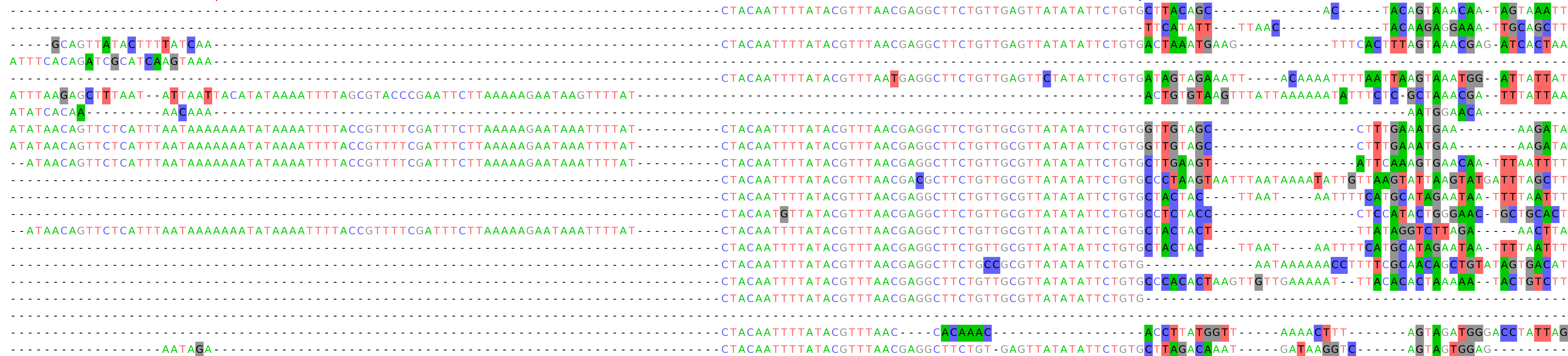


MSA length = 5096

Start crop Point

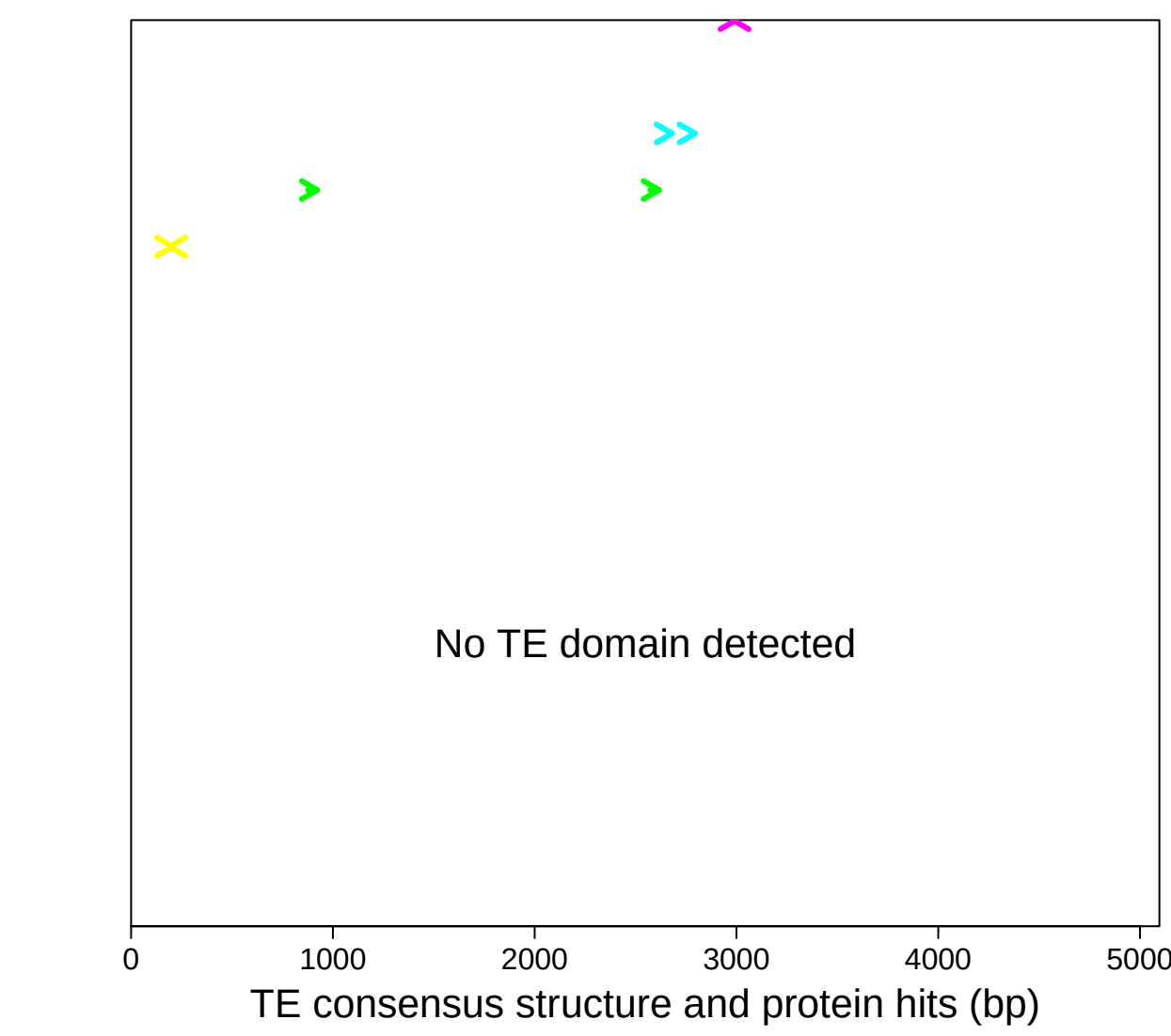
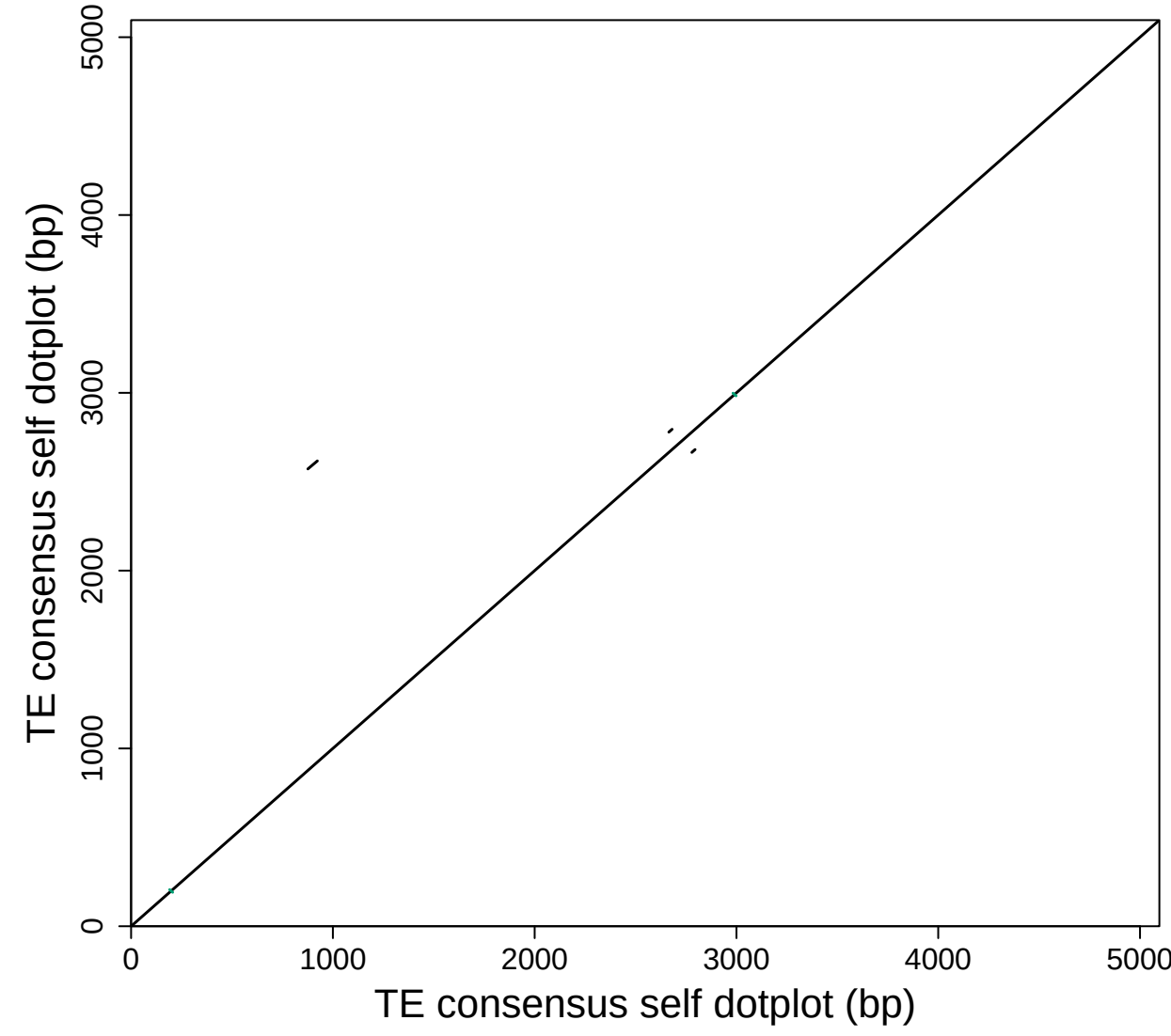
