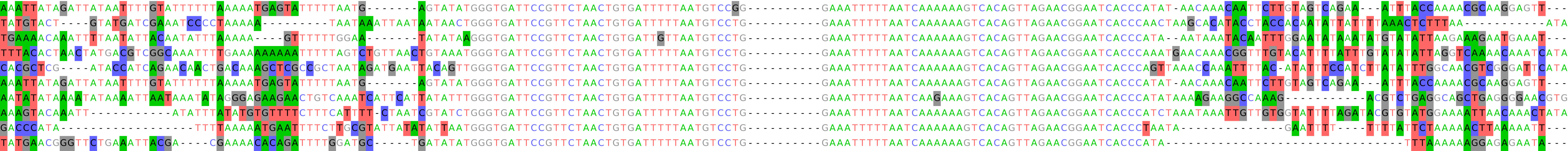


Start Crop Point

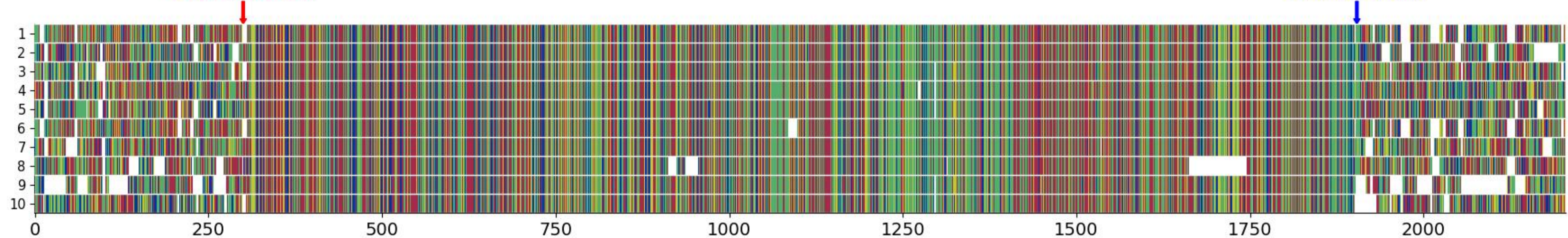
End Crop Point

MSA length = 1605



