

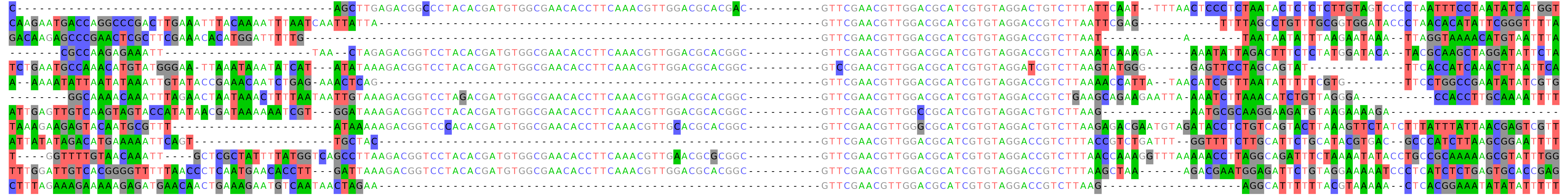
Start crop Point



End crop Point

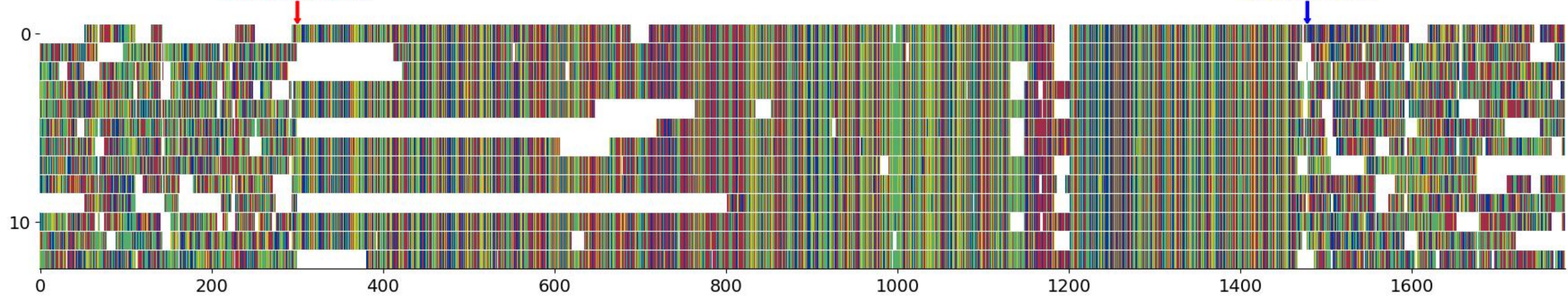


MSA length = 1179



