

MSA length = 2510

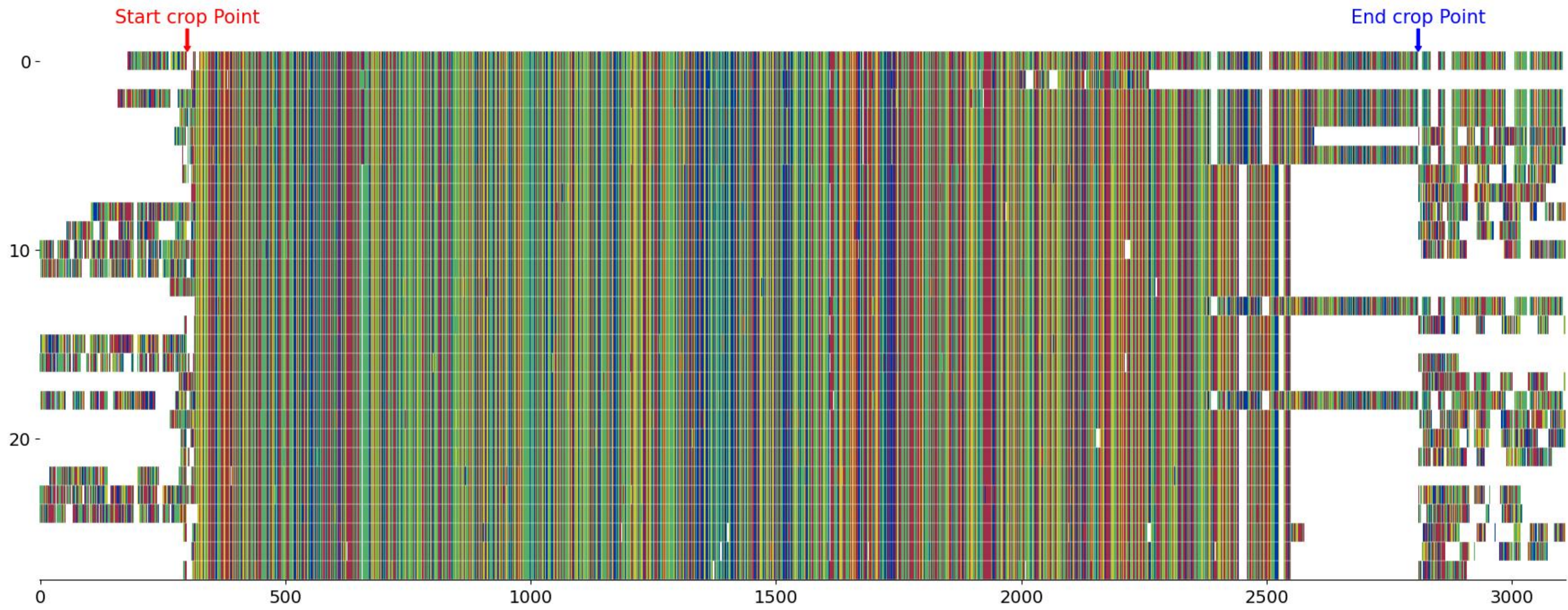
Start crop Point

End crop Point



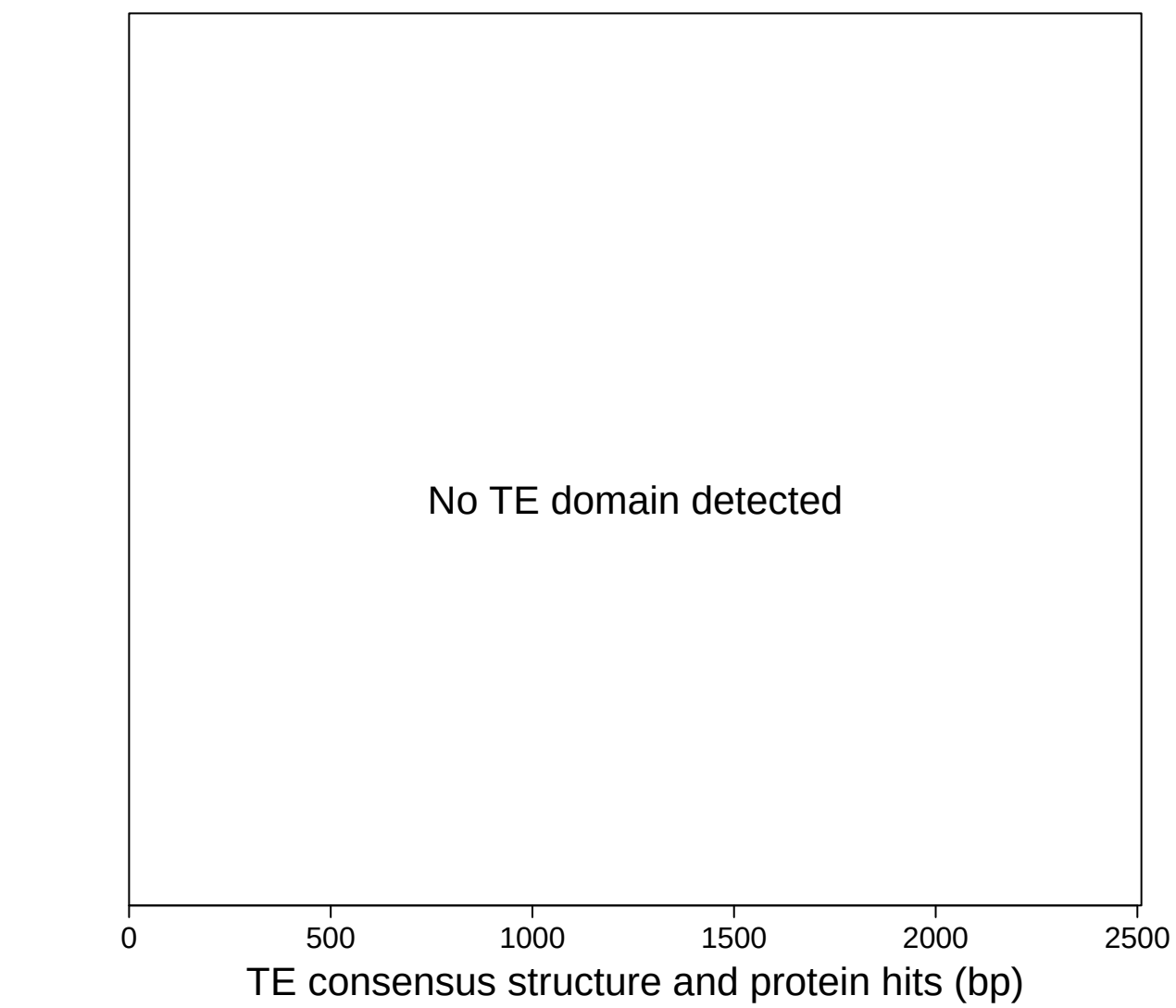
