

Start crop Point



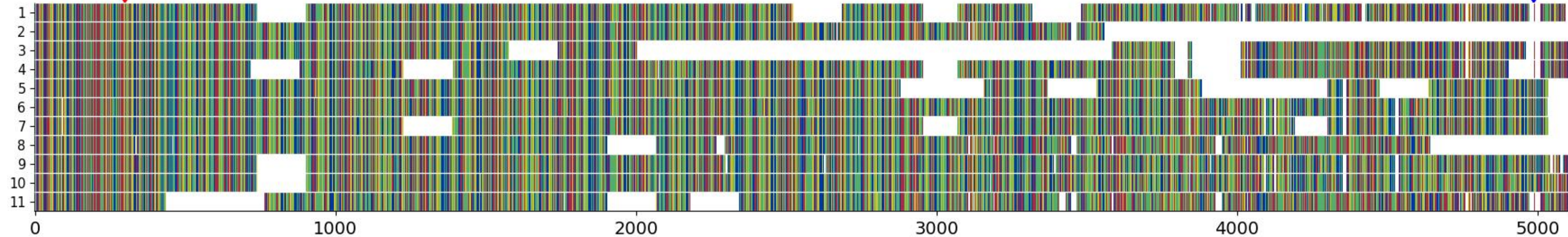
MSA length = 4688

End crop Point



