

Start crop Point

End crop Point

0

10

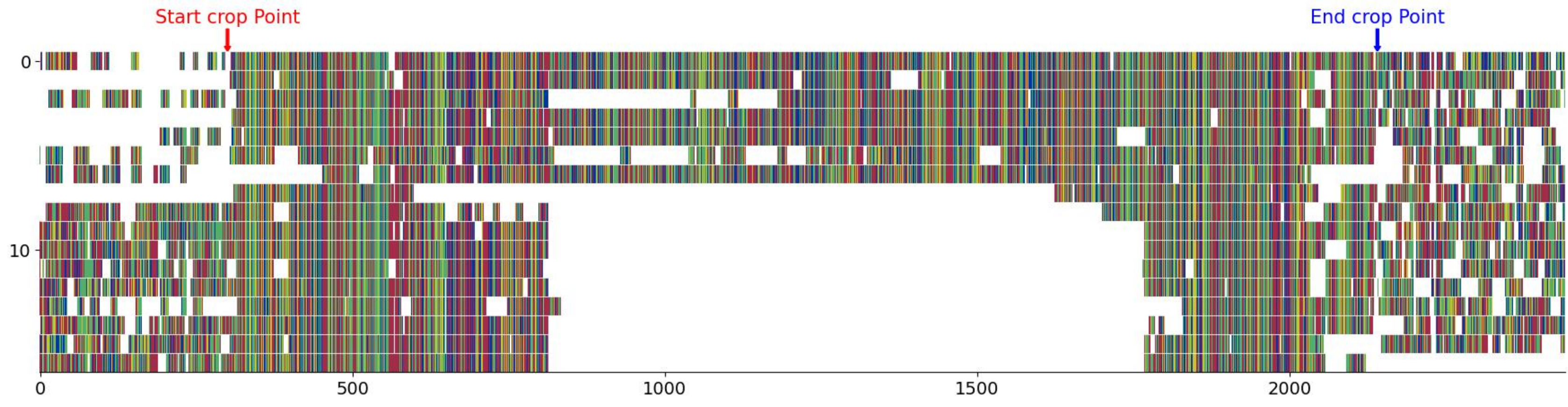
0

500

1000

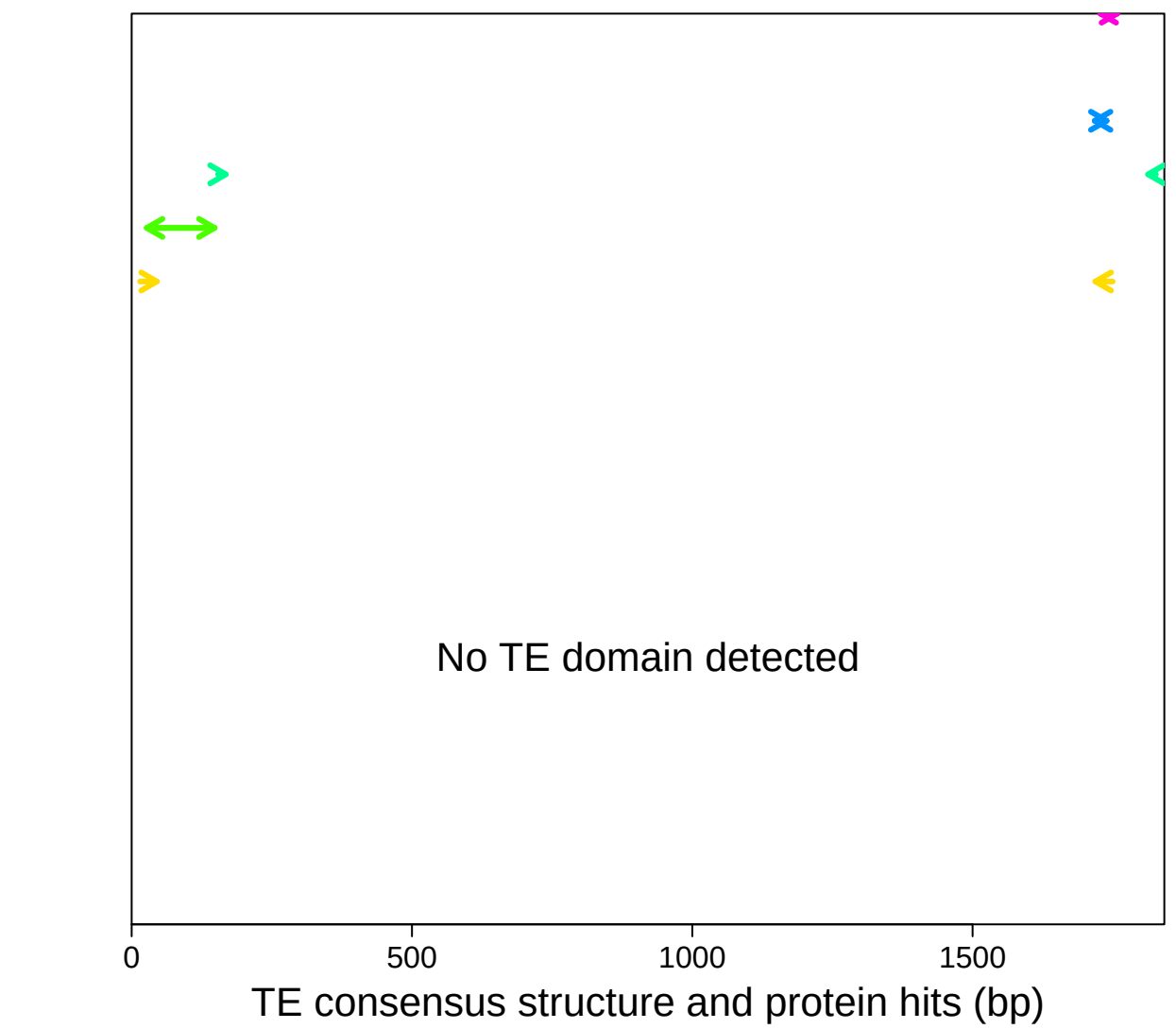
