

Start crop Point



End crop Point



MSA length = 1746



Start crop Point

End crop Point

0

10

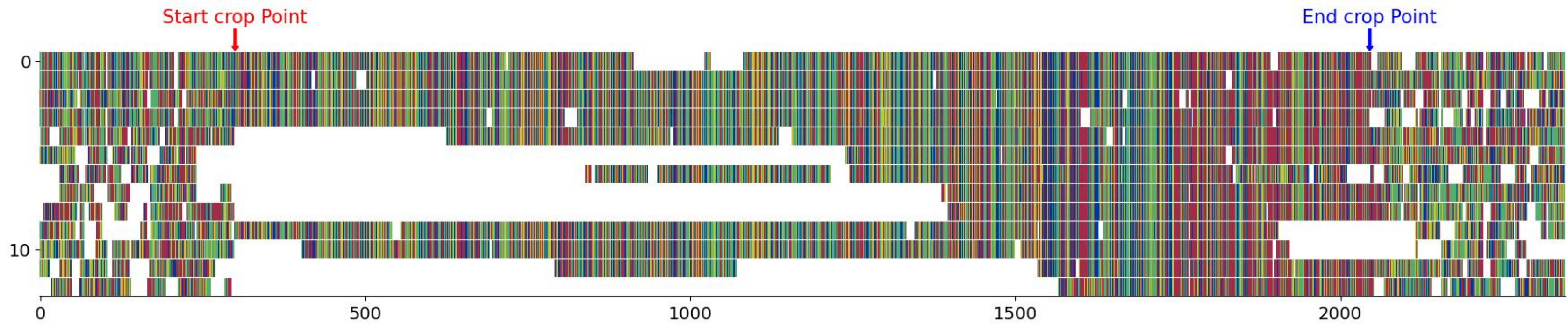
0

500

1000

