

Start crop Point

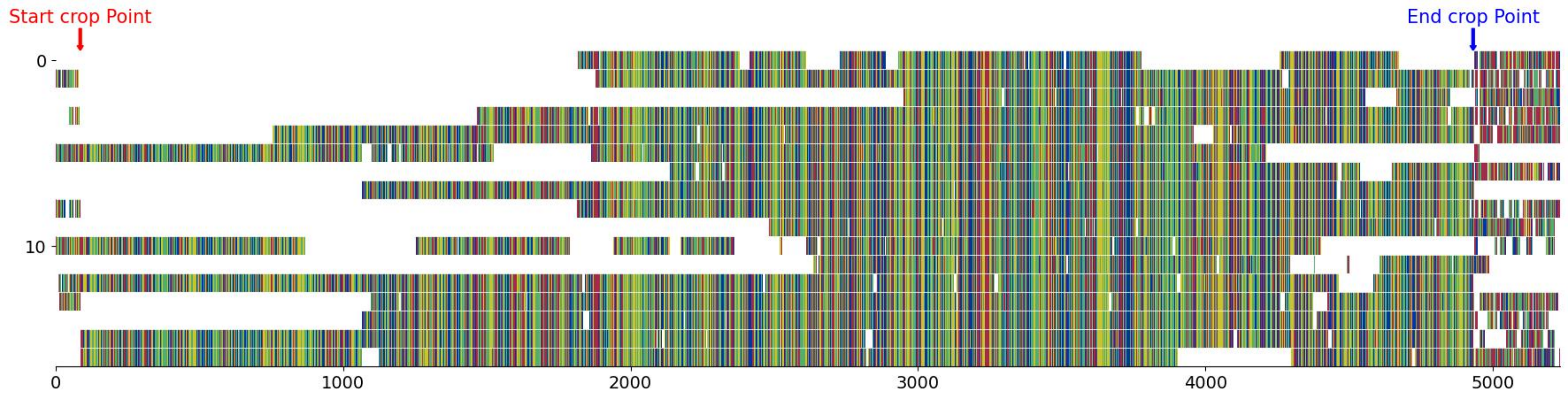


End crop Point



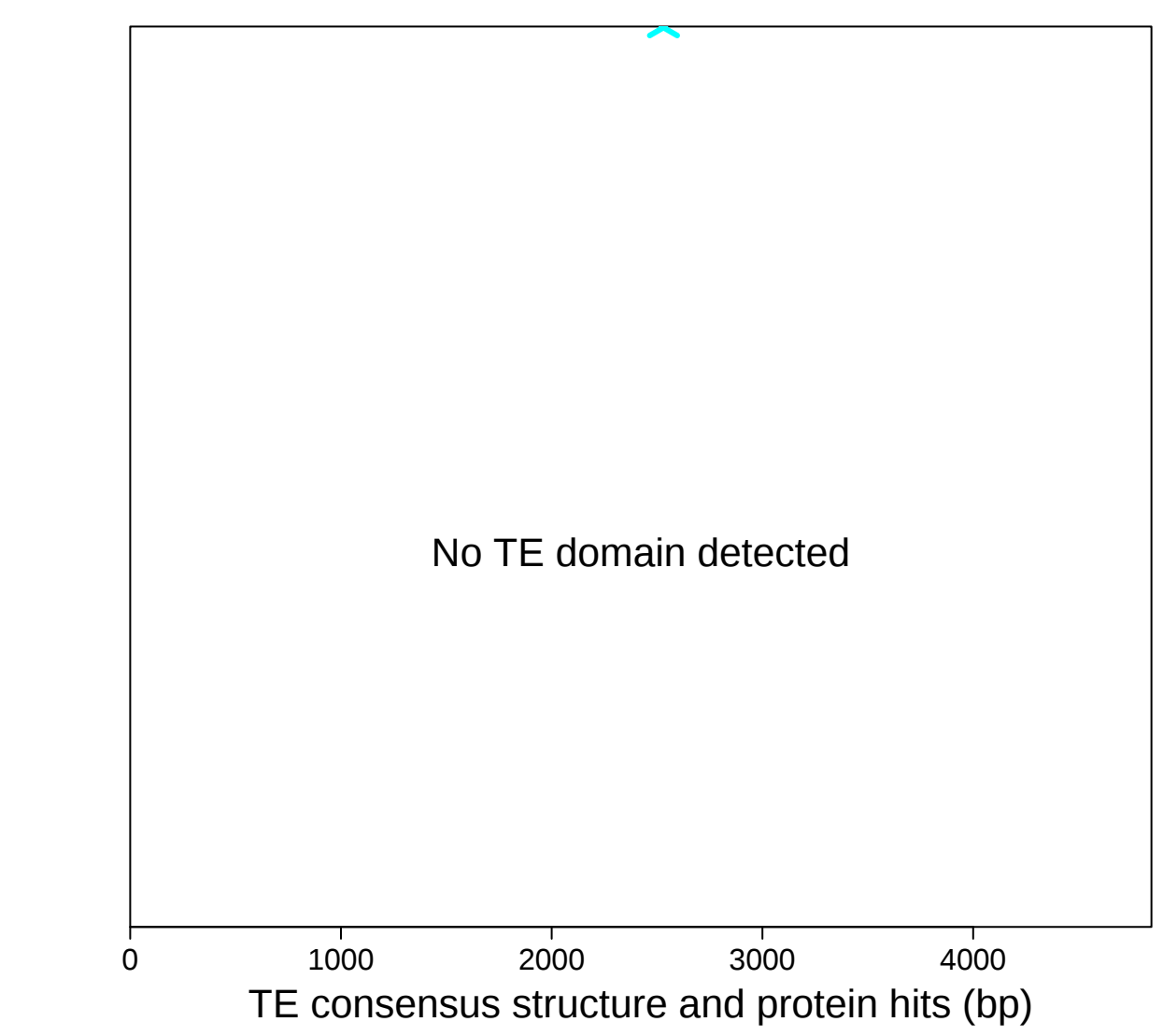
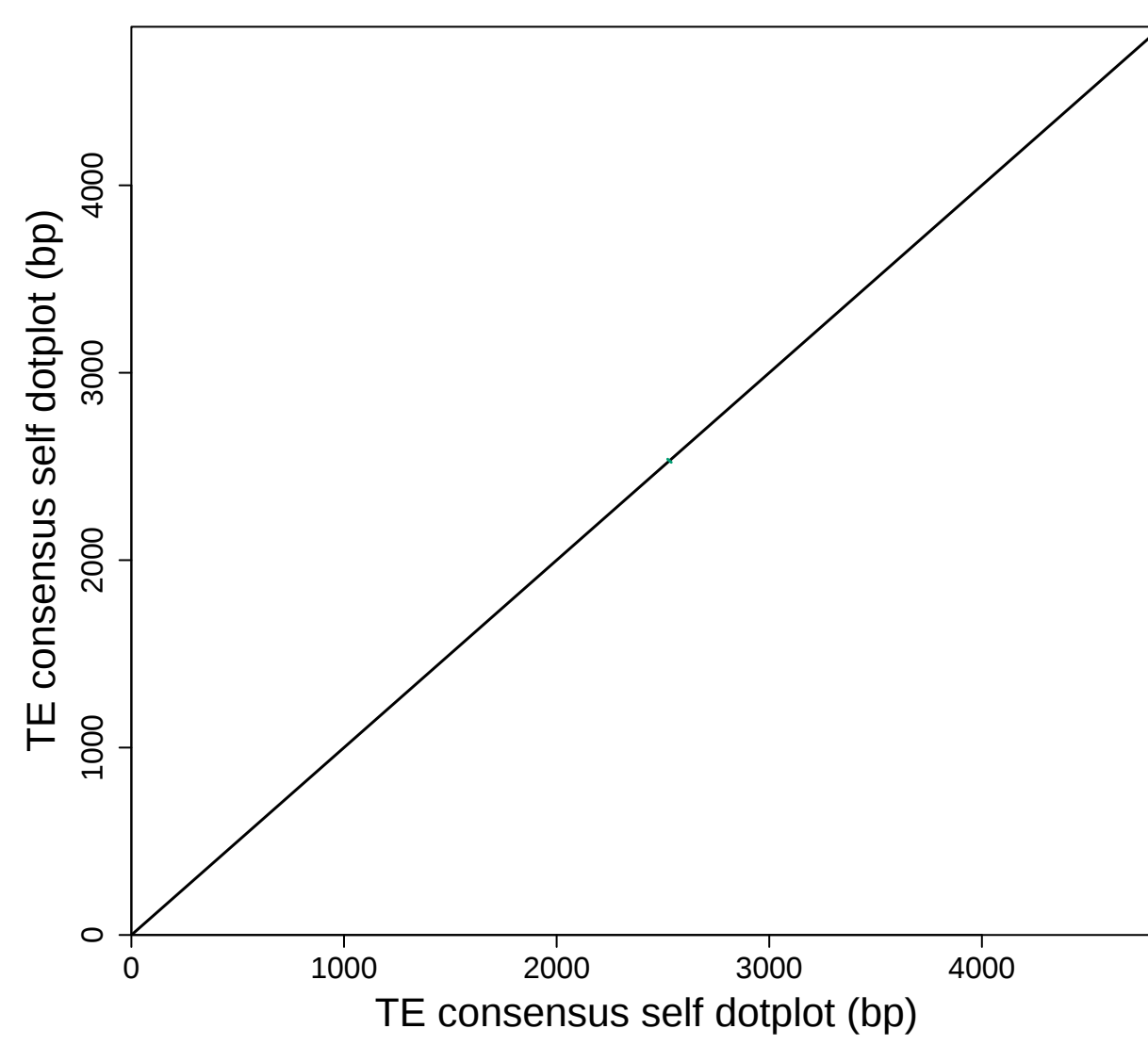