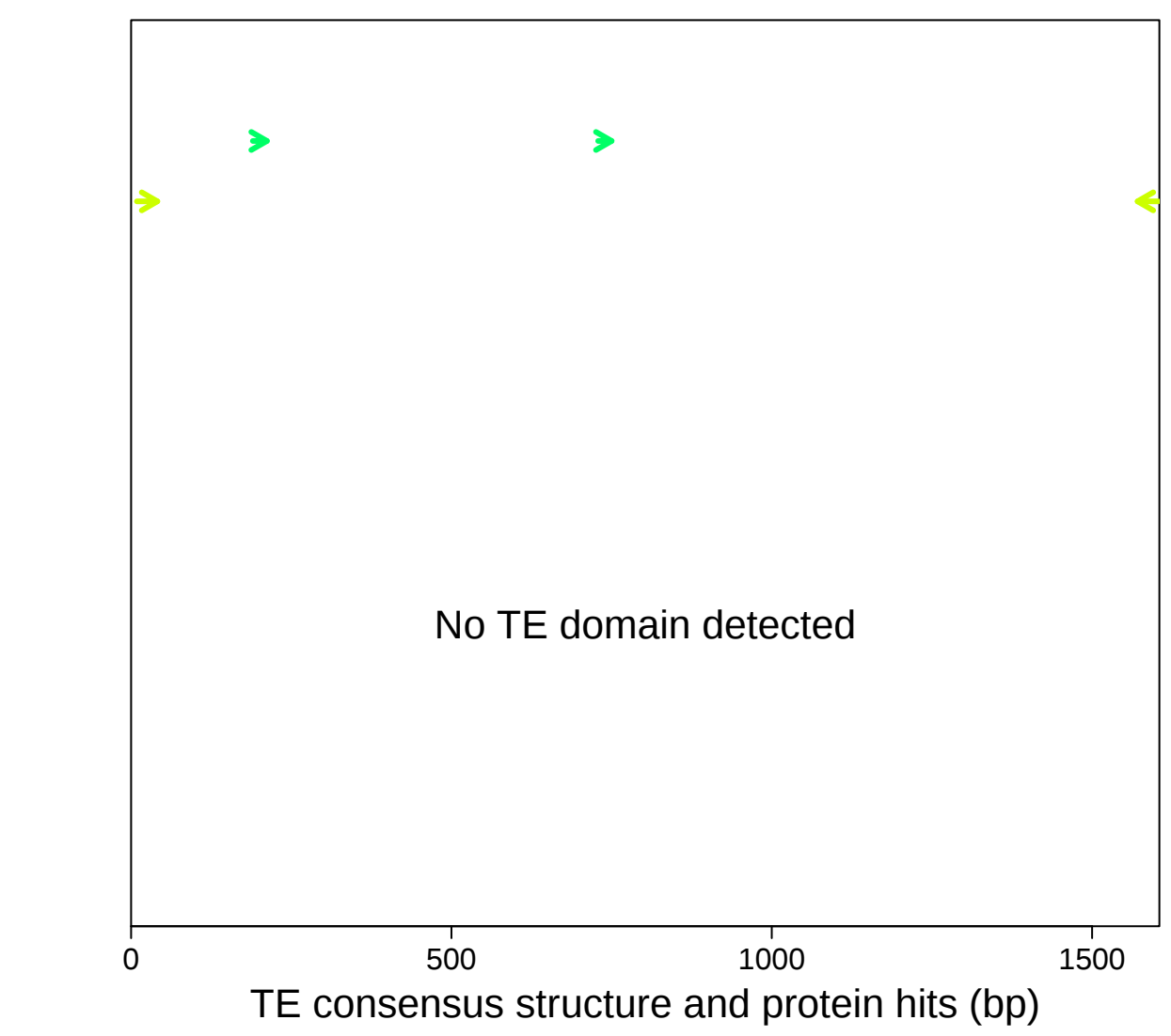
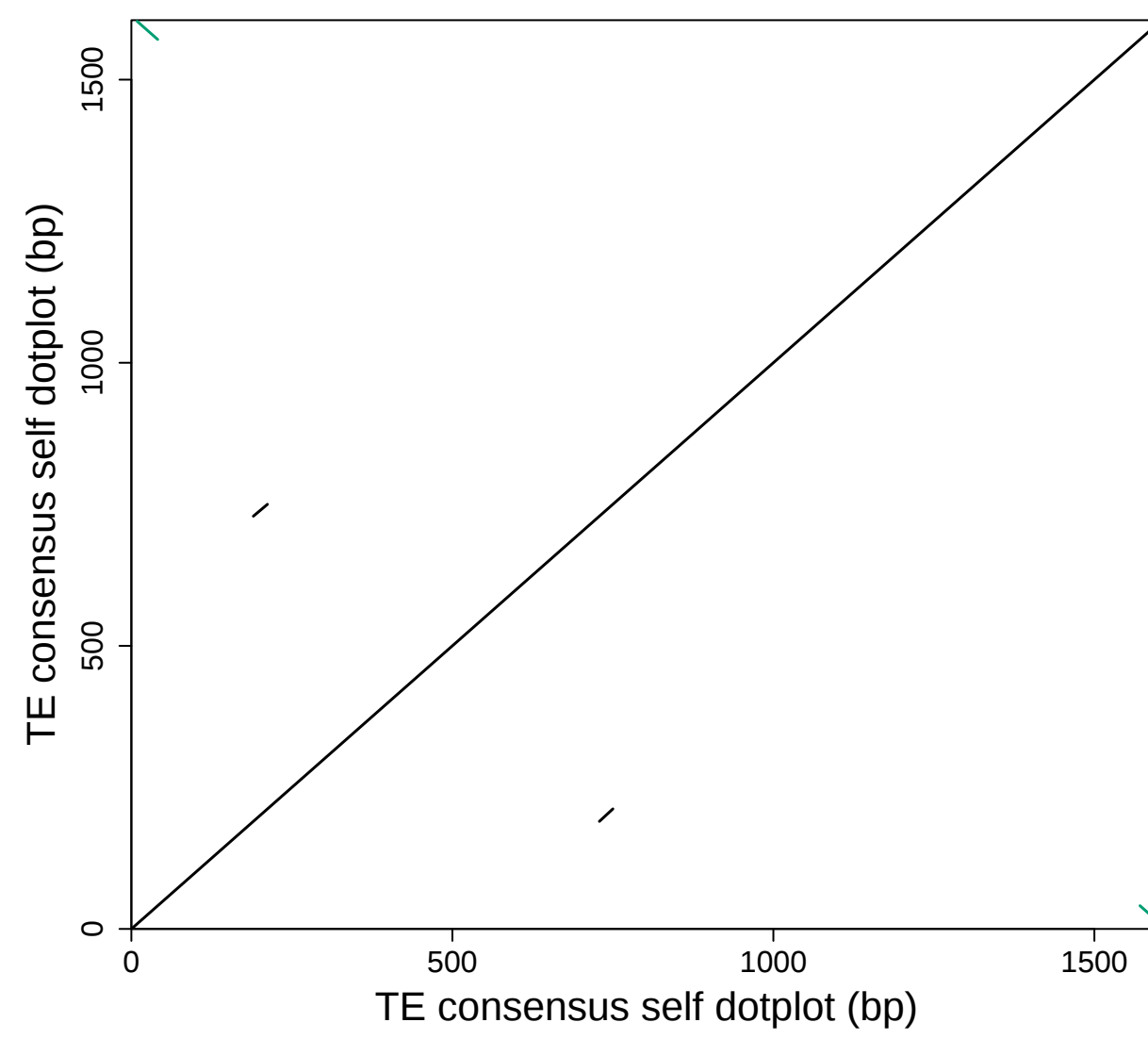
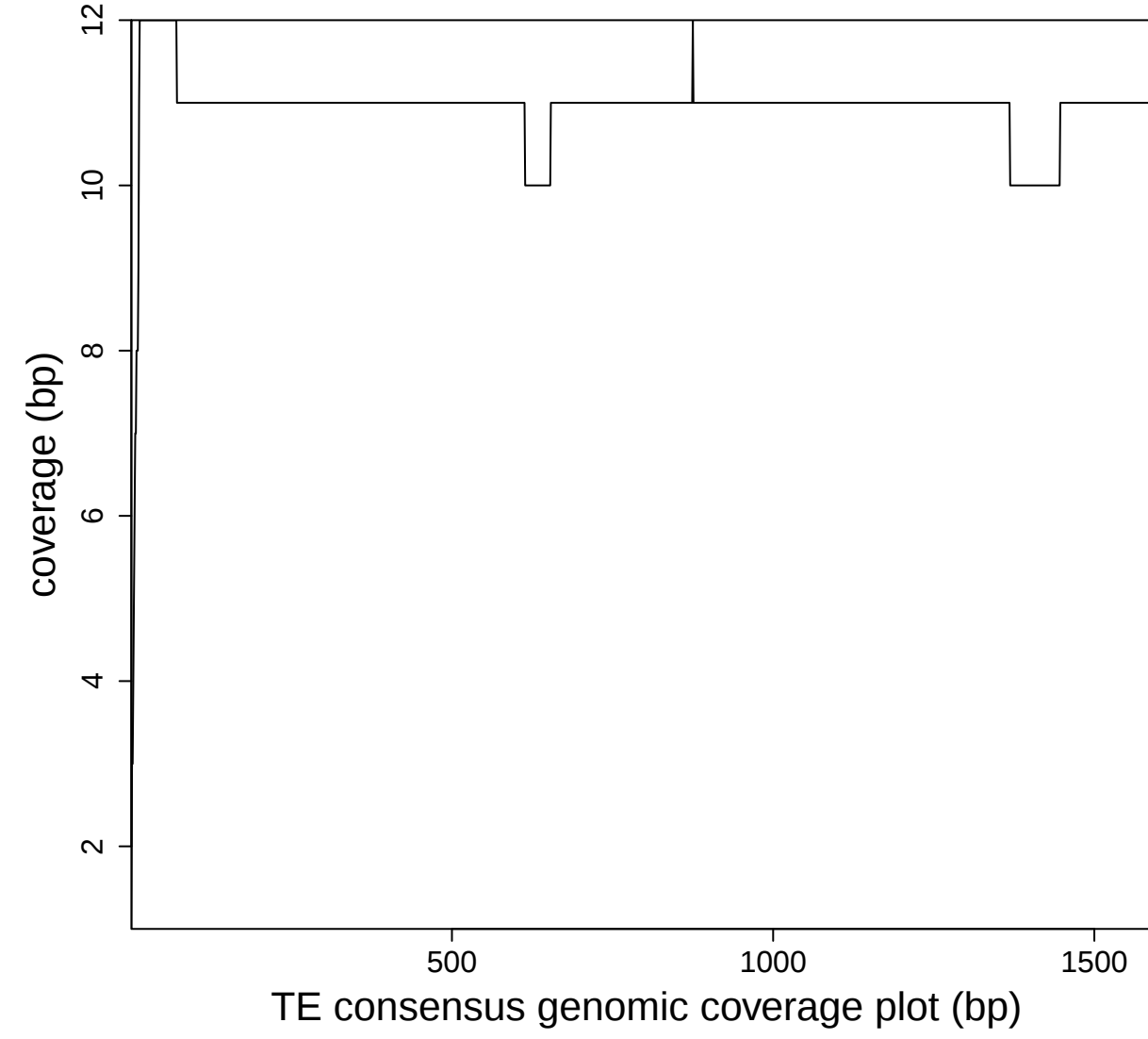
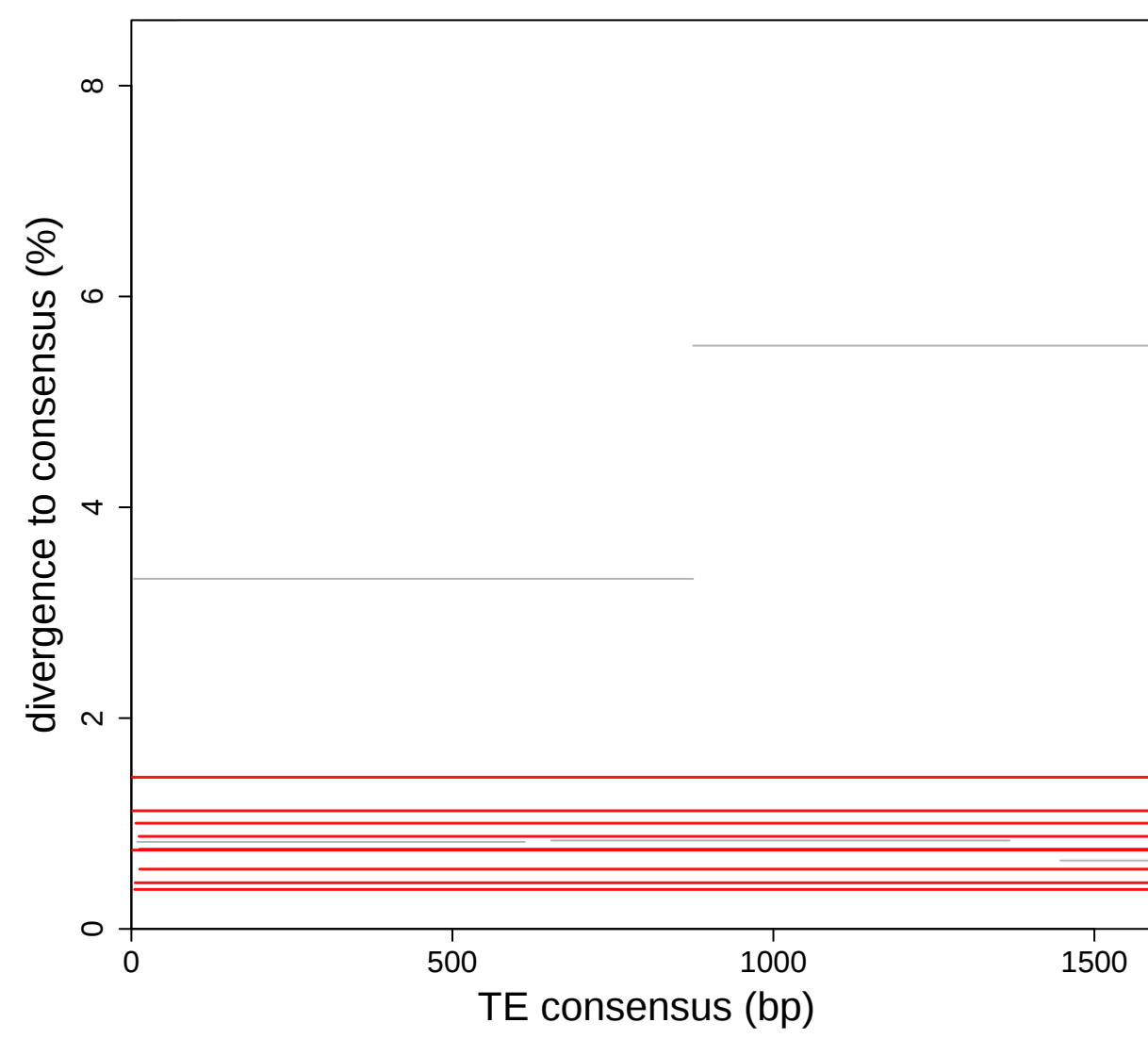
