

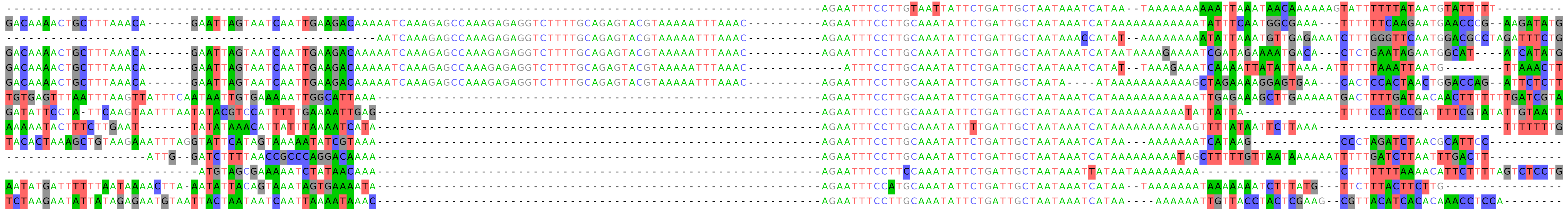
Start crop Point



End crop Point



MSA length = 1792



Start crop Point



End crop Point



0 -

10

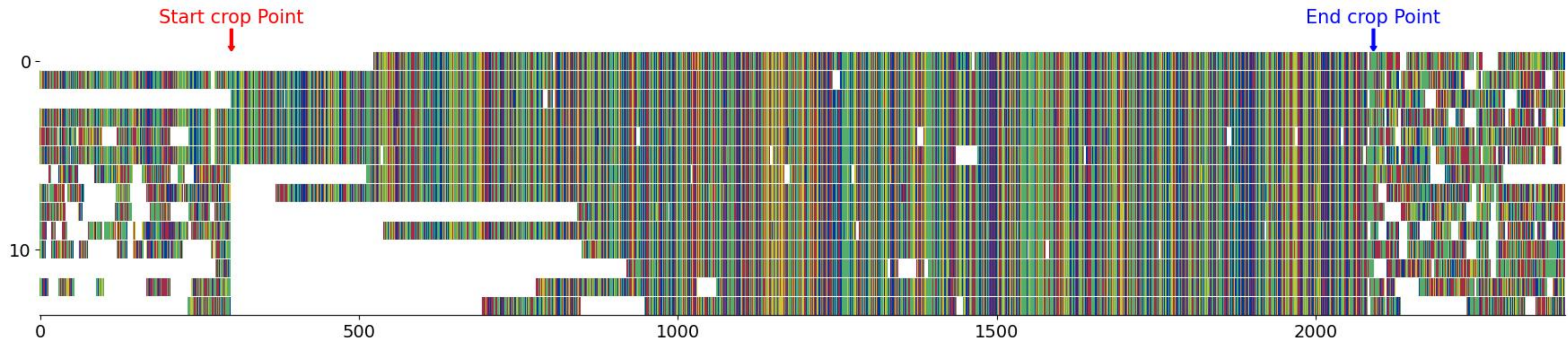
0

500

1000

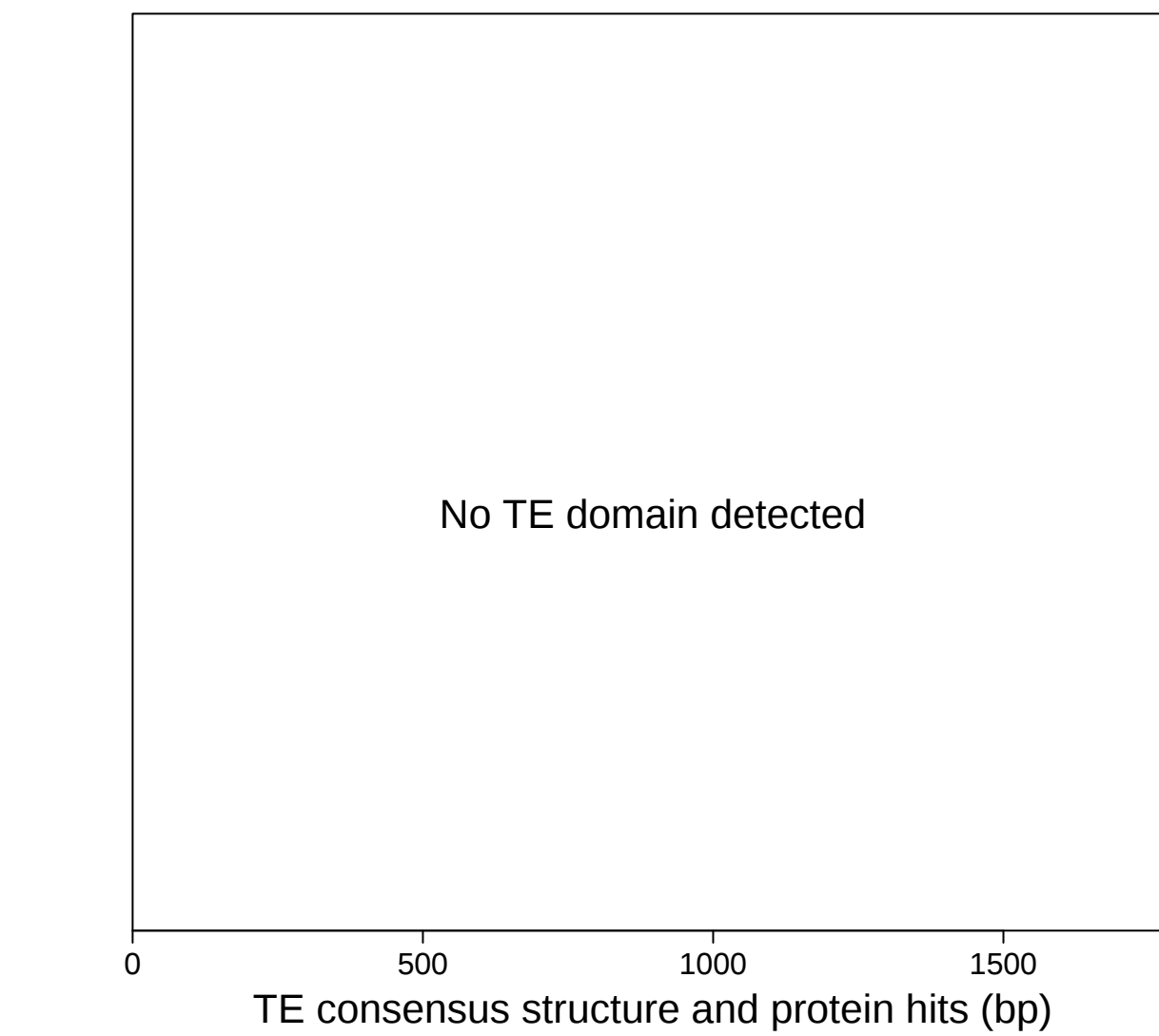
