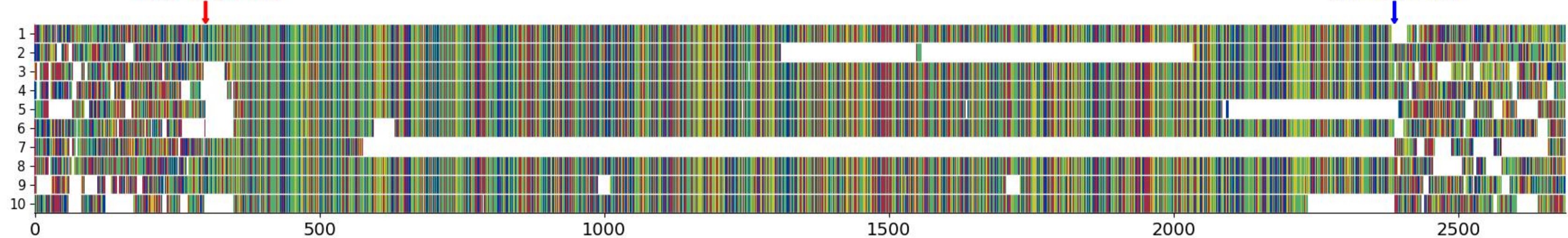


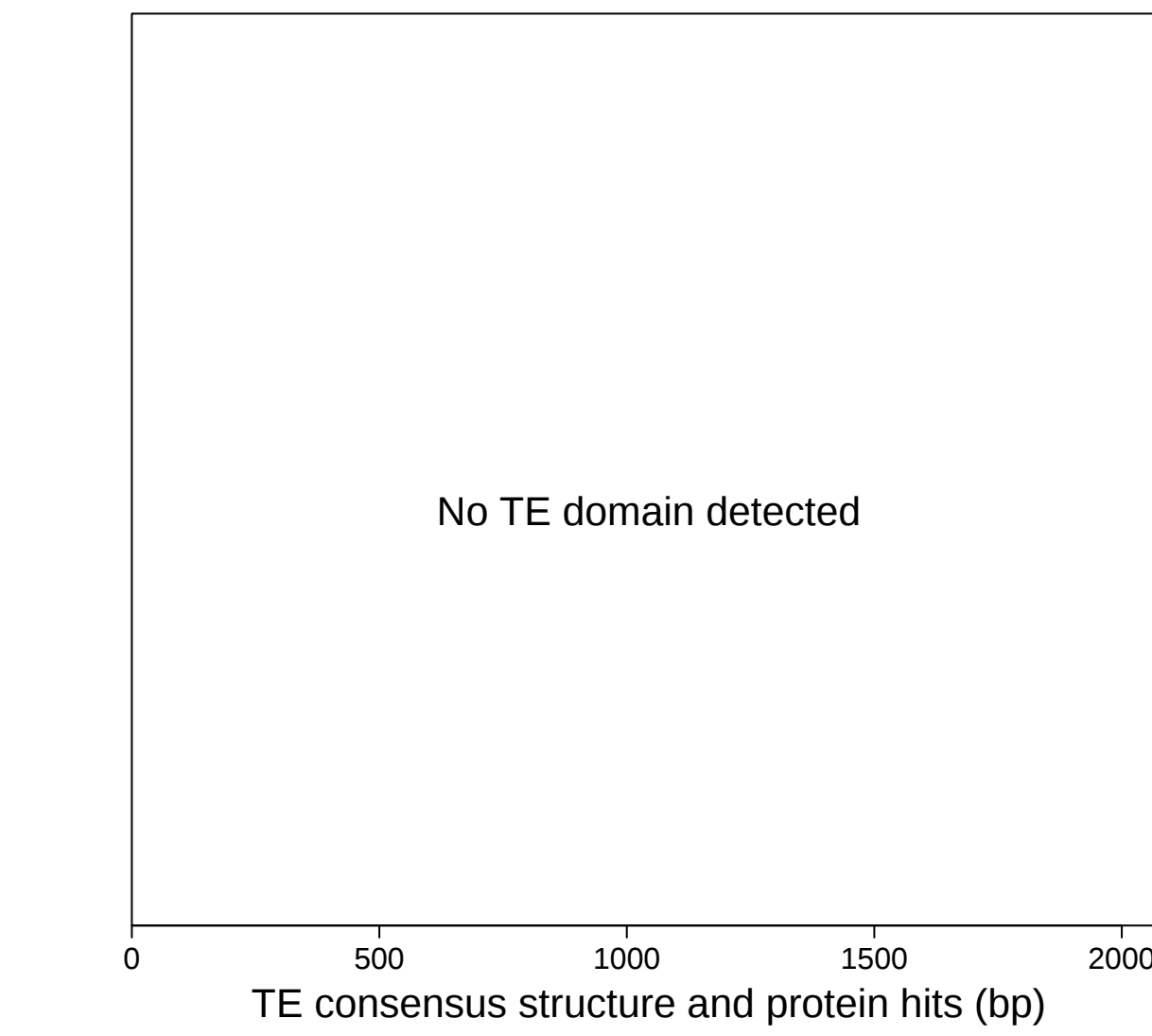
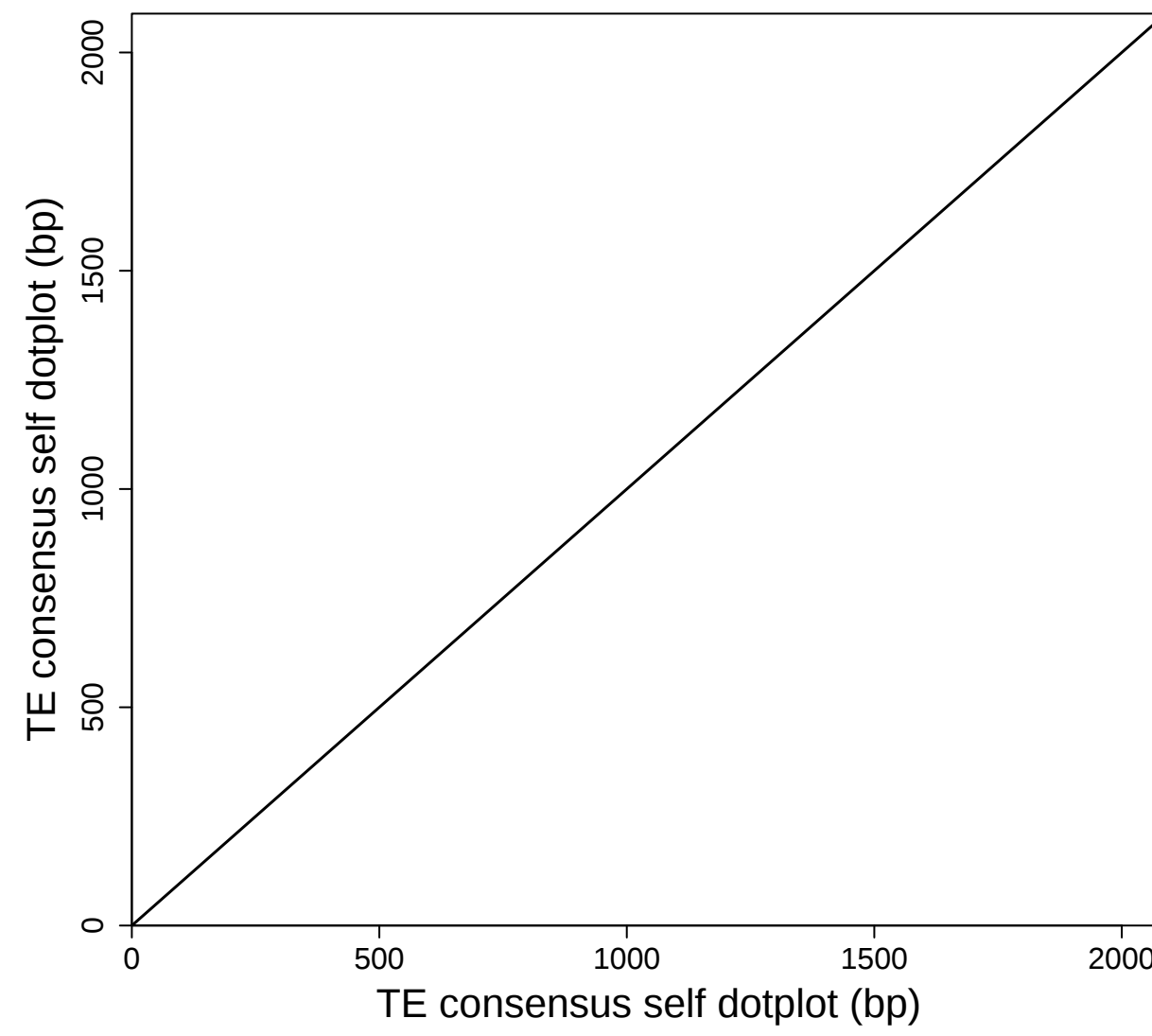
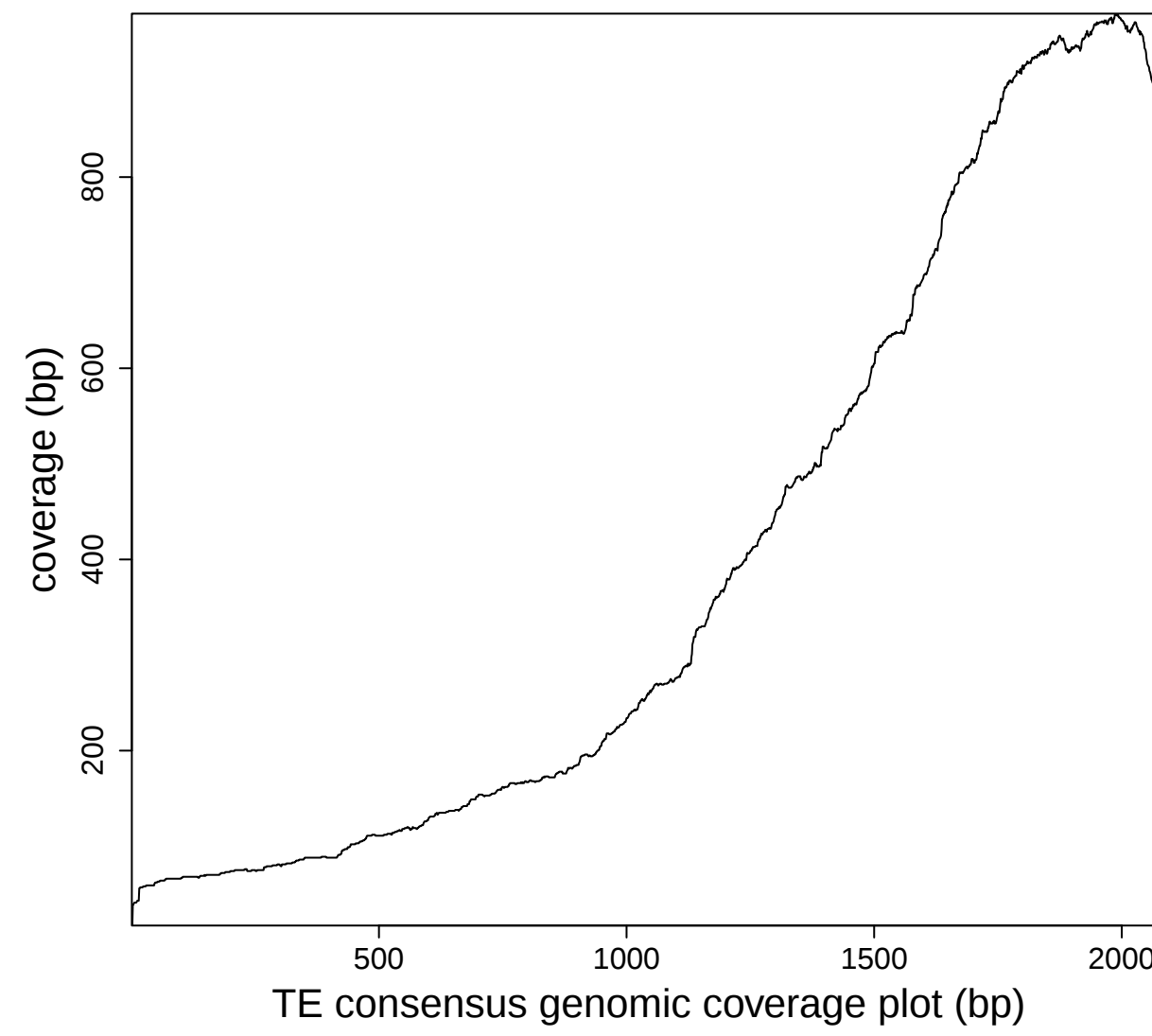
