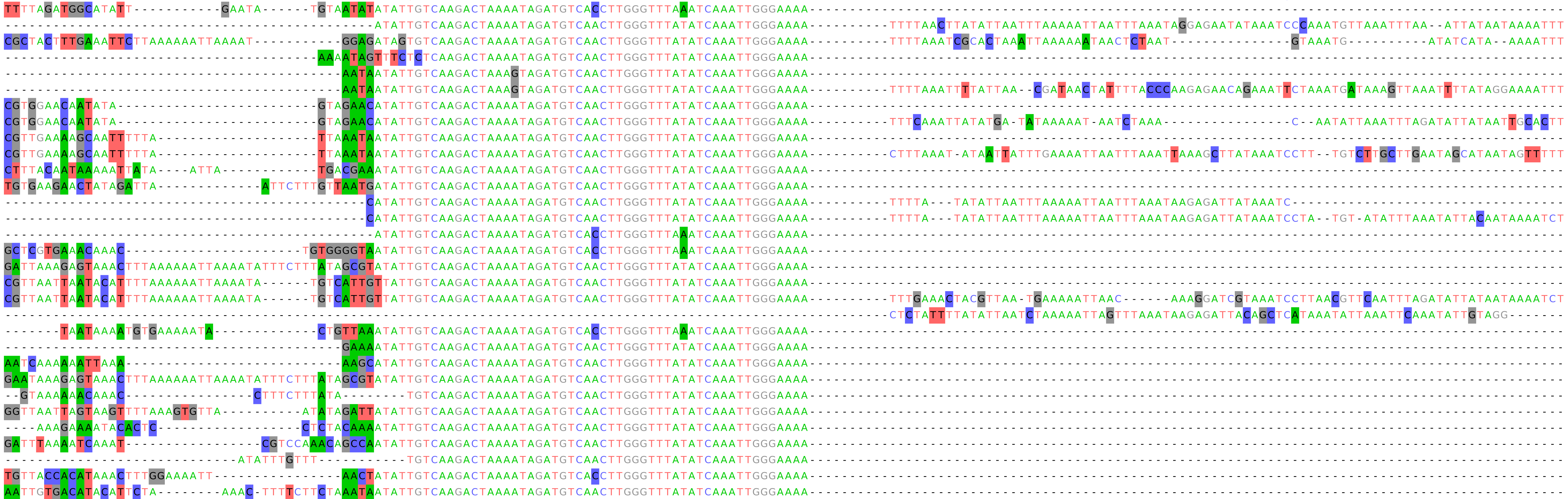


MSA length = 6768

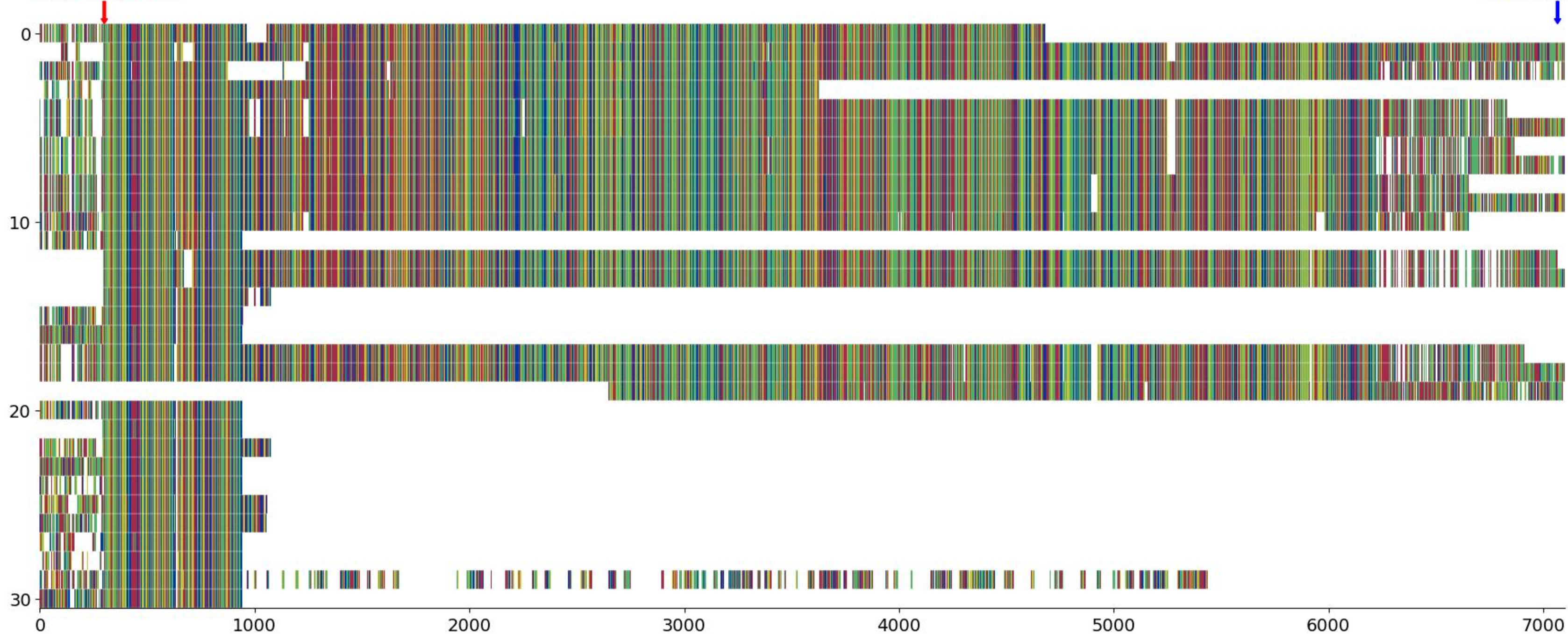
Start crop Point

End crop Point



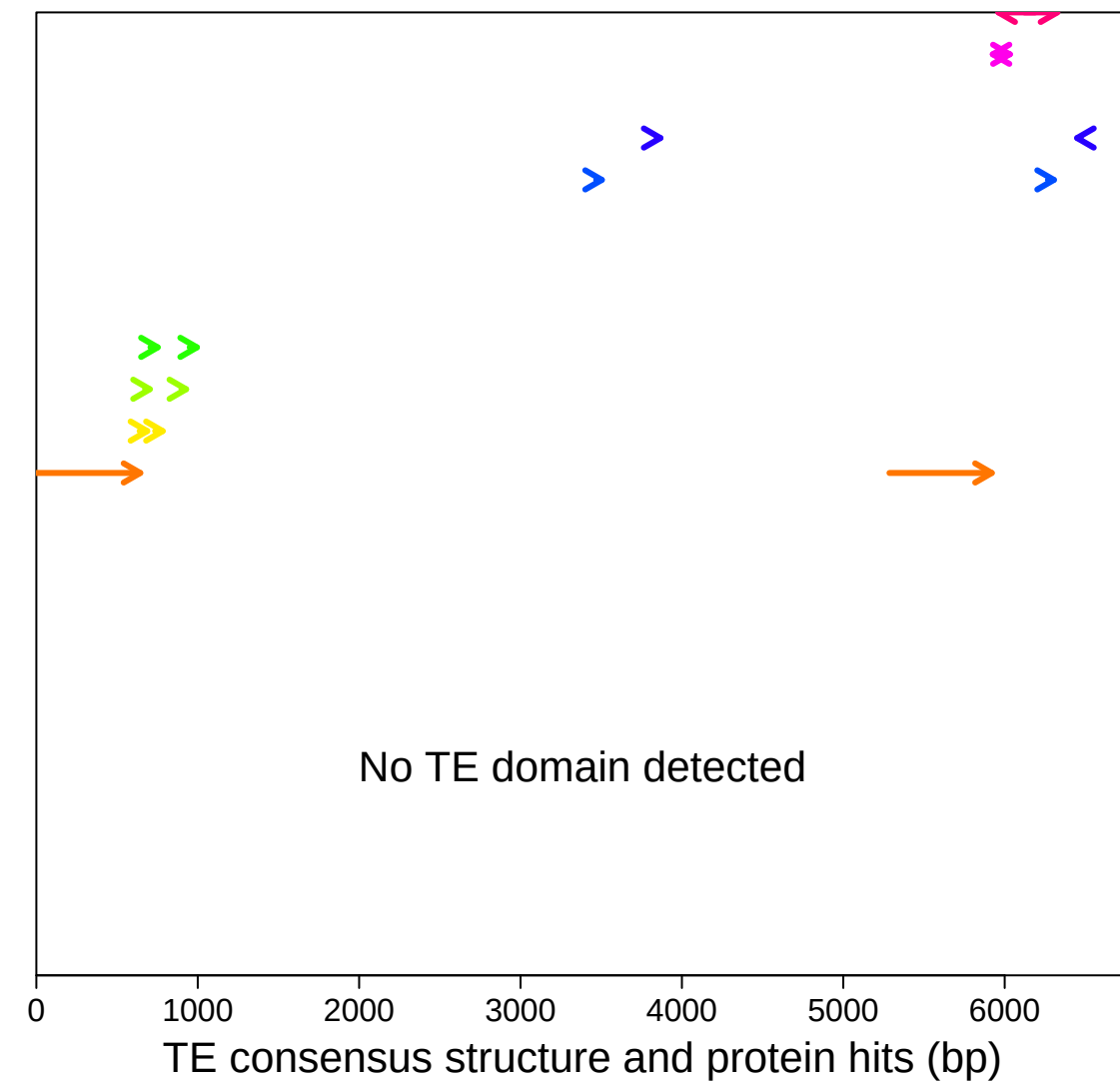
