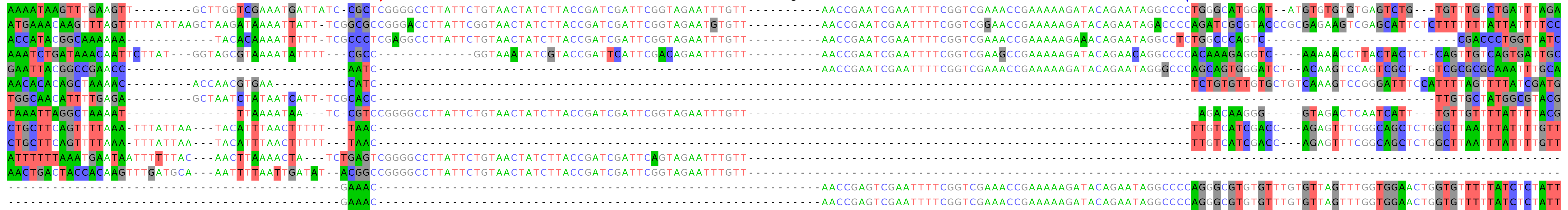


Start crop Point

End crop Point

MSA length = 1318



Start crop Point

End crop Point

0

10

0

250

500

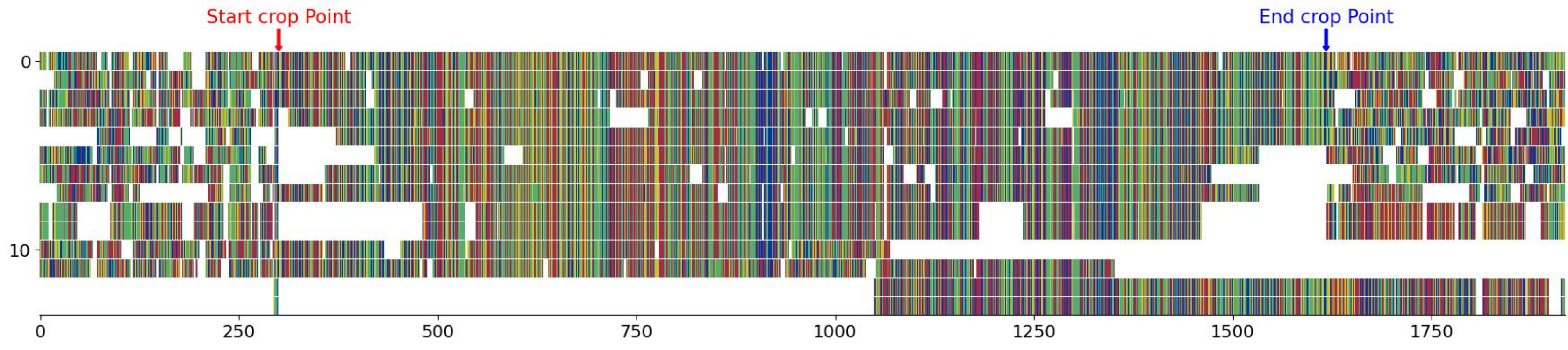
750

1000

1250

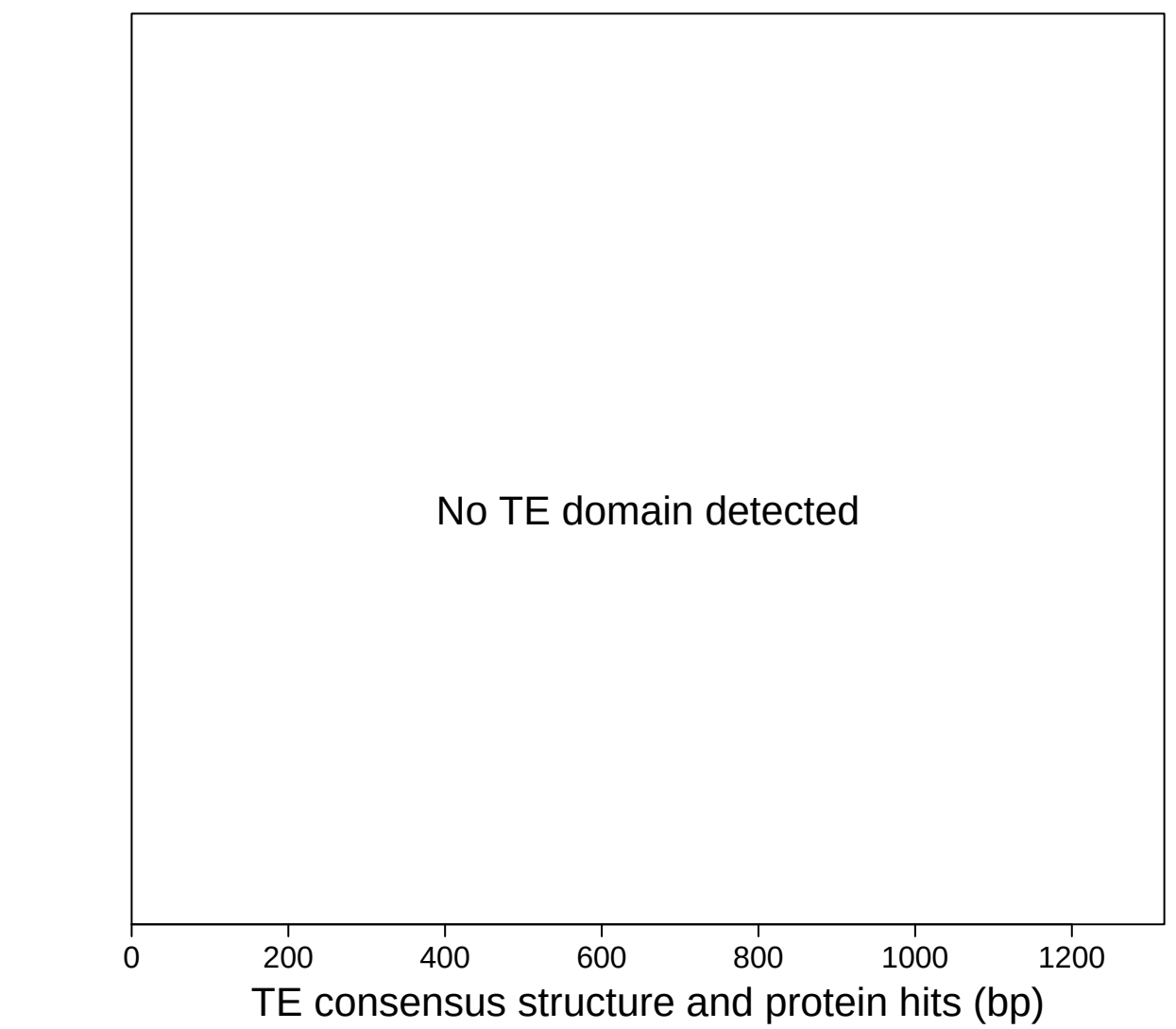