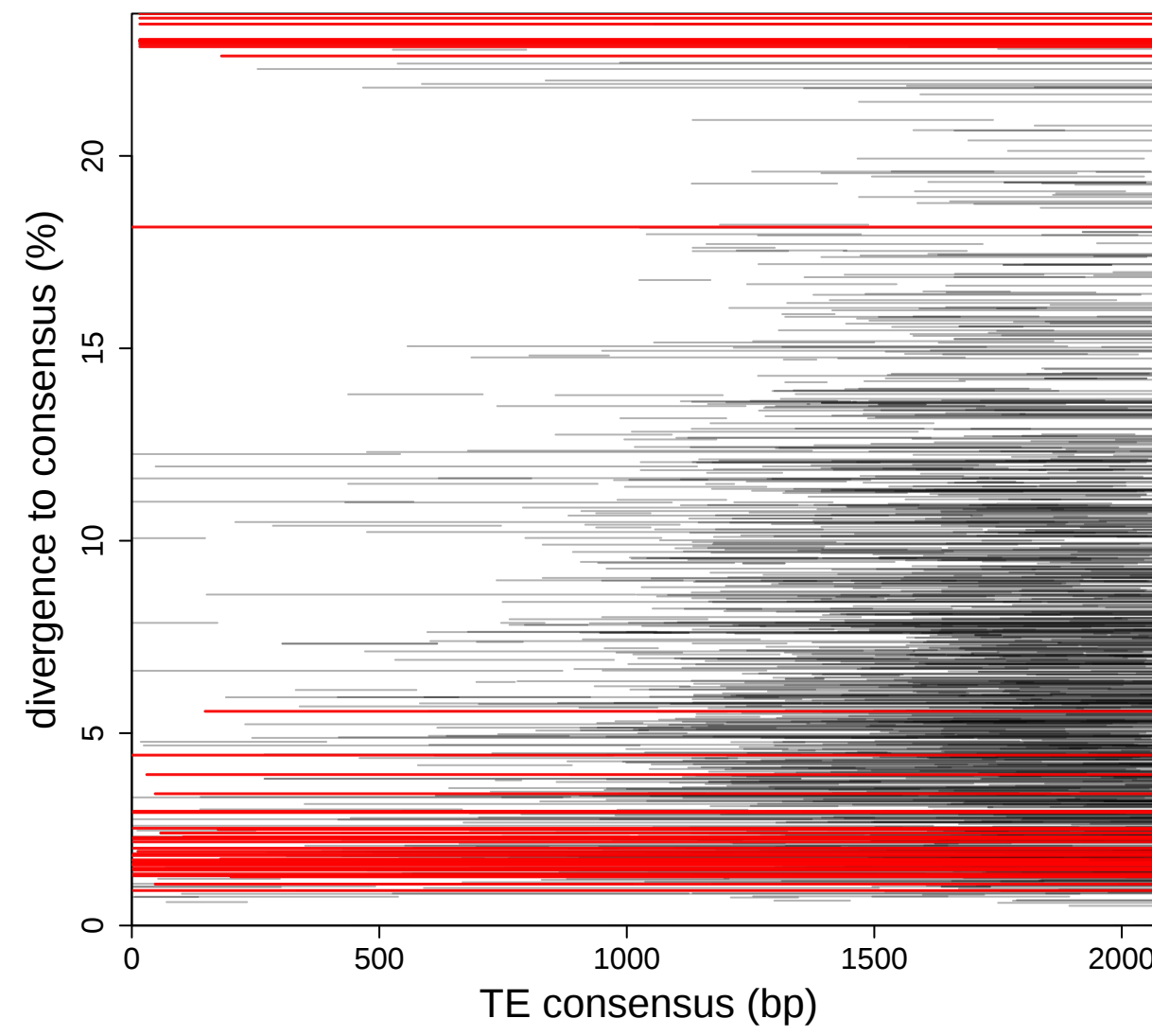
Start crop Point

End crop Point



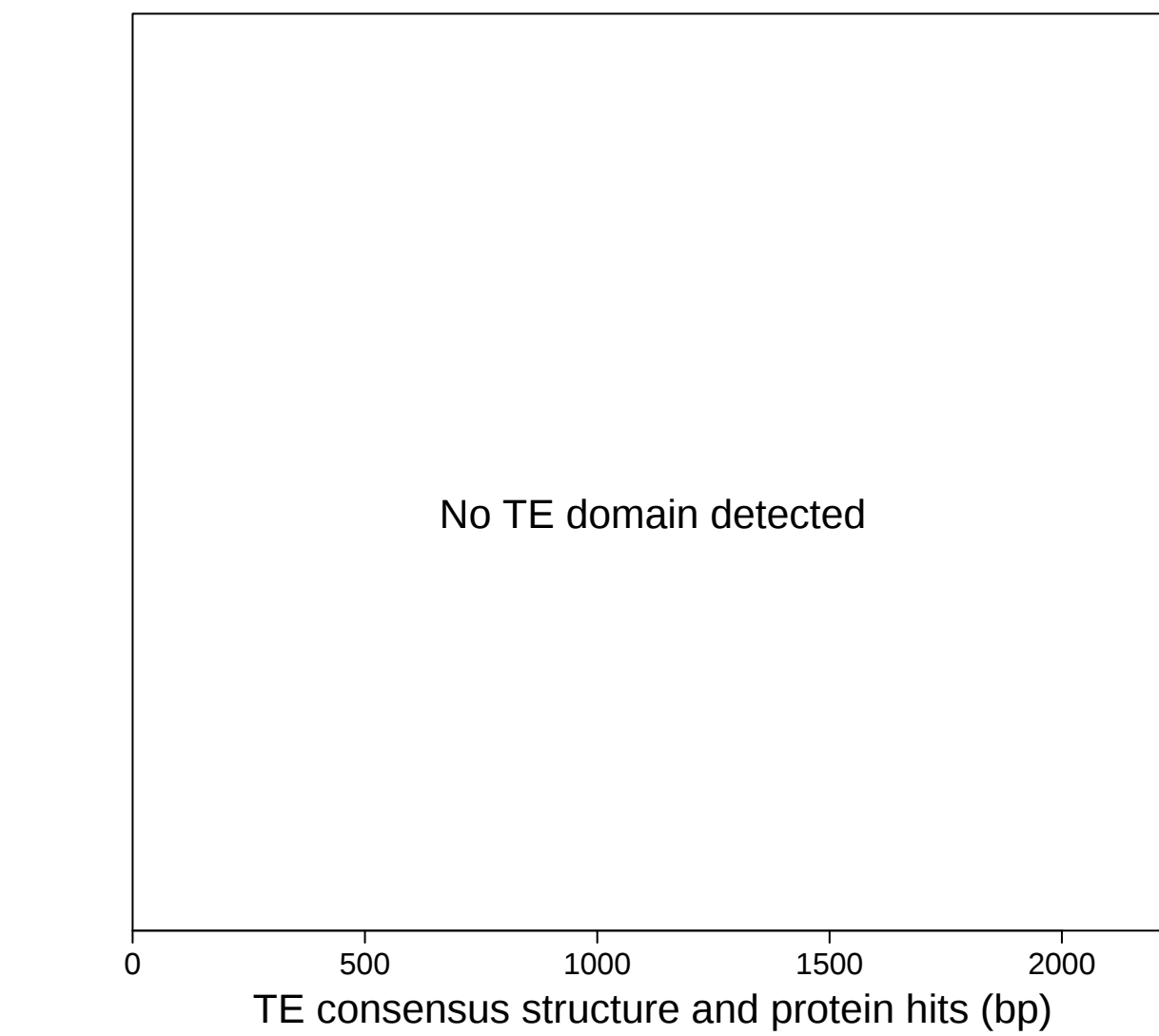
