

Start Crop Point



End Crop Point

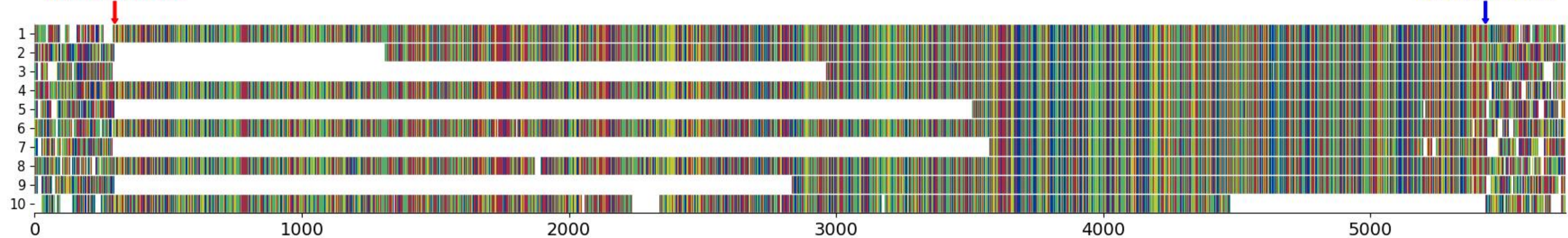


MSA length = 5133



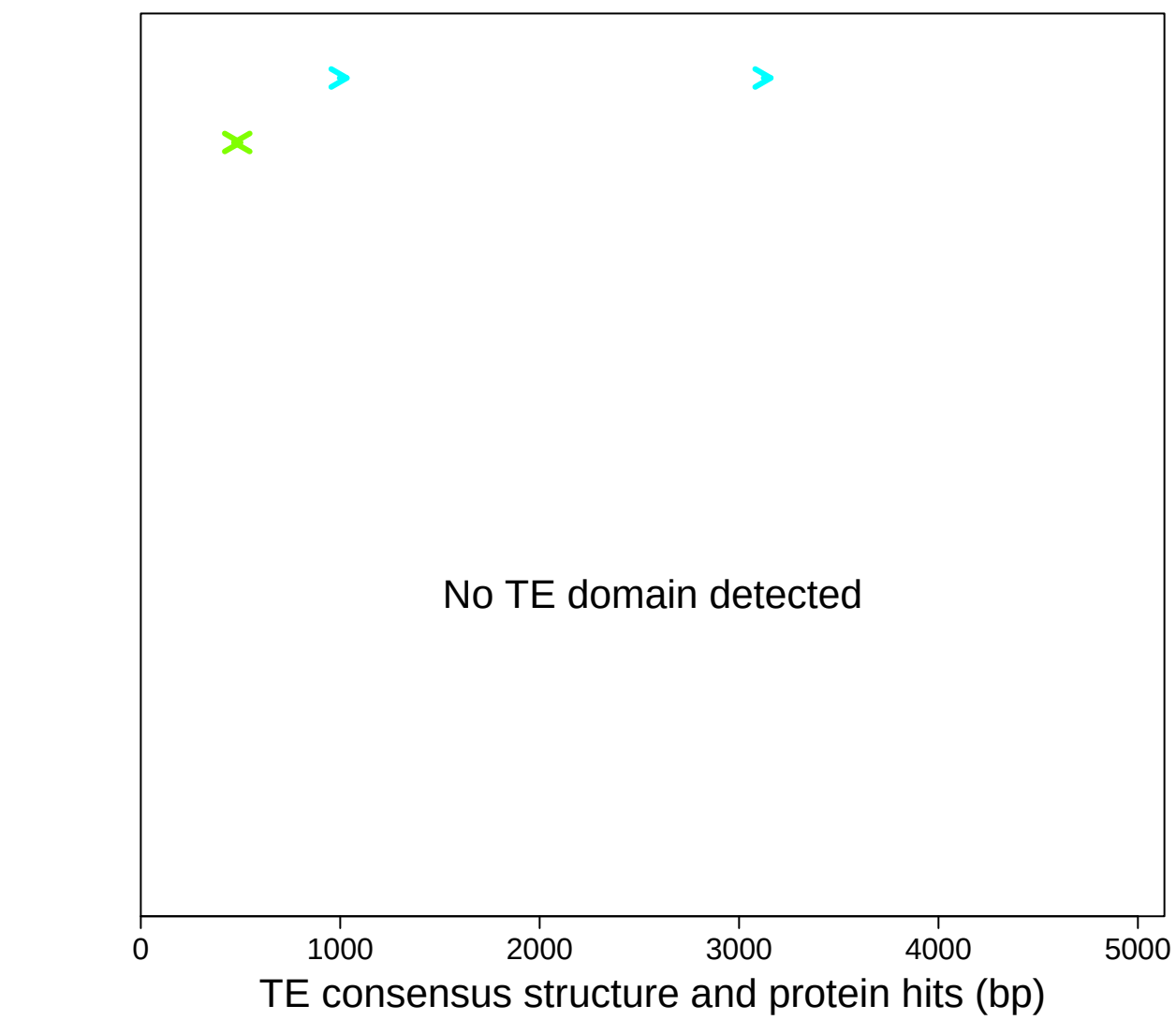
