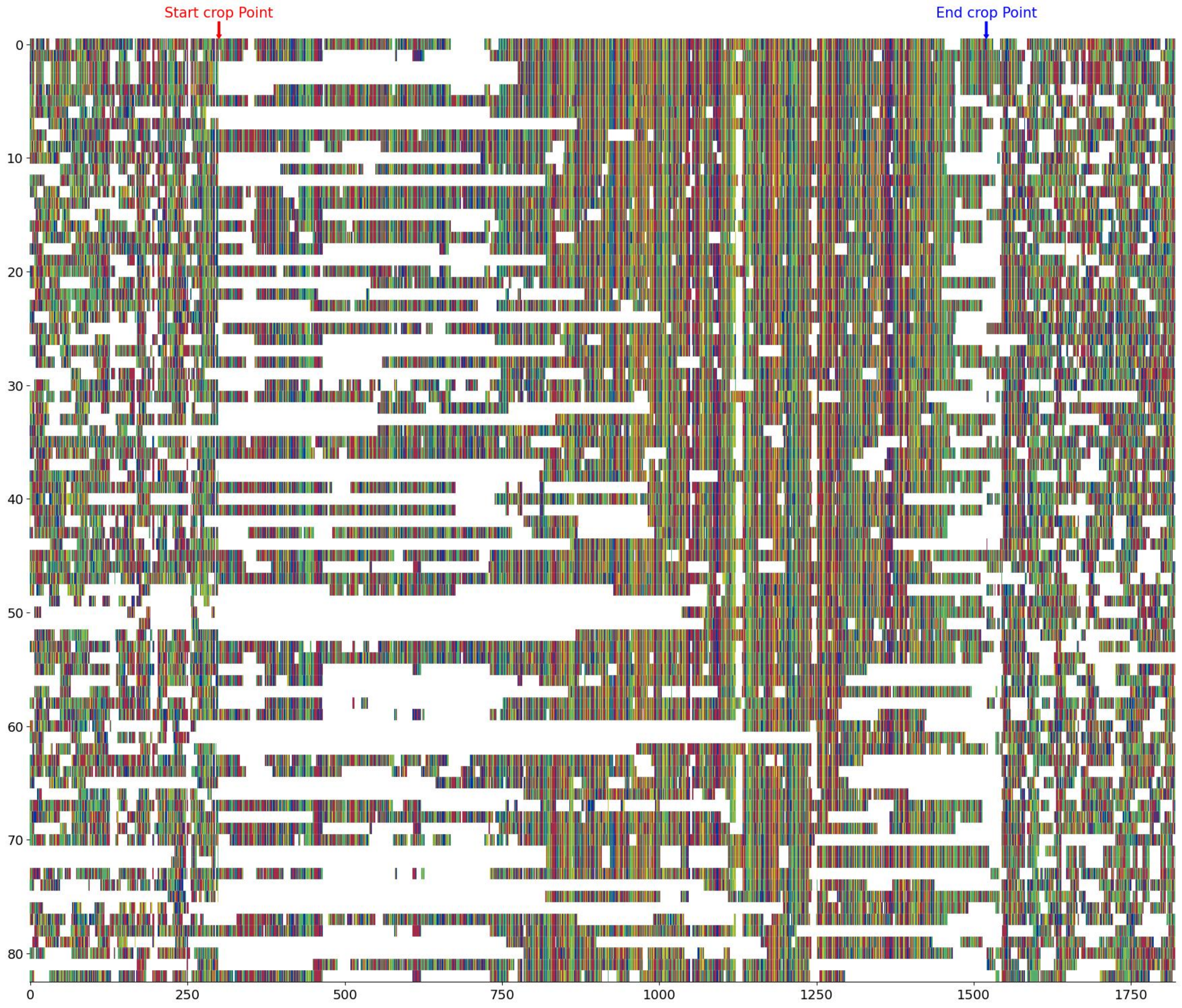




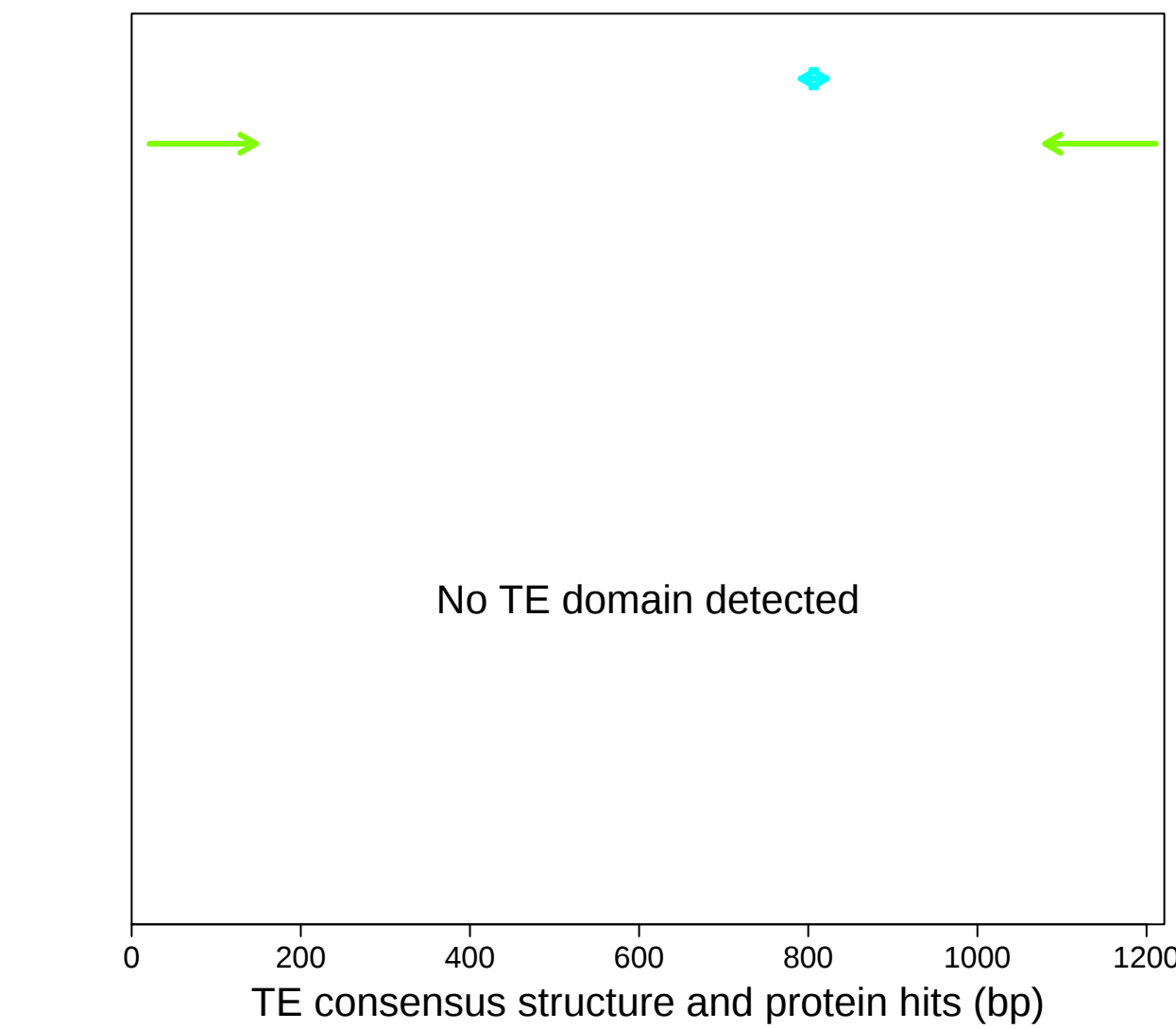
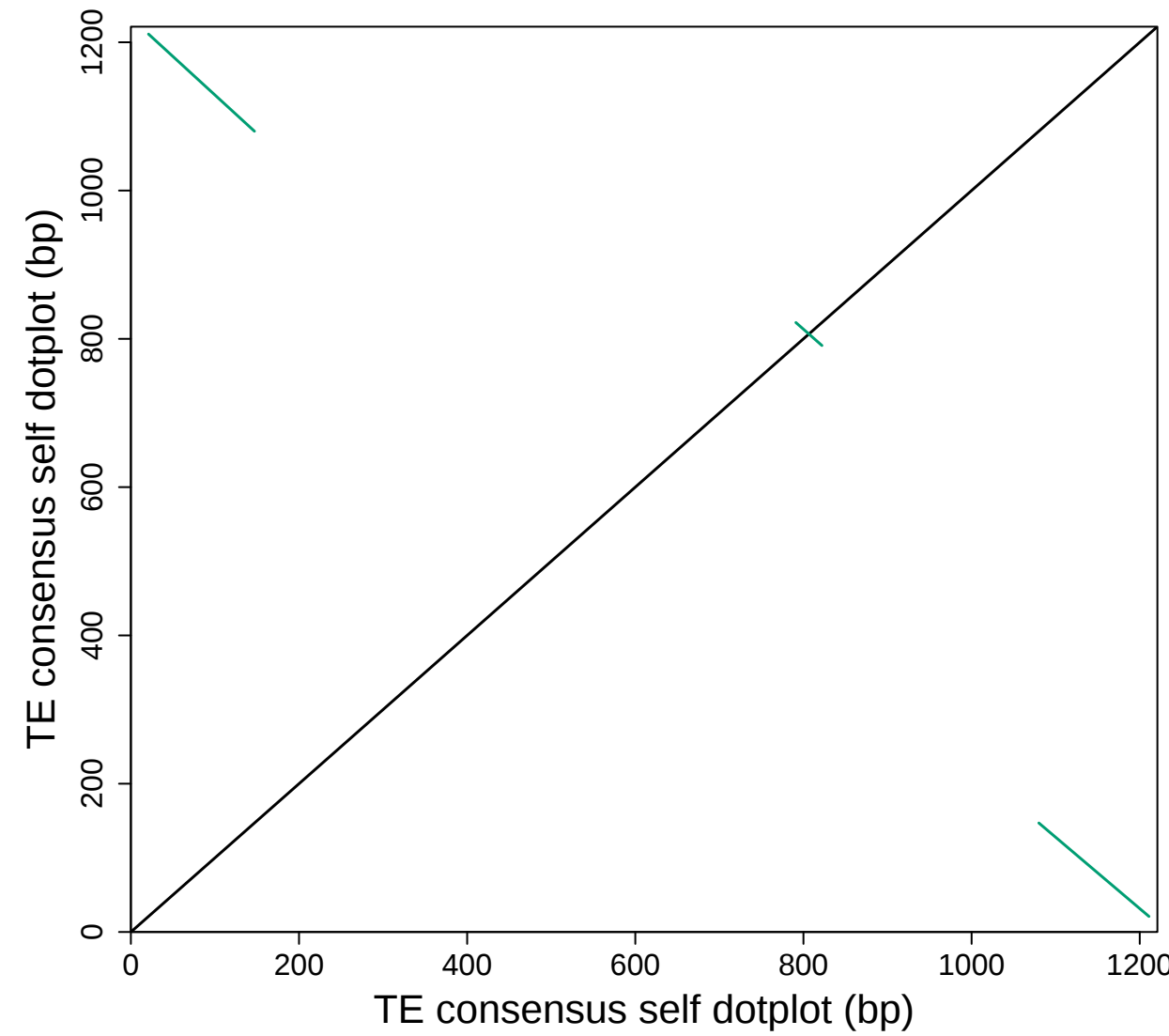
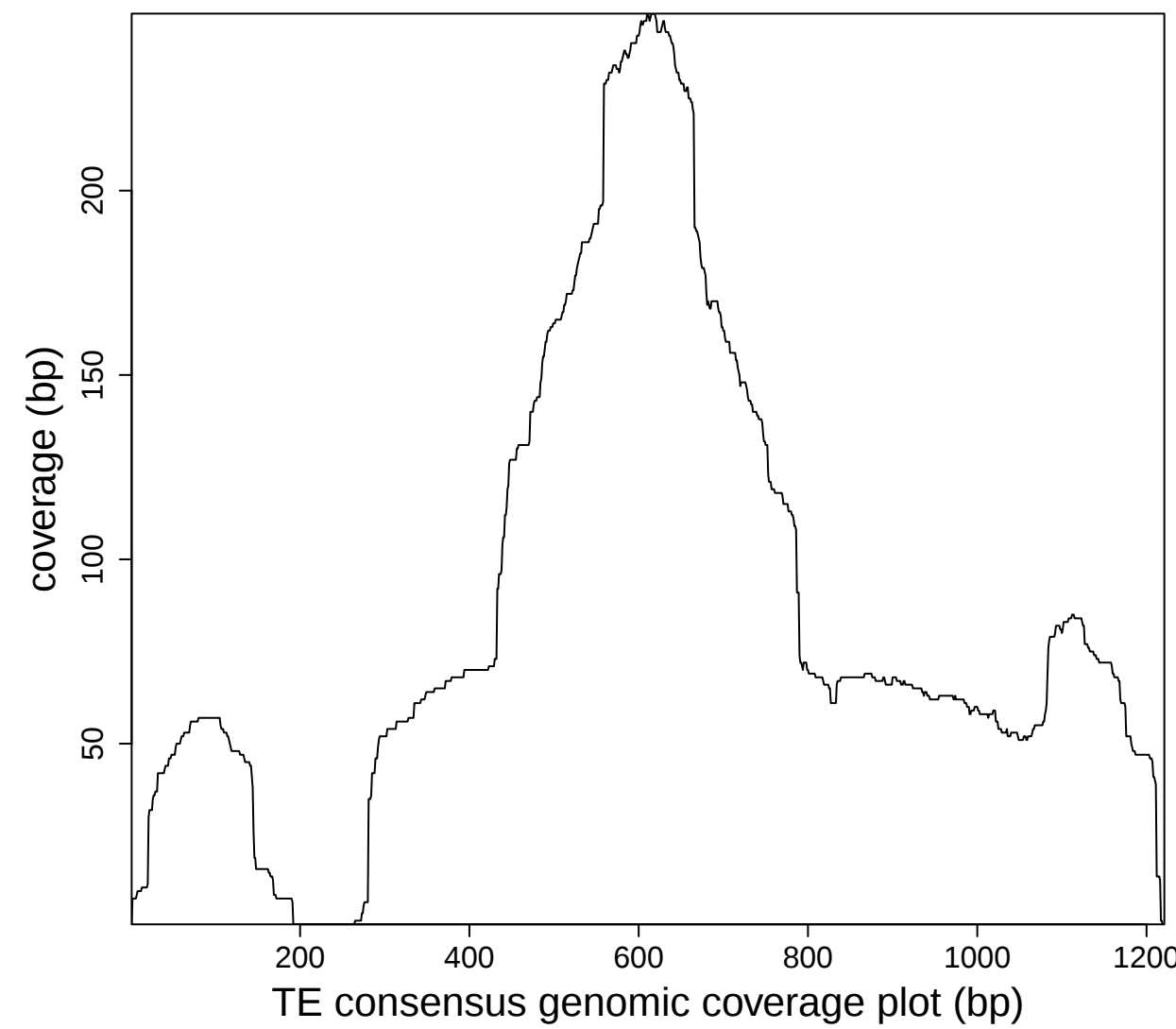
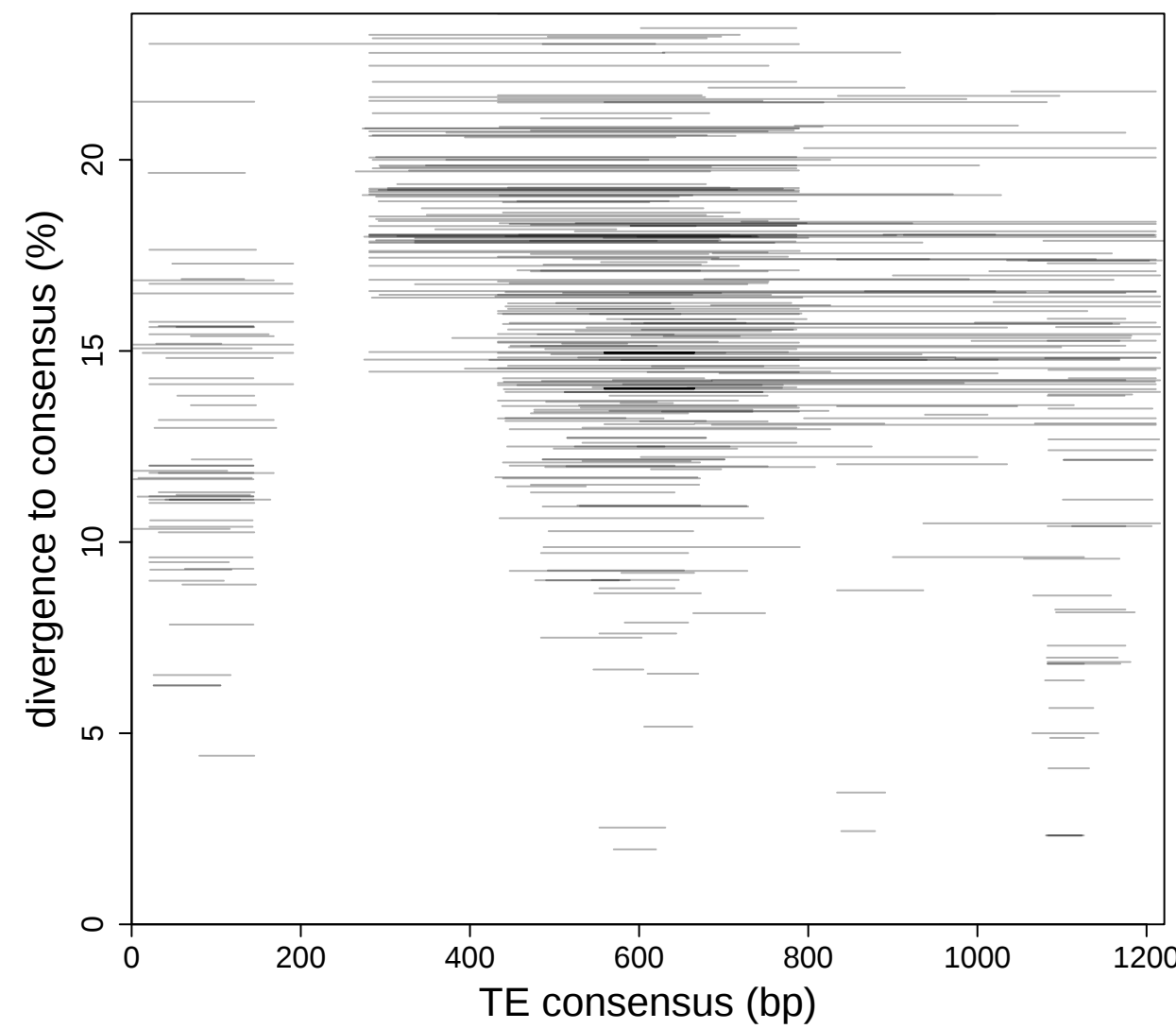
Start crop Point

End crop Point



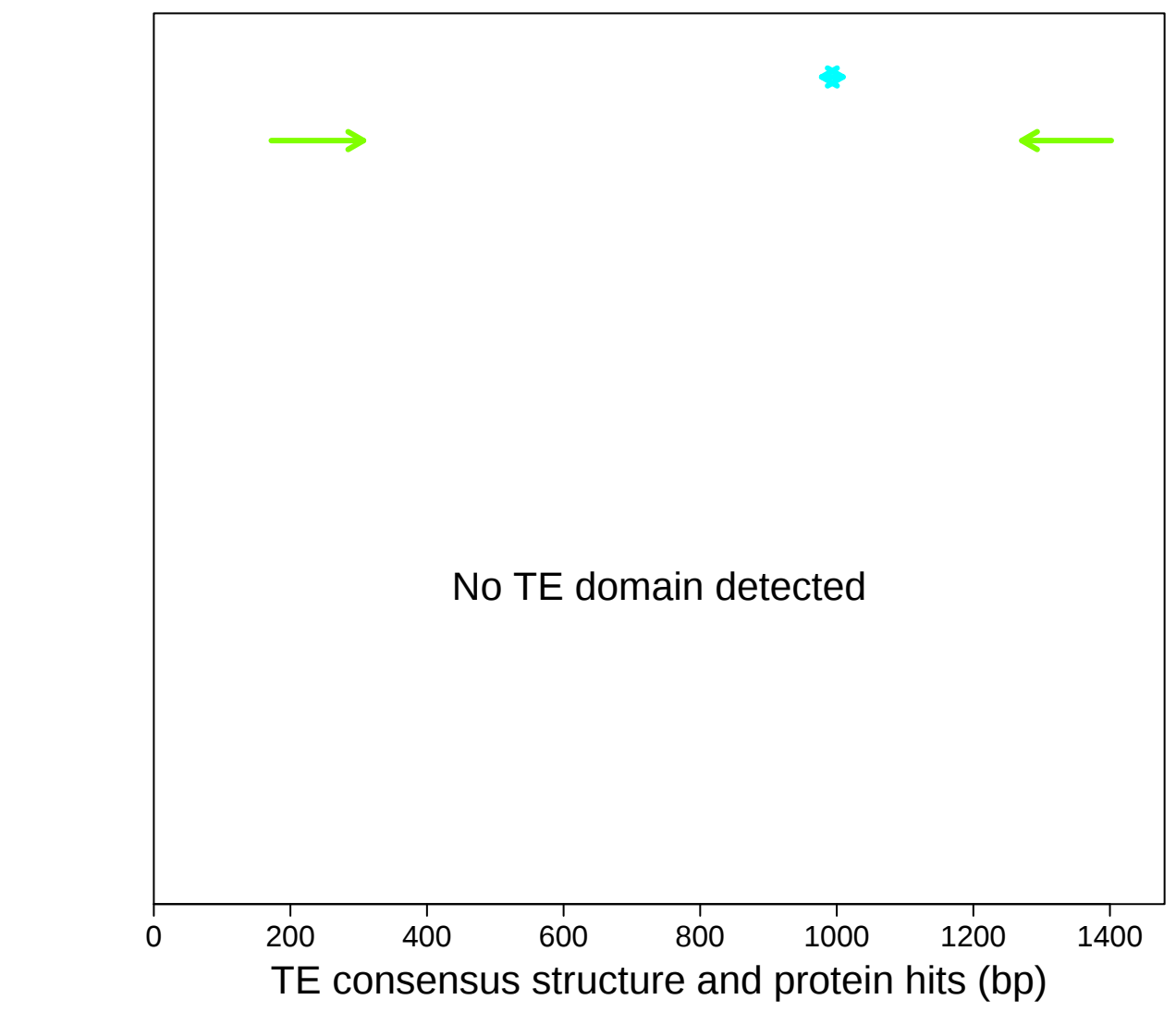
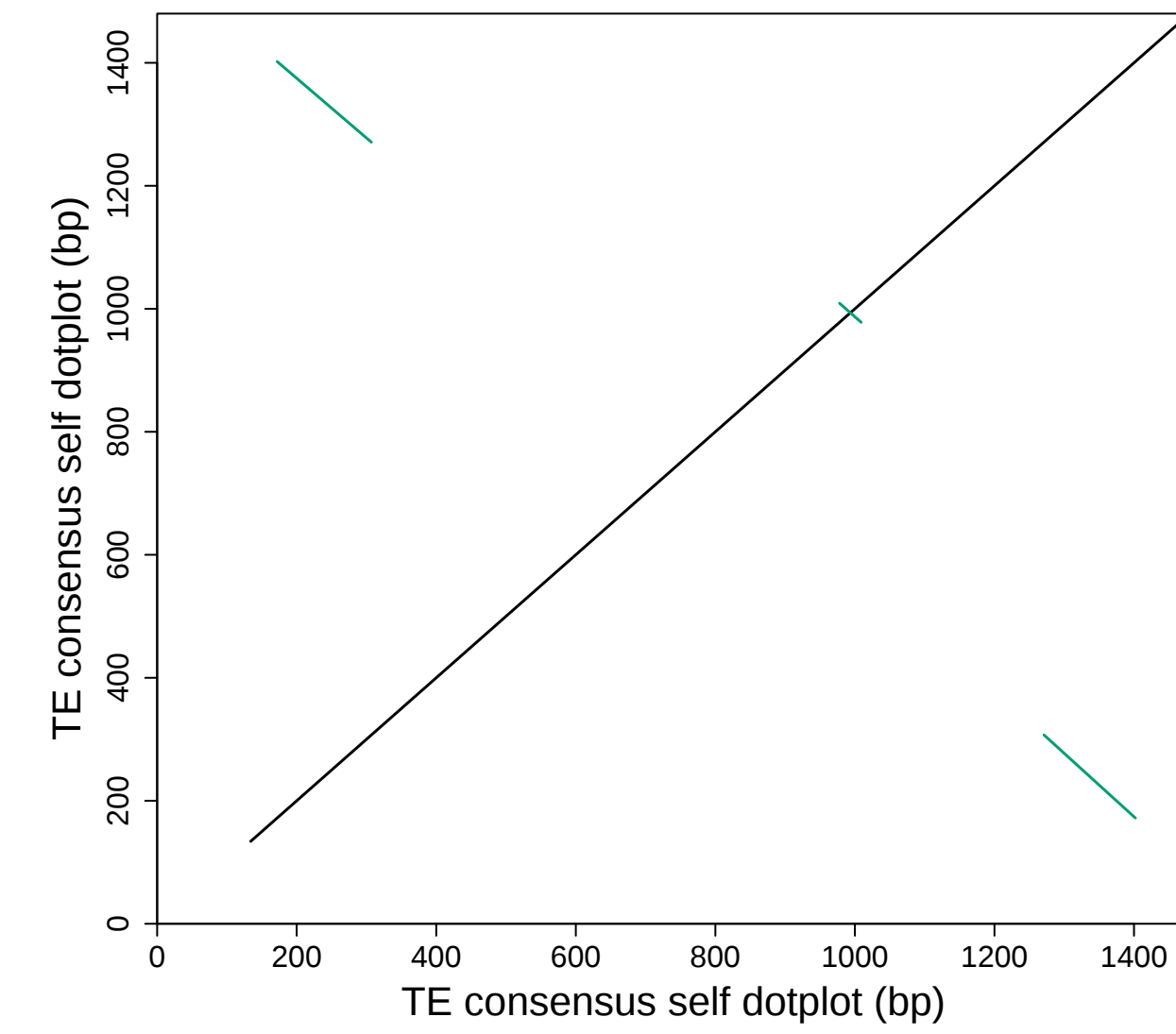
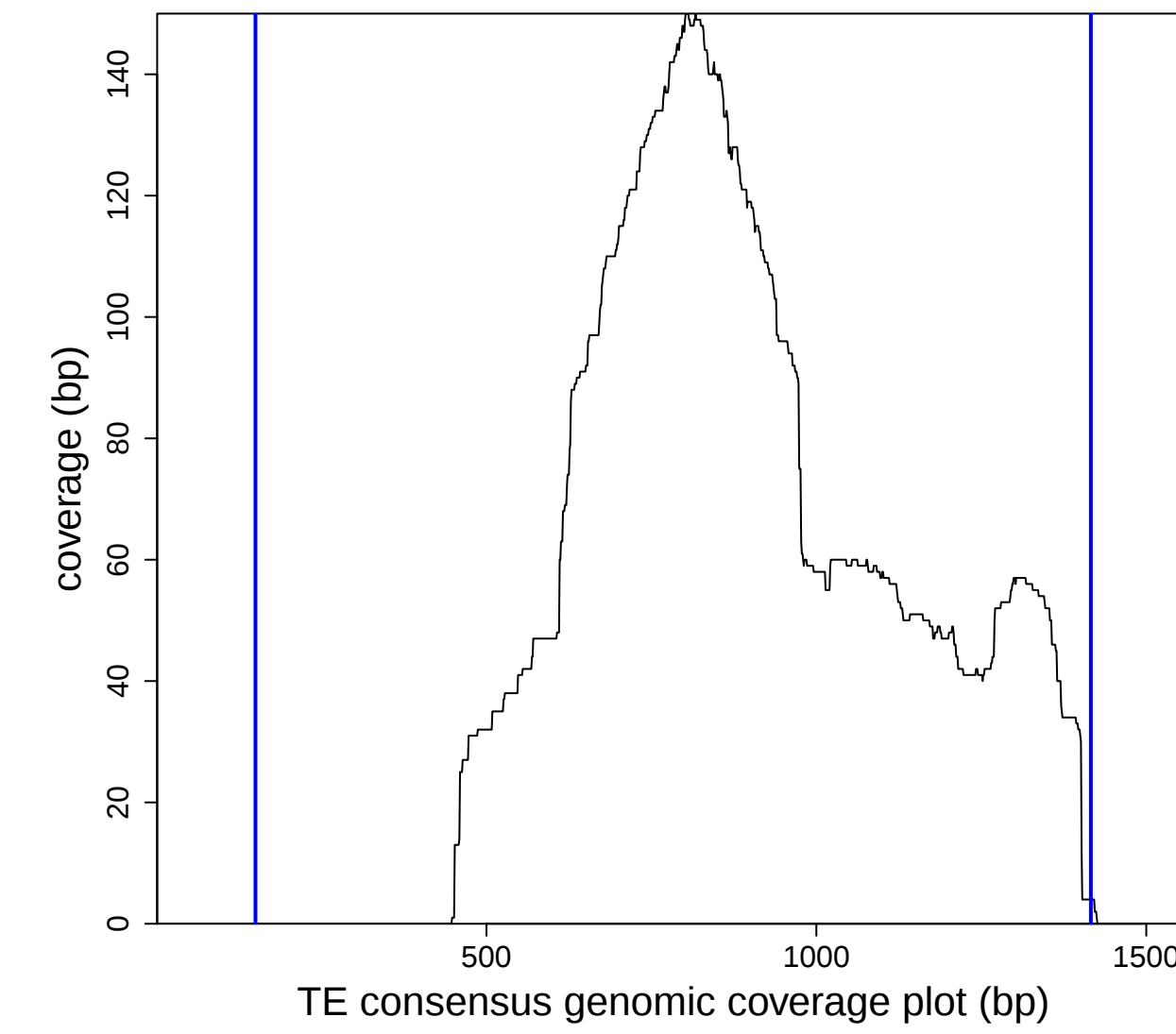
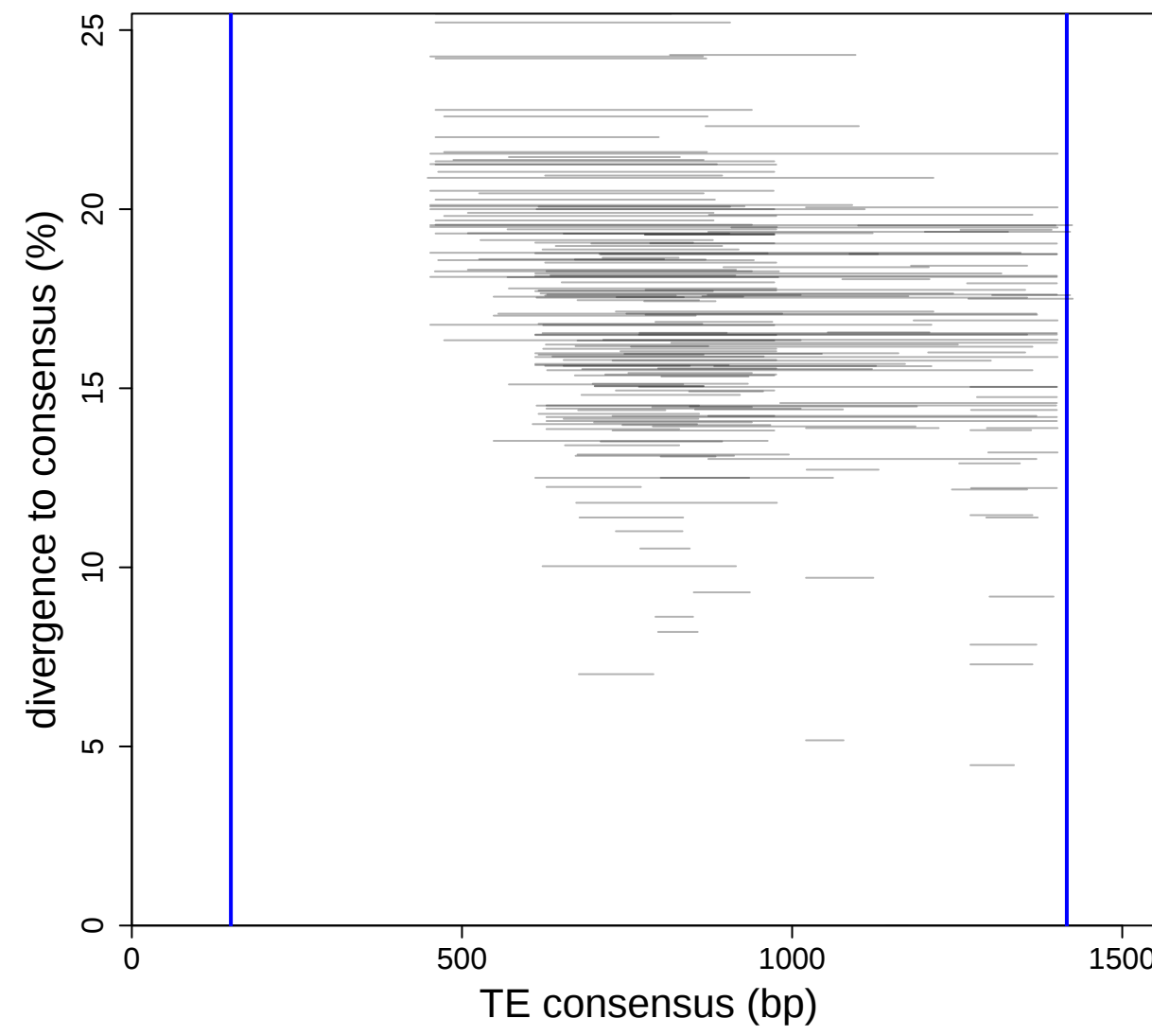
ed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 1221bp; fragments: 393; full length: 0 (>=1098.9bp)

After TEtrimmer 1221 bp

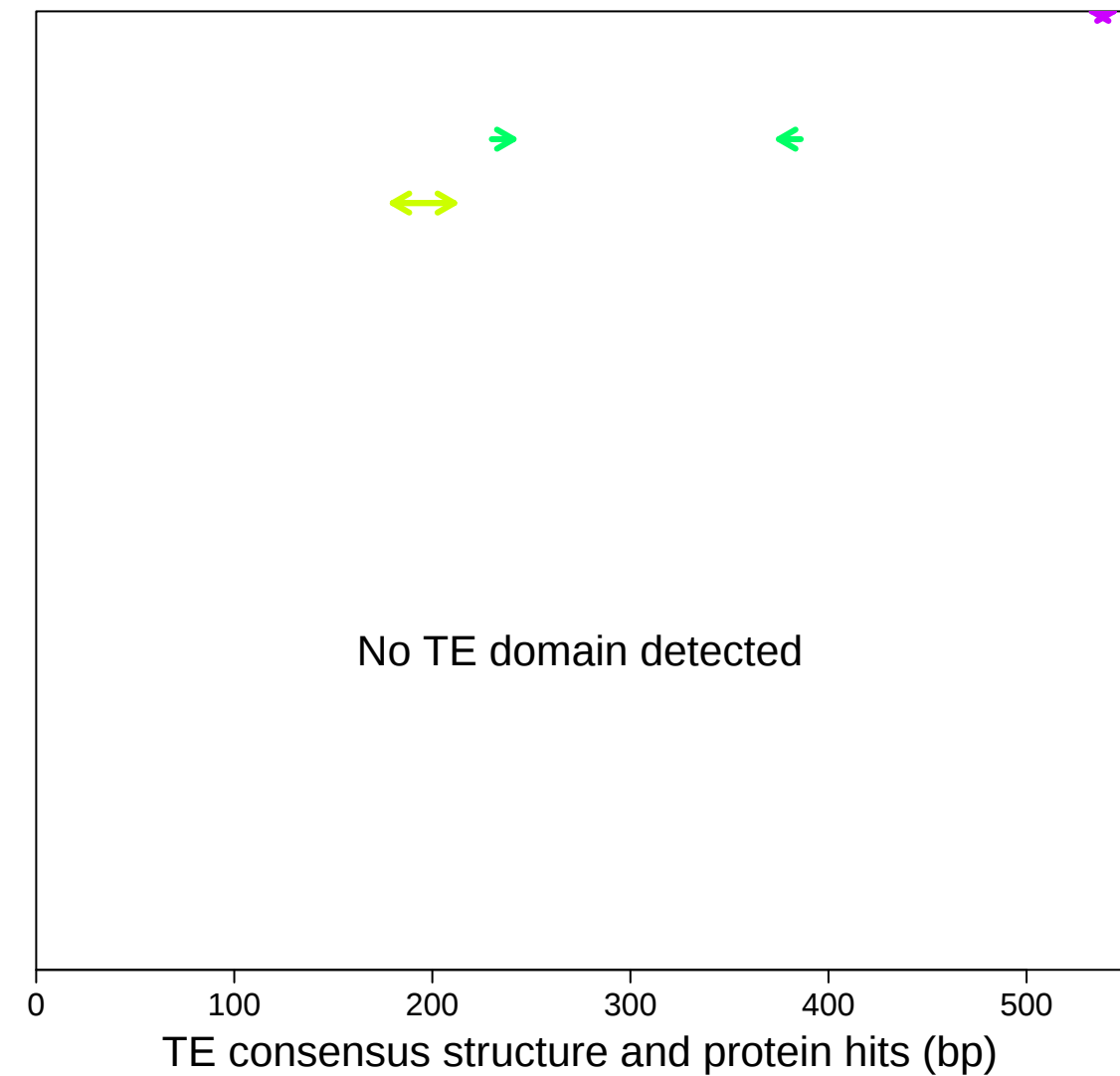
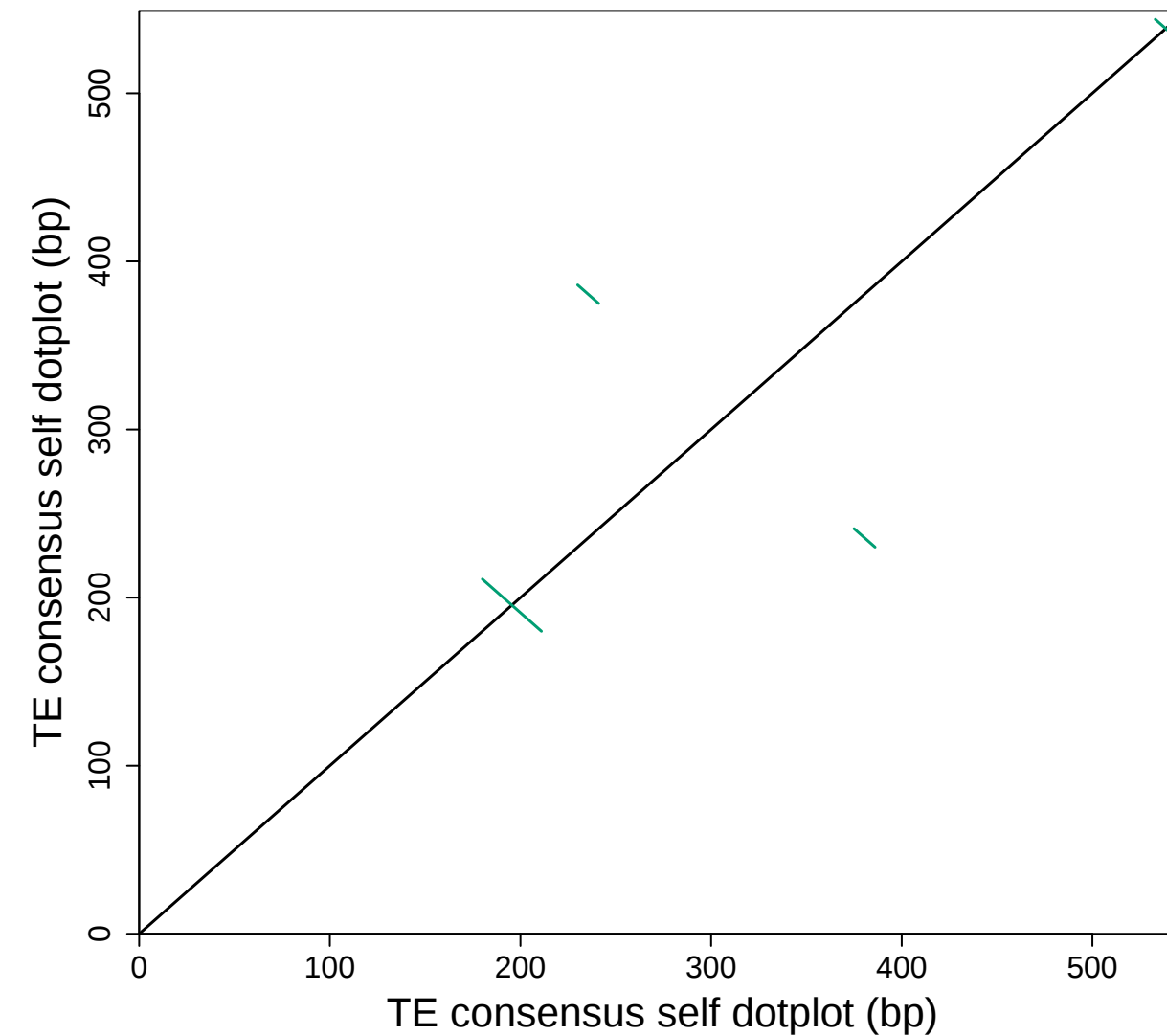
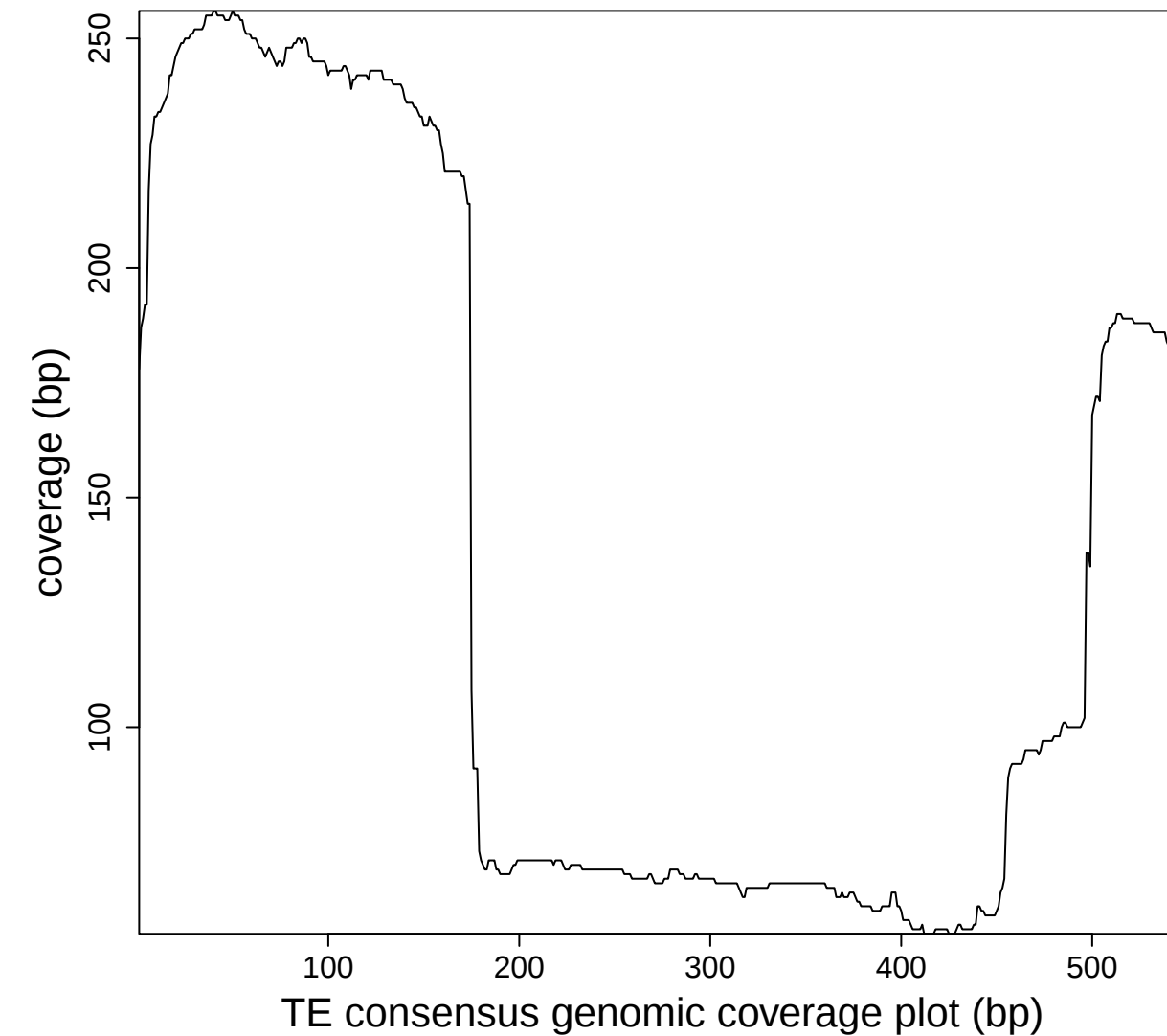
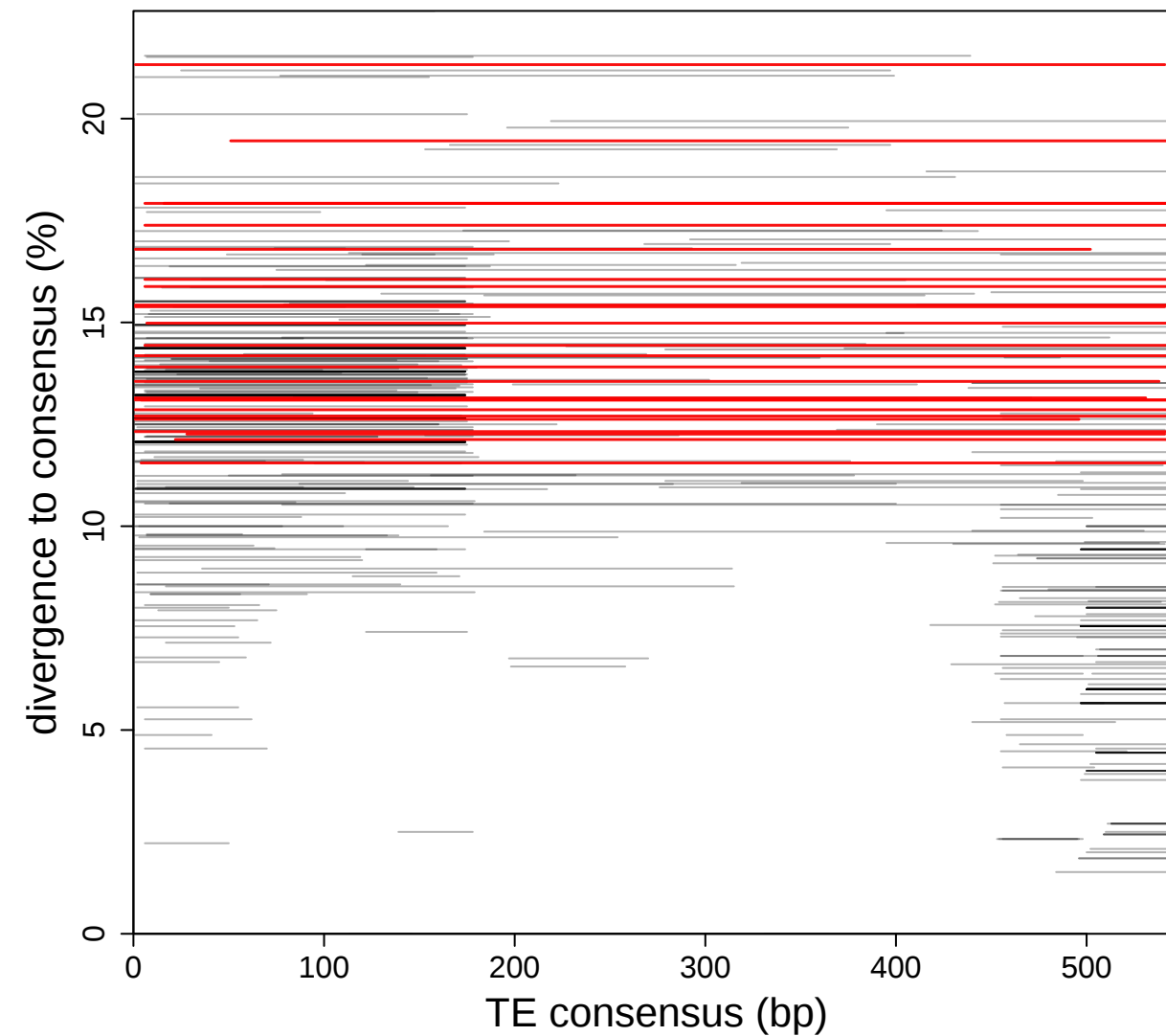


hily_76.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcIn.fa_aln.fa
size: 1566bp; fragments: 203; full length: 0 (>=1409.4bp)

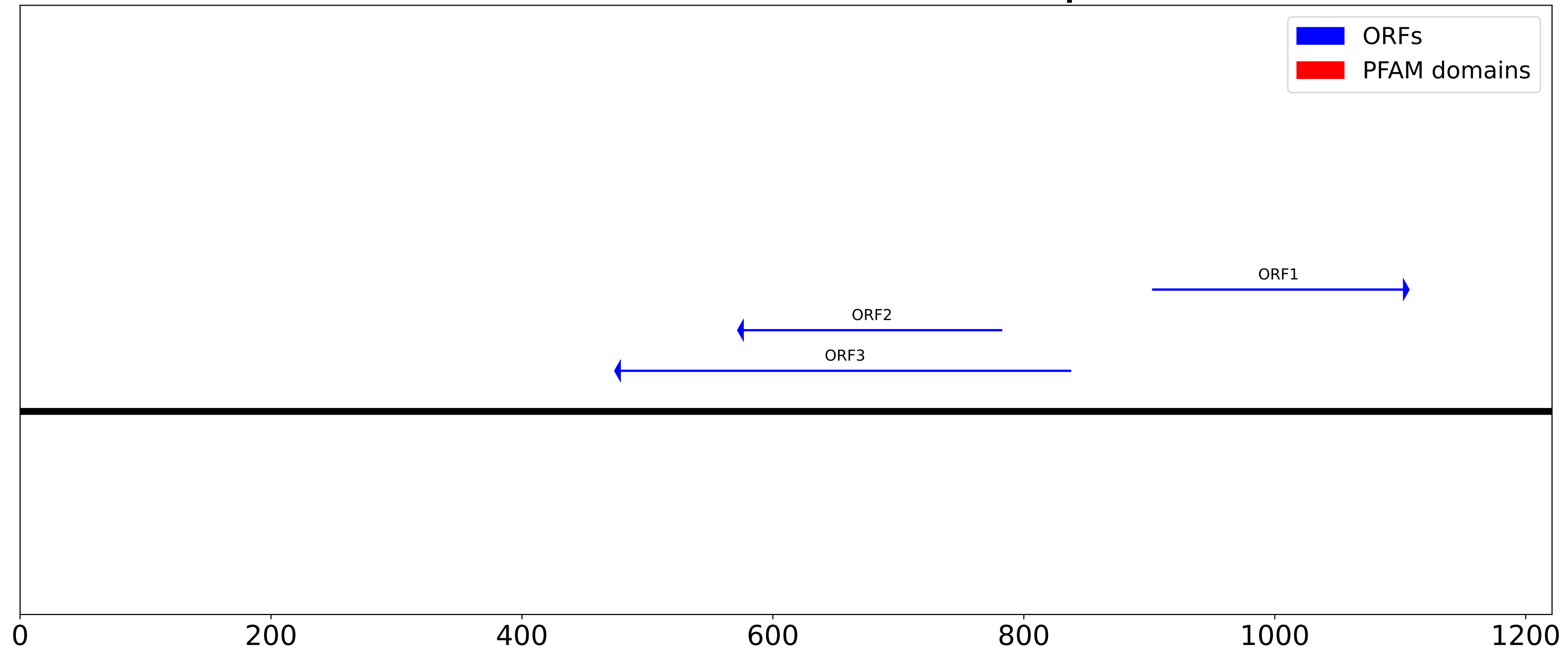
After TEtrimmer Extended plot Blue lines are boundaries



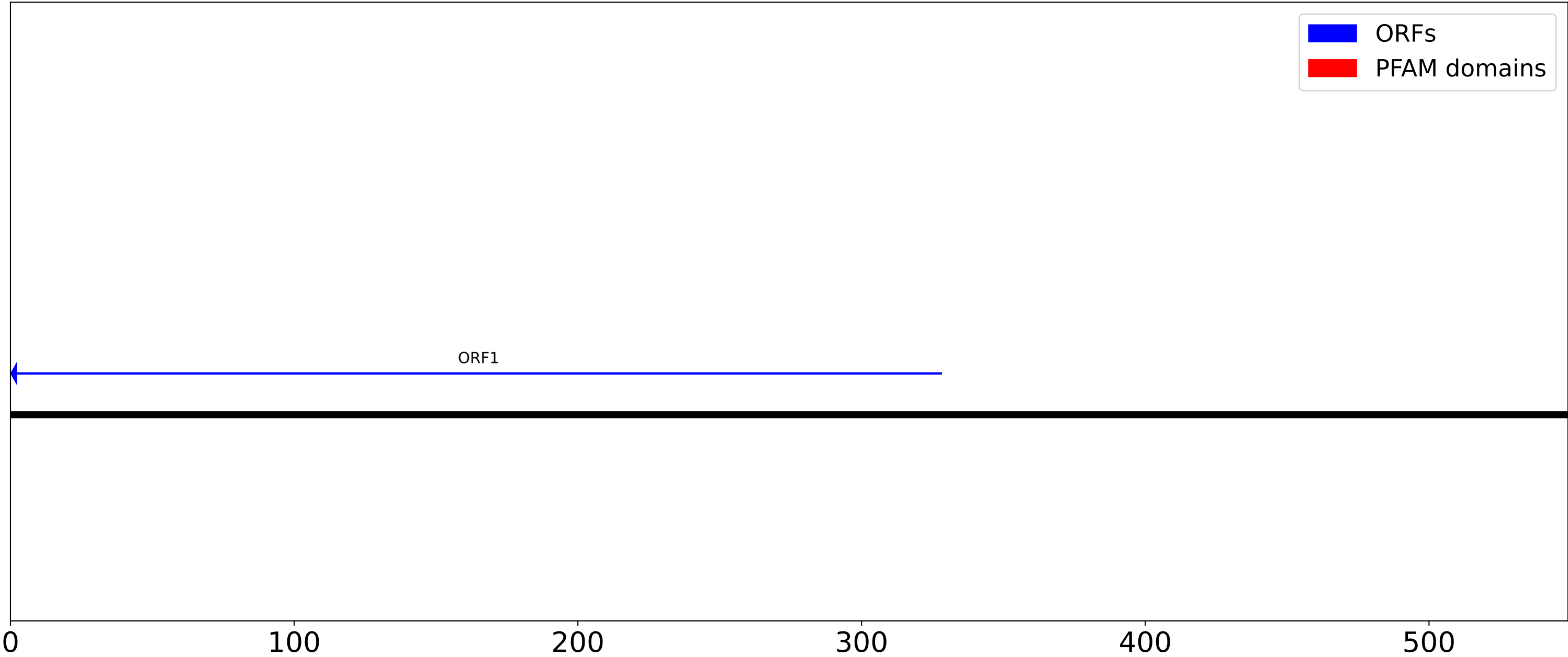
TE: rnd_5_family_76
size: 549bp; fragments: 468; full length: 25 (≥ 494.1 bp)



After TEtrimmer ORF and PFAM domain plot



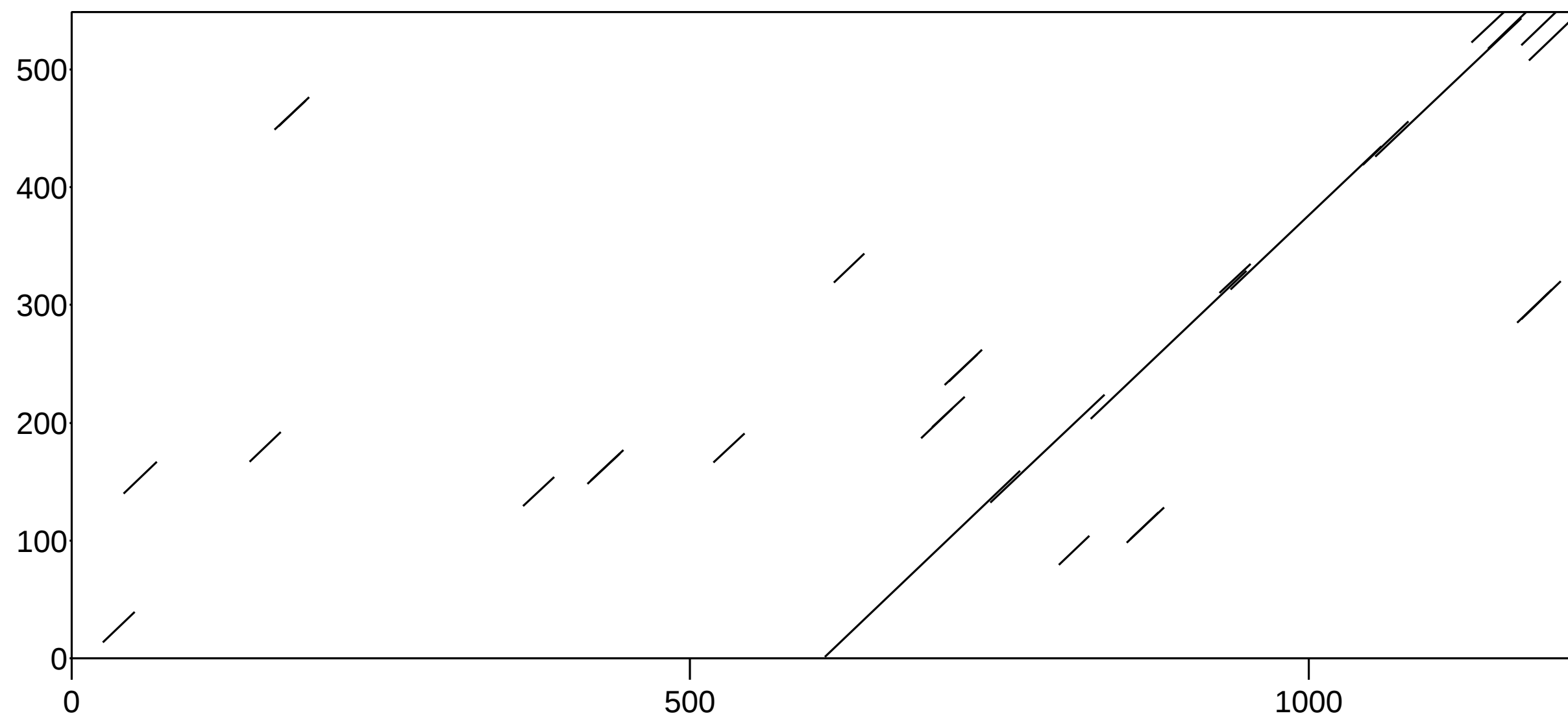
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)