

Start crop Point



End crop Point



MSA length = 5915



Start crop Point

End crop Point

0

10

0

1000

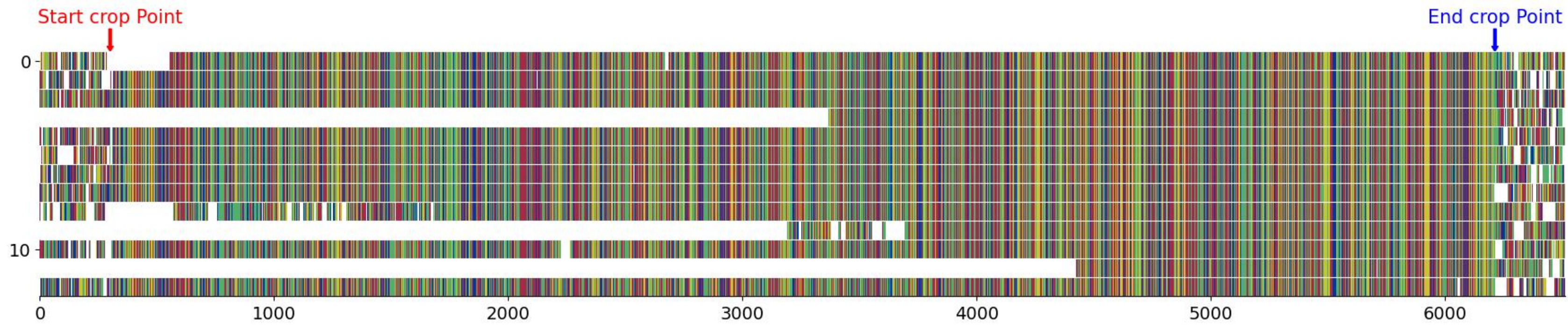
2000

3000

4000

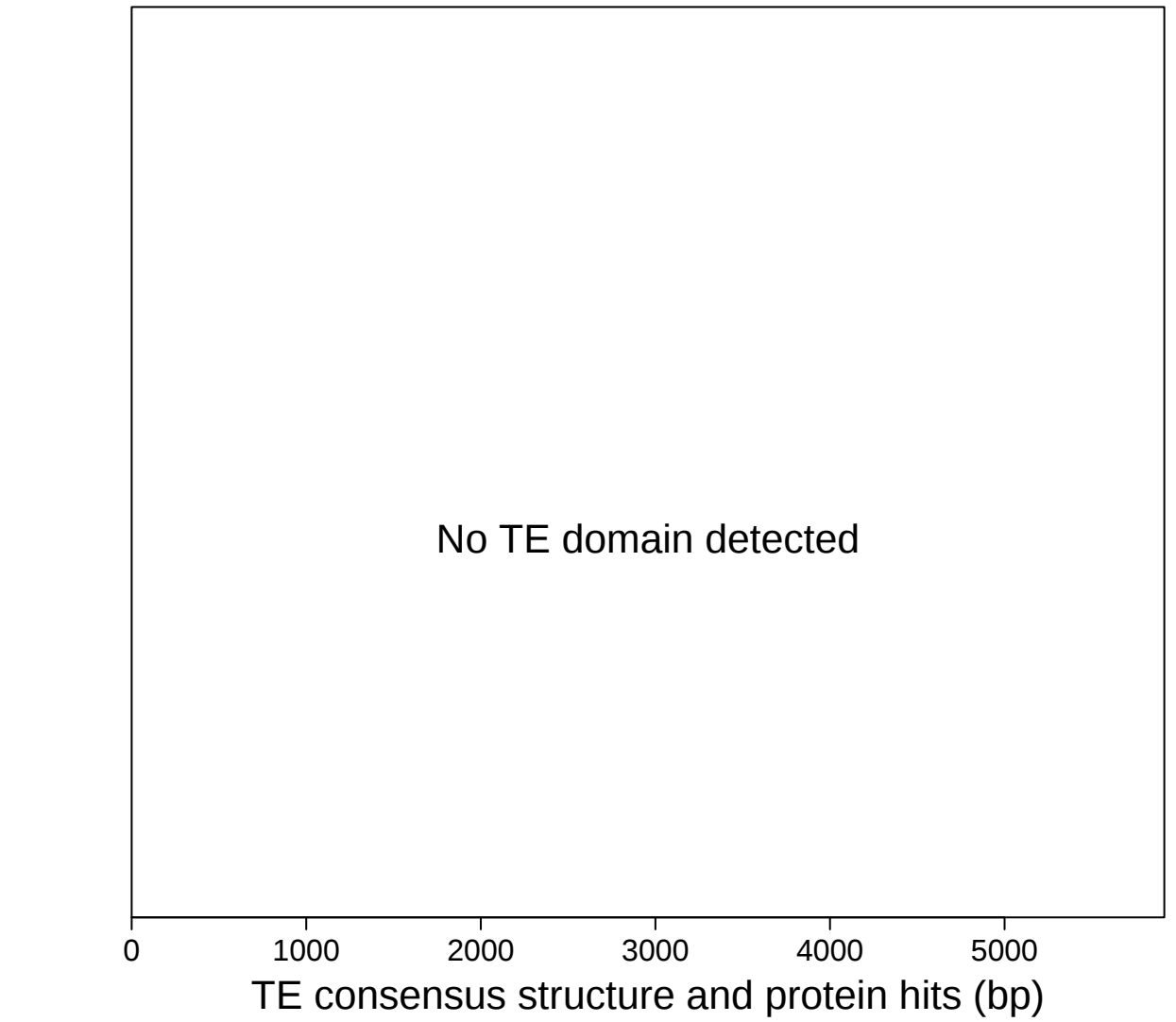
