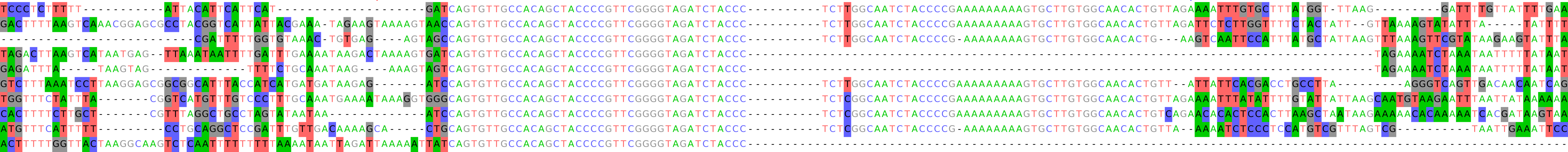


Start Crop Point

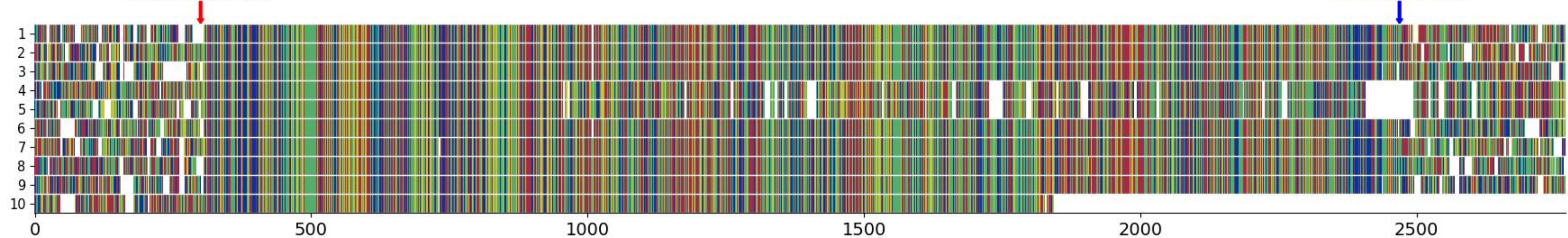
