

Start crop Point

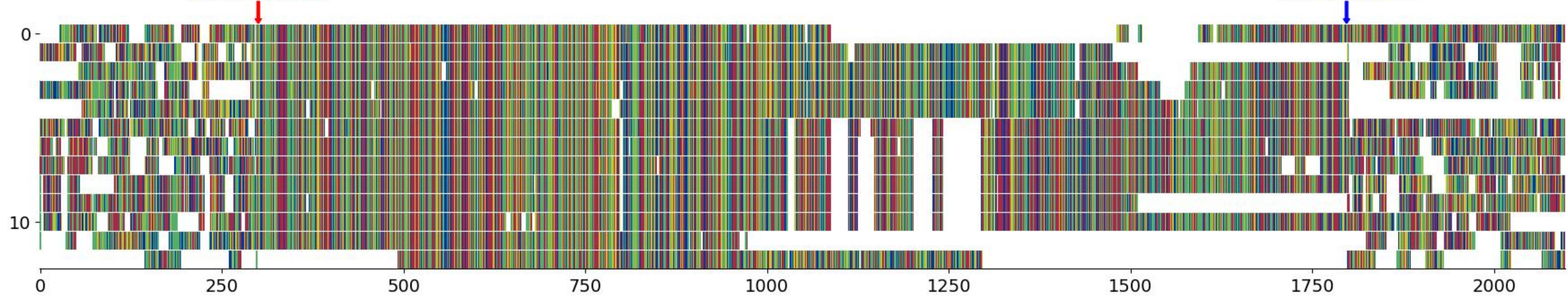
End crop Point

MSA length = 1498



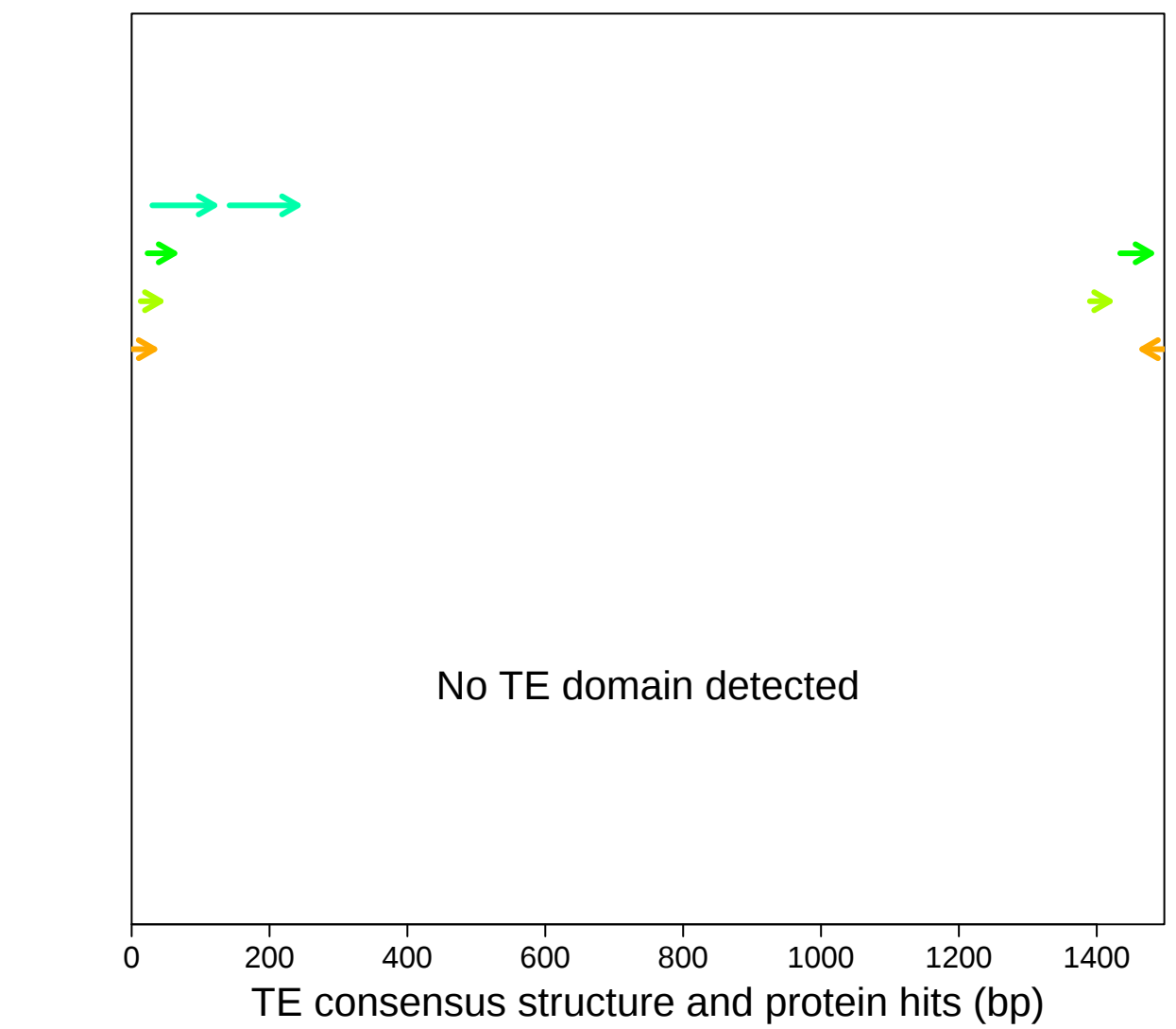
