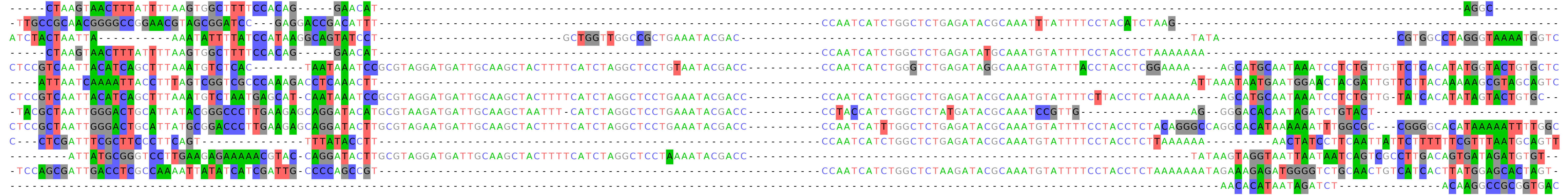


Start crop Point

End crop Point

MSA length = 1451



Start crop Point

End crop Point

0

10

0

250

500

750

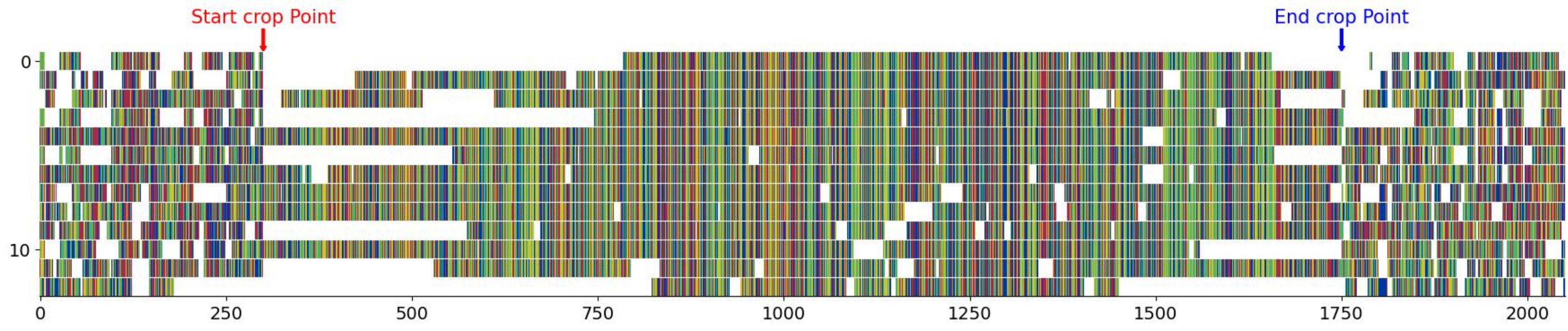
