

Start crop Point



End crop Point

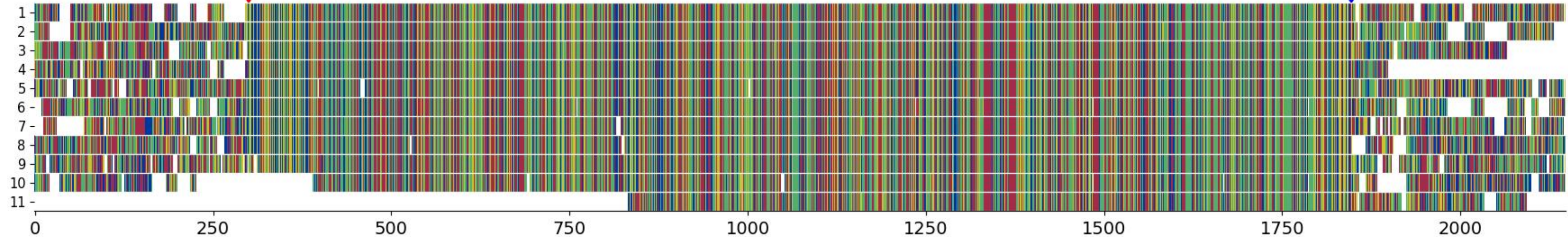


MSA length = 1548



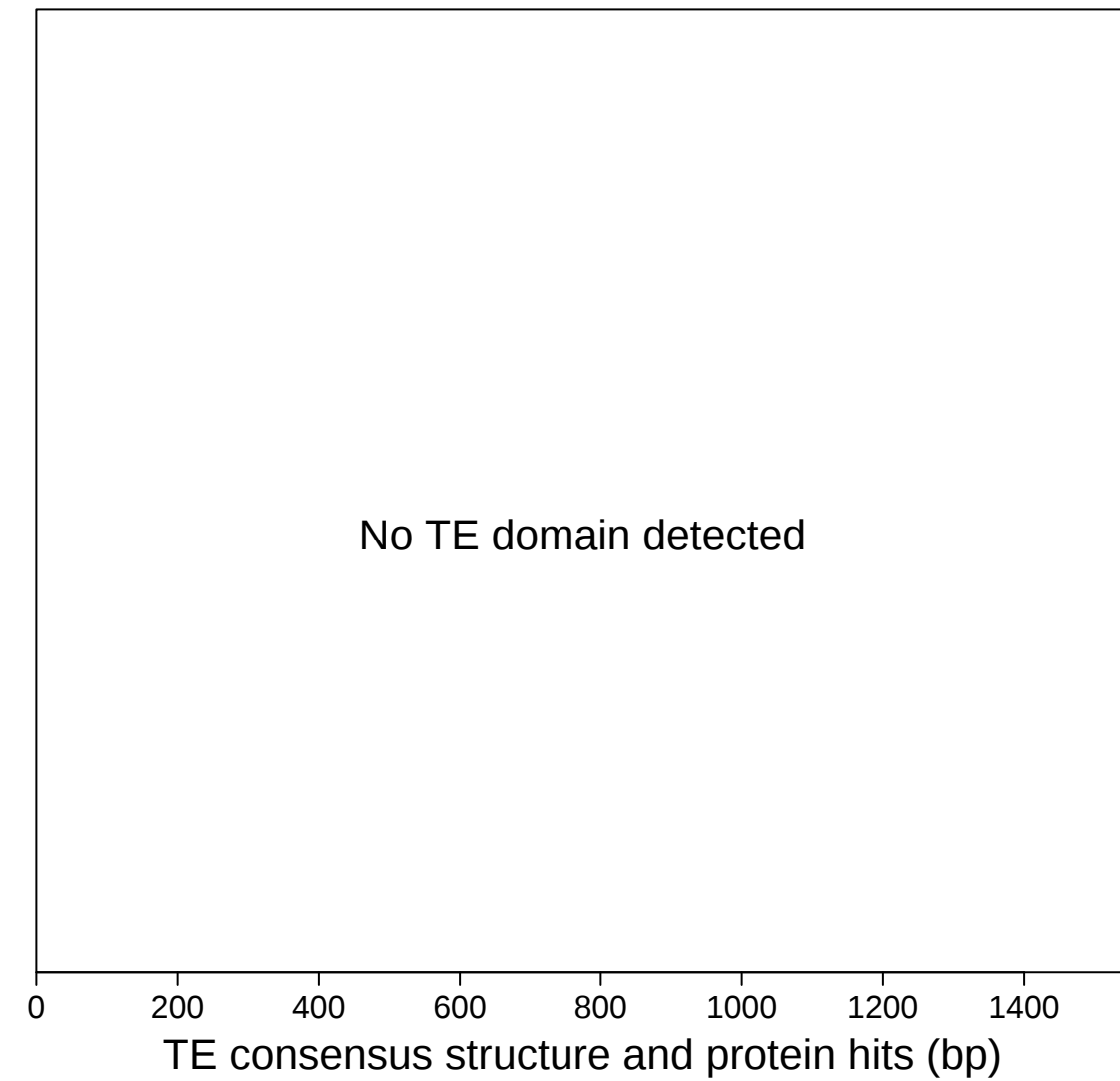
