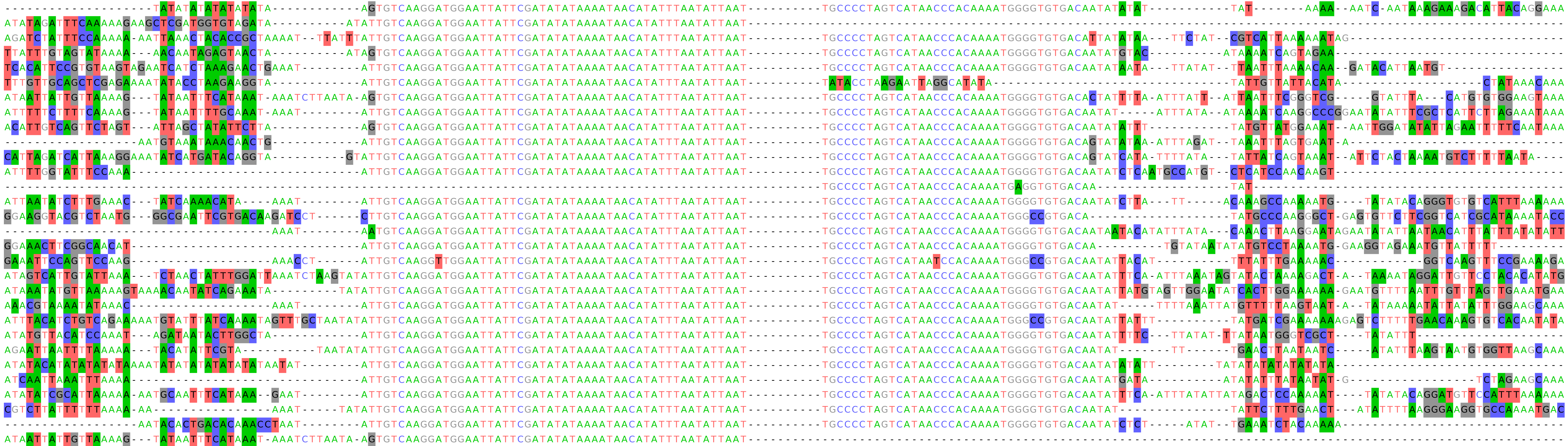


MSA length = 6040

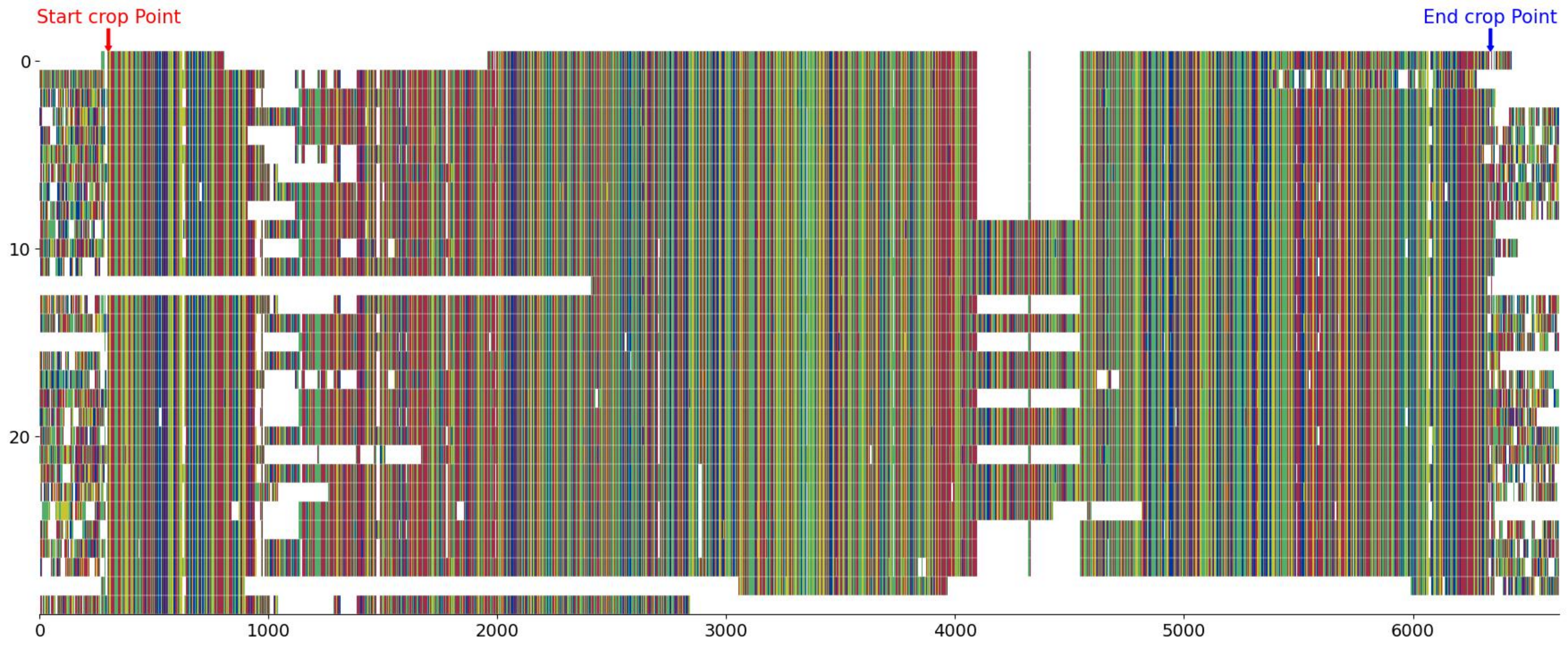
Start crop Point

End crop Point