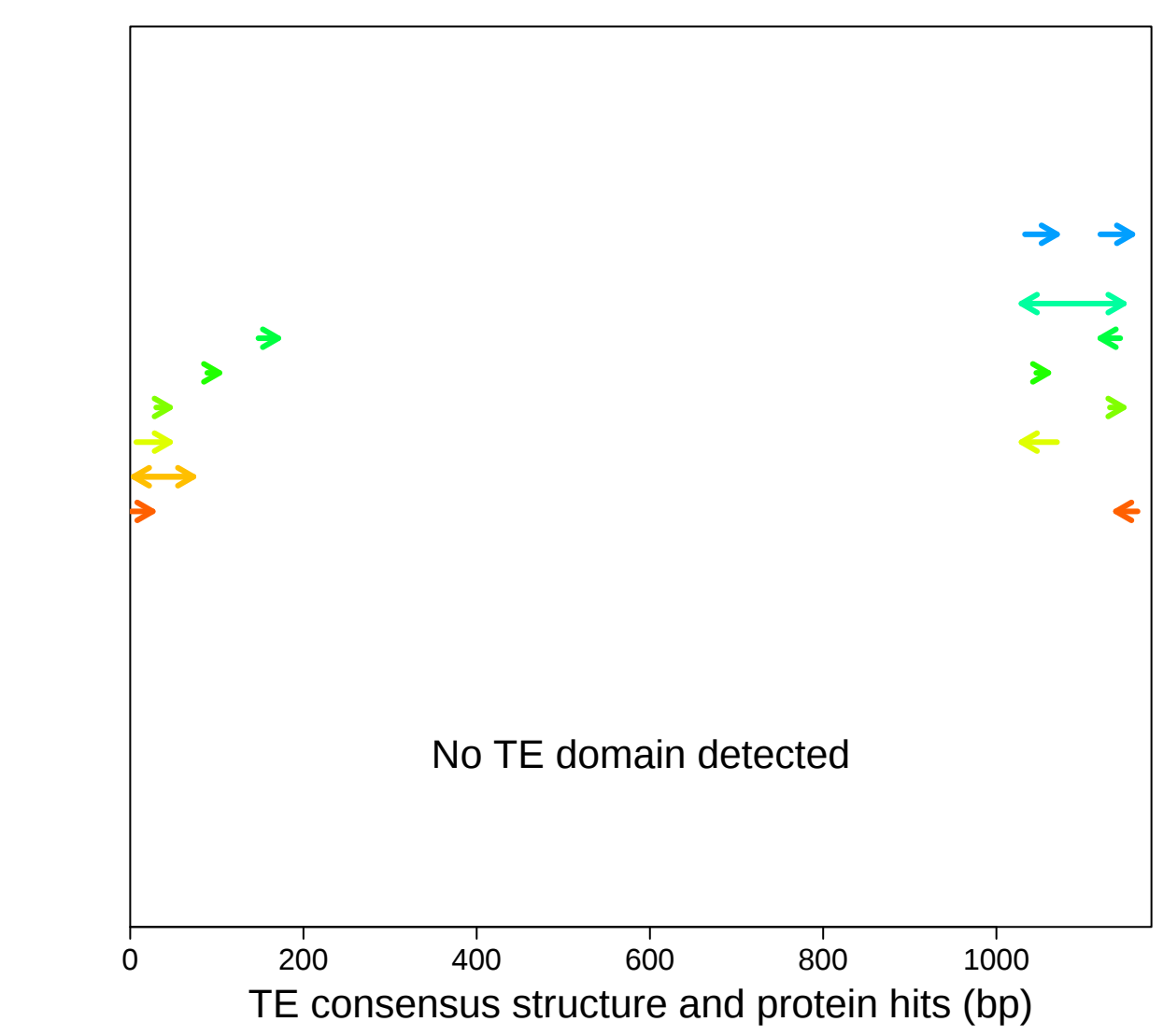
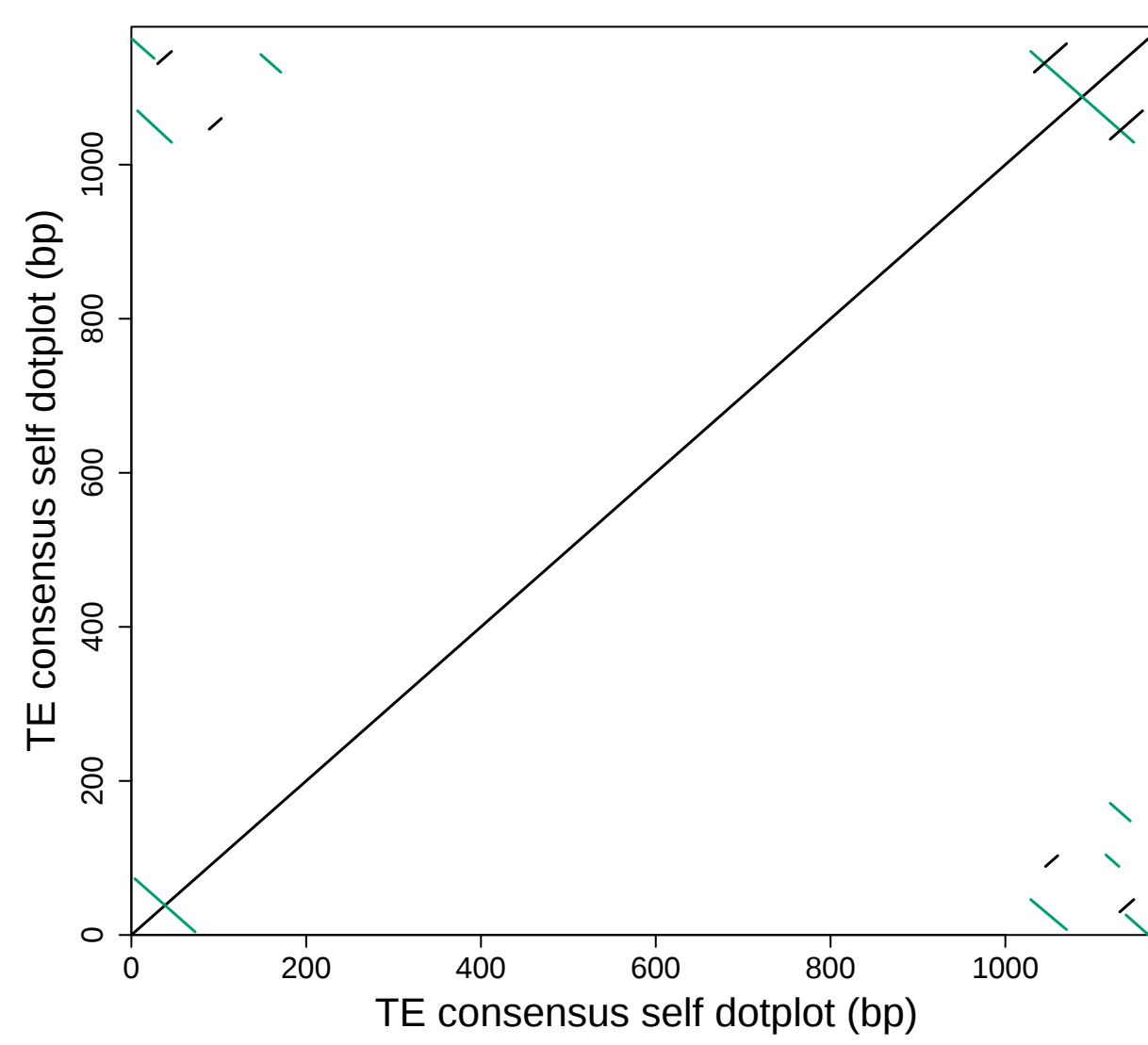
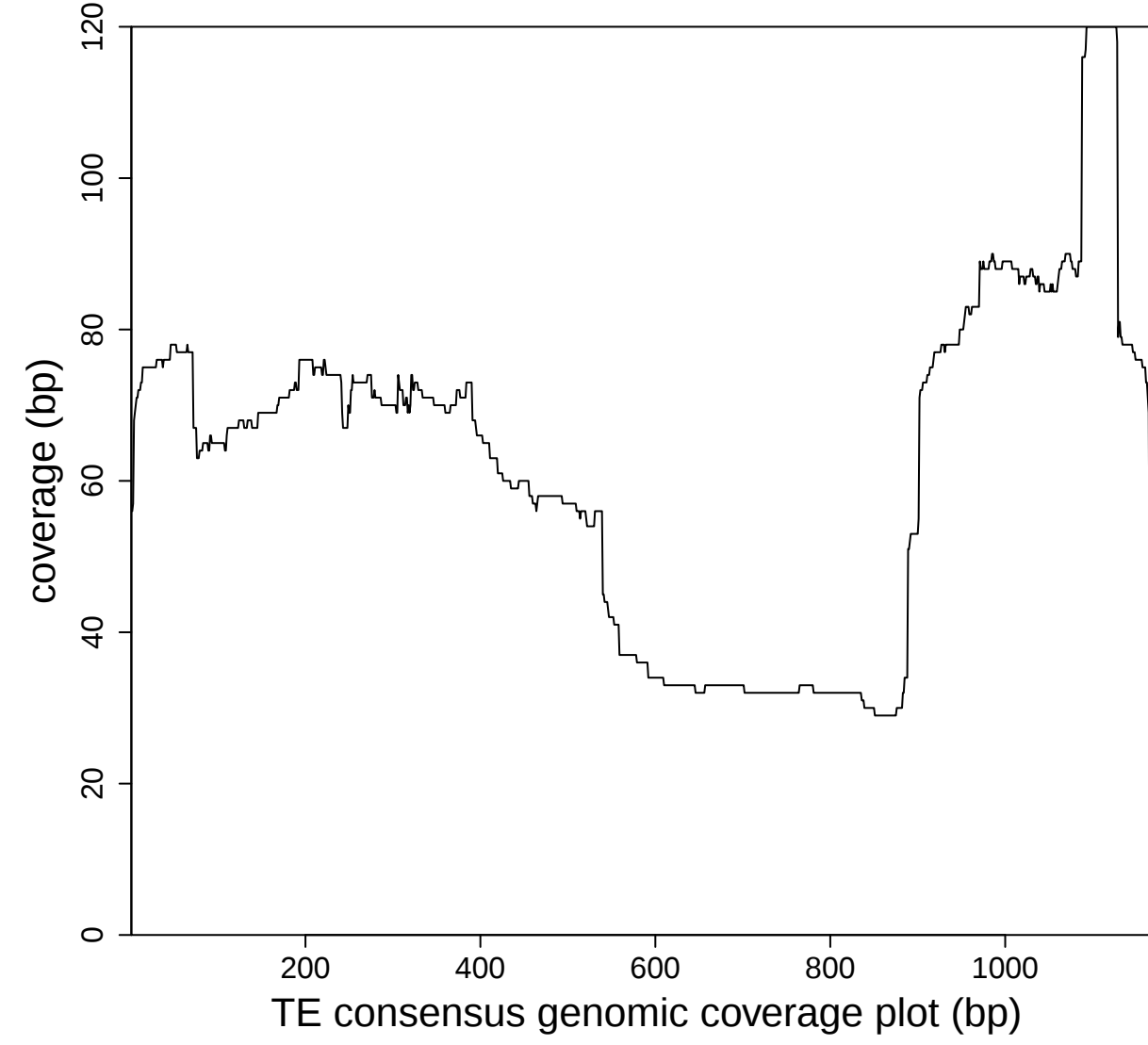
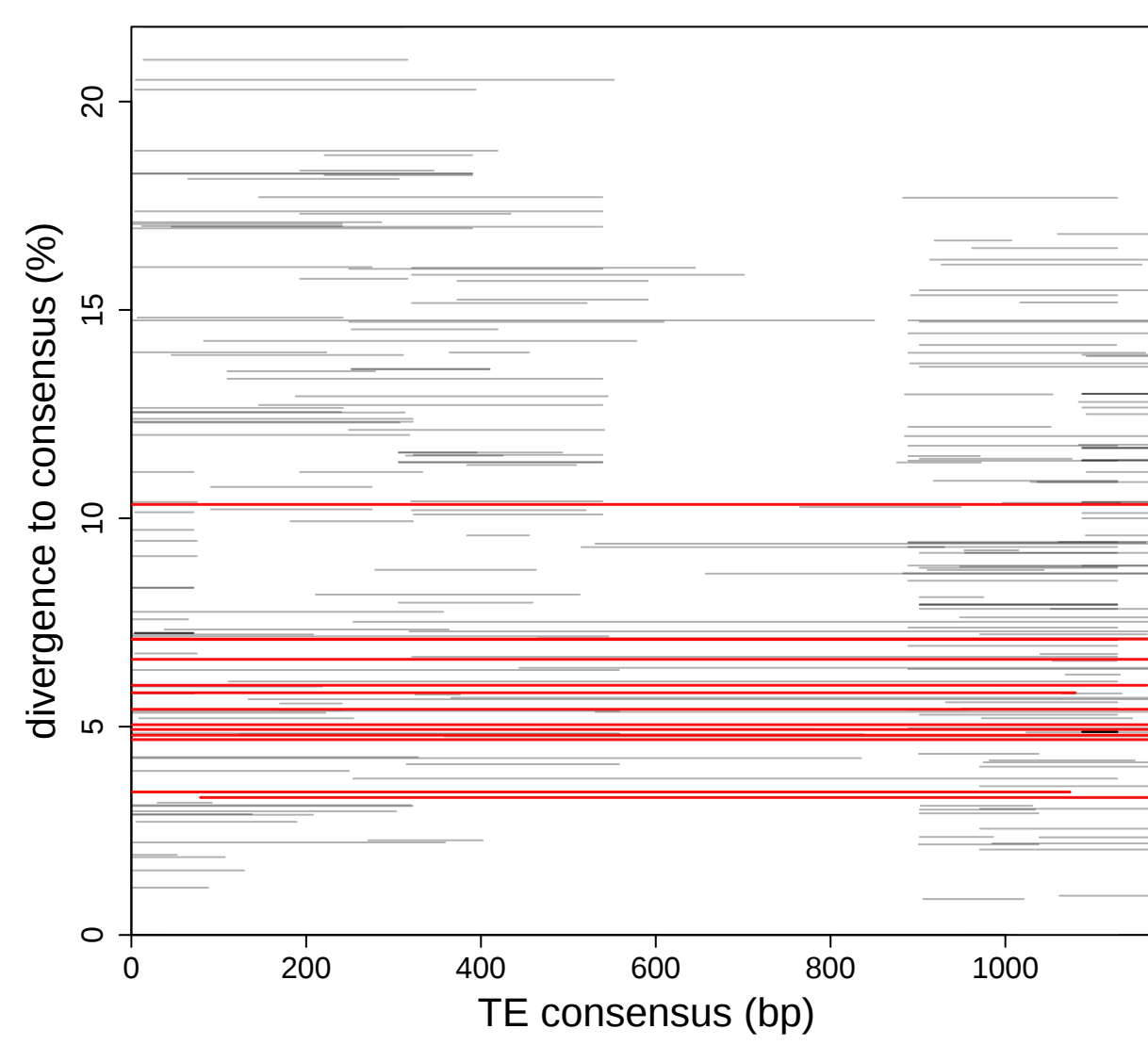
Start crop Point

