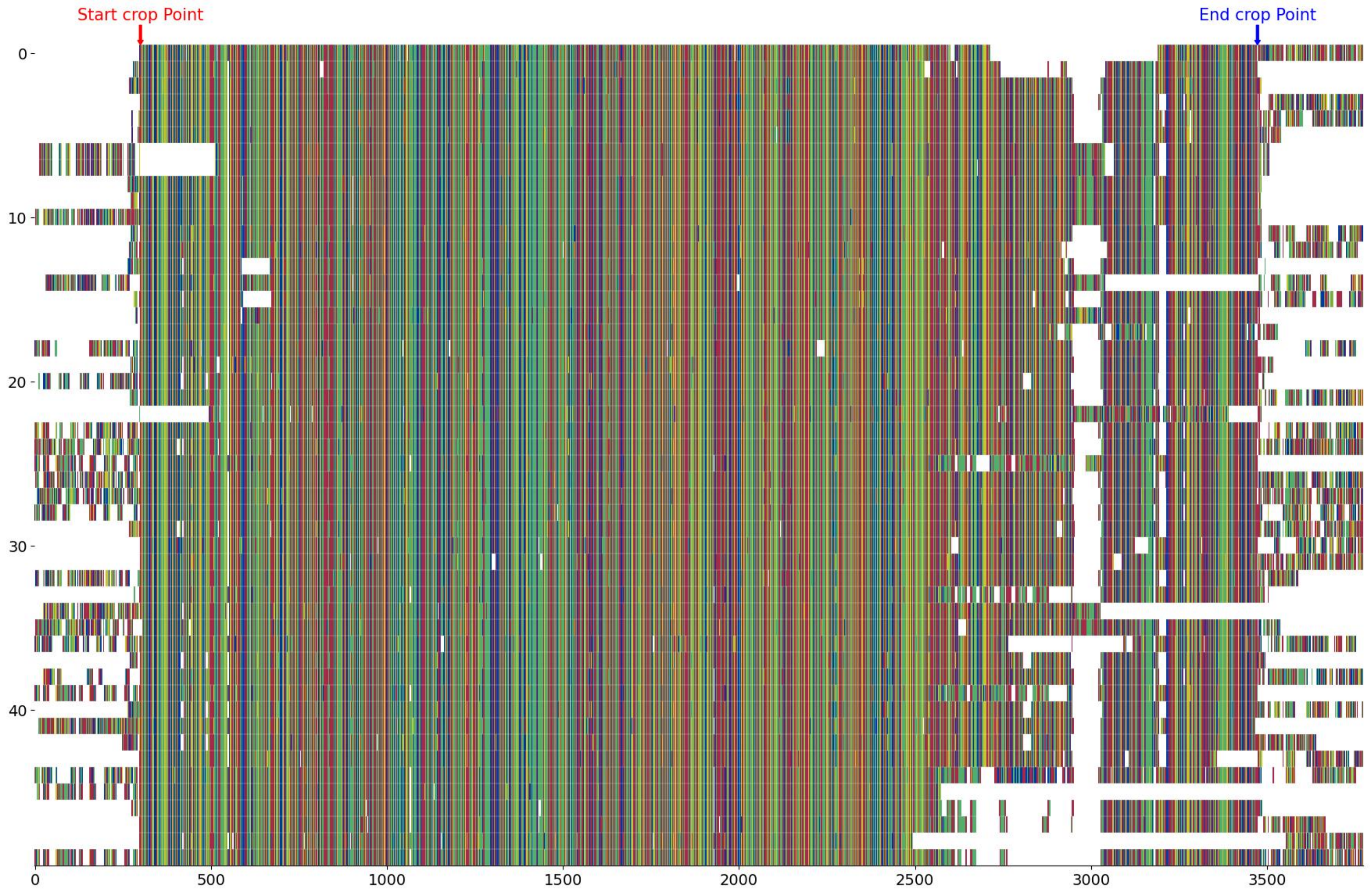




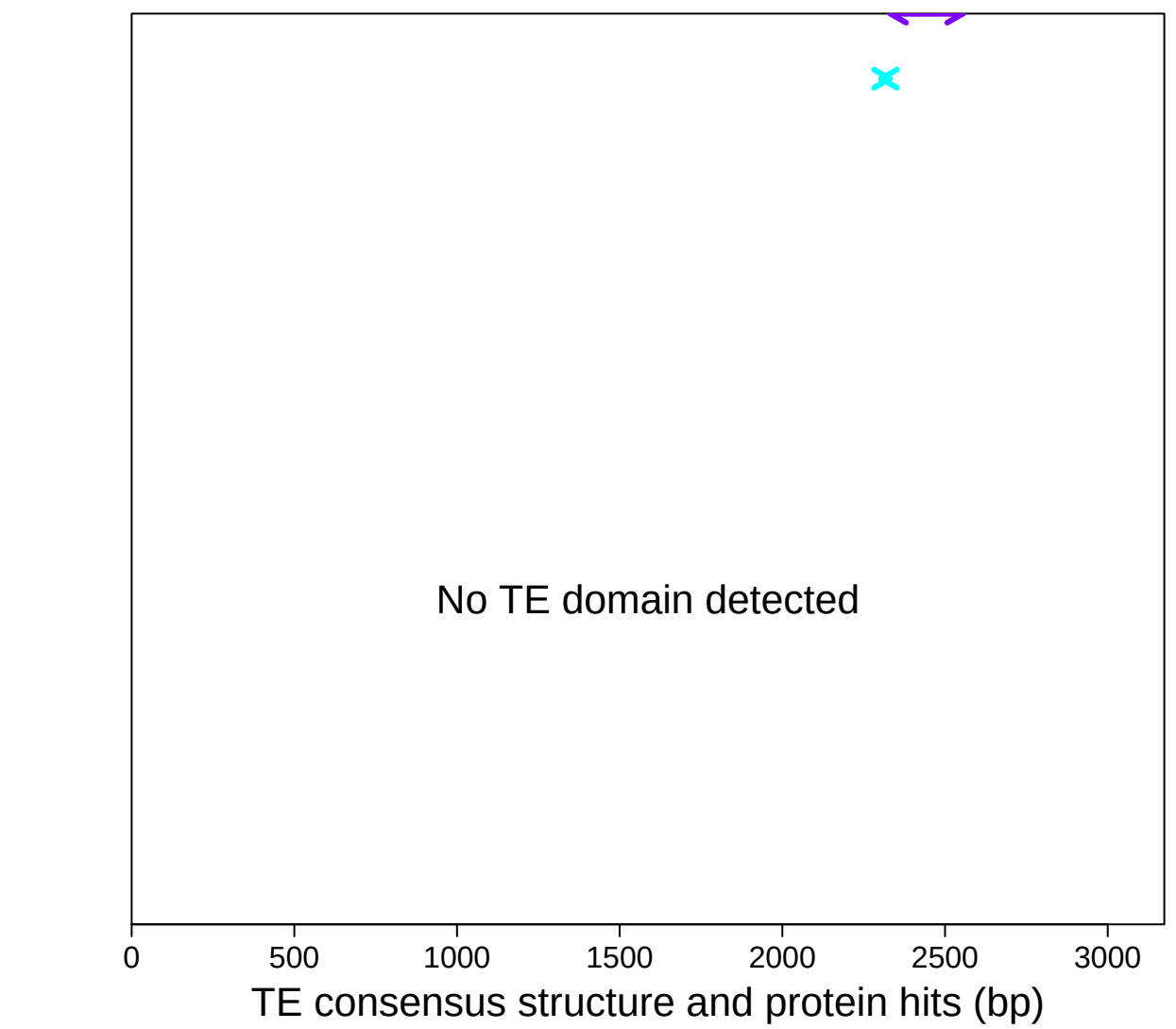
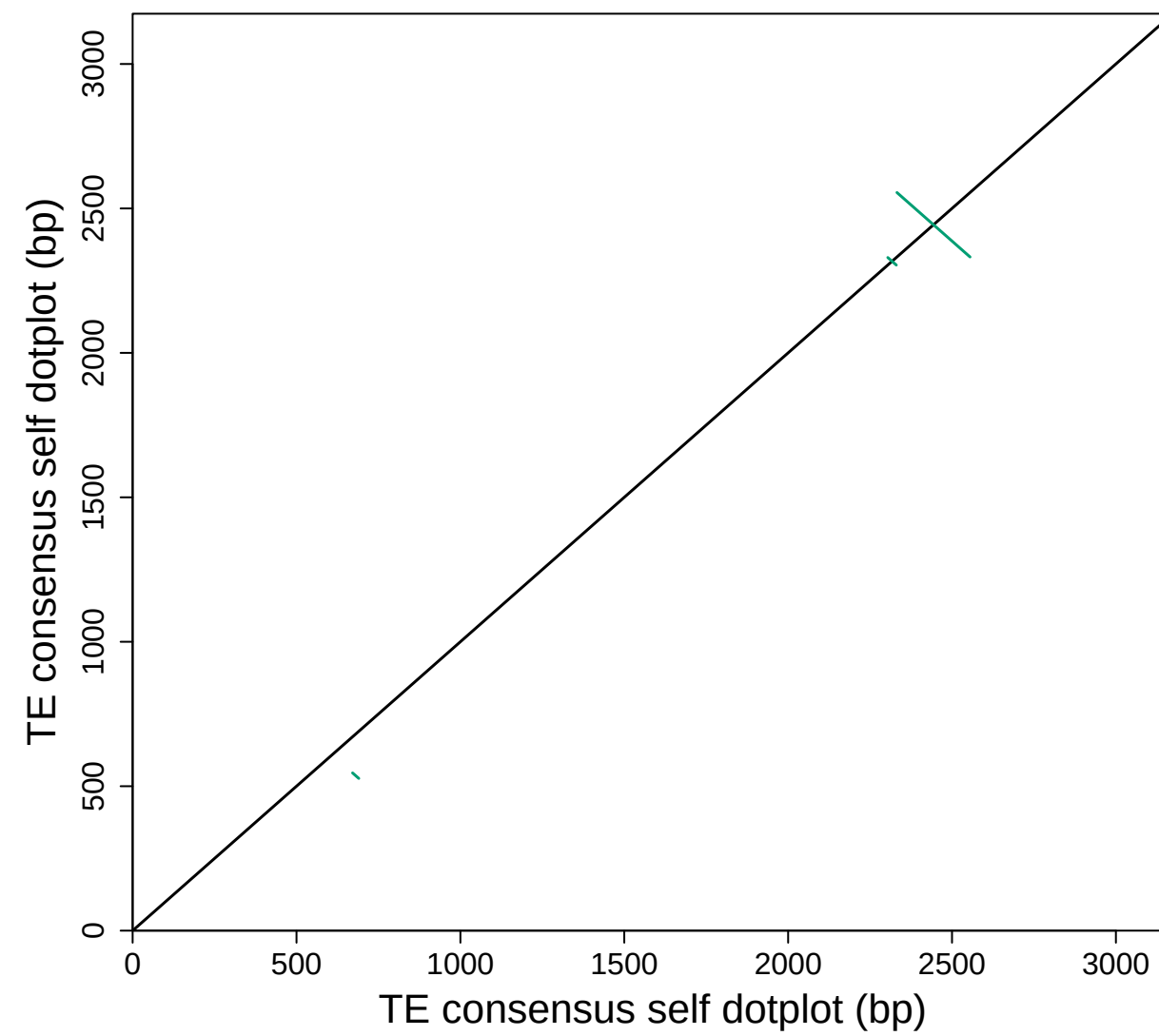
