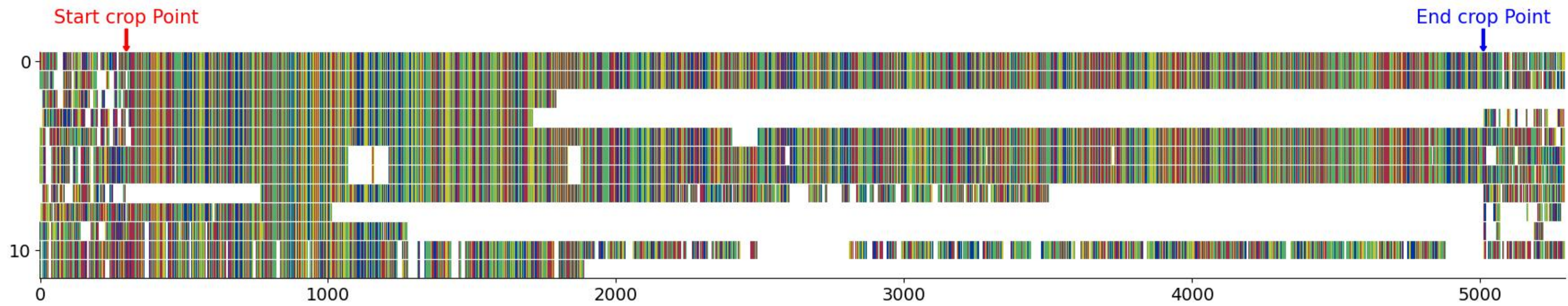


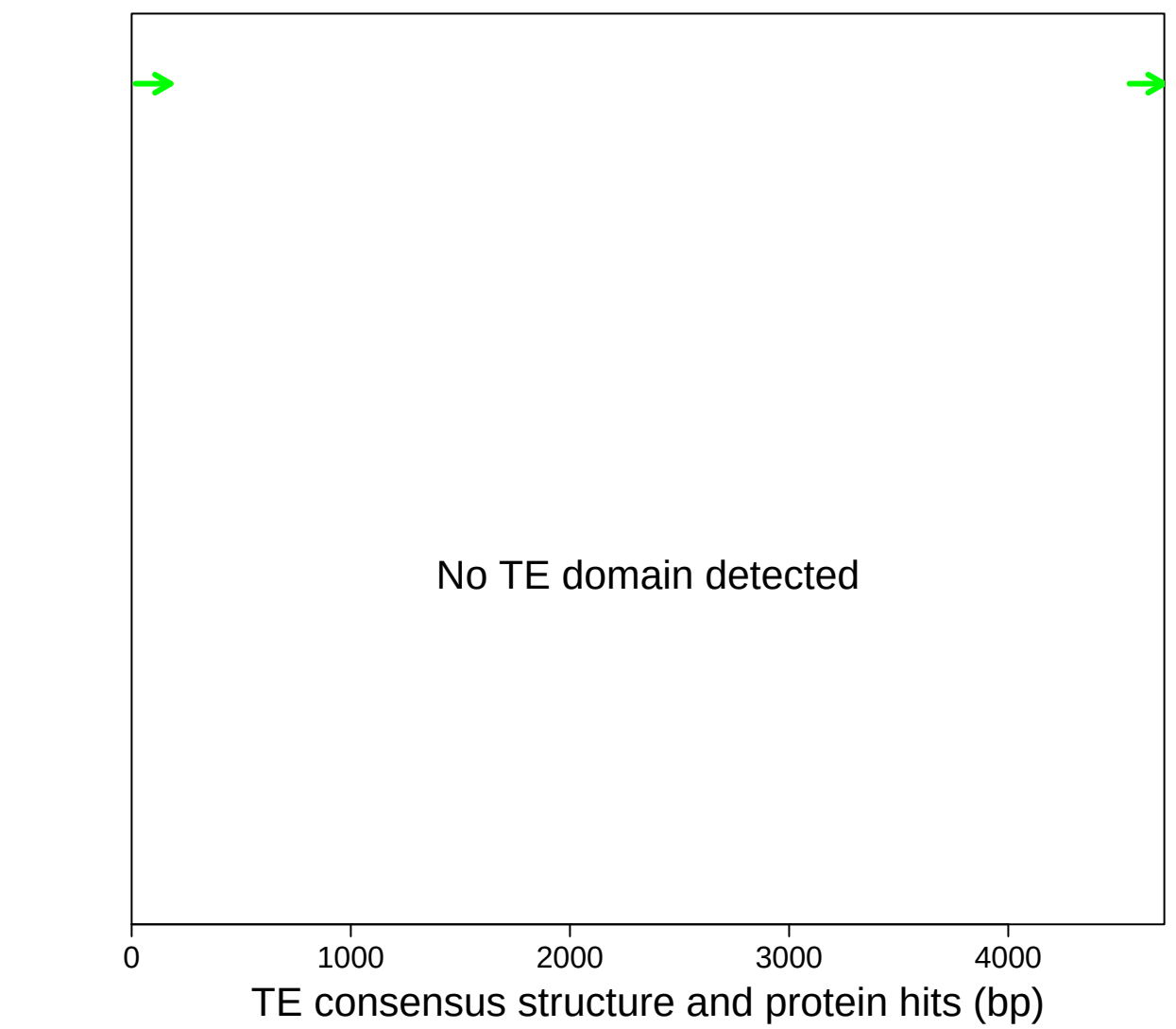
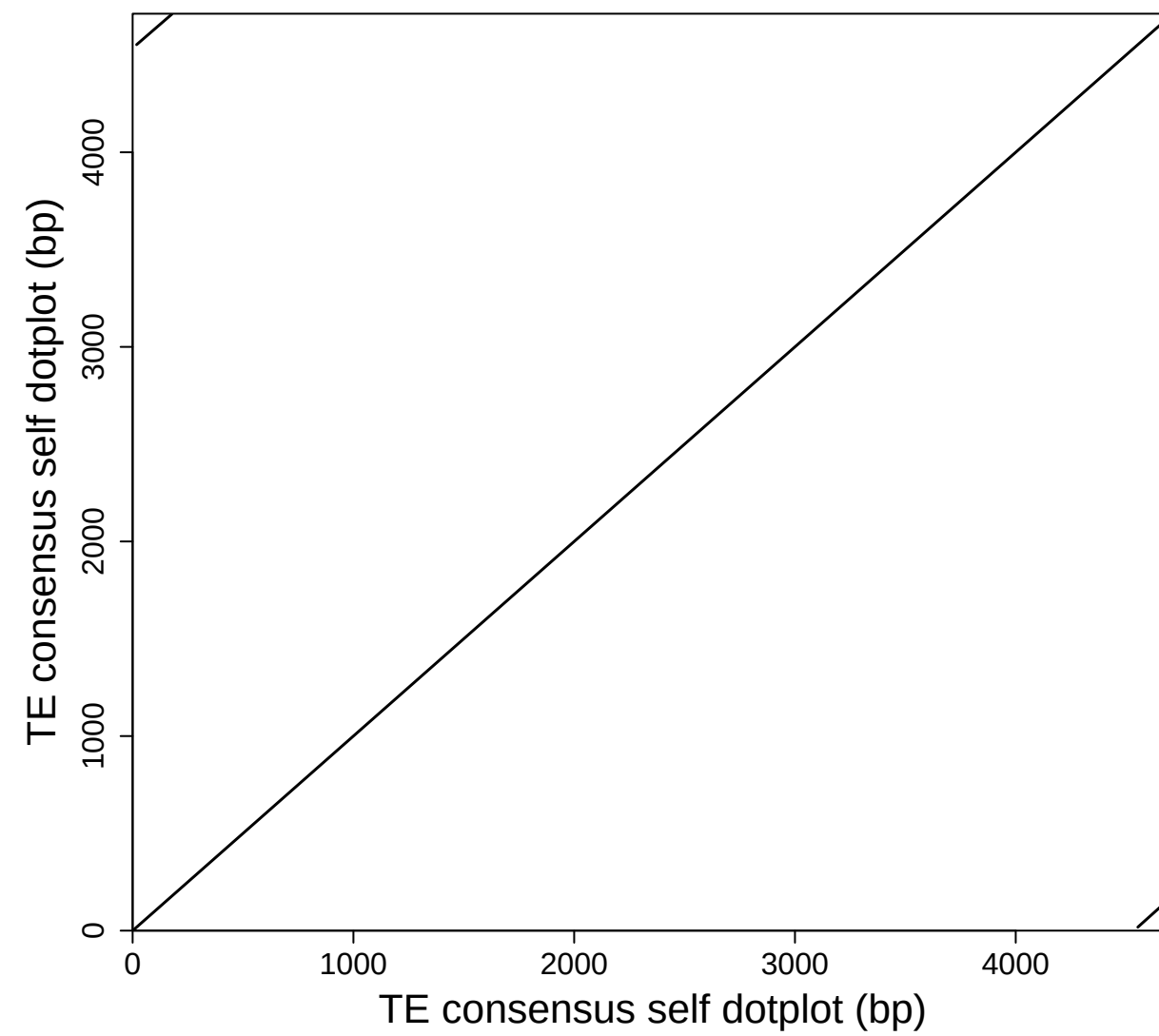
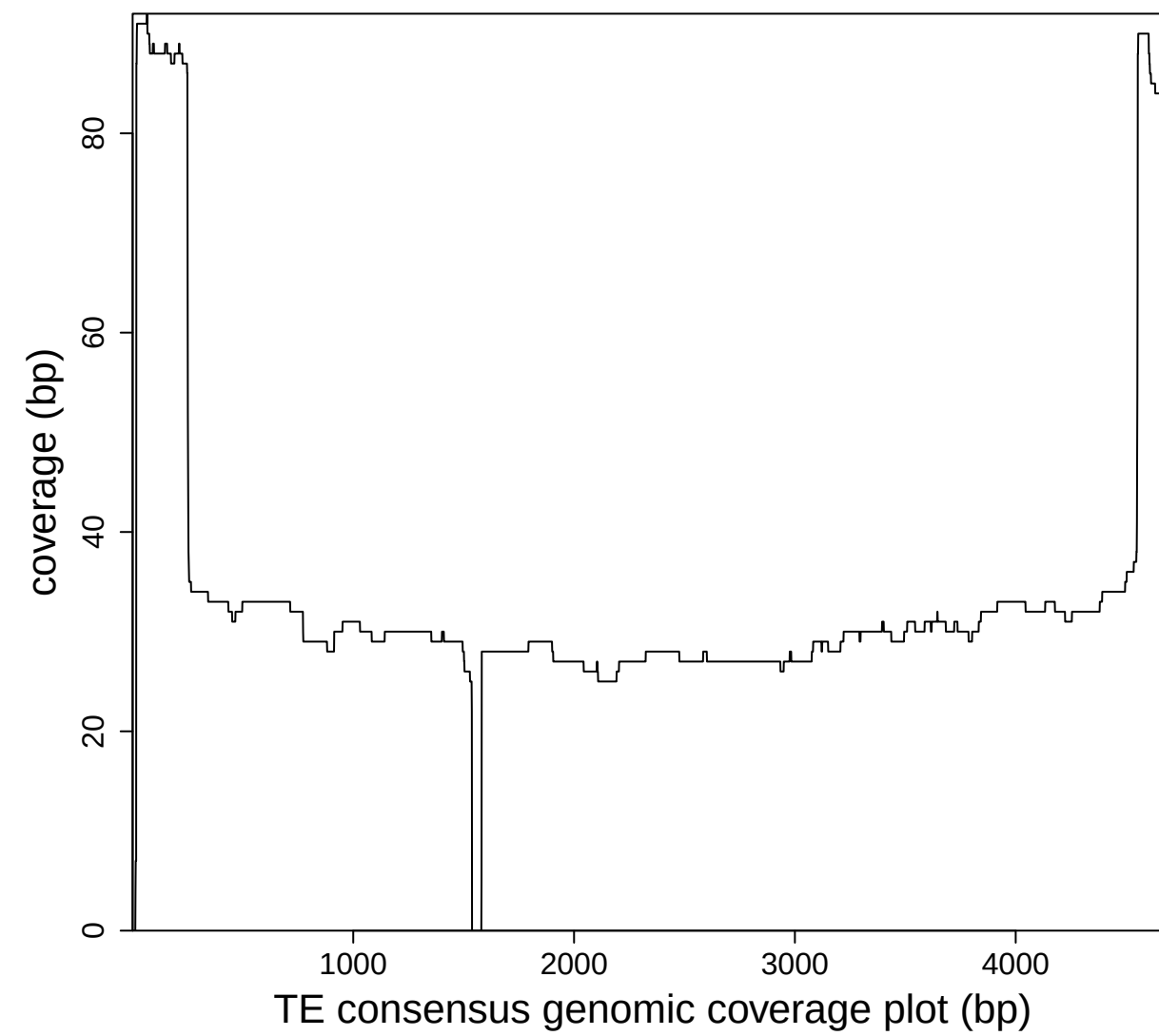