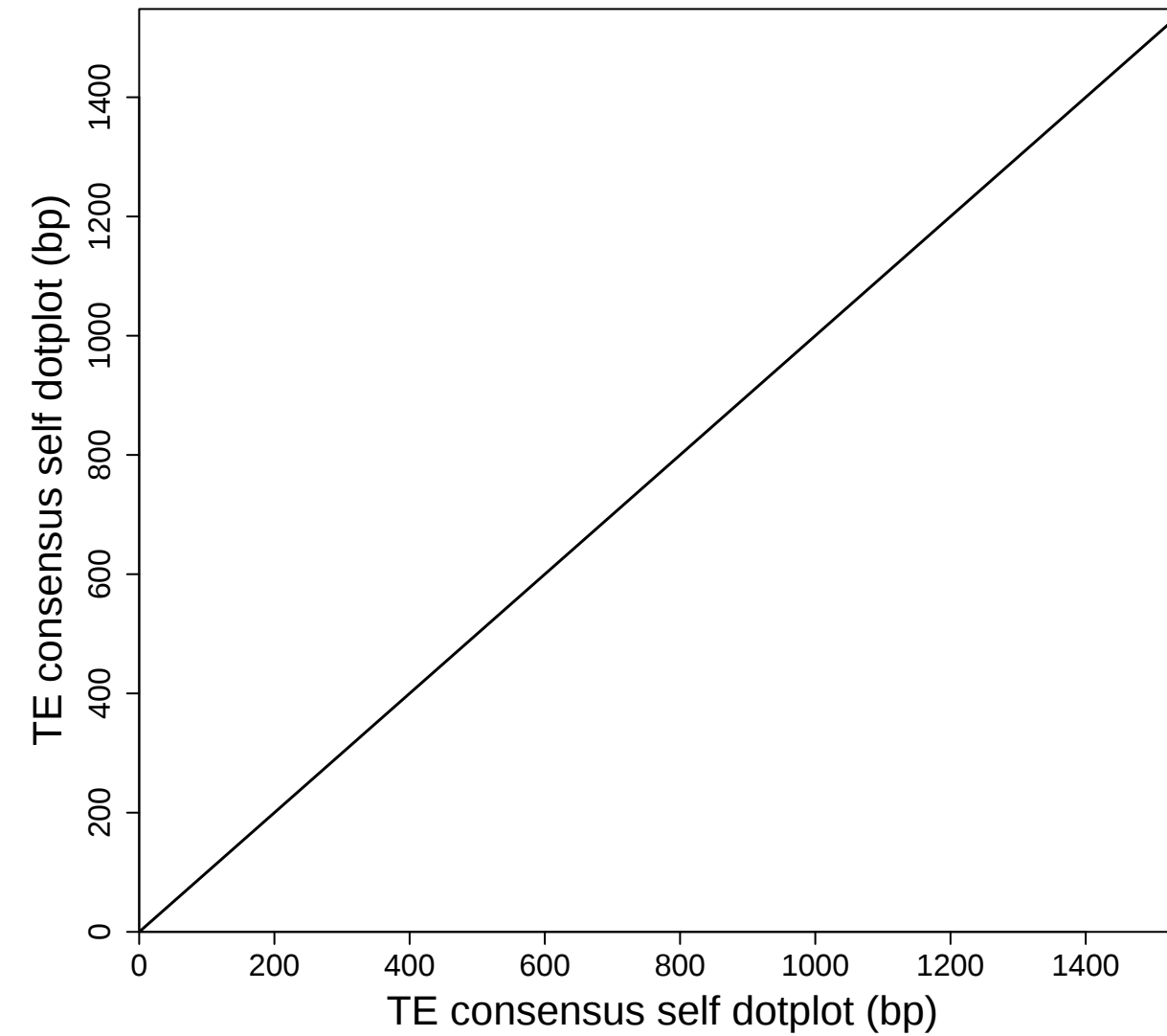
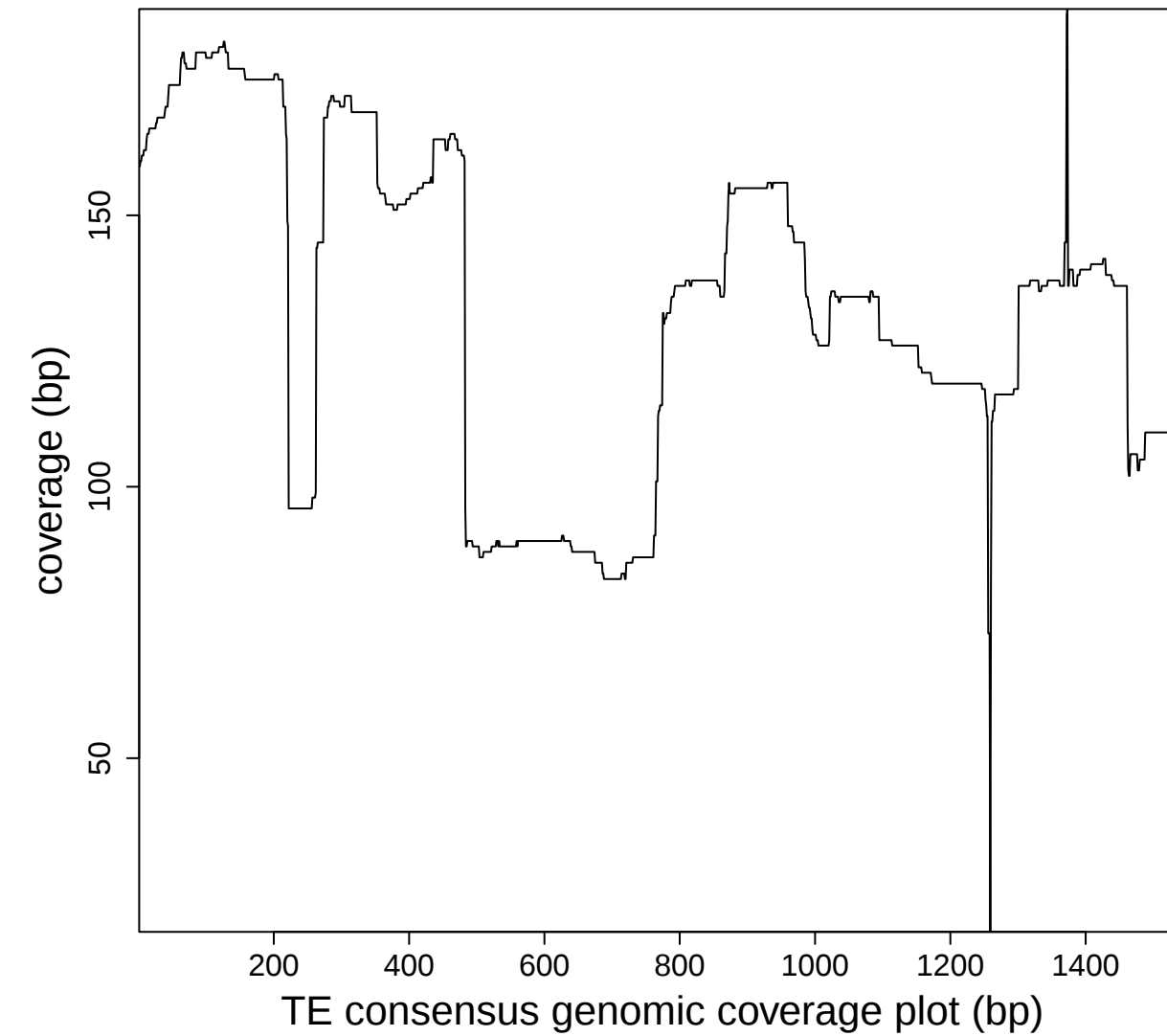
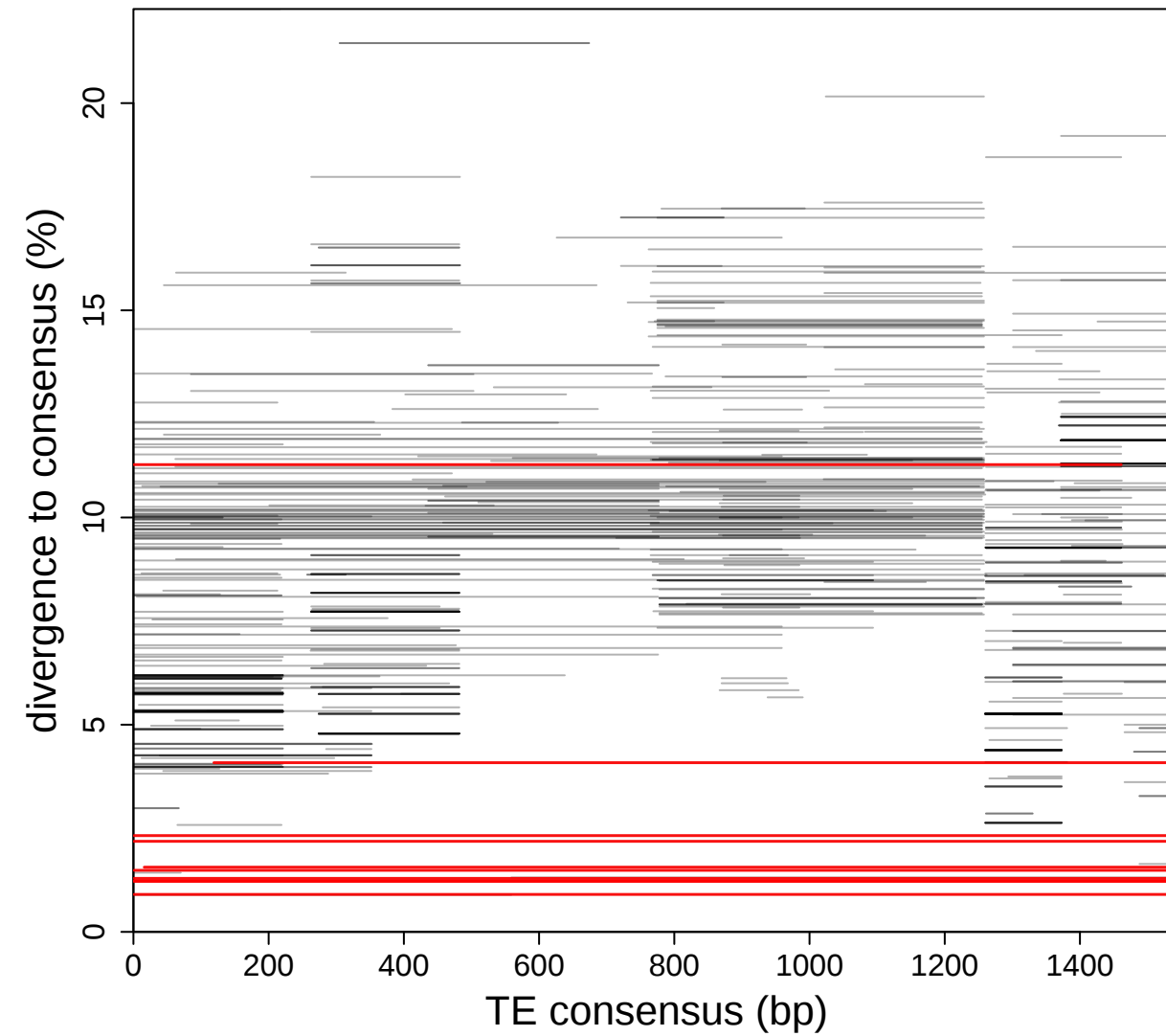
Start crop Point