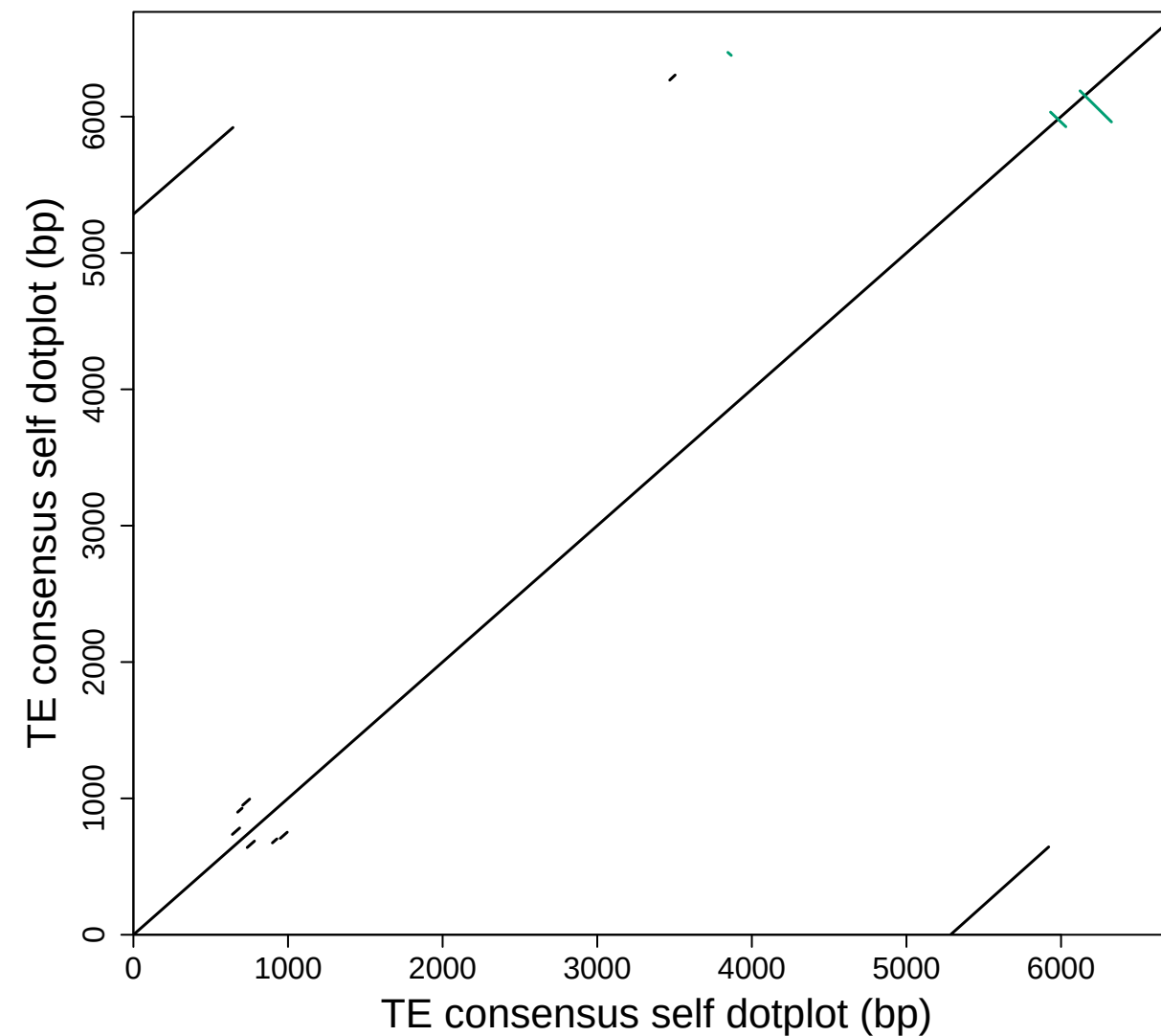
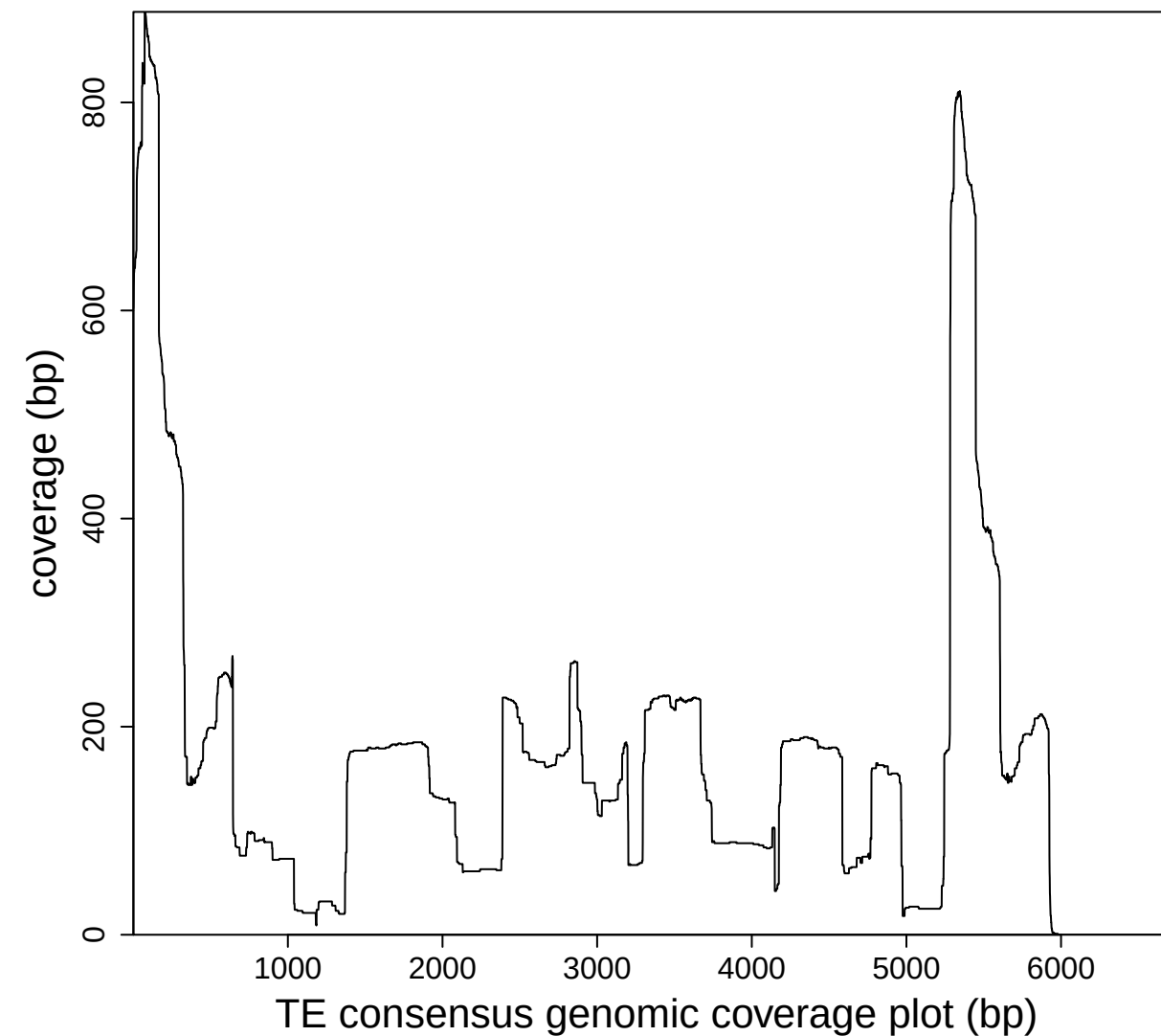
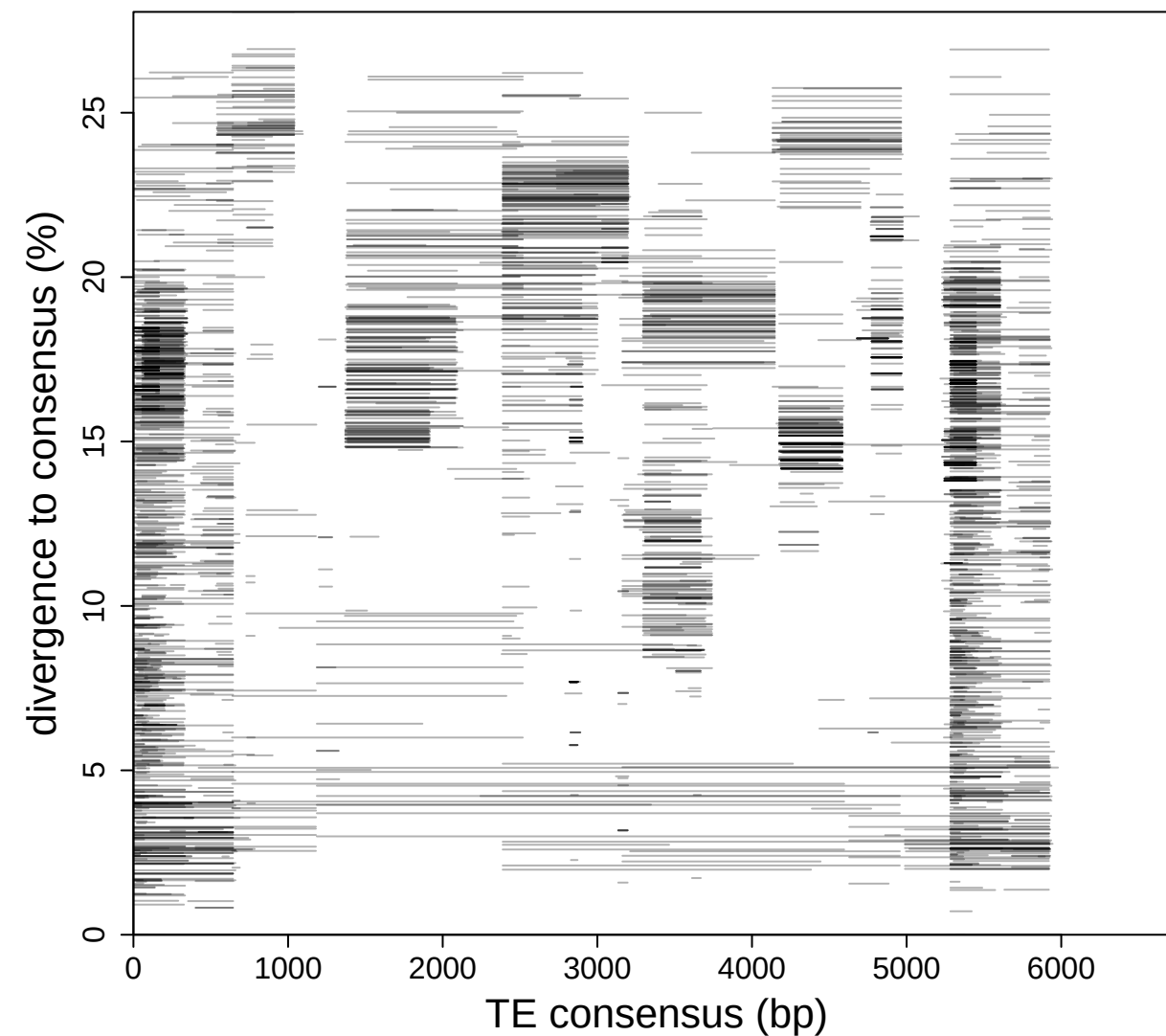
Start crop Point

End crop Point



