

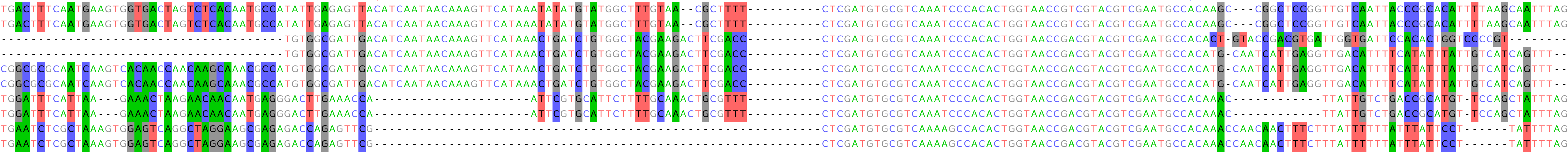
Start Crop Point



End Crop Point

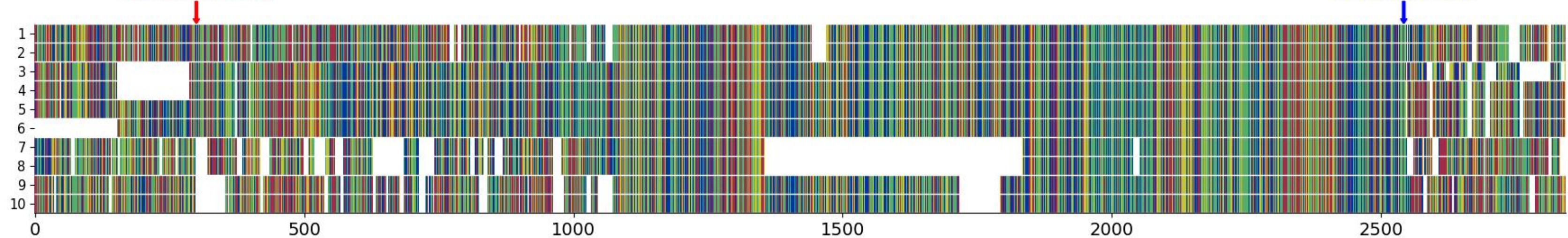


MSA length = 2243



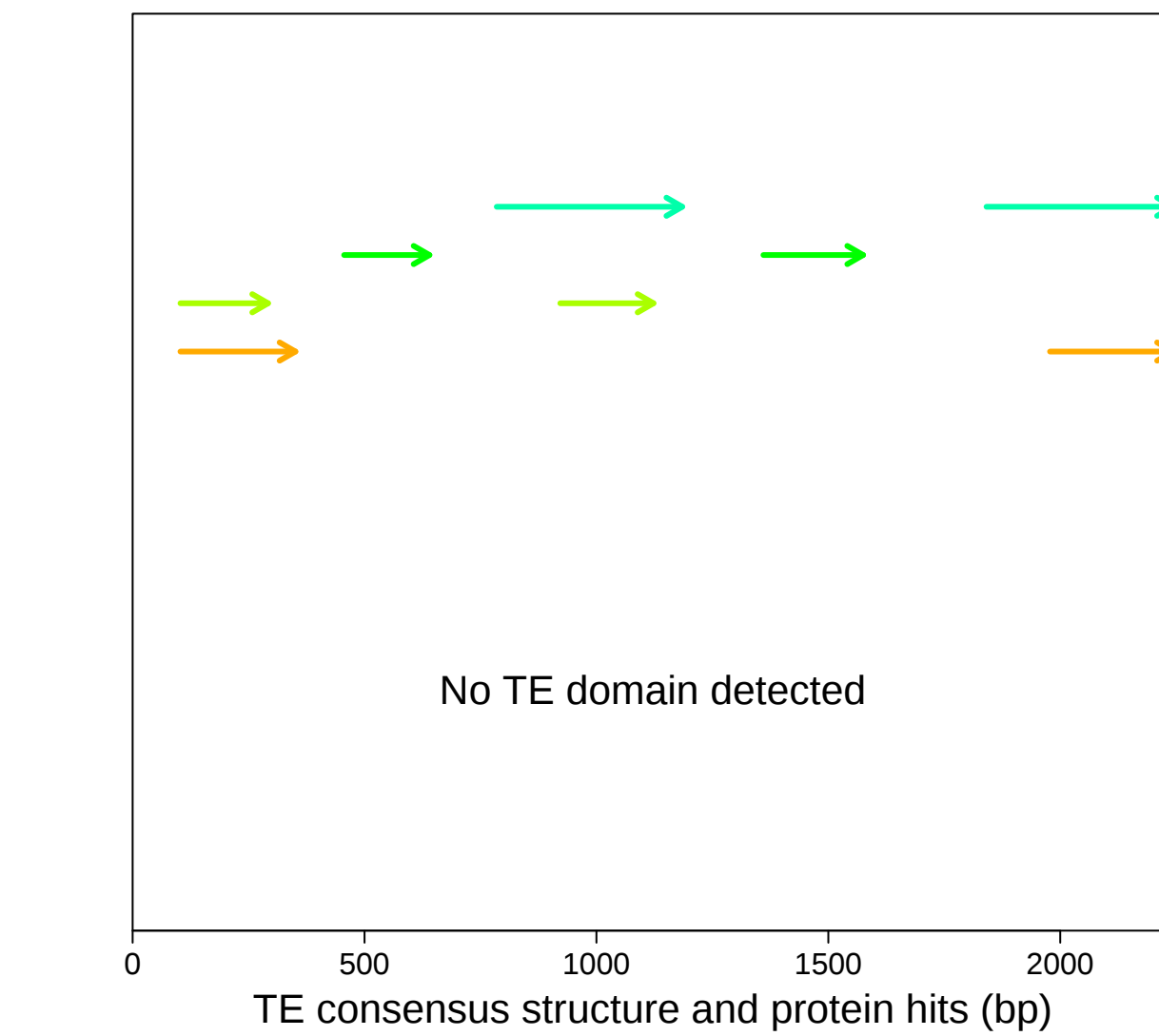
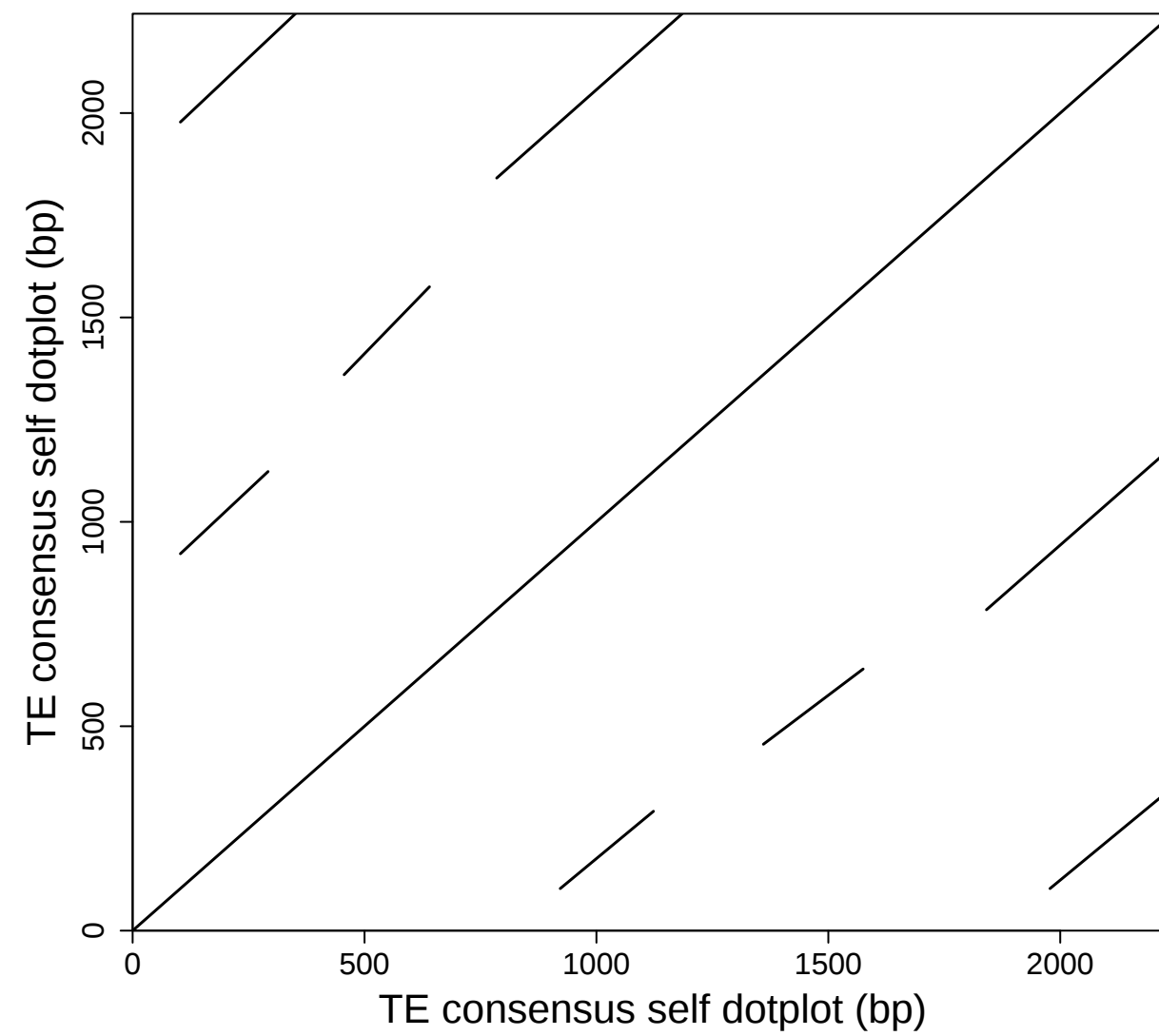
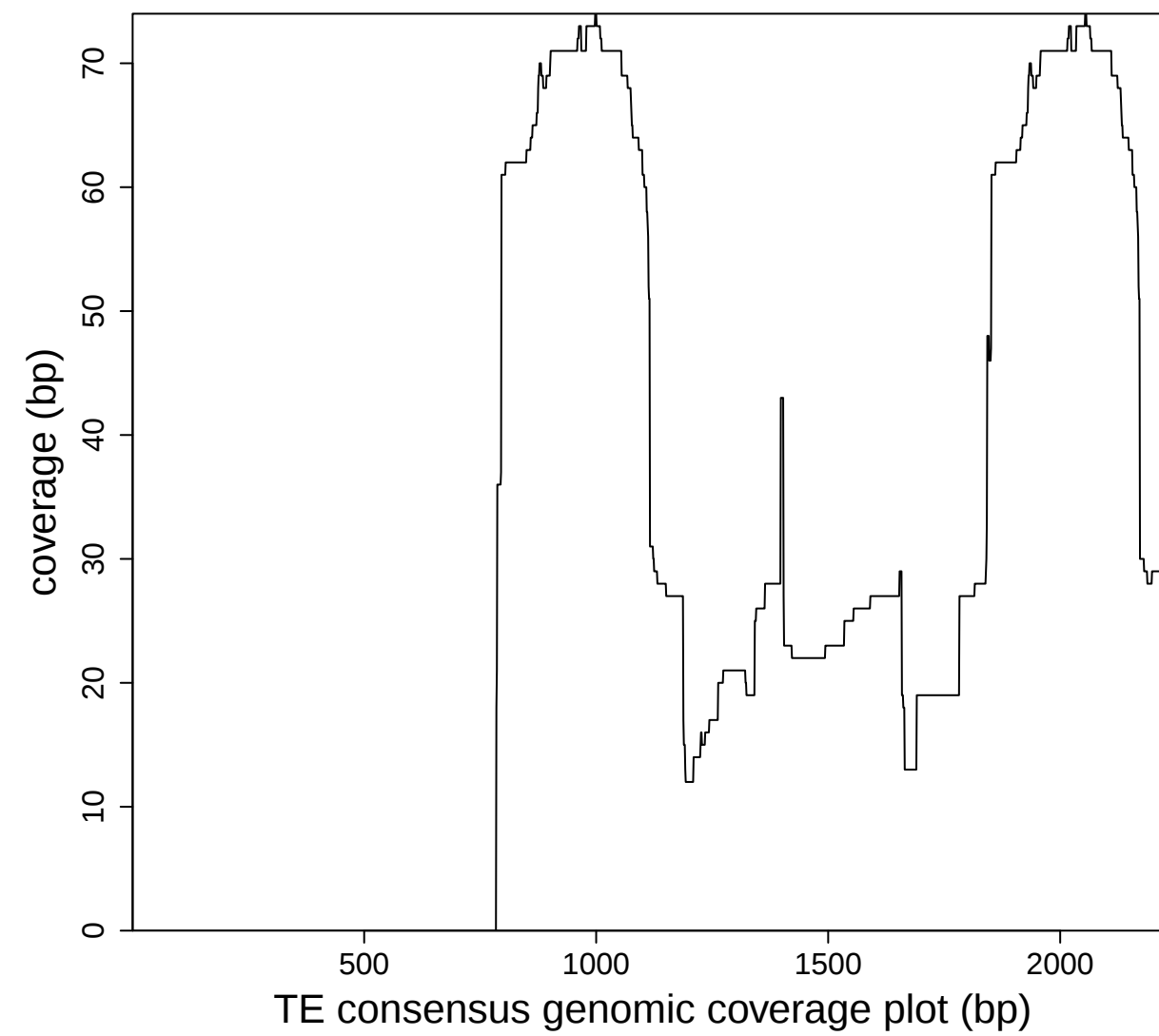
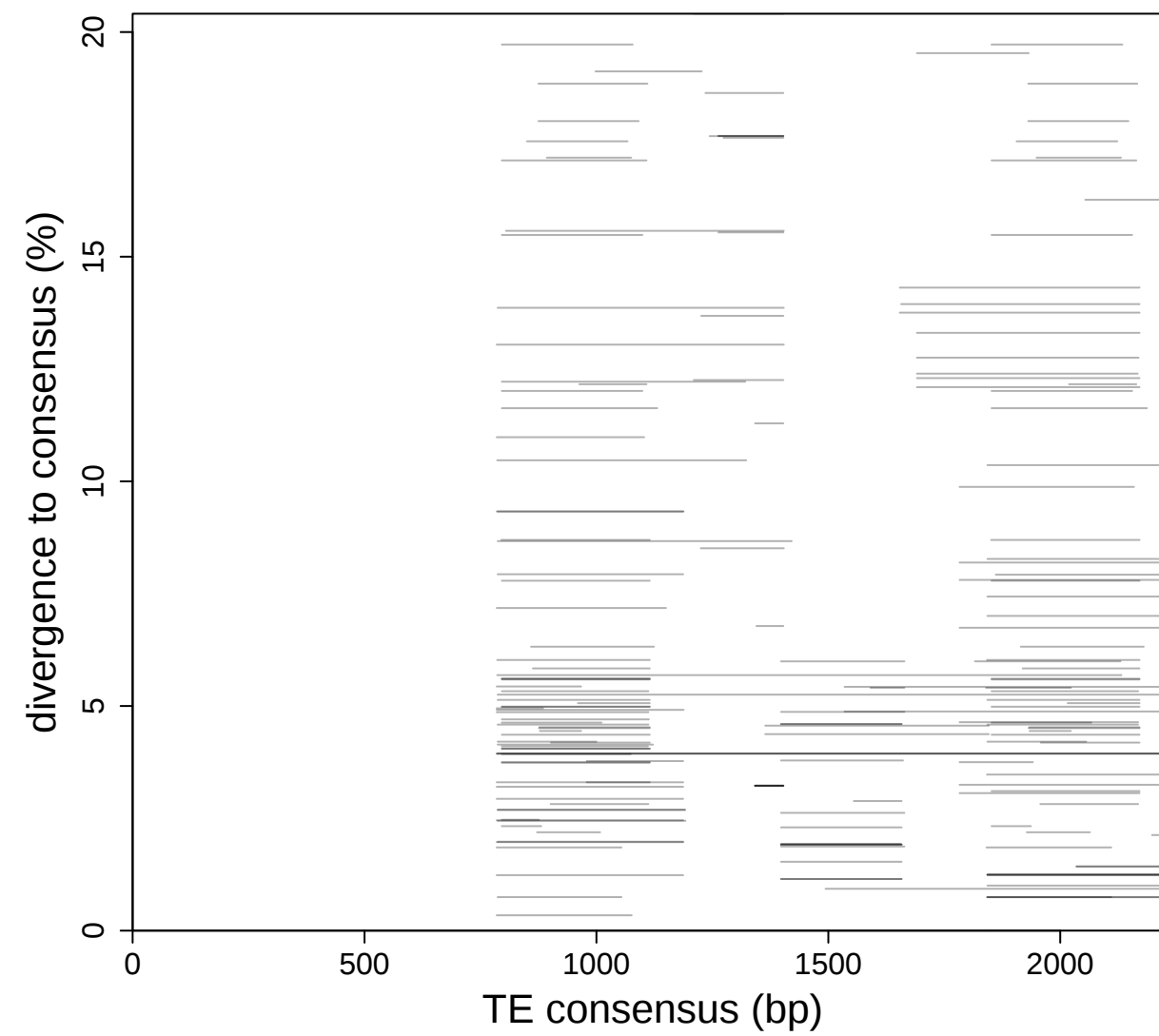
Start crop Point

End crop Point



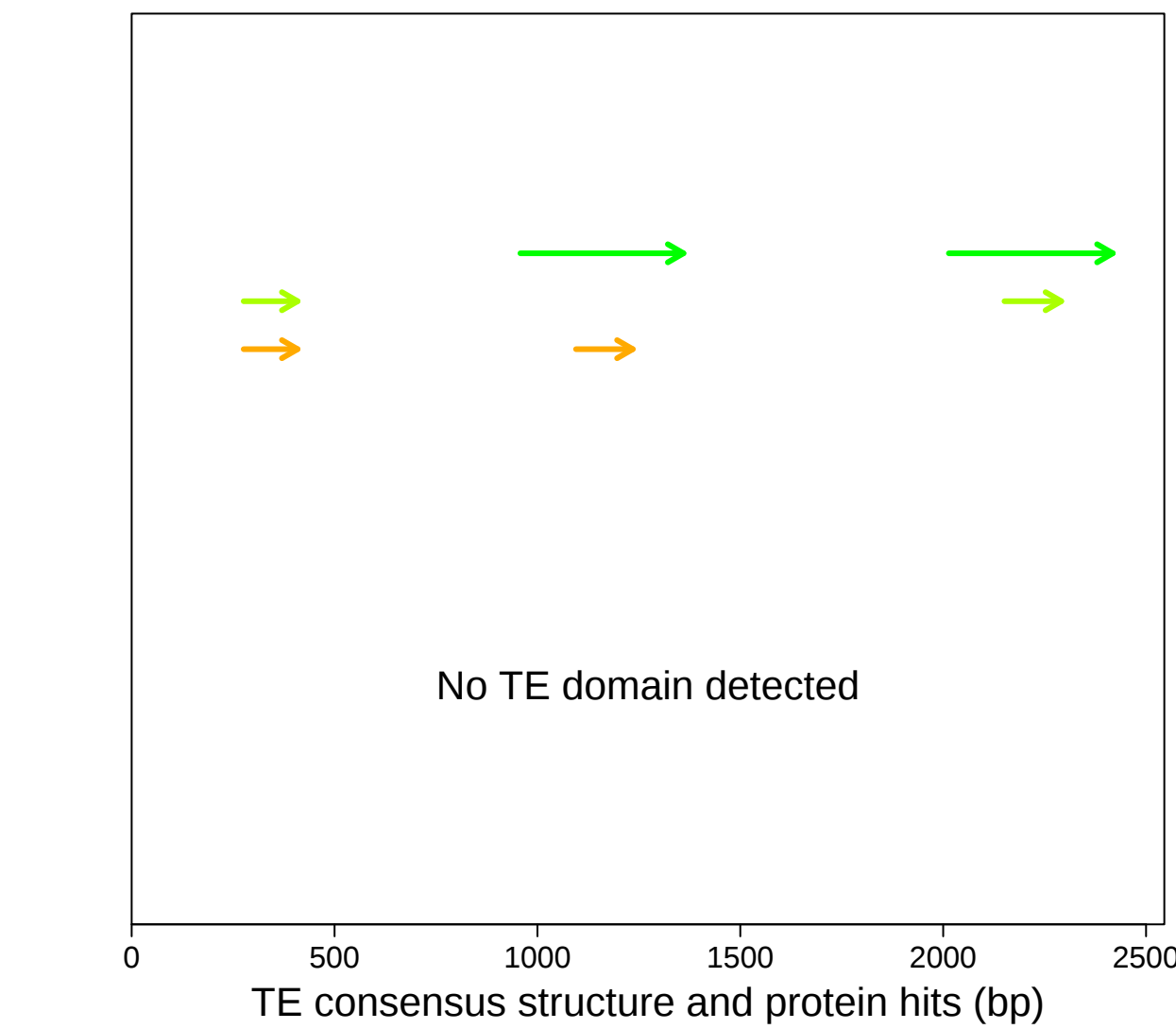
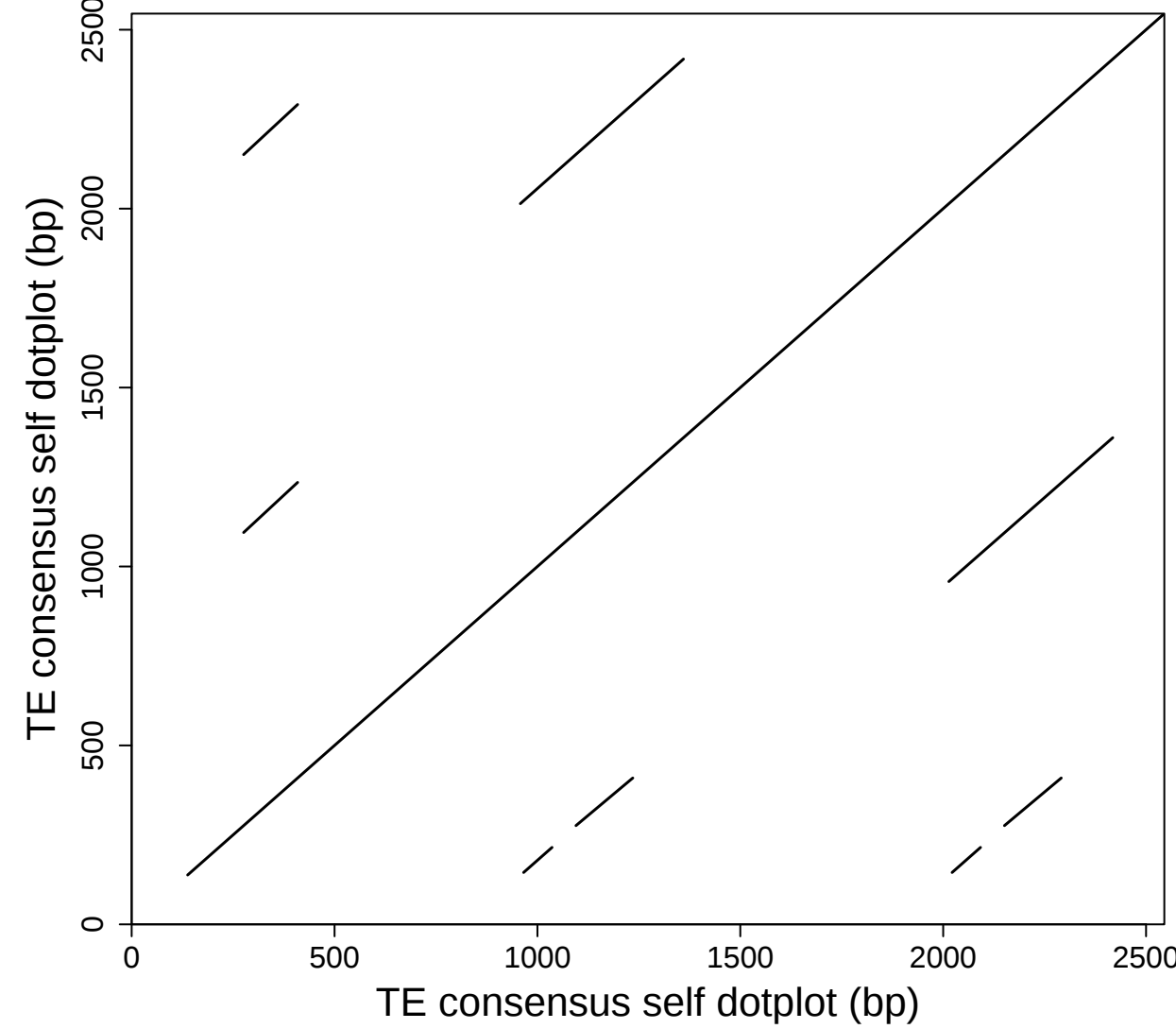
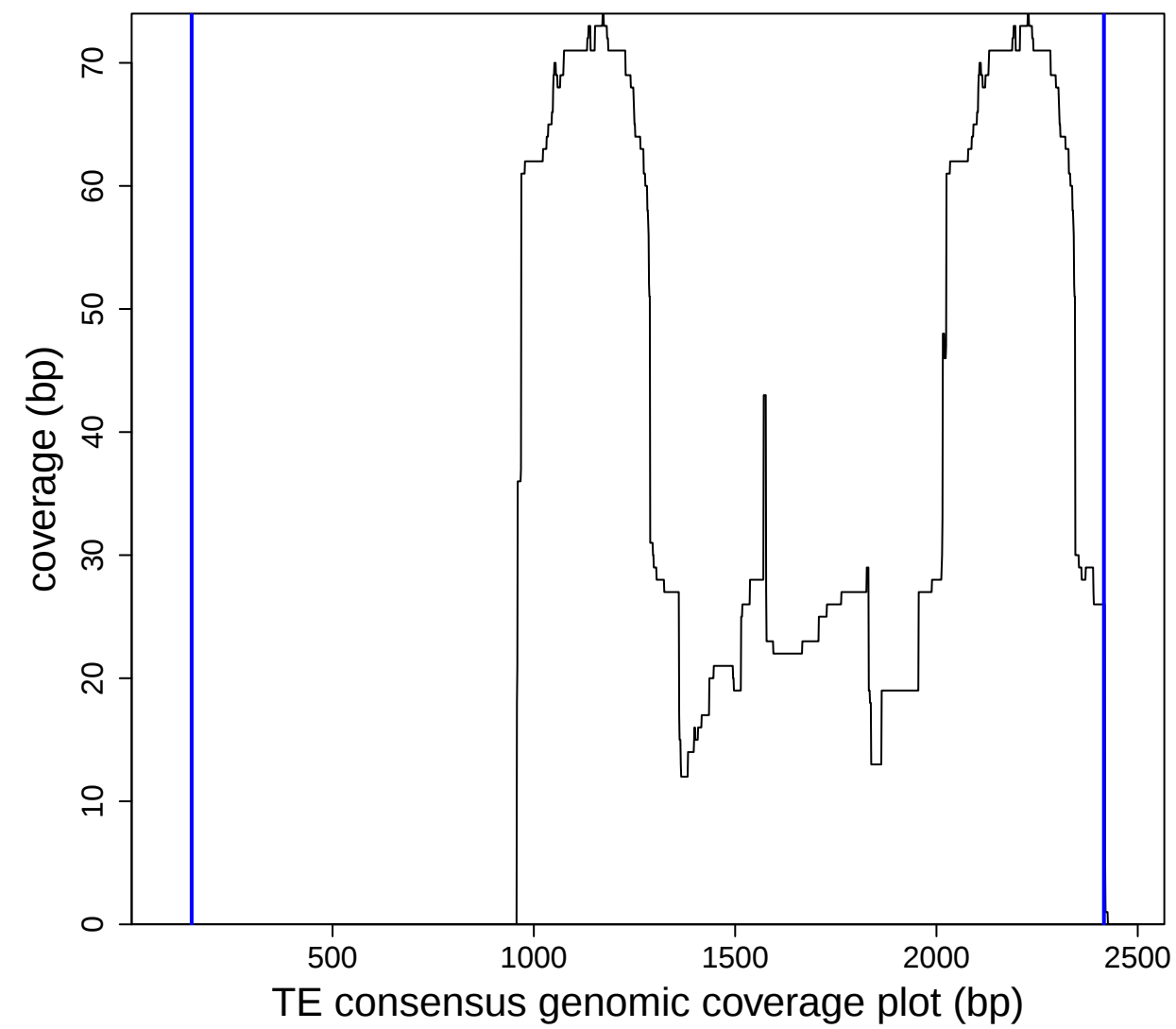
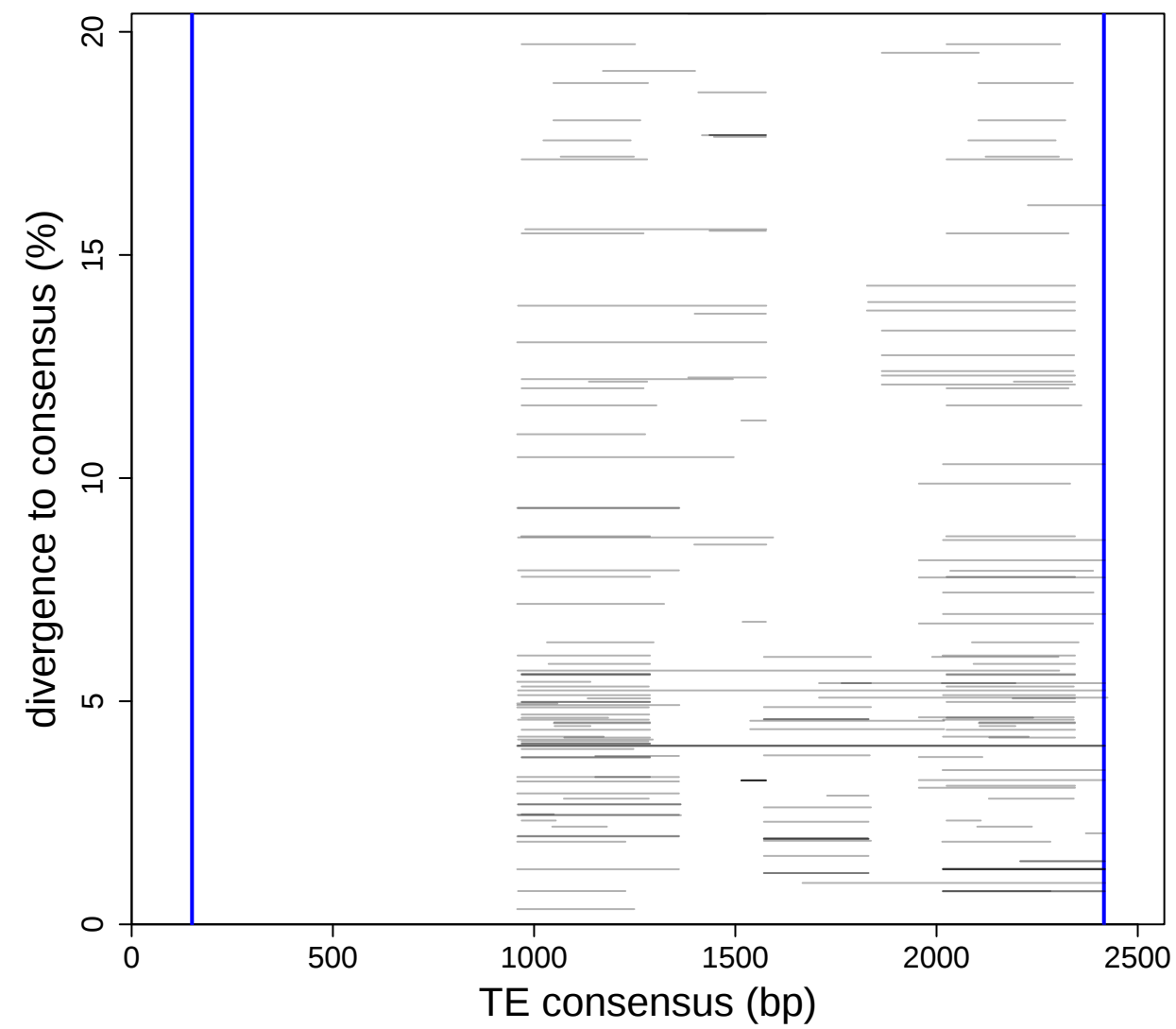
bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.fa
size: 2243bp; fragments: 190; full length: 0 (>=2018.7bp)

After TETrimmer 2243 bp

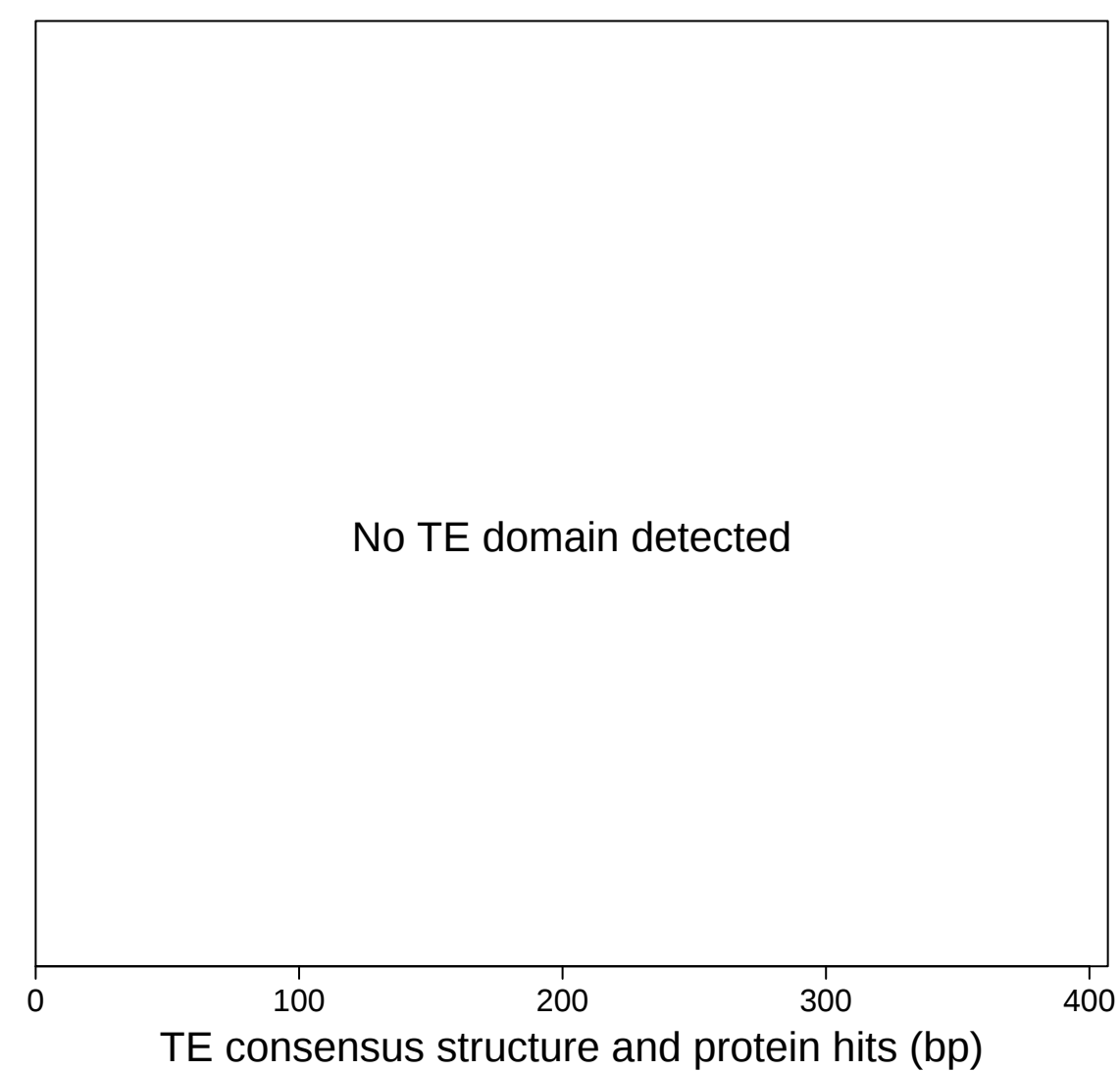
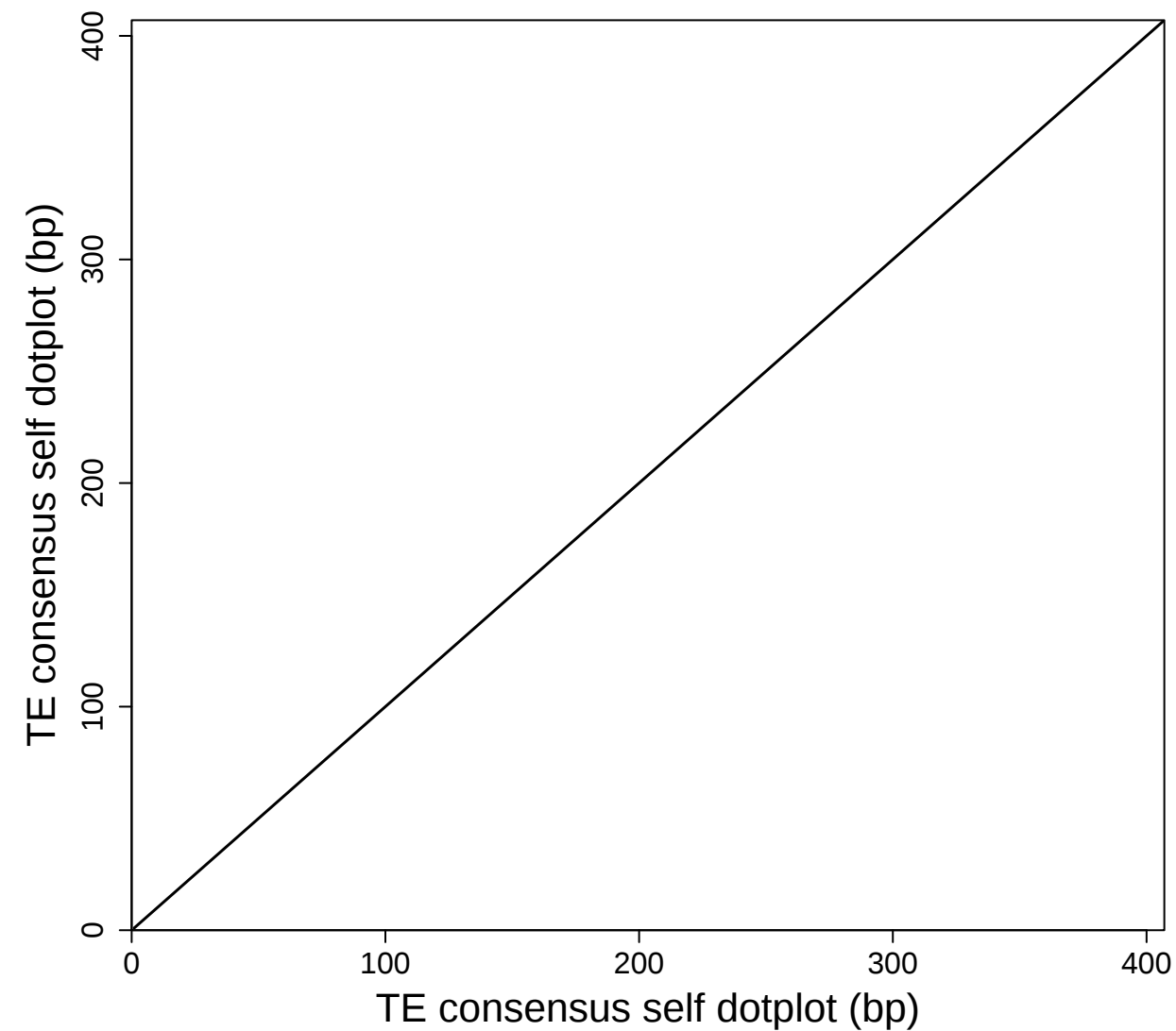
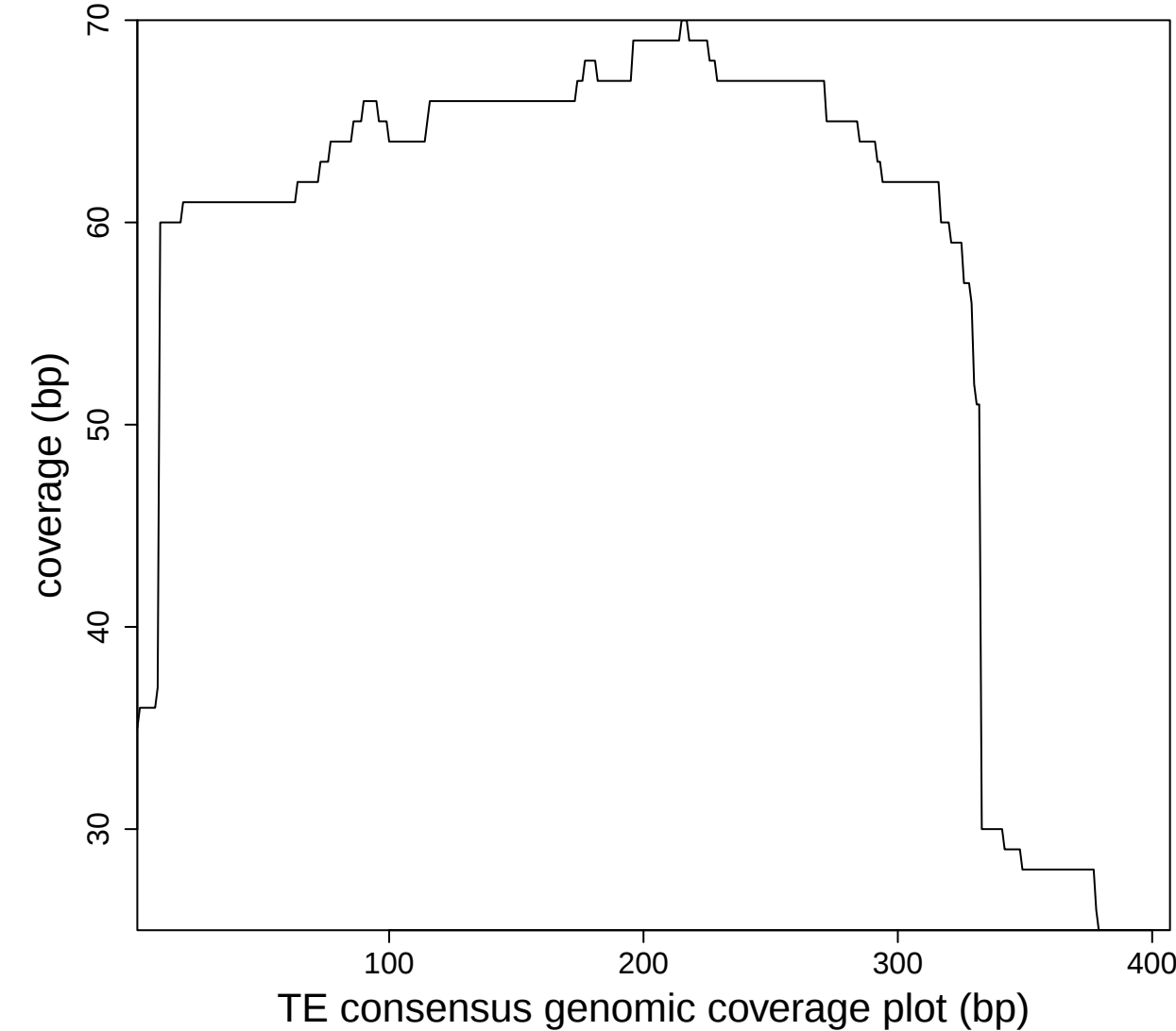
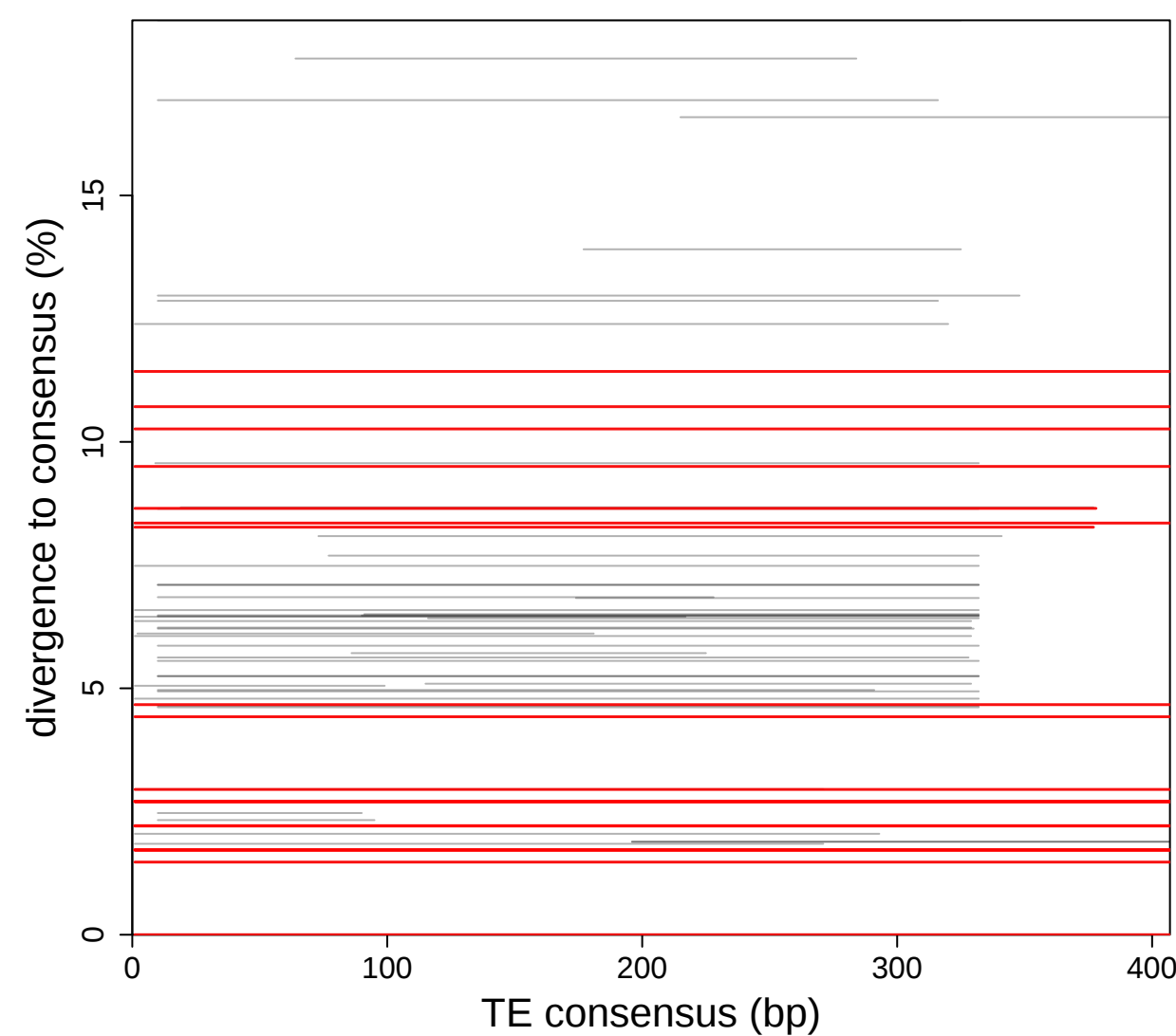


ed_uf.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_
size: 2566bp; fragments: 190; full length: 0 (>=2309.4bp)

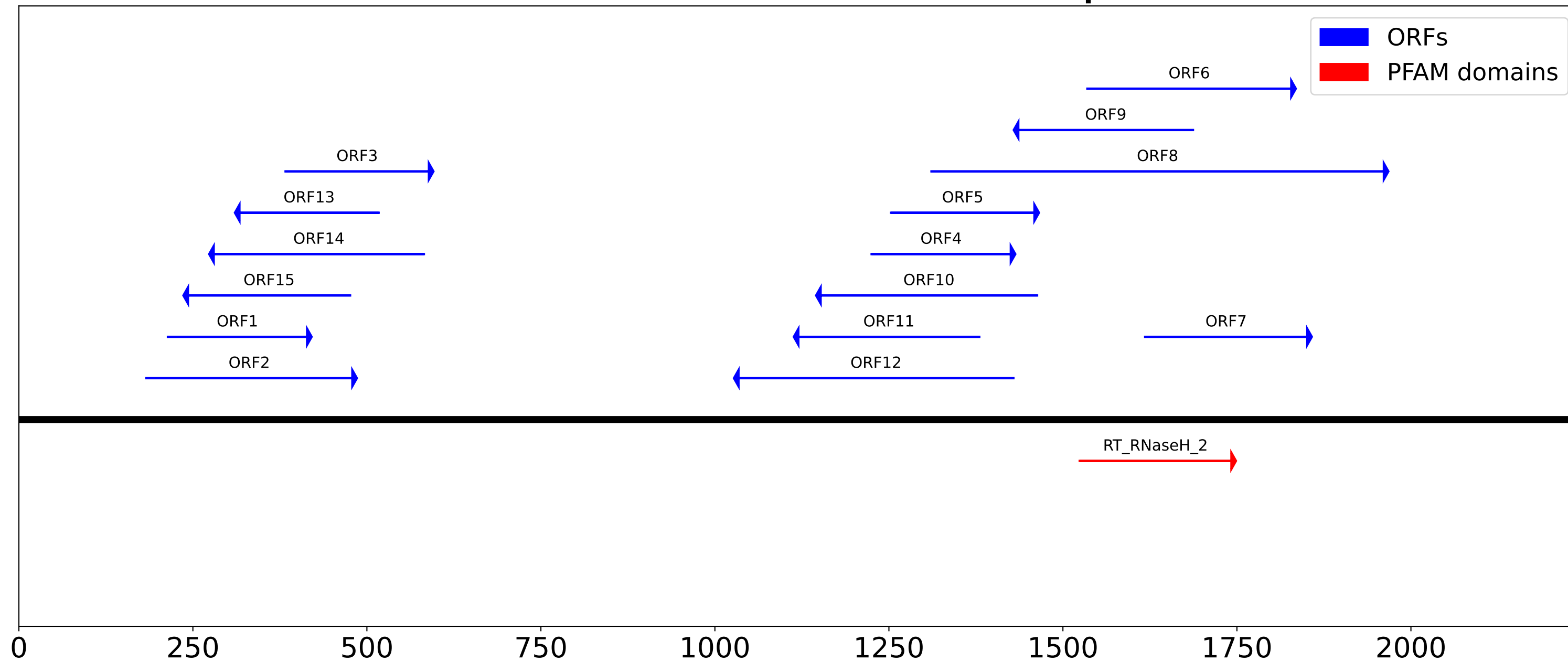
After TEtrimmer Extended plot Blue lines are boundaries



TE: ltr_1_family_349
size: 407bp; fragments: 74; full length: 24 (≥ 366.3 bp)



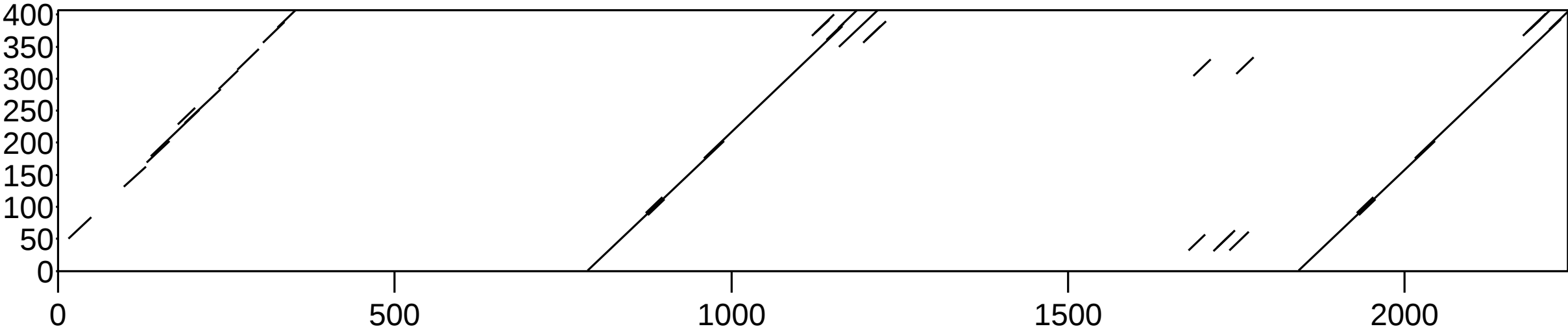
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)