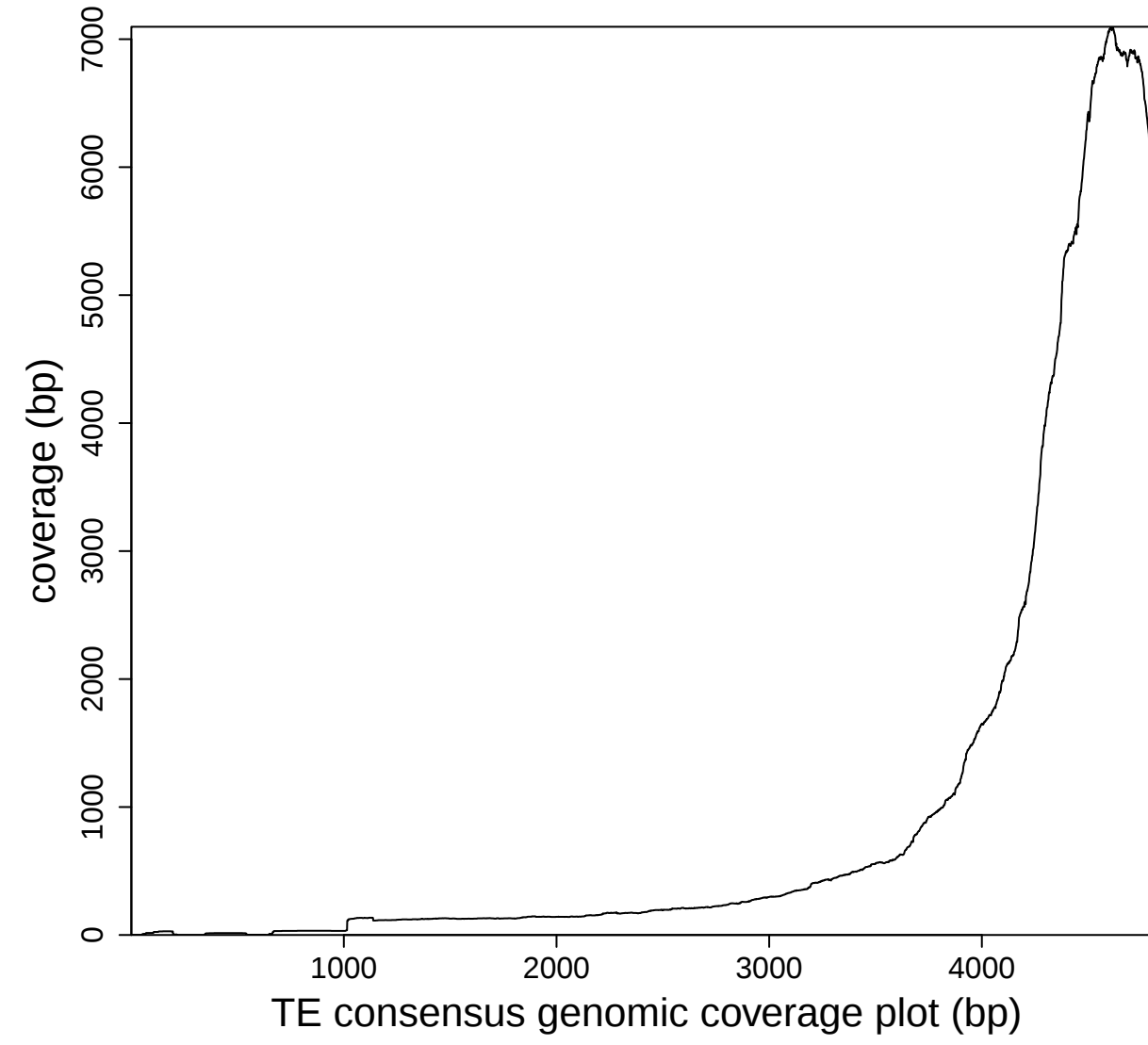
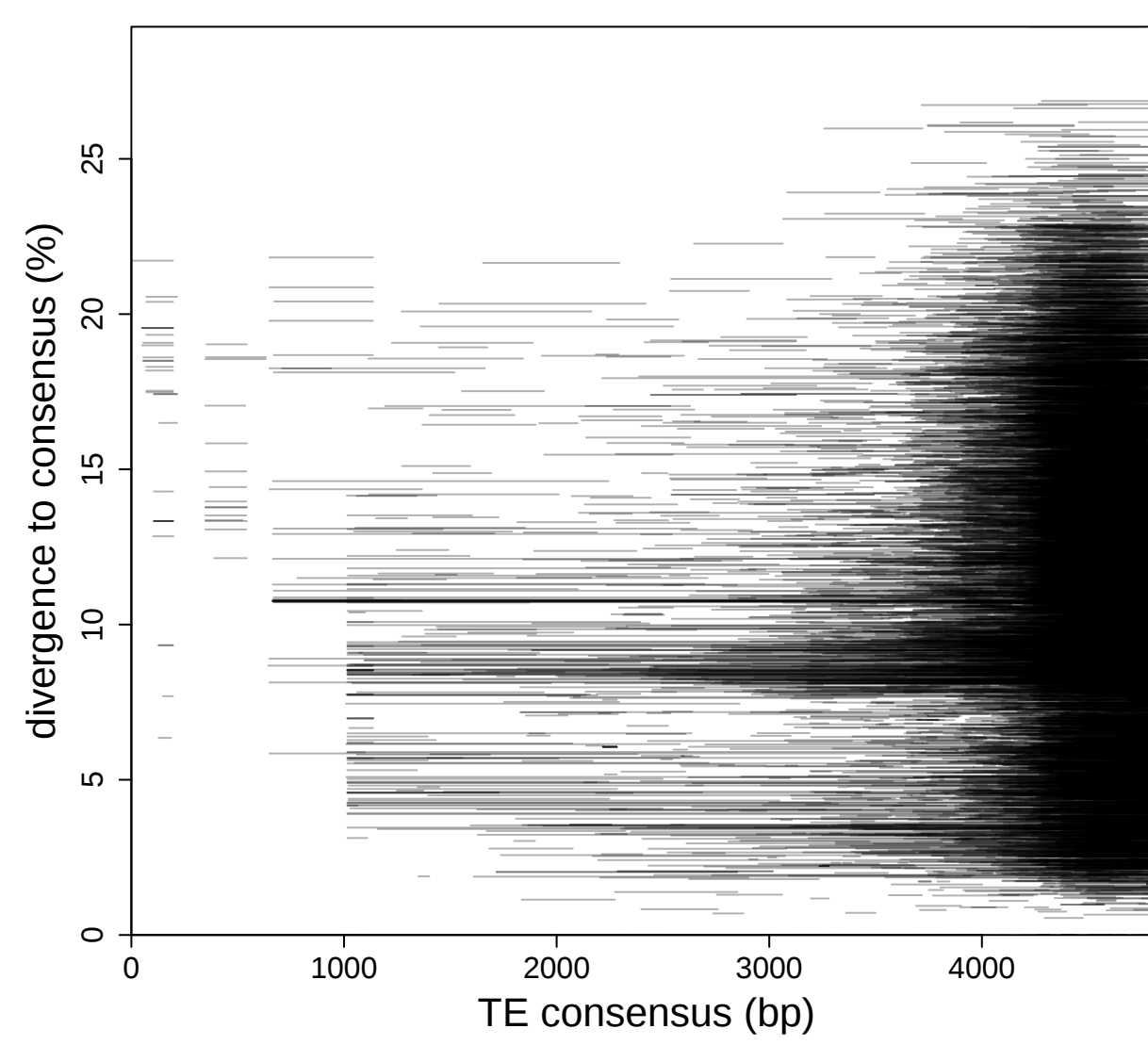
MSA length = 4846





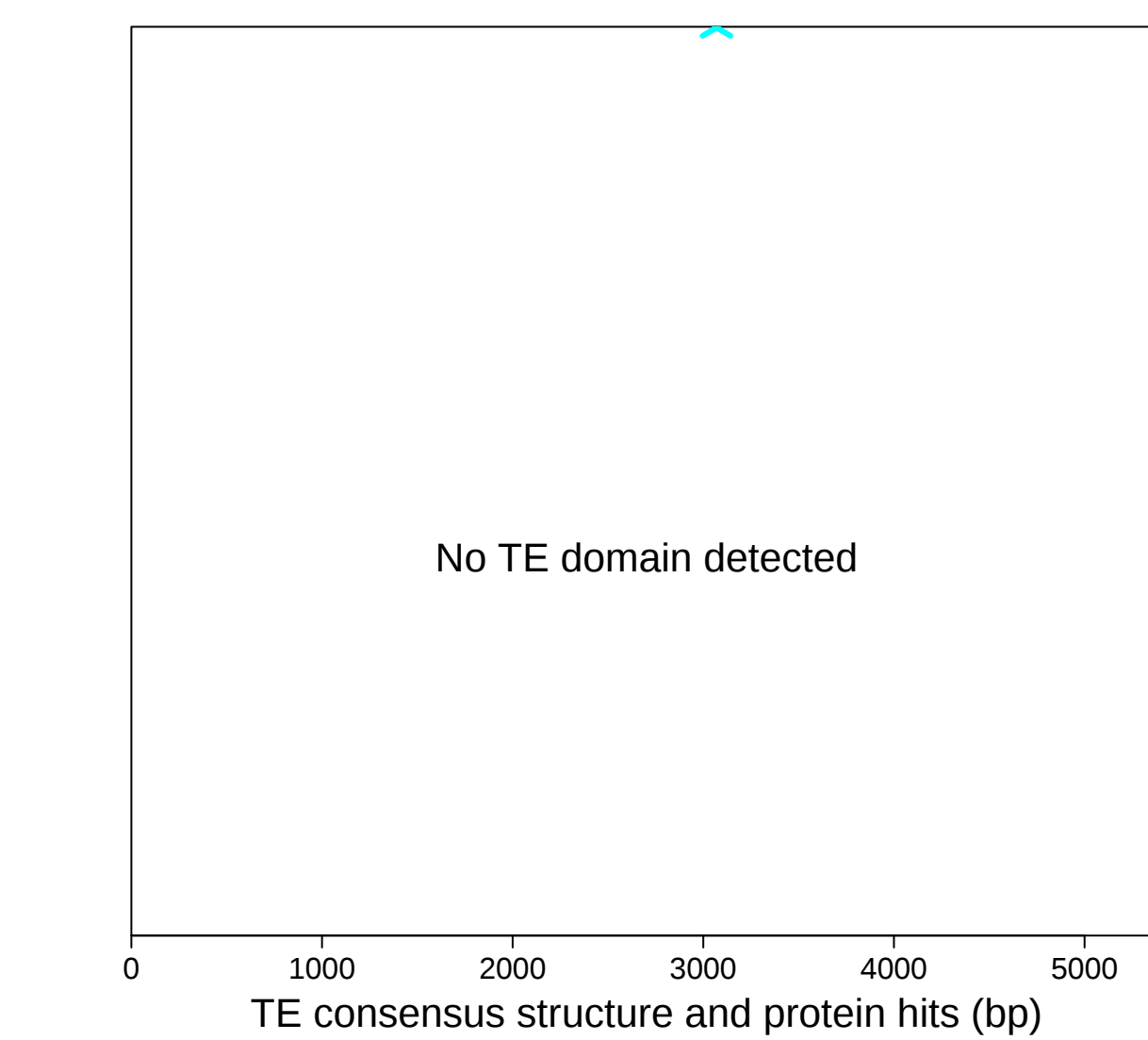
