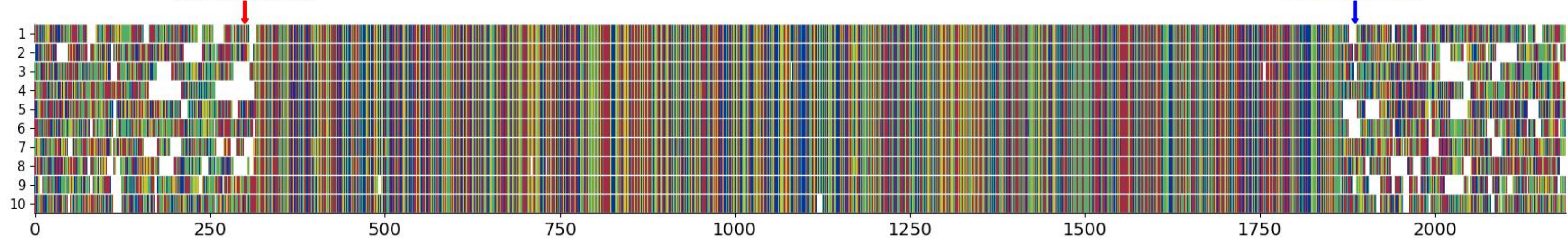


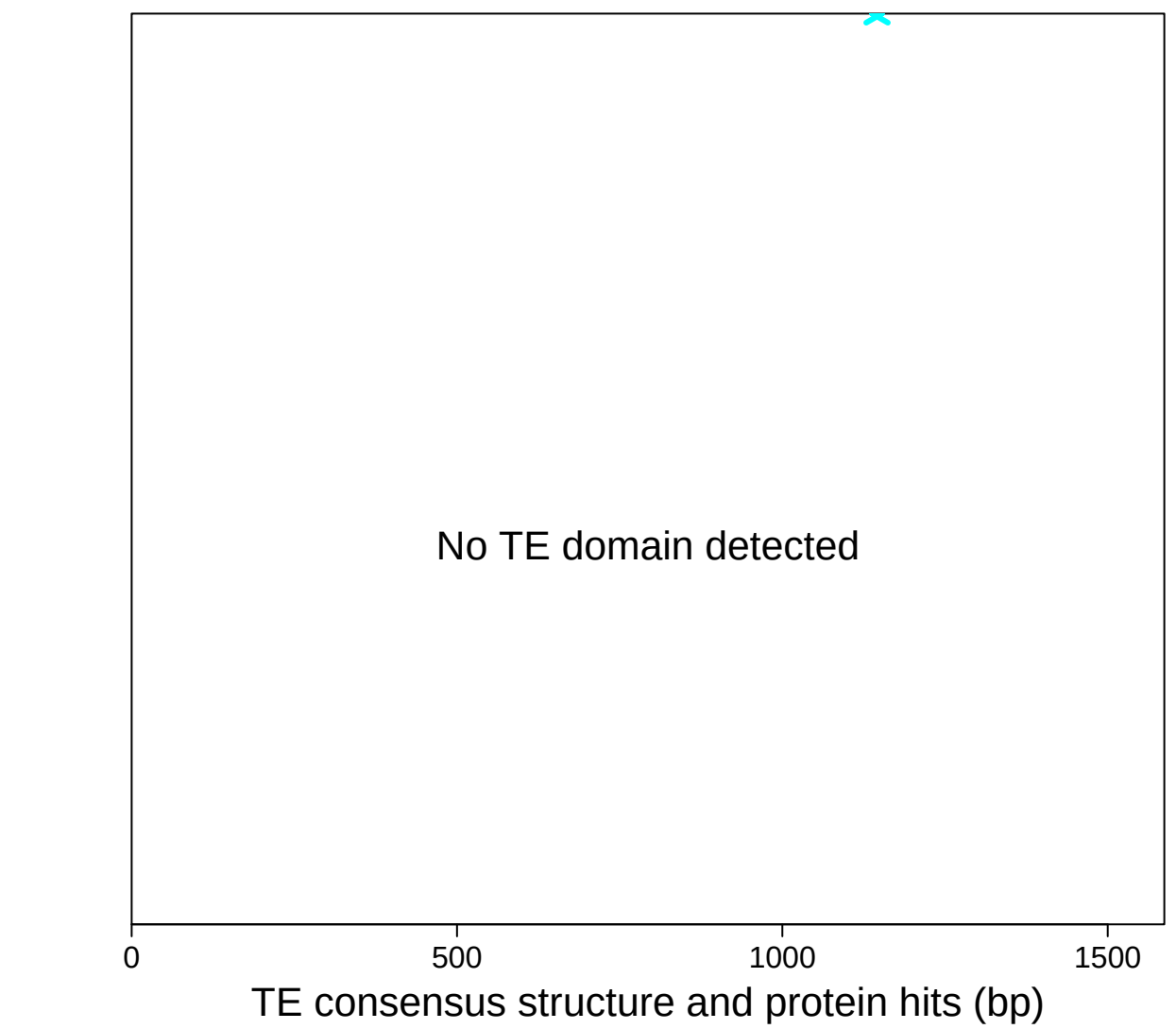
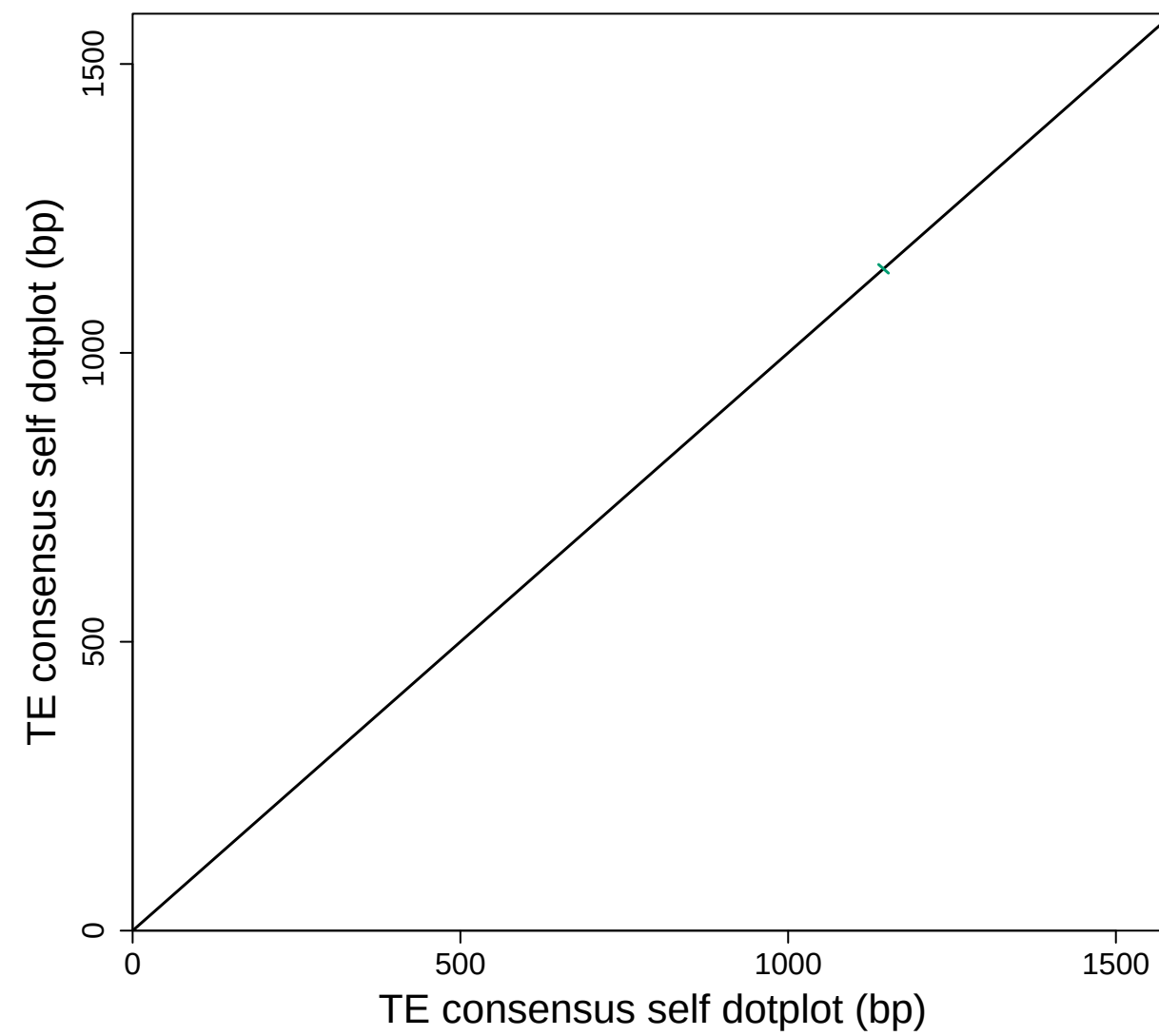
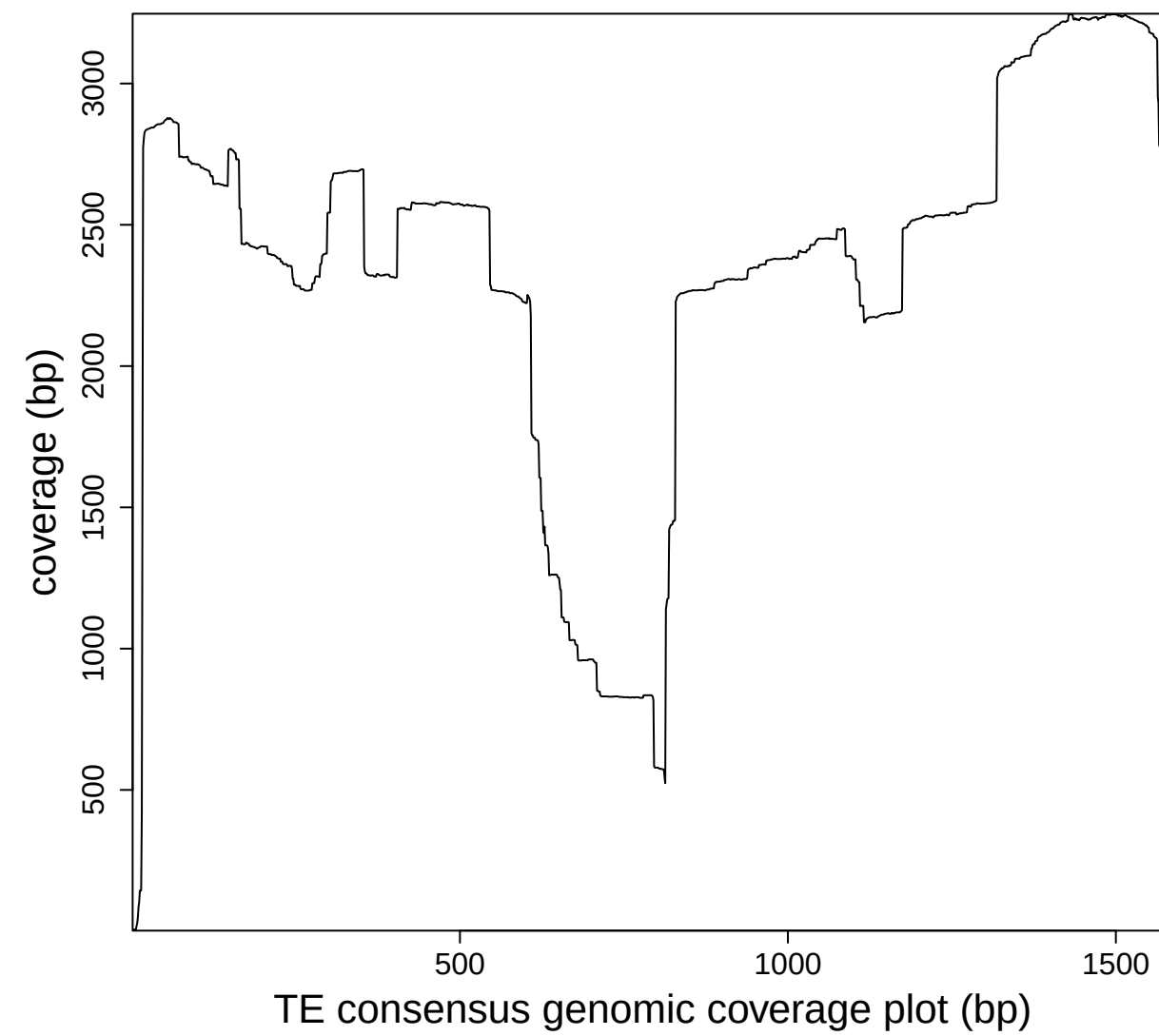