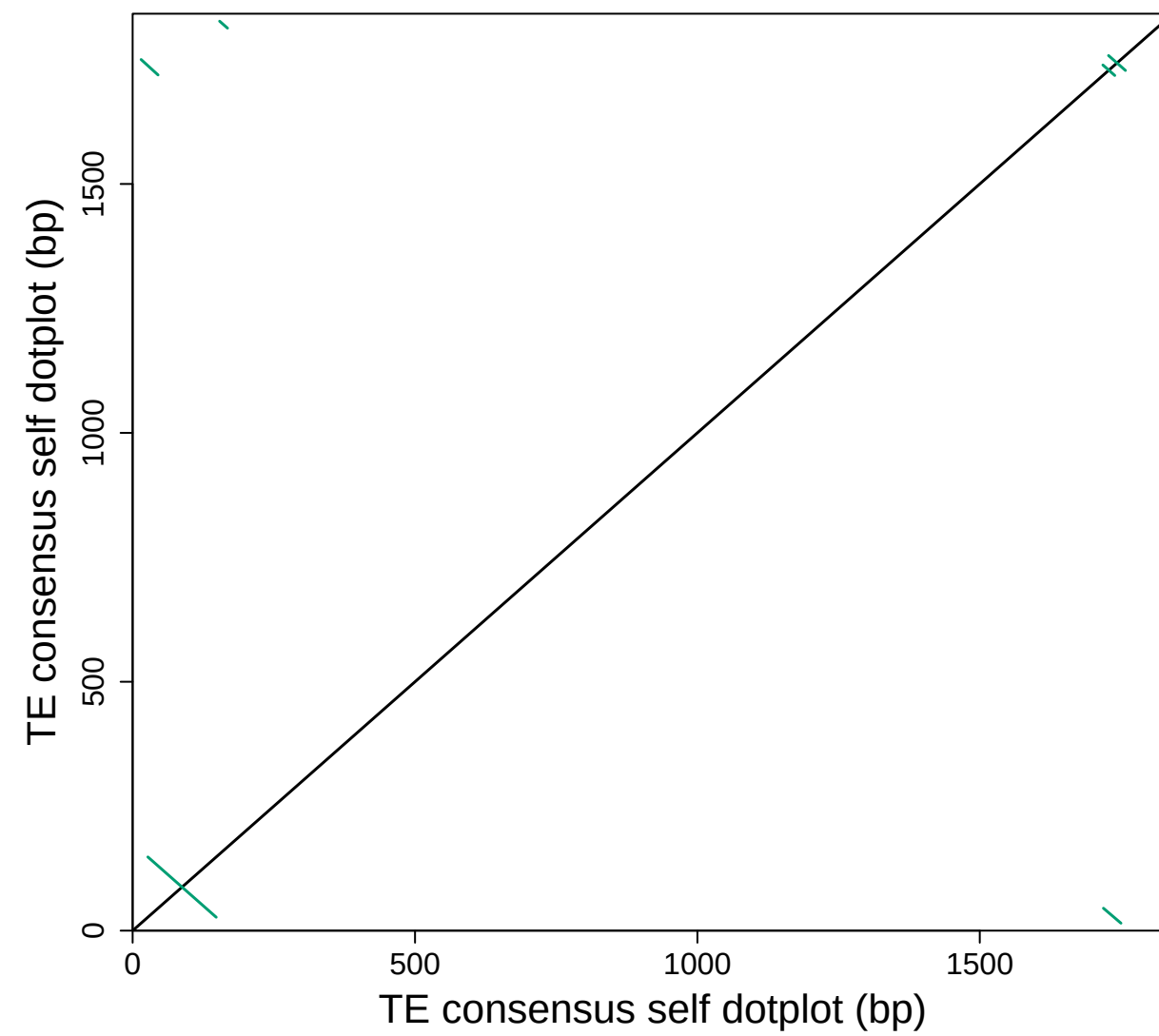
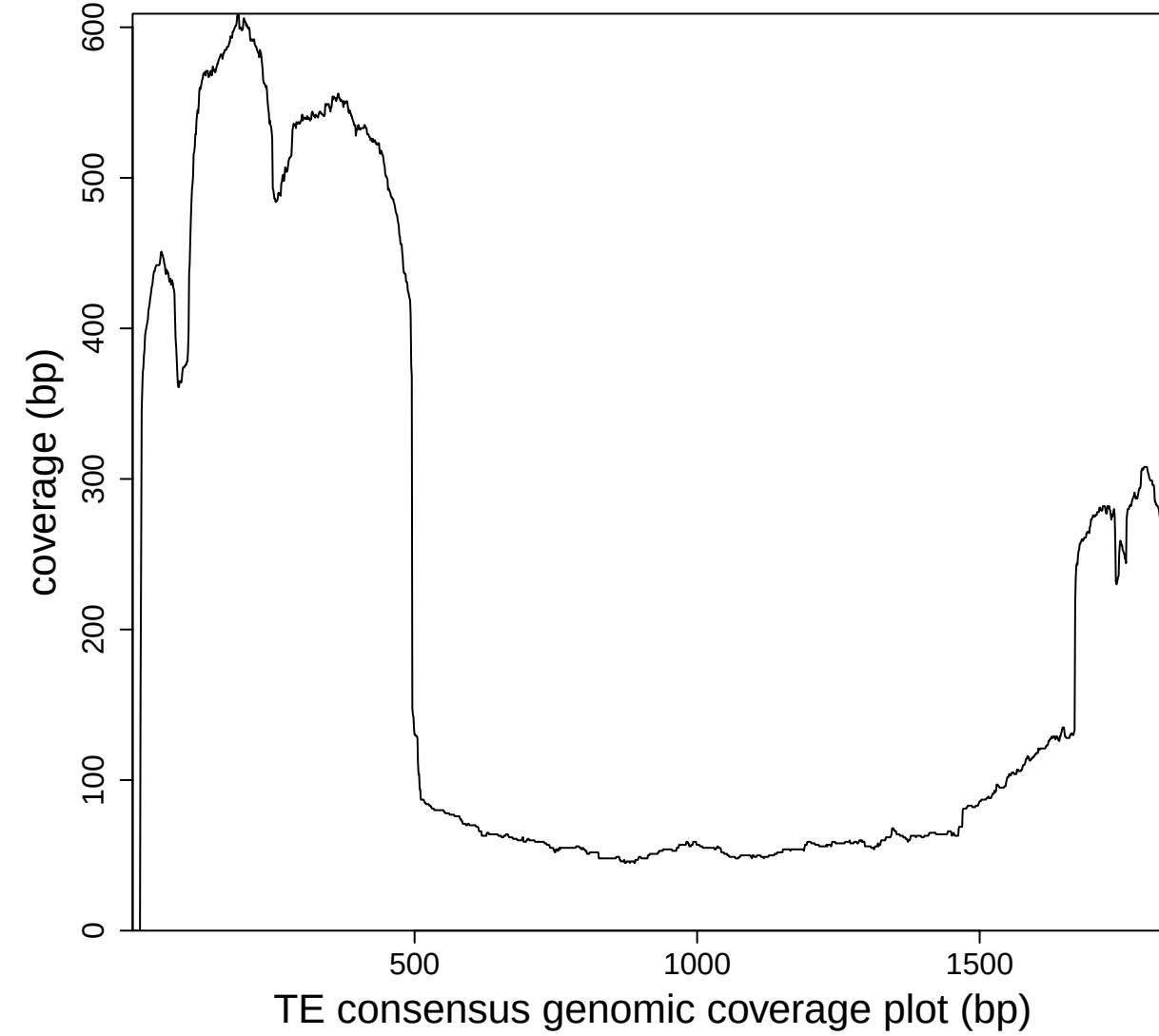
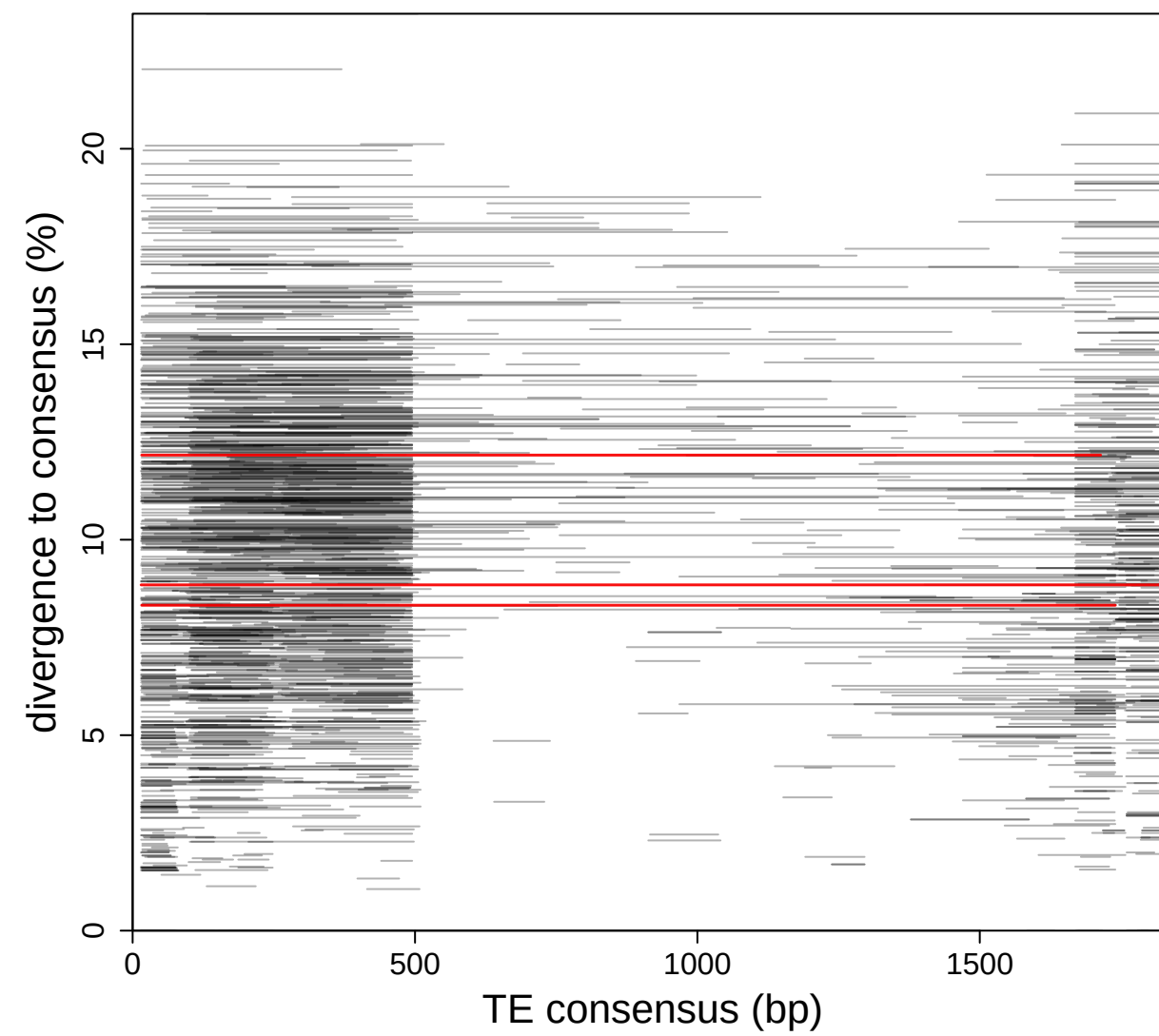
1500

