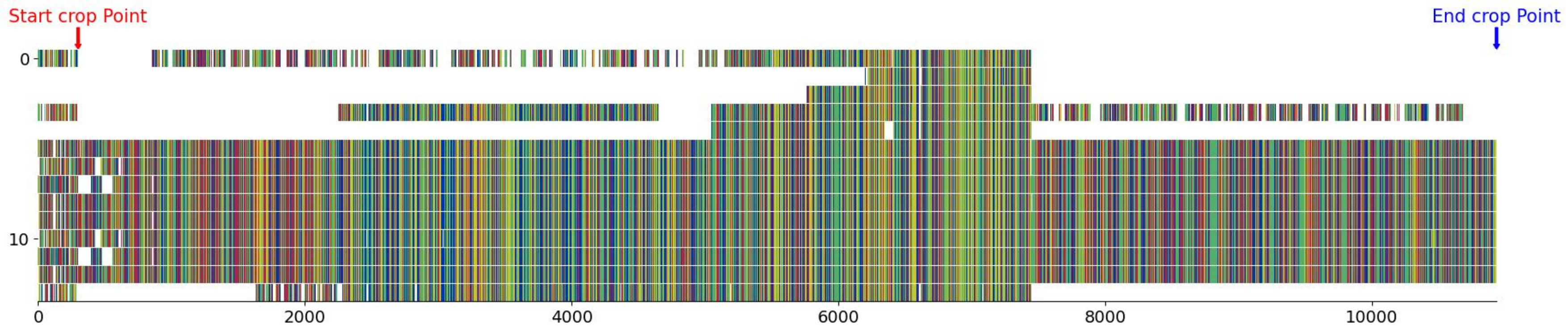


Start crop Point

End crop Point

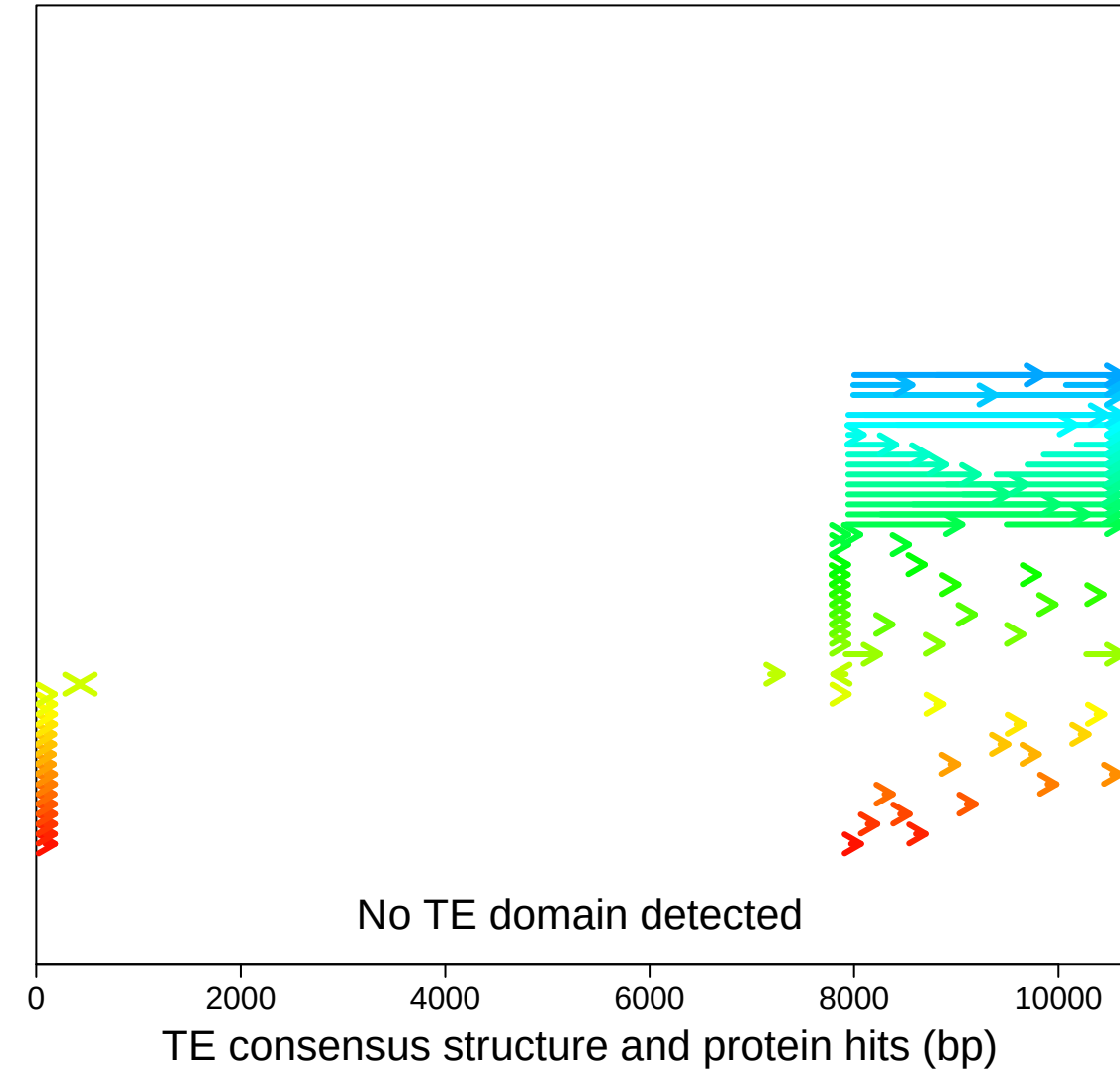
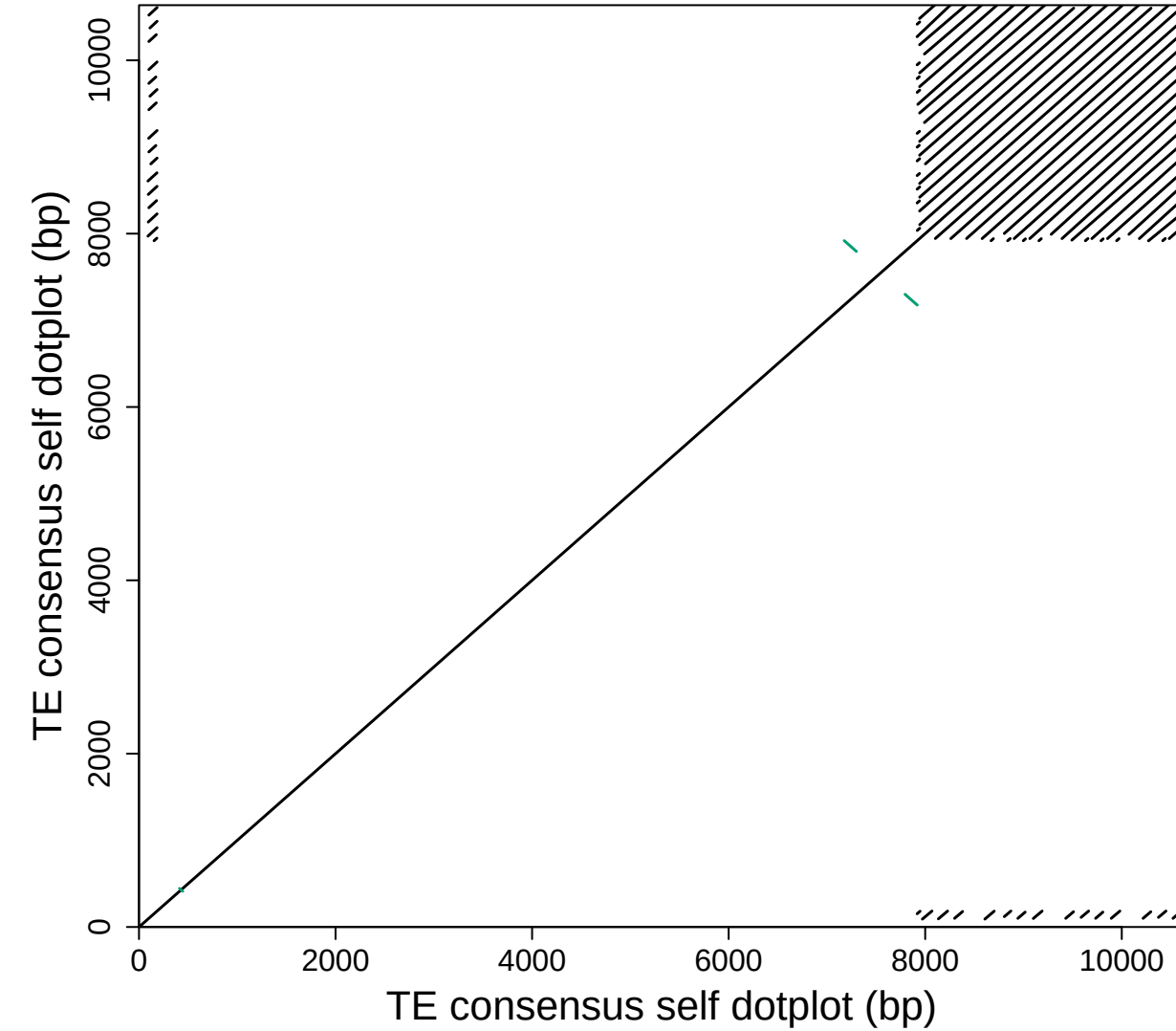
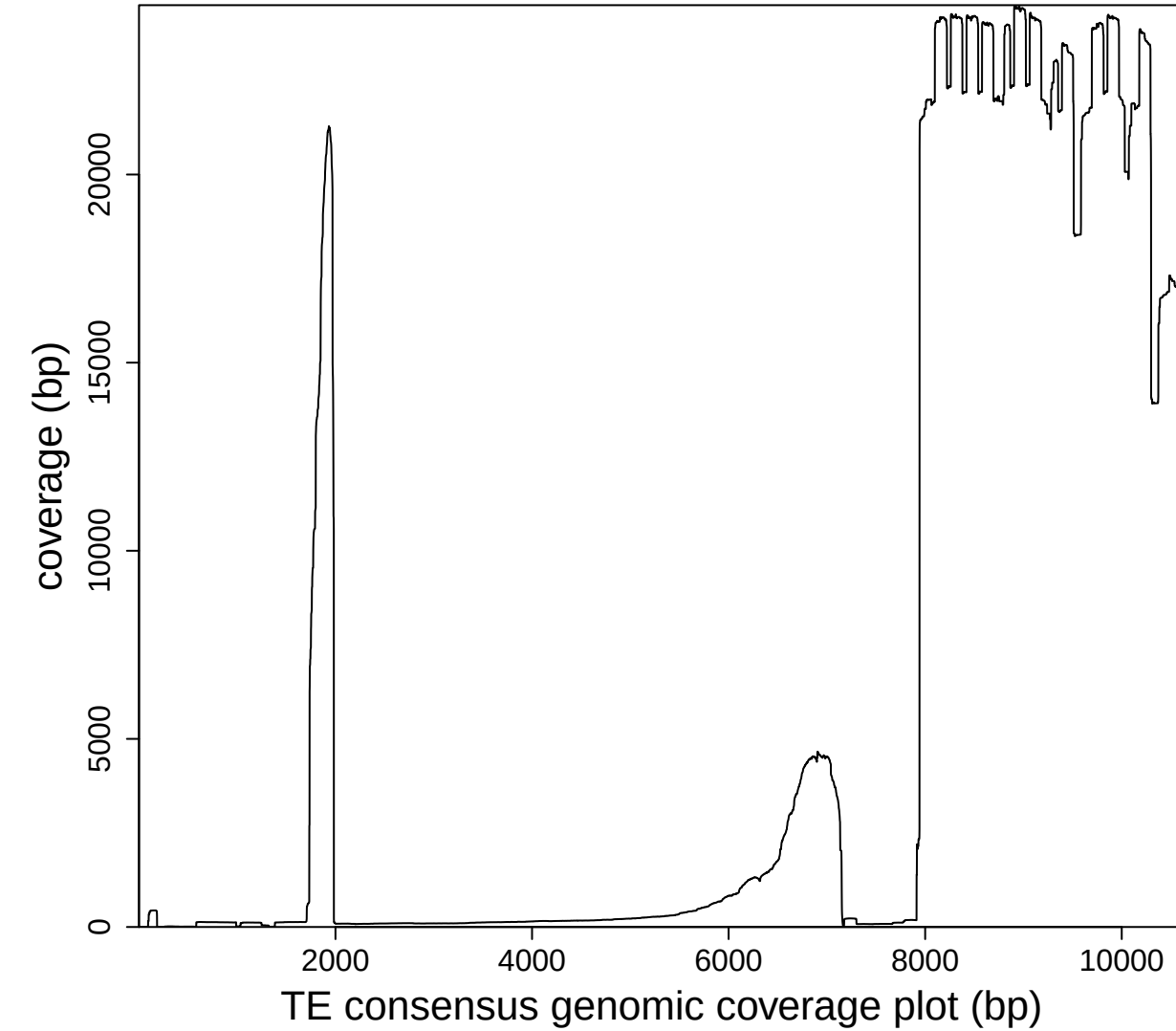
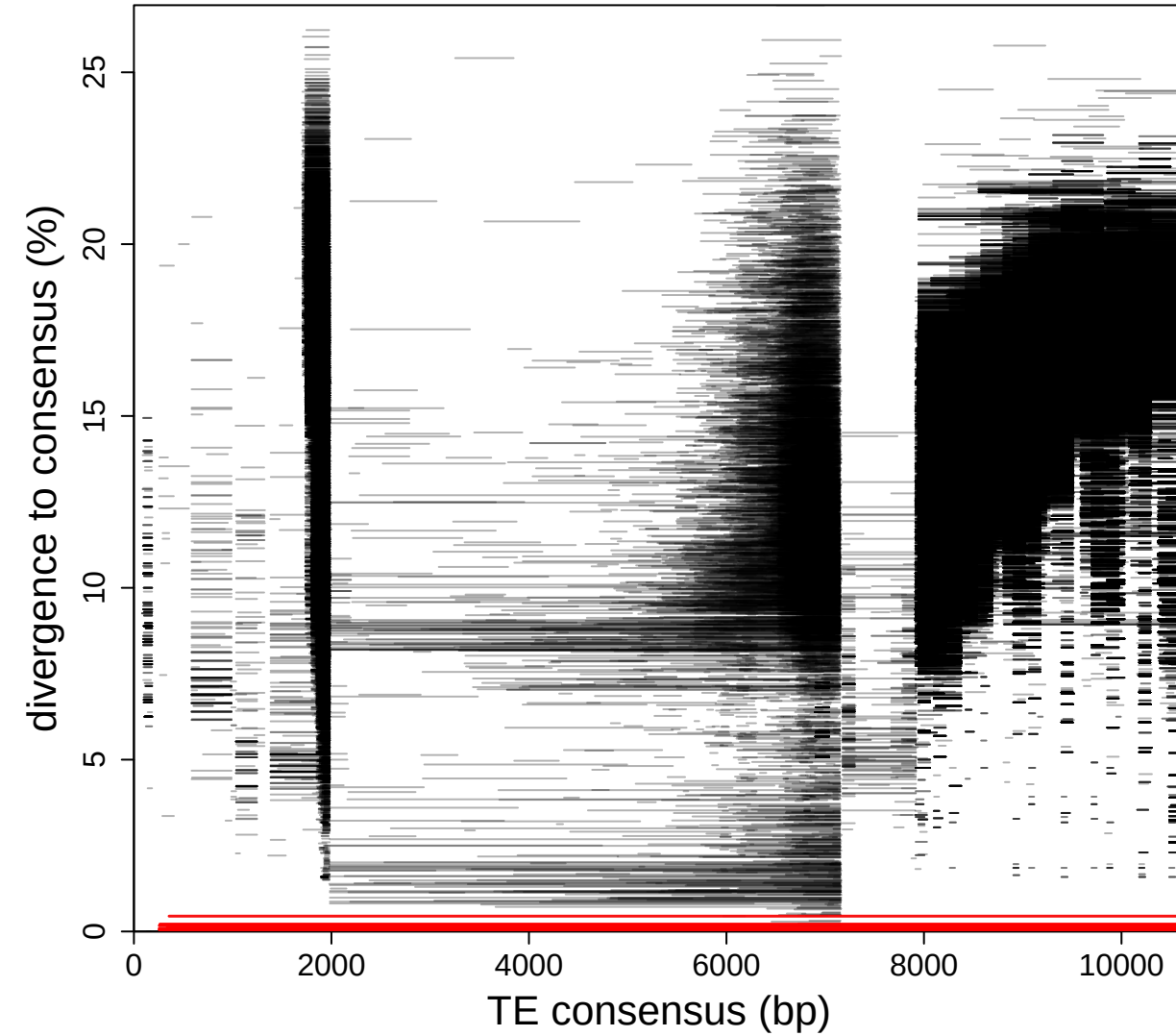
MSA length = 10636





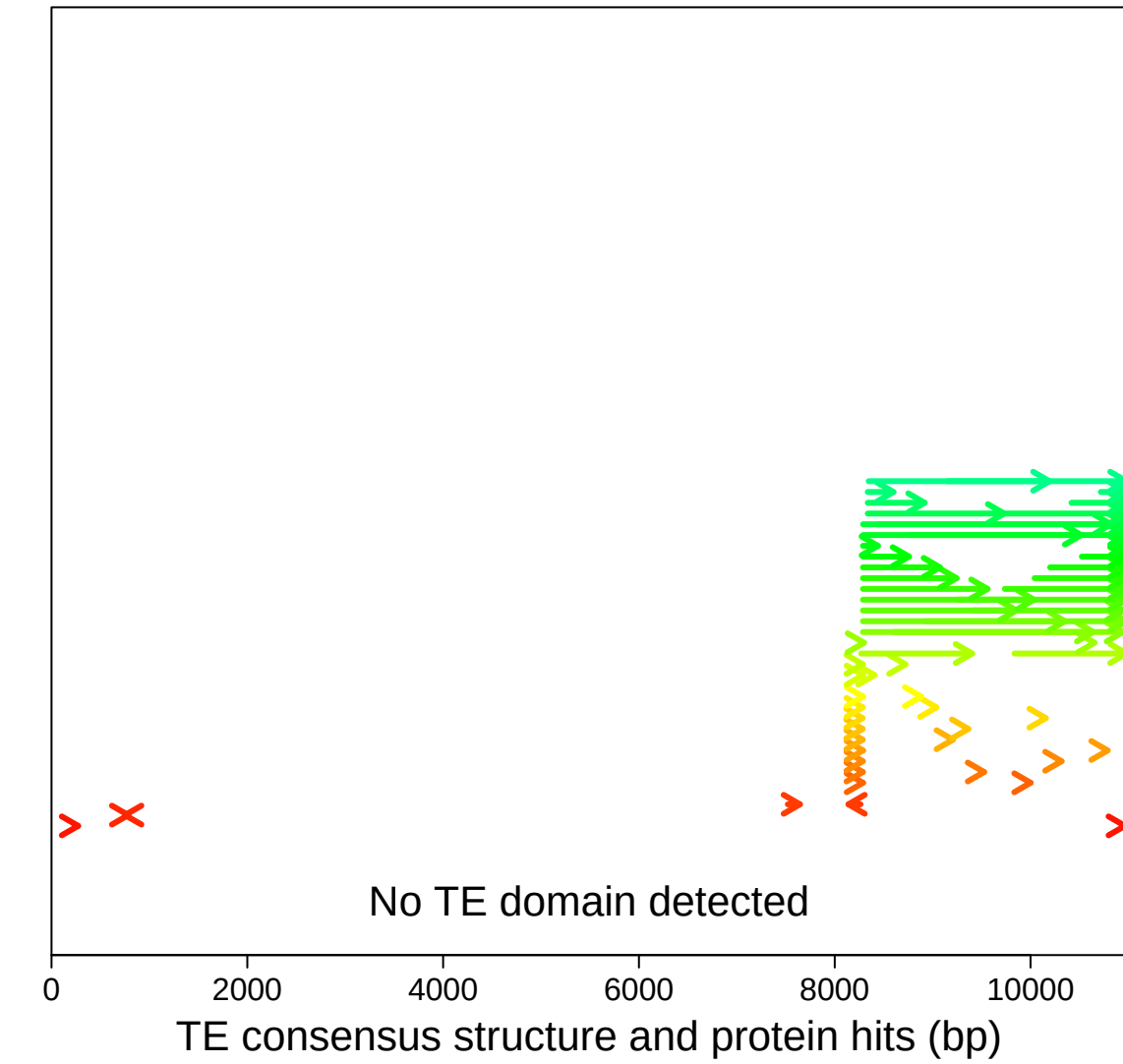
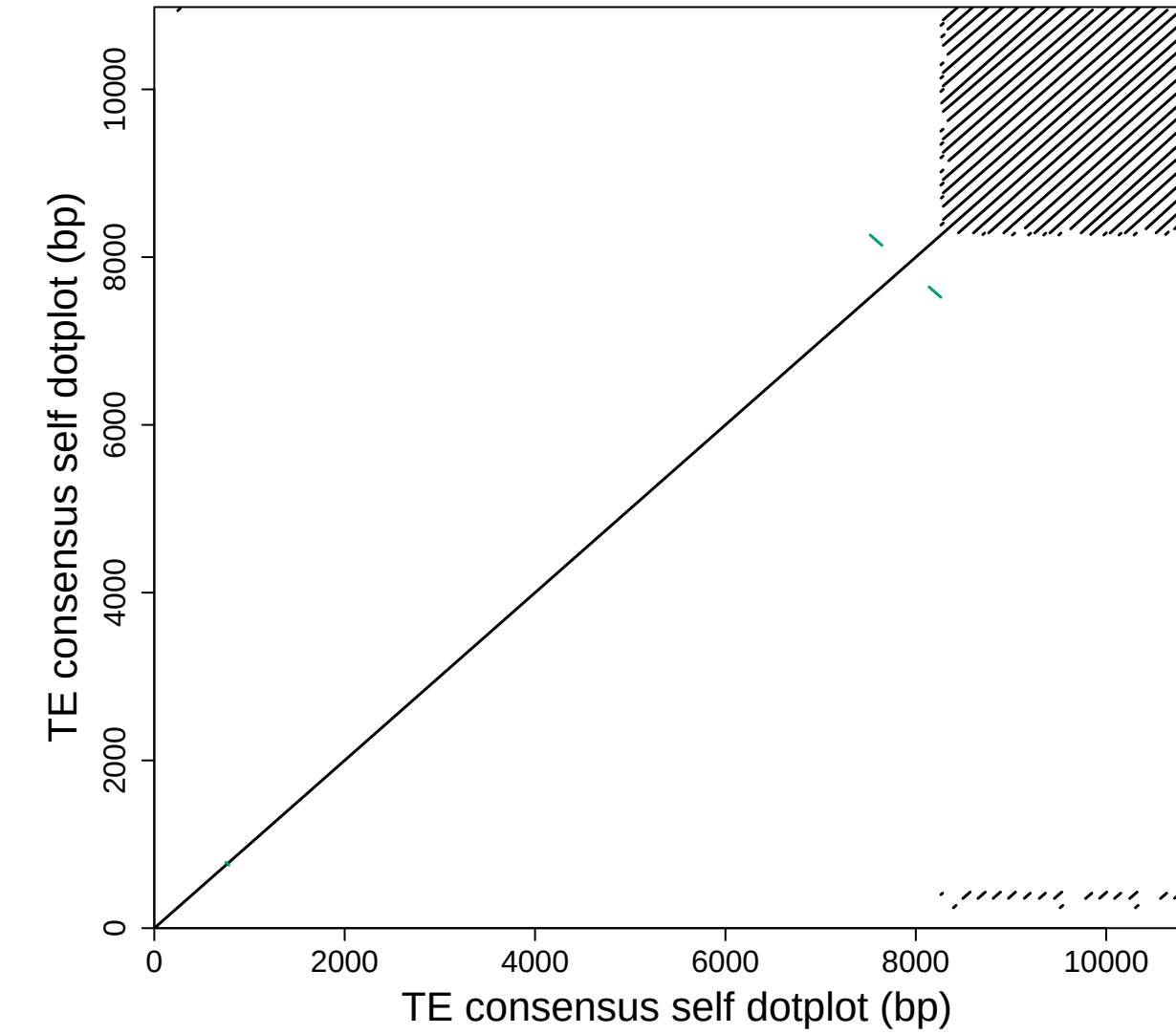
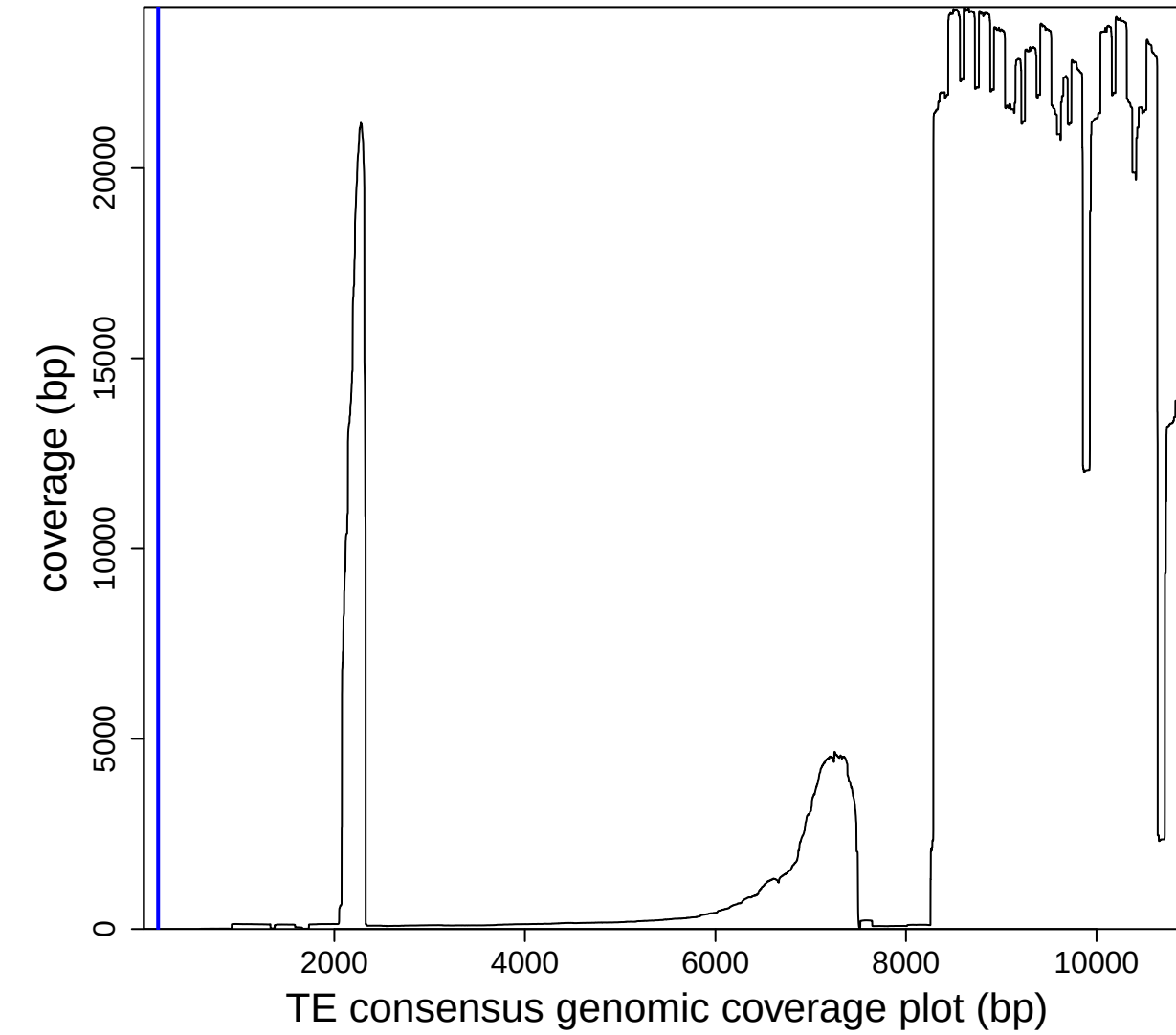
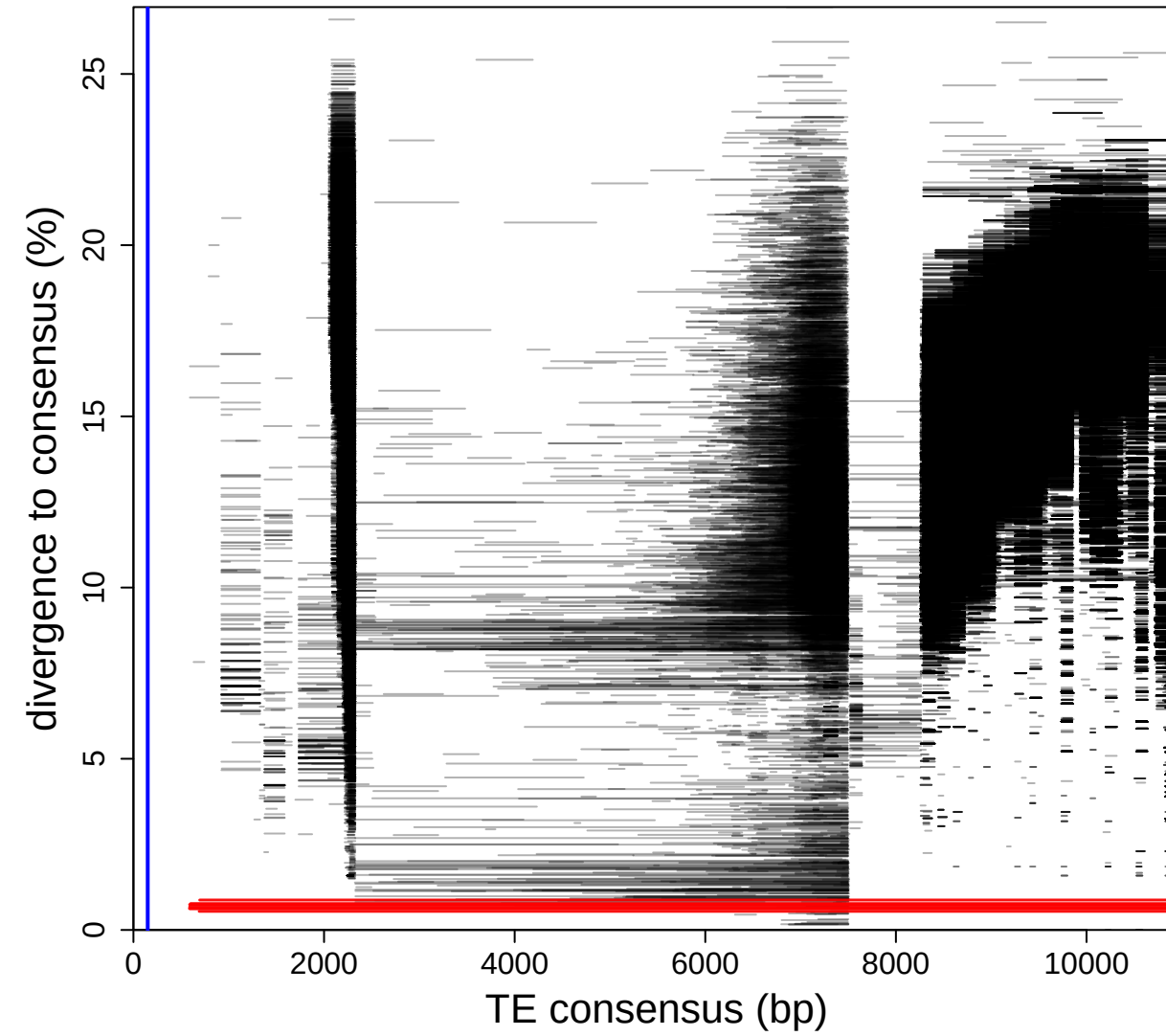
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 10636bp; fragments: 88300; full length: 8 (>=9572.4bp)

After TEtrimmer 10636 bp



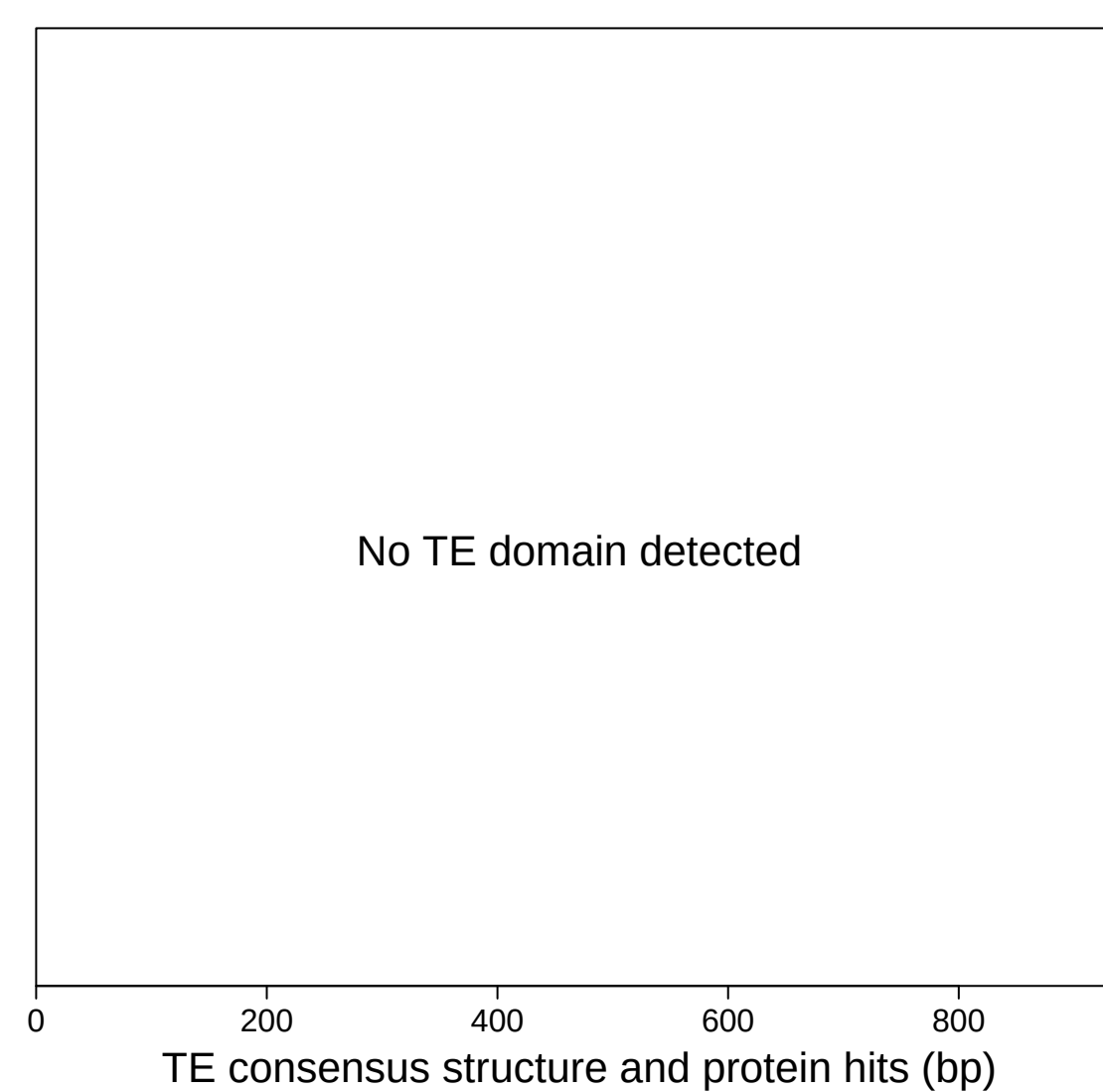
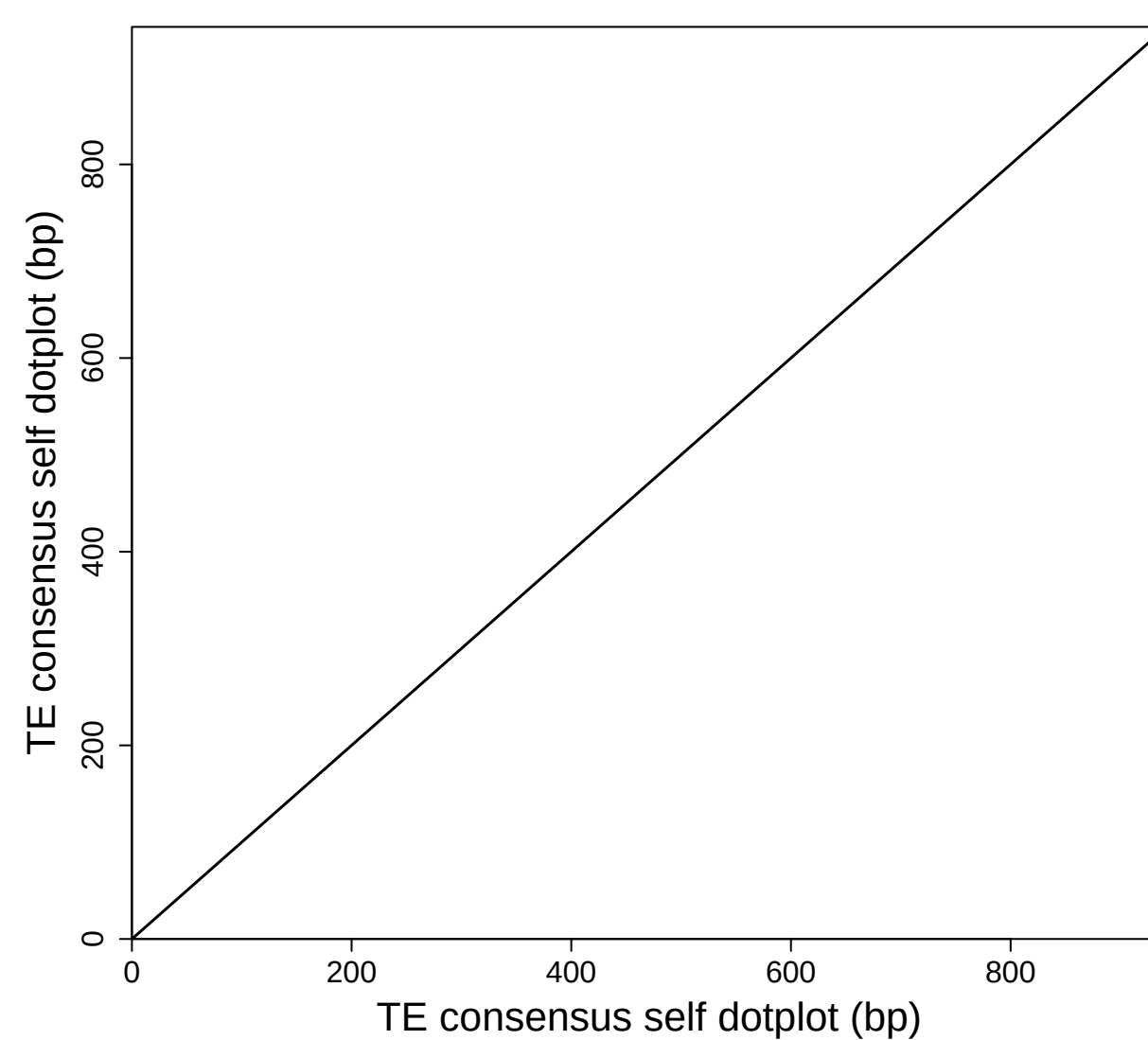
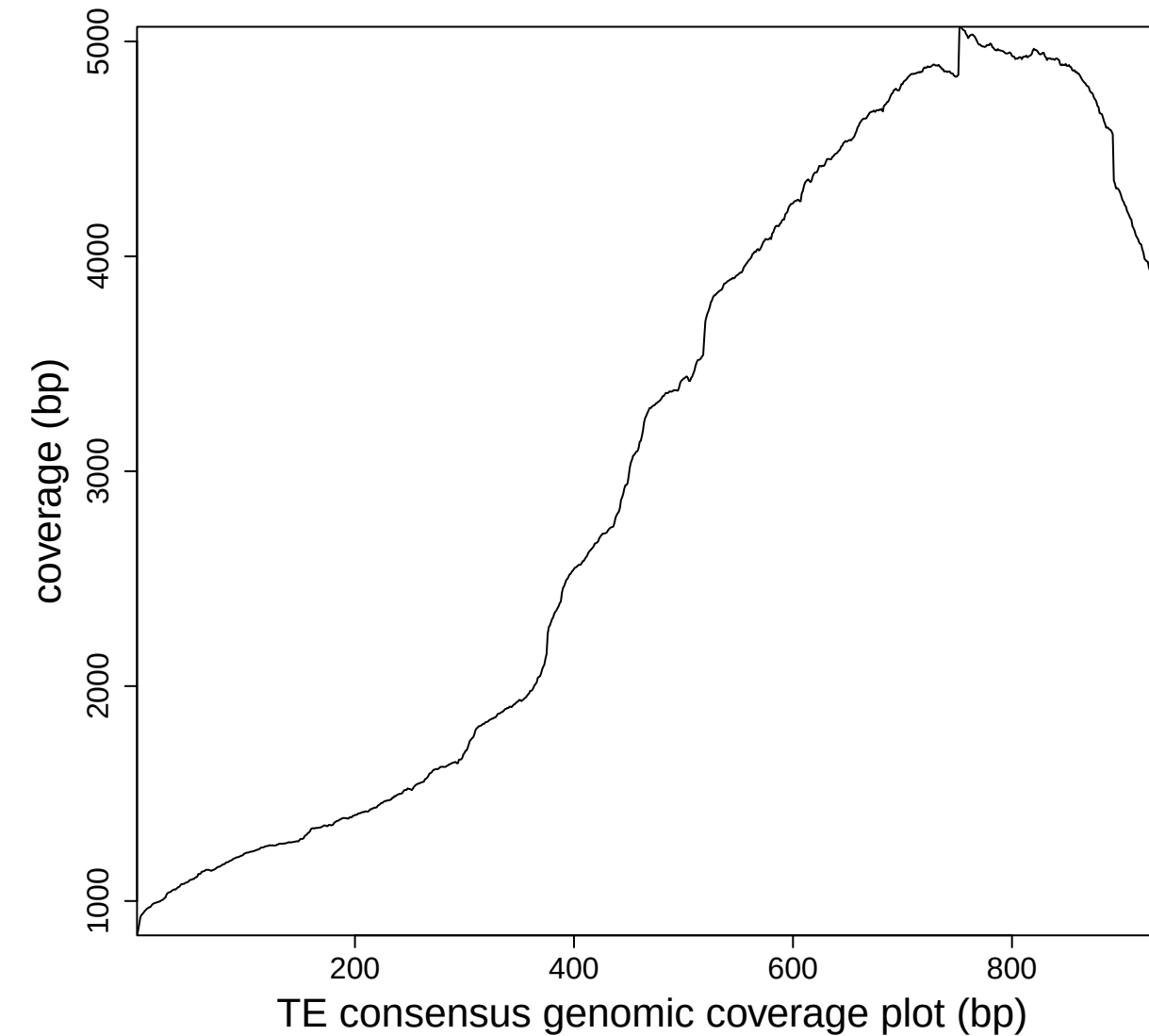
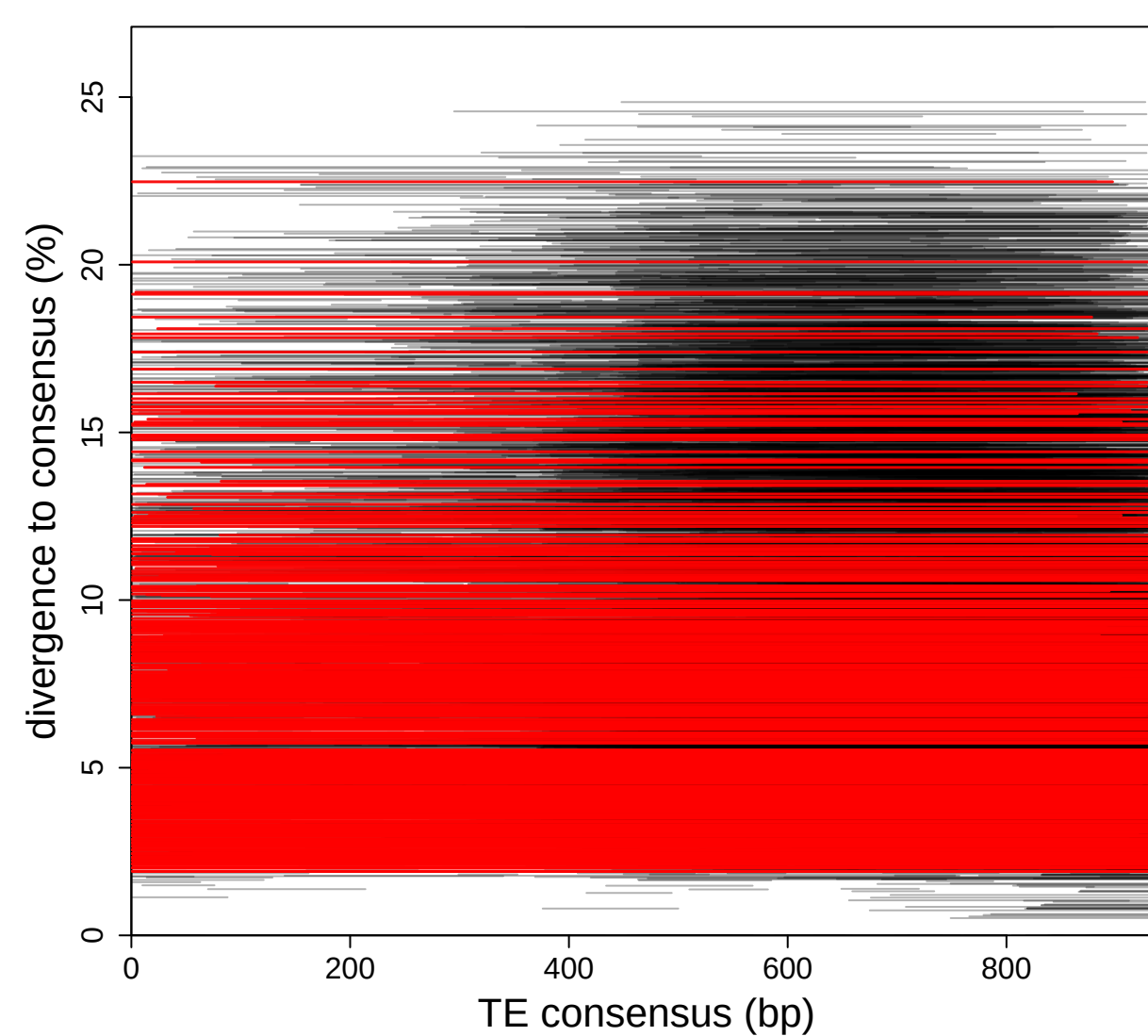
3.fasta.b.bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 10981bp; fragments: 99653; full length: 8 (>=9882.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

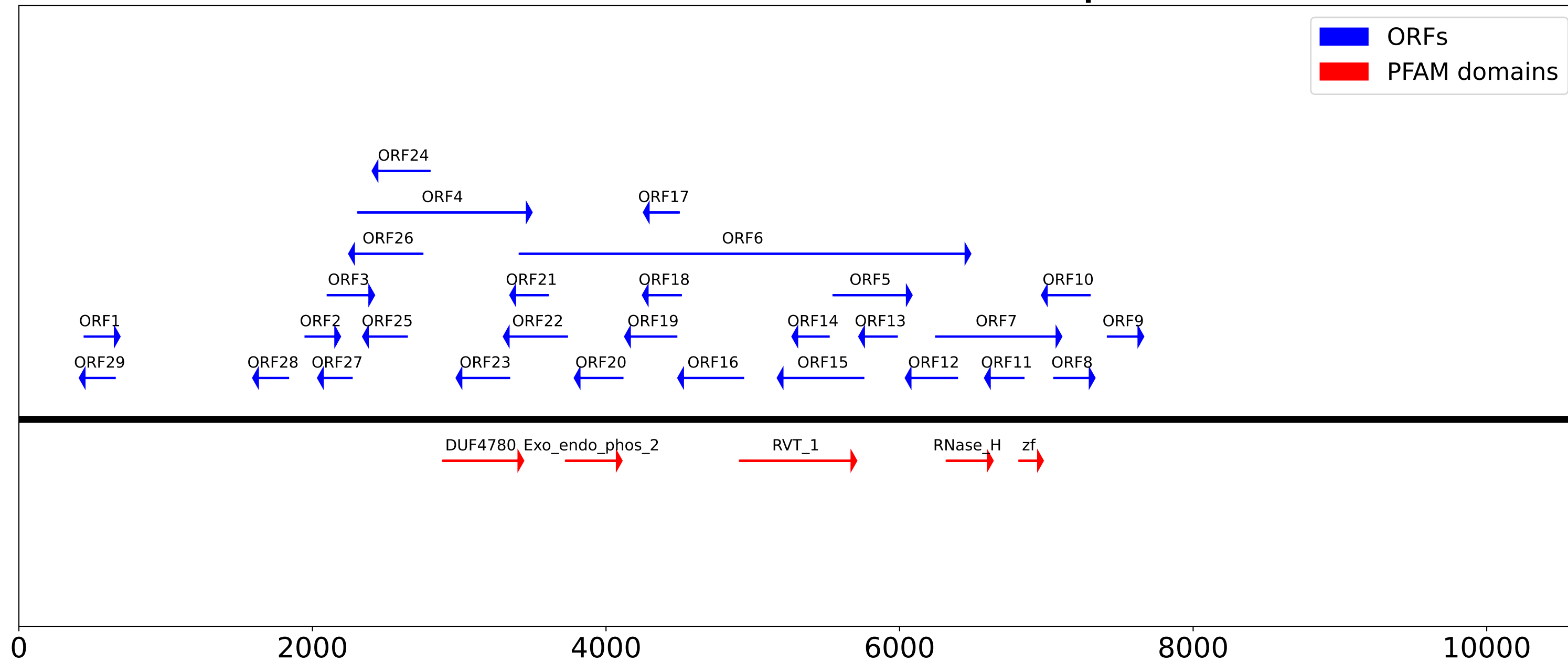


TE: rnd_1_family_383
size: 942bp; fragments: 7208; full length: 683 (≥ 847.8 bp)

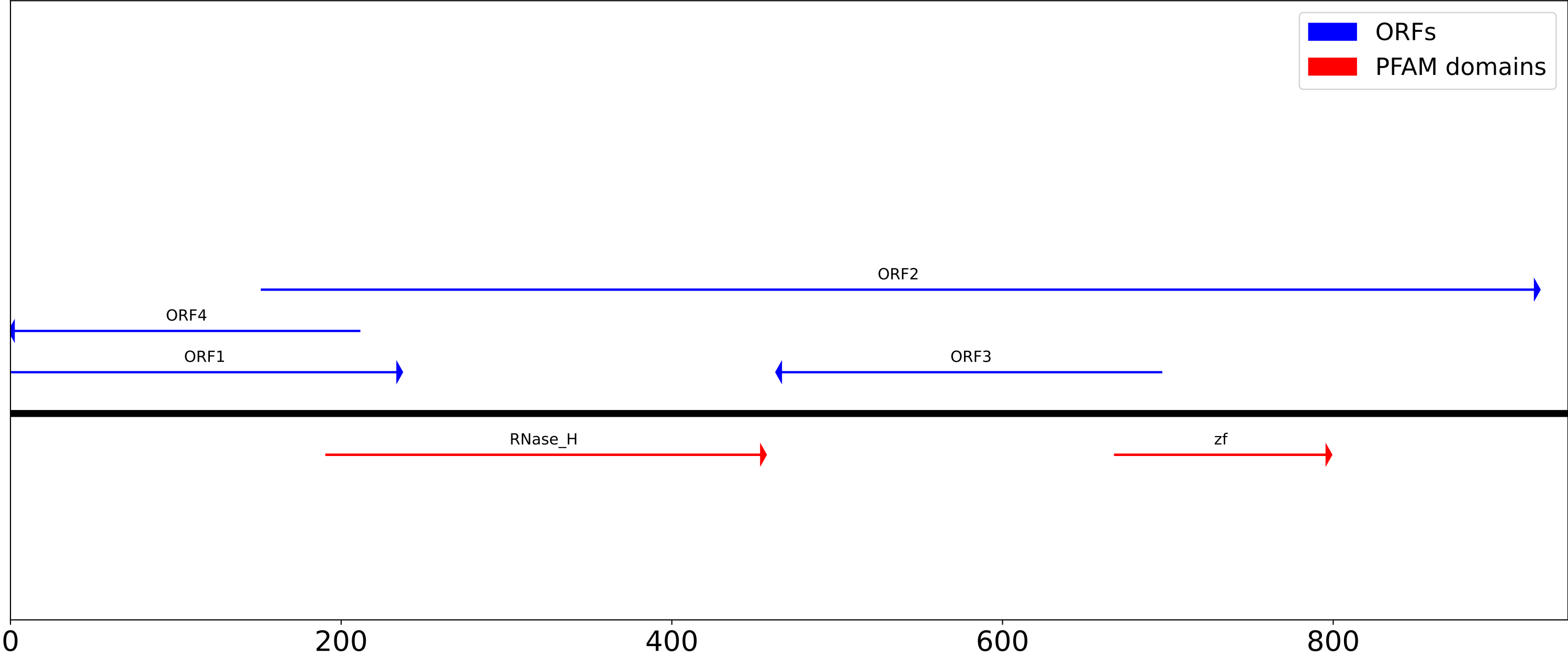
Before TEtrimmer 942 bp



After TEtrimmer ORF and PFAM domain plot



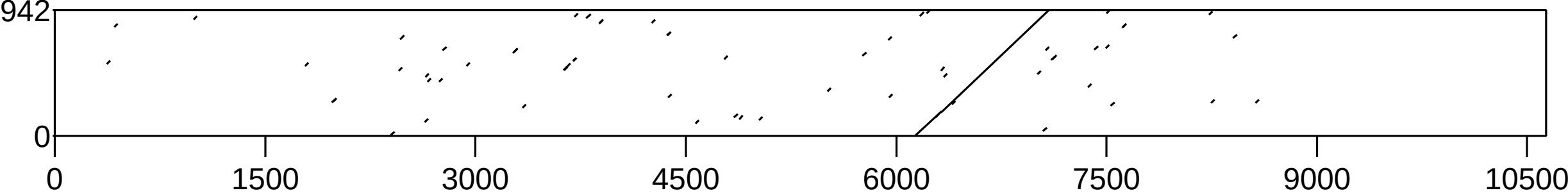
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

ensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)