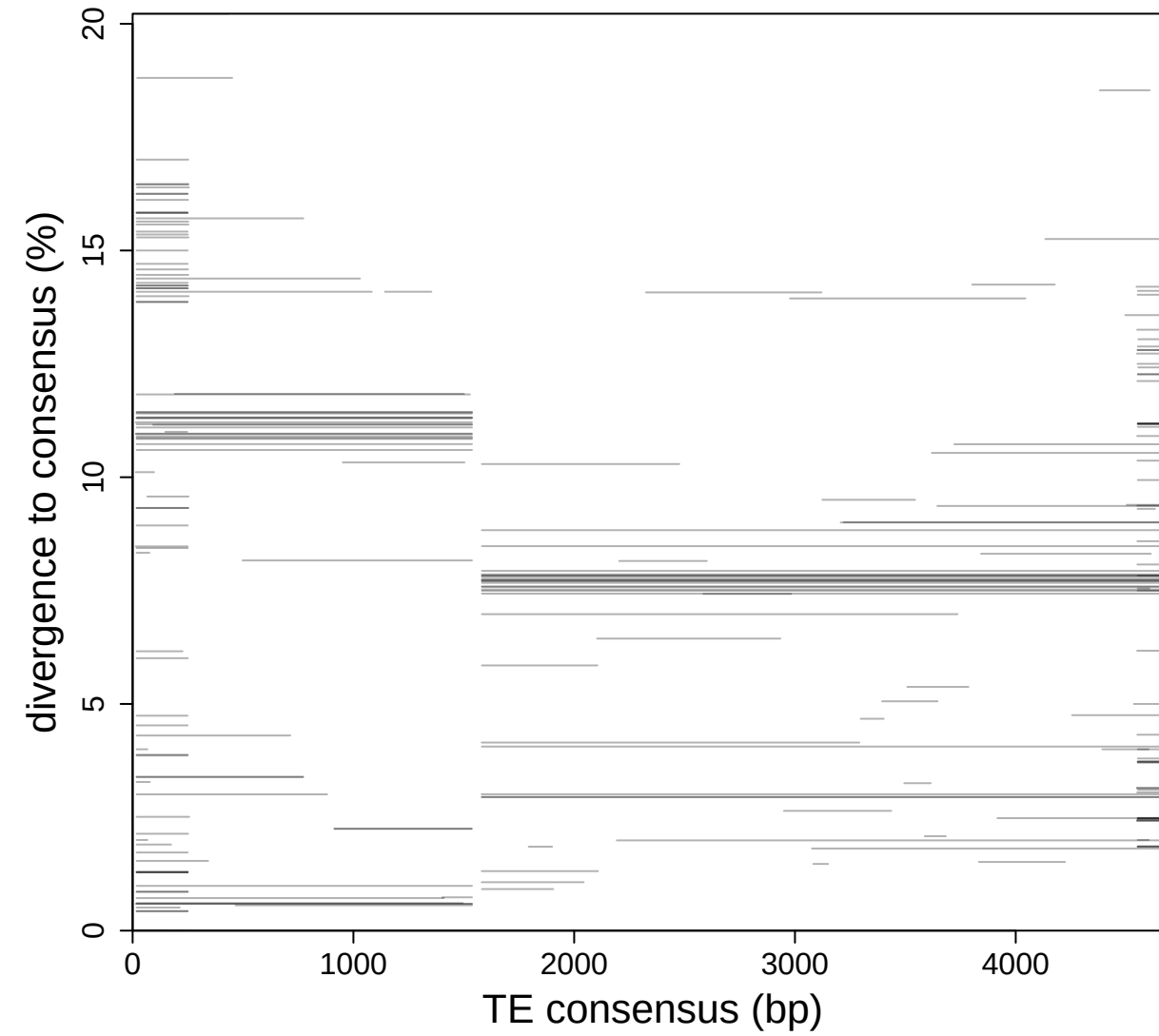
MSA length = 4712





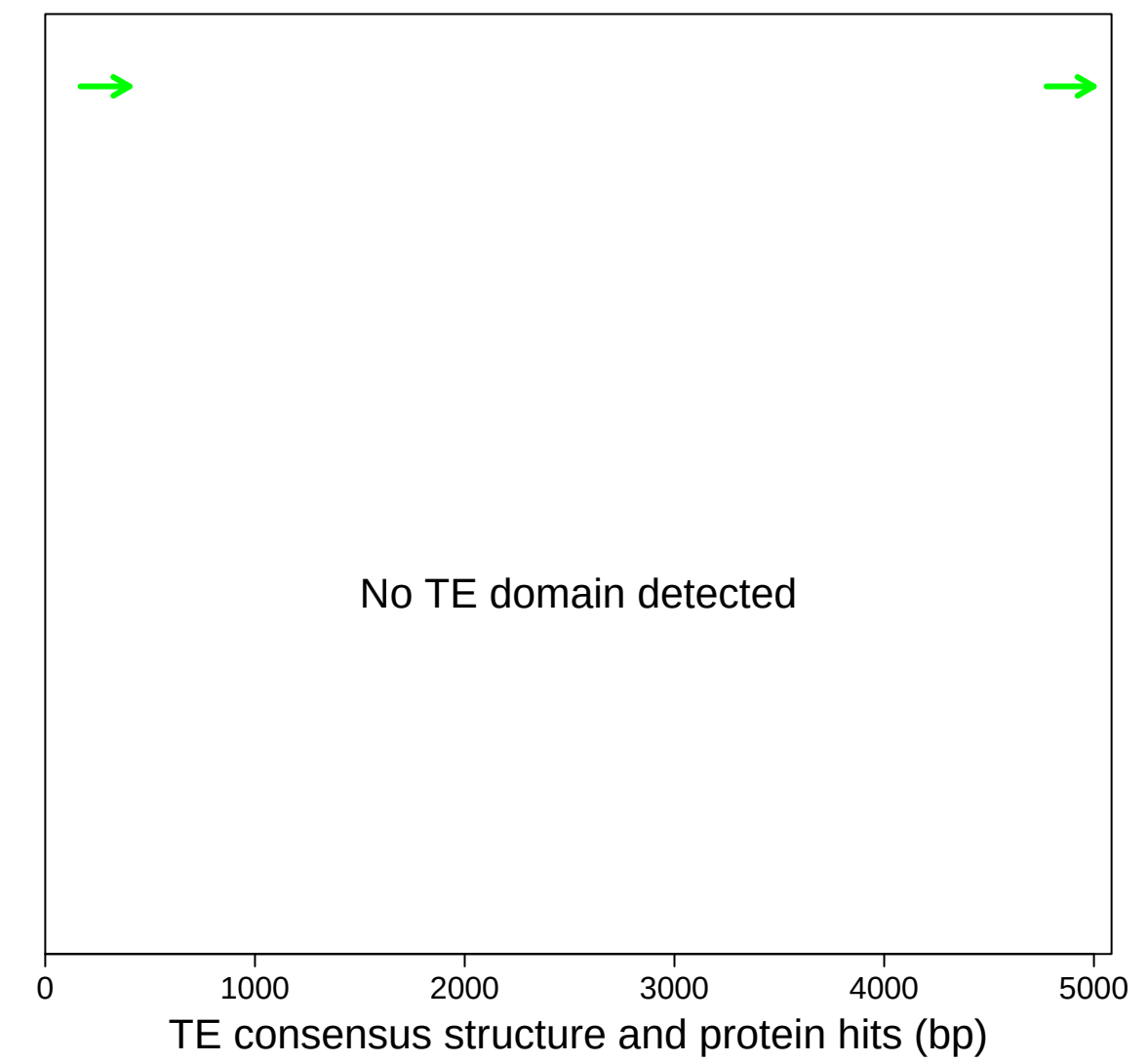
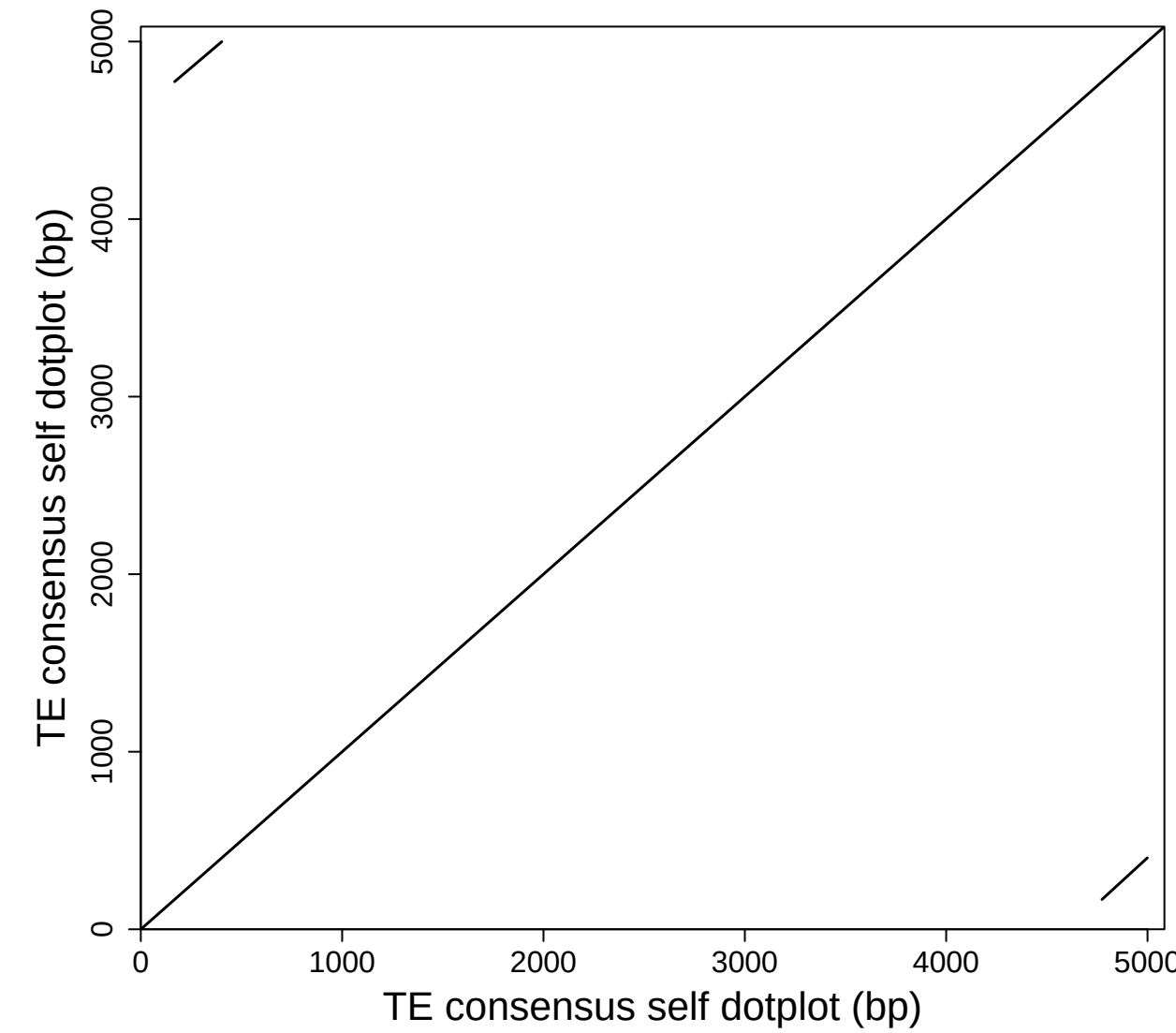
