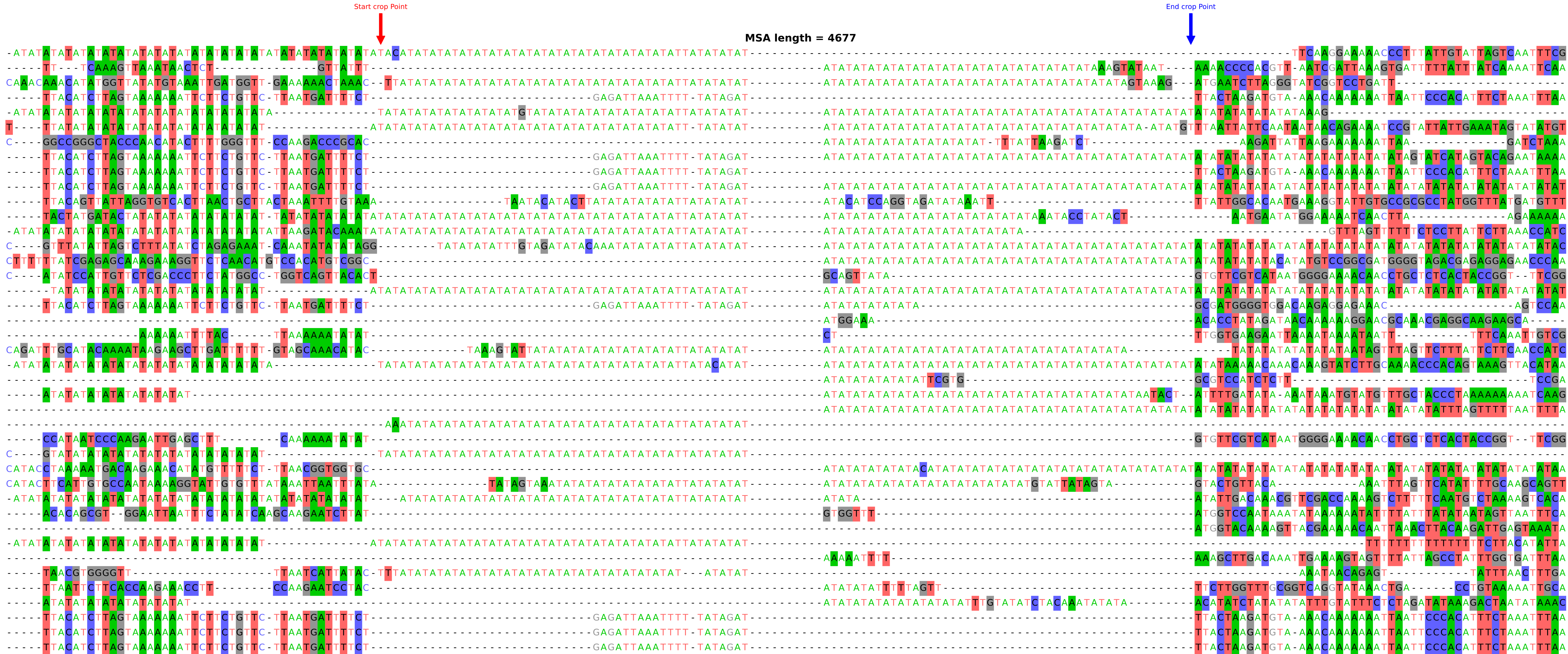
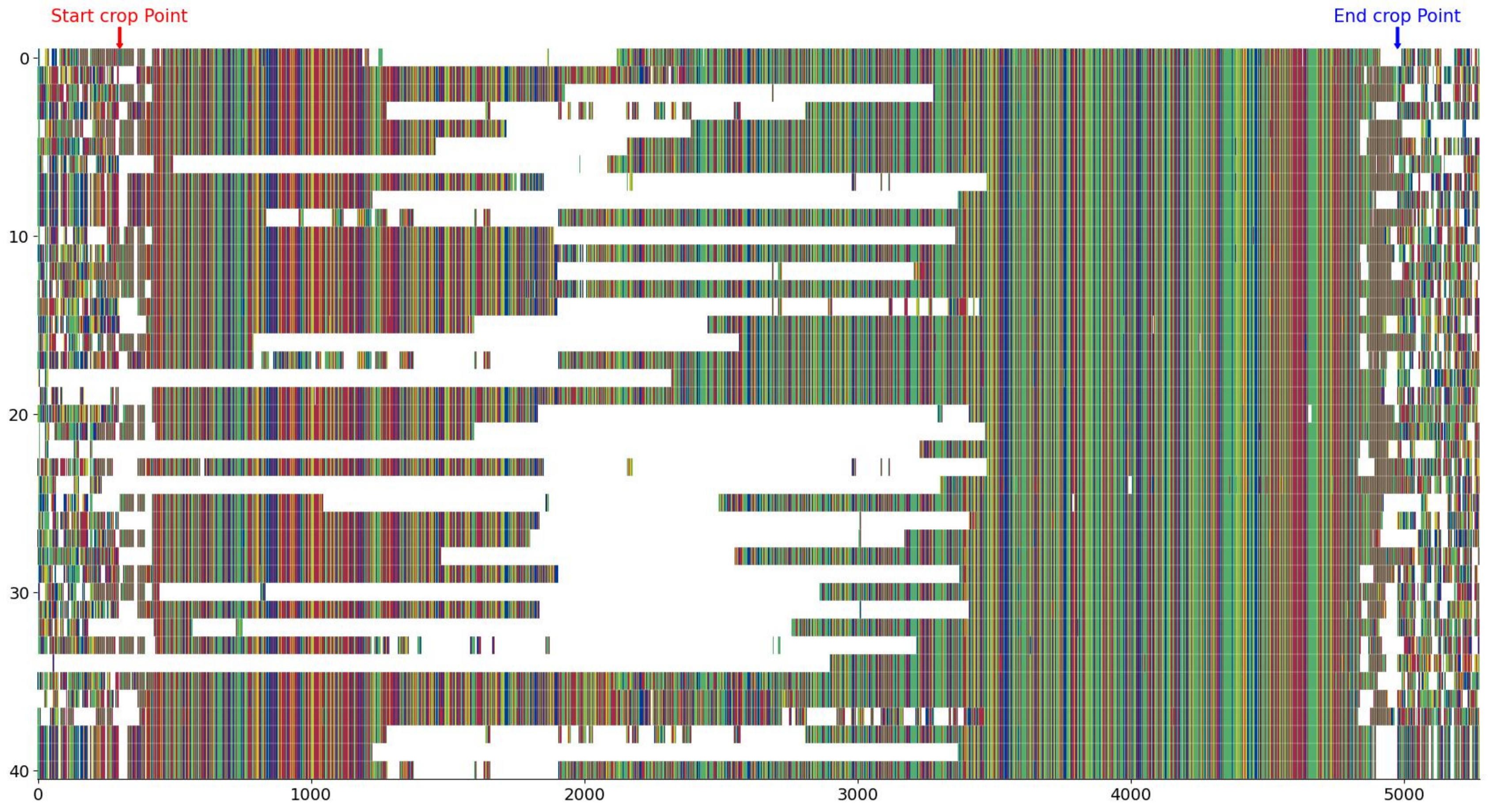
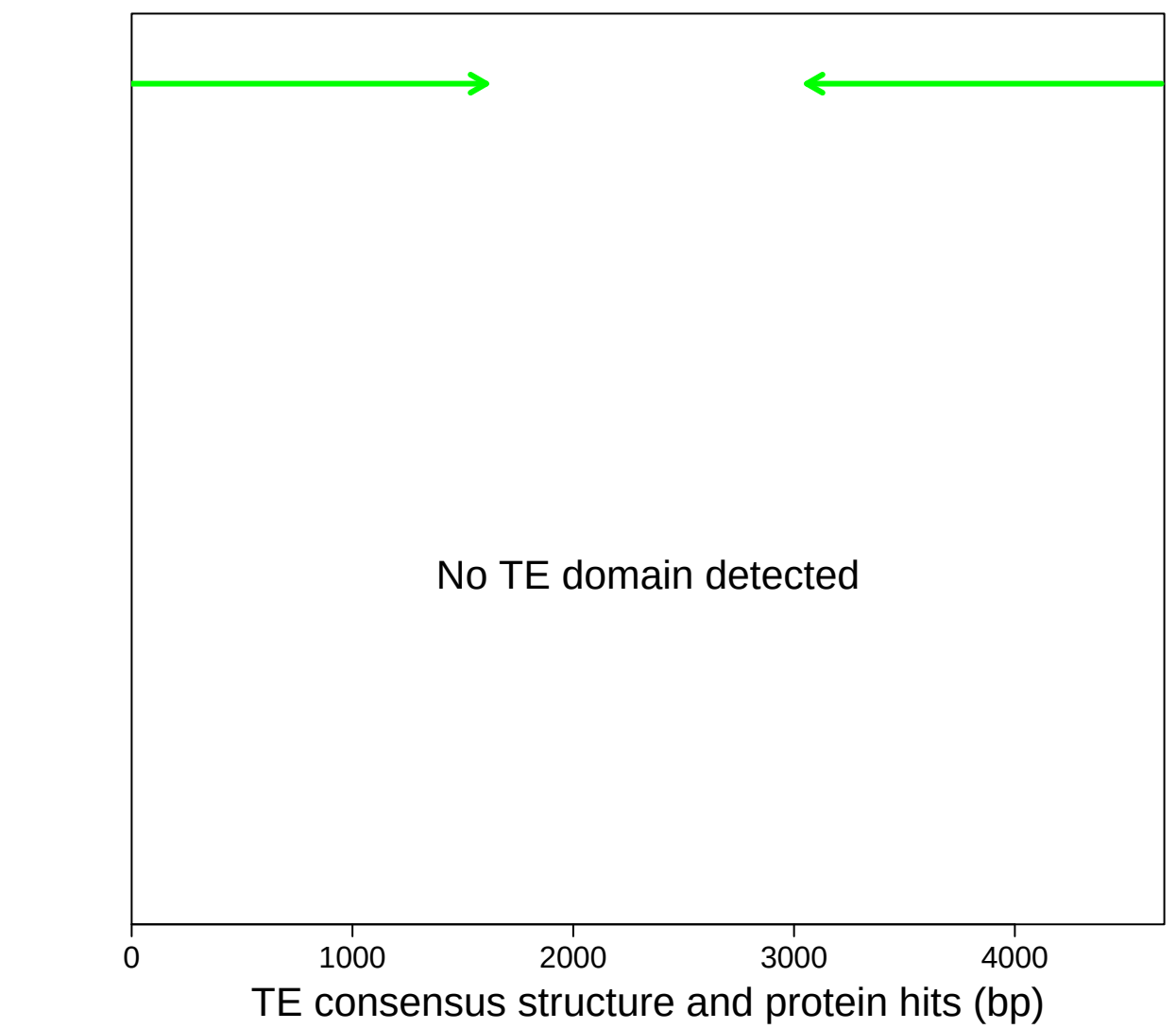
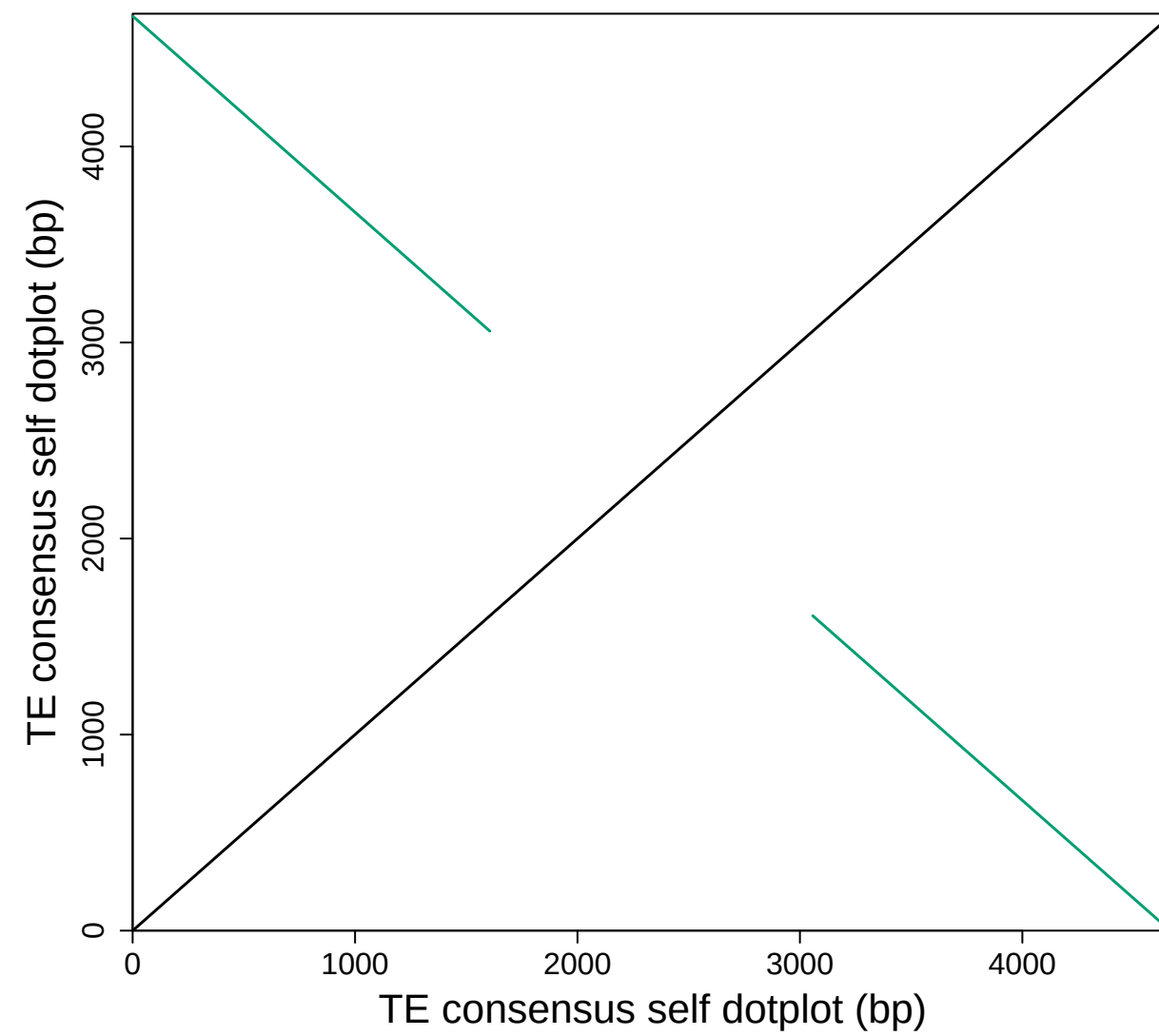
