

Start crop Point

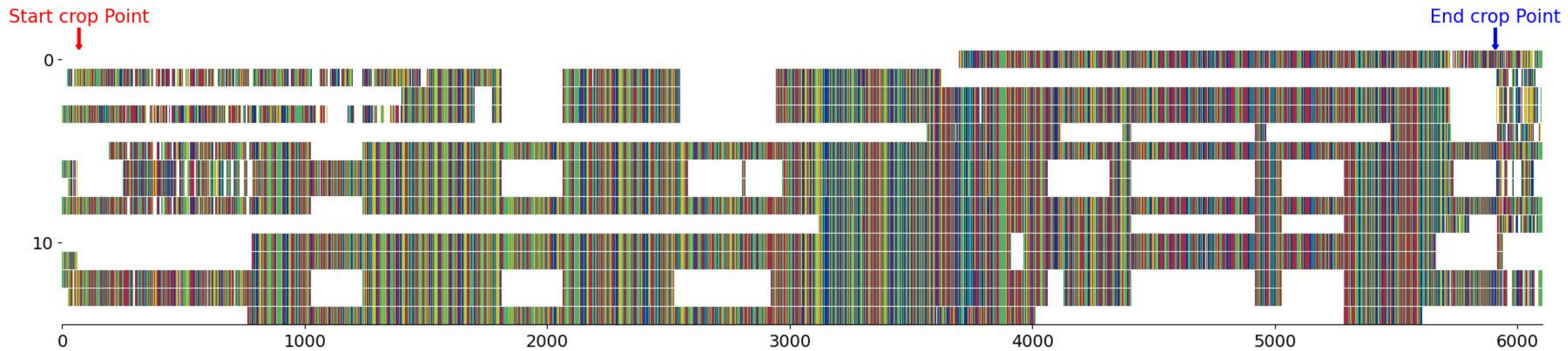


End crop Point



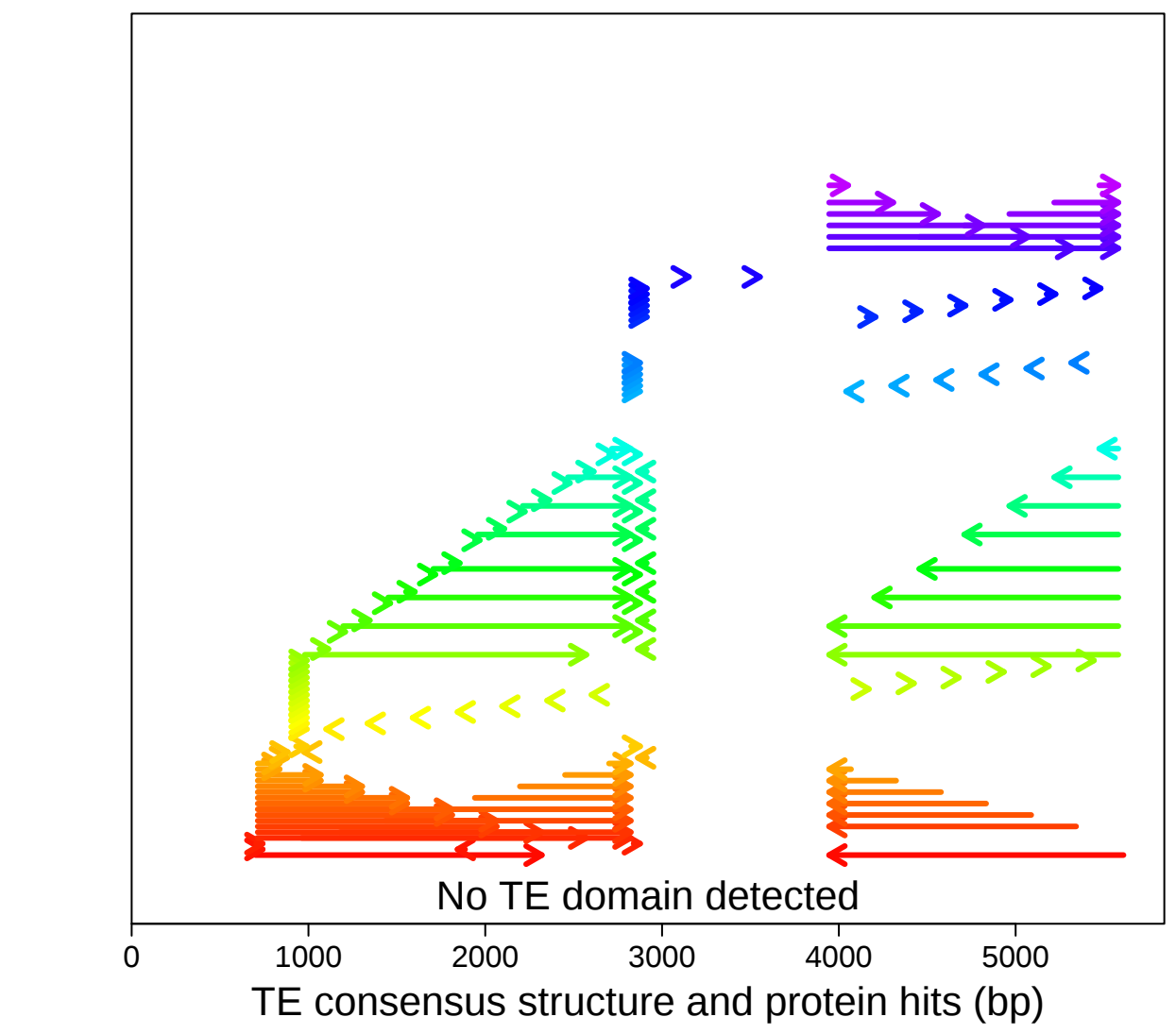
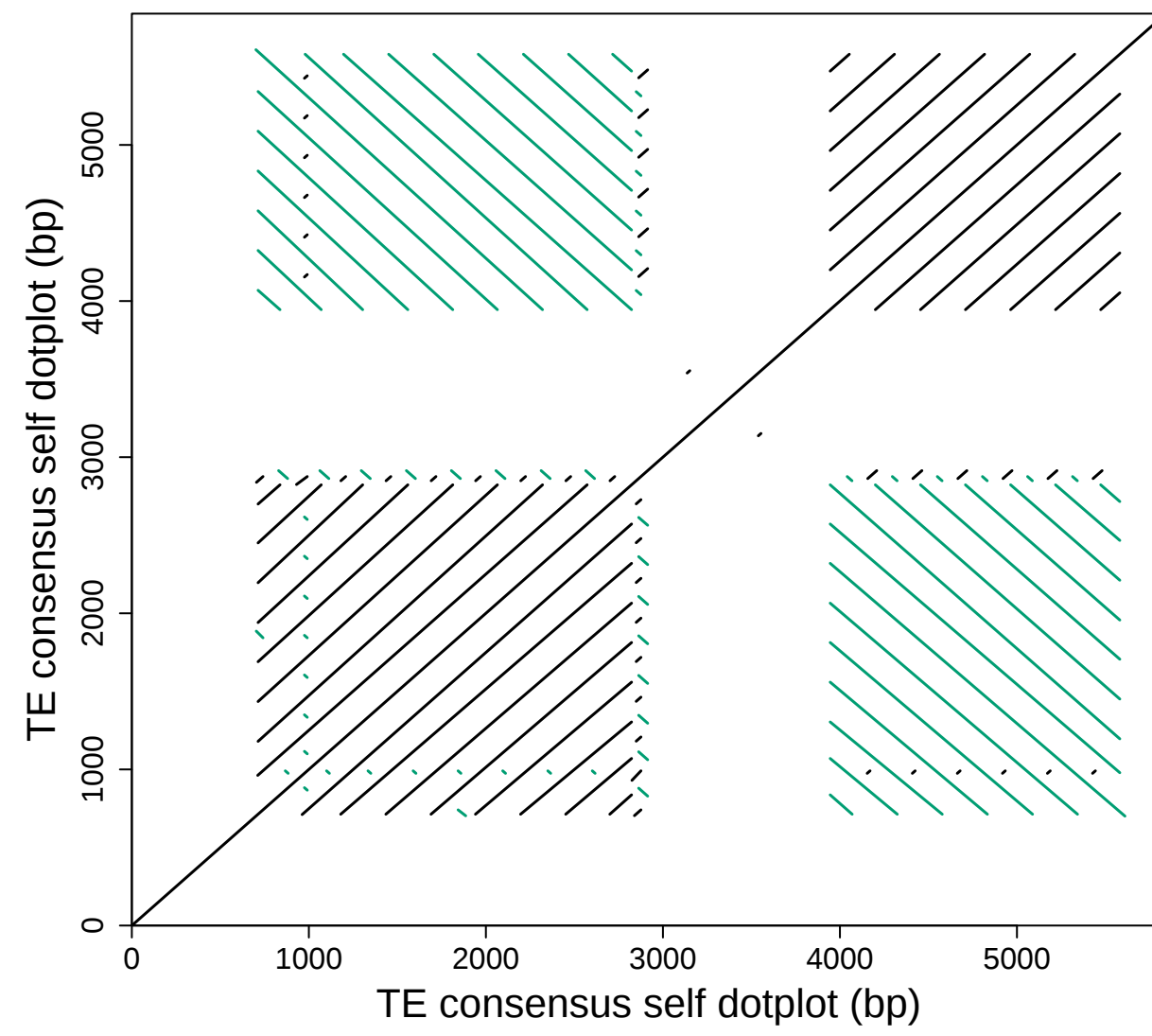
