

Start crop Point

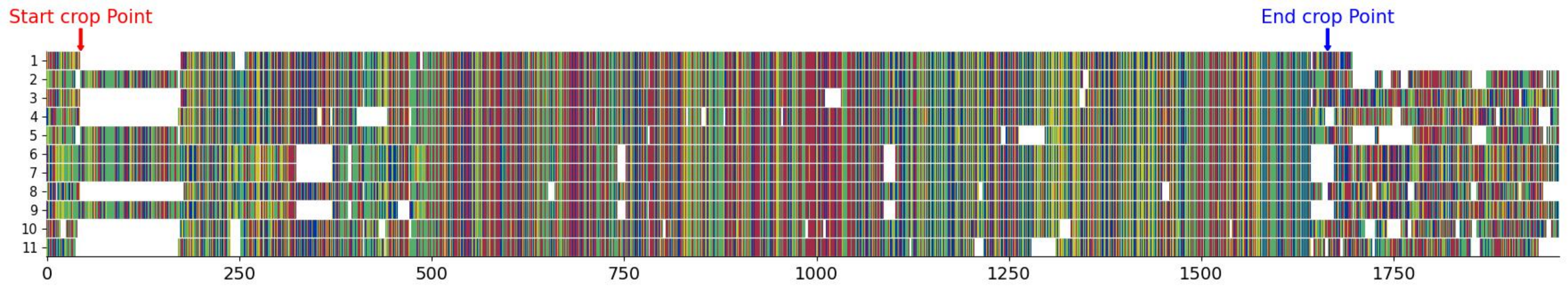


End crop Point



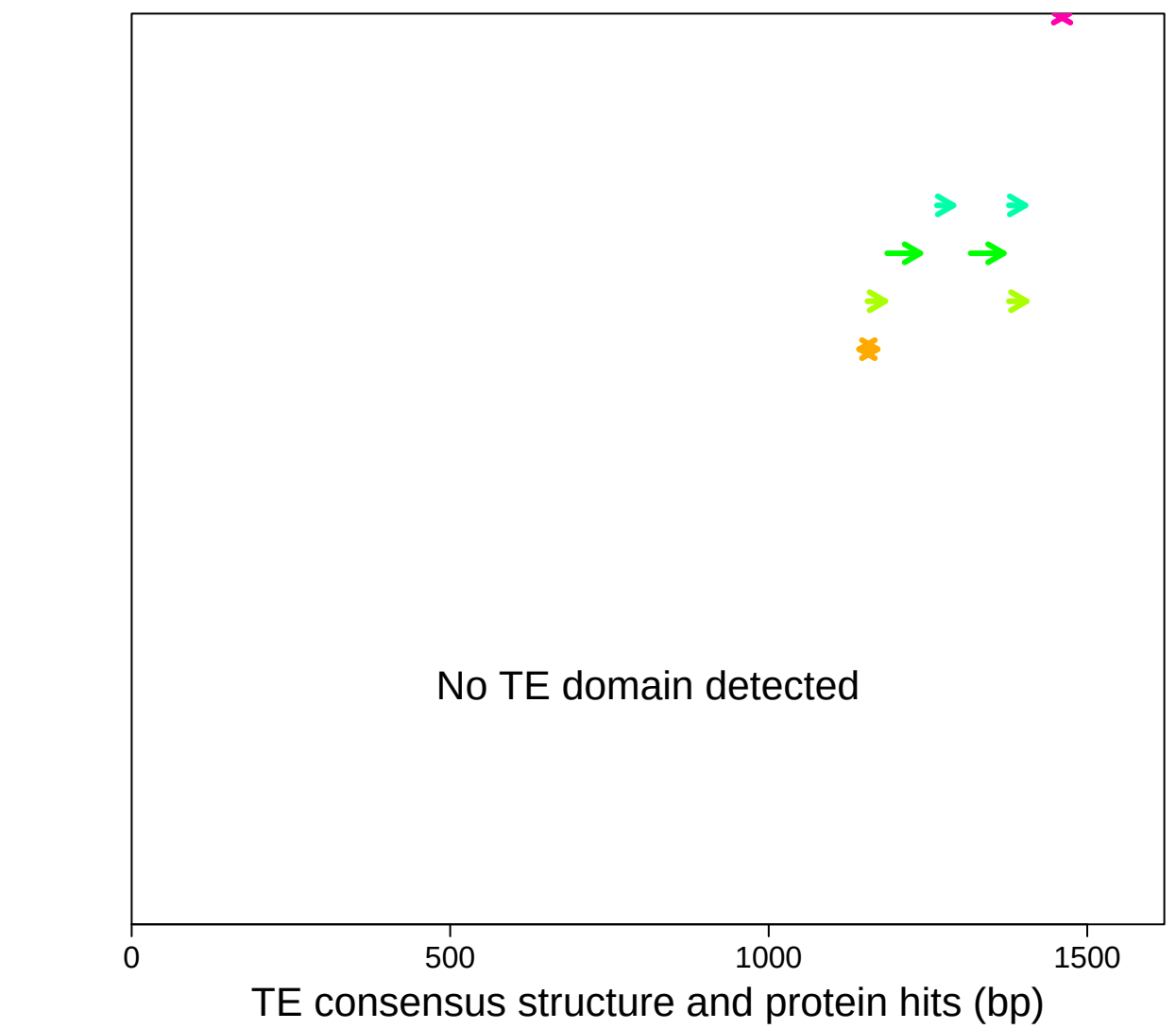
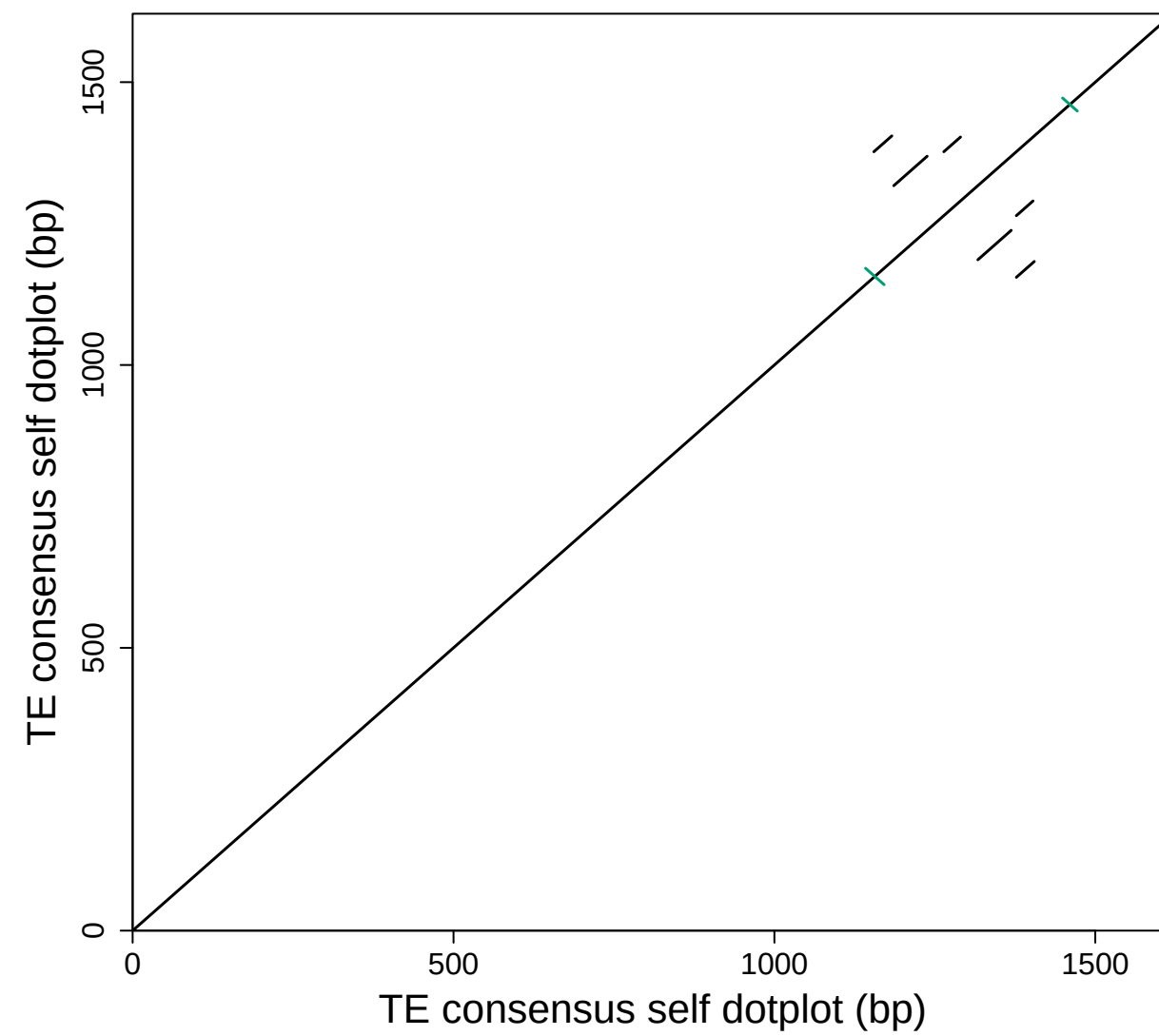
