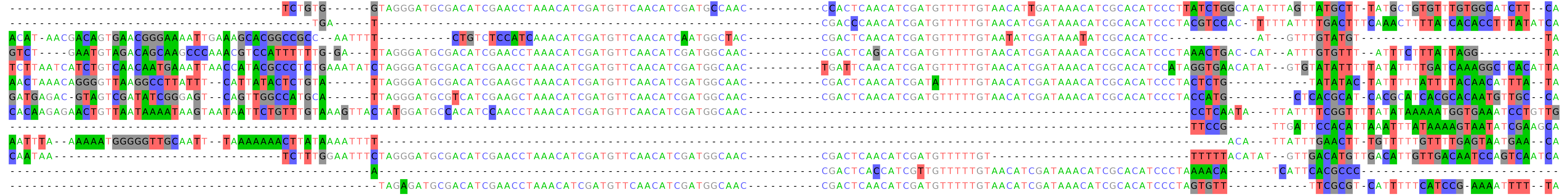


Start crop Point

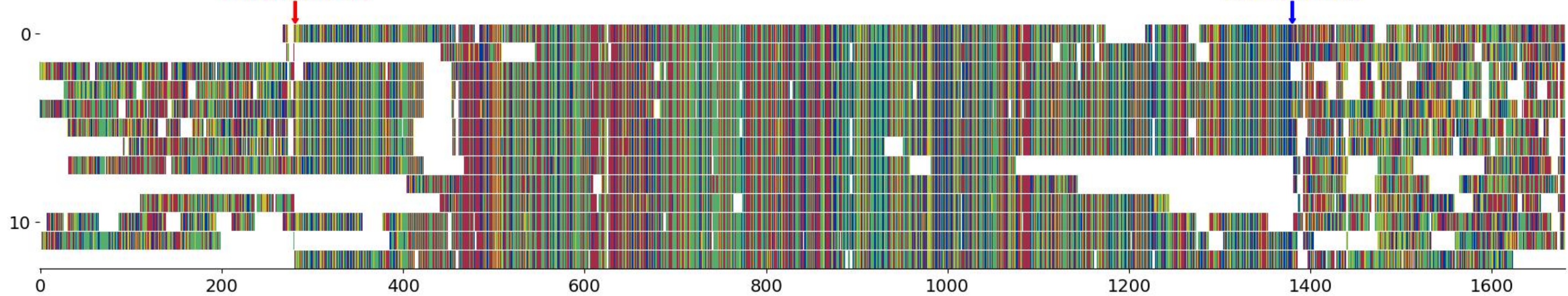
End crop Point

MSA length = 1100