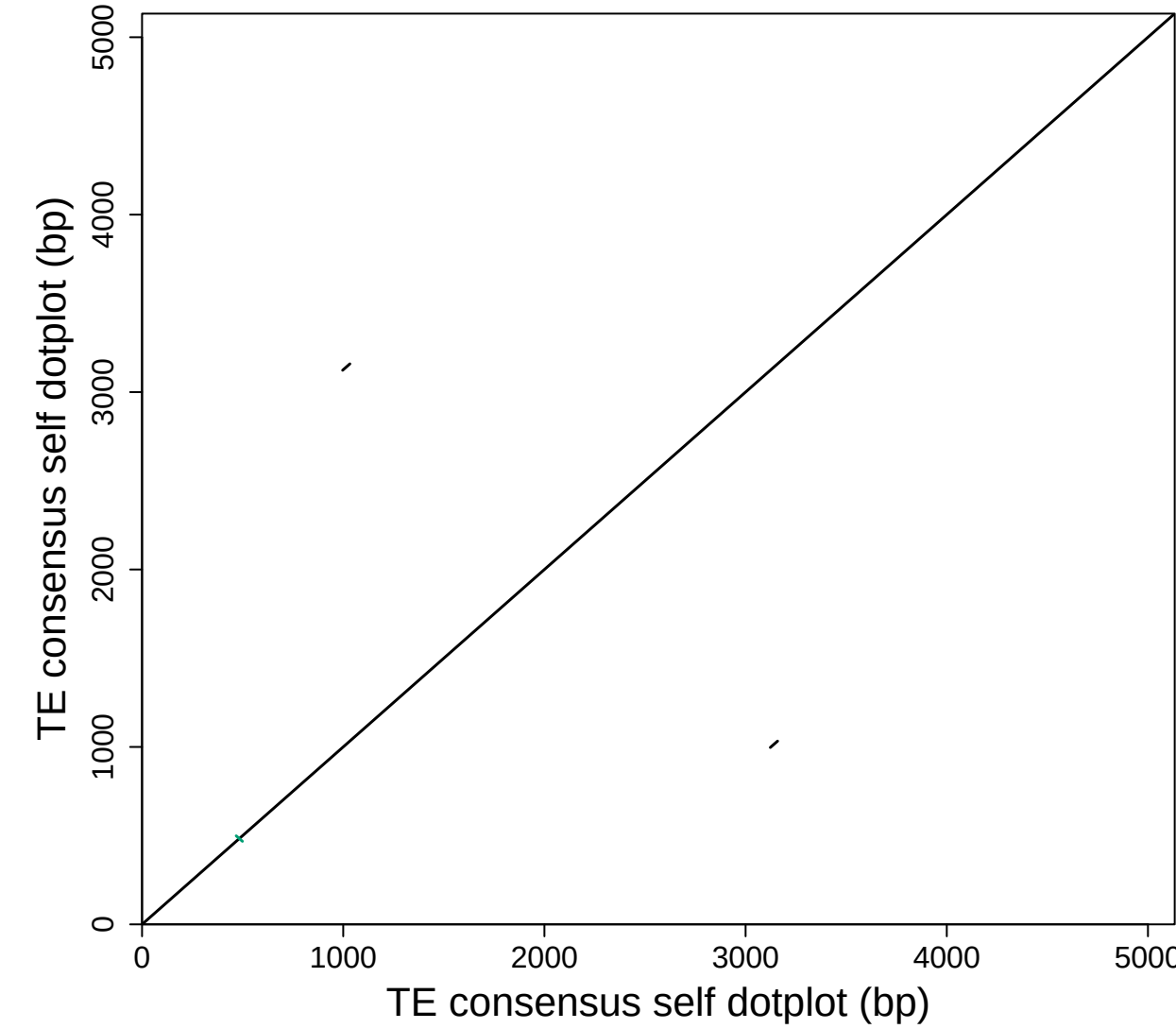
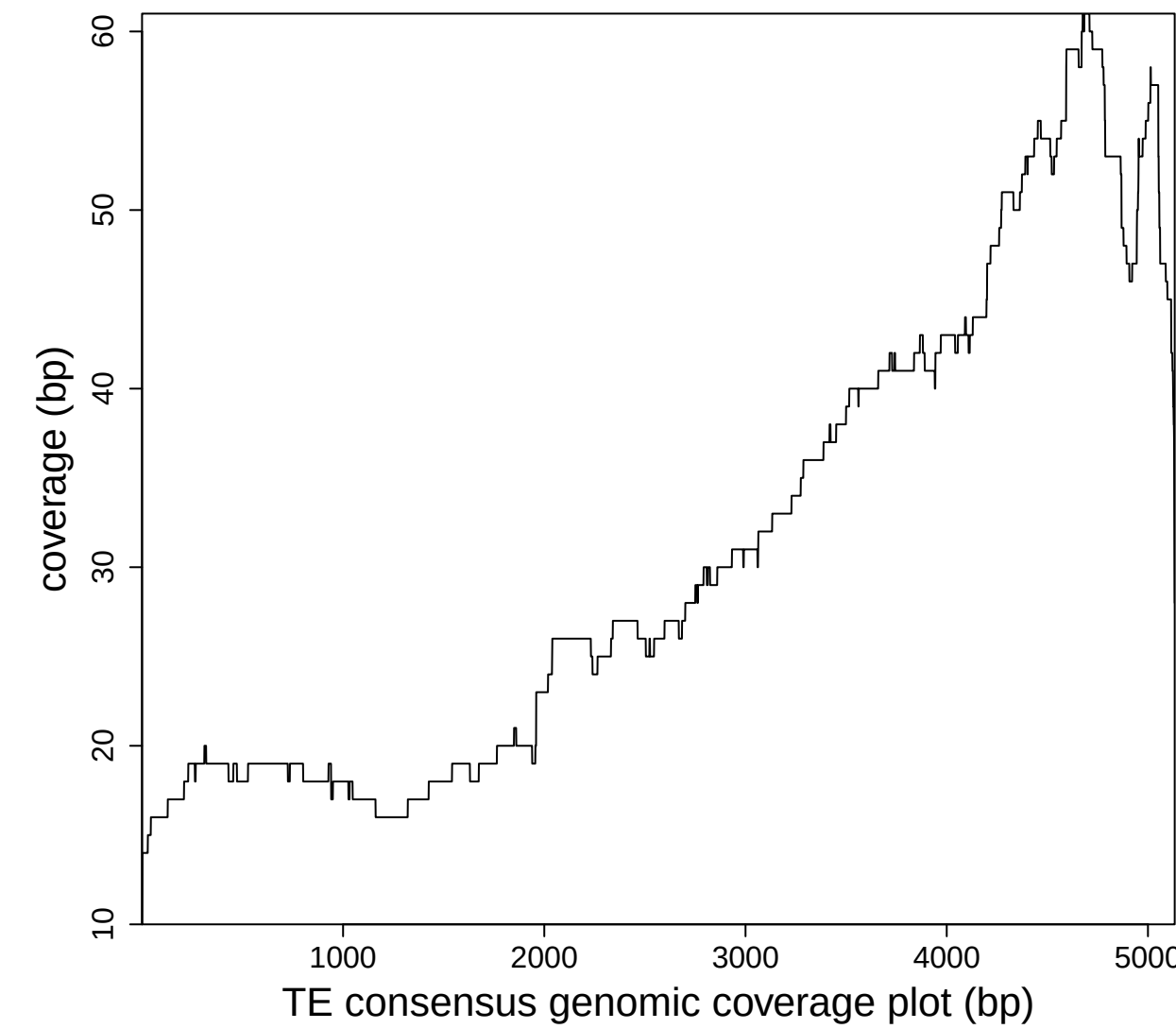
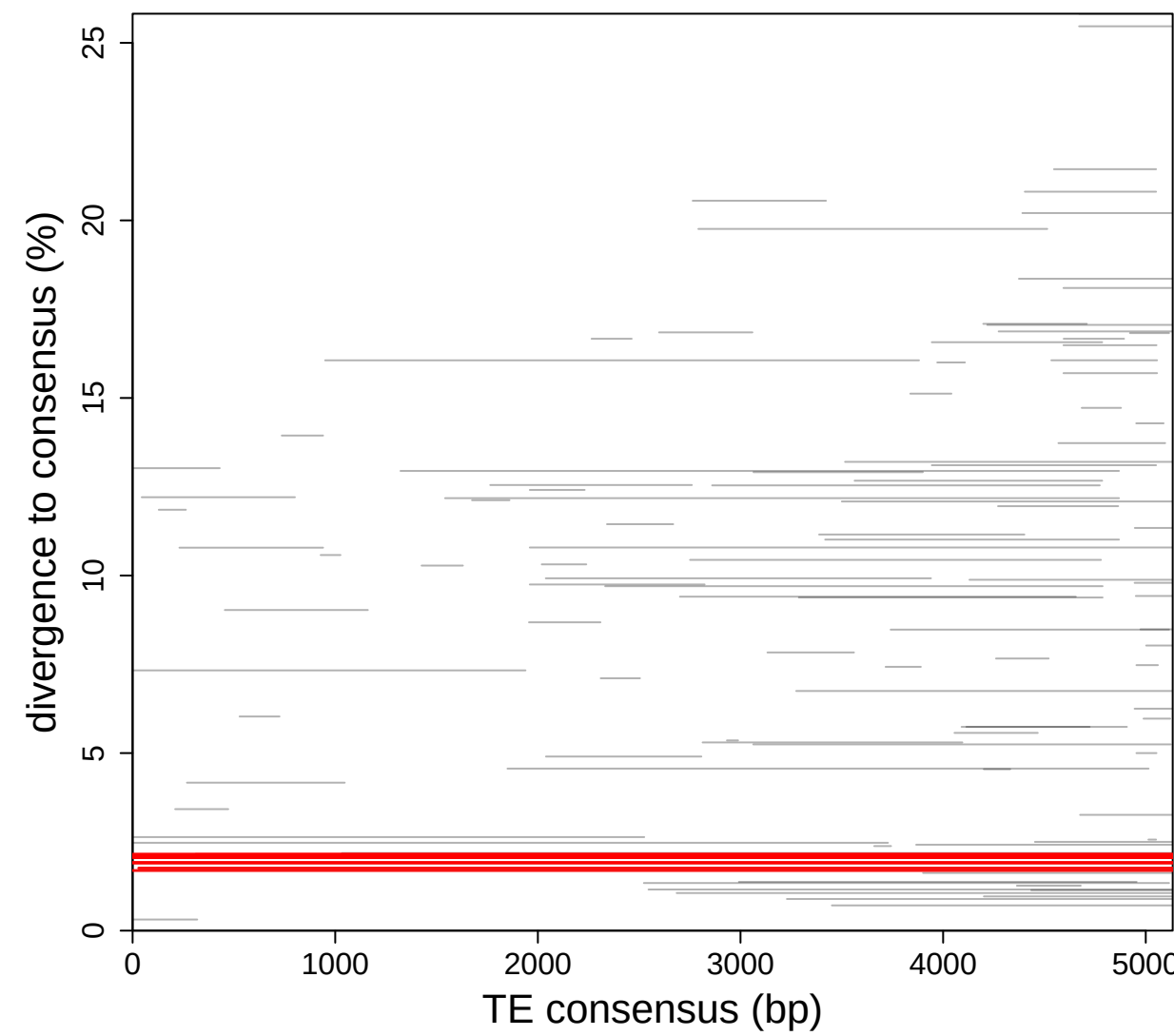
Start crop Point