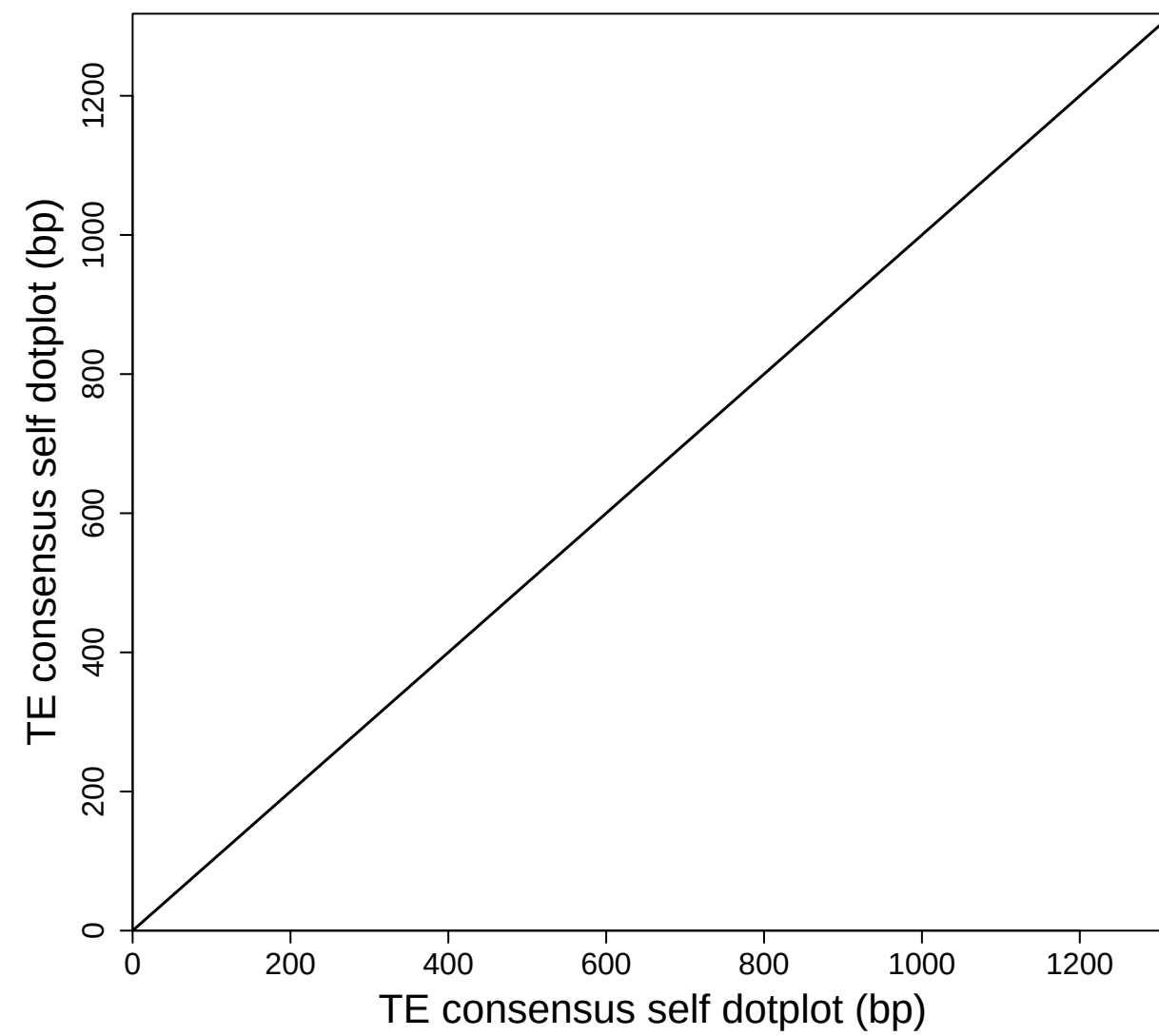
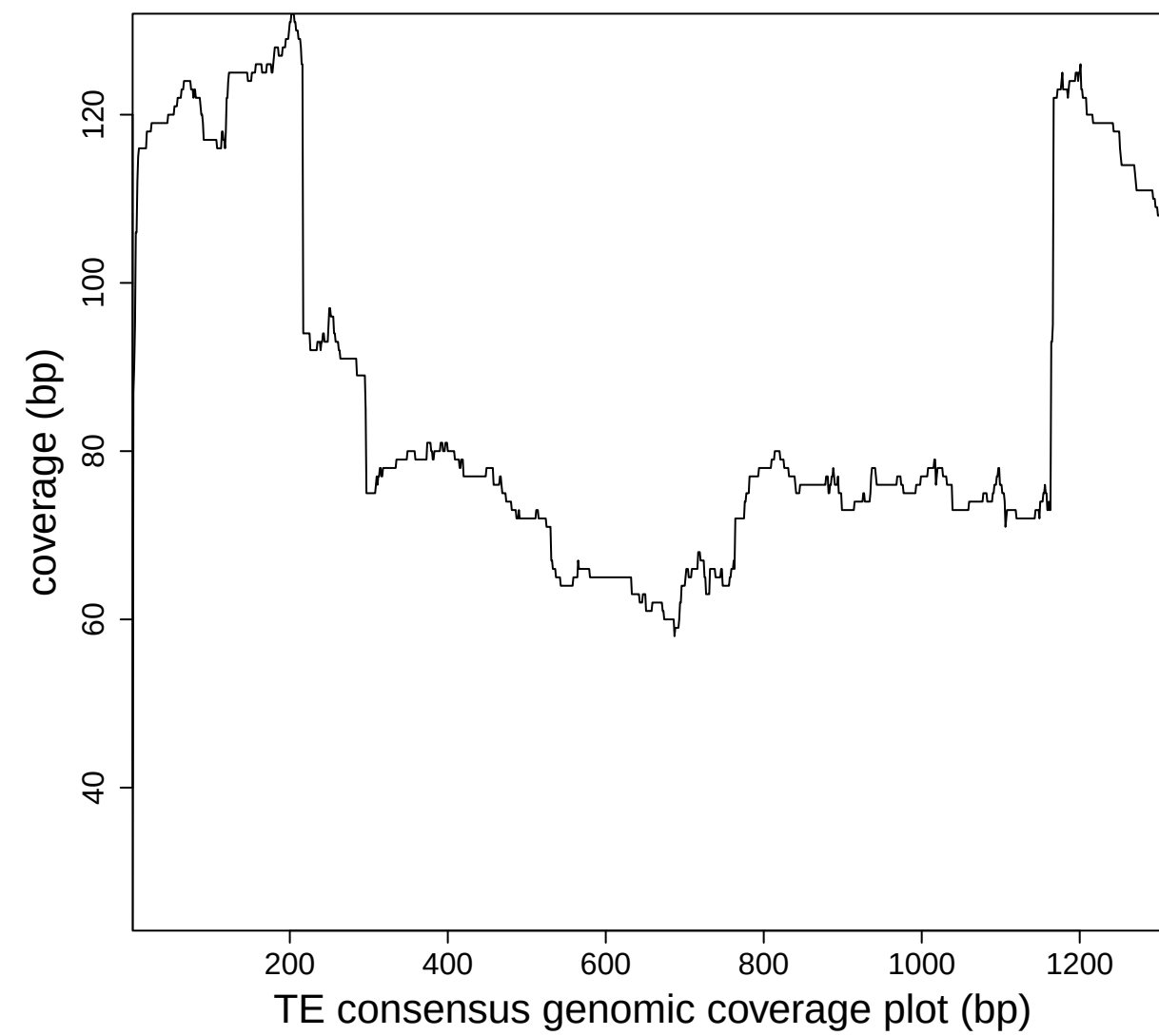
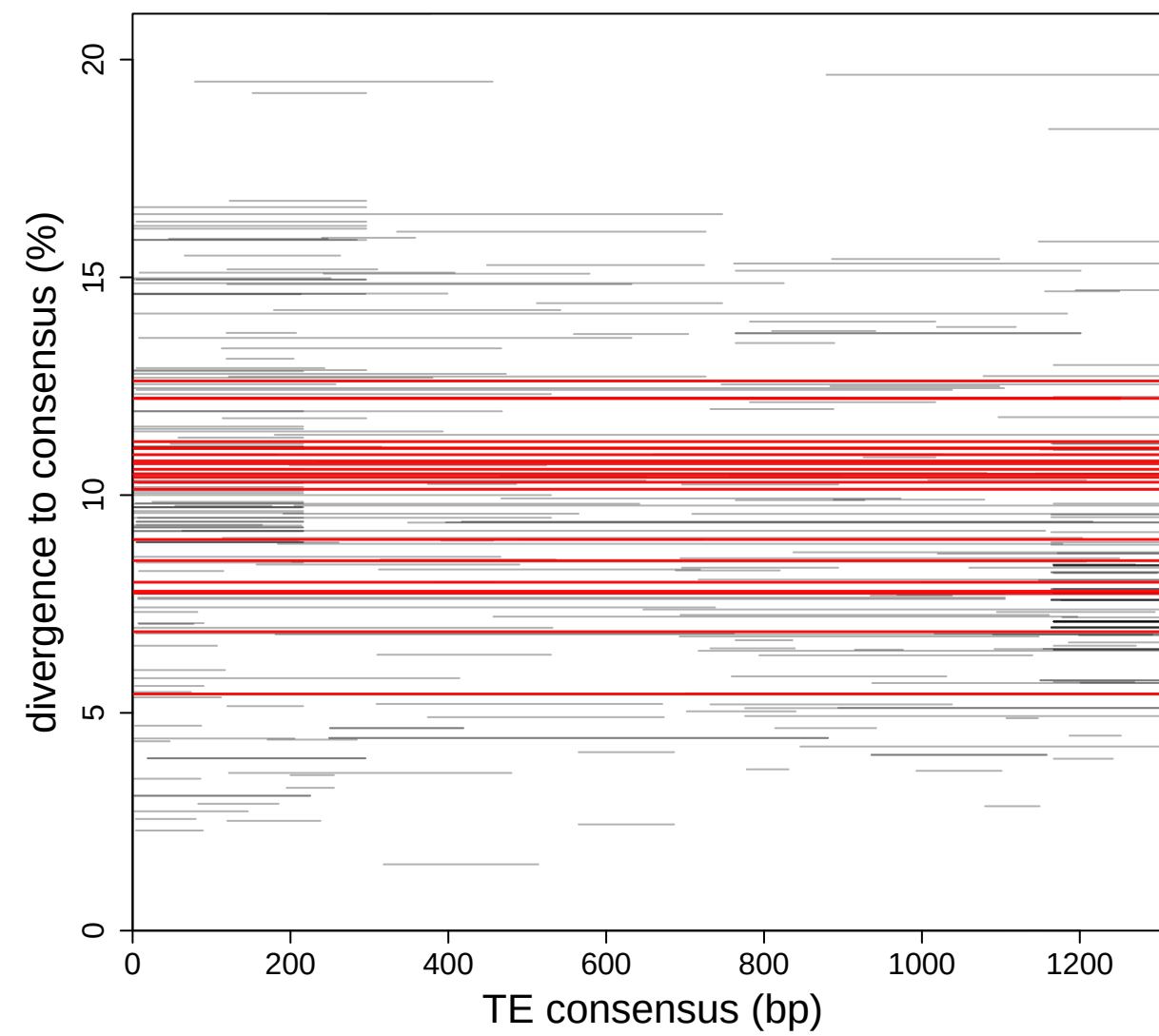
1500