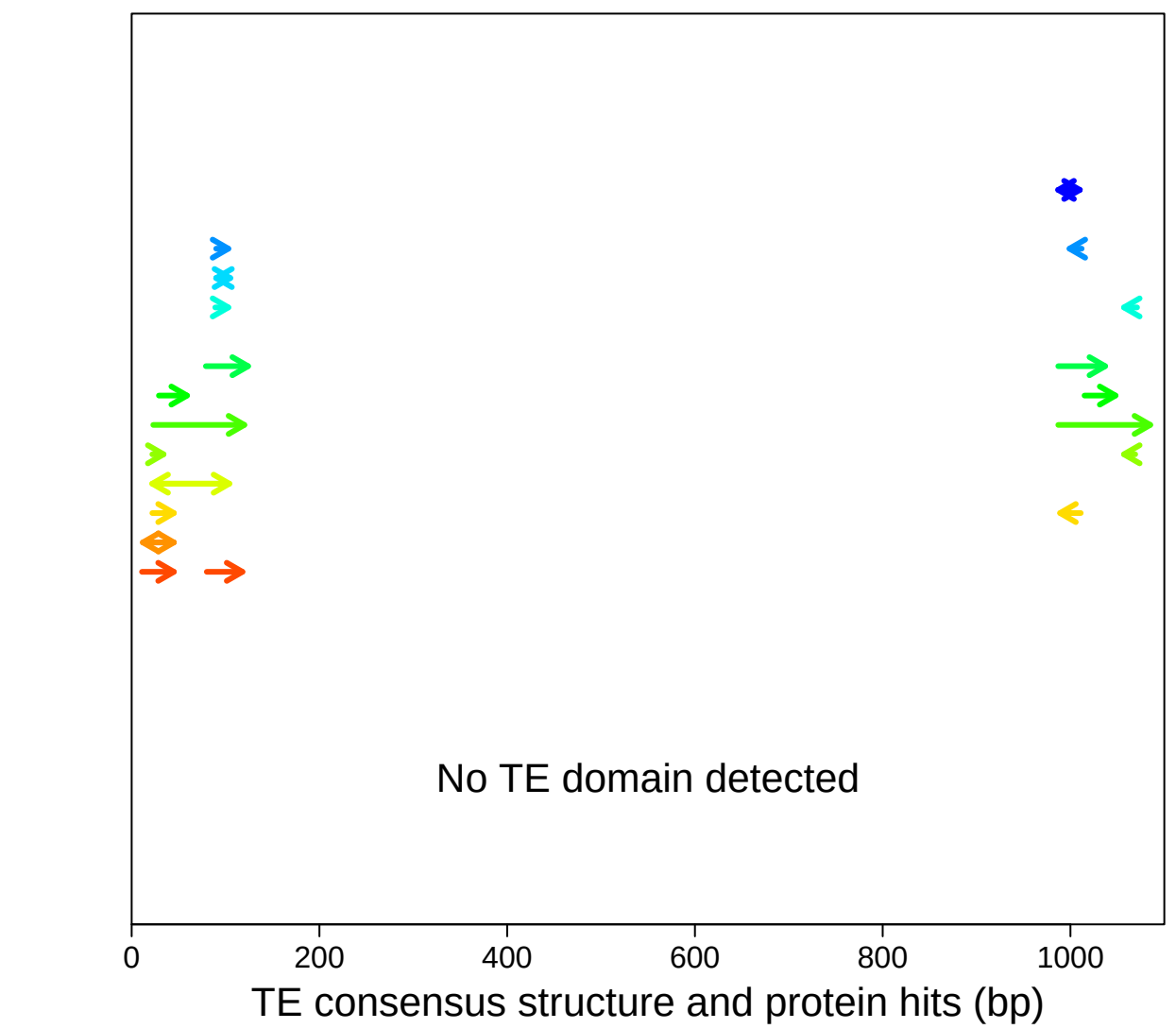
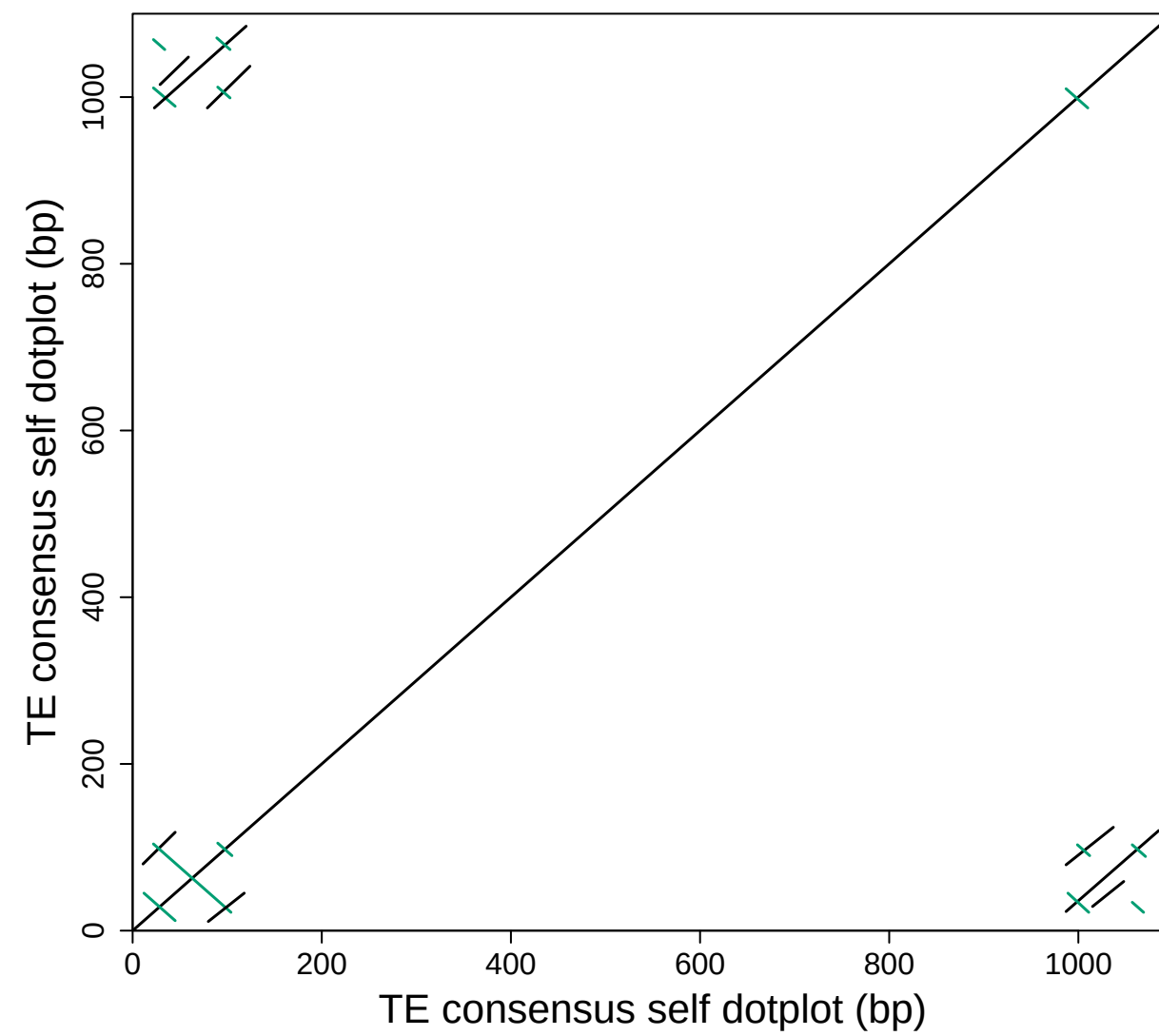
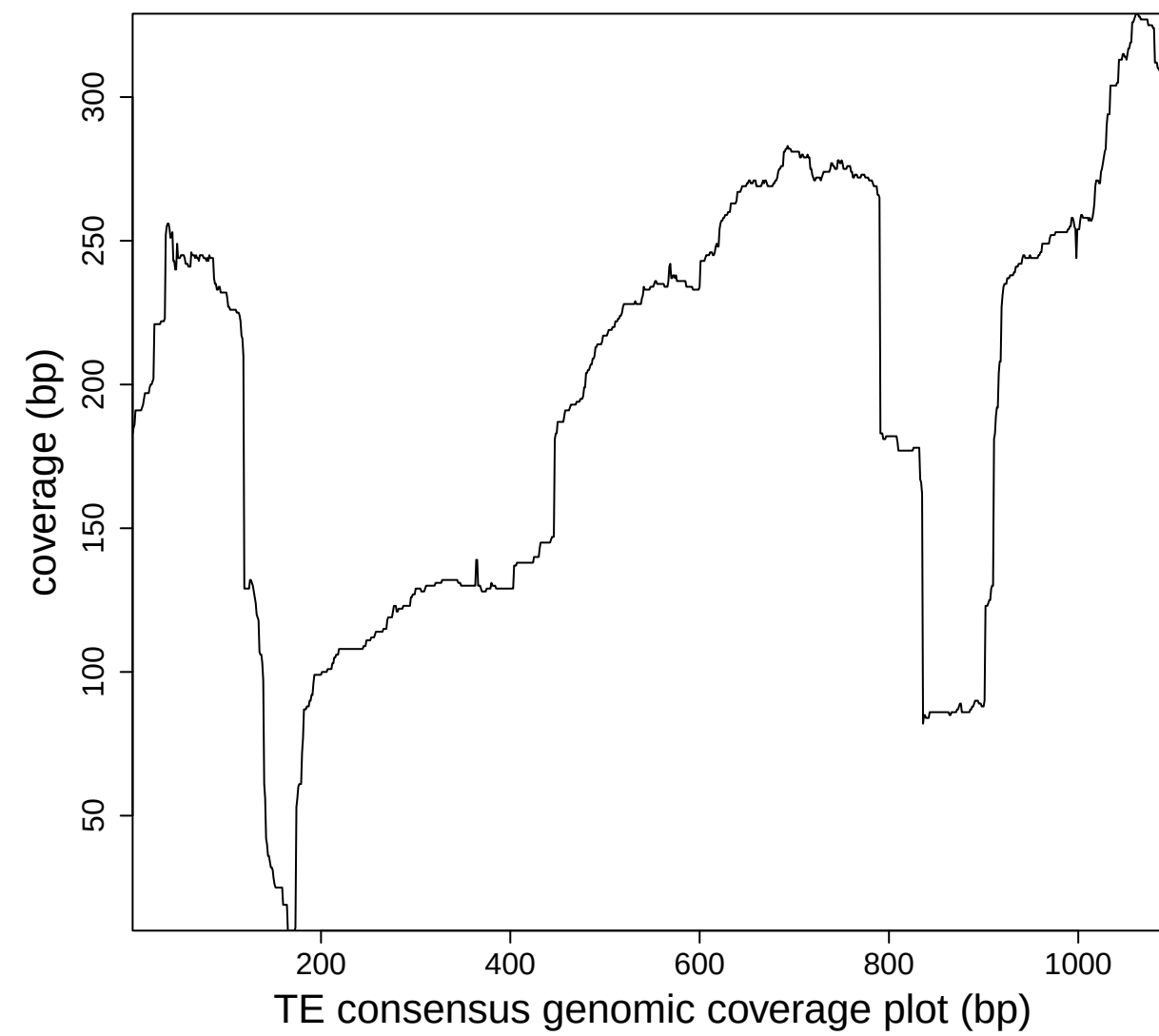
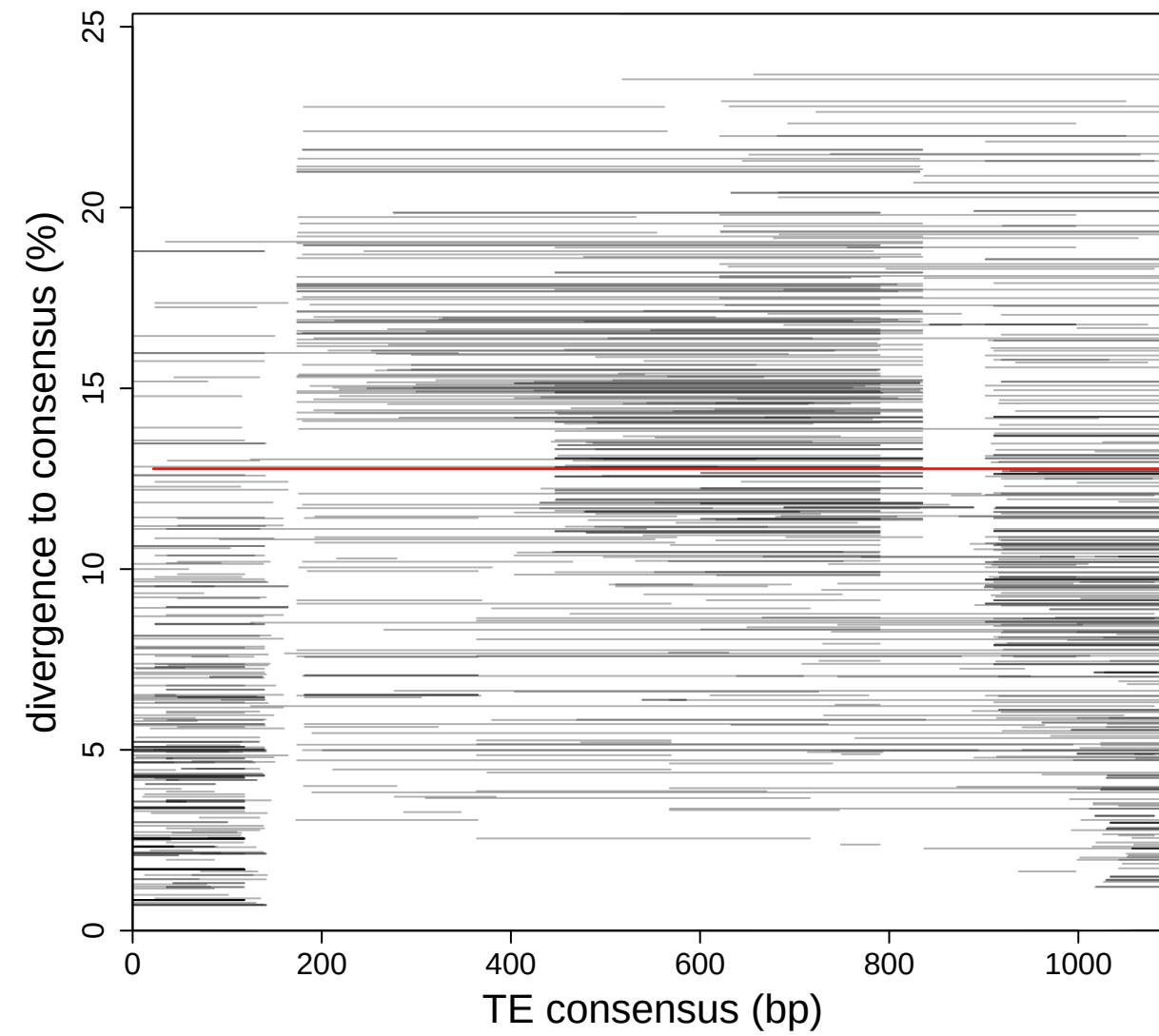