End Crop Point

MSA length = 2170



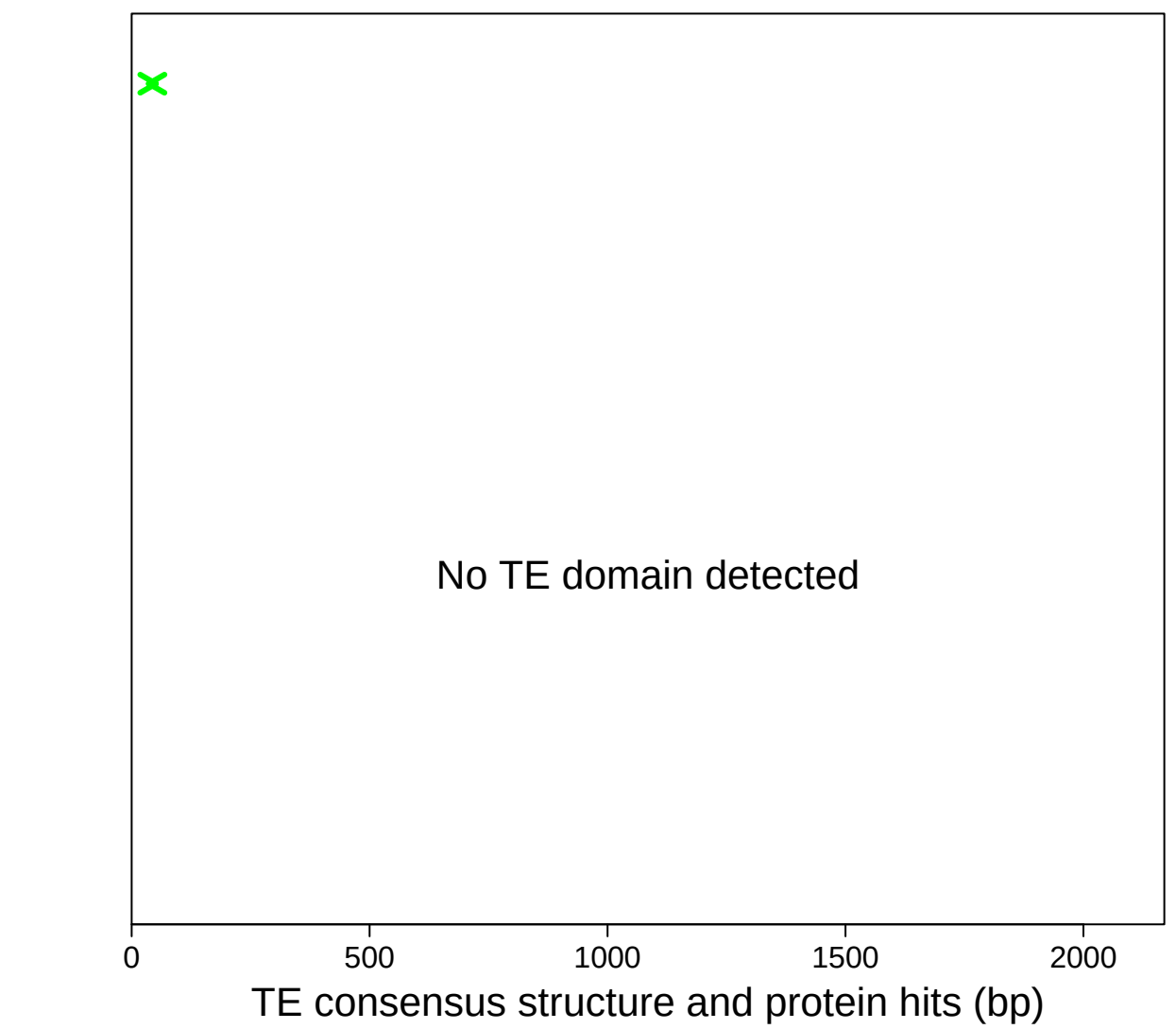
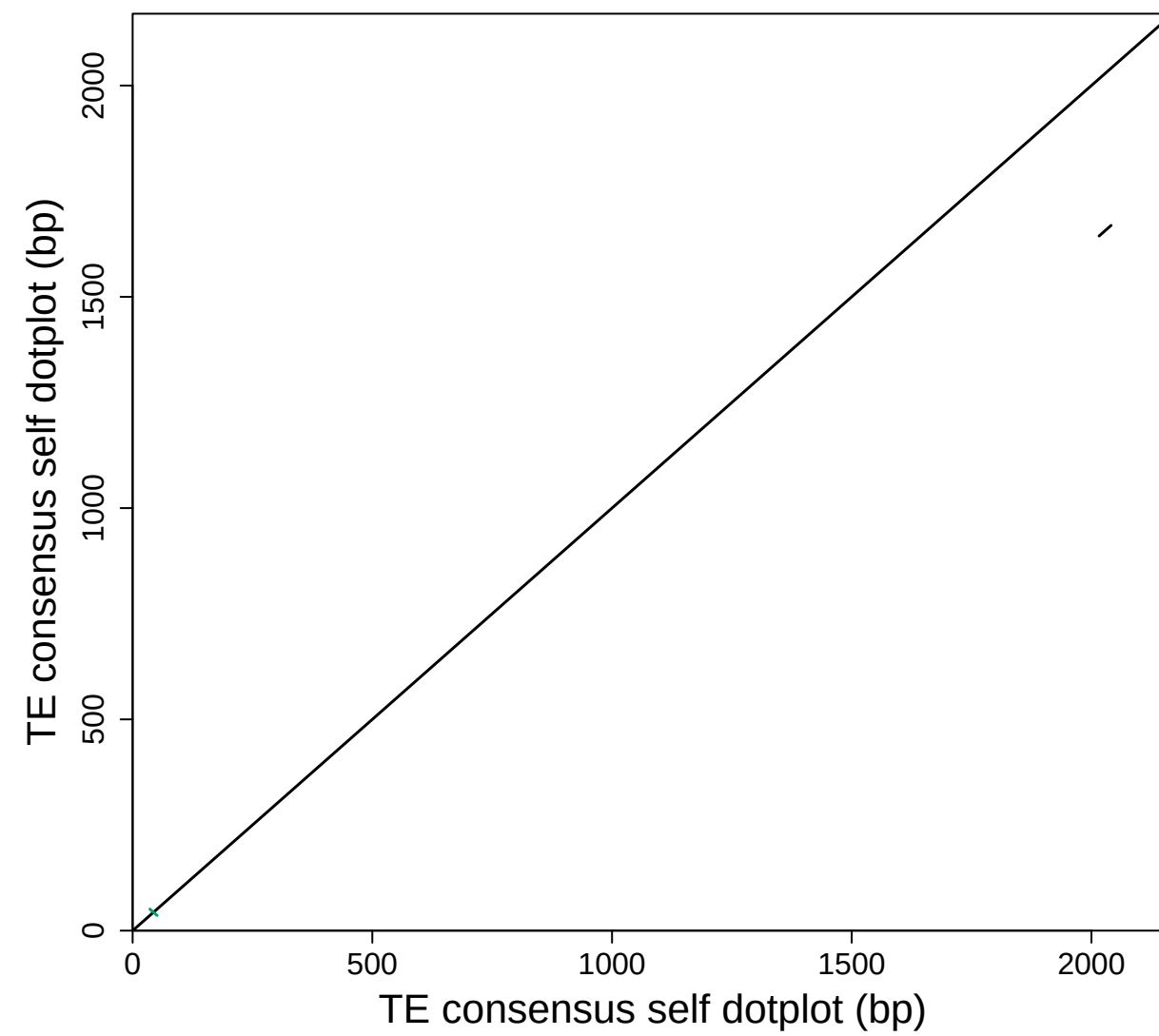
