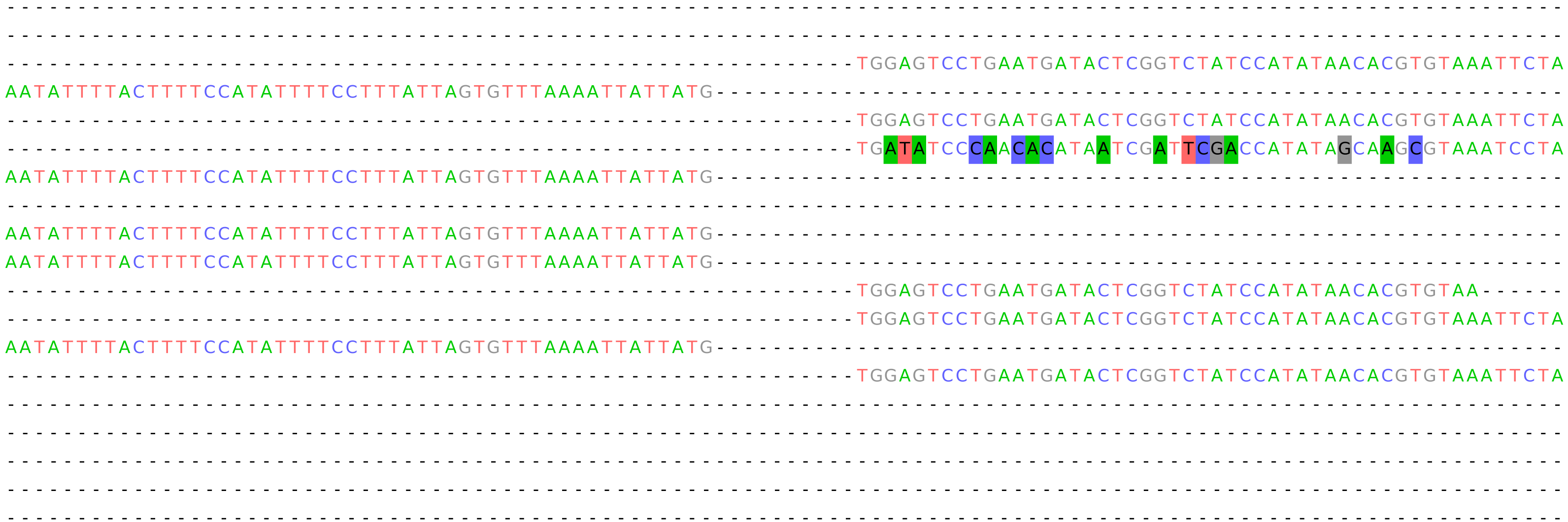
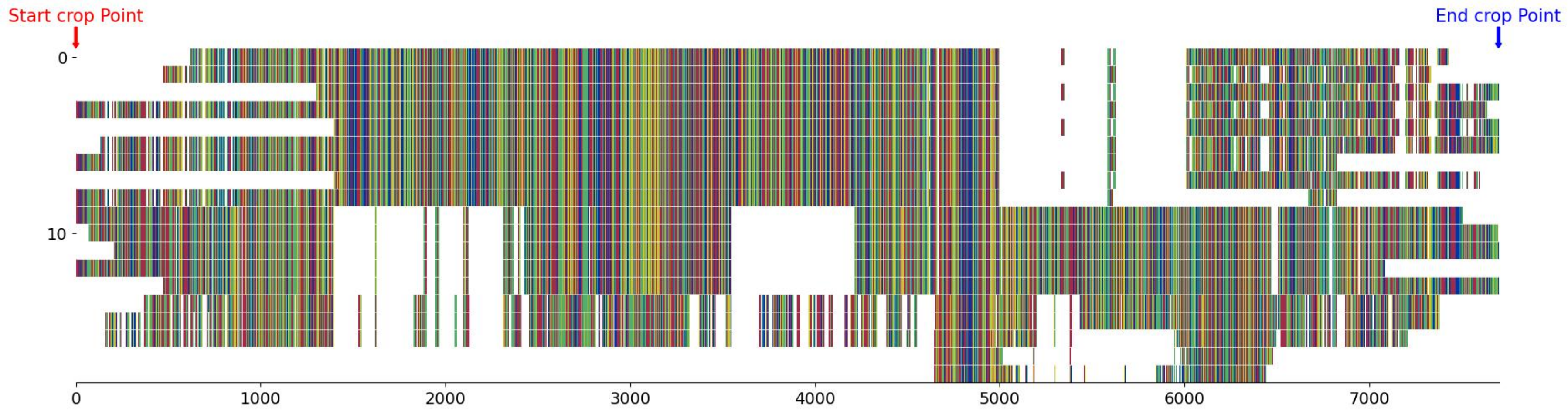


Start crop Point

End crop Point

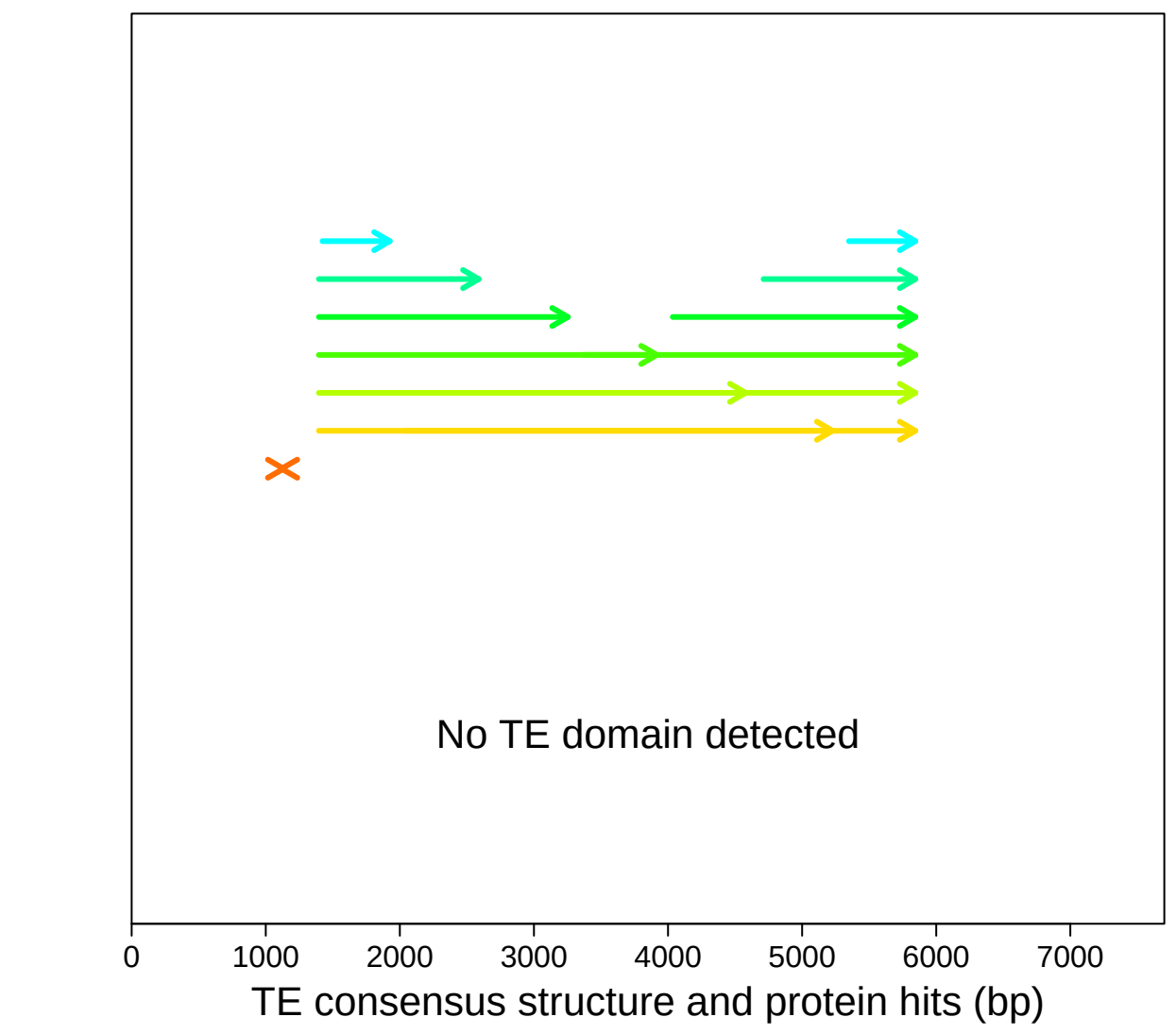
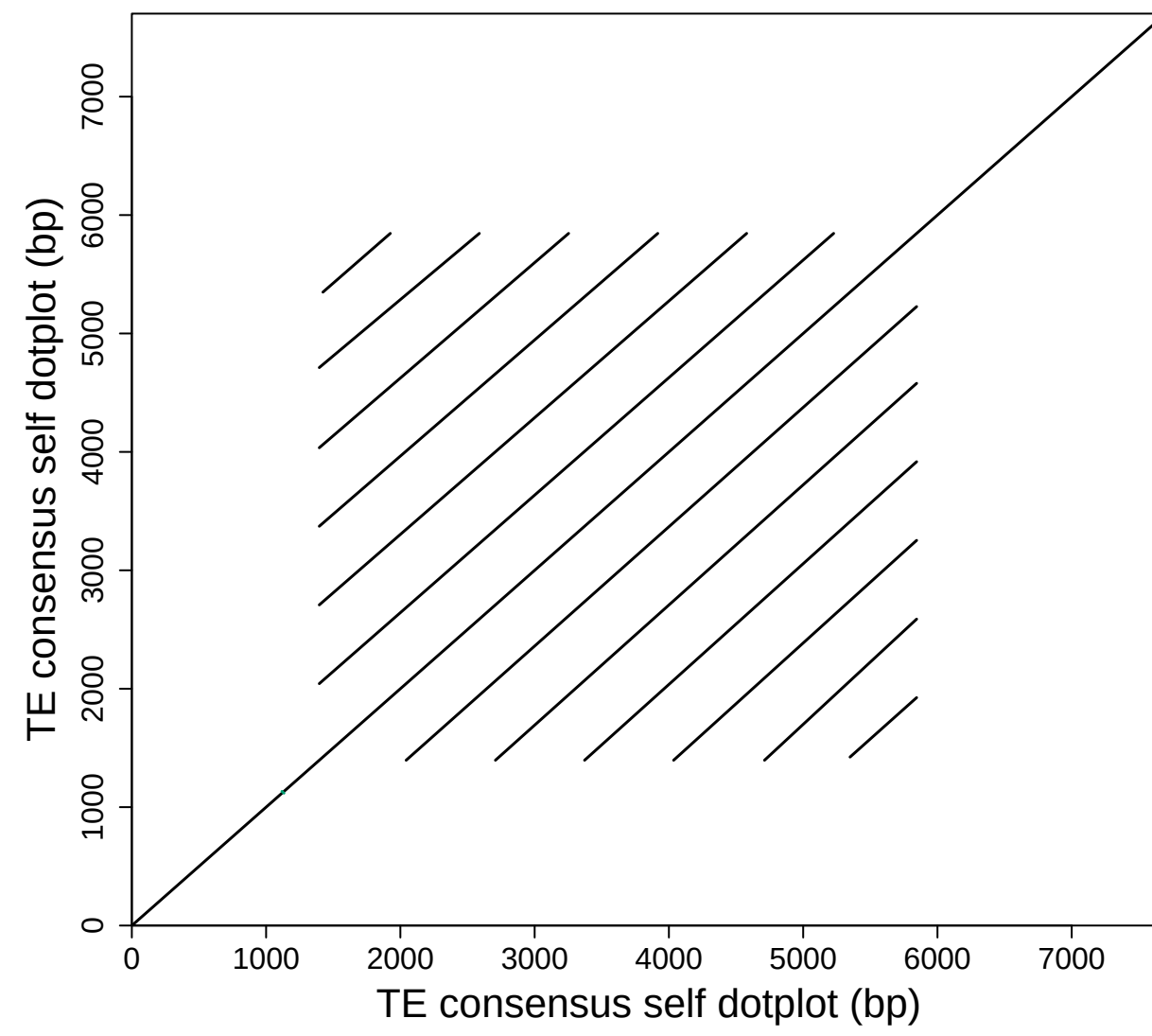
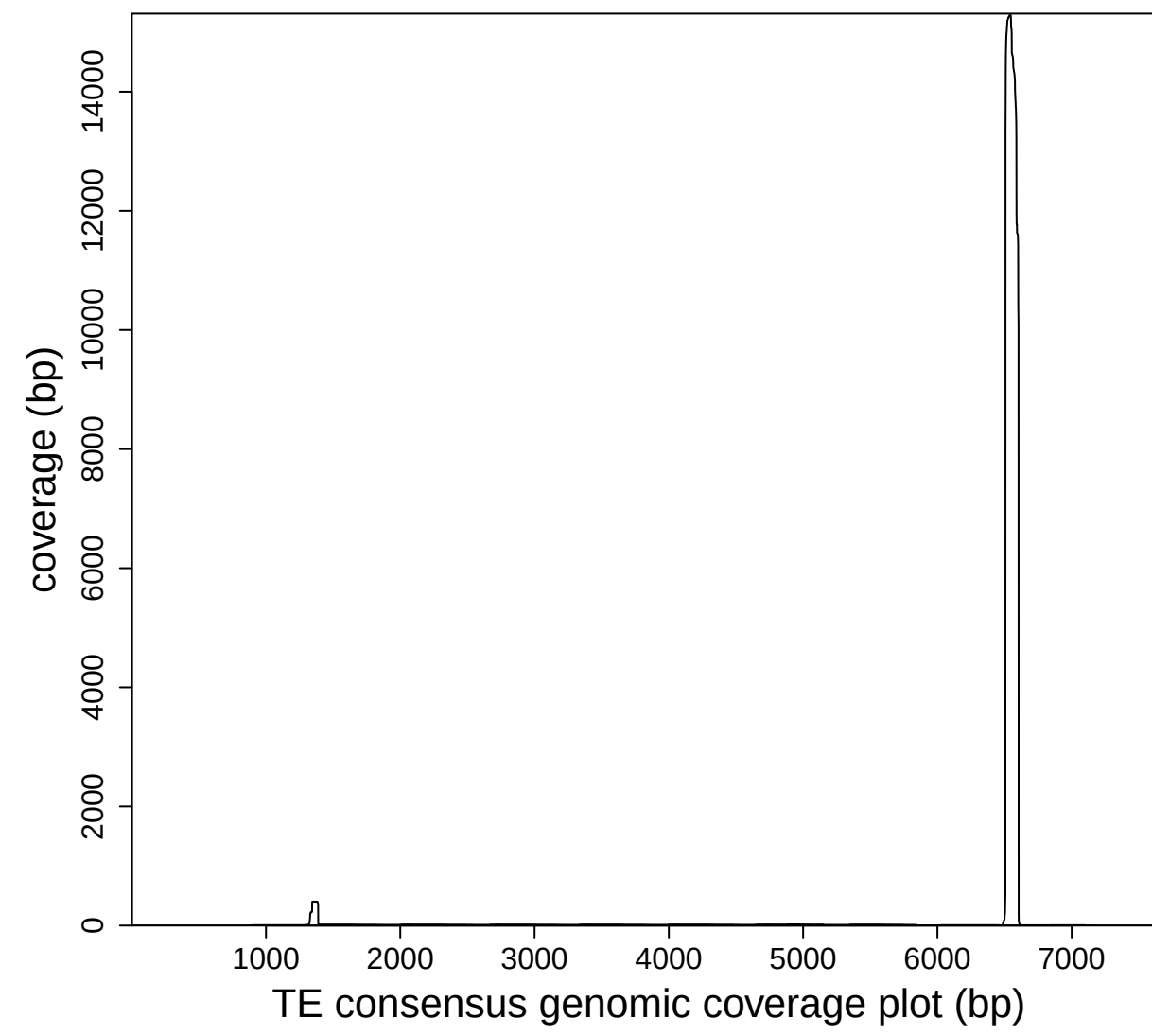
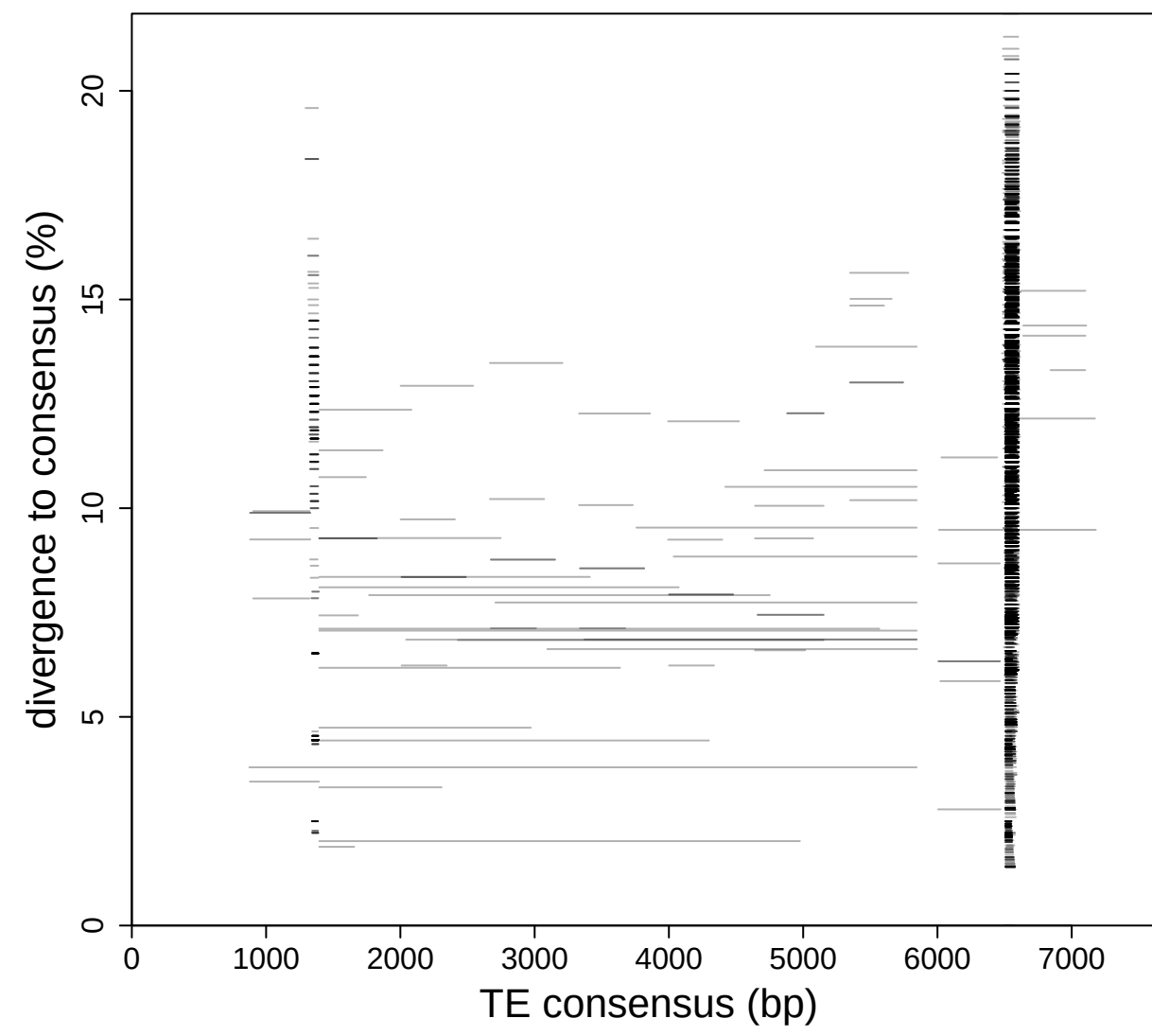
MSA length = 7701





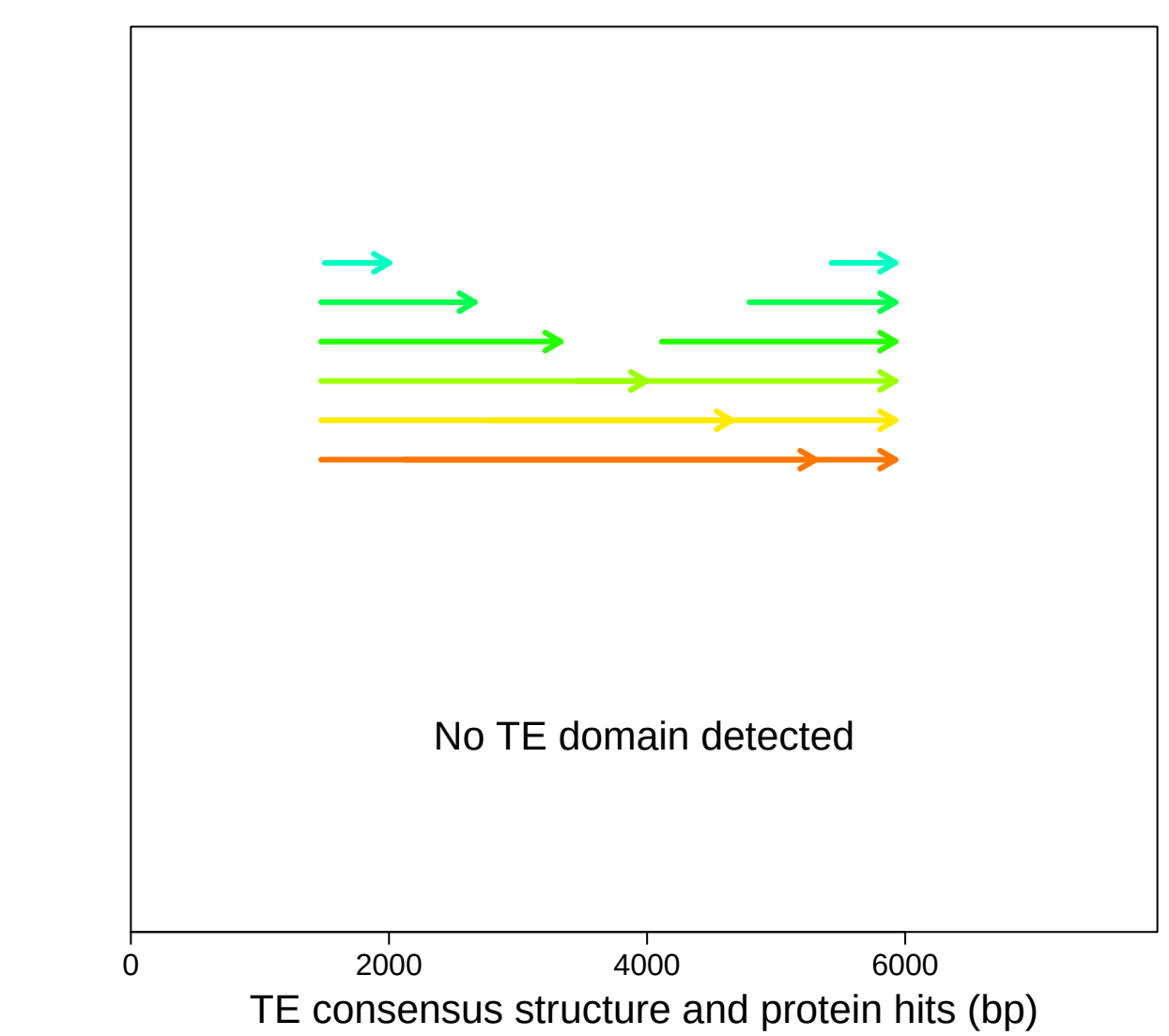
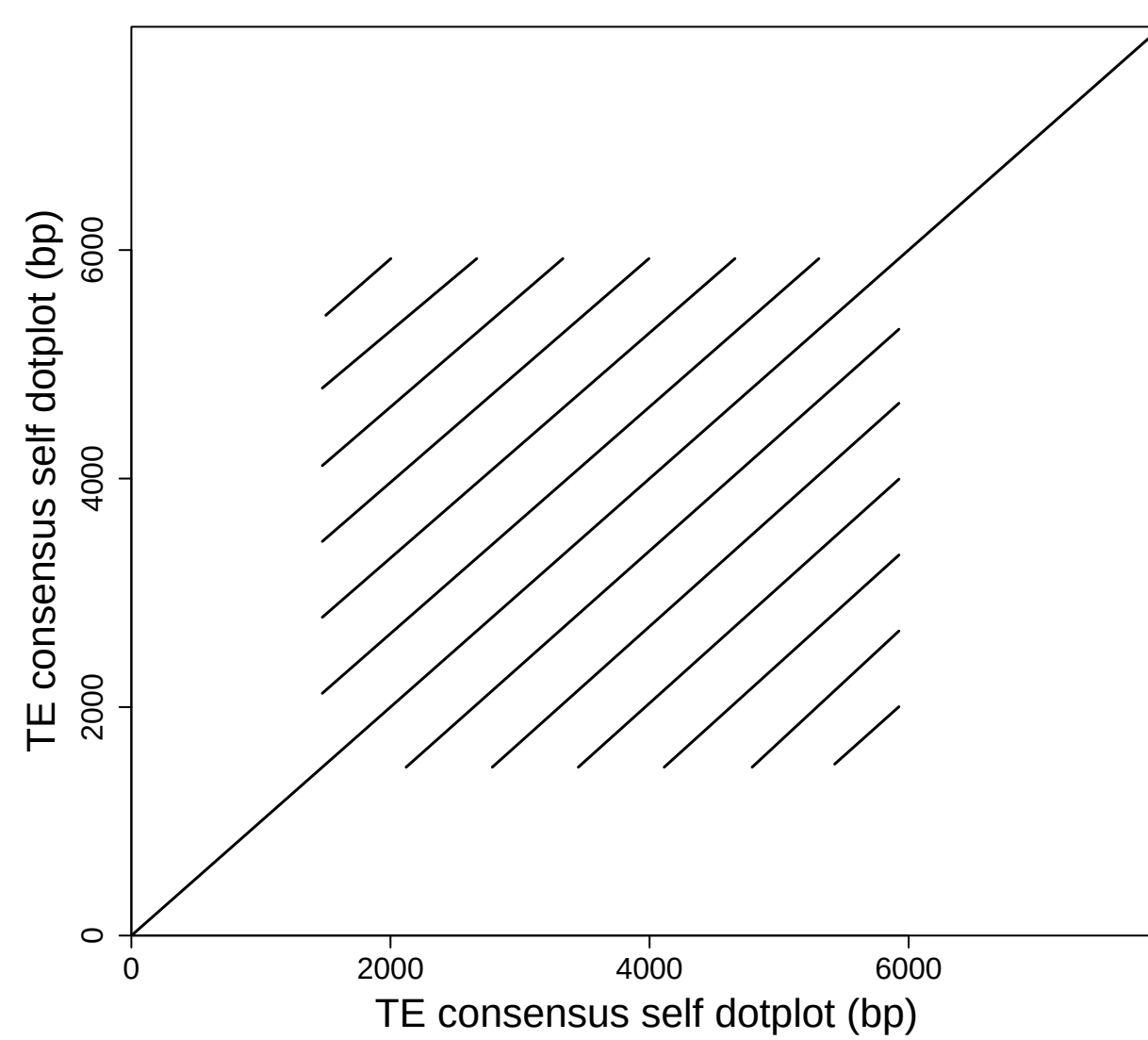
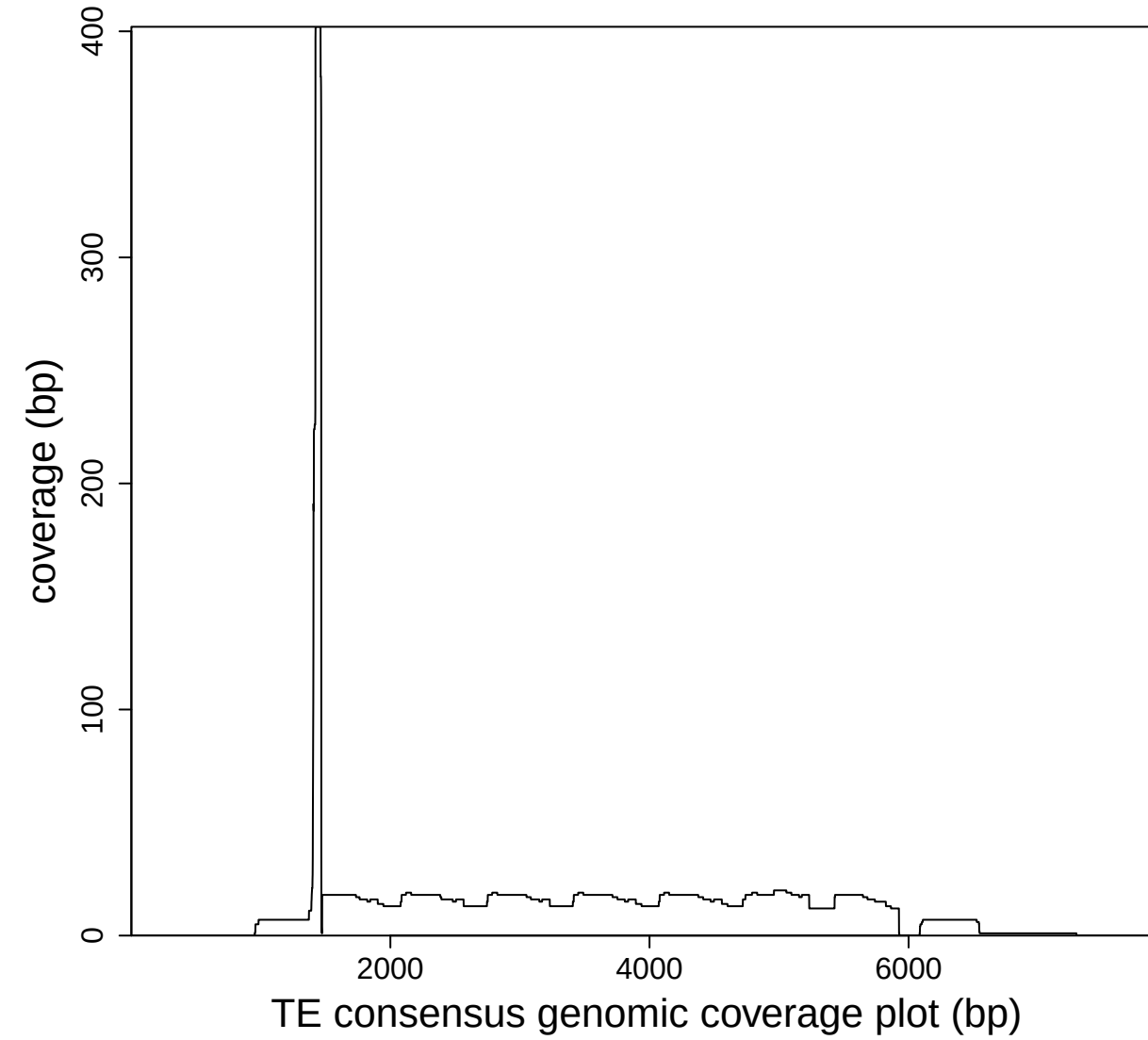
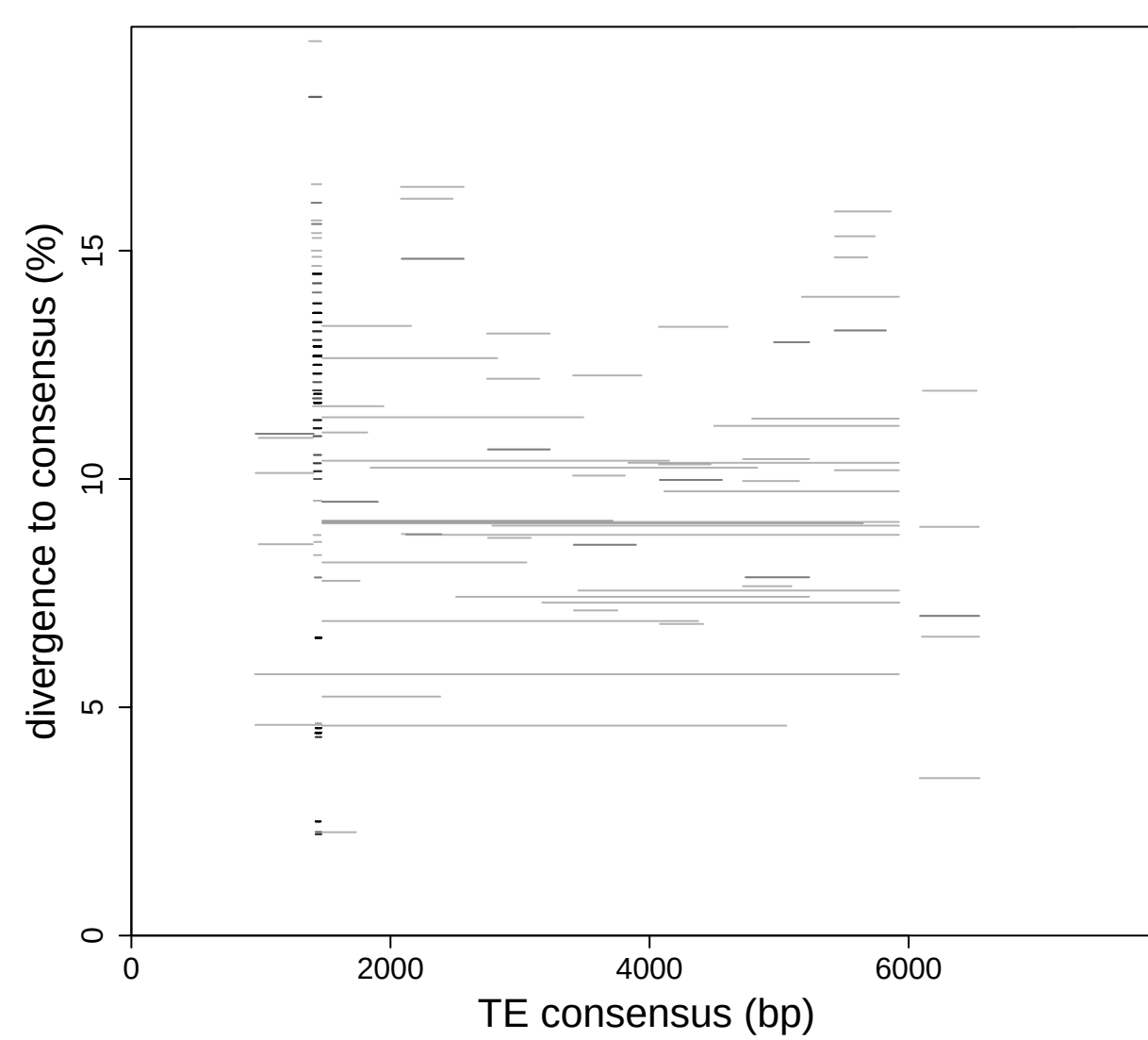
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 7701bp; fragments: 15803; full length: 0 (>=6930.9bp)

After TEtrimmer 7701 bp

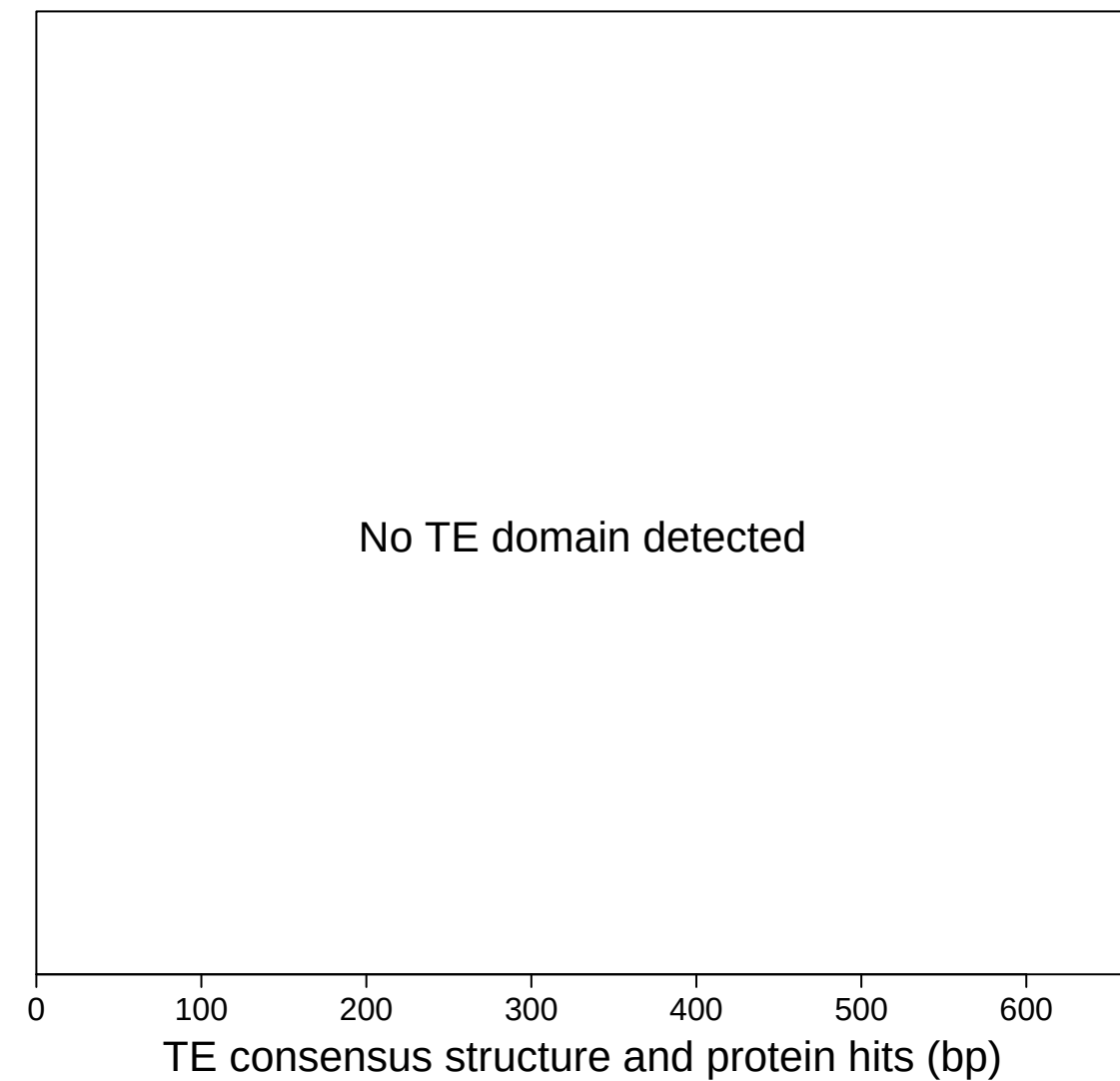
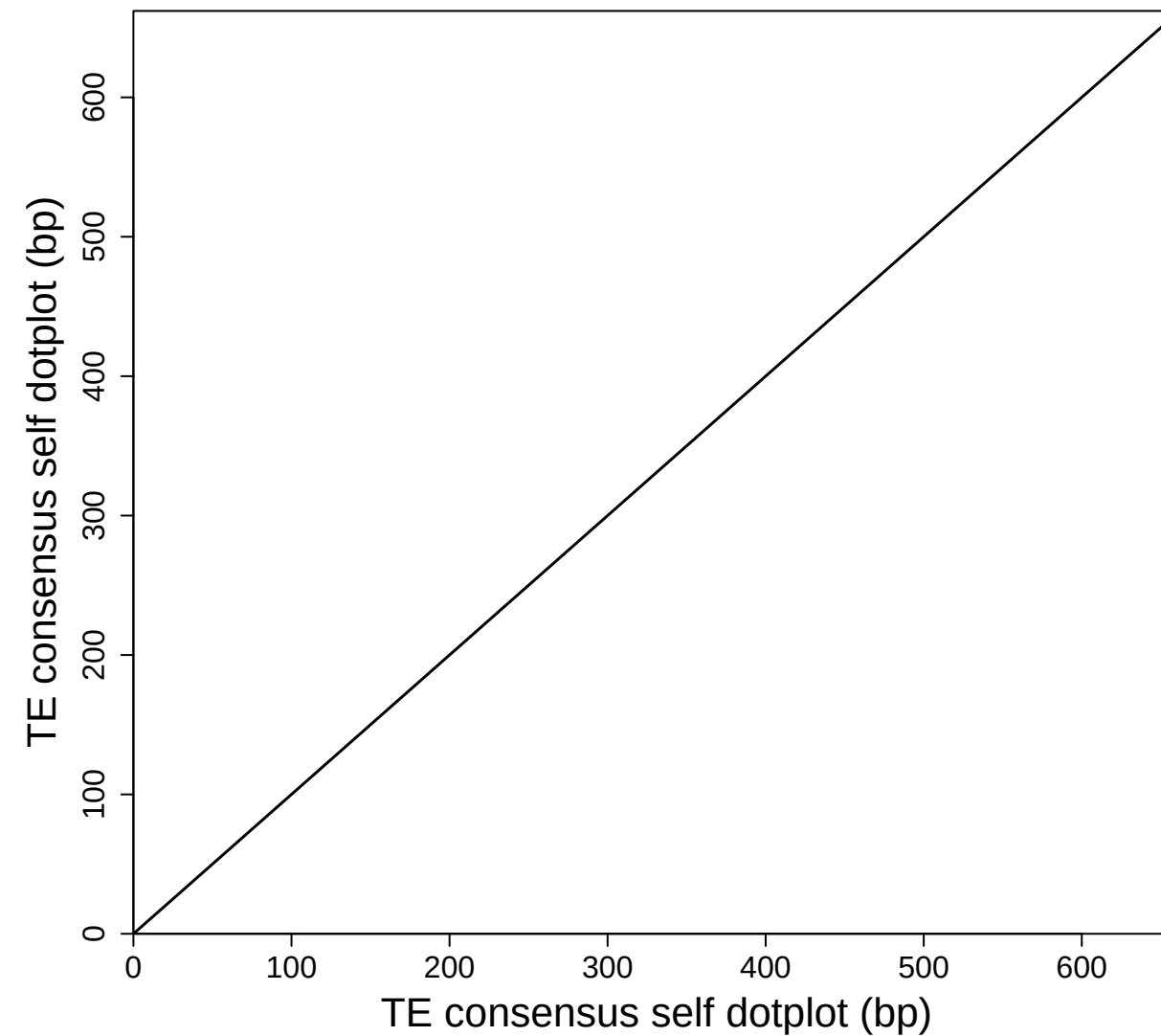
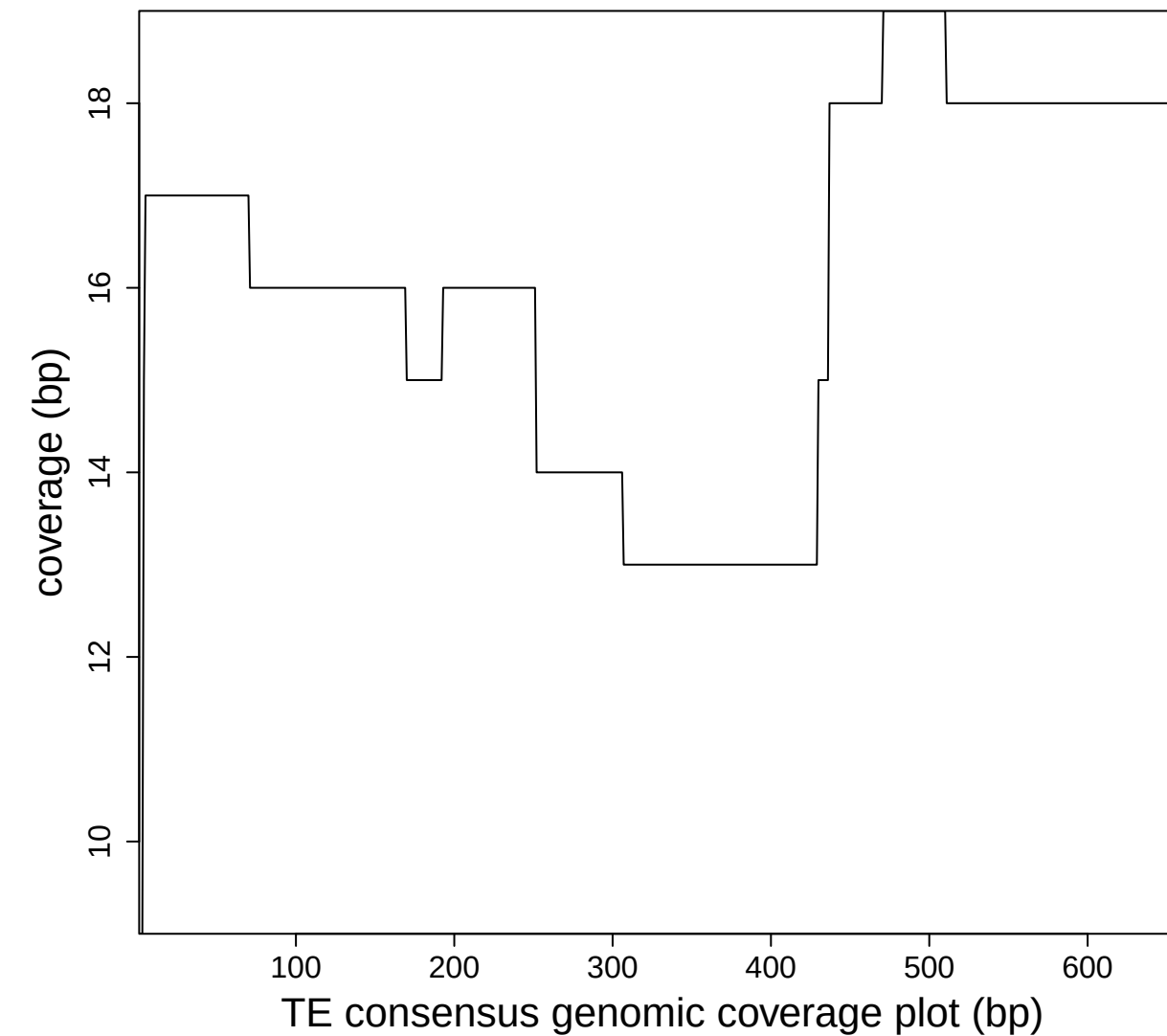
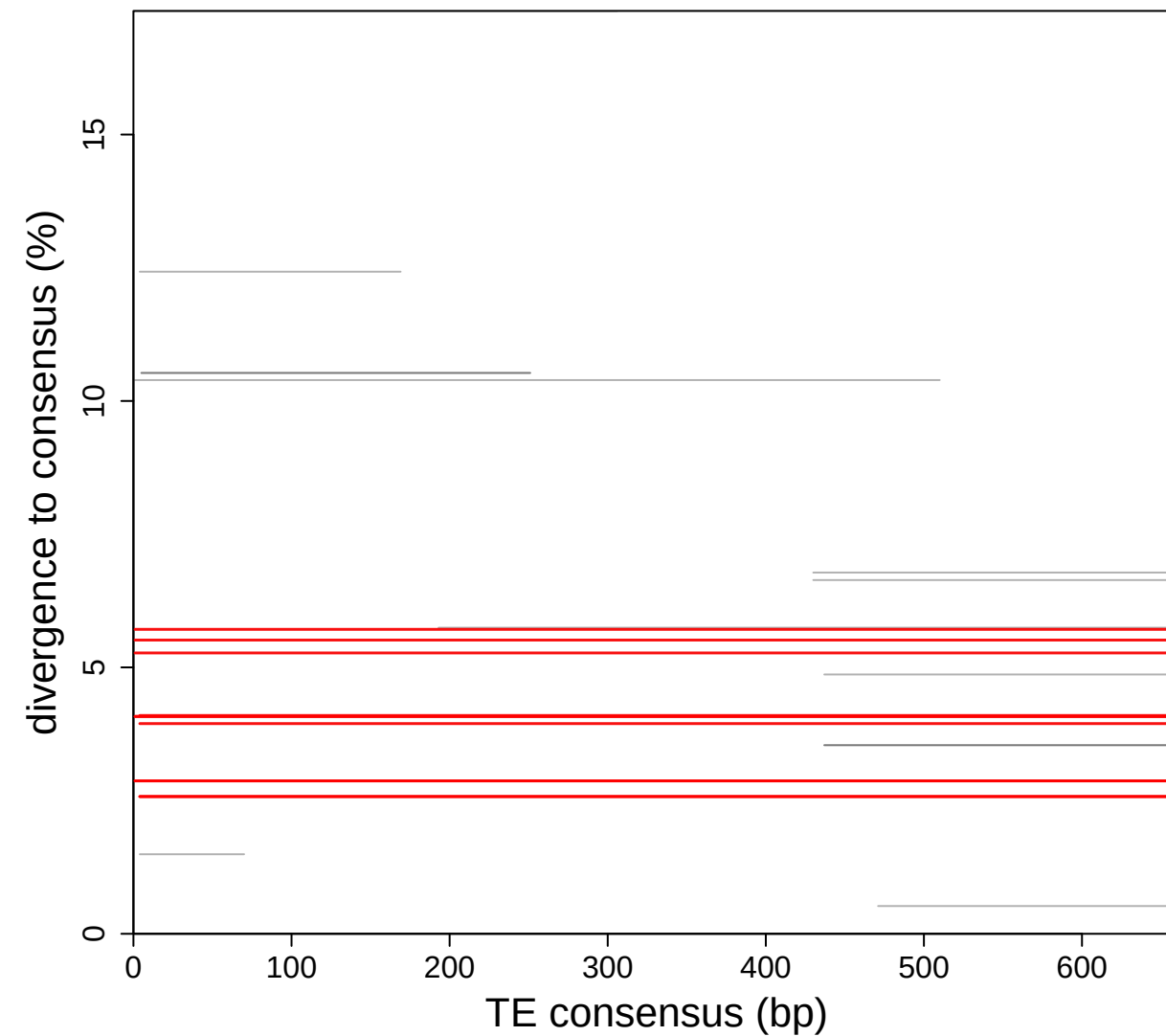


1.fasta.b.bed uf.bed g_1.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 7954bp; fragments: 475; full length: 0 (>=7158.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

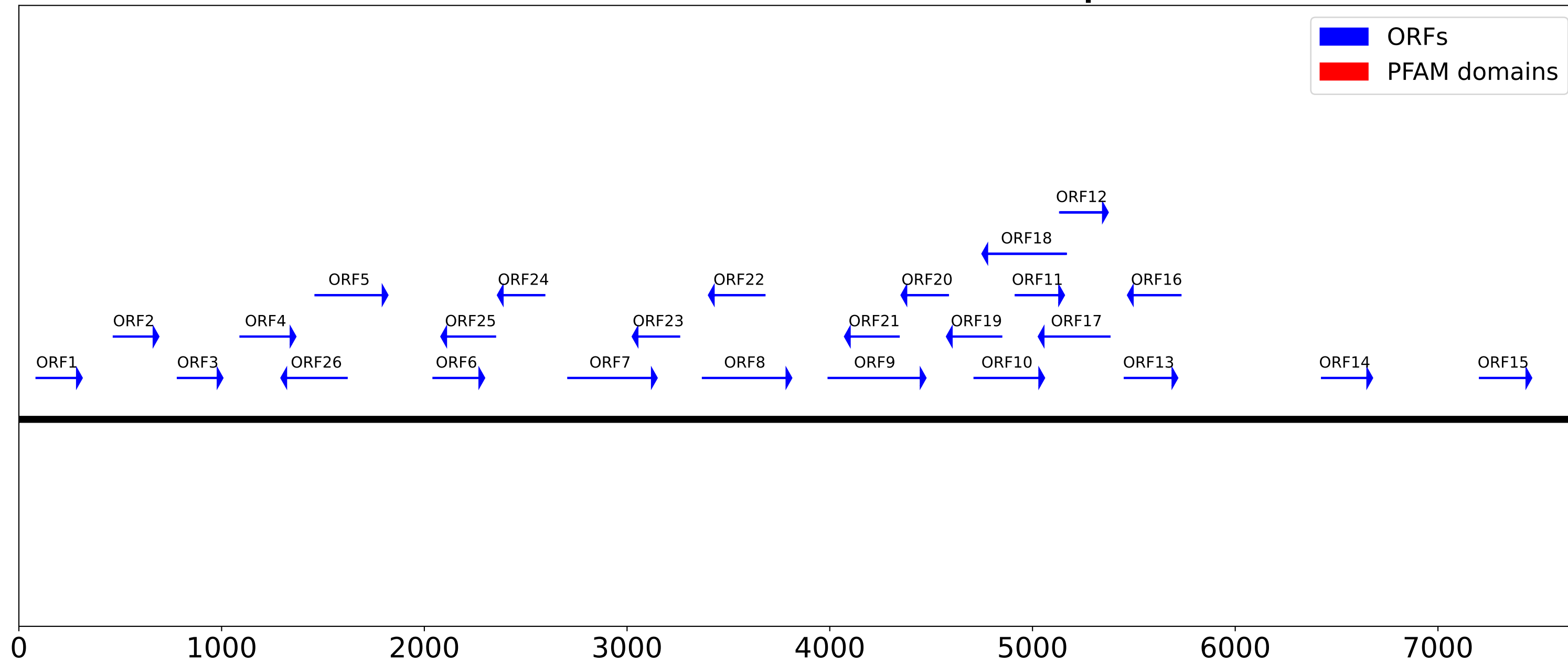


TE: rnd_5_family_8571
size: 662bp; fragments: 24; full length: 11 (≥ 595.8 bp)

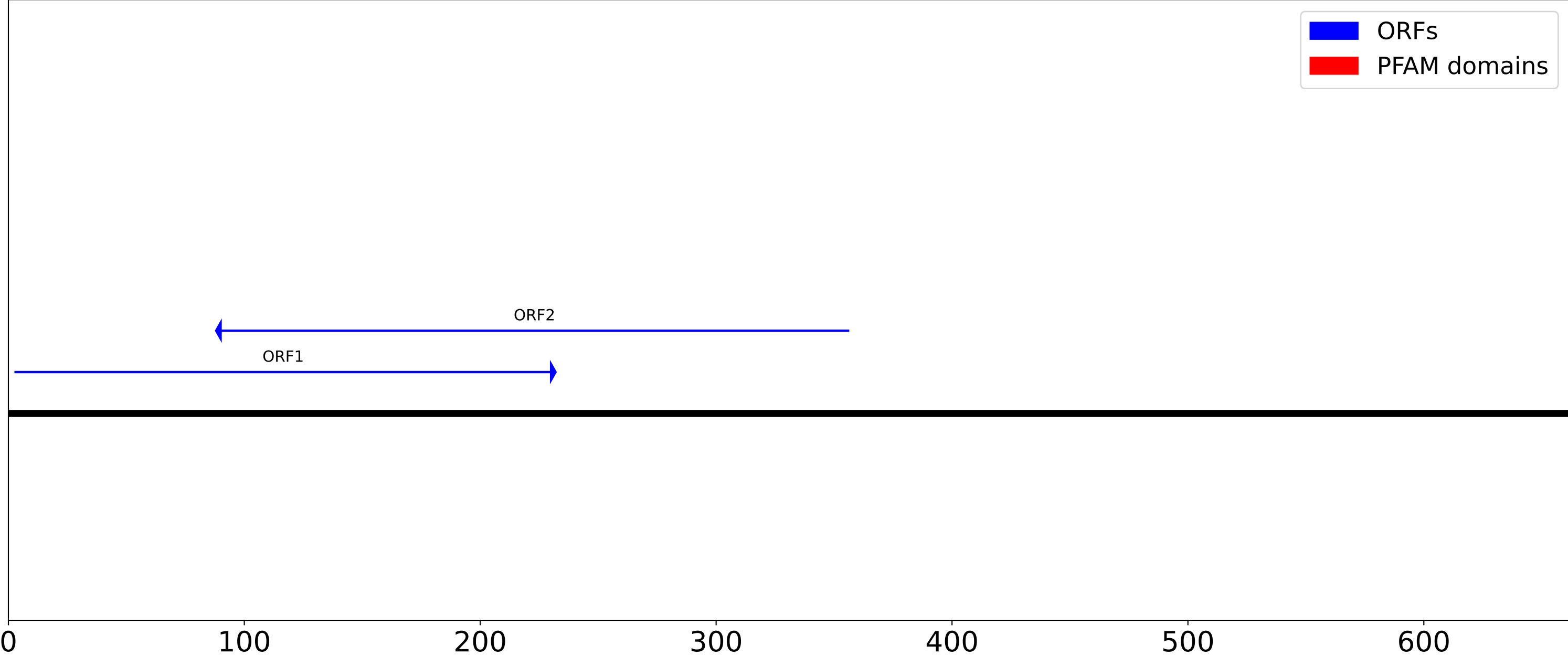


Before TEtrimmer 662 bp

After TEtrimmer ORF and PFAM domain plot



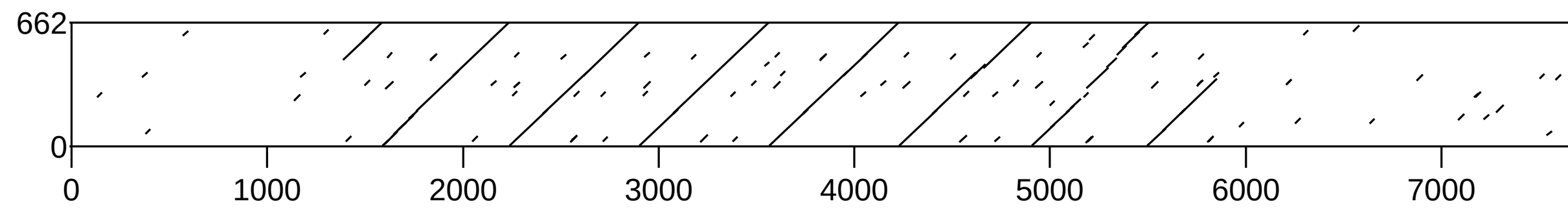
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

ensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)