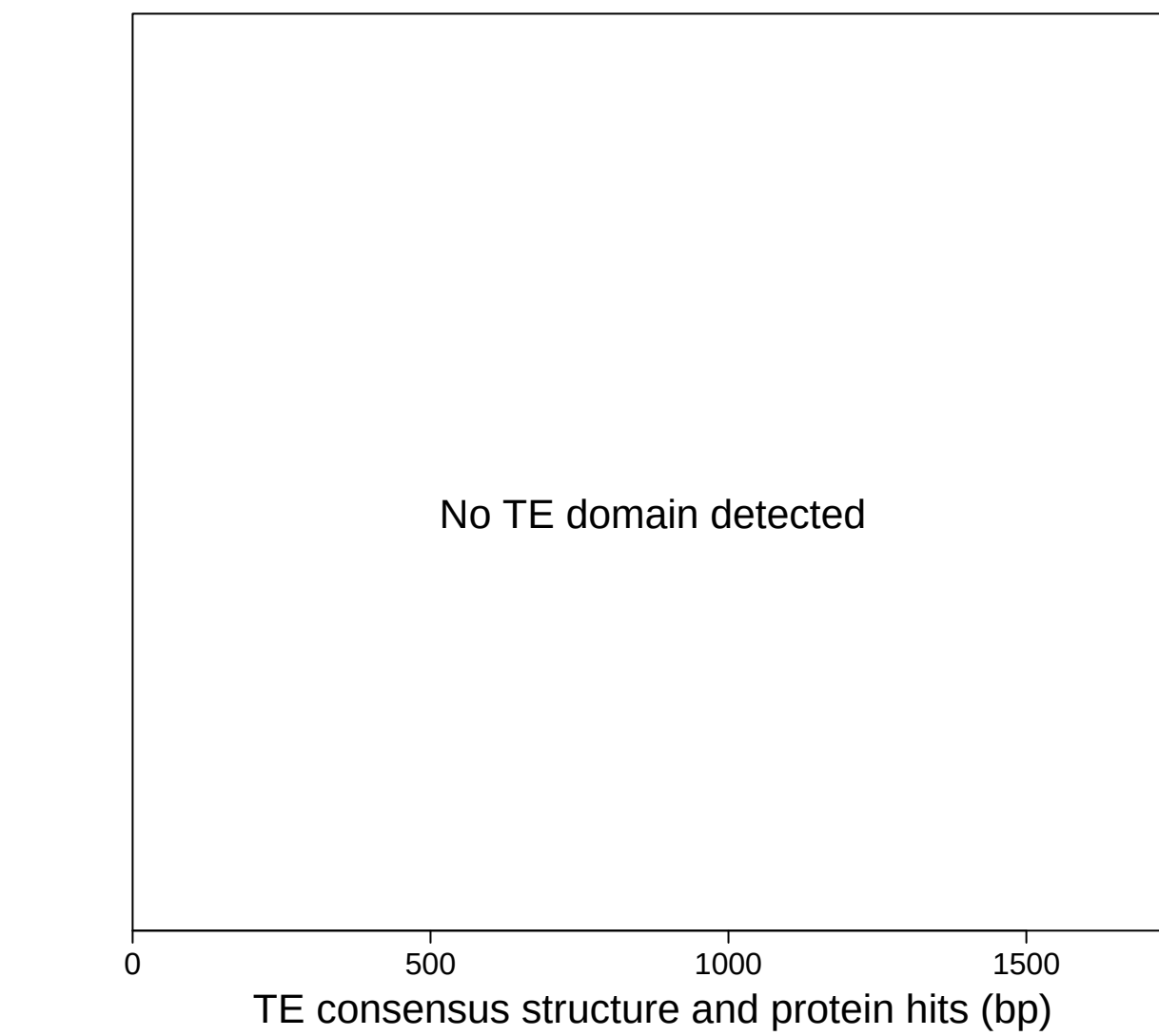
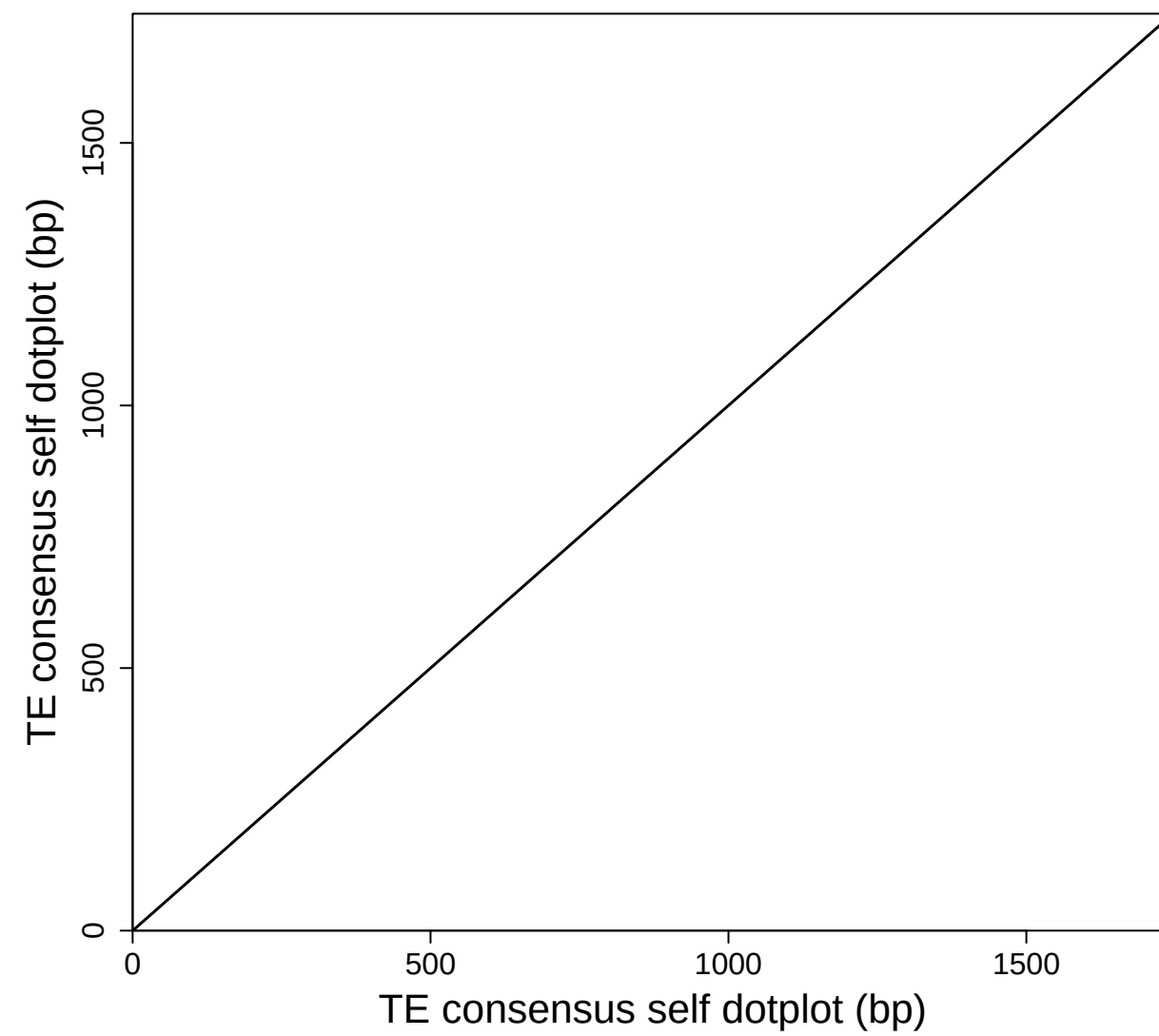
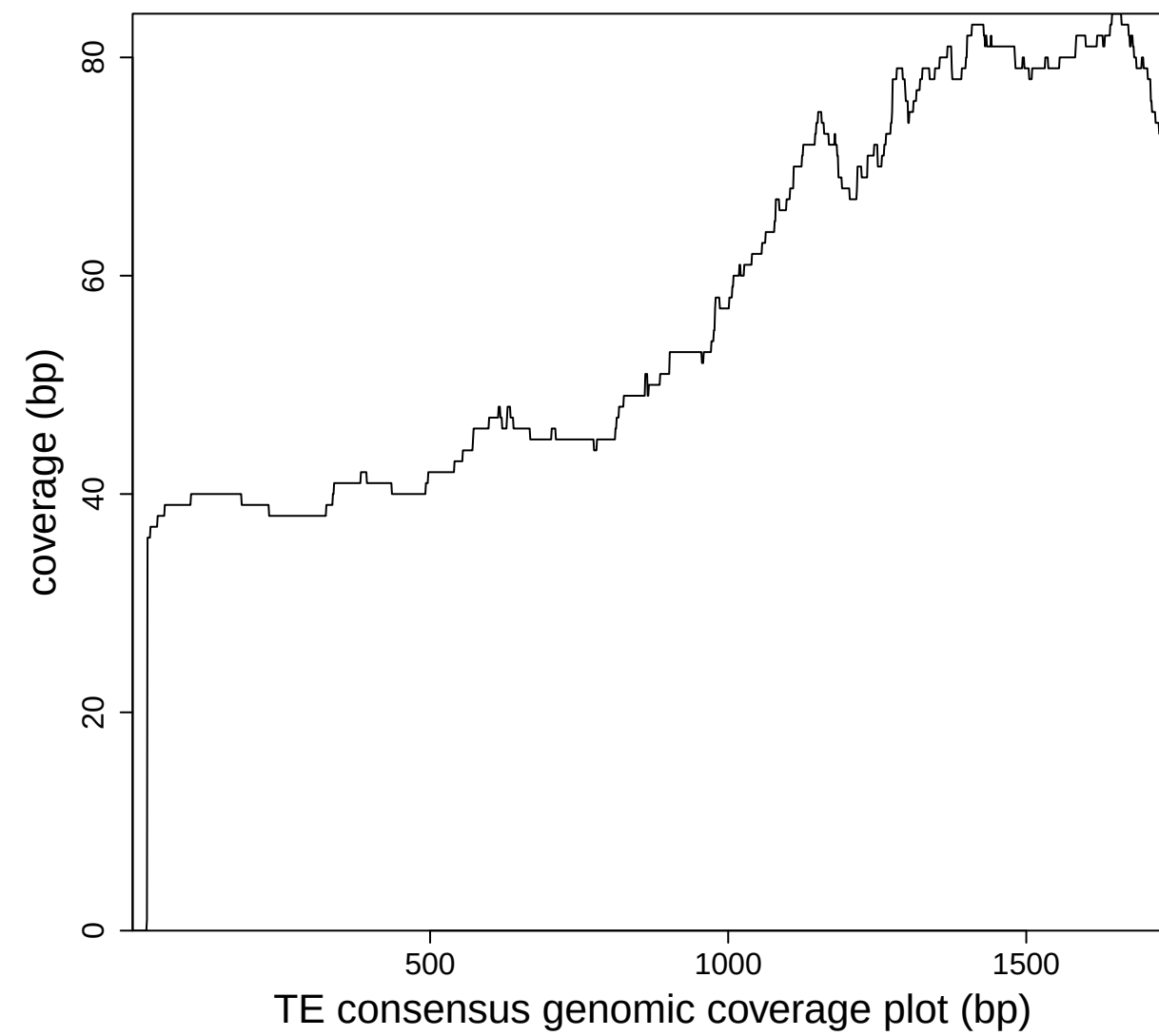
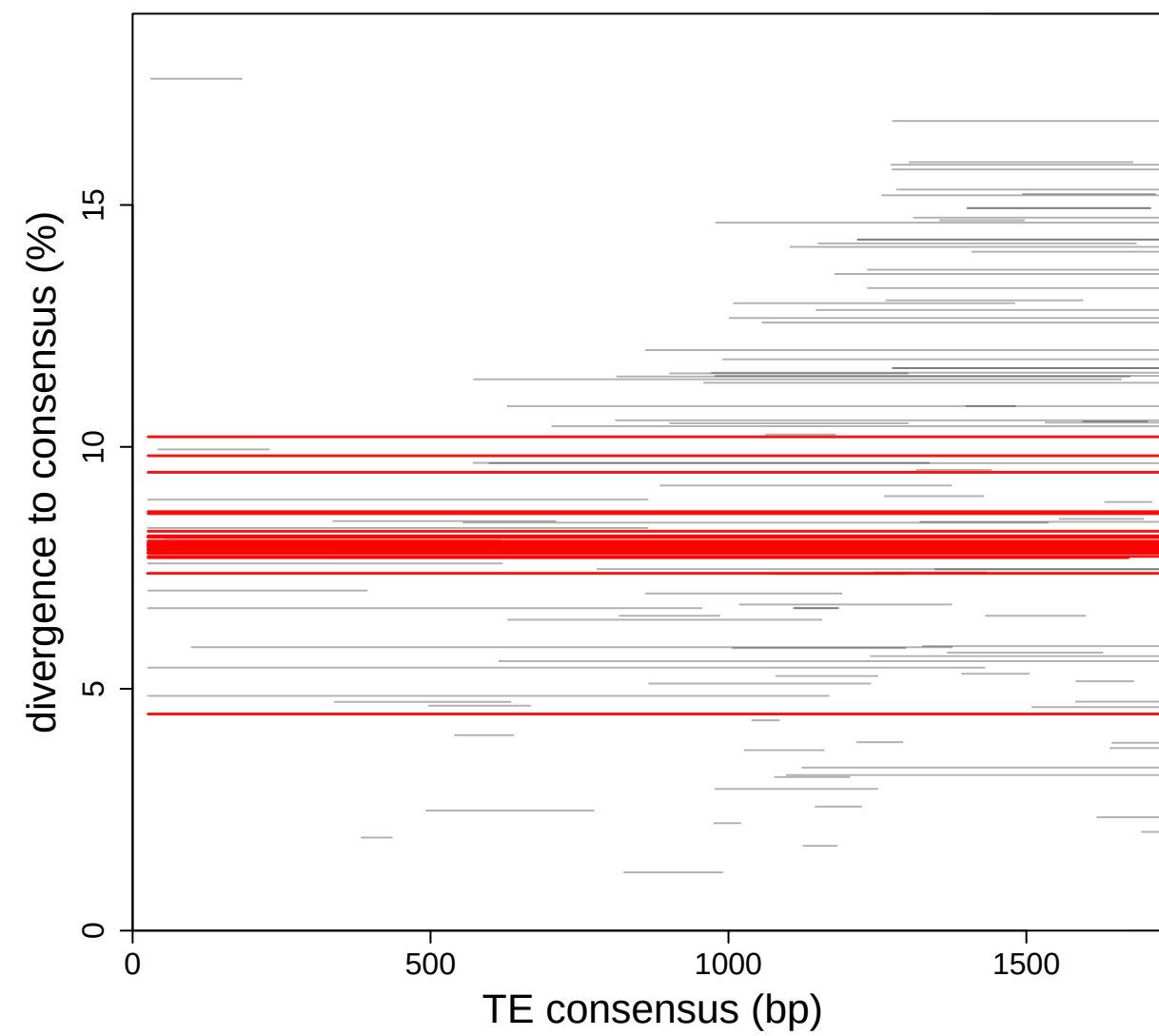
1500

