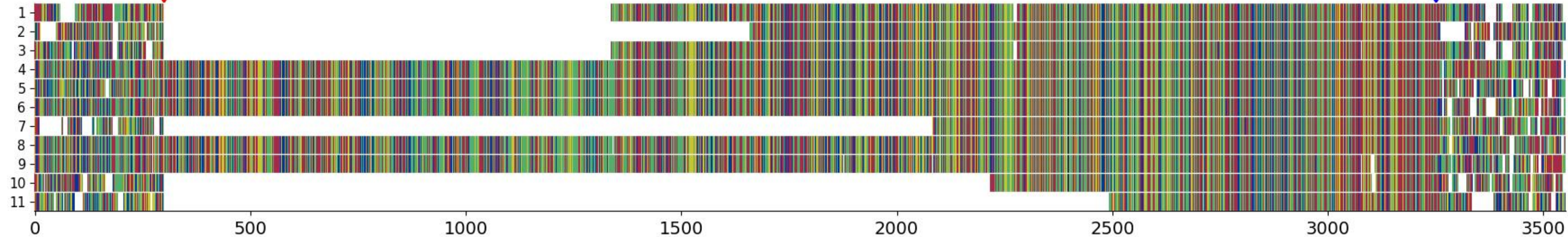


eng

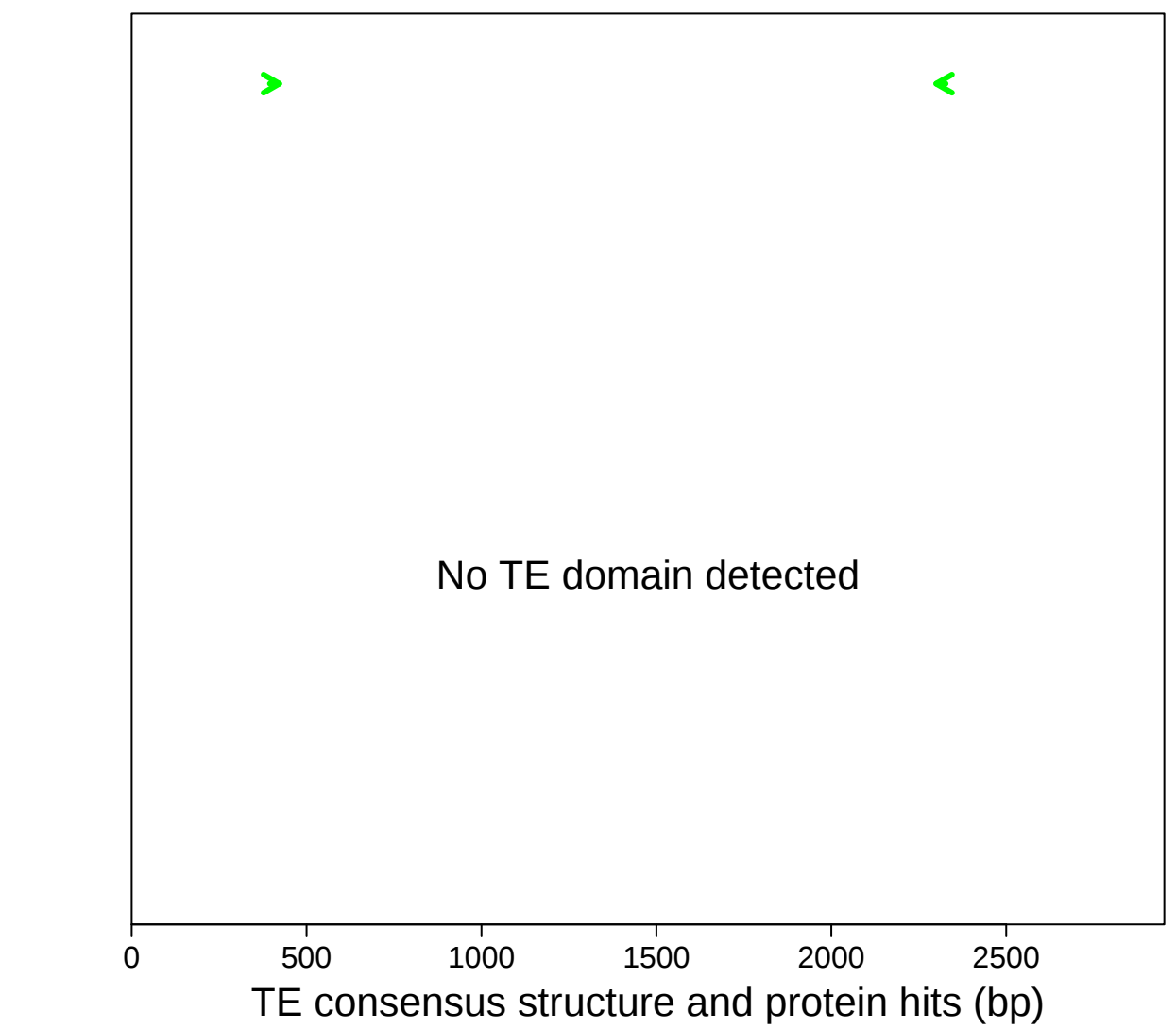
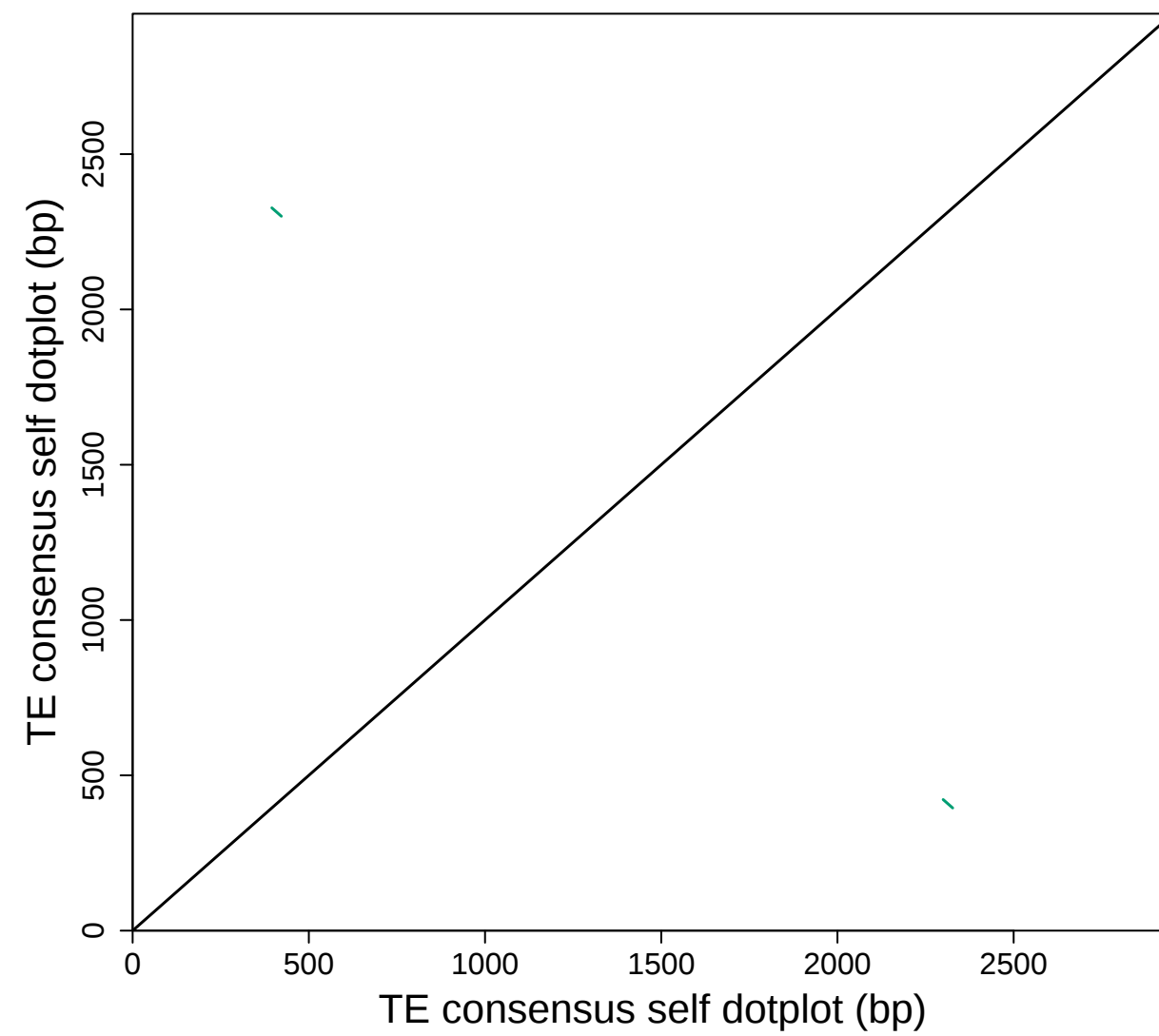
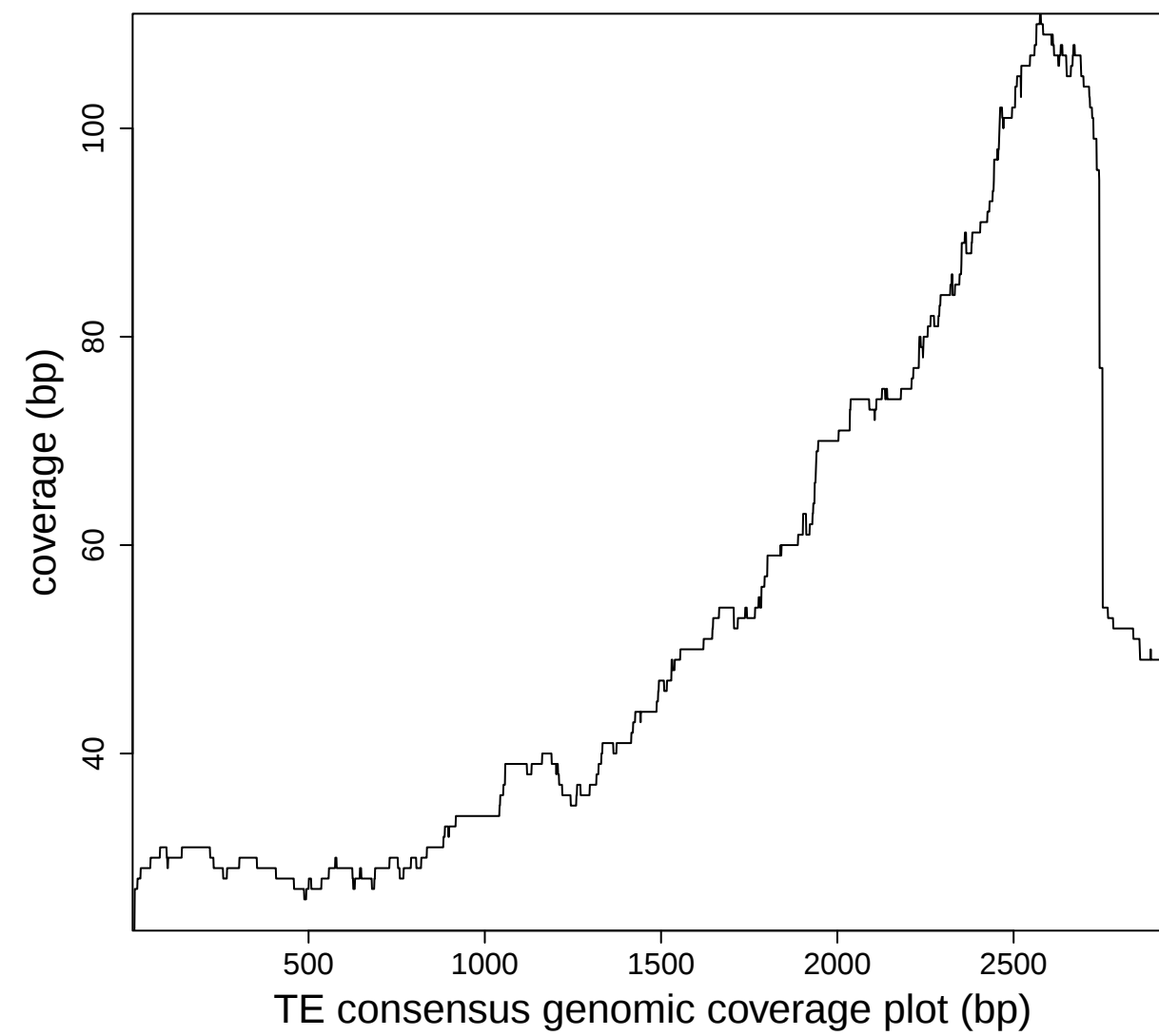
