

Start crop Point

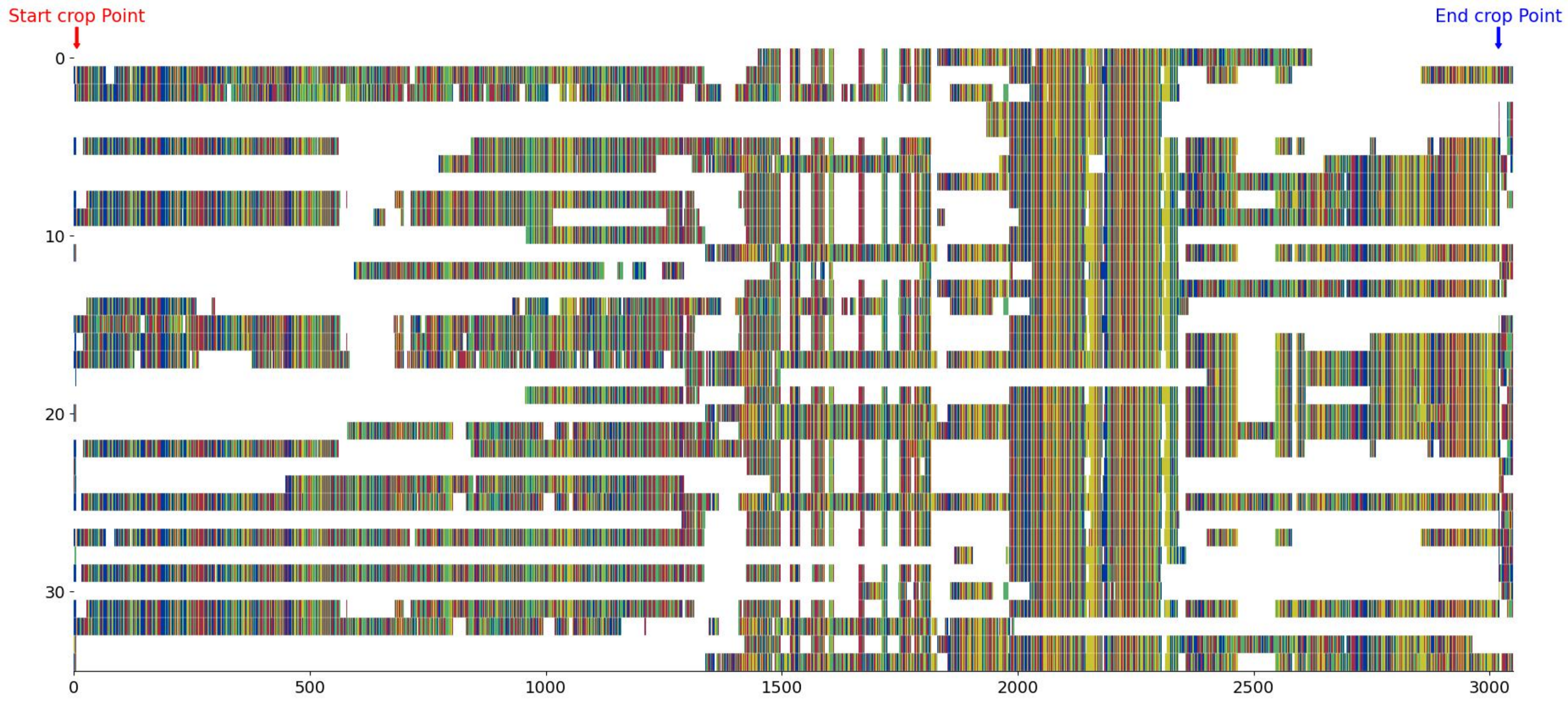


End crop Point



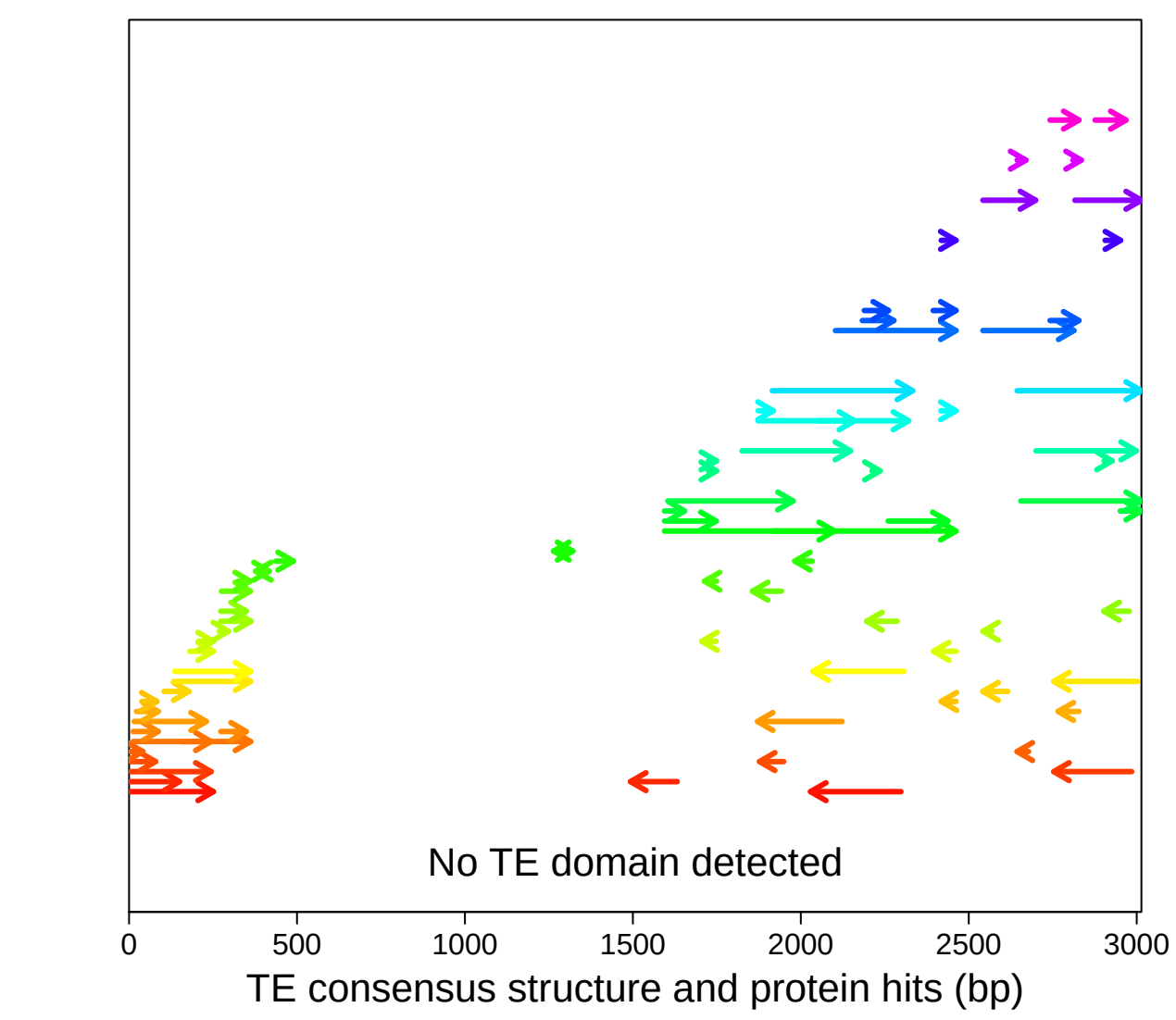
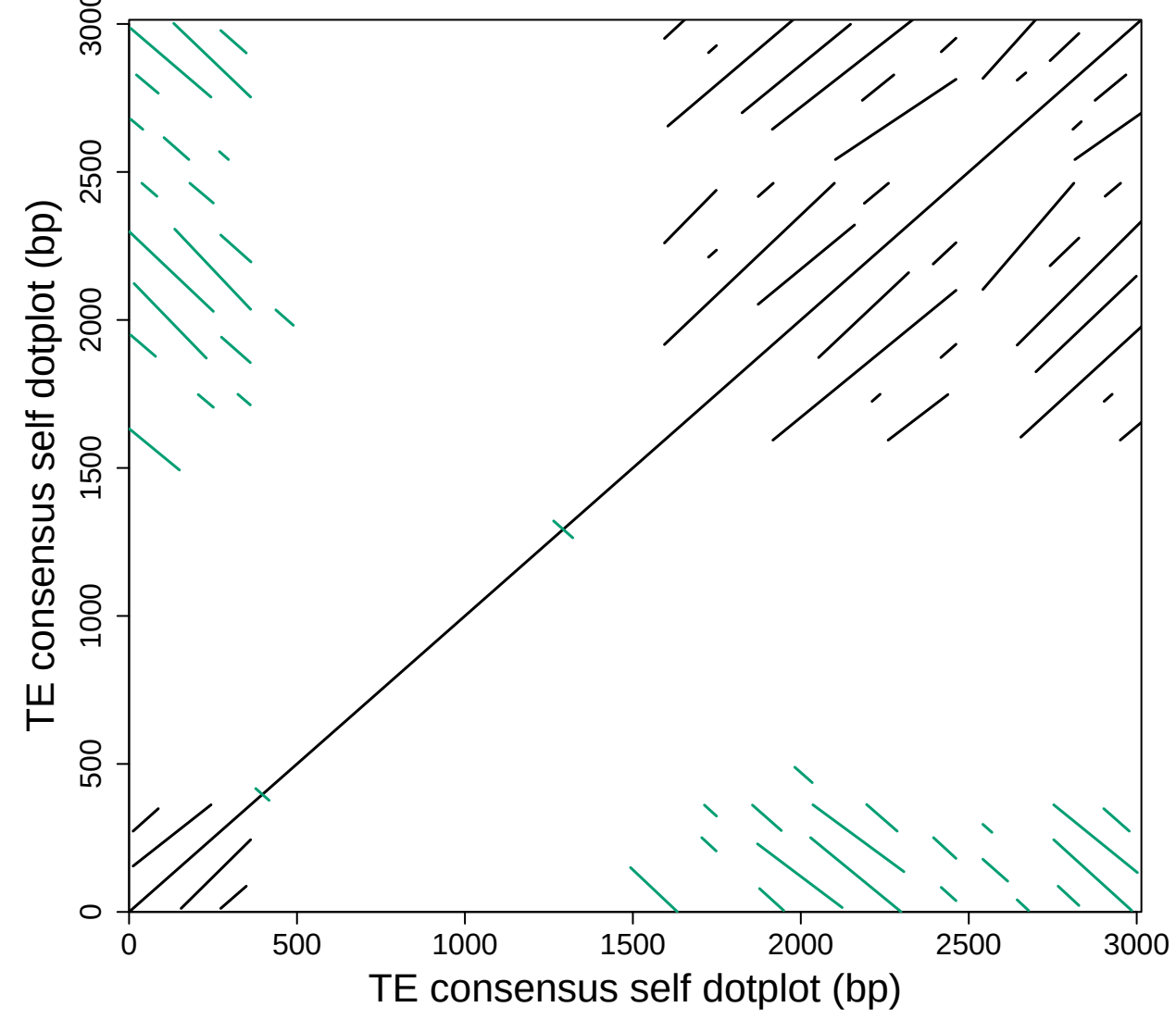