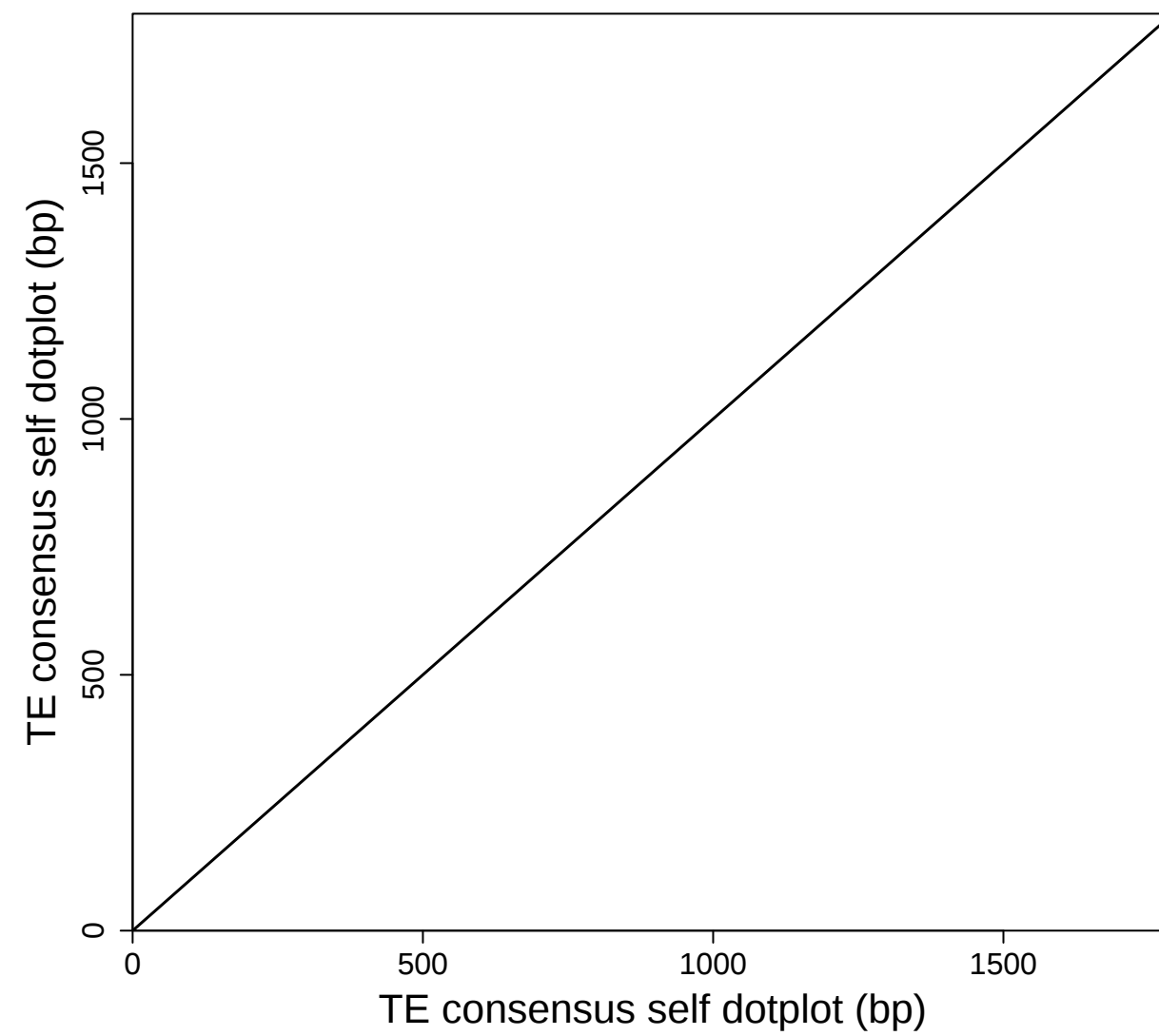
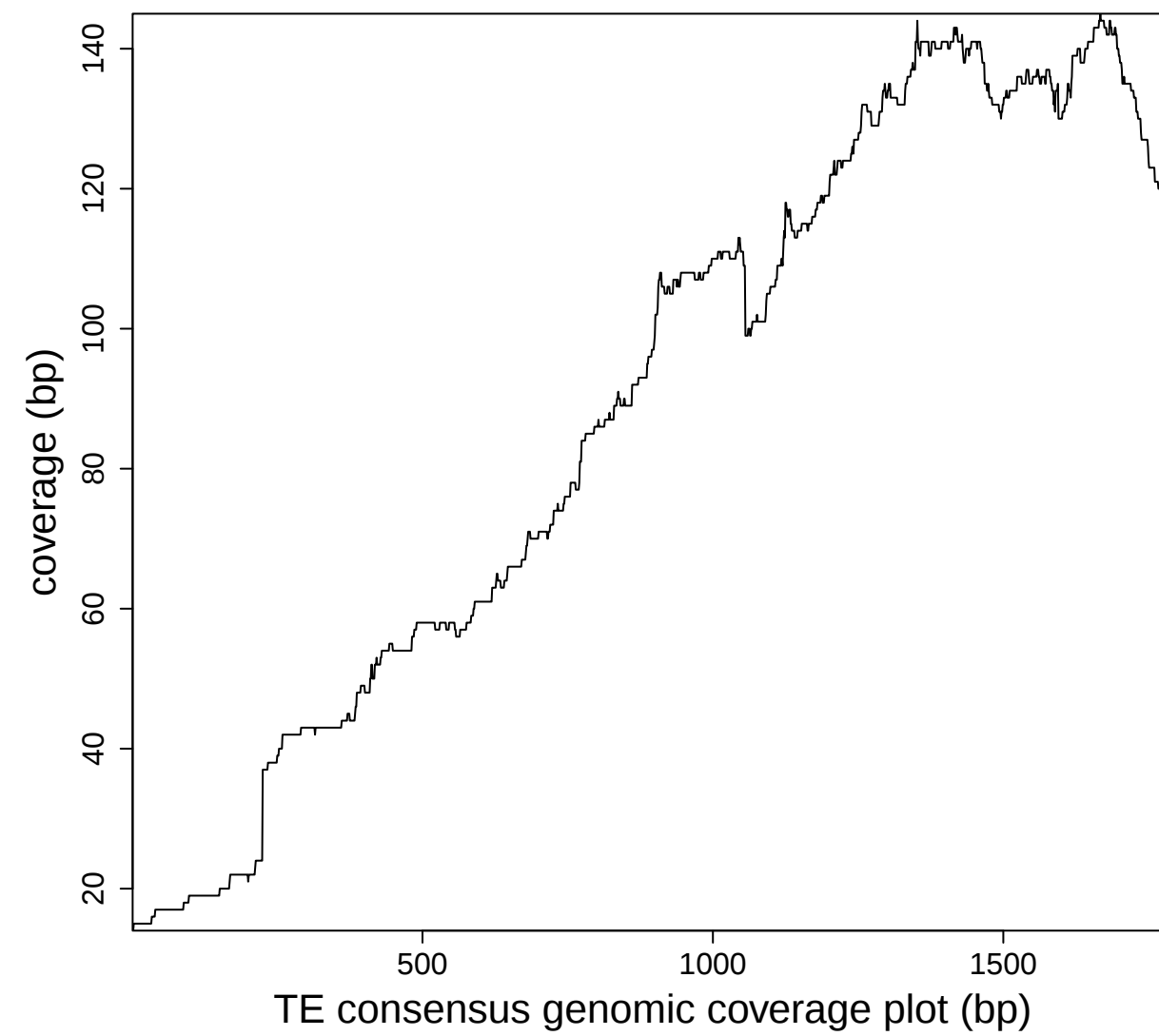
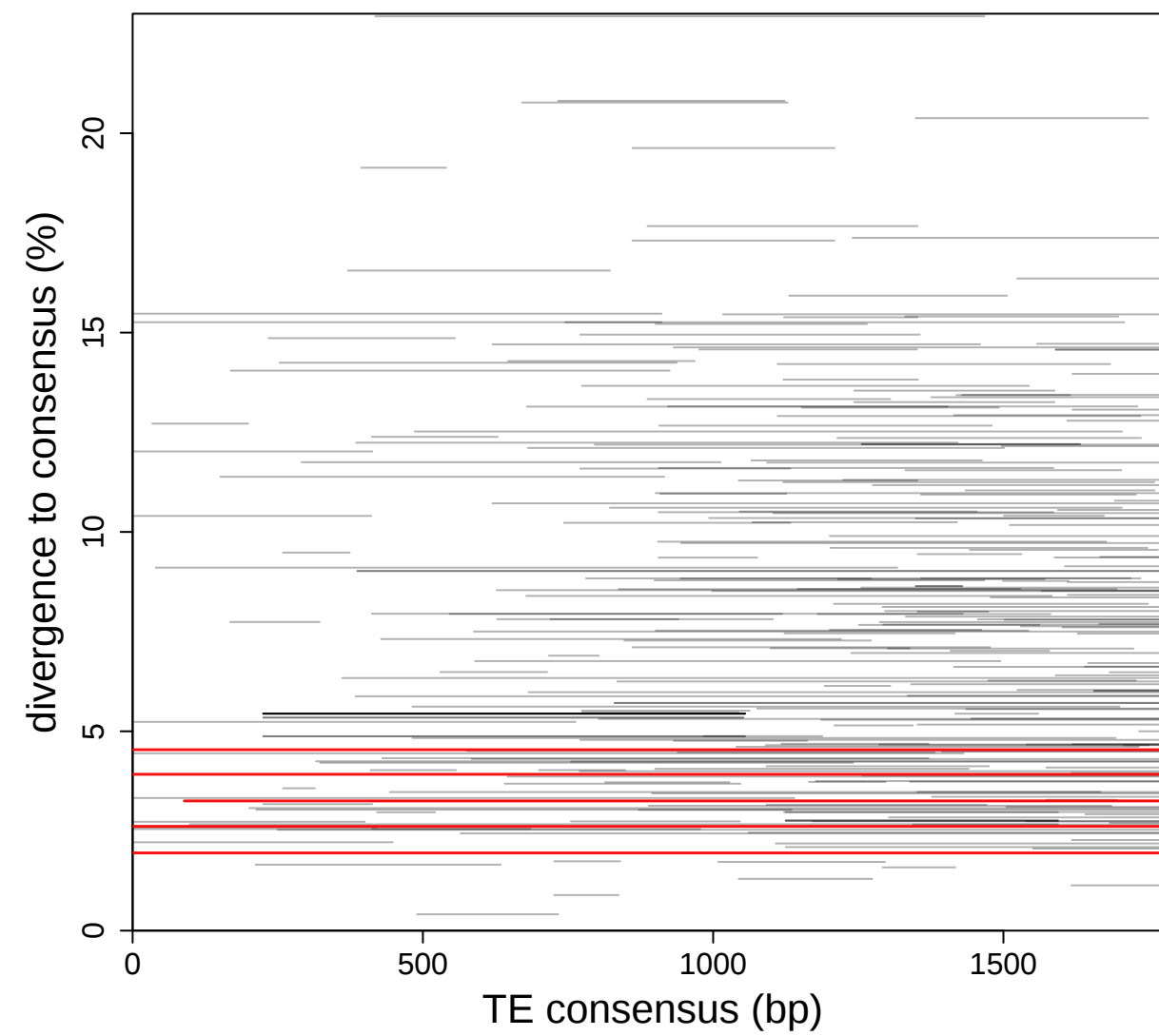
1500