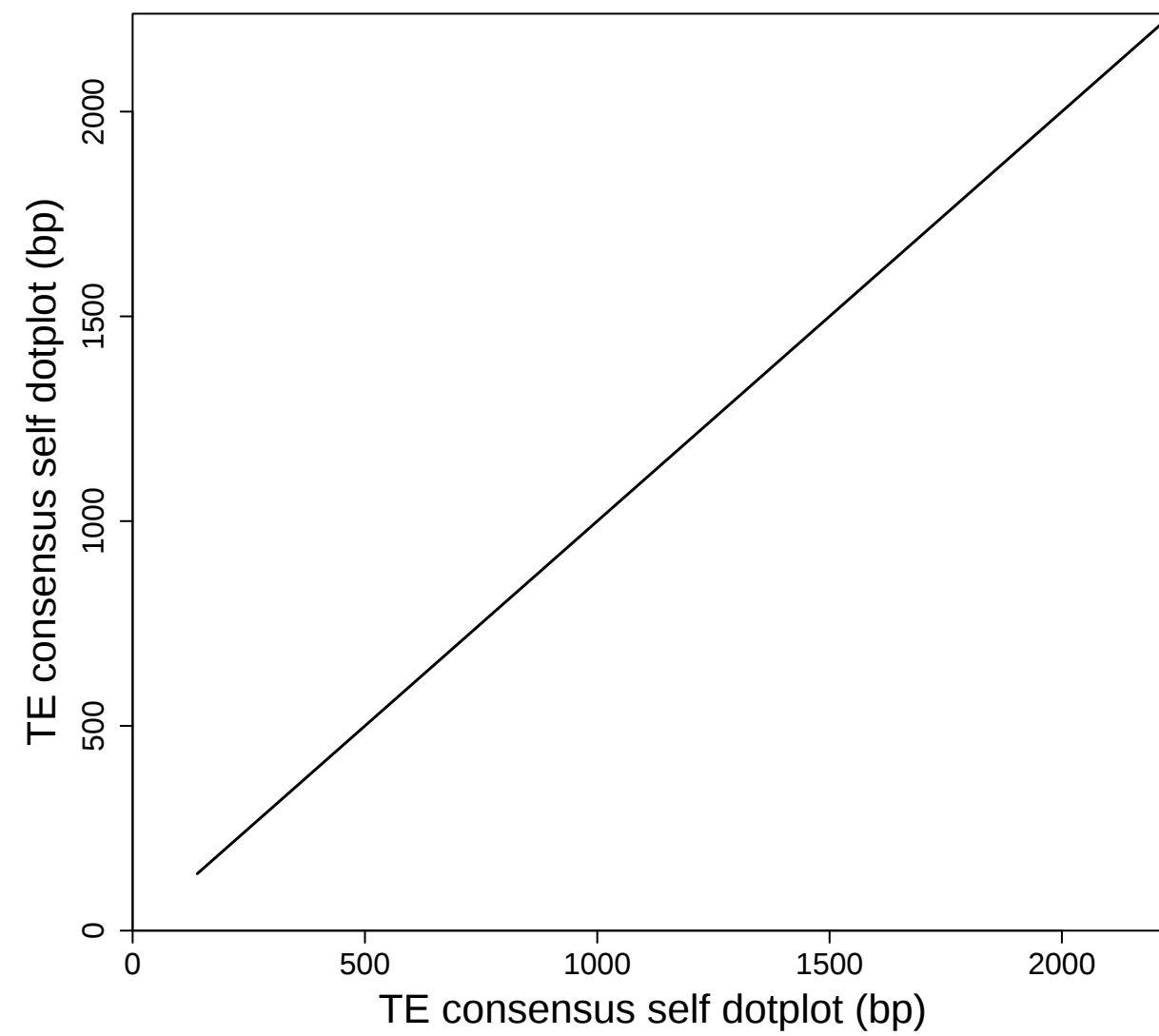
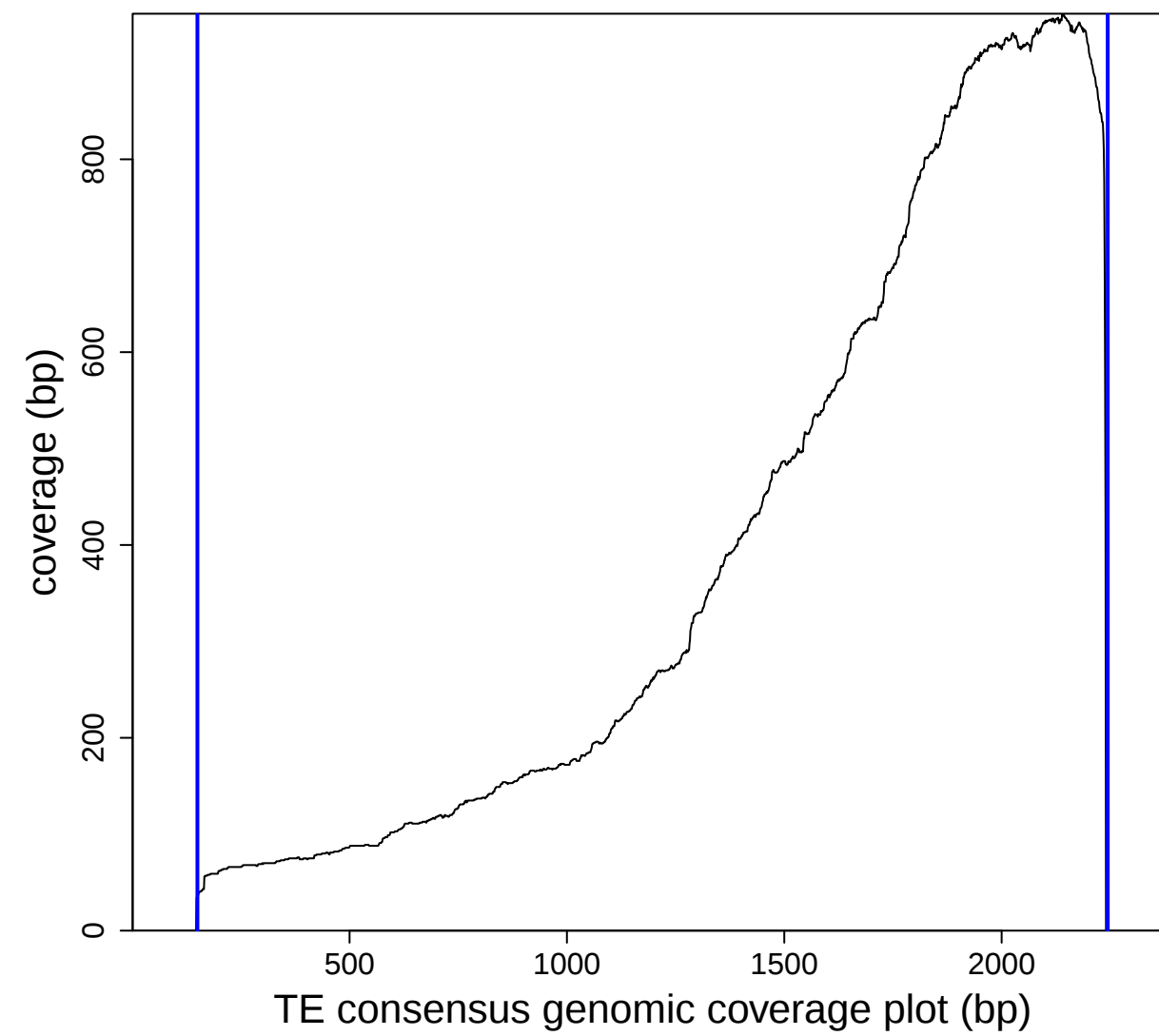
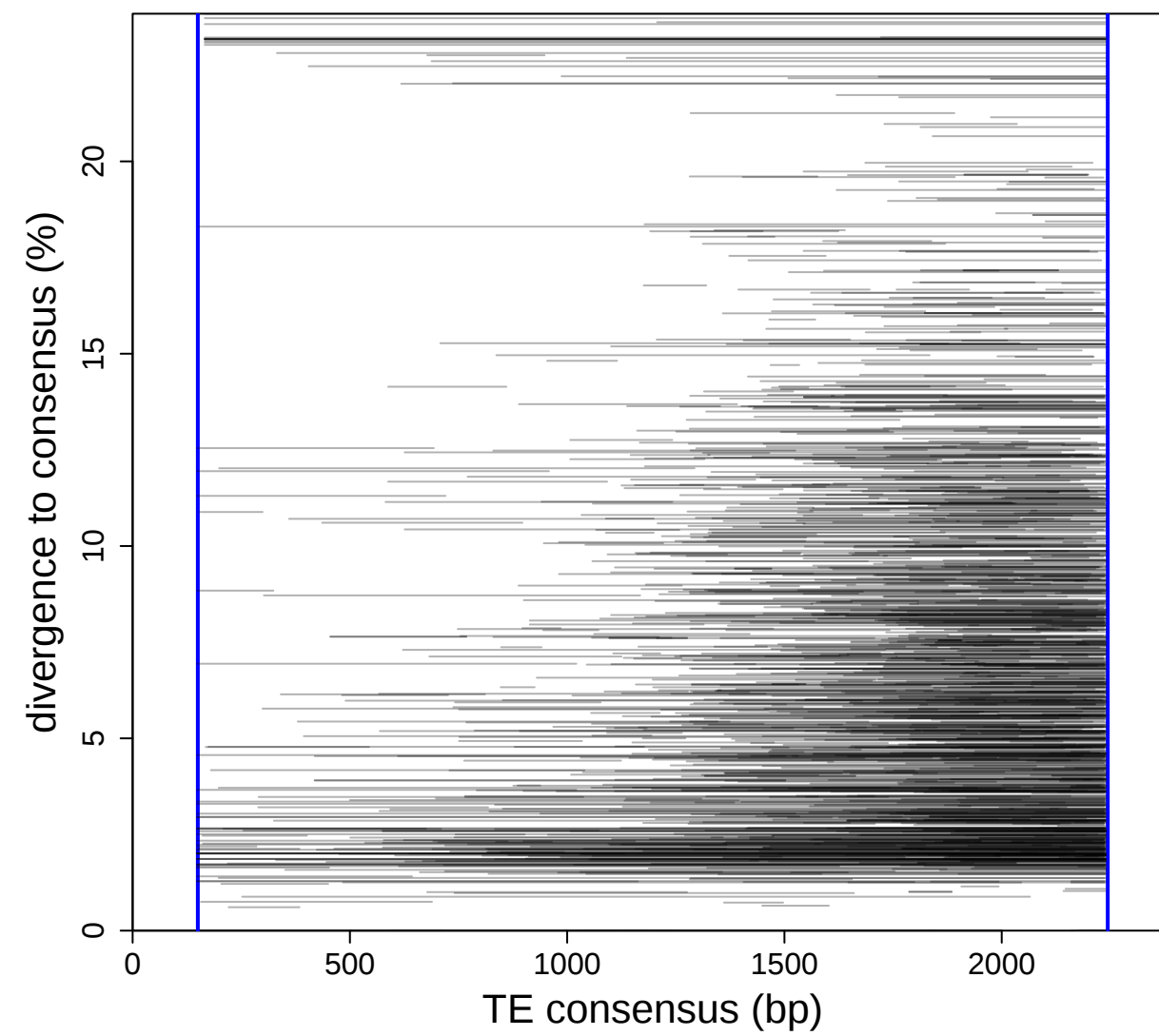
bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 