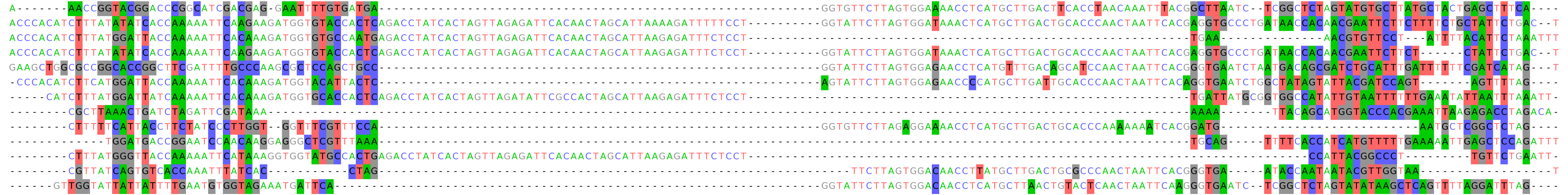


Start crop Point

End crop Point

MSA length = 1418



Start crop Point



End crop Point



0

10

0

250

500

750

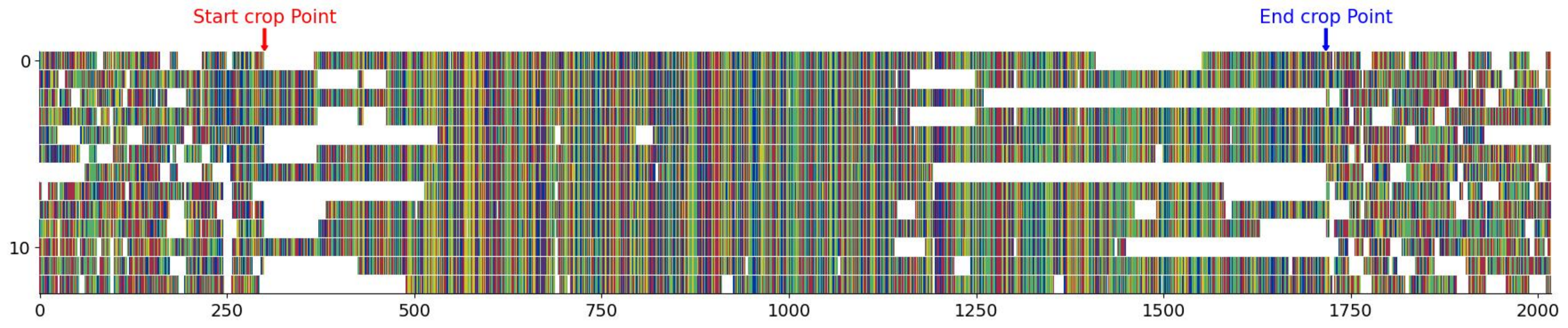
1000

1250

1500

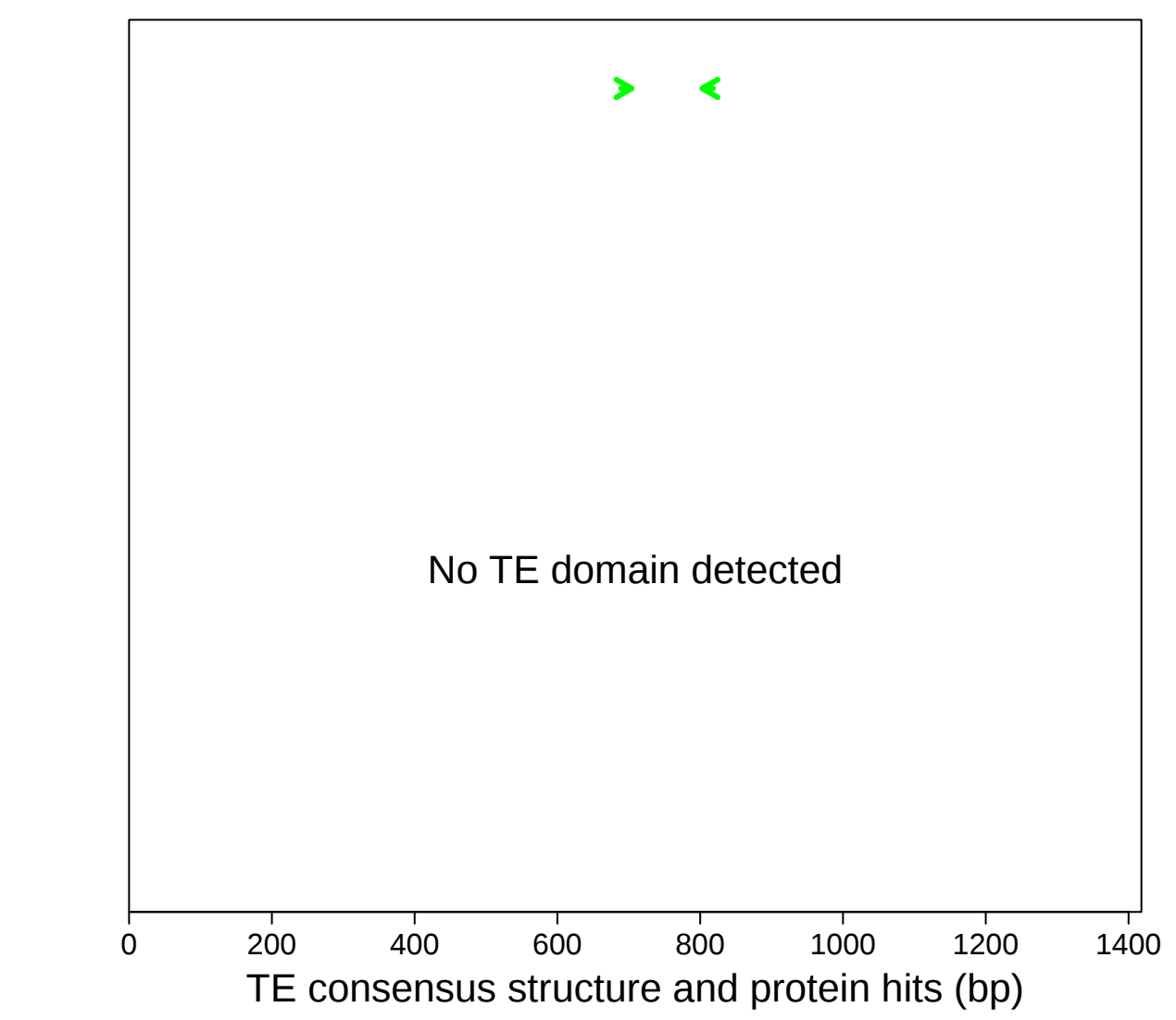
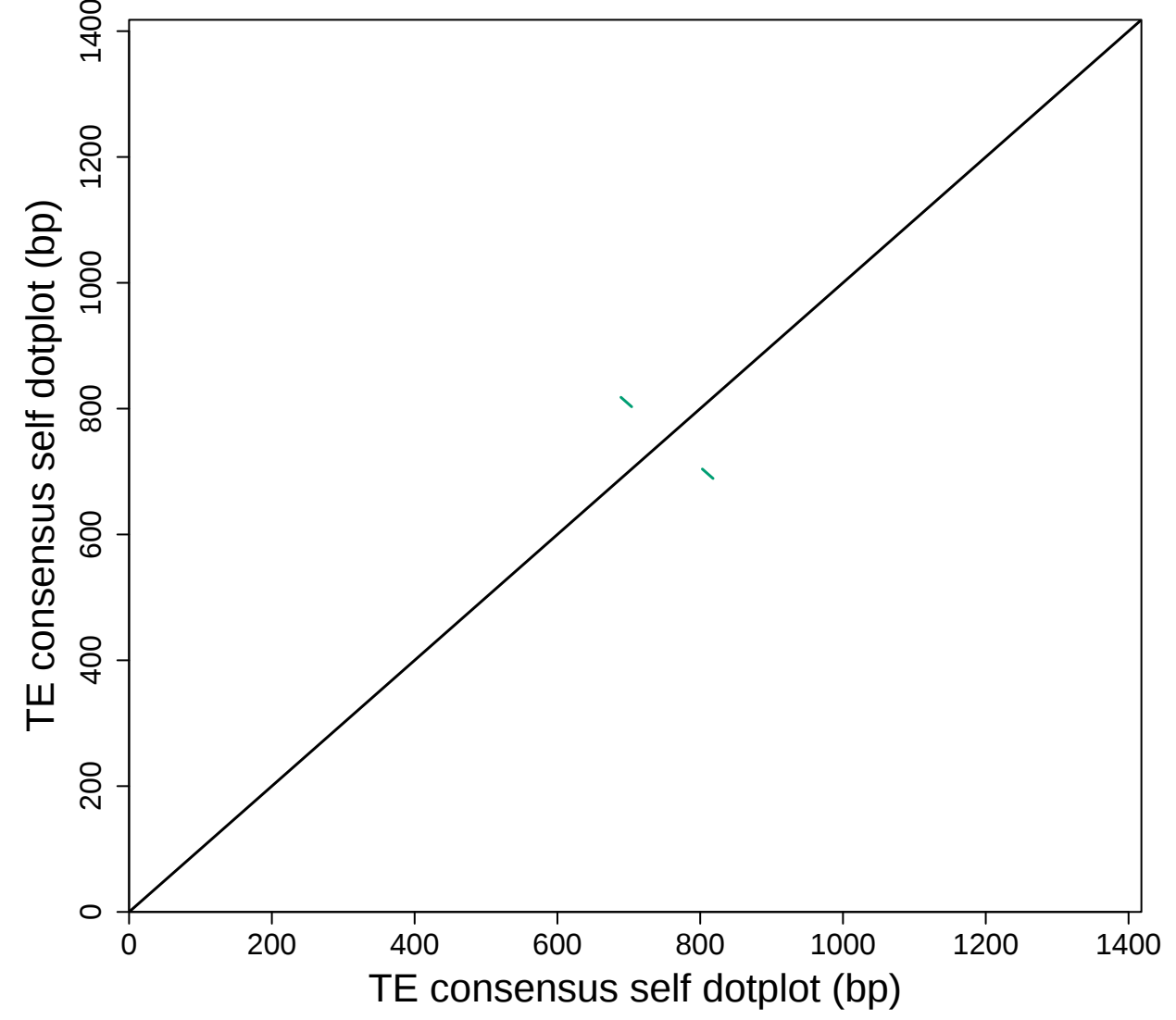
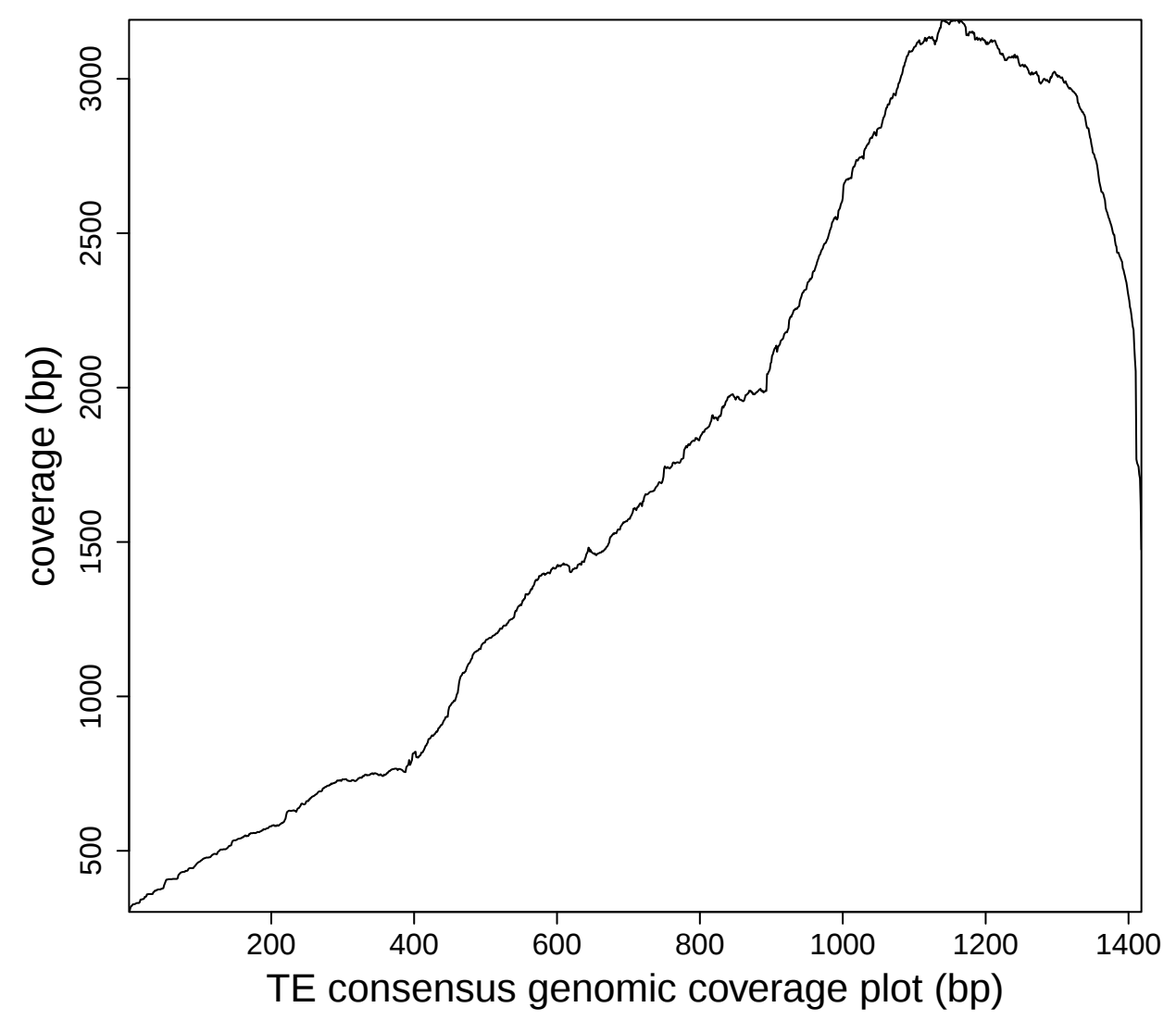
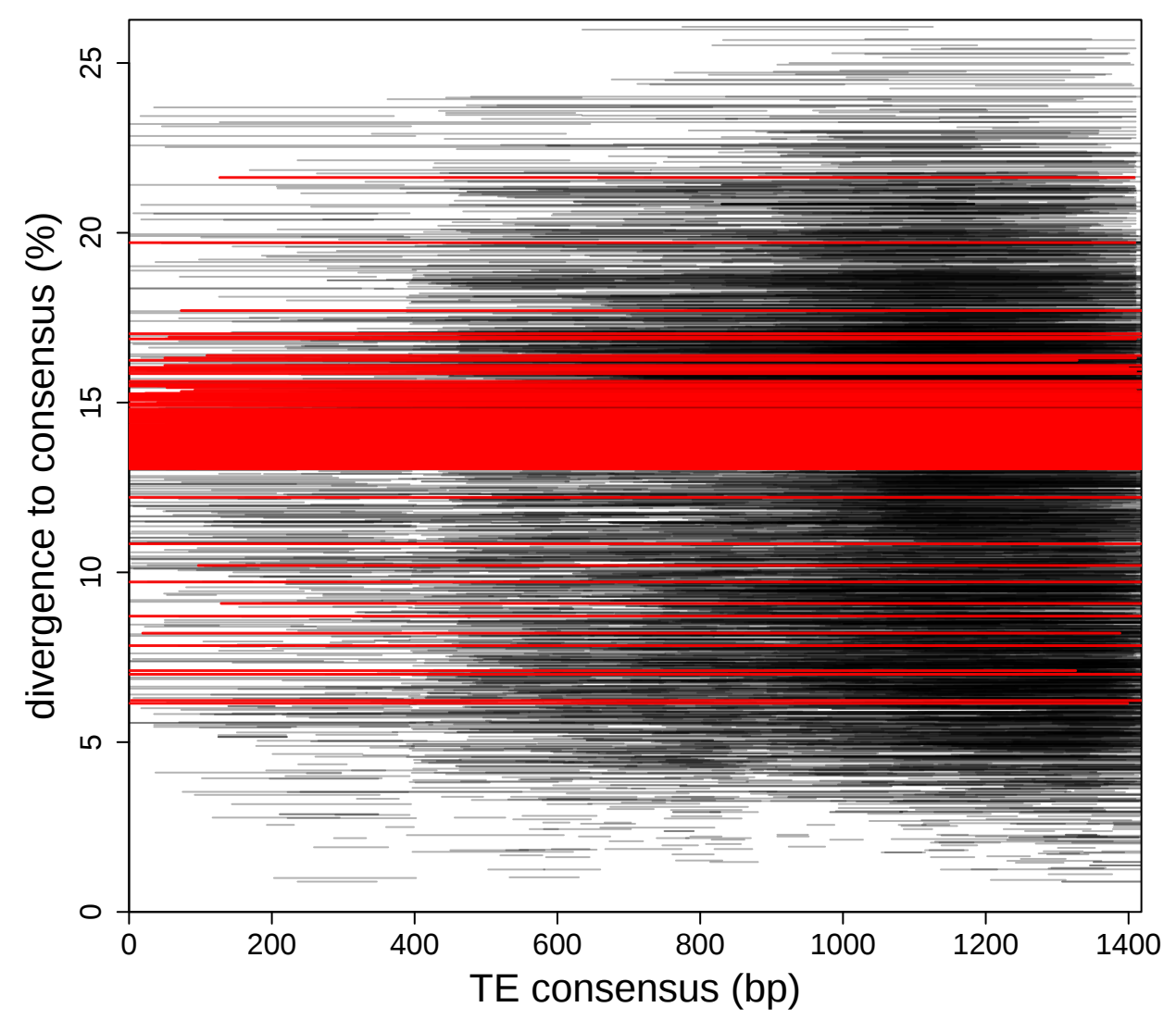
1750

2000



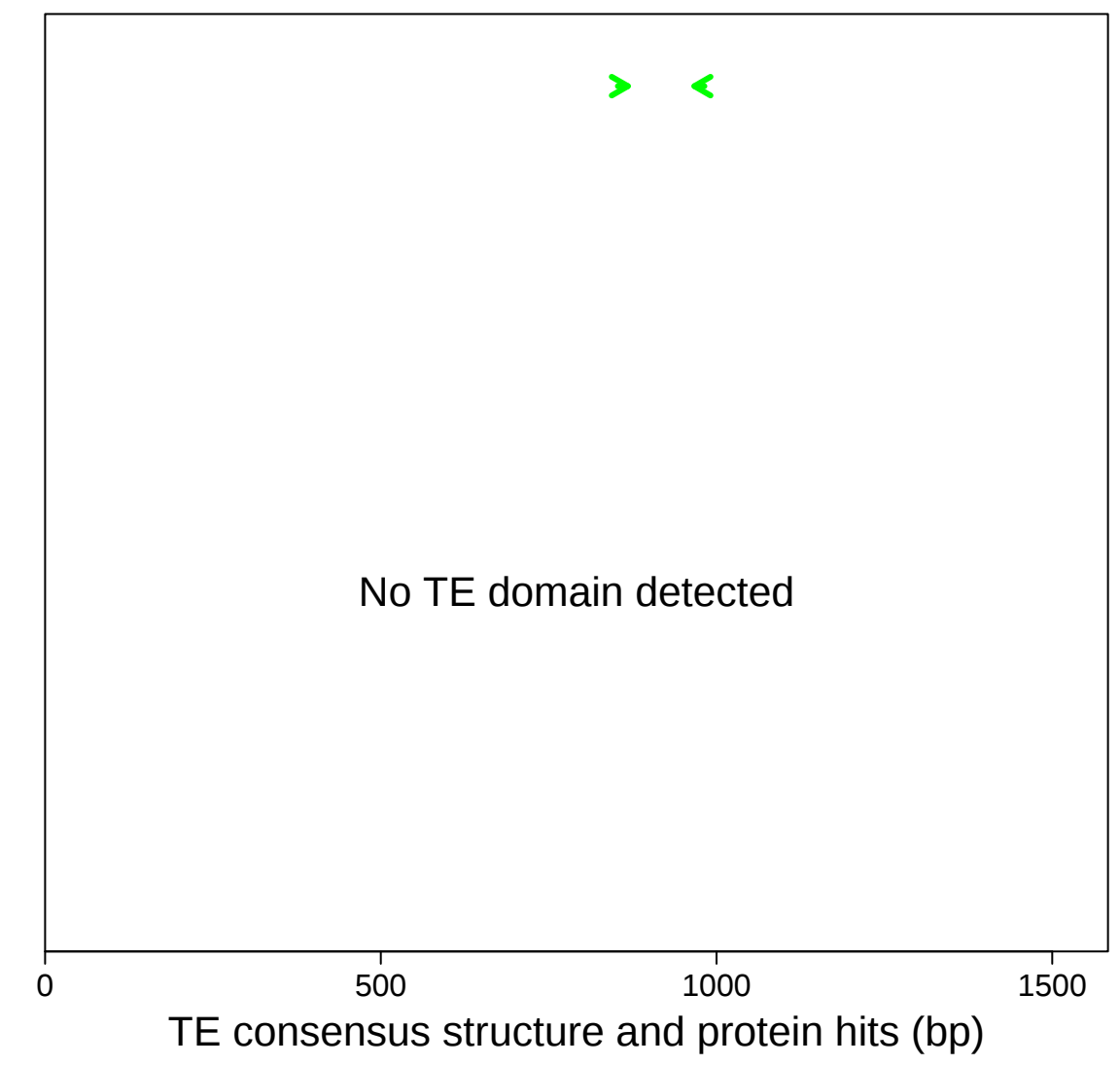
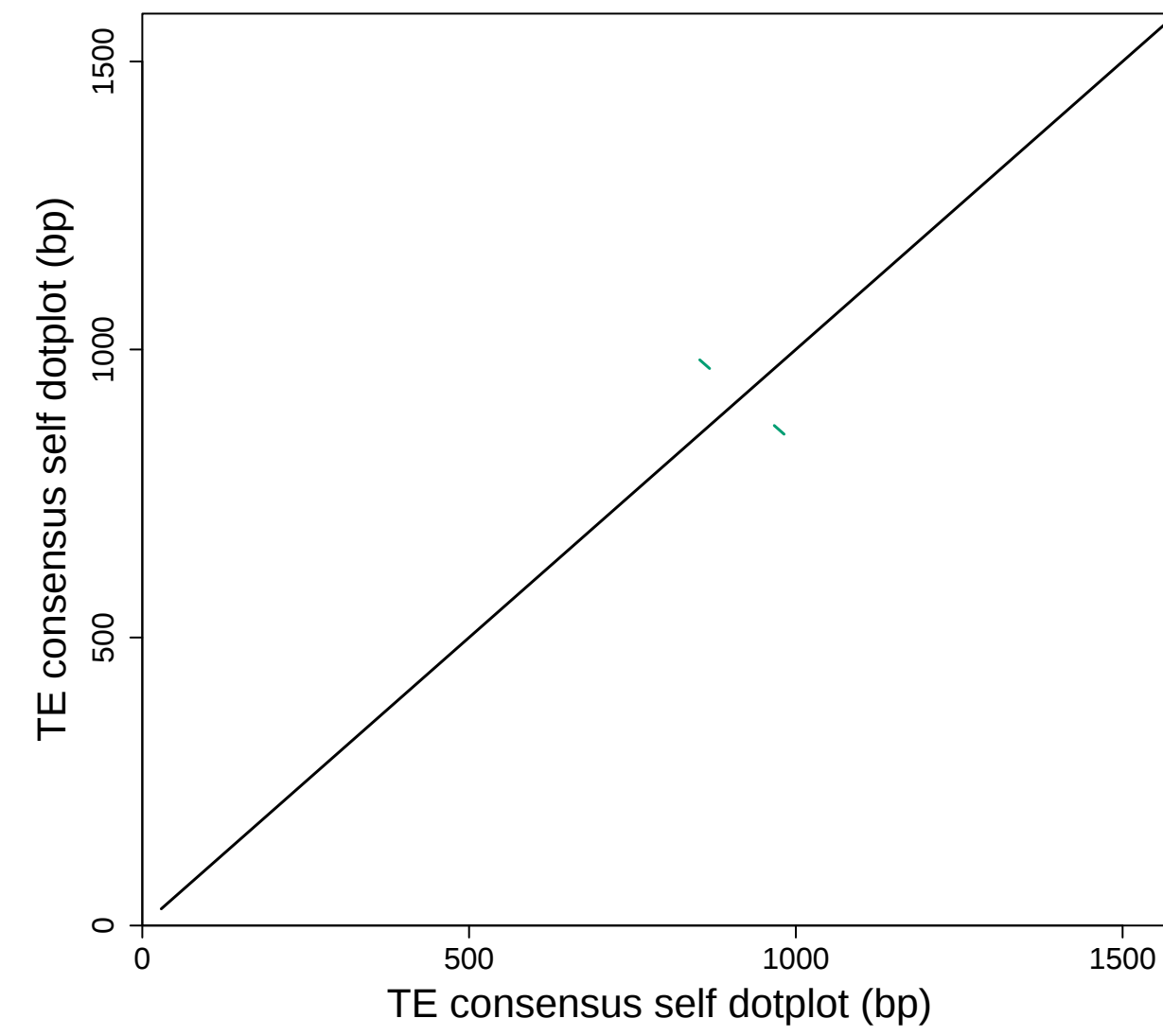
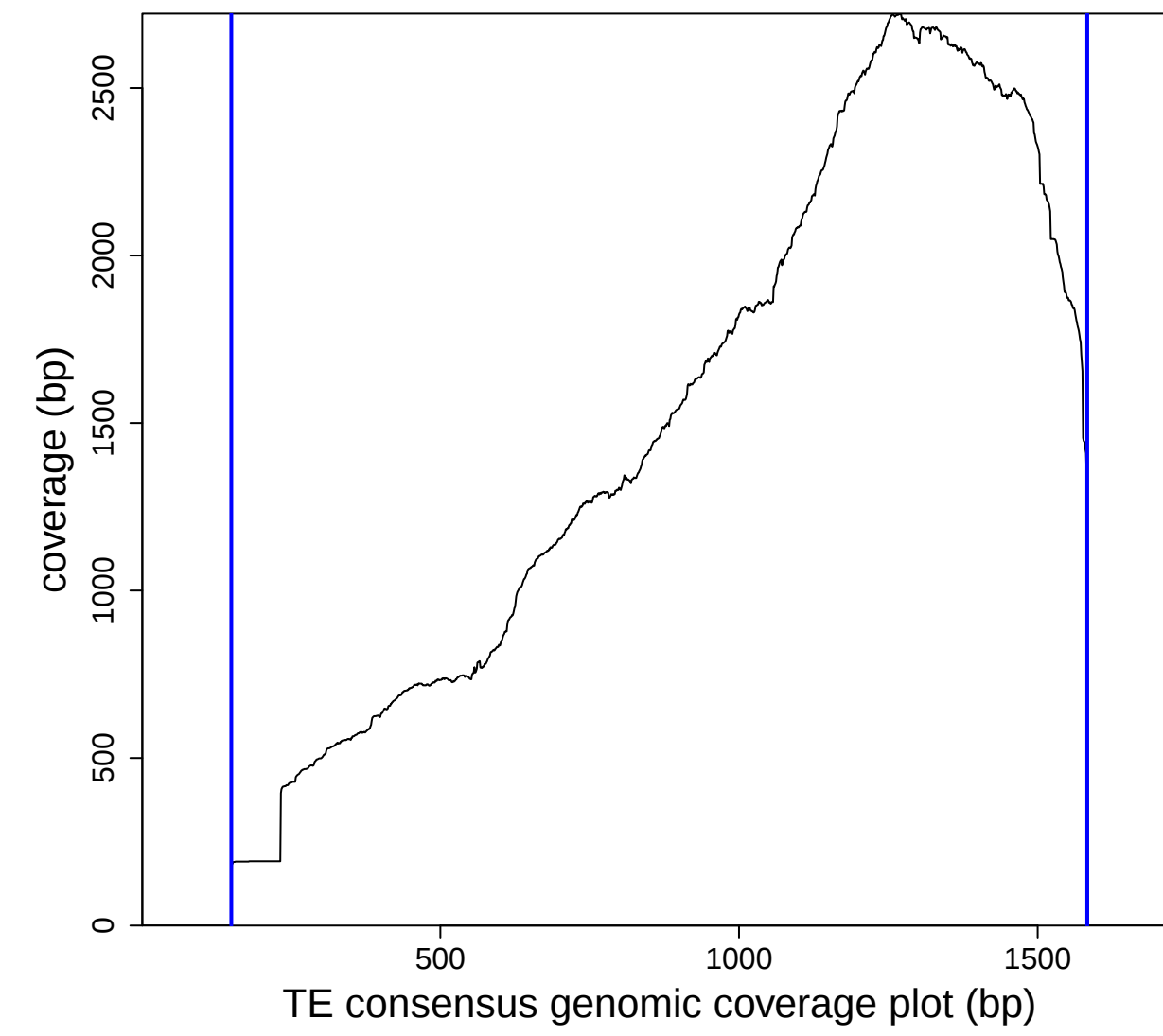
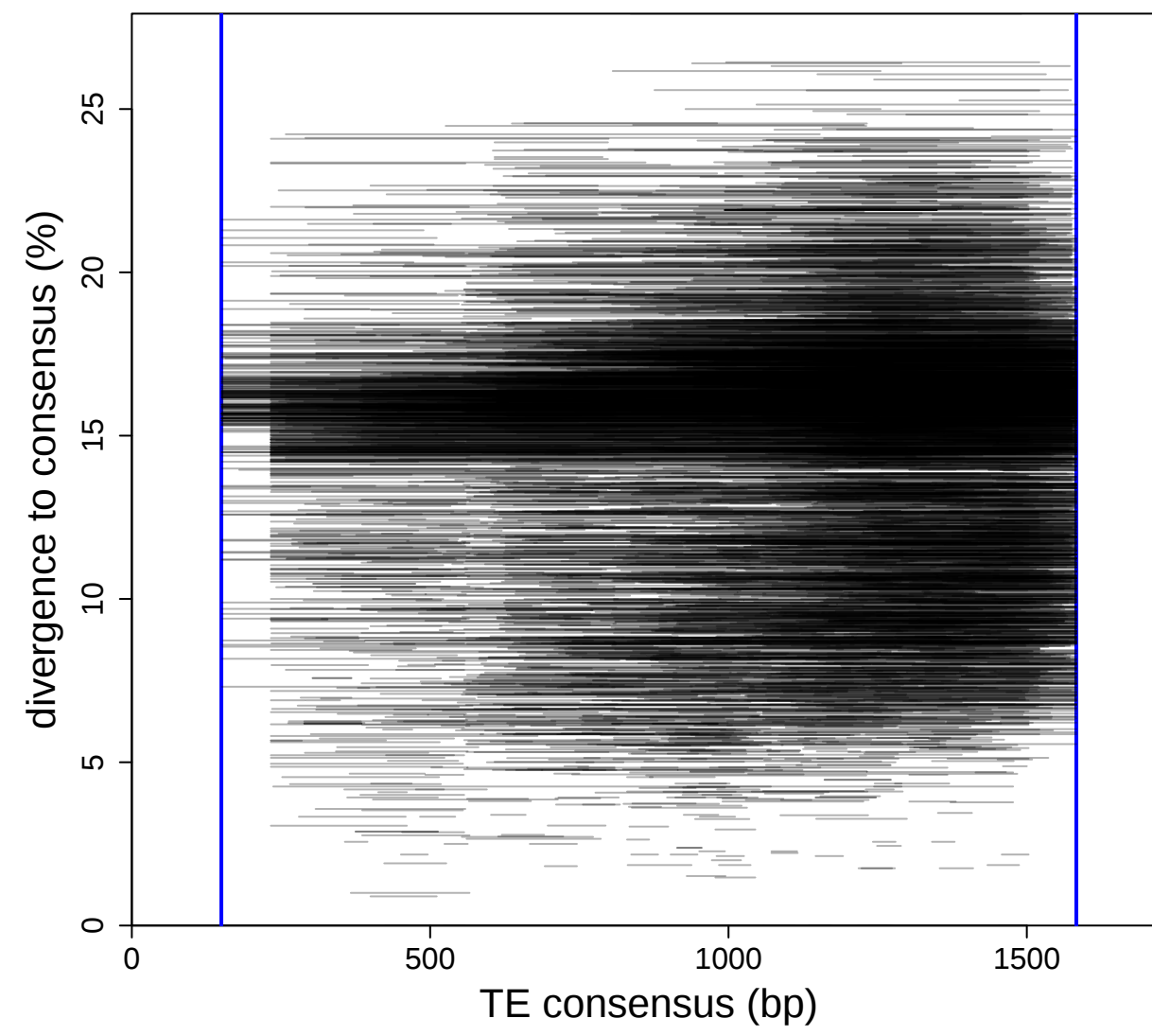
ed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 1418bp; fragments: 5597; full length: 269 (≥ 1276.2 bp)

After TEtrimmer 1418 bp

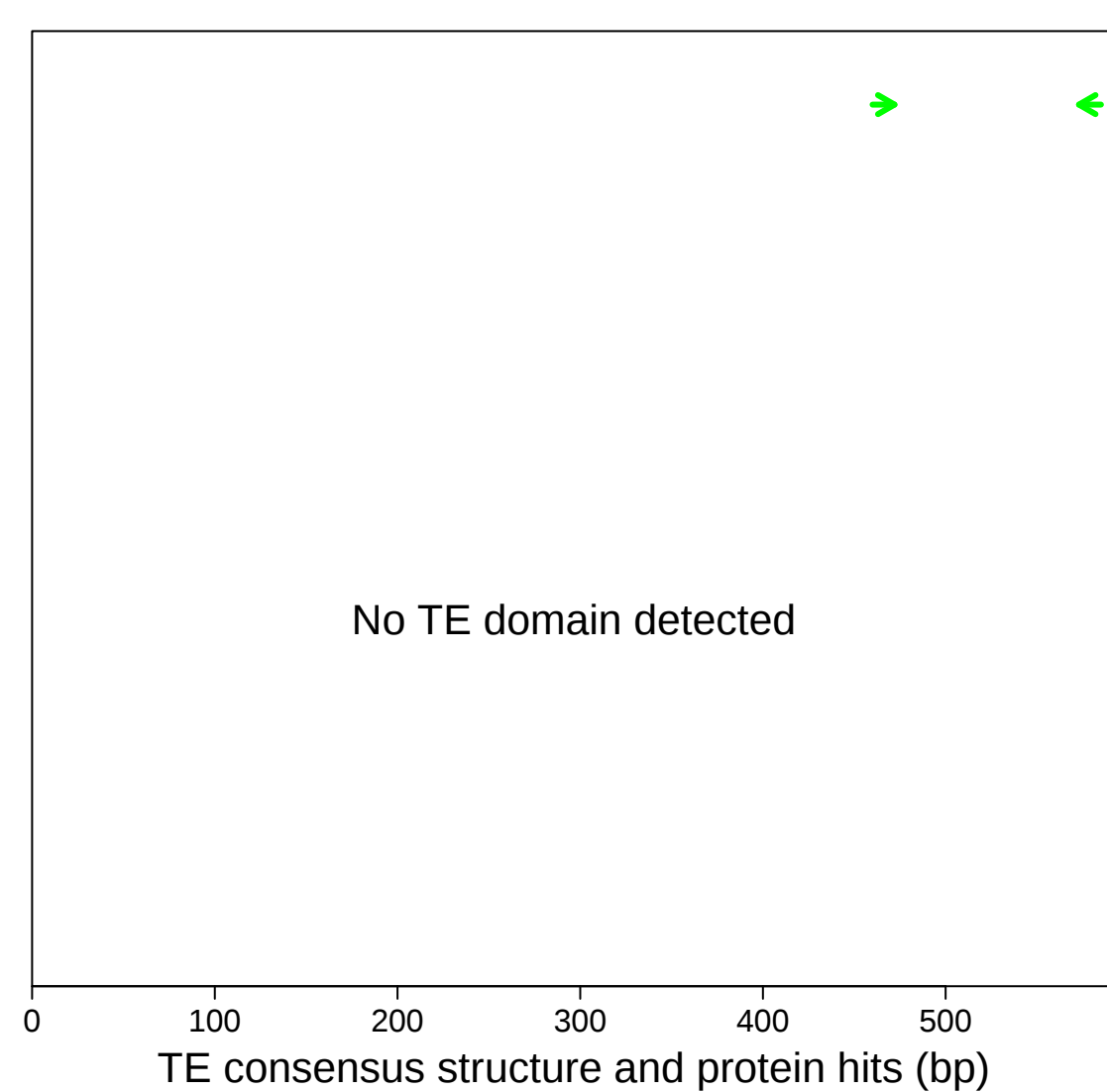
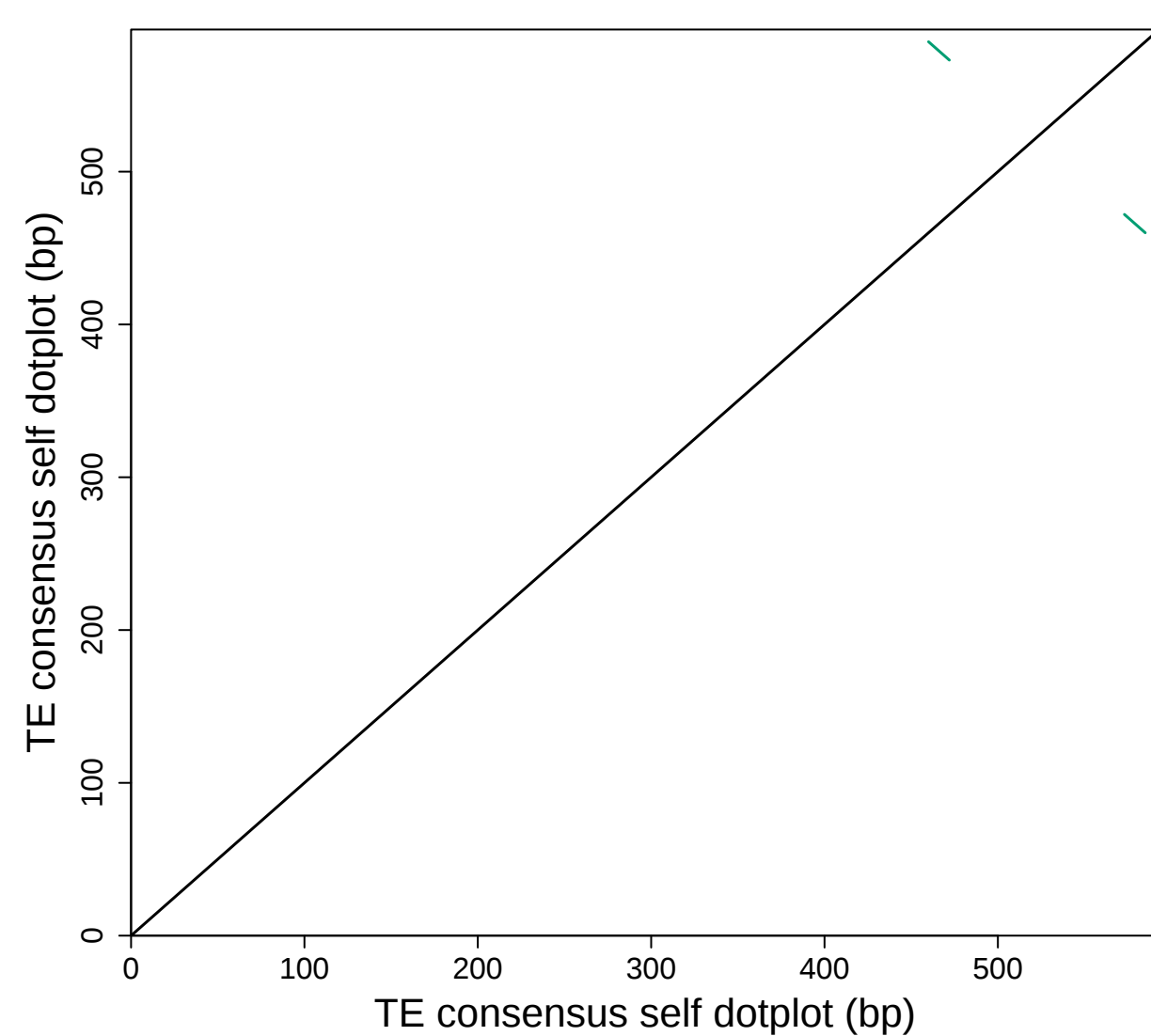
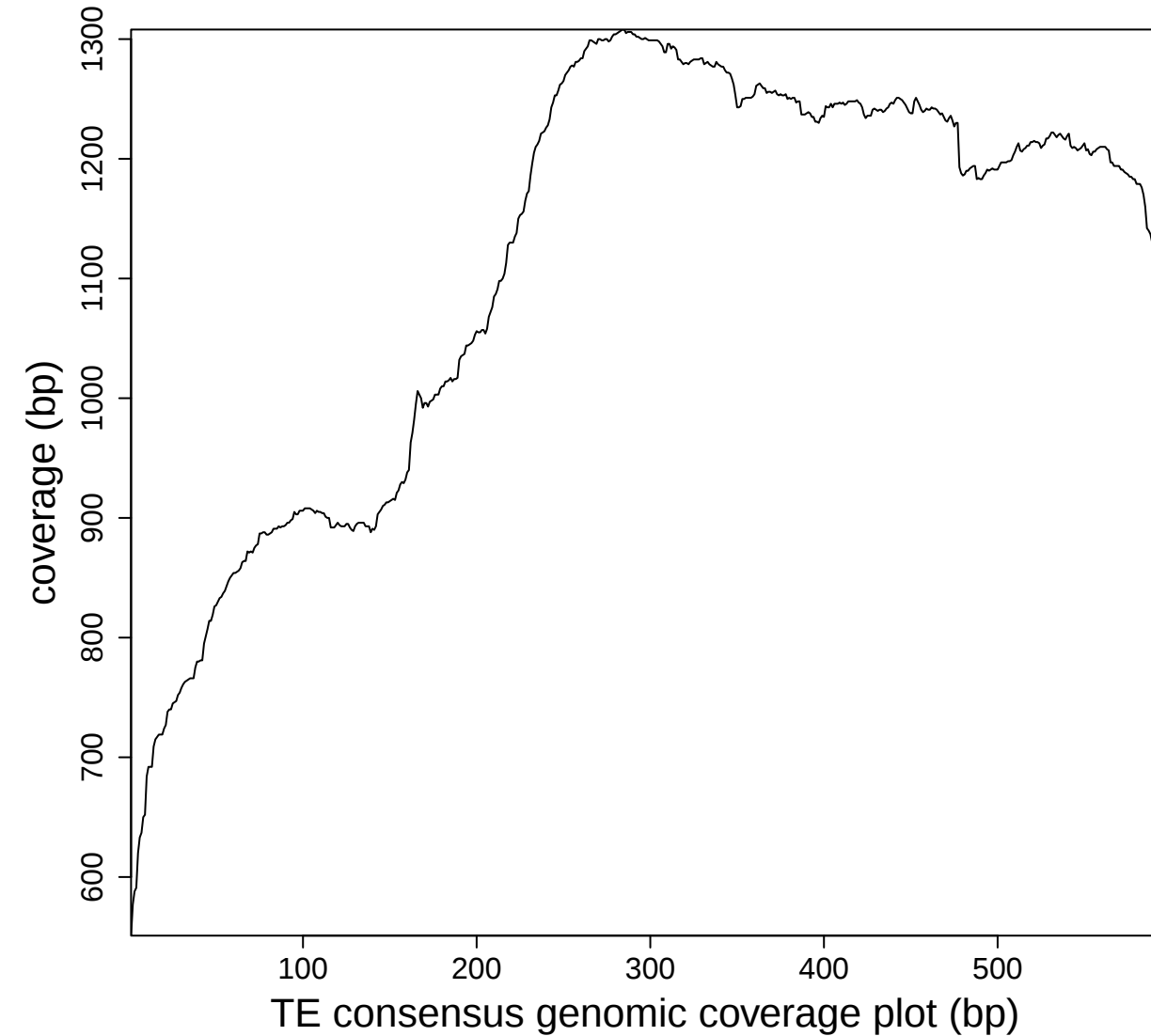
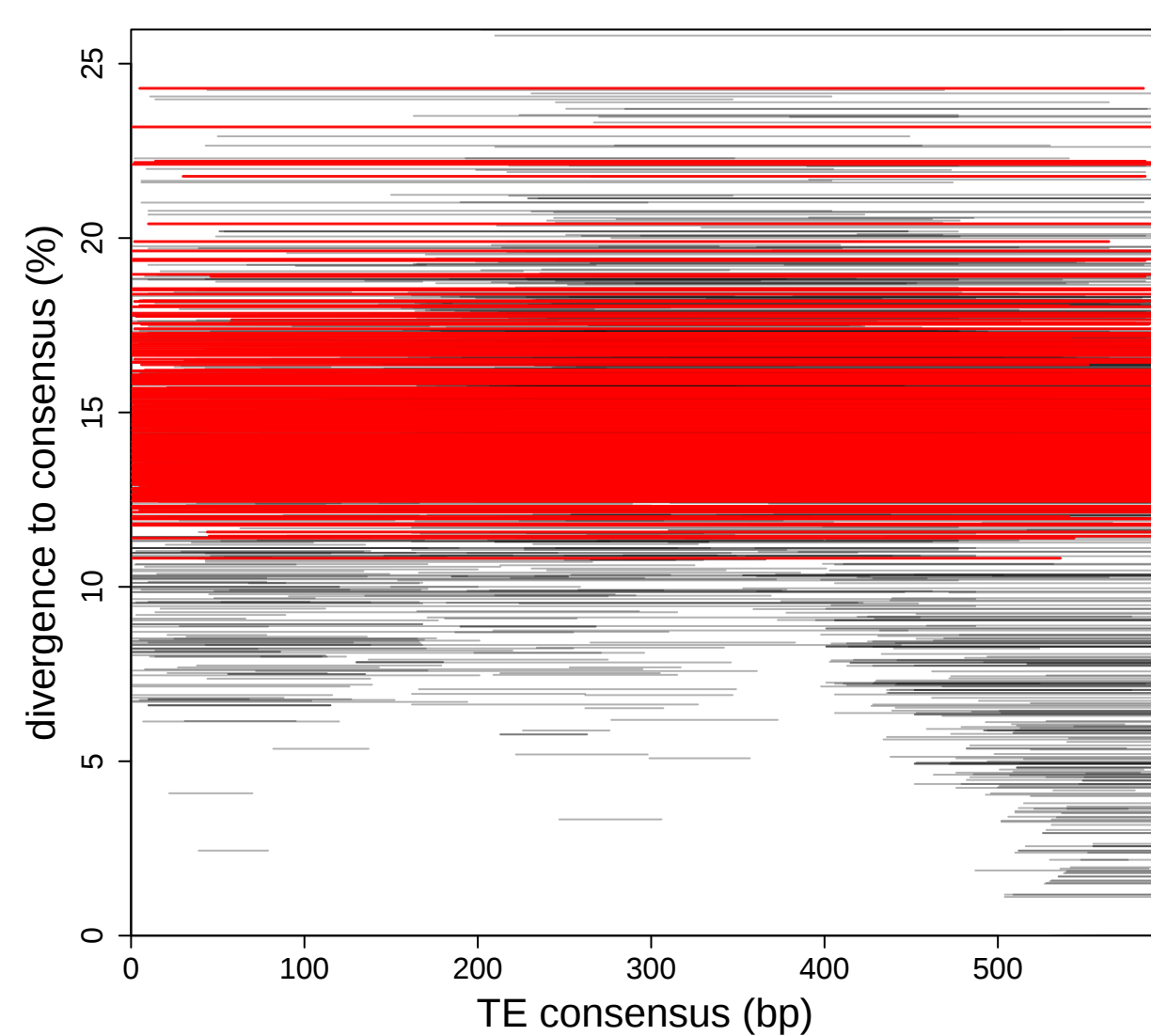


8.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1733bp; fragments: 4635; full length: 0 (>=1559.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

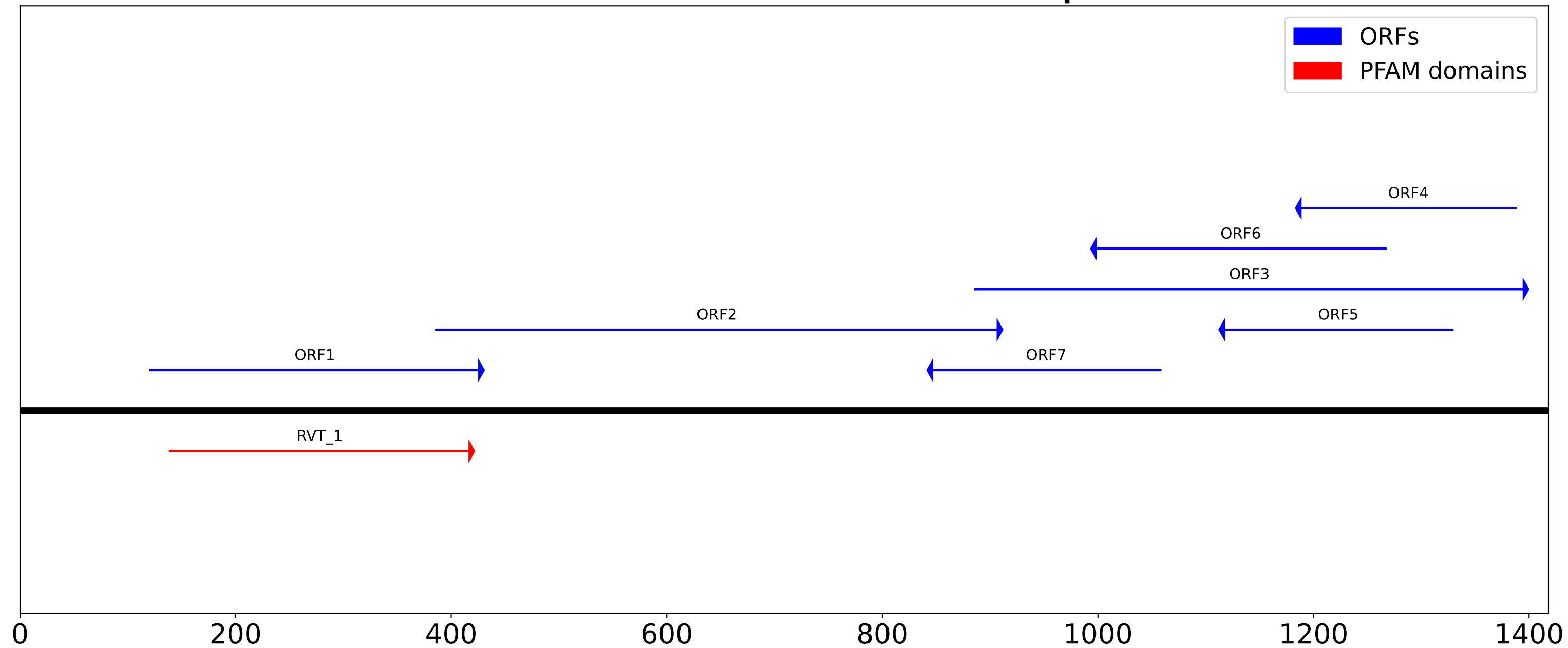


TE: rnd_1_family_208
size: 593bp; fragments: 1811; full length: 543 (>=533.7bp)

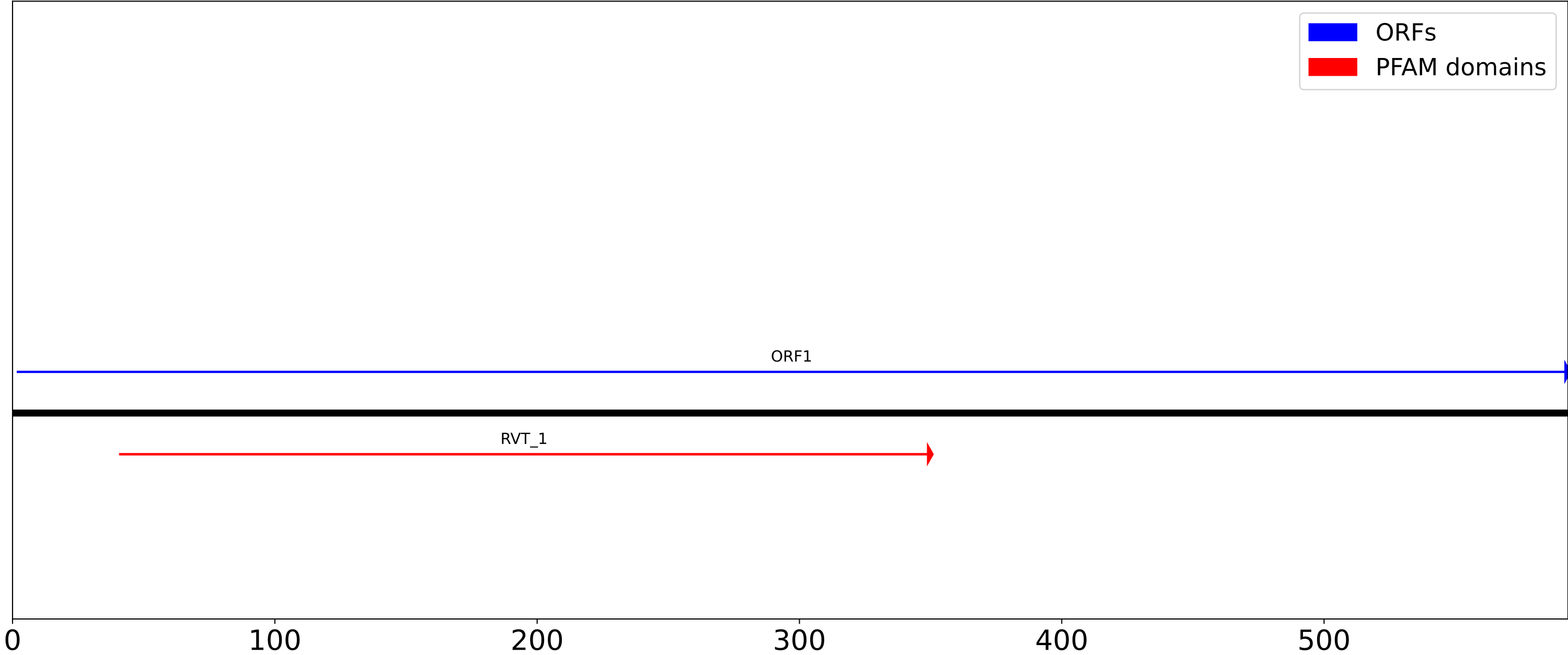


Before TEtrimmer 593 bp

After TEtrimmer ORF and PFAM domain plot



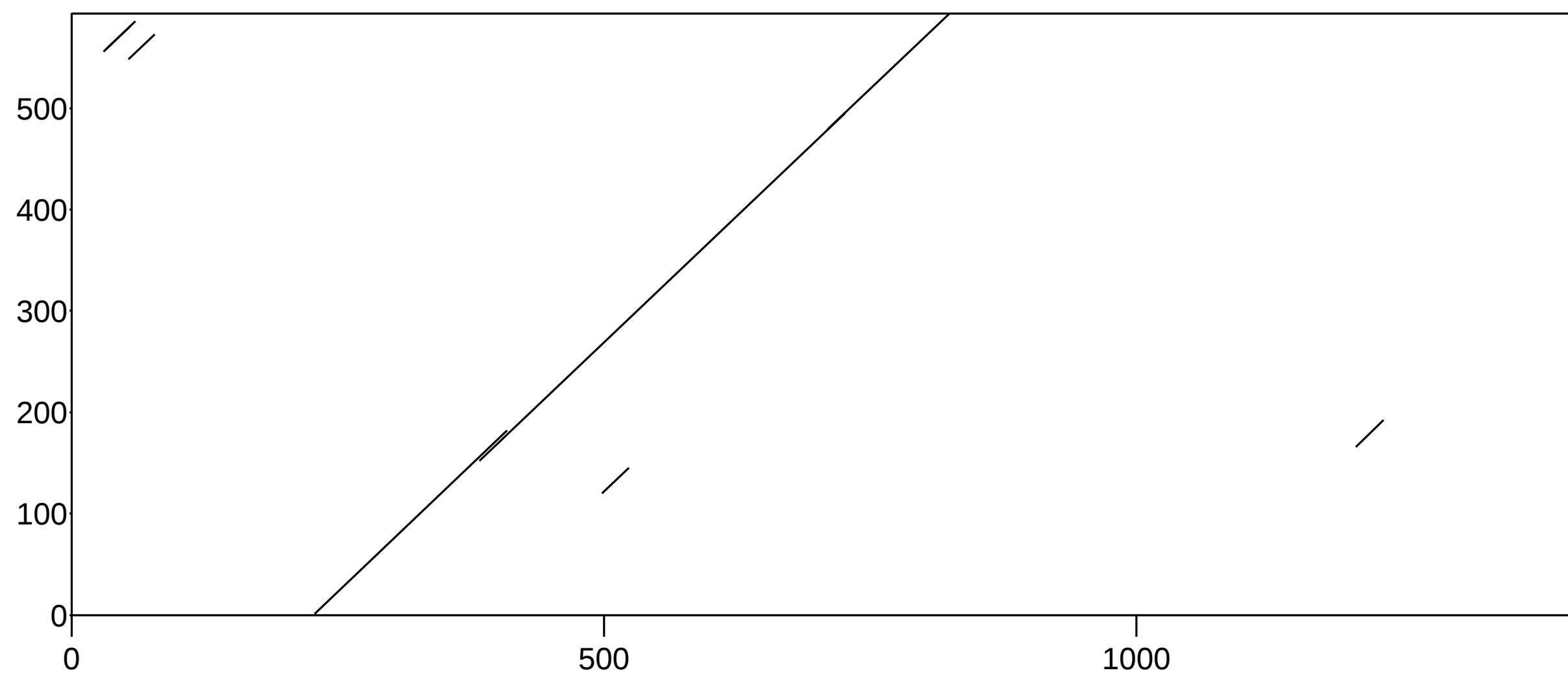
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)