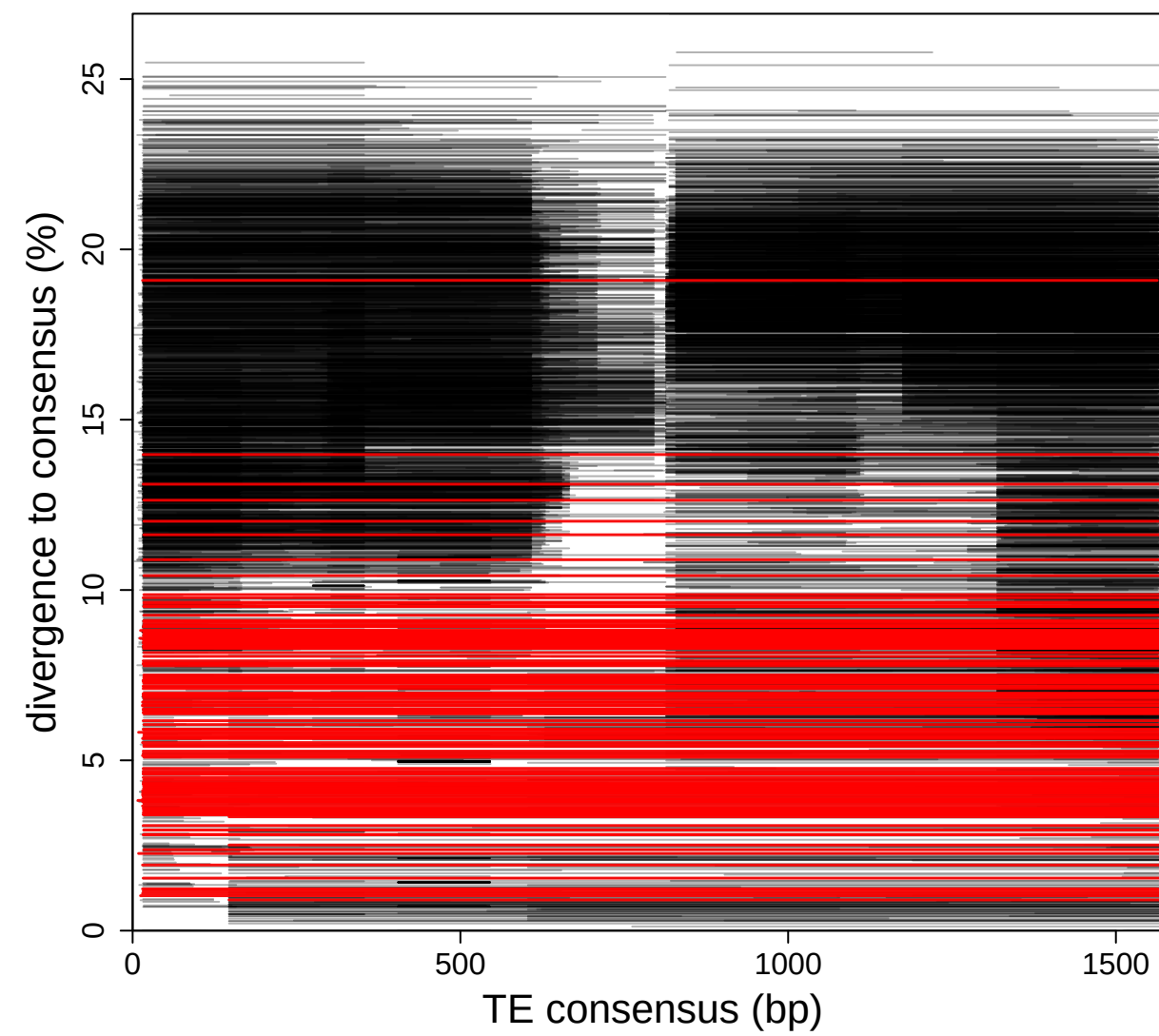
Start crop Point

End crop Point



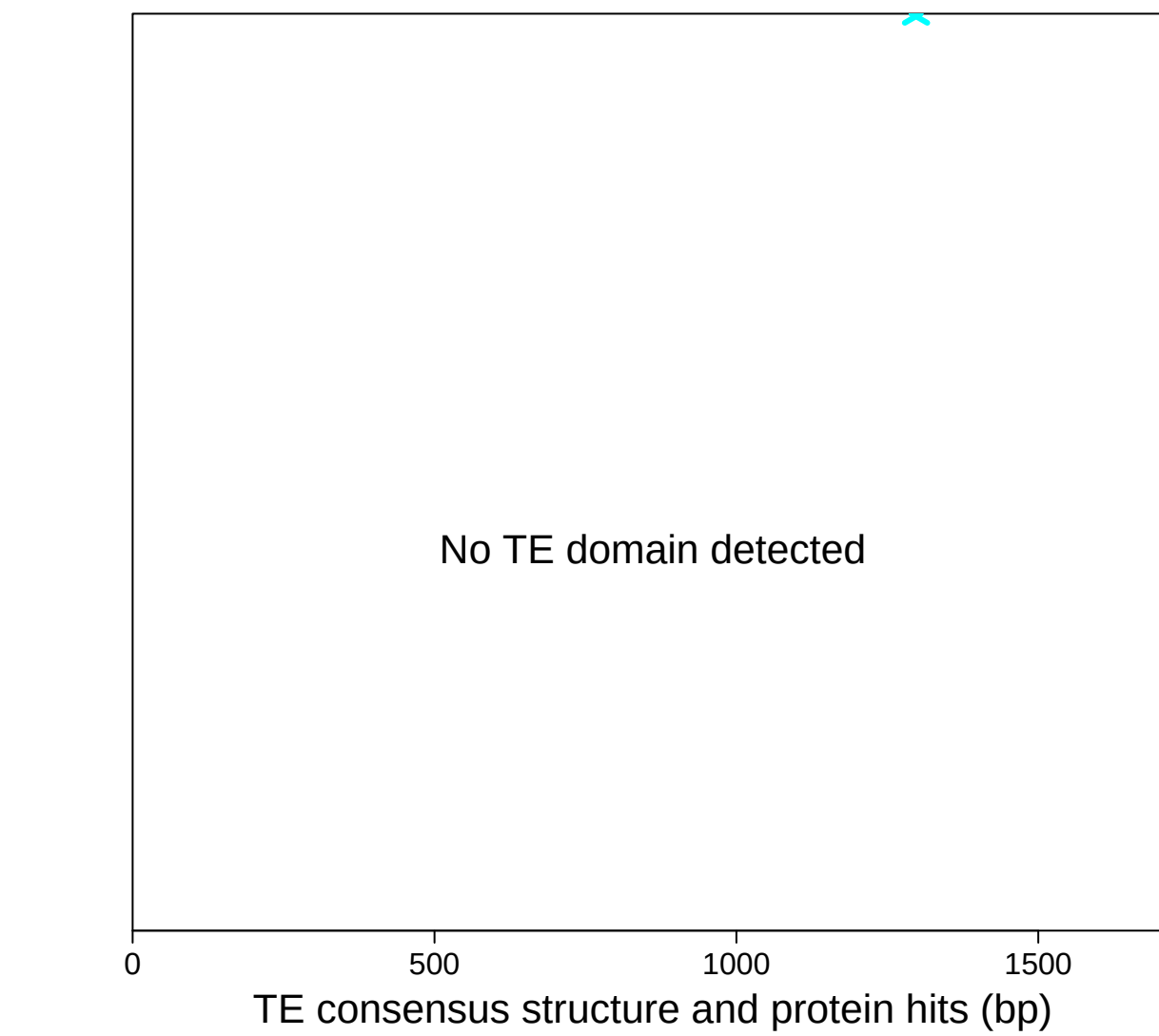