End crop Point



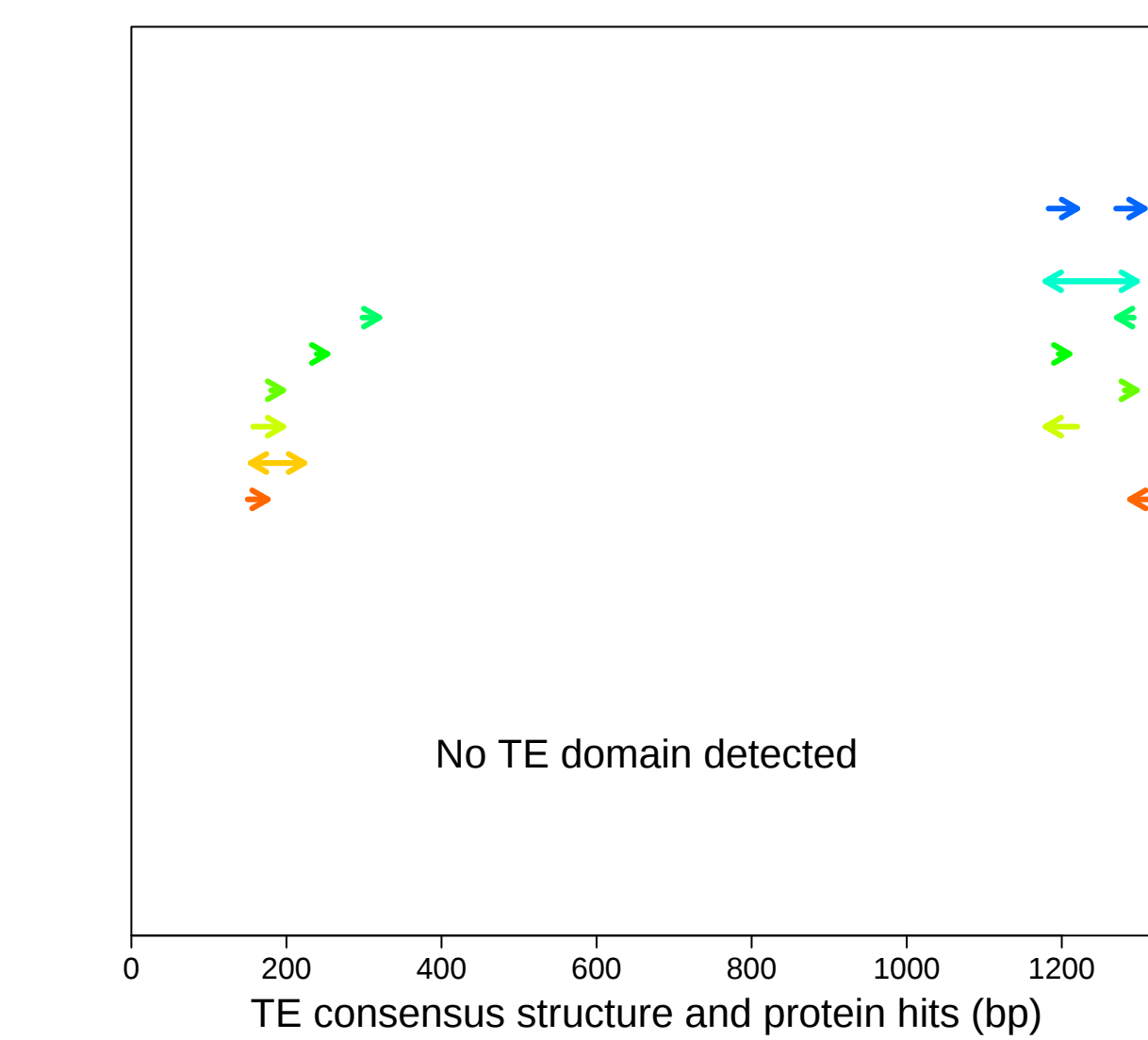
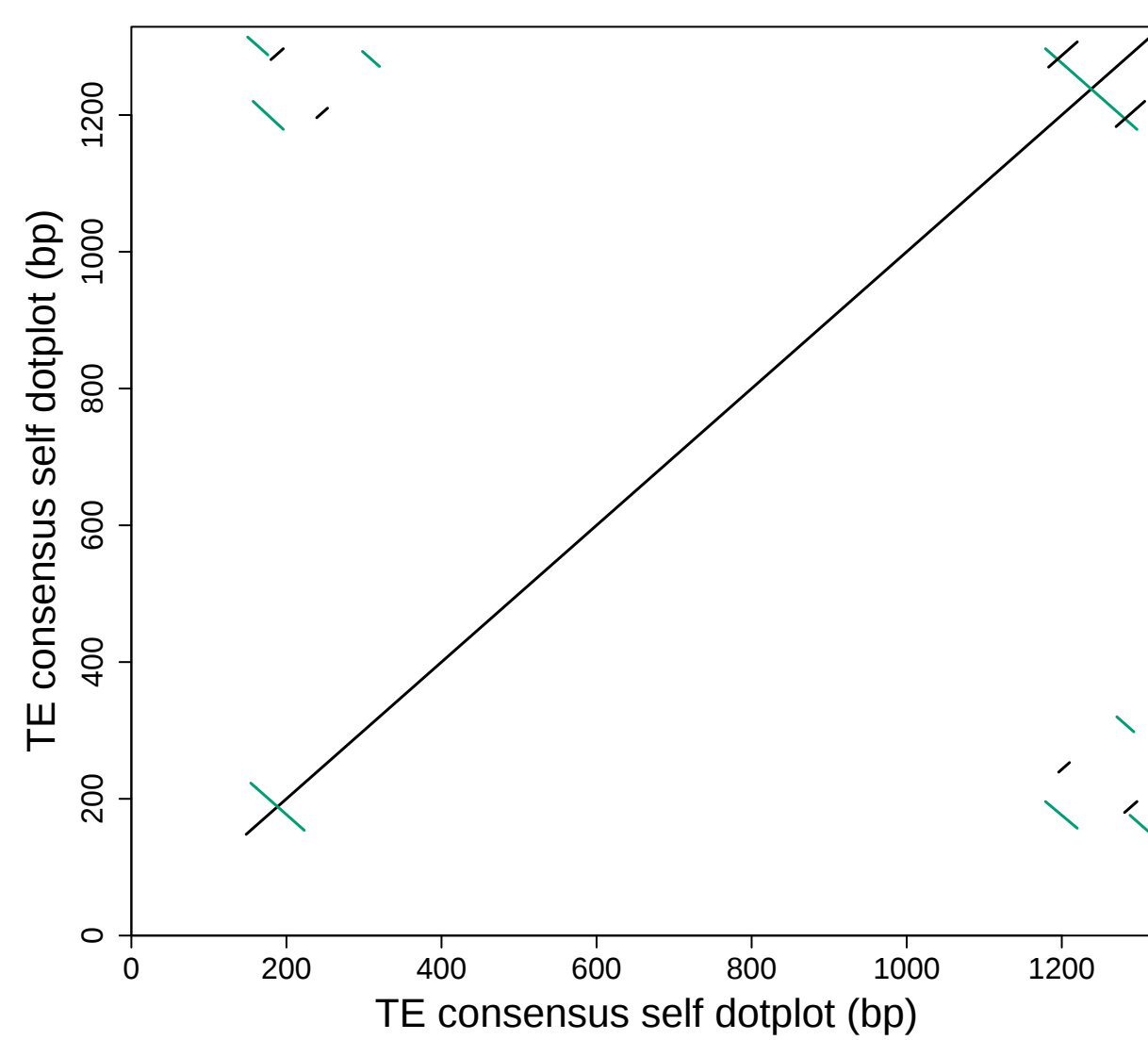
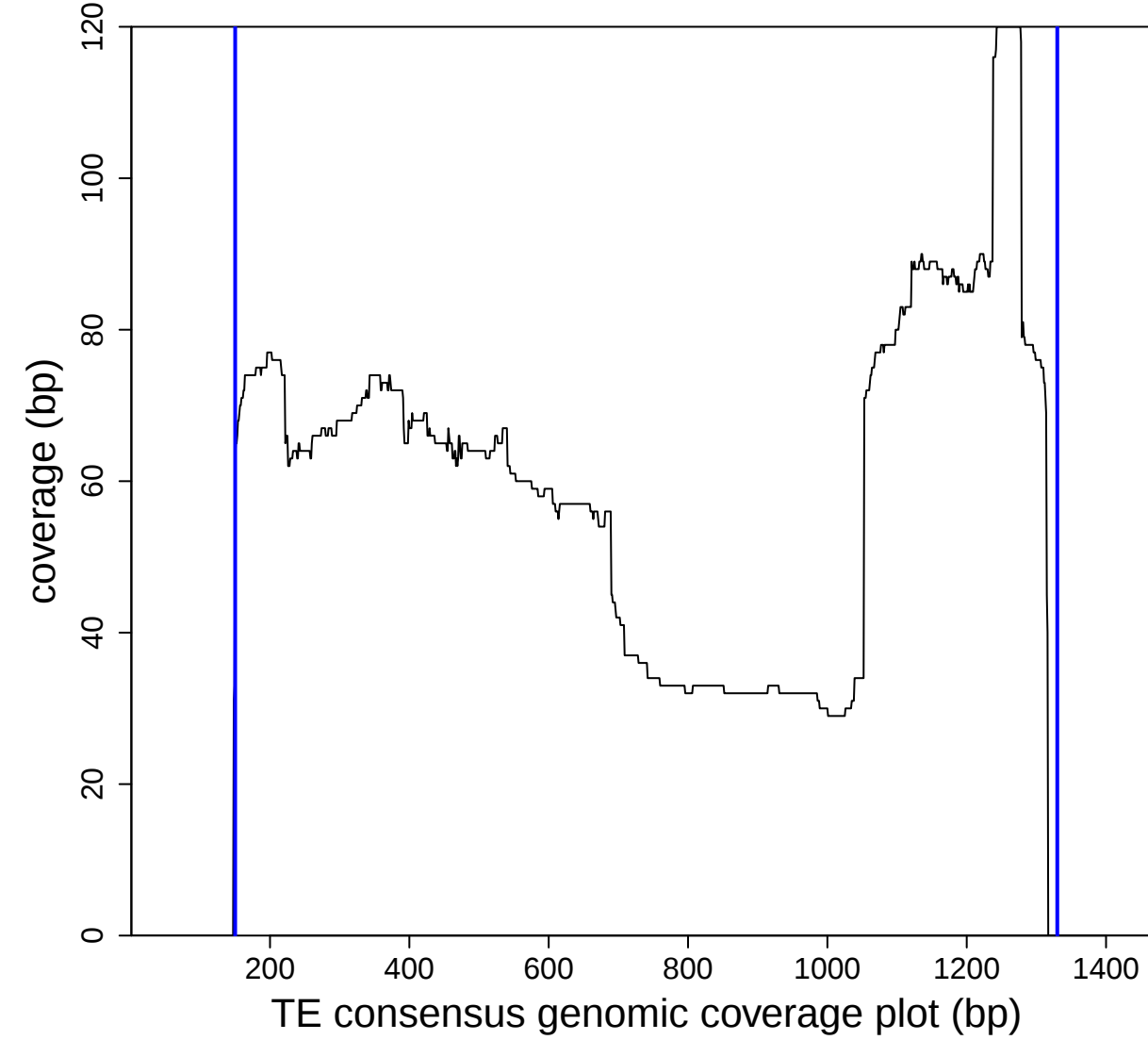