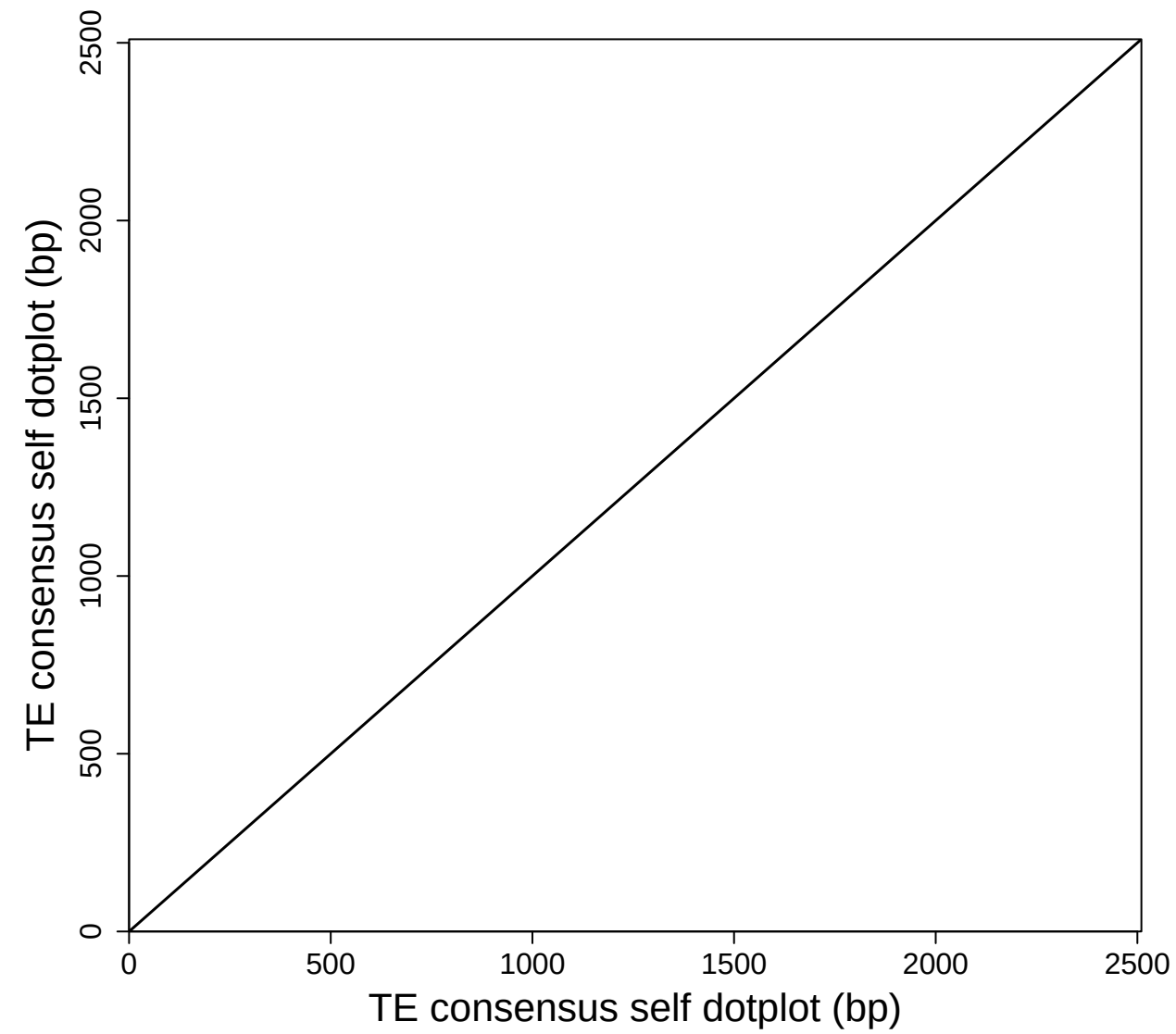
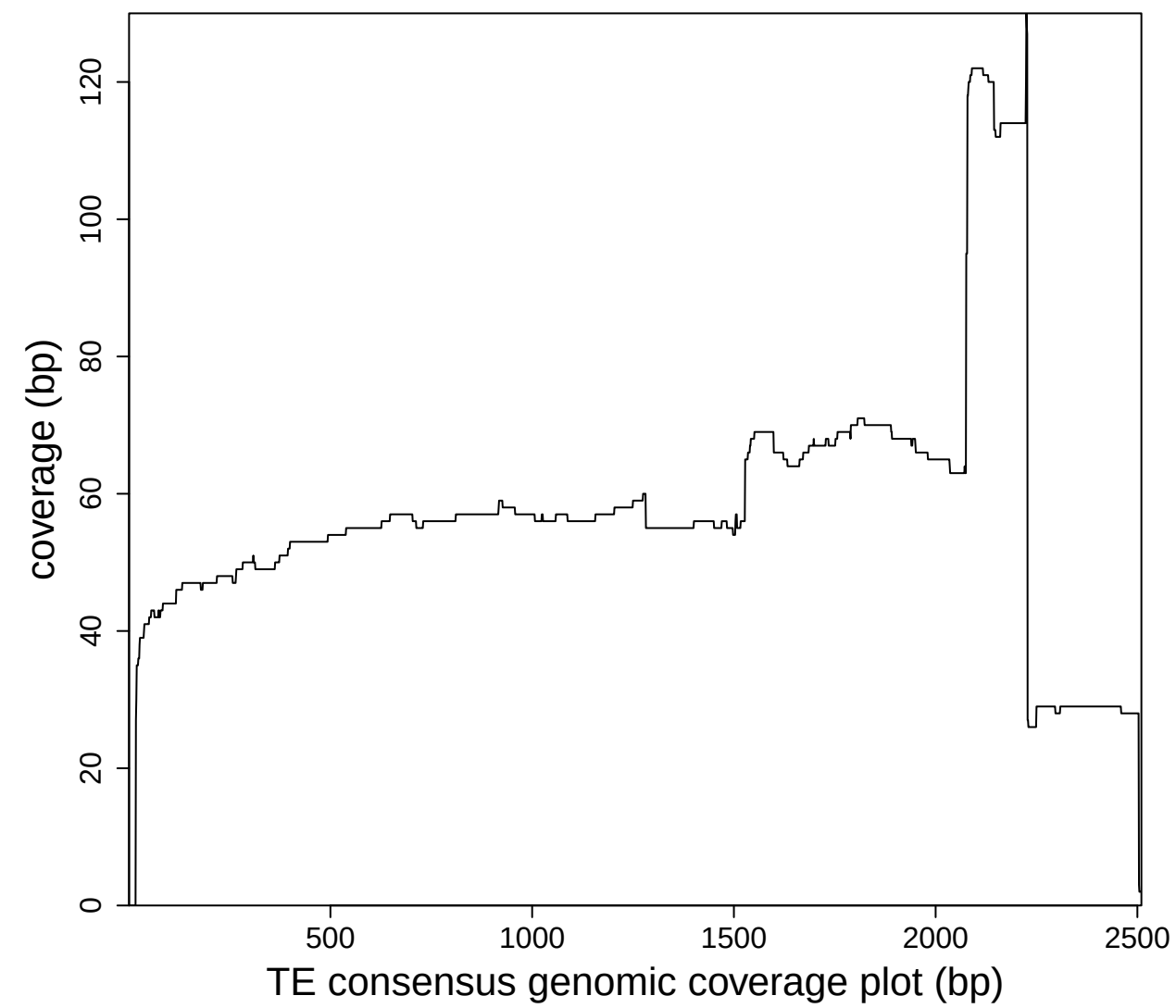
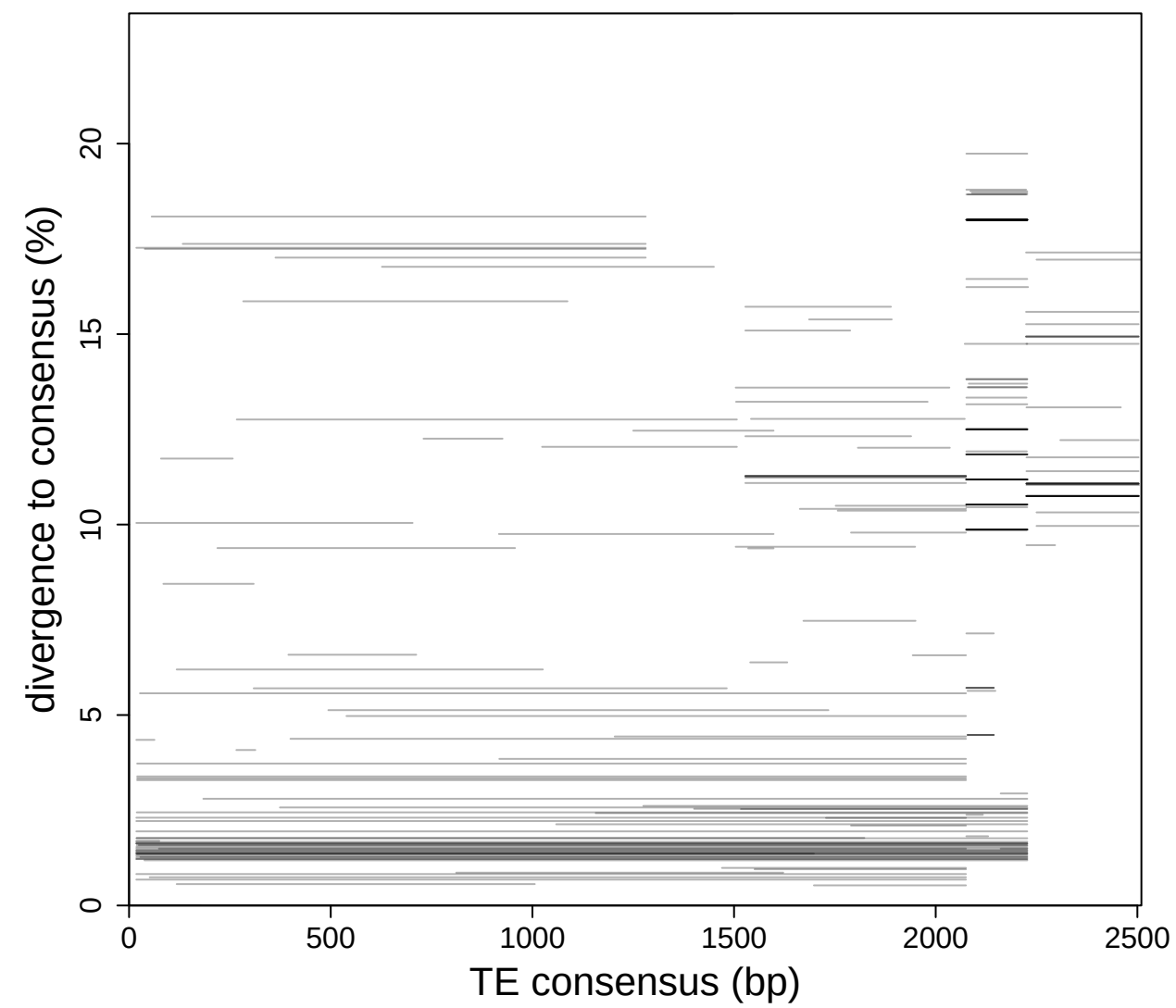
Start crop Point

End crop Point



