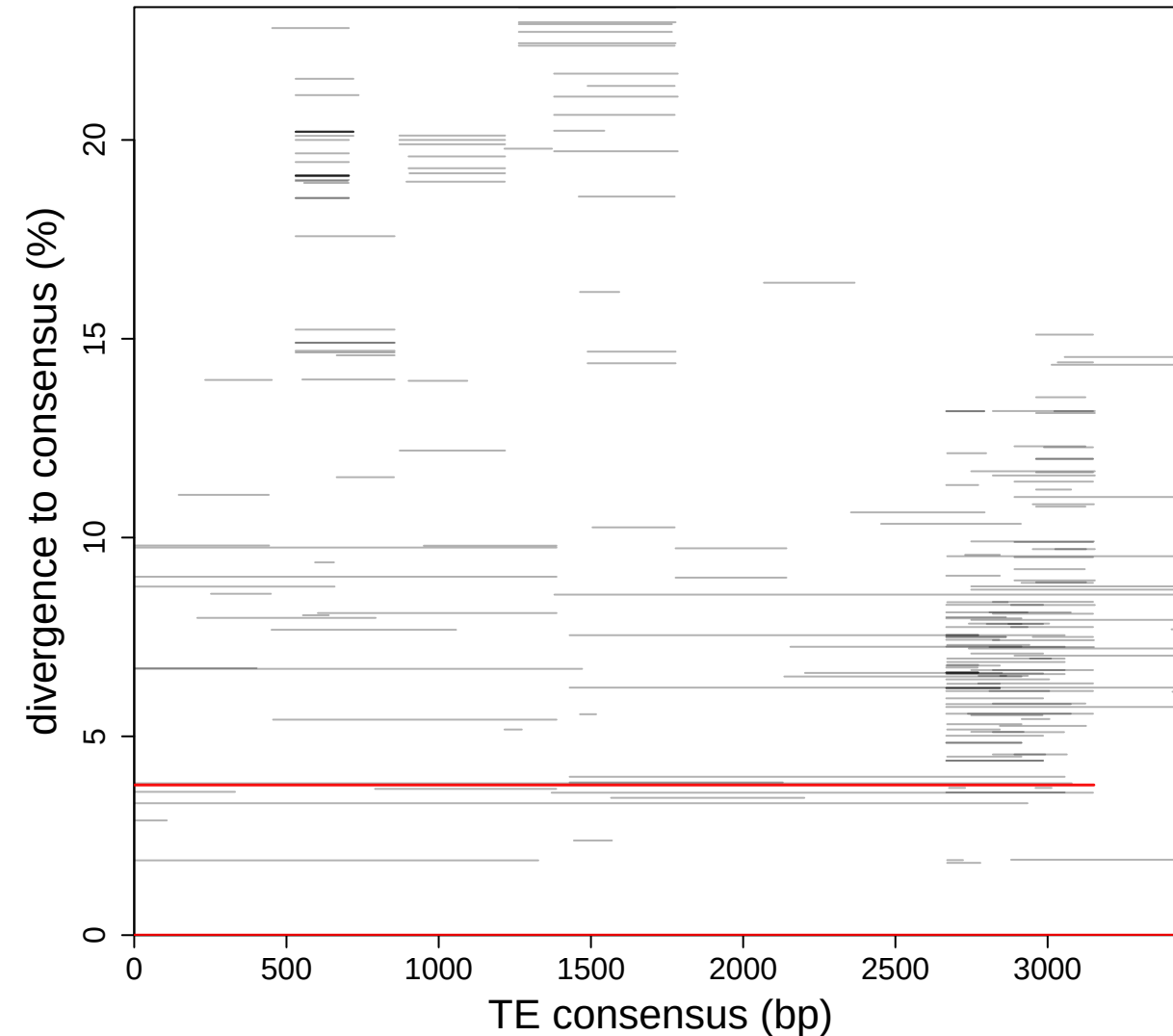
bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 5936bp; fragments: 266; full length: 0 (>=5342.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

