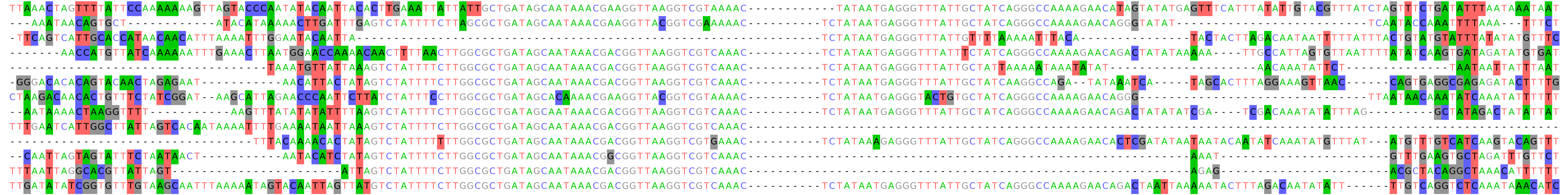


Start crop Point

End crop Point

MSA length = 2174



Start crop Point



End crop Point



0

10

0

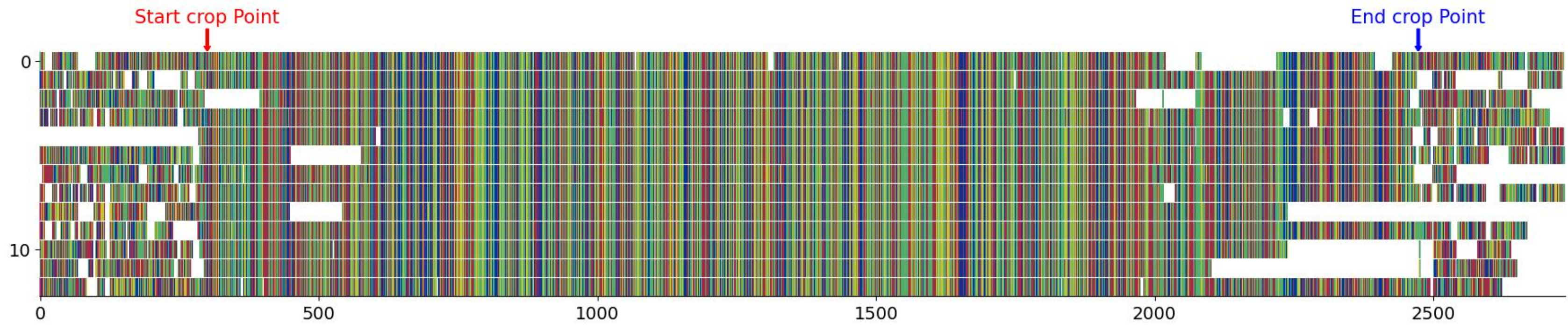
500

1000

1500

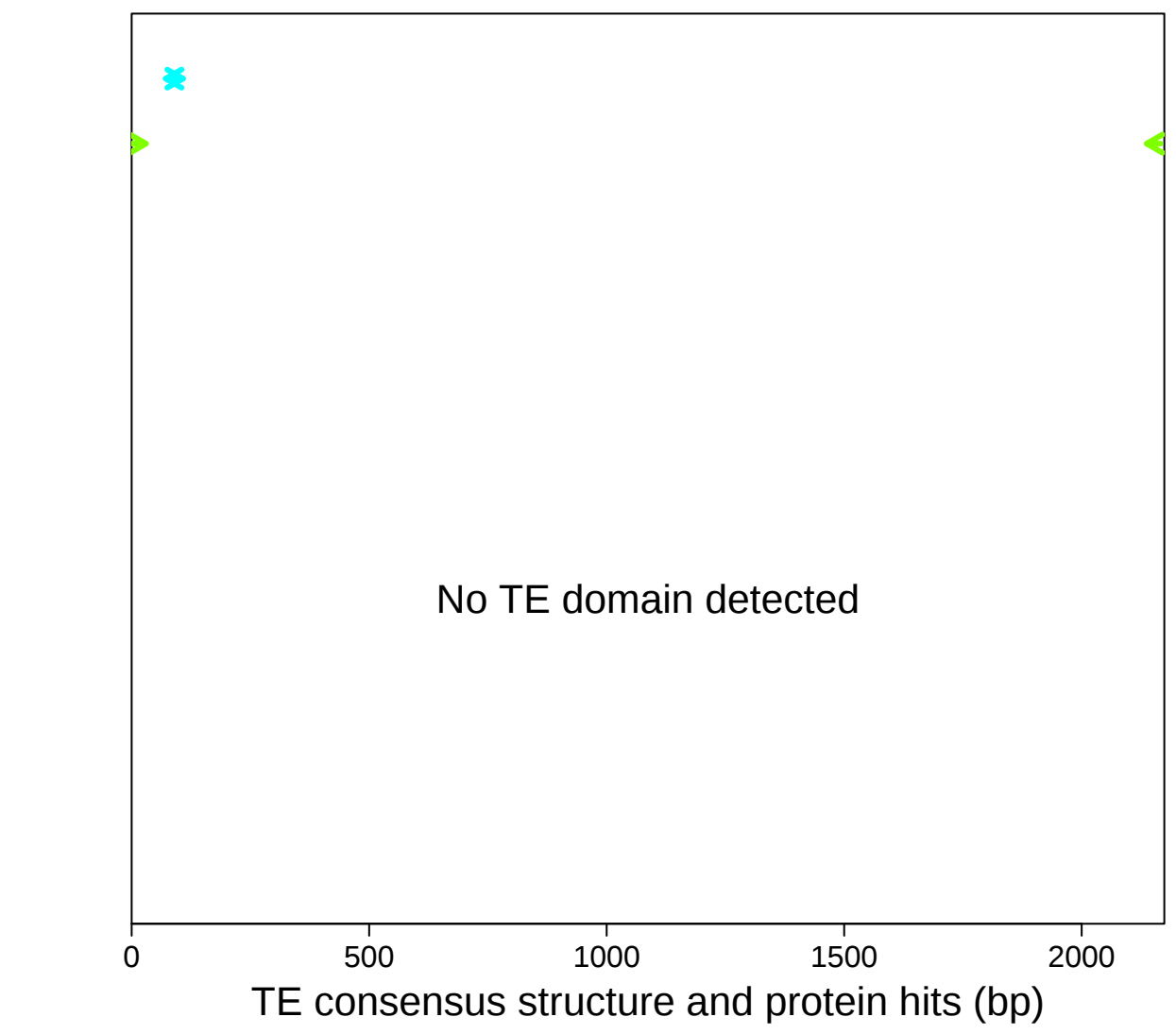