2000



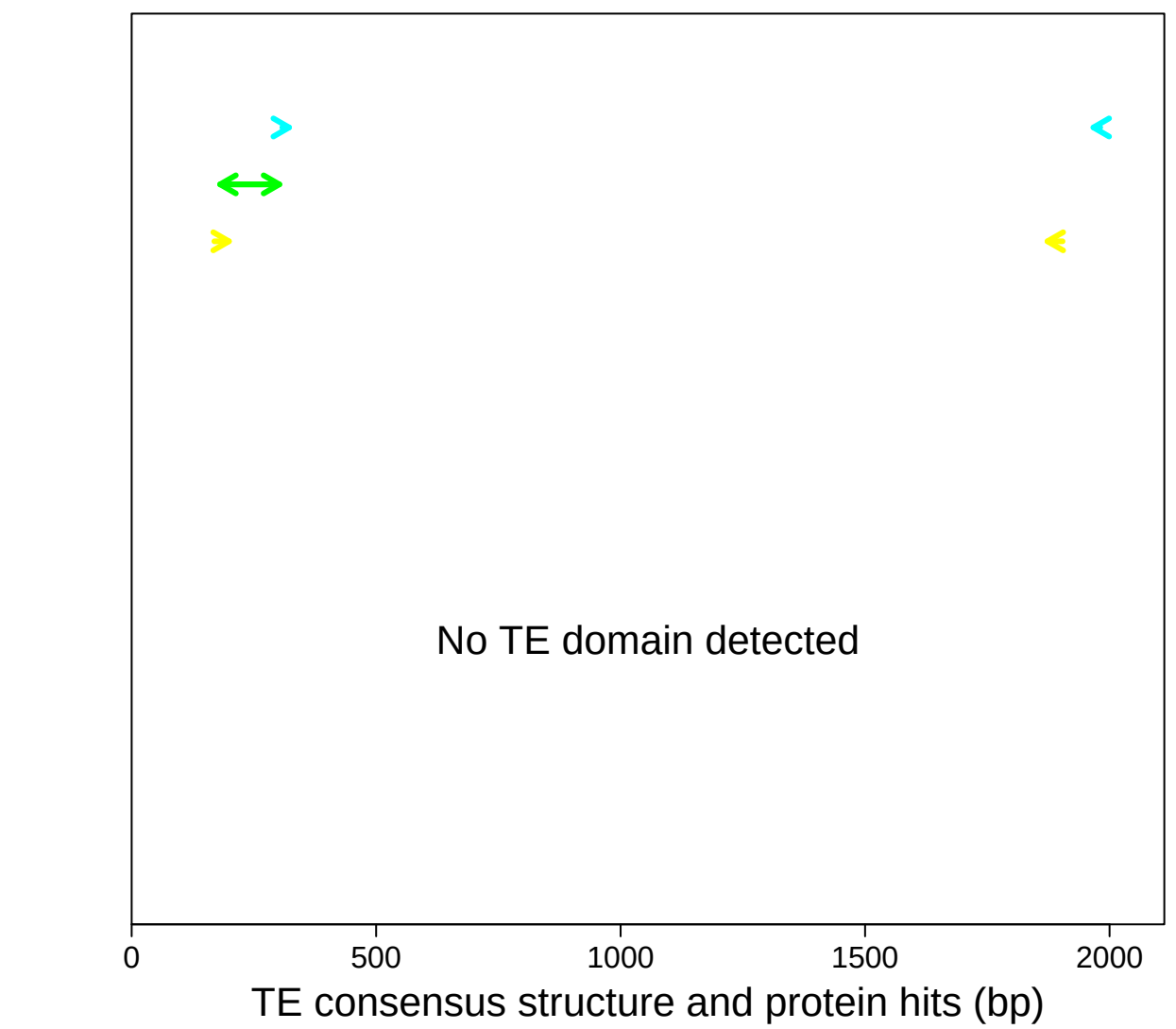
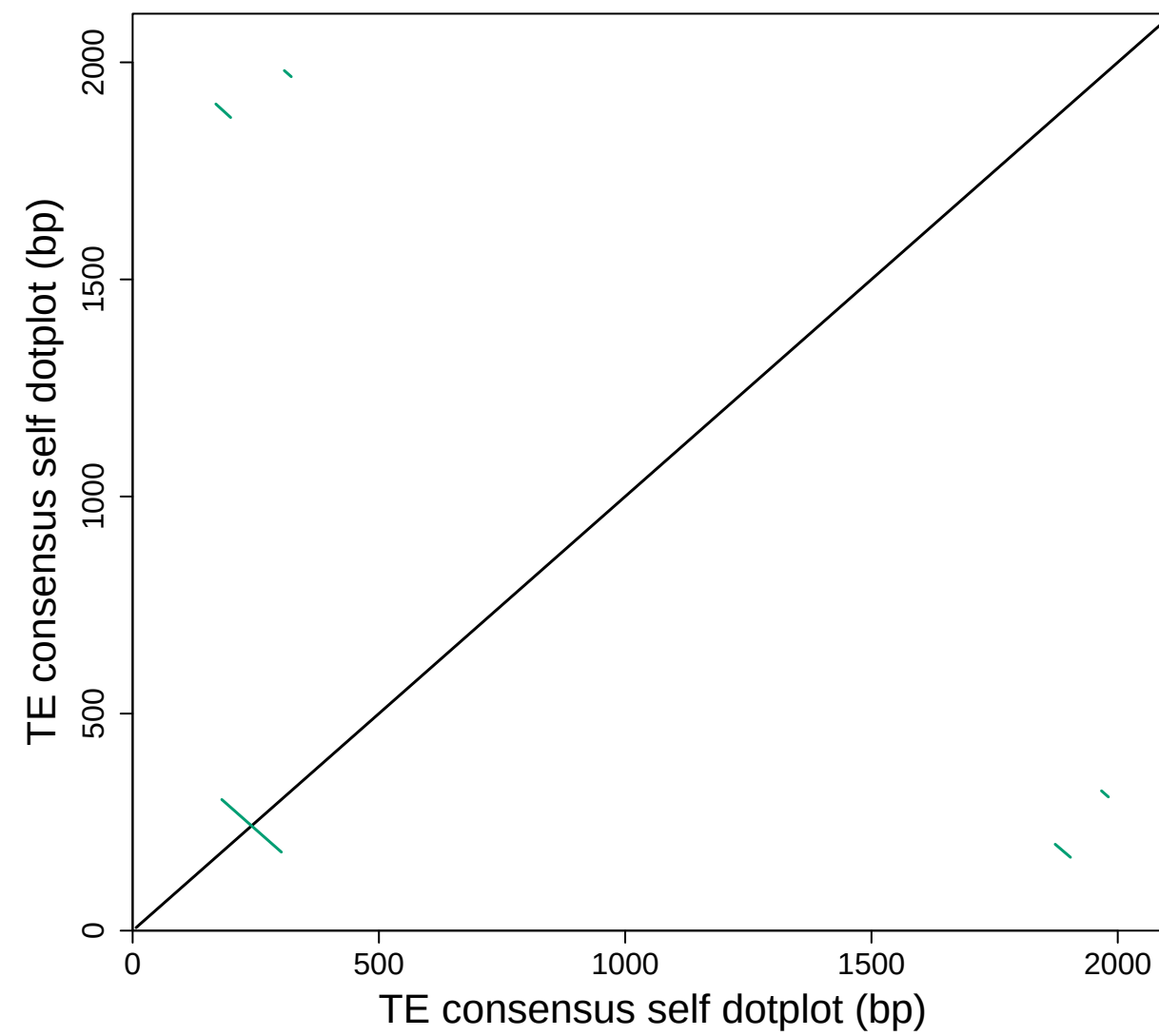
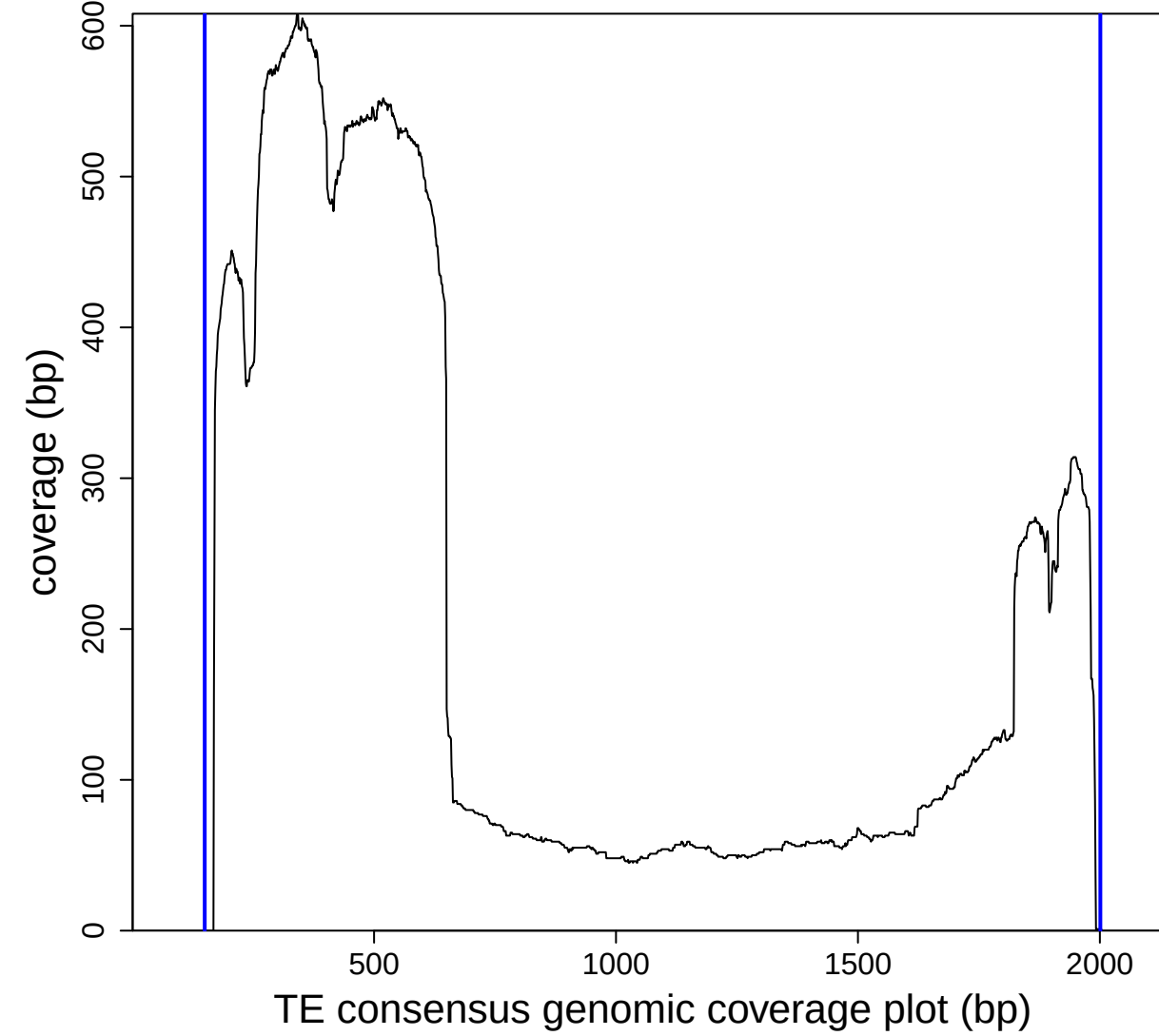
