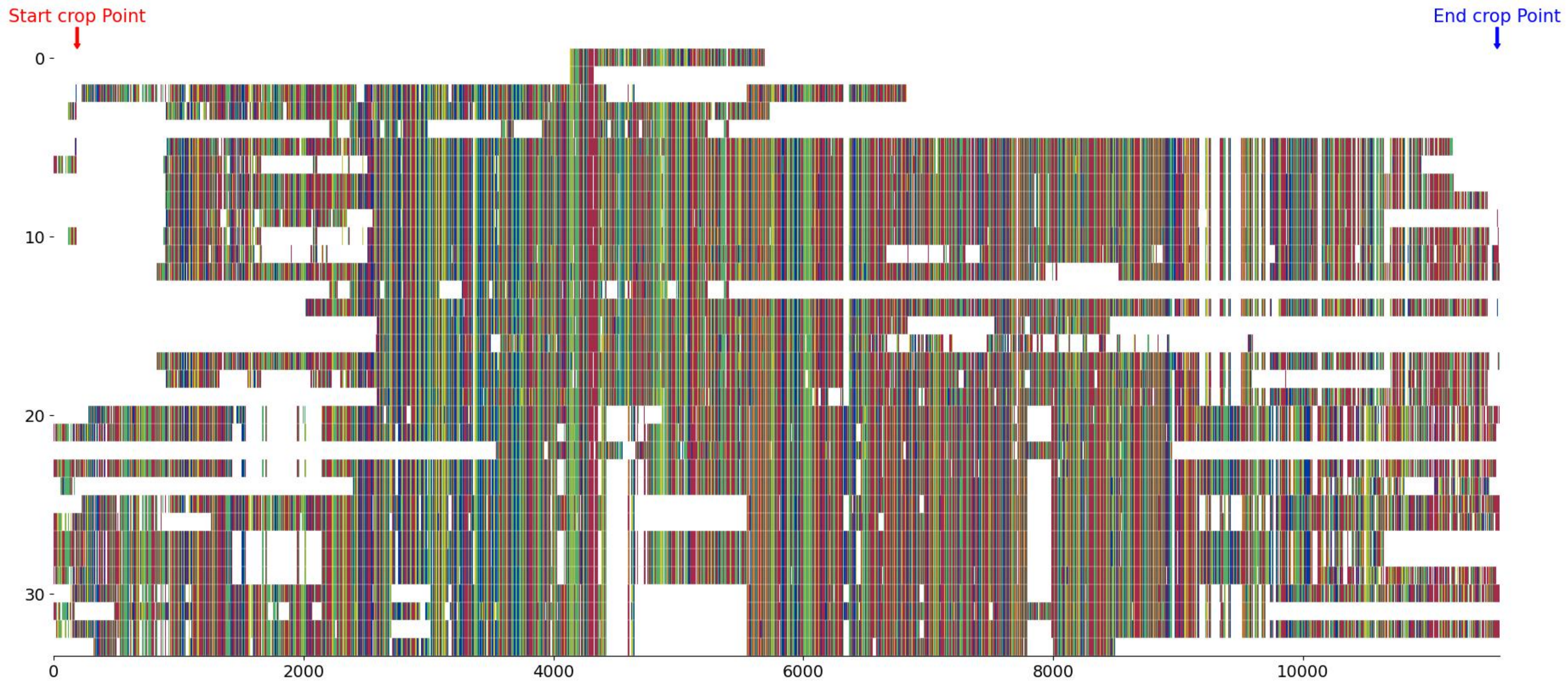


Start crop Point

End crop Point

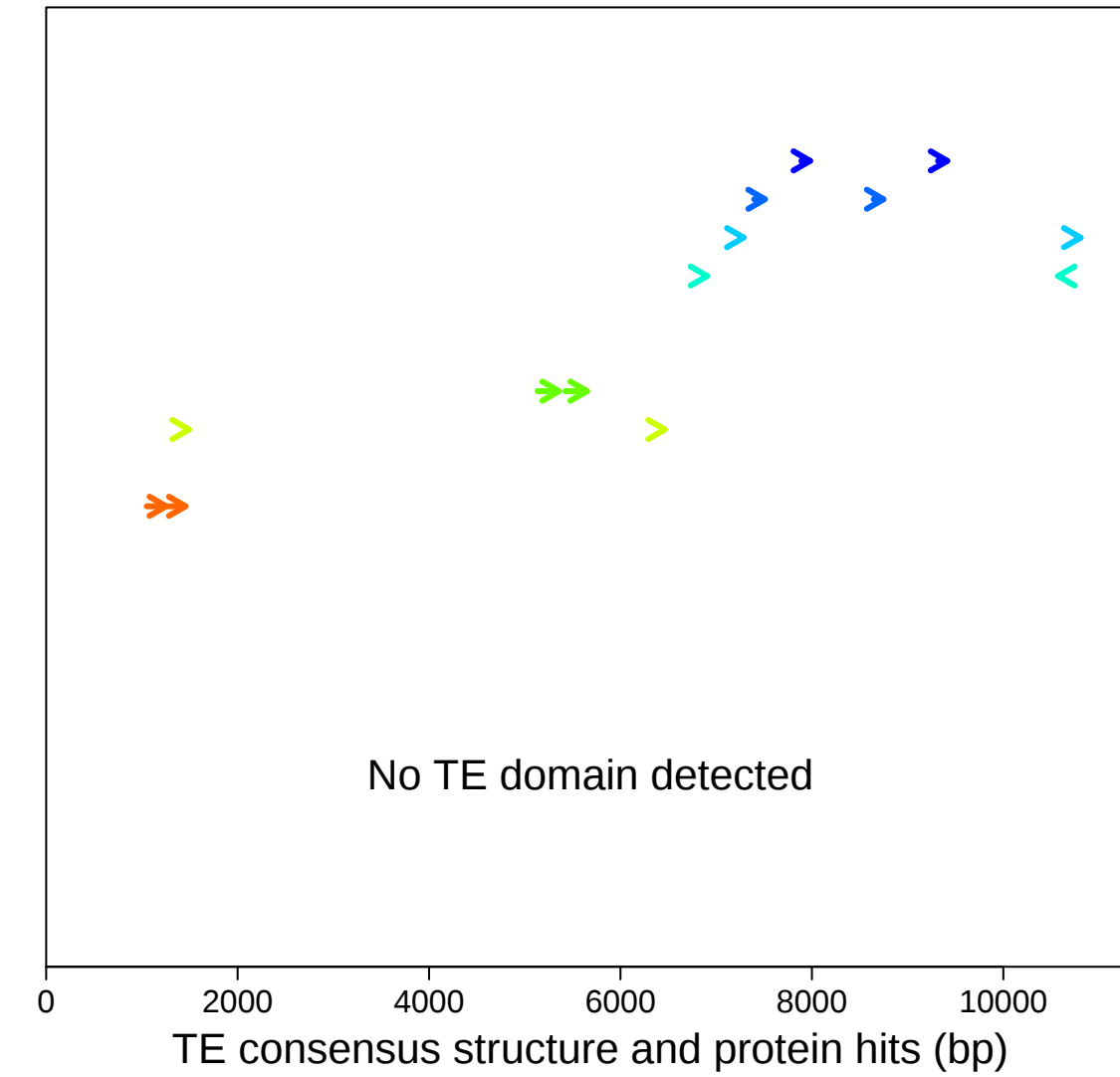
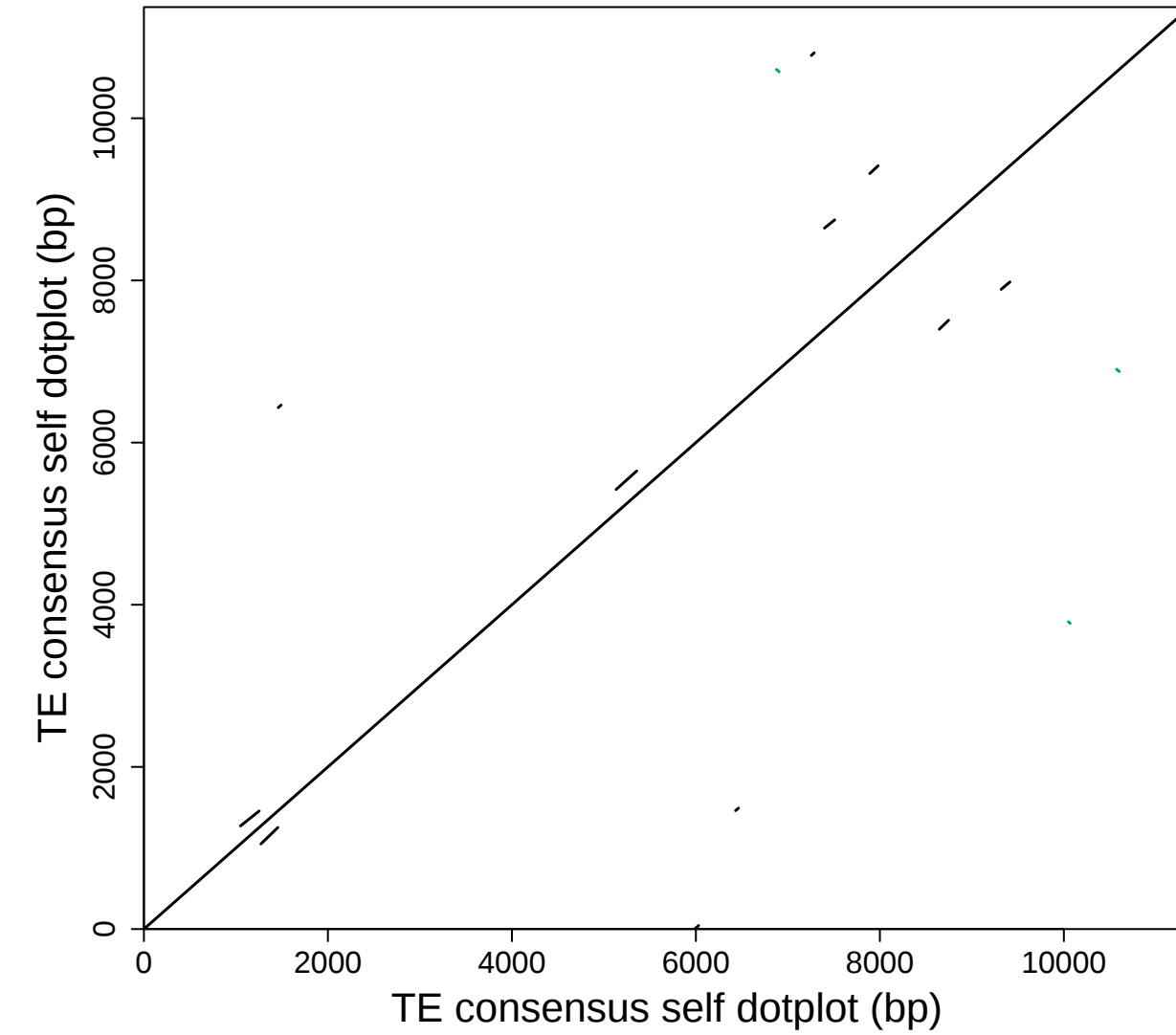
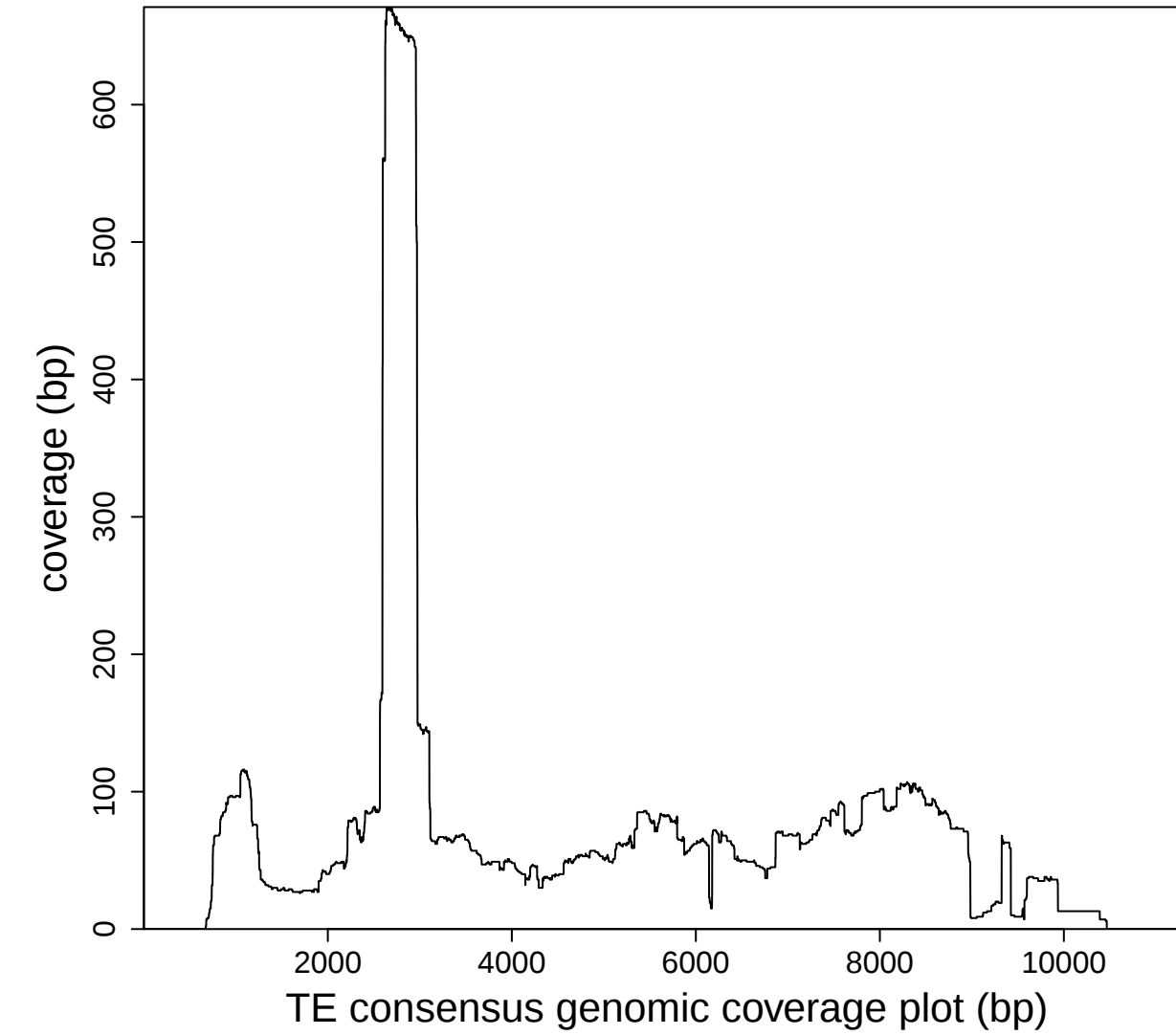
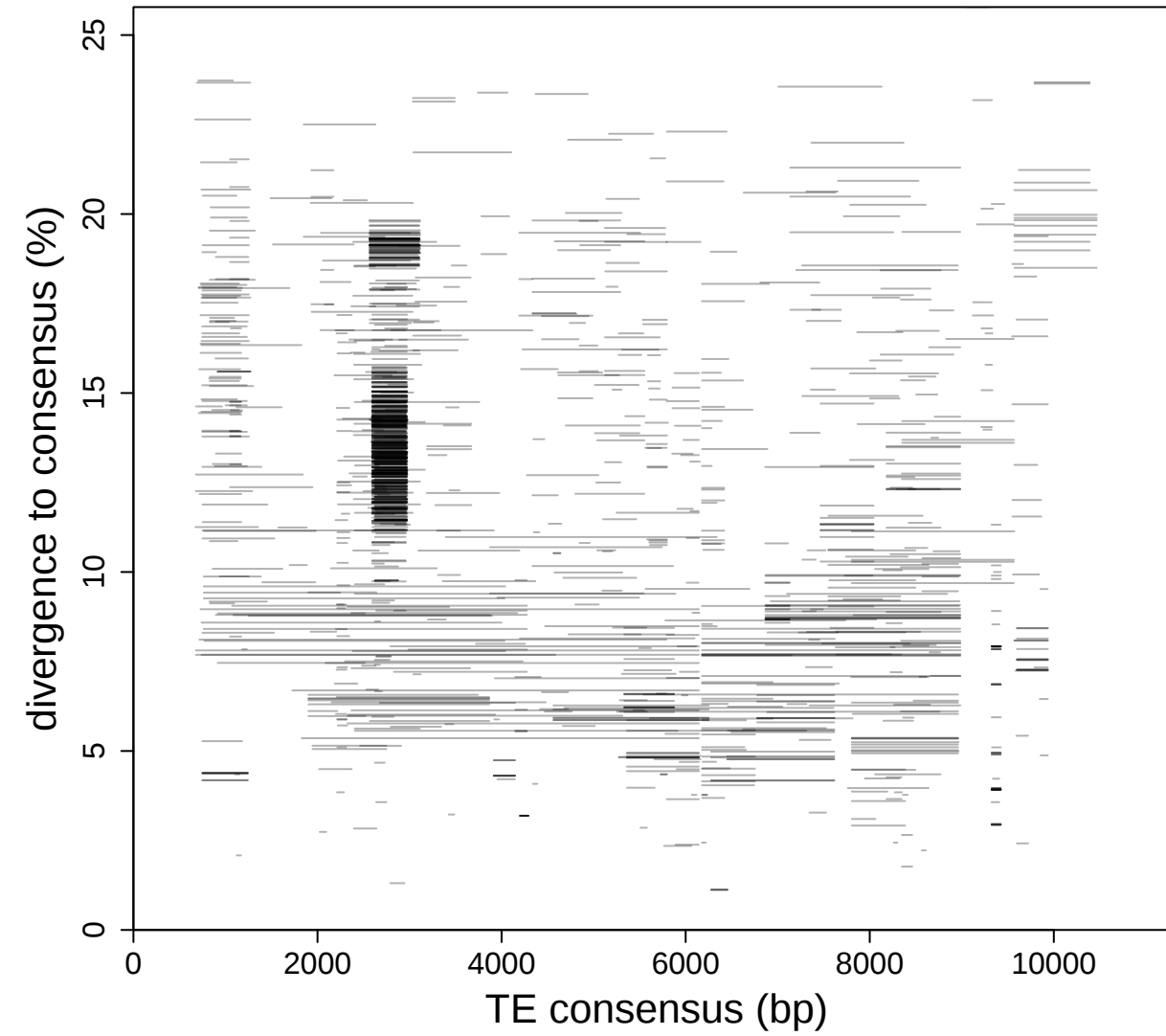
MSA length = 11371





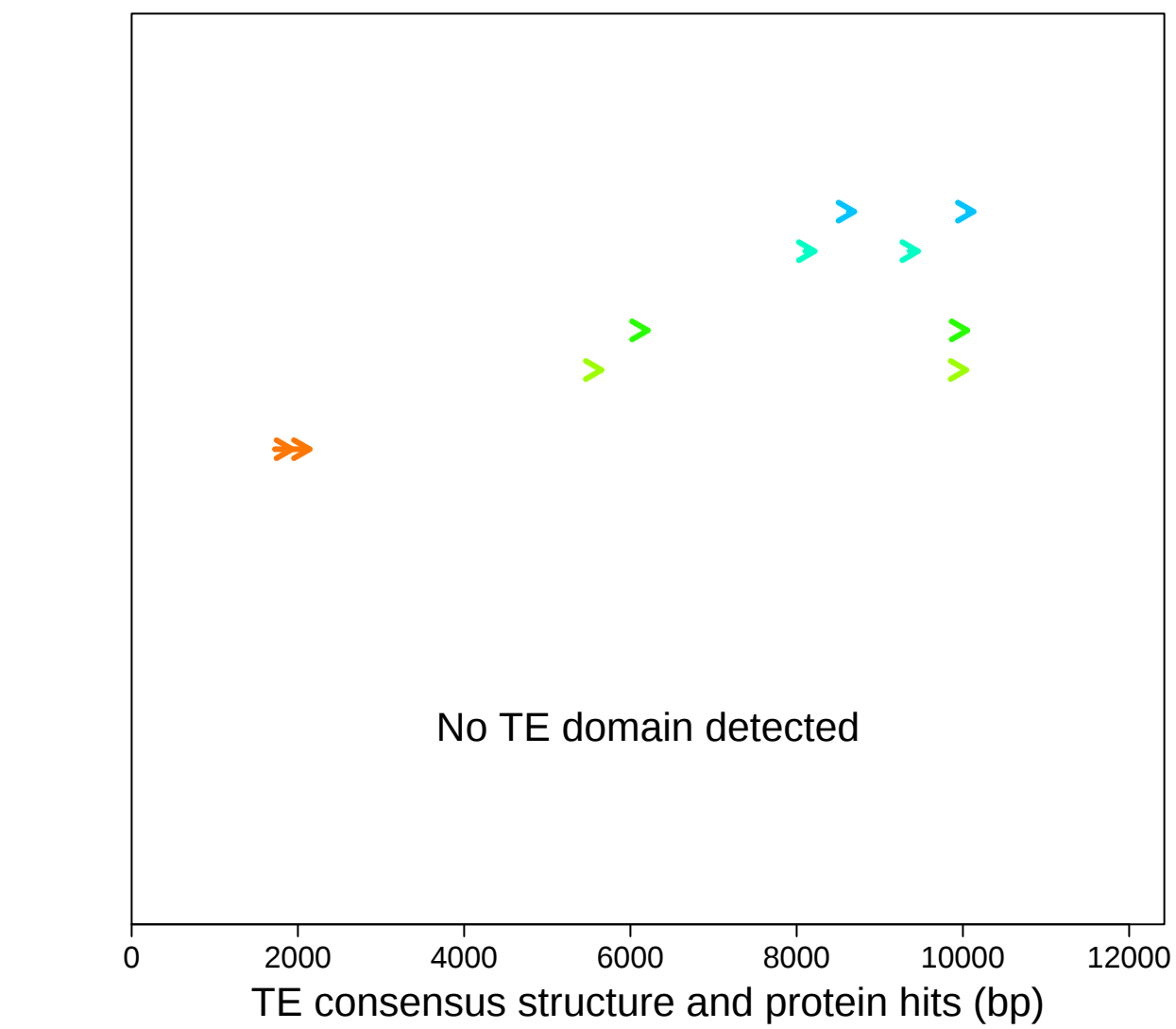
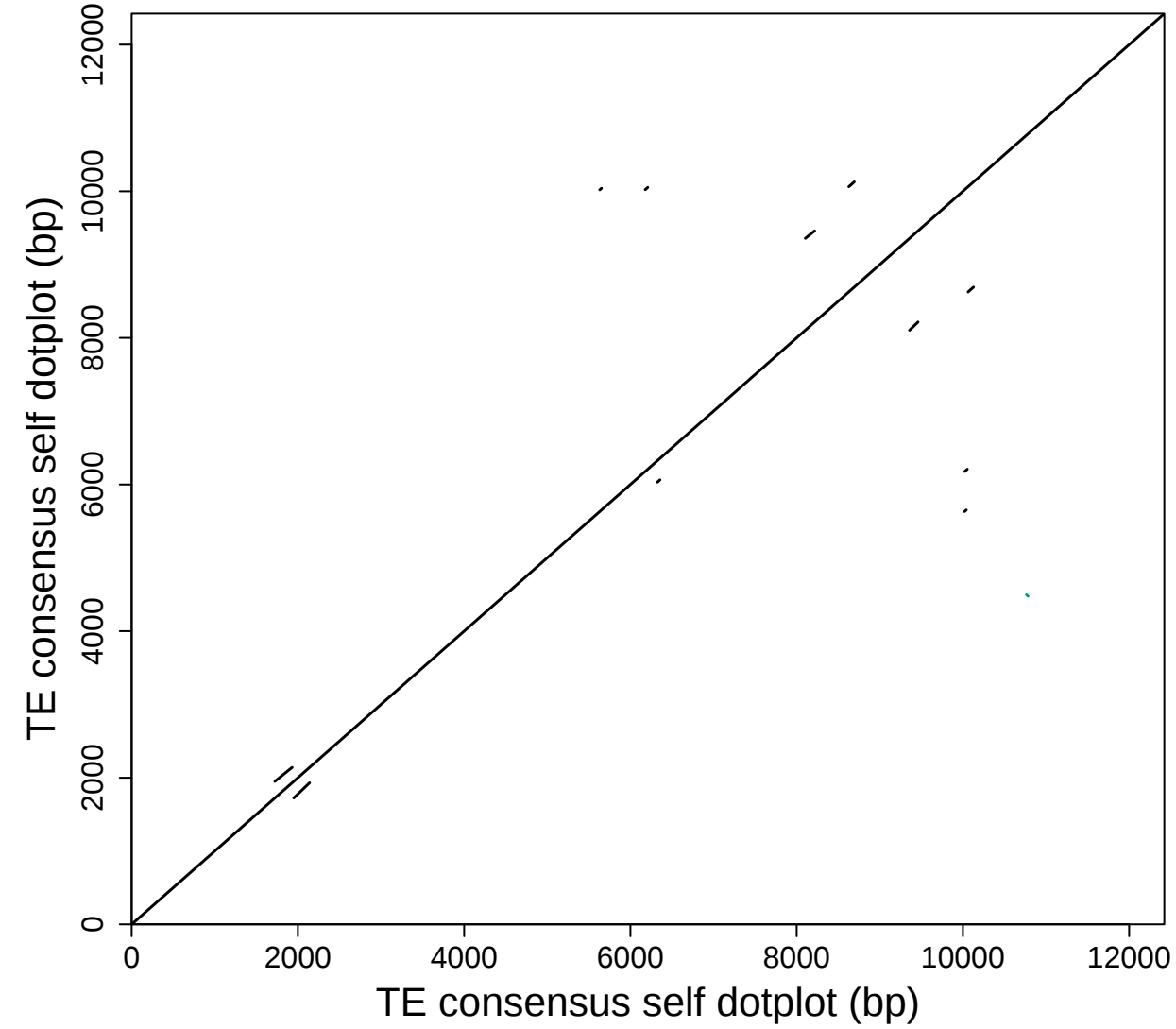
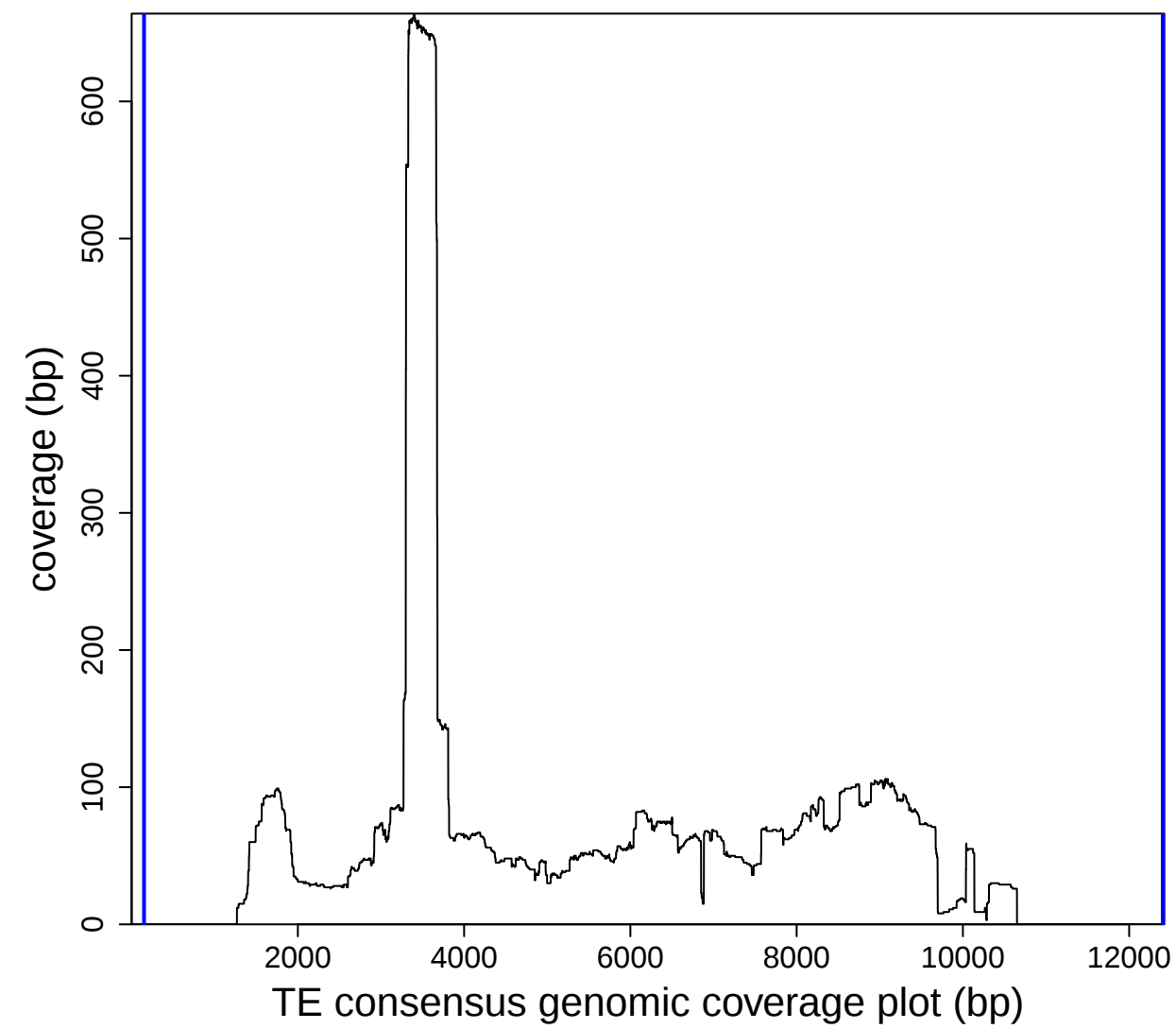
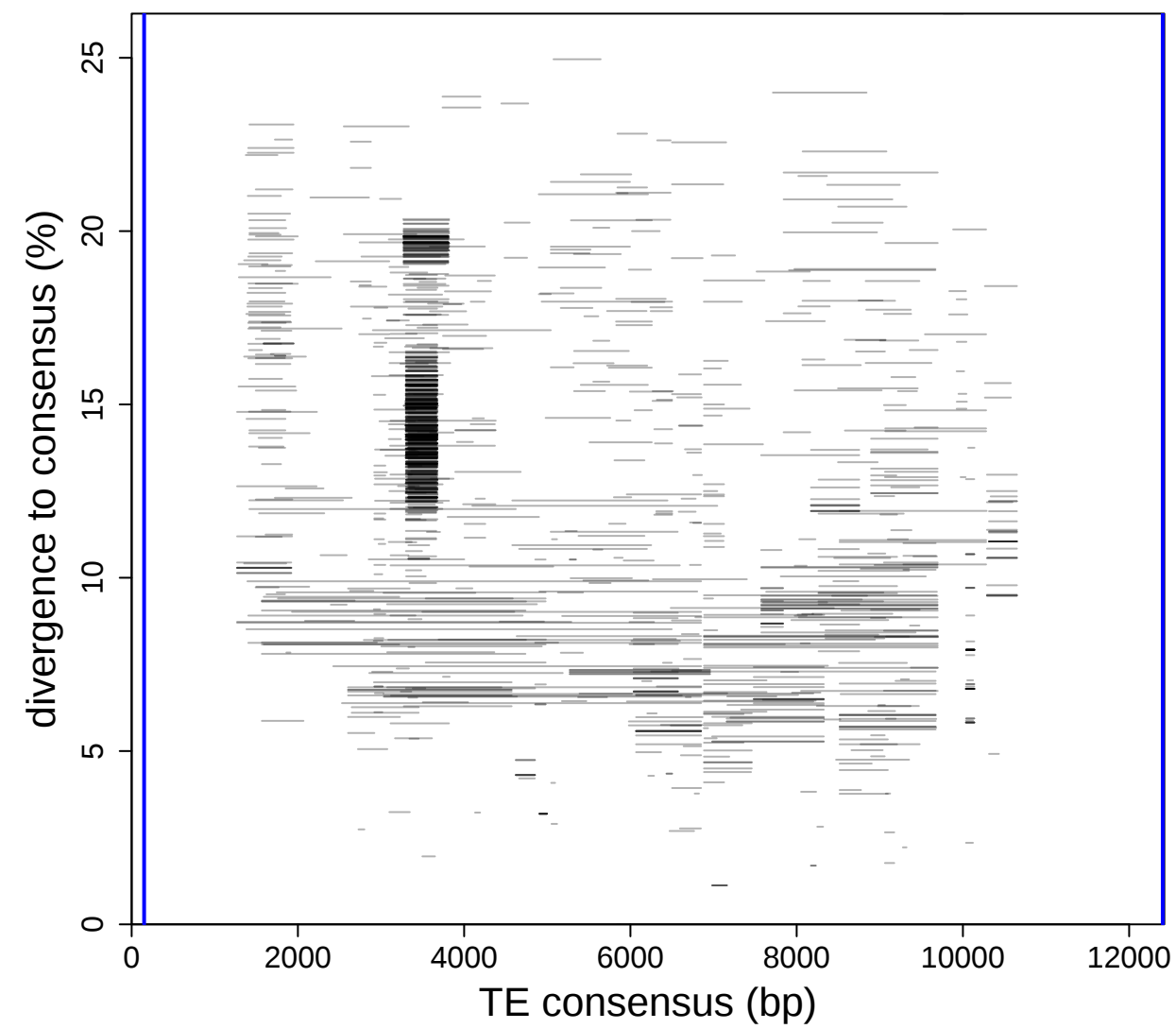
b.bed uf.bed g_1.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa
size: 11371bp; fragments: 1474; full length: 0 (>=10233.9bp)

After TEtrimmer 11371 bp

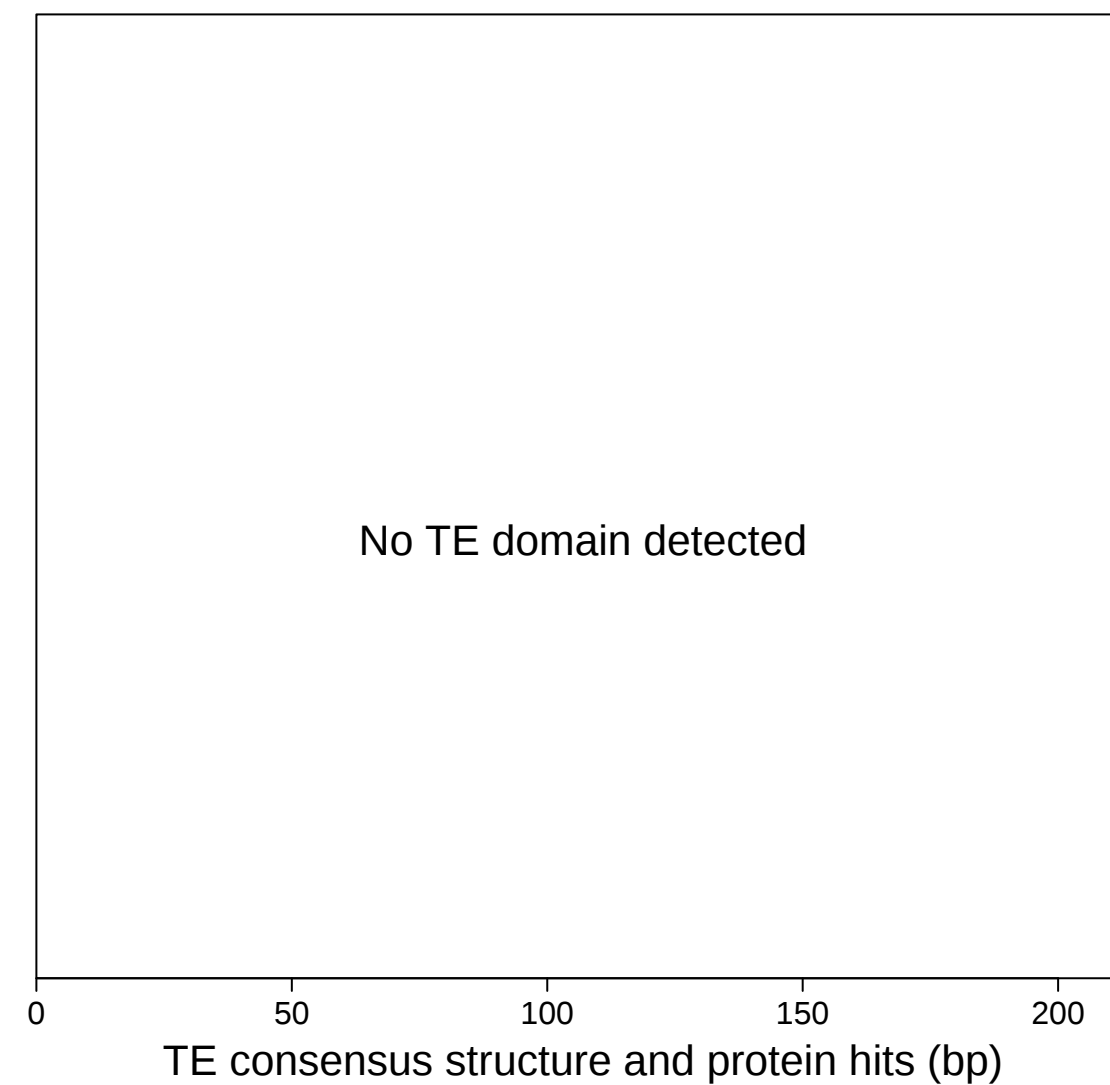
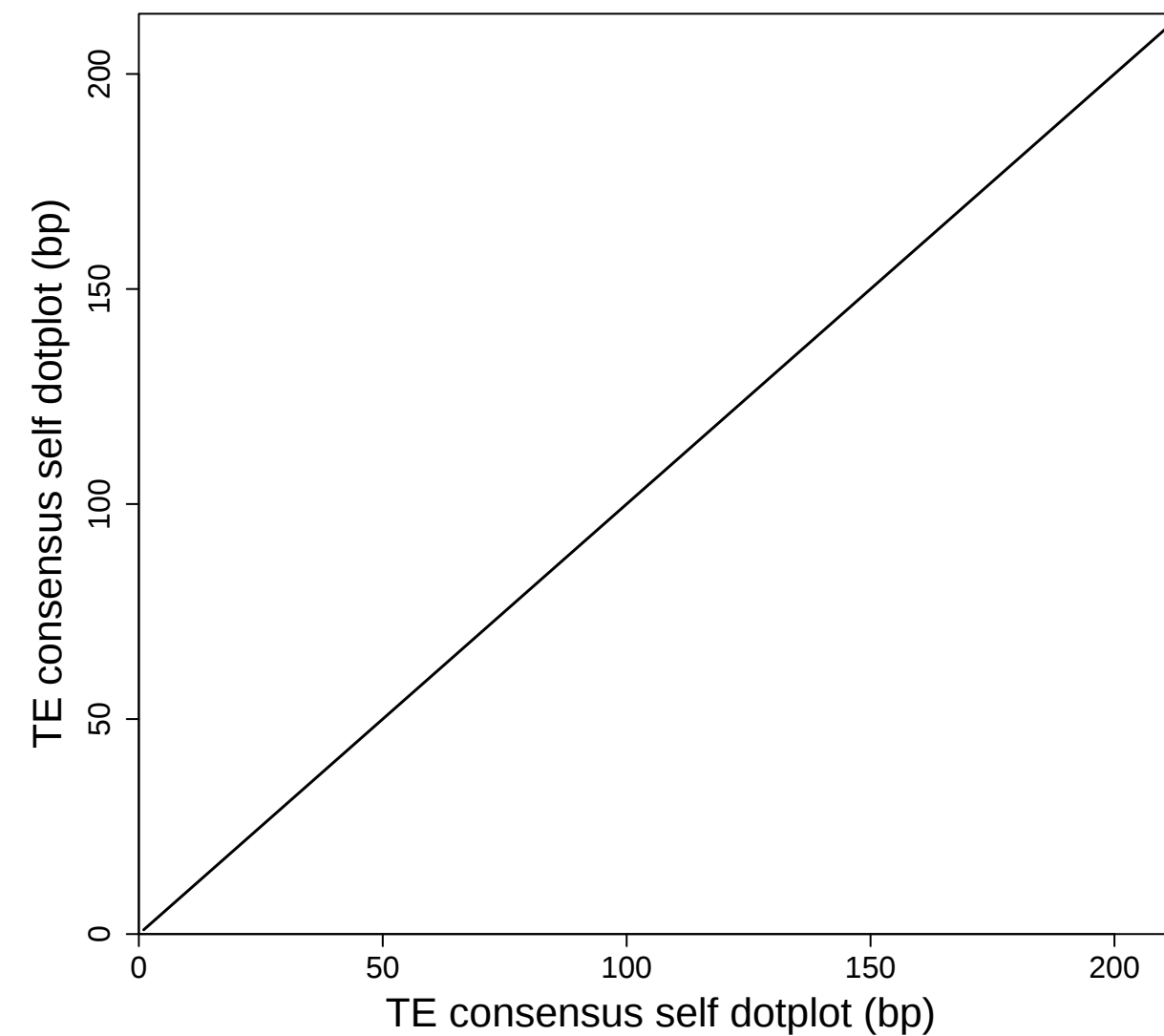
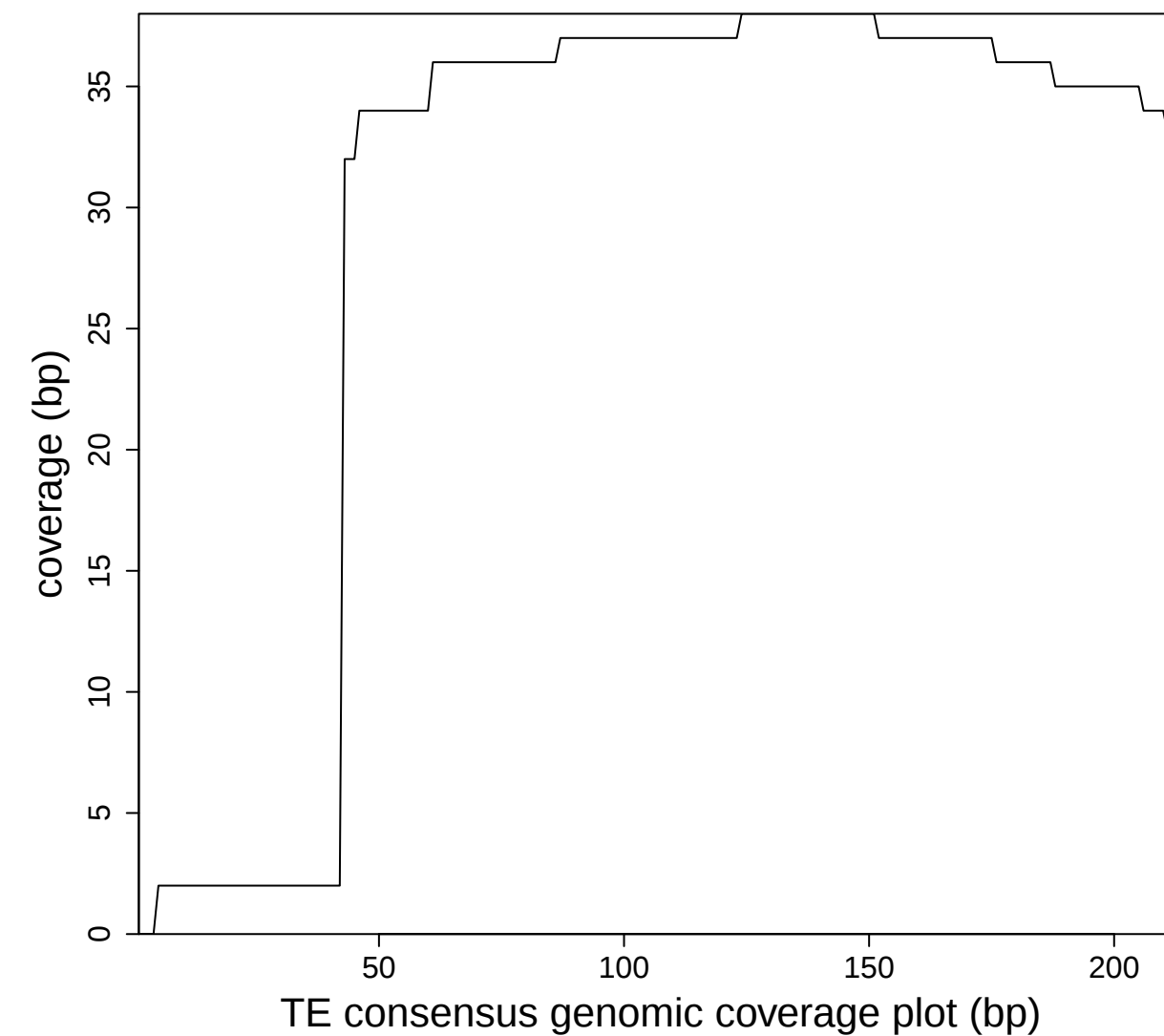
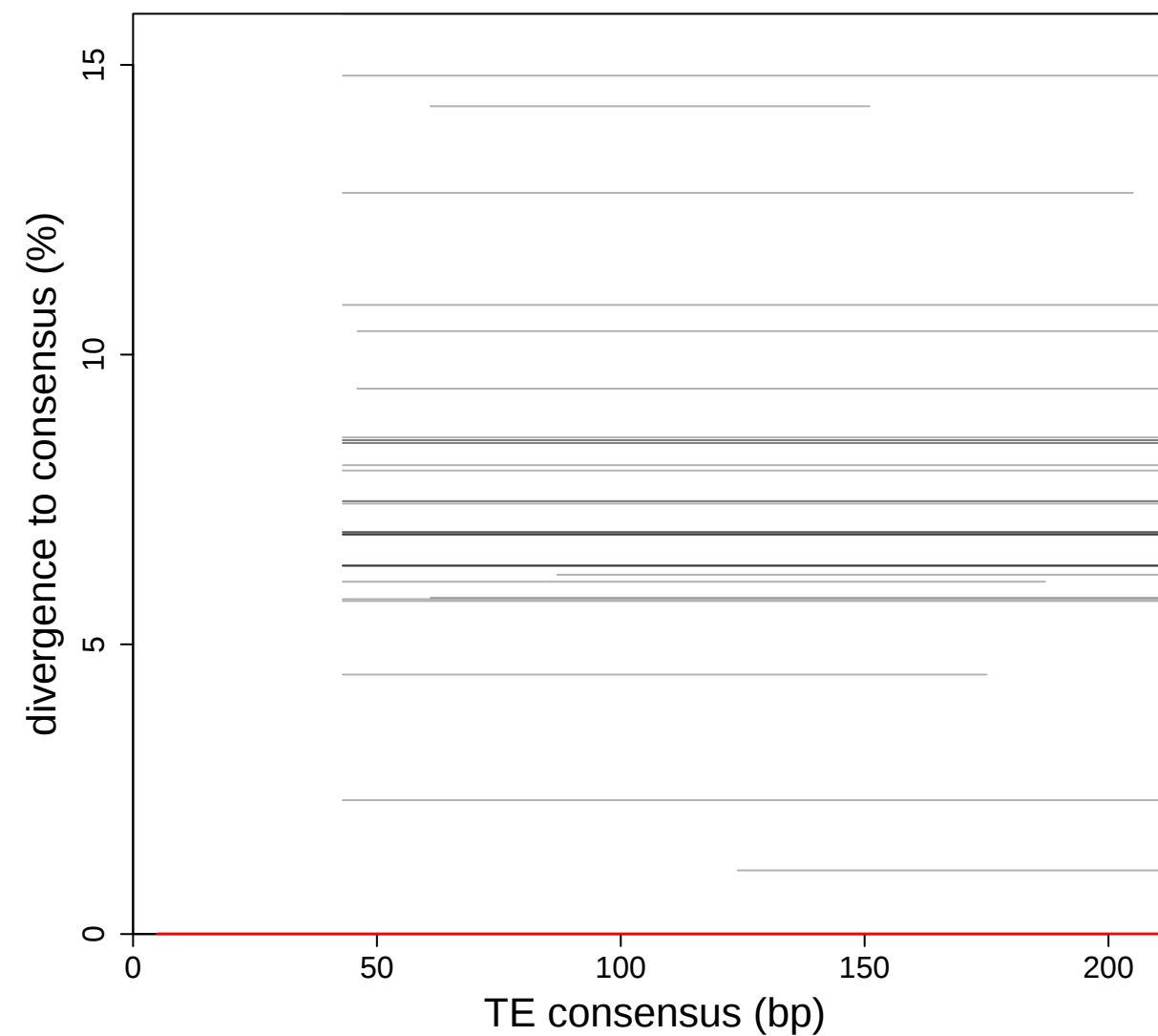


3.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 12423bp; fragments: 1393; full length: 0 (>=11180.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

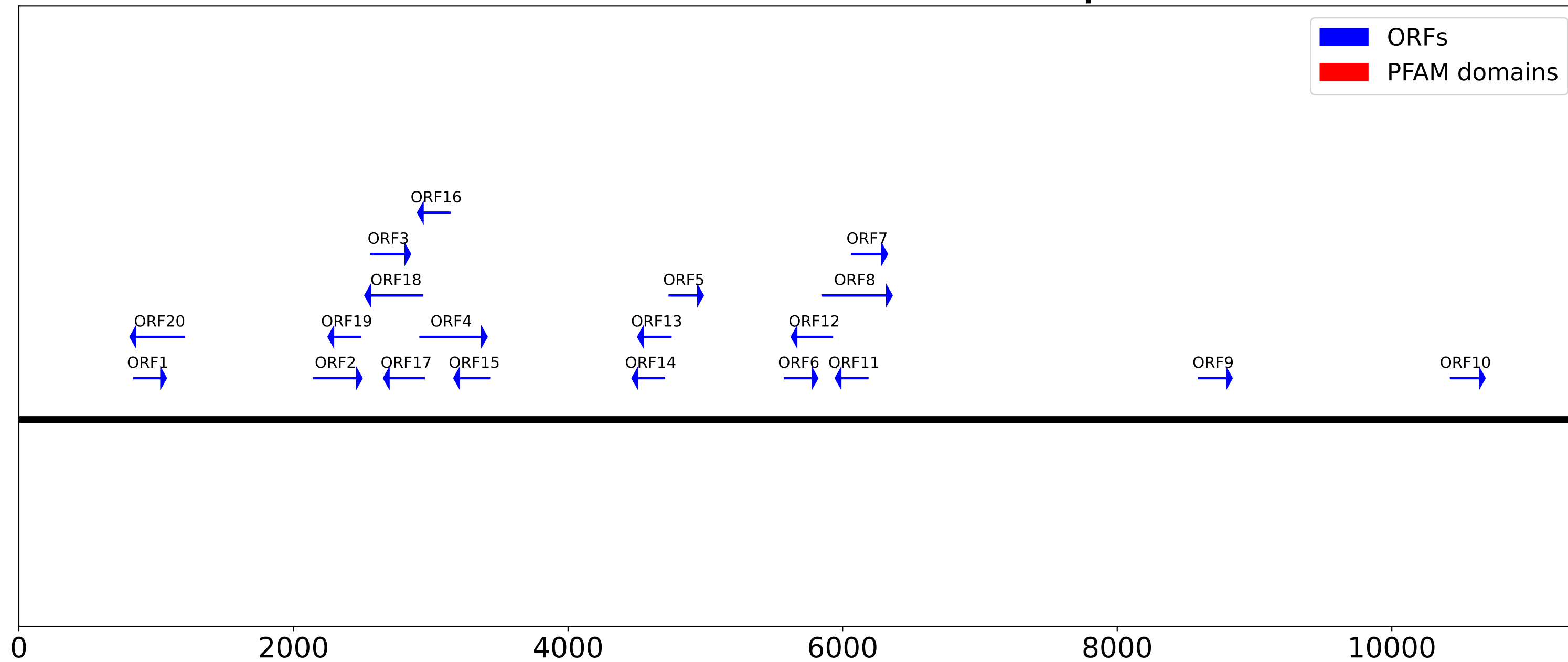


TE: ltr_1_family_363
size: 214bp; fragments: 38; full length: 2 (≥ 192.6 bp)



Before TEtrimmer 214 bp

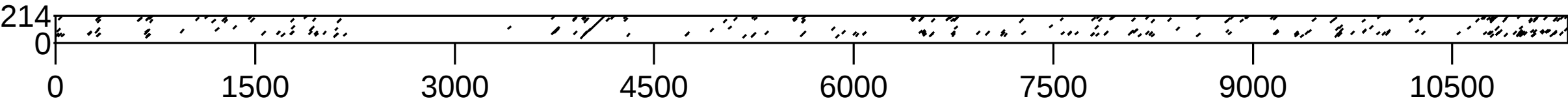
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

Consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)