

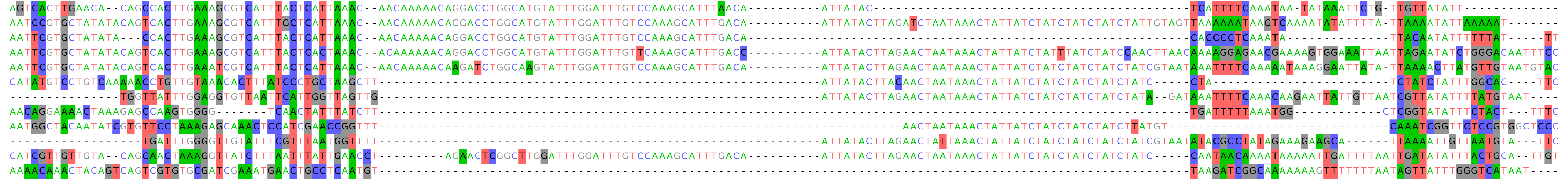
Start crop Point



End crop Point



MSA length = 1397



Start crop Point



End crop Point



0

10

0

250

500

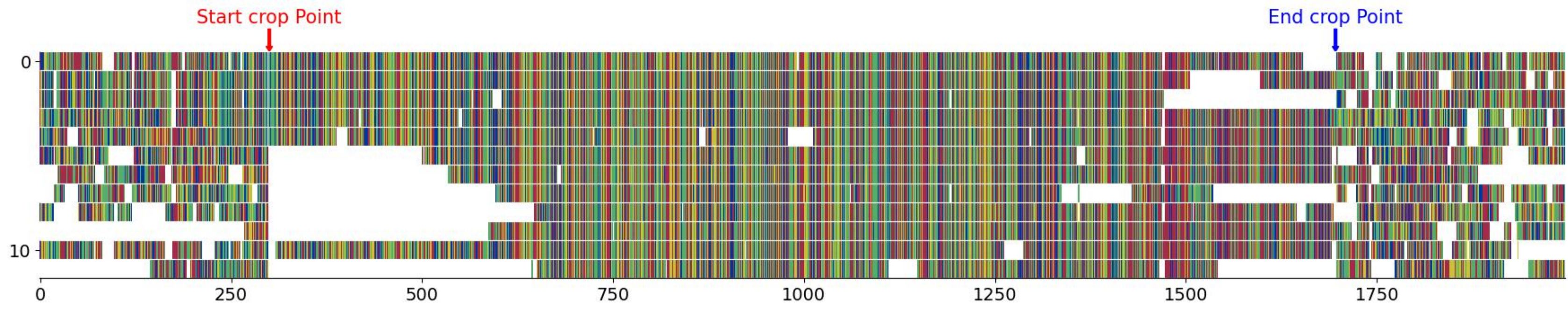
750

1000

1250

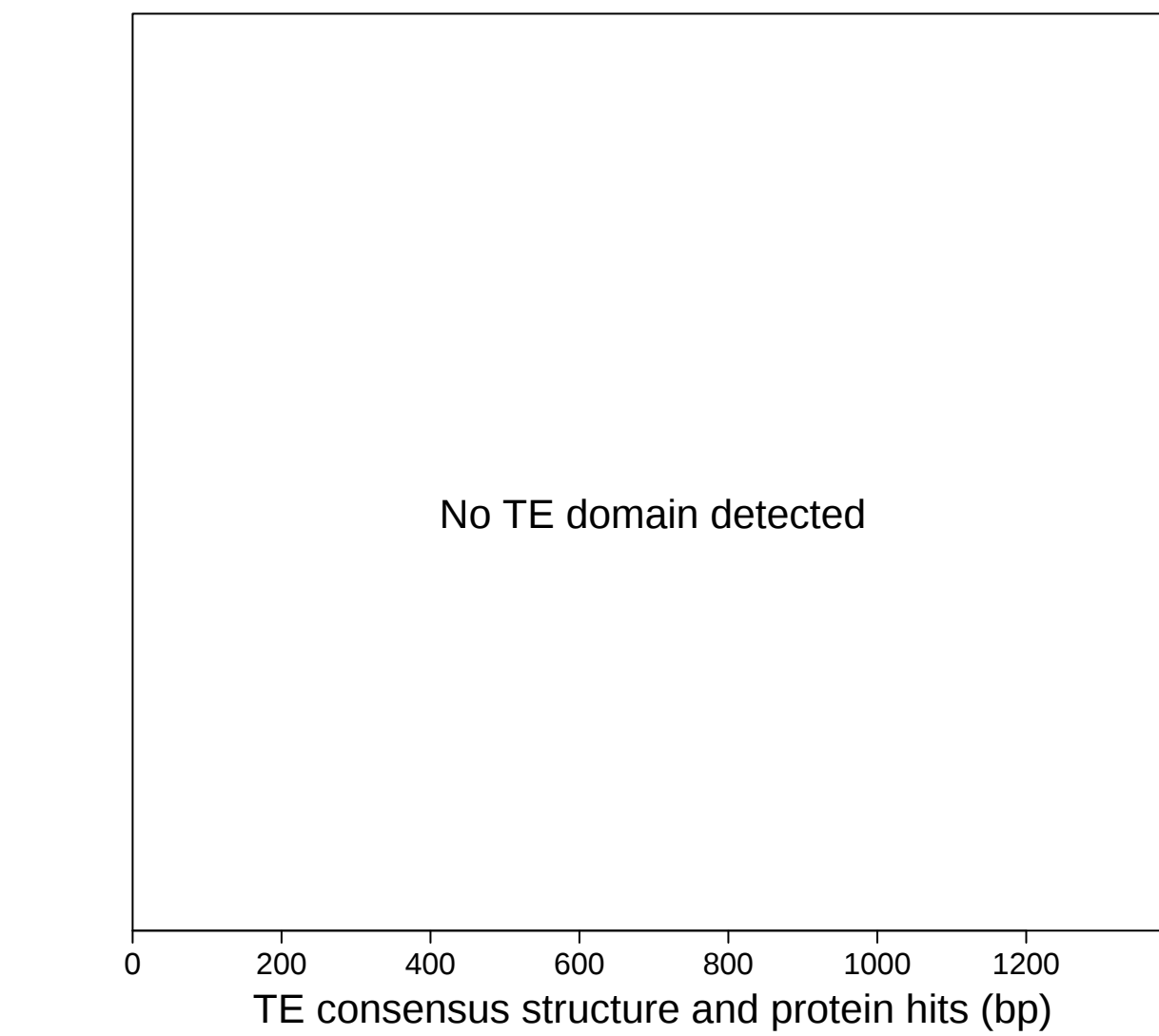