End crop Point



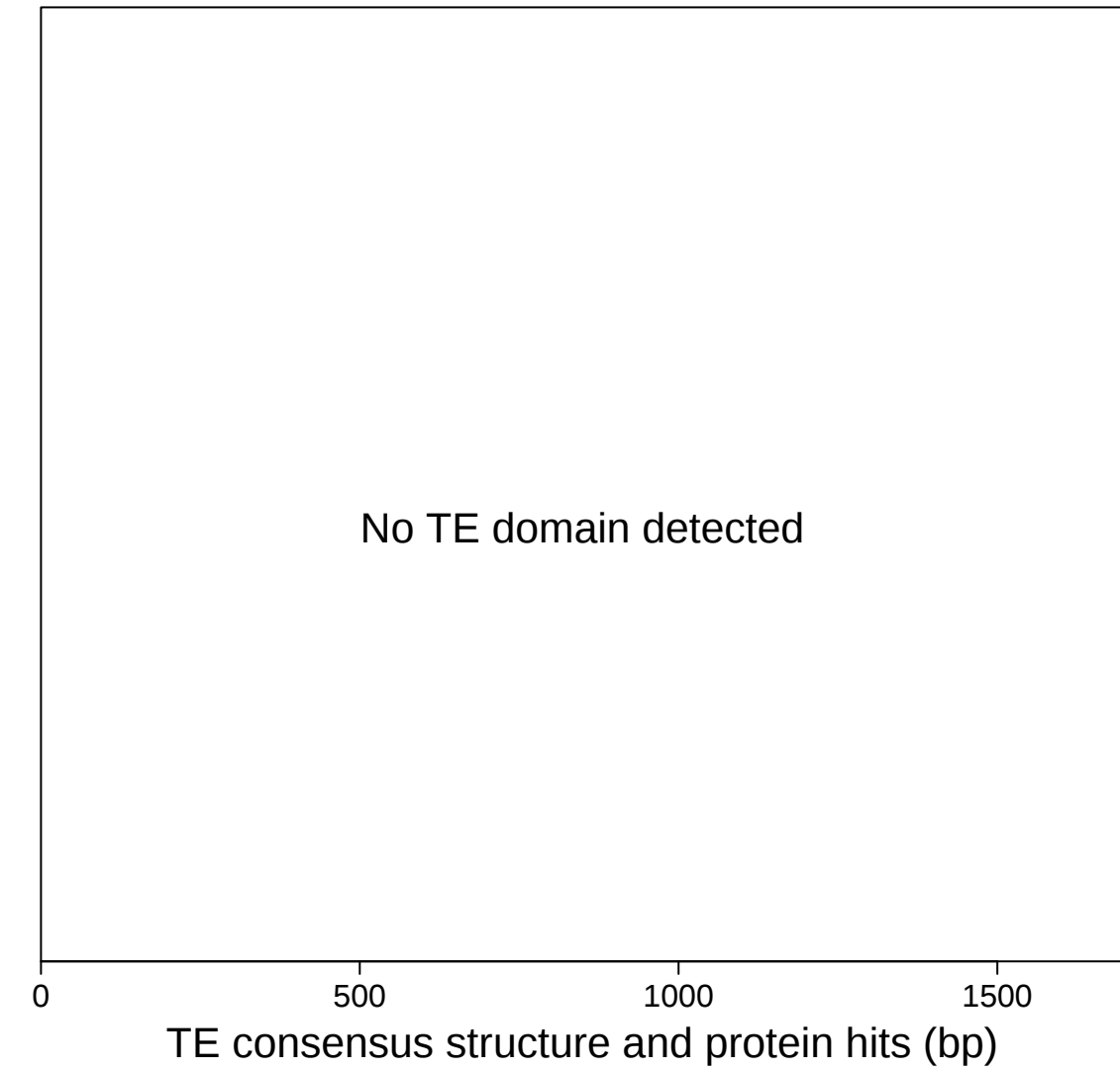
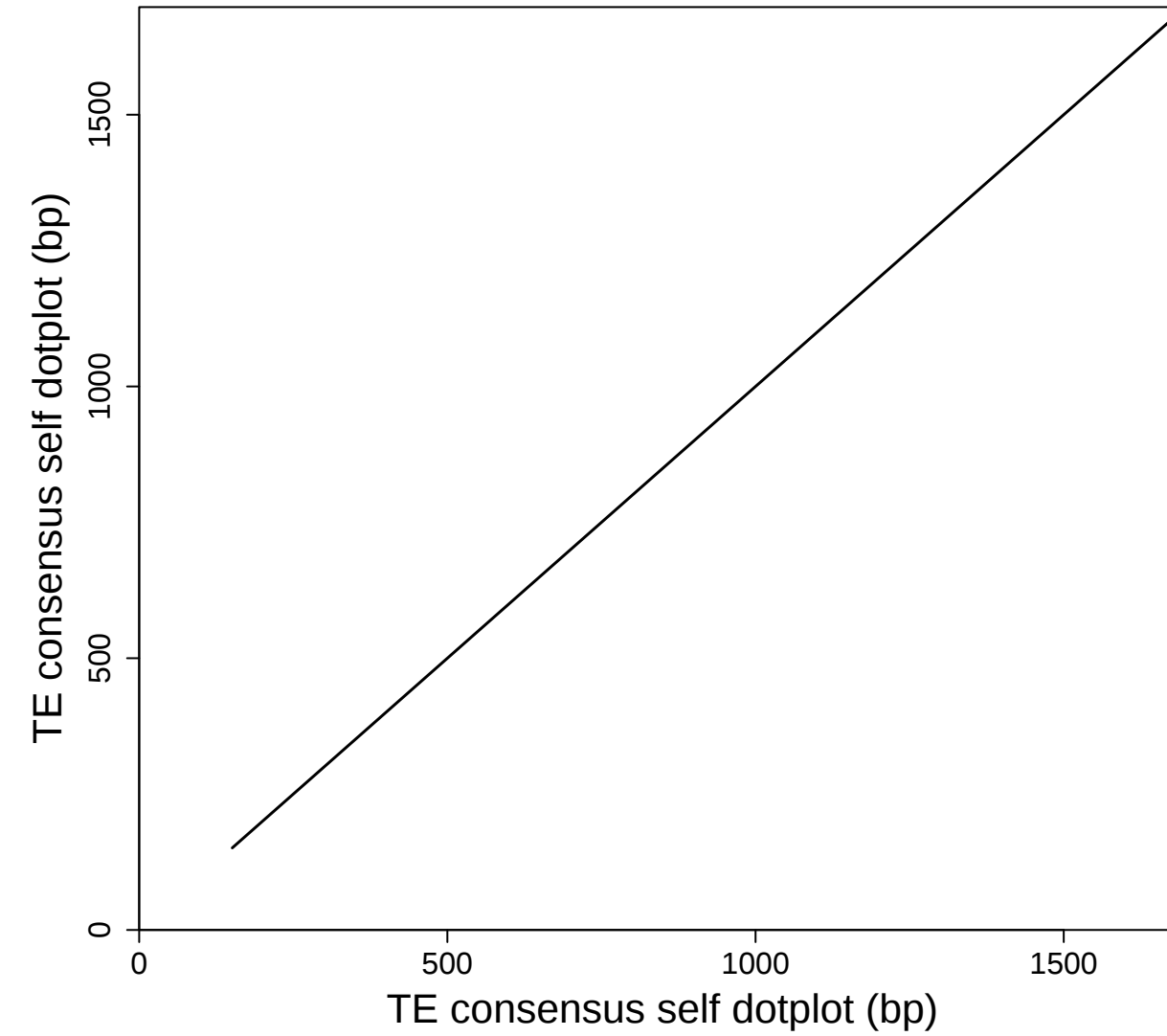
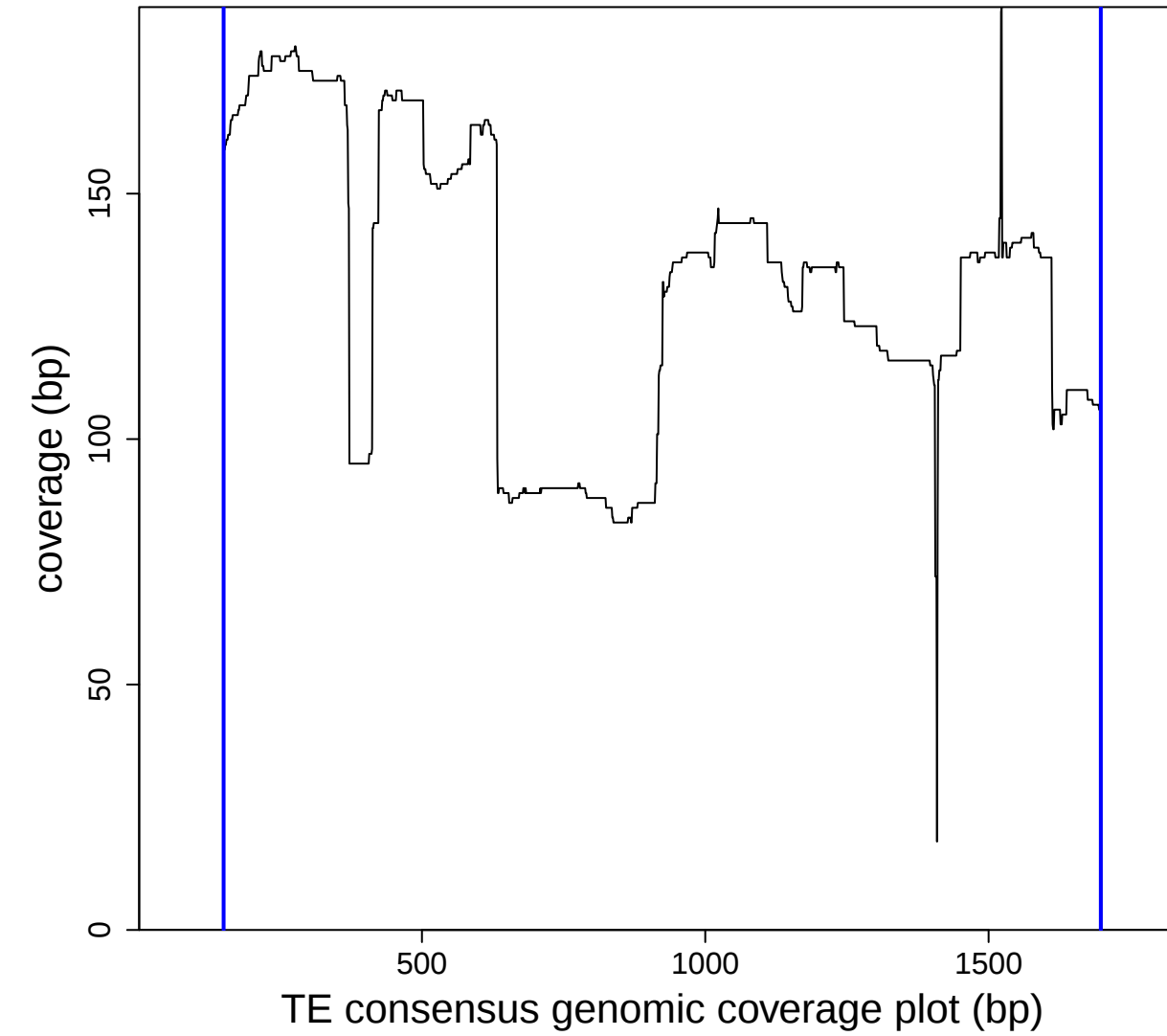