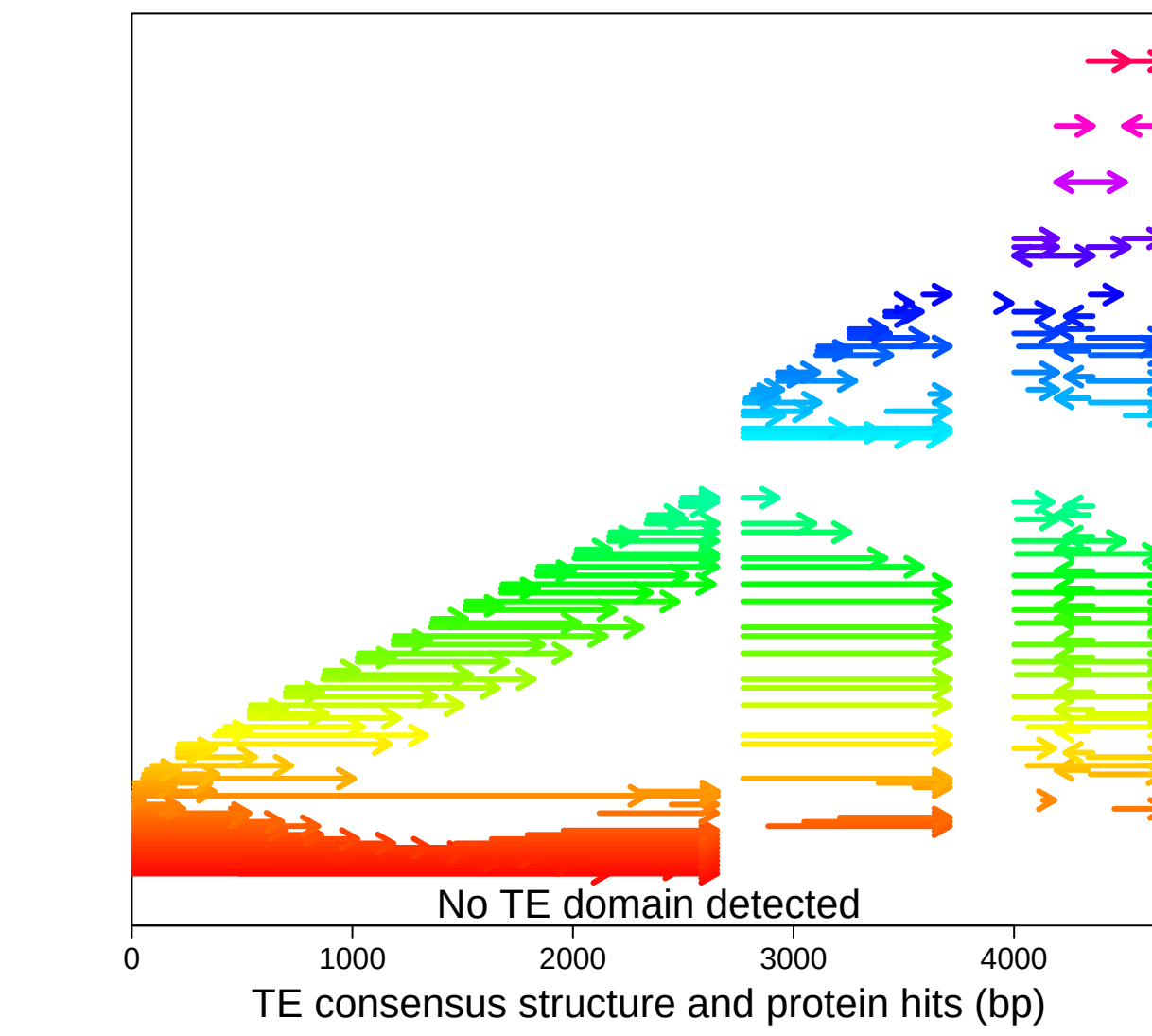
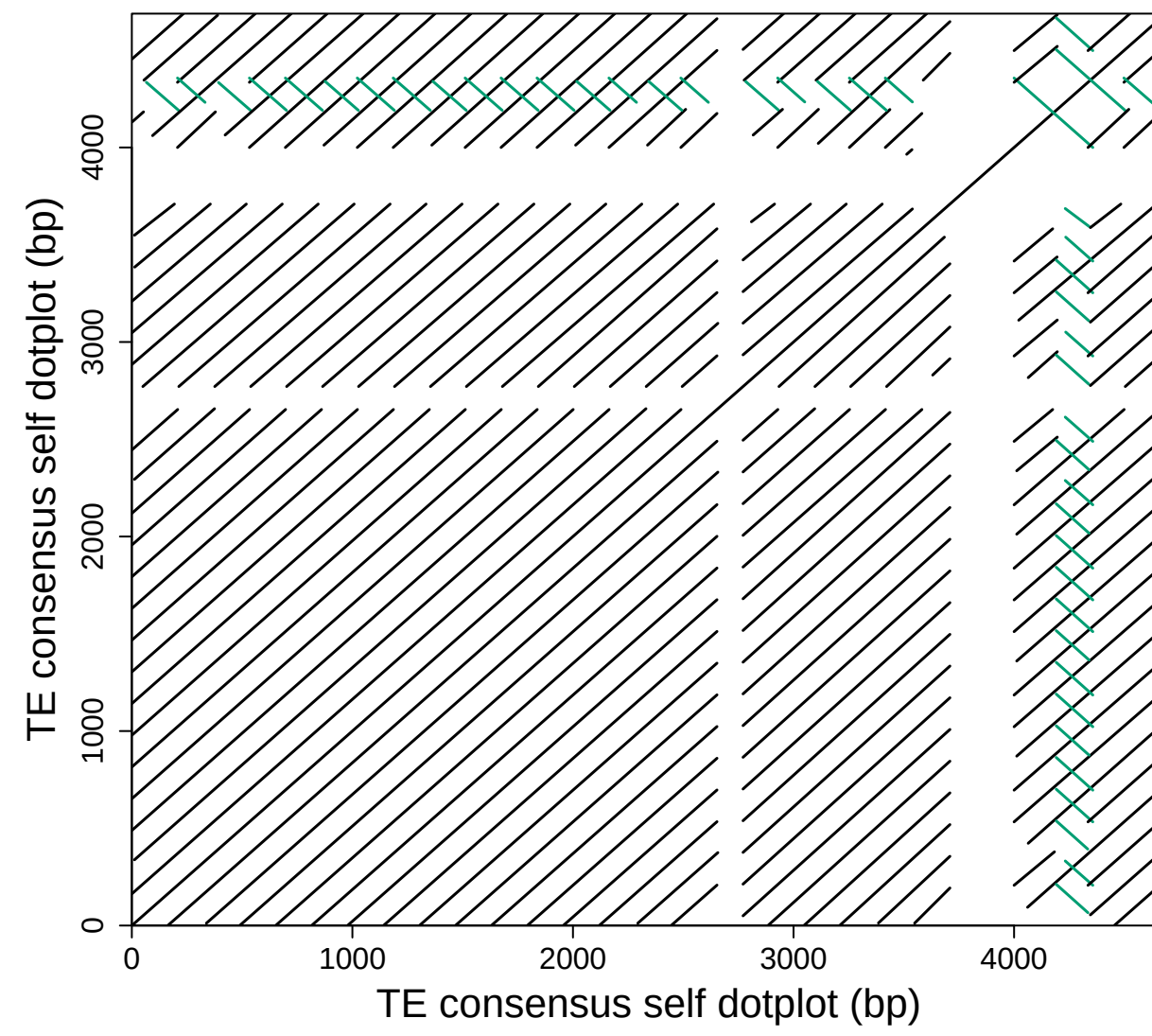