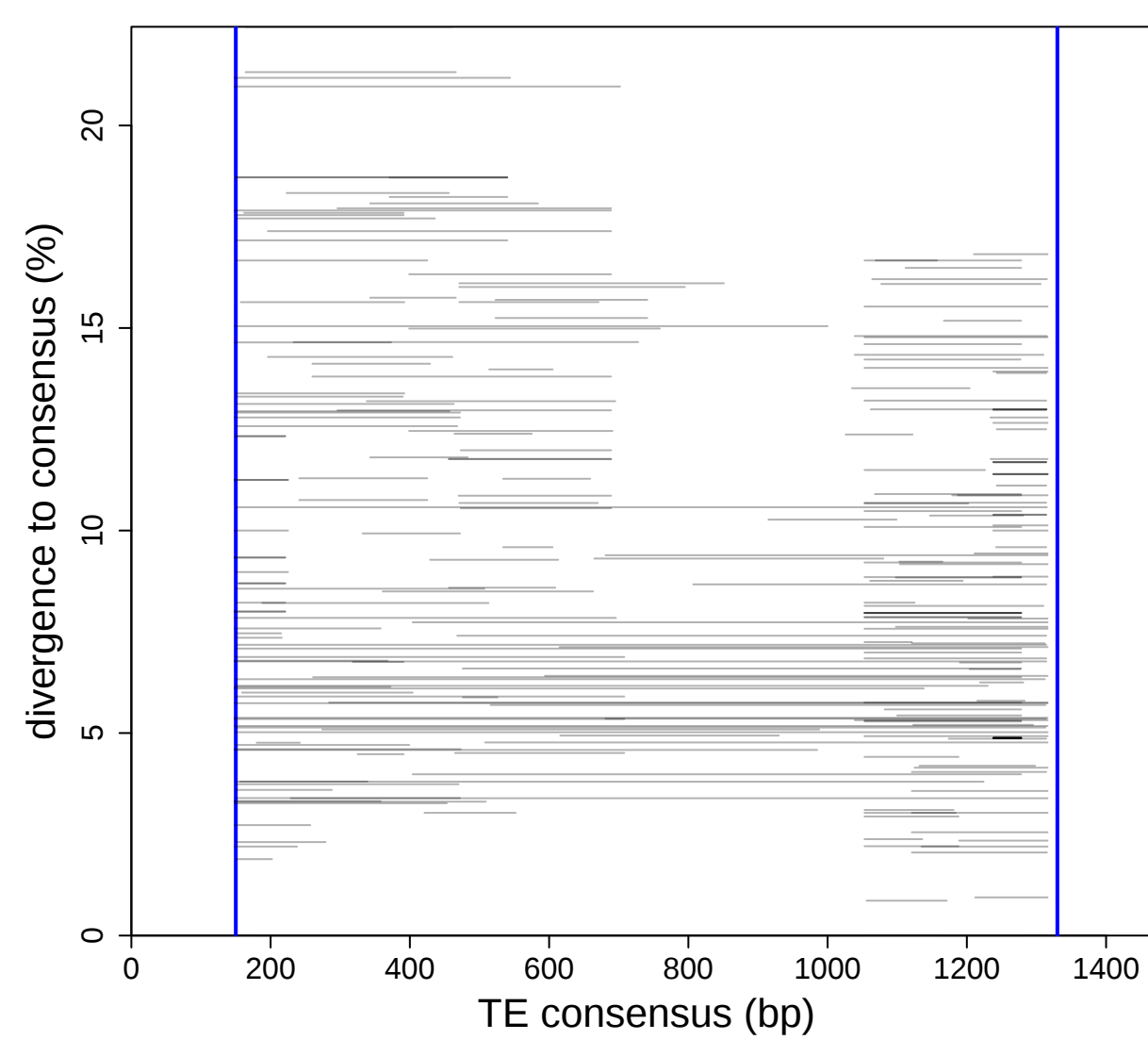
g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_3
size: 1179bp; fragments: 260; full length: 13 (>=1061.1bp)