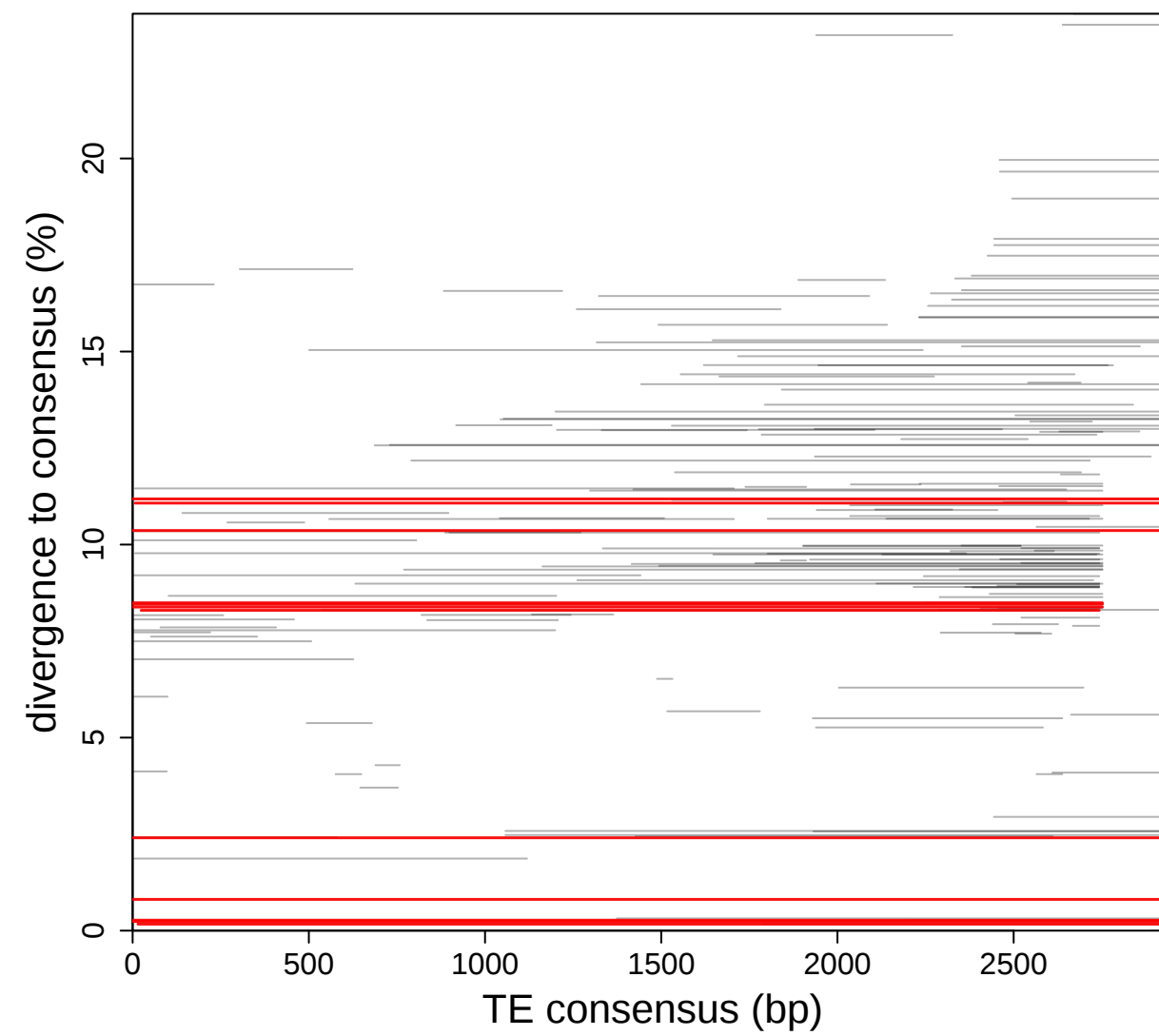
Start crop Point

End crop Point



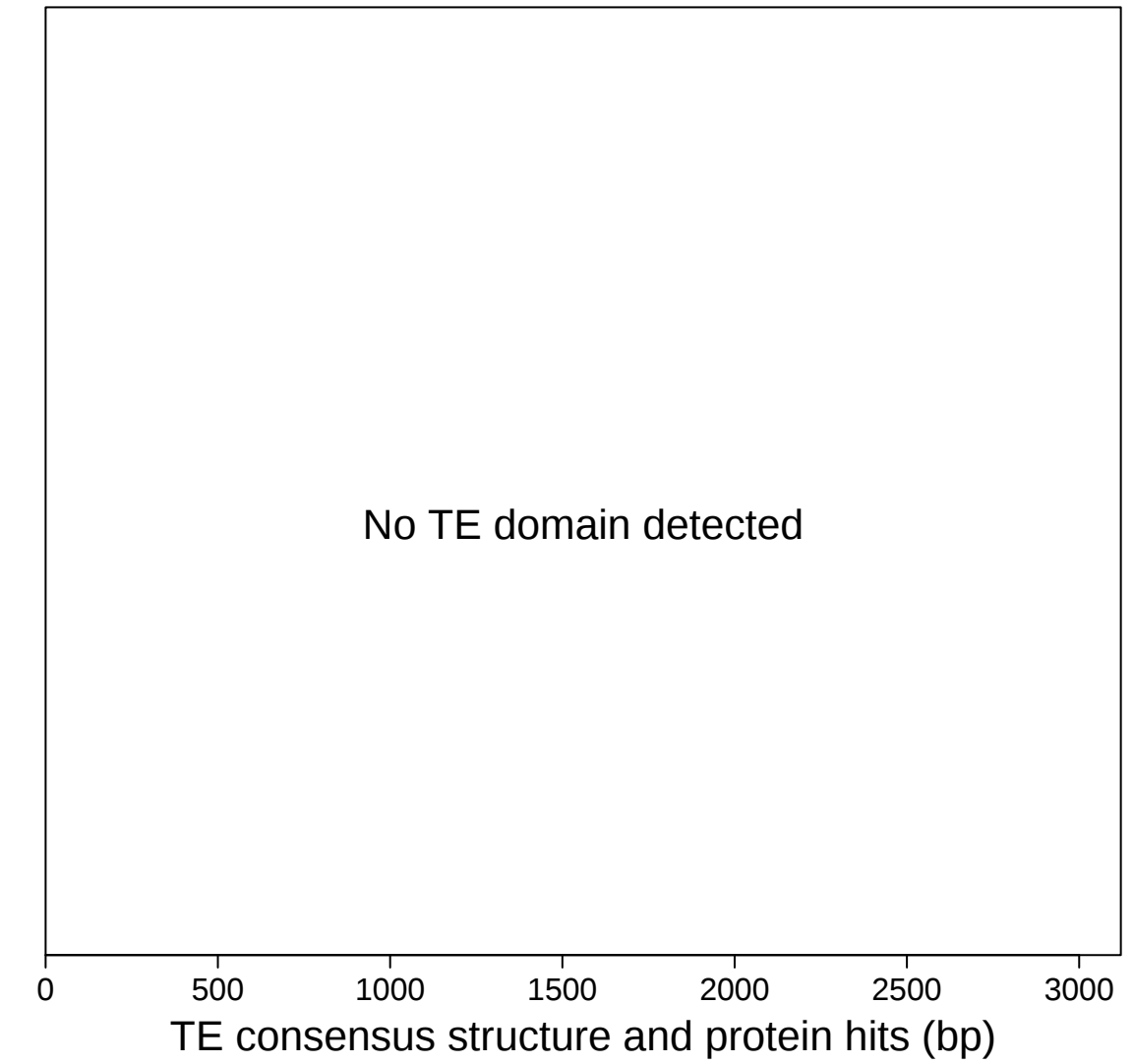
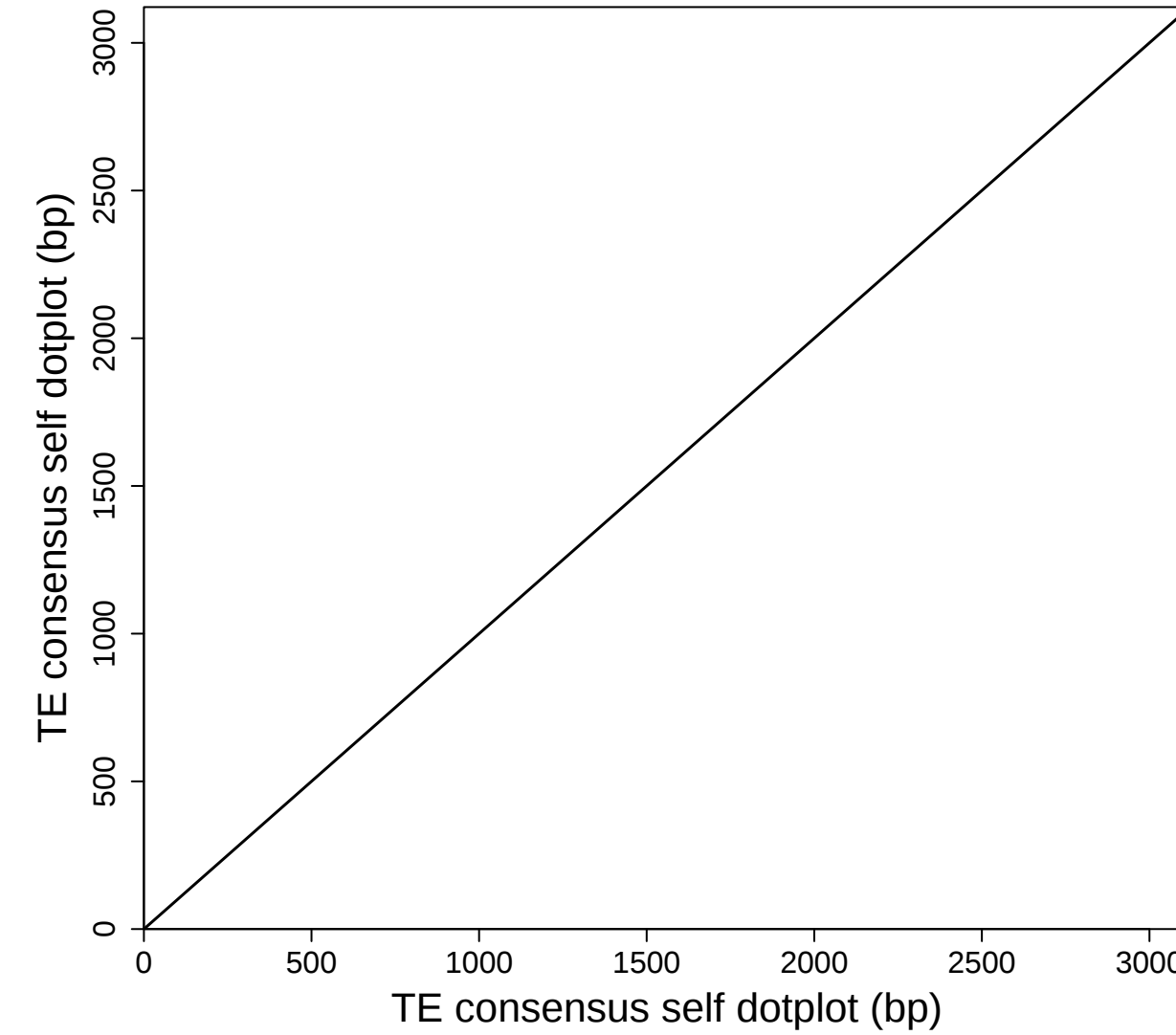