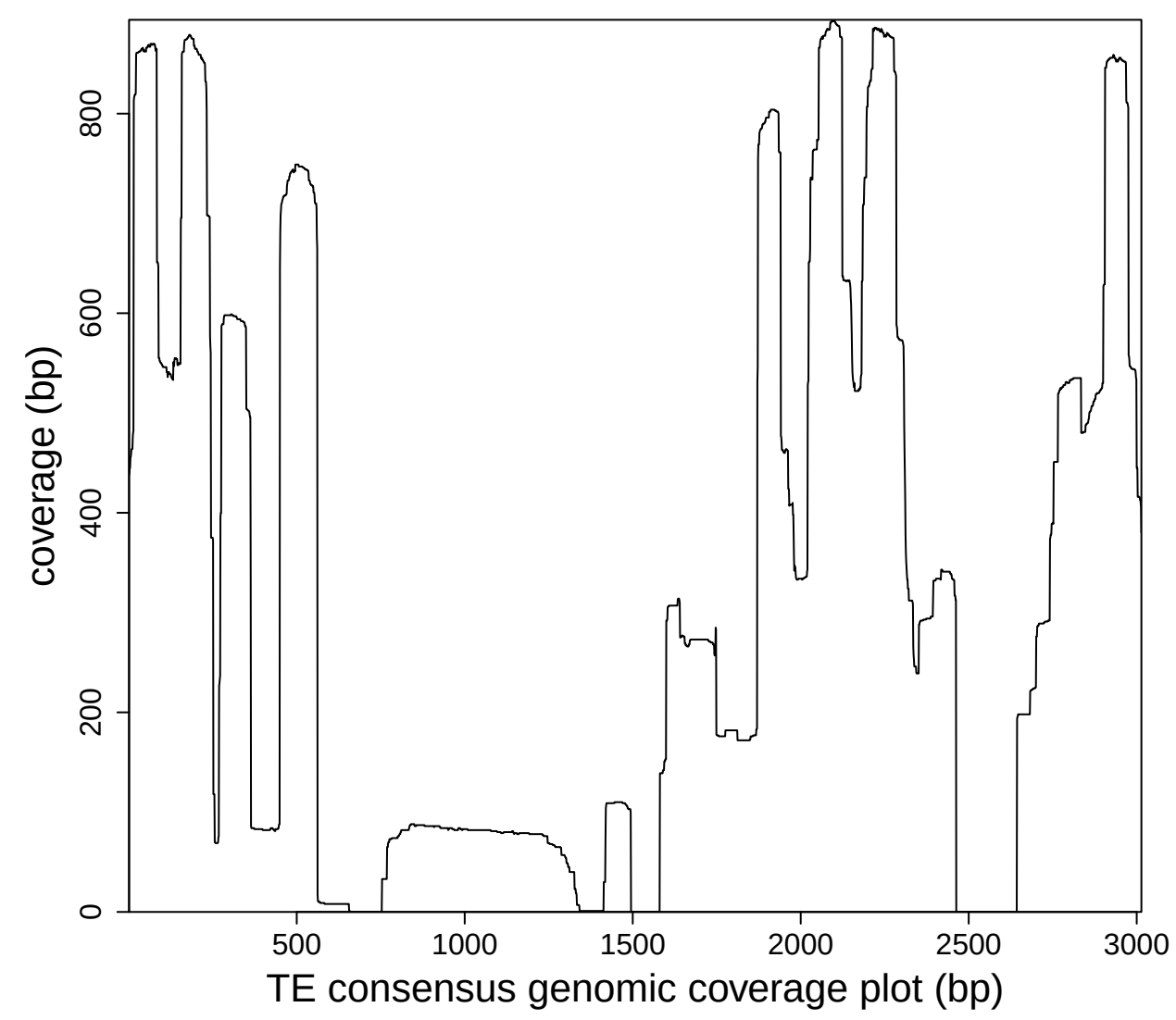
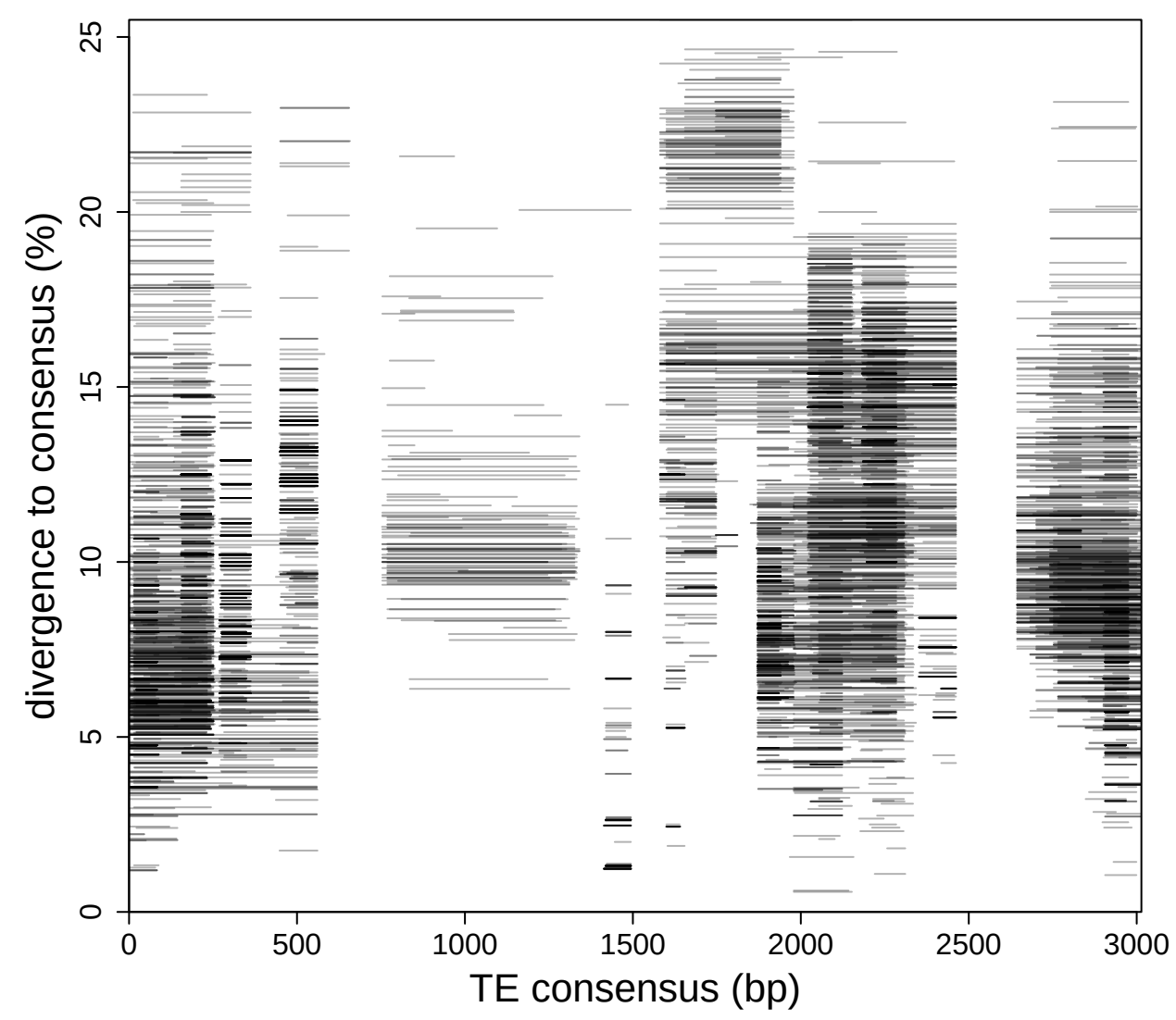
MSA length = 3014





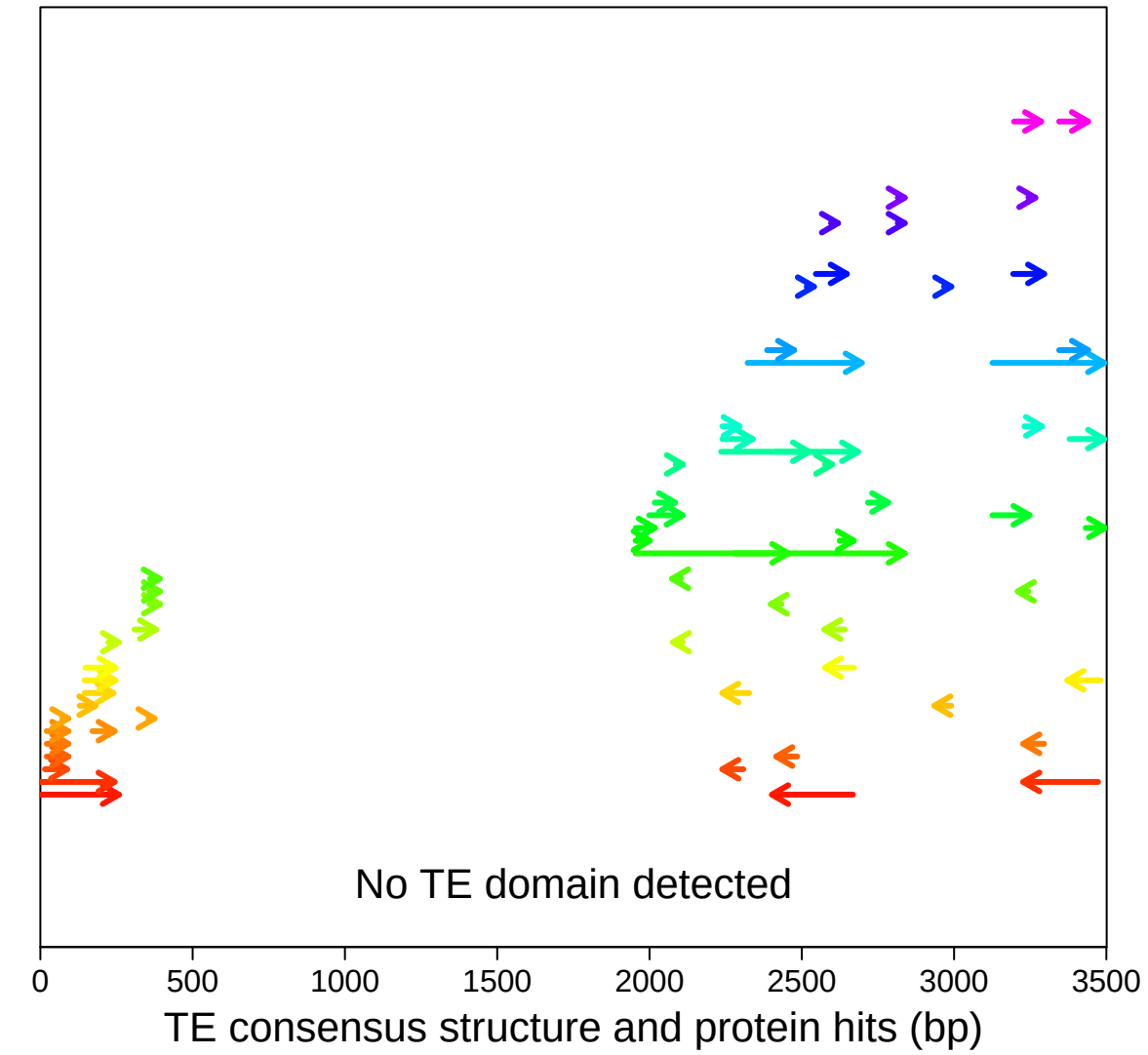
