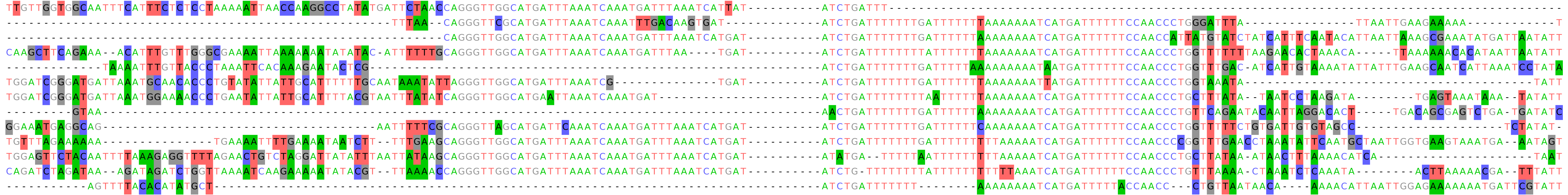


Start crop Point

End crop Point

MSA length = 1266



Start crop Point



End crop Point



0 -

10 -

0

250

500

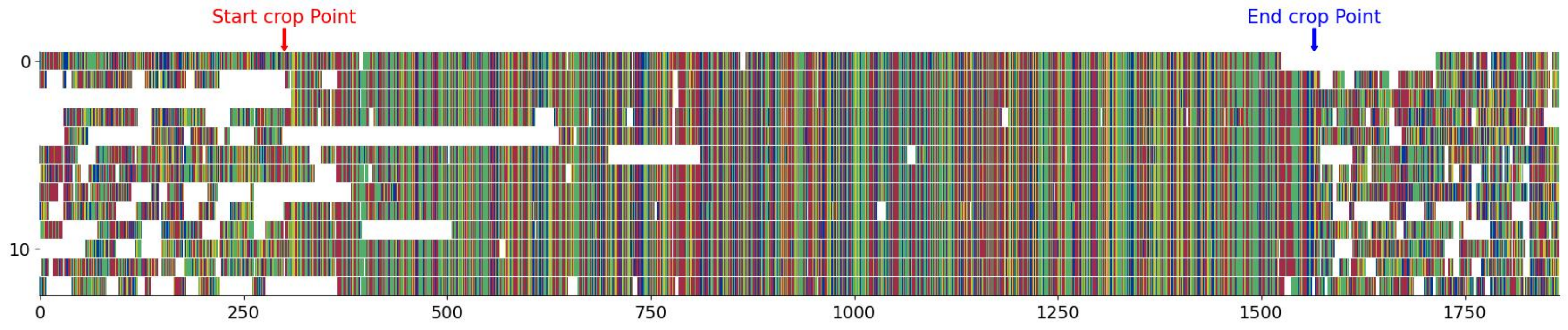
750

1000

1250

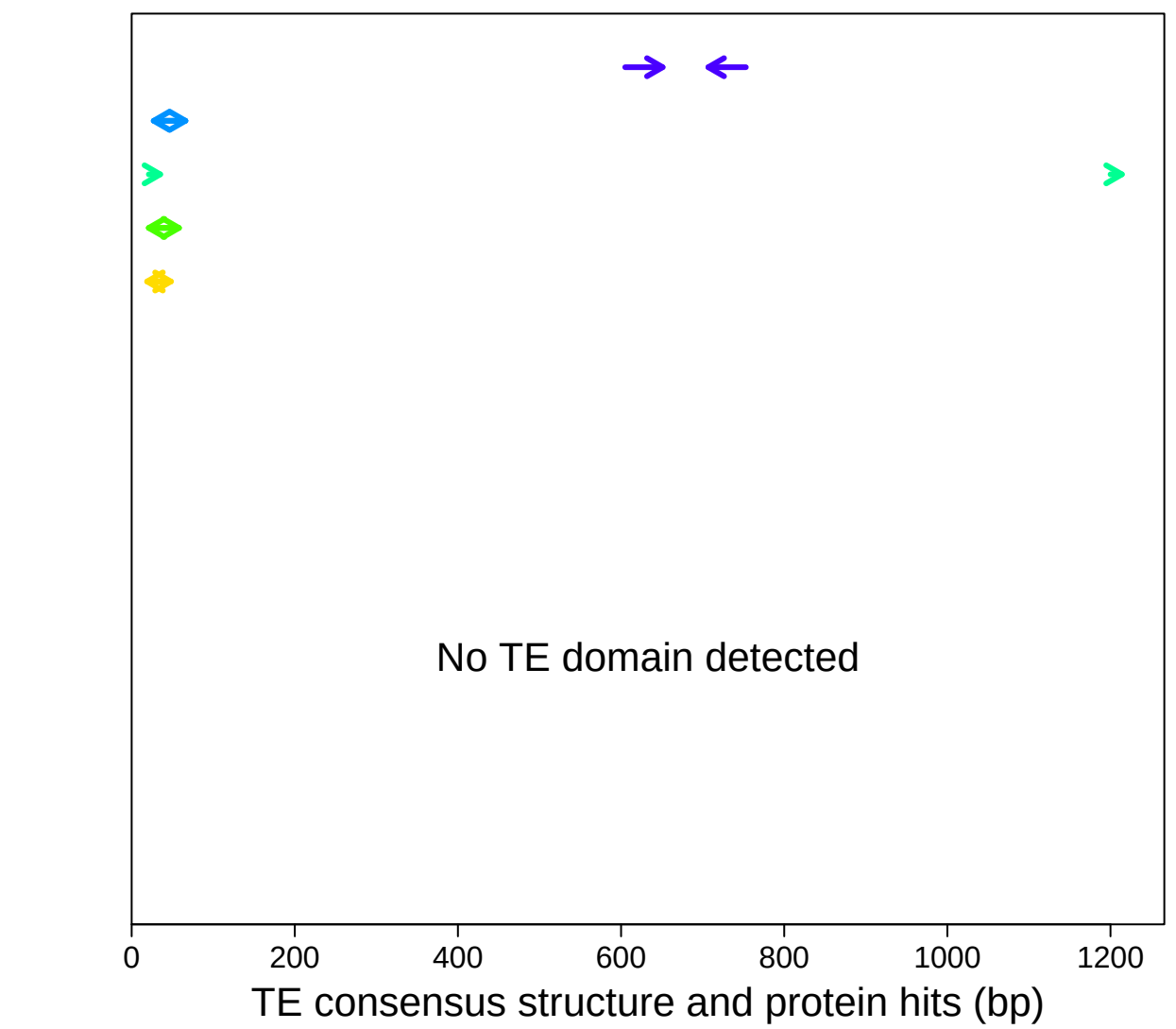