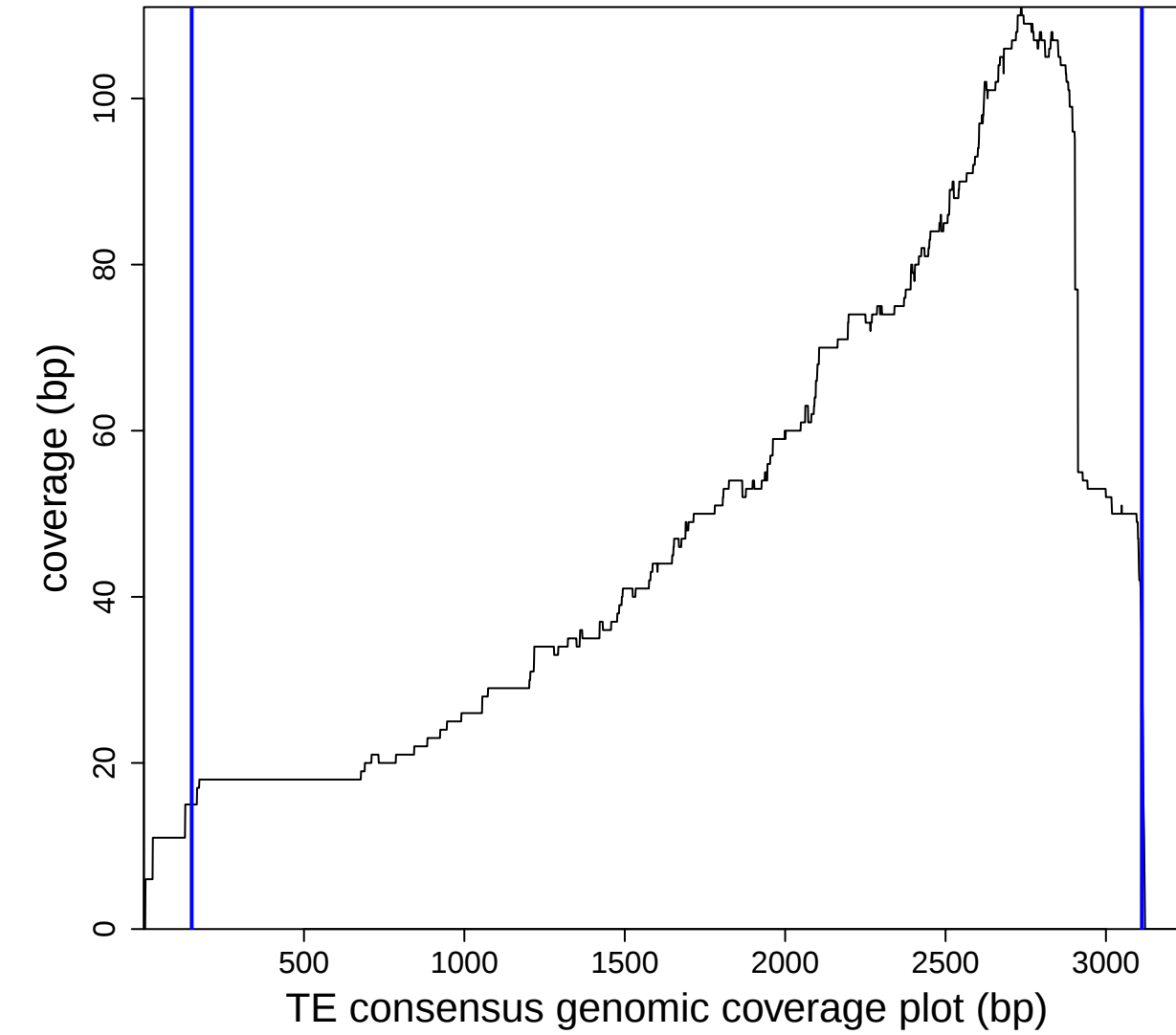
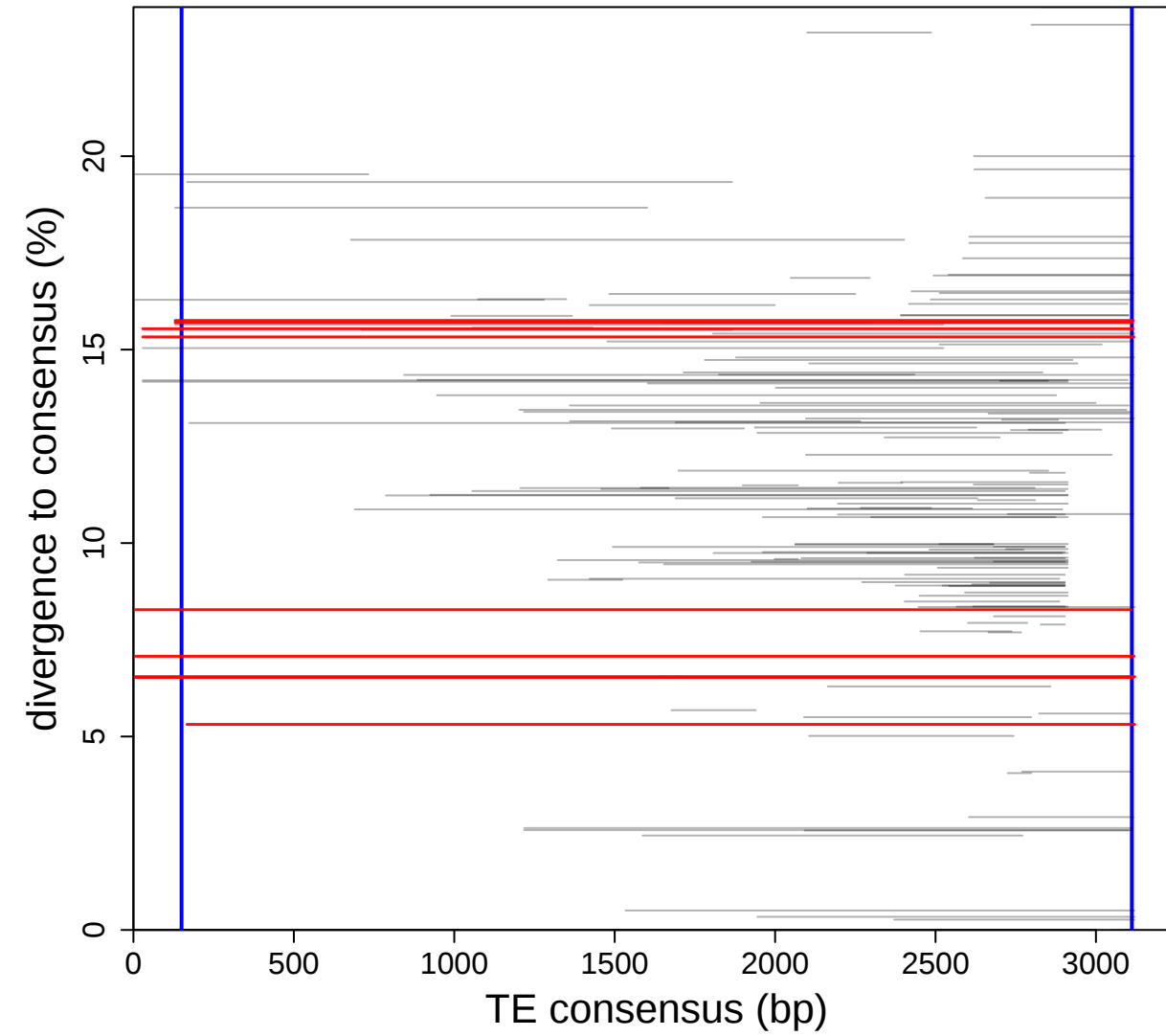
bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 2952bp; fragments: 