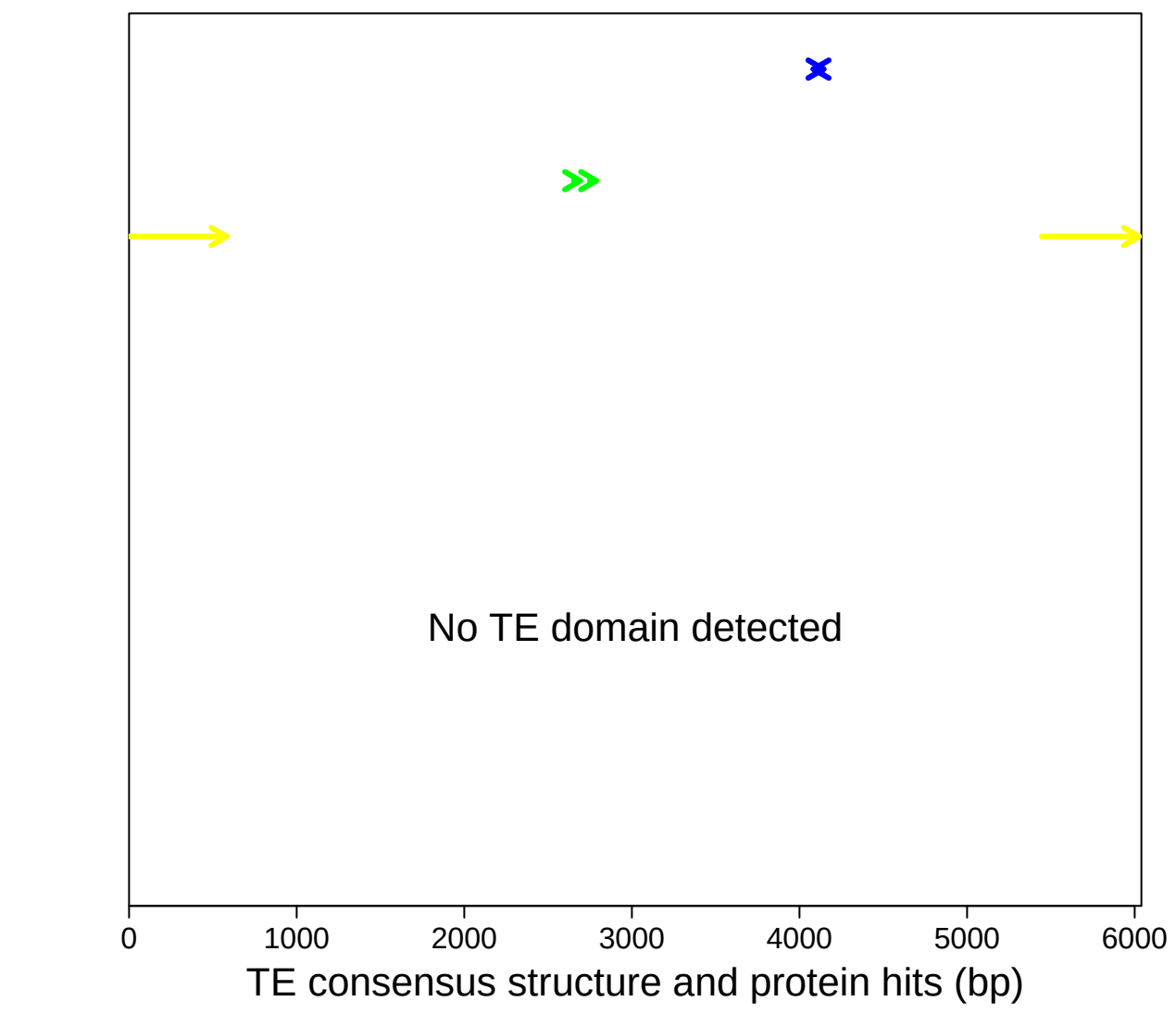
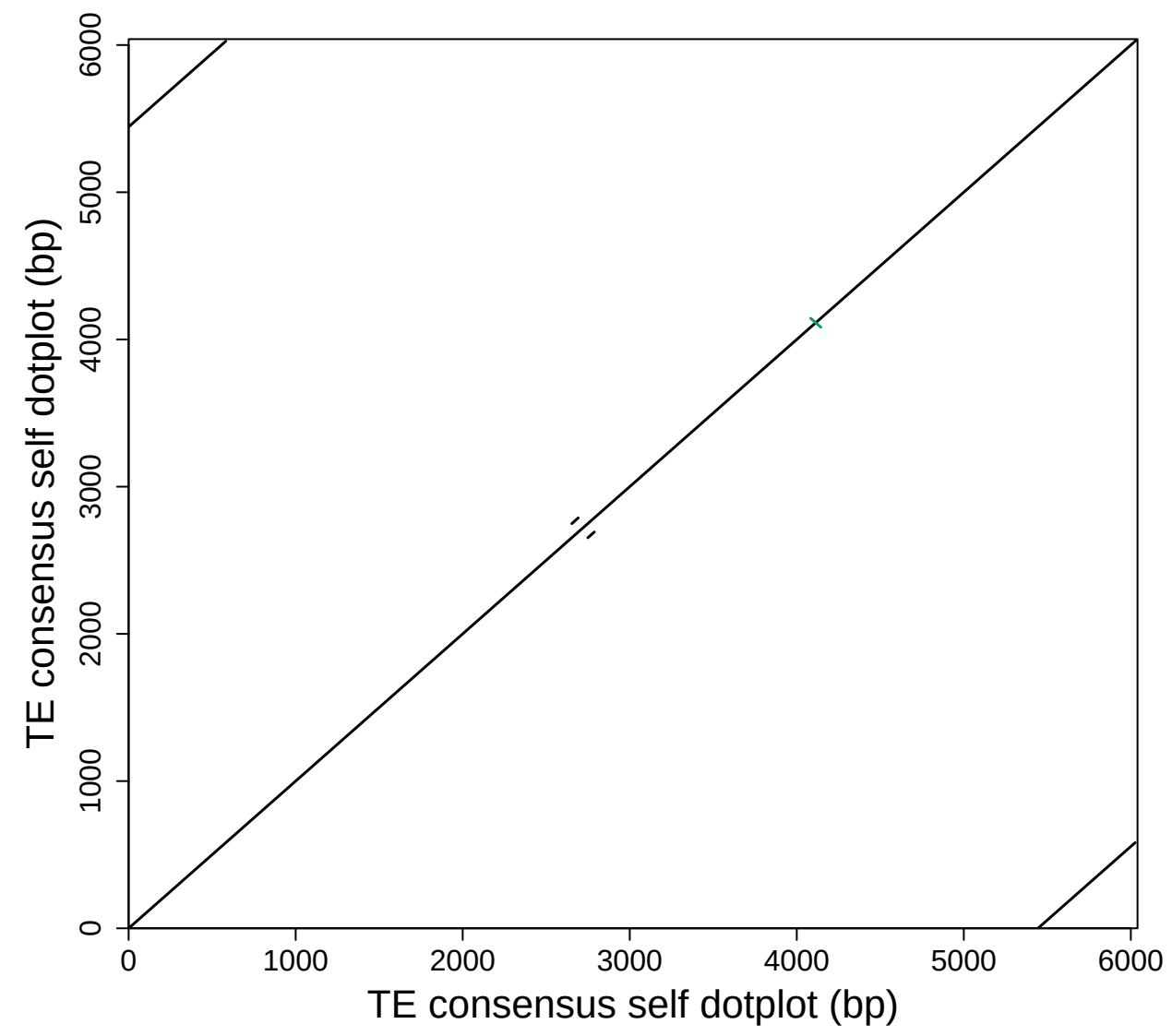
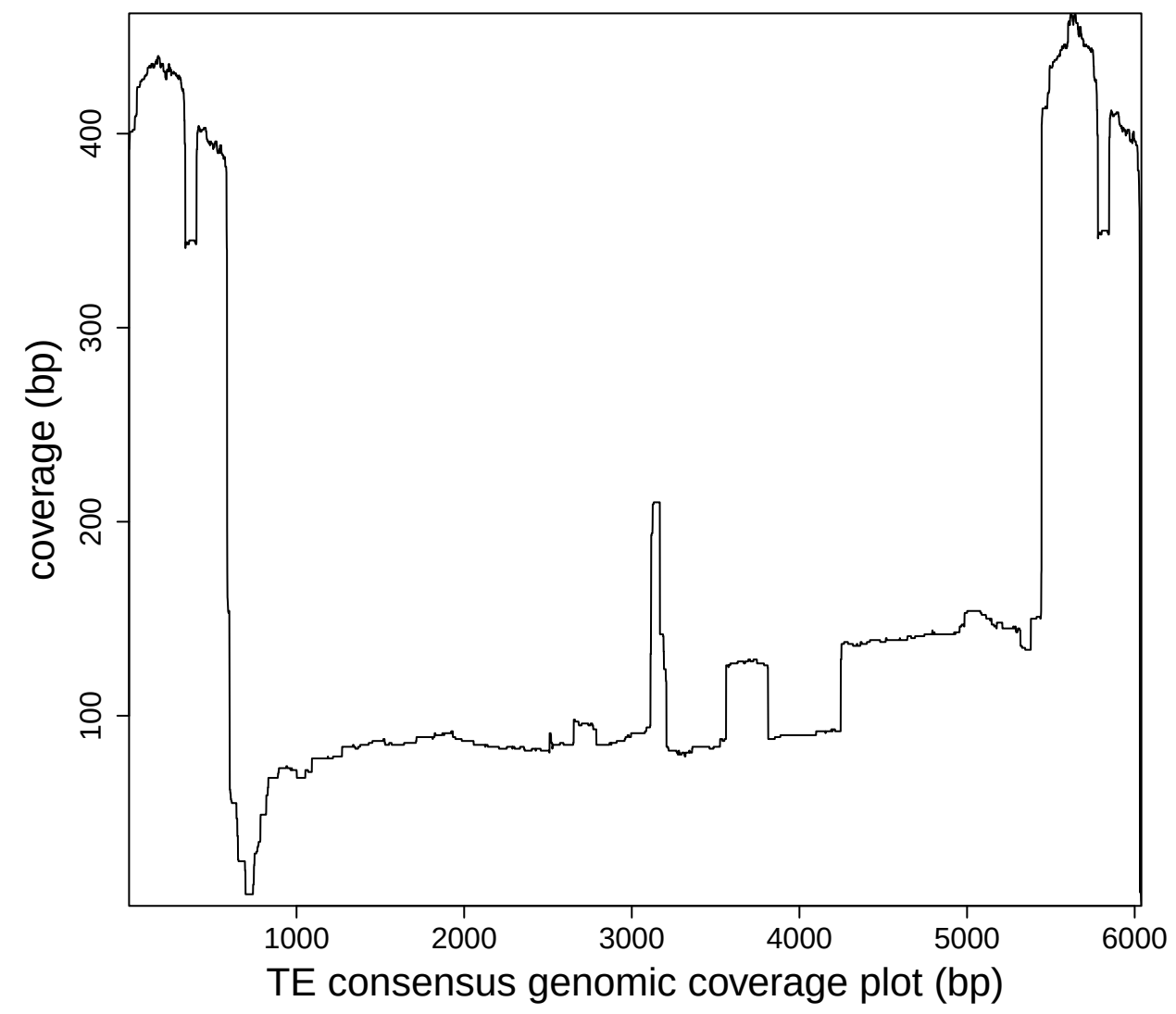
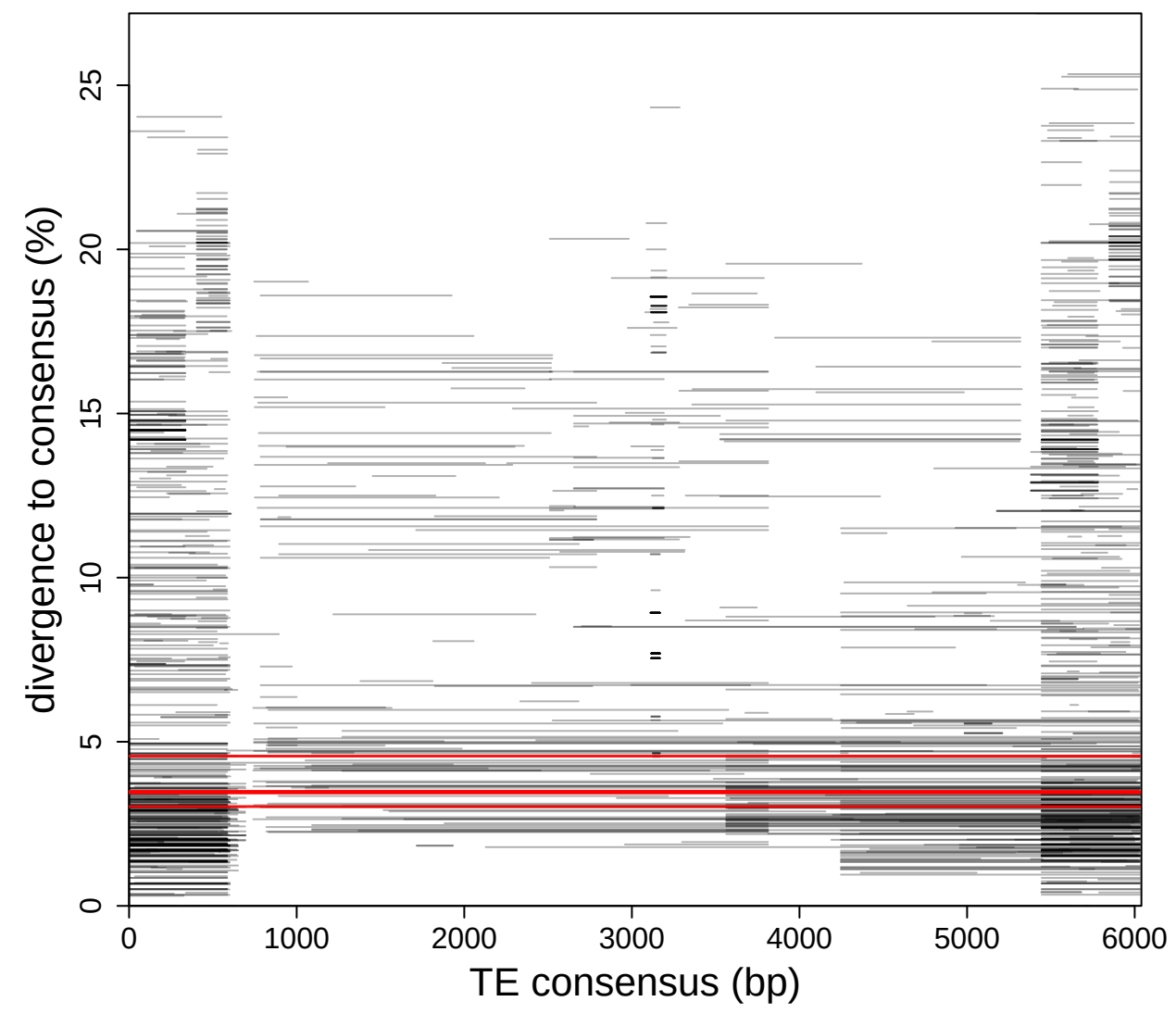