After TETrimmer 1179 bp

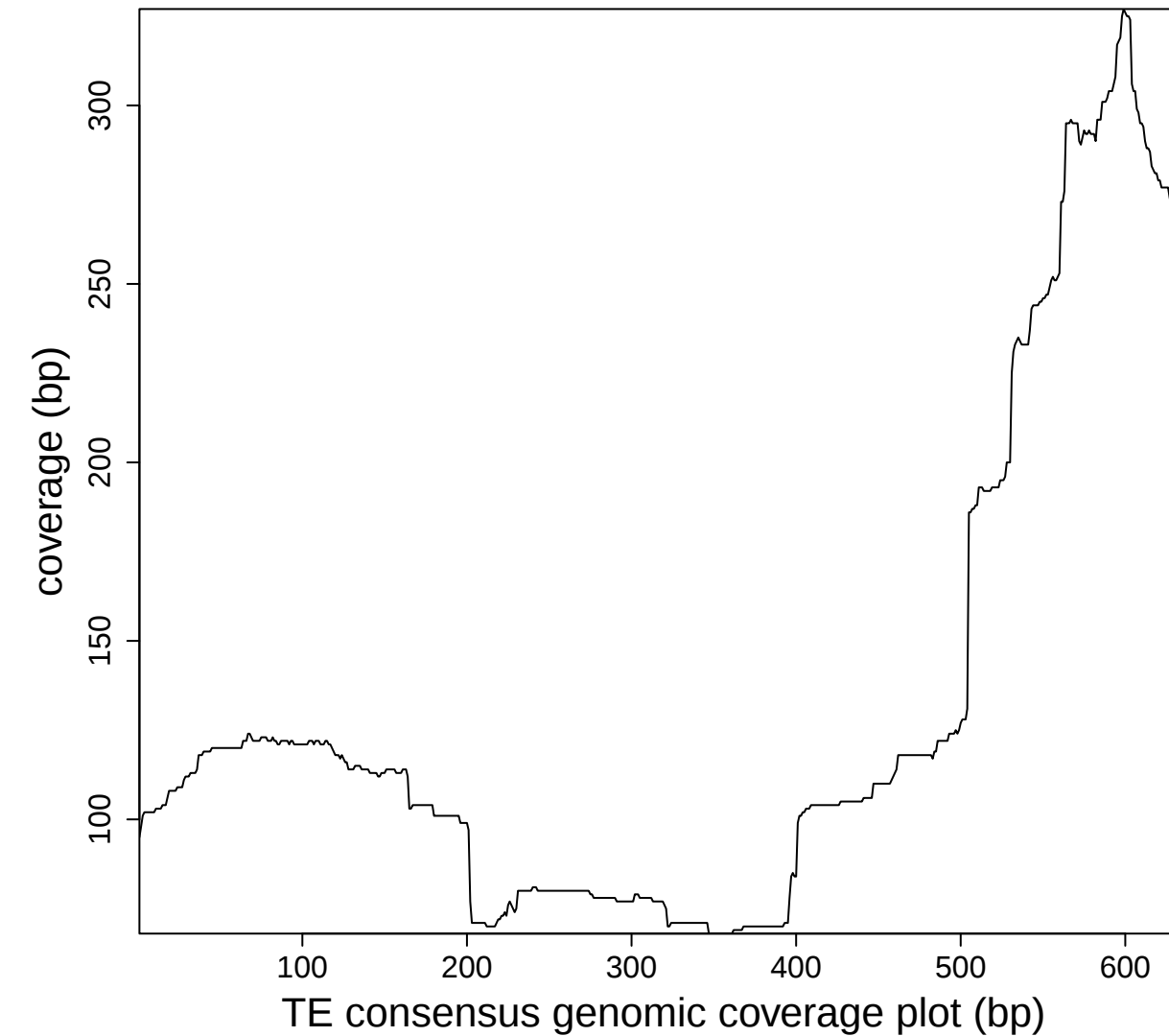
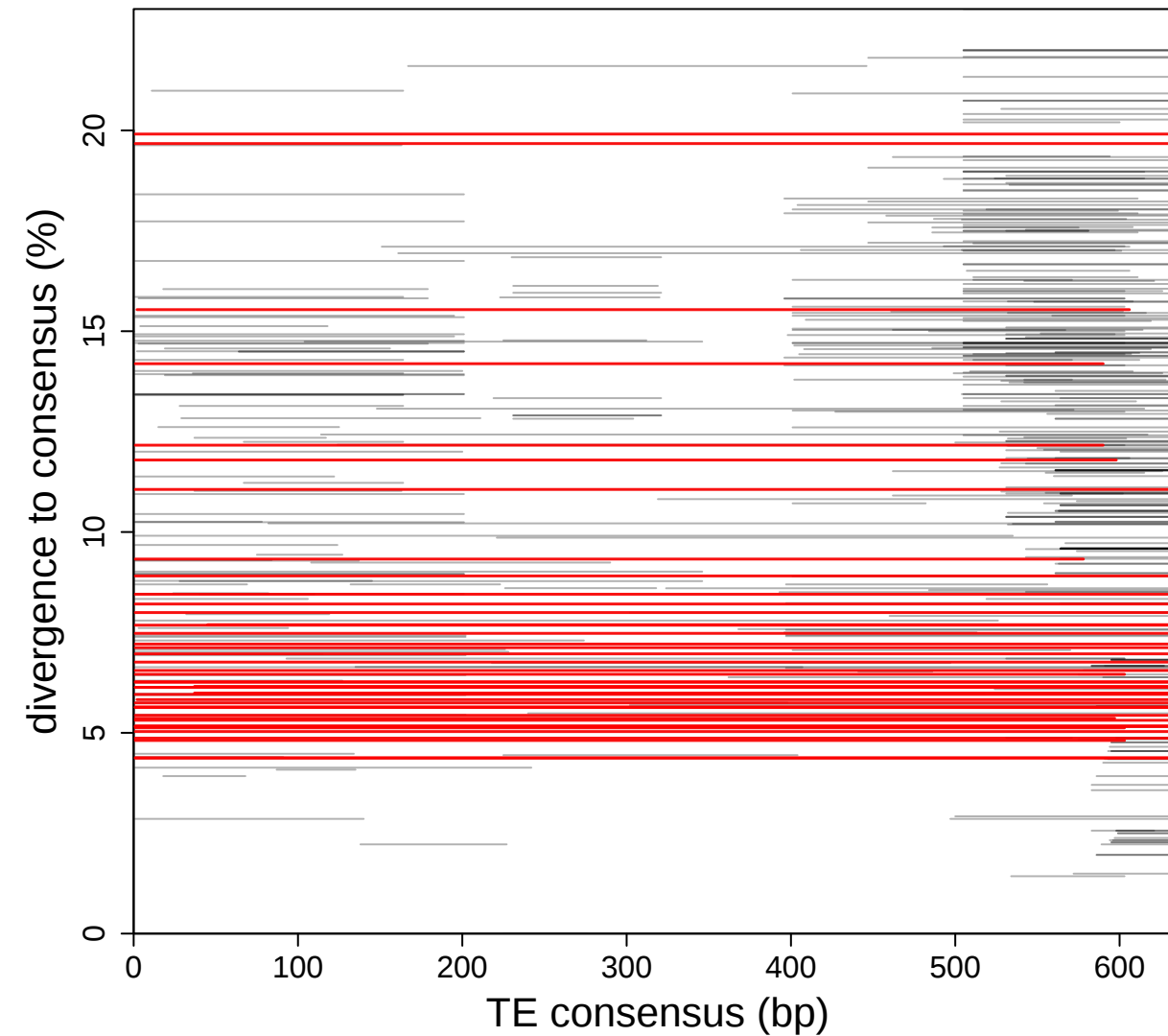


f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 1480bp; fragments: 253; full length: 0 (>=1332bp)