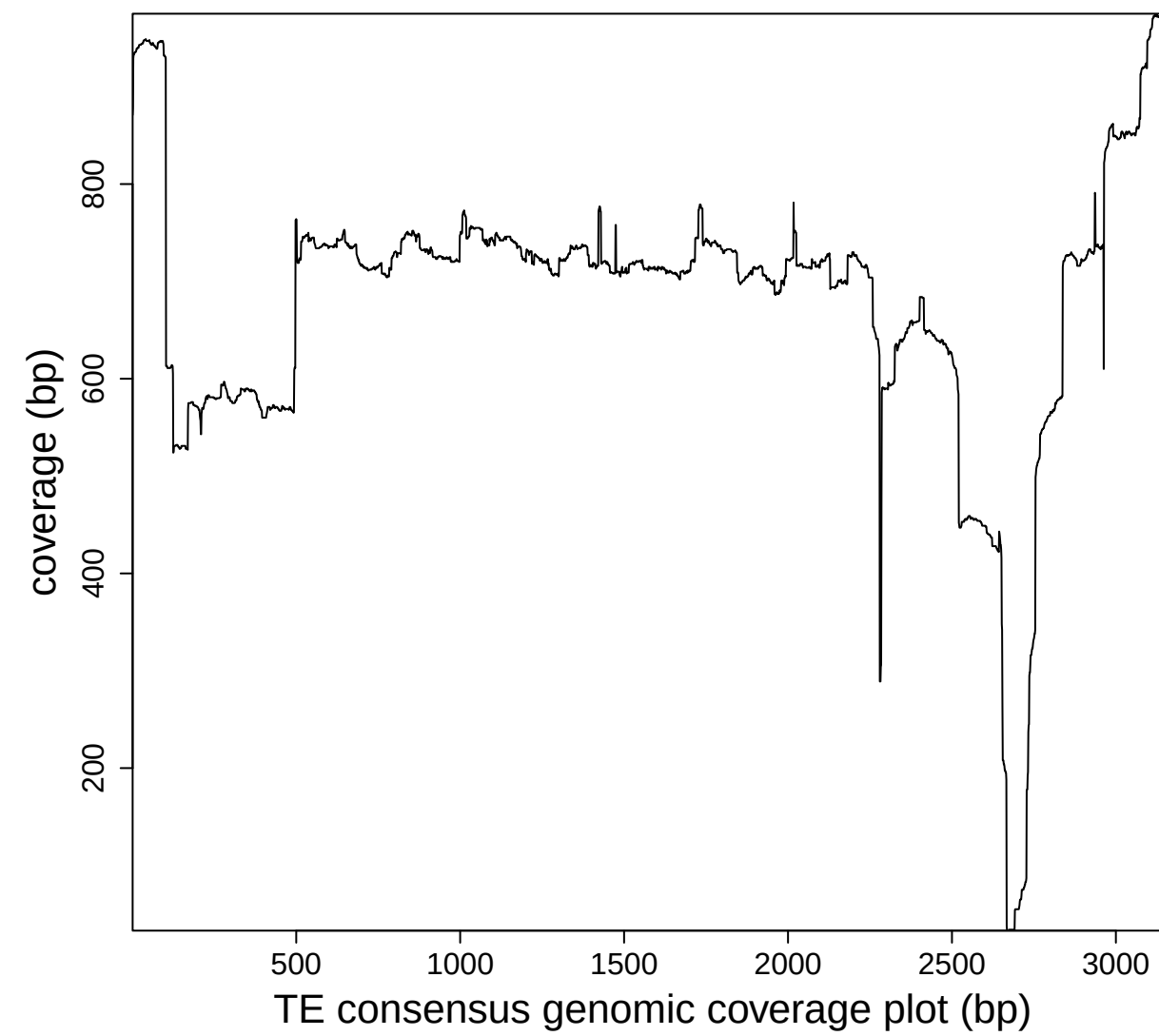
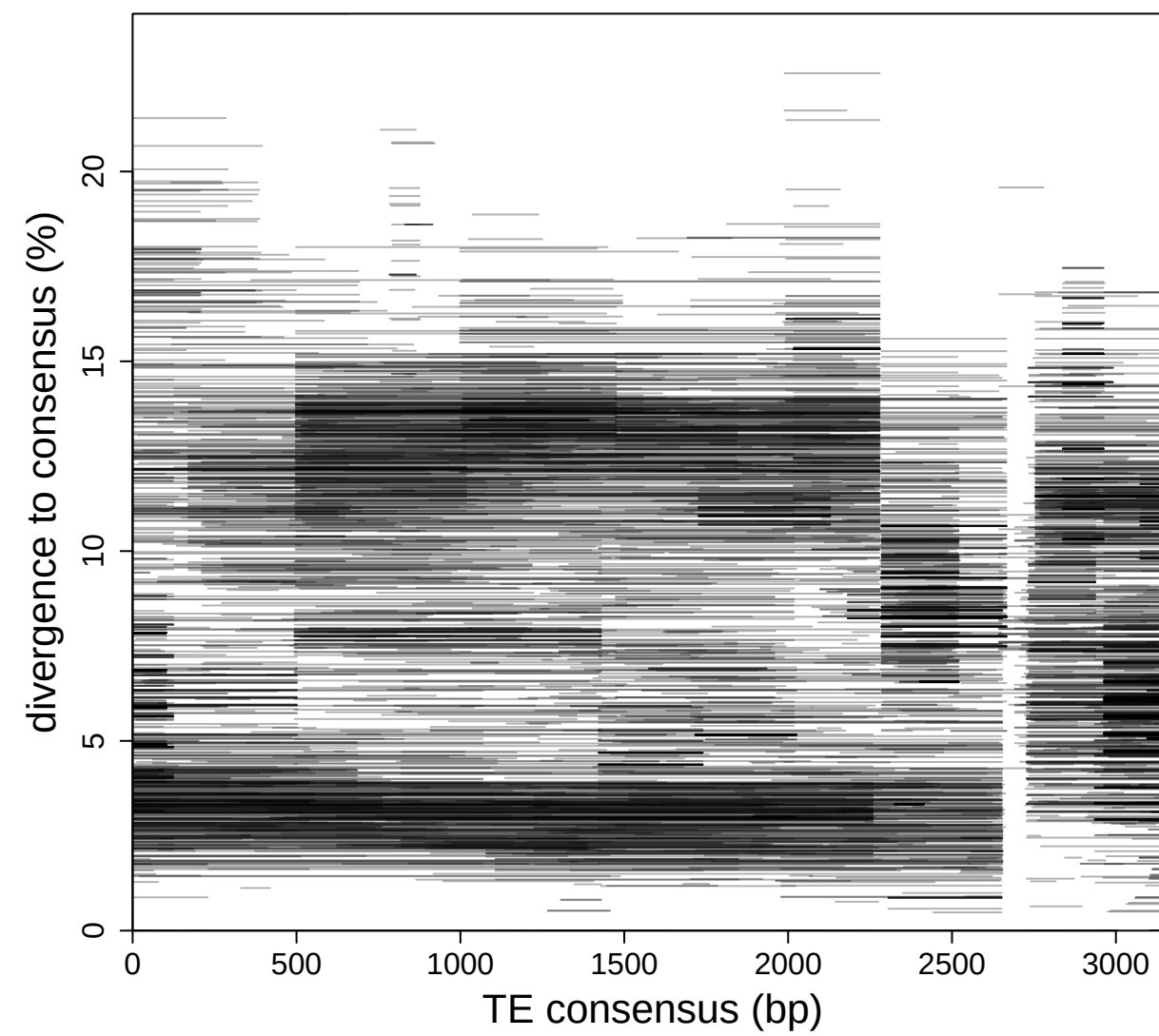
Start crop Point

End crop Point



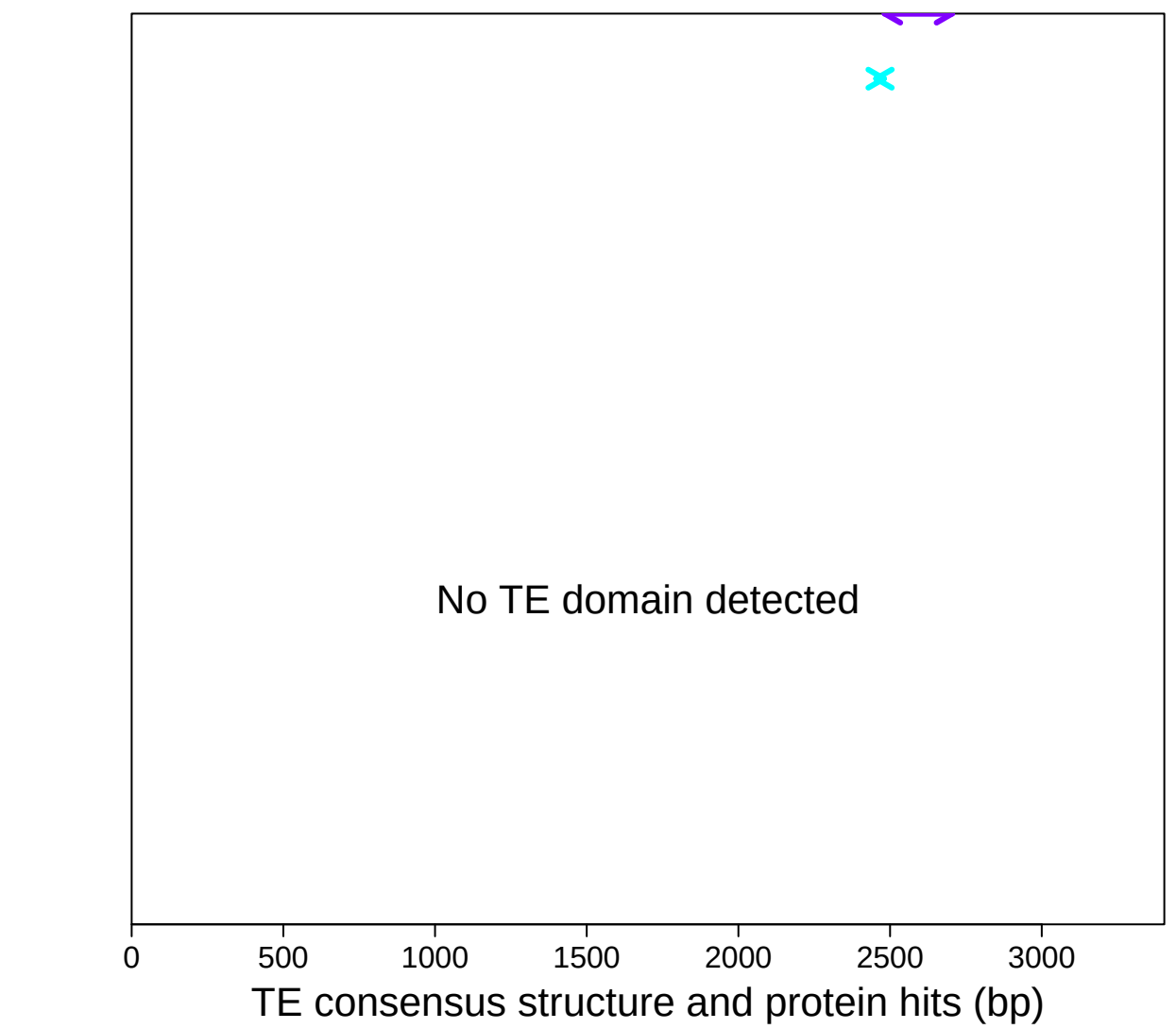