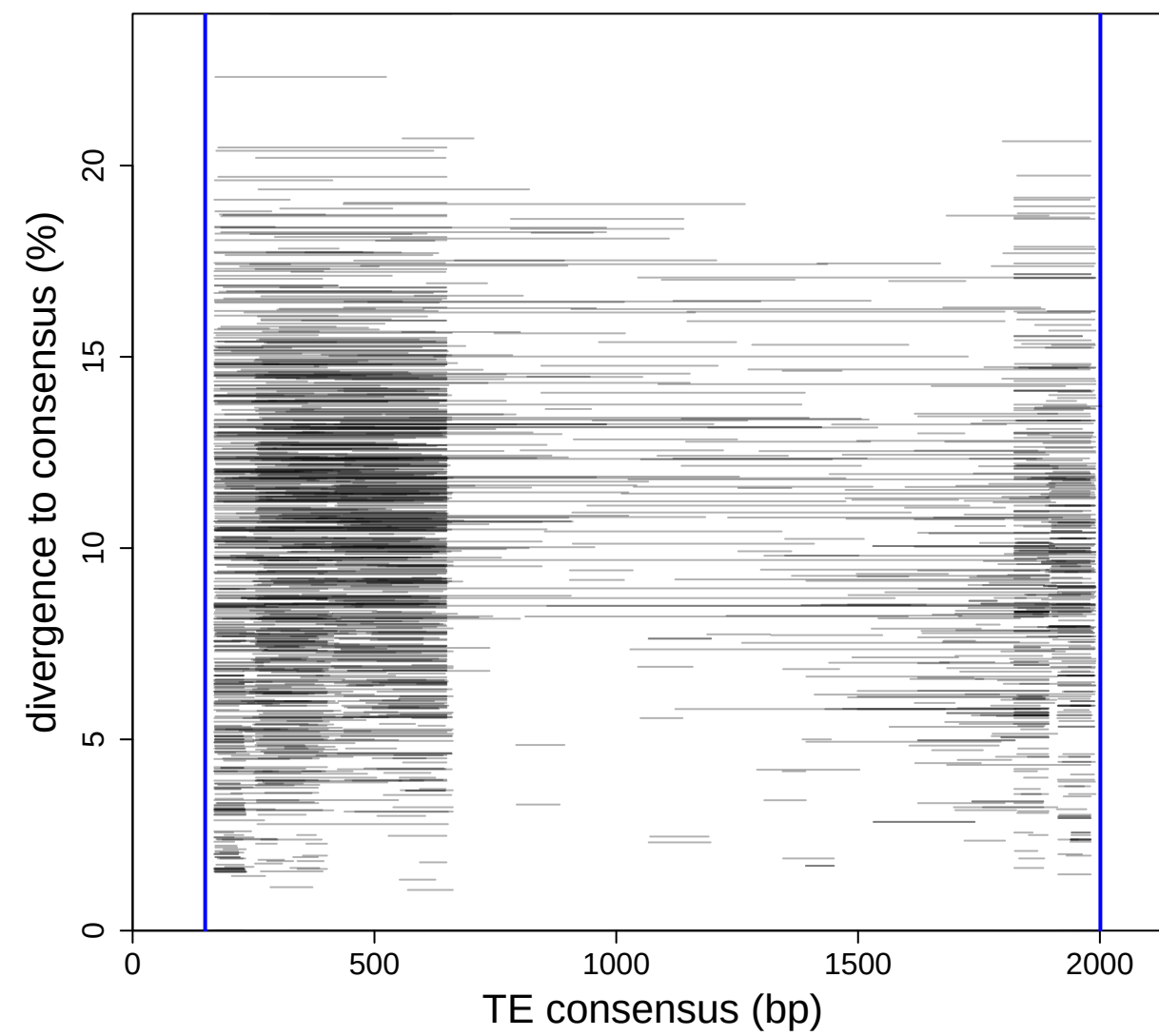
g_1.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0 0 bcln.fa ε
size: 1842bp; fragments: 1634; full length: 3 (>=1657.8bp)