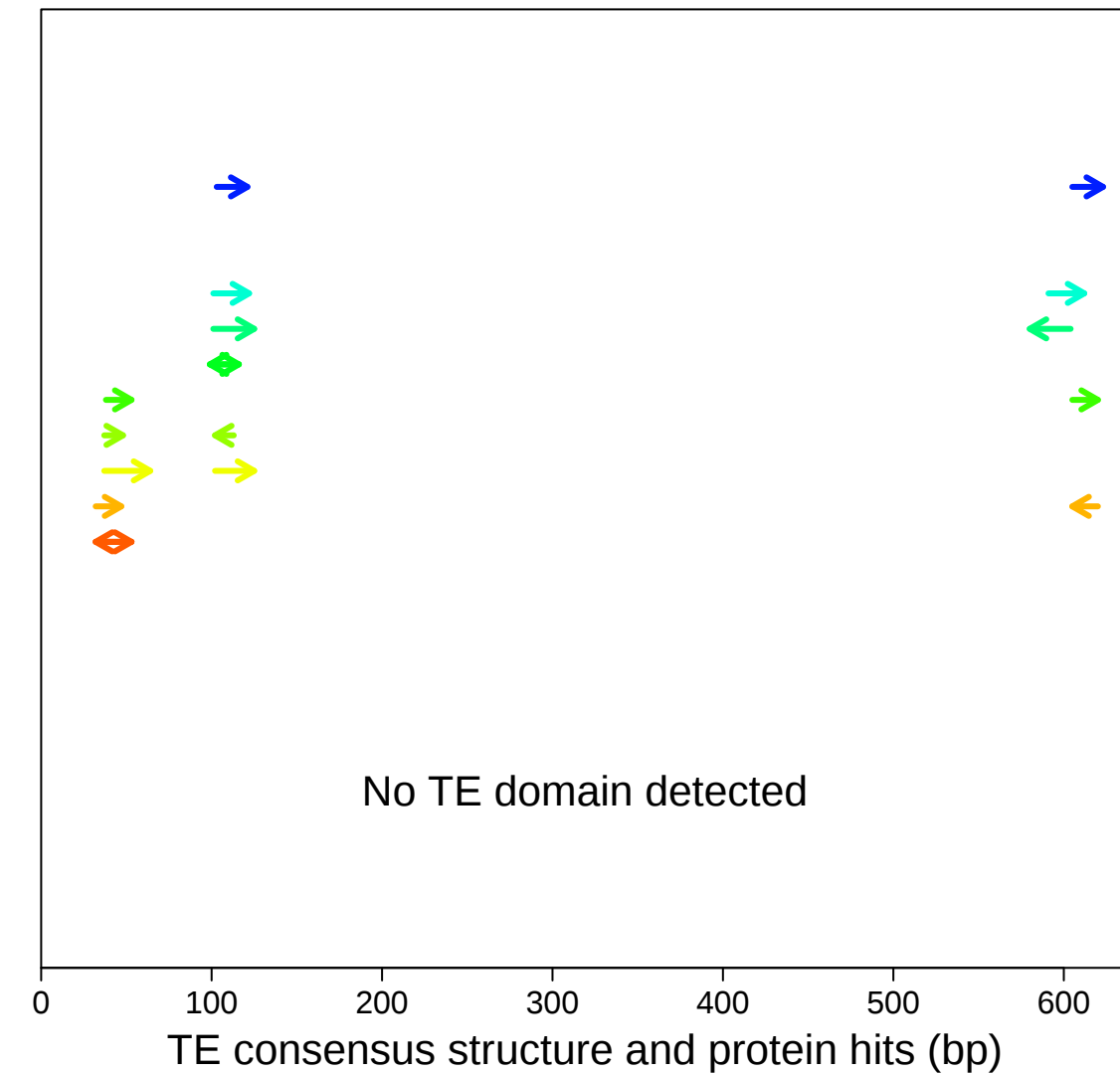
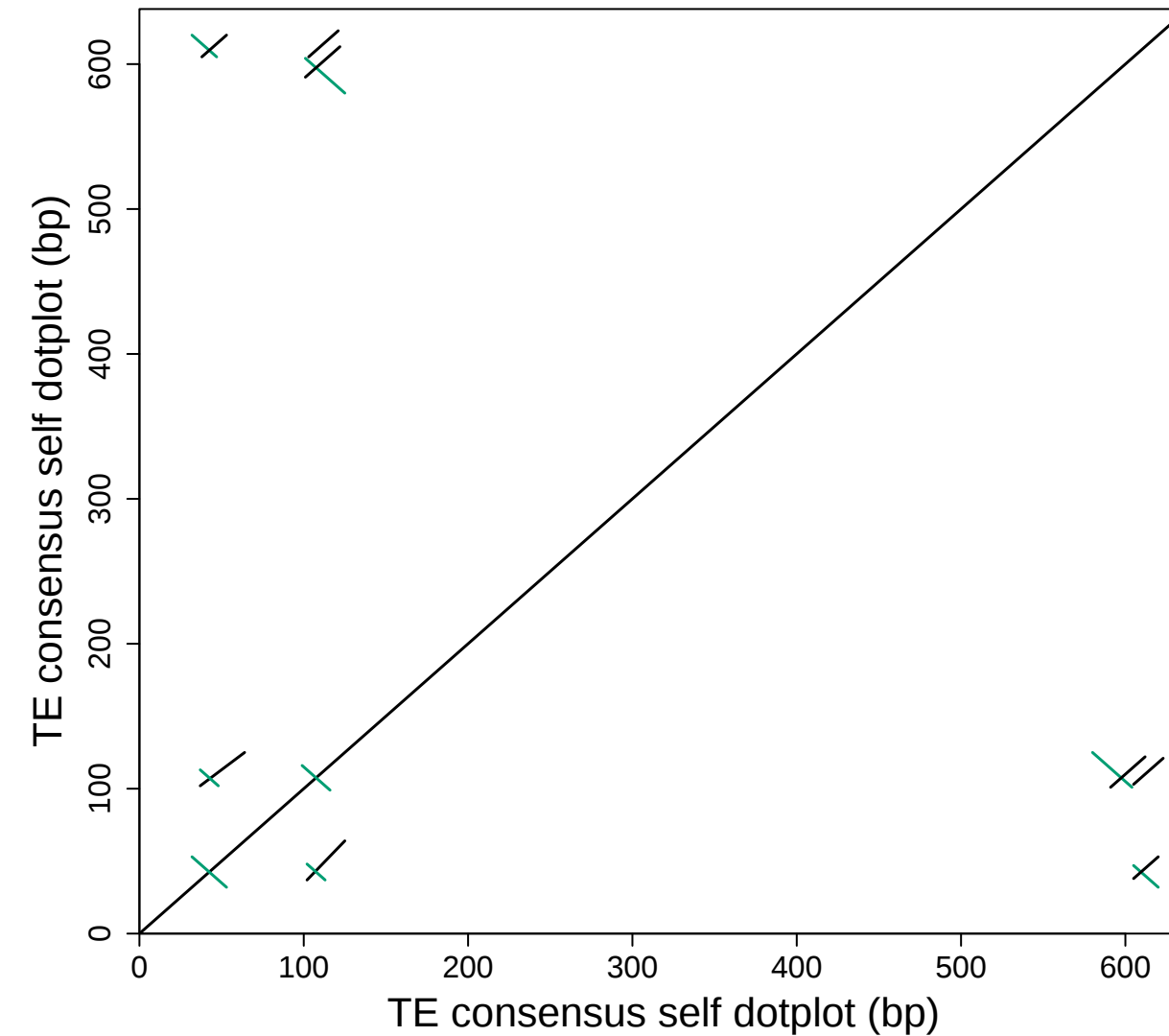
After TEtrimmer Extended plot Blue lines are boundaries



