

Start crop Point

End crop Point

0

10

0

250

500

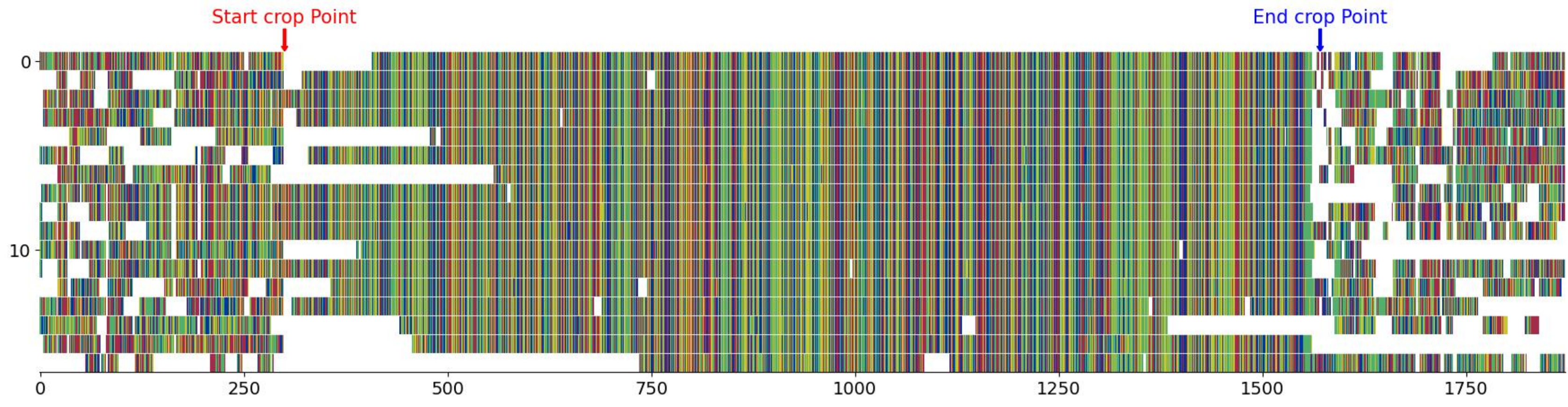