Start crop Point

End crop Point



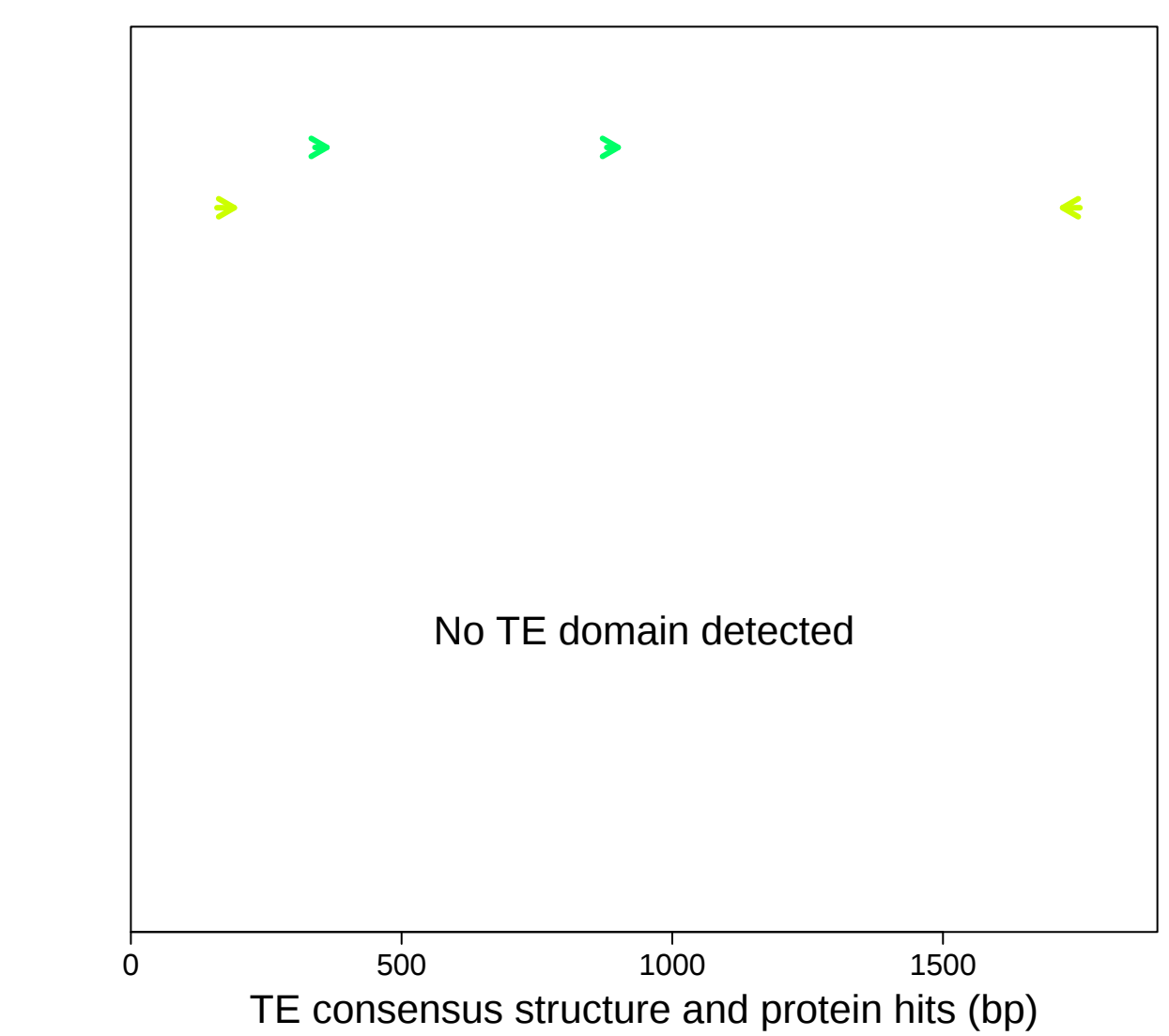
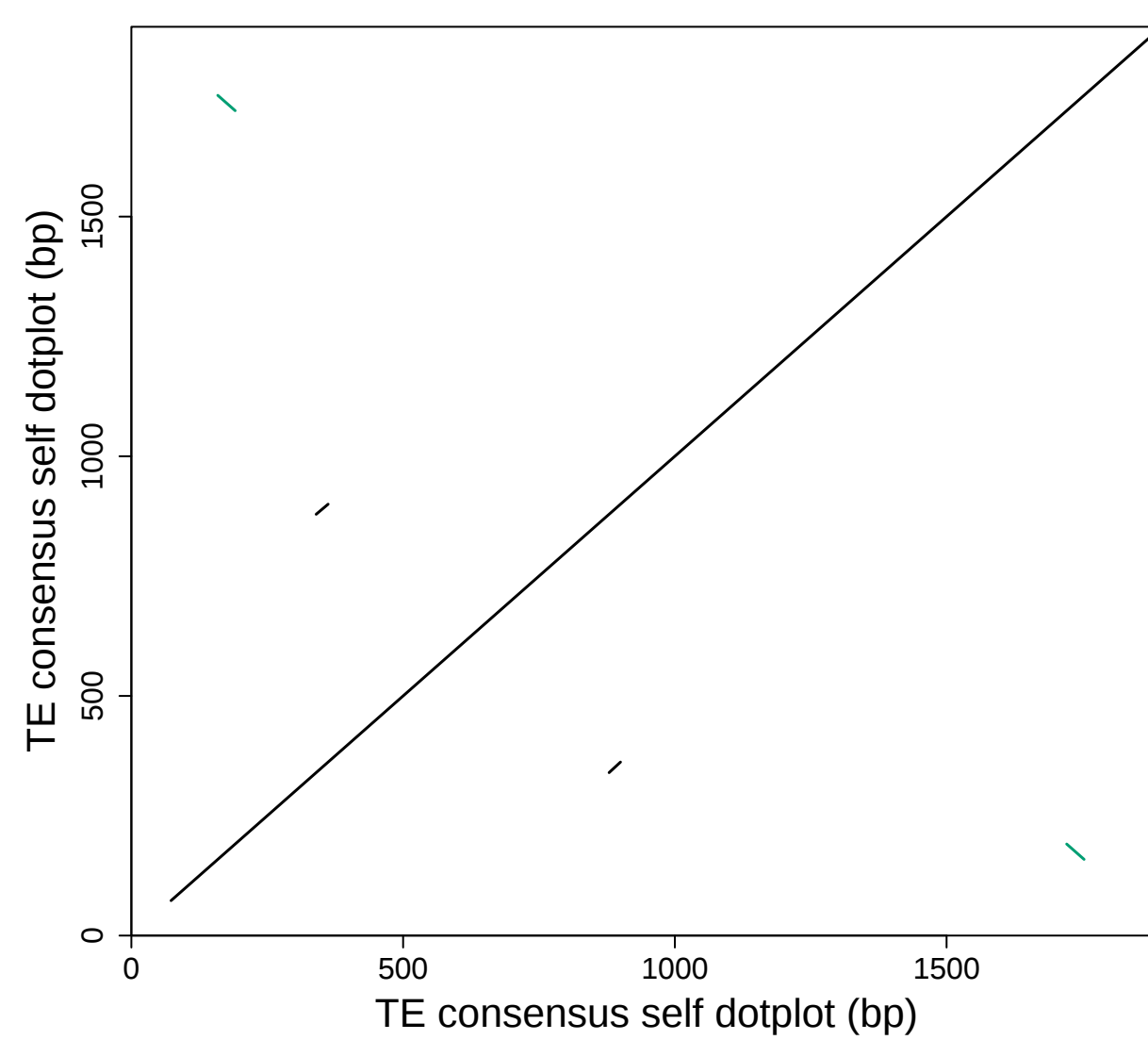
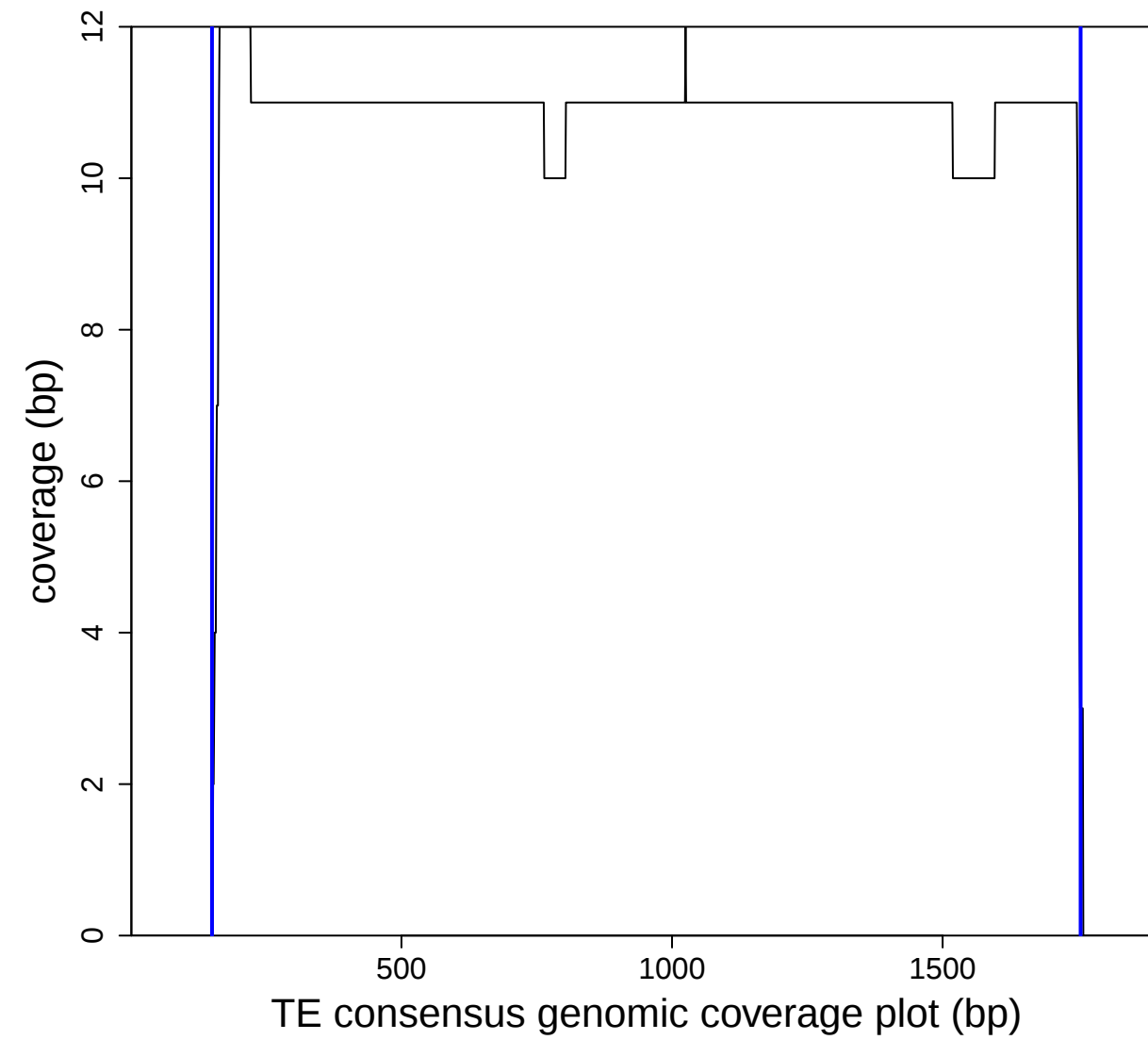