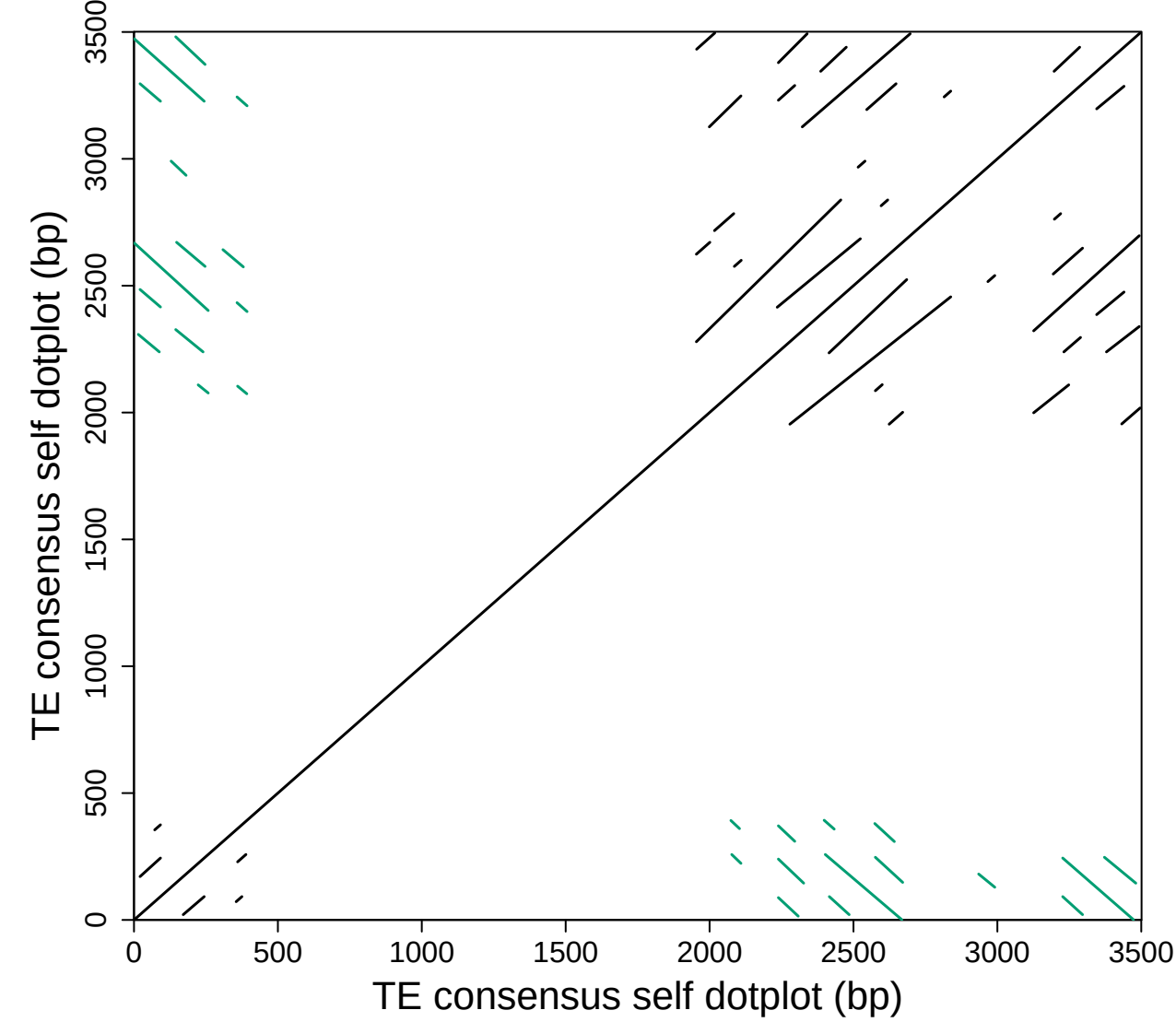
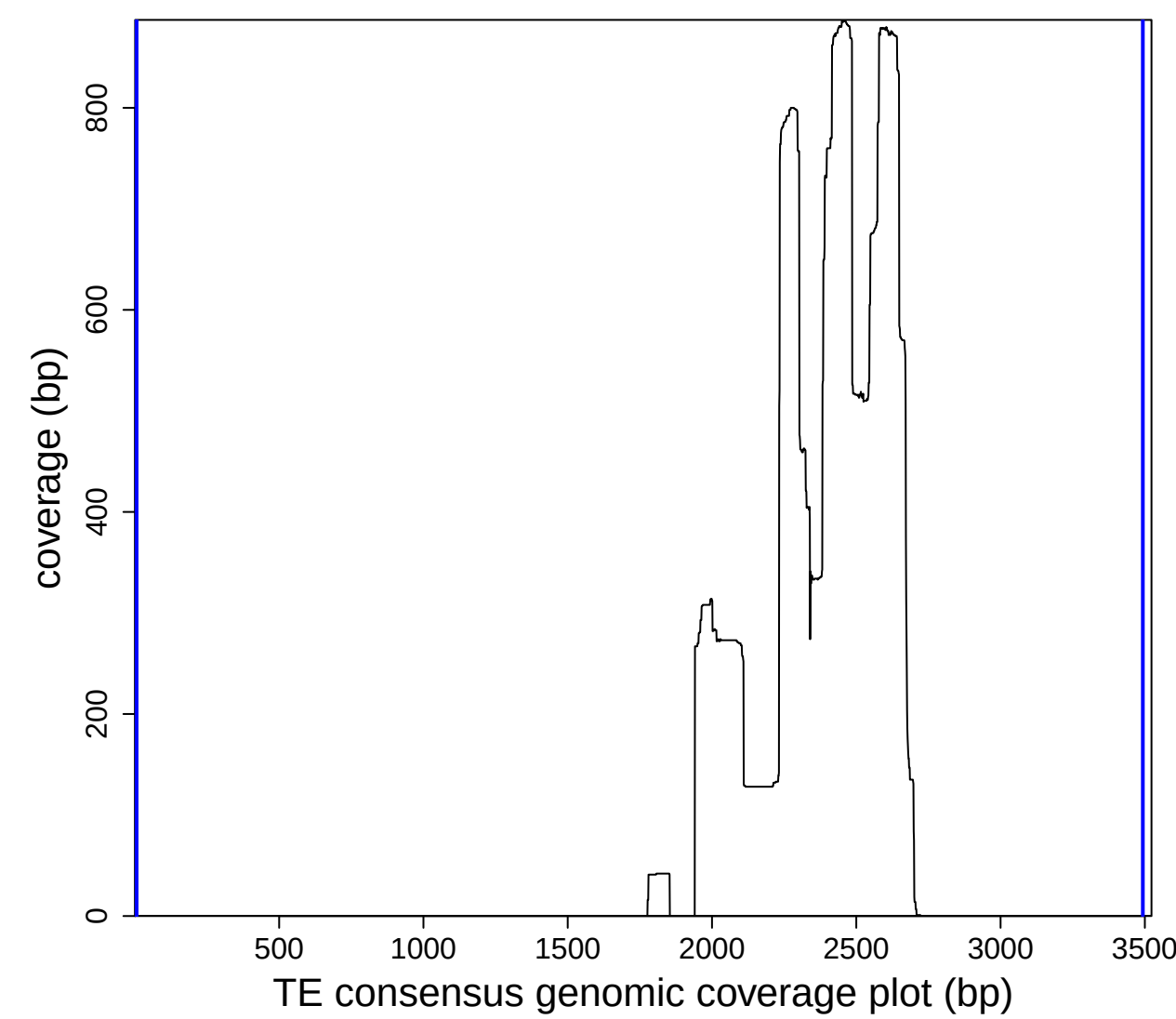
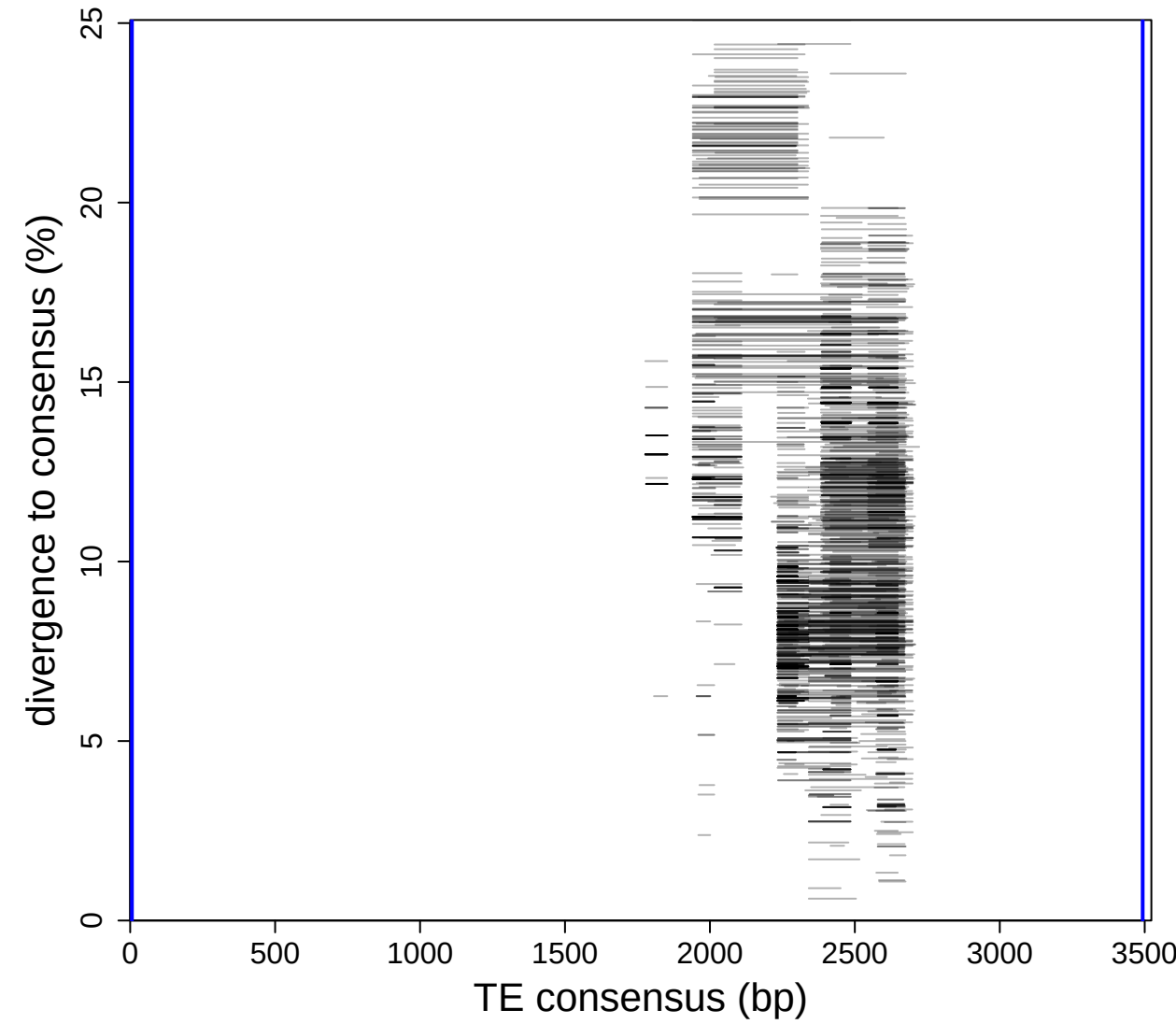
g_1.bed fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa a