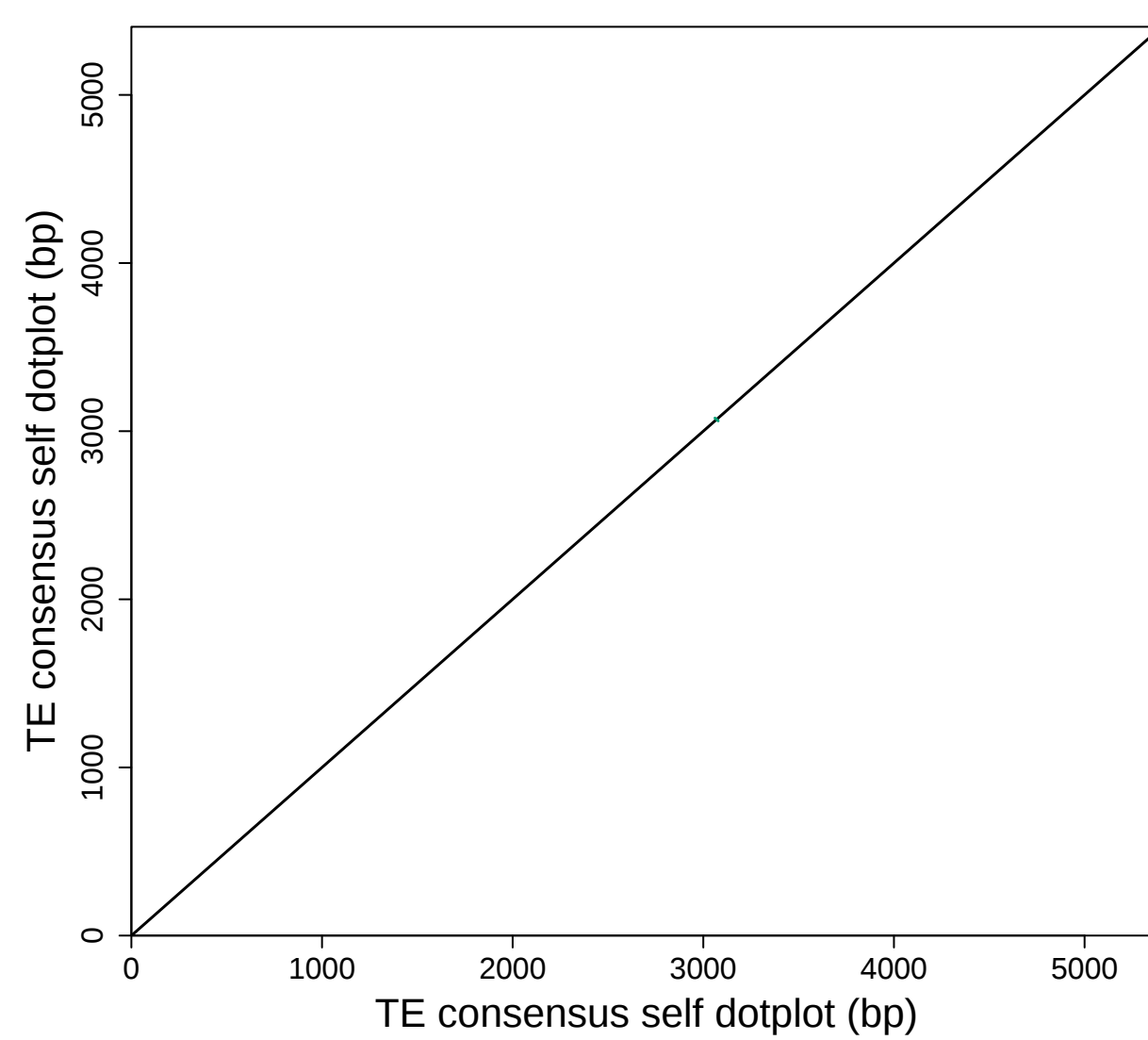
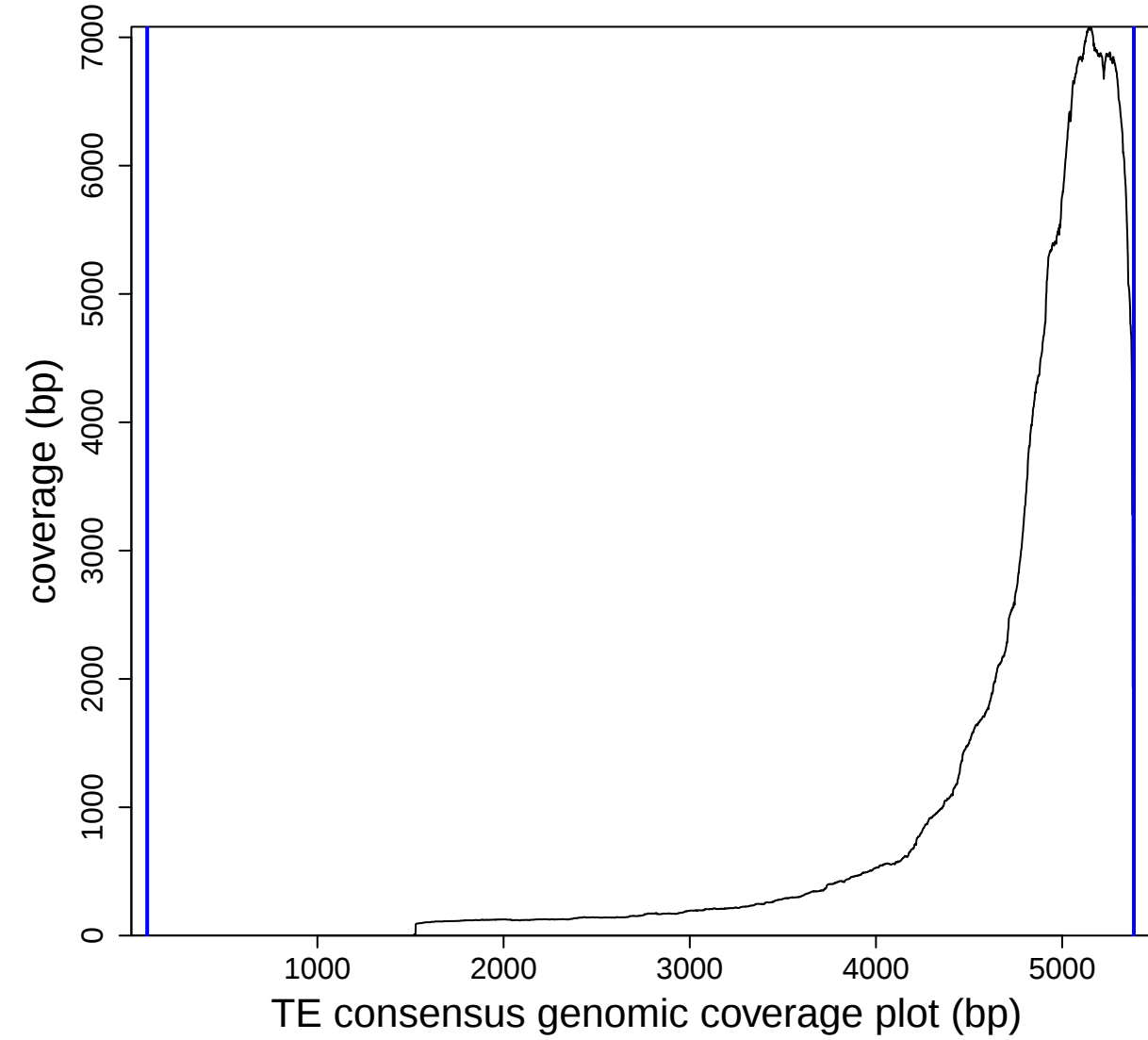
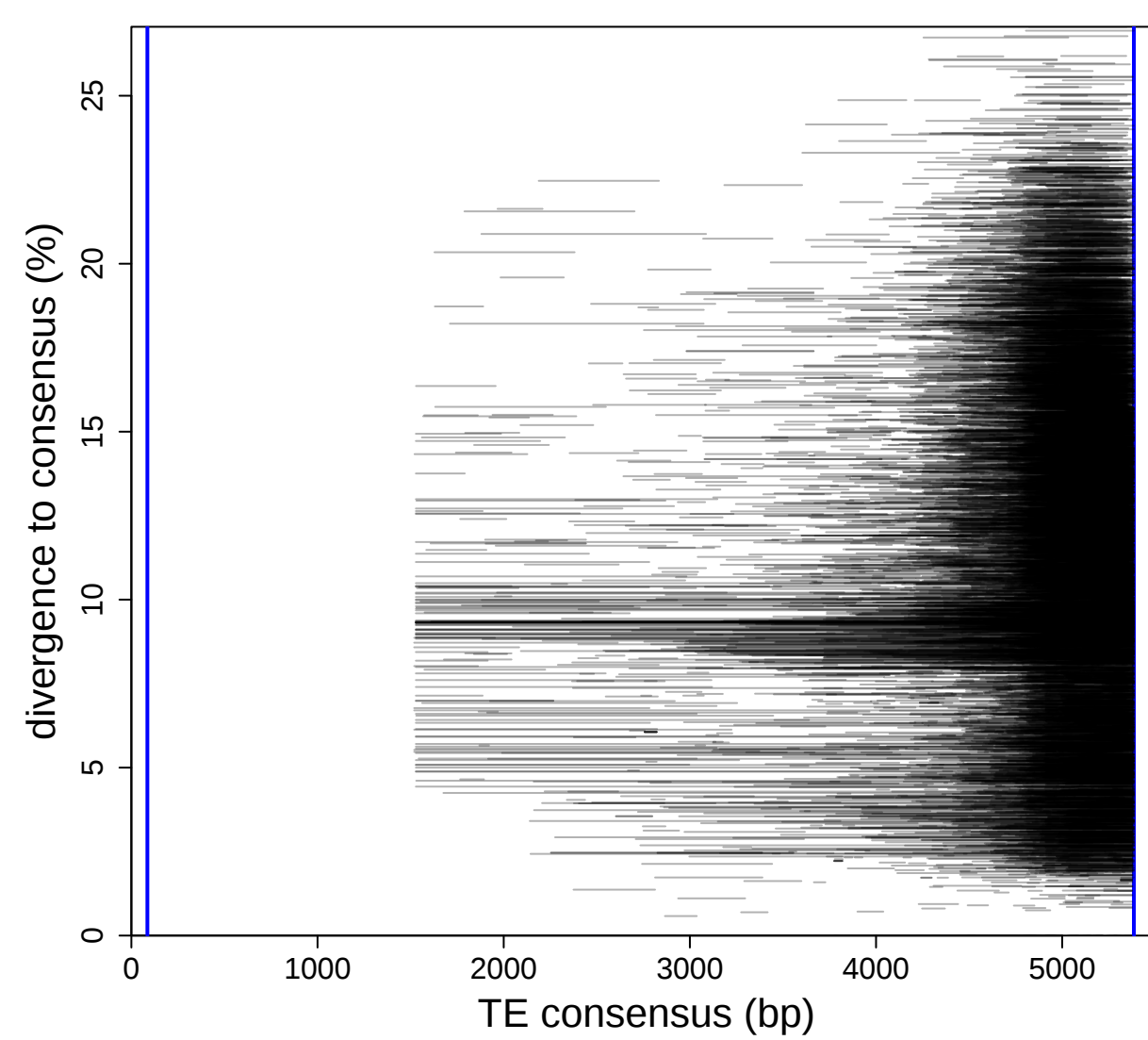
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 