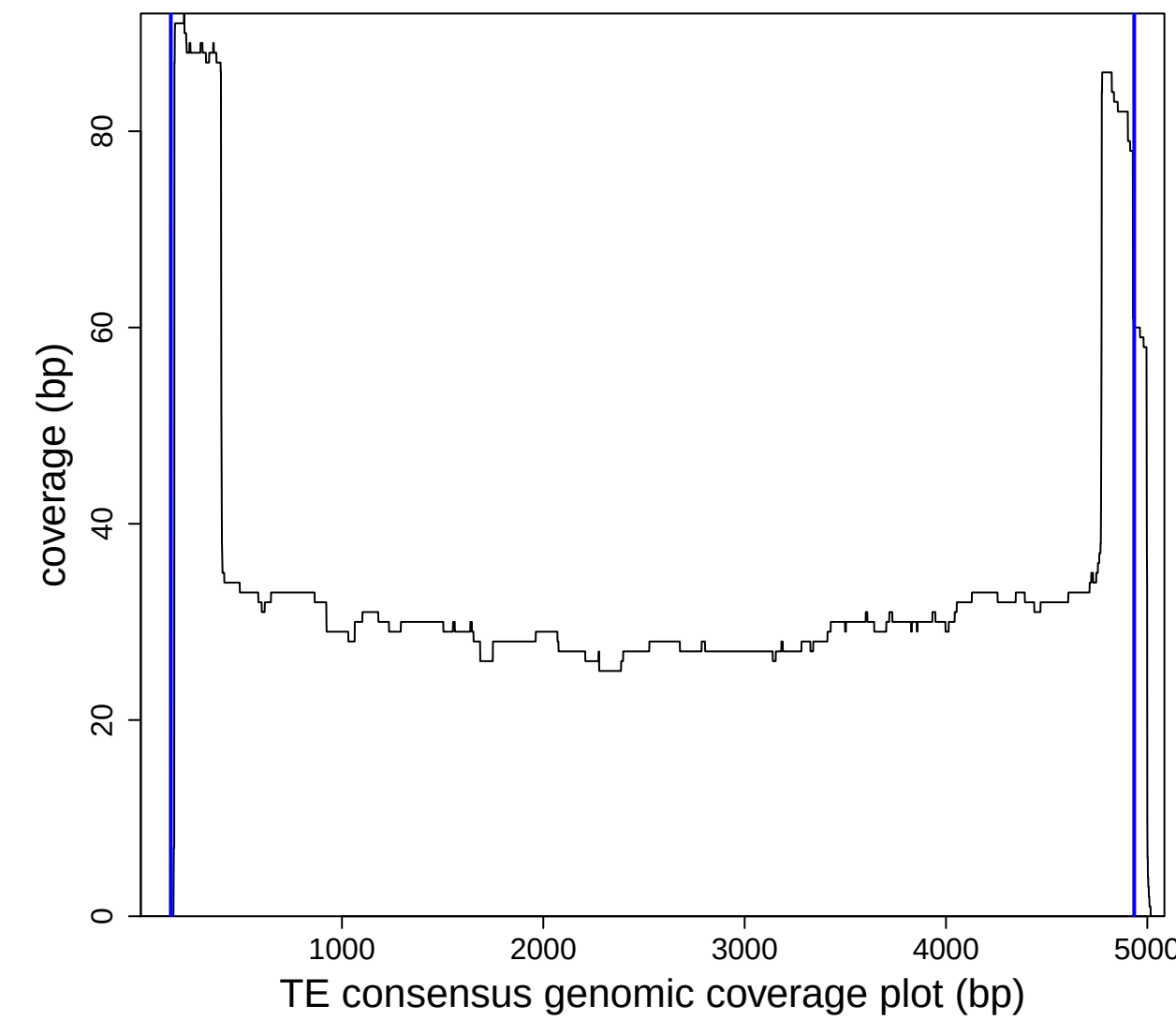
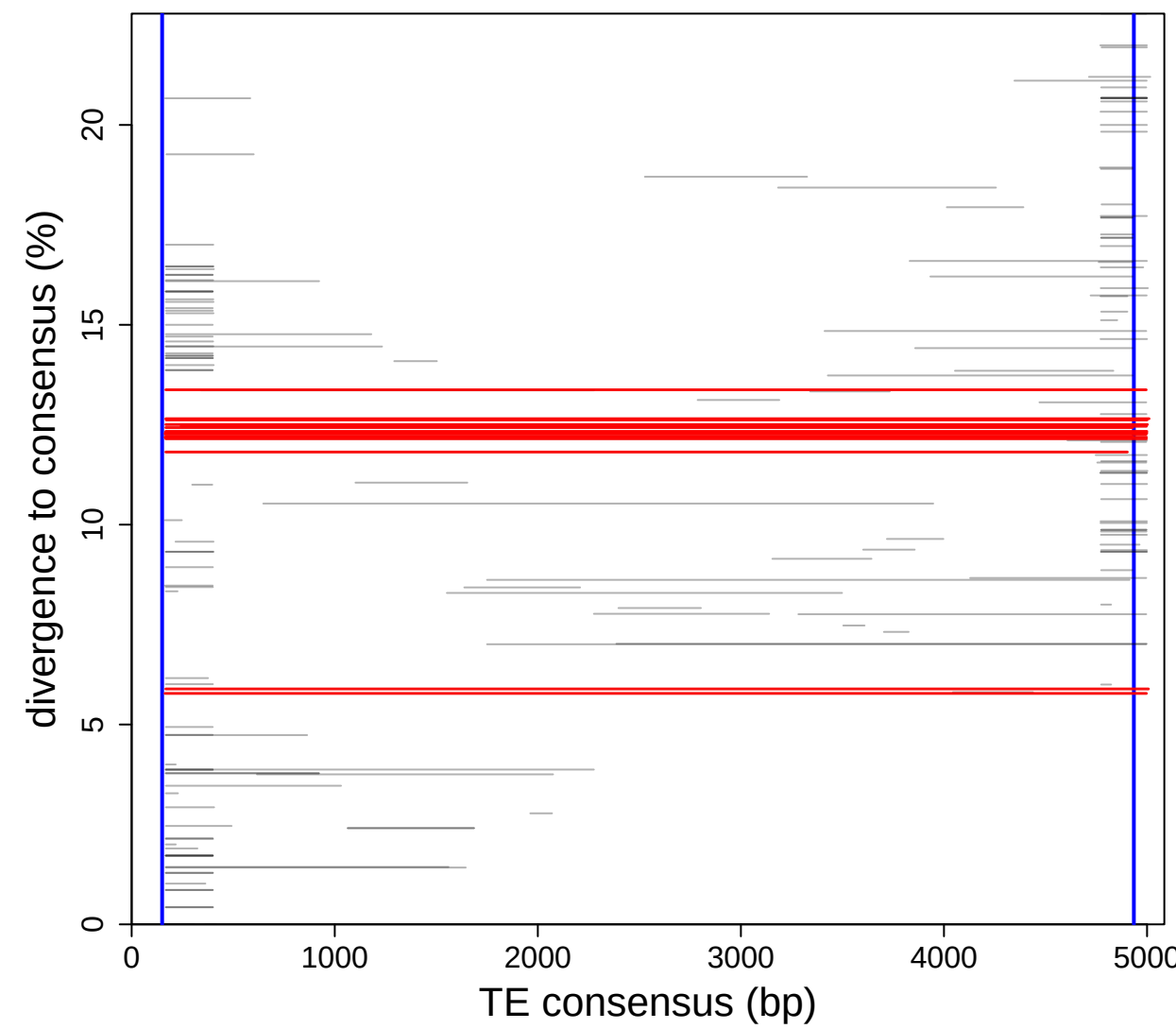
ed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 