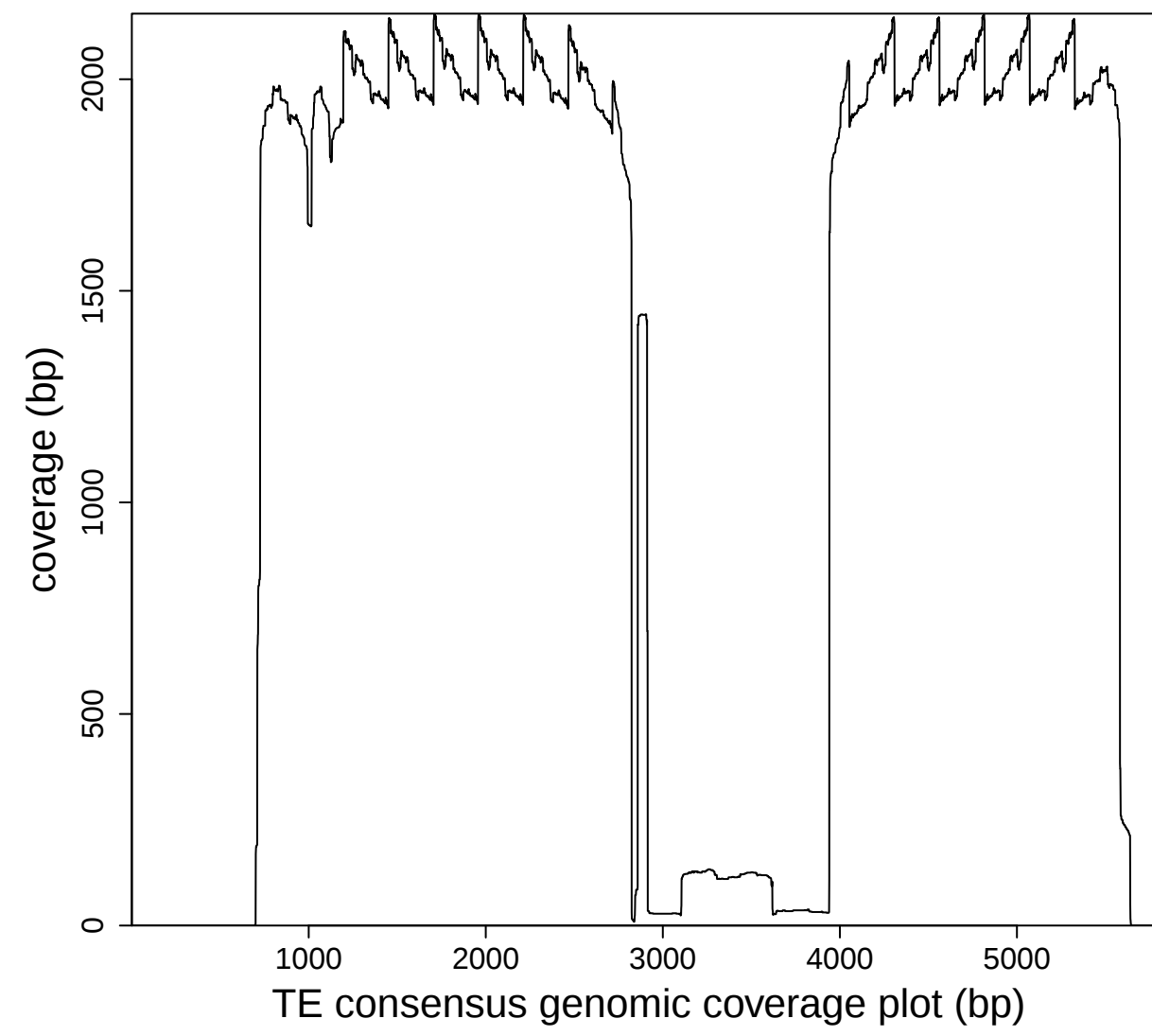
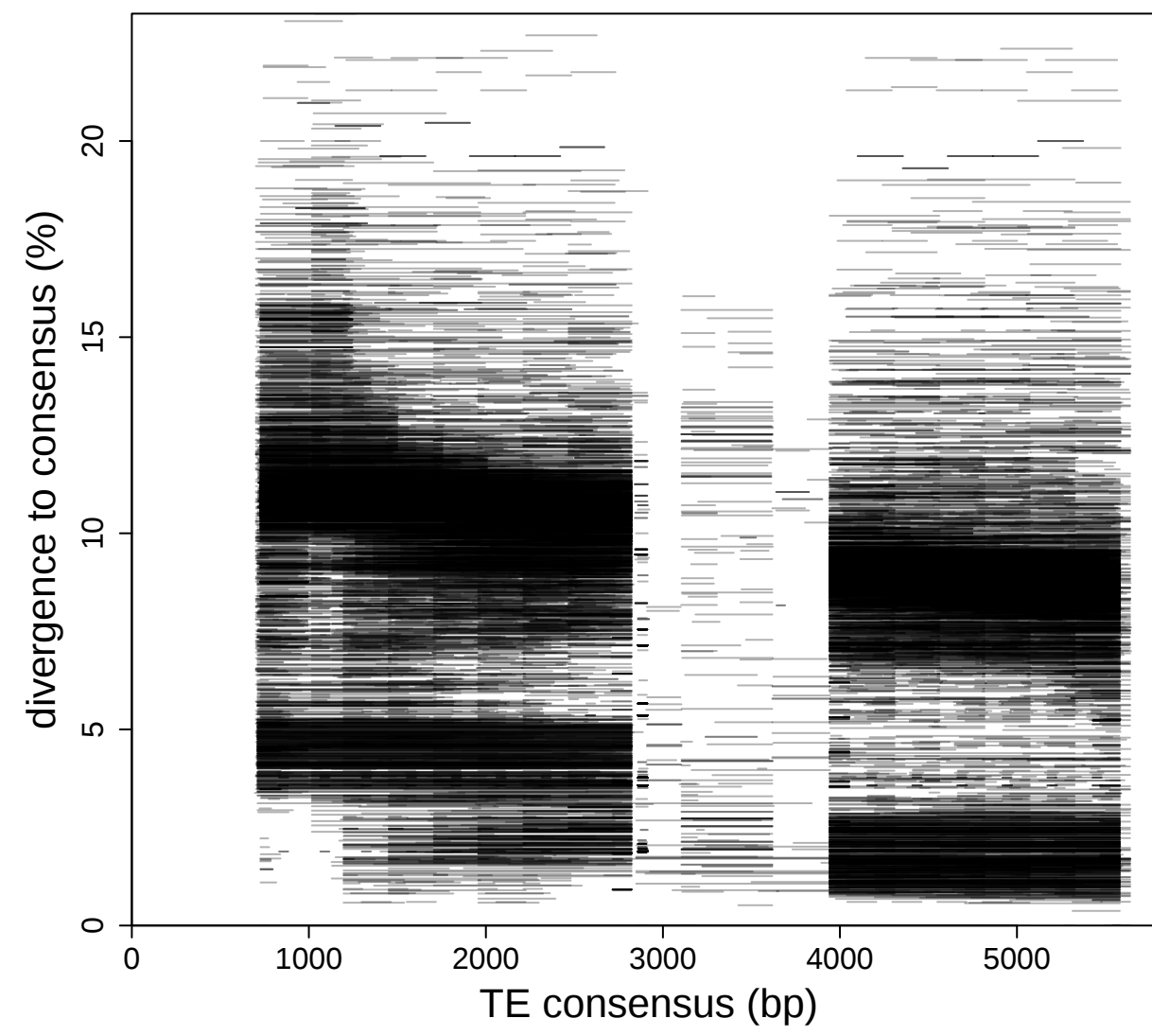
MSA length = 5841





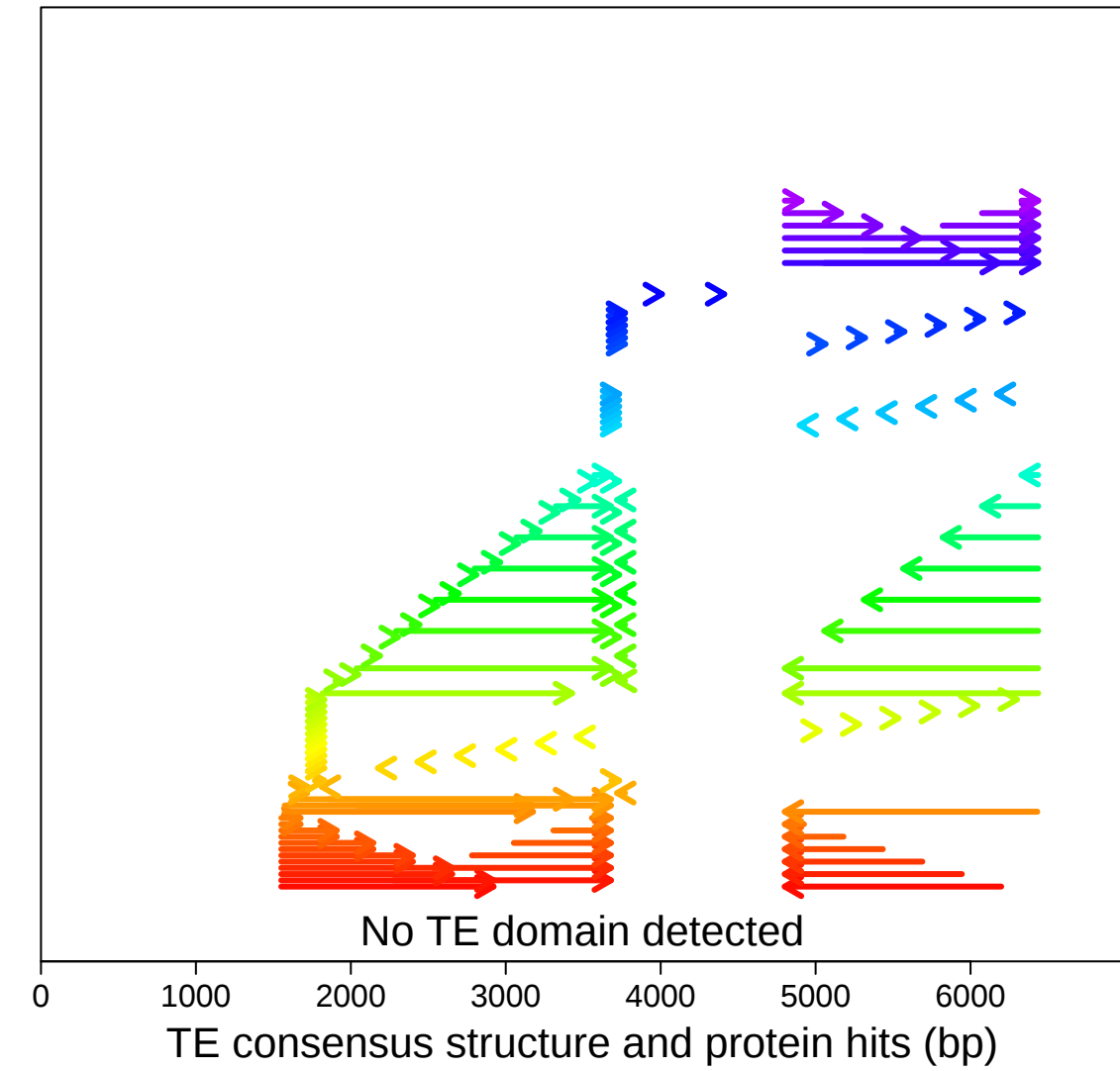