1000

1250

1500

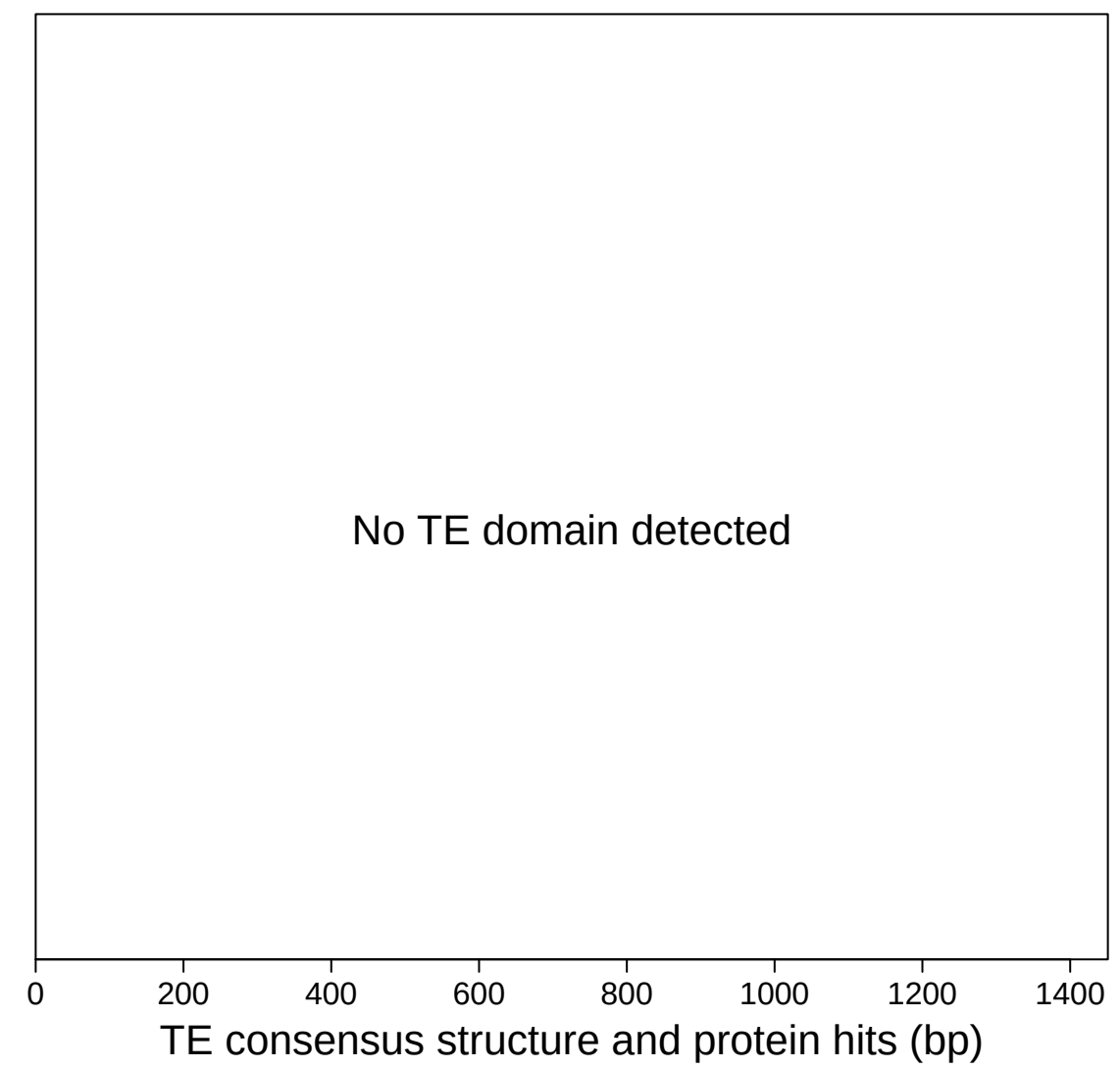
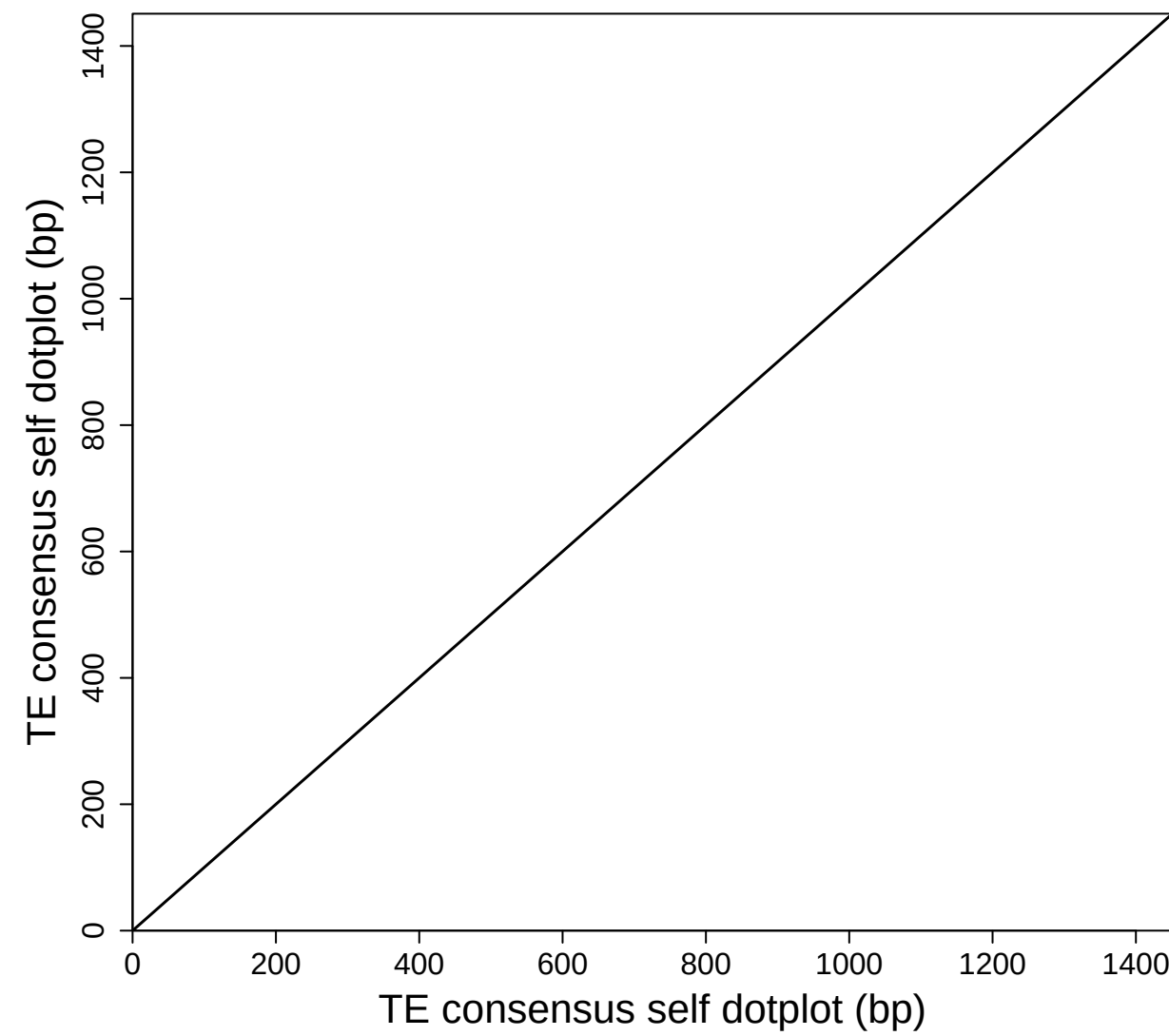
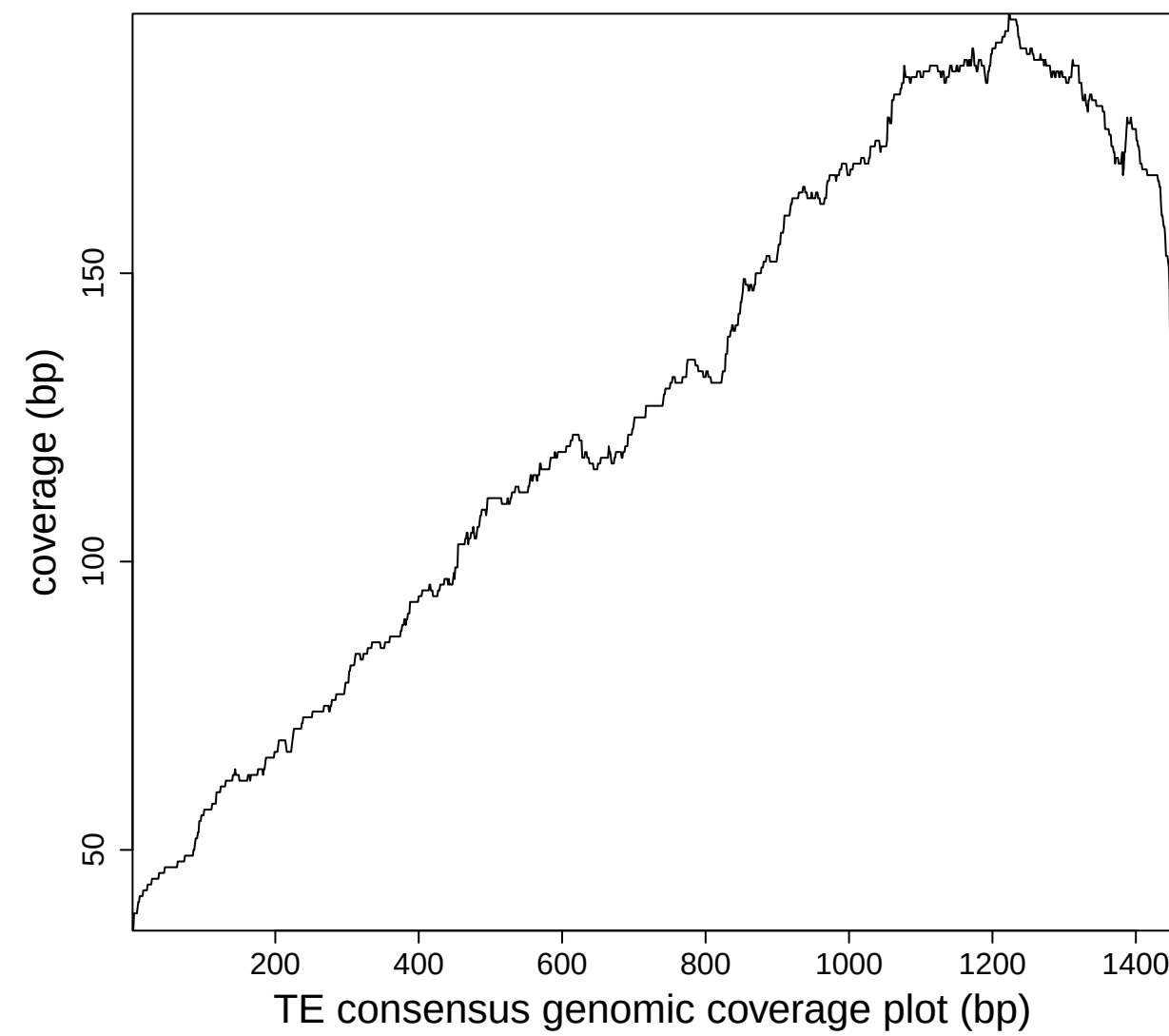
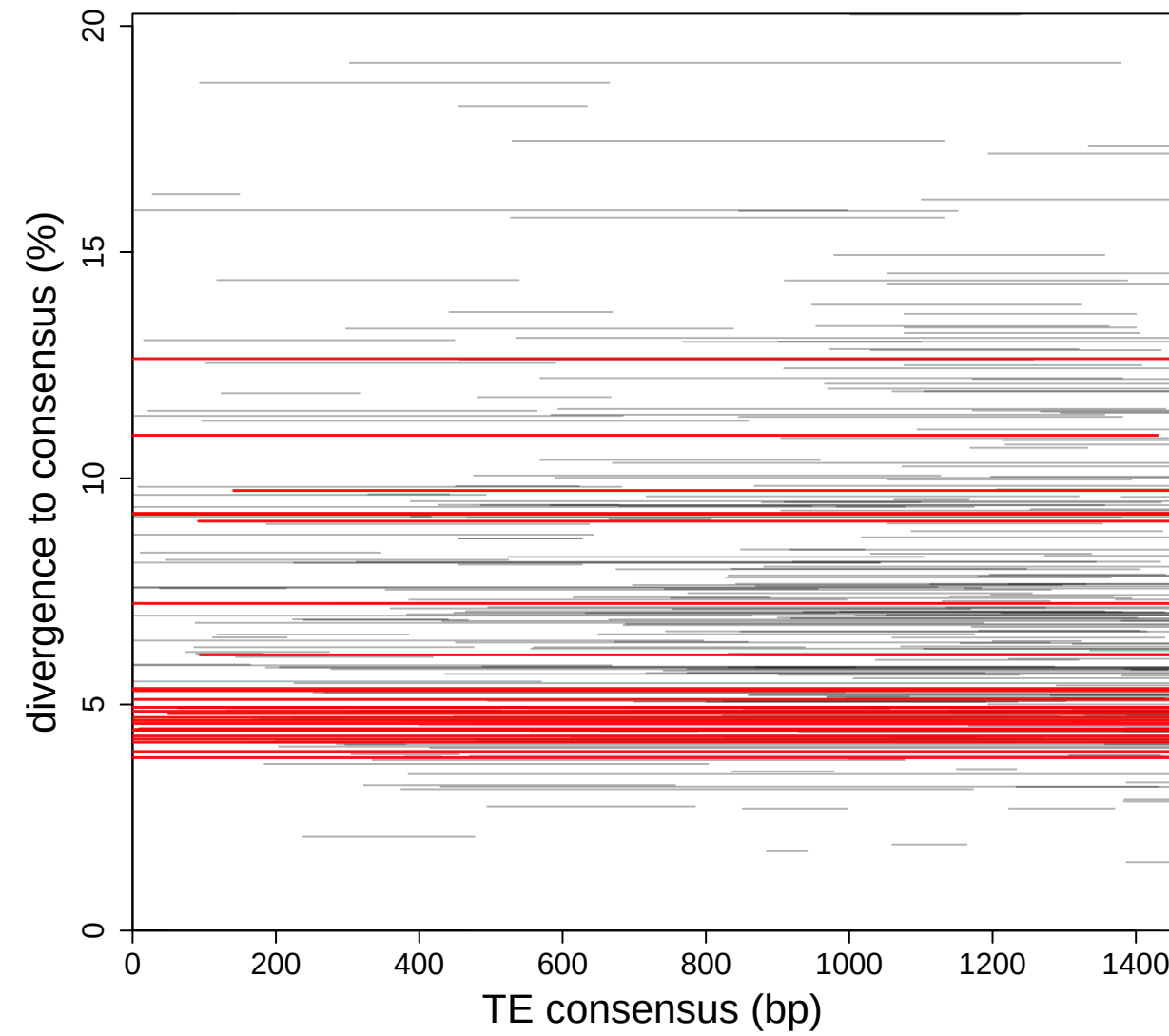
1750