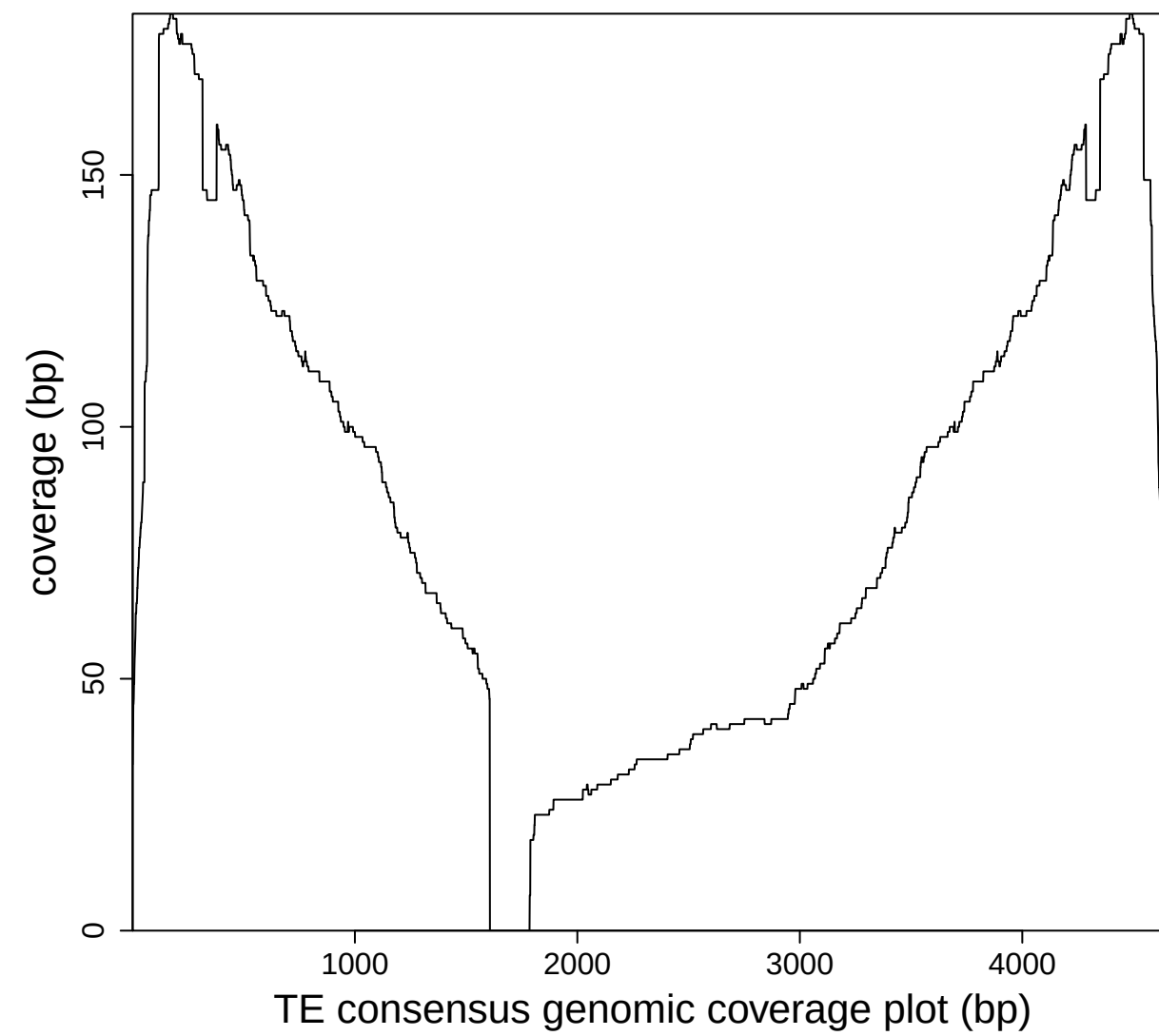
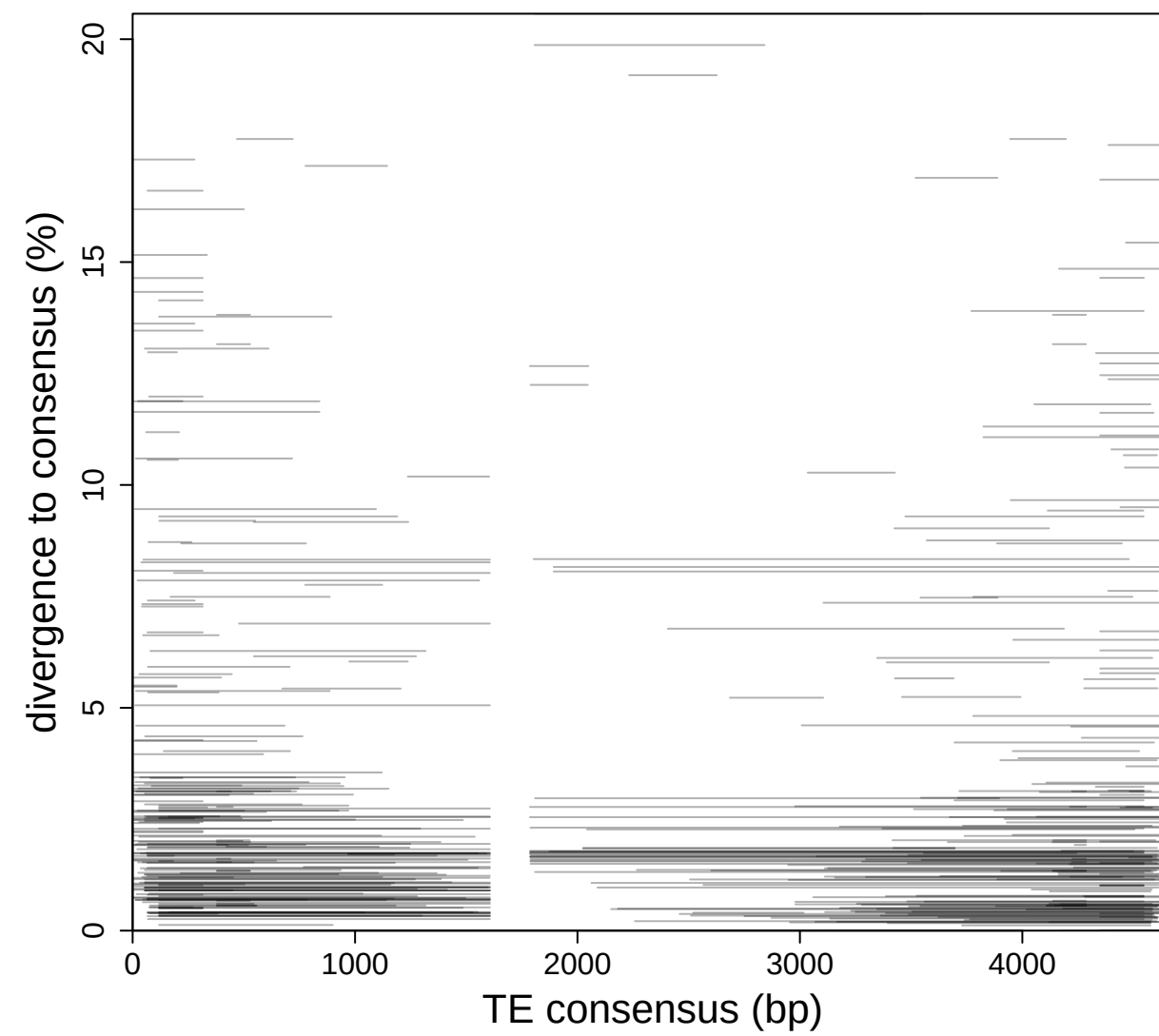






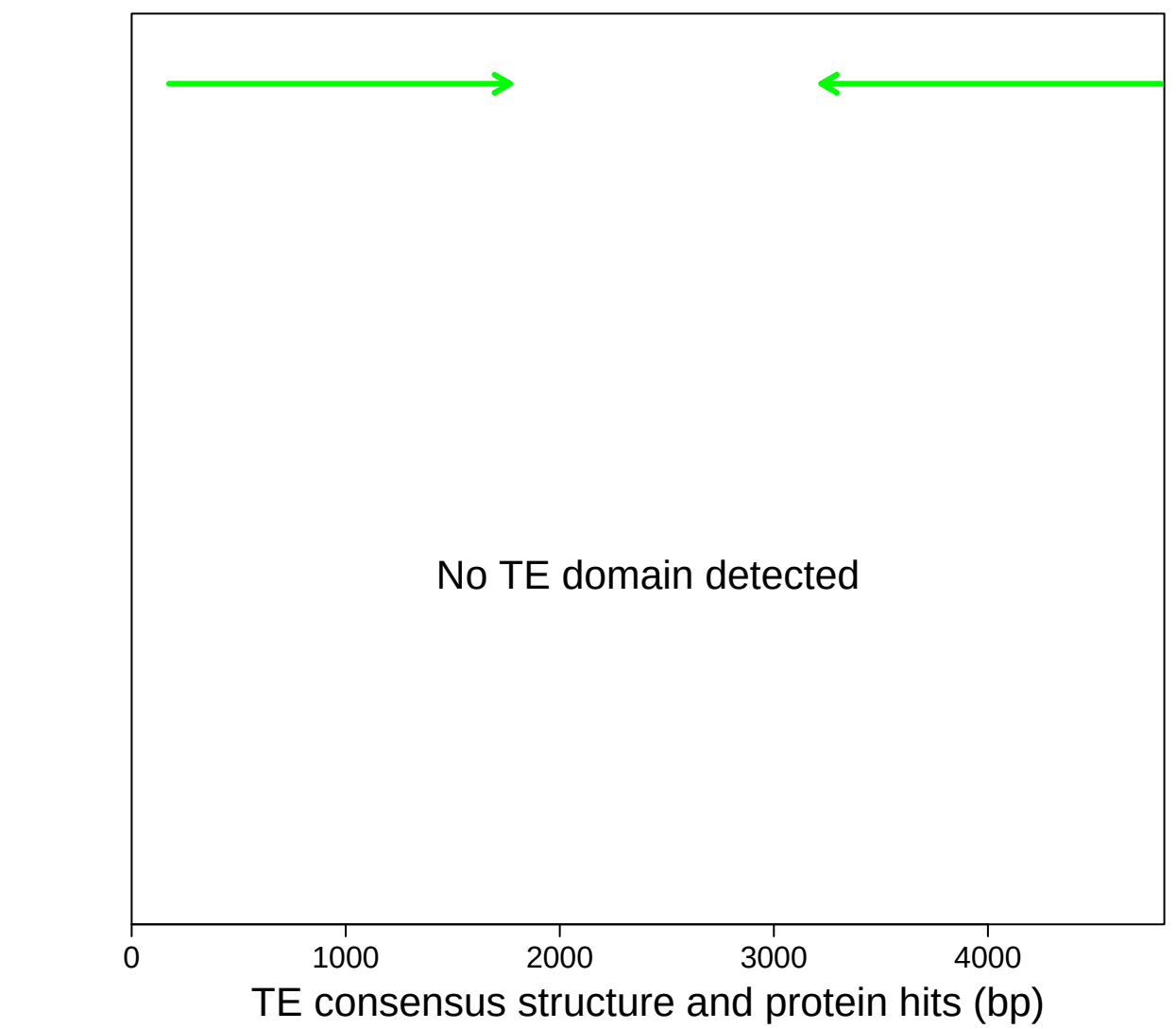