Start crop Point

End crop Point



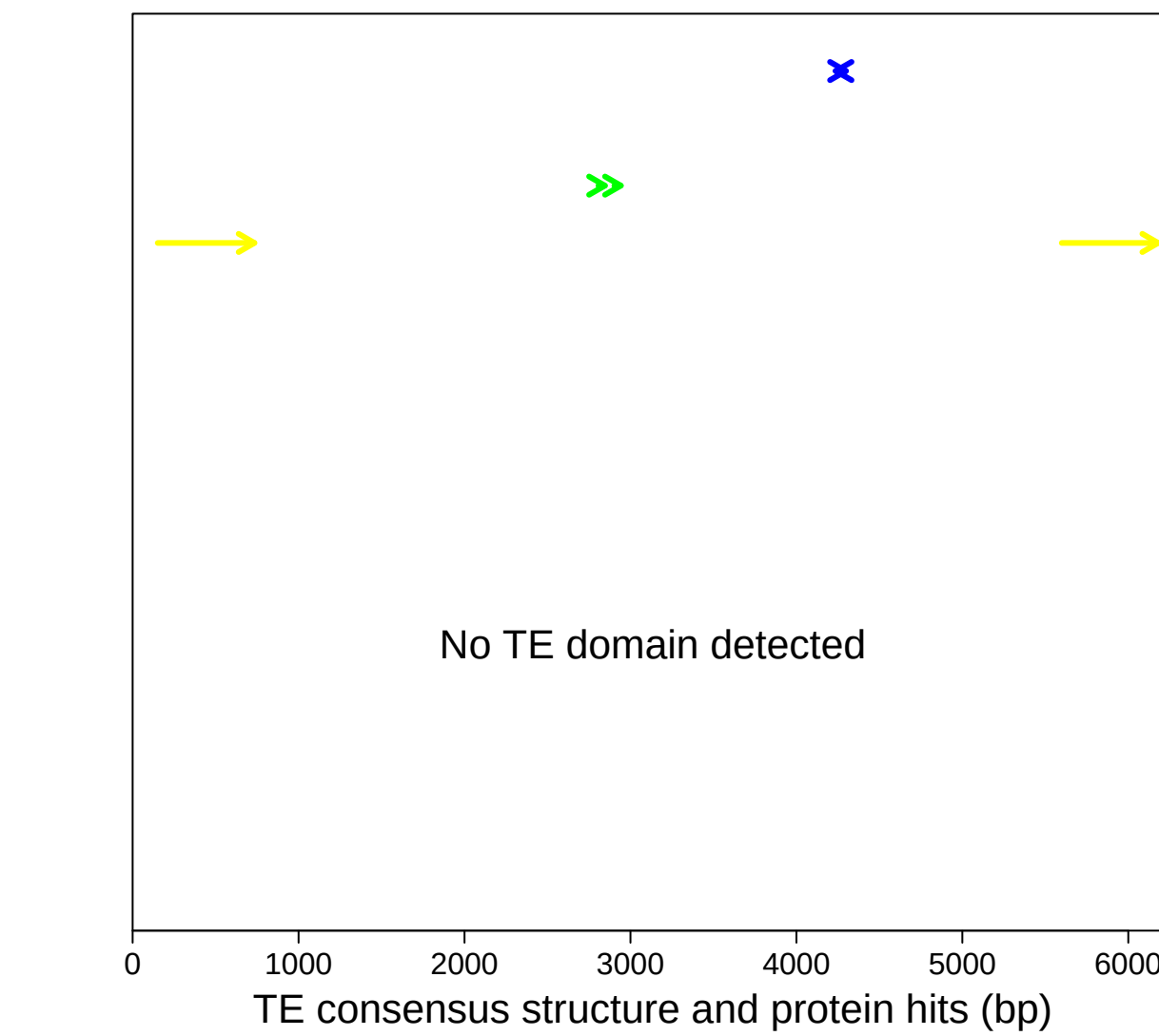
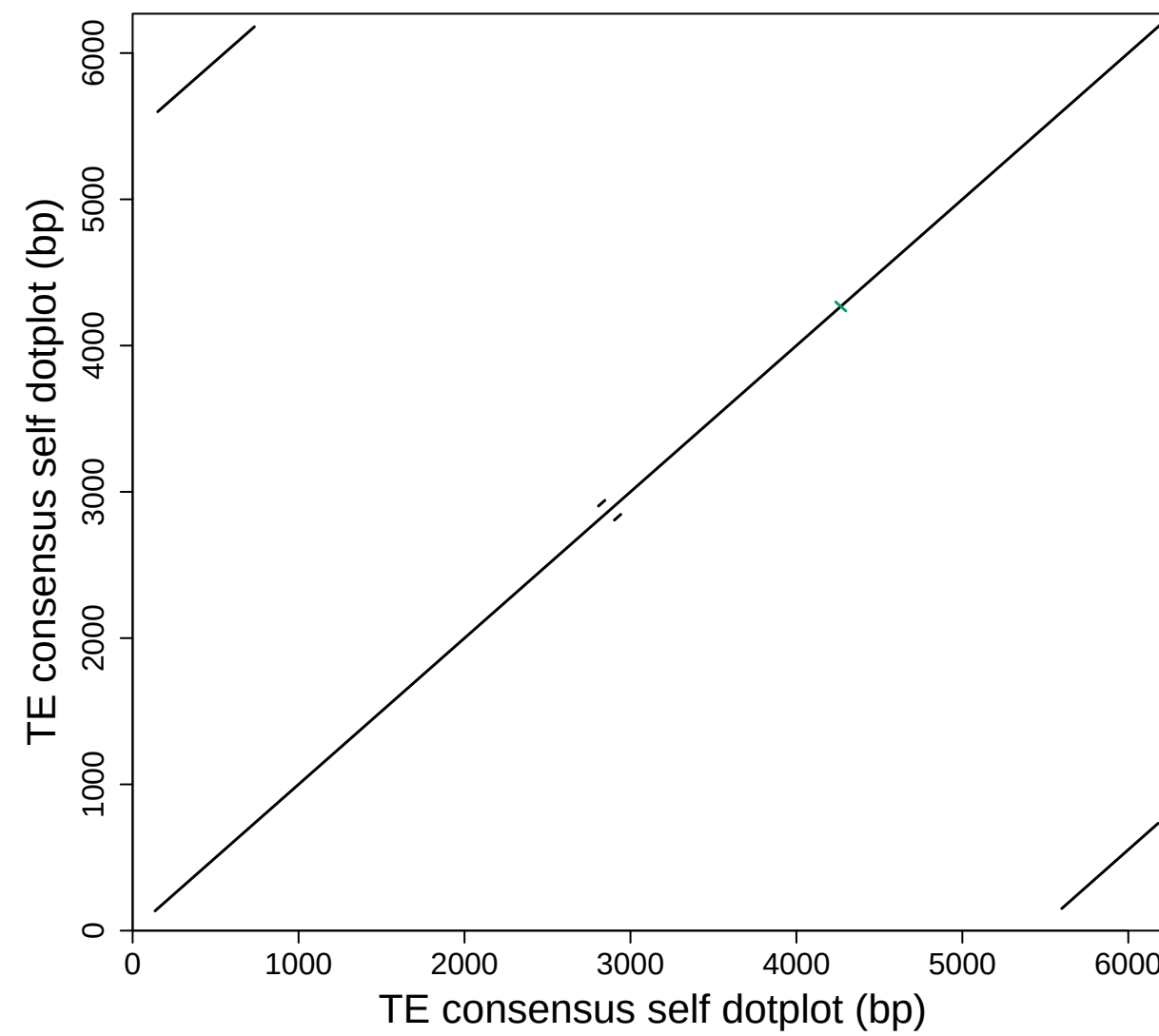
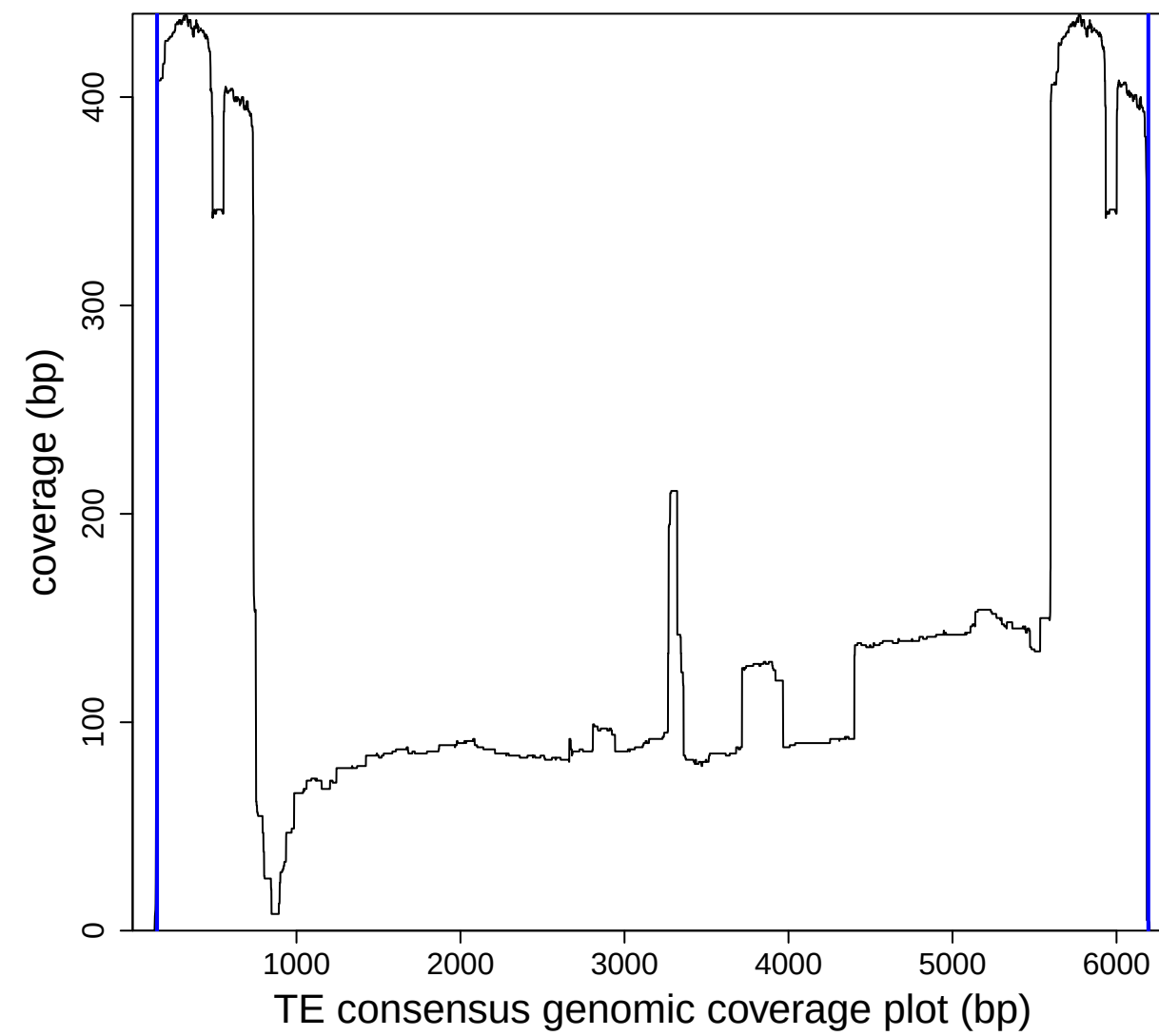