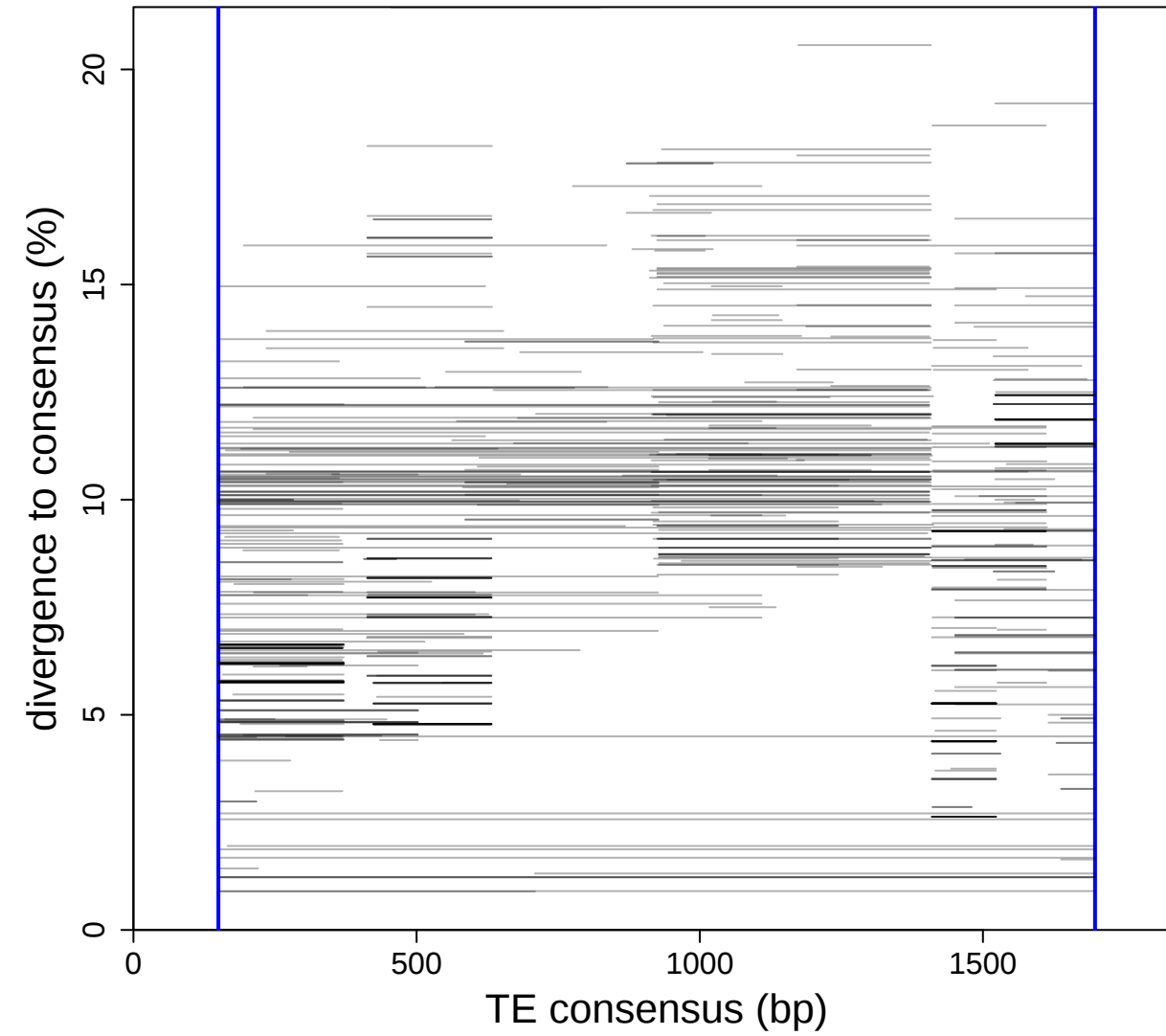
a.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1548bp; fragments: 597; full length: 11 (≥ 1393.2 bp)