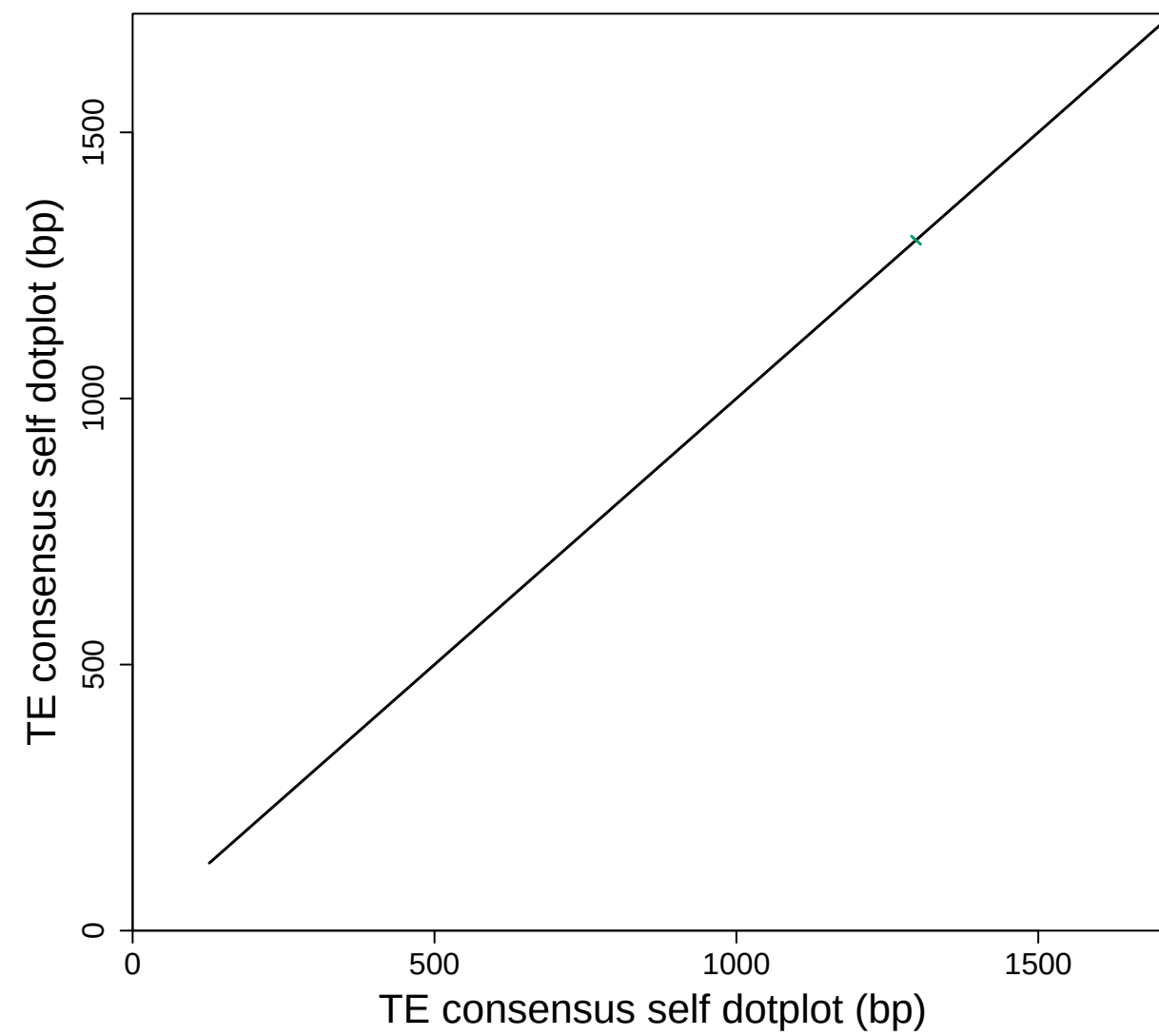
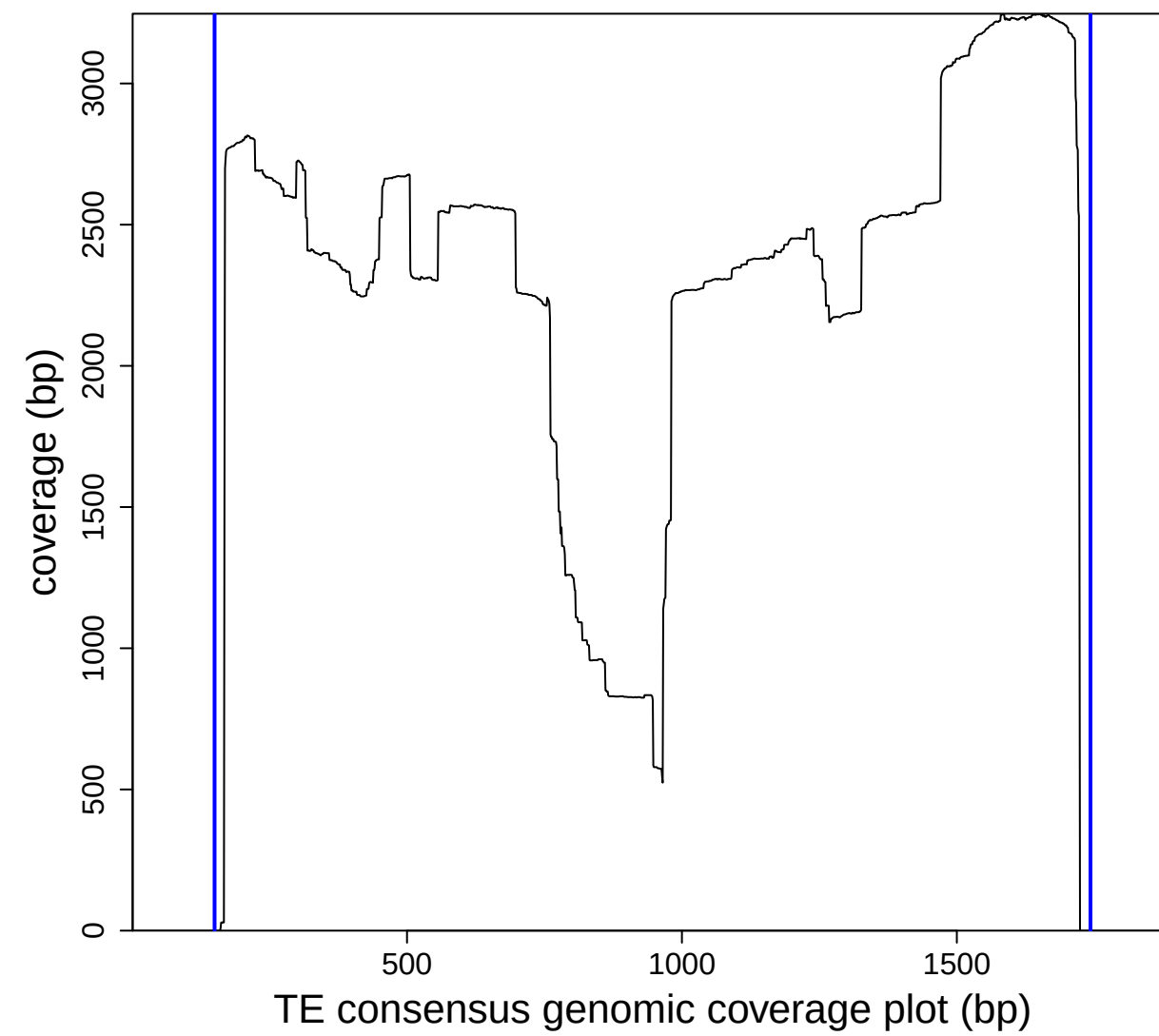
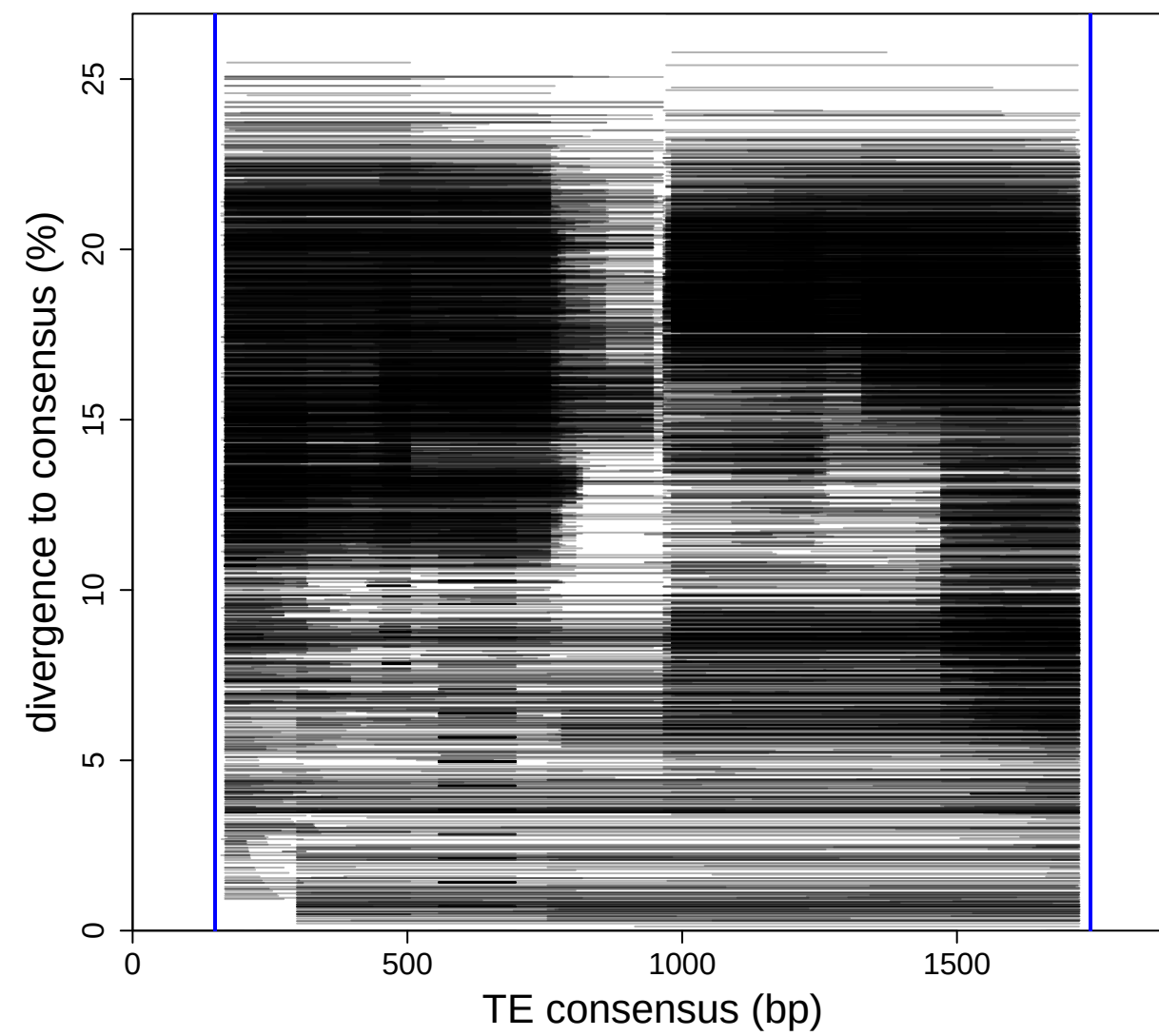
.l.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 