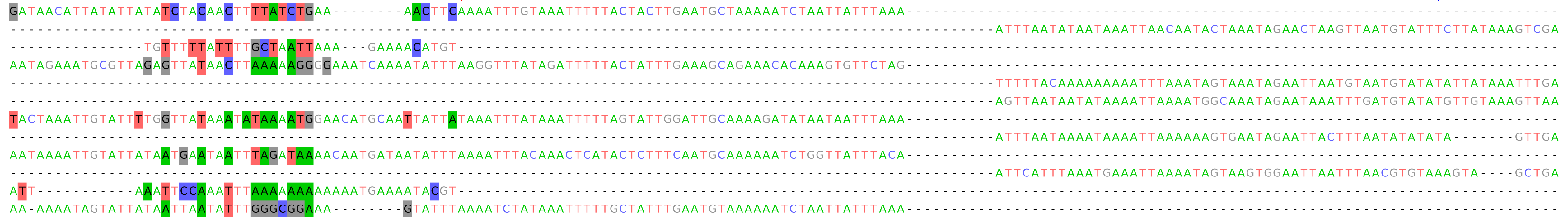
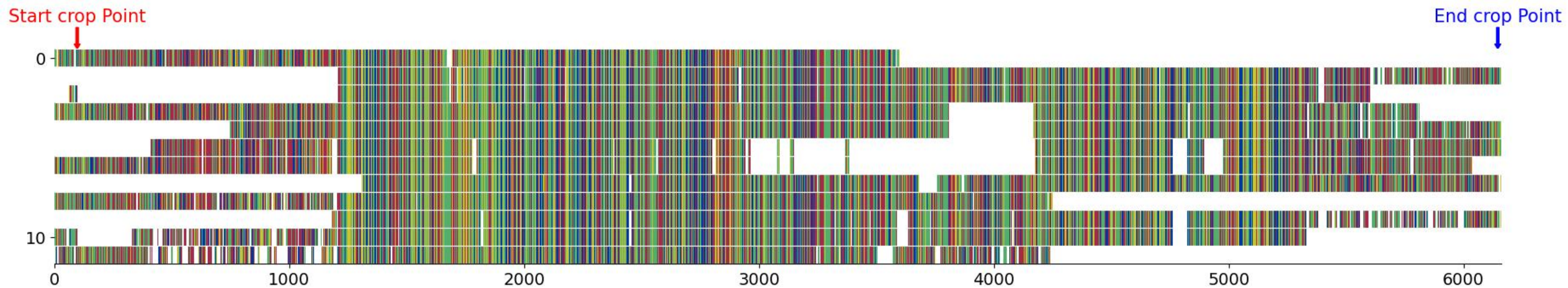


Start crop Point

End crop Point

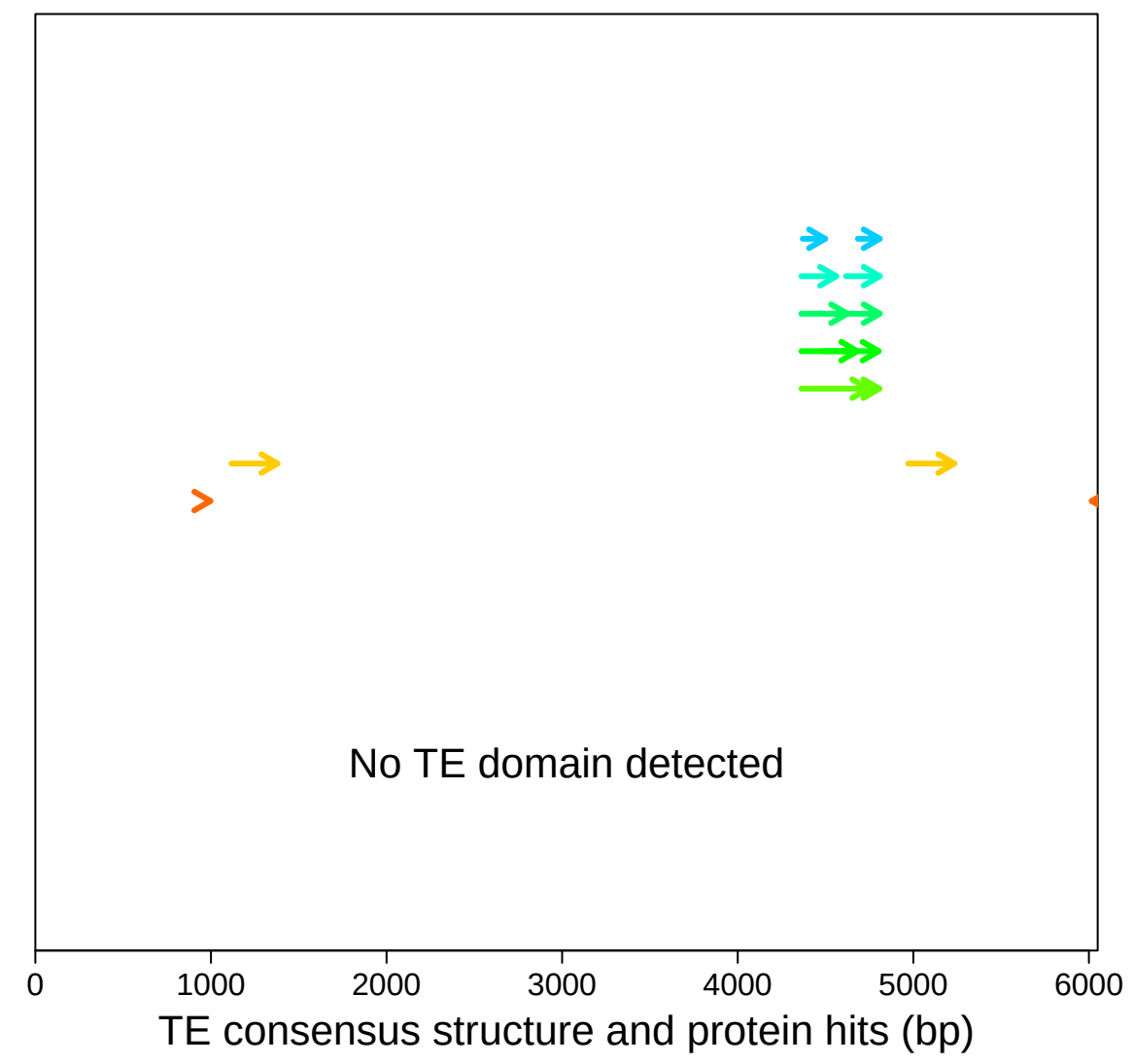