2000



2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.
size: 1451bp; fragments: 325; full length: 29 (>=1305.9bp)

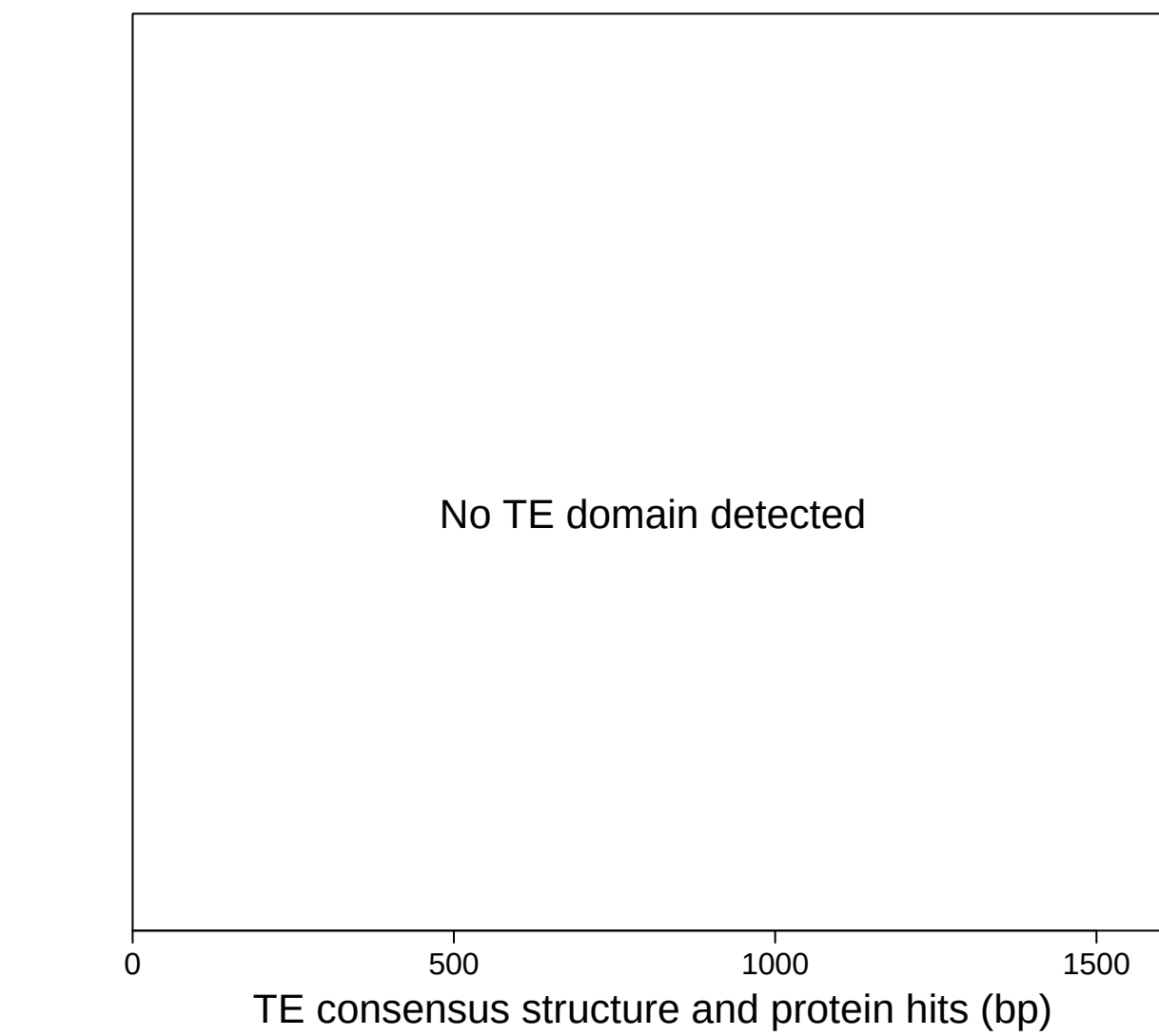