2000



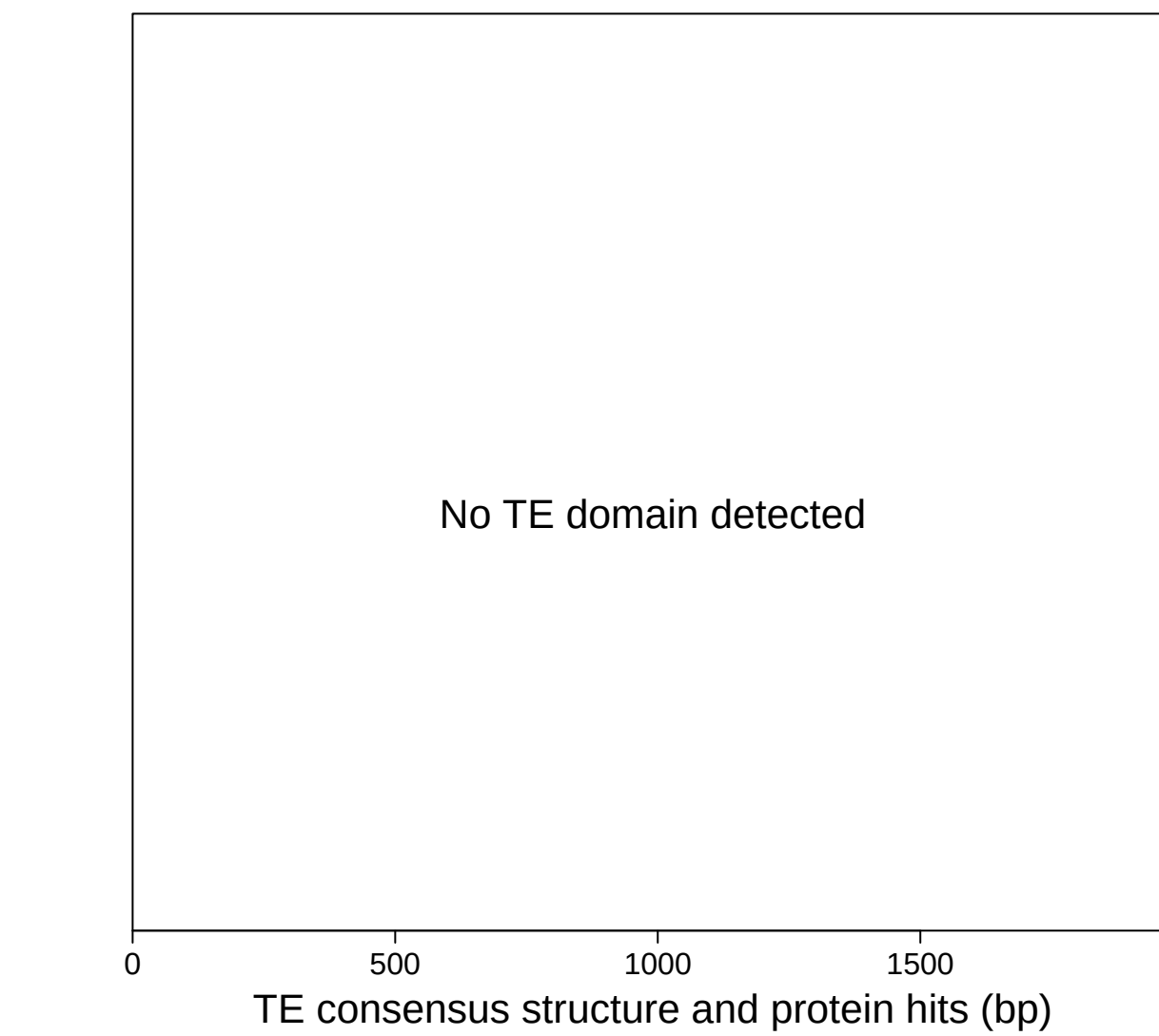
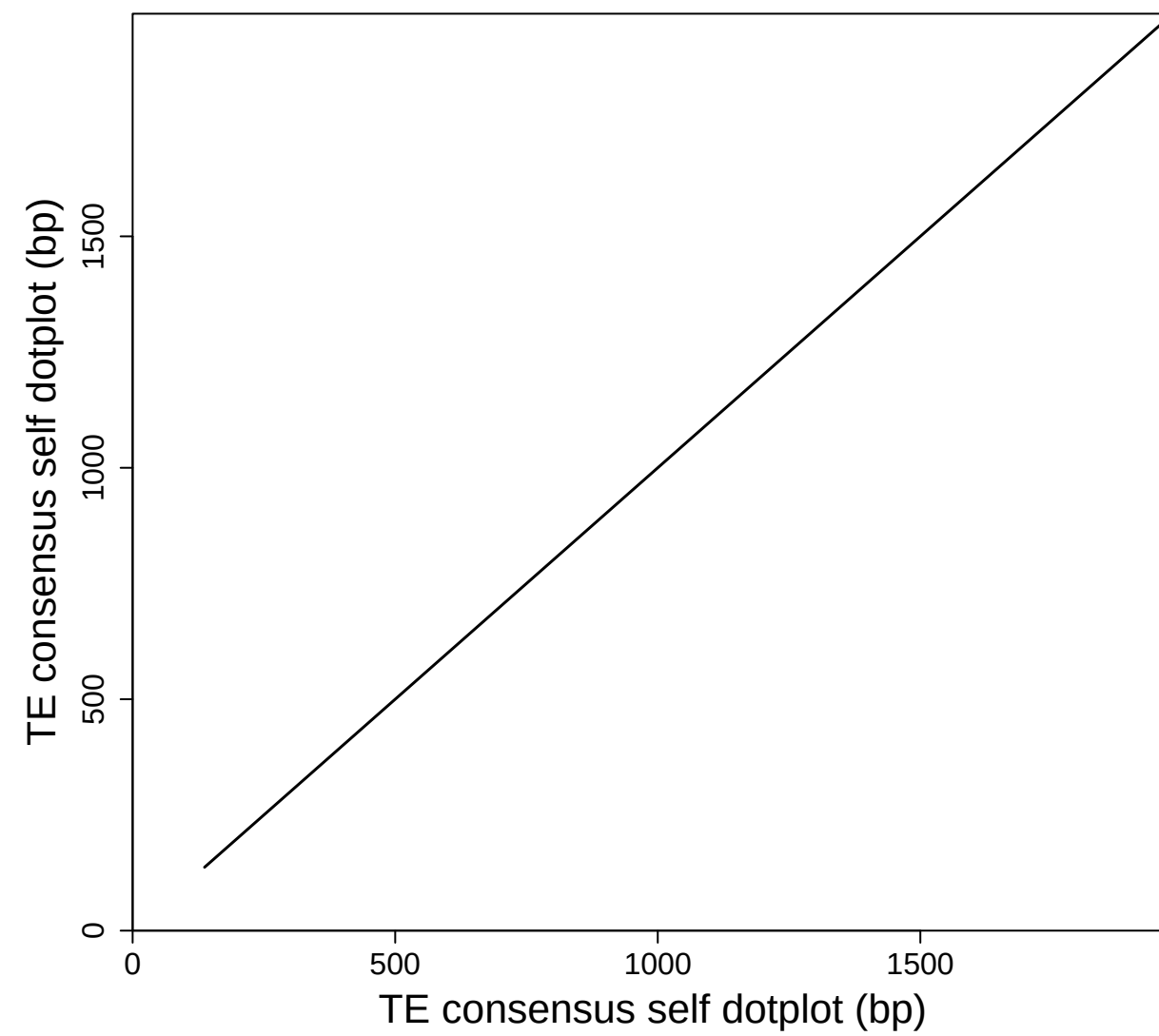
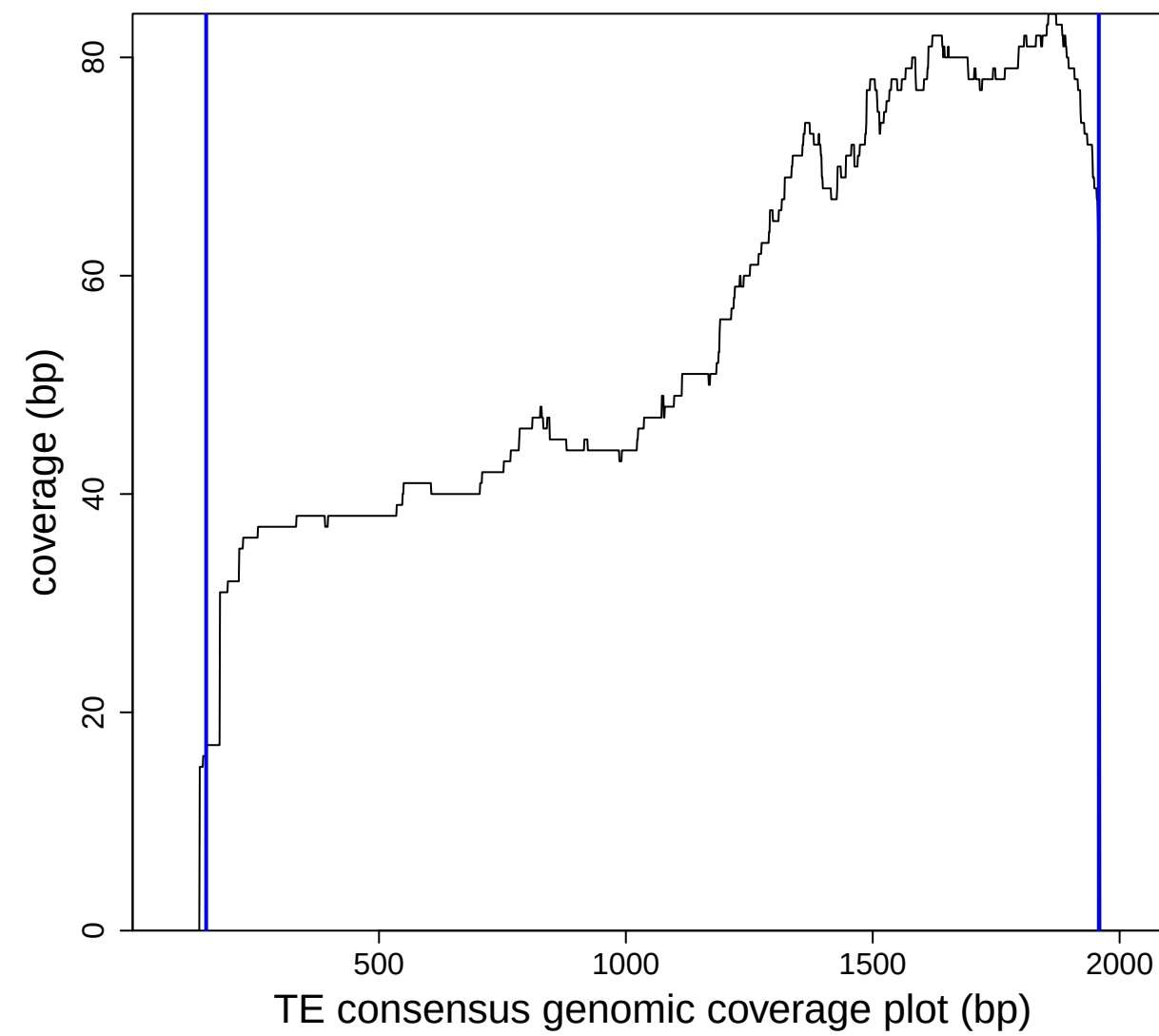
