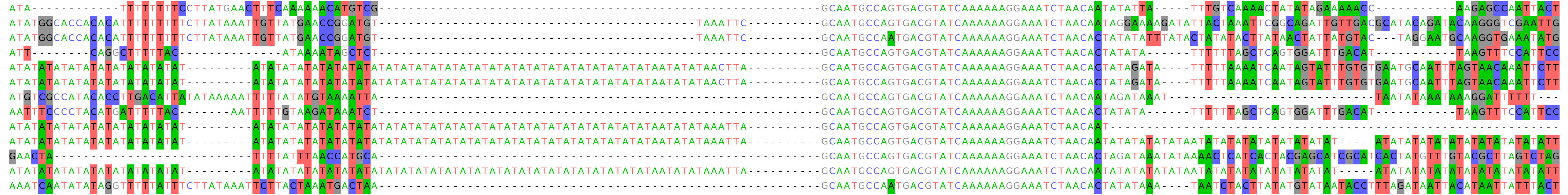
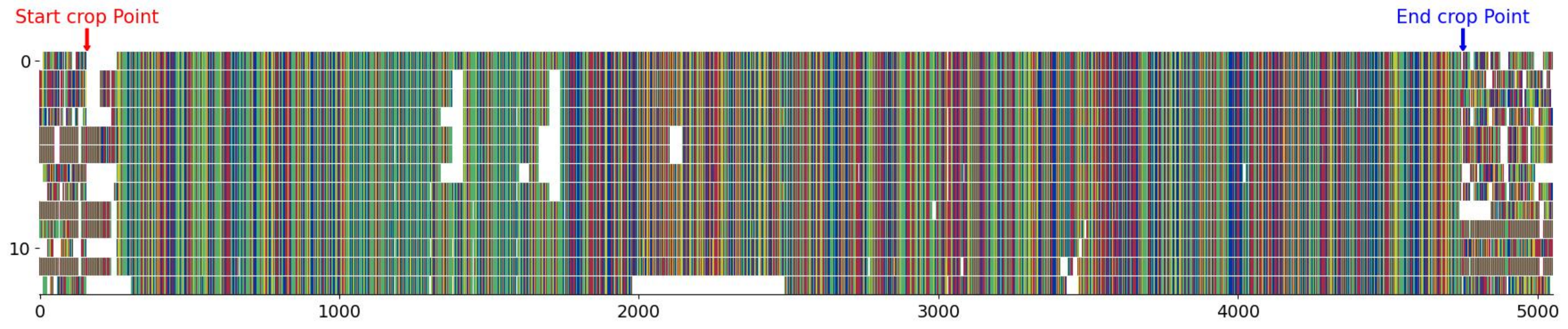


Start crop Point

End crop Point

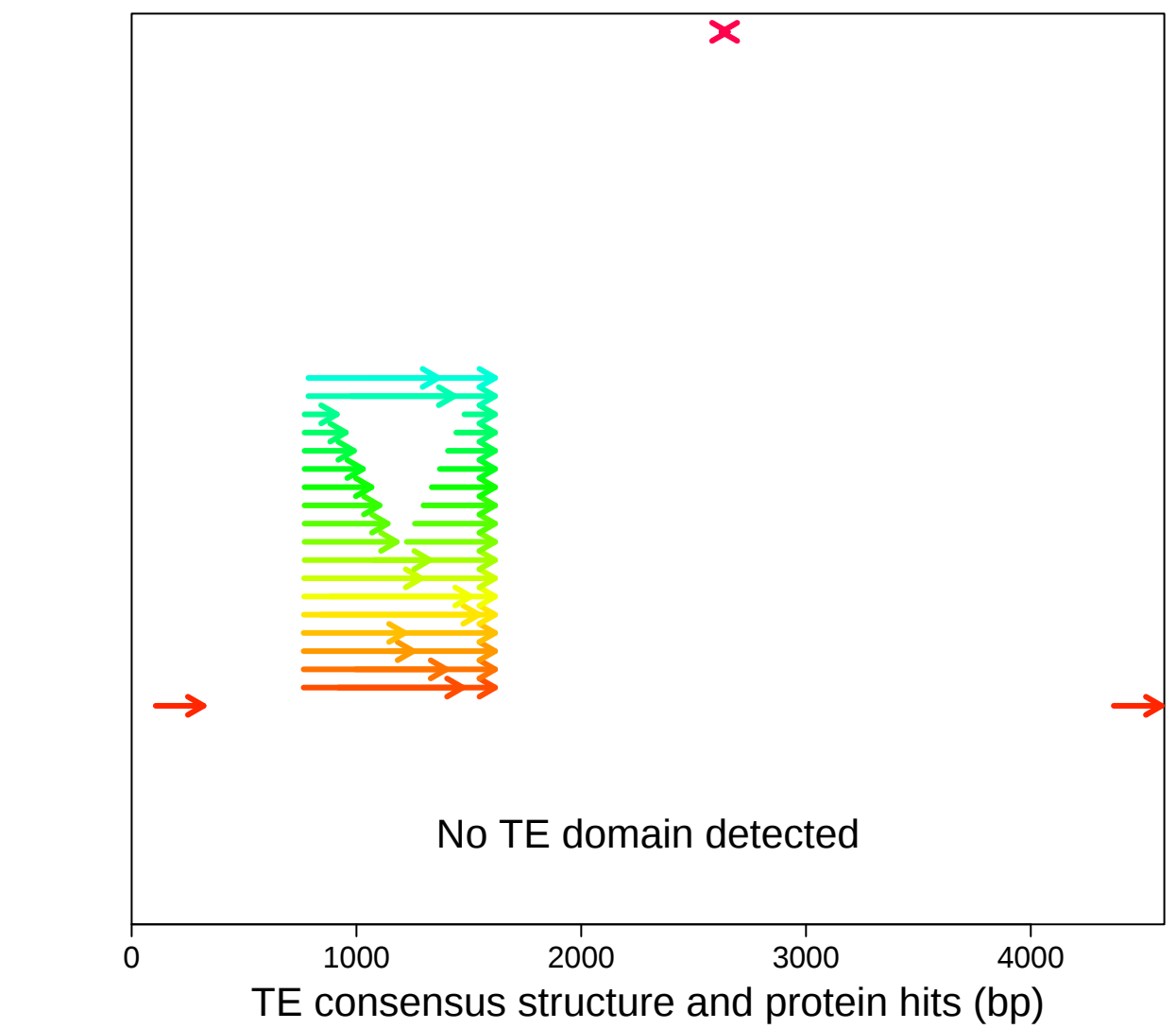