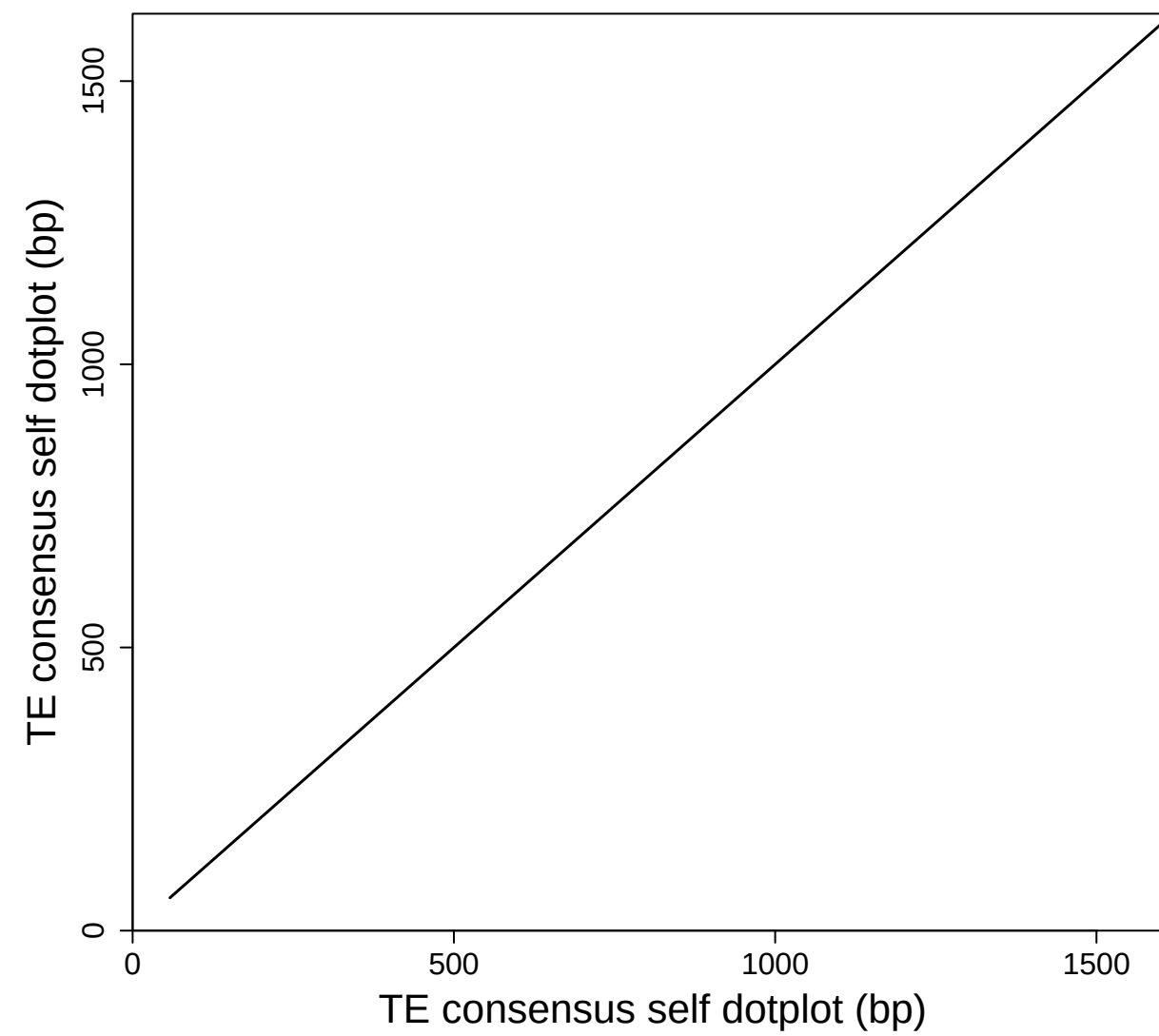
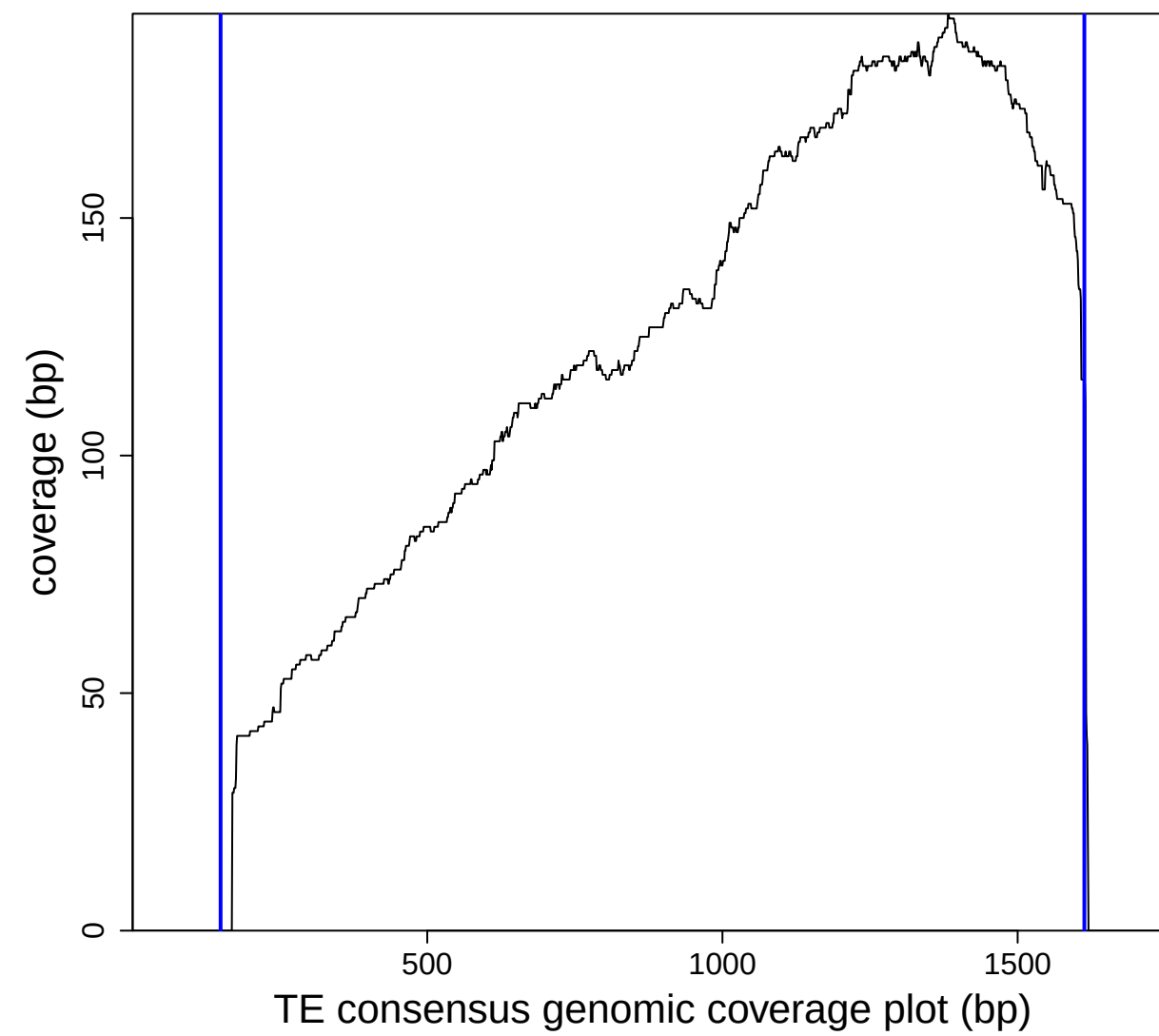
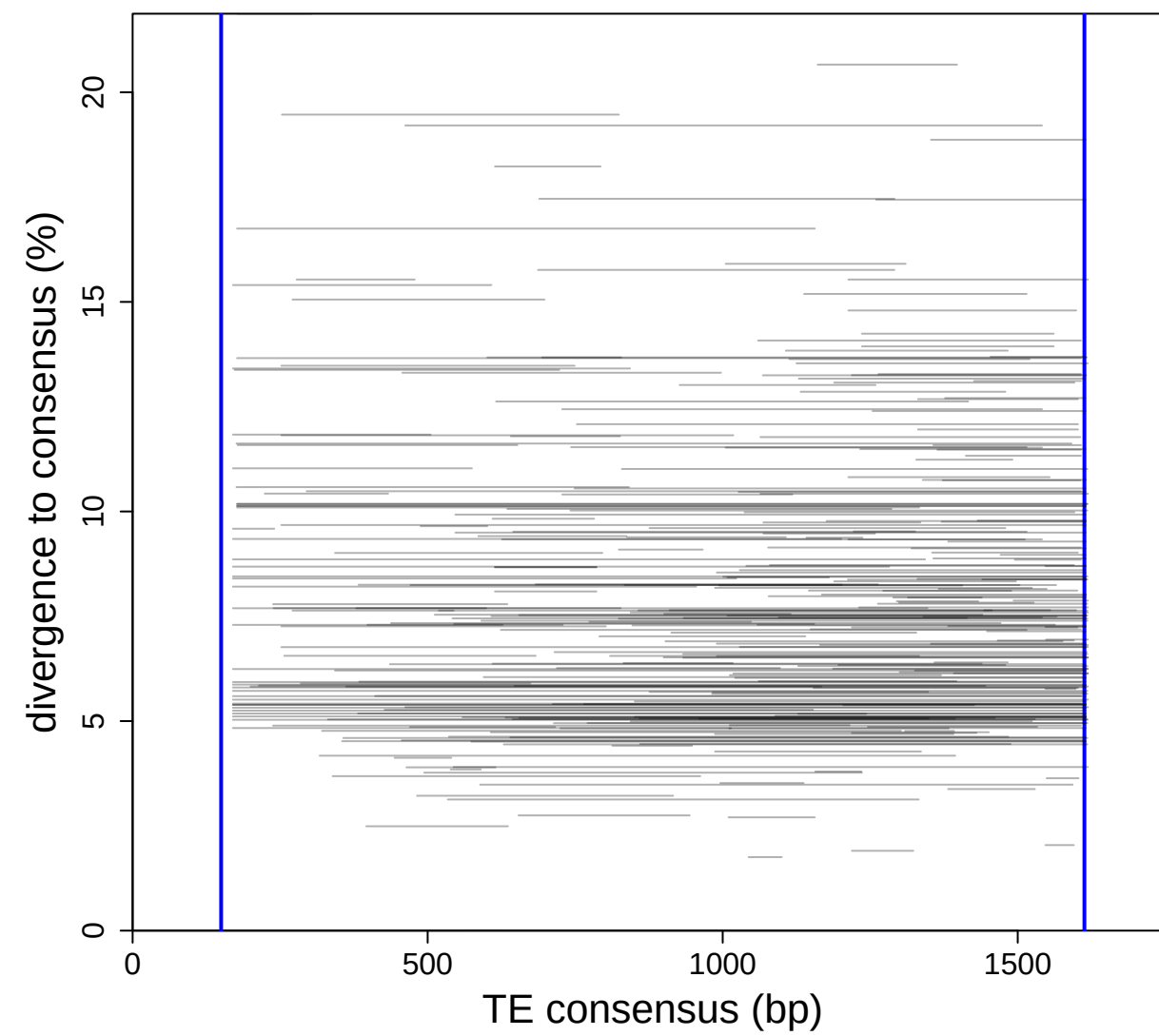
After TEtrimmer 1451 bp



