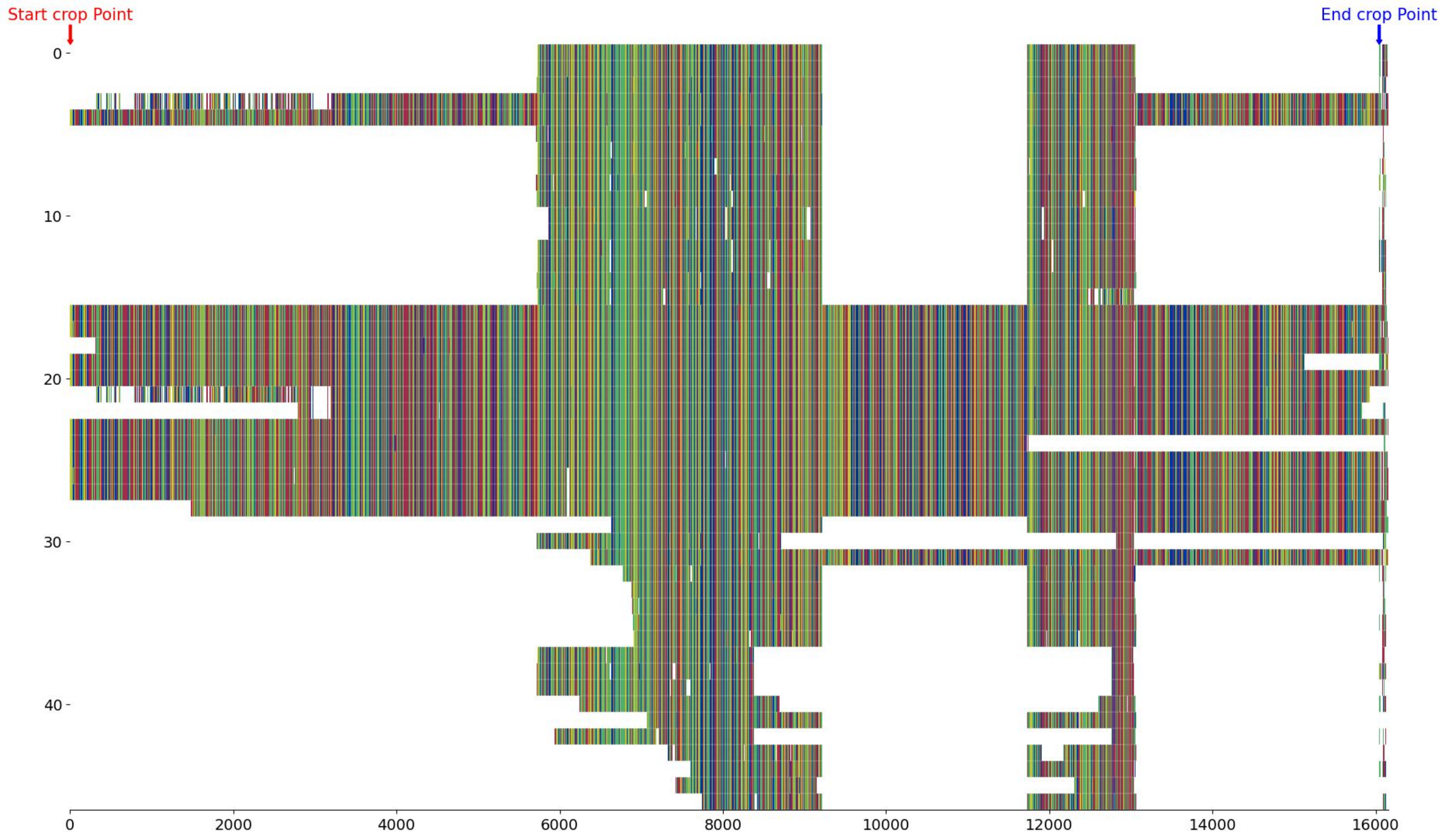


Start crop Point

End crop Point

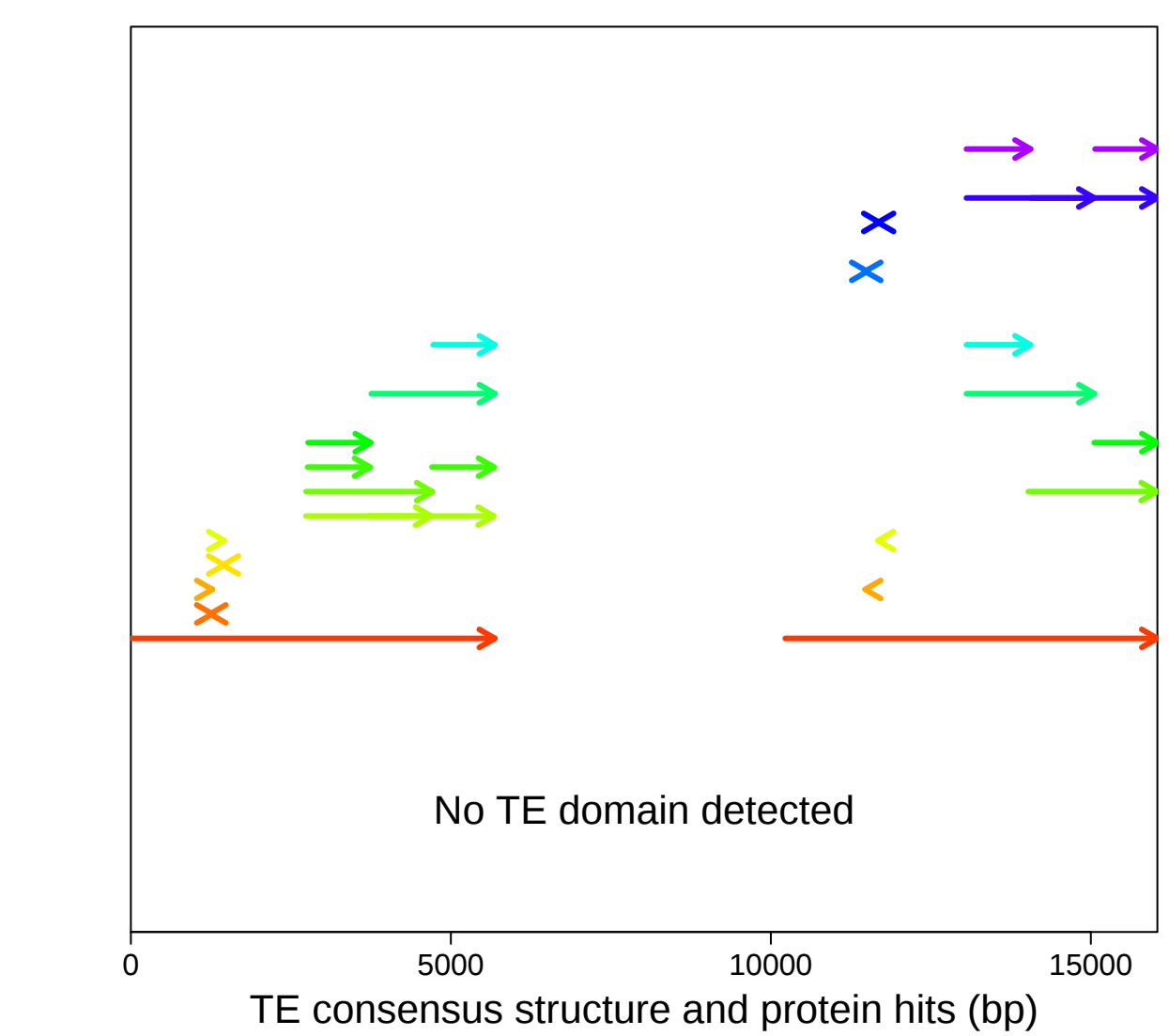
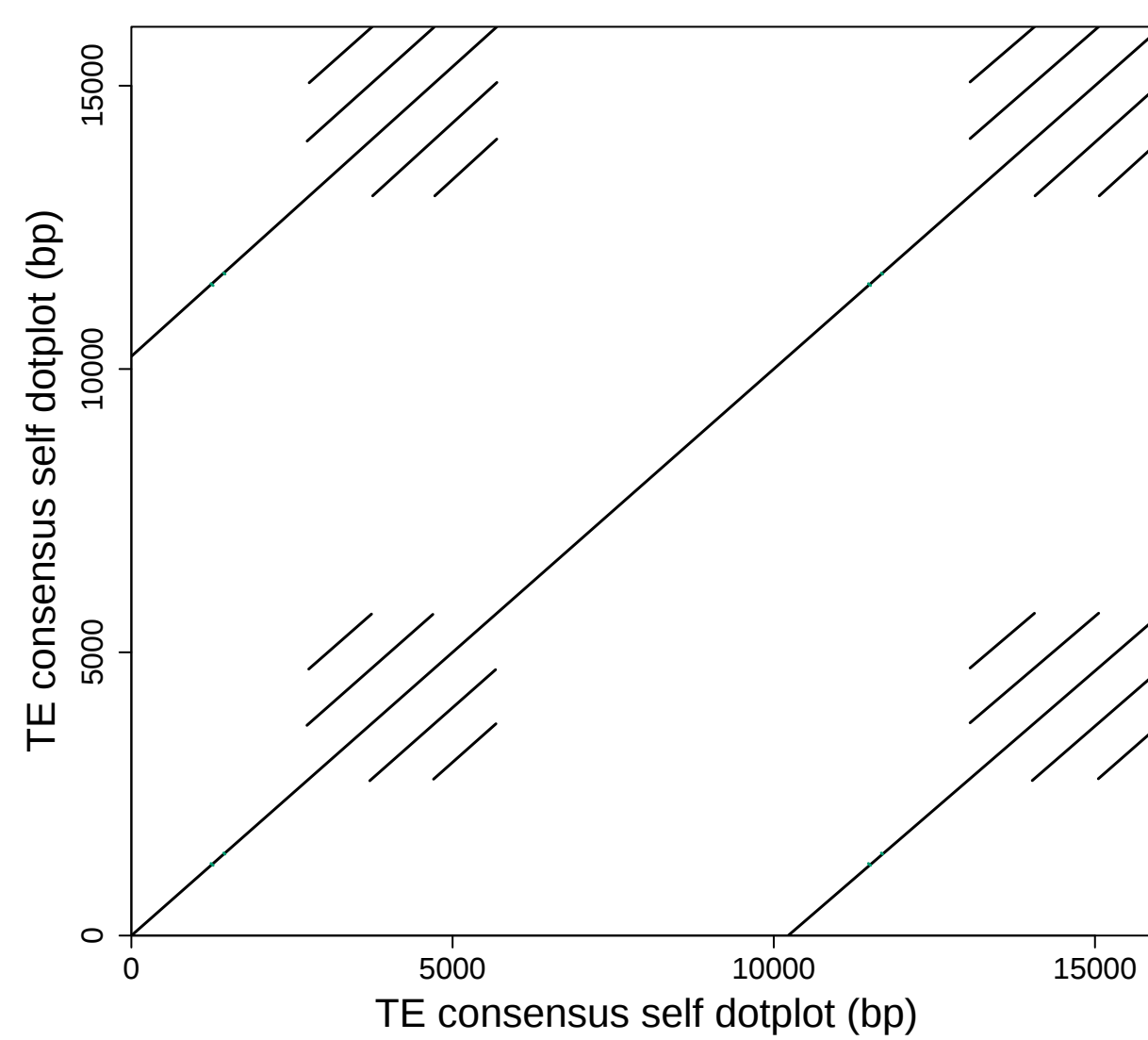