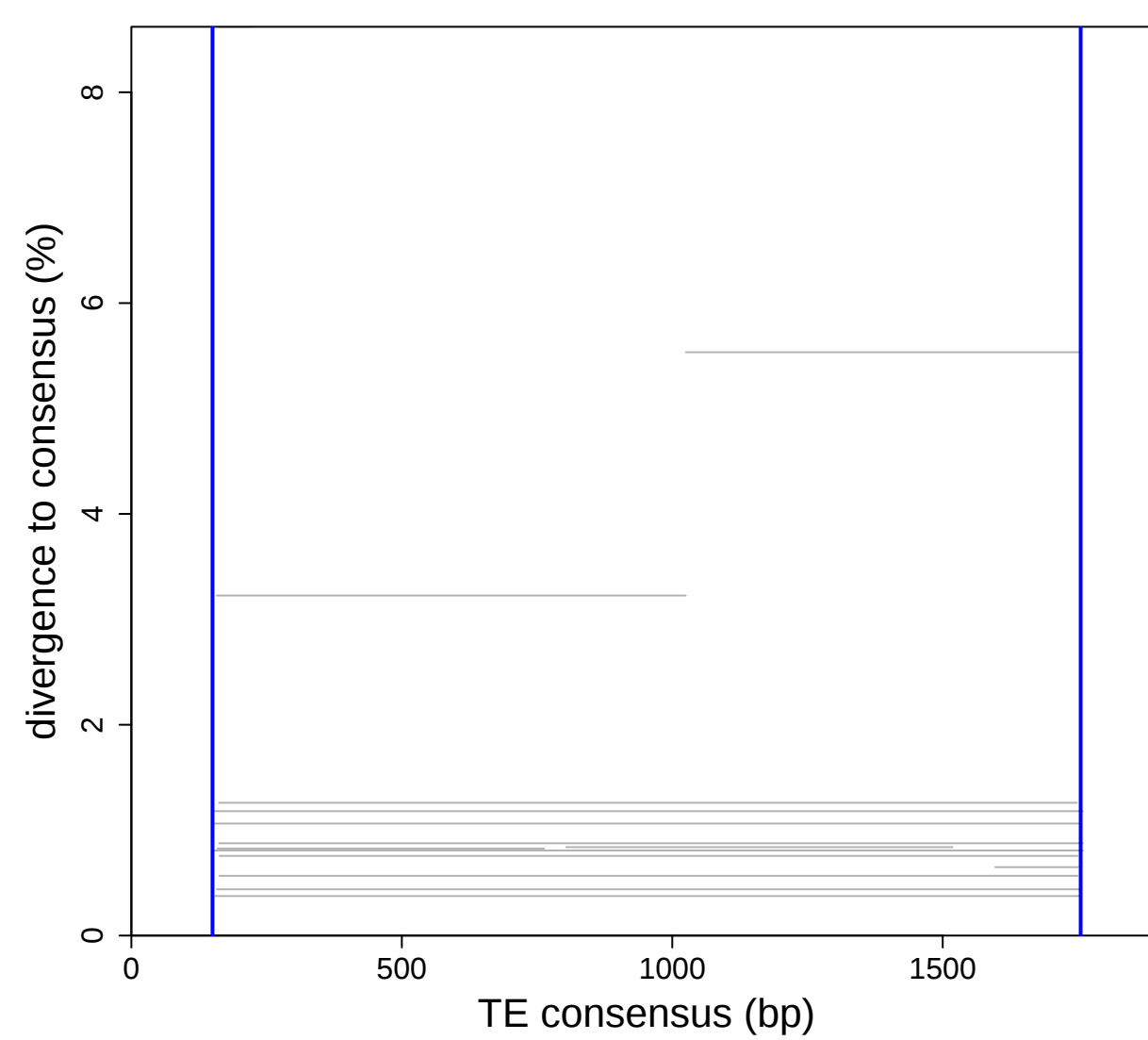
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 1605bp; fragments: 15; full length: 9 (≥ 1444.5 bp)