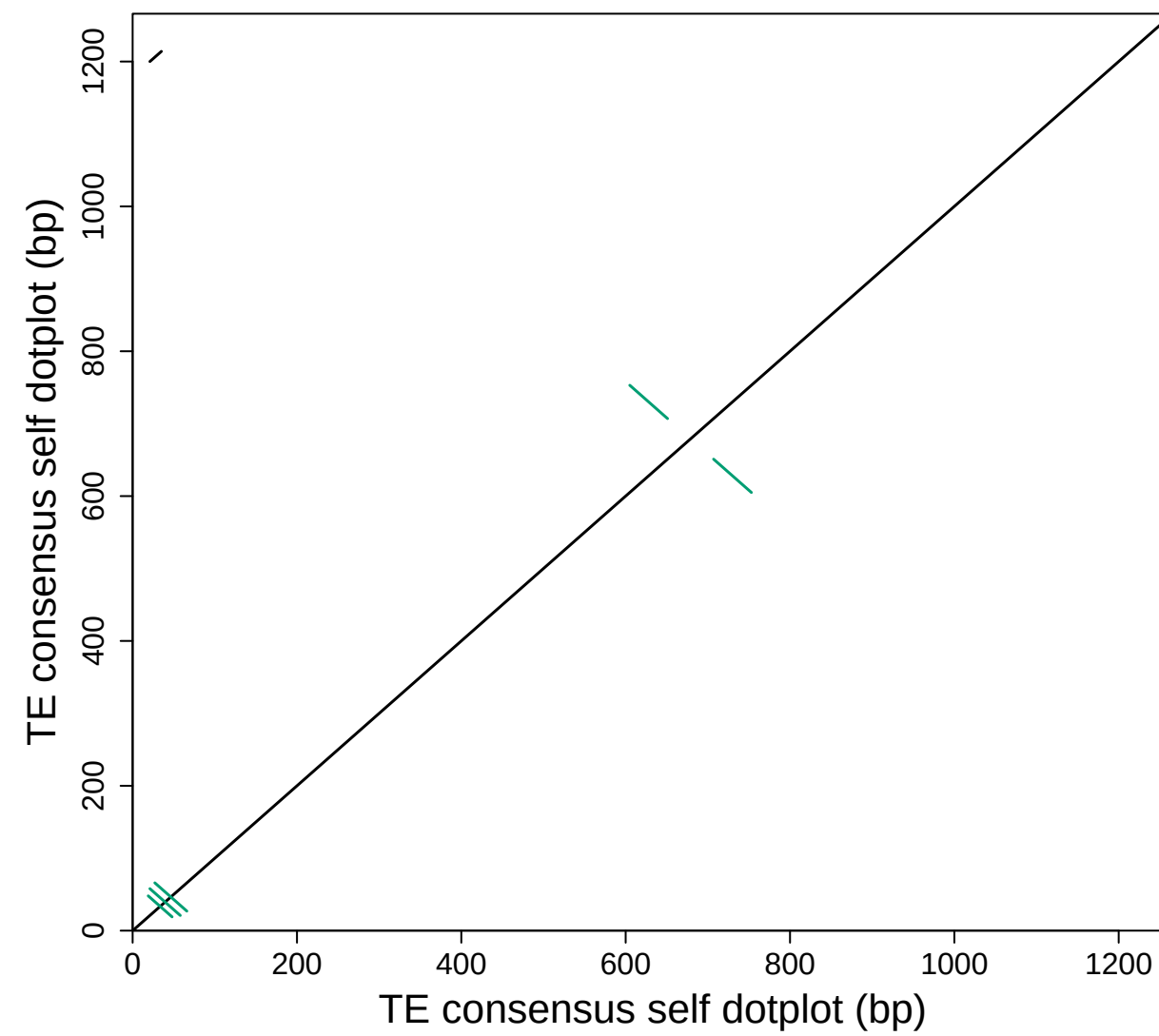
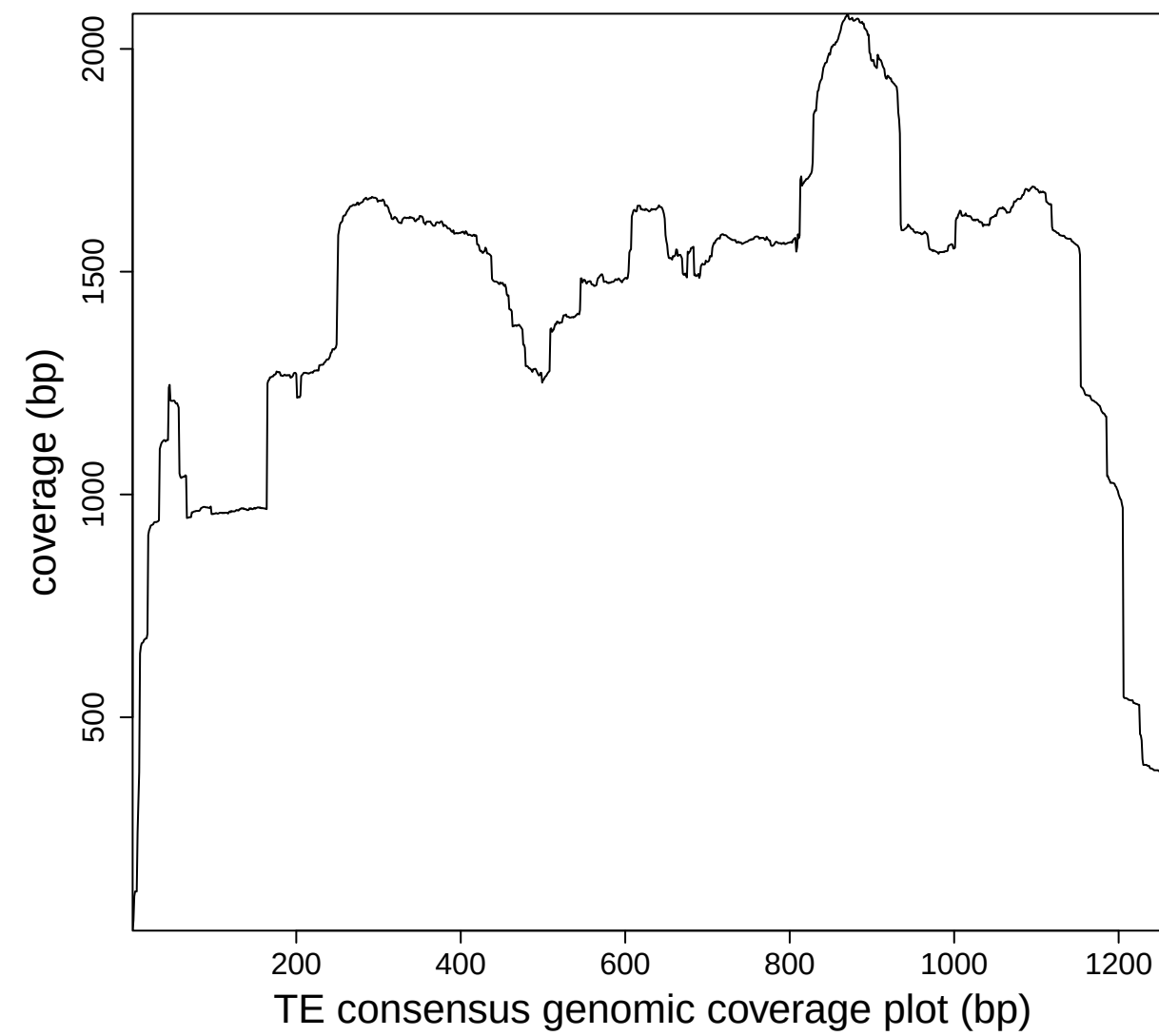
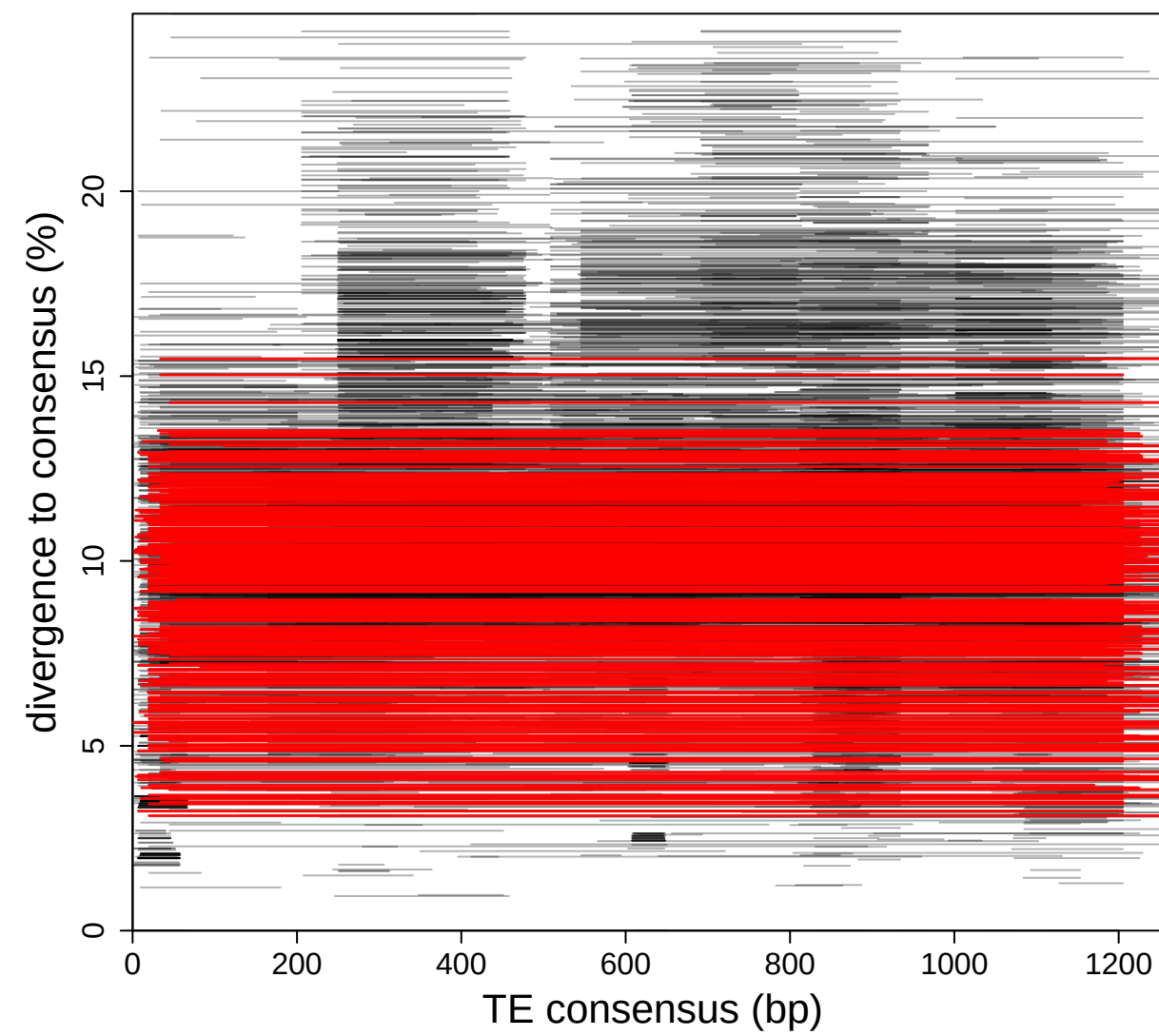
1500