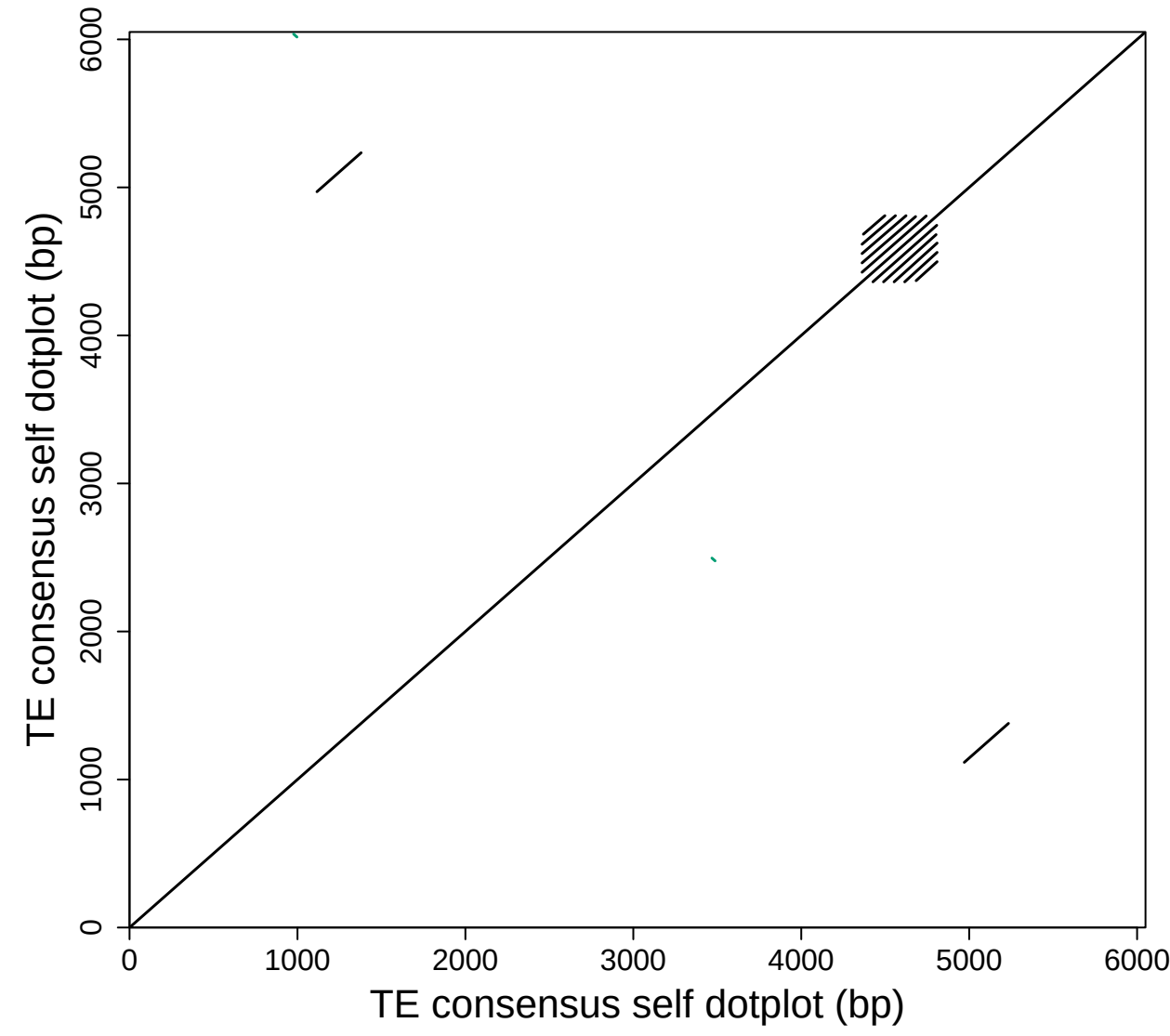
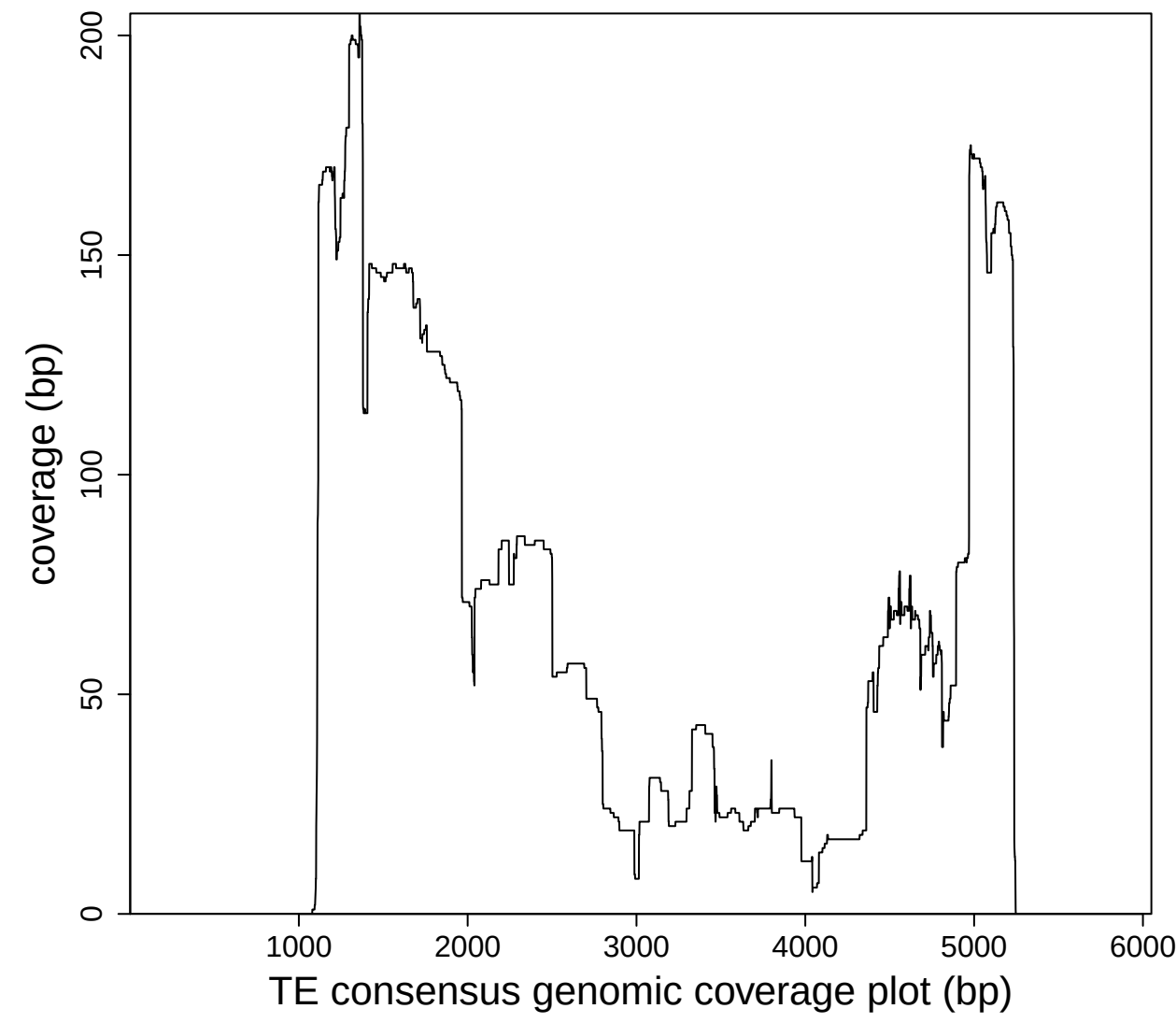
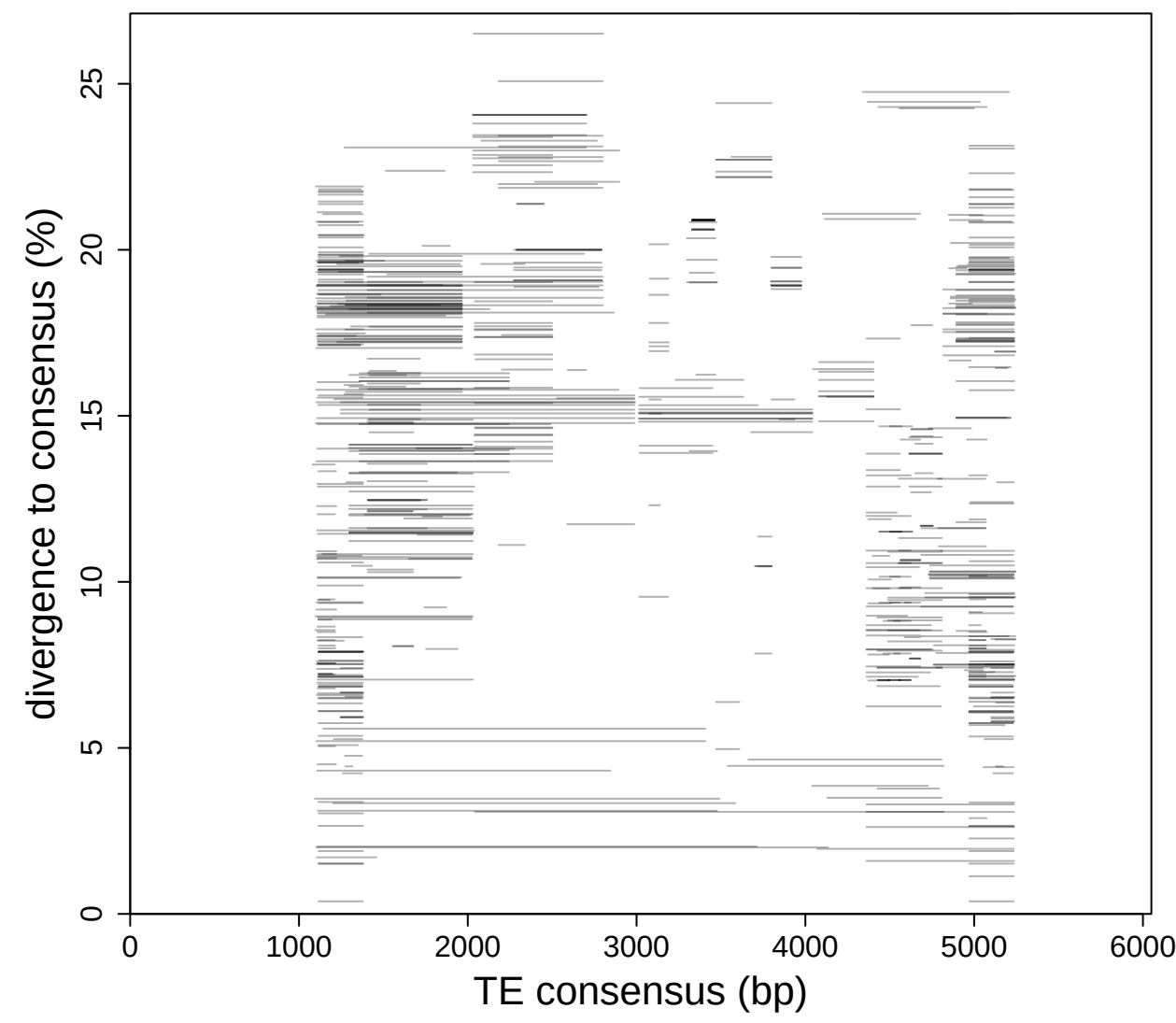
MSA length = 6050