Start crop Point

End crop Point



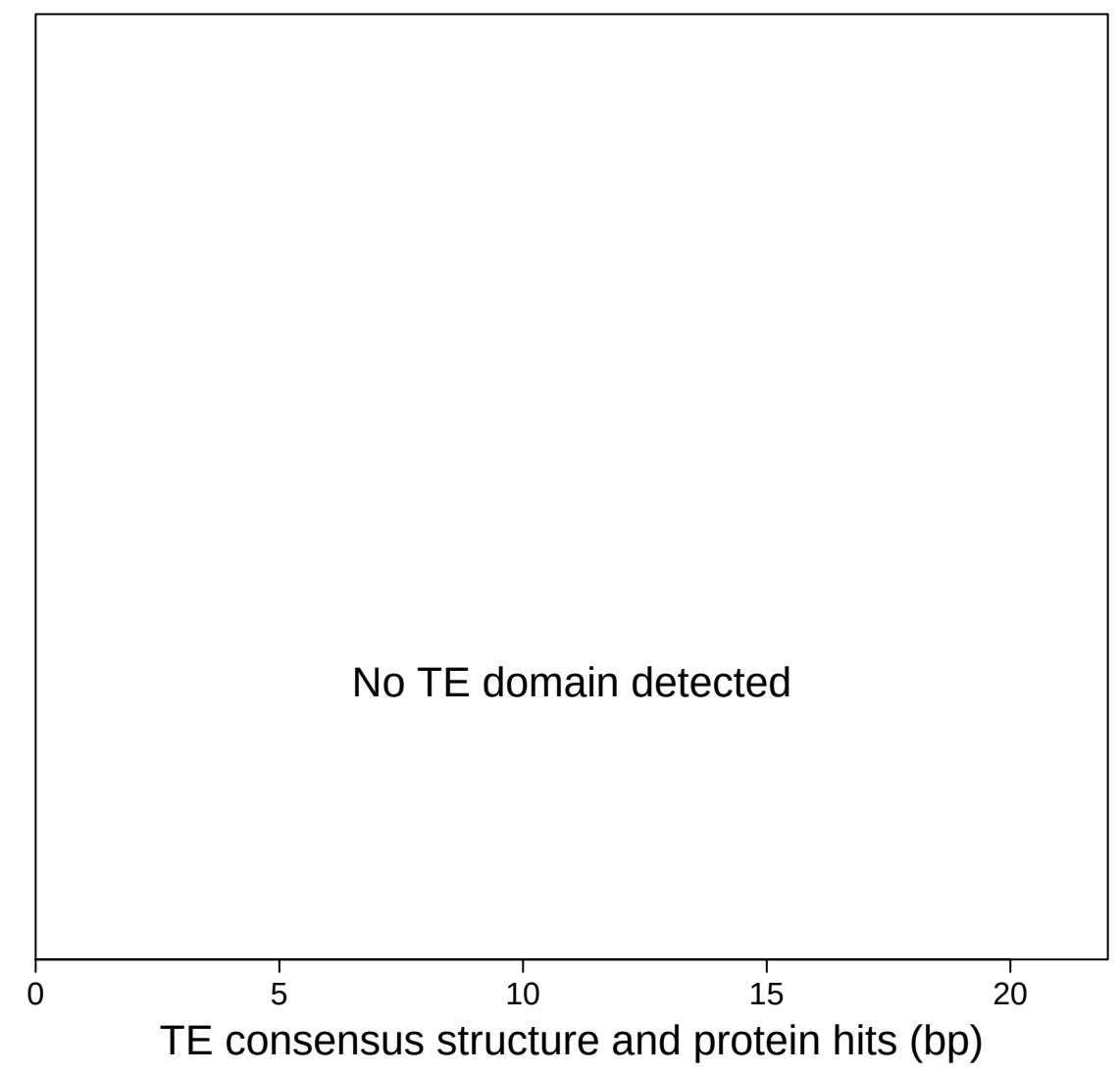
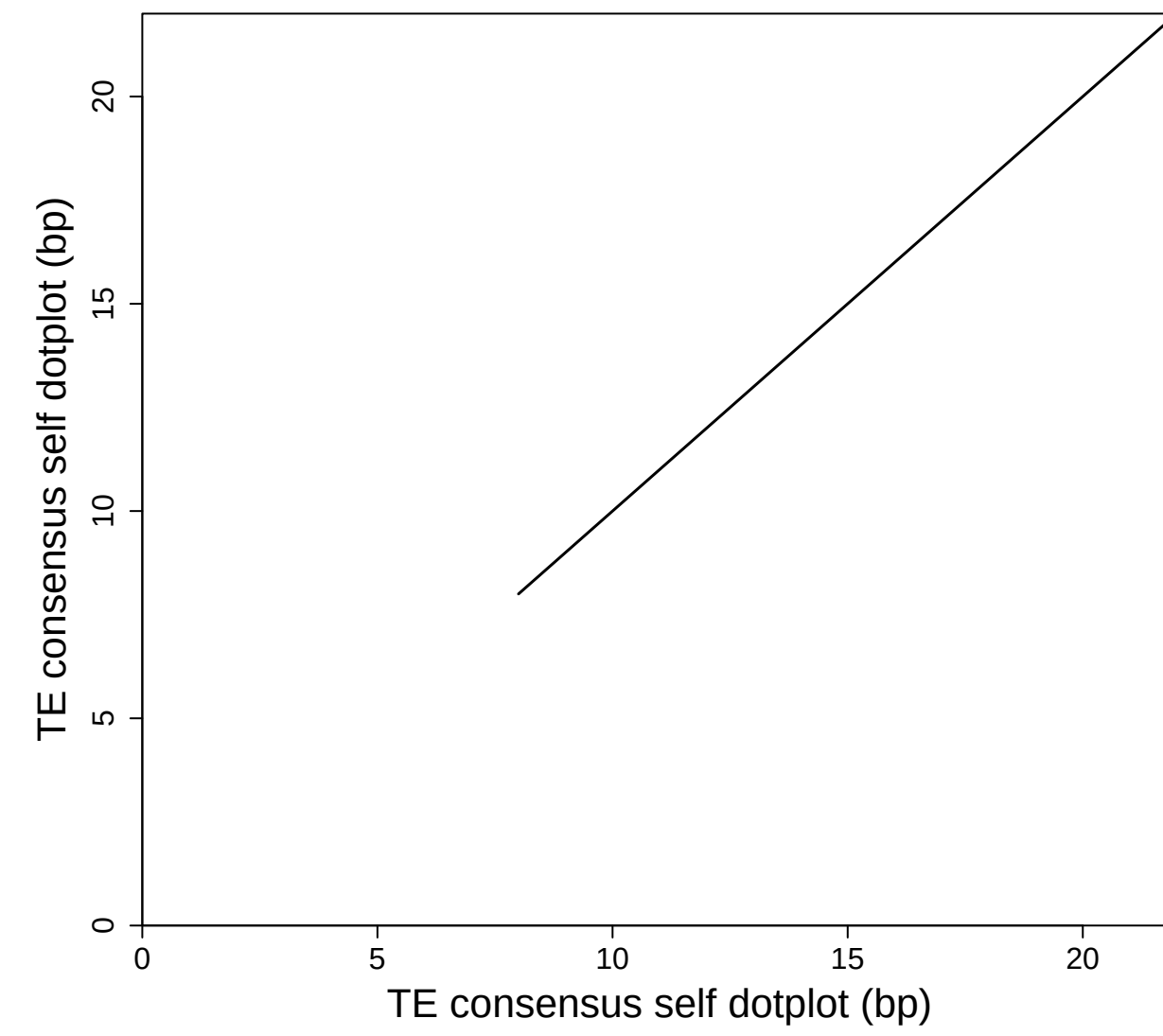
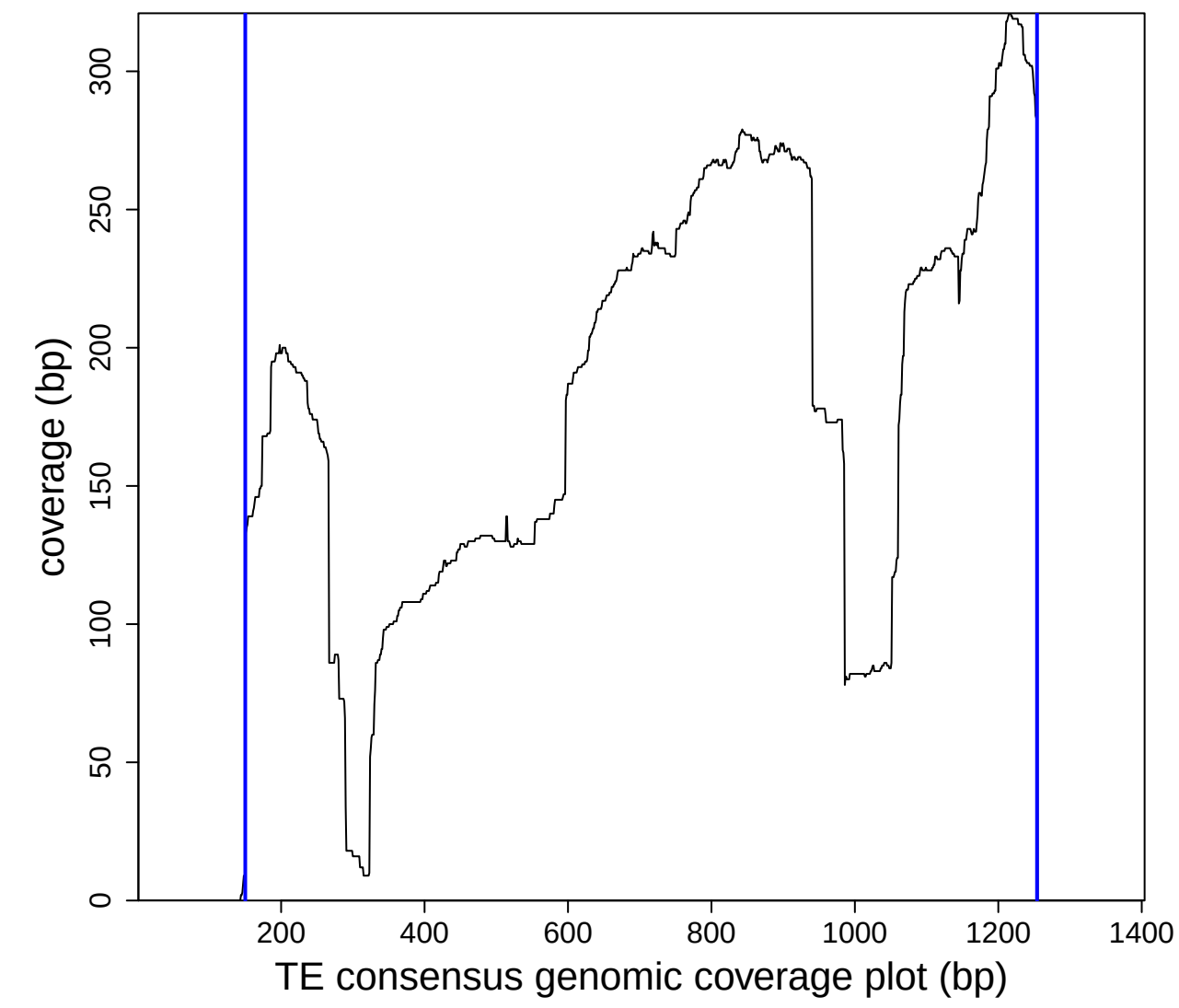