4846bp; fragments: 12240; full length: 0 (>=4361.4bp)

After TEtrimmer 4846 bp

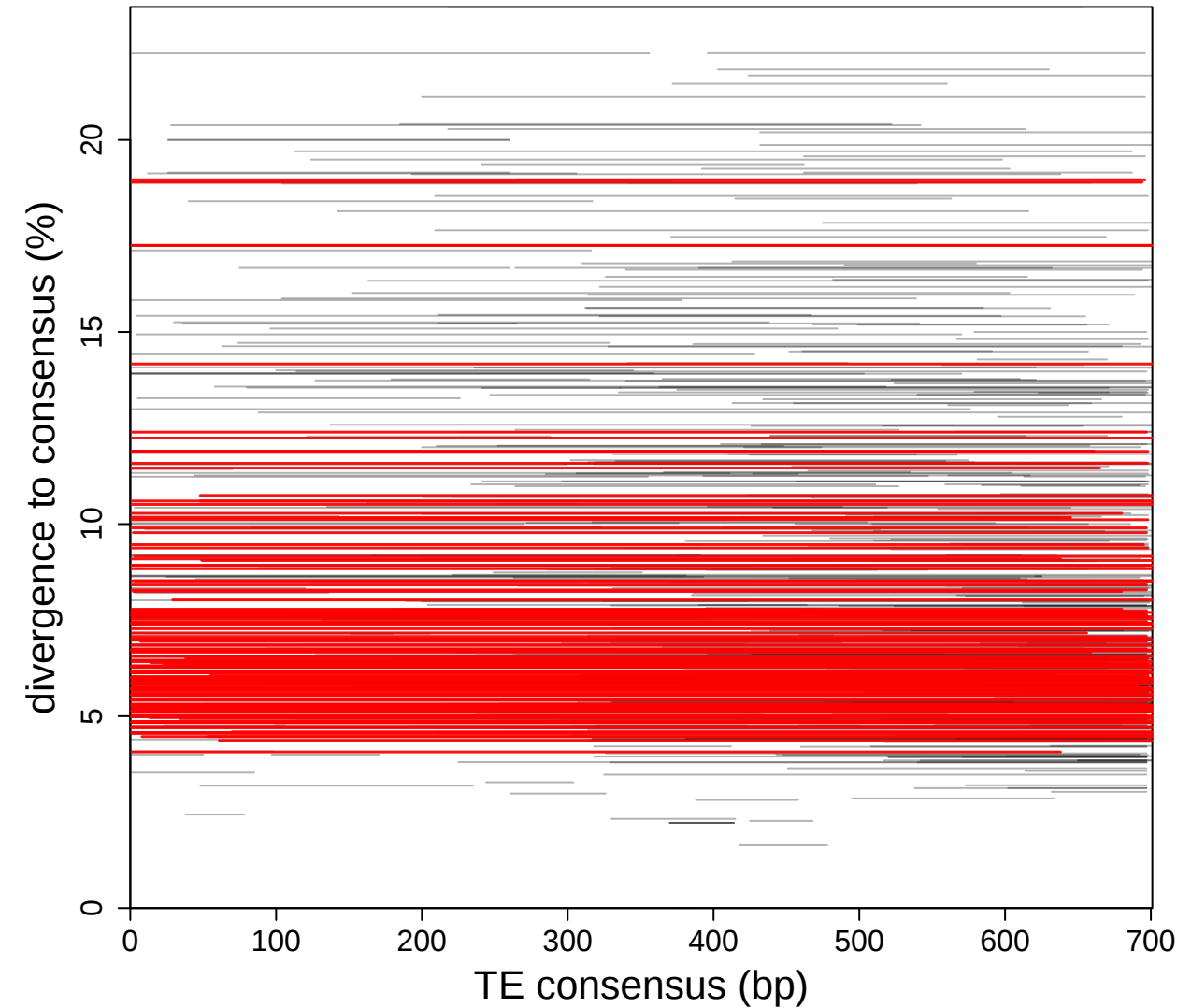


0.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 