End crop Point



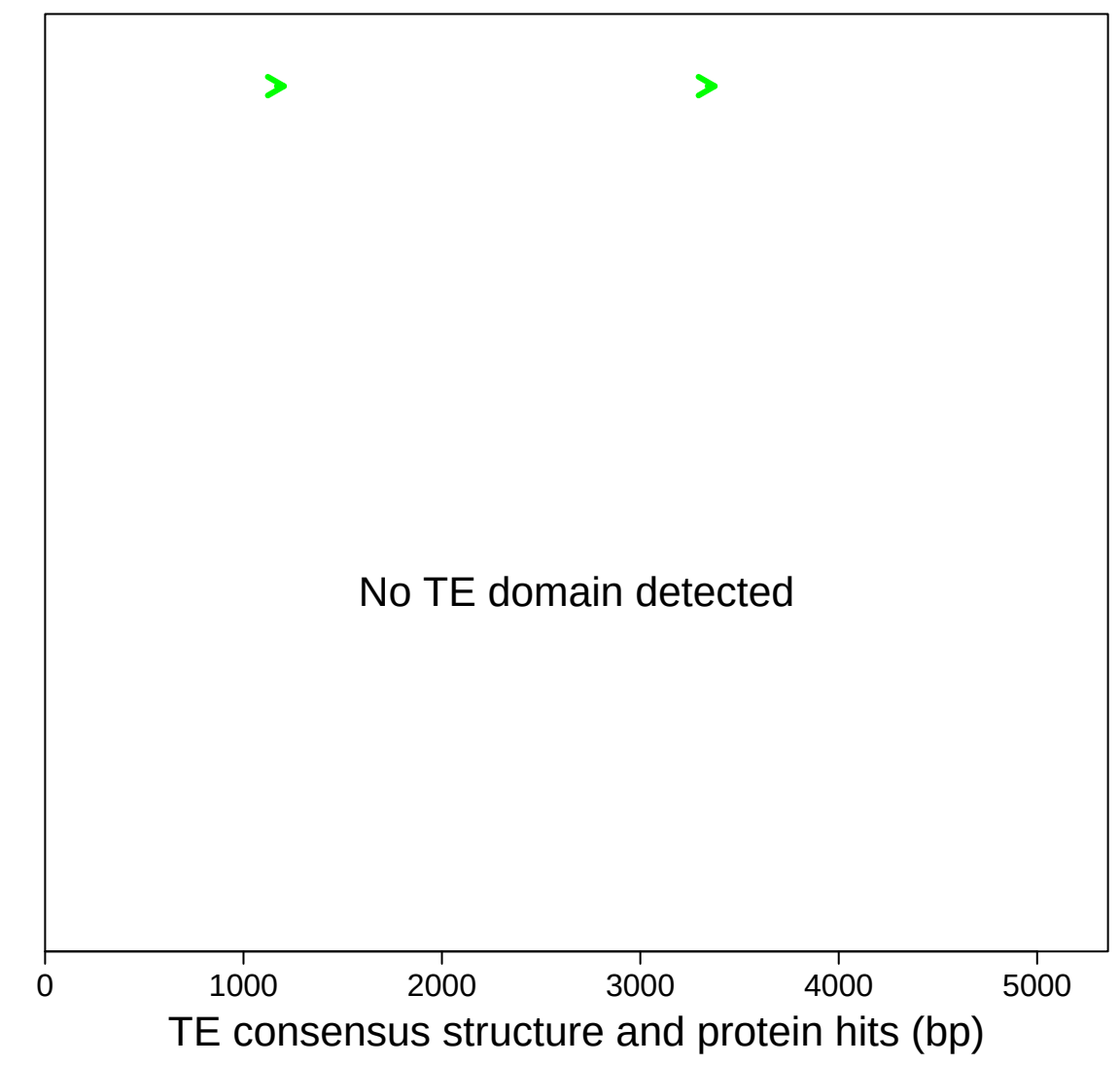
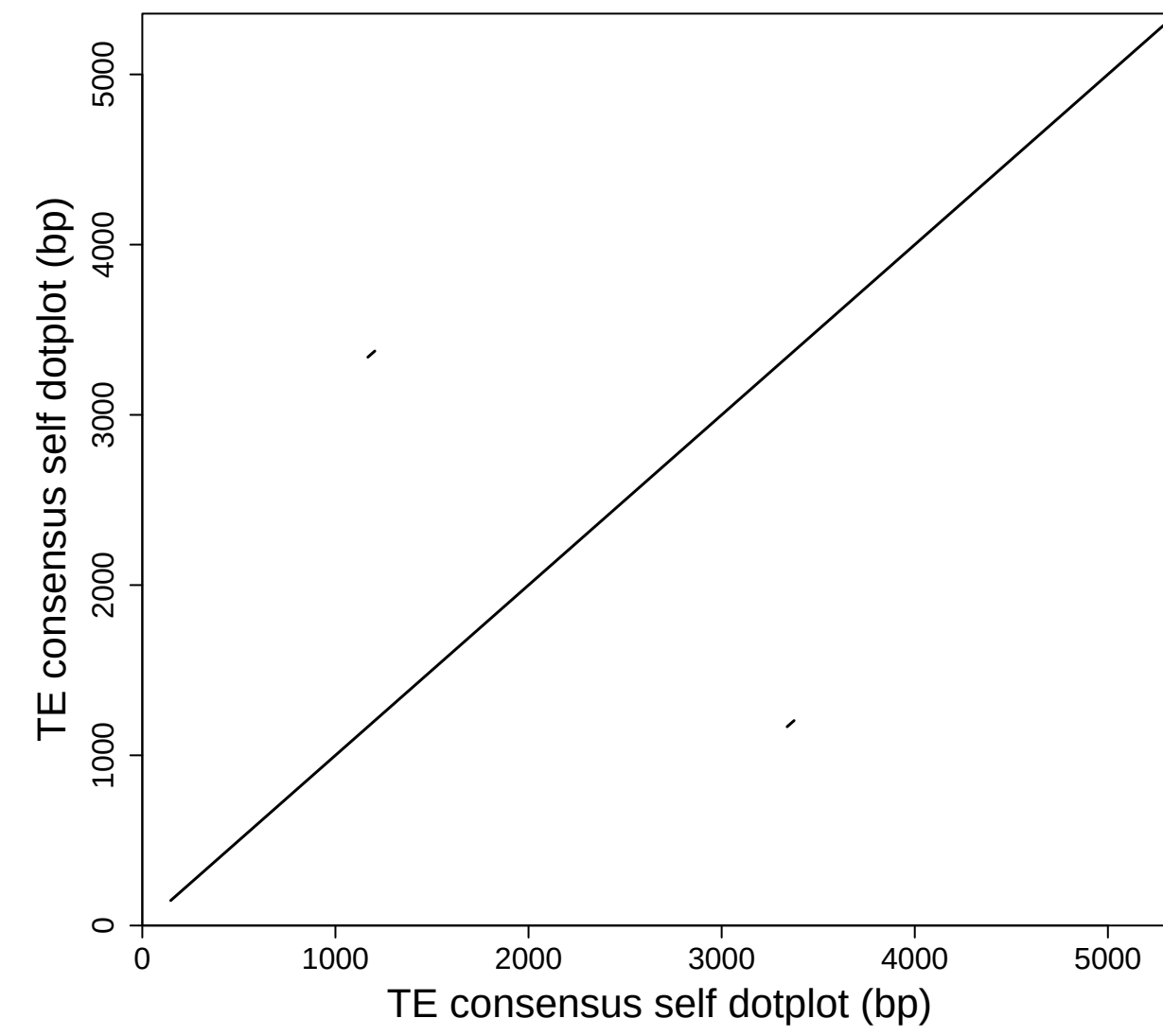
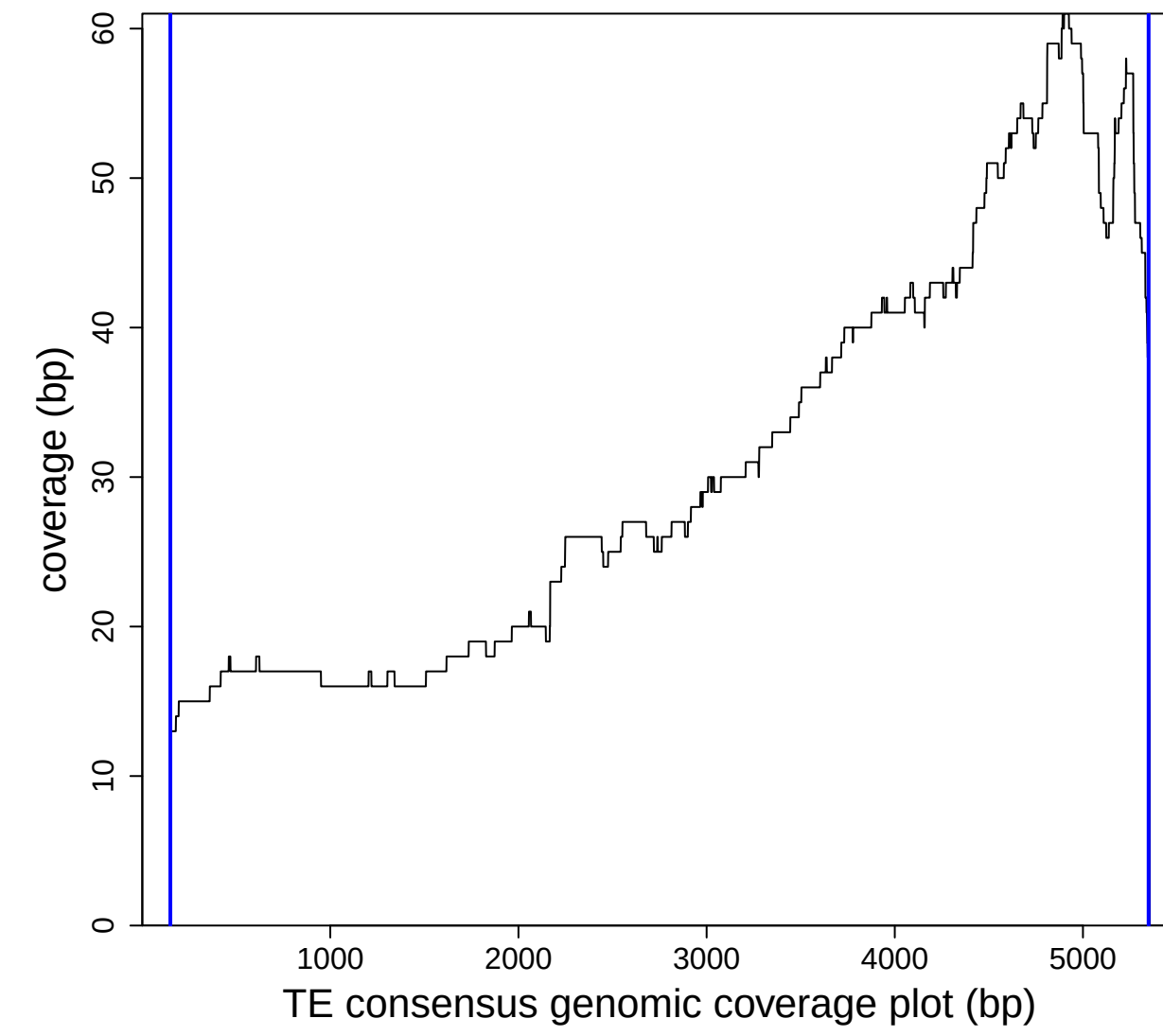
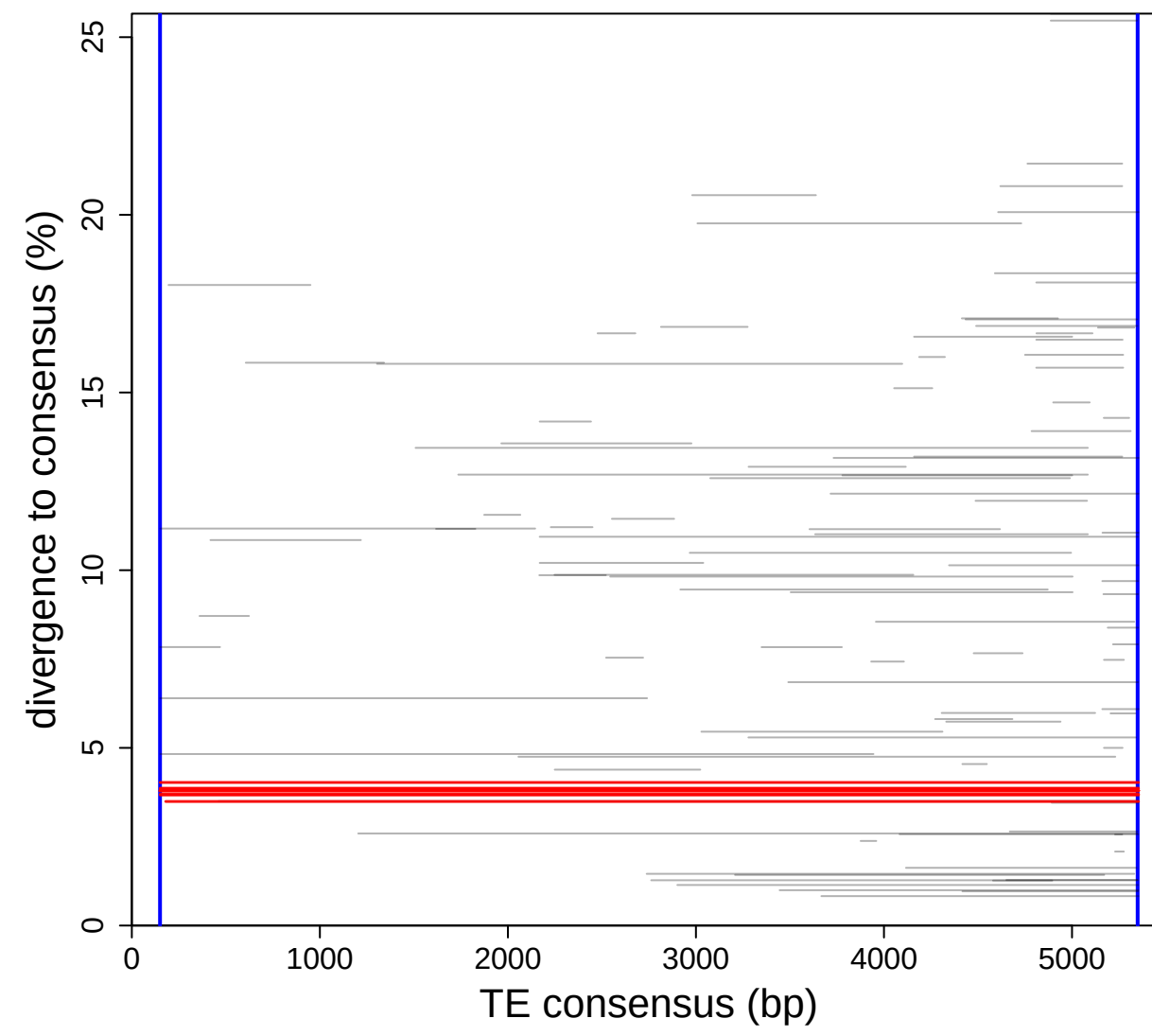
bed_uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa c
size: 5133bp; fragments: 118; full length: 11 (≥ 4619.7 bp)

After TEtrimmer 5133 bp

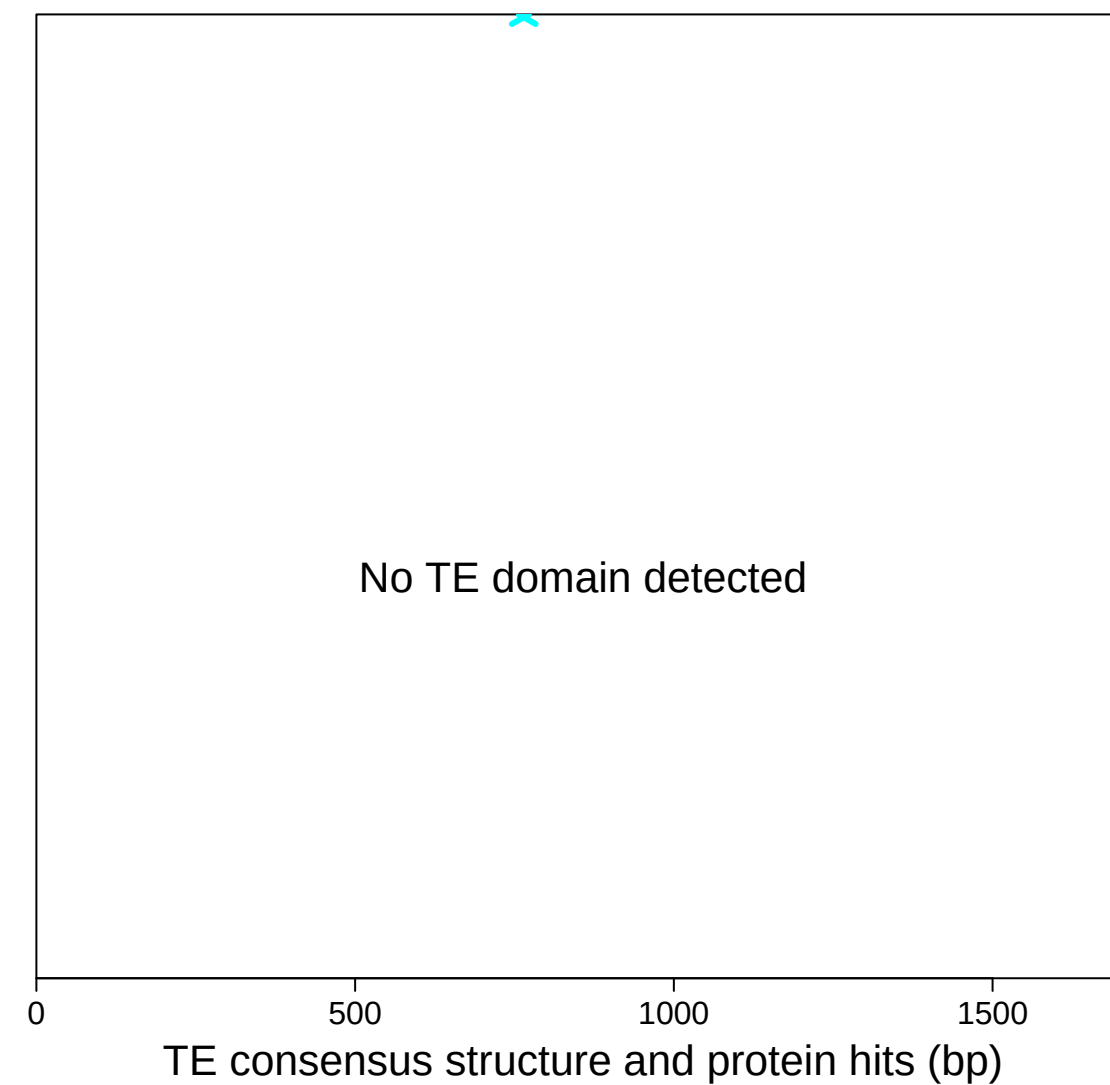
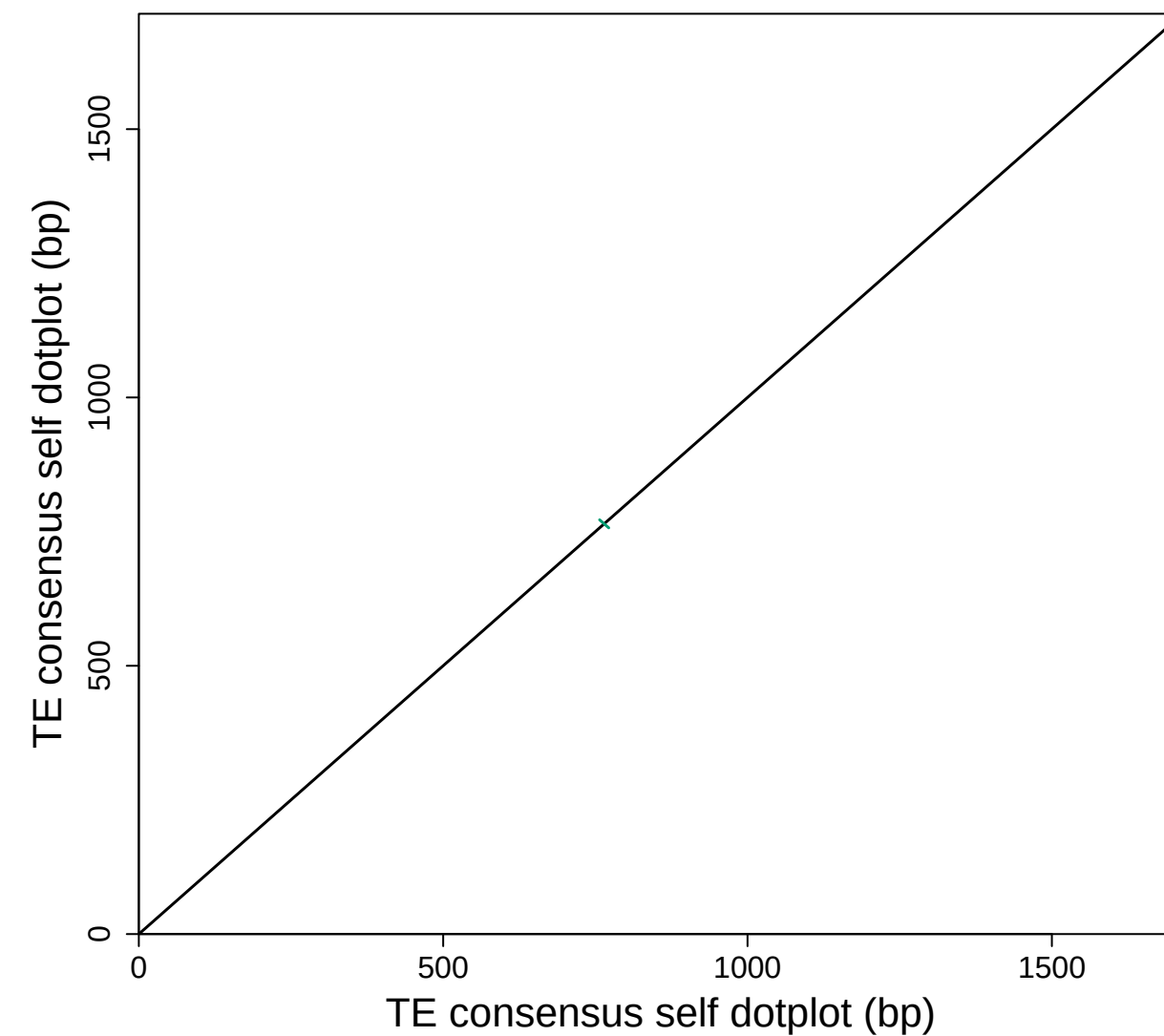
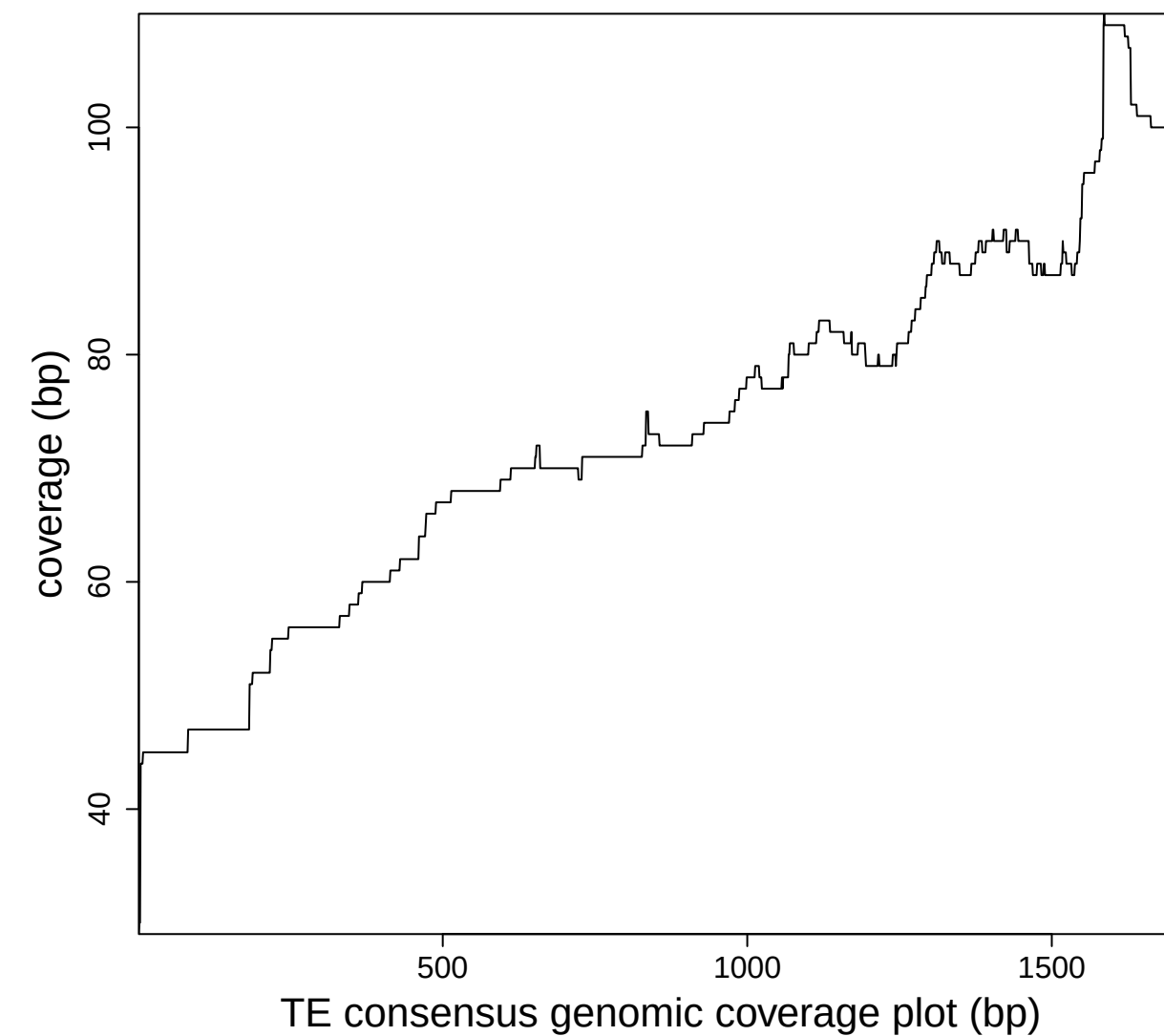
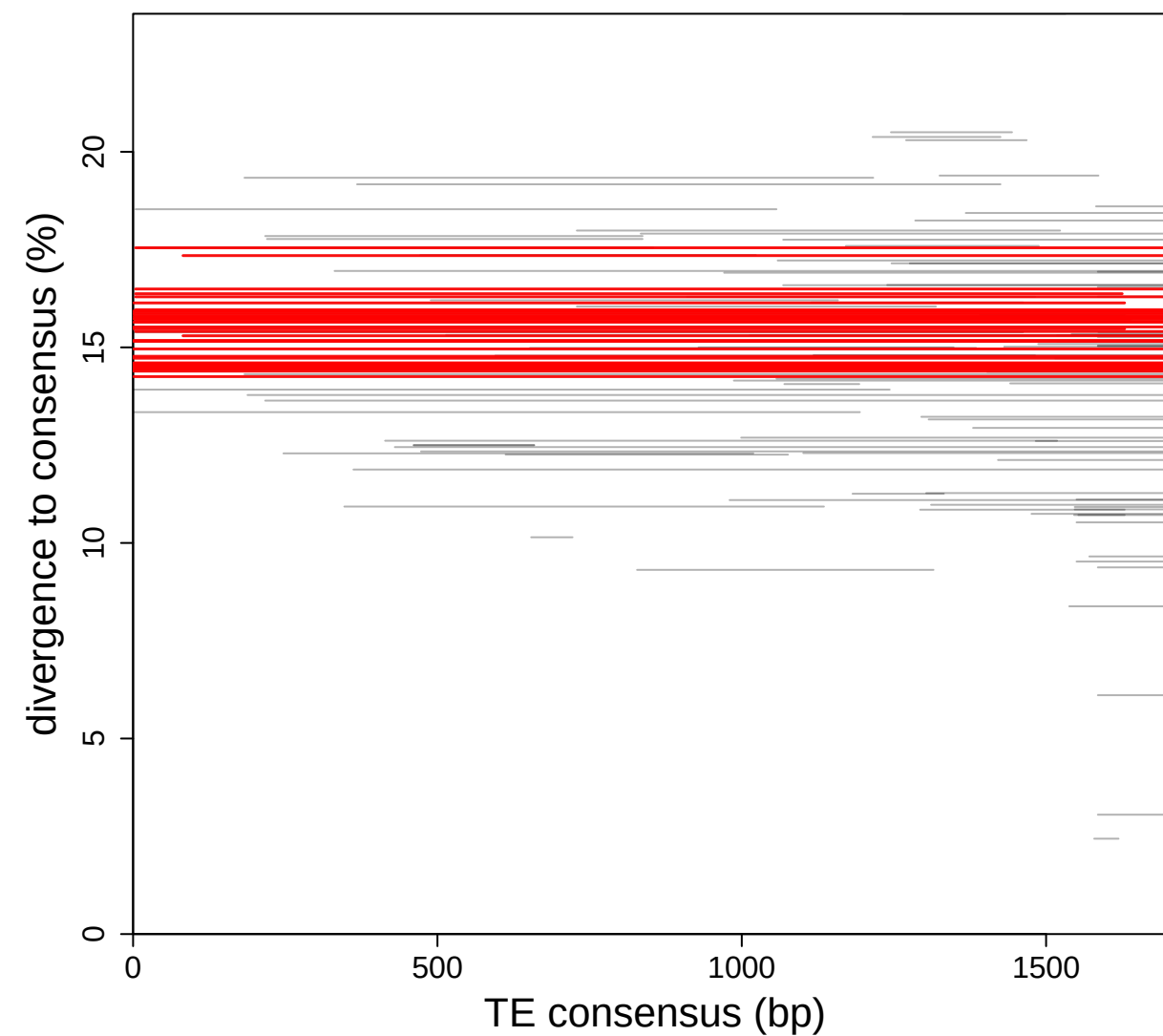


2.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 5499bp; fragments: 111; full length: 10 (>=4949.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

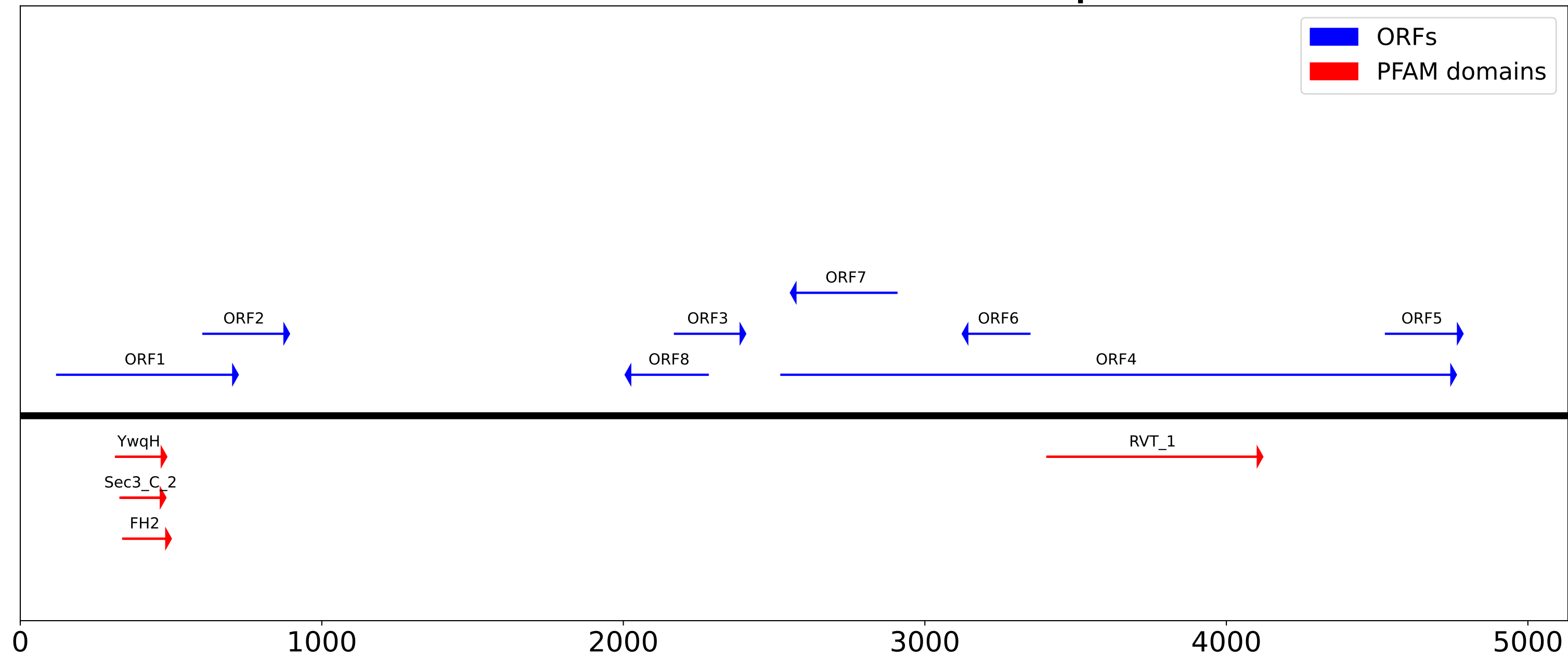


TE: rnd_4_family_1352
size: 1715bp; fragments: 146; full length: 42 (≥ 1543.5 bp)

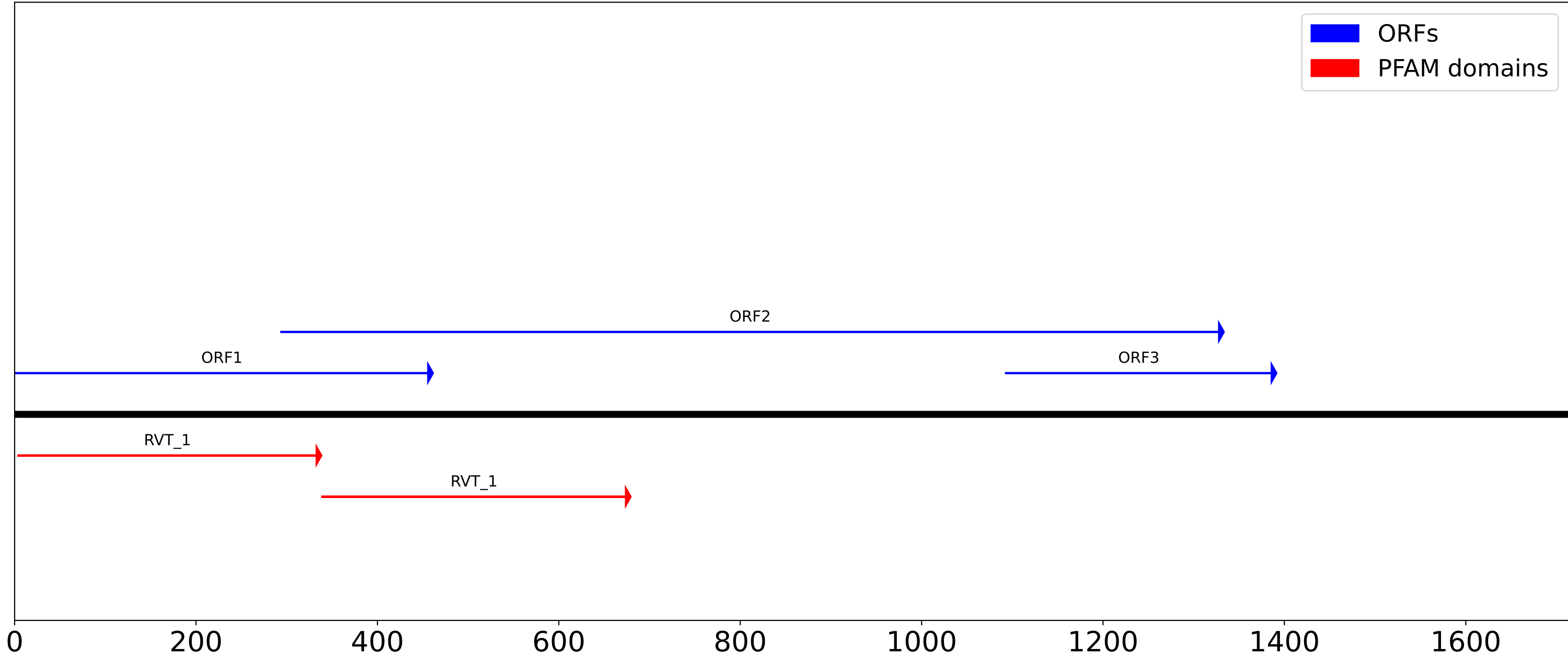


Before TEtrimmer 1715 bp

After TEtrimmer ORF and PFAM domain plot



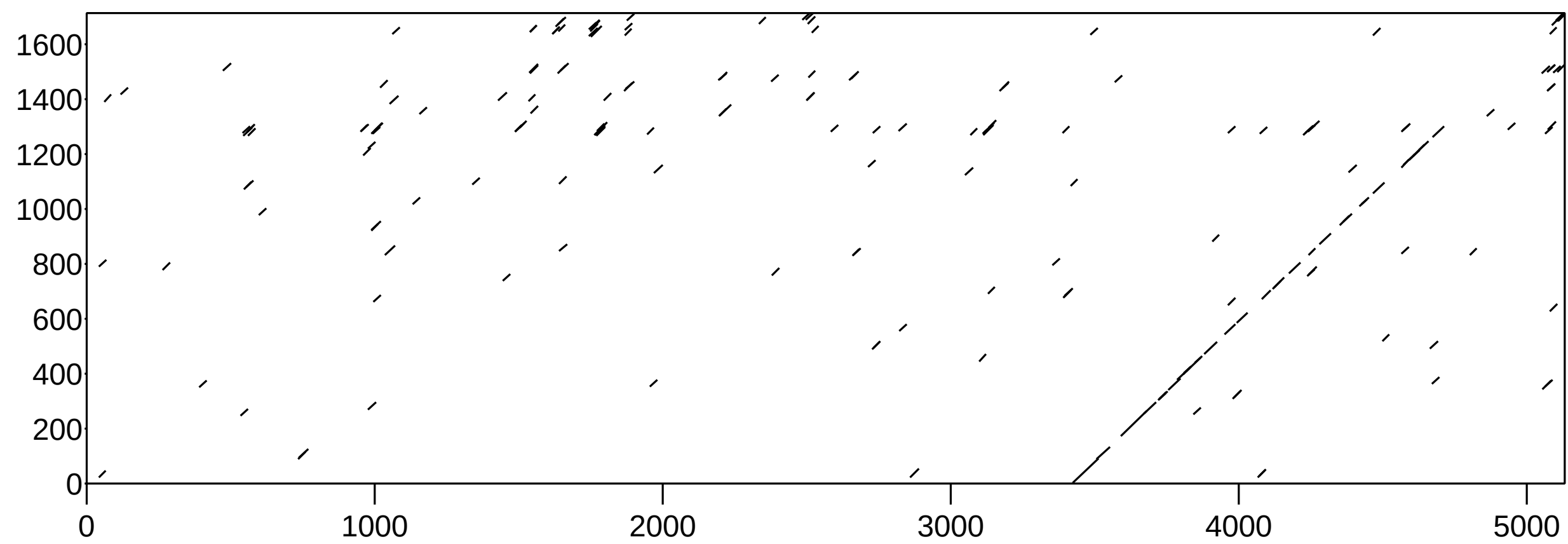
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)