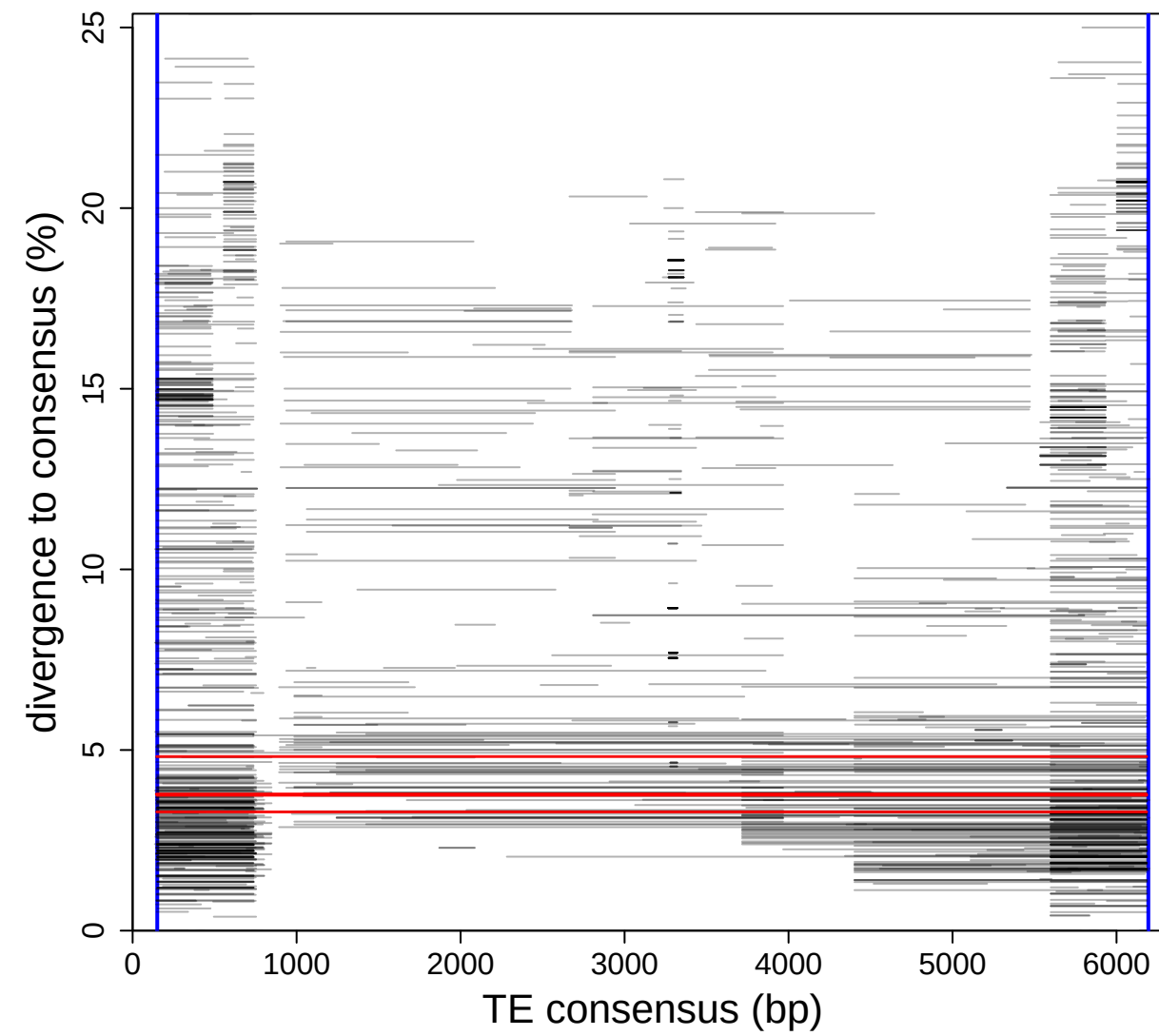
g_1.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln.fa a
size: 6040bp; fragments: 1426; full length: 5 (>=5436bp)