After TEtrimmer 1605 bp

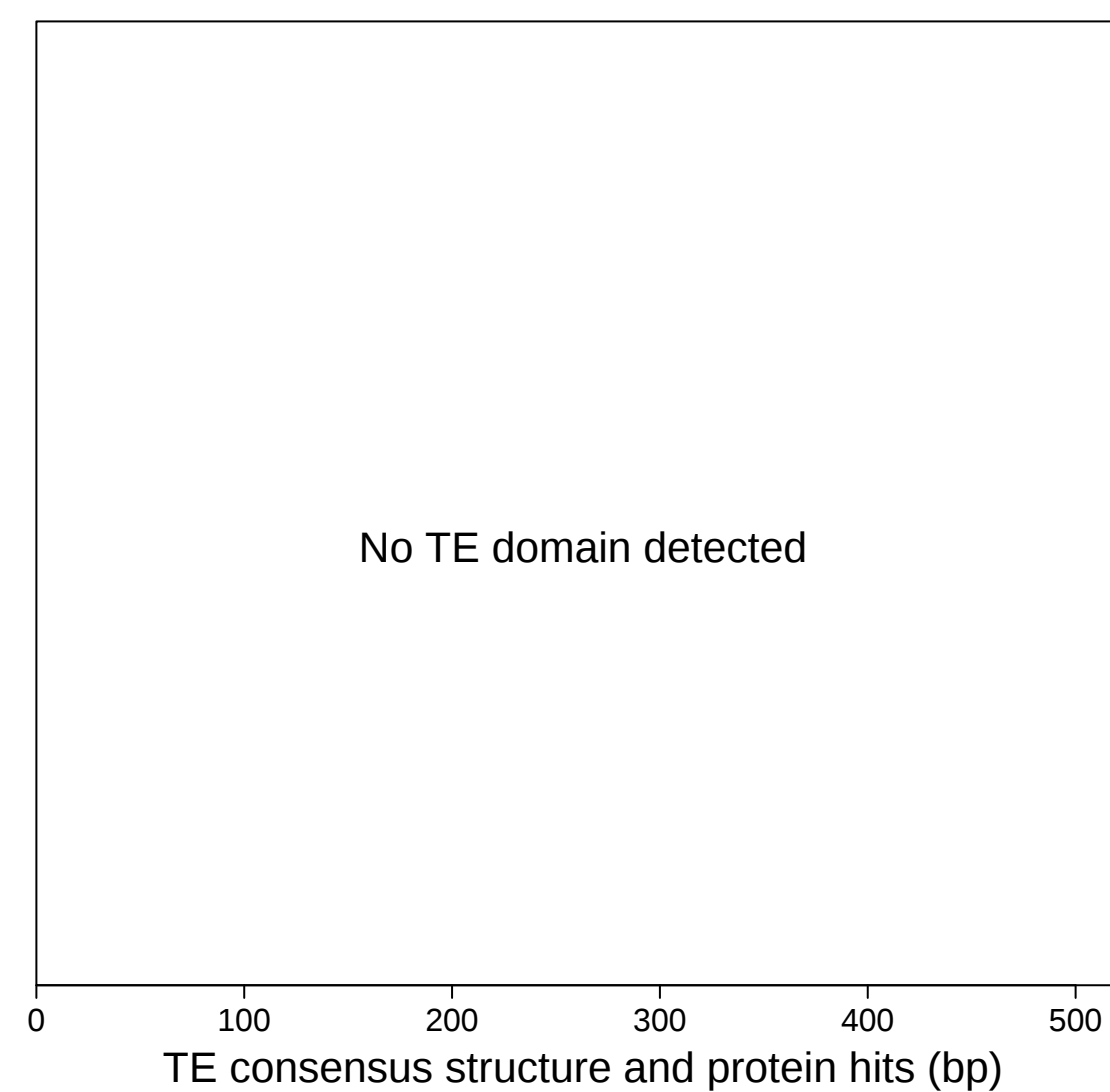
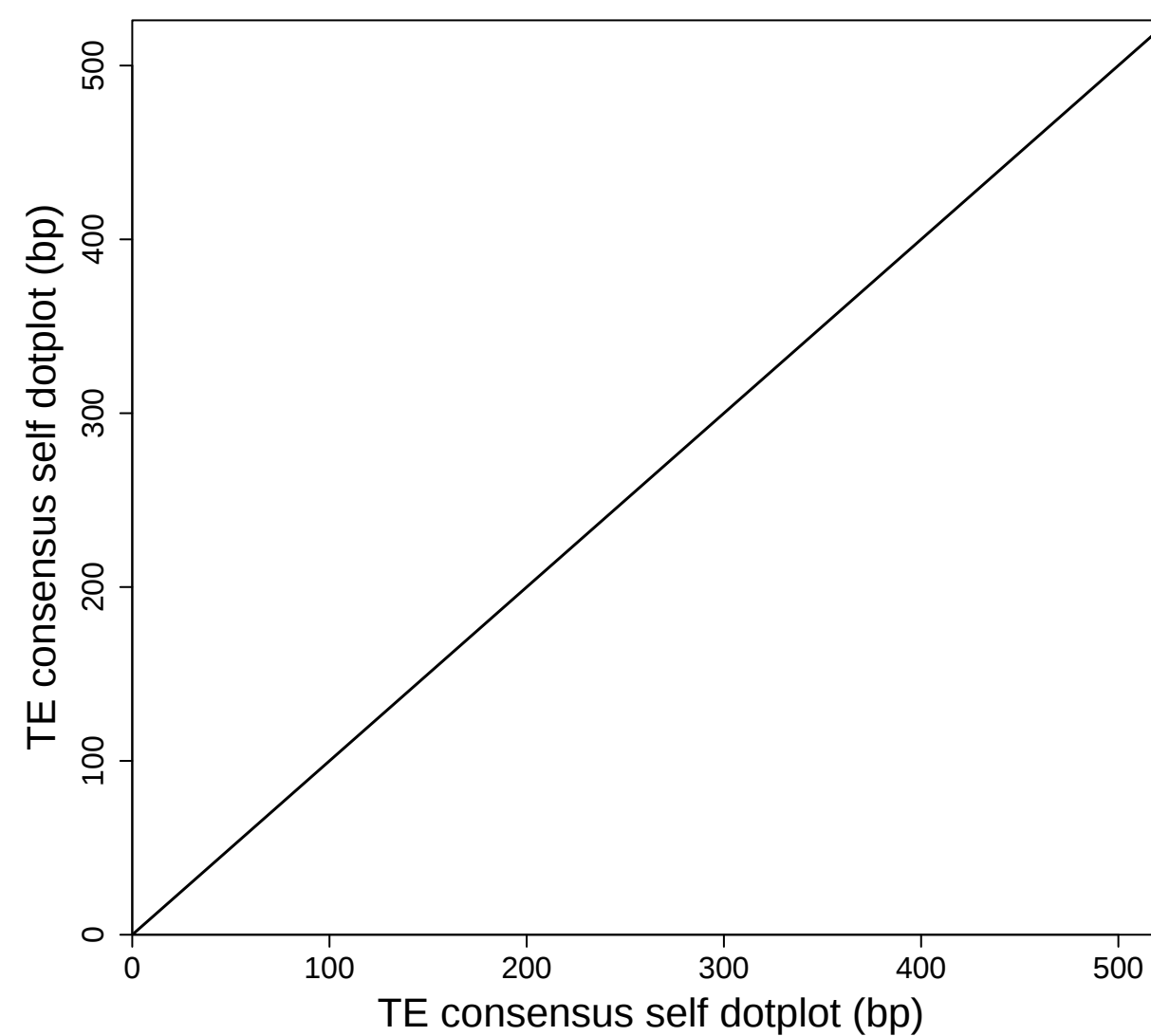
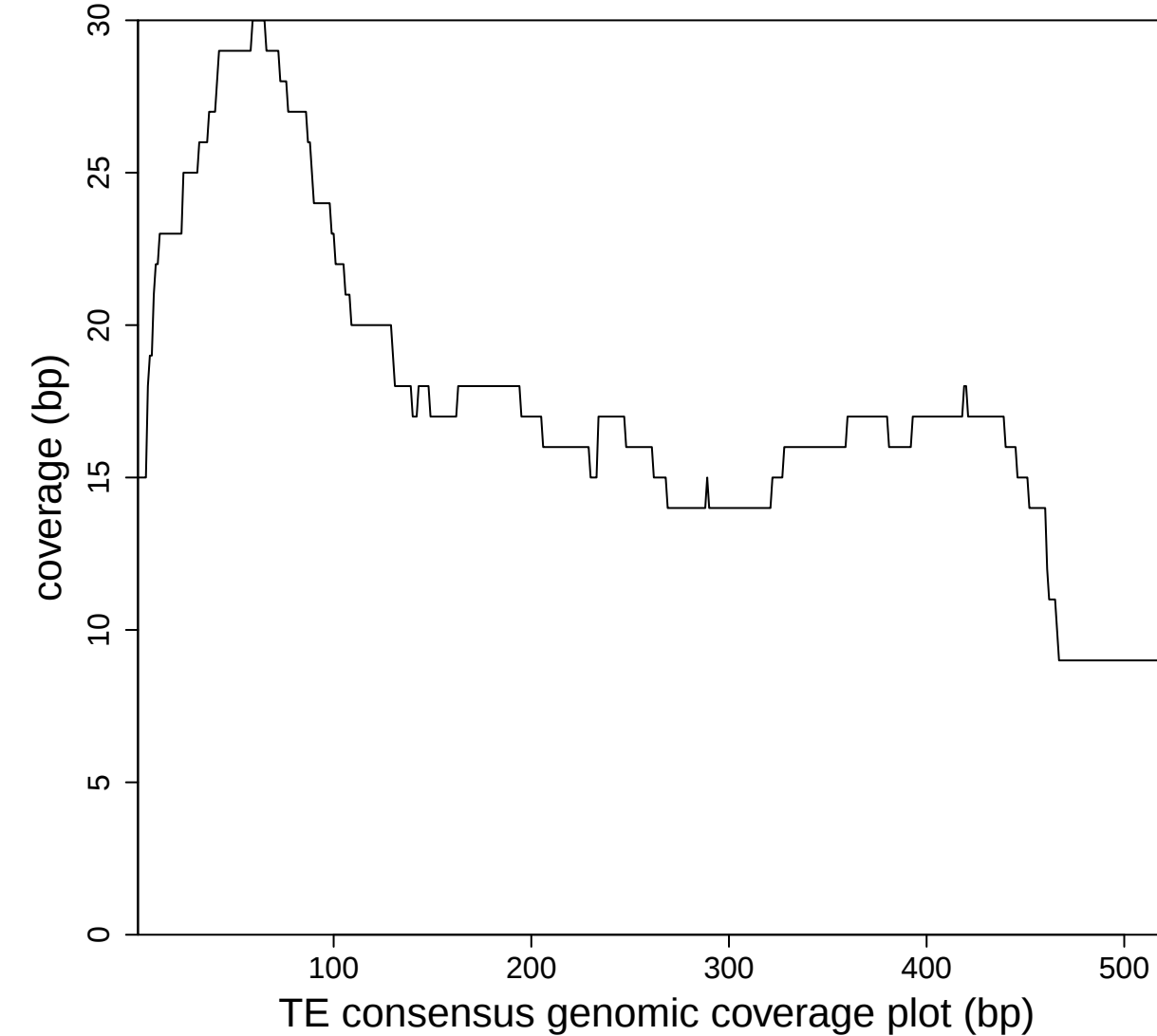