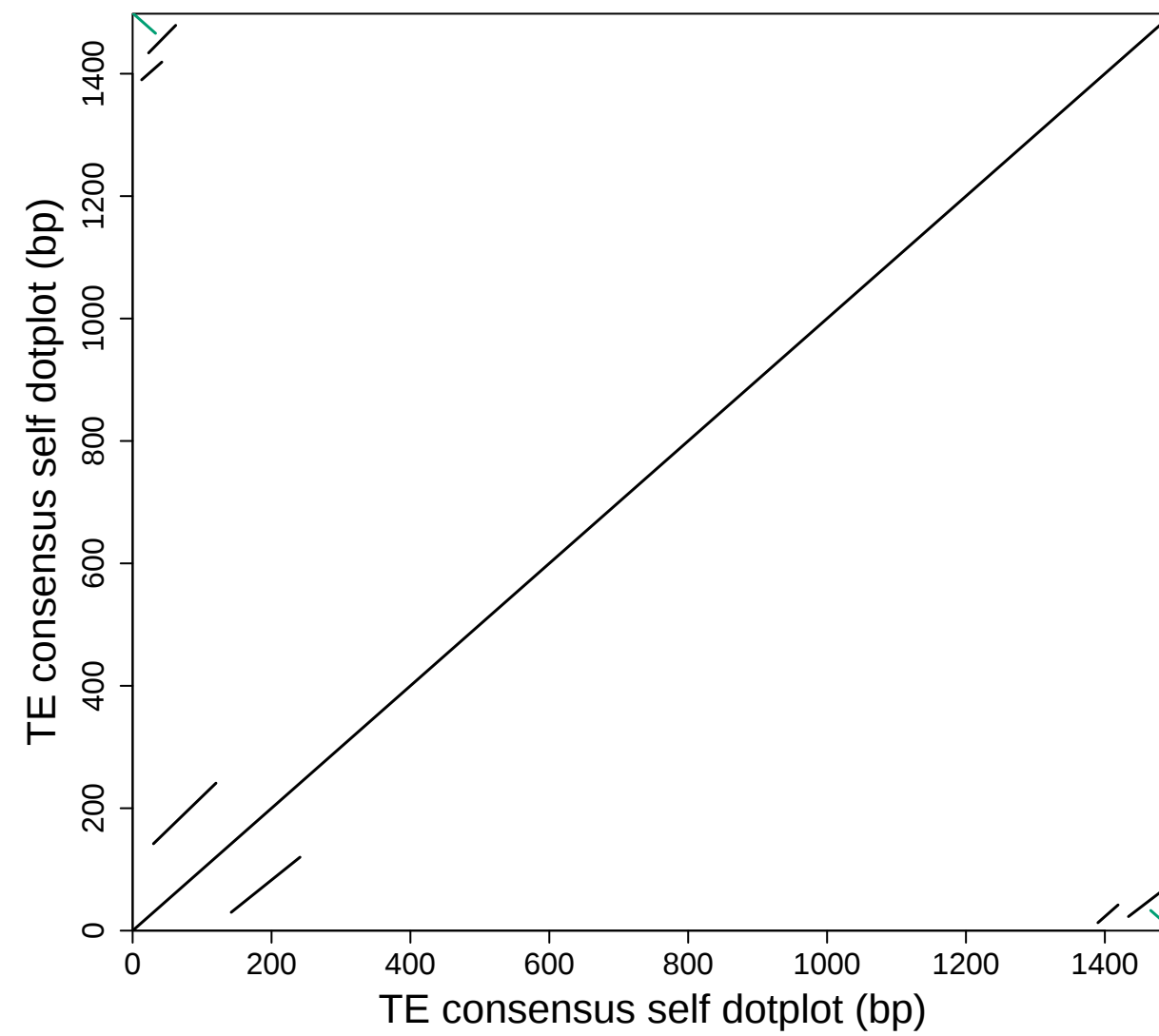
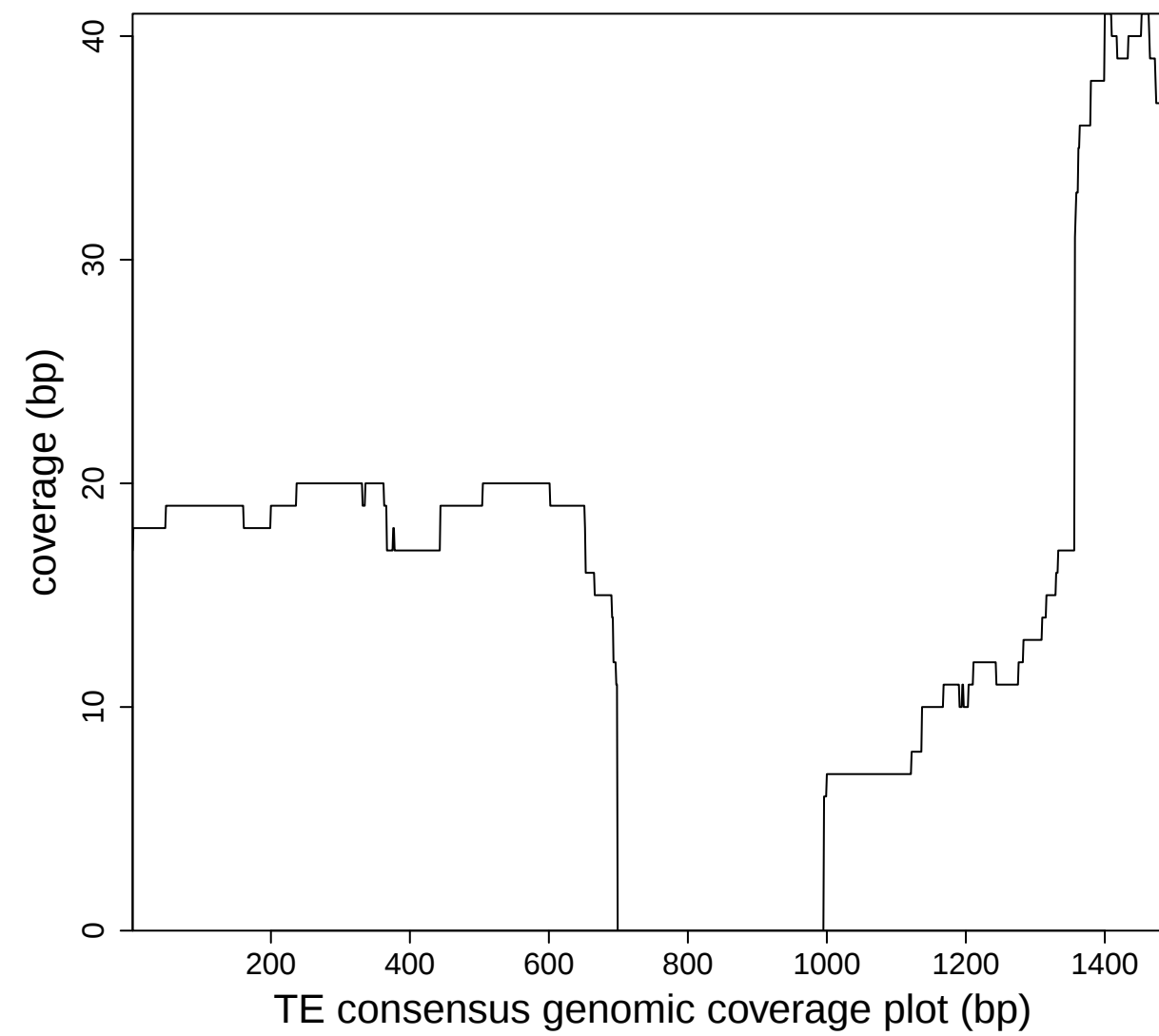
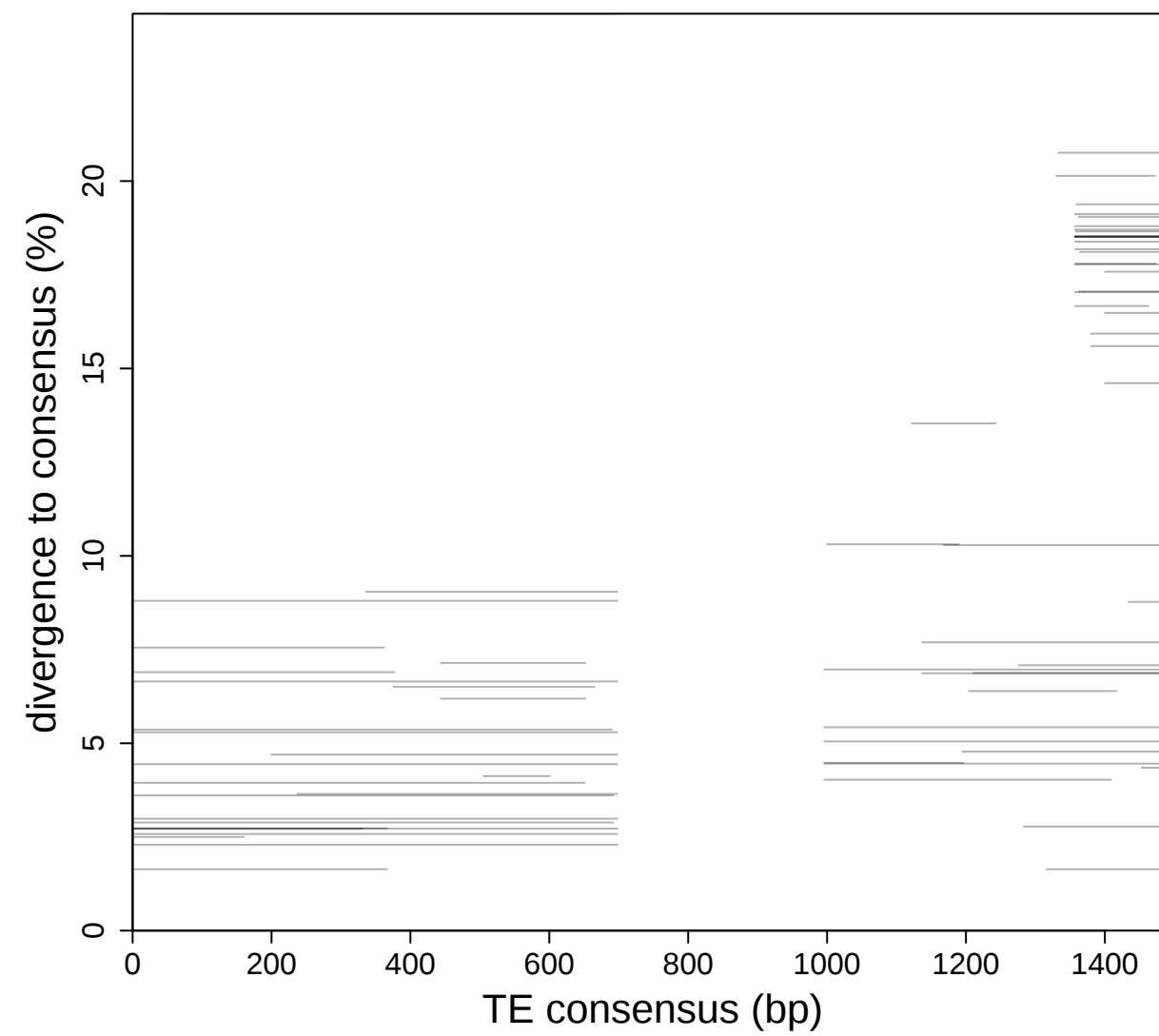
Start crop Point