5535bp; fragments: 12116; full length: 0 (>=4981.5bp)

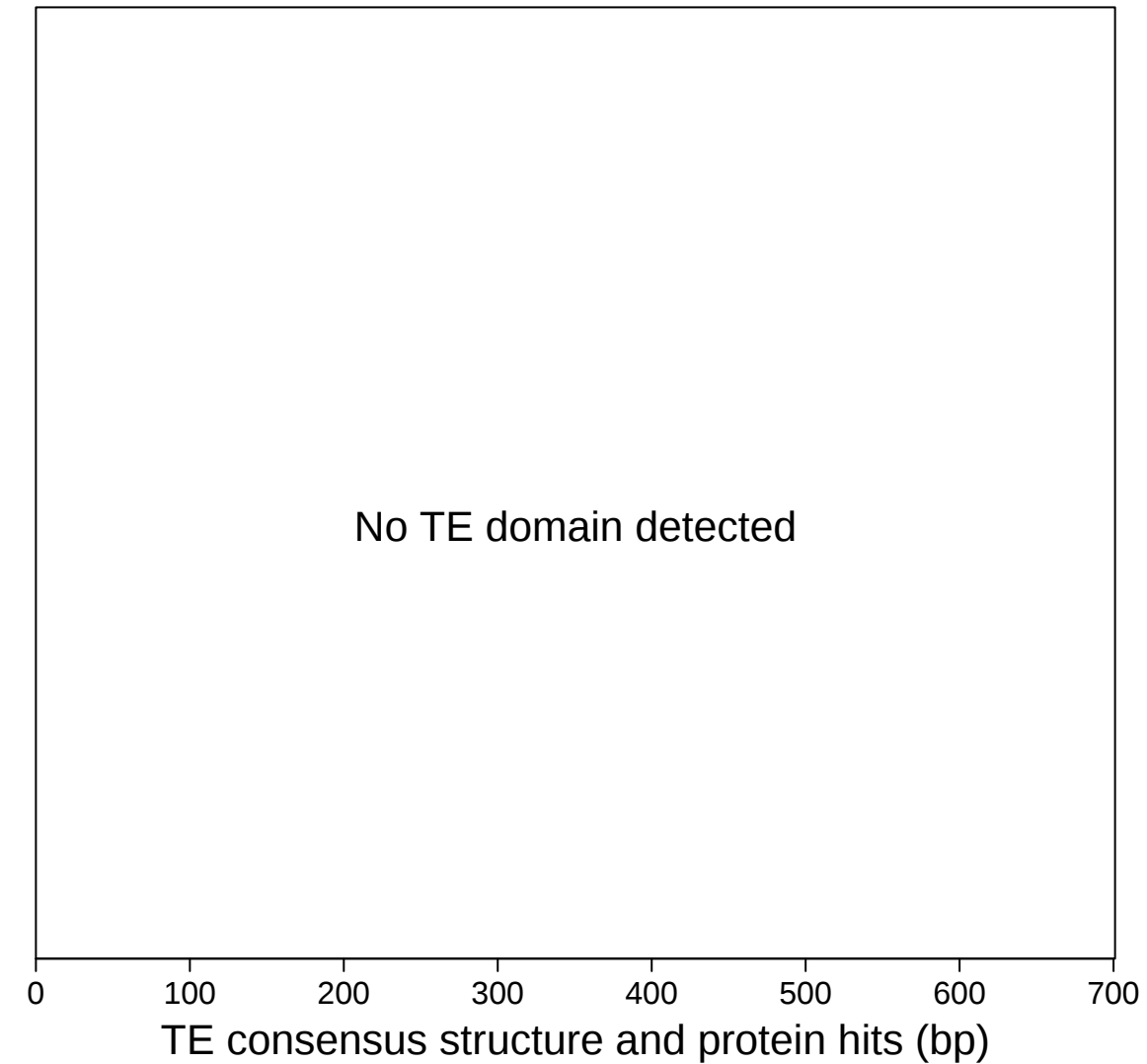
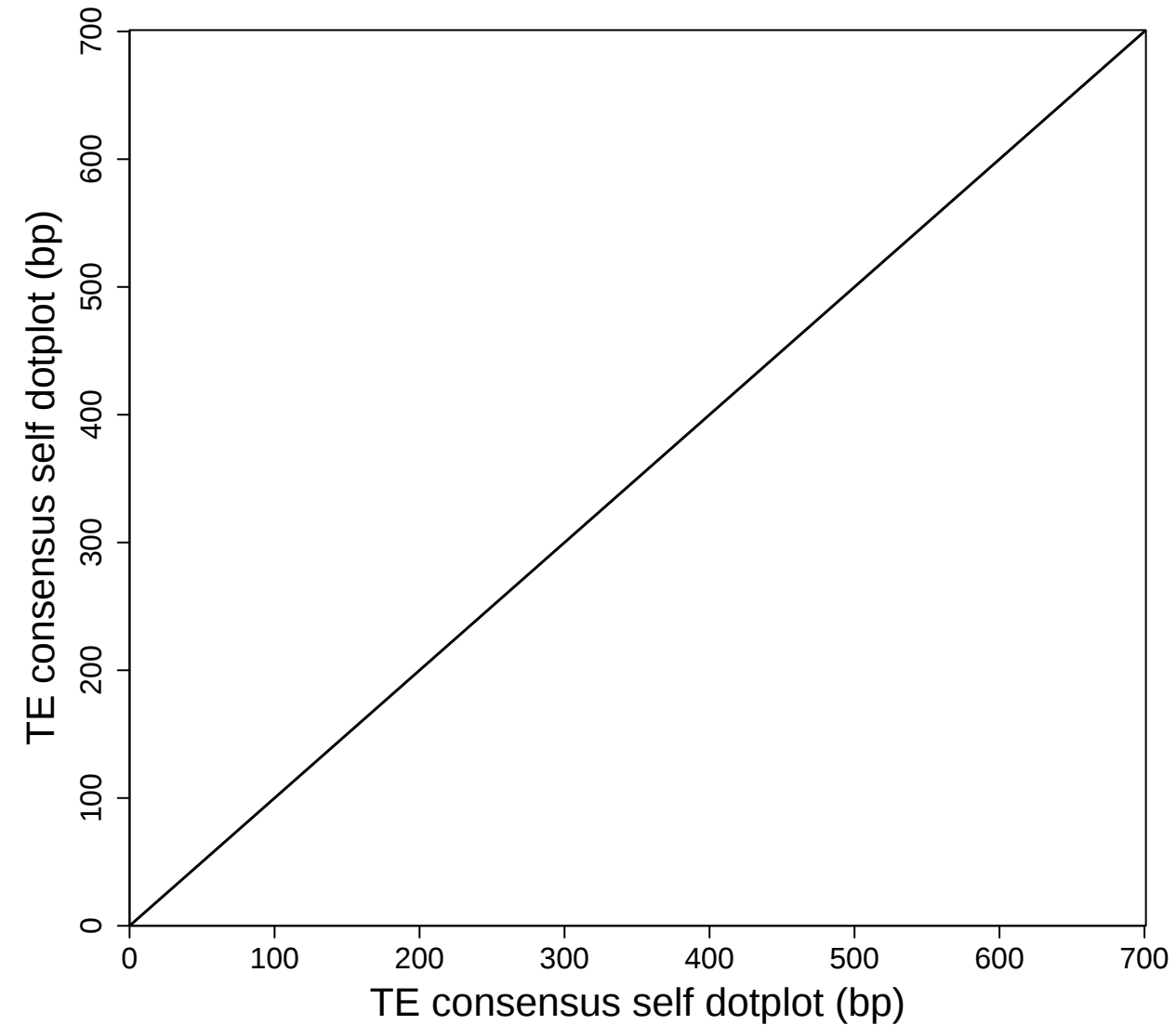