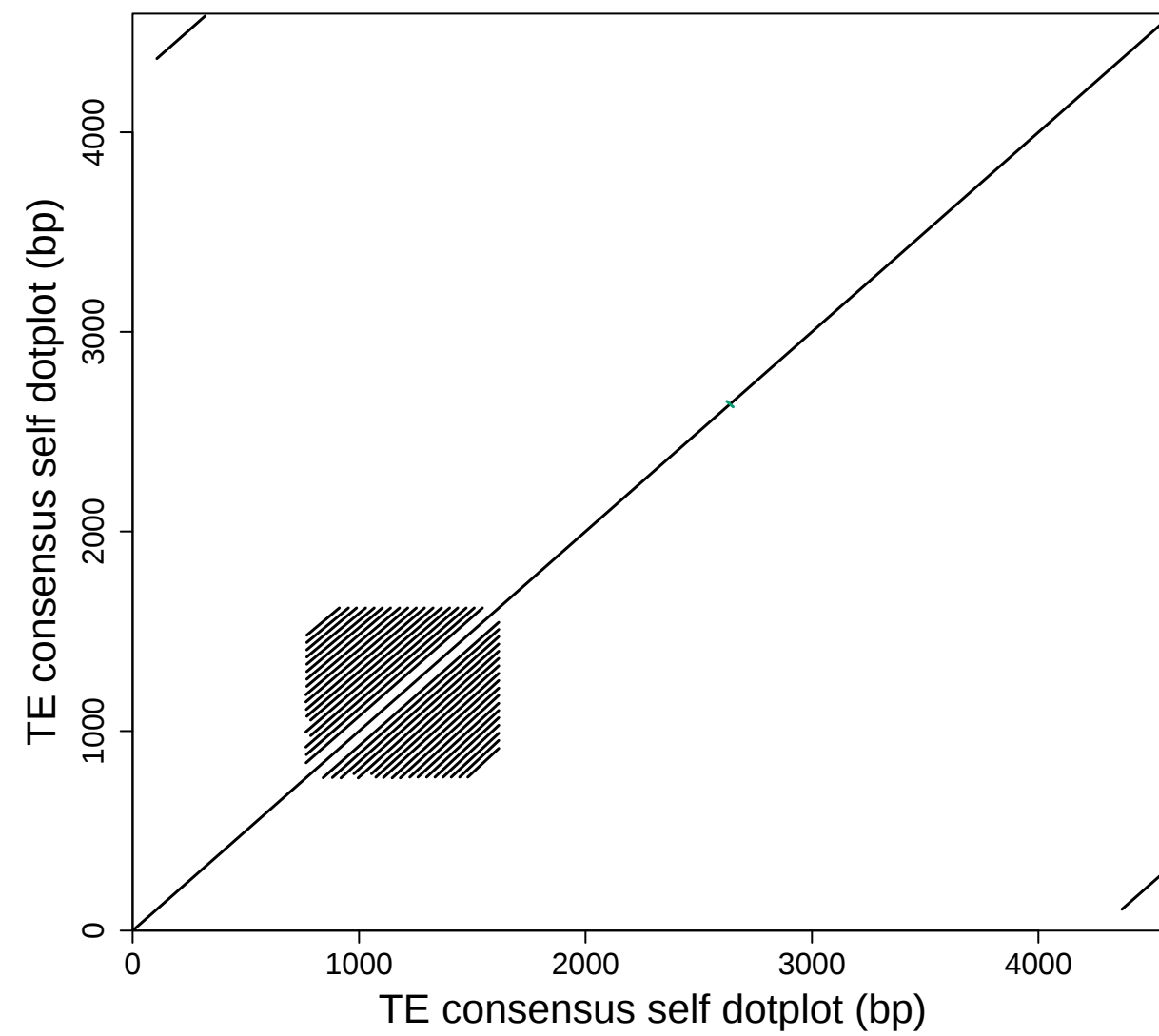
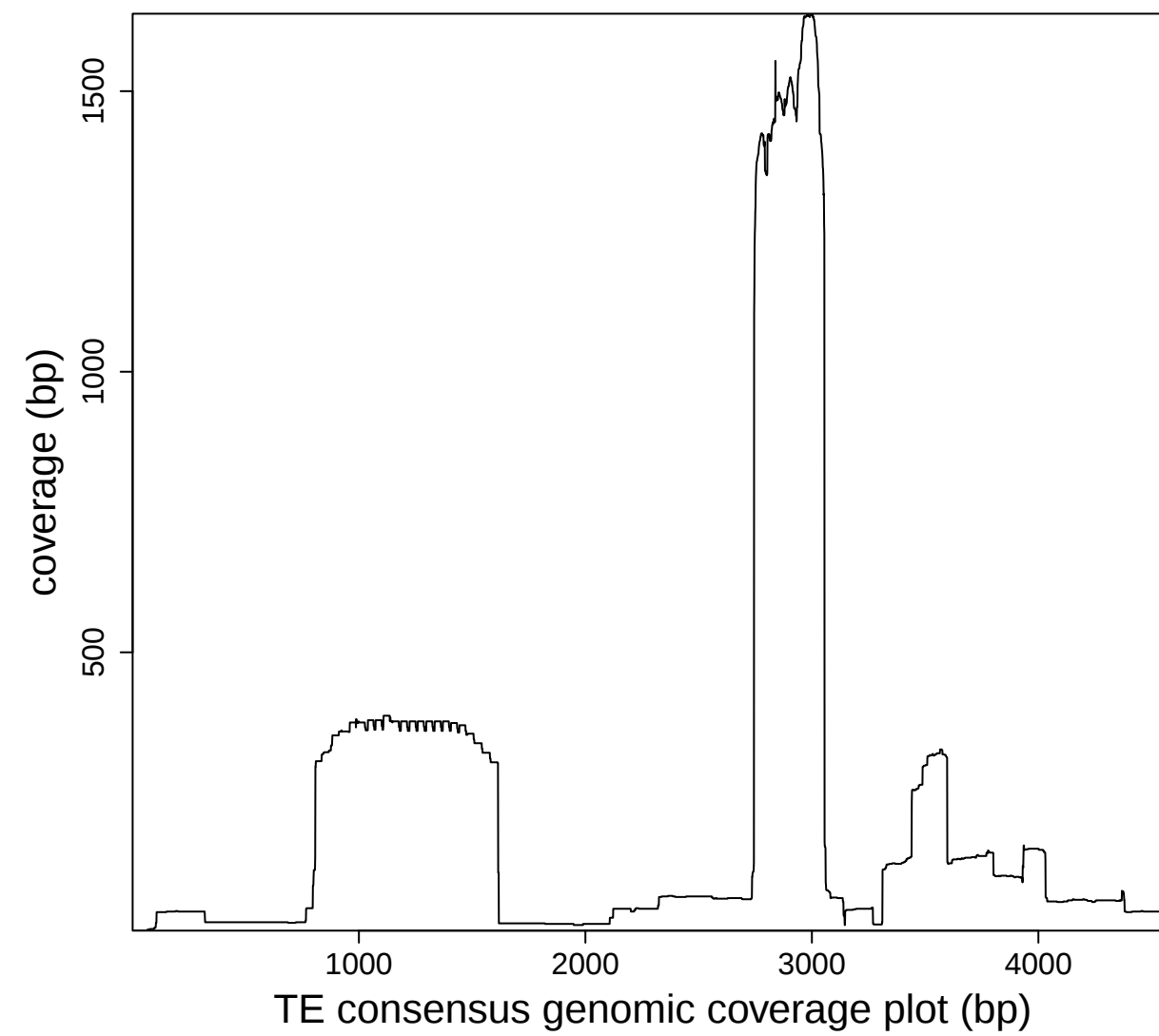
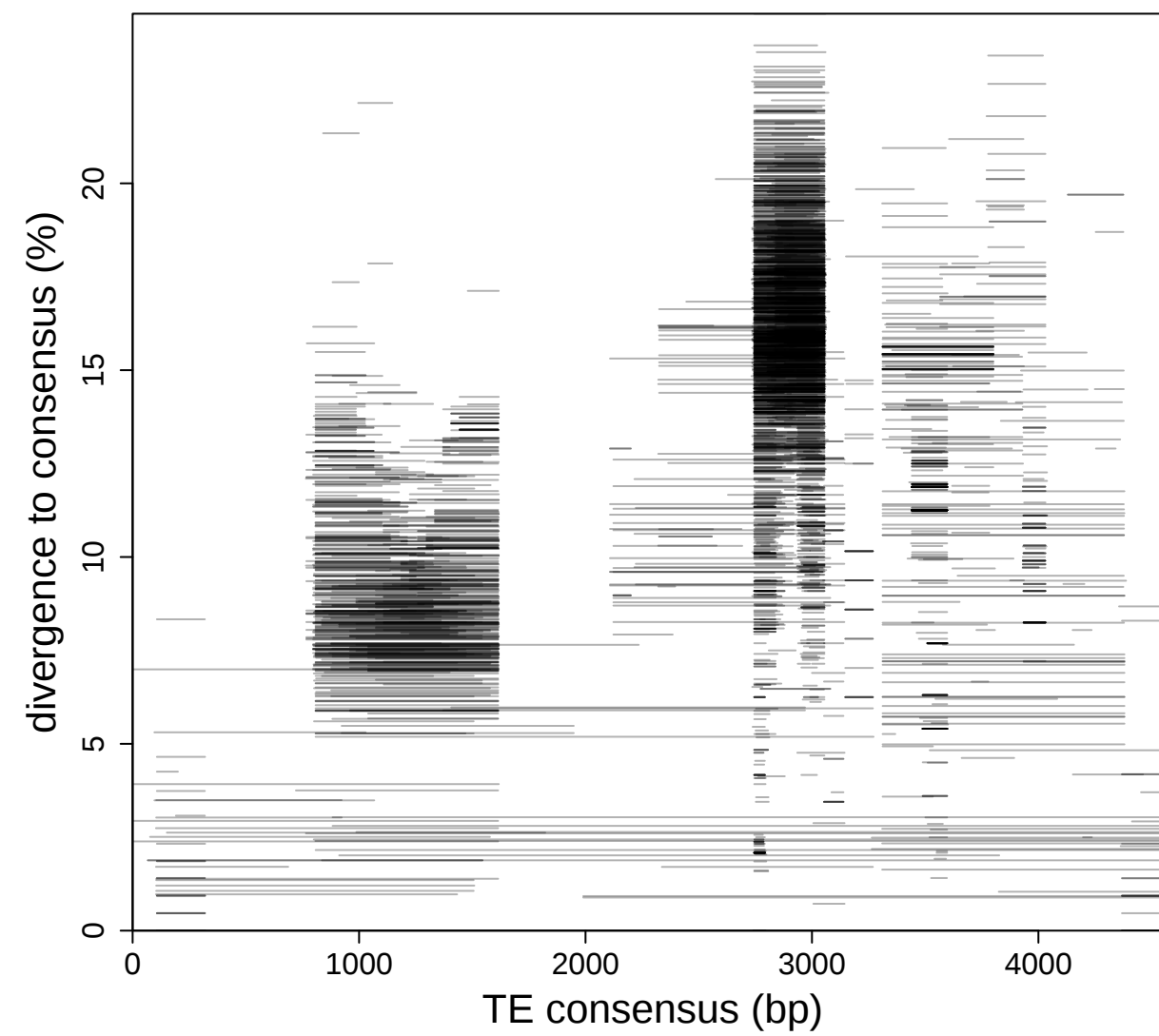