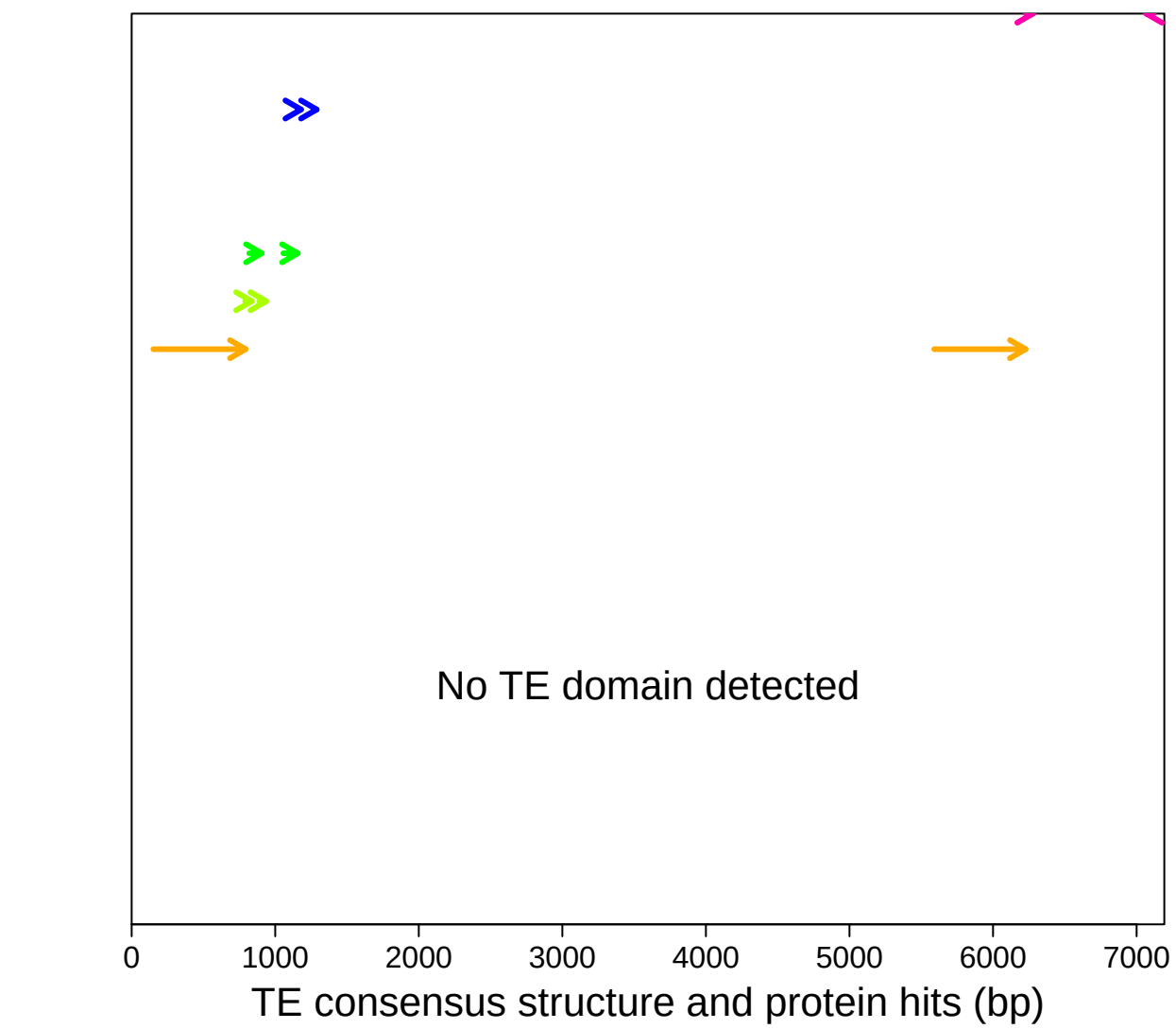
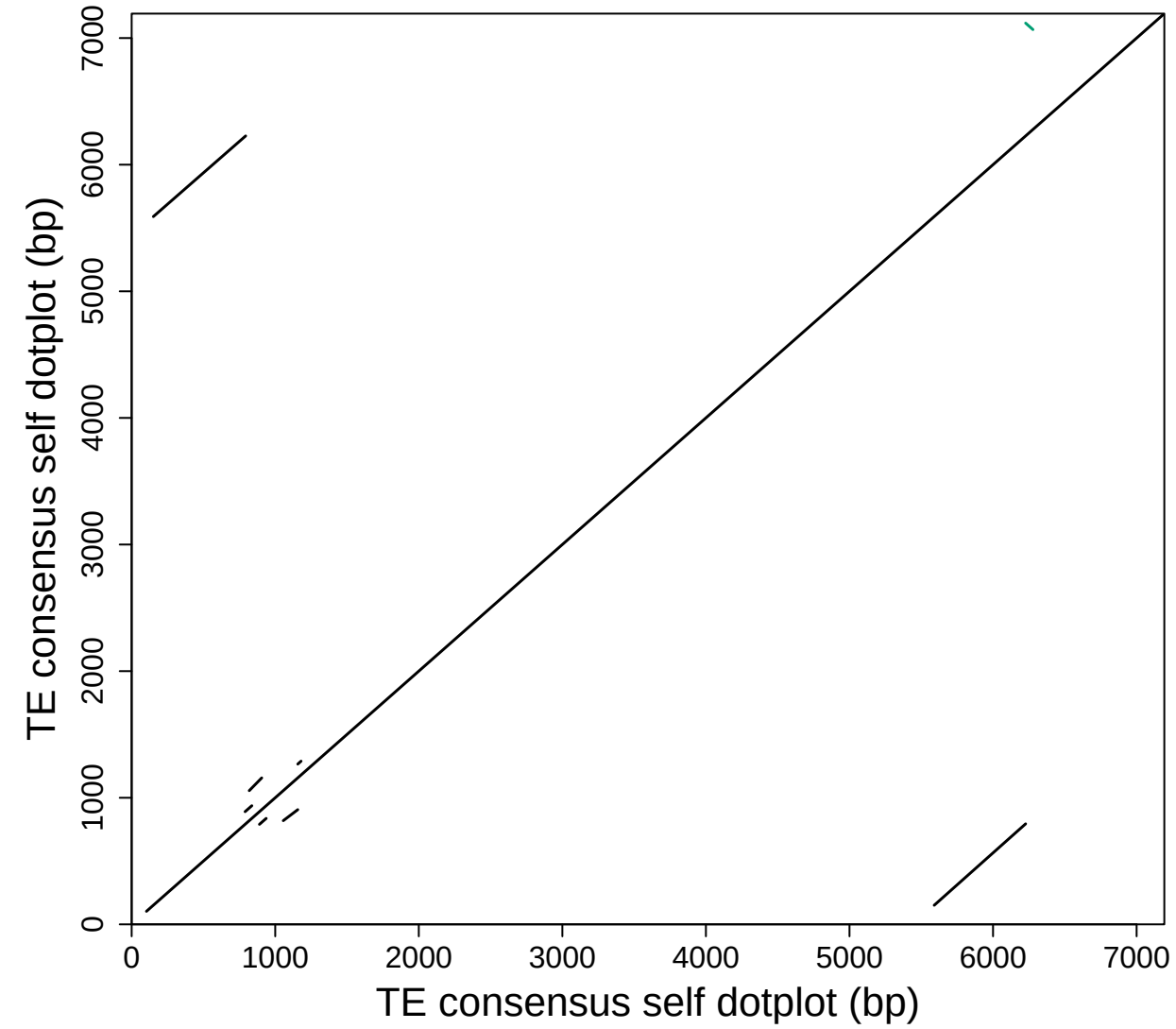
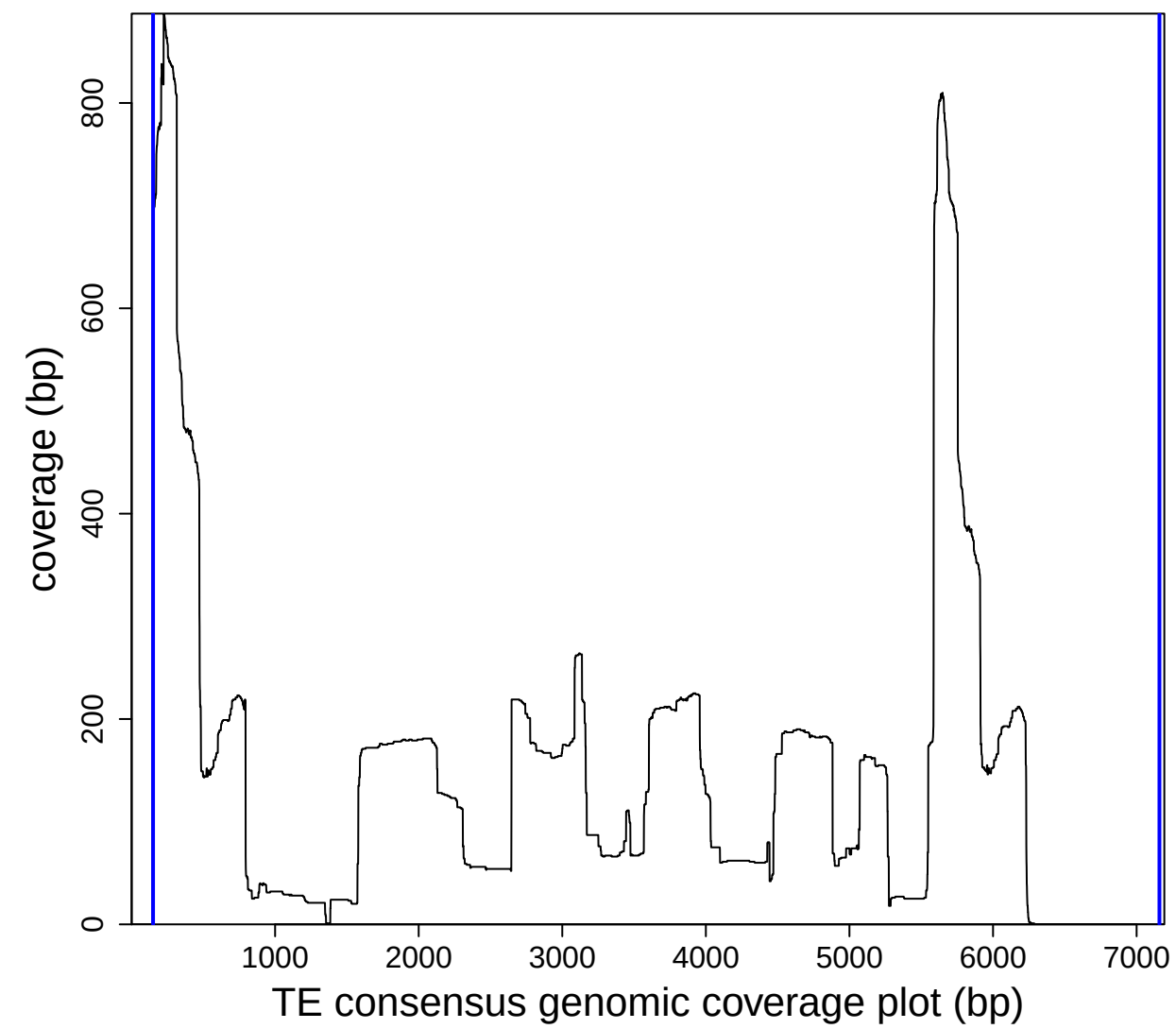
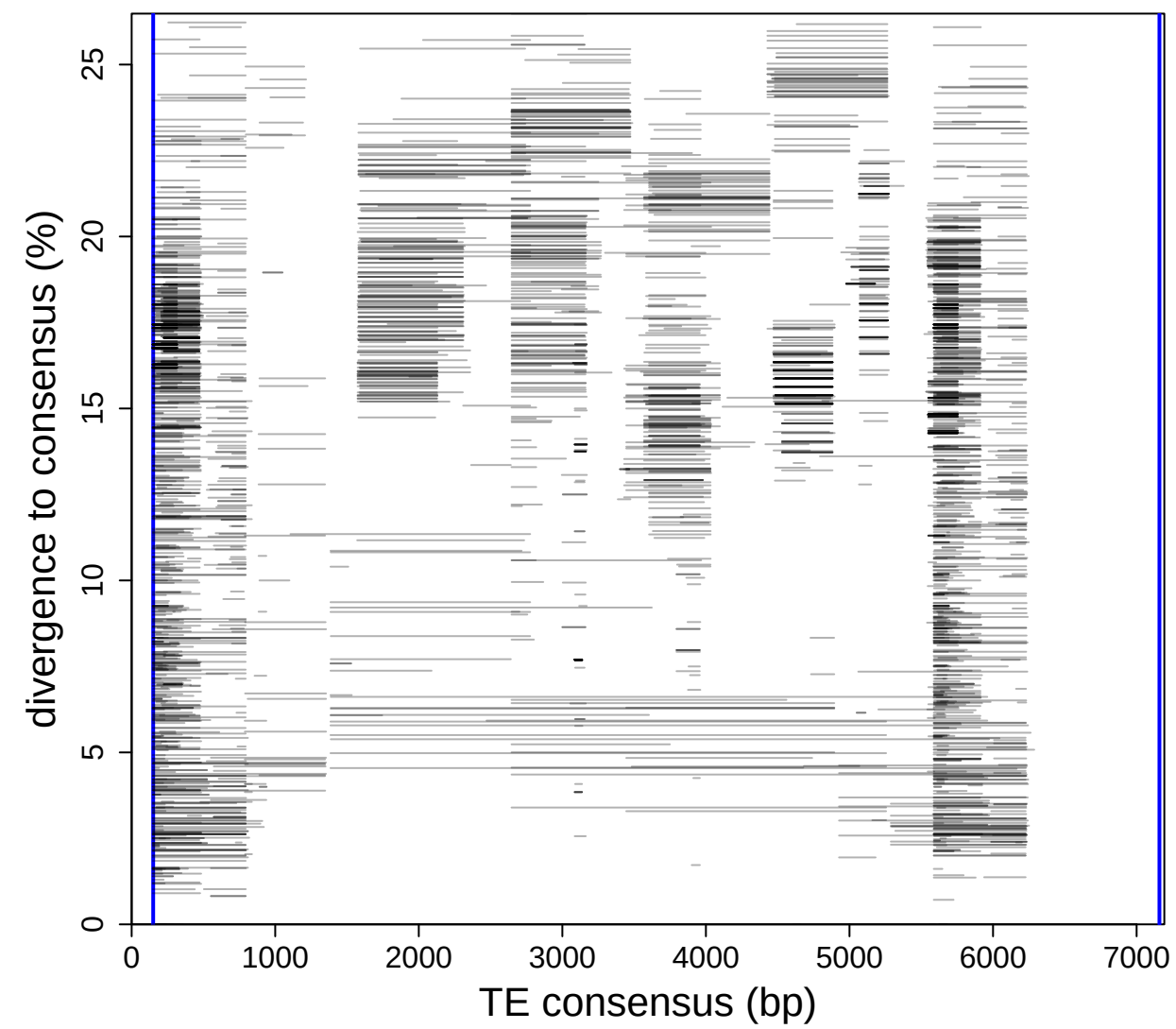