750

1000

1250

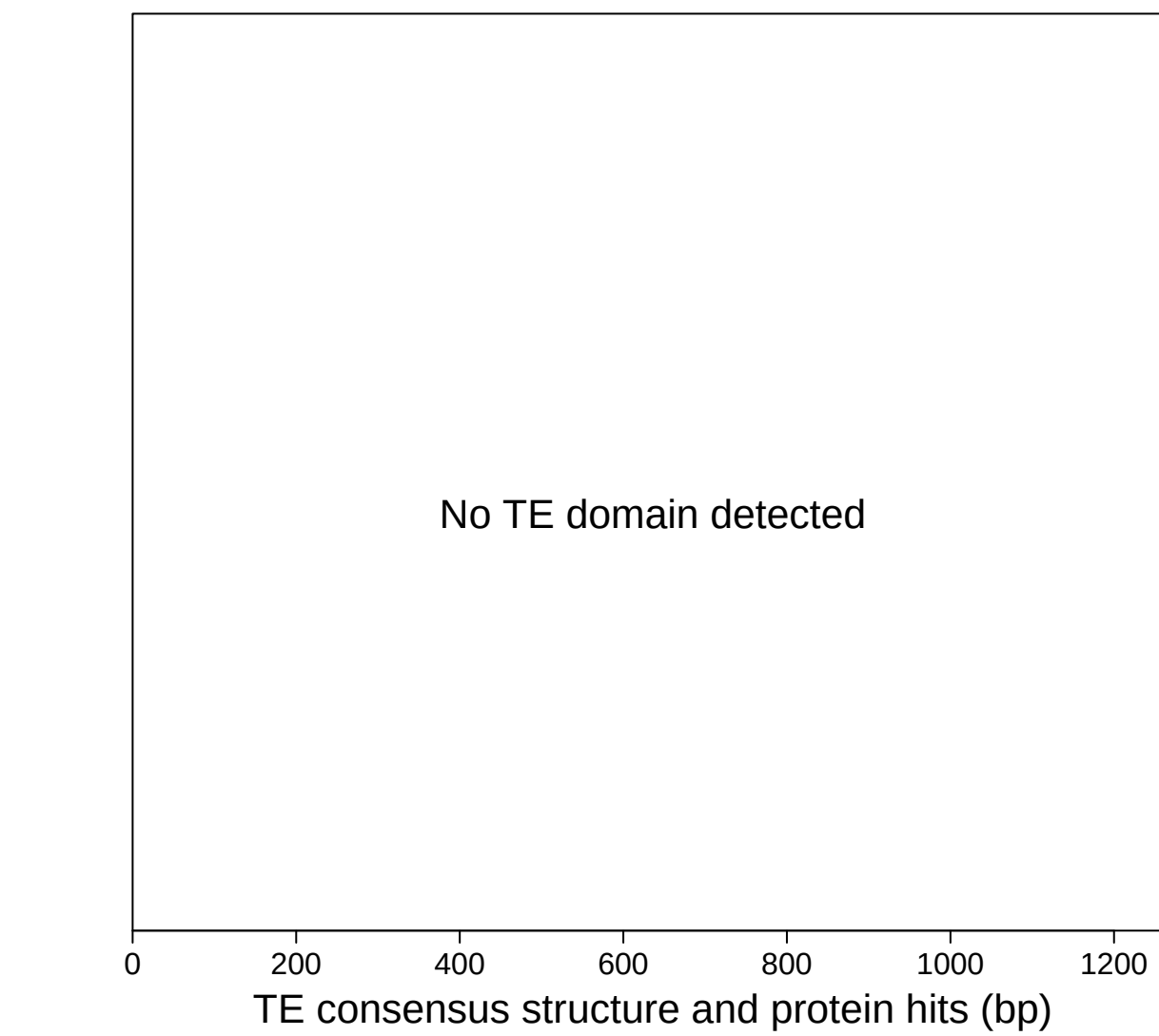
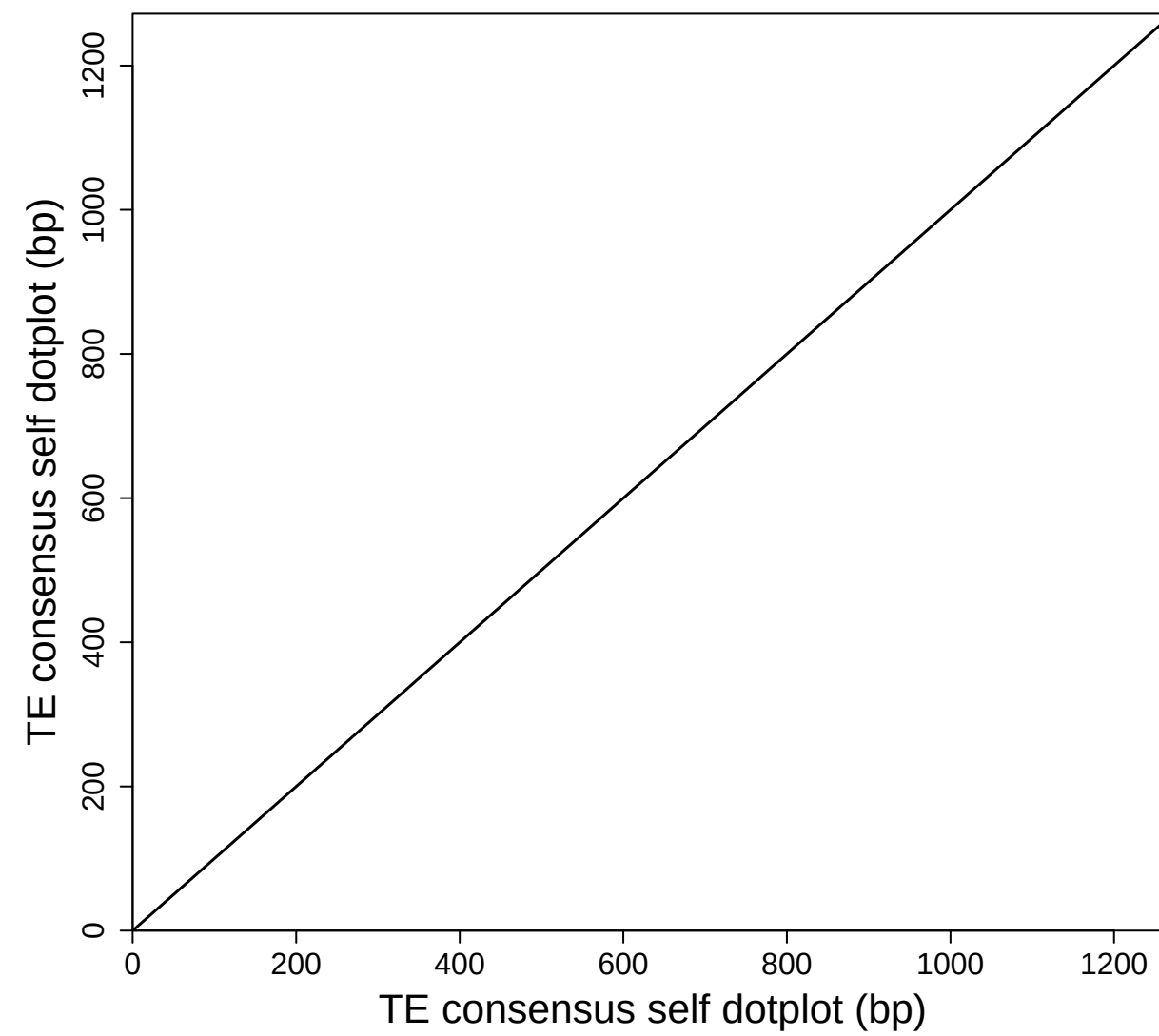