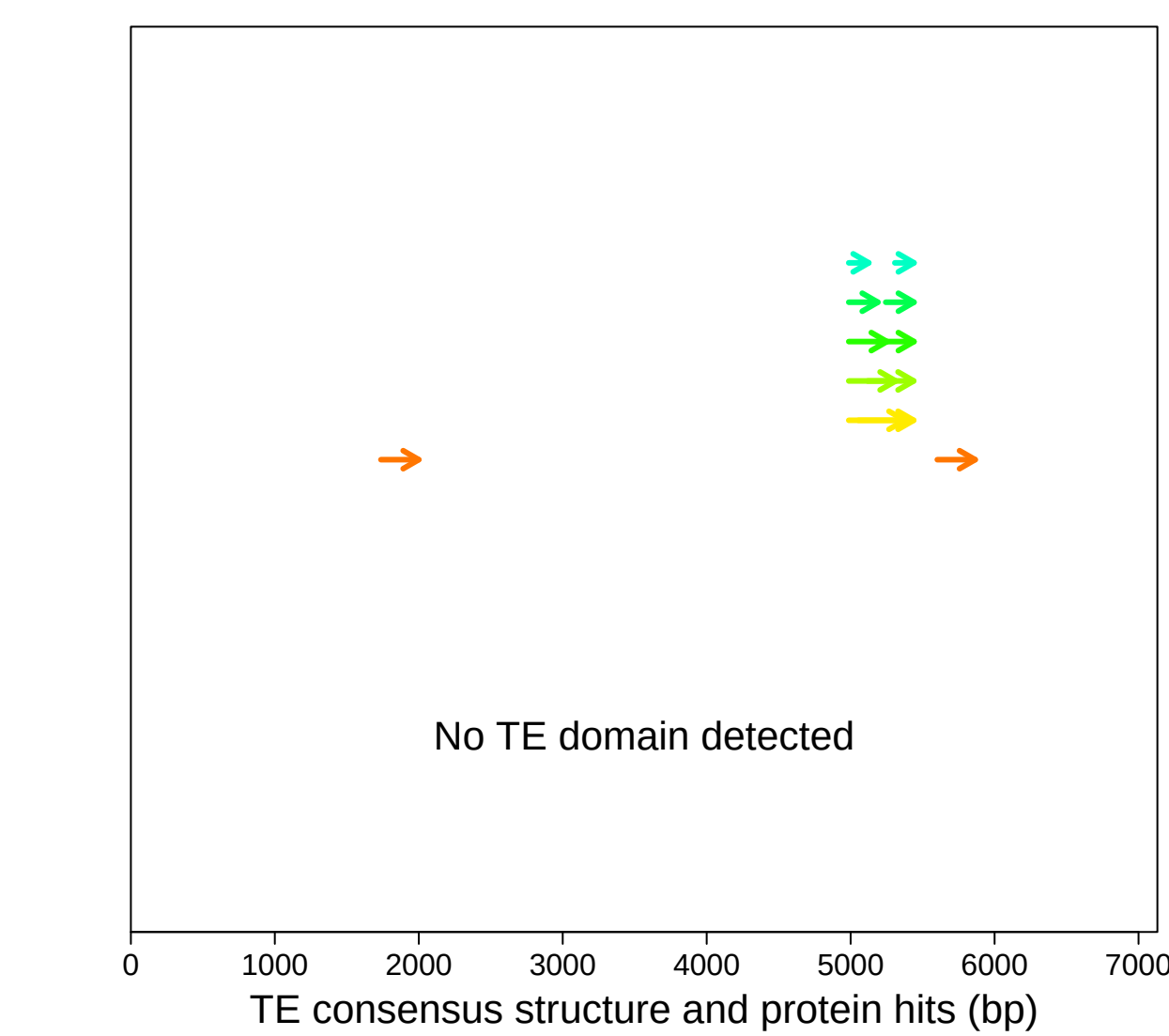
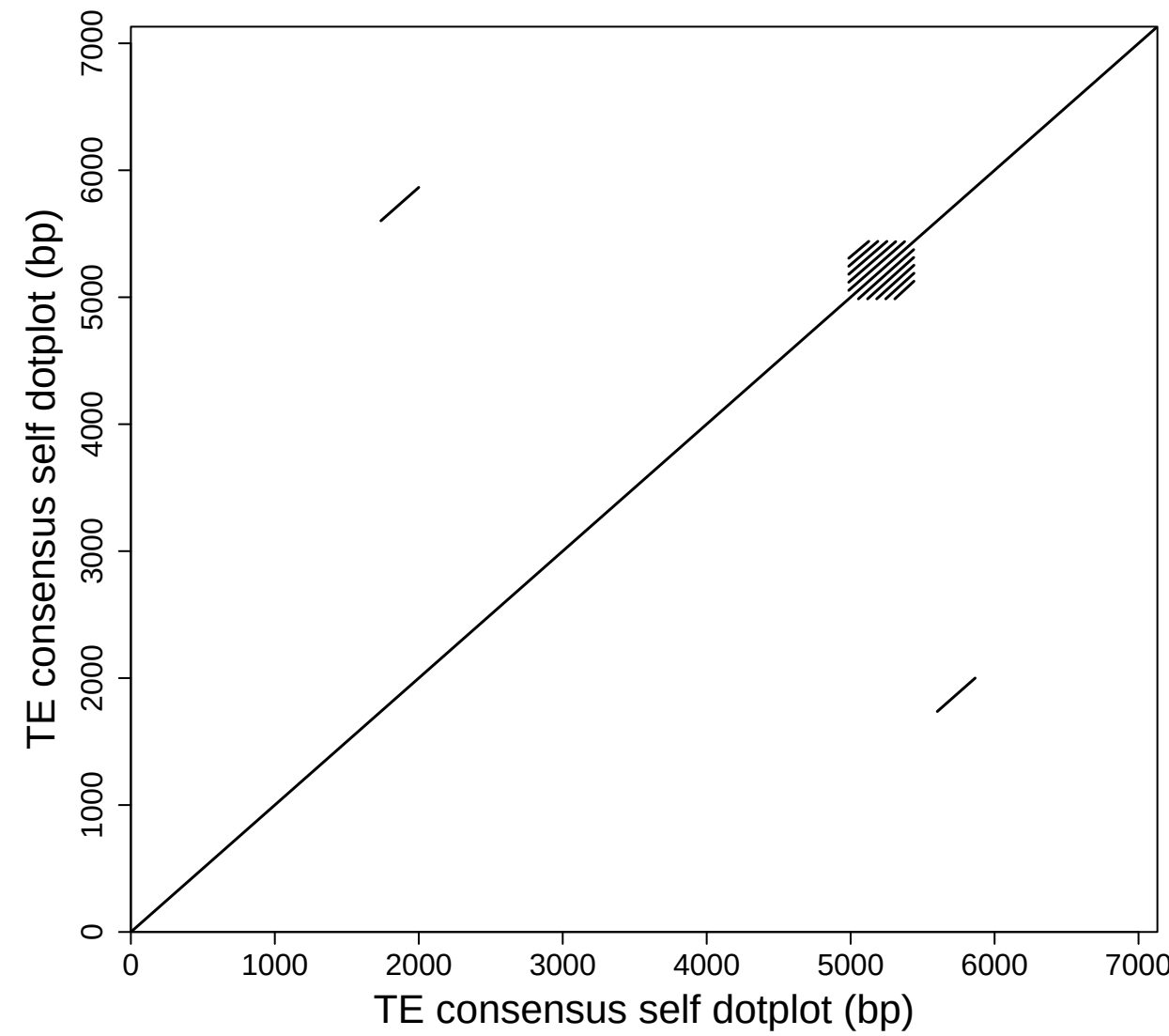
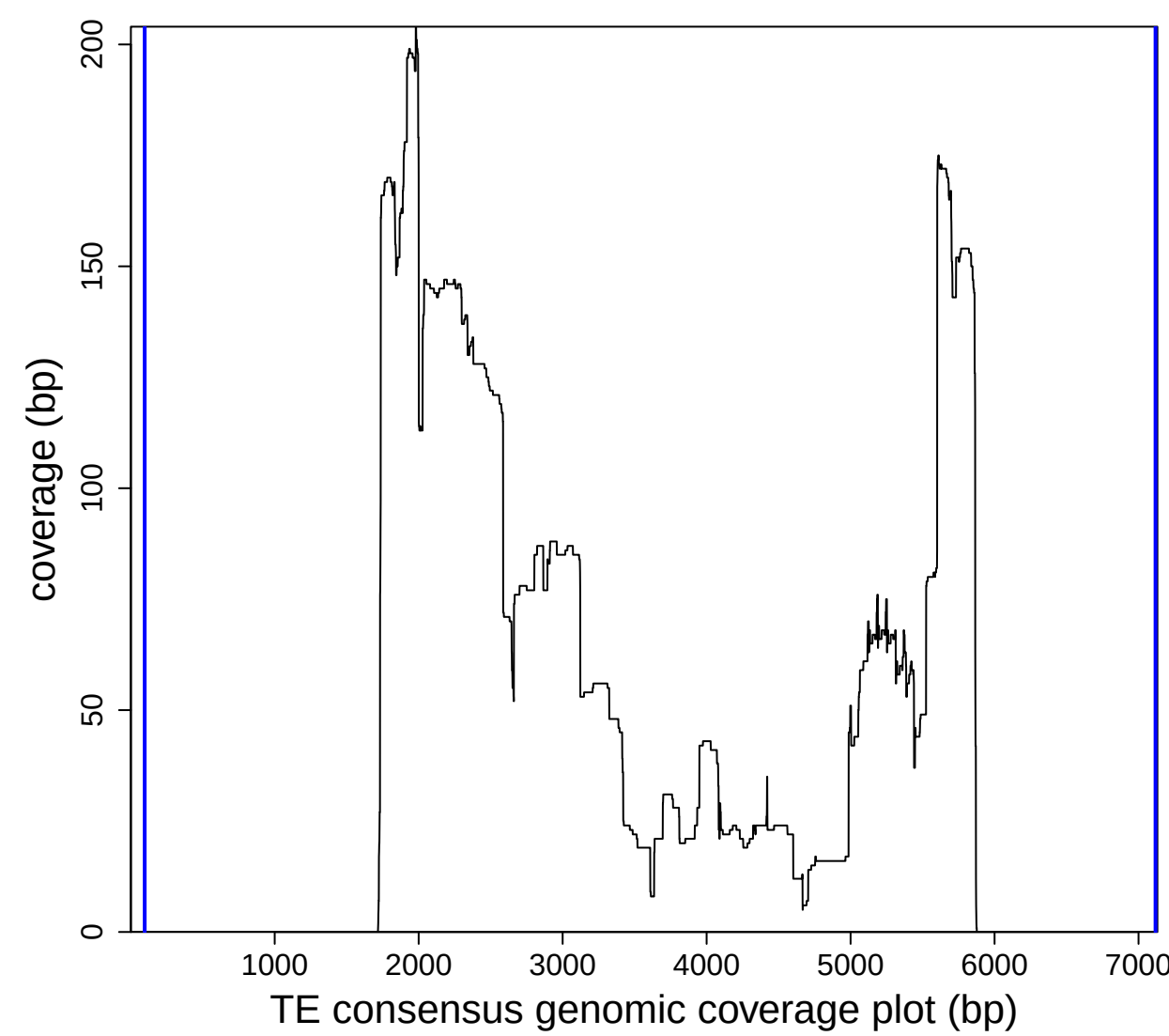
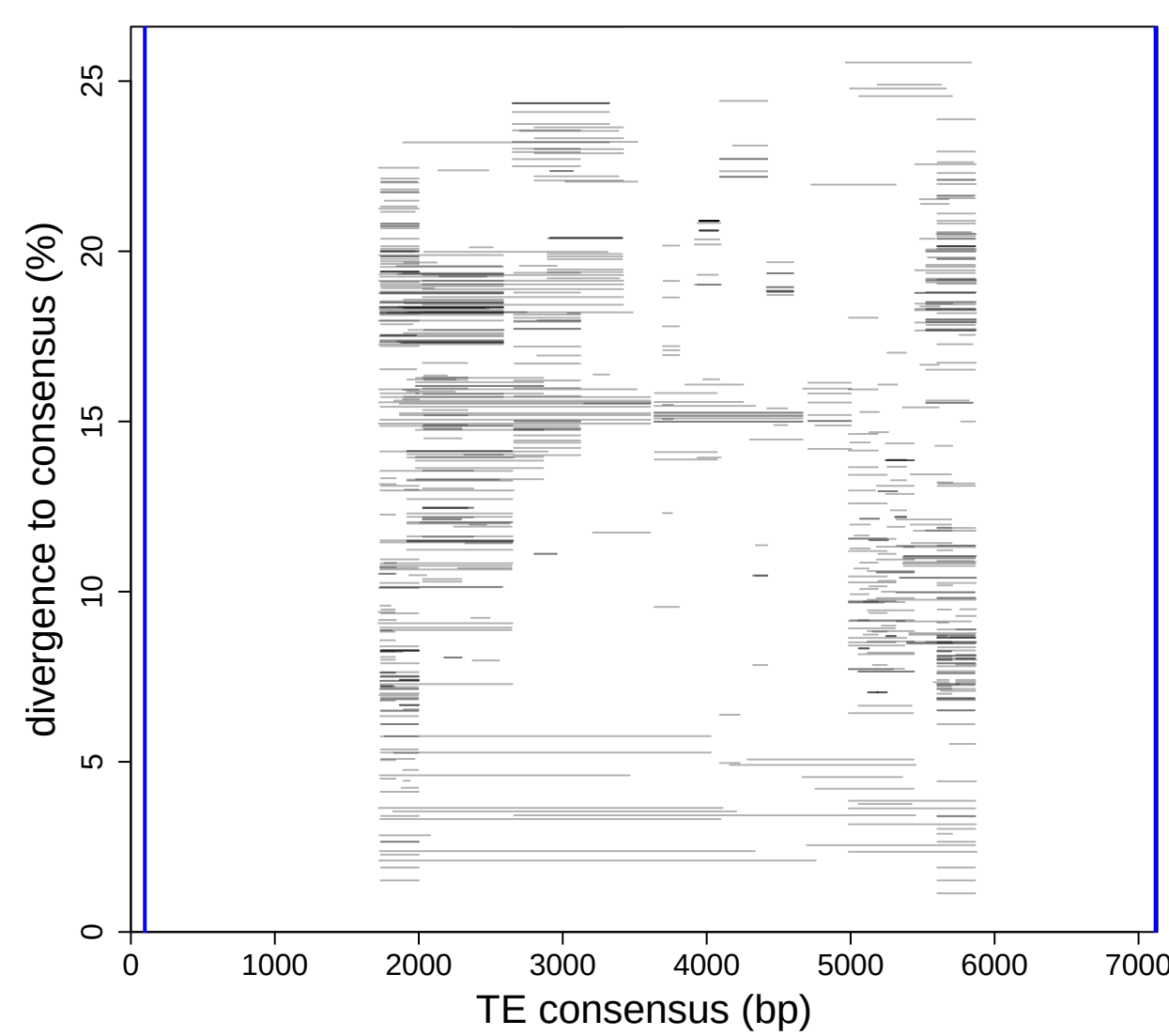
