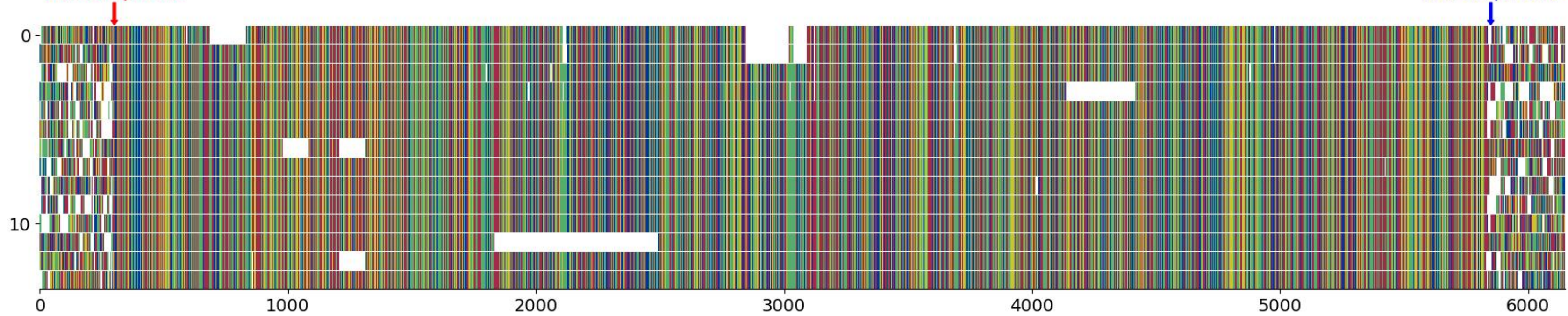


MSA length = 5551



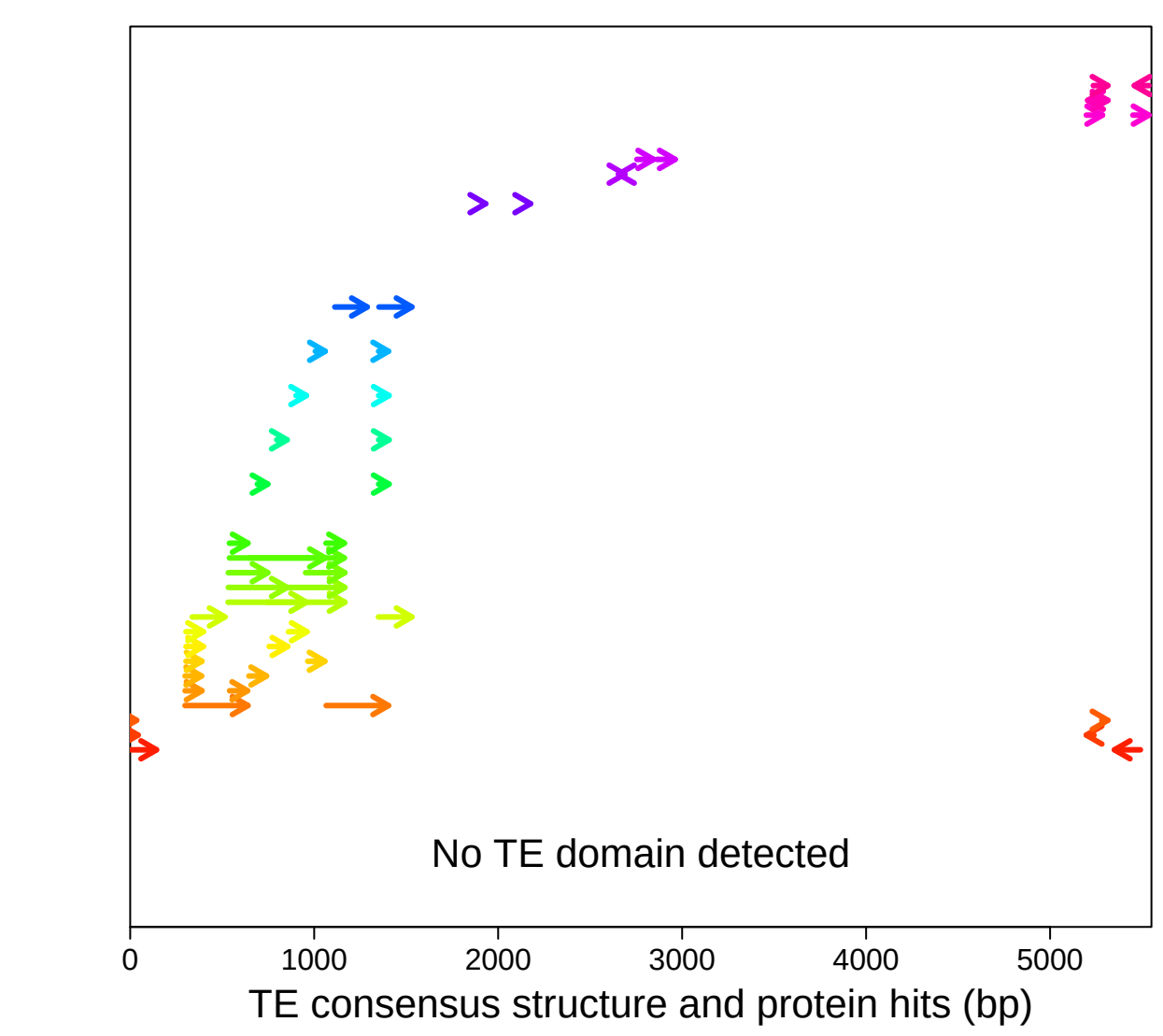
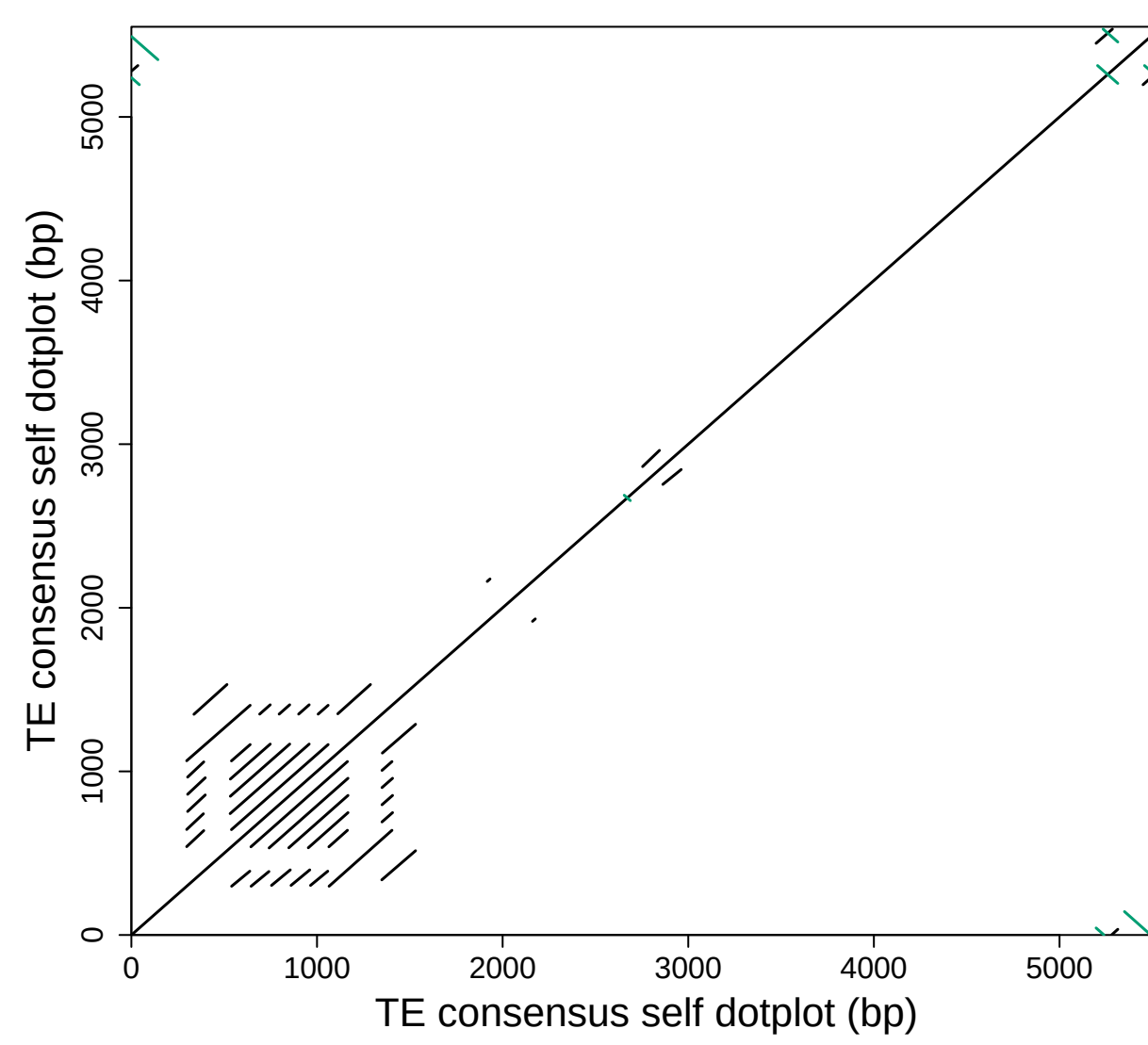
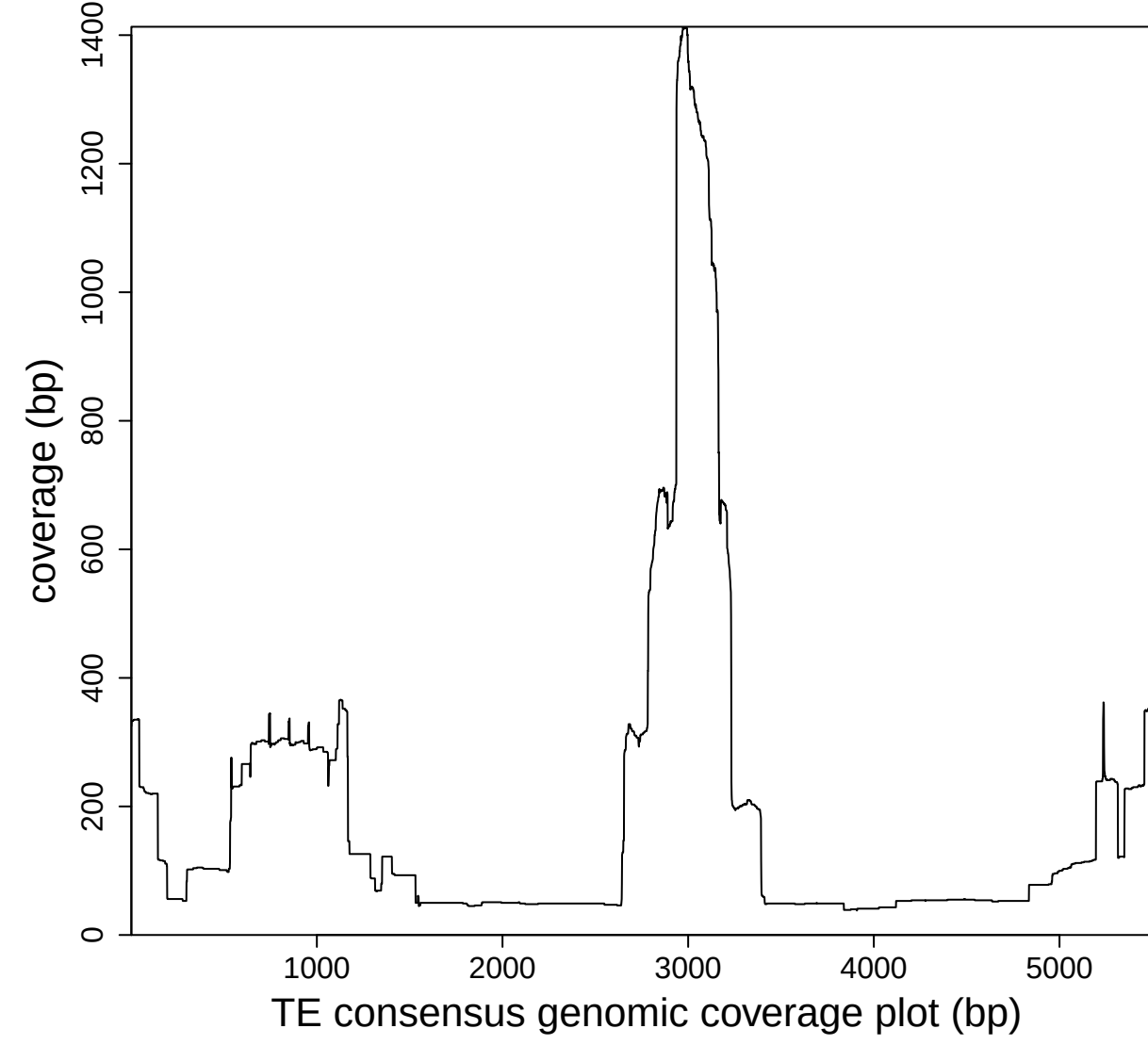
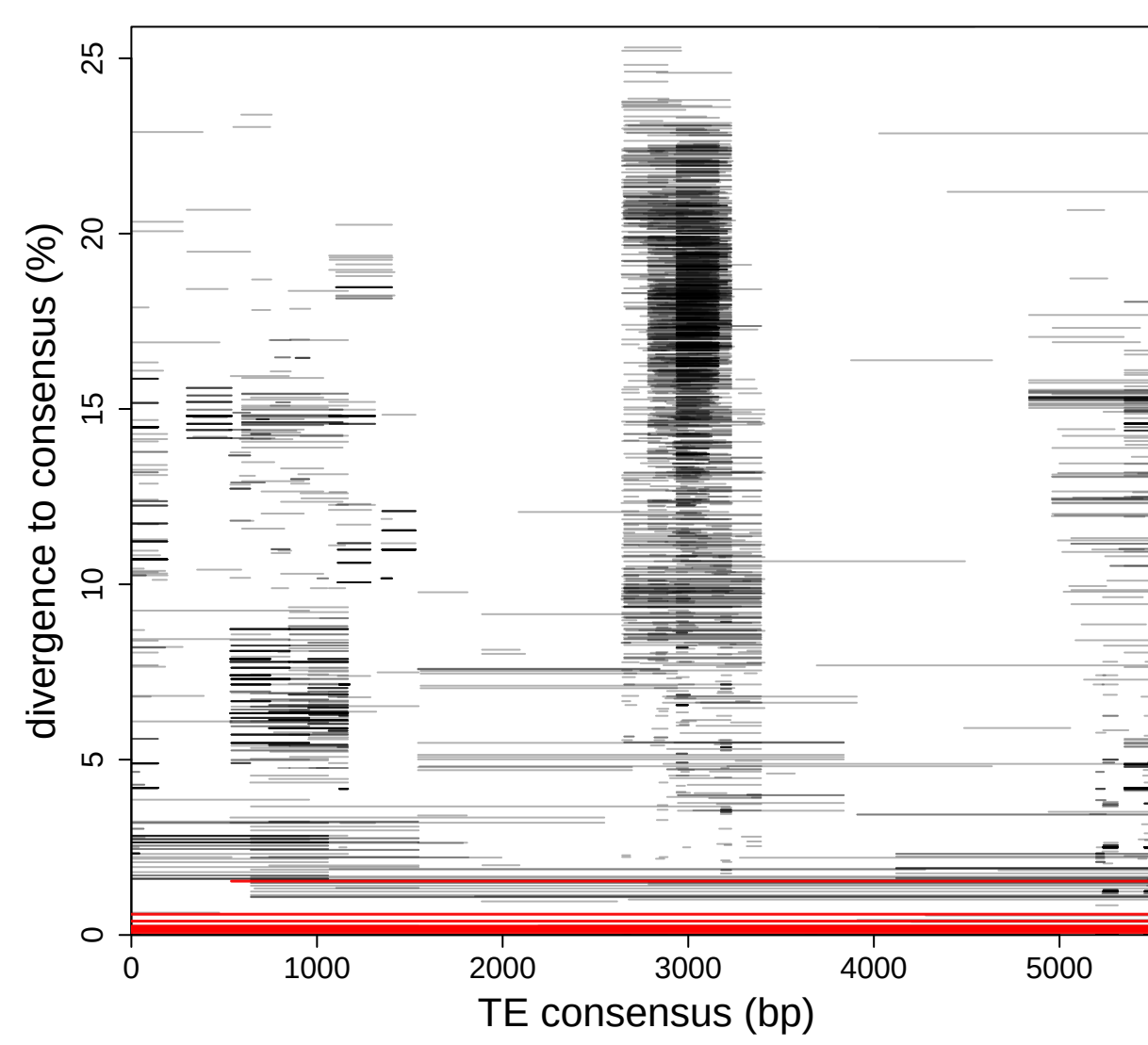
Start crop Point

End crop Point



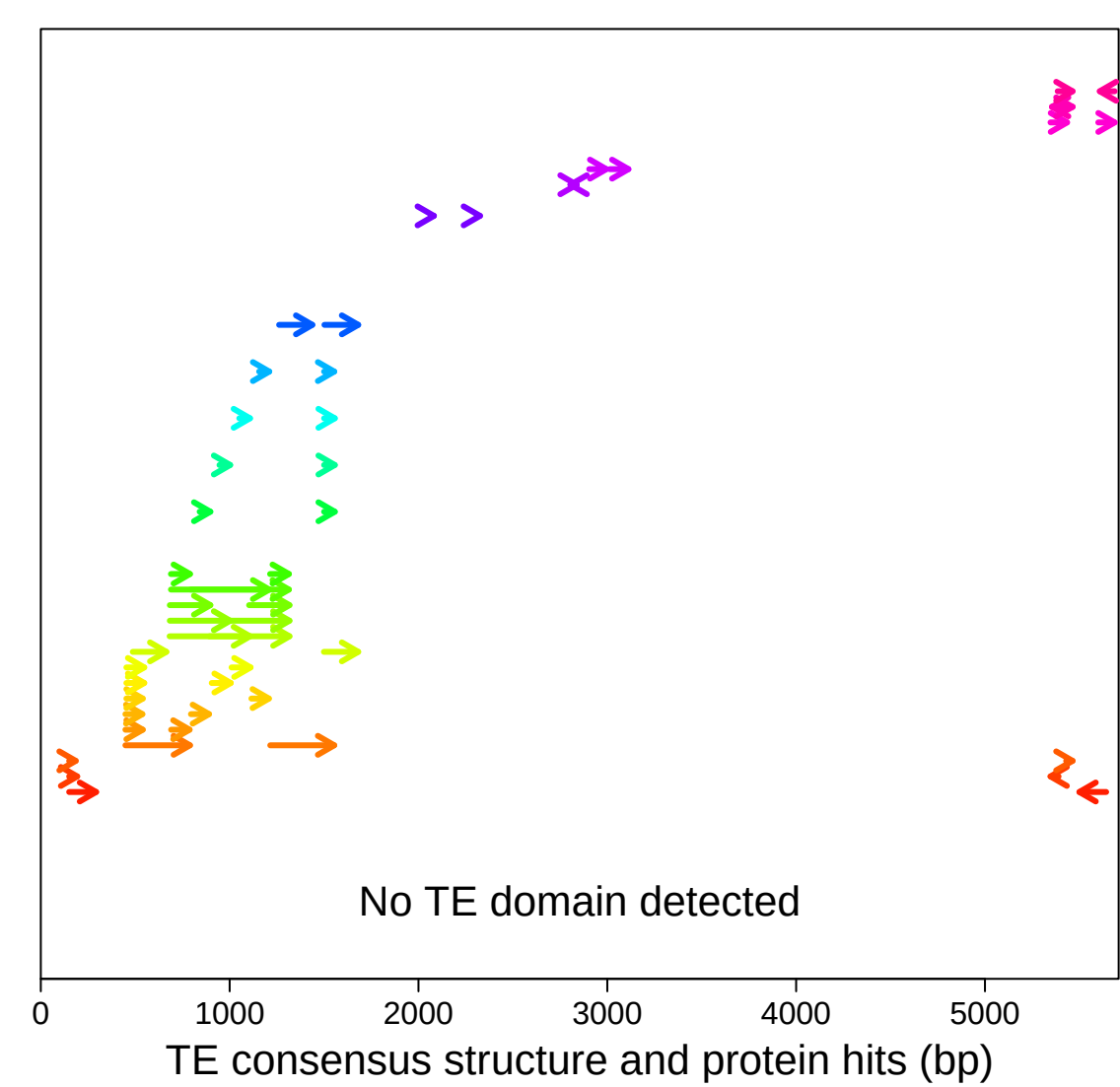
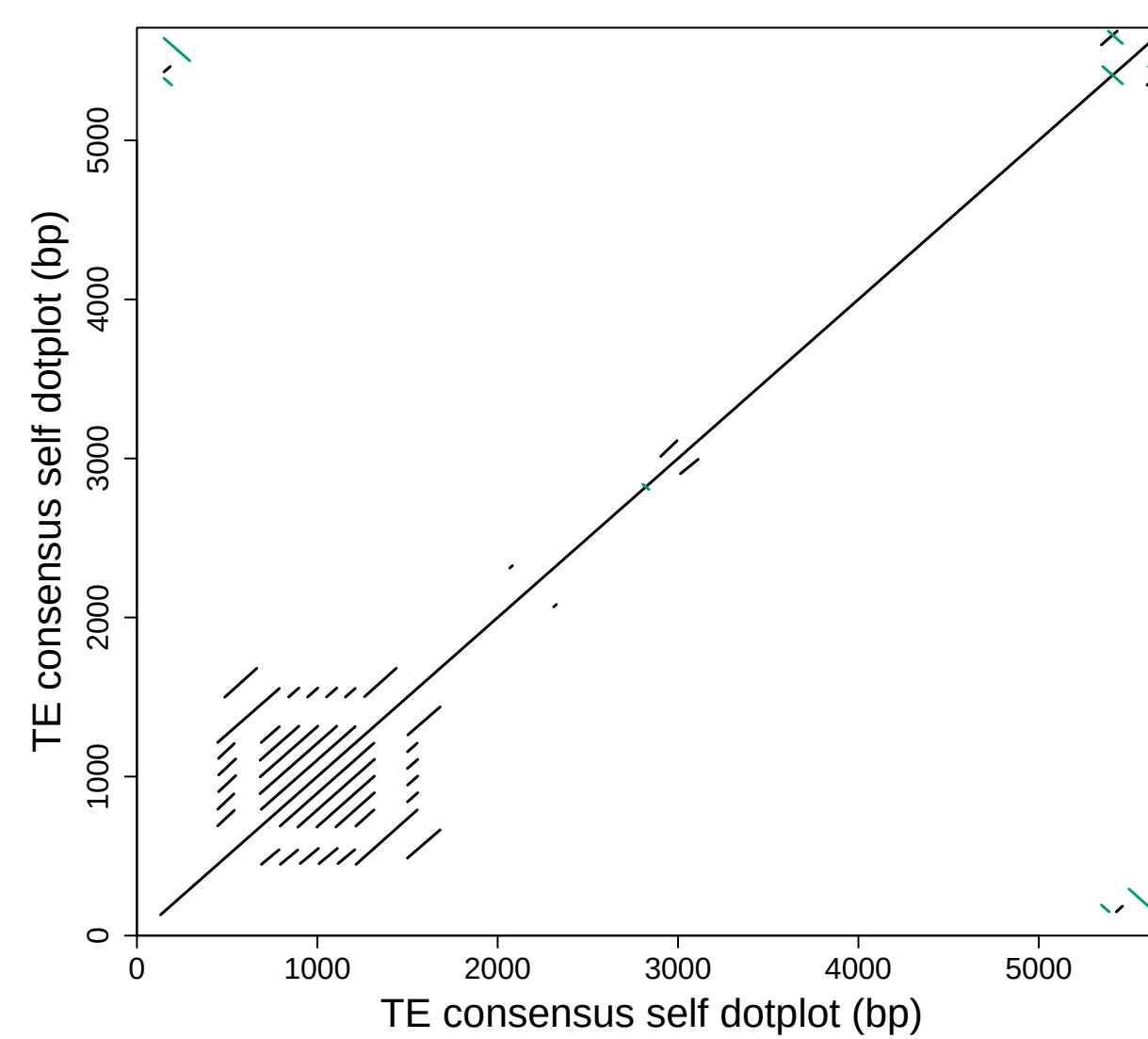
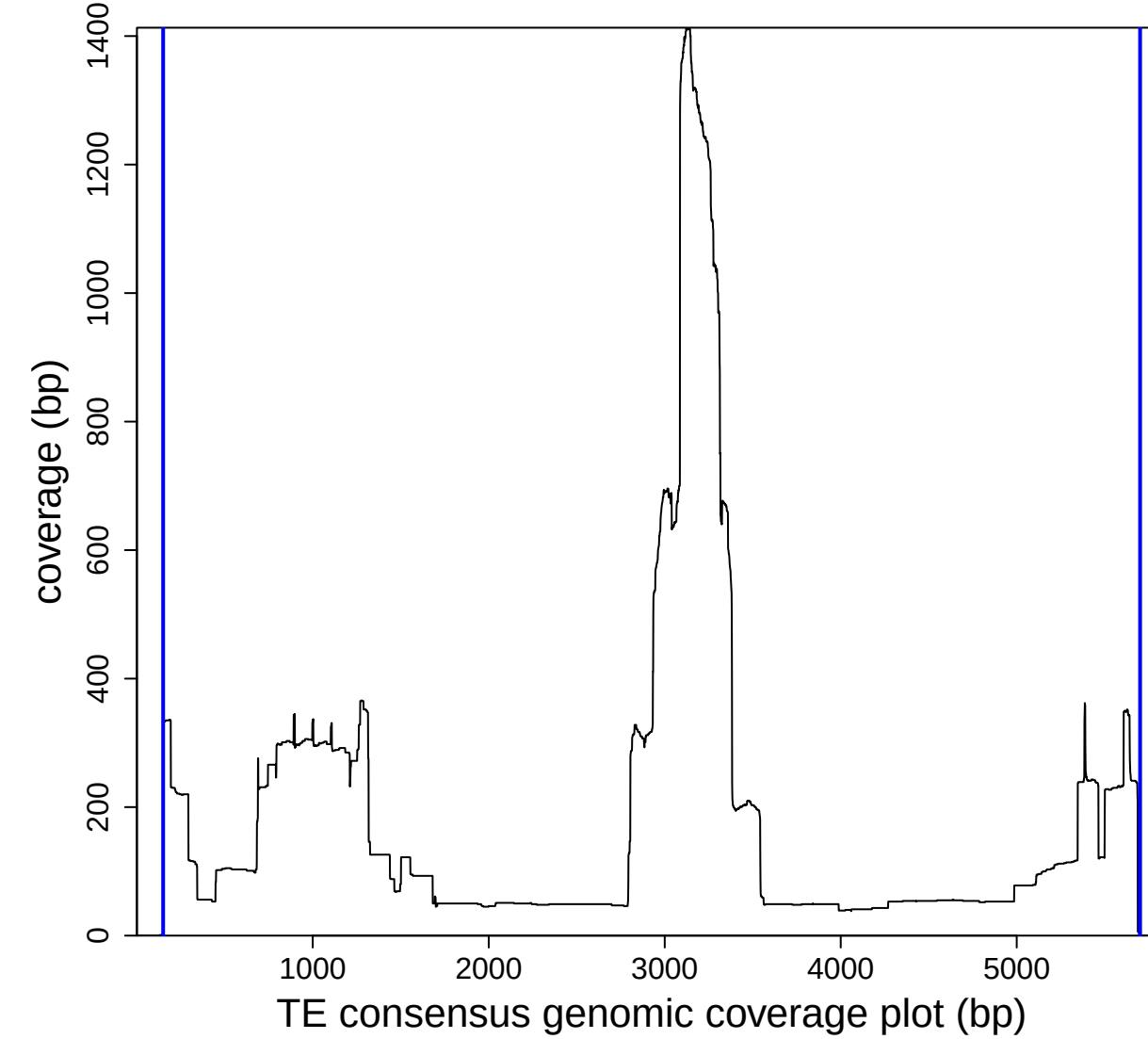
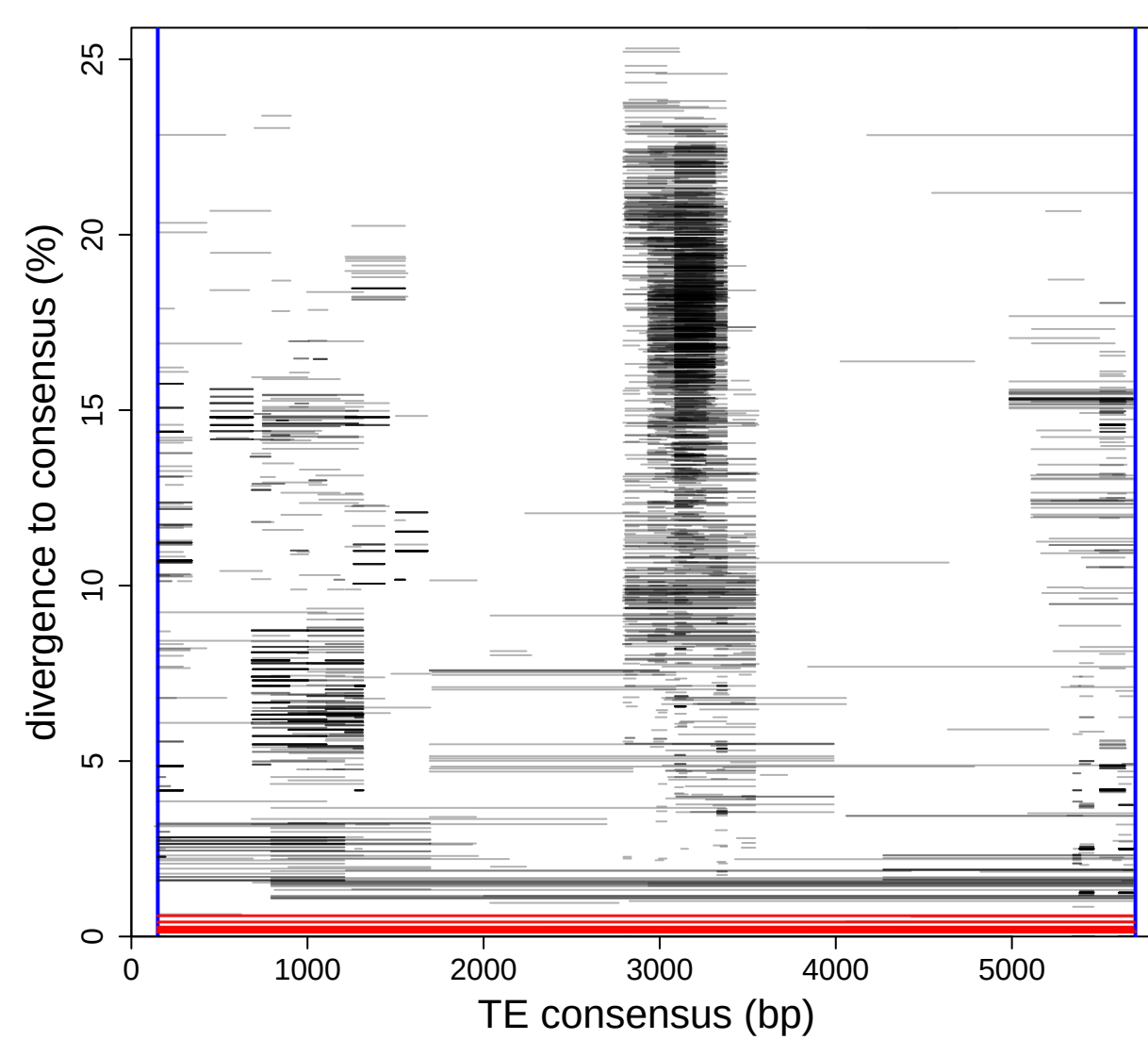
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 5551bp; fragments: 3386; full length: 10 (>=4995.9bp)

After TEtrimmer 5551 bp



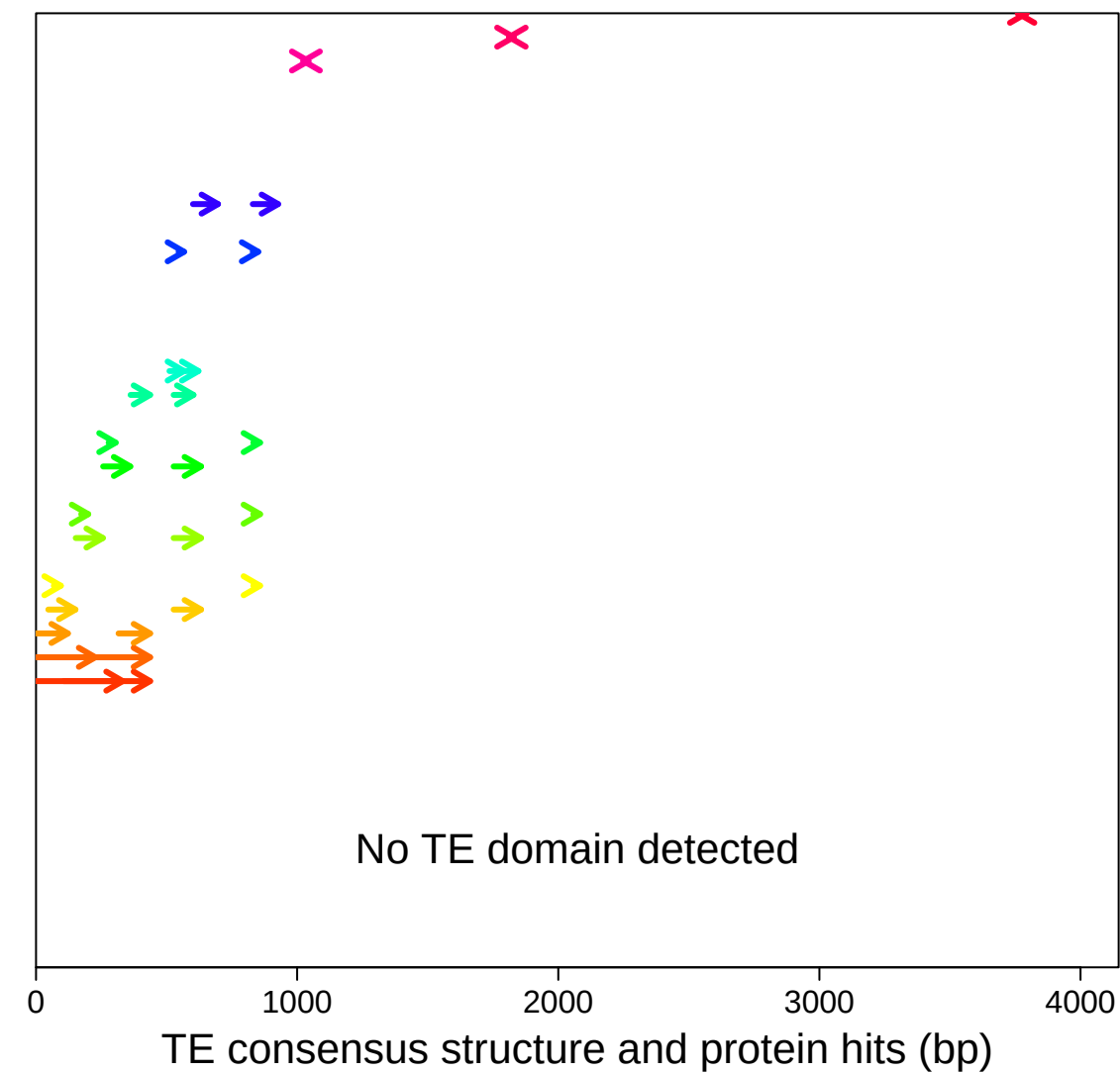
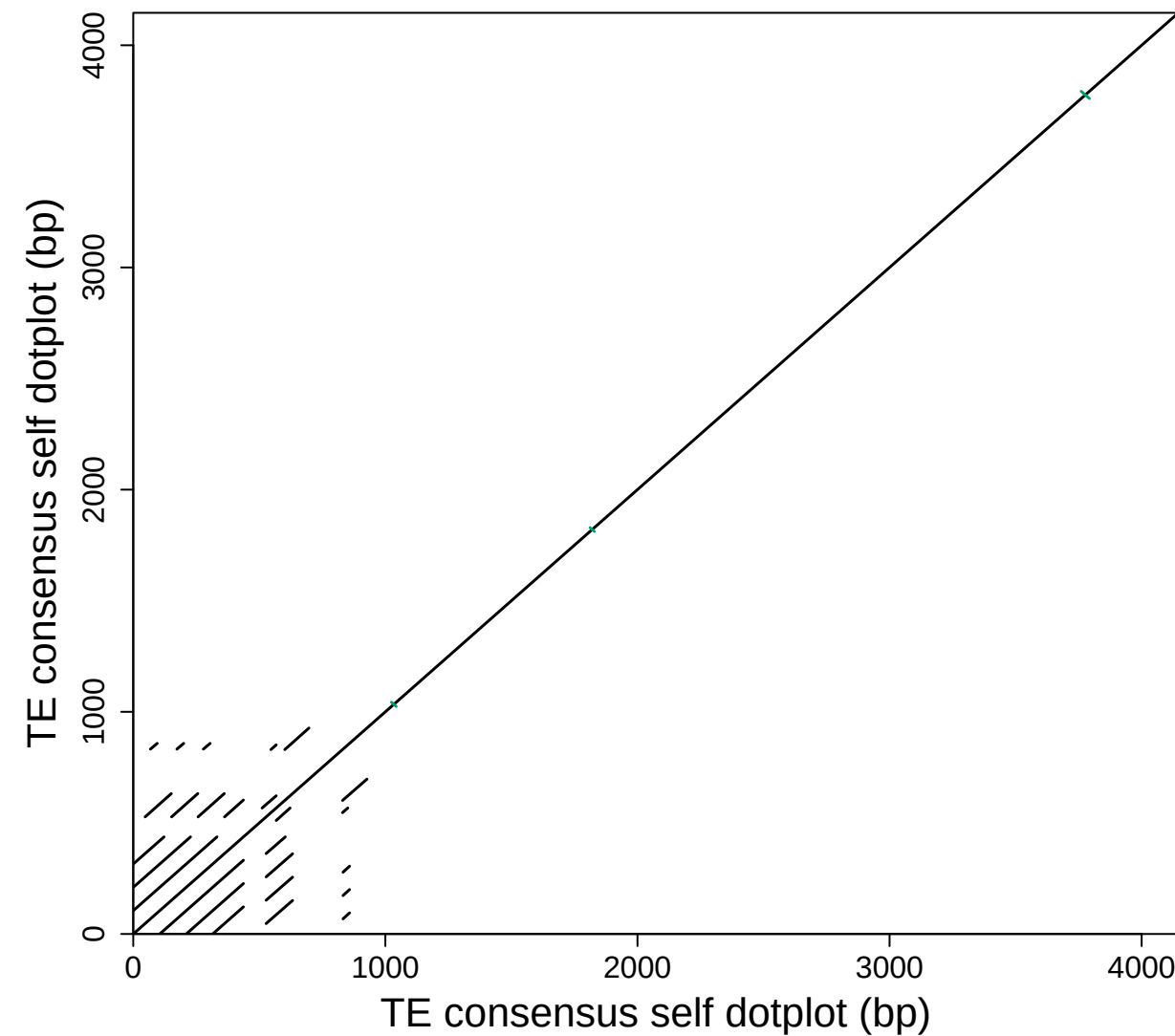
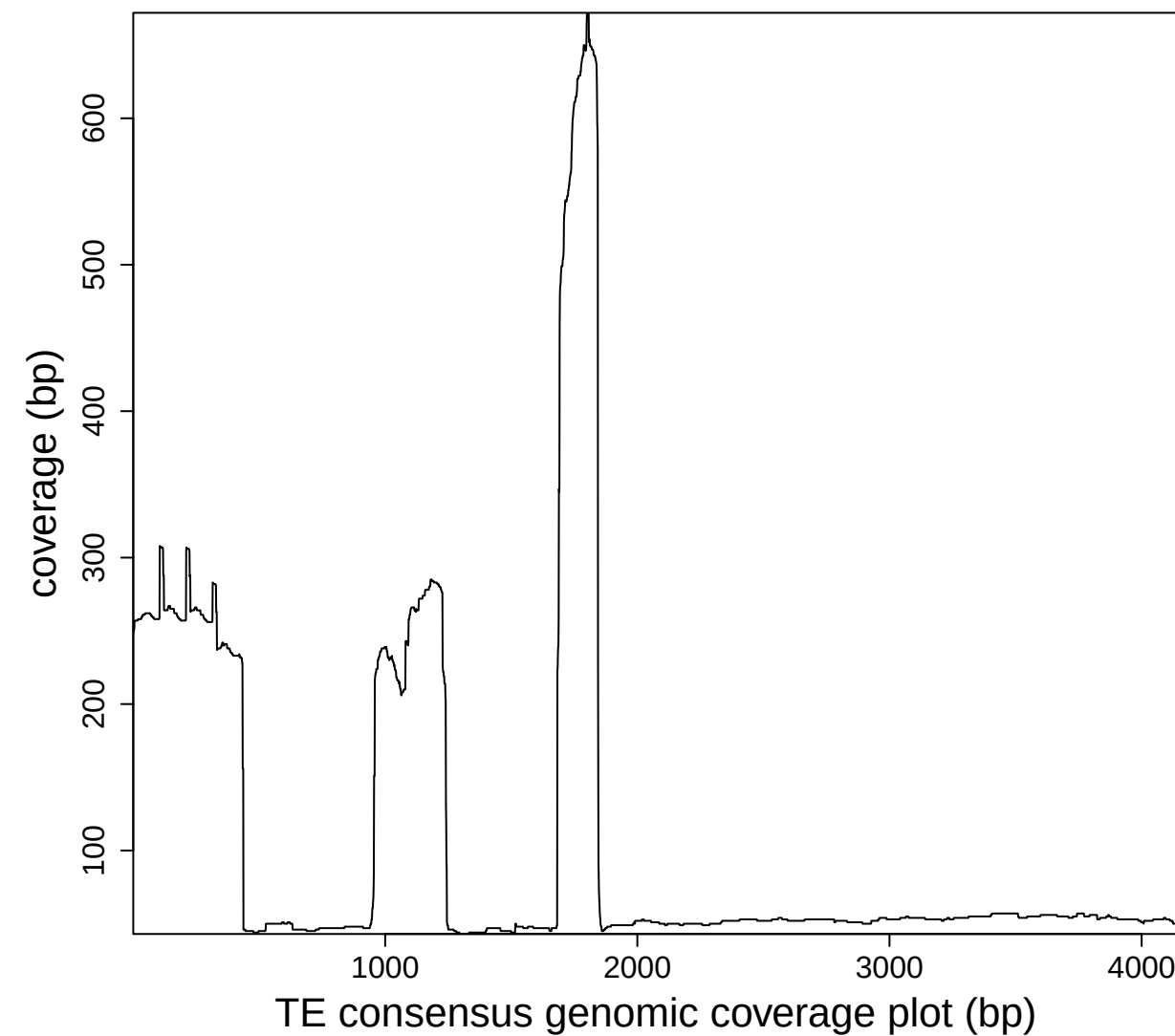
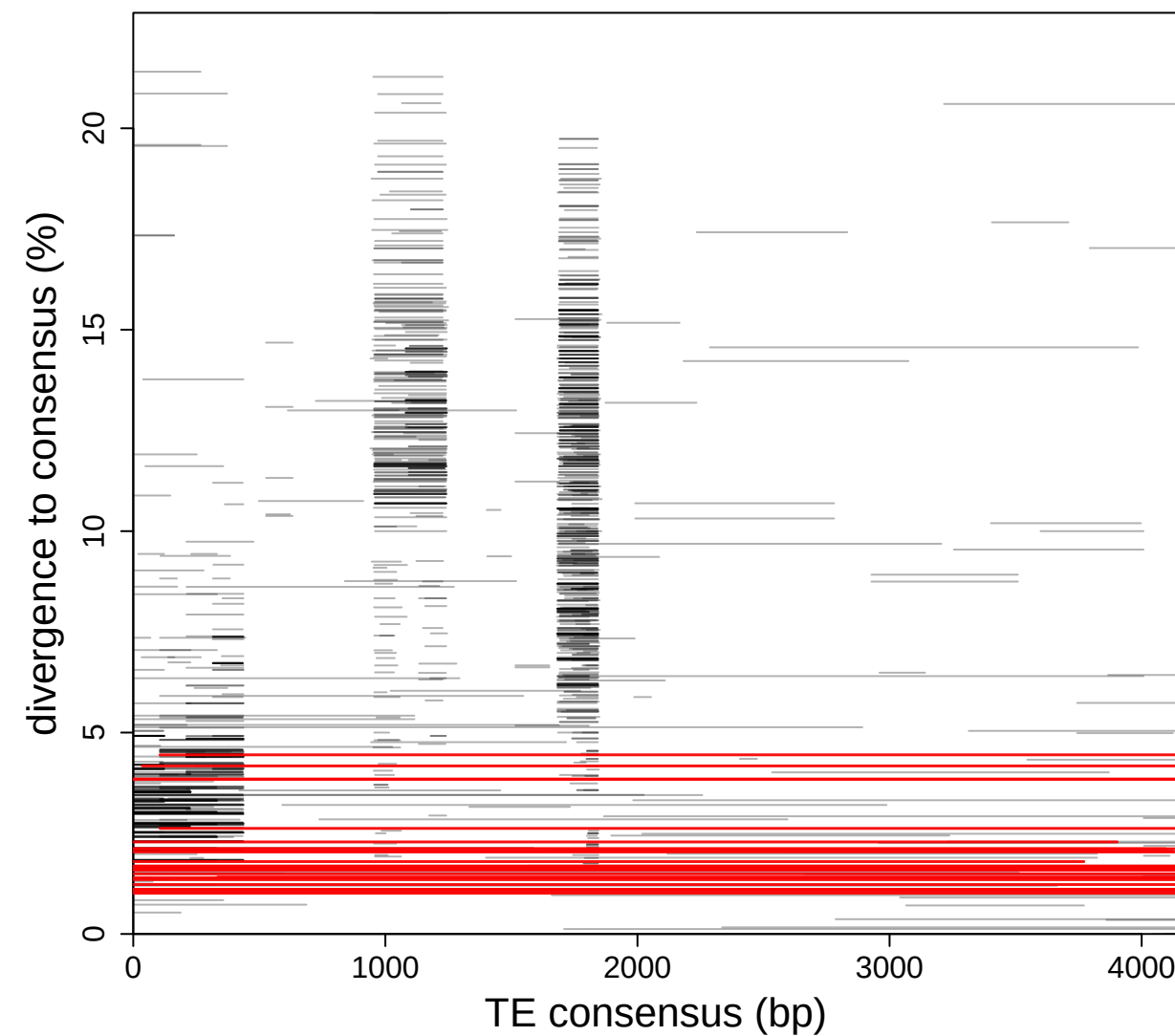
2.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcIn.fa aln.fa cl.f
size: 5851bp; fragments: 3386; full length: 9 (≥ 5265.9 bp)

After TEtrimmer Extended plot Blue lines are boundaries

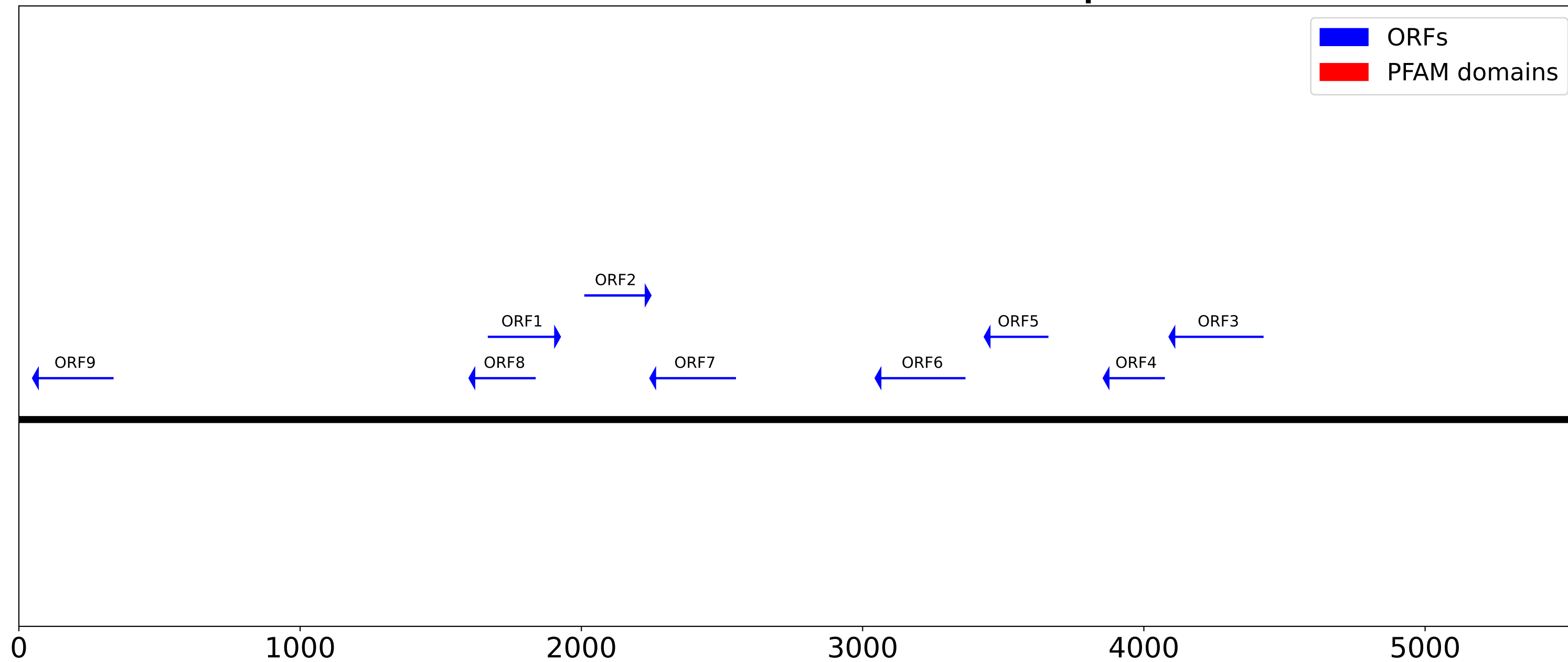


TE: rnd_5_family_8462
size: 4146bp; fragments: 1412; full length: 27 (≥ 3731.4 bp)

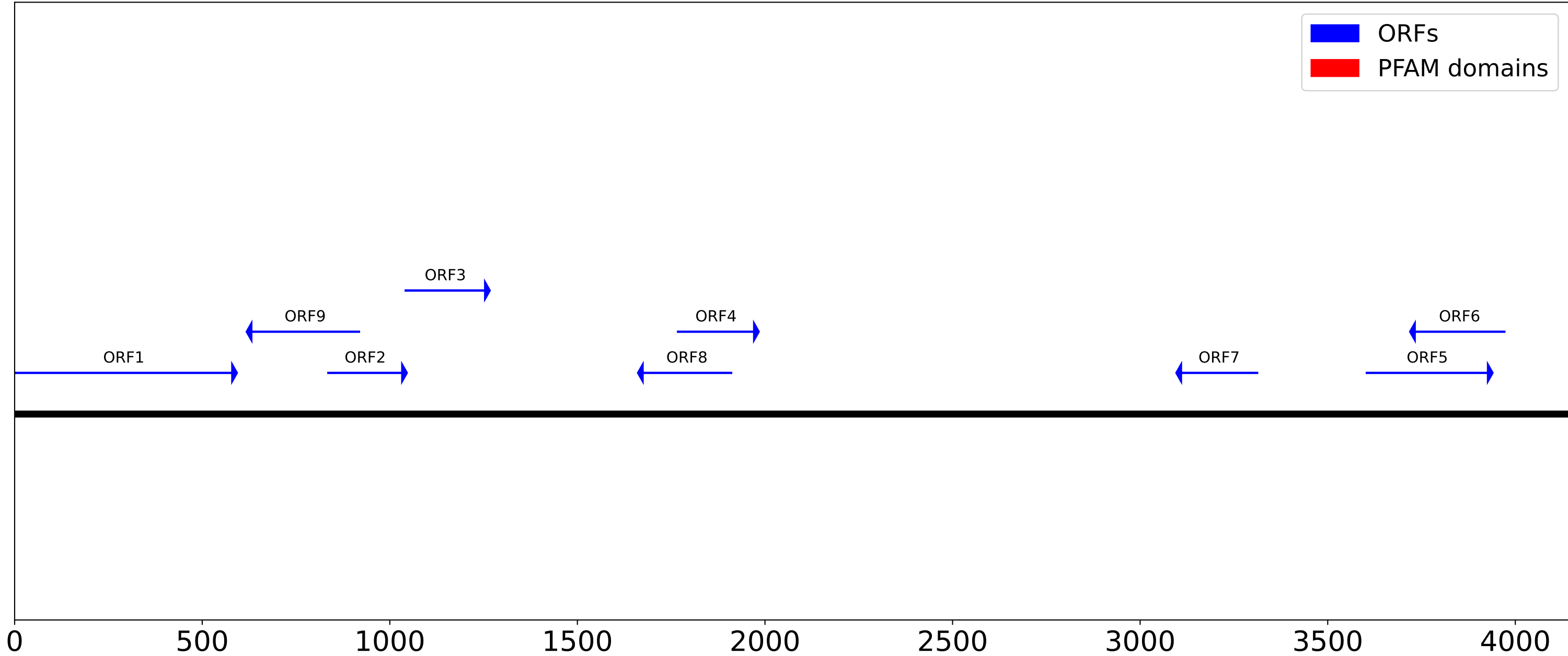
Before TEtrimmer 4146 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