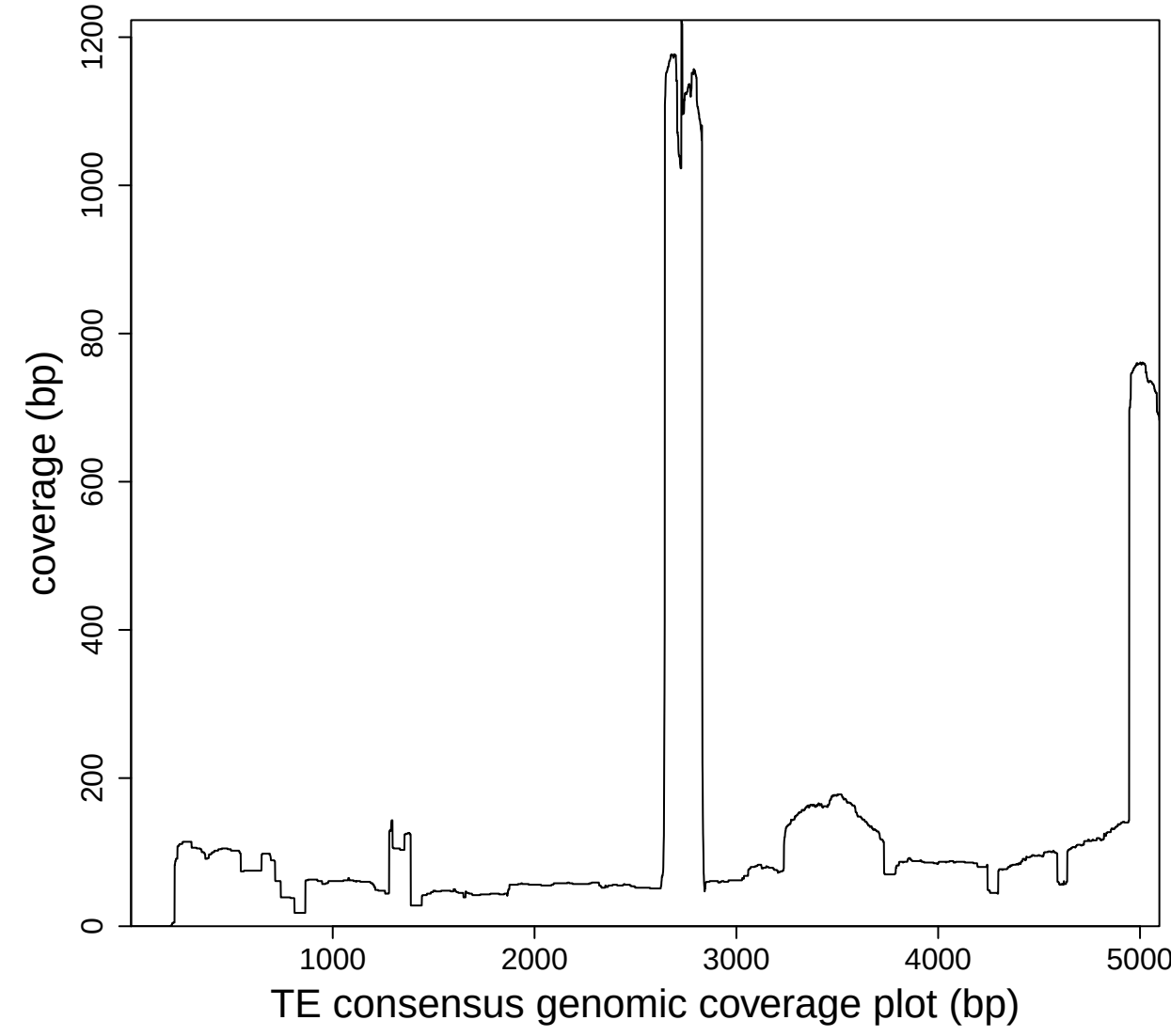
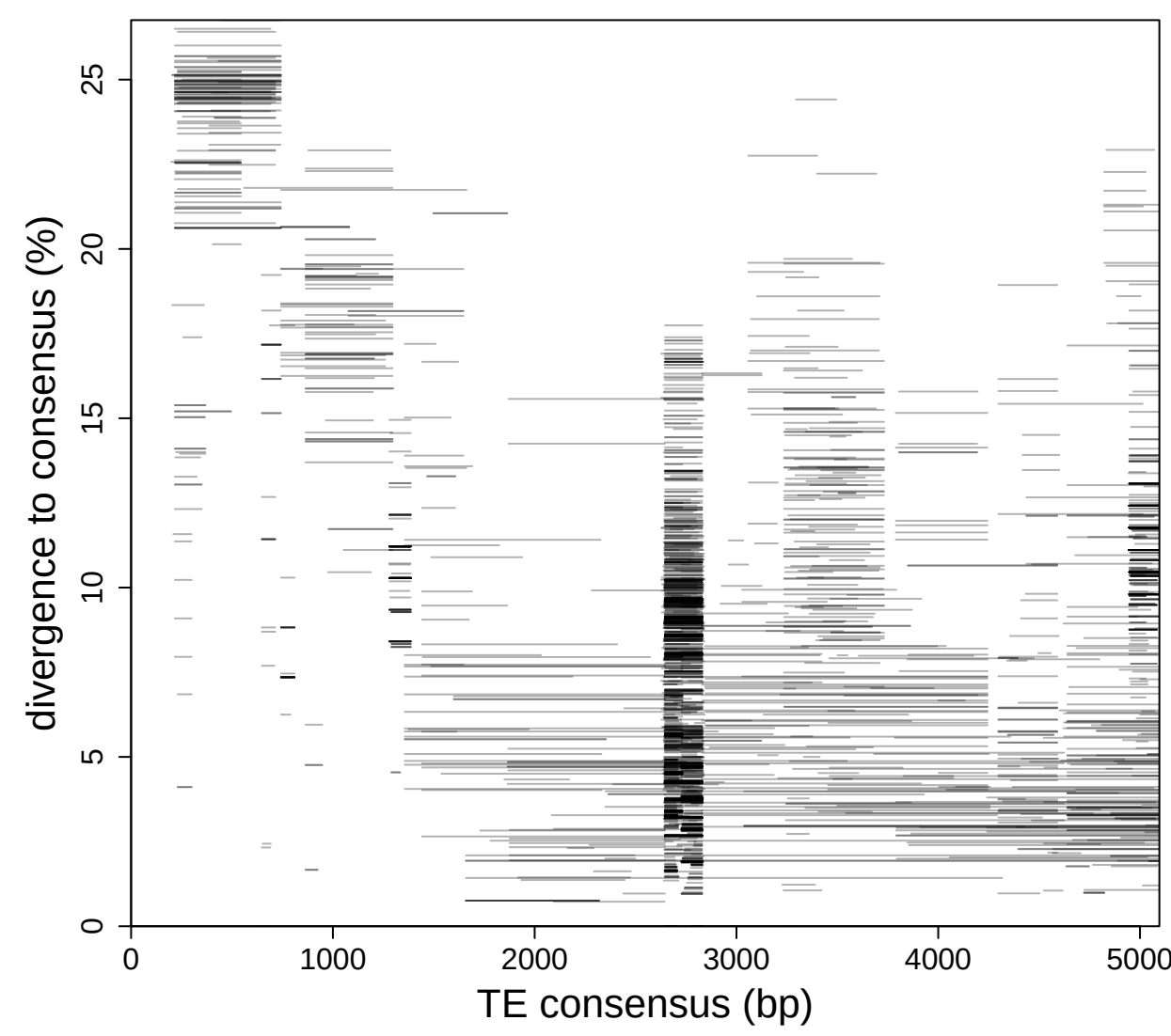
End crop Point





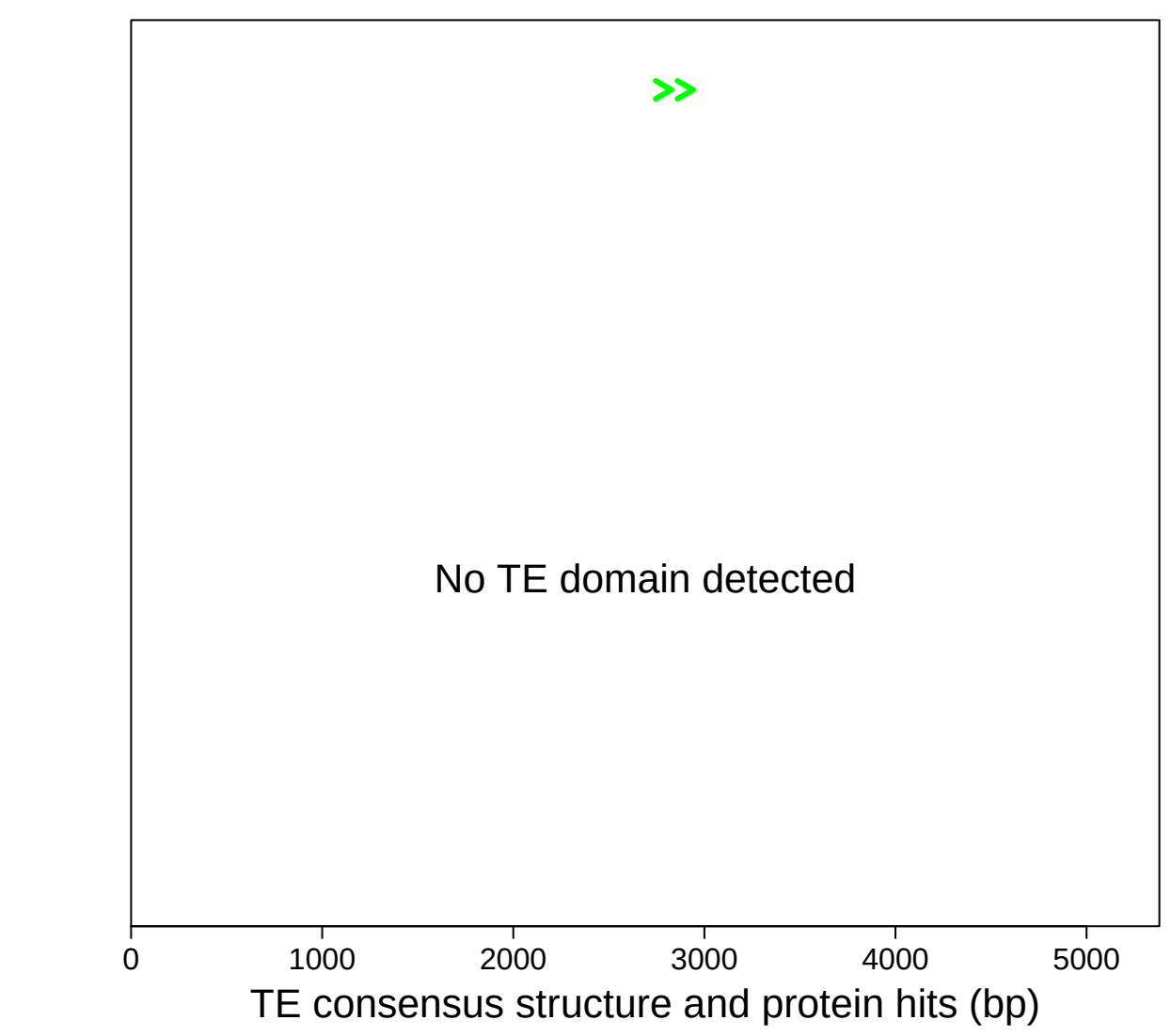
