

Start crop Point



End crop Point

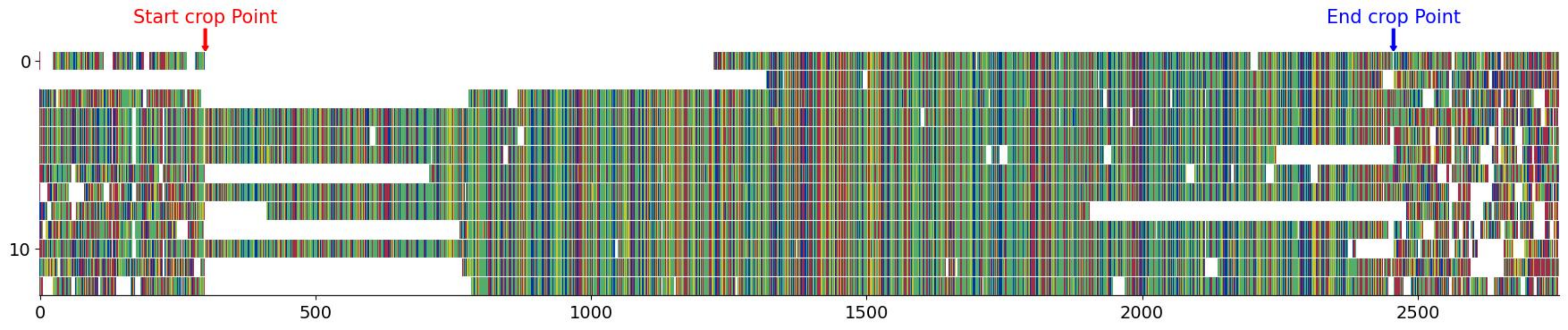


MSA length = 2157



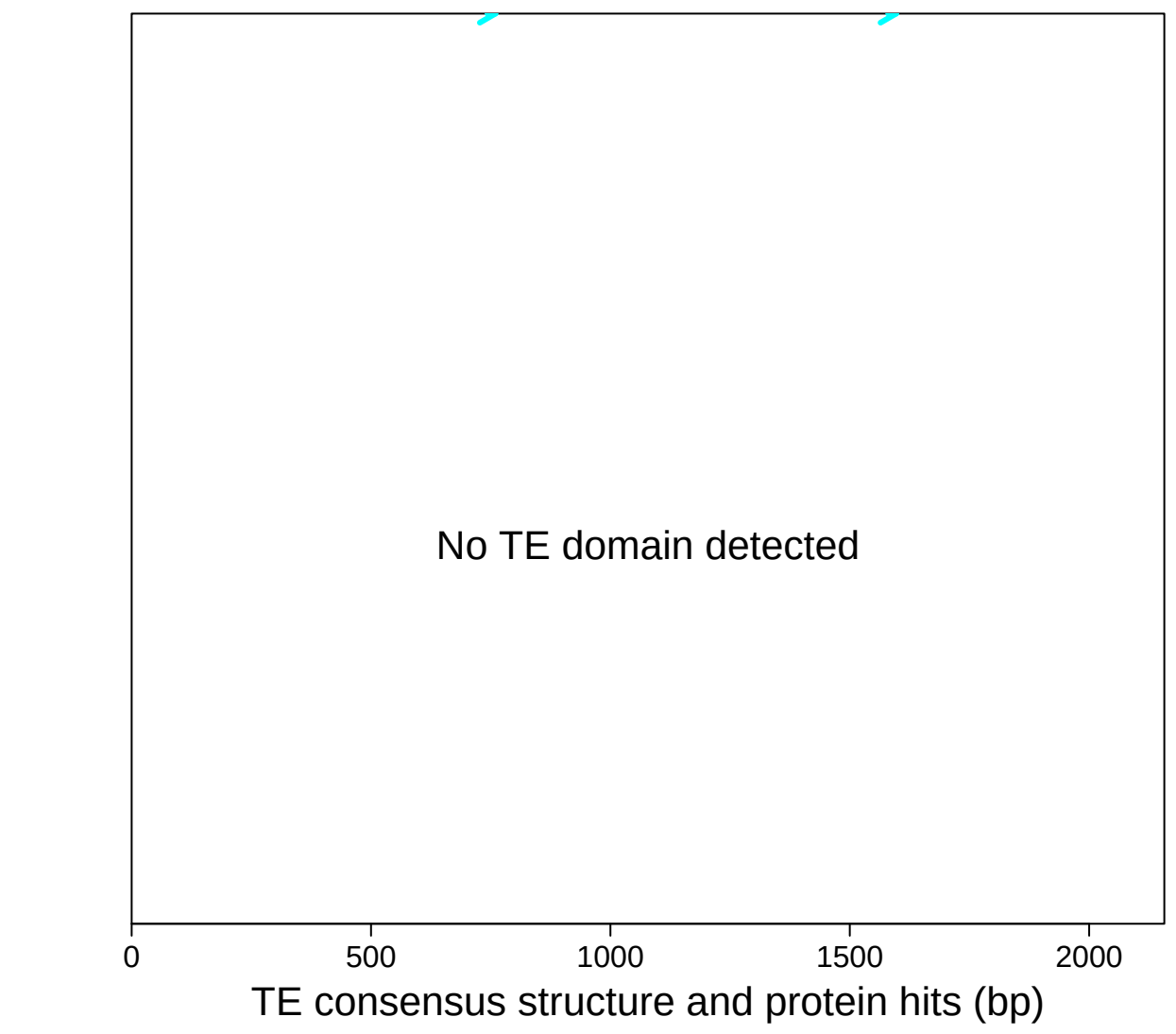
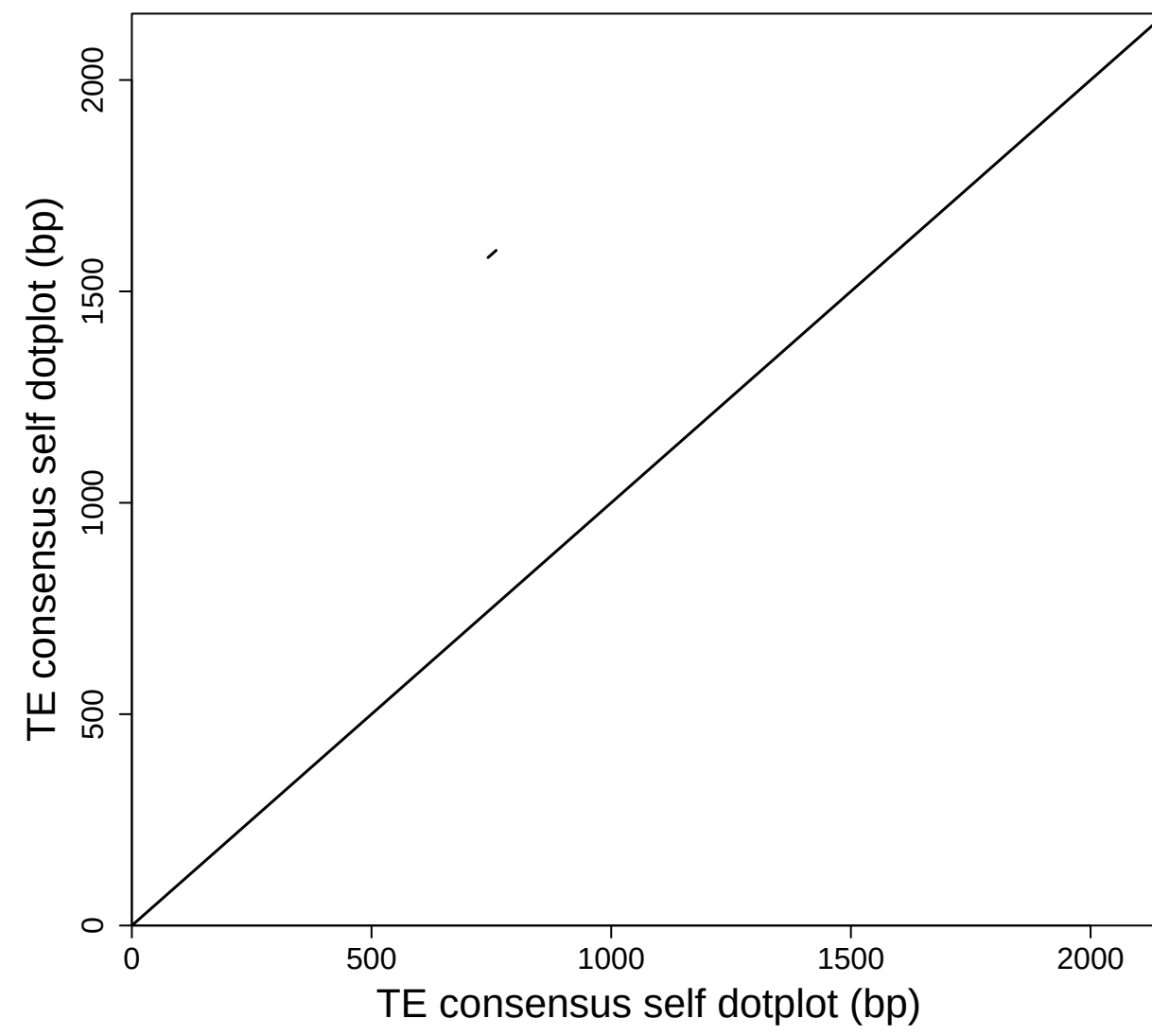
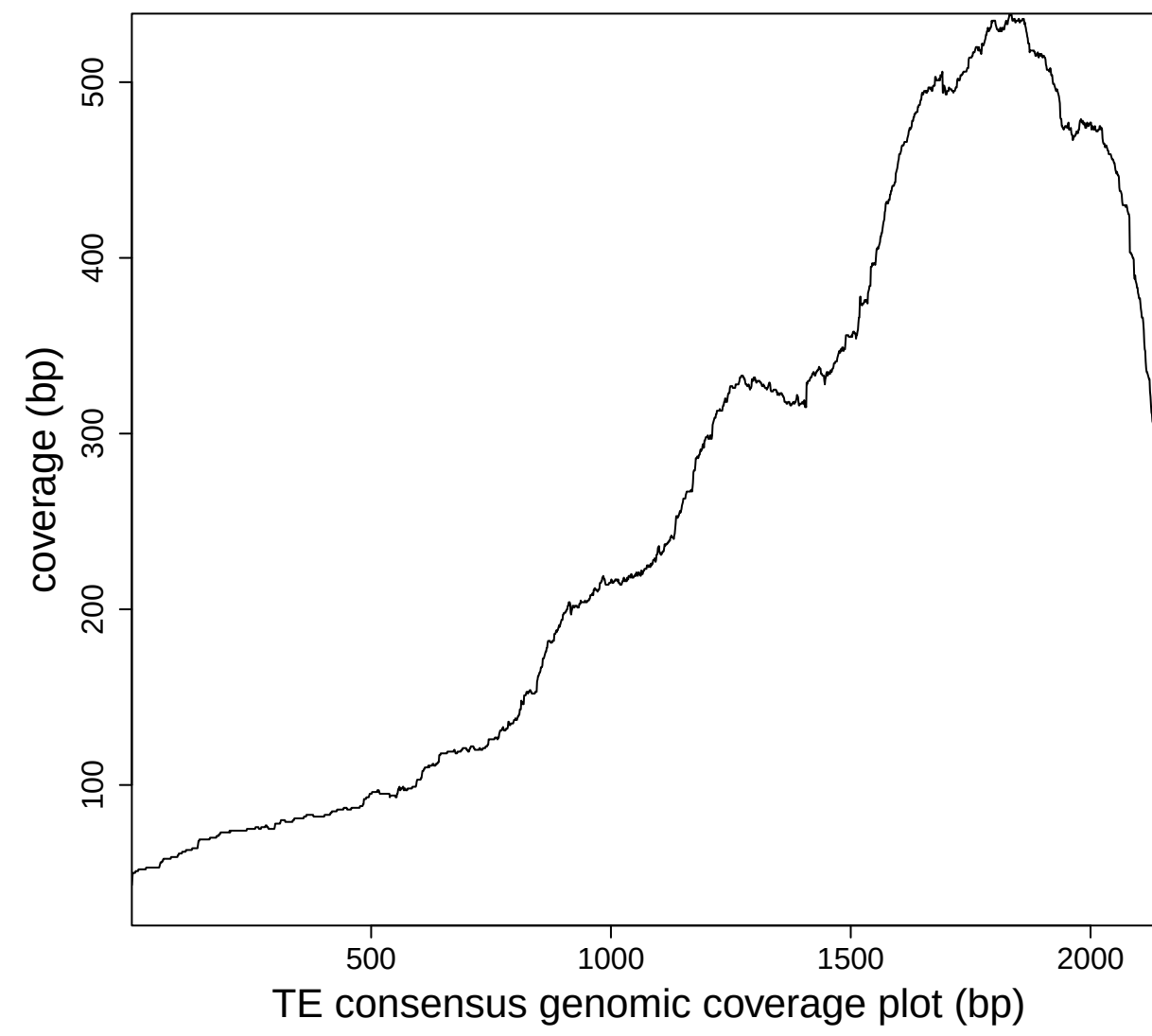
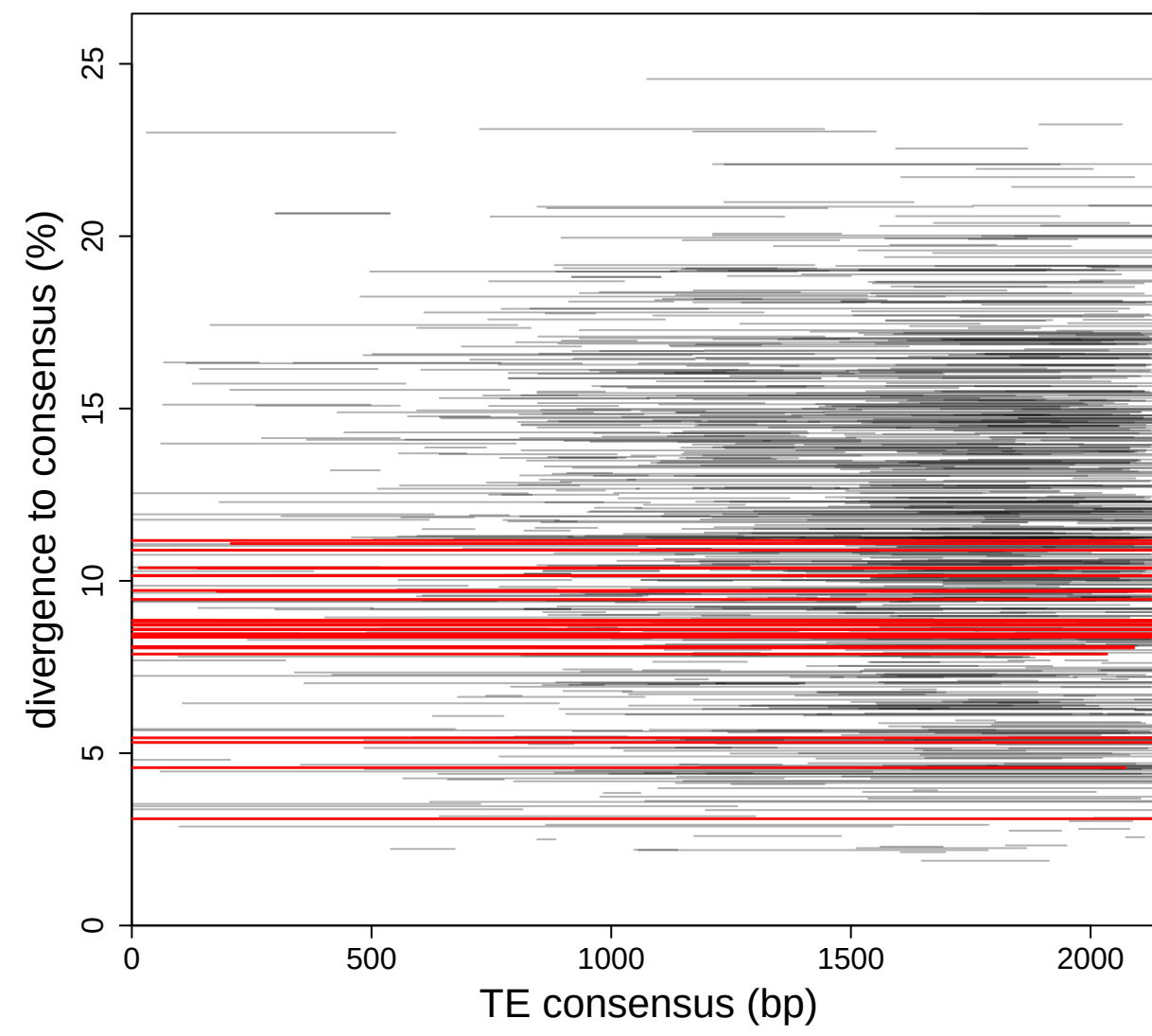
Start crop Point

End crop Point



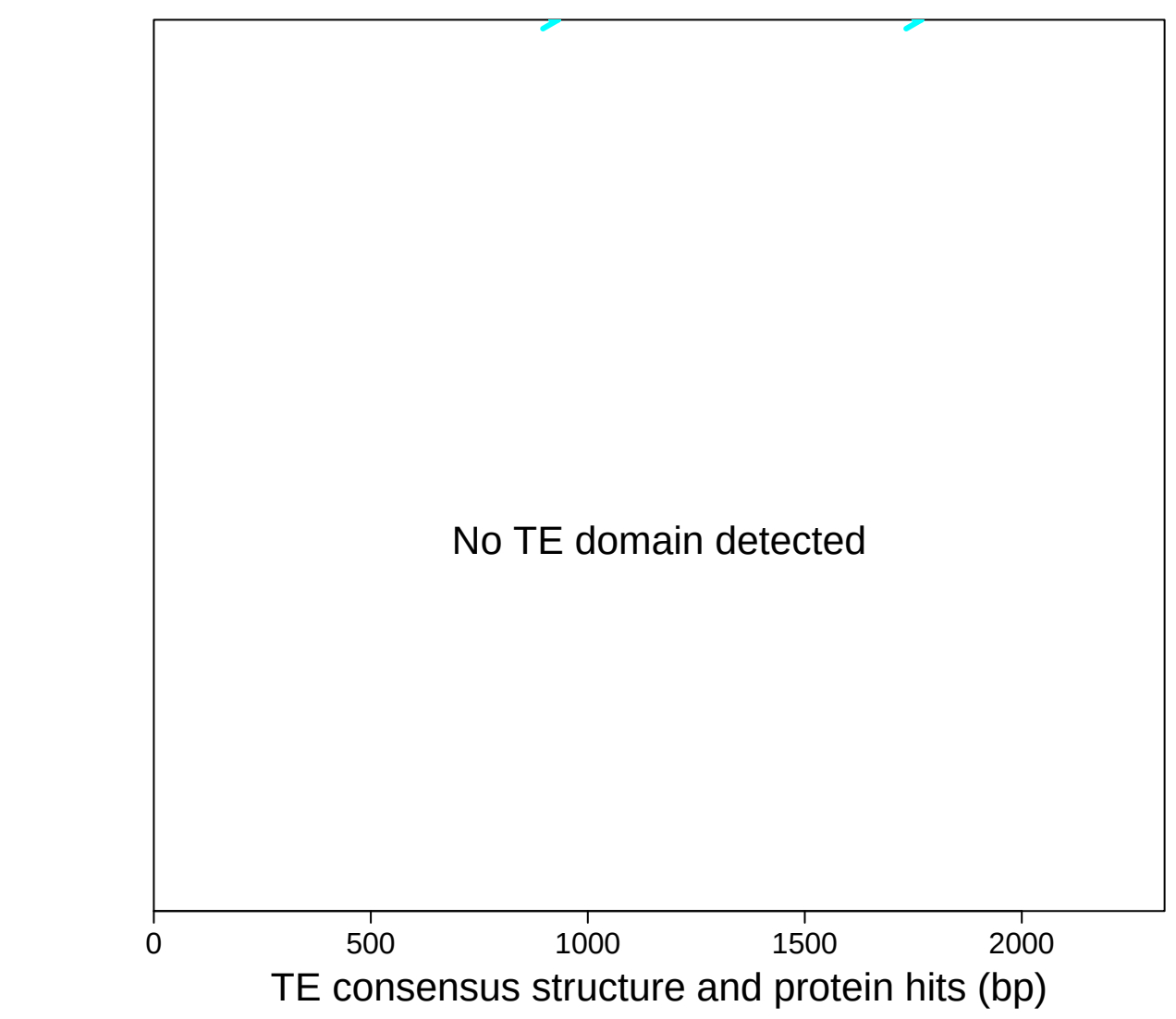
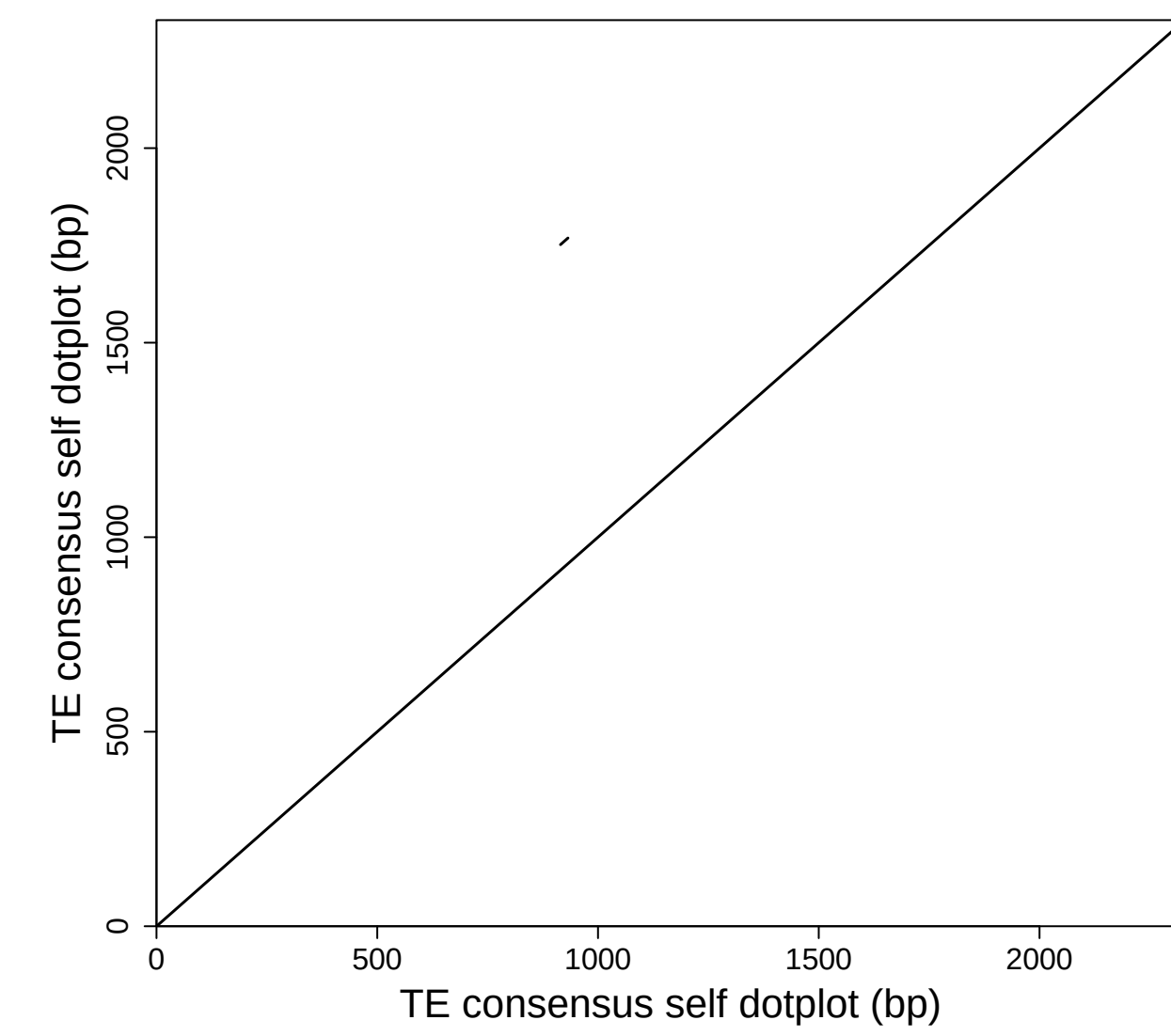
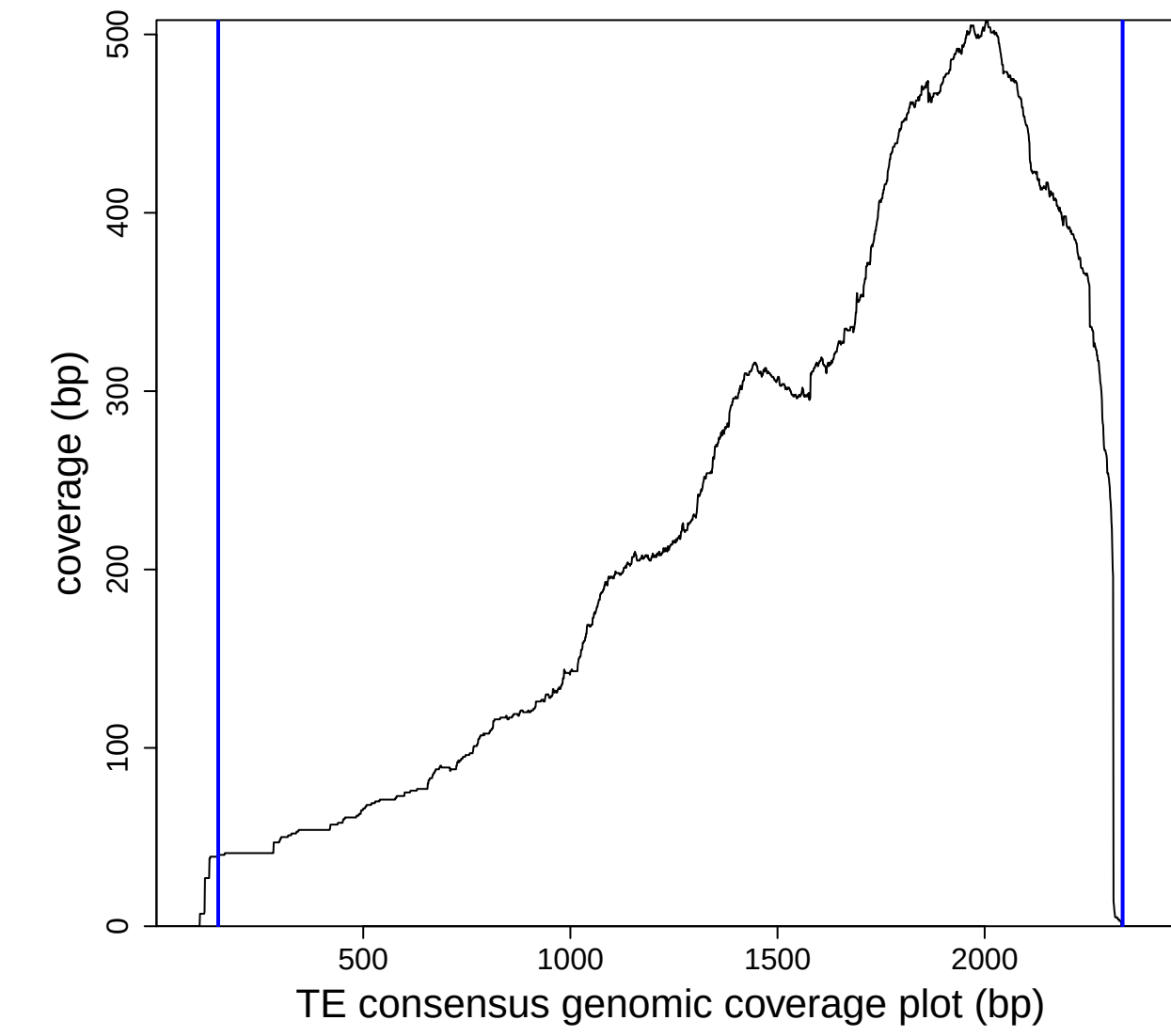
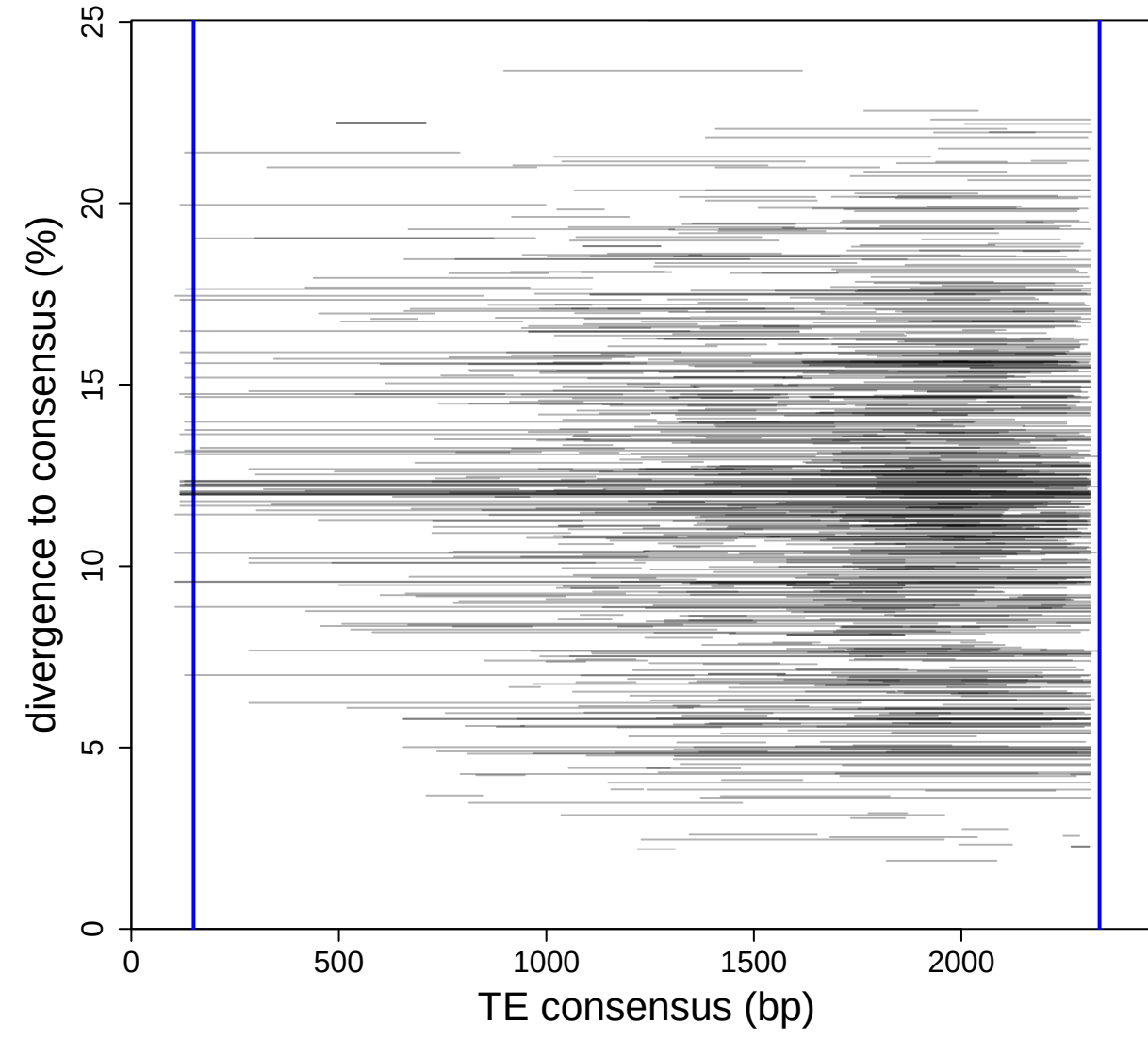
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 2157bp; fragments: 991; full length: 31 (>=1941.3bp)

After TEtrimmer 2157 bp



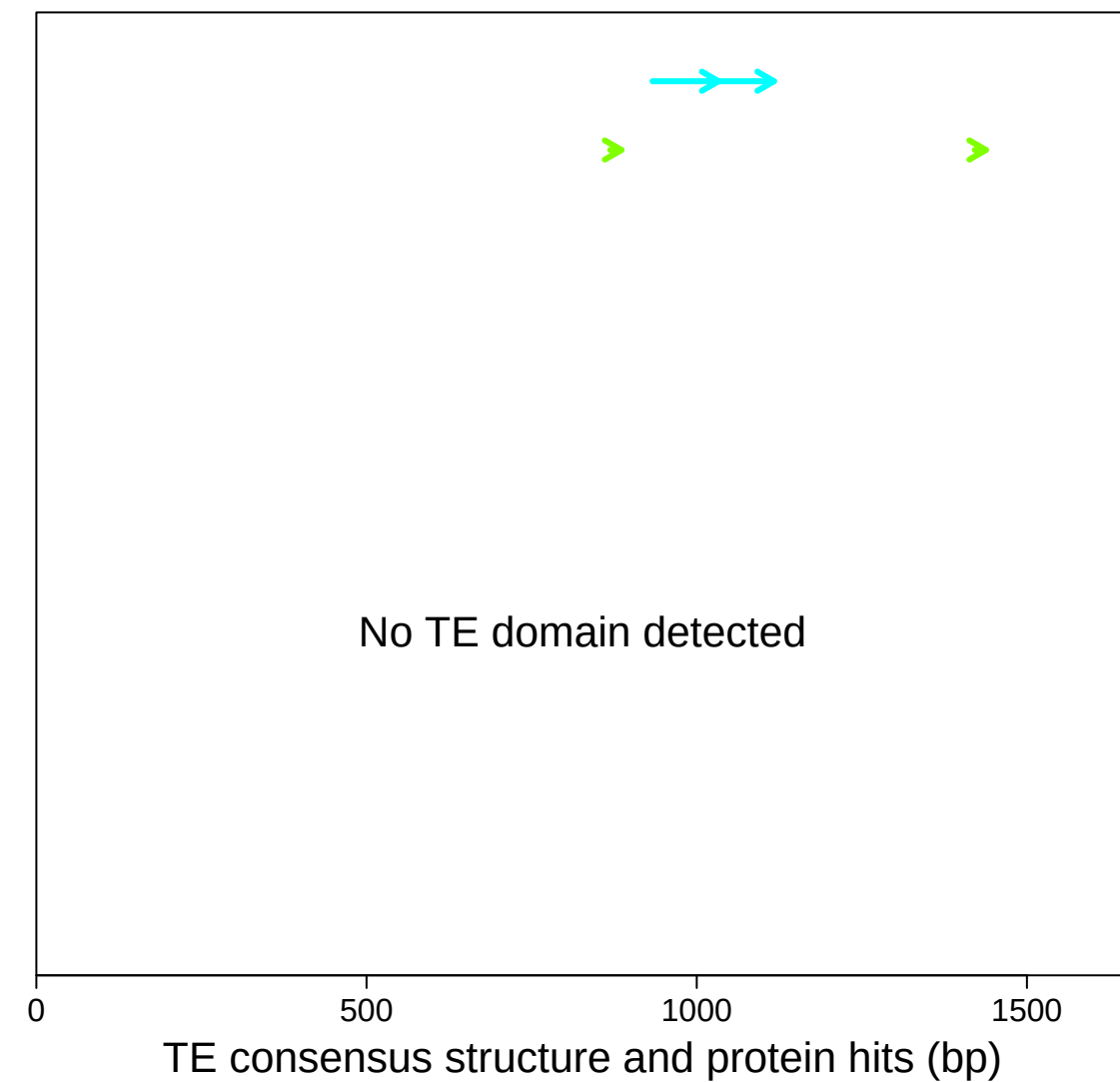
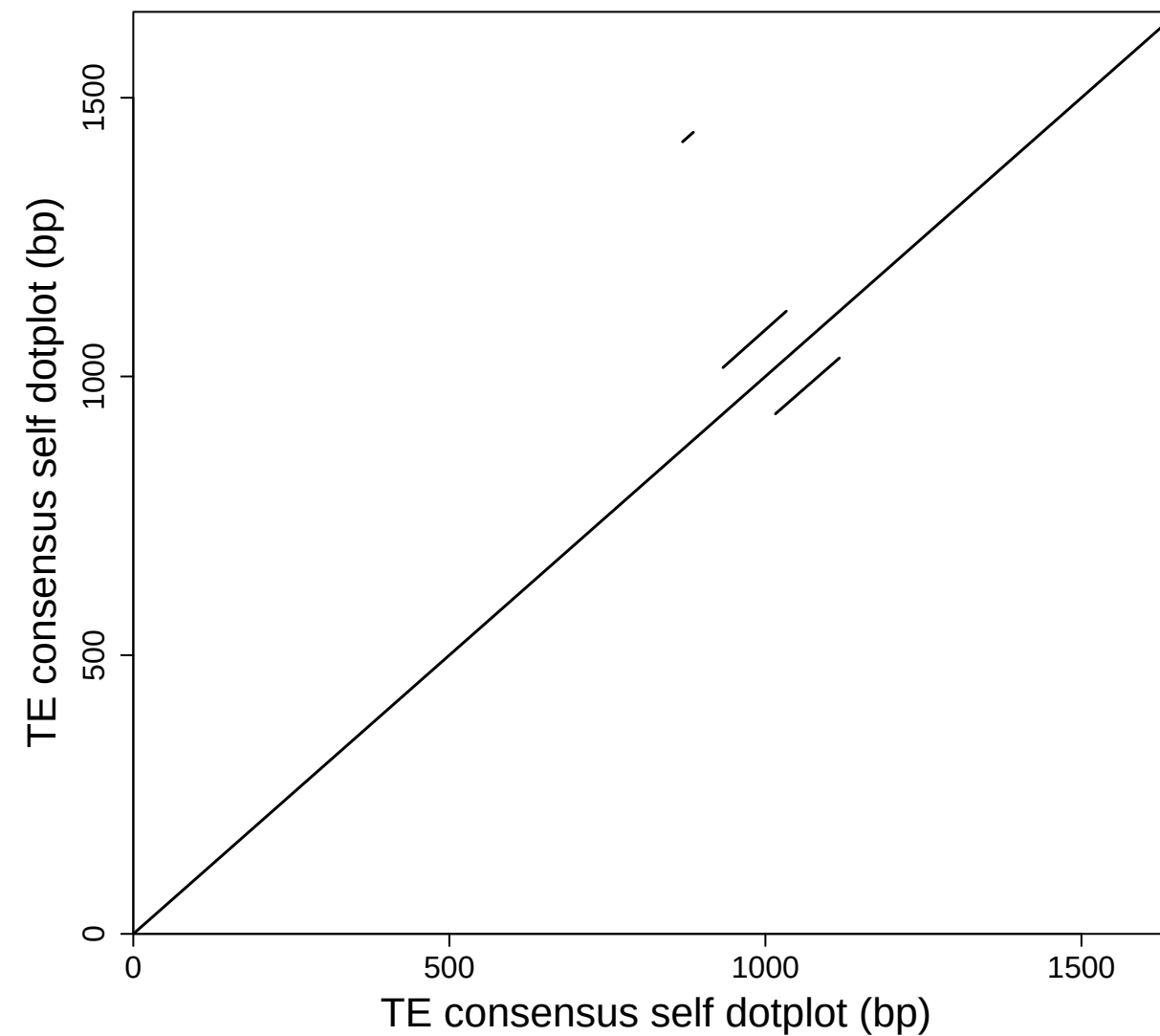
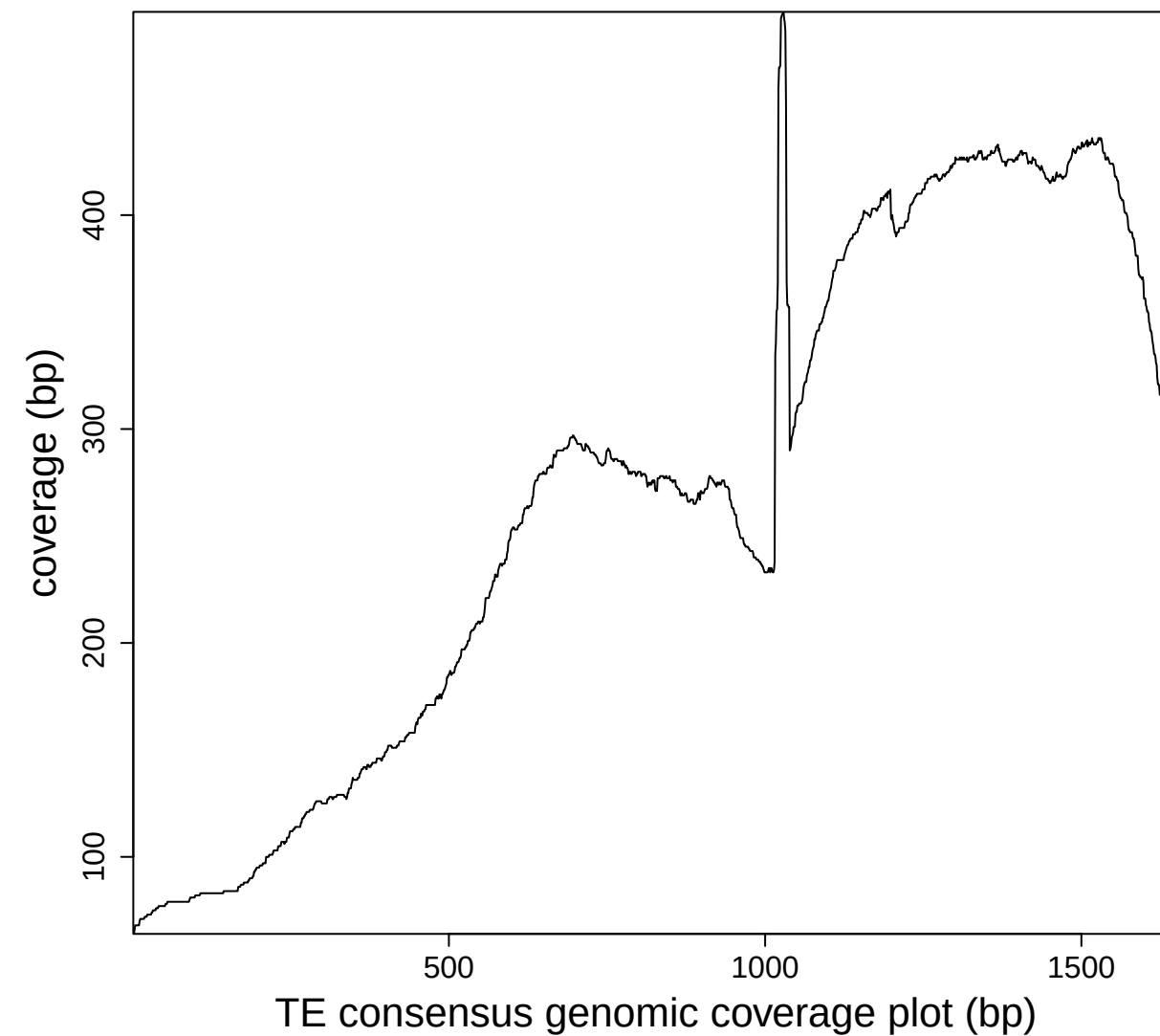
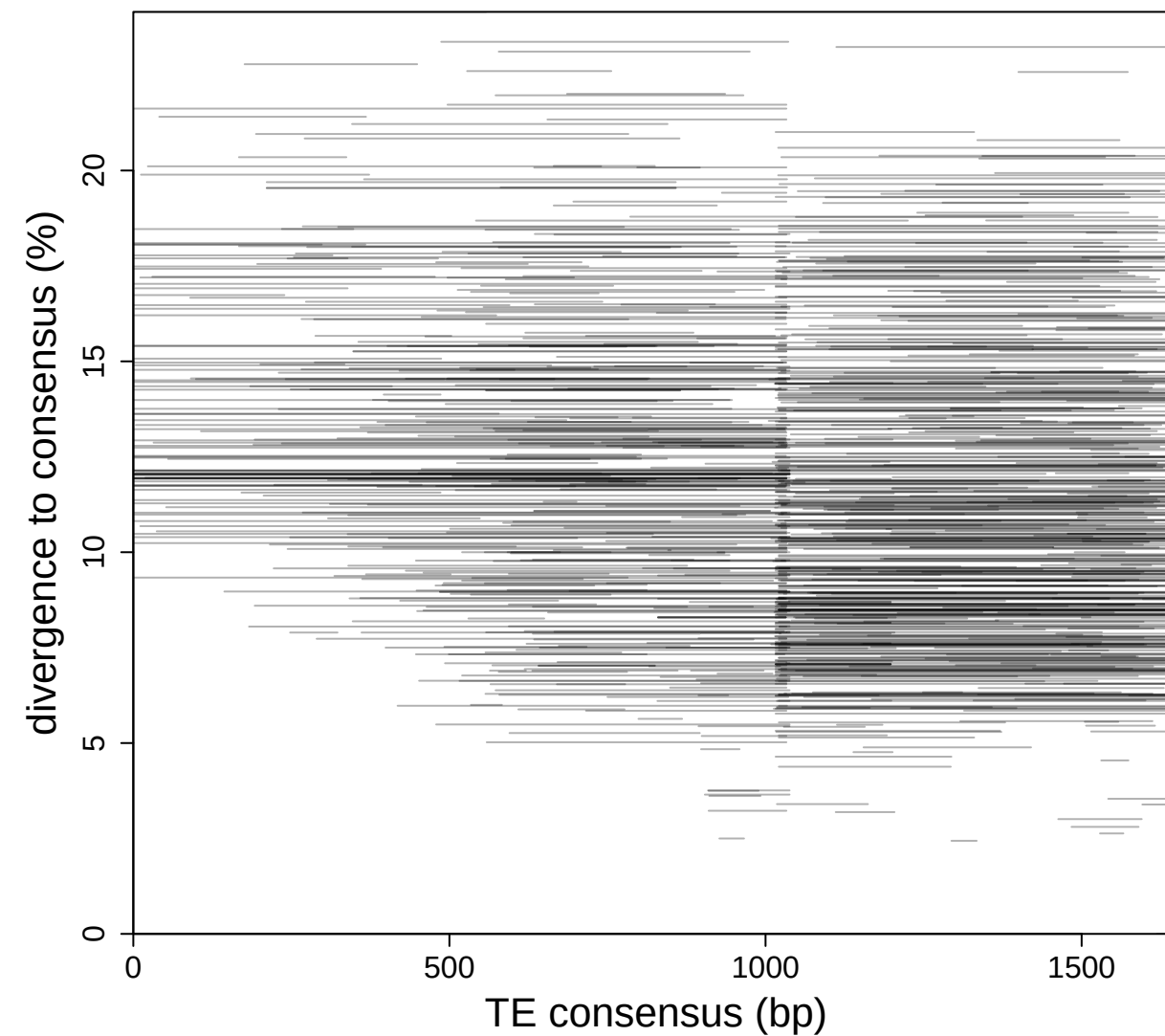
6.fasta.b.bed uf.bed g_3.bed fm_1.bed_0_0 bcln.fa aln.fa cl.fa
size: 2483bp; fragments: 878; full length: 0 (>=2234.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

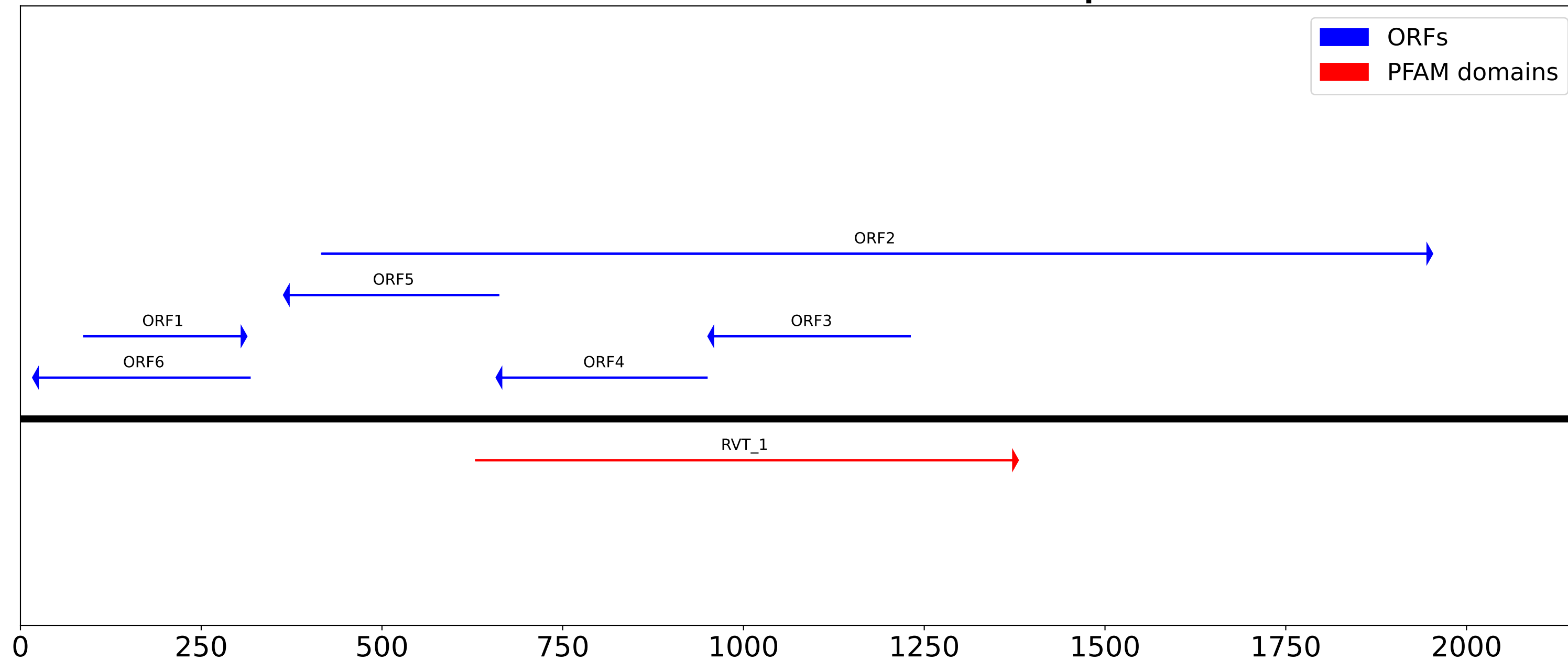


TE: rnd_3_family_436
size: 1654bp; fragments: 1004; full length: 0 (≥ 1488.6 bp)

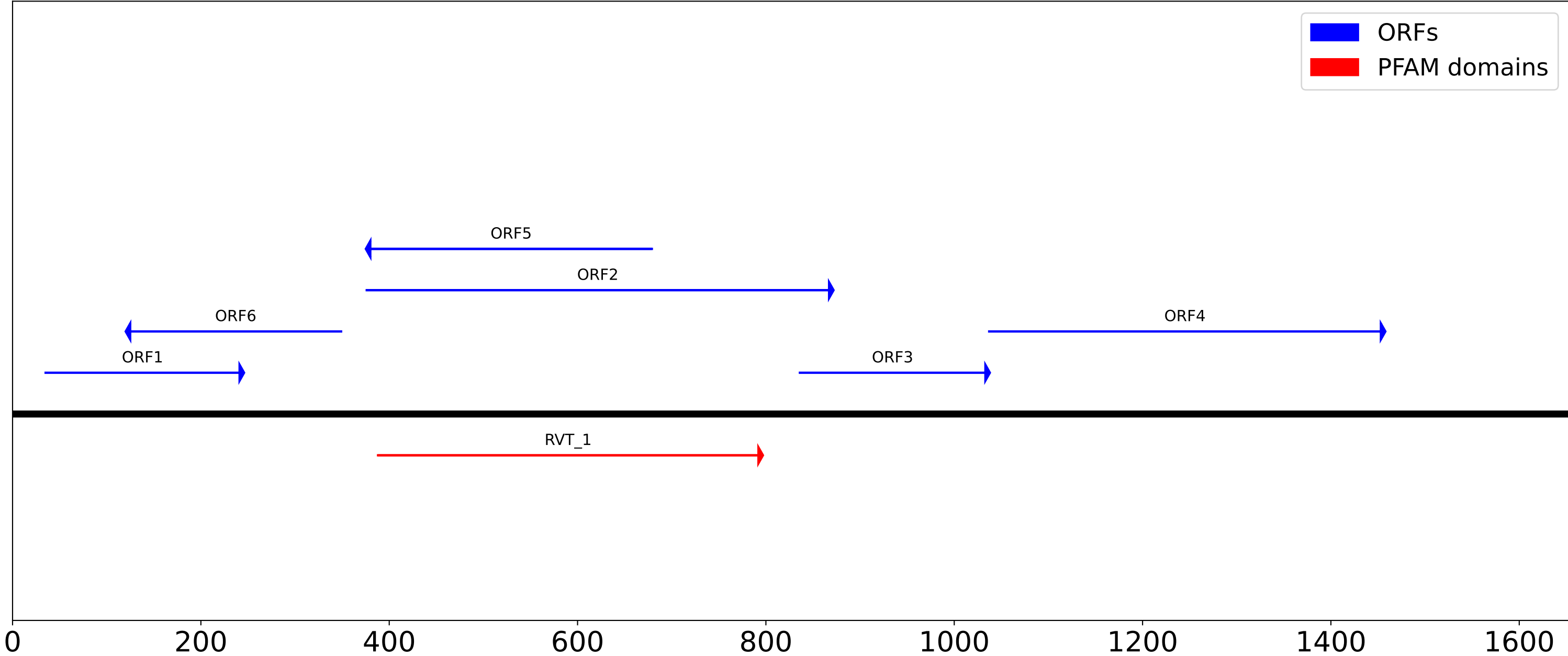
Before TEtrimmer 1654 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