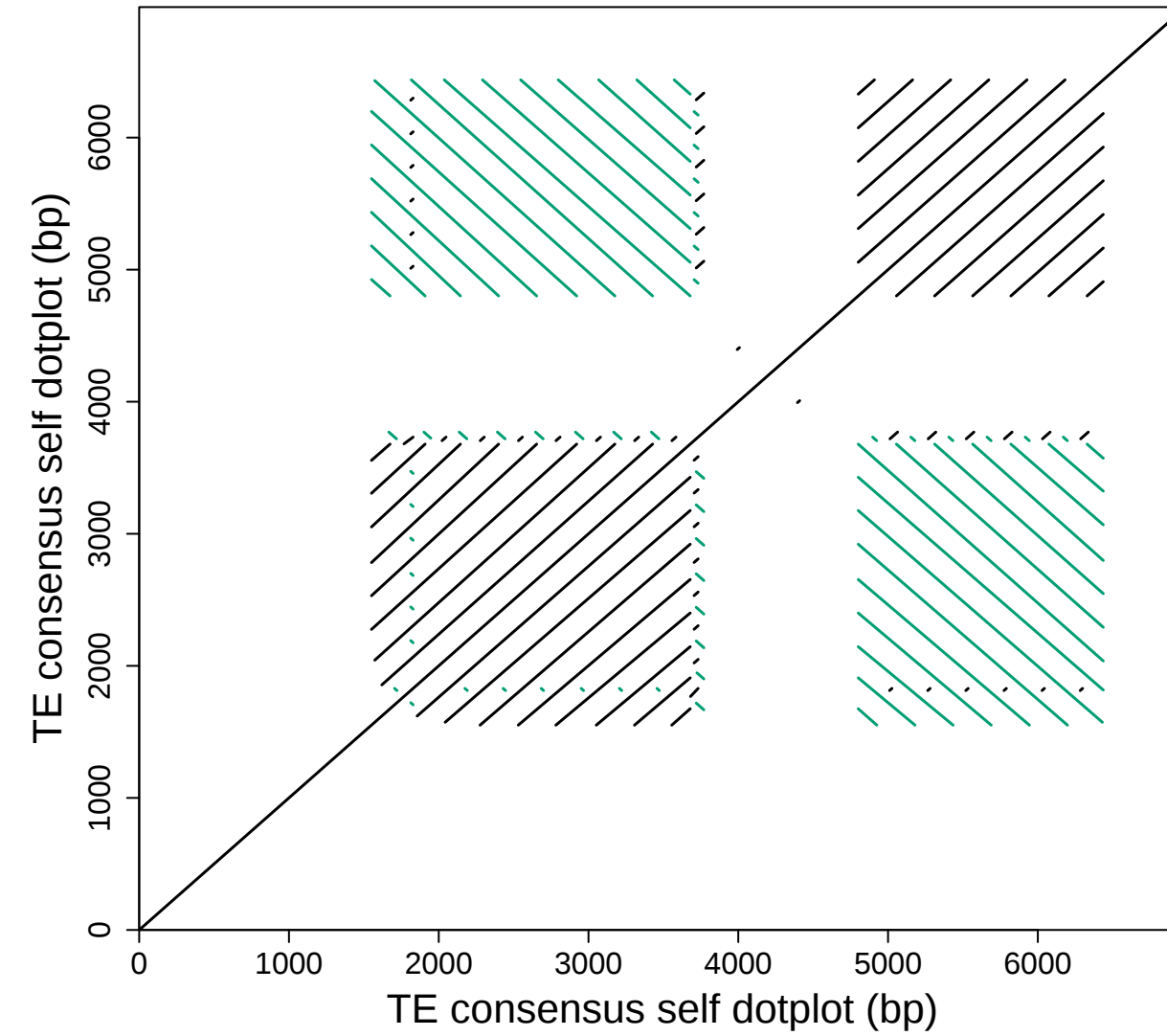
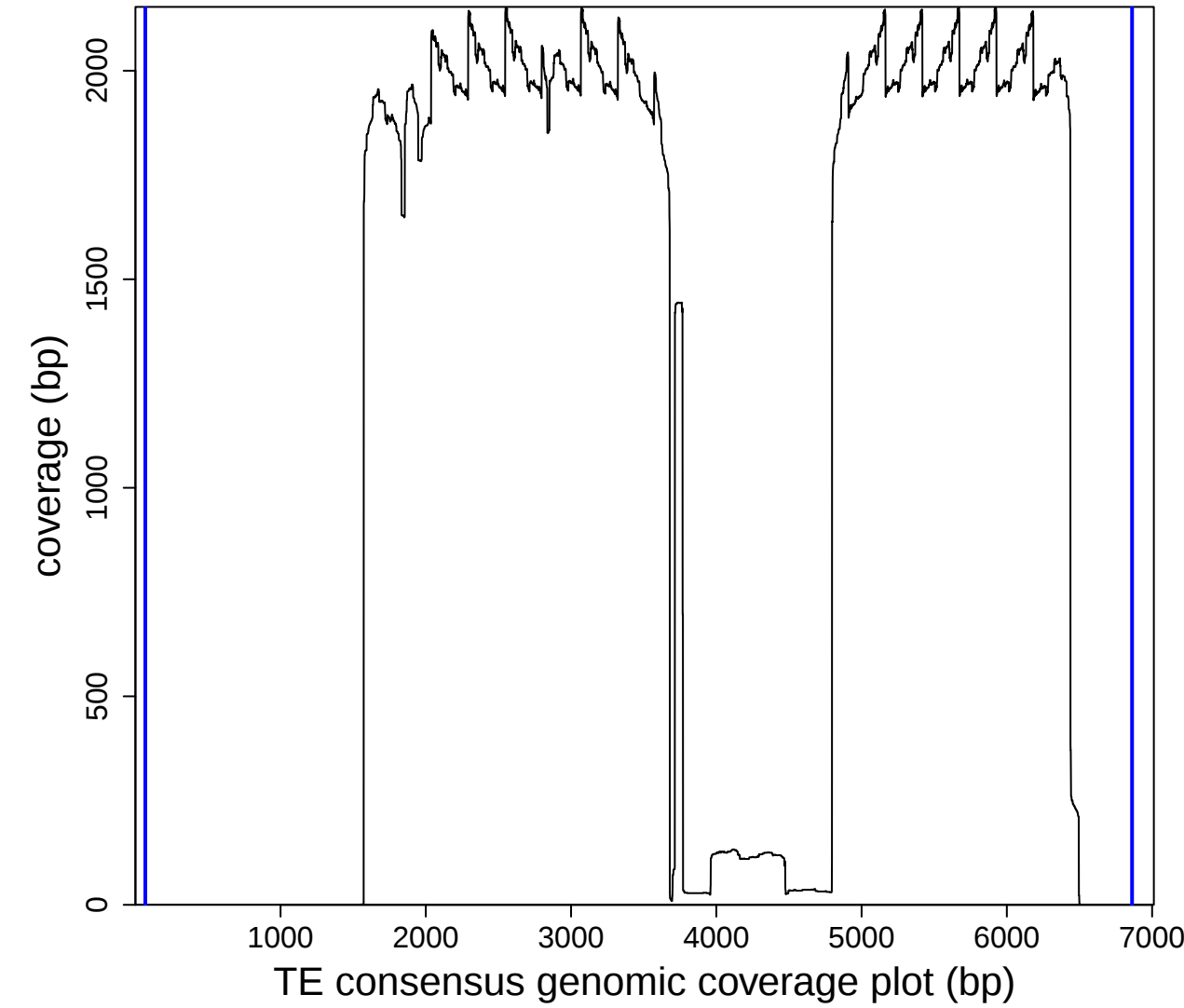
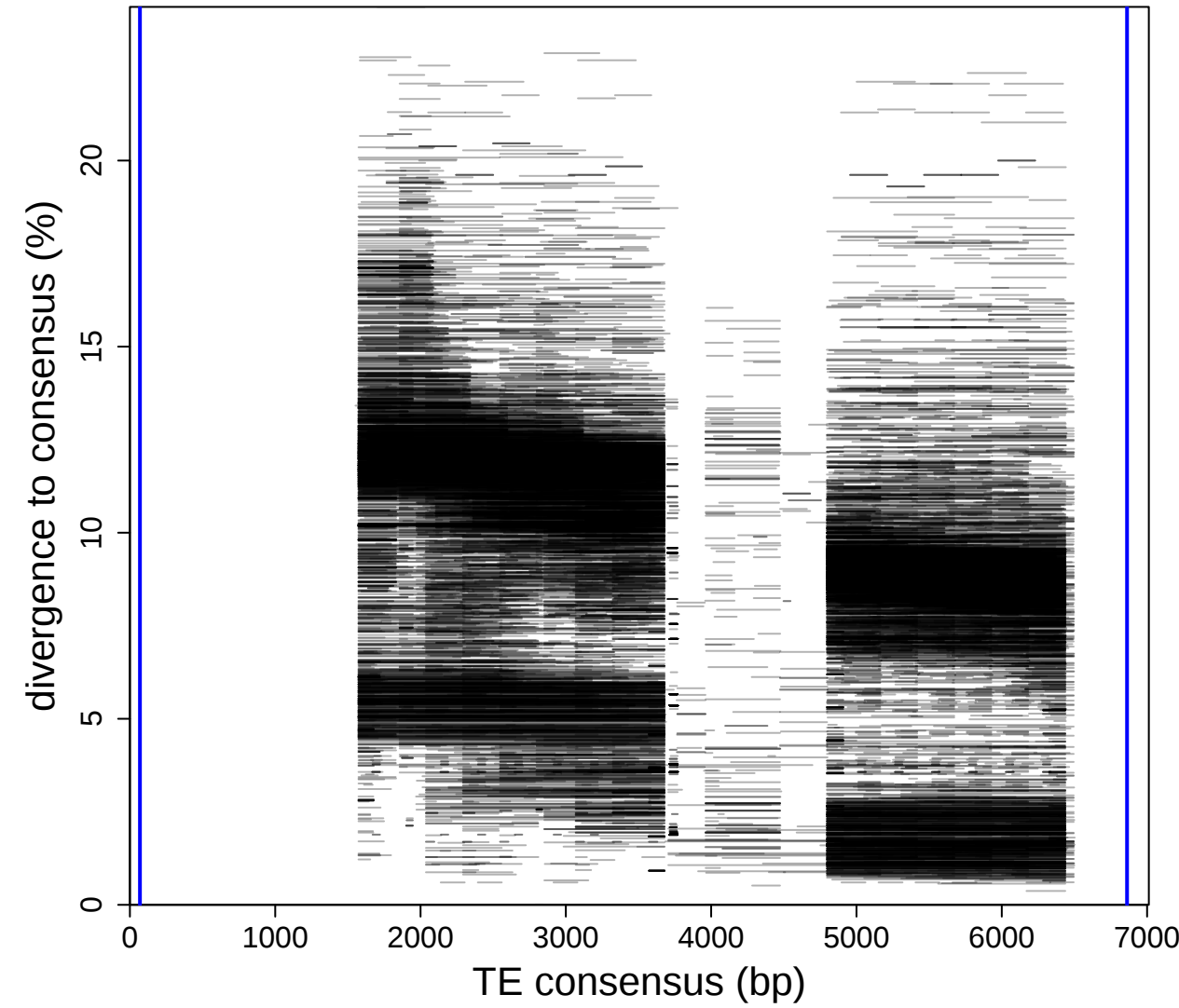
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_al