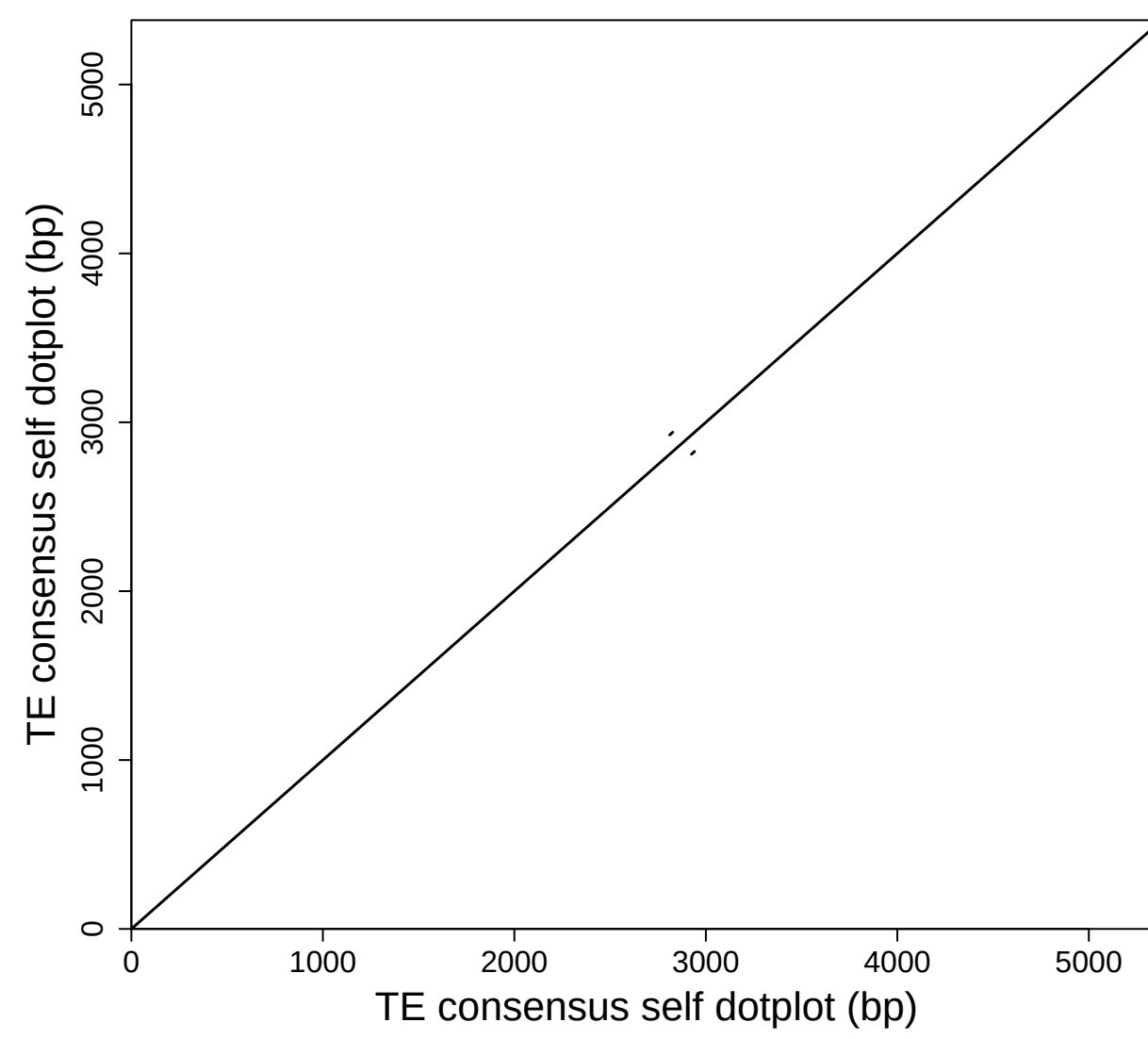
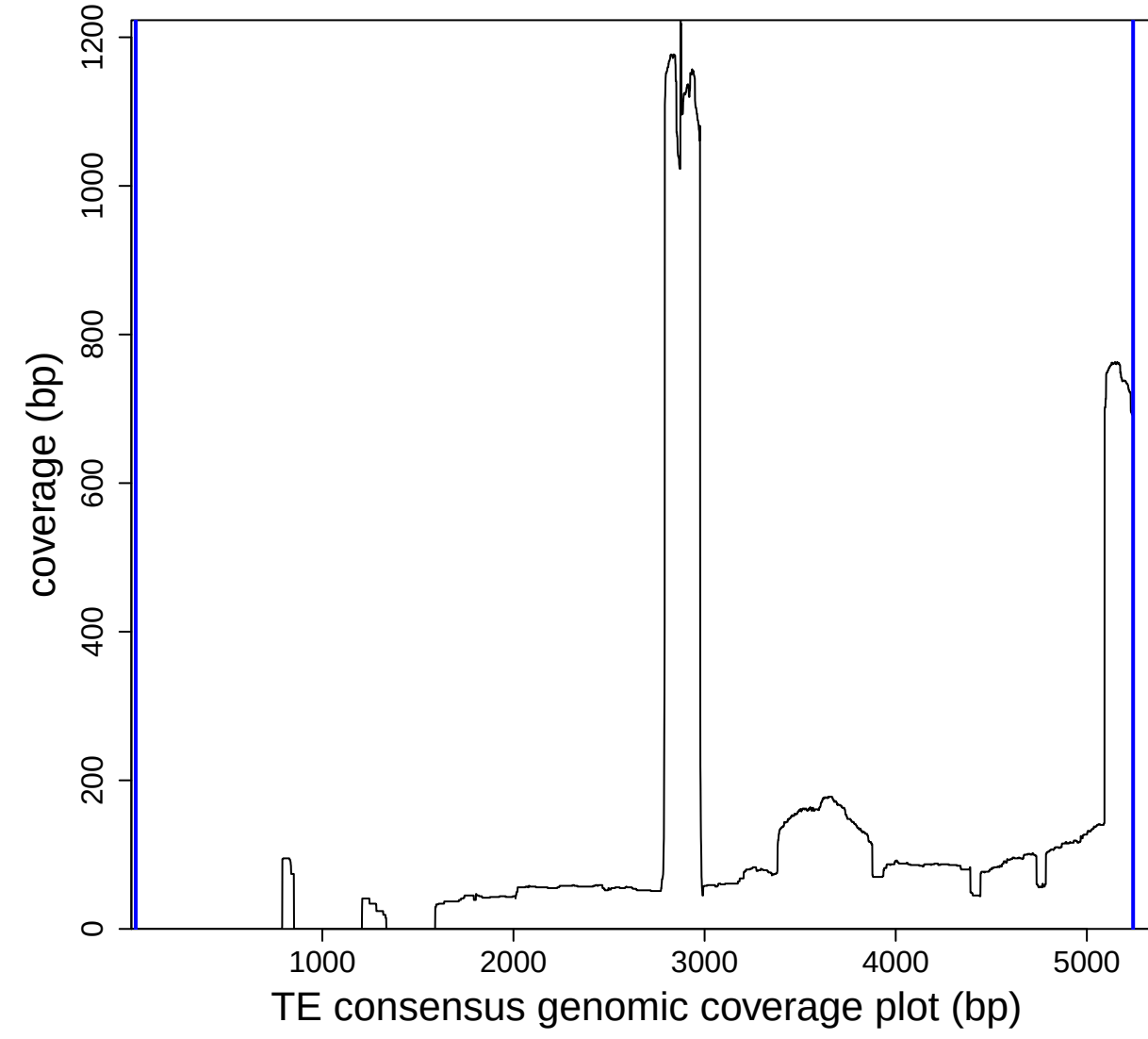
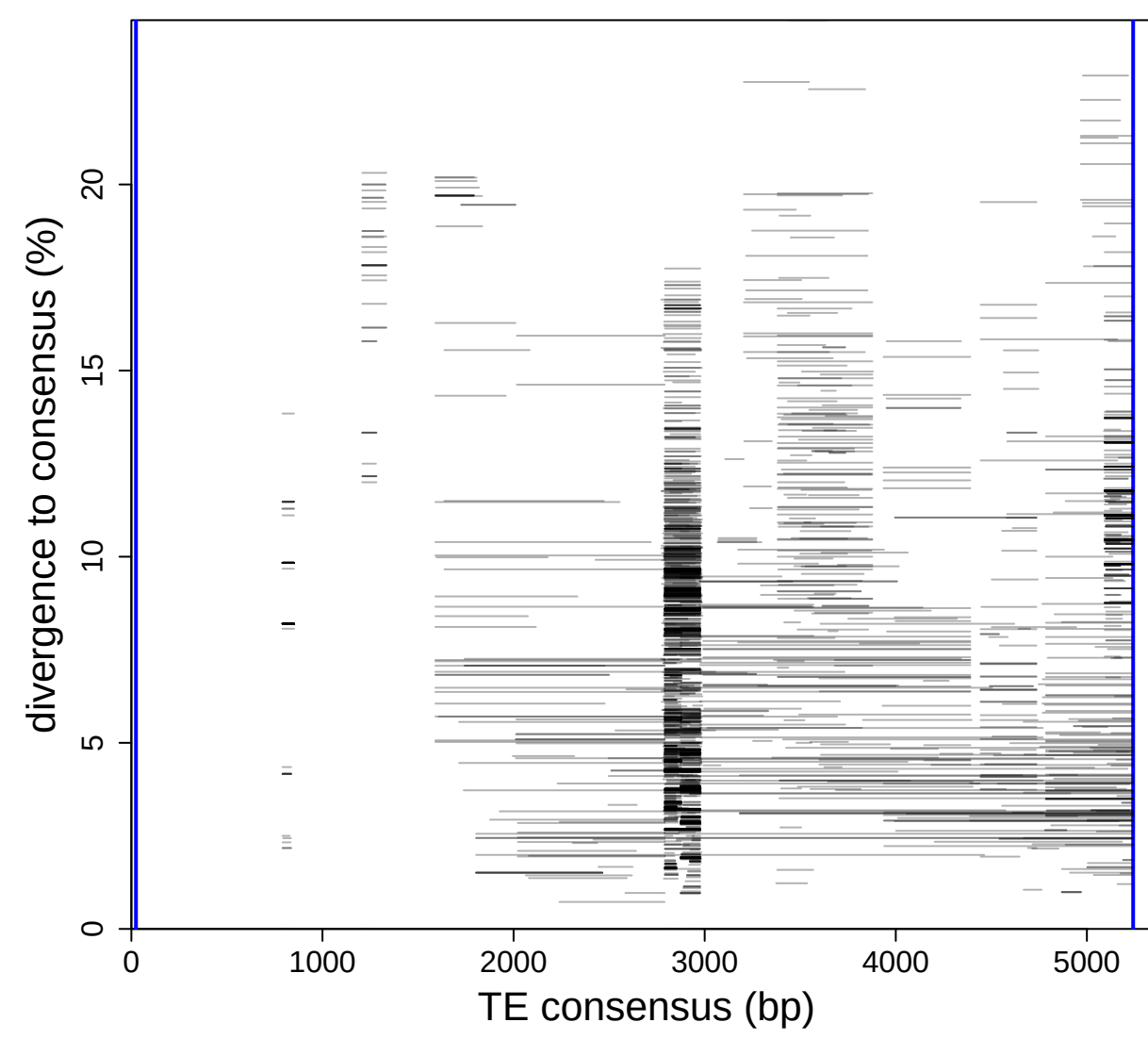
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 