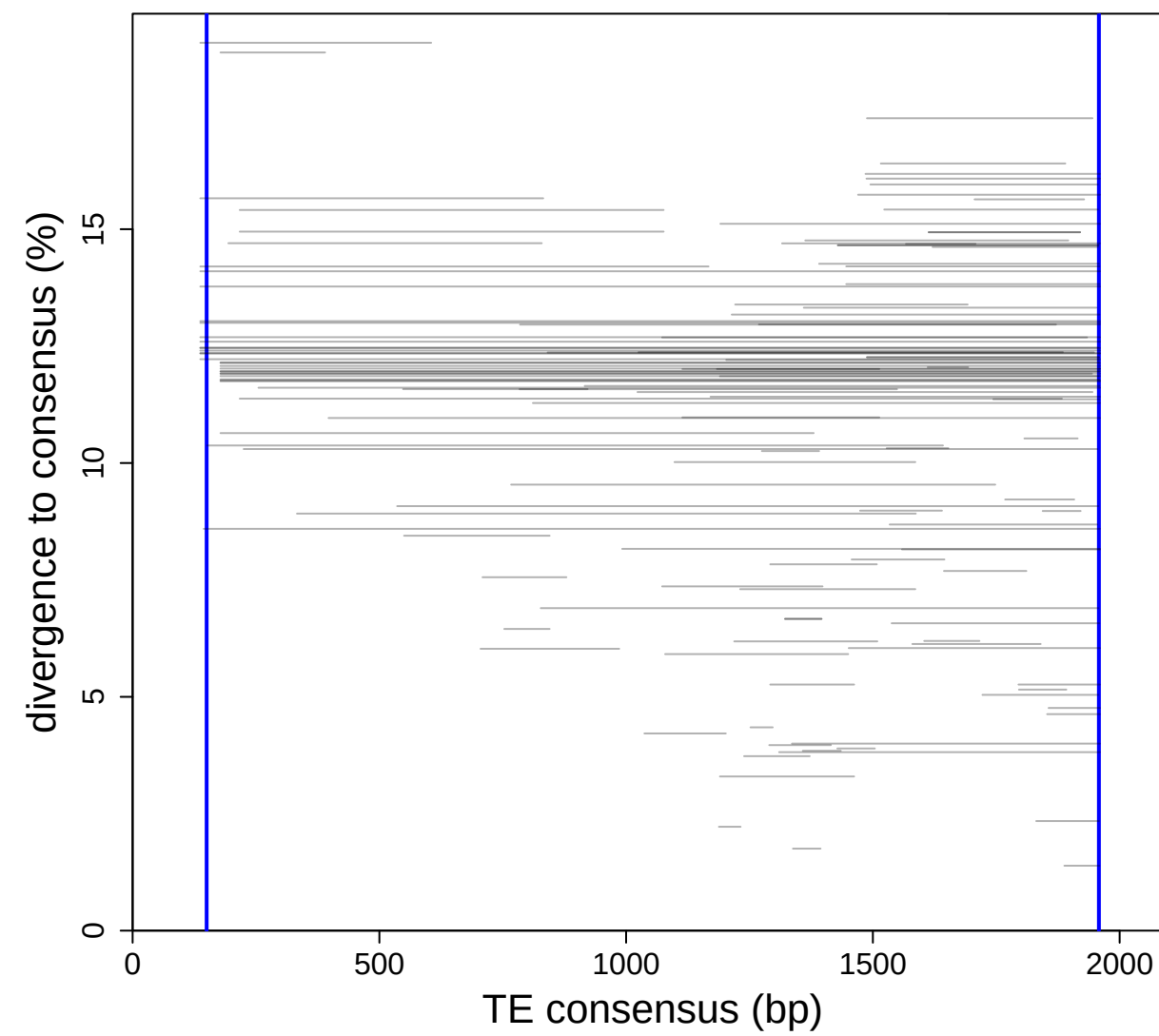
2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.
size: 1746bp; fragments: 138; full length: 29 (≥ 1571.4 bp)