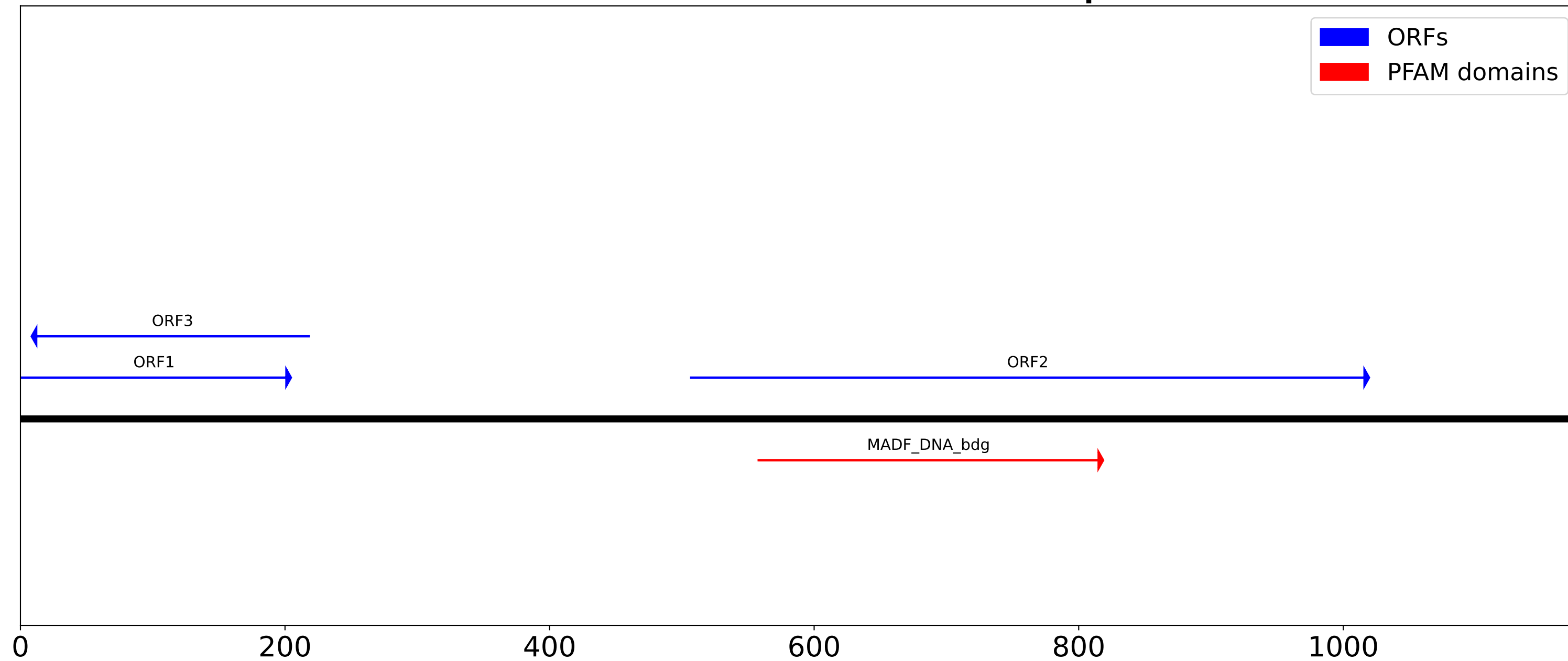
TE: rnd_5_family_2161
size: 638bp; fragments: 454; full length: 45 (≥ 574.2 bp)



