

Start Crop Point



End Crop Point

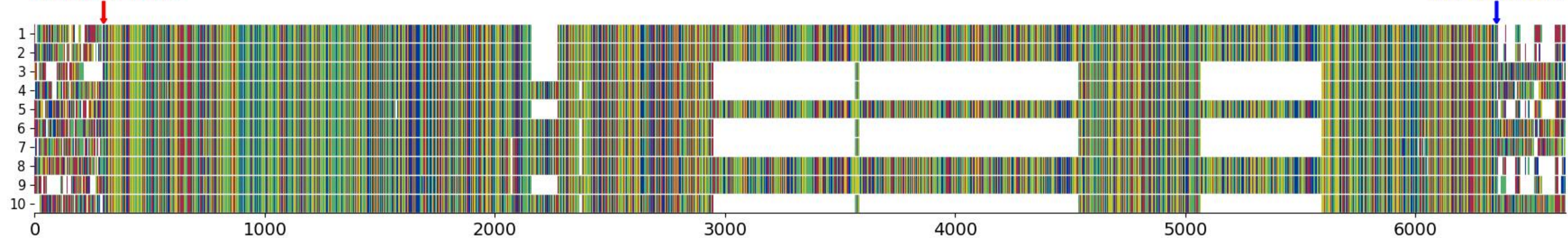


MSA length = 6054



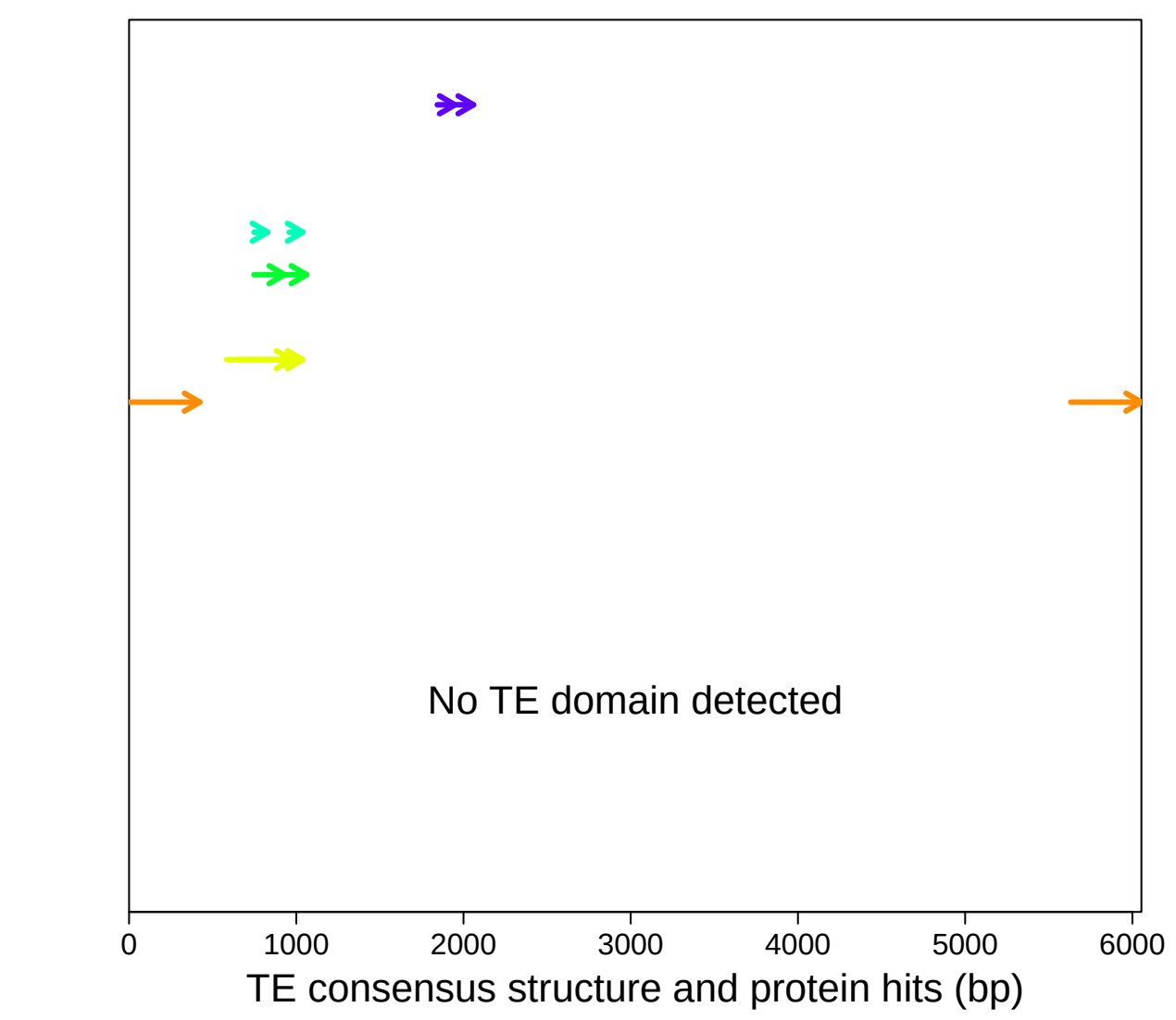
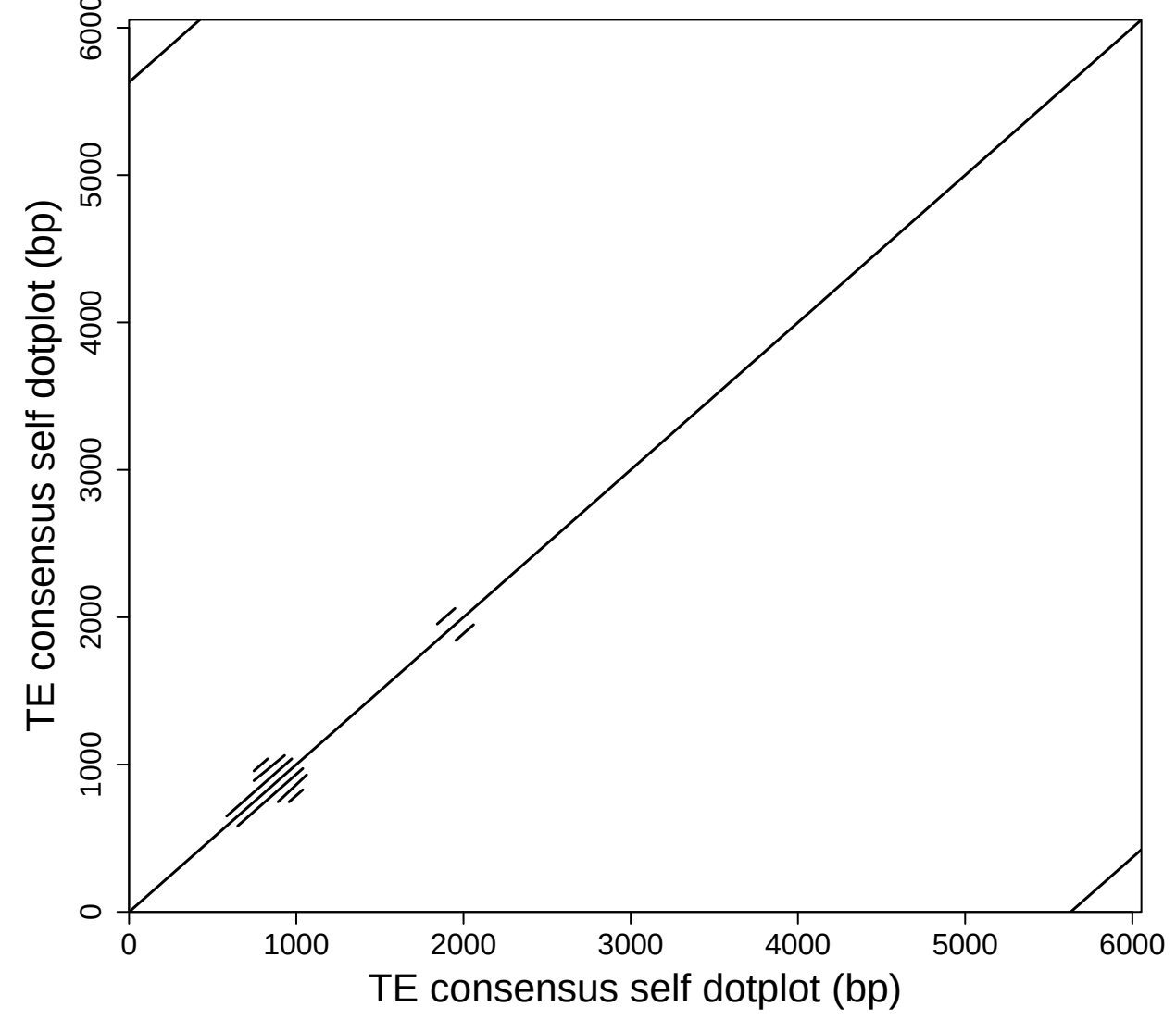
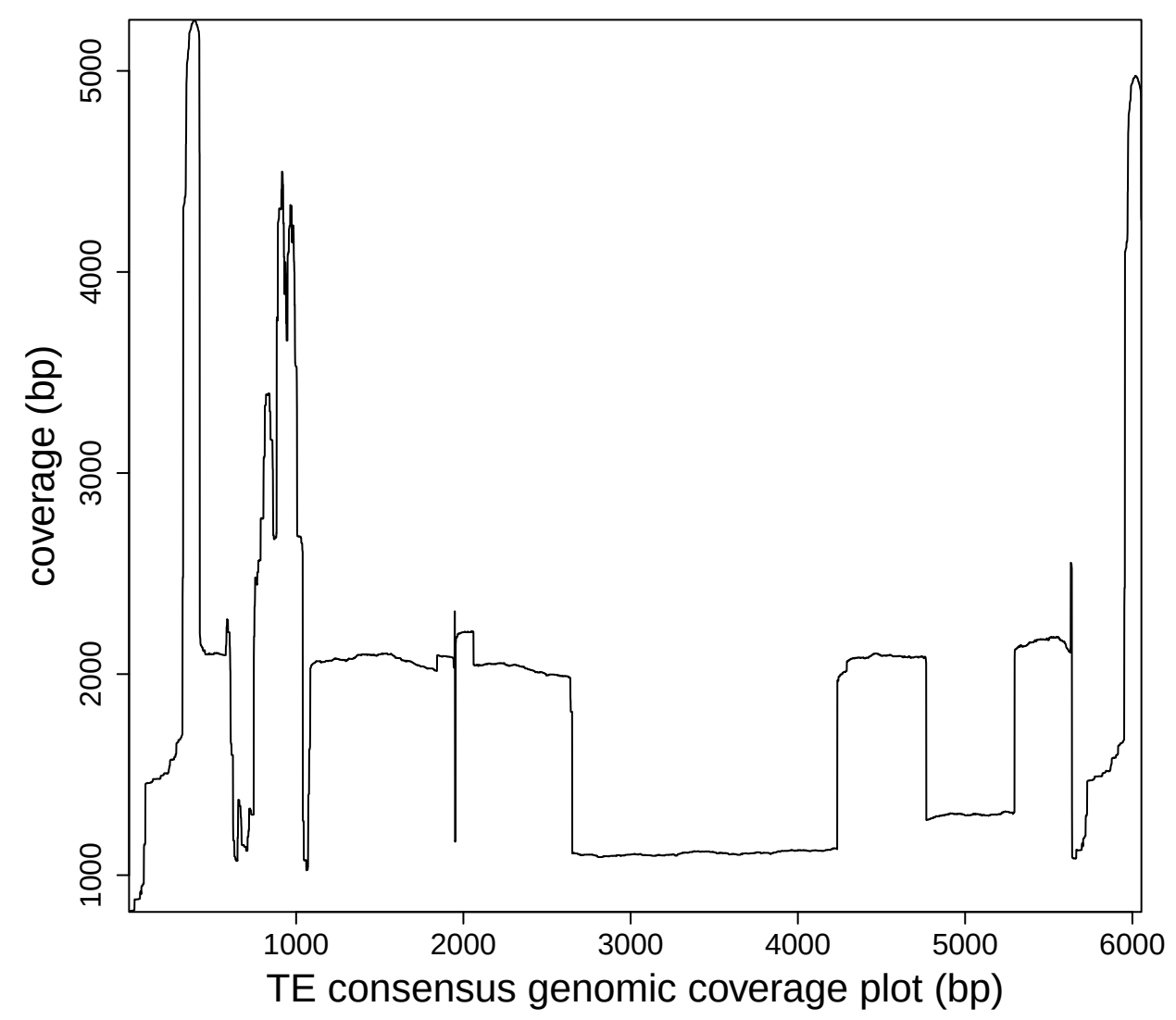
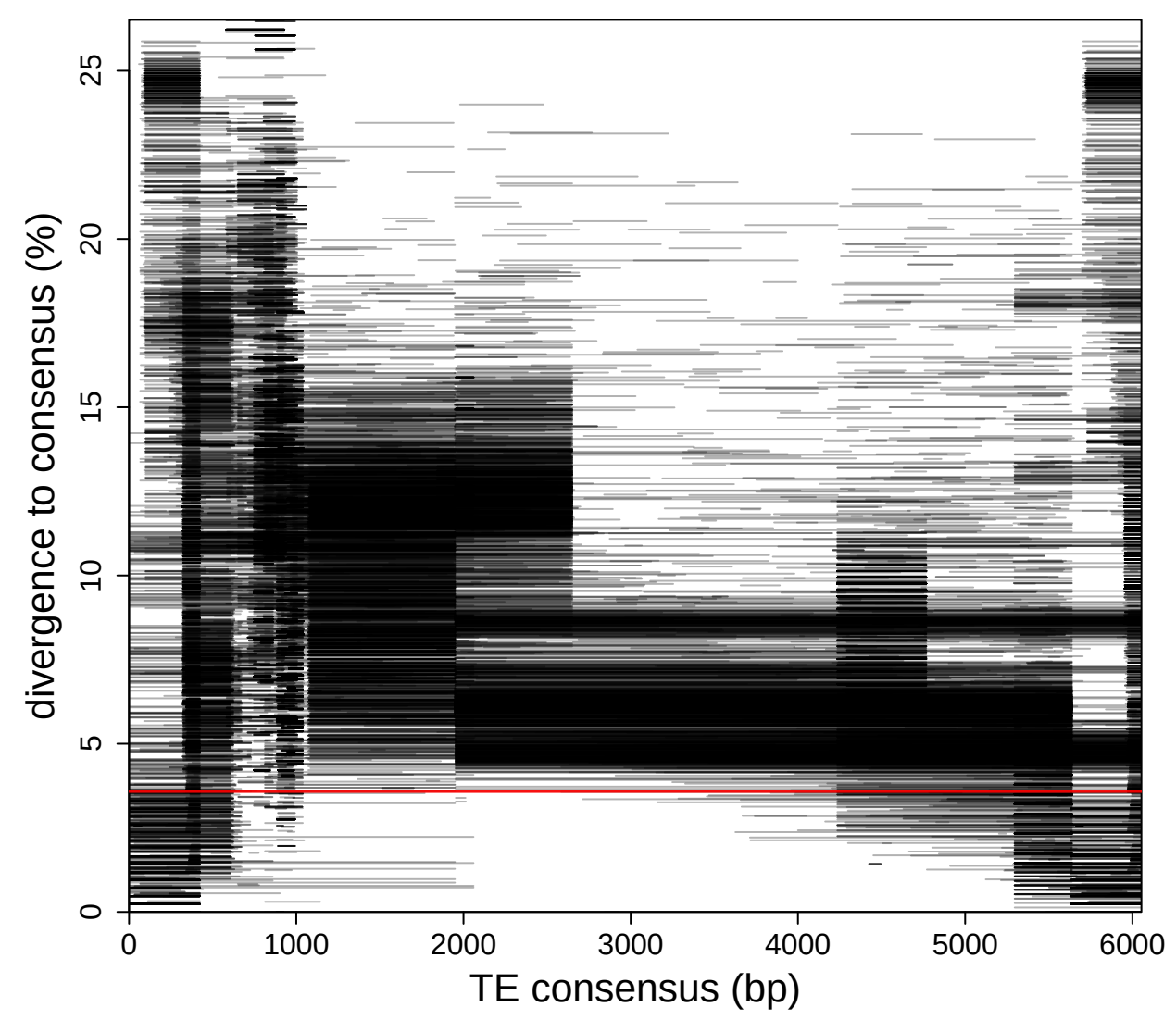
Start crop Point

End crop Point



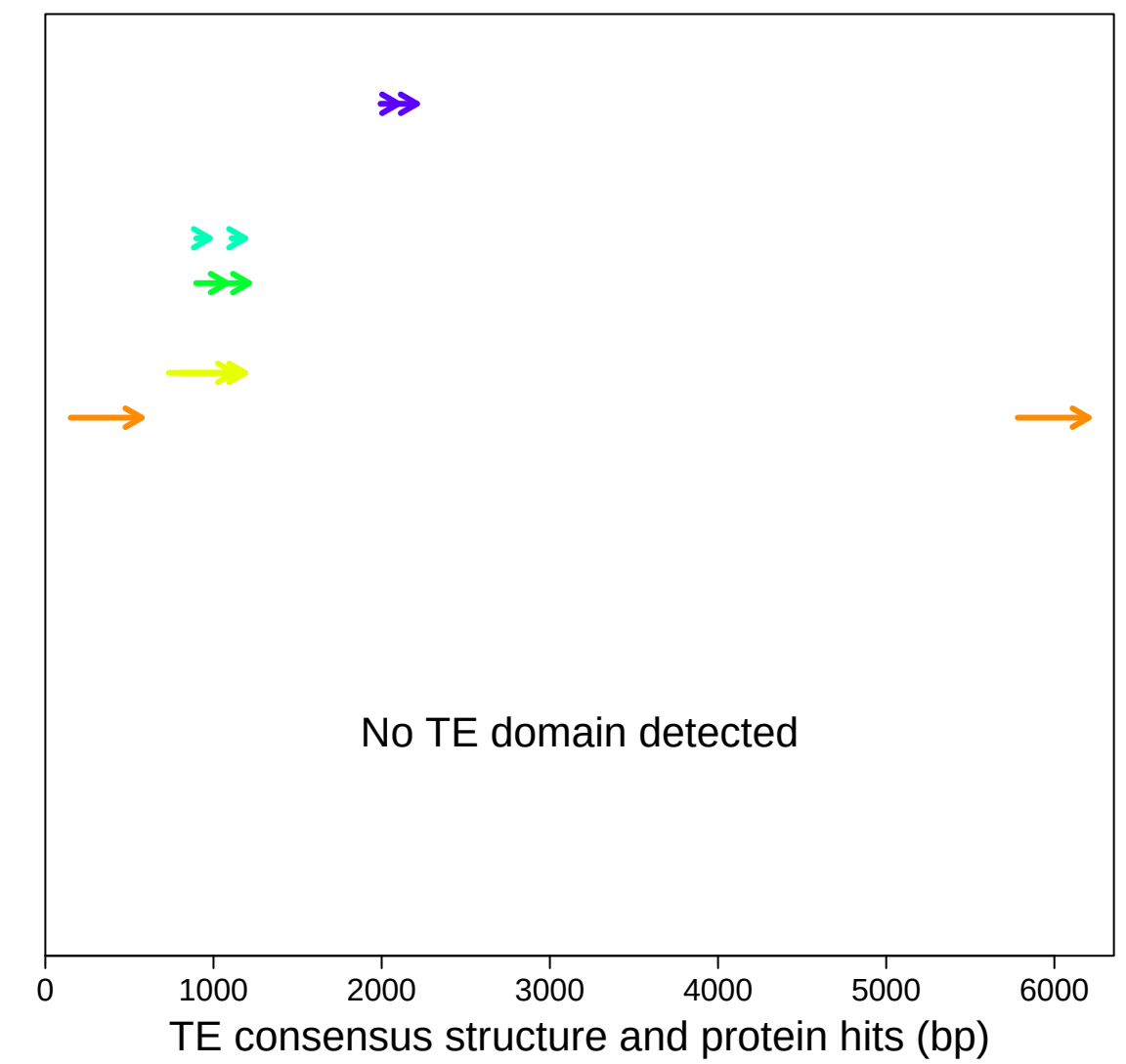
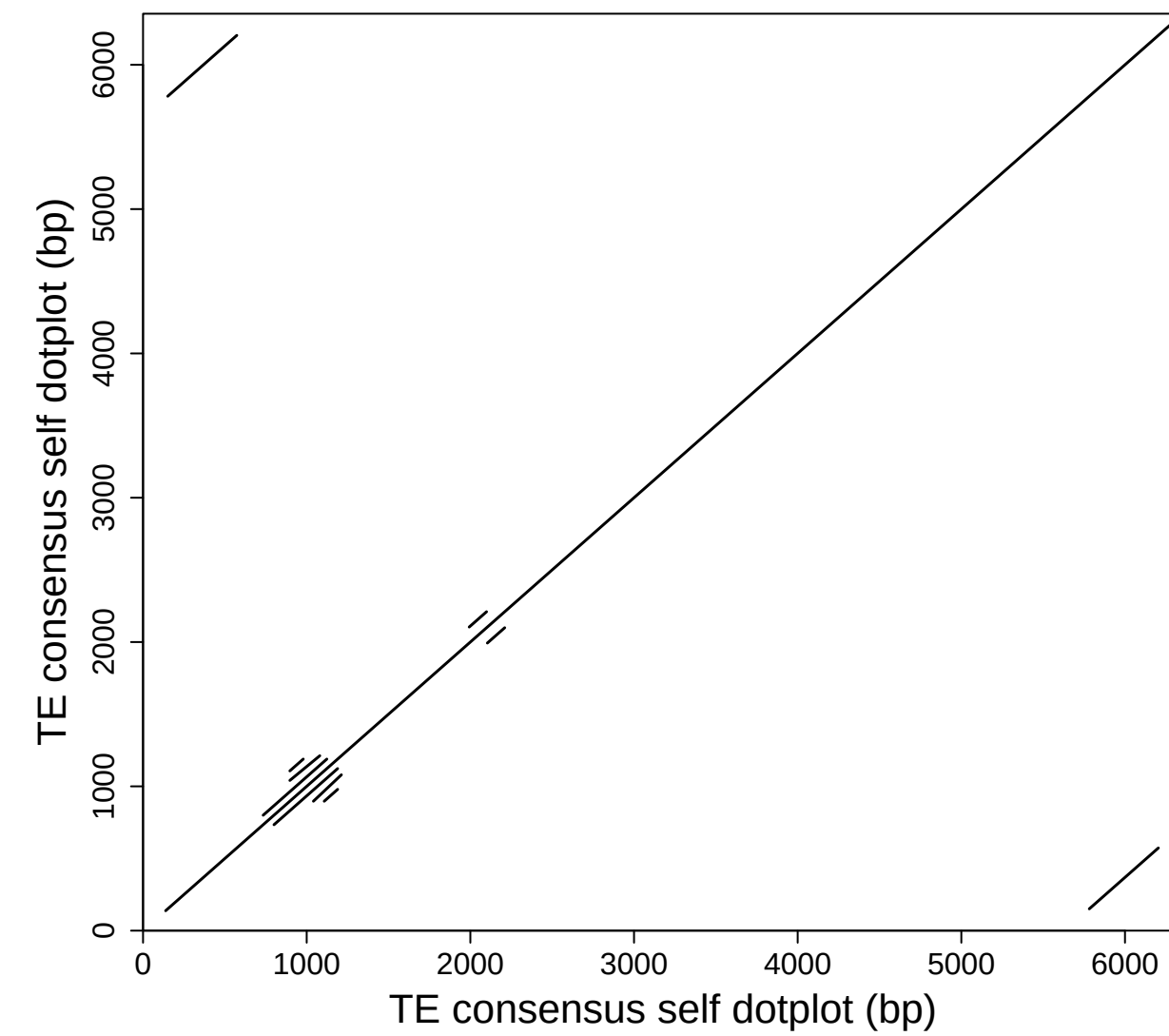
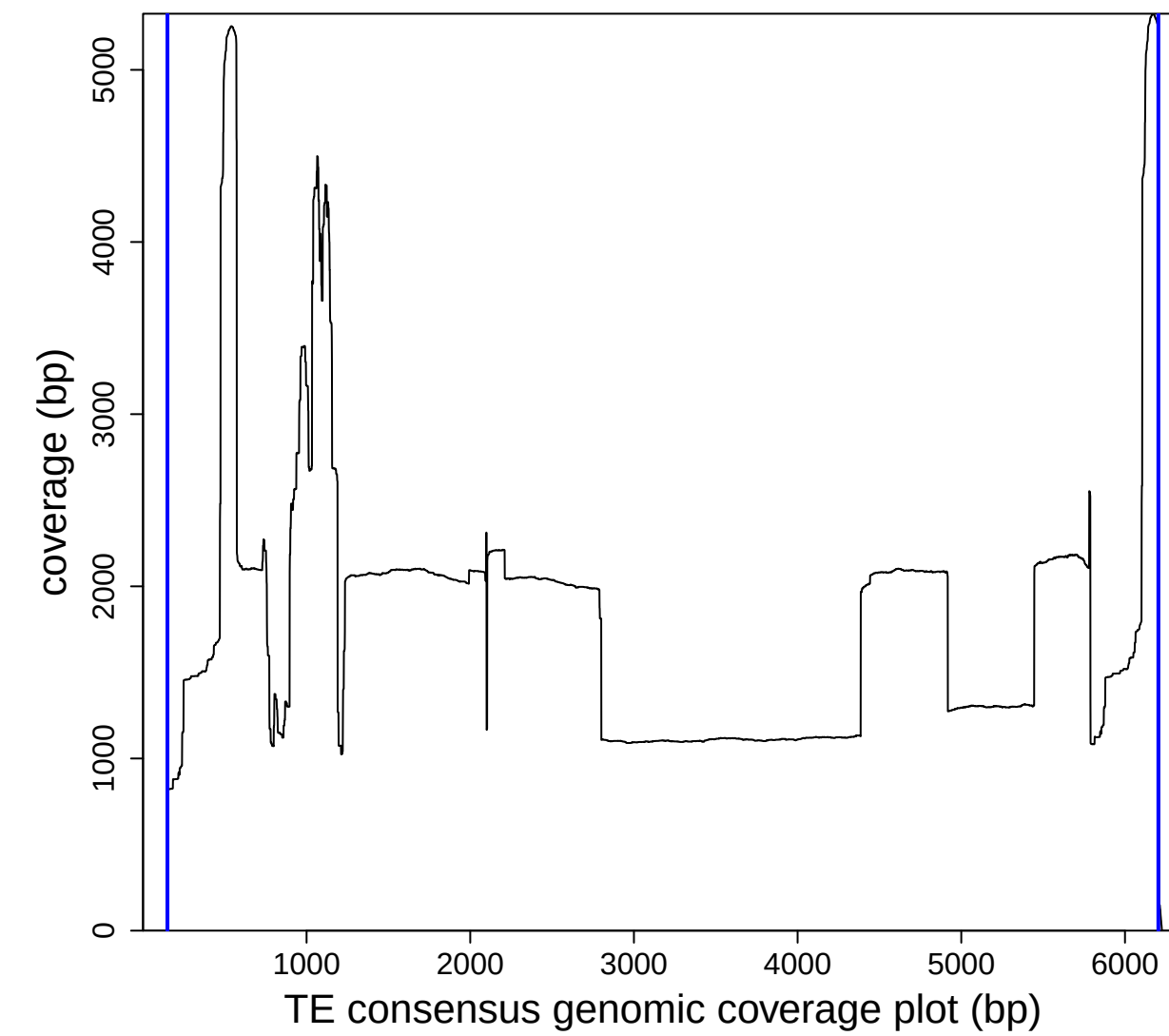
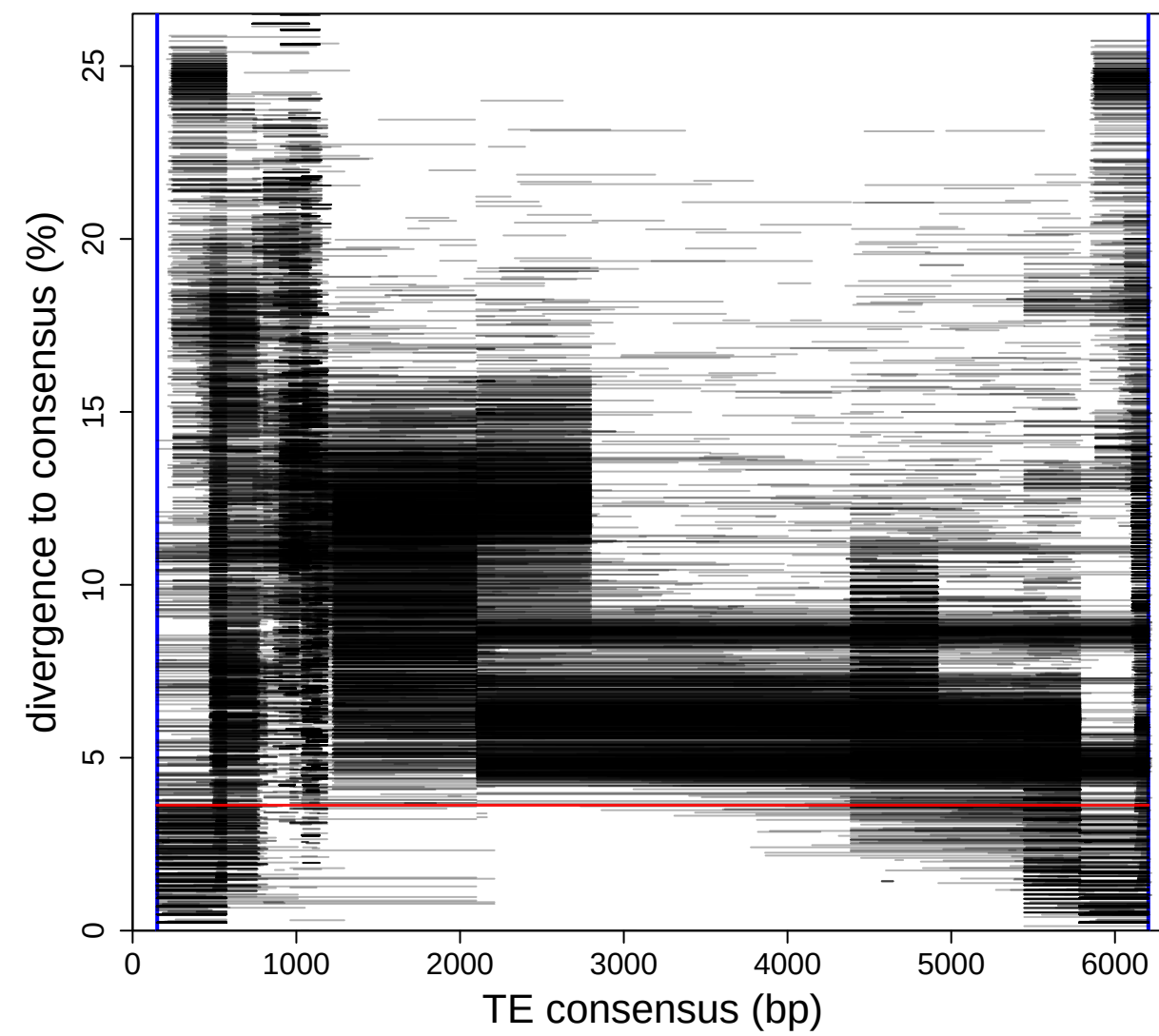
bed_uf.bed g_4.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa c
size: 6054bp; fragments: 20671; full length: 1 (>=5448.6bp)

After TETrimmer 6054 bp



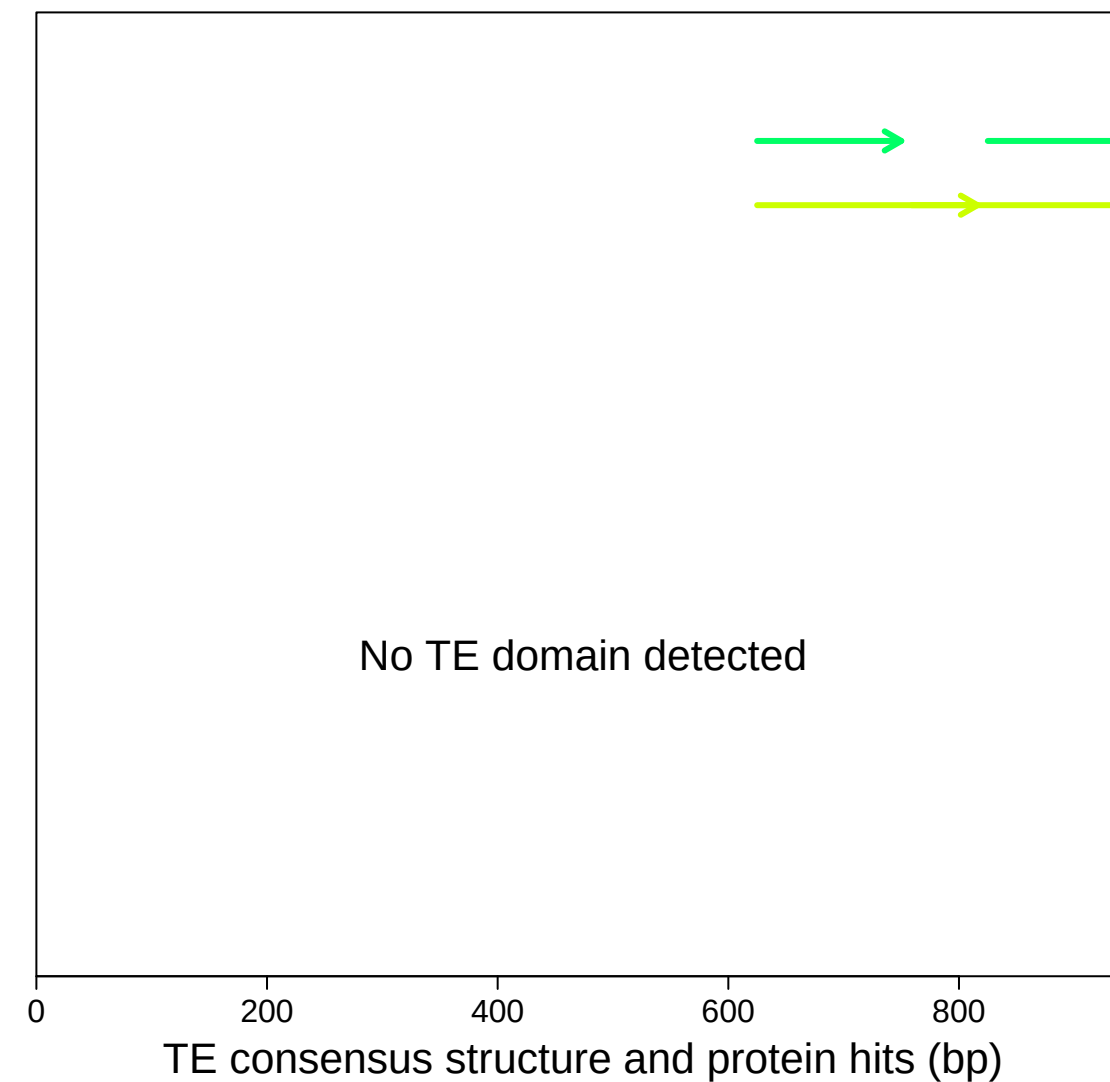
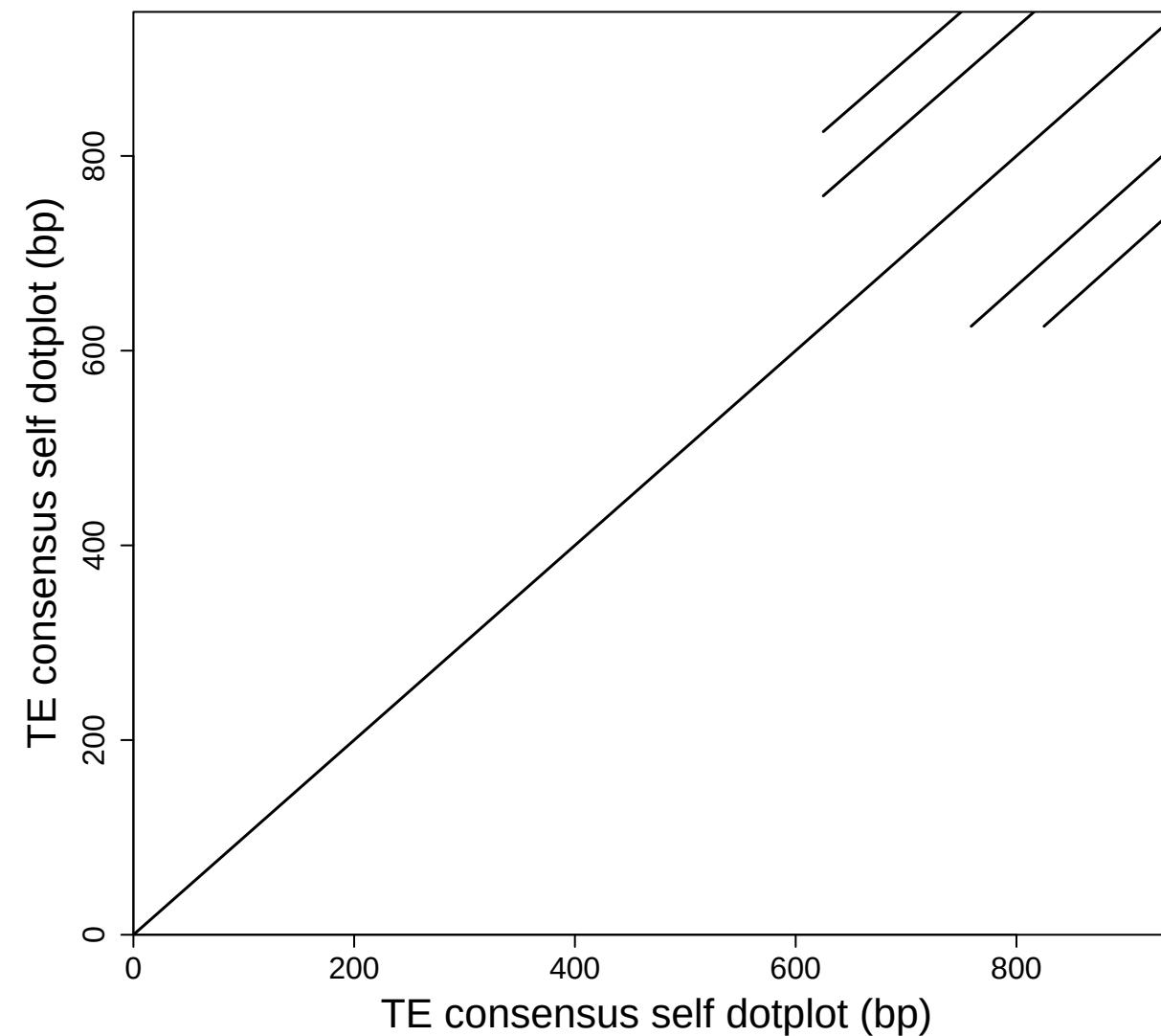
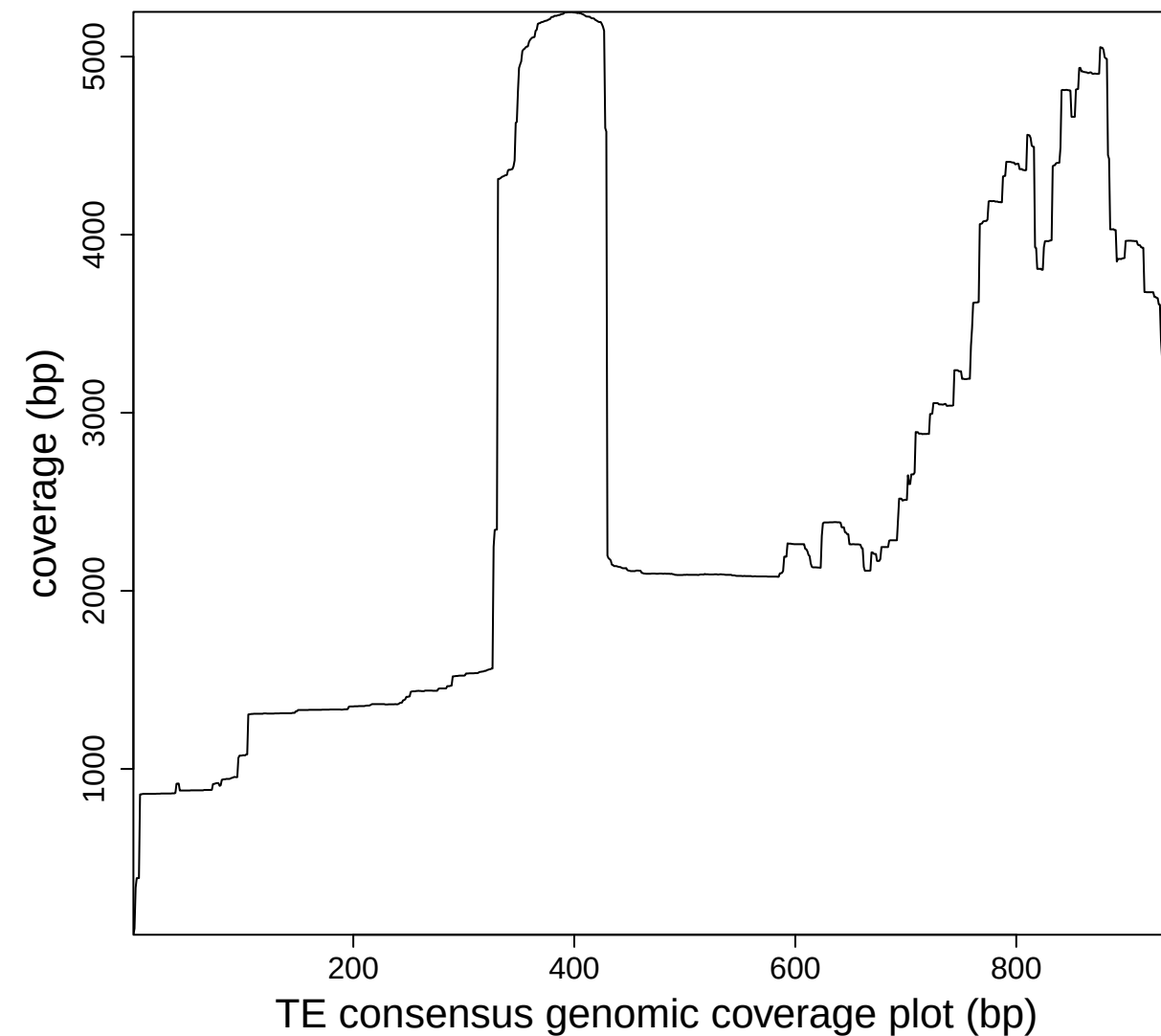
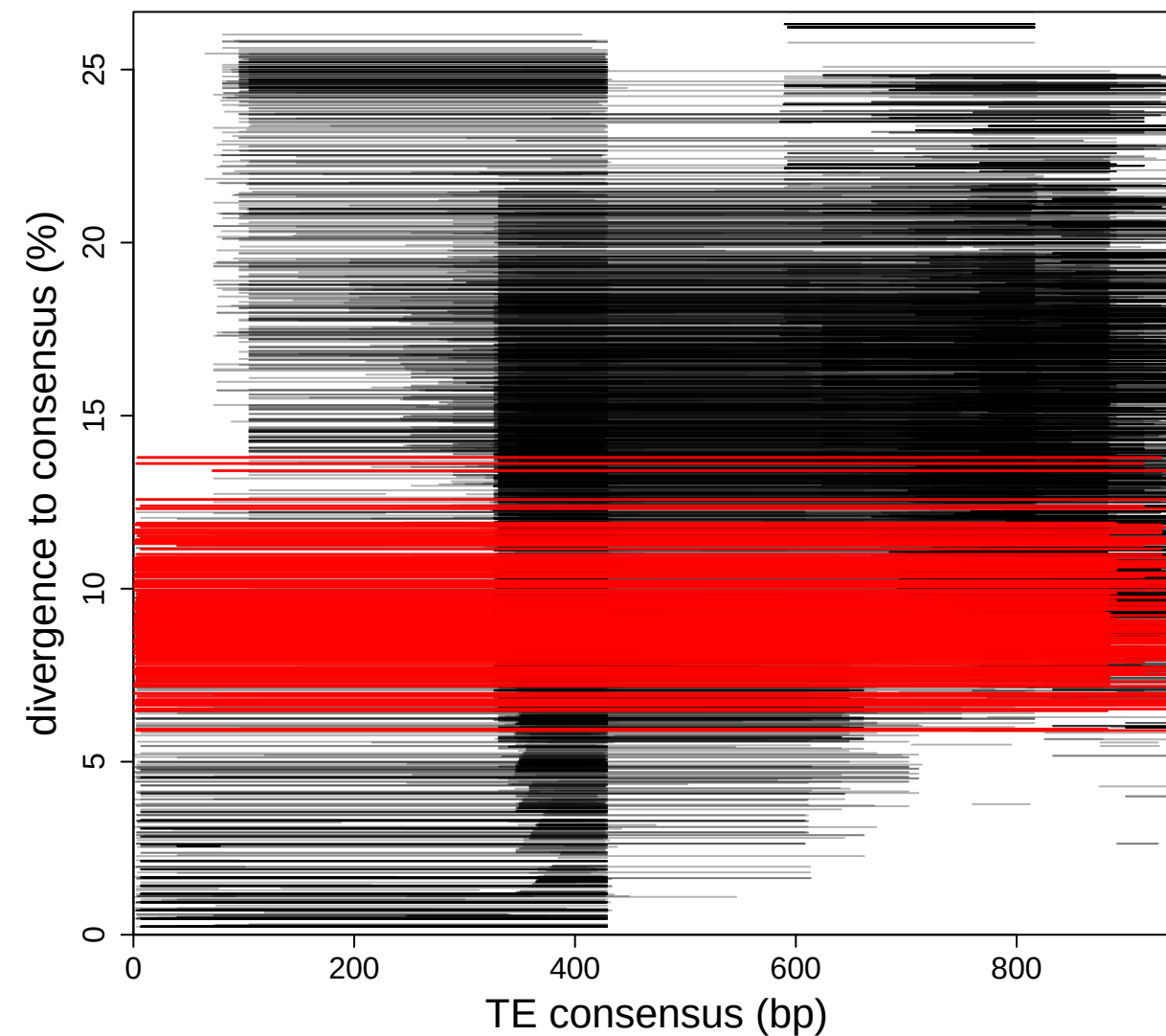
ly_1353.fasta.b.bed uf.bed g 4.bed fm 1.bed 0 0 bcln.fa aln.f
size: 6354bp; fragments: 21022; full length: 1 (>=5718.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

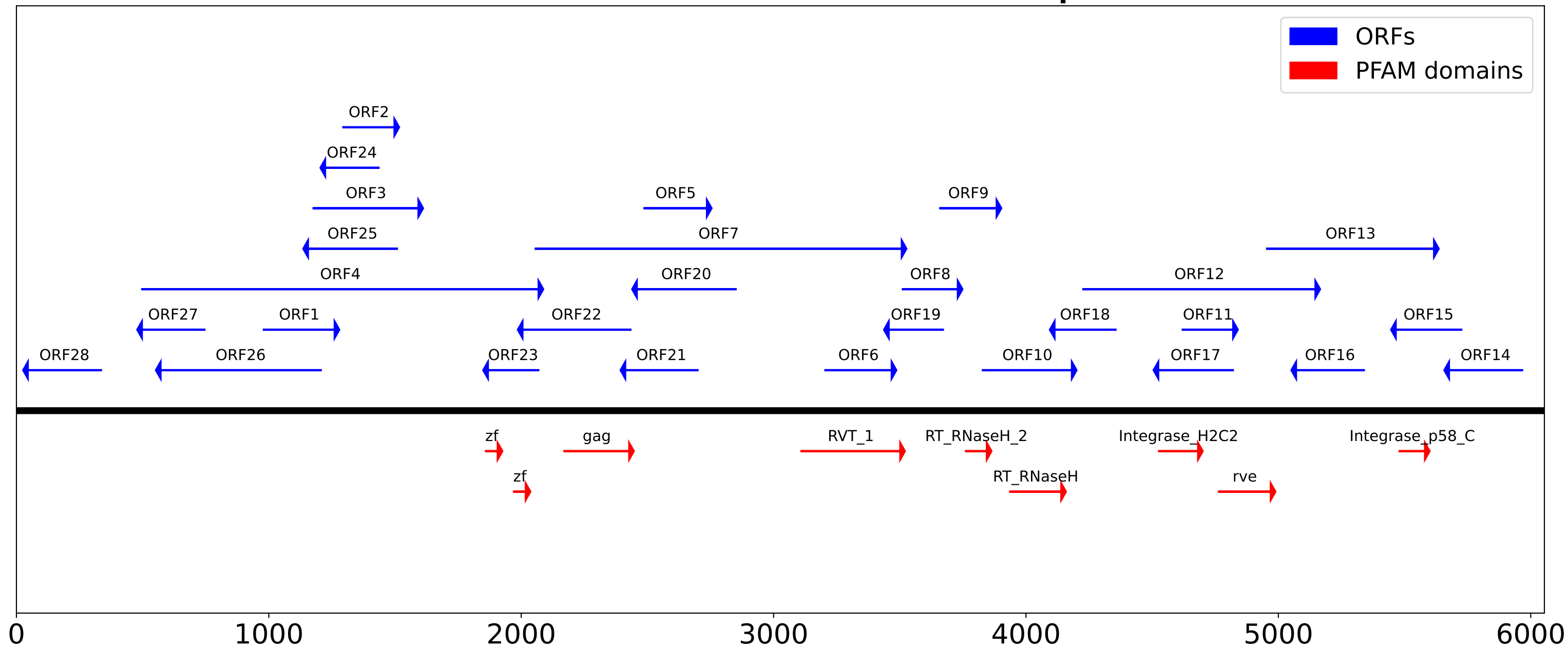


TE: rnd_5_family_1353
size: 948bp; fragments: 10200; full length: 222 (≥ 853.2 bp)

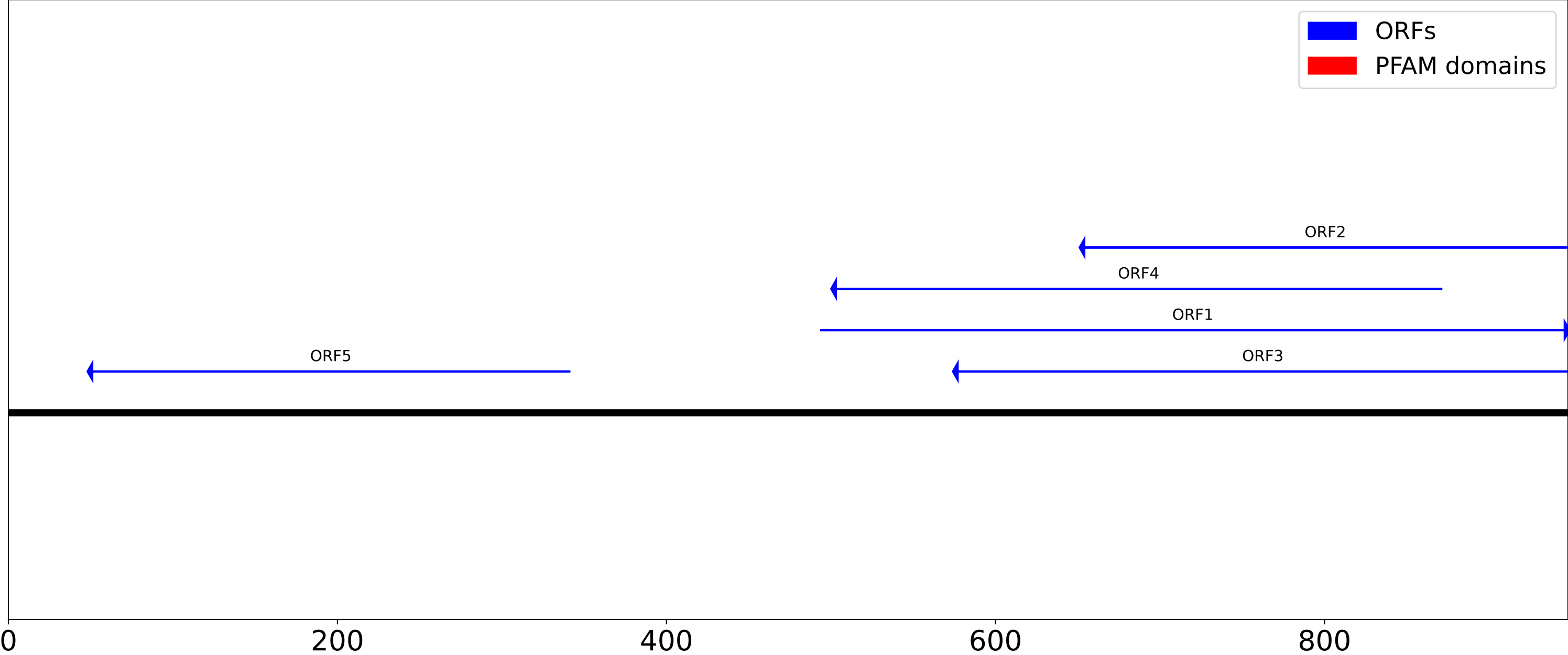
Before TEtrimmer 948 bp



After TEtrimmer ORF and PFAM domain plot



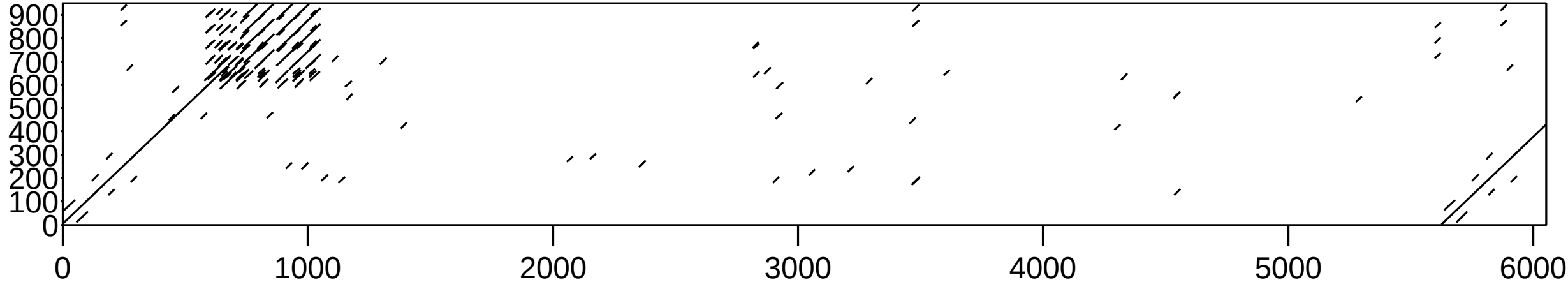
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

1sensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)