1587bp; fragments: 7395; full length: 180 (≥ 1428.3 bp)

After TEtrimmer 1587 bp

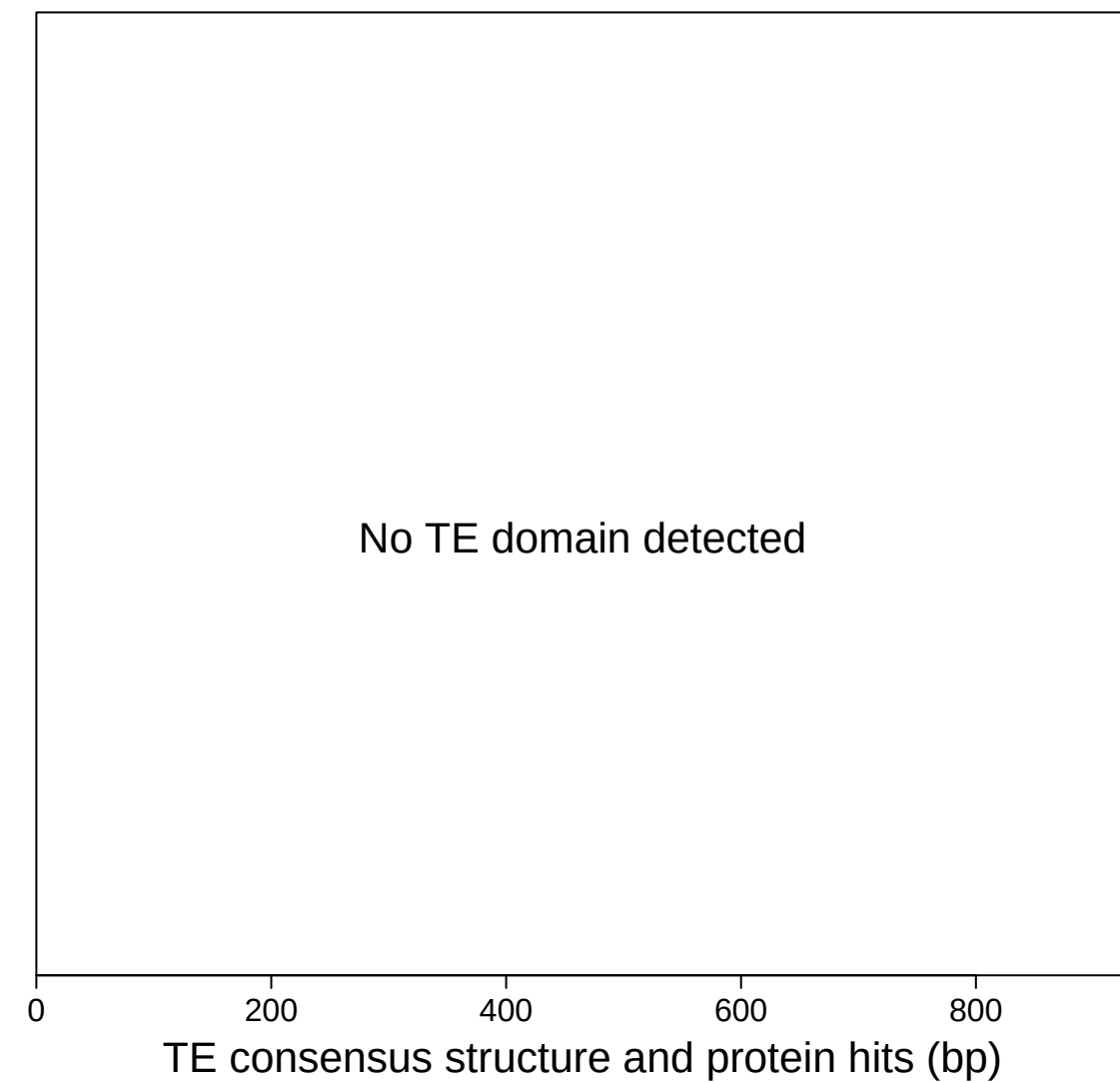
