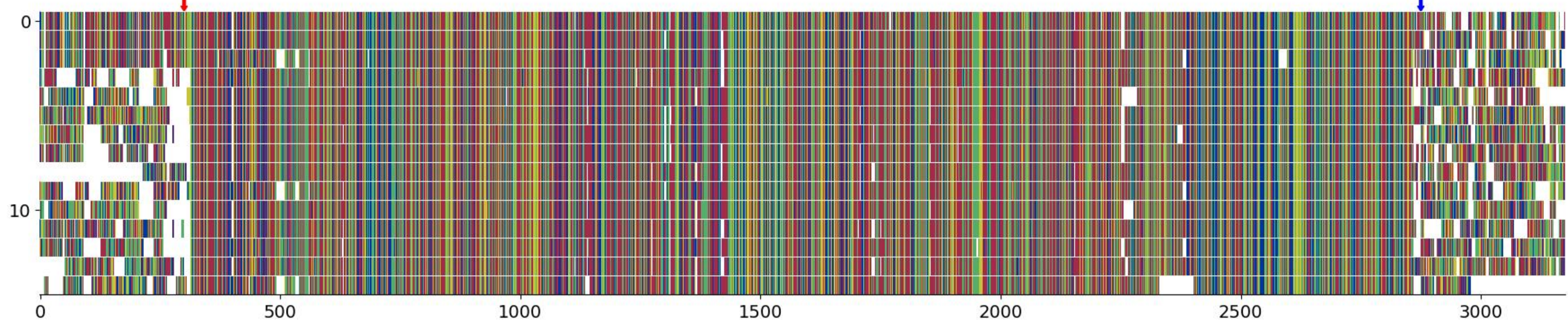


MSA length = 2576



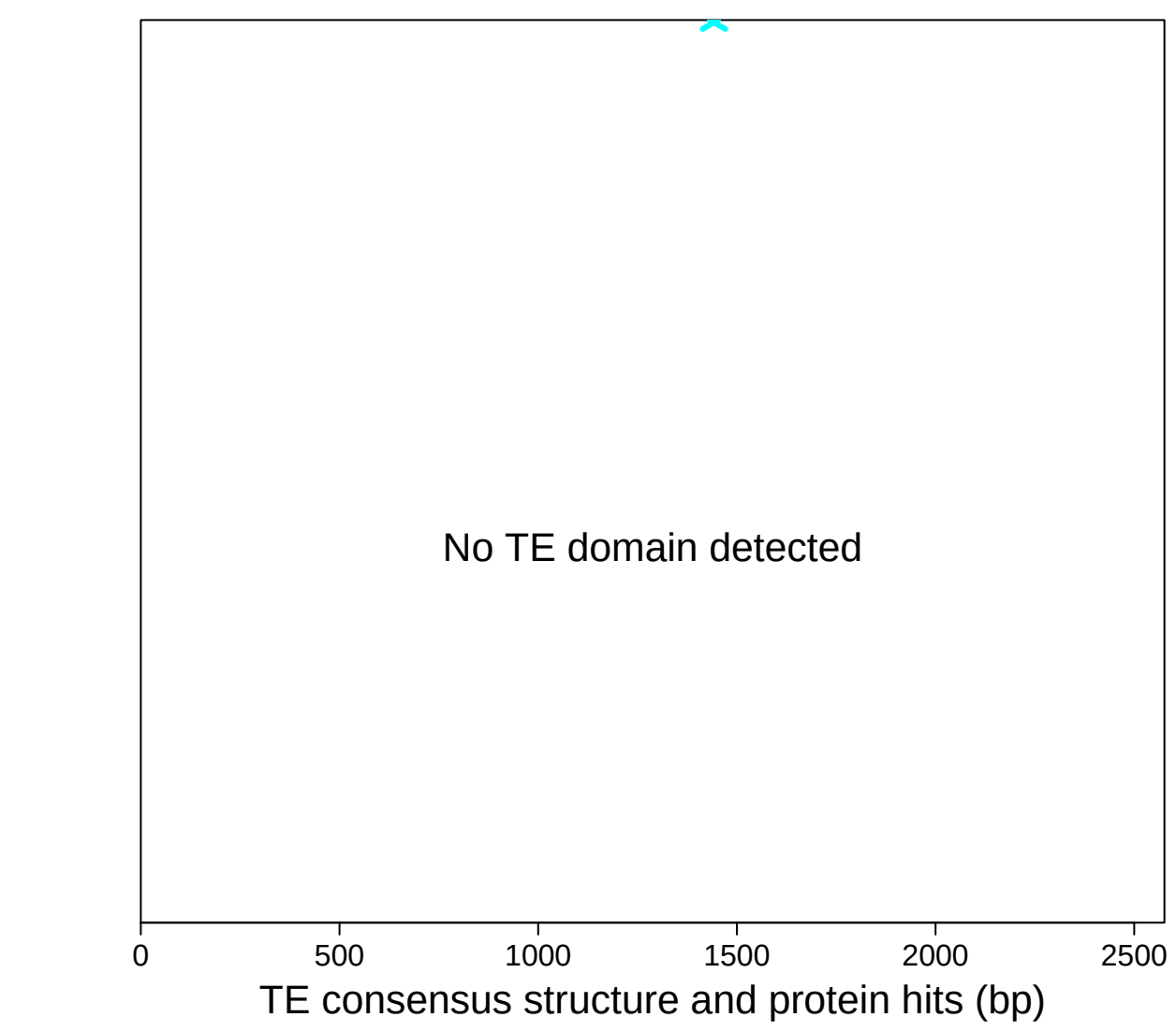
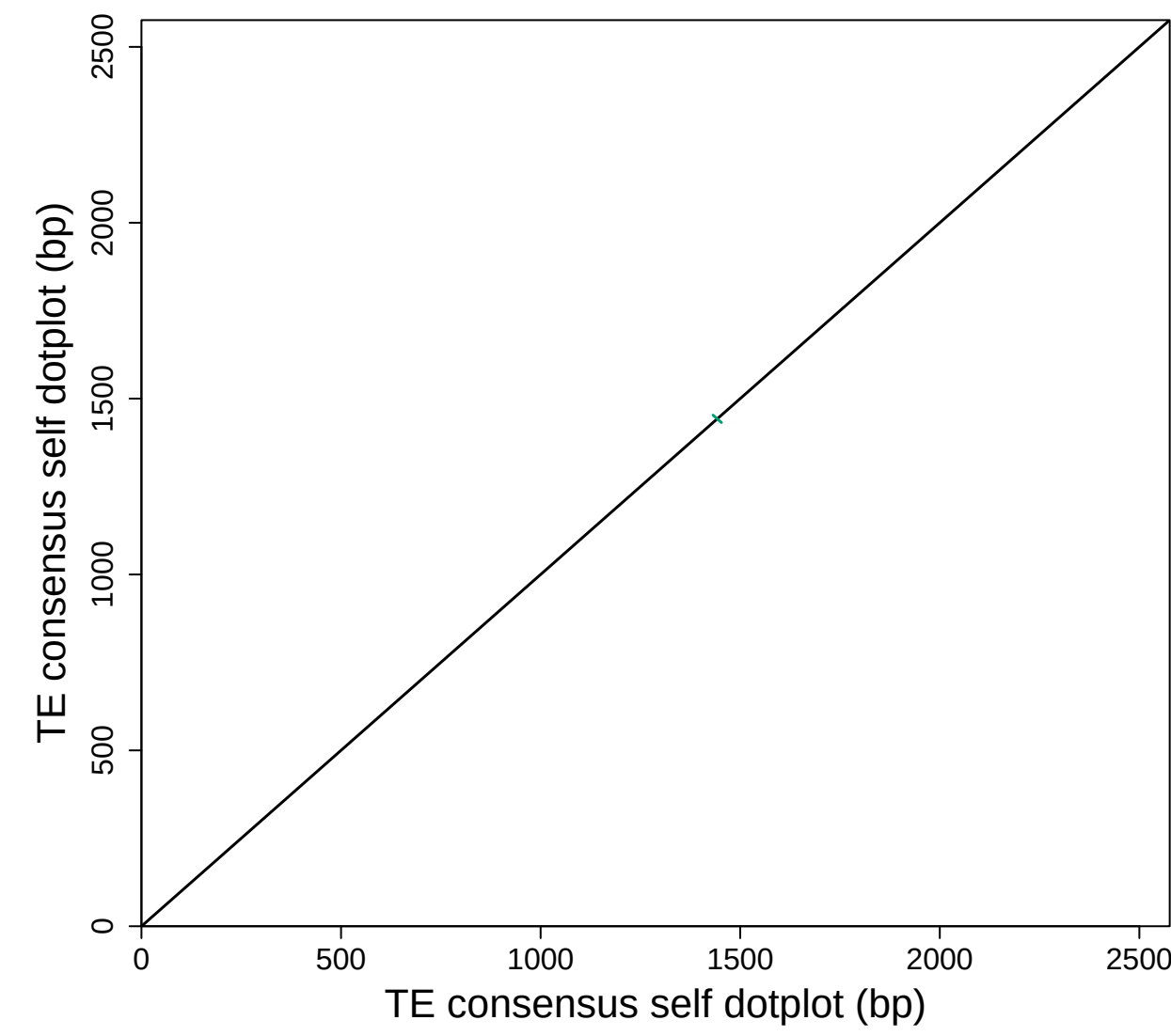