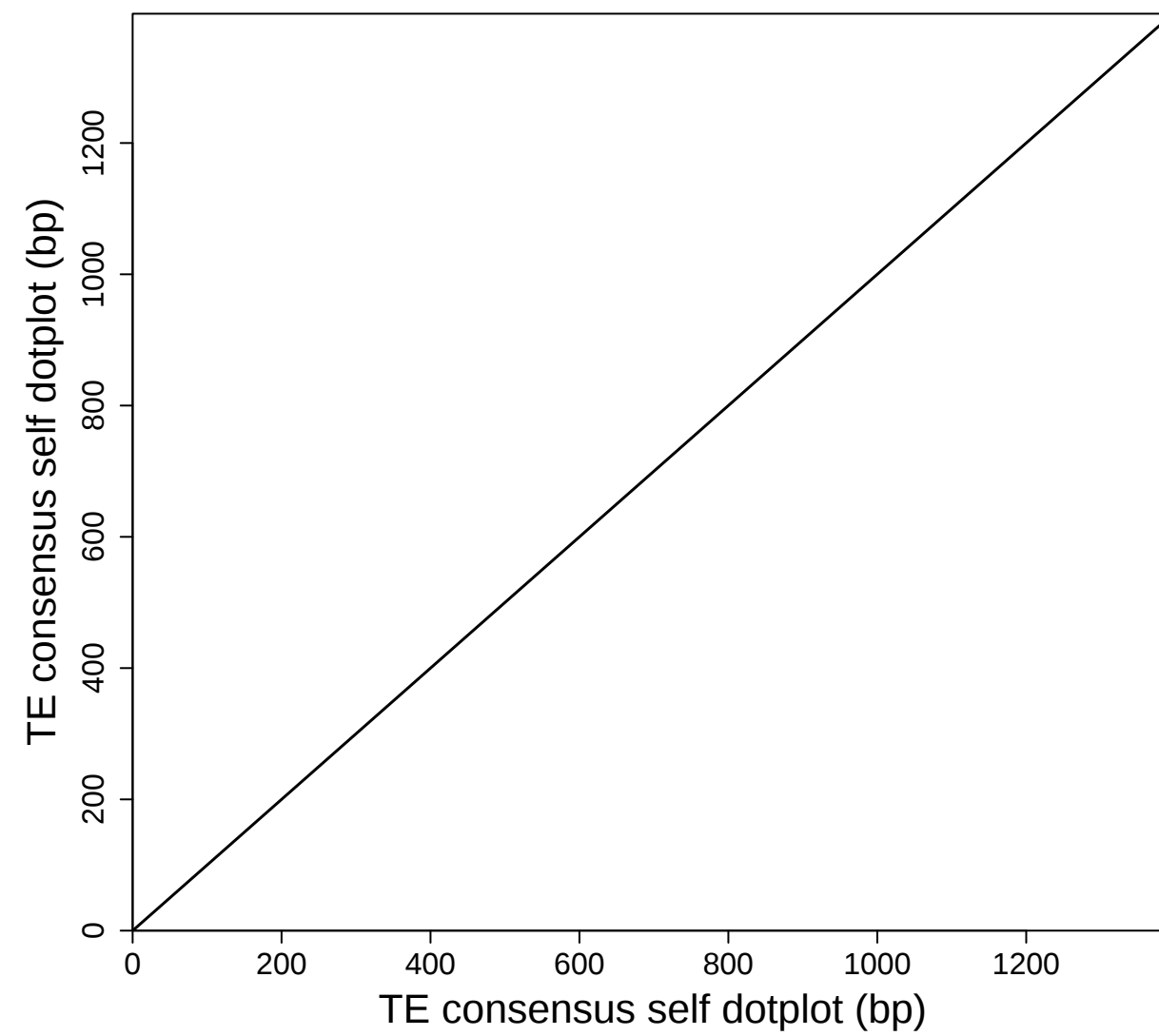
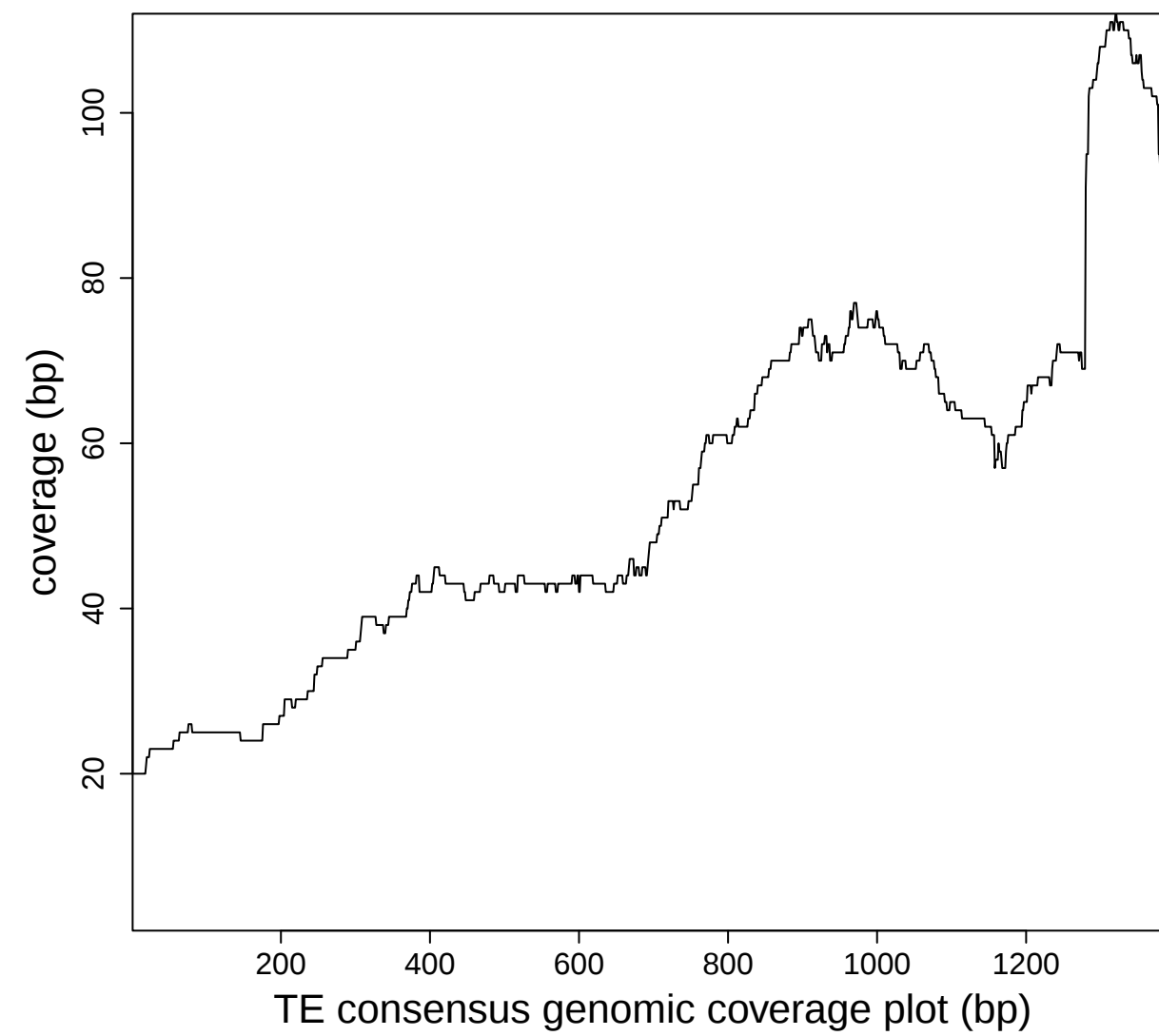
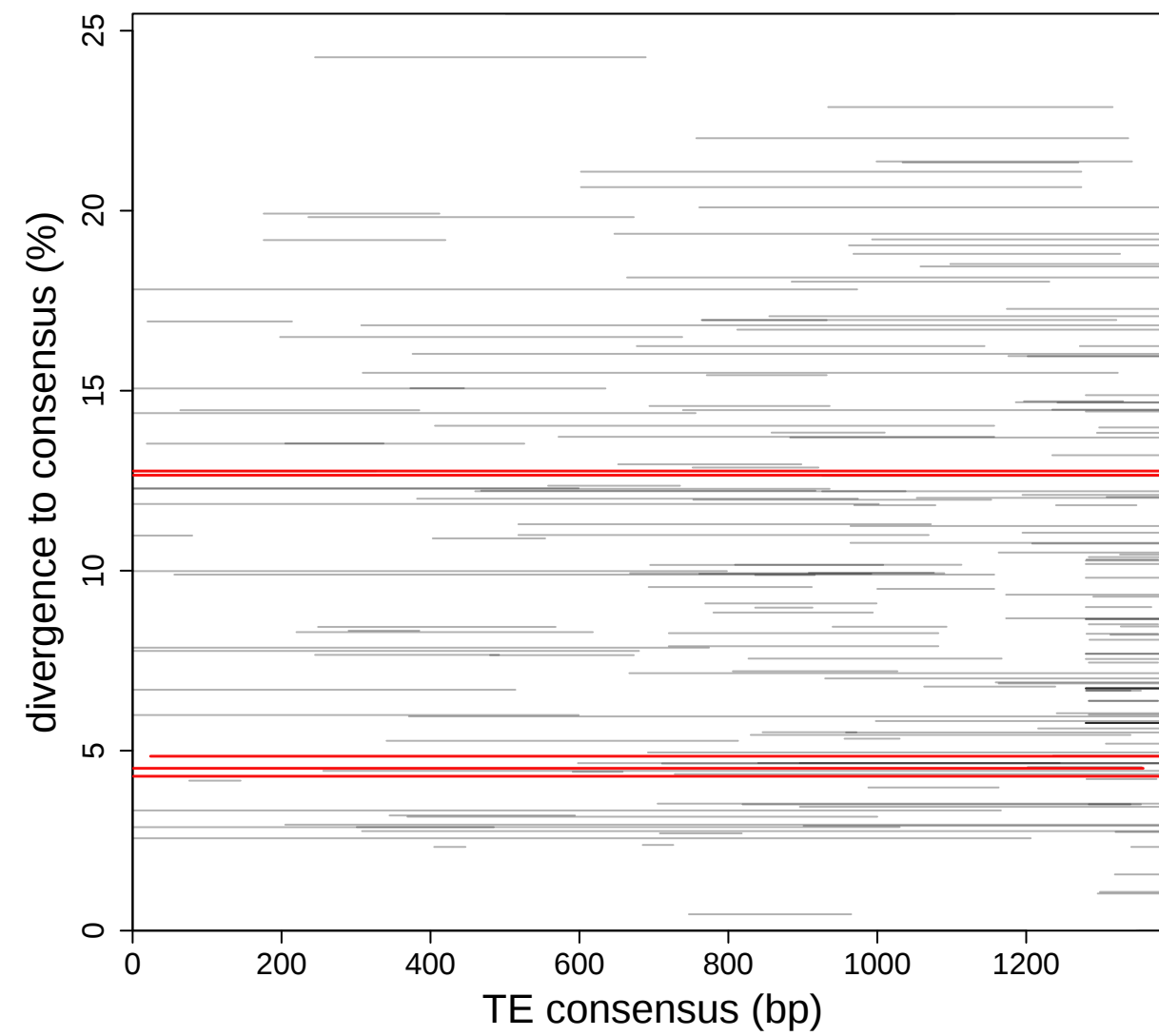
1500

