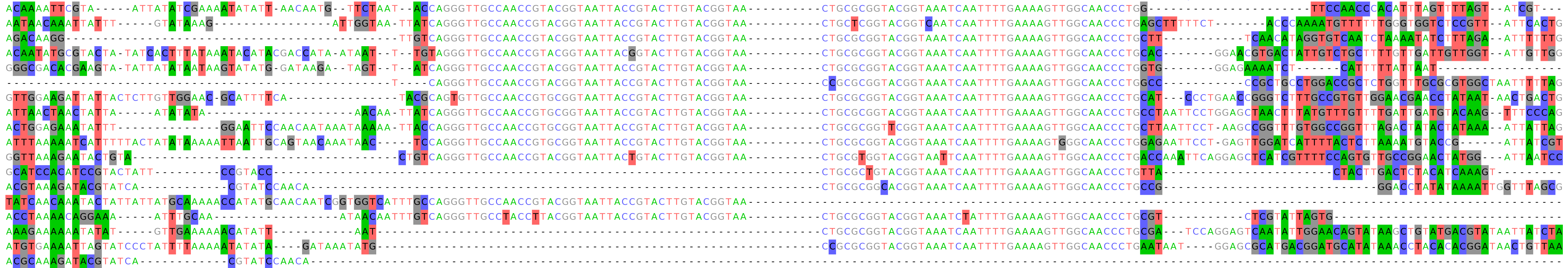


MSA length = 857

Start crop Point

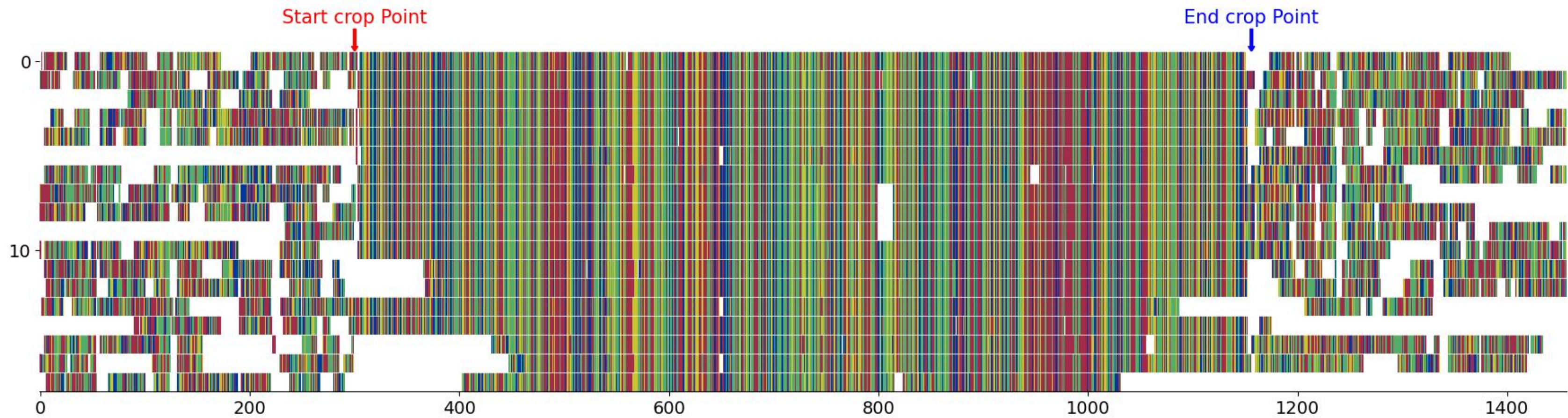


End crop Point



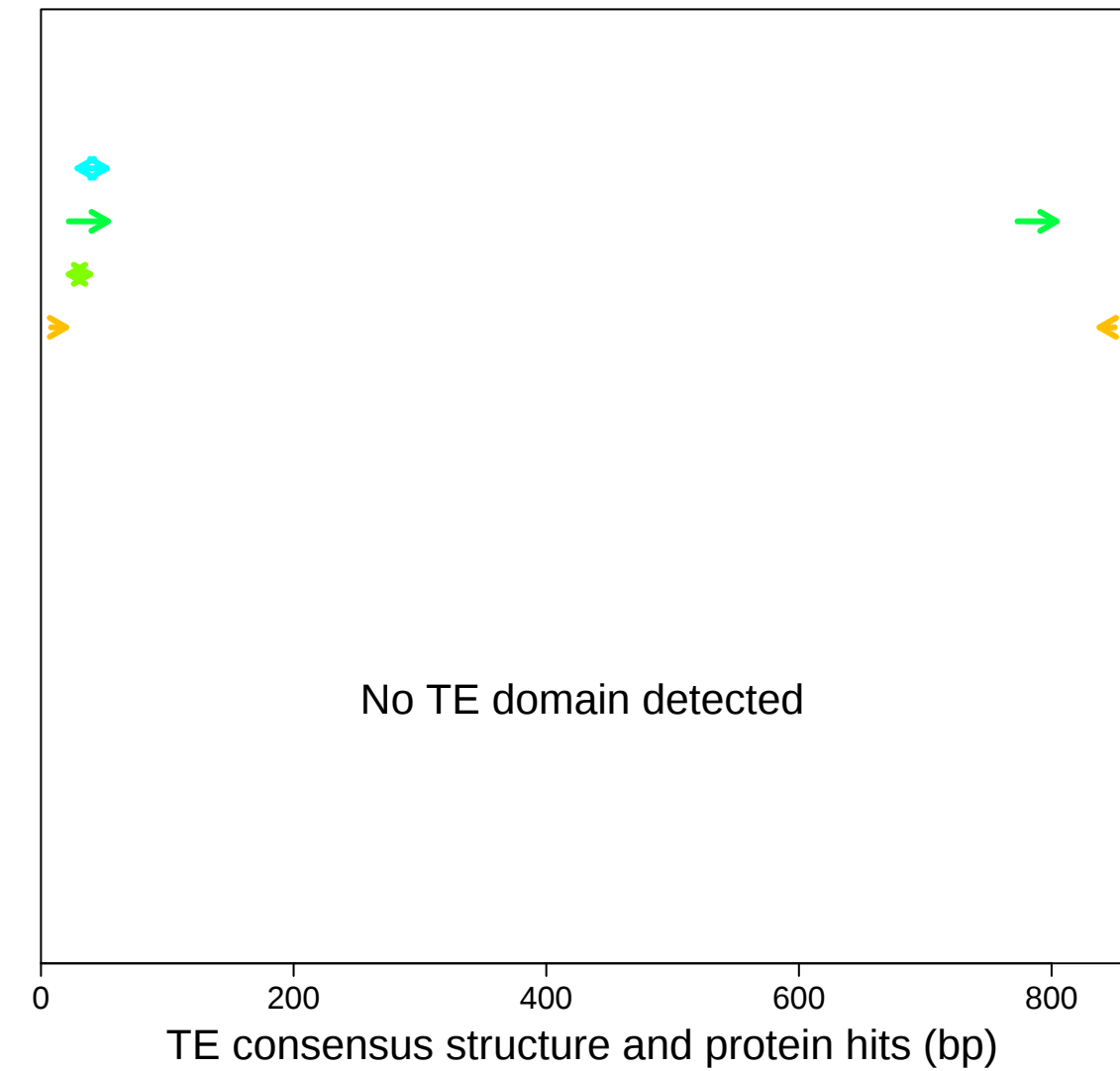
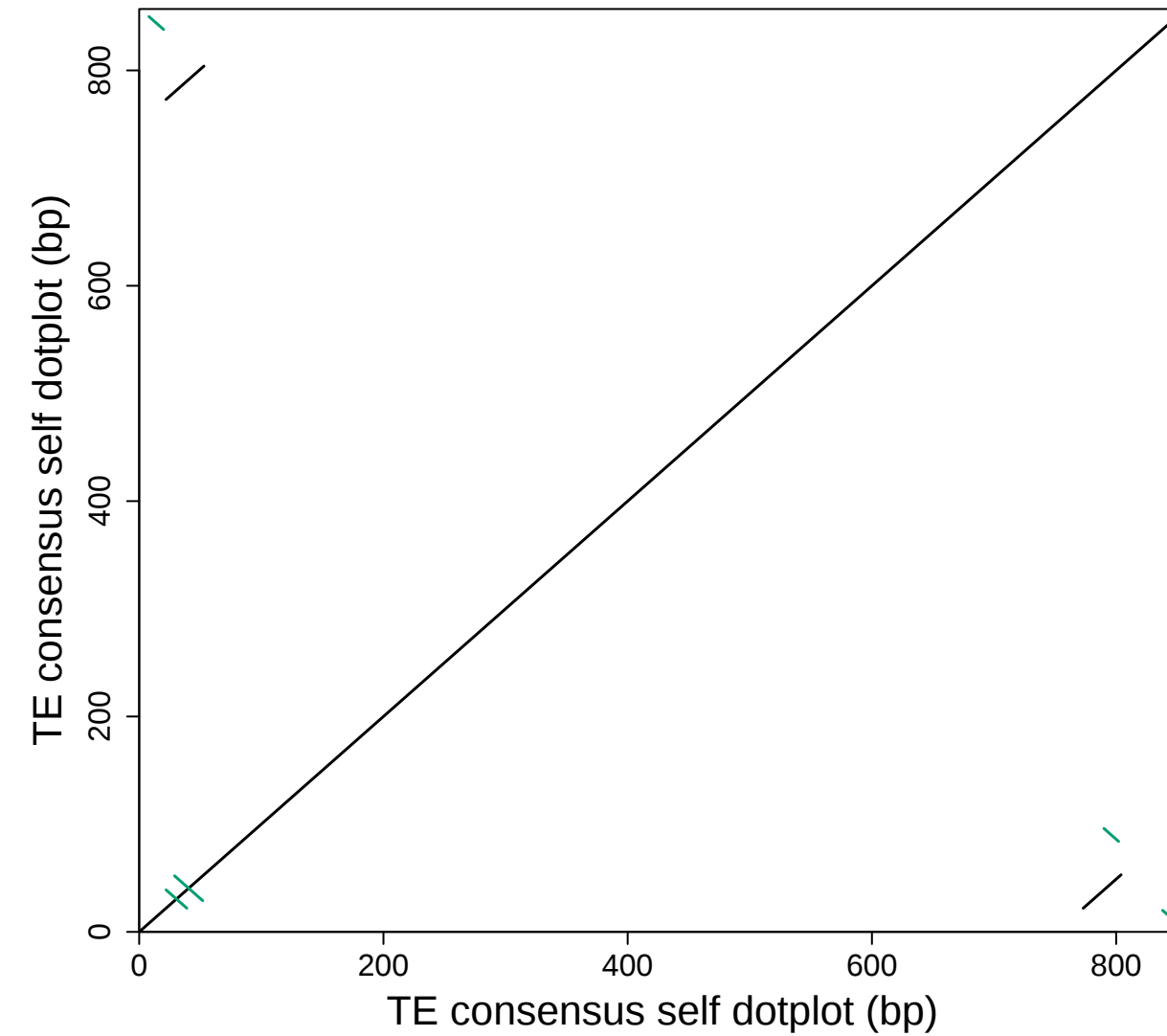
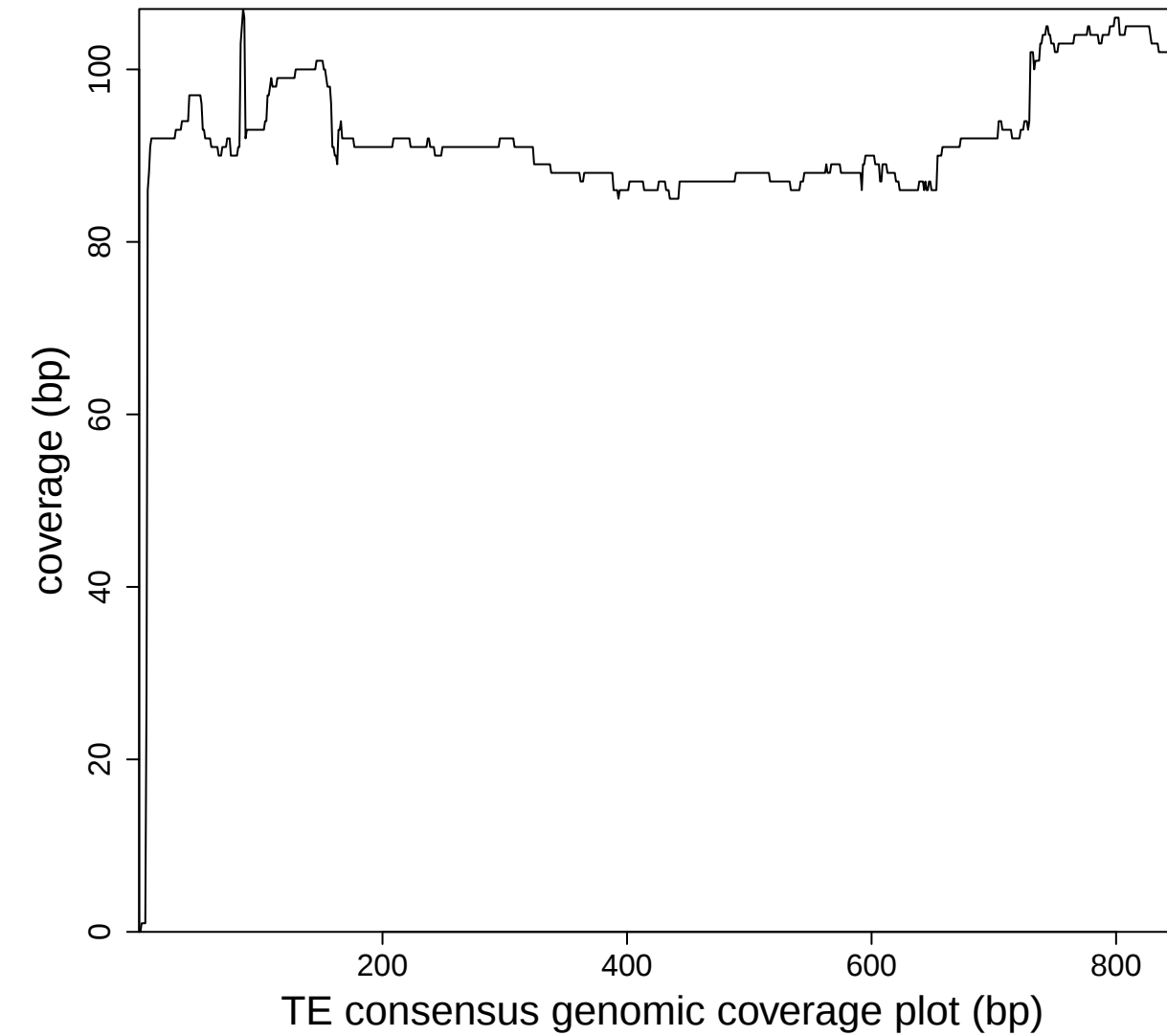
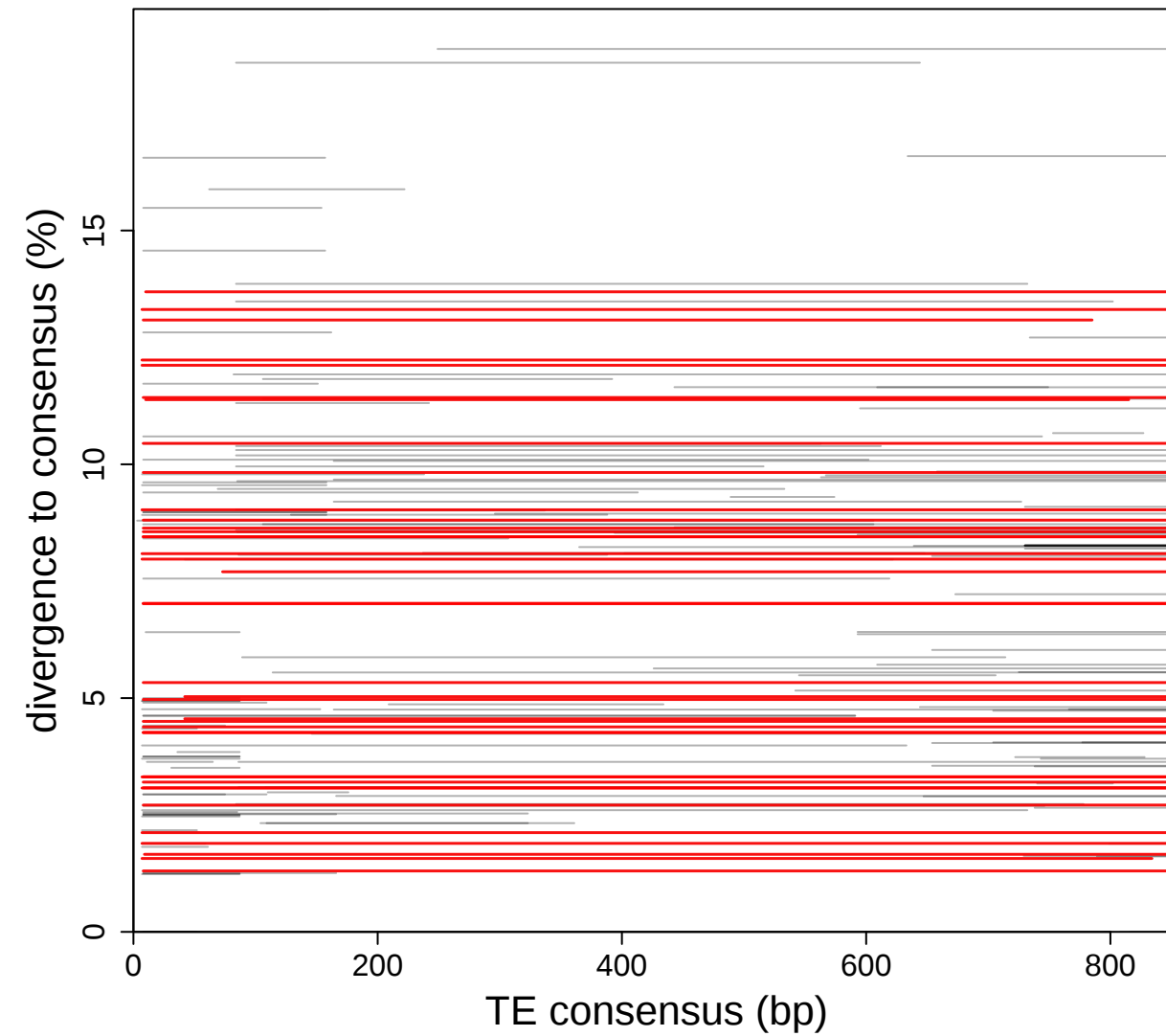
Start crop Point

End crop Point



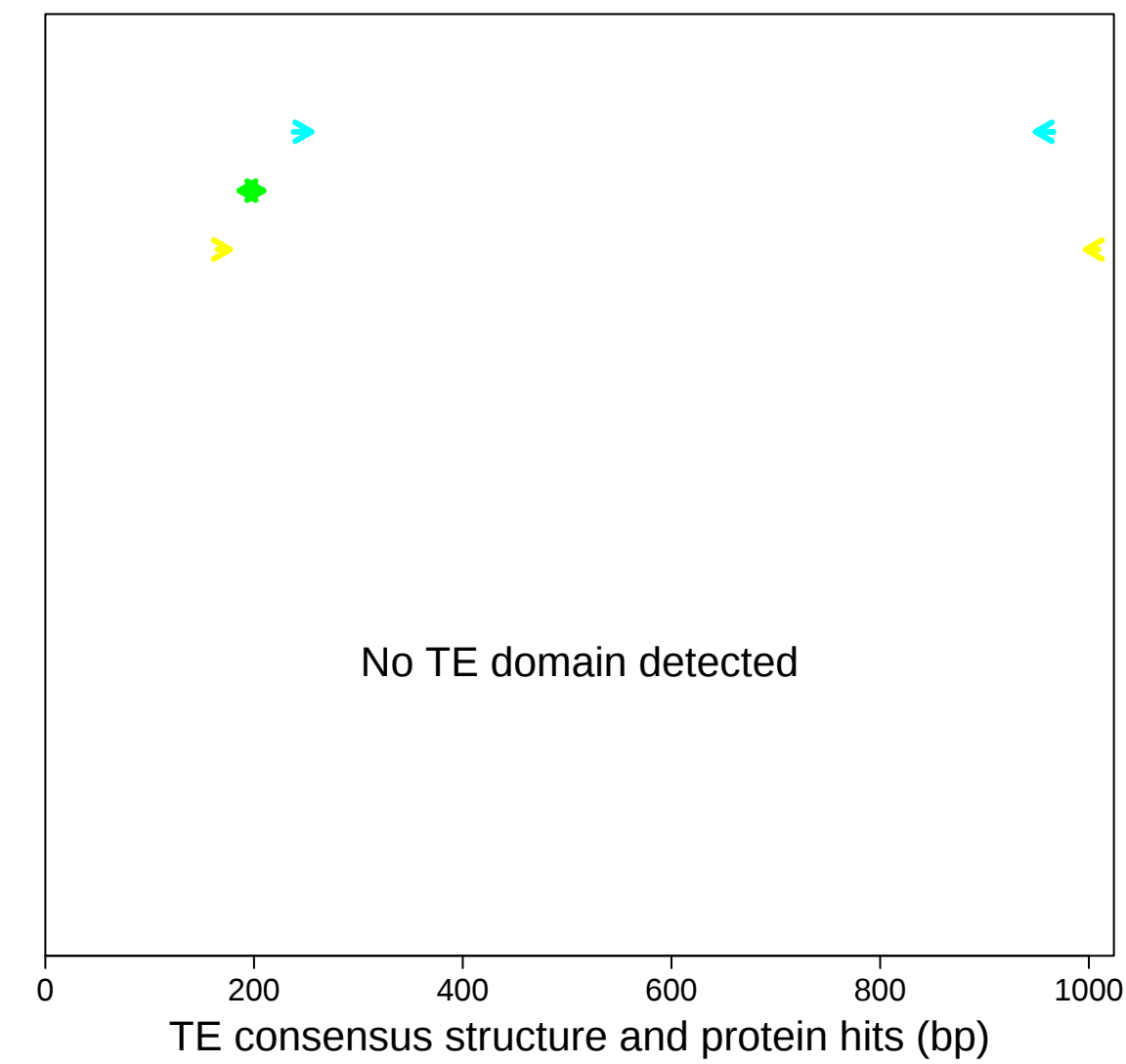
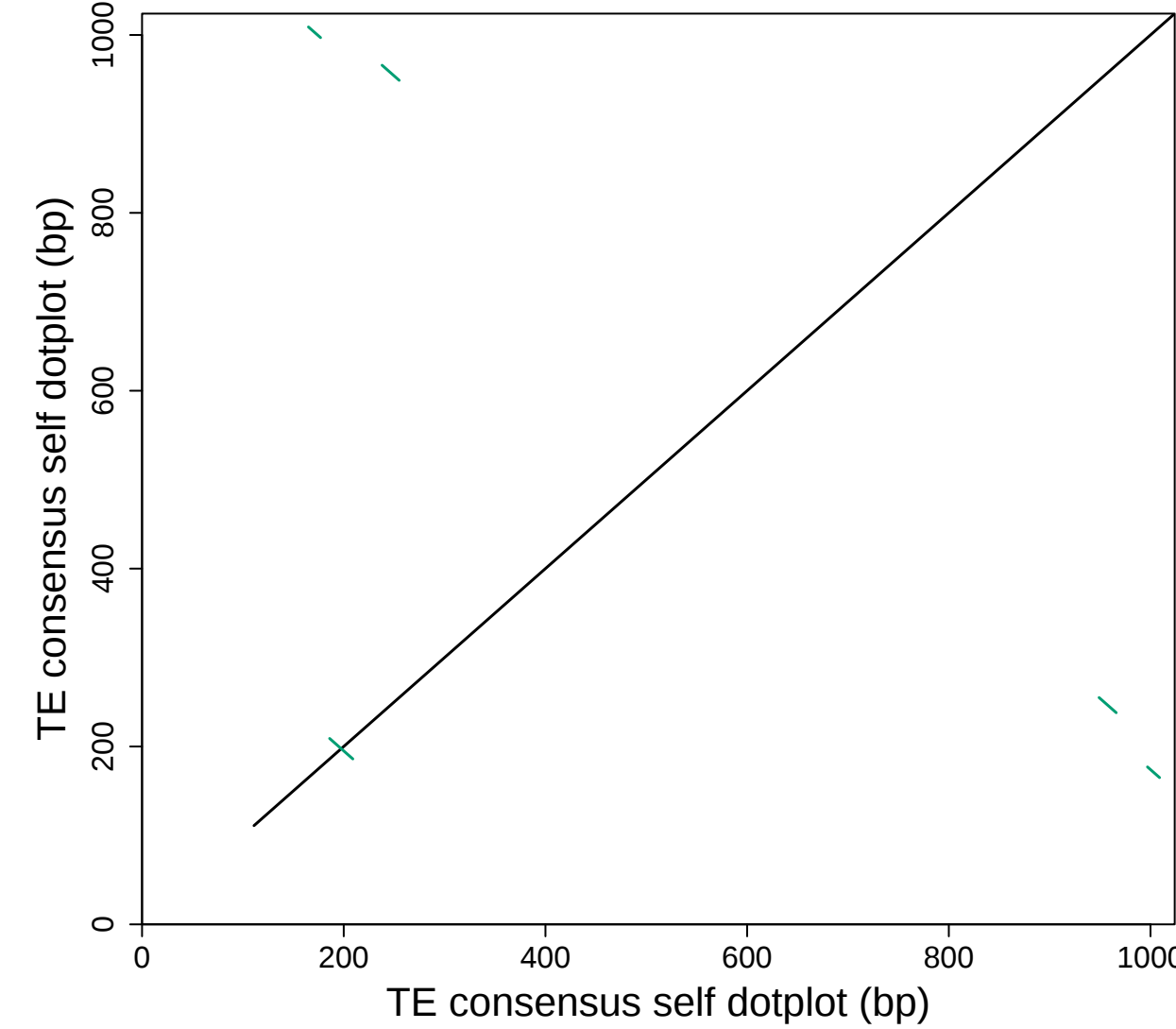
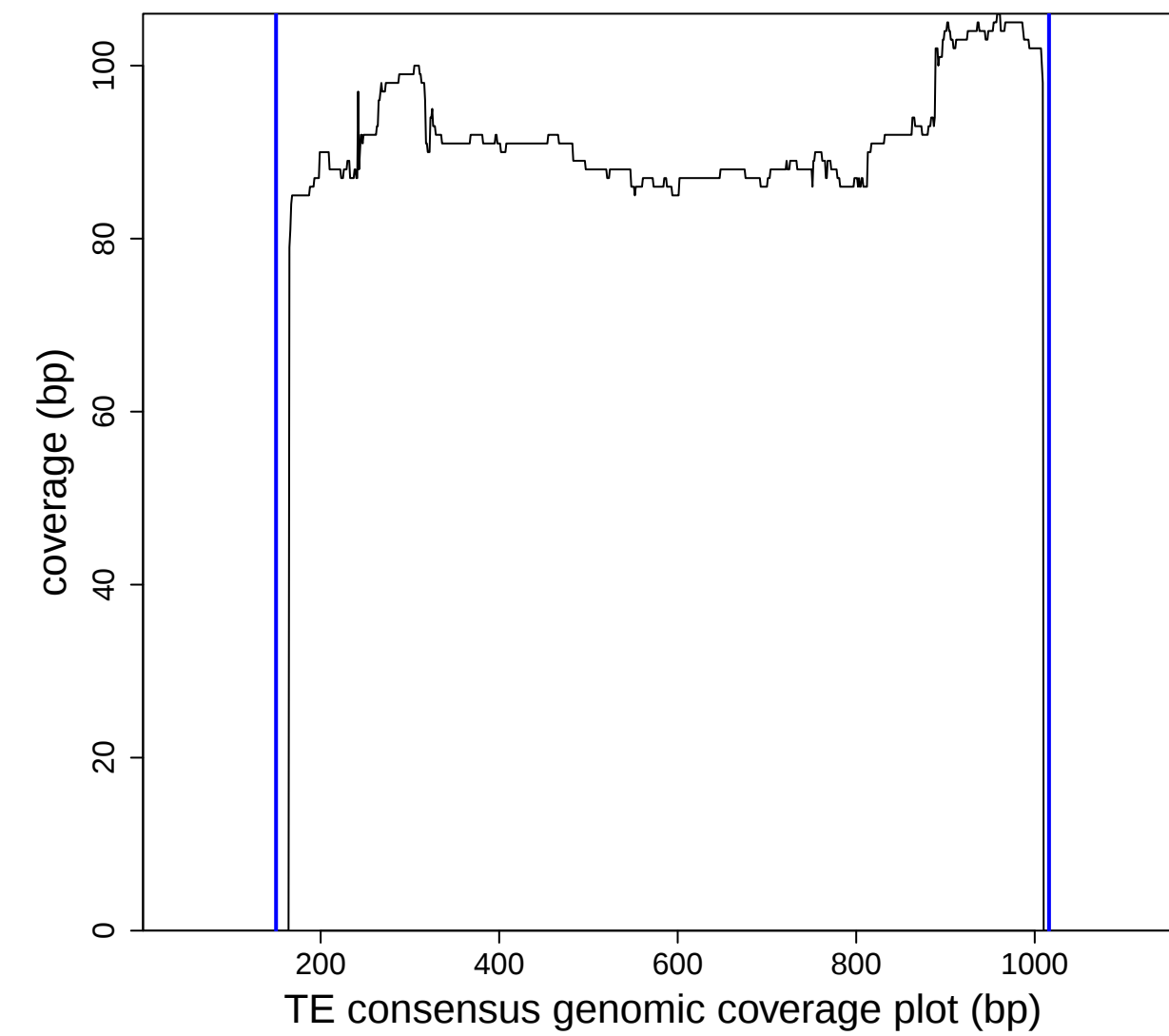
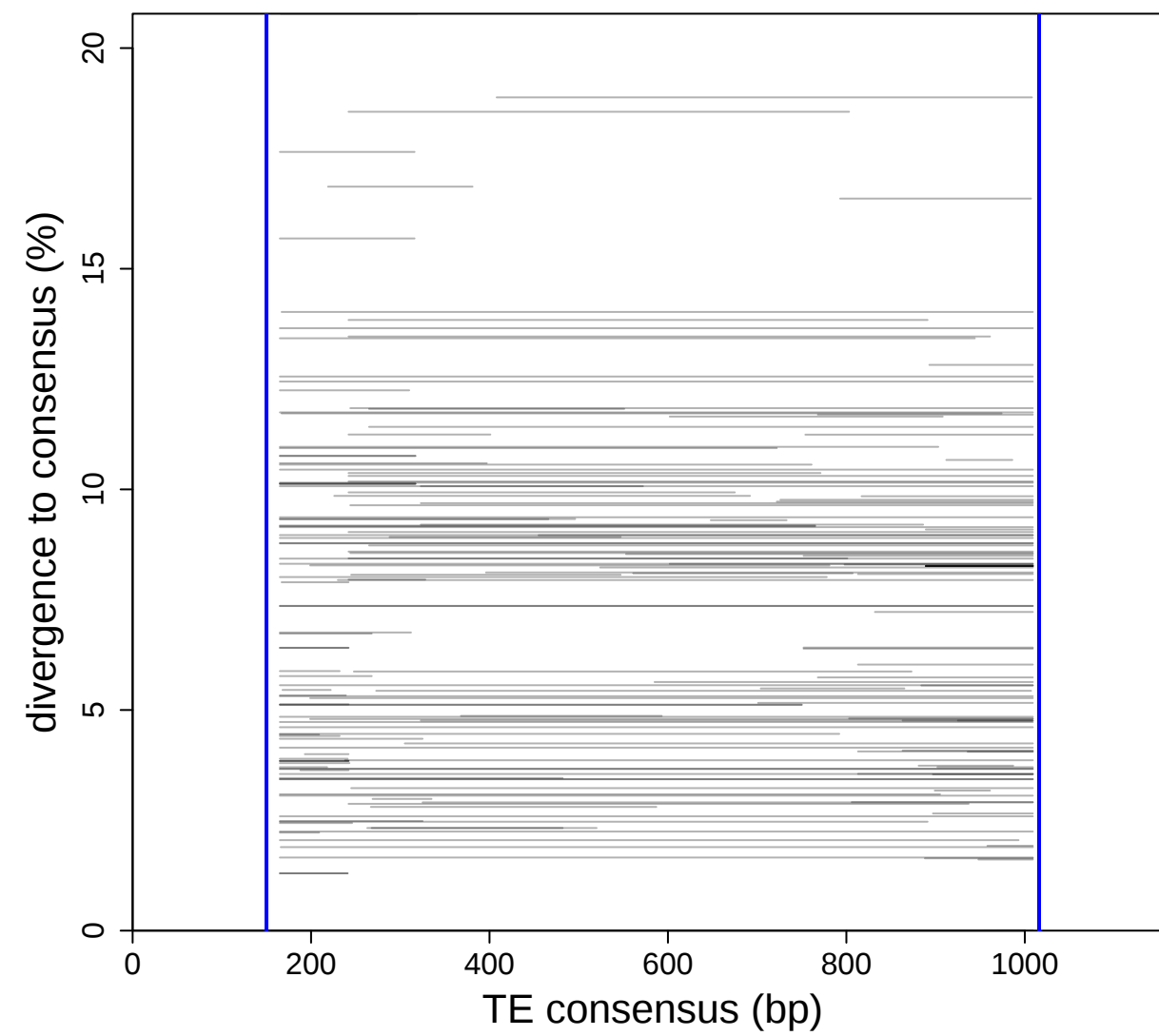
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 857bp; fragments: 191; full length: 39 (≥ 771.3 bp)

After TEtrimmer 857 bp

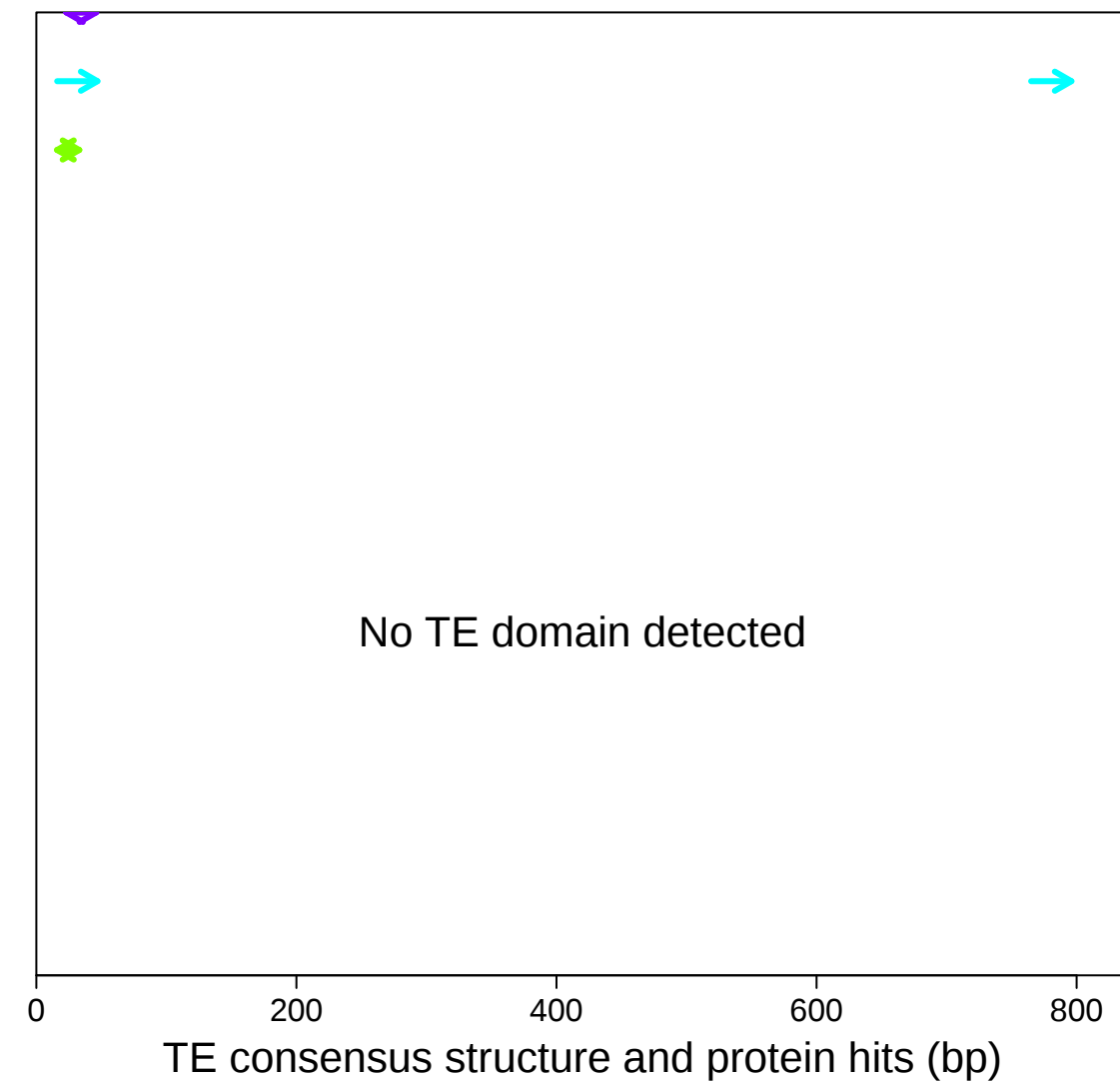
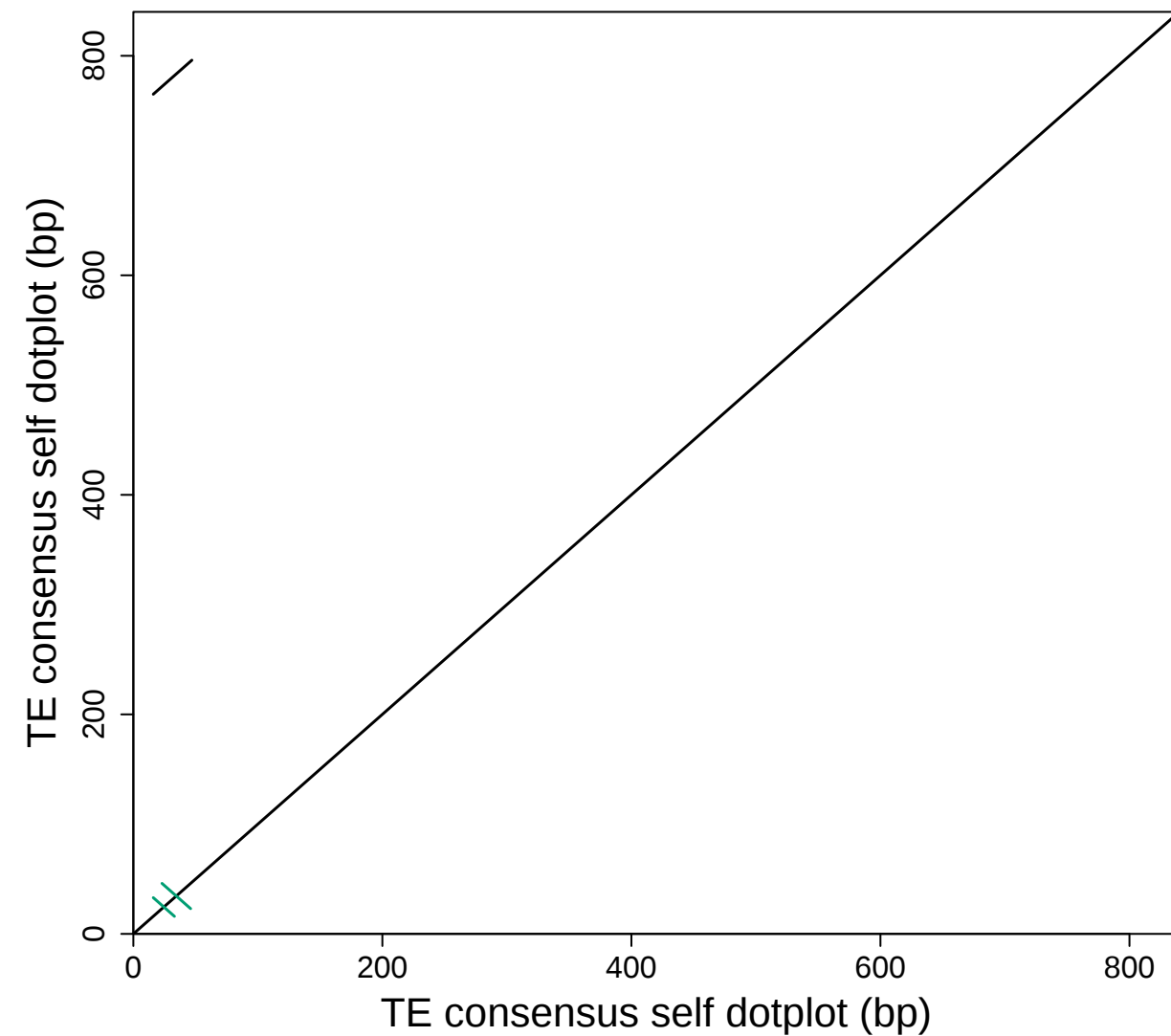
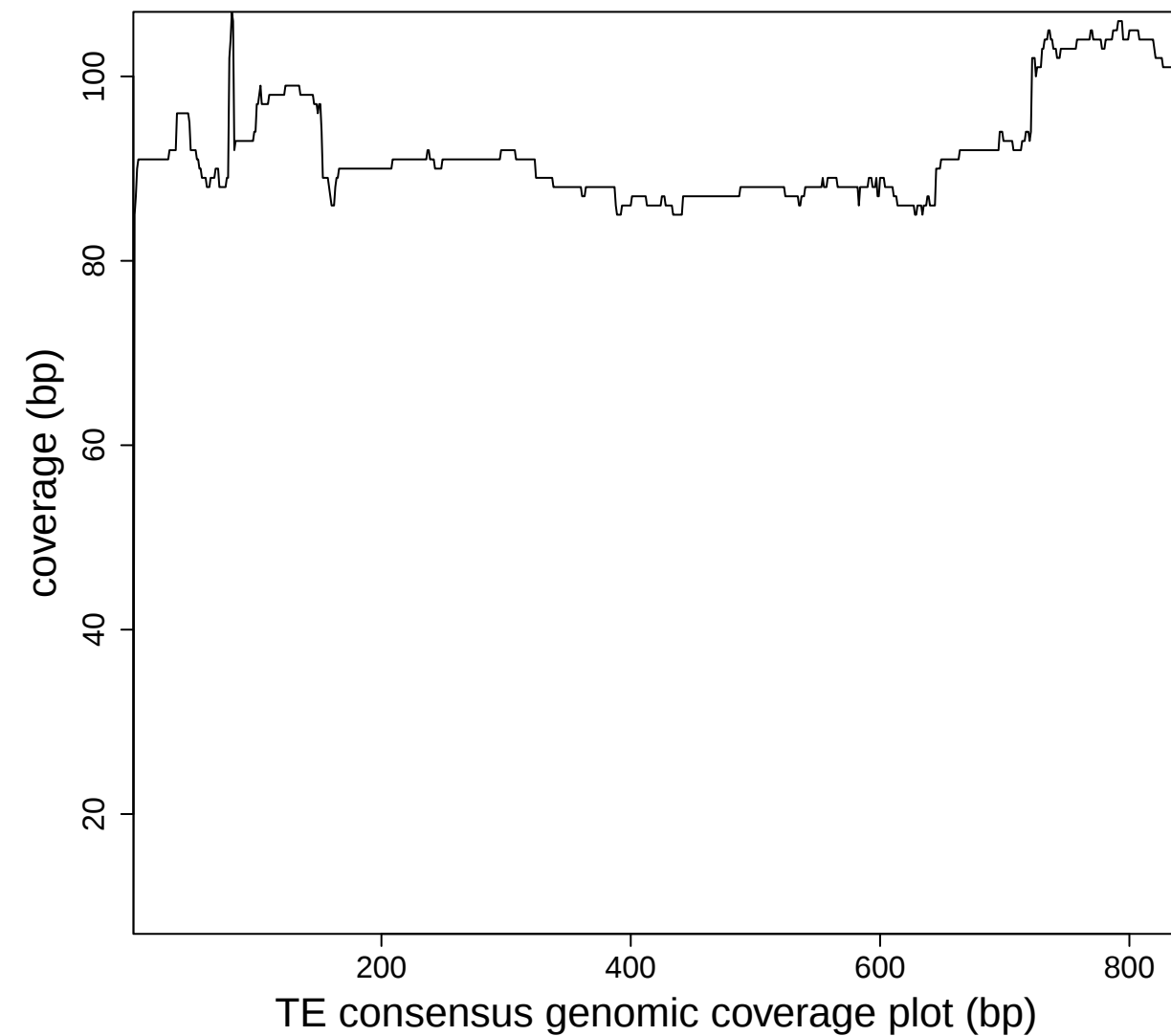
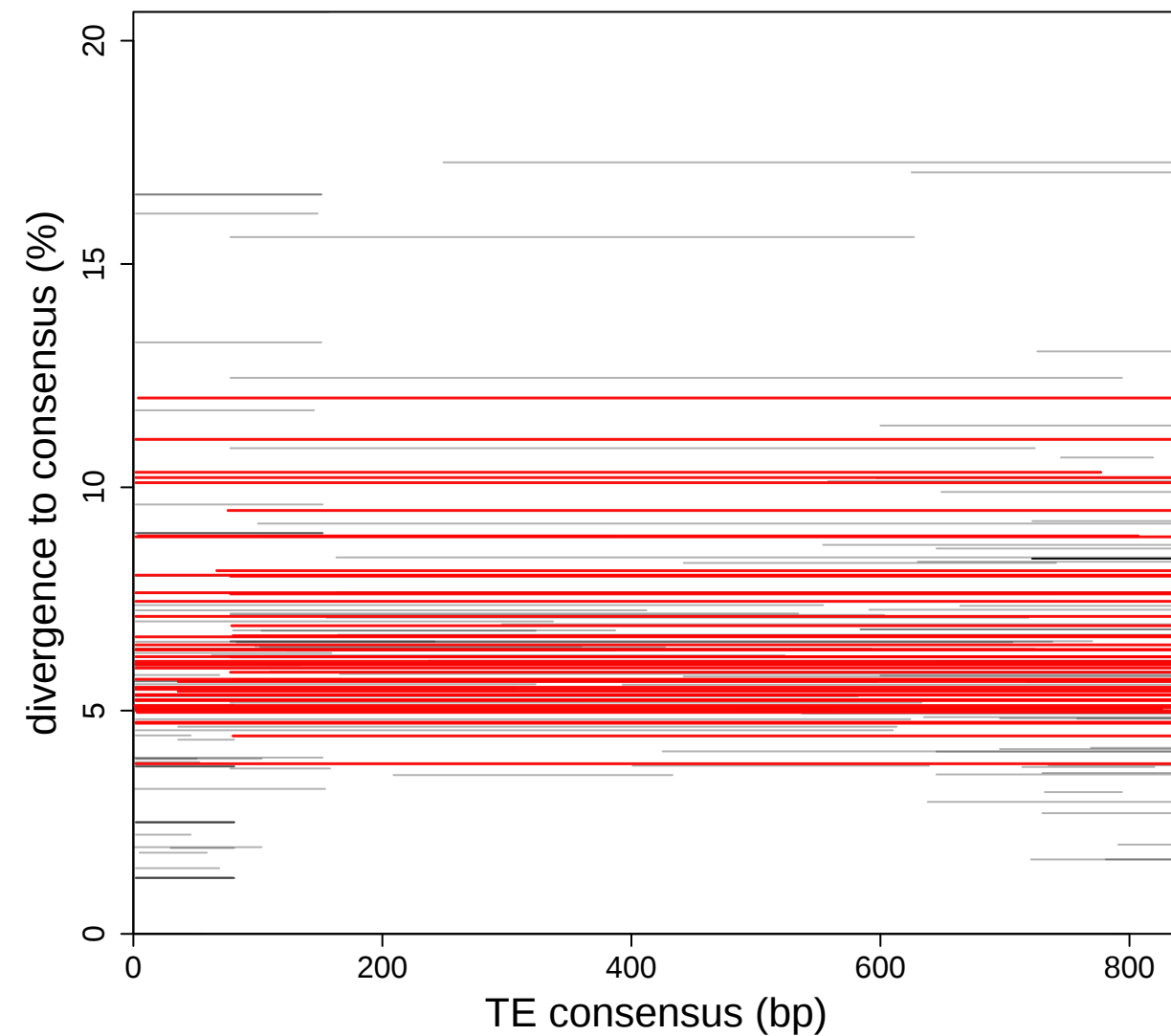


5.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 1166bp; fragments: 185; full length: 0 (>=1049.4bp)

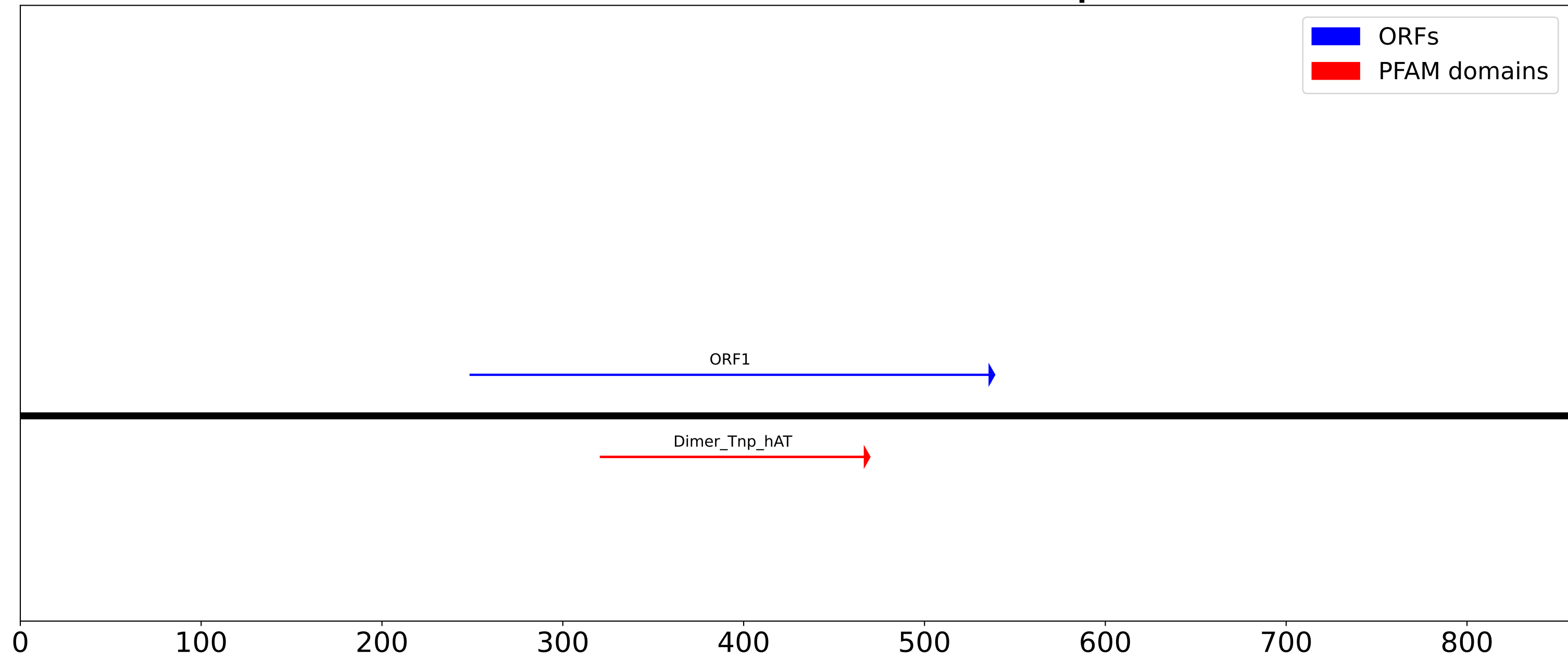
After TEtrimmer Extended plot Blue lines are boundaries



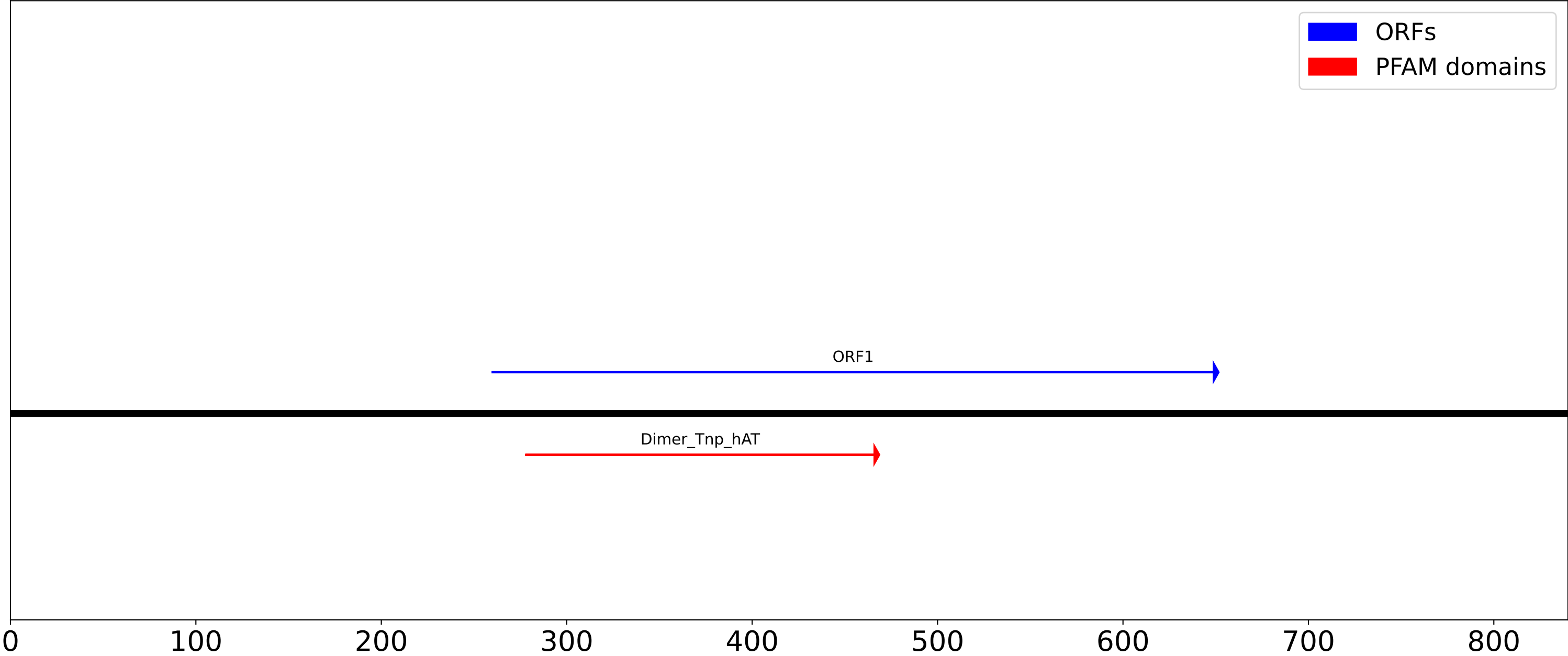
TE: rnd_5_family_6425
size: 840bp; fragments: 188; full length: 48 (≥ 756 bp)



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