2089bp; fragments: 1472; full length: 48 (≥ 1880.1 bp)

After TEtrimmer 2089 bp

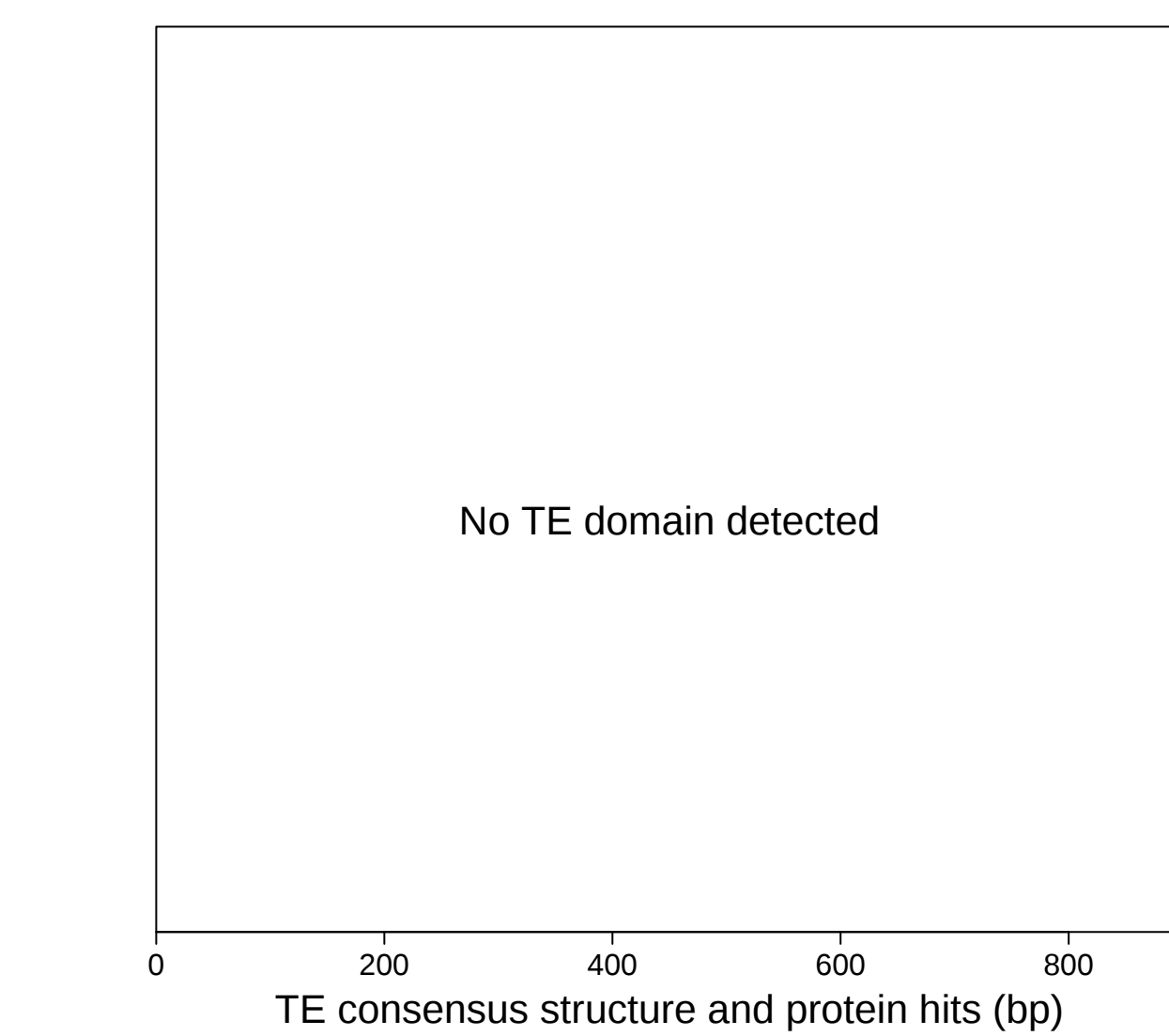