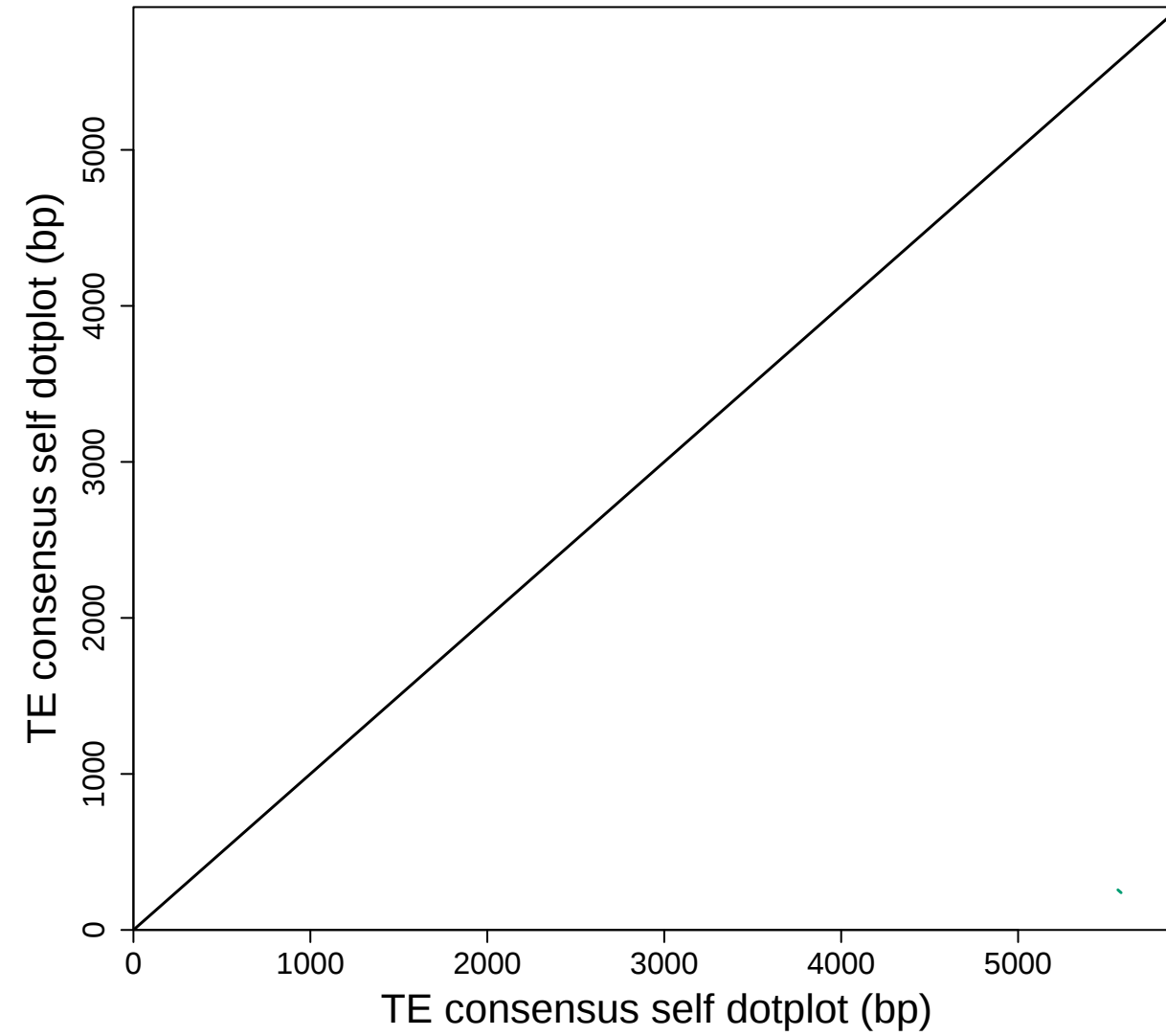
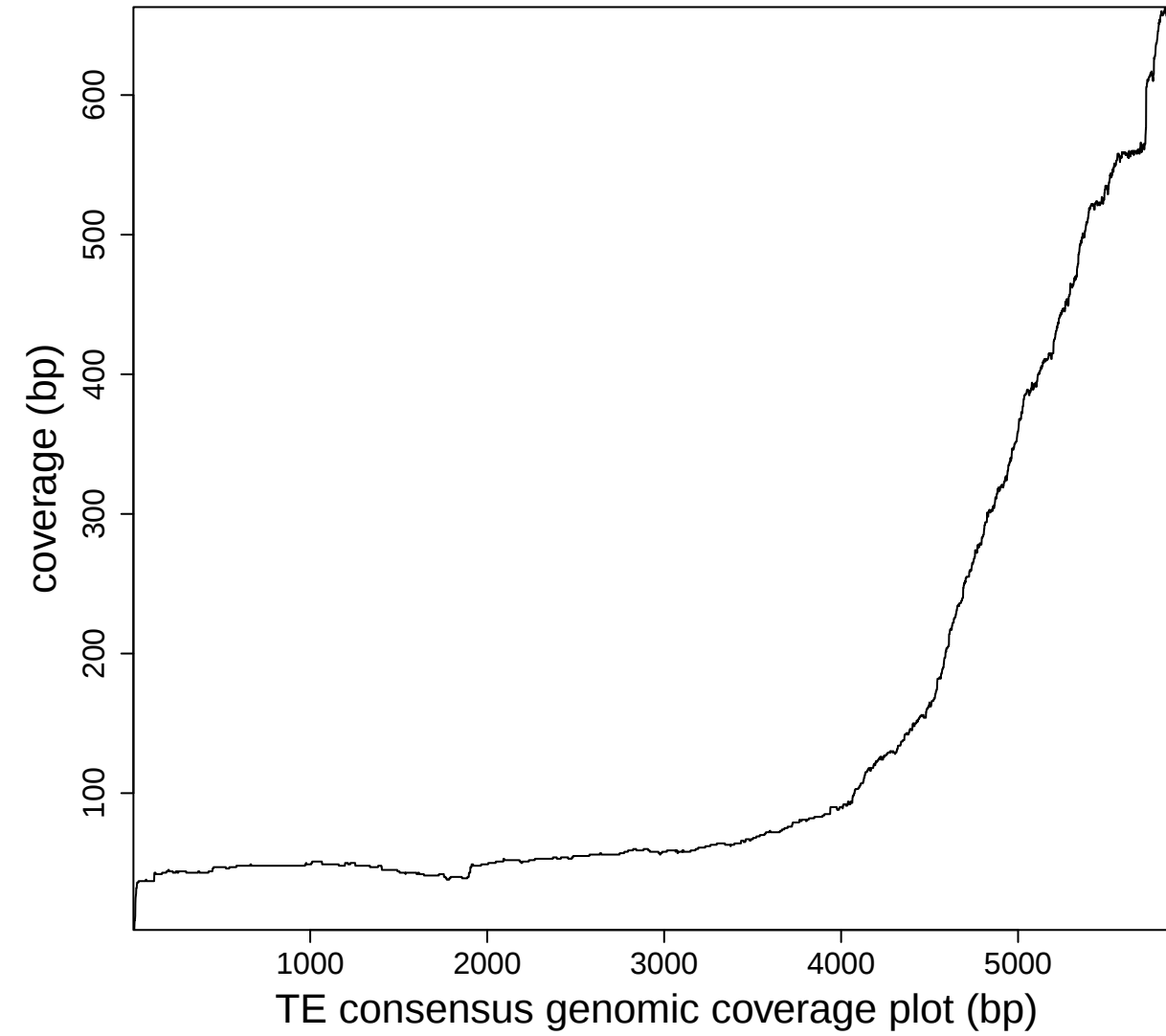
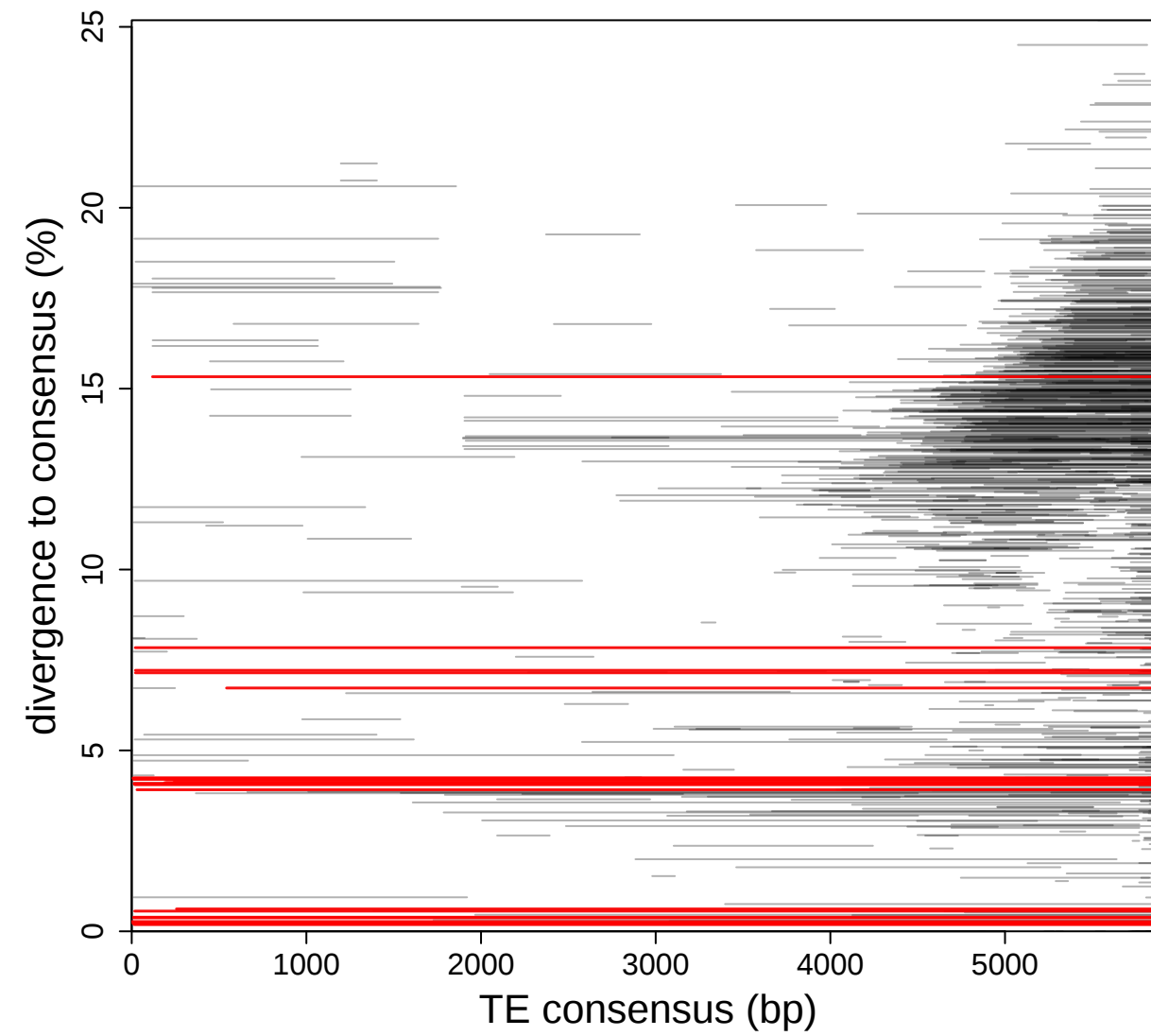
5000