After TETrimmer 1842 bp

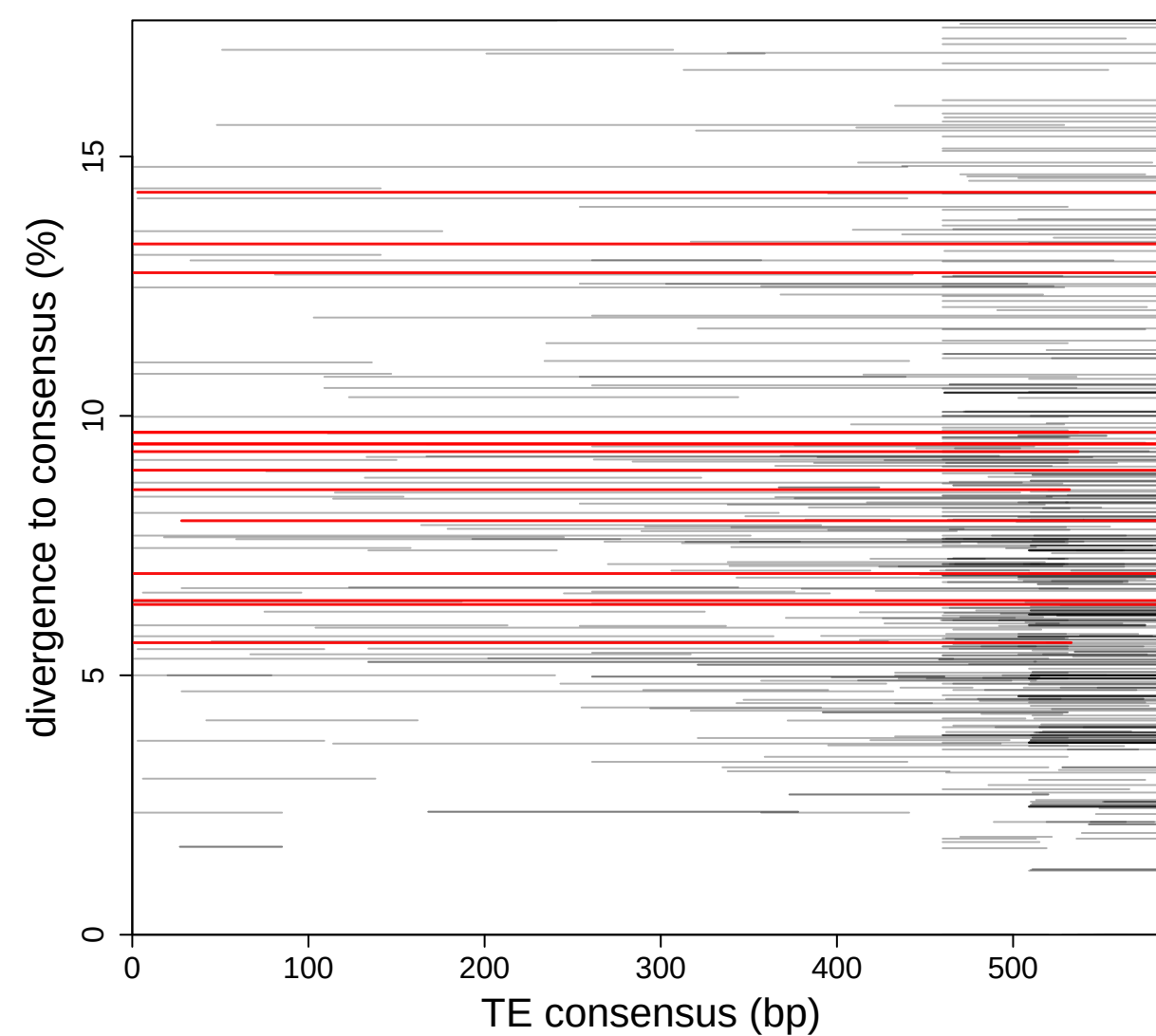


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 2151bp; fragments: 1630; full length: 0 (>=1935.9bp)