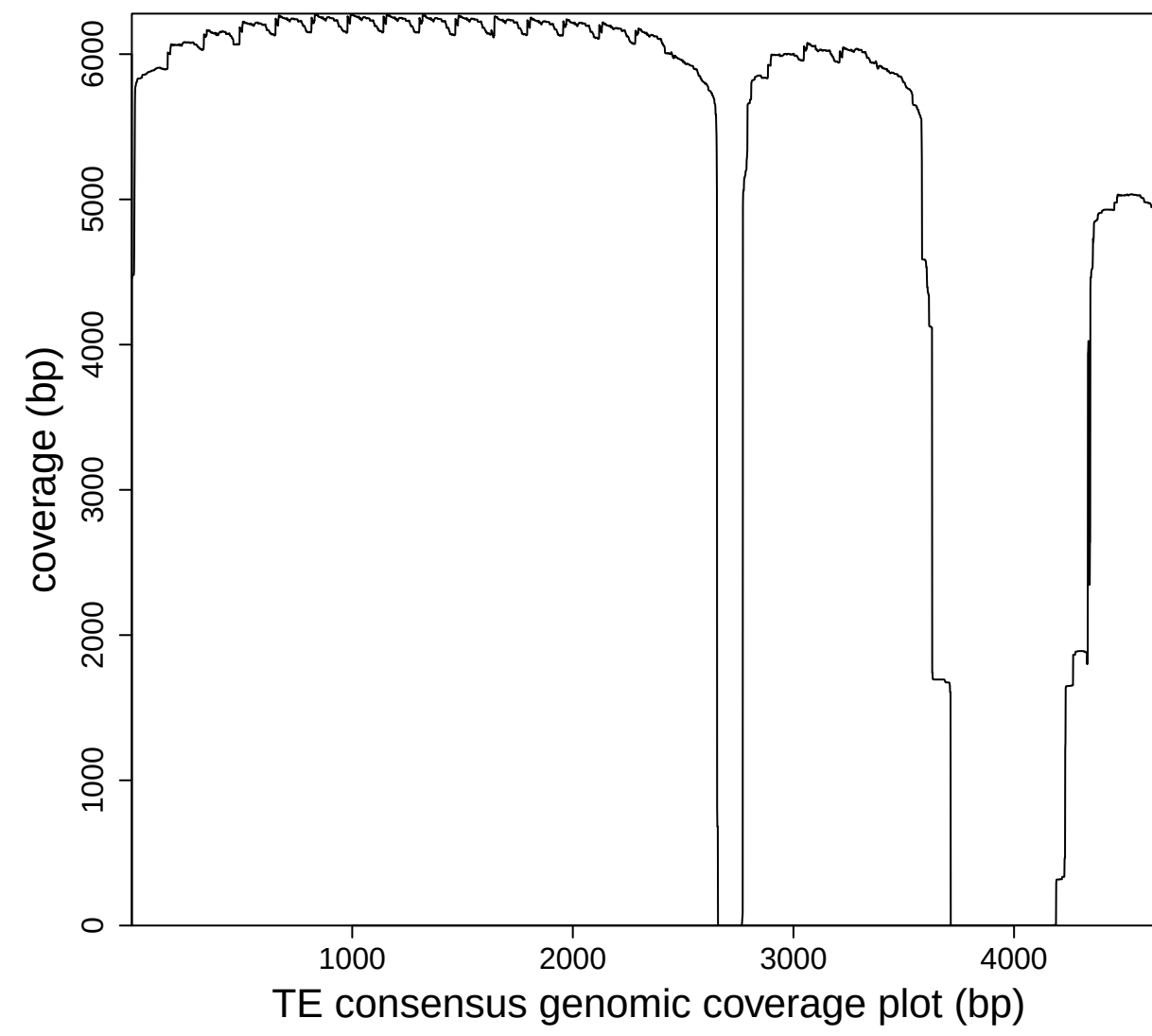
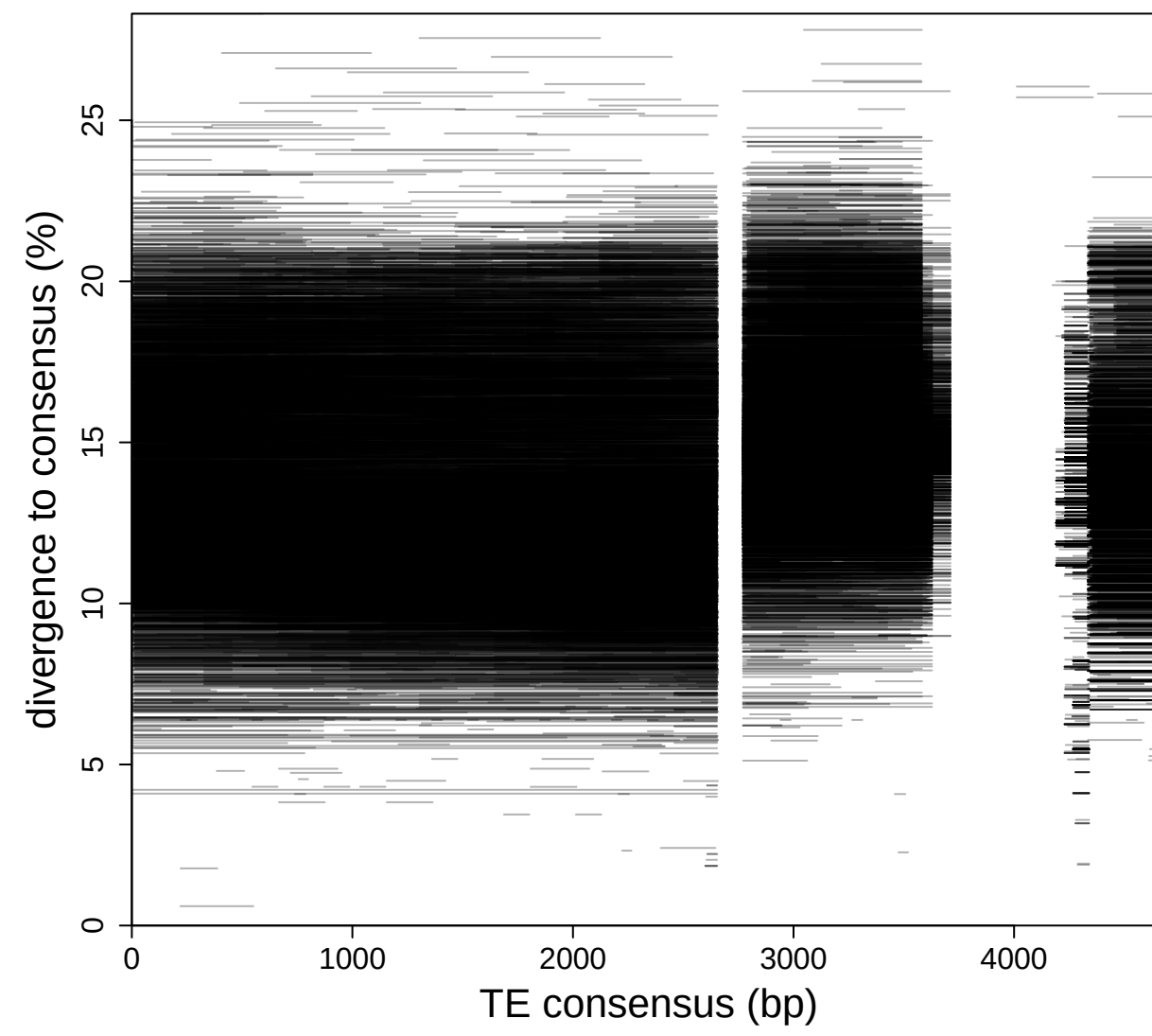
Start crop Point

End crop Point



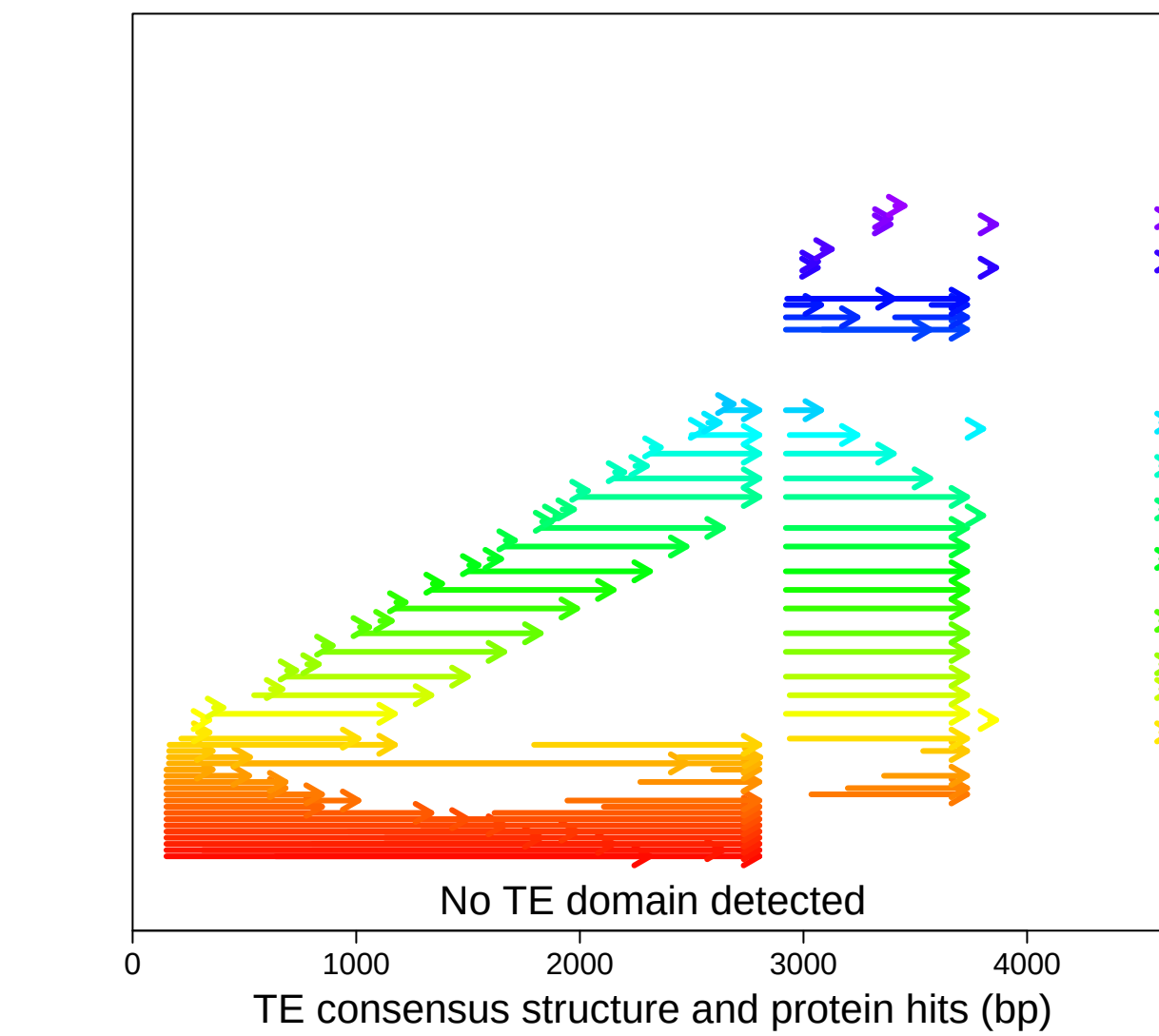
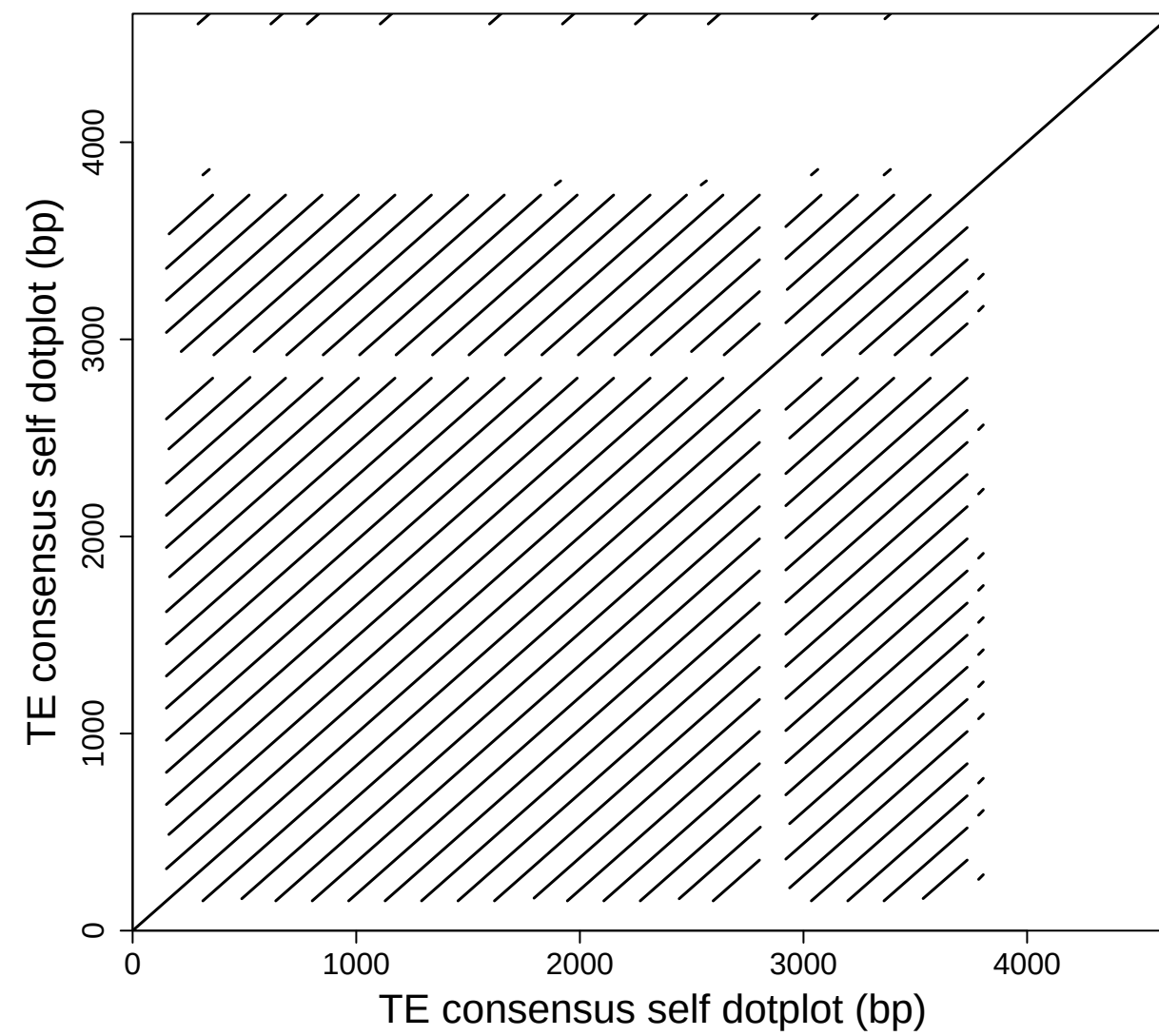
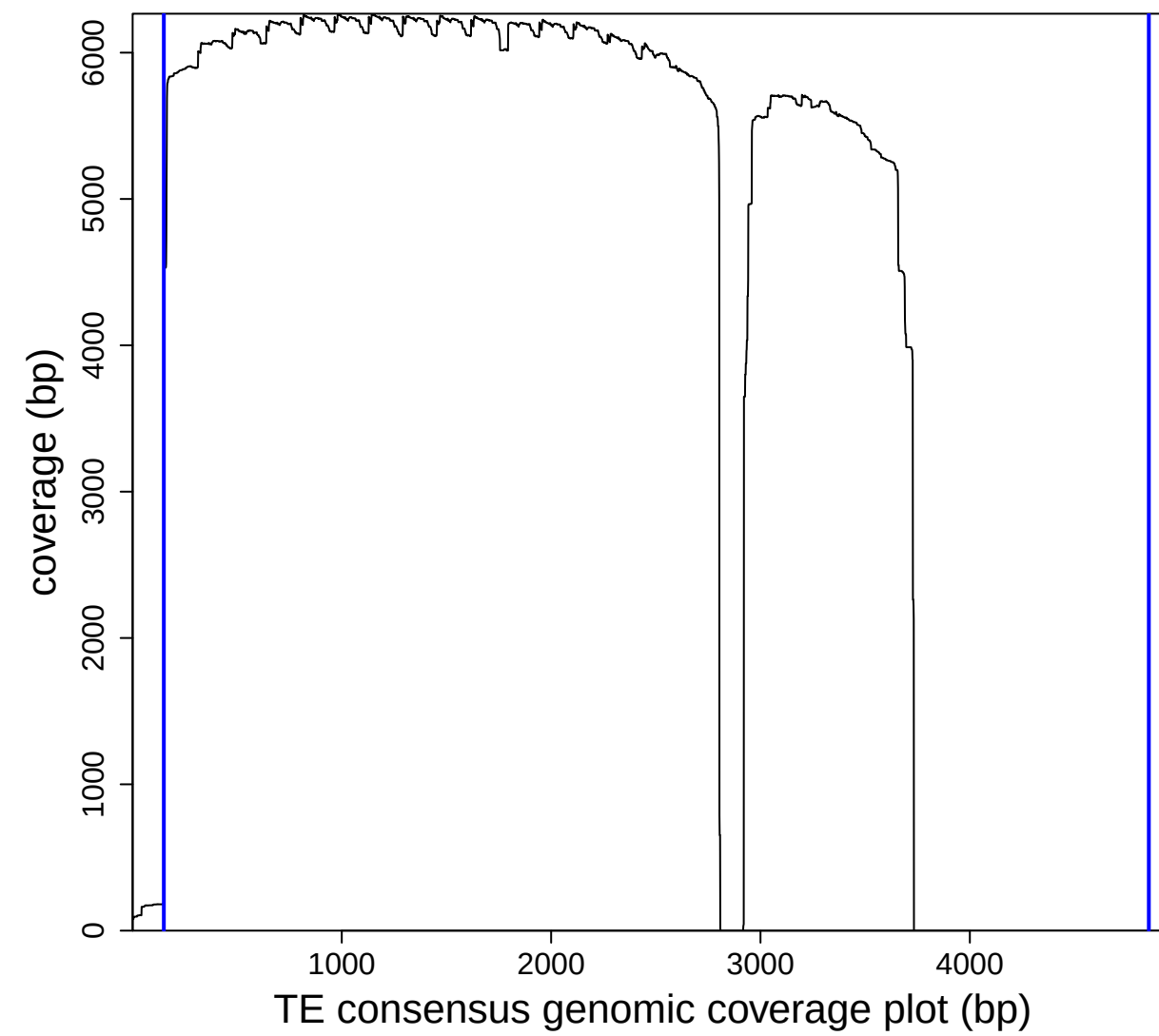
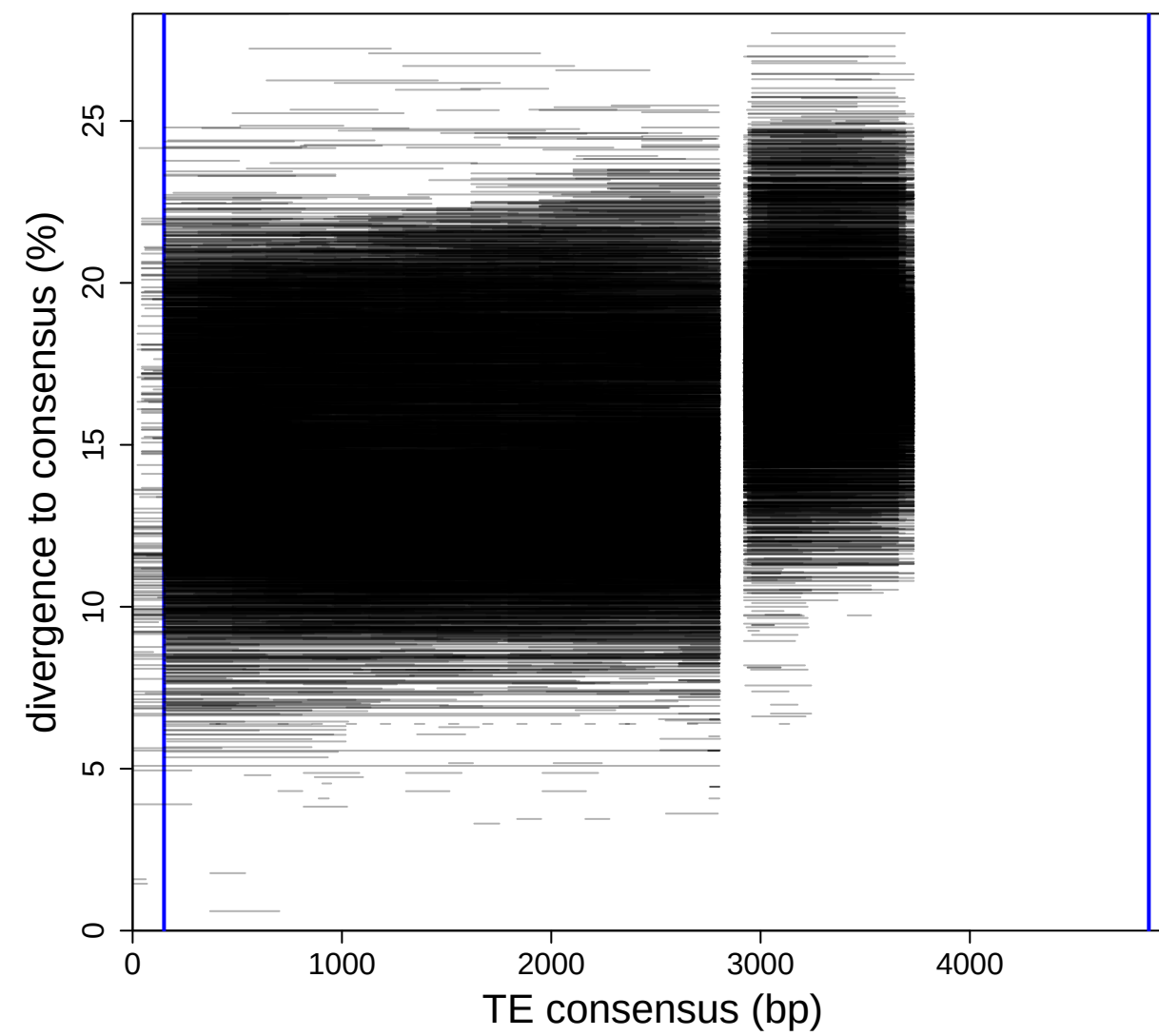
bed_uf.bed g_4.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa c
size: 4688bp; fragments: 23441; full length: 0 (>=4219.2bp)

After TEtrimmer 4688 bp

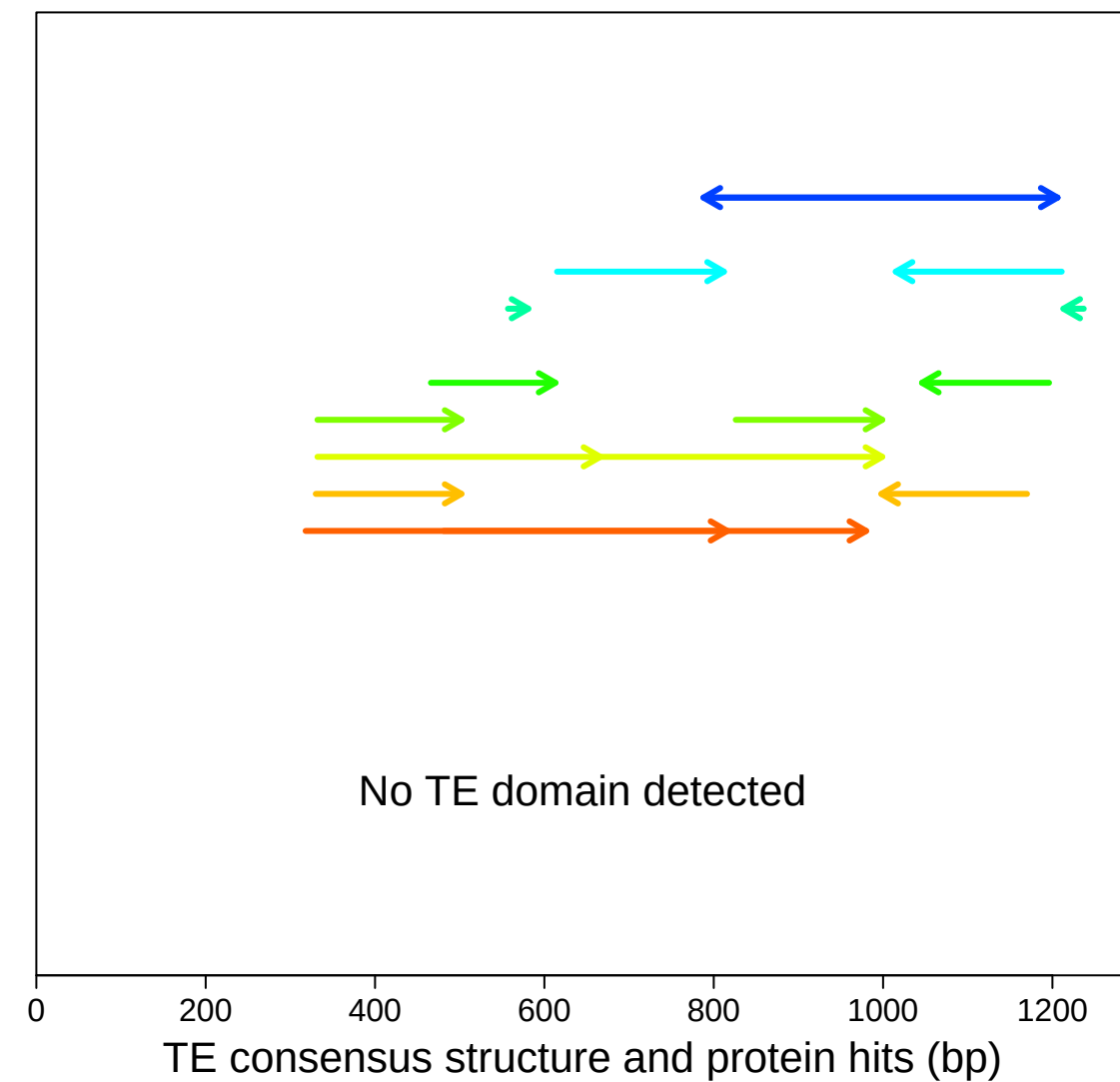
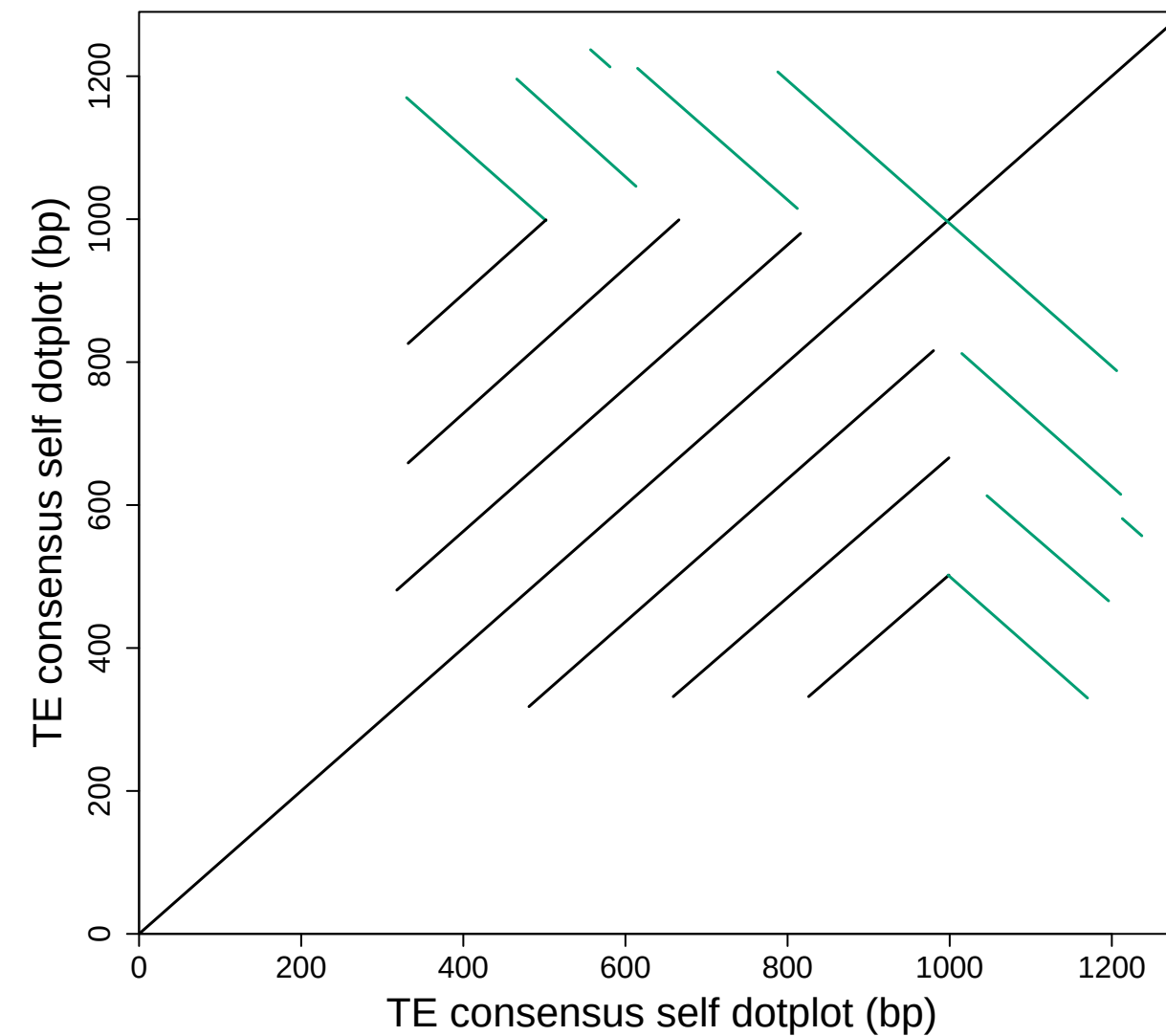
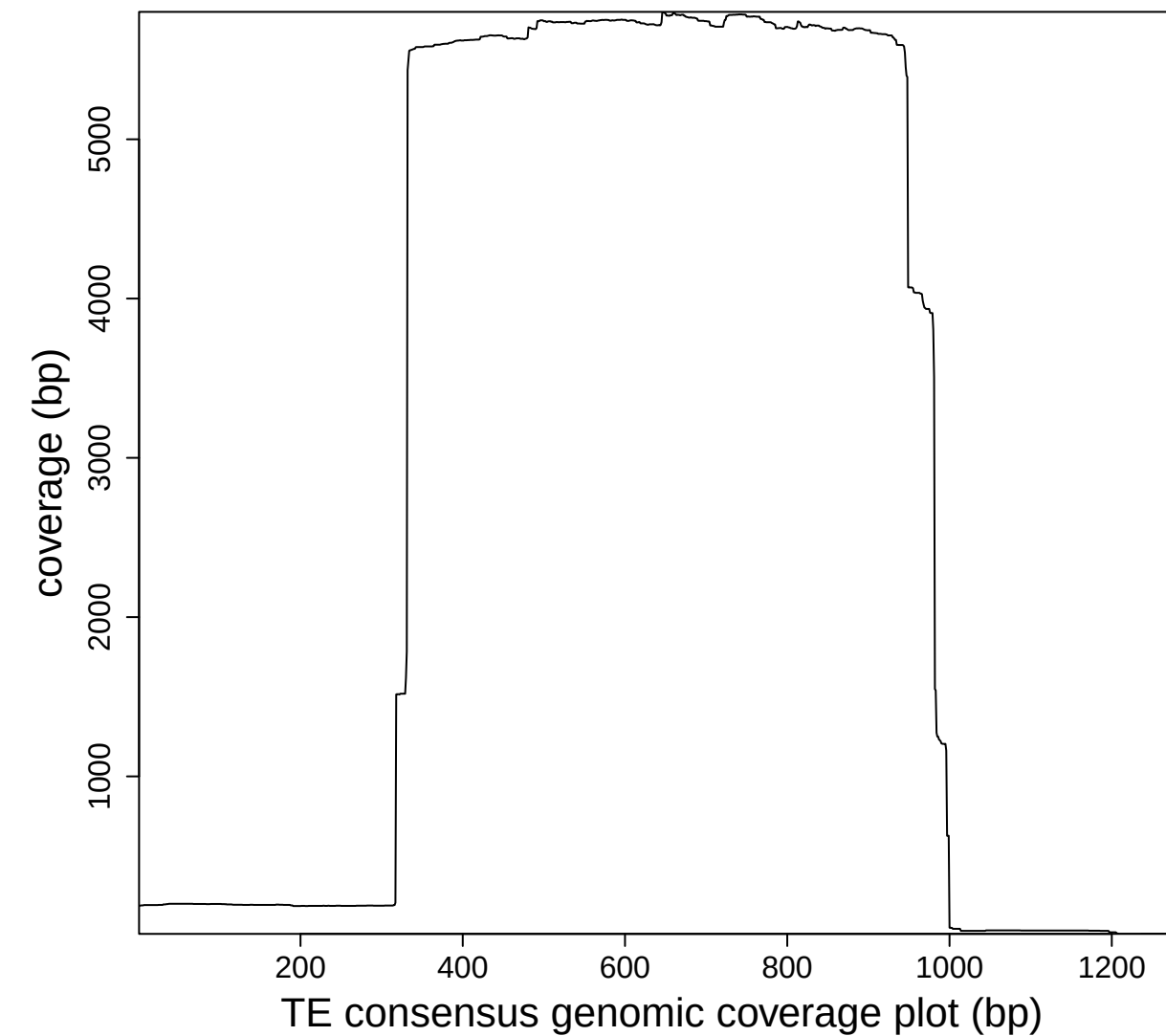
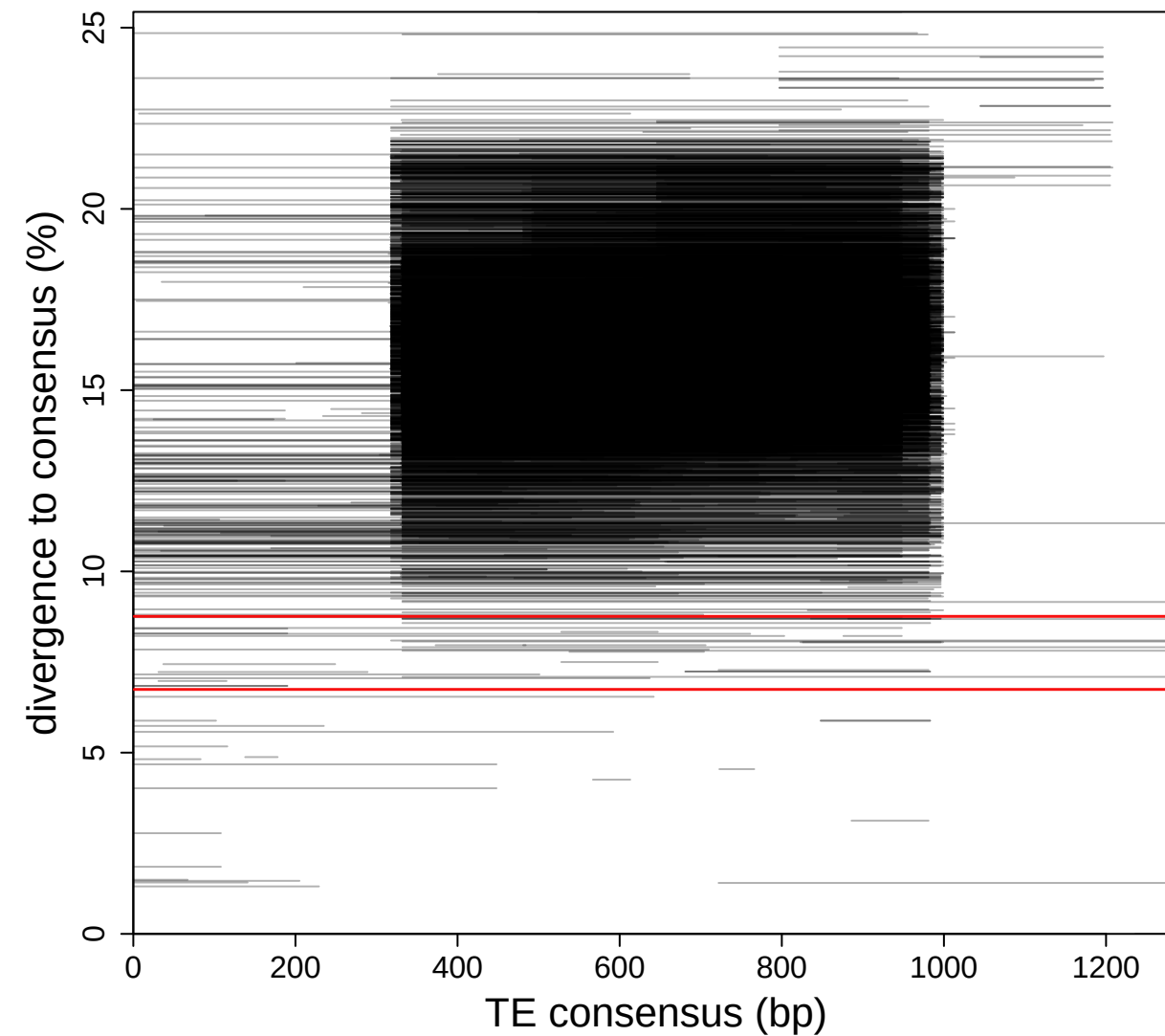


ly_8603.fasta.b.bed uf.bed g_4.bed fm_1.bed_0_0 bcln.fa aln.f
size: 4970bp; fragments: 15698; full length: 0 (>=4473bp)

After TEtrimmer Extended plot Blue lines are boundaries

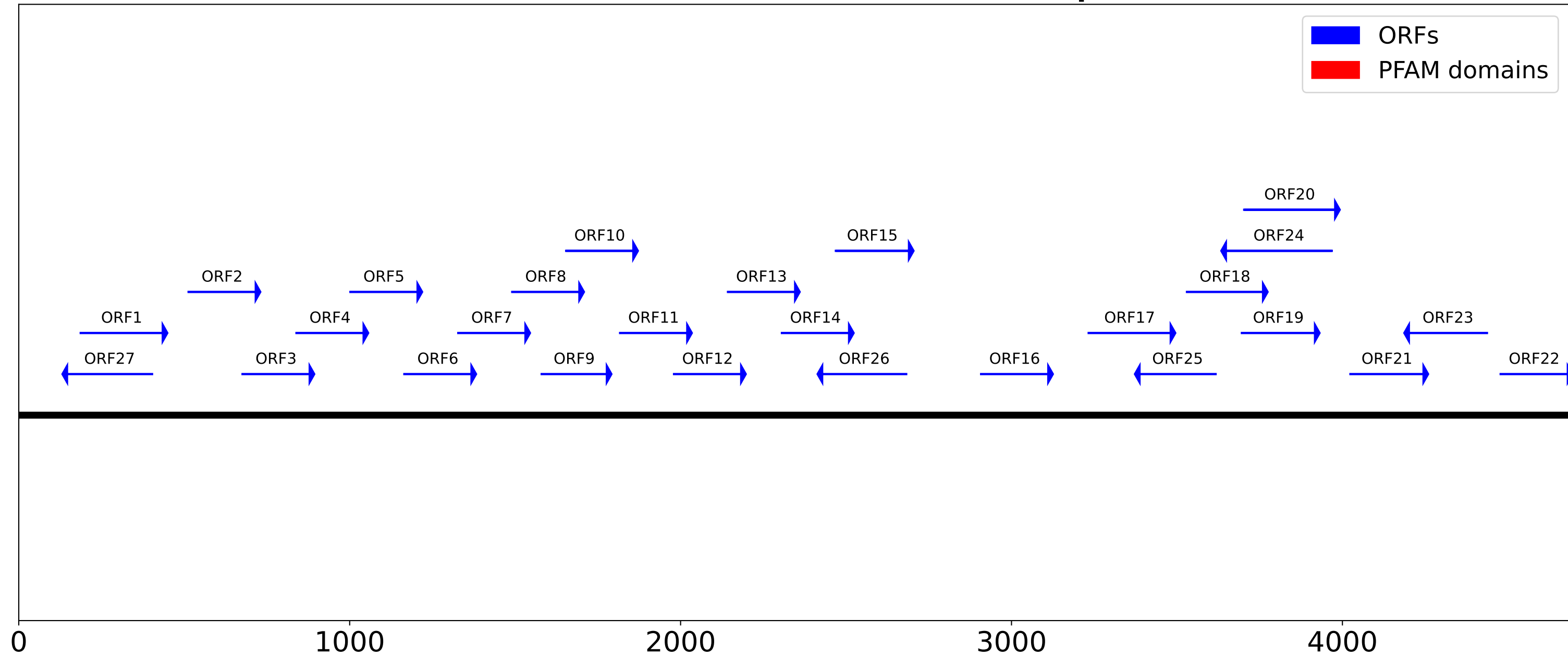


TE: rnd_5_family_8603
size: 1290bp; fragments: 6250; full length: 2 (≥ 1161 bp)

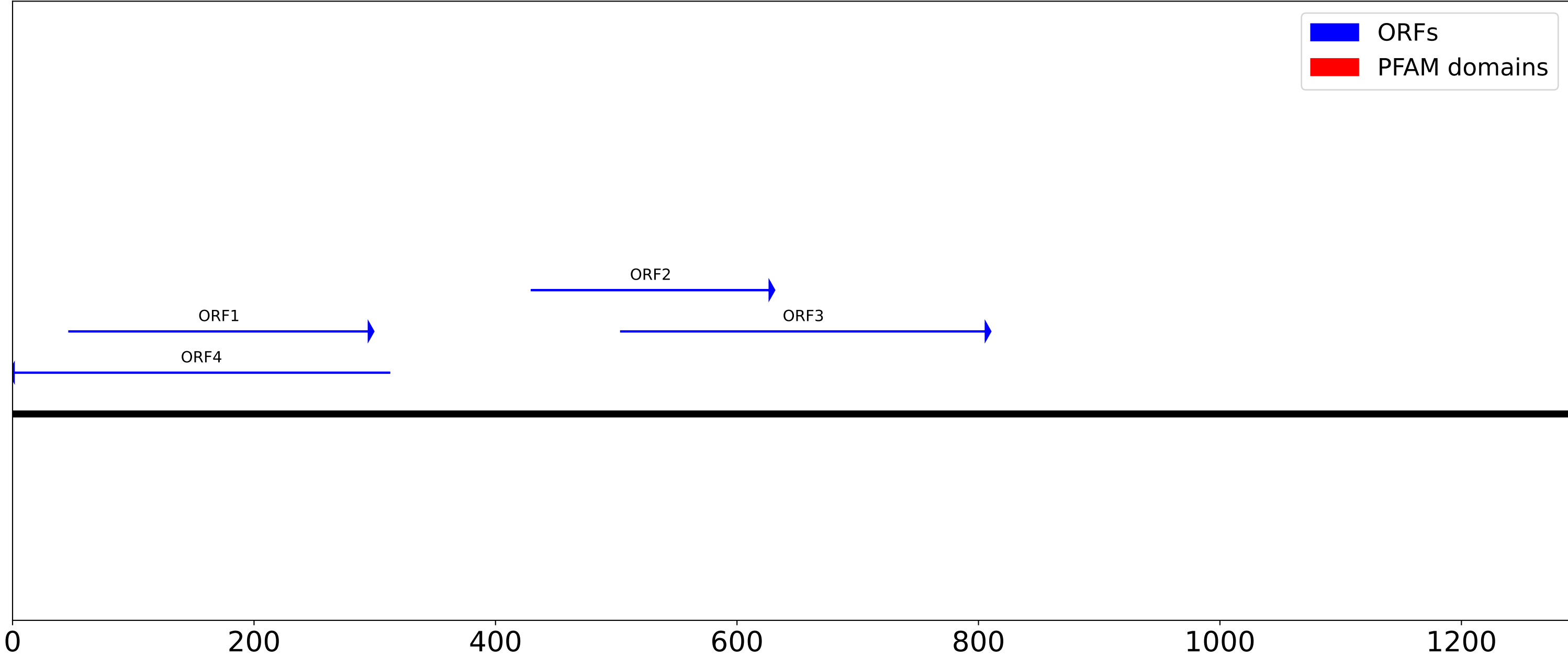


Before TEtrimmer 1290 bp

After TEtrimmer ORF and PFAM domain plot



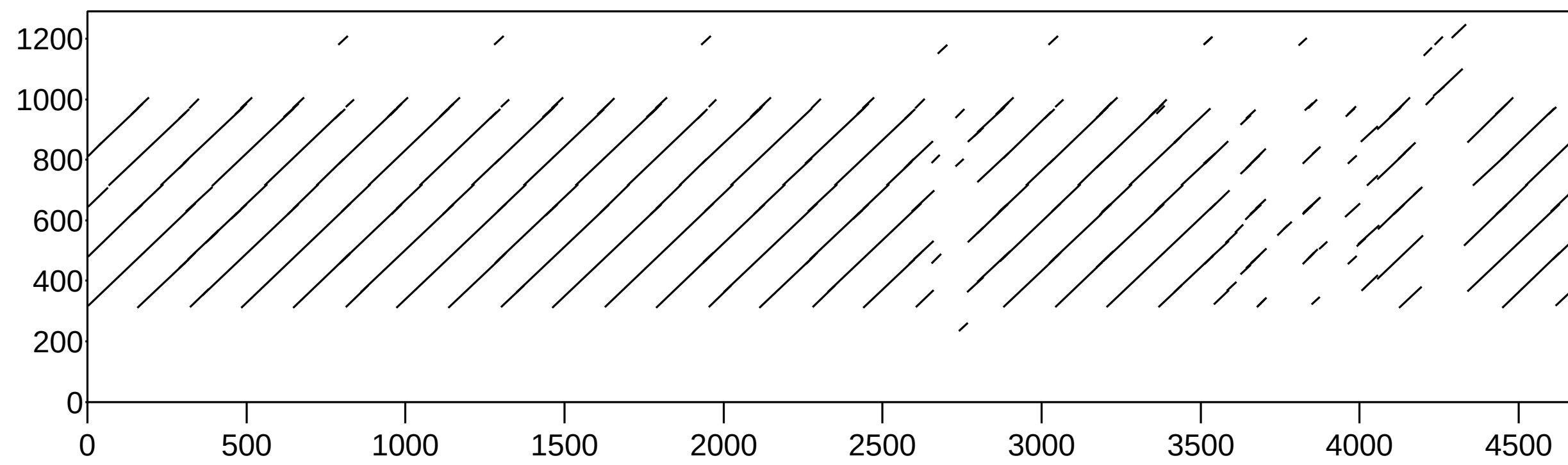
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)