

Start Crop Point



End Crop Point

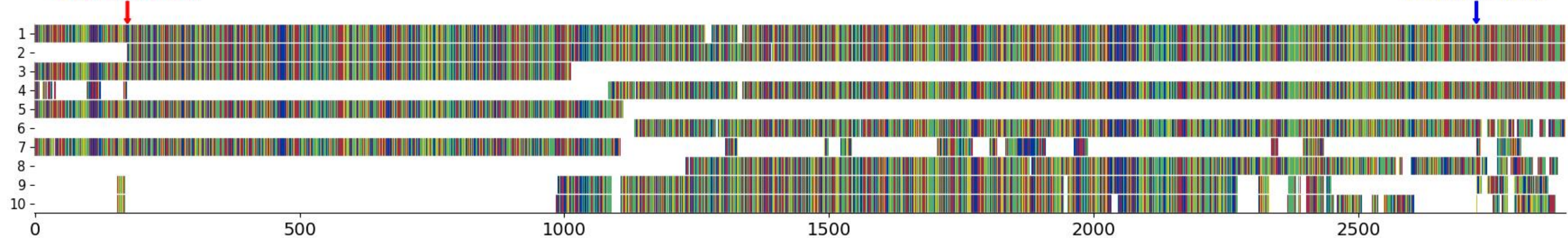


MSA length = 2549



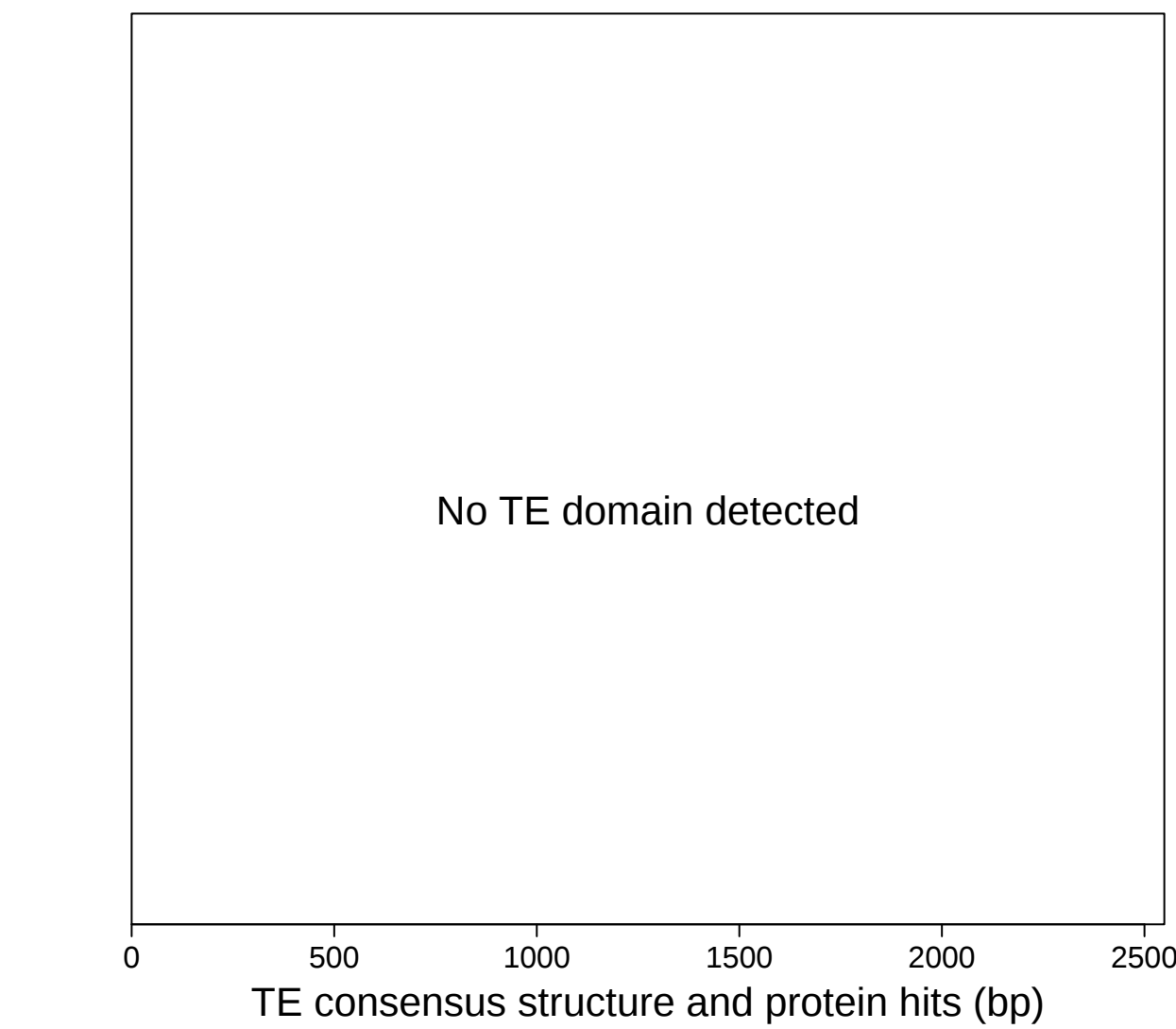
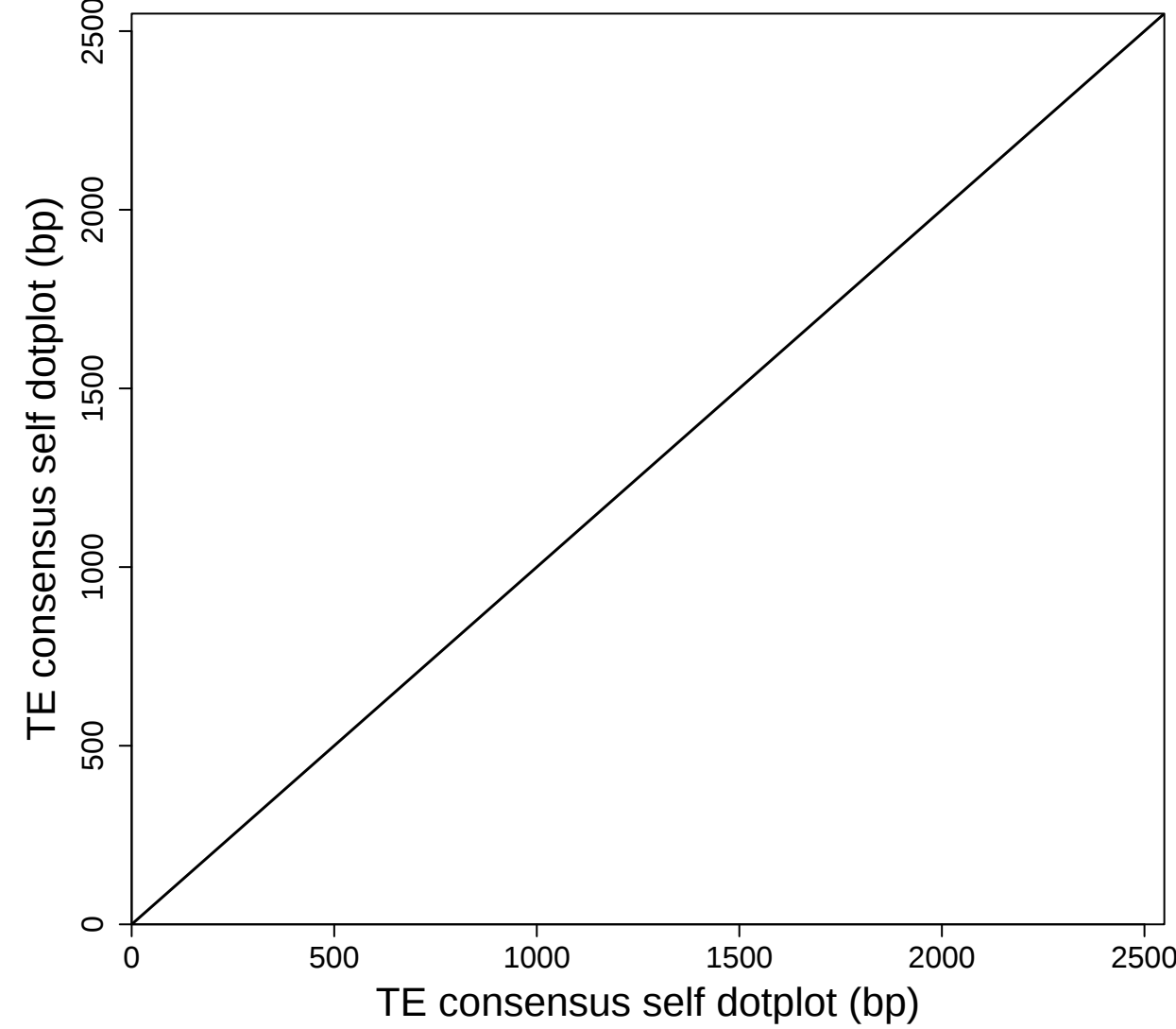
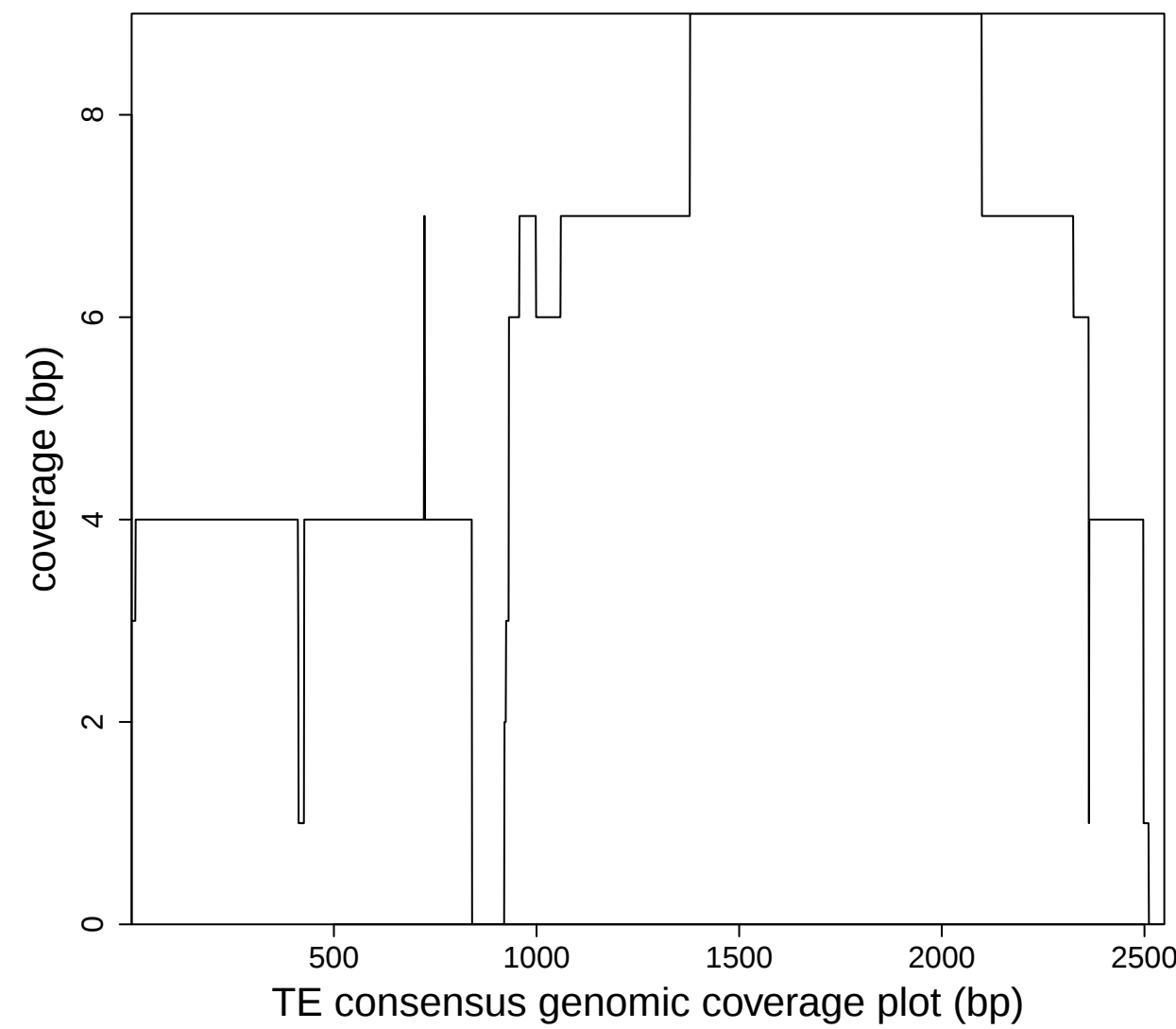
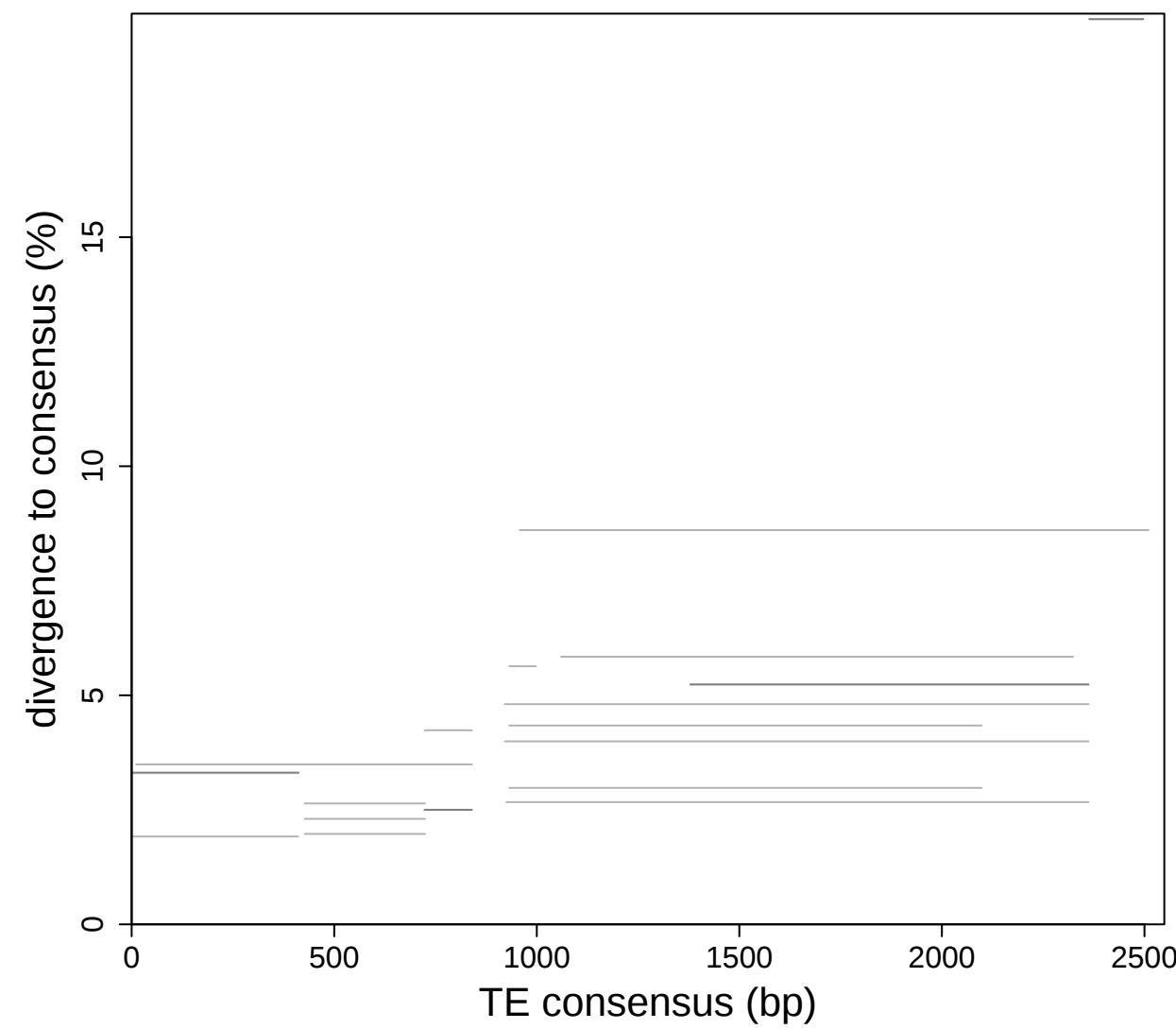
Start crop Point

End crop Point



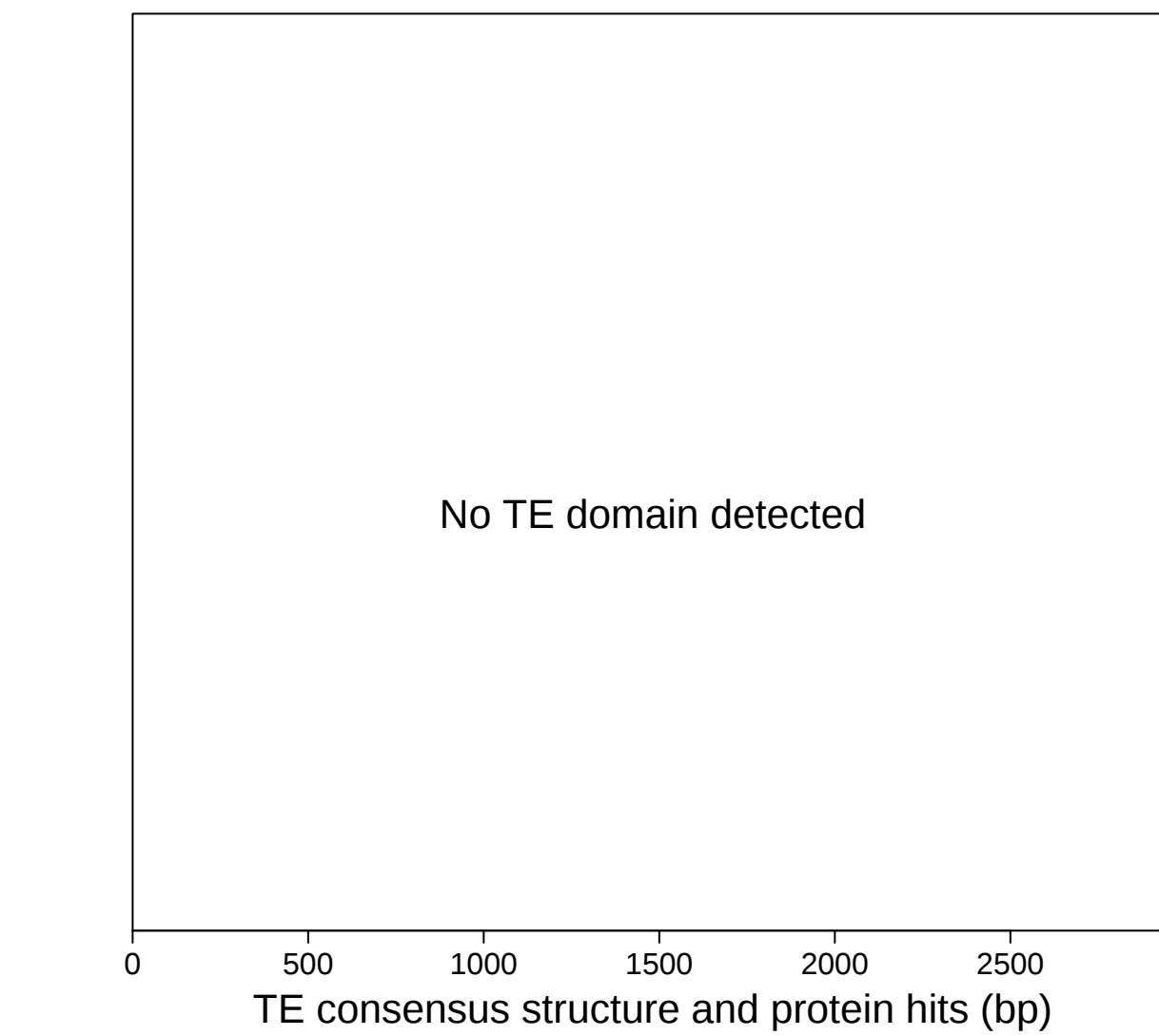
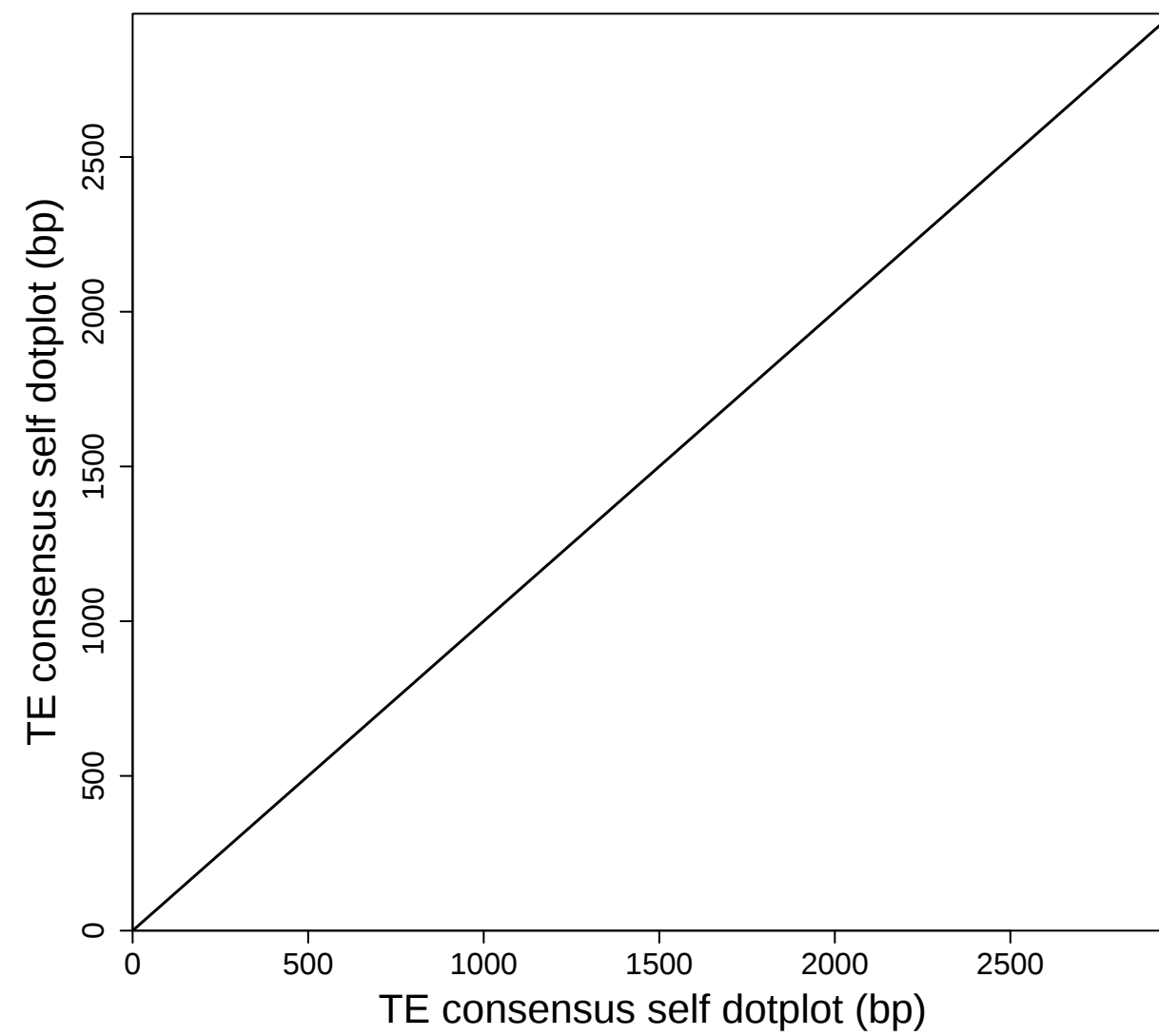
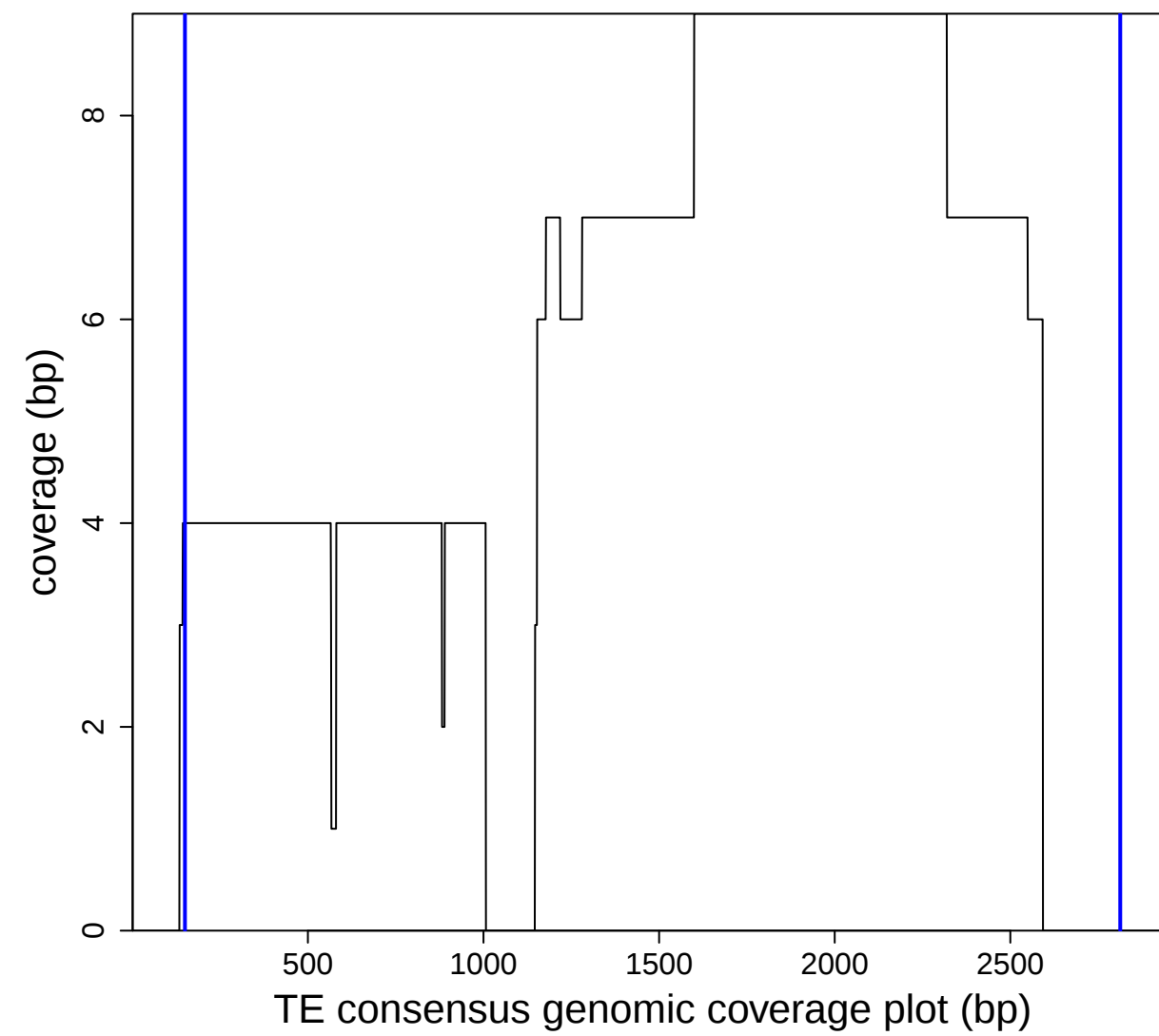
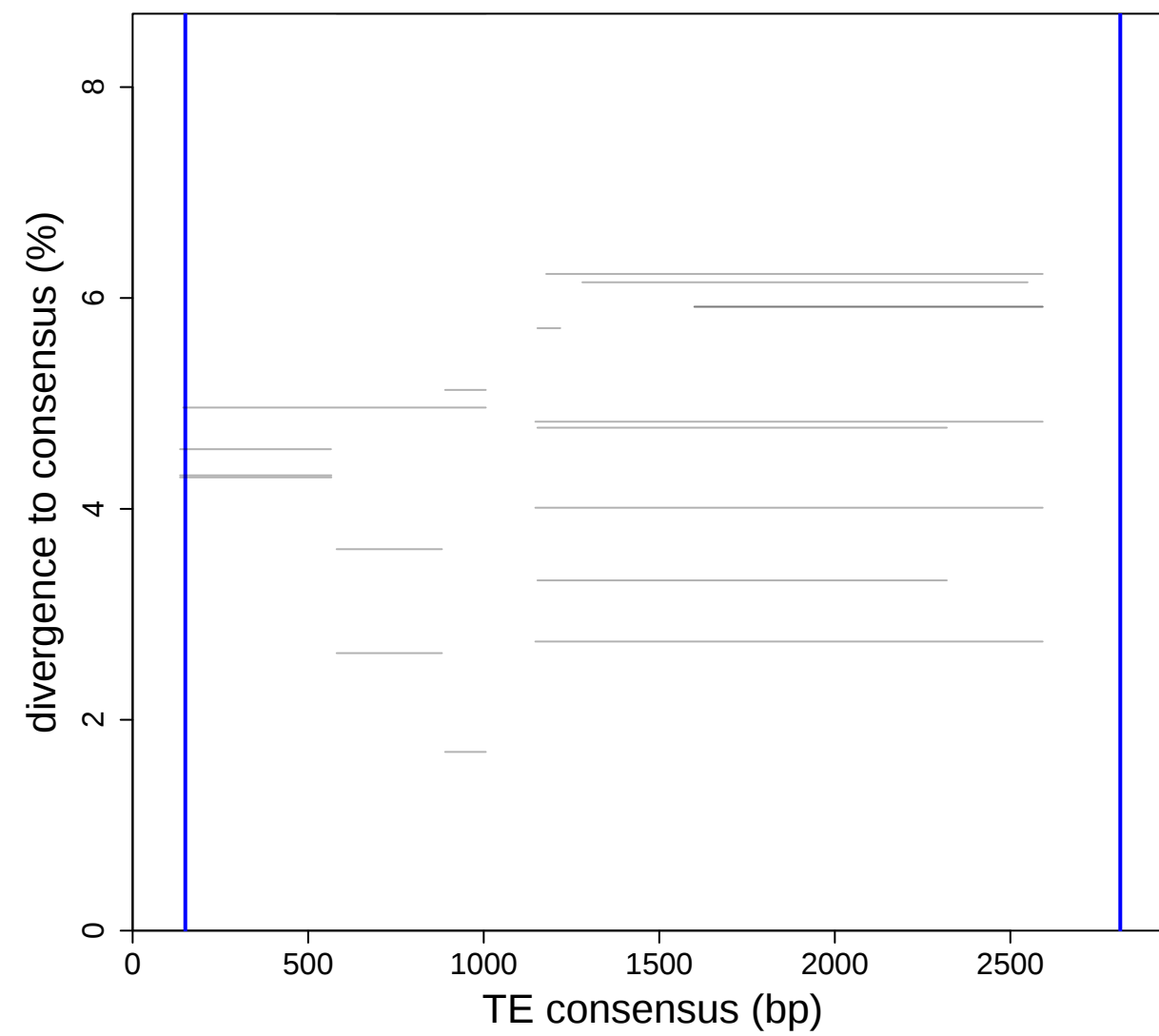
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 2549bp; fragments: 23; full length: 0 (>=2294.1bp)

After TETrimmer 2549 bp



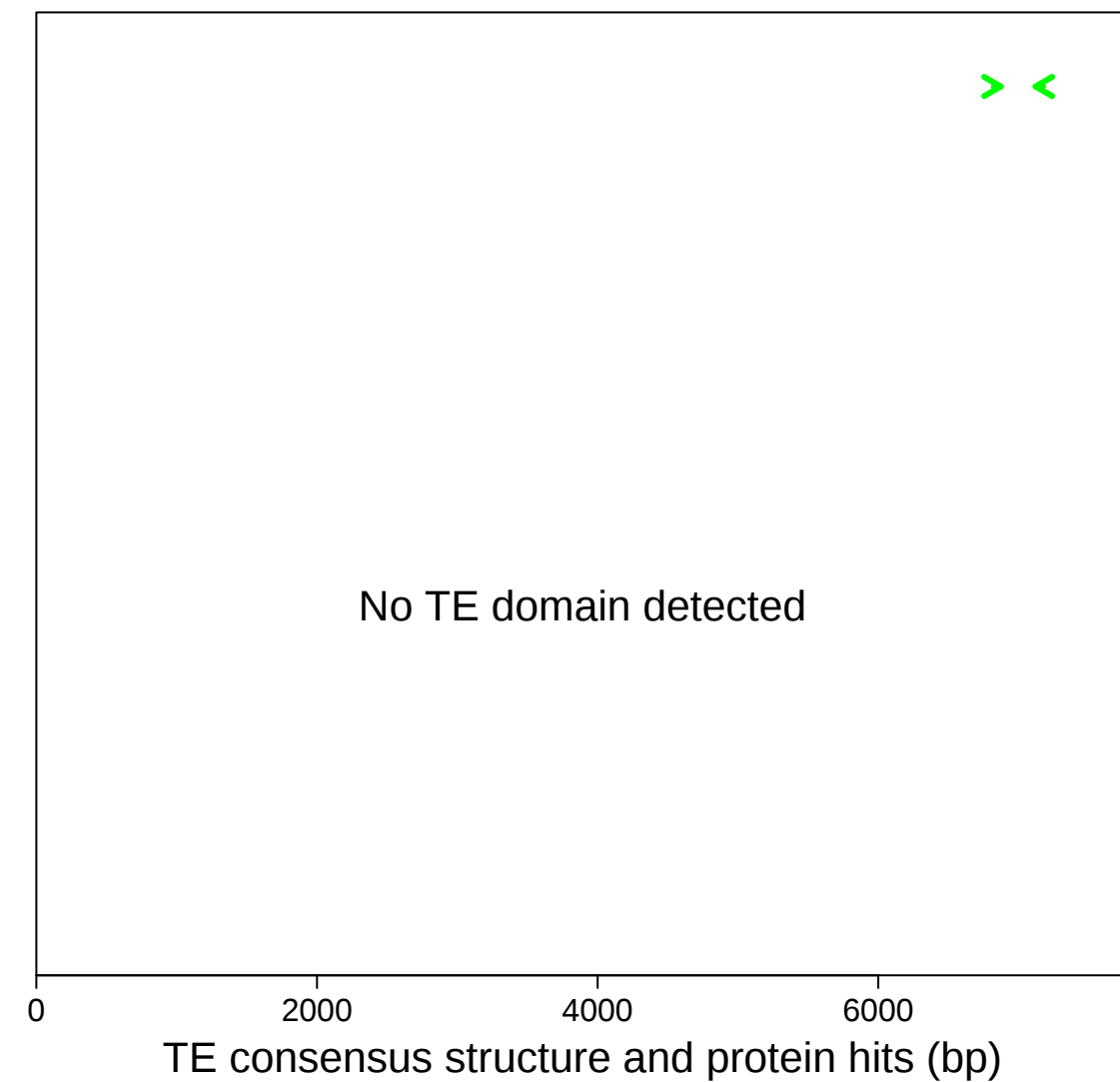
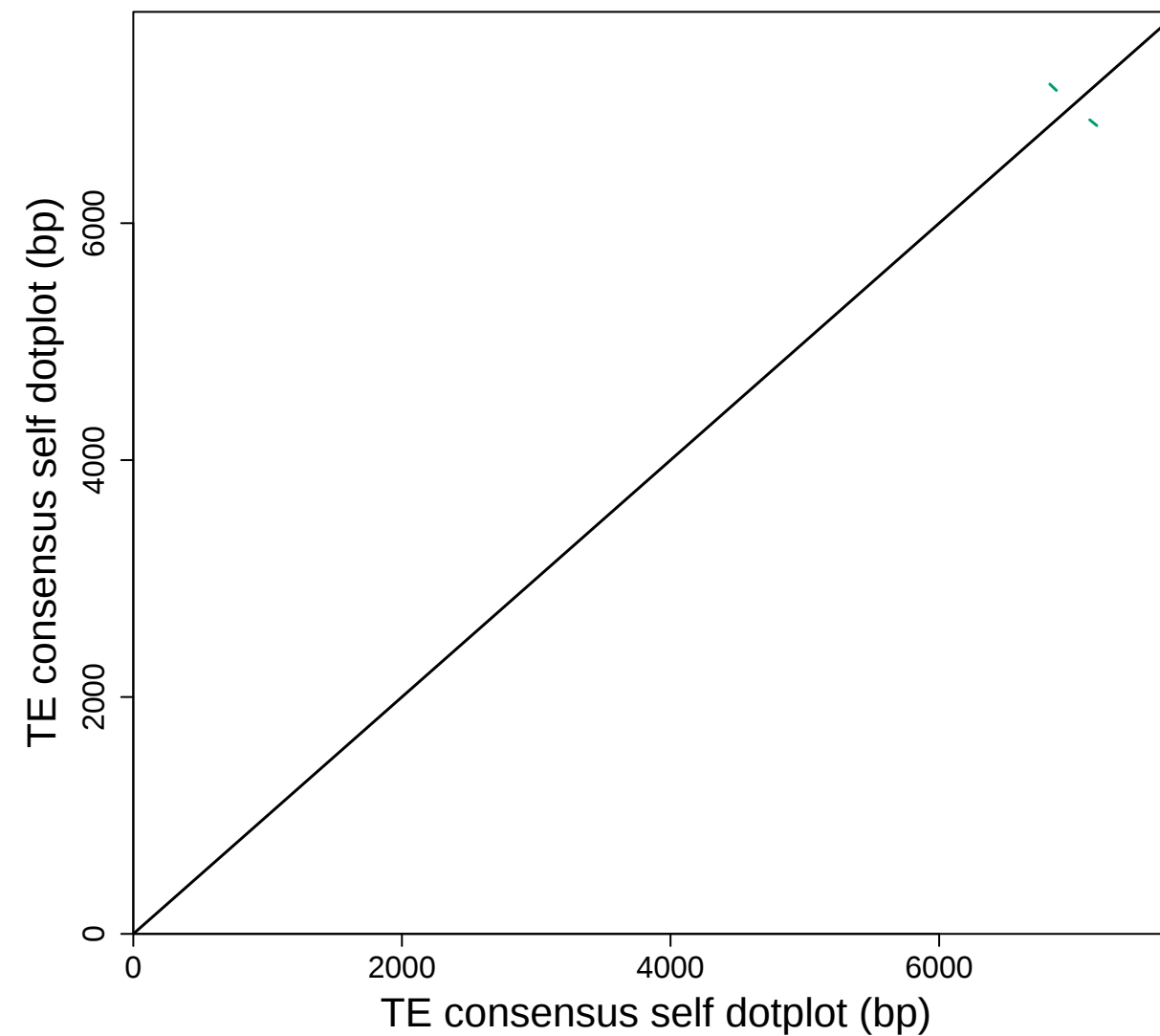
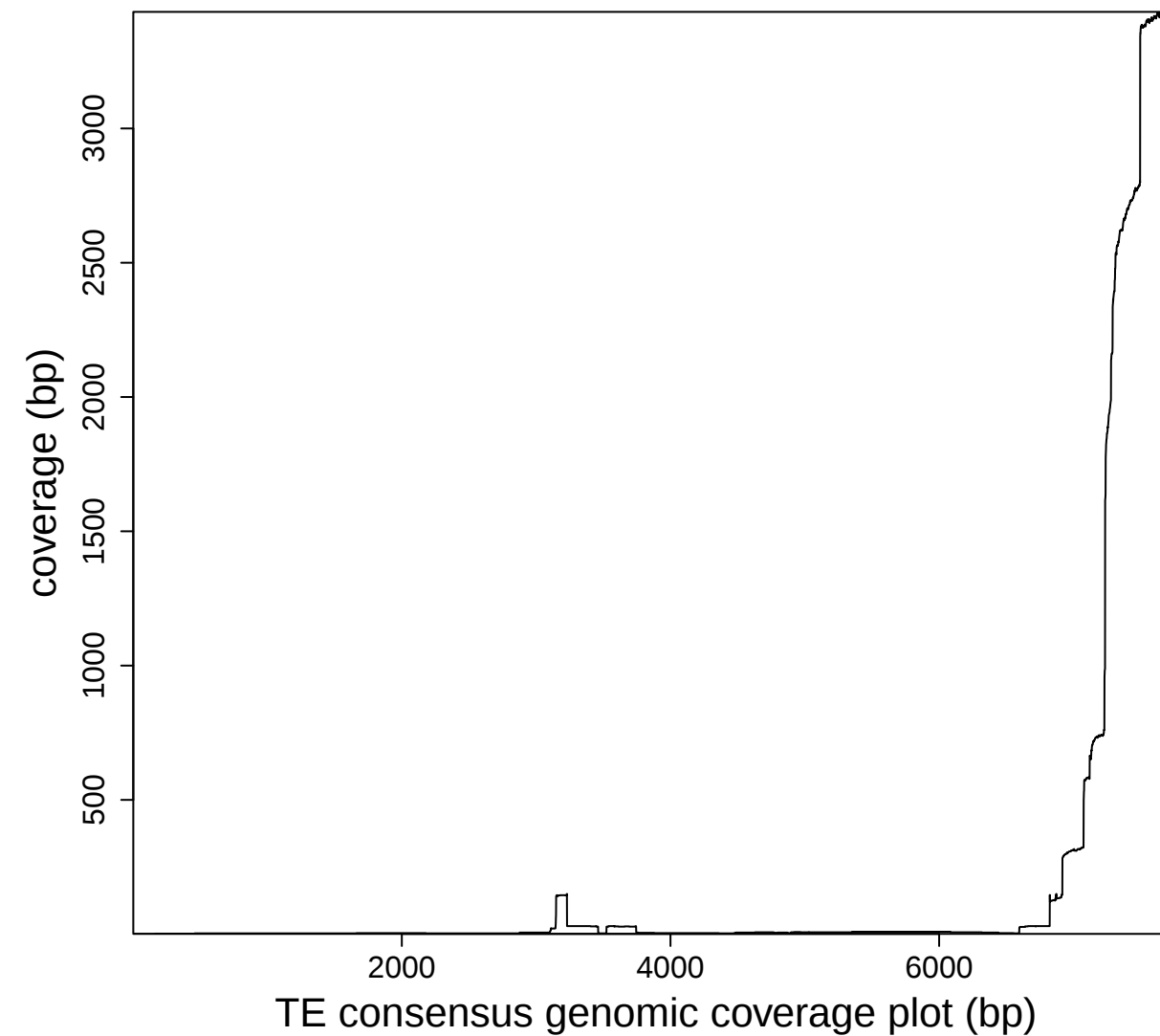
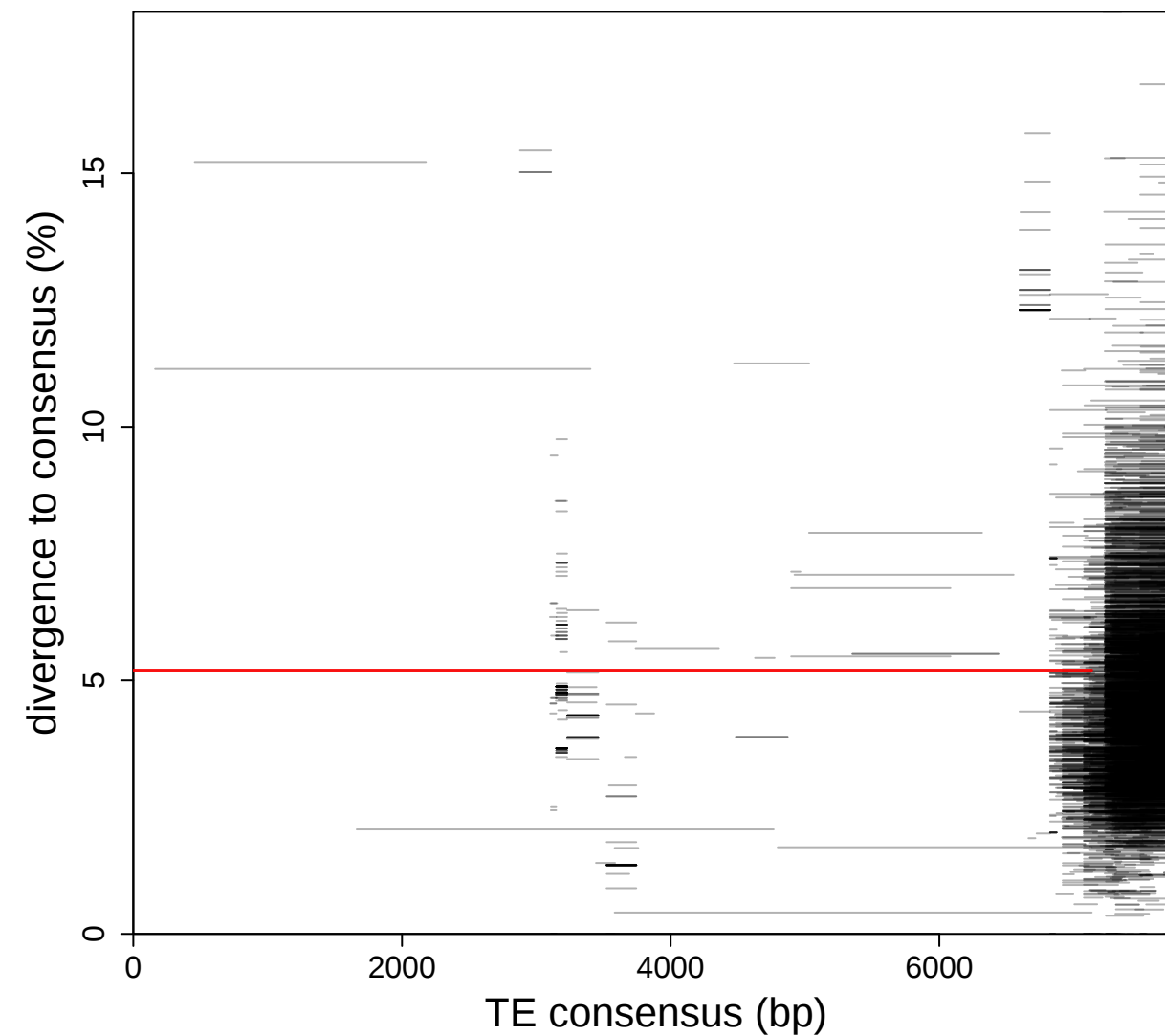
6.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 2963bp; fragments: 19; full length: 0 (>=2666.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

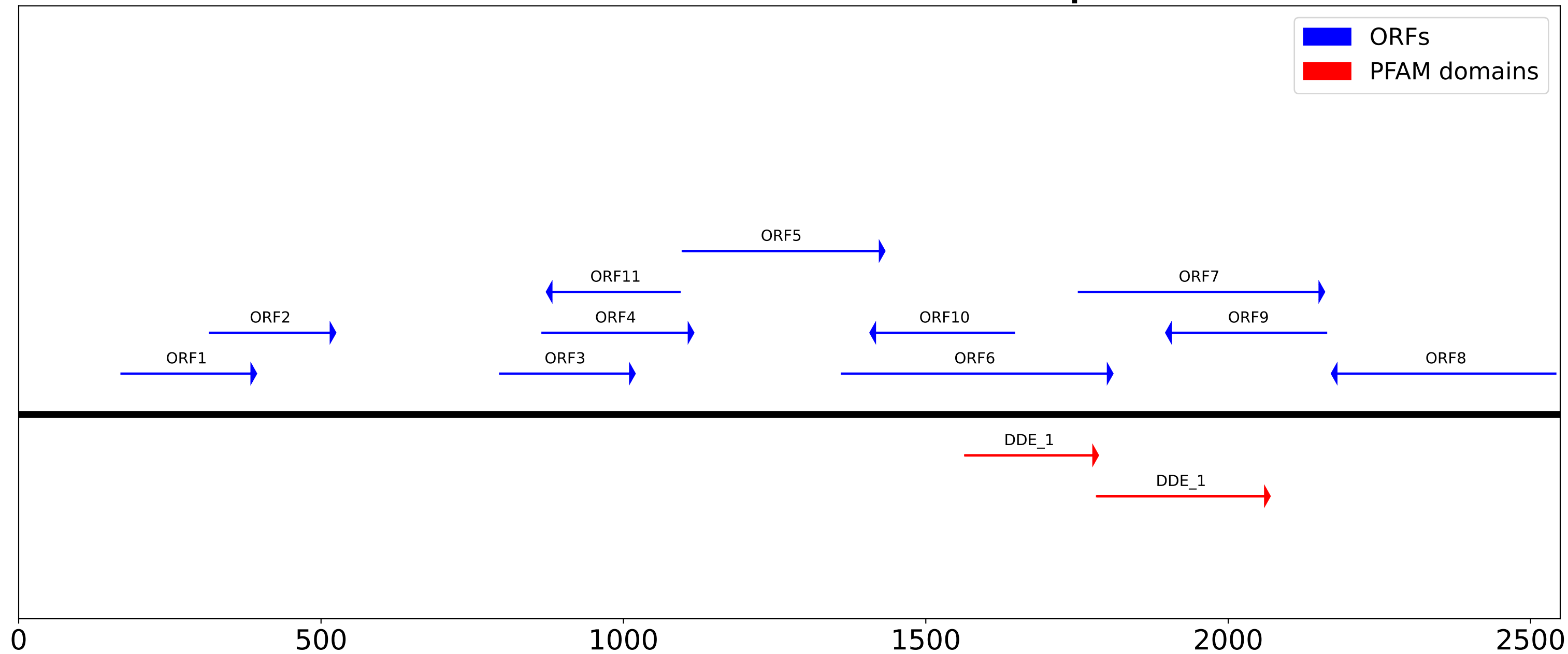


TE: rnd_5_family_1206
size: 7784bp; fragments: 4510; full length: 1 (≥ 7005.6 bp)

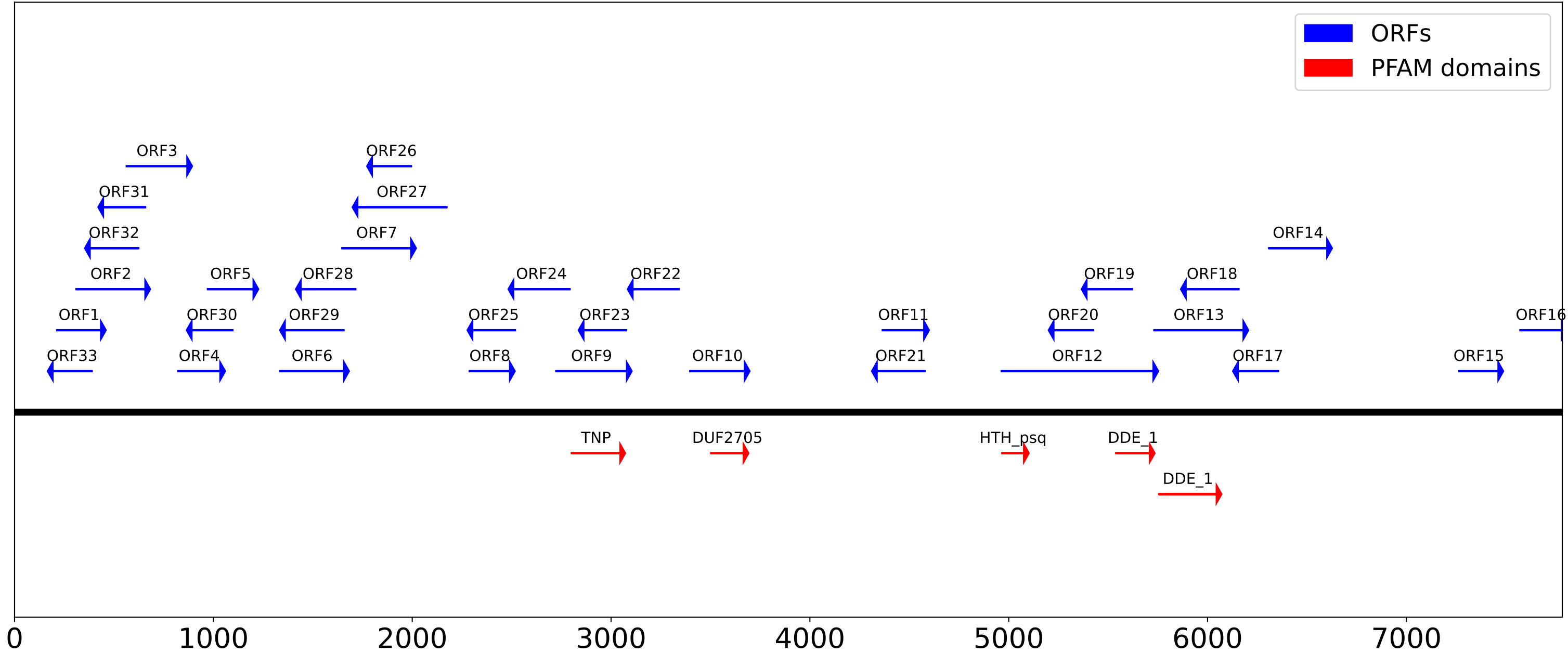
Before TEtrimmer 7784 bp



After TETrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