2000



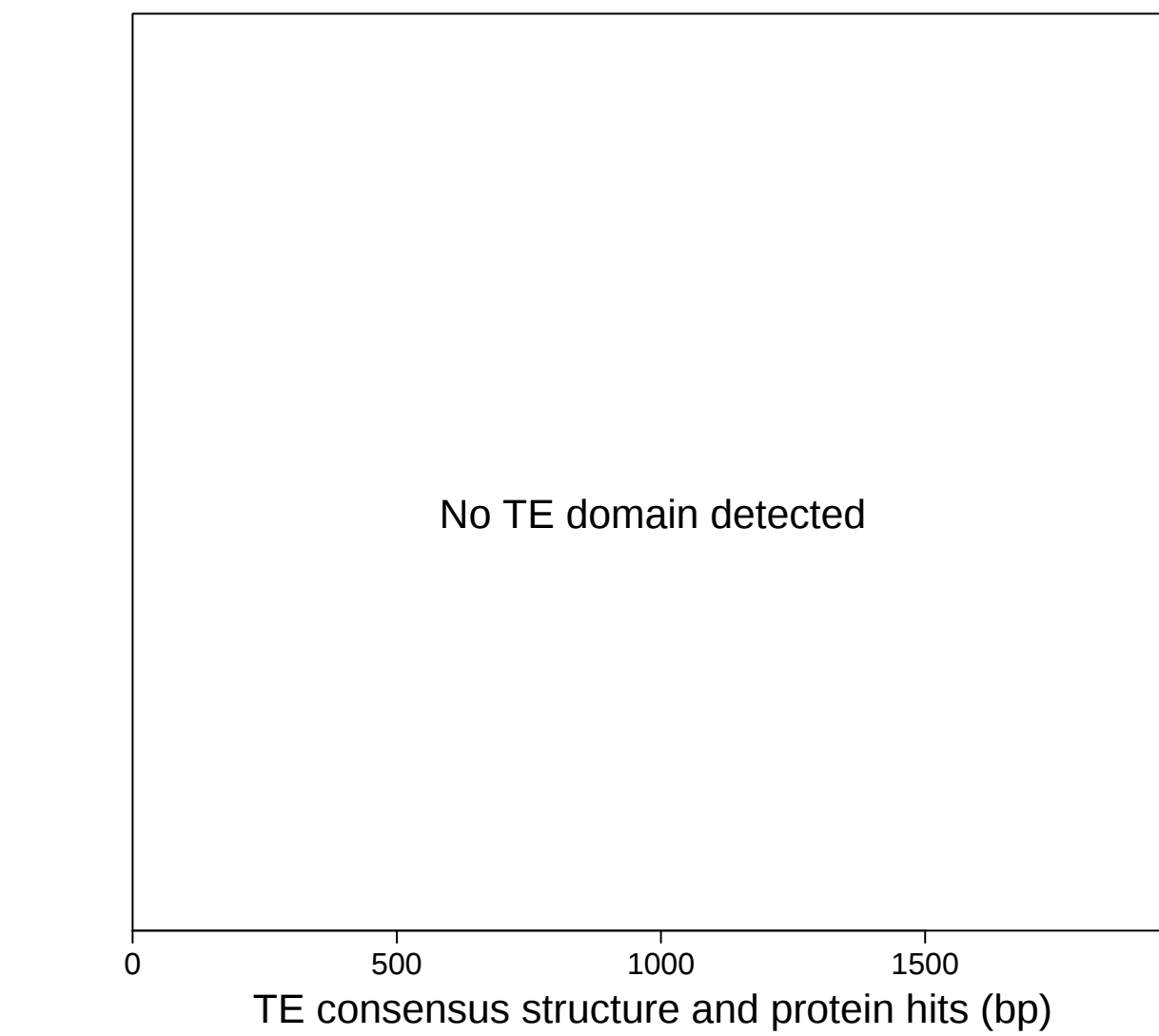
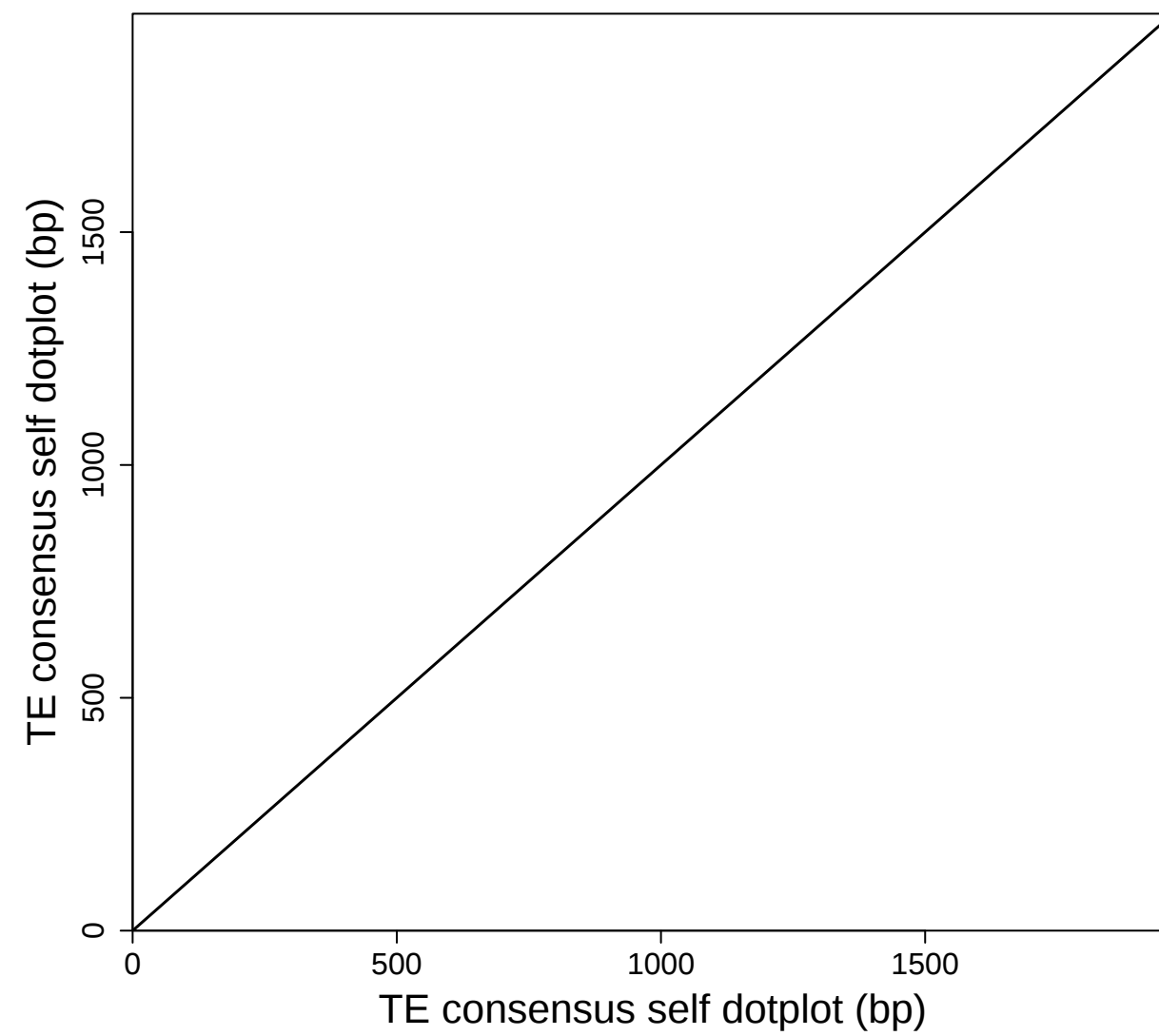
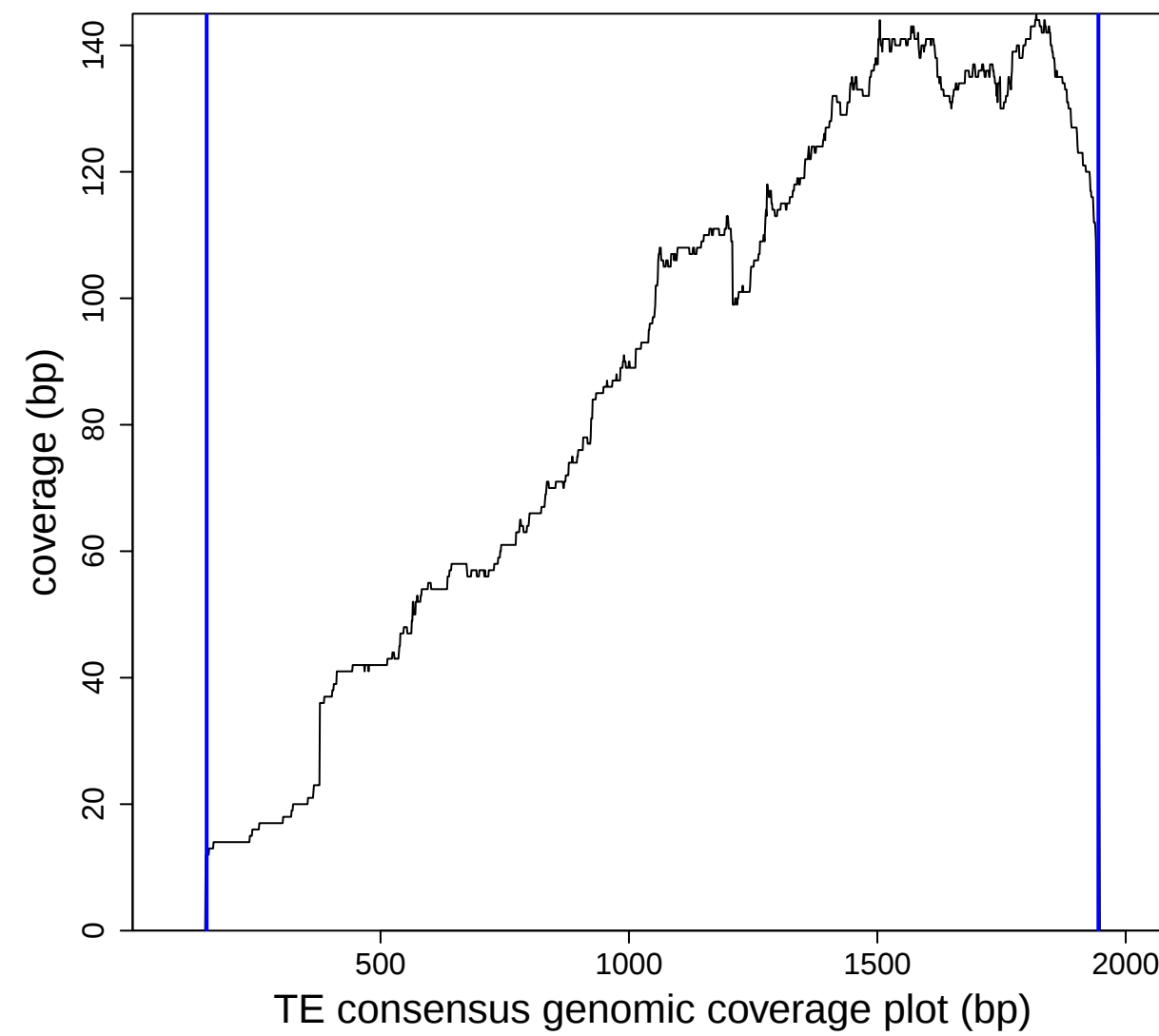