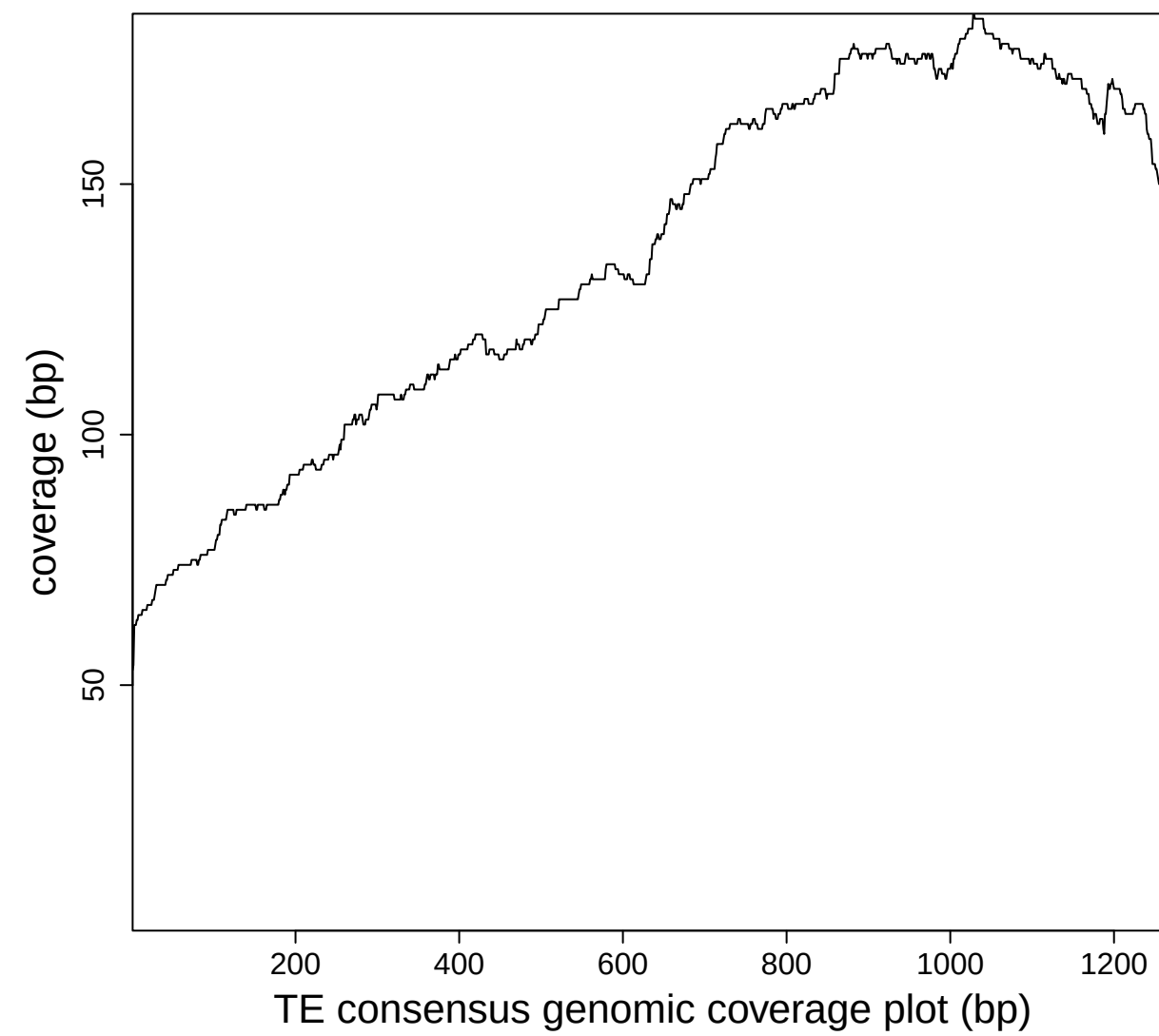
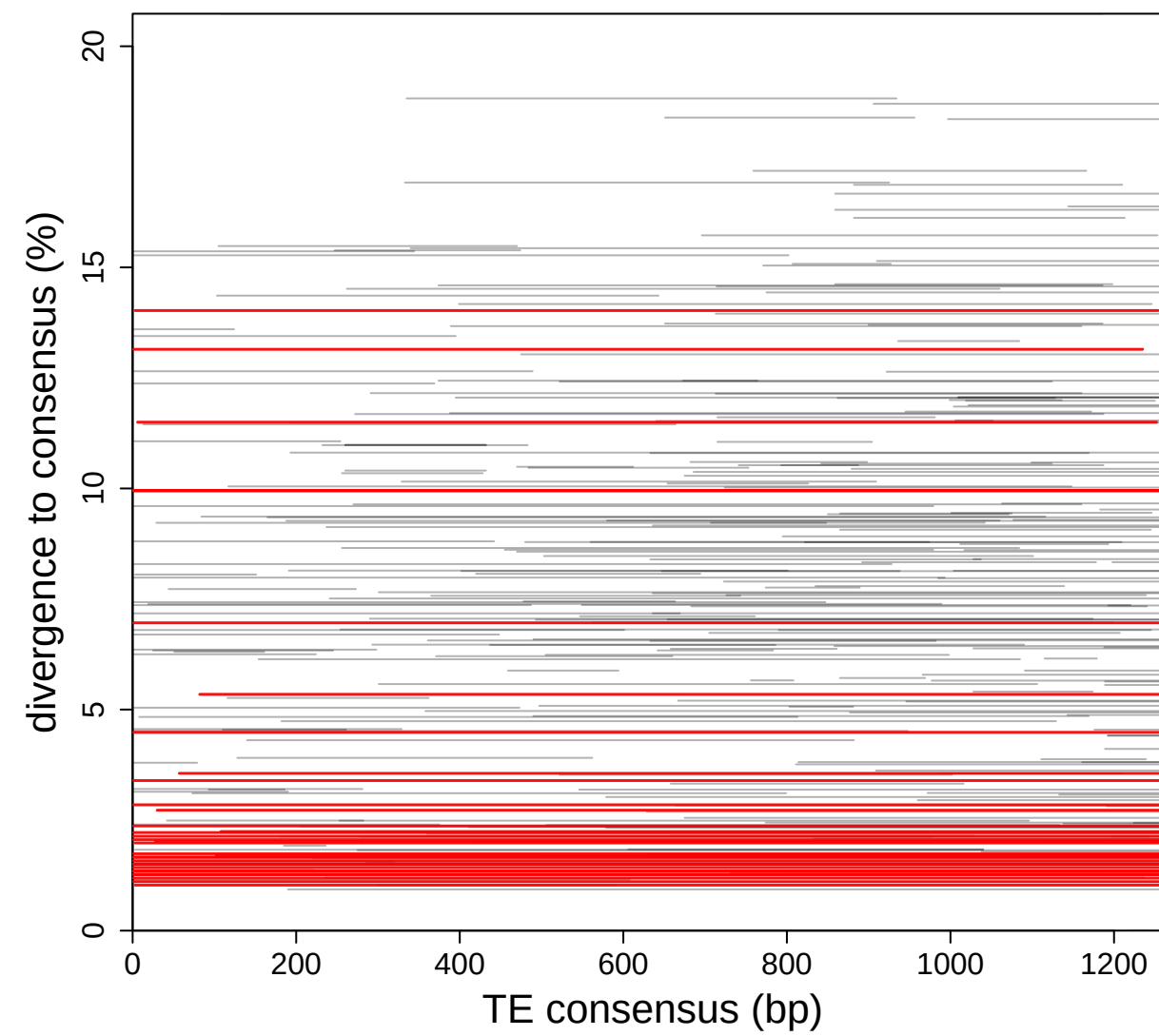
1500