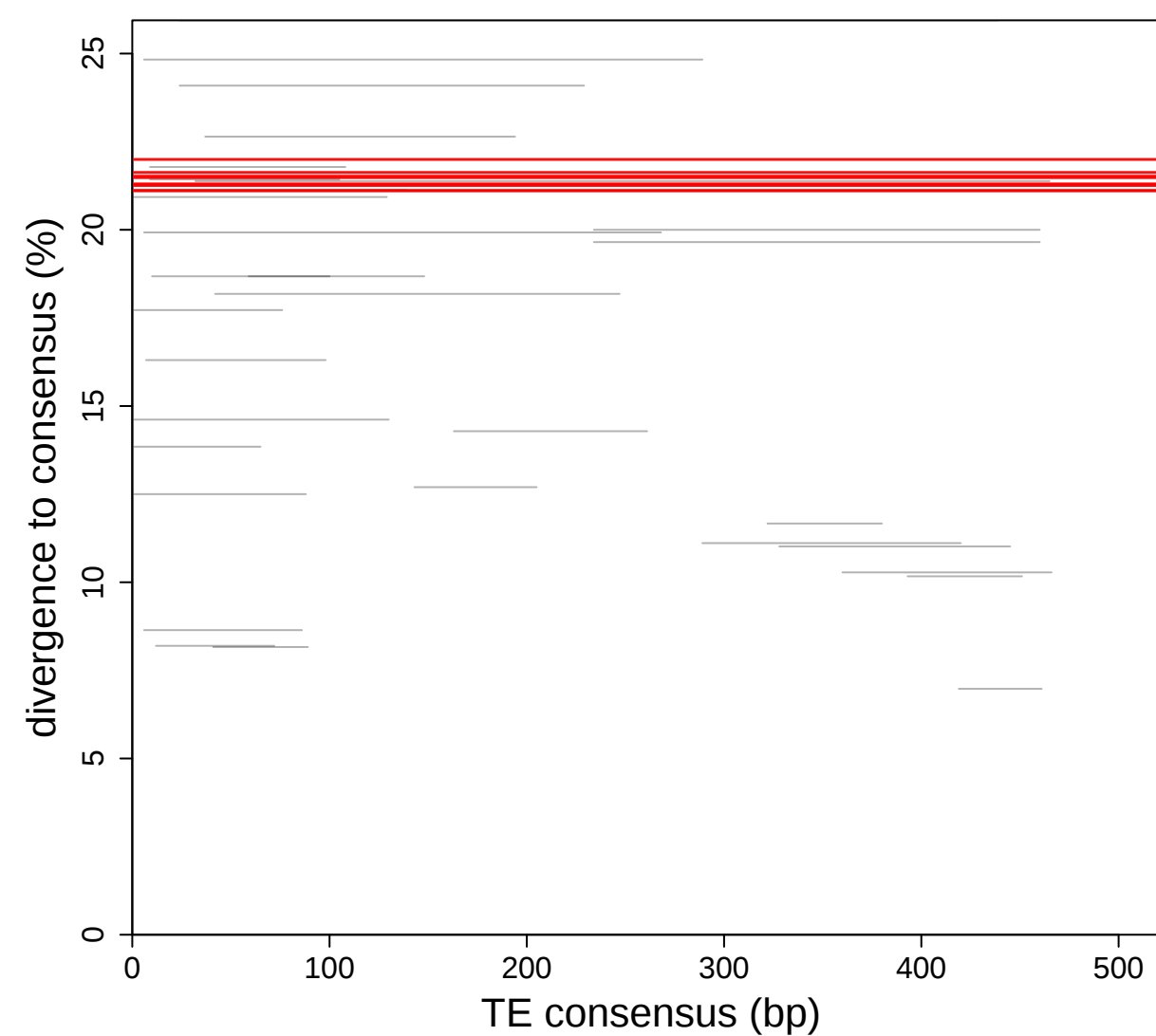


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 1905bp; fragments: 15; full length: 0 (>=1714.5bp)