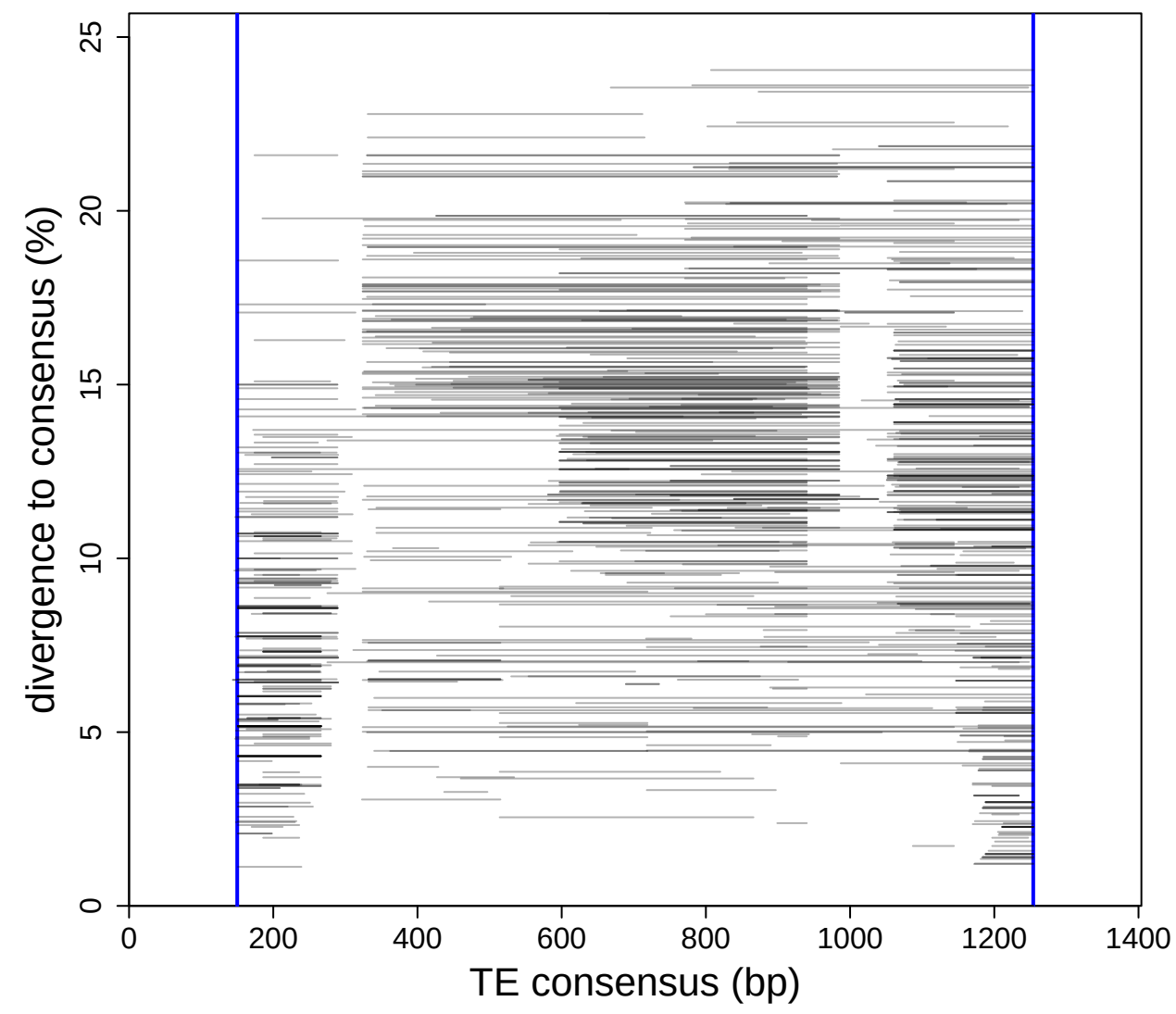
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1100bp; fragments: 917; full length: 1 (>=990bp)