After TEtrimmer 6040 bp

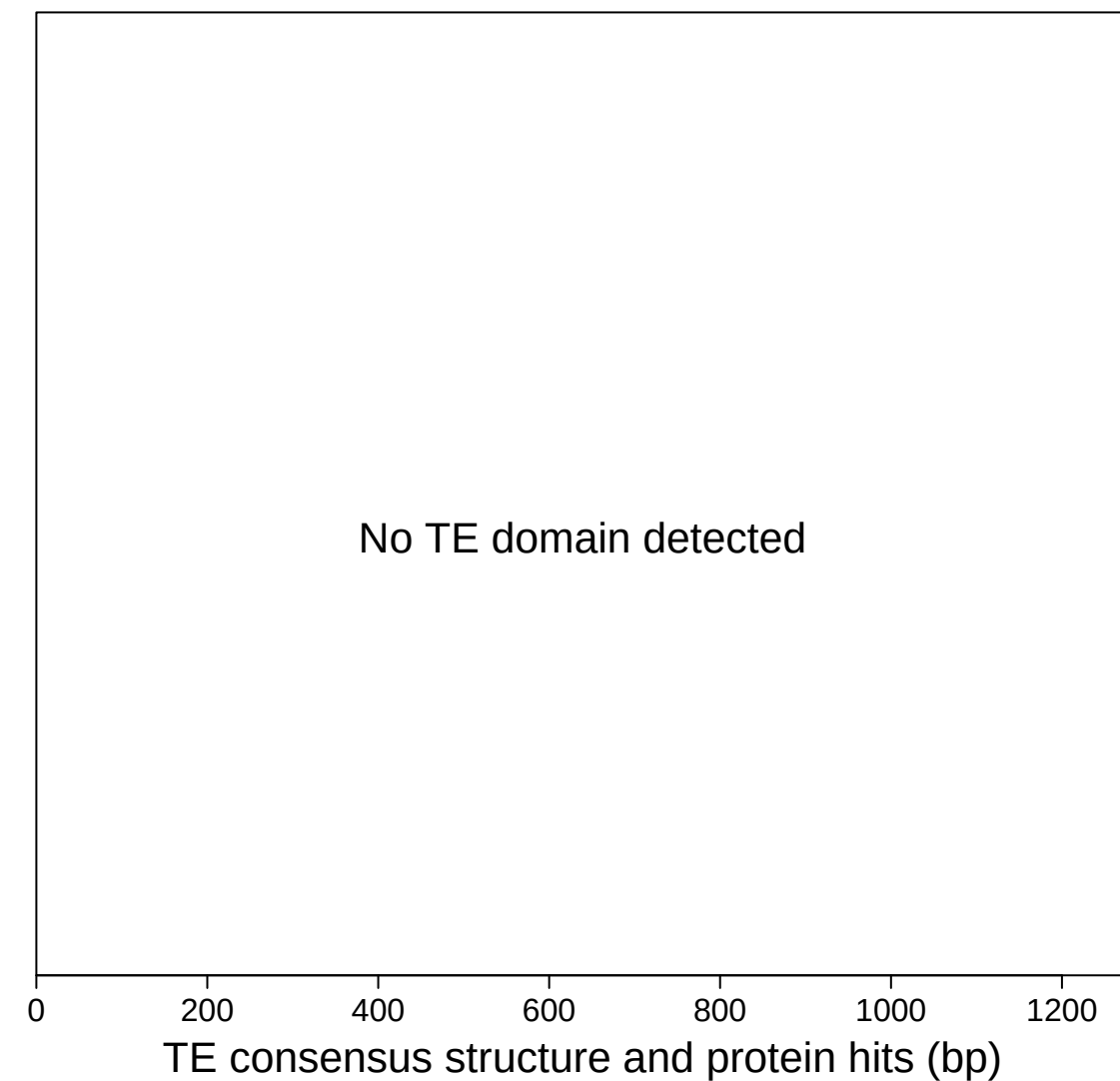
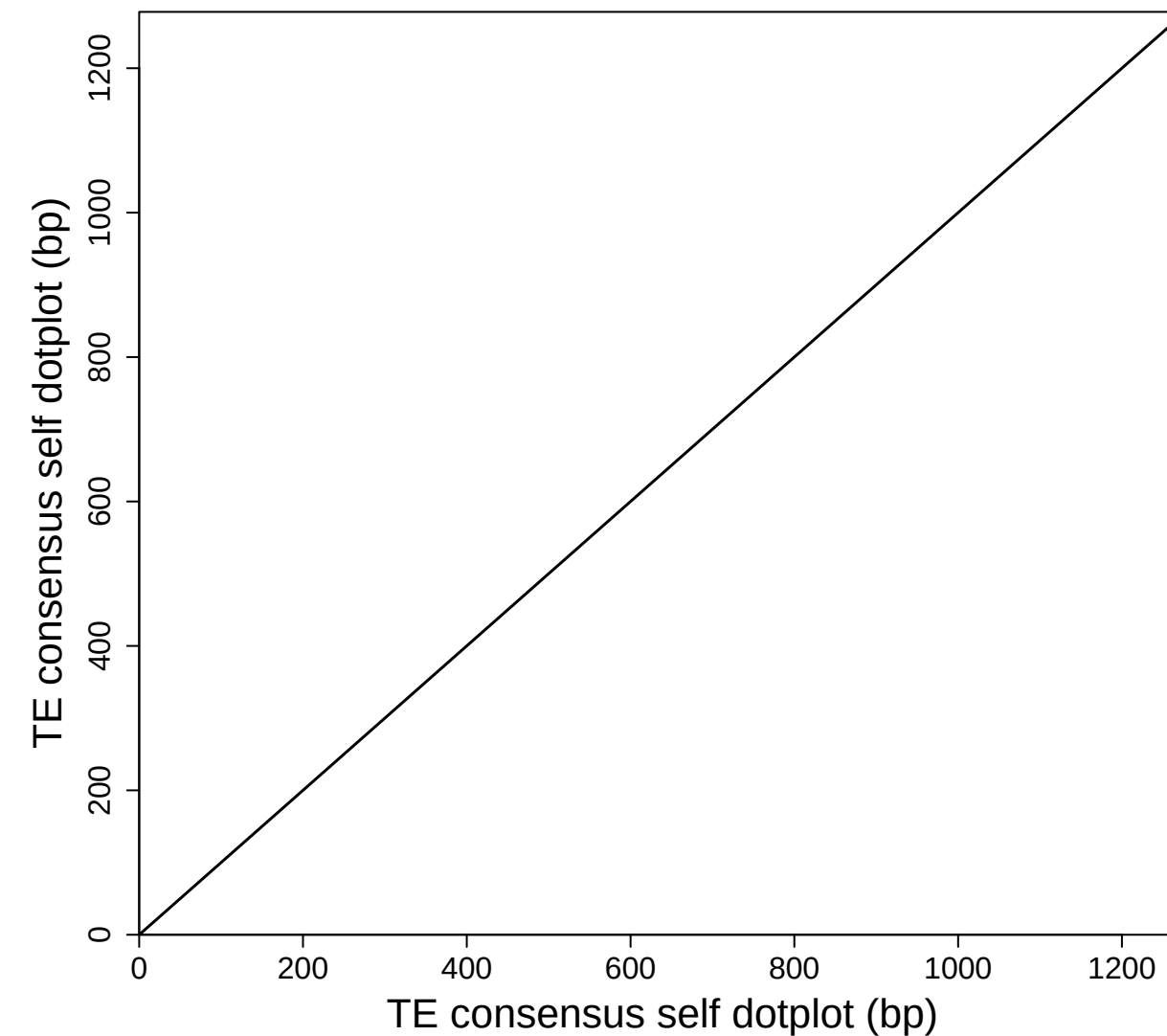
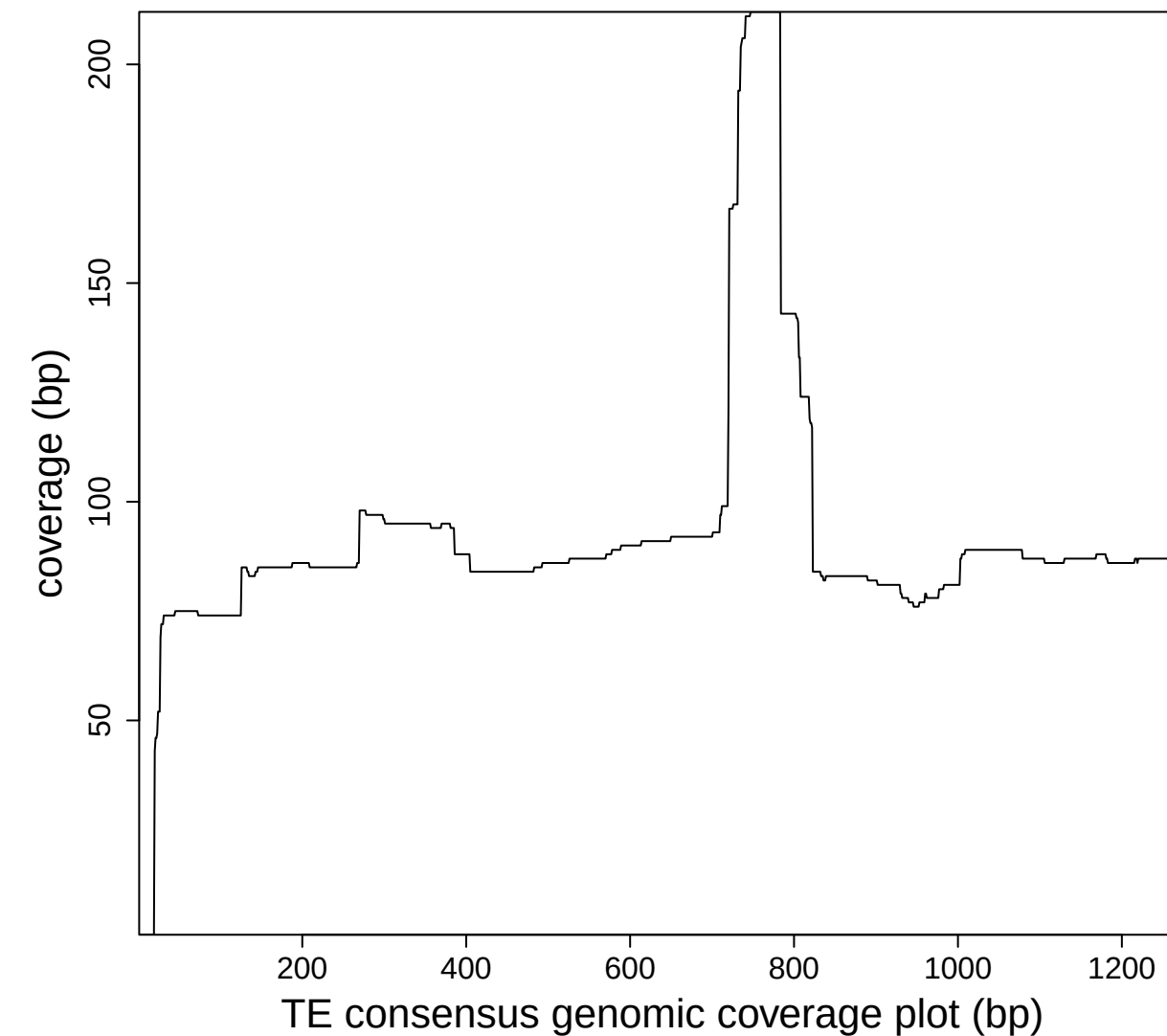