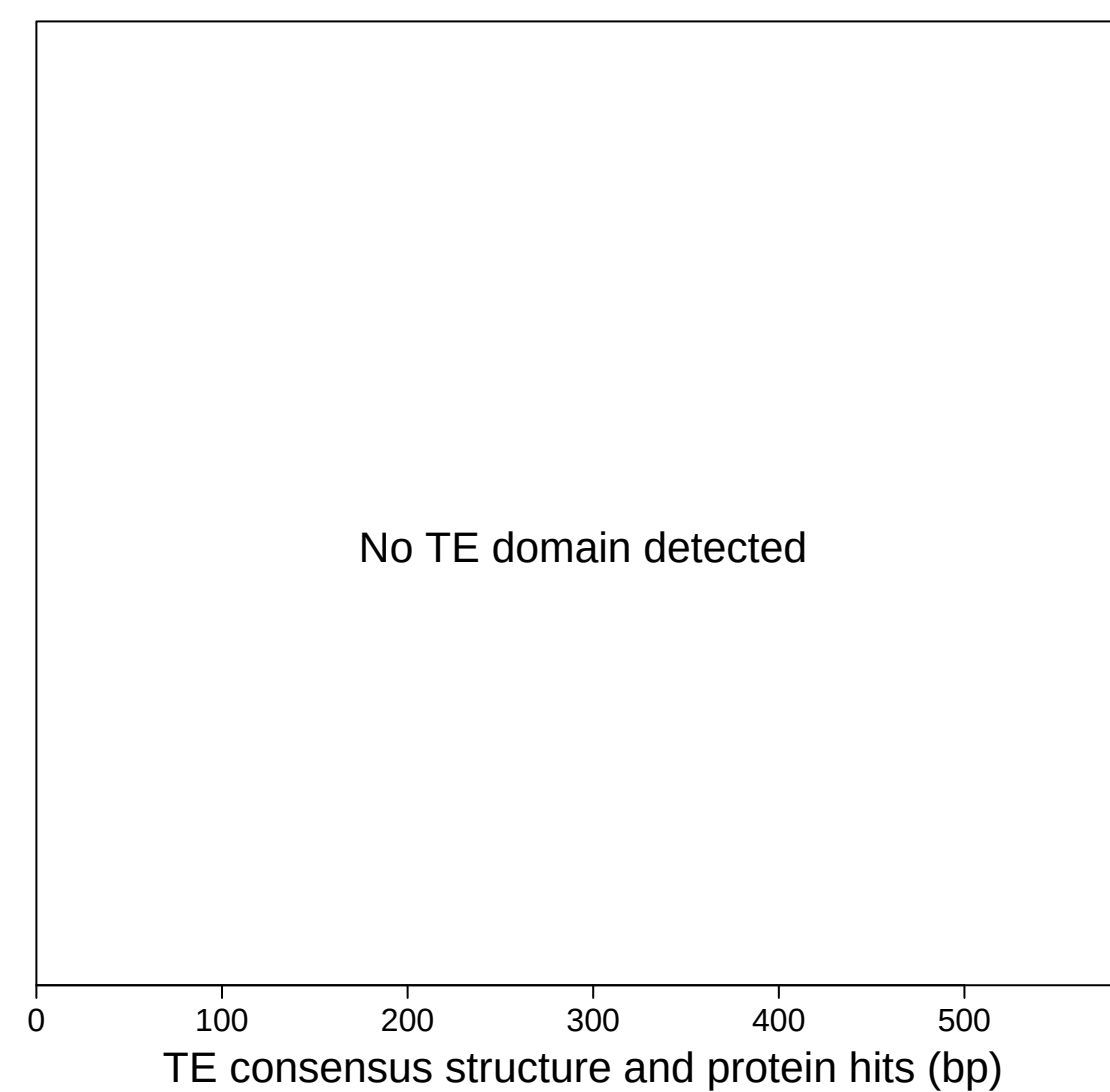
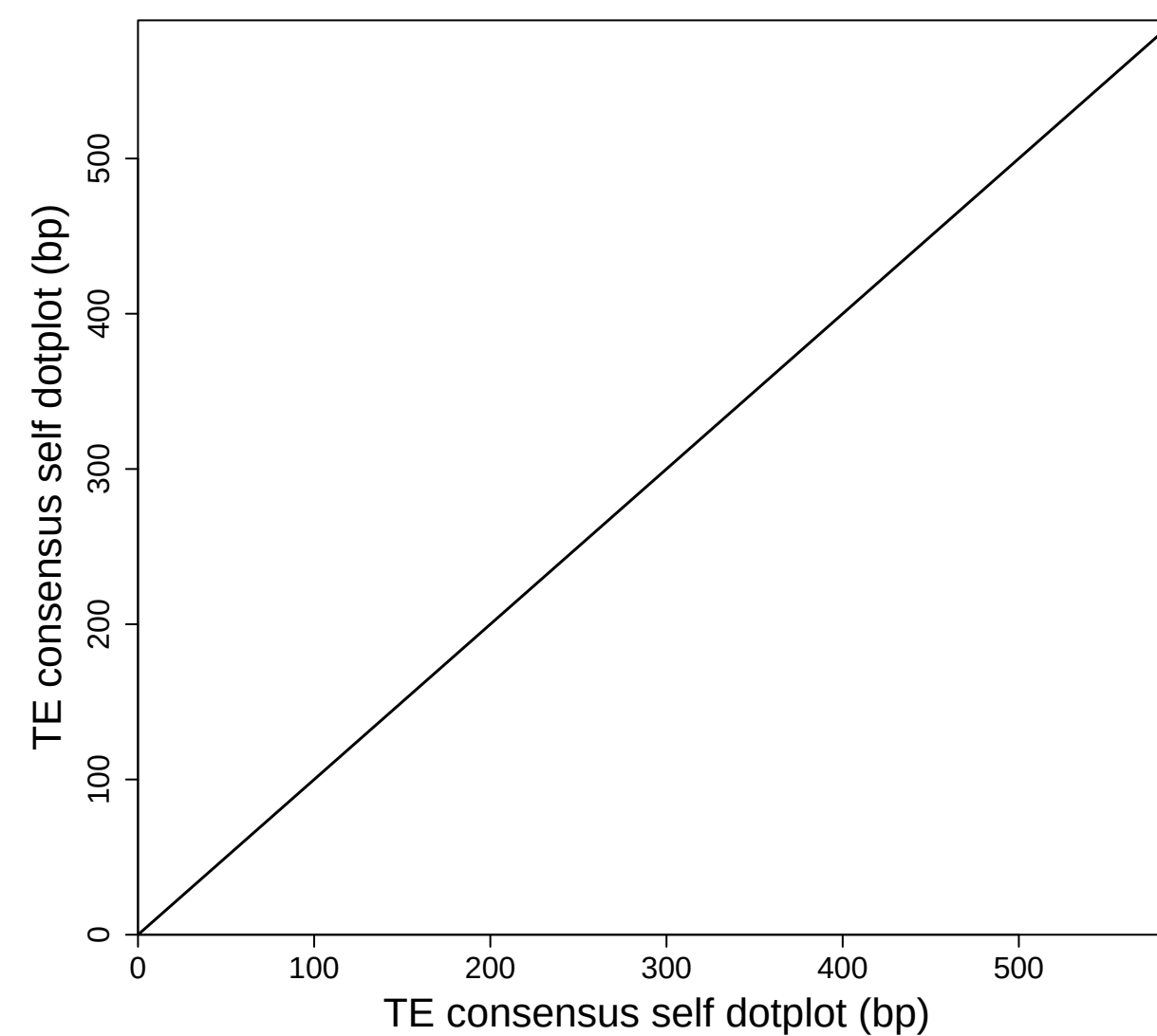
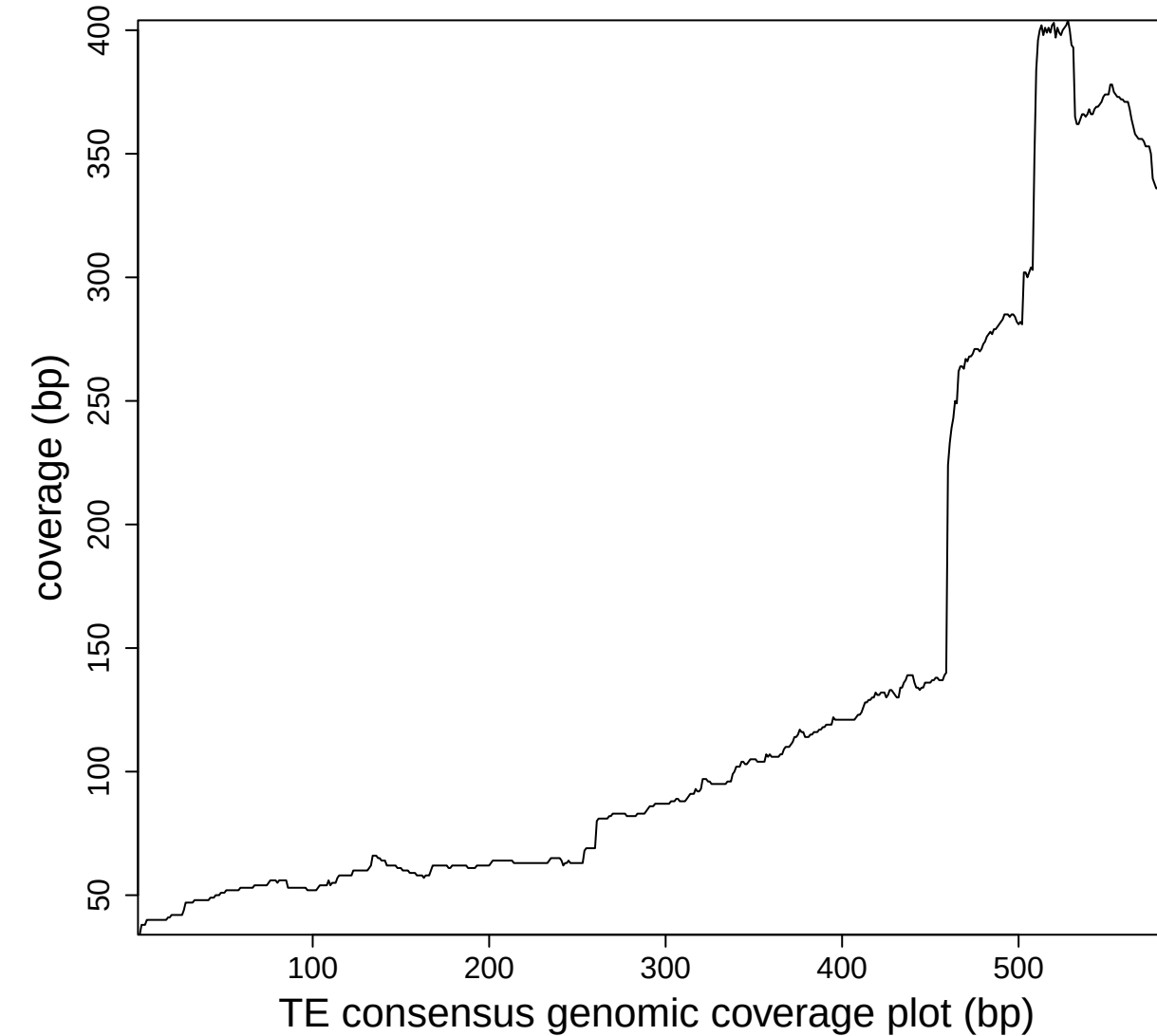
After TEtrimmer Extended plot Blue lines are boundaries