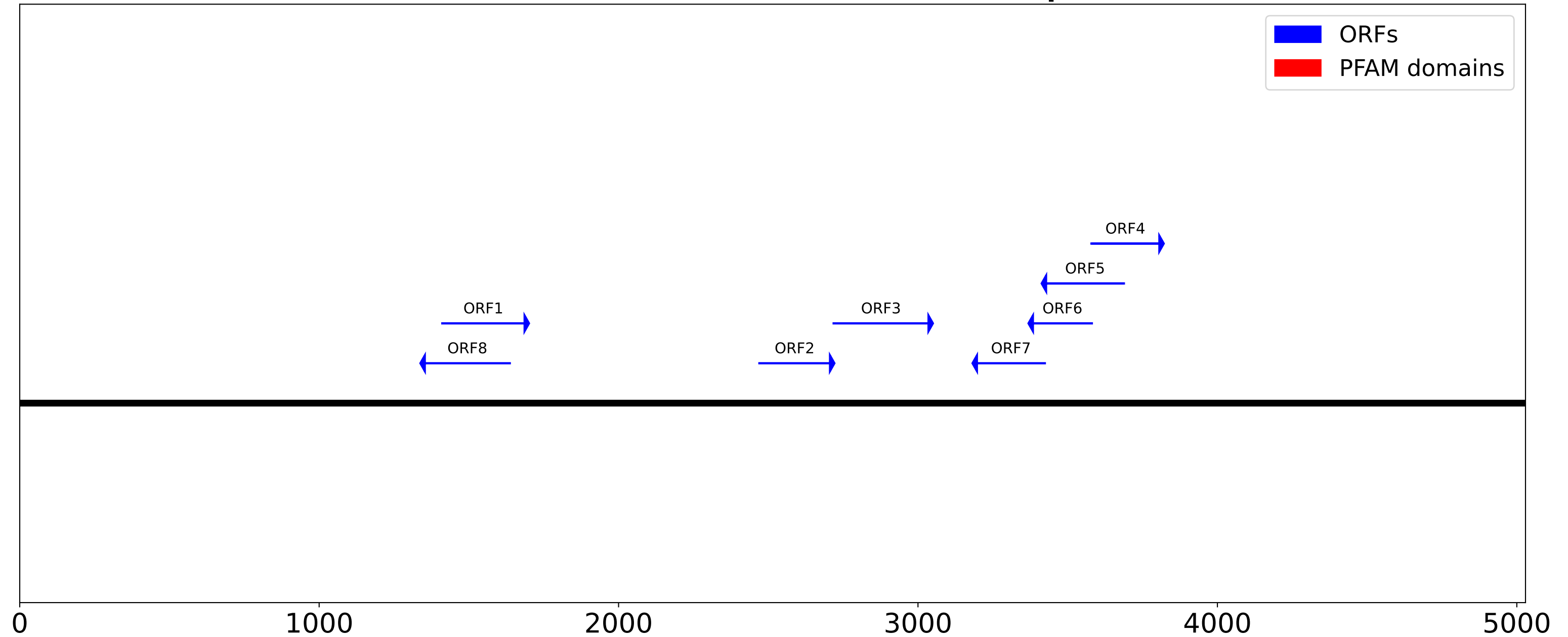


TE: ltr_1_family_918
size: 3458bp; fragments: 221; full length: 2 (>=3112.2bp)

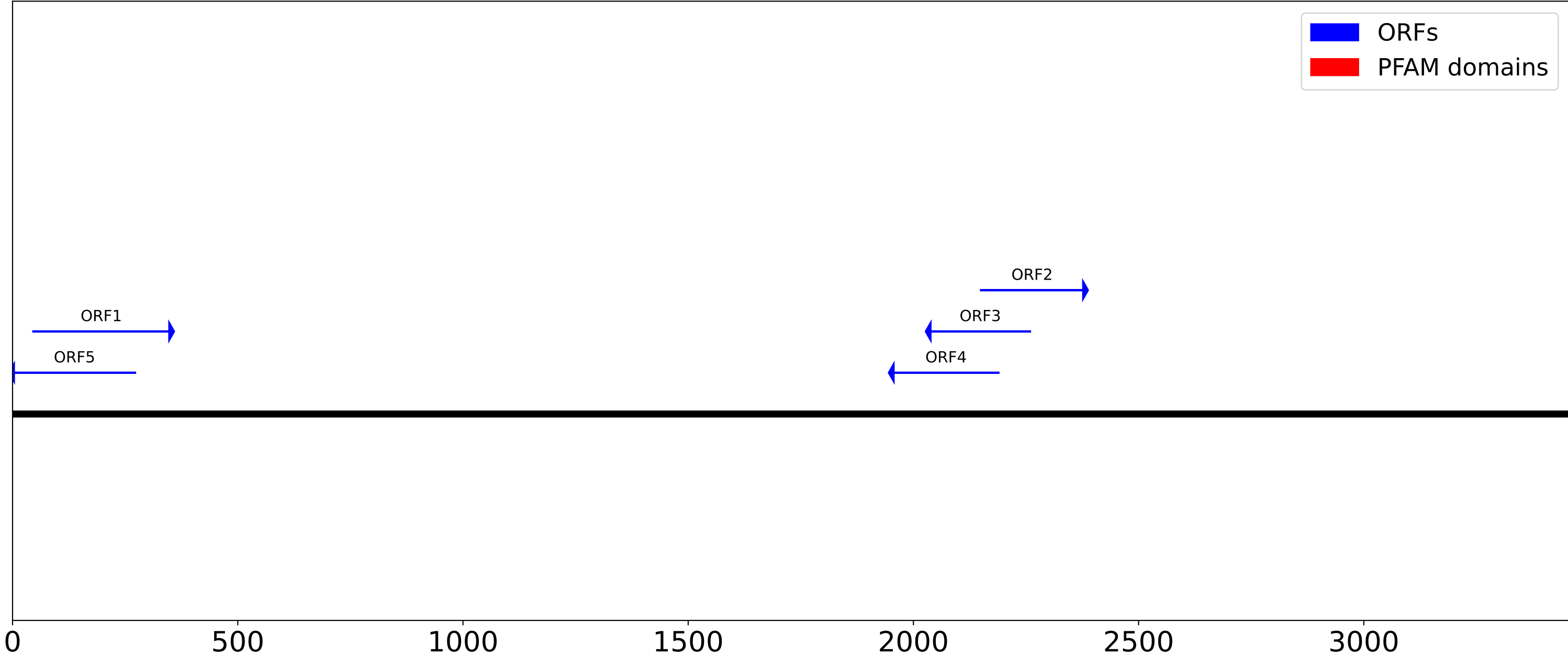
Before TEtrimmer 3458 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