1750



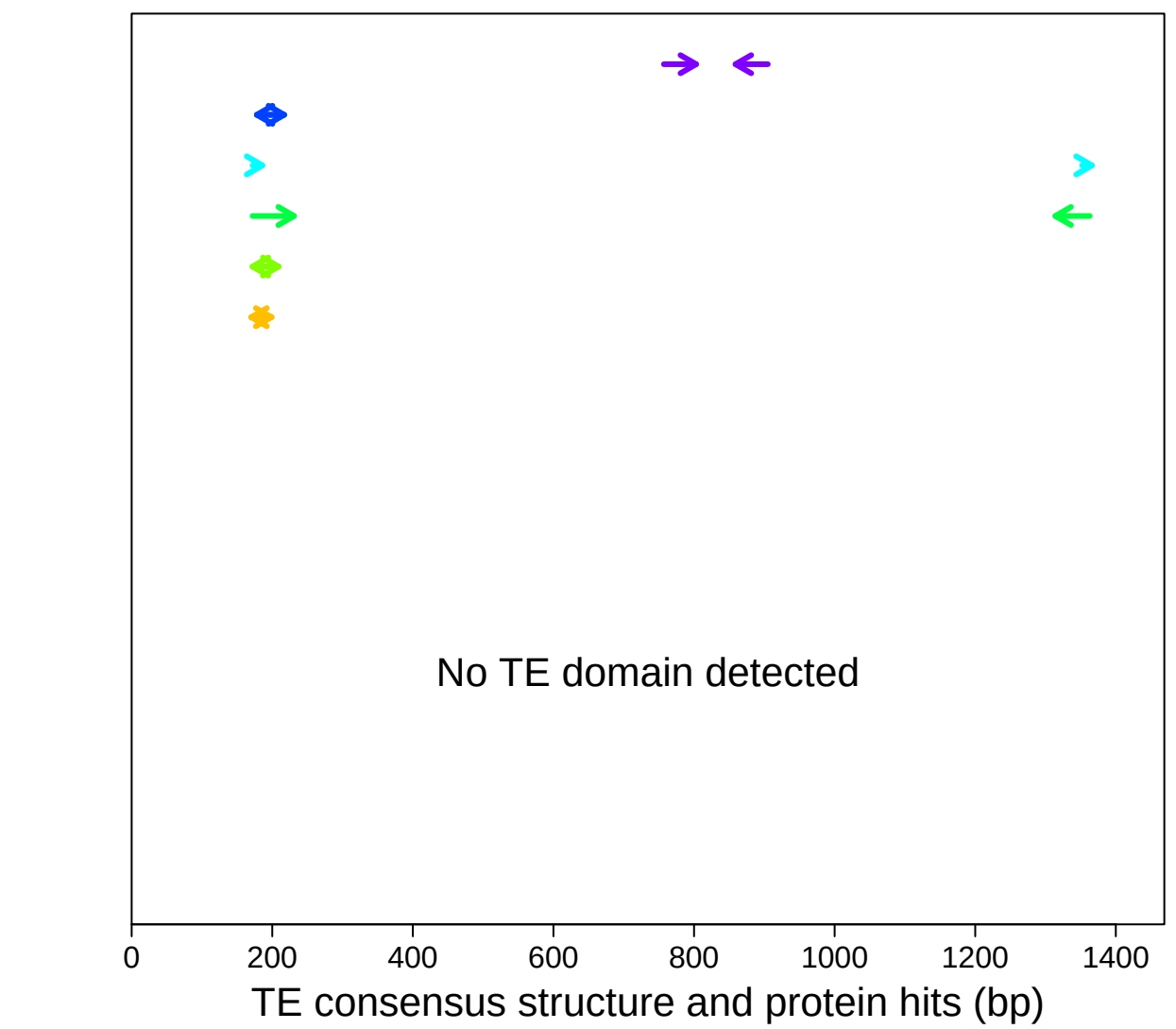
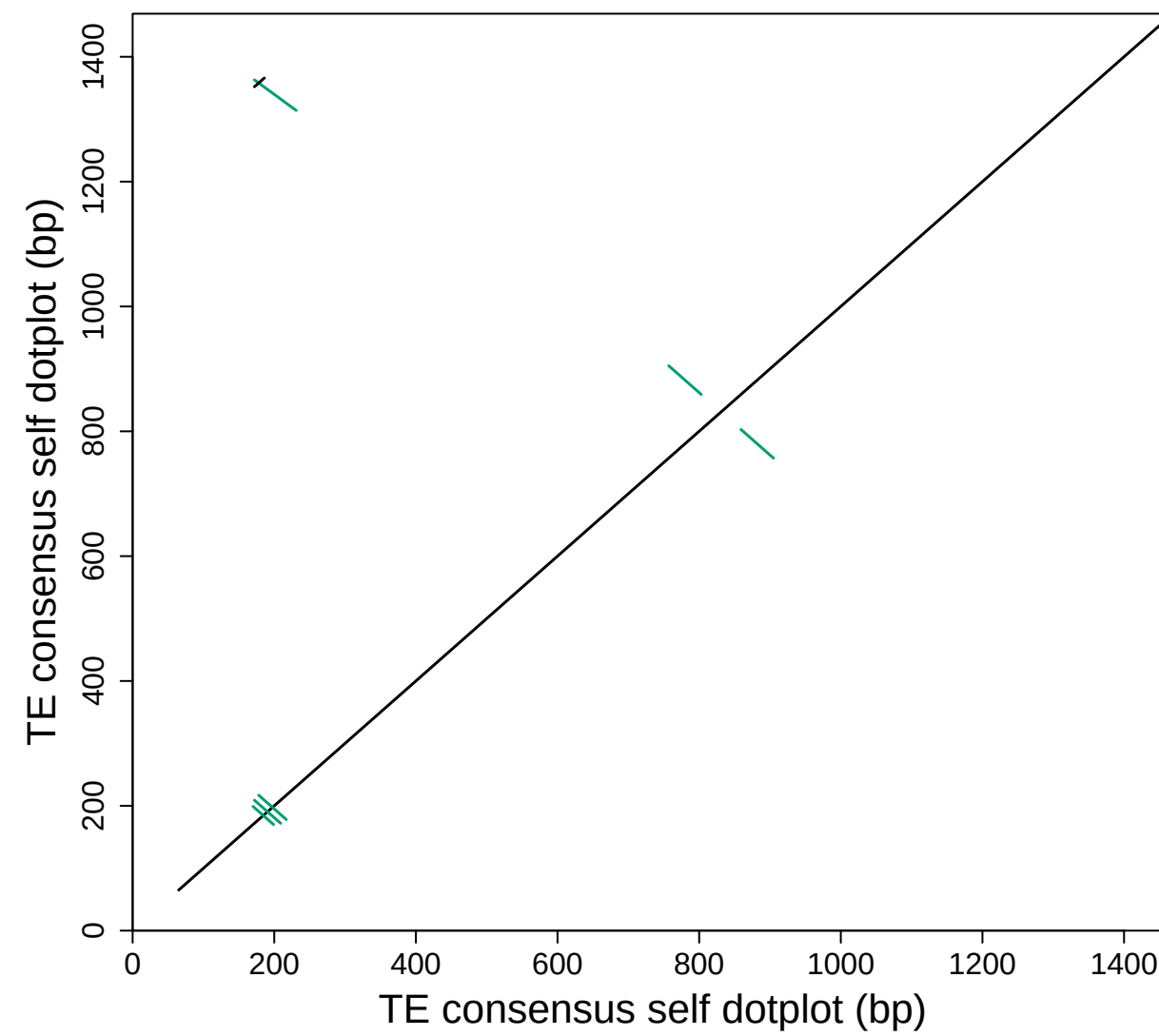
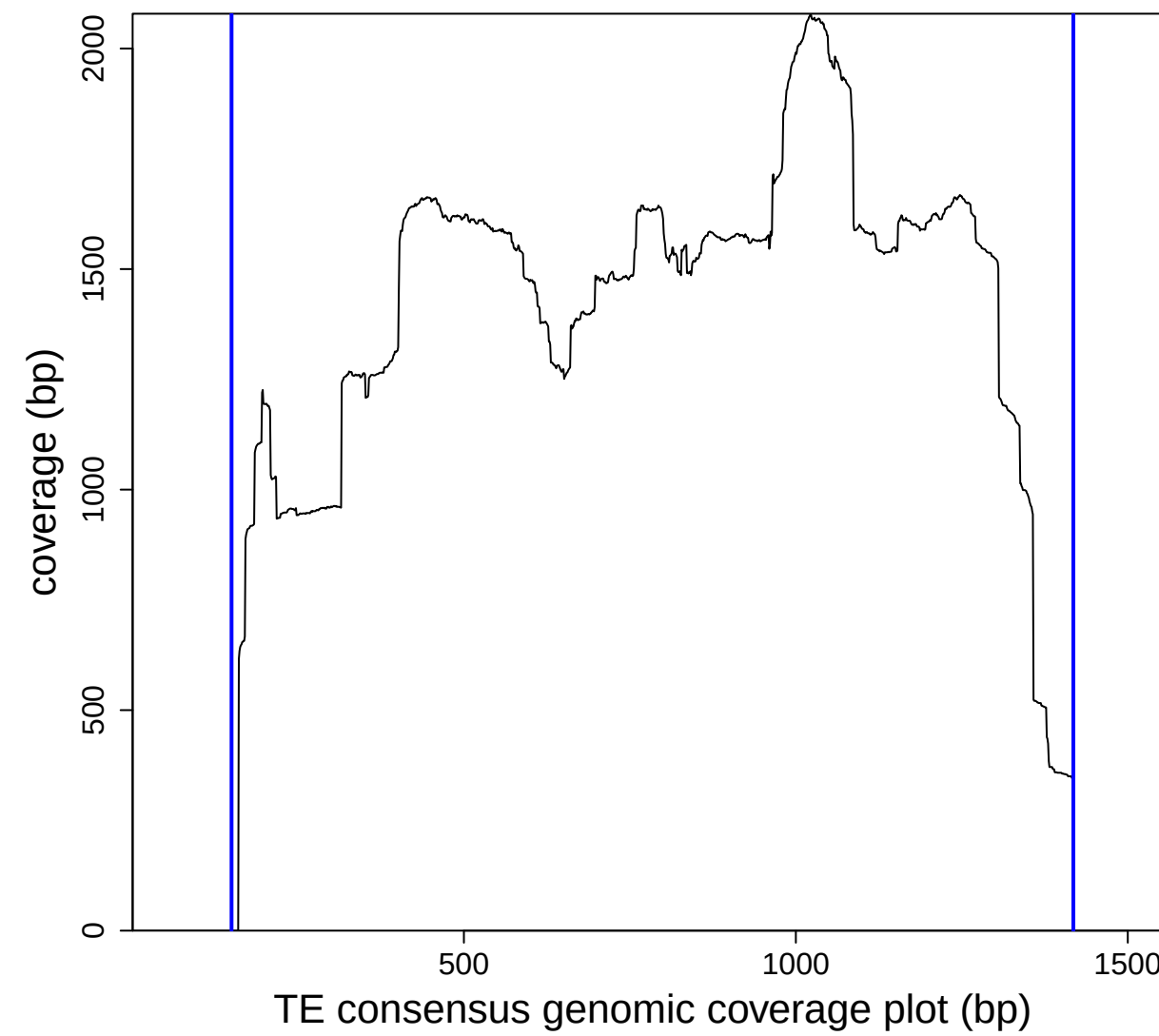