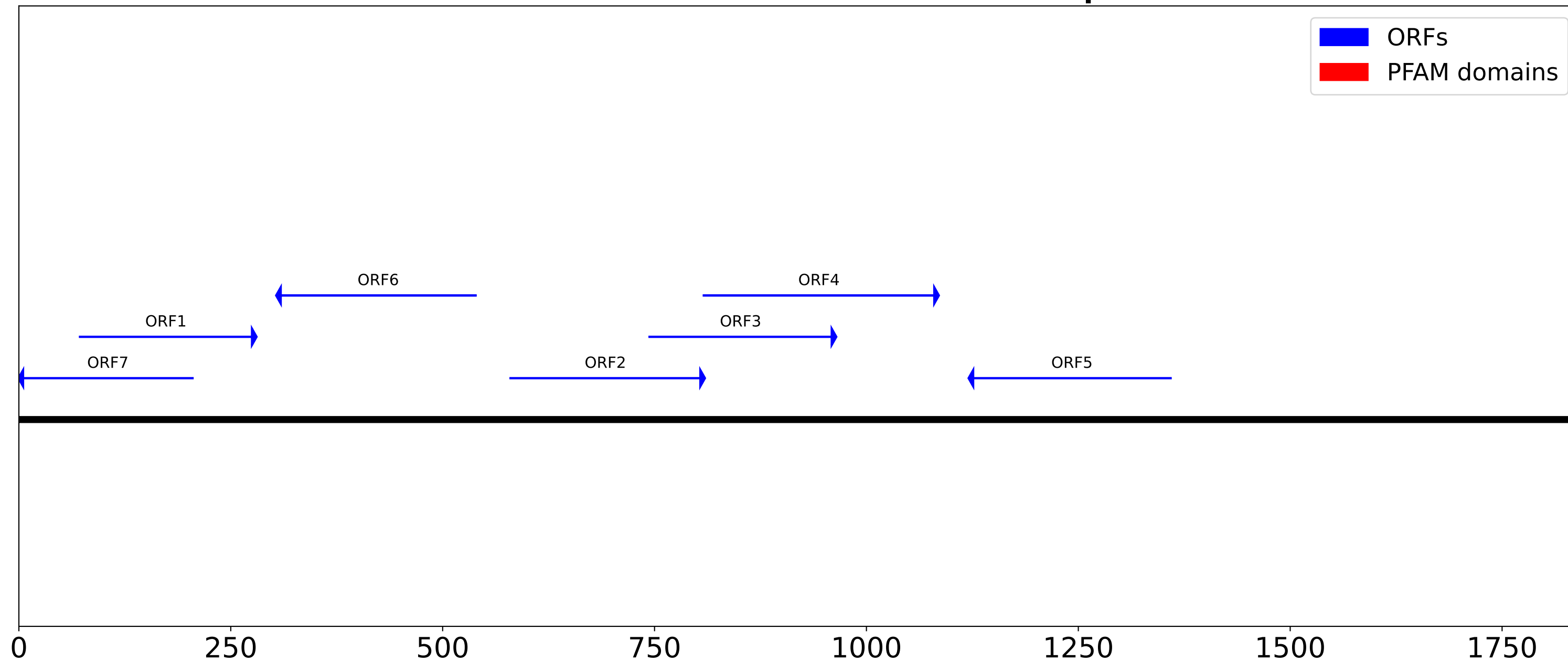
TE: rnd_5_family_7682
size: 589bp; fragments: 546; full length: 15 (≥ 530.1 bp)