size: 5841bp; fragments: 11876; full length: 0 (>=5256.9bp)

After TEtrimmer 5841 bp

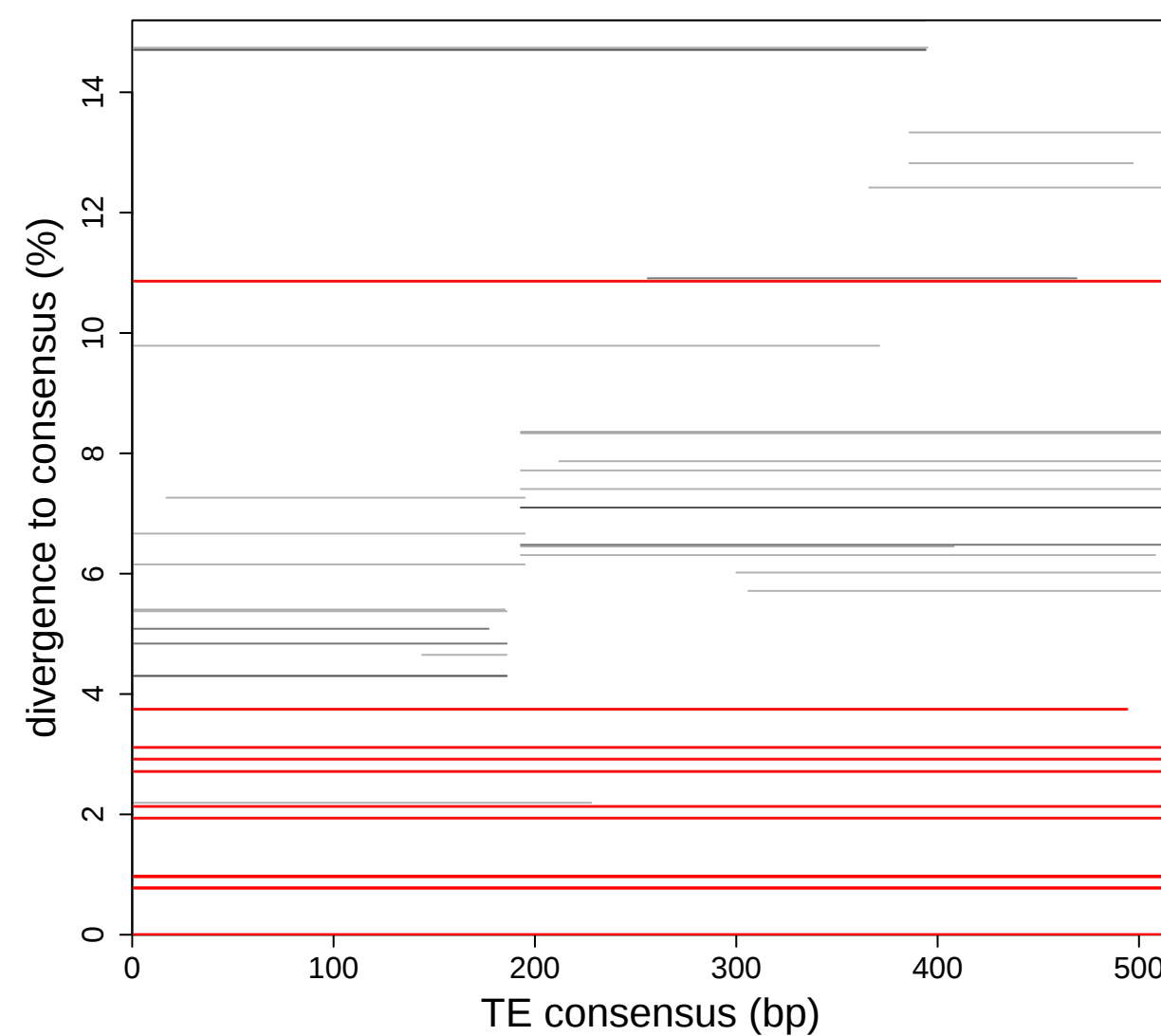


bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