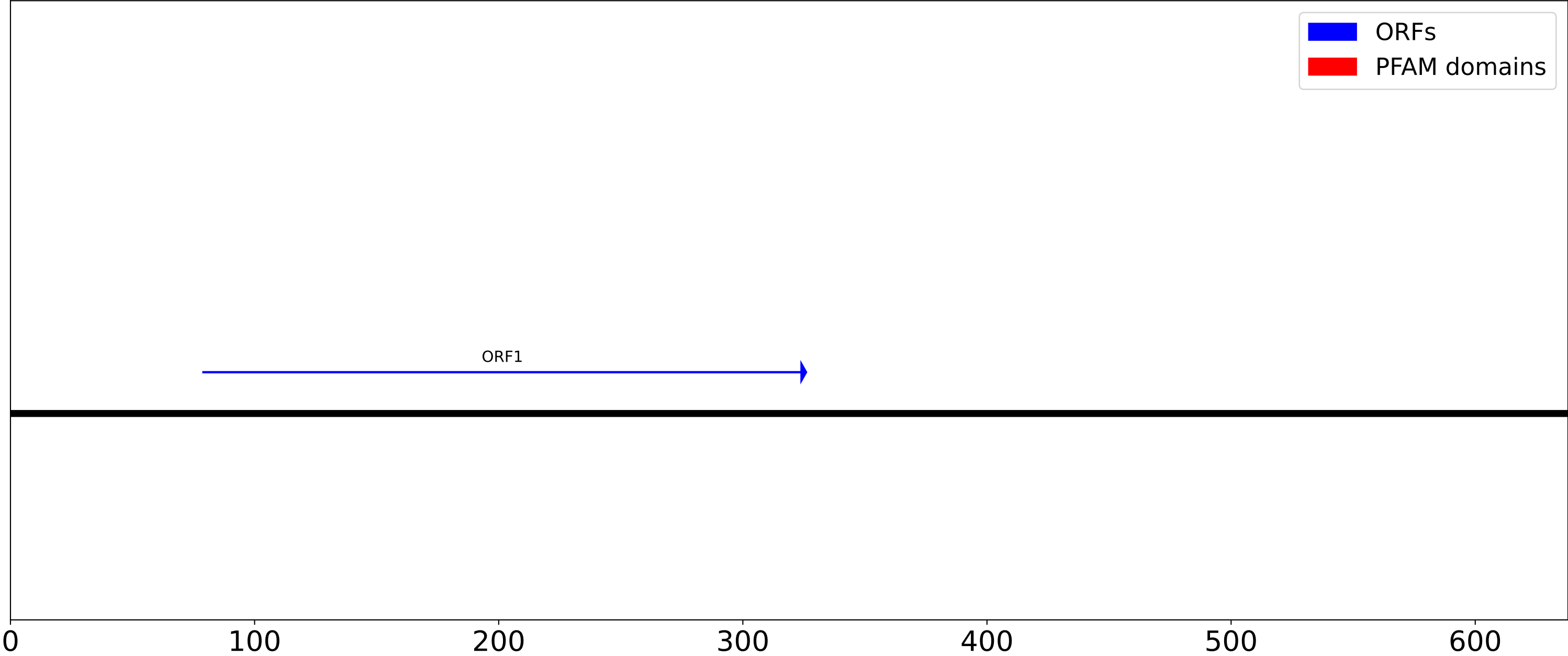
Before TEtrimmer 638 bp



After TEtrimmer ORF and PFAM domain plot



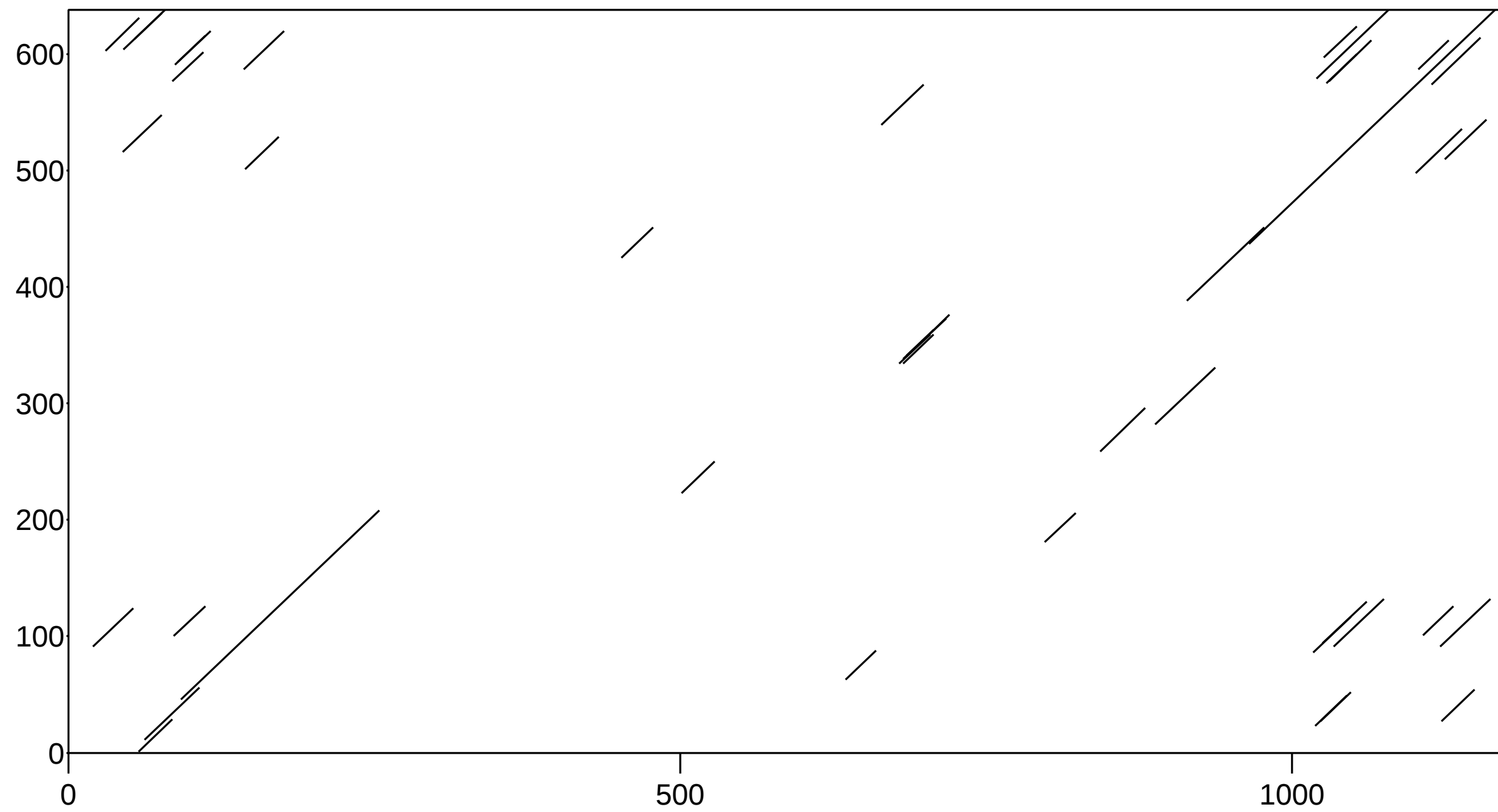
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)