1750



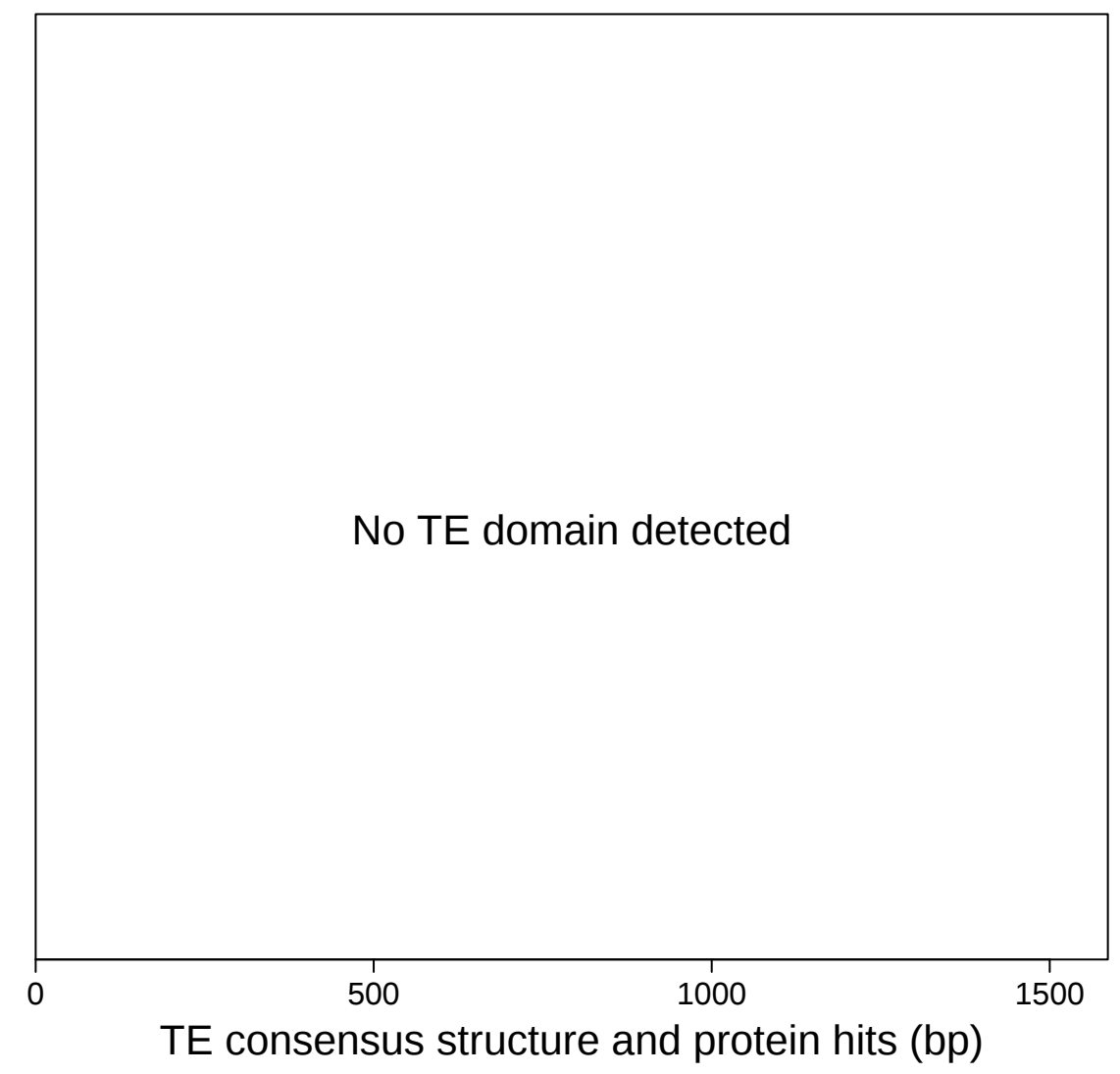
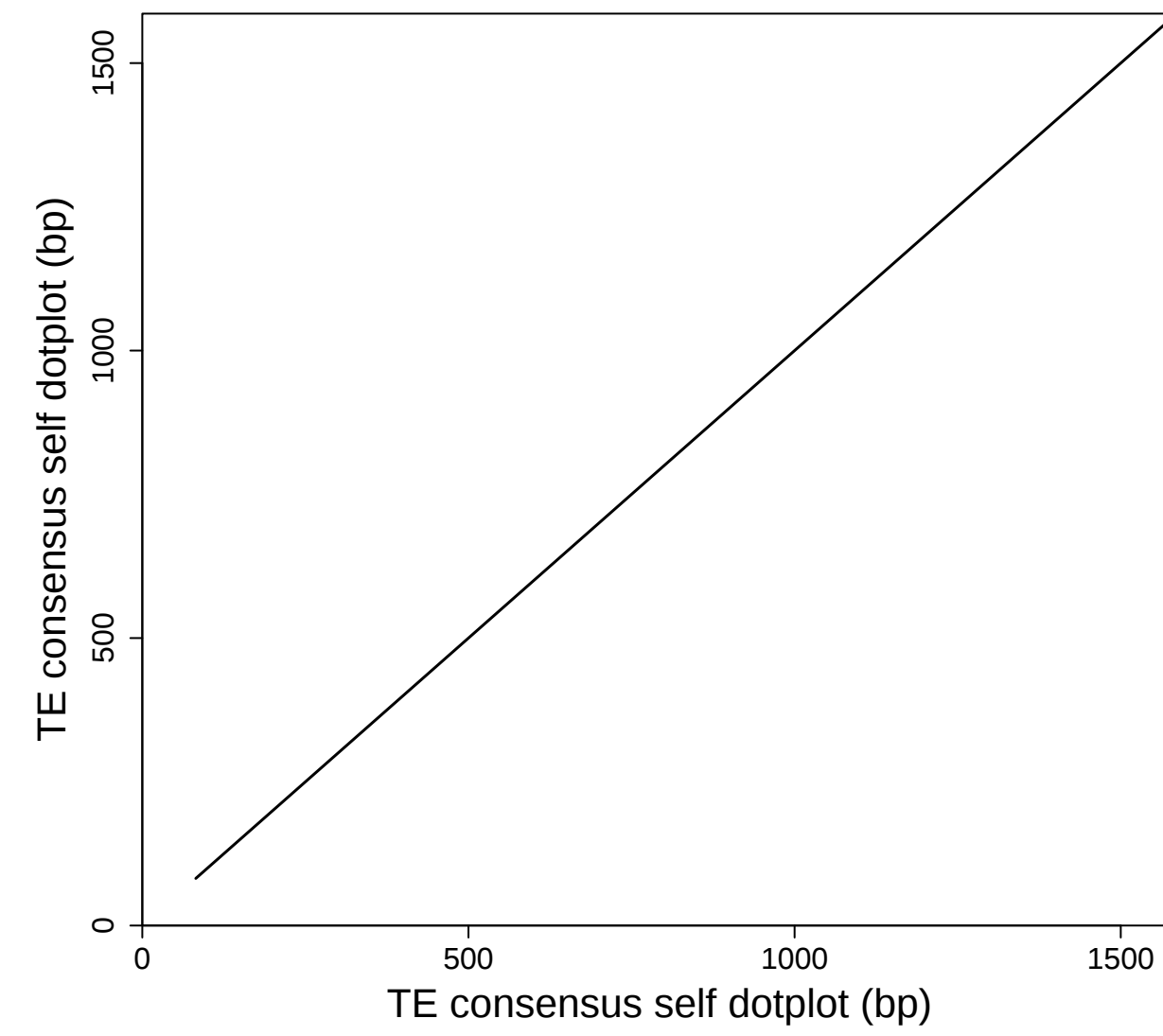
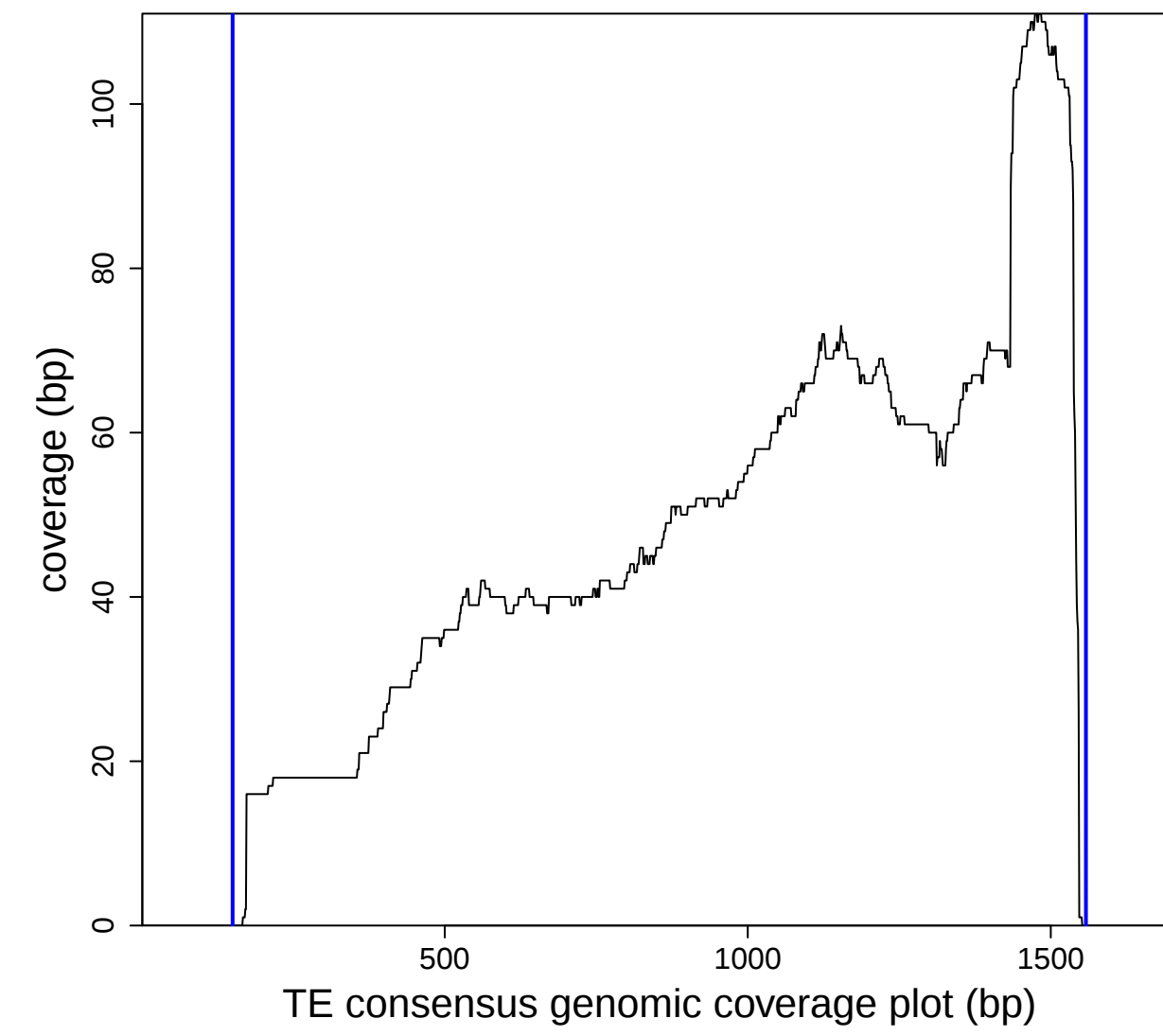