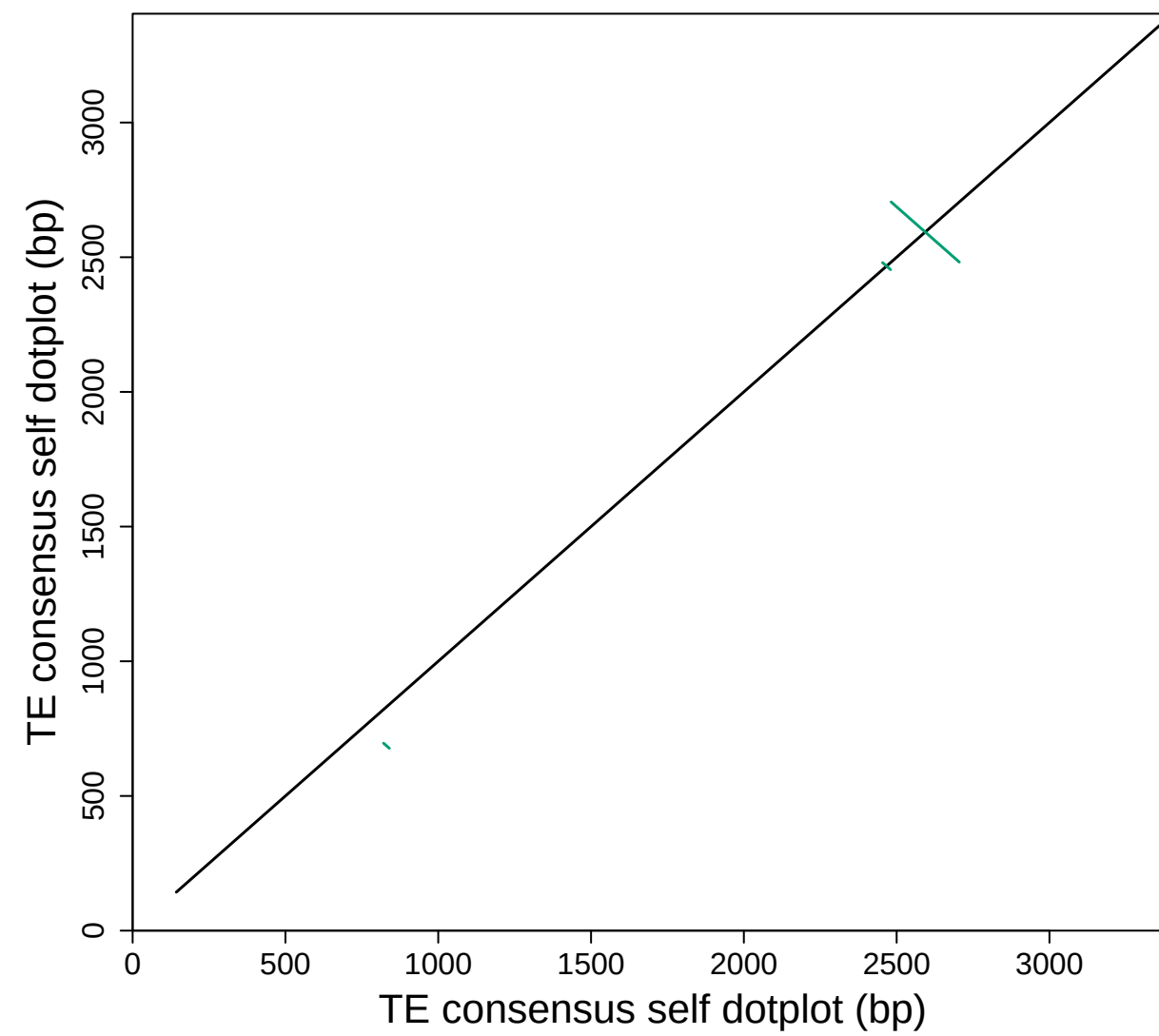
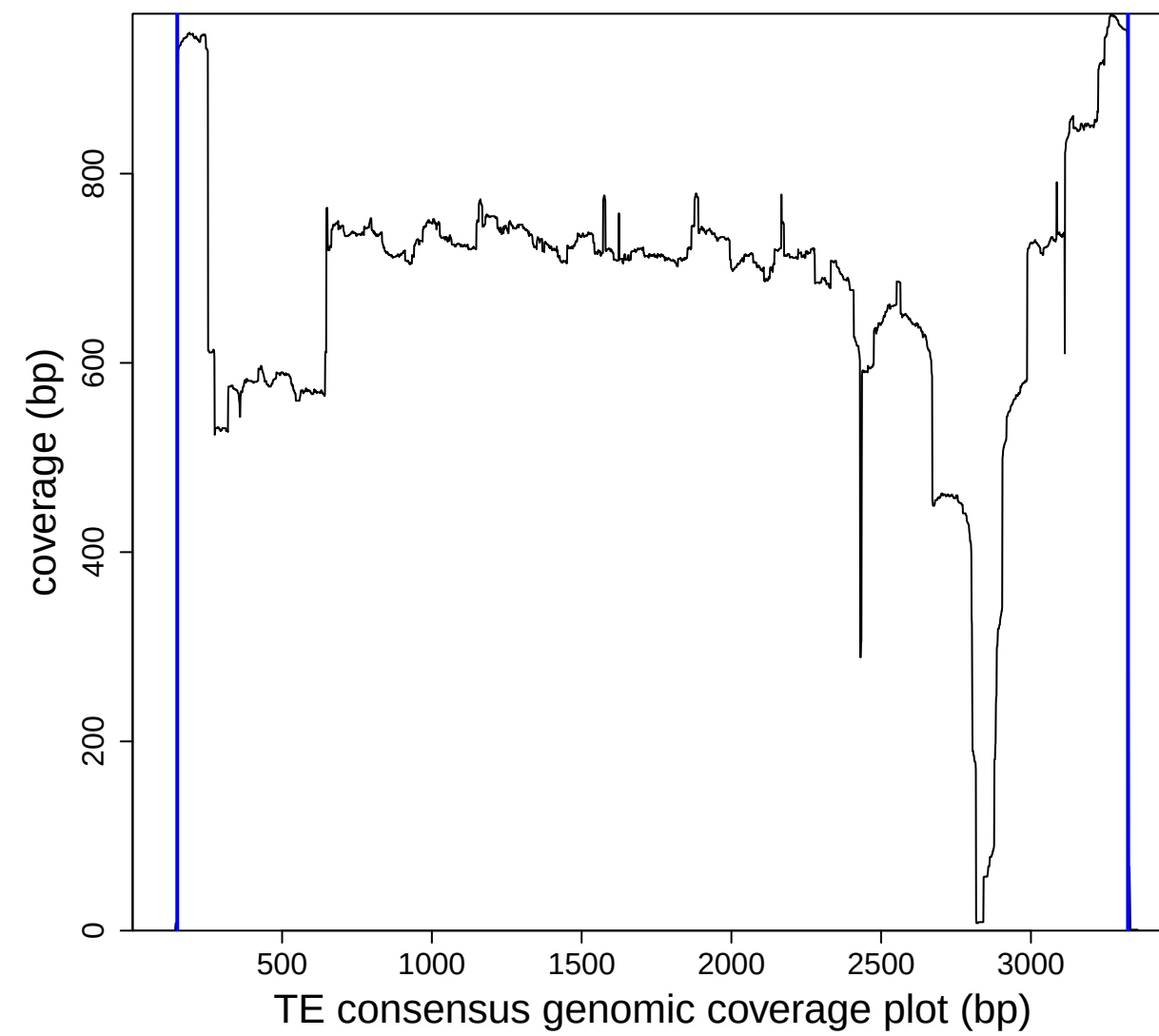
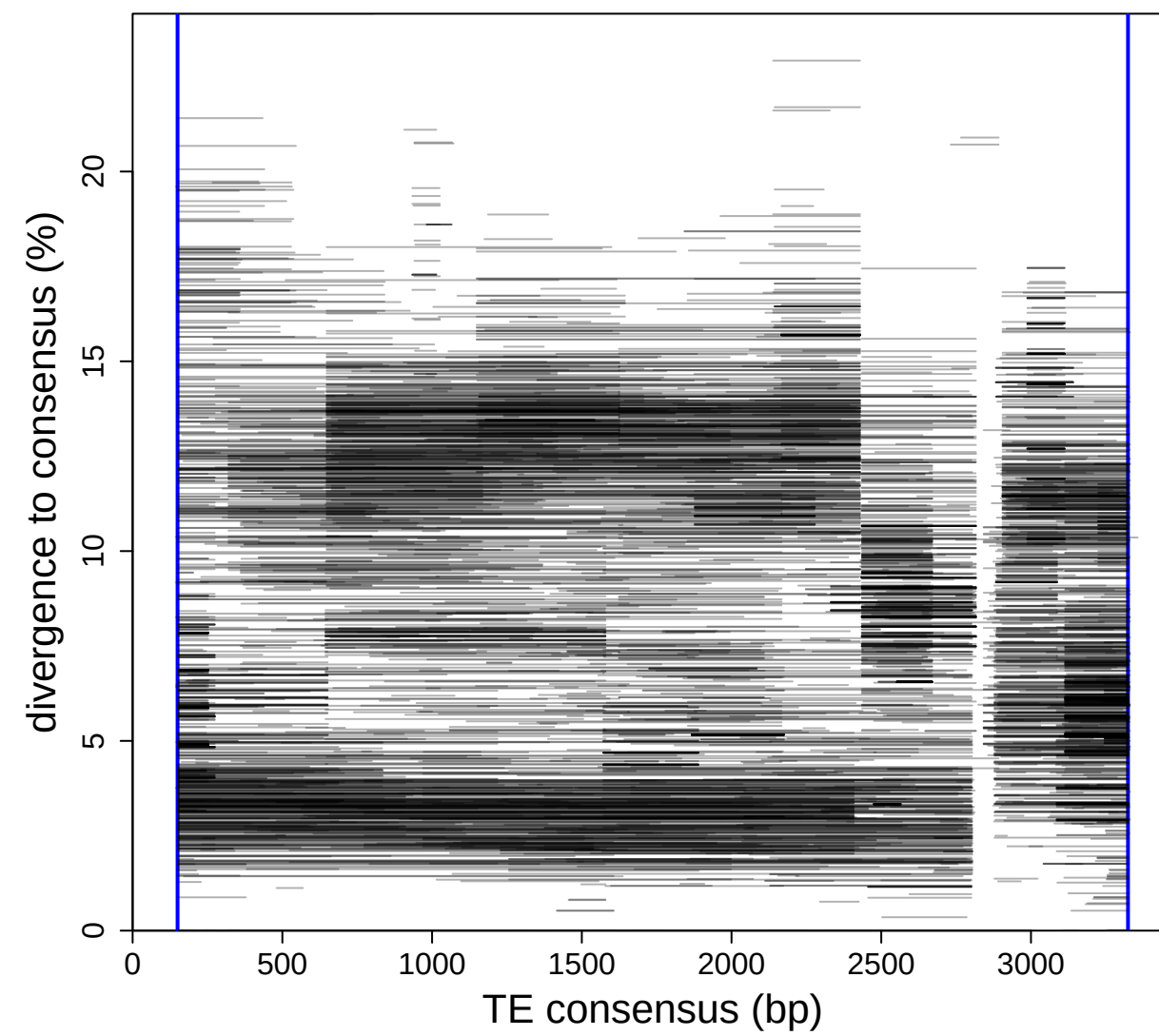