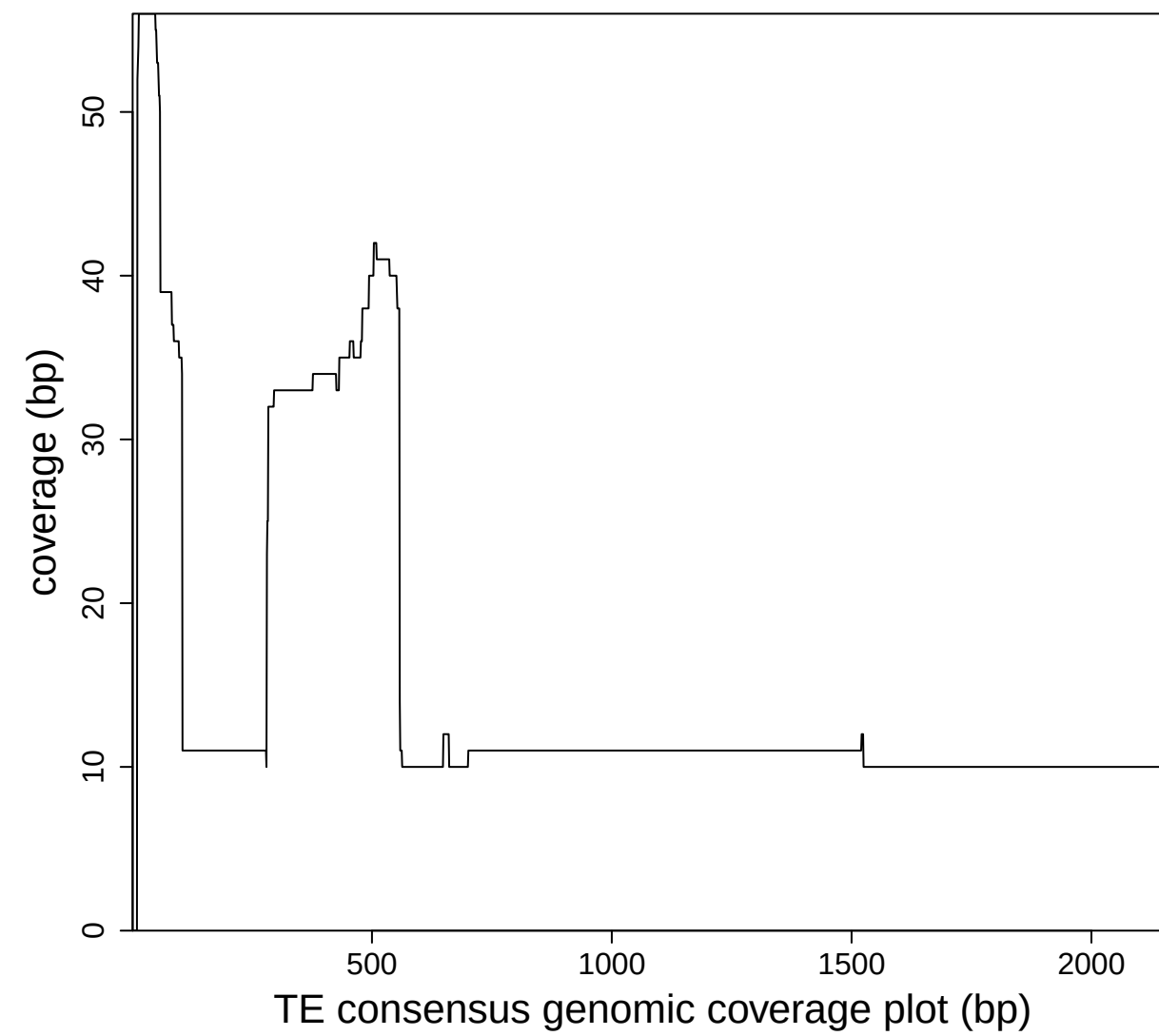
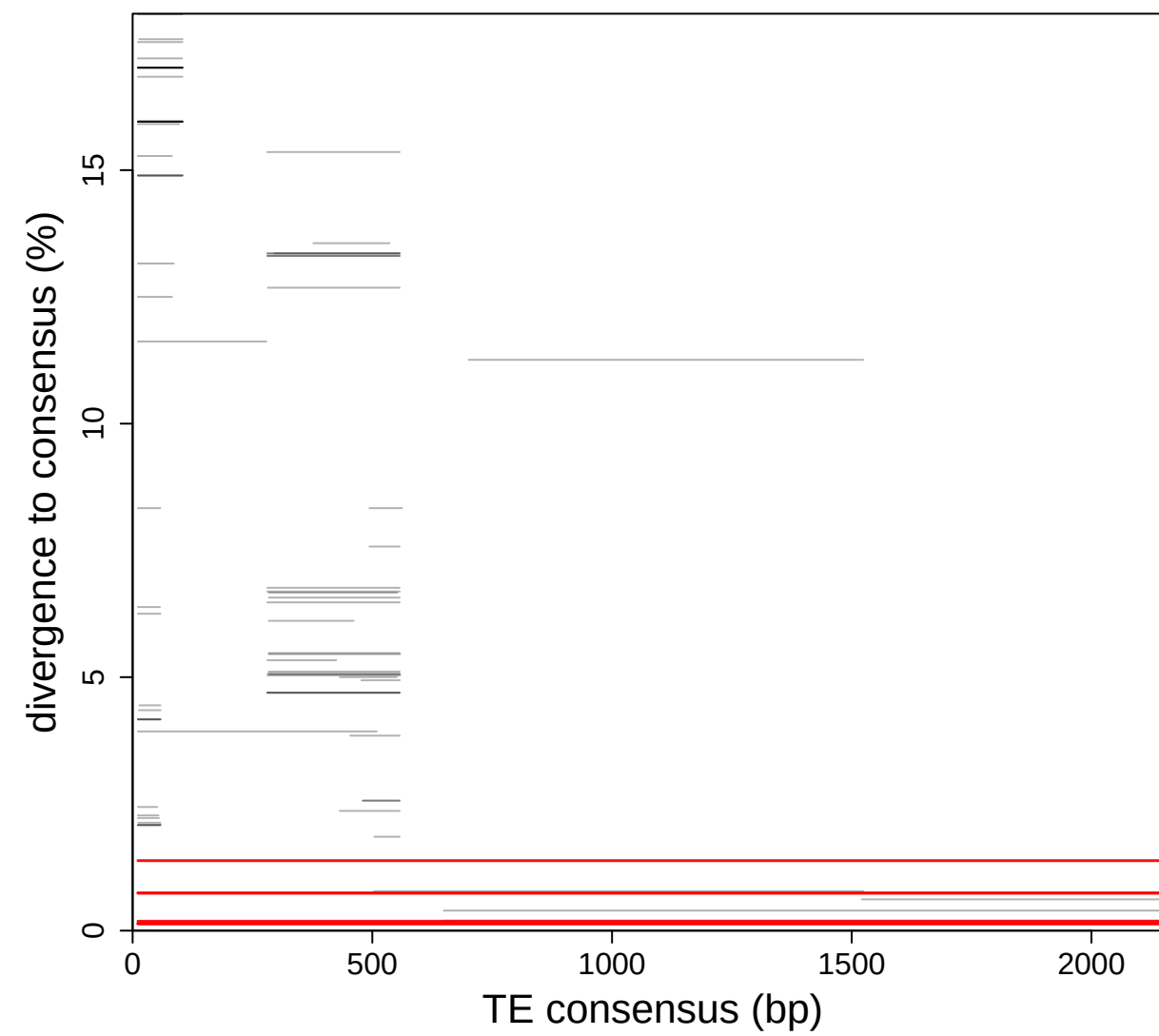
Start crop Point