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa al
size: 6768bp; fragments: 3087; full length: 0 (>=6091.2bp)

After TEtrimmer 6768 bp



bed_g_1.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bclr
size: 7193bp; fragments: 2977; full length: 0 (>=6473.7bp)

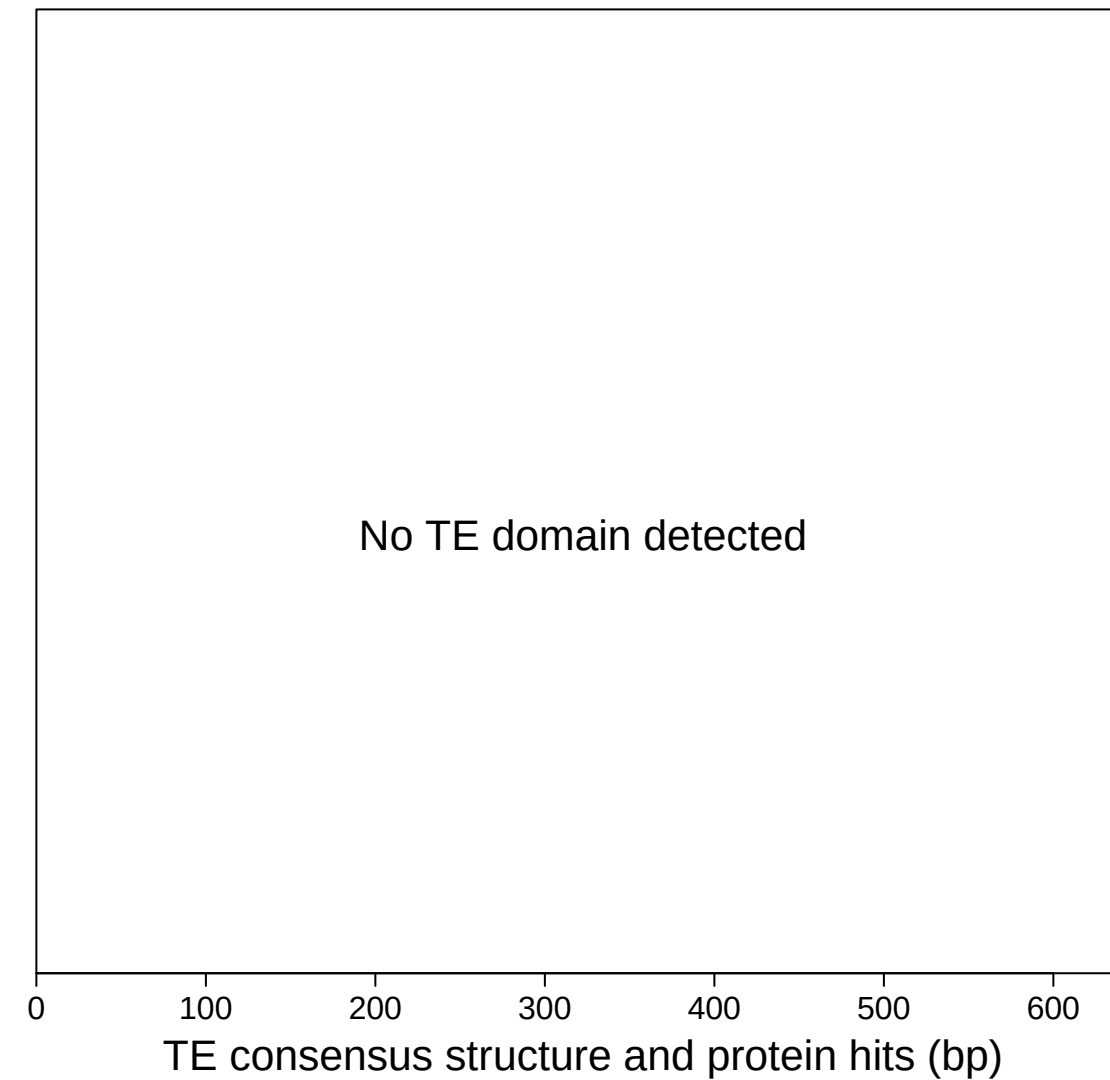