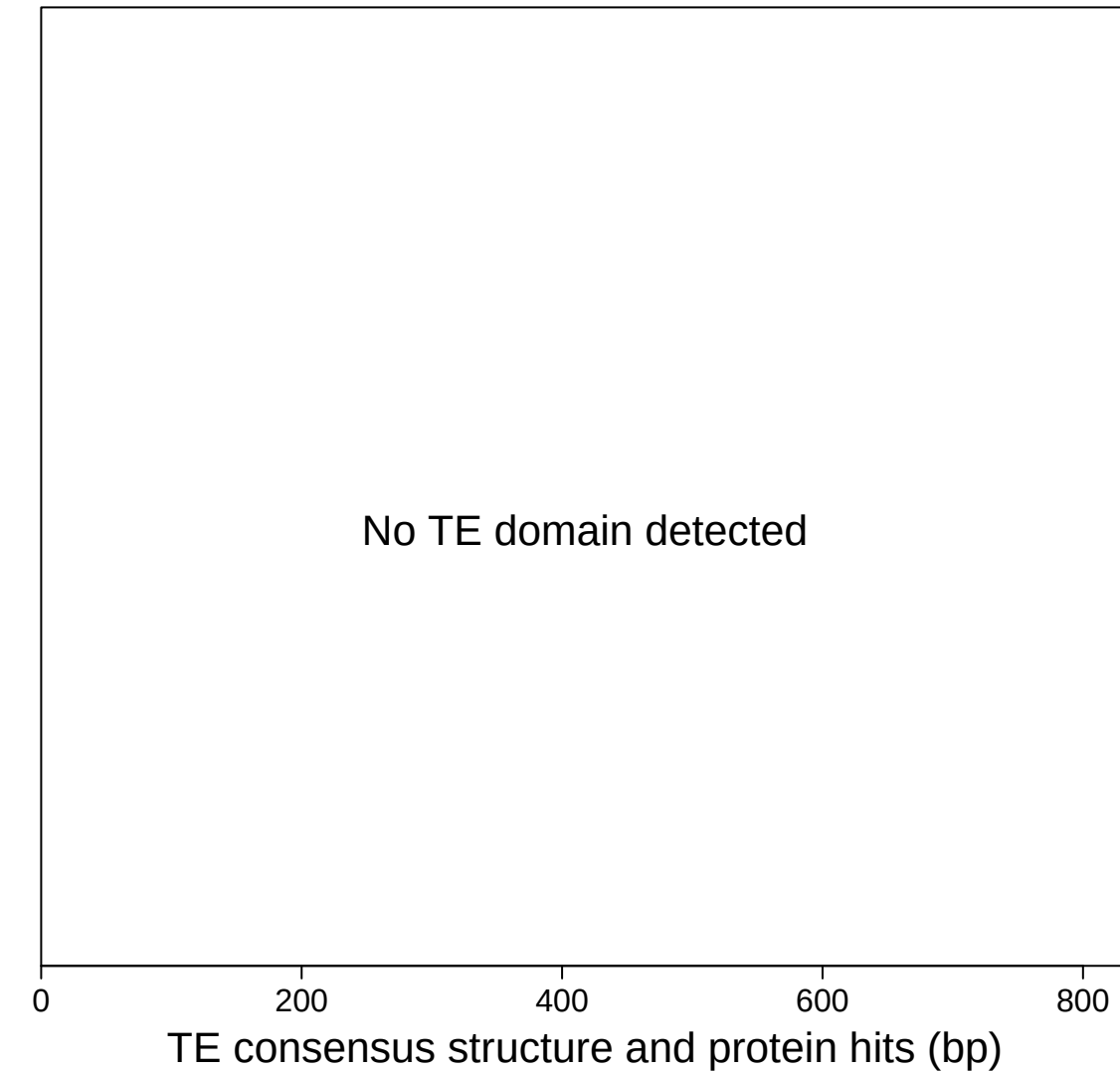
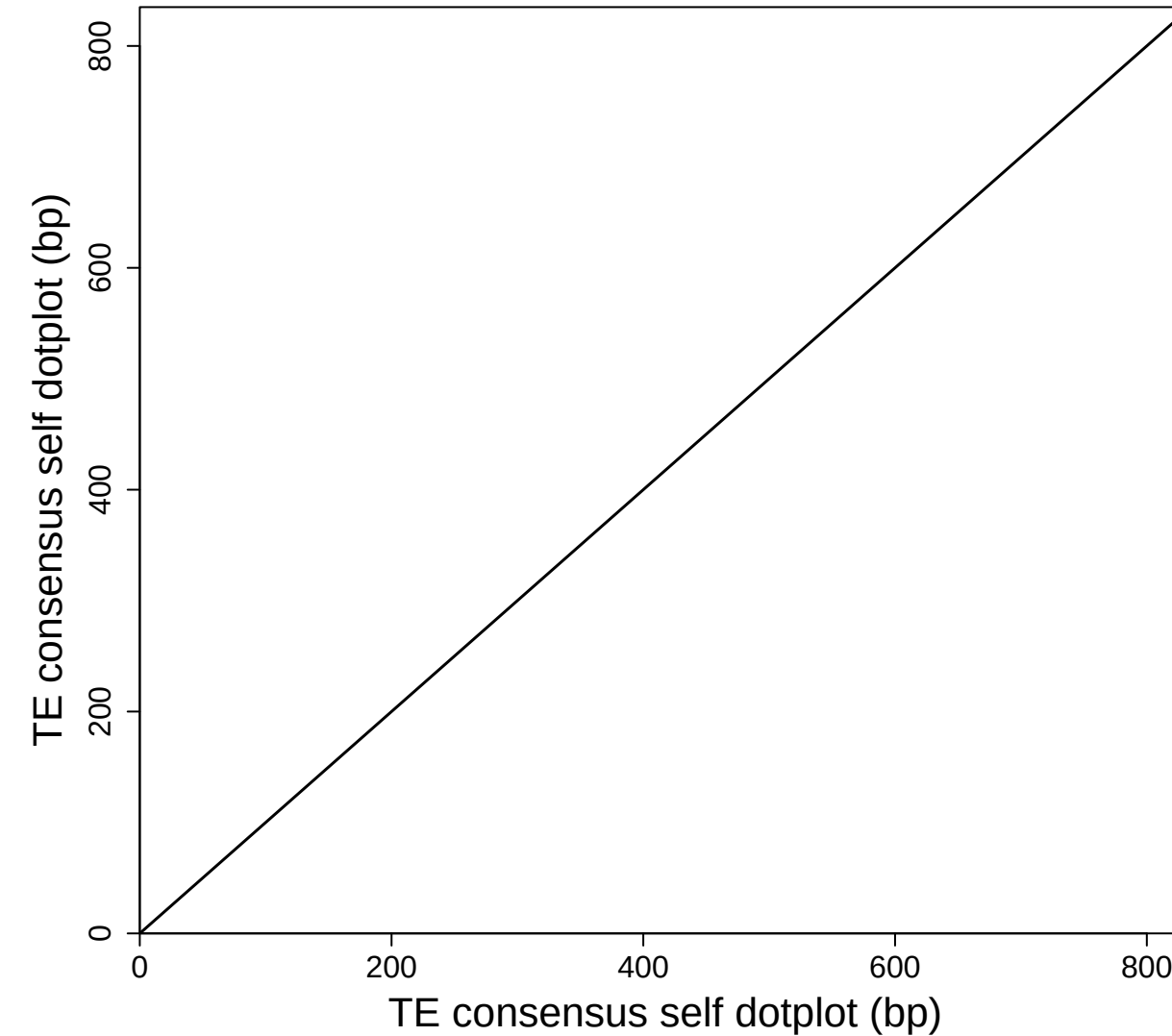
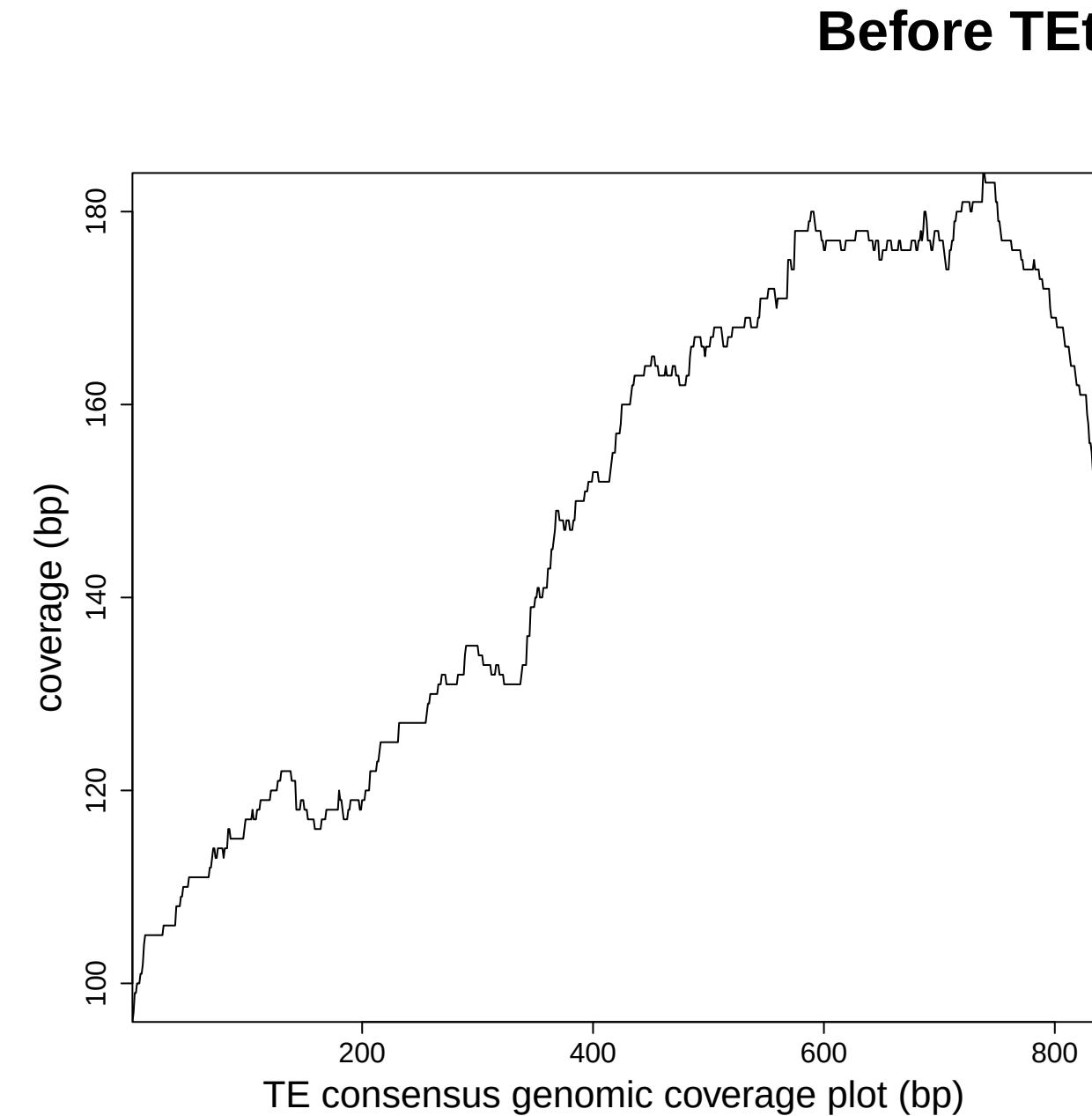