5096bp; fragments: 3026; full length: 0 (>=4586.4bp)

After TETrimmer 5096 bp



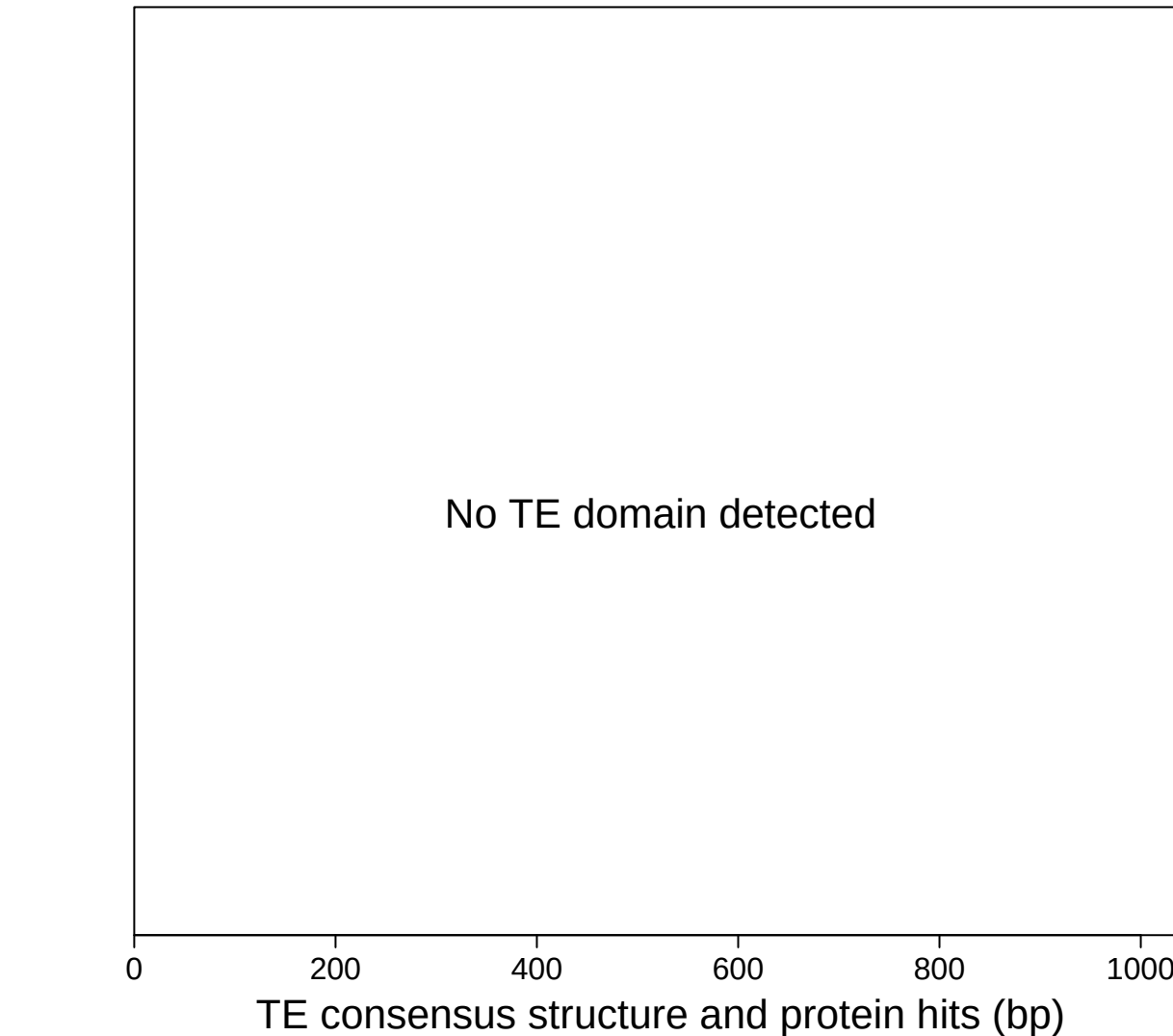