End crop Point



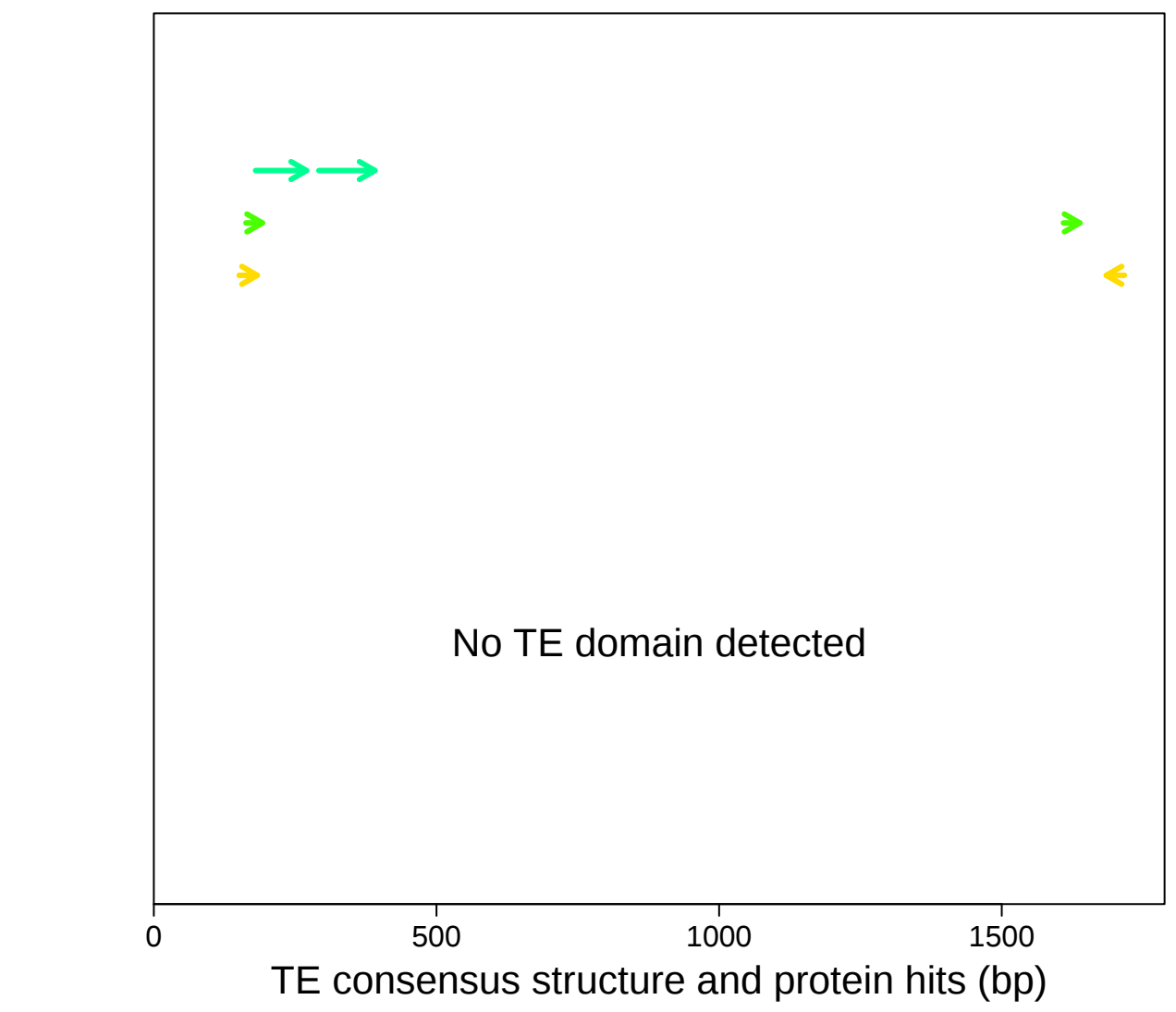
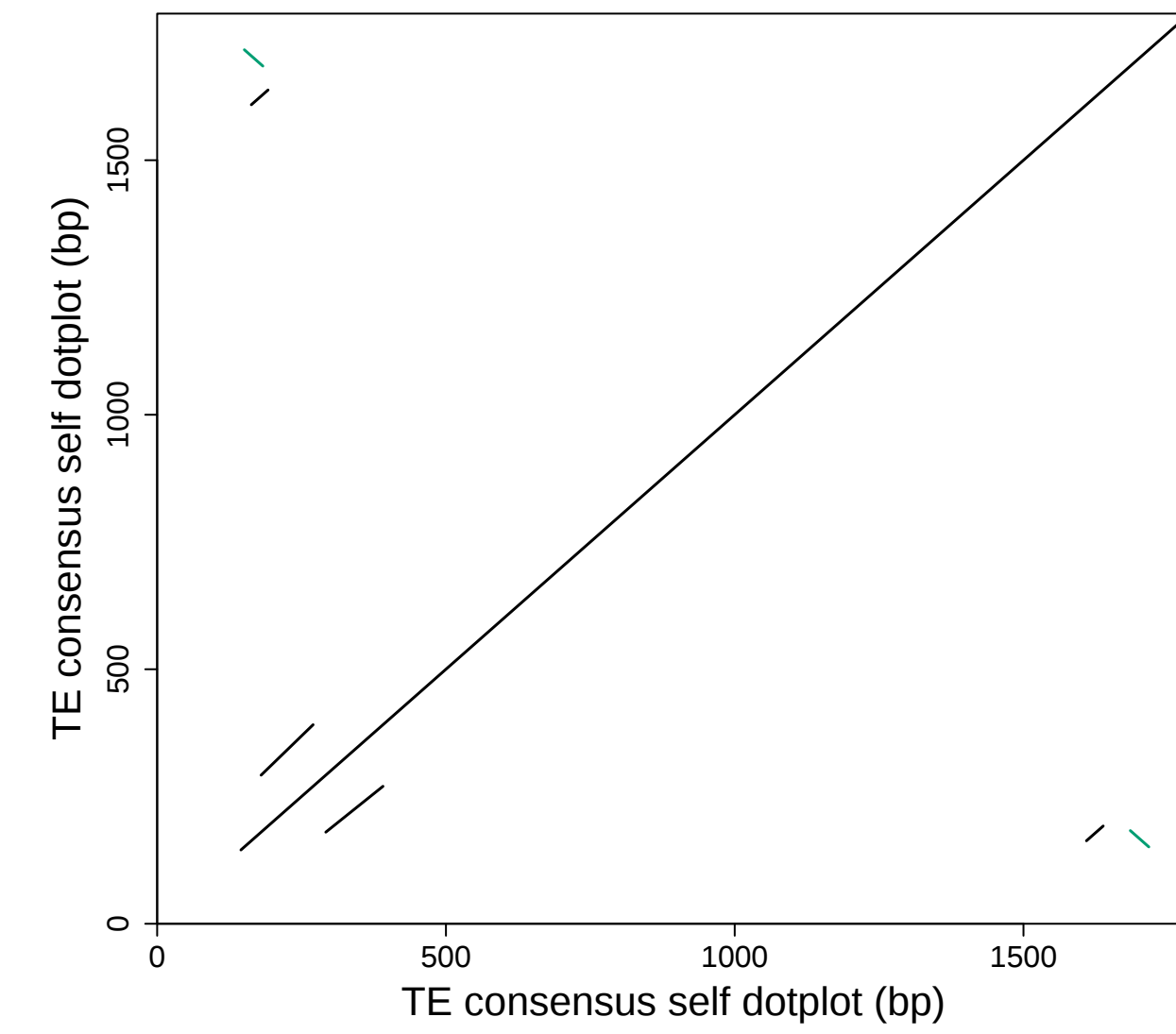
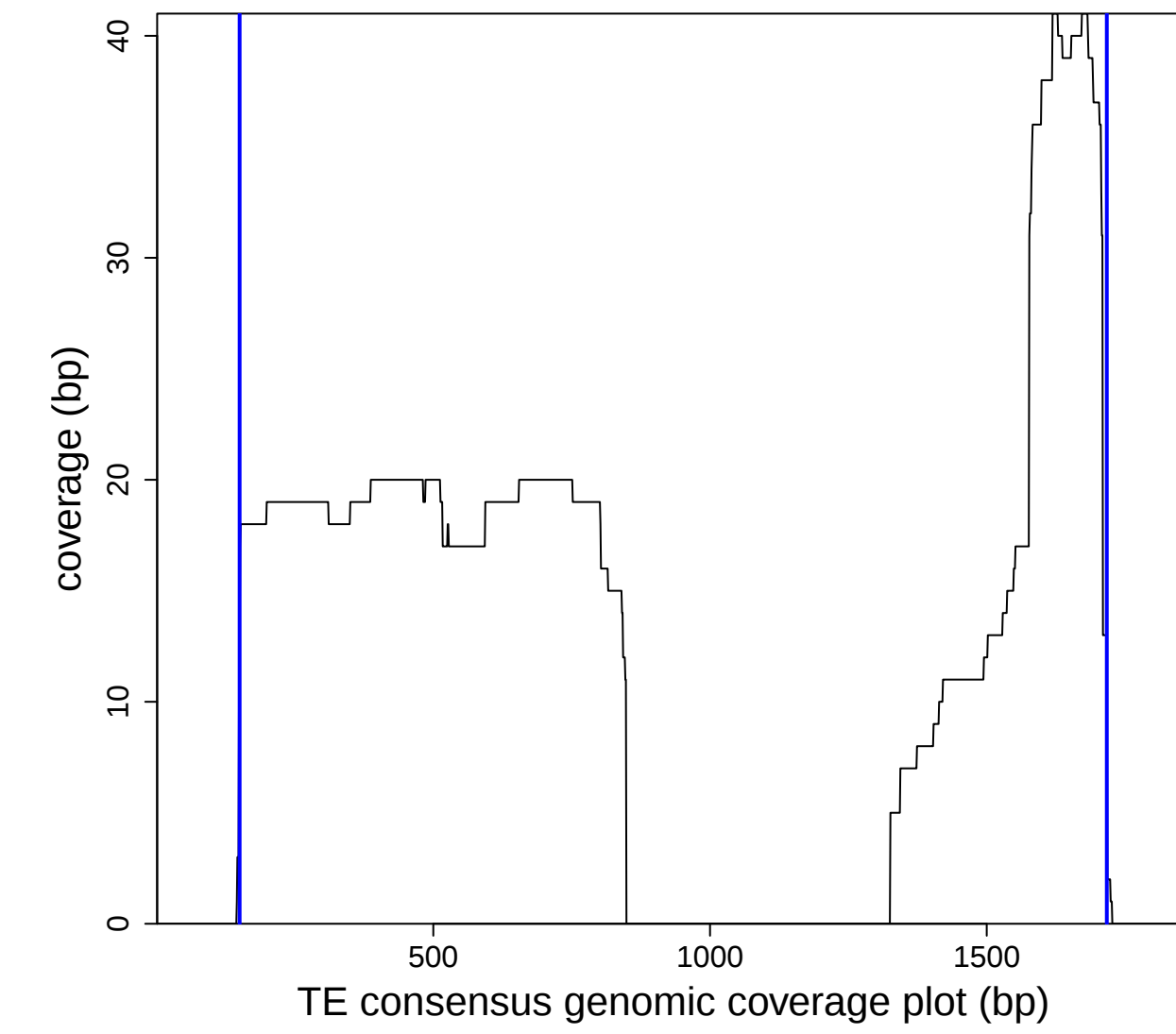