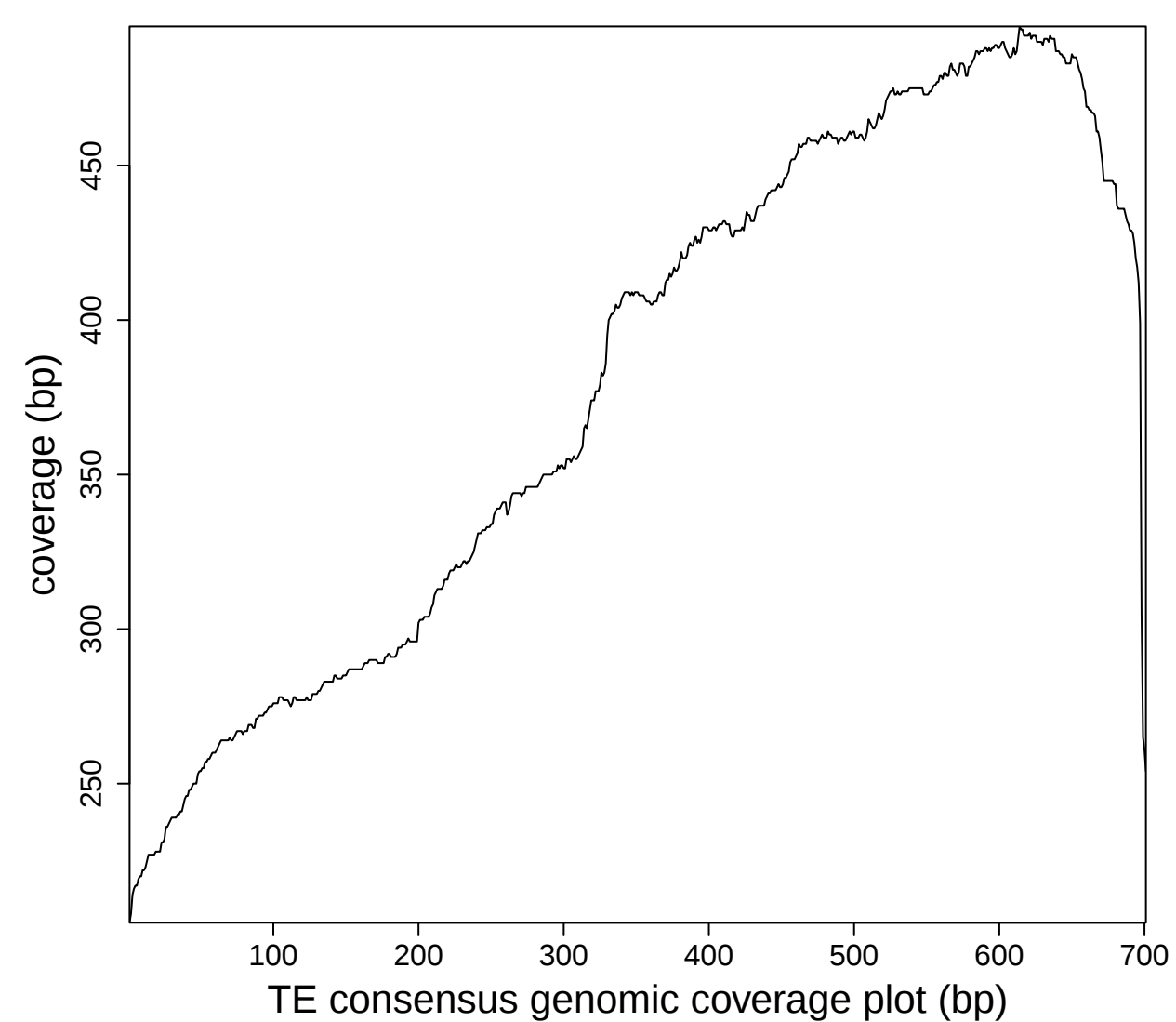
After TEtrimmer Extended plot Blue lines are boundaries



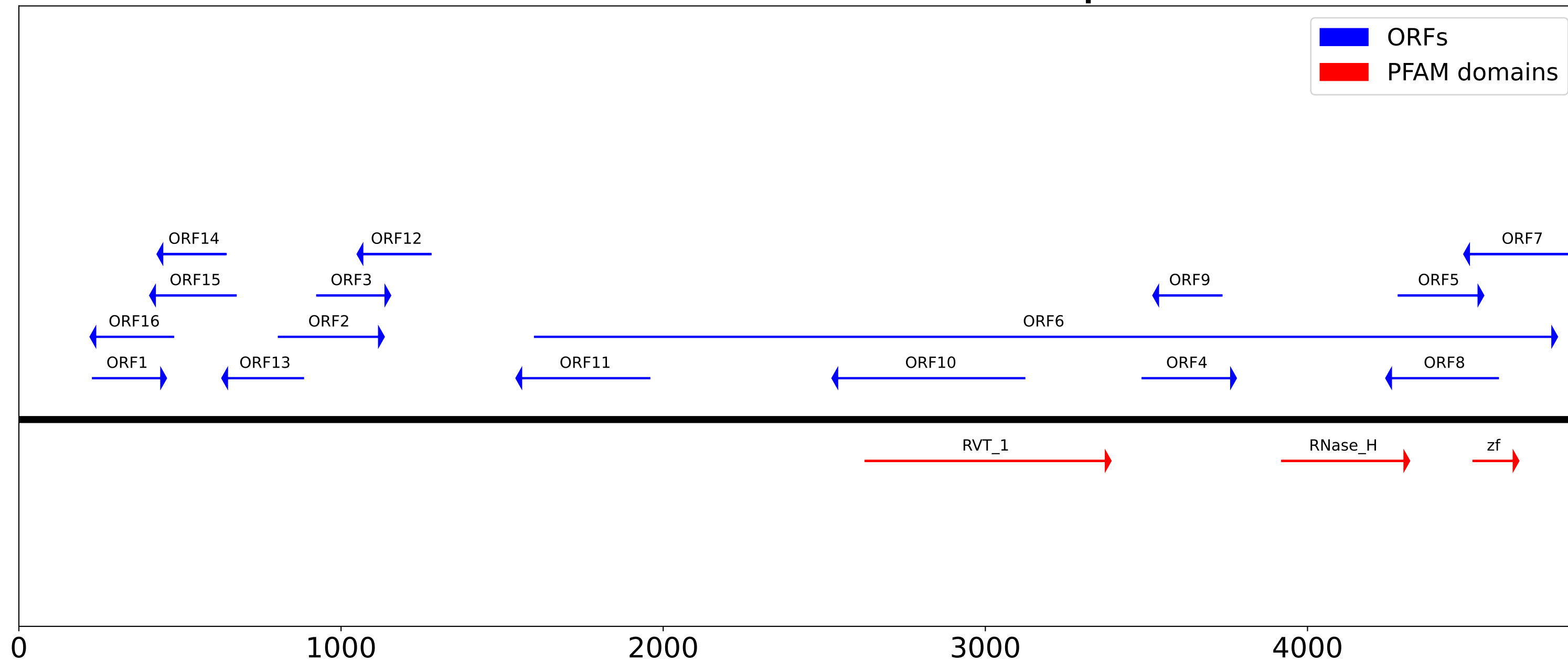
TE: rnd_4_family_4550