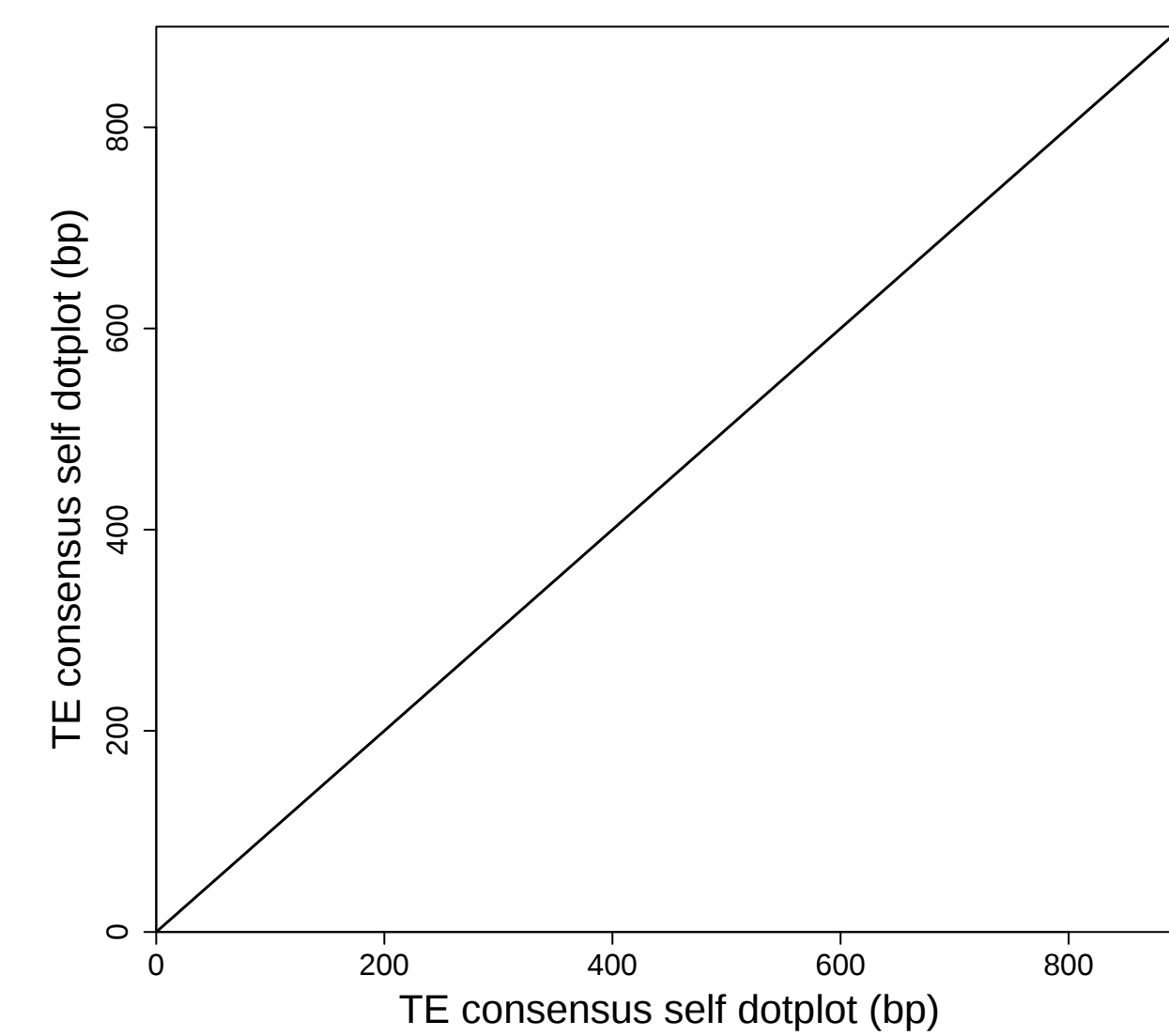
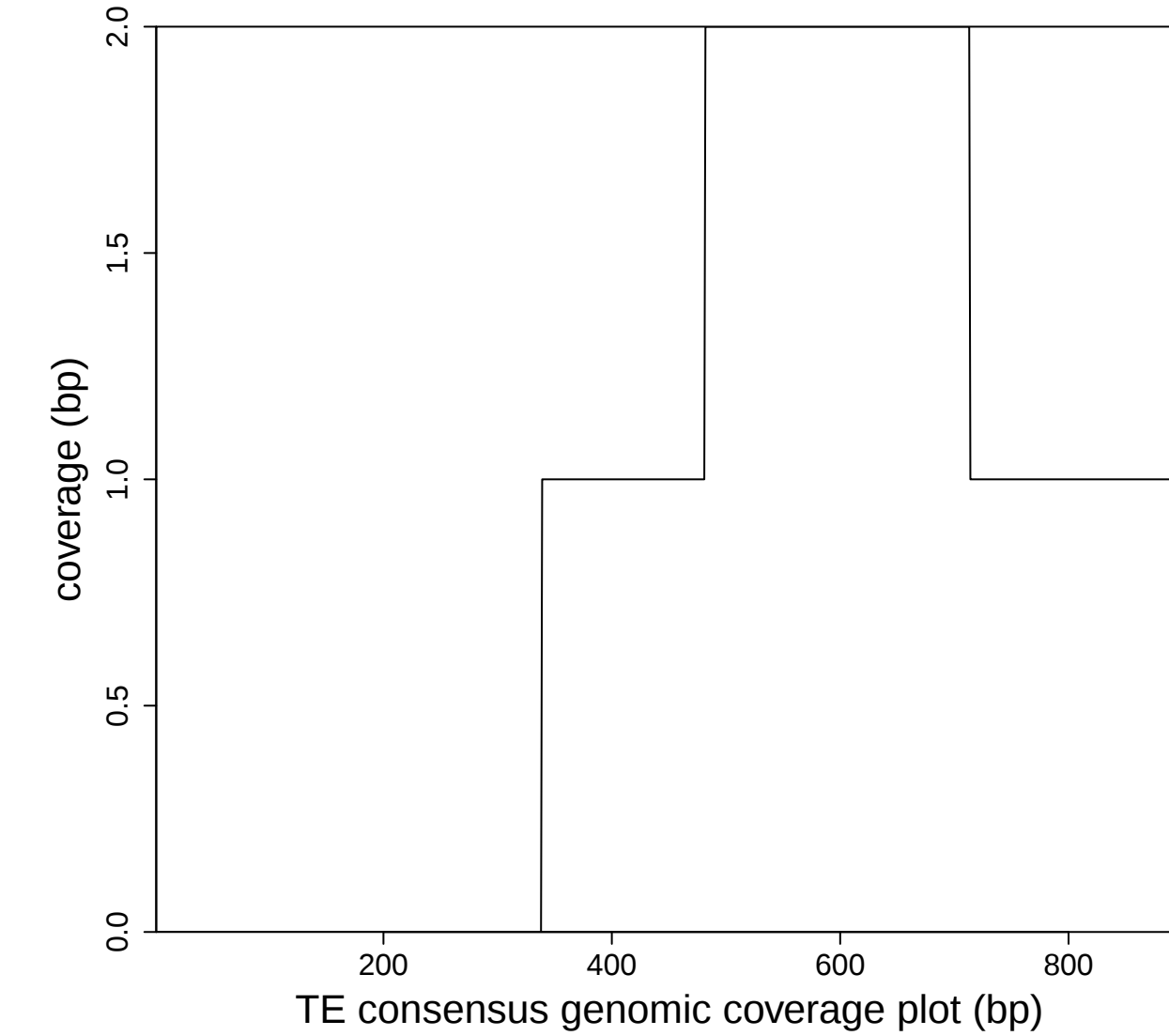
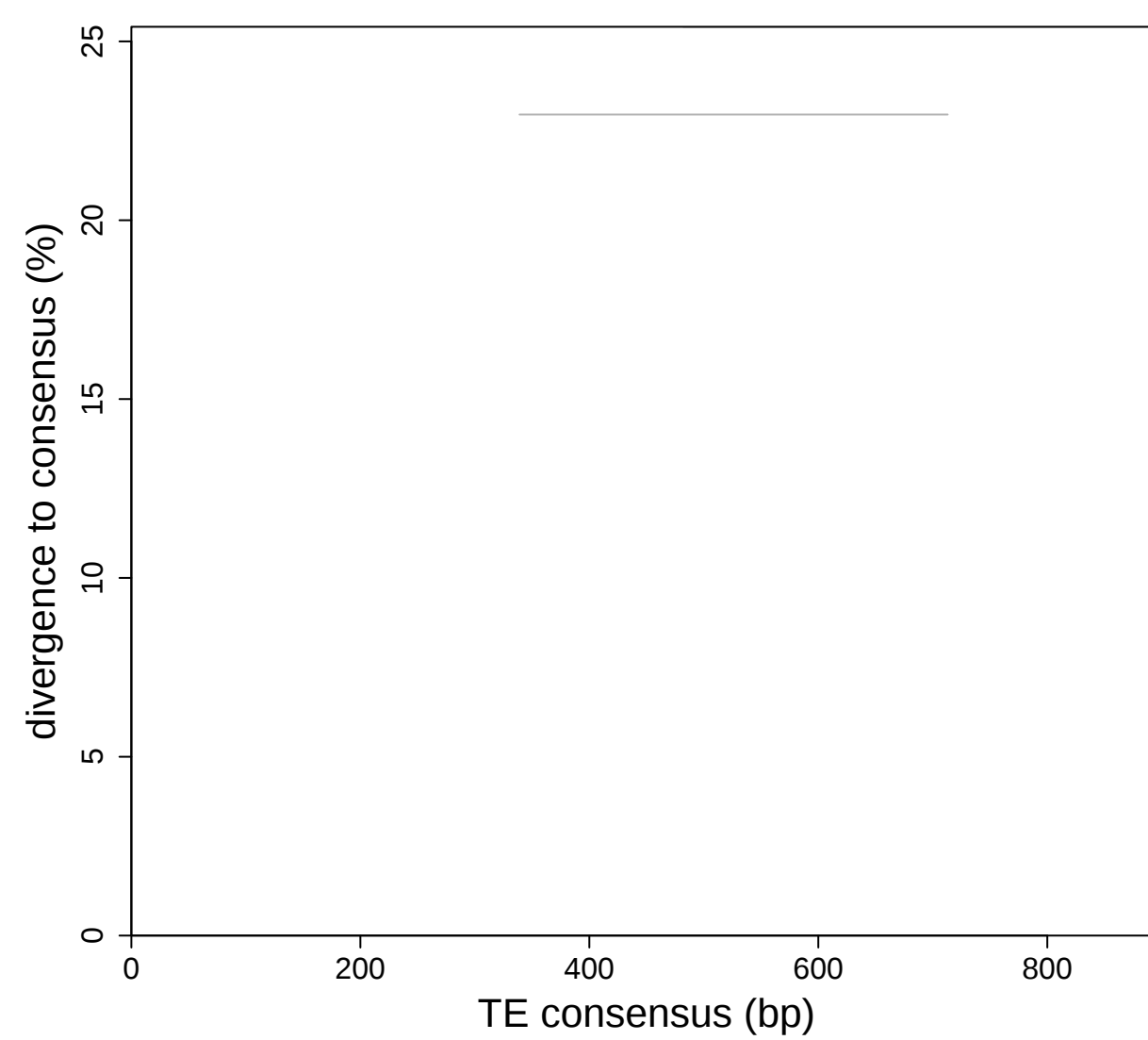


2.fasta.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 