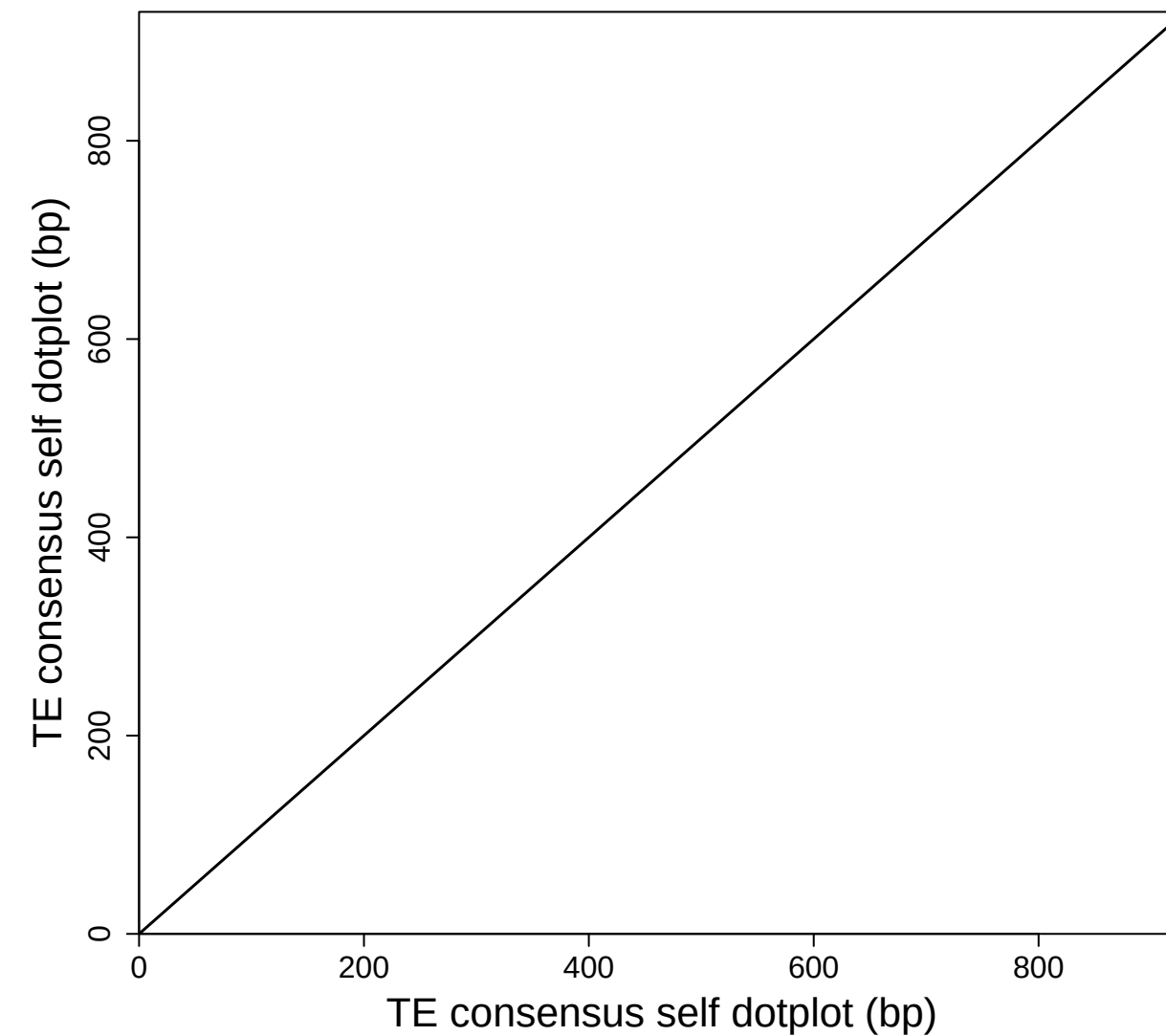
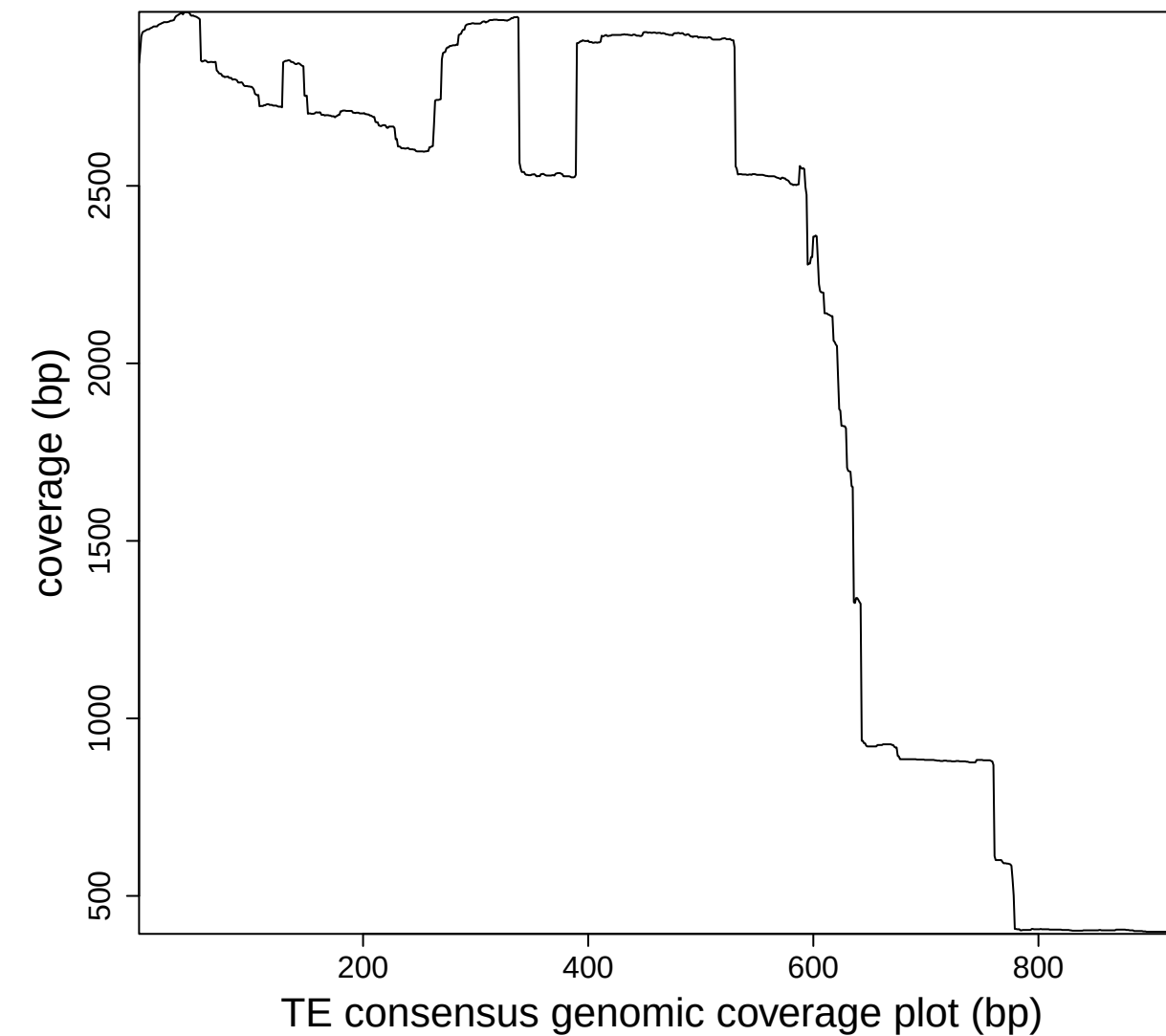
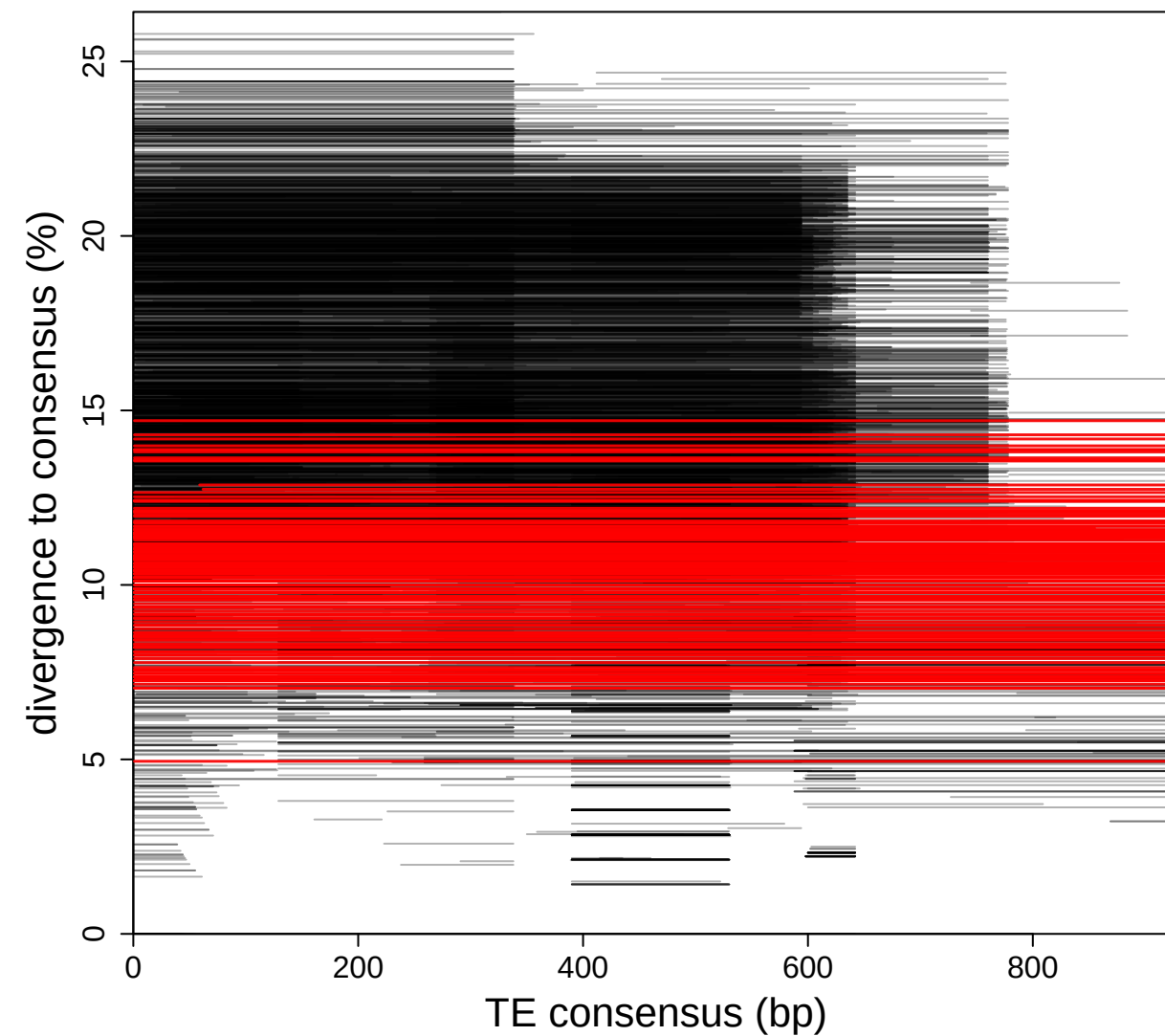


0.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 