175; full length: 12 (>=2656.8bp)

After TEtrimmer 2952 bp

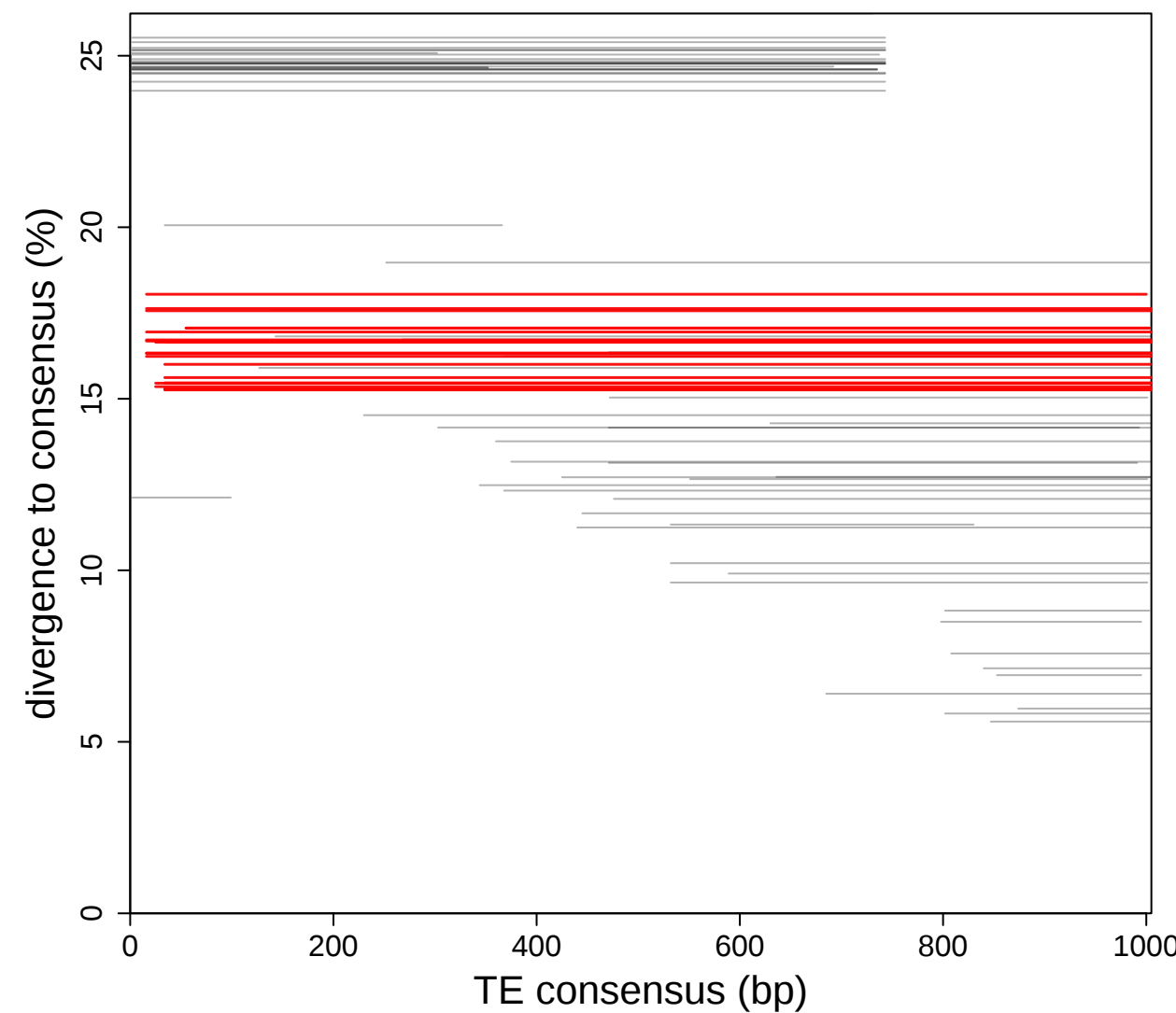


3.fasta.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.fa