End crop Point



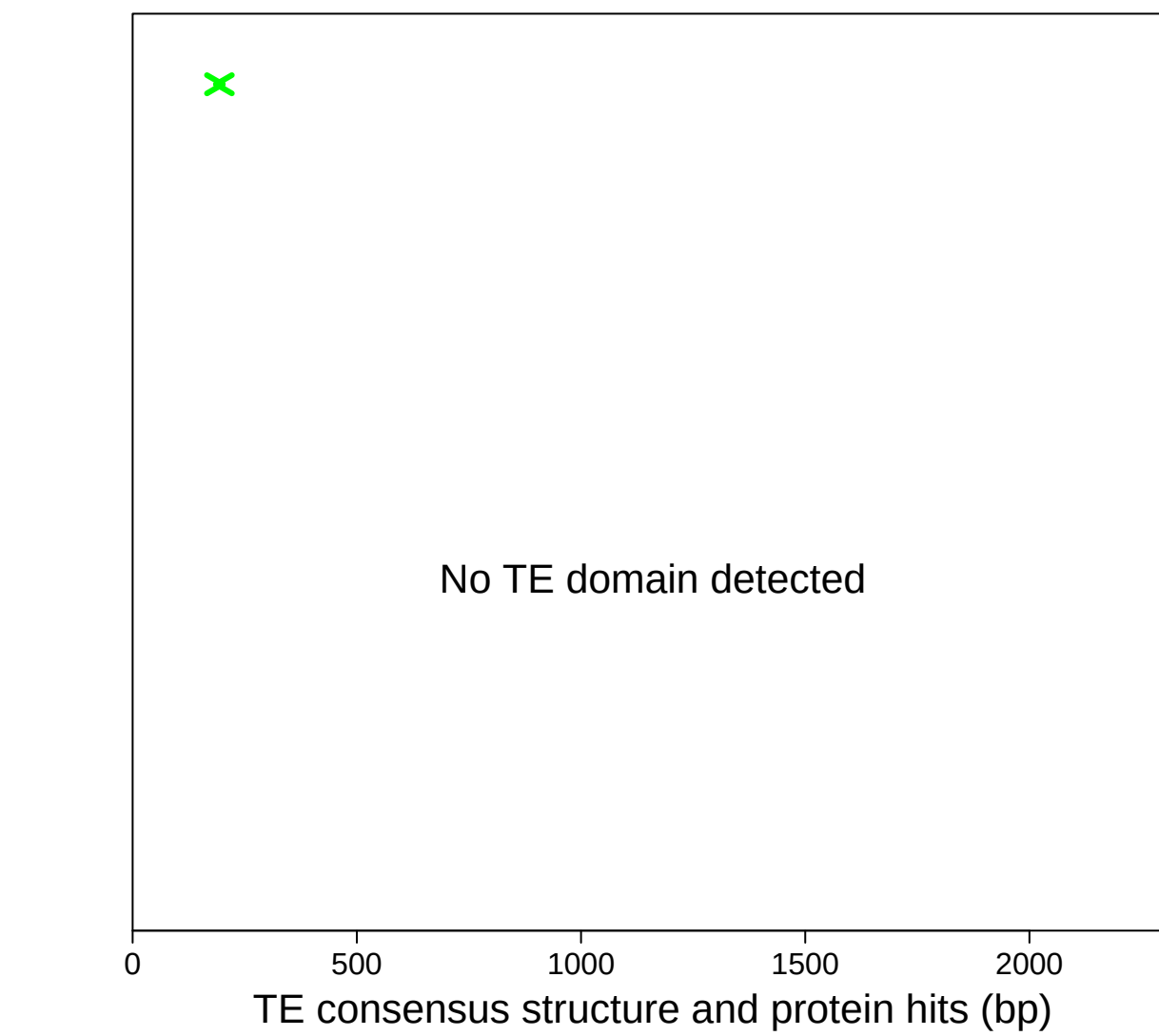
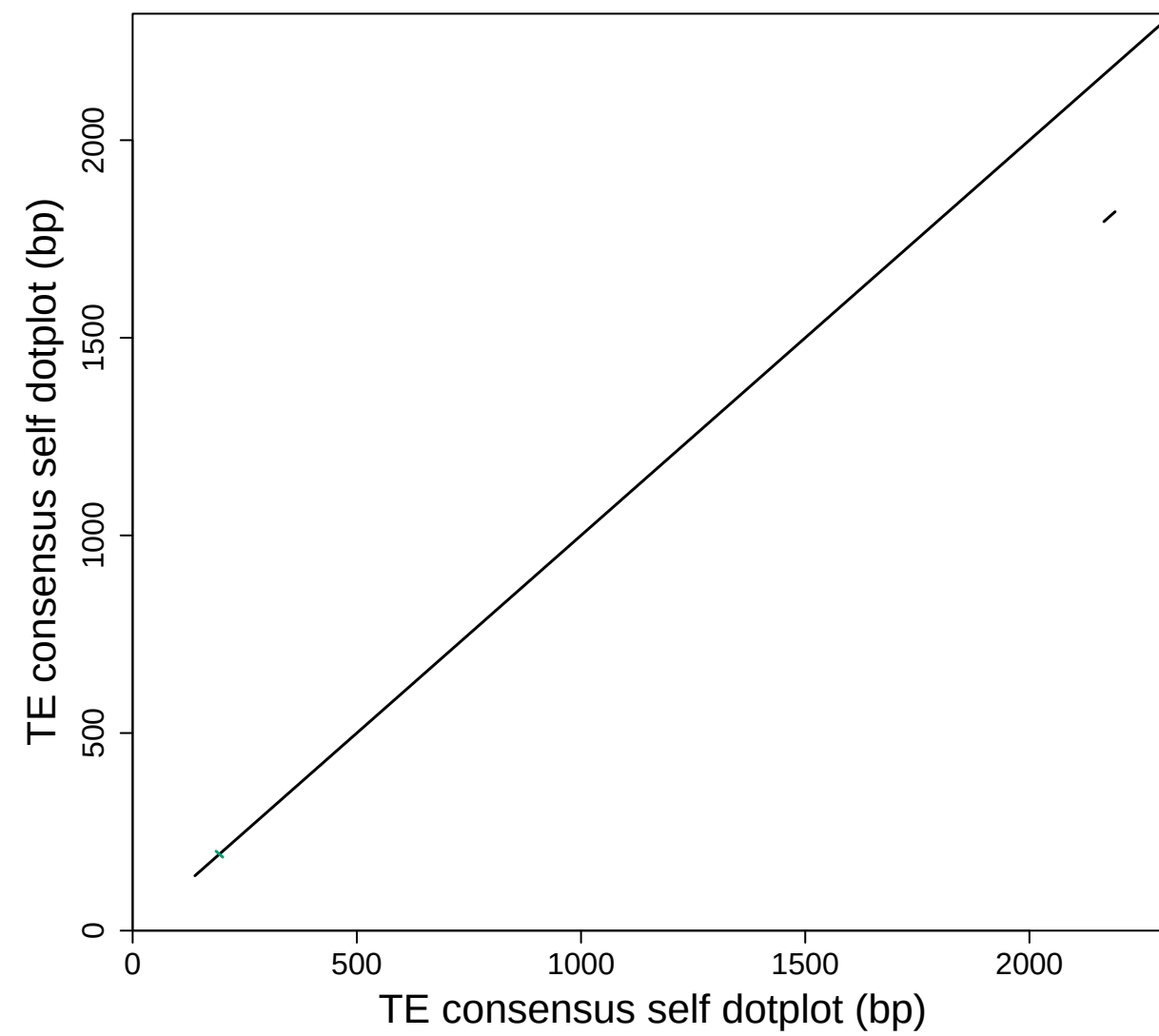
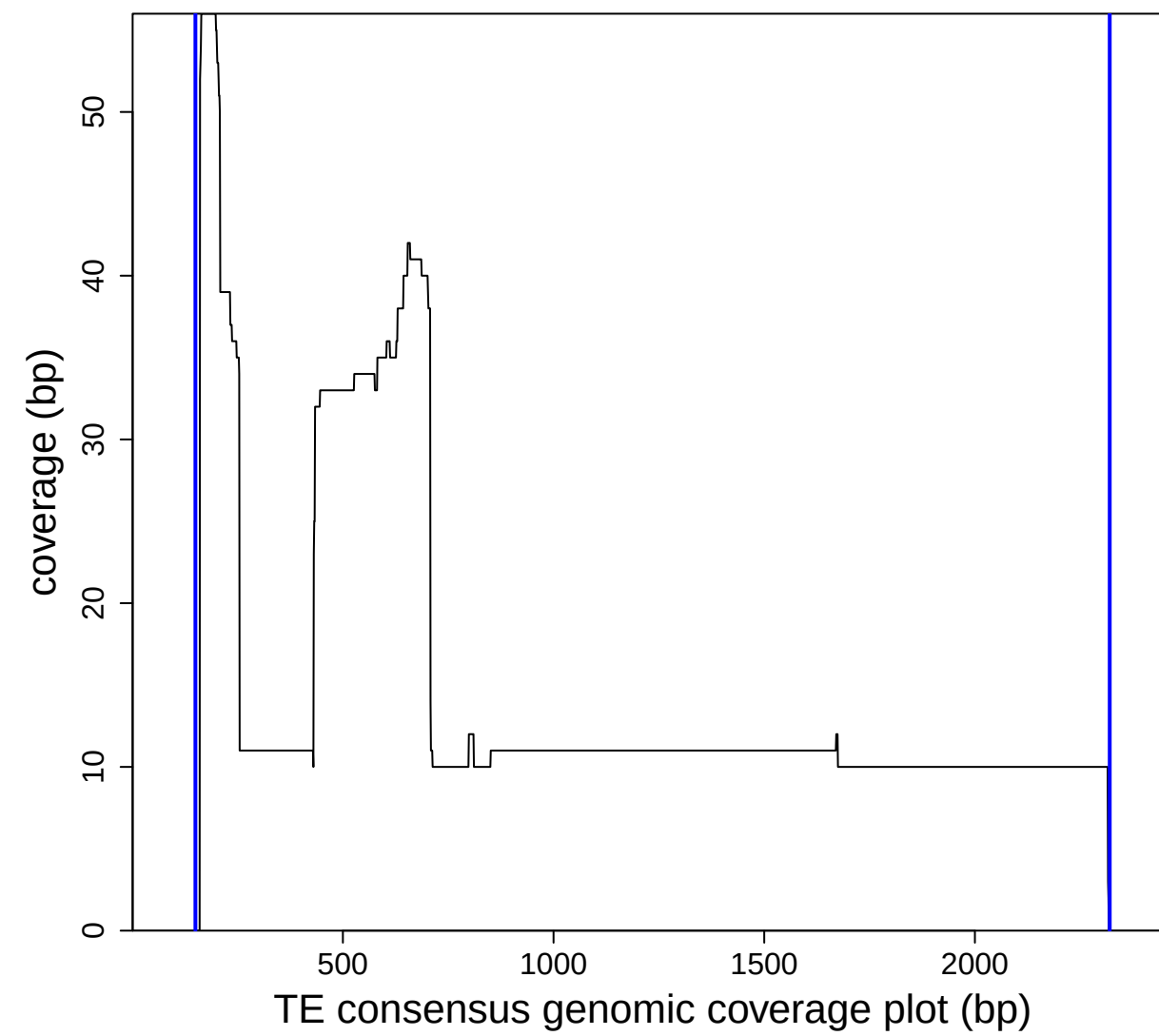