6000



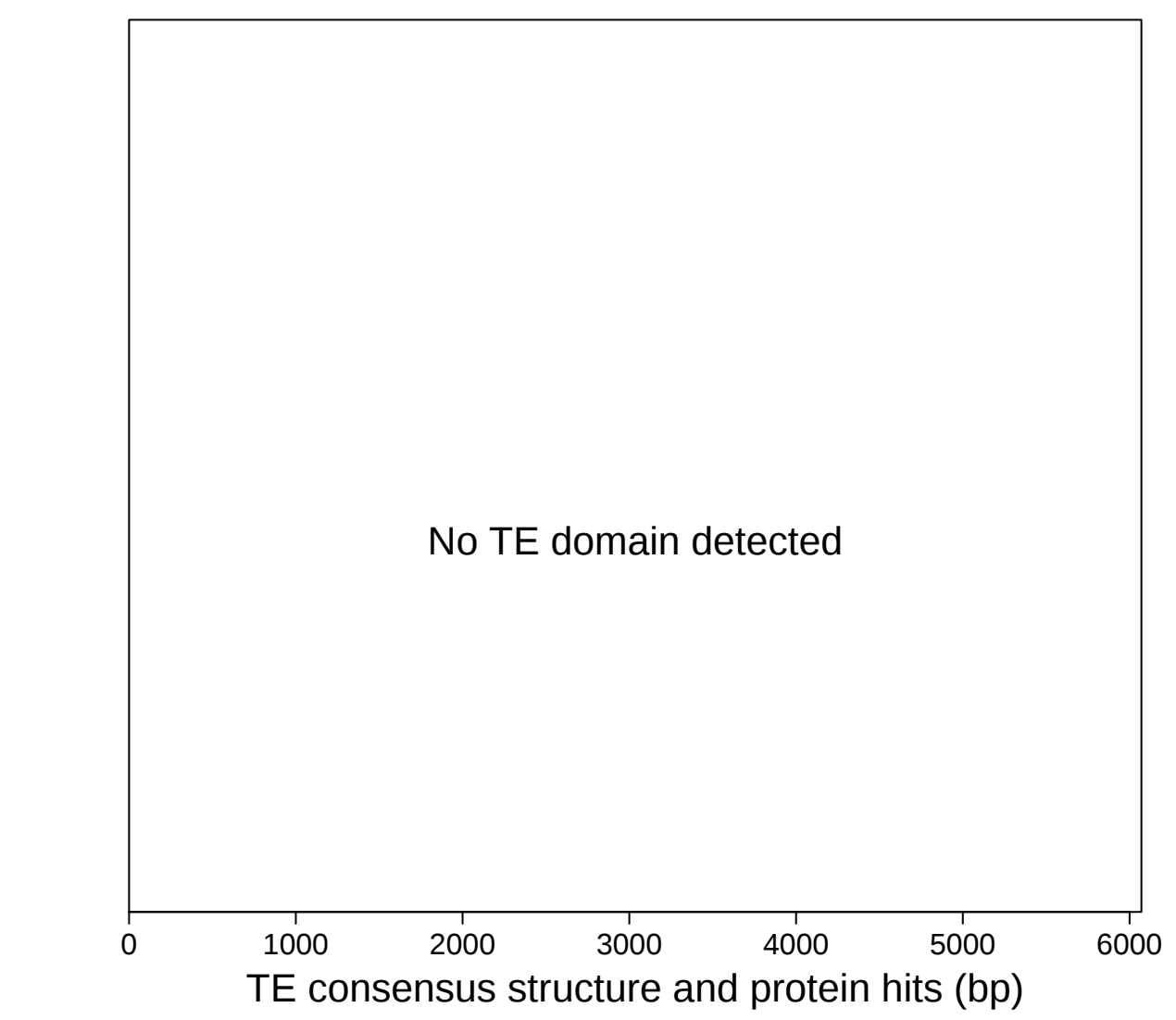
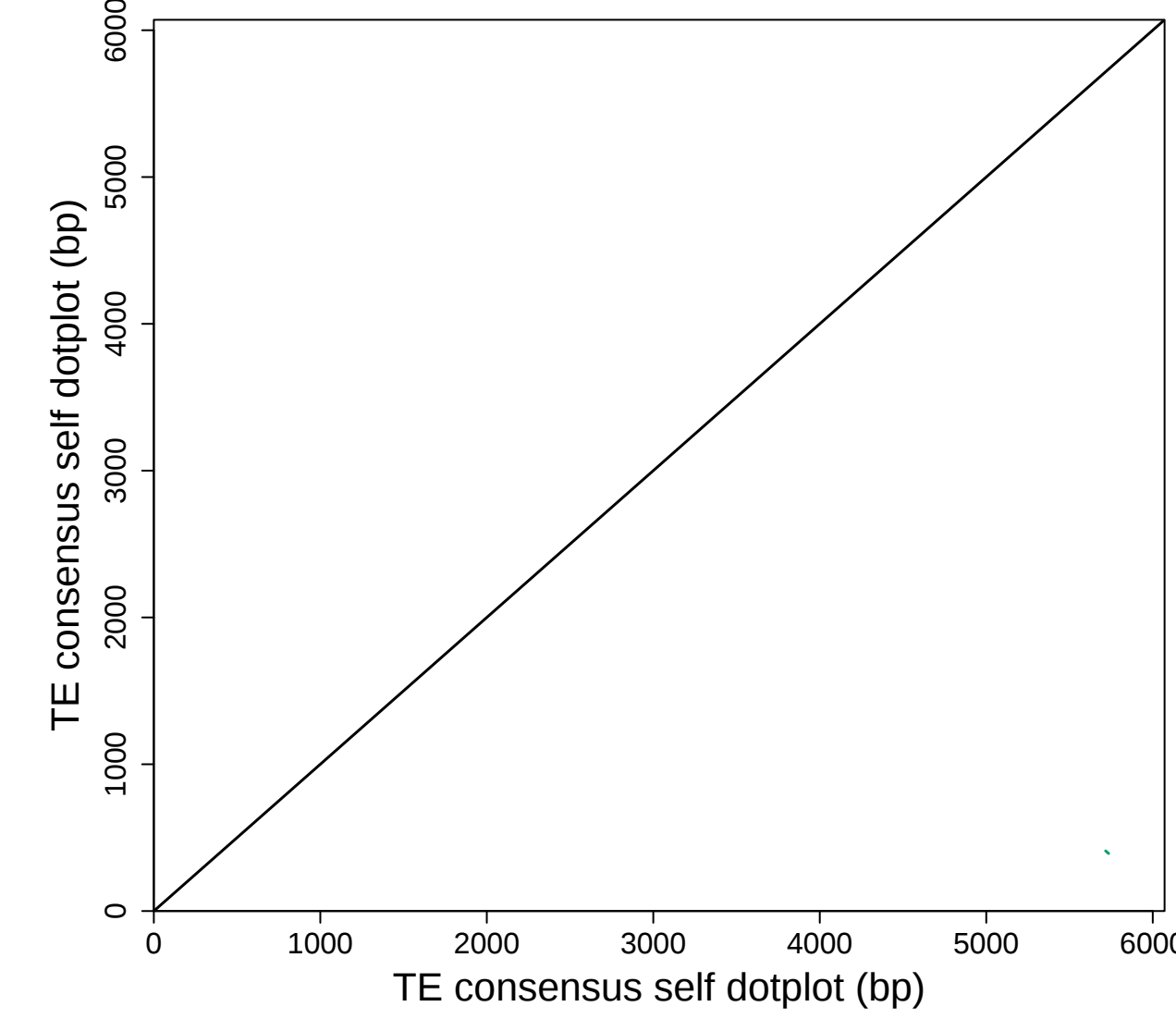
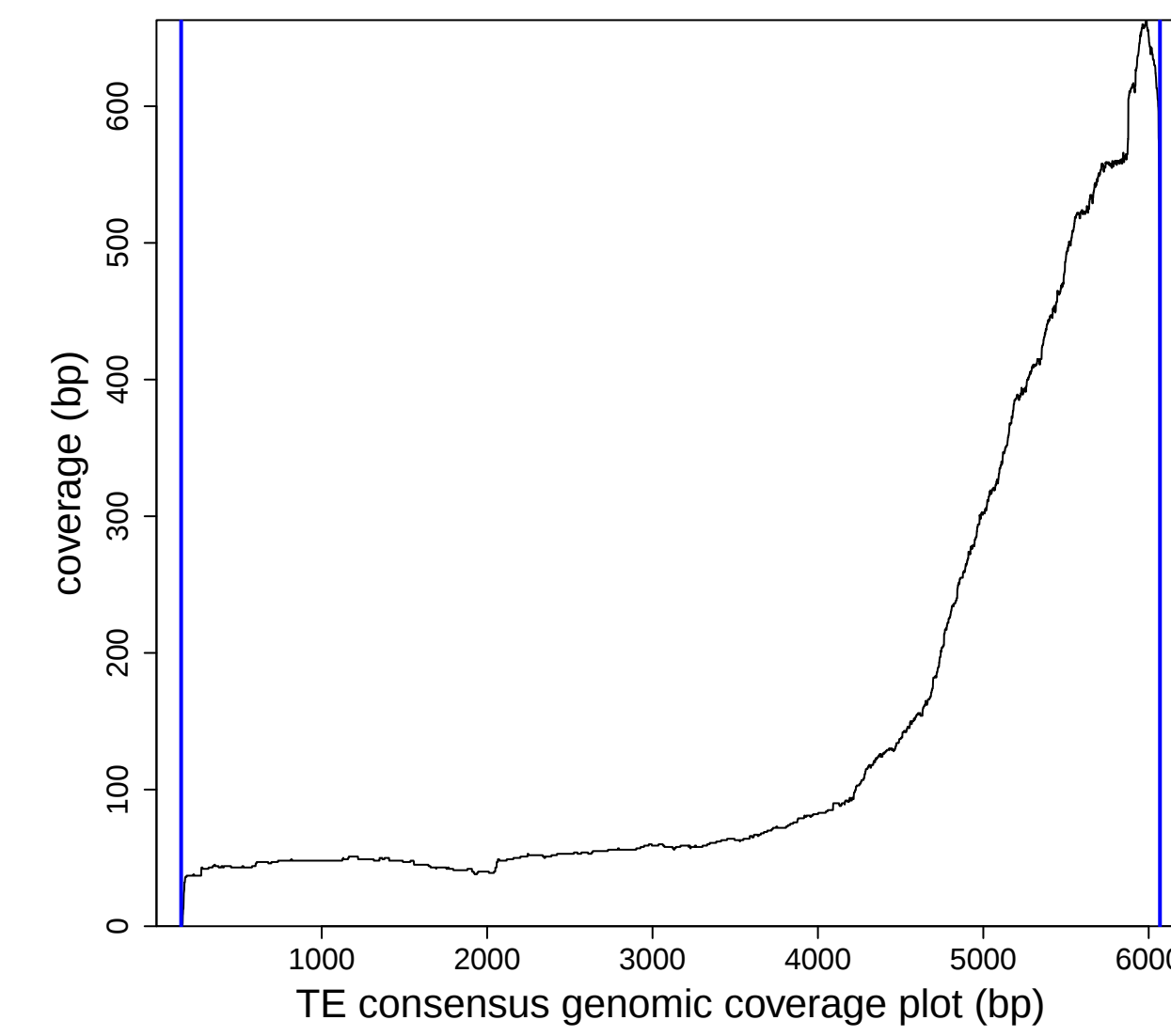