size: 3262bp; fragments: 152; full length: 9 (>=2935.8bp)

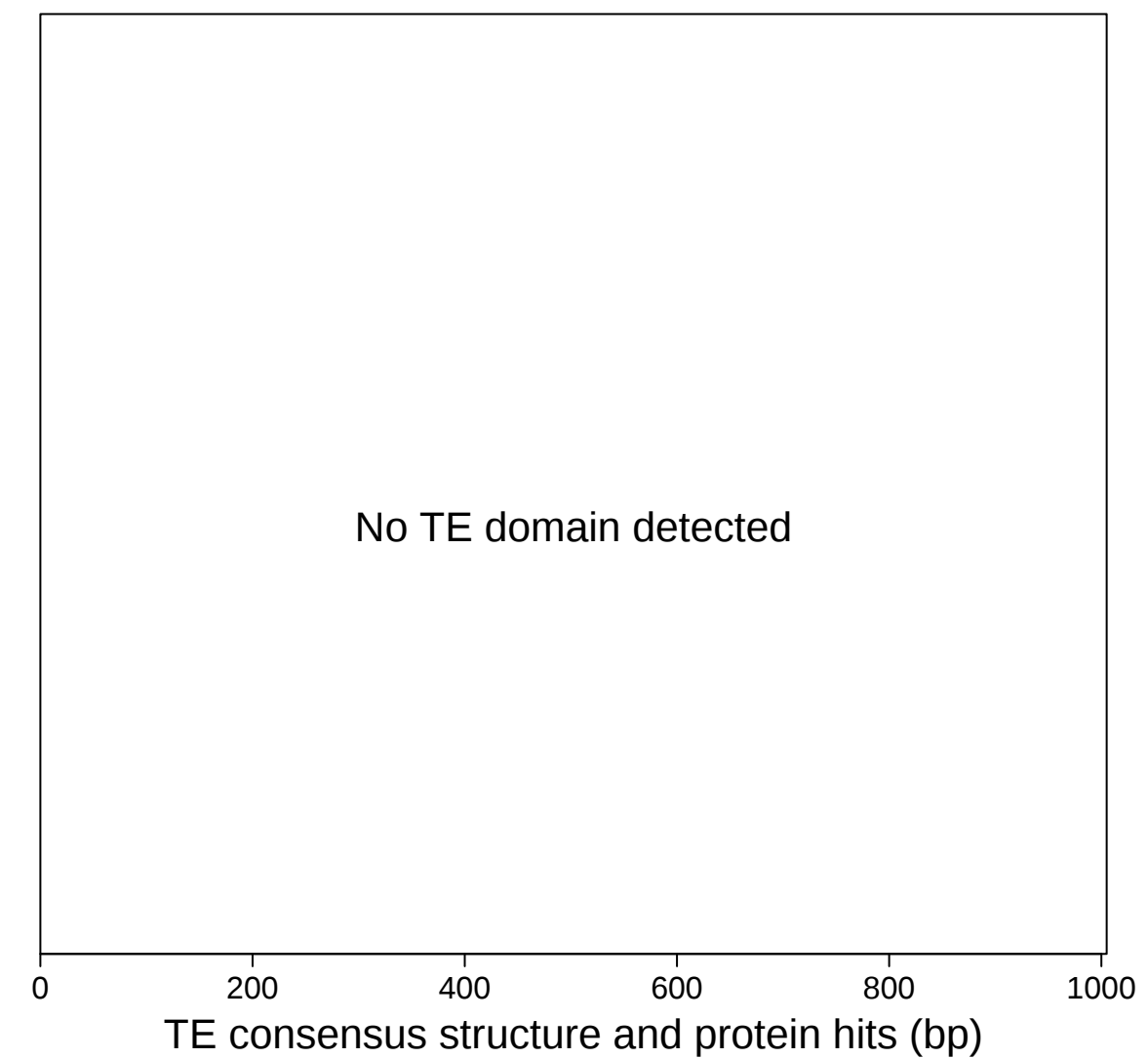
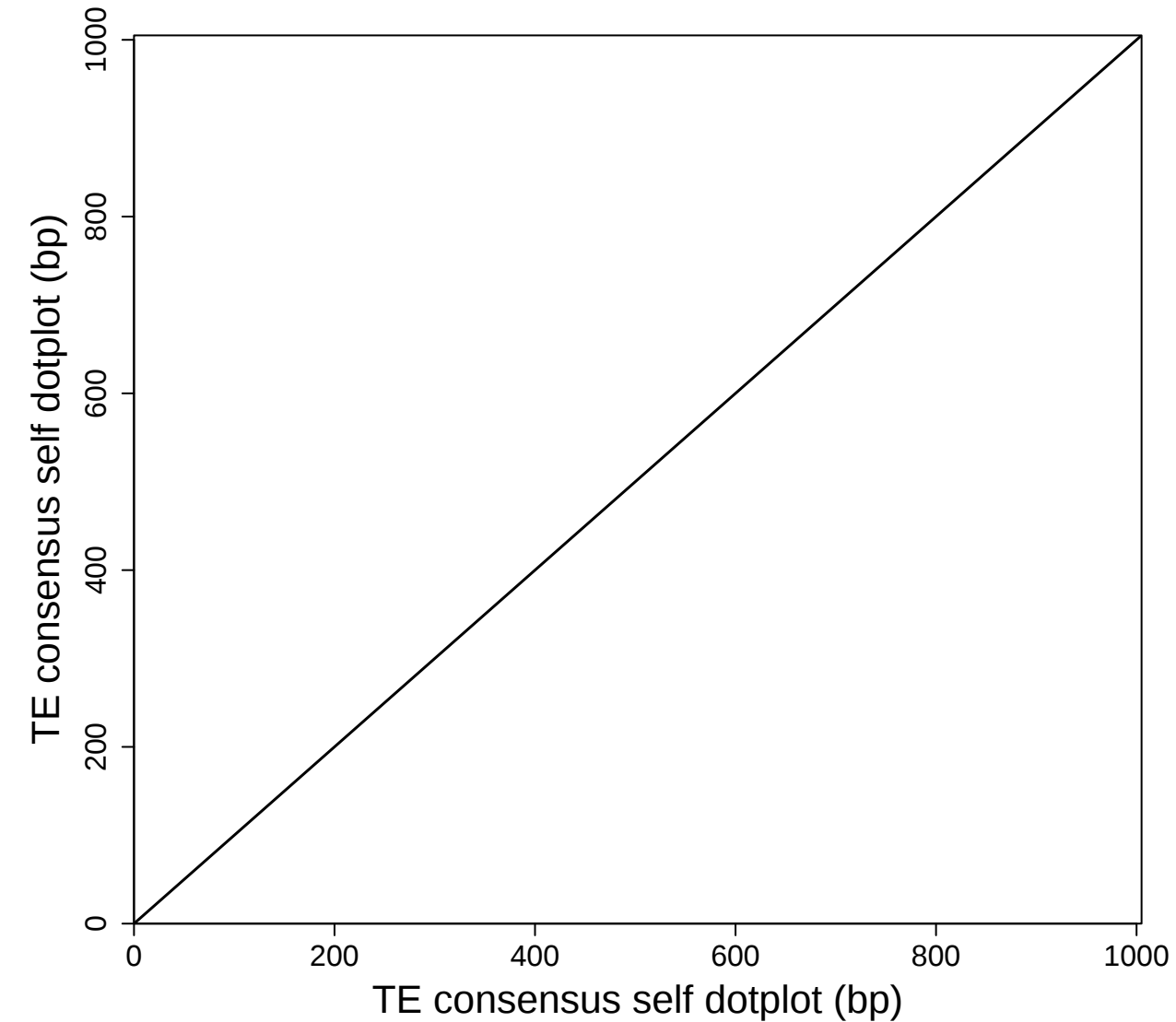