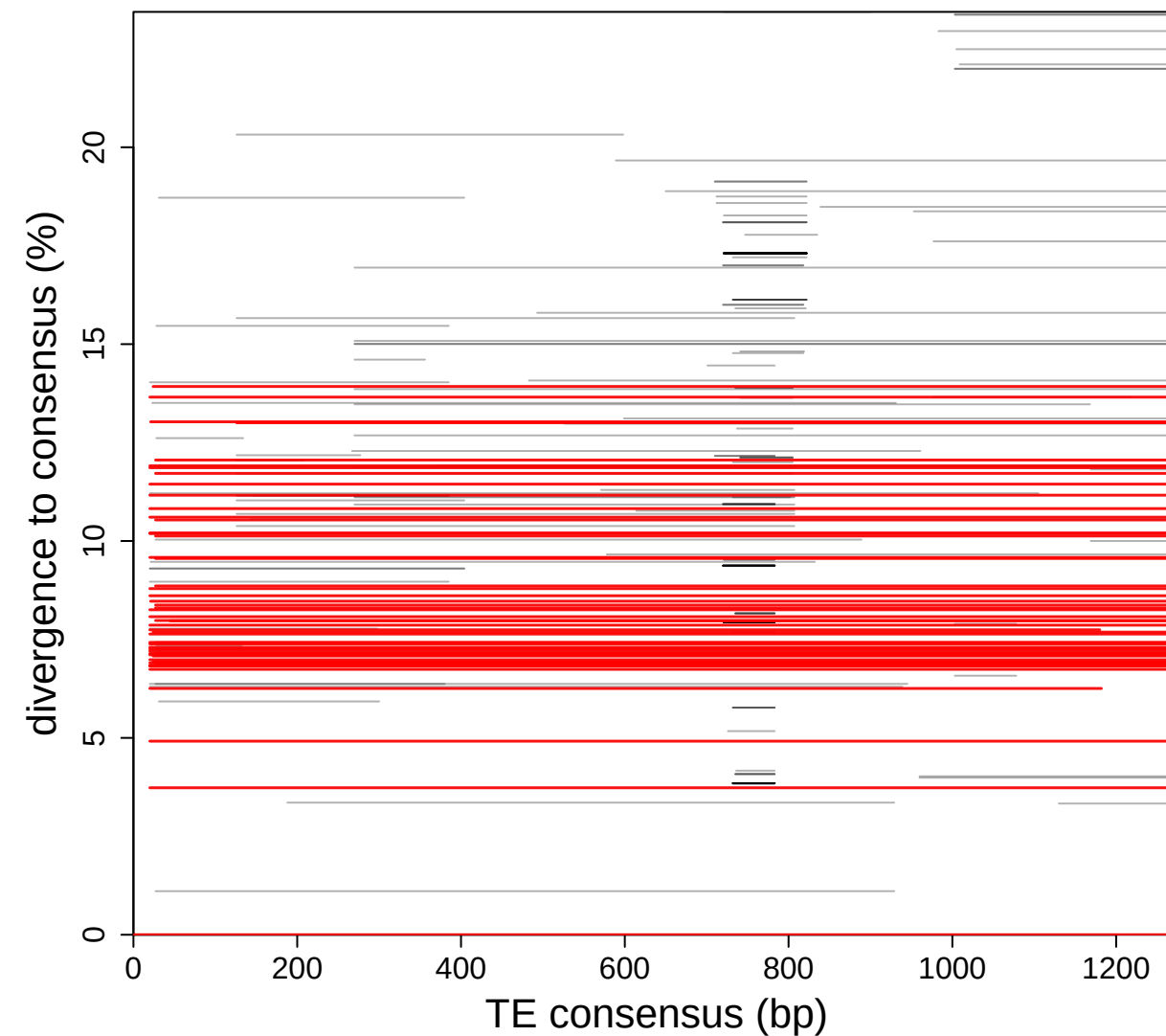


.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 6344bp; fragments: 1410; full length: 5 (>=5709.