4712bp; fragments: 216; full length: 0 (>=4240.8bp)

After TETrimmer 4712 bp



ly_768.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcIn.fa_aln.fa
size: 5085bp; fragments: 186; full length: 18 (≥ 4576.5 bp)

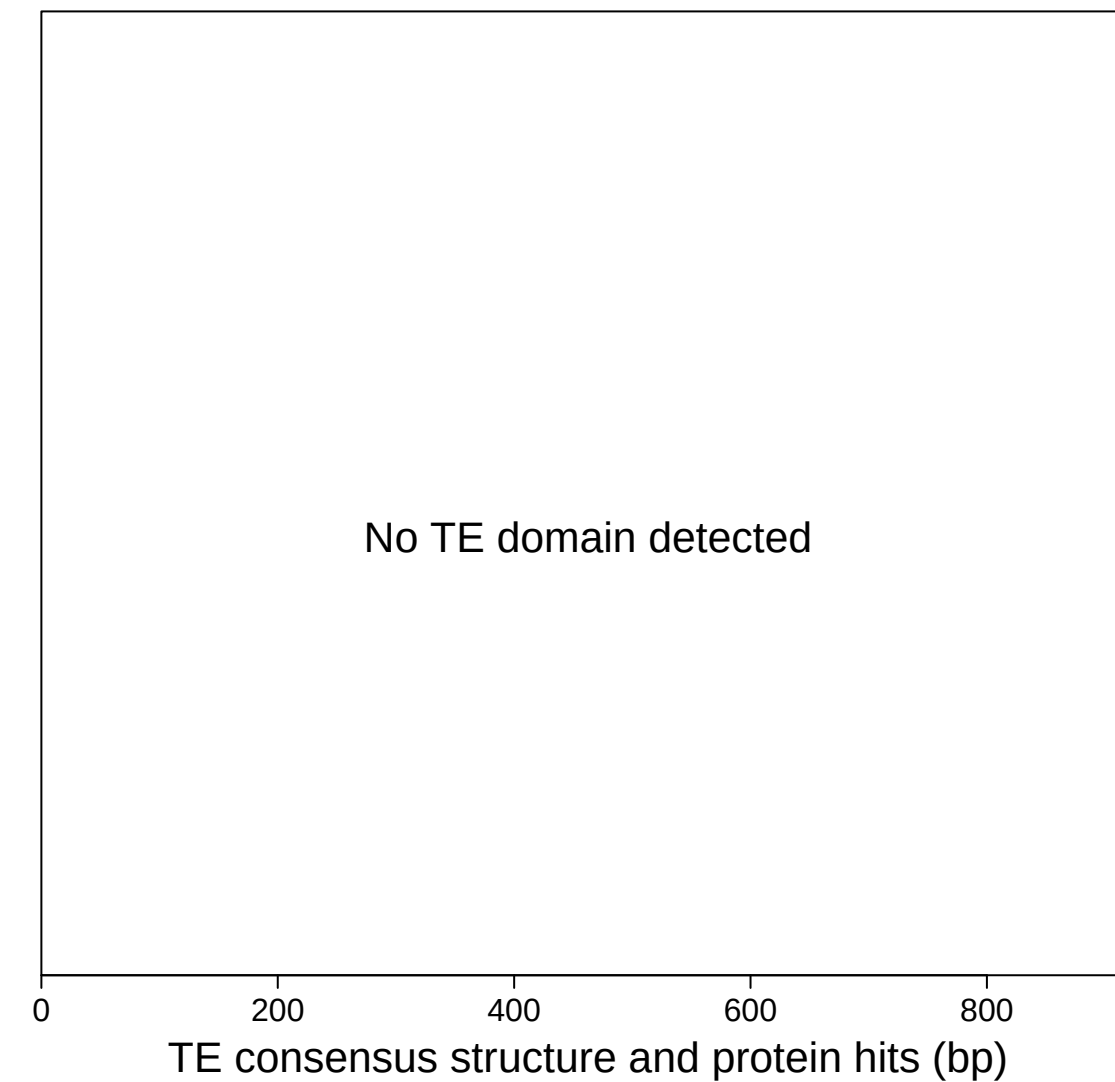
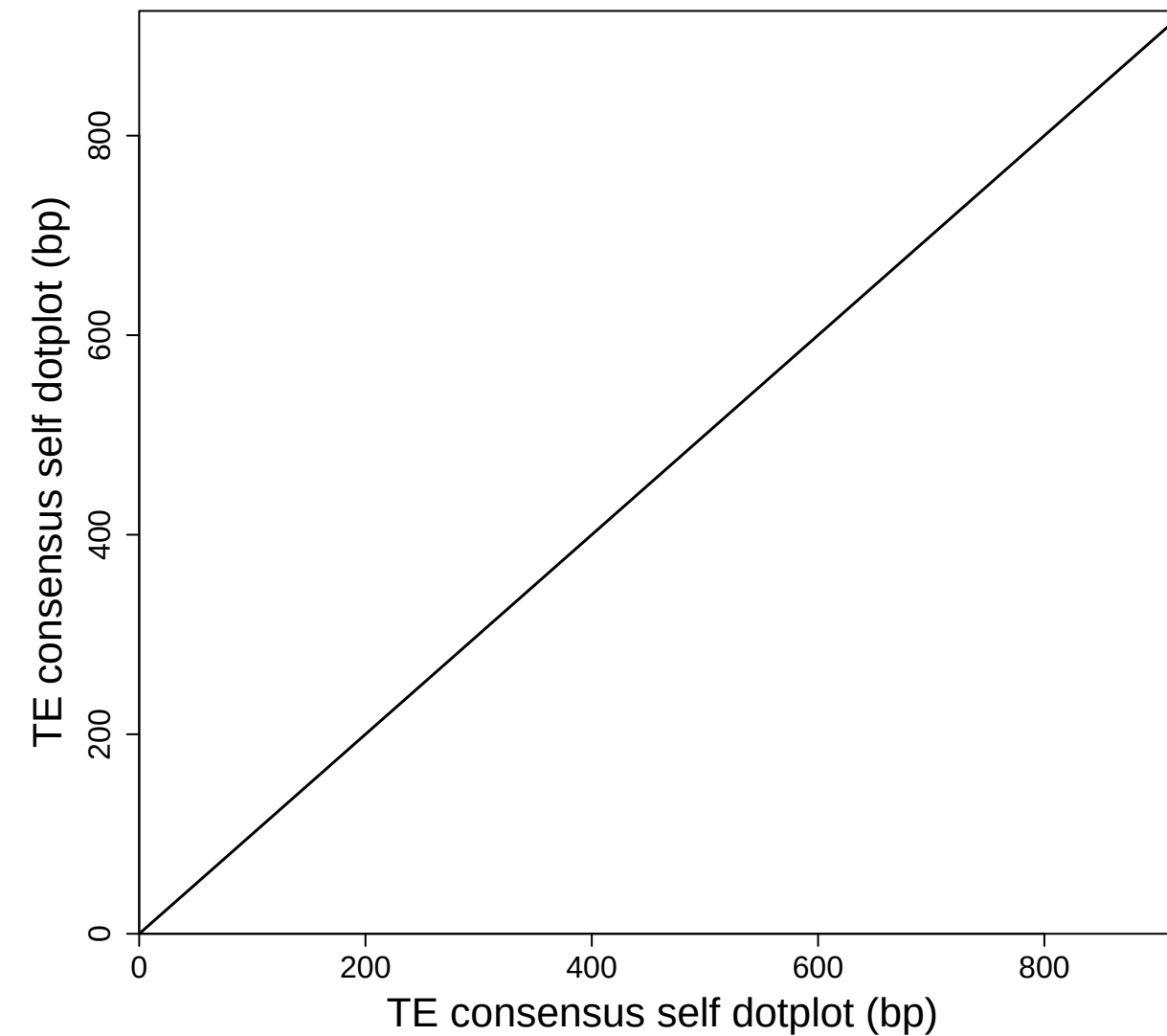
