

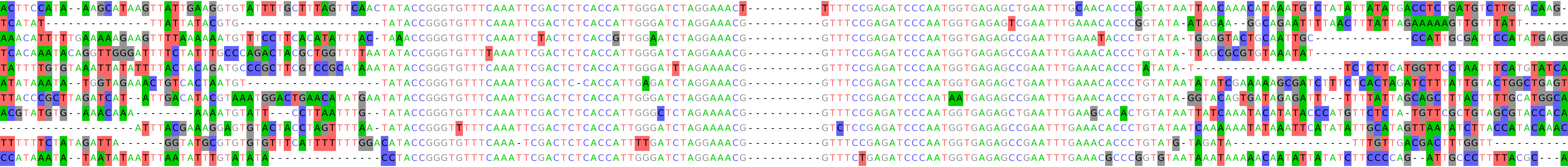
Start crop Point



End crop Point

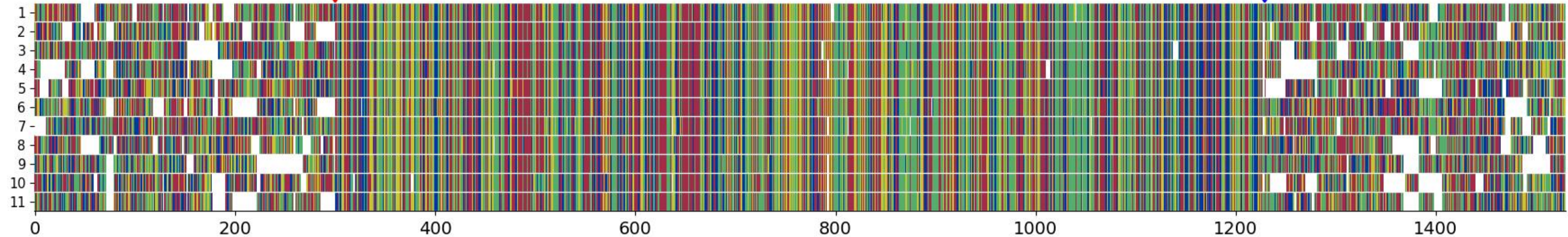


MSA length = 930



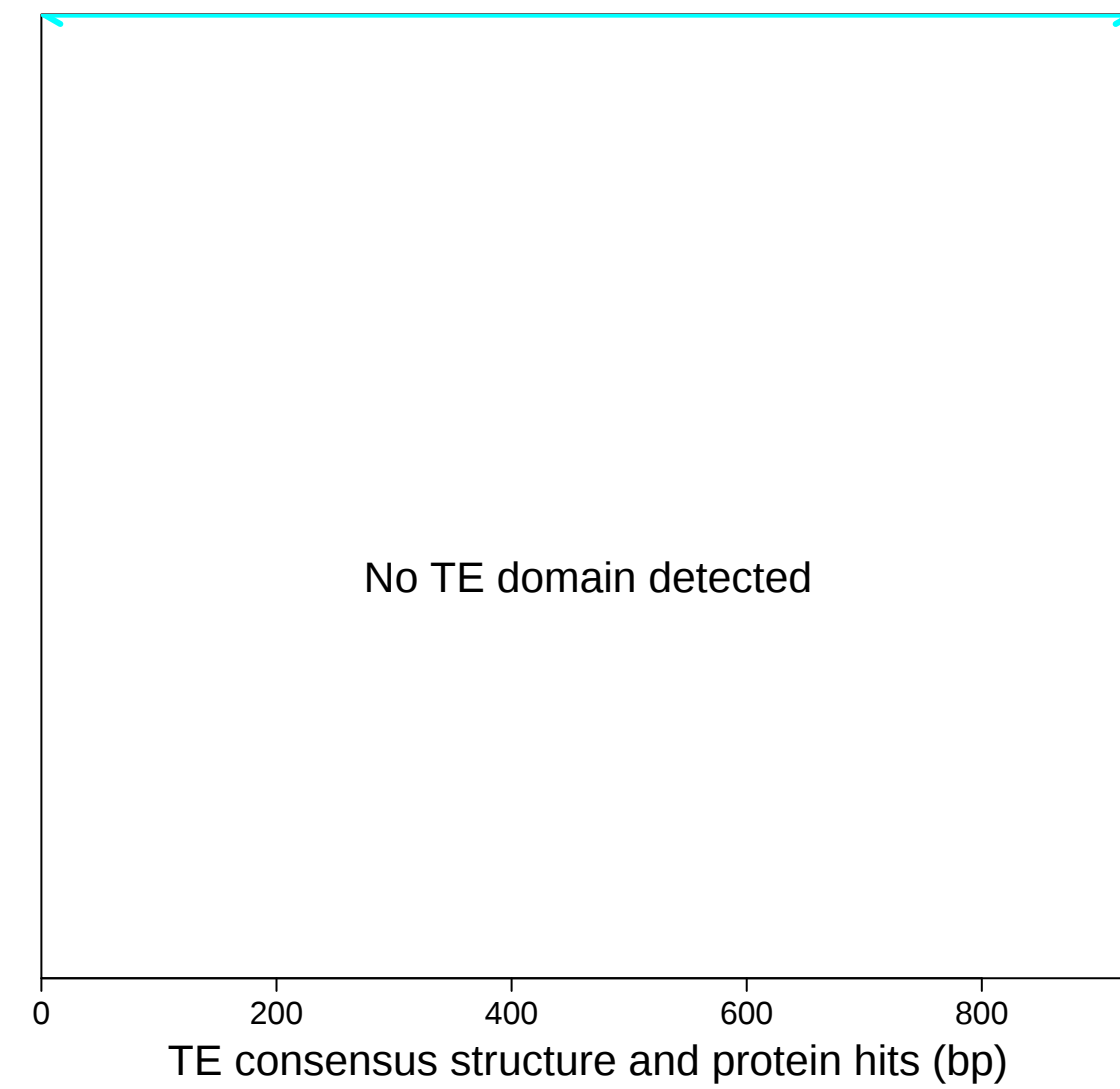
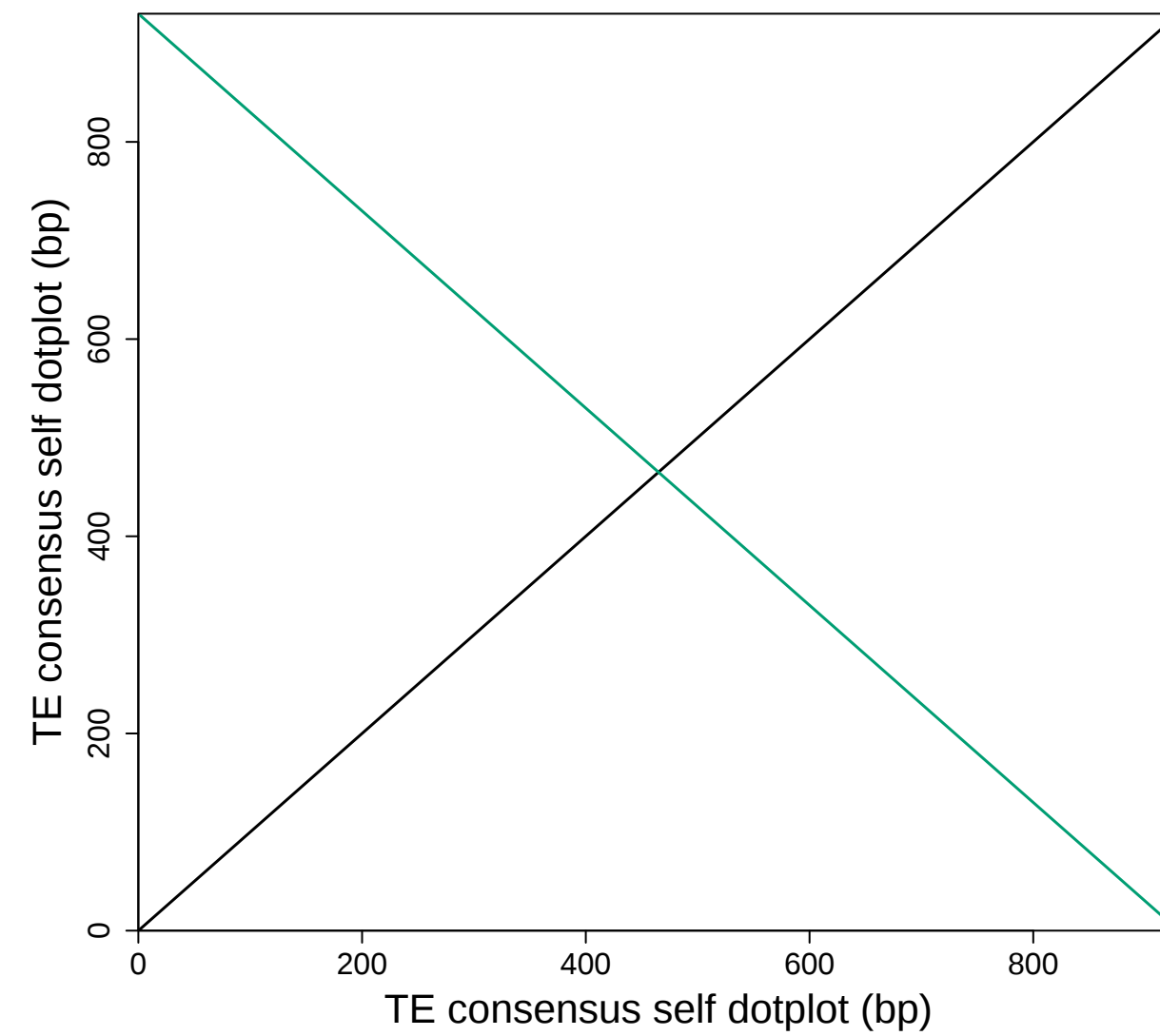
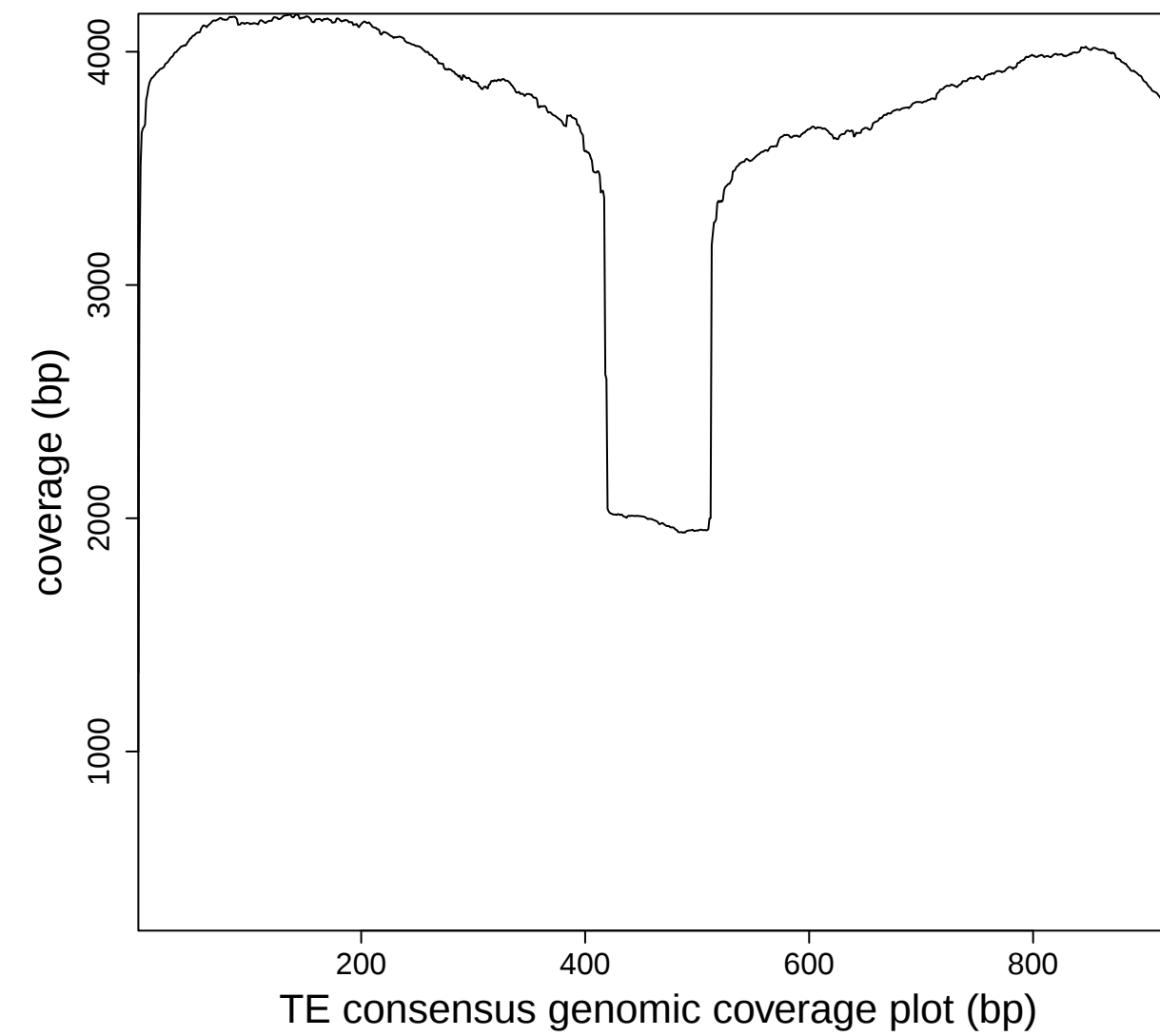
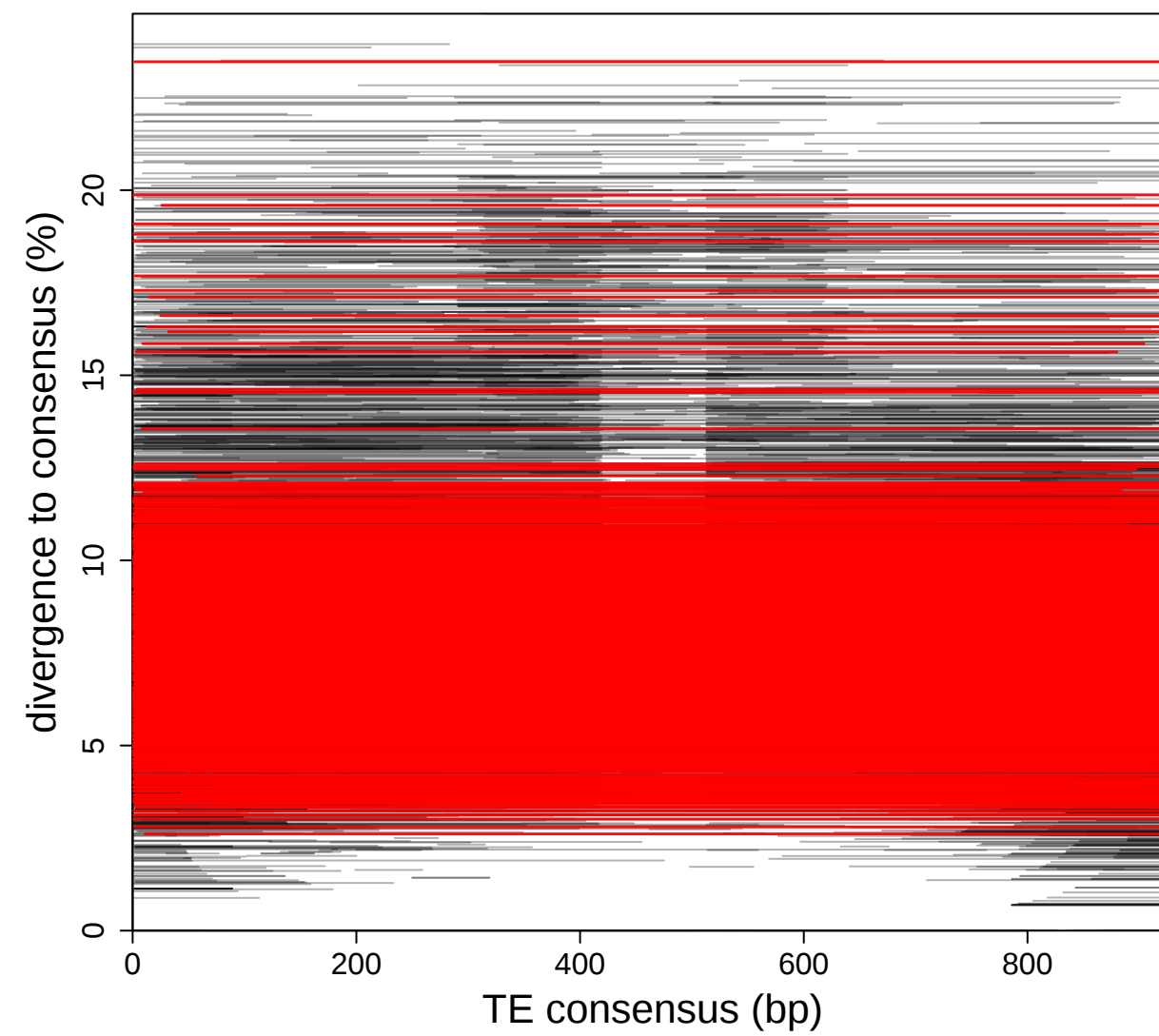
Start crop Point

End crop Point



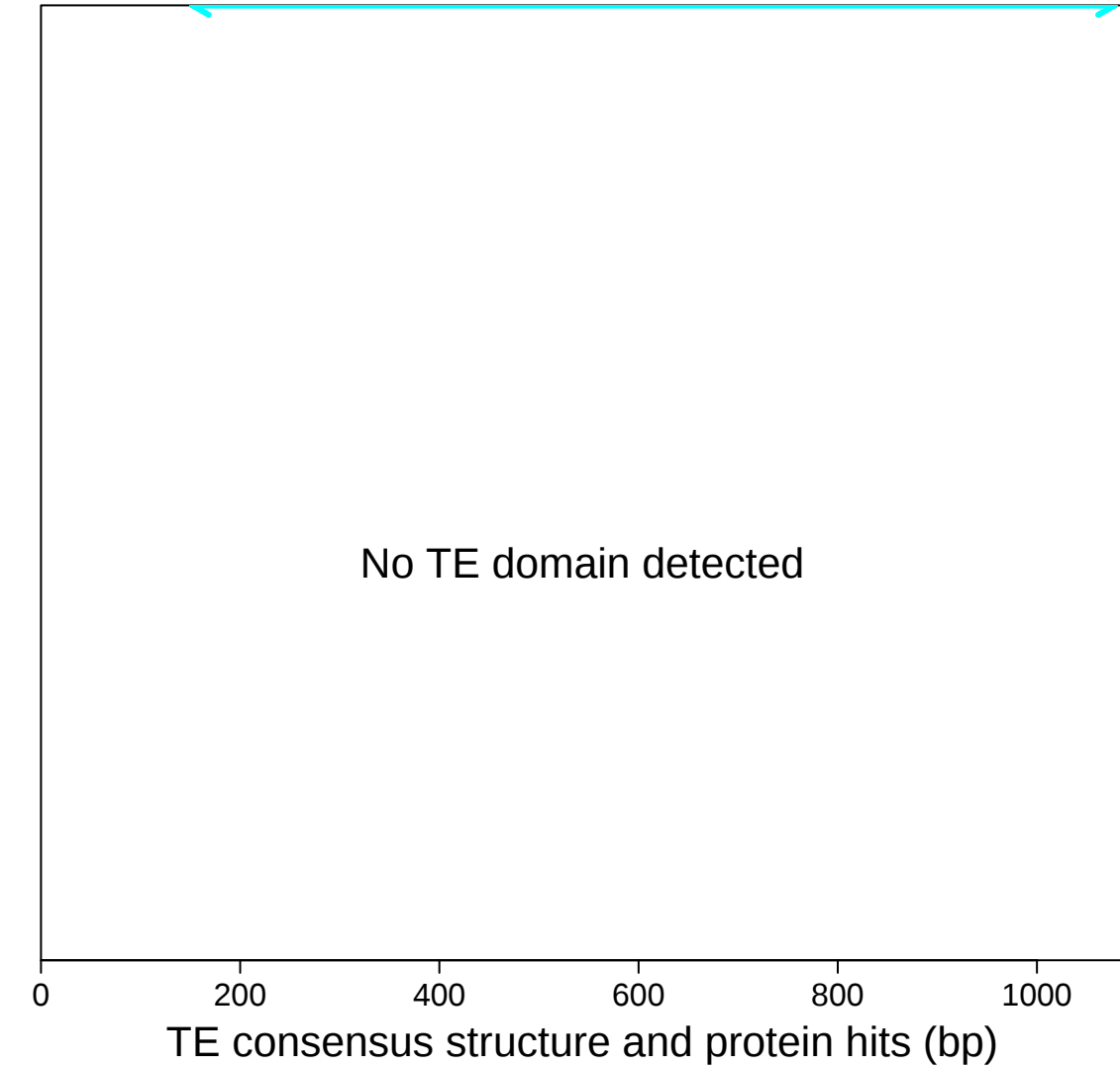
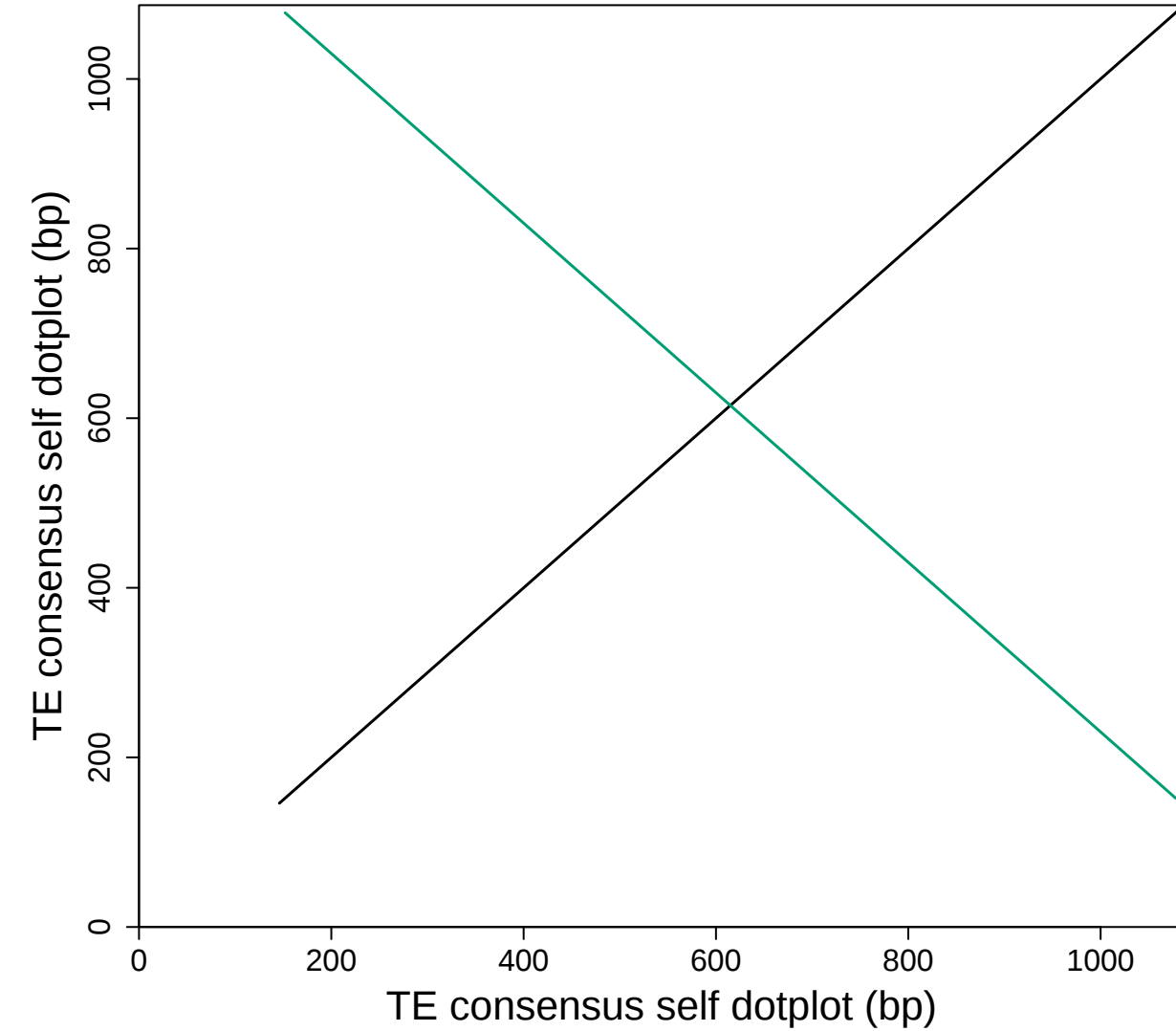
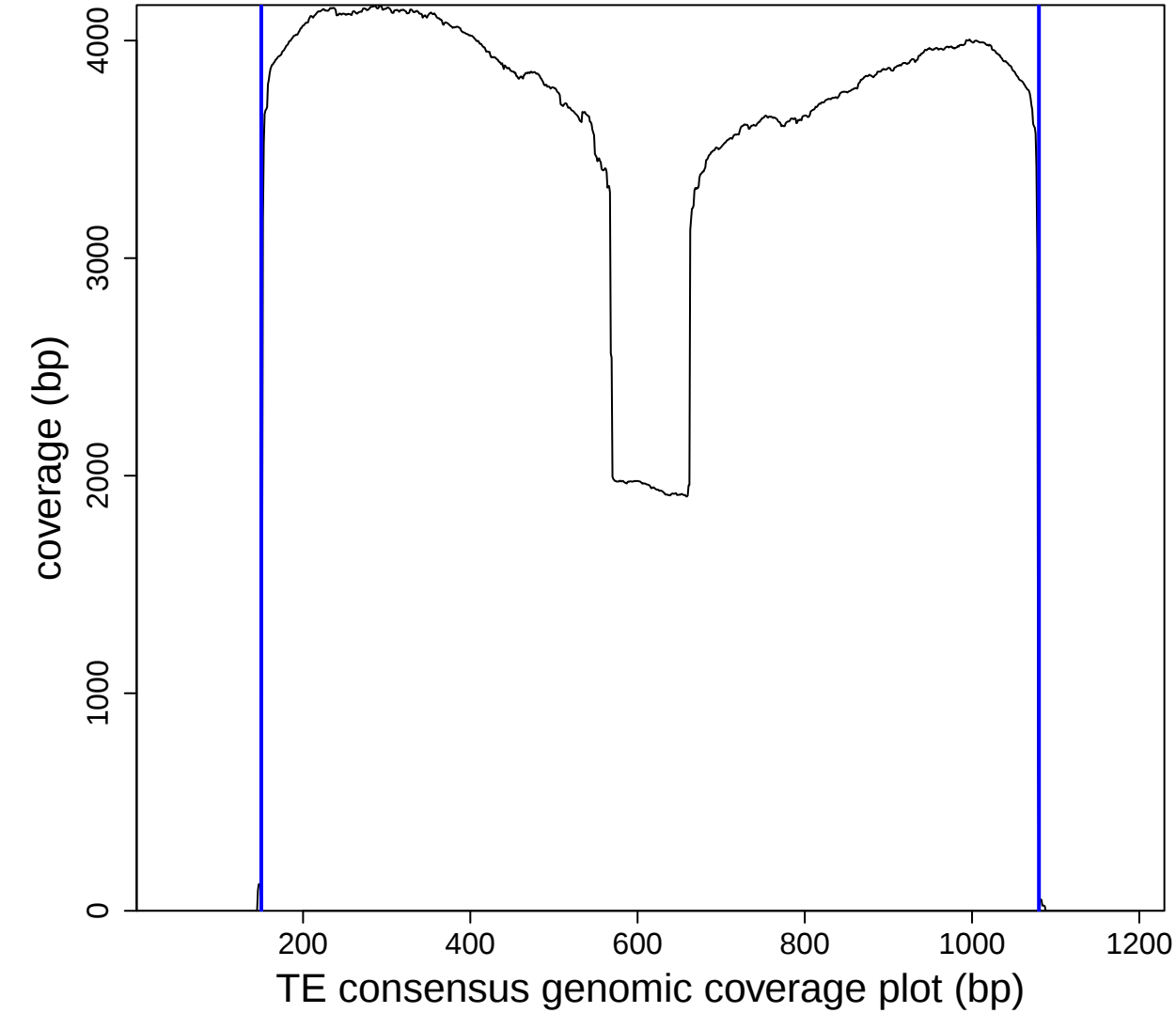
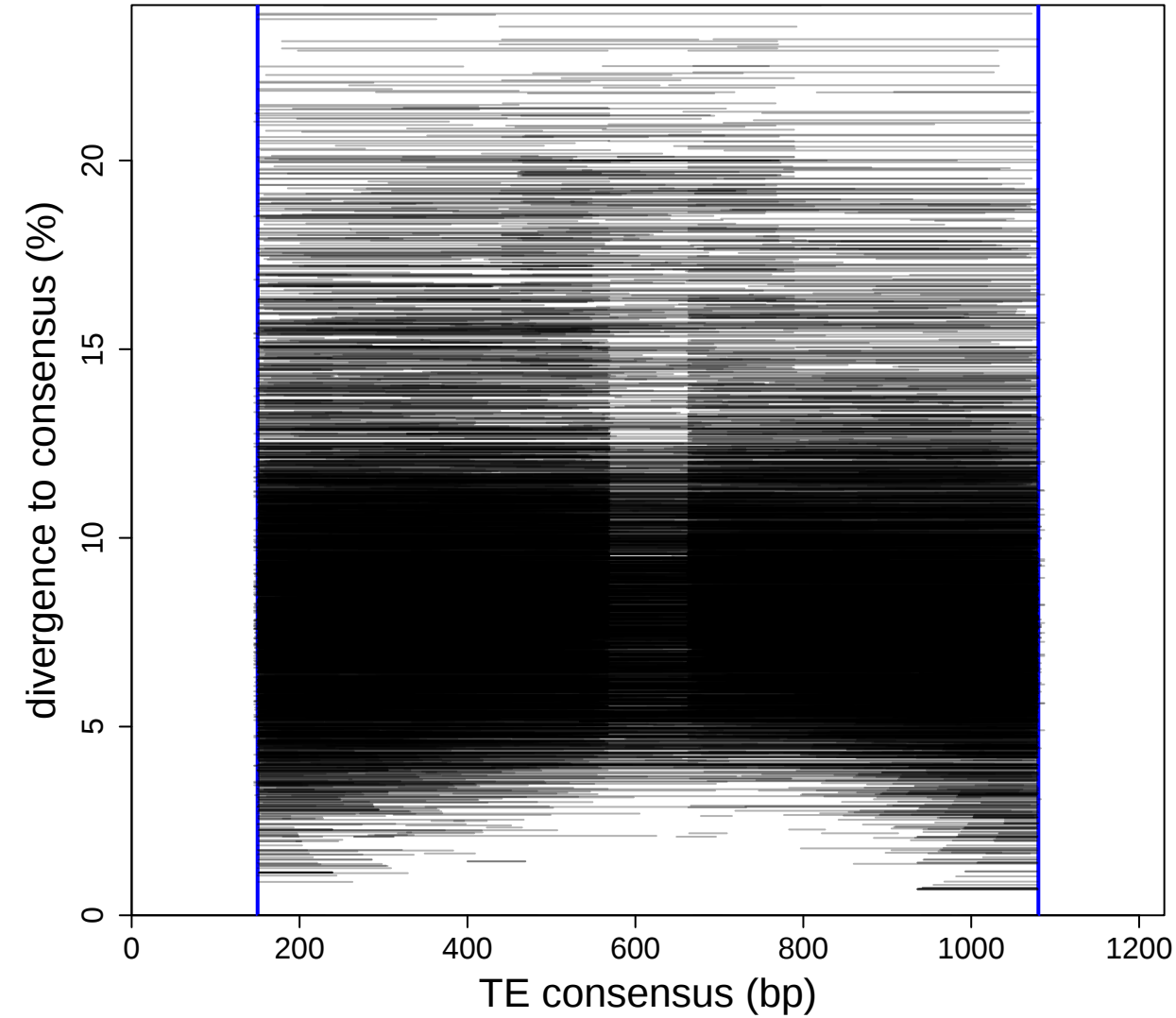
l.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bc\n.fa_aln.fa_cl.fa_gs.fa
size: 930bp; fragments: 8282; full length: 1057 (>=837bp)

After TEtrimmer 930 bp

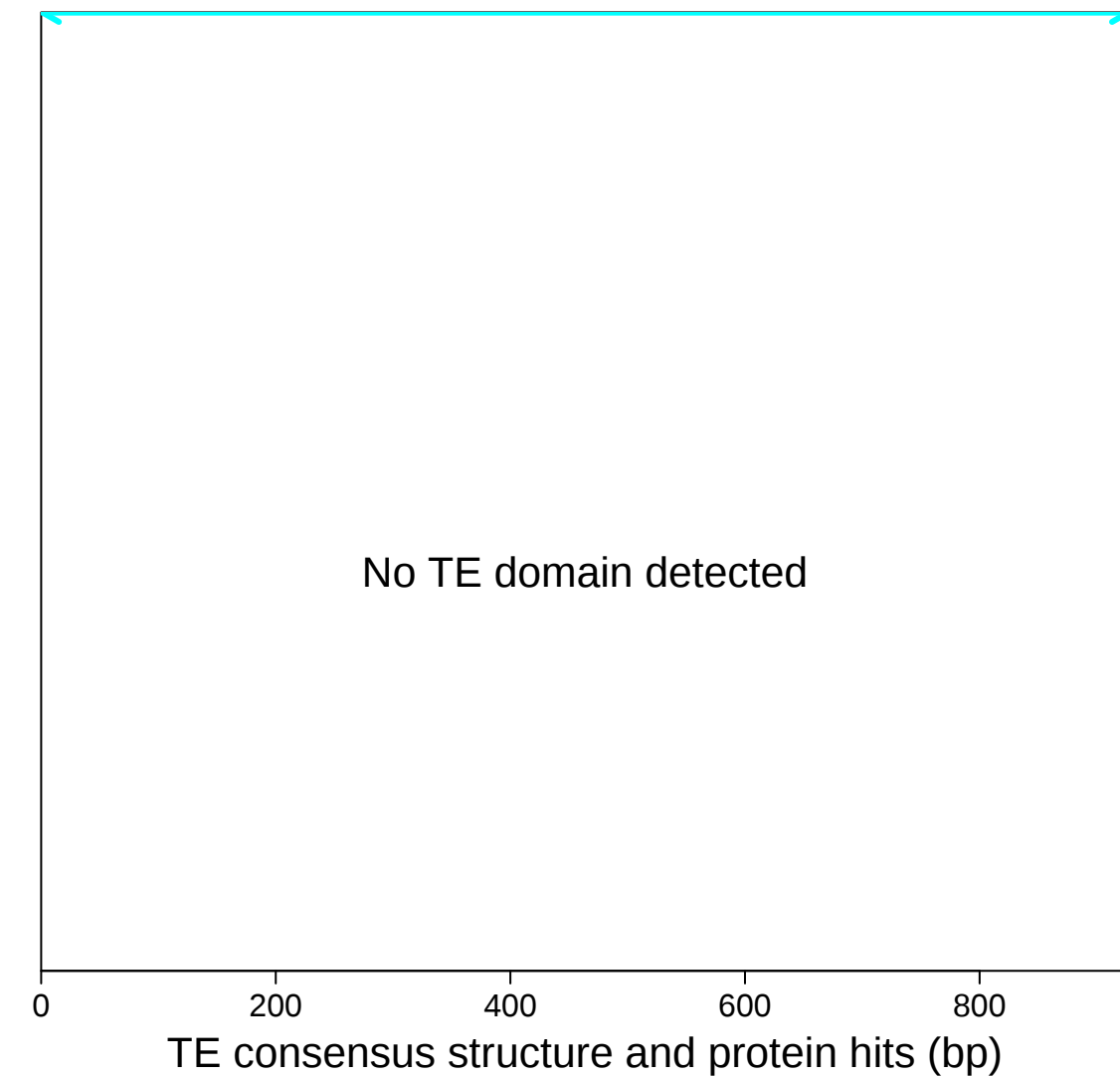
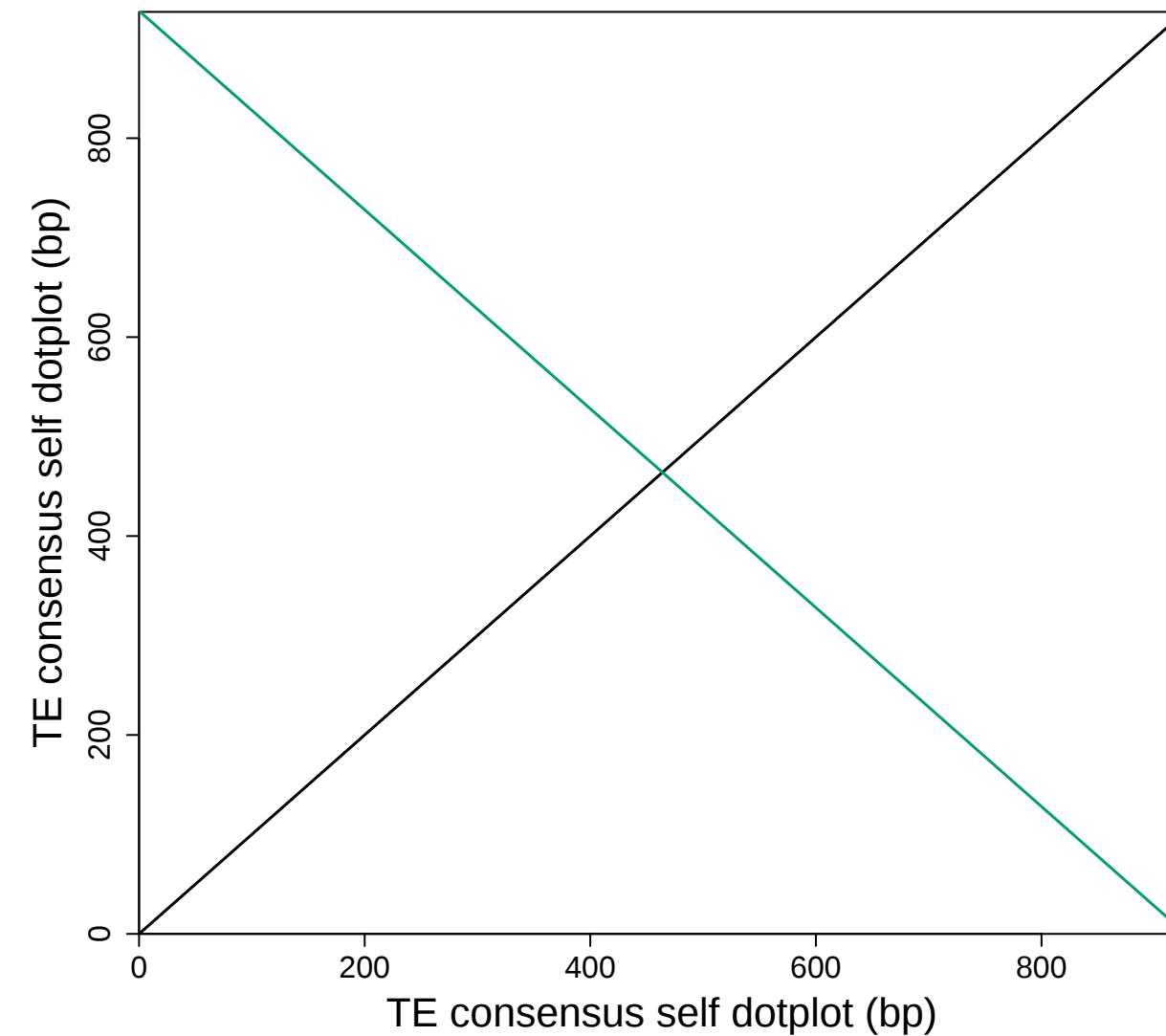
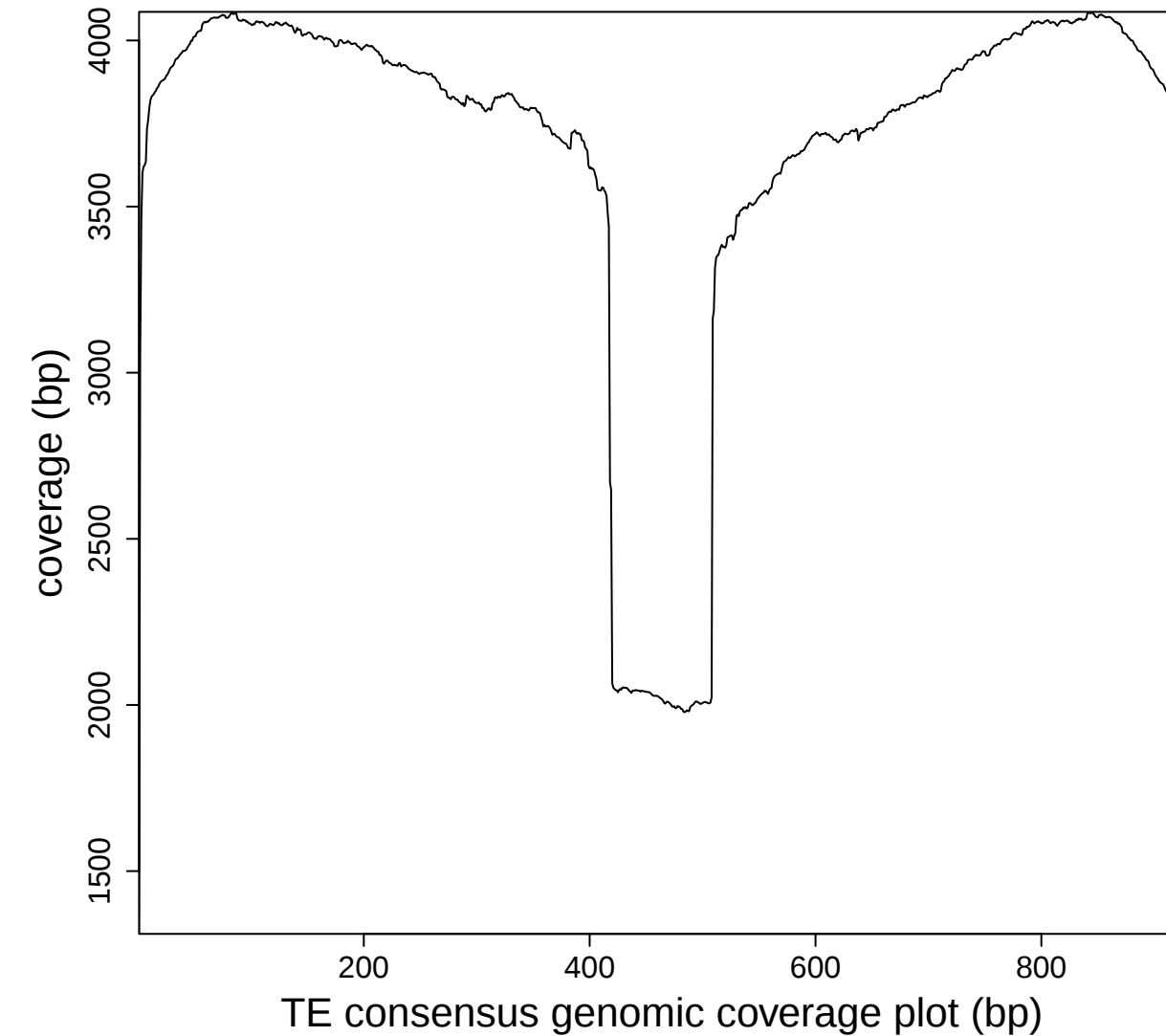
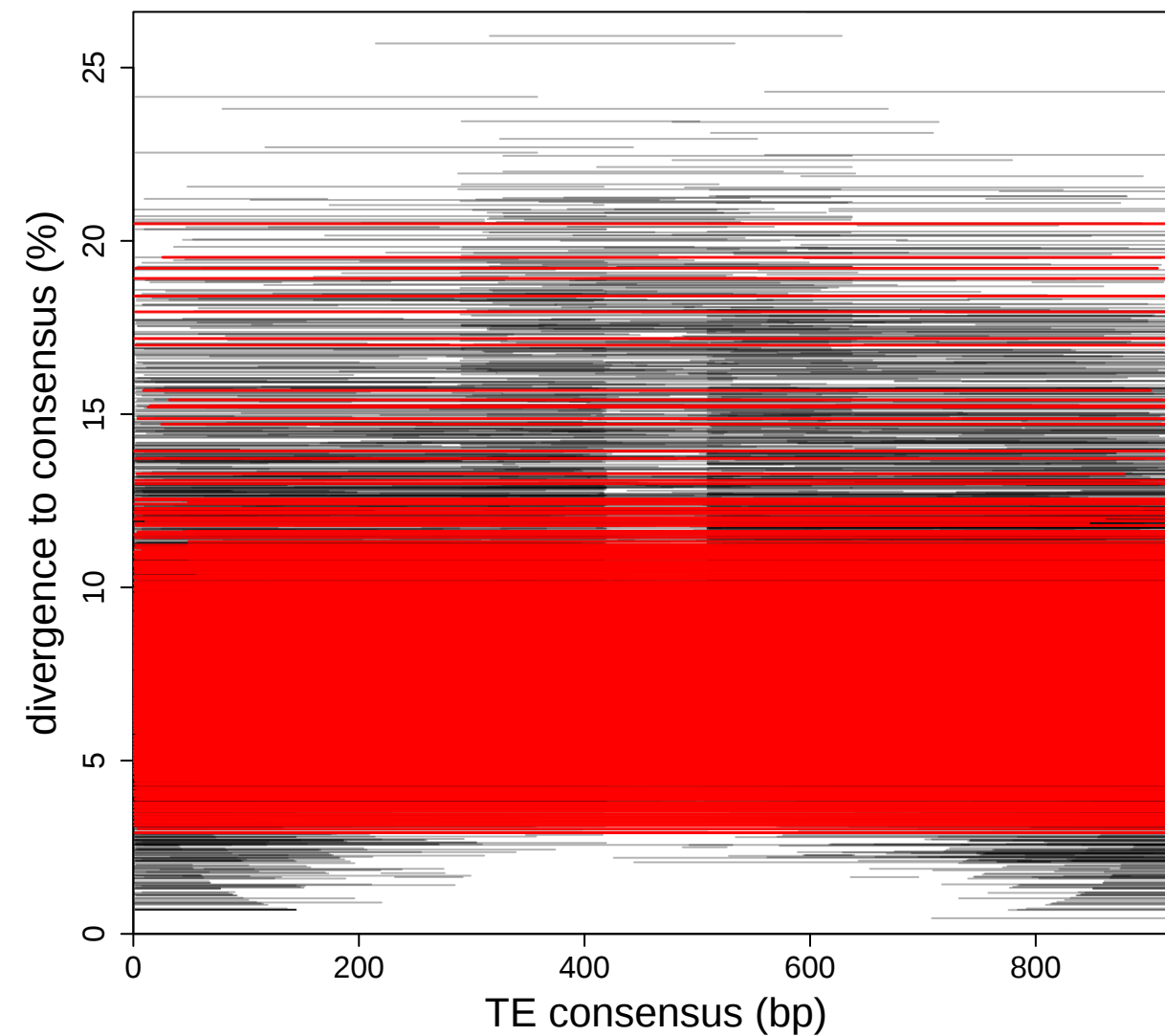


6.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1230bp; fragments: 8206; full length: 0 (>=1107bp)

After TEtrimmer Extended plot Blue lines are boundaries

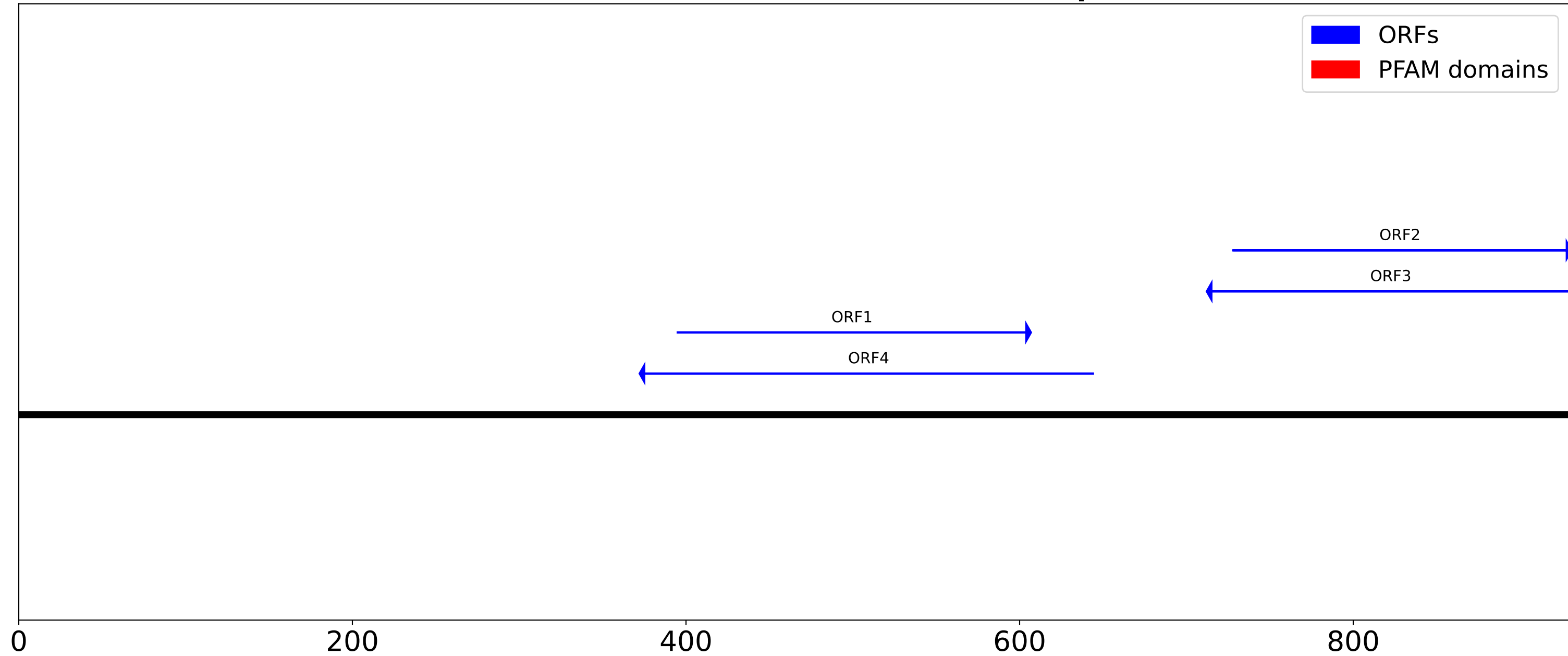


TE: rnd_1_family_126
size: 927bp; fragments: 8154; full length: 1056 (≥ 834.3 bp)

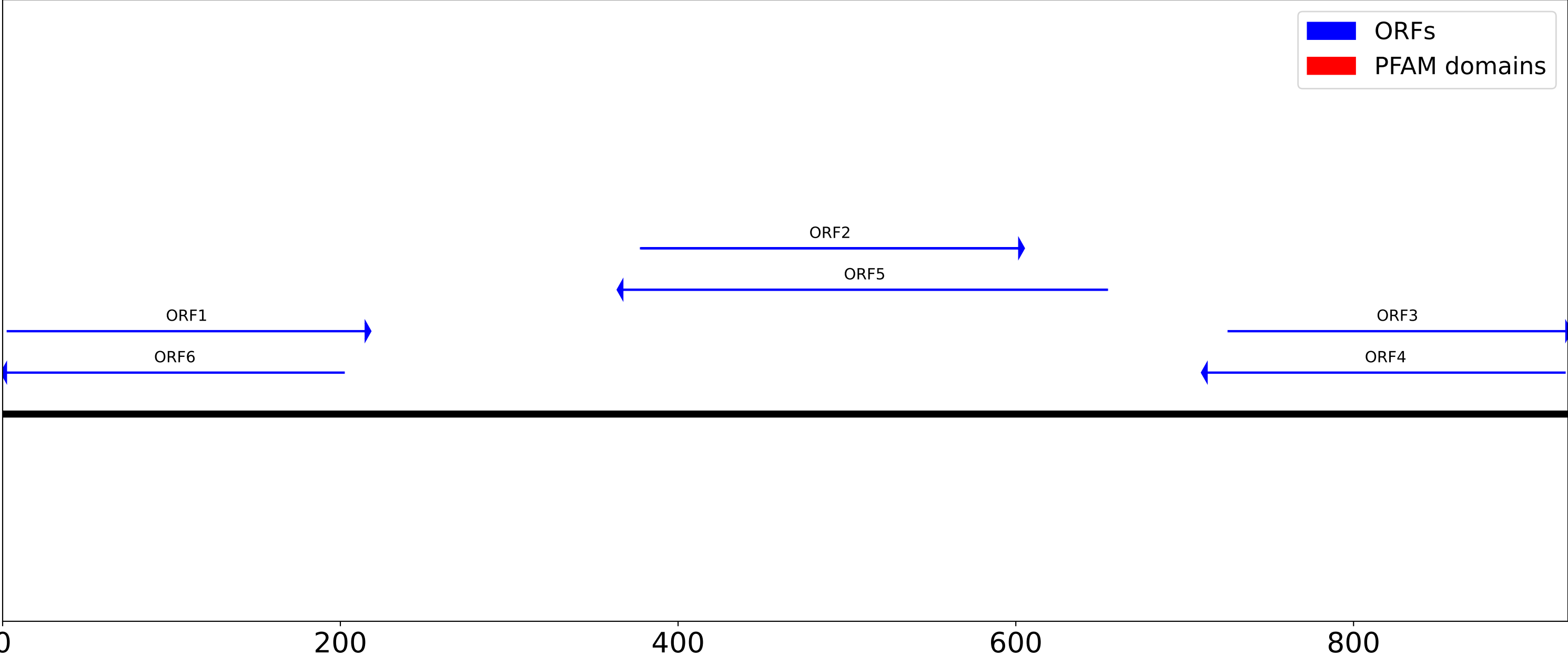


Before TEtrimmer 927 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