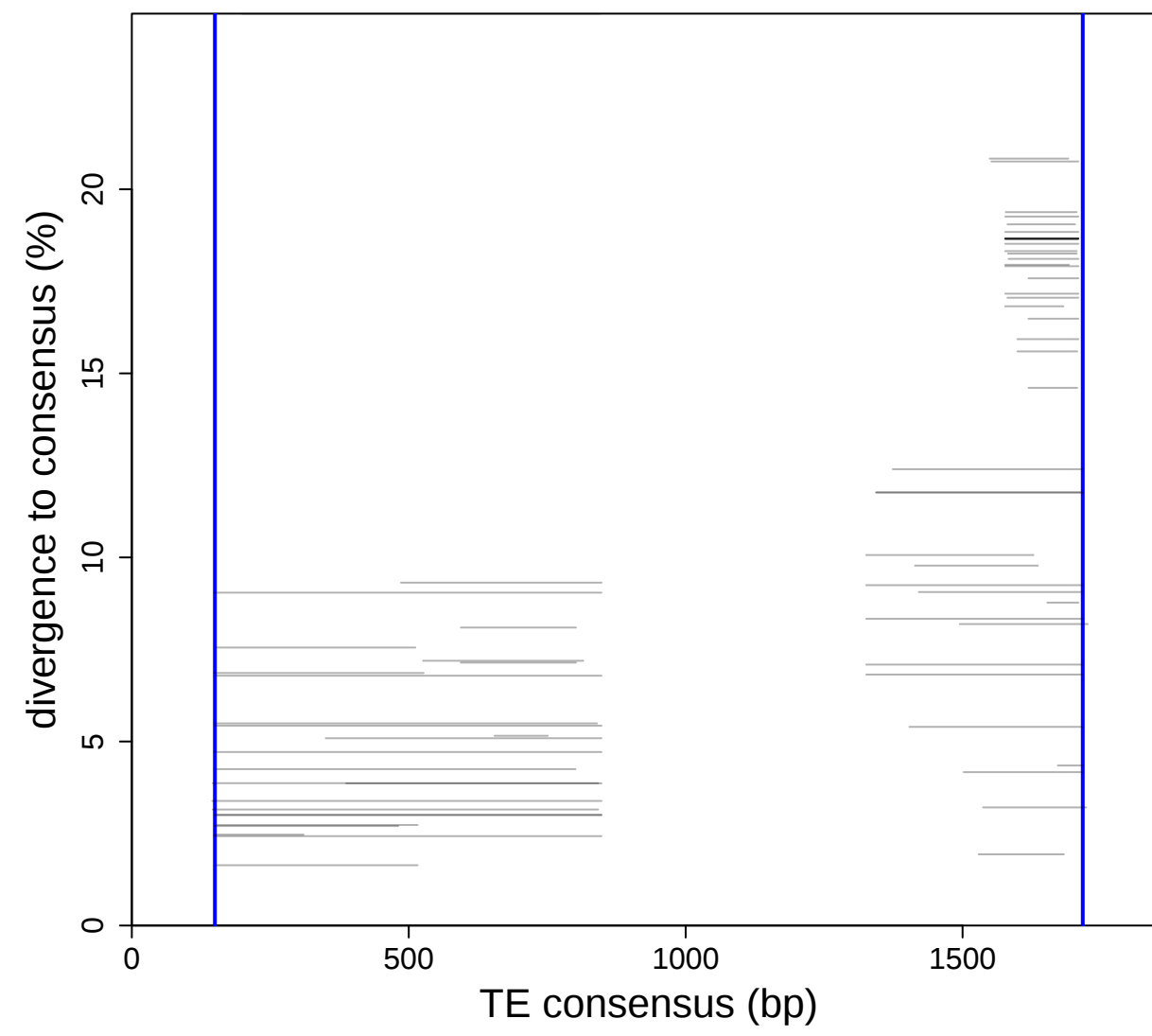
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1498bp; fragments: 72; full length: 0 (>=1348.2bp)