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 3174bp; fragments: 4257; full length: 0 (>=2856.6bp)

After TEtrimmer 3174 bp



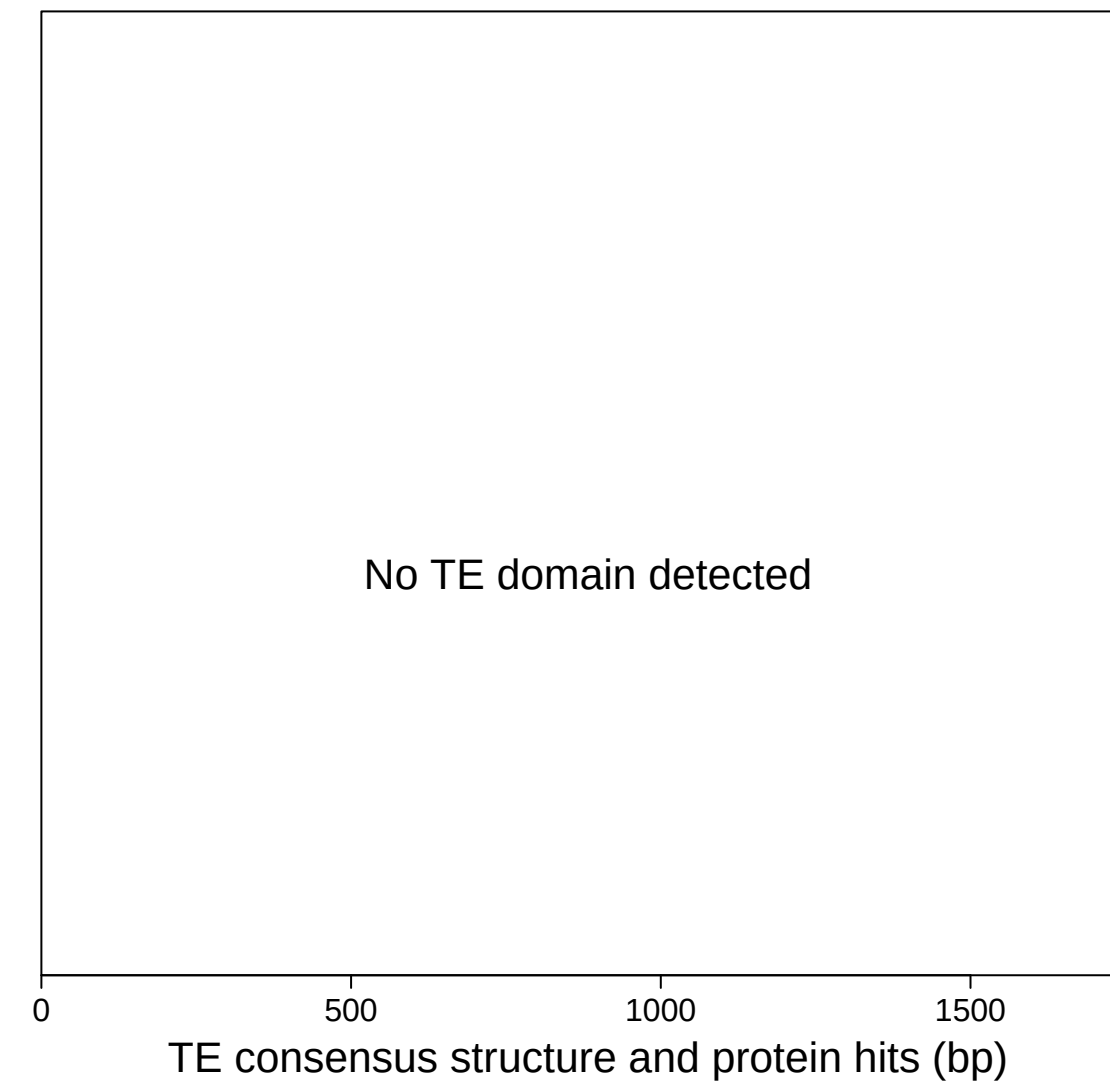
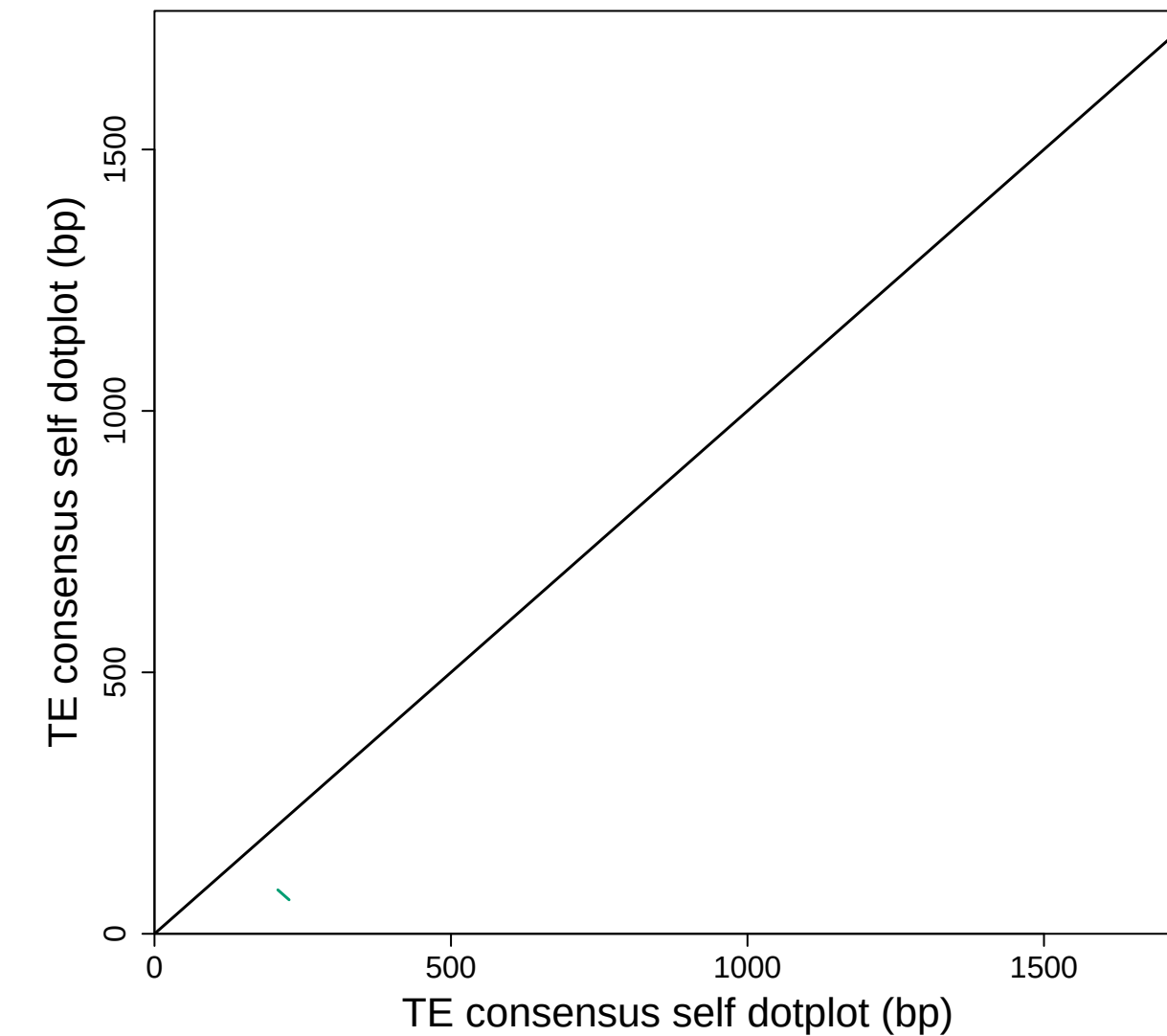
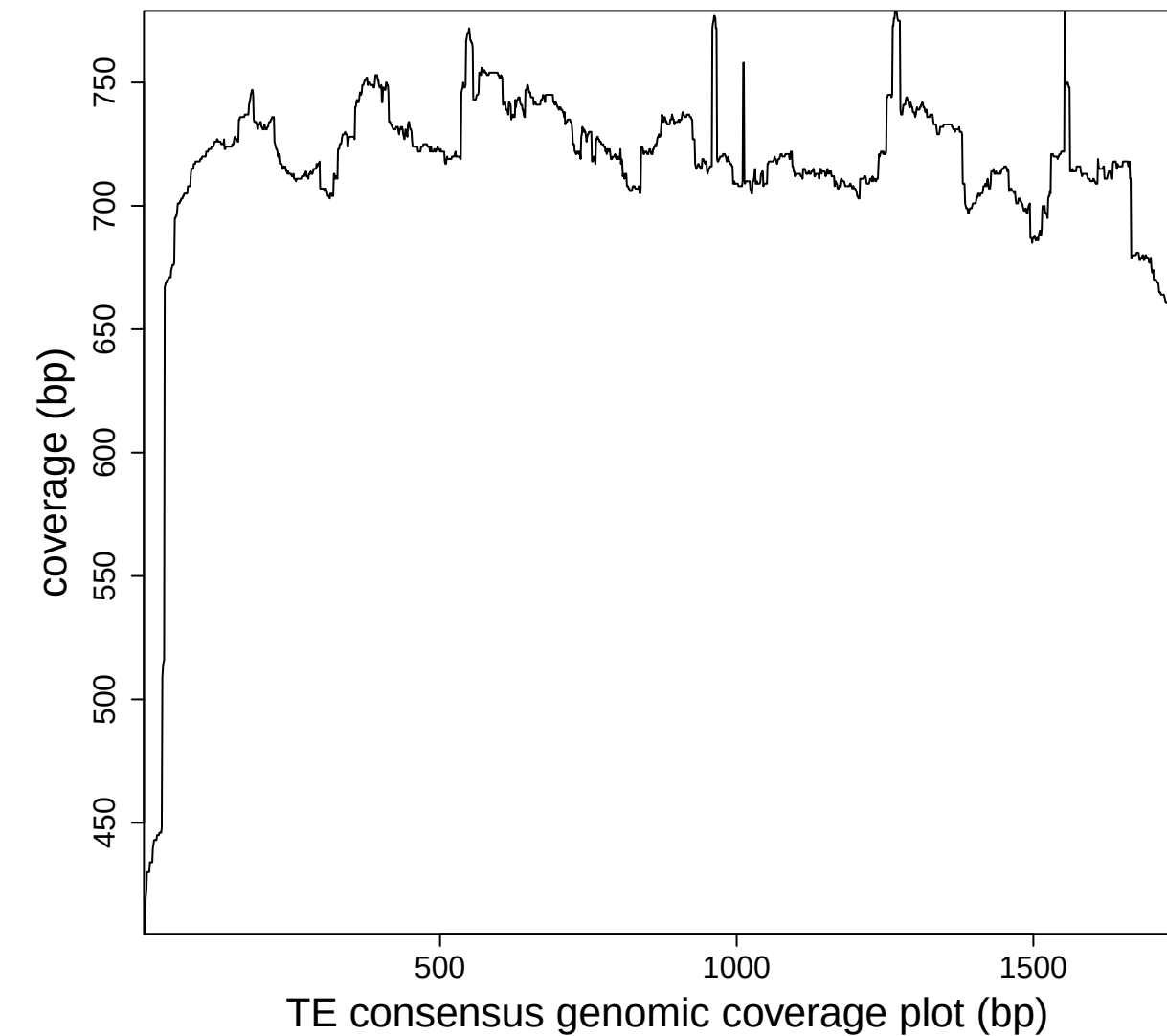
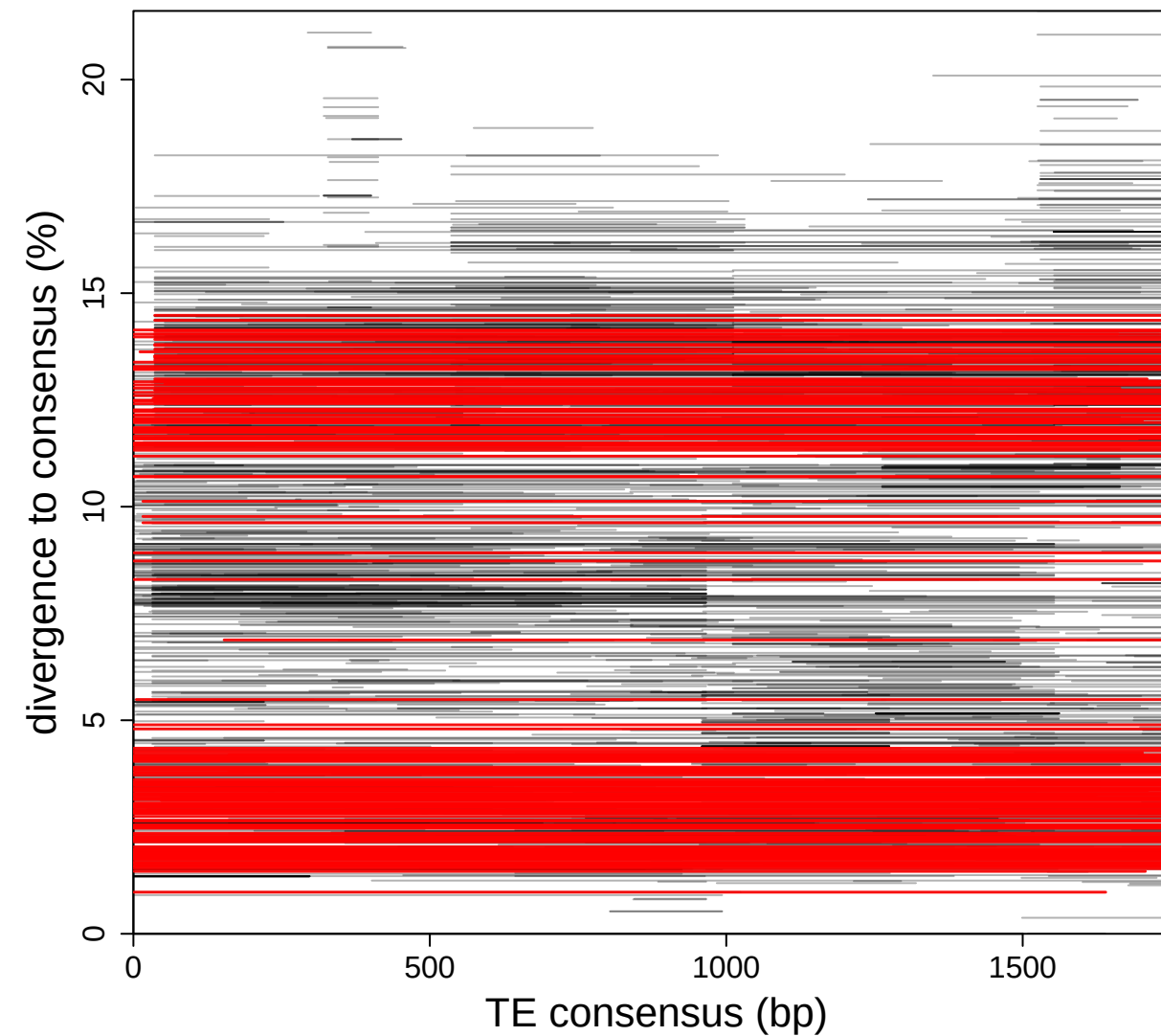