MSA length = 4594





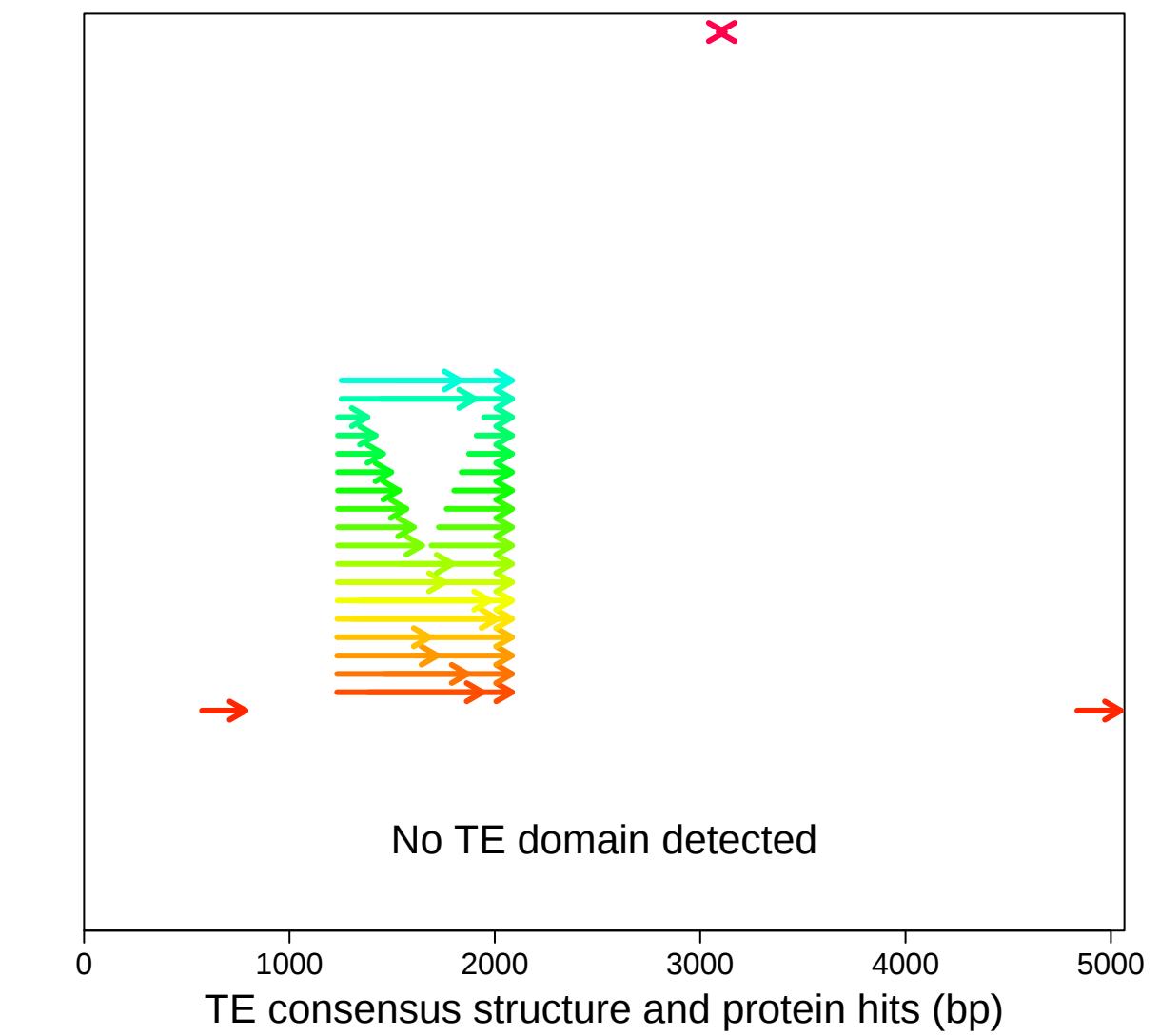
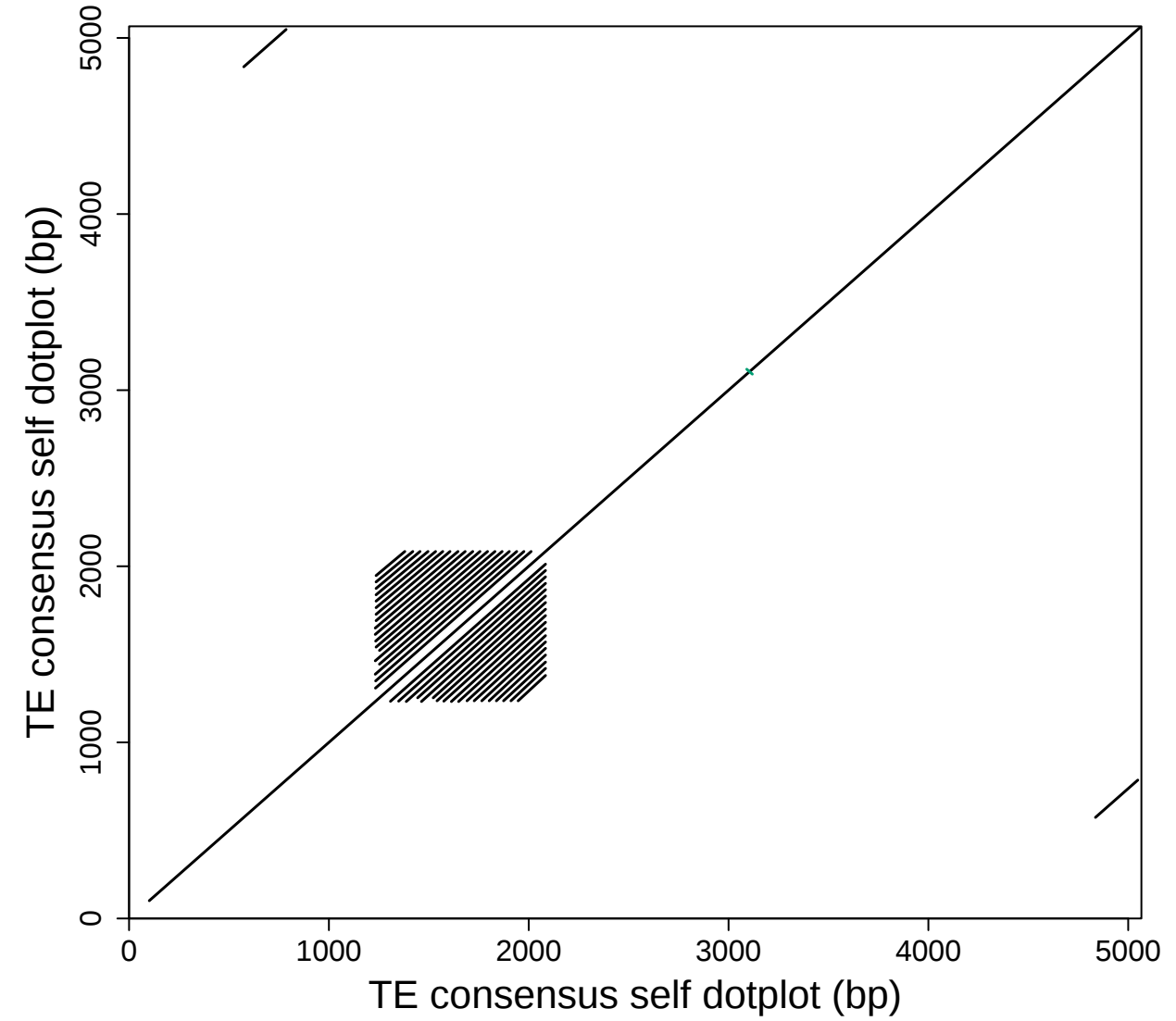
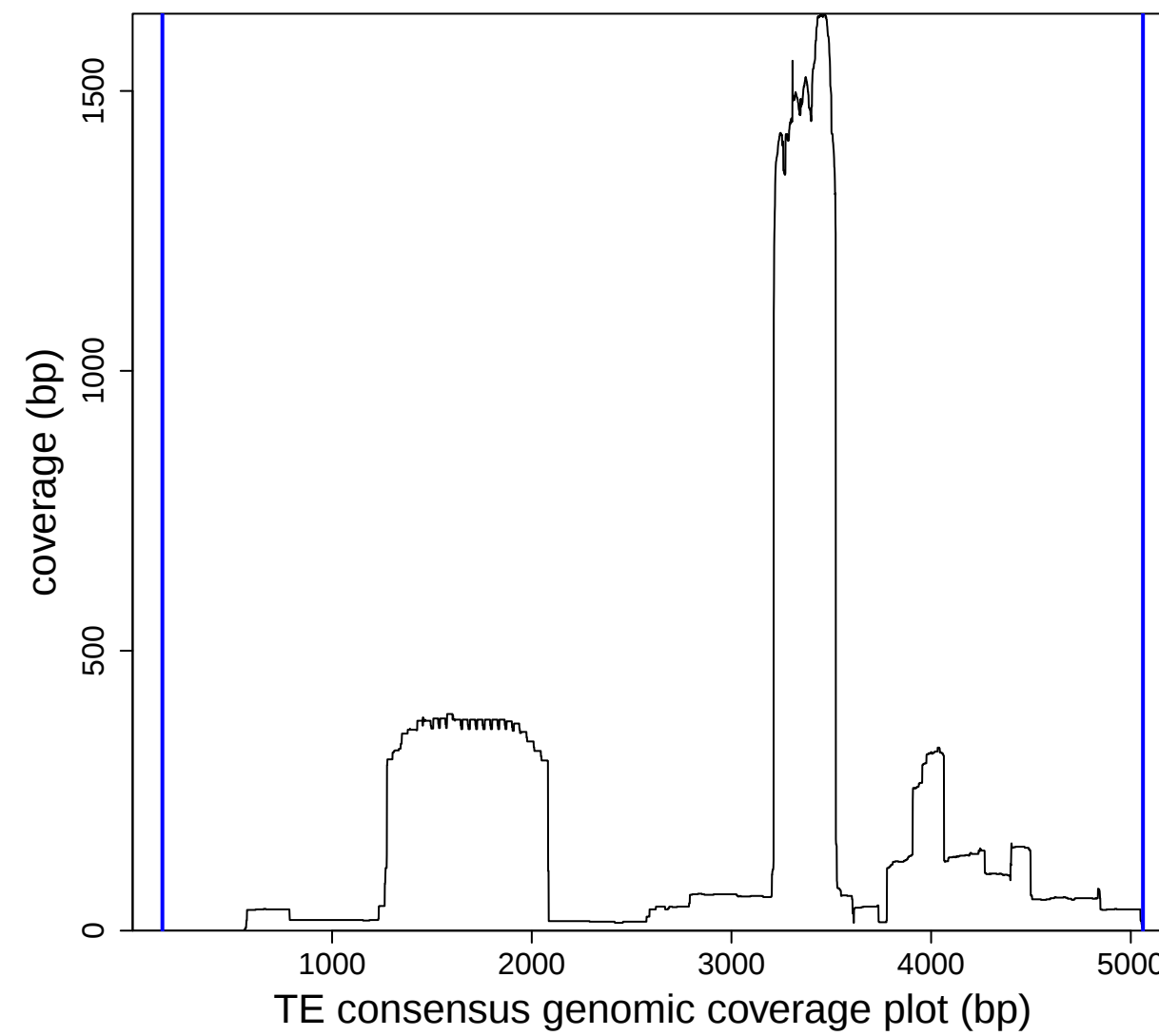