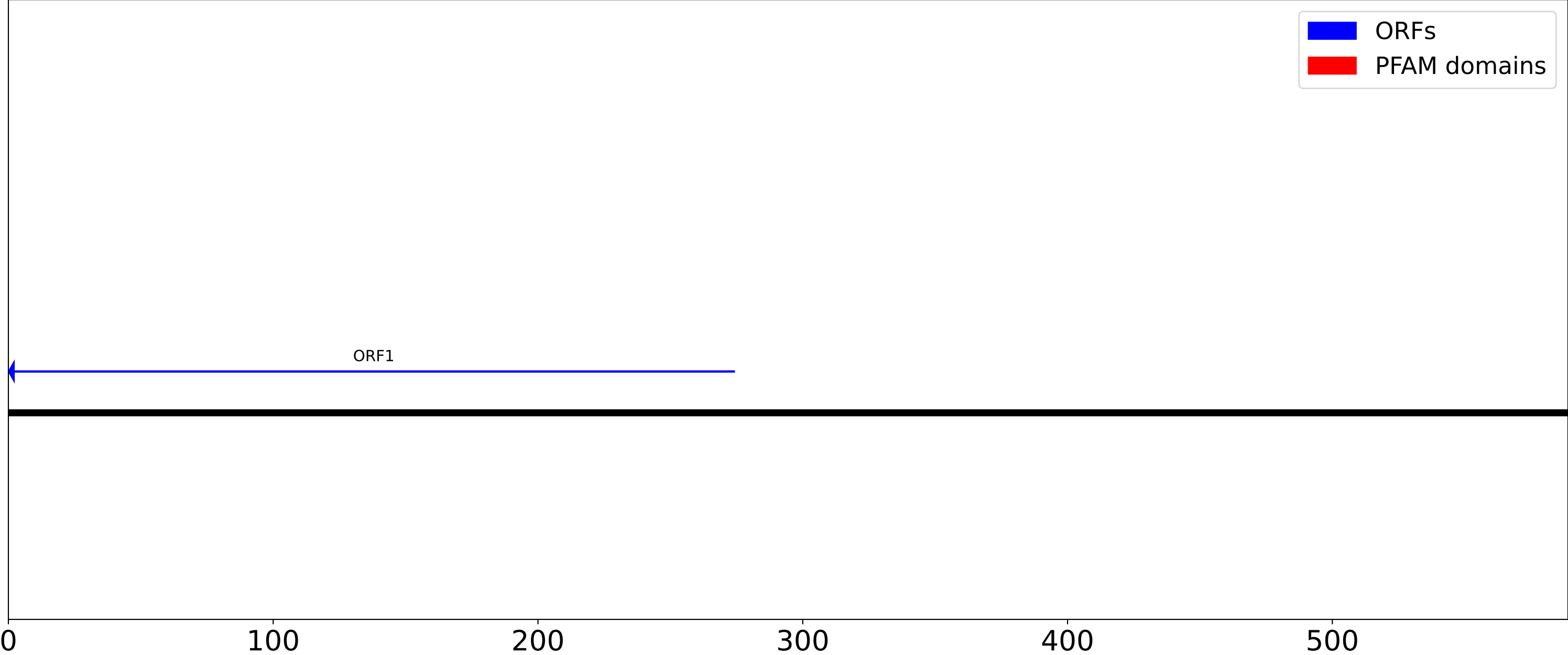
Before TEtrimmer 589 bp



After TEtrimmer ORF and PFAM domain plot



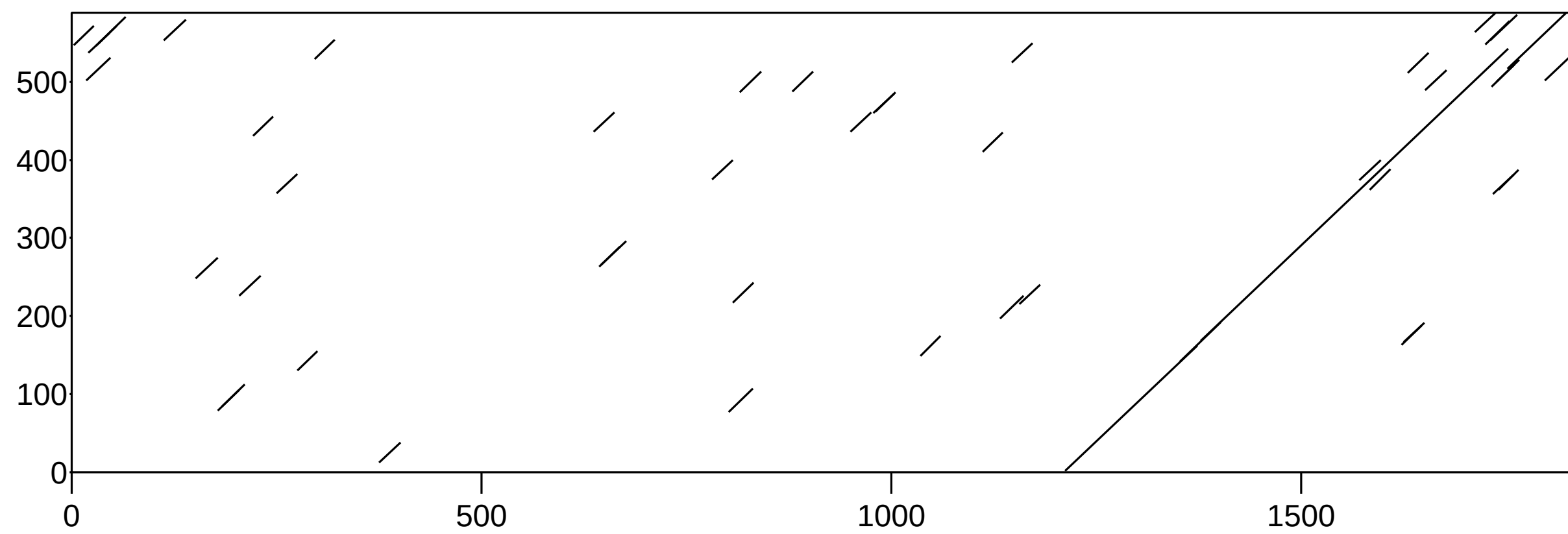
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)