After TETrimmer 1548 bp

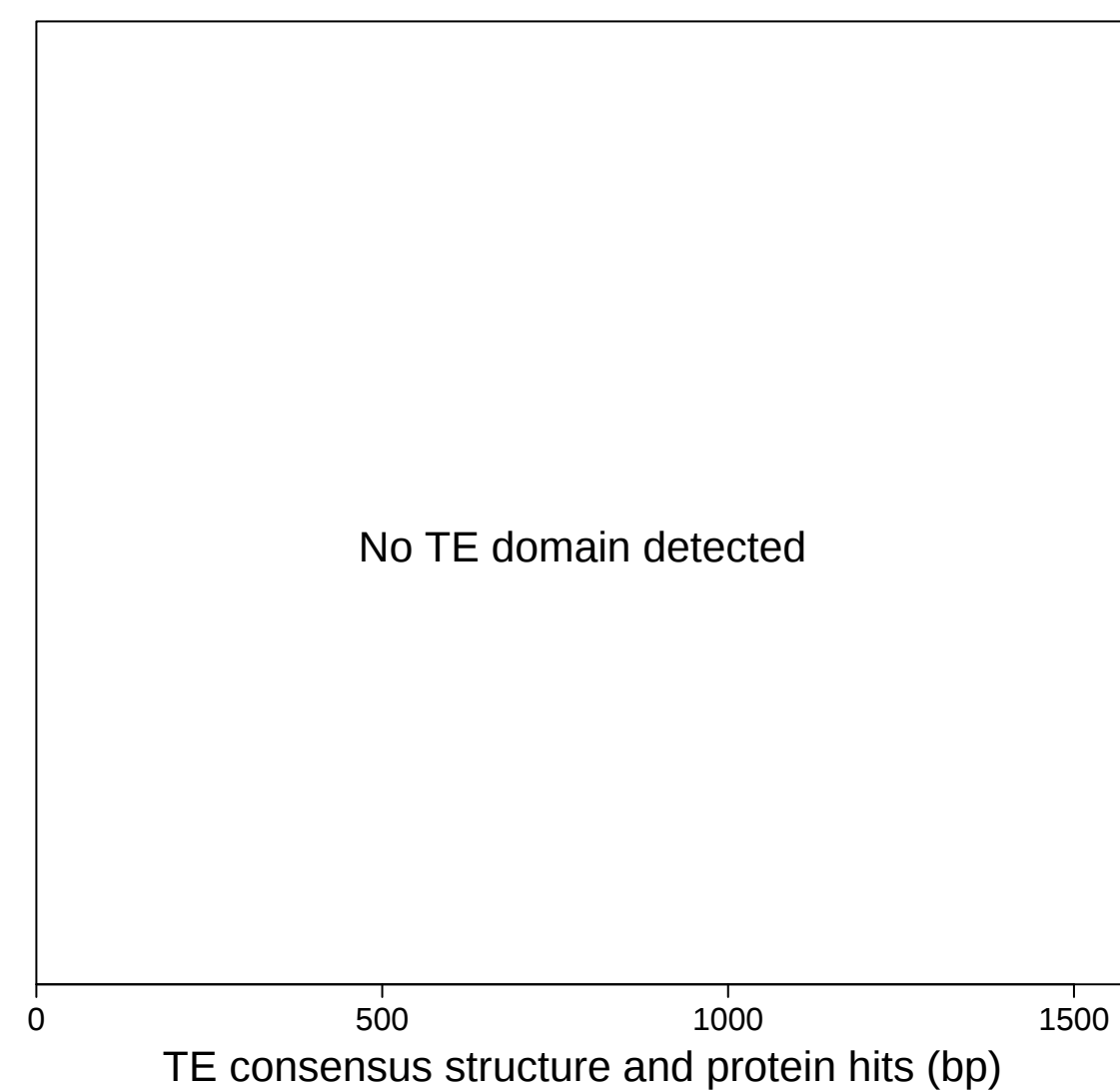
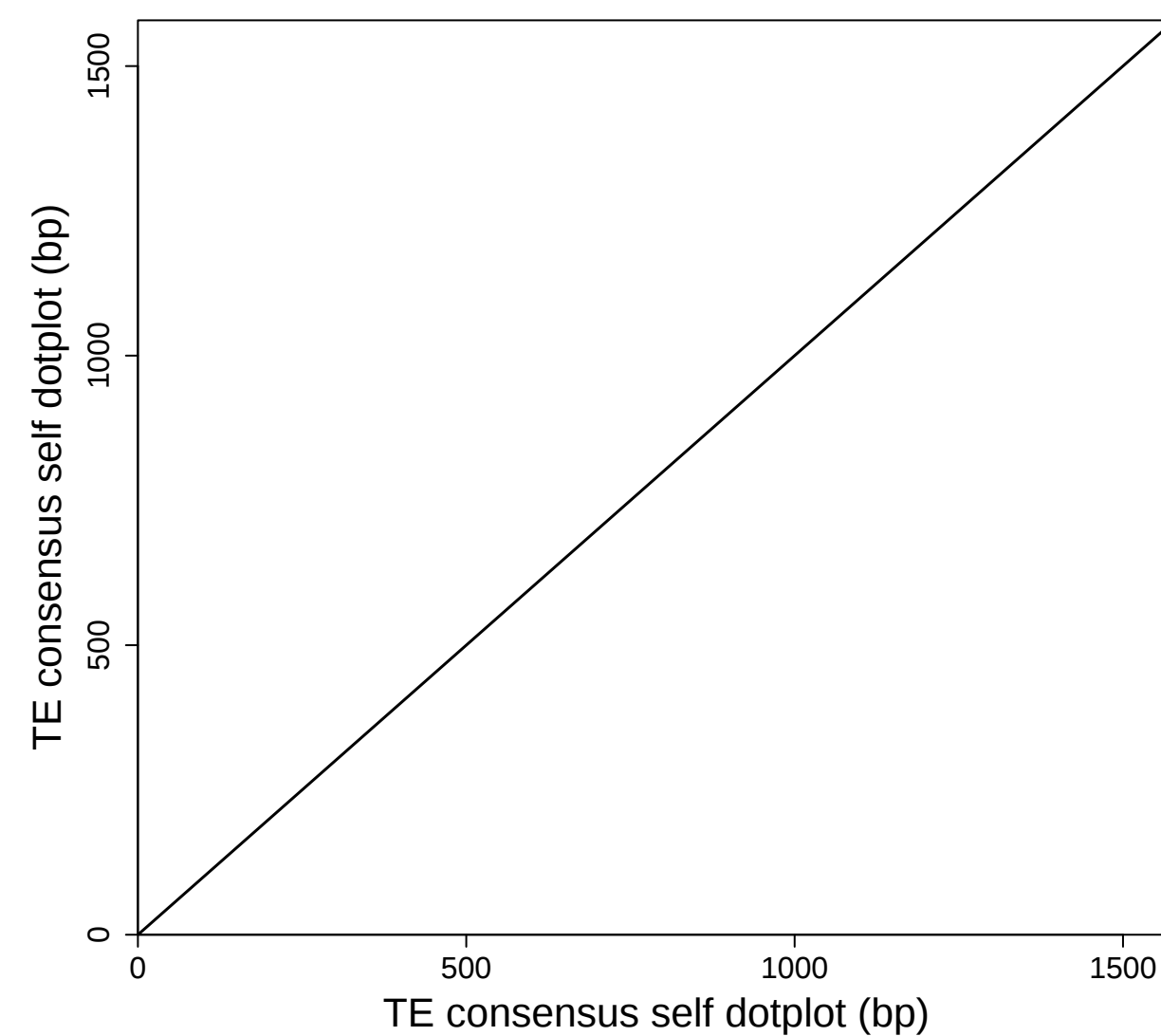
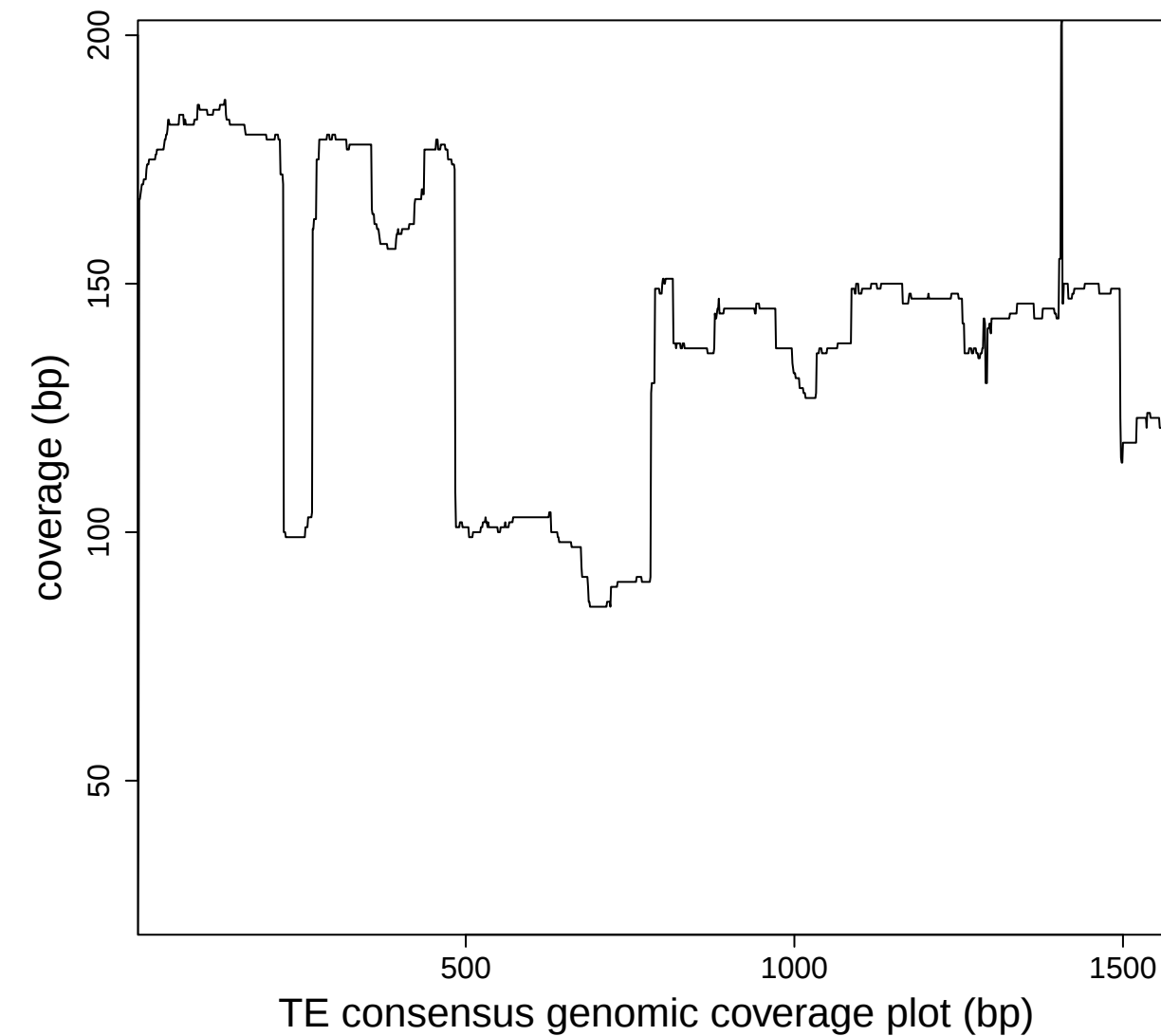