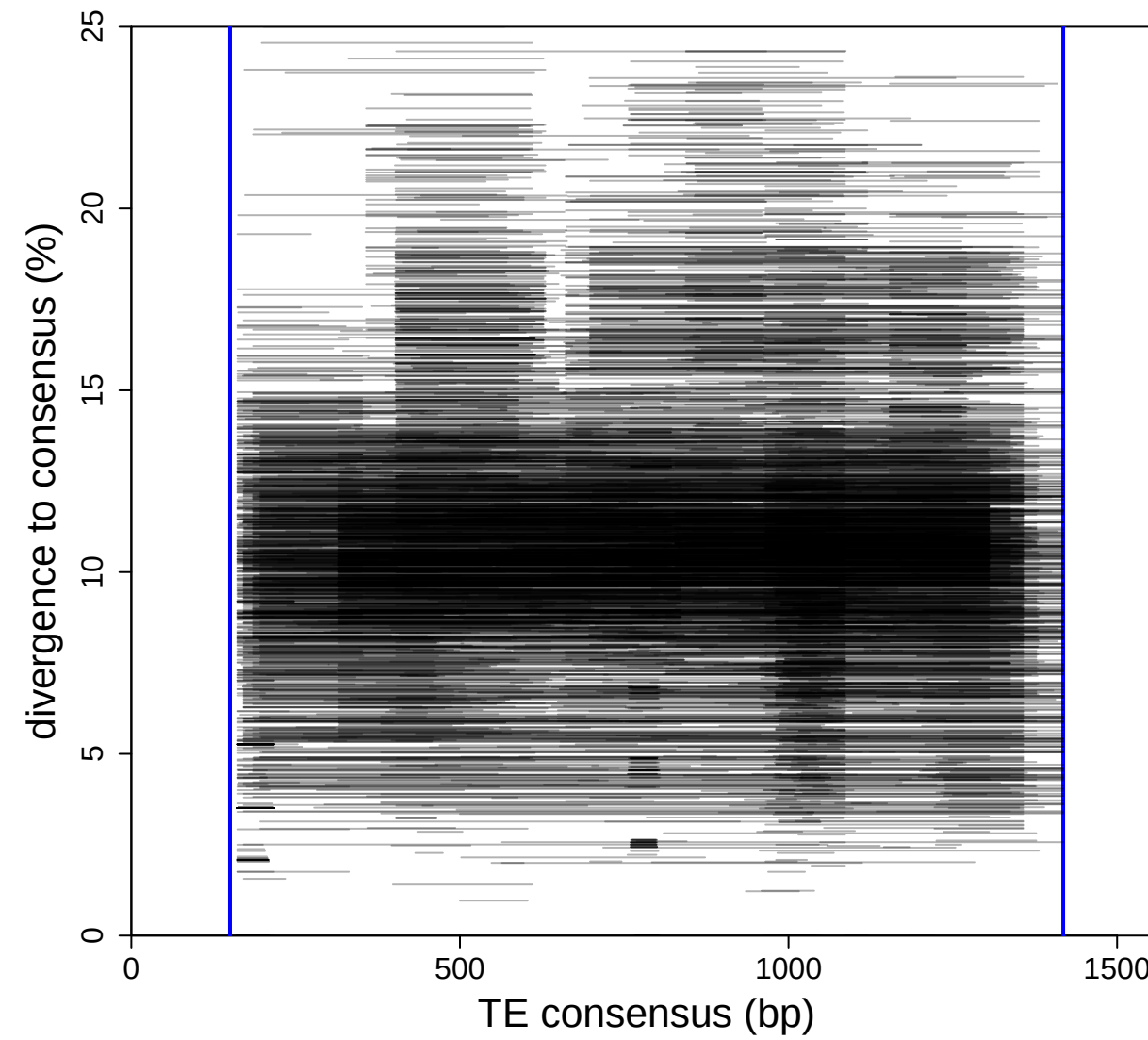
g_3.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln.fa ε
size: 1266bp; fragments: 4558; full length: 299 (>=1139.4bp)