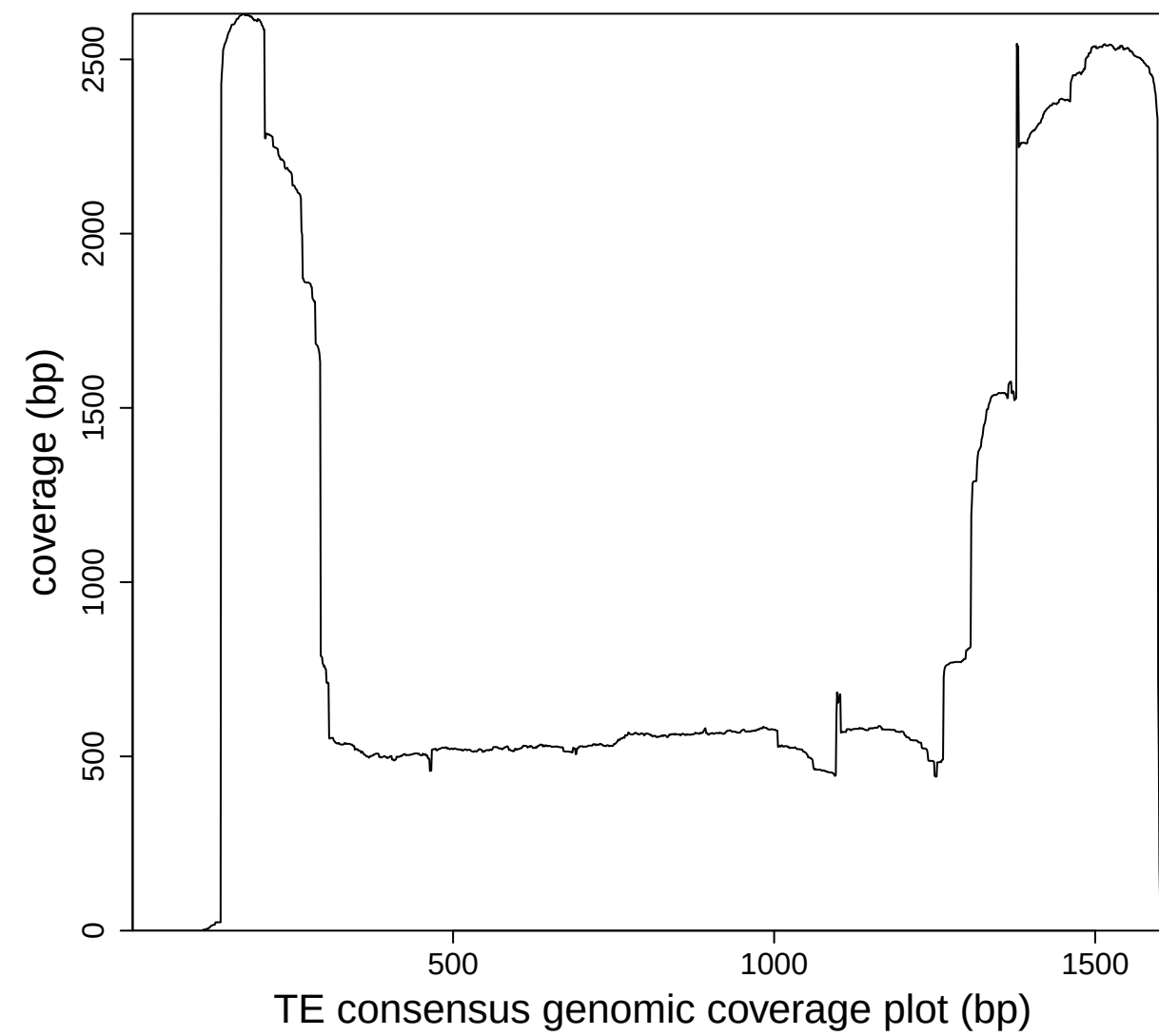
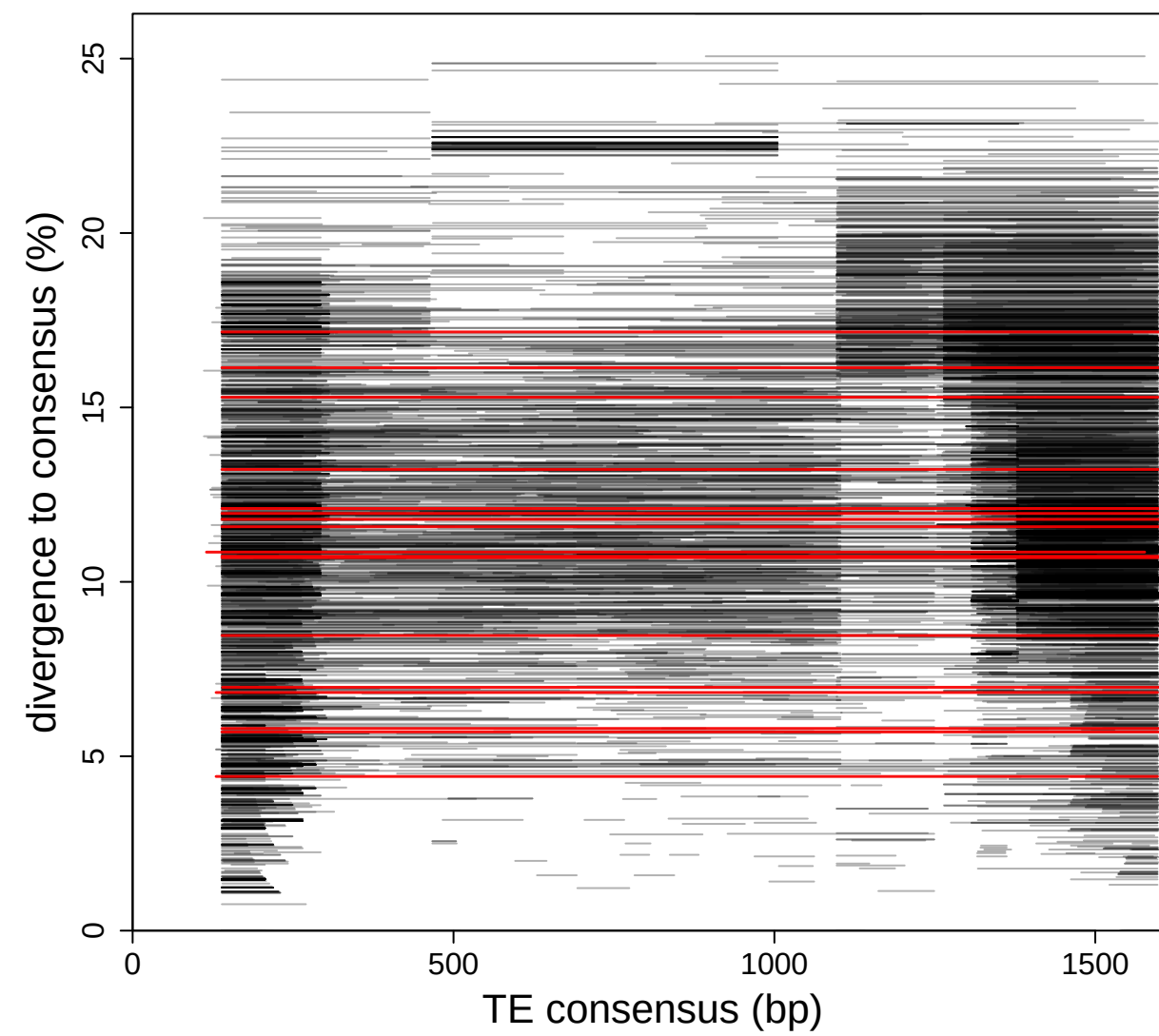
MSA length = 1621





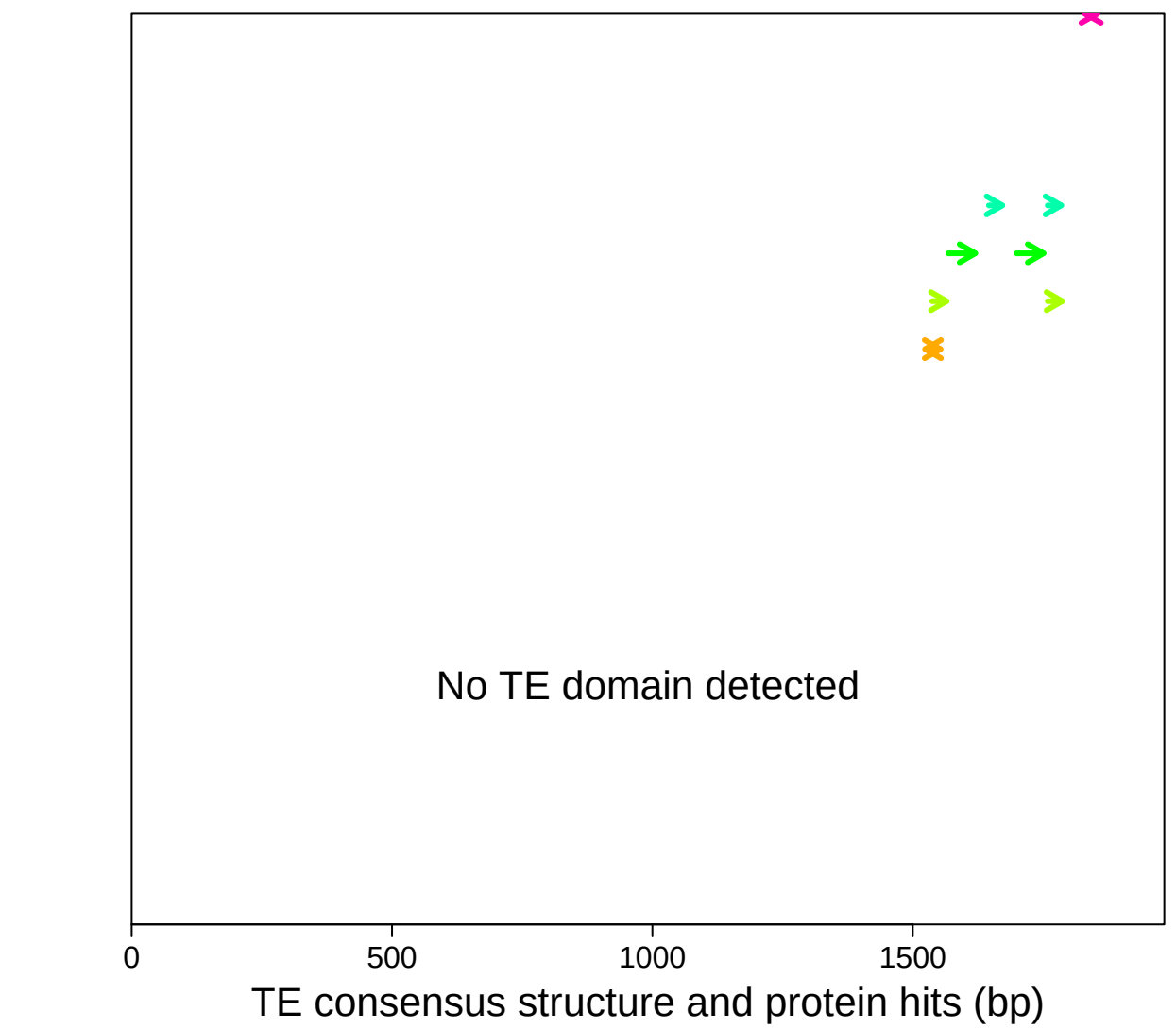