1750



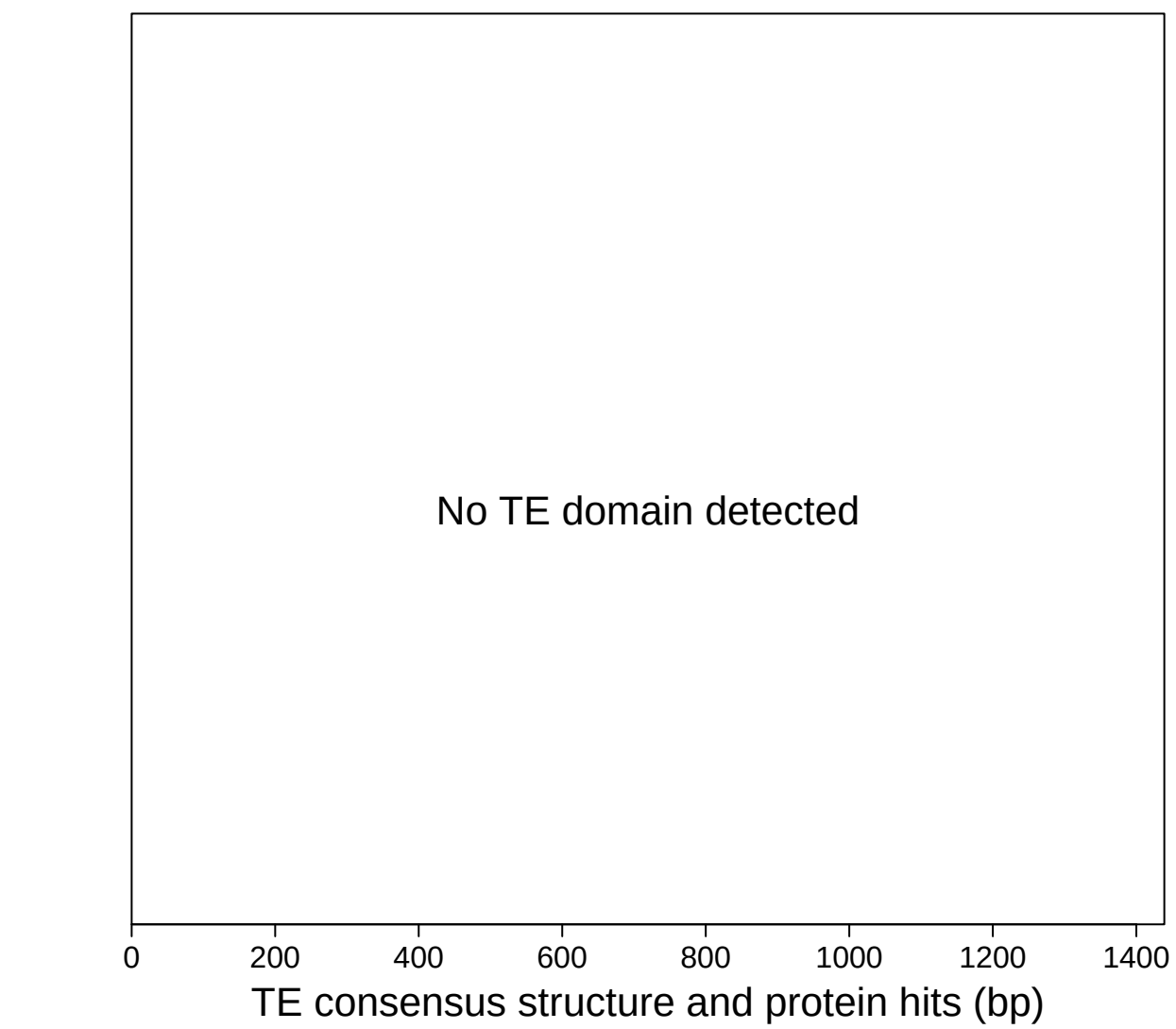
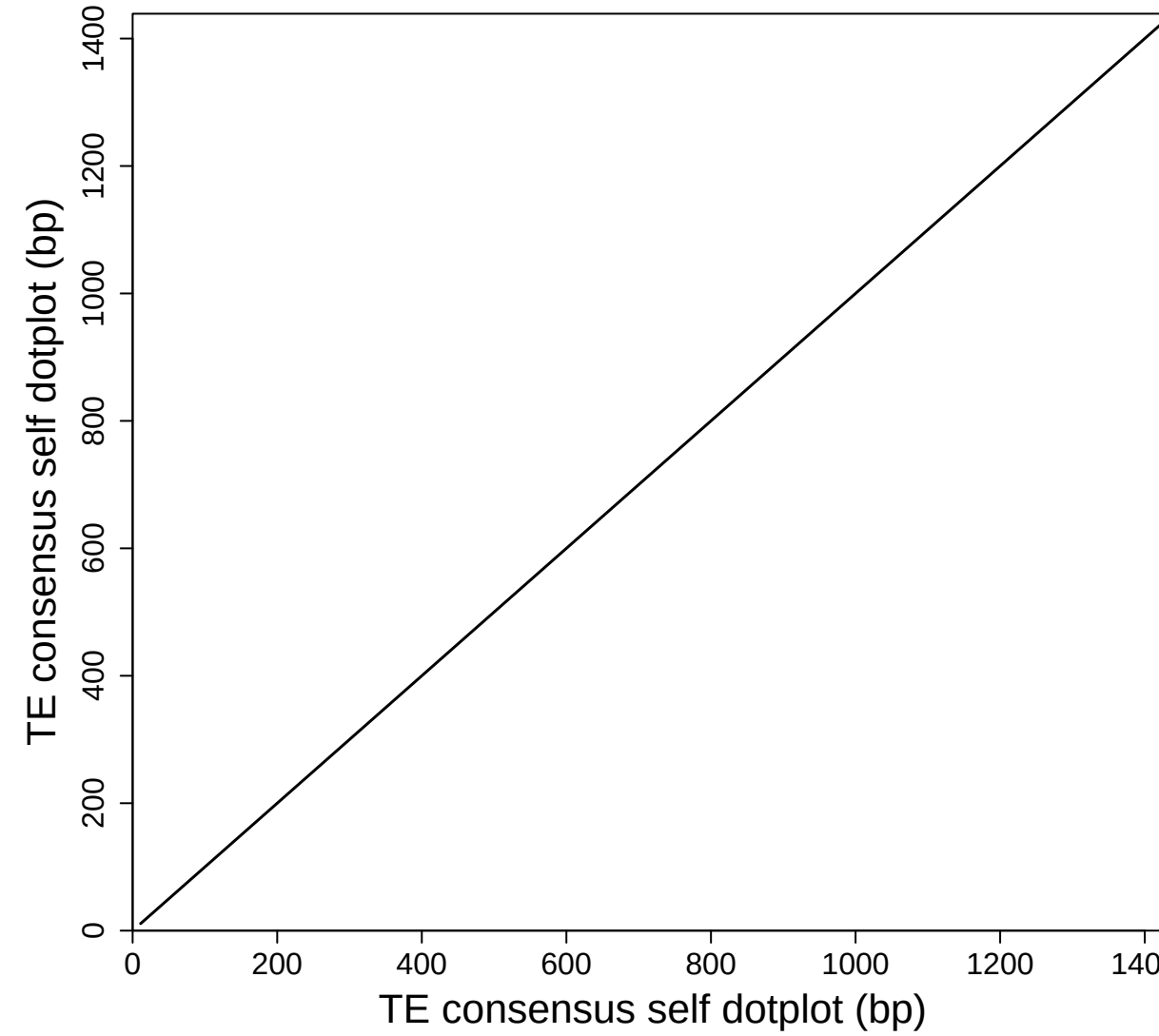
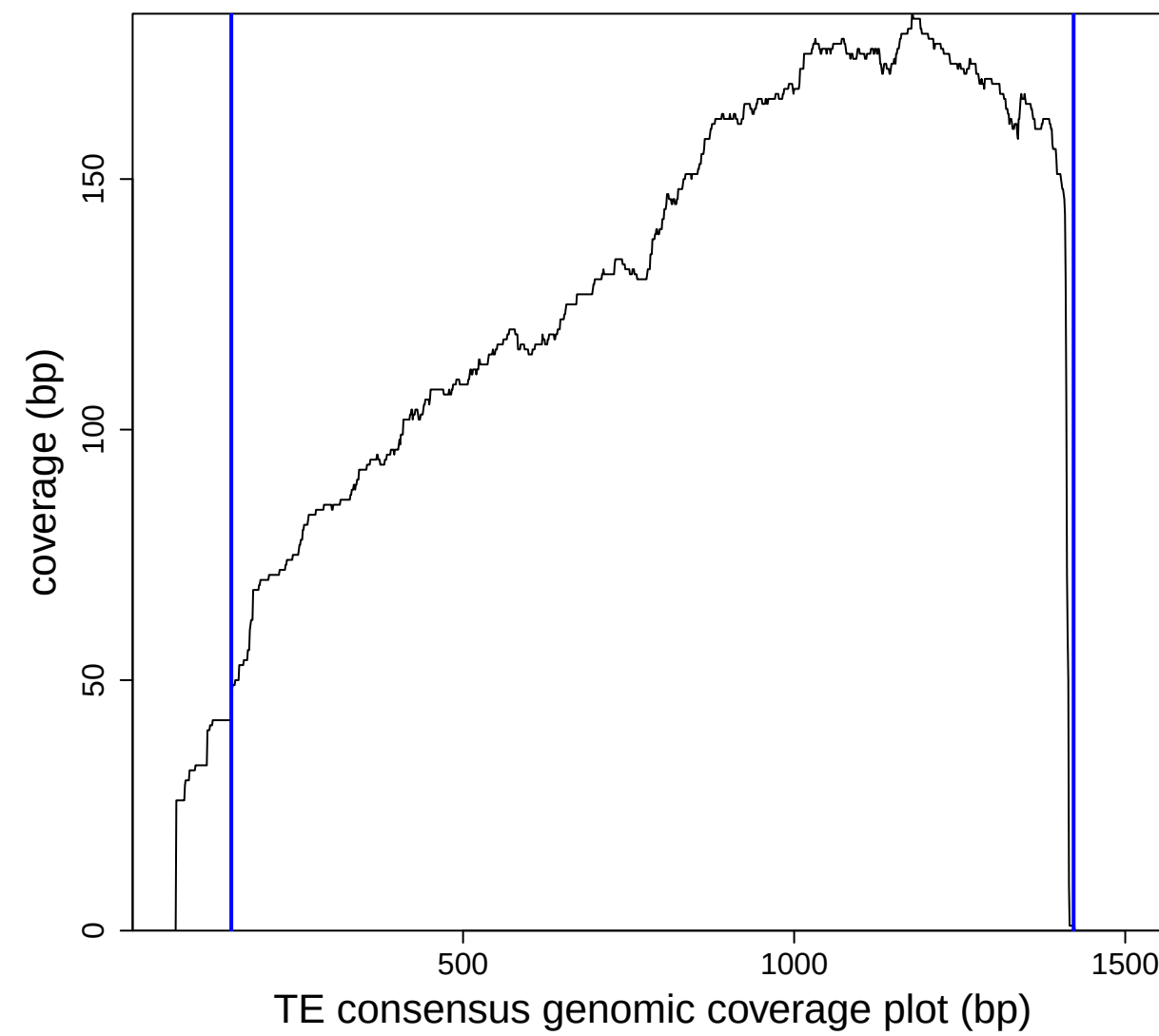