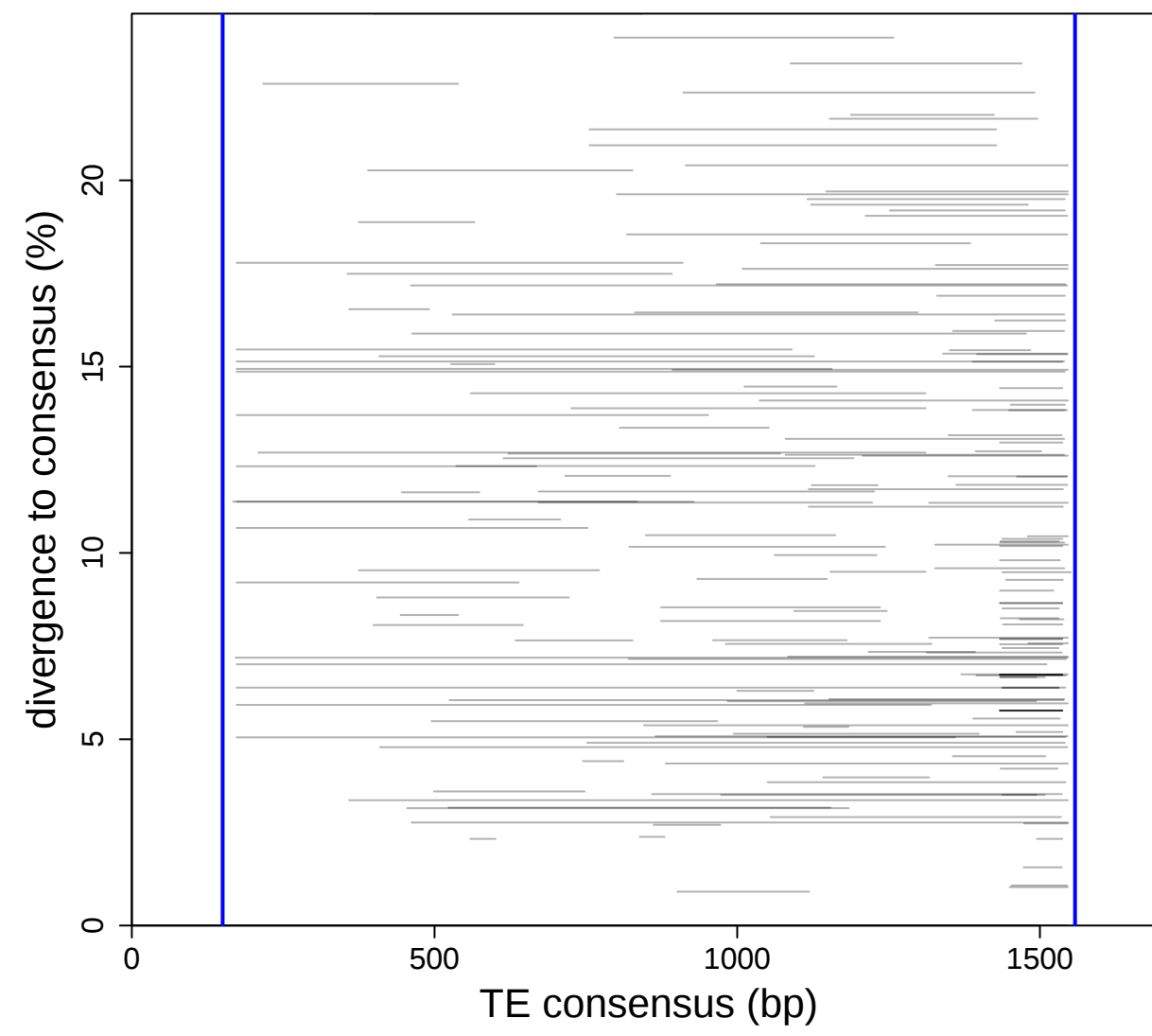
bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 1397bp; fragments: 204; full length: 5 (≥ 1257.3 bp)