After TETrimmer 1100 bp

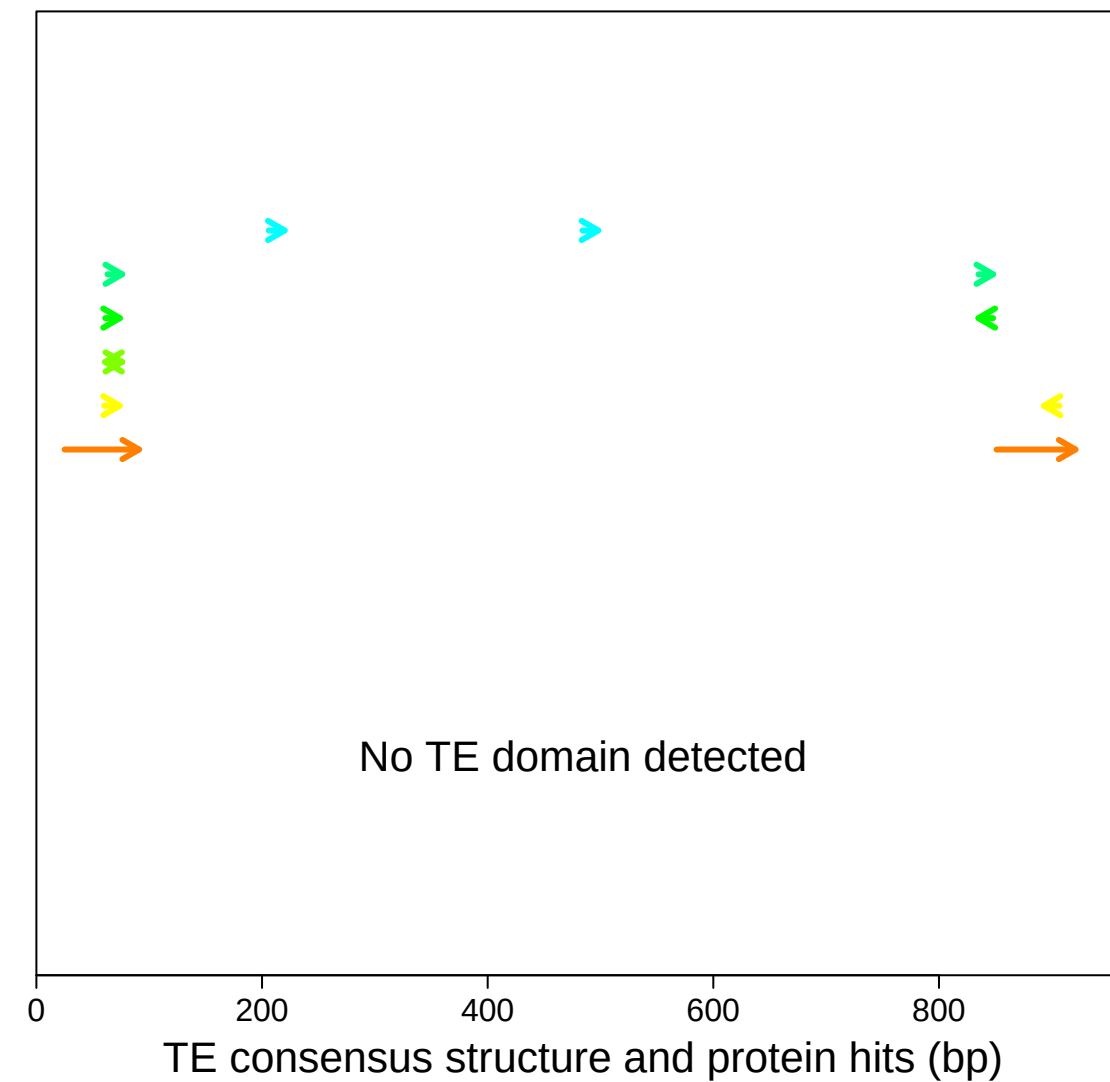
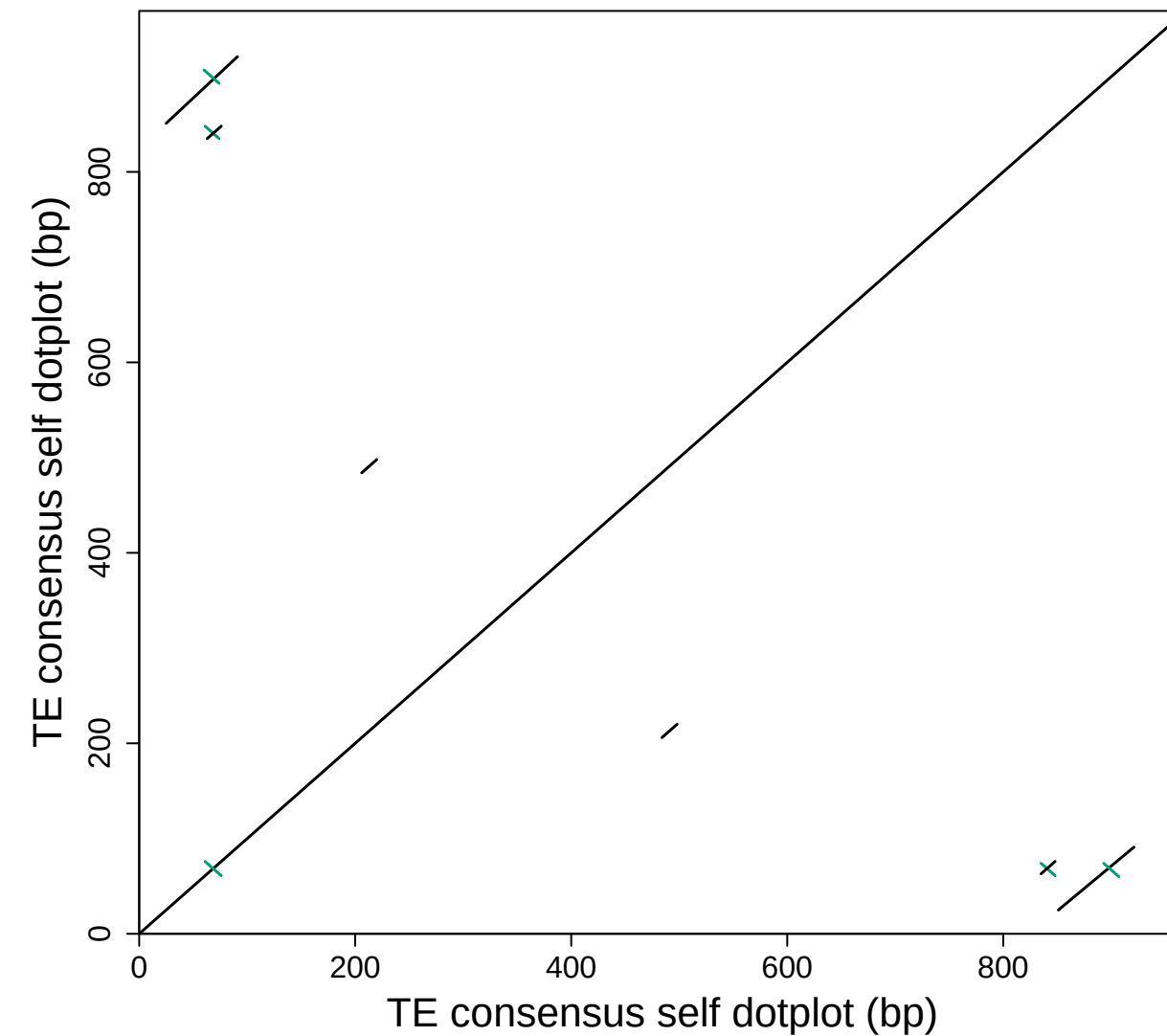
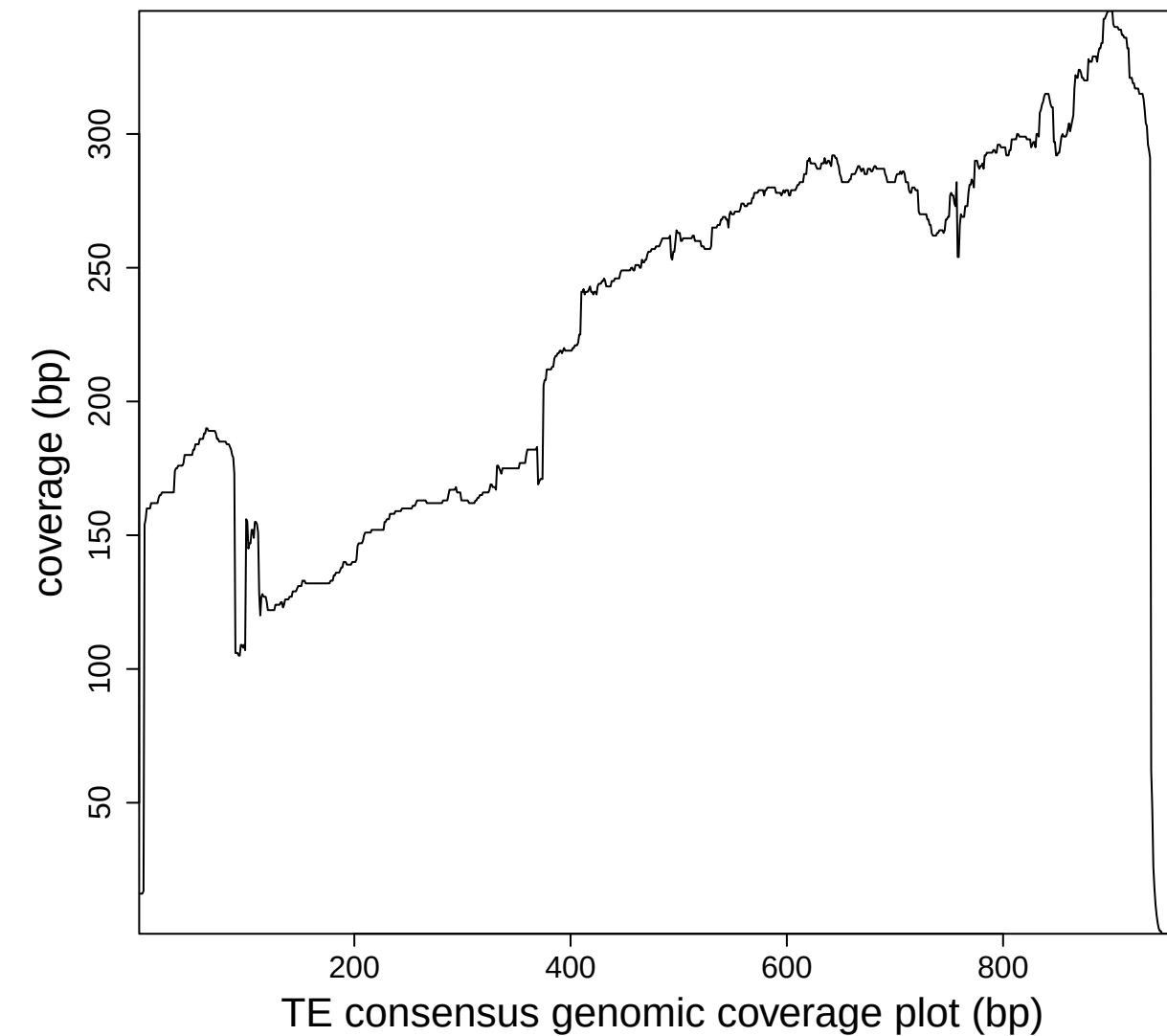