1750



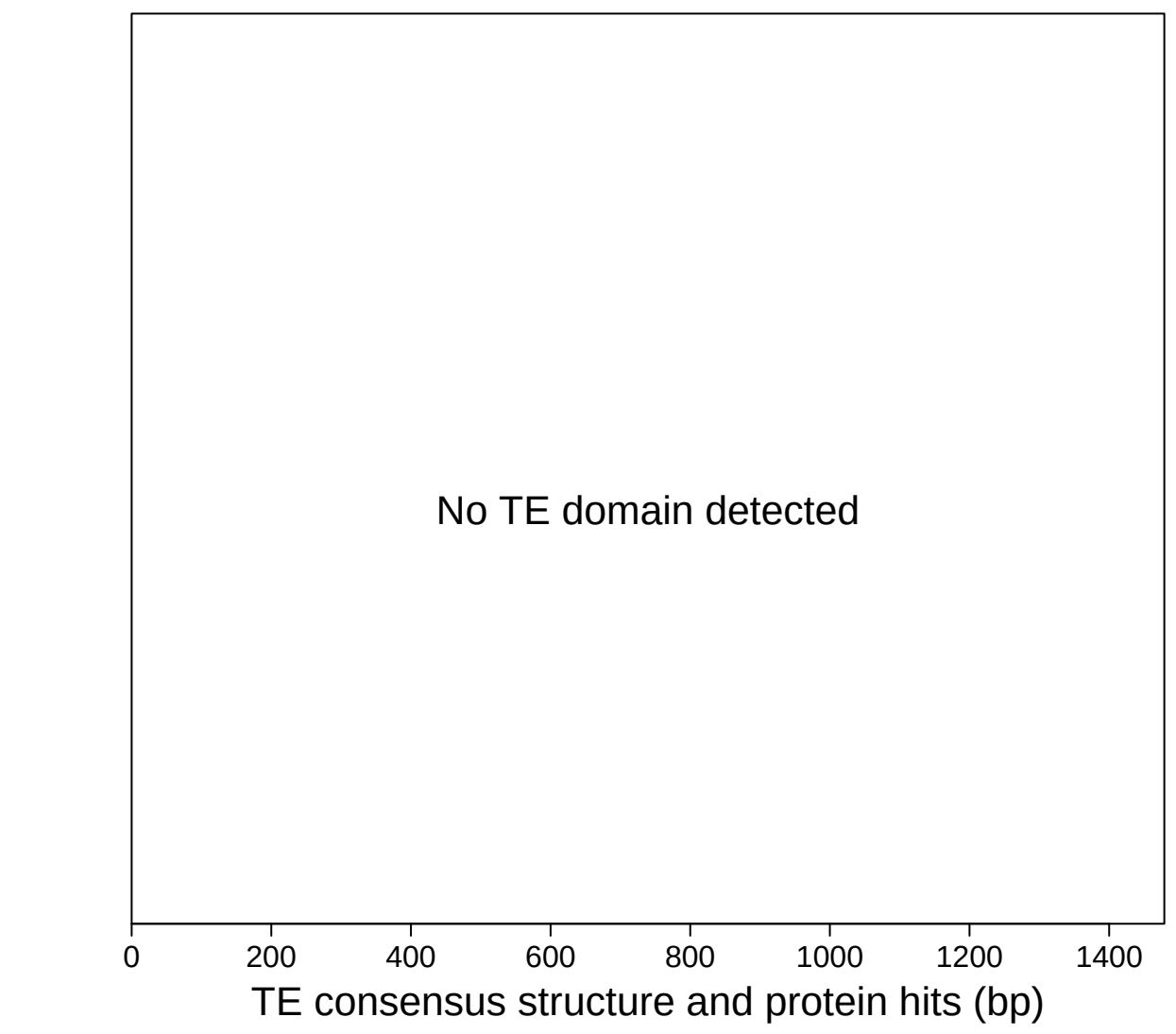
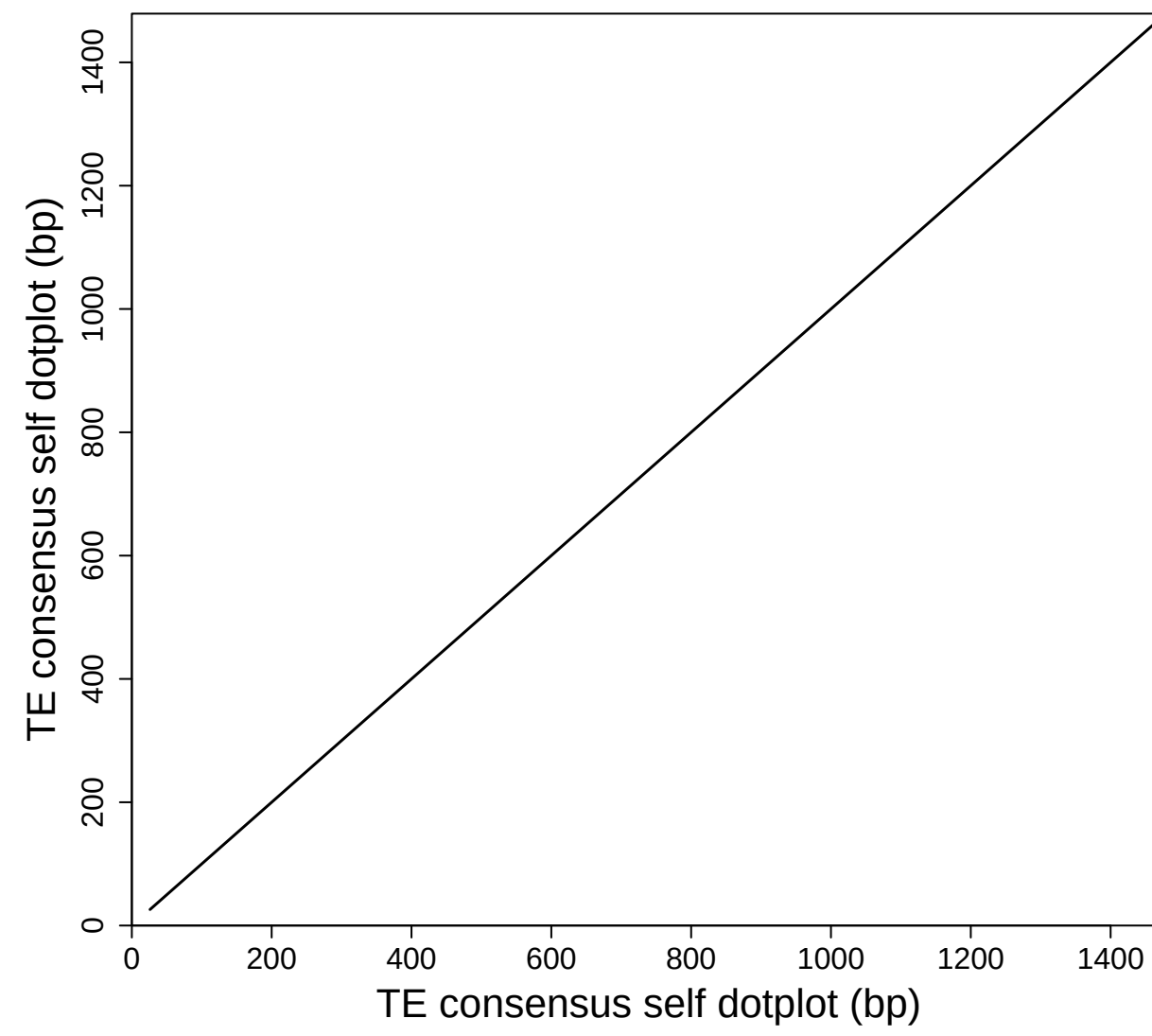
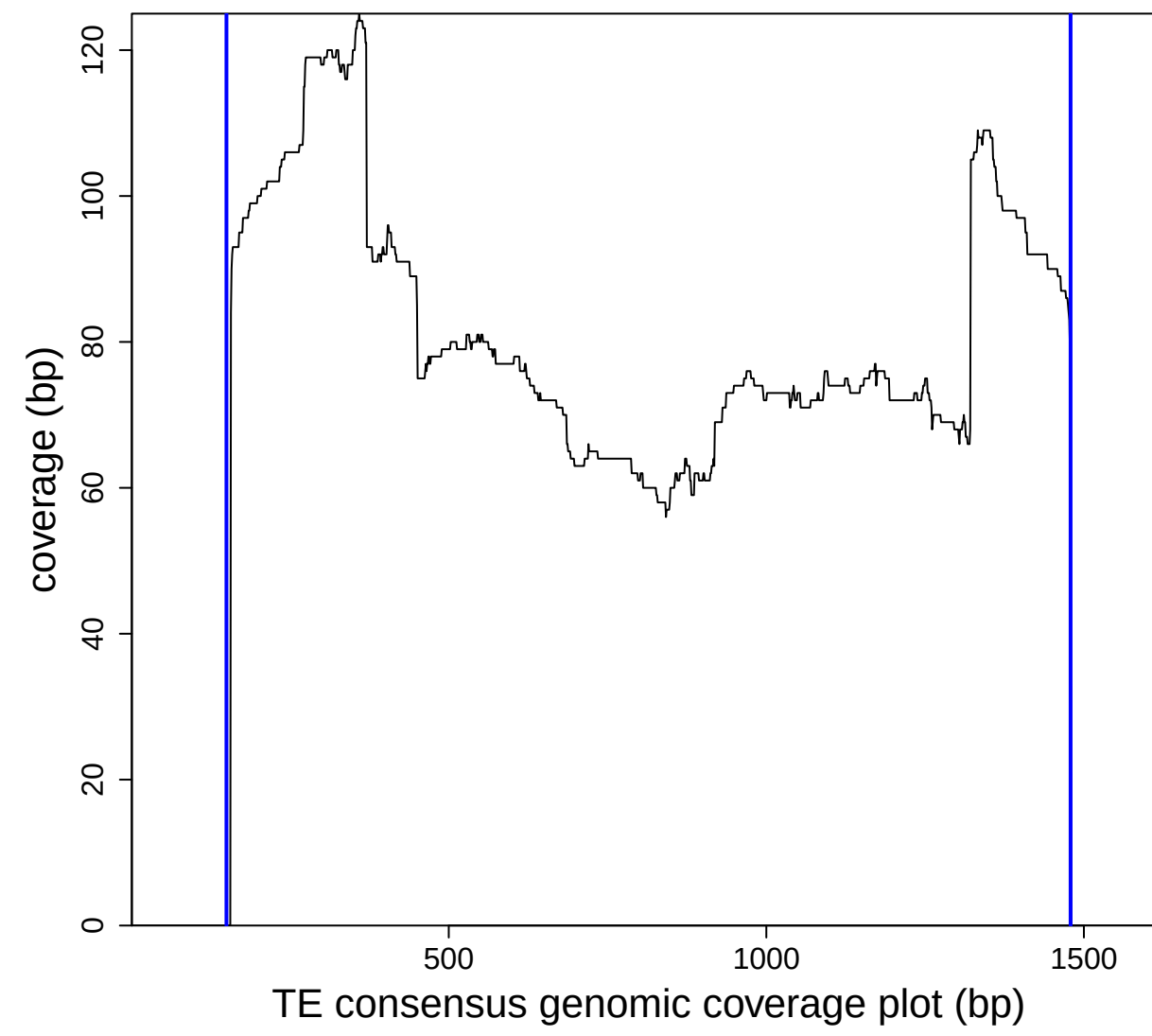