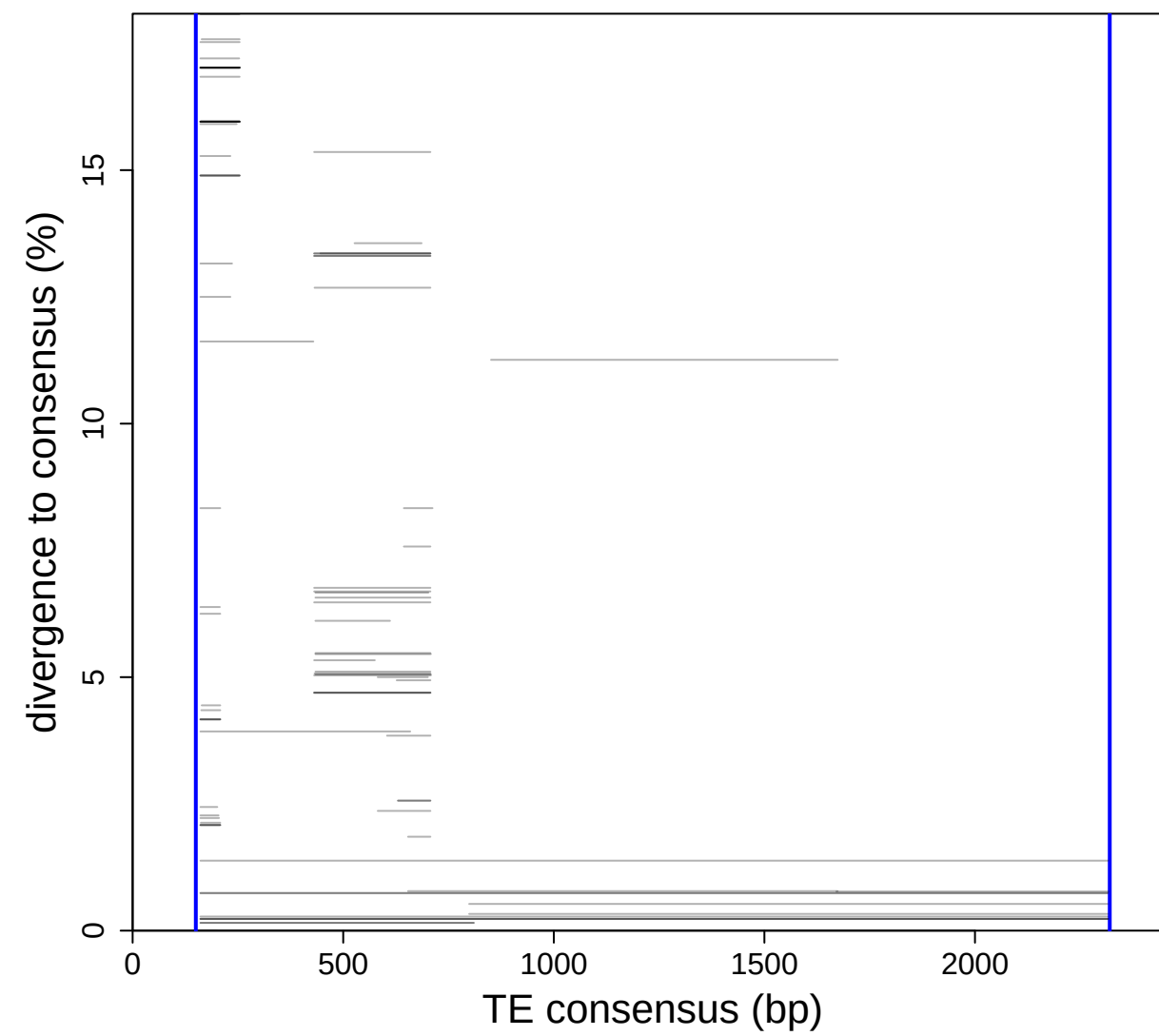
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_al
size: 2170bp; fragments: 94; full length: 7 (>=1953bp)