After TEtrimmer 1498 bp

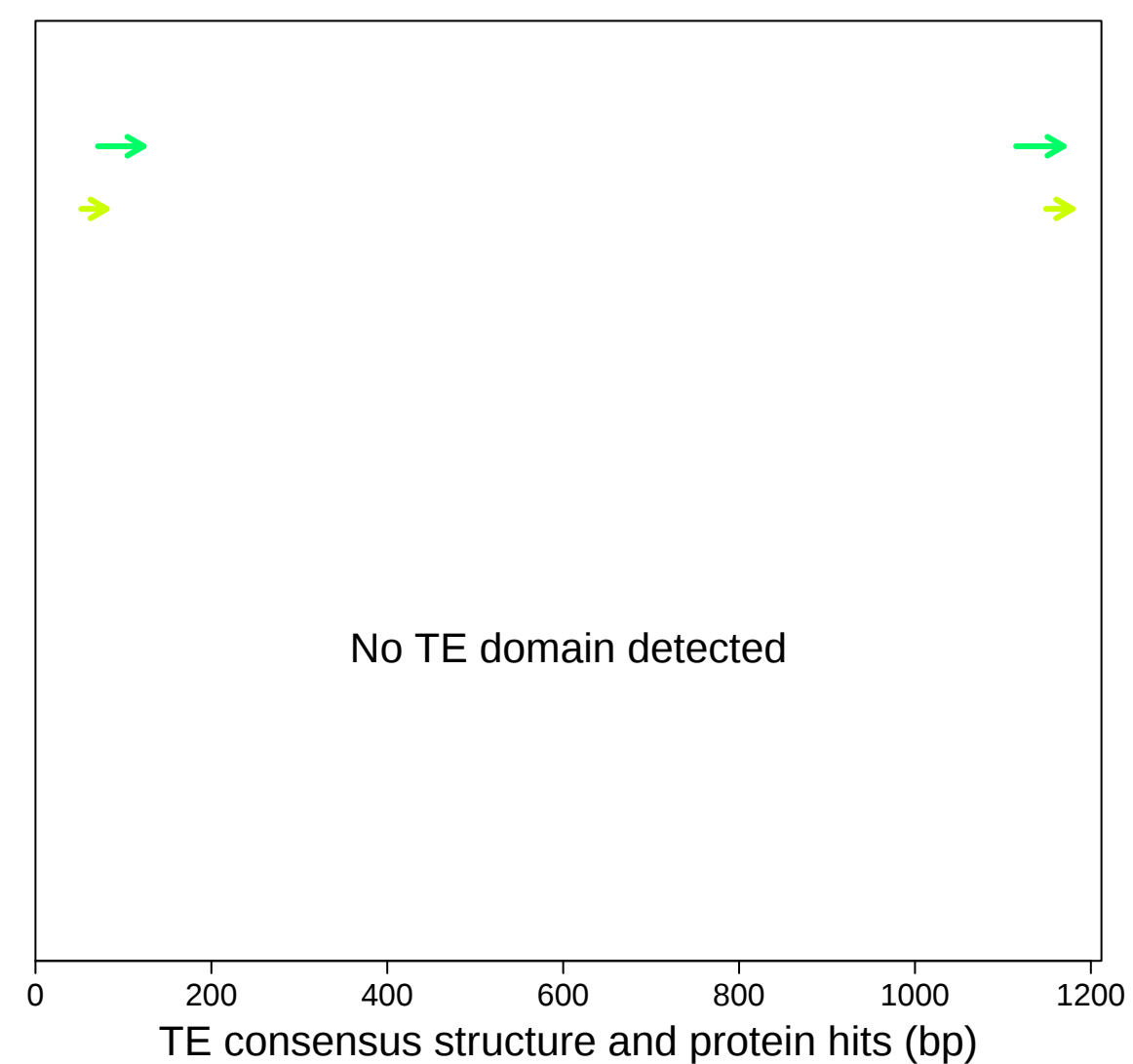
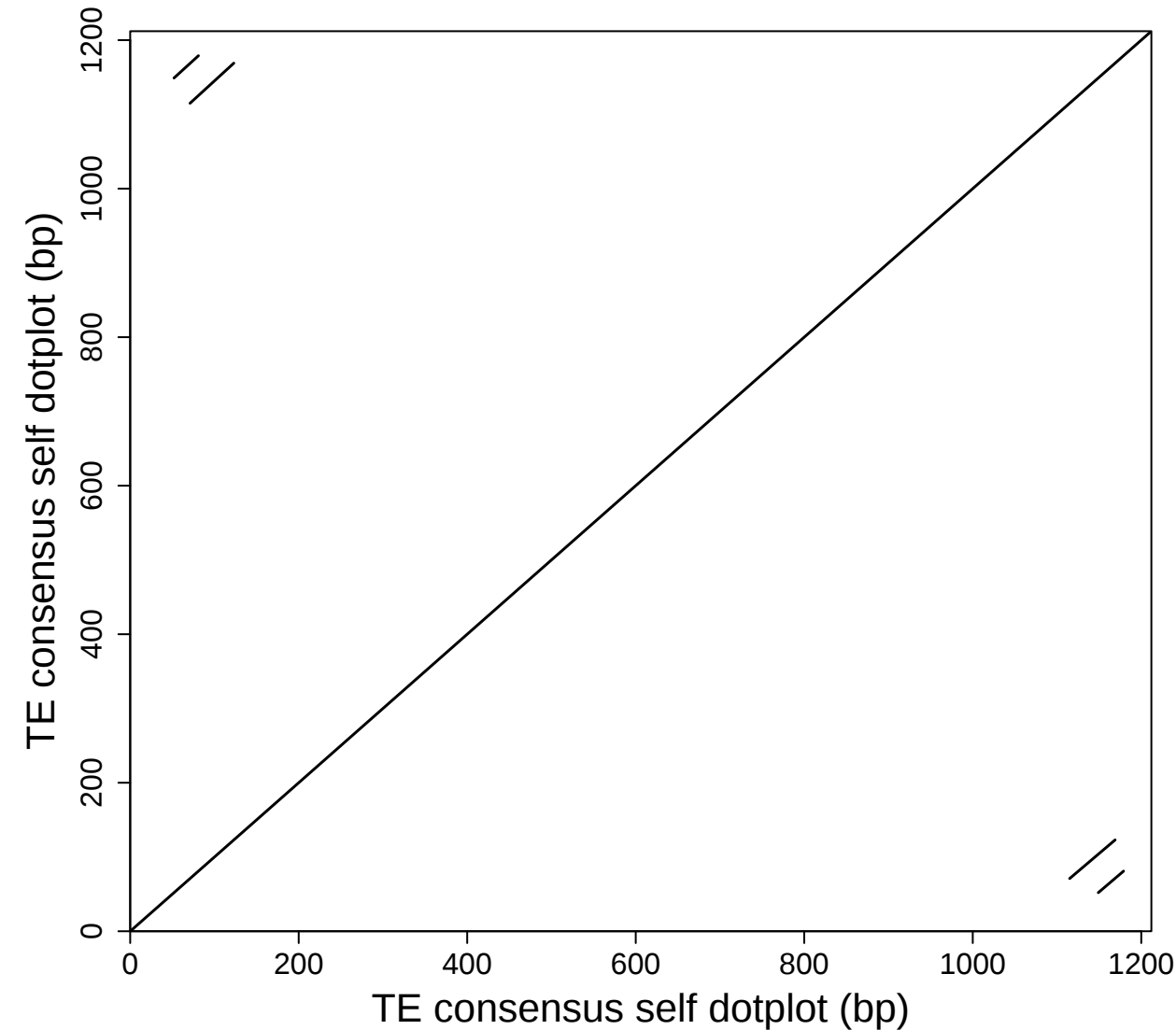
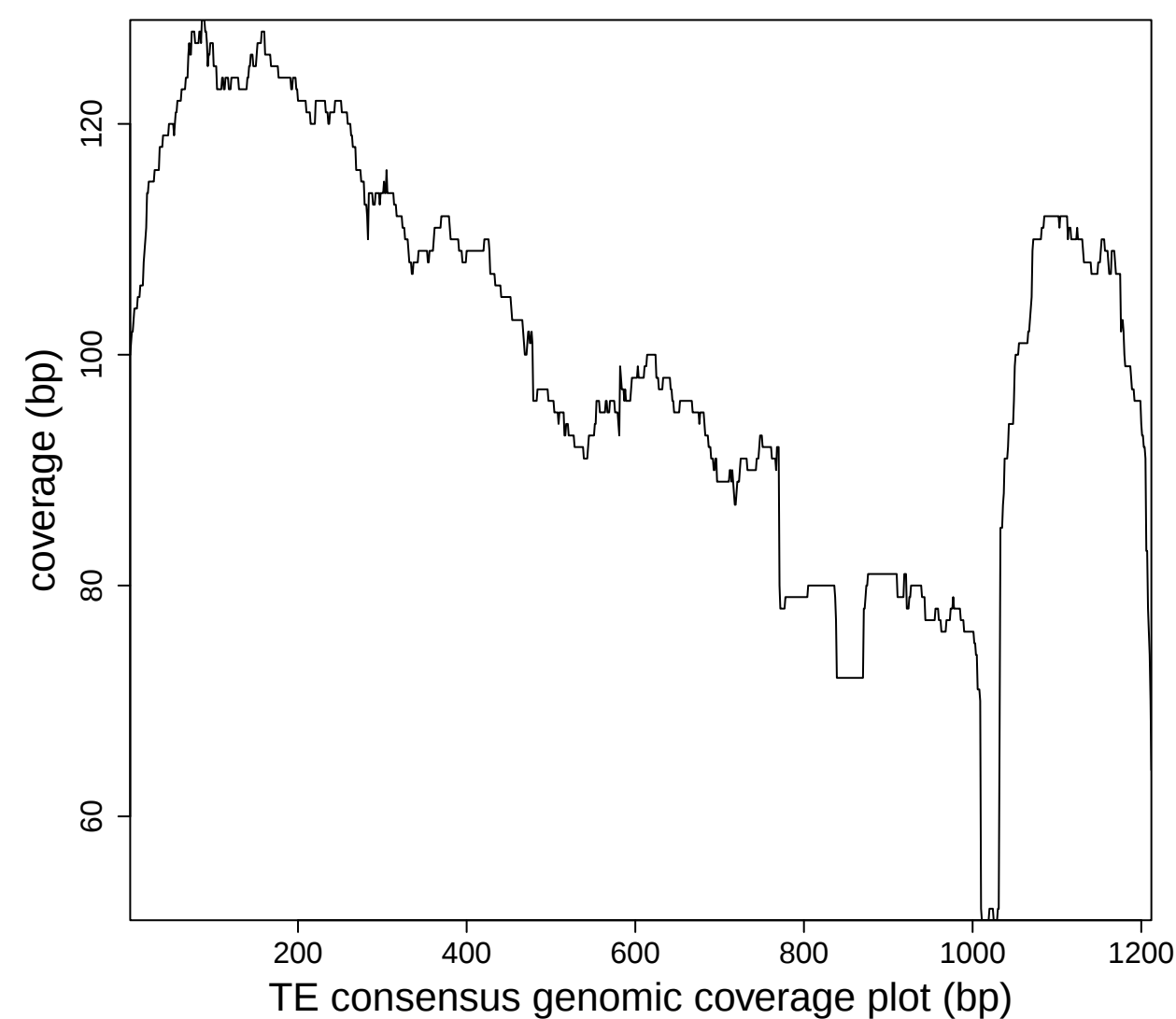