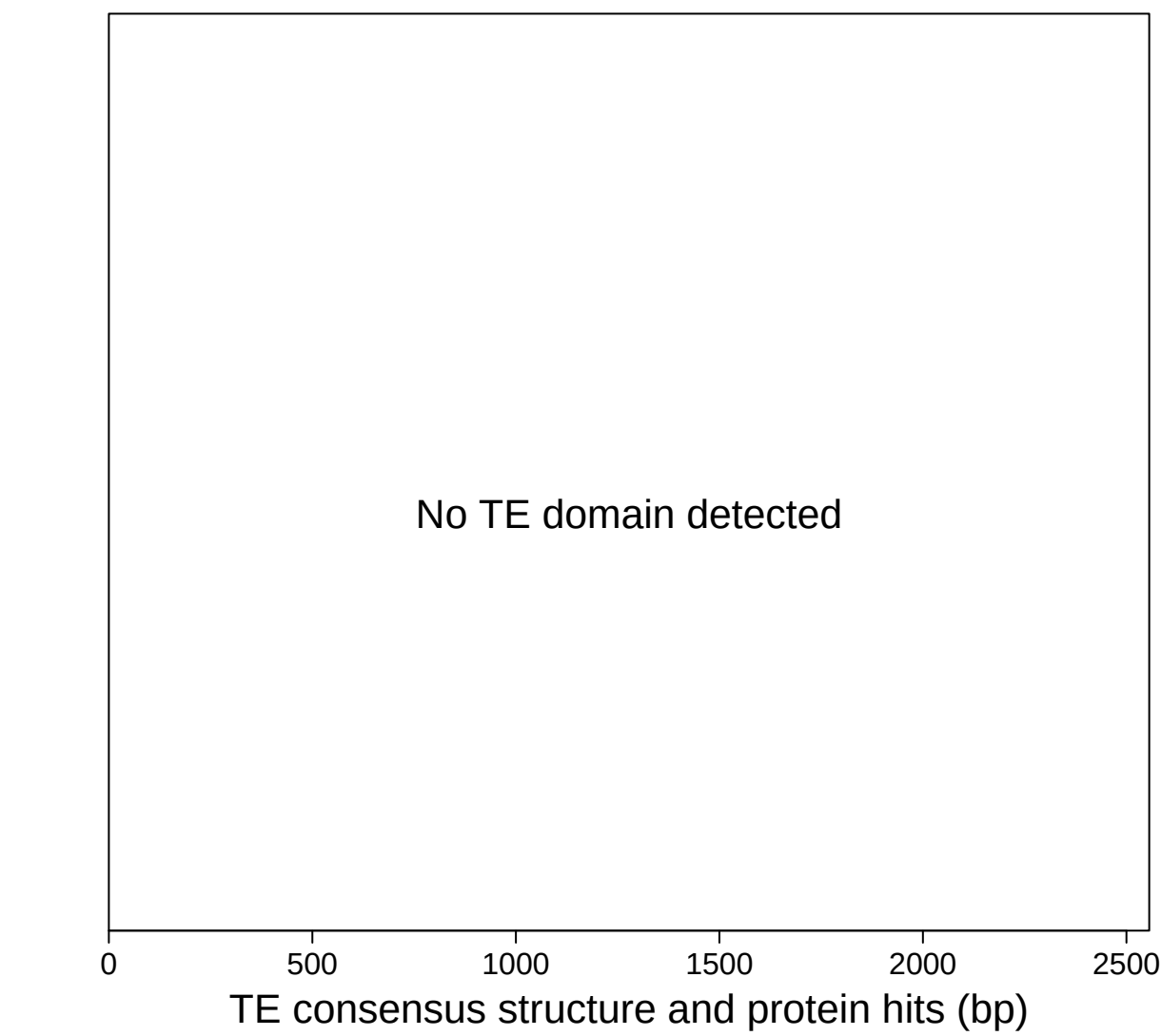
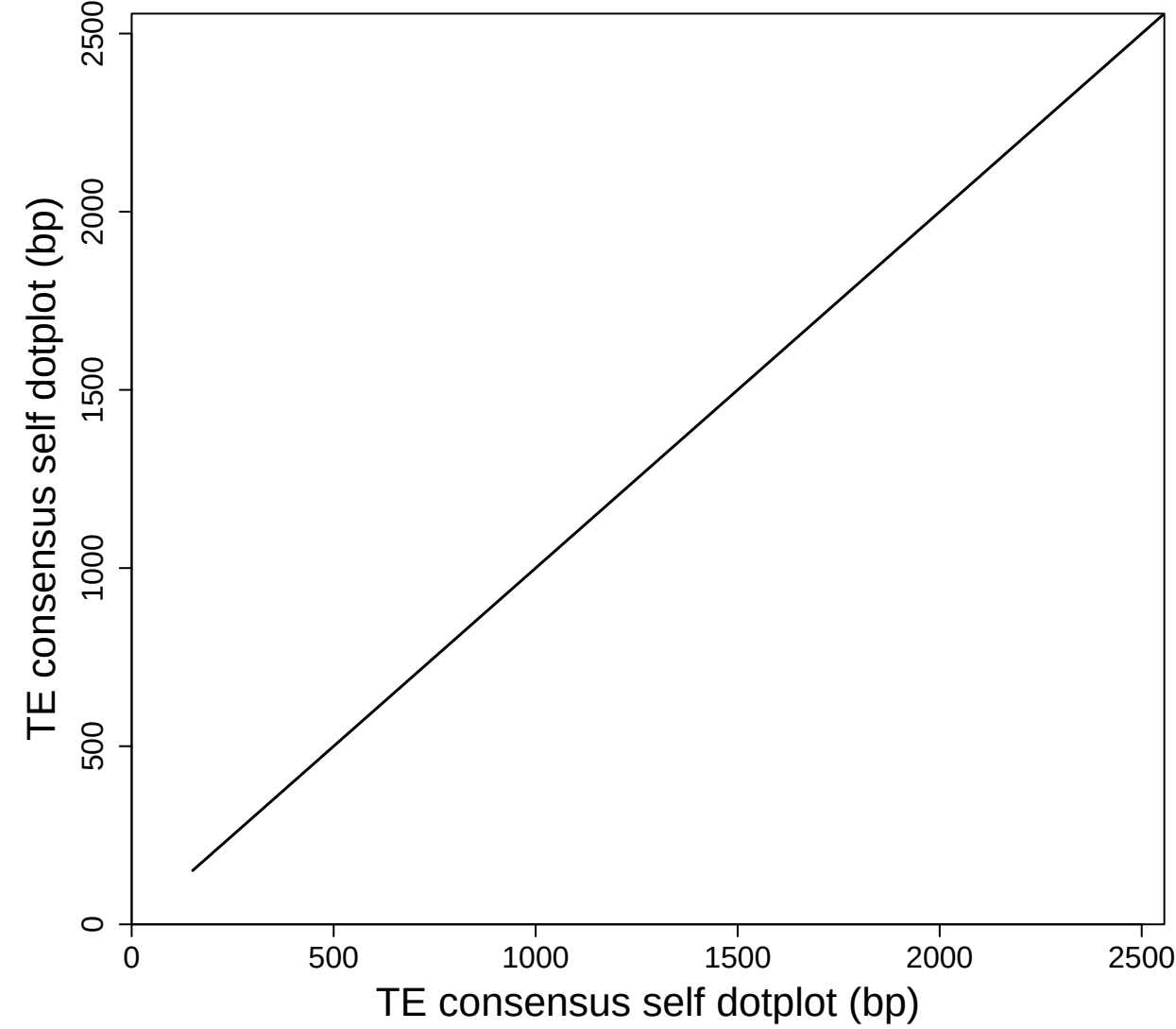
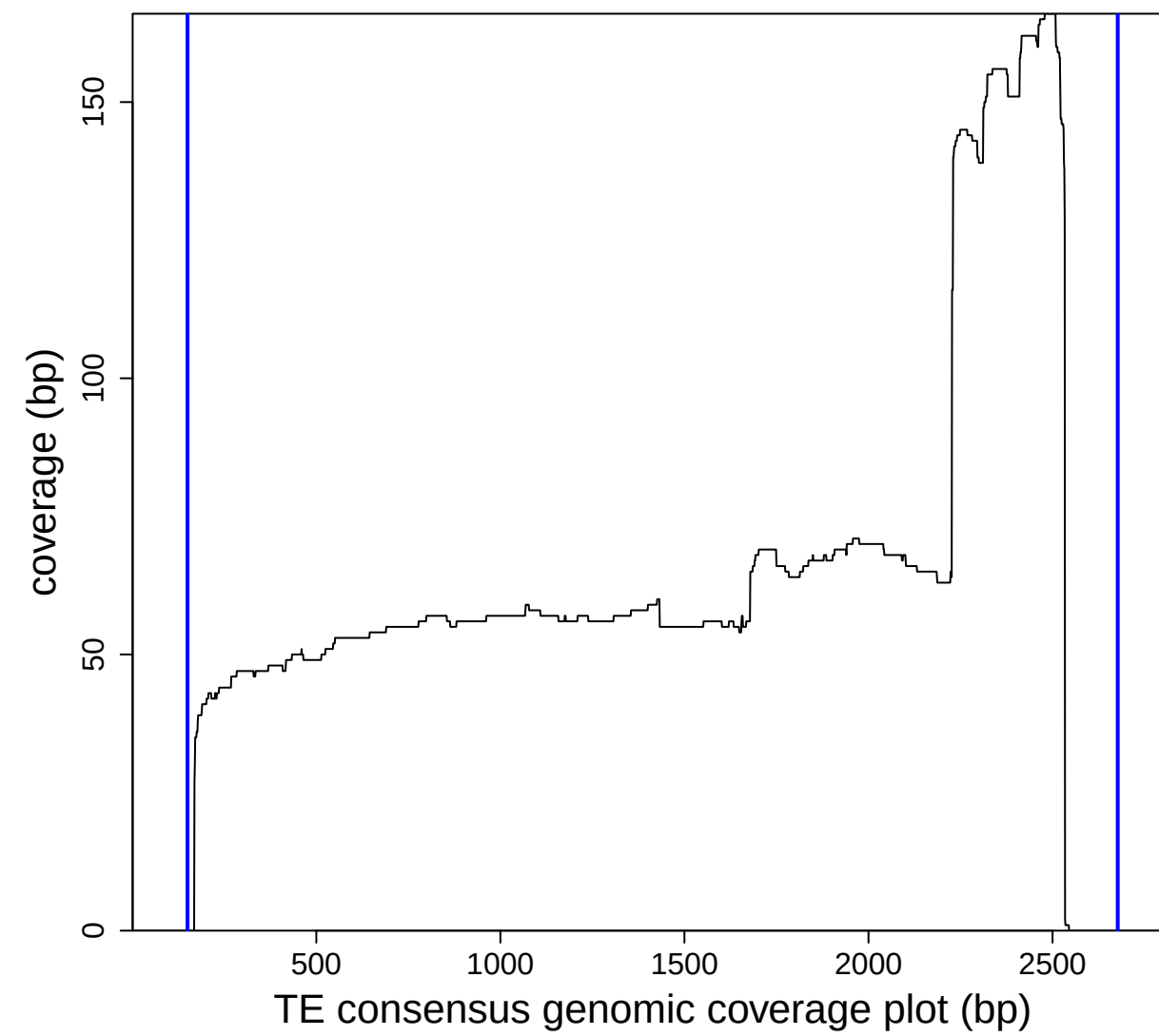
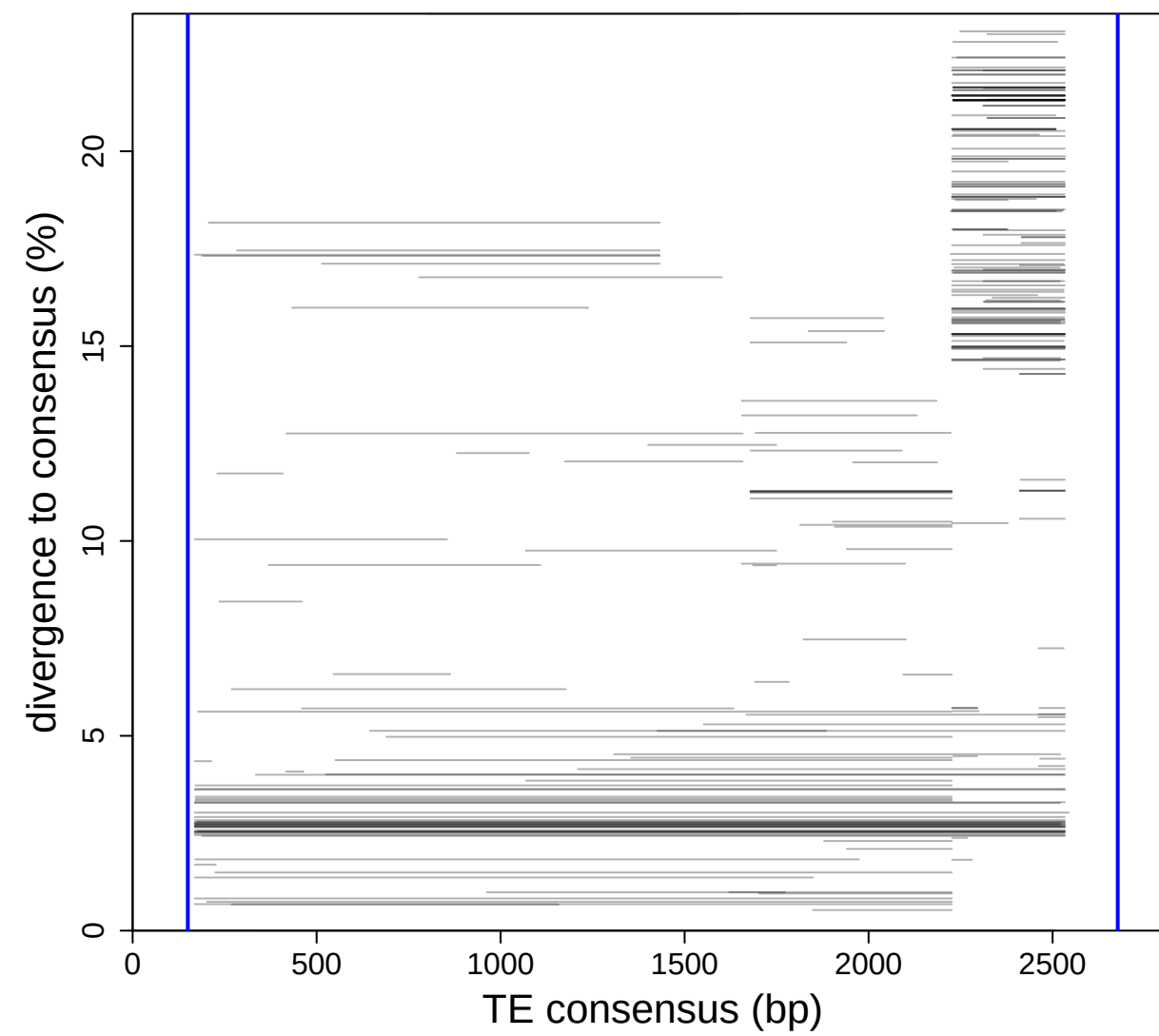