2394bp; fragments: 1449; full length: 0 (>=2154.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

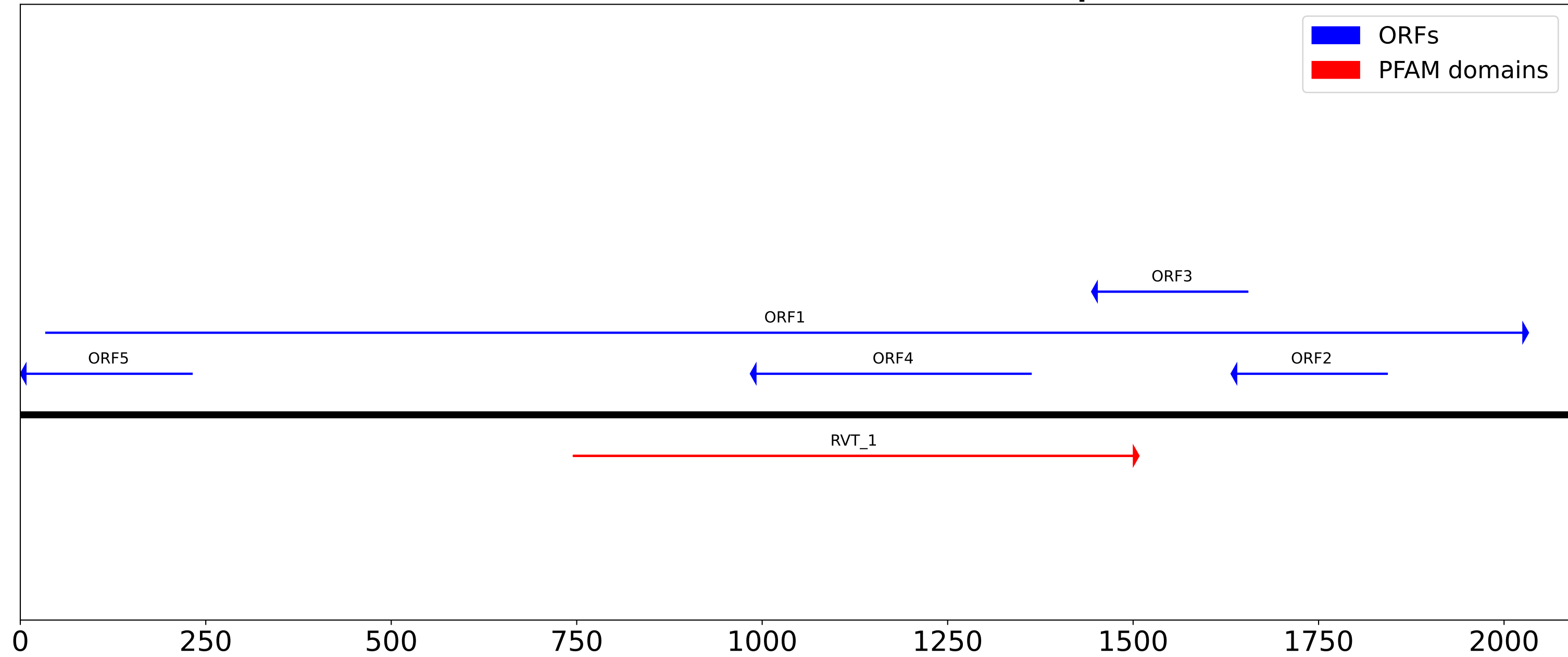