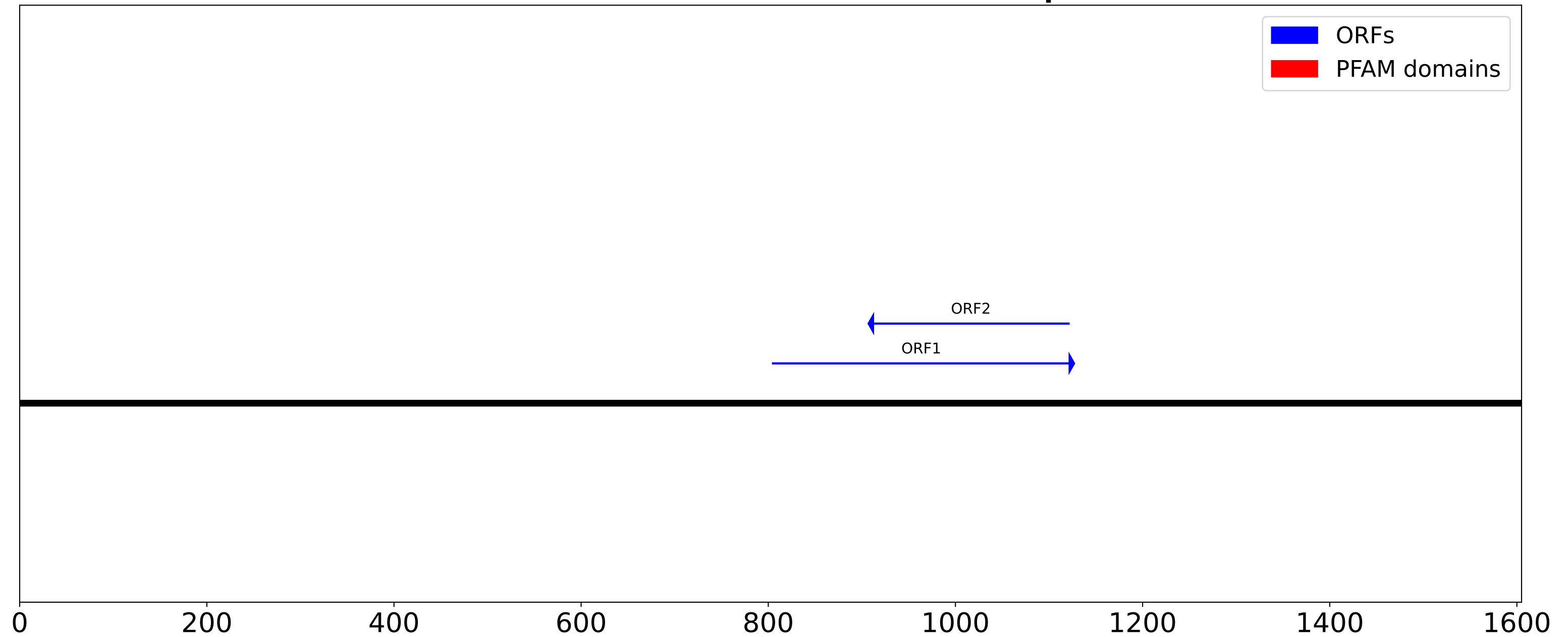
After TEtrimmer Extended plot Blue lines are boundaries



TE: rnd_4_family_184