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 3474bp; fragments: 4238; full length: 0 (>=3126.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

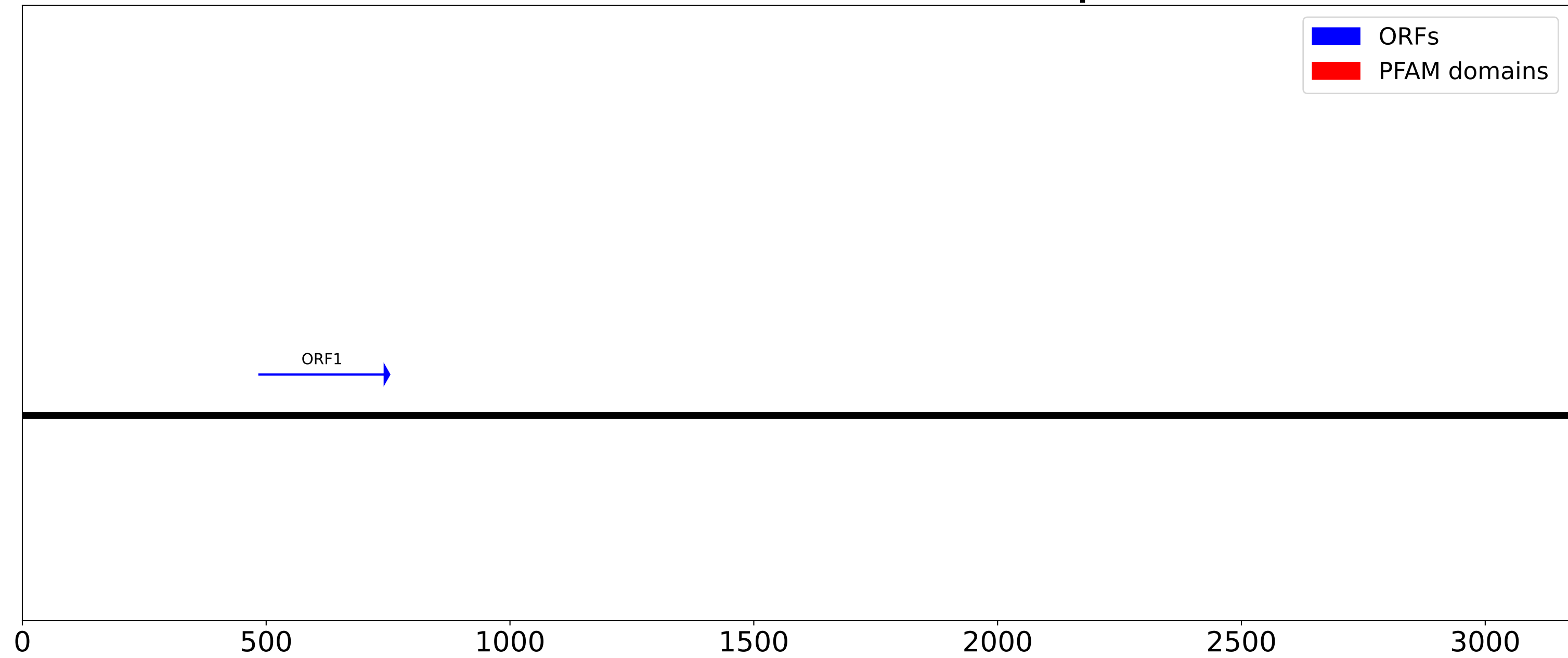


TE: rnd_1_family_400
size: 1765bp; fragments: 1734; full length: 200 (≥ 1588.5 bp)