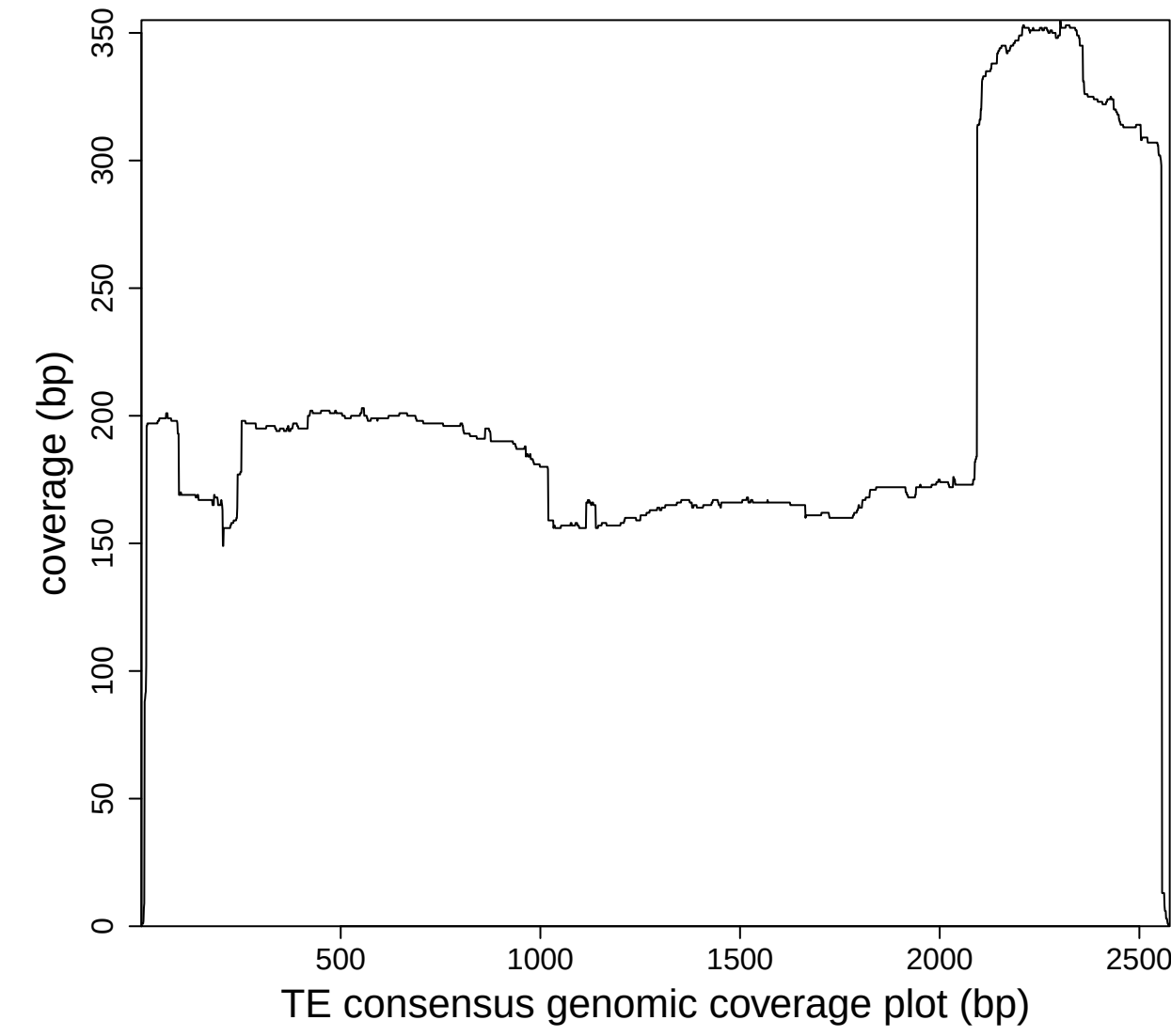
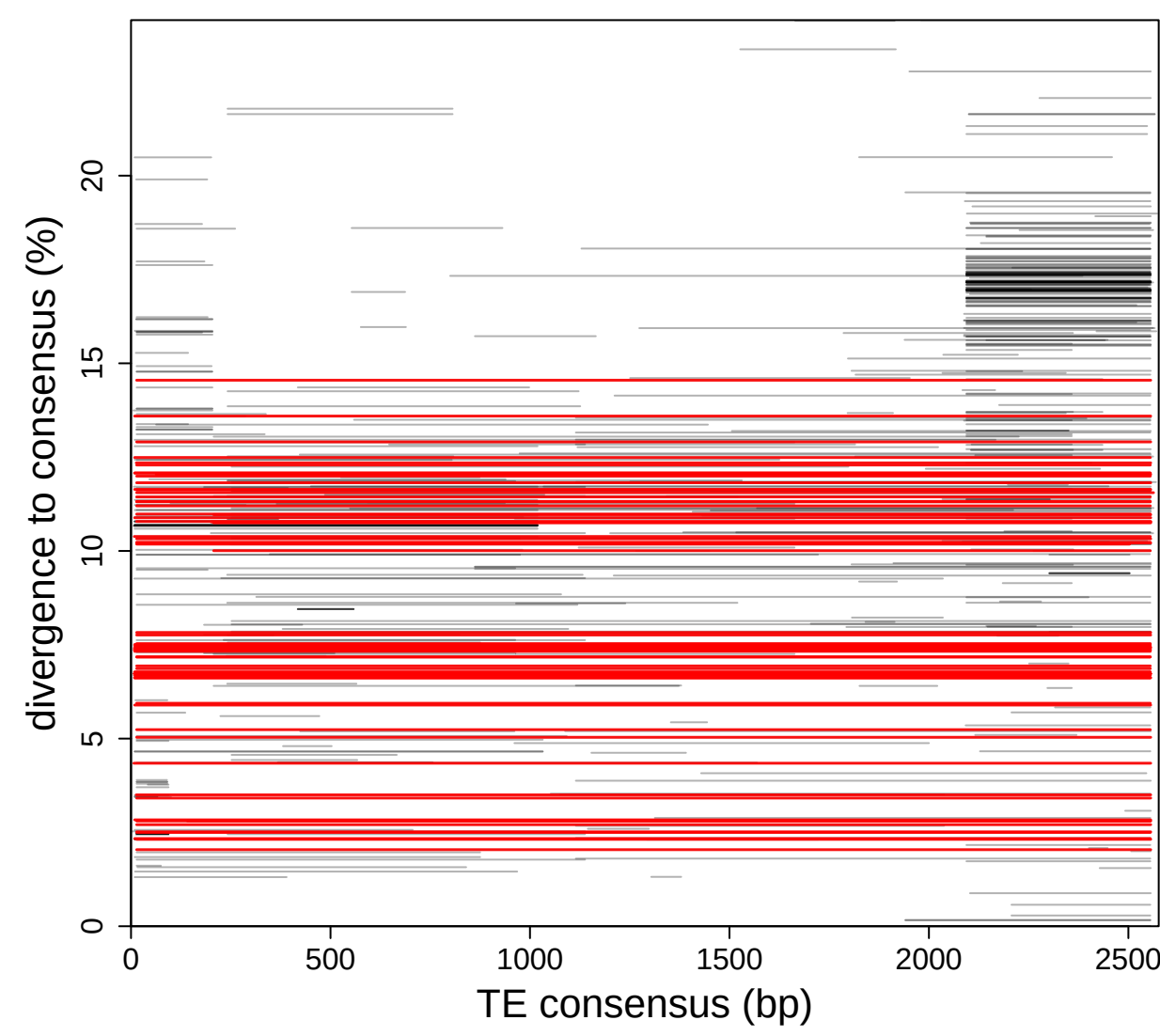
Start crop Point

End crop Point