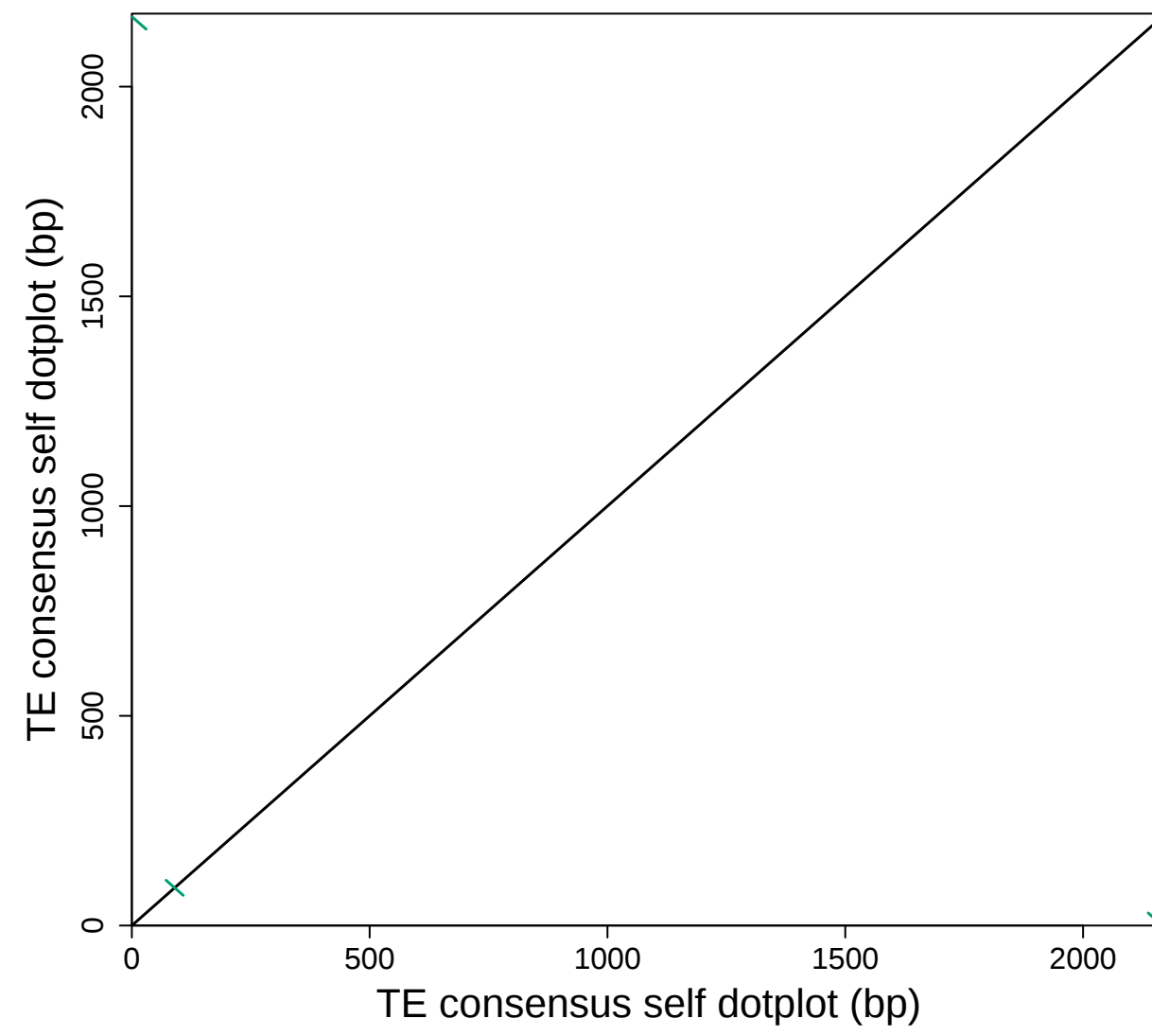
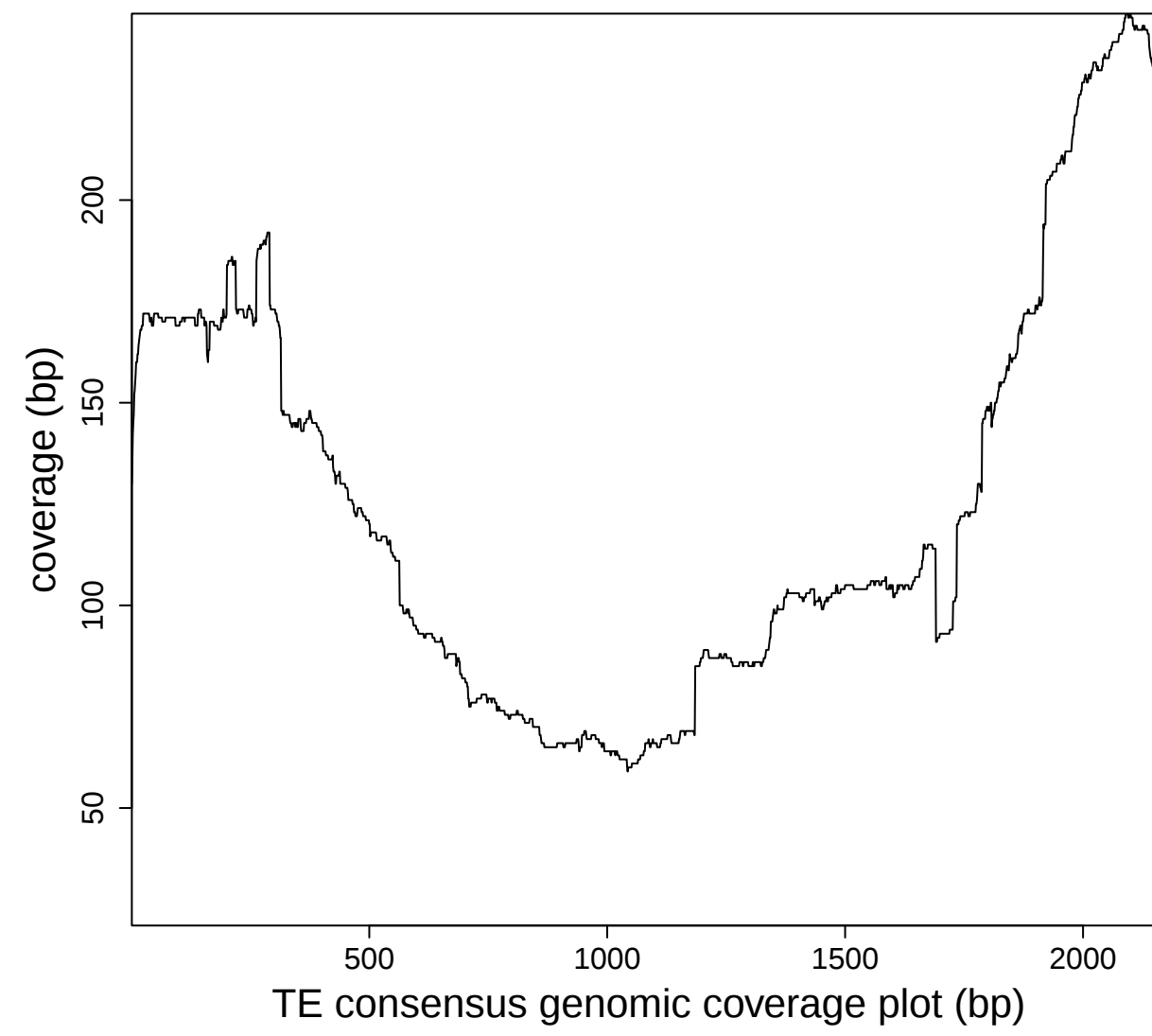
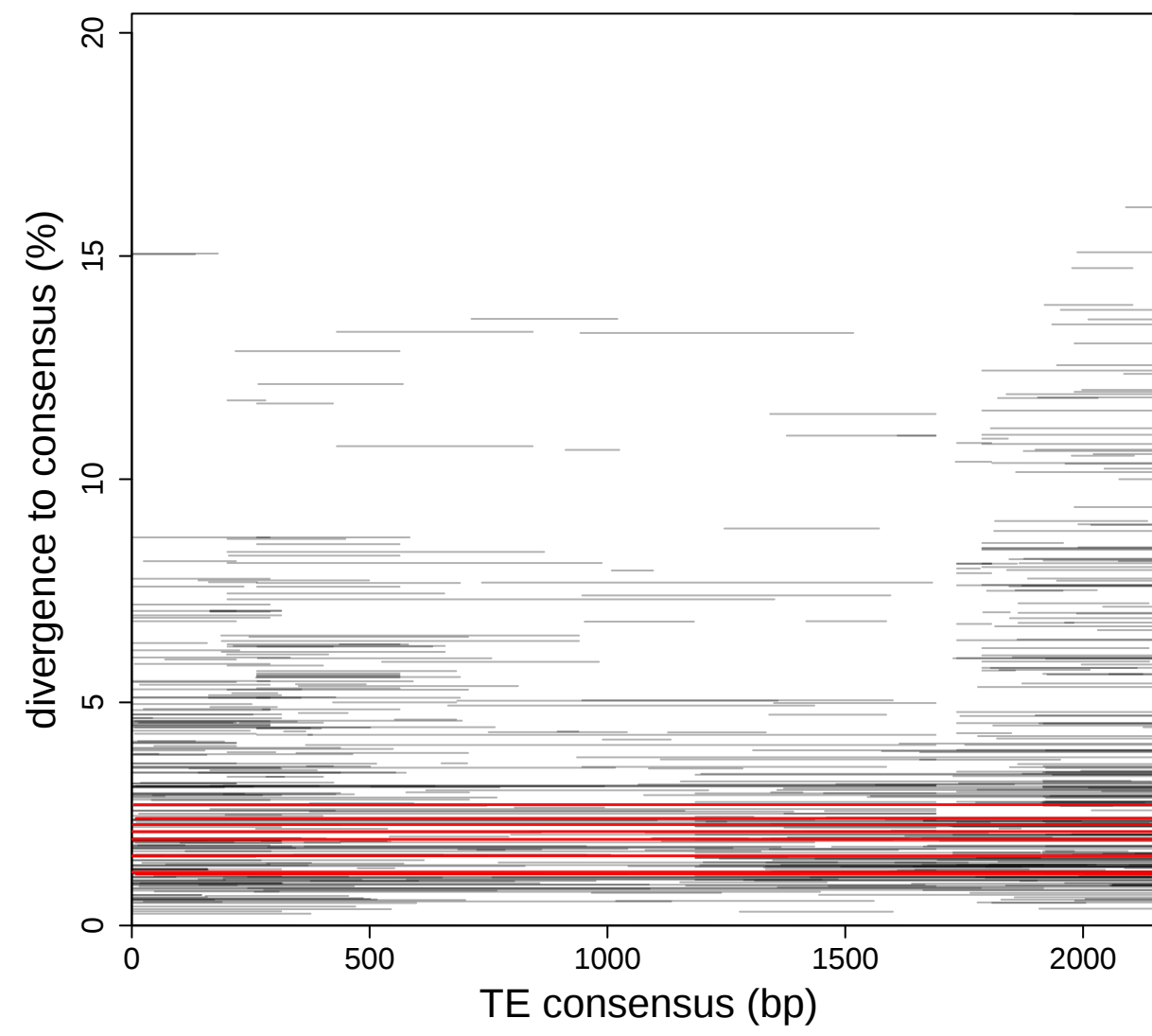
2000

