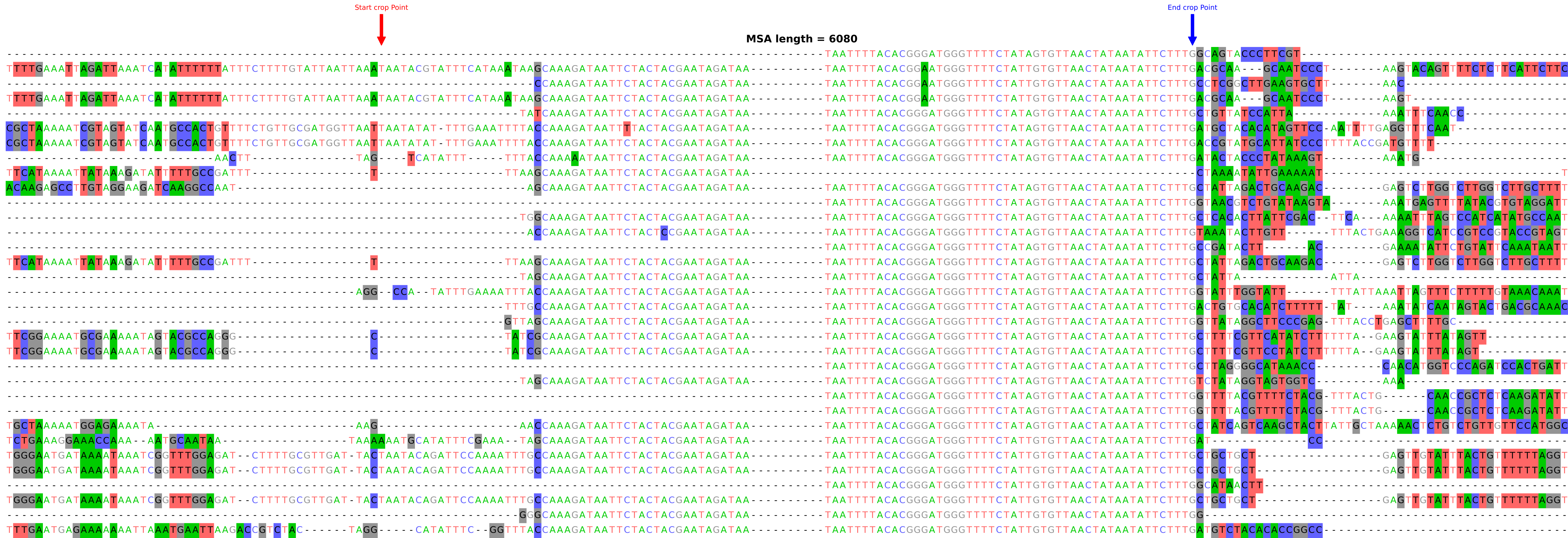
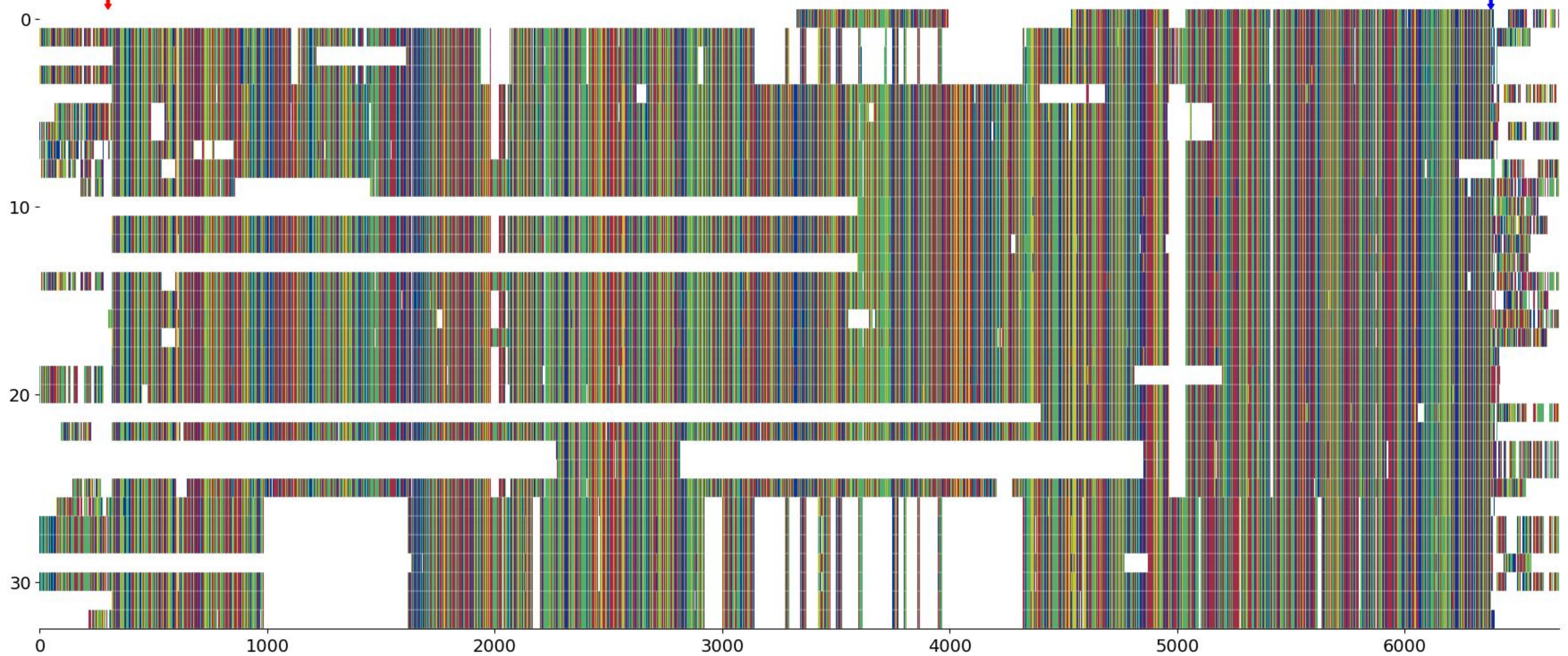




Start crop Point

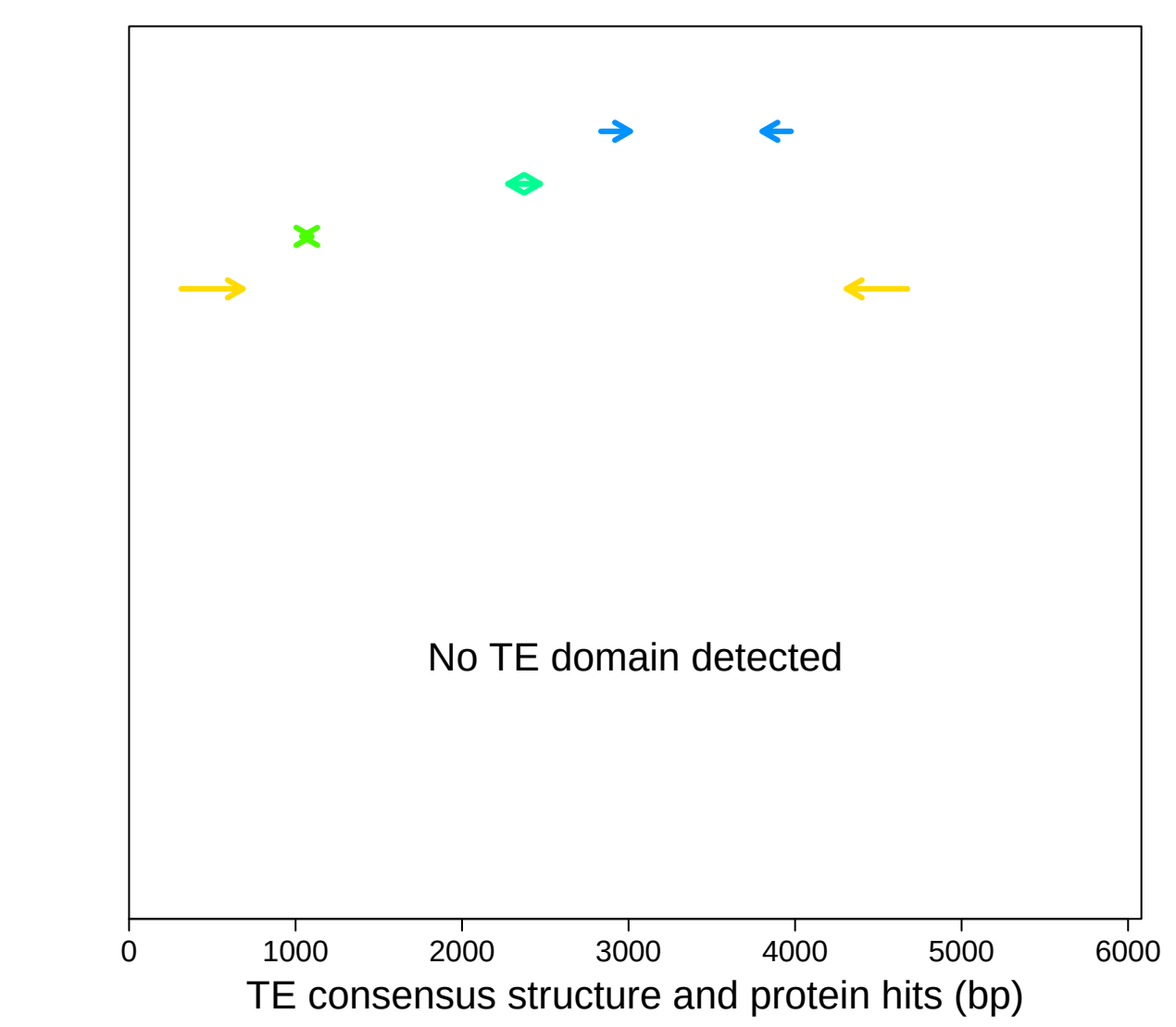
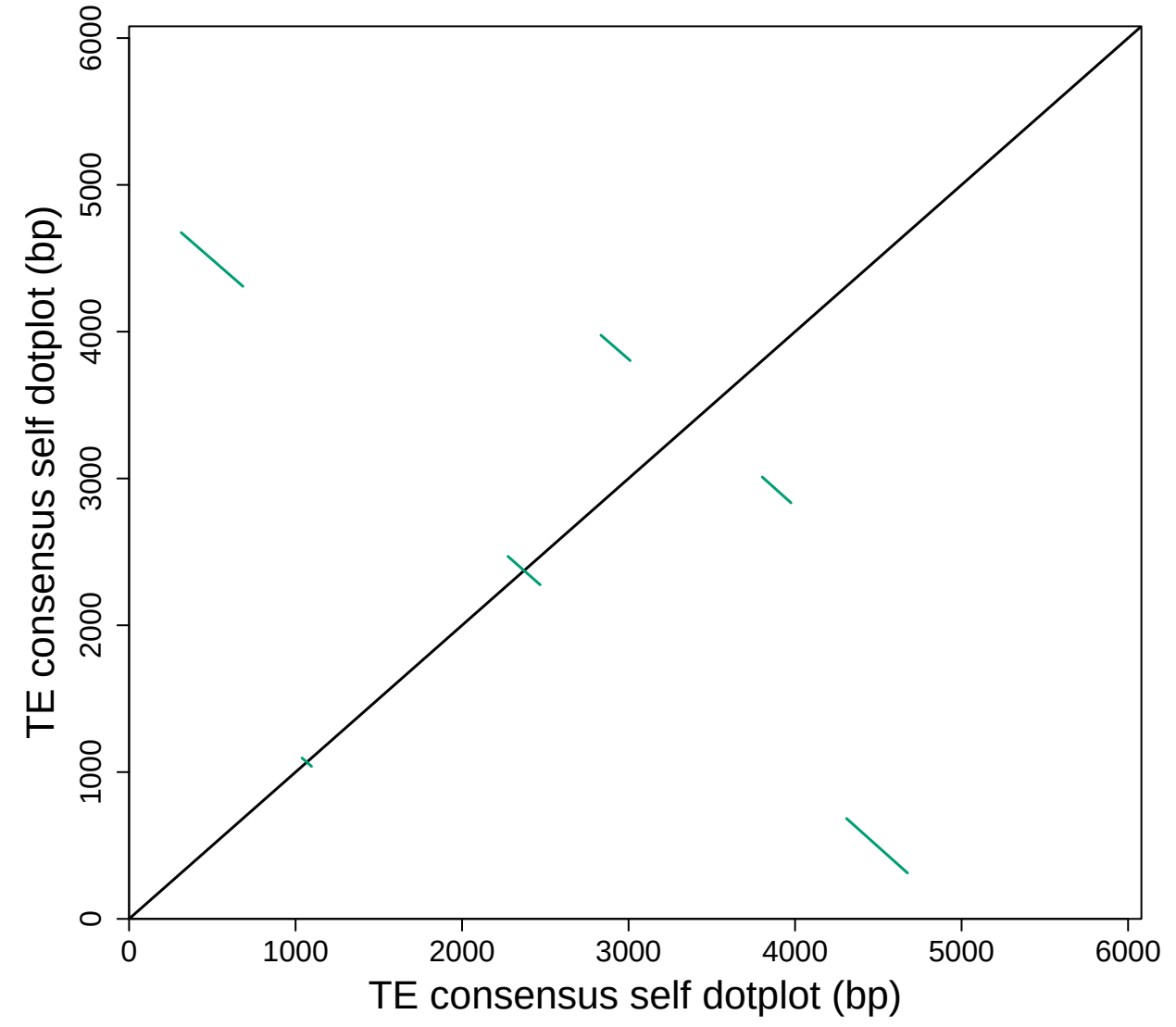
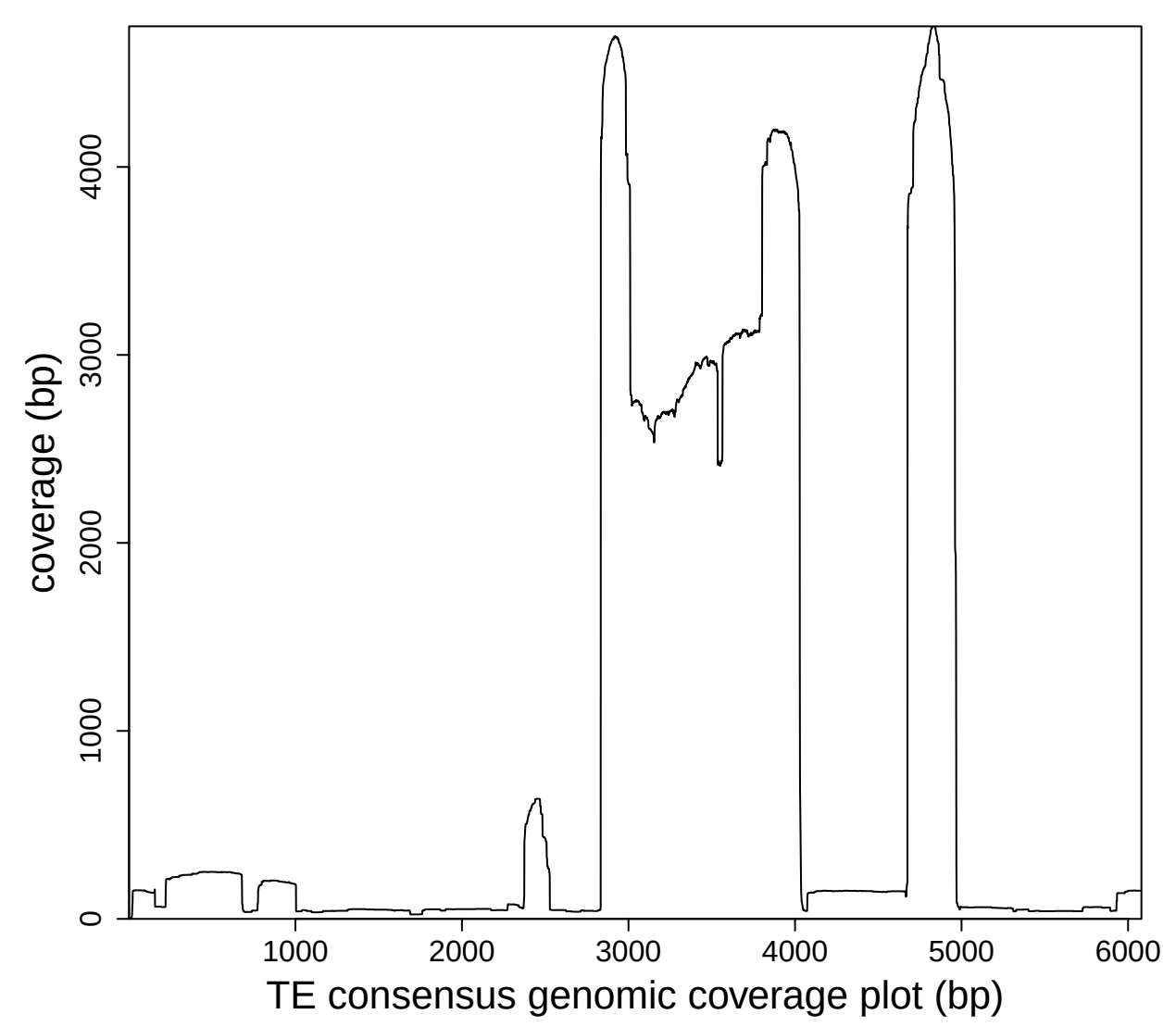
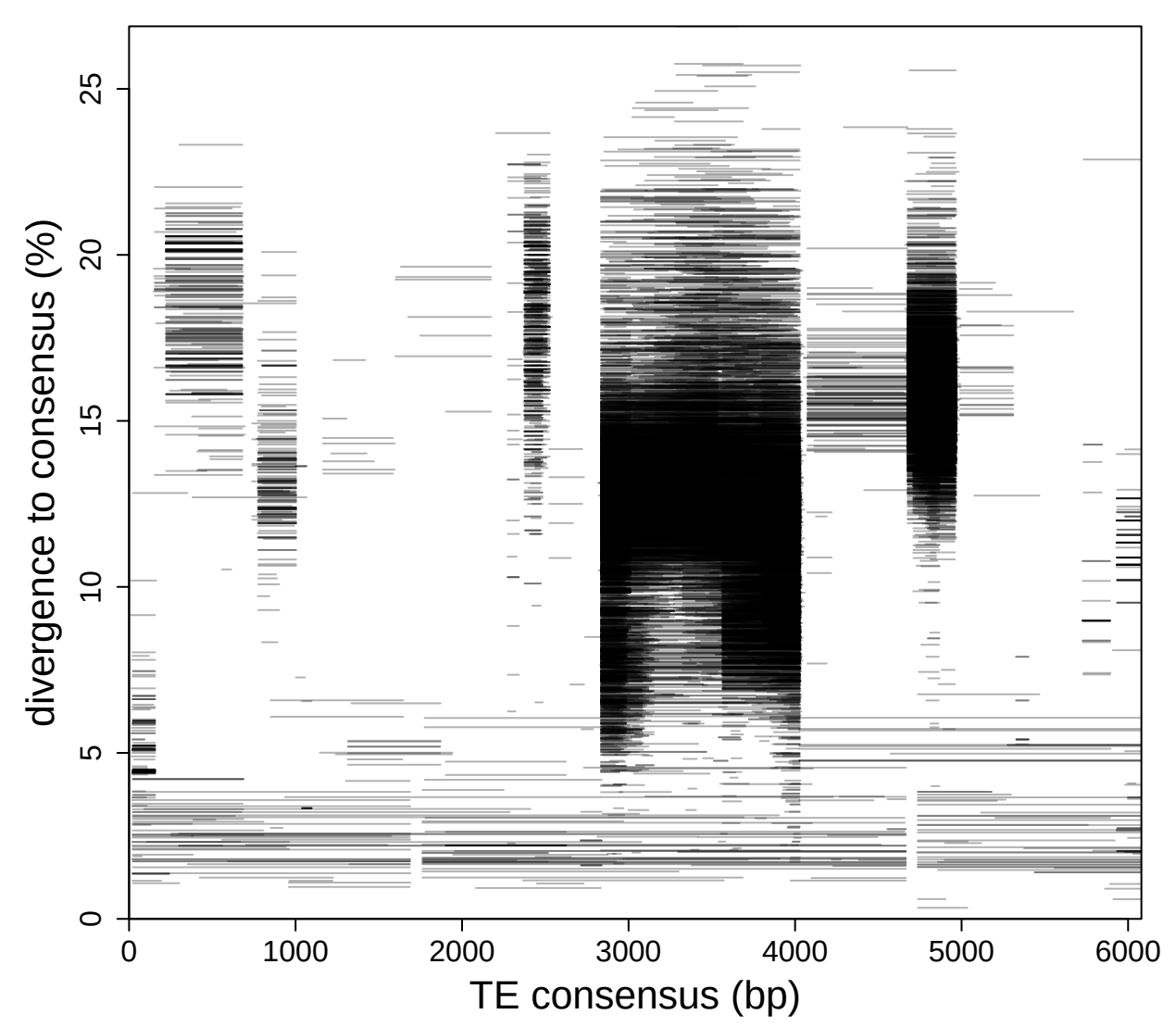


End crop Point



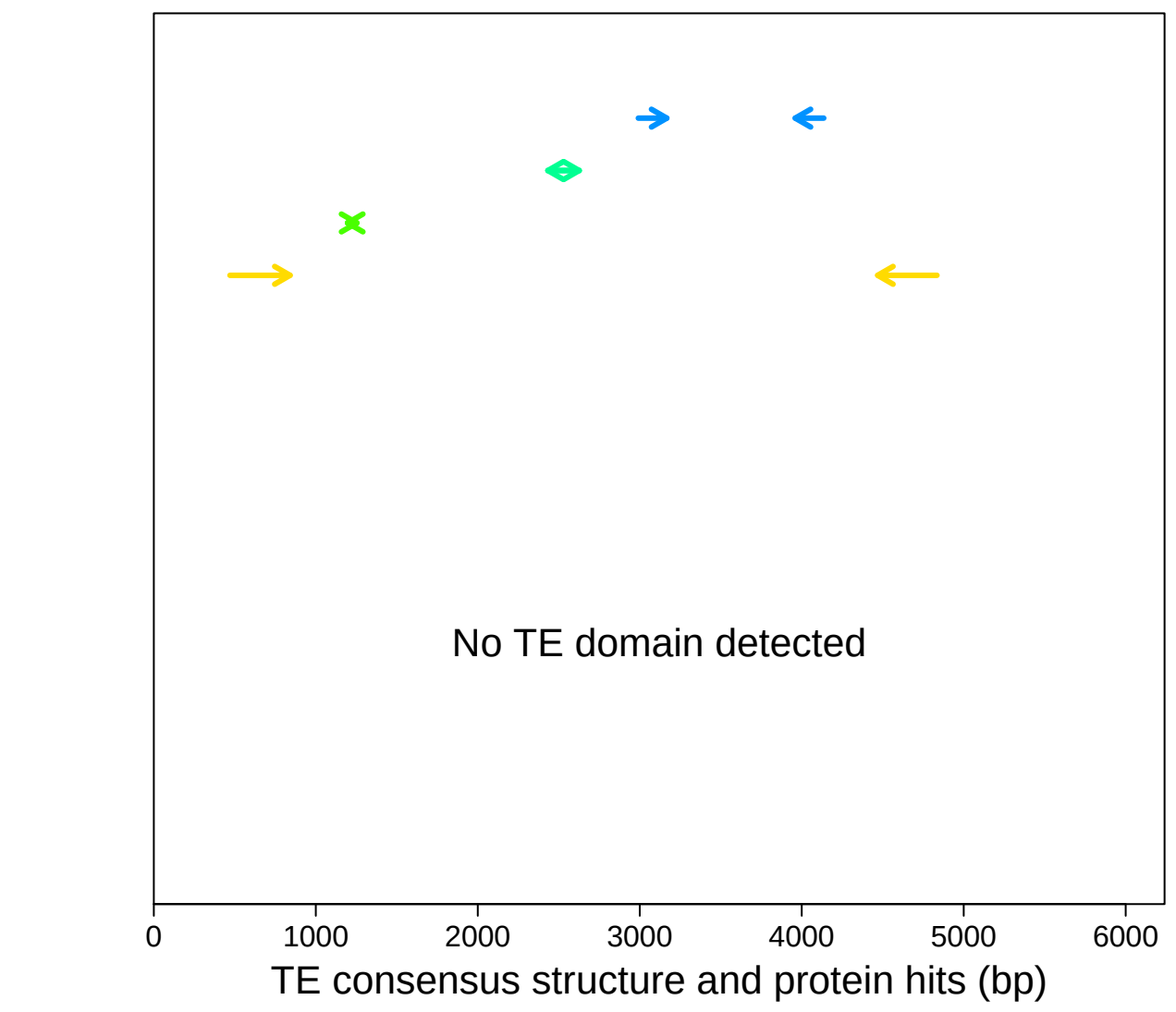
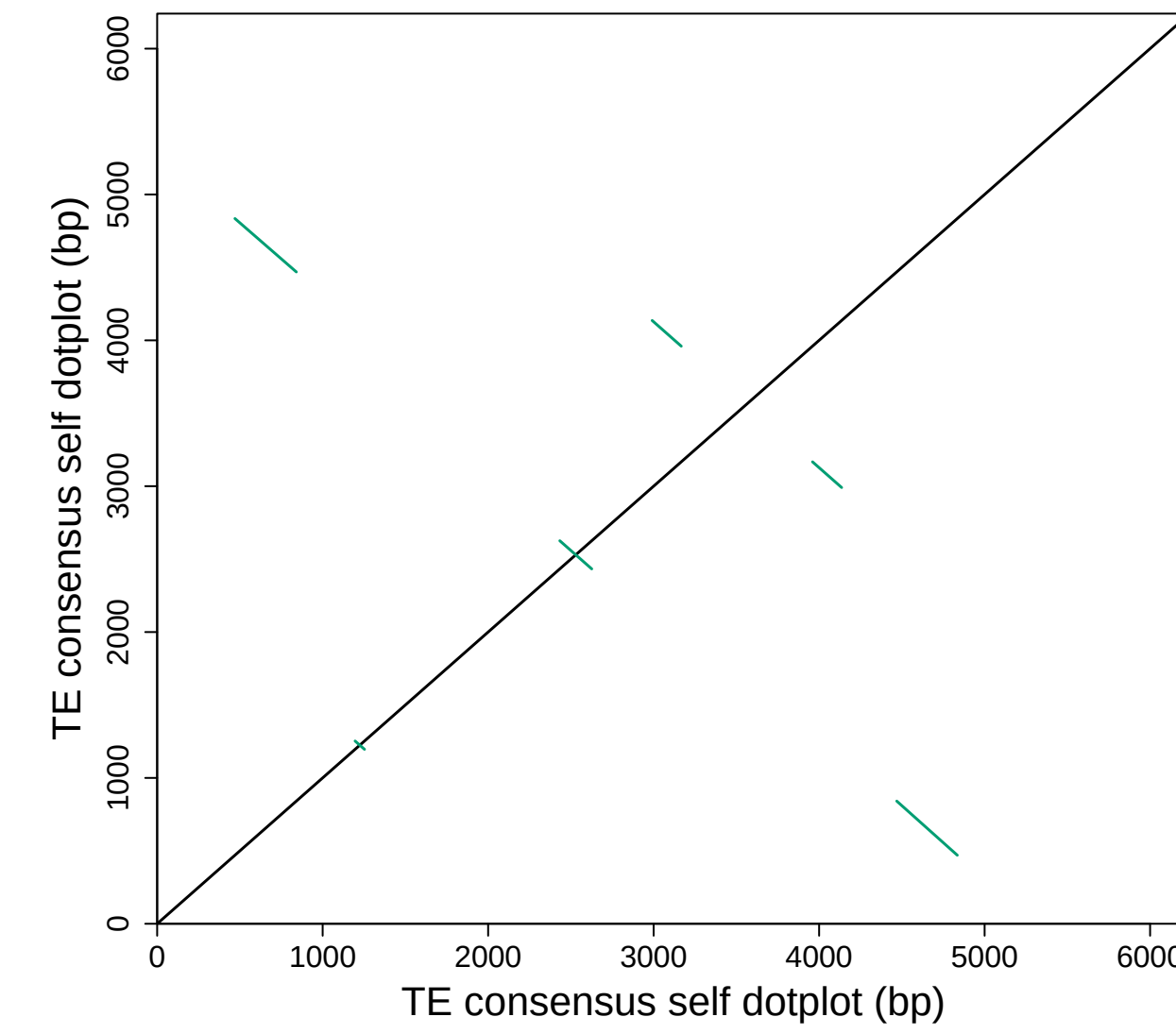
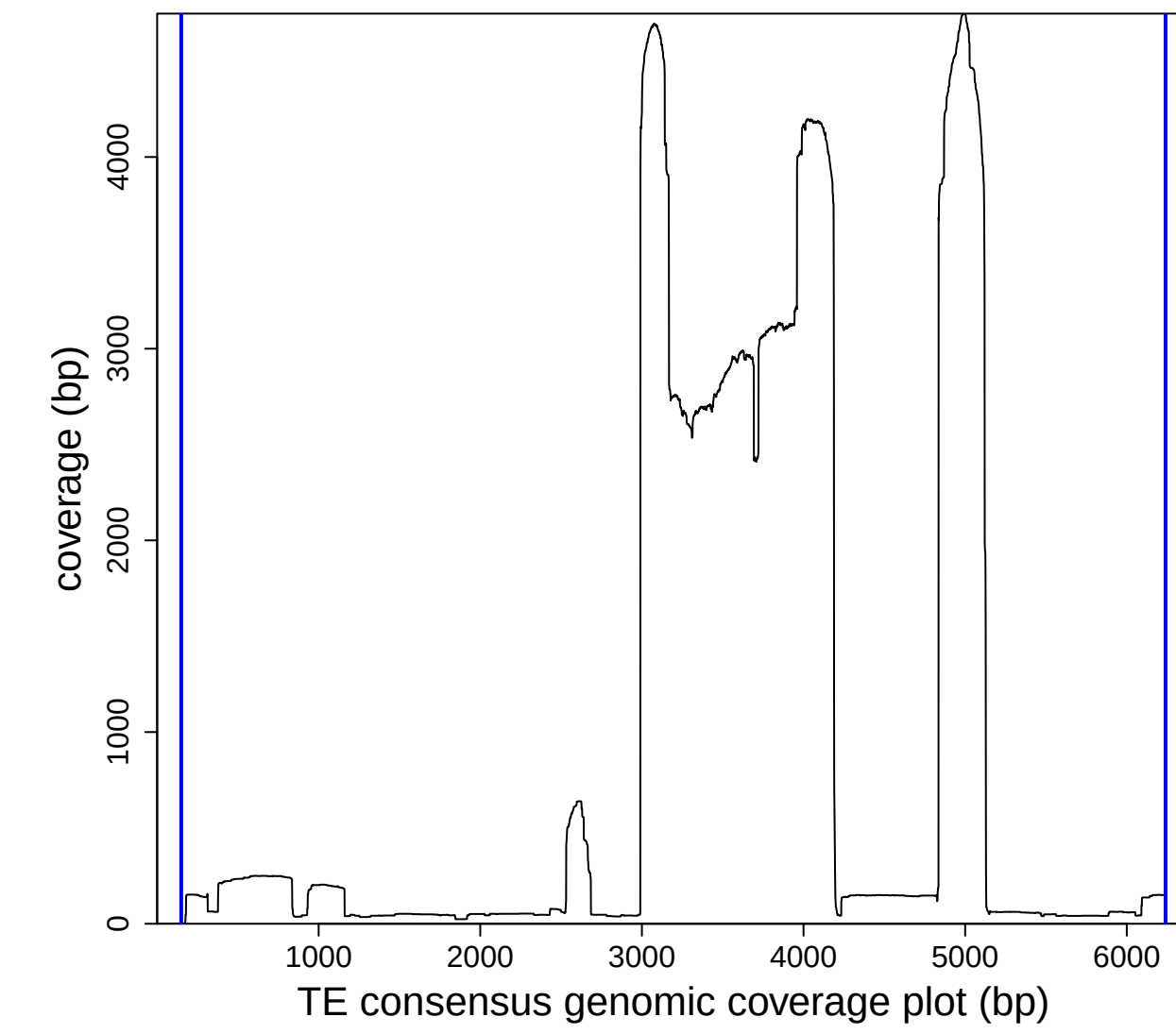
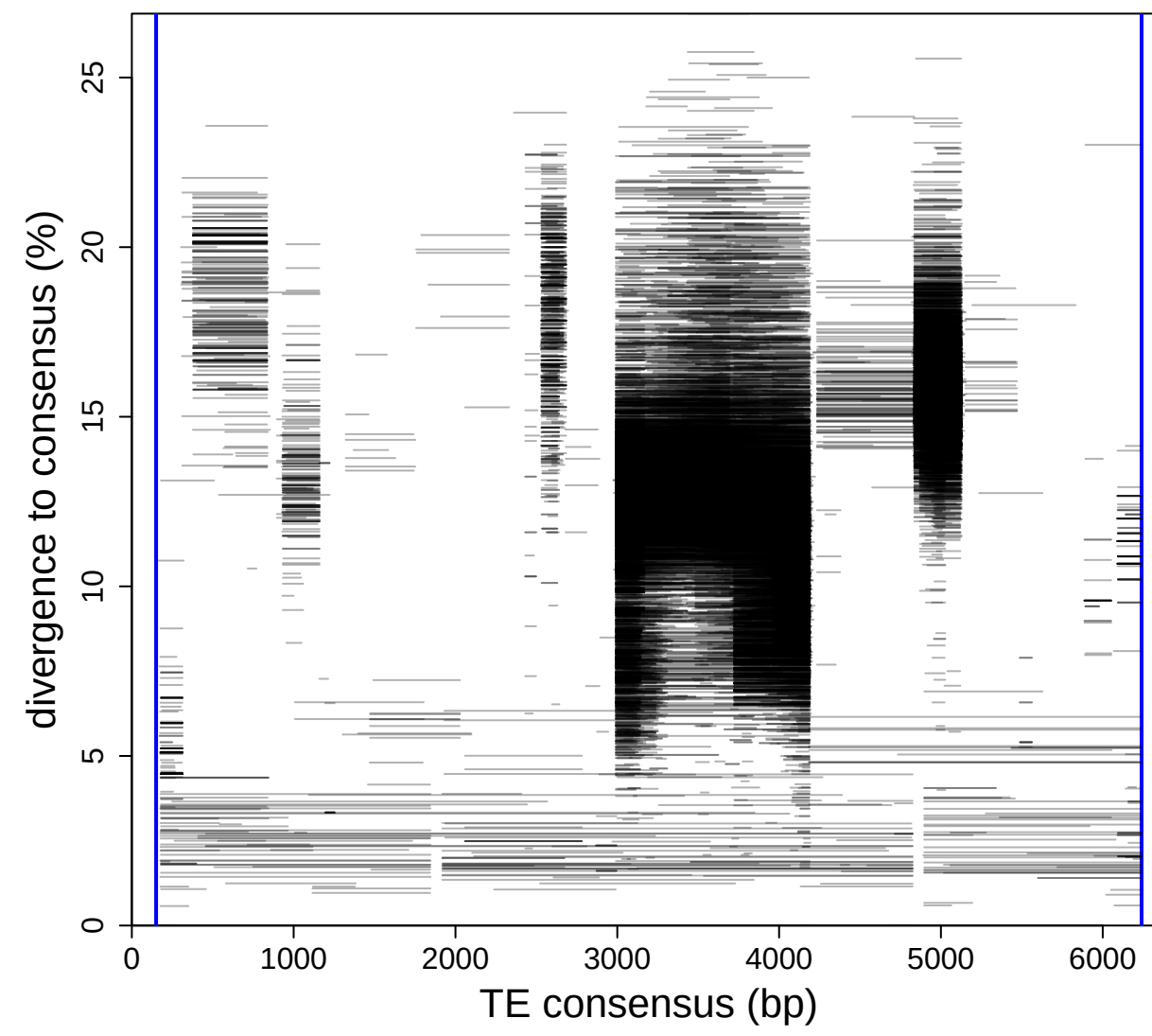
ta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.f
size: 6080bp; fragments: 15131; full length: 0 (>=5472bp)

After TEtrimmer 6080 bp

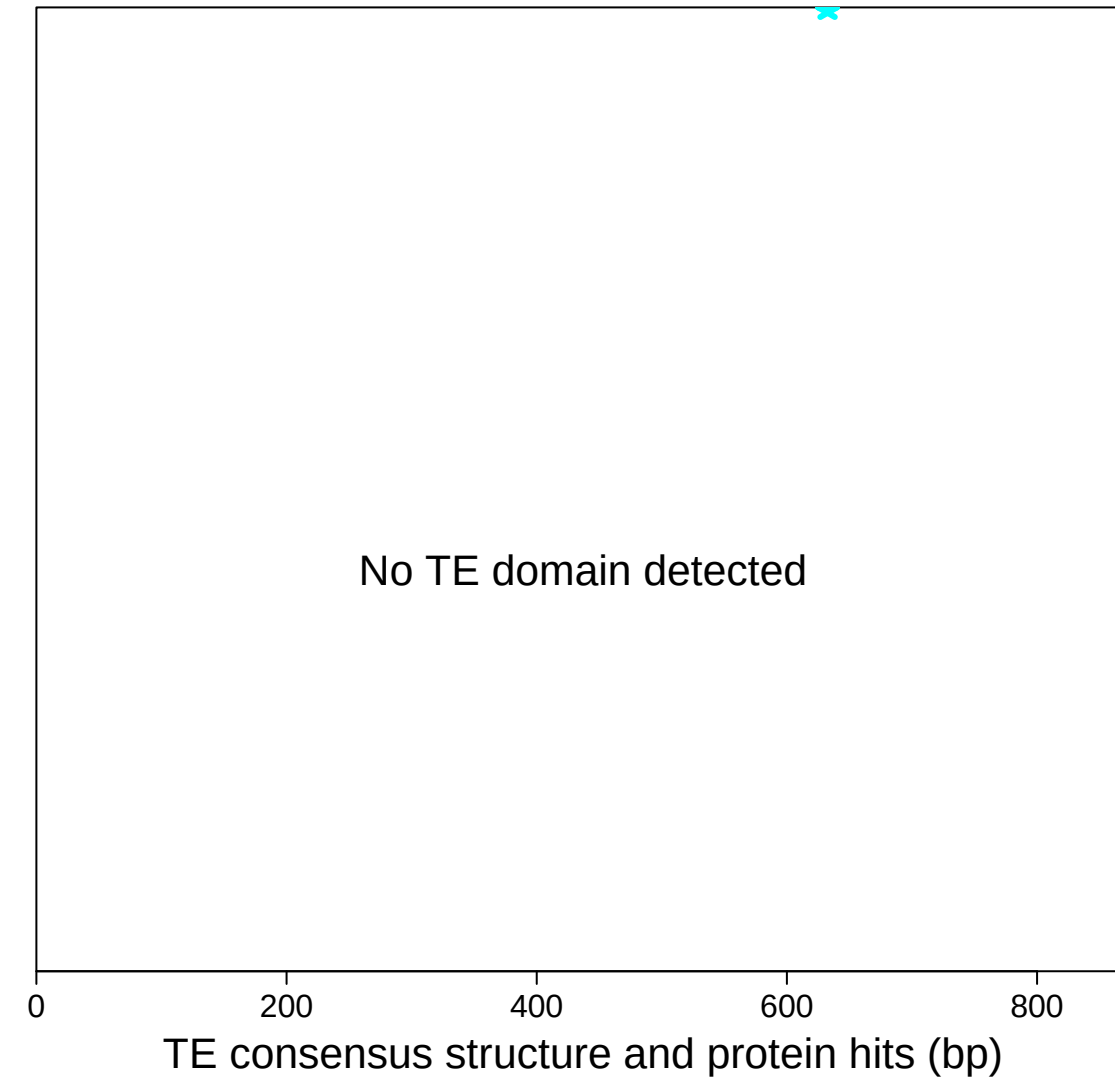
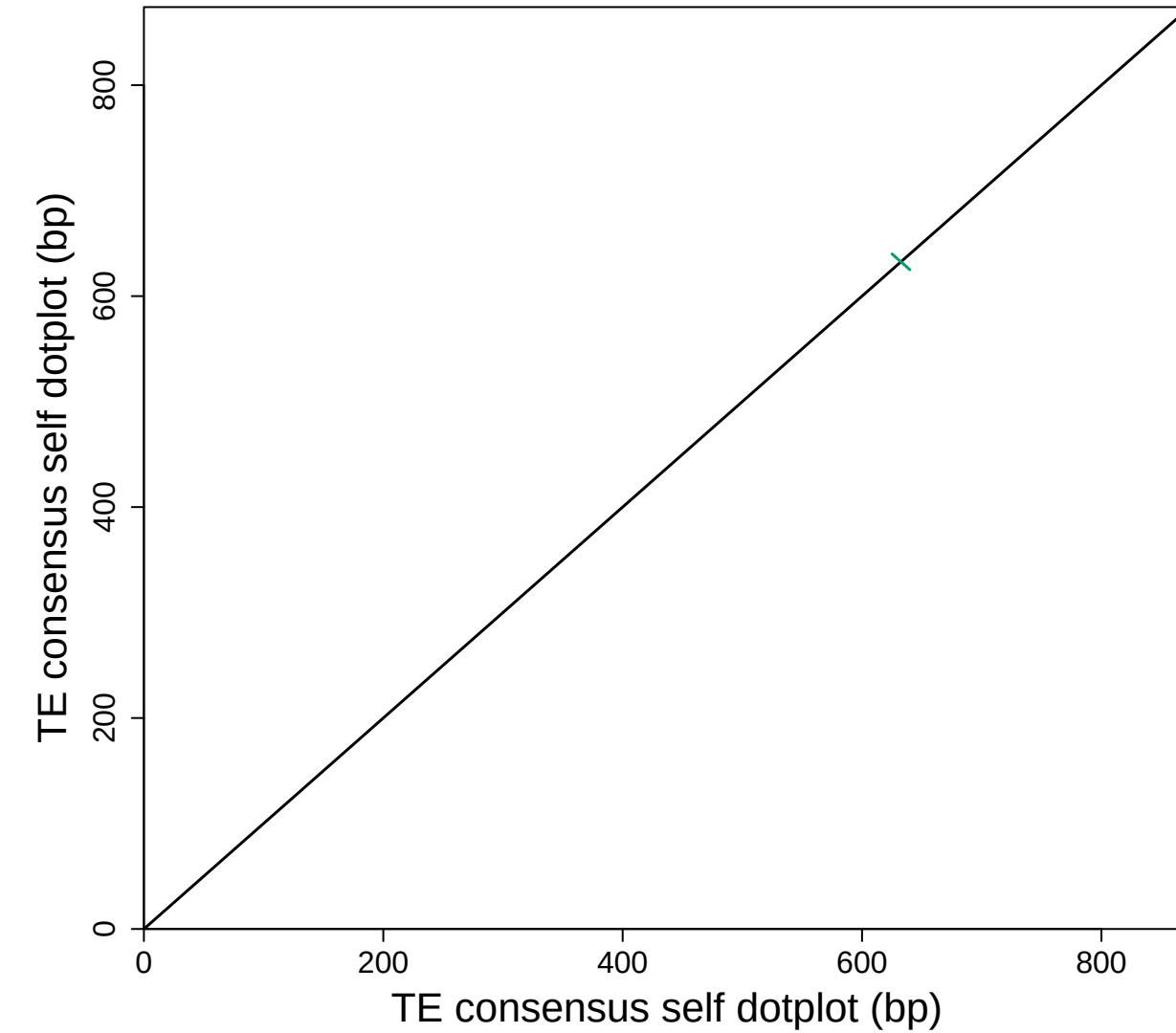
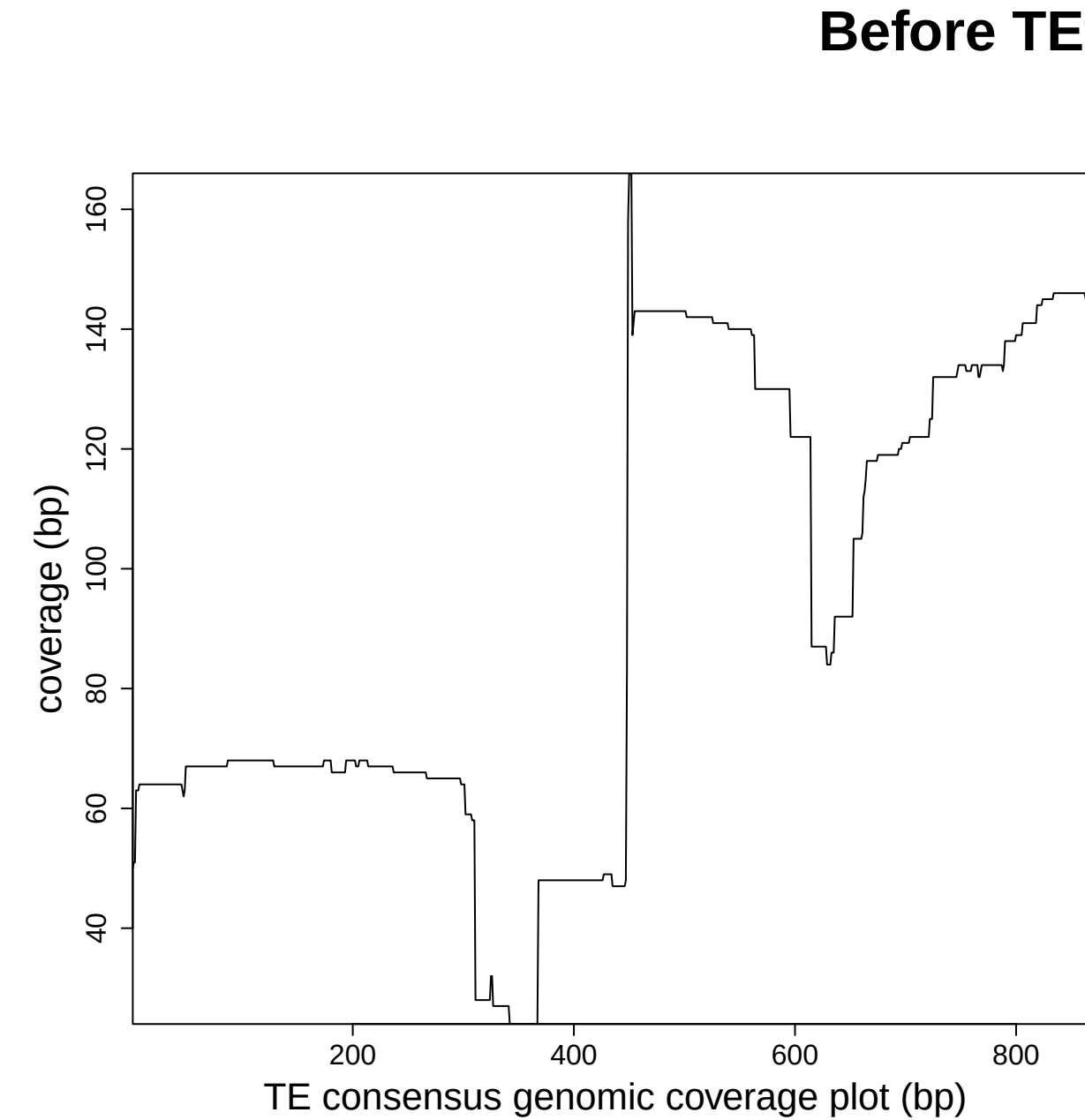
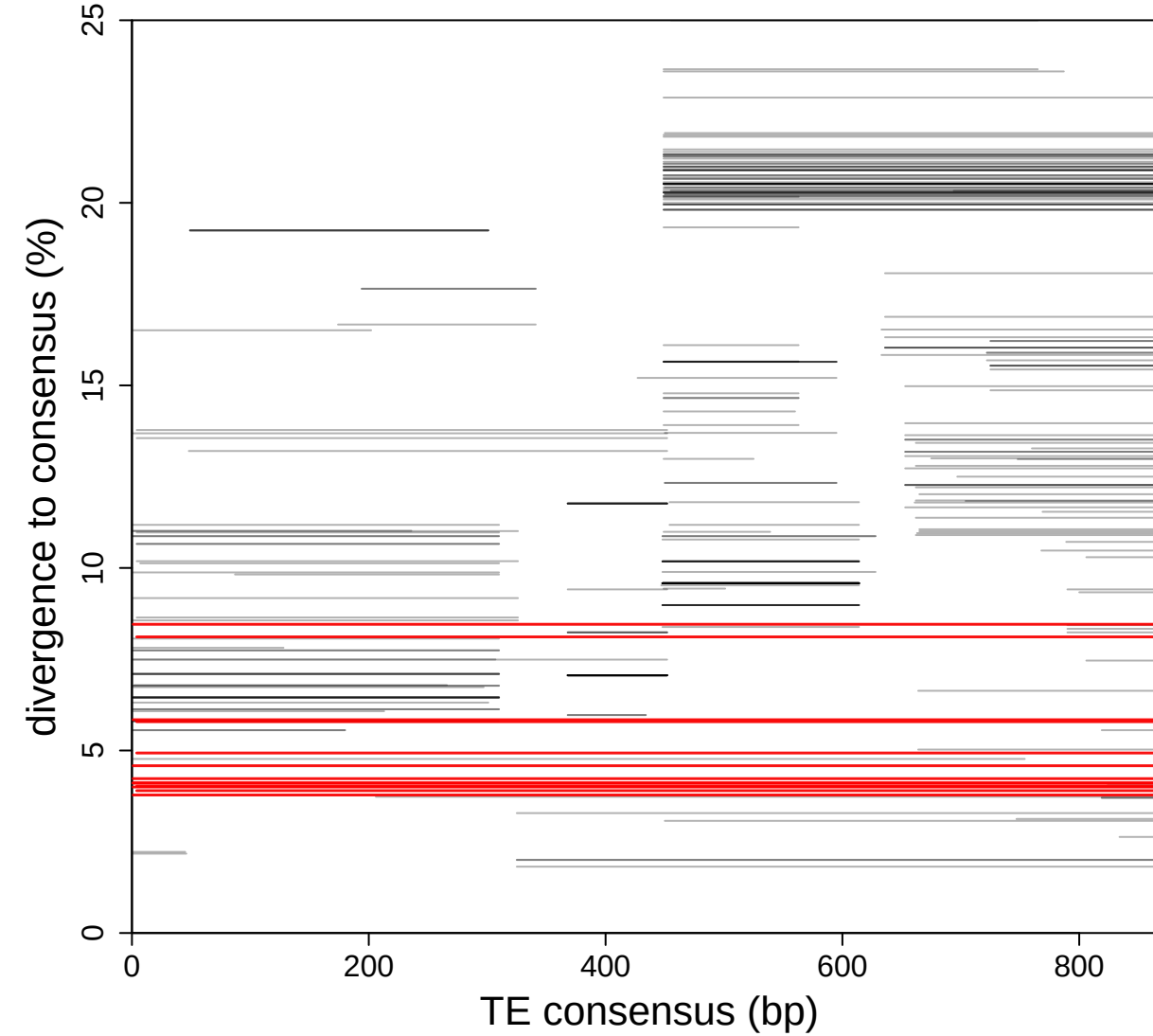


08.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.1
size: 6390bp; fragments: 15139; full length: 0 (>=5751bp)

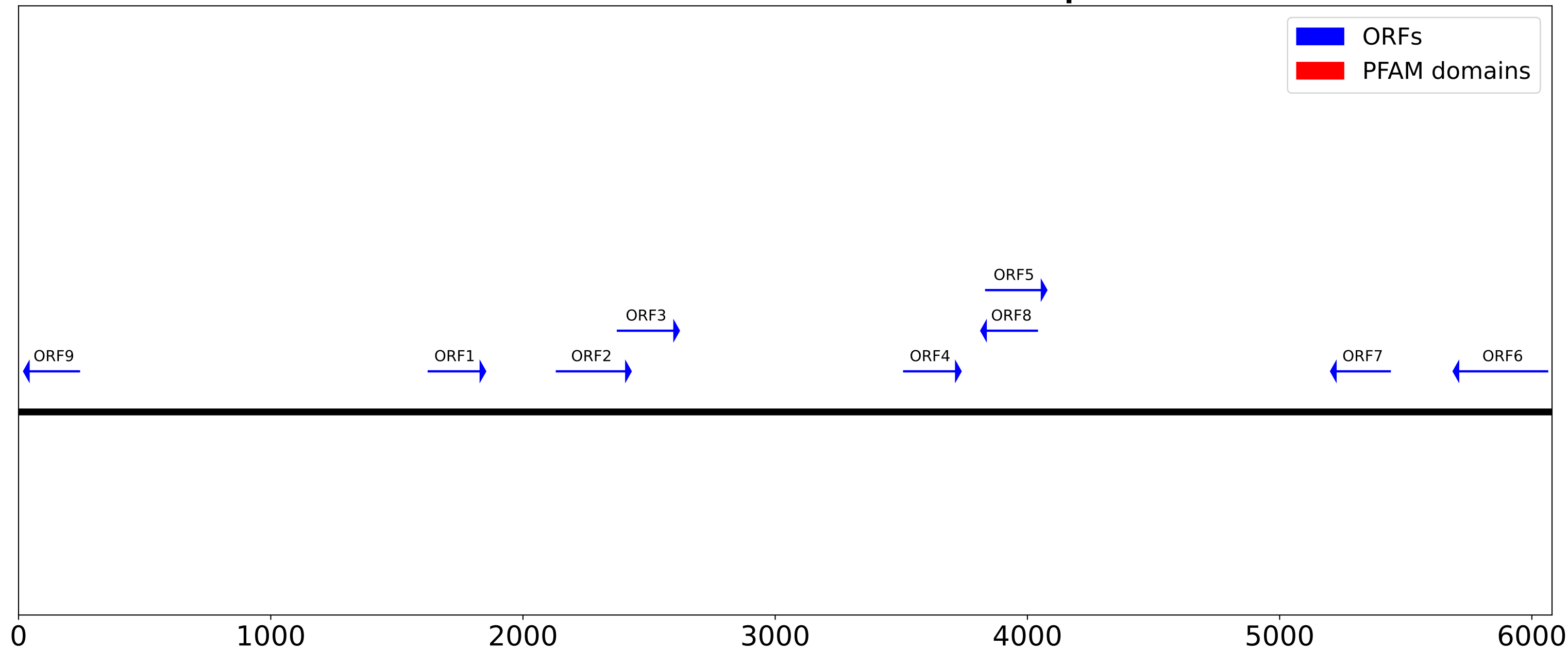
After TEtrimmer Extended plot Blue lines are boundaries



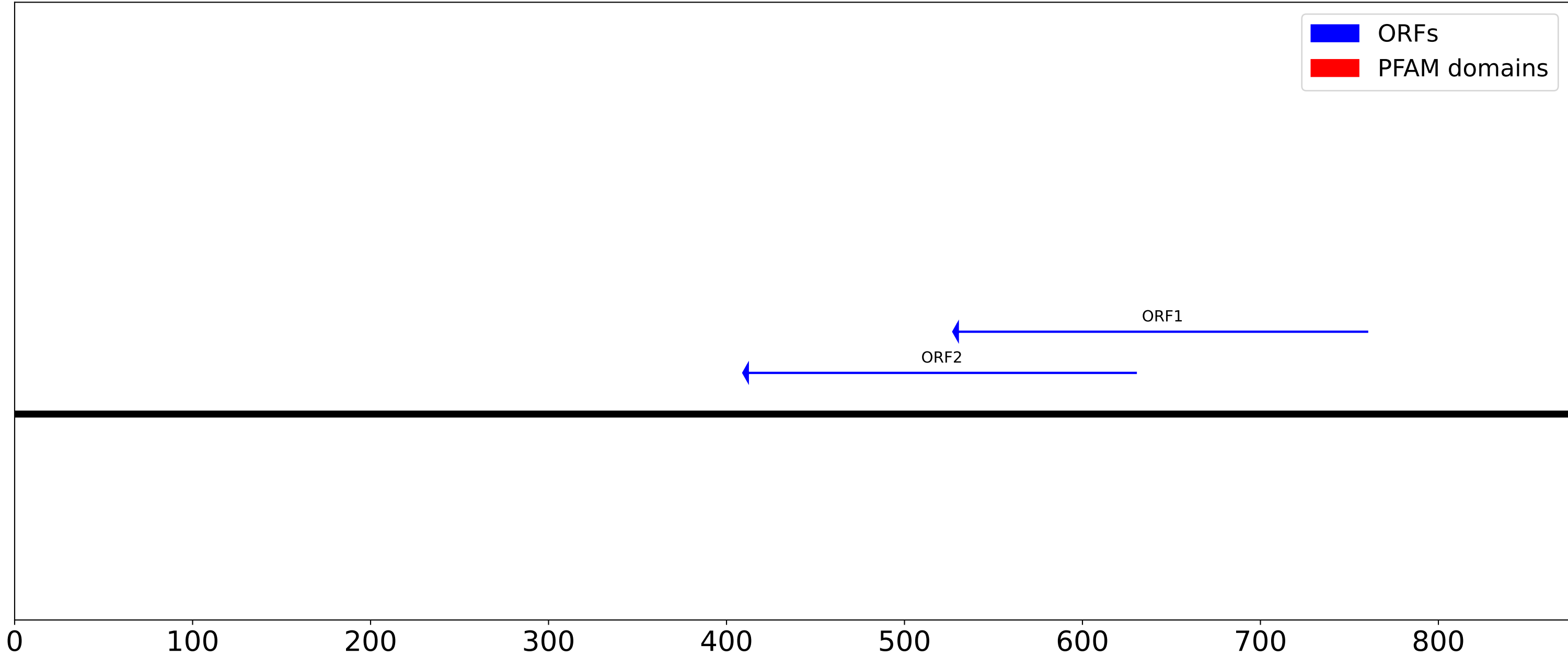
TE: rnd_5_family_11408
size: 874bp; fragments: 292; full length: 13 (≥ 786.6 bp)



After TEtrimmer ORF and PFAM domain plot



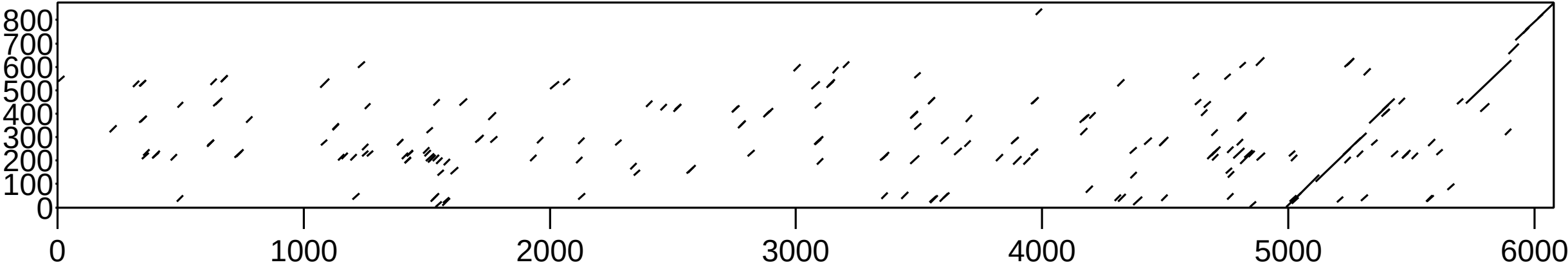
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

isensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)