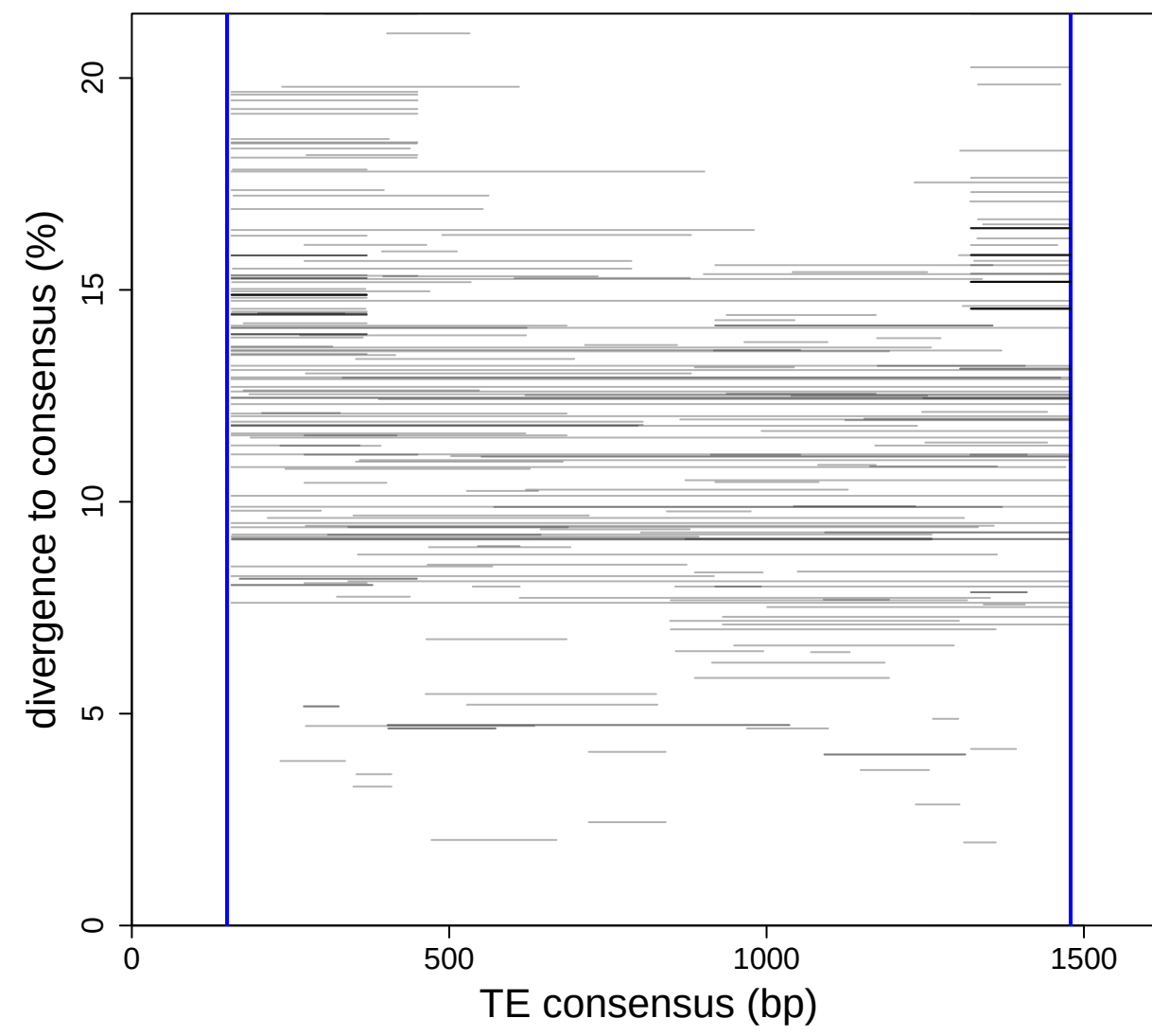
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1318bp; fragments: 312; full length: 21 (≥ 1186.2 bp)

