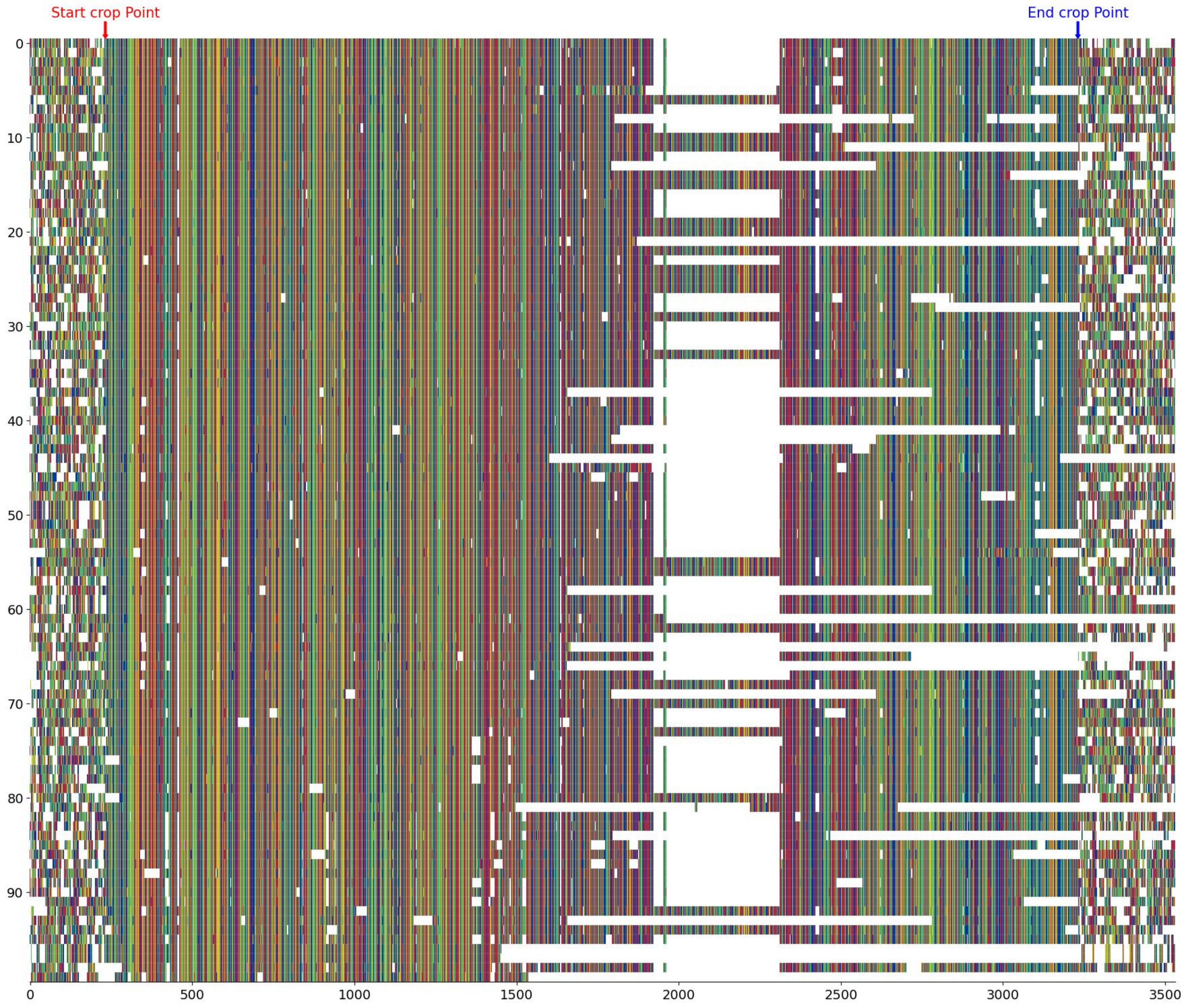
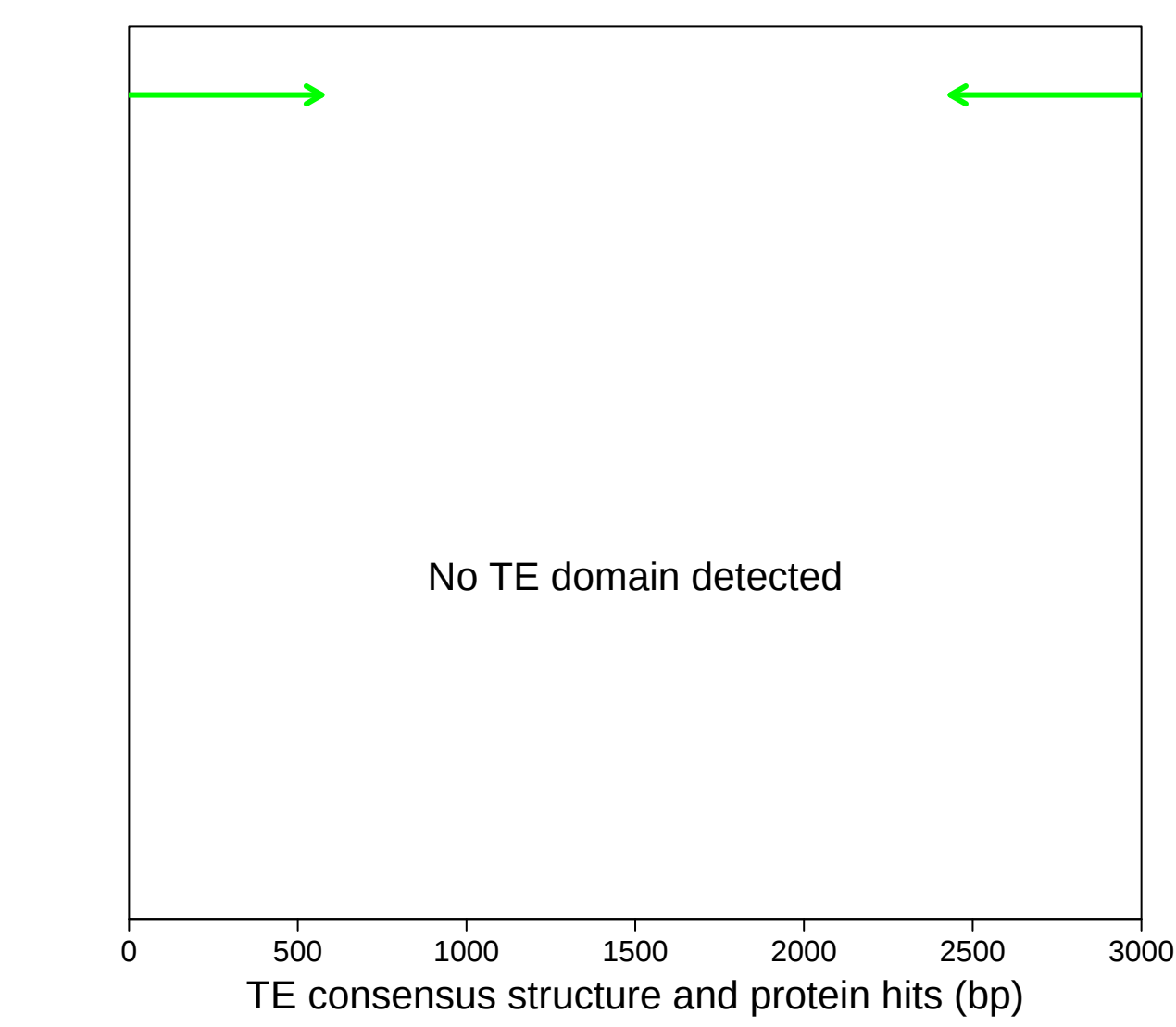
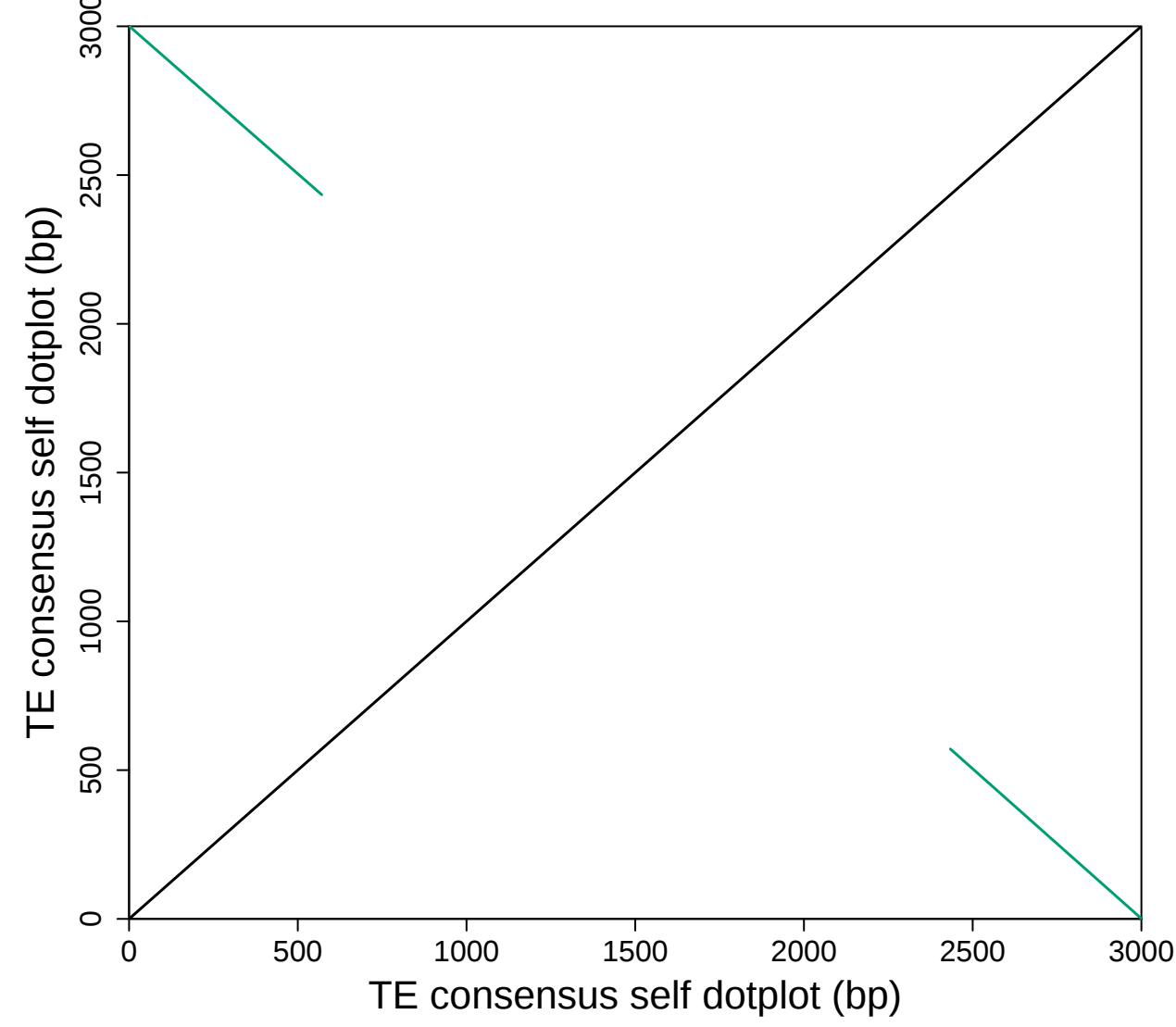
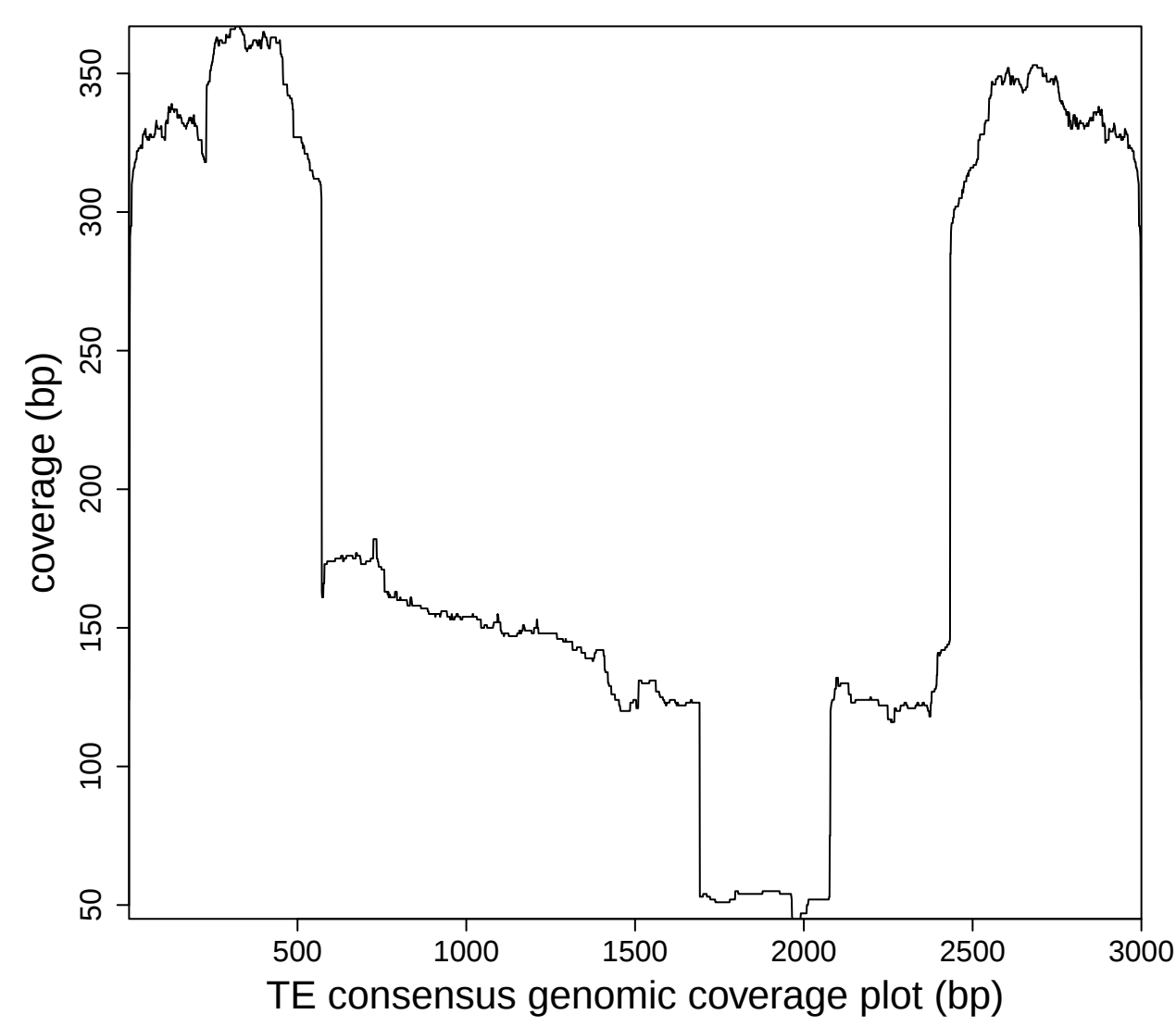
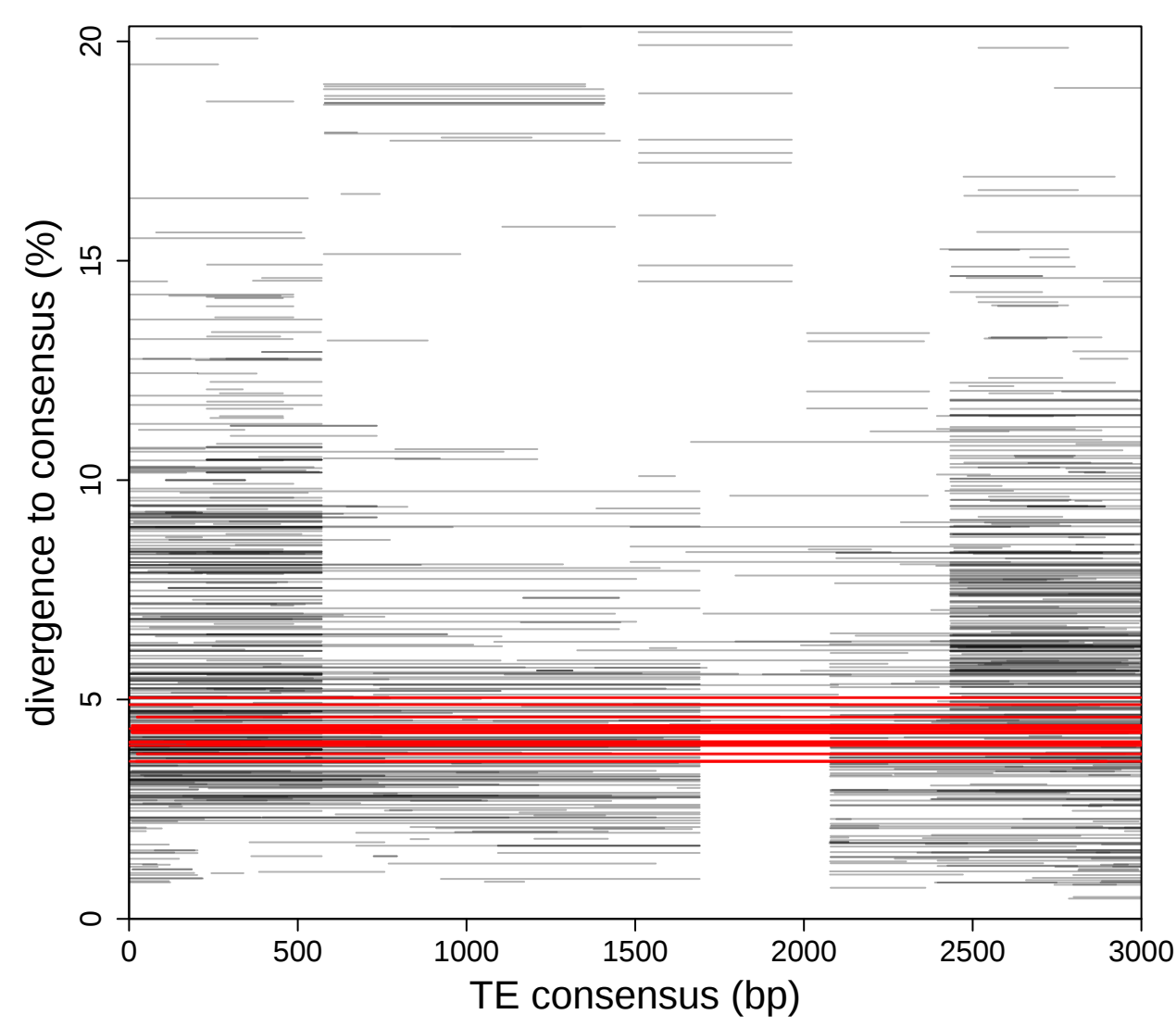






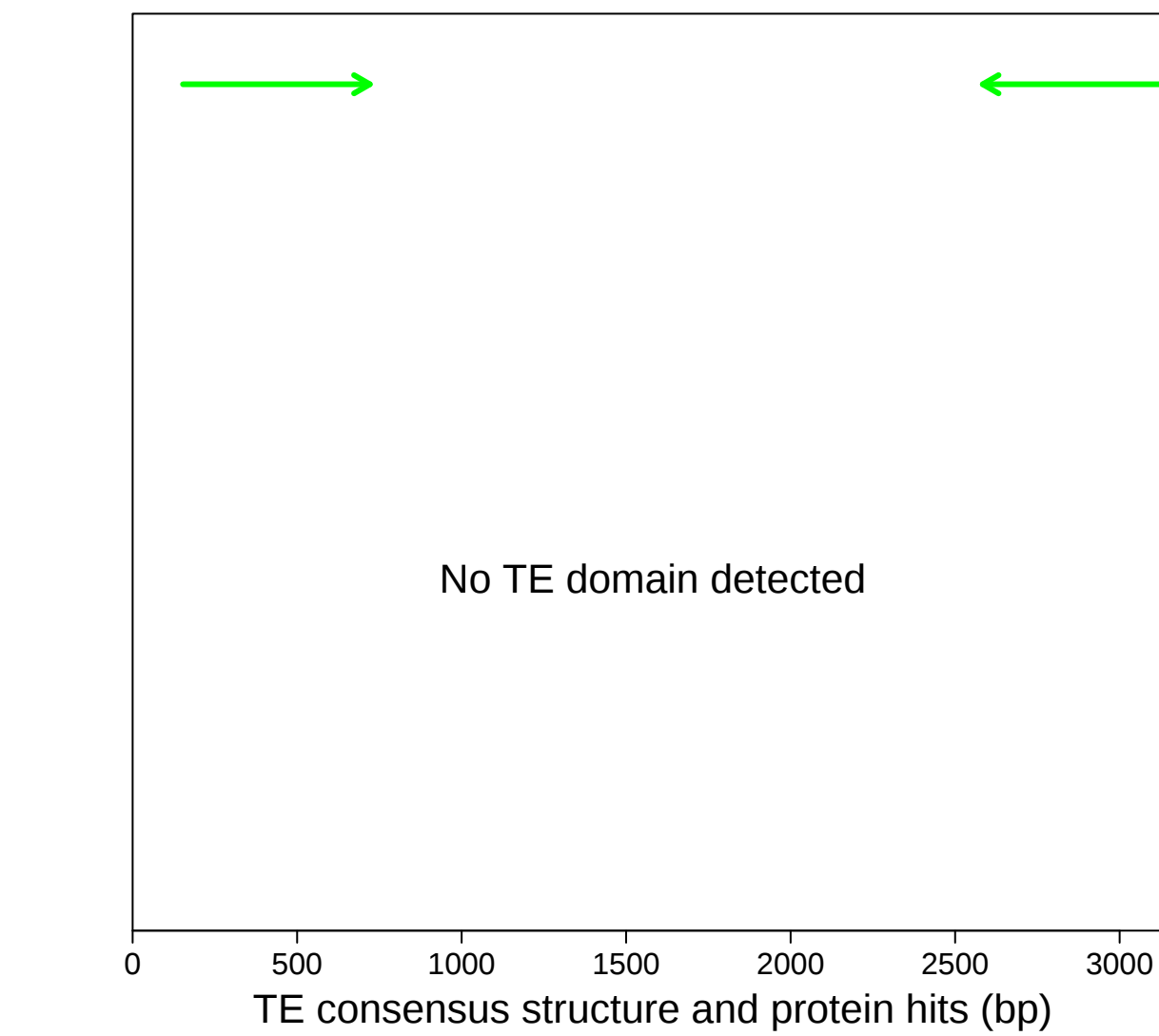
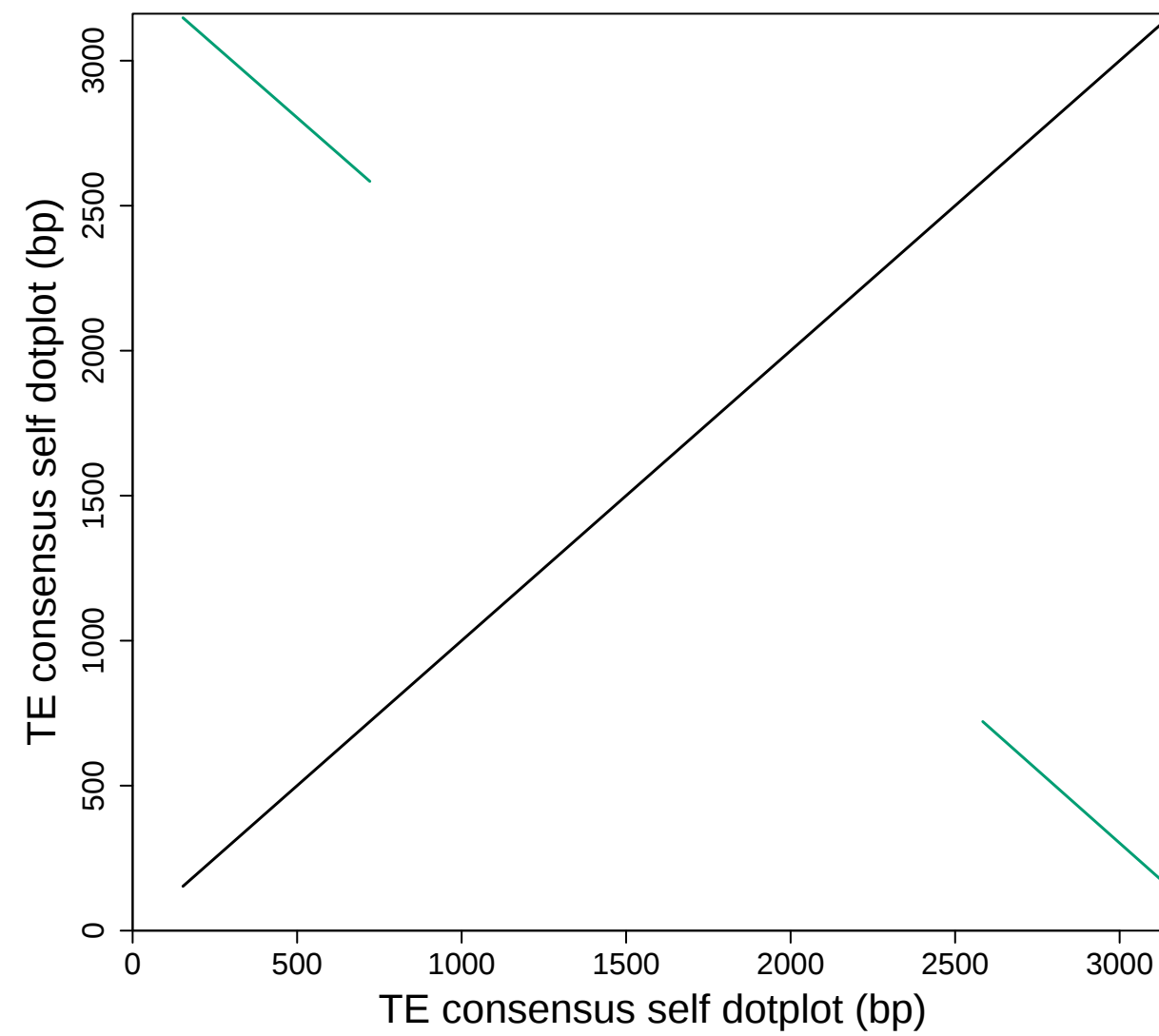
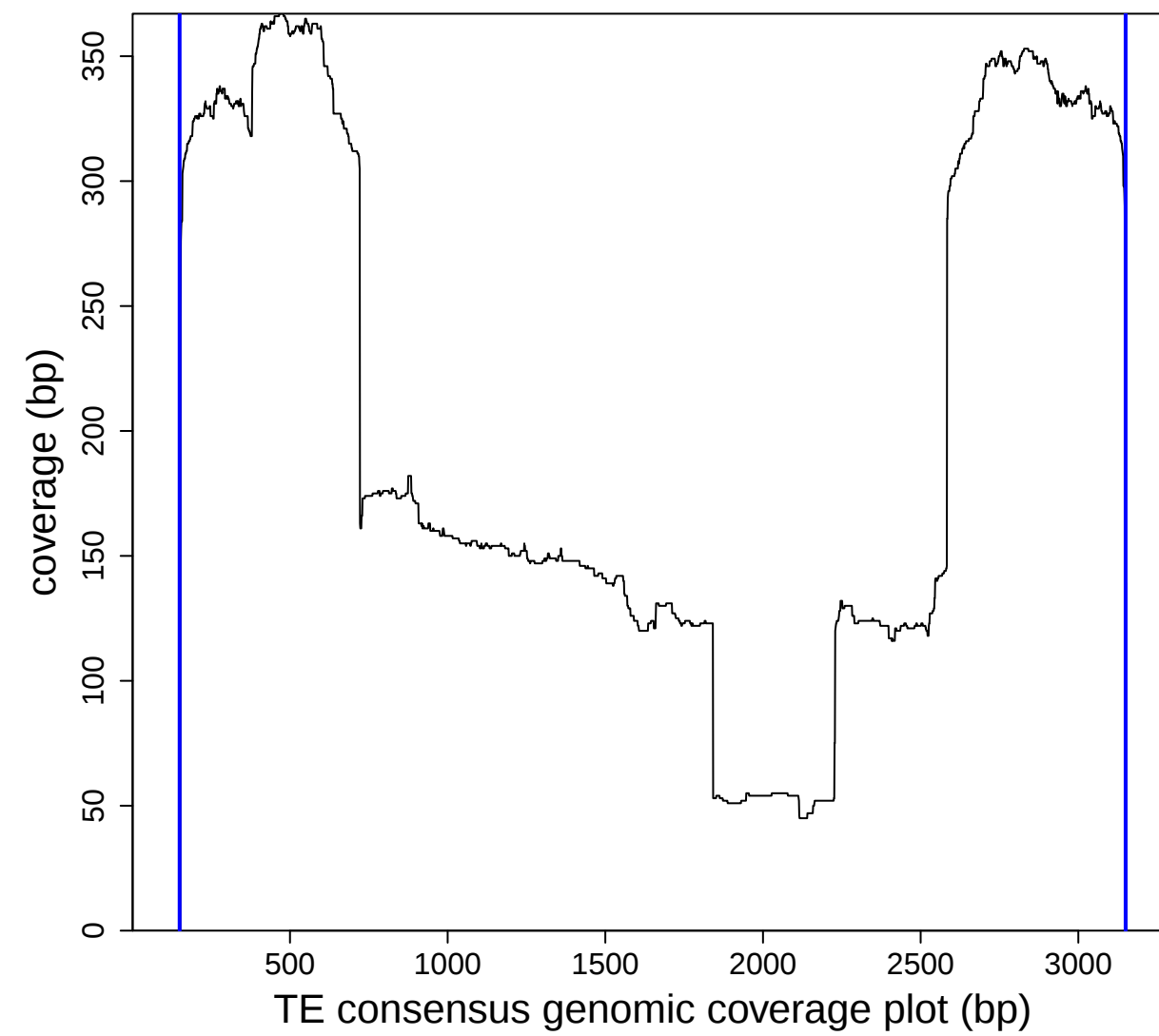
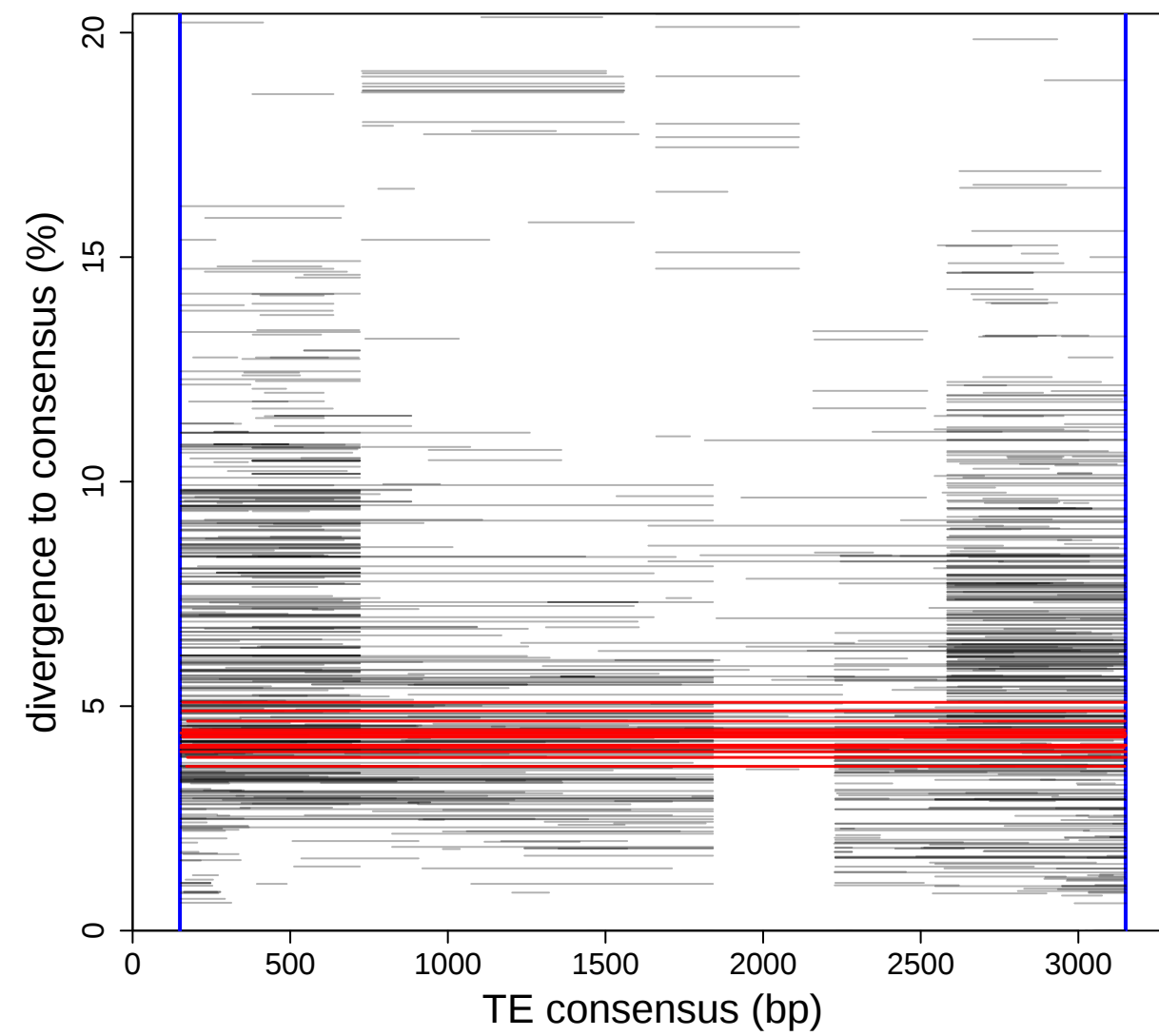
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 3000bp; fragments: 990; full length: 20 (≥ 2700 bp)

After TEtrimmer 3000 bp



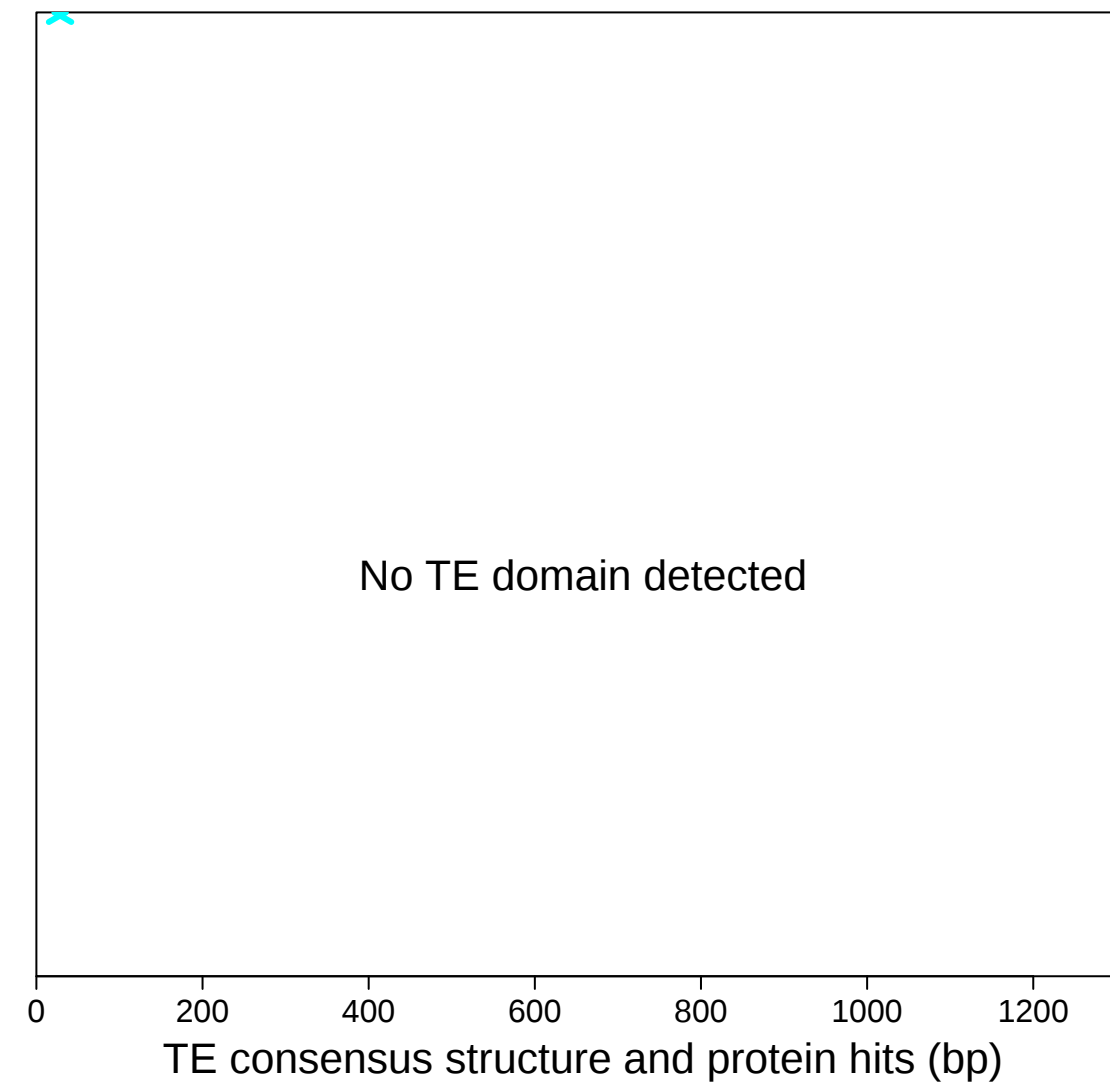
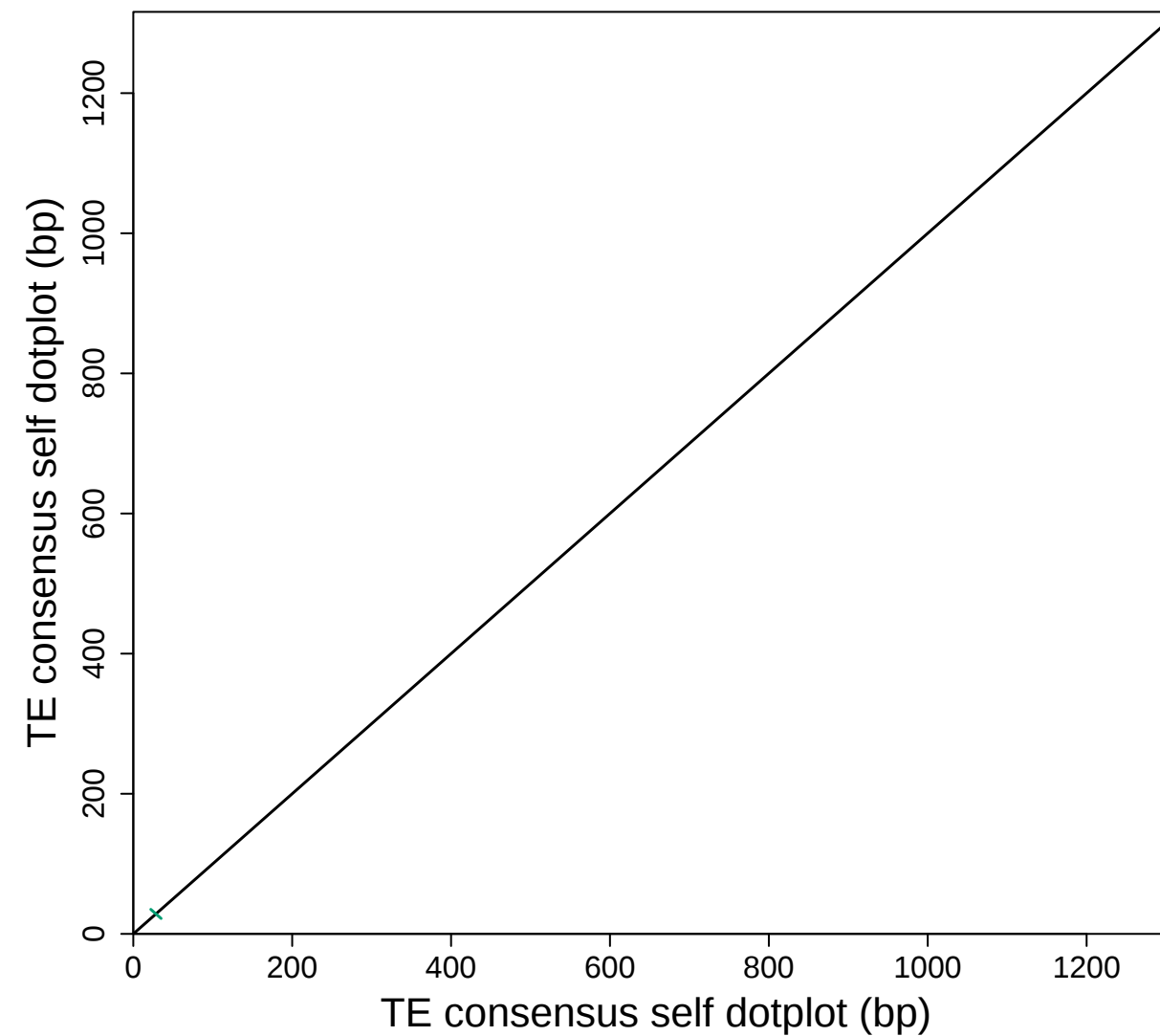
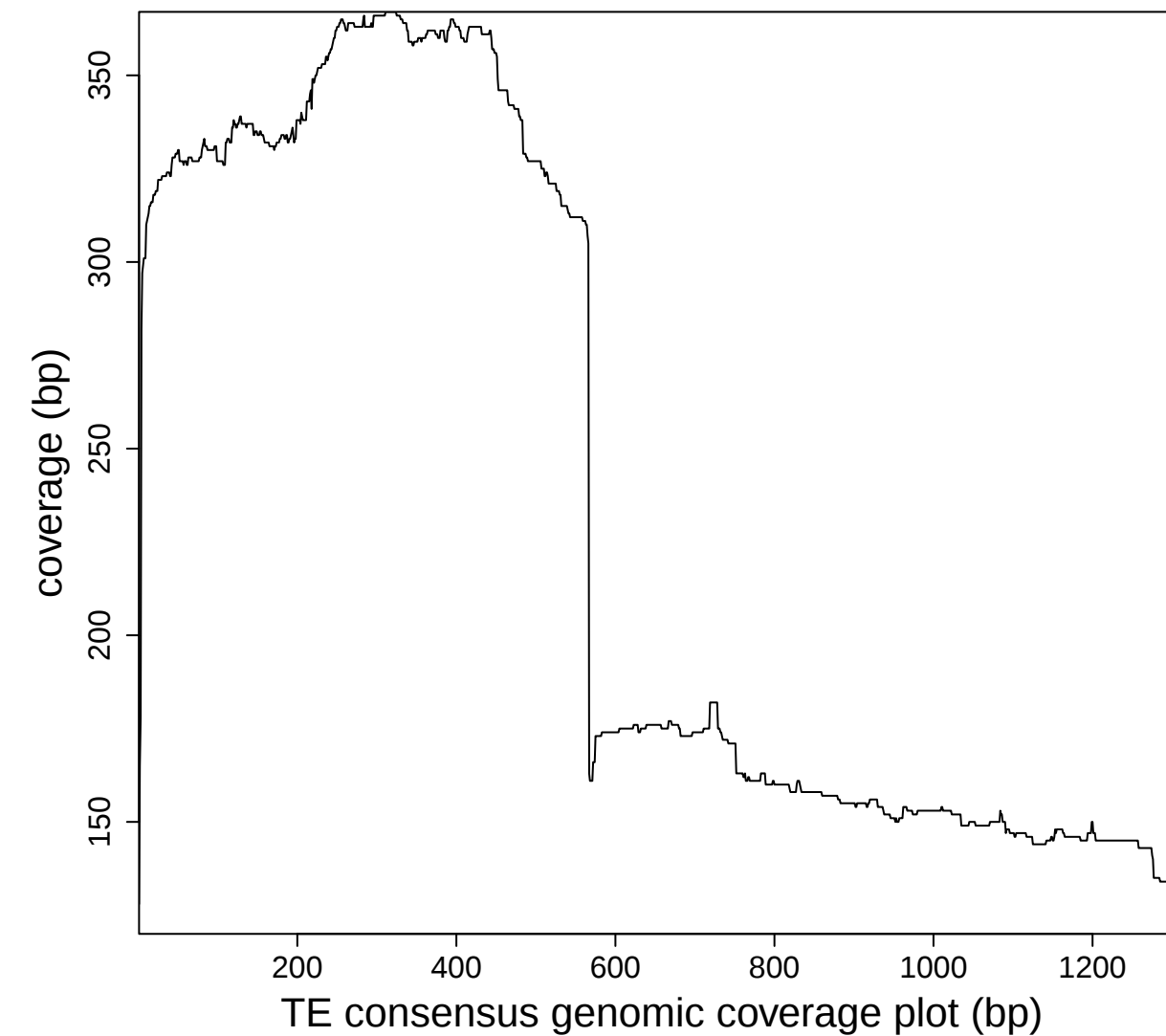
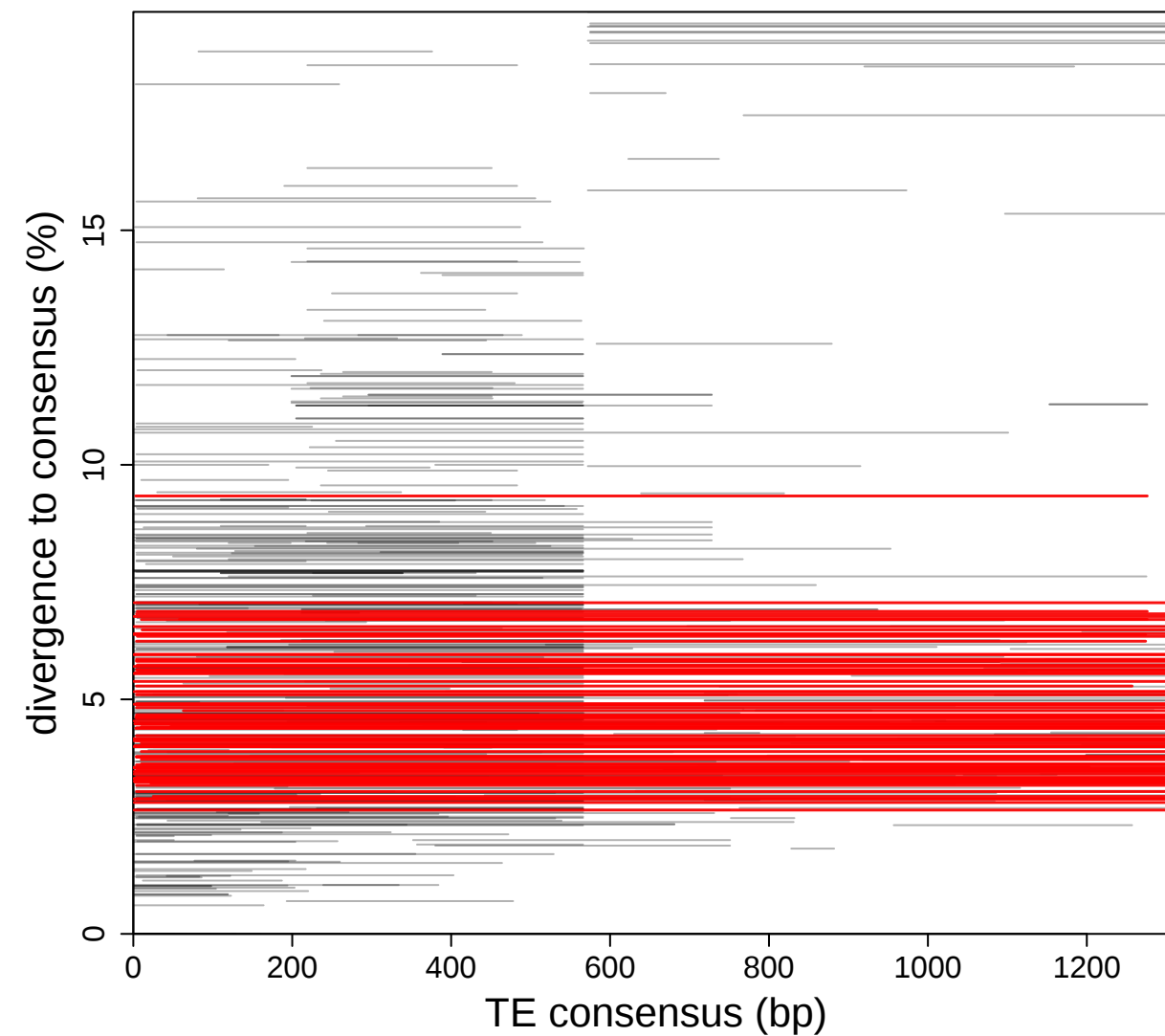
ly_1577.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 3300bp; fragments: 986; full length: 16 (>=2970bp)

After TEtrimmer Extended plot Blue lines are boundaries

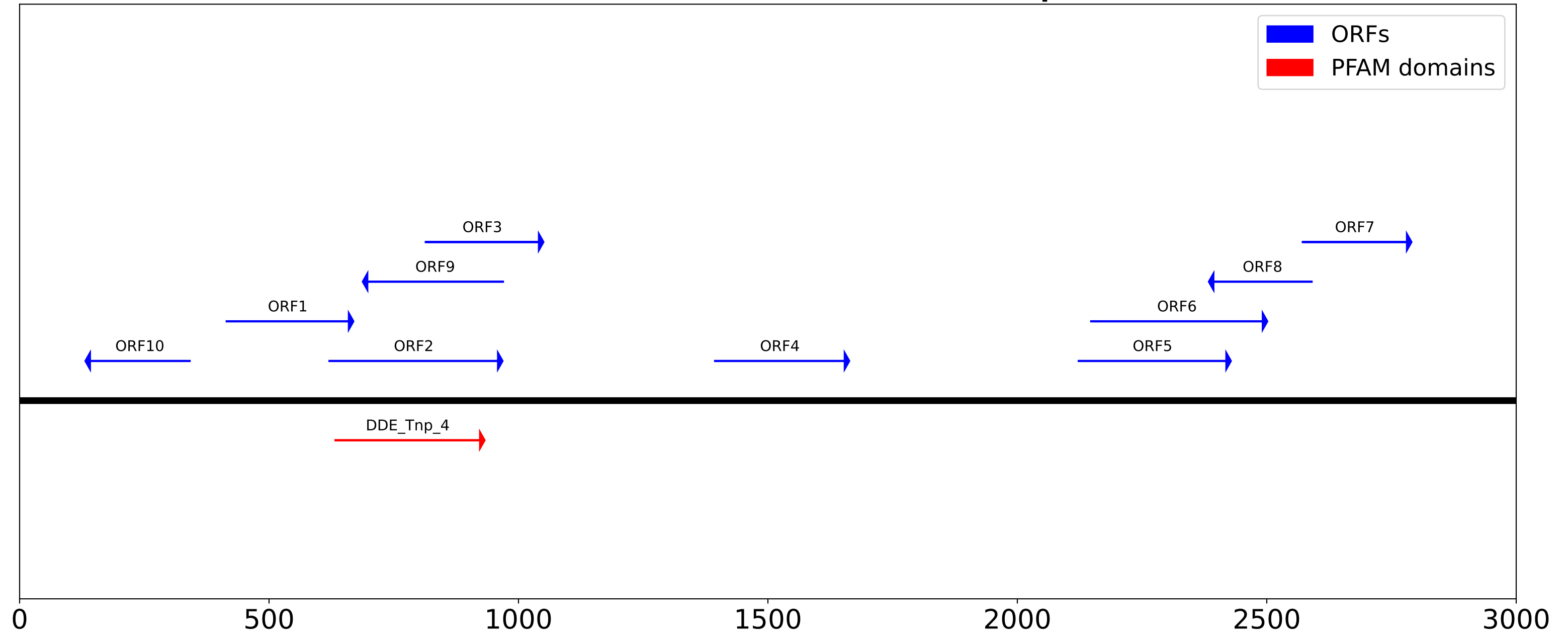


TE: rnd_5_family_1577
size: 1316bp; fragments: 529; full length: 87 (≥ 1184.4 bp)

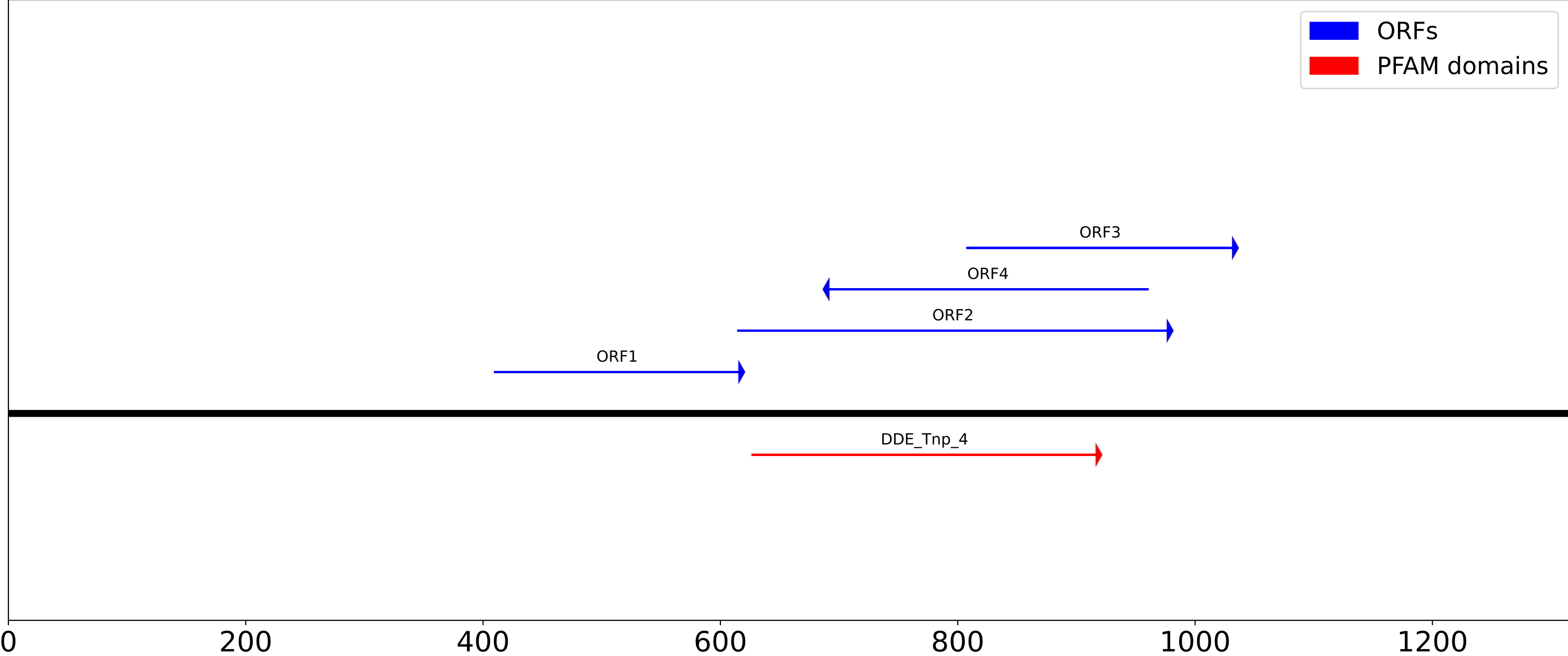
Before TEtrimmer 1316 bp



After TEtrimmer ORF and PFAM domain plot



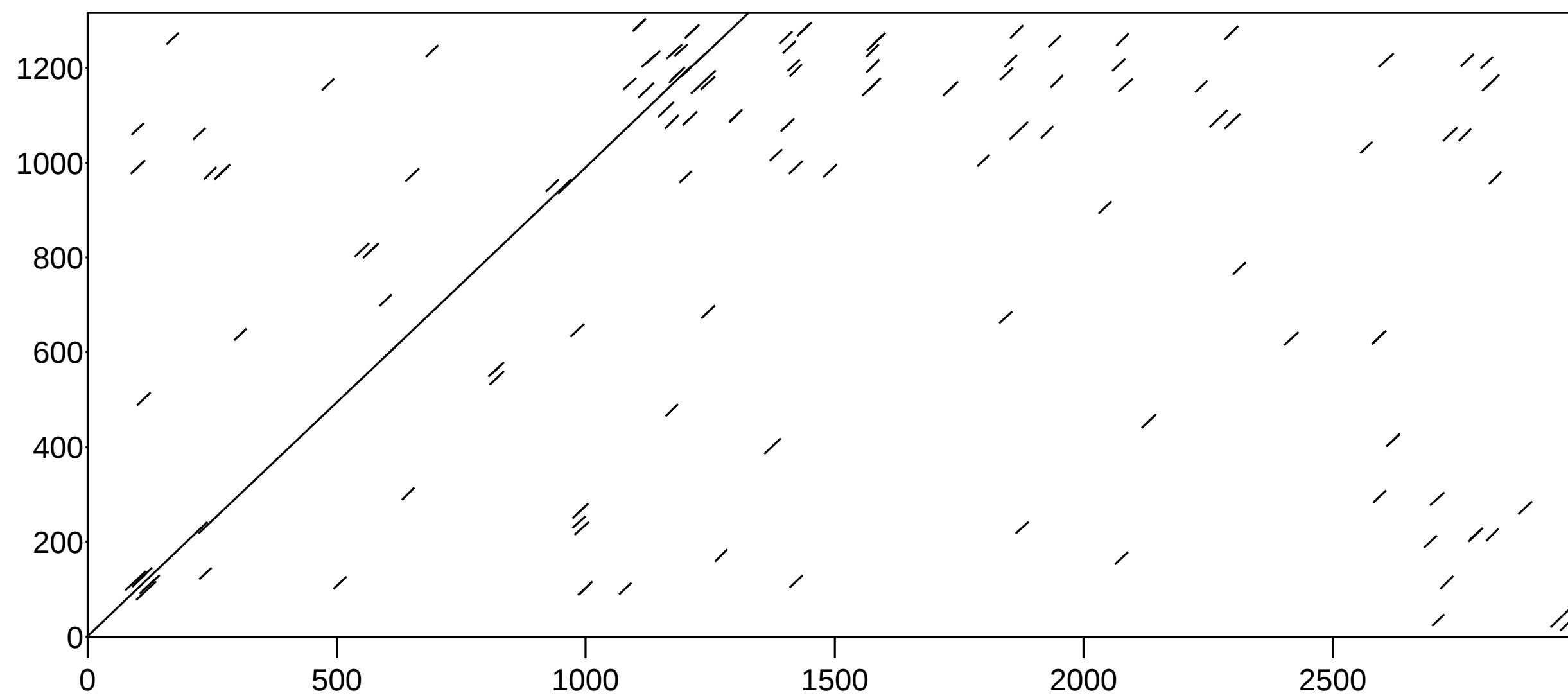
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)