After TEtrimmer 1746 bp

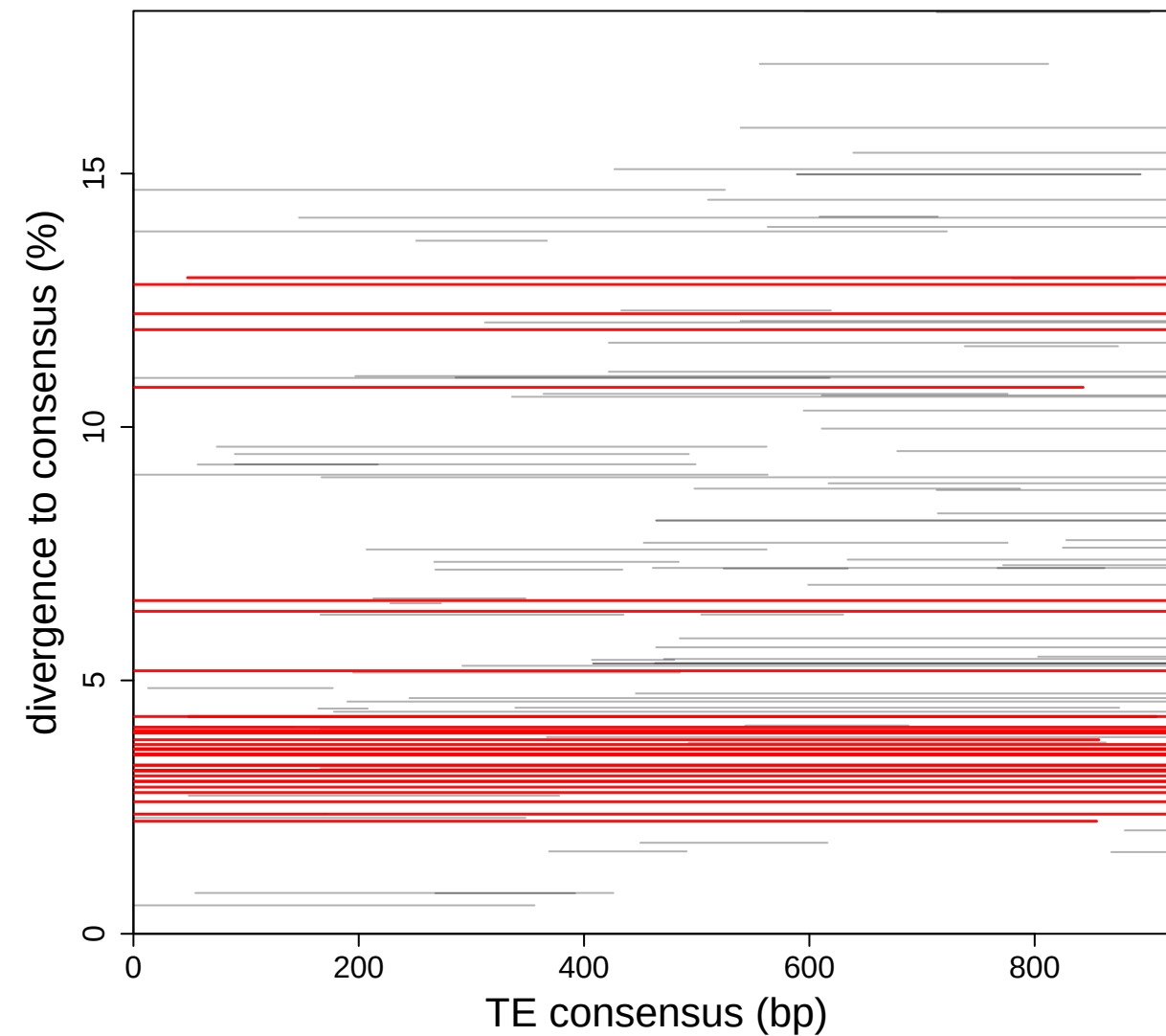


uf.bed g 2.bed fm 1.bed 0 0 n.bed g 1.bed fm 2.bed 0 0 b
size: 2108bp; fragments: 132; full length: 0 (>=1897.2bp)

