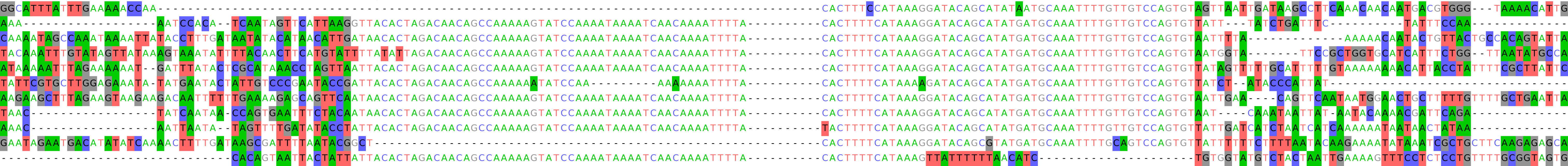
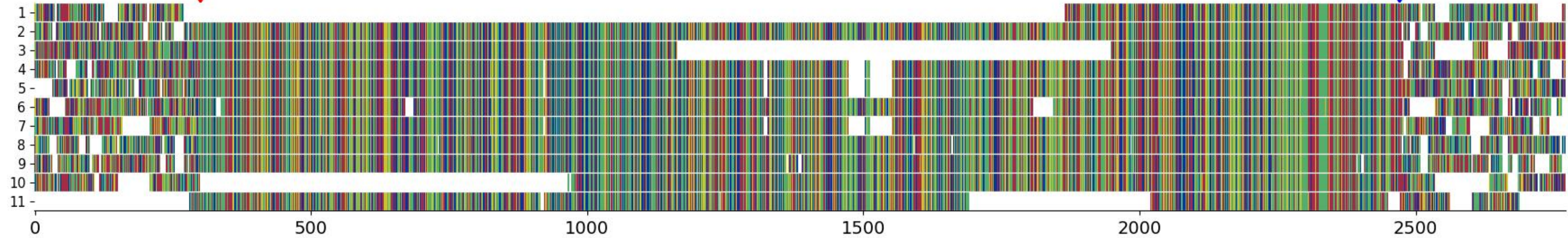


A length



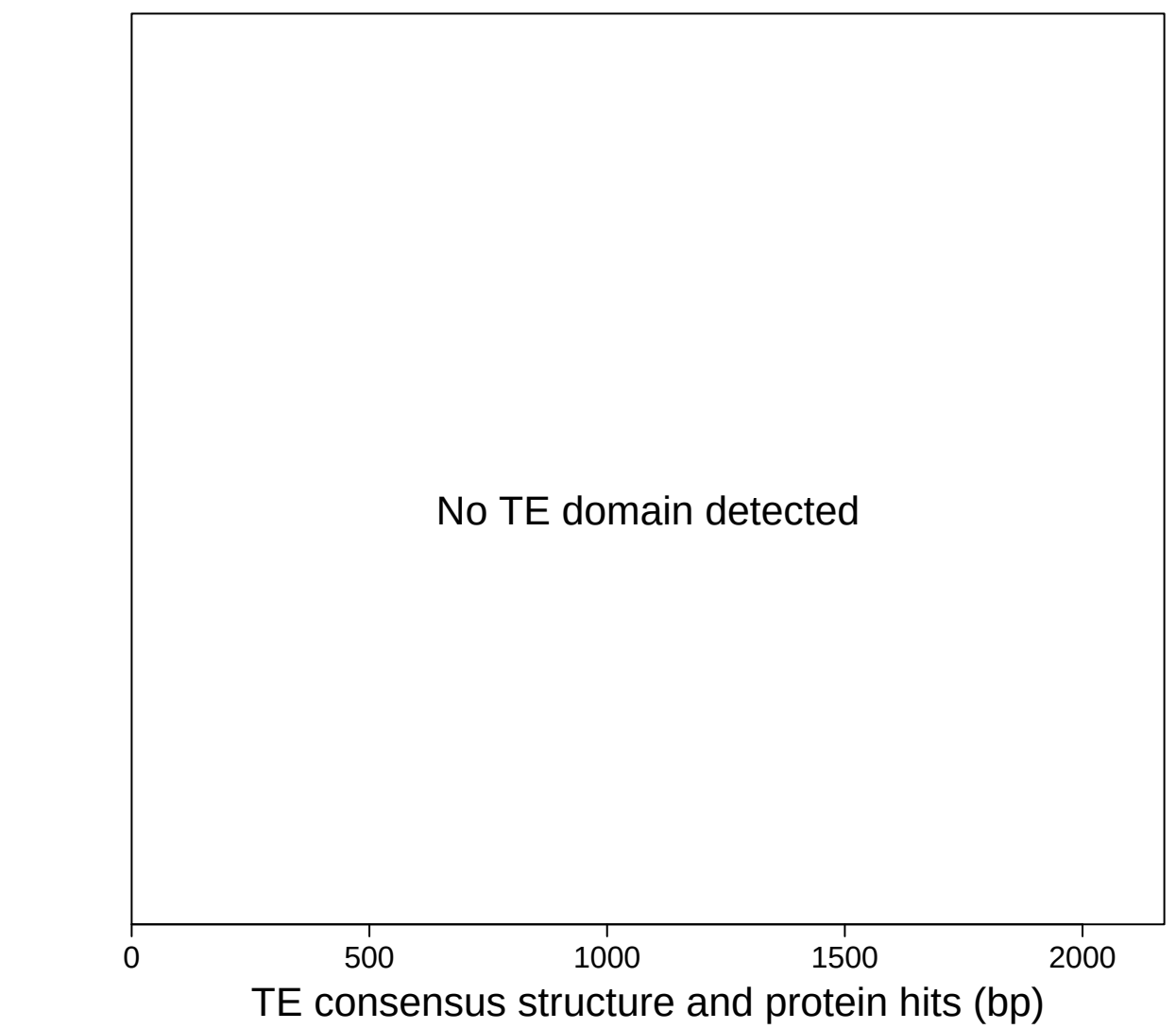
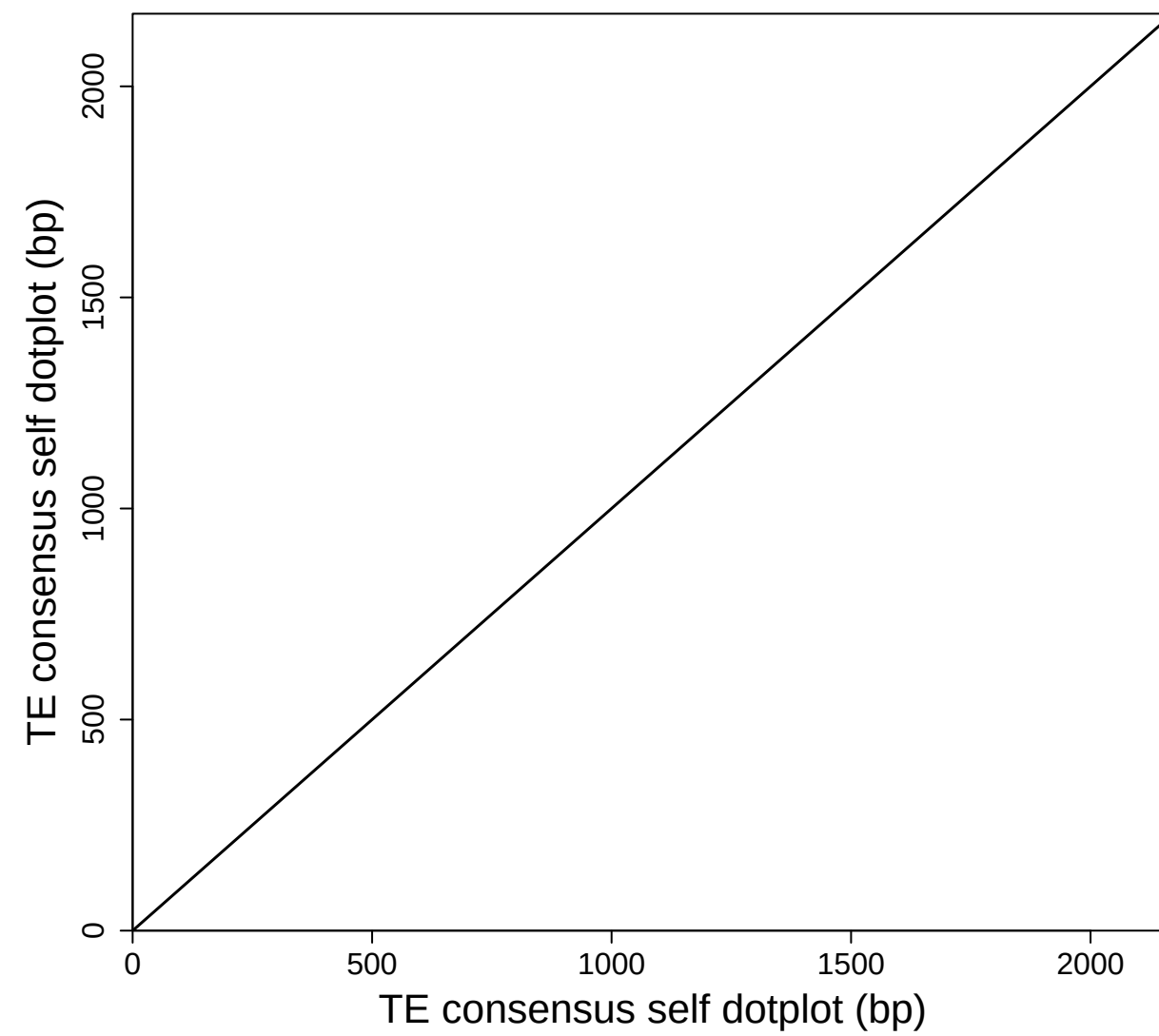
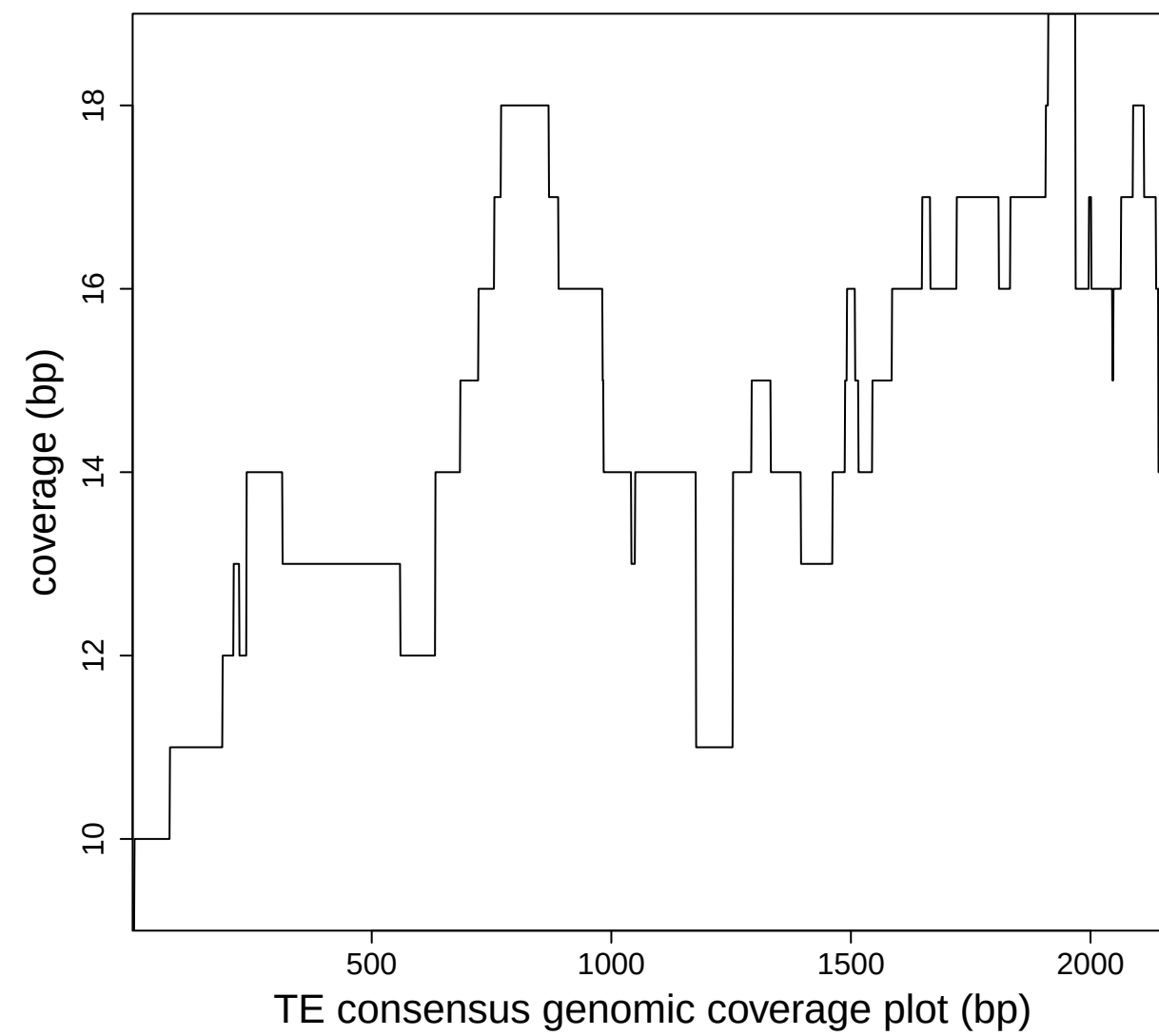
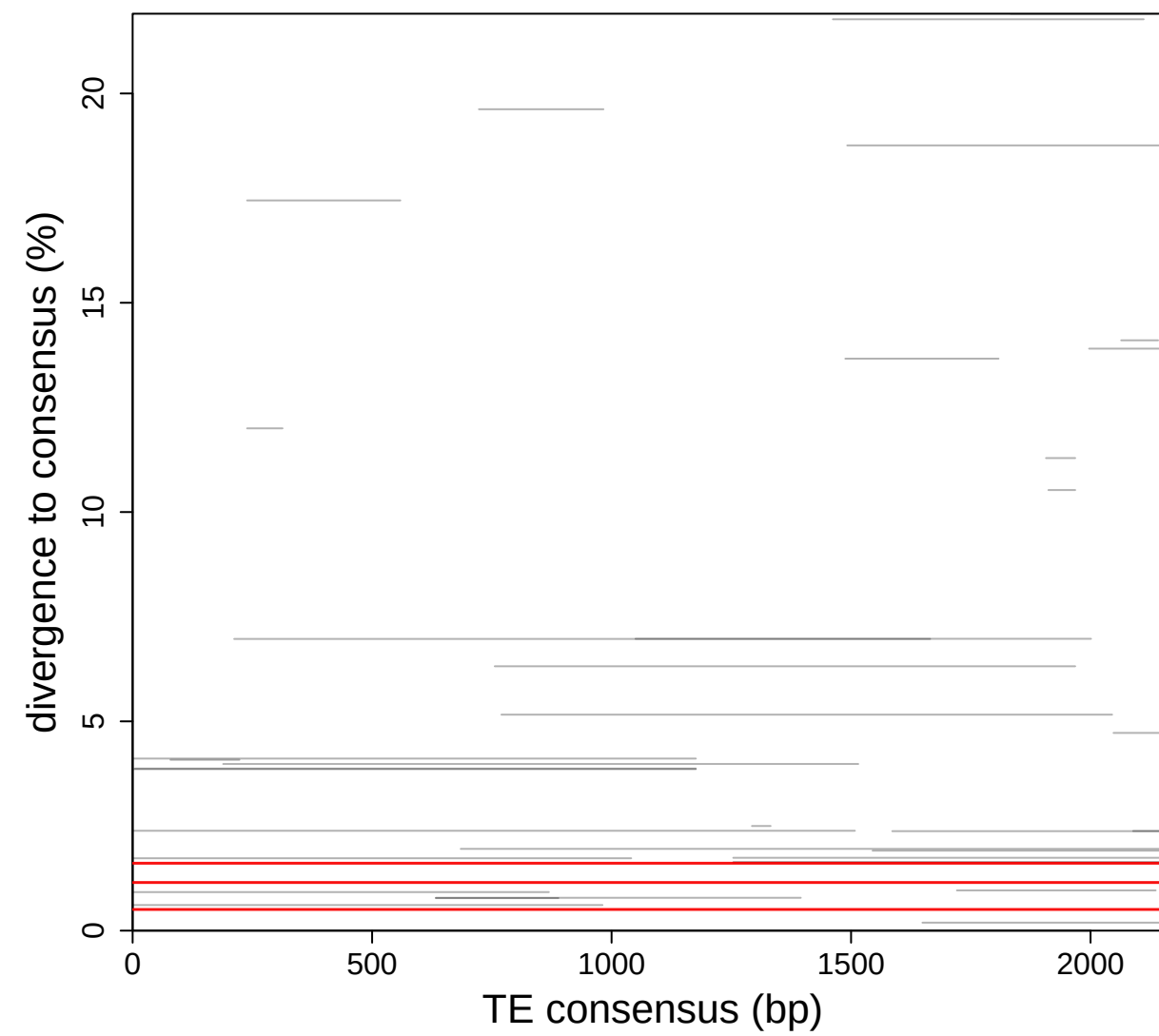
Start crop Point

End crop Point



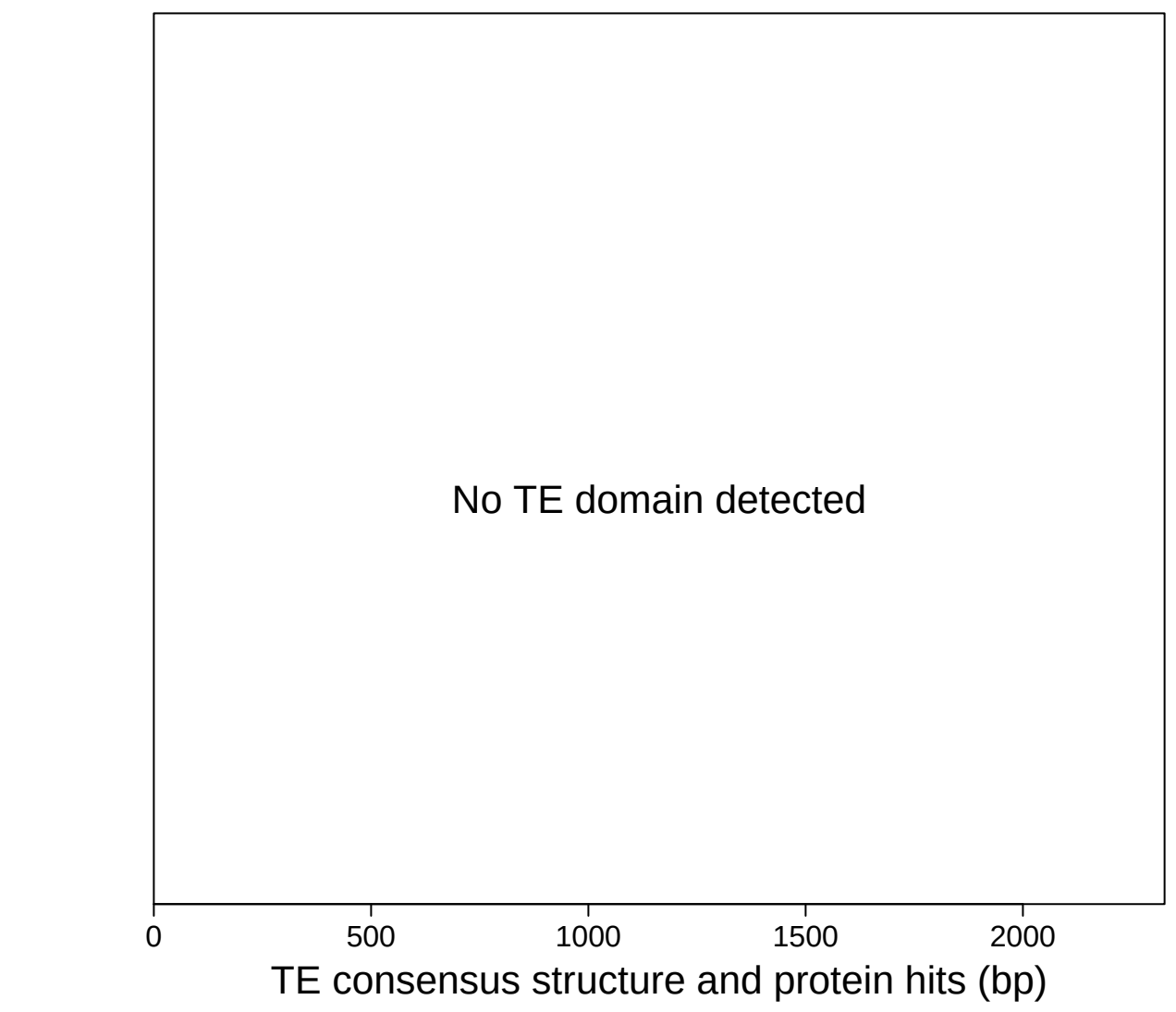
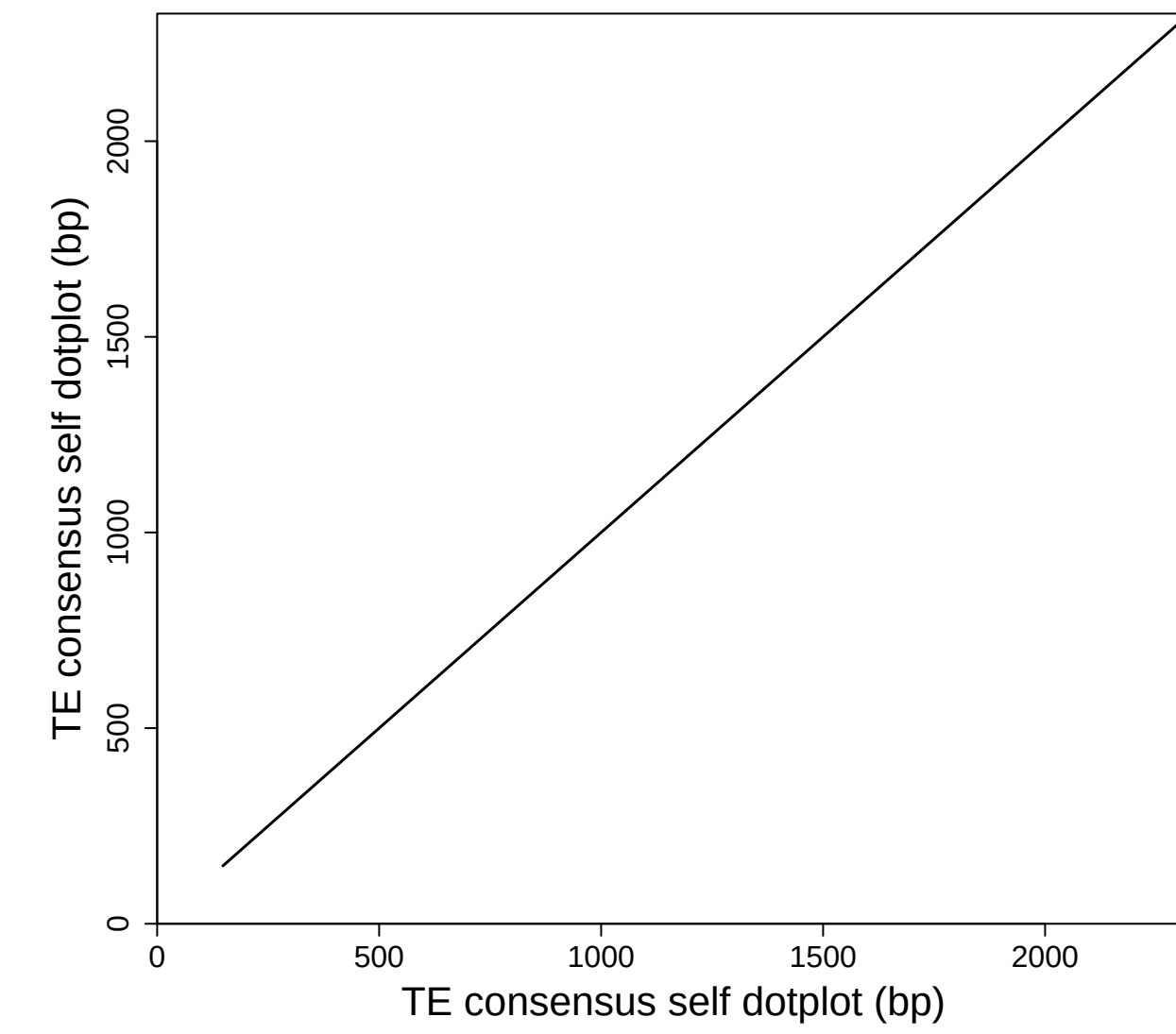
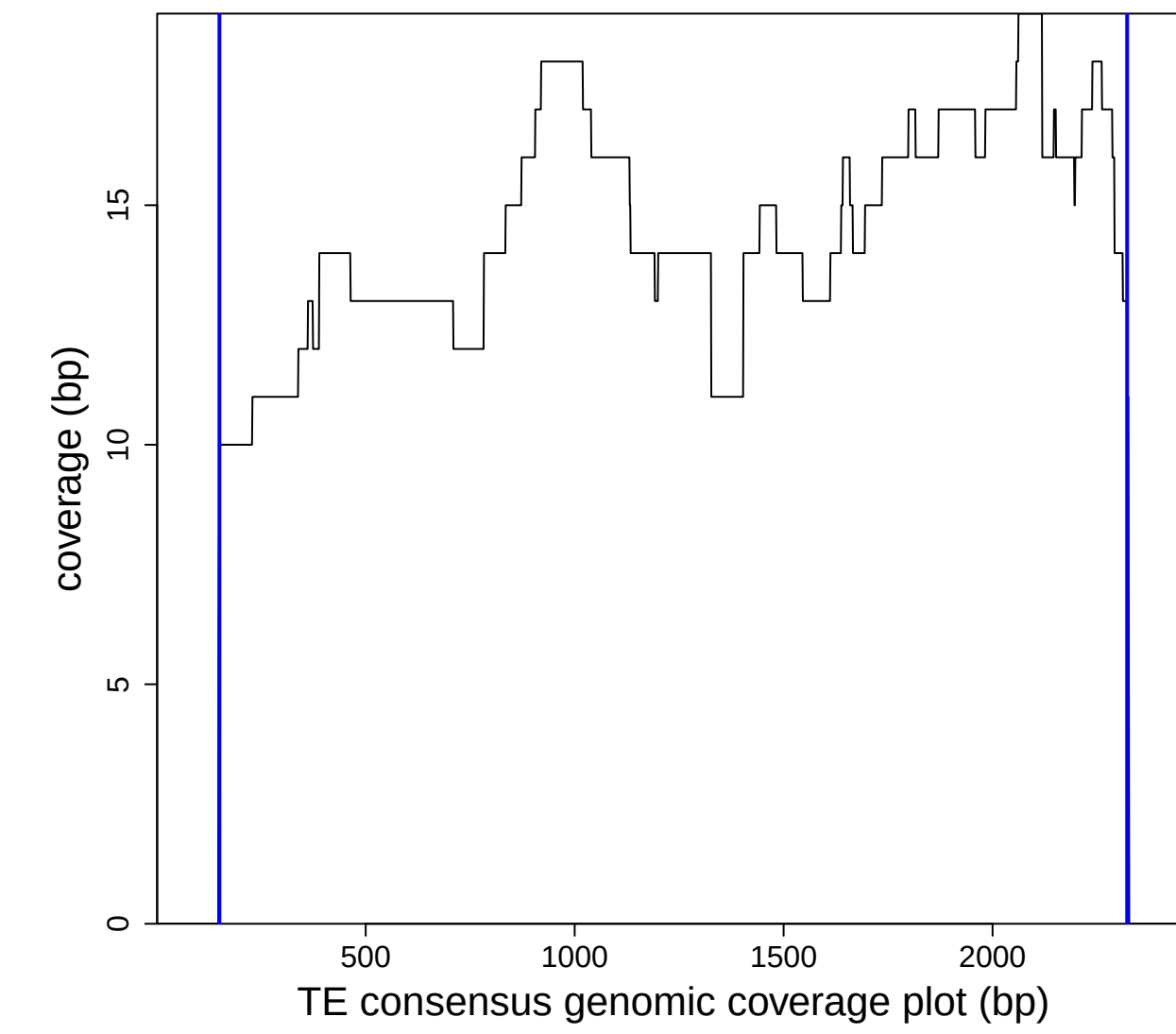
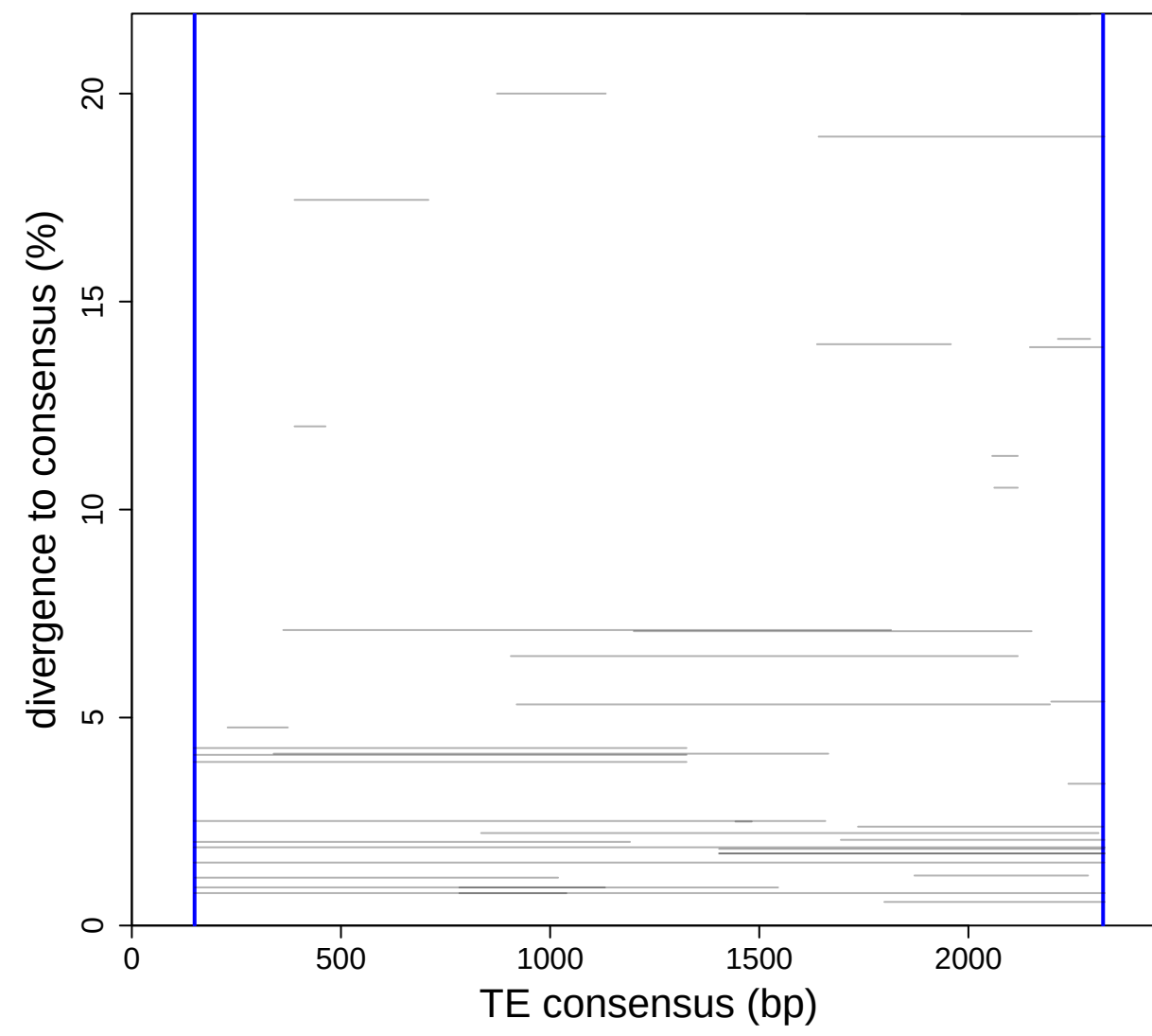
l.b.bed uf.bed g_1.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa
size: 2172bp; fragments: 40; full length: 3 (>=1954.8bp)

After TEtrimmer 2172 bp

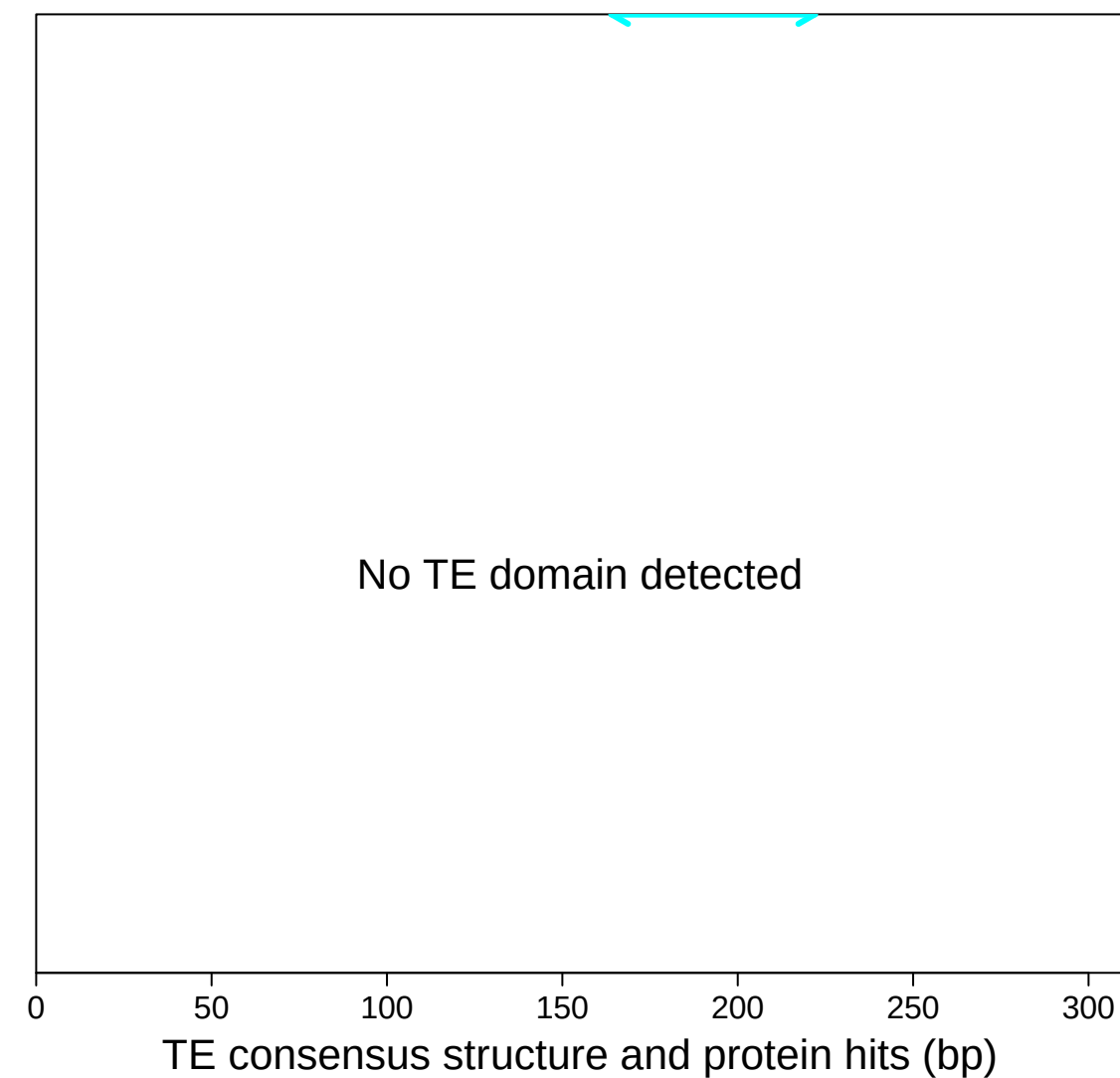
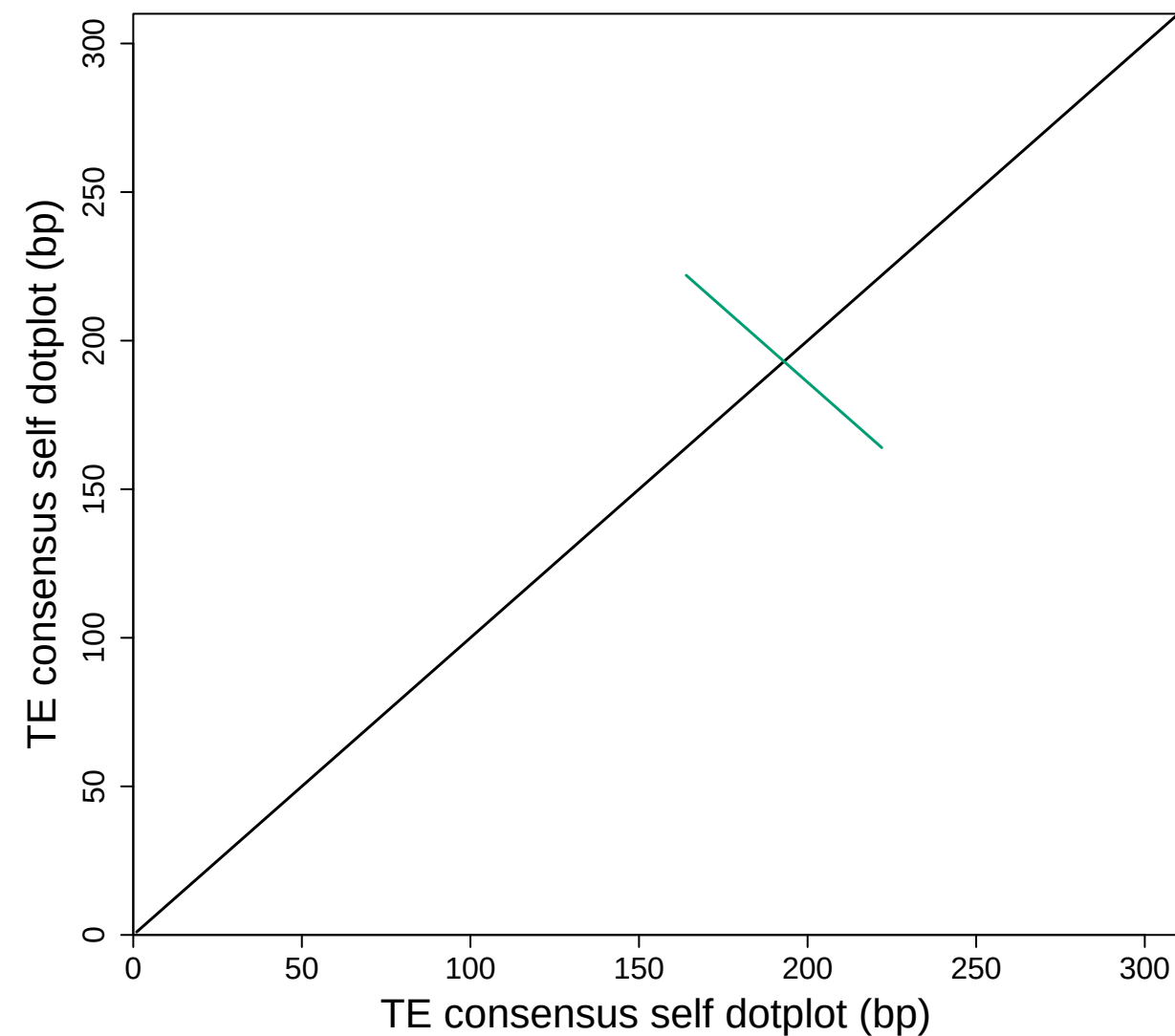
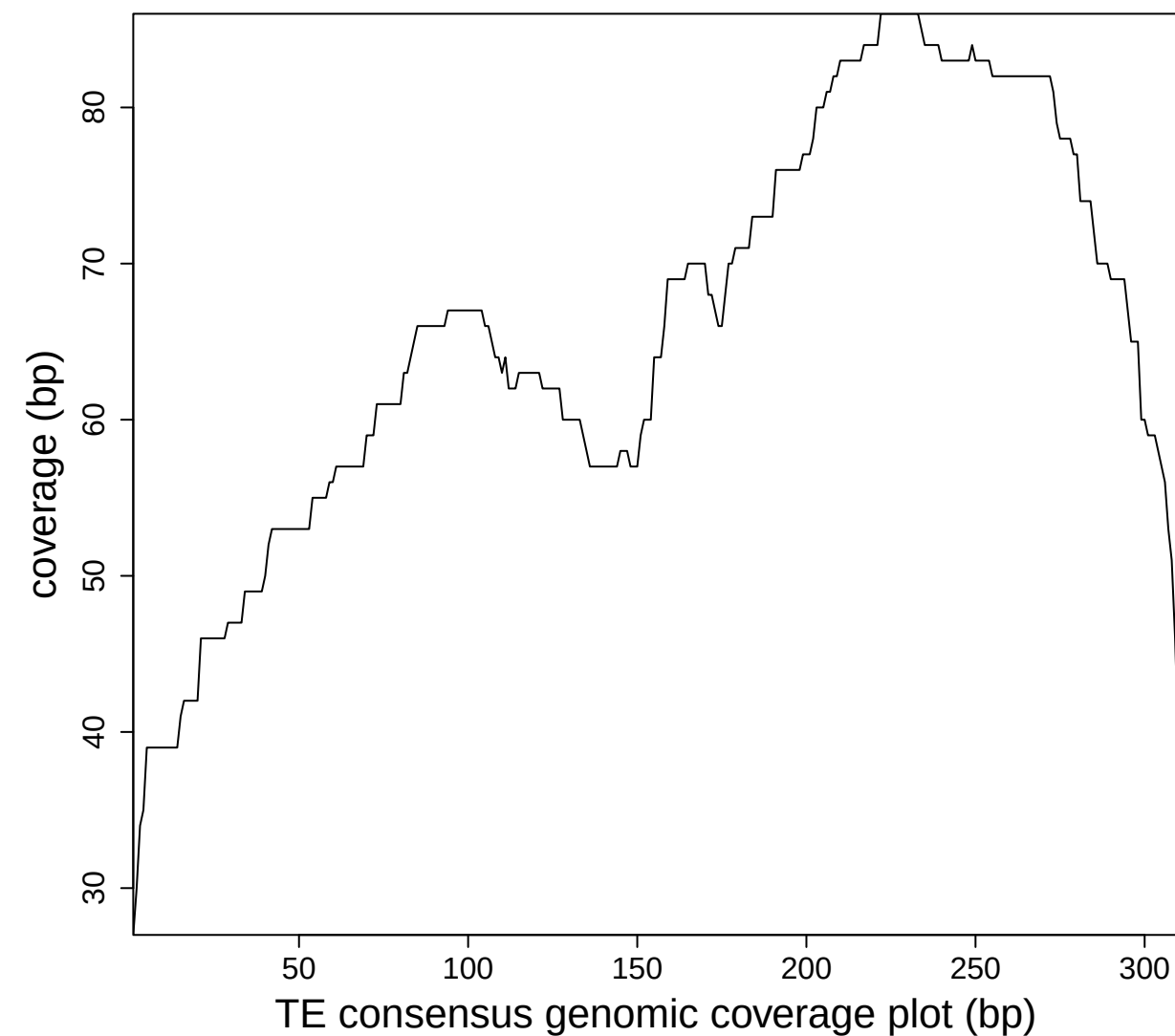
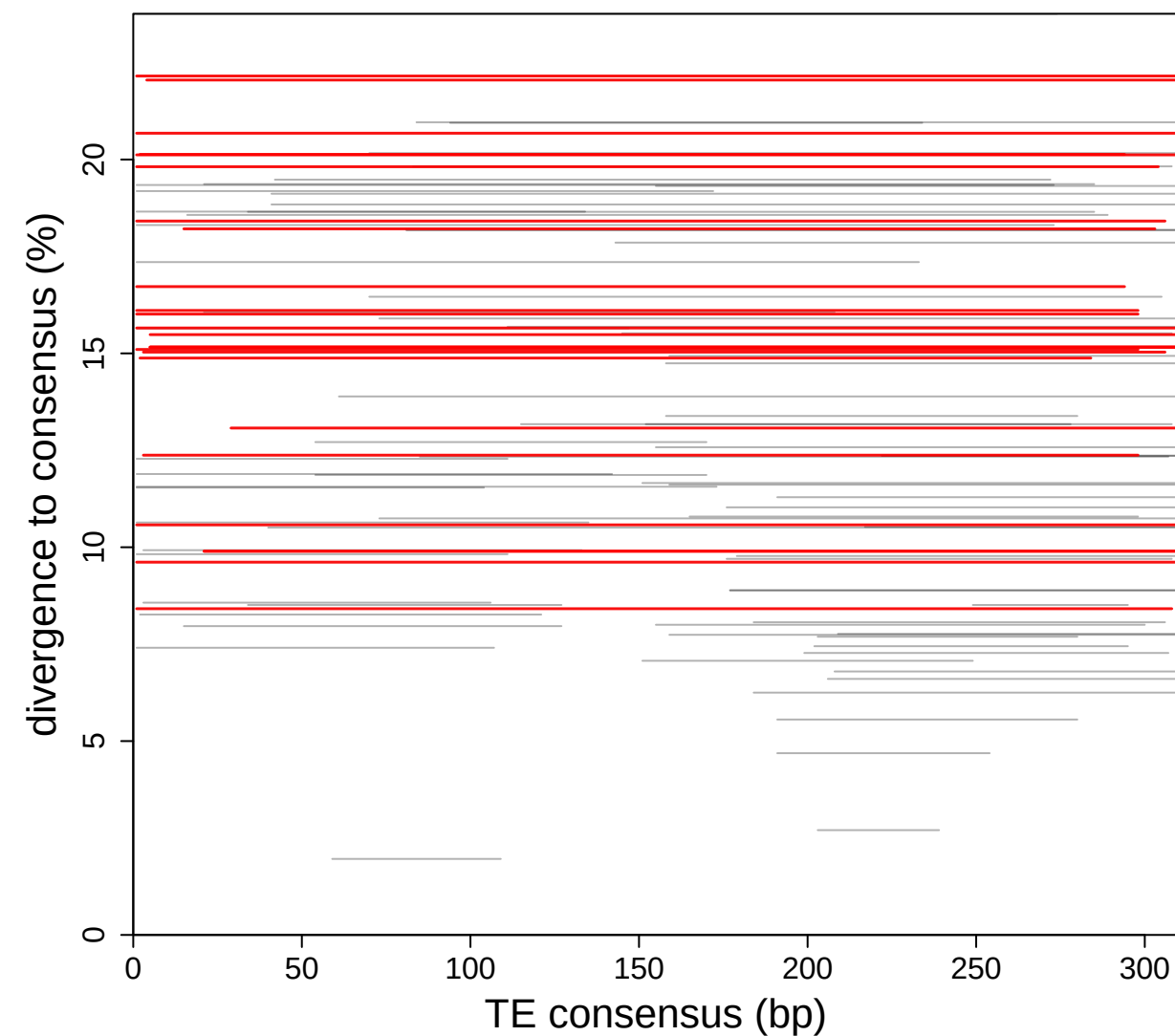


4.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 2472bp; fragments: 40; full length: 0 (>=2224.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

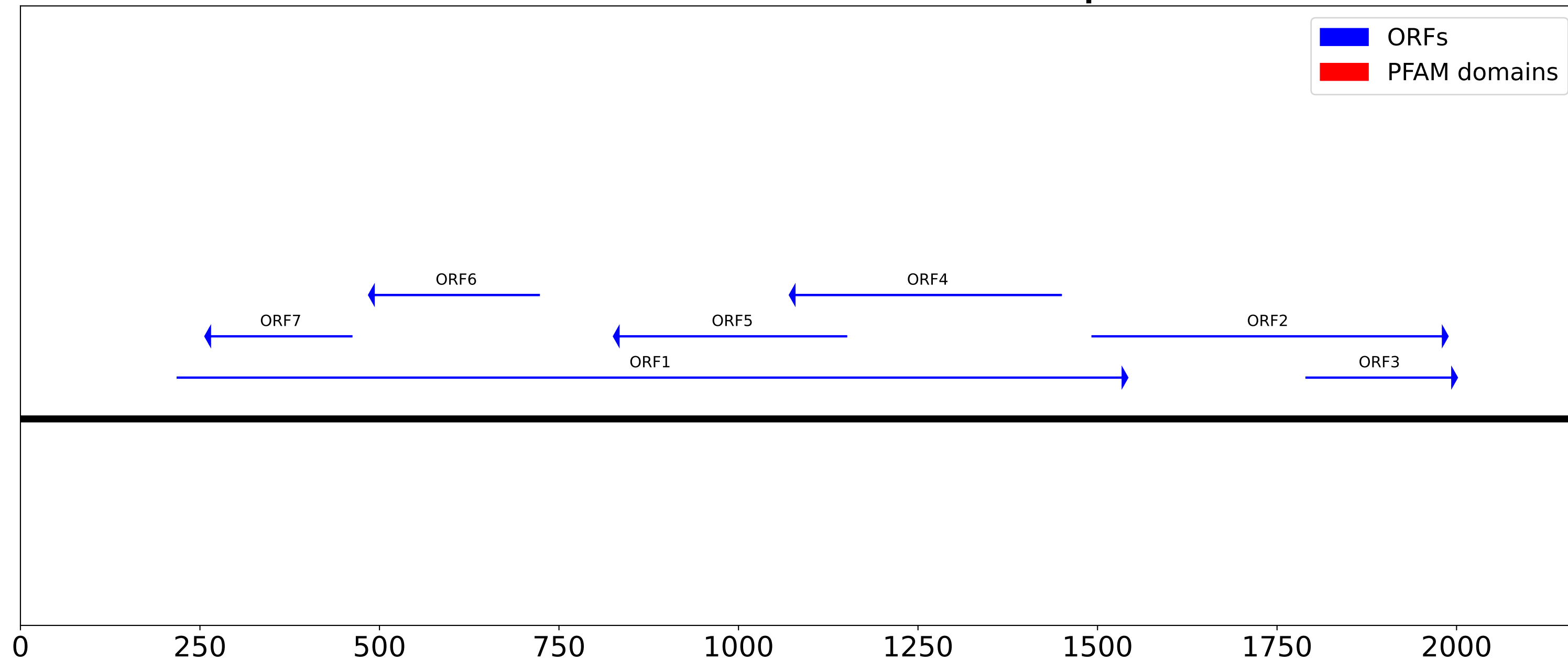


TE: rnd_5_family_574
size: 310bp; fragments: 106; full length: 26 (≥ 279 bp)



Before TEtrimmer 310 bp

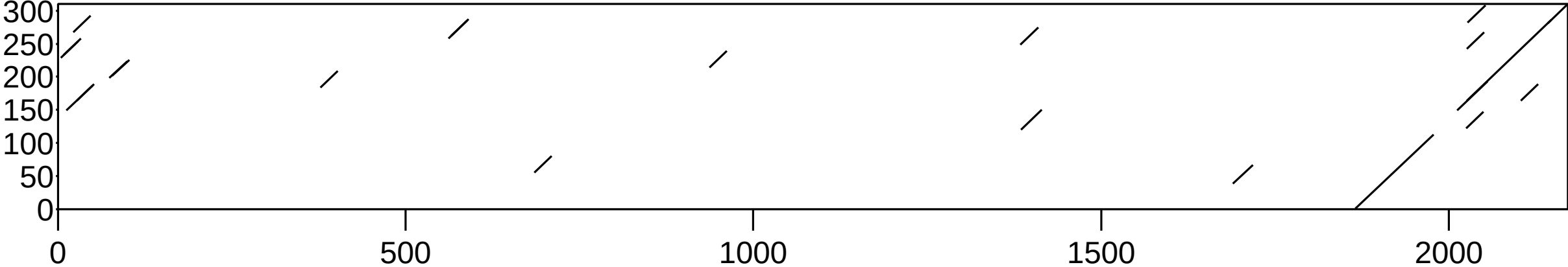
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

isensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)