_4.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_al
size: 6050bp; fragments: 726; full length: 0 (>=5445bp)

After TEtrimmer 6050 bp

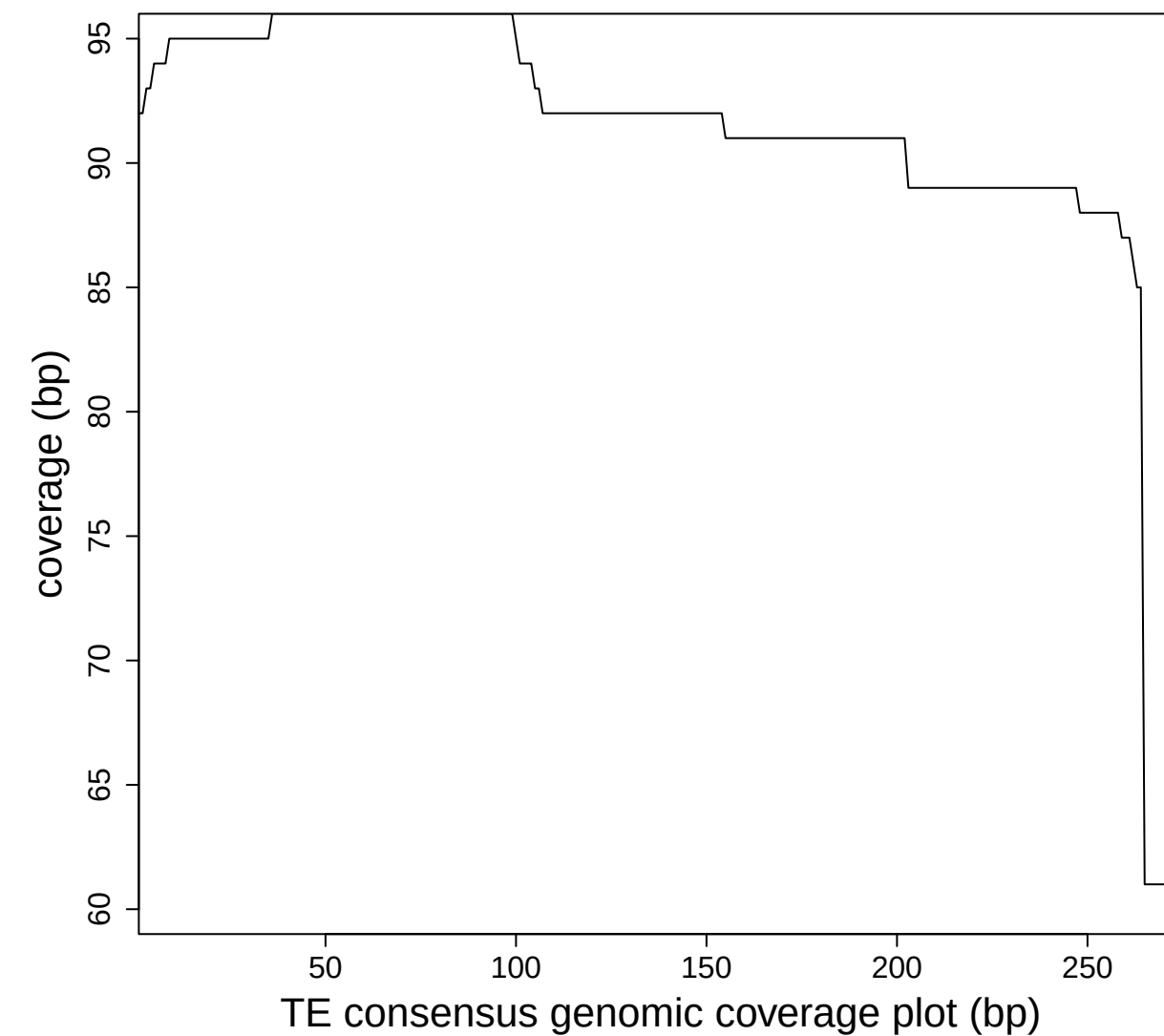