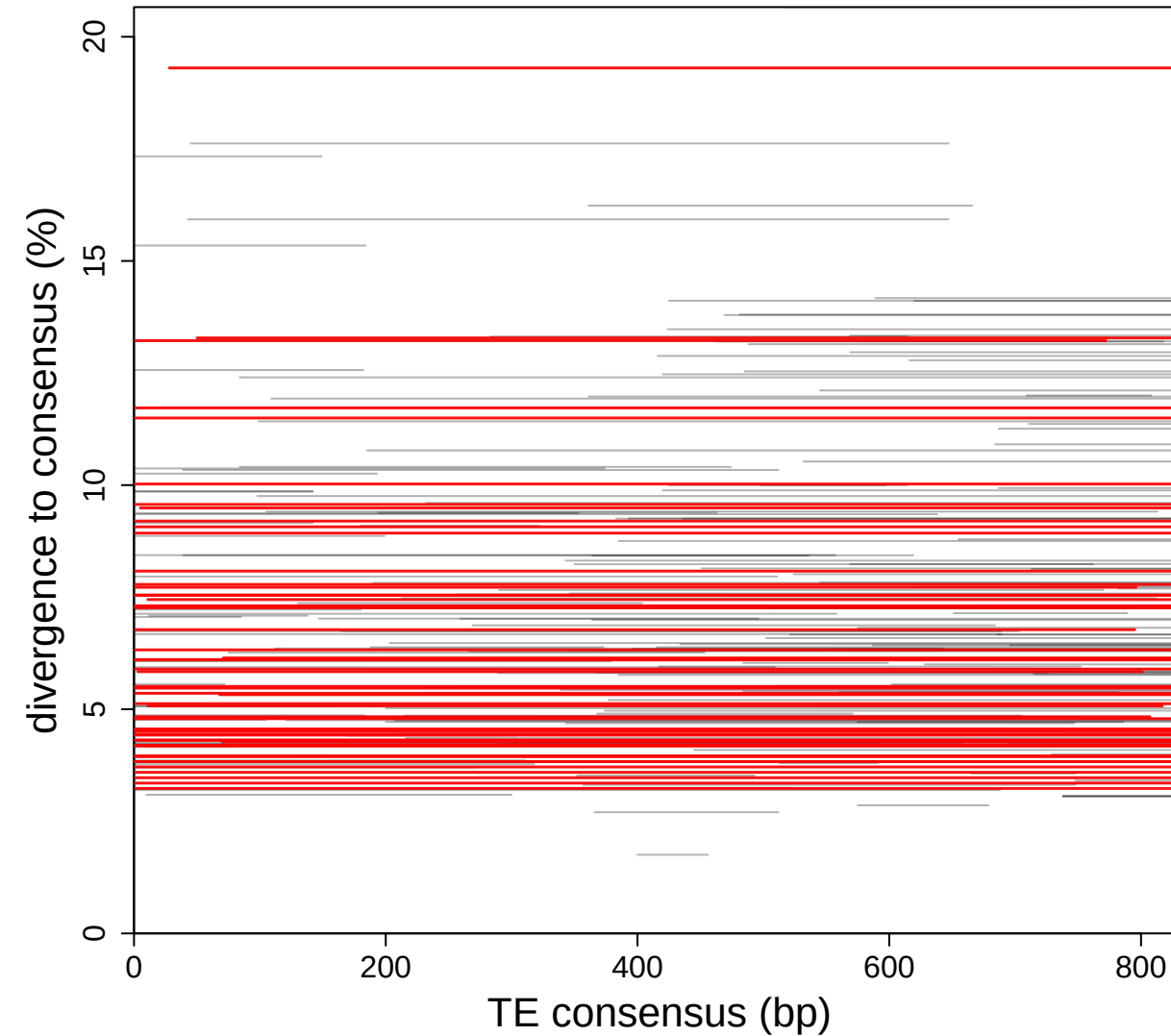
uf.bed g 2.bed fm 1.bed 0 0 n.bed g 1.bed fm 2.bed 0 0 b
size: 1763bp; fragments: 304; full length: 0 (>=1586.7bp)