size: 526bp; fragments: 40; full length: 9 (≥ 473.4 bp)



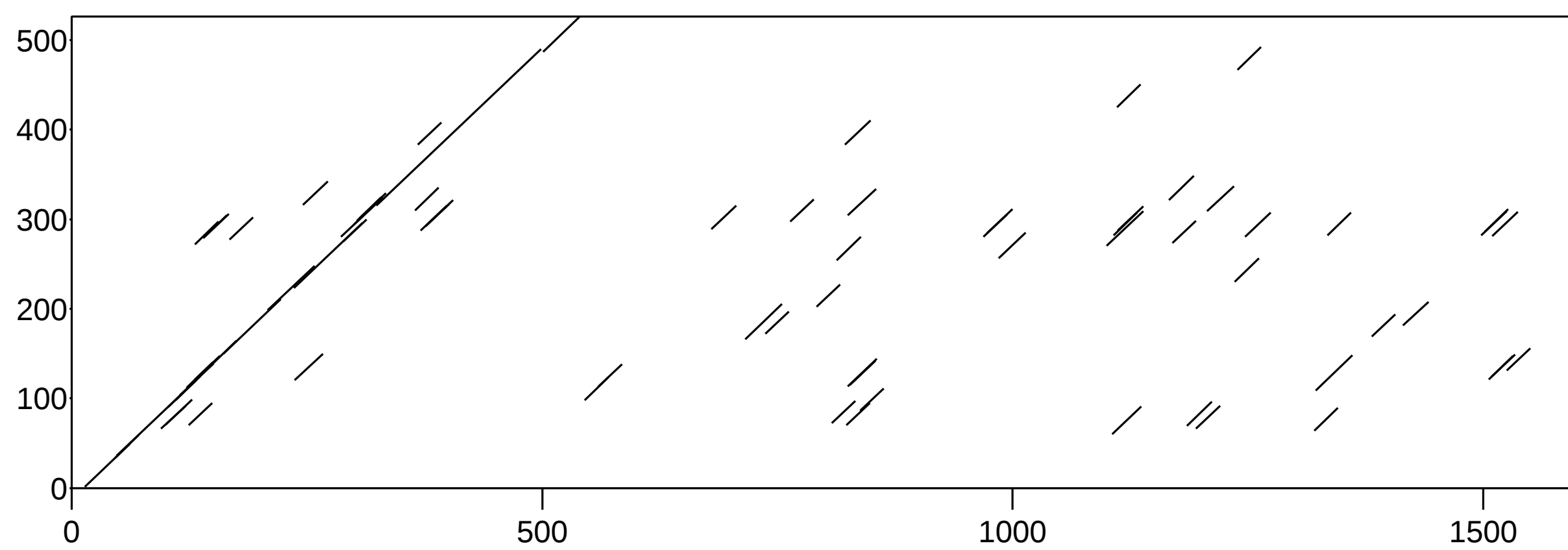
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)