2500



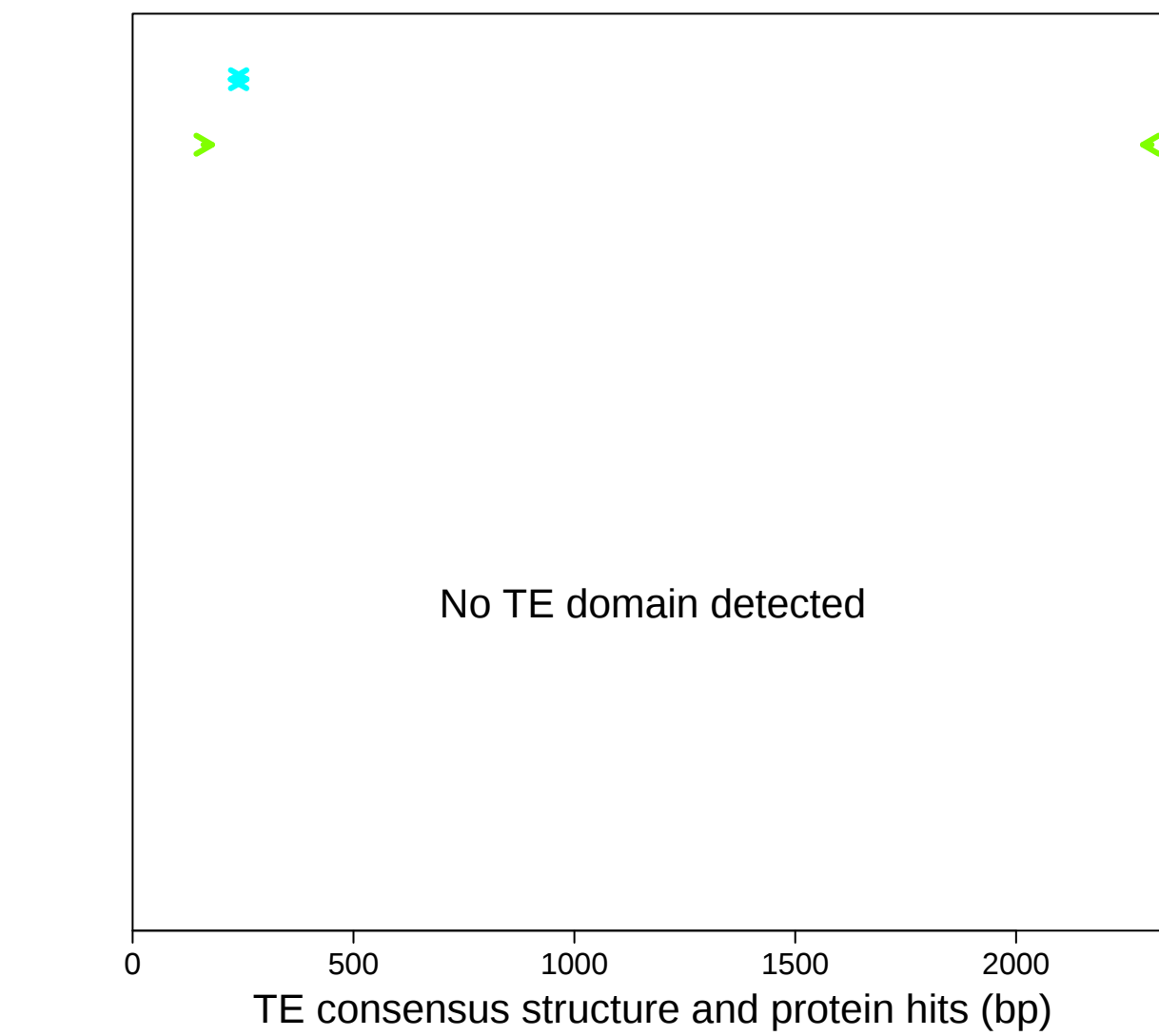
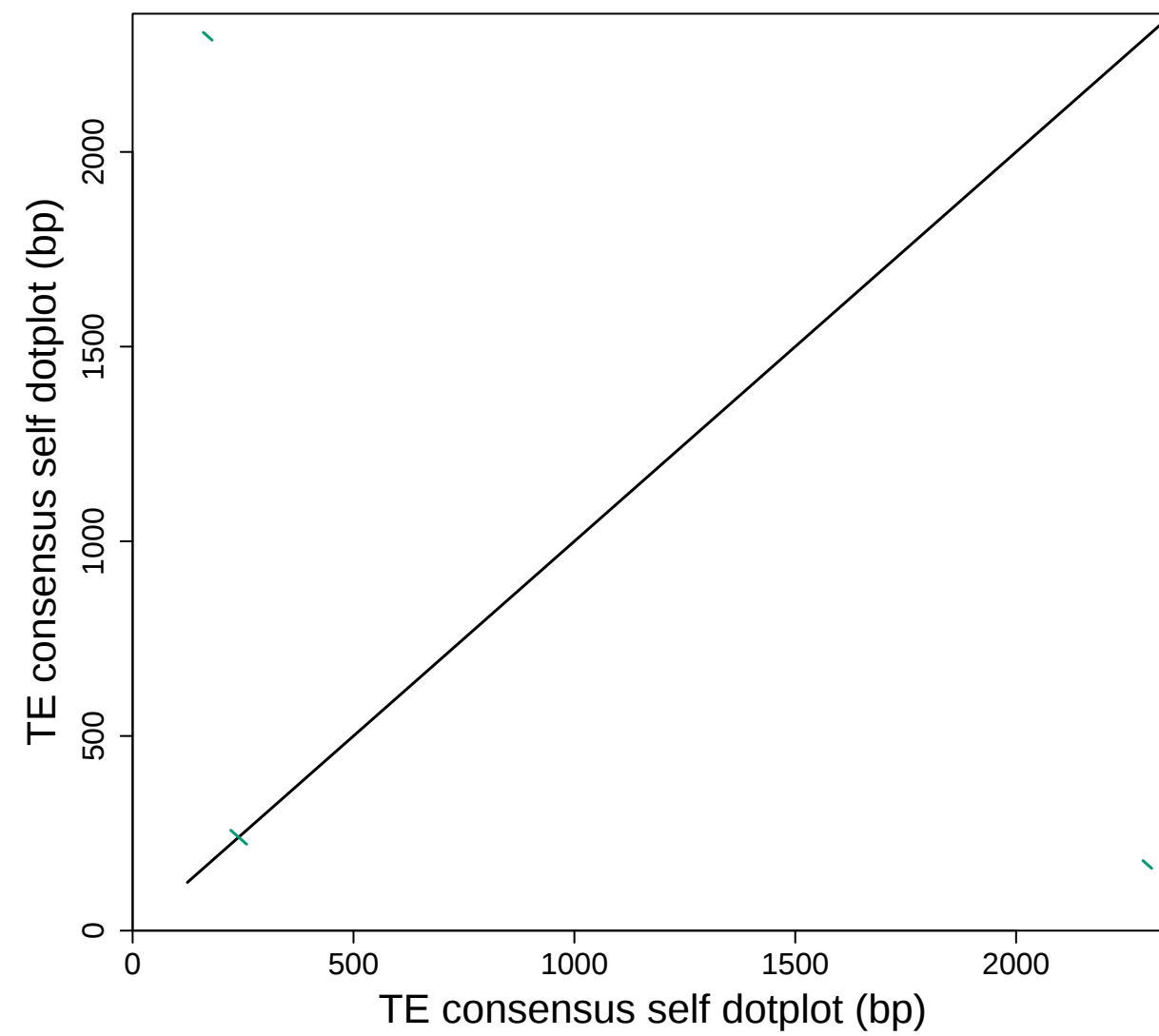
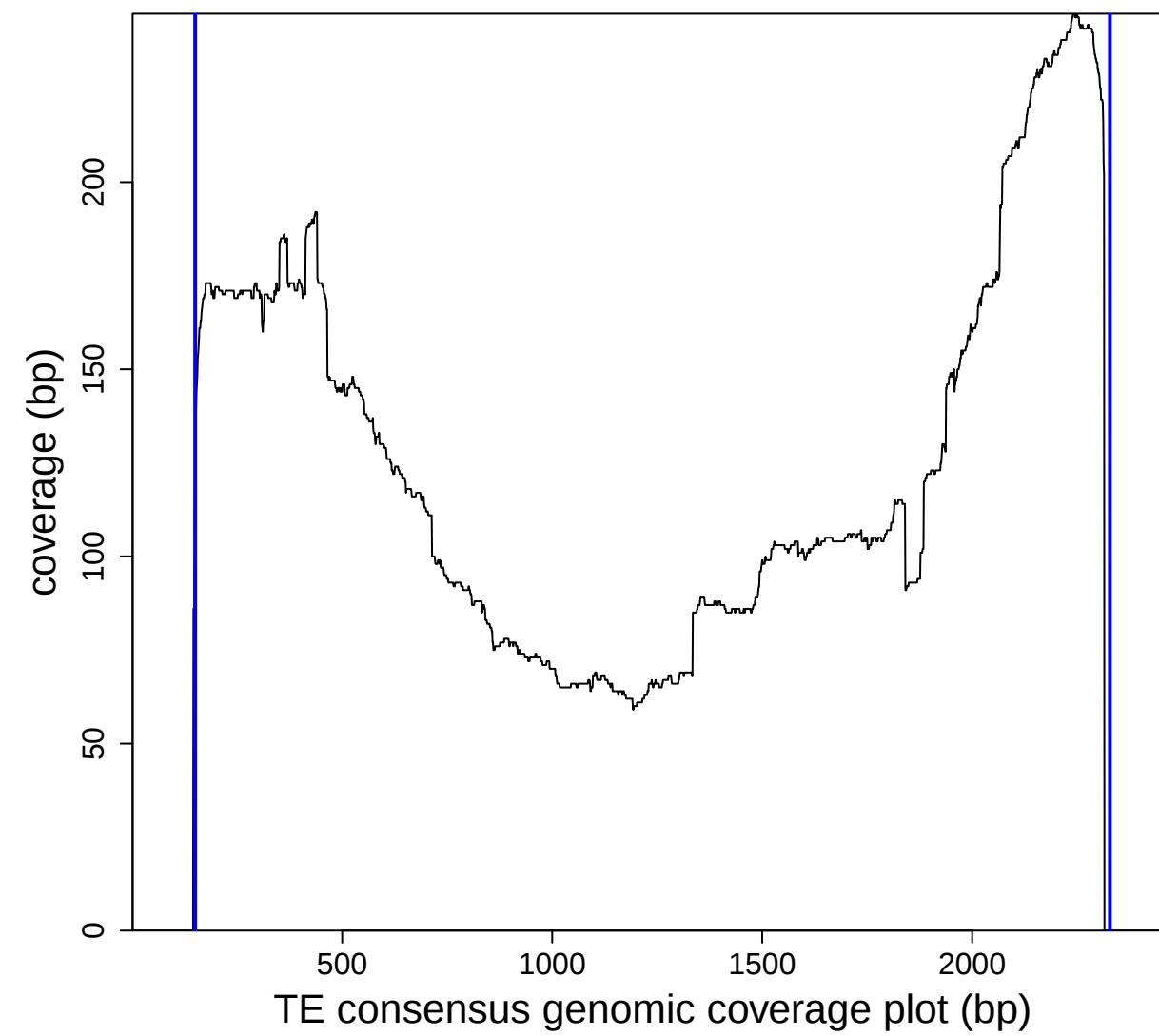