Before TEtrimmer 1765 bp



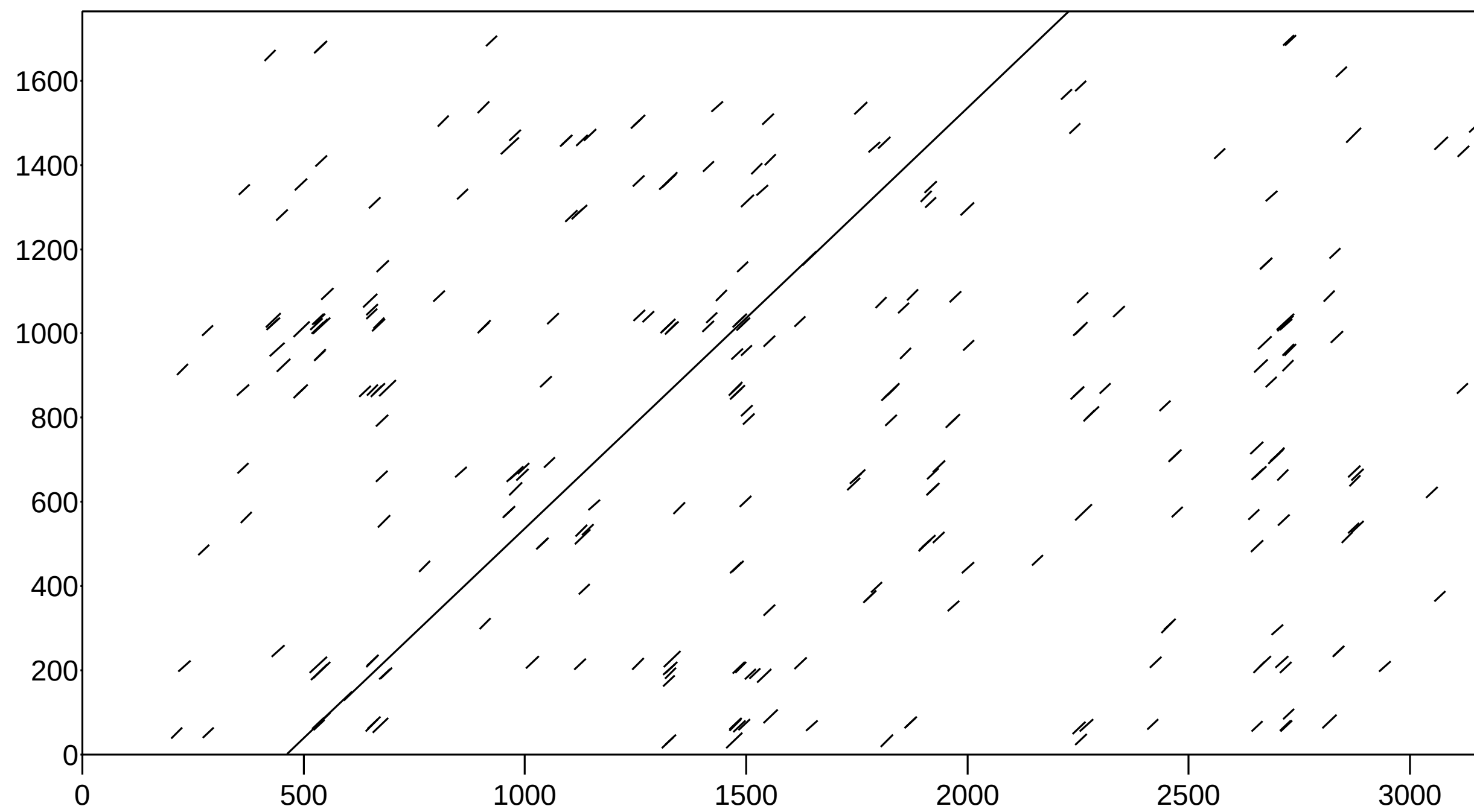
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)