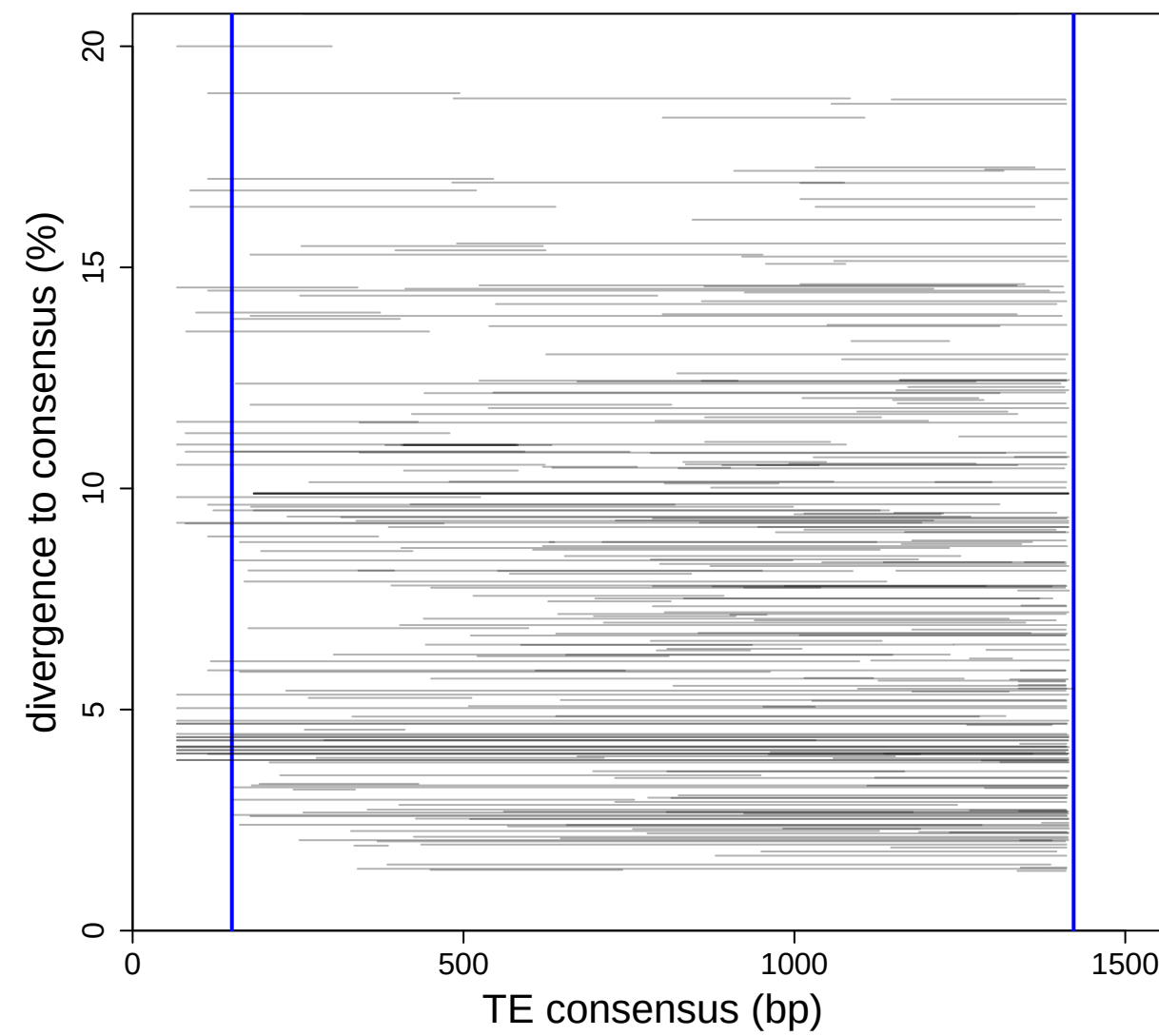
bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa c  
size: 1272bp; fragments: 306; full length: 39 (>=1144.8bp)