After TEtrimmer 1397 bp

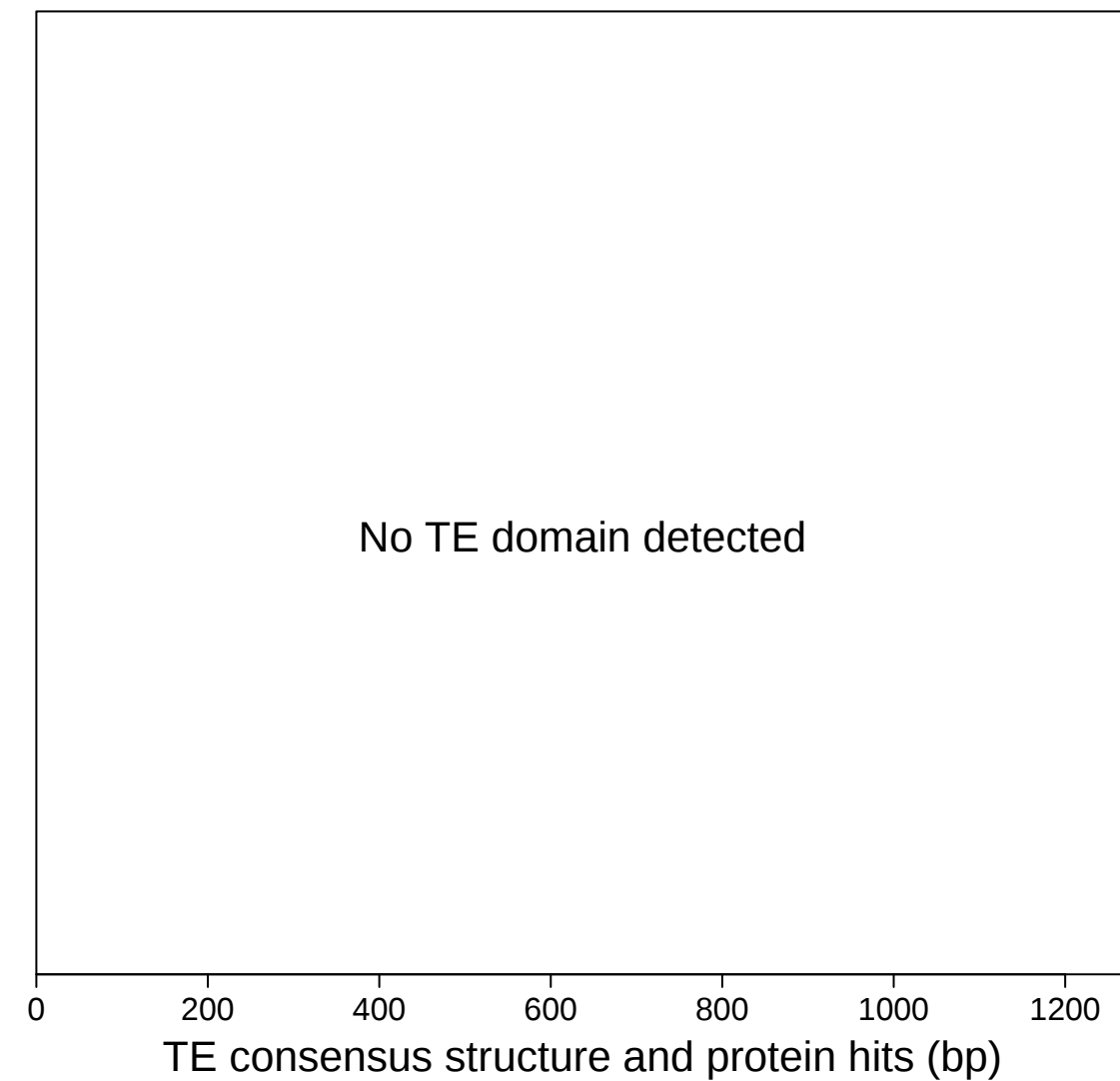
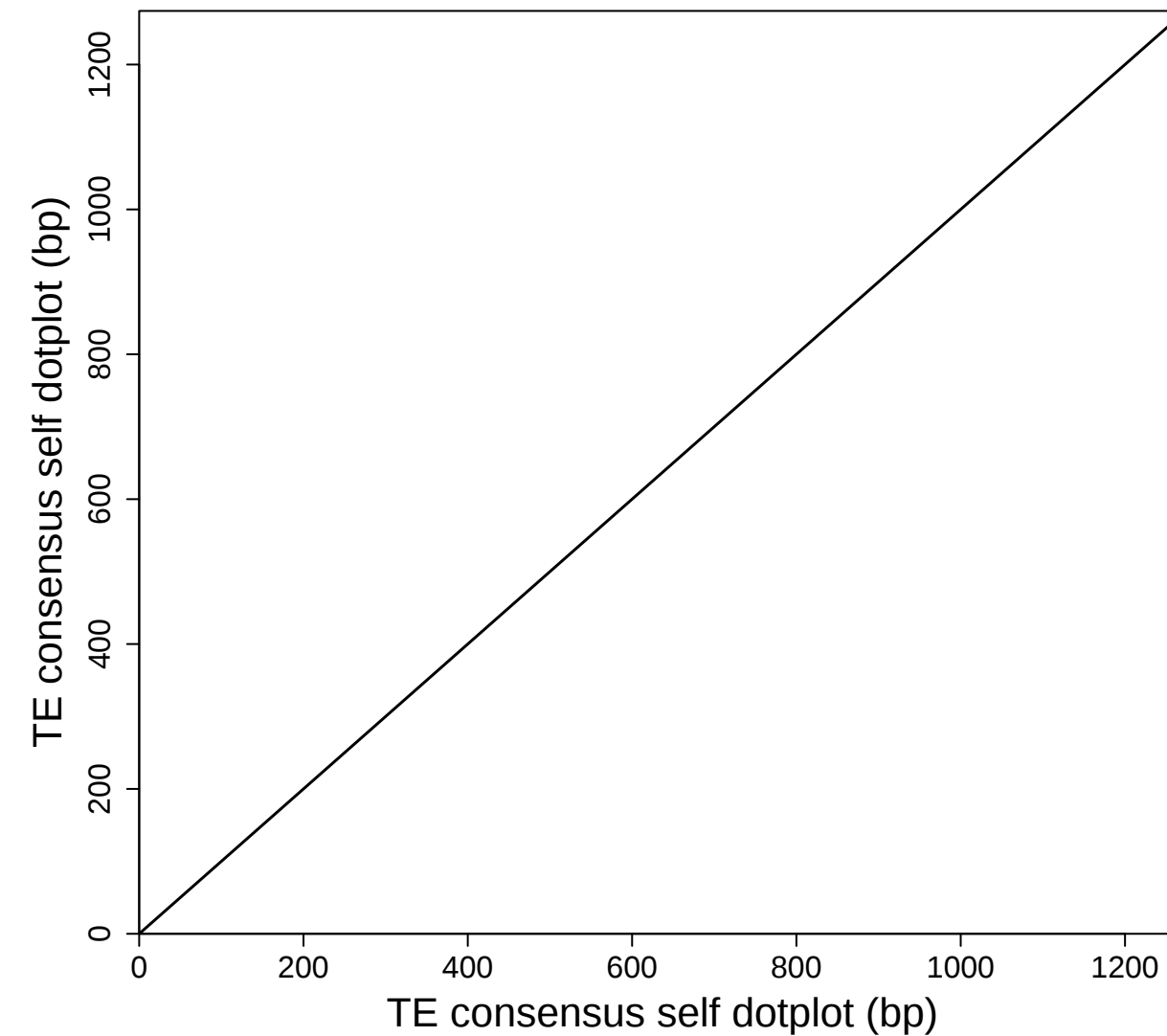