After TEtrimmer 2170 bp



bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 2470bp; fragments: 94; full length: 0 (>=2223bp)

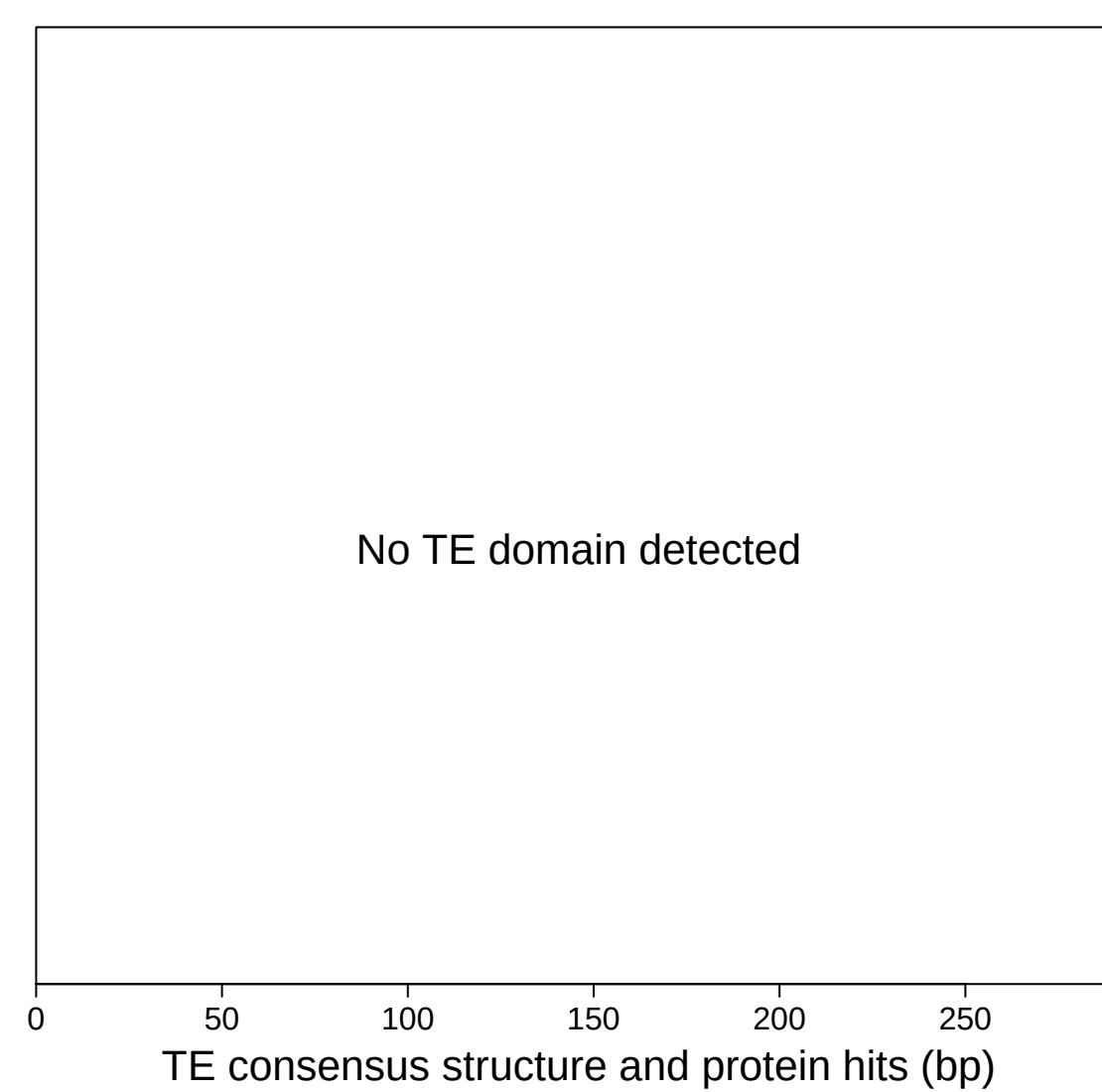