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

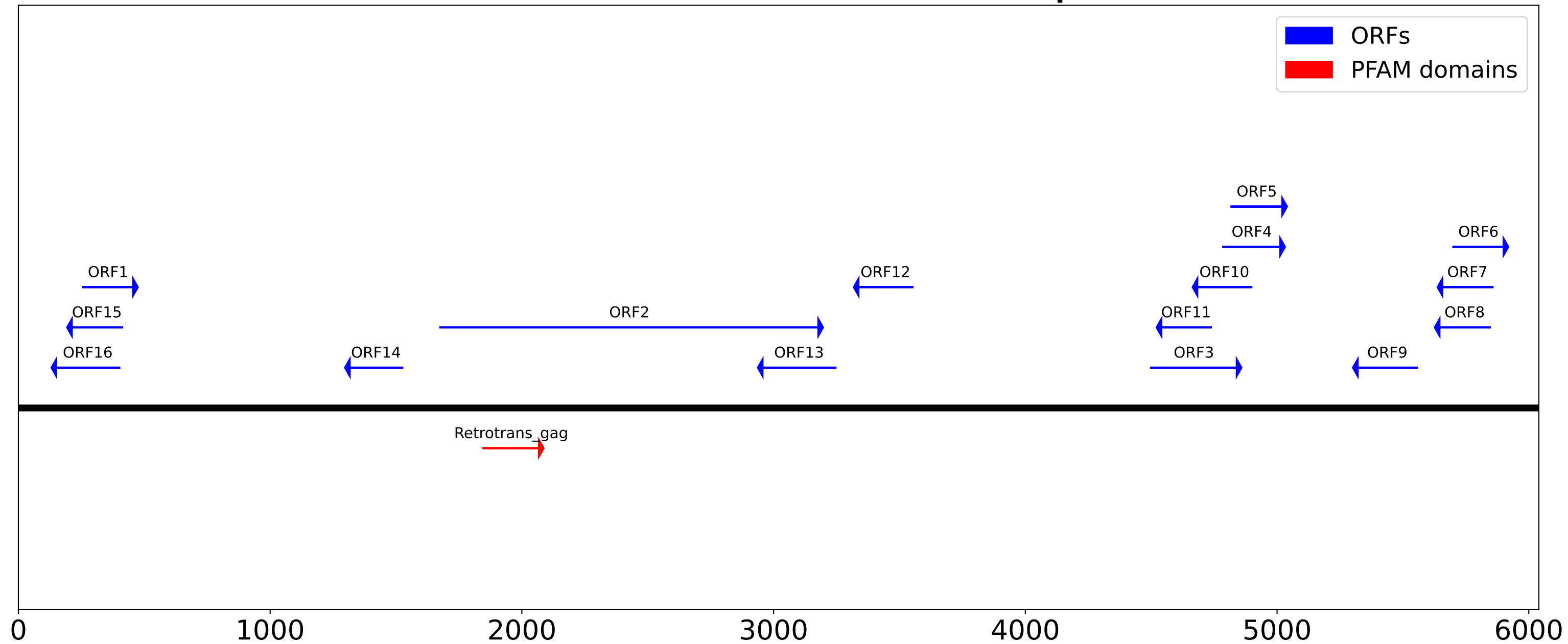


TE: ltr_1_family_1030
size: 1278bp; fragments: 