 7012bp; fragments: 11813; full length: 0 (>=6310.8bp)

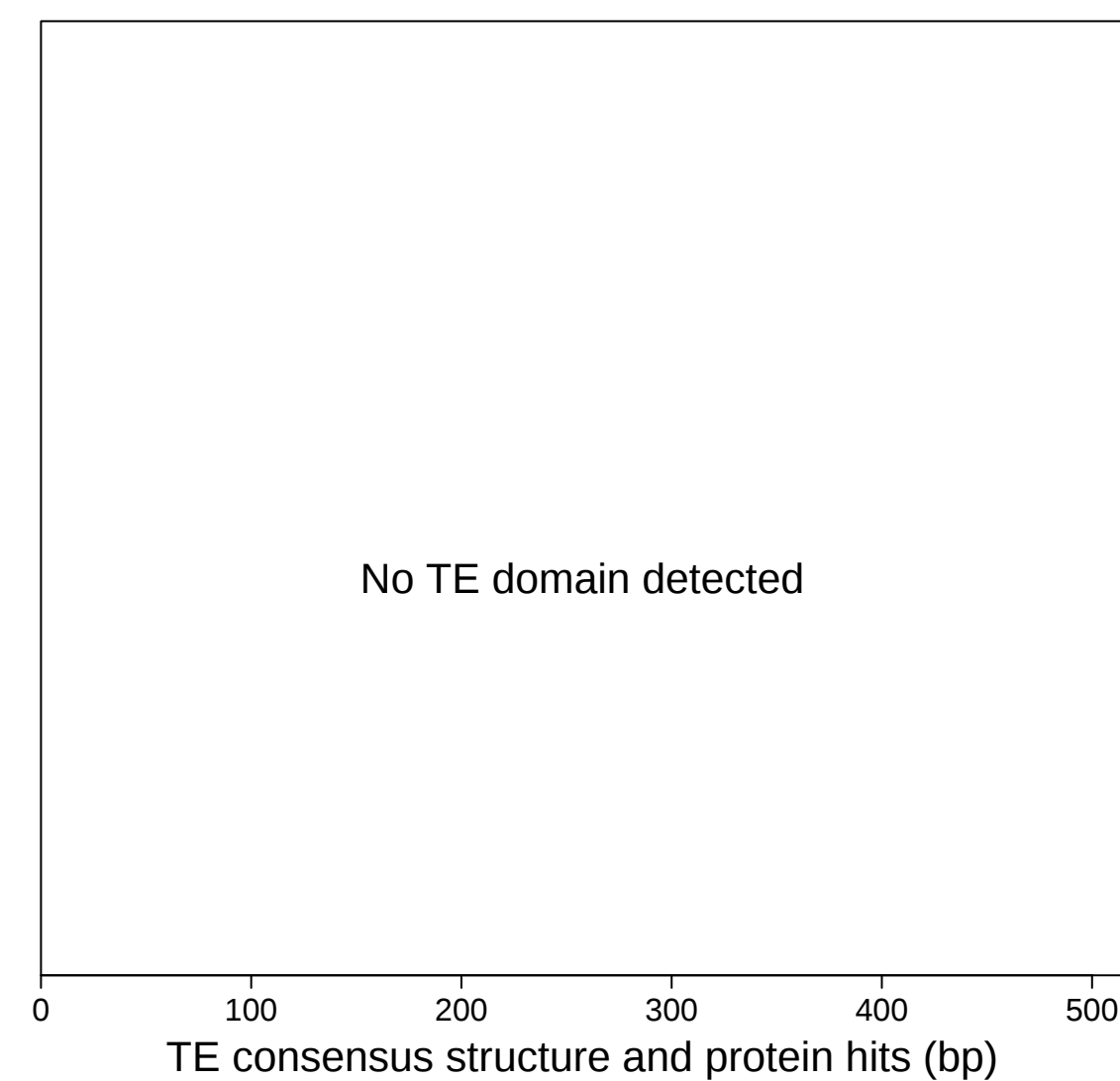
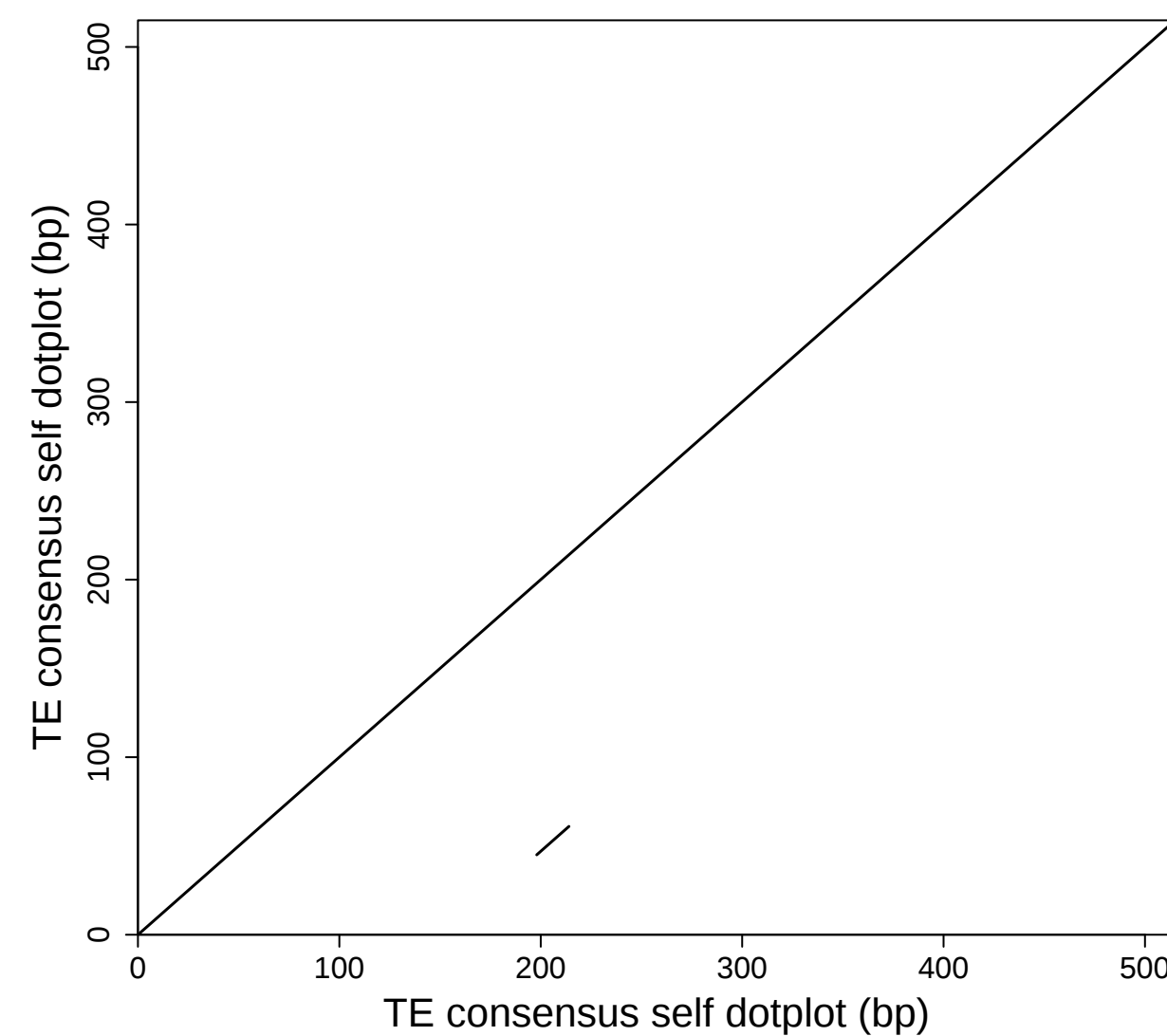