TE: rnd_4_family_1612
size: 900bp; fragments: 2; full length: 0 (≥ 810 bp)

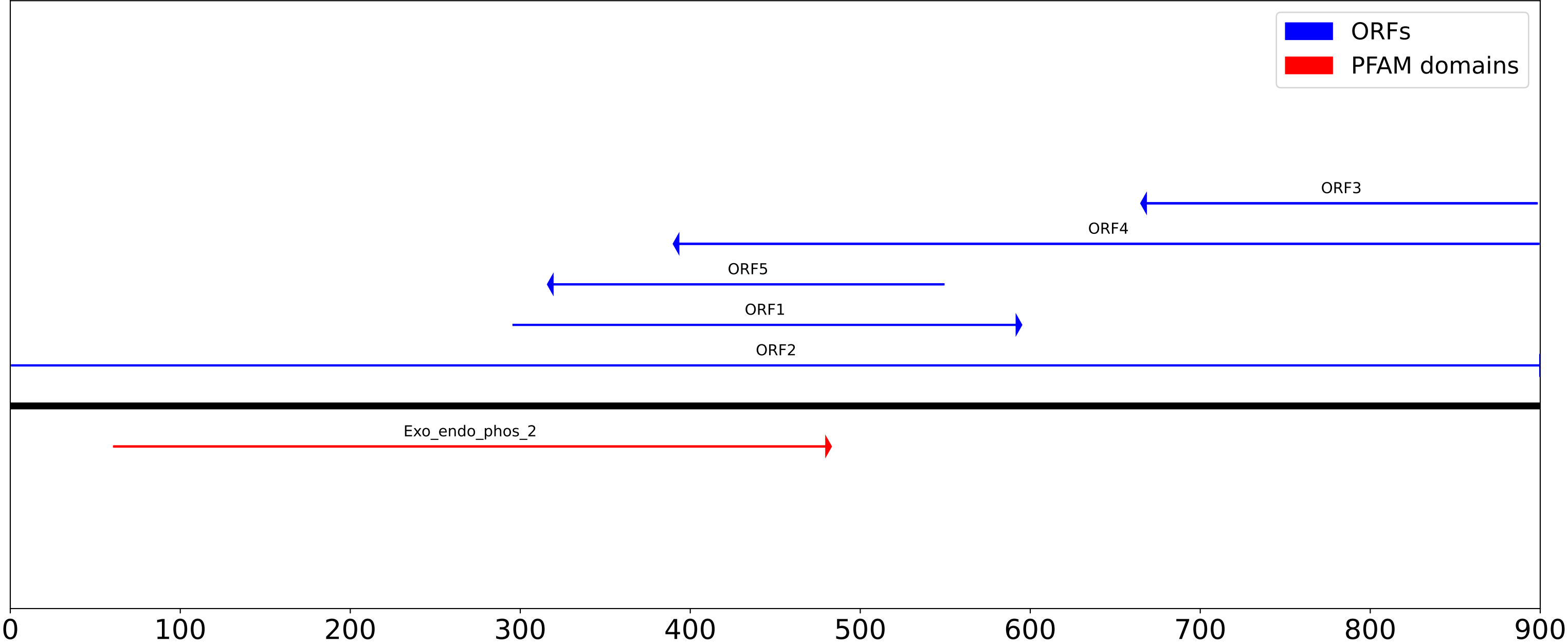


Before TEtrimmer 900 bp