size: 3014bp; fragments: 5810; full length: 0 (>=2712.6bp)

After TEtrimmer 3014 bp



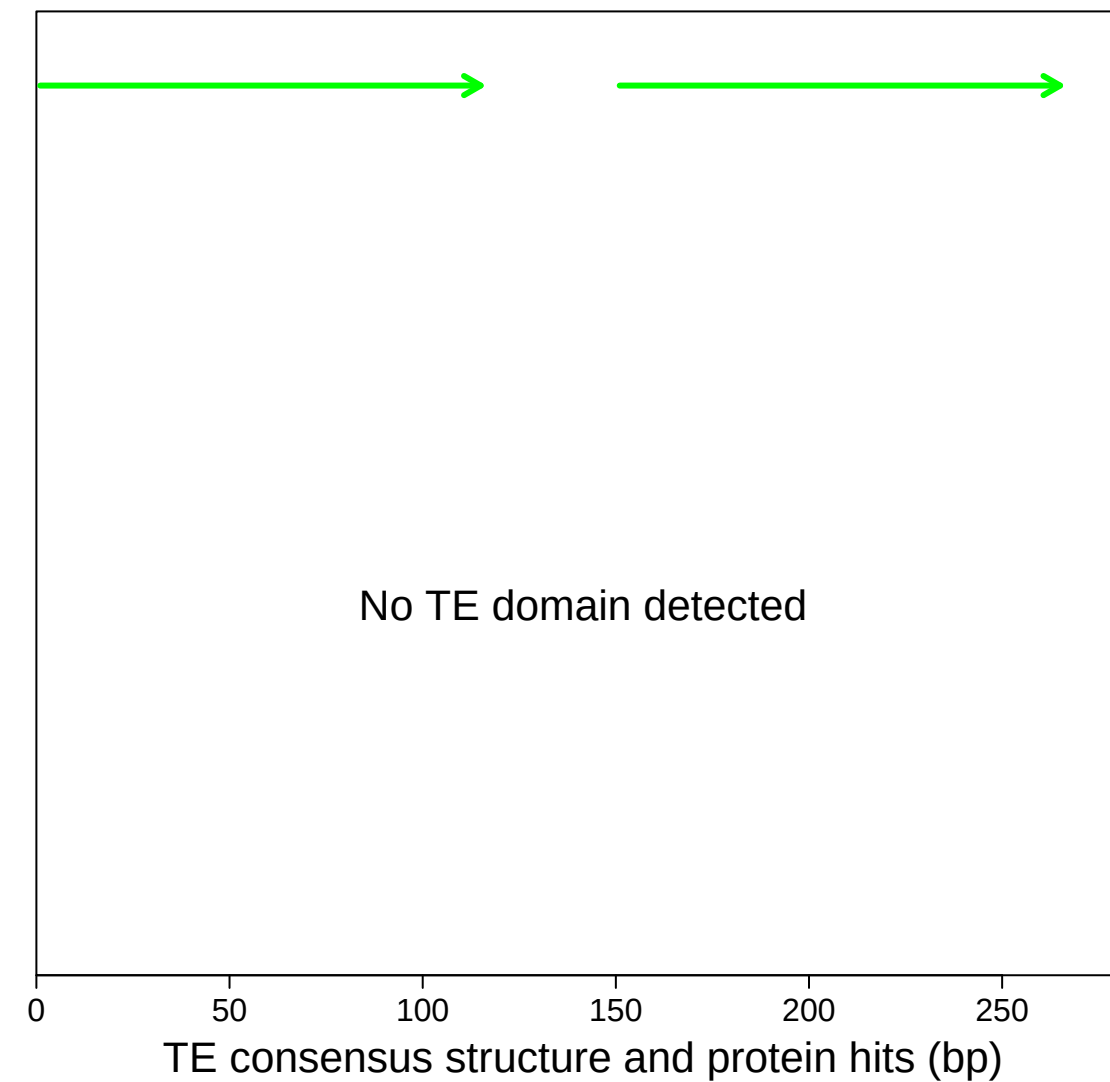
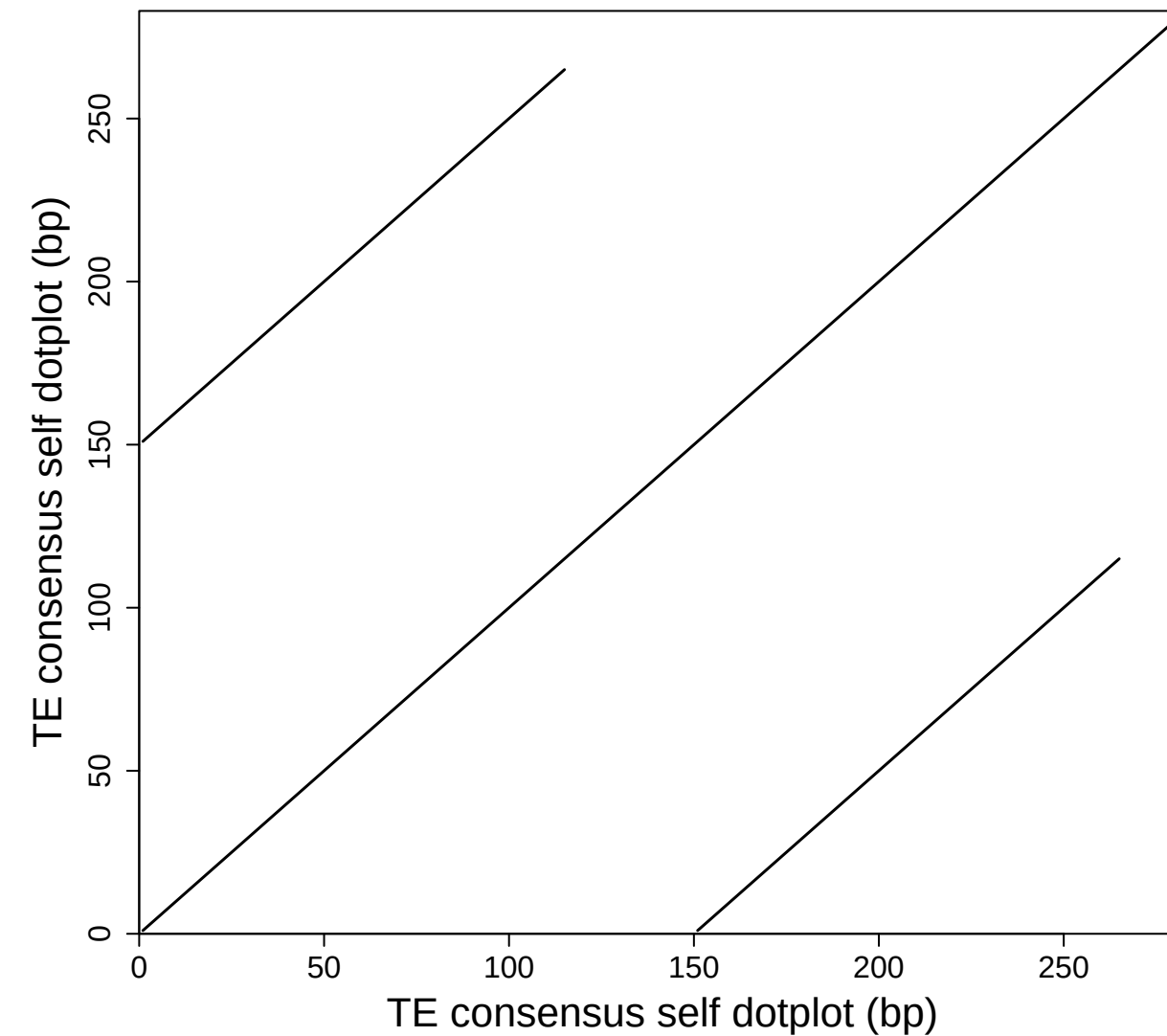