After TEtrimmer Extended plot

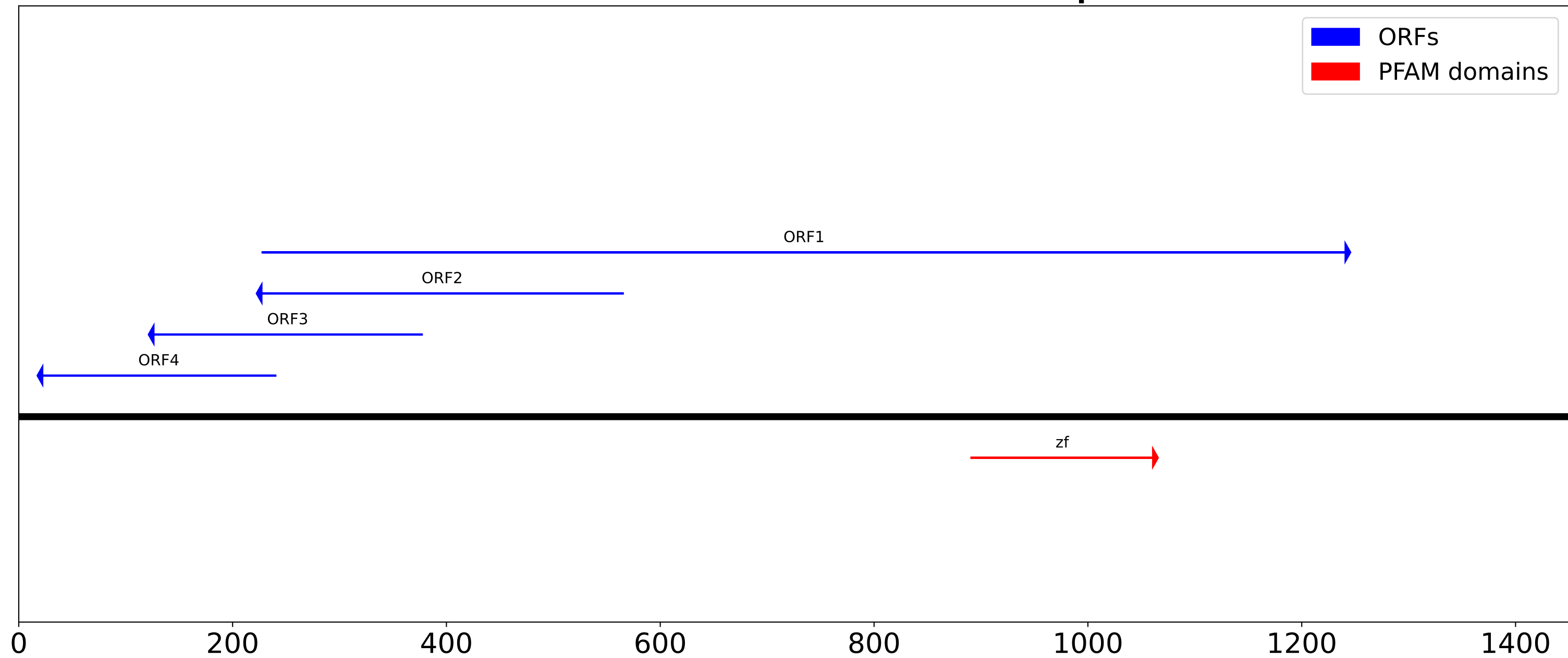
Blue lines are boundaries



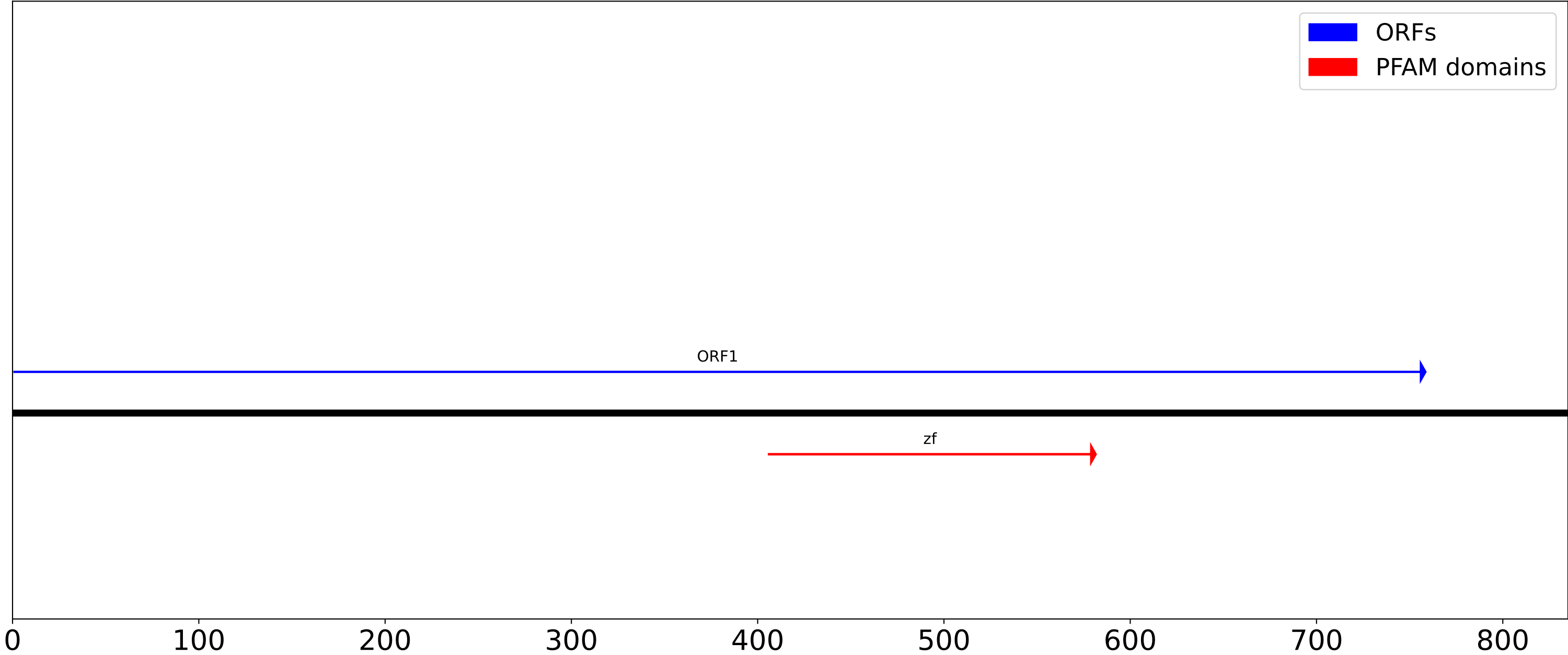
TE: rnd_5_family_19694
