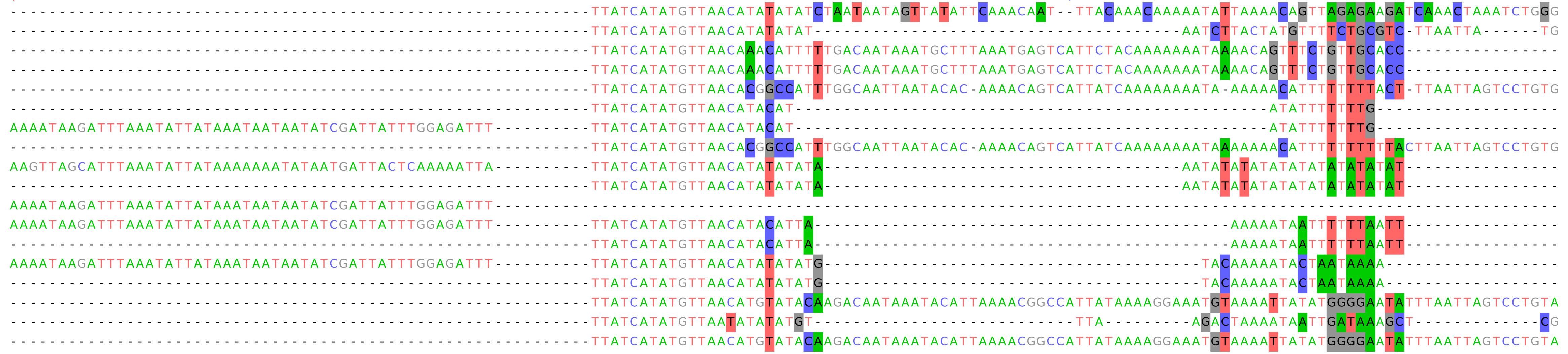
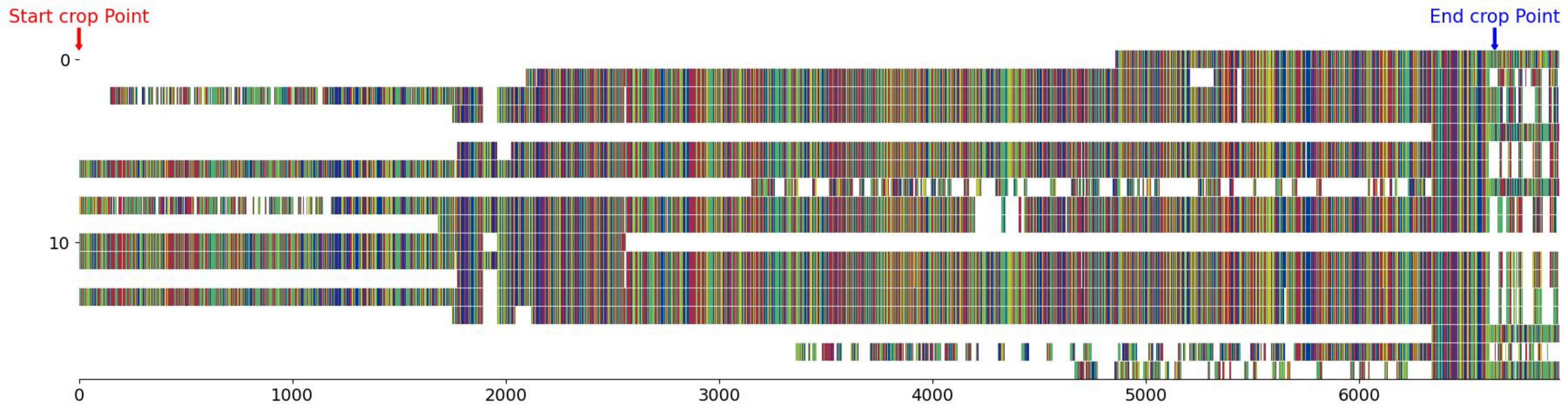


Start crop Point

End crop Point

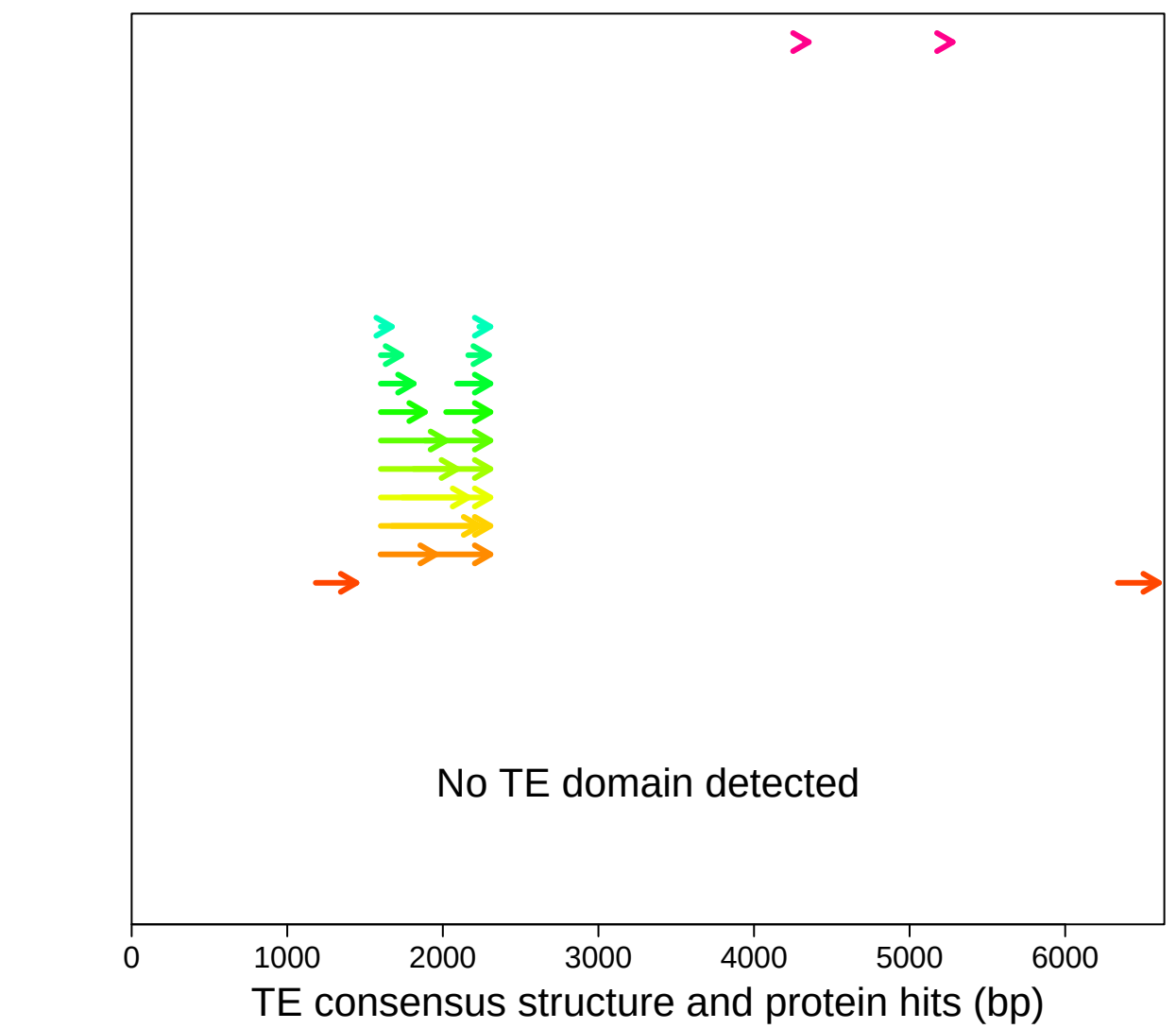
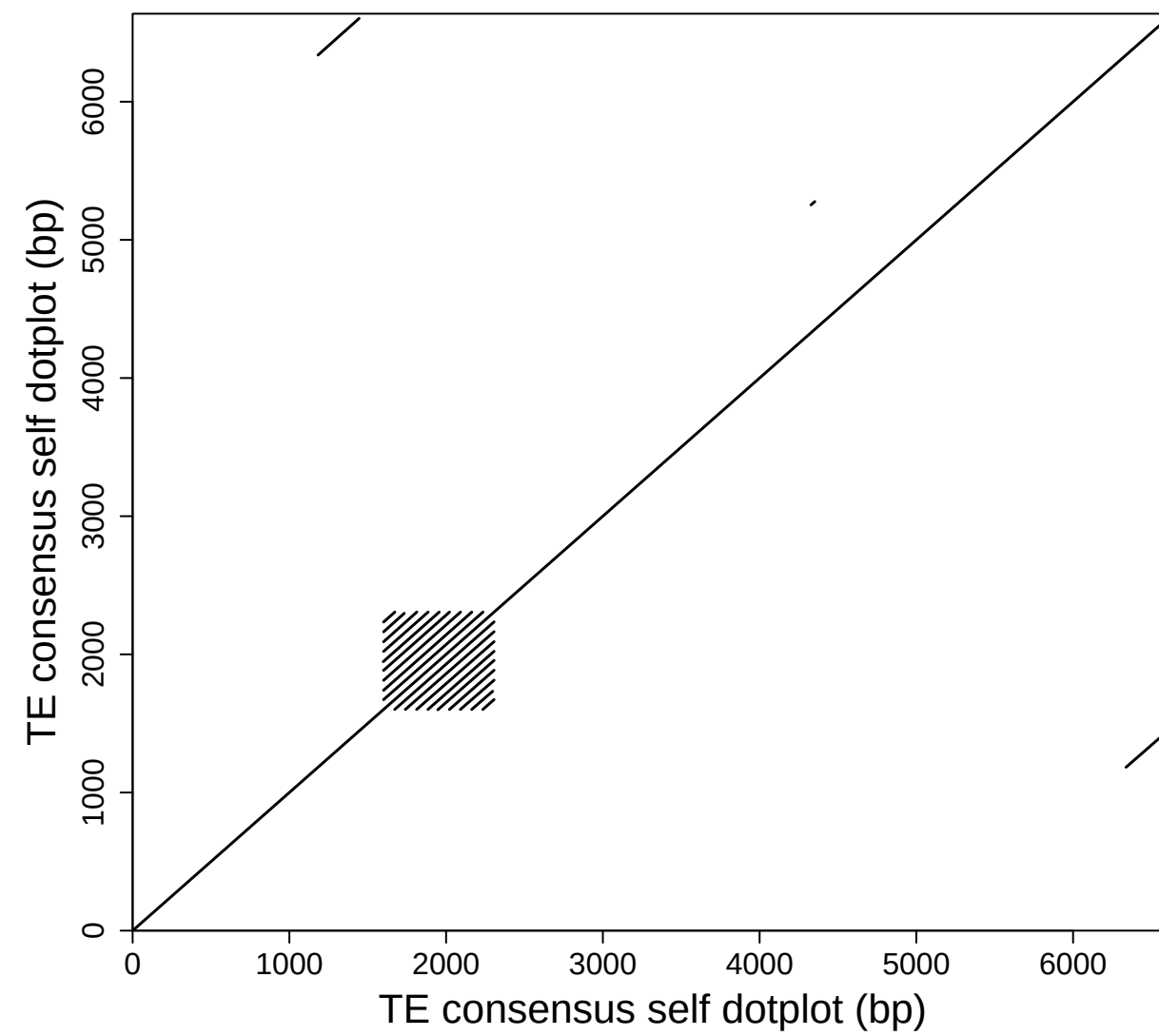
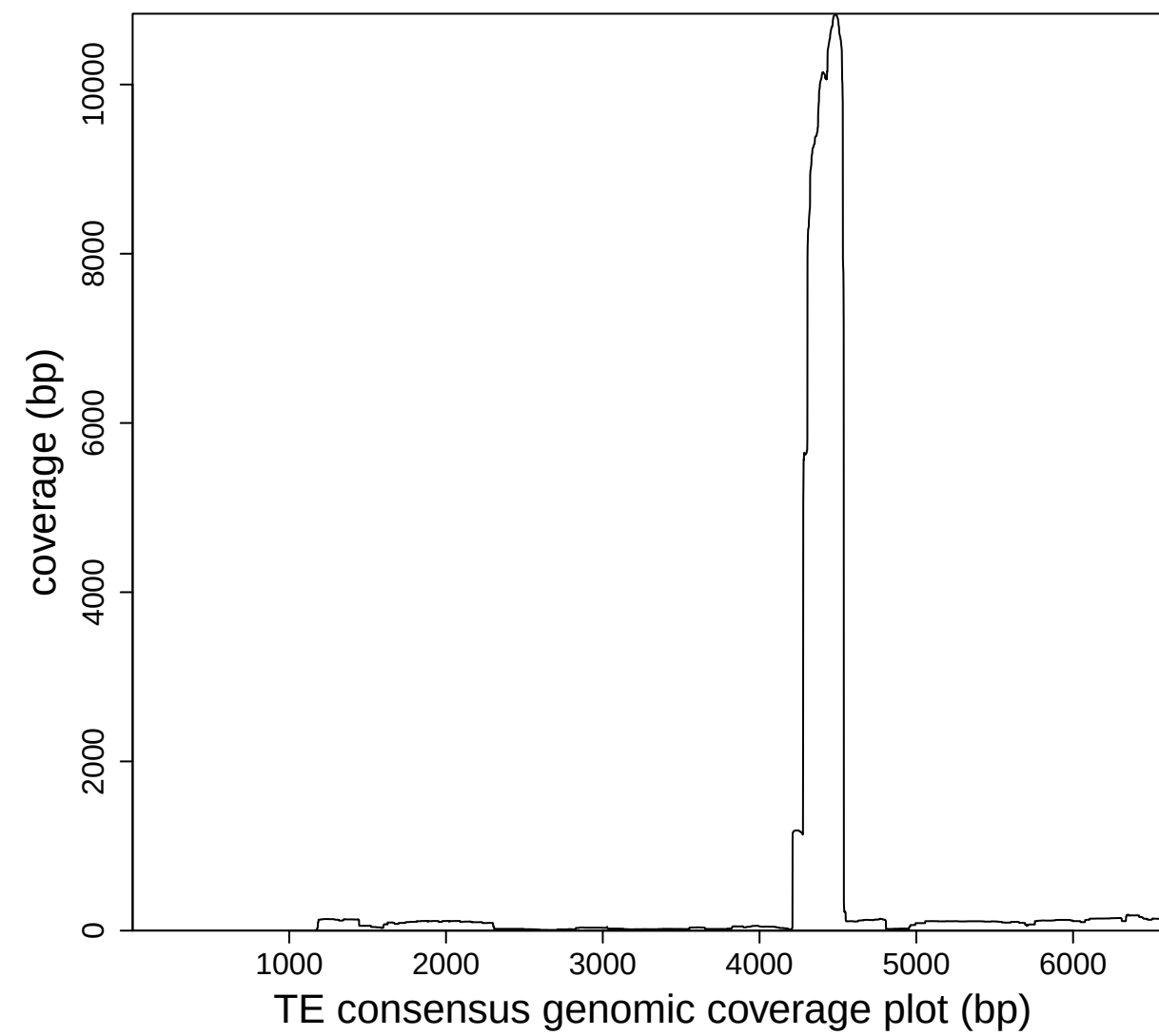
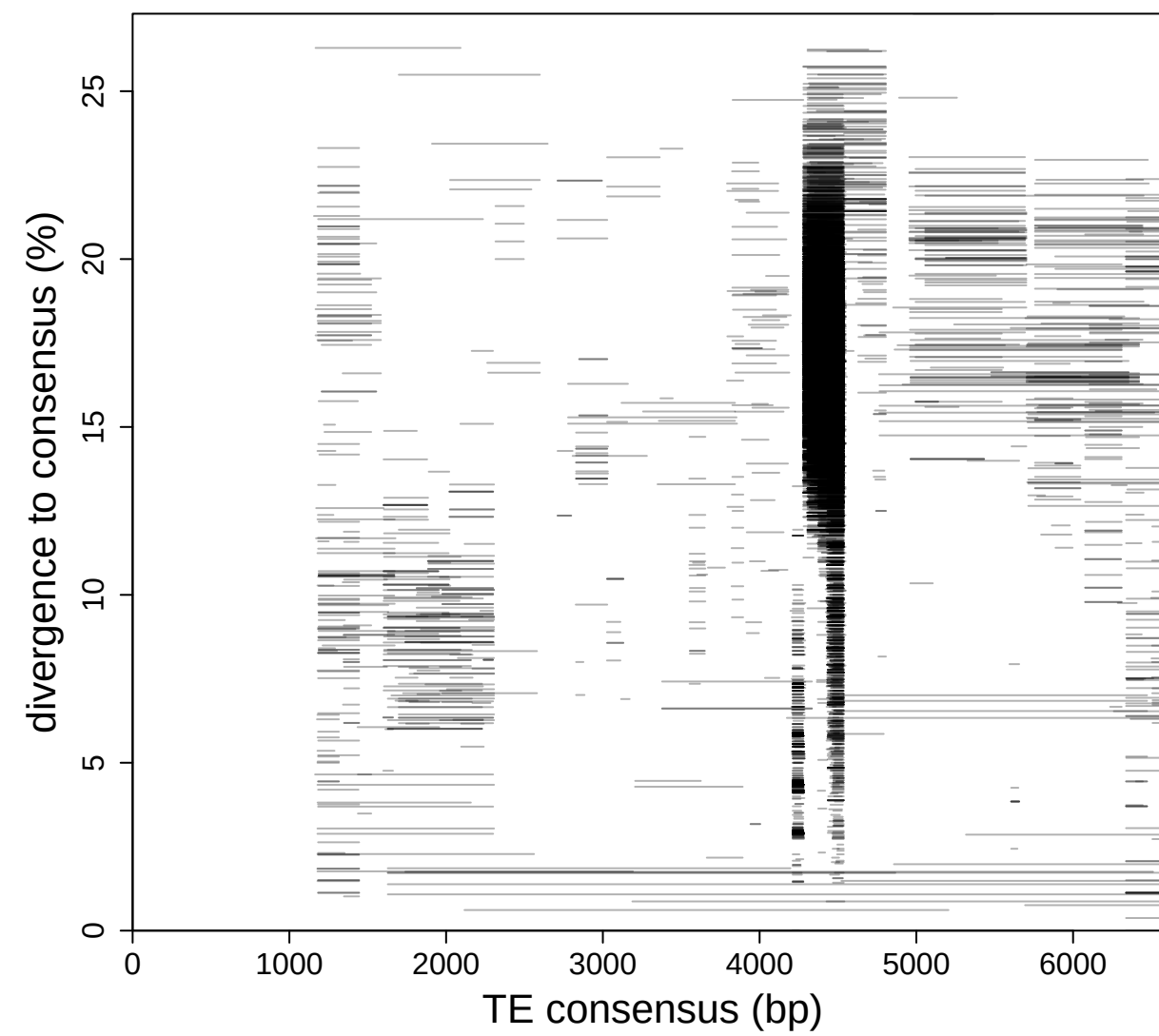
MSA length = 6637





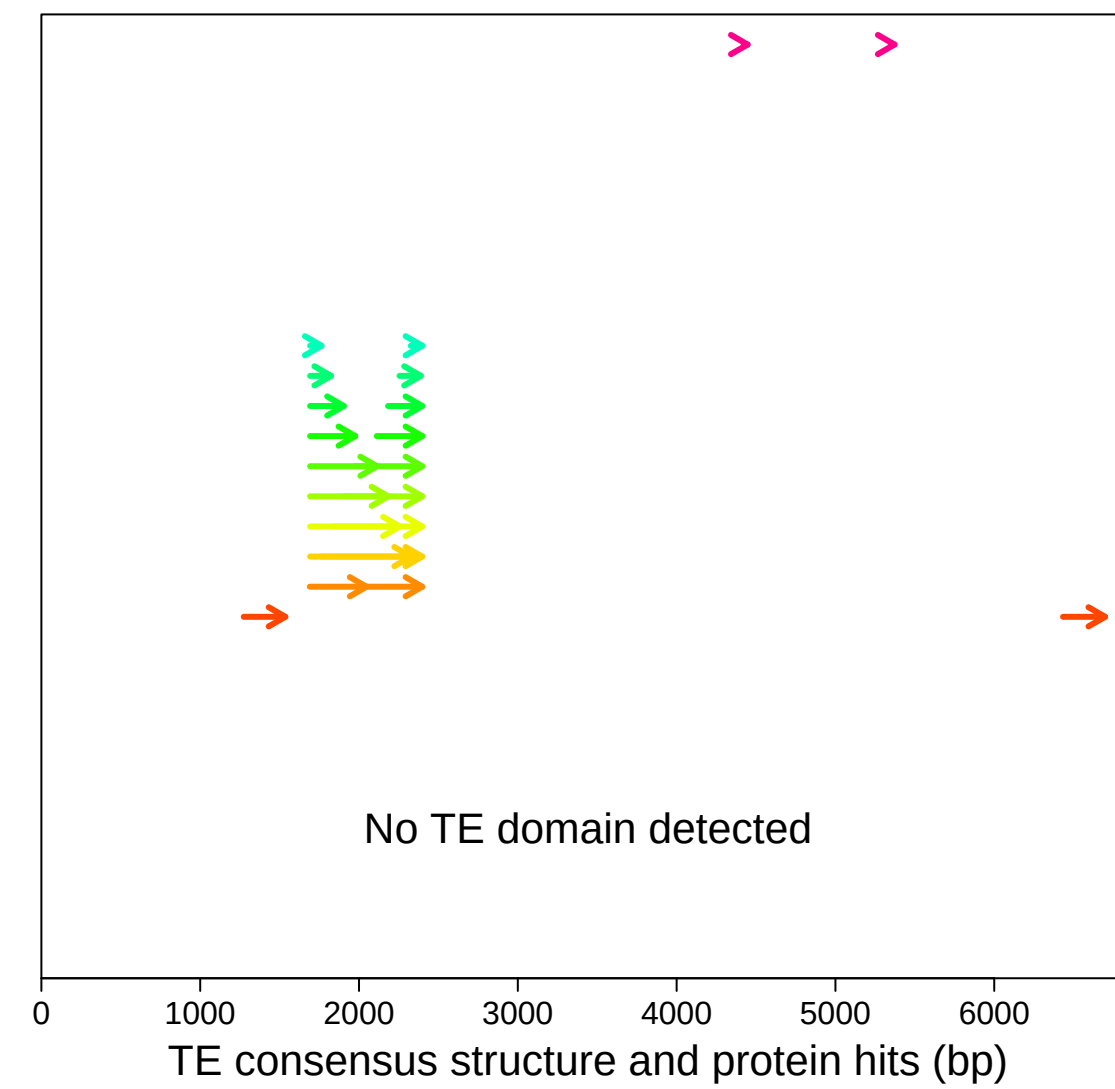
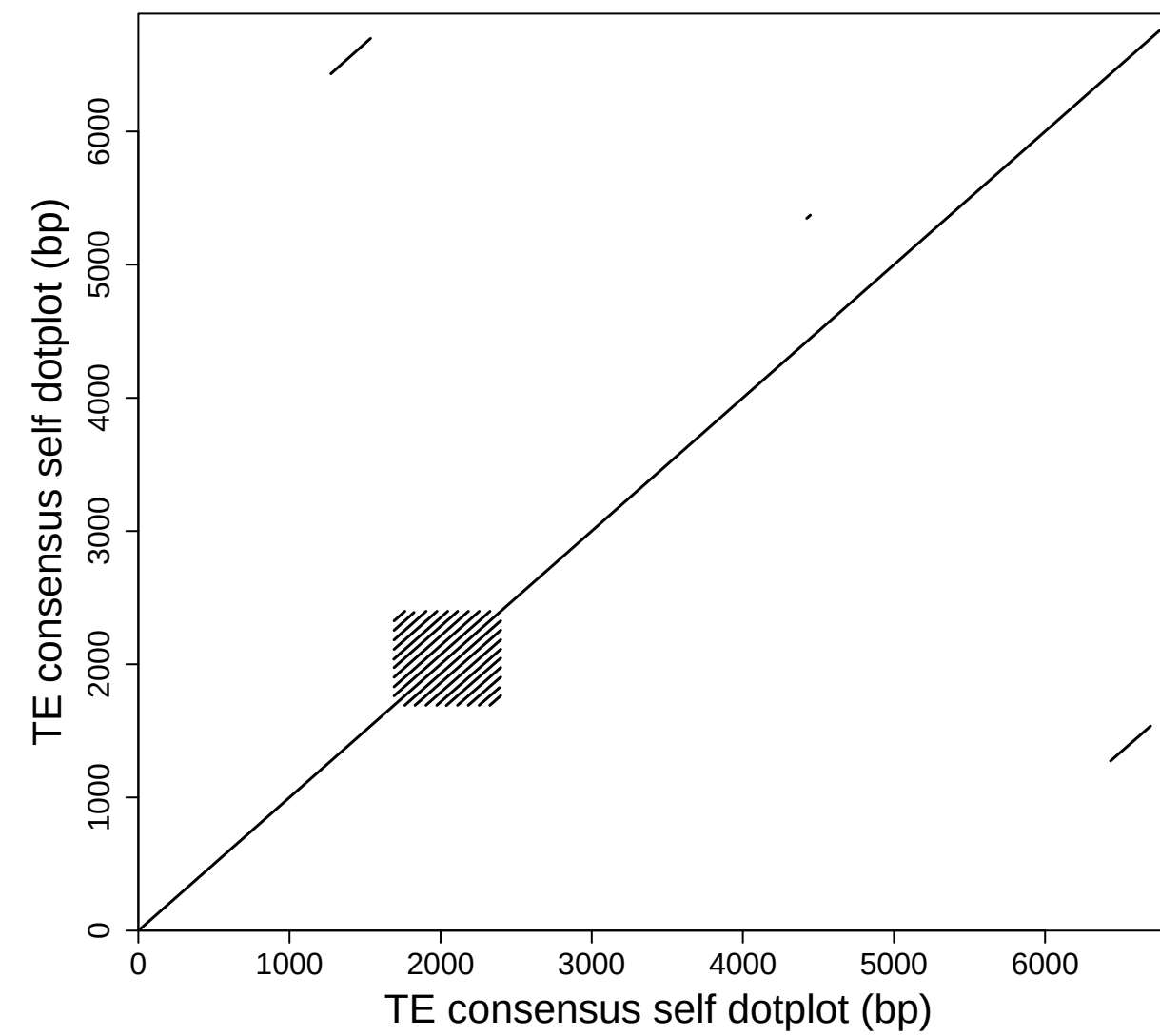
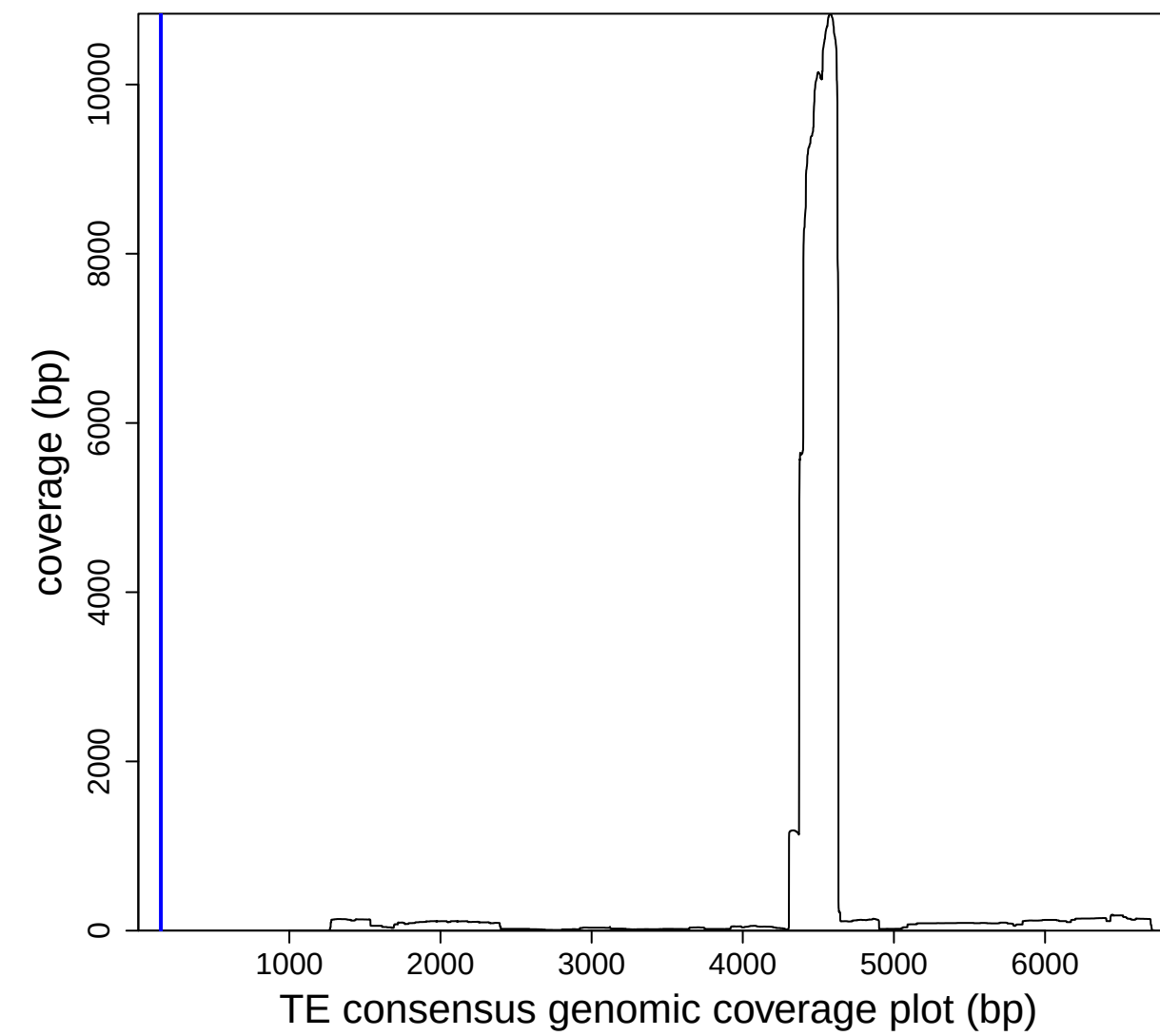
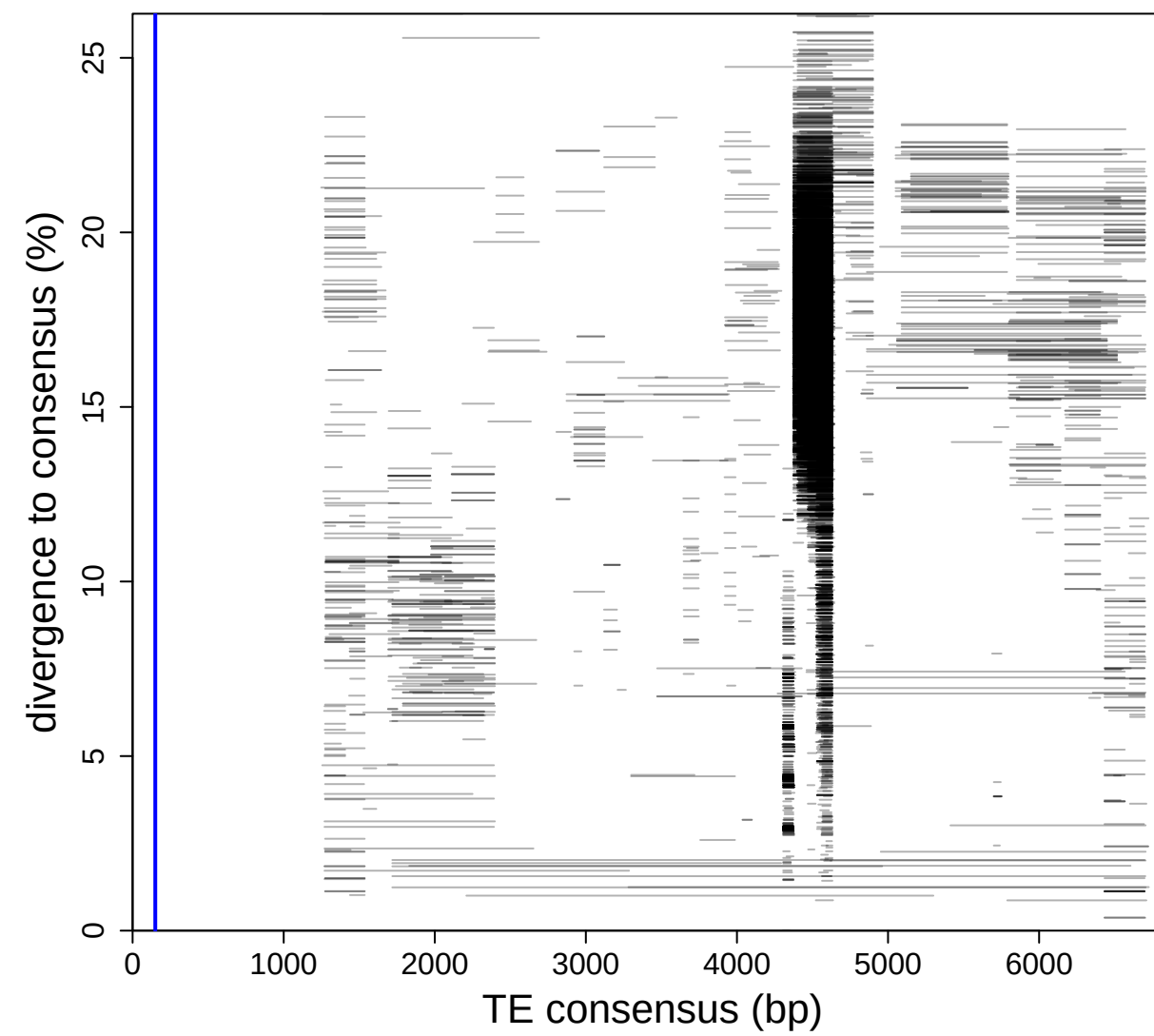
_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa al
size: 6637bp; fragments: 13181; full length: 0 (>=5973.3bp)

After TEtrimmer 6637 bp

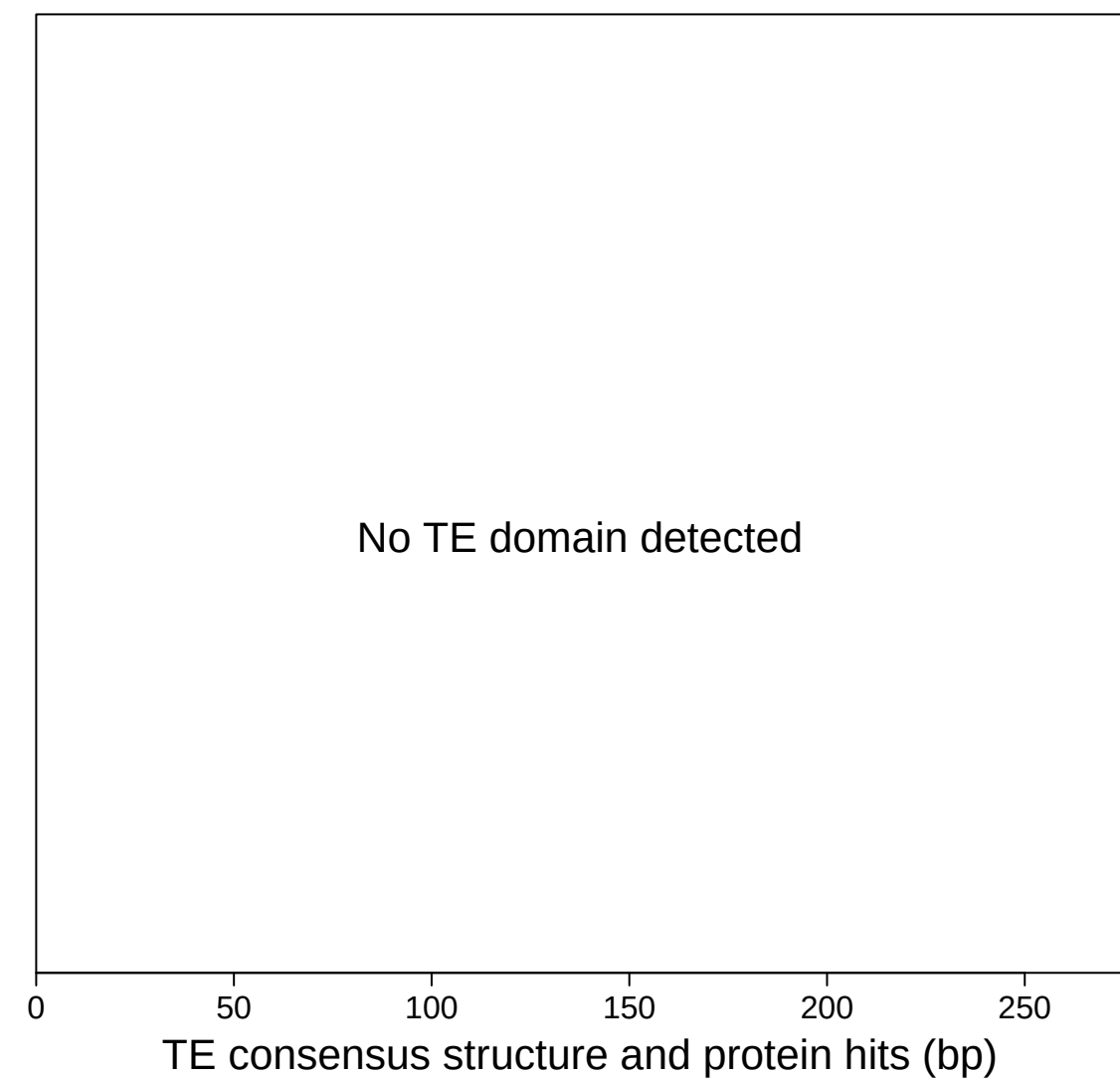
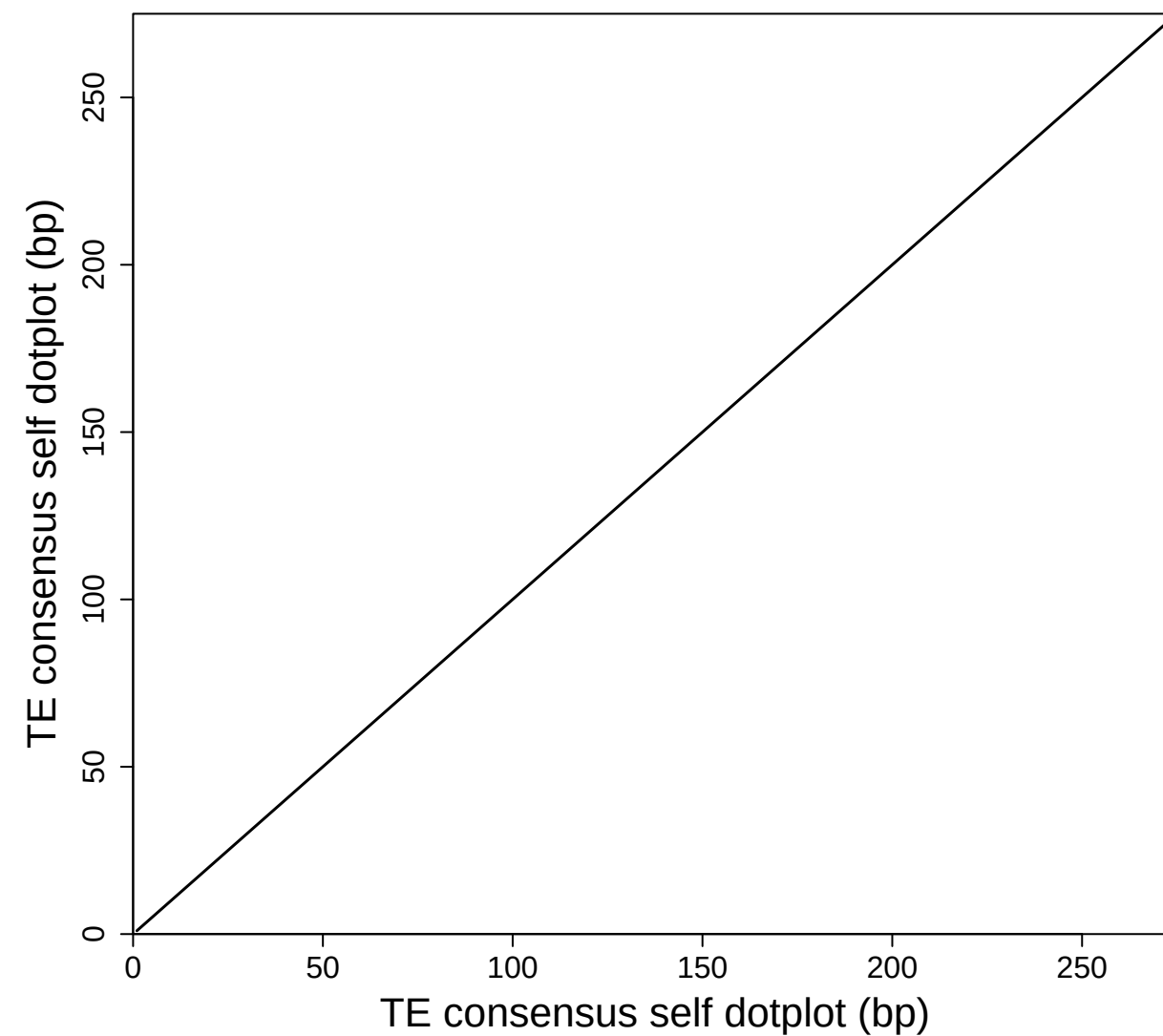
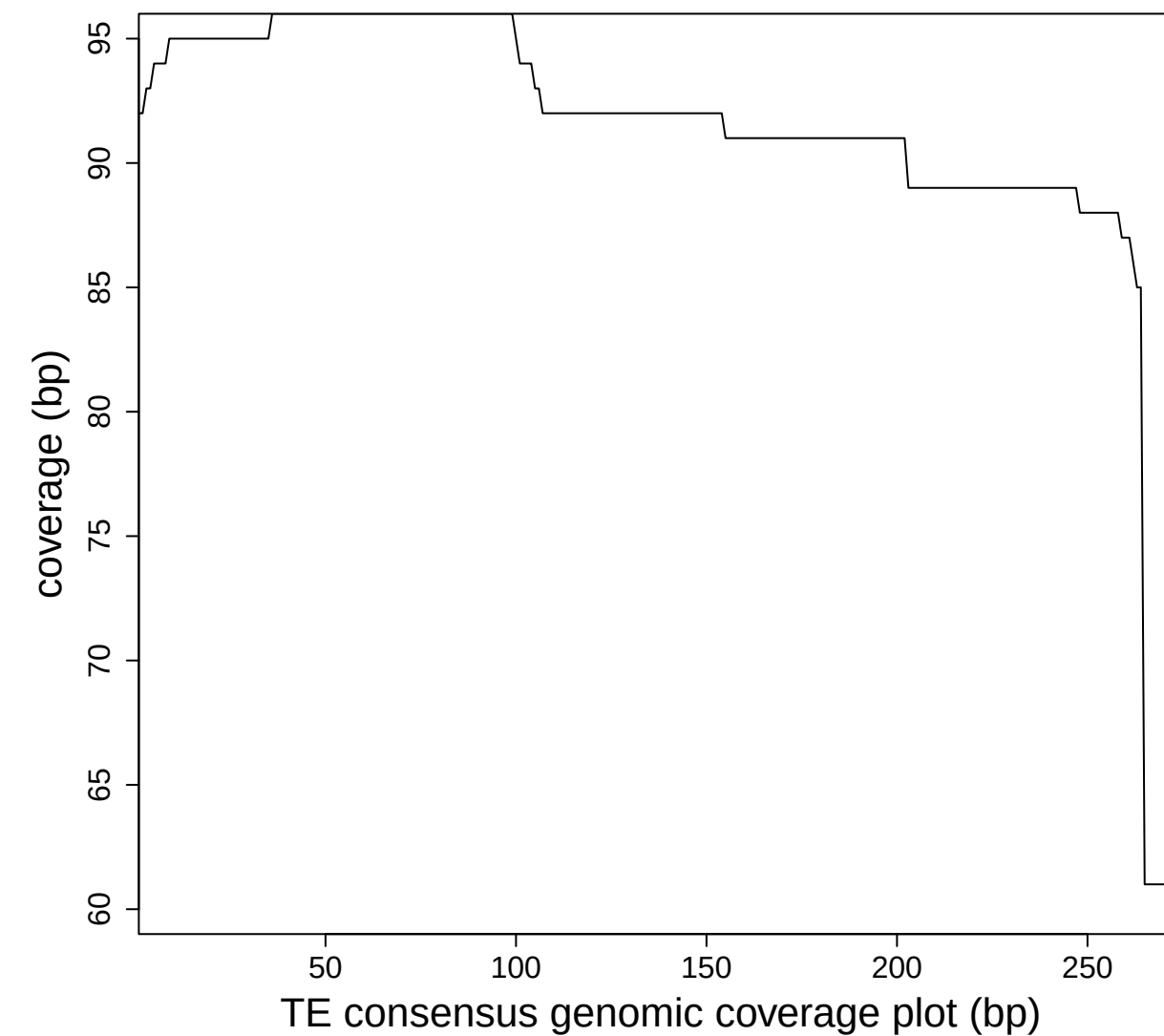
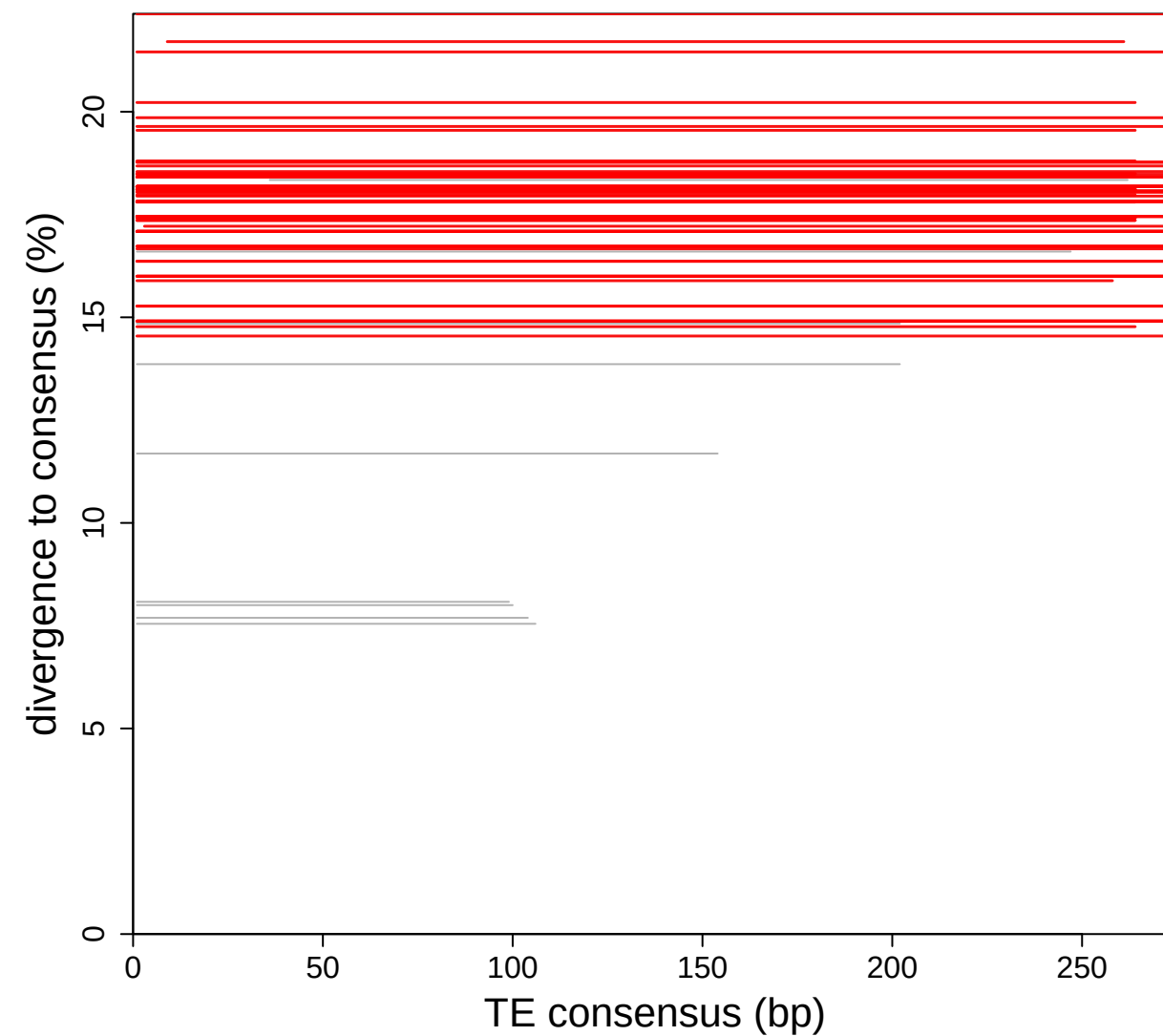


d_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa
size: 6884bp; fragments: 13153; full length: 0 (>=6195.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

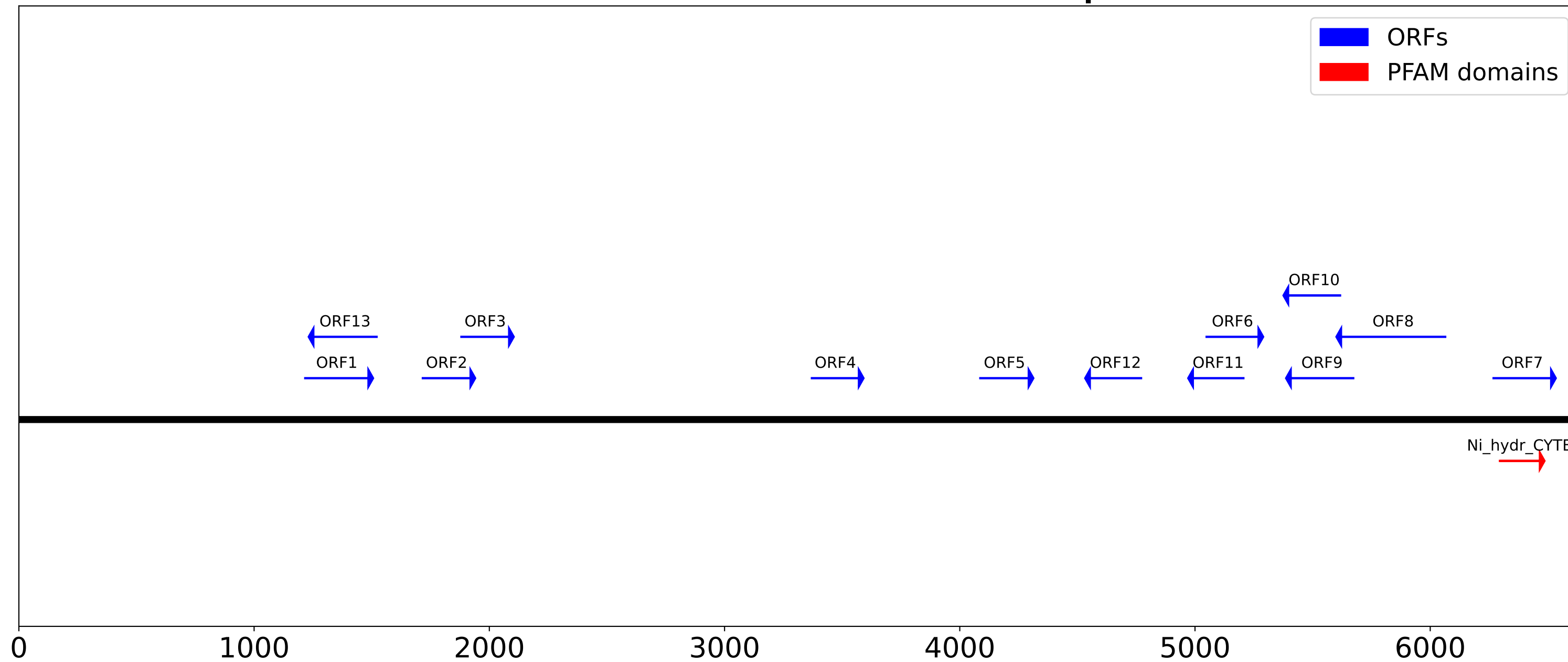


TE: ltr_1_family_307
size: 275bp; fragments: 96; full length: 87 (≥ 247.5 bp)

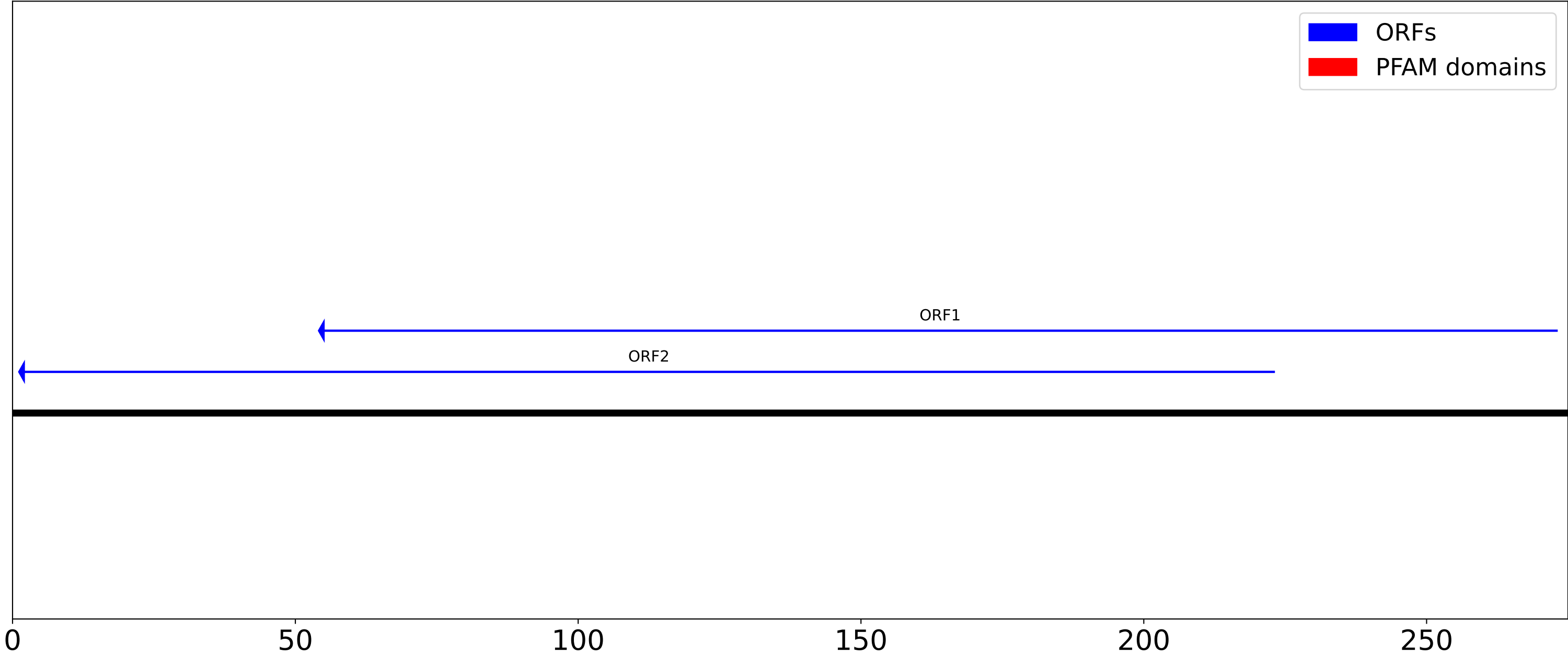


Before TEtrimmer 275 bp

After TEtrimmer ORF and PFAM domain plot



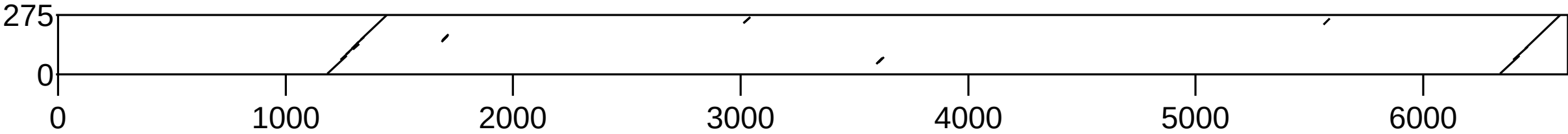
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)