1893bp; fragments: 7331; full length: 0 (>=1703.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

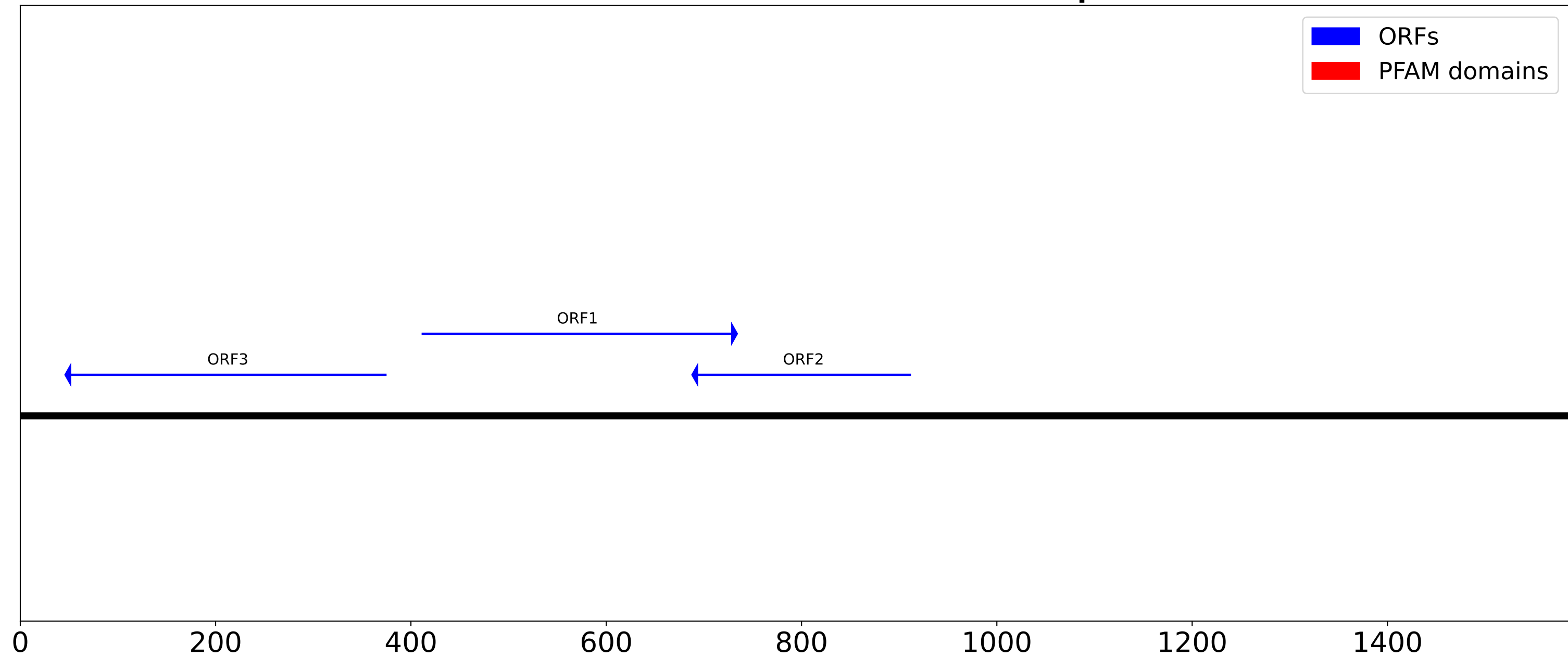