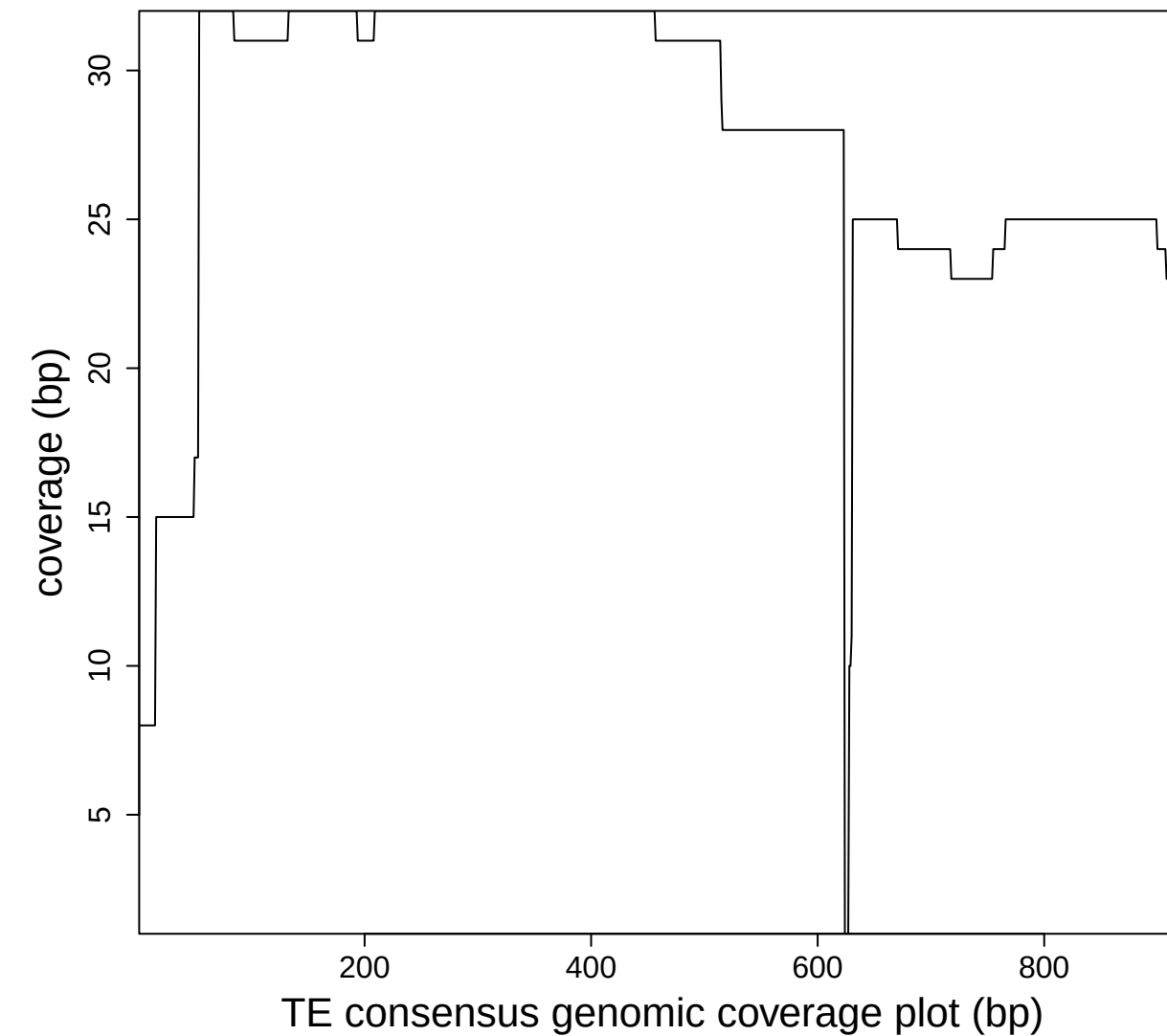
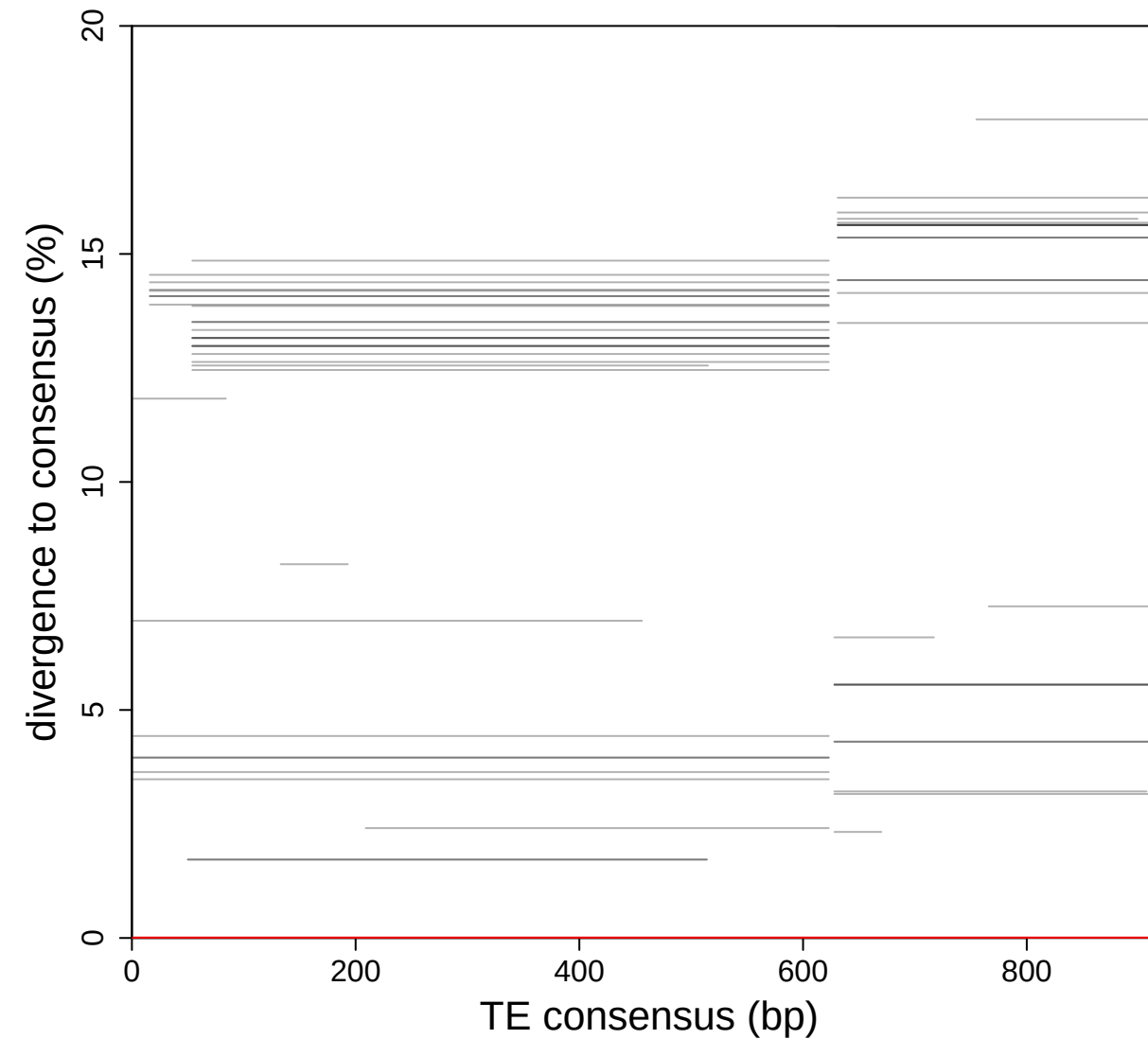
After TEtrimmer Extended plot Blue lines are boundaries



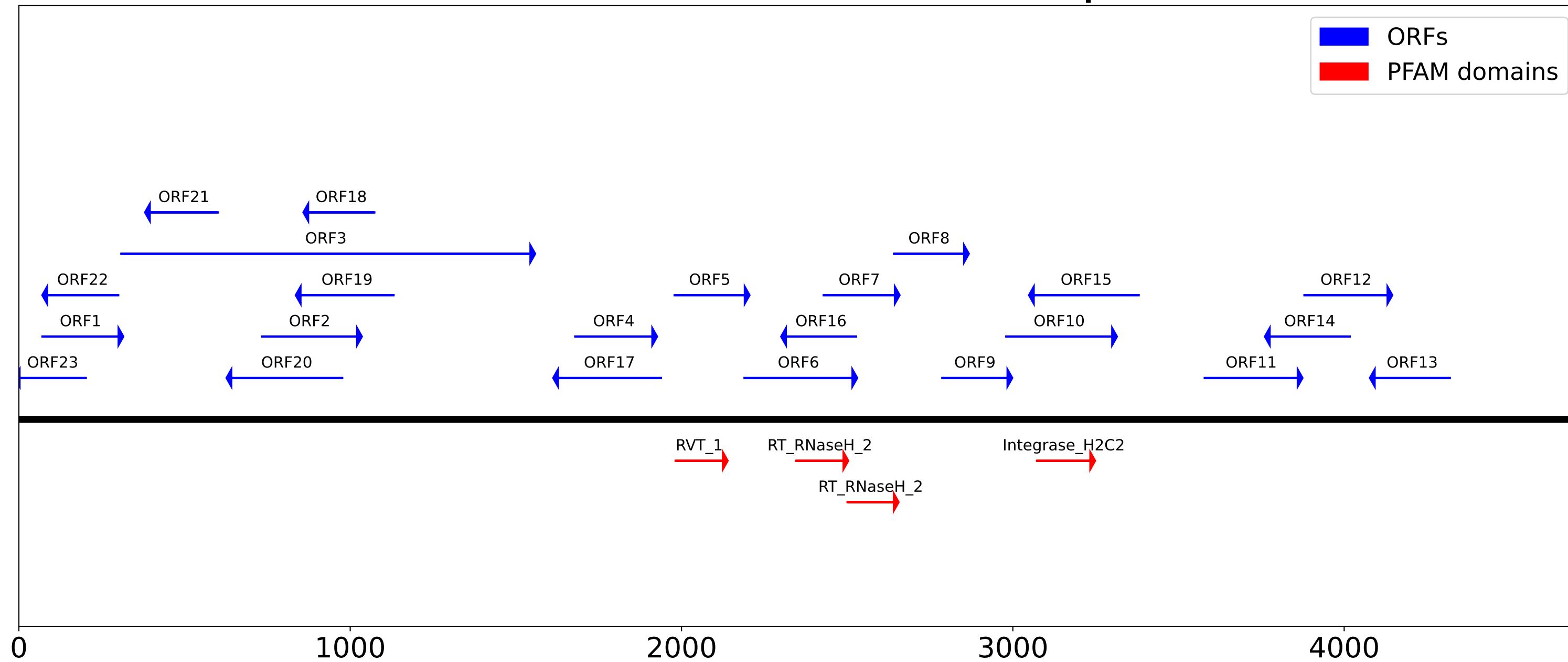