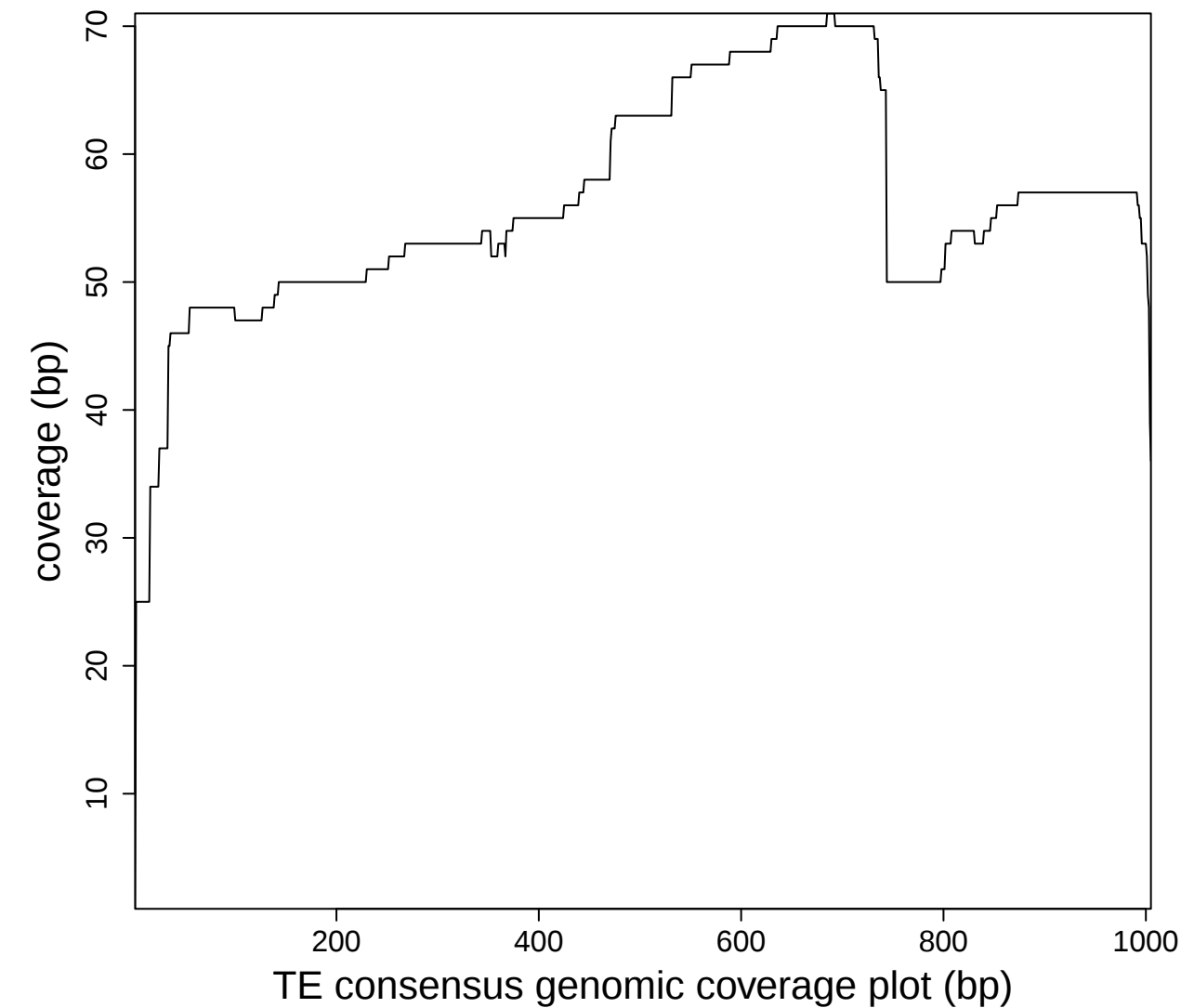
After TEtrimmer Extended plot Blue lines are boundaries



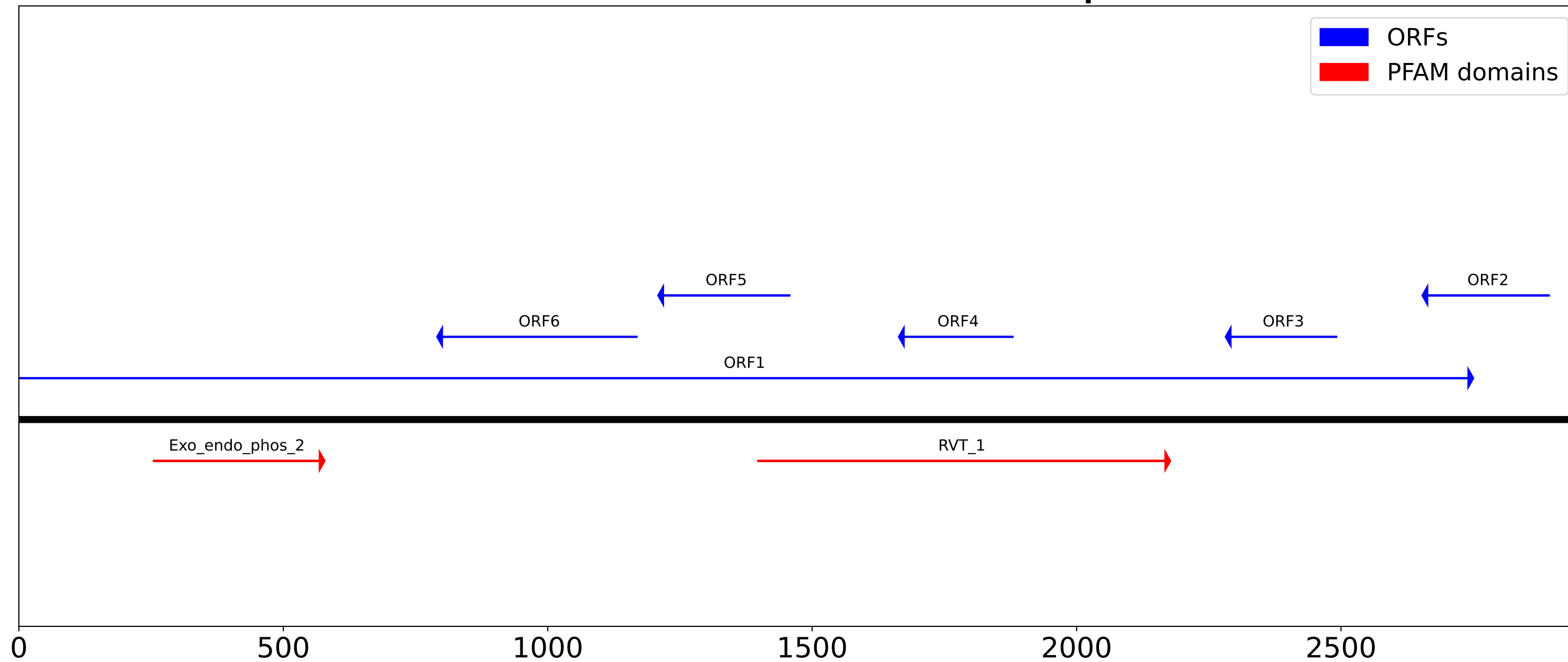
TE: rnd_4_family_143