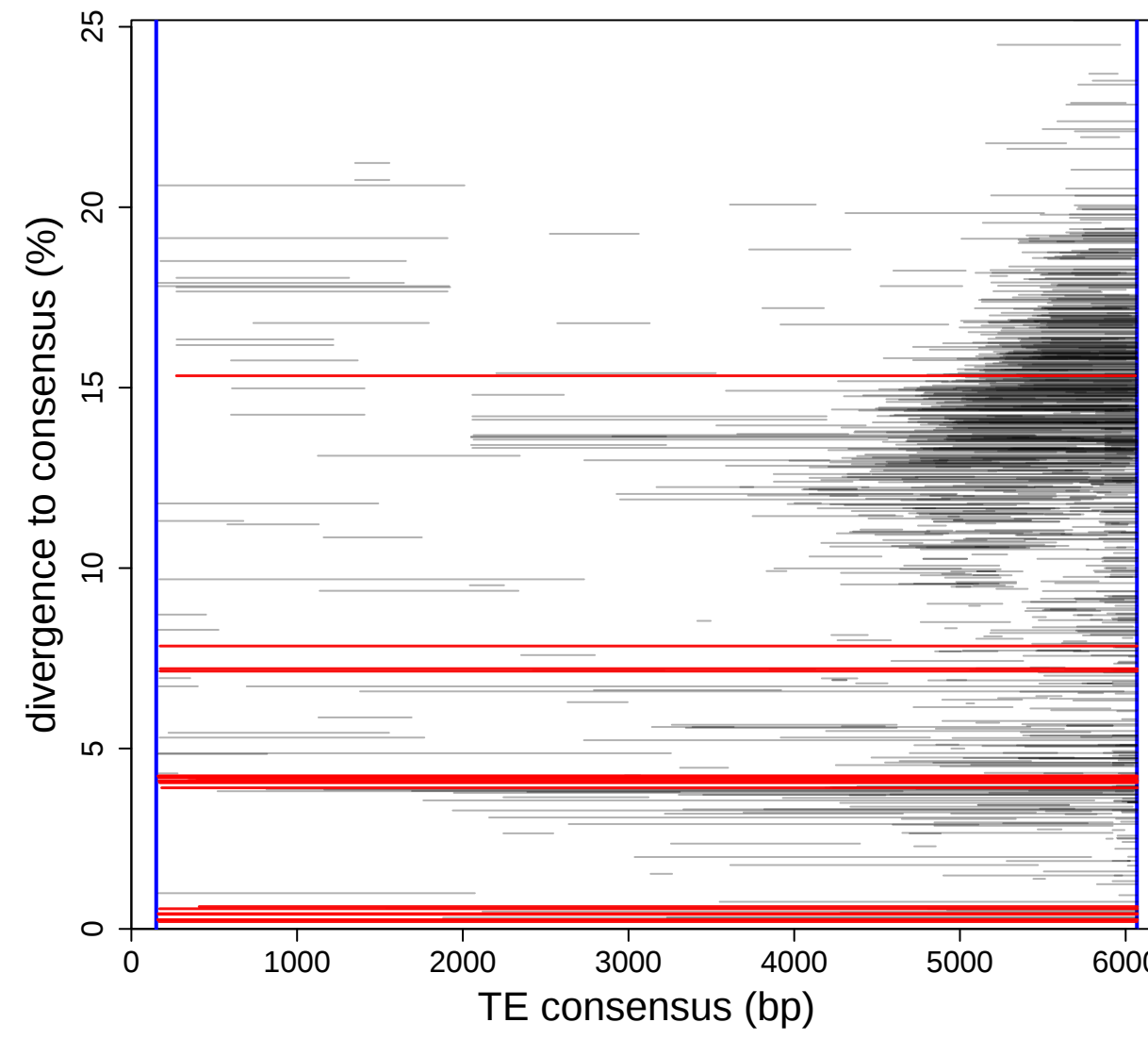
3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.f
size: 5915bp; fragments: 1030; full length: 24 (>=5323.5bp)