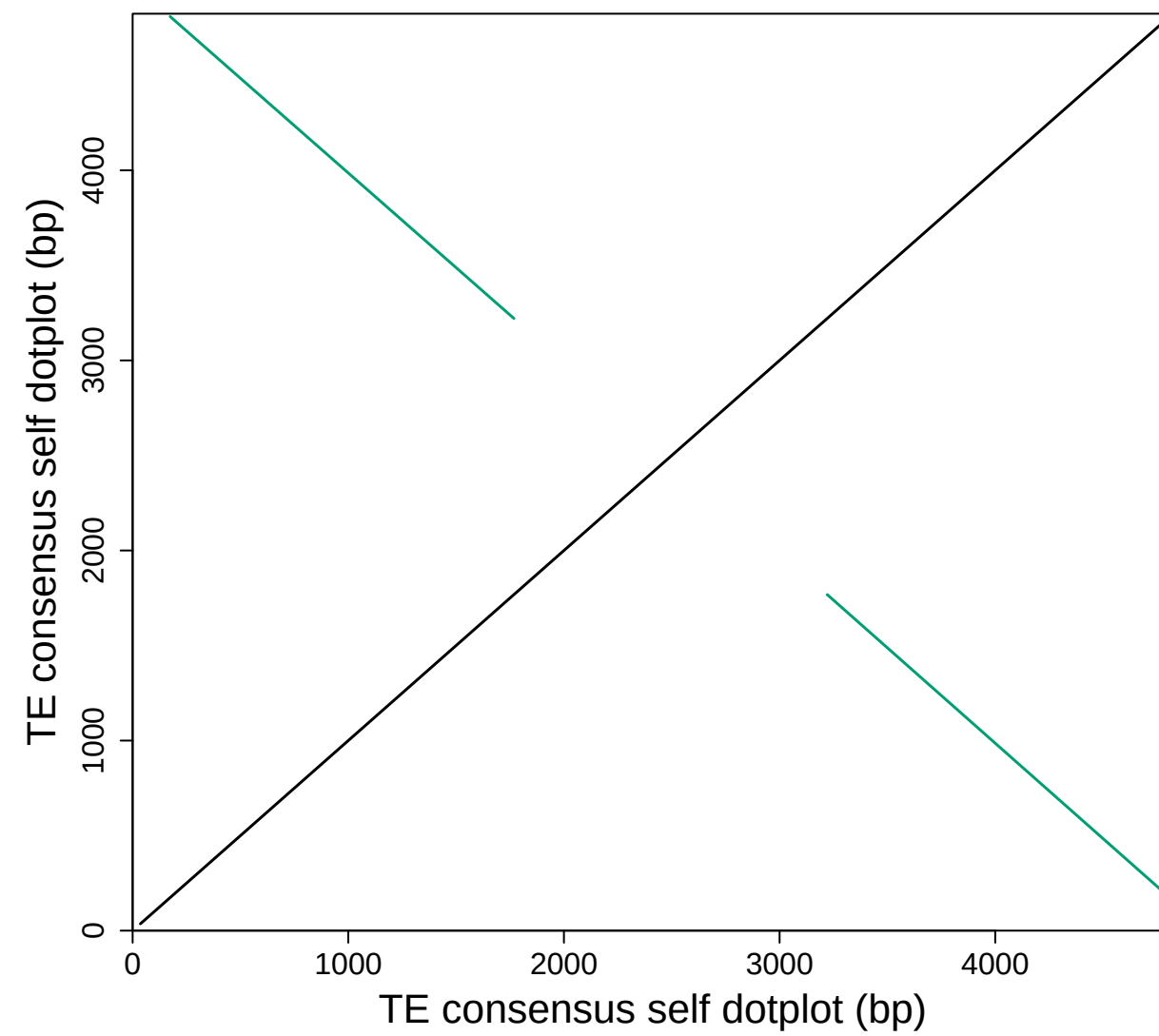
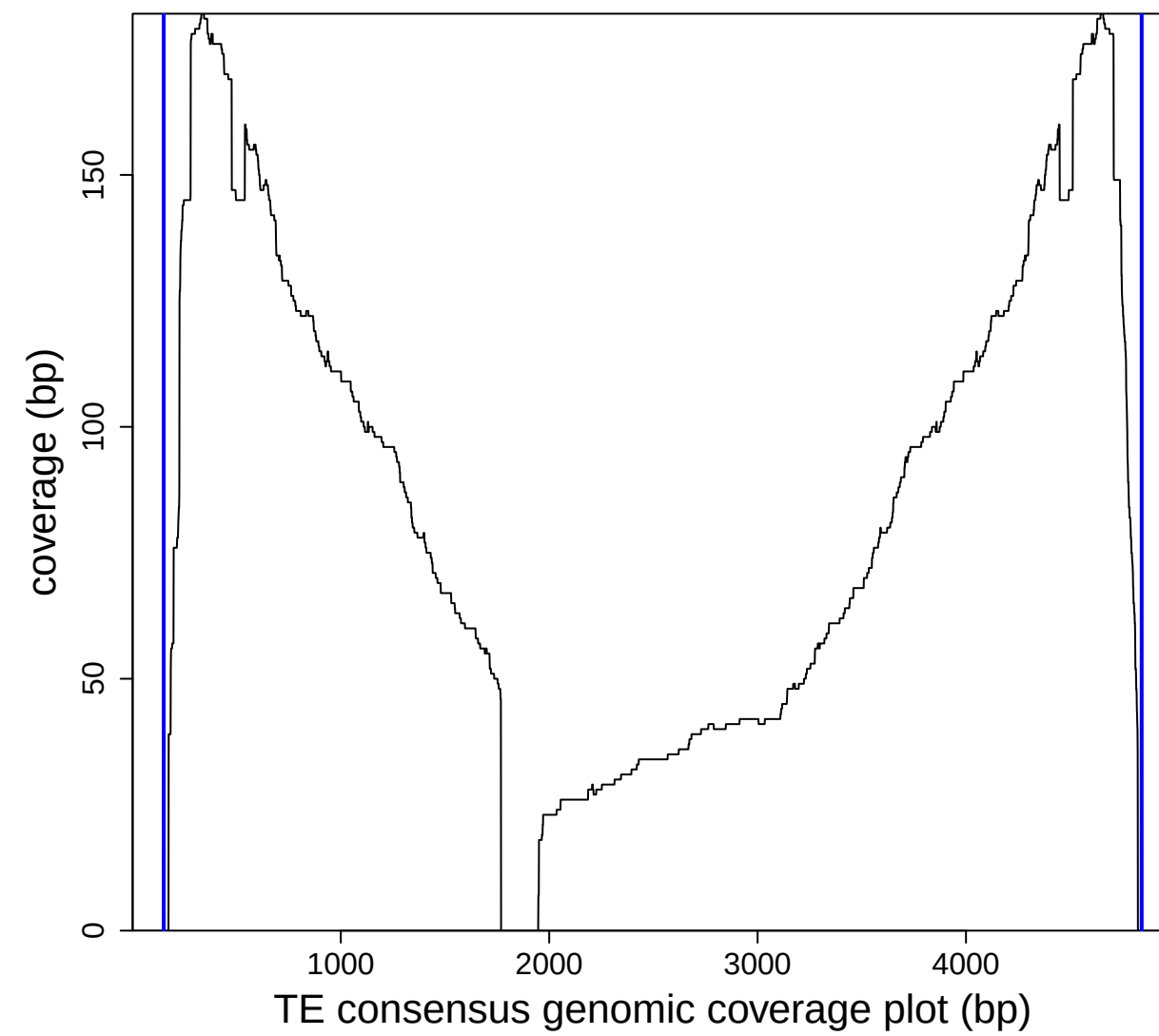
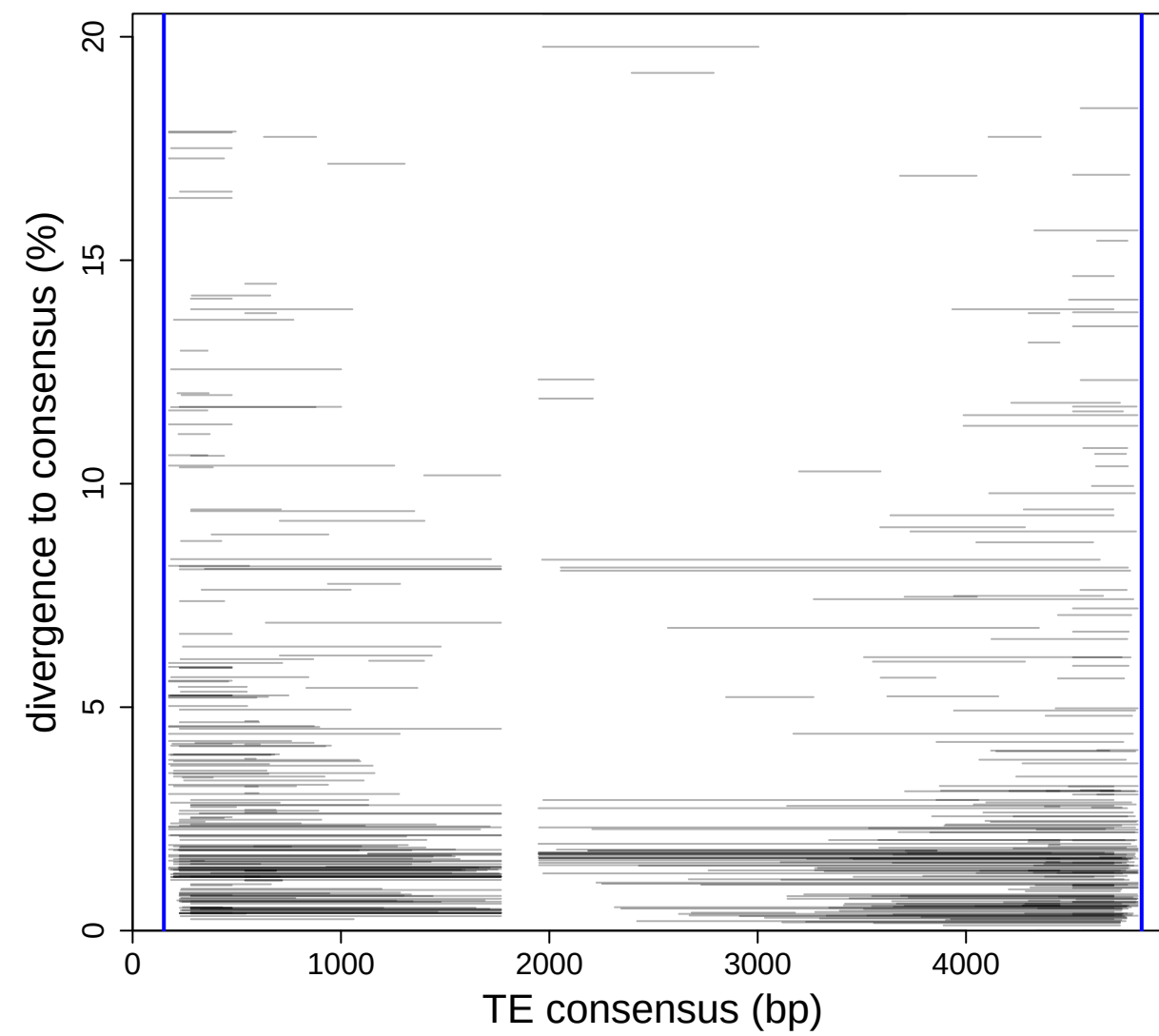
bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa ce
size: 4677bp; fragments: 435; full length: 0 (>=4209.3bp)

After TETrimmer 4677 bp

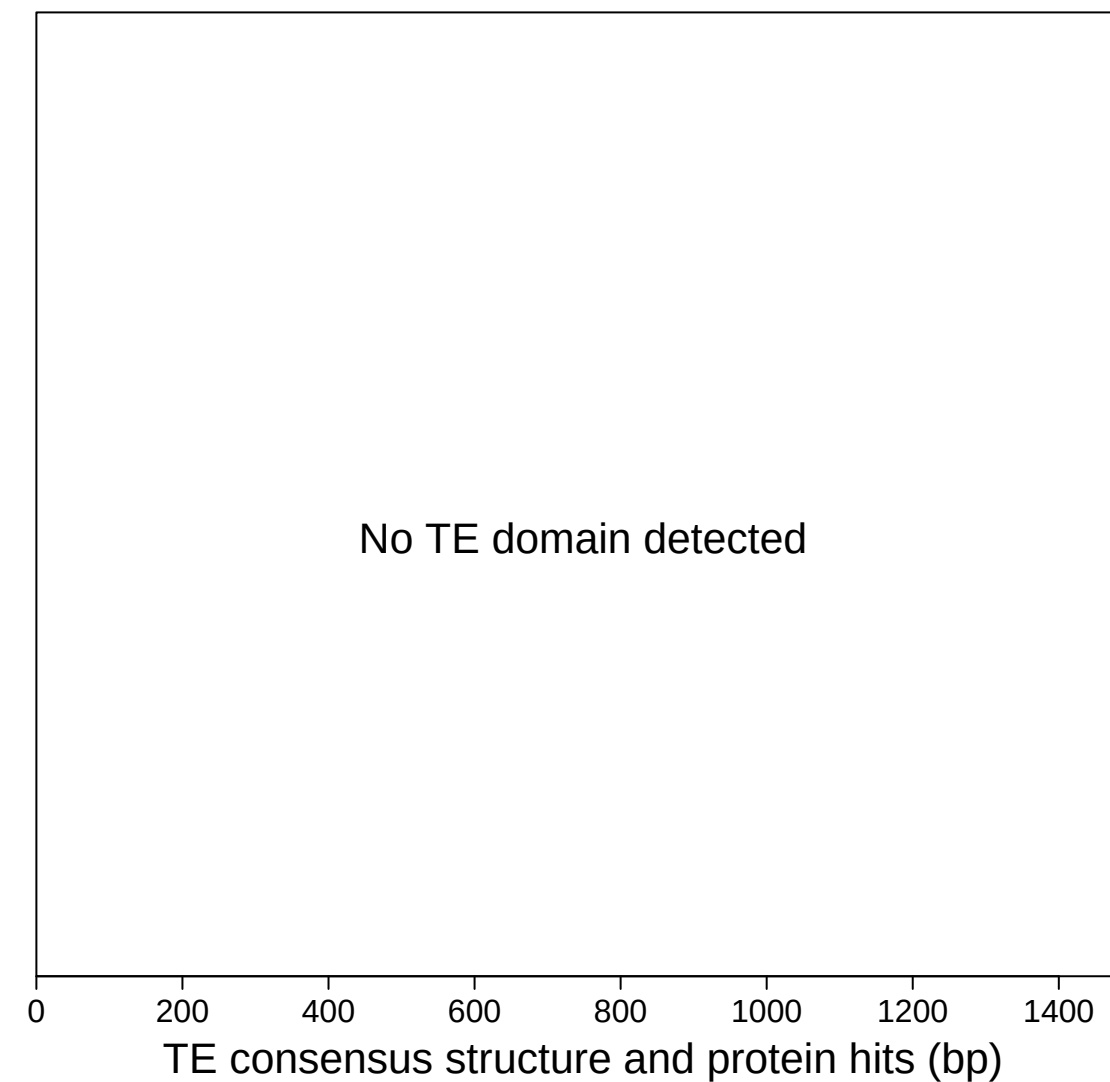