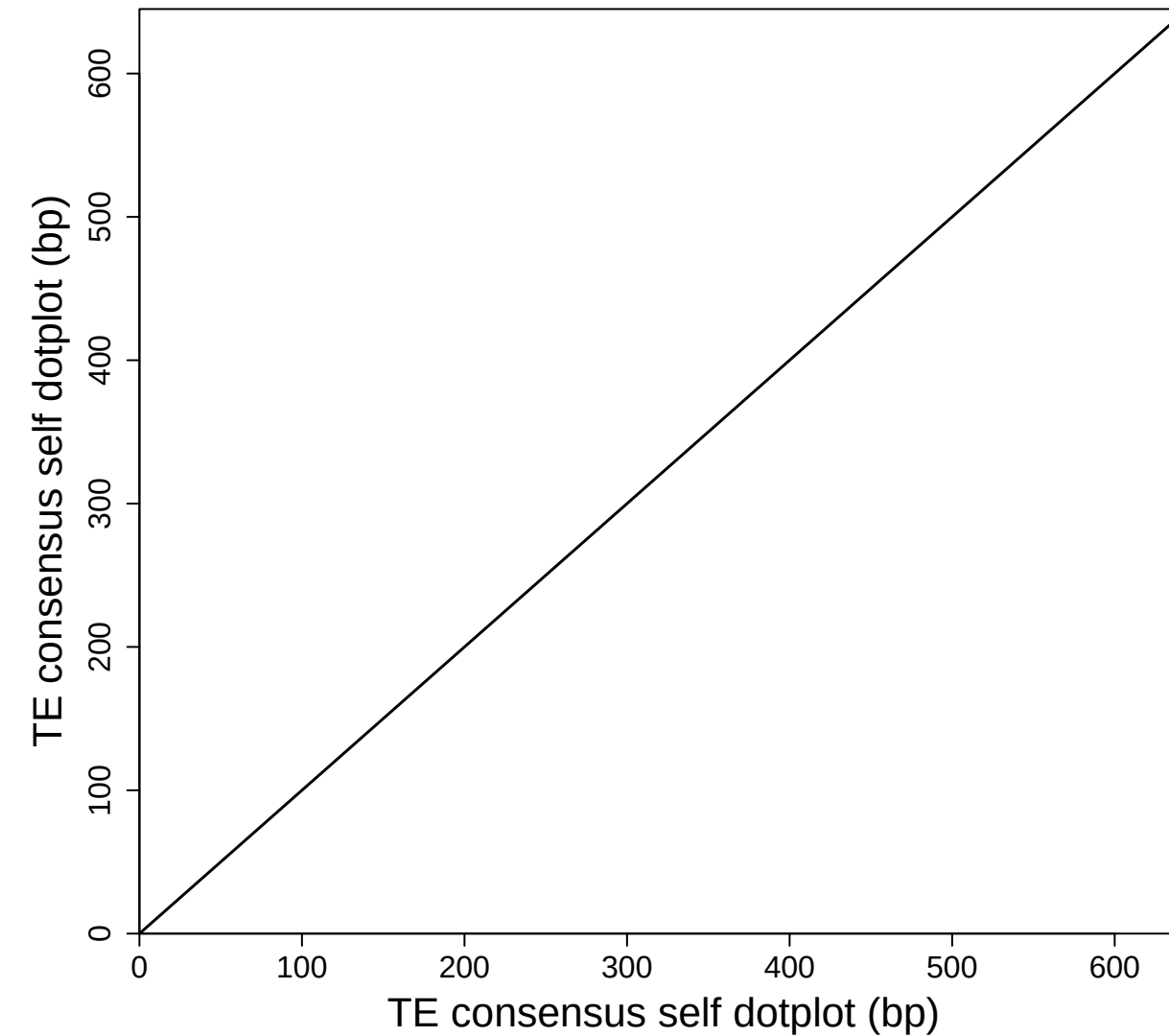
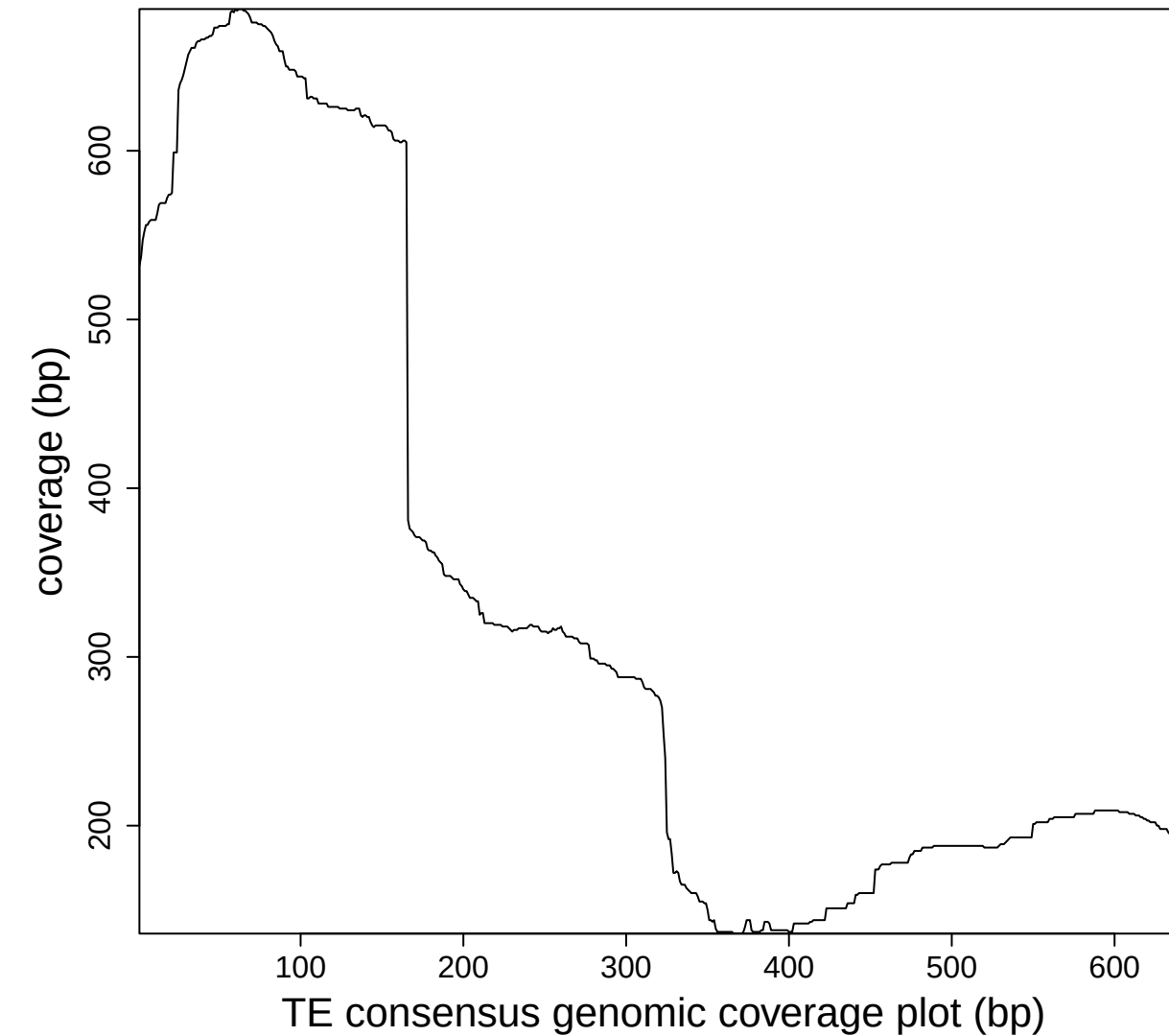
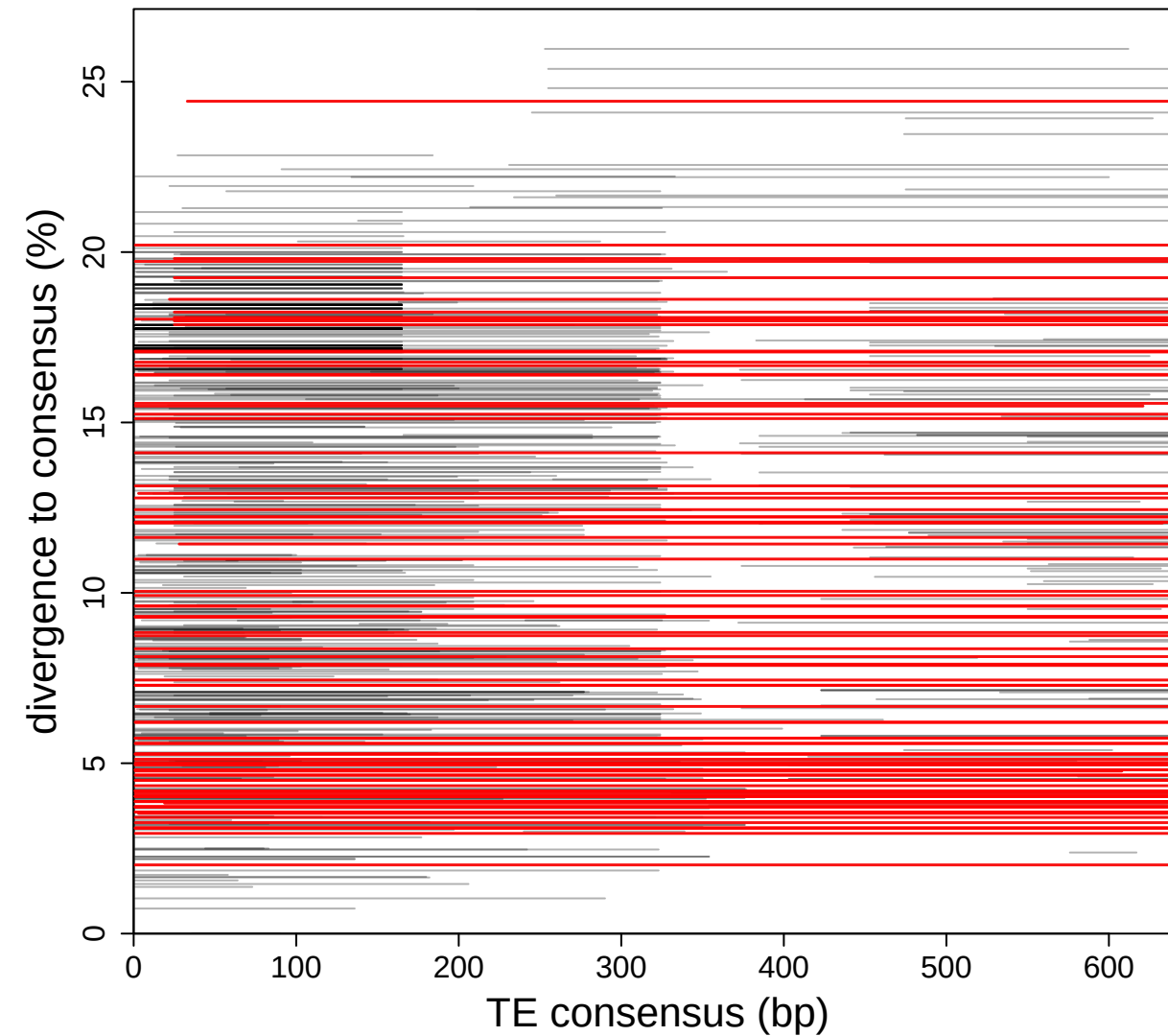
After TEtrimmer Extended plot Blue lines are boundaries