After TETrimmer 1266 bp



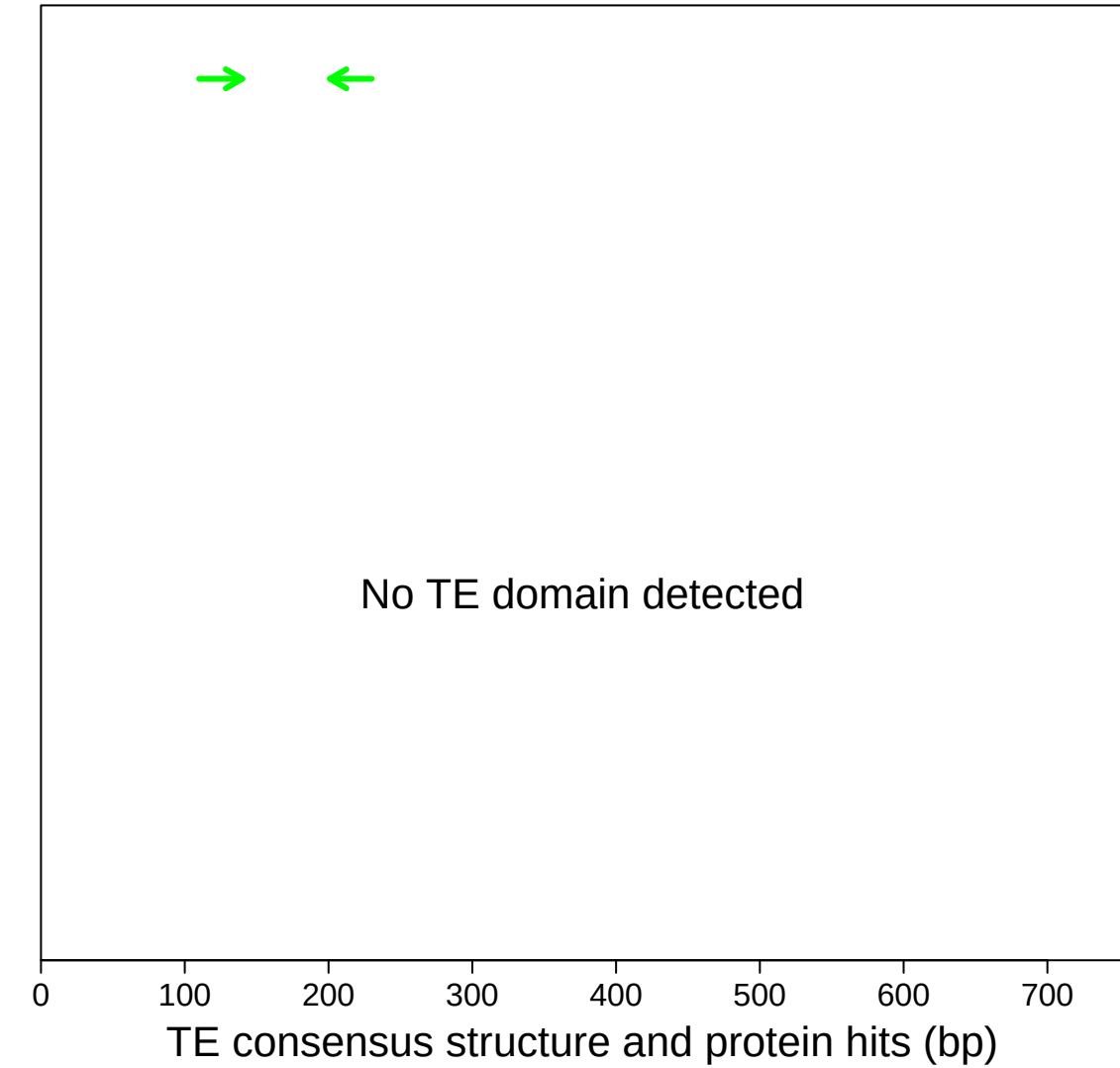
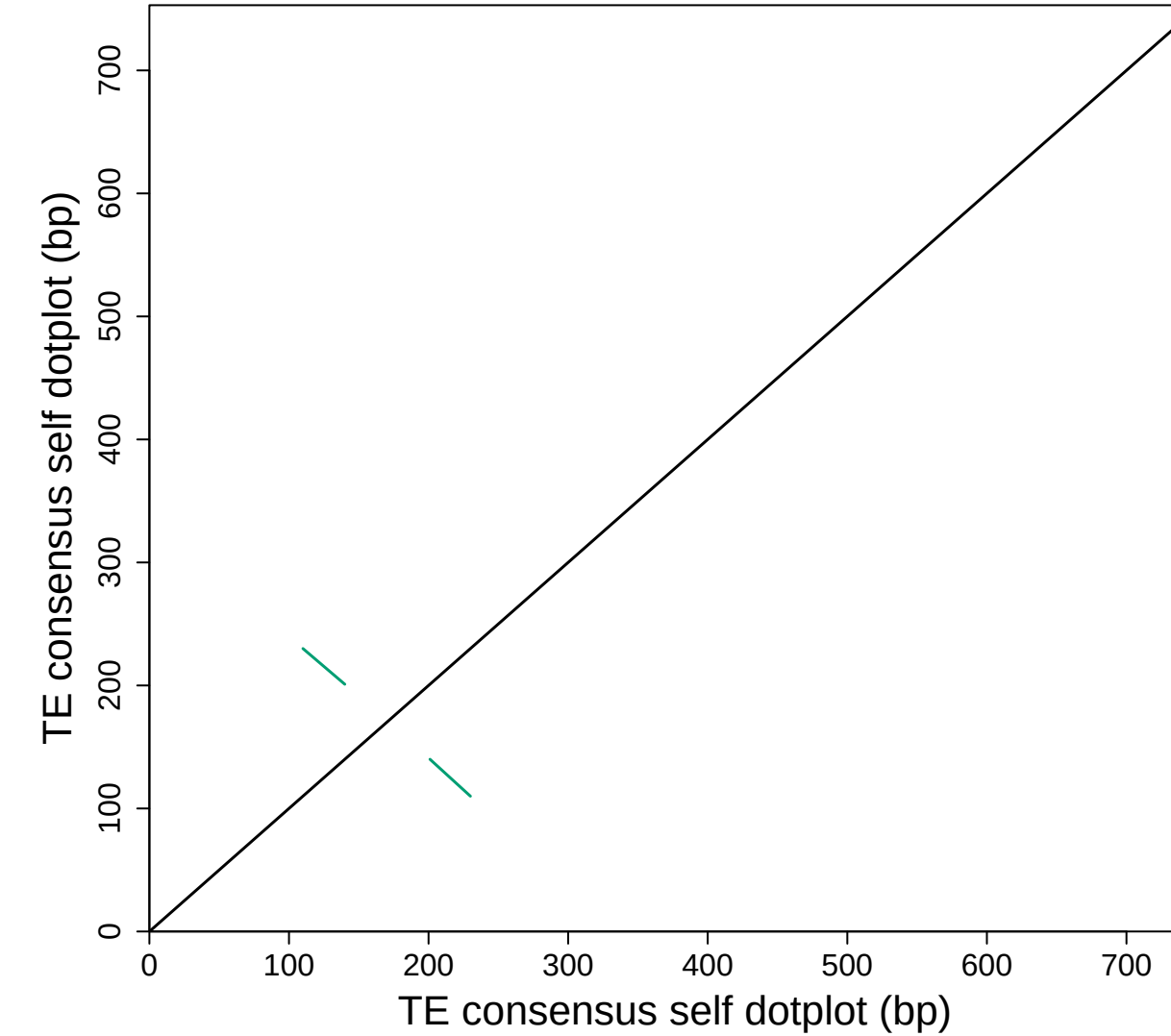
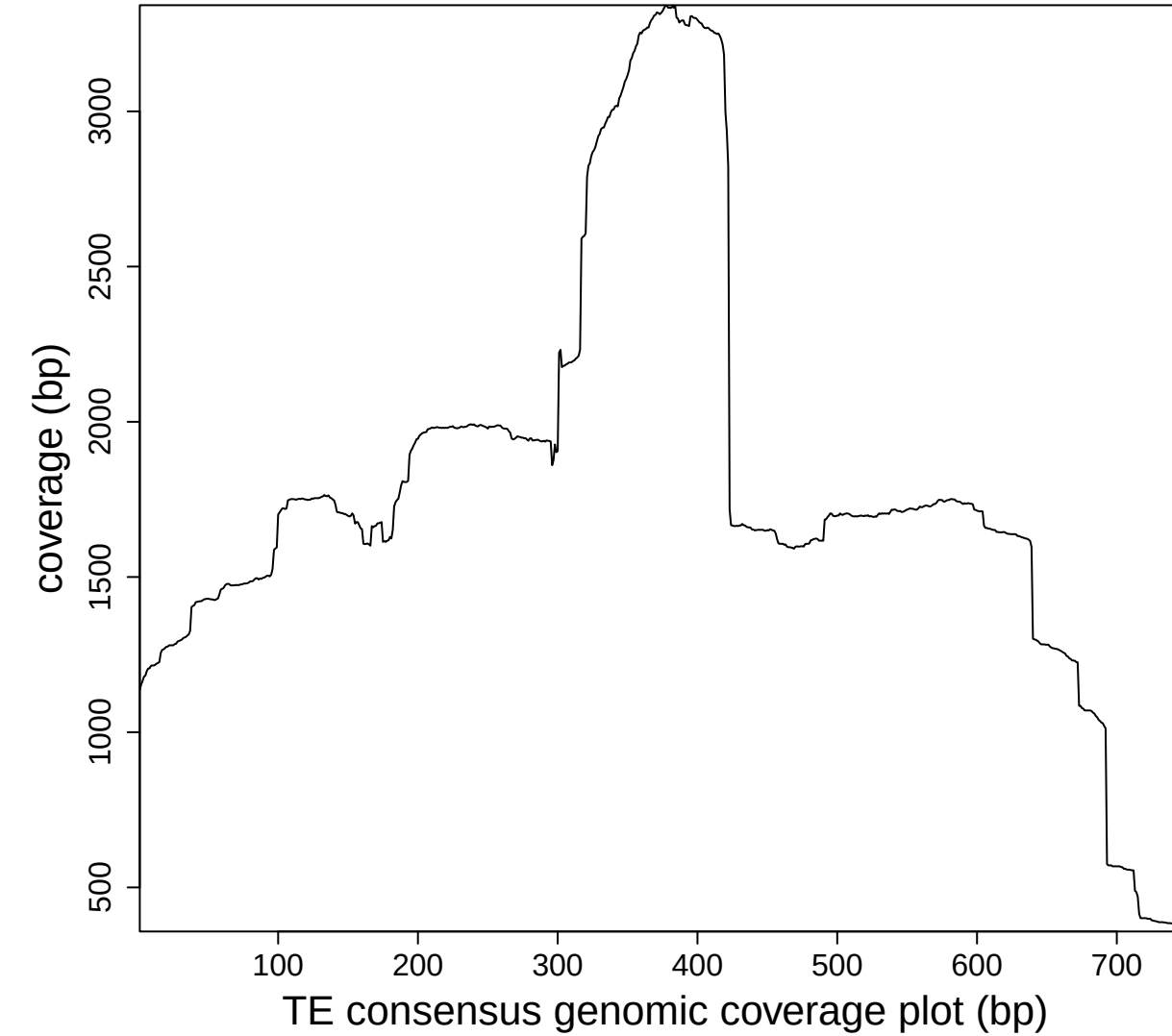
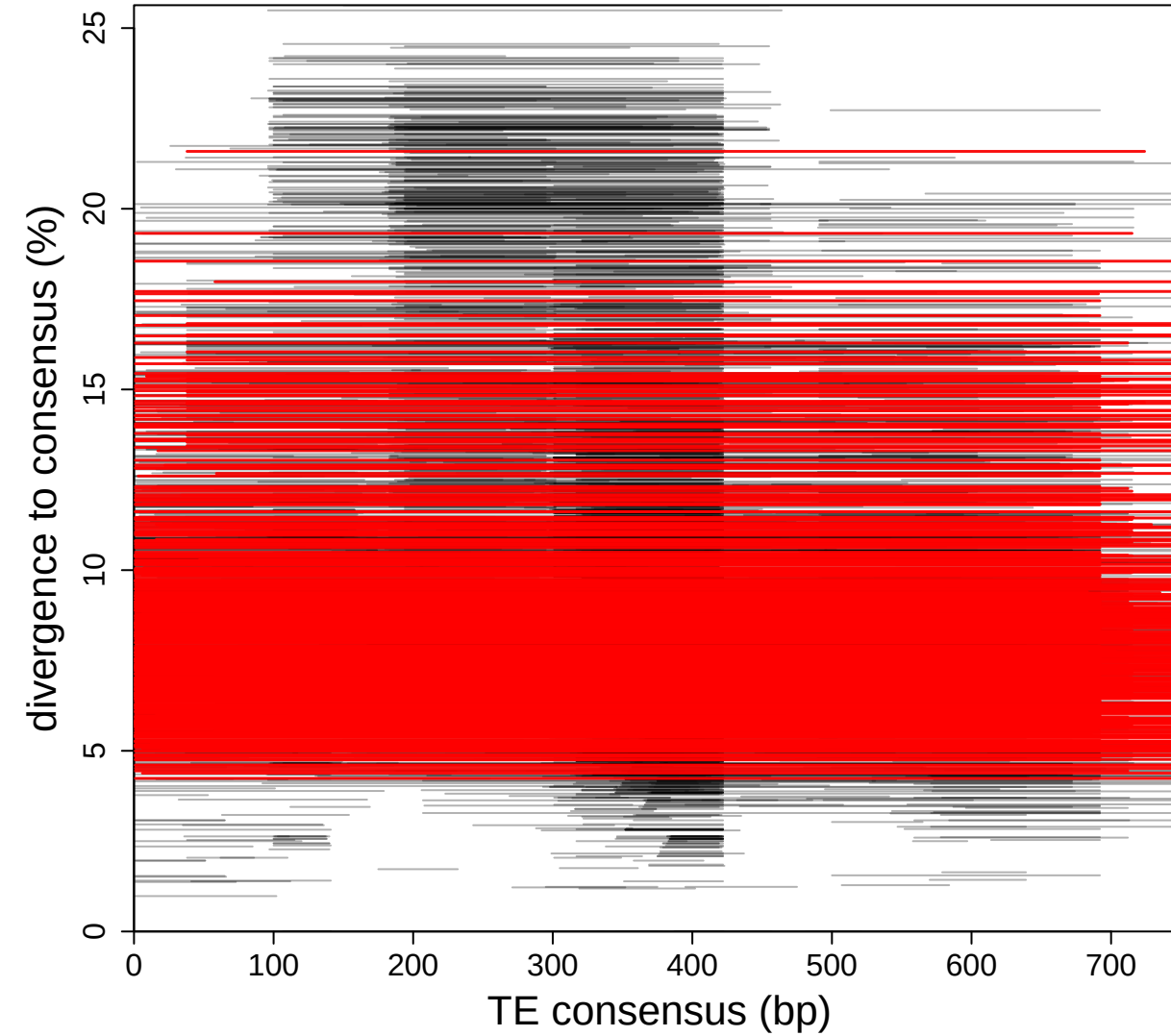
f.bed_g_3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 1568bp; fragments: 4500; full length: 0 (>=1411.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