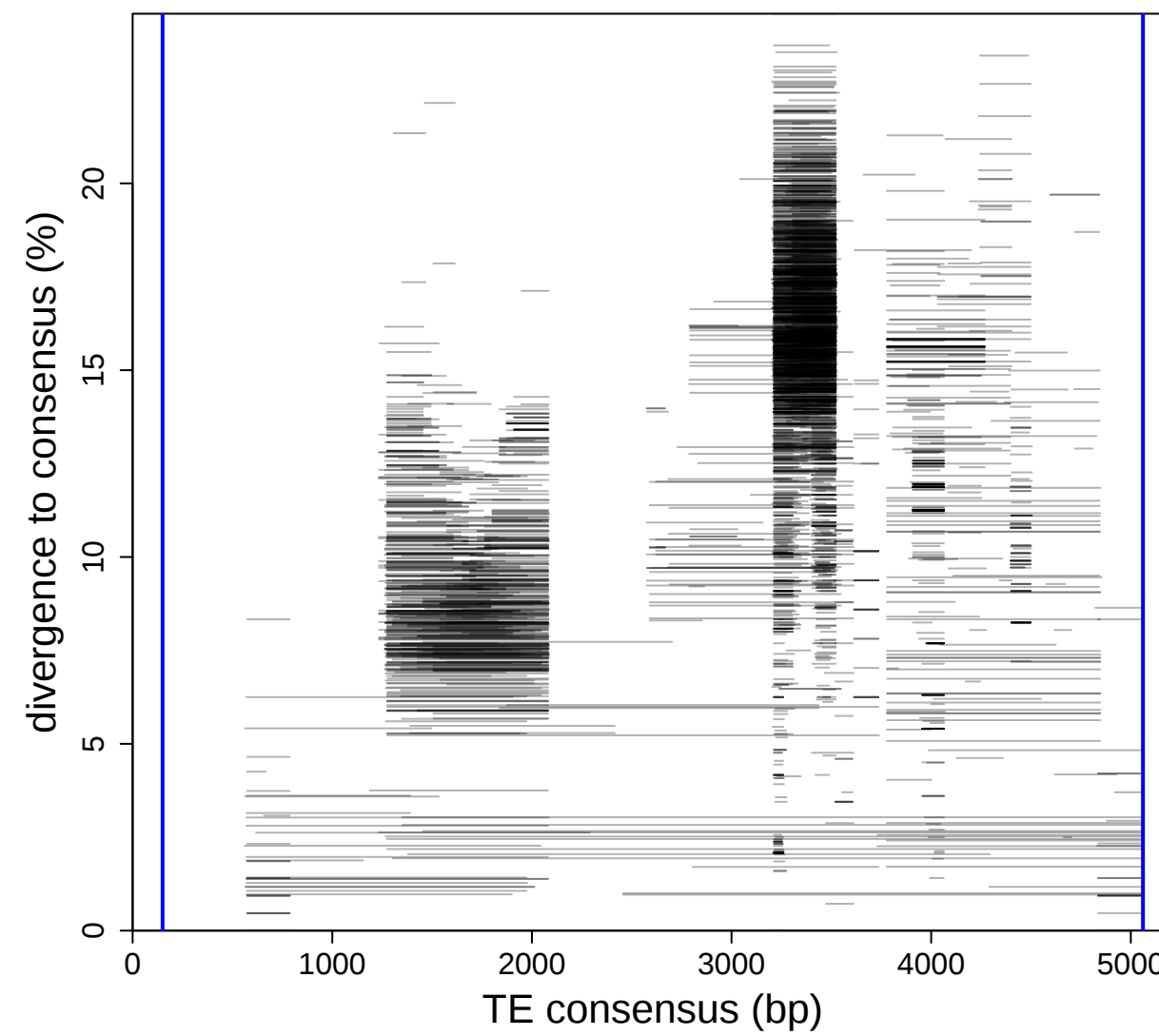
g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_ε
size: 4594bp; fragments: 3241; full length: 0 (>=4134.6bp)