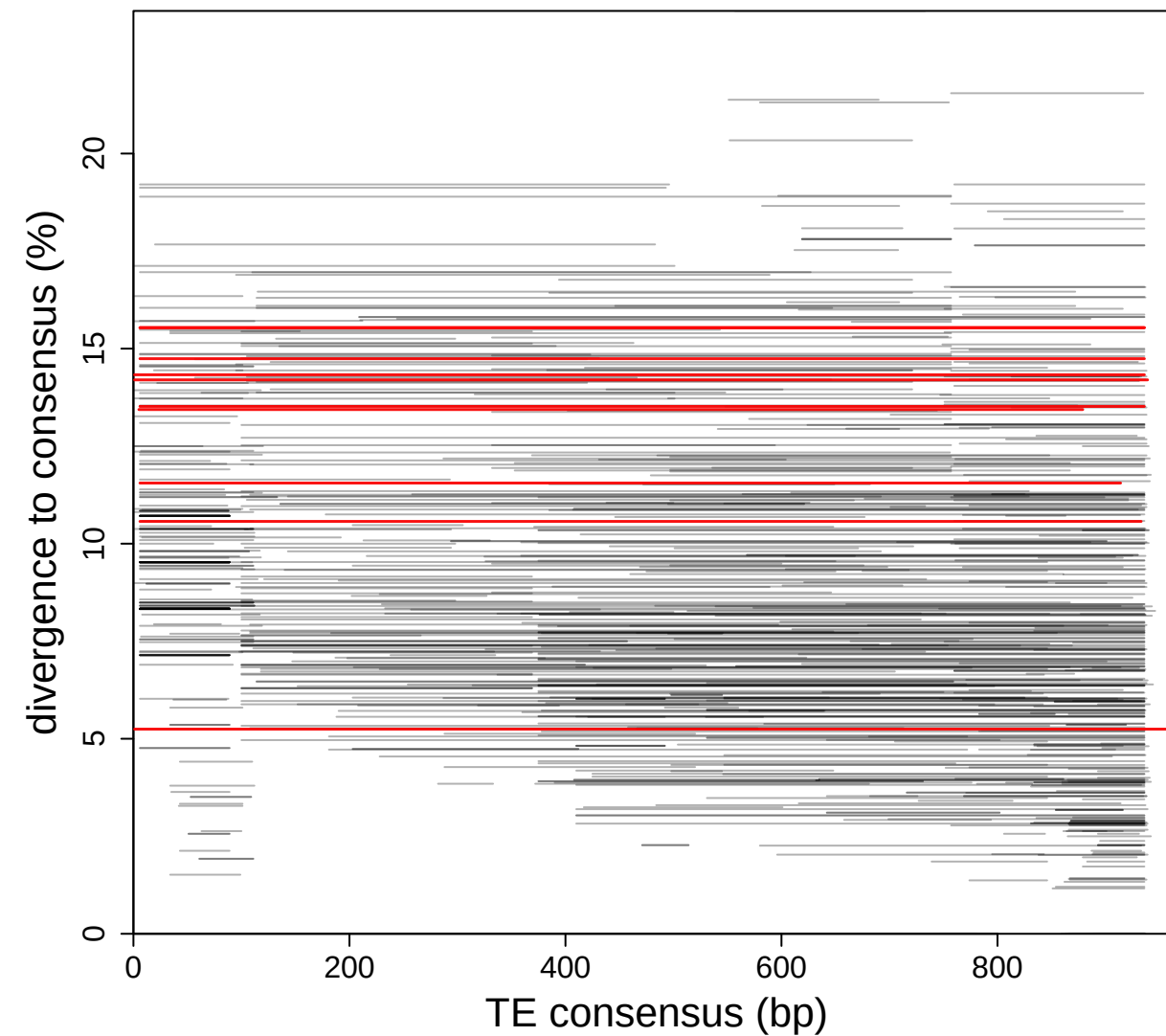


3.fasta.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 1404bp; fragments: 828; full length: 0 (>=1263.