size: 1005bp; fragments: 84; full length: 22 (≥ 904.5 bp)



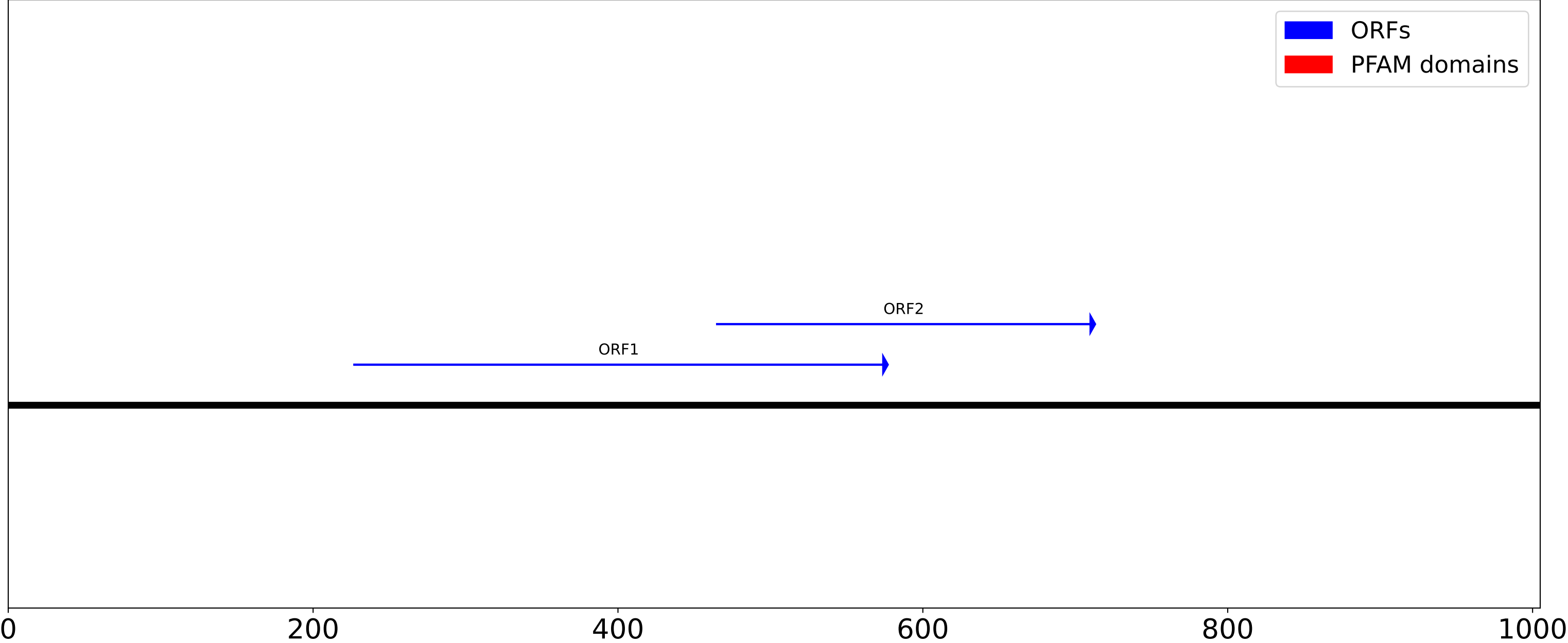
Before TEtrimmer 1005 bp



After TEtrimmer ORF and PFAM domain plot



