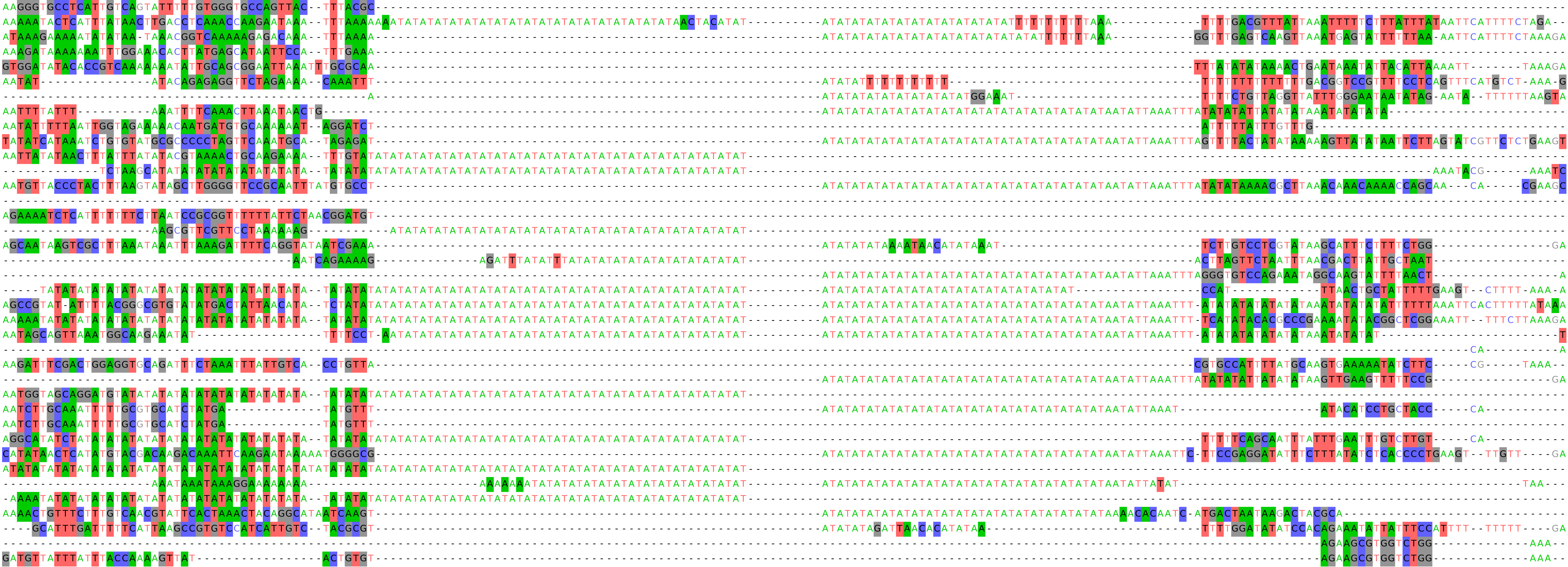


MSA length = 5594

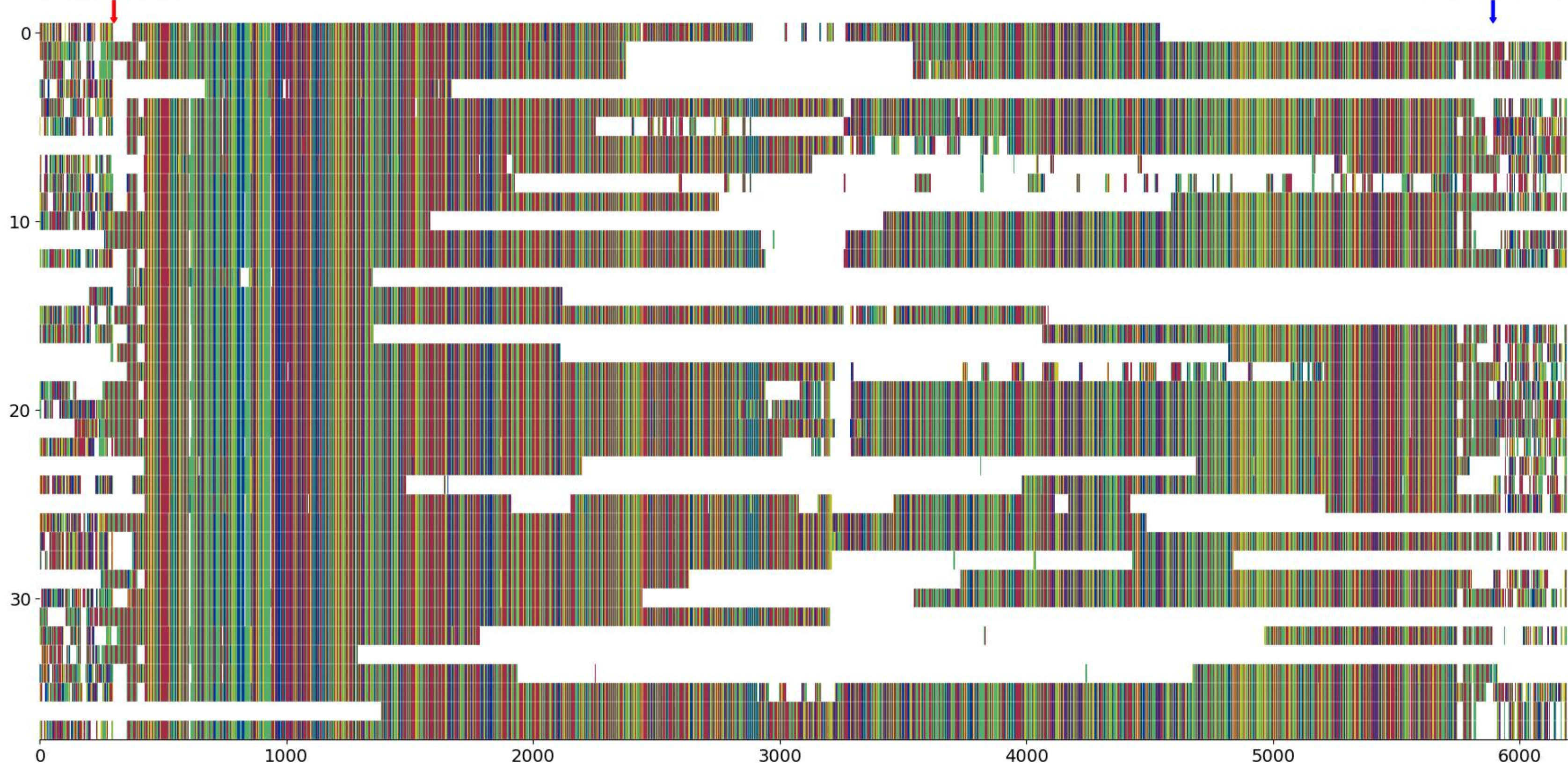
Start crop Point

End crop Point



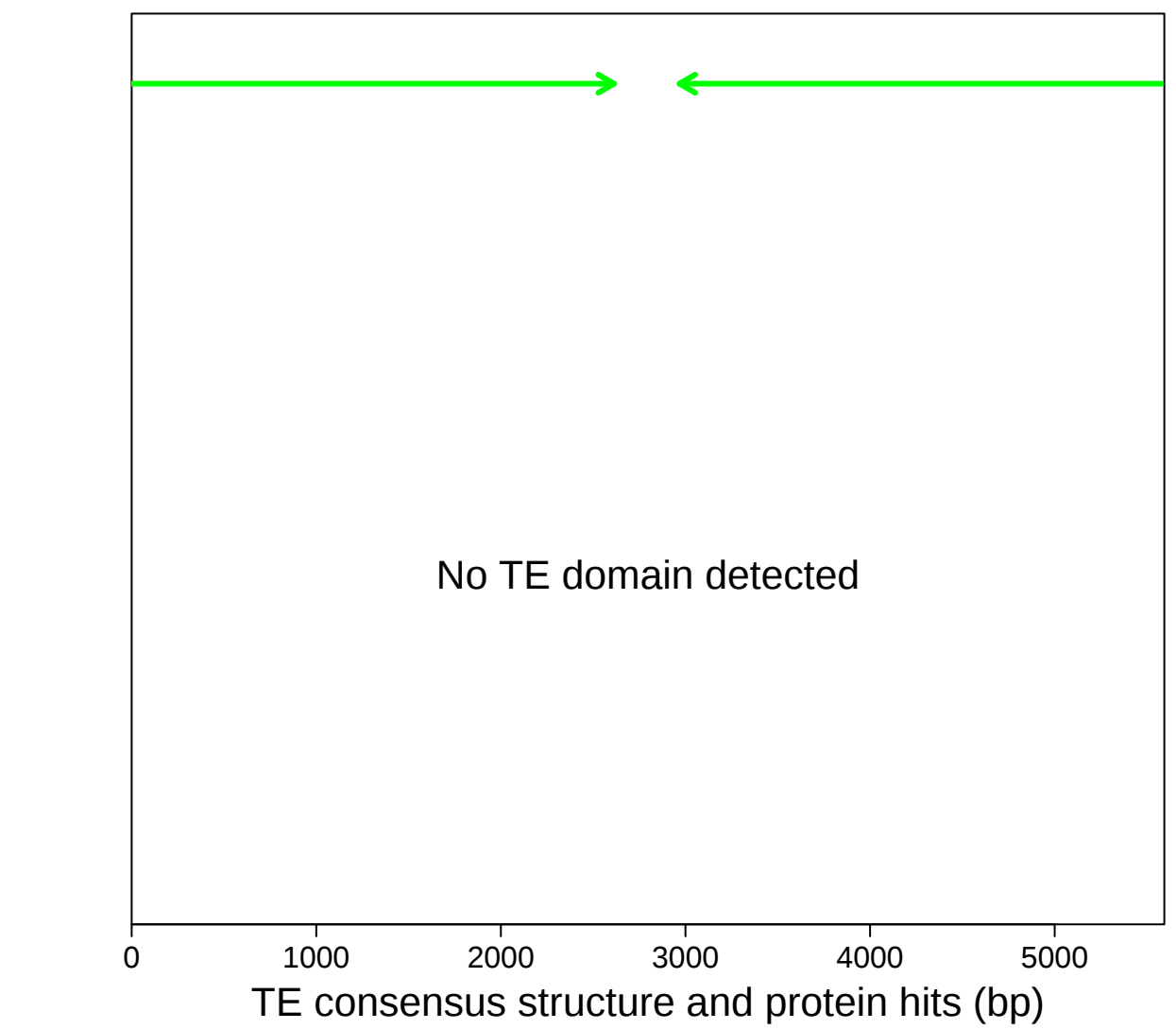
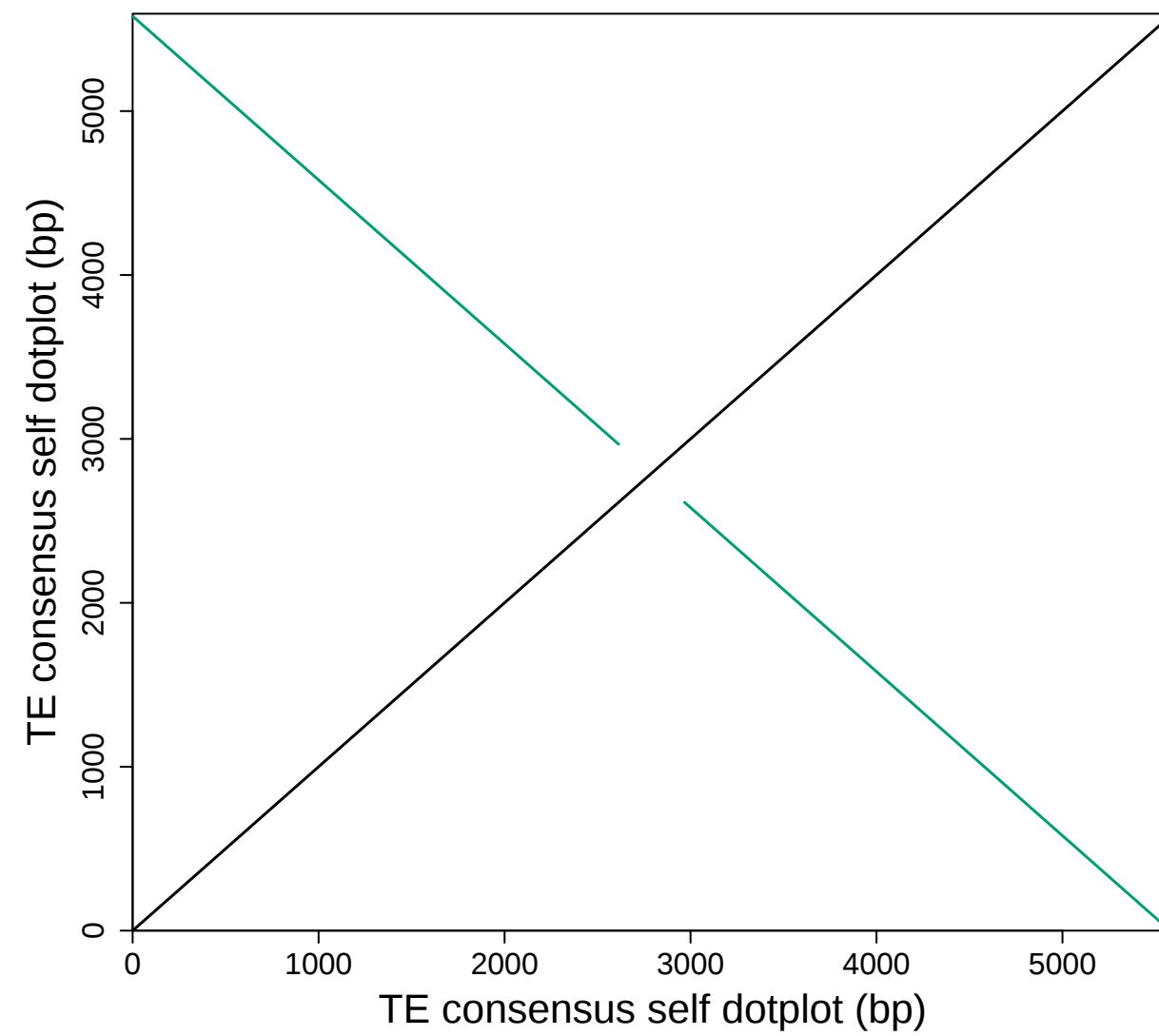
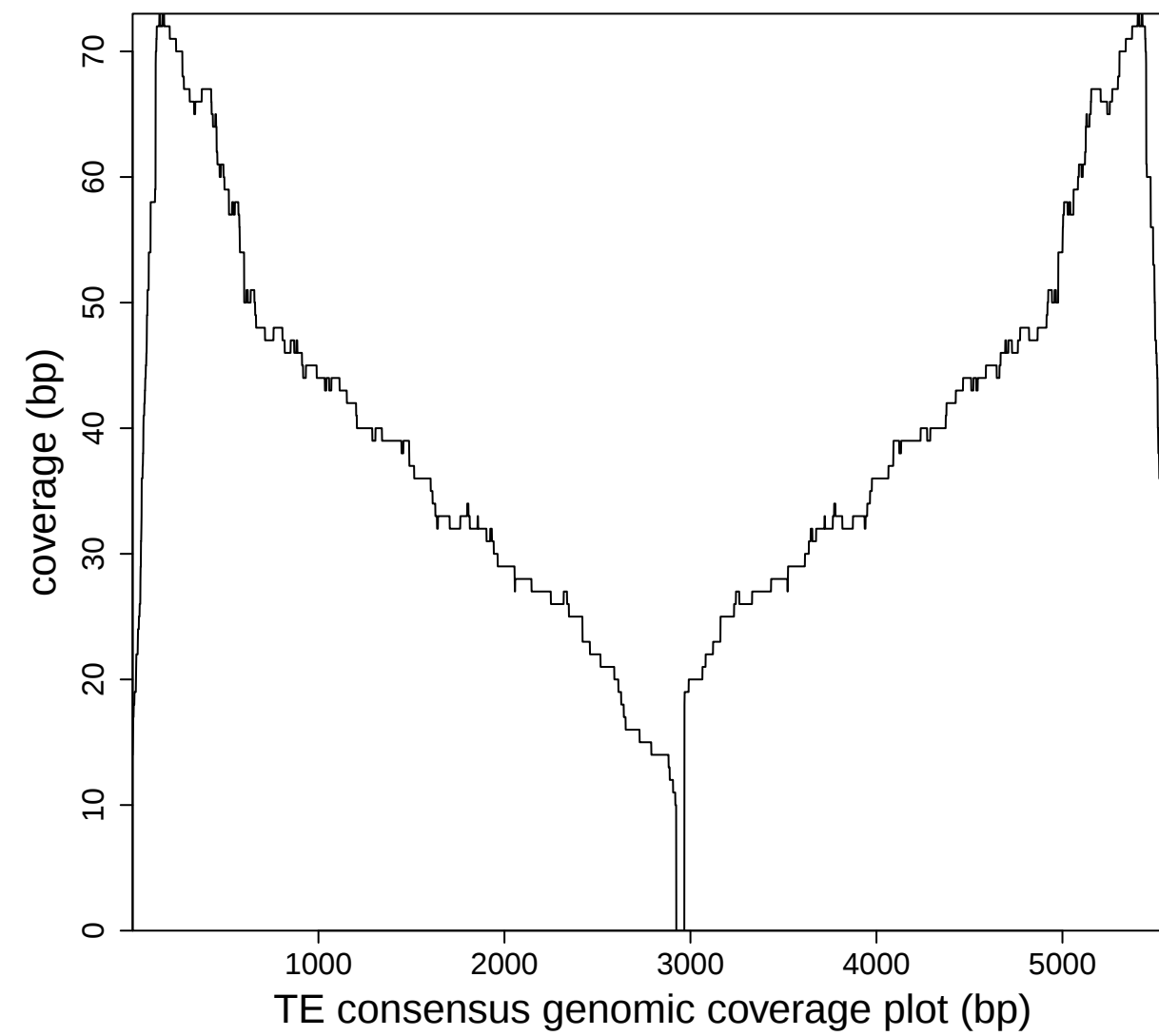
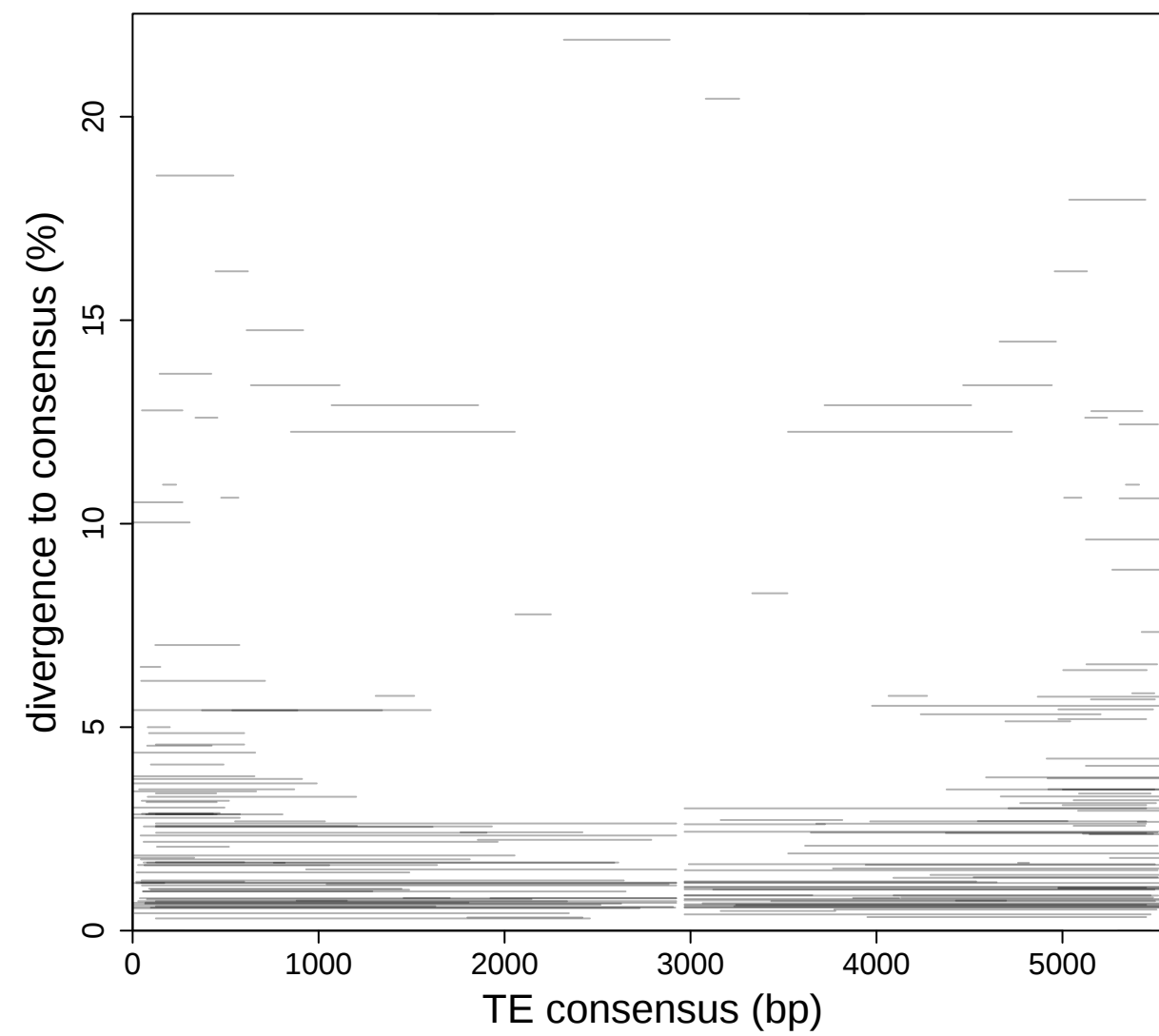
Start crop Point

End crop Point



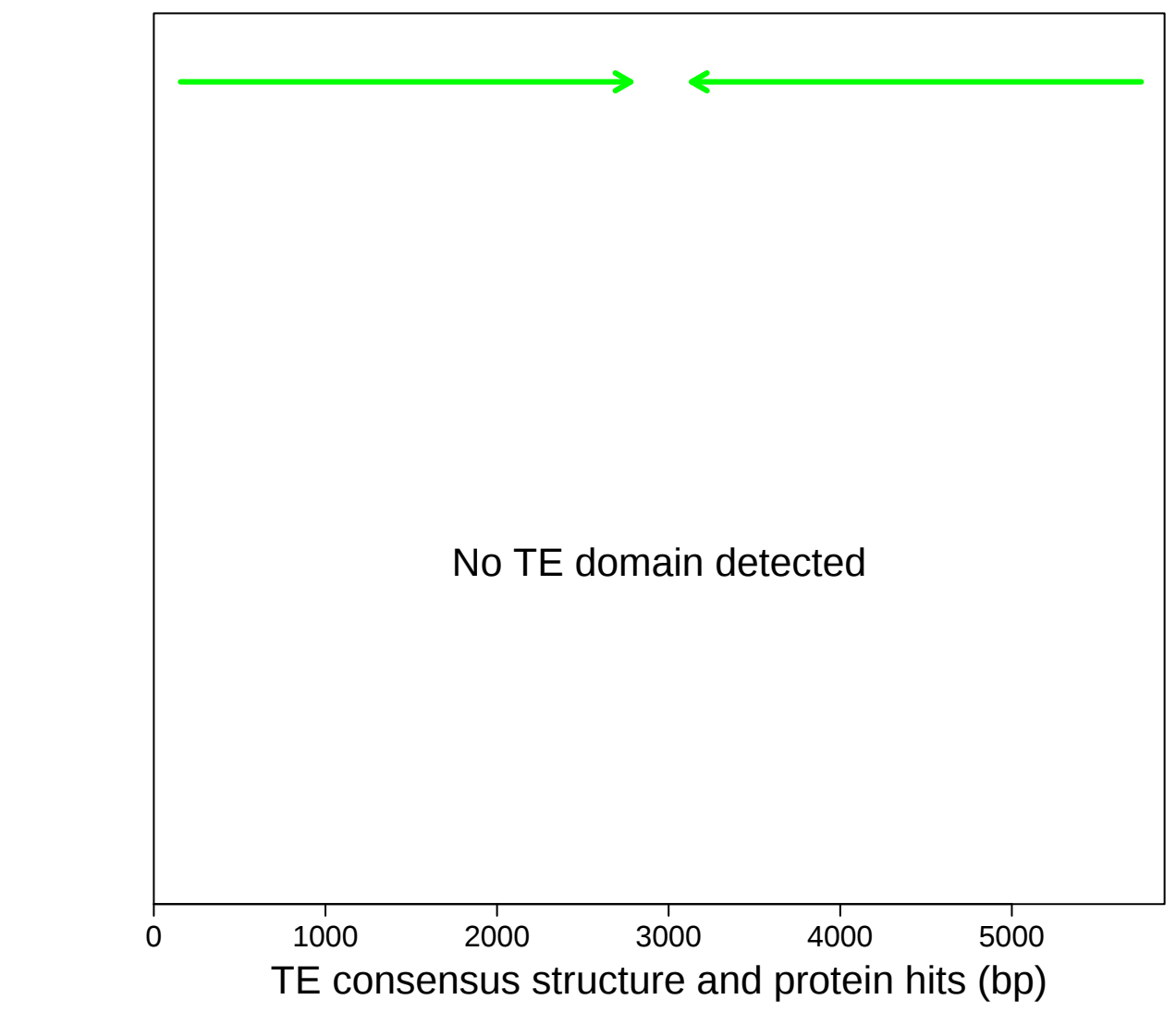
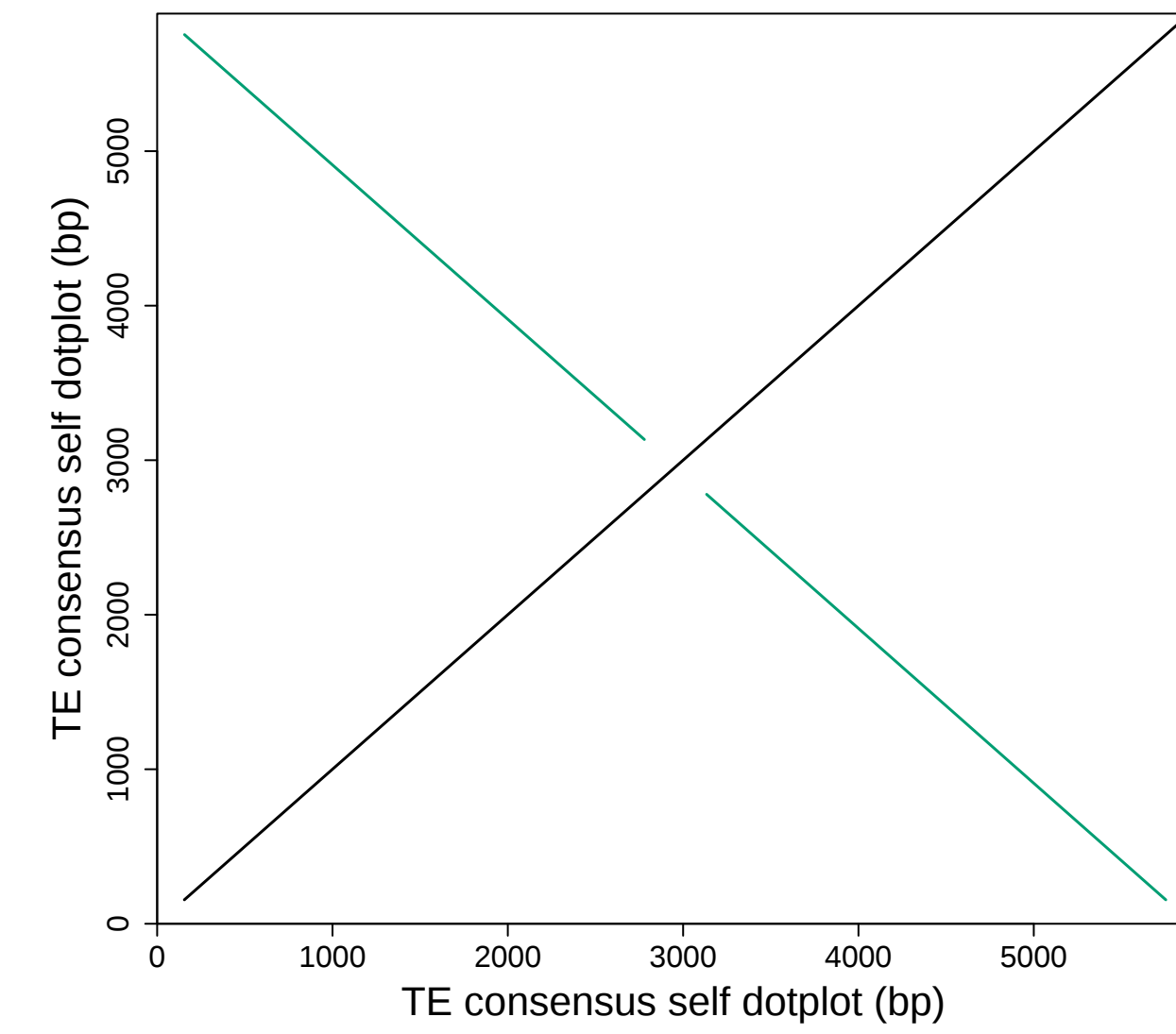
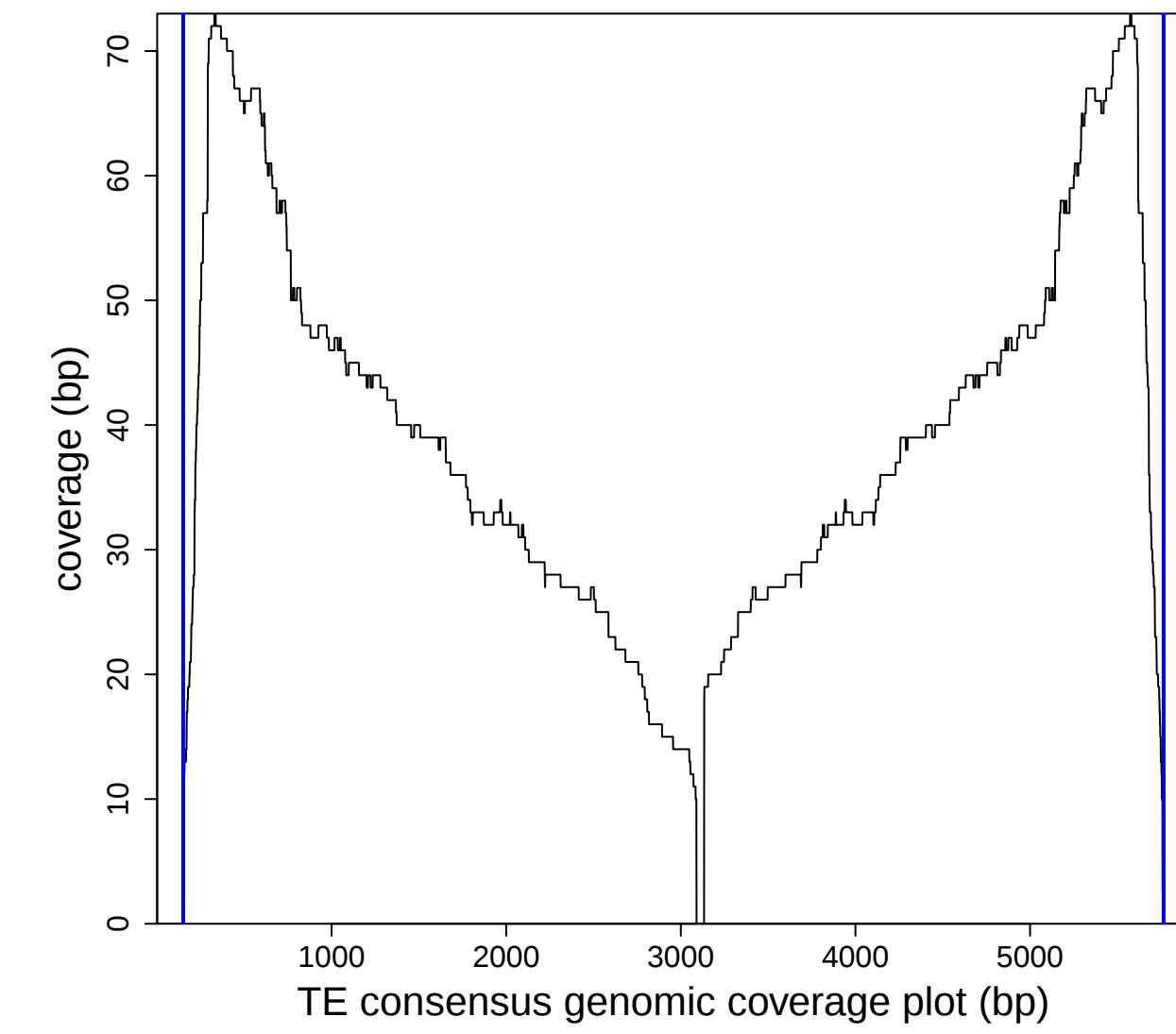
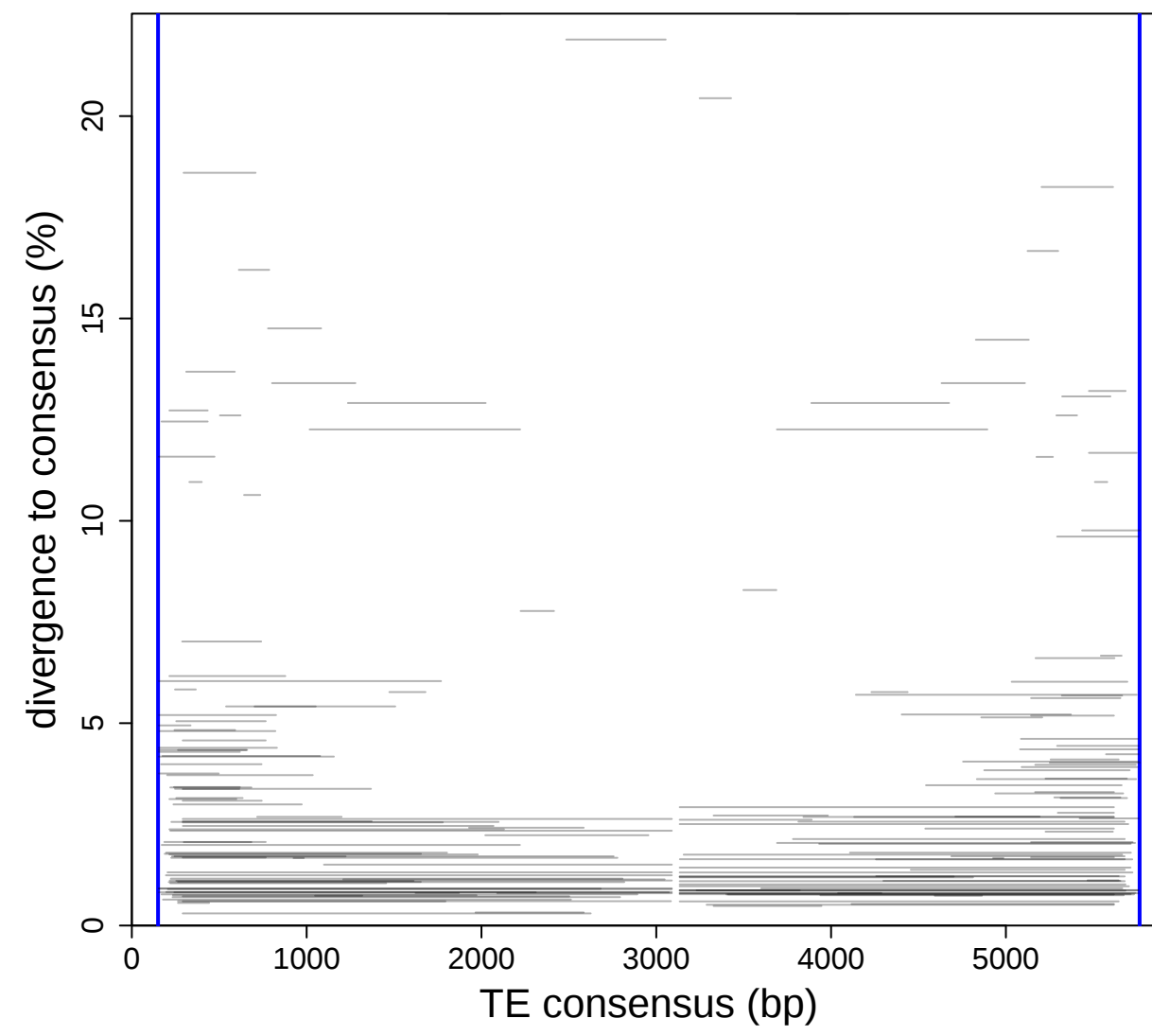
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 5594bp; fragments: 194; full length: 0 (>=5034.6bp)

After TETrimmer 5594 bp

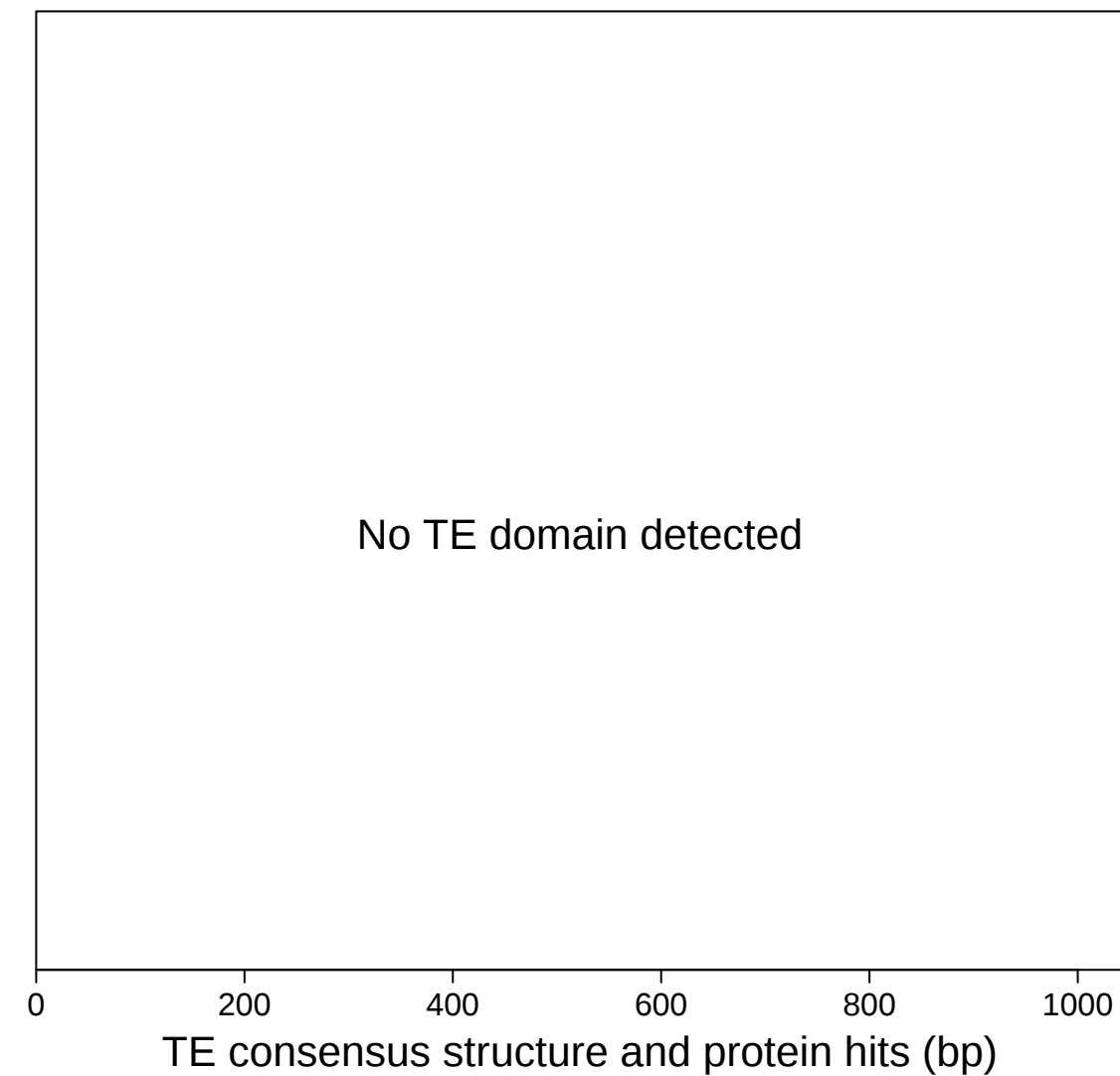
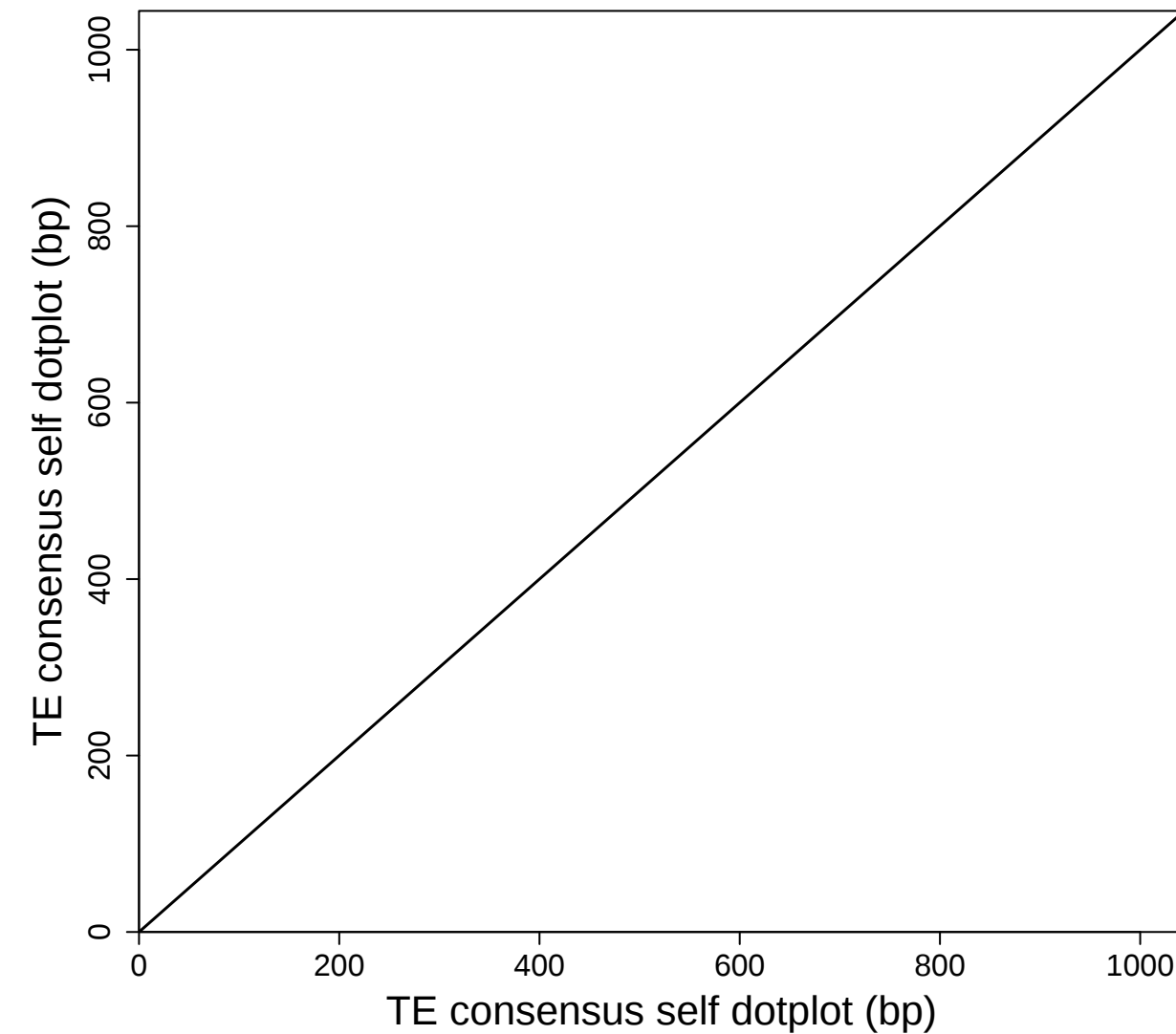
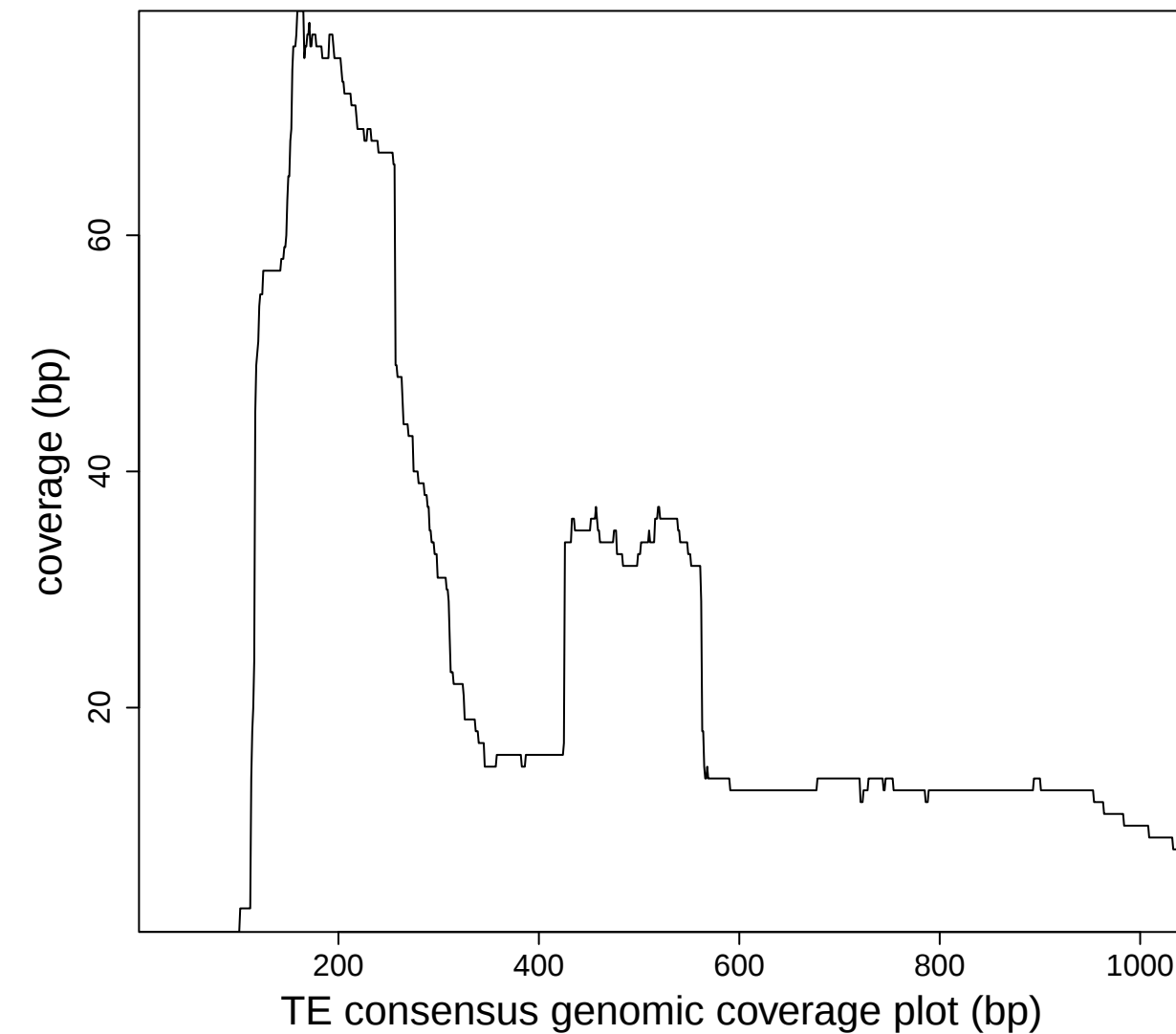
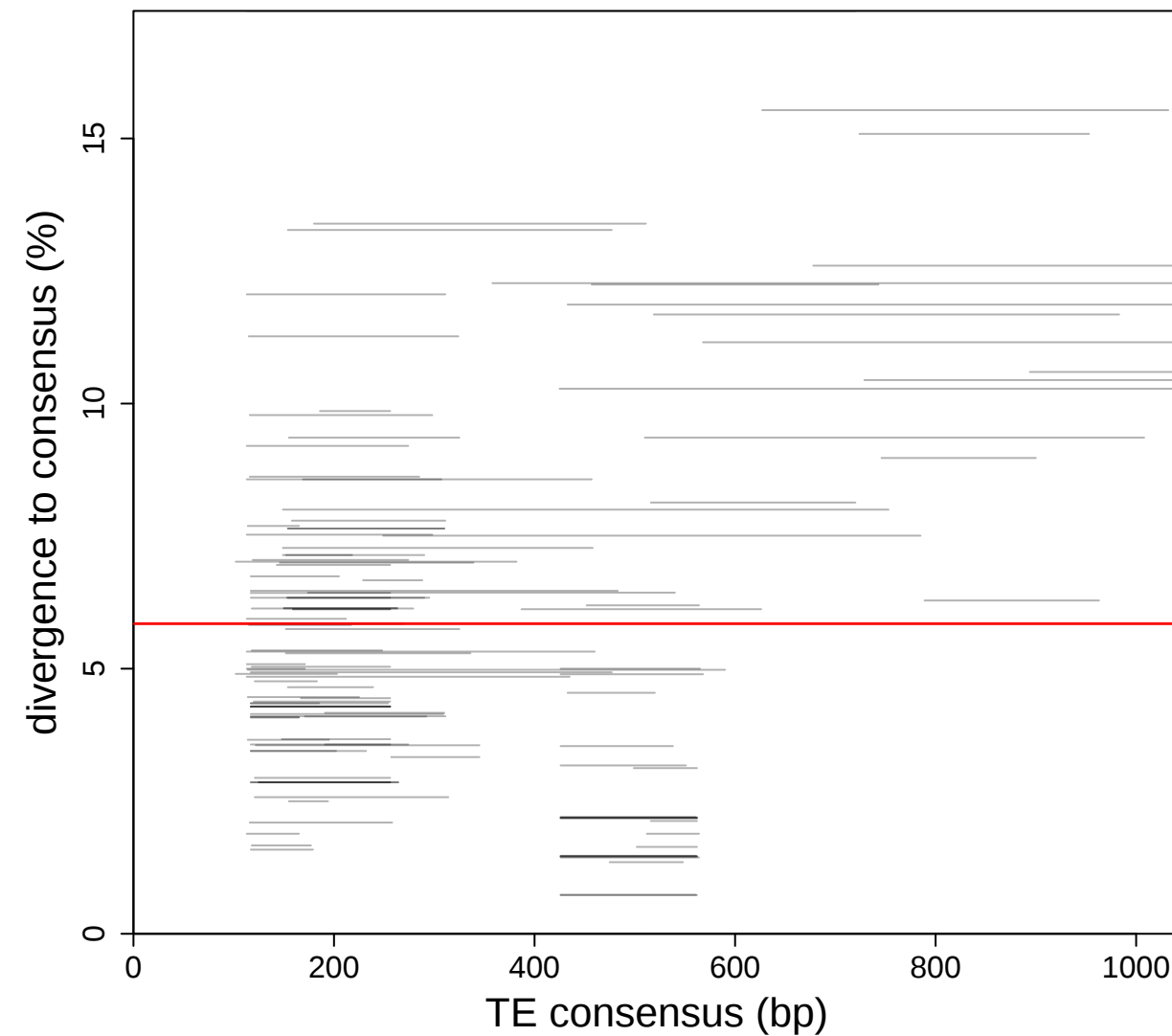


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 5915bp; fragments: 192; full length: 0 (>=5323.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

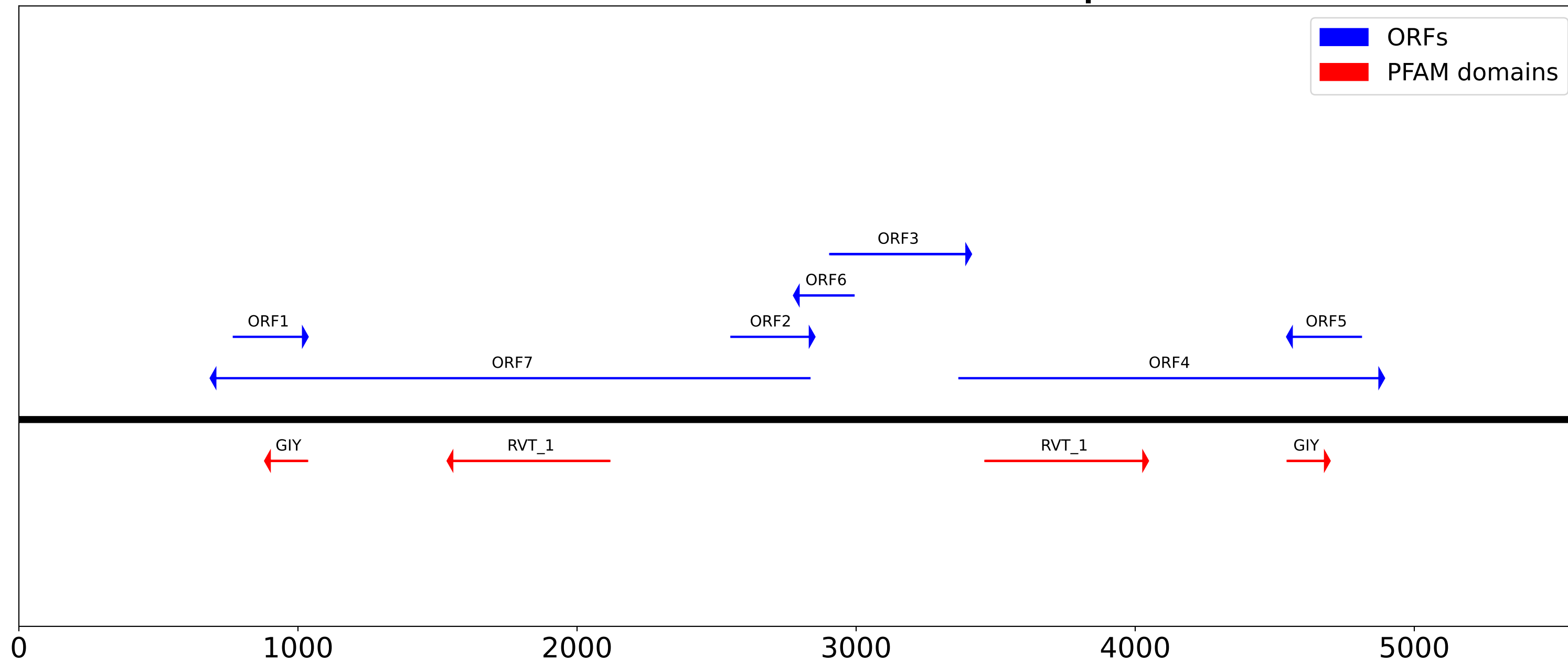


TE: rnd_5_family_998
size: 1044bp; fragments: 130; full length: 1 (≥ 939.6 bp)

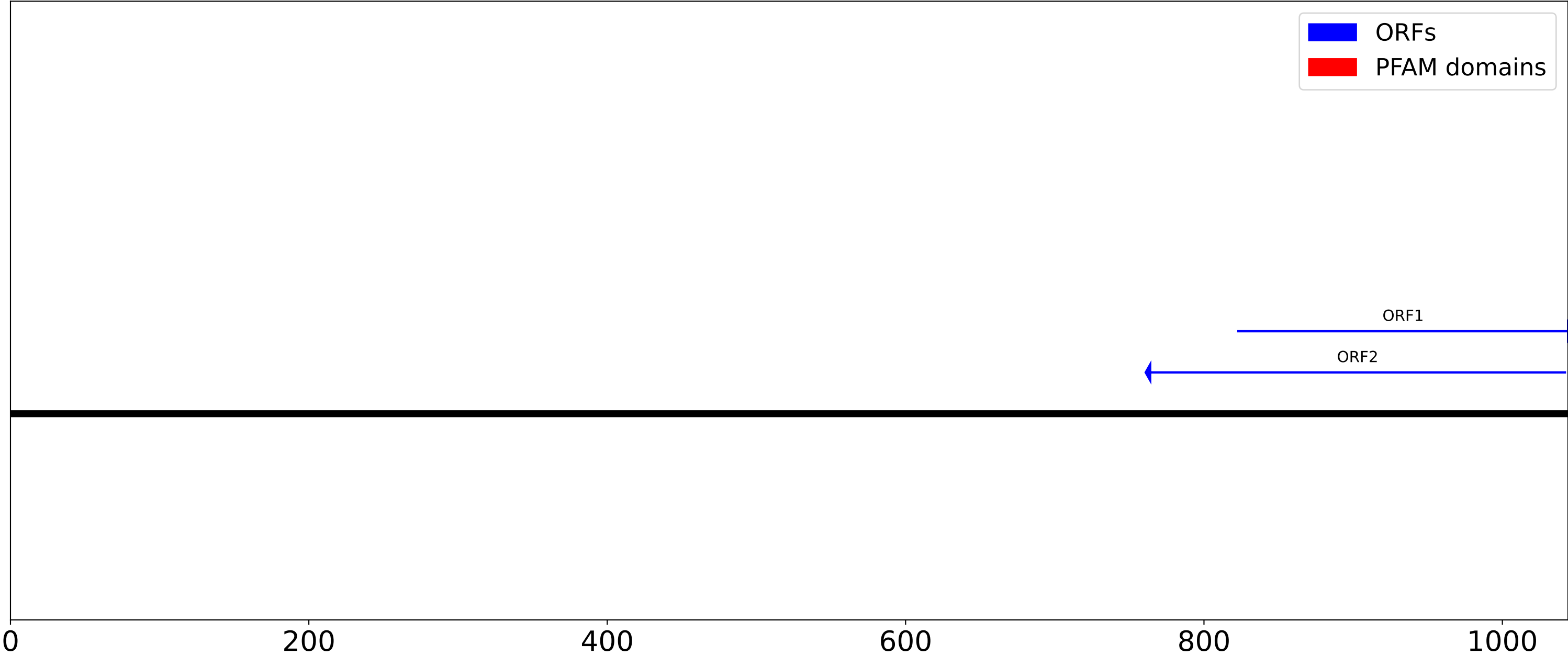


Before TEtrimmer 1044 bp

After TEtrimmer ORF and PFAM domain plot



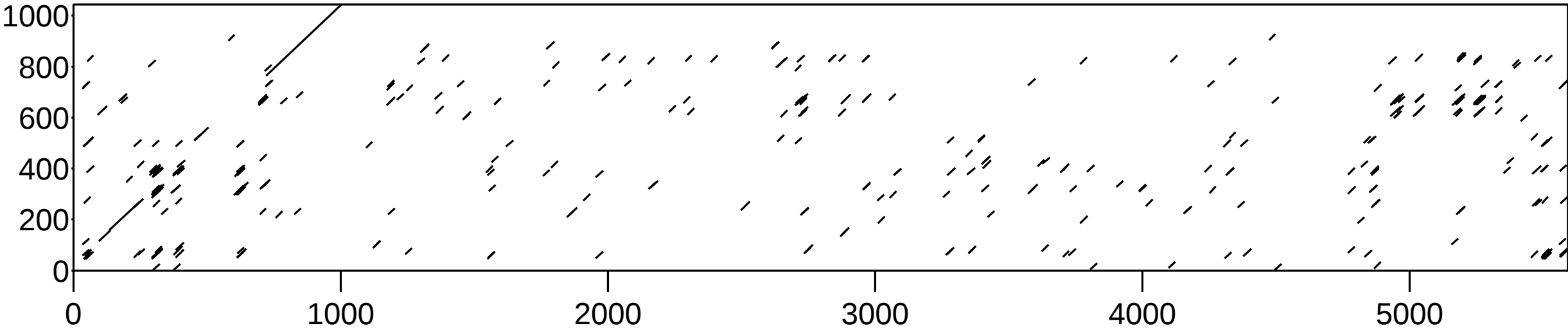
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)