size: 701bp; fragments: 686; full length: 189 (≥ 630.9 bp)



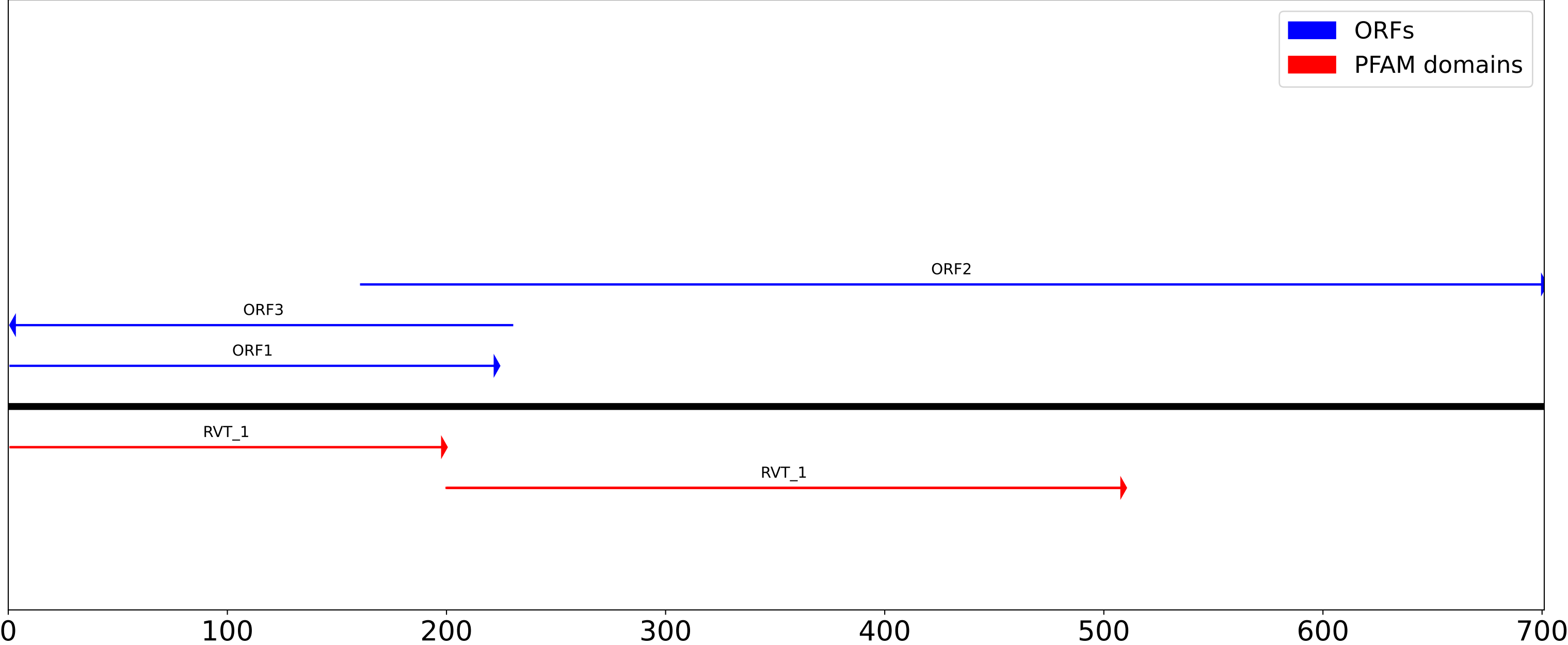
Before TEtrimmer 701 bp



After TEtrimmer ORF and PFAM domain plot