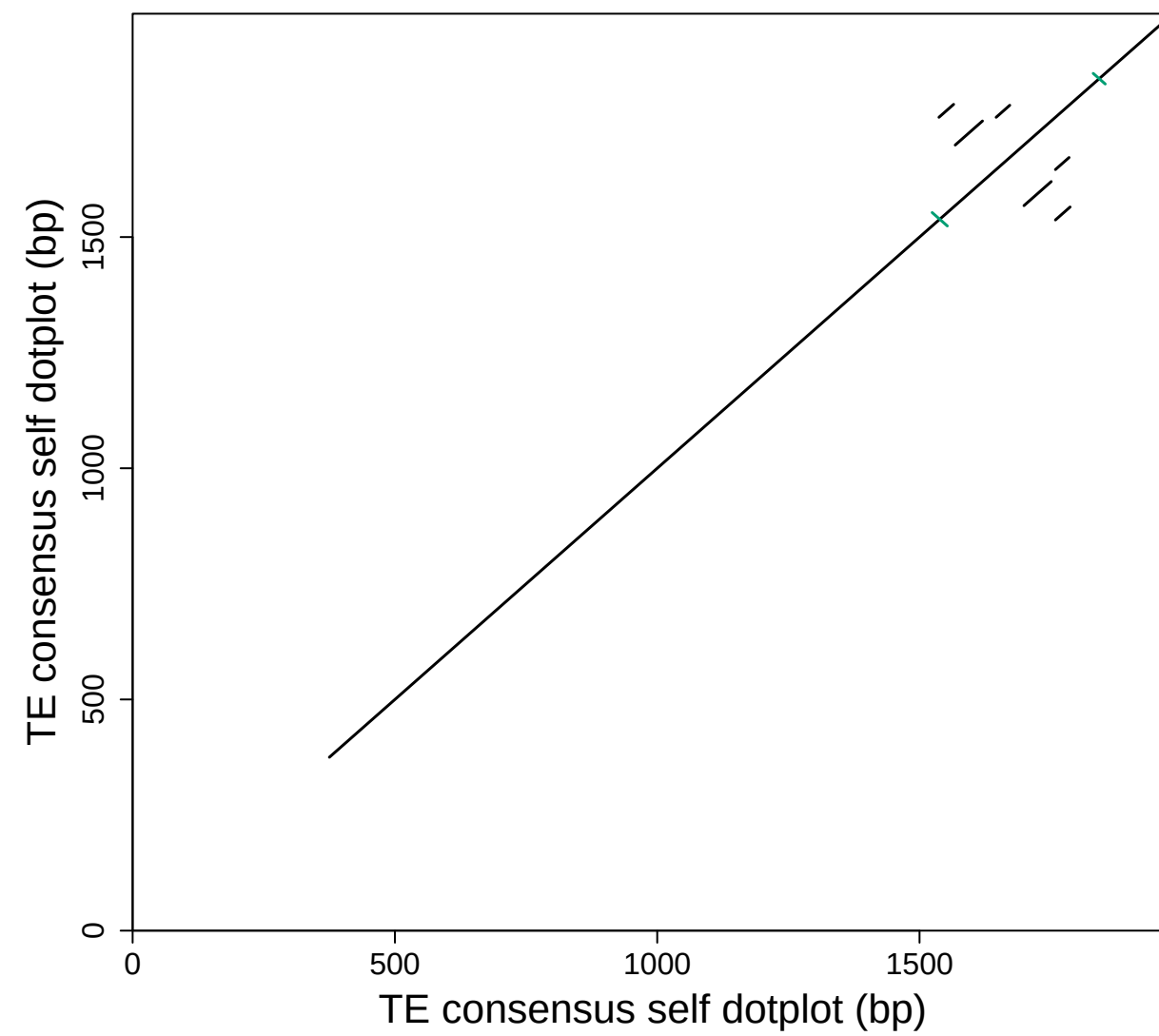
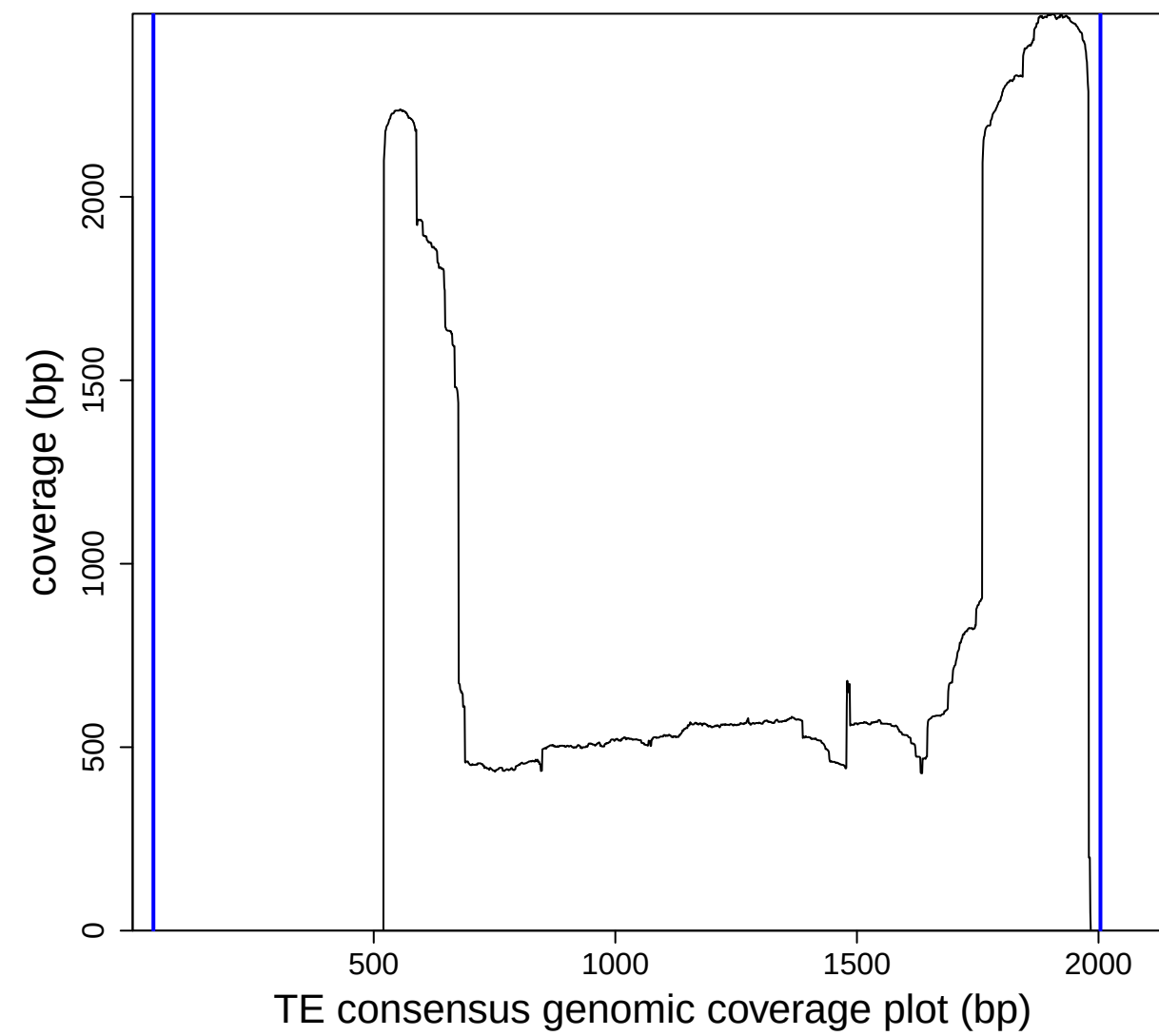
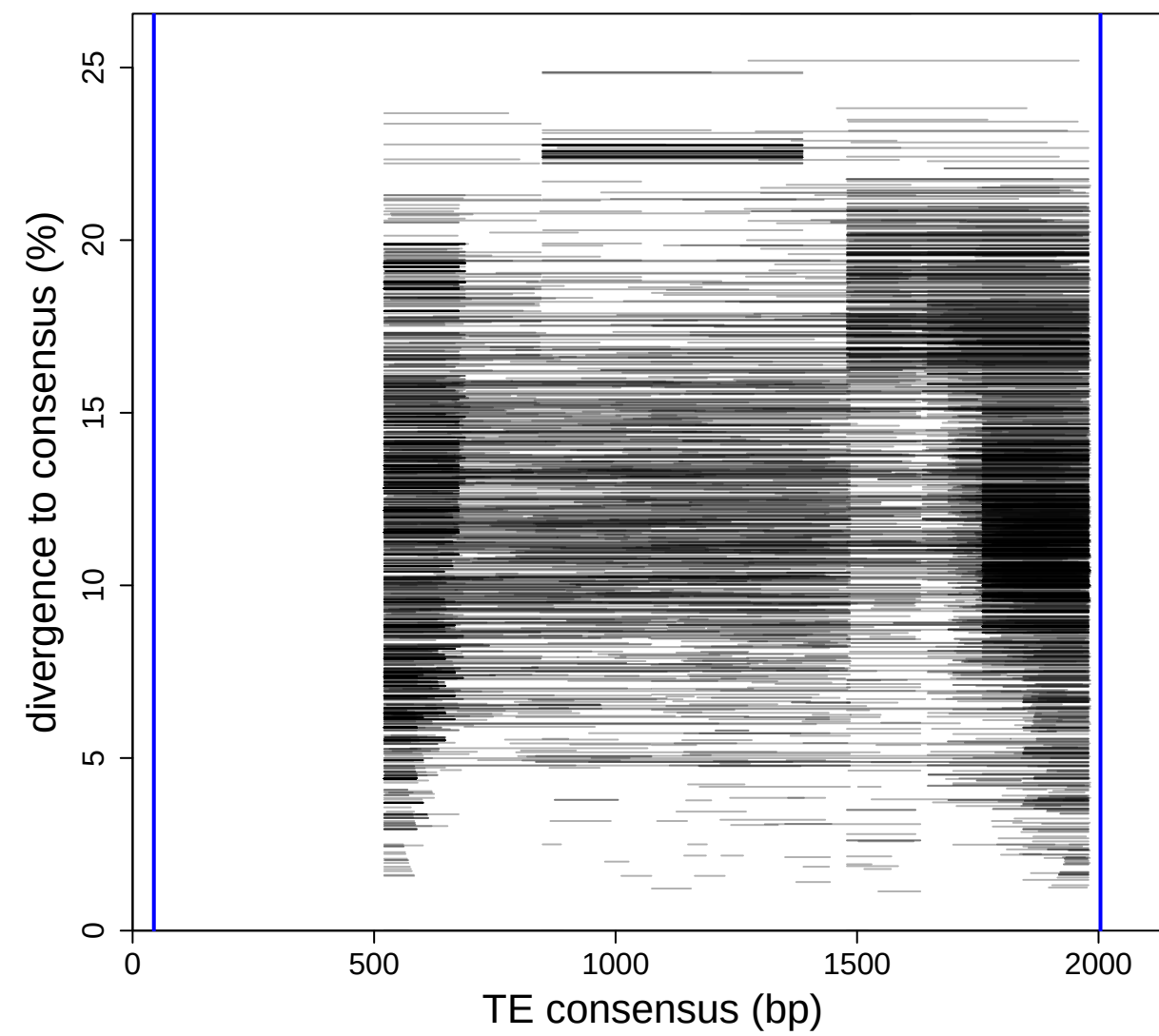
