

Start crop Point

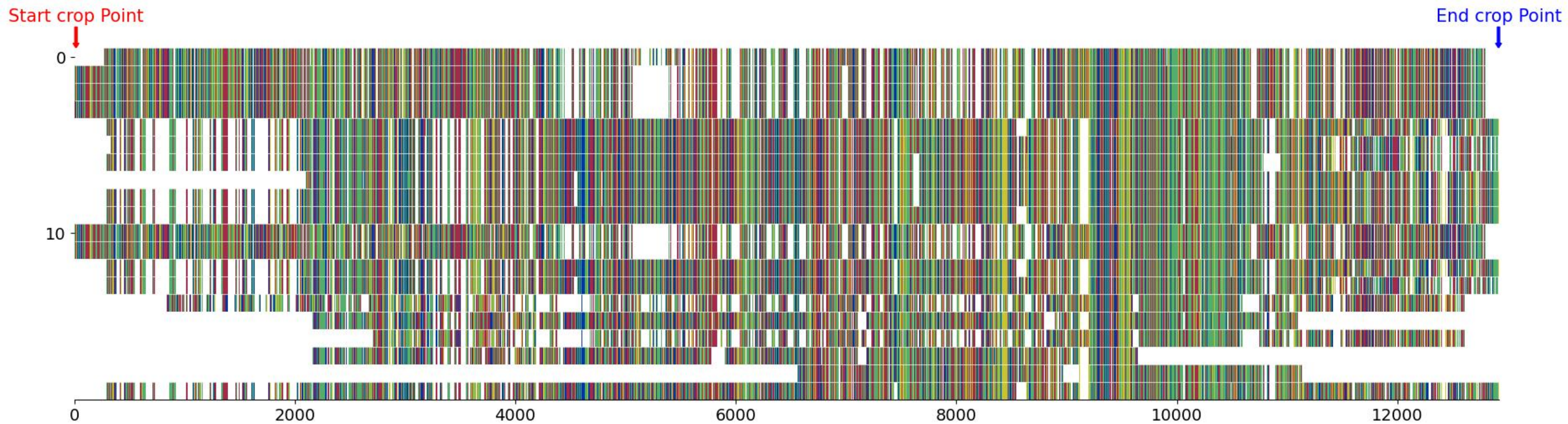


End crop Point



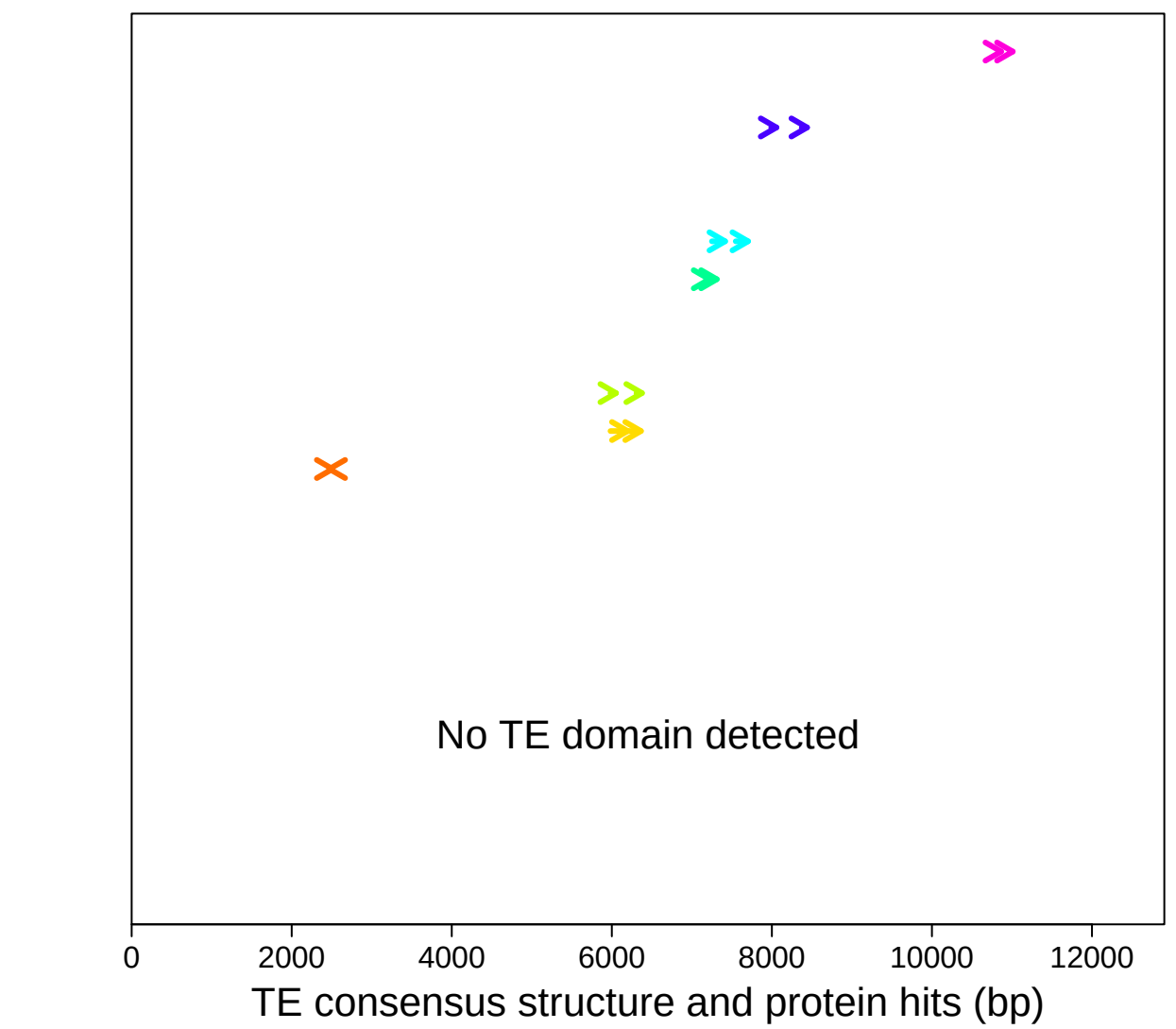
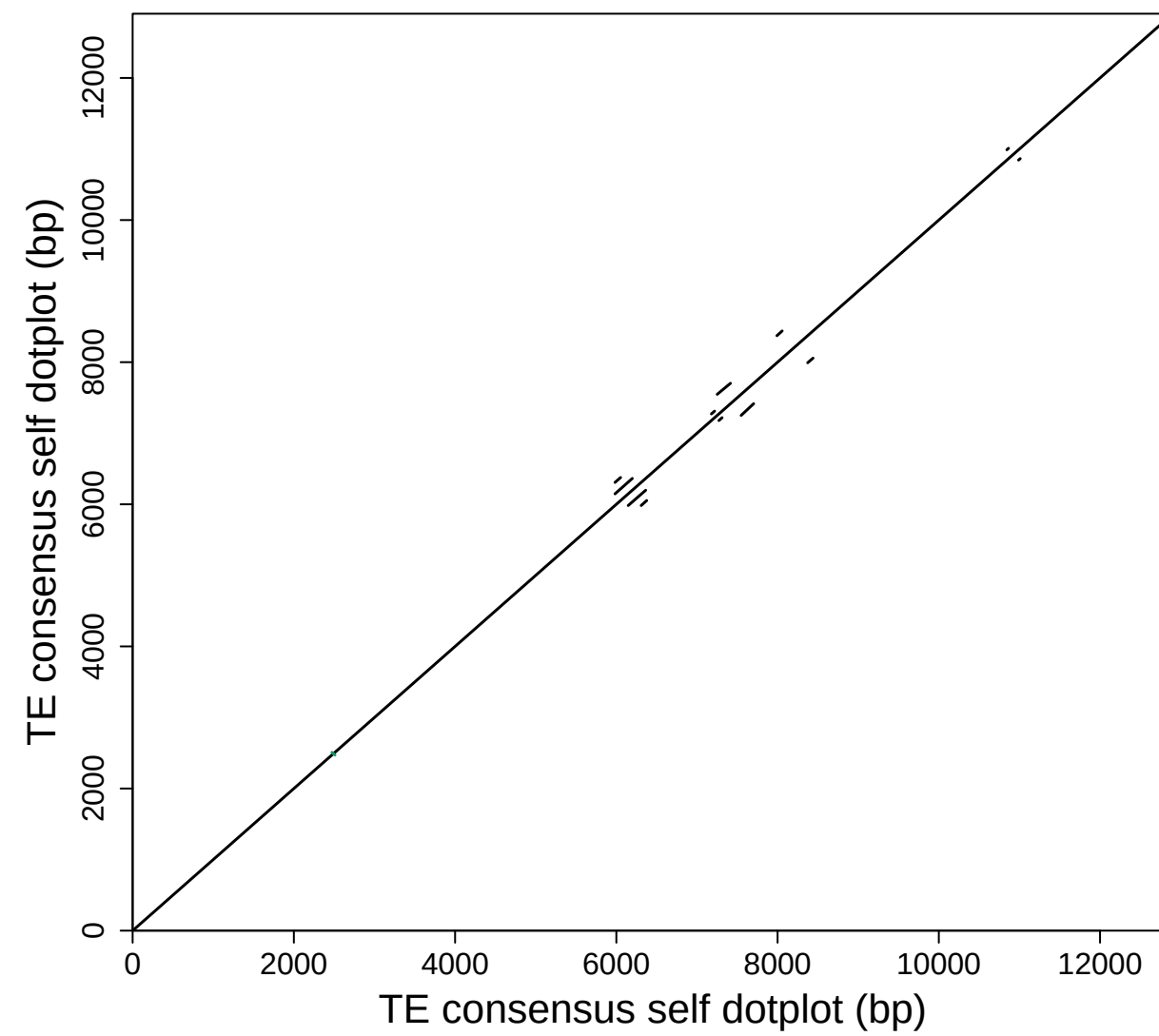
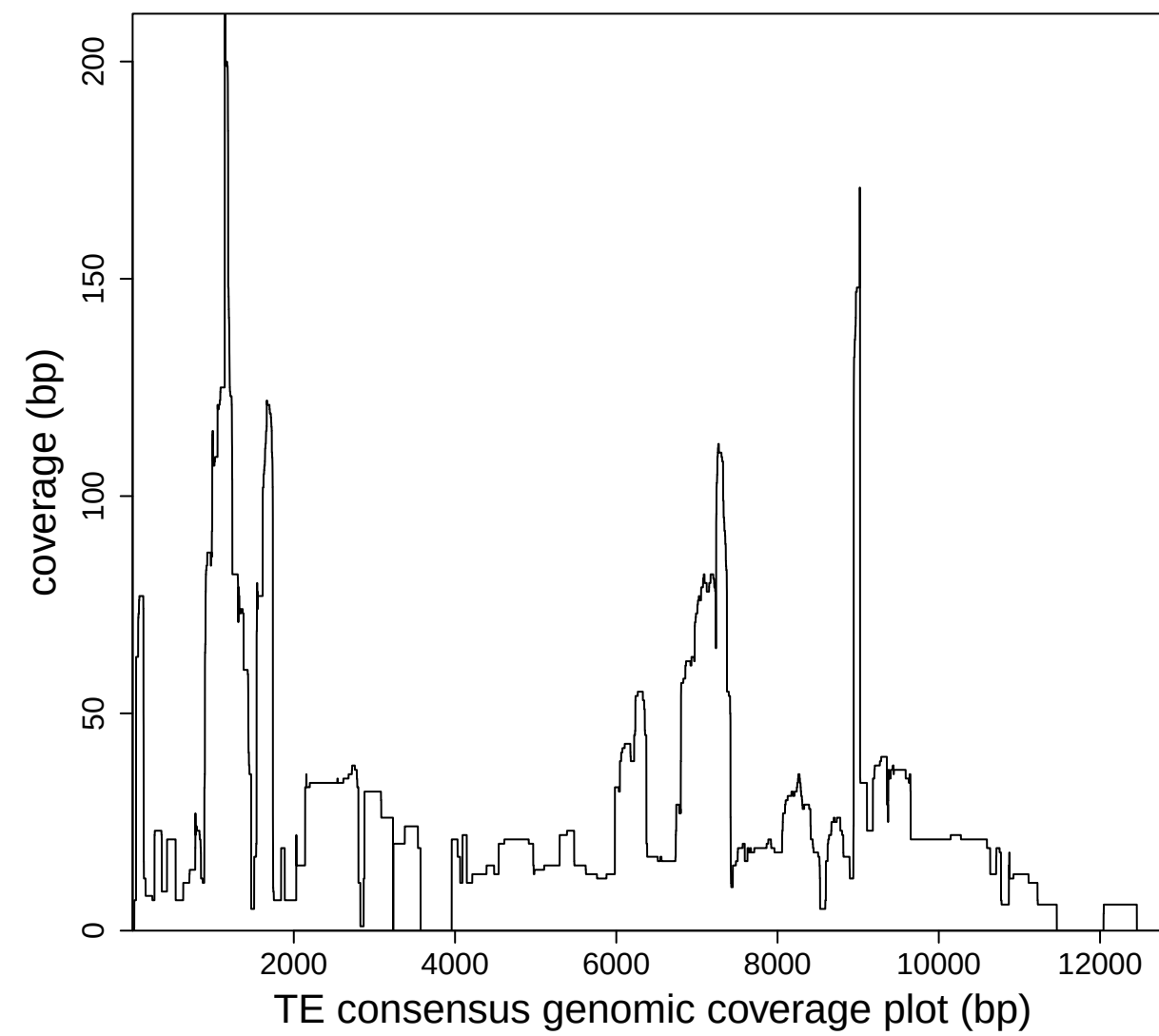
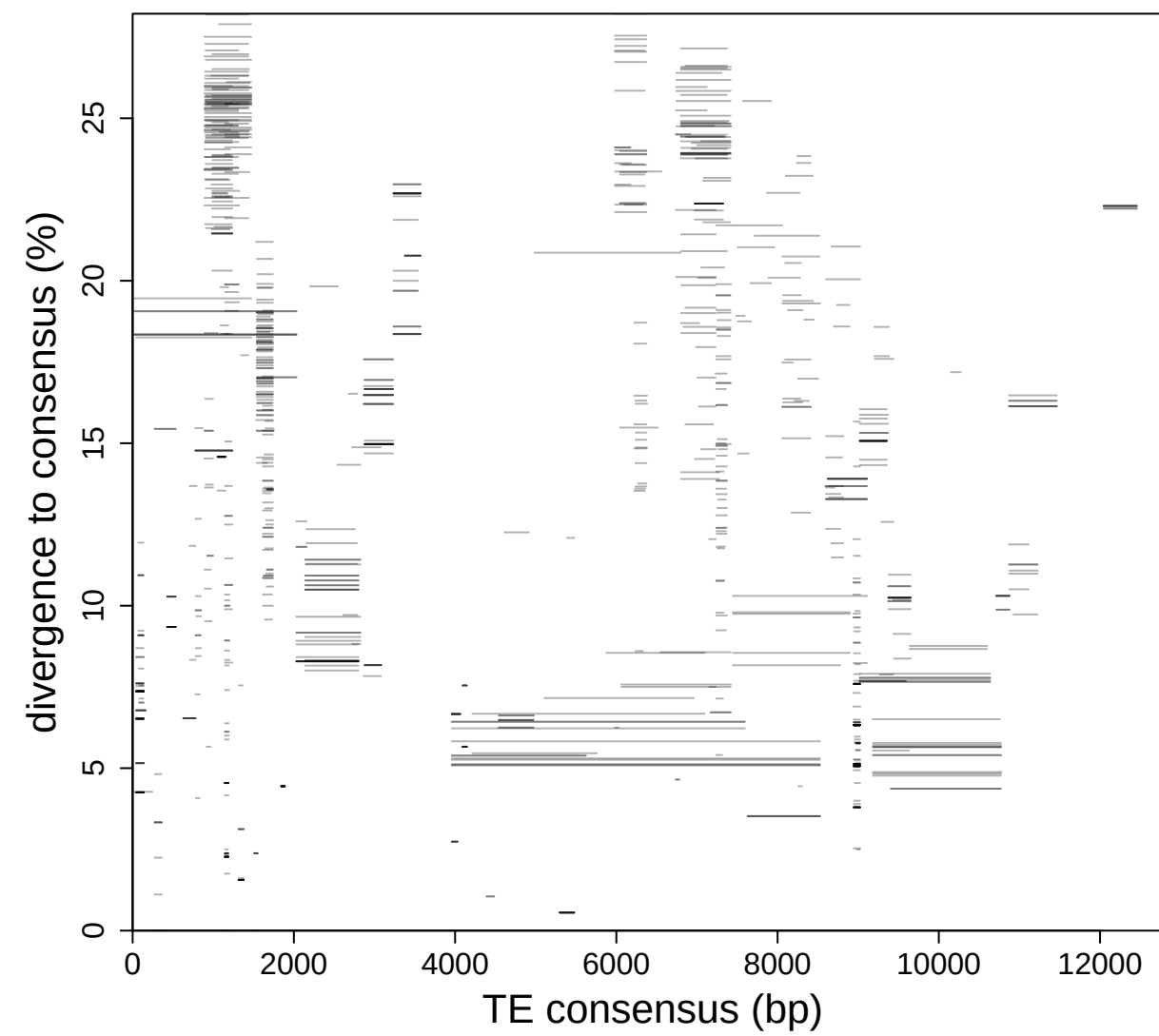
MSA length = 12904





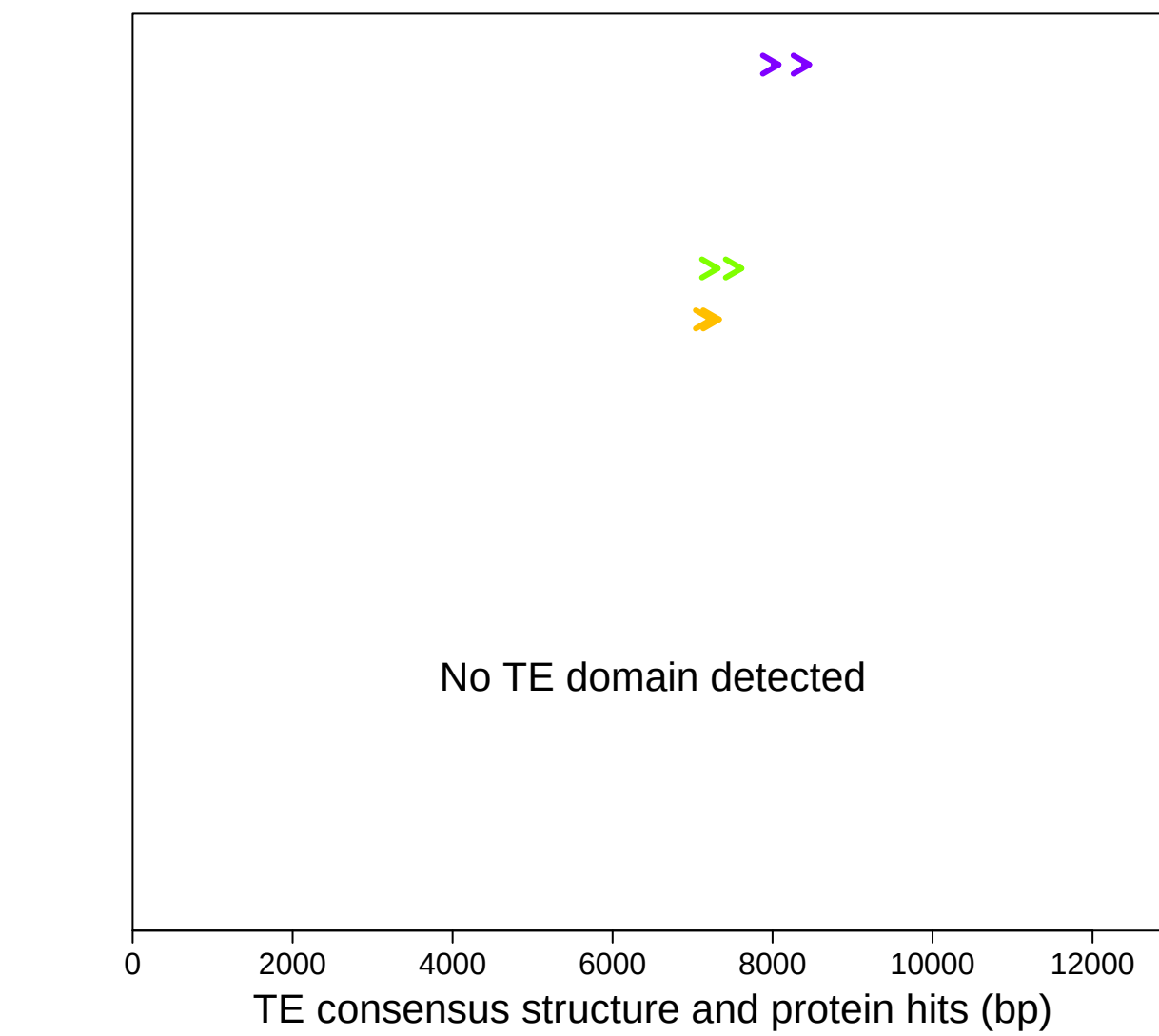
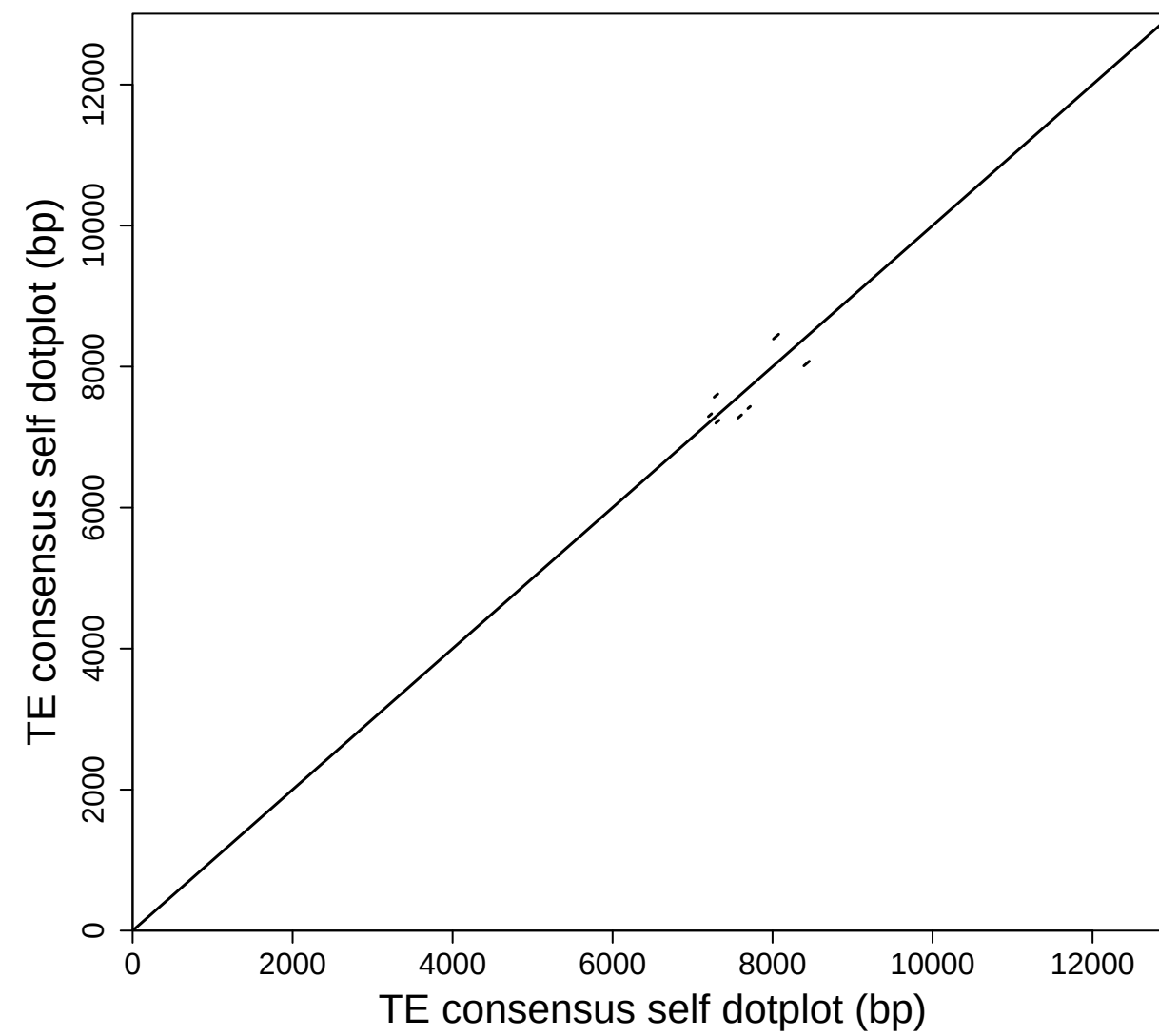
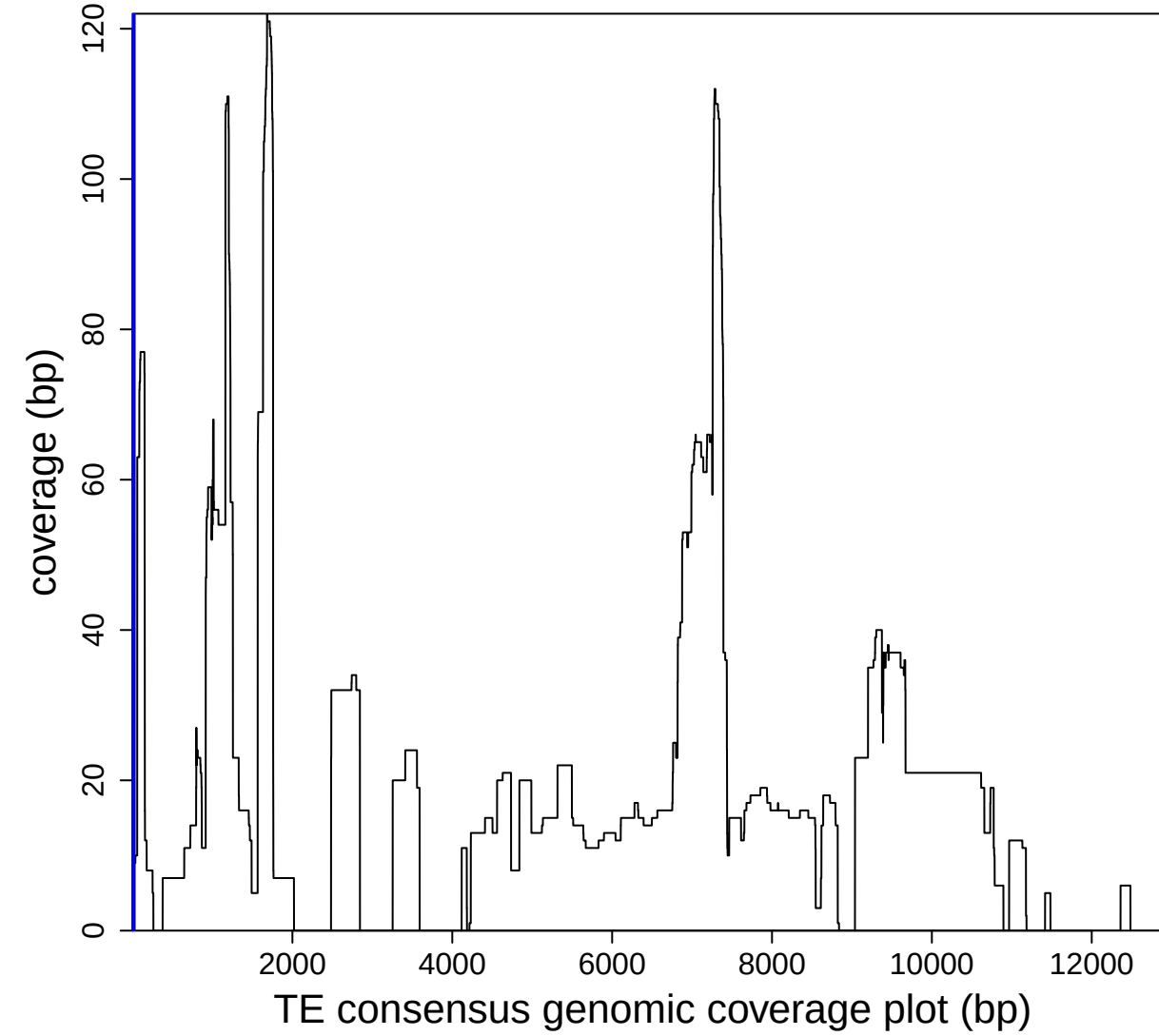
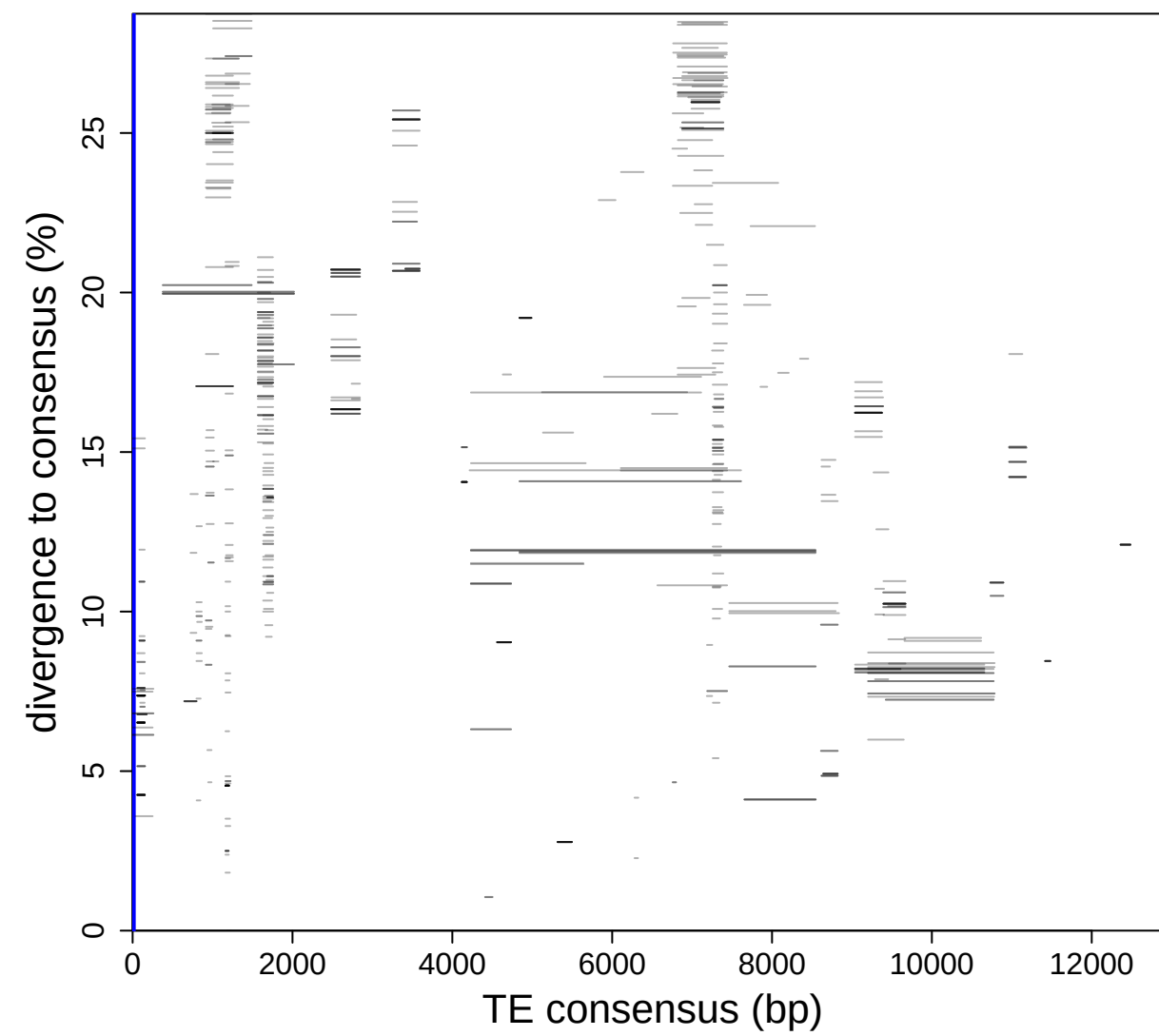
b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 12904bp; fragments: 1103; full length: 0 (>=11613.6bp)

After TEtrimmer 12904 bp

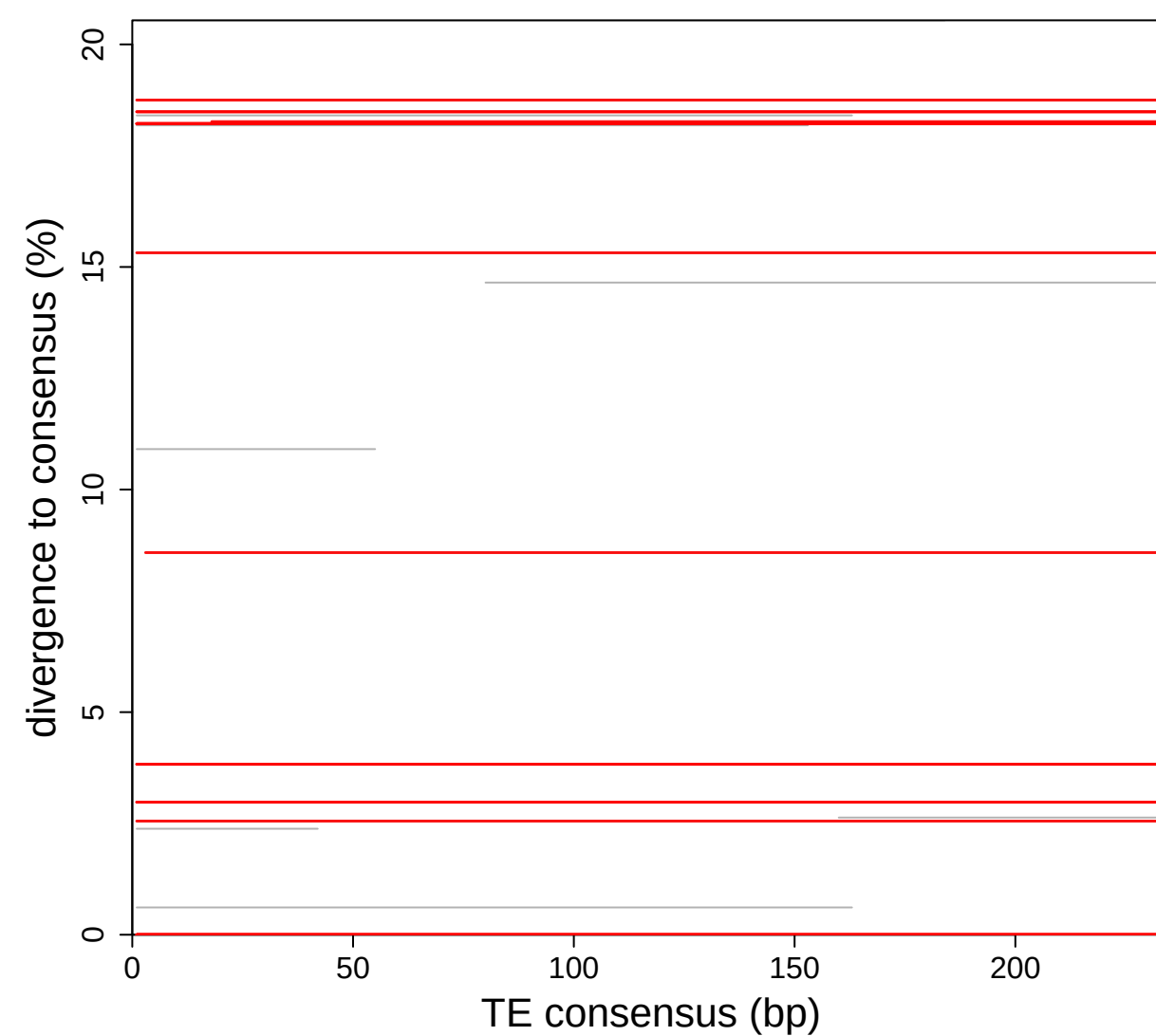


fasta.b.bed uf.bed g_1.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa
size: 13016bp; fragments: 700; full length: 0 (>=11714.4bp)

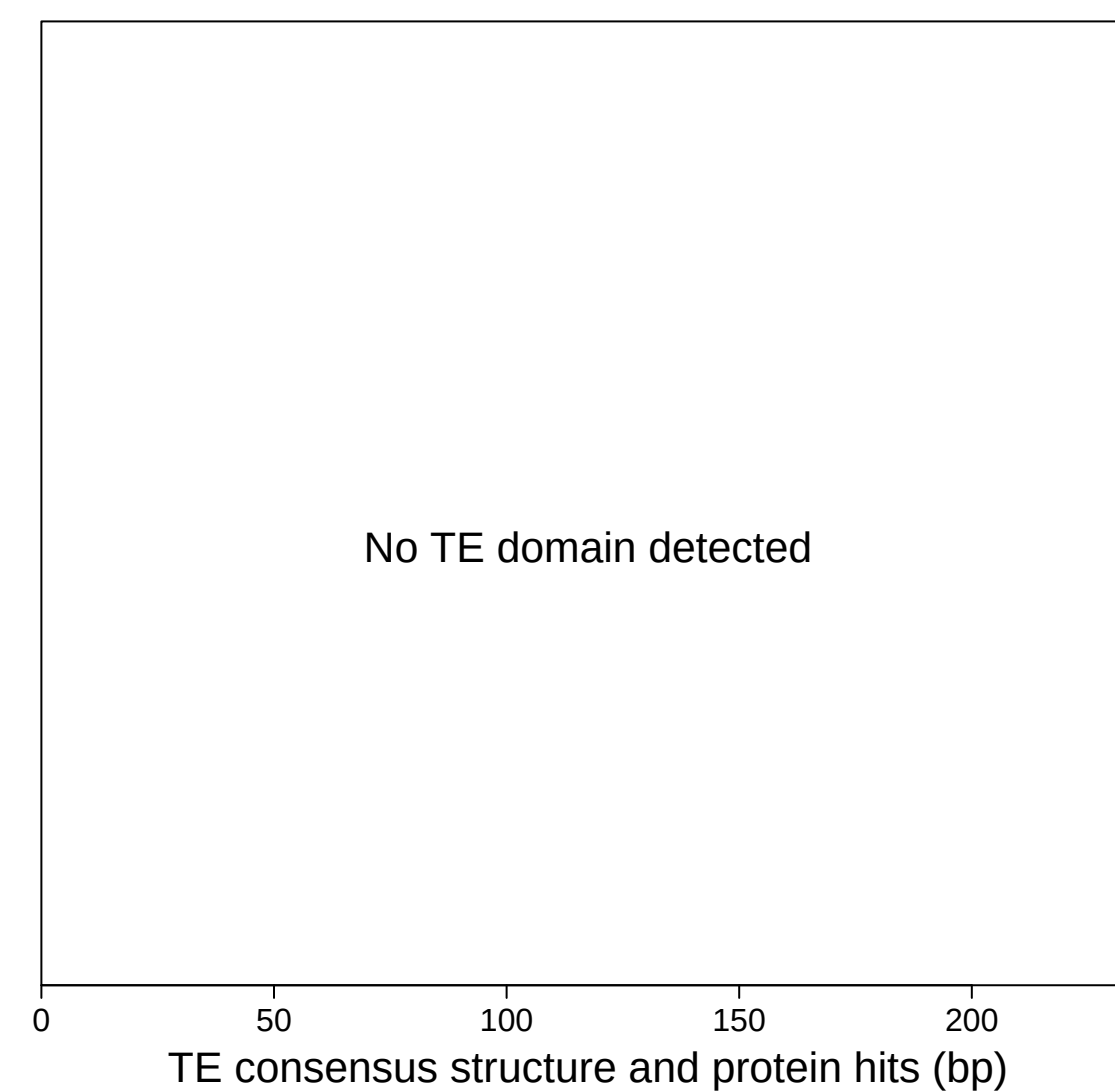
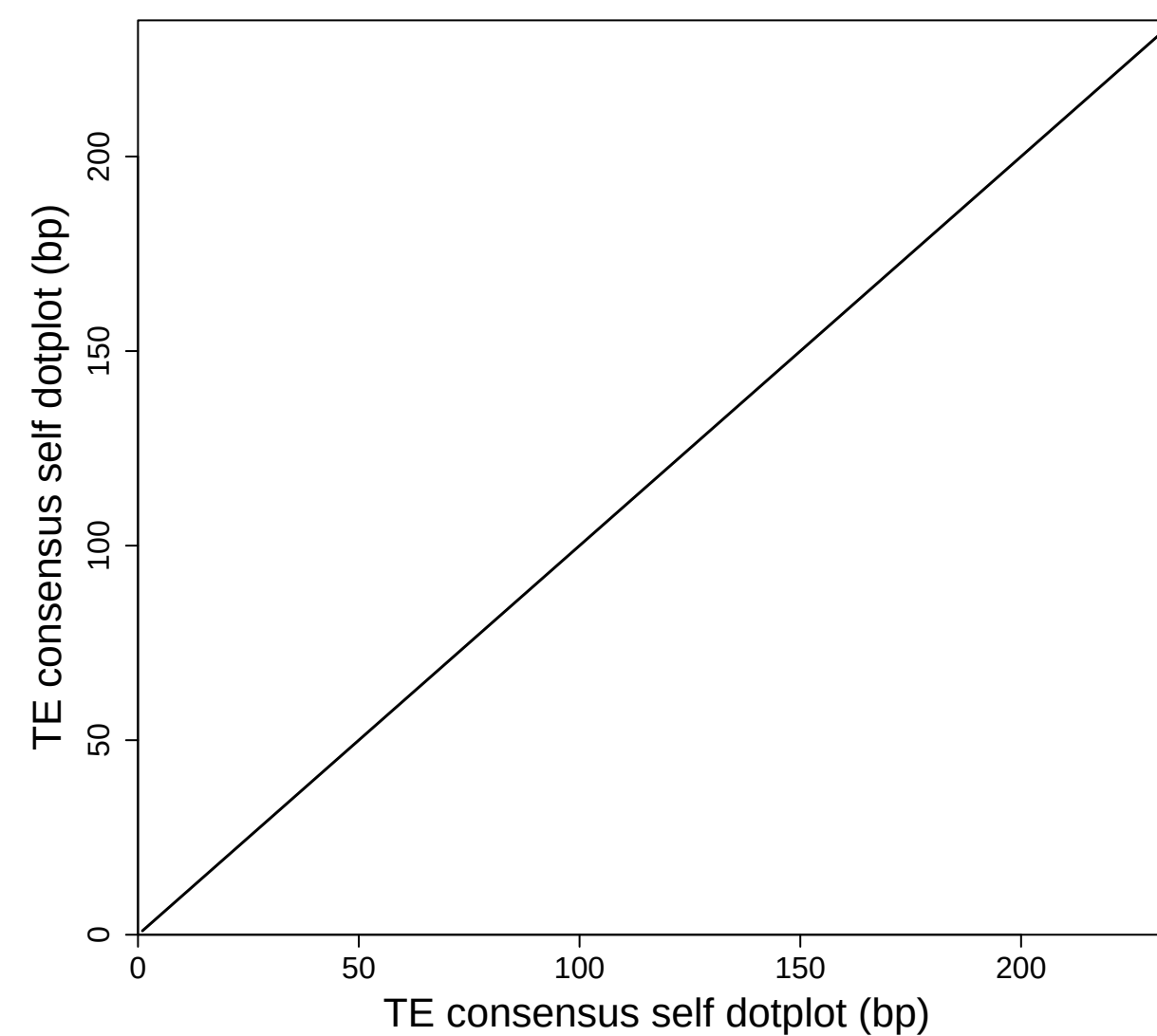
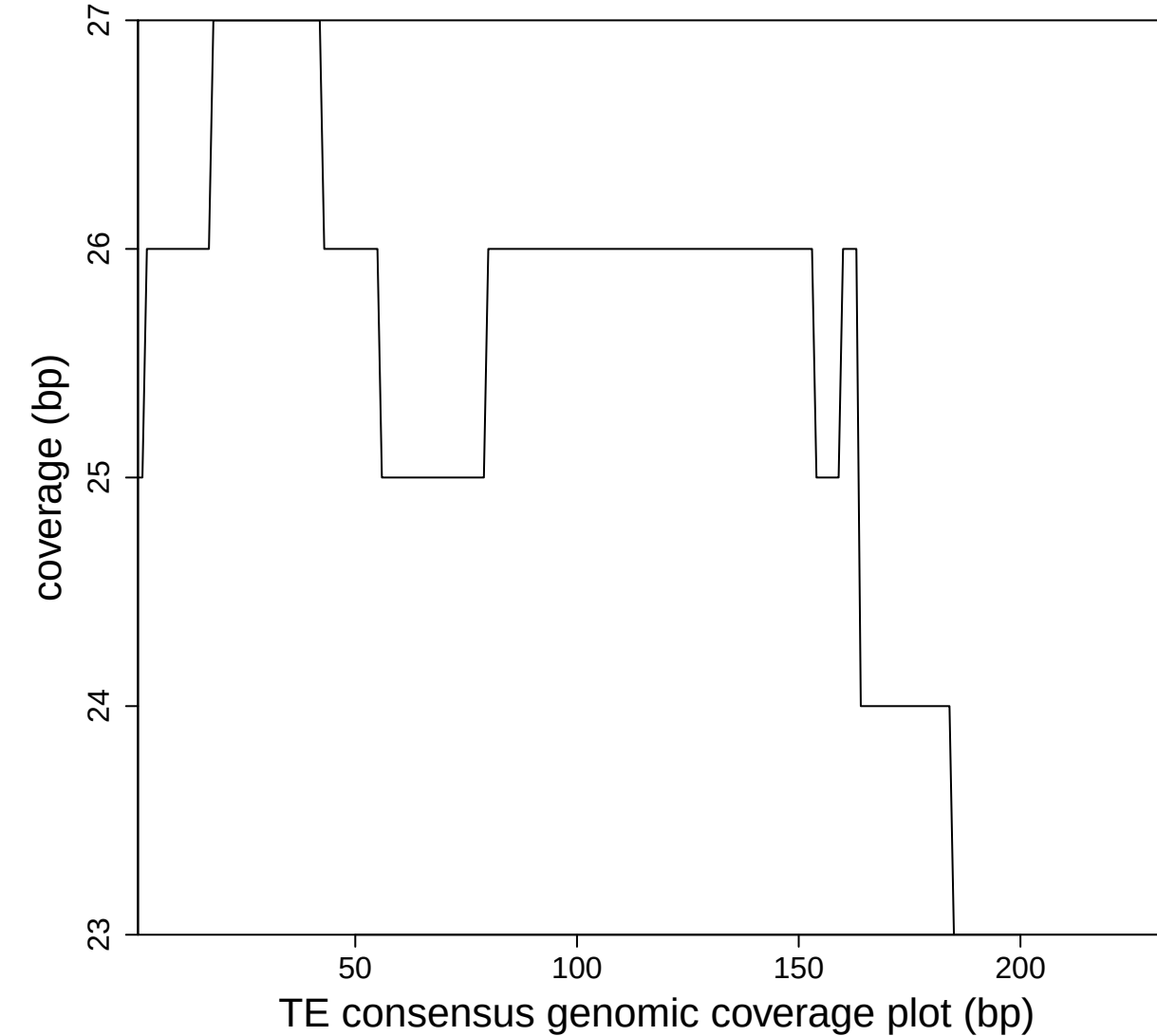
After TEtrimmer Extended plot Blue lines are boundaries



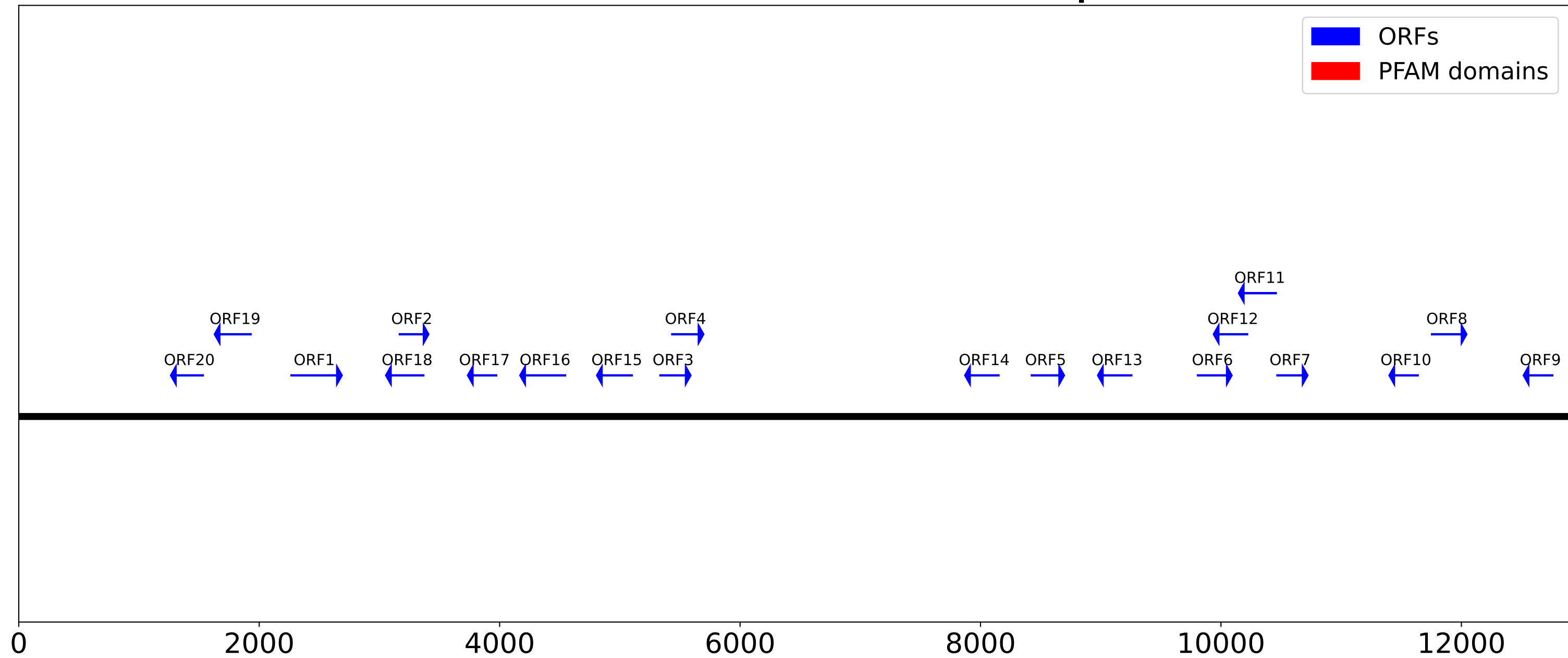
TE: ltr_1_family_75
size: 235bp; fragments: 29; full length: 21 (≥ 211.5 bp)



Before TEtrimmer 235 bp



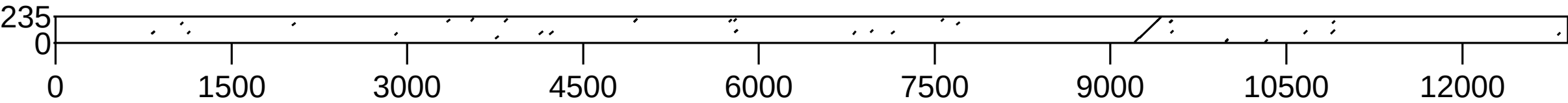
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)