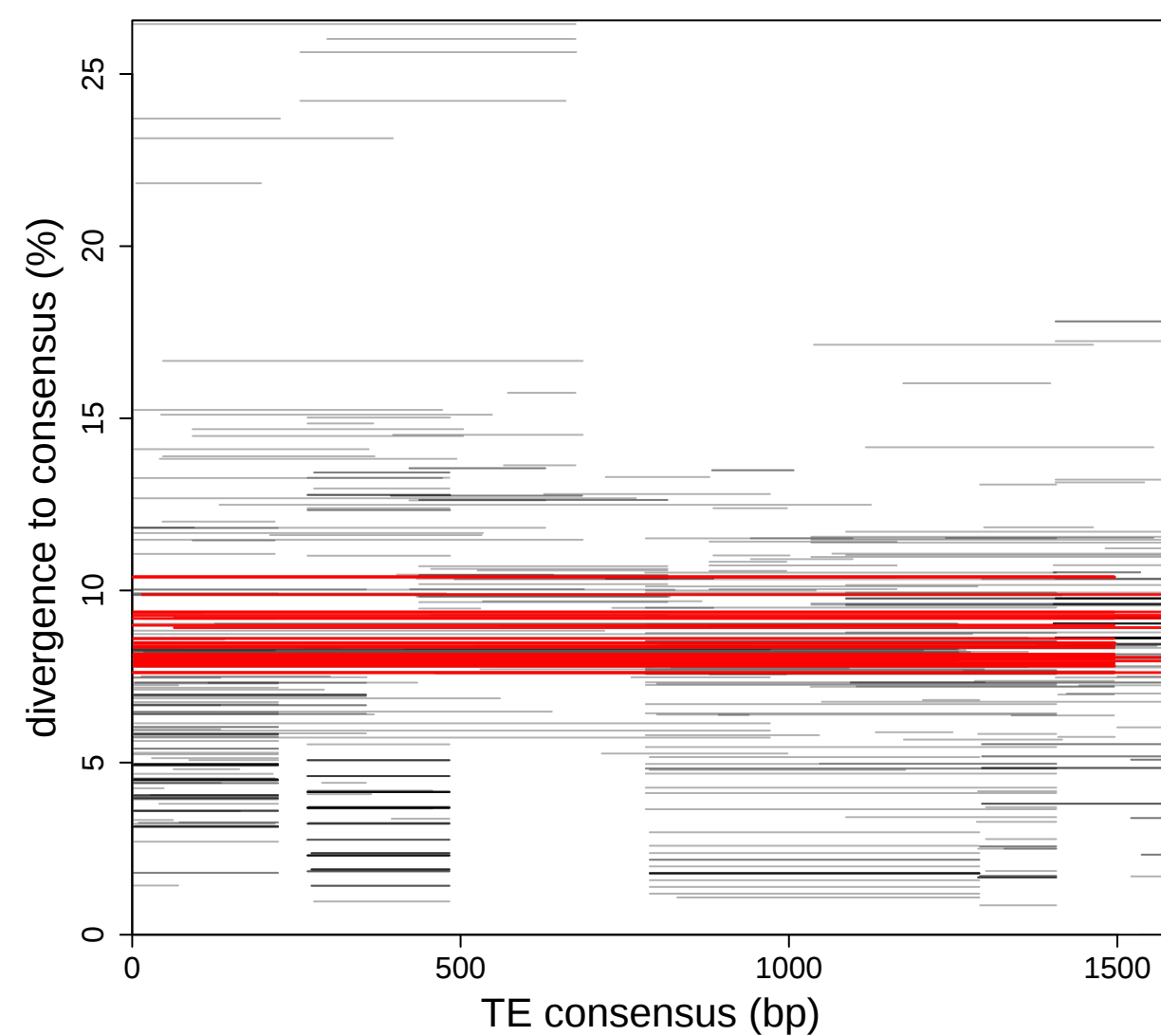


9.fasta.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 1848bp; fragments: 584; full length: 0 (>=1663.2bp)