# After TEtrimmer 1272 bp

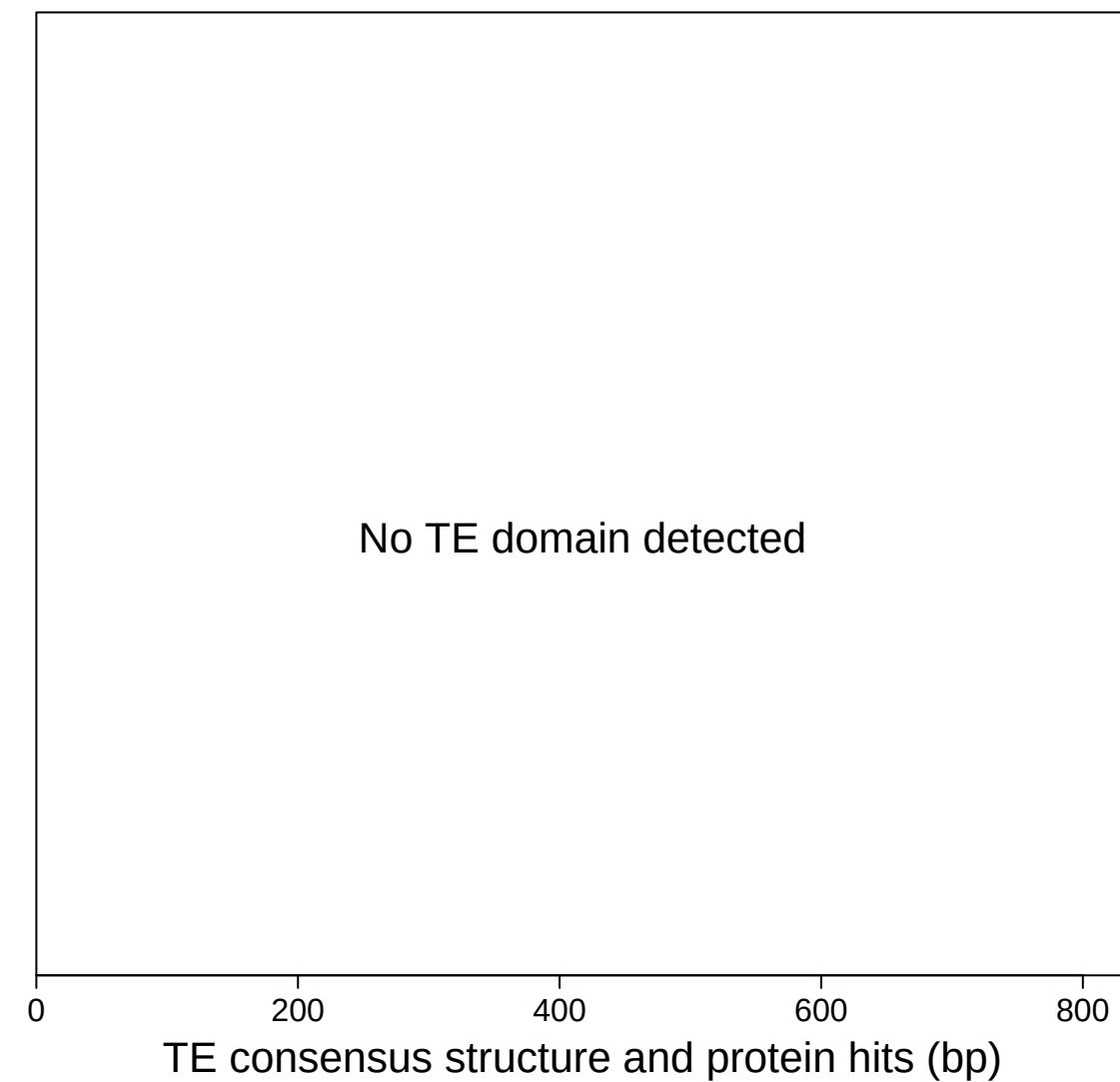
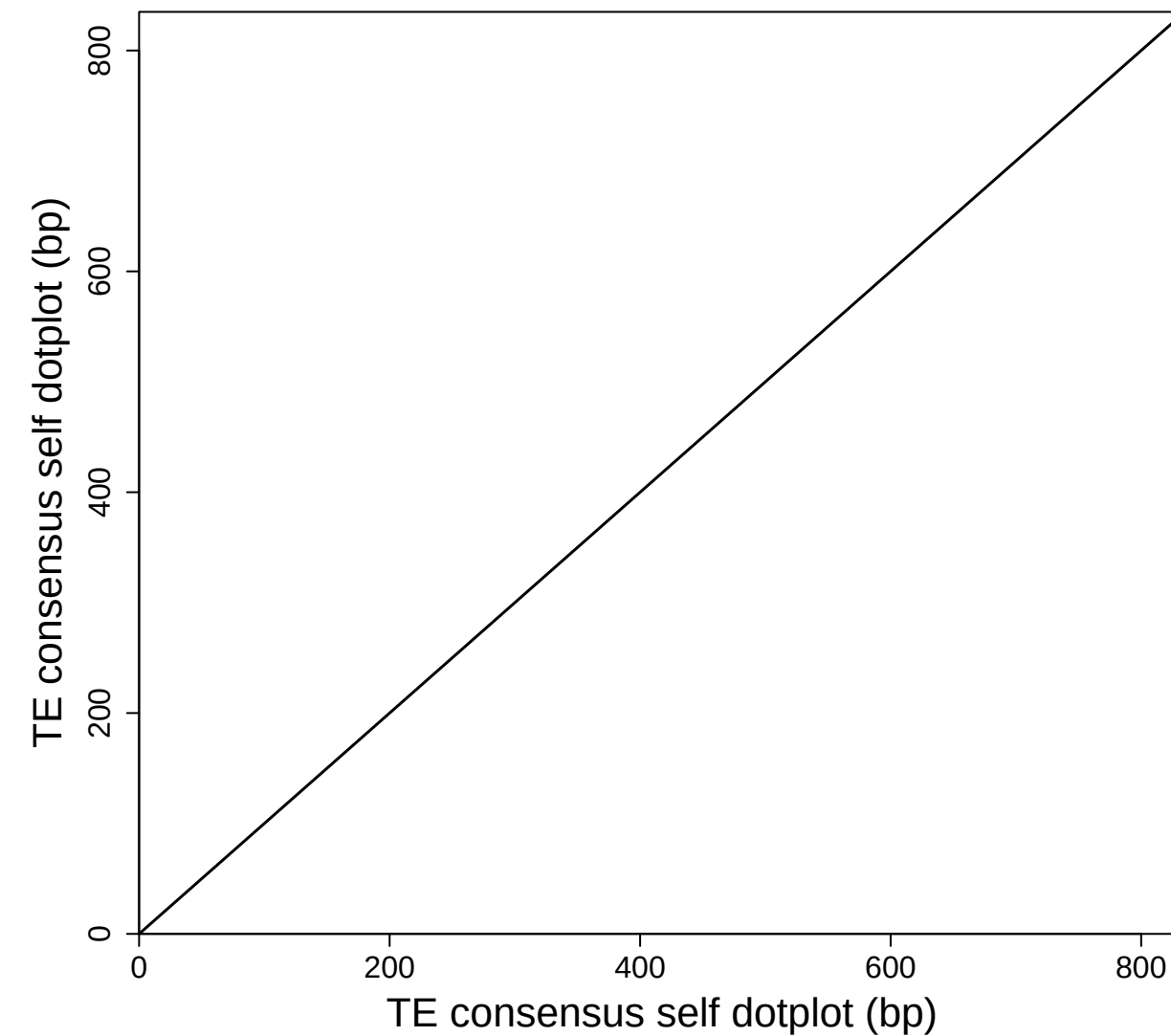
