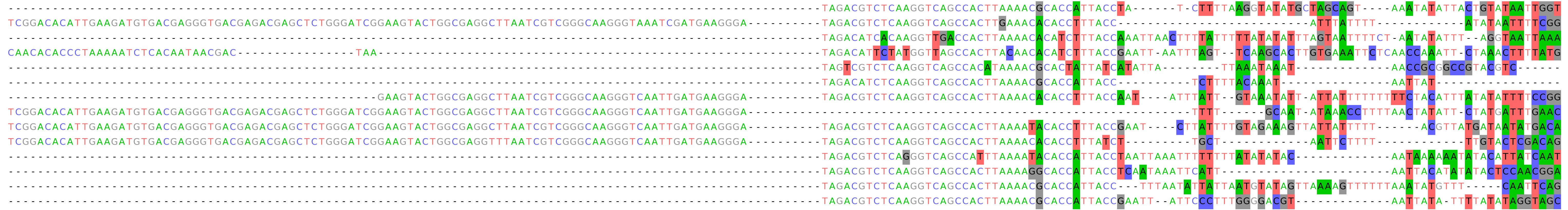
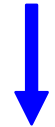


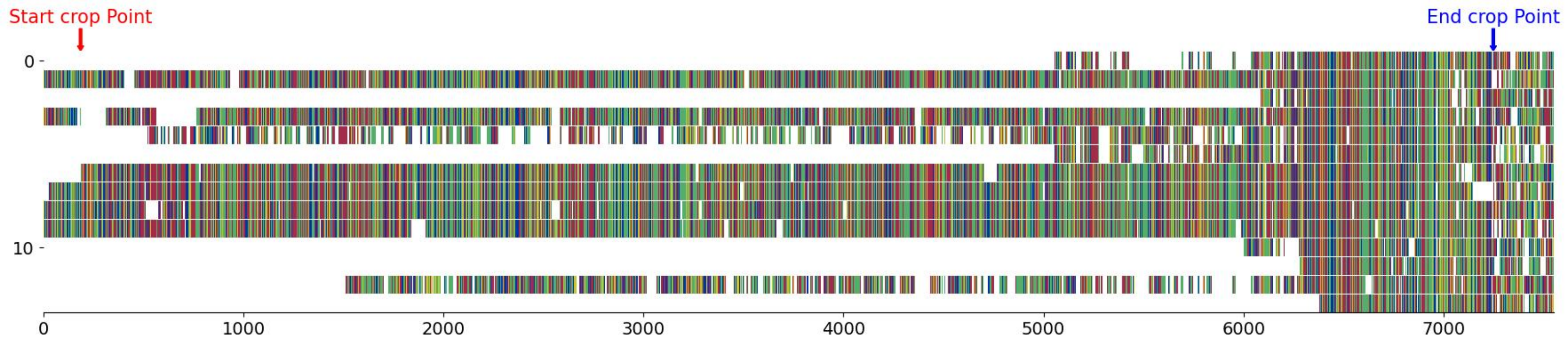
Start crop Point



MSA length = 7064

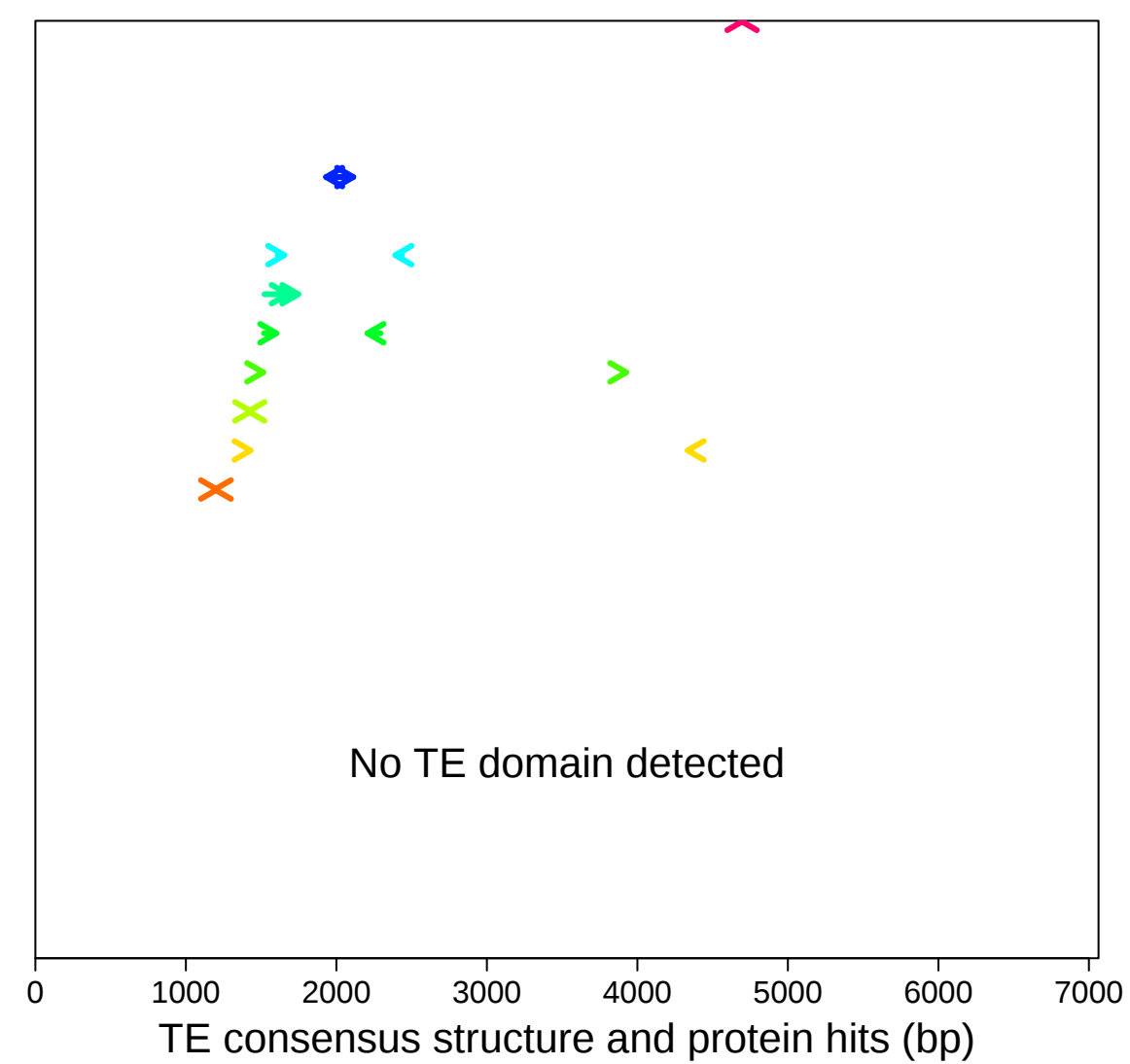
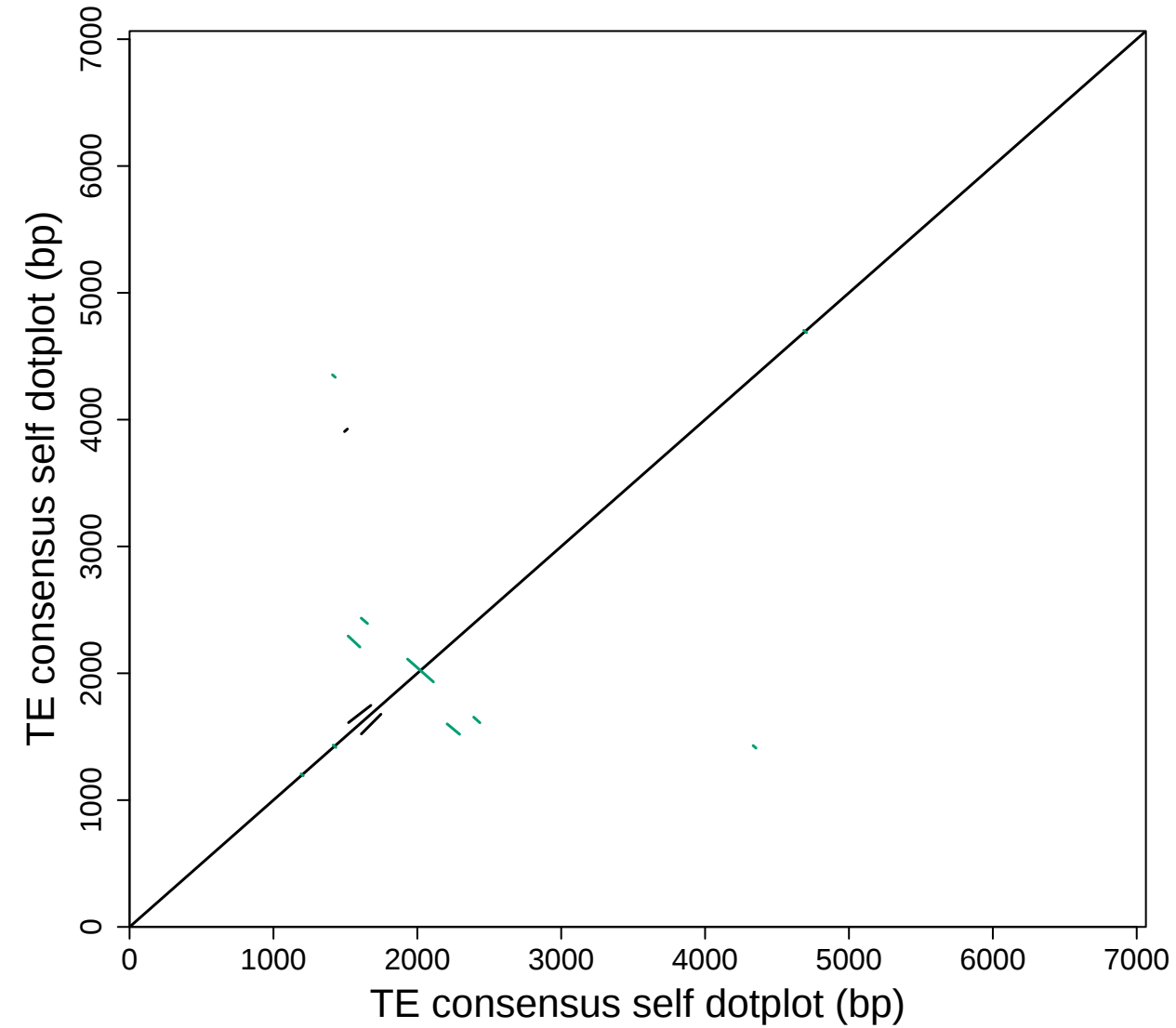
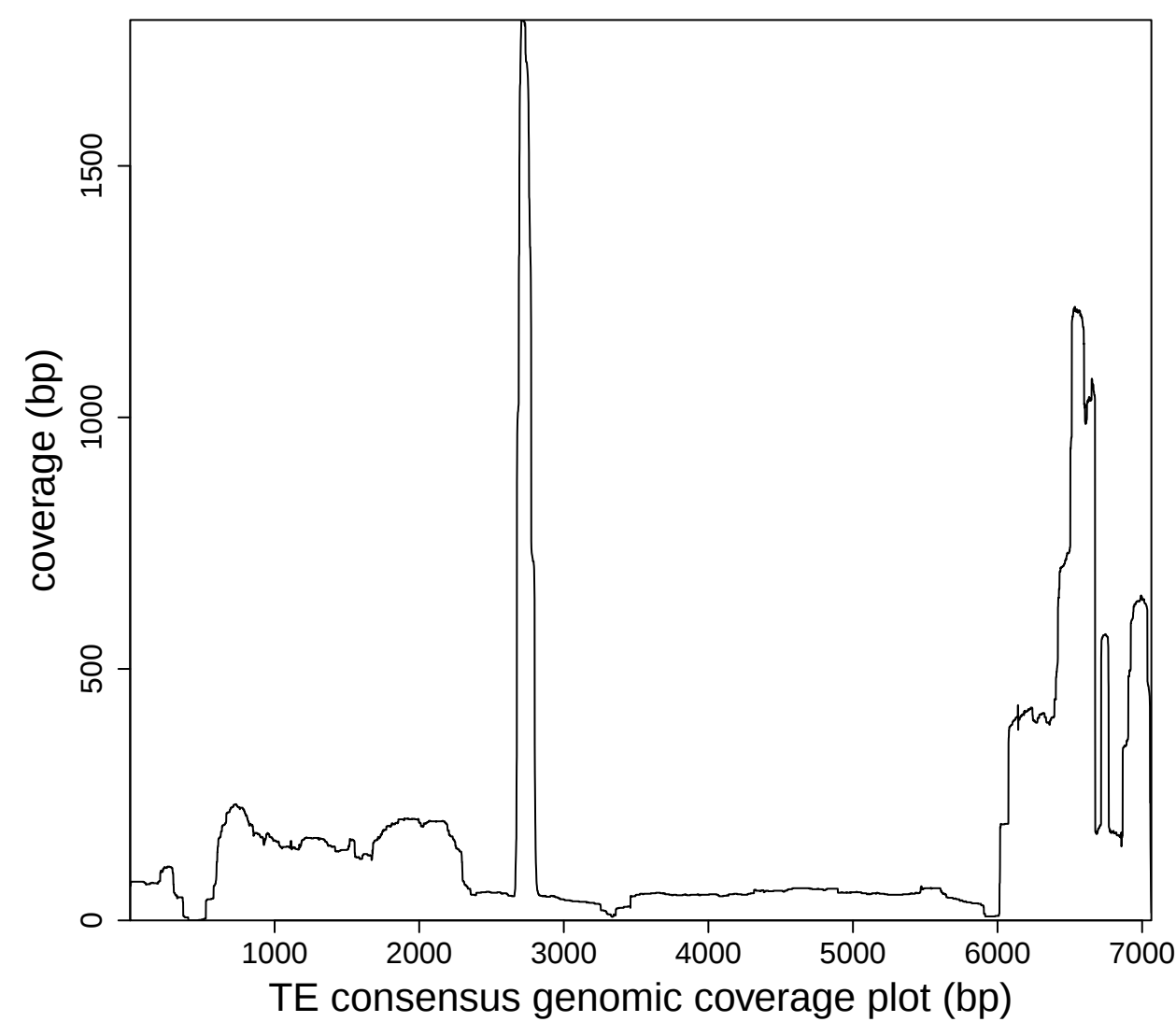
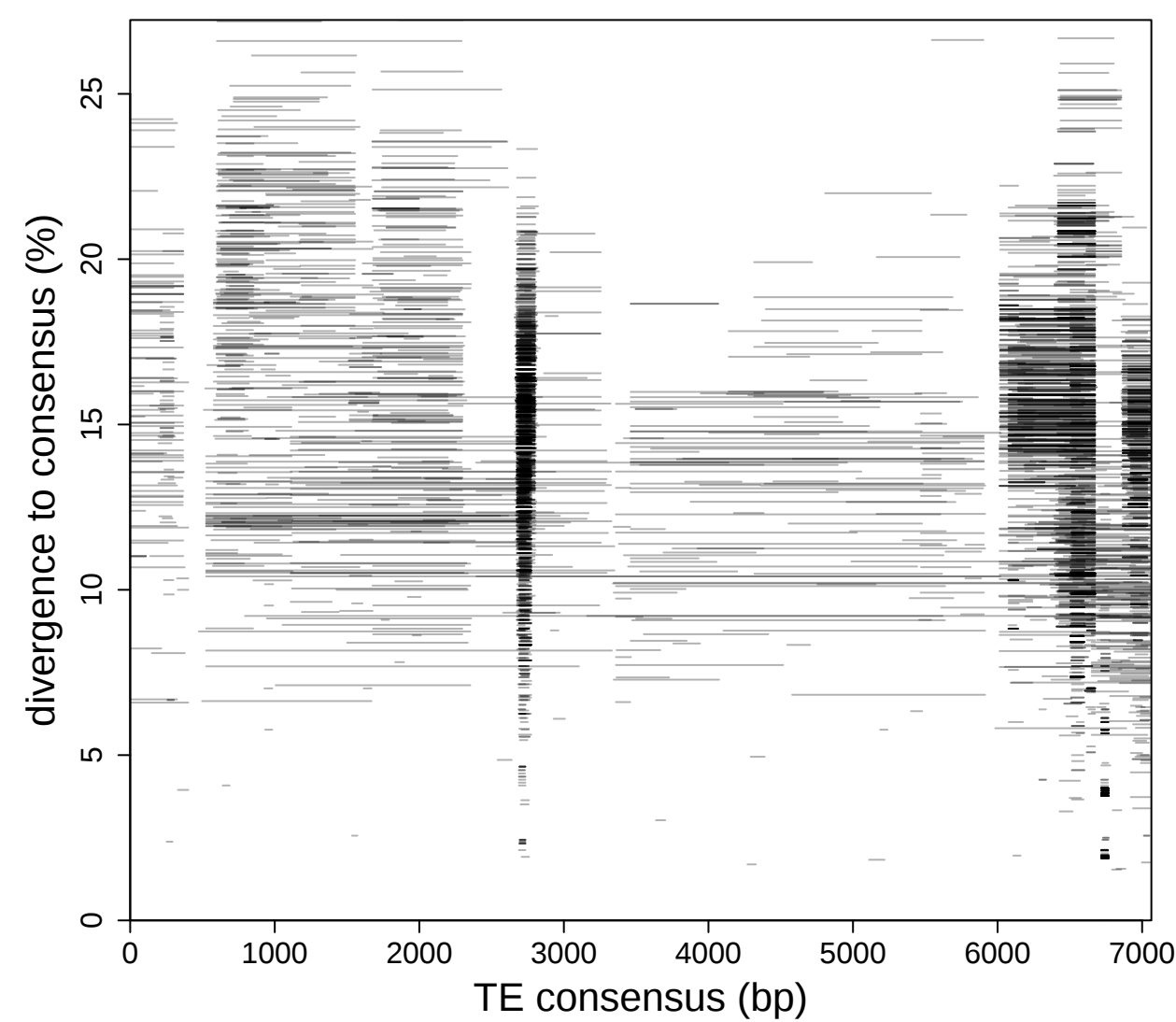
End crop Point





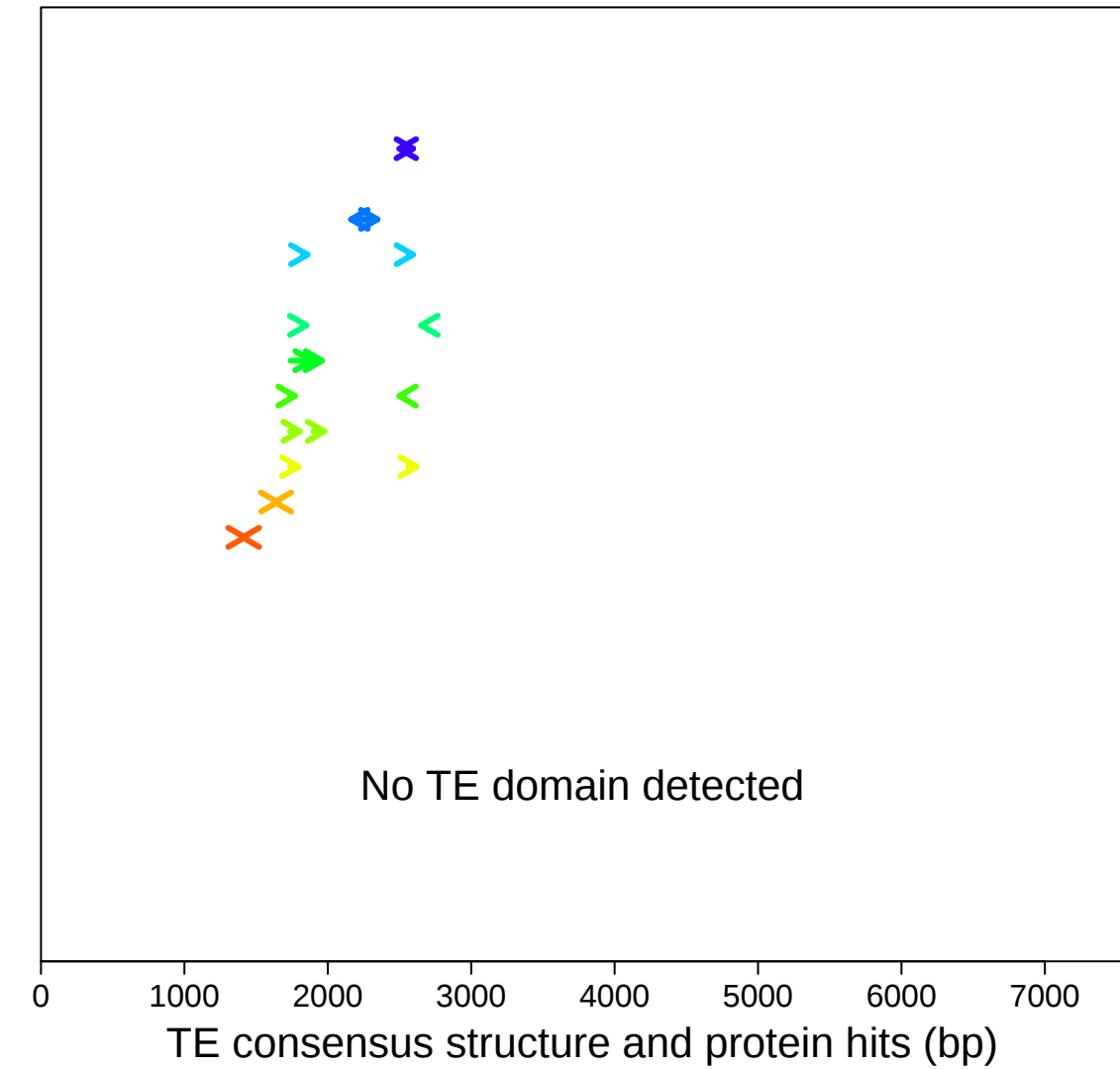
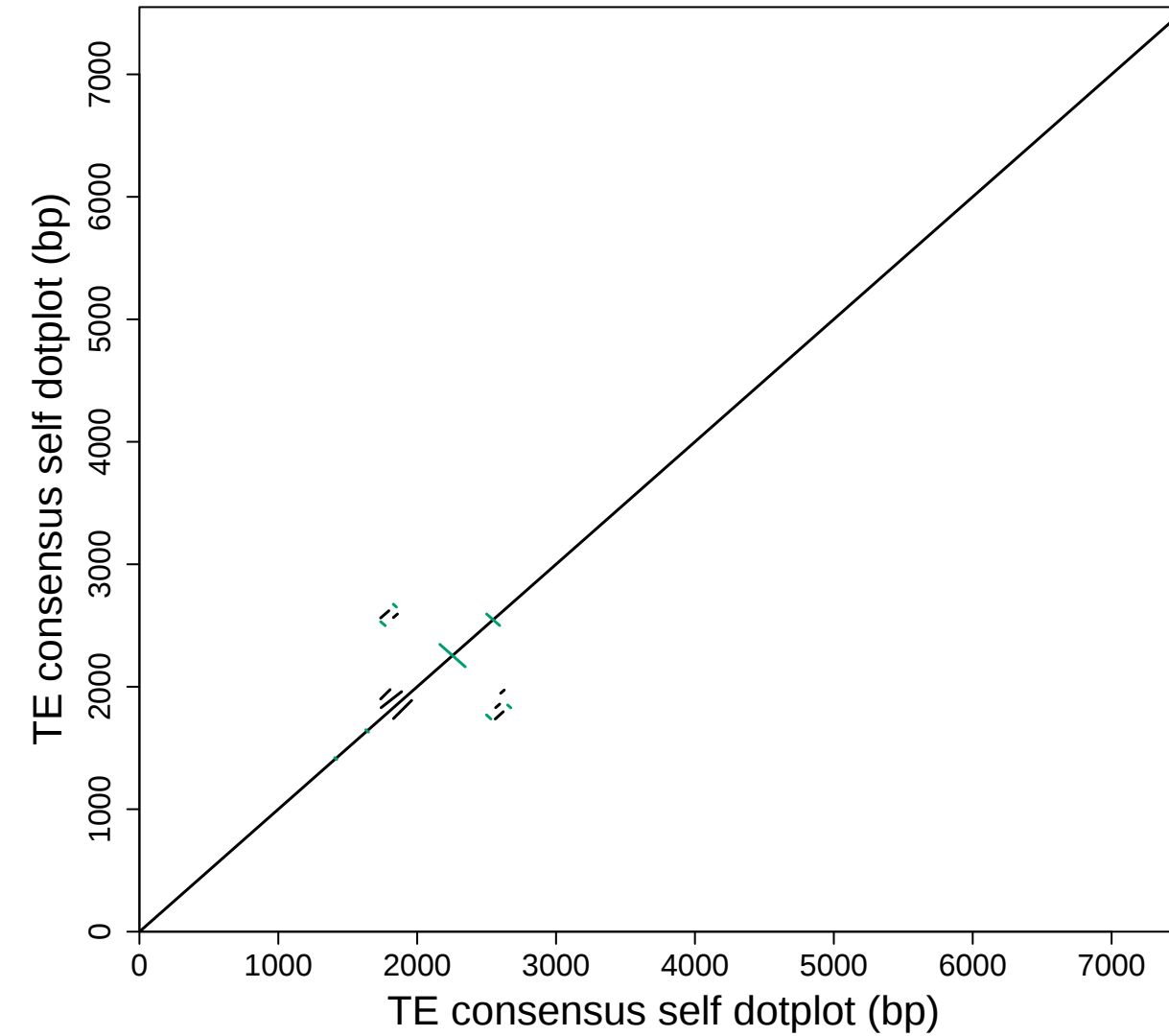
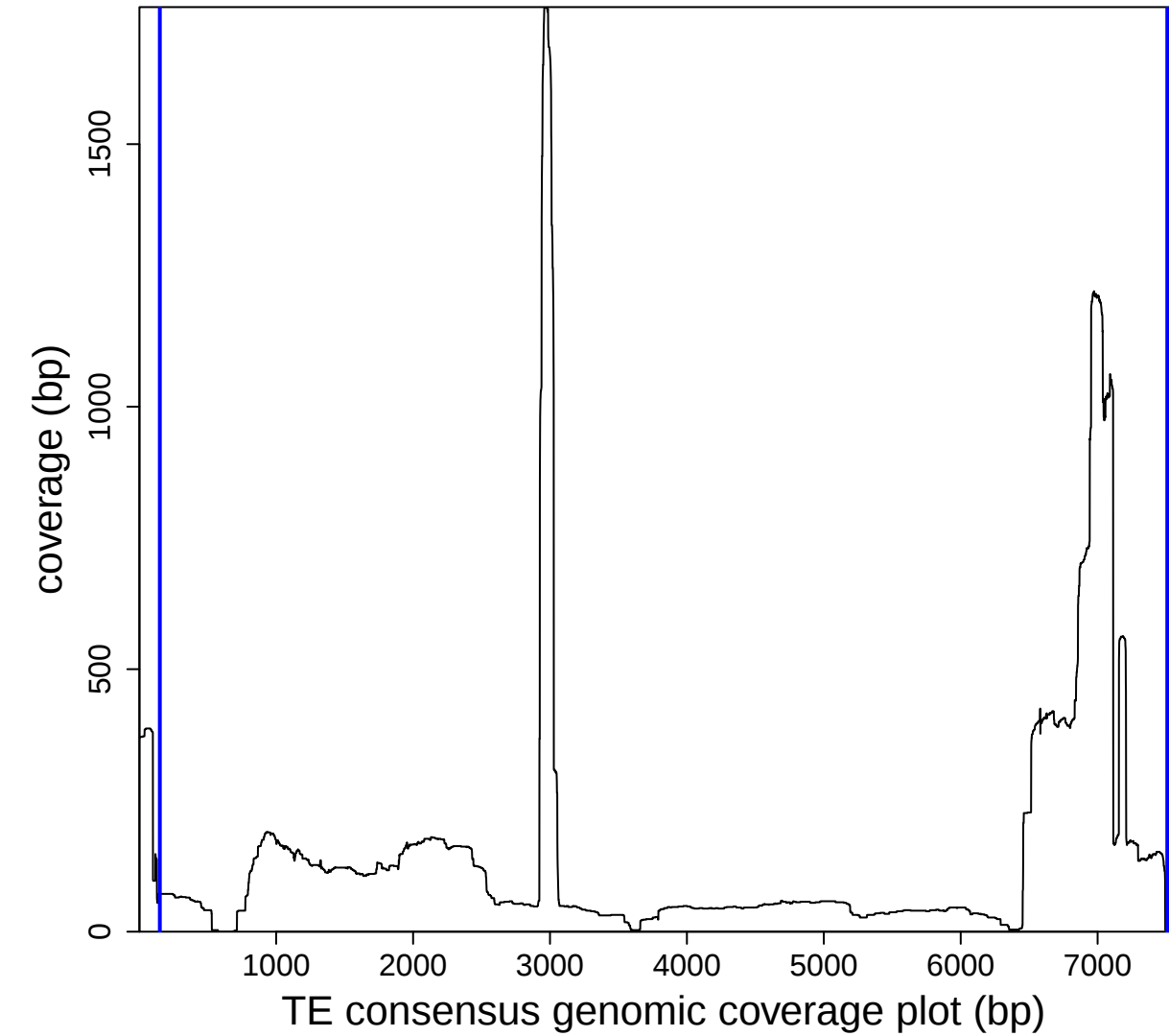
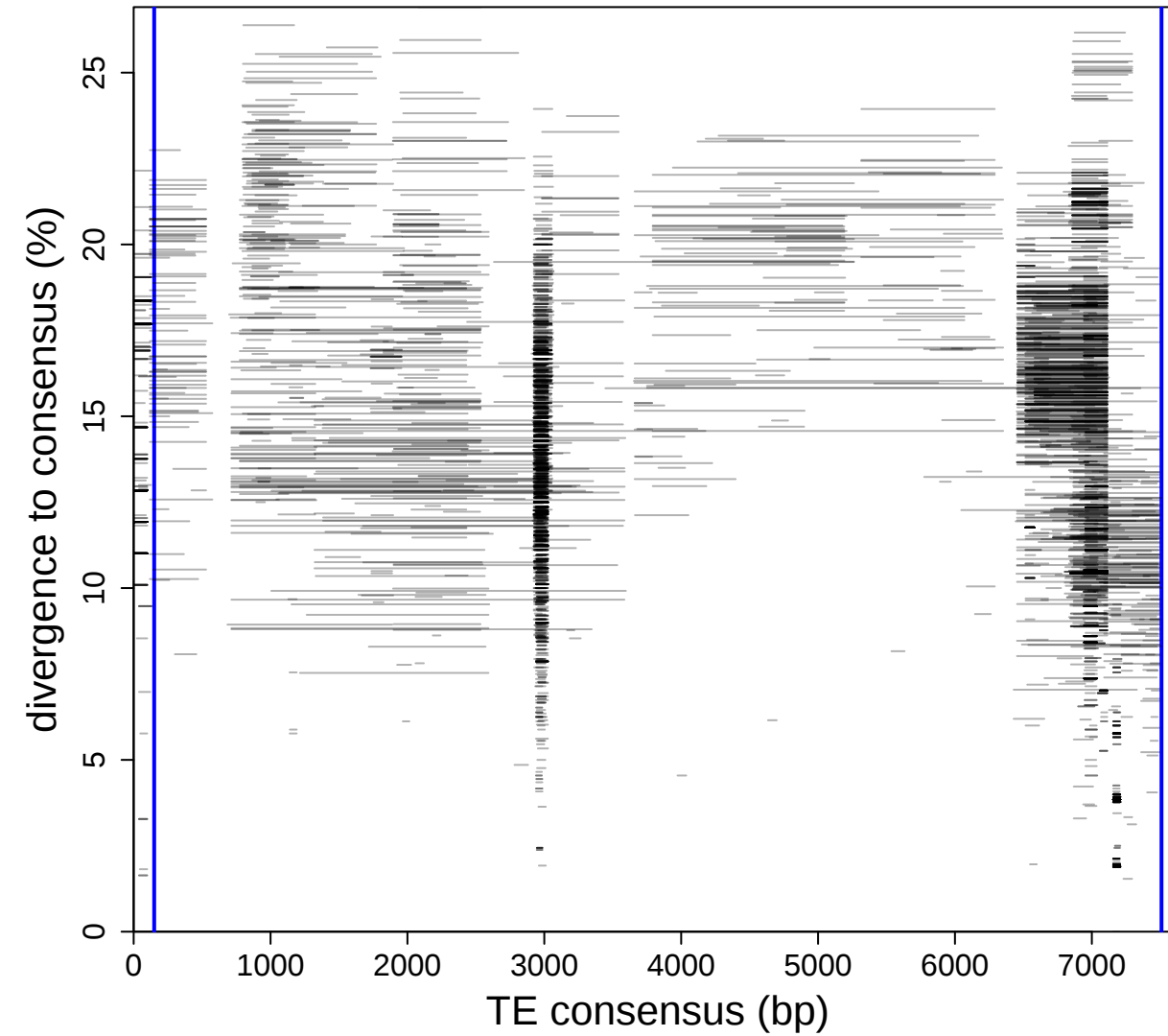
l.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 7064bp; fragments: 5058; full length: 0 (>=6357.6bp)

After TEtrimmer 7064 bp

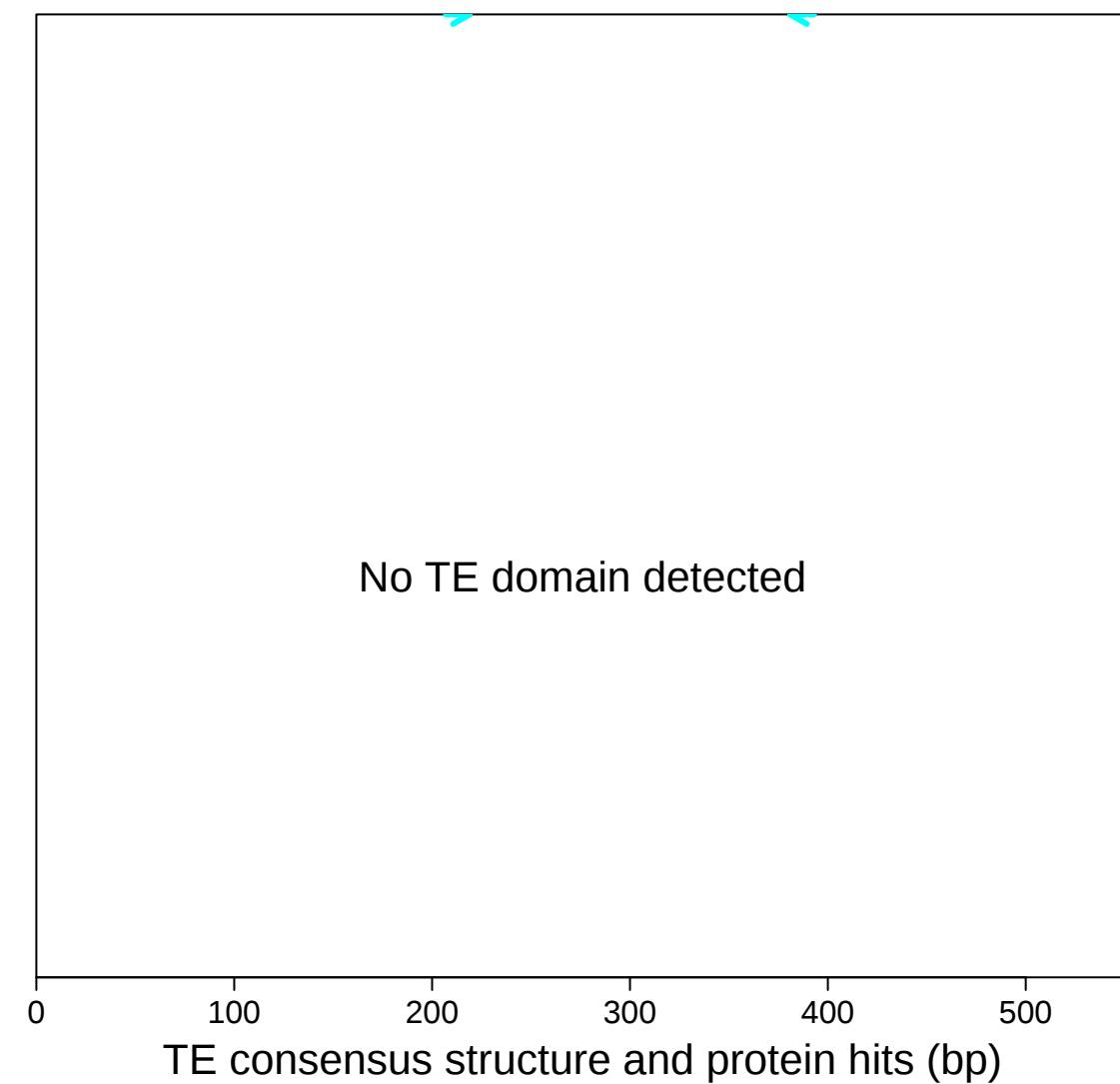
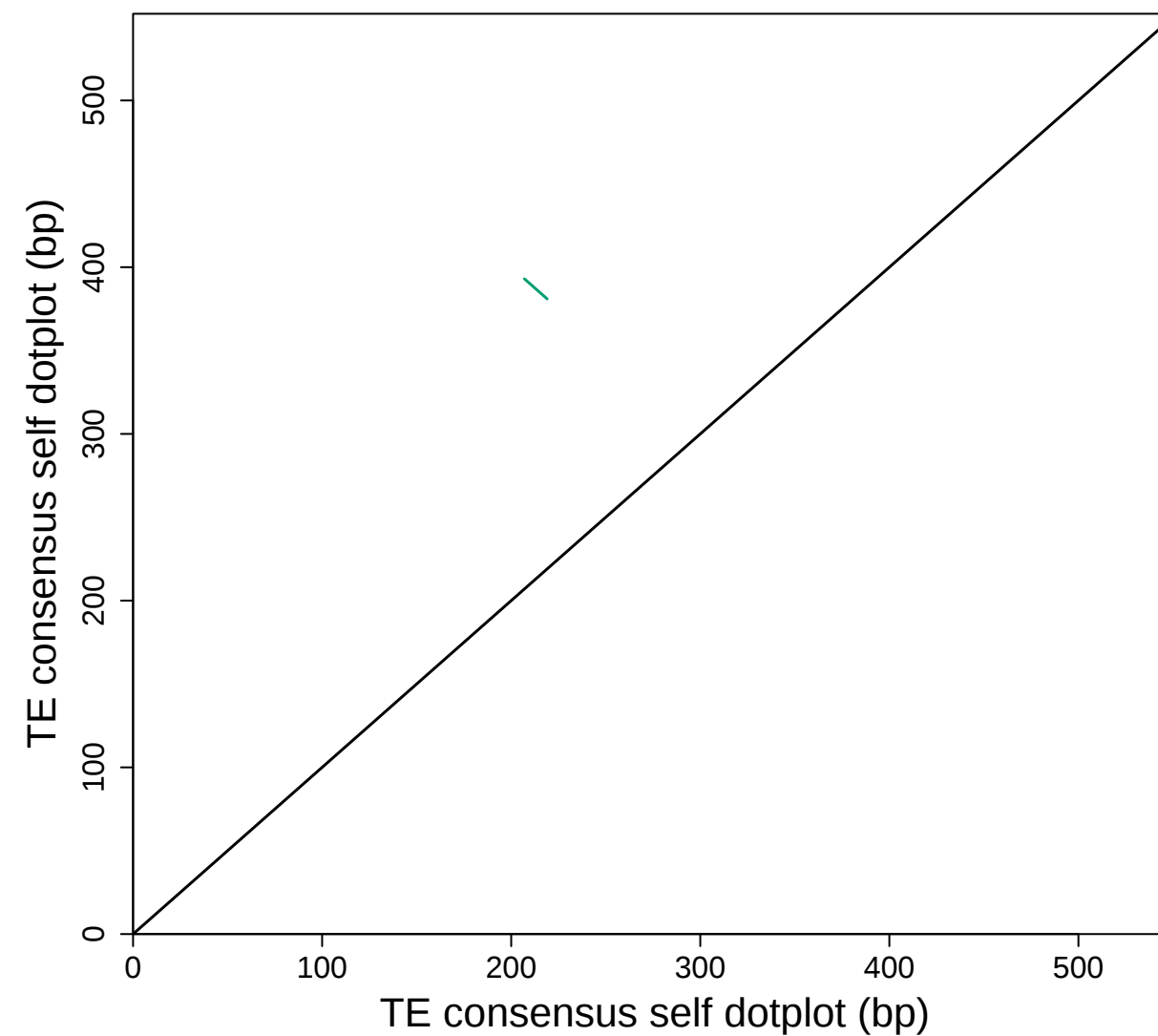
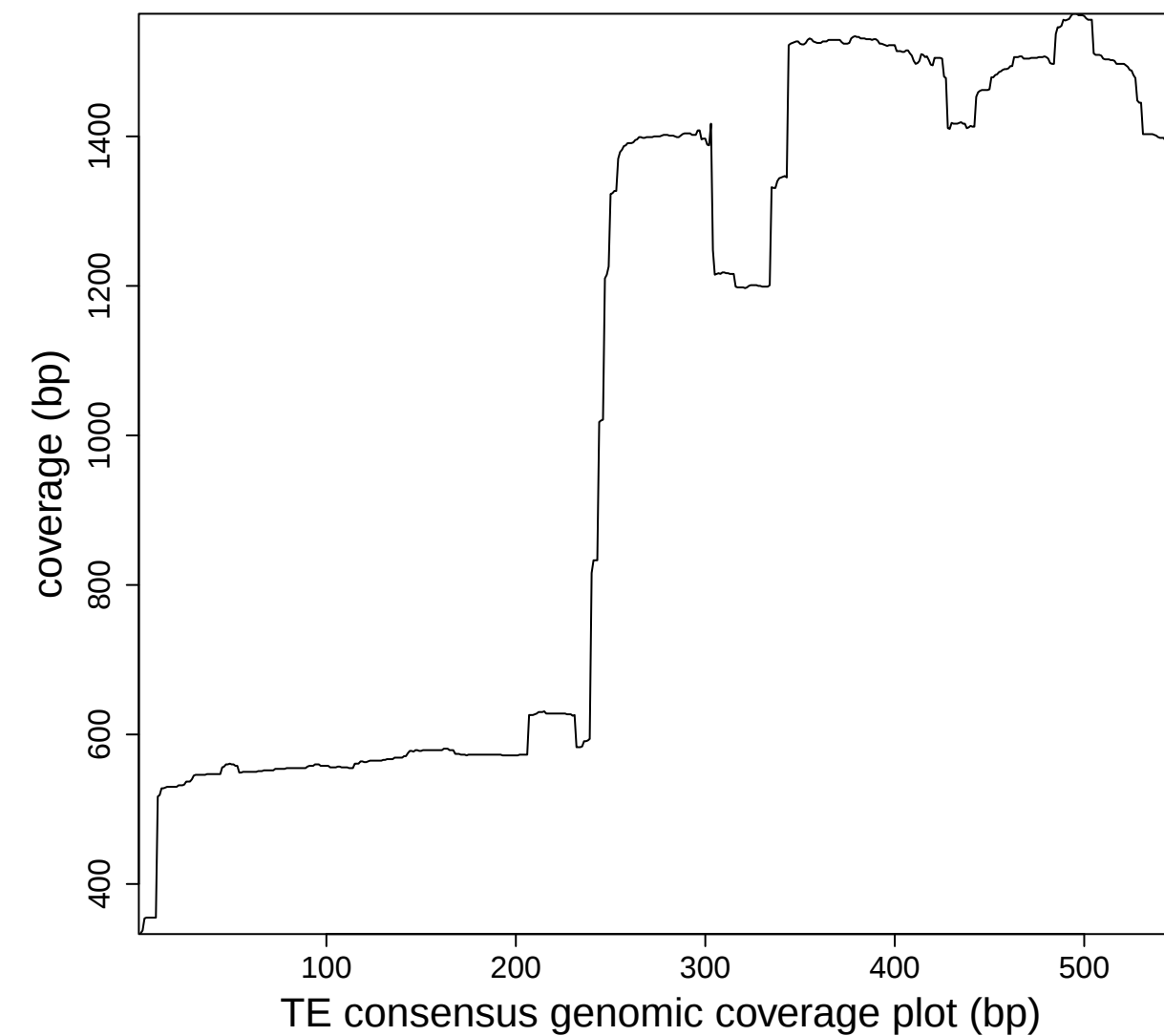
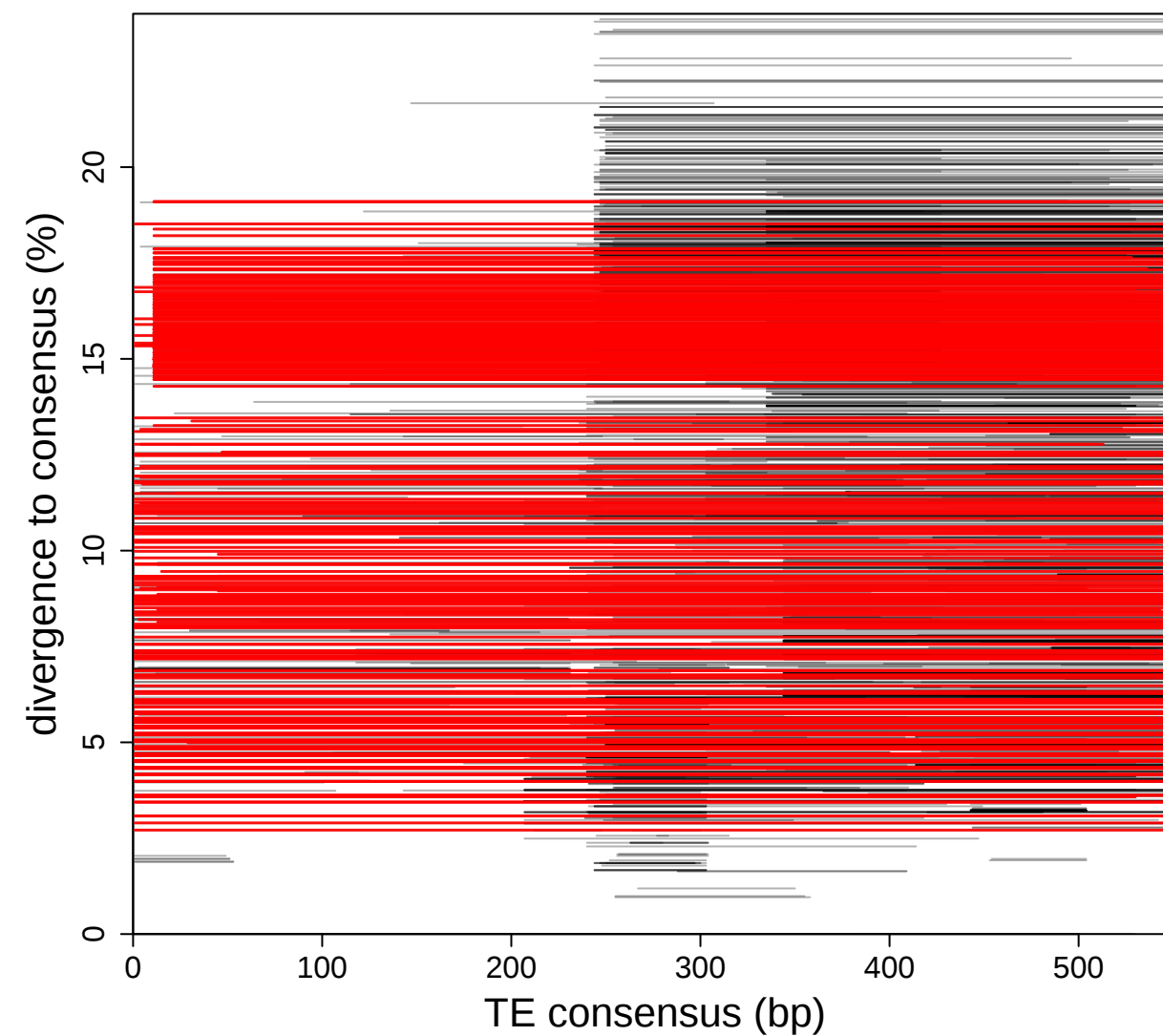


9.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 7659bp; fragments: 4735; full length: 0 (>=6893.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

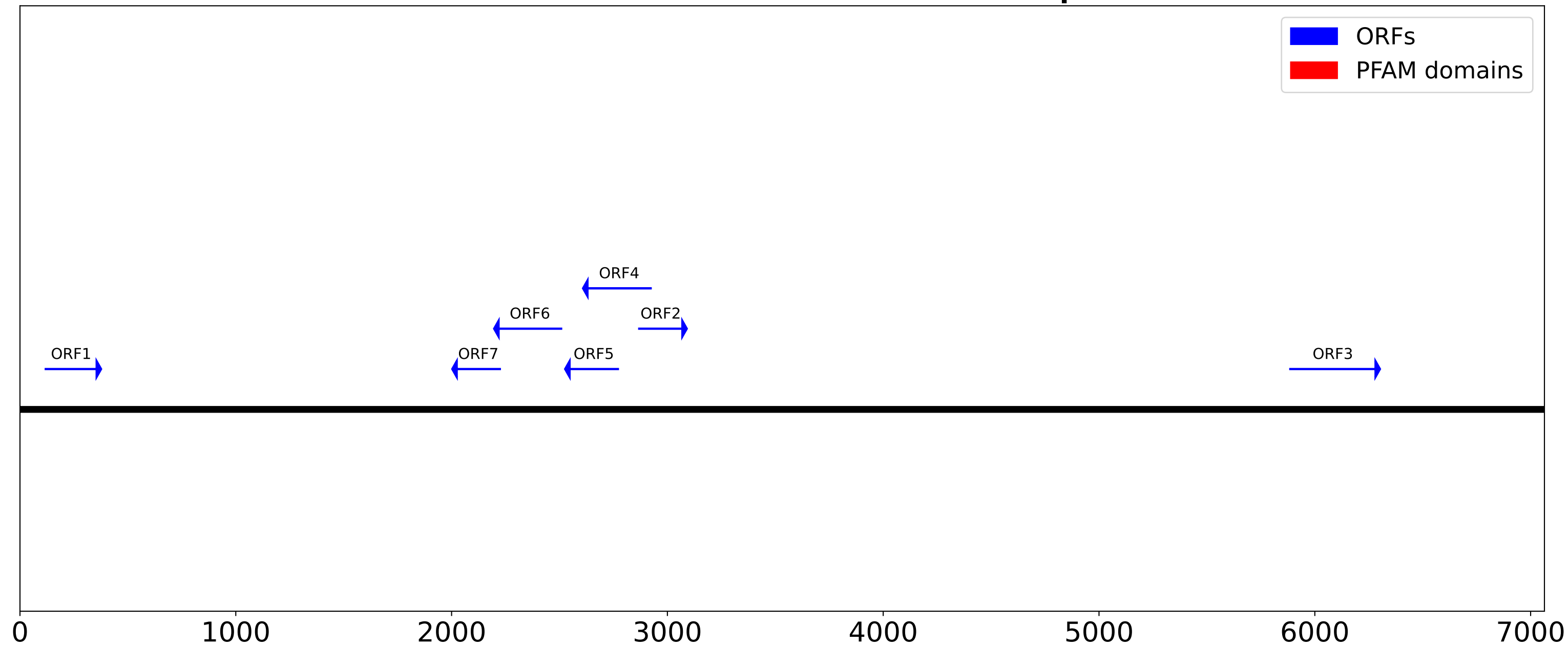


TE: rnd_1_family_879
size: 552bp; fragments: 2173; full length: 408 (≥ 496.8 bp)



Before TEtrimmer 552 bp

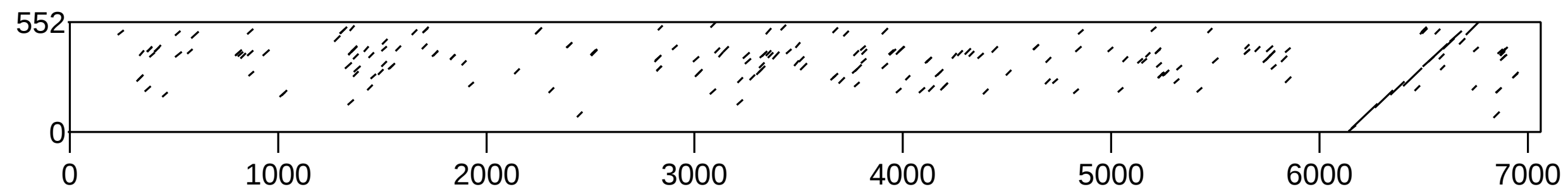
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

ensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)