252; full length: 57 (≥ 1150.2 bp)

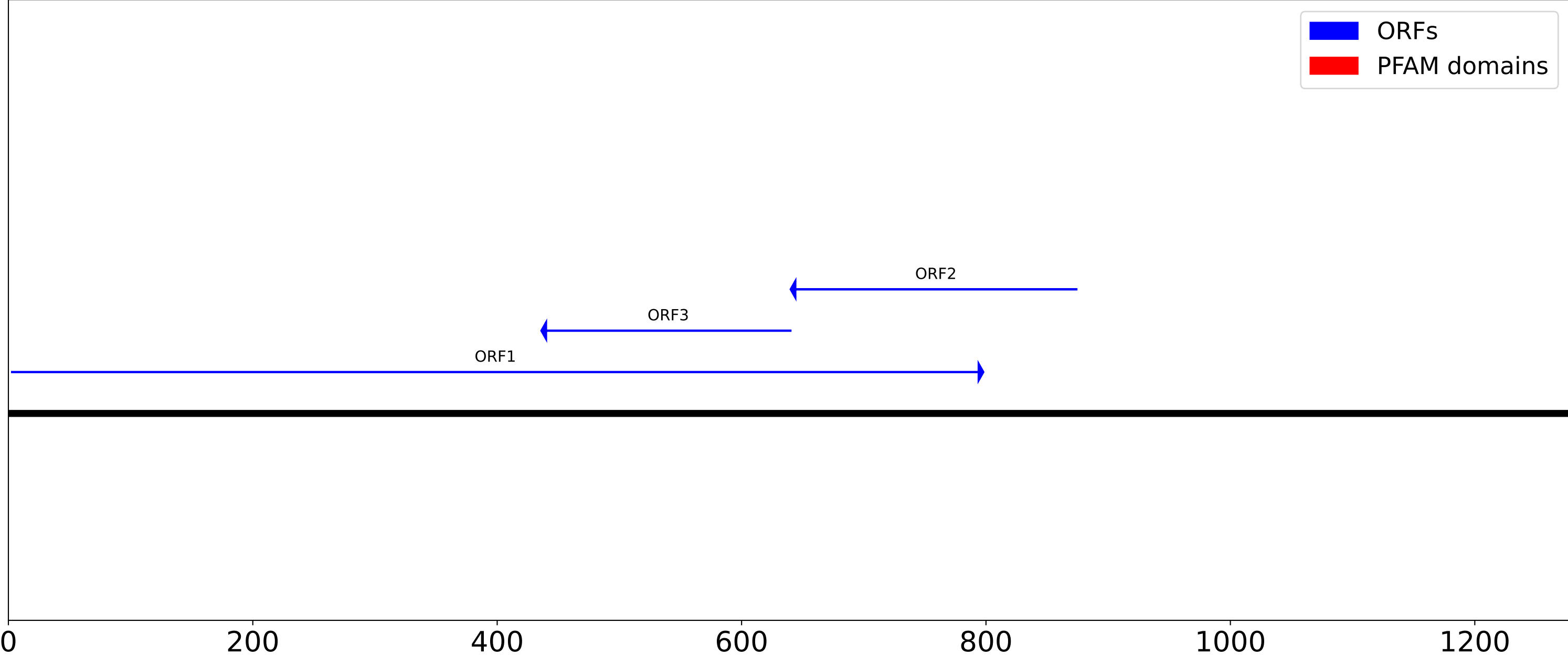


Before TEtrimmer 1278 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)