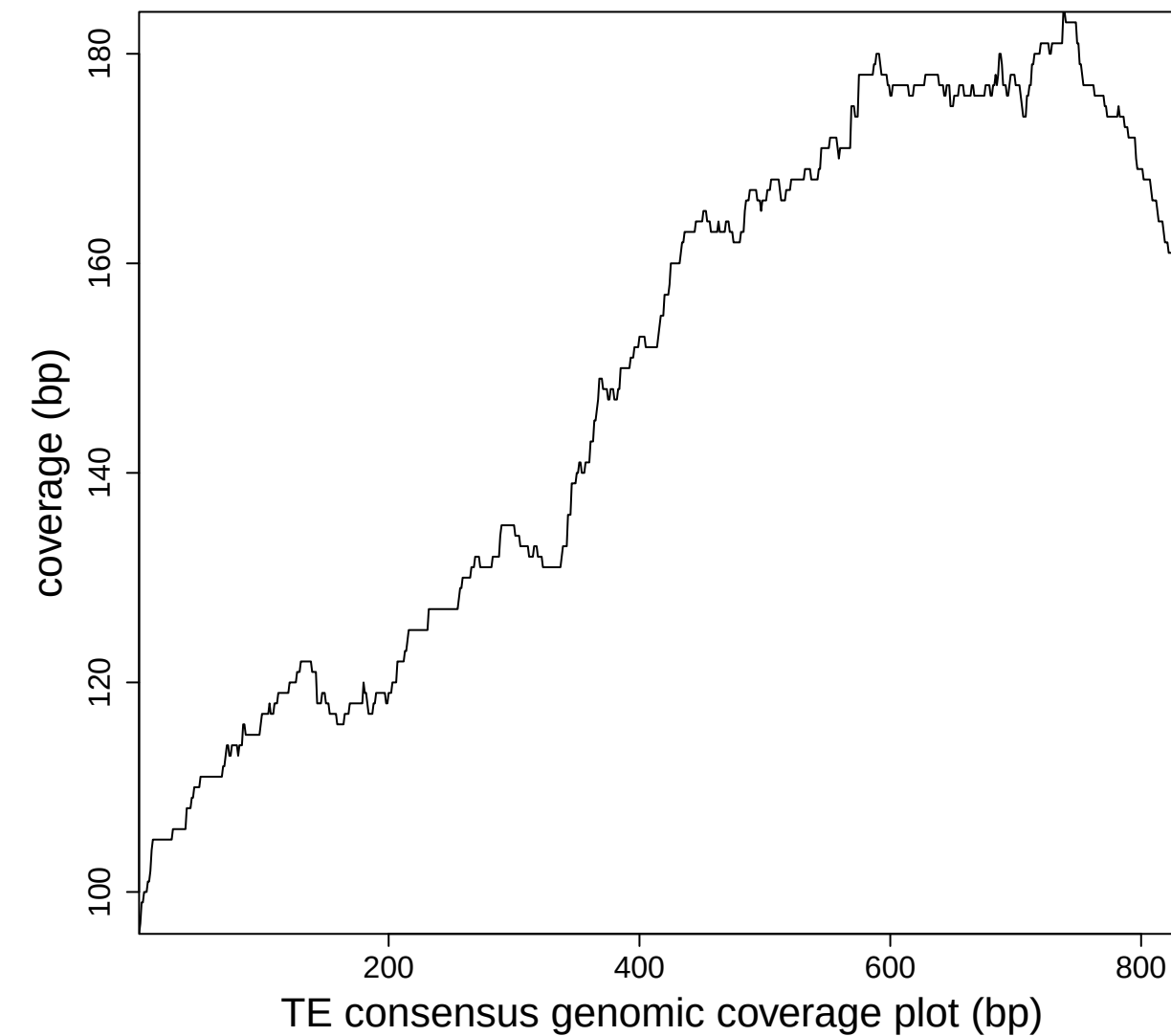
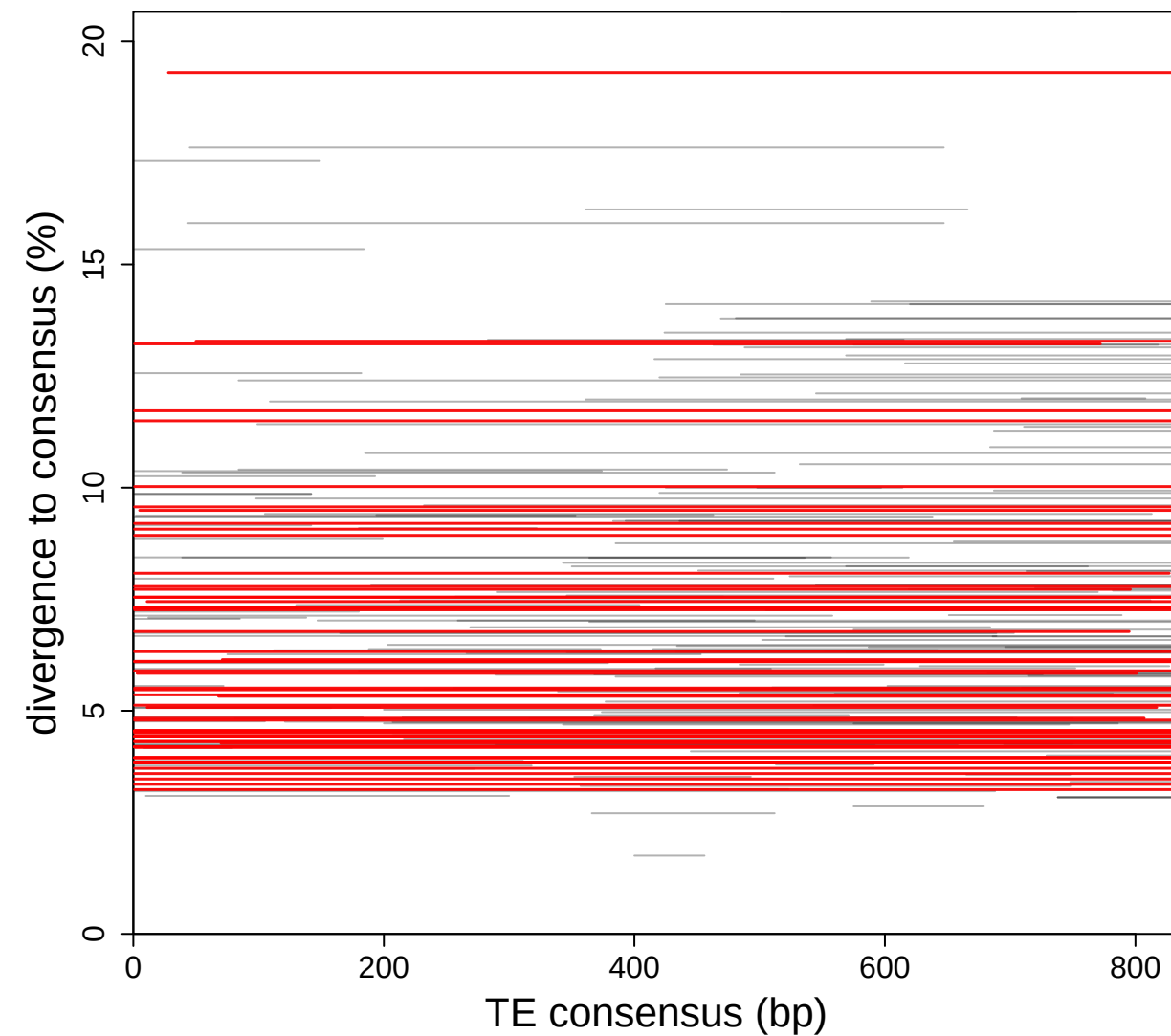