Before TEtrimmer 925 bp

TE: ltr_1_family_768

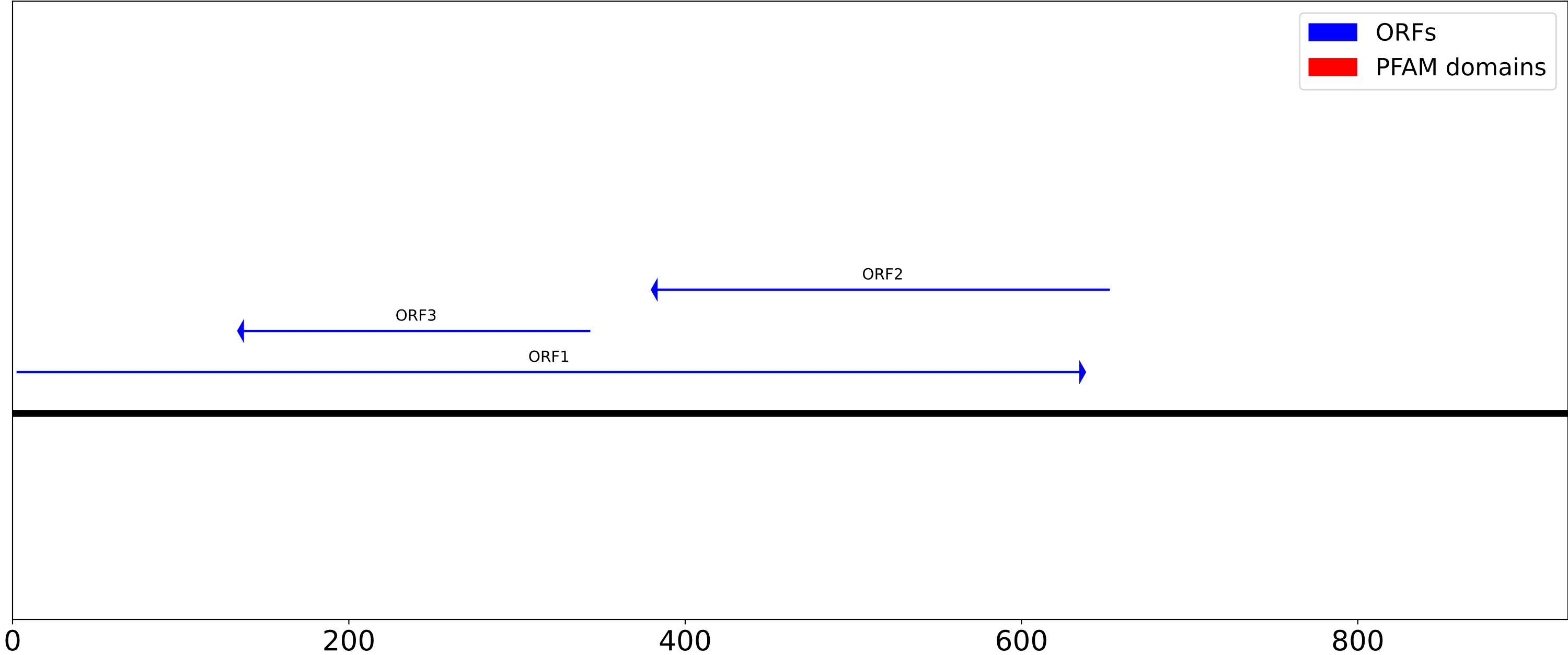
size: 925bp; fragments: 60; full length: 1 (≥ 832.5 bp)