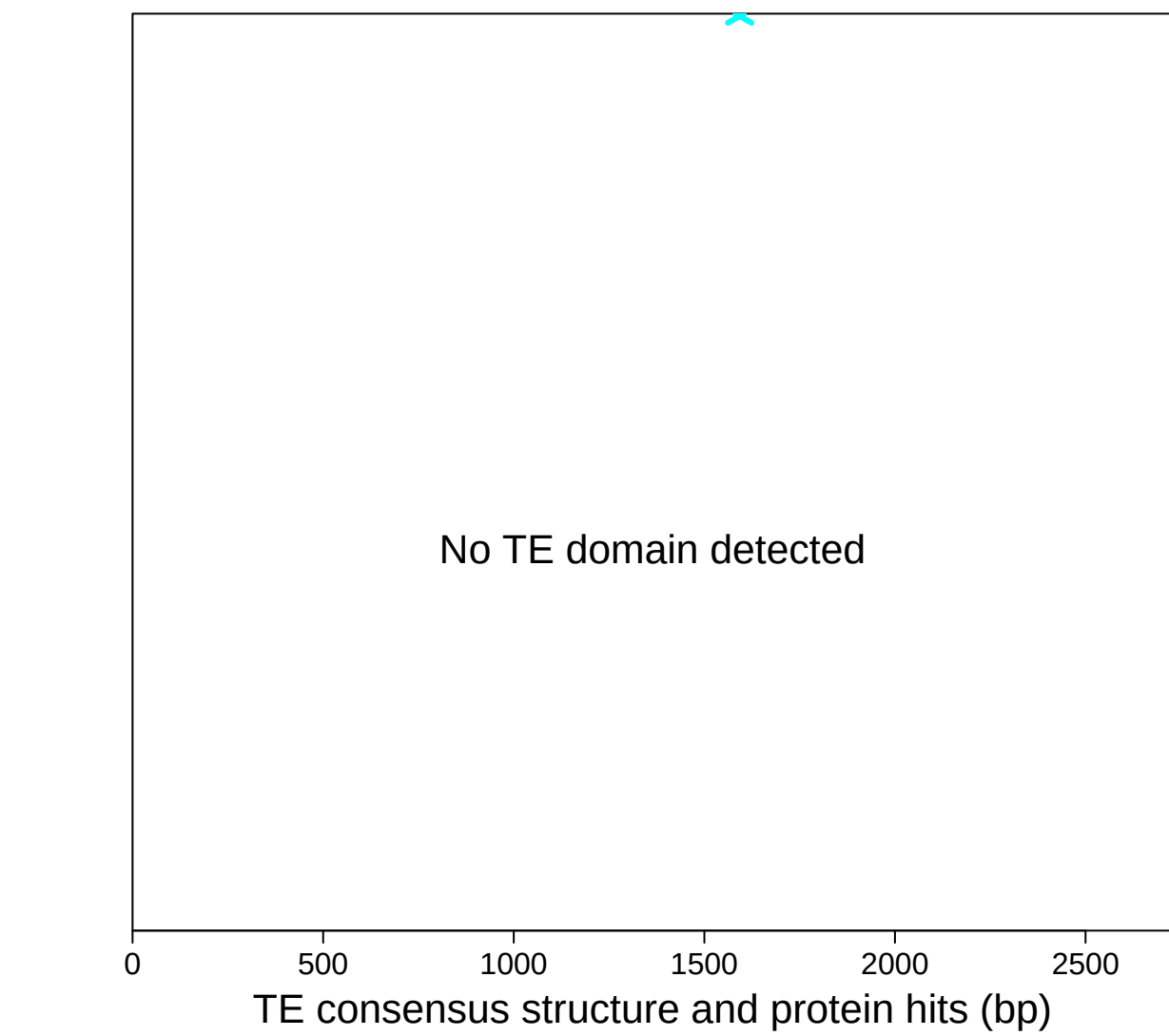
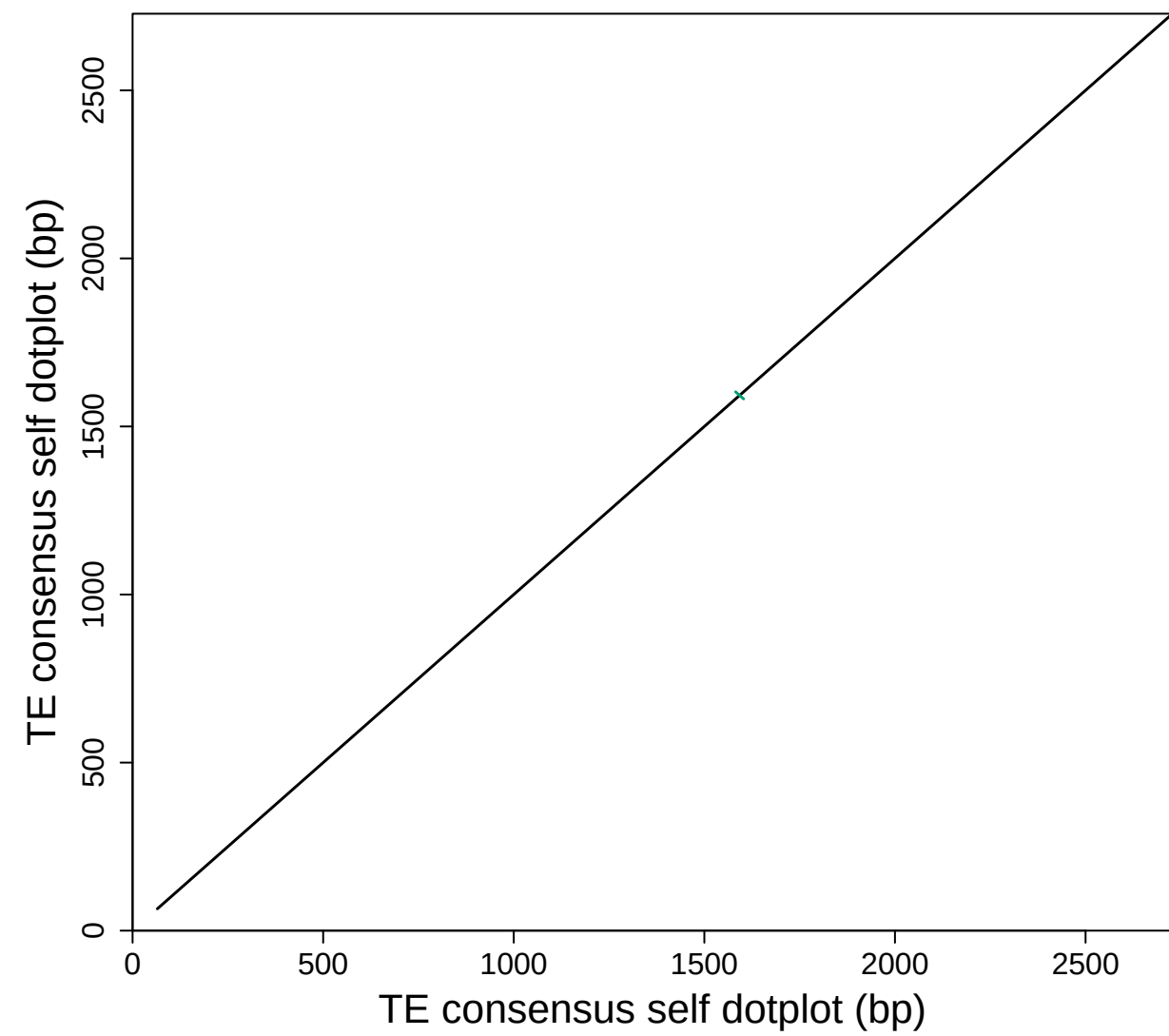
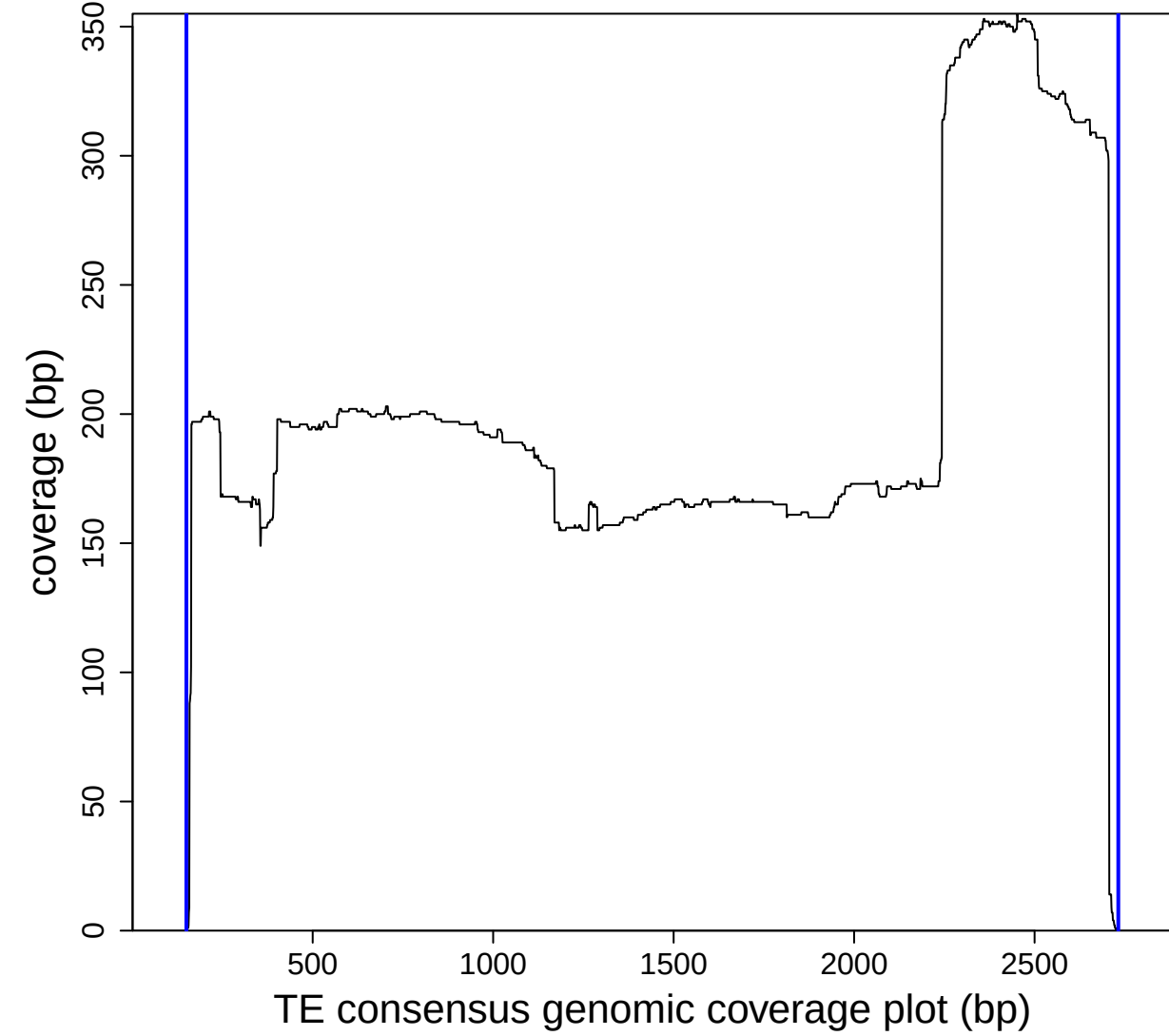
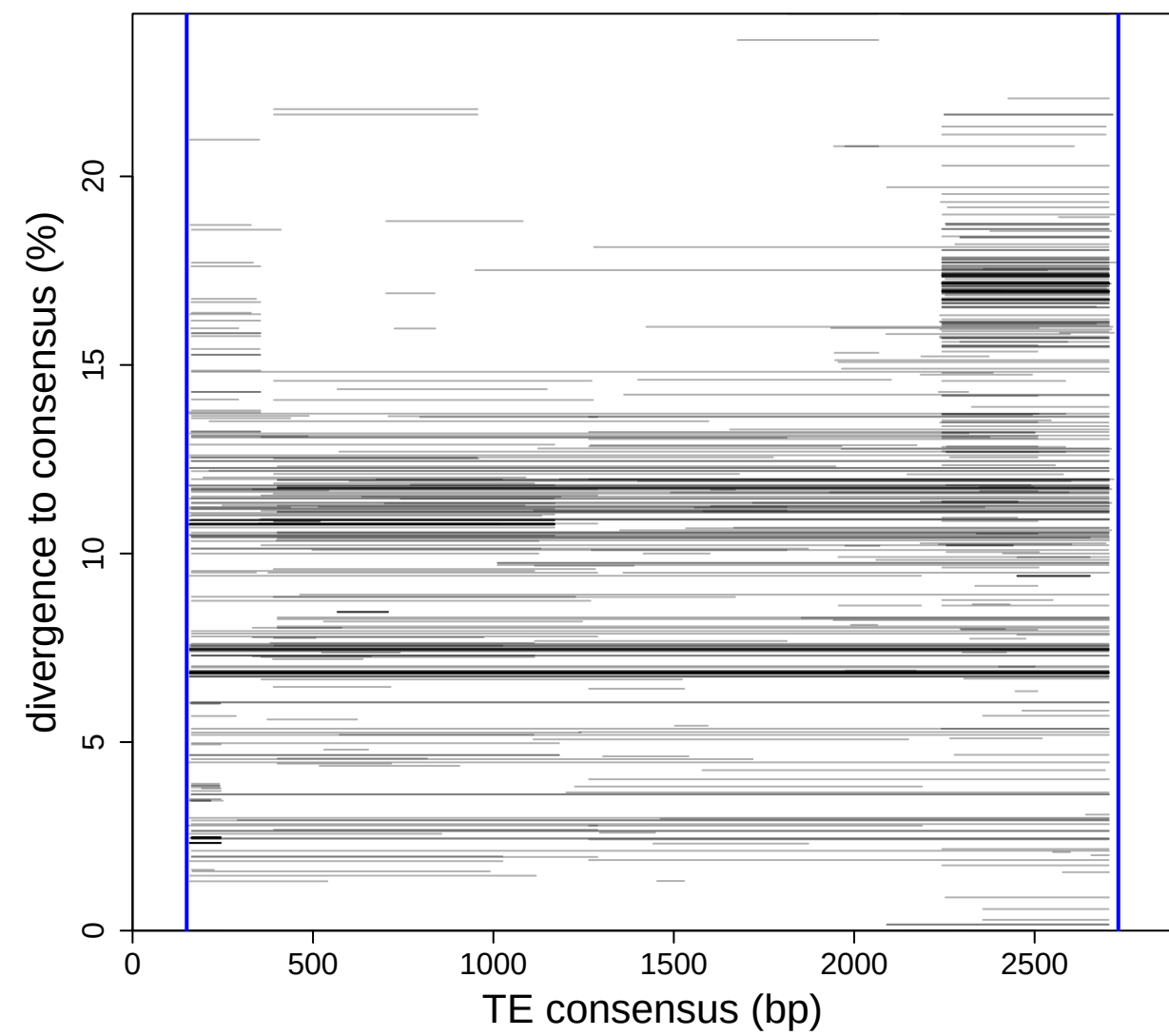