94.fasta.b.bed\_uf.bed\_g\_3.bed\_fm\_1.bed\_0\_0\_bcln.fa\_aln.fa\_cl.1  
size: 1572bp; fragments: 299; full length: 0 (>=1414.8bp)

# After TEtrimmer Extended plot Blue lines are boundaries

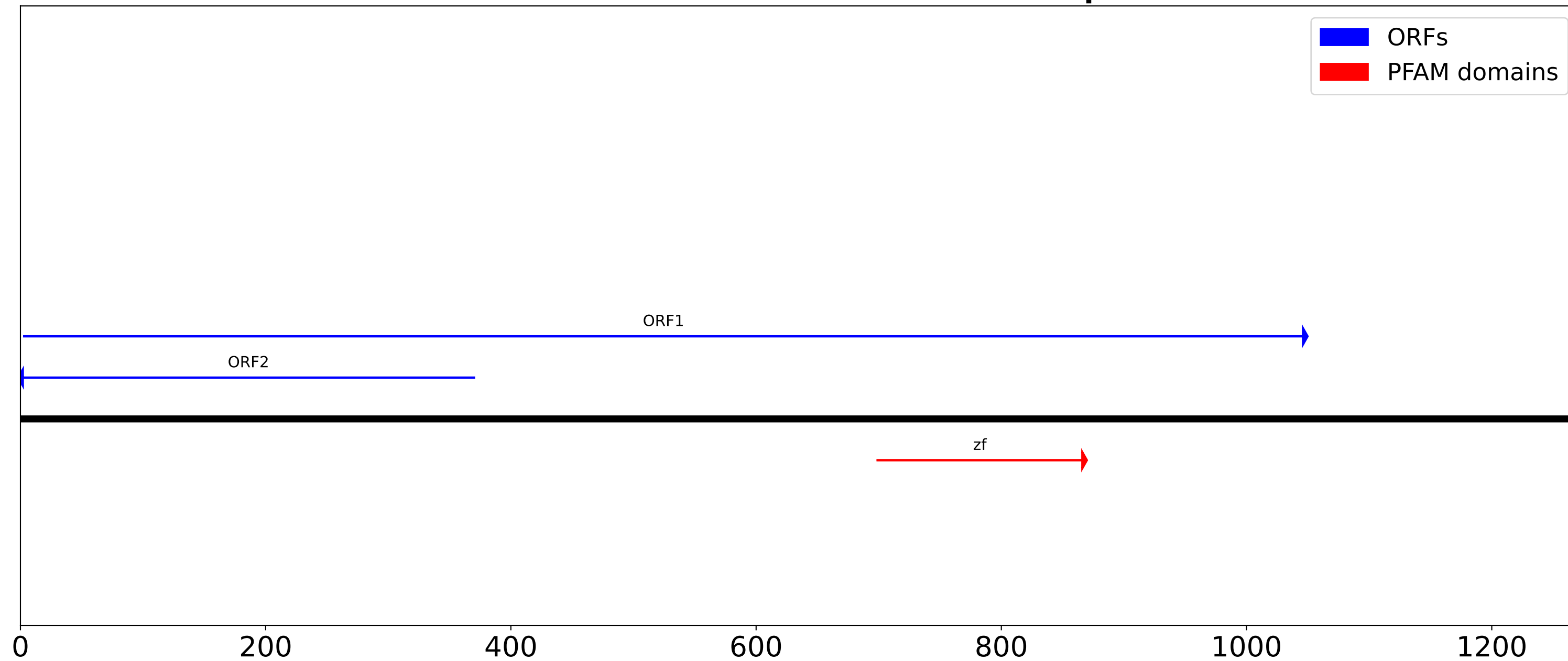


TE: rnd\_5\_family\_19694  
size: 835bp; fragments: 256; full length: 72 ( $\geq 751.5$ bp)

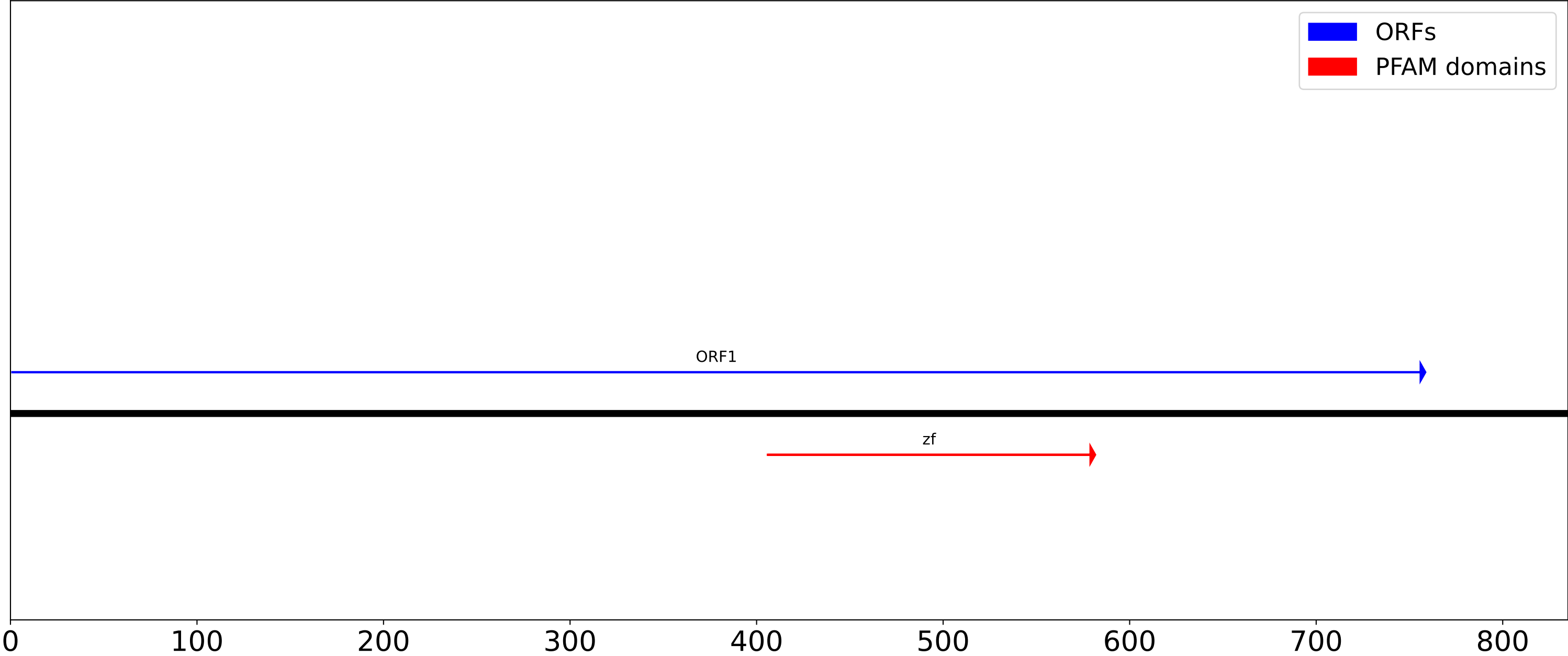


Before TEtrimmer 835 bp

# After TEtrimmer ORF and PFAM domain plot



**Before TEtrimmer ORF and PFAM domain plot**





# Dotplot

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