After TETrimmer 4594 bp

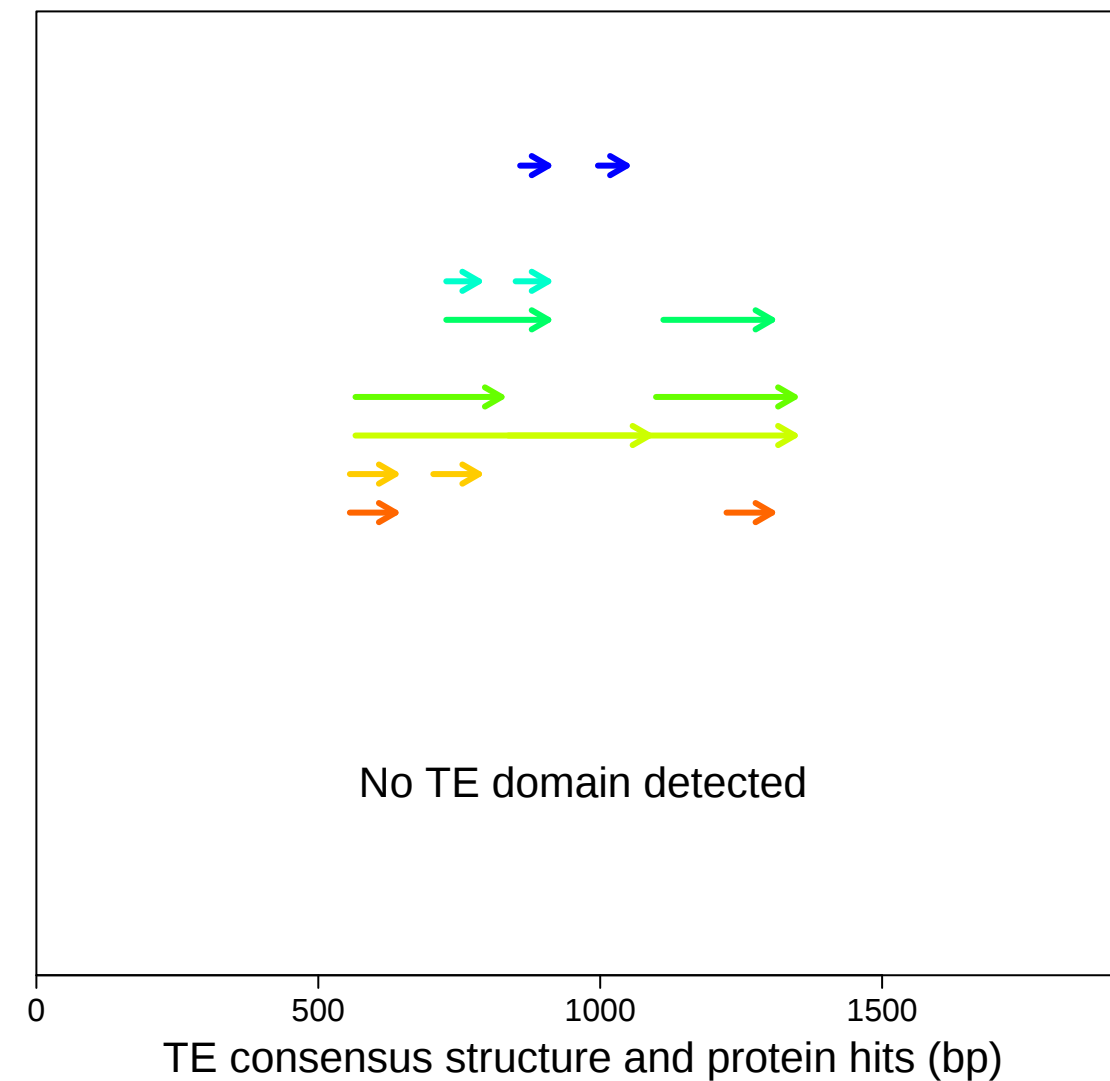
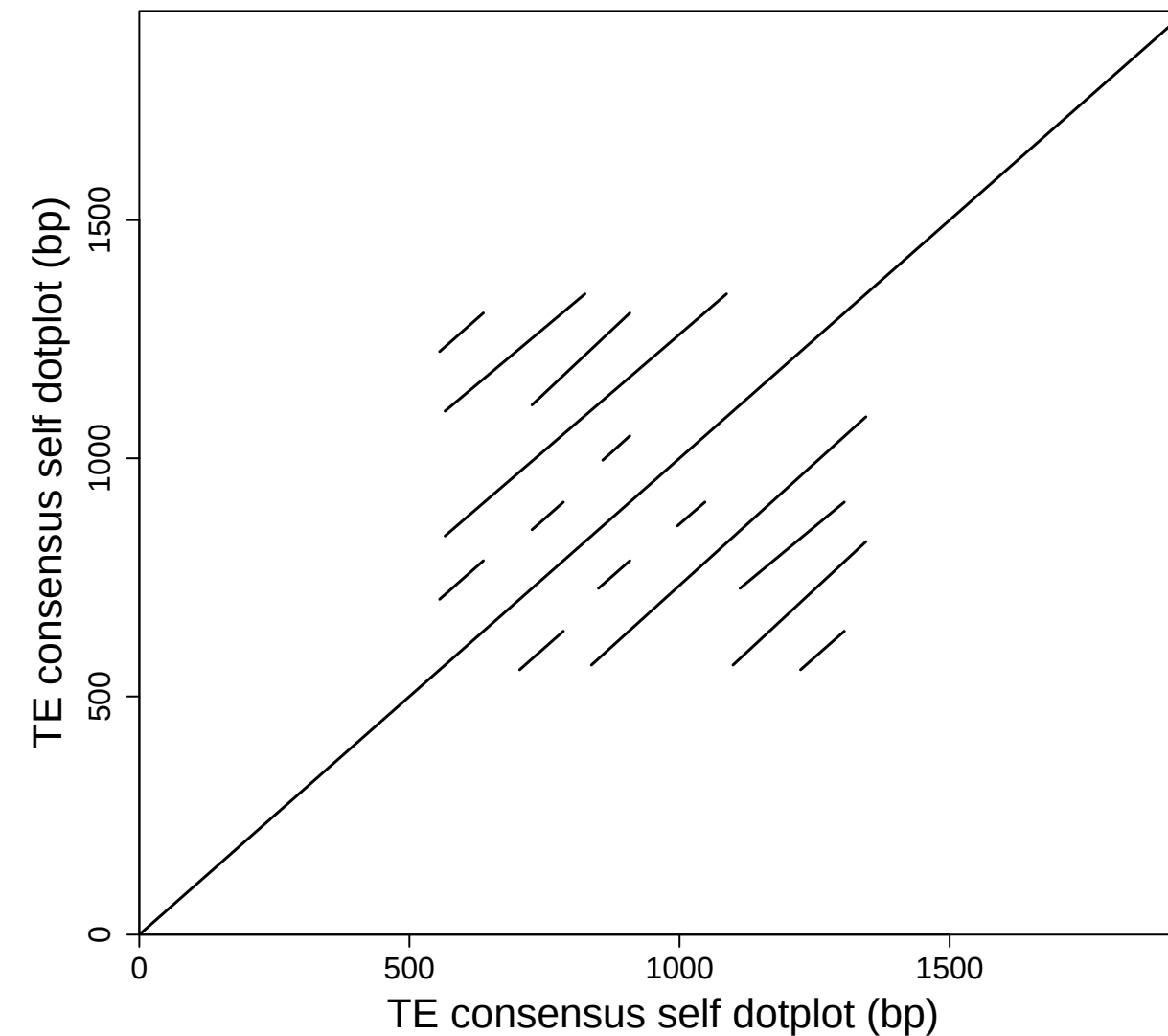
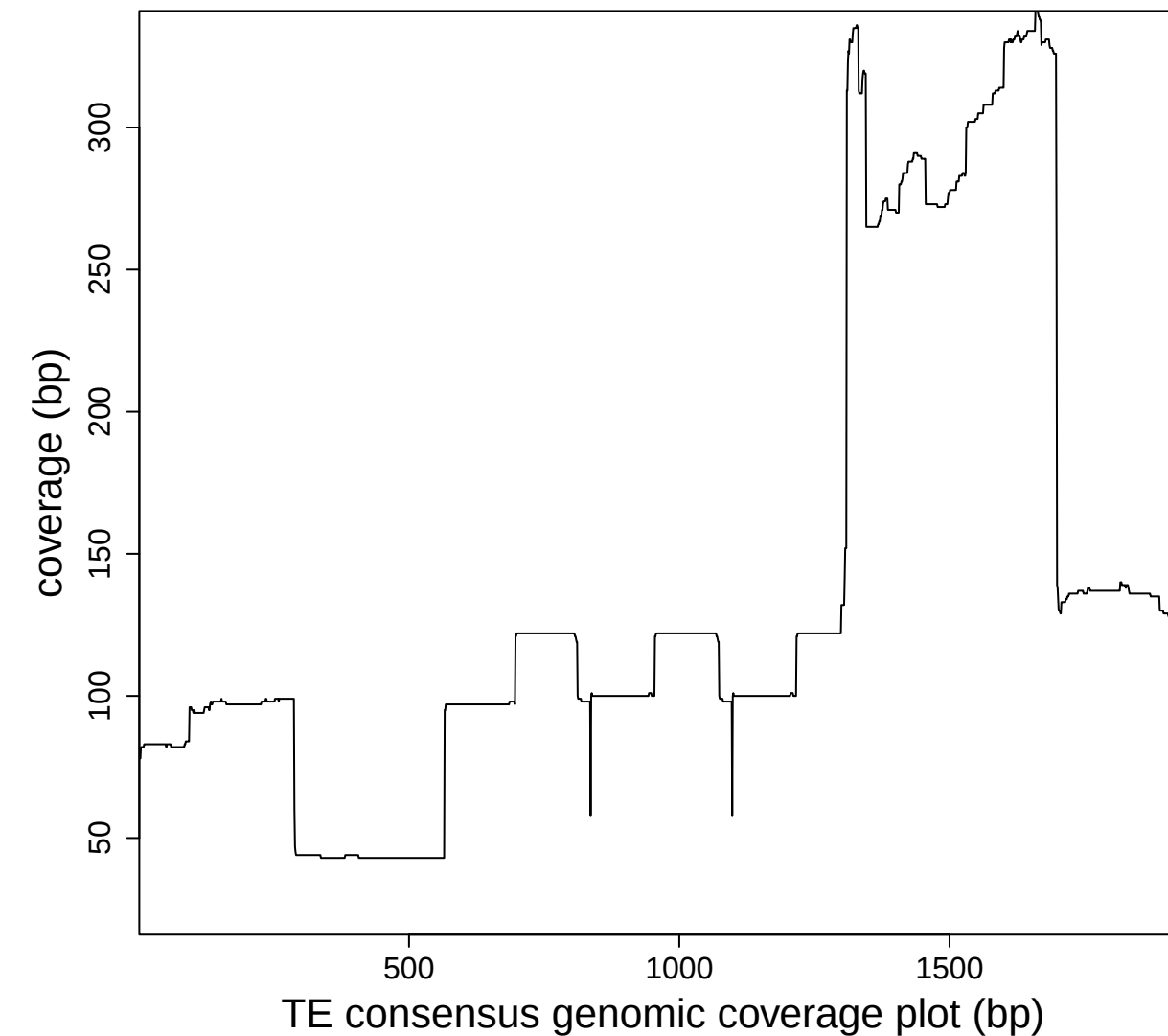