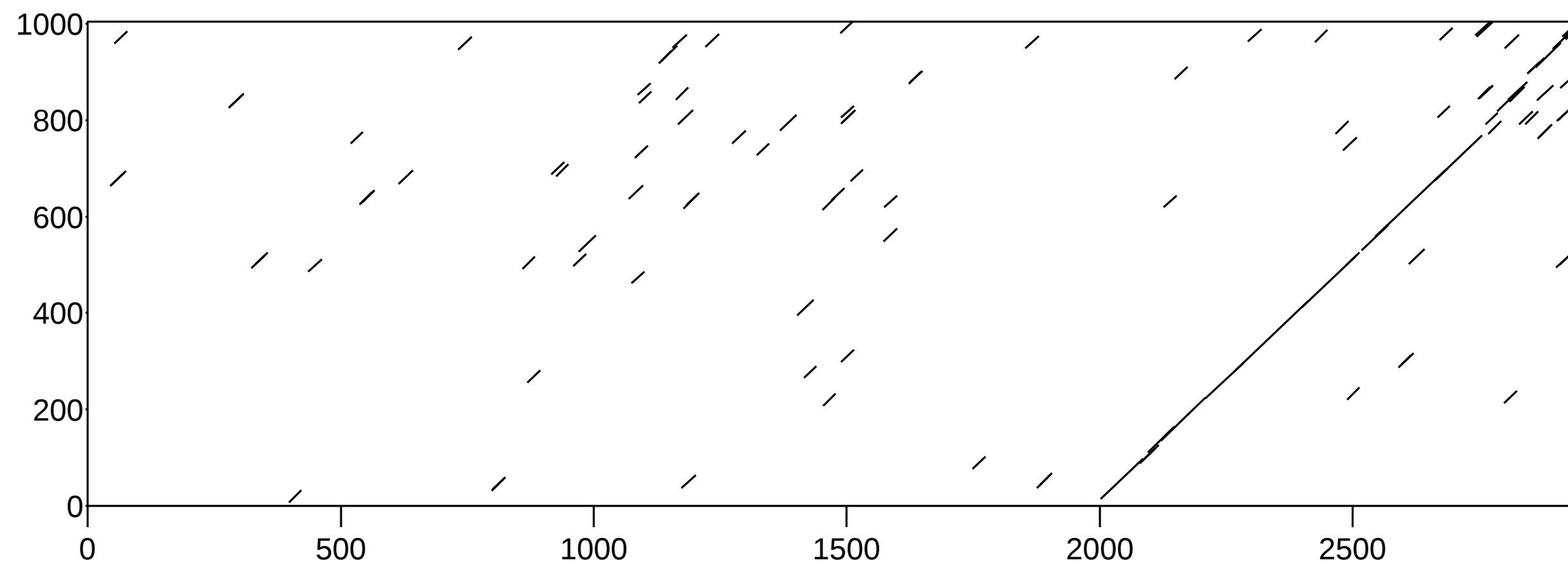
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

E consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)