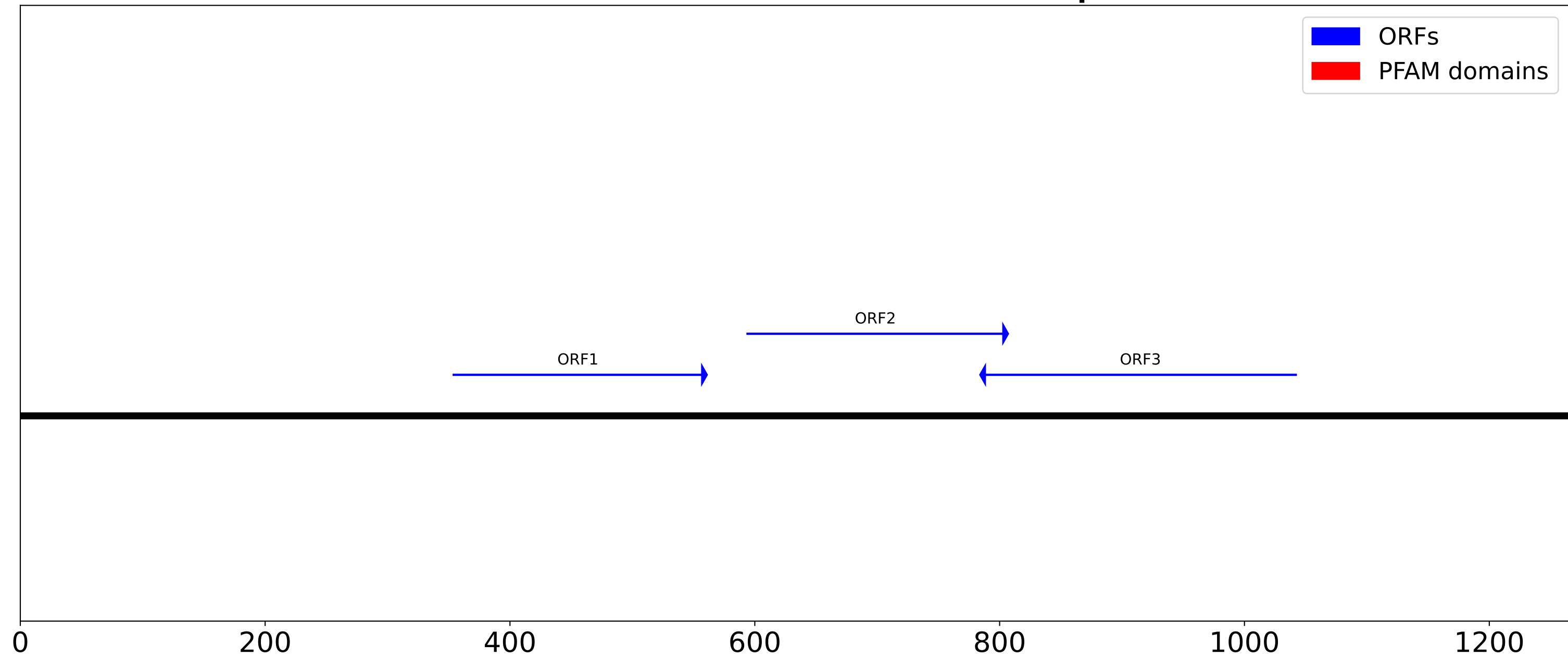


Before TEtrimmer 753 bp

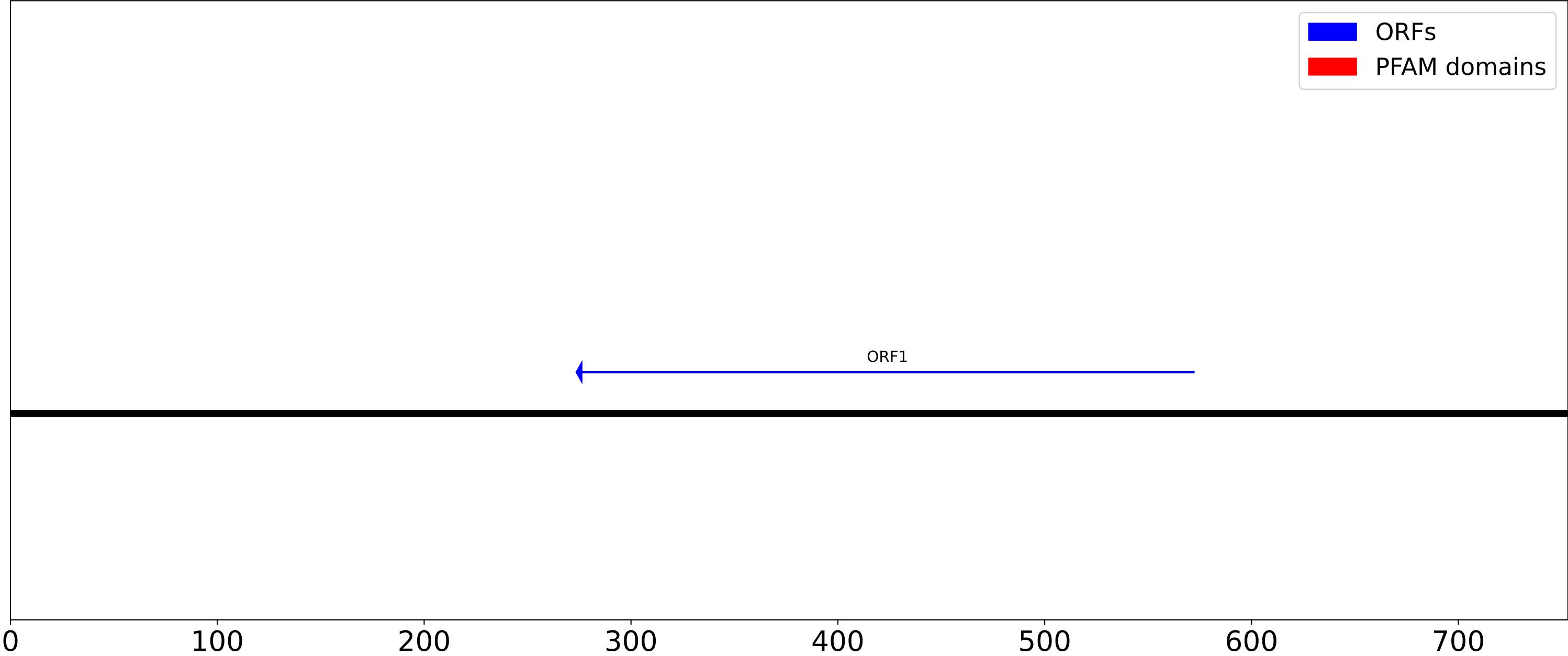
TE: rnd_4_family_1198
size: 753bp; fragments: 4826; full length: 498 (≥ 677.7 bp)



After TEtrimmer ORF and PFAM domain plot



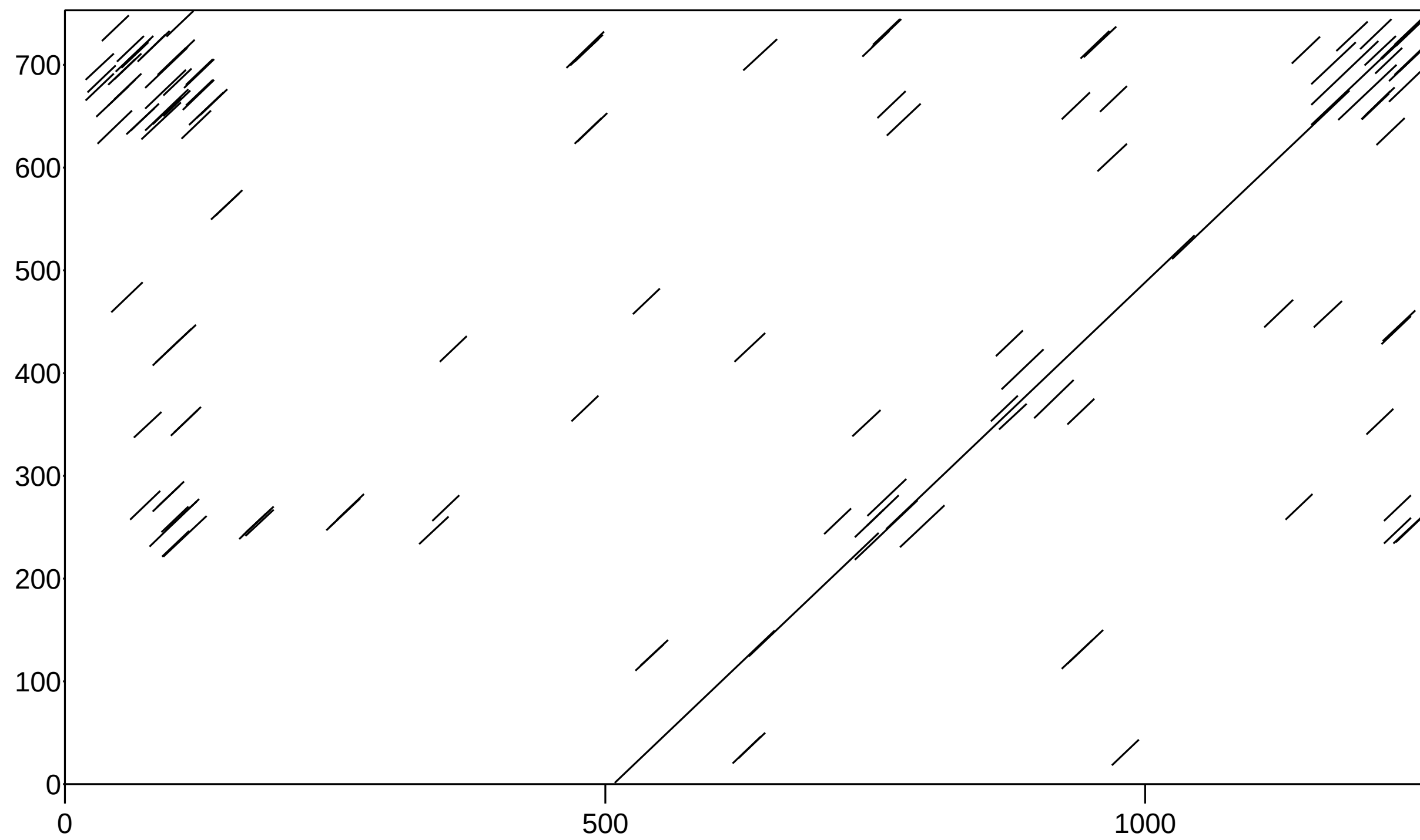
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)