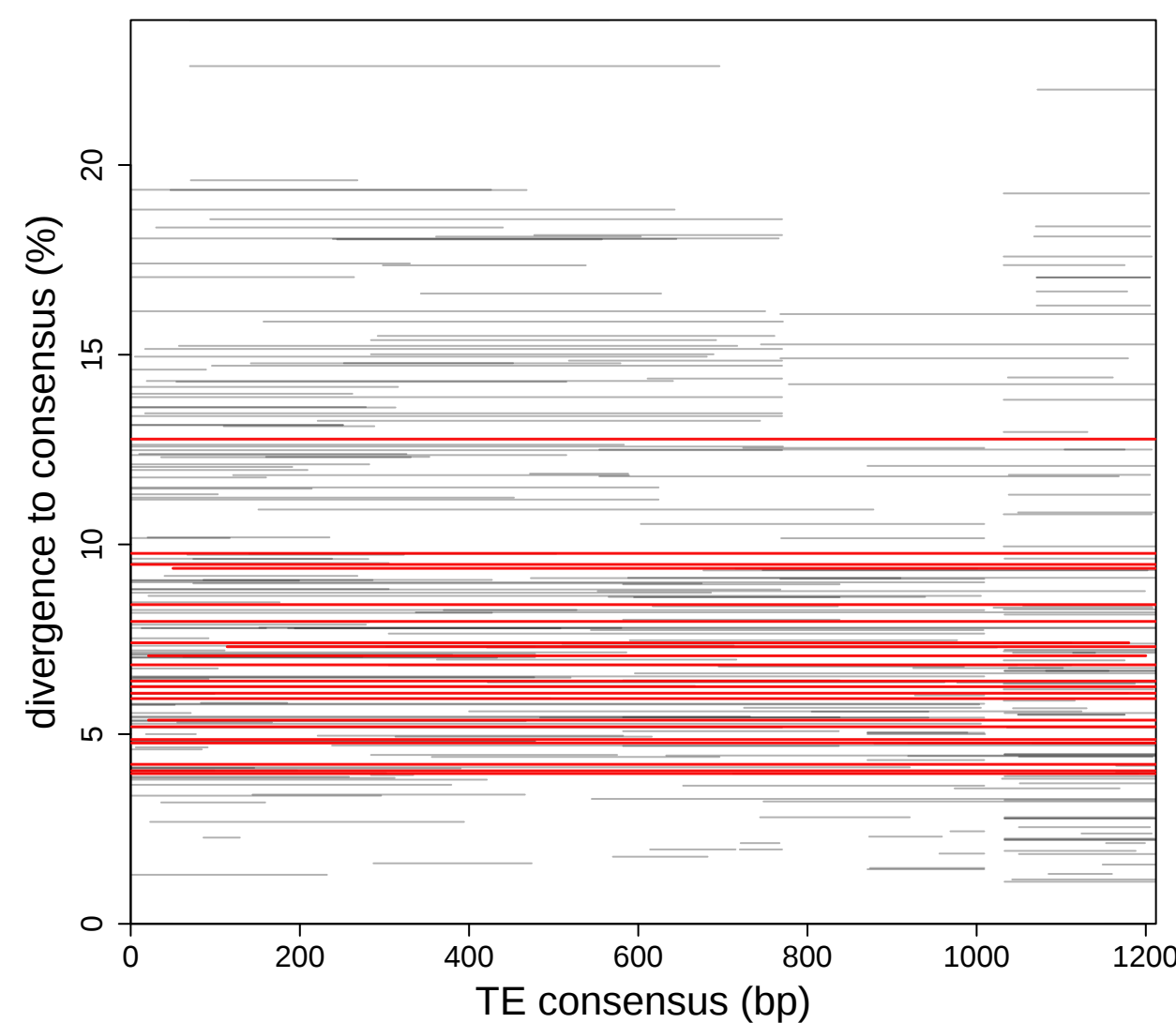


5.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 1867bp; fragments: 69; full length: 0 (>=1680.3bp)