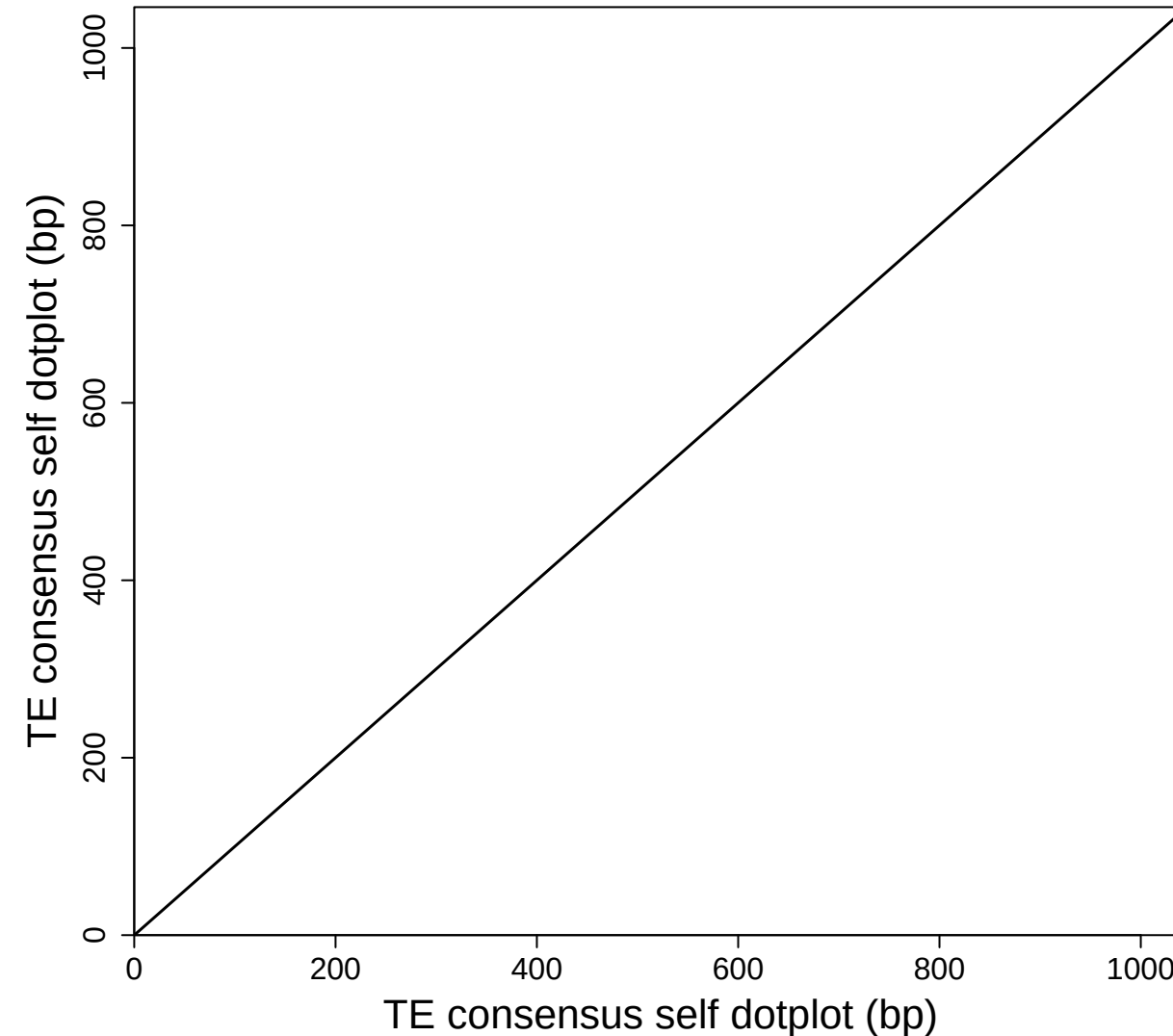
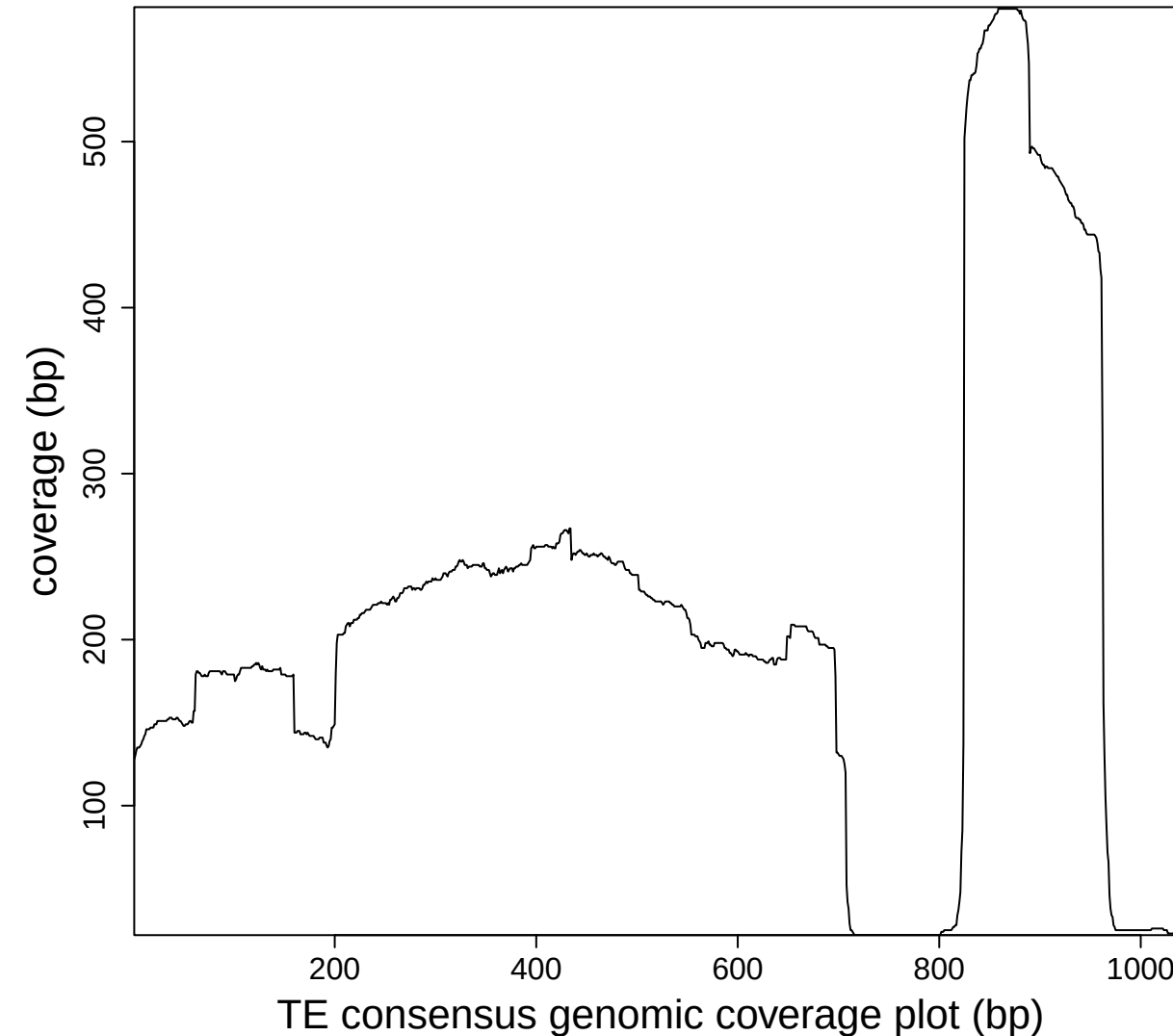
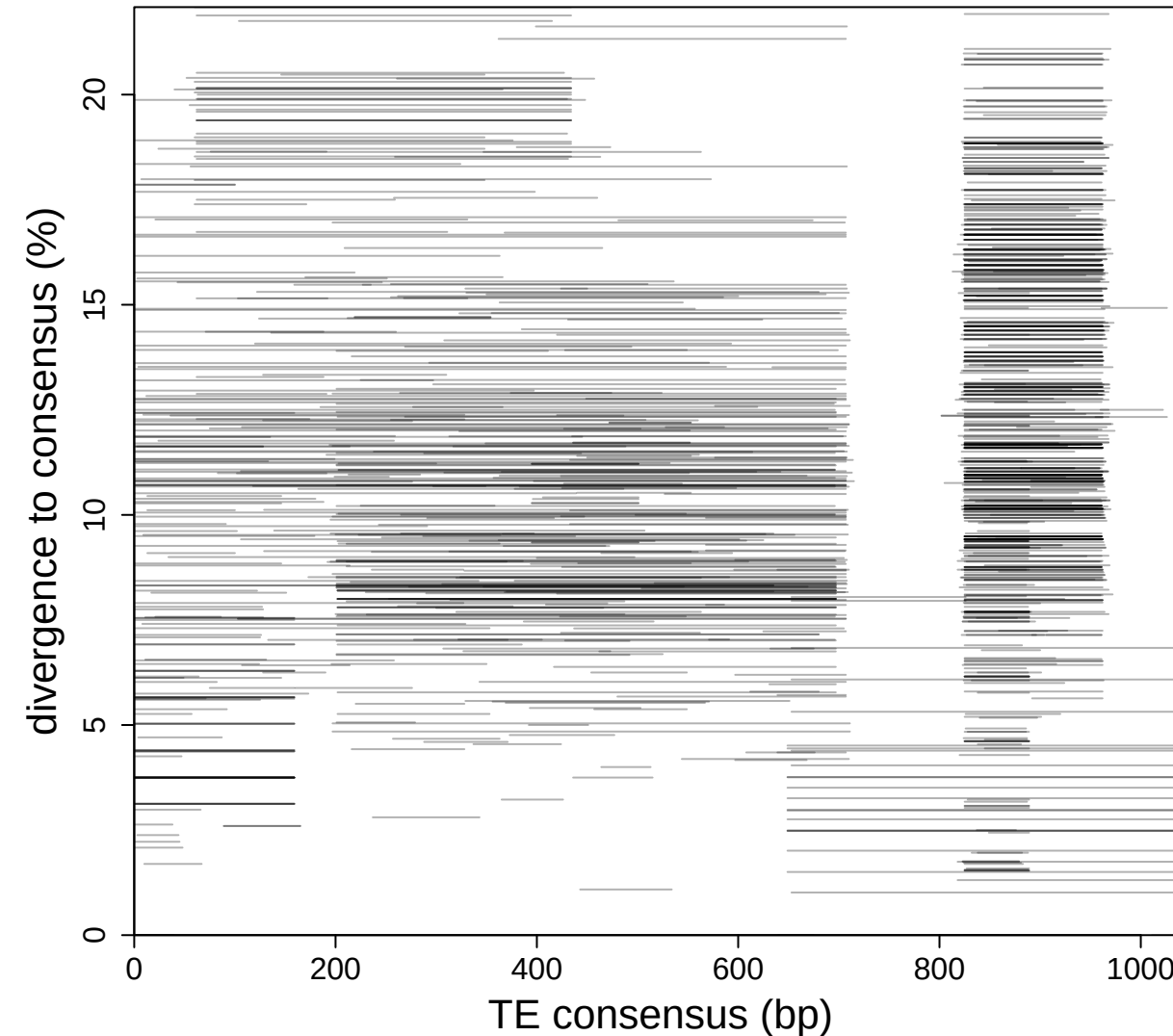
6.fasta.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcIn.fa