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1621bp; fragments: 6406; full length: 17 (>=1458.9bp)

After TETrimmer 1621 bp

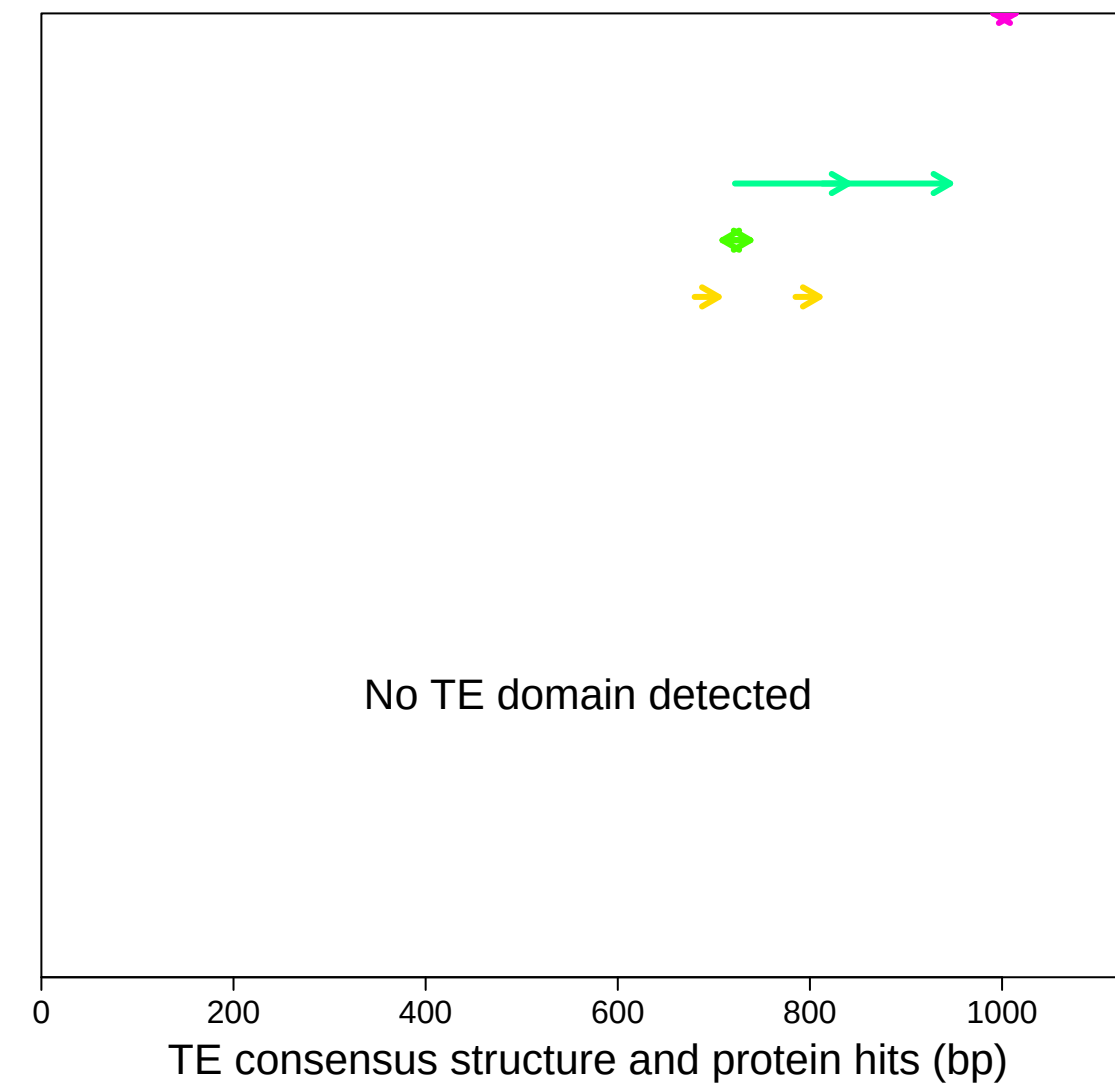
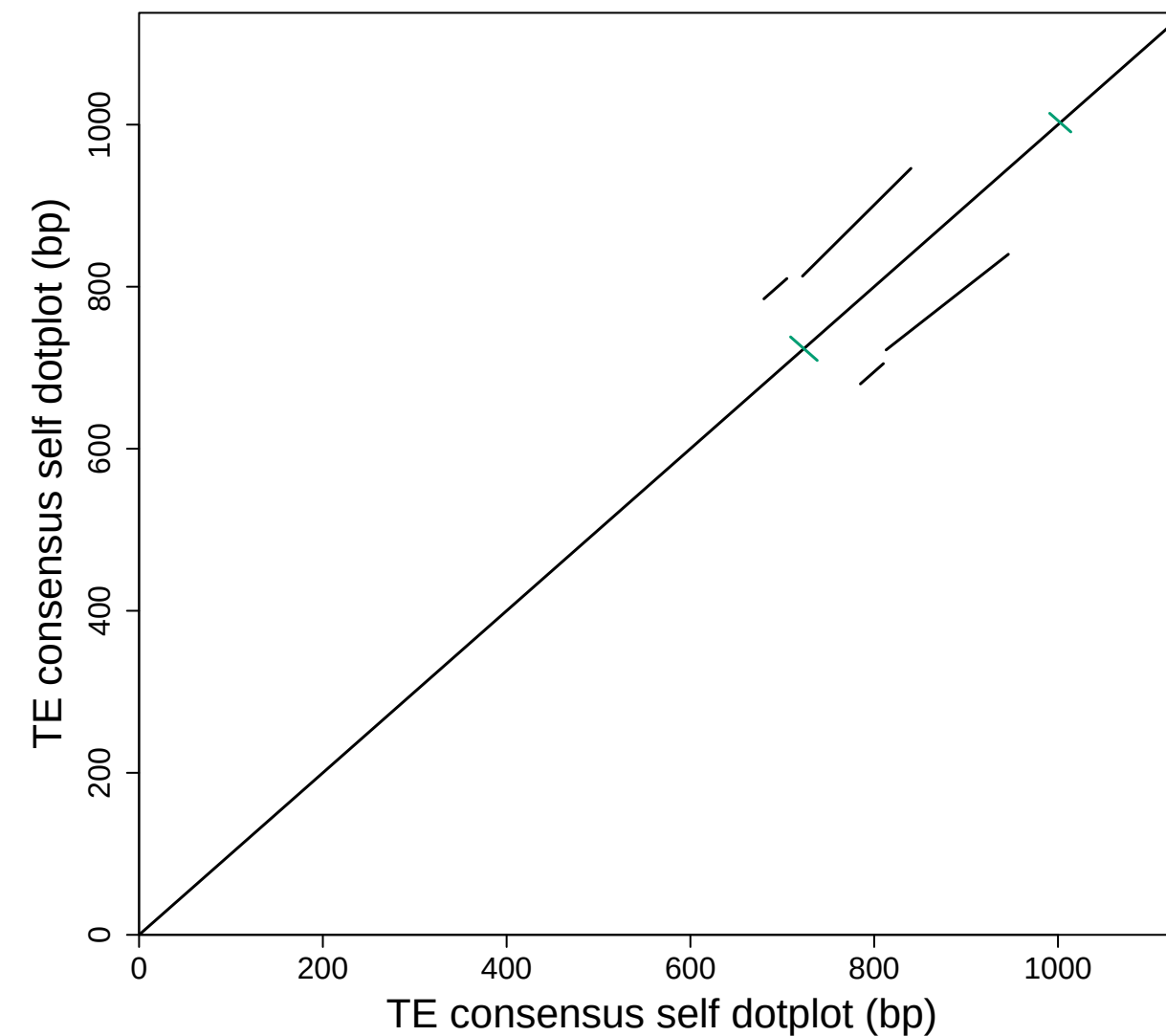
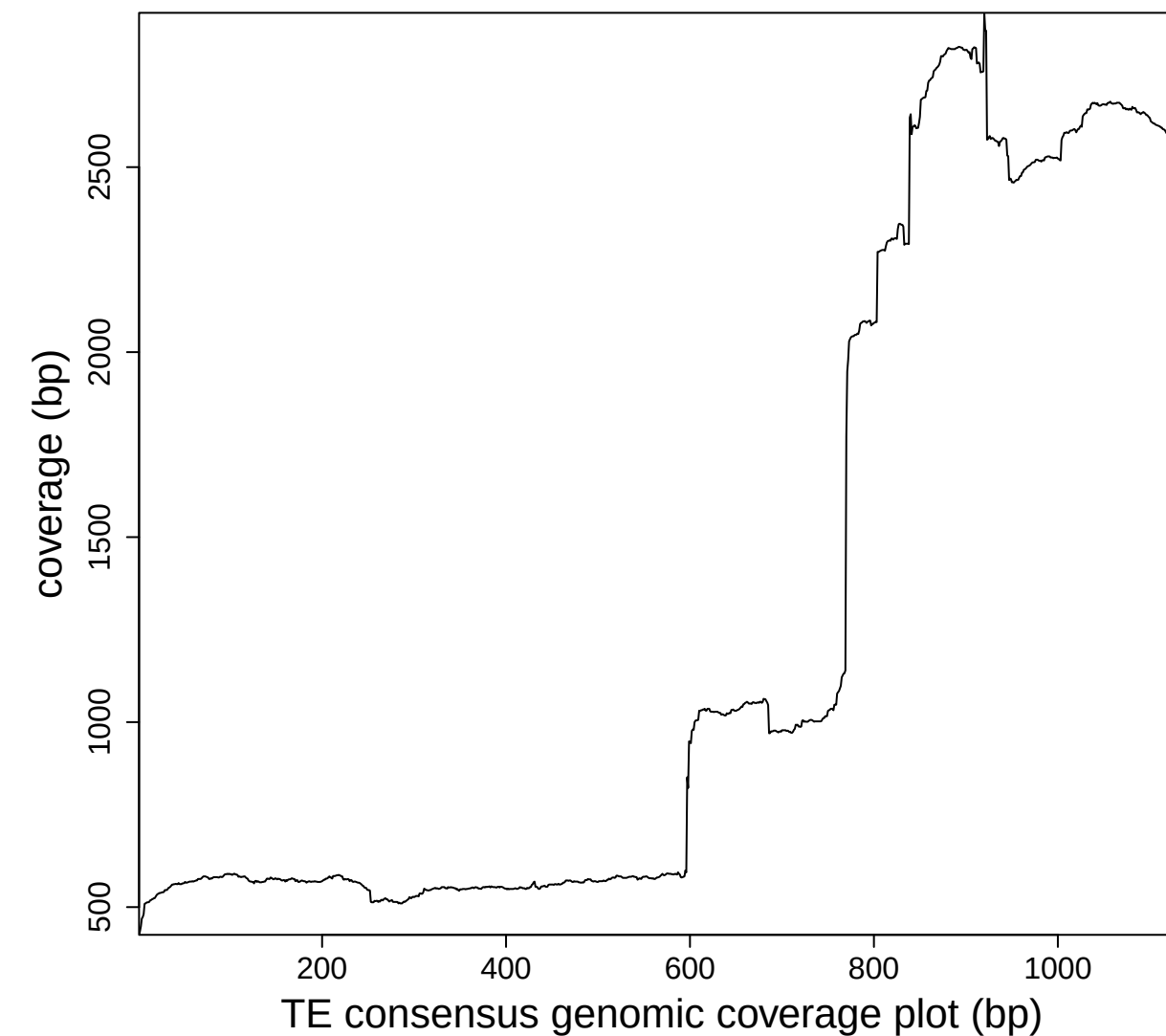
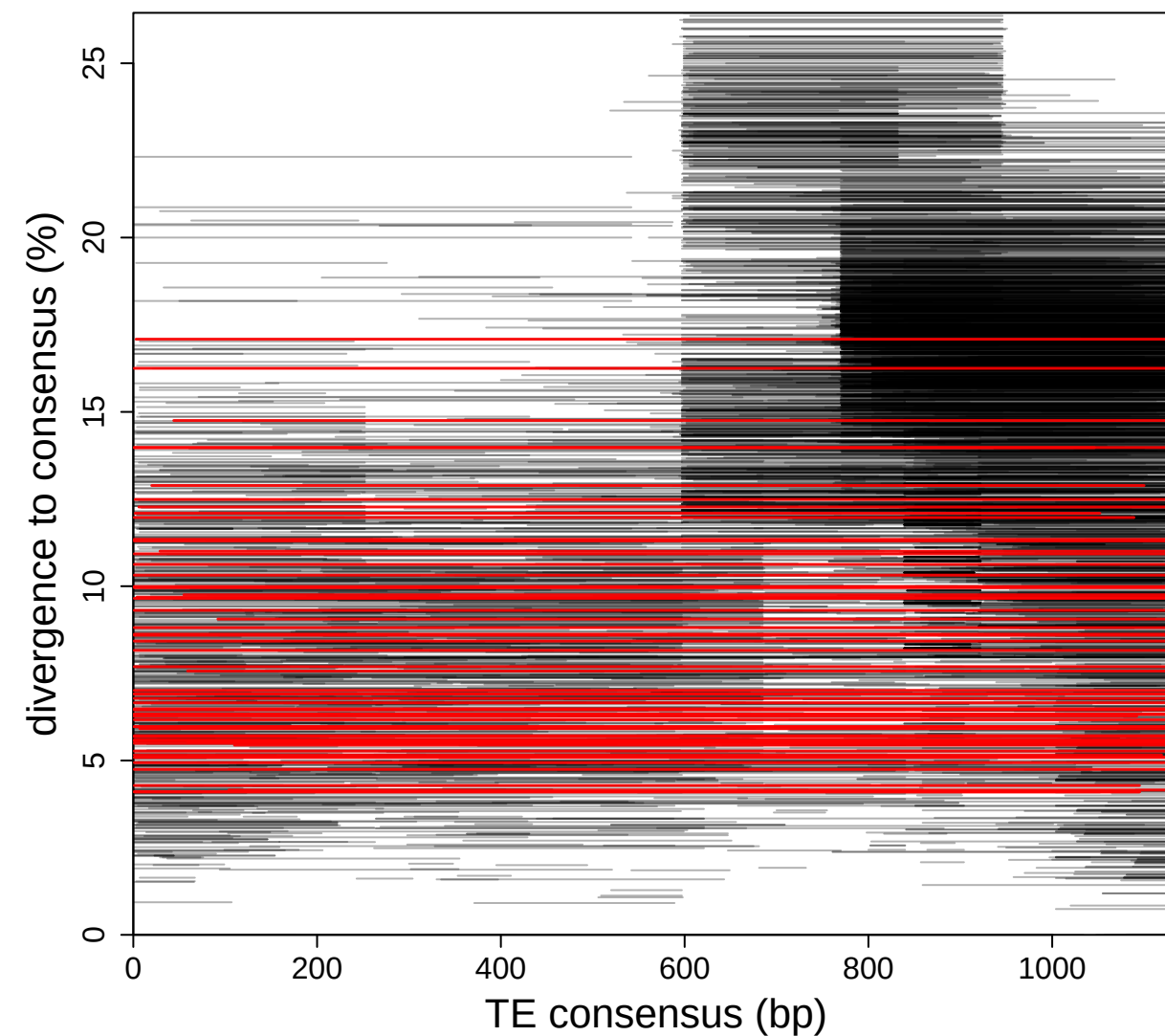


