

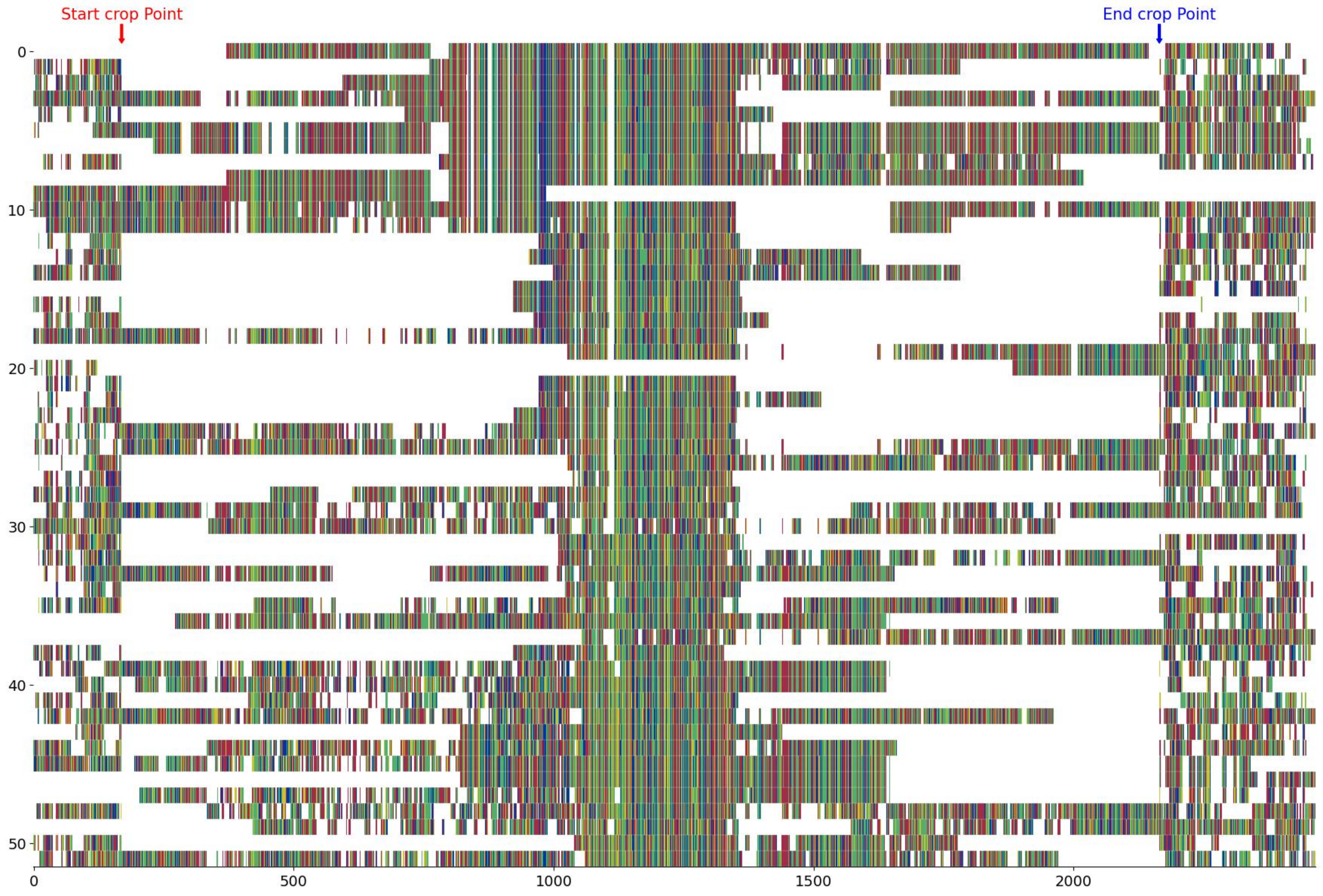
MSA length = 1997

Start crop Point

End crop Point

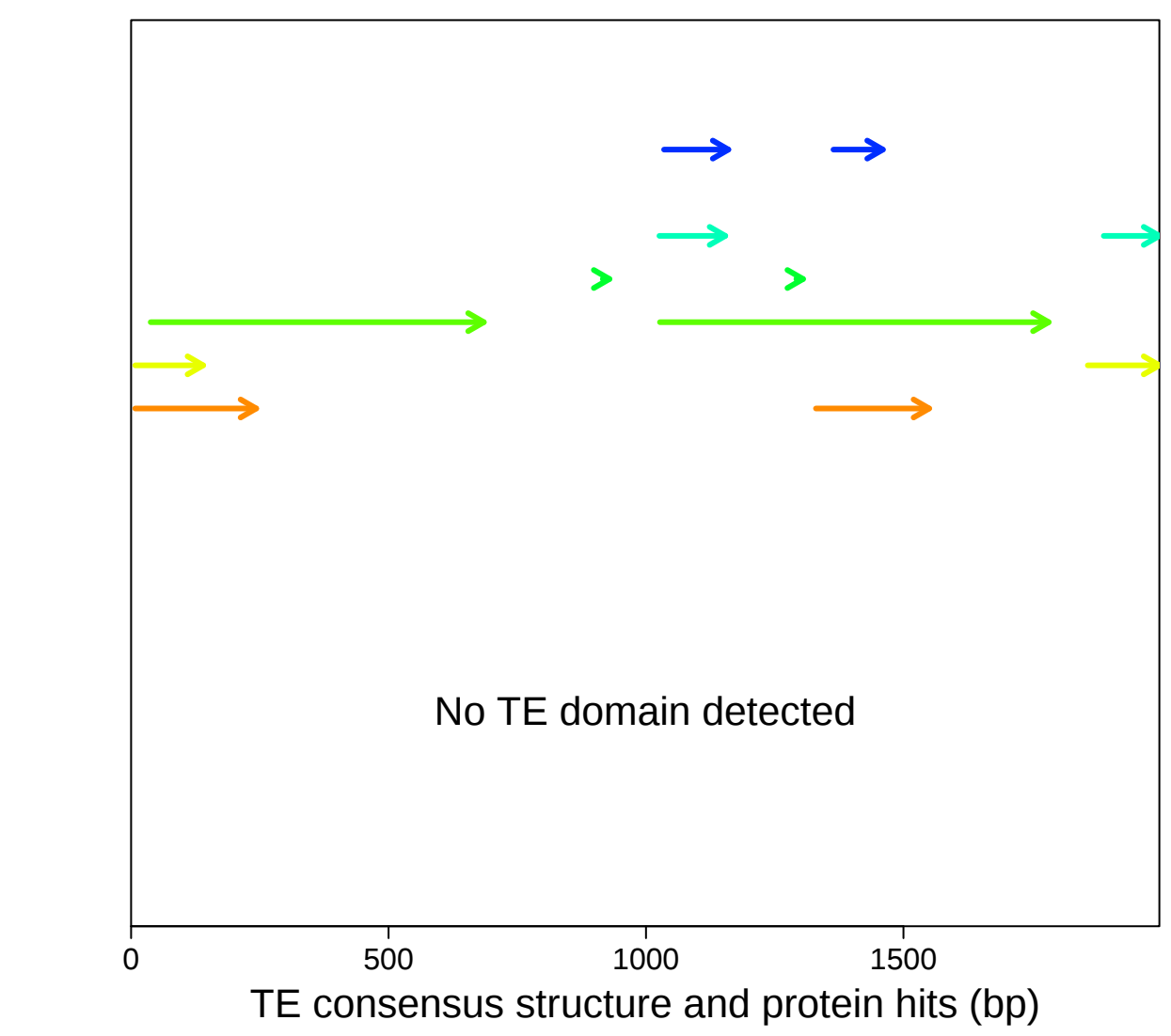
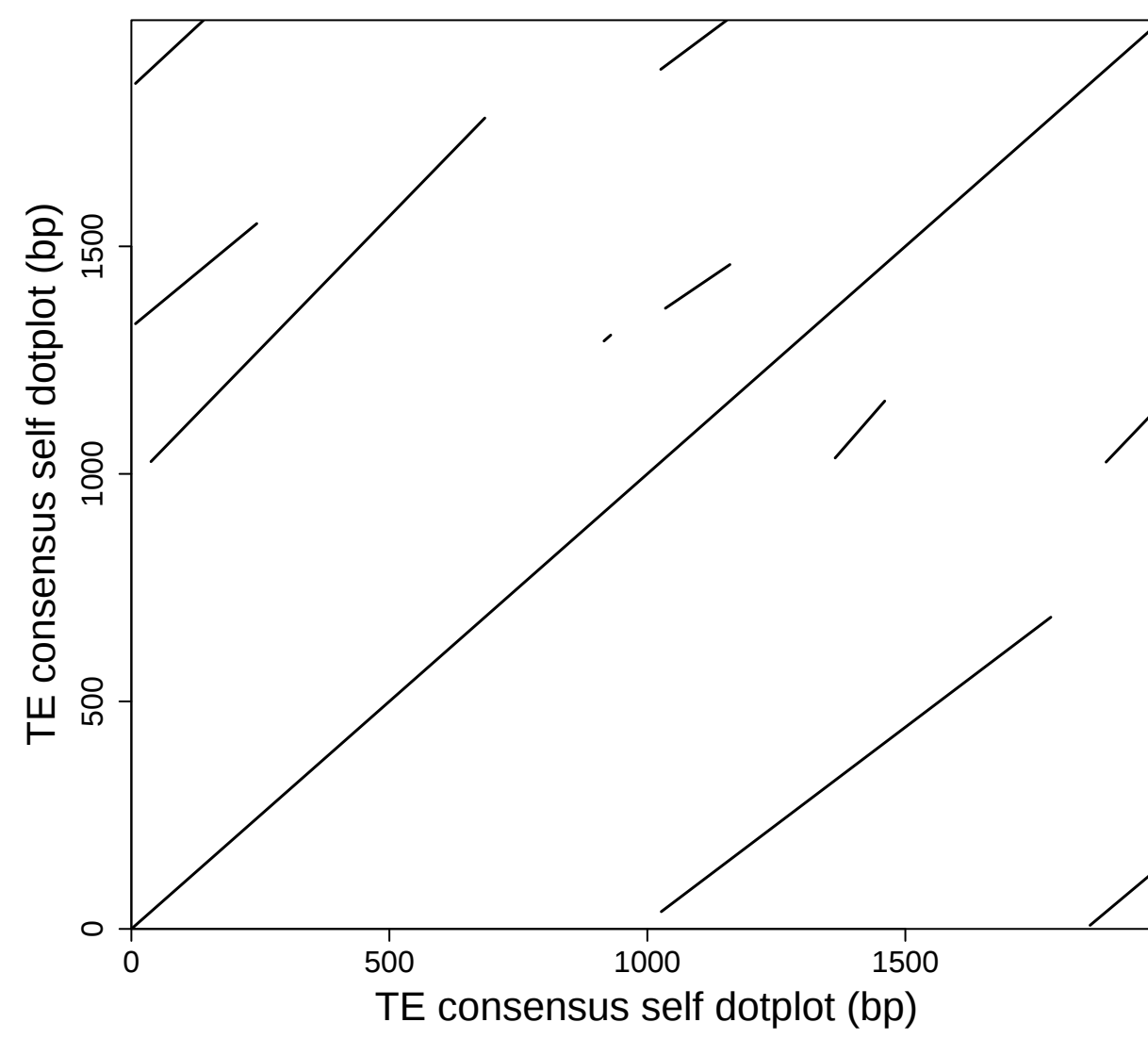
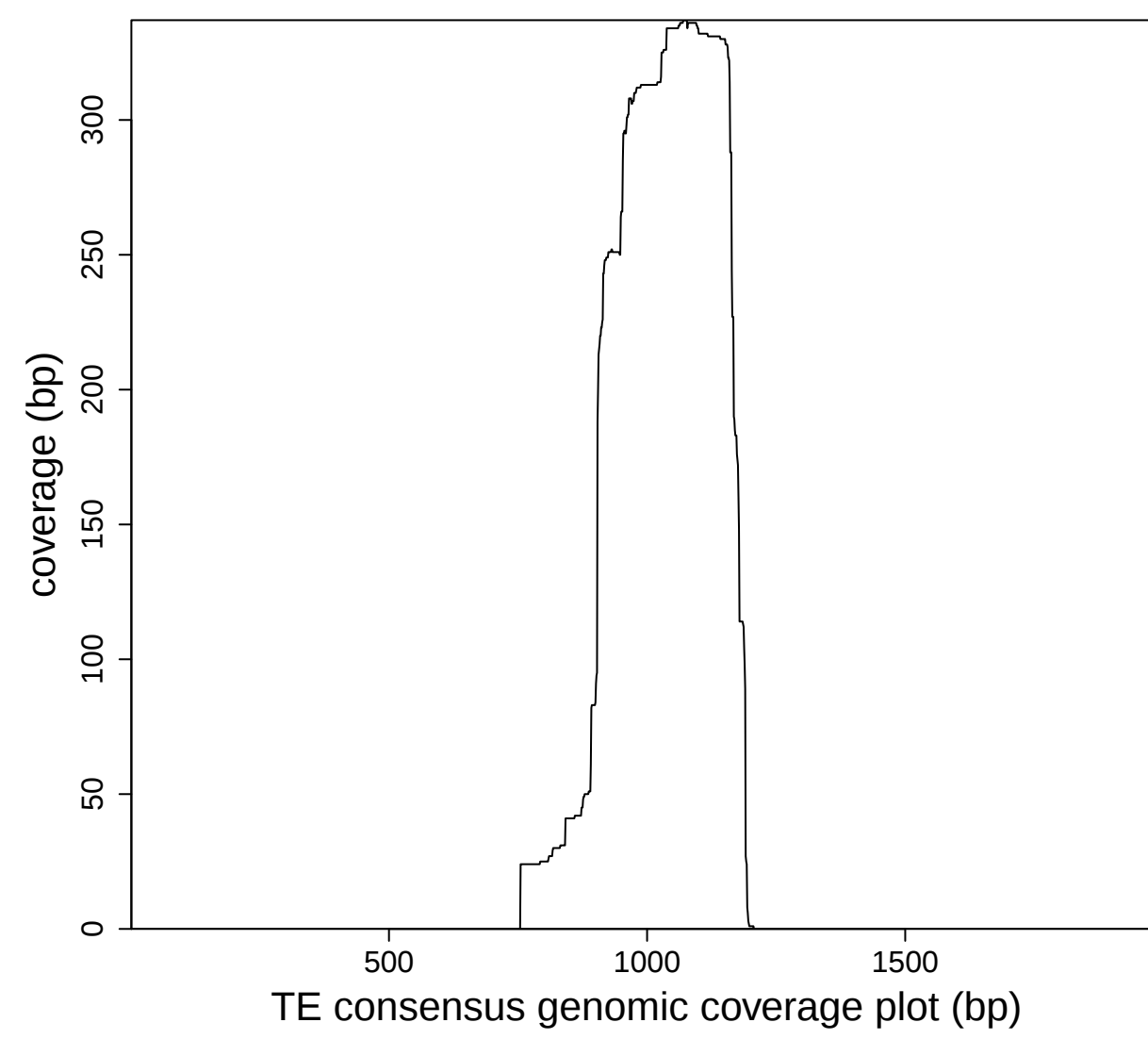
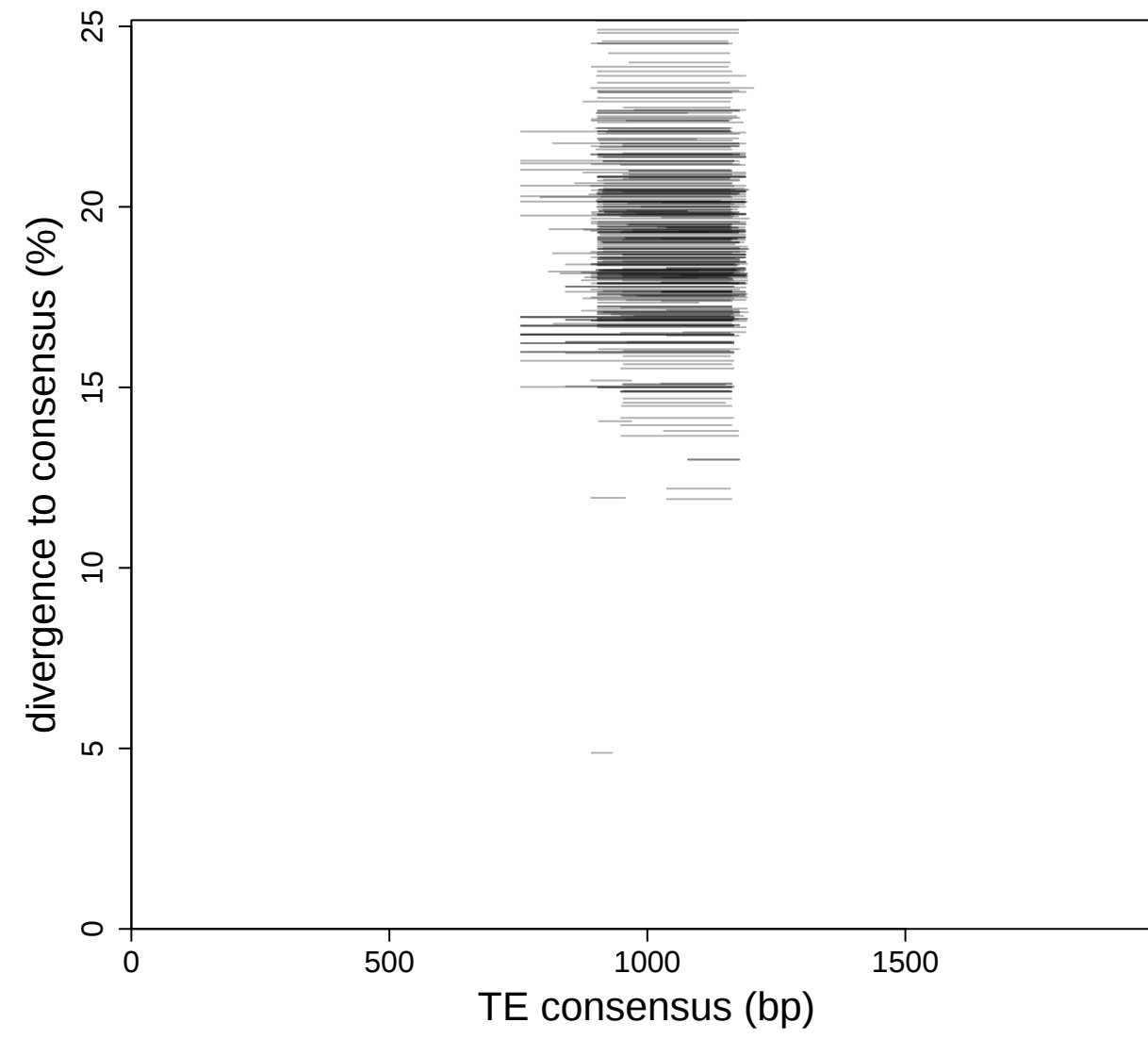
Start crop Point

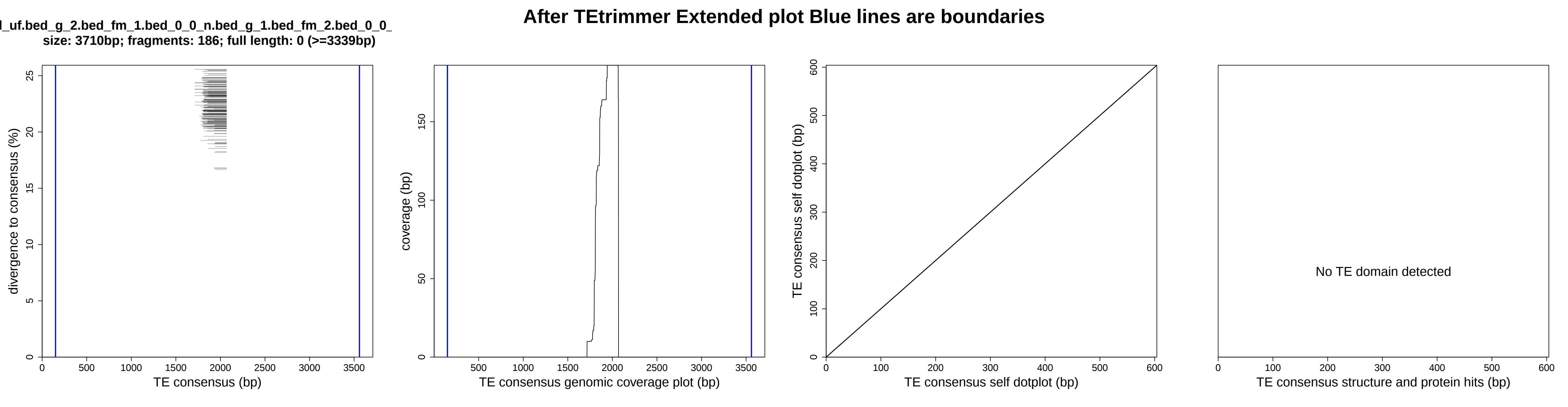
End crop Point



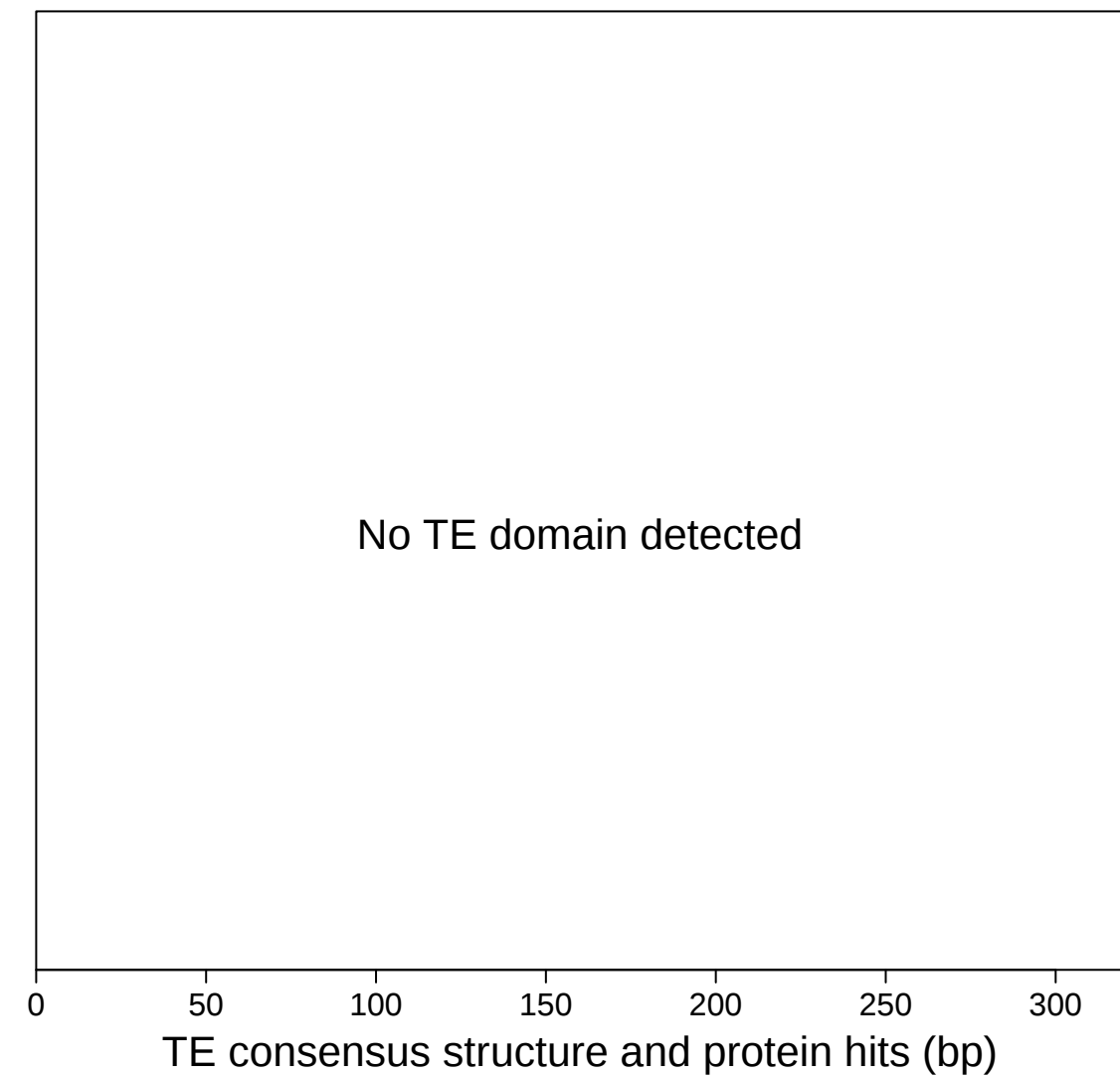
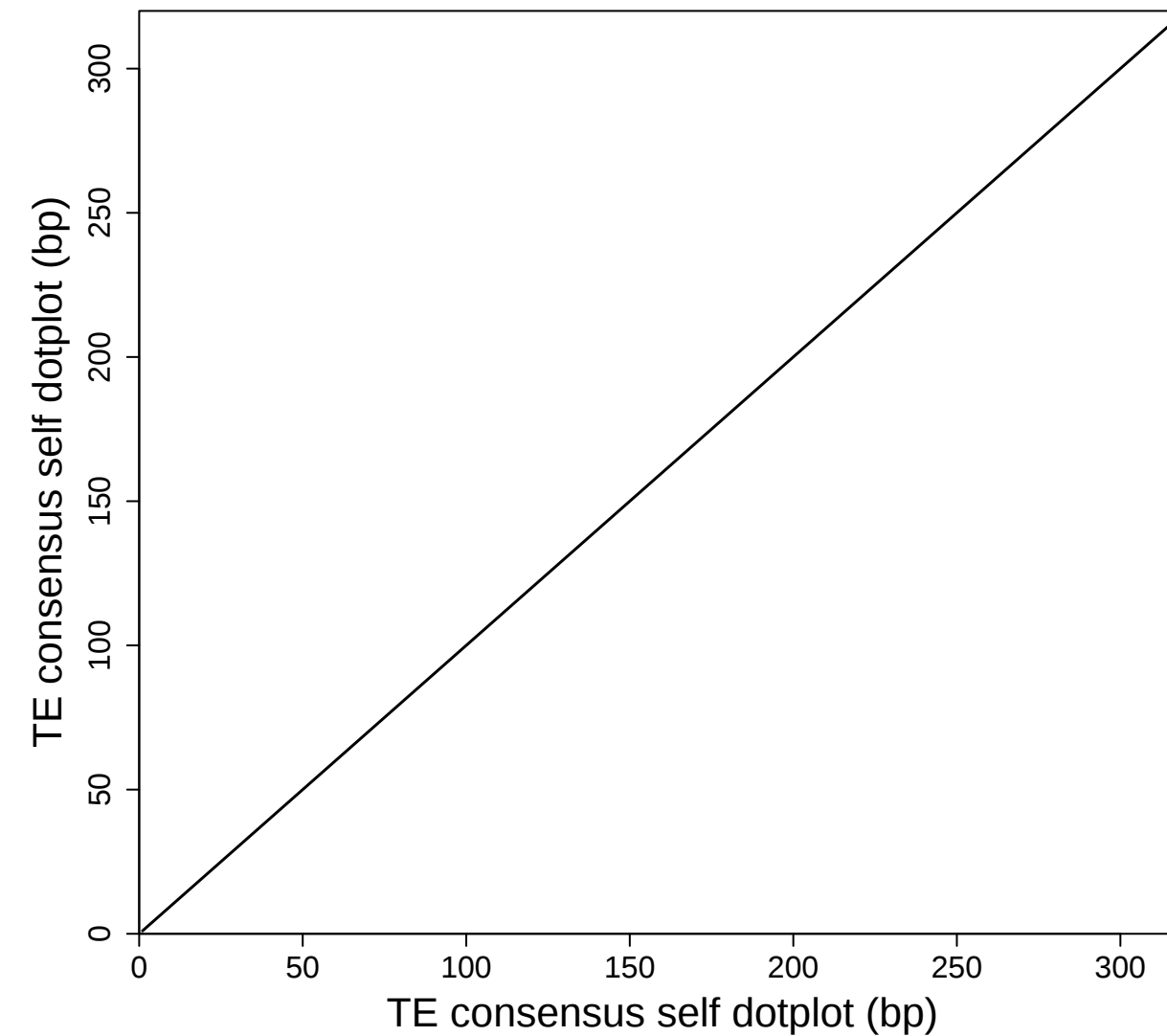
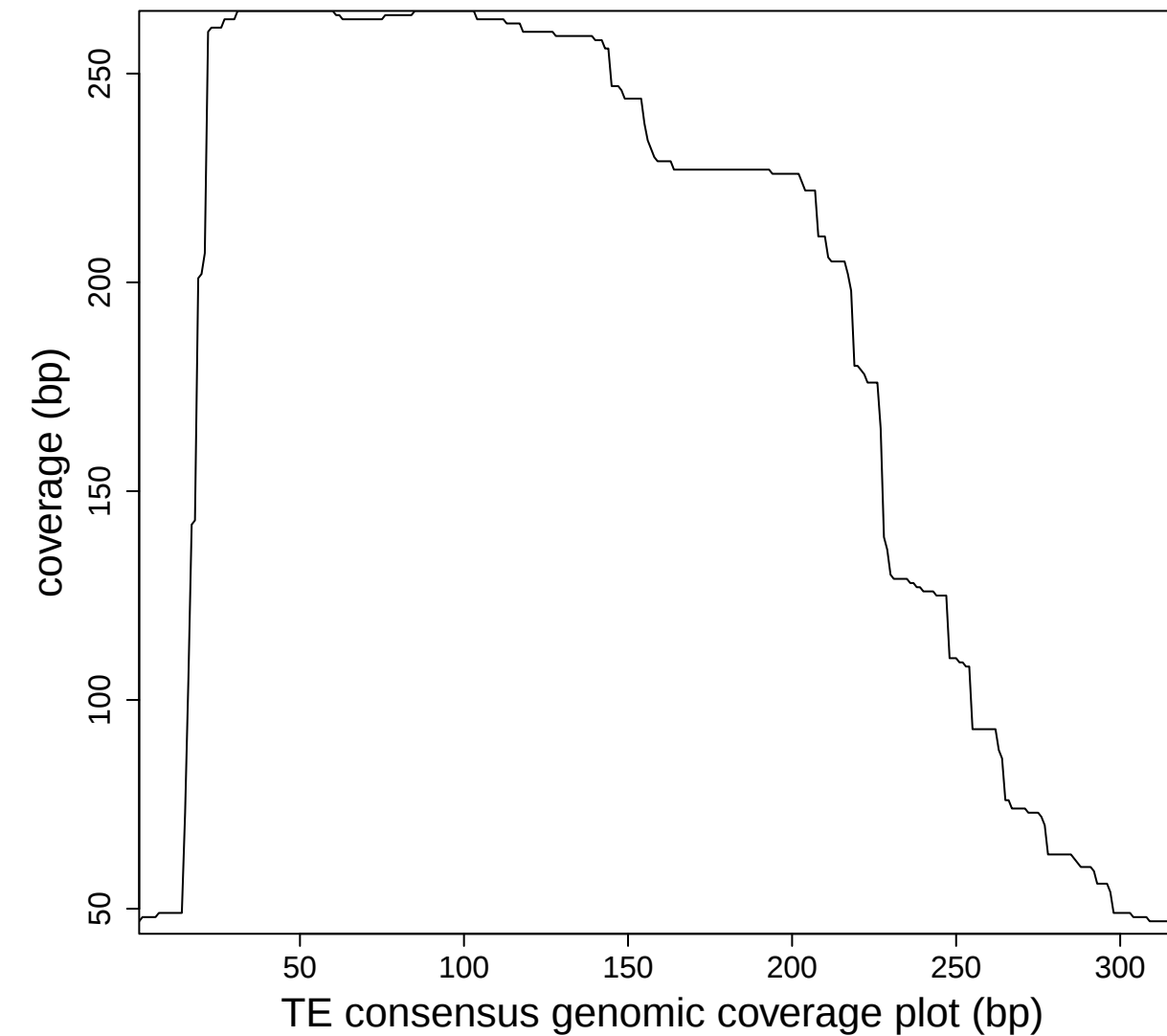
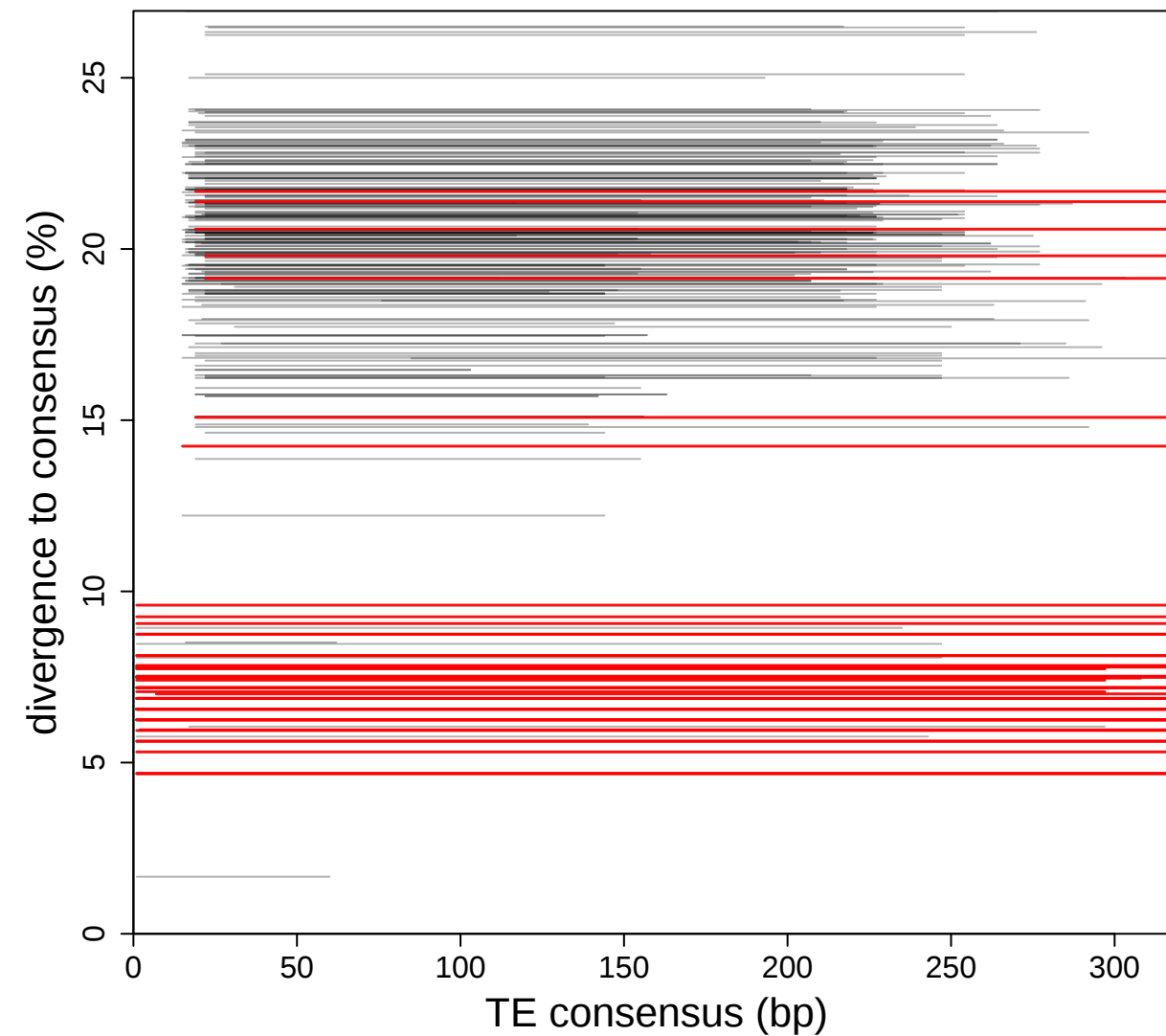
bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.fa
size: 1997bp; fragments: 344; full length: 0 (>=1797.3bp)

After TEtrimmer 1997 bp



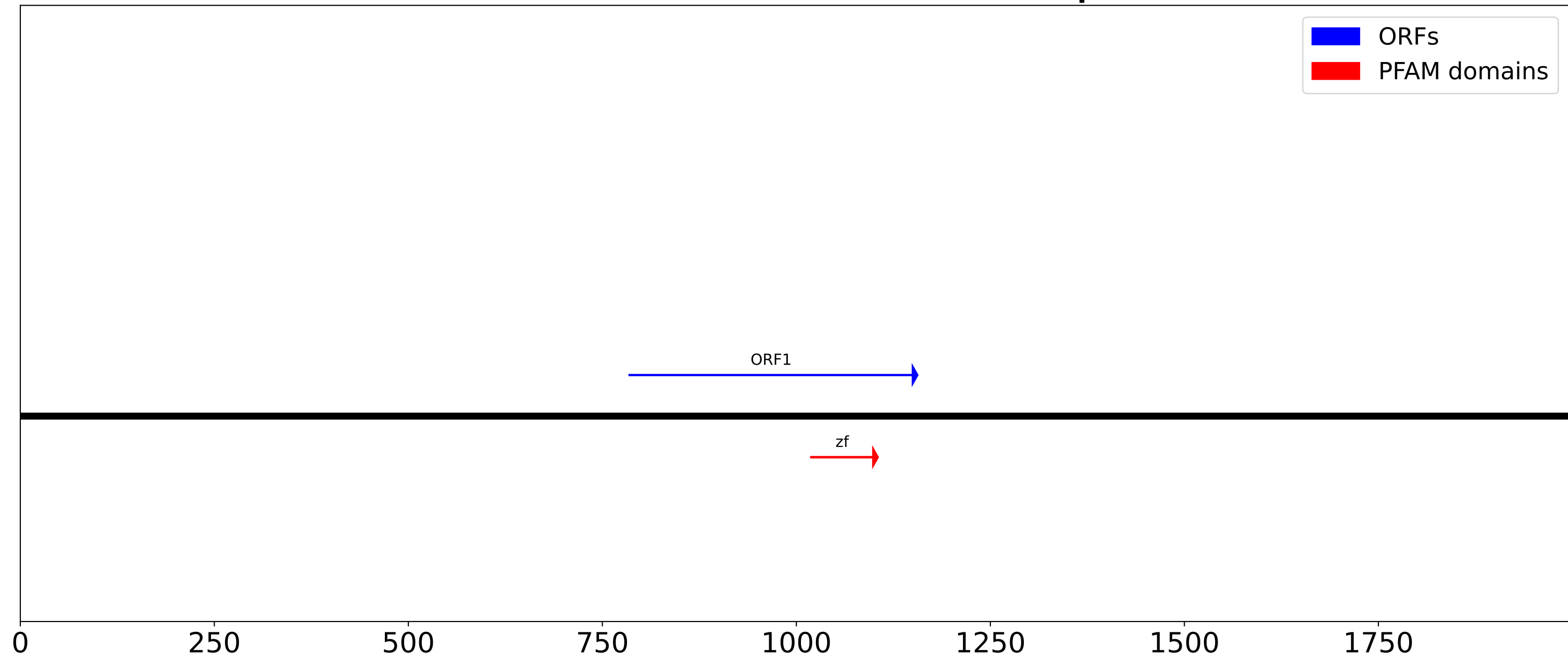


TE: rnd_1_family_947
size: 320bp; fragments: 267; full length: 51 (≥ 288 bp)

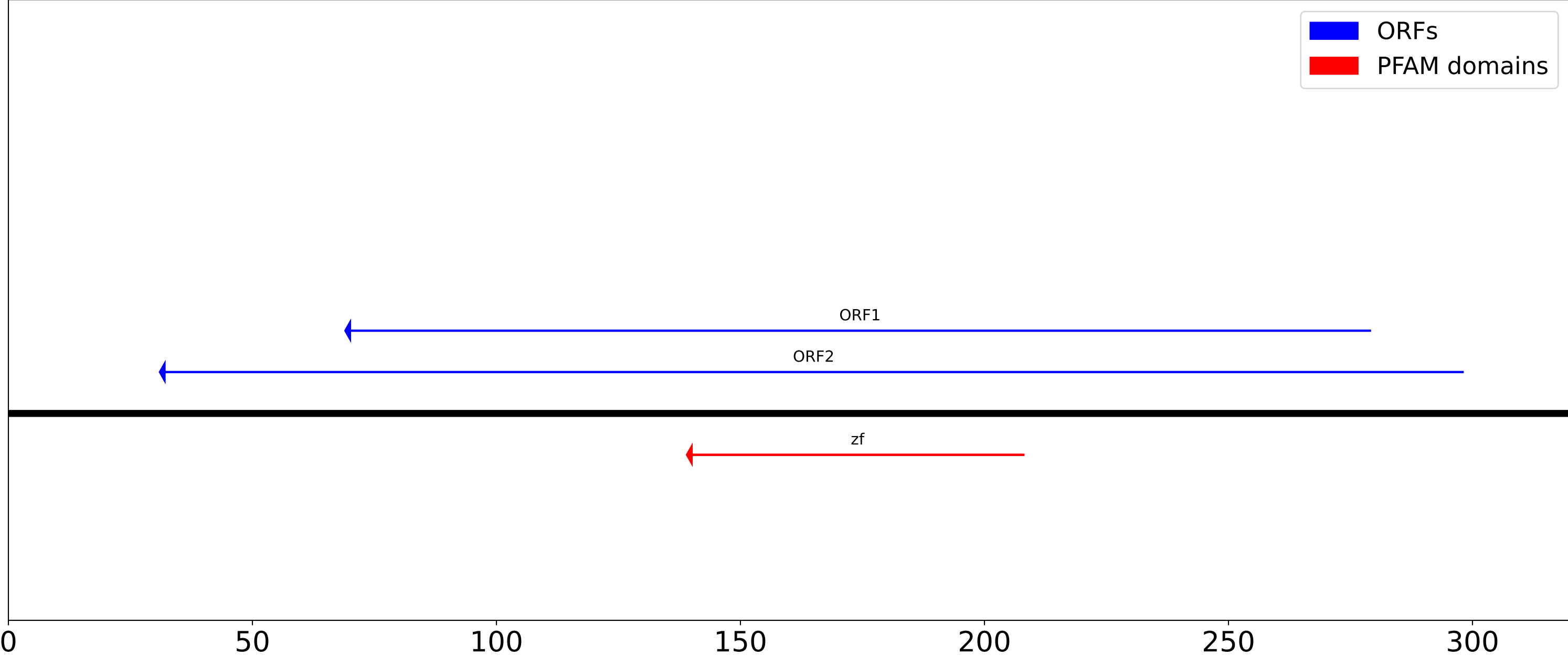


Before TEtrimmer 320 bp

After TEtrimmer ORF and PFAM domain plot



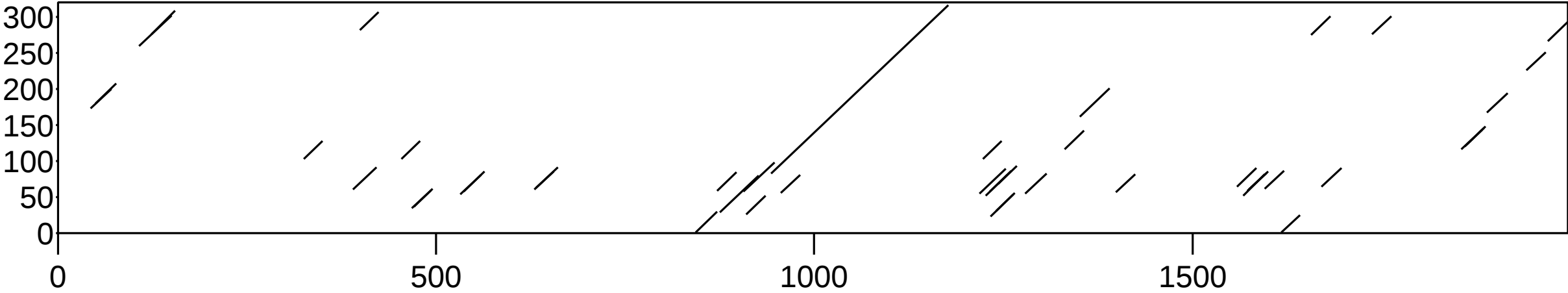
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)