After TEtrimmer 5915 bp



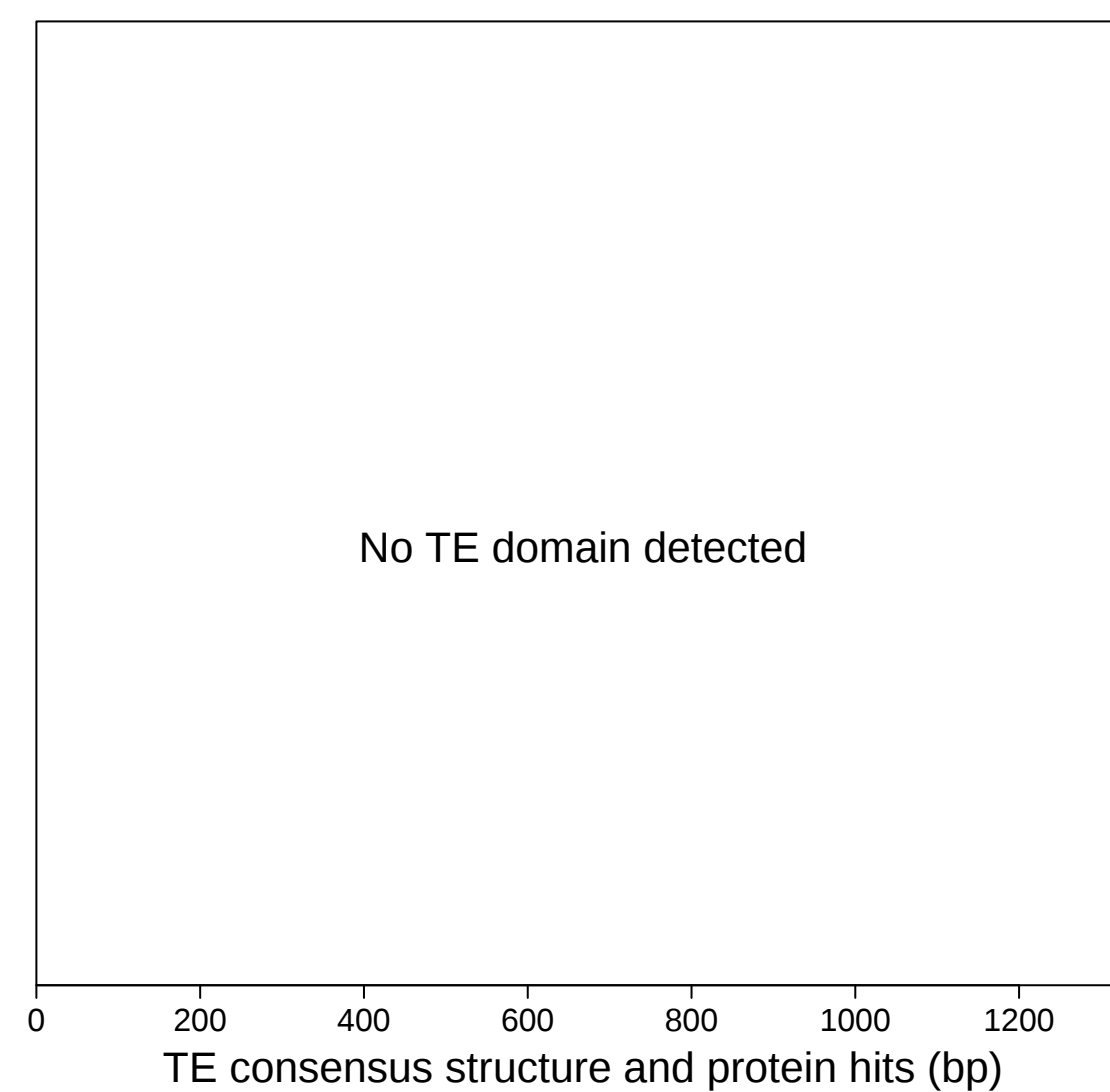
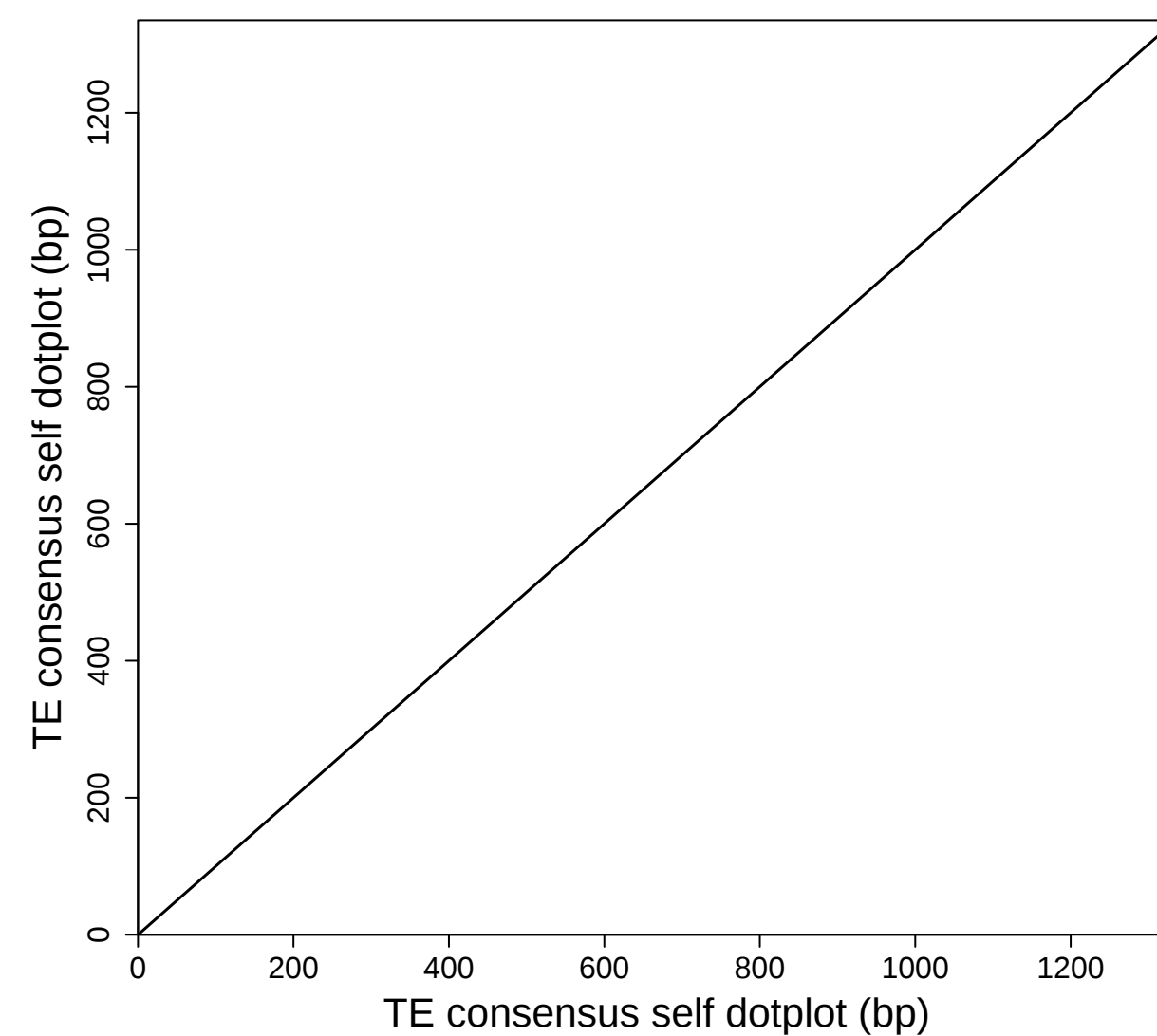
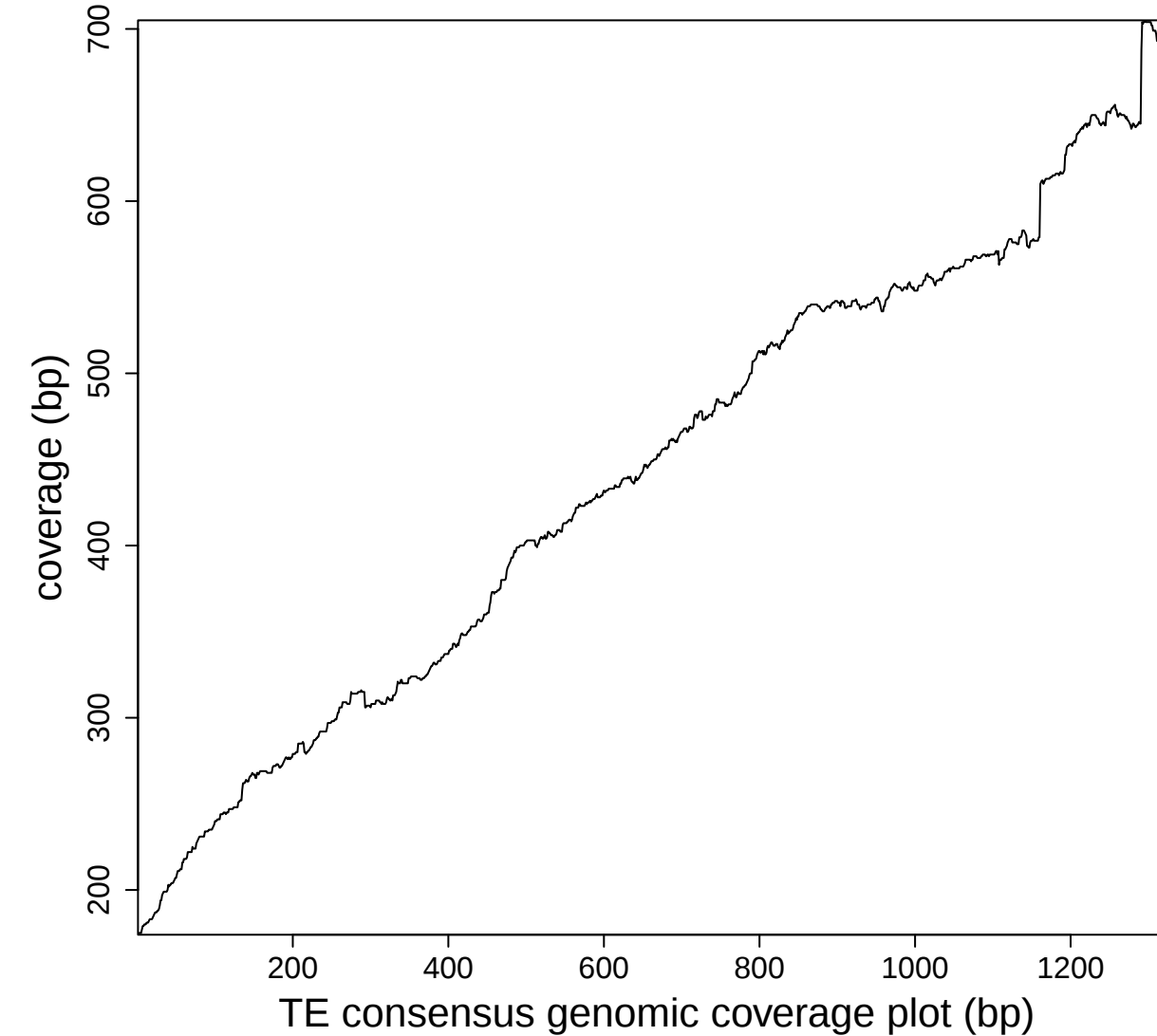