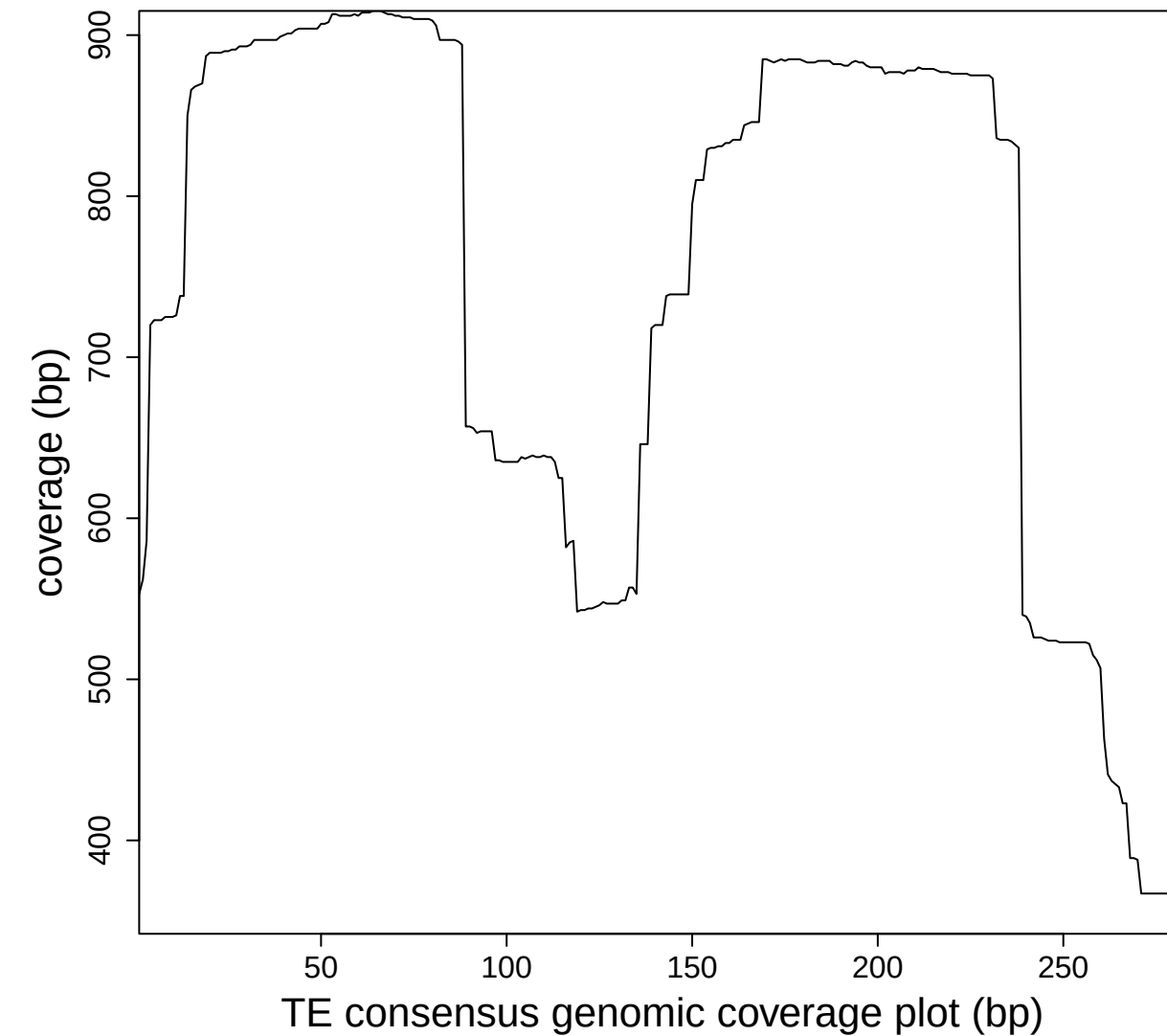
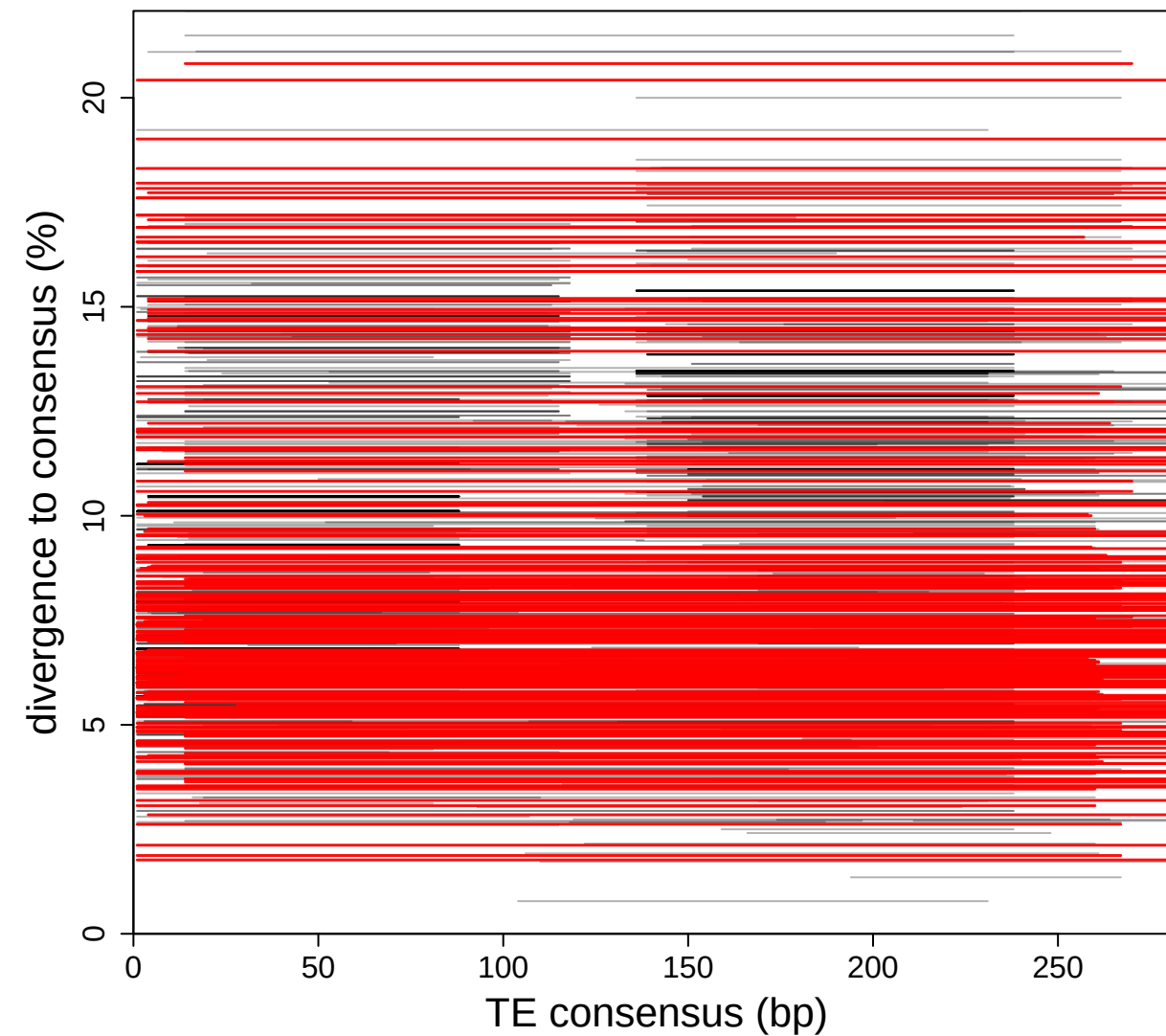
f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 