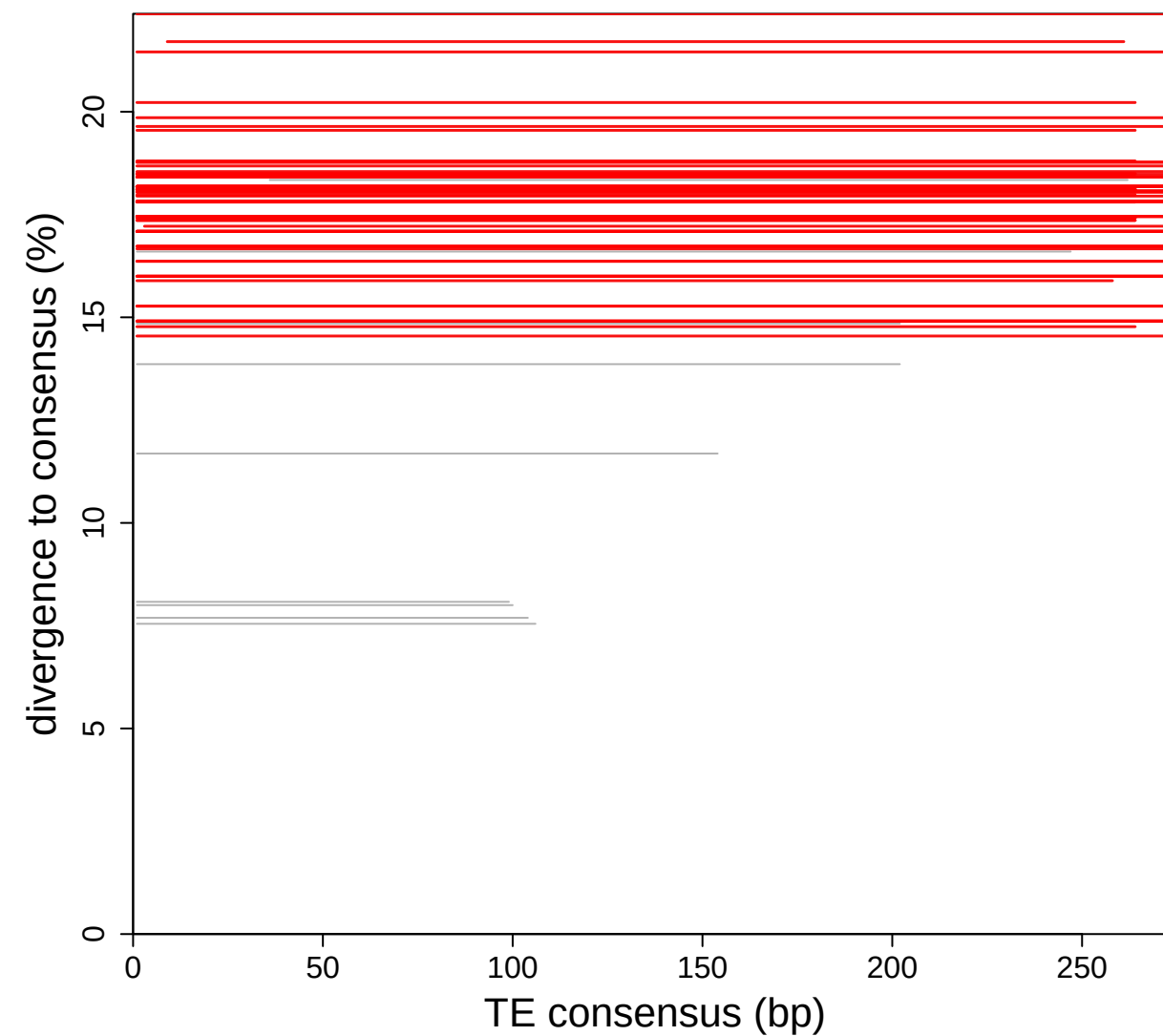


bed_g_4.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 7131bp; fragments: 721; full length: 0 (>=6417.9bp)