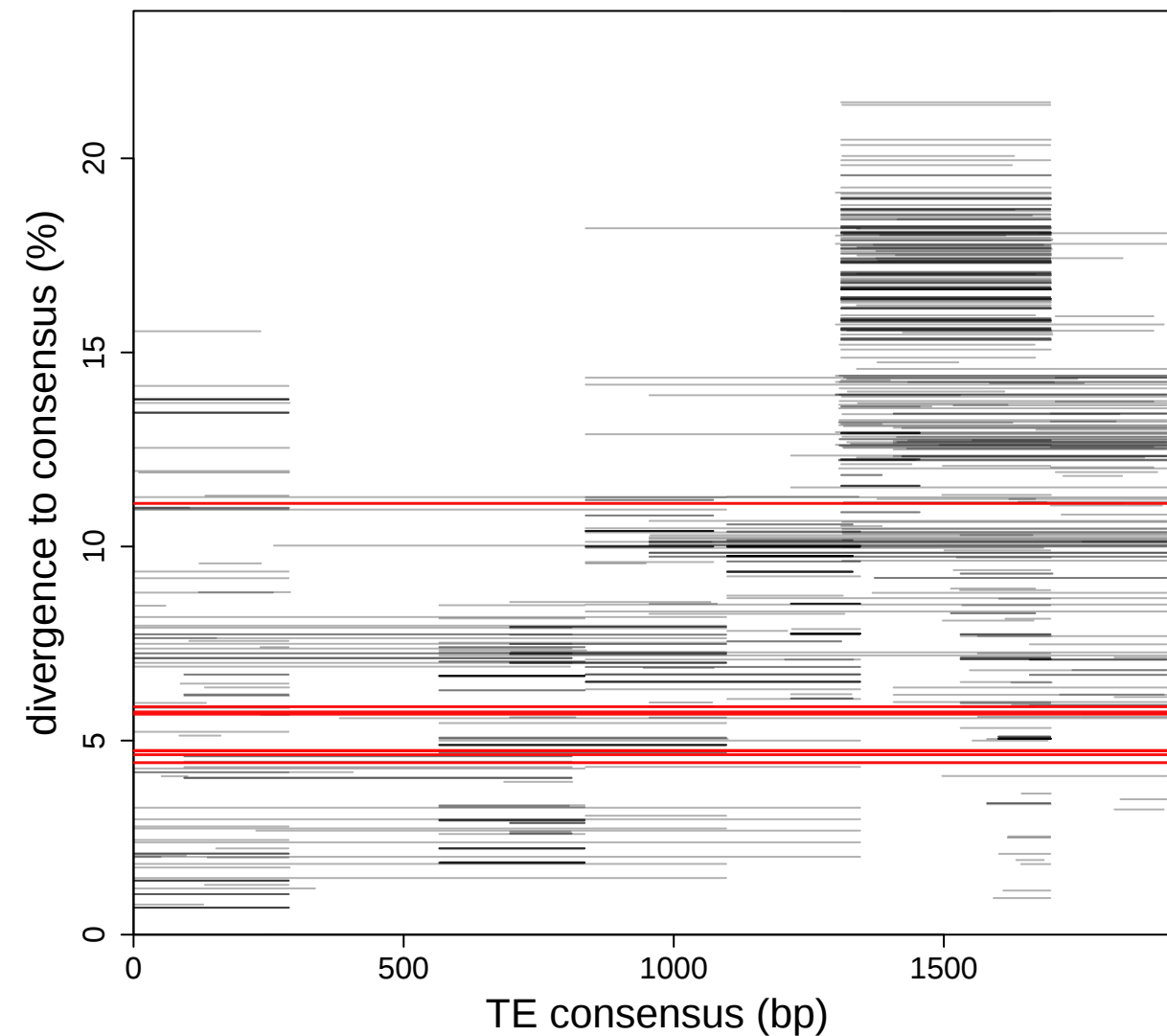


f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 5211bp; fragments: 3242; full length: 0 (>=4689.9bp)