3523bp; fragments: 2144; full length: 0 (>=3170.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

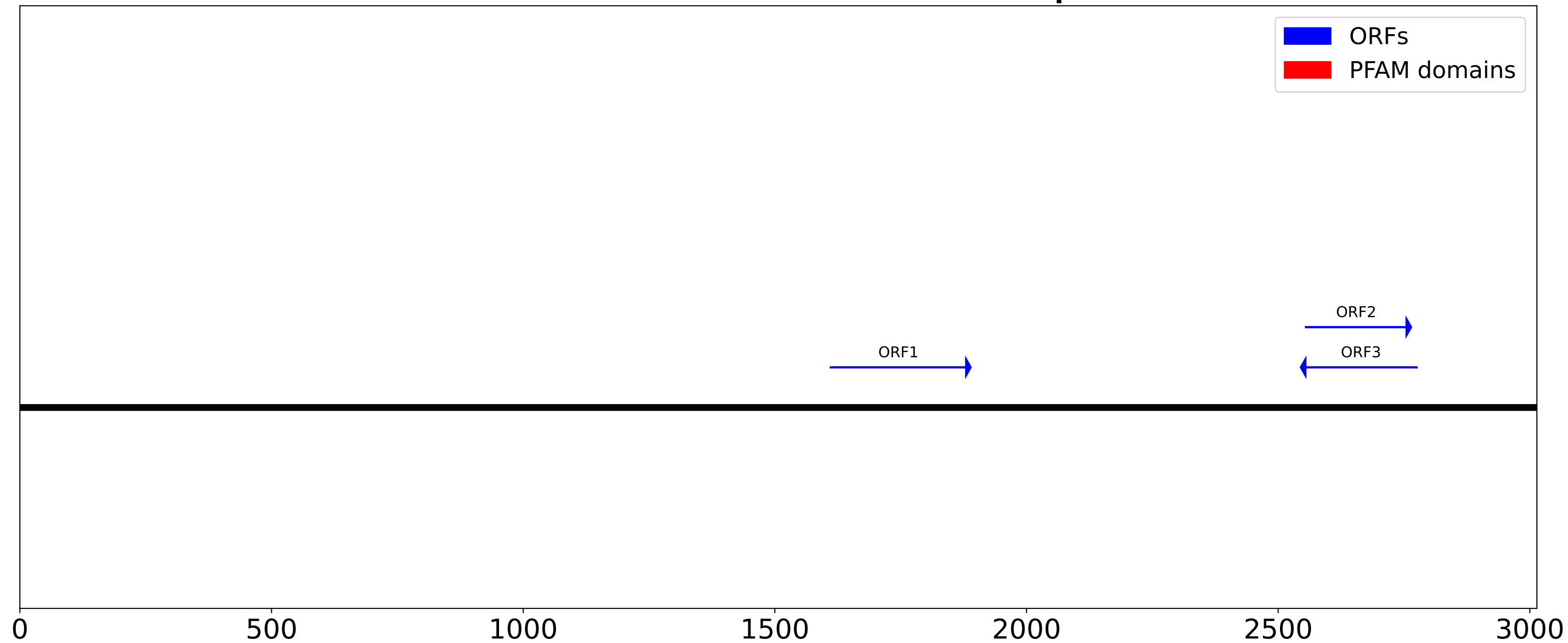


TE: rnd_1_family_917