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 2510bp; fragments: 224; full length: 0 (>=2259bp)

After TEtrimmer 2510 bp

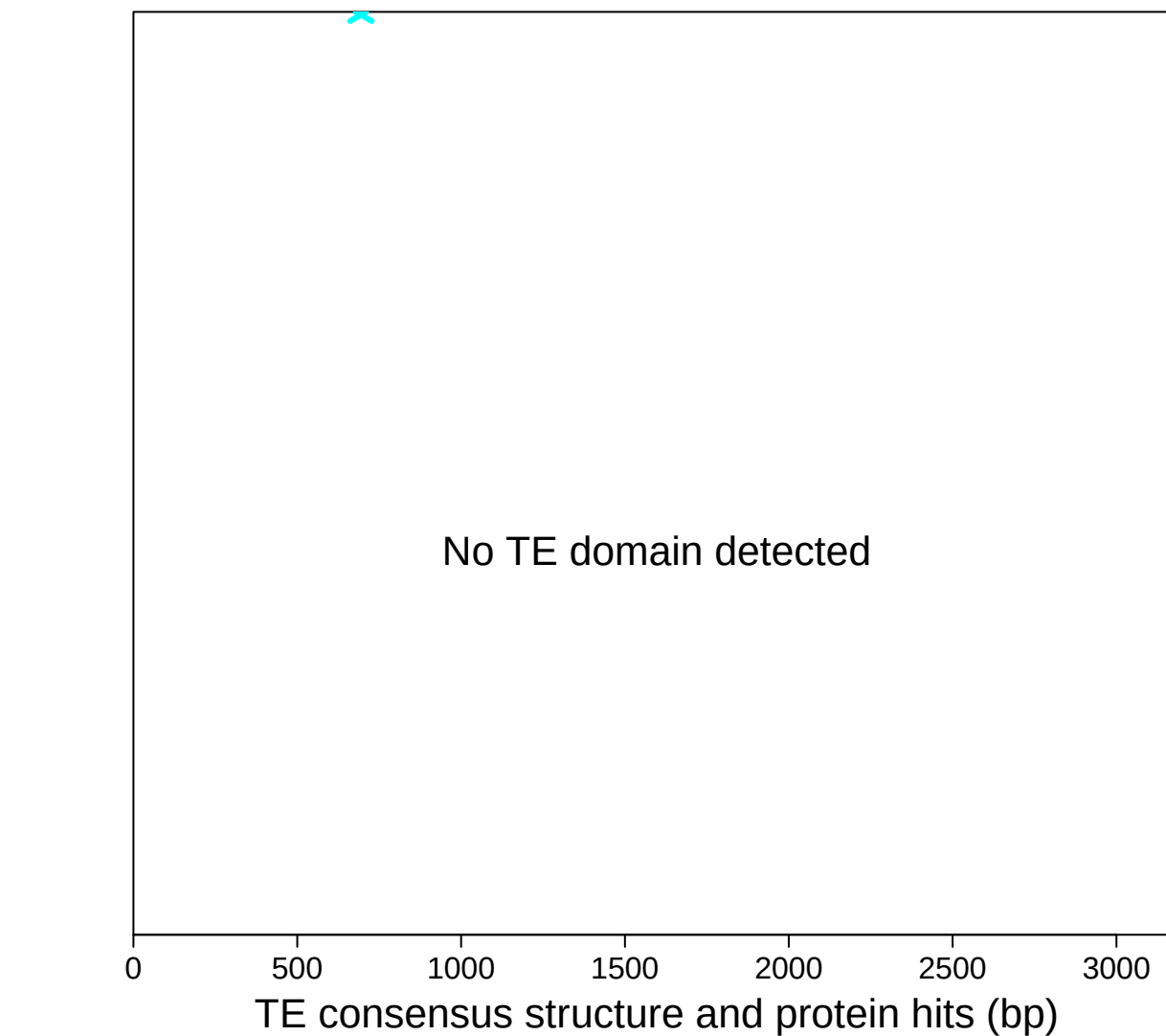
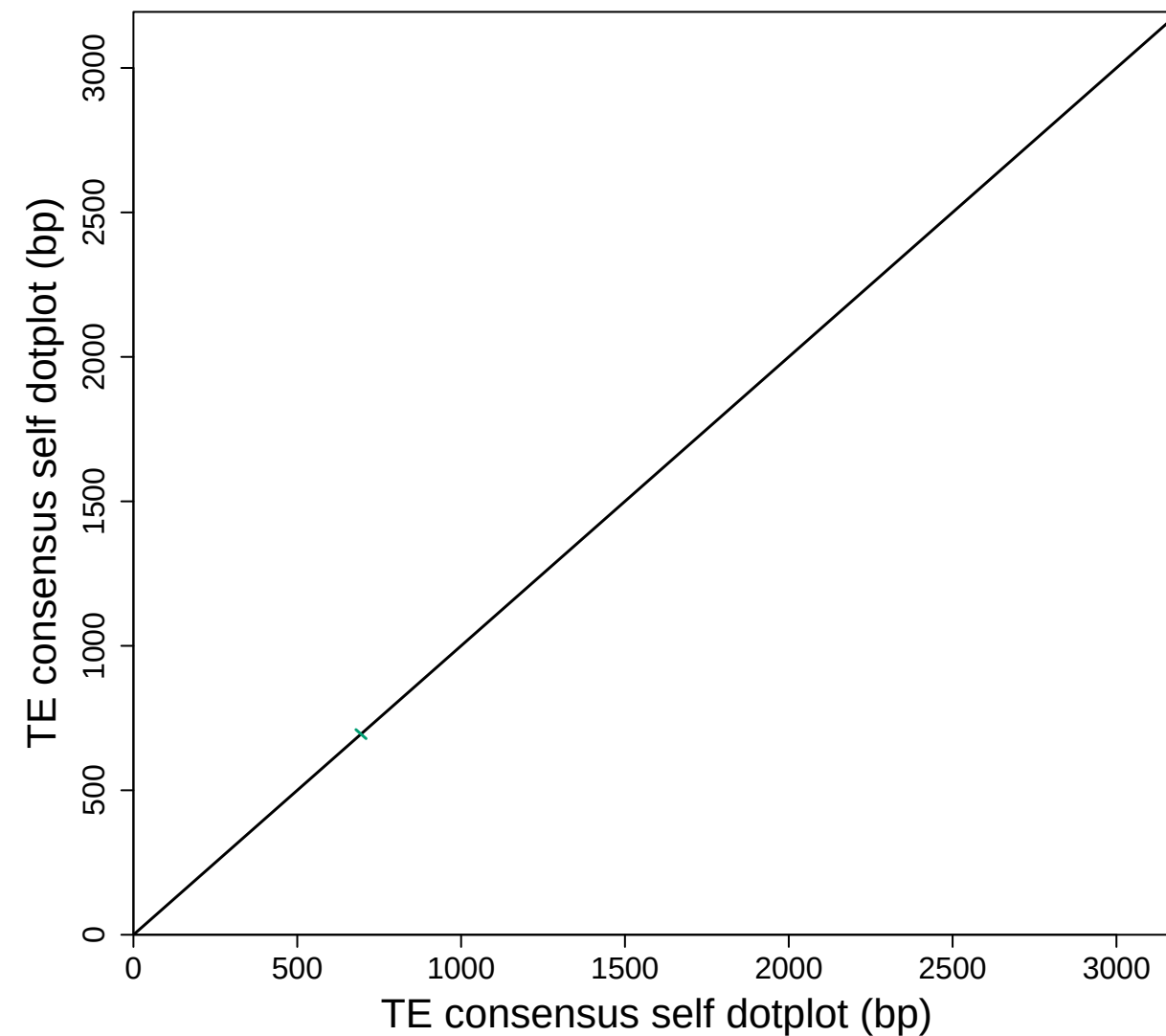
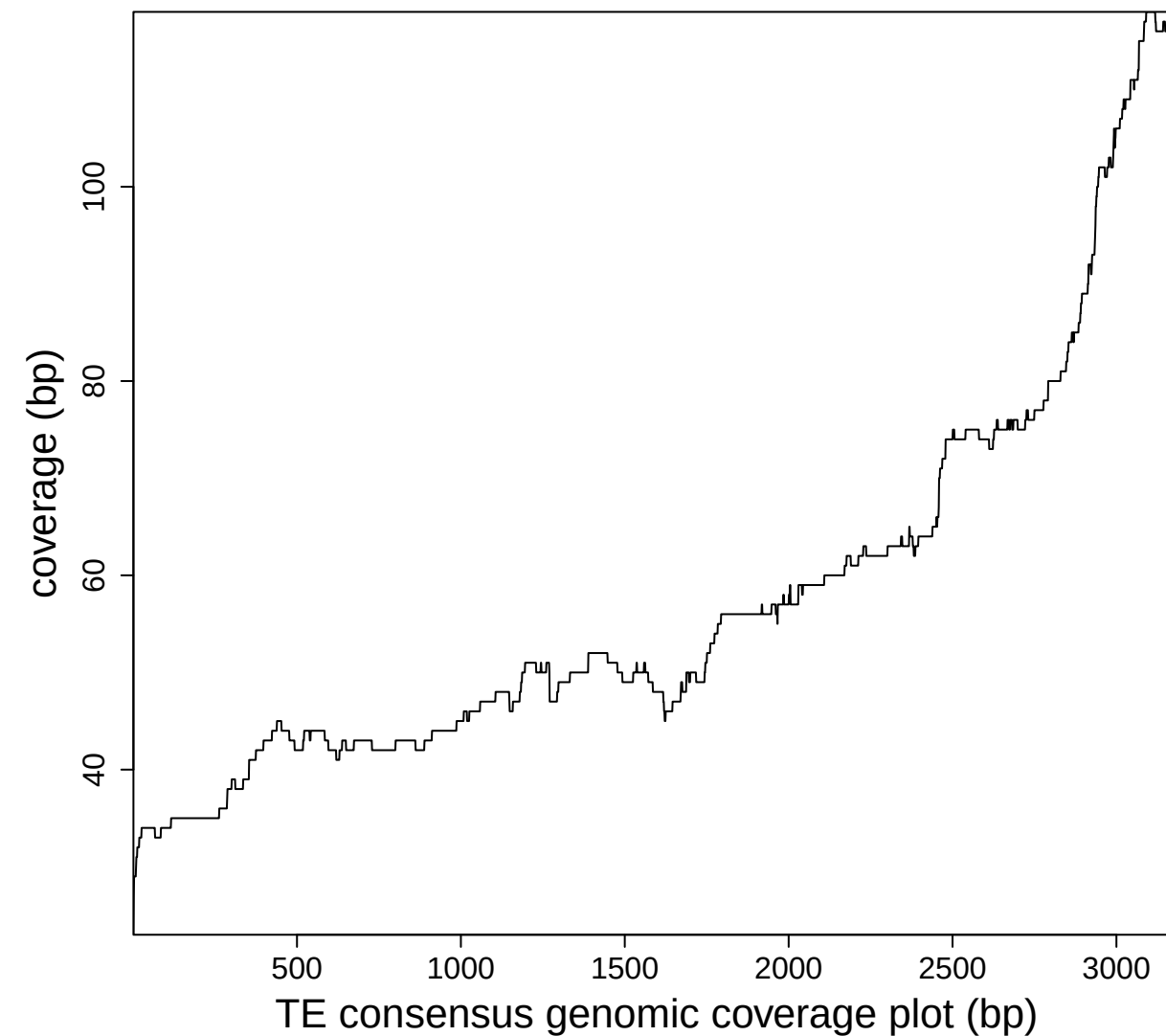
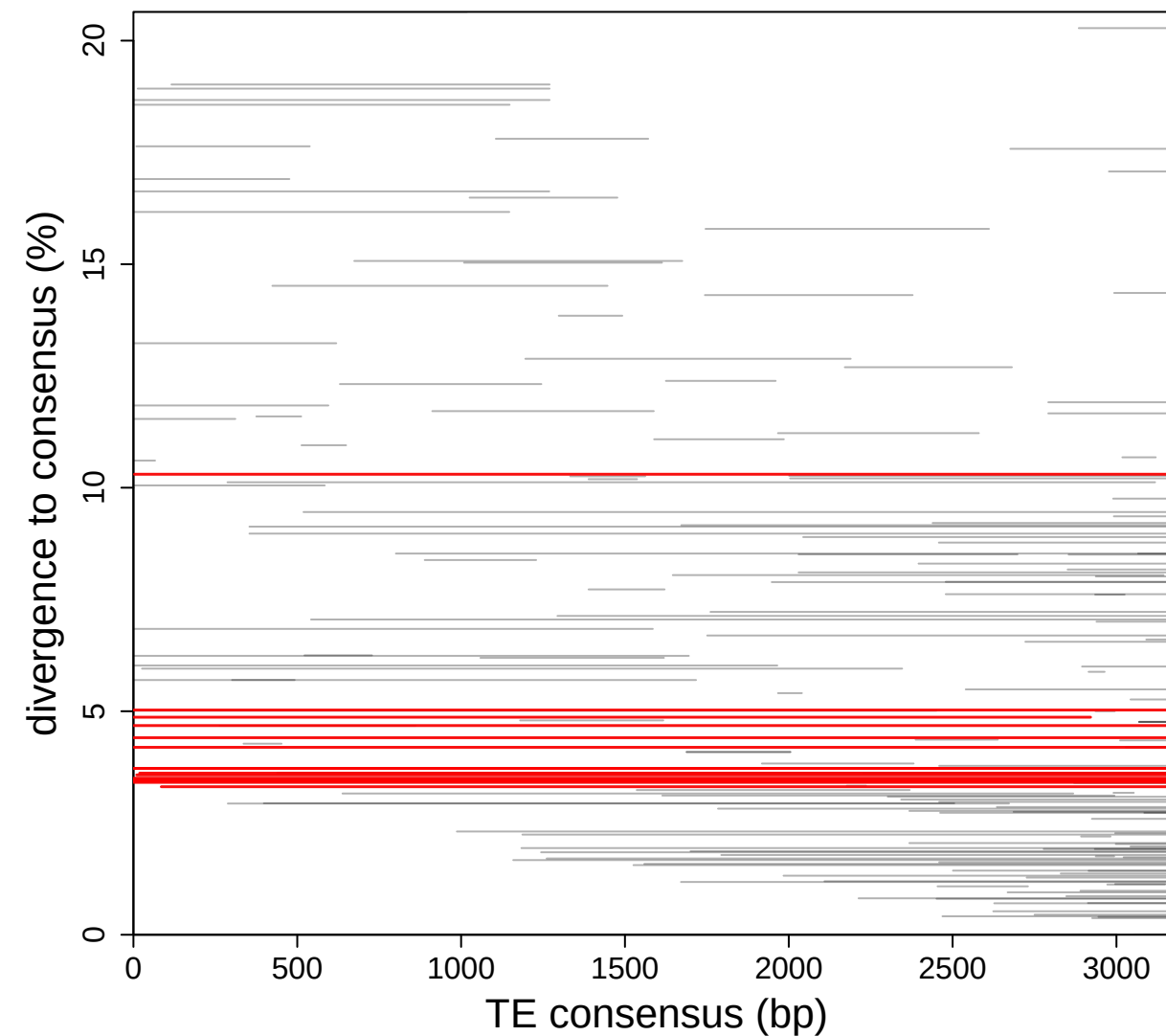


8.fasta.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcIn.fa aln.fa cl.f
size: 2827bp; fragments: 250; full length: 0 (>=2544.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

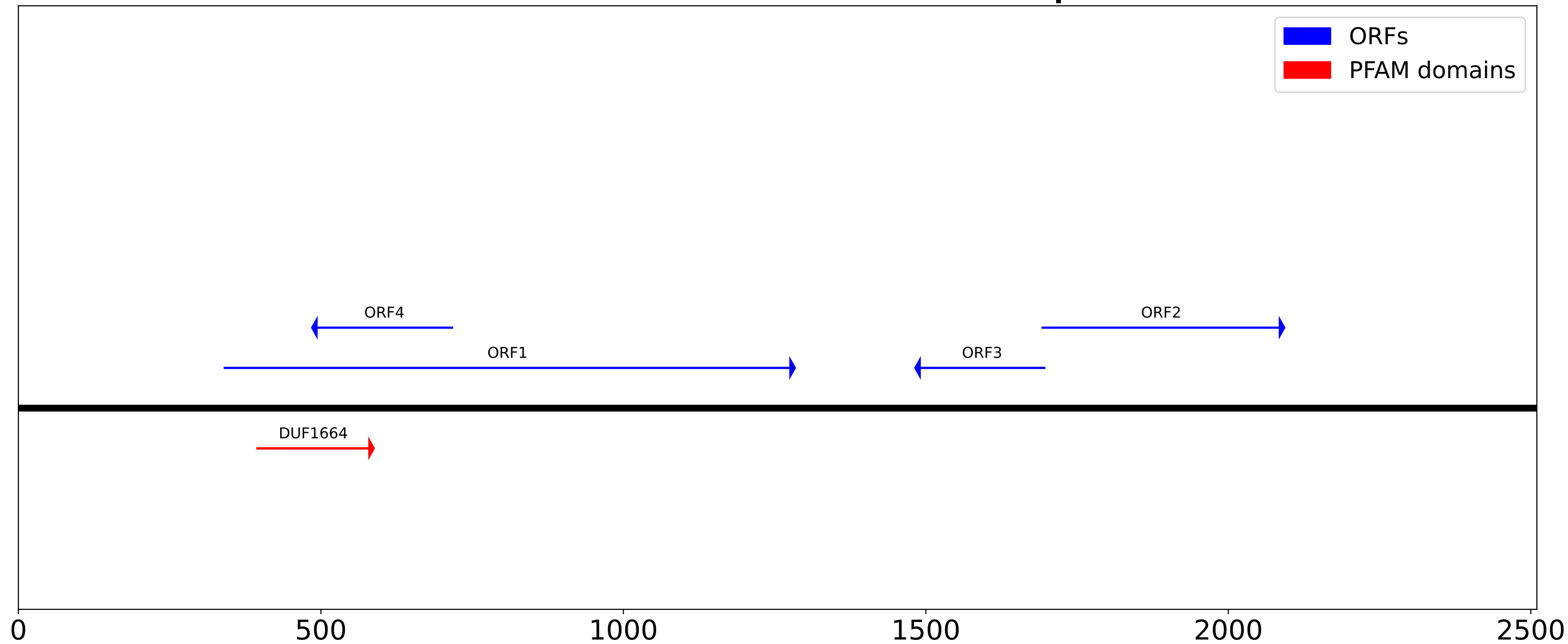


TE: rnd_5_family_5268
size: 3194bp; fragments: 188; full length: 16 (≥ 2874.6 bp)

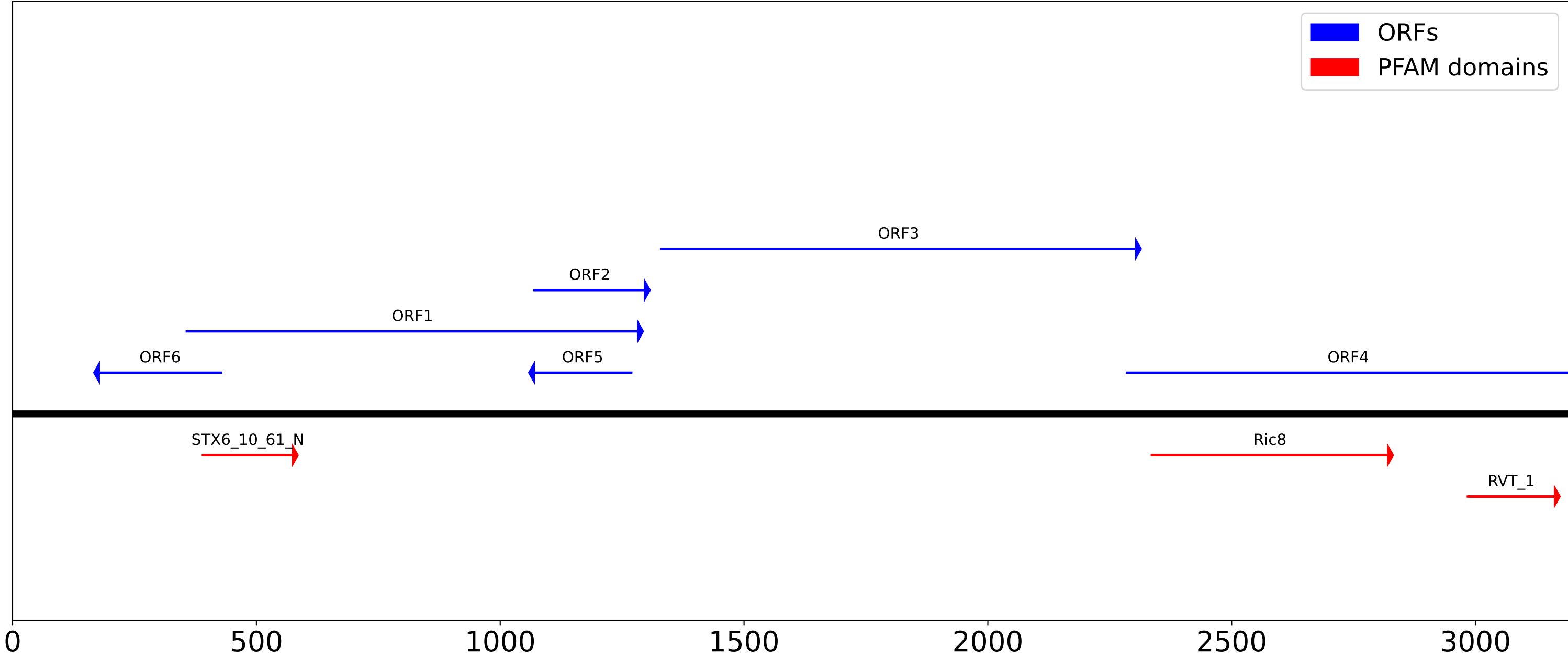


Before TEtrimmer 3194 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