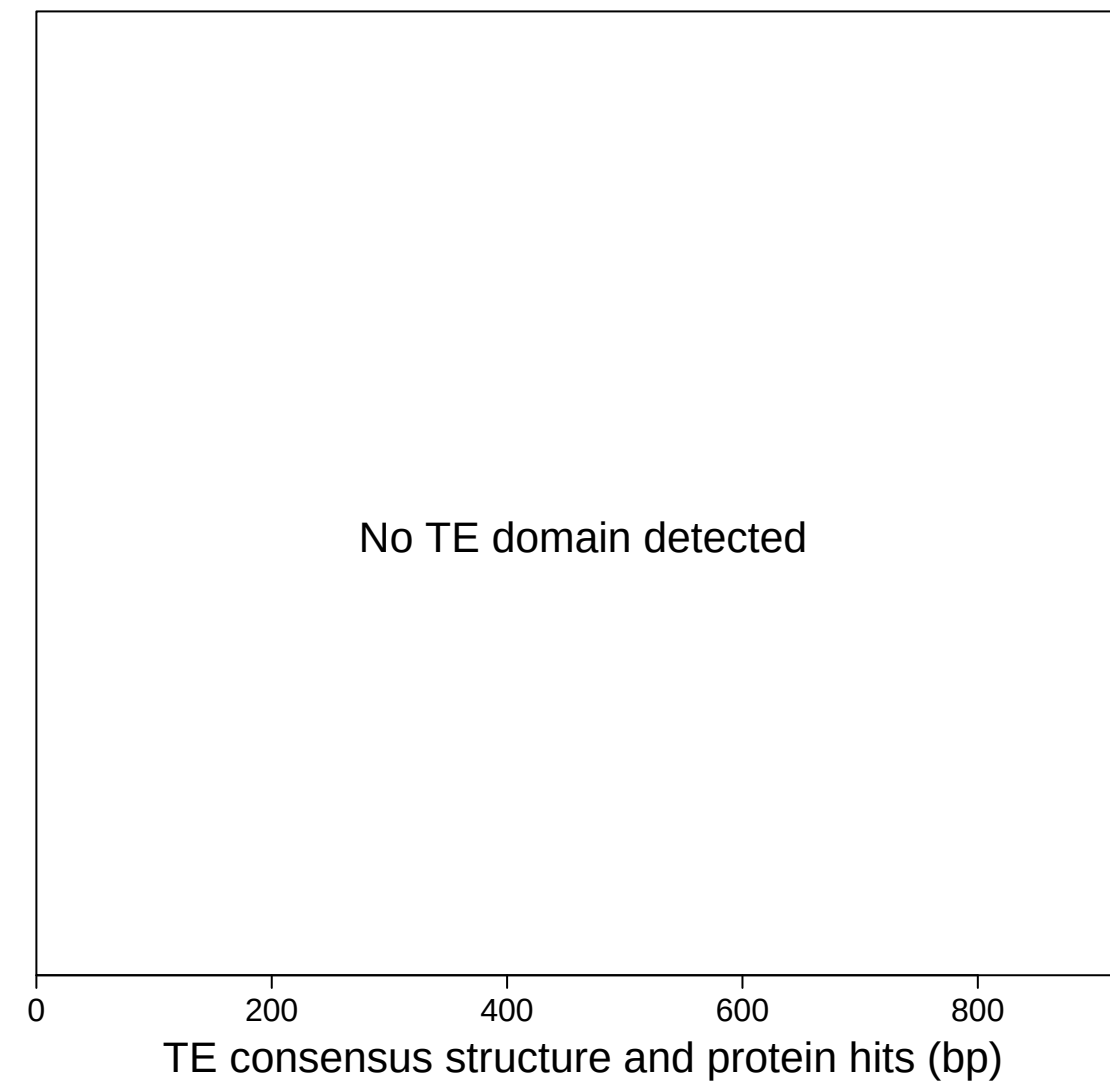
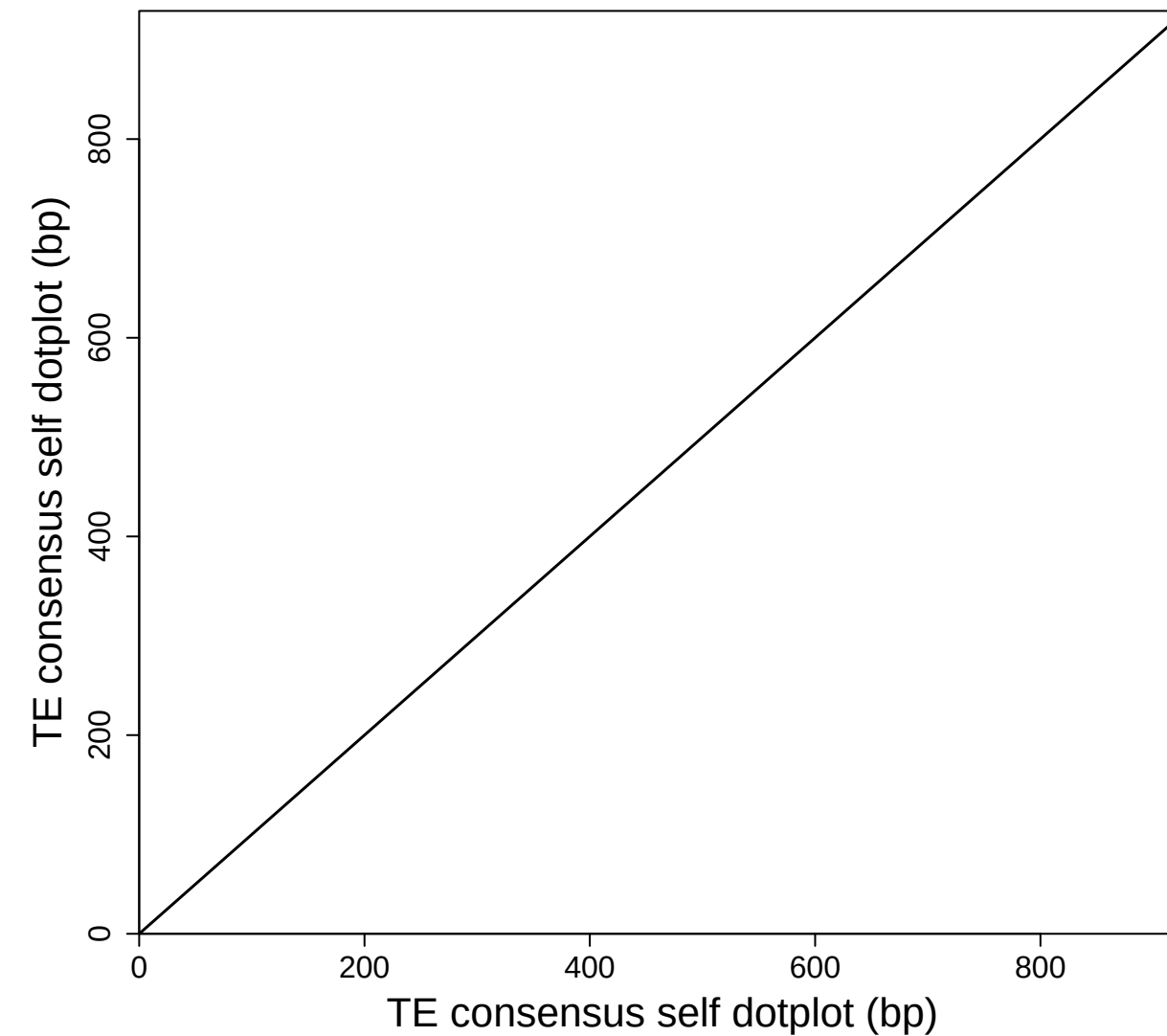
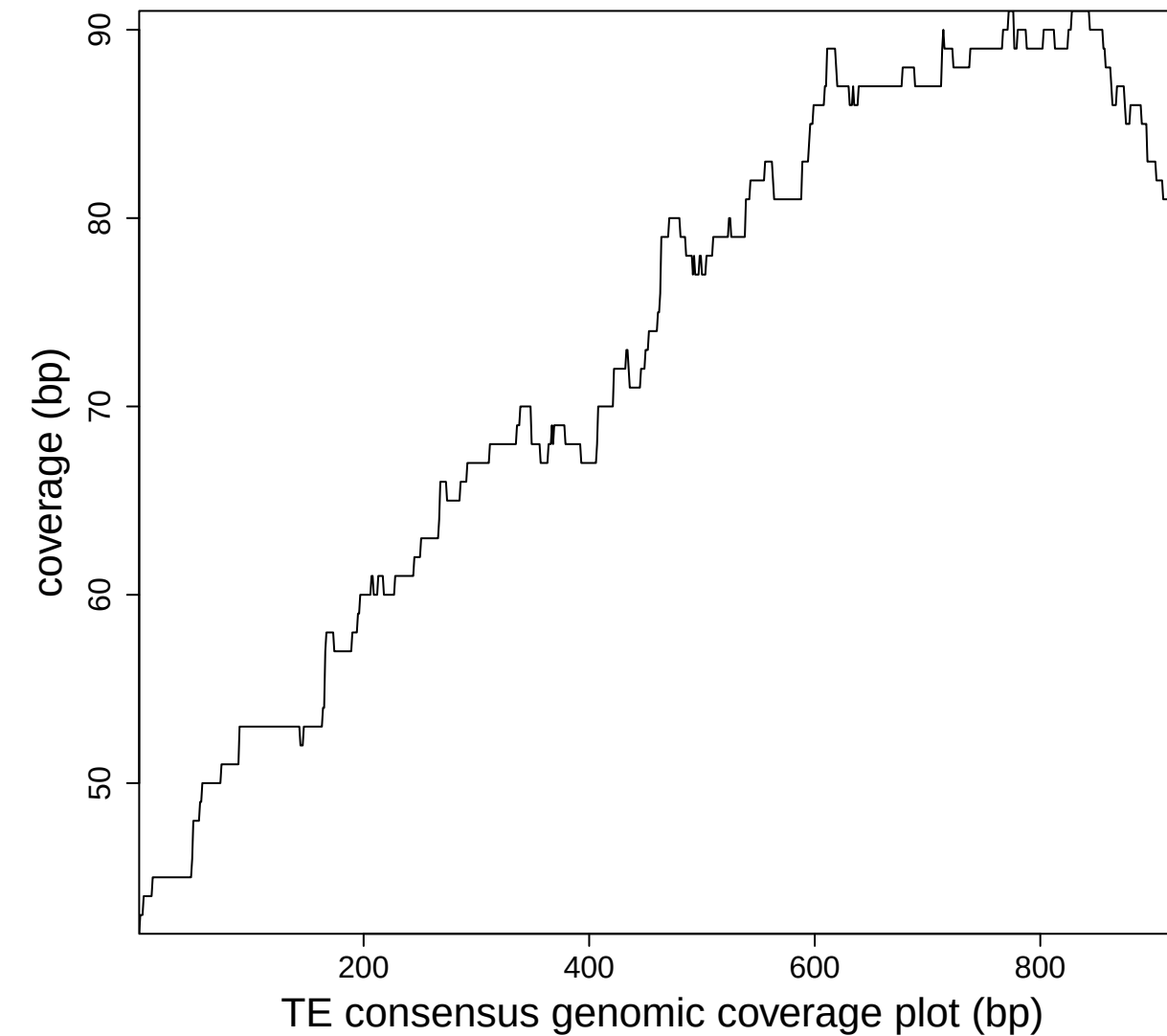
After TEtrimmer Extended plot Blue lines are boundaries