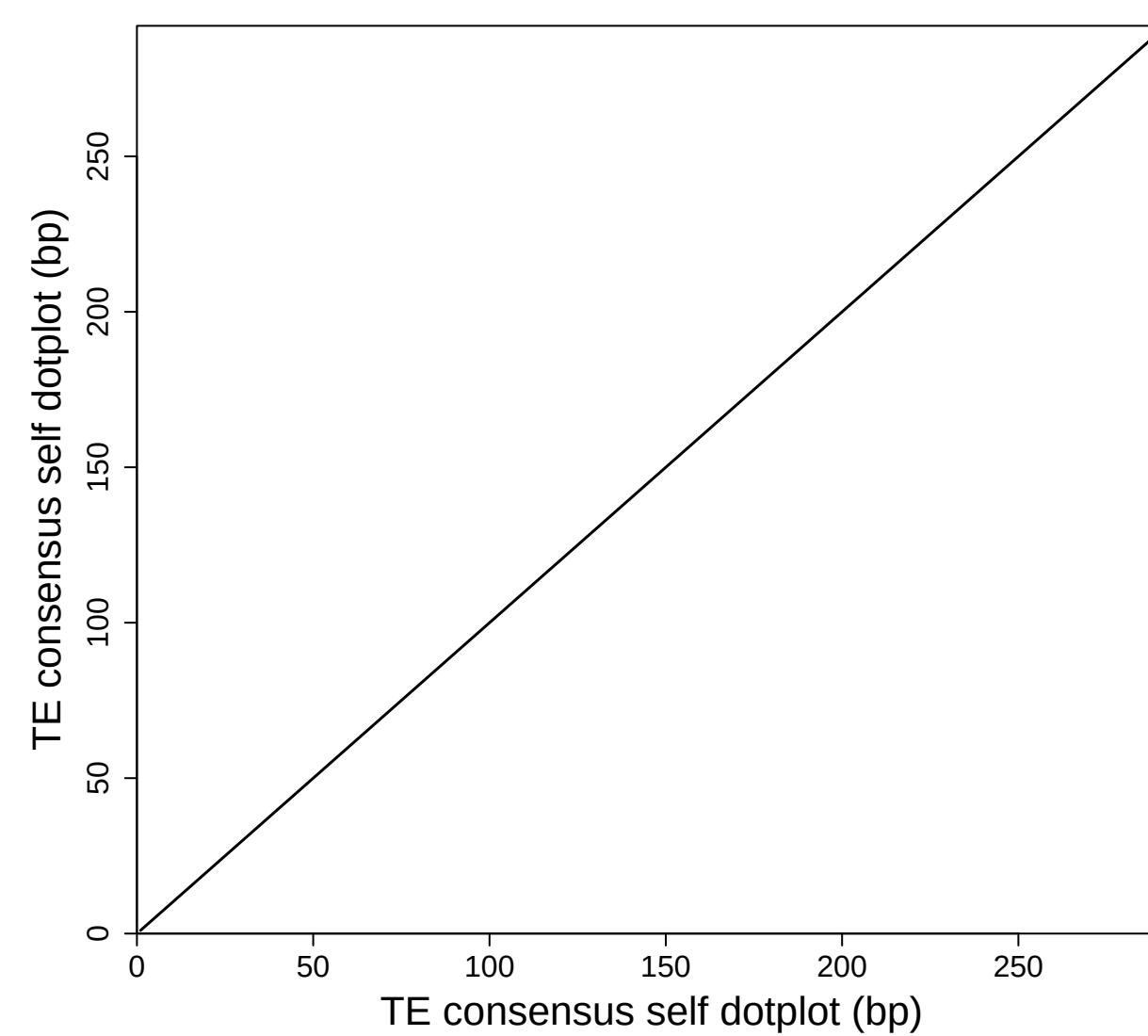
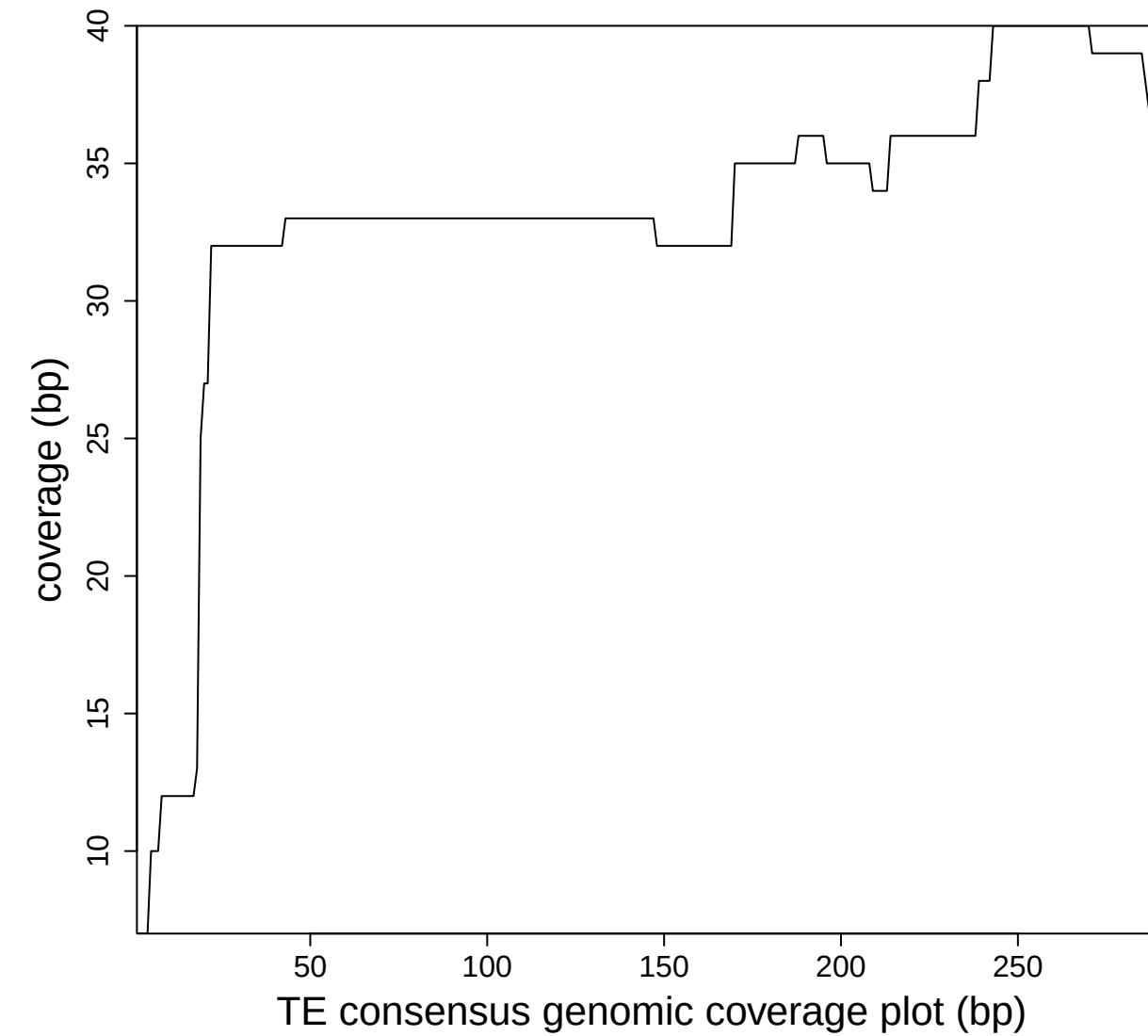
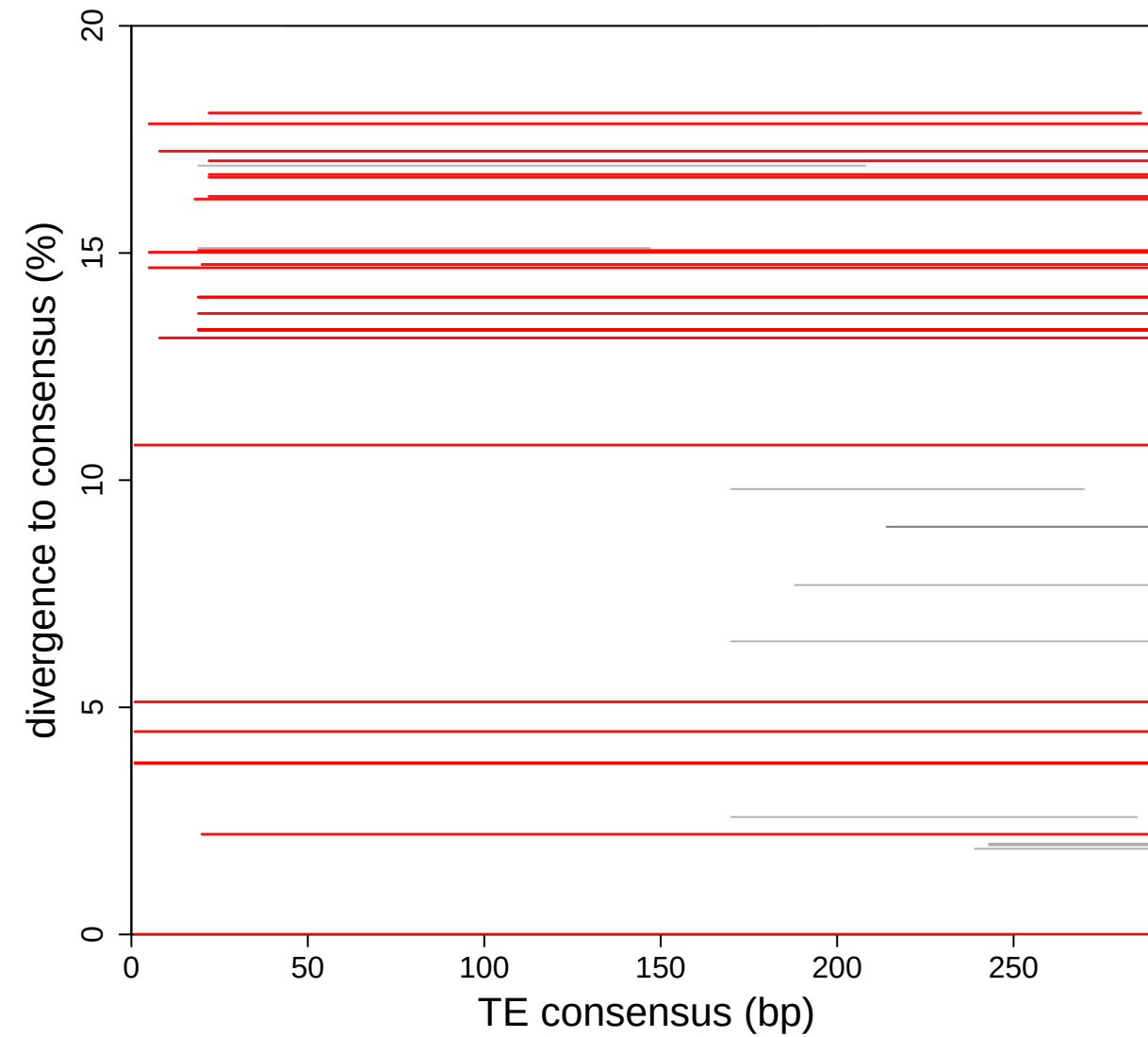
After TEtrimmer Extended plot Blue lines are boundaries