After TEtrimmer 1318 bp

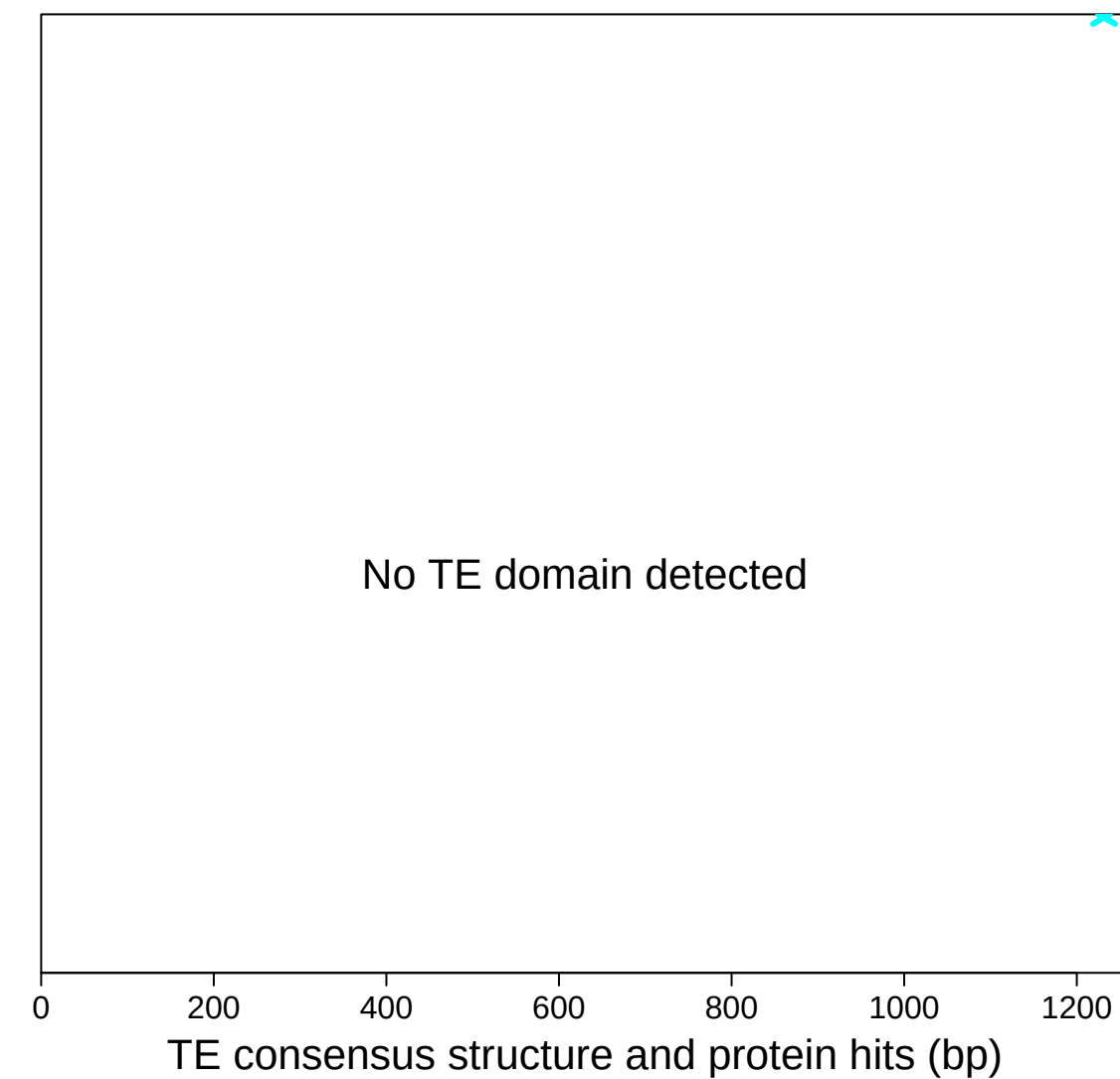
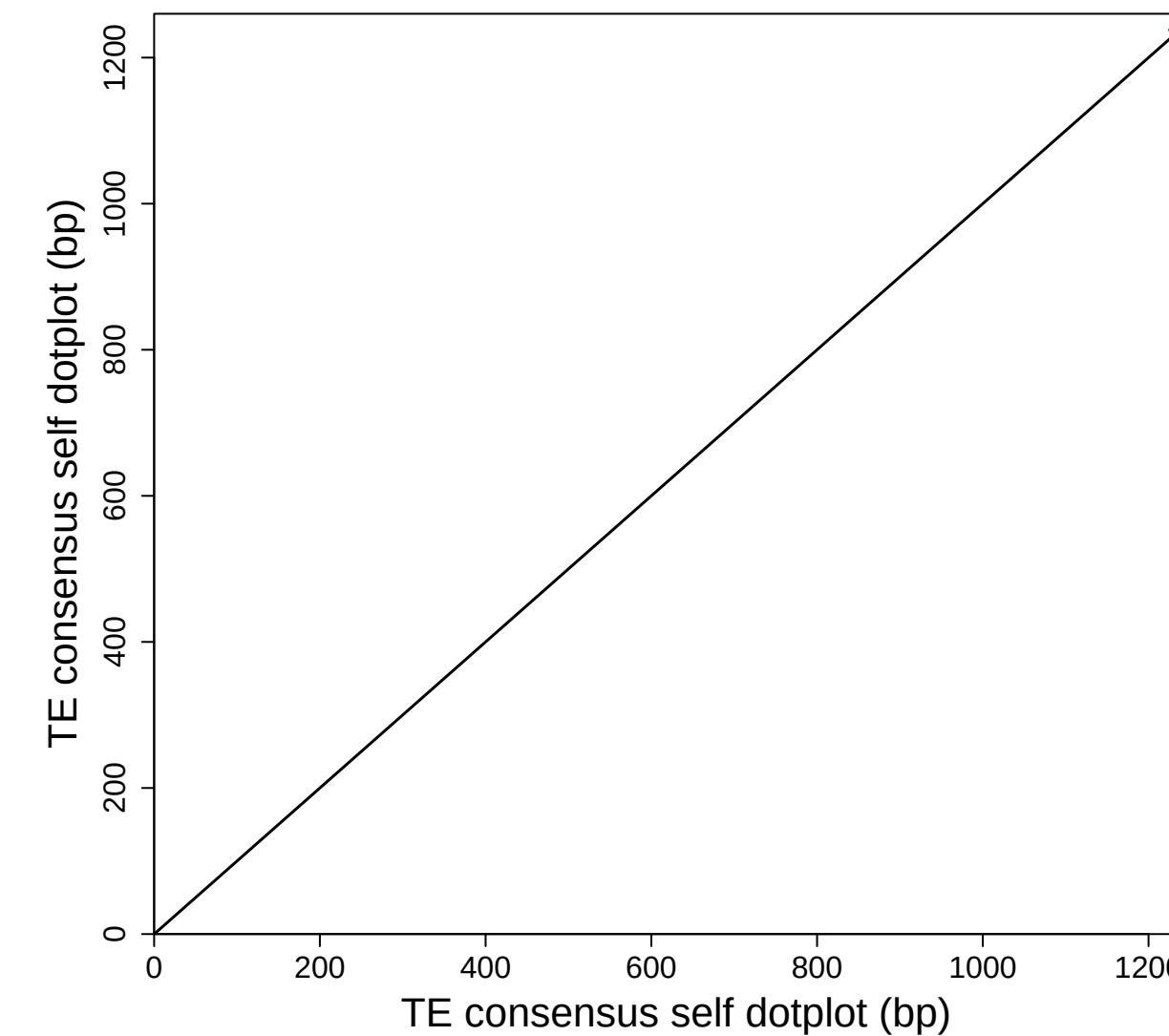