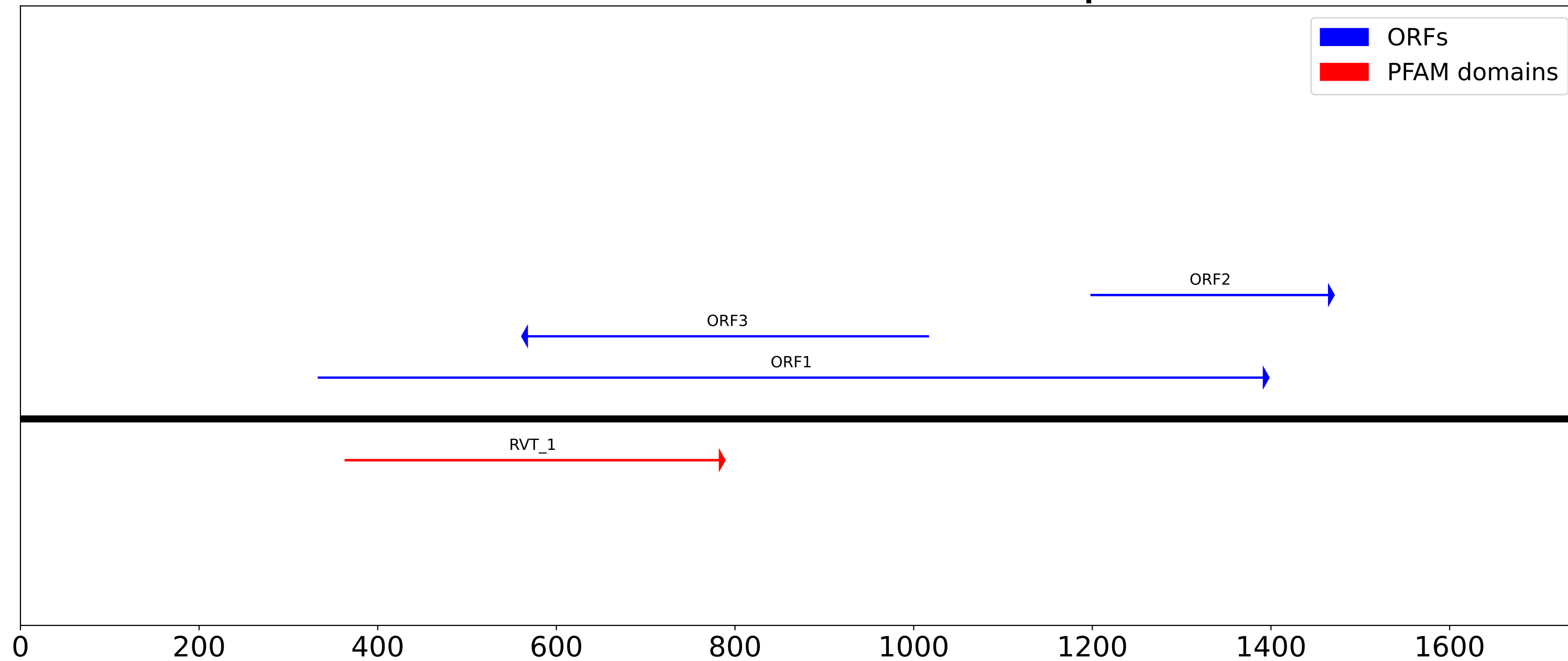
TE: rnd_5_family_19222
size: 929bp; fragments: 130; full length: 38 (≥ 836.1 bp)