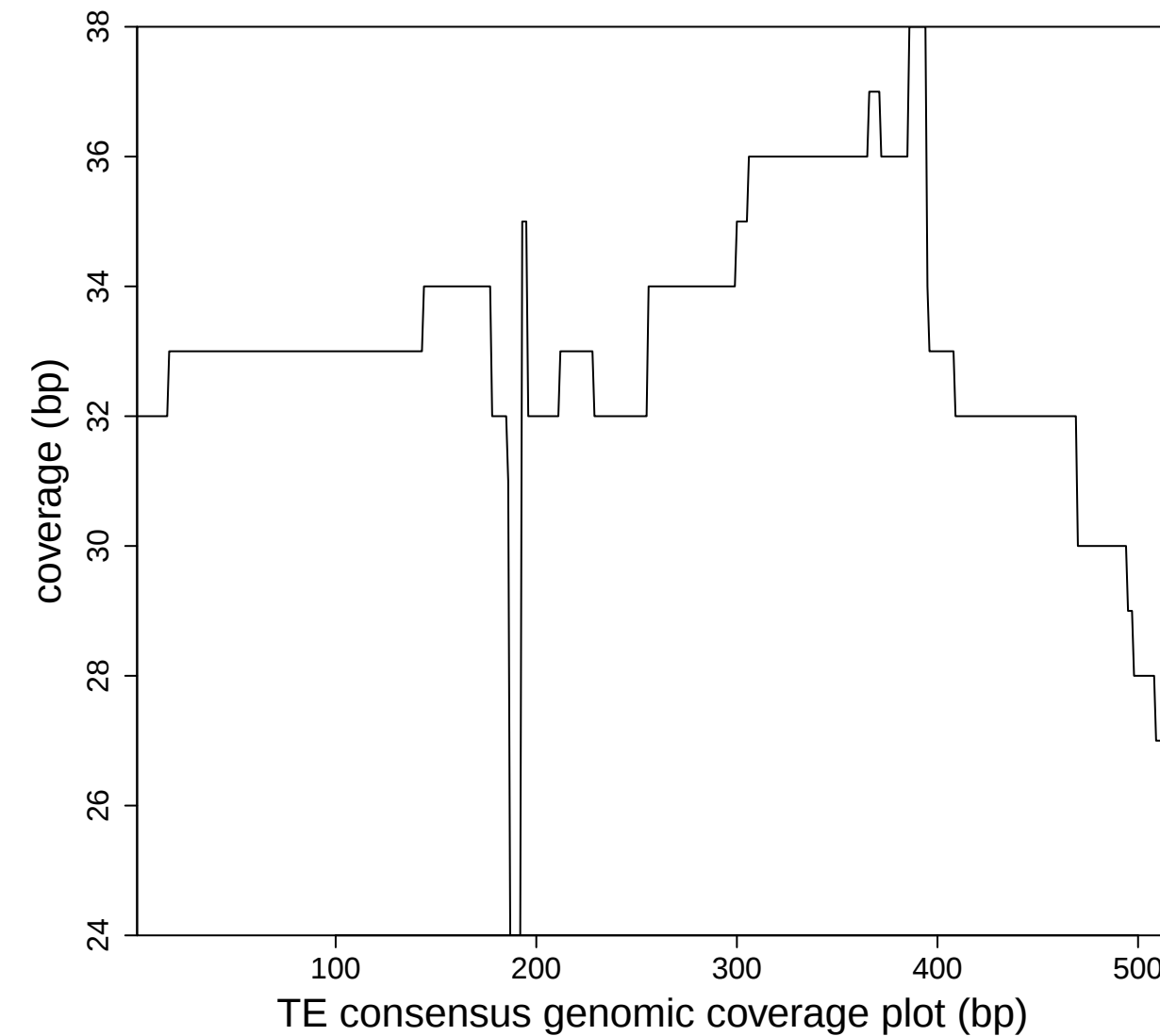
After TEtrimmer Extended plot Blue lines are boundaries



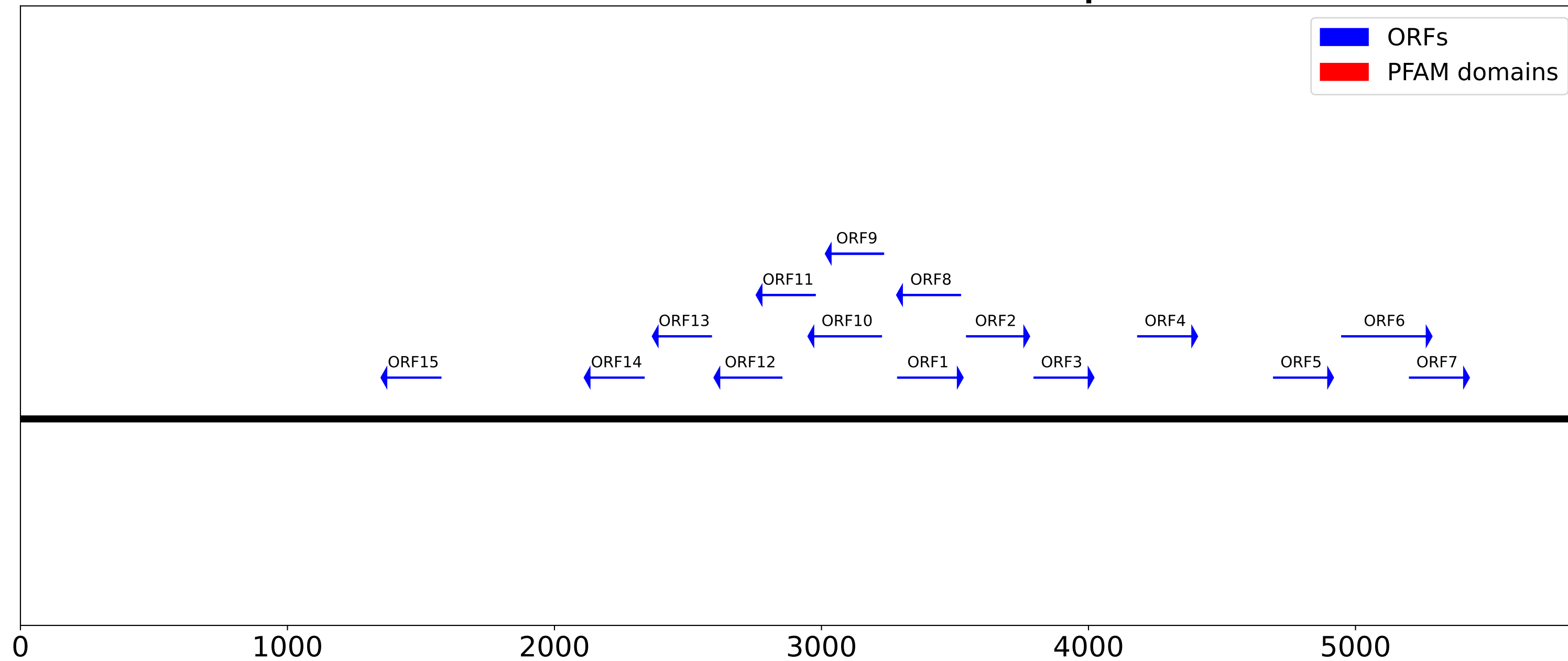
TE: ltr_1_family_229