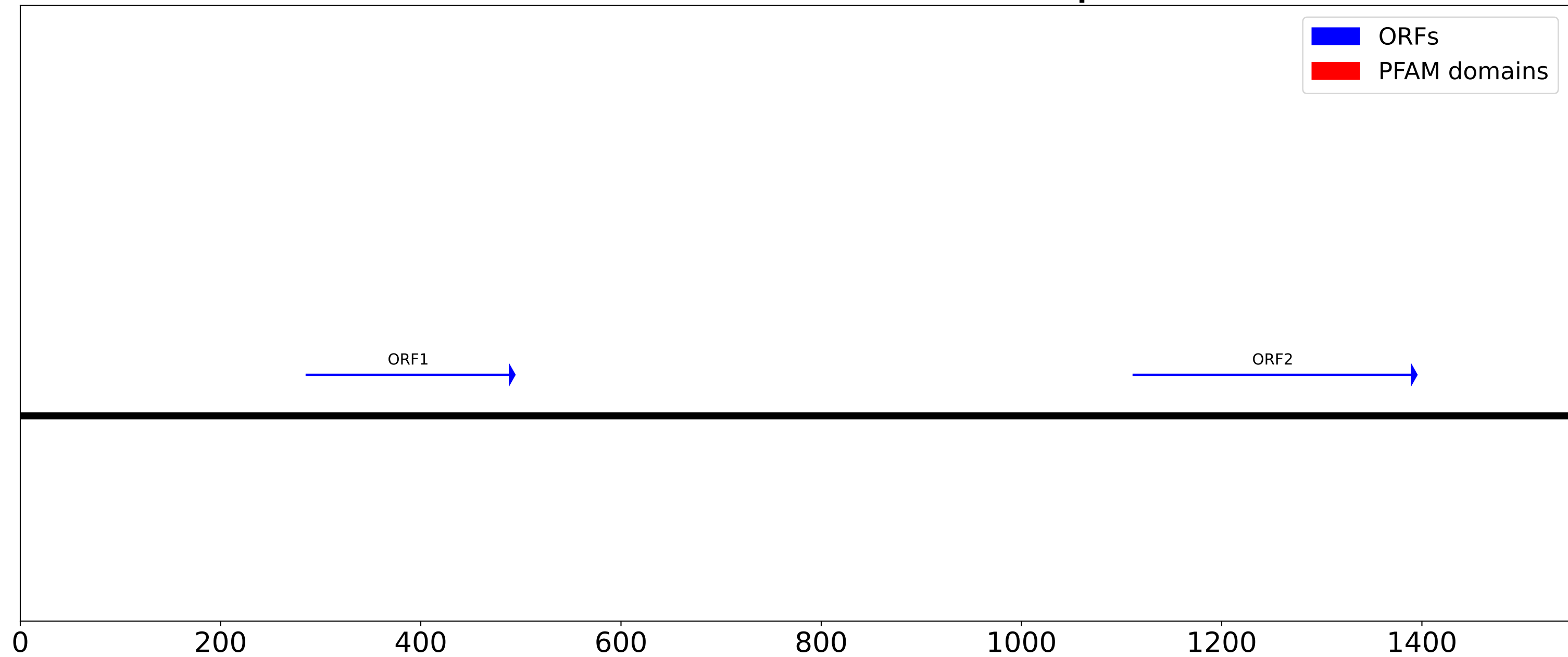
After TEtrimmer Extended plot Blue lines are boundaries



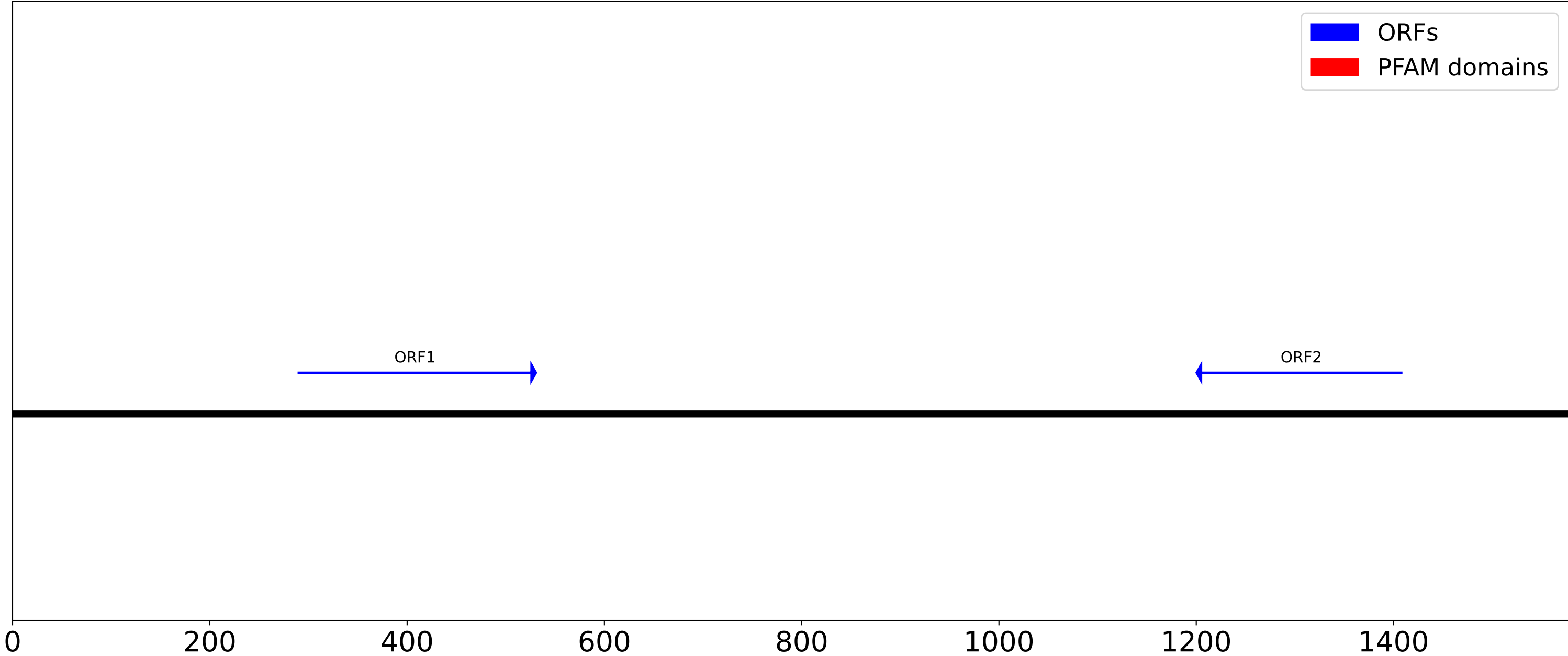
TE: rnd_4_family_2019
size: 1579bp; fragments: 552; full length: 31 (≥ 1421.1 bp)



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