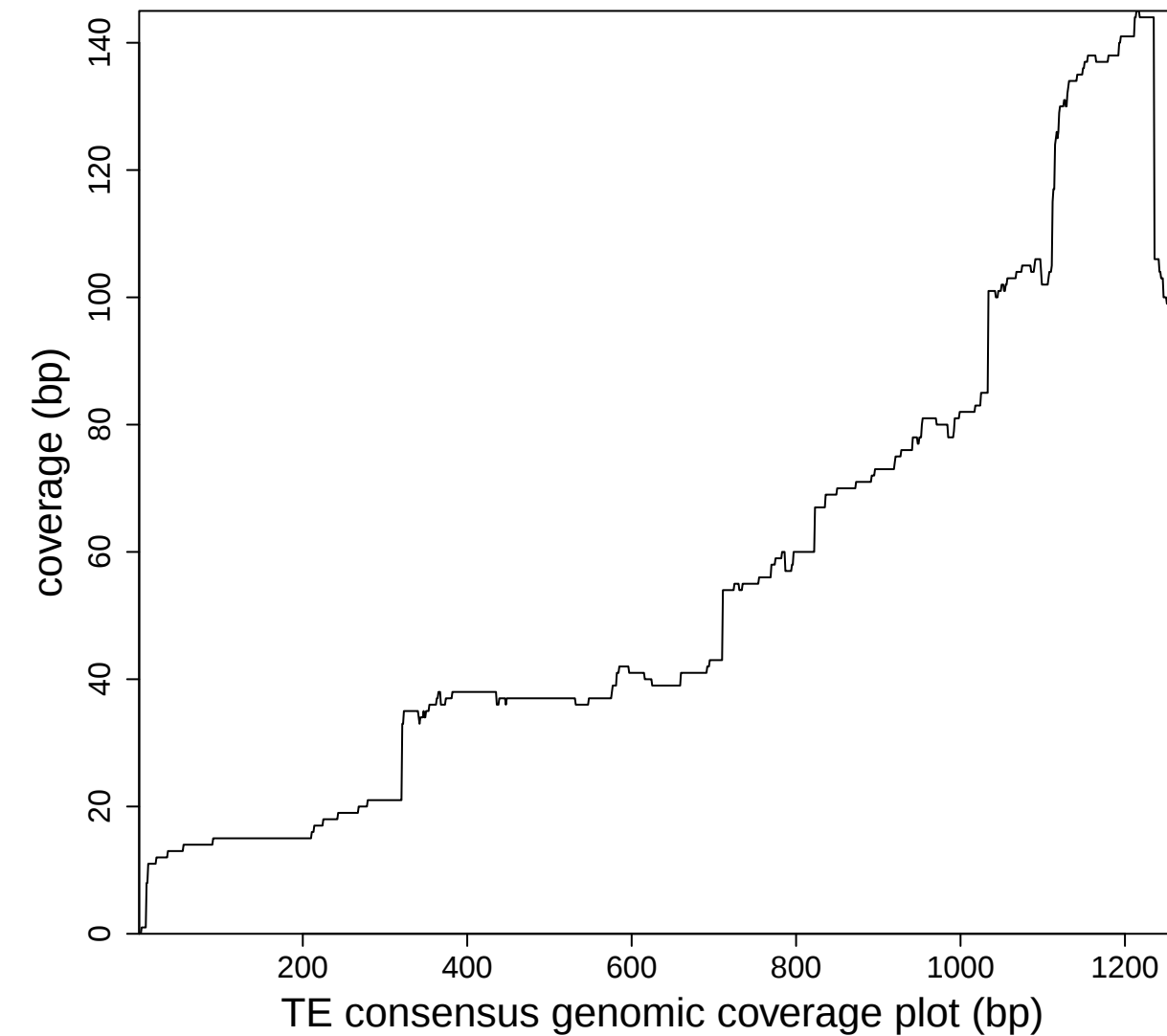
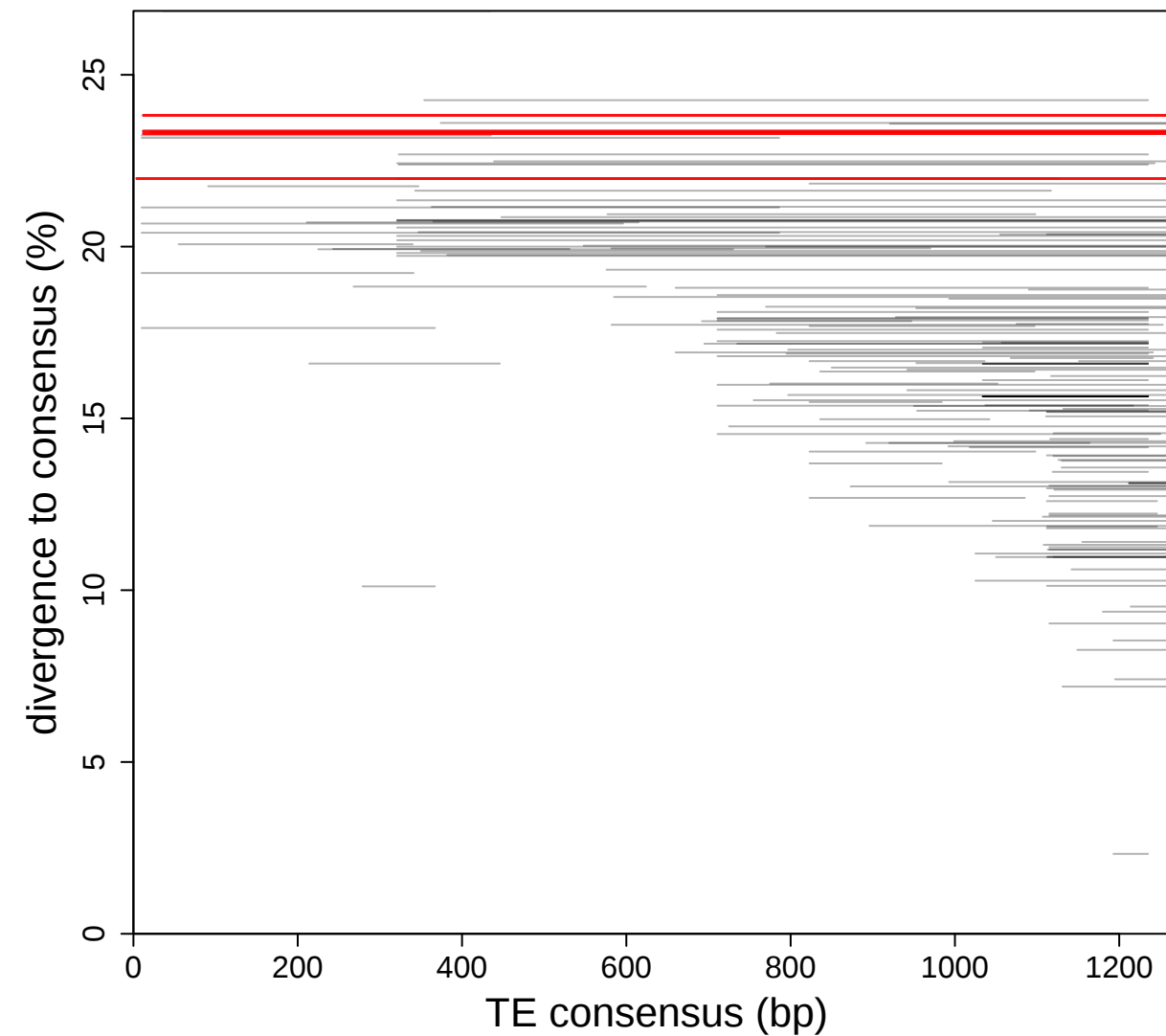