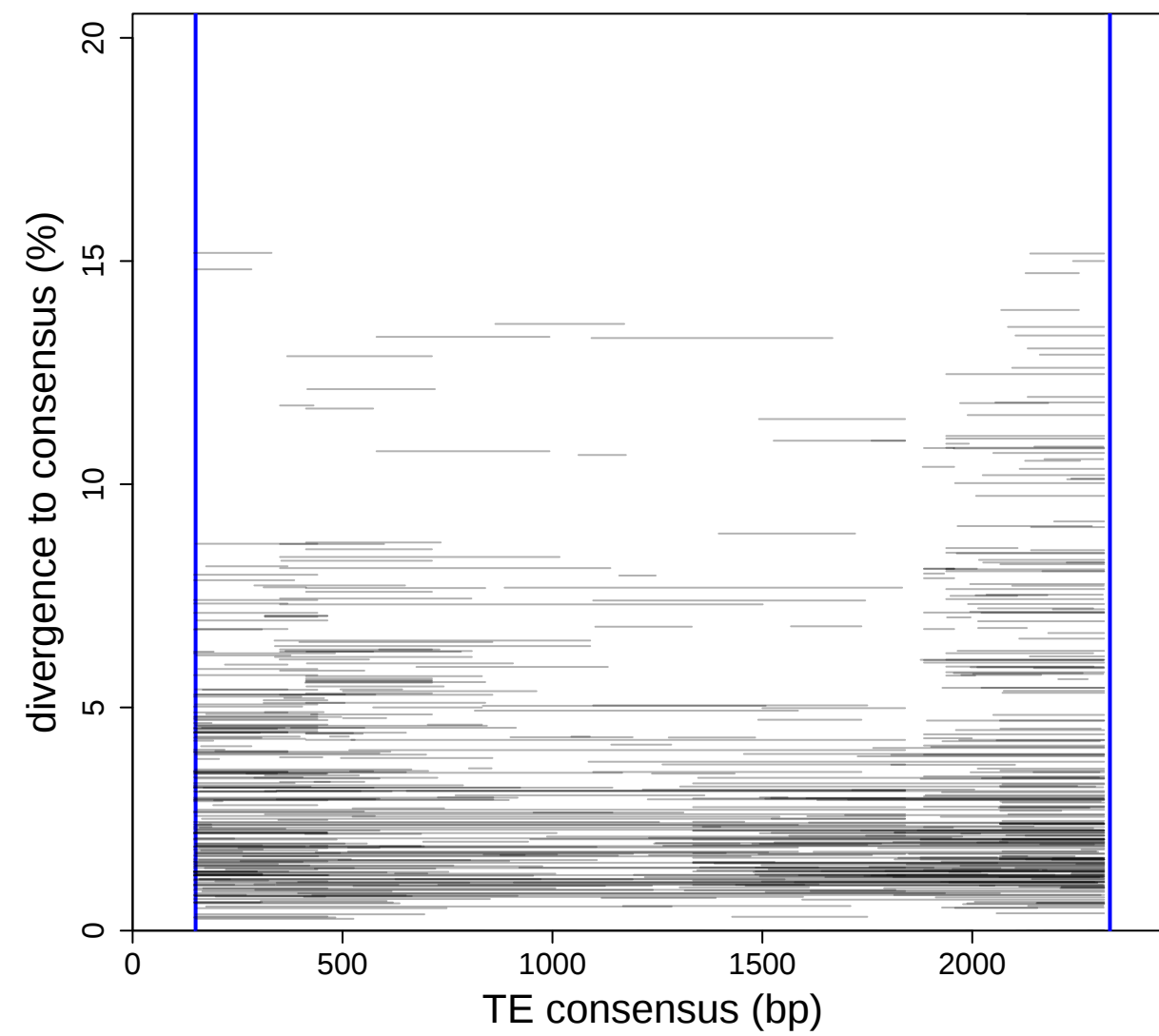
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 2174bp; fragments: 621; full length: 8 (>=1956.6bp)