size: 515bp; fragments: 53; full length: 14 (≥ 463.5 bp)



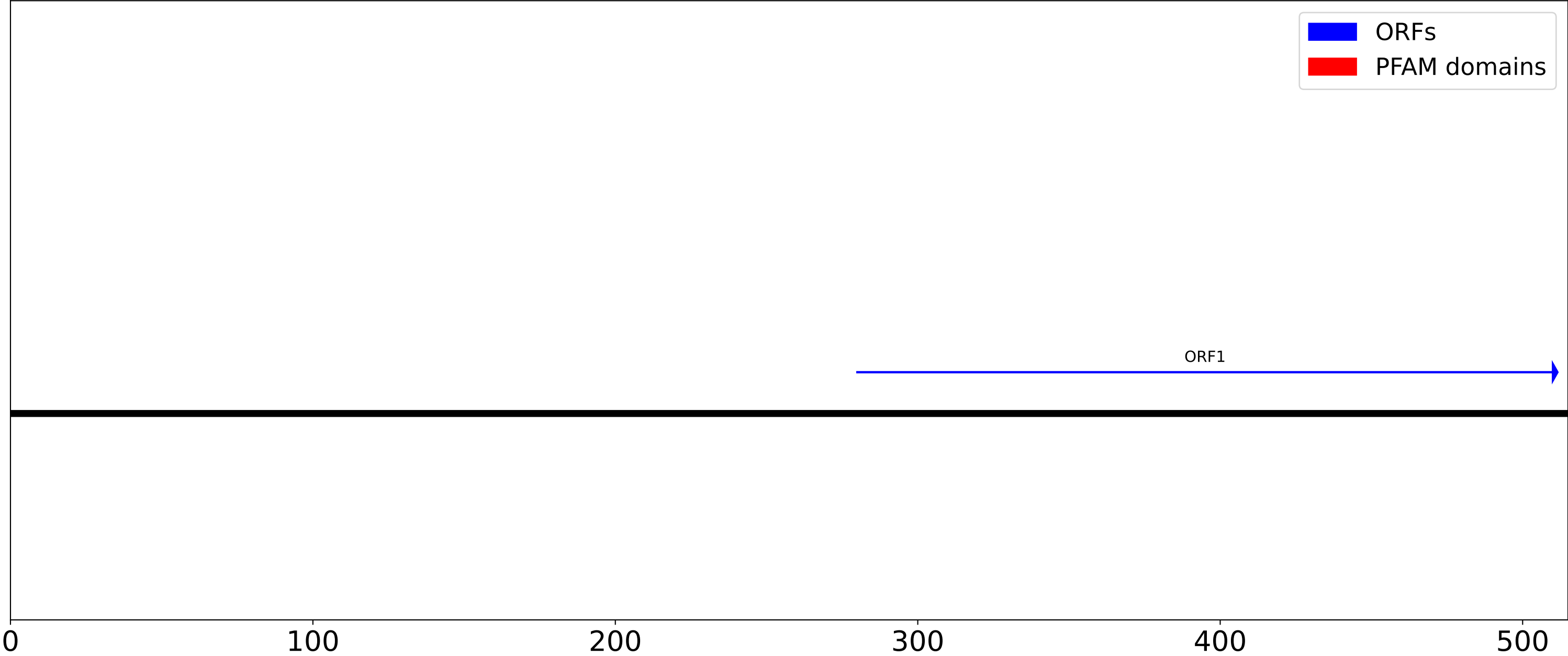
Before TEtrimmer 515 bp



After TEtrimmer ORF and PFAM domain plot



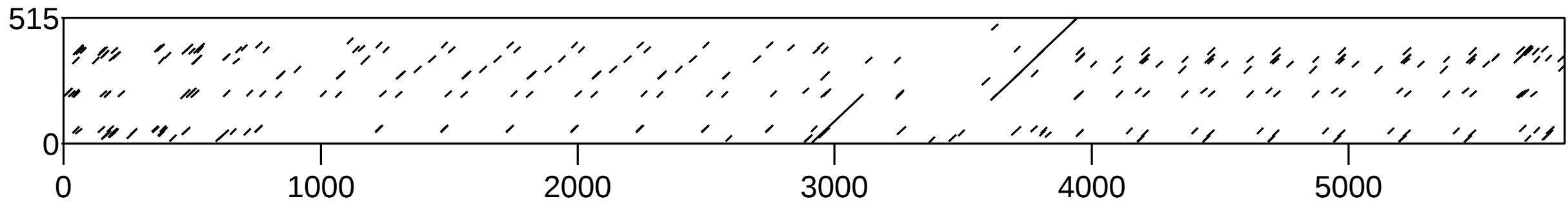
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

ensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)