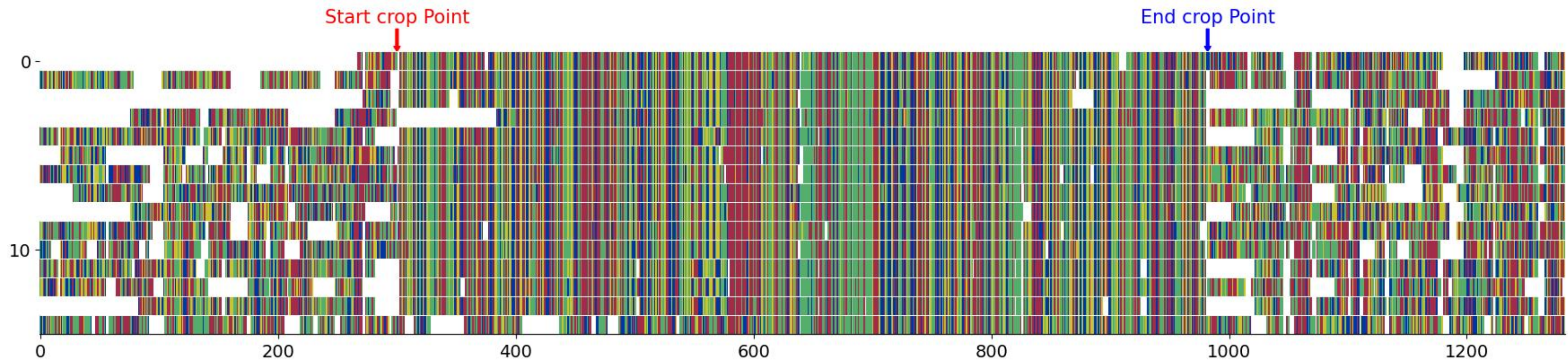
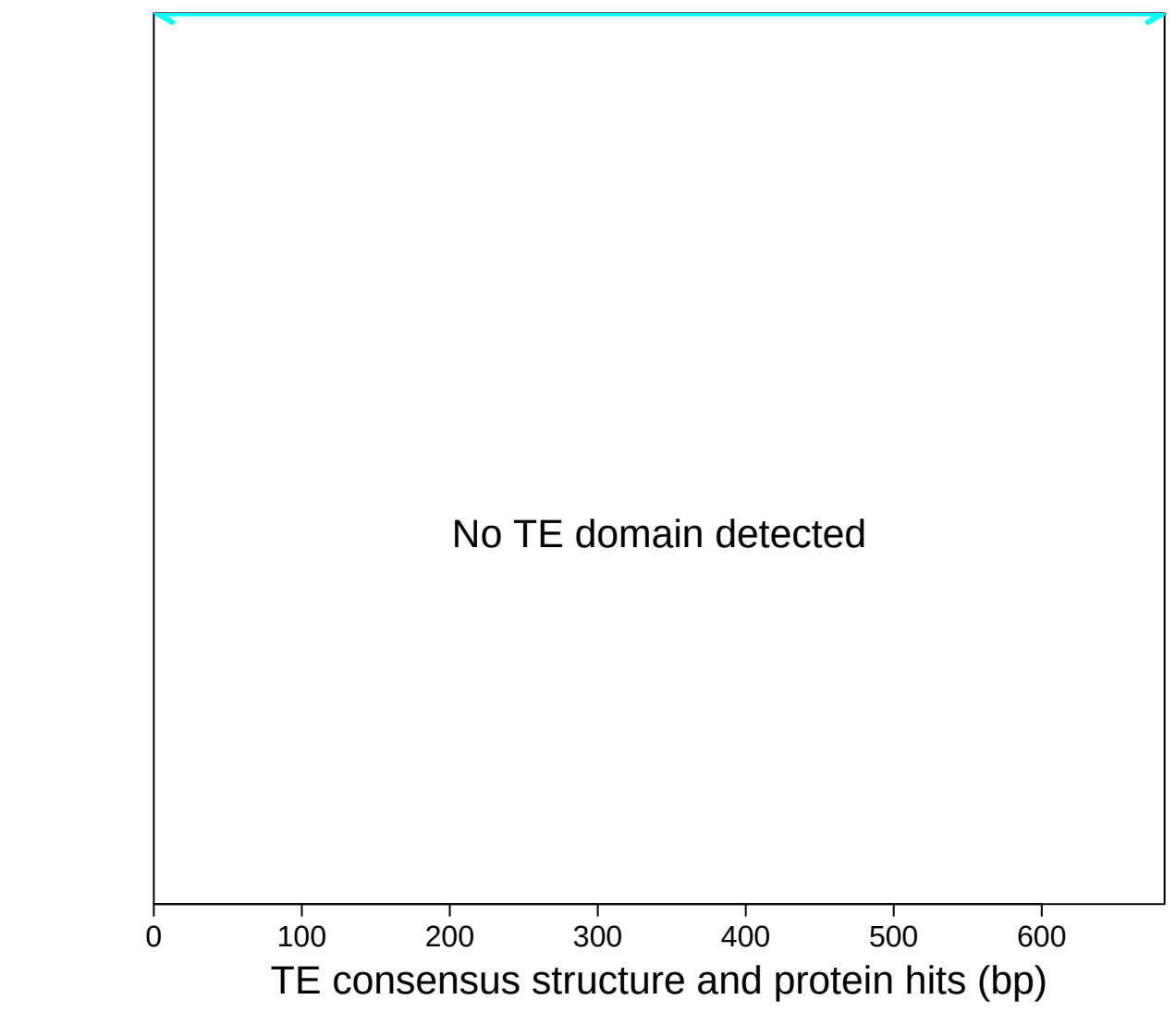
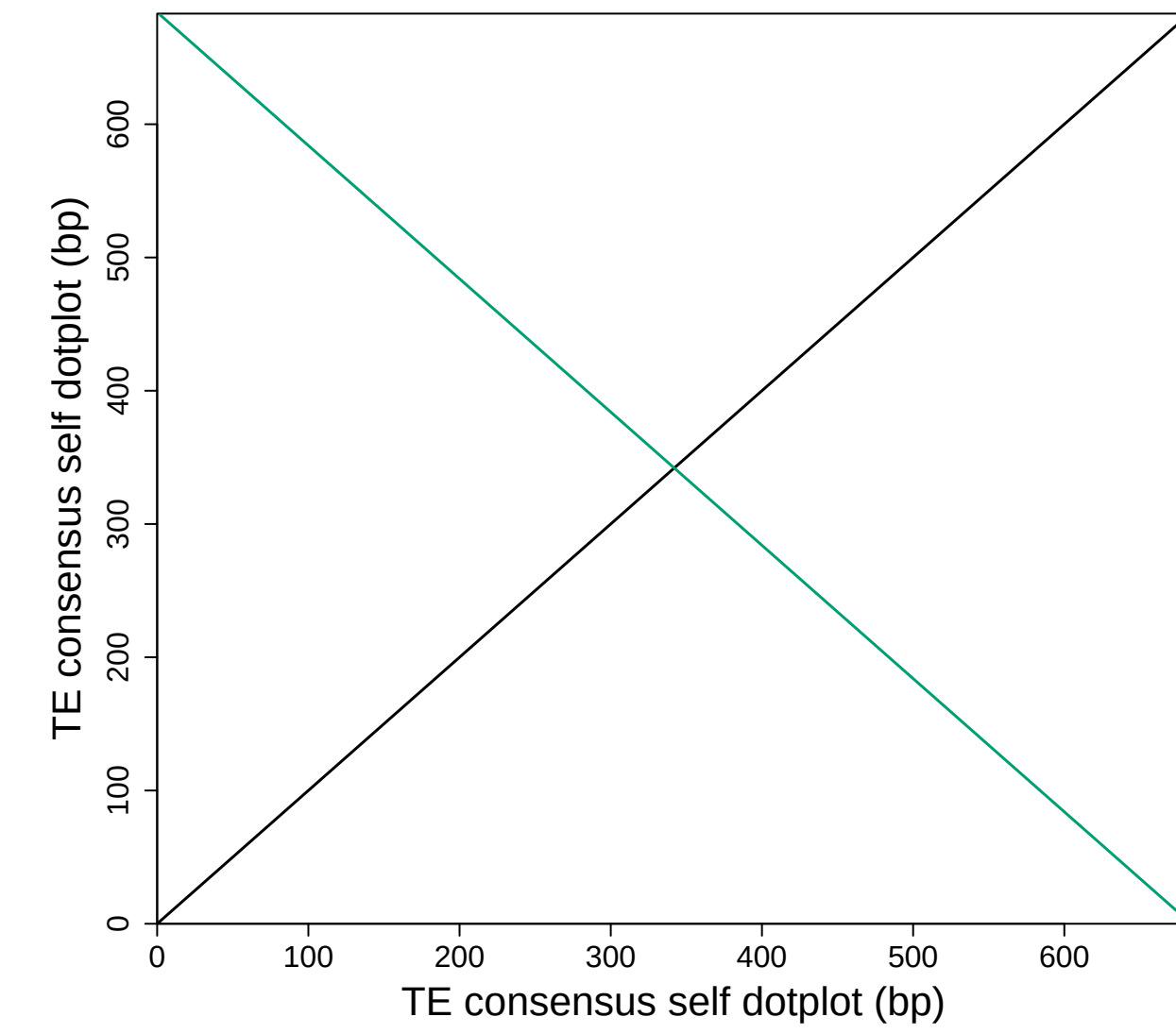
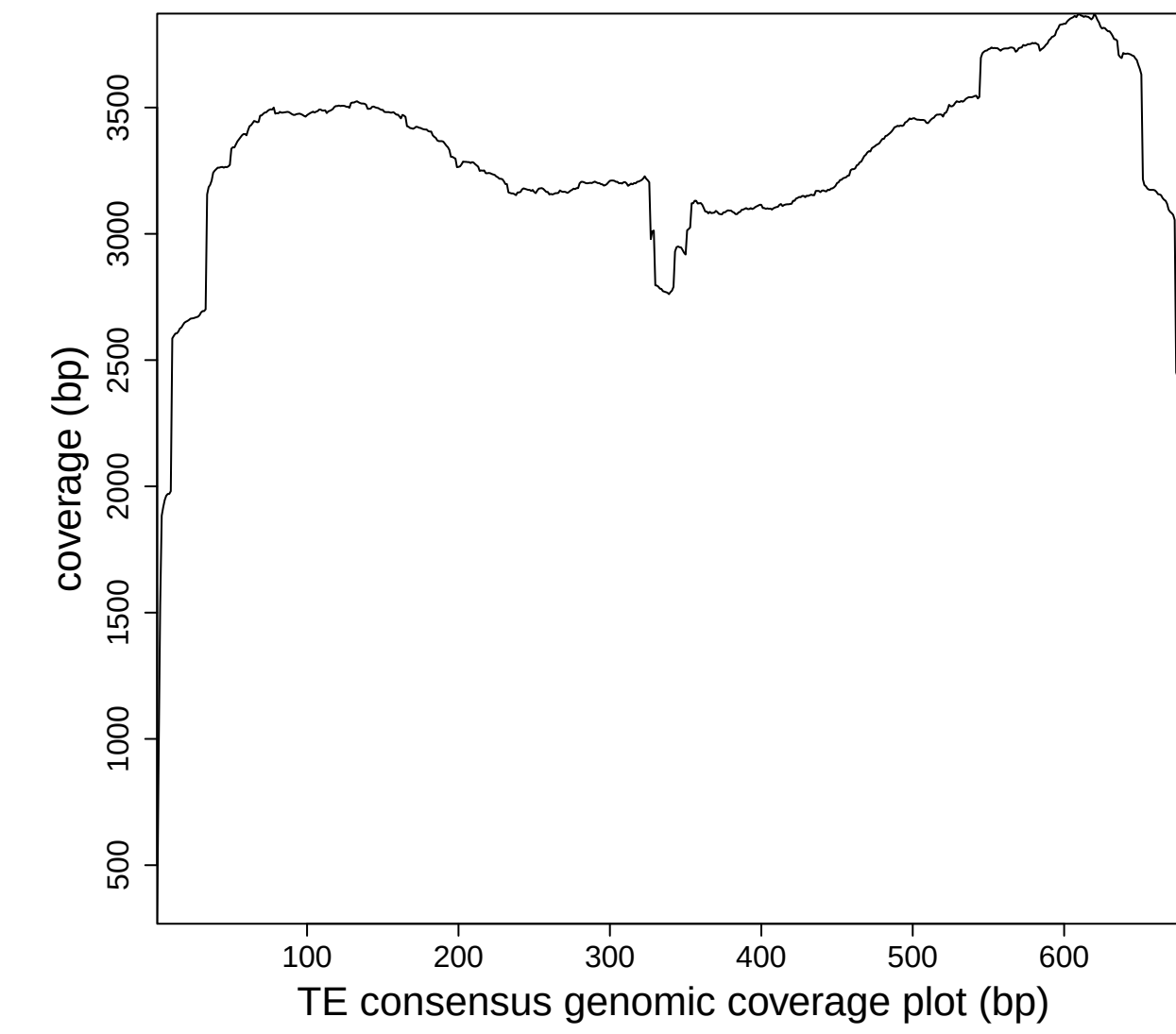
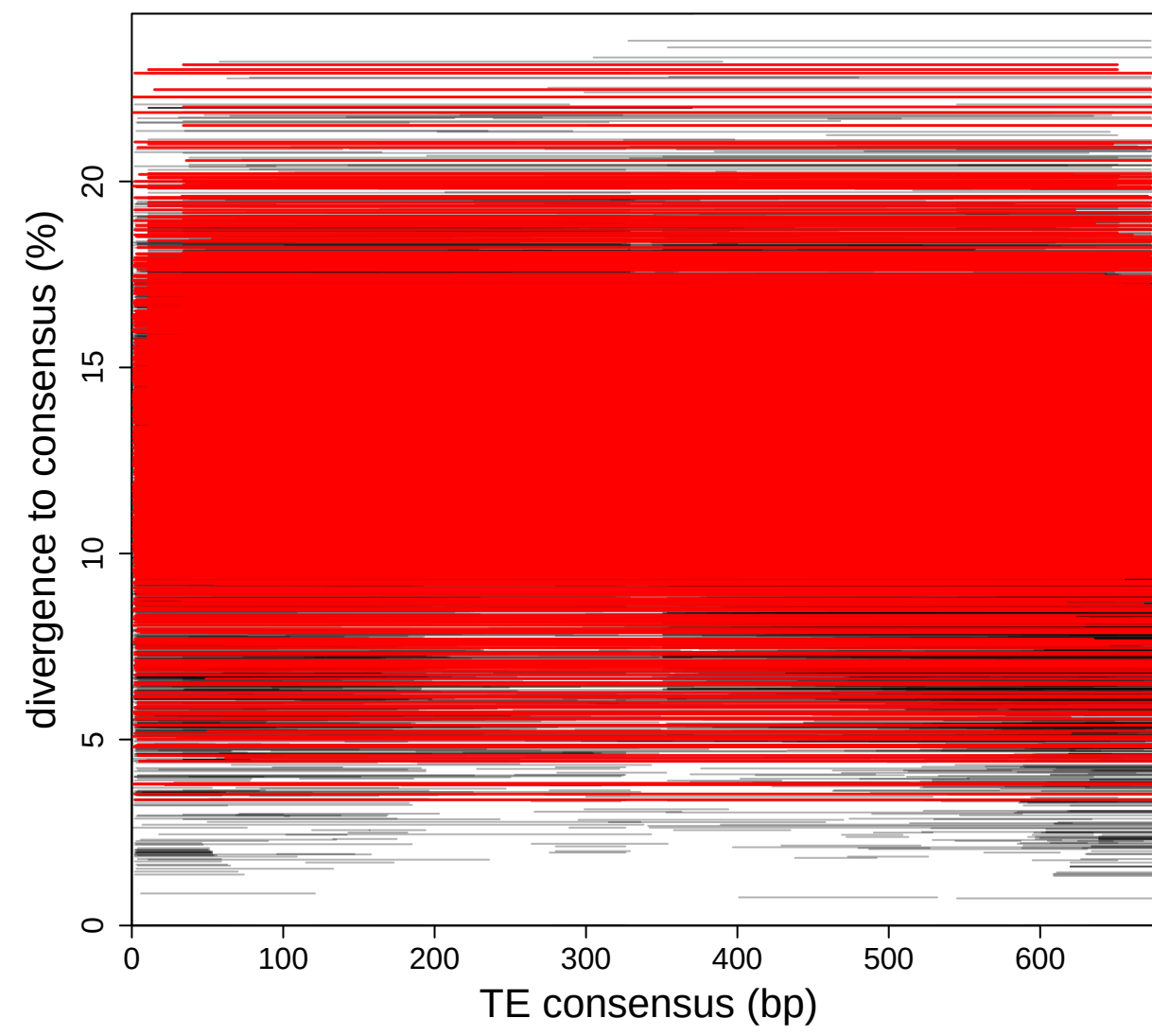






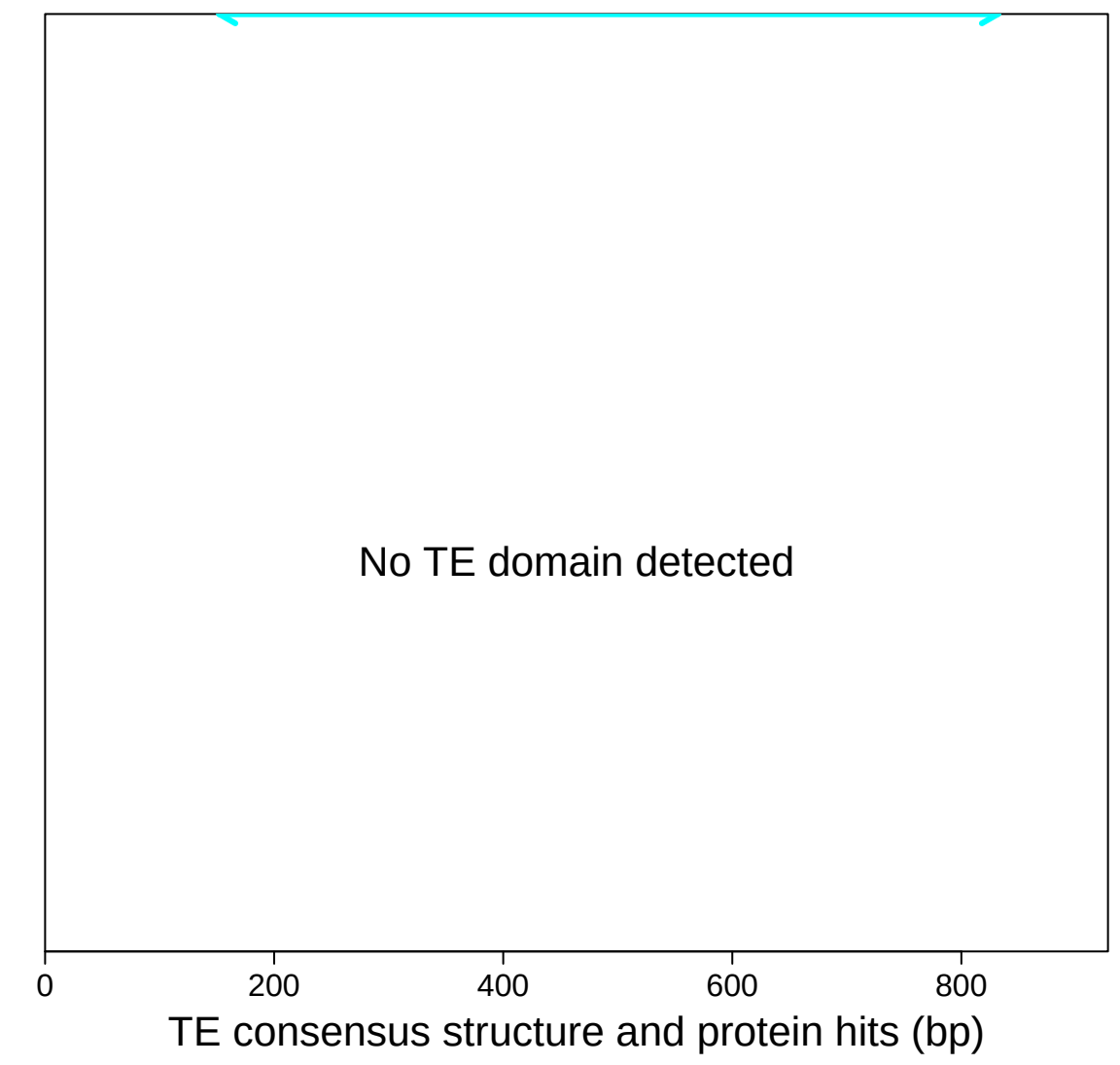
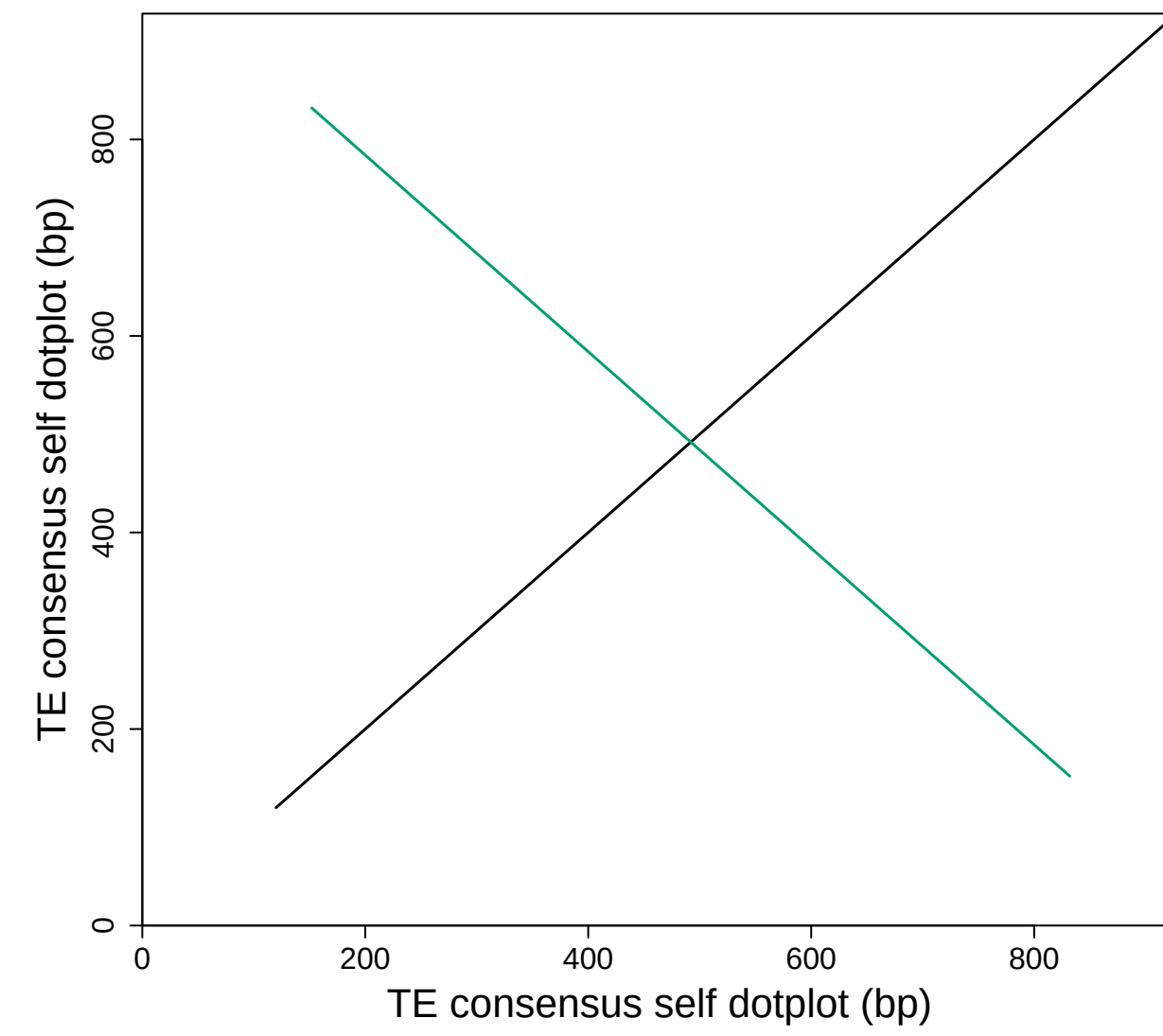
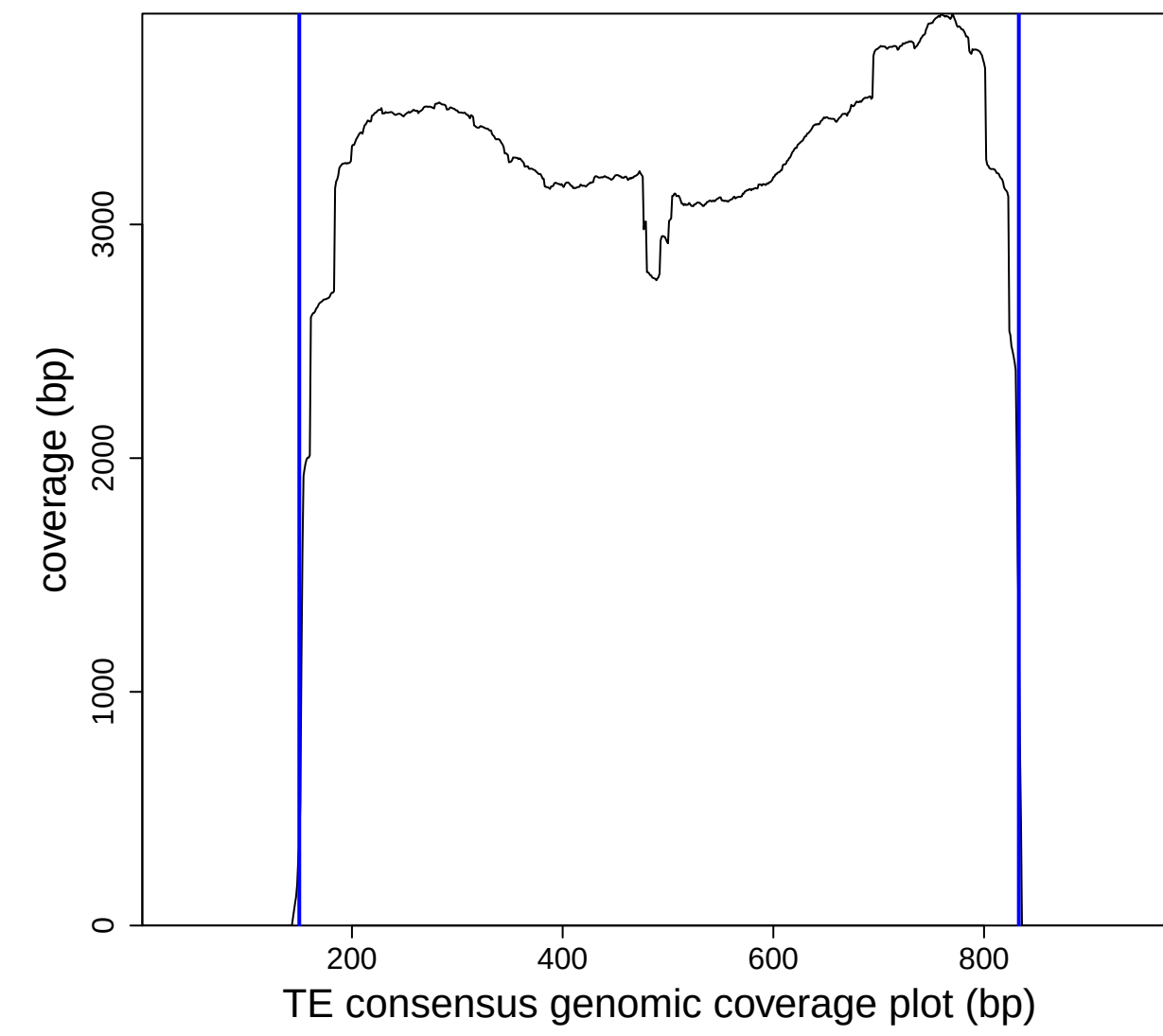
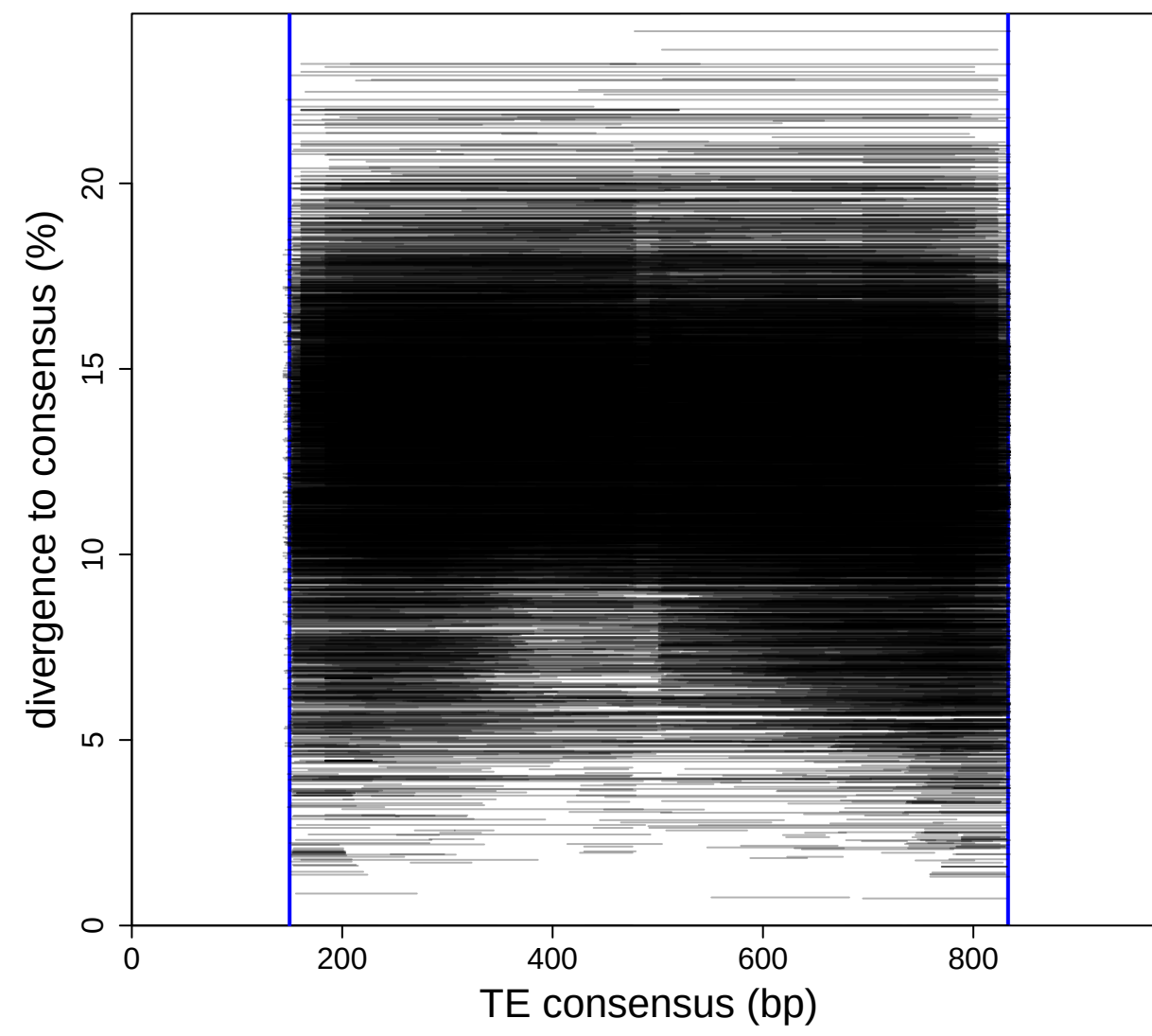
l.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 683bp; fragments: 6505; full length: 1326 (≥ 614.7 bp)

After TEtrimmer 683 bp



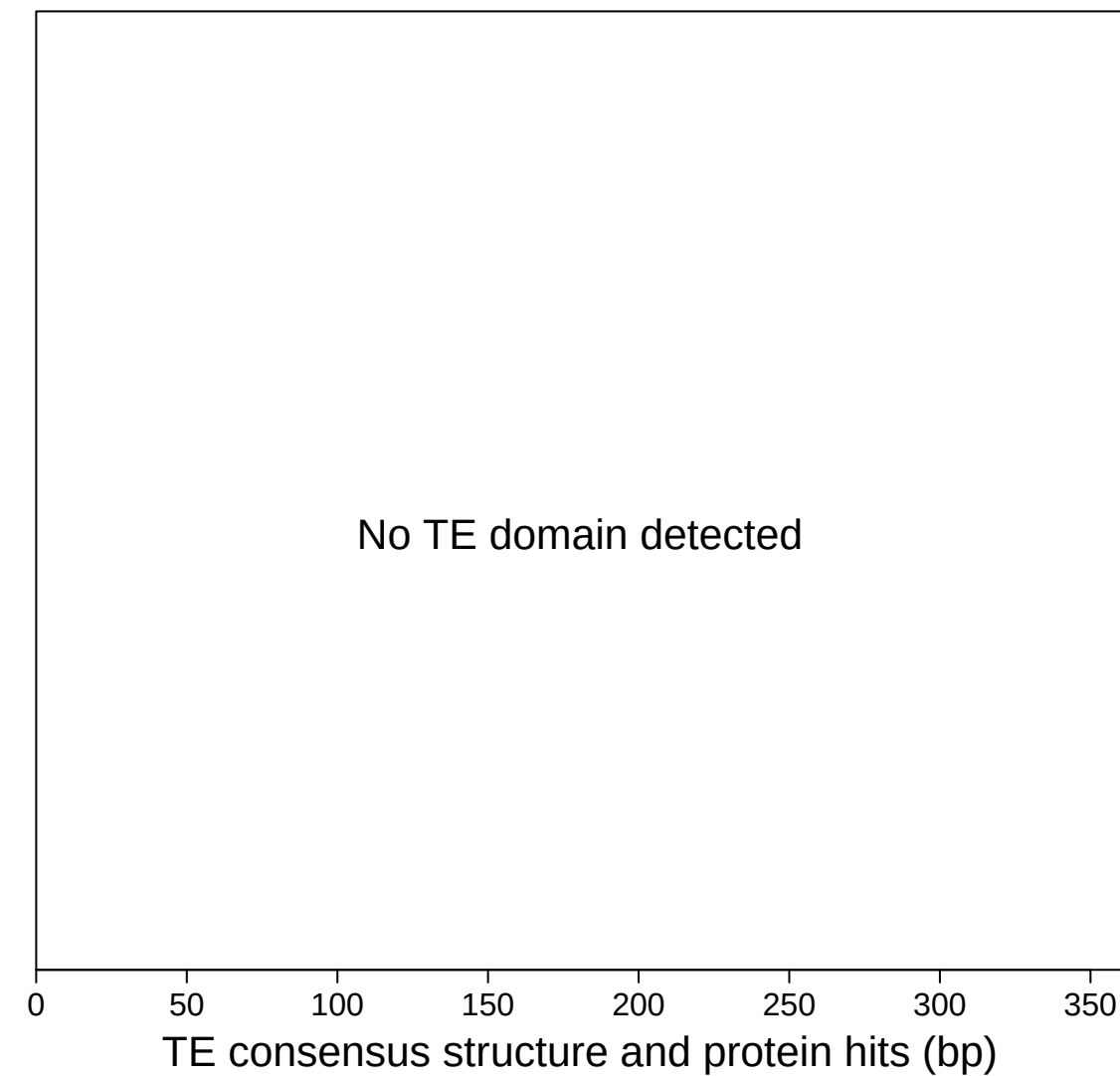
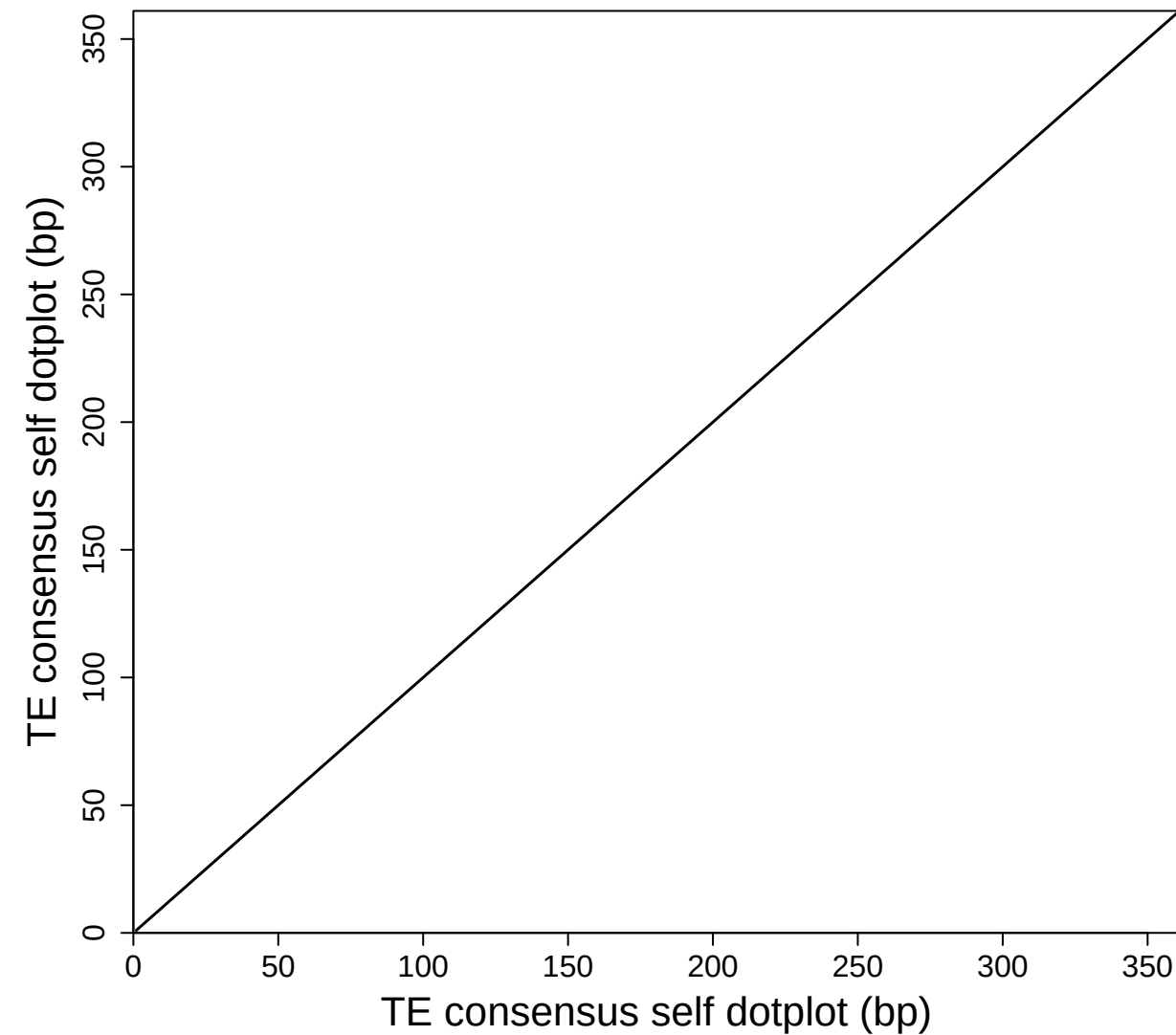
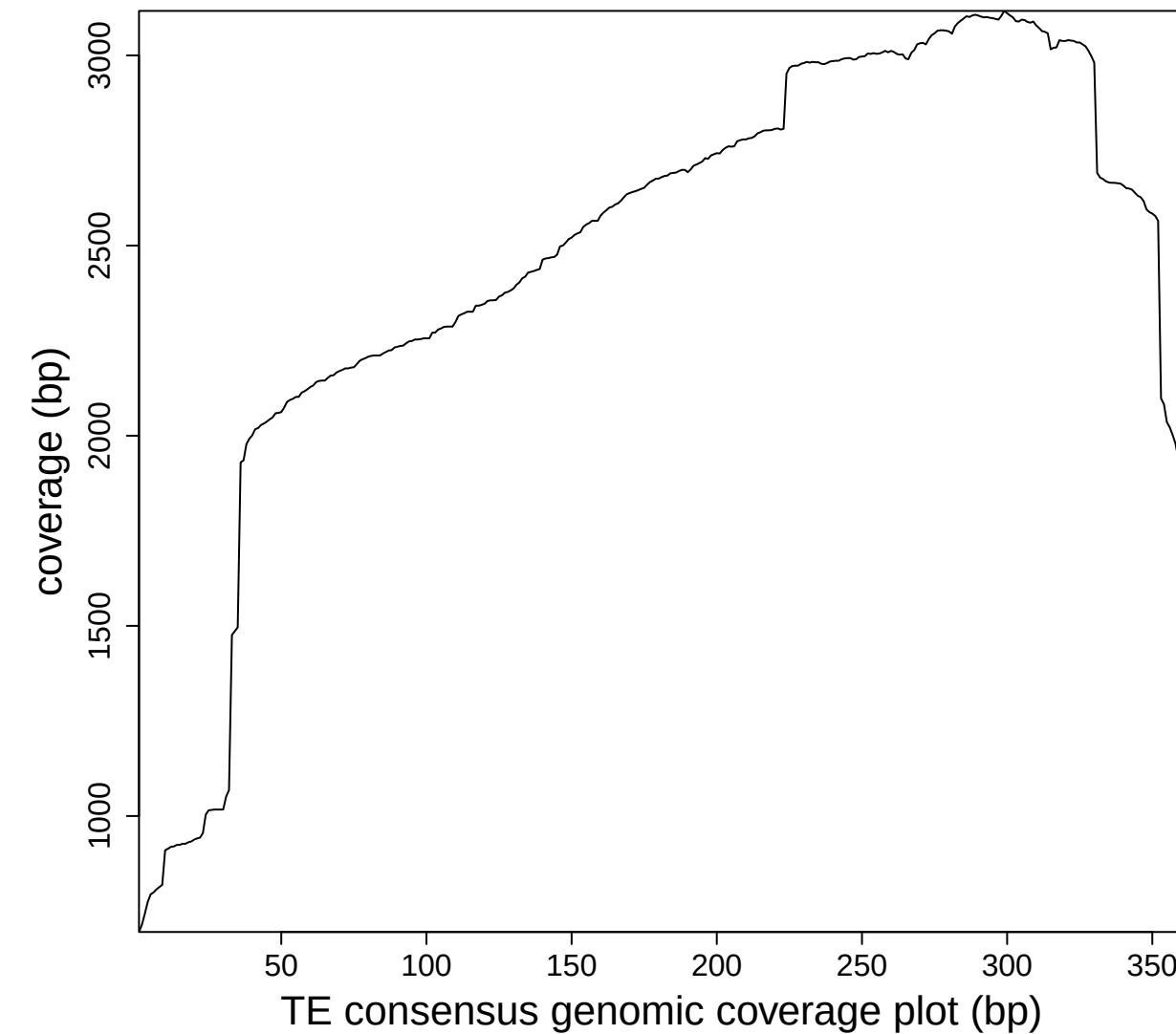
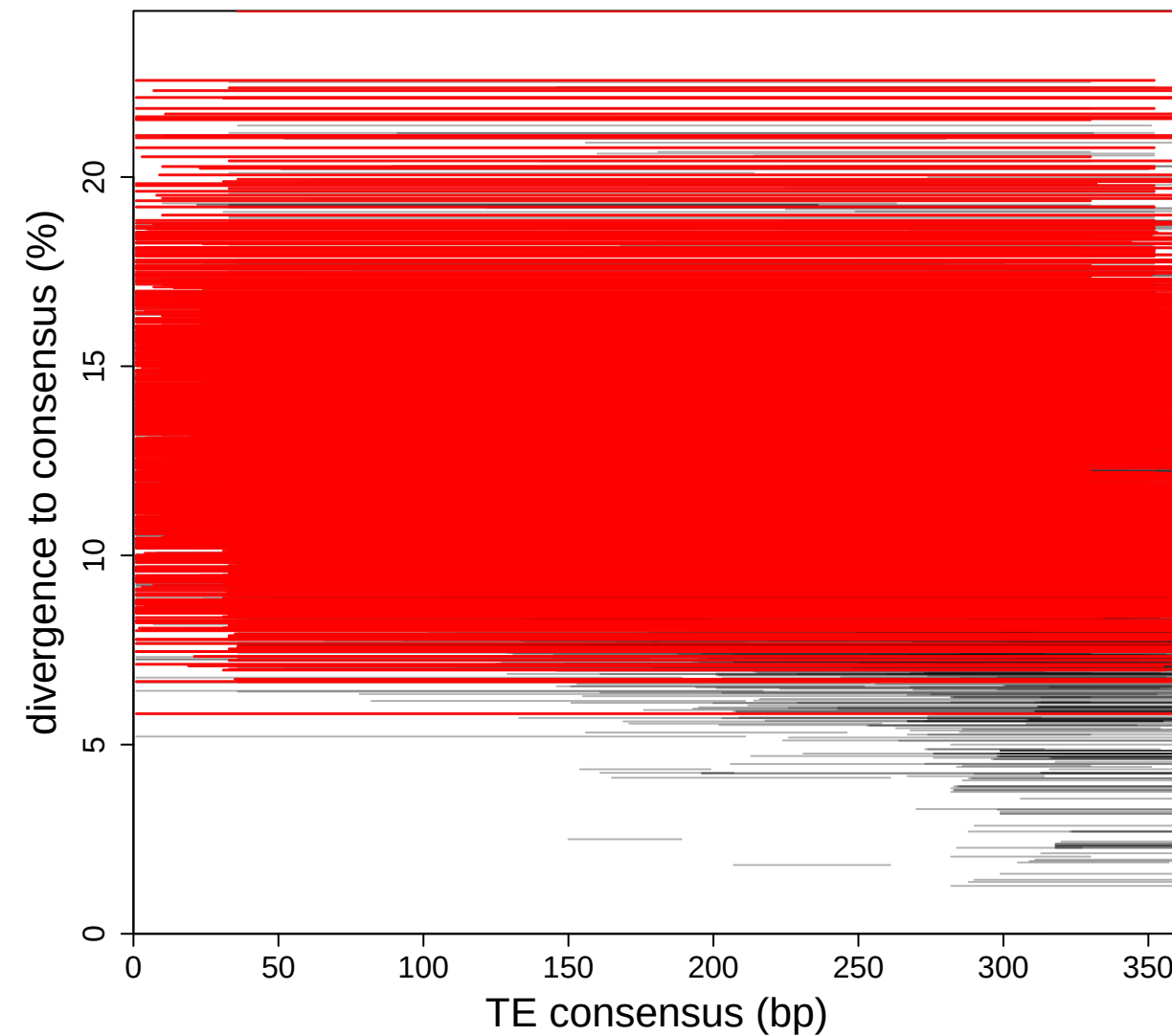
1.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 983bp; fragments: 6538; full length: 0 (>=884.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

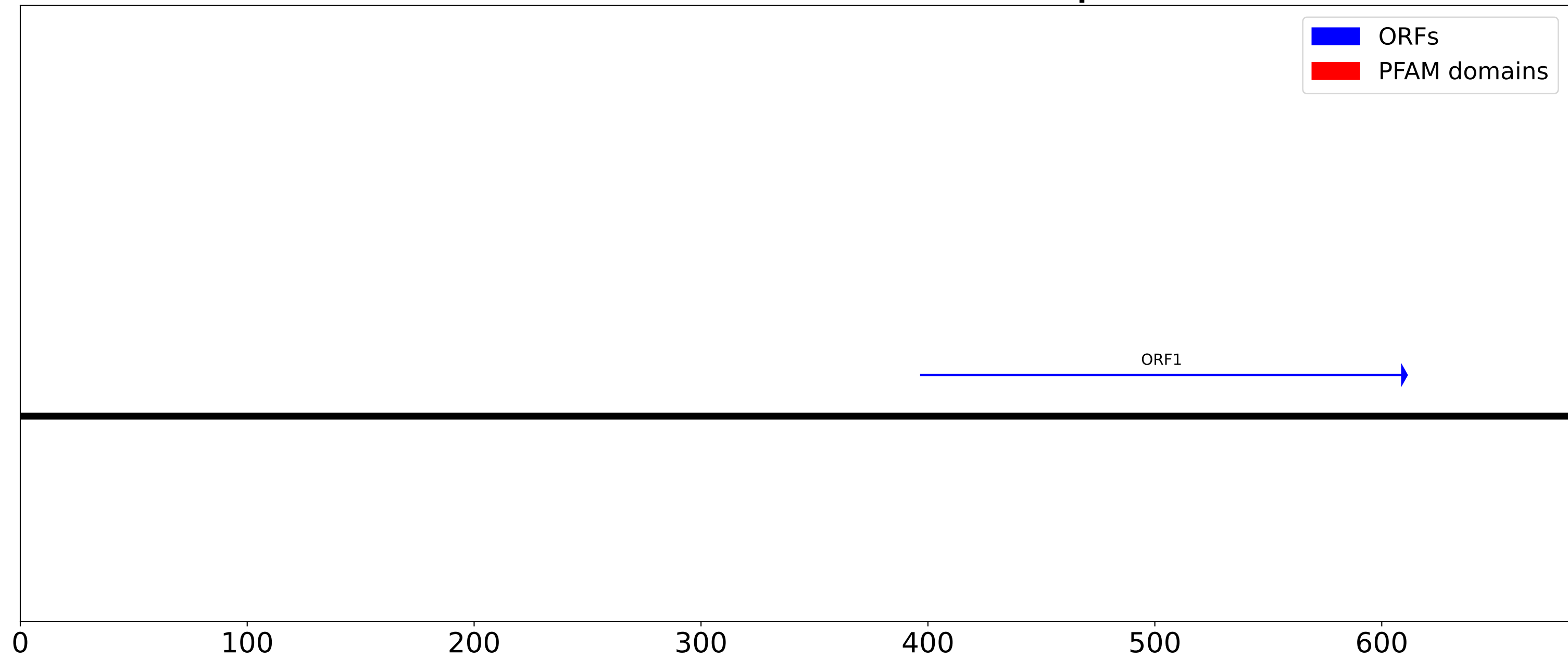


TE: rnd_1_family_131
size: 361bp; fragments: 3402; full length: 1313 (≥ 324.9 bp)

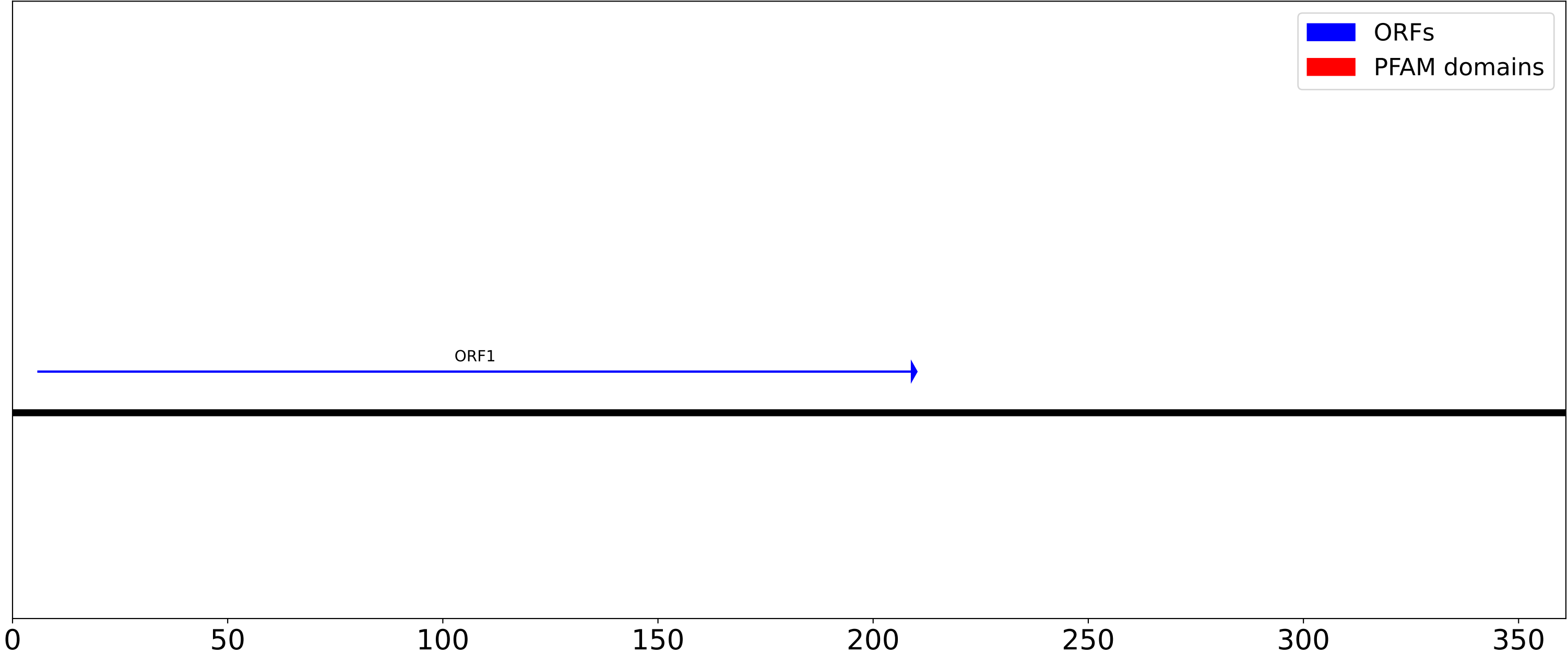
Before TEtrimmer 361 bp



After TEtrimmer ORF and PFAM domain plot



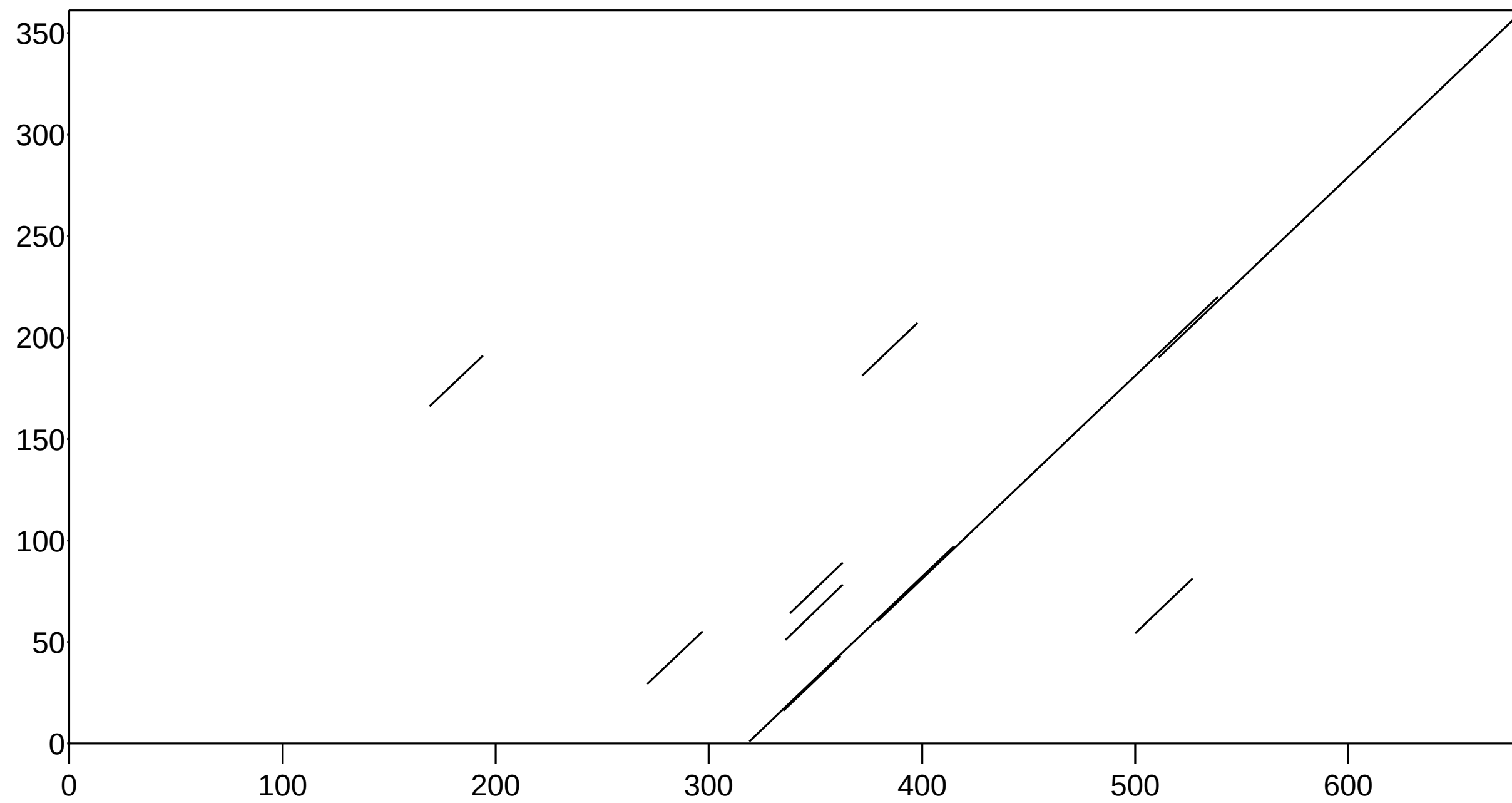
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)