0.fasta.b.bed uf.bed g_2.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 2154bp; fragments: 5487; full length: 0 (>=1938.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

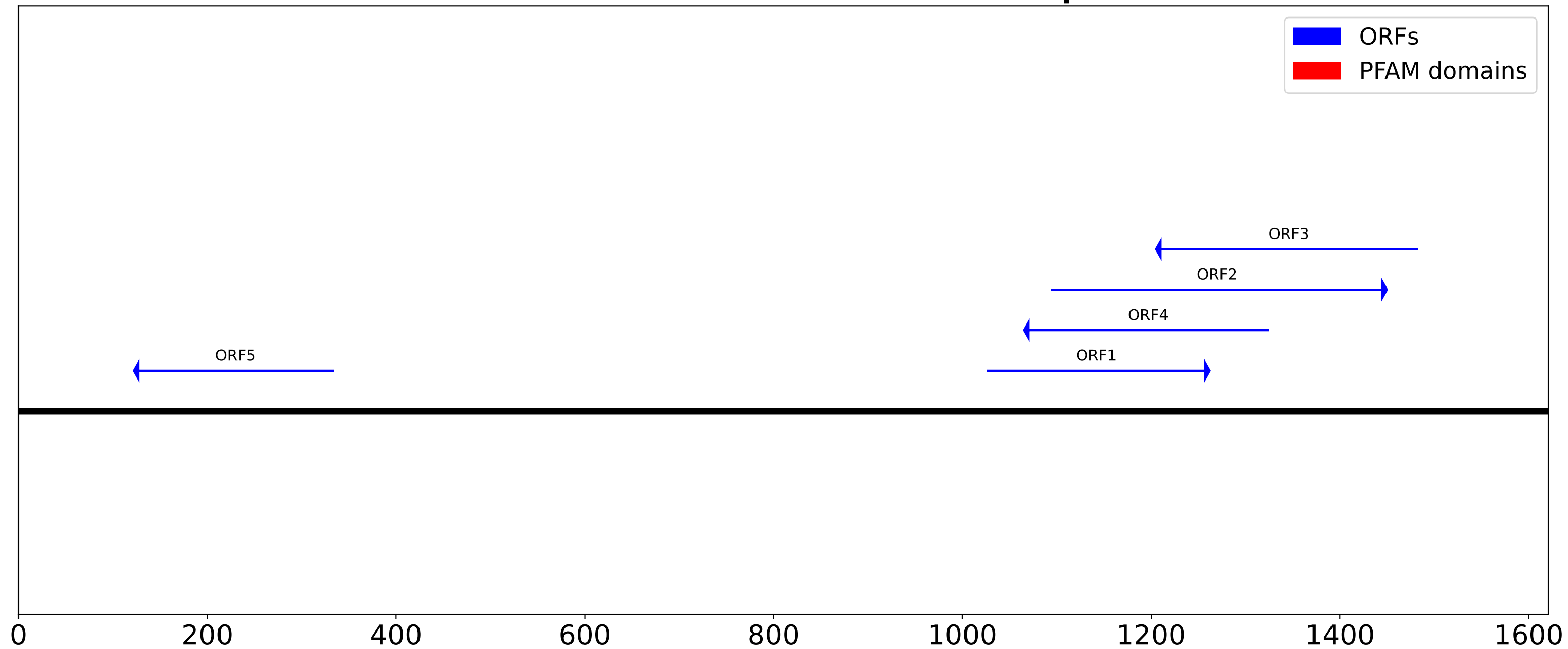


TE: rnd_4_family_1130
size: 1138bp; fragments: 4591; full