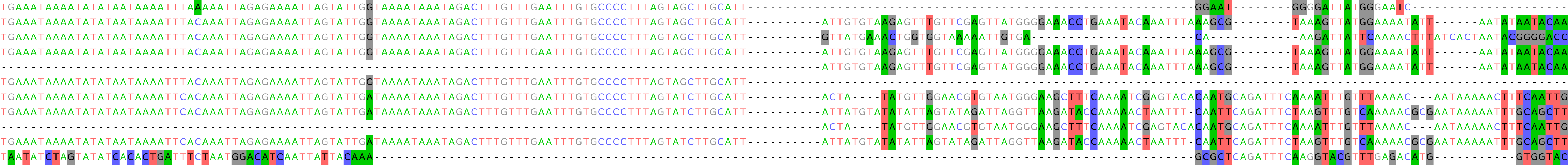


Start crop Point



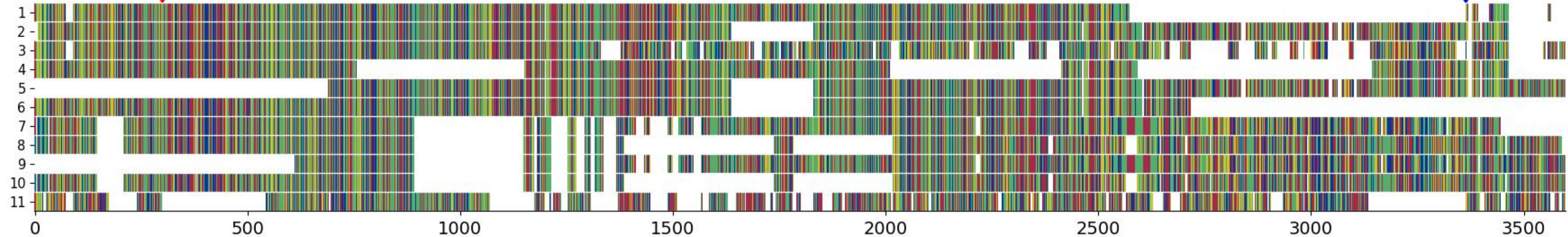
MSA length = 3065

End crop Point



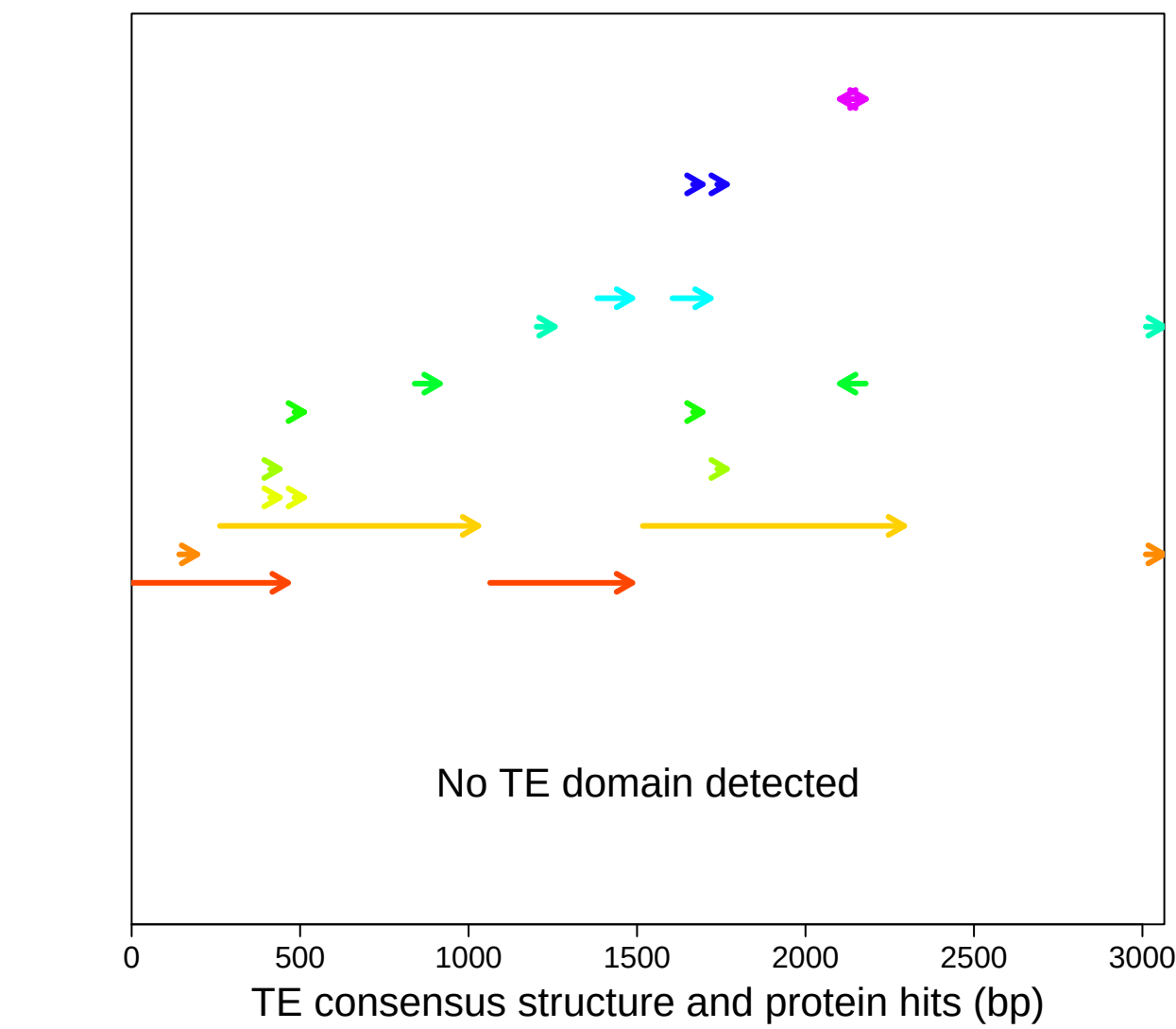
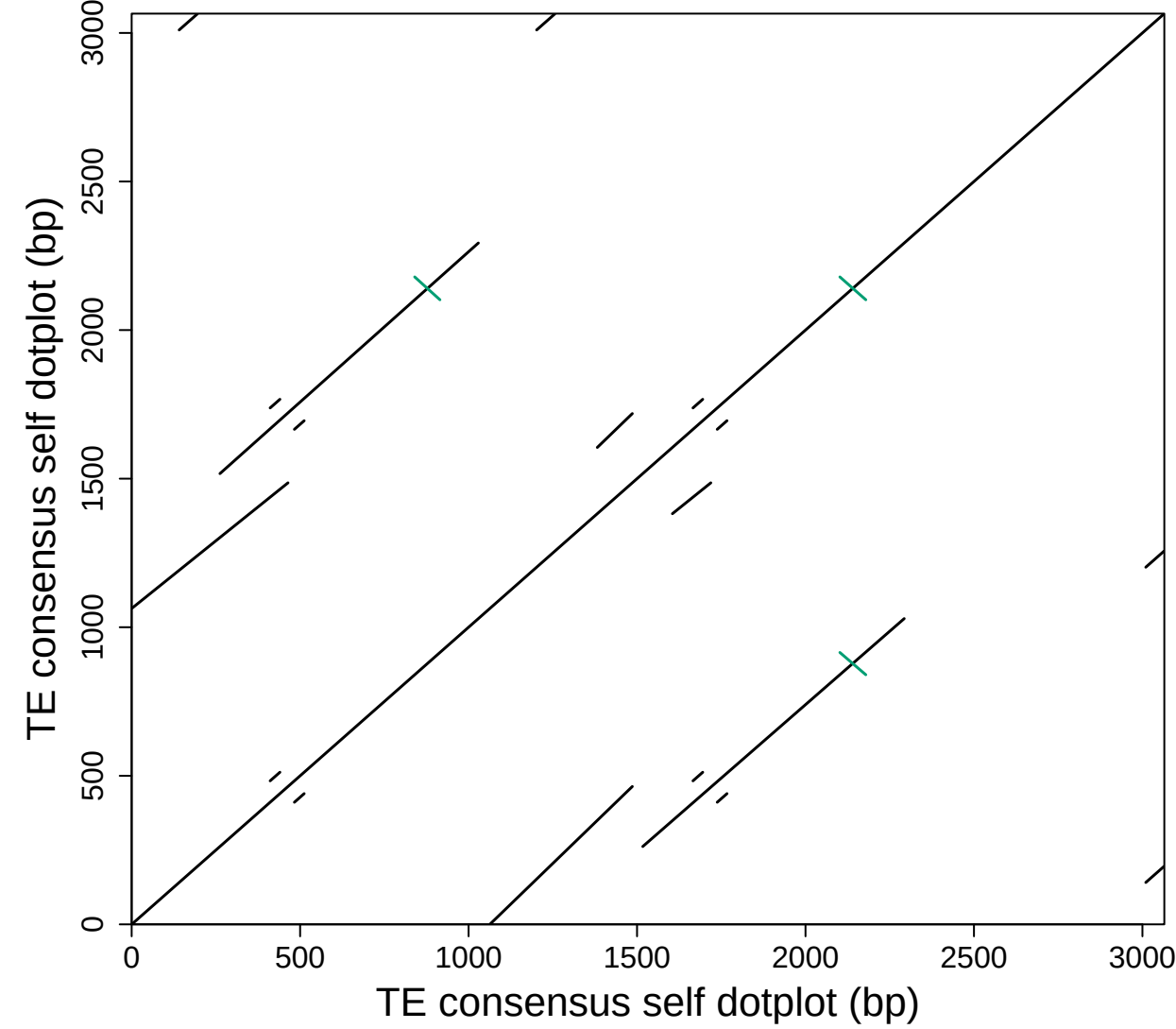
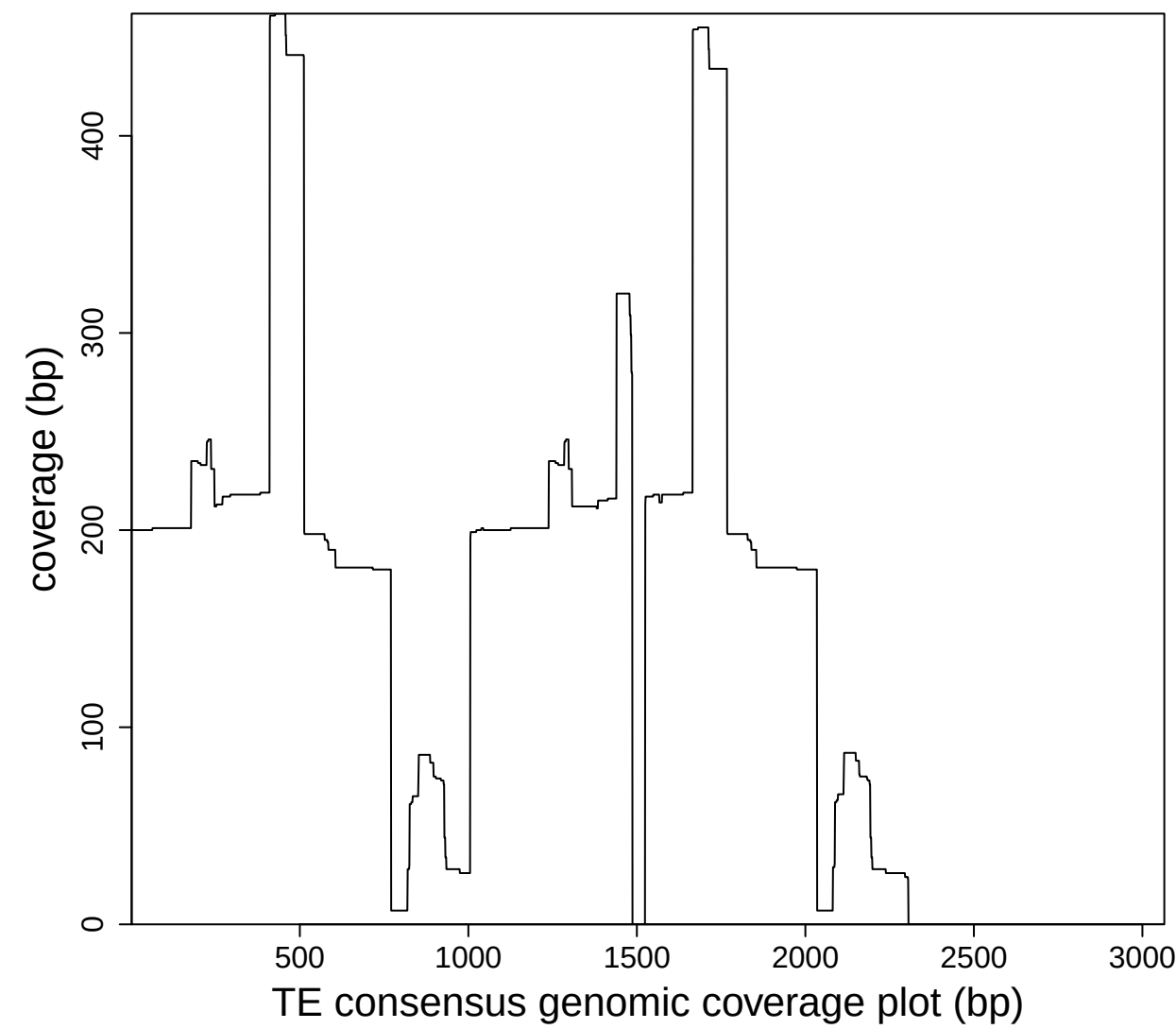
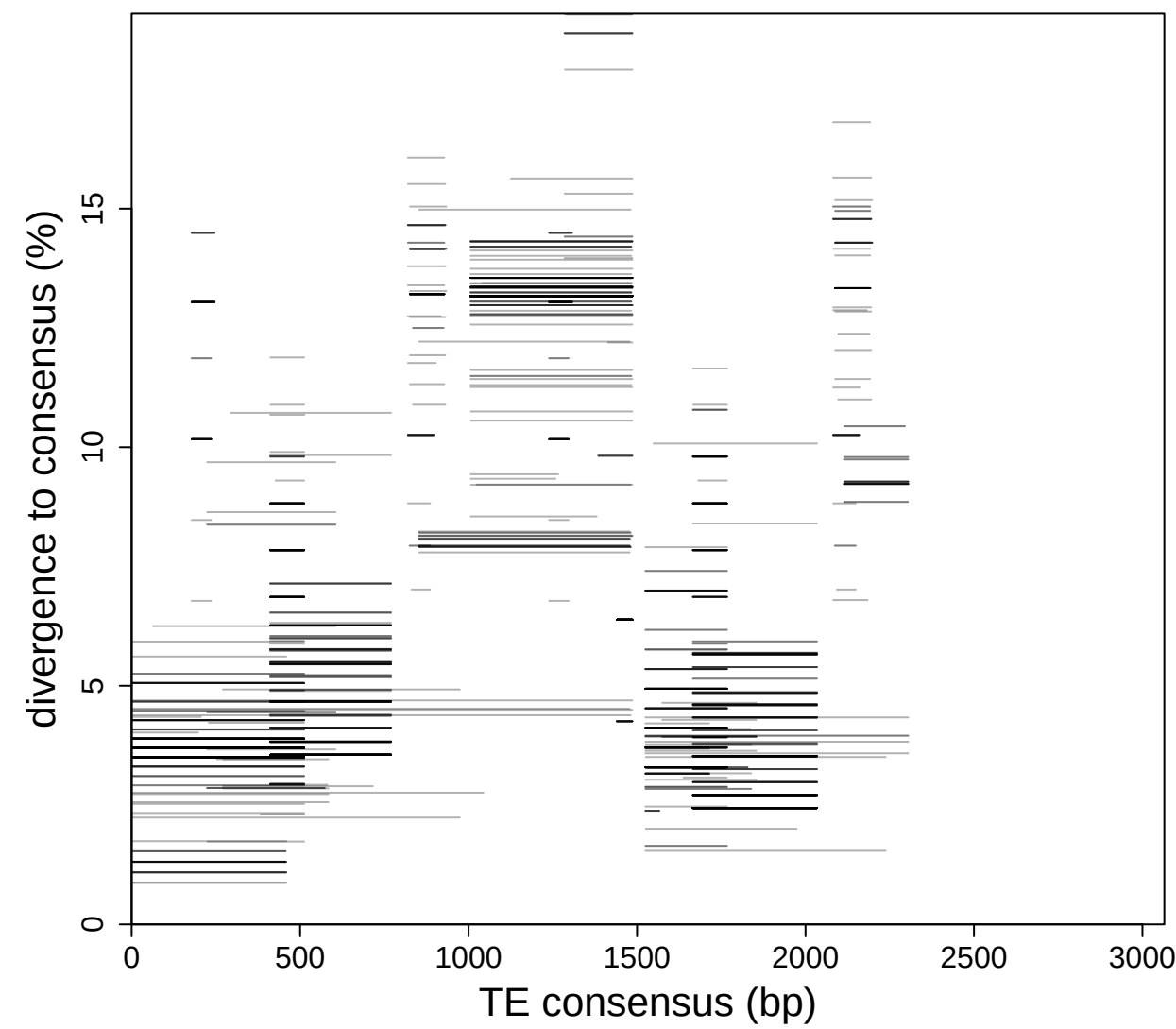
Start crop Point

End crop Point



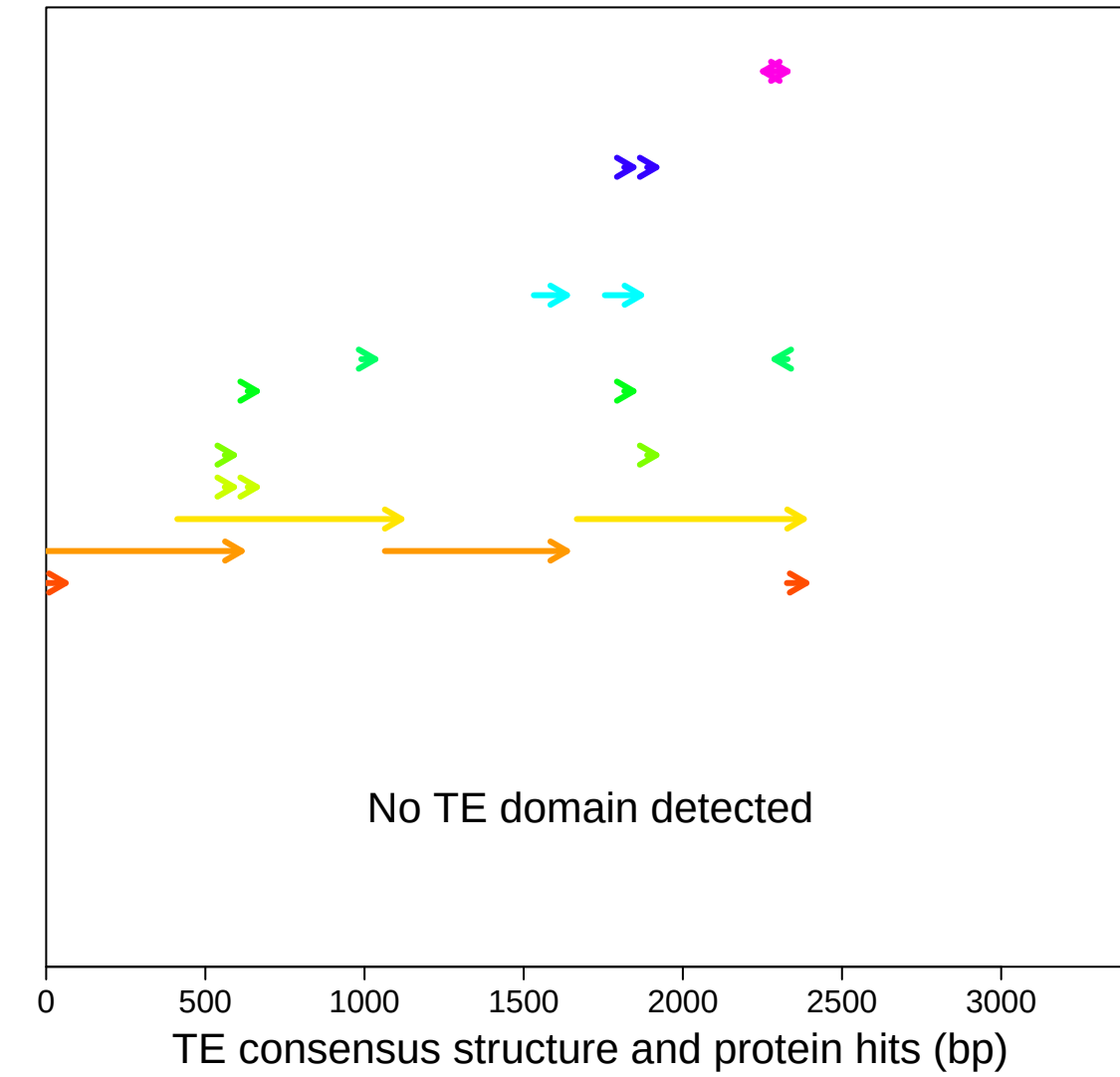
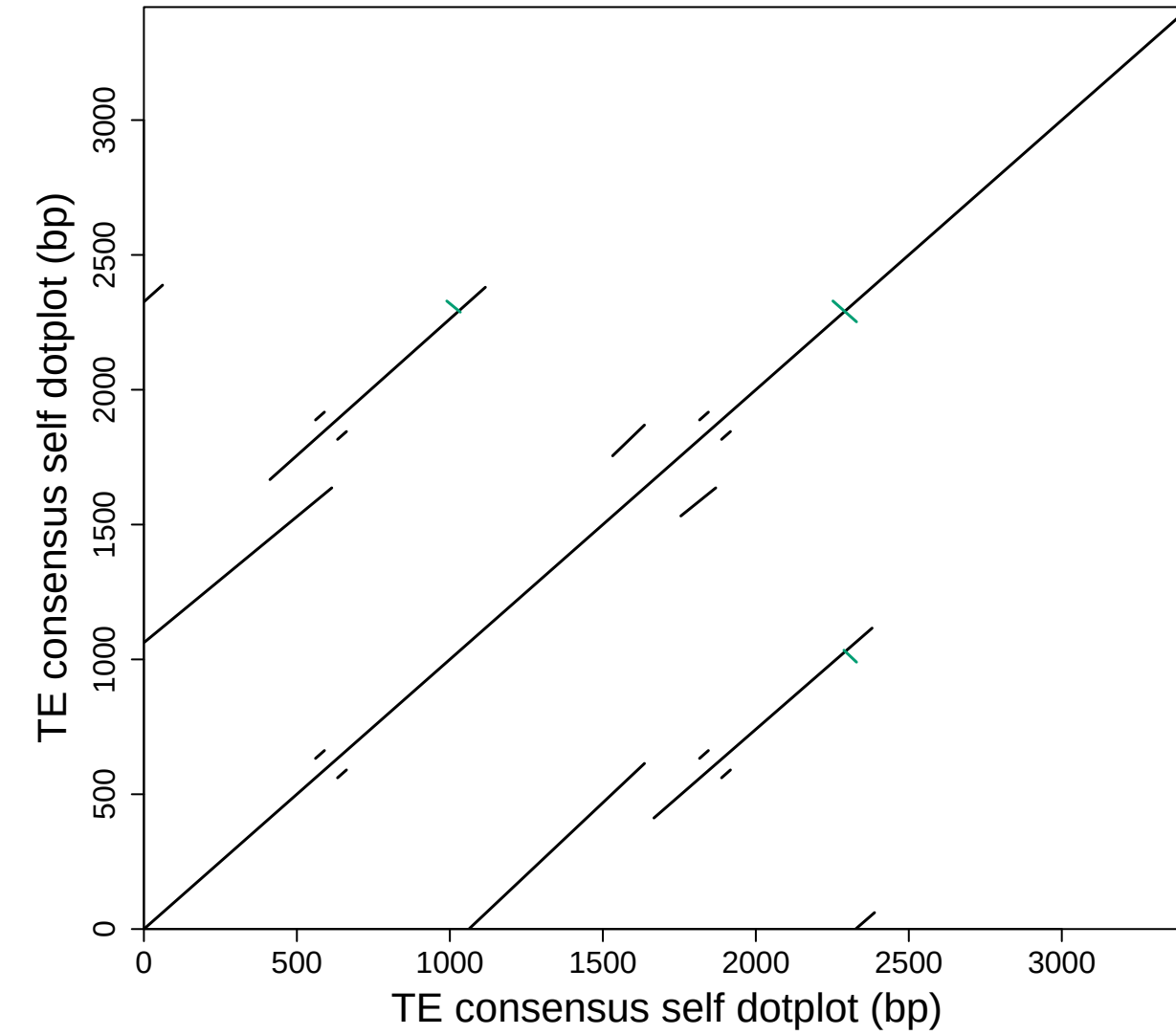
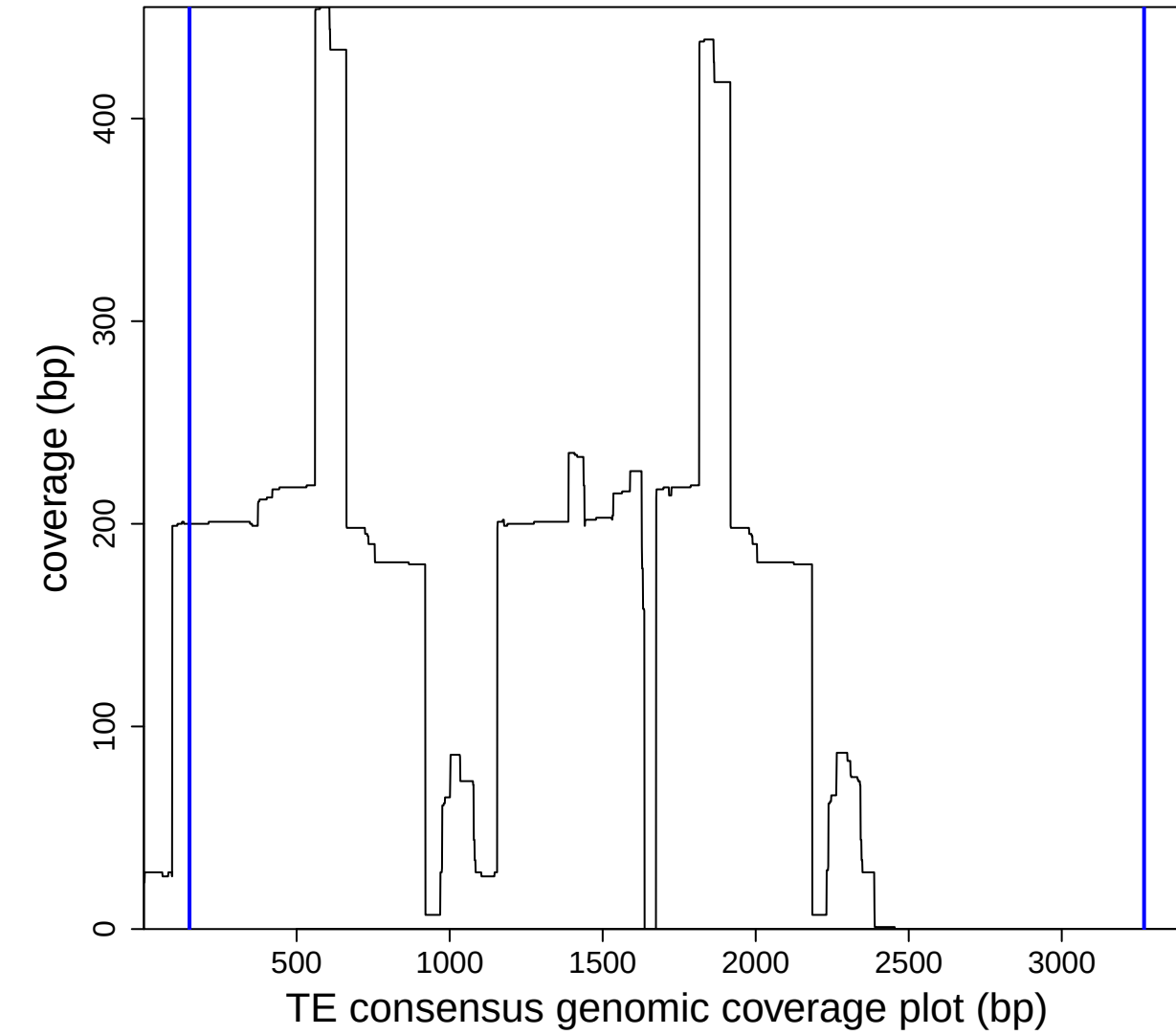
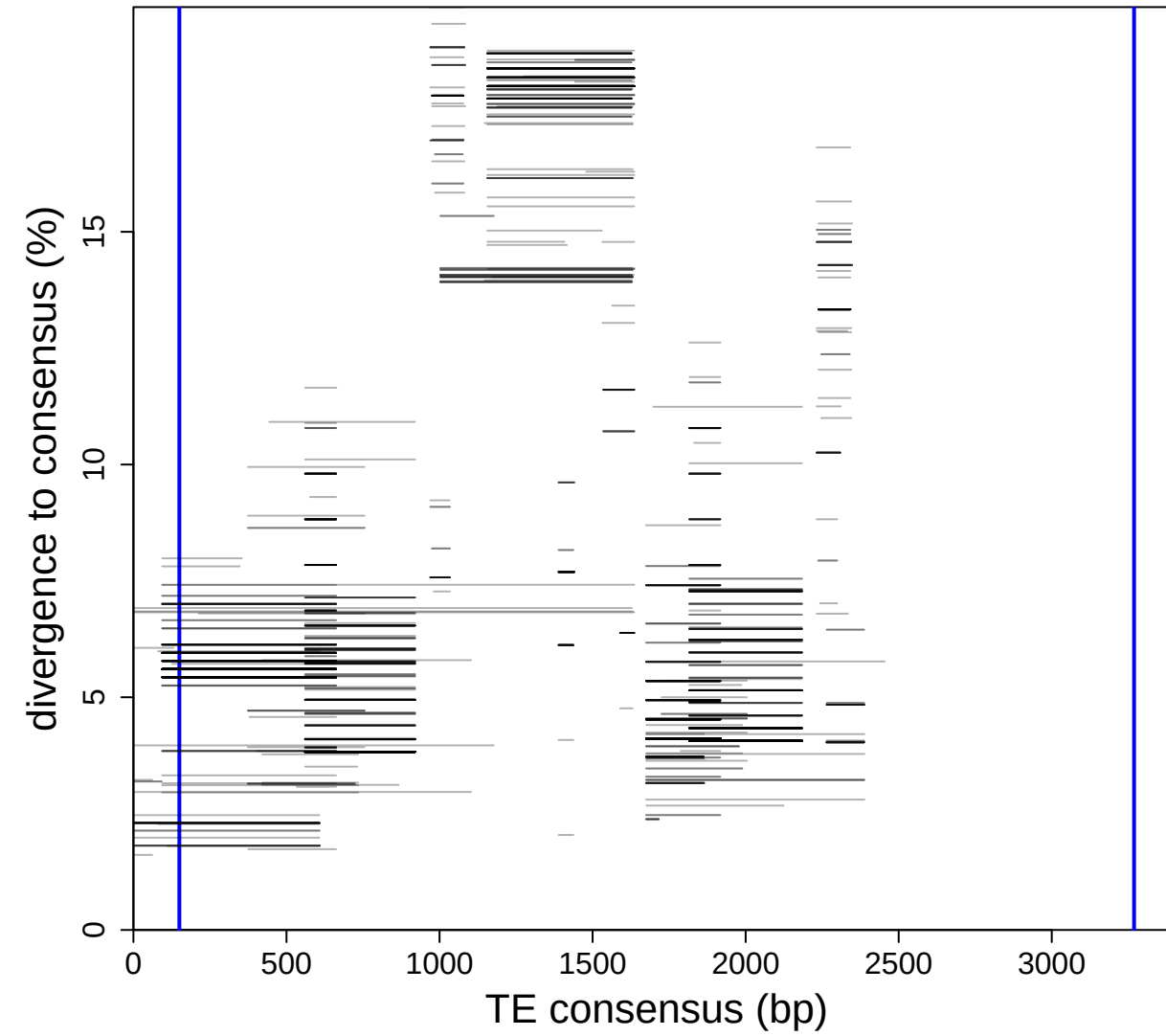
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 3065bp; fragments: 1448; full length: 0 (>=2758.5bp)

After TEtrimmer 3065 bp



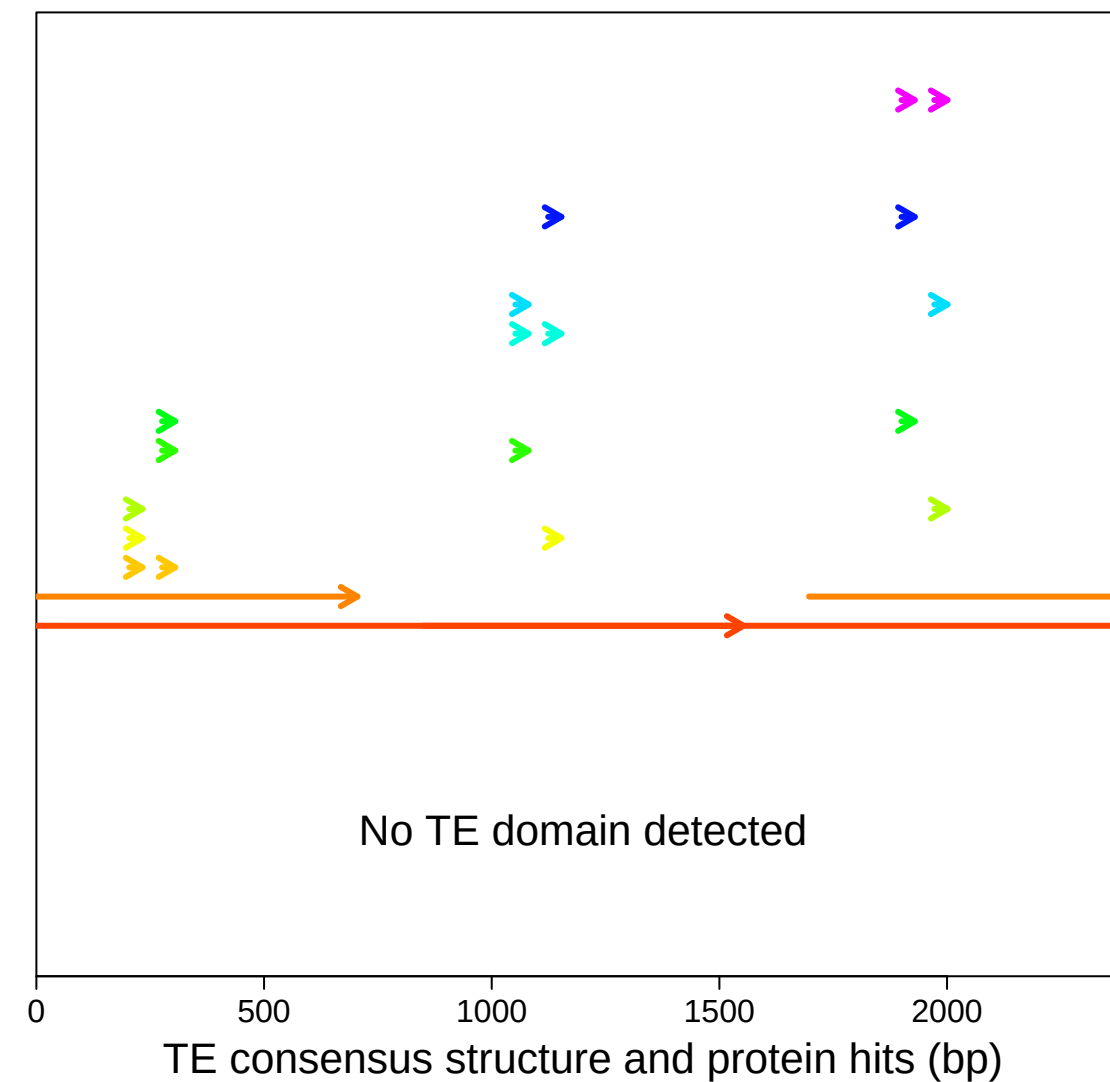
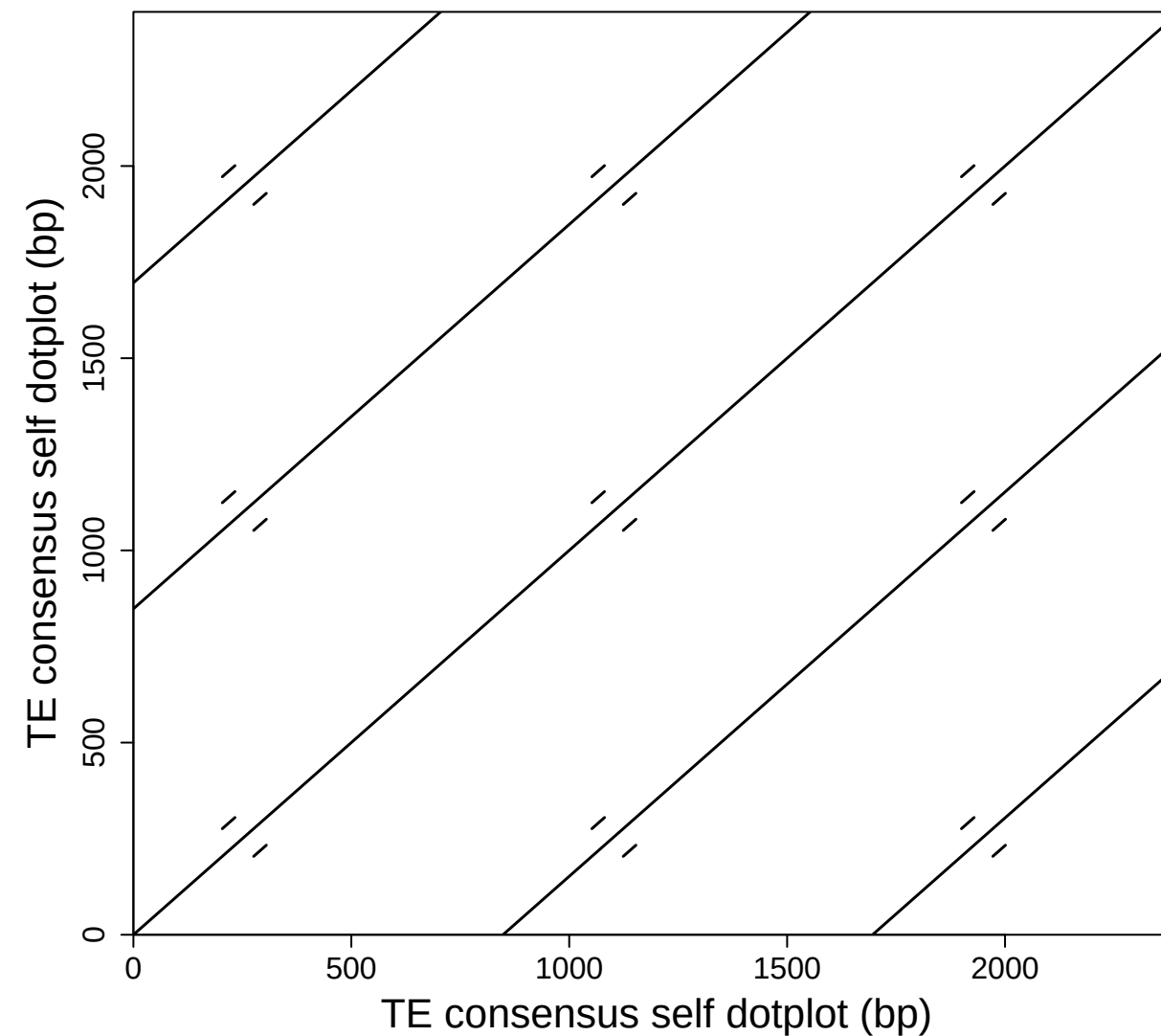
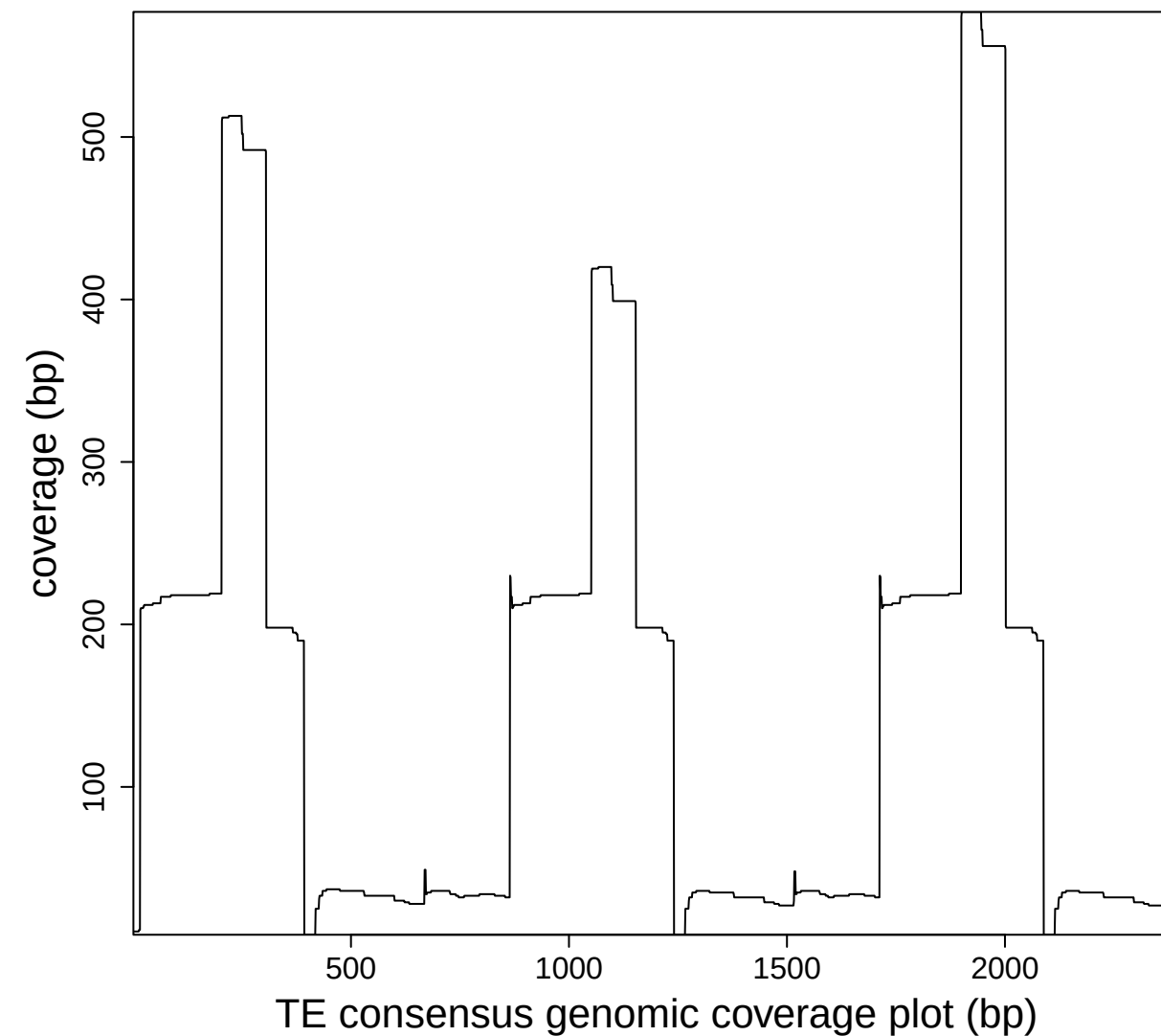
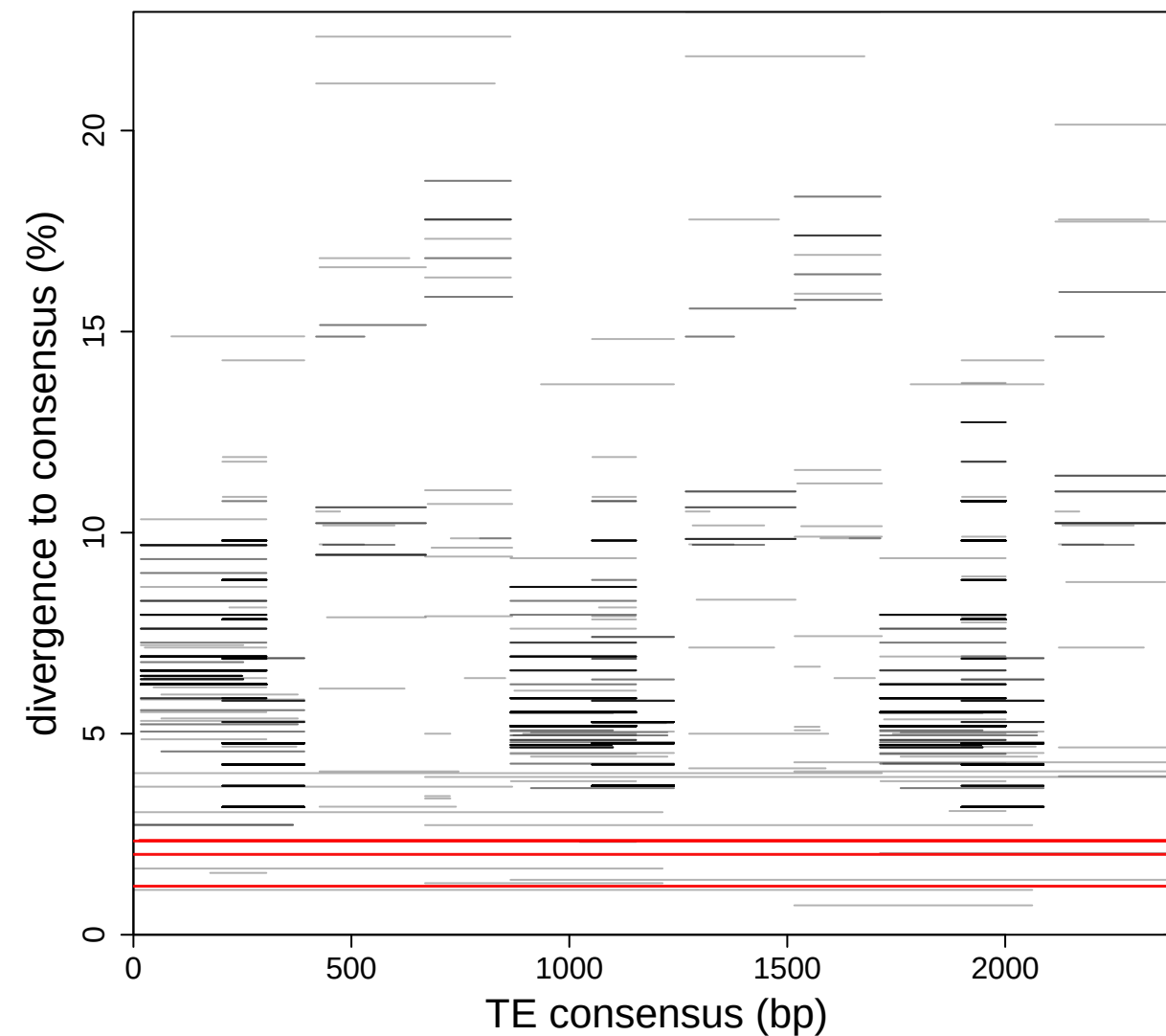
ily_313.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 3419bp; fragments: 1304; full length: 0 (>=3077.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

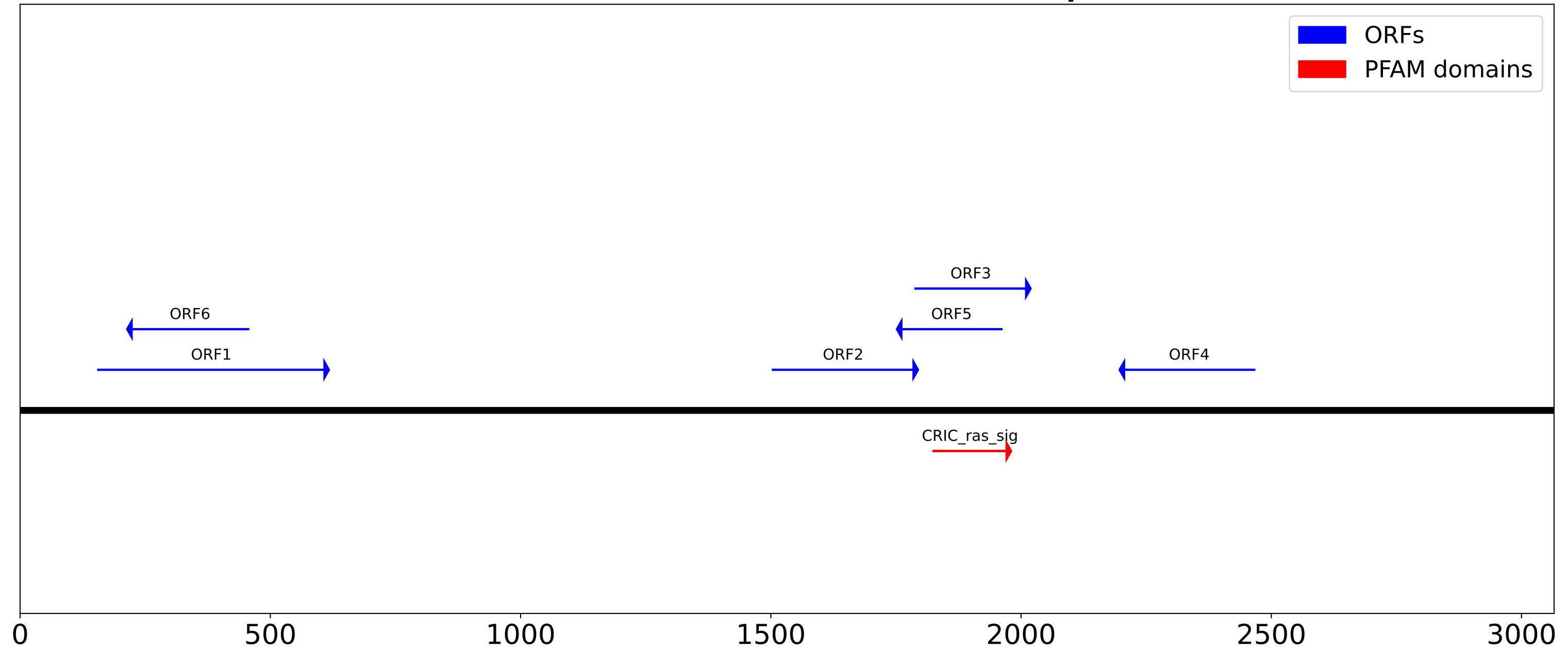


TE: rnd_1_family_313
size: 2401bp; fragments: 1627; full length: 4 (≥ 2160.9 bp)

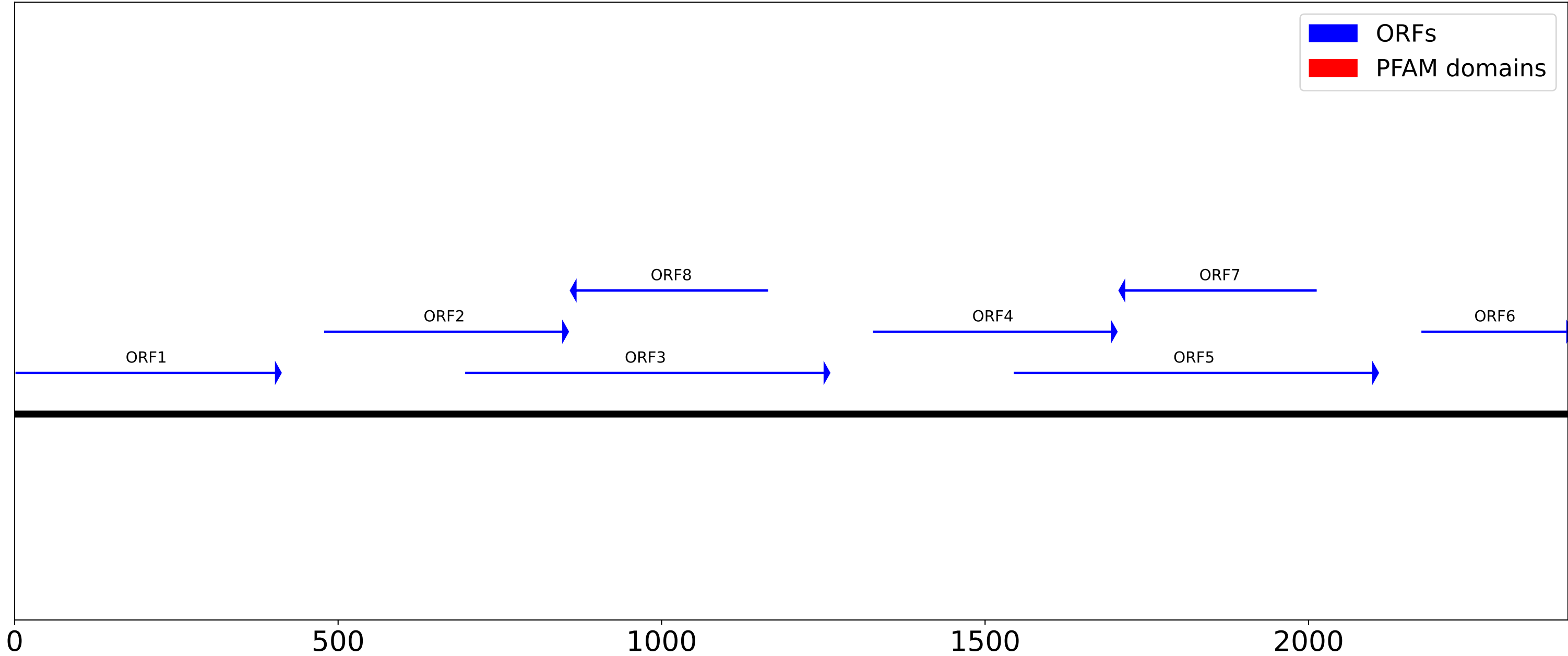
Before TEtrimmer 2401 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