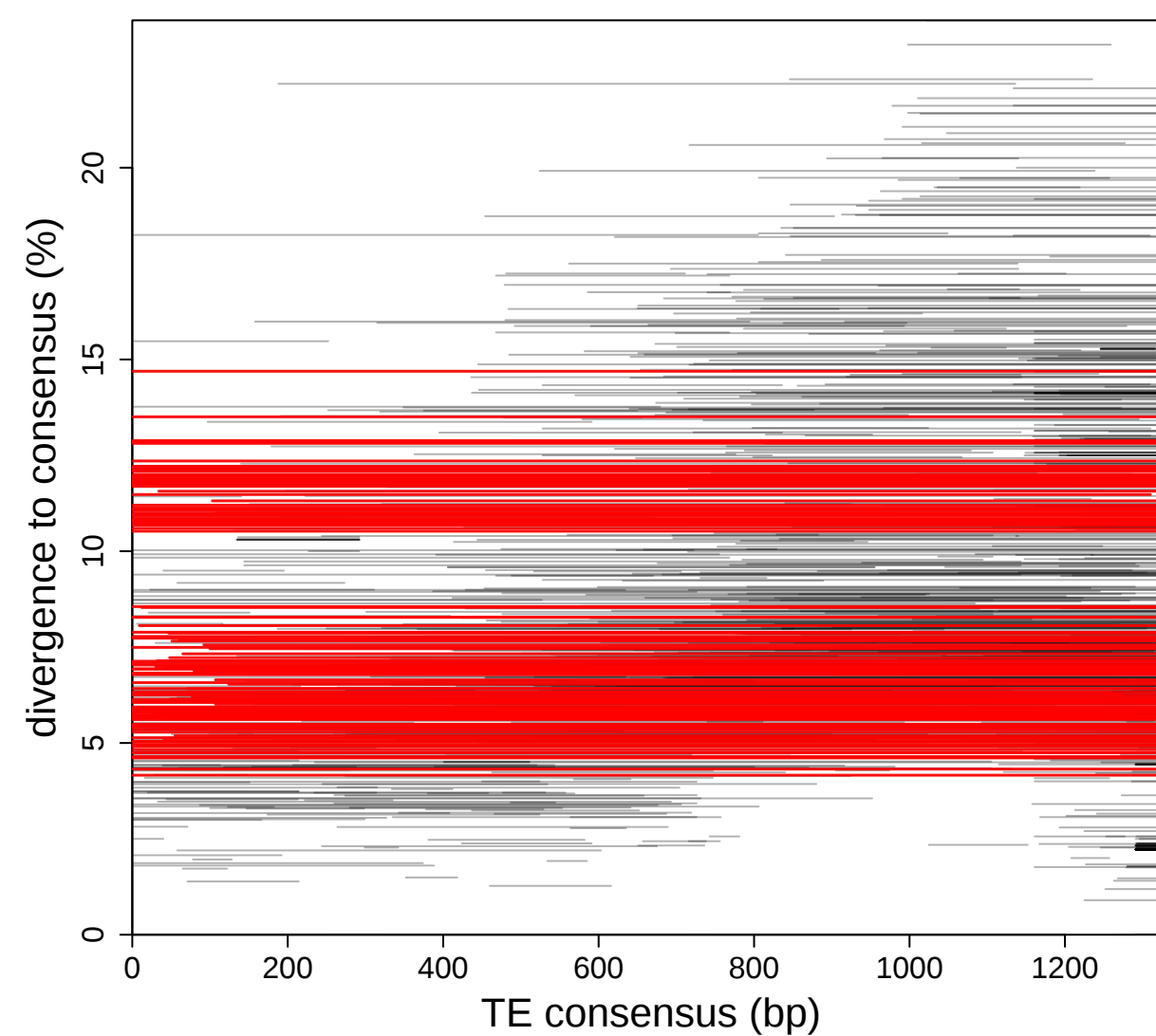
f.bed_g_3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 6218bp; fragments: 1030; full length: 23 (≥ 5596.2 bp)