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

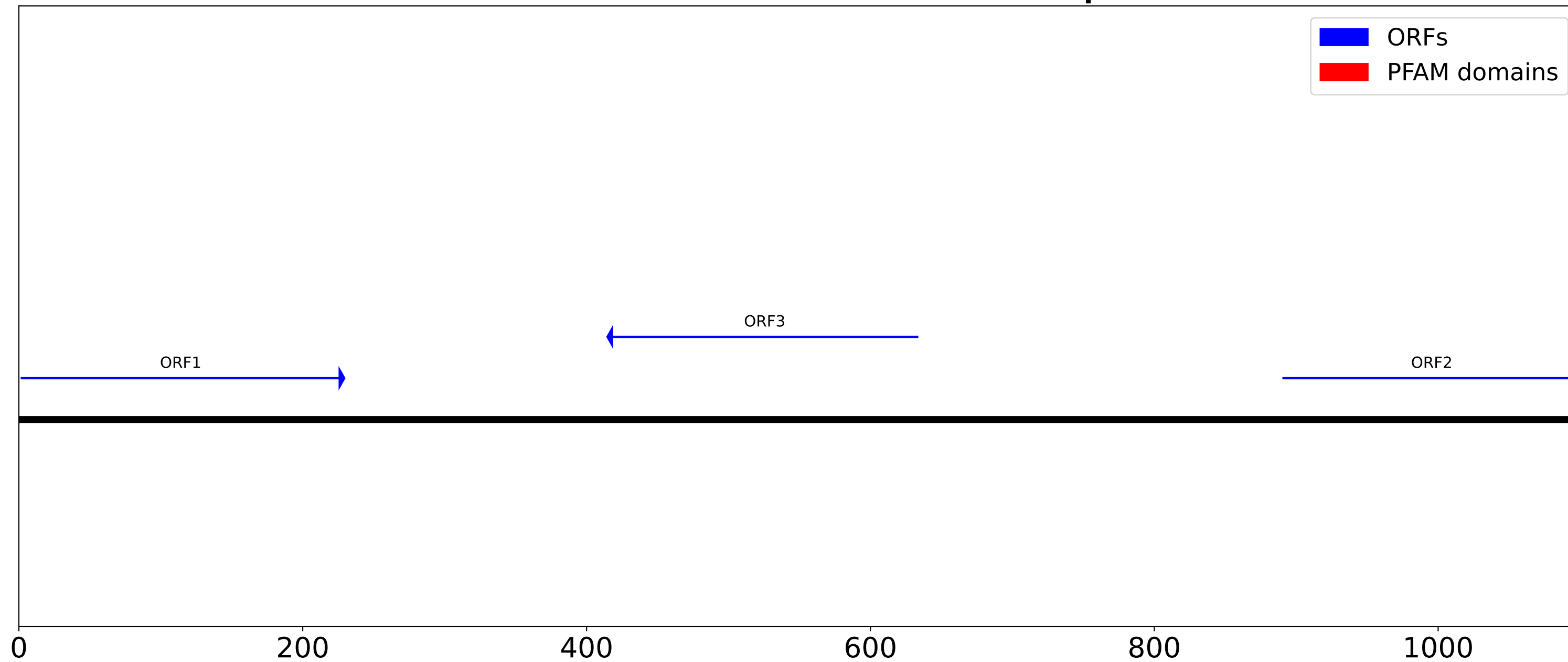


TE: rnd_4_family_2263
size: 969bp; fragments: 