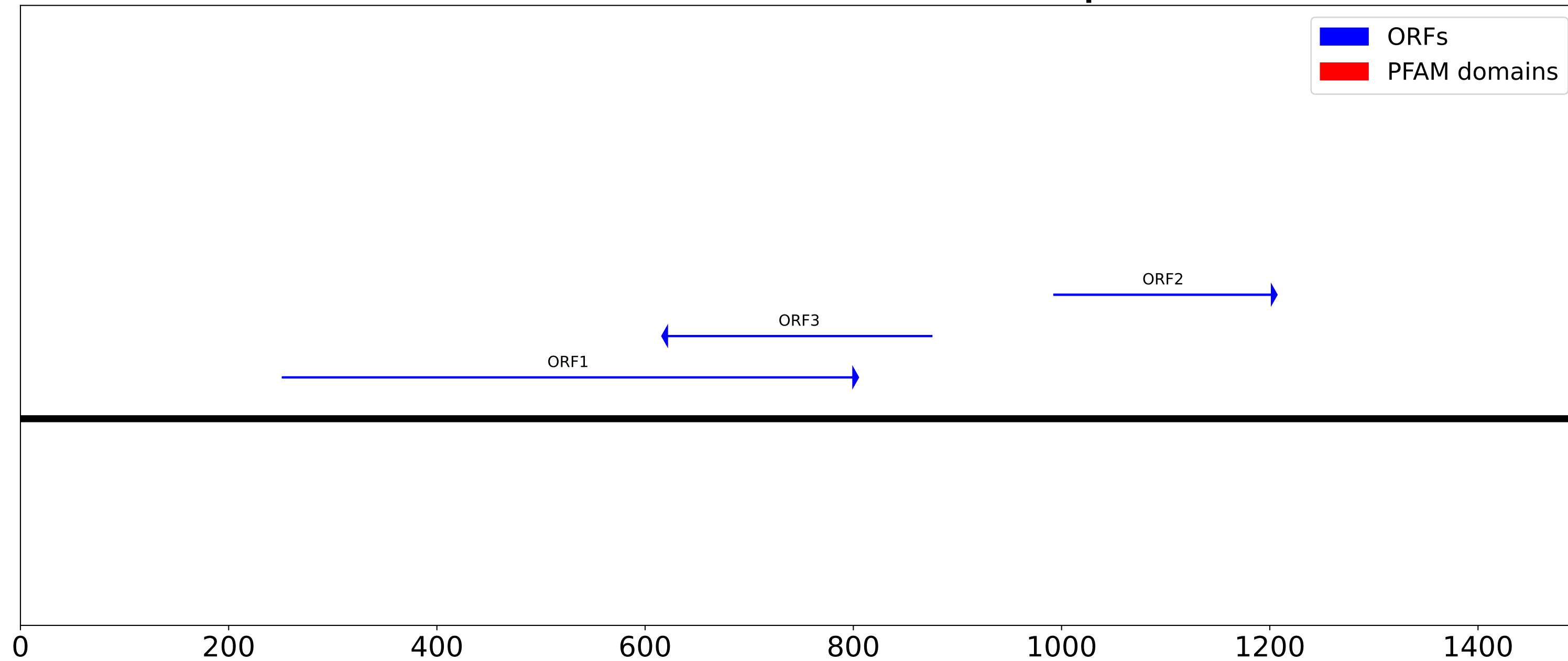
After TEtrimmer Extended plot Blue lines are boundaries



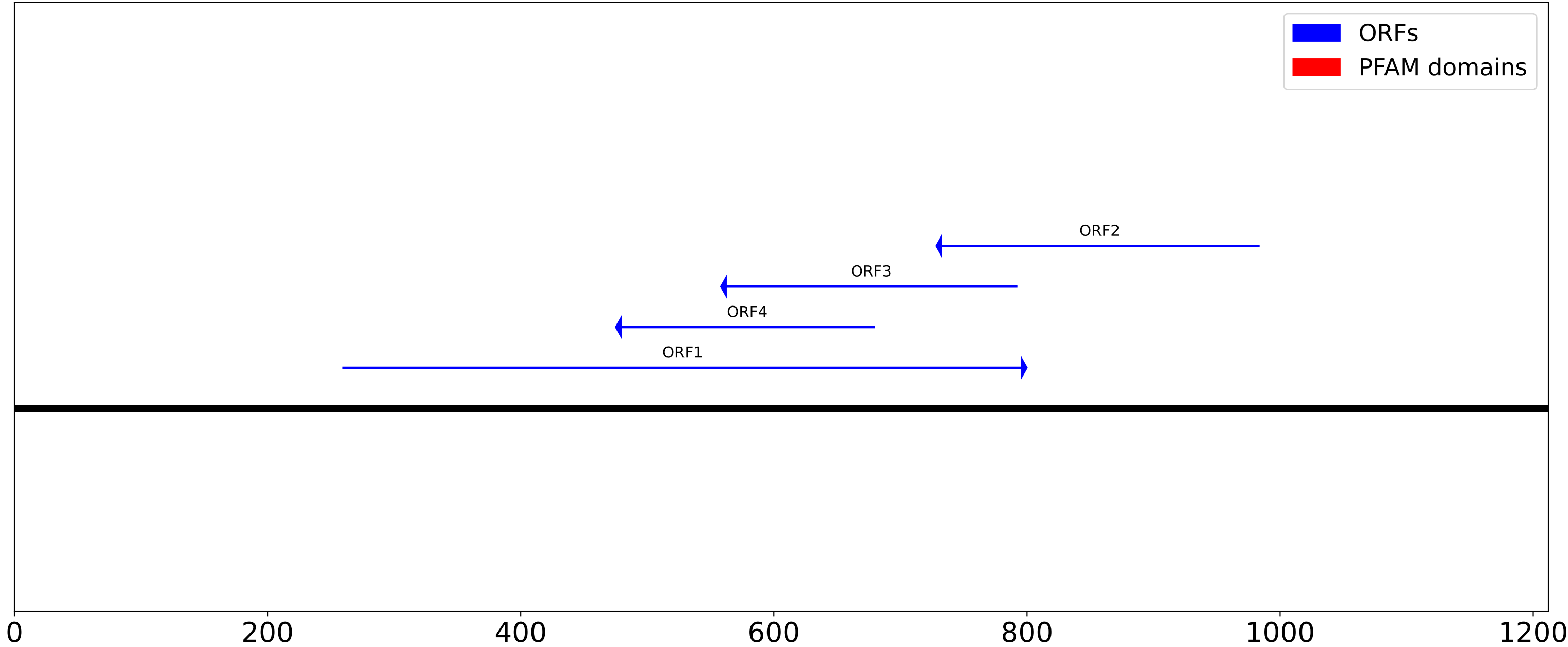
TE: rnd_5_family_2525
size: 1212bp; fragments: 311; full length: 21 (≥ 1090.8 bp)



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