After TEtrimmer Extended plot Blue lines are boundaries

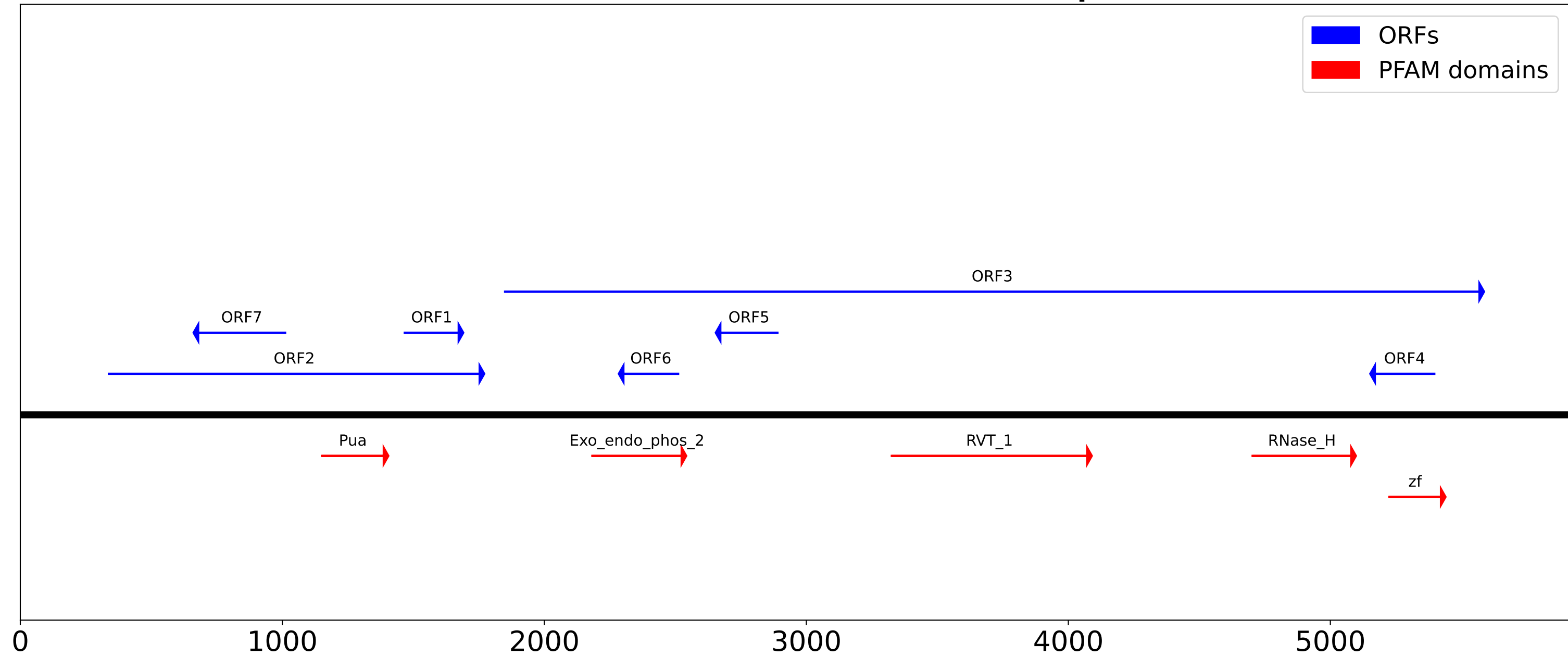


TE: rnd_4_family_3724
size: 1335bp; fragments: 1039; full length: 151 (≥ 1201.5 bp)

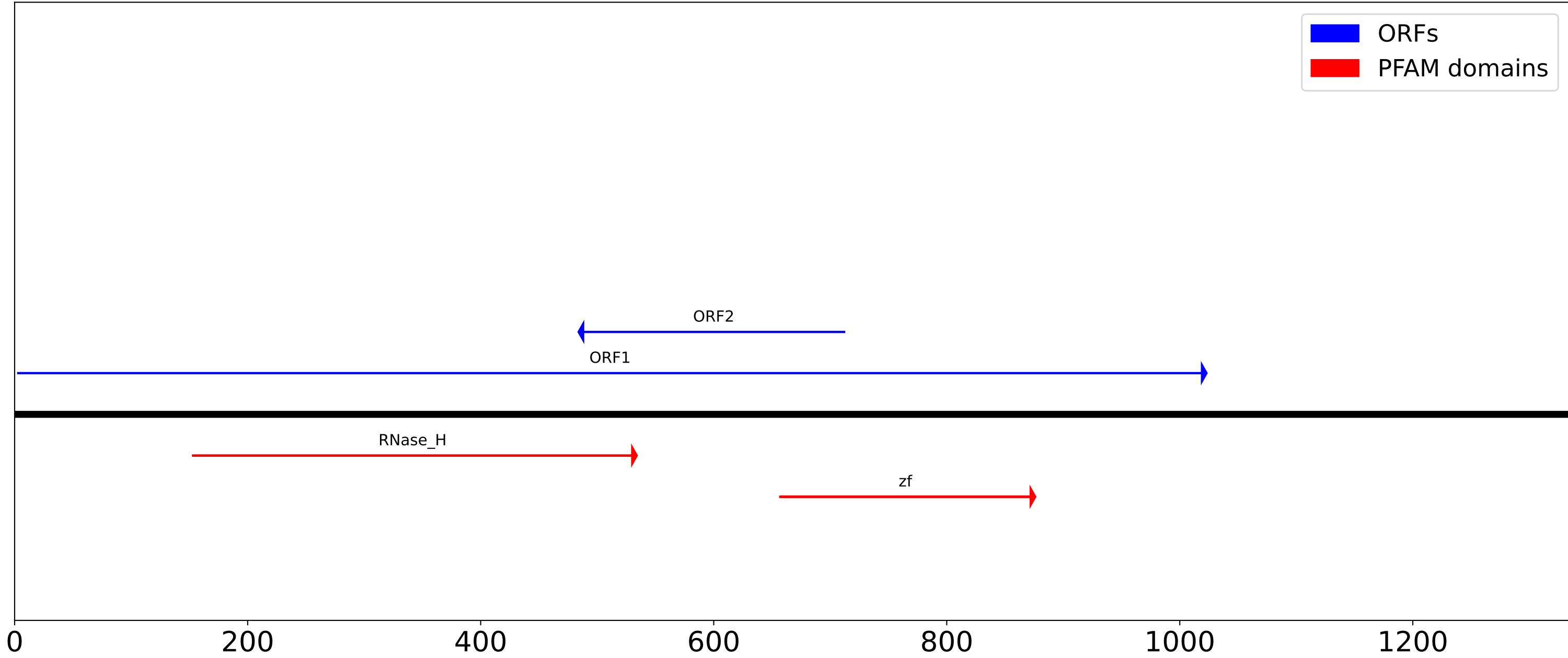
Before TEtrimmer 1335 bp



After TEtrimmer ORF and PFAM domain plot



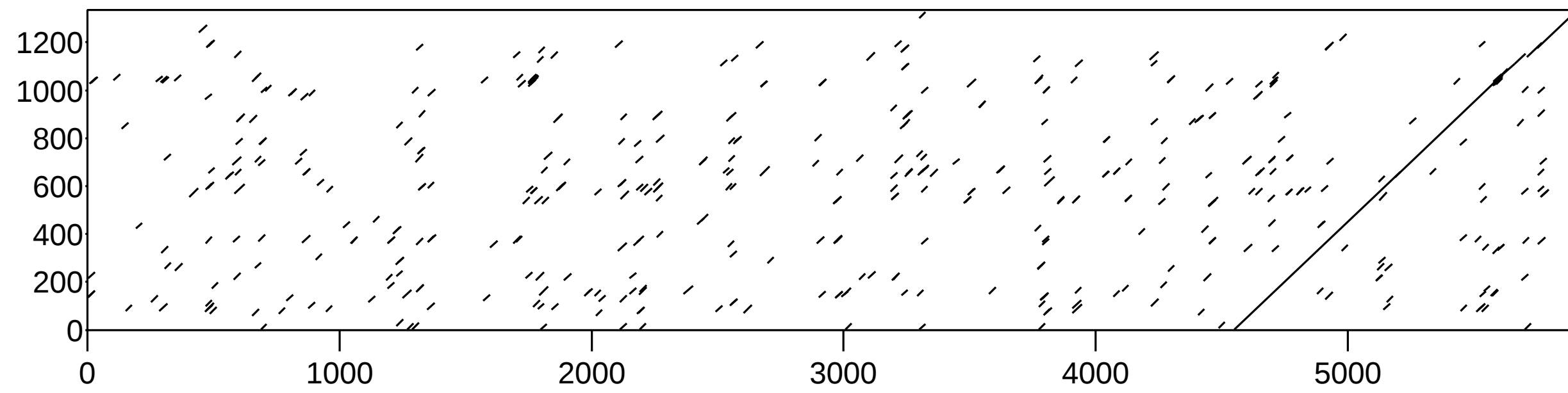
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)