TE: rnd_1_family_660
size: 930bp; fragments: 4359; full length: 193 (≥ 837 bp)

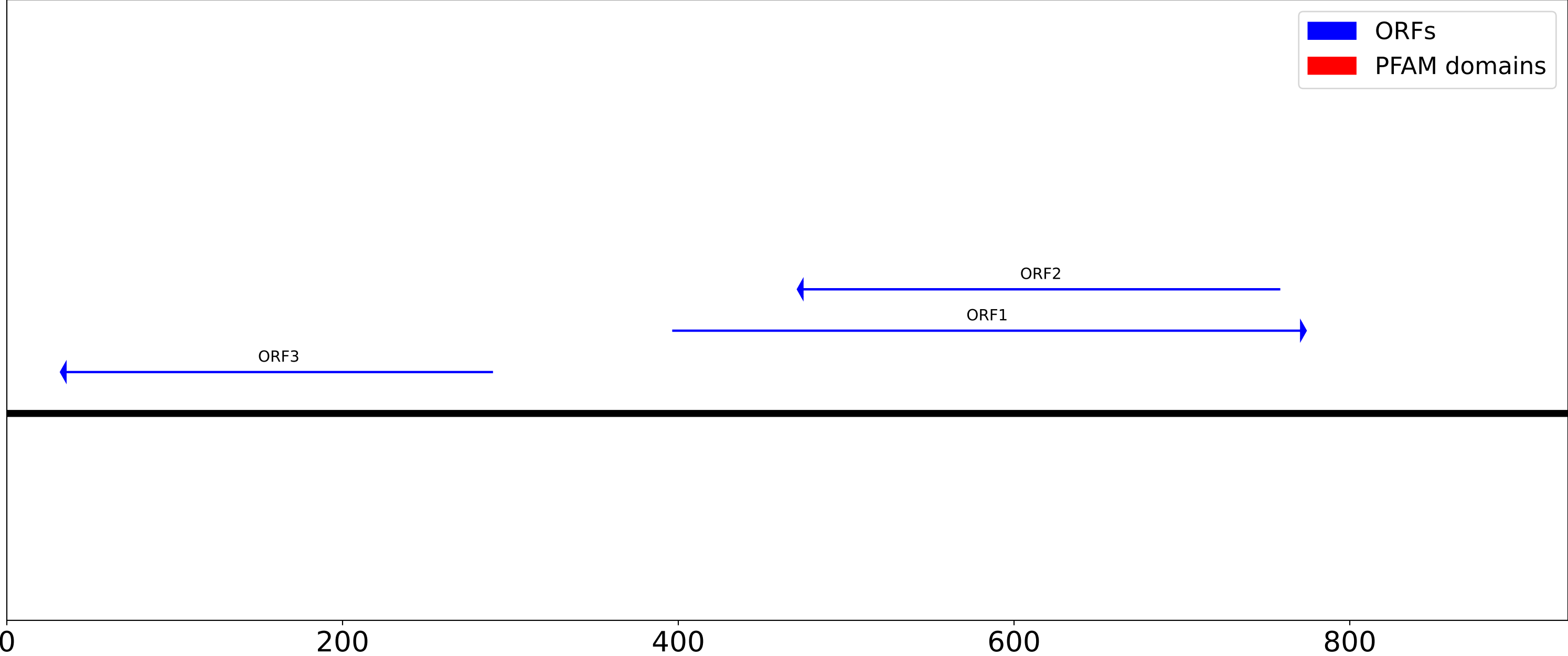


Before TEtrimmer 930 bp