After TEtrimmer 2174 bp



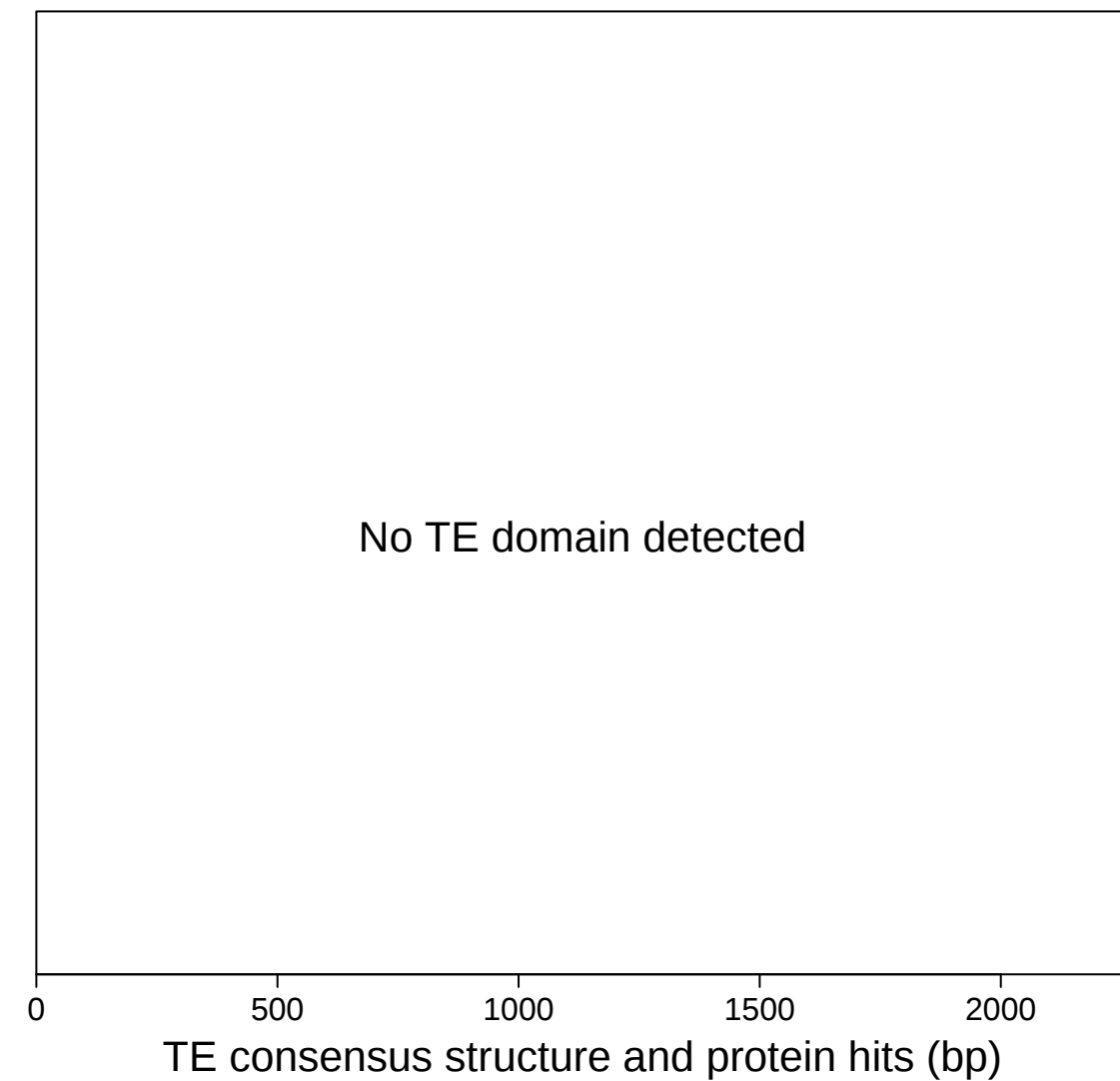
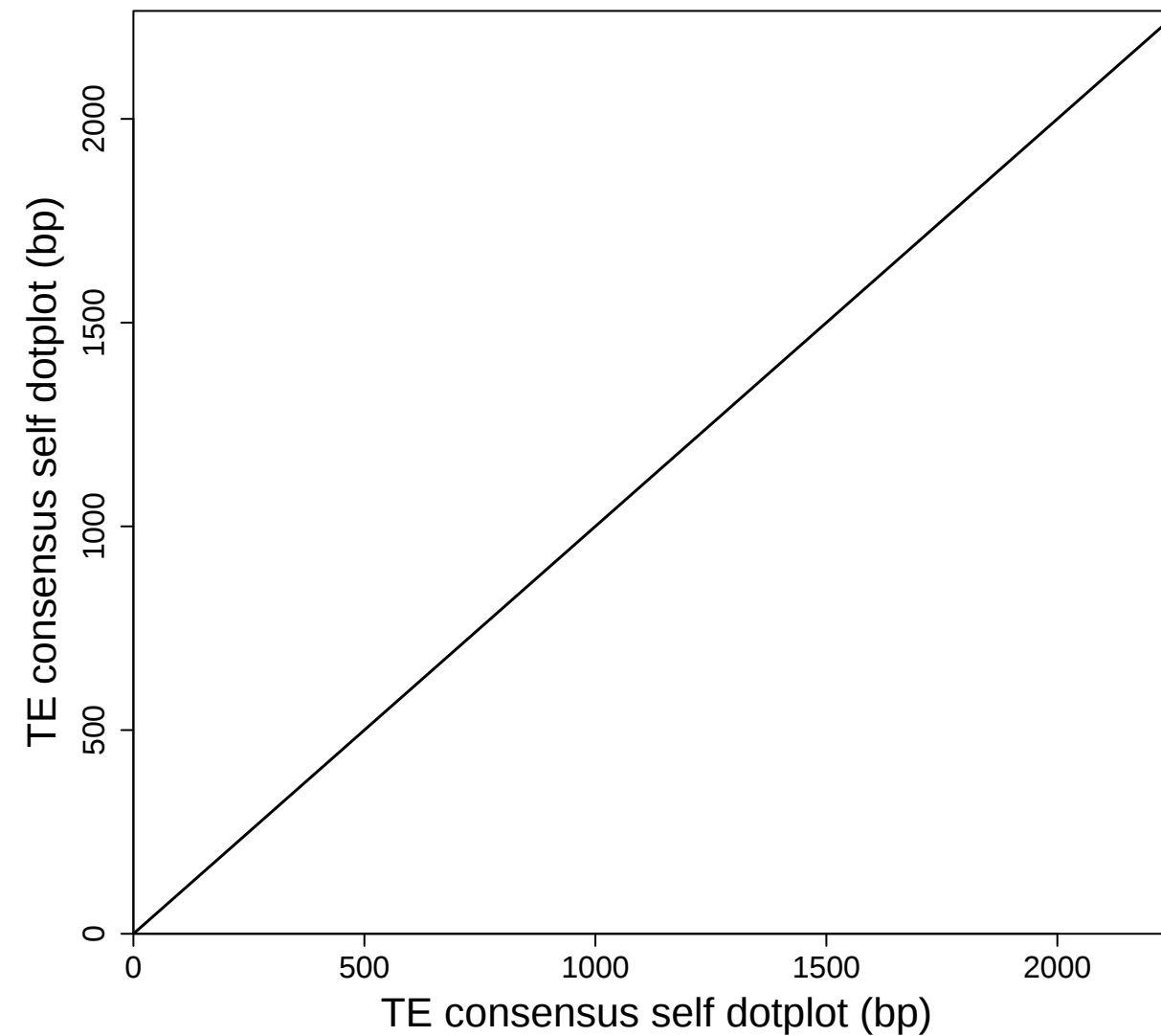
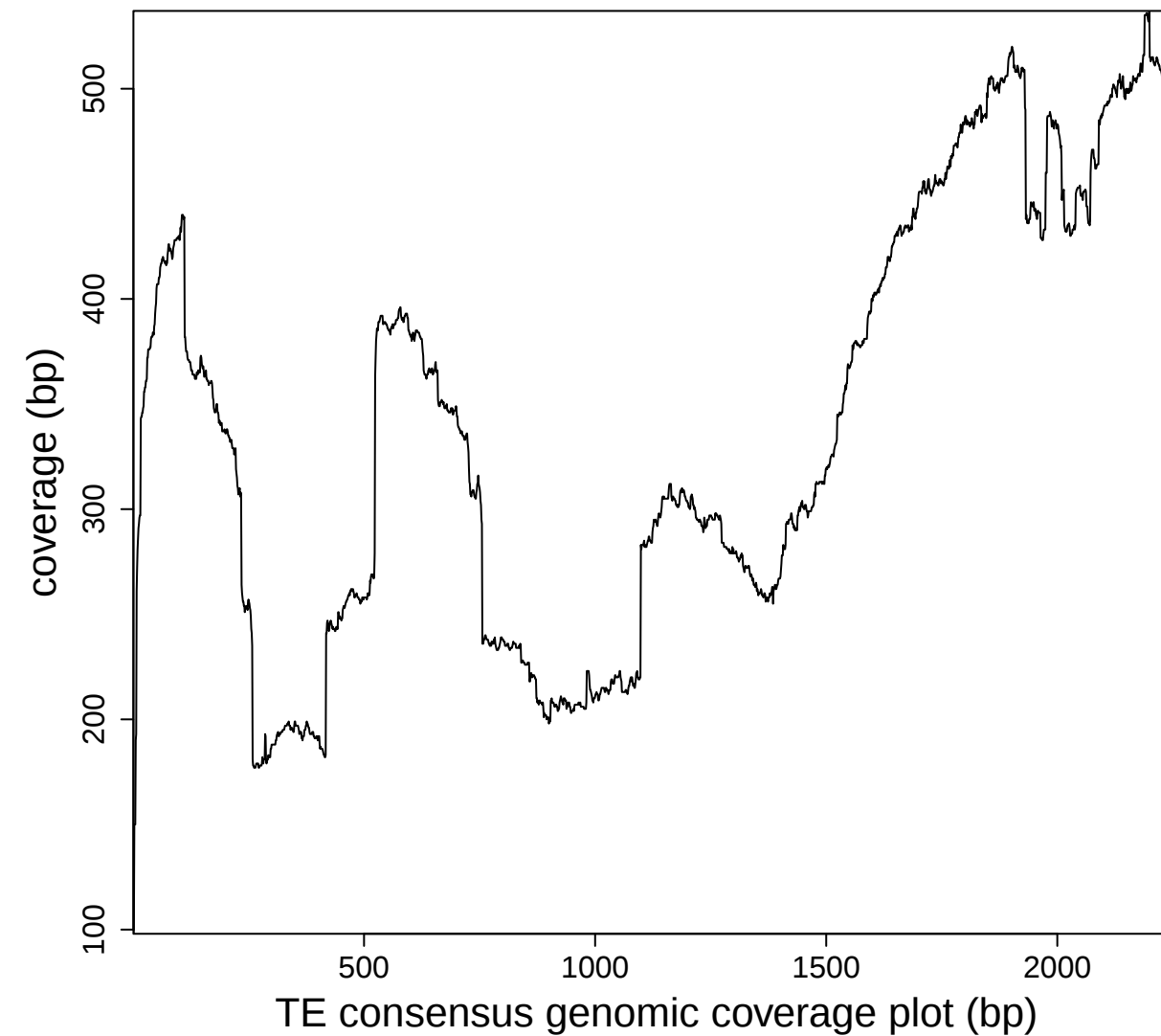
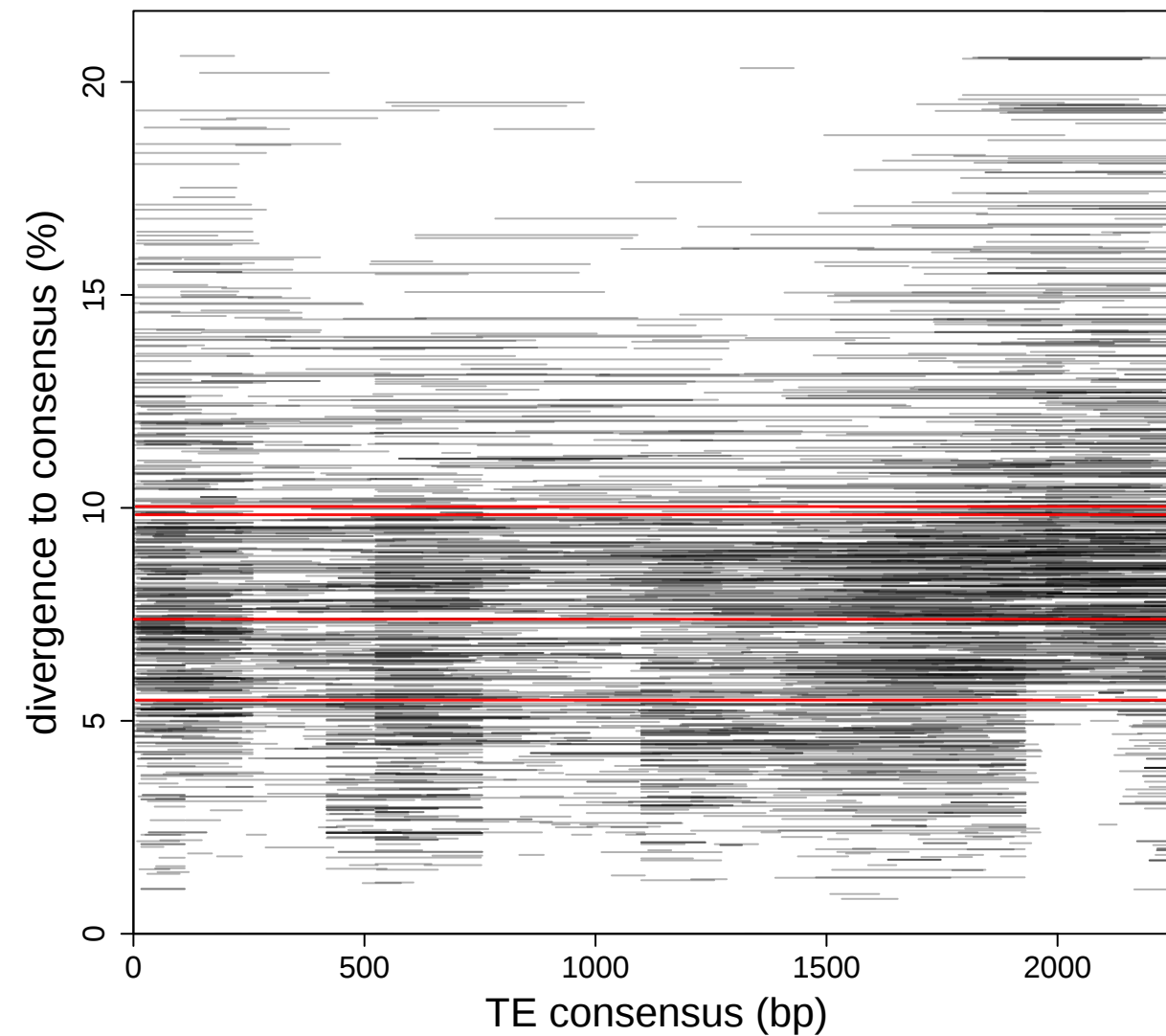
5.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 2478bp; fragments: 621; full length: 0 (>=2230.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