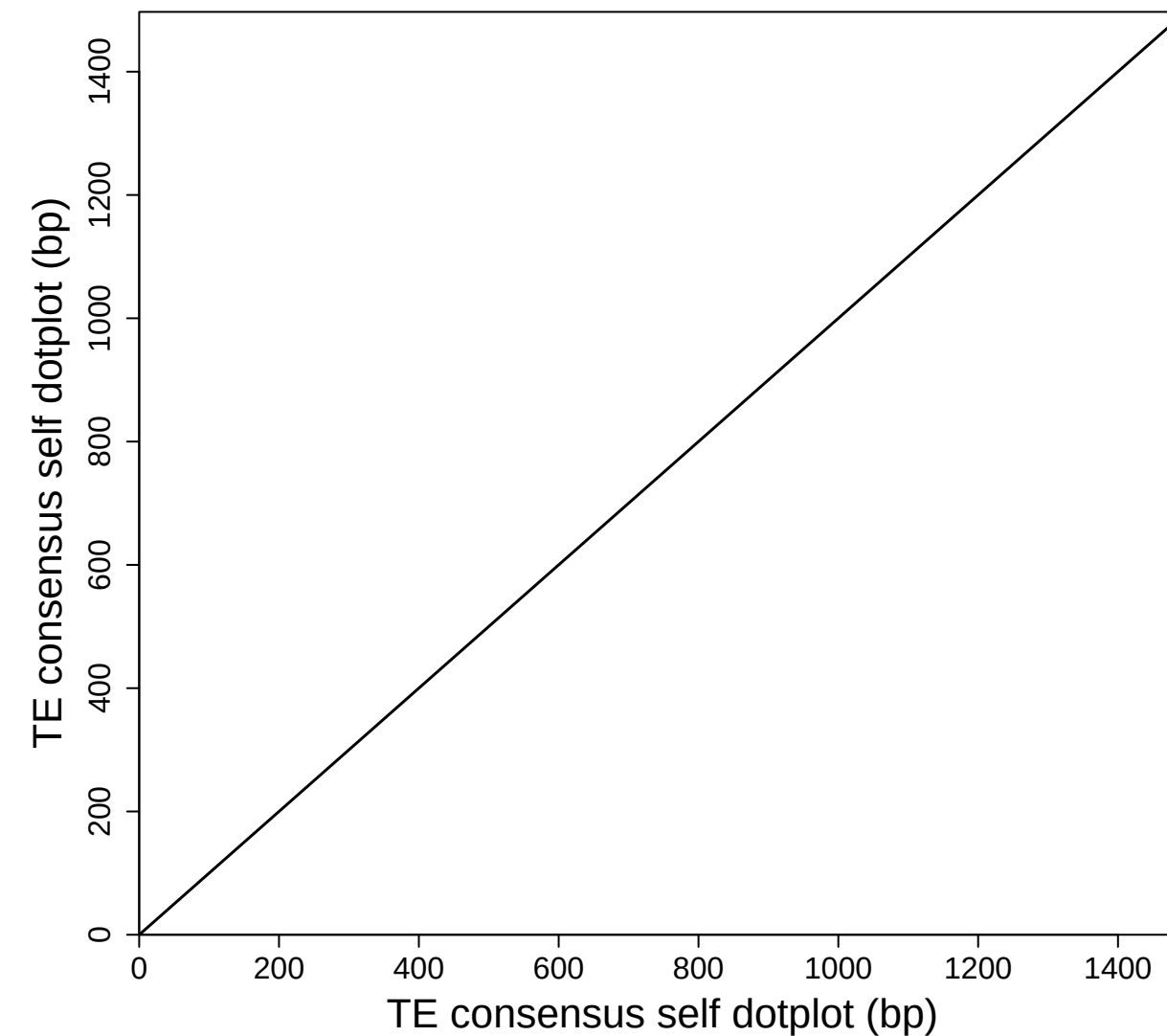
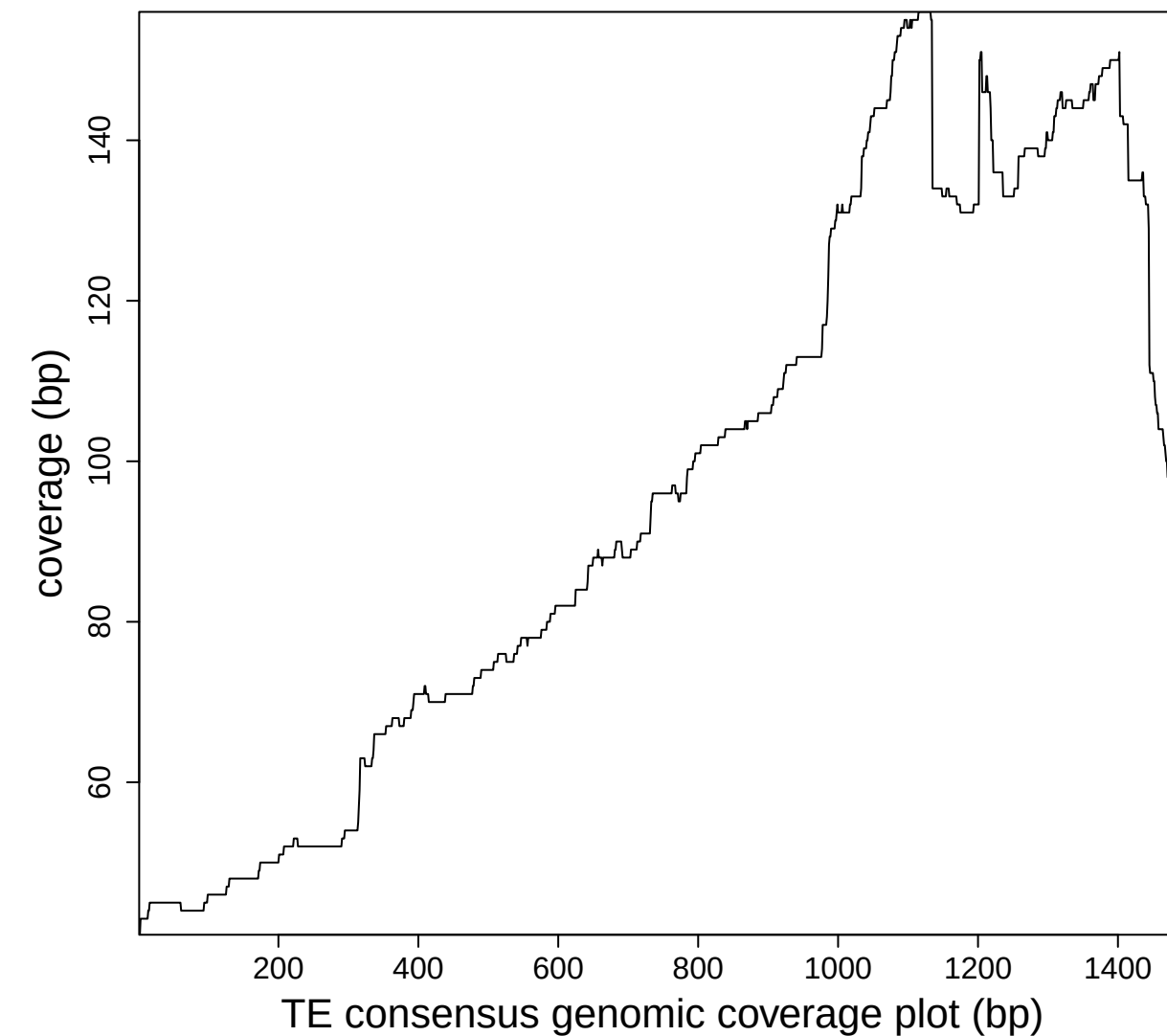
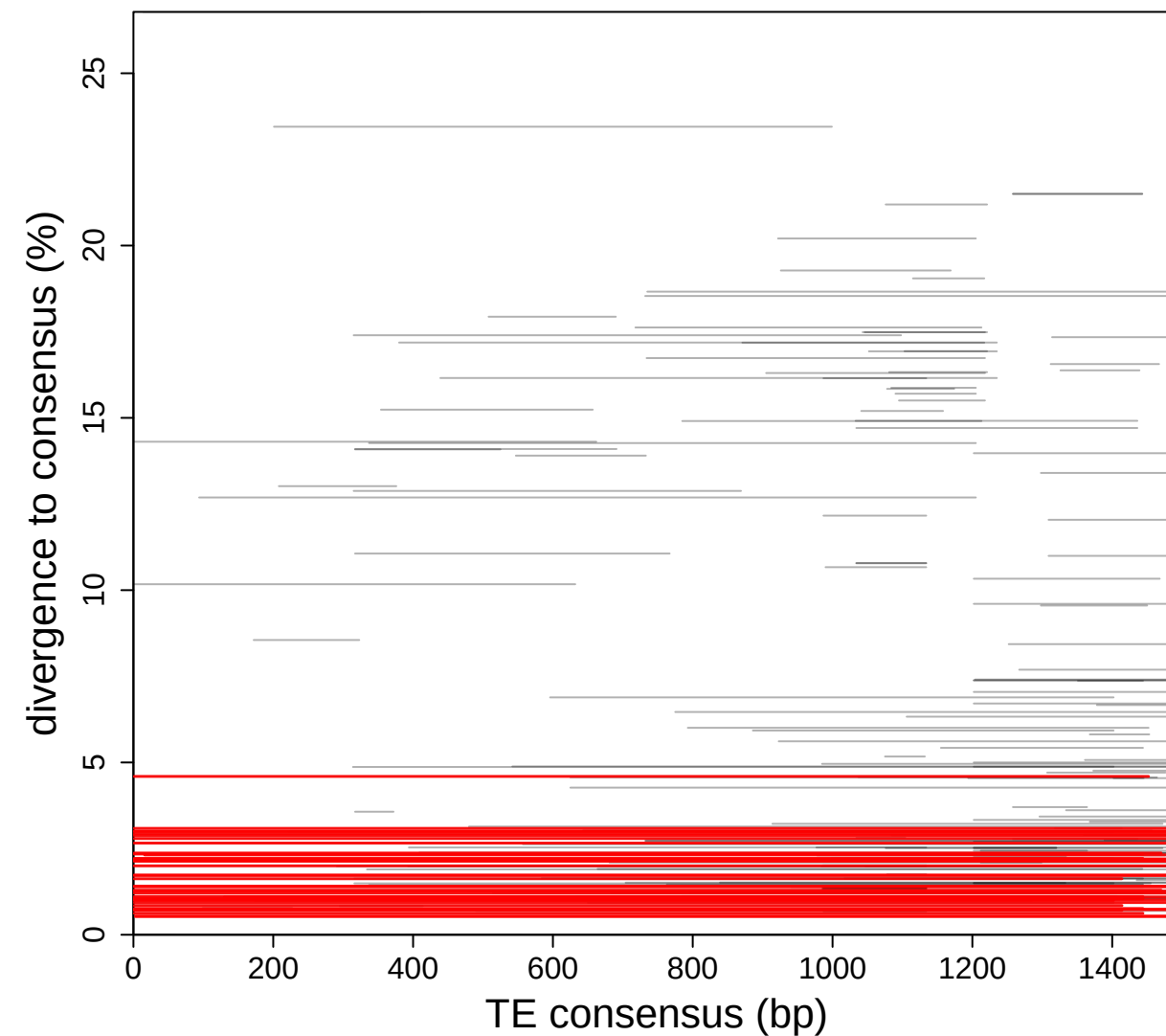


ily_874.