After TETrimmer ORF and PFAM domain plot



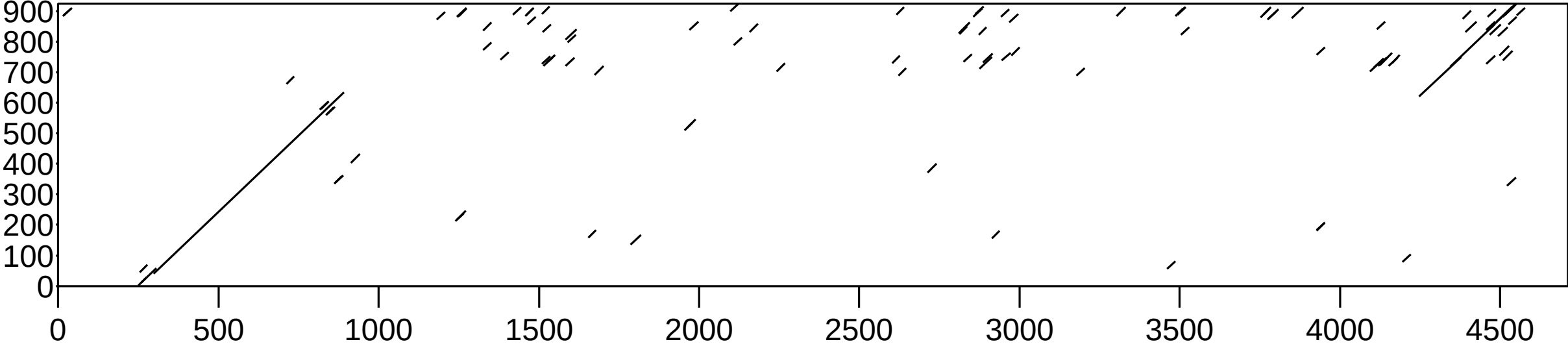
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)