a.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 2576bp; fragments: 566; full length: 87 (>=2318.4bp)

After TEtrimmer 2576 bp

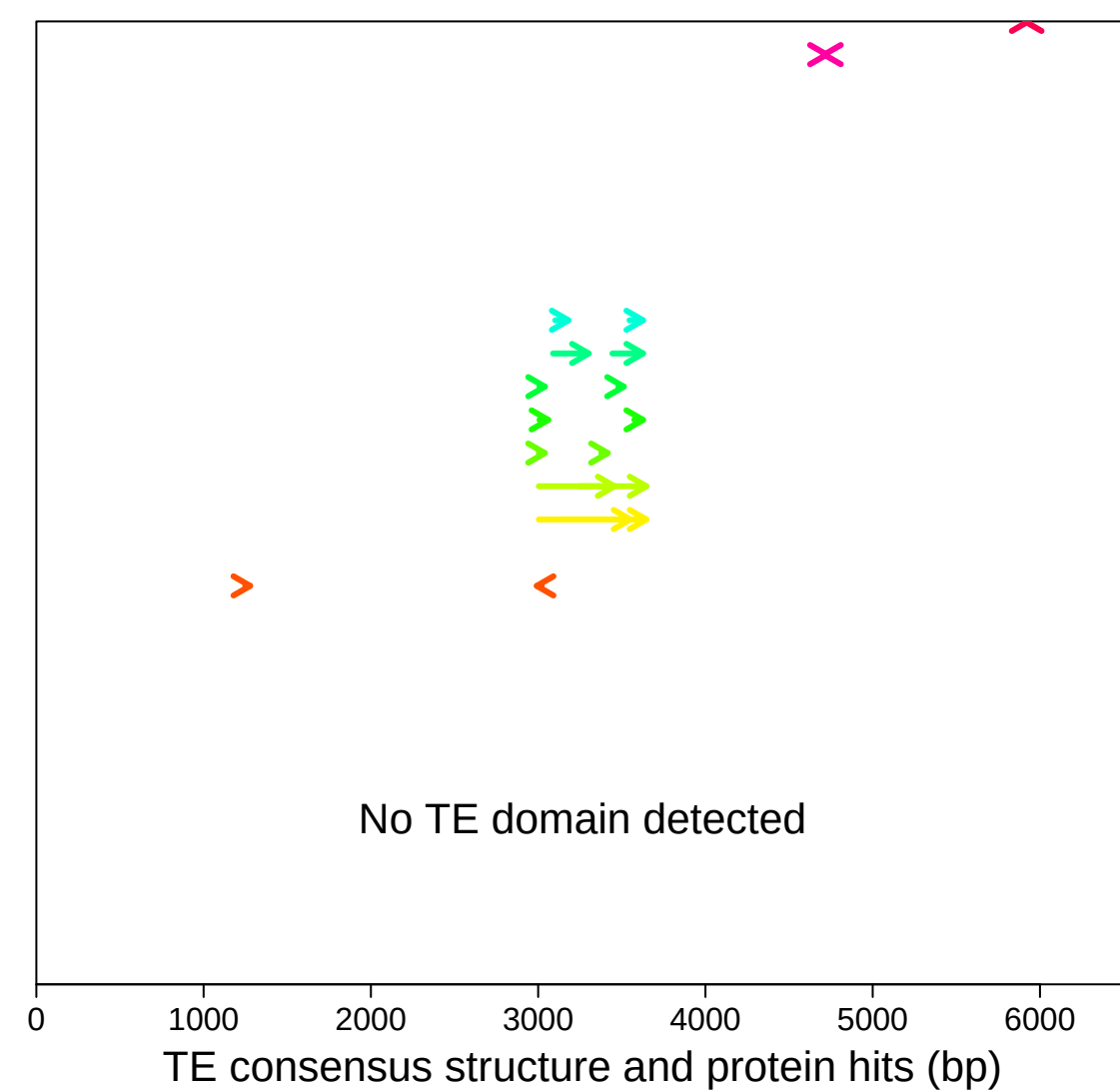
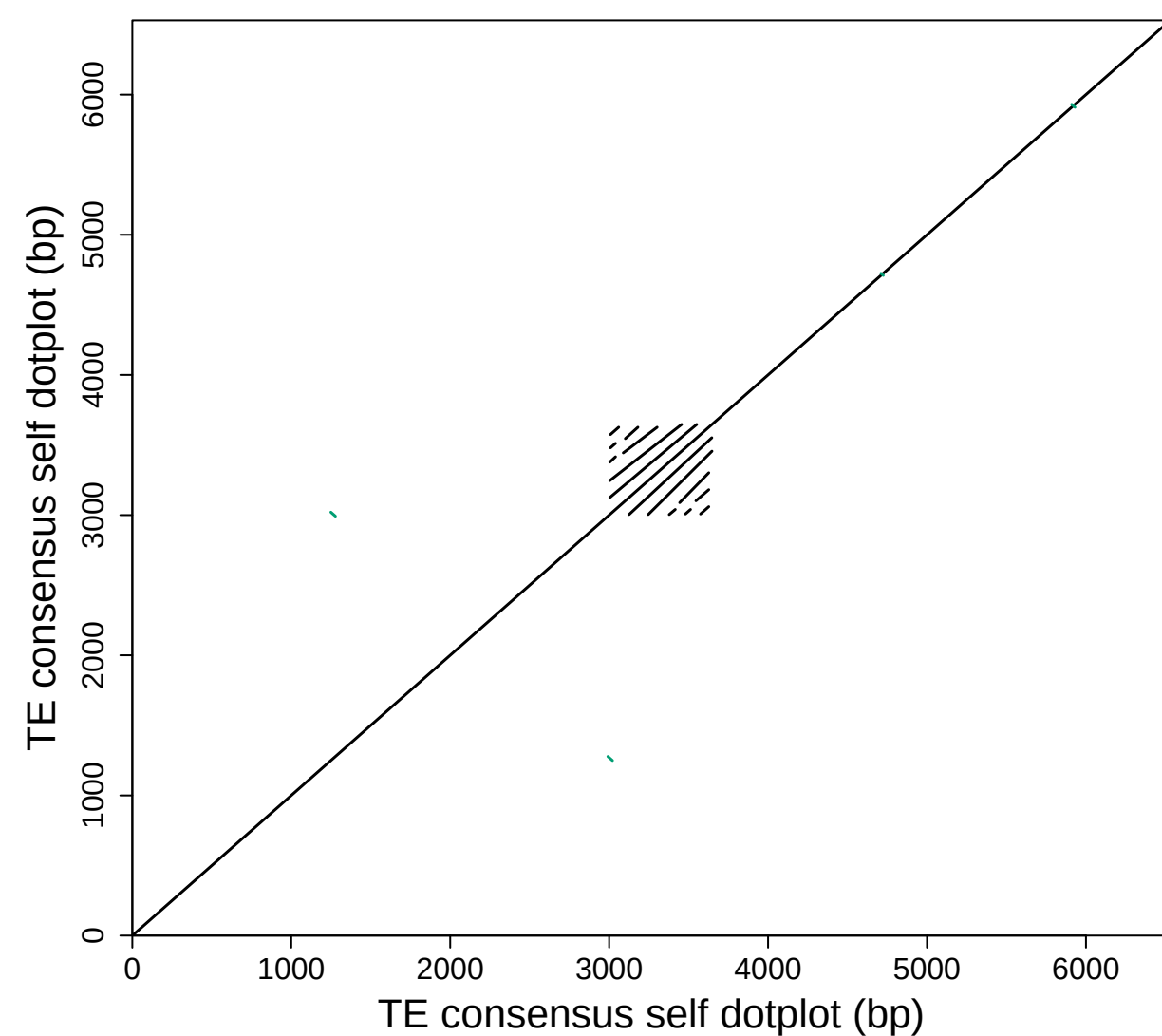
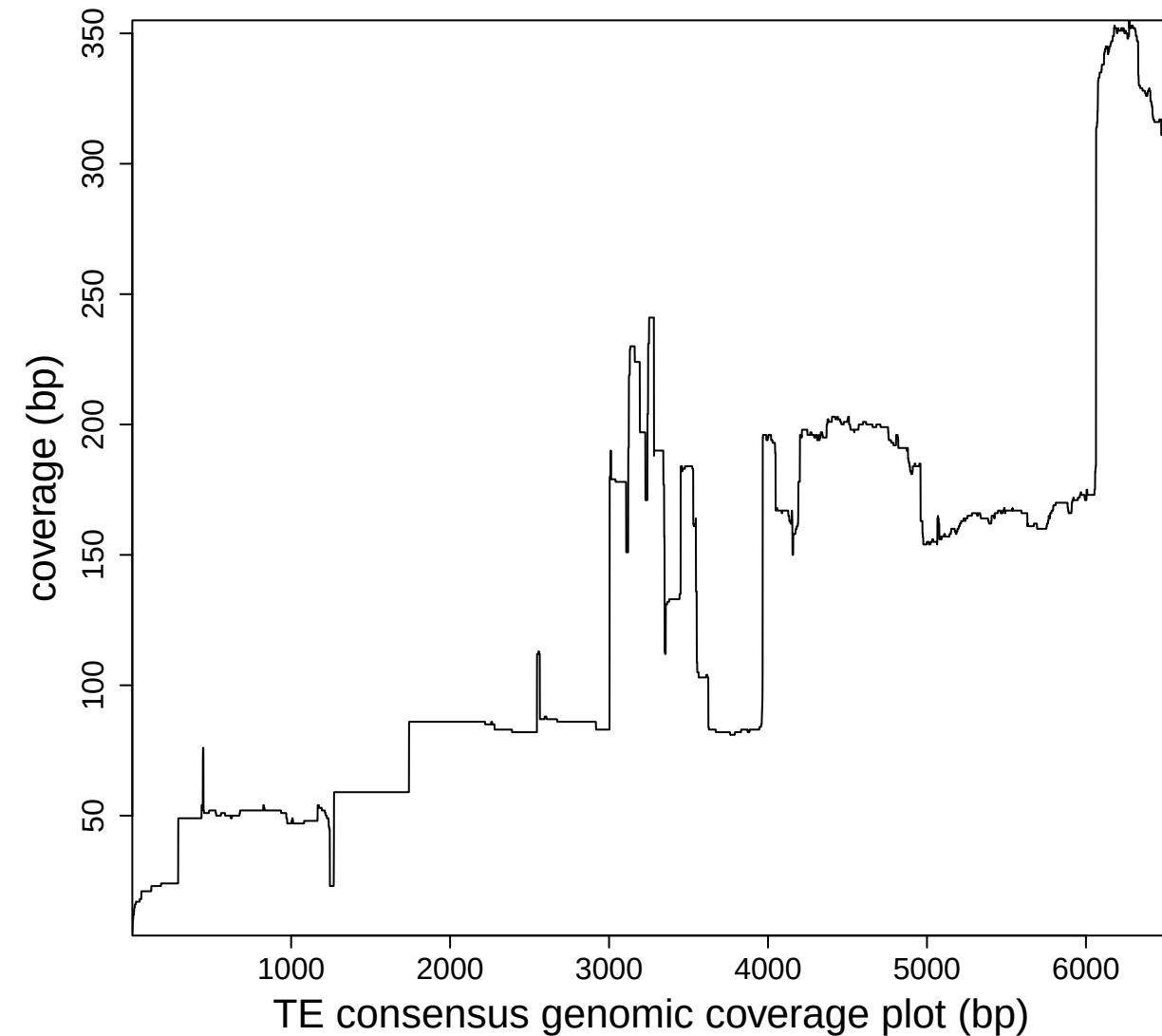
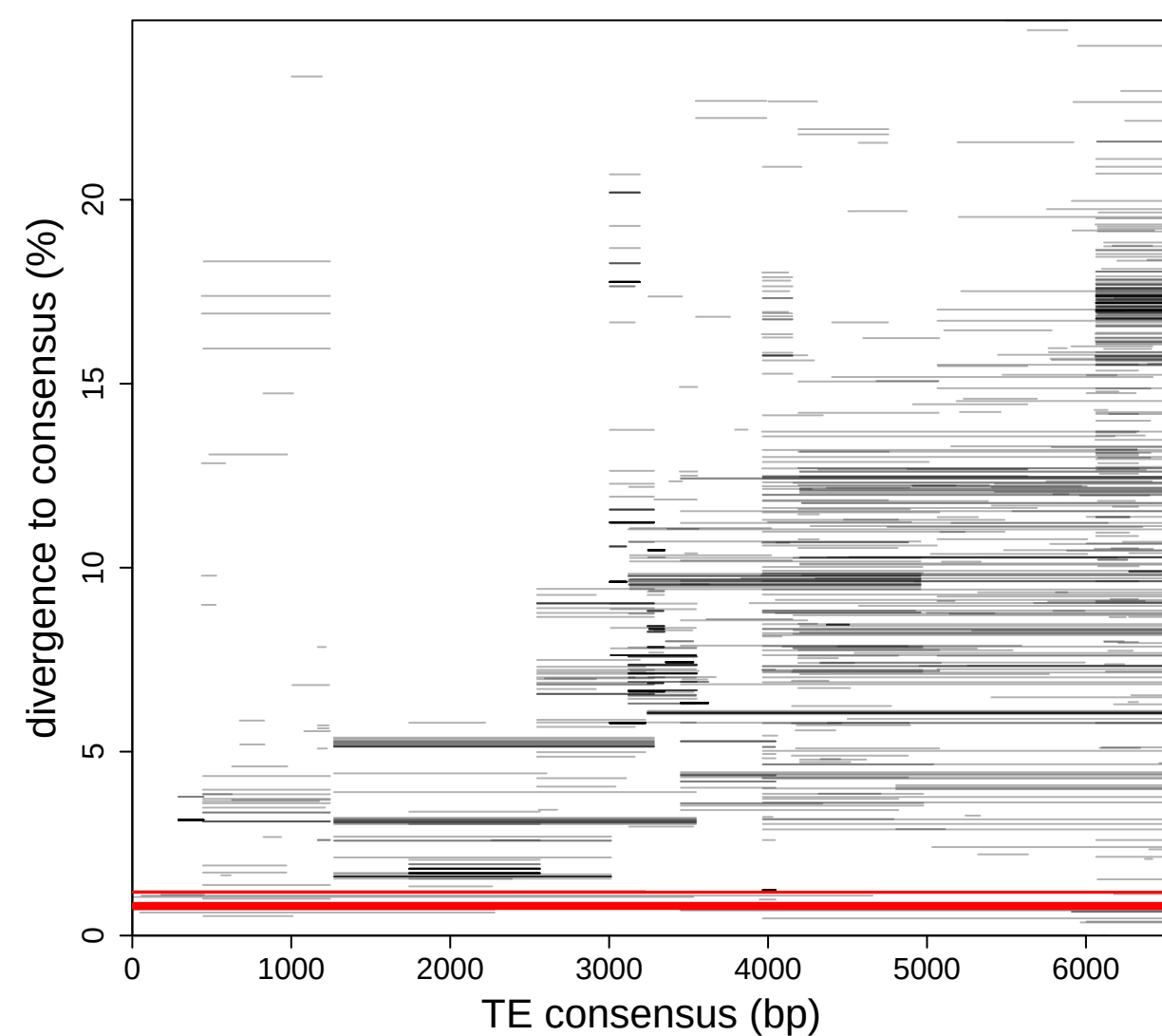