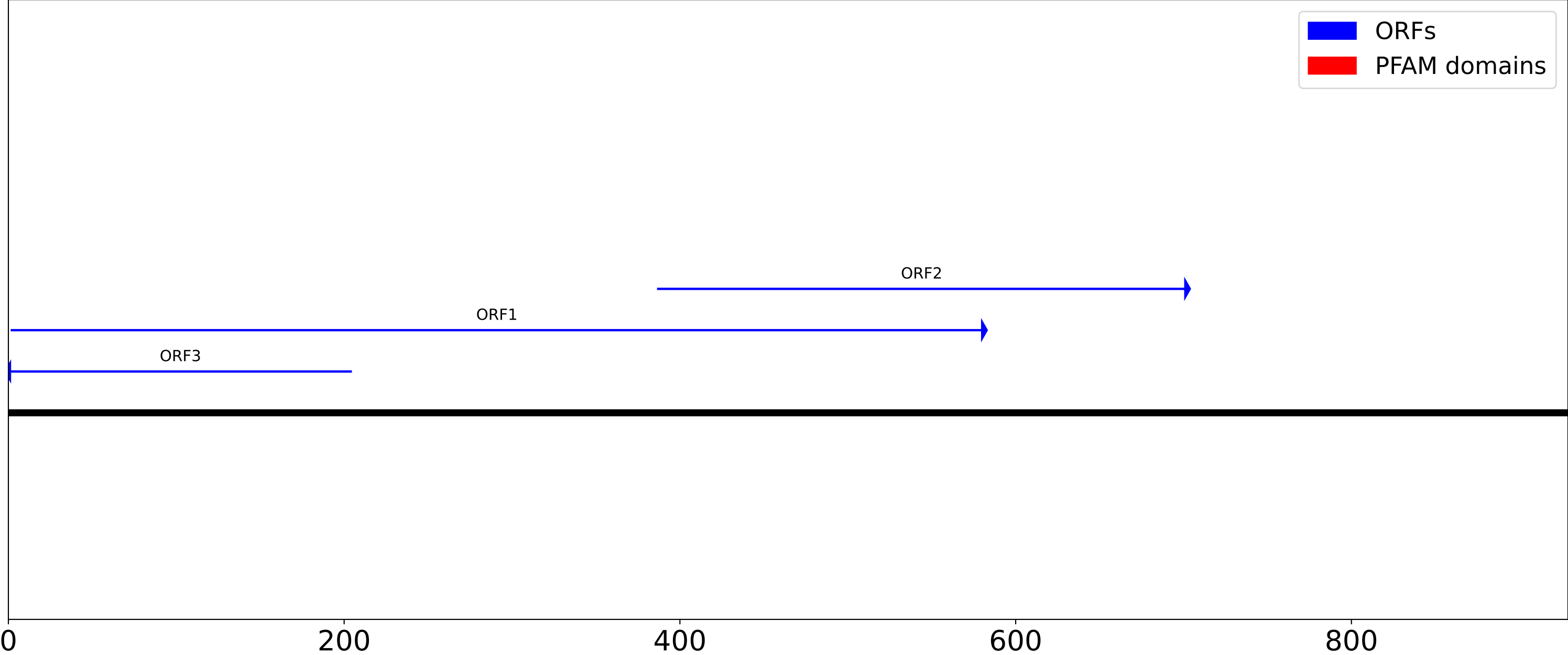
Before TEtrimmer 929 bp



After TEtrimmer ORF and PFAM domain plot



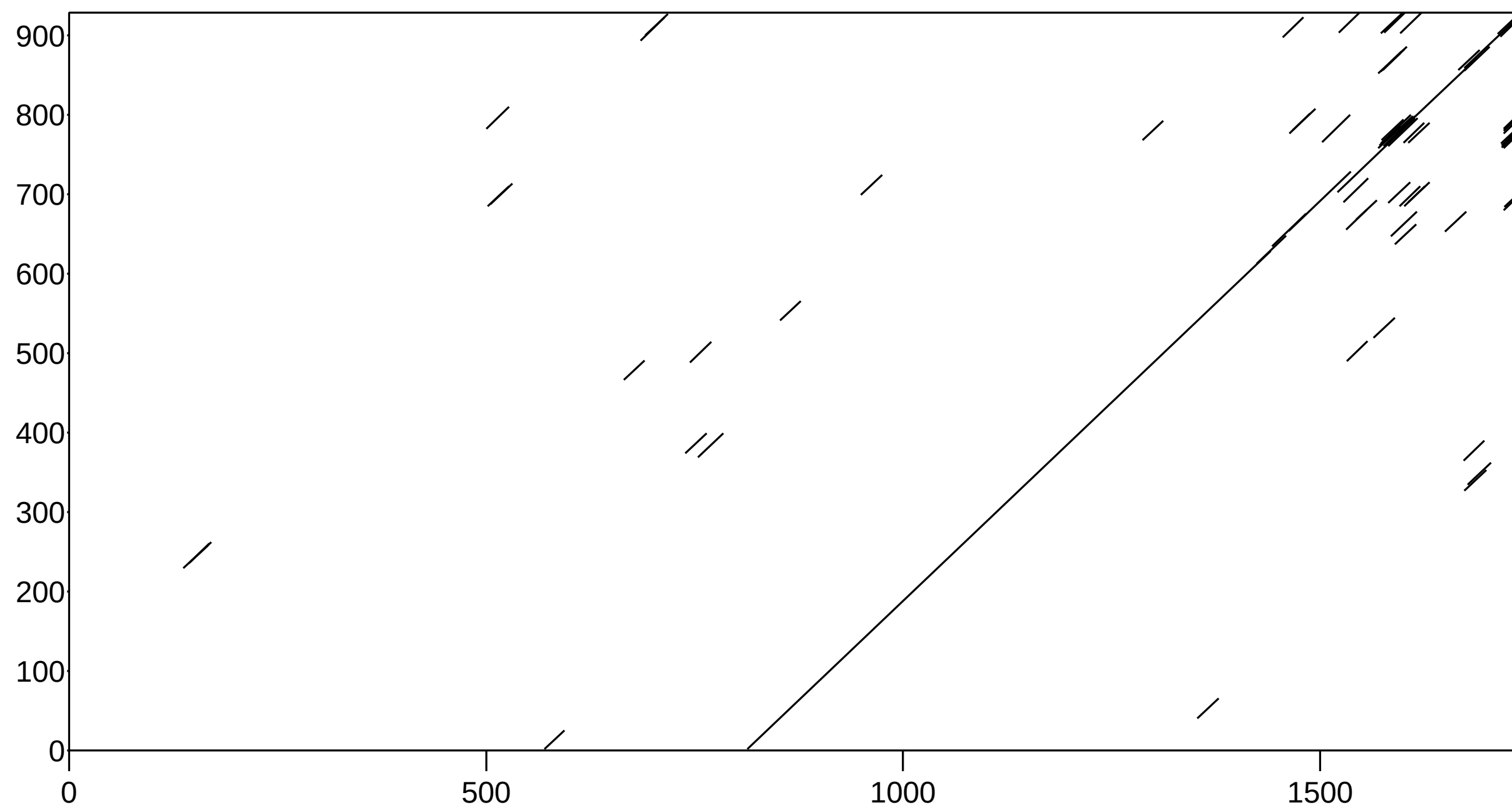
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)