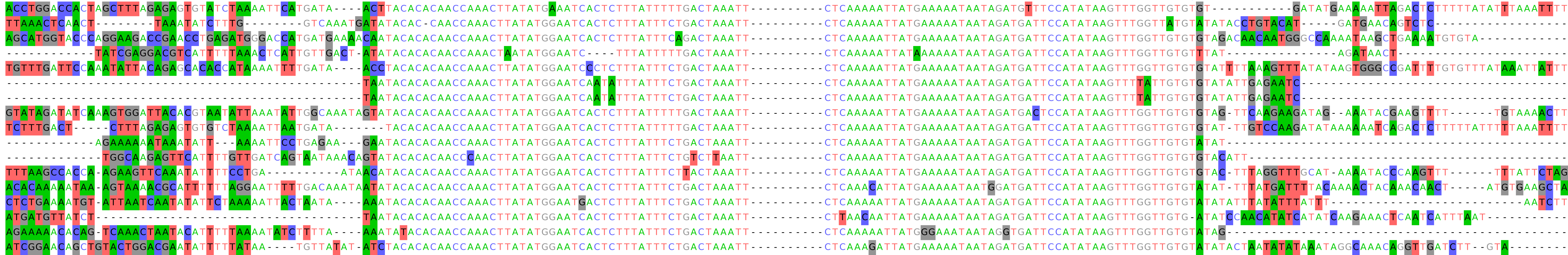


Start crop Point

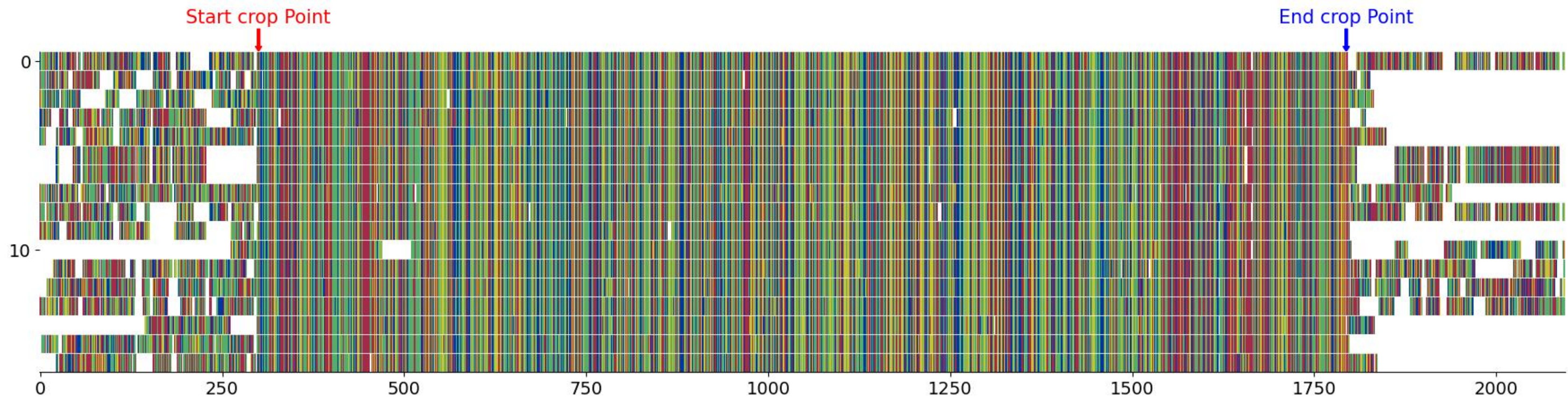
End crop Point

MSA length = 1495



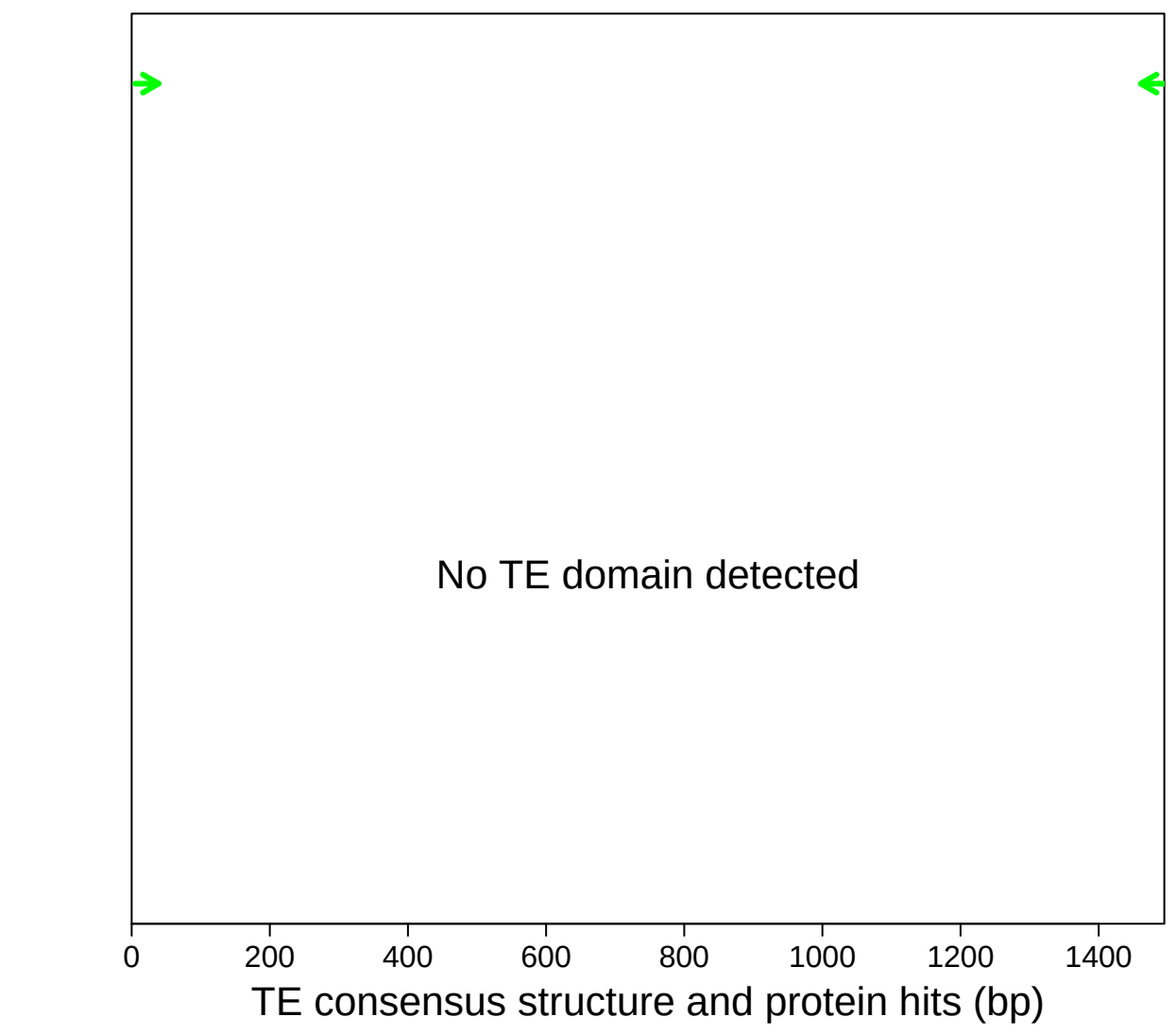
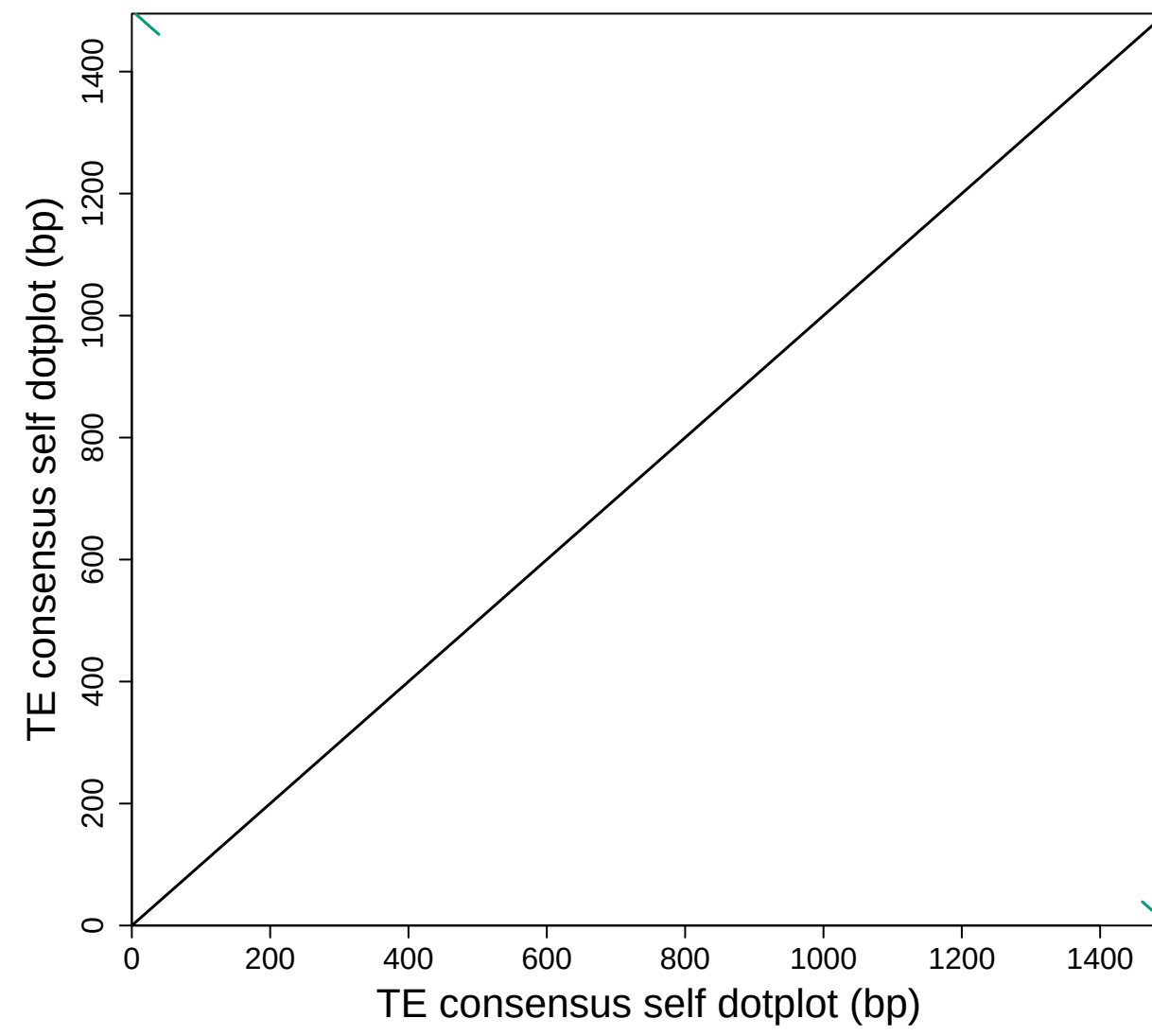
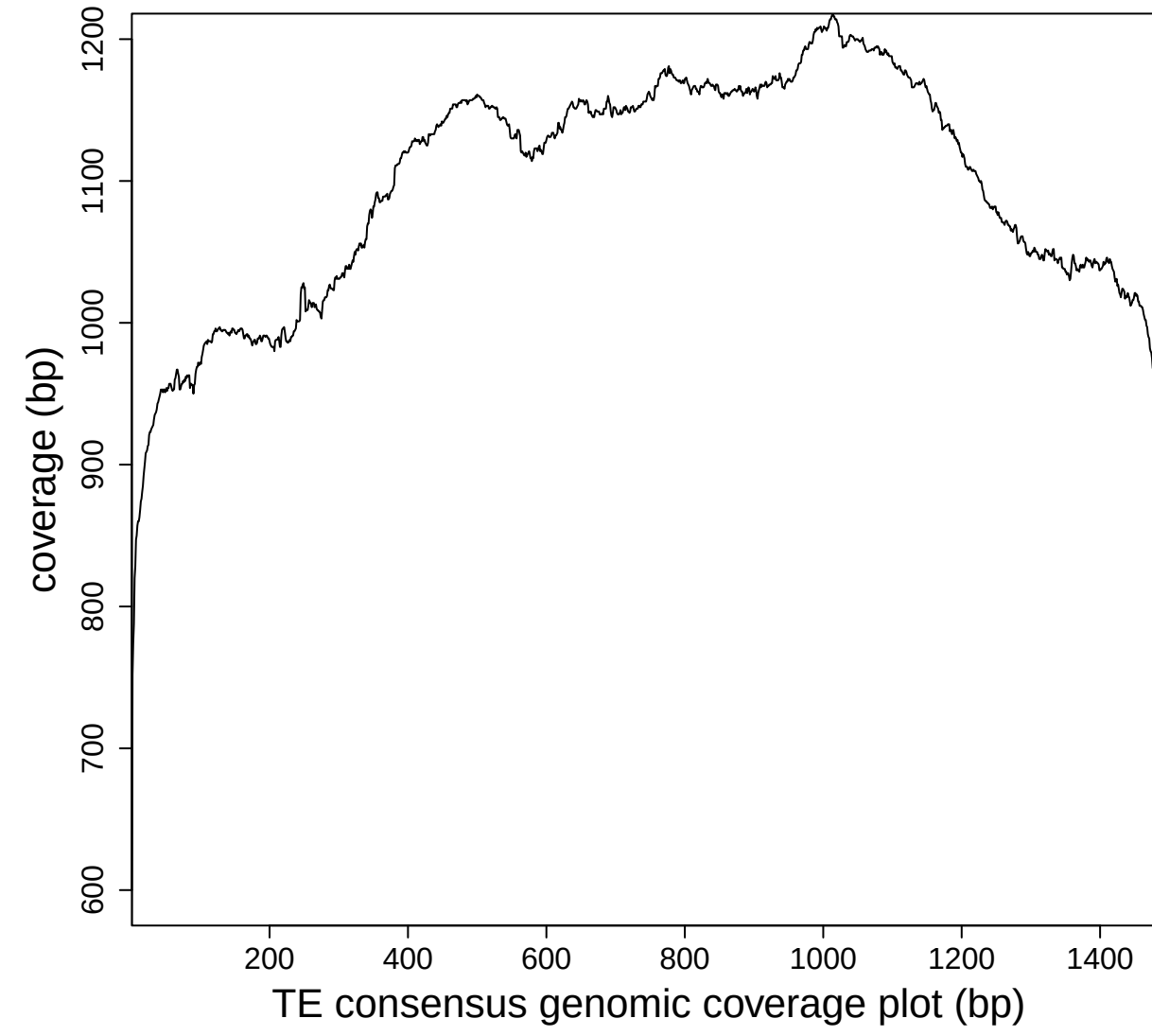
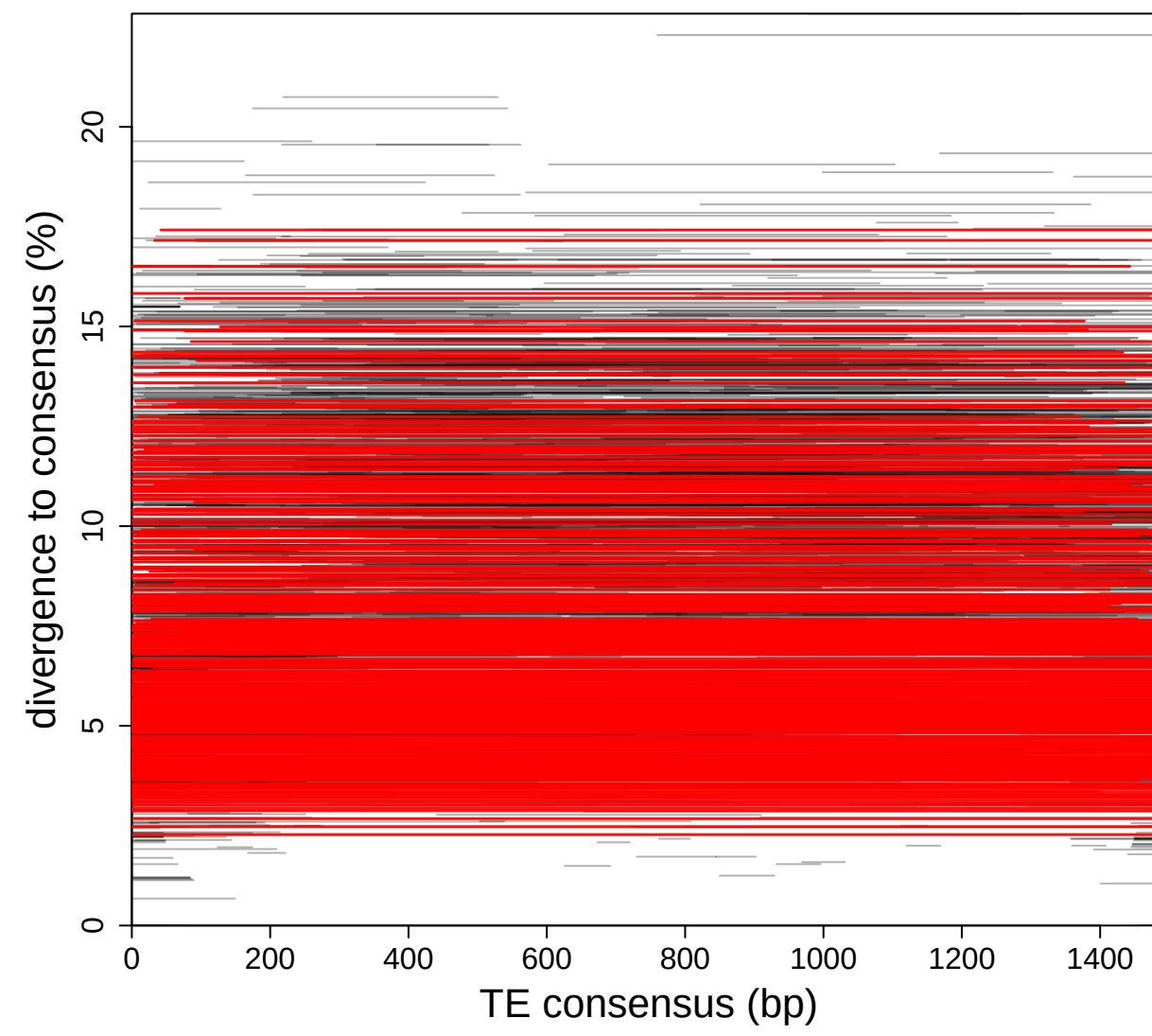
Start crop Point

End crop Point



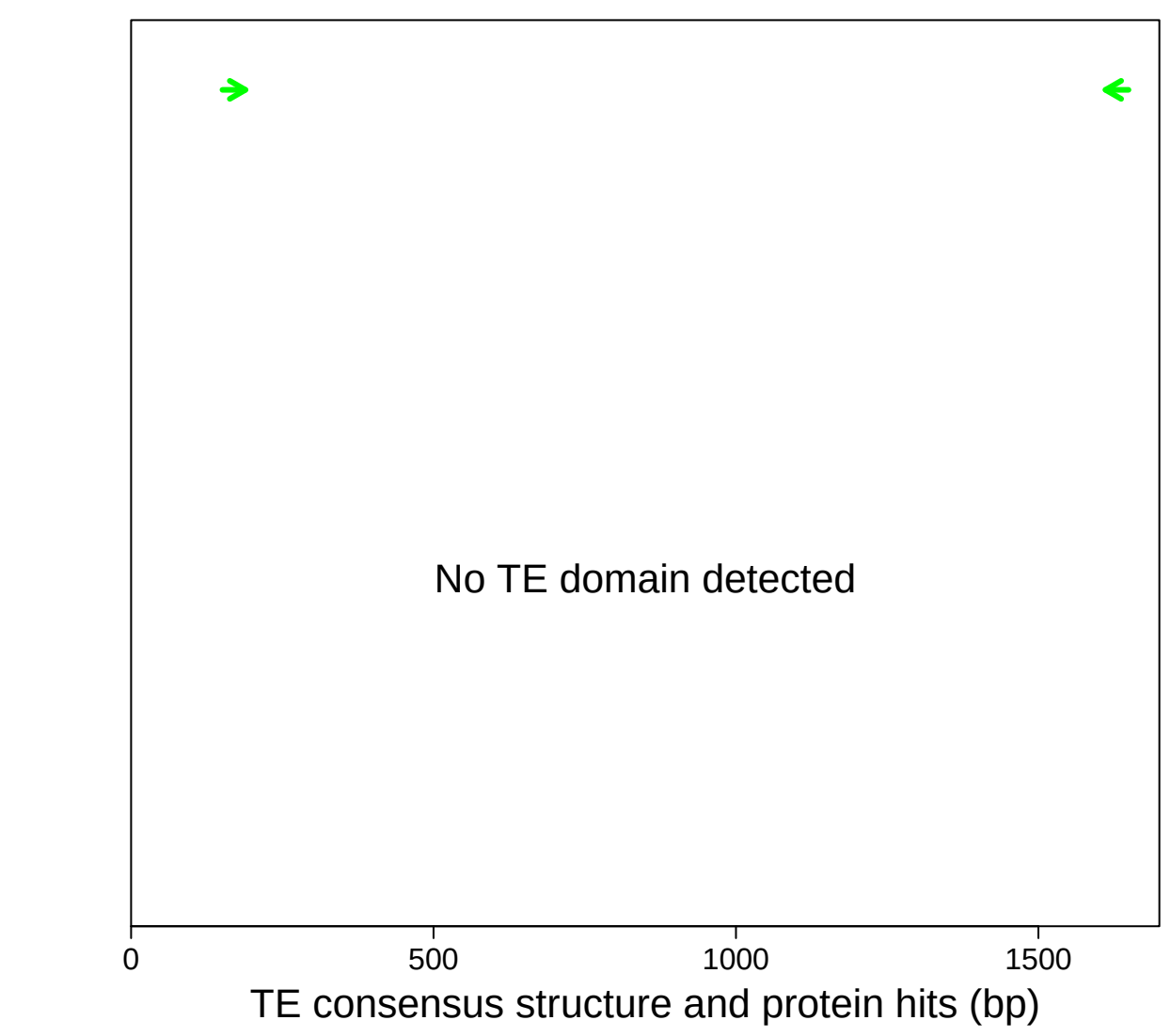
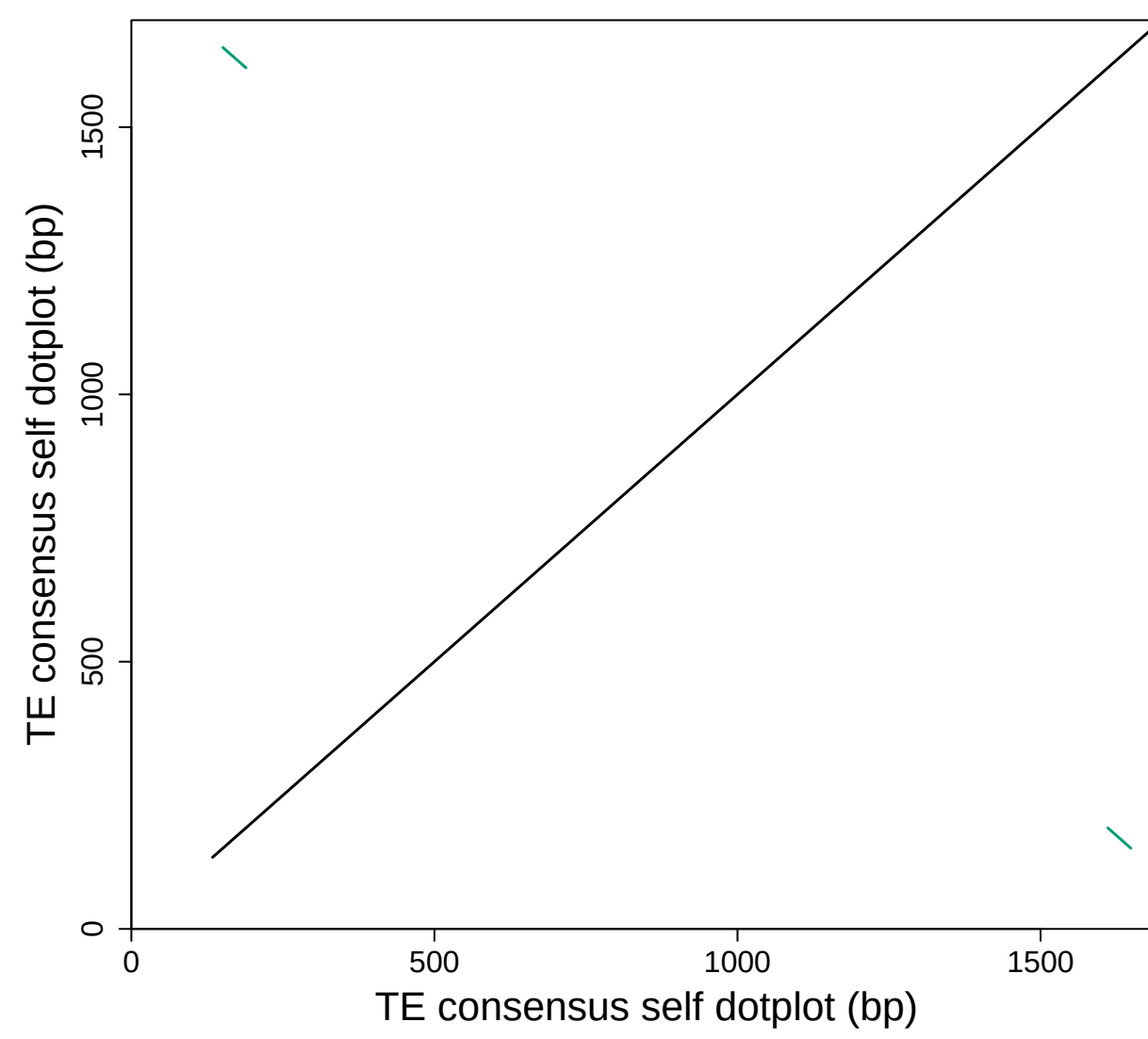
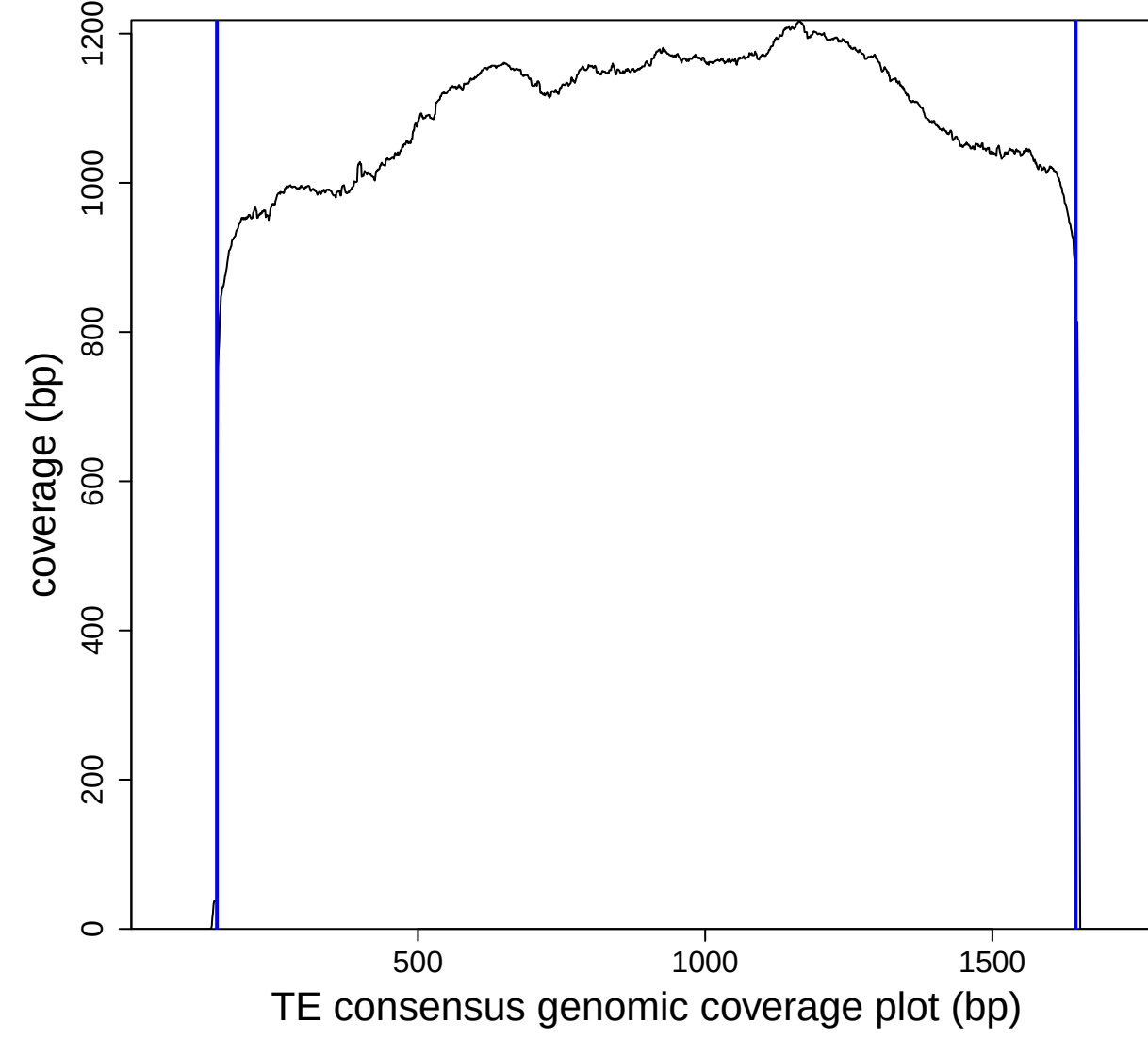
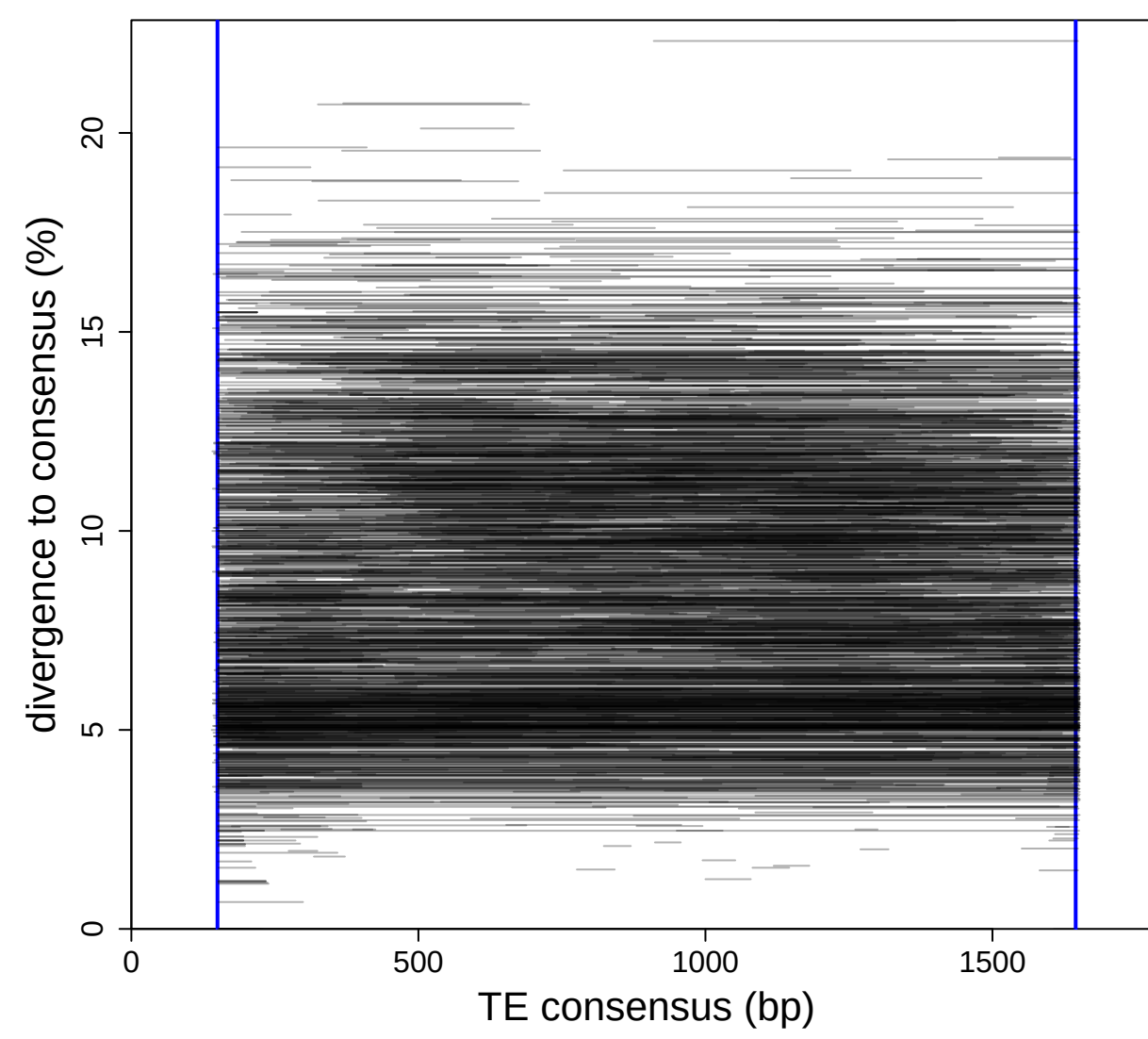
l.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcIn.fa aln.fa cl.fa gs.fa
size: 1495bp; fragments: 2548; full length: 379 (≥ 1345.5 bp)

After TEtrimmer 1495 bp

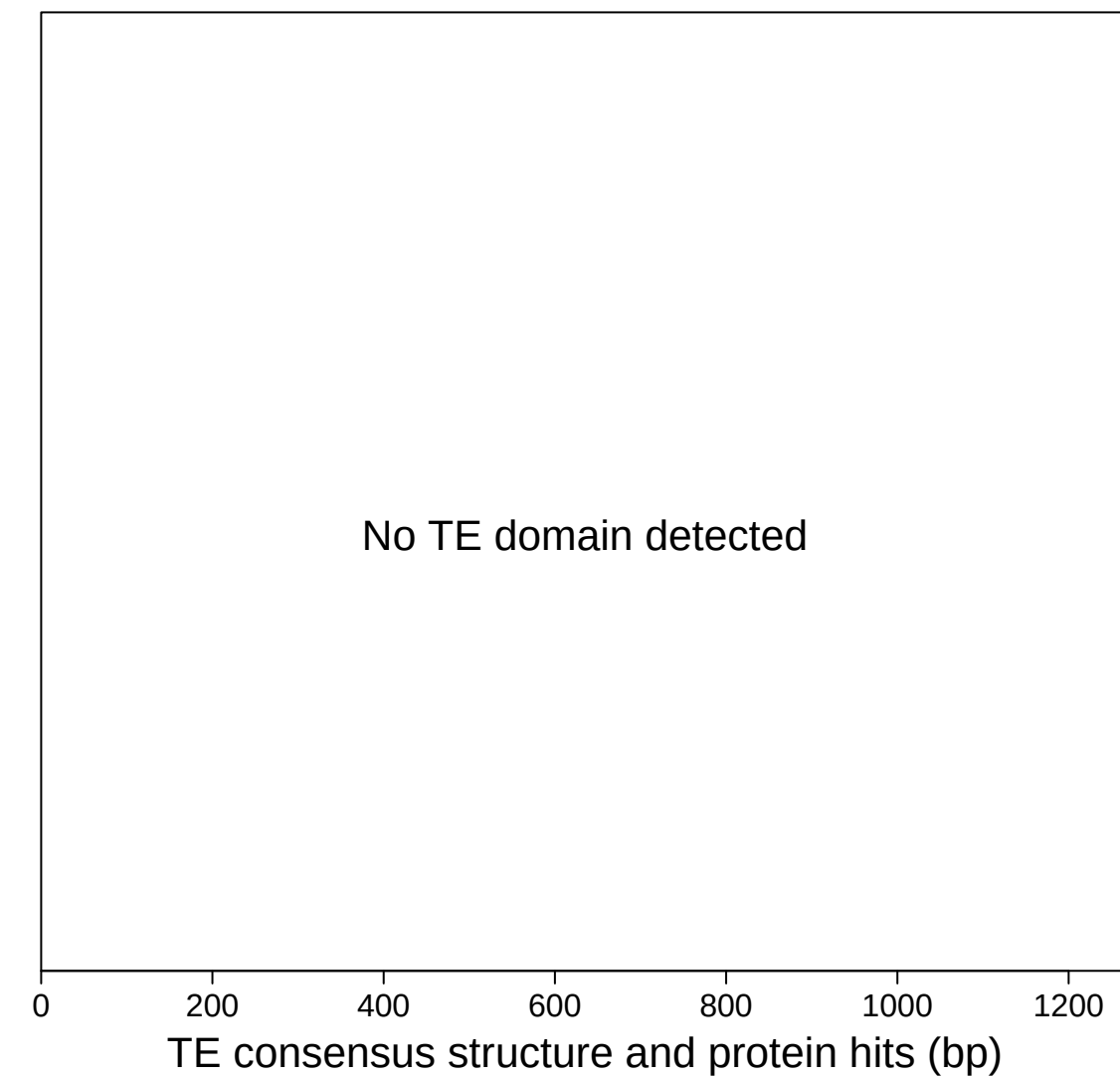
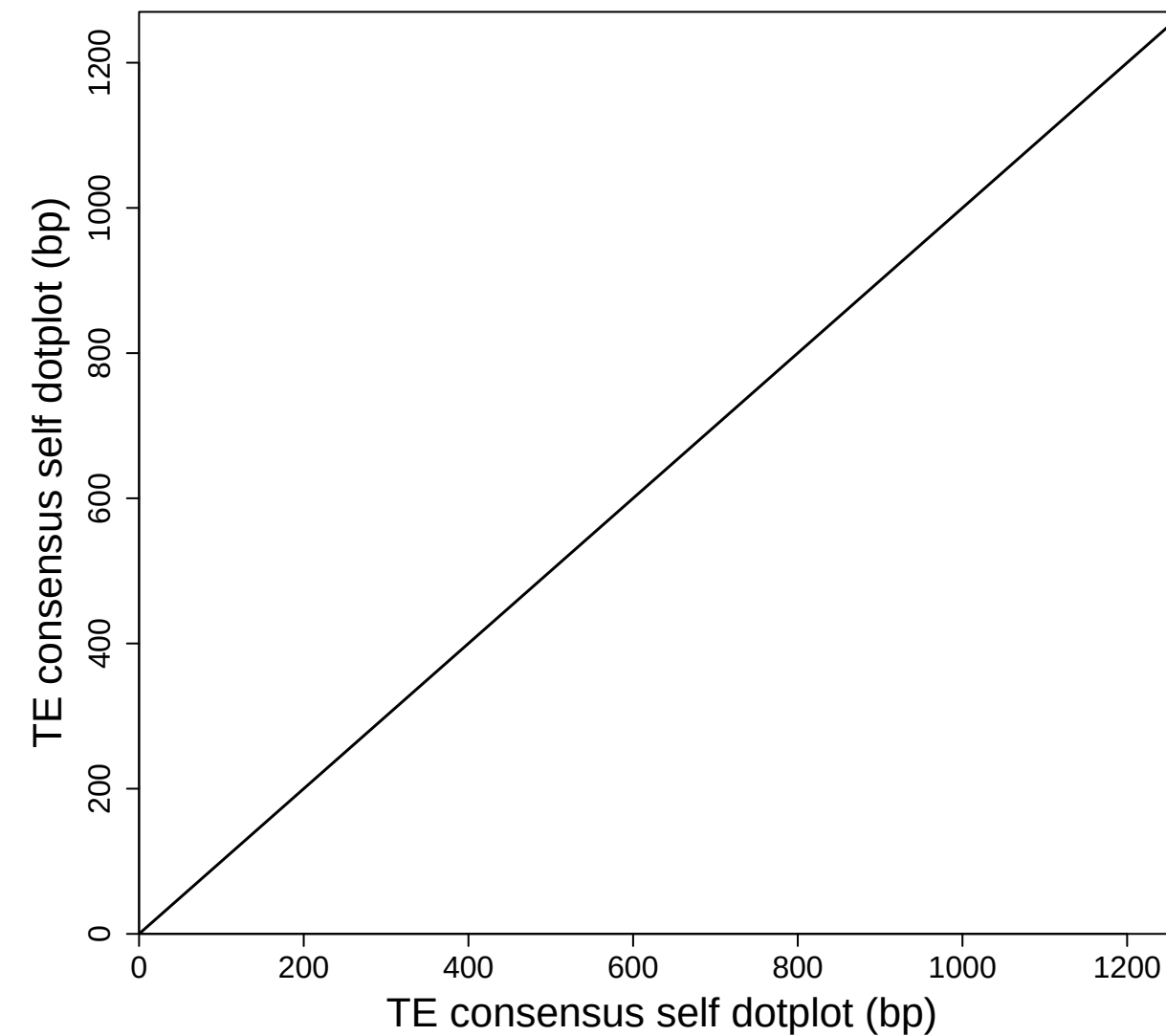
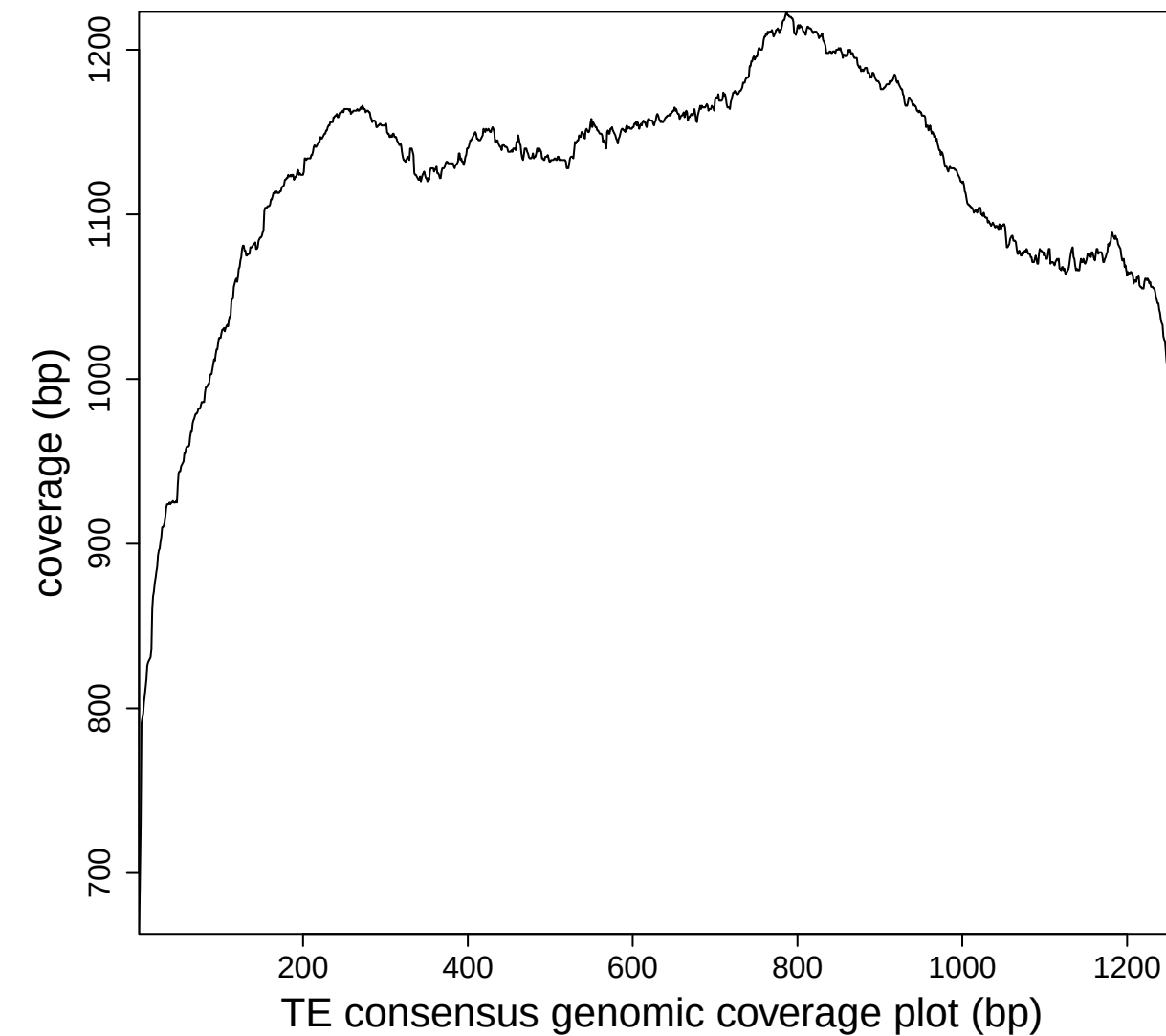
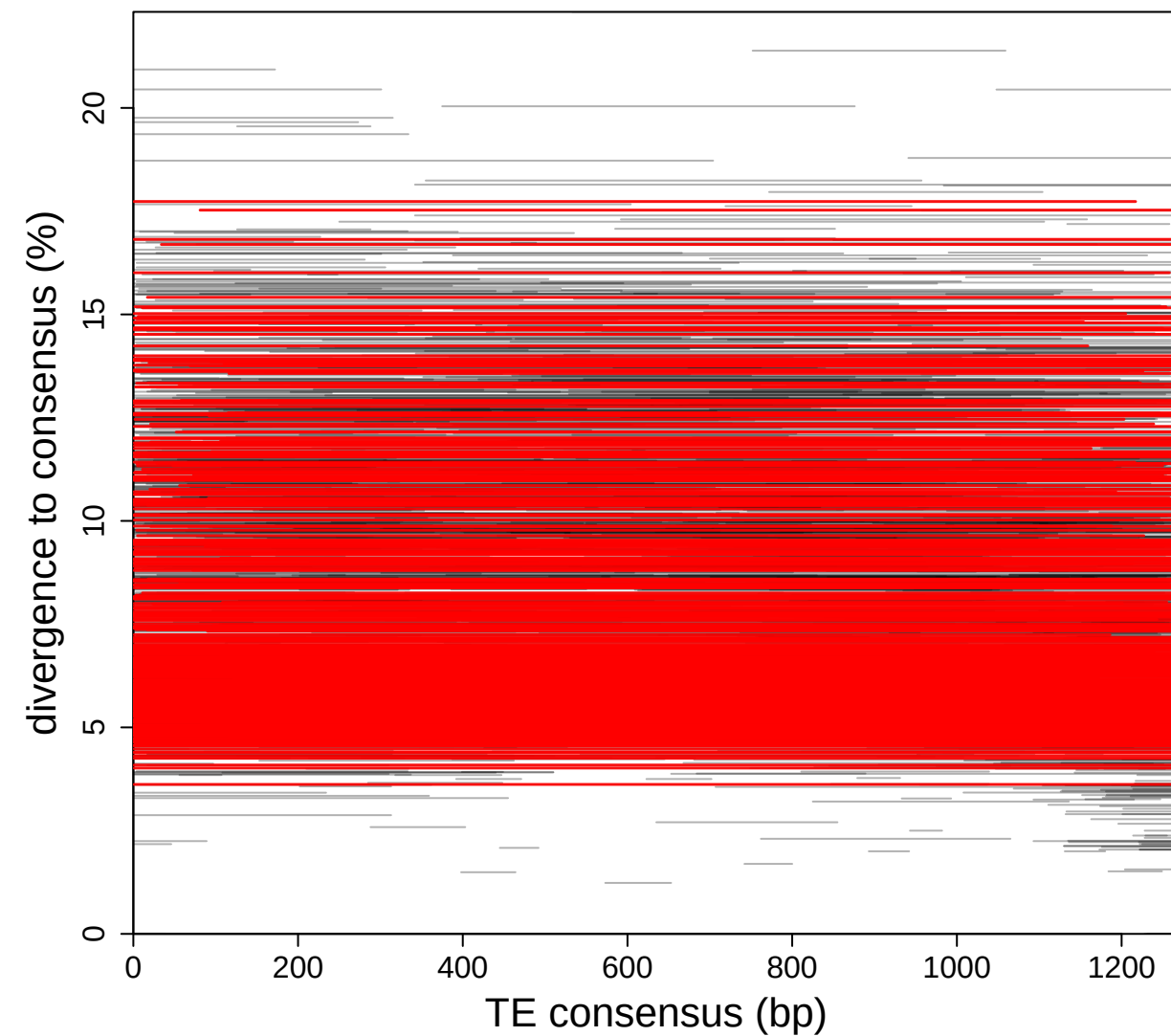


0.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 1795bp; fragments: 2549; full length: 0 (>=1615.5bp)

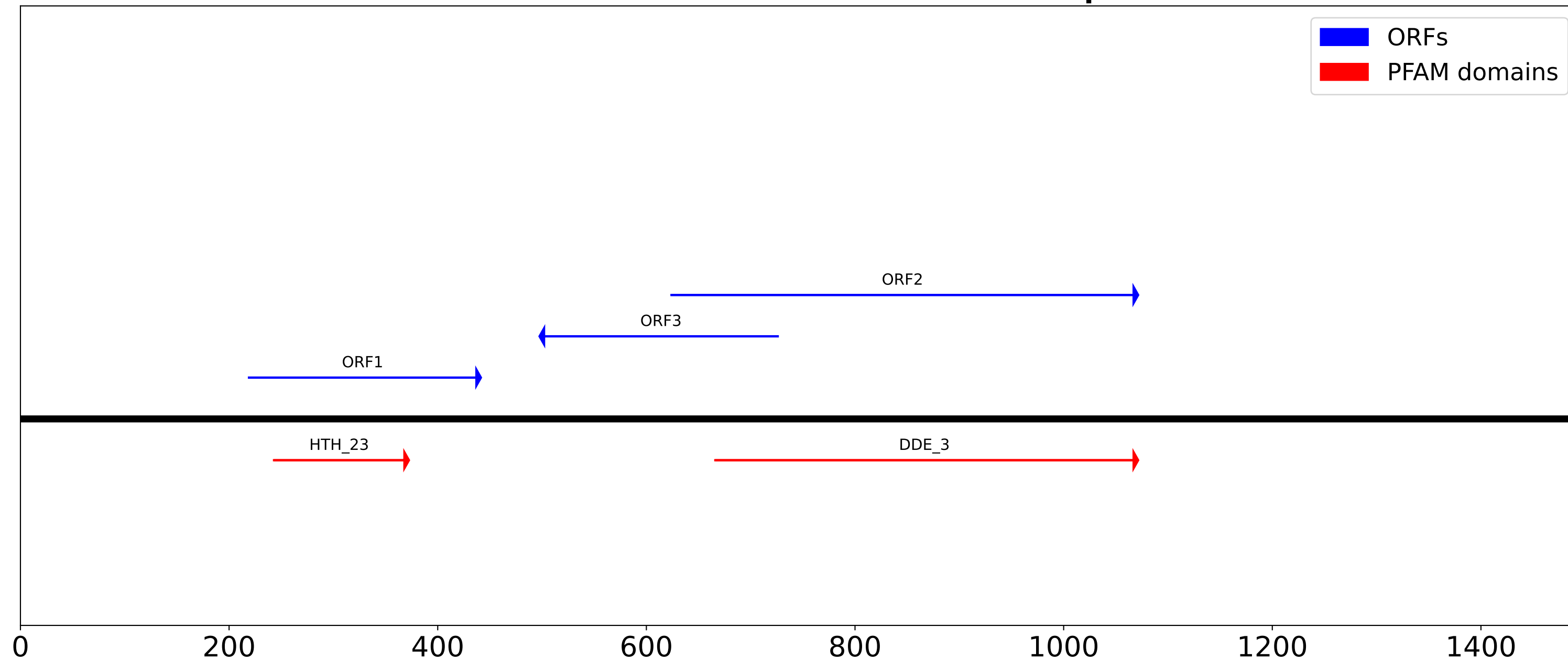
After TEtrimmer Extended plot Blue lines are boundaries



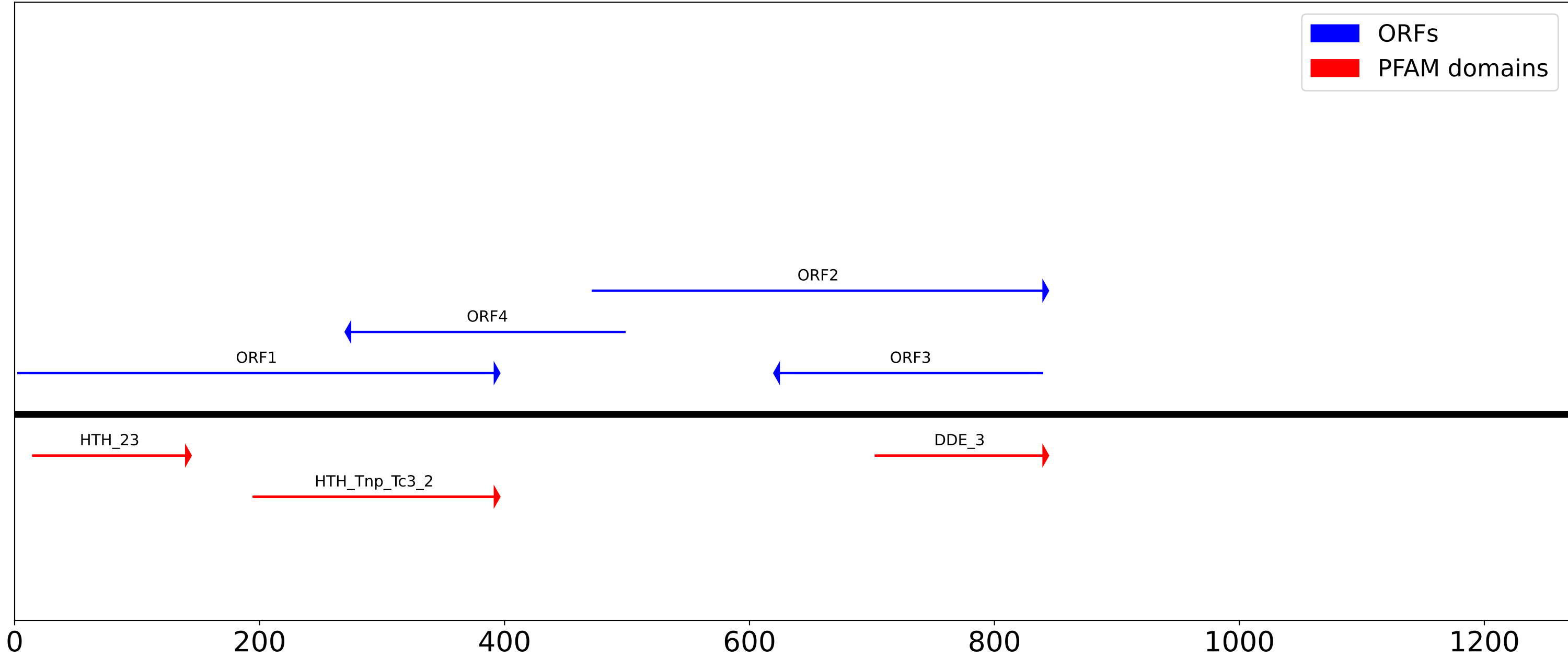
TE: rnd_1_family_740
size: 1270bp; fragments: 2263; full length: 467 (≥ 1143 bp)



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