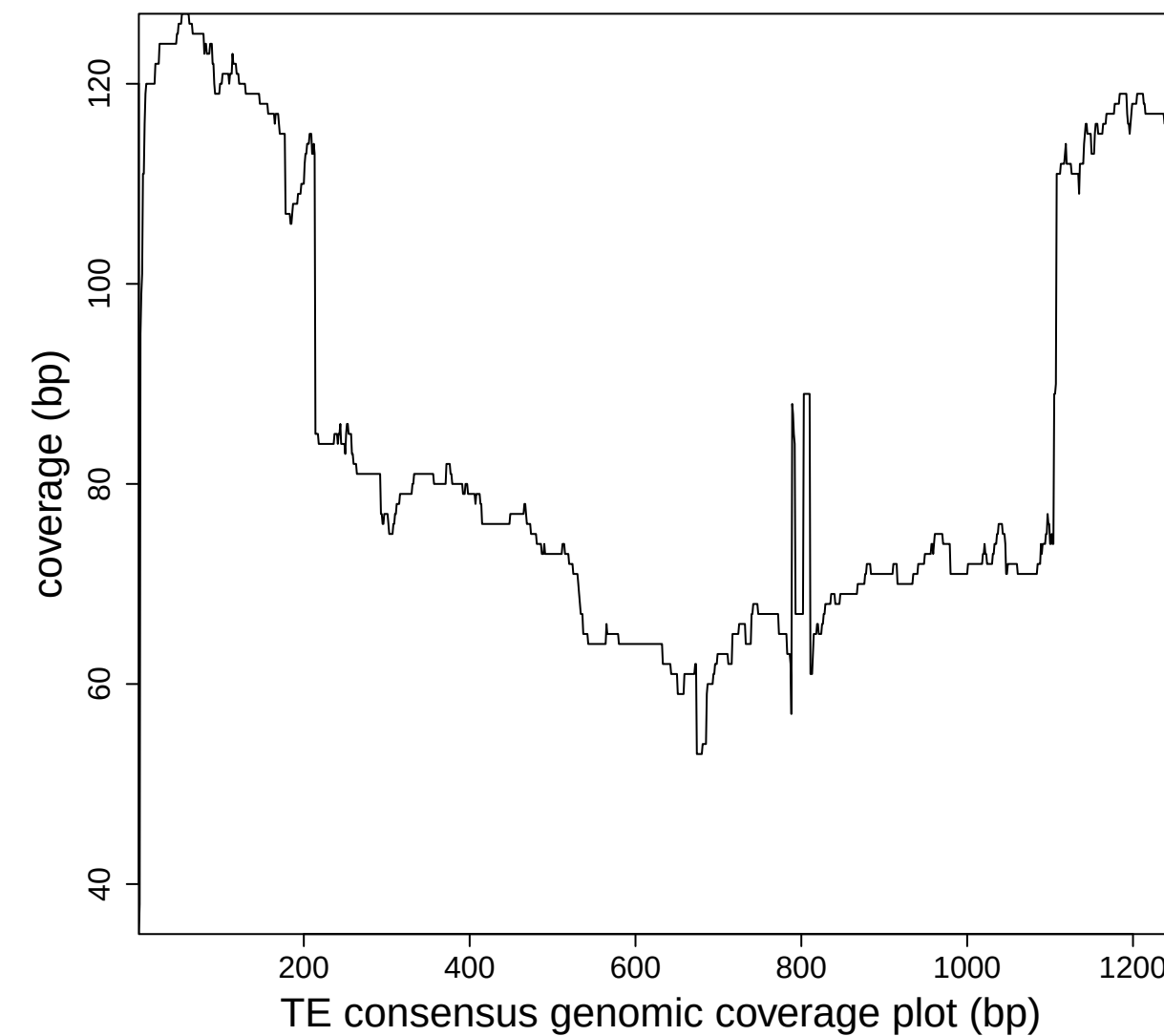
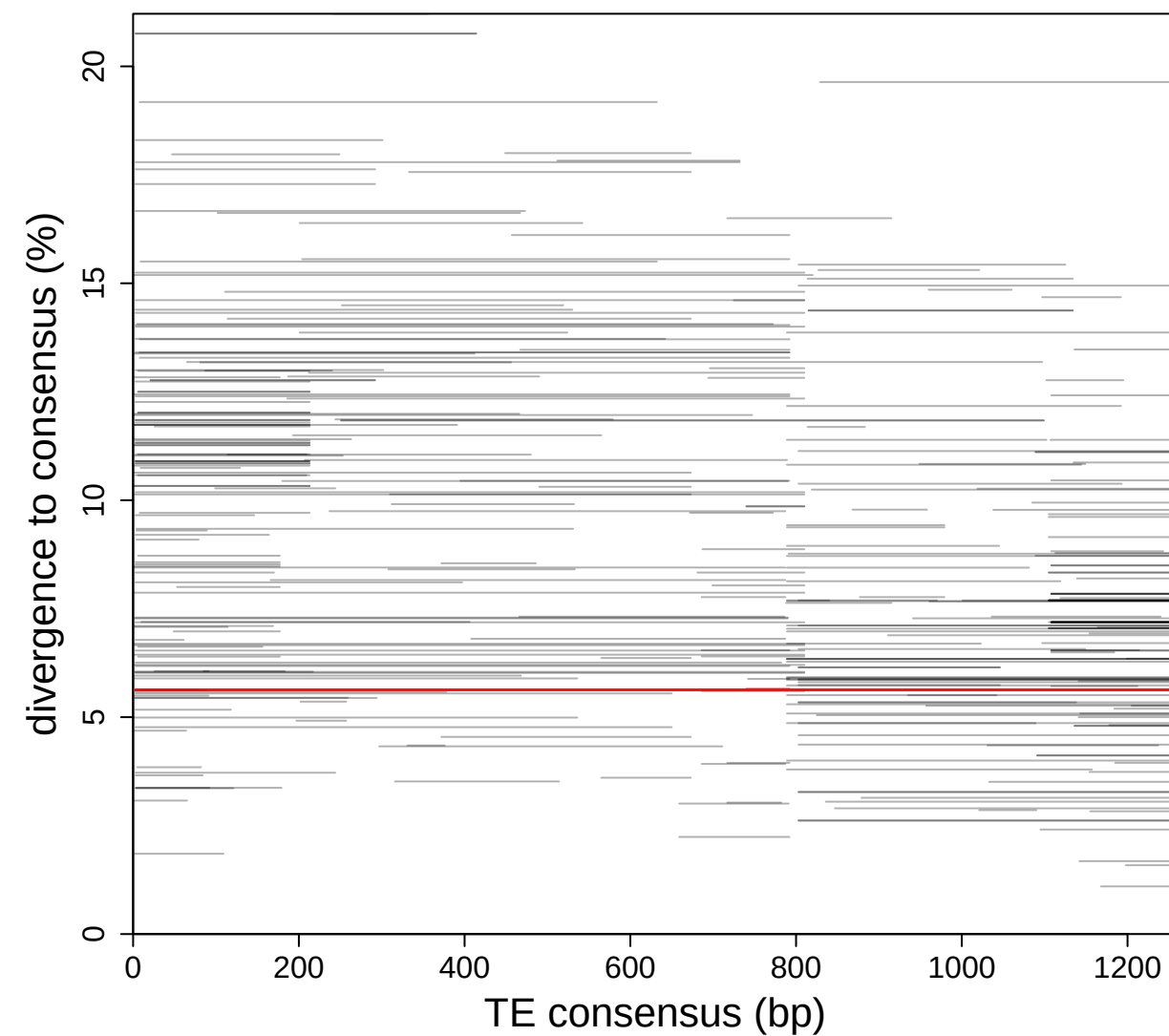