aln.fa cl.f
size: 5392bp; fragments: 2816; full length: 0 (>=4852.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

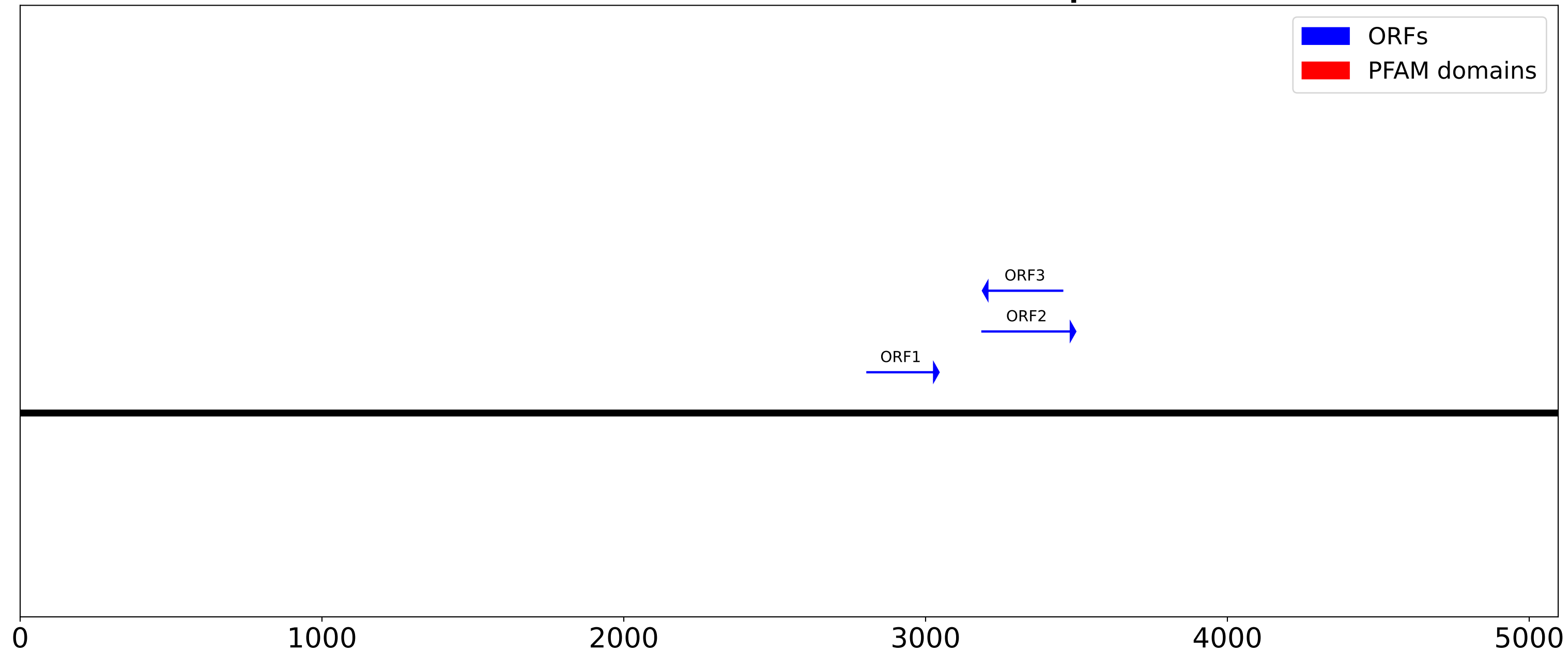


TE: rnd_5_family_6946
size: 1046bp; fragments: 1076; full length: 0 (>=941.4bp)

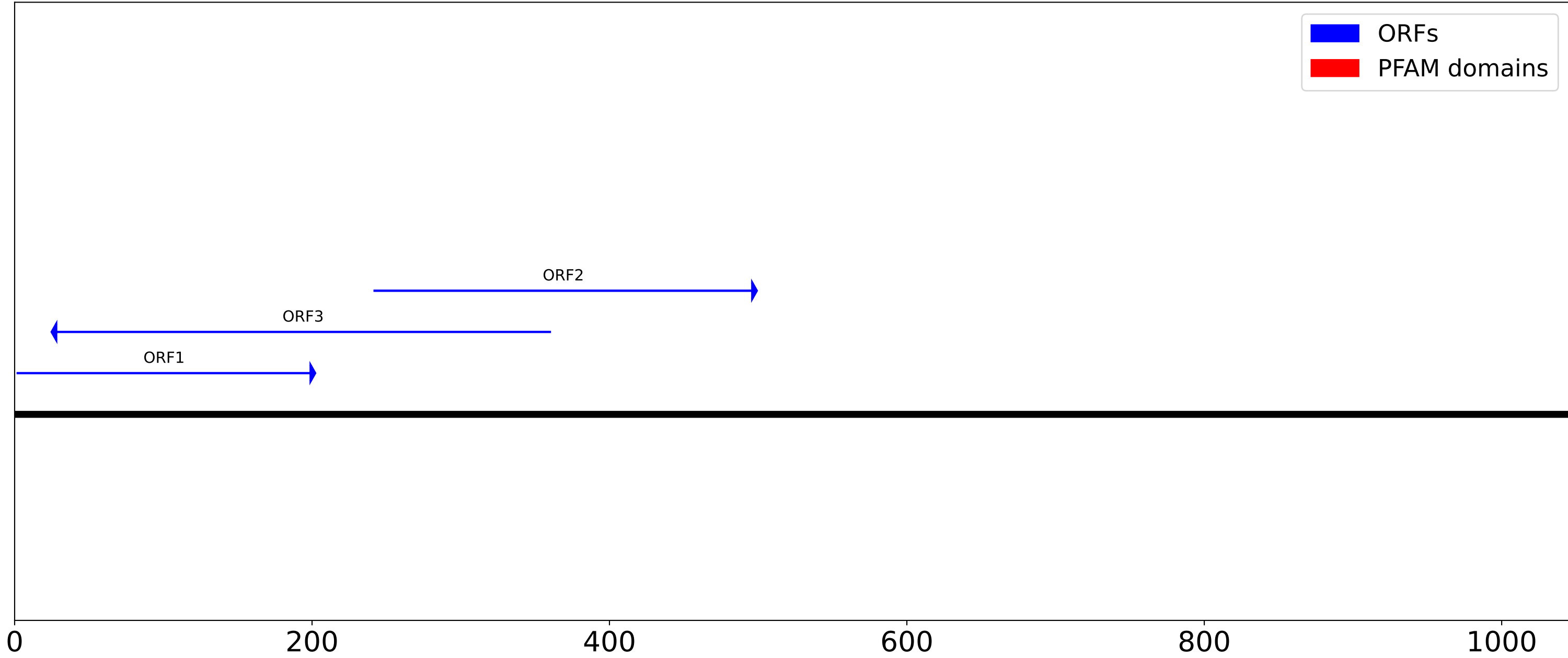