763; full length: 10 (≥ 872.1 bp)

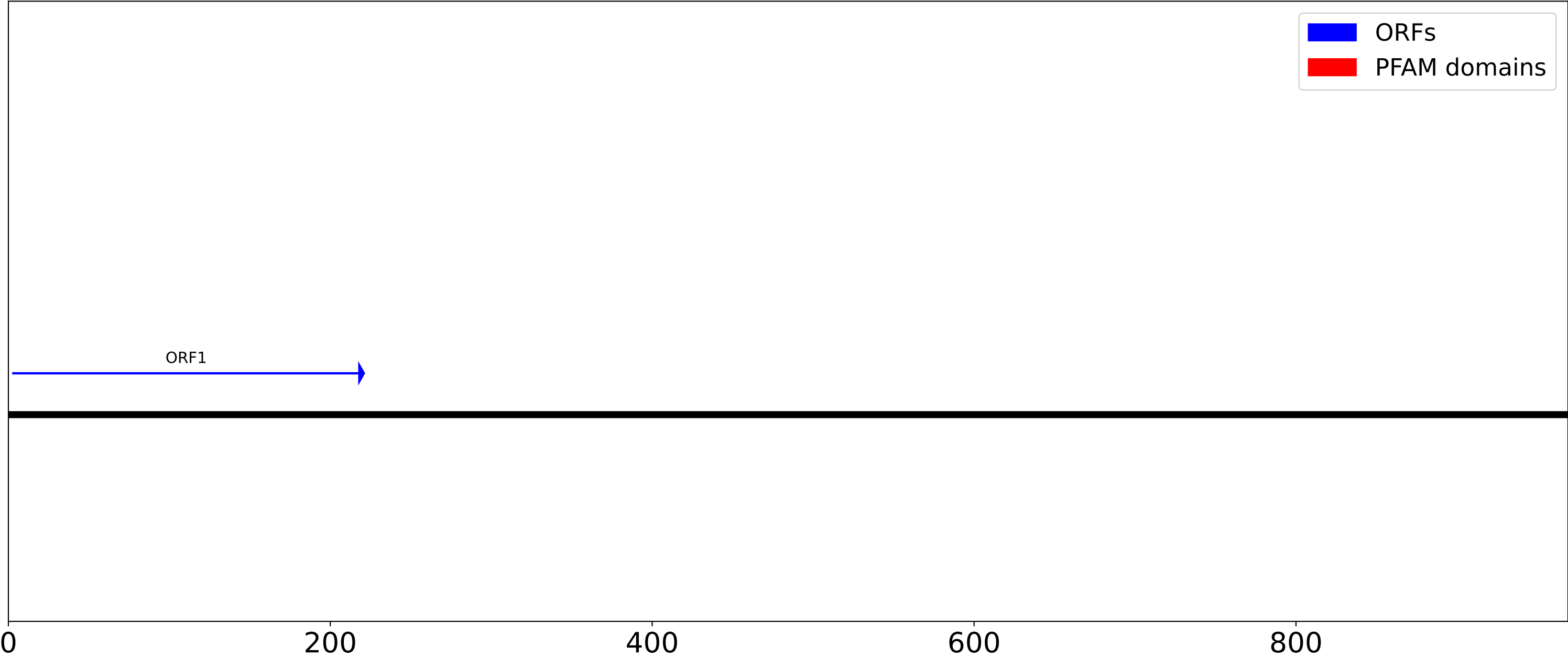


Before TEtrimmer 969 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