size: 283bp; fragments: 1302; full length: 339 (≥ 254.7 bp)

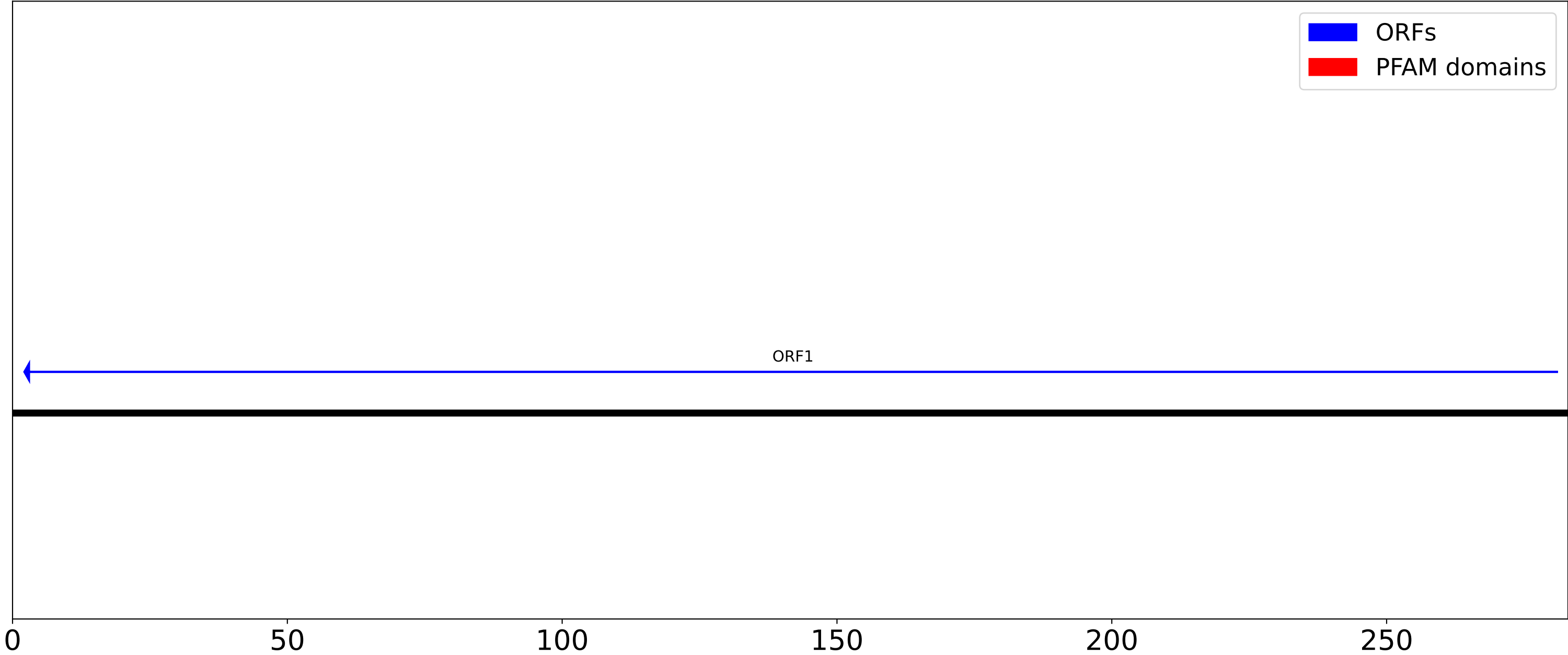
Before TEtrimmer 283 bp



After TEtrimmer ORF and PFAM domain plot



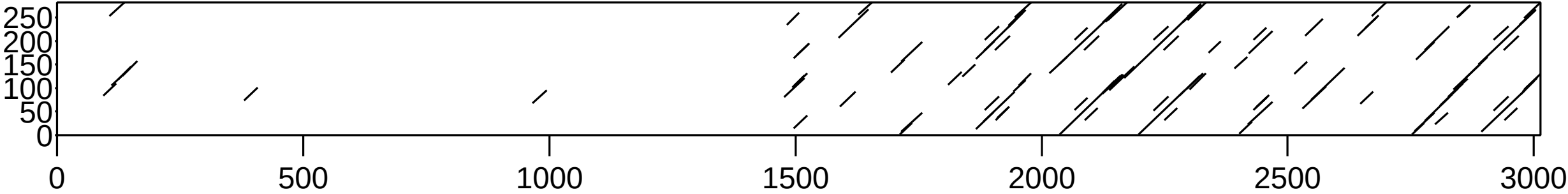
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

ensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)