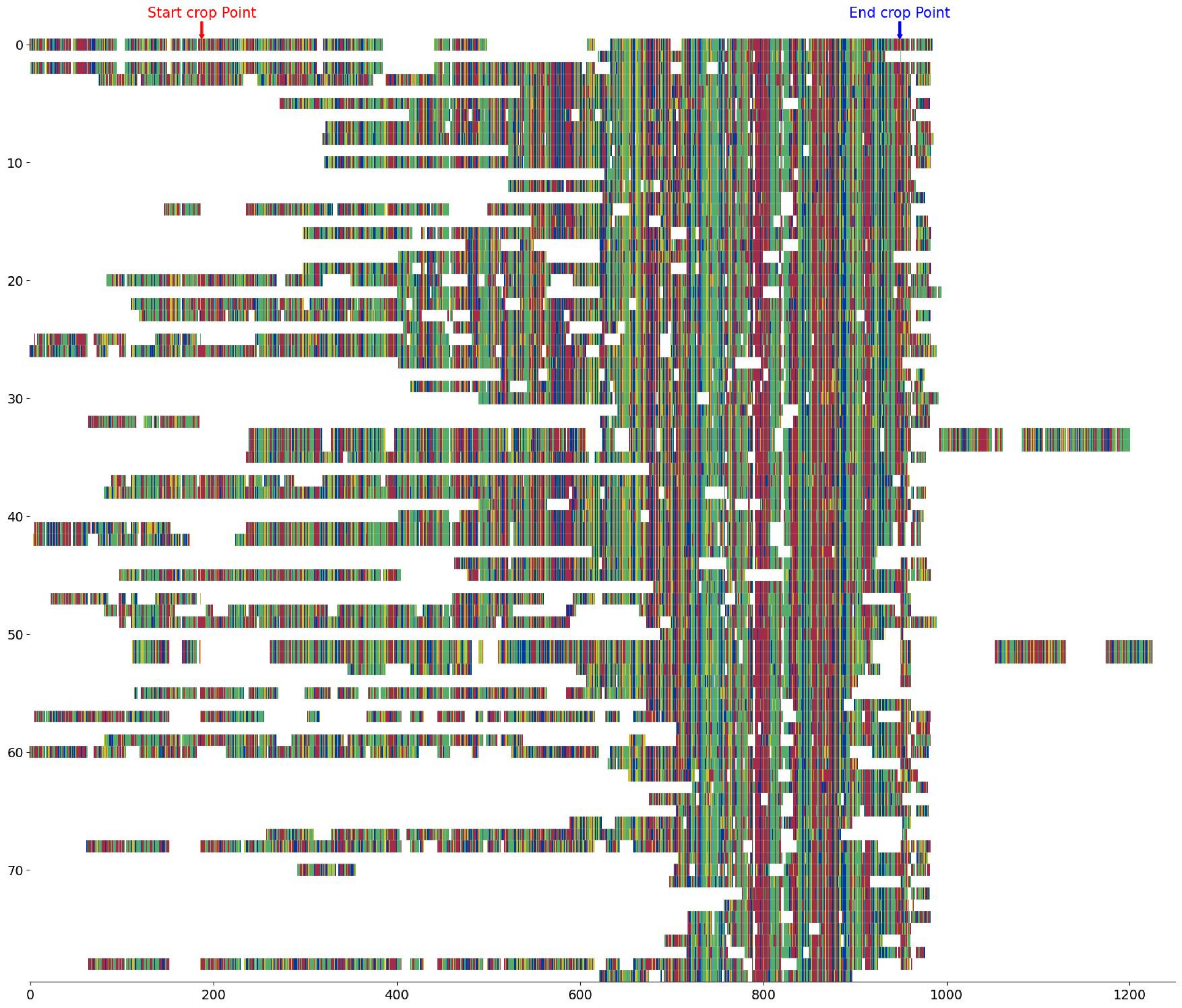
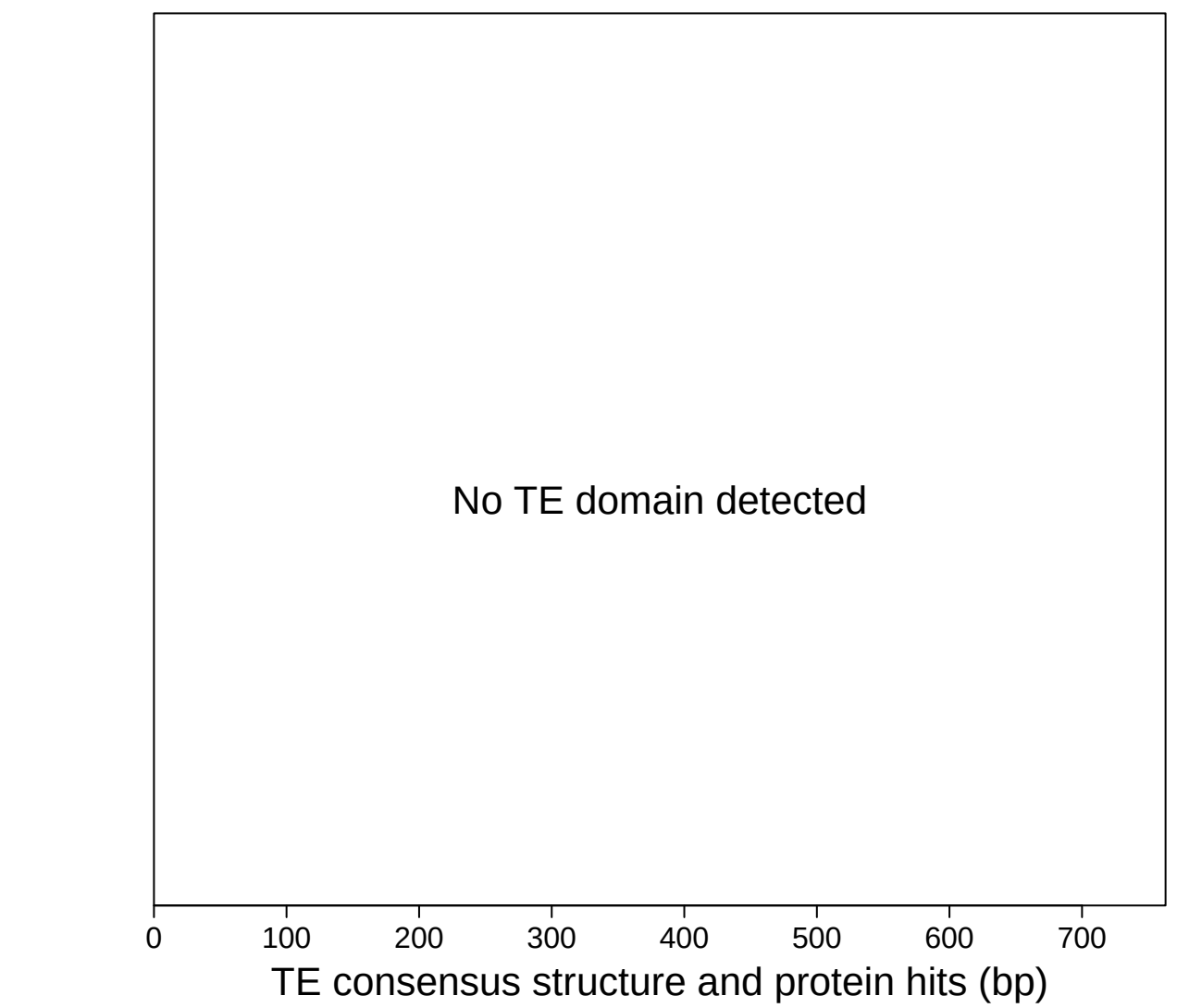
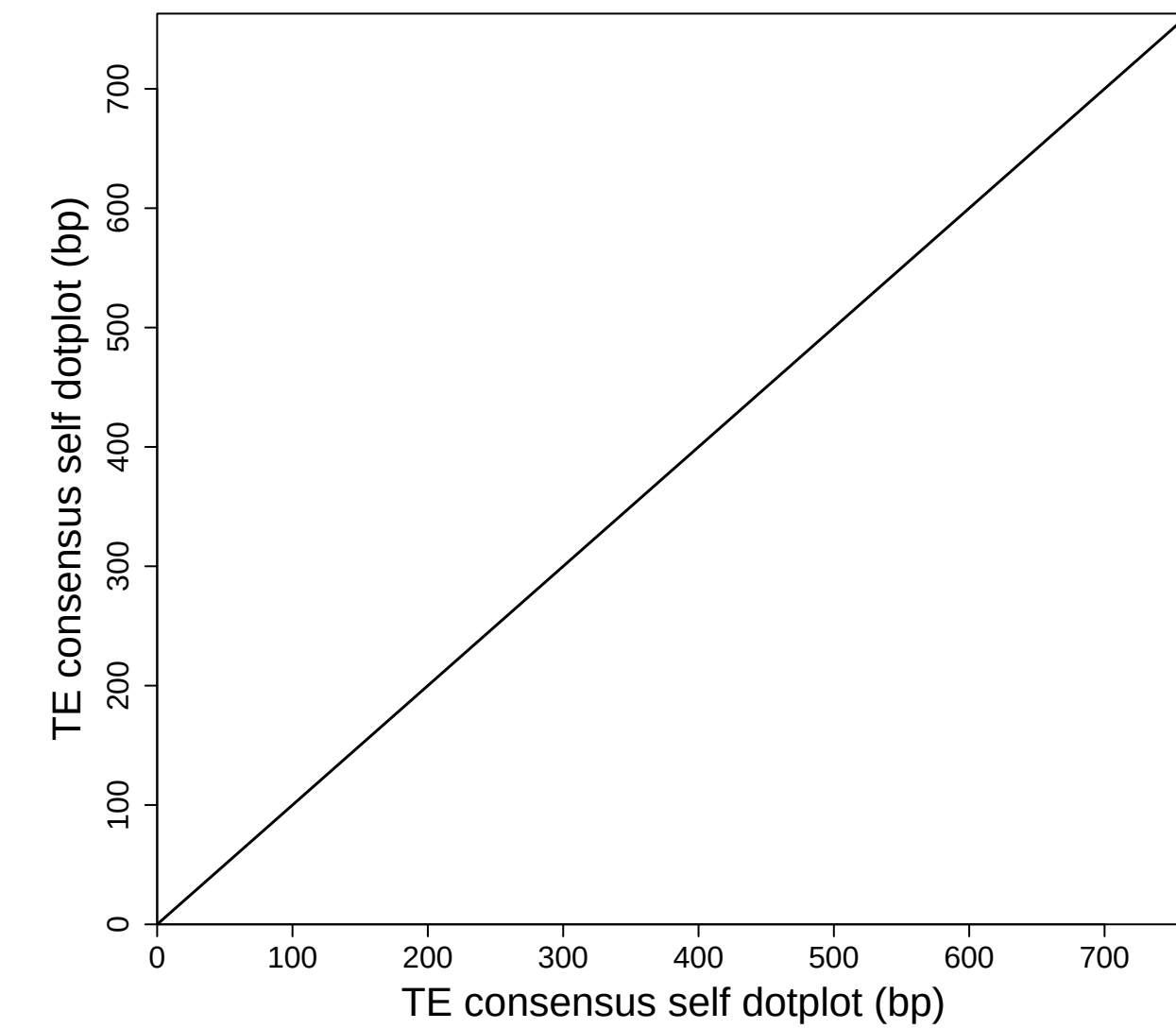
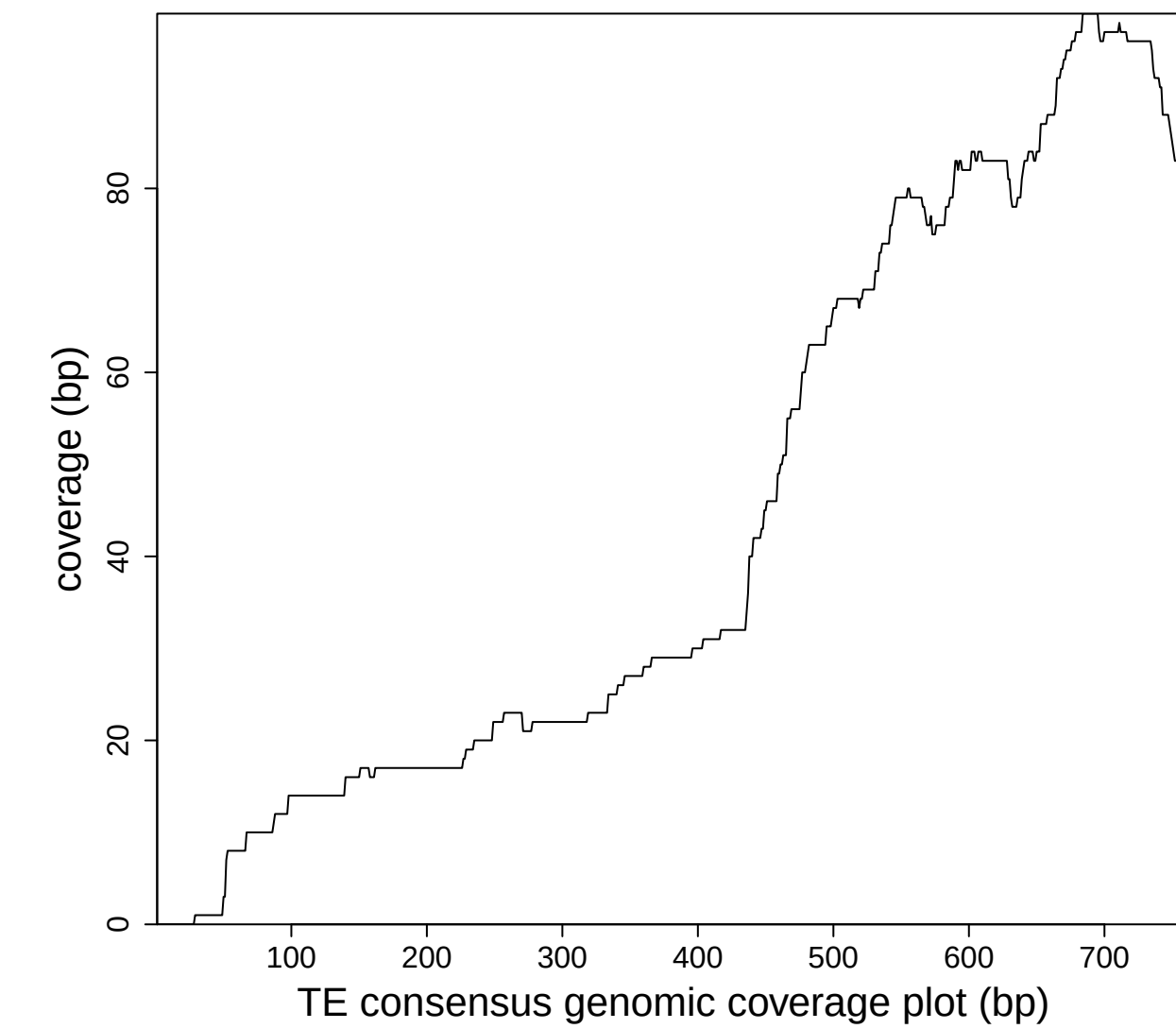
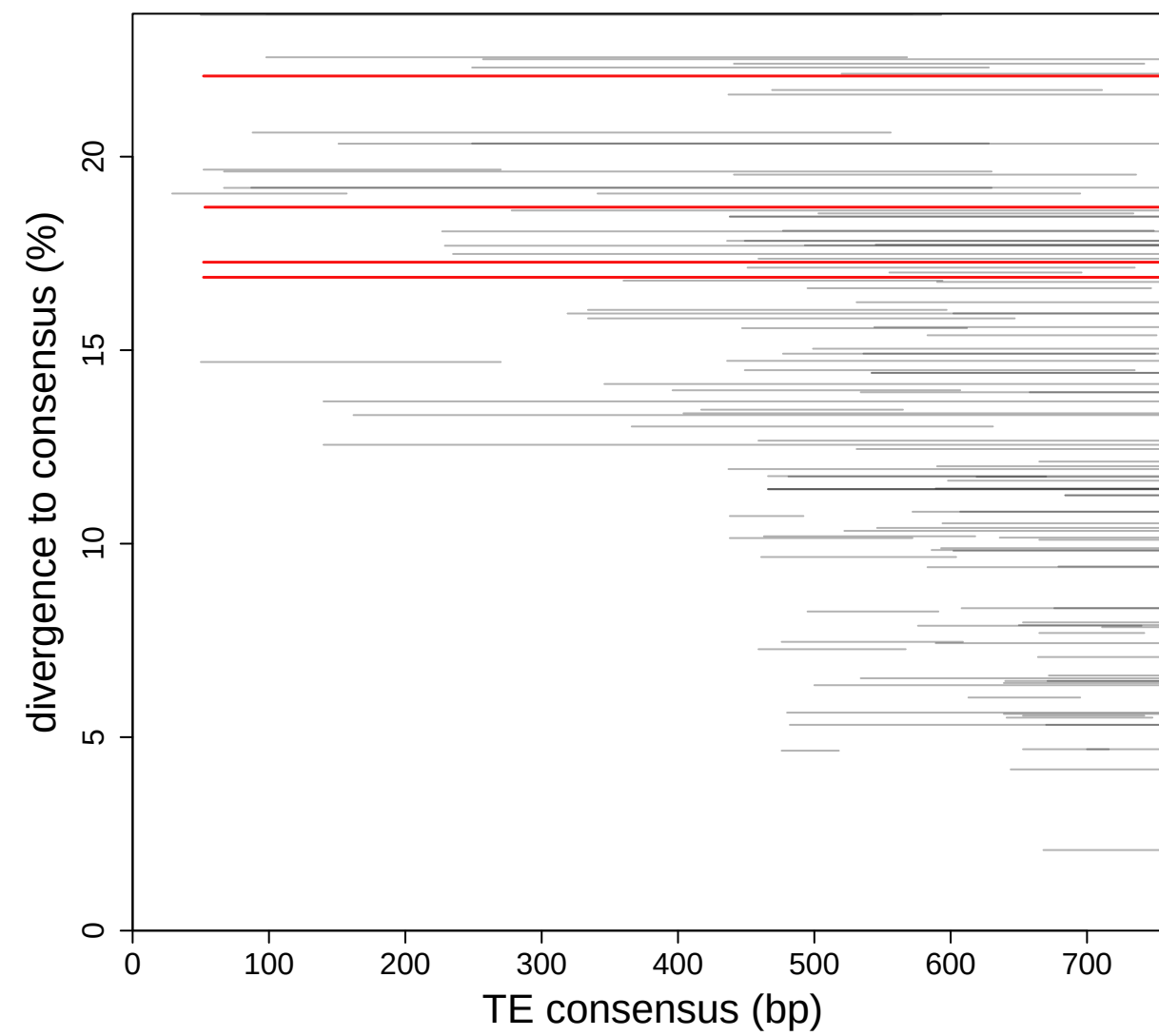






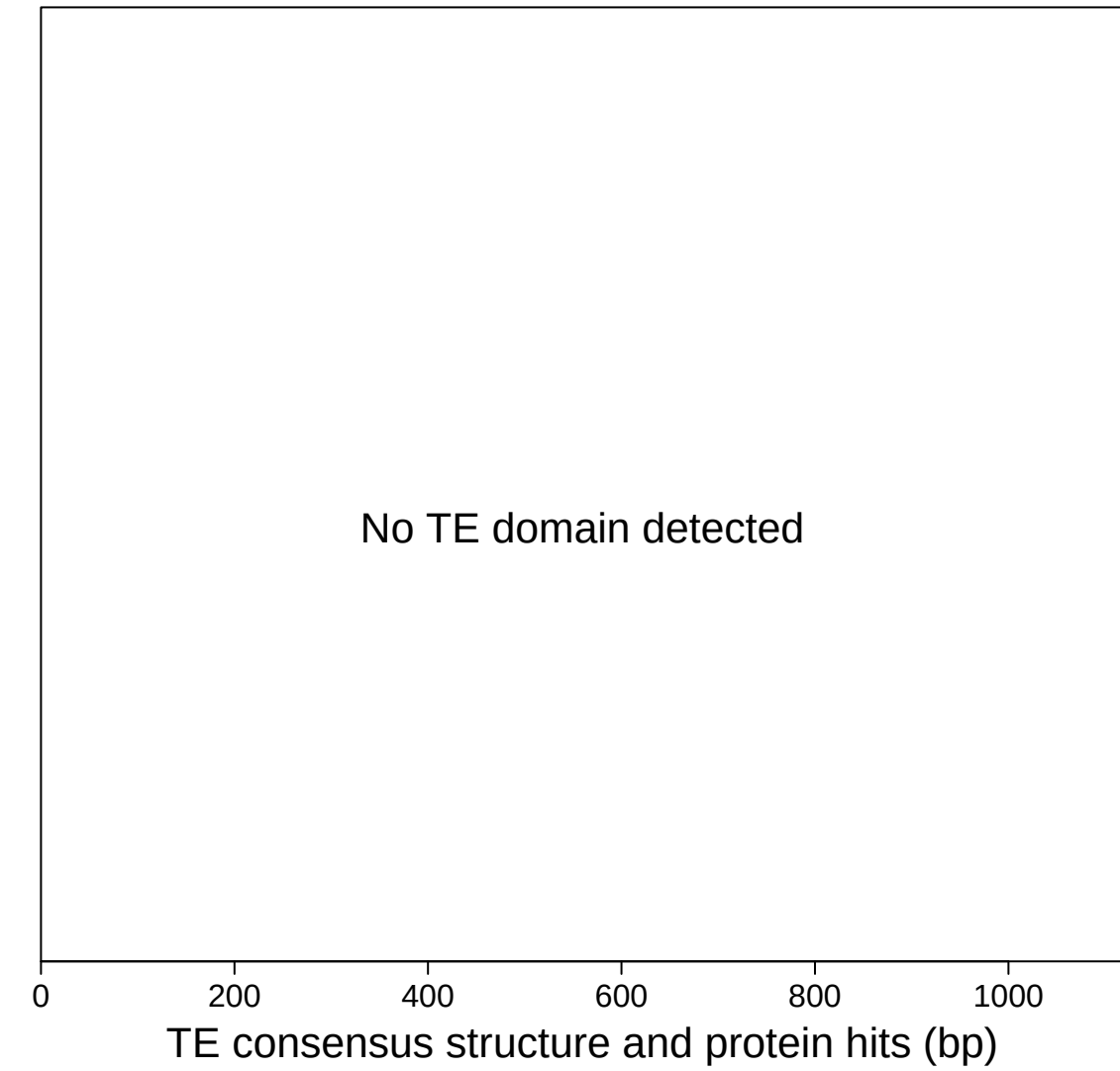
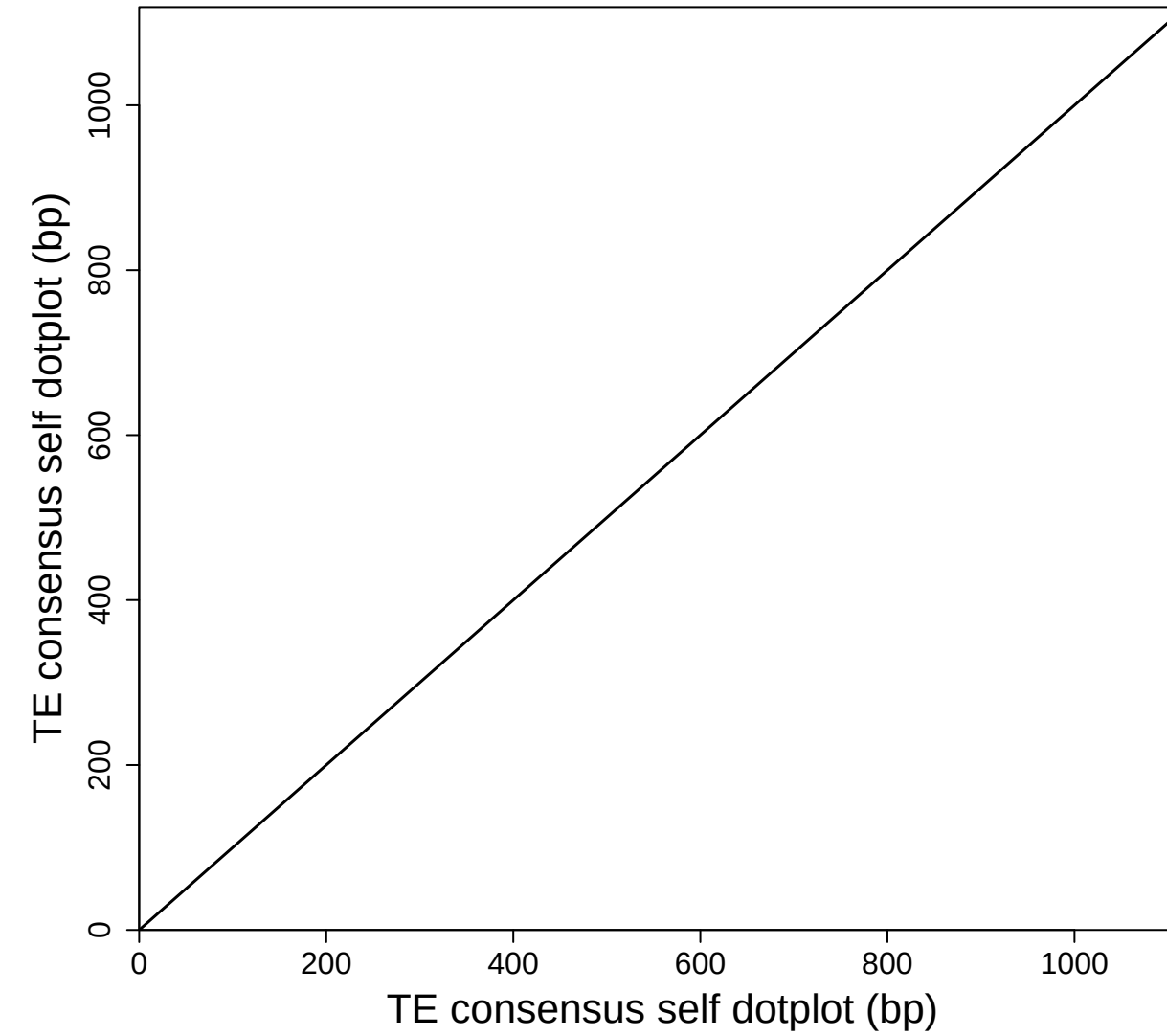
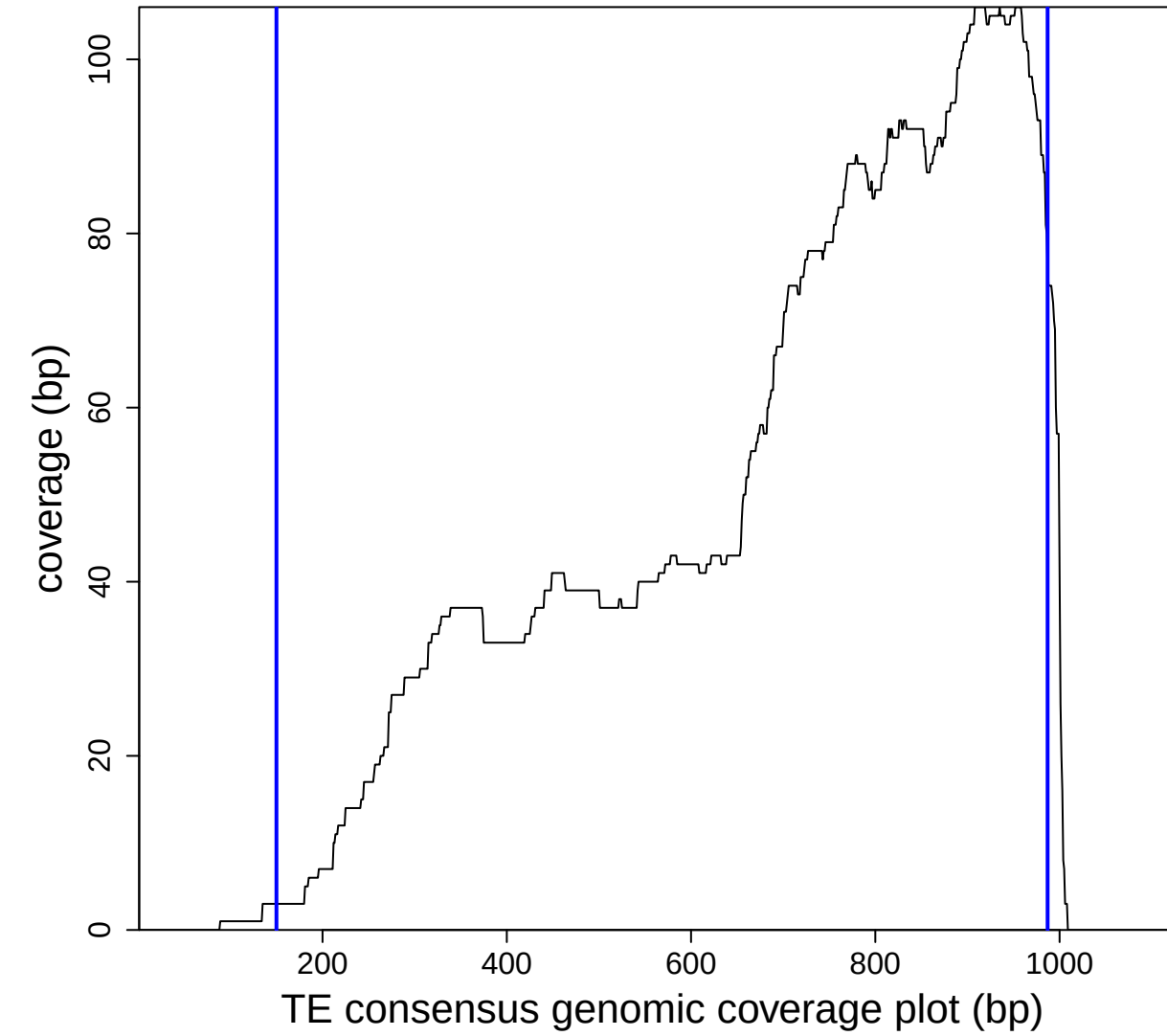
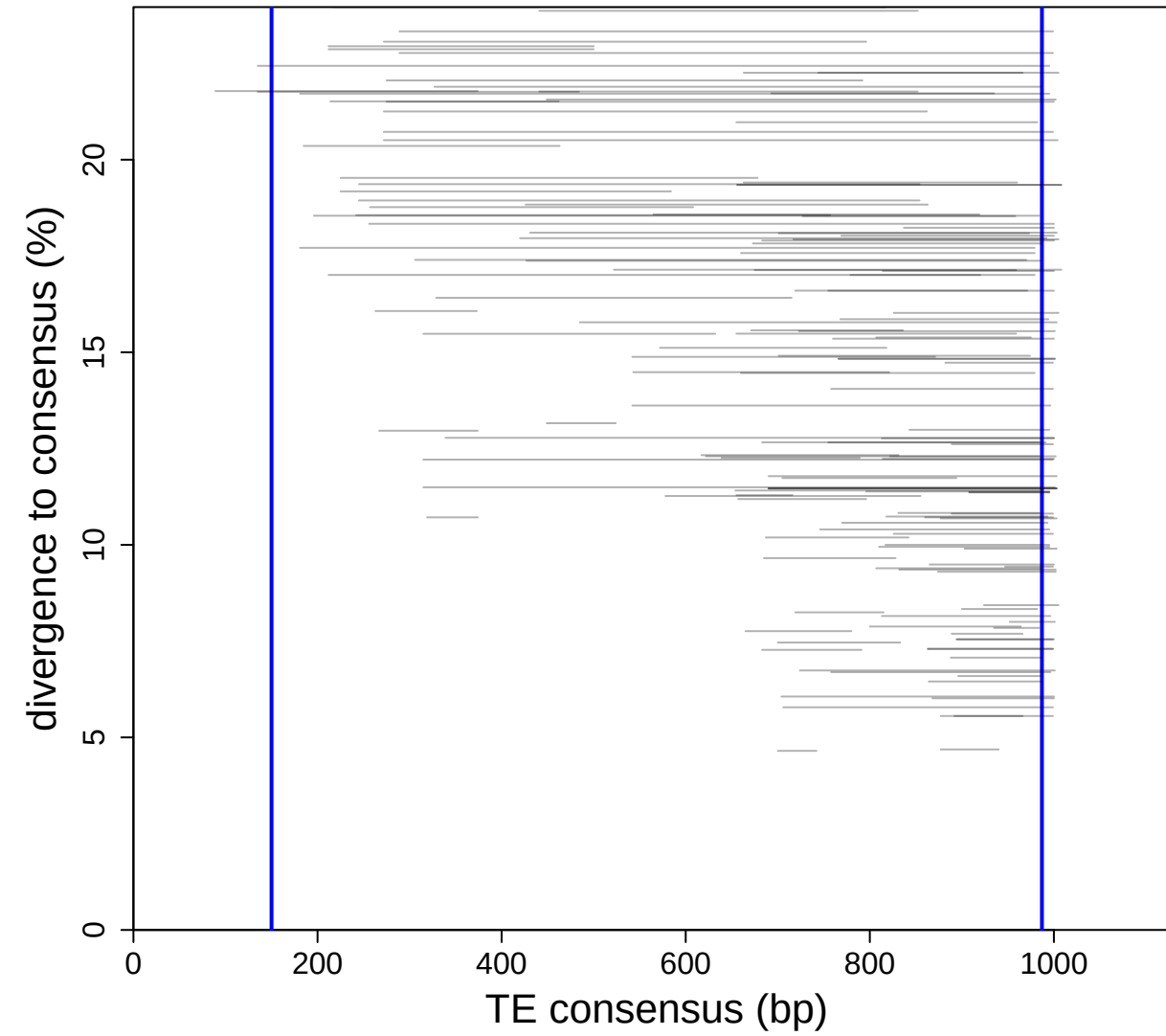
l.b.bed uf.bed g_2.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa
size: 763bp; fragments: 128; full length: 4 (>=686.7bp)

After TEtrimmer 763 bp

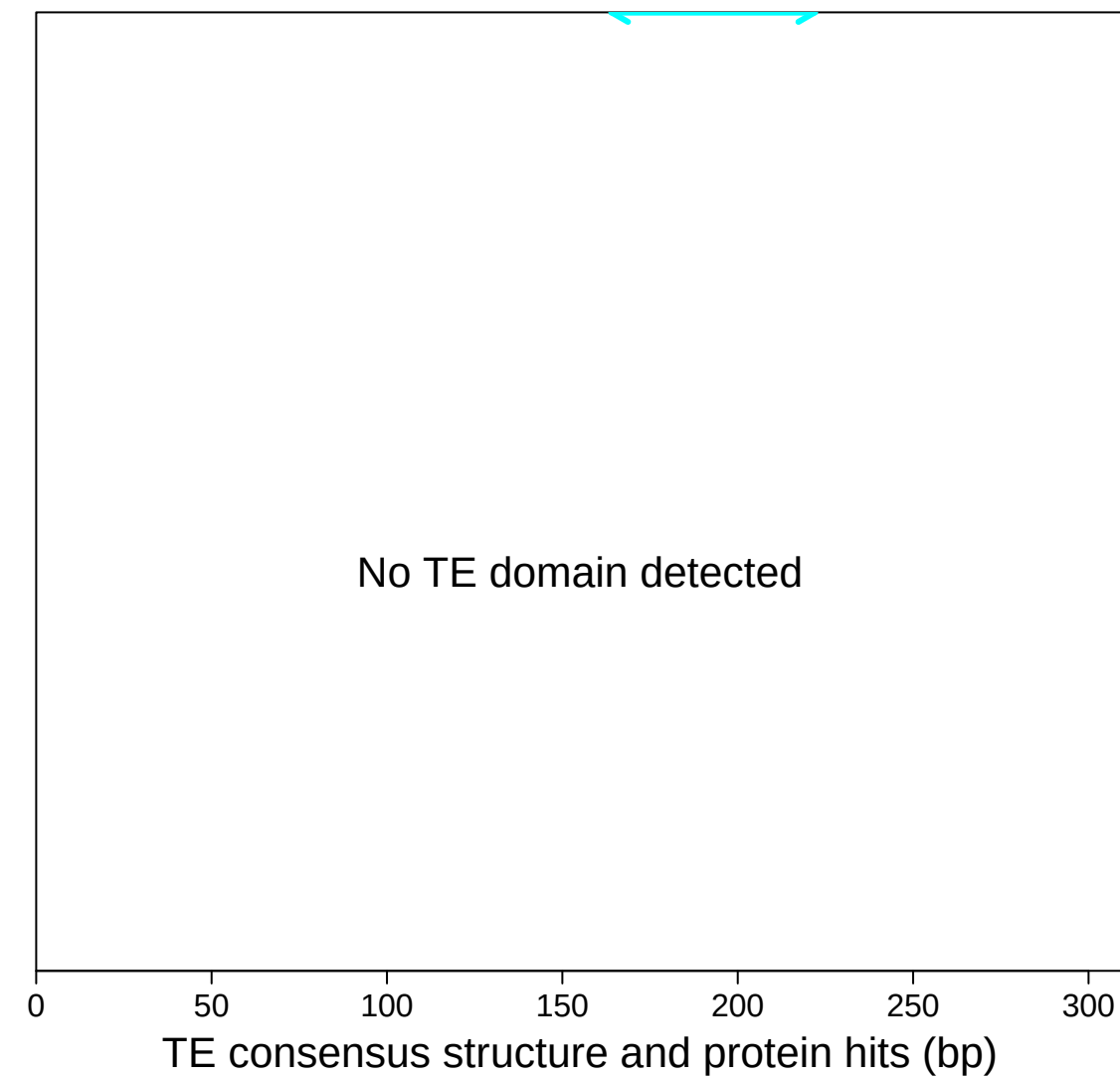
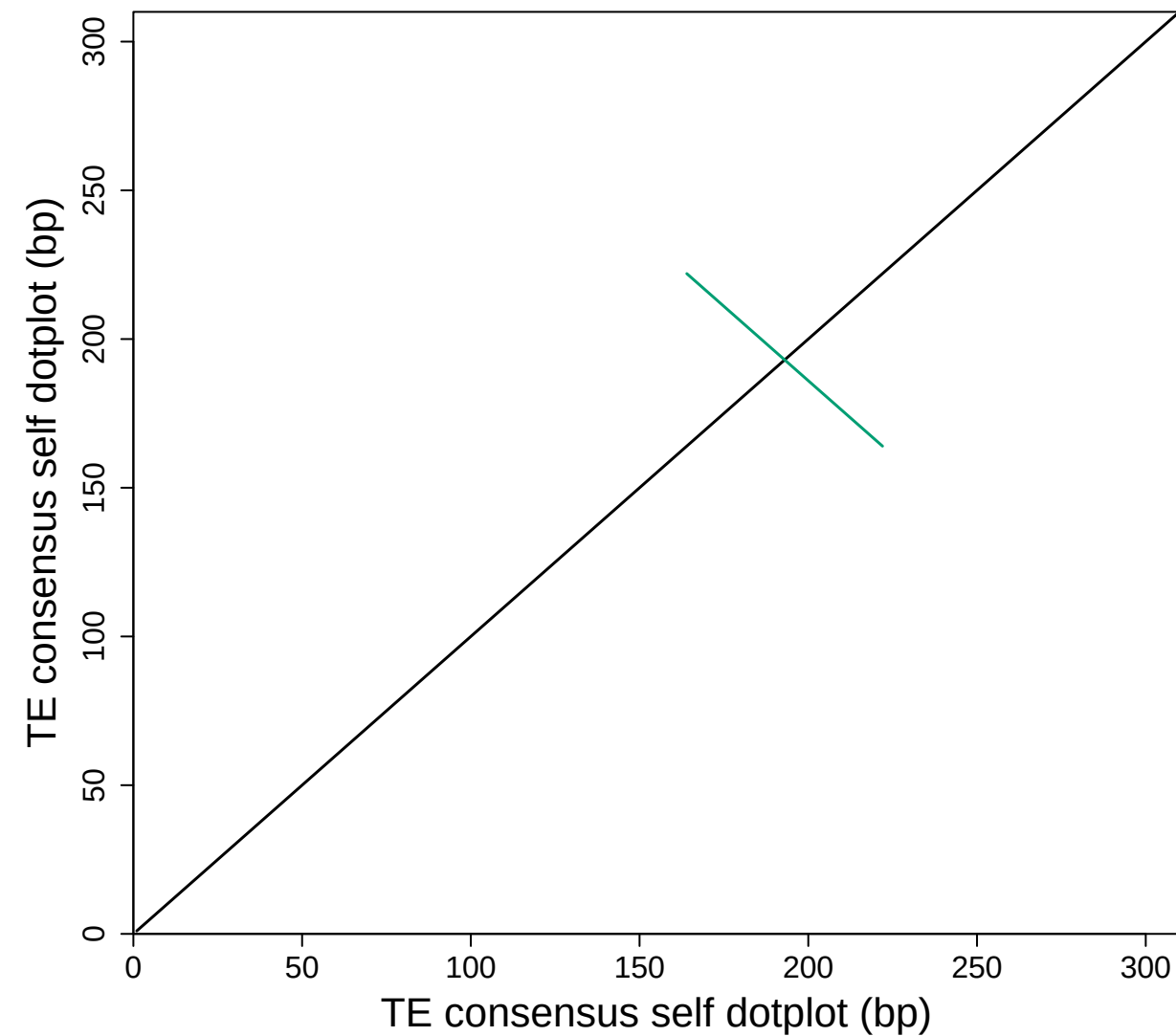
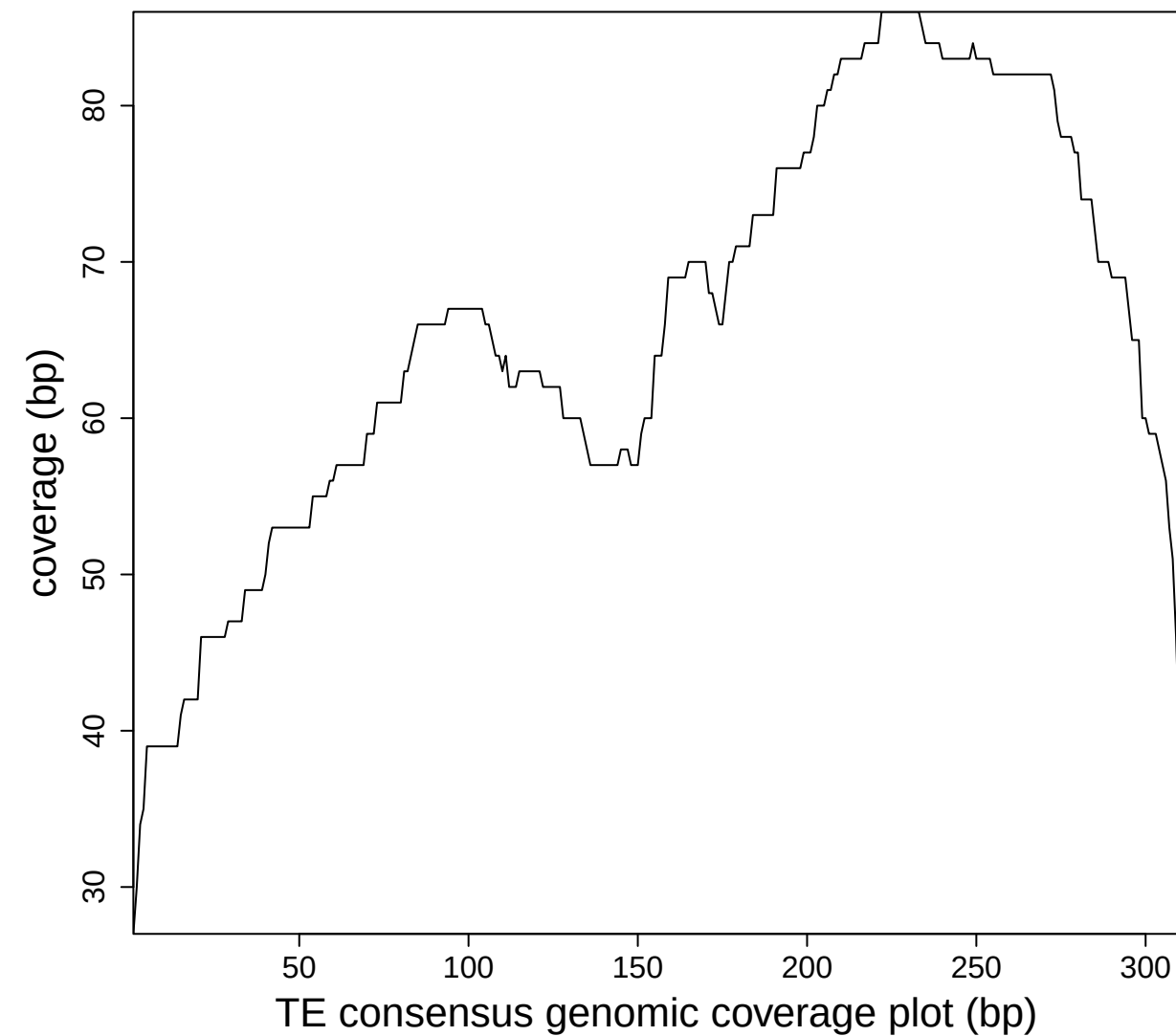
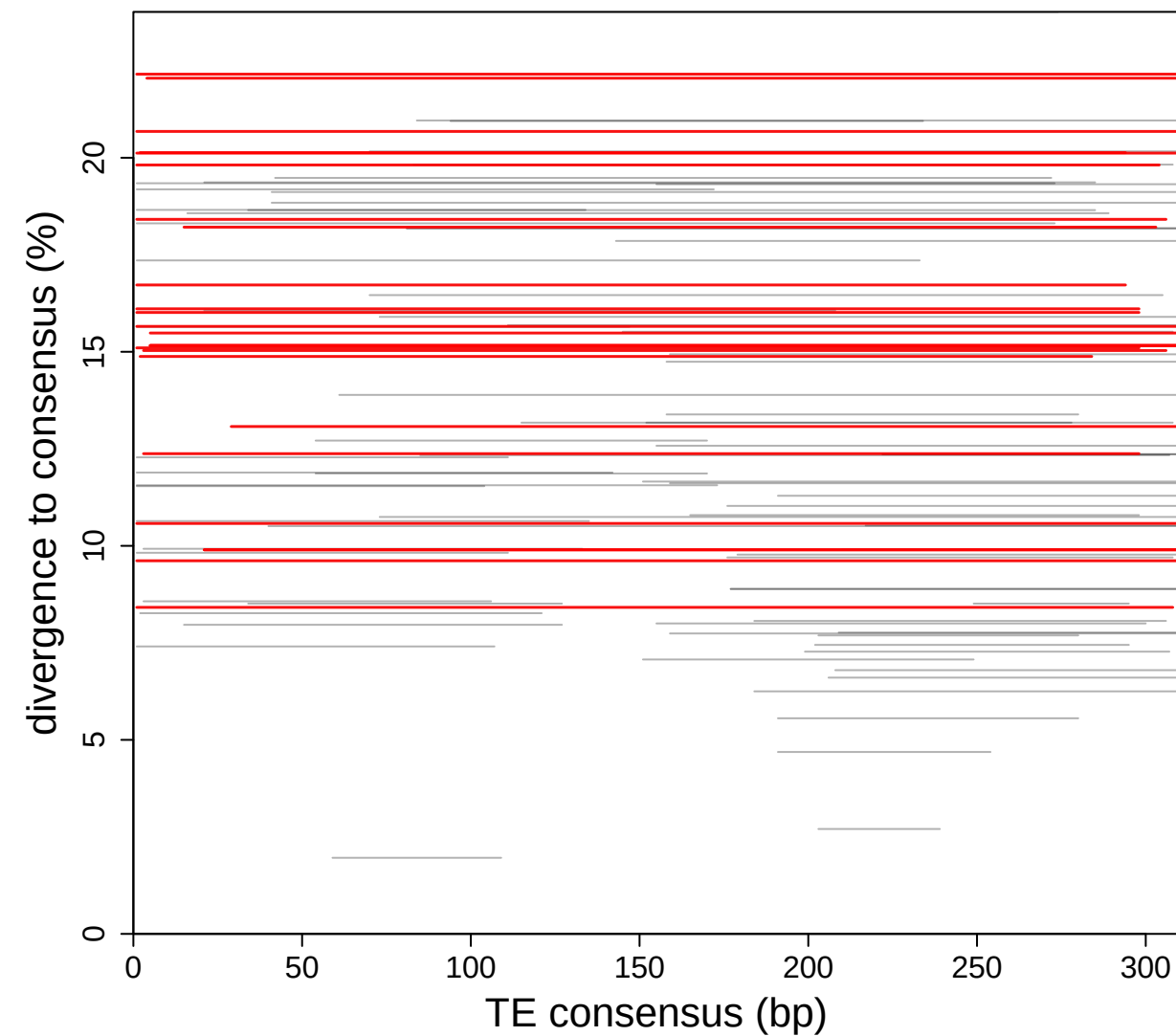


4.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 1137bp; fragments: 152; full length: 0 (>=1023.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

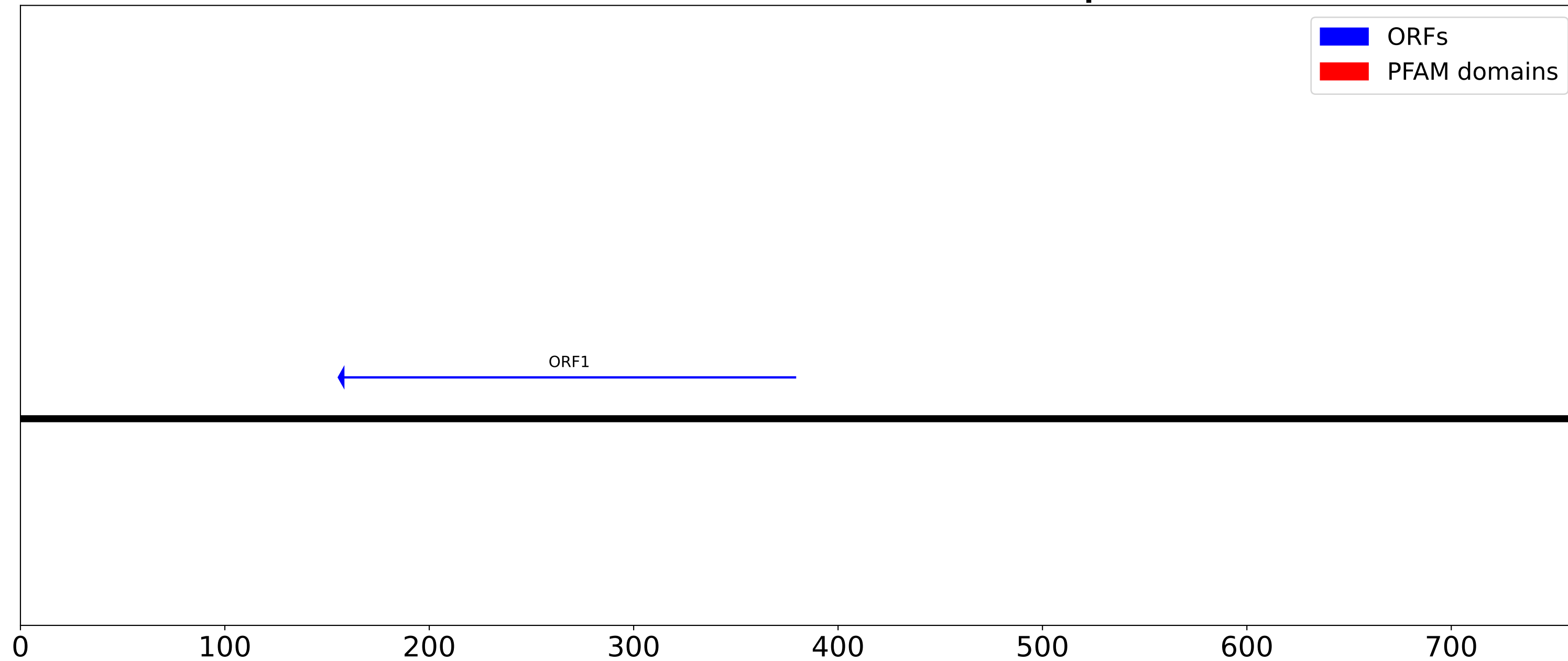


TE: rnd_5_family_574
size: 310bp; fragments: 106; full length: 26 (≥ 279 bp)



Before TEtrimmer 310 bp

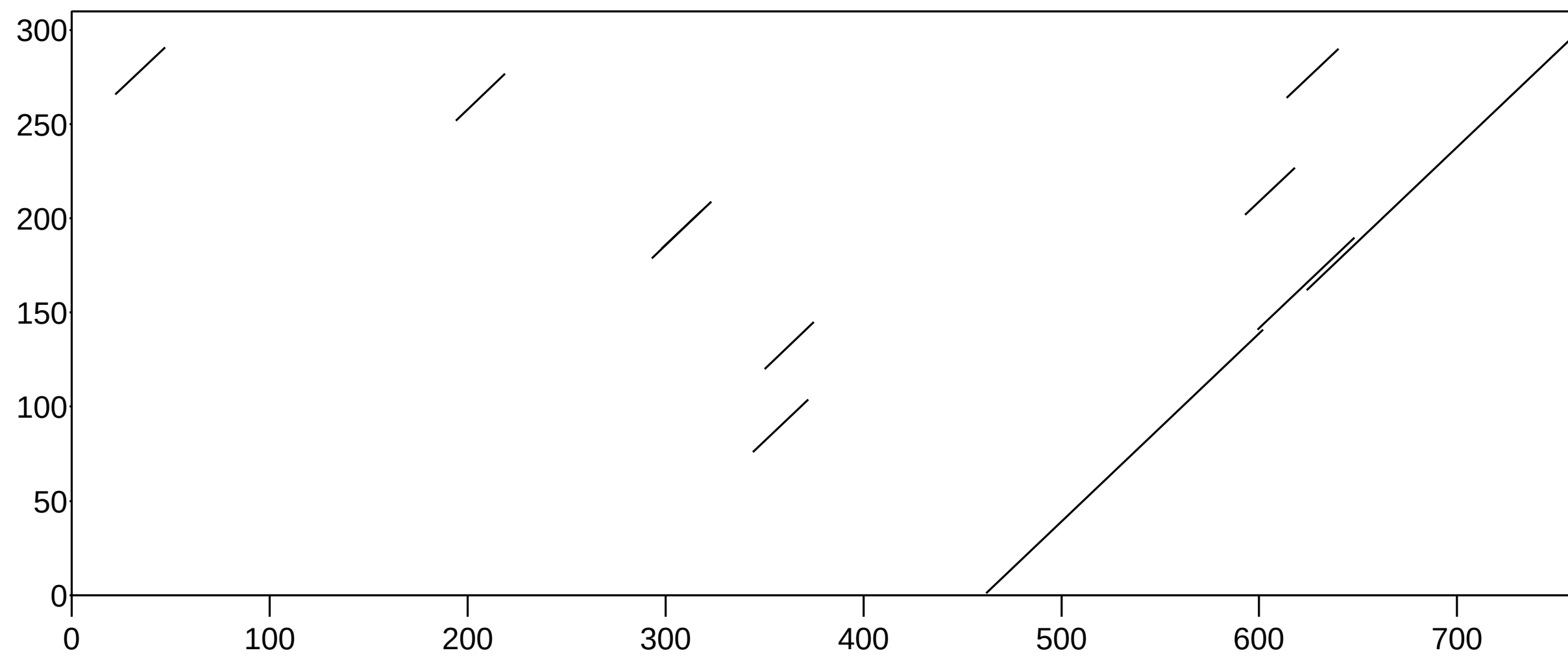
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)