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 4993bp; fragments: 435; full length: 0 (>=4493.7bp)

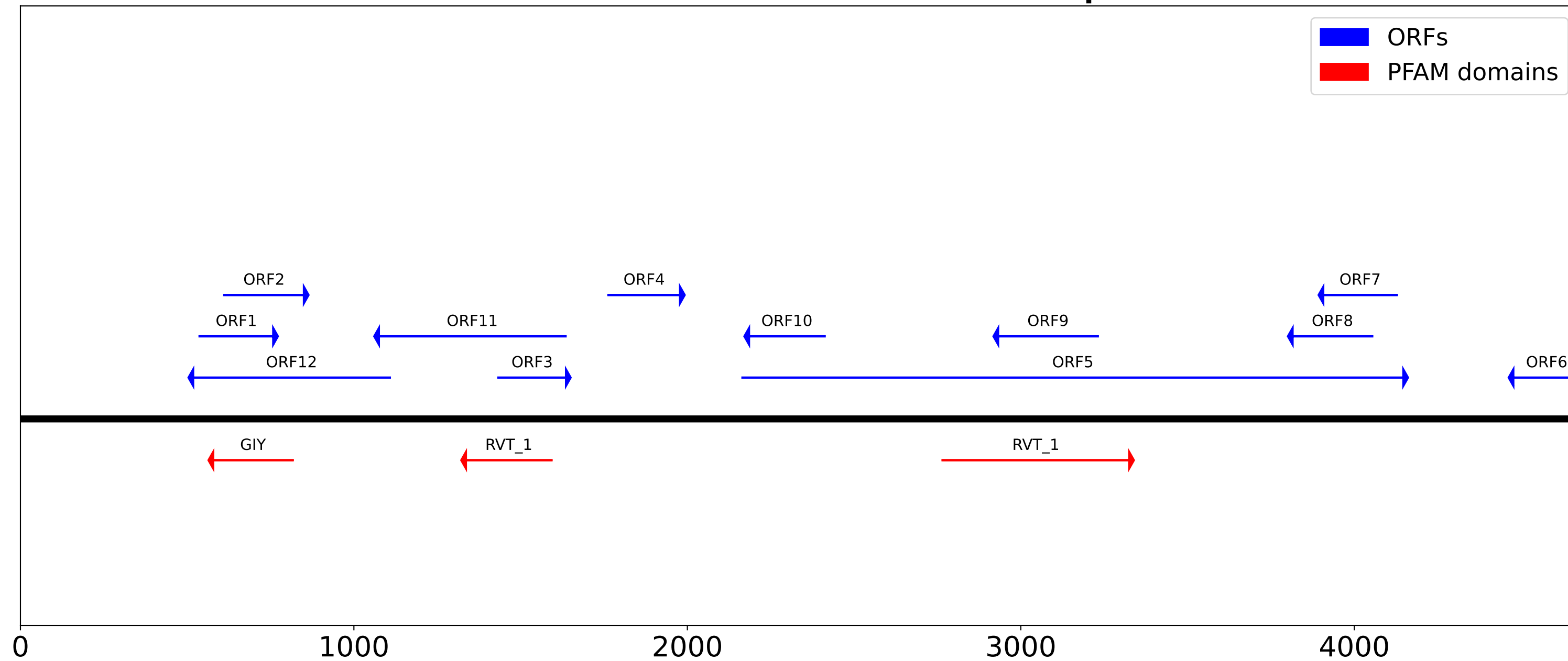
