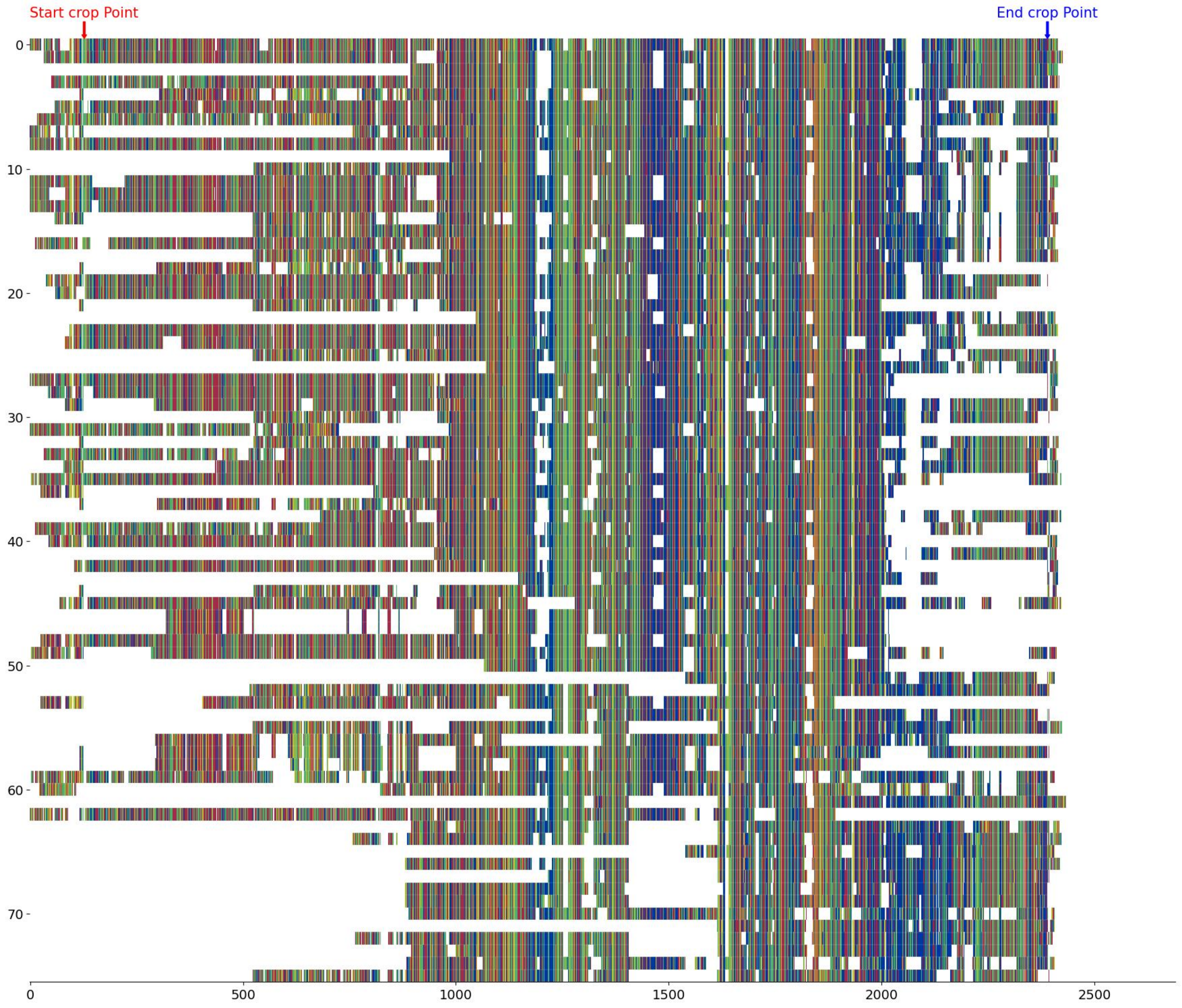




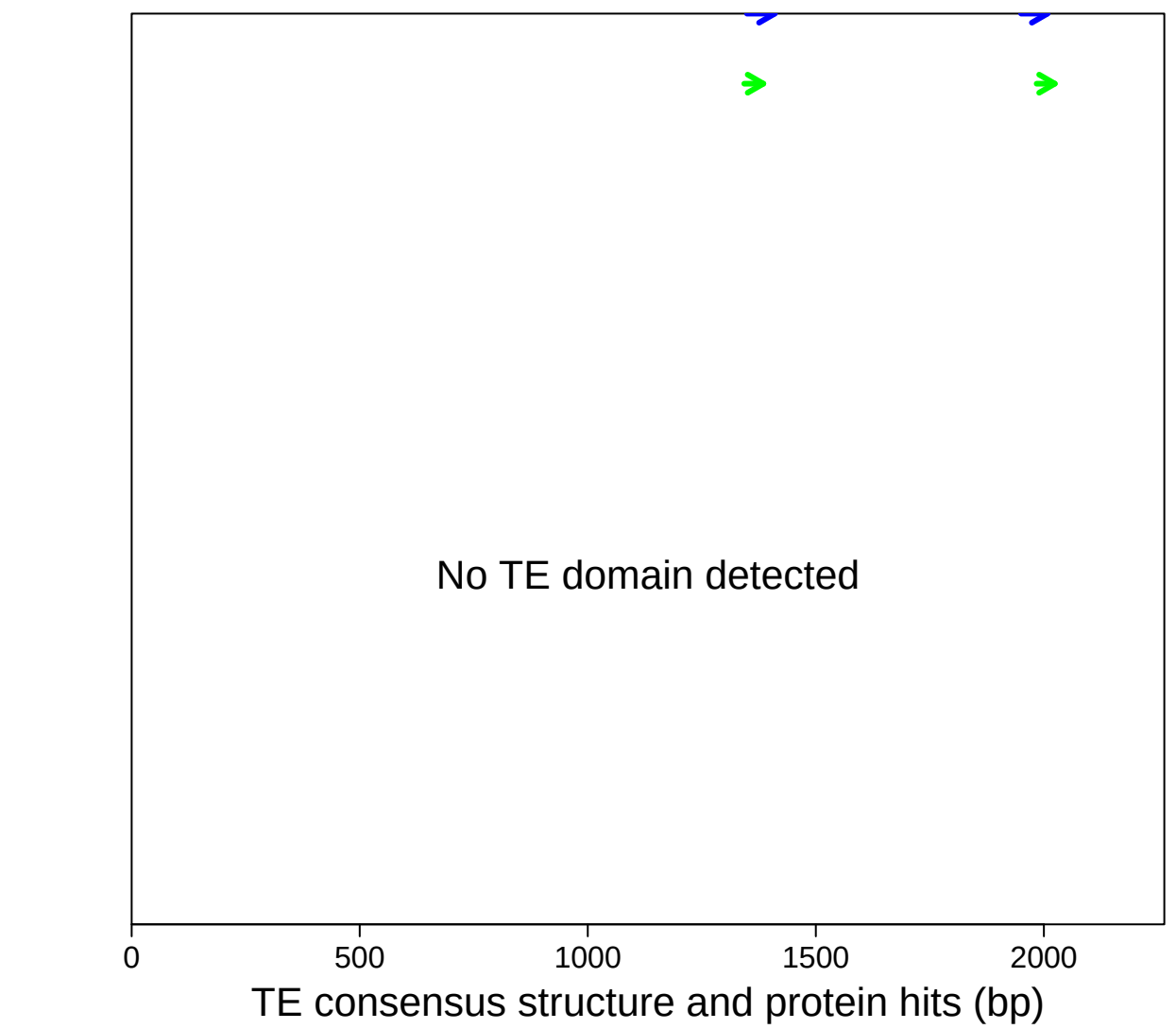
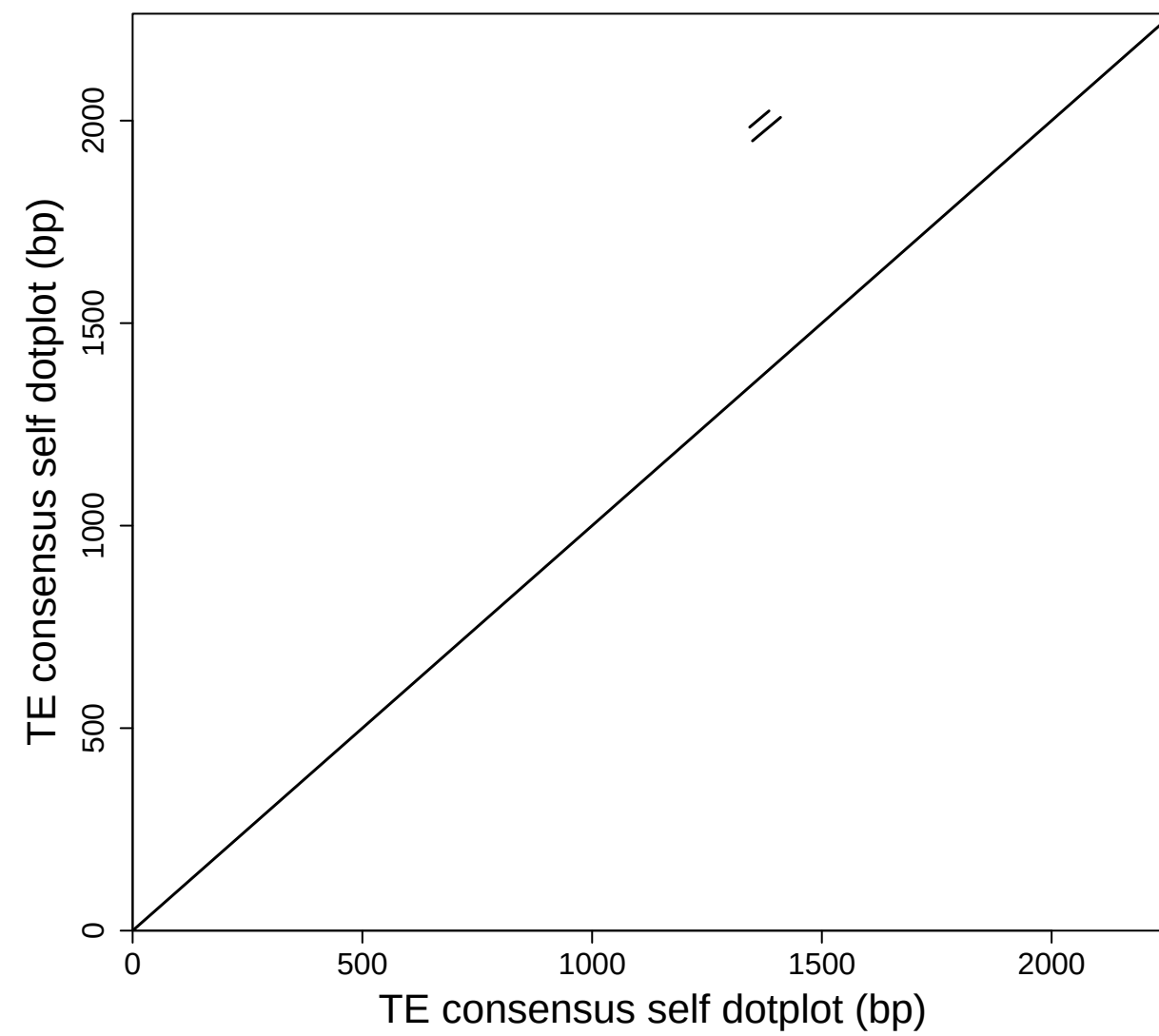
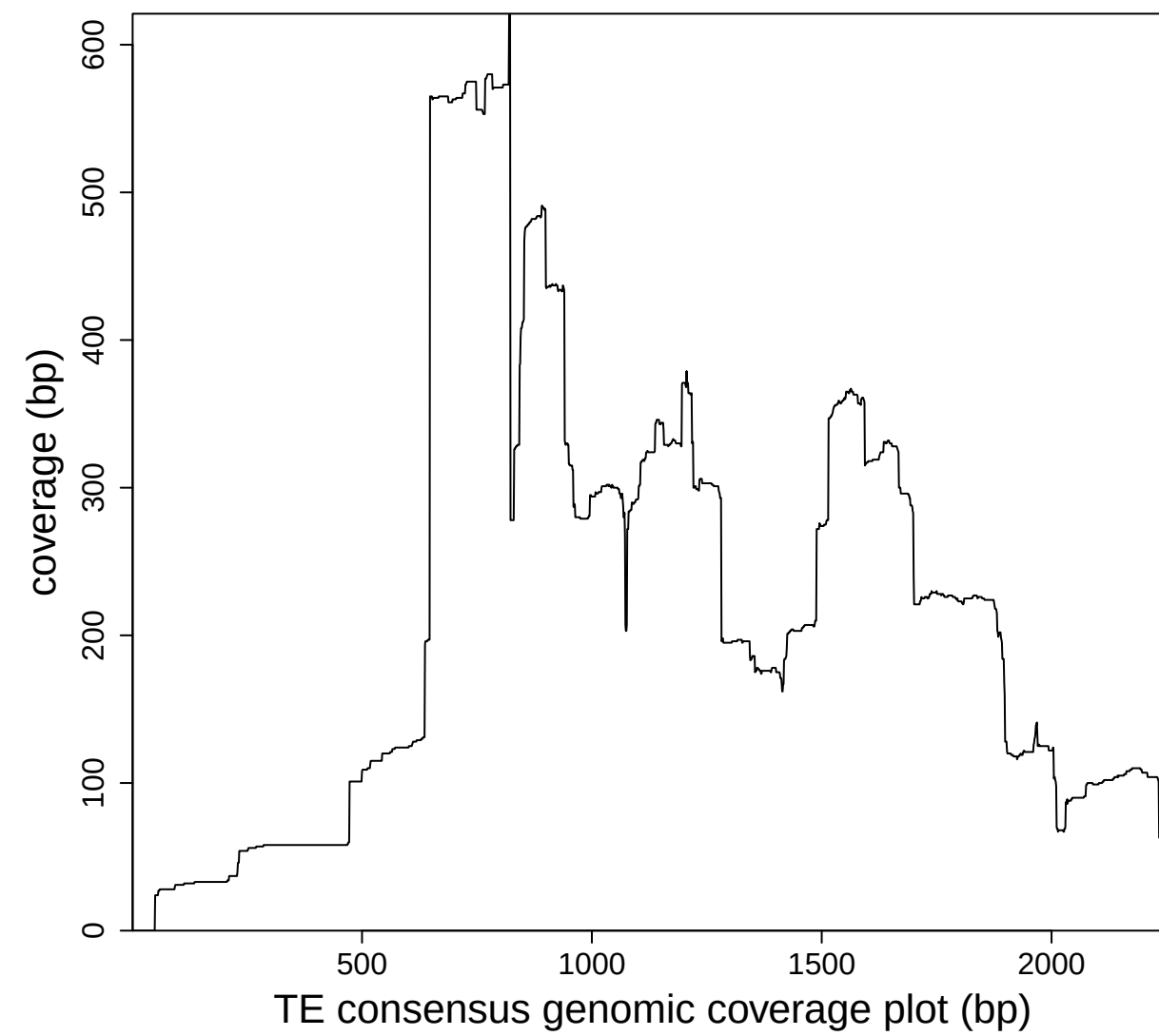
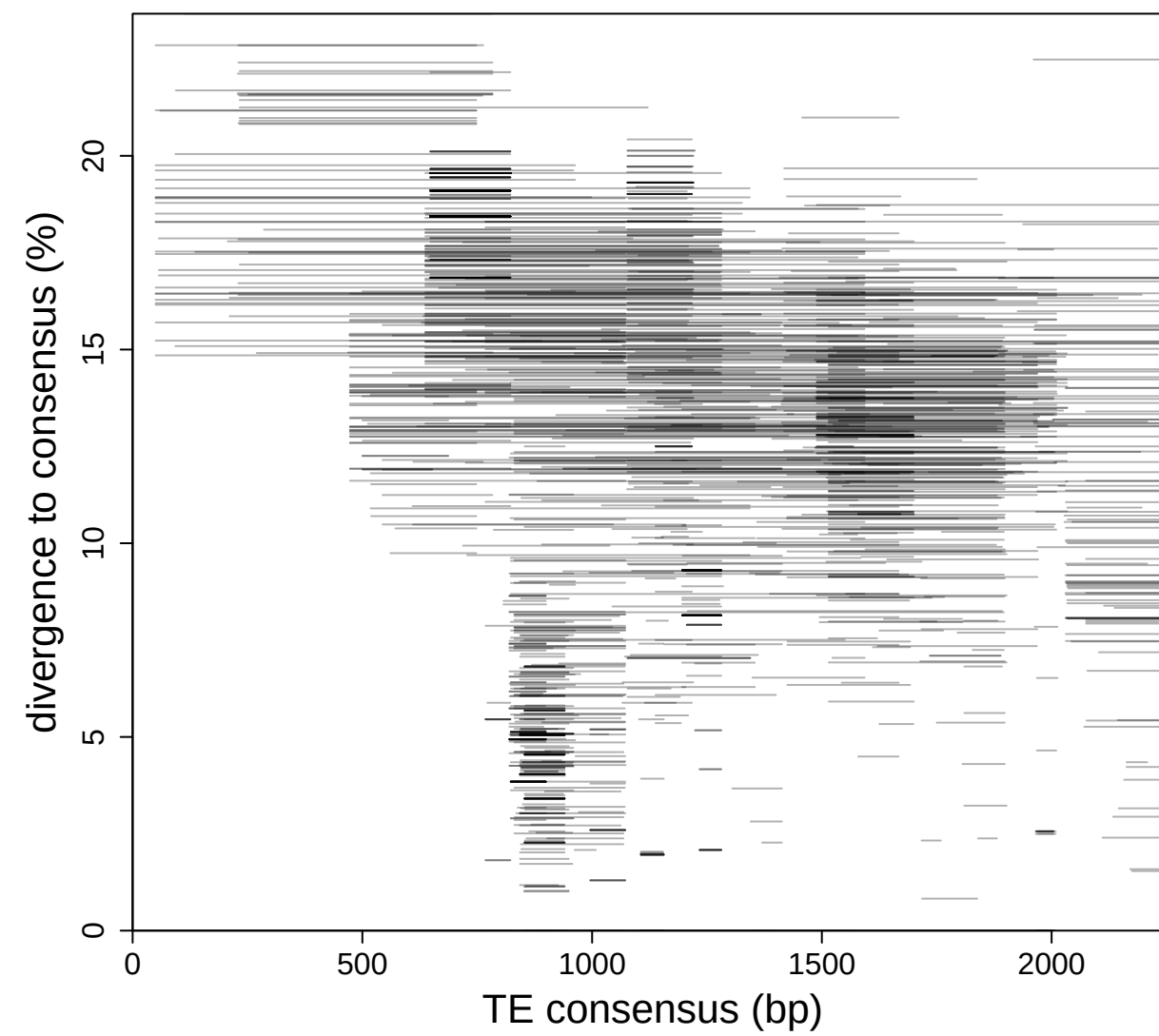
Start crop Point

End crop Point



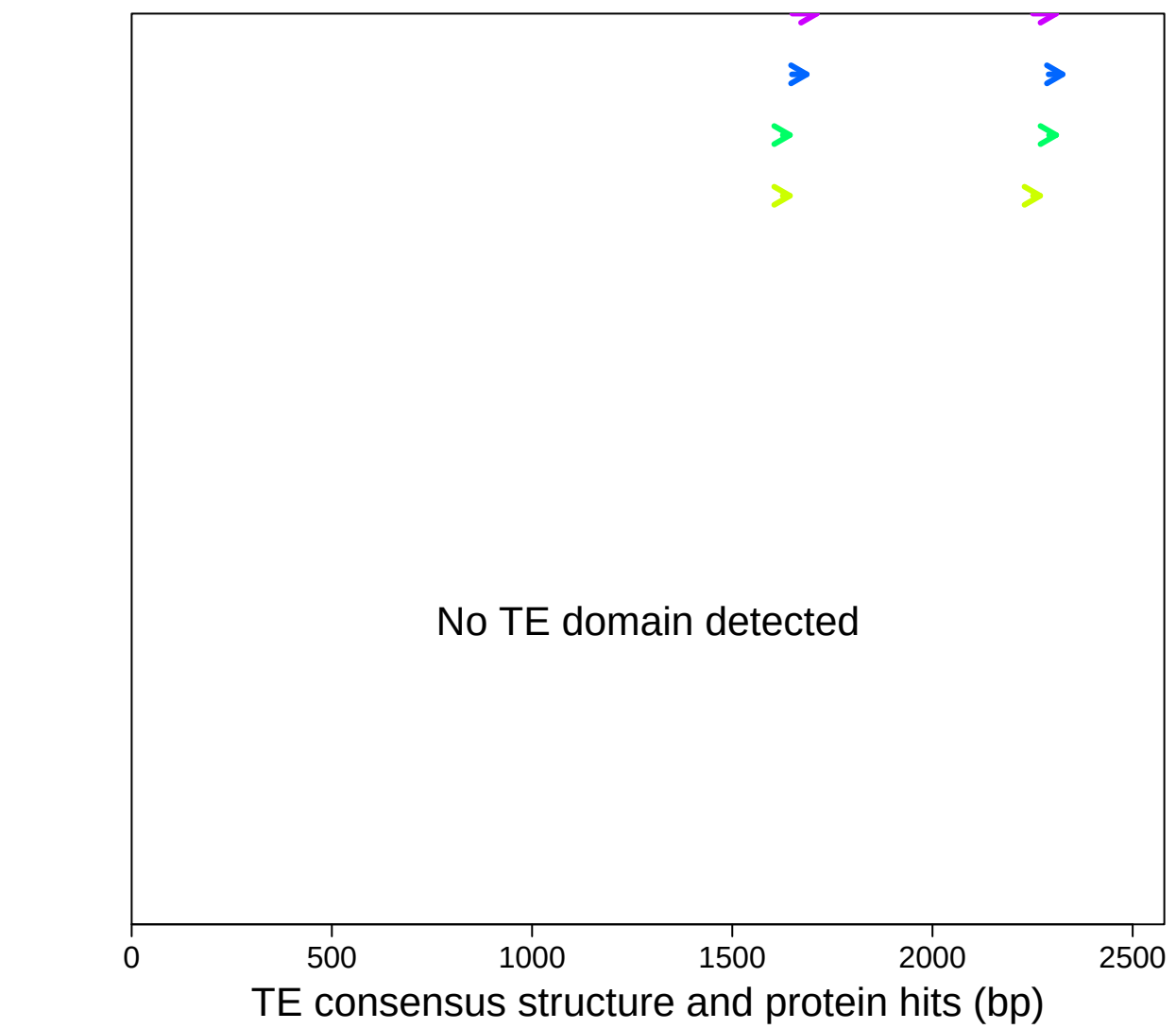
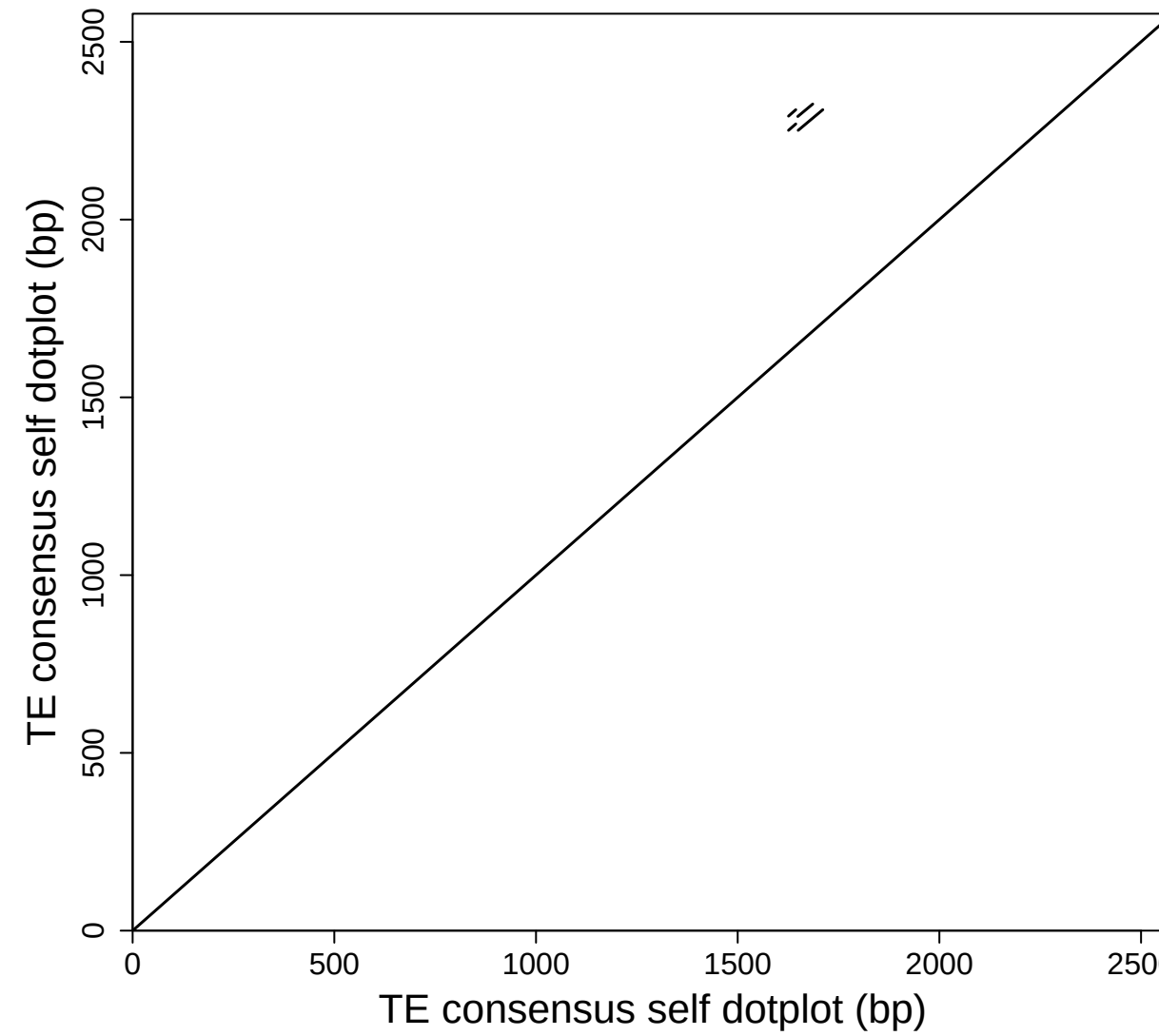
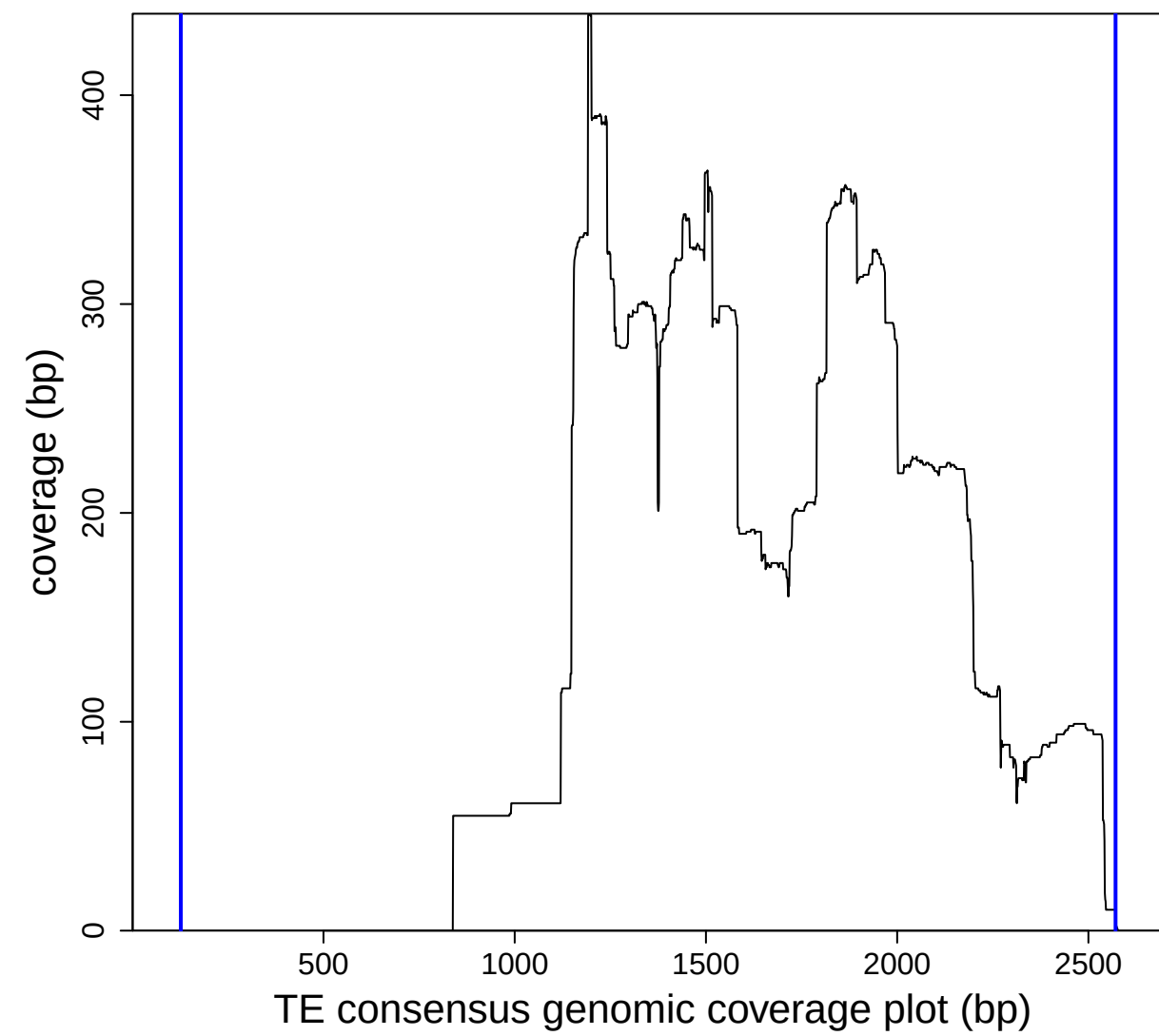
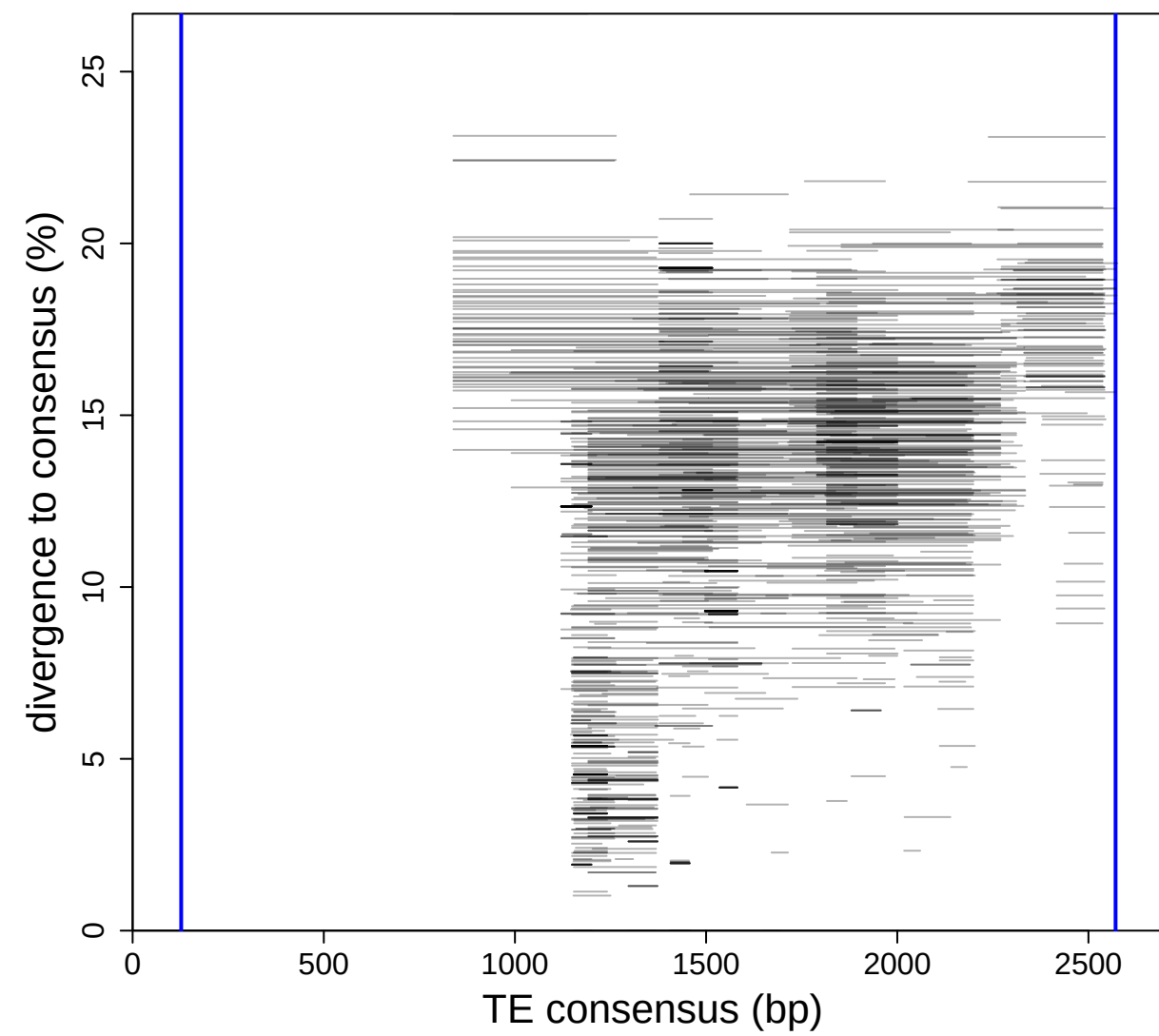
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 2264bp; fragments: 1609; full length: 0 (>=2037.6bp)

After TEtrimmer 2264 bp

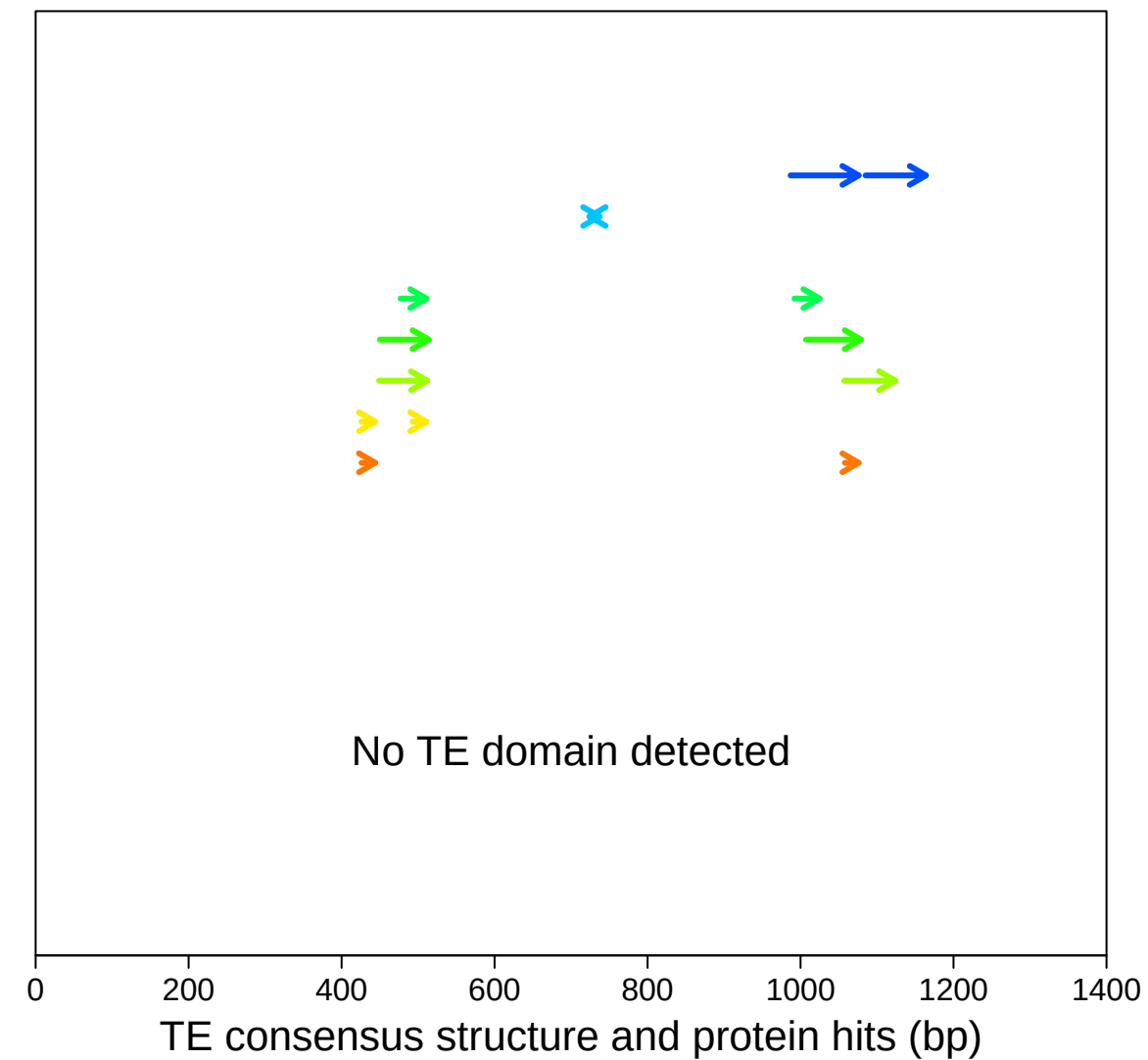
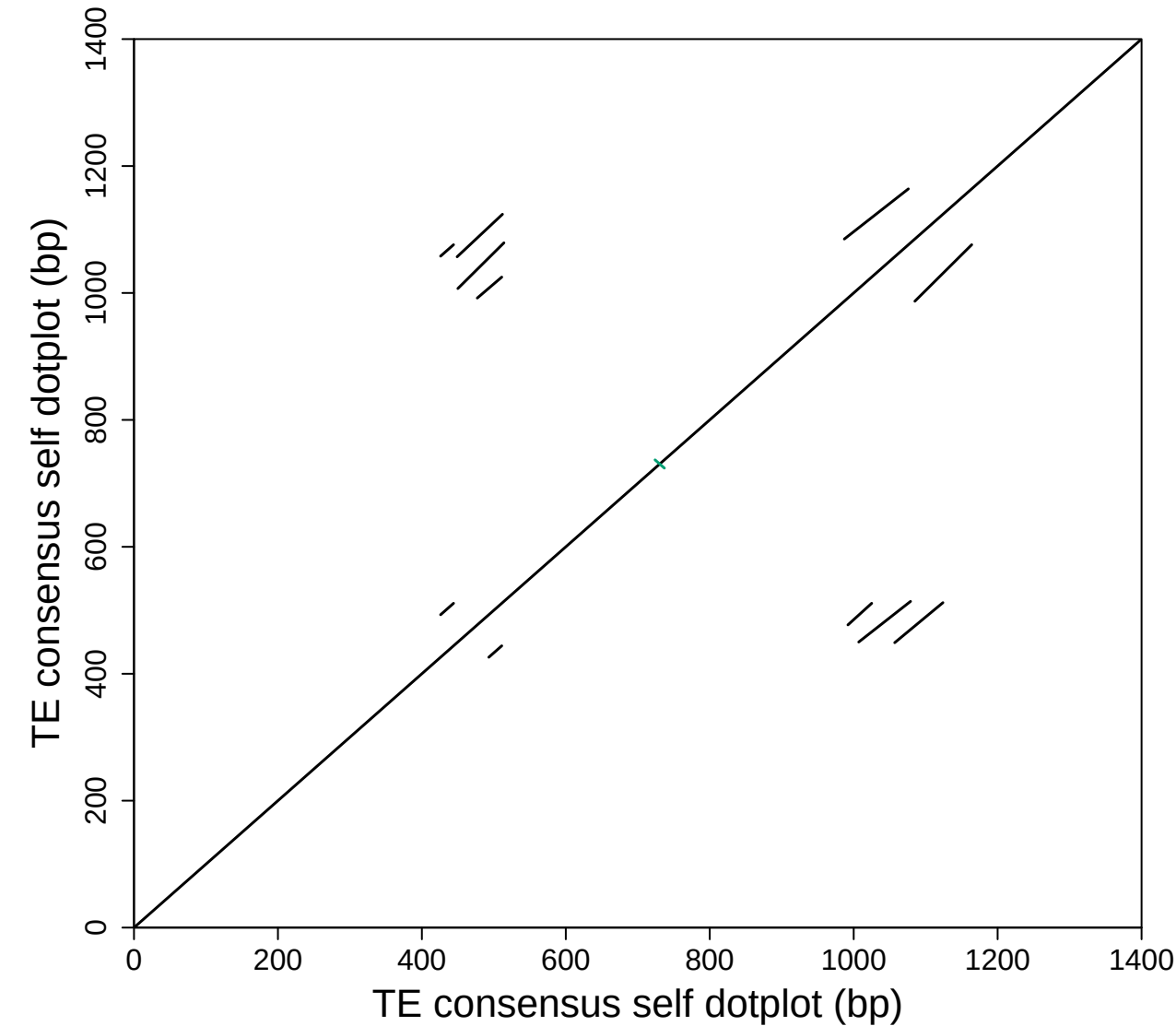
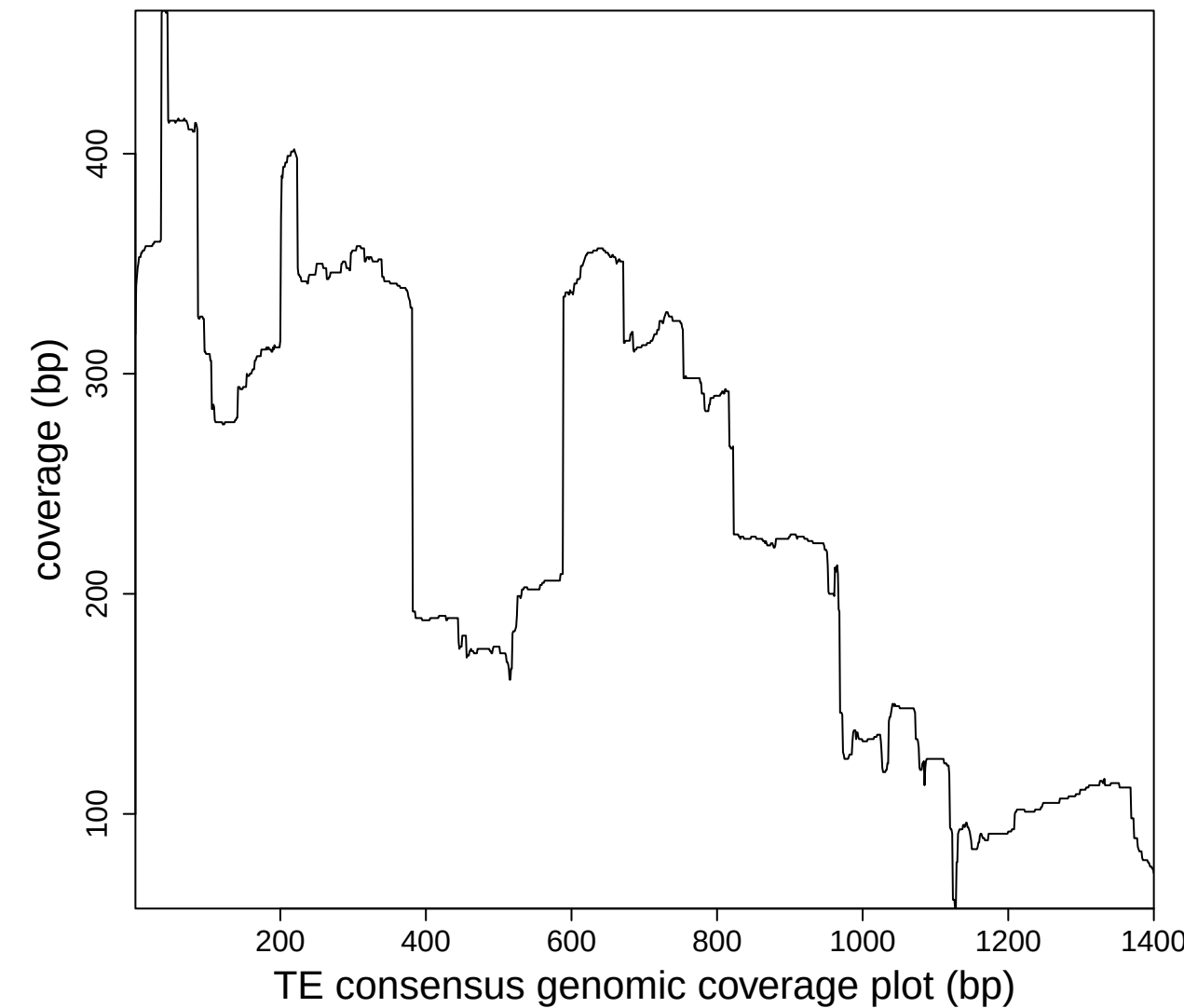
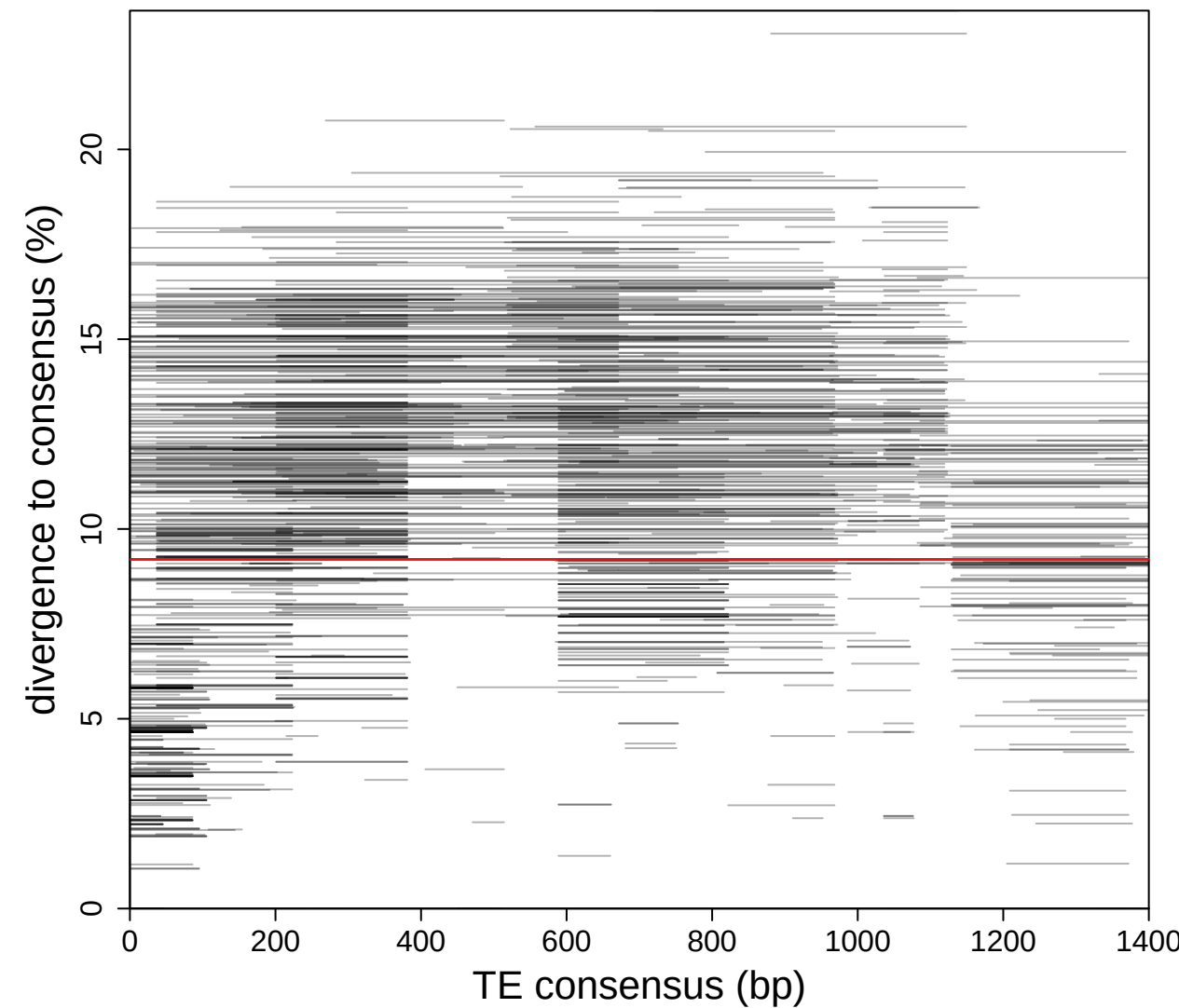


1.fasta.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 2721bp; fragments: 1079; full length: 0 (>=2448.9bp)

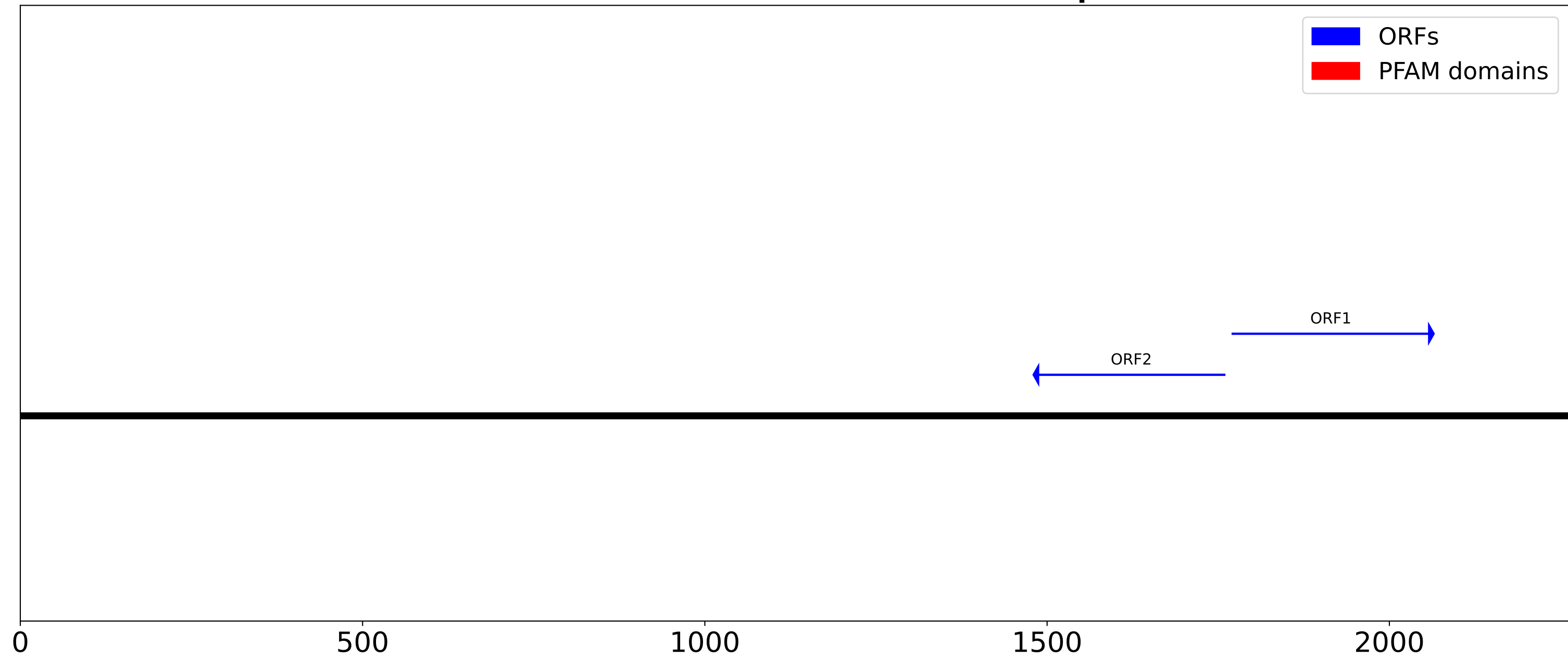
After TEtrimmer Extended plot Blue lines are boundaries



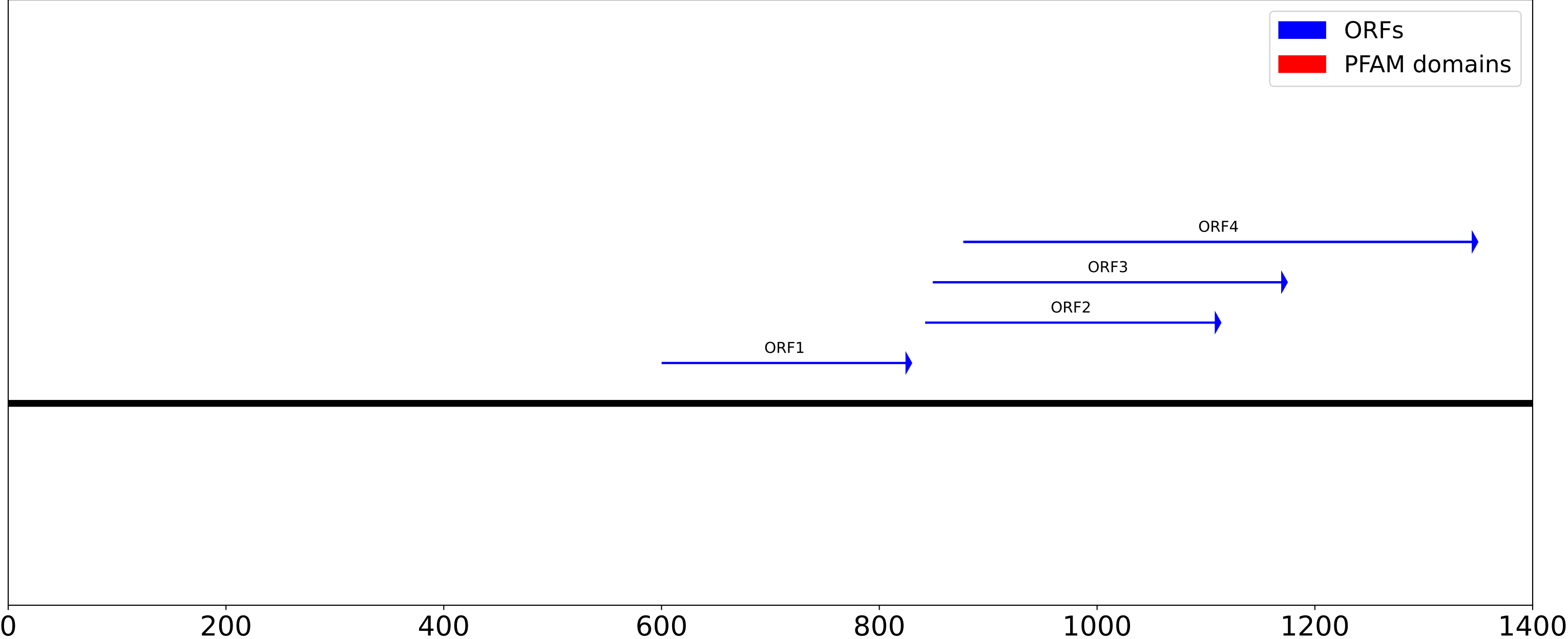
TE: rnd_5_family_1431
size: 1400bp; fragments: 1115; full length: 1 (>=1260bp)



After TEtrimmer ORF and PFAM domain plot



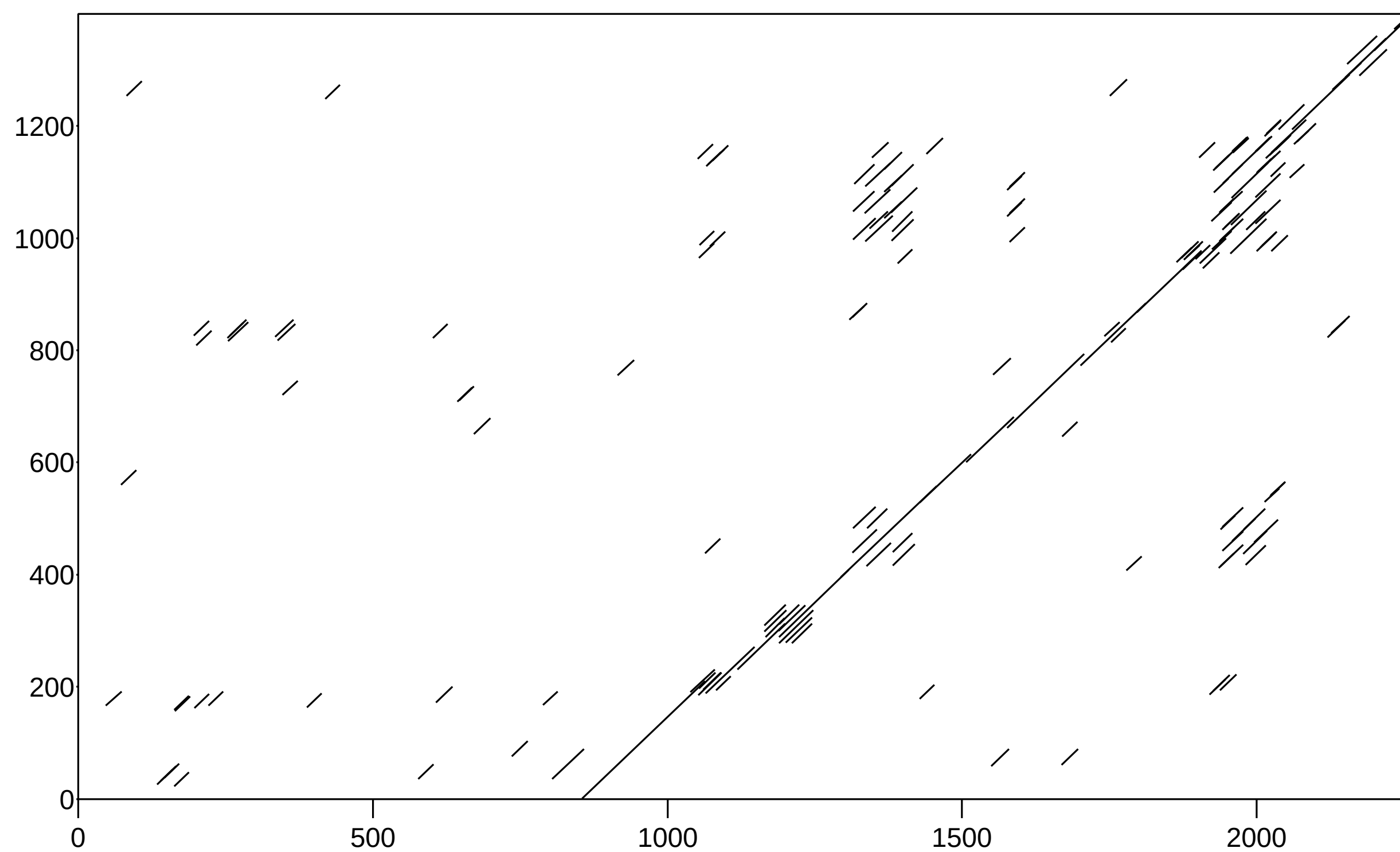
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)