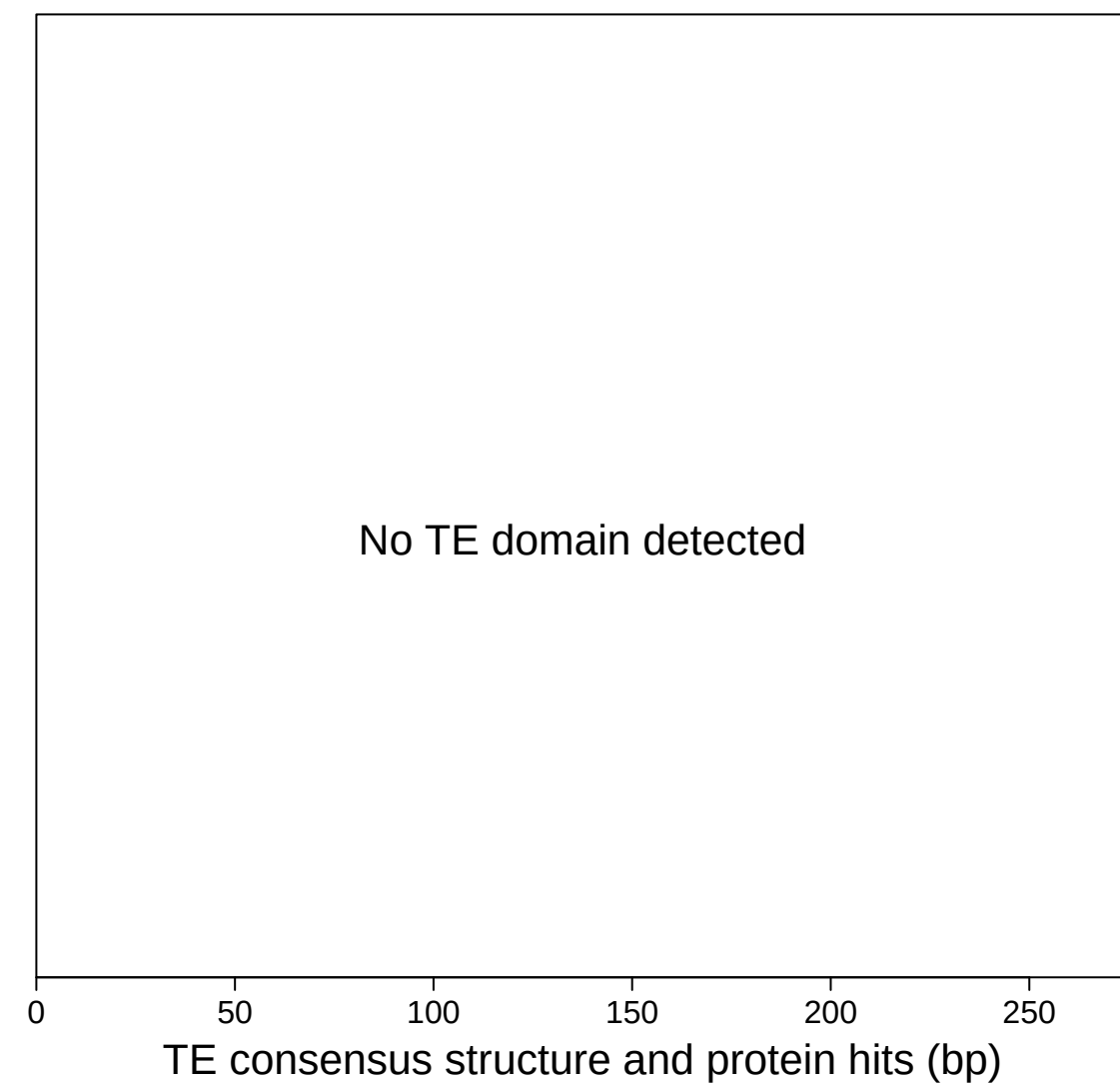
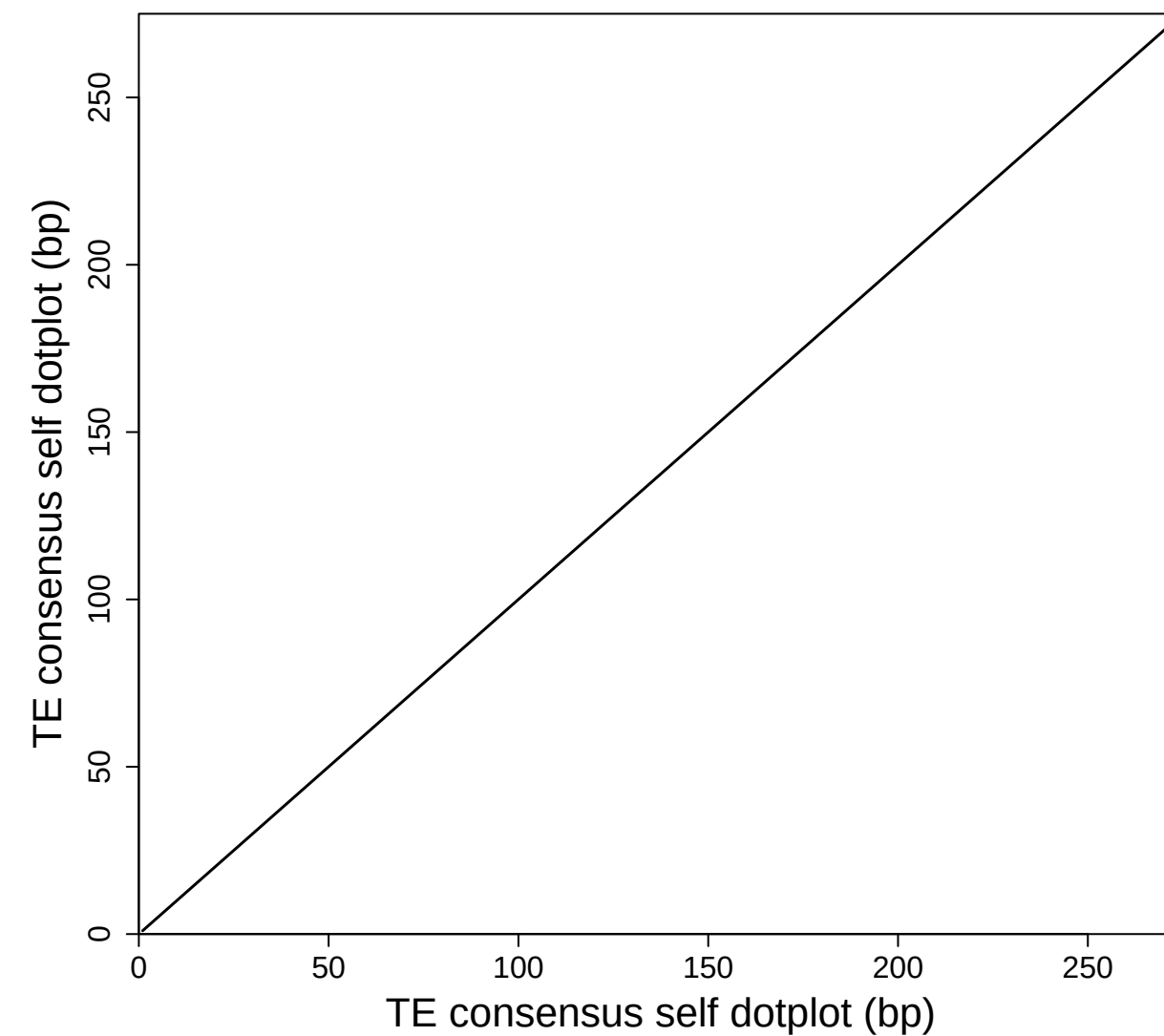
After TEtrimmer Extended plot Blue lines are boundaries