After TEtrimmer Extended plot Blue lines are boundaries



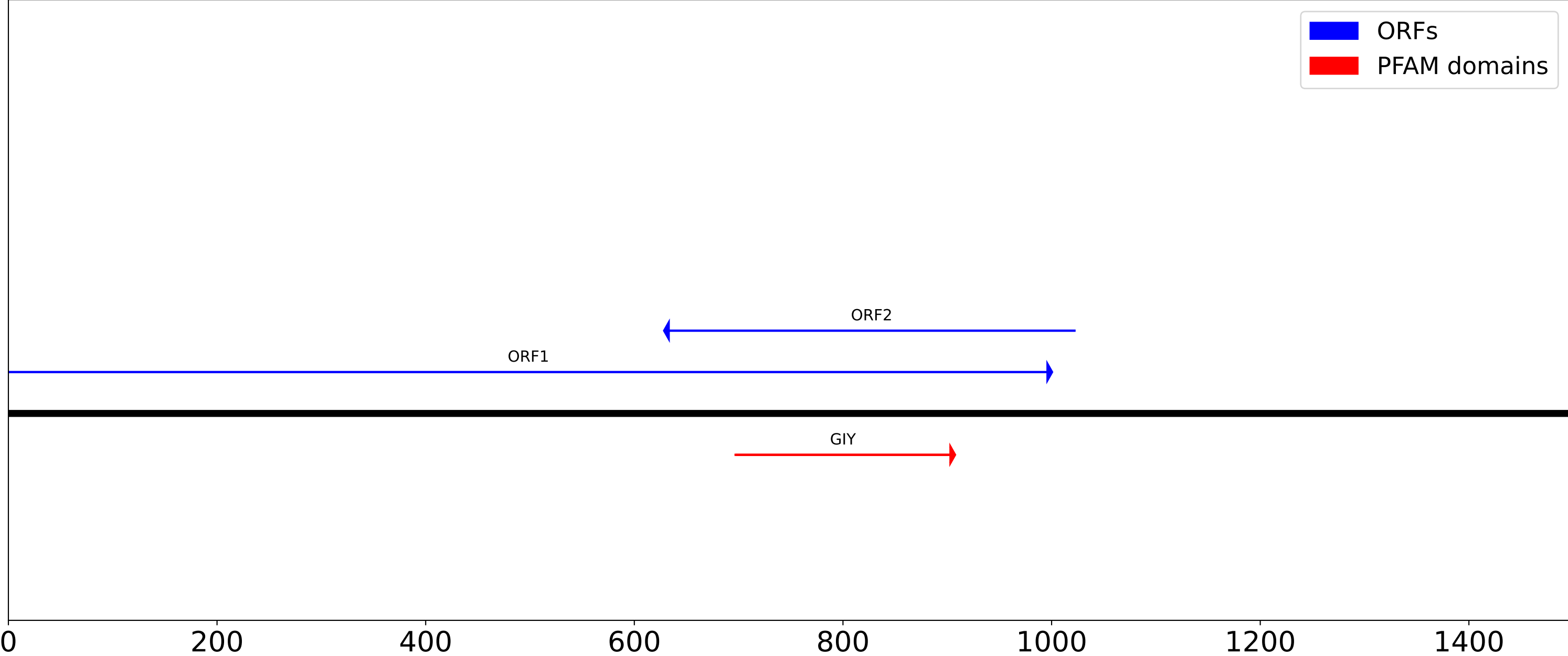
TE: rnd_1_family_874
size: 1497bp; fragments: 228; full length: 42 (≥ 1347.3 bp)