2.fasta.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1708bp; fragments: 185; full length: 0 (>=1537.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

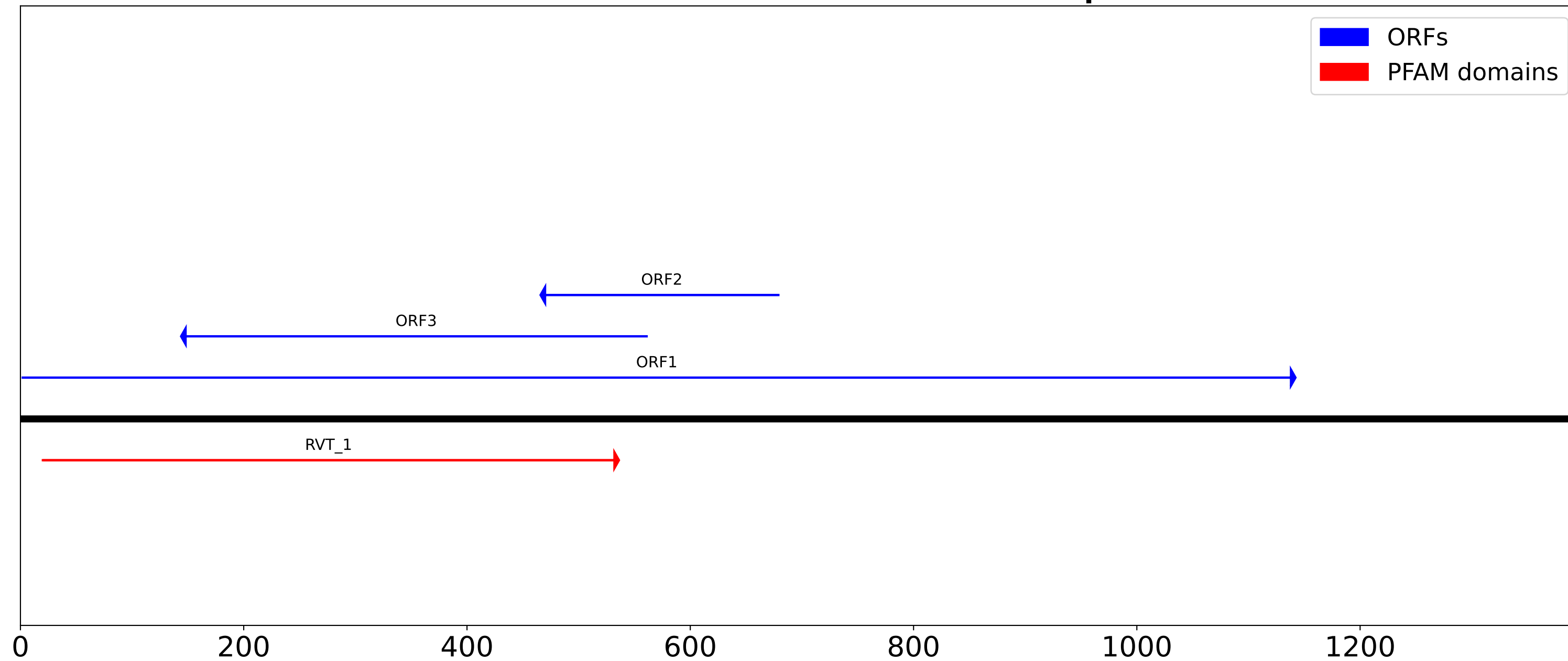


TE: rnd_3_family_682
size: 1274bp; fragments: 176; full length: 5 (≥ 1146.6 bp)

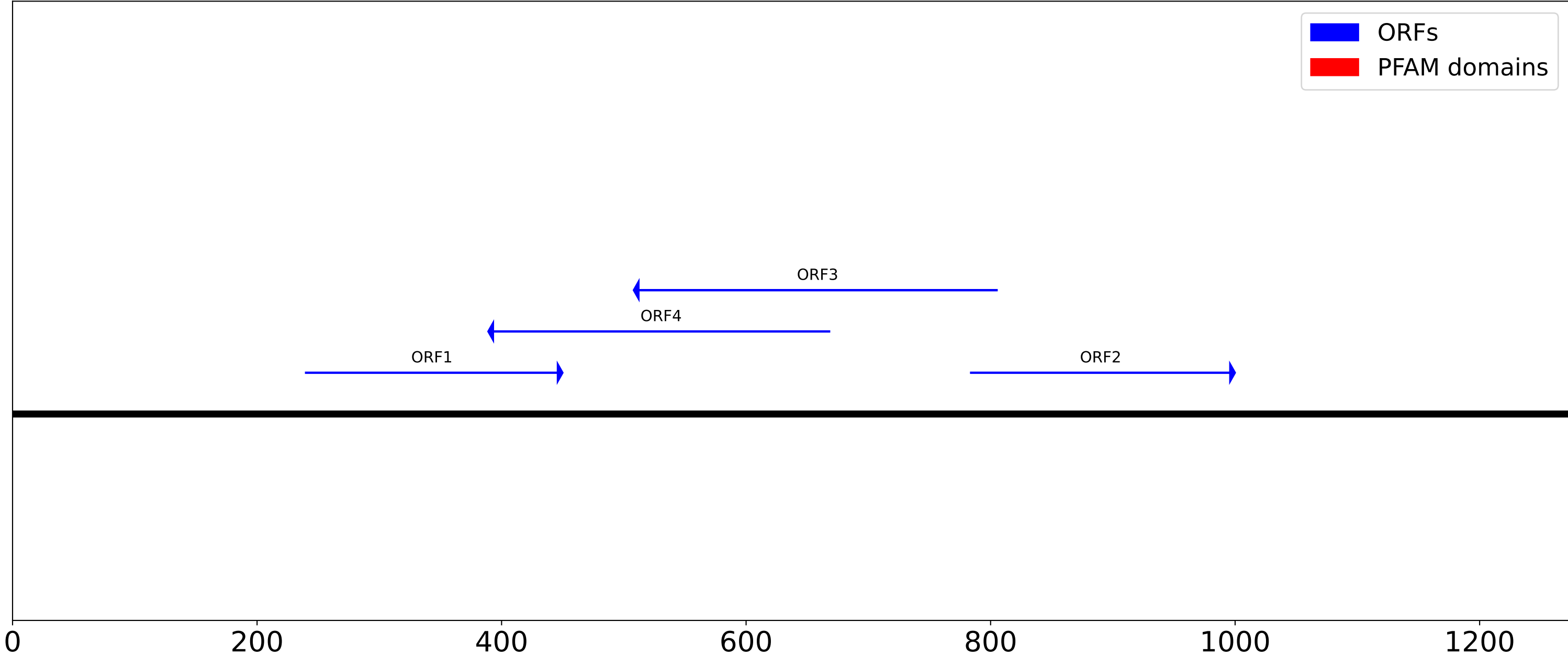


Before TEtrimmer 1274 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