size: 835bp; fragments: 256; full length: 72 (≥ 751.5 bp)



After TEtrimmer ORF and PFAM domain plot



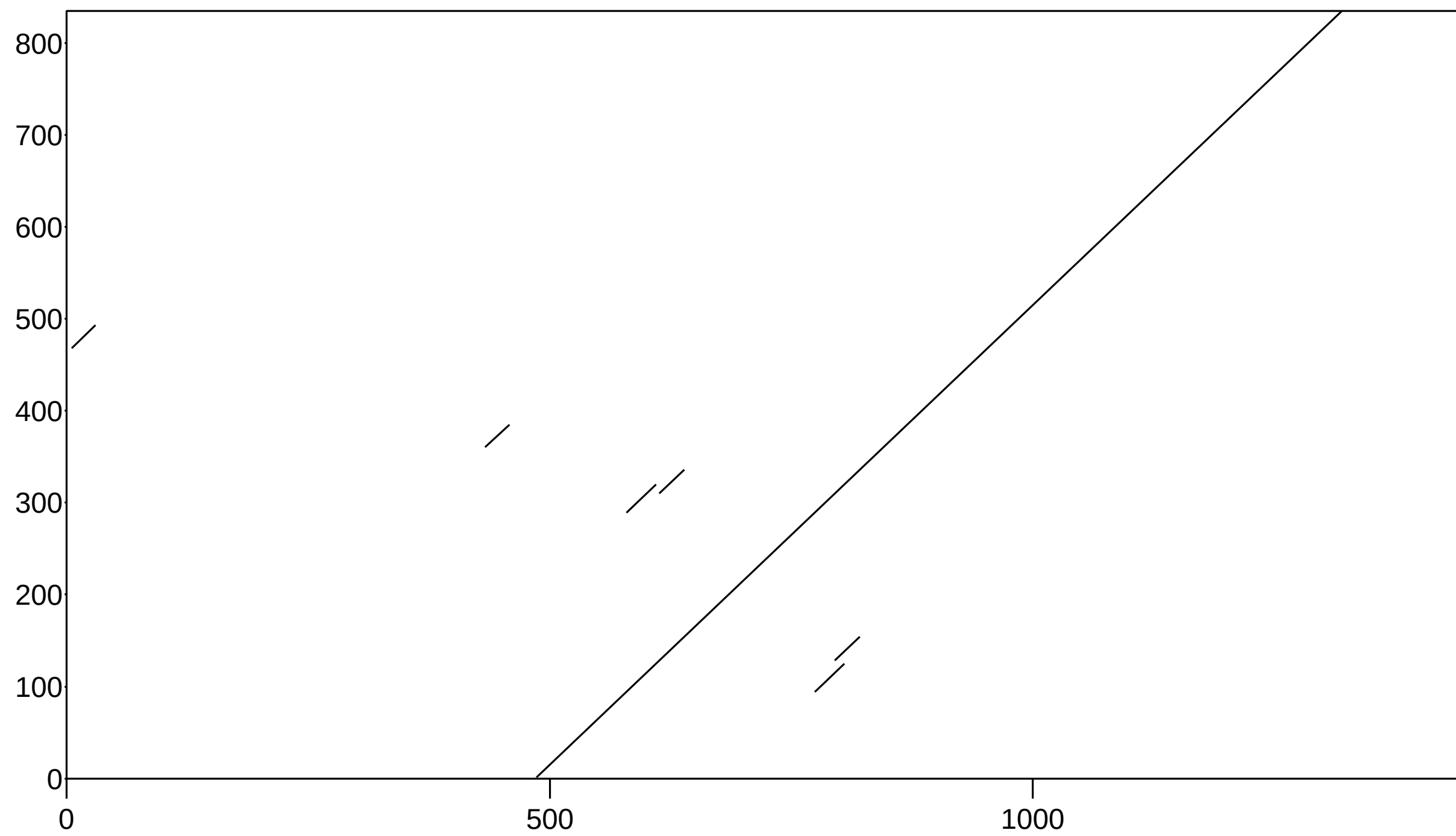
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)