After TEtrimmer ORF and PFAM domain plot



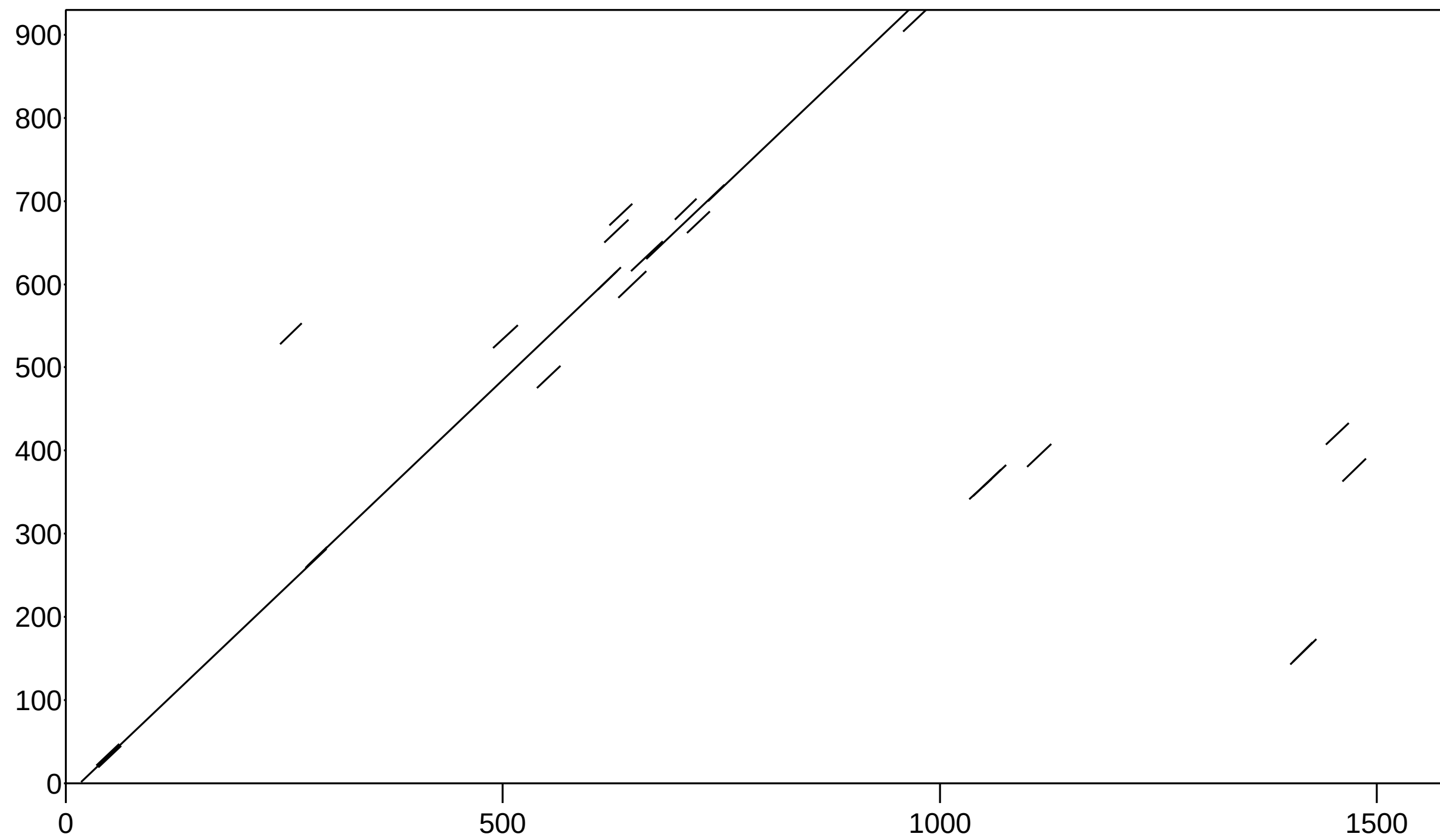
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)