

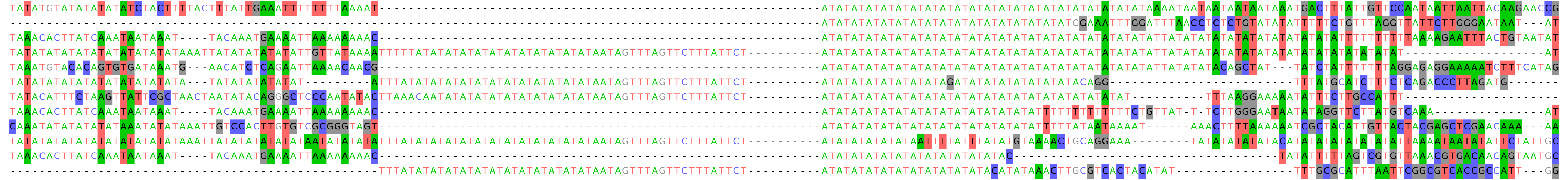
Start crop Point



End crop Point



MSA length = 2542



Start crop Point

End crop Point

0

10

0

500

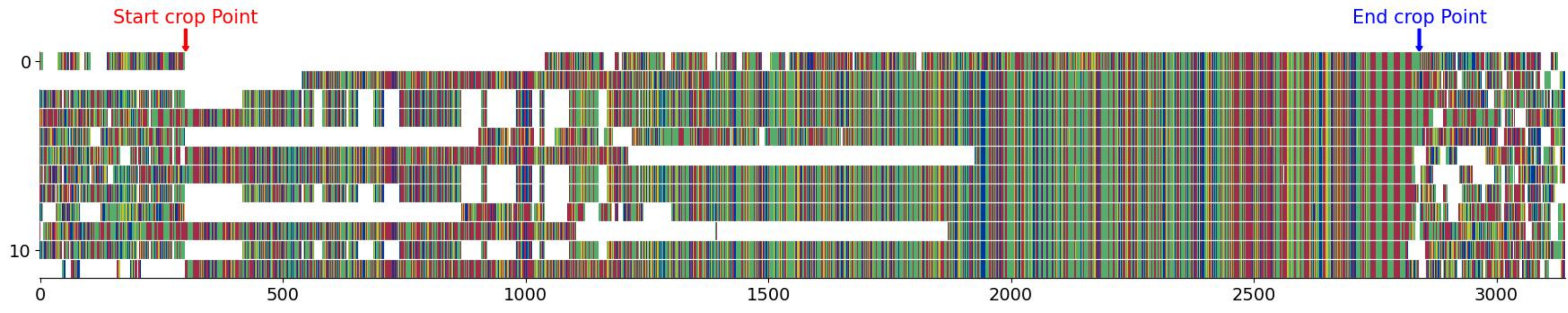
1000

1500

2000

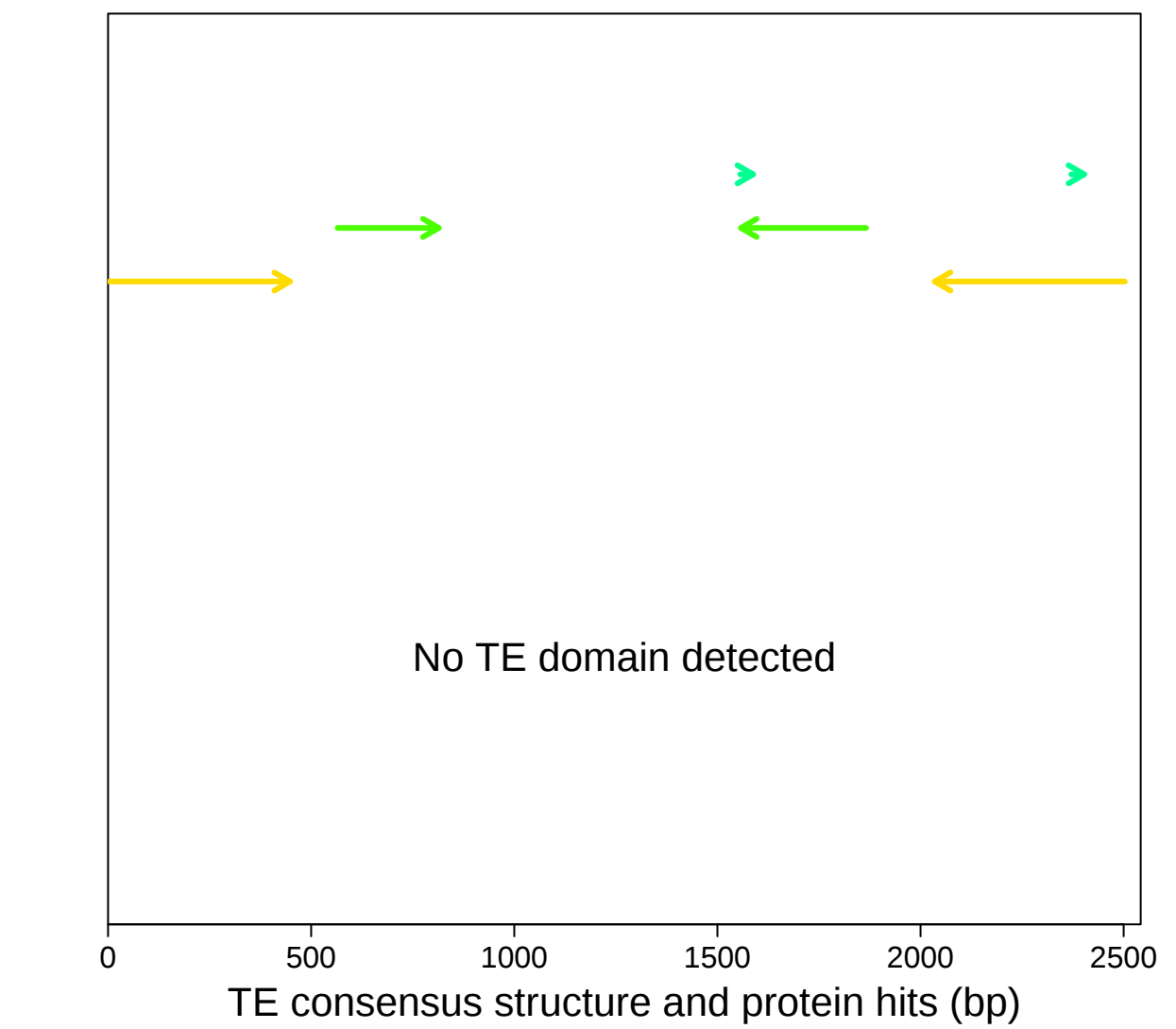
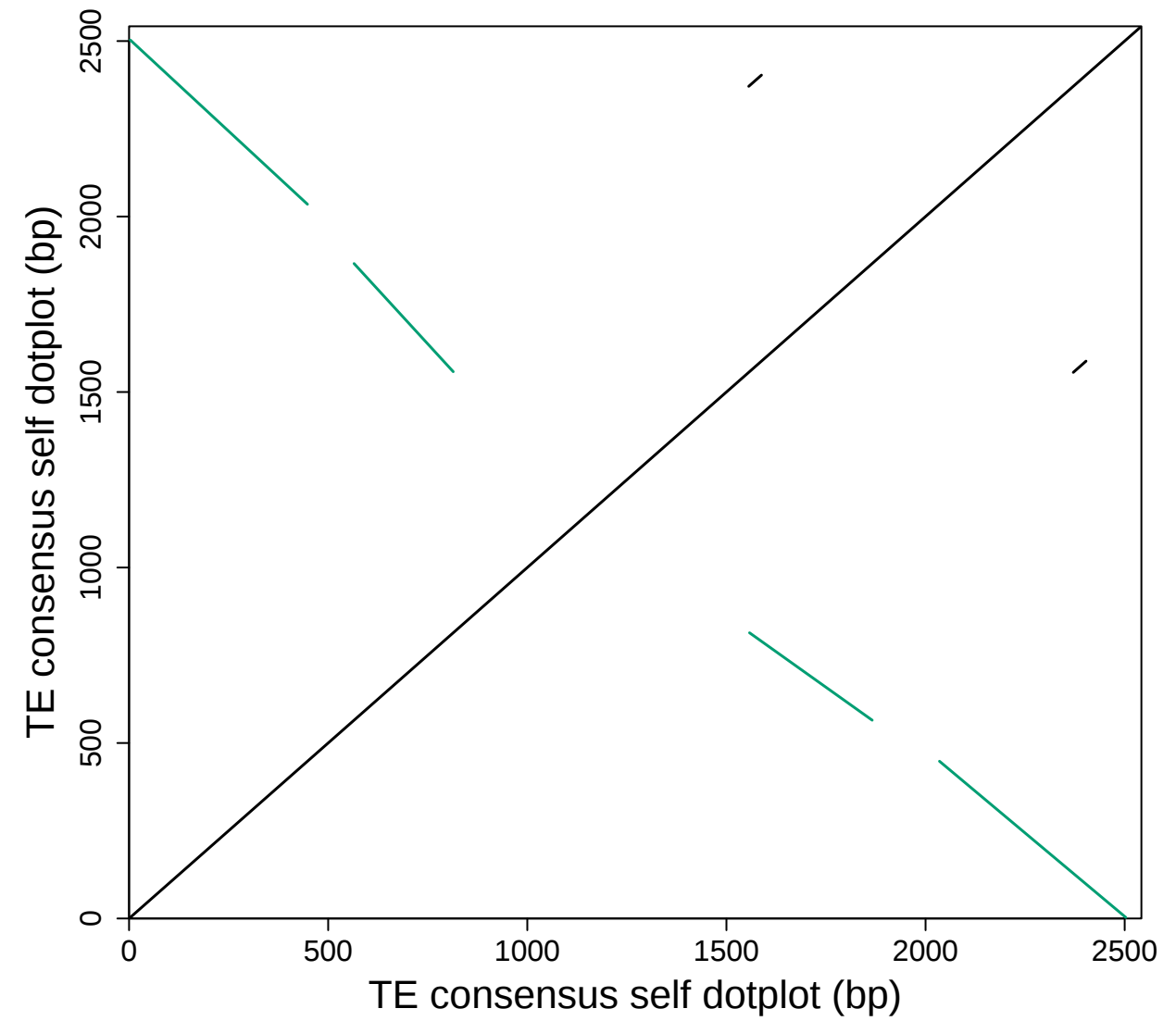
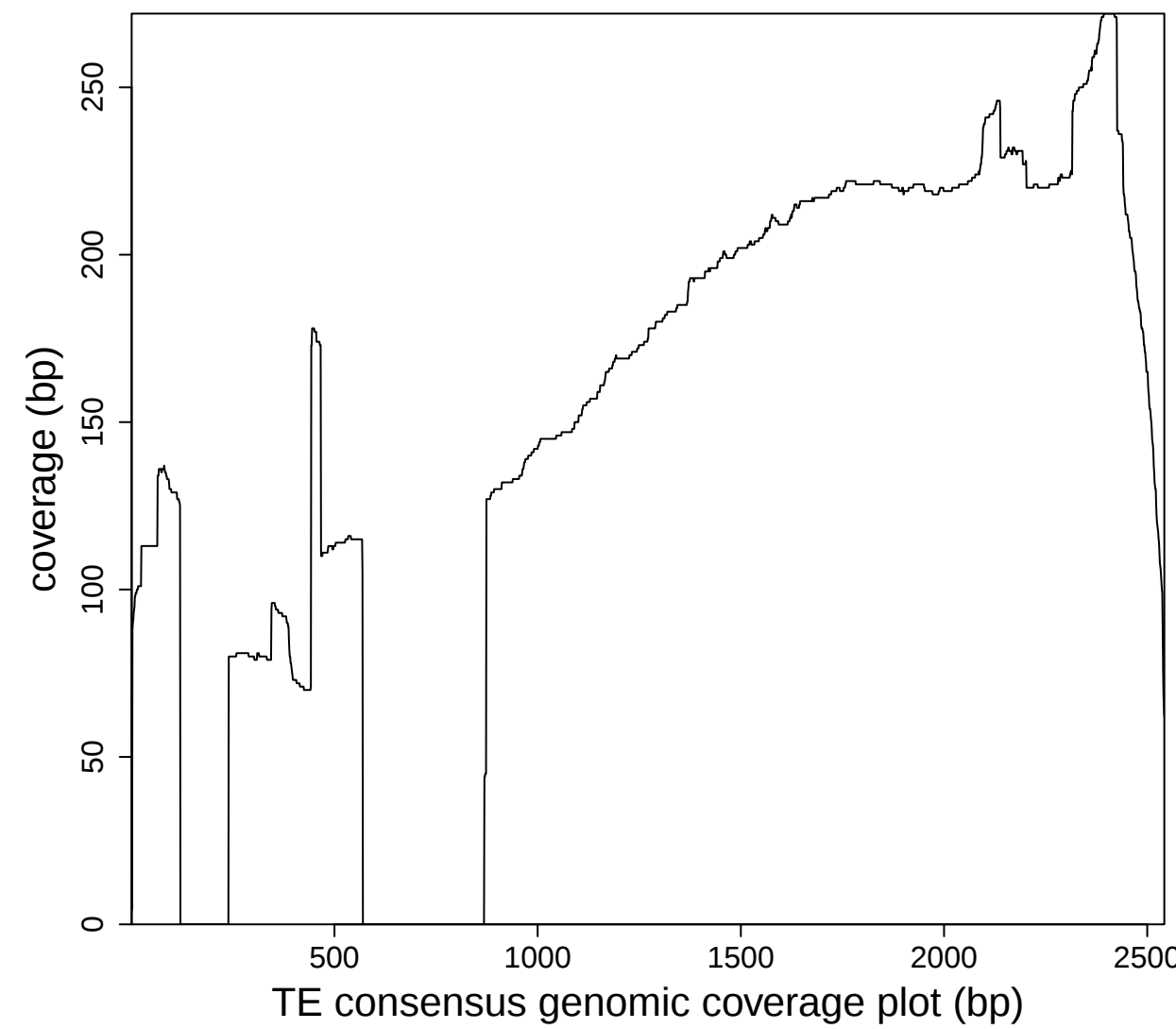
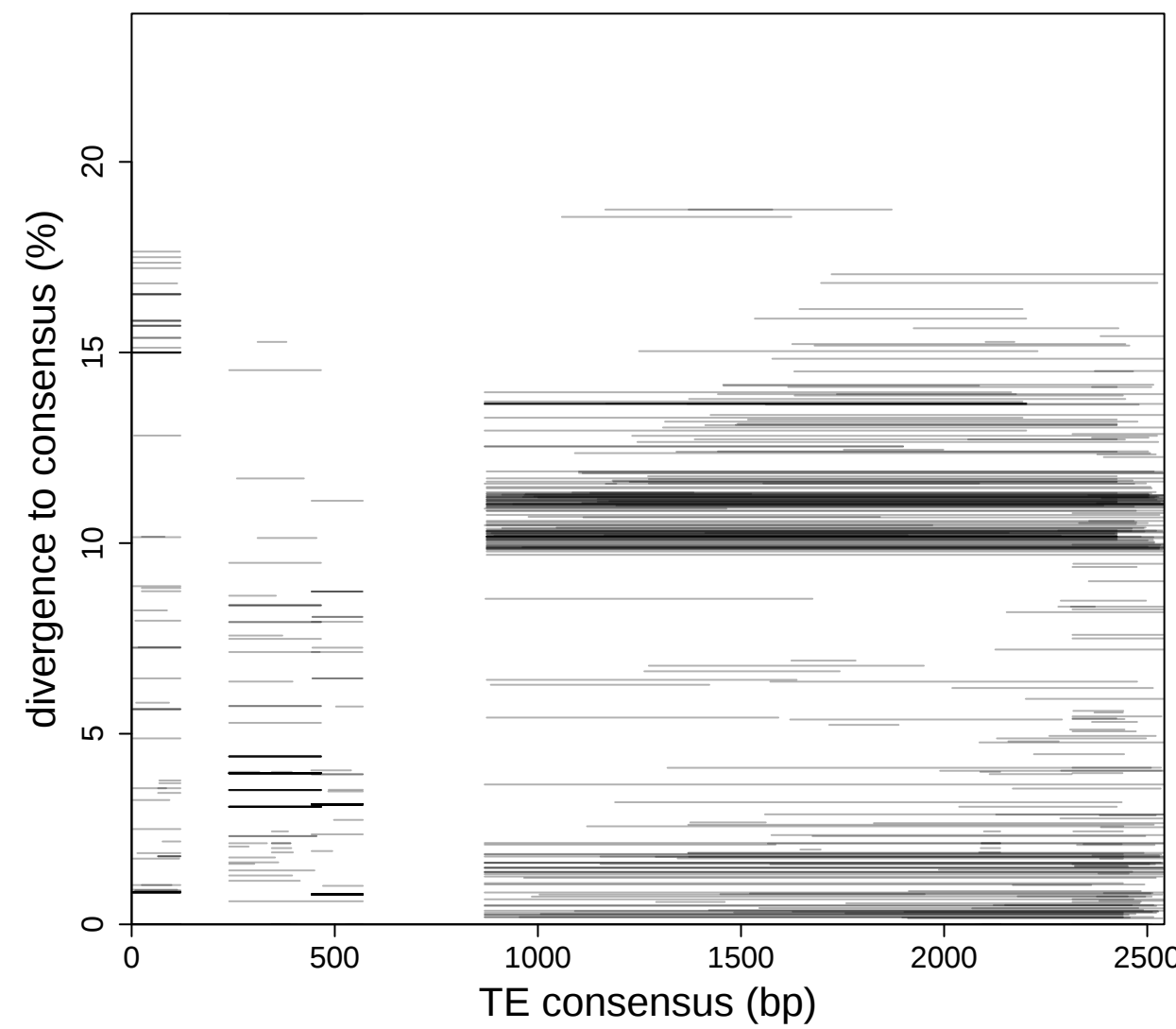
2500

3000



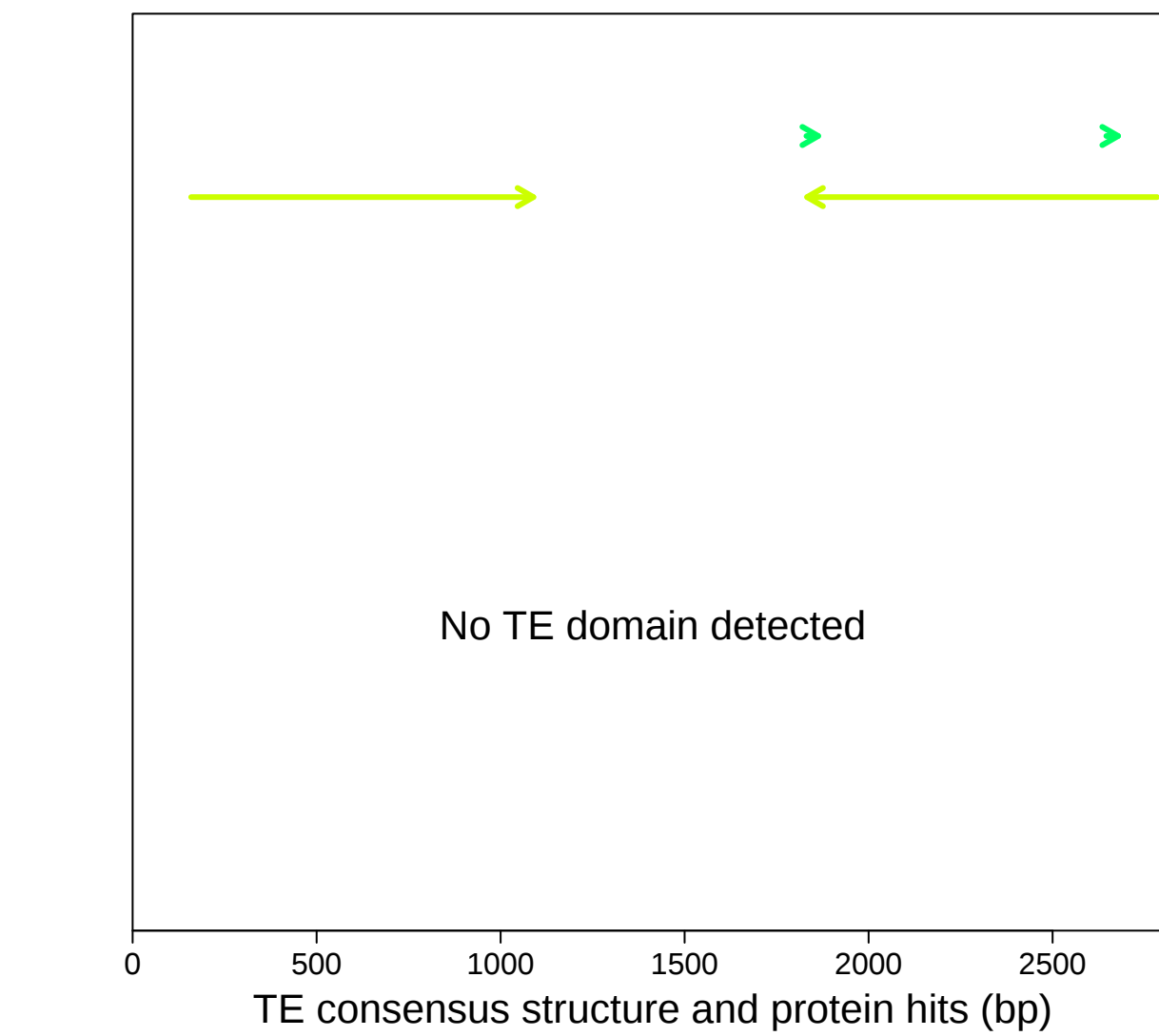
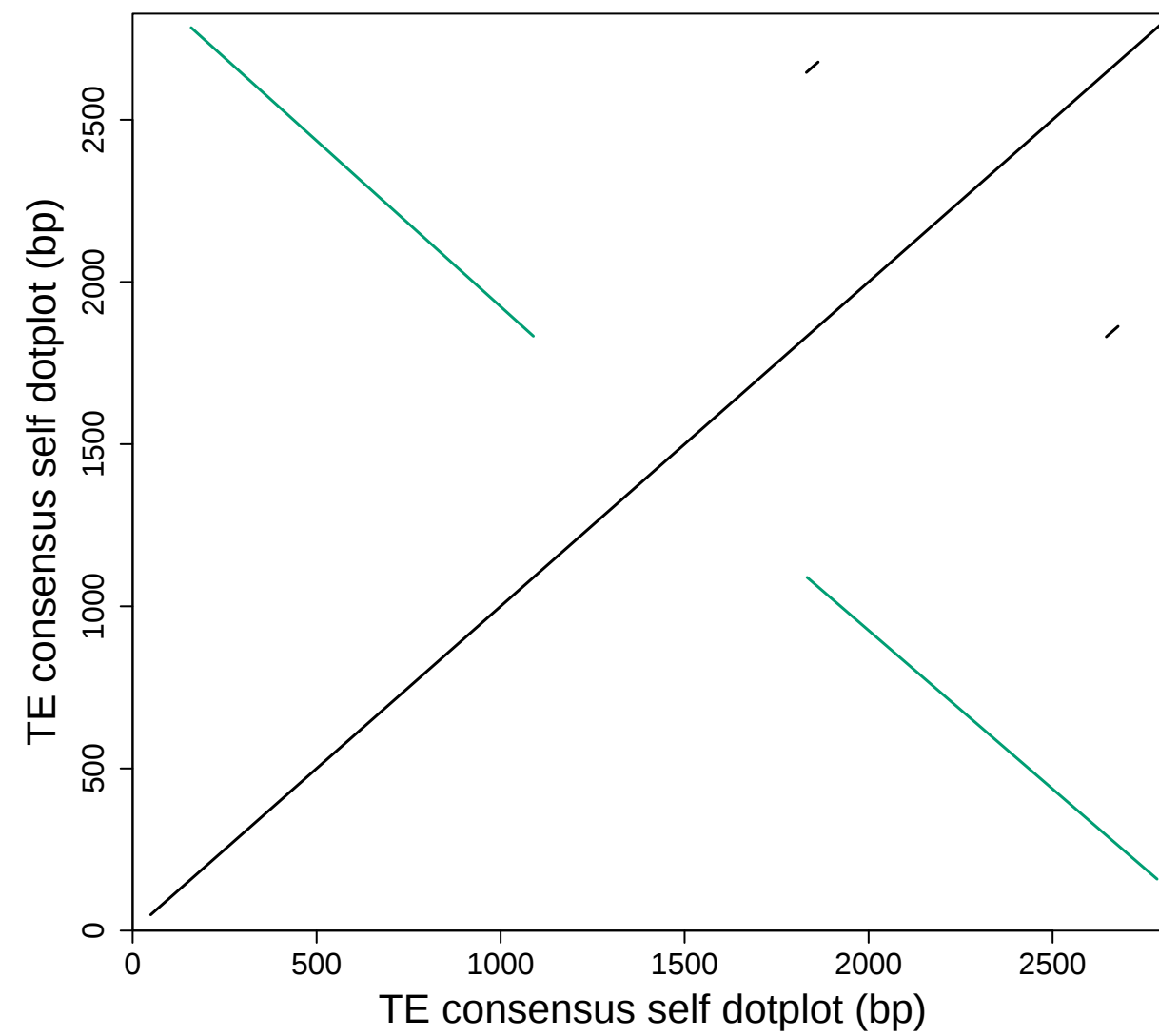
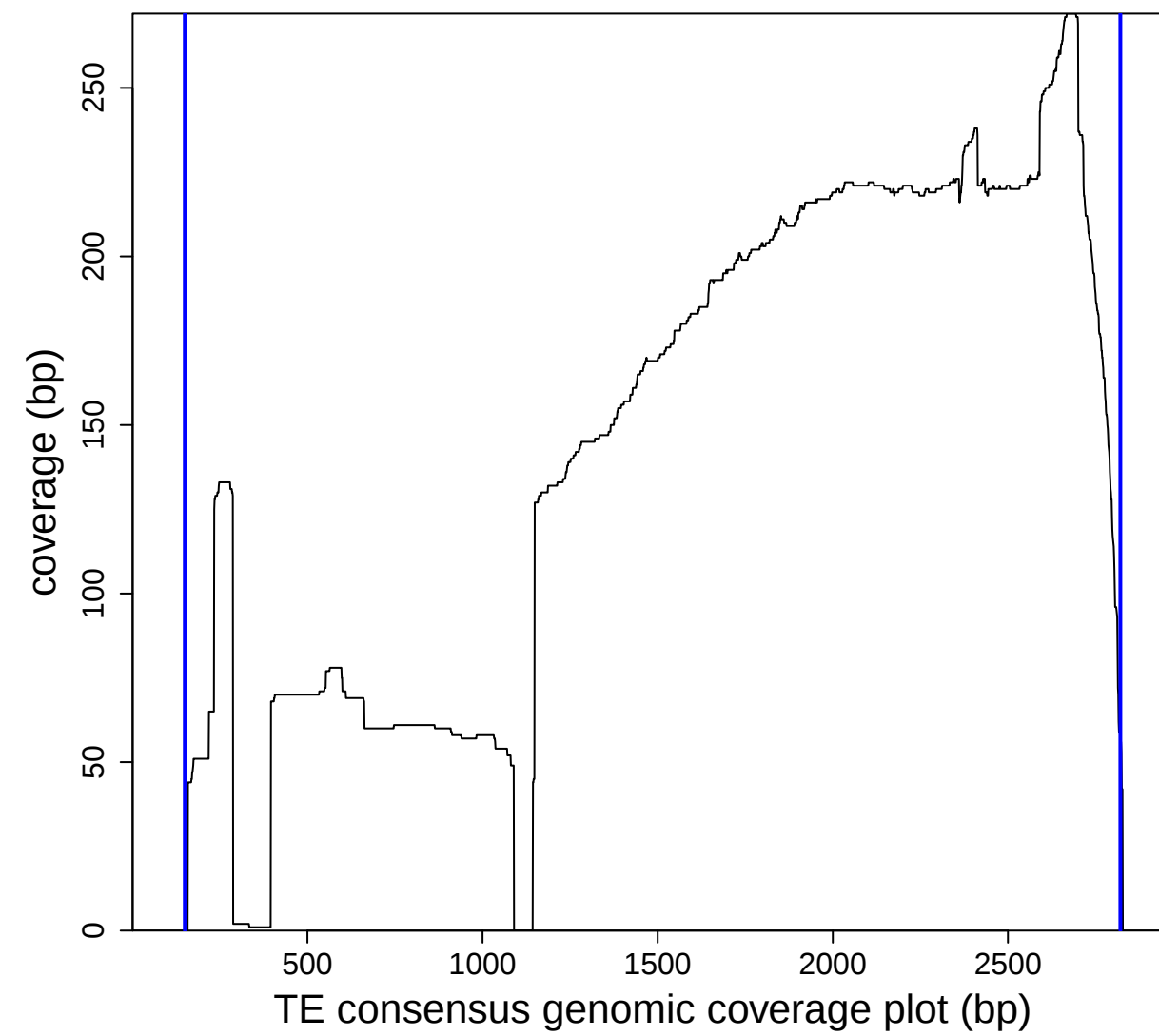
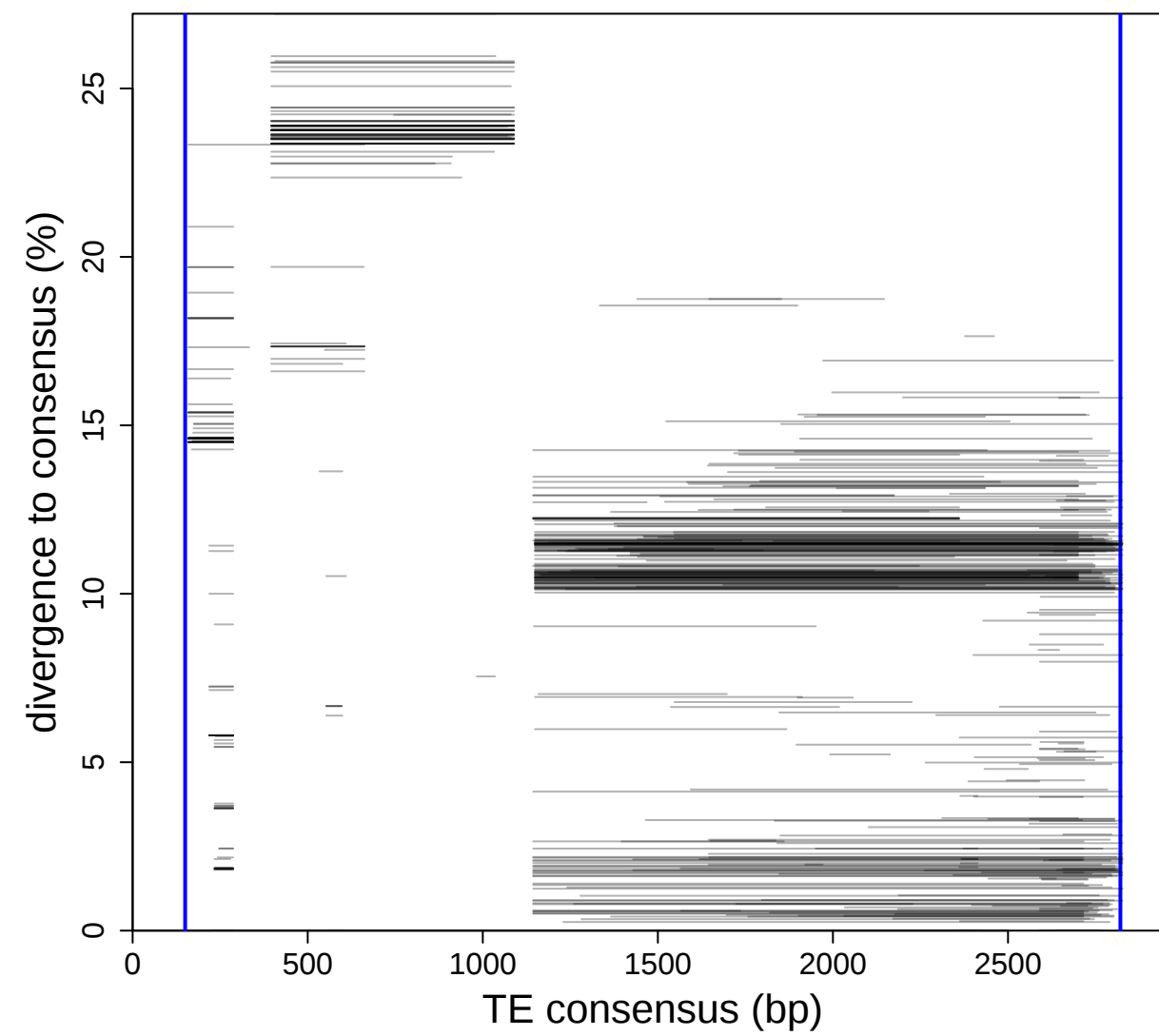
bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 2542bp; fragments: 694; full length: 0 (>=2287.8bp)

After TETrimmer 2542 bp

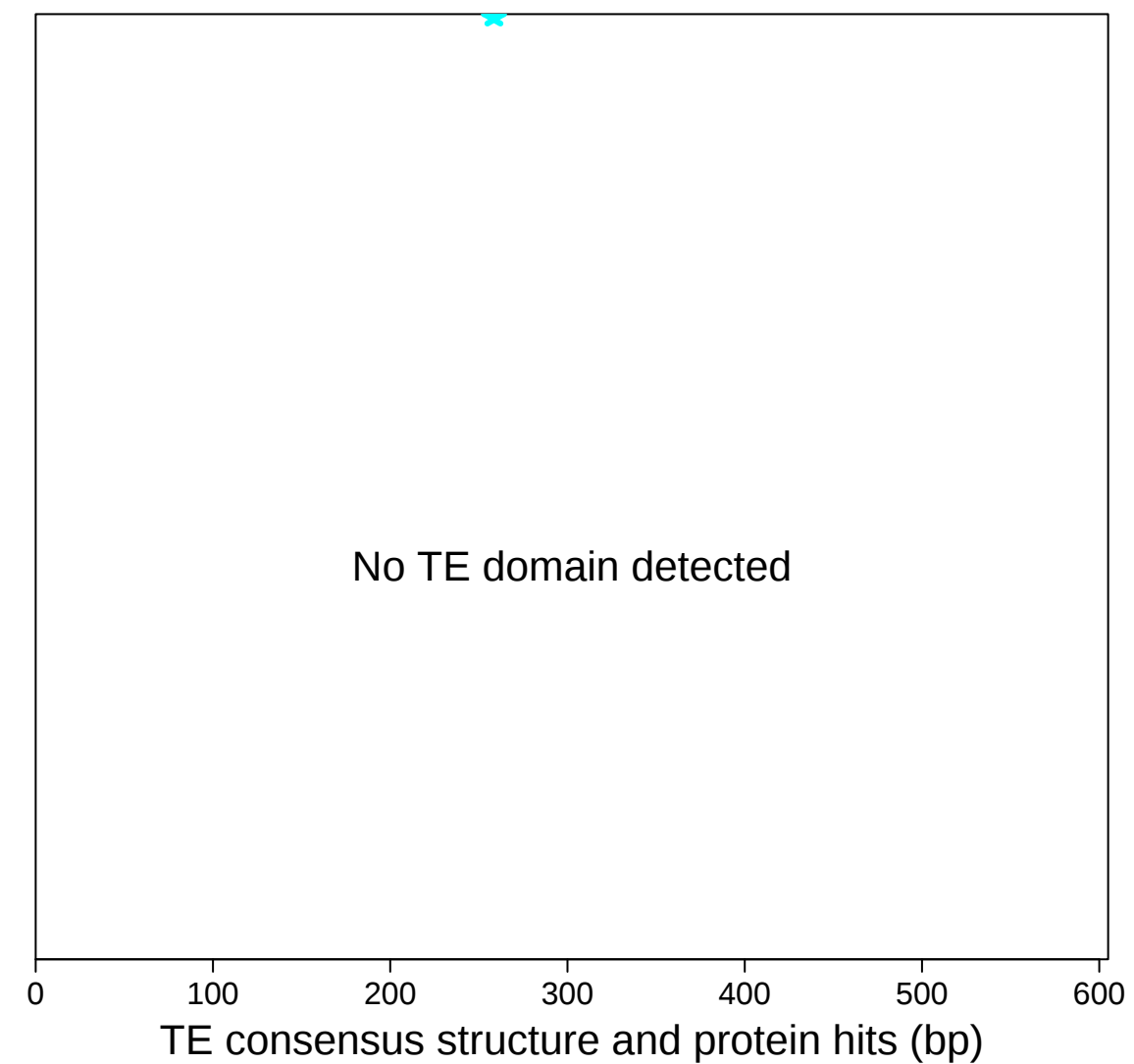
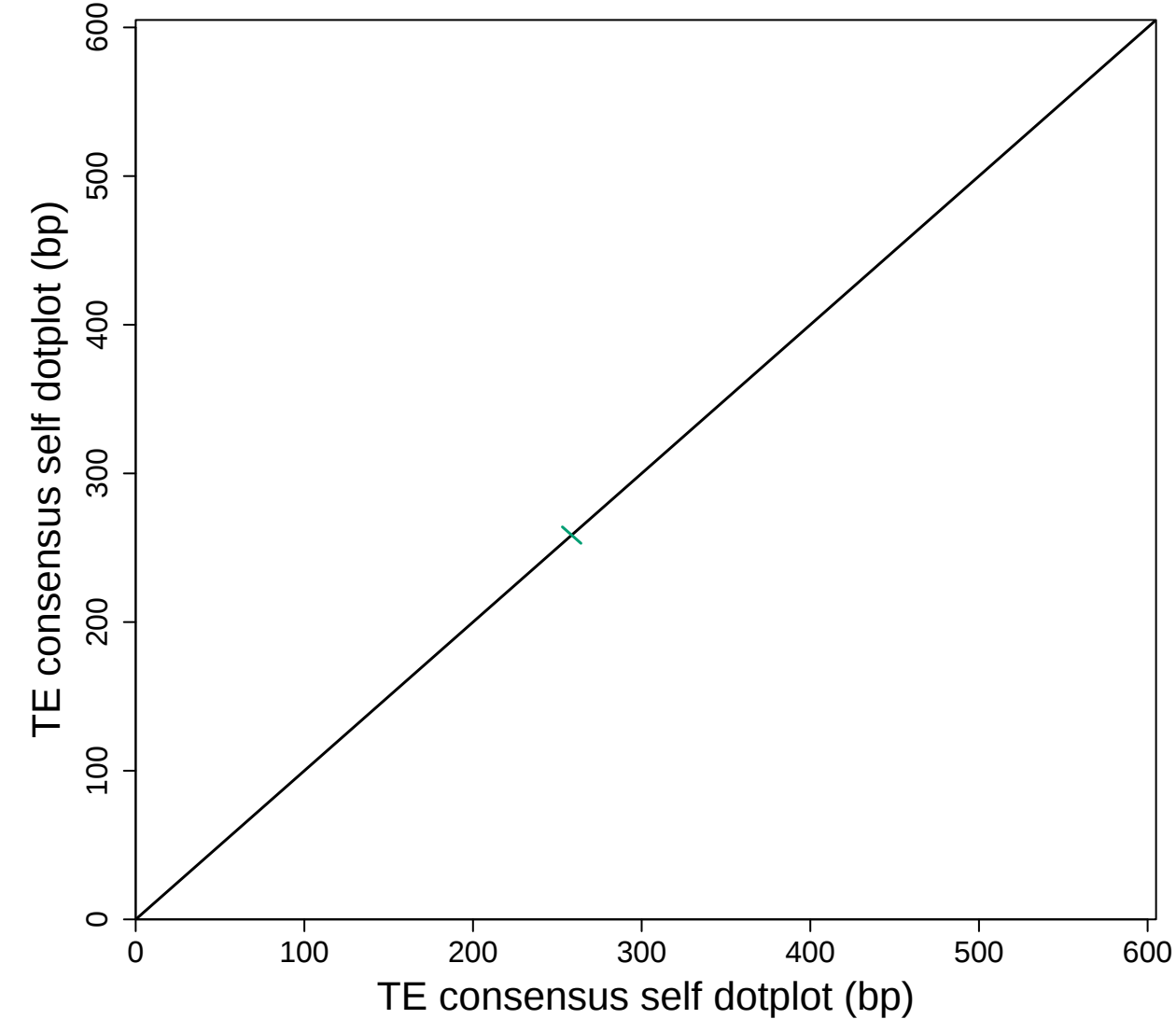
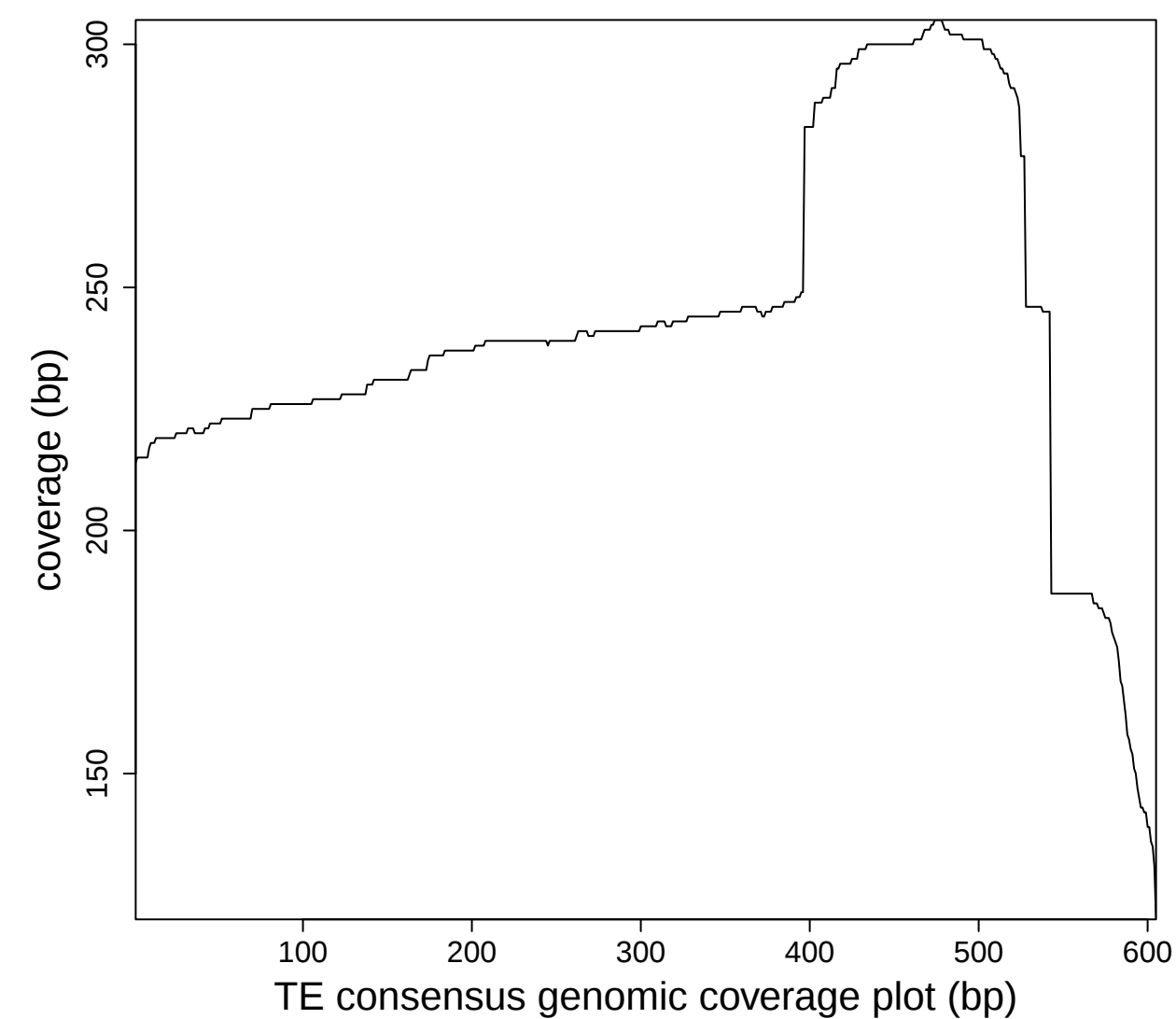
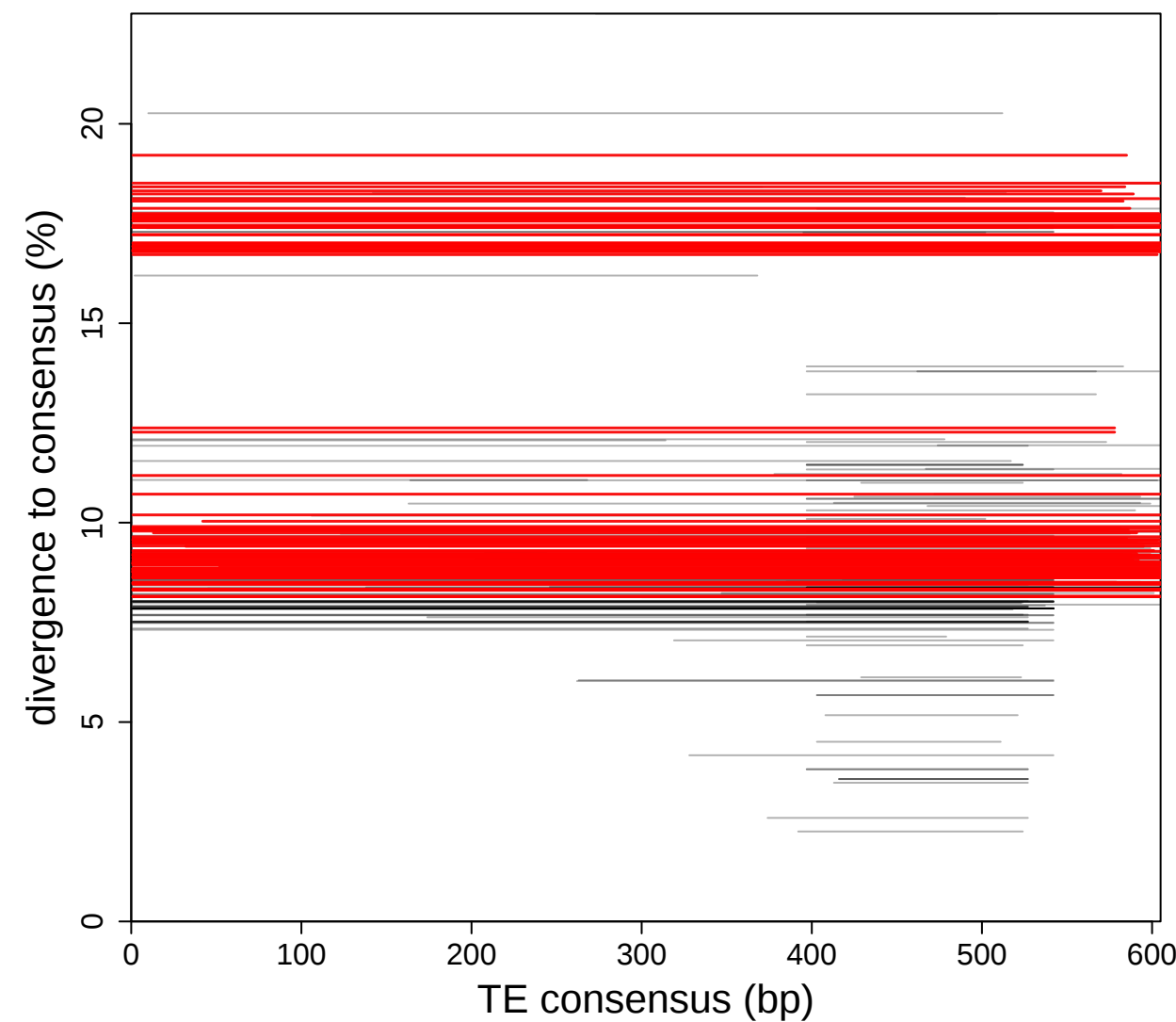


ly_1855.fasta.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 2971bp; fragments: 553; full length: 0 (>=2673.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

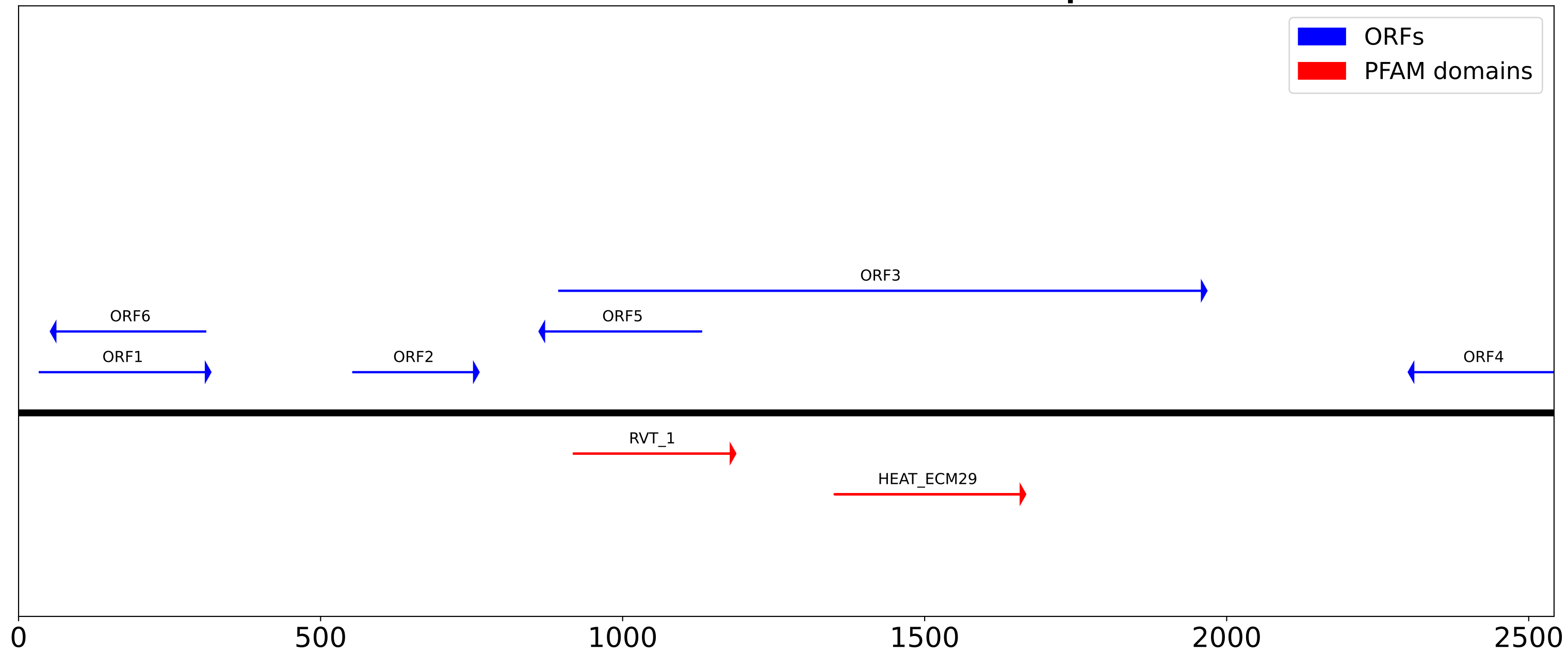


TE: rnd_4_family_1855
size: 605bp; fragments: 311; full length: 143 (≥ 544.5 bp)



Before TEtrimmer 605 bp

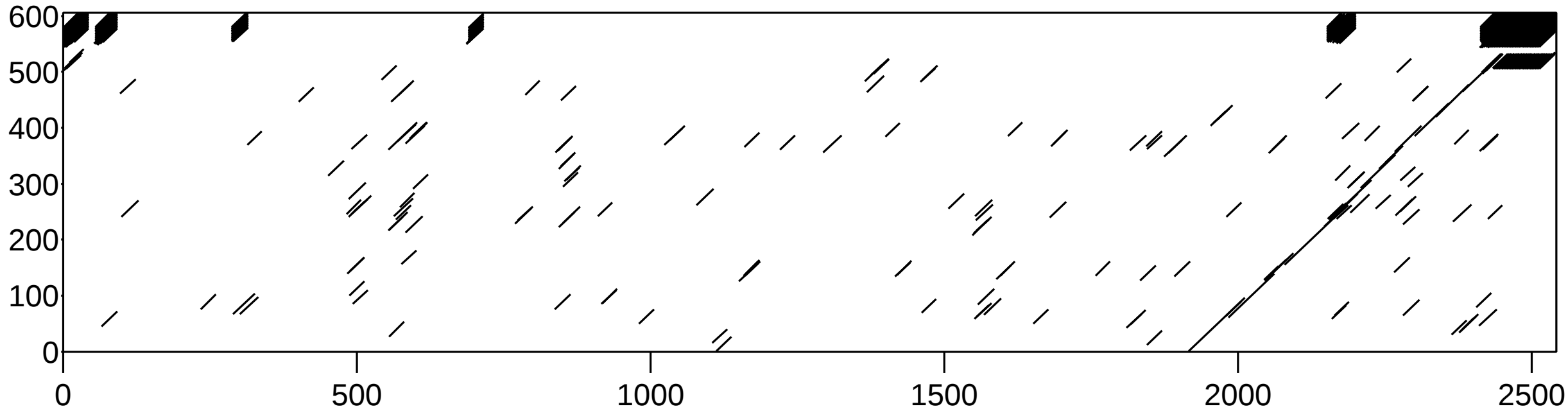
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)