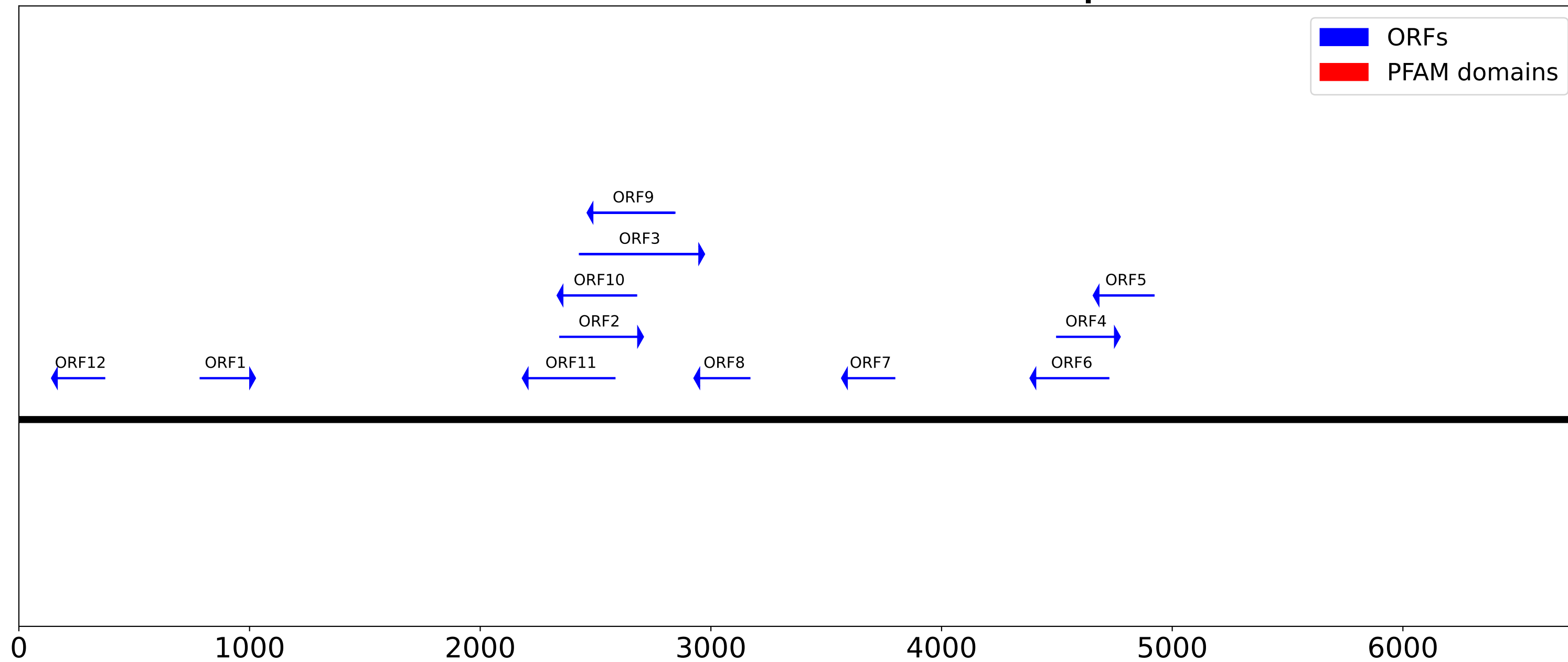
Before TEtrimmer 645 bp

TE: ltr_1_family_116

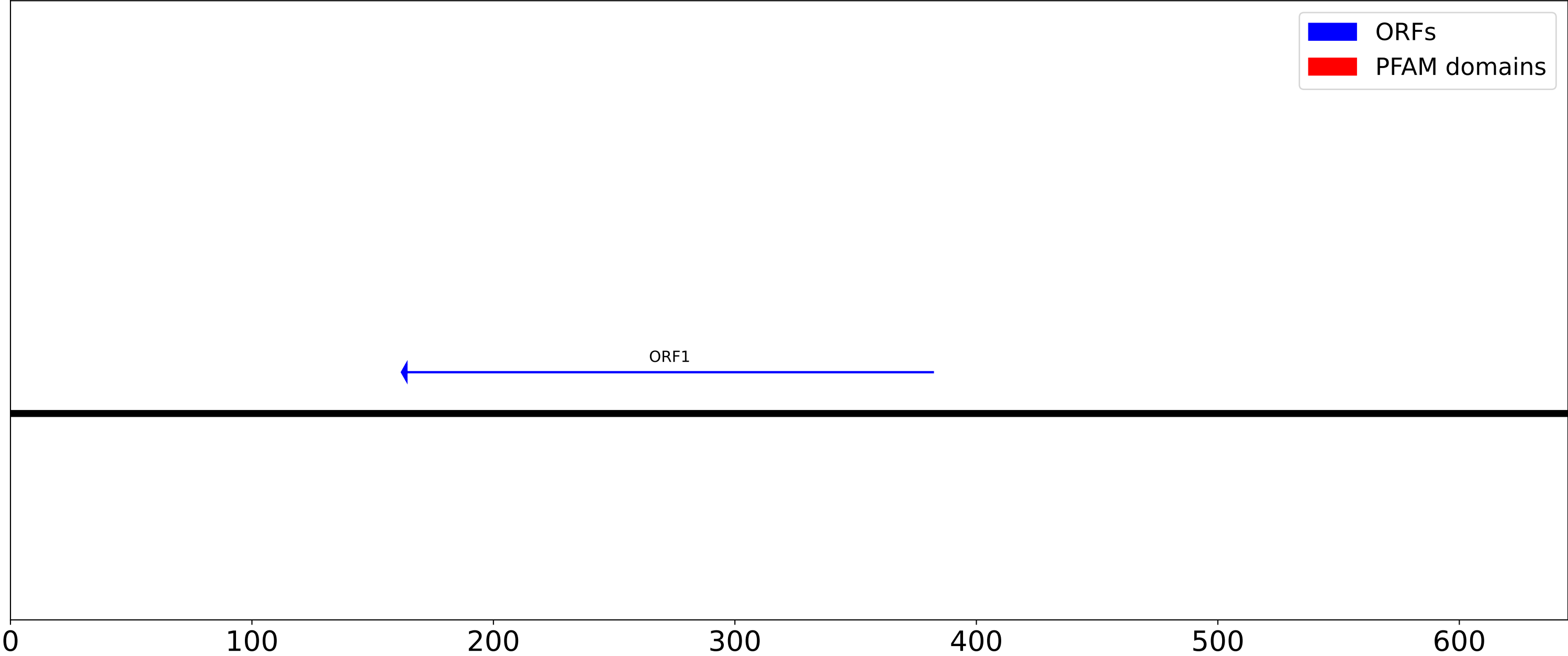
size: 645bp; fragments: 803; full length: 105 (≥ 580.5 bp)



After TEtrimmer ORF and PFAM domain plot



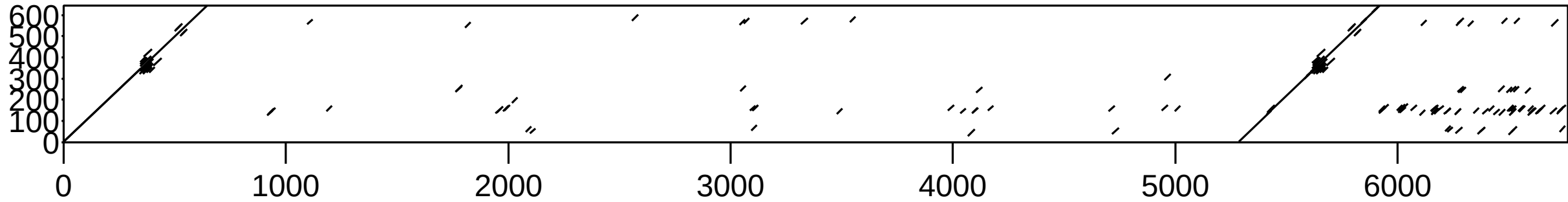
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)