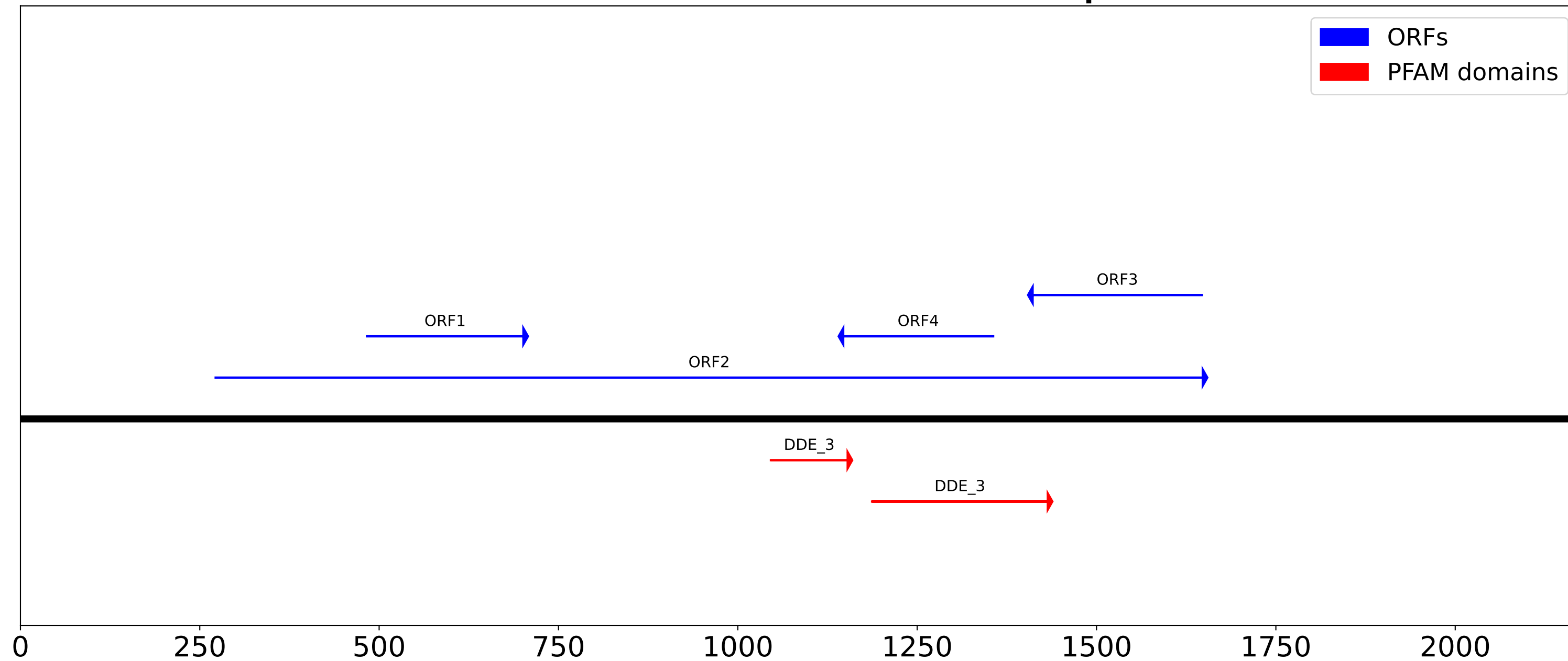


TE: rnd_4_family_1445
size: 2265bp; fragments: 2413; full length: 4 (≥ 2038.5 bp)

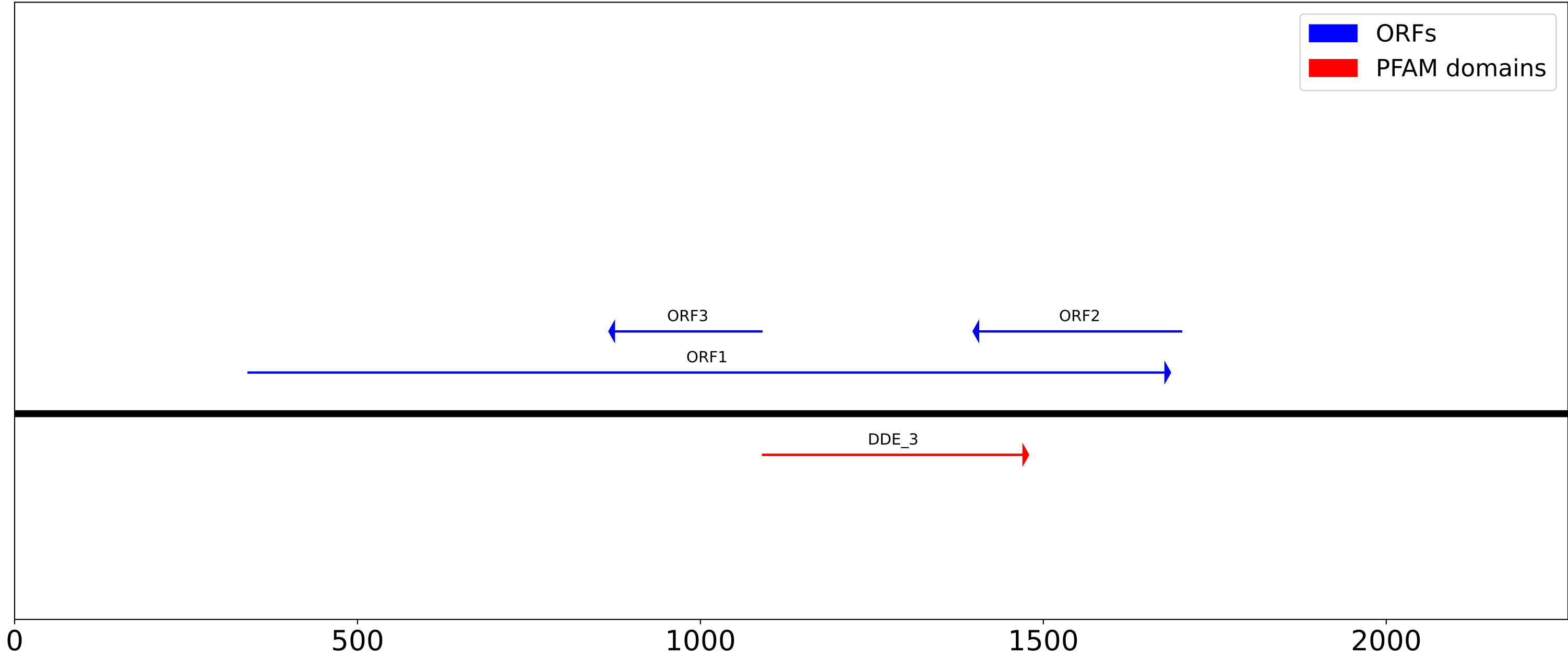
Before TEtrimmer 2265 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