After TEtrimmer ORF and PFAM domain plot



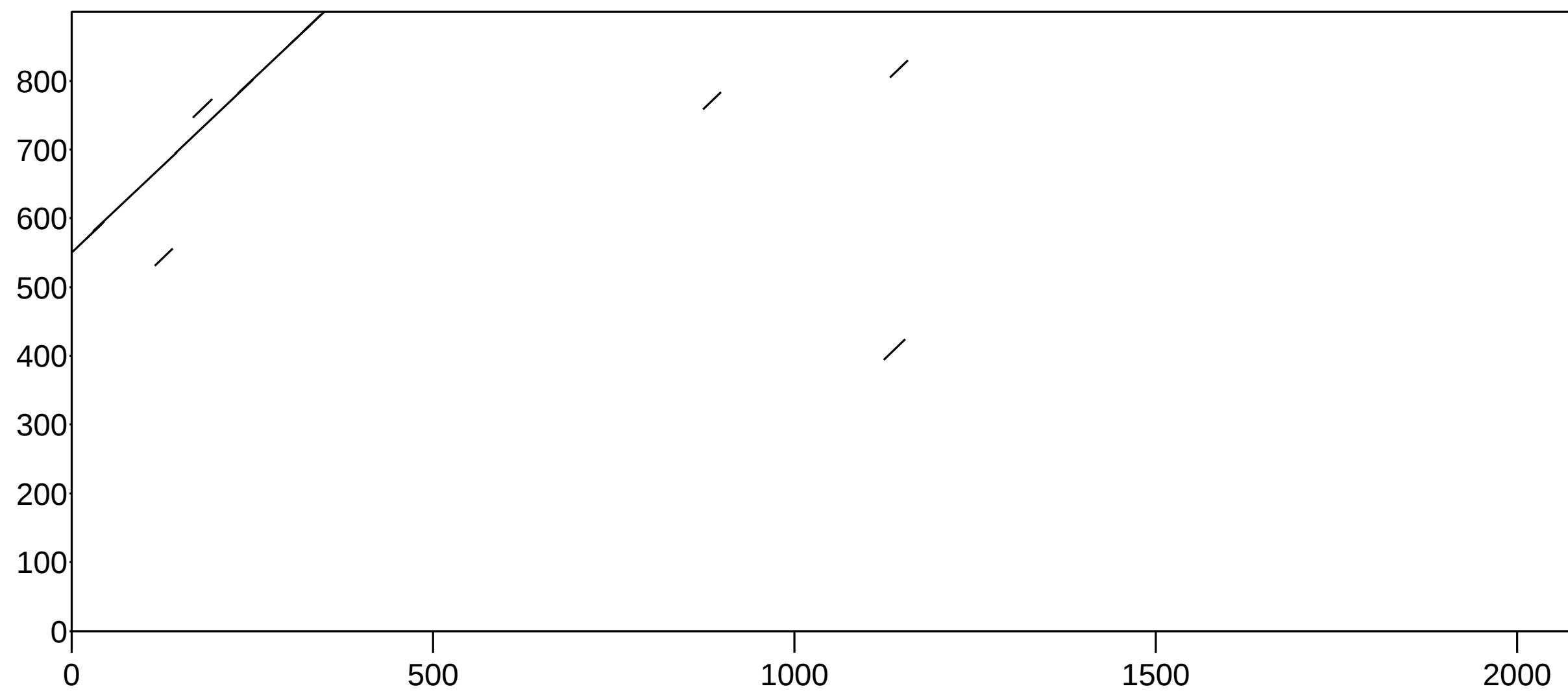
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)