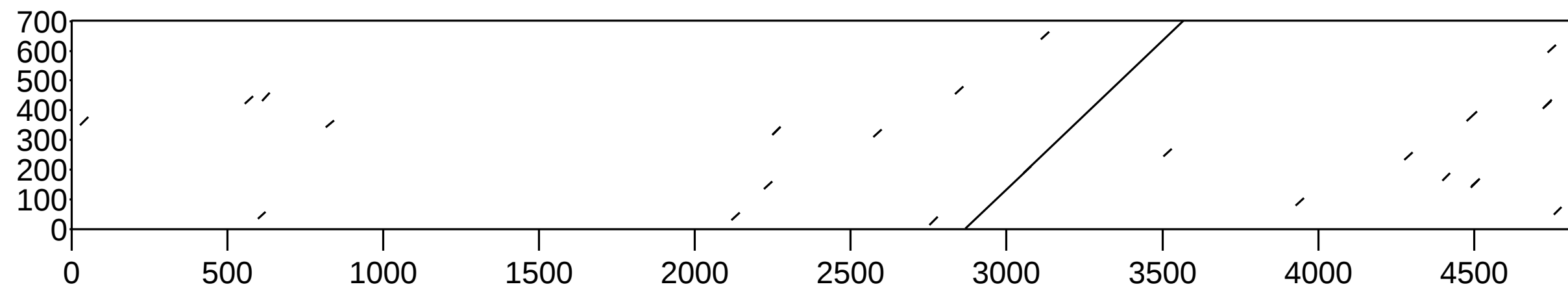
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

isensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)