length: 62 (≥ 1024.2 bp)

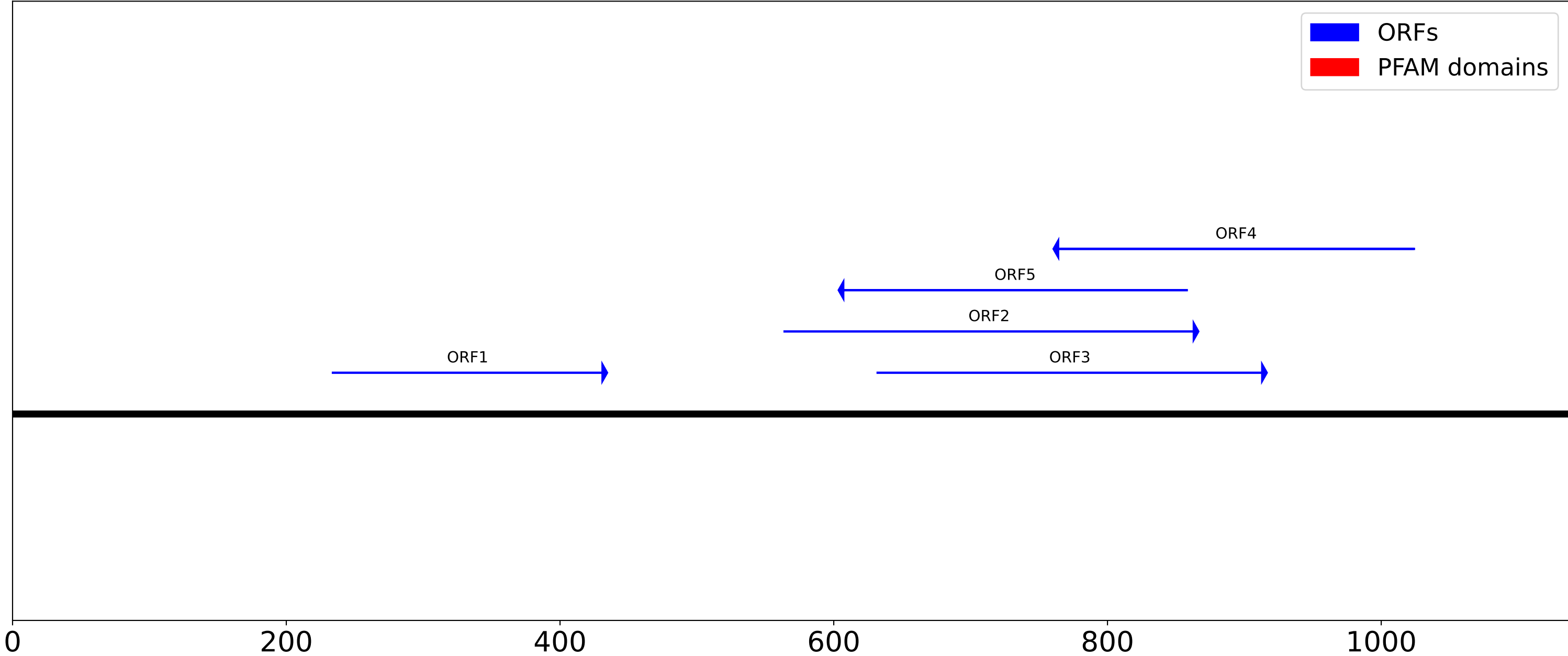


Before TEtrimmer 1138 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

