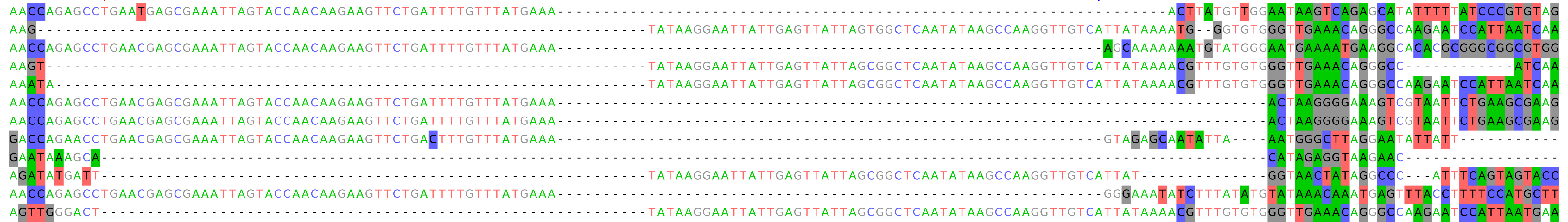
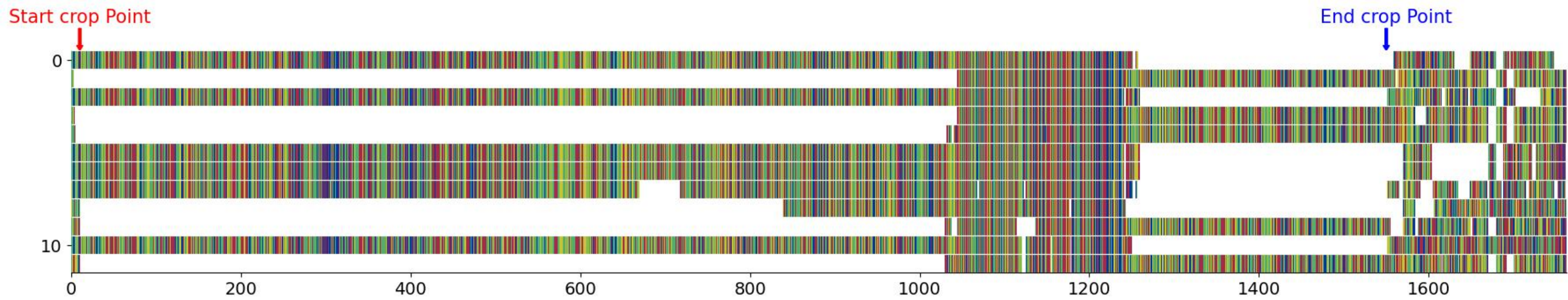


Start crop Point

End crop Point

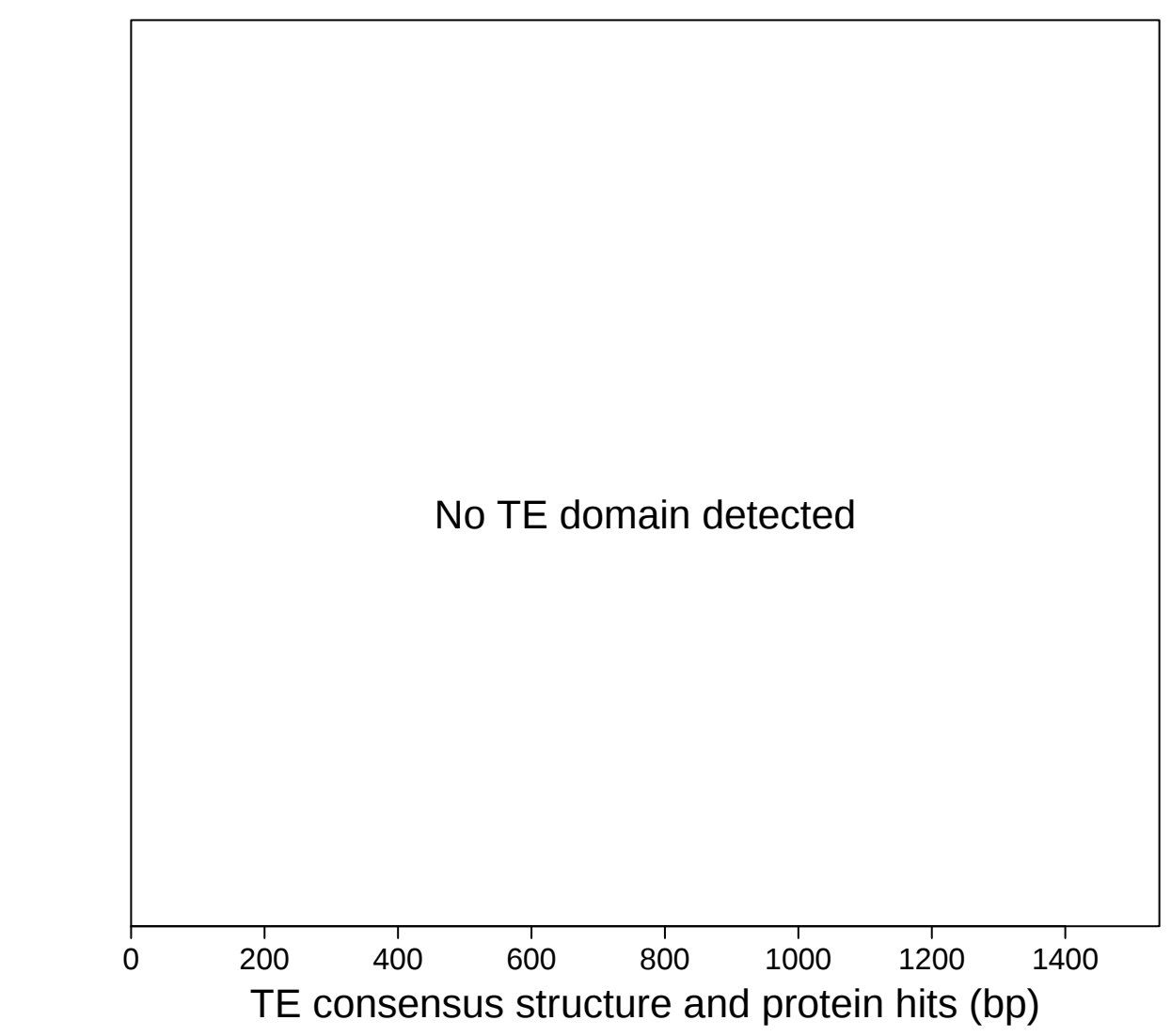
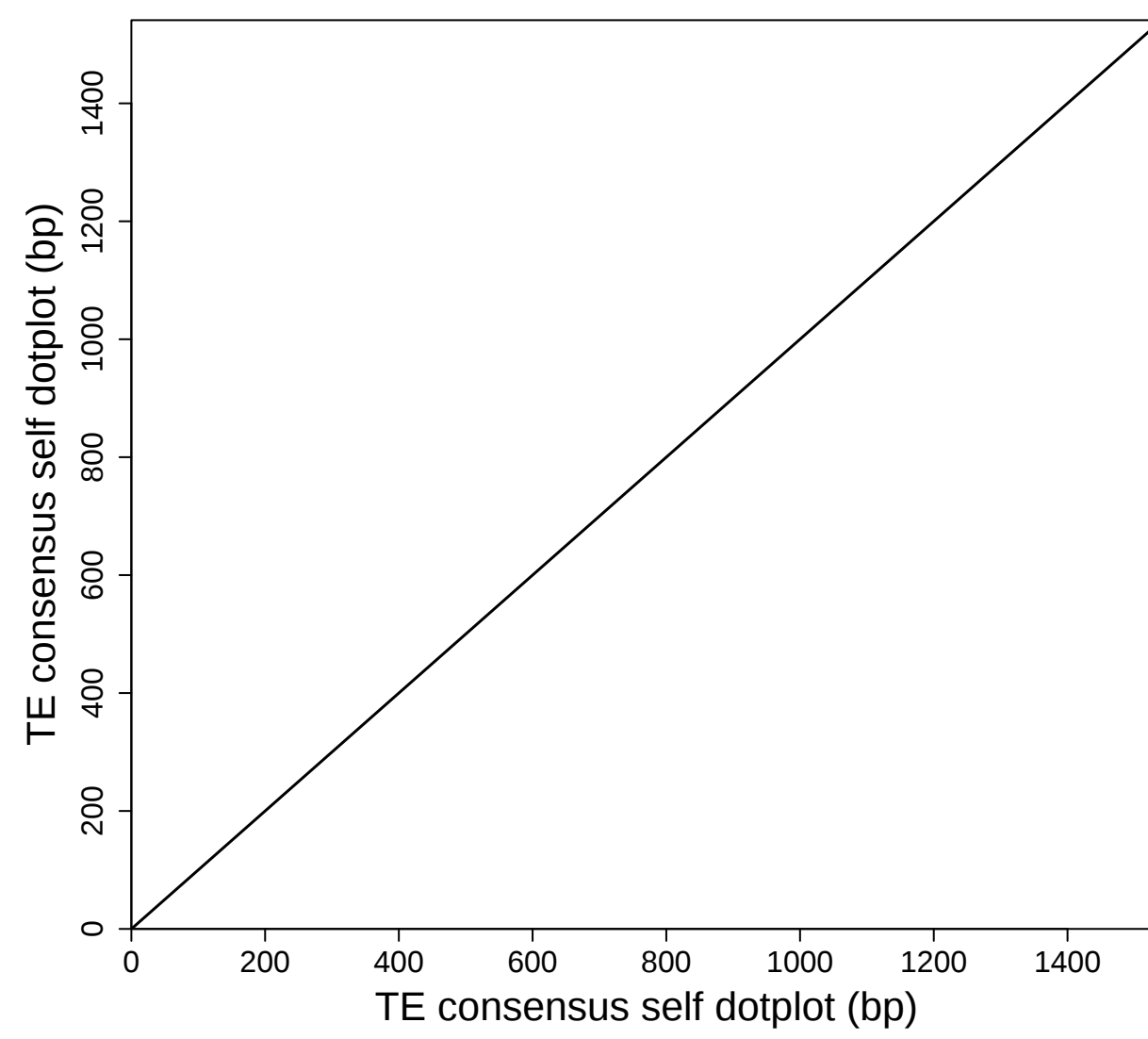
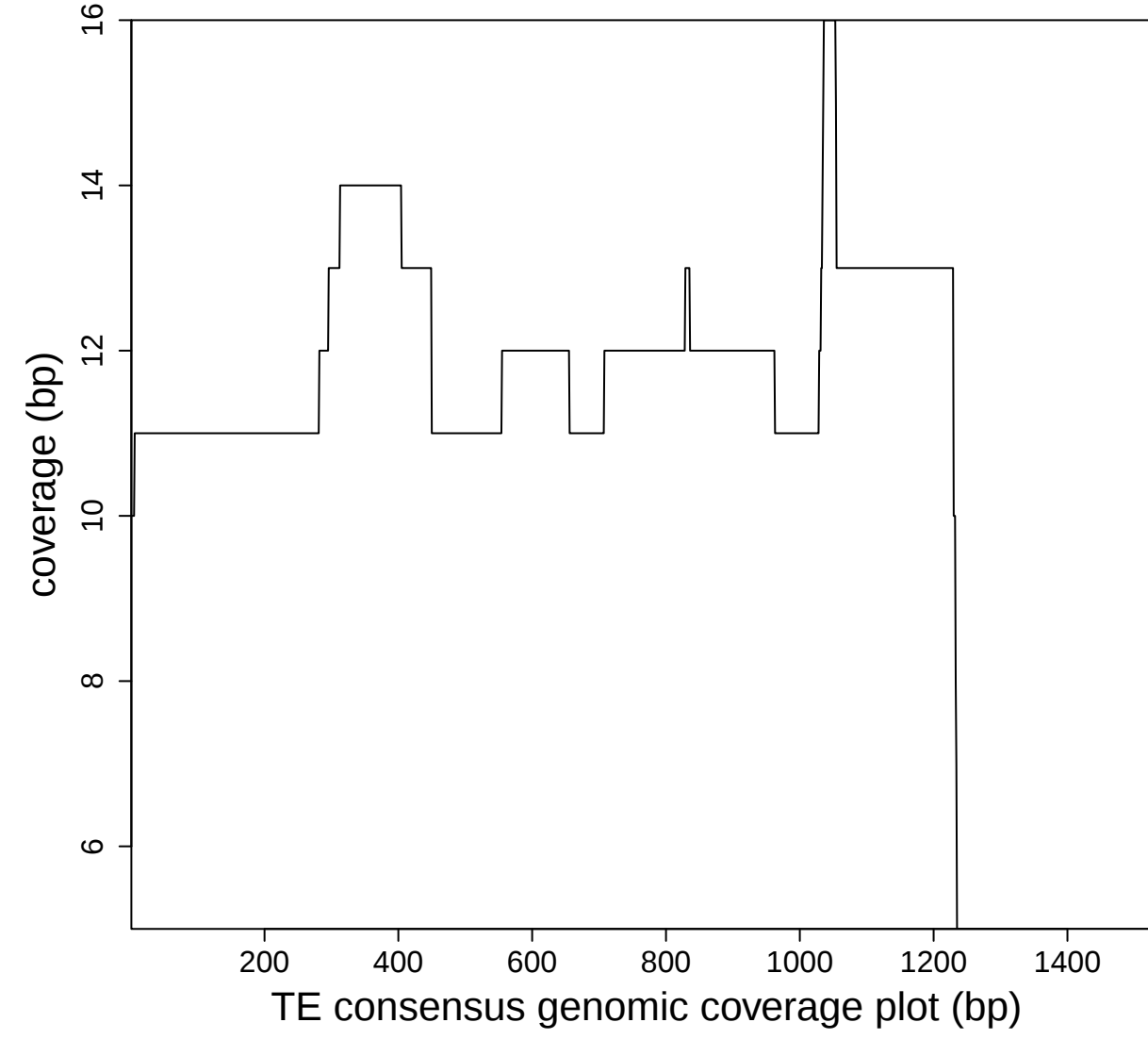
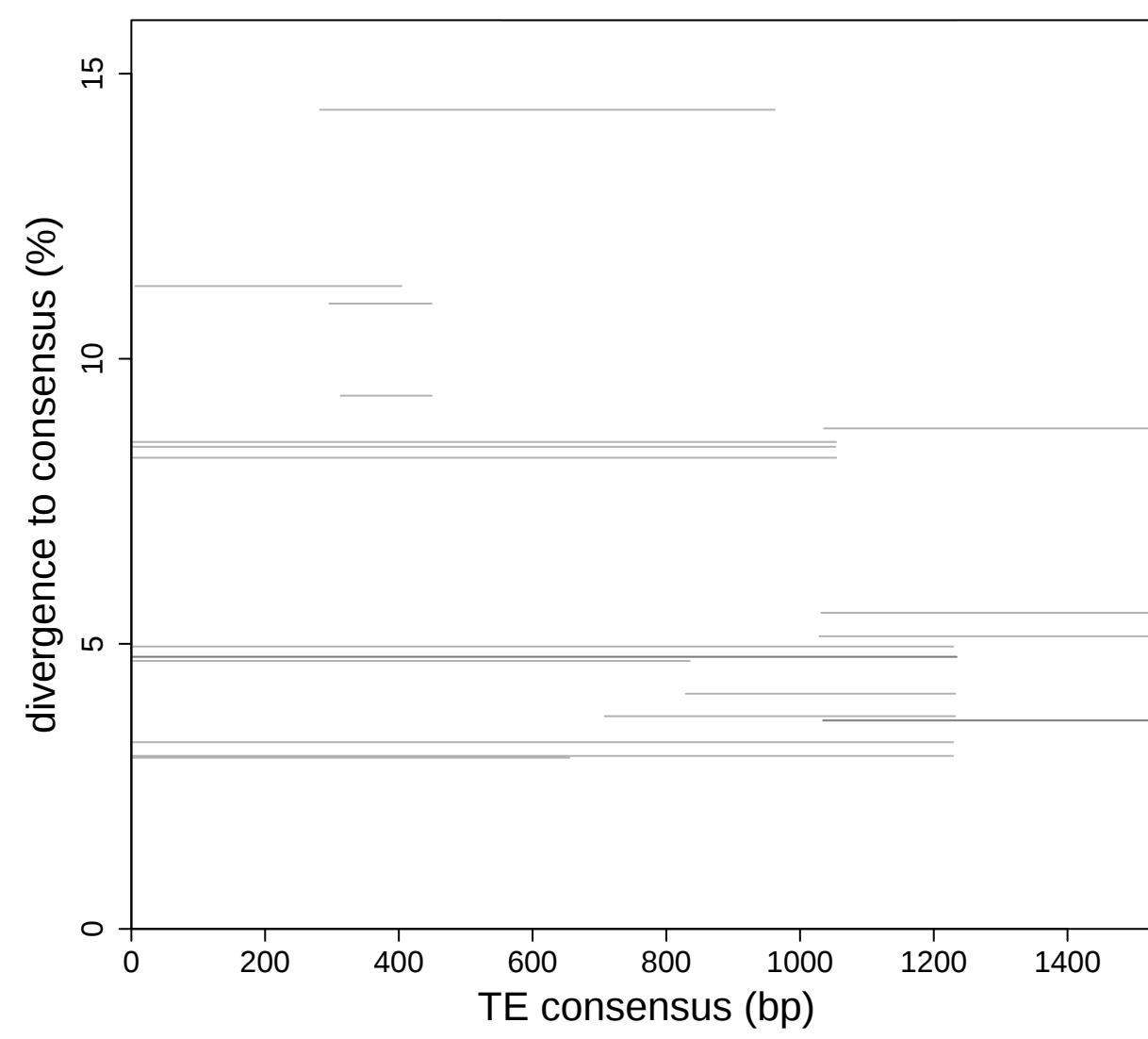
MSA length = 1541





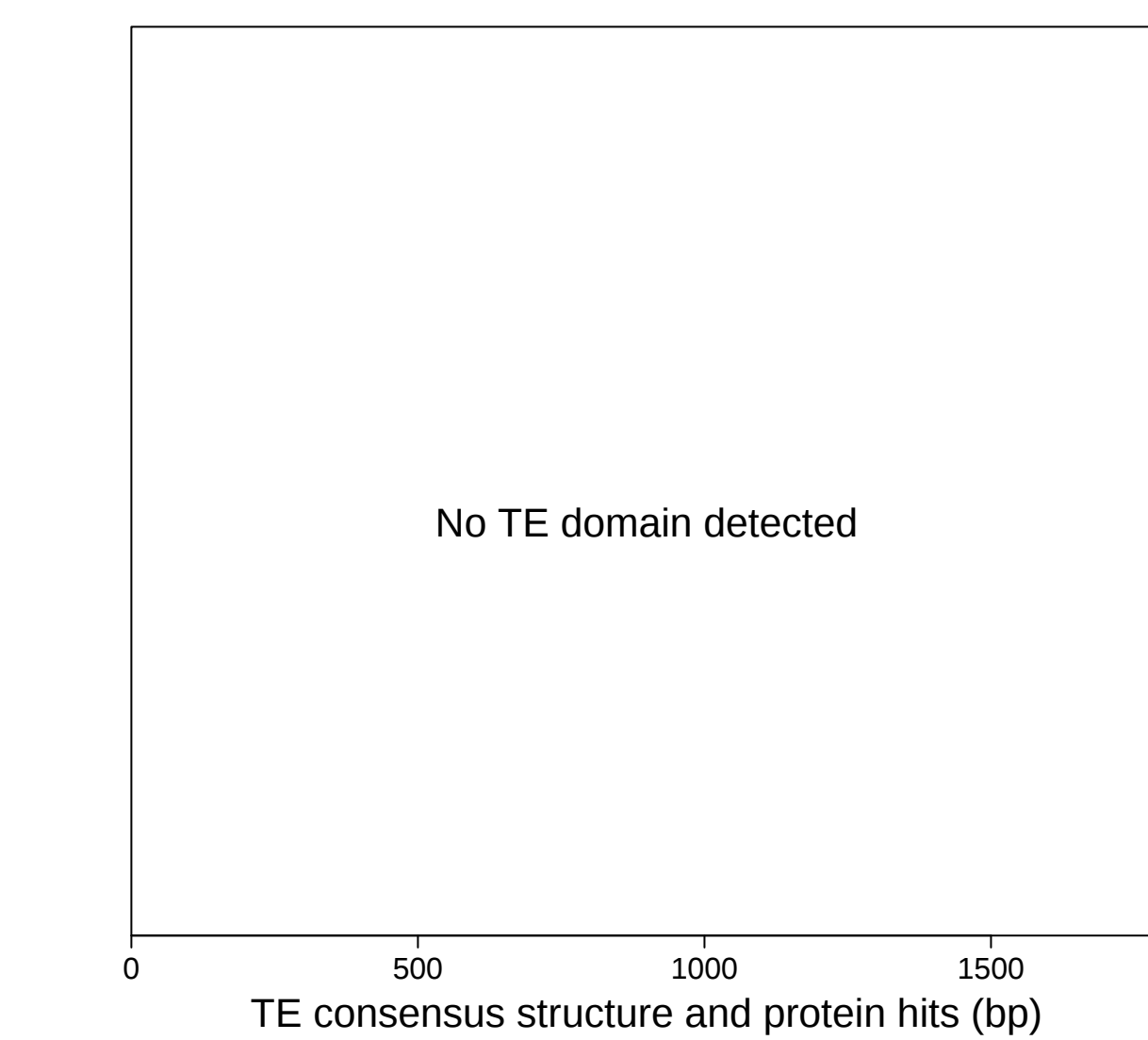
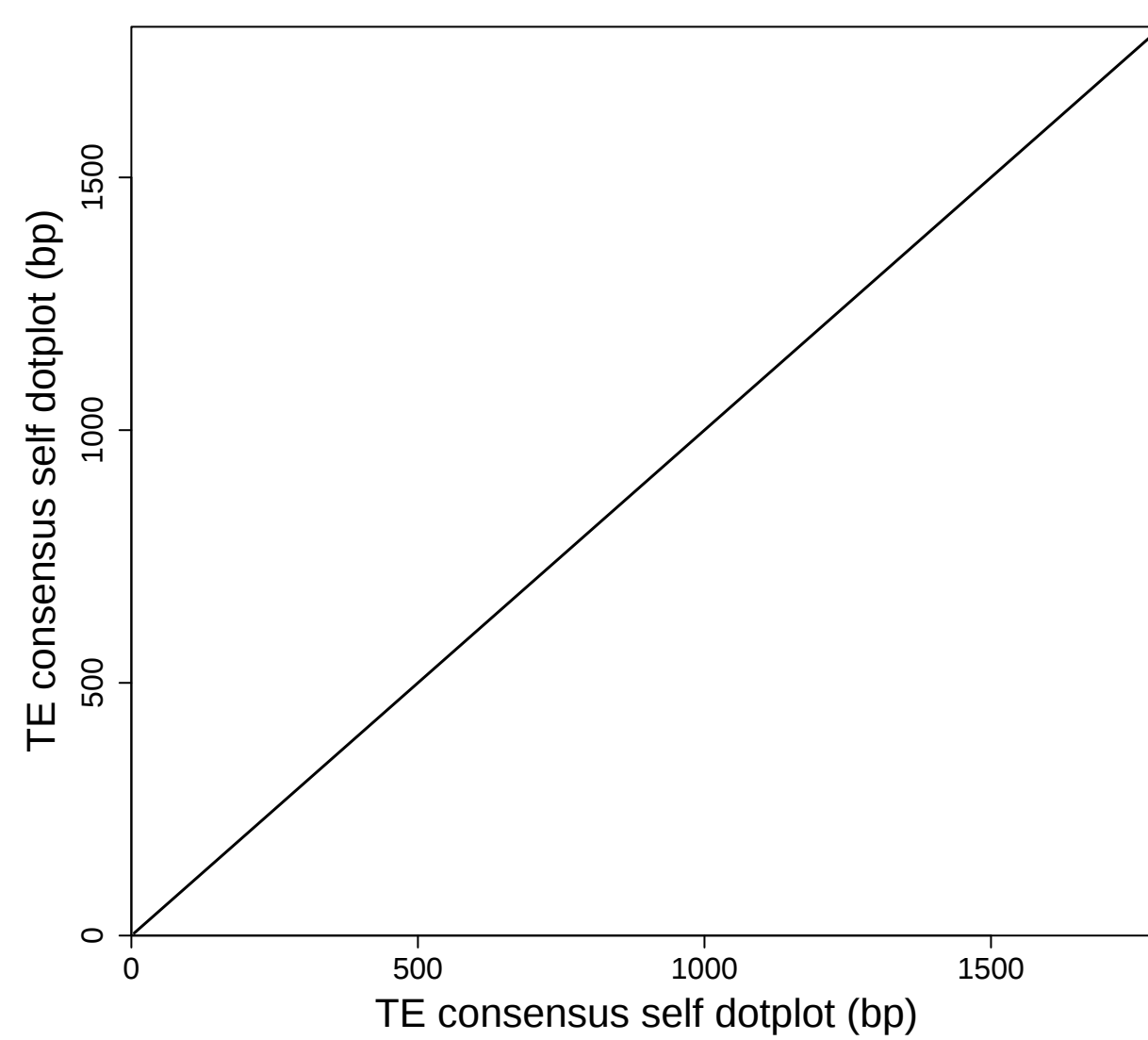
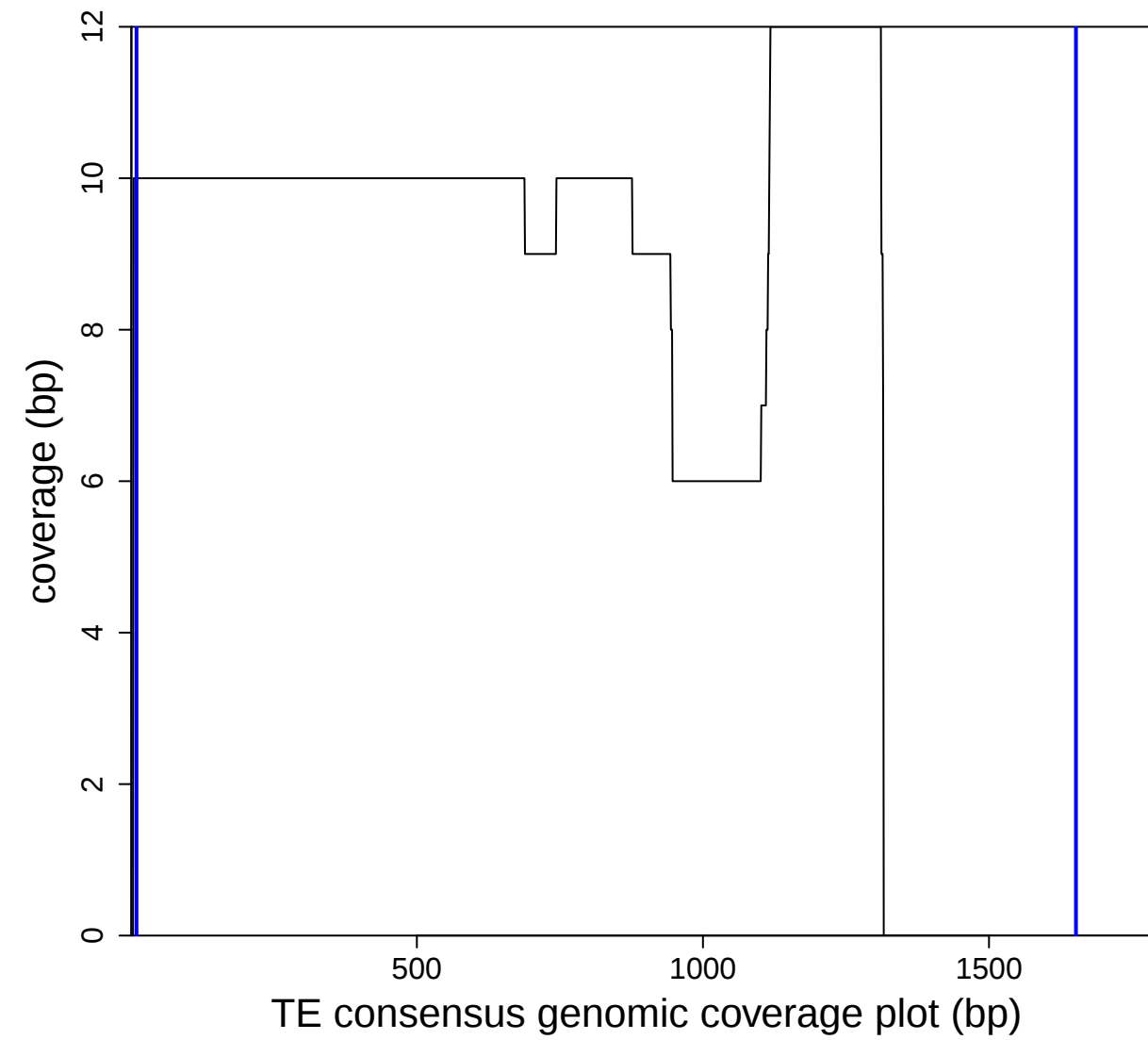
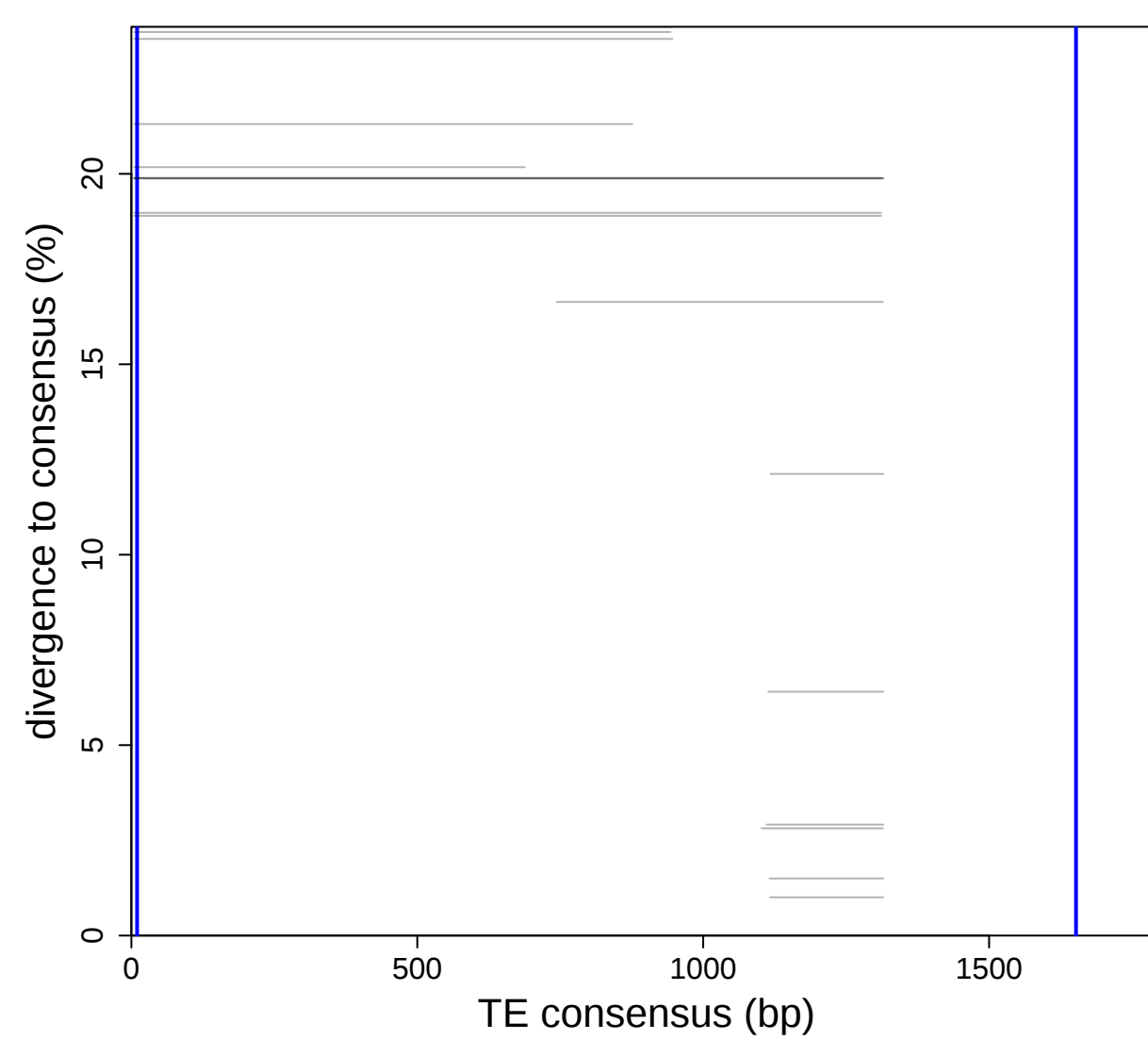
b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 1541bp; fragments: 22; full length: 0 (>=1386.9bp)

After TEtrimmer 1541 bp



2.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1802bp; fragments: 17; full length: 0 (>=1621.8bp)

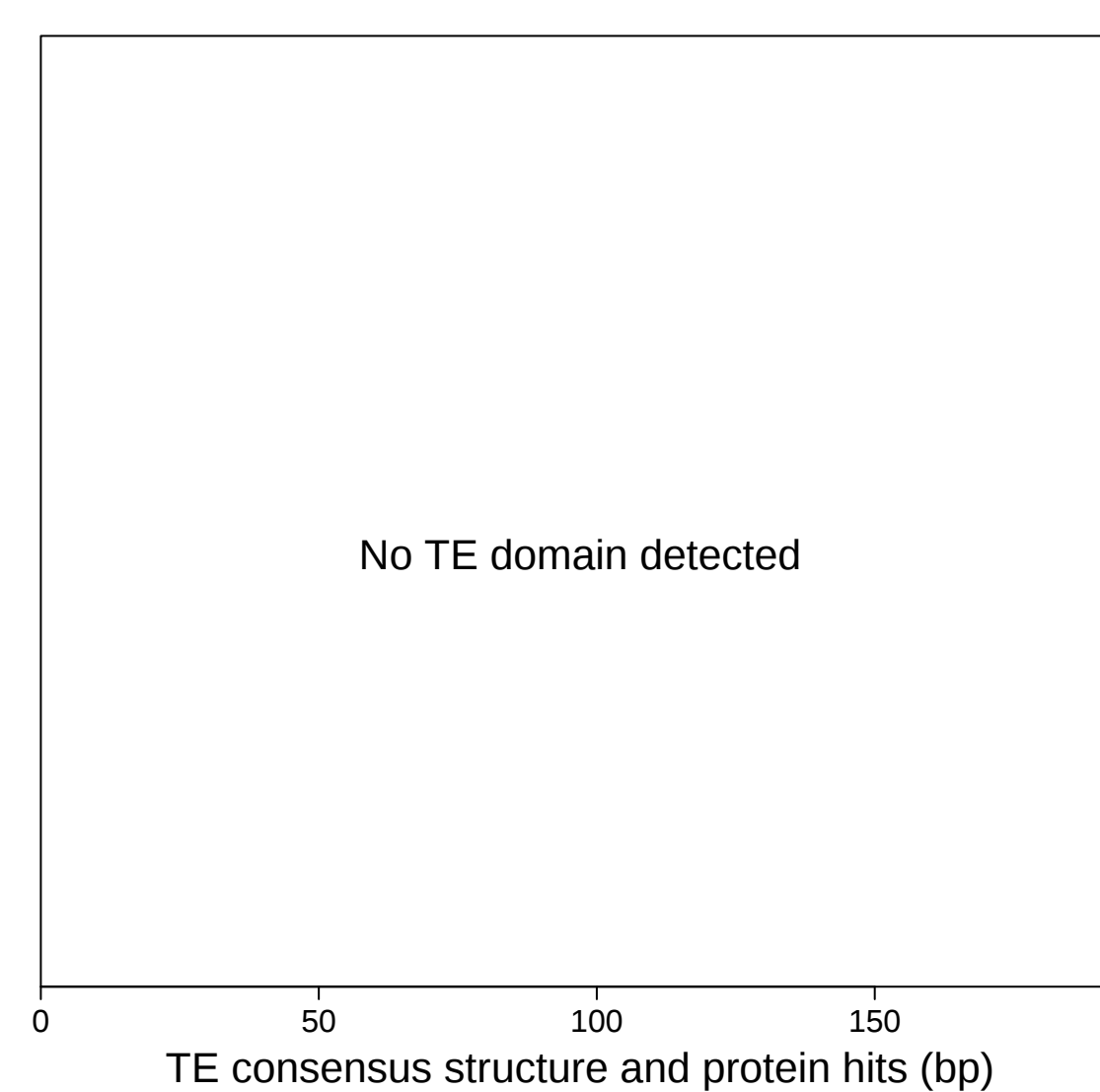
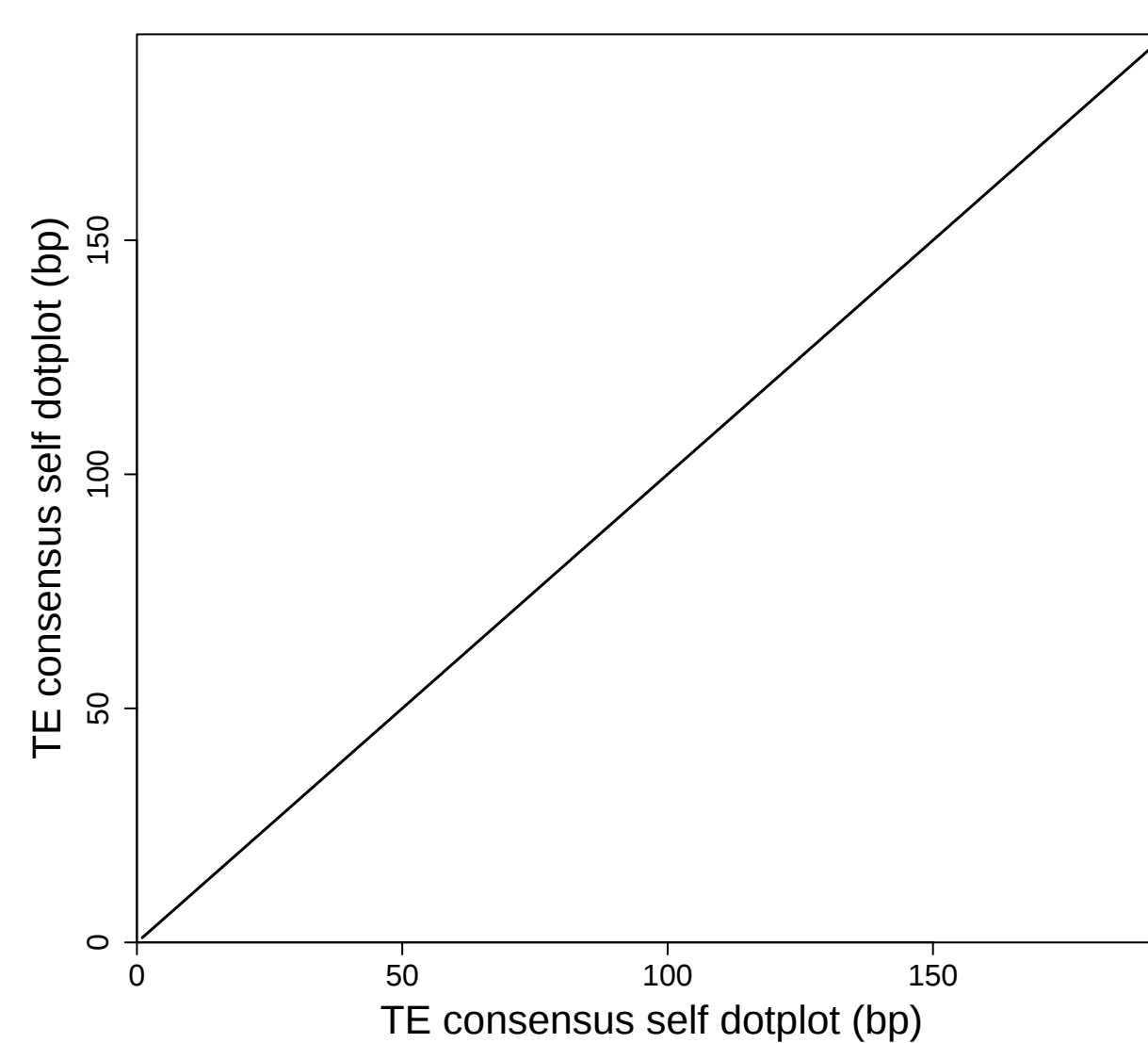
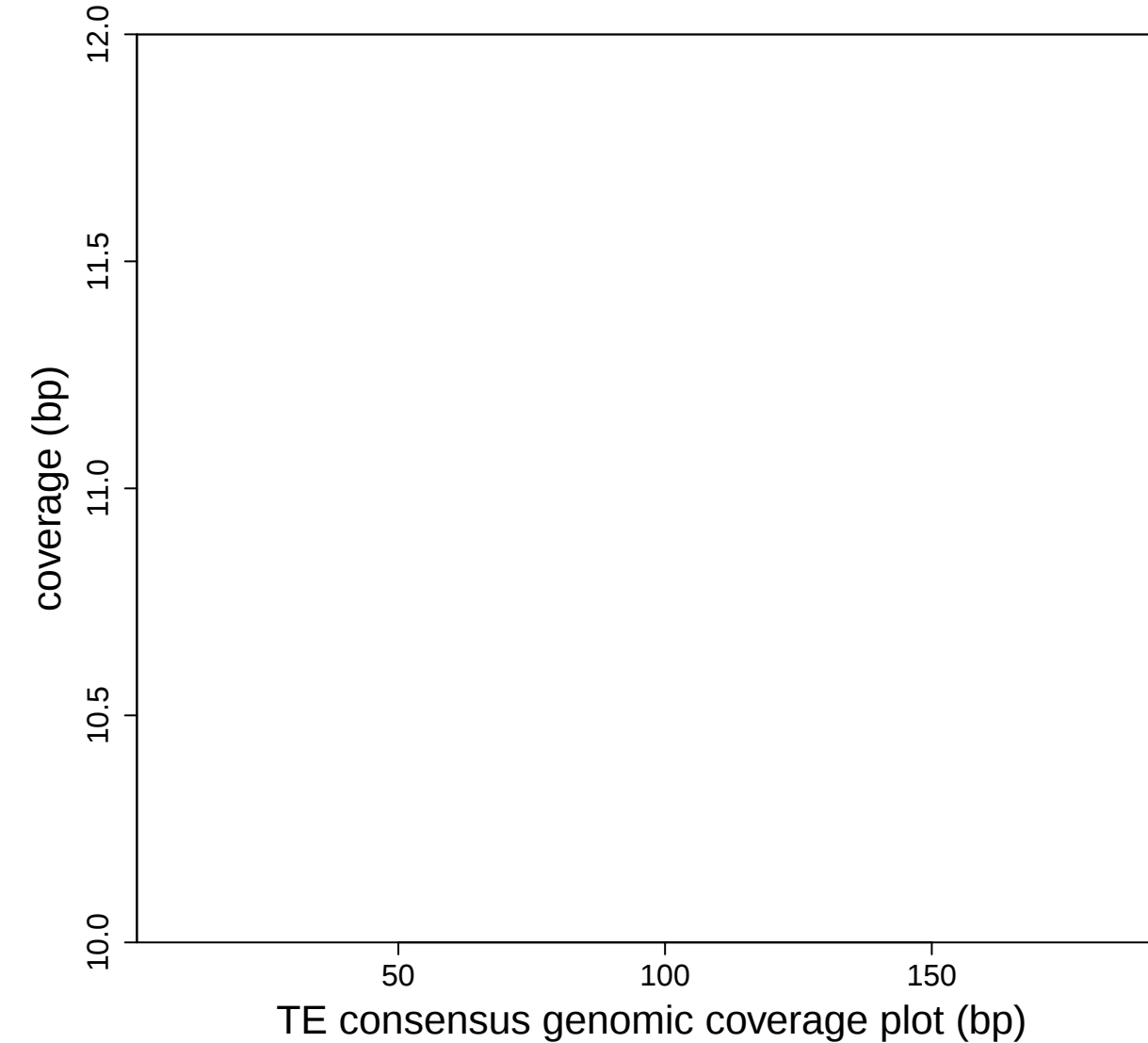
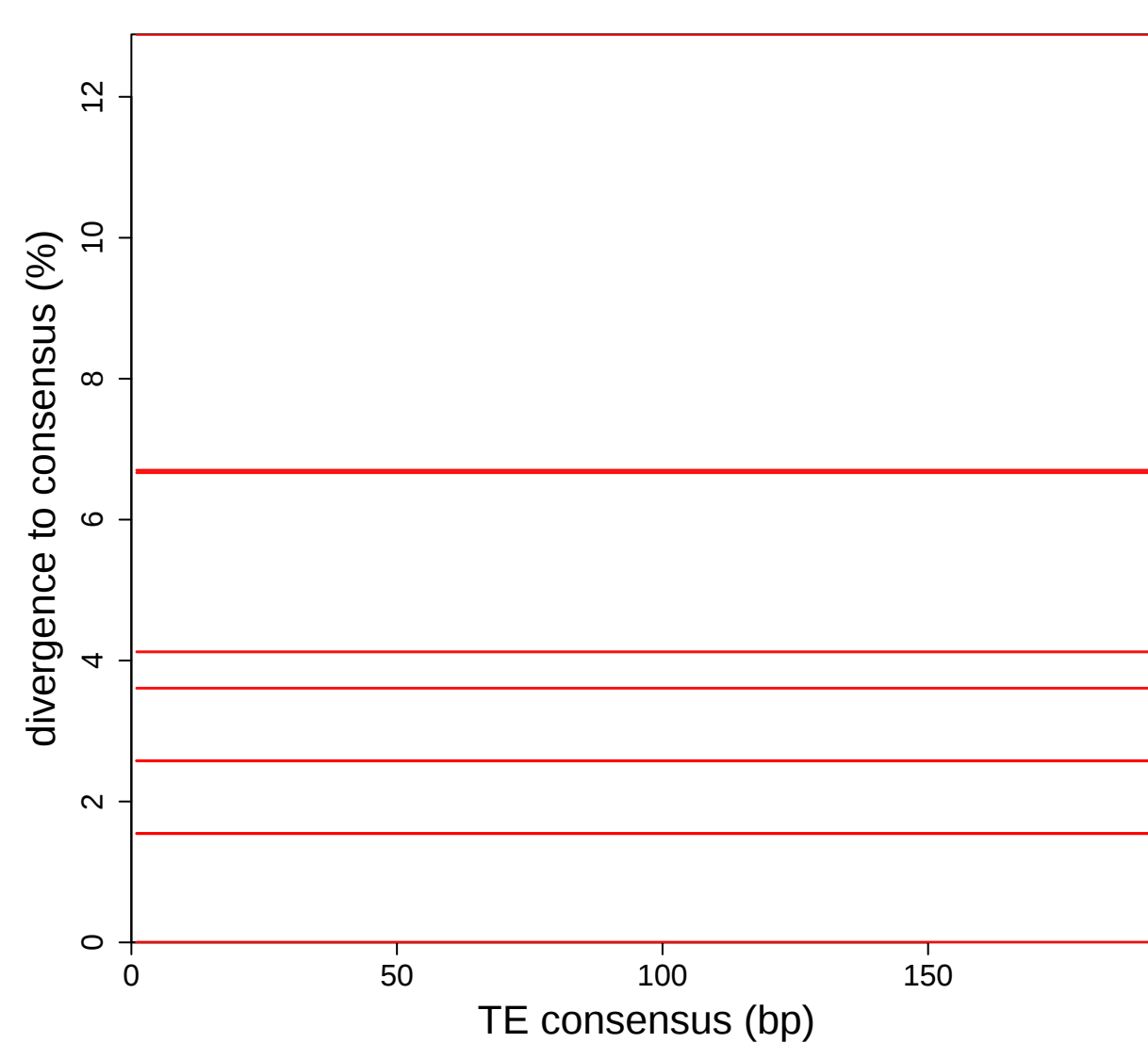
After TEtrimmer Extended plot Blue lines are boundaries



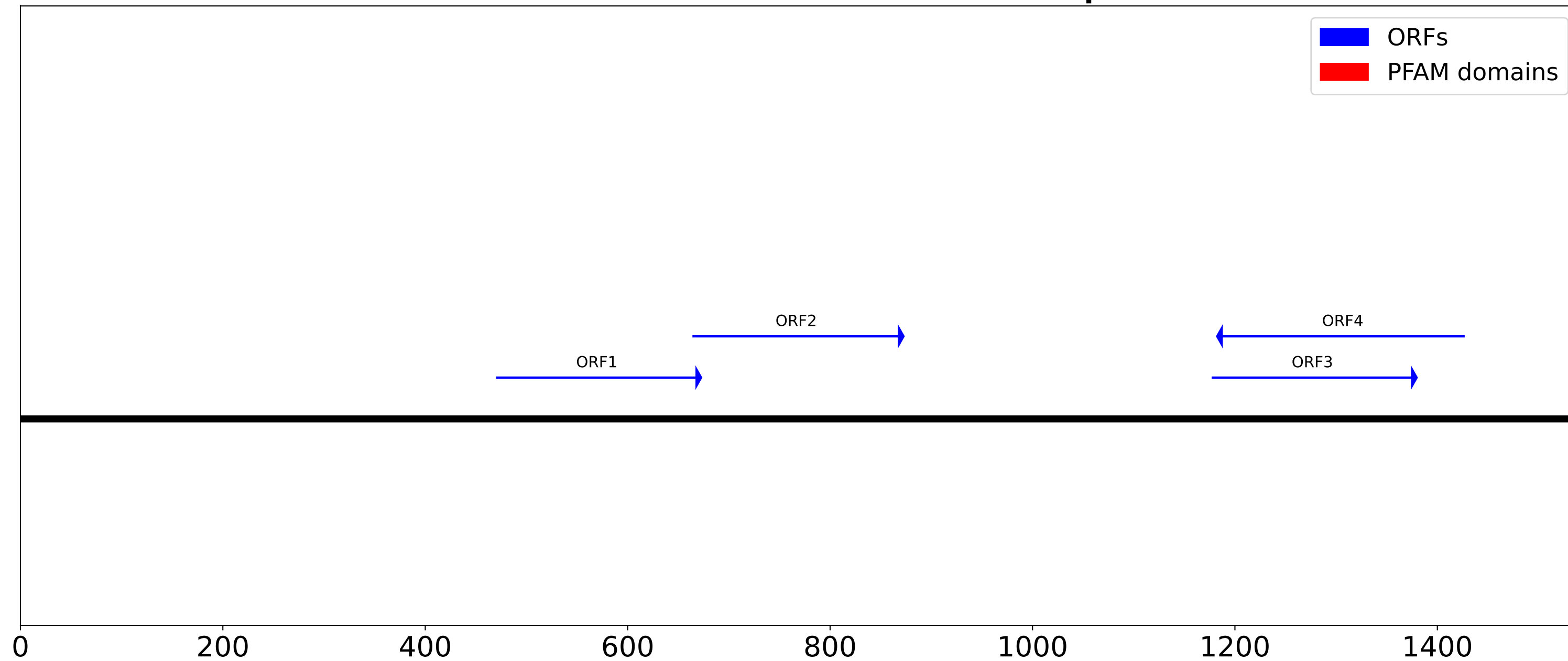
Before TEtrimmer 194 bp

TE: ltr_1_family_272

size: 194bp; fragments: 12; full length: 12 (≥ 174.6 bp)



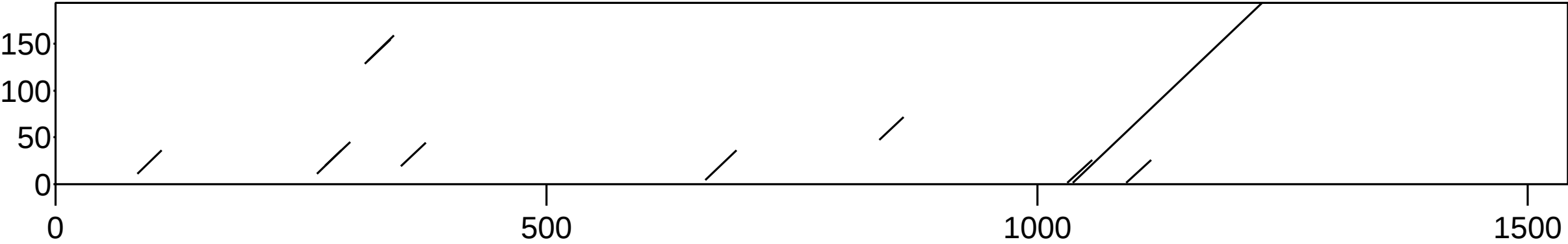
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

sensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)