

MSA length = 4393

Start crop Point

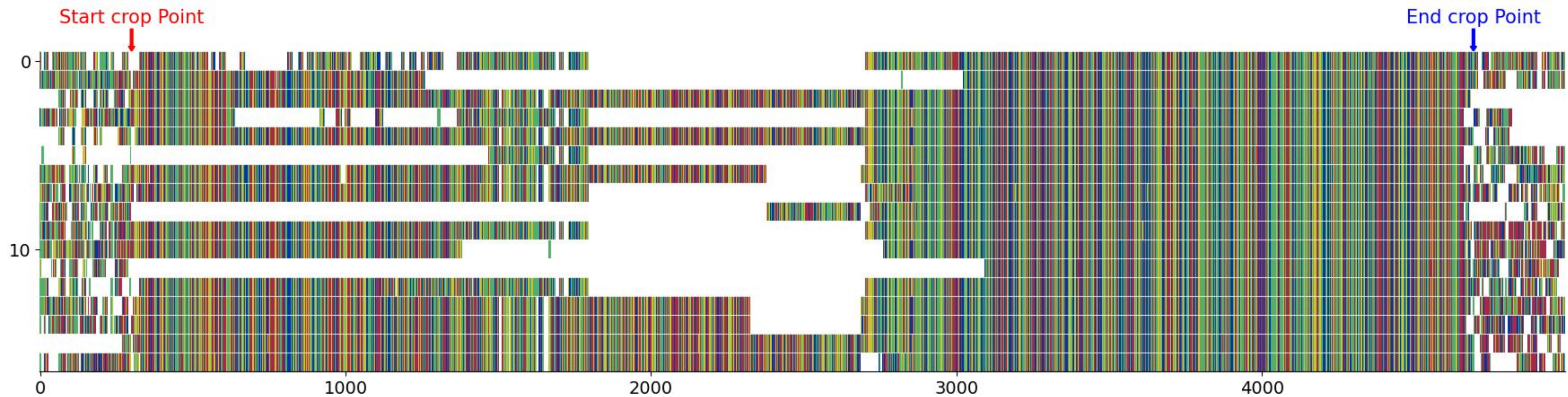


End crop Point



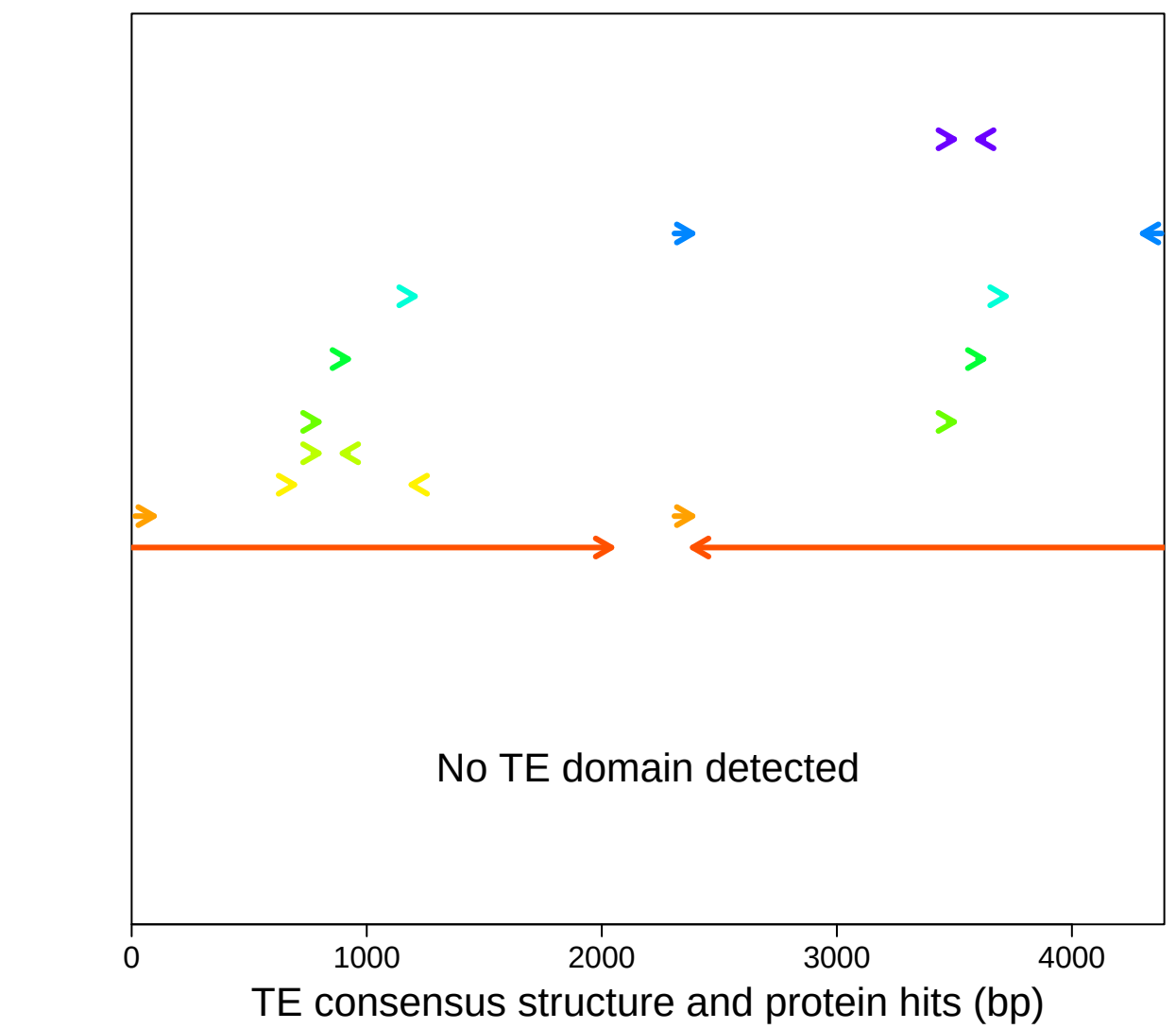
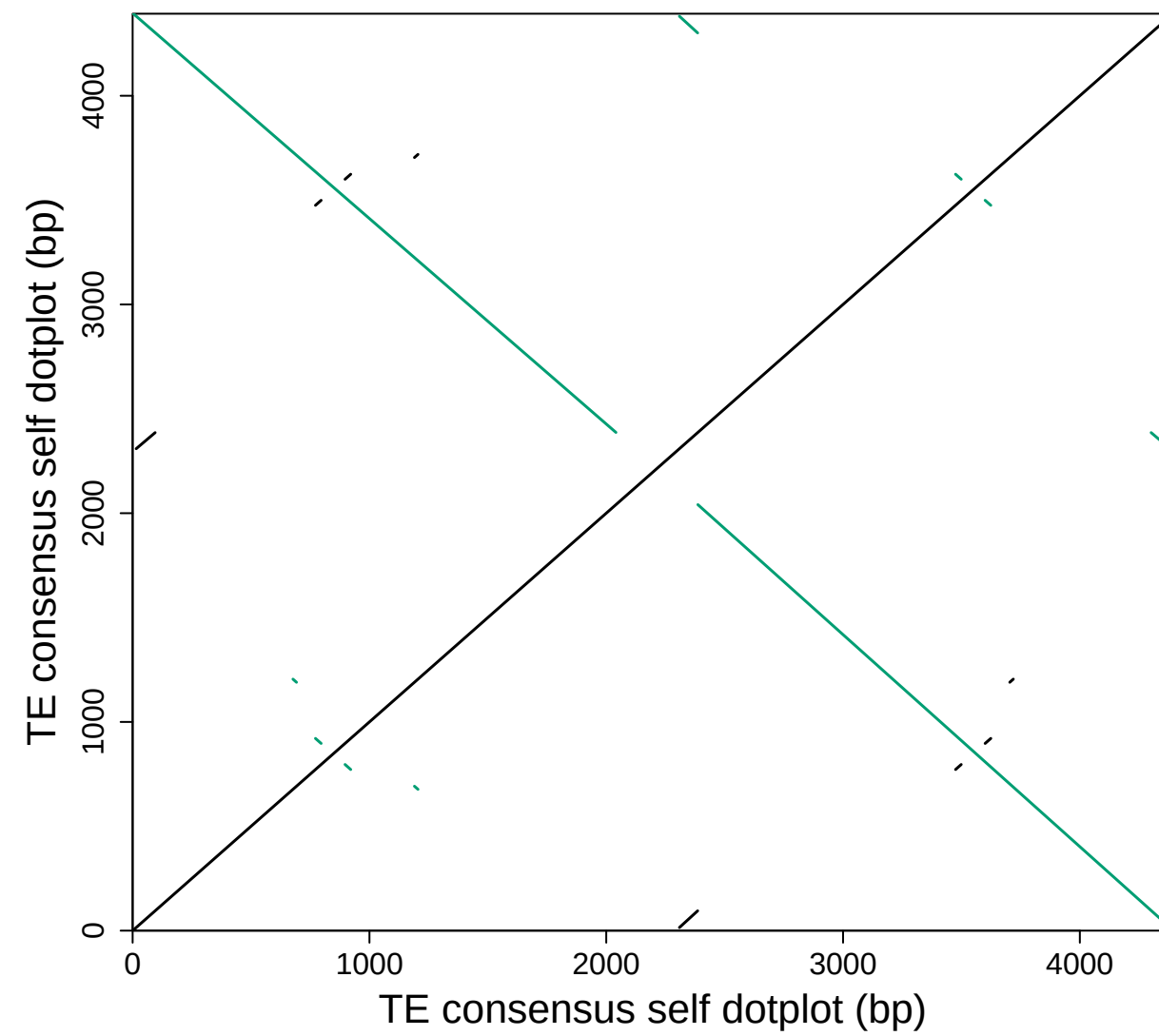
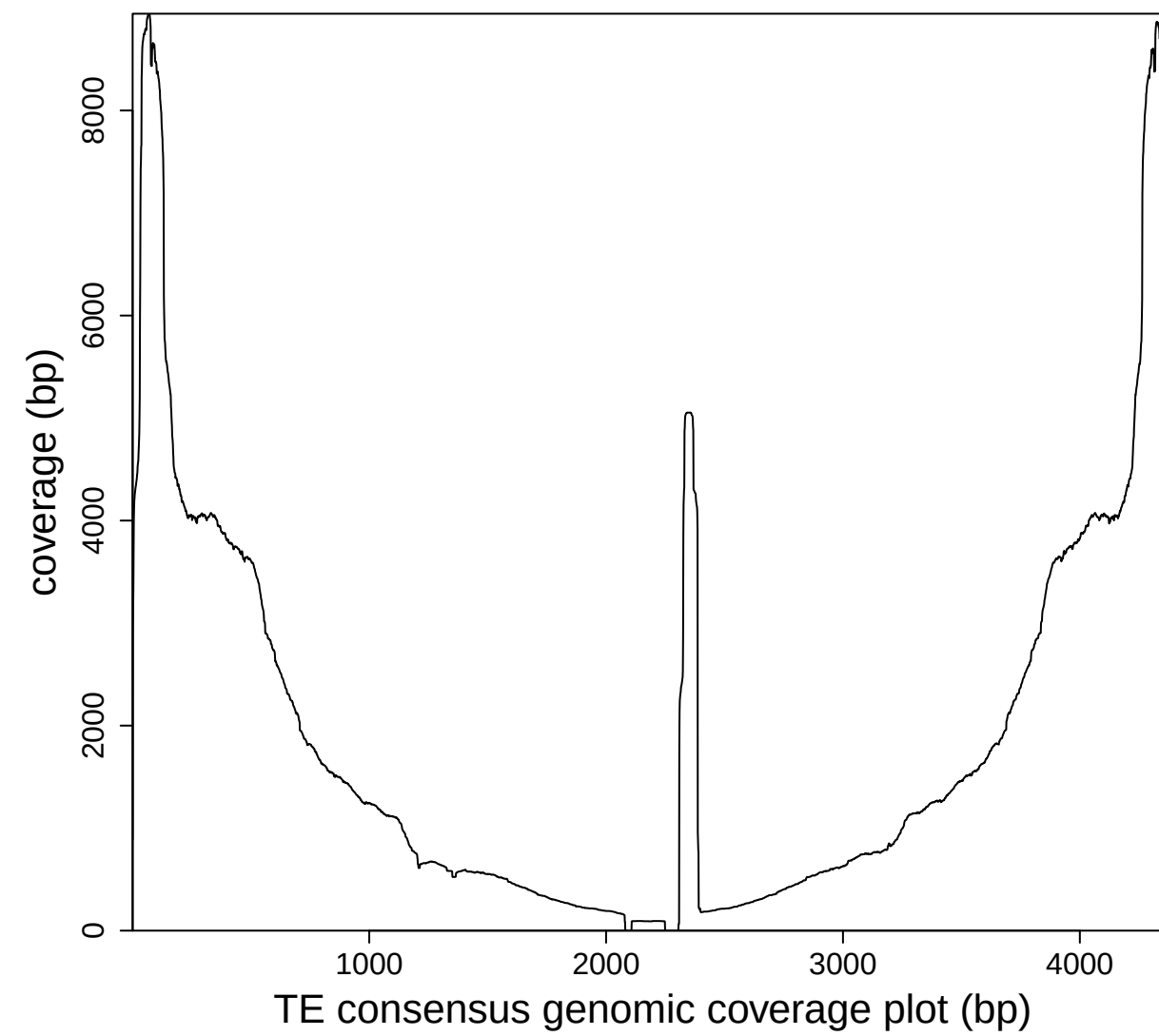
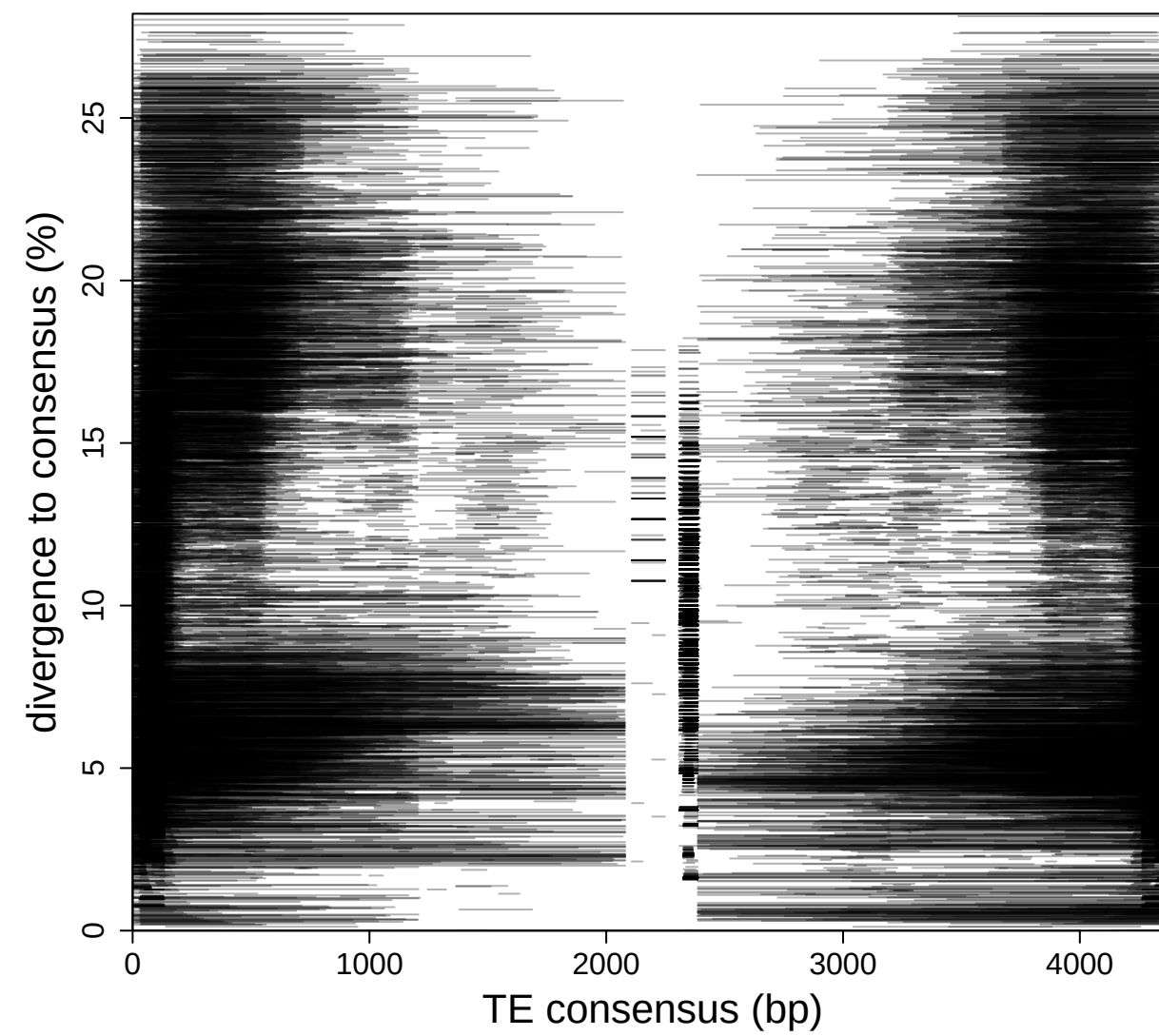
Start crop Point

End crop Point



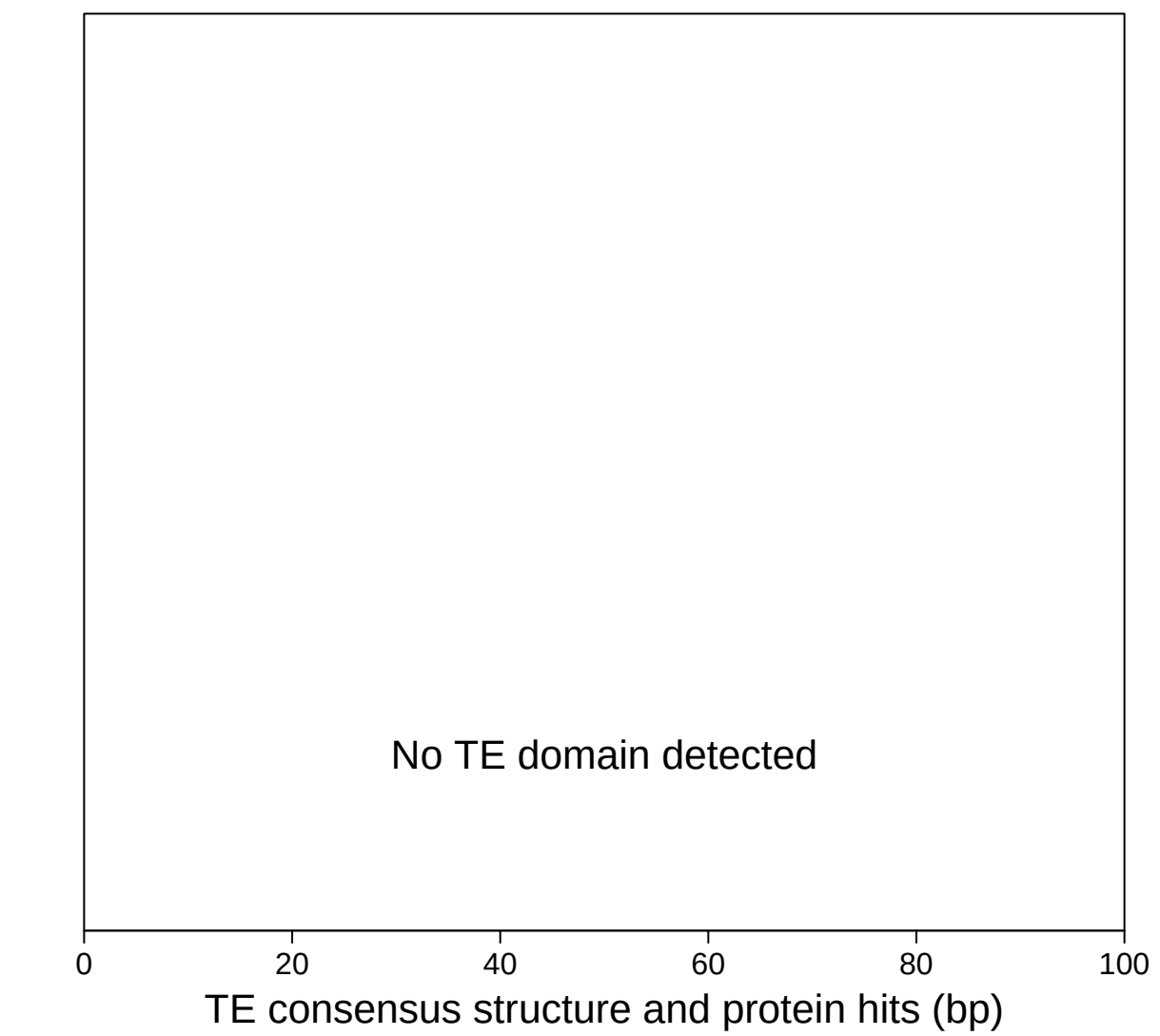
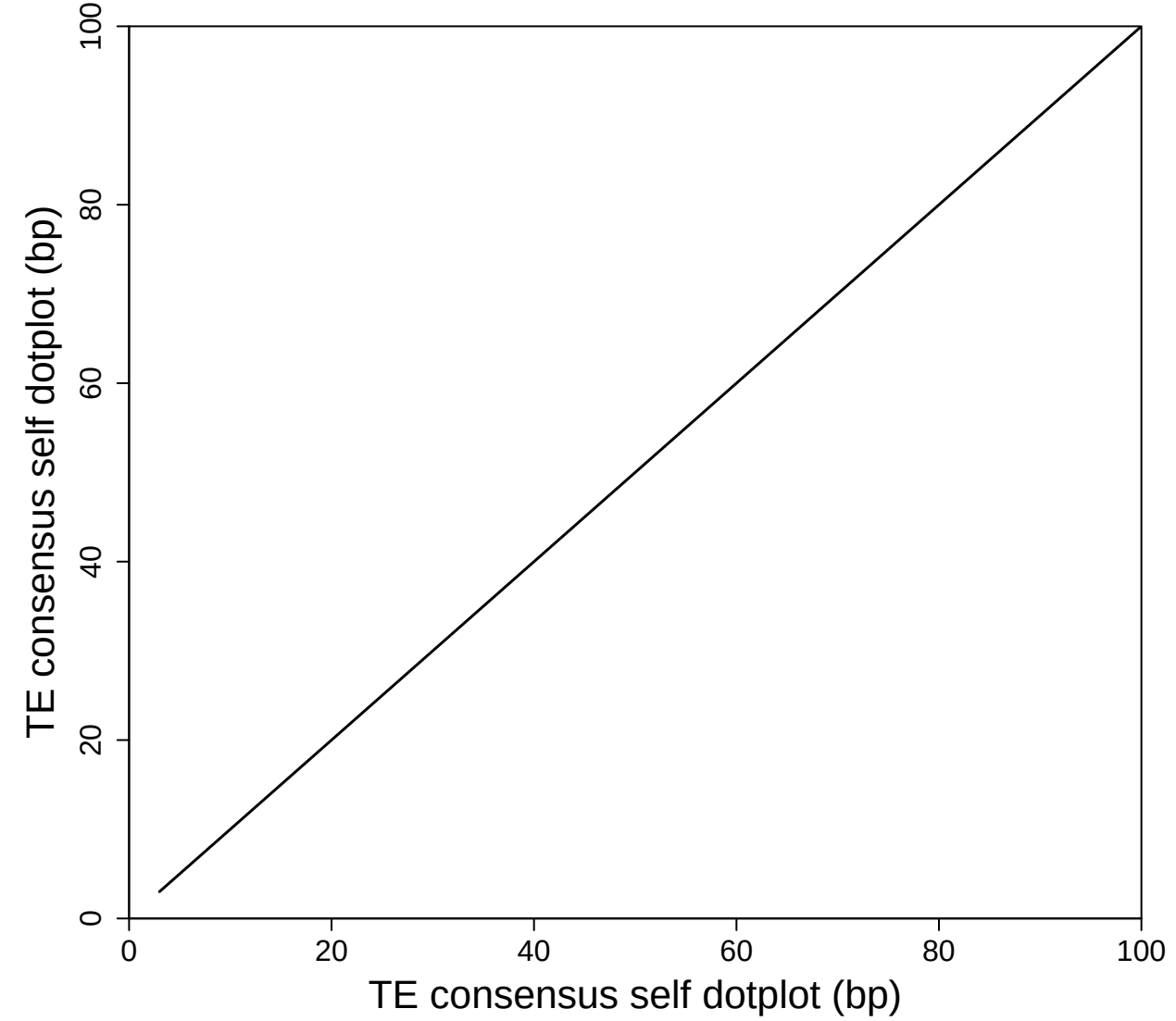
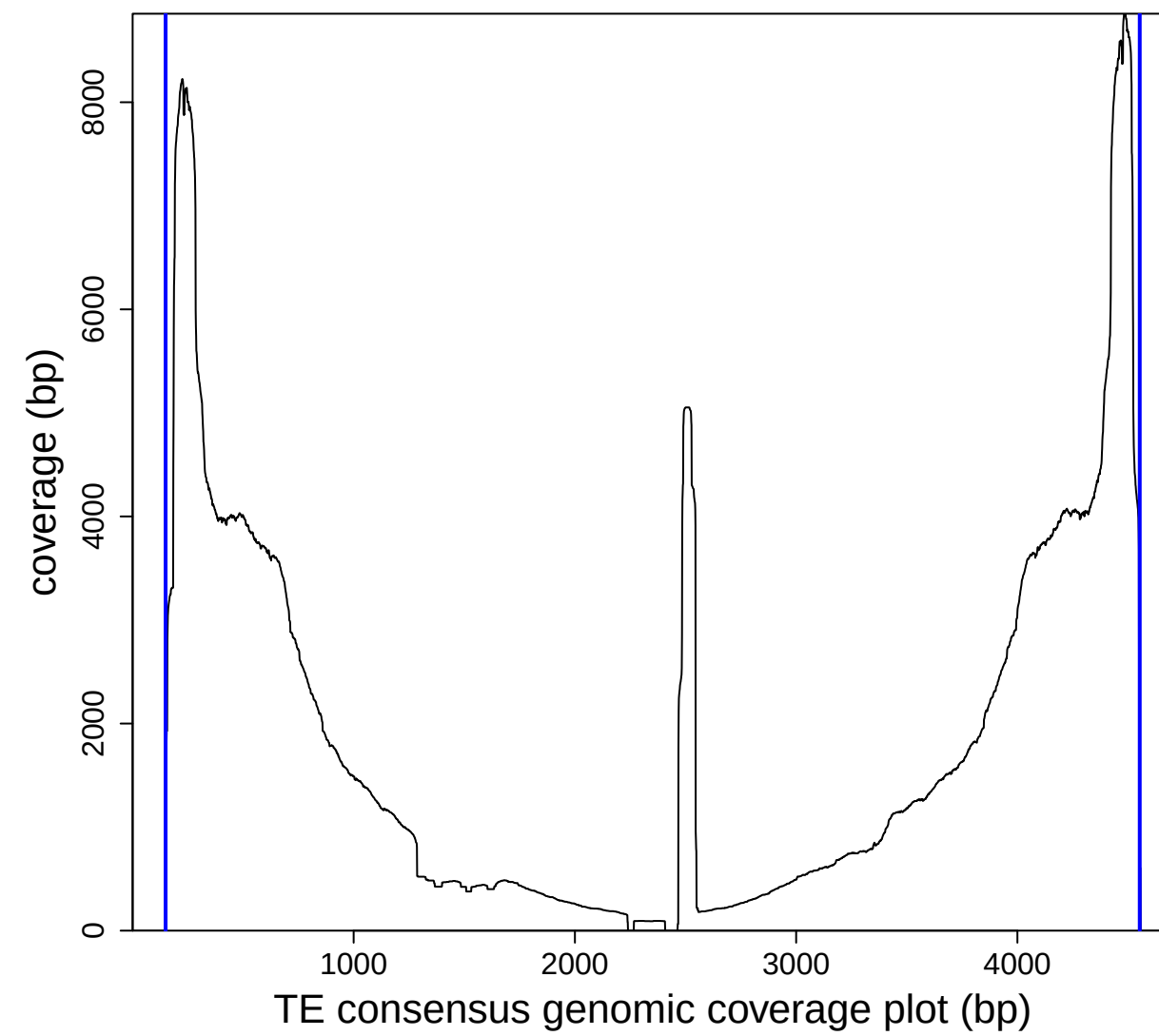
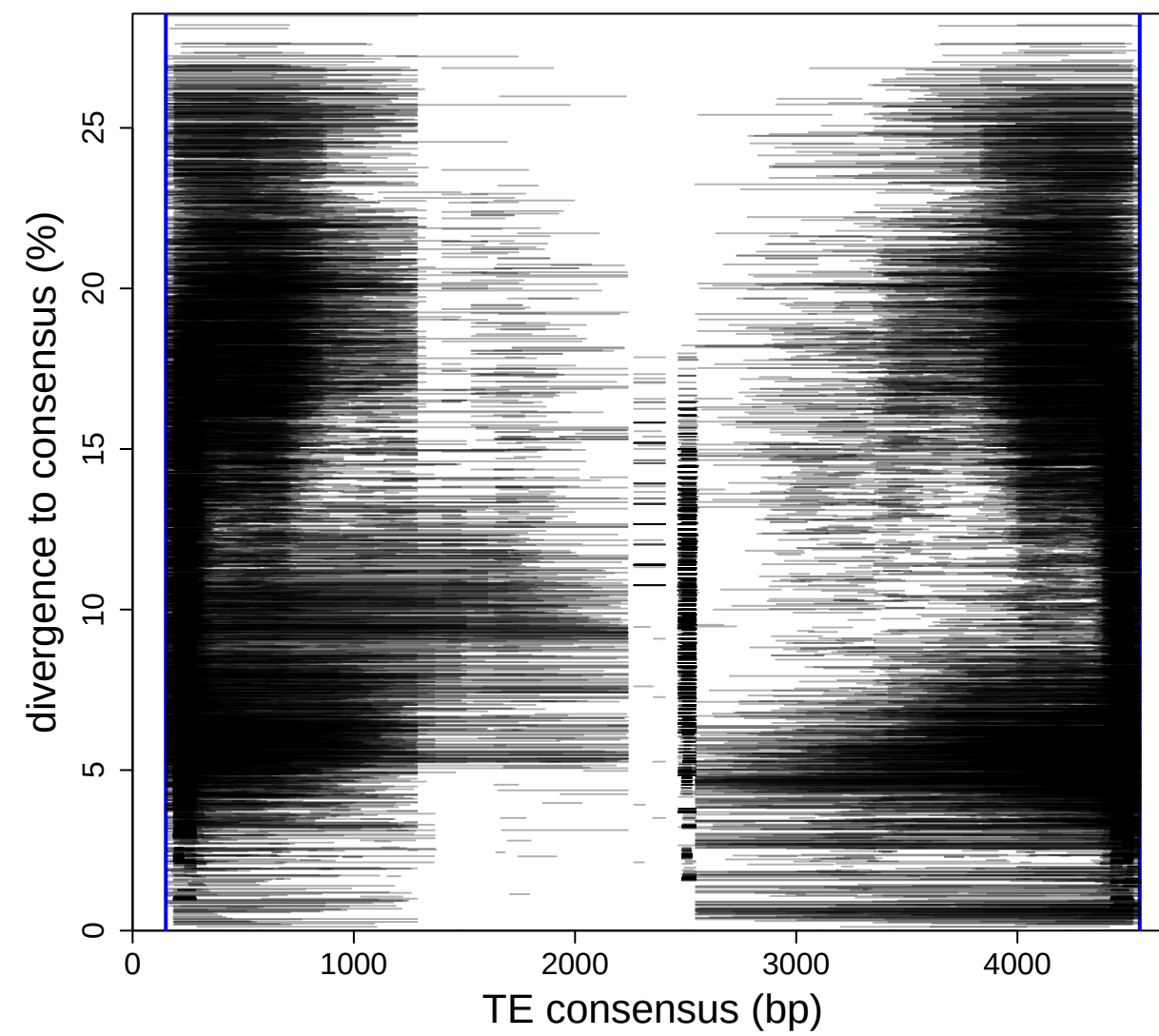
bed_uf.bed g_3.bed fm_1.bed 0_0 bcln.fa aln.fa cl.fa gs.fa c
size: 4393bp; fragments: 31223; full length: 0 (>=3953.7bp)

After TEtrimmer 4393 bp



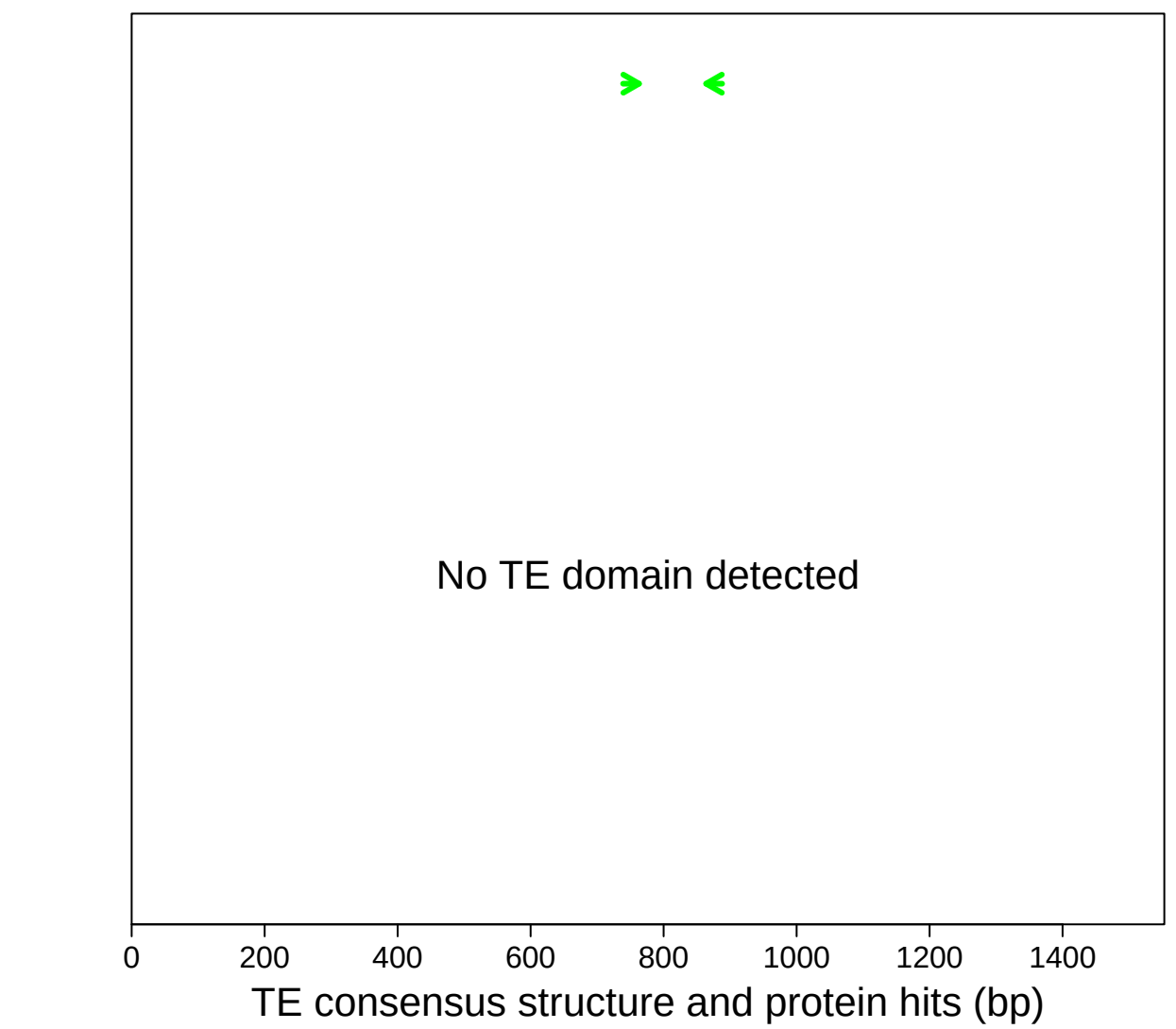
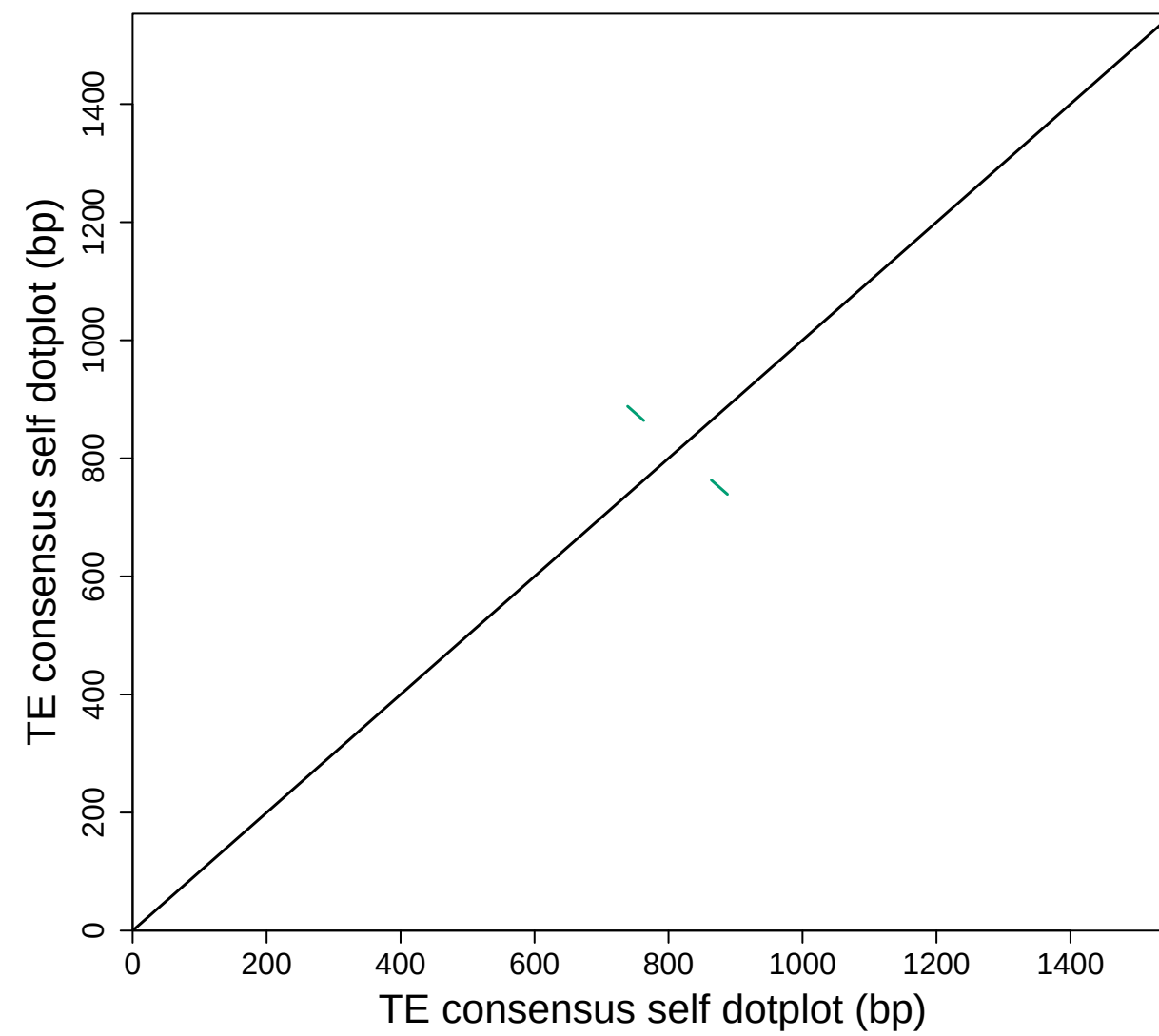
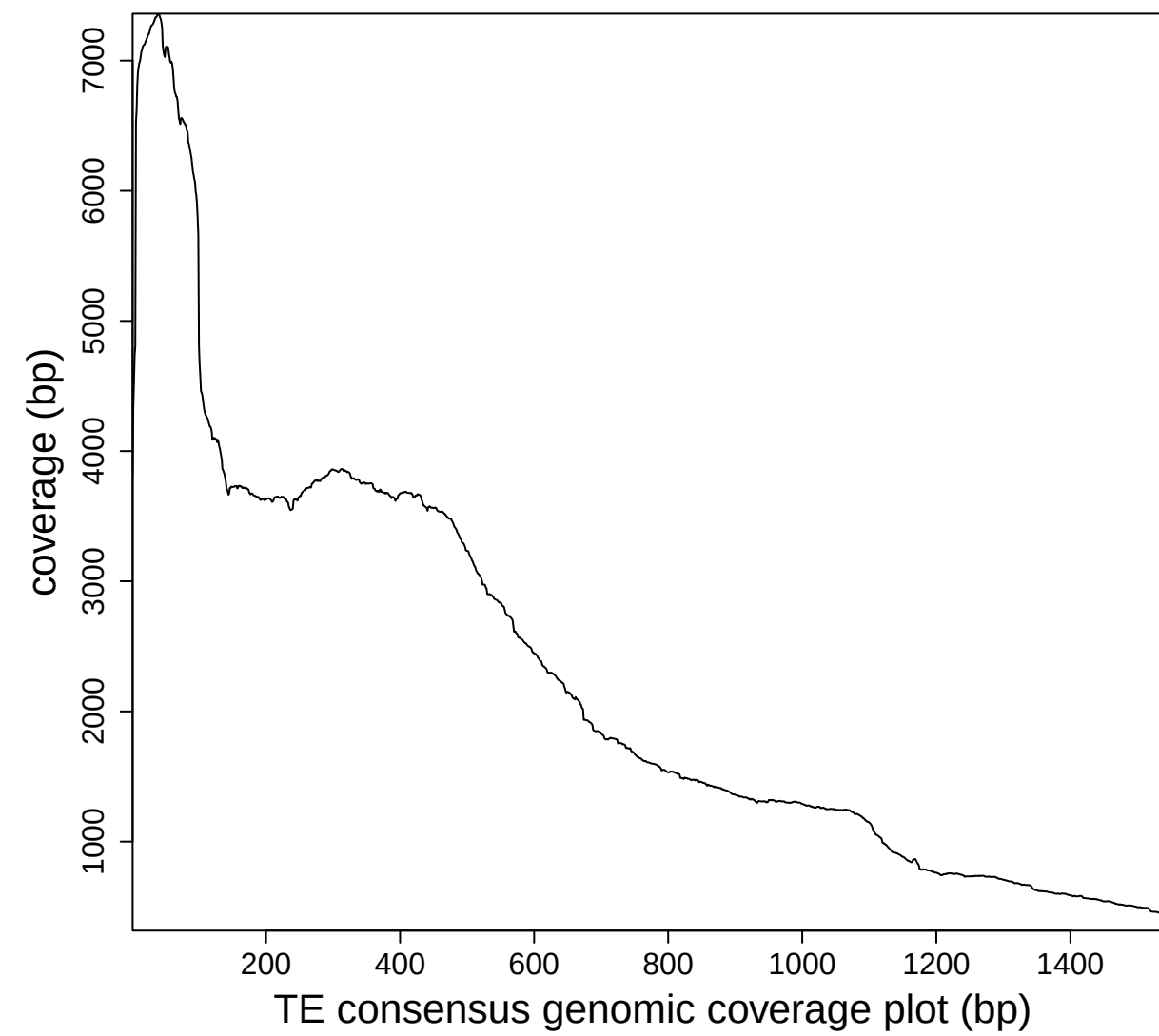
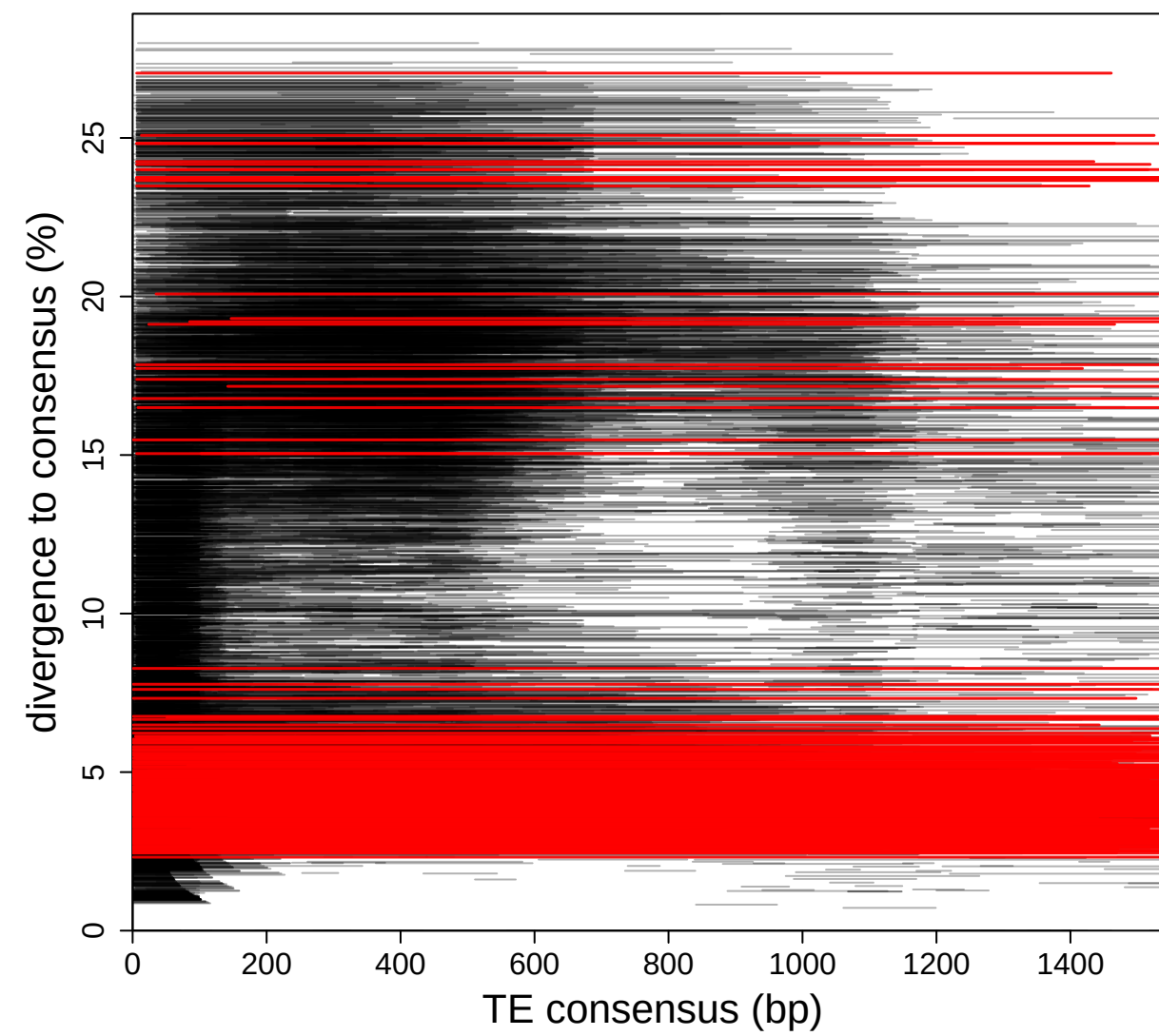
asta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_c
size: 4703bp; fragments: 30018; full length: 0 (>=4232.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

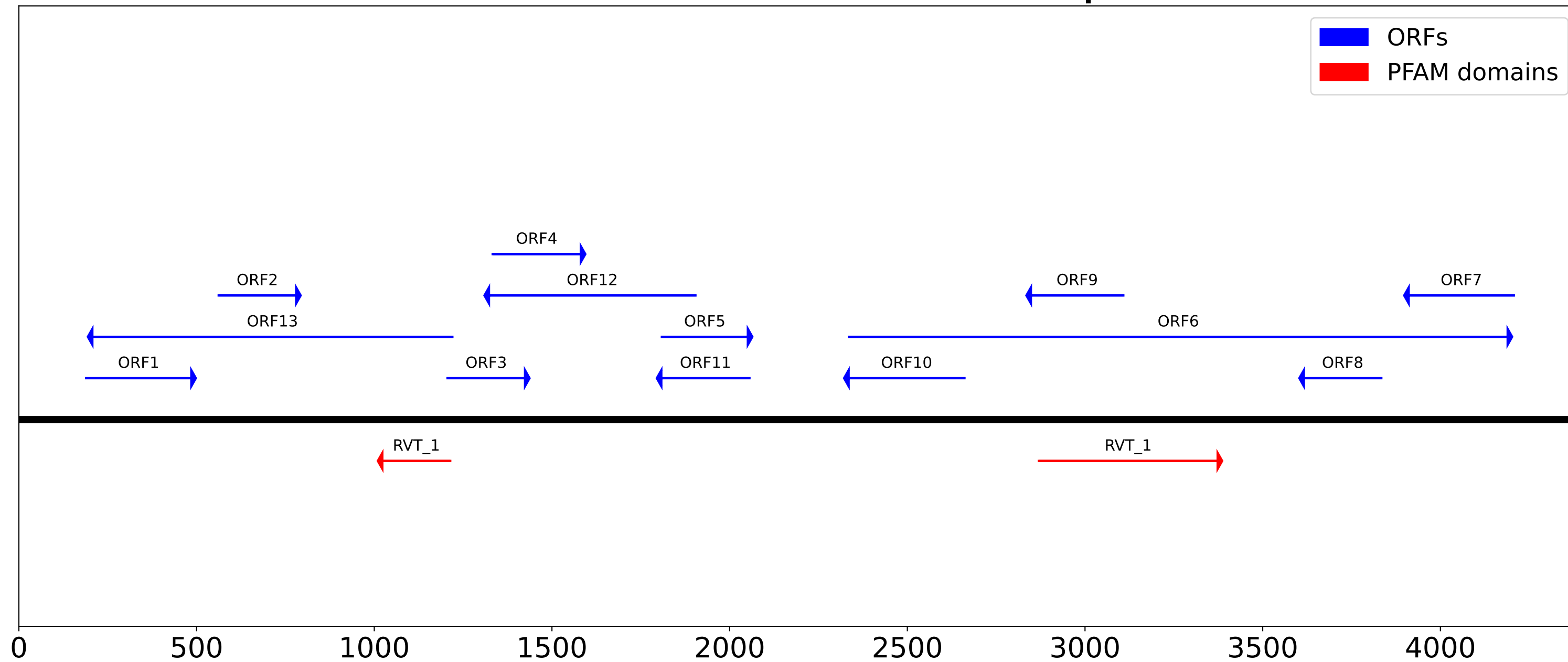


TE: rnd_4_family_2225
size: 1553bp; fragments: 10883; full length: 282 (≥ 1397.7 bp)

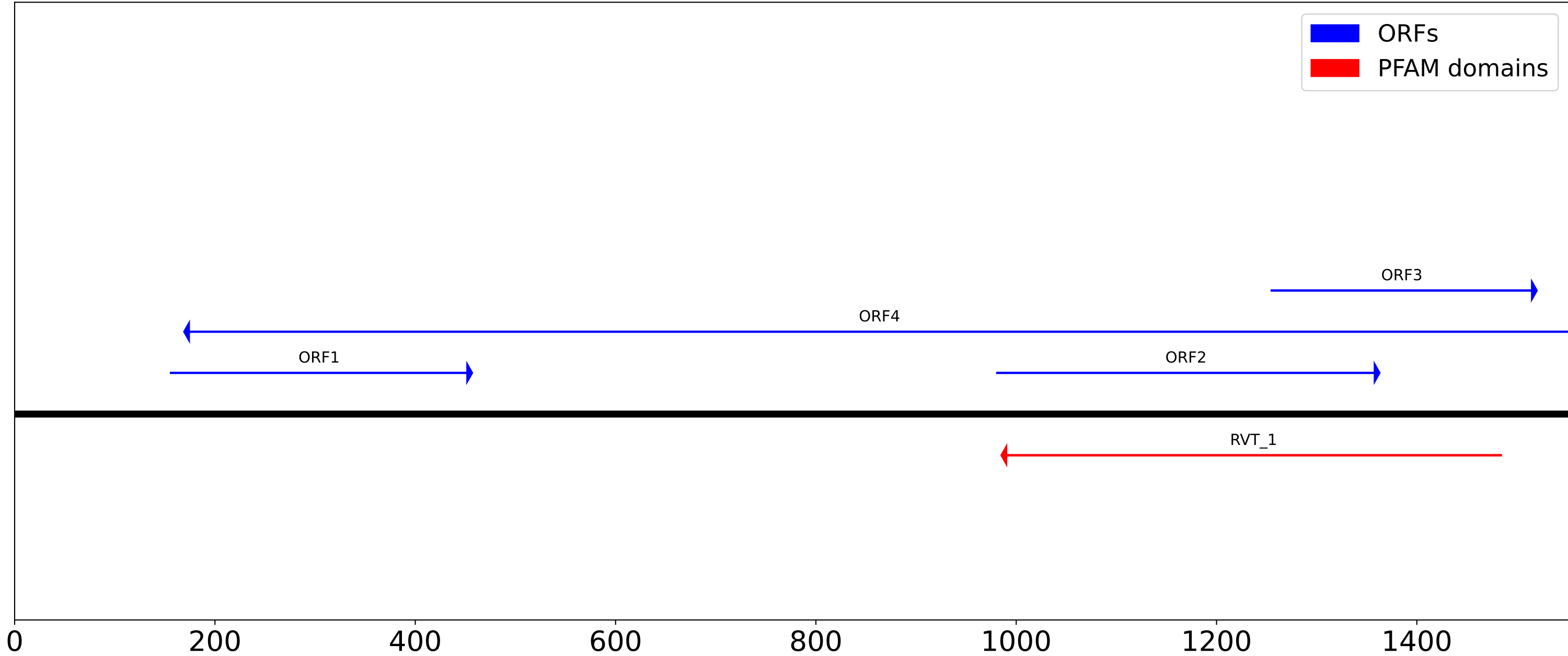
Before TEtrimmer 1553 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

E consensus before TEtrimmer (bp)

