

Start crop Point



End crop Point

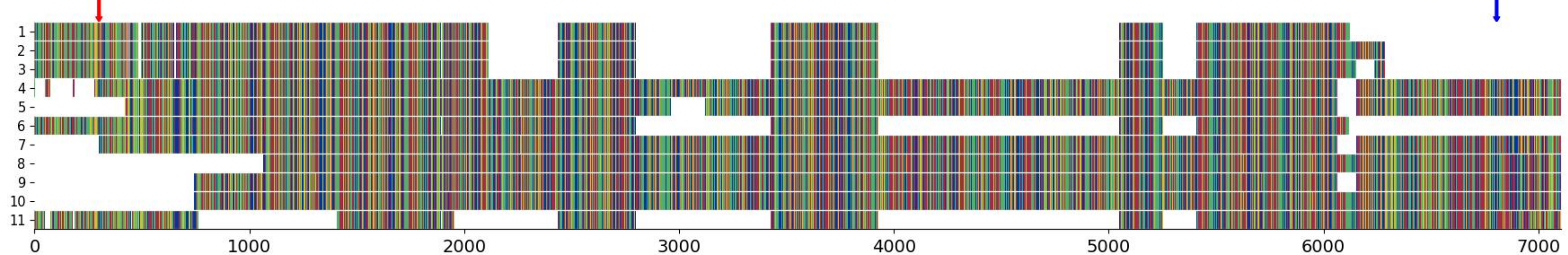


MSA length = 6507



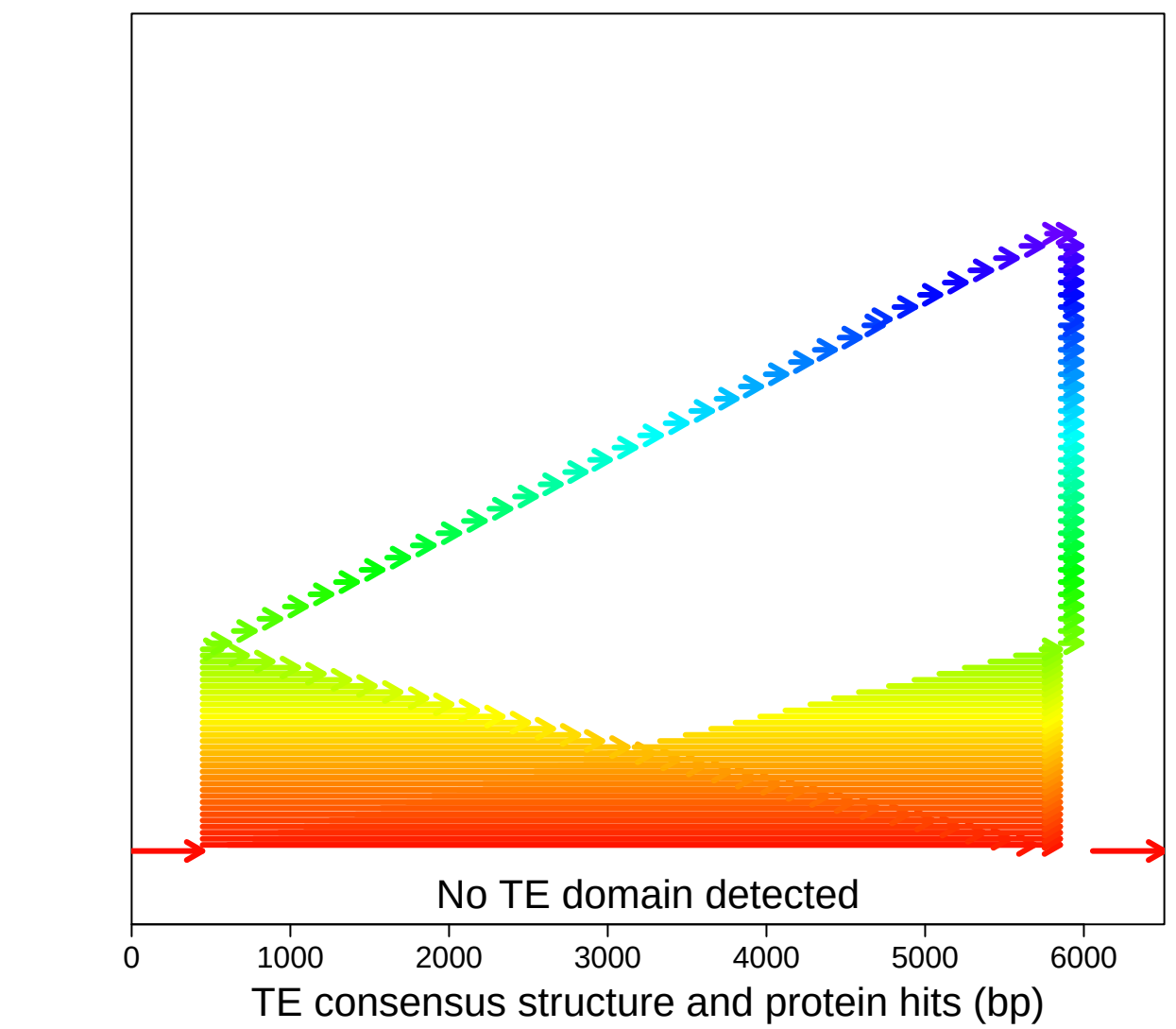
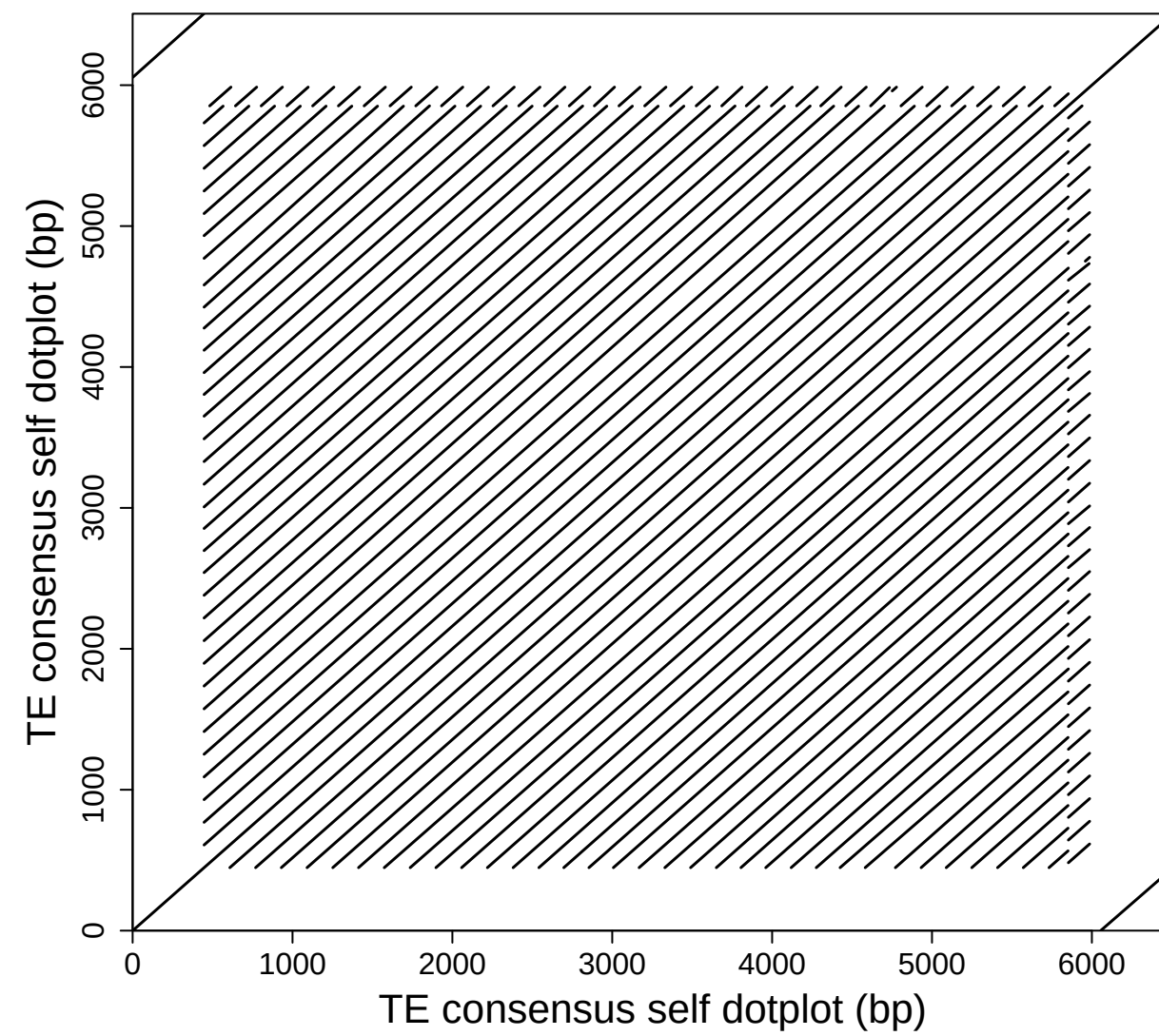
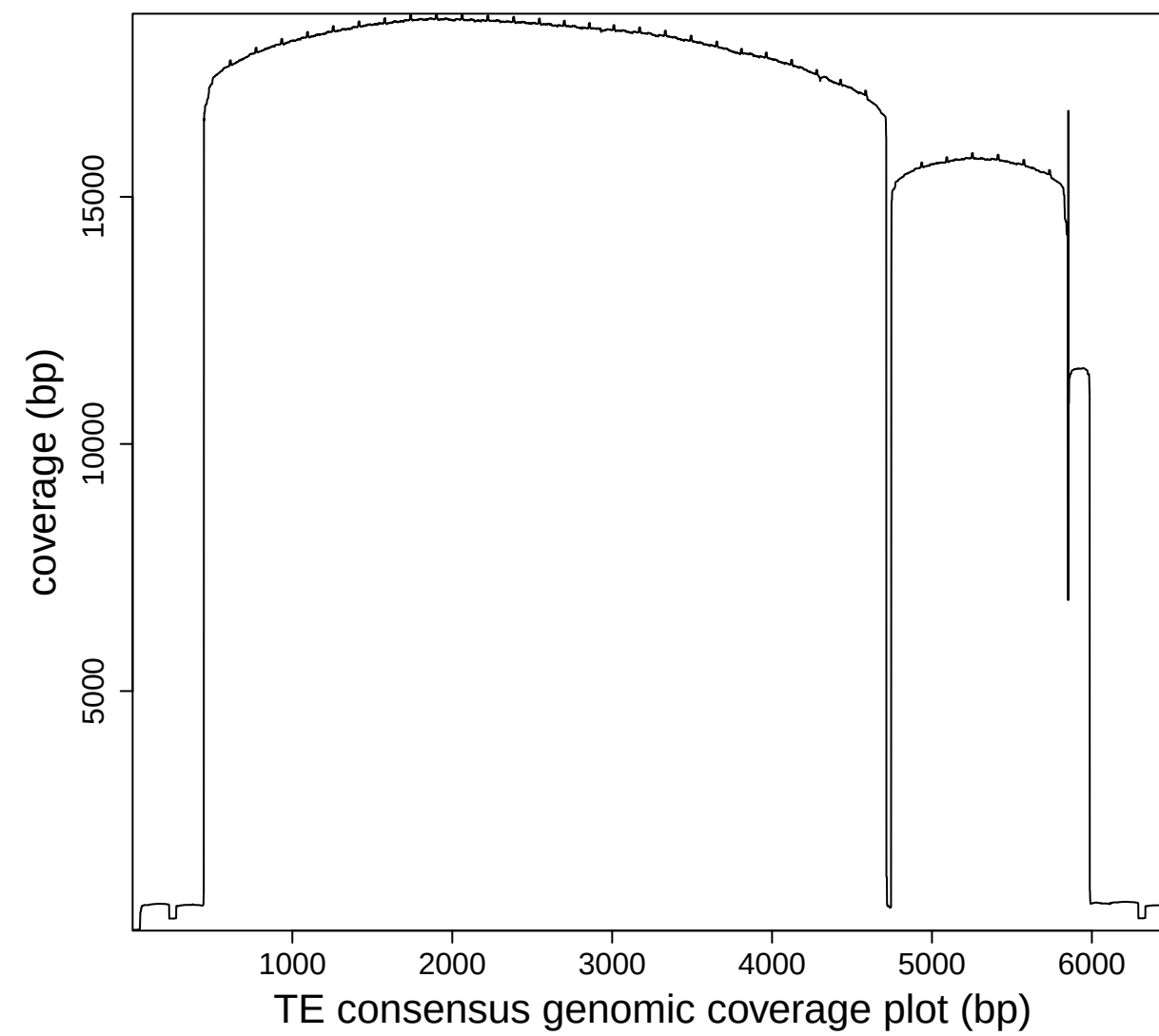
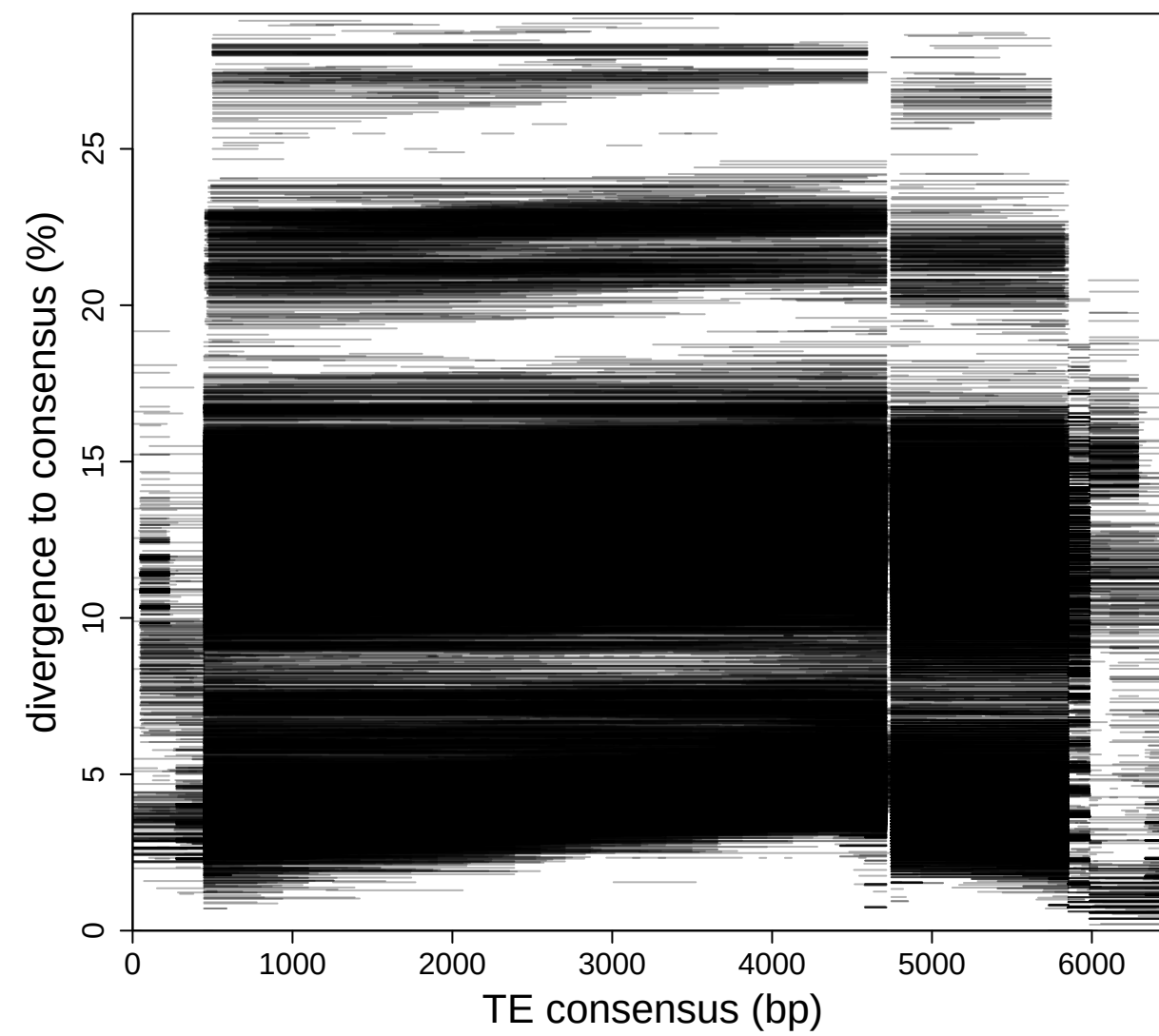
Start crop Point

End crop Point



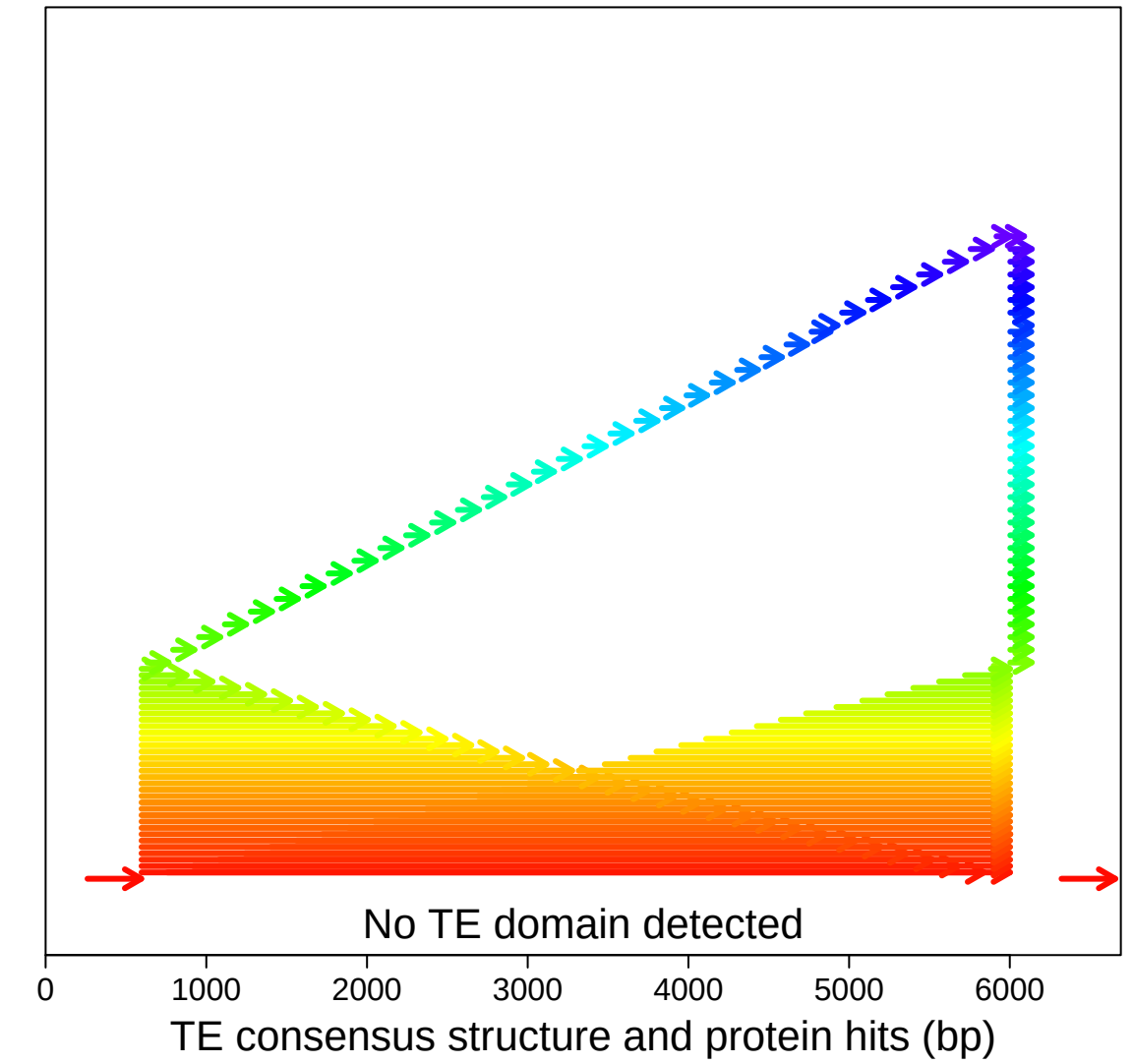
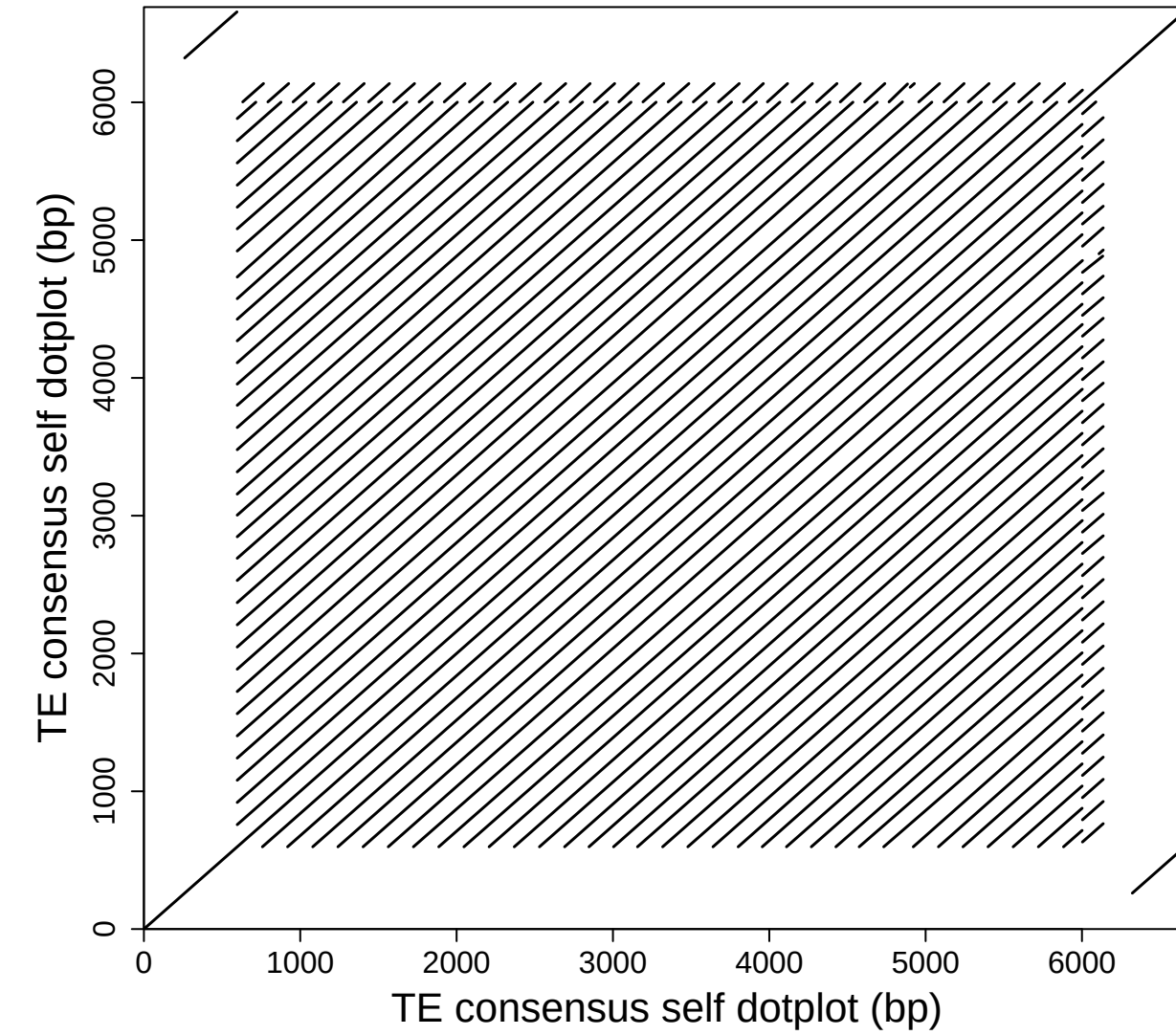
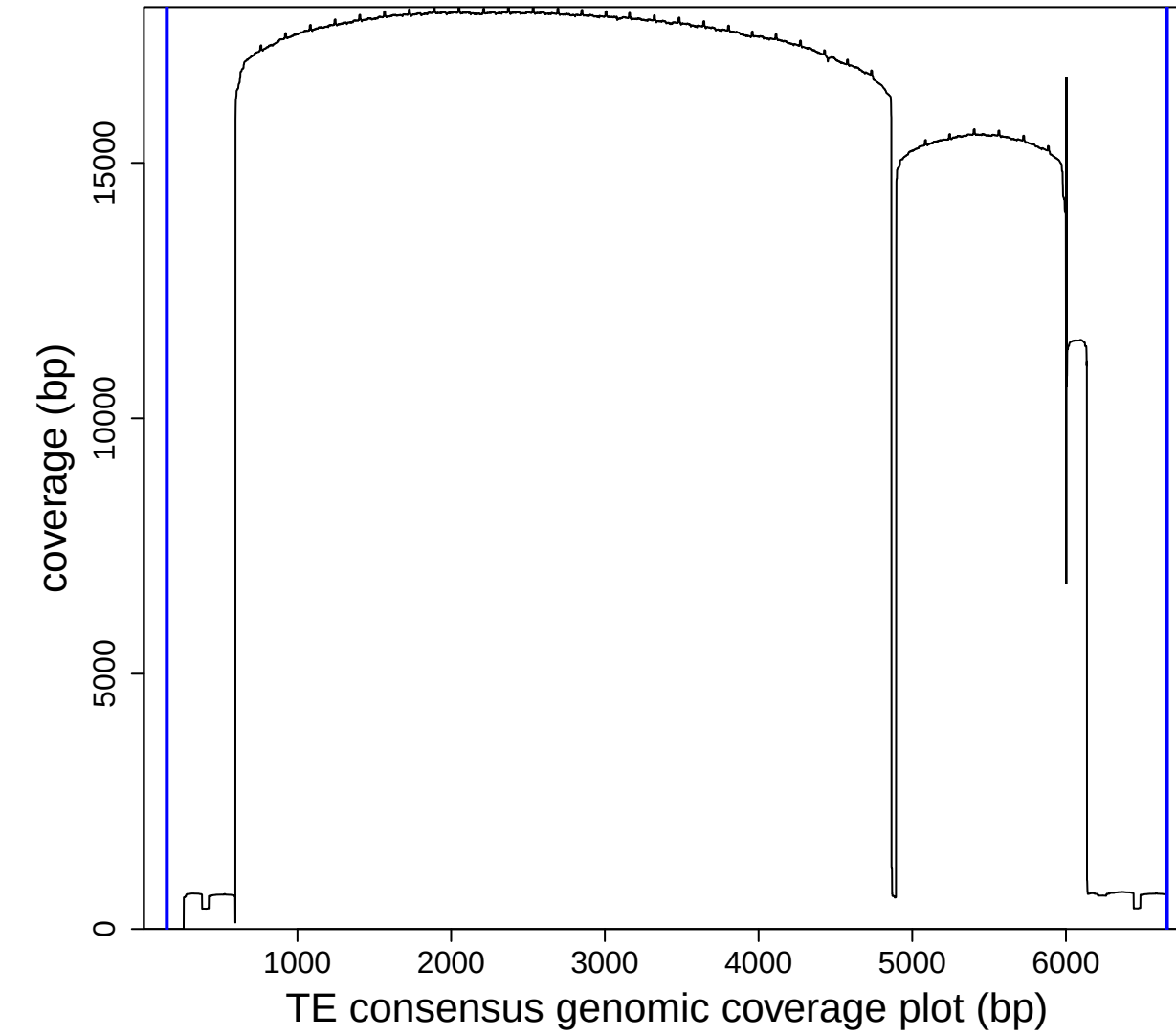
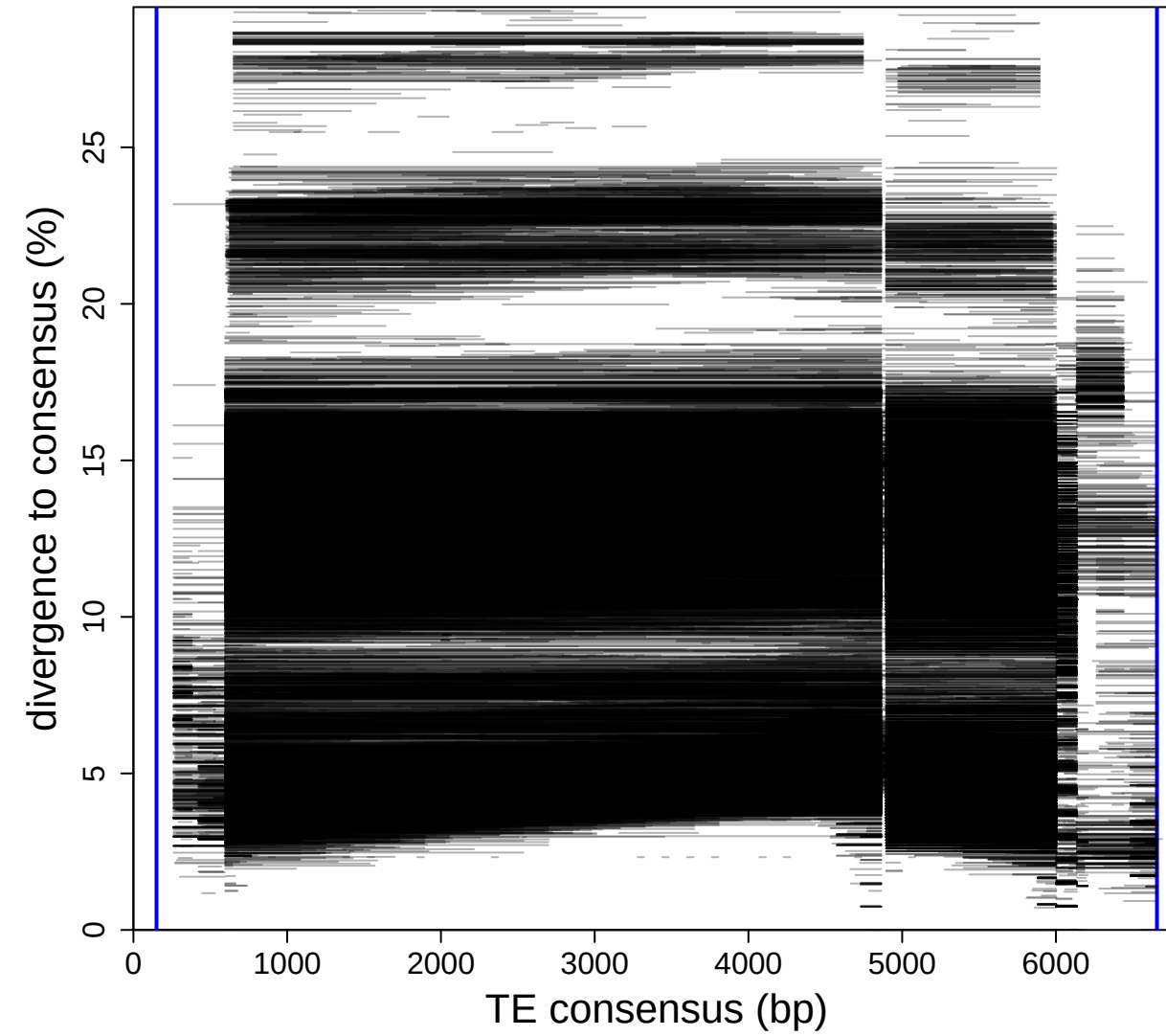
bed uf.bed g_3.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa ce
size: 6507bp; fragments: 61825; full length: 0 (>=5856.3bp)

After TEtrimmer 6507 bp



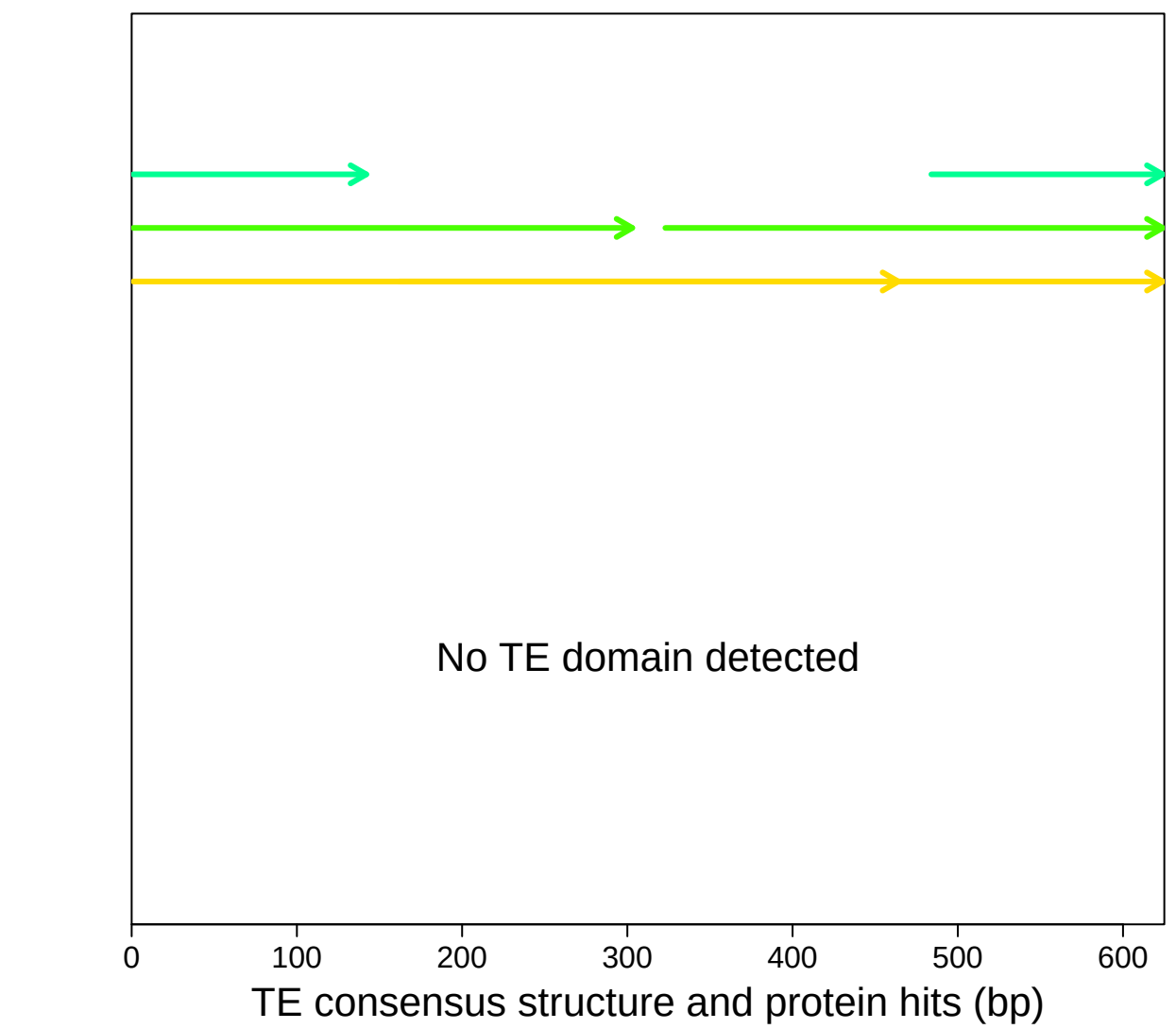
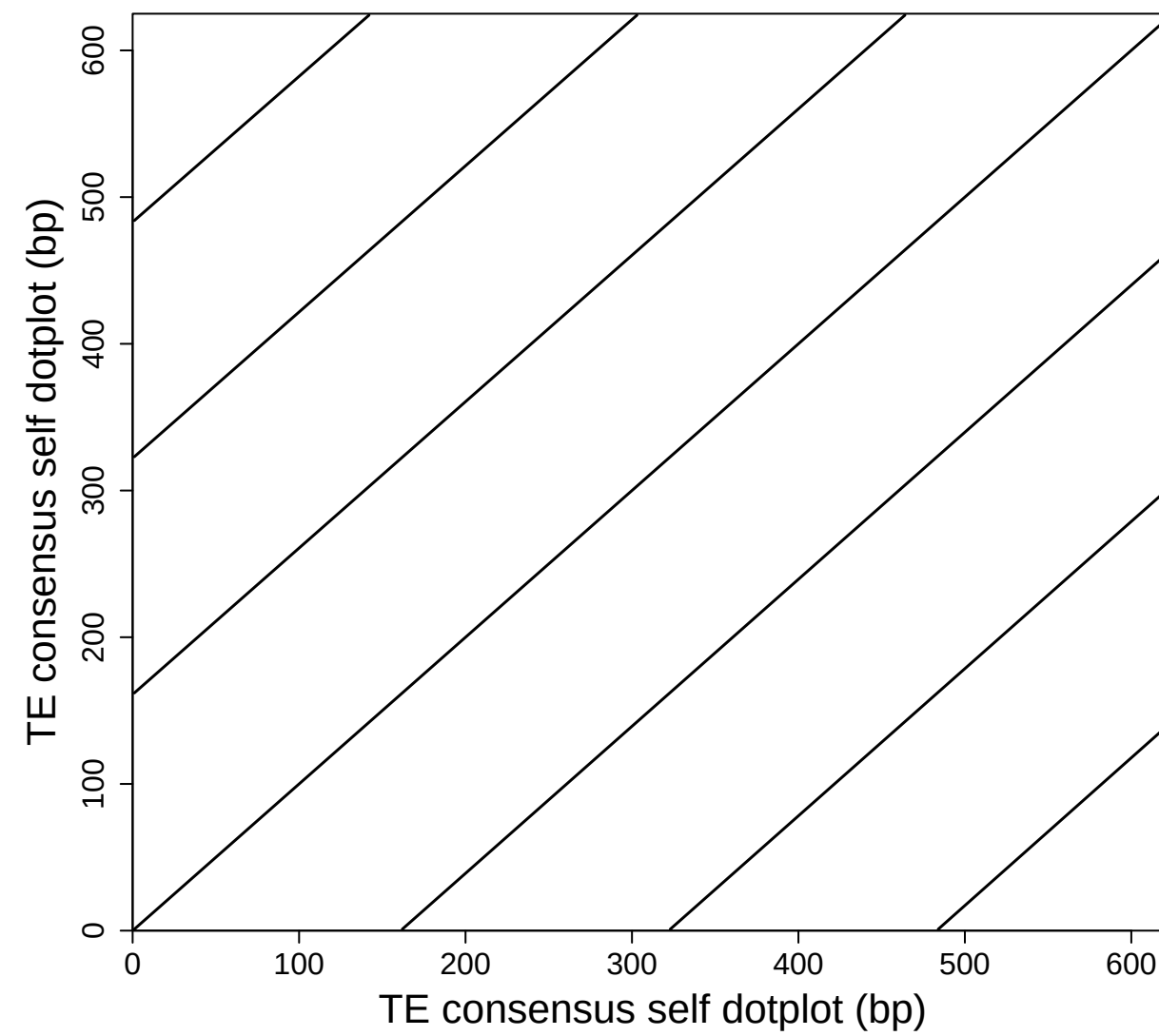
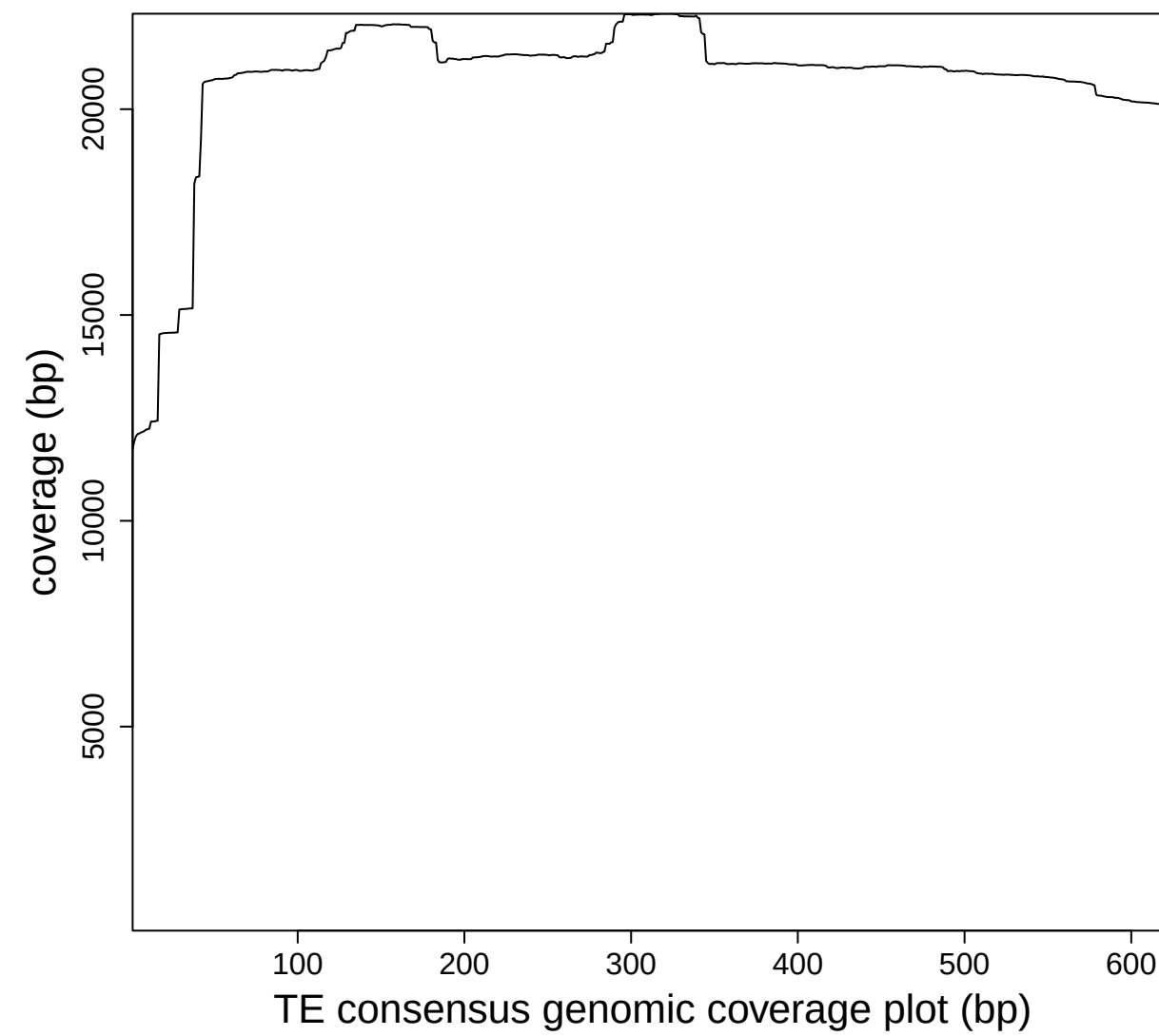
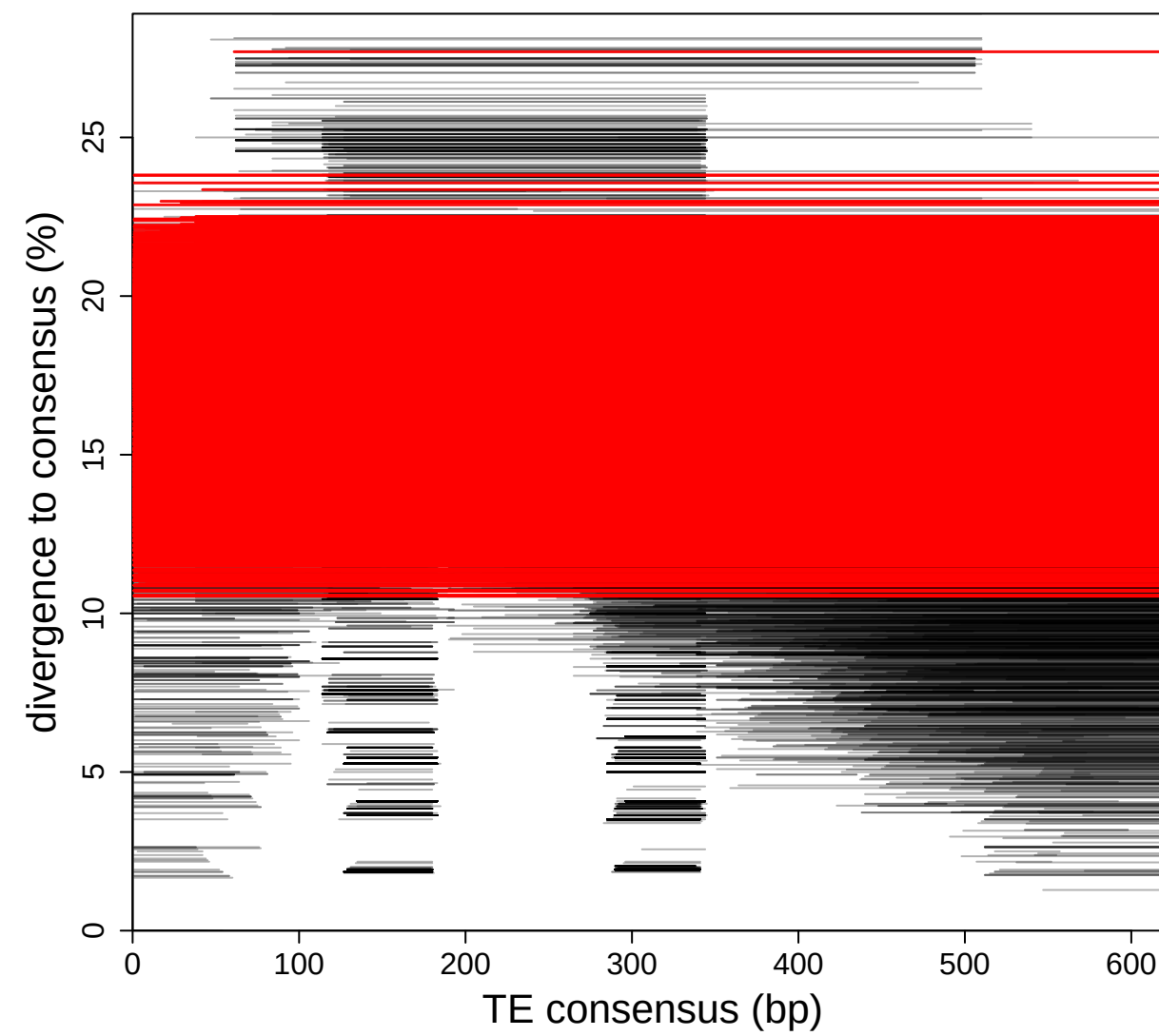
ily_588.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 6807bp; fragments: 60705; full length: 0 (>=6126.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

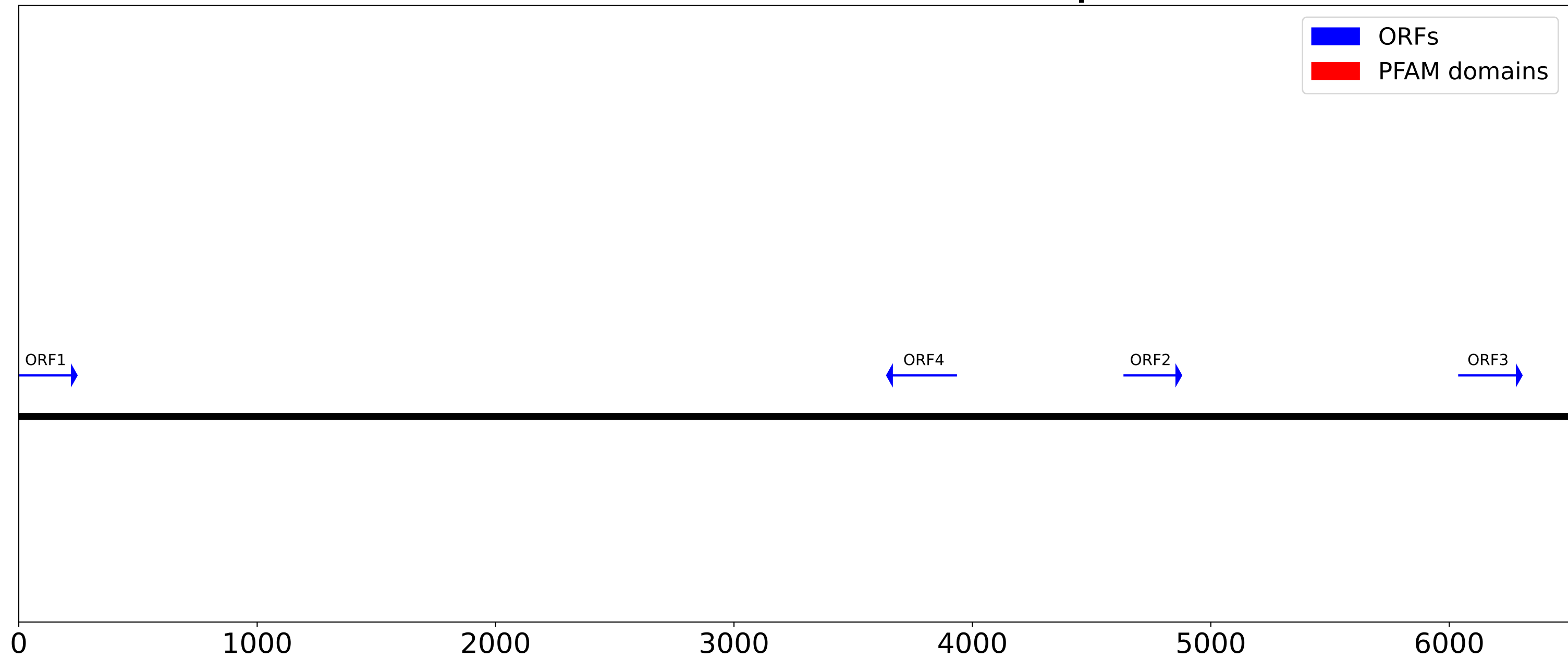


TE: rnd_4_family_588
size: 625bp; fragments: 24743; full length: 18599 (≥ 562.5 bp)

Before TEtrimmer 625 bp



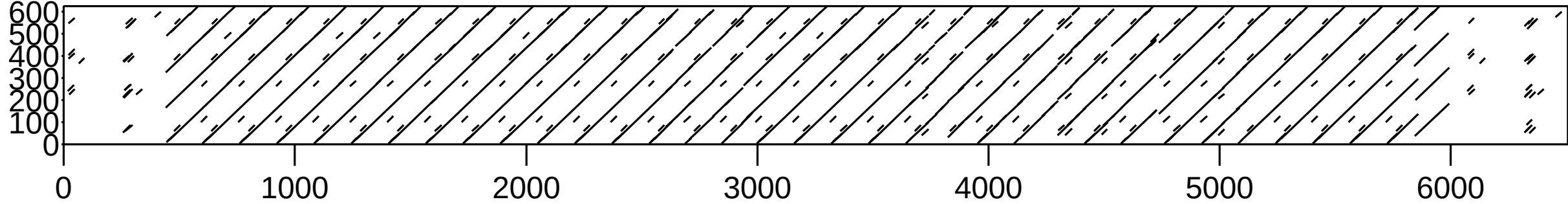
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)