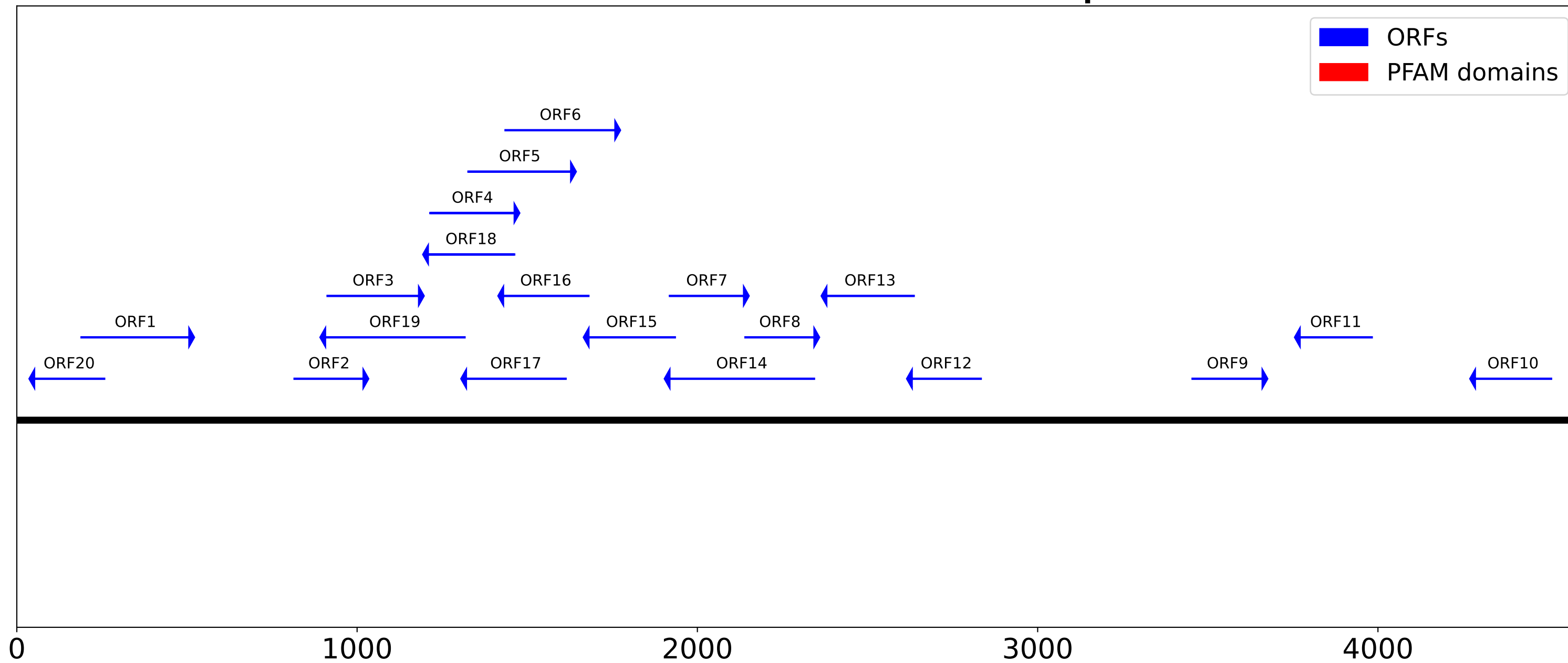
After TEtrimmer Extended plot Blue lines are boundaries



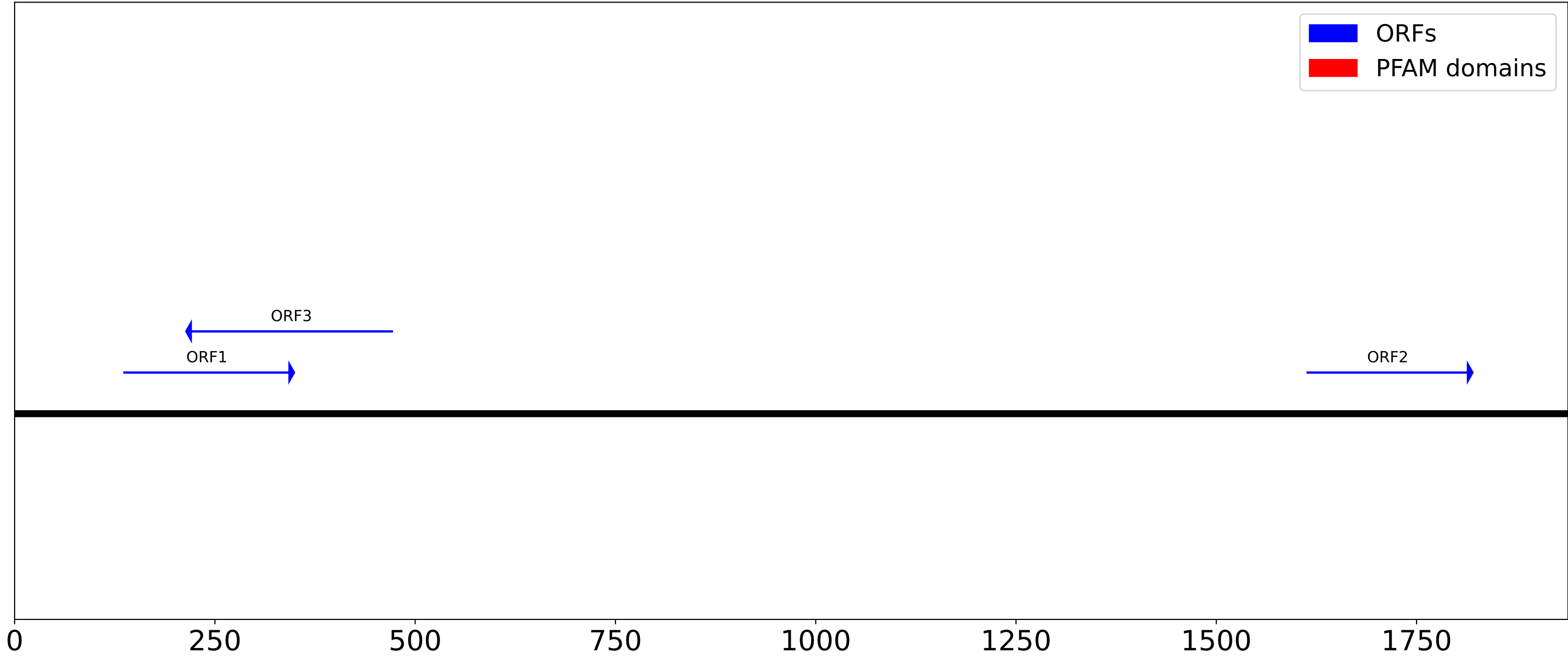
TE: rnd_4_family_1188
size: 1939bp; fragments: 680; full length: 8 (≥ 1745.1 bp)



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)