9.fasta.b.bed uf.bed g_2.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 1629bp; fragments: 268; full length: 0 (>=1466.1bp)

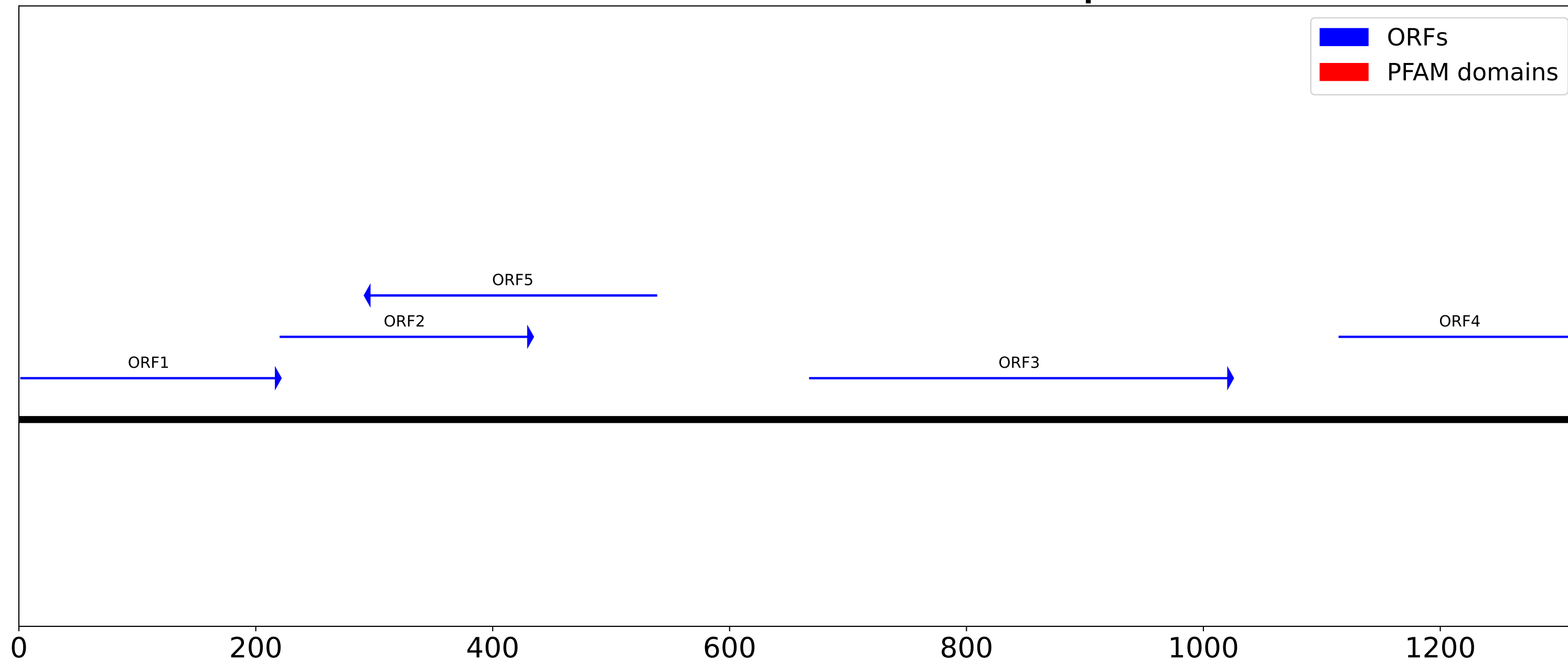
After TEtrimmer Extended plot Blue lines are boundaries



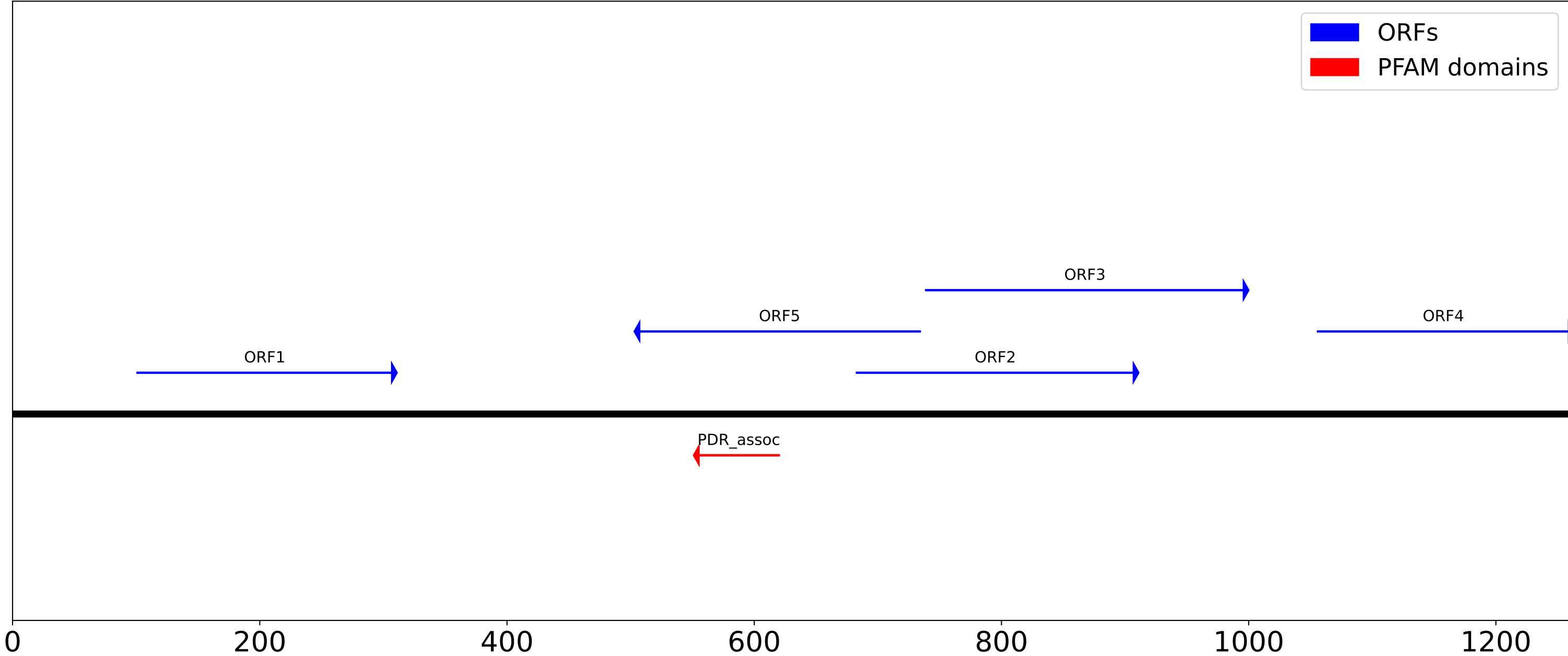
TE: rnd_5_family_7239
size: 1260bp; fragments: 349; full length: 1 (>=1134bp)



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