After TEtrimmer ORF and PFAM domain plot



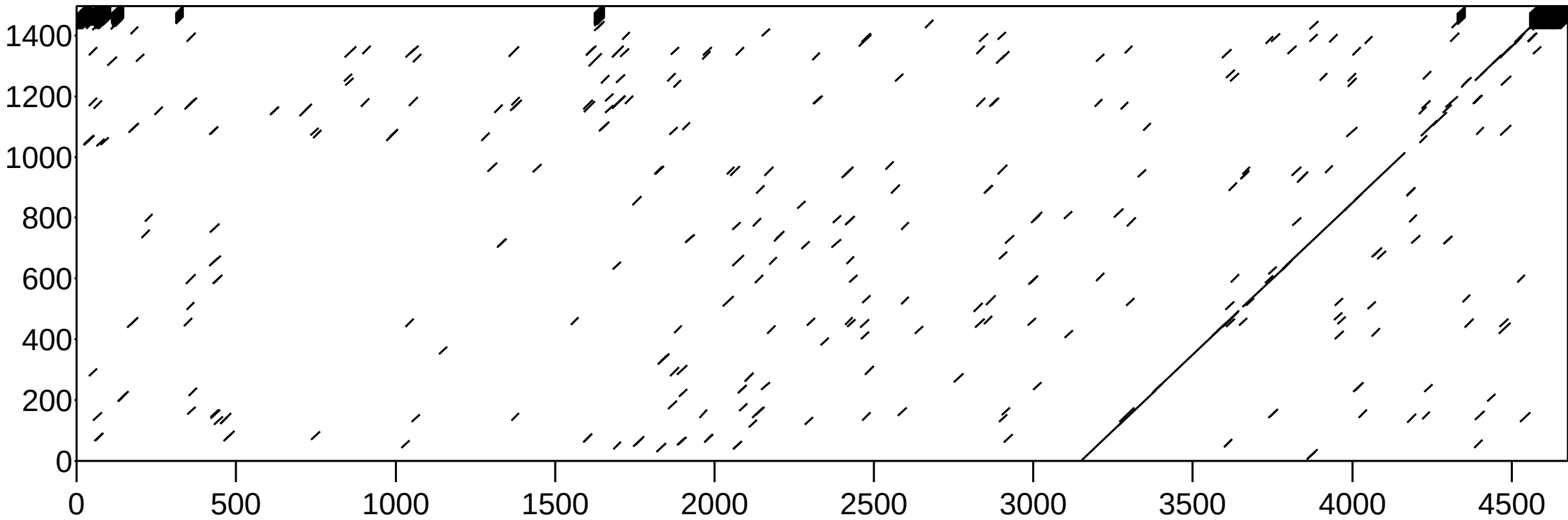
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)