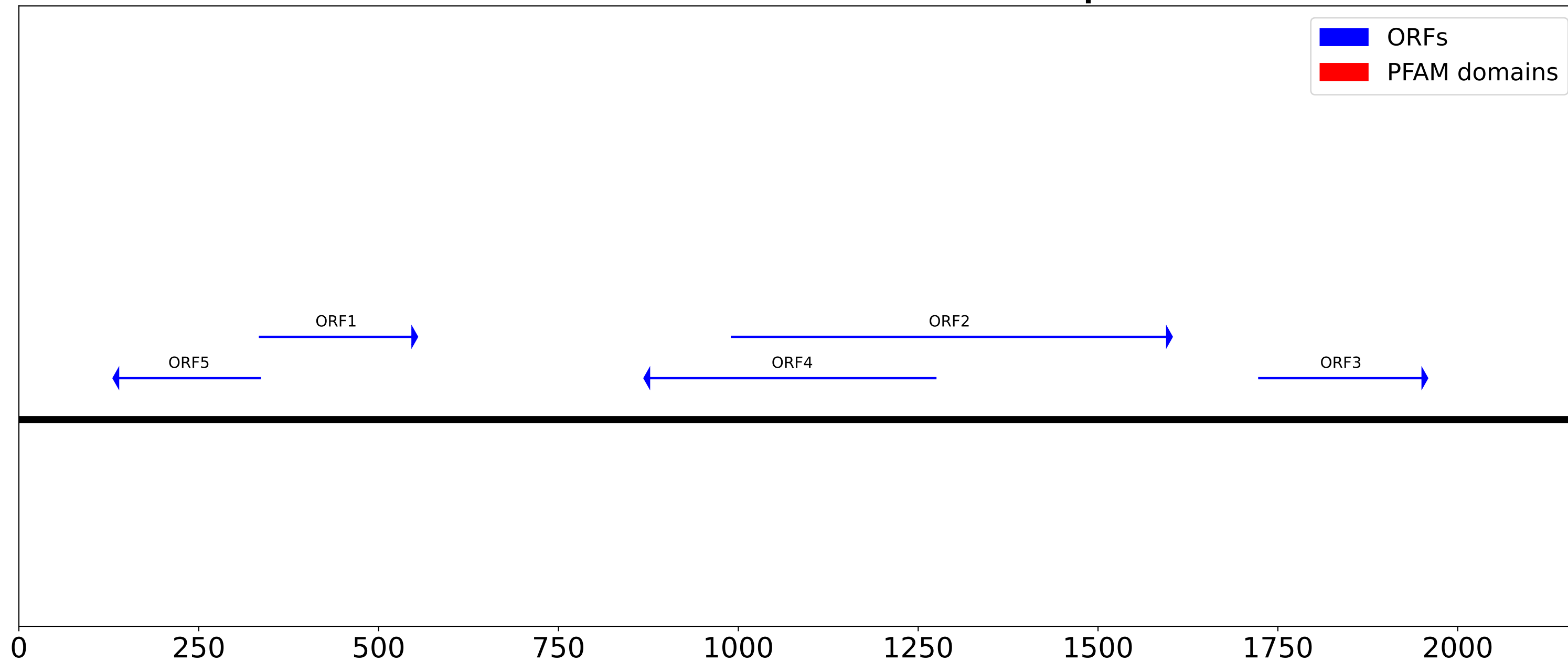
Before TEtrimmer 292 bp

TE: ltr_1_family_101

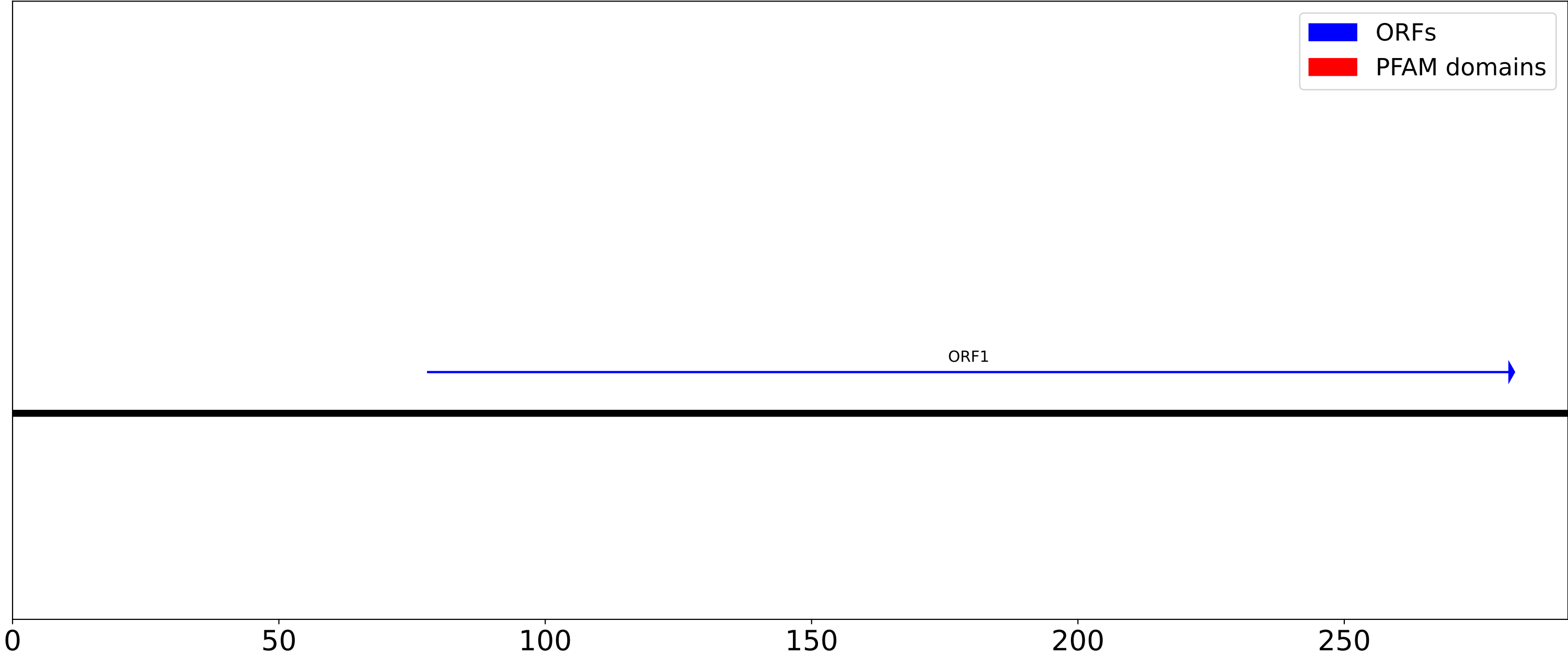
size: 292bp; fragments: 43; full length: 30 (≥ 262.8 bp)



After TEtrimmer ORF and PFAM domain plot



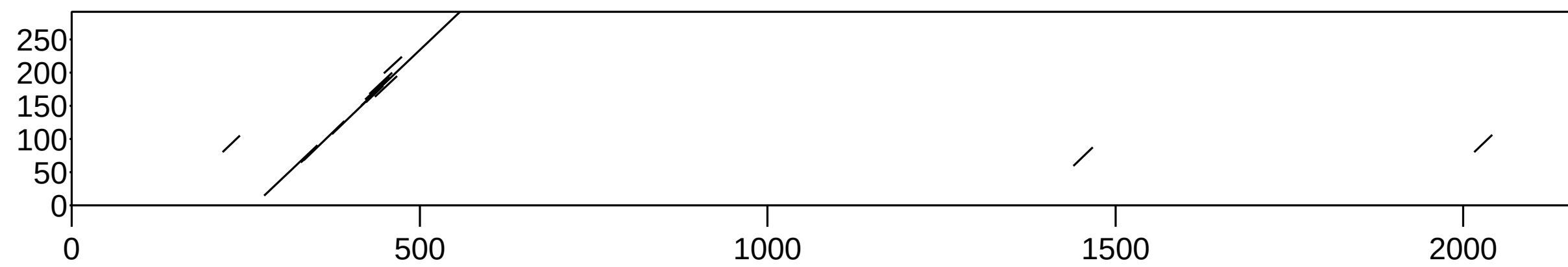
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

sensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)