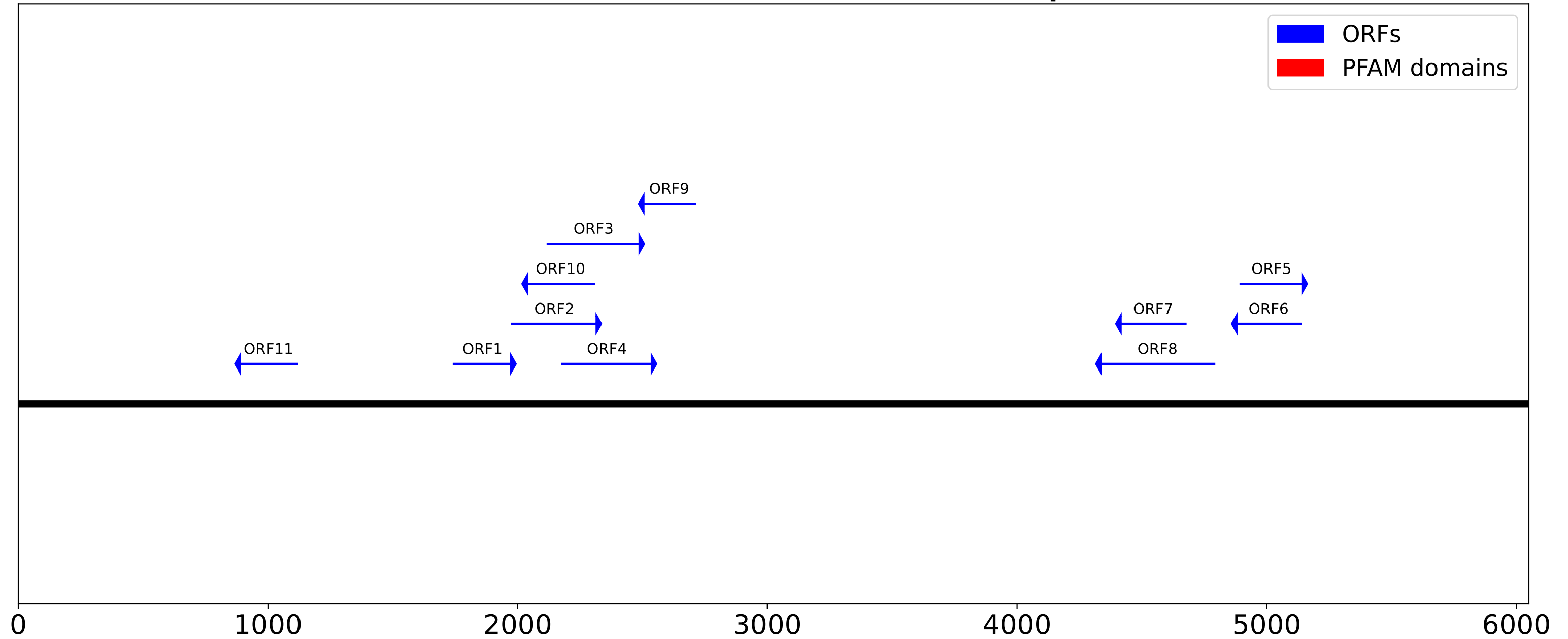
TE: ltr_1_family_307
size: 275bp; fragments: 96; full length: 87 (≥ 247.5 bp)



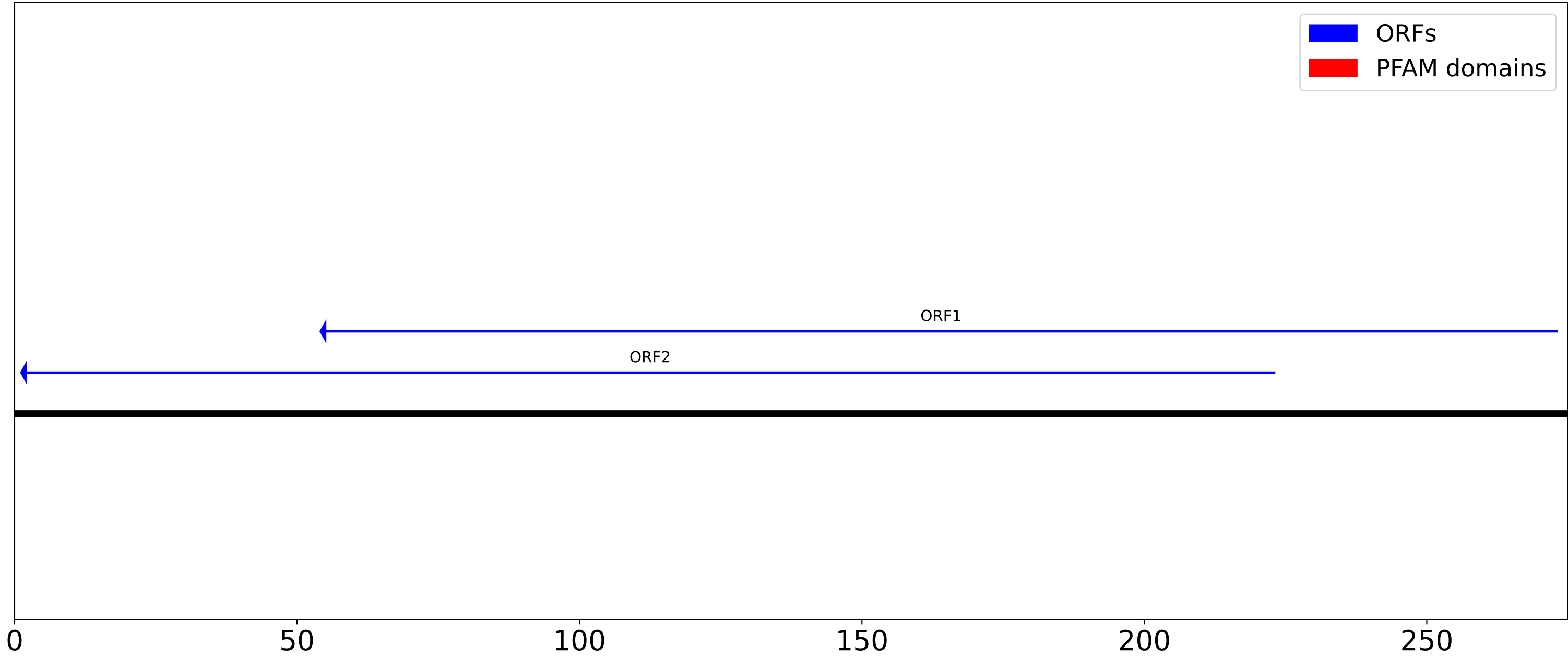
Before TEtrimmer 275 bp



After TETrimmer ORF and PFAM domain plot



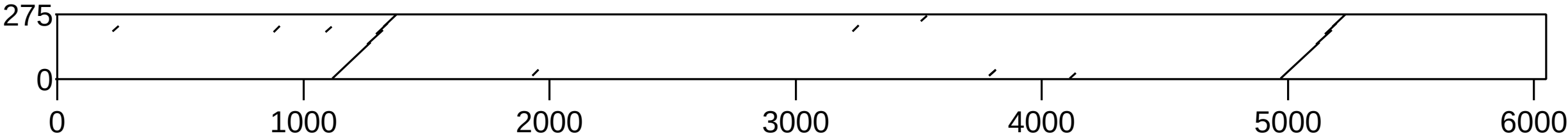
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)