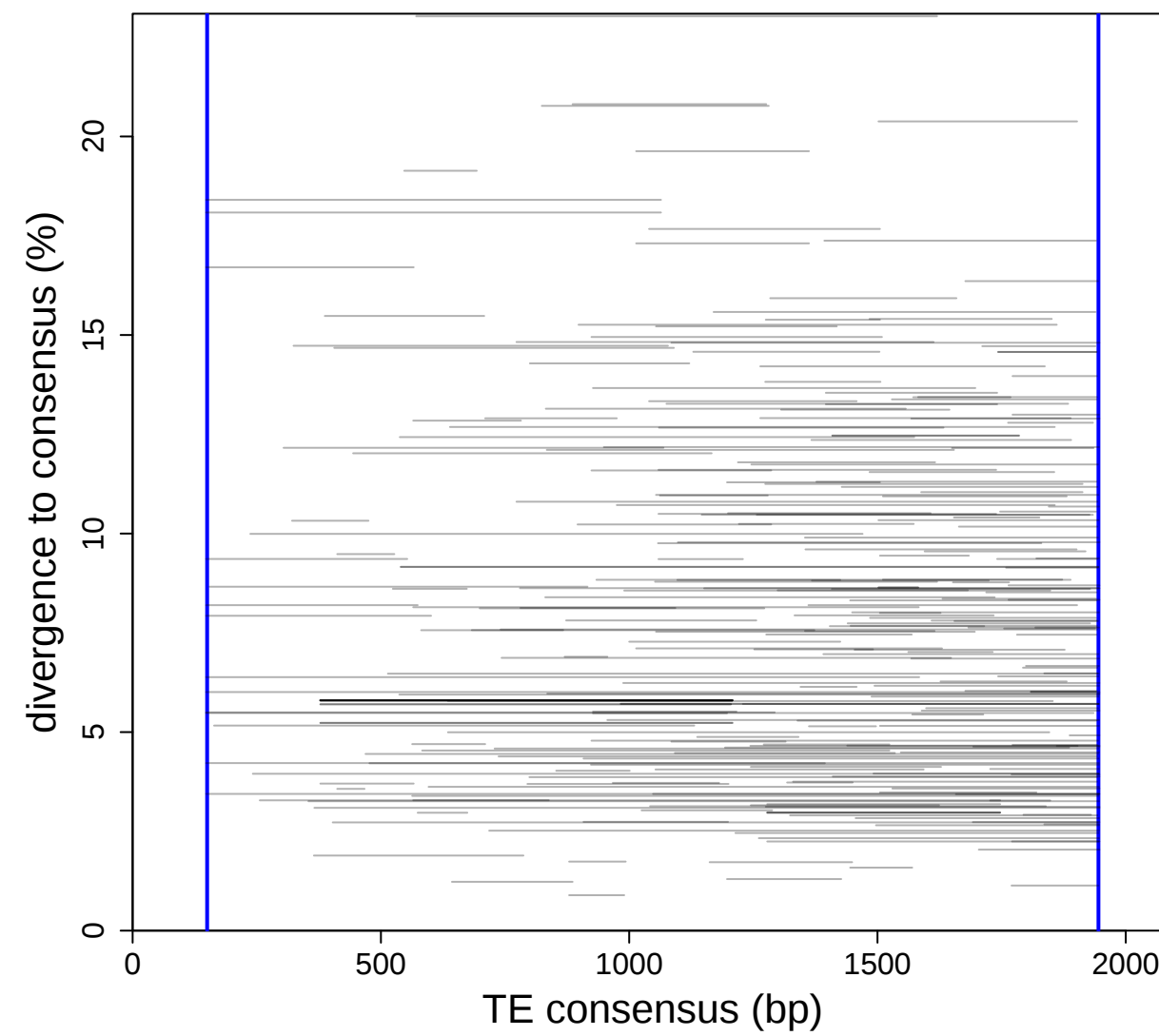
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 1792bp; fragments: 286; full length: 5 (>=1612.8bp)

After TEtrimmer 1792 bp



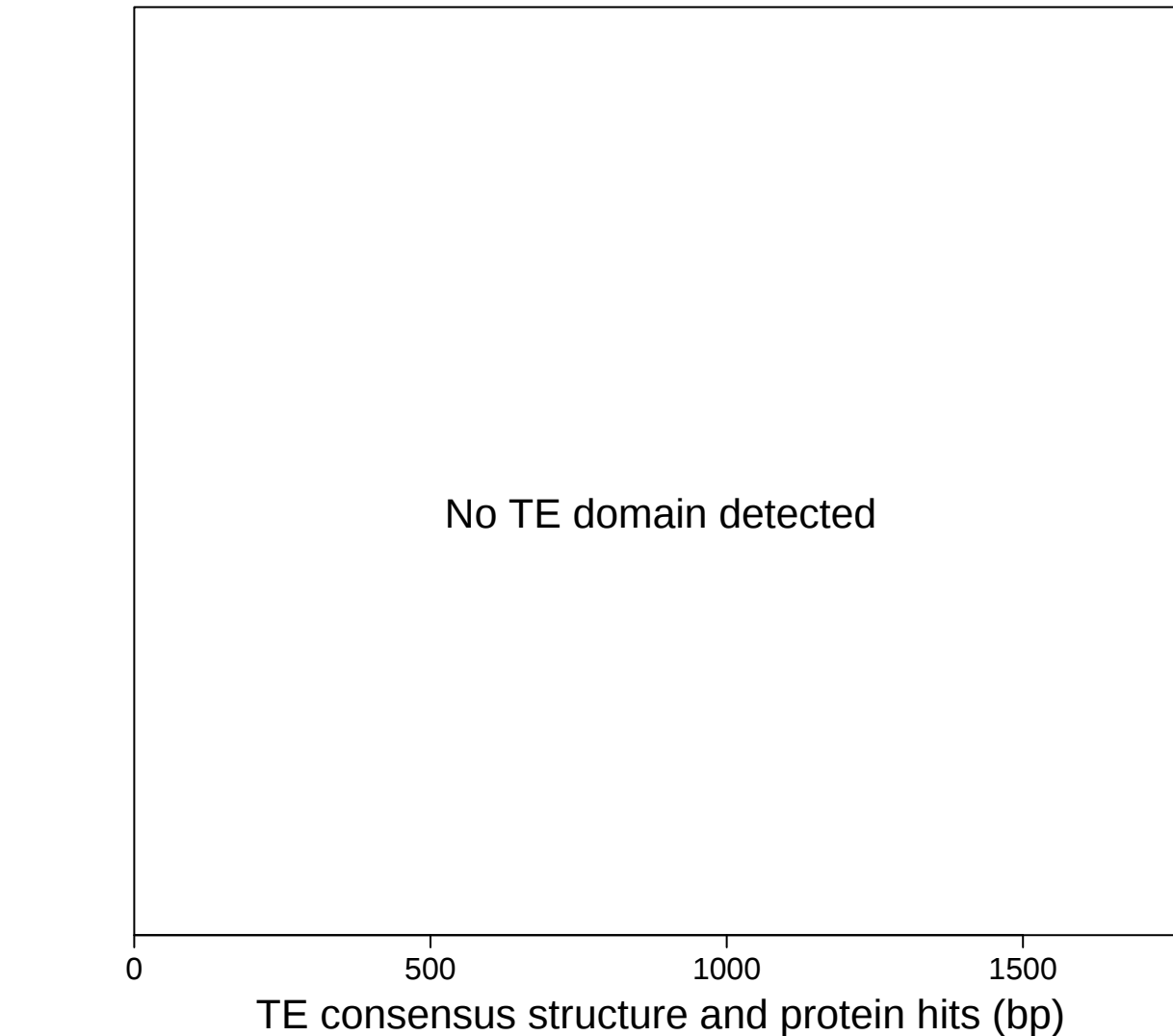
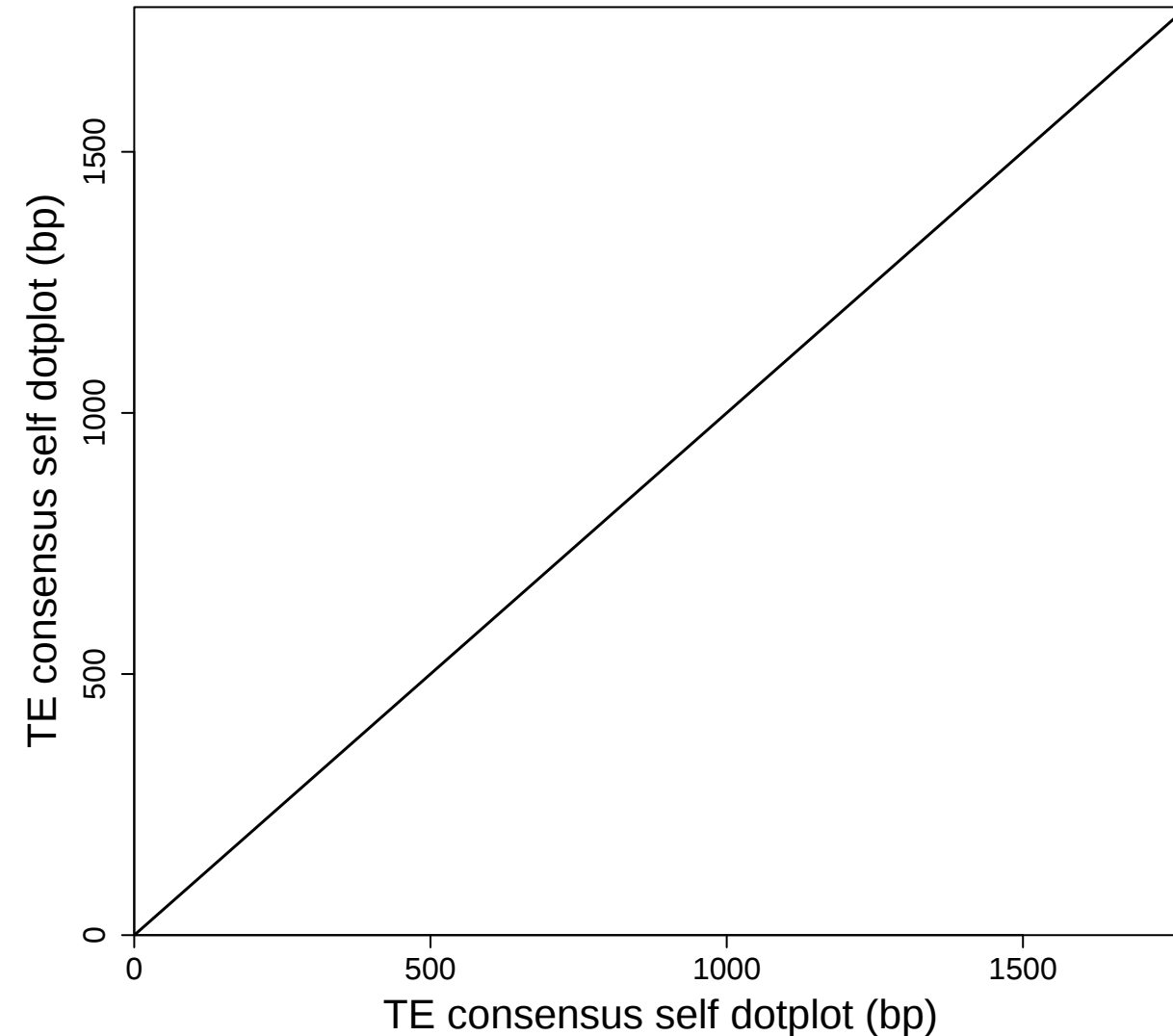
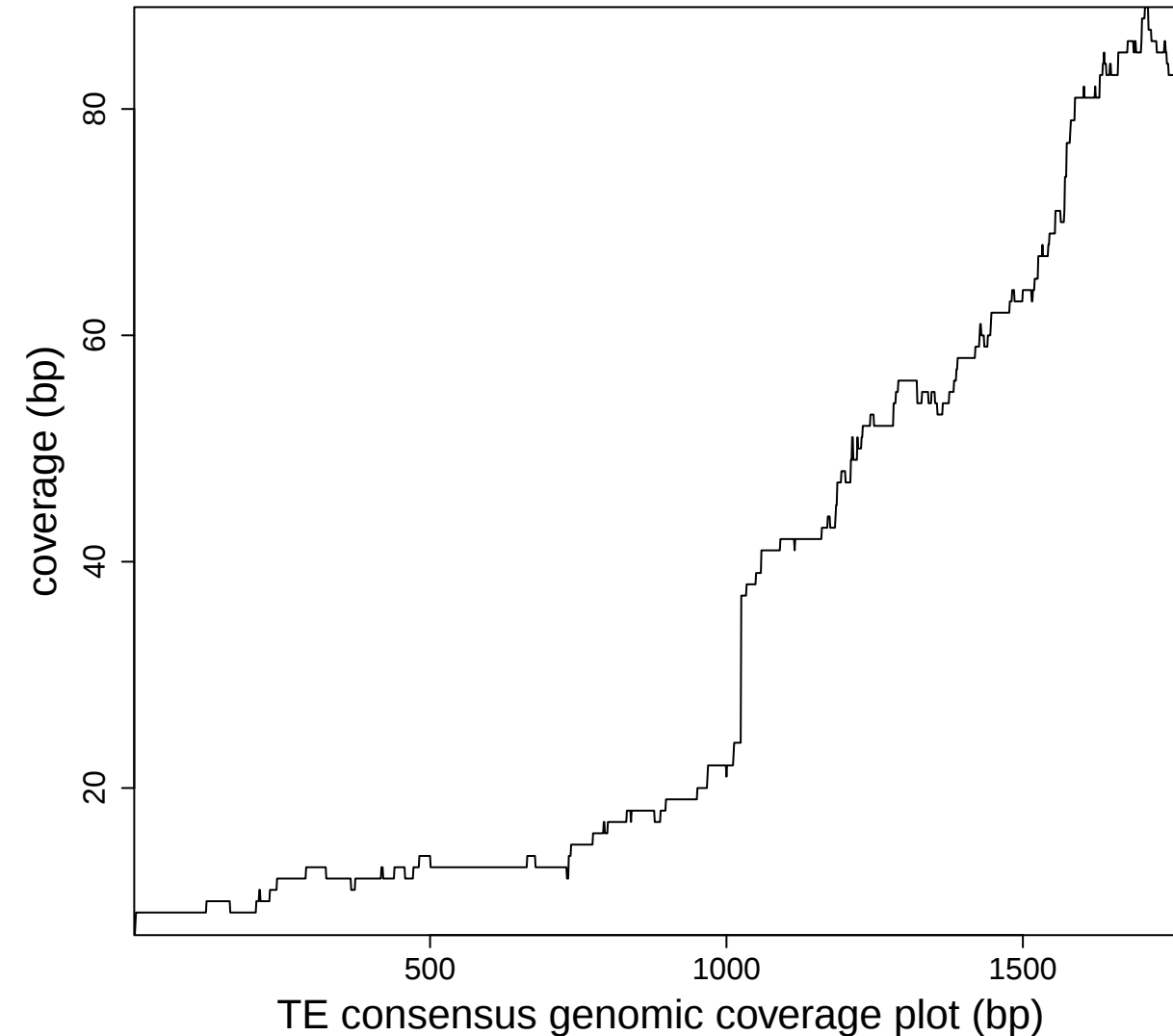
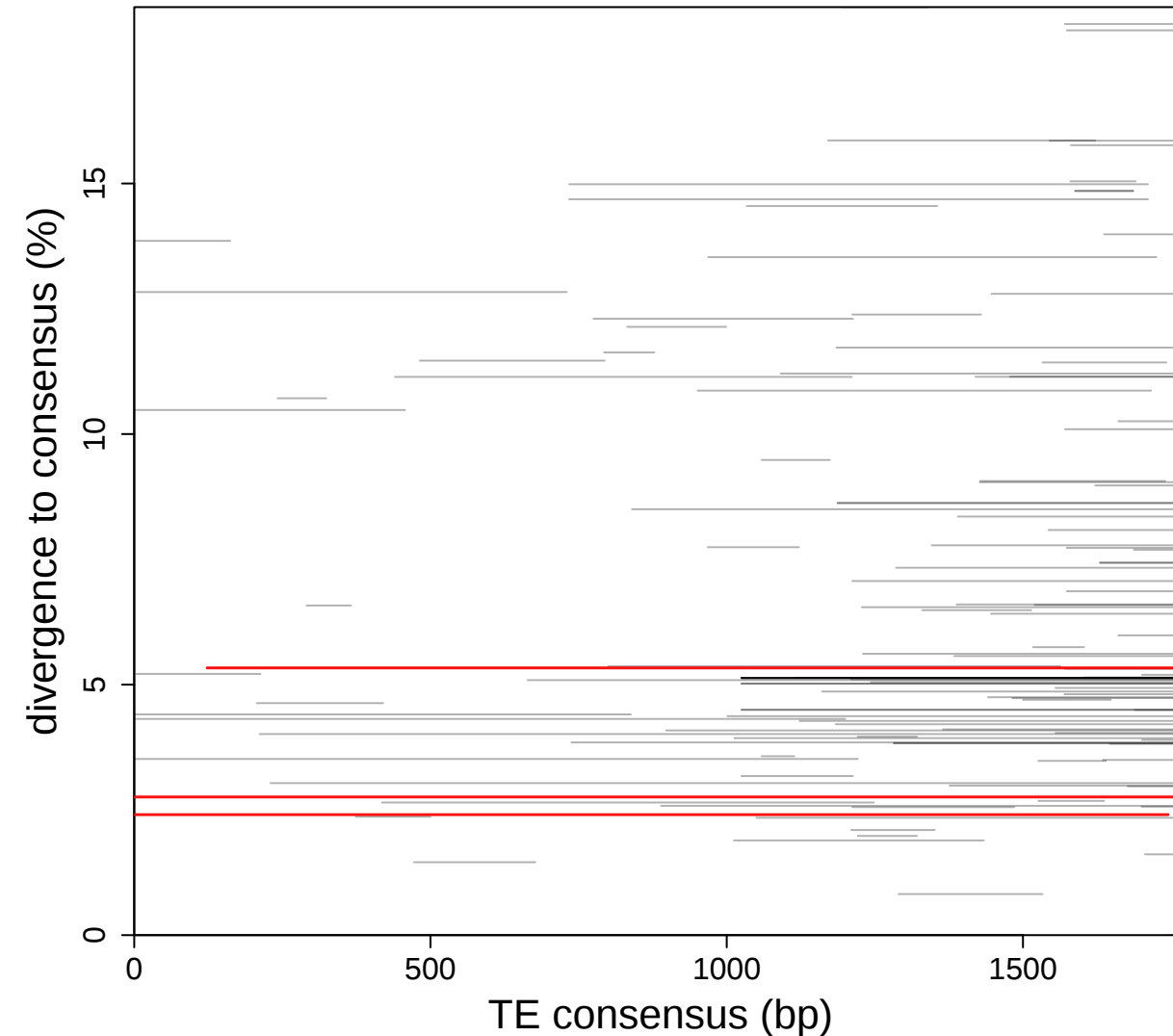
2.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 2095bp; fragments: 285; full length: 0 (>=1885.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

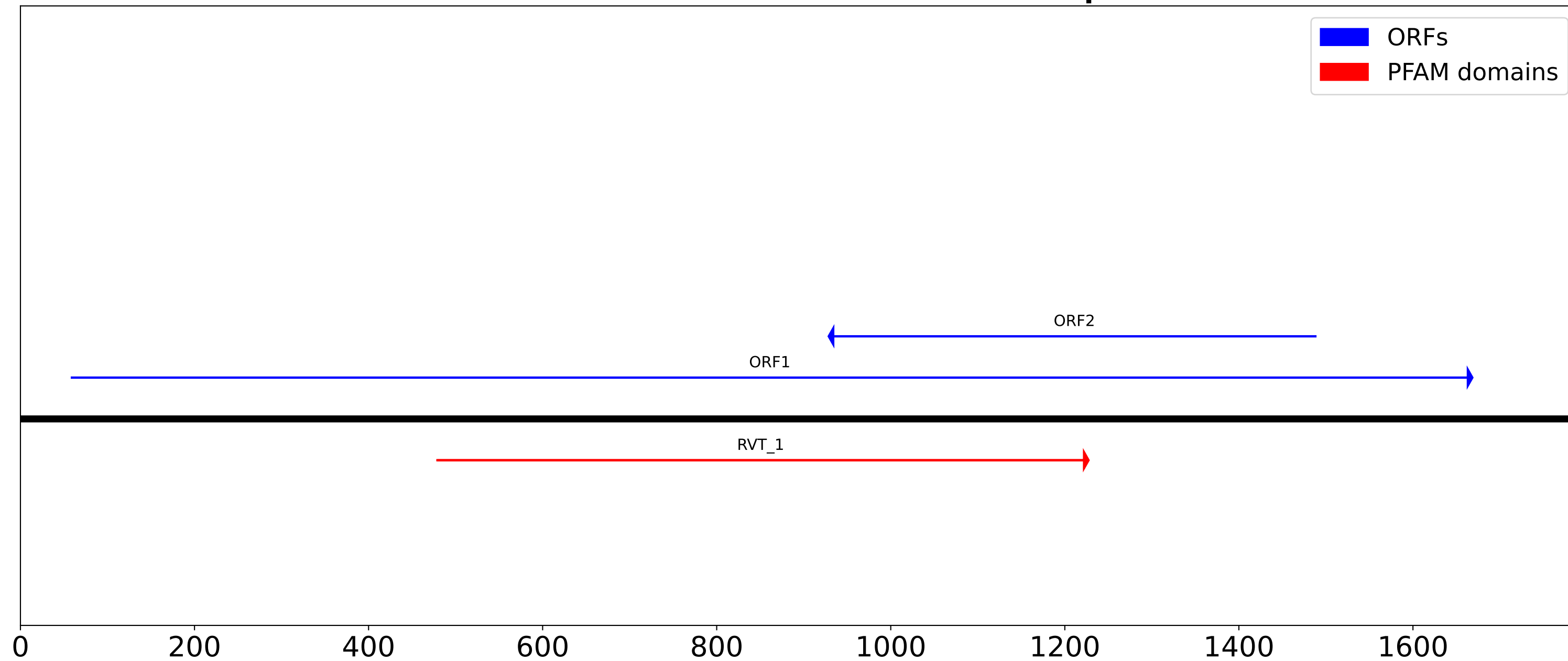


TE: rnd_5_family_2262
size: 1777bp; fragments: 131; full length: 3 (>=1599.3bp)

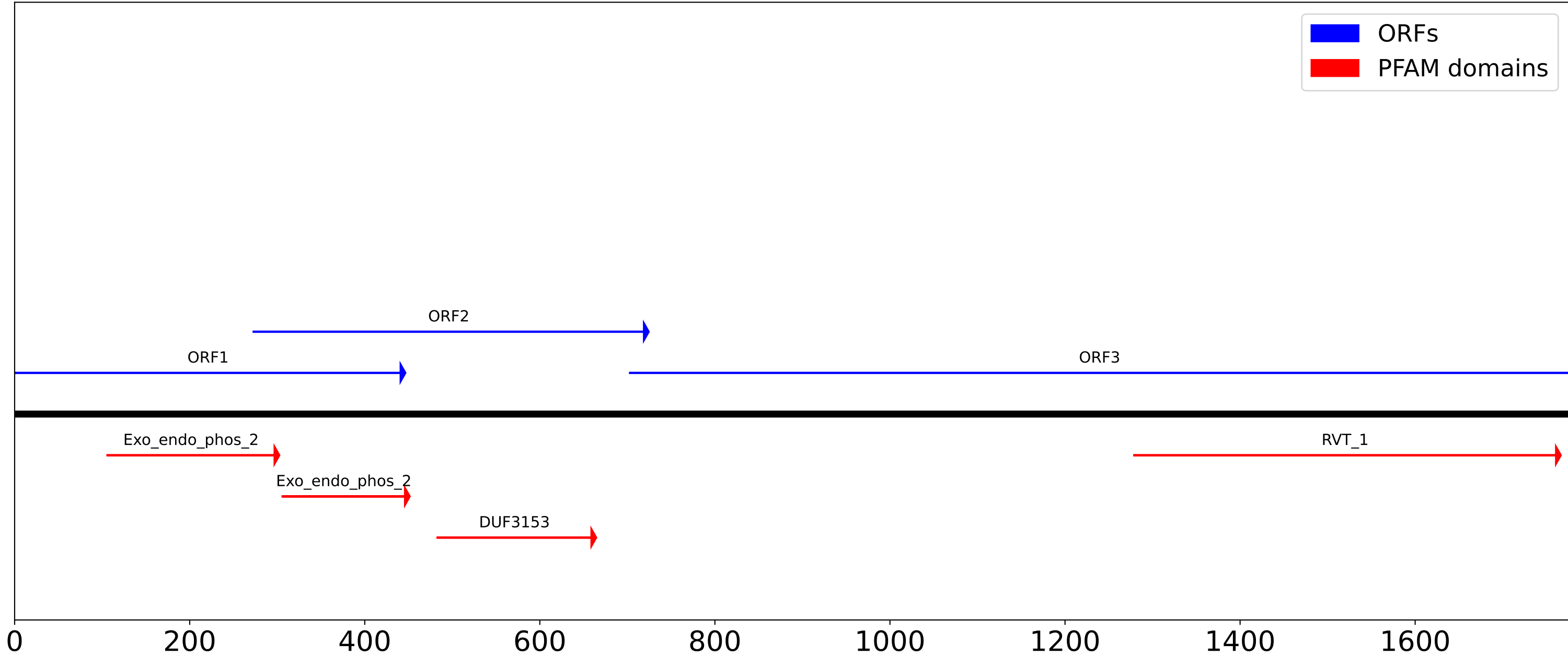
Before TEtrimmer 1777 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