Before TEtrimmer 1046 bp



After TEtrimmer ORF and PFAM domain plot



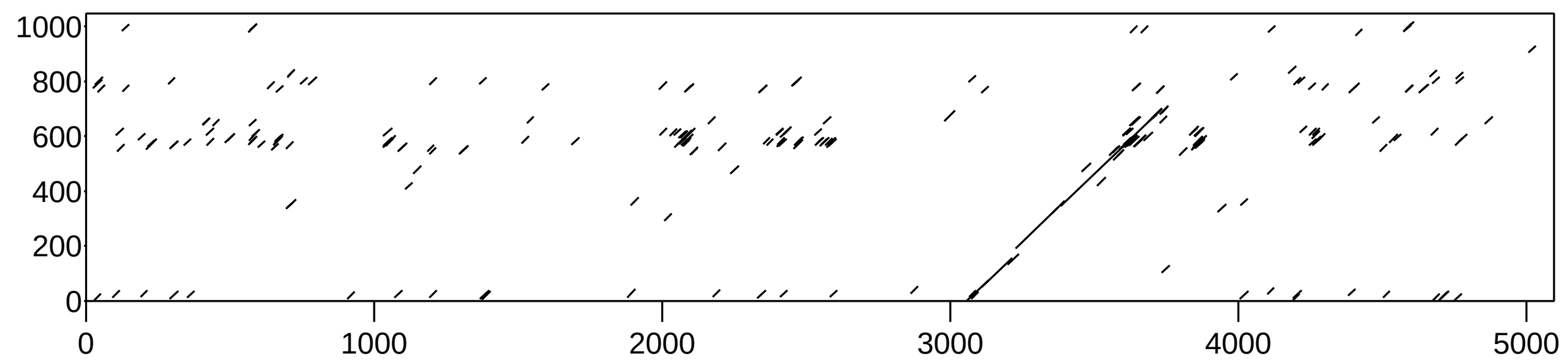
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)