5.fasta.b.bed uf.bed g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 2882bp; fragments: 566; full length: 0 (>=2593.8bp)

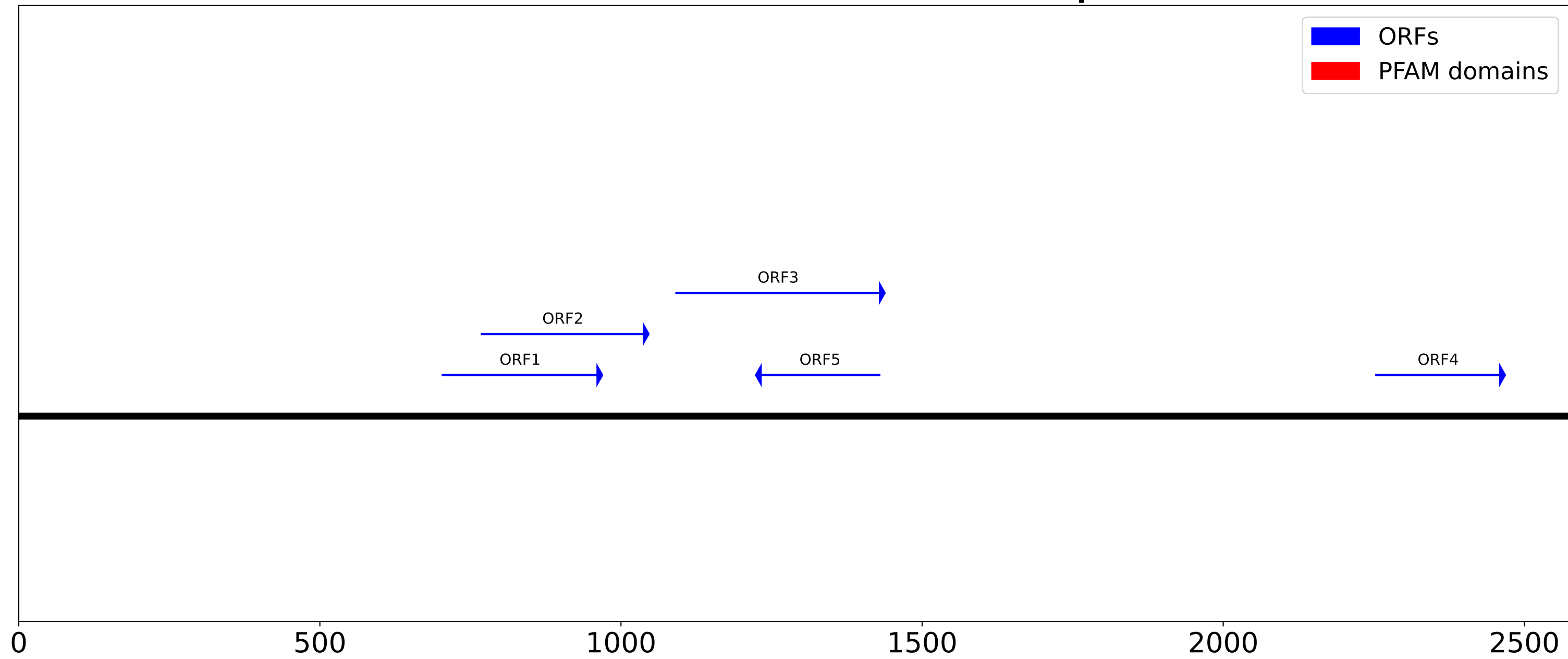
After TEtrimmer Extended plot Blue lines are boundaries



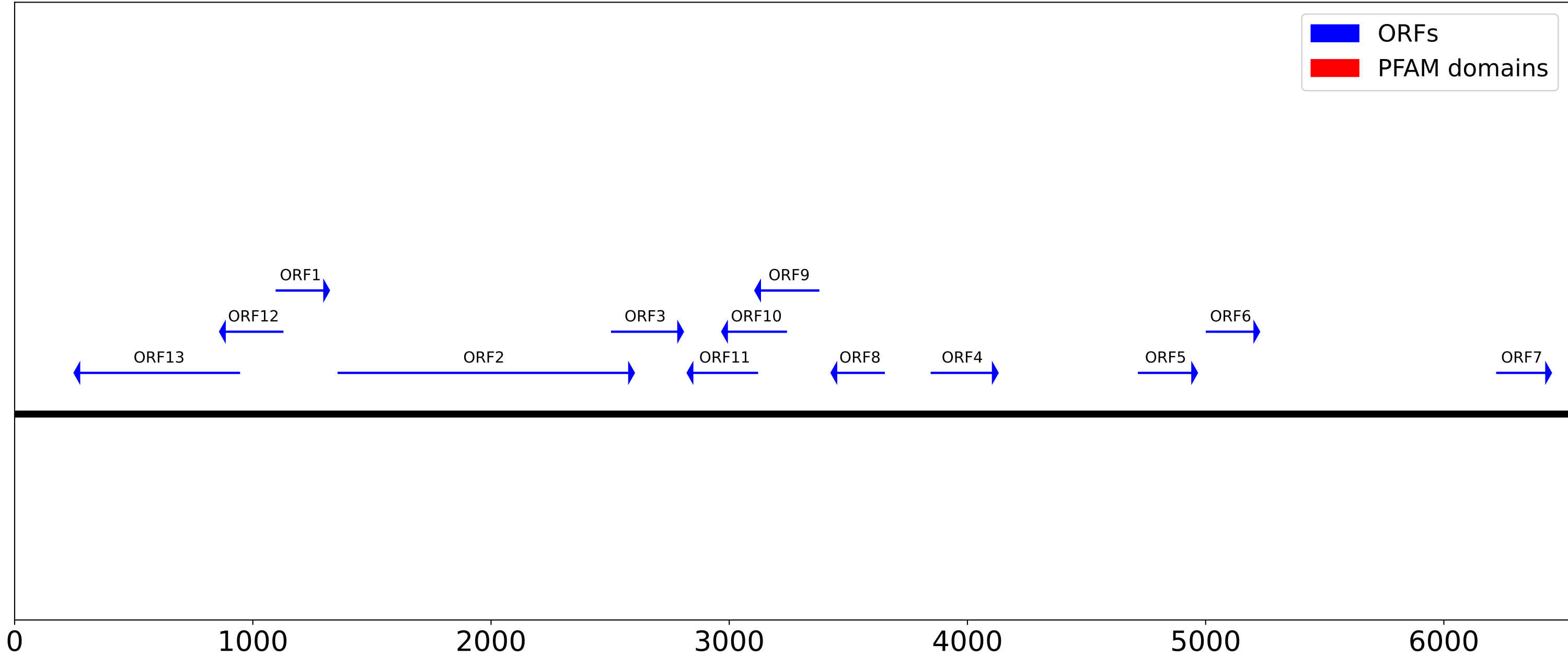
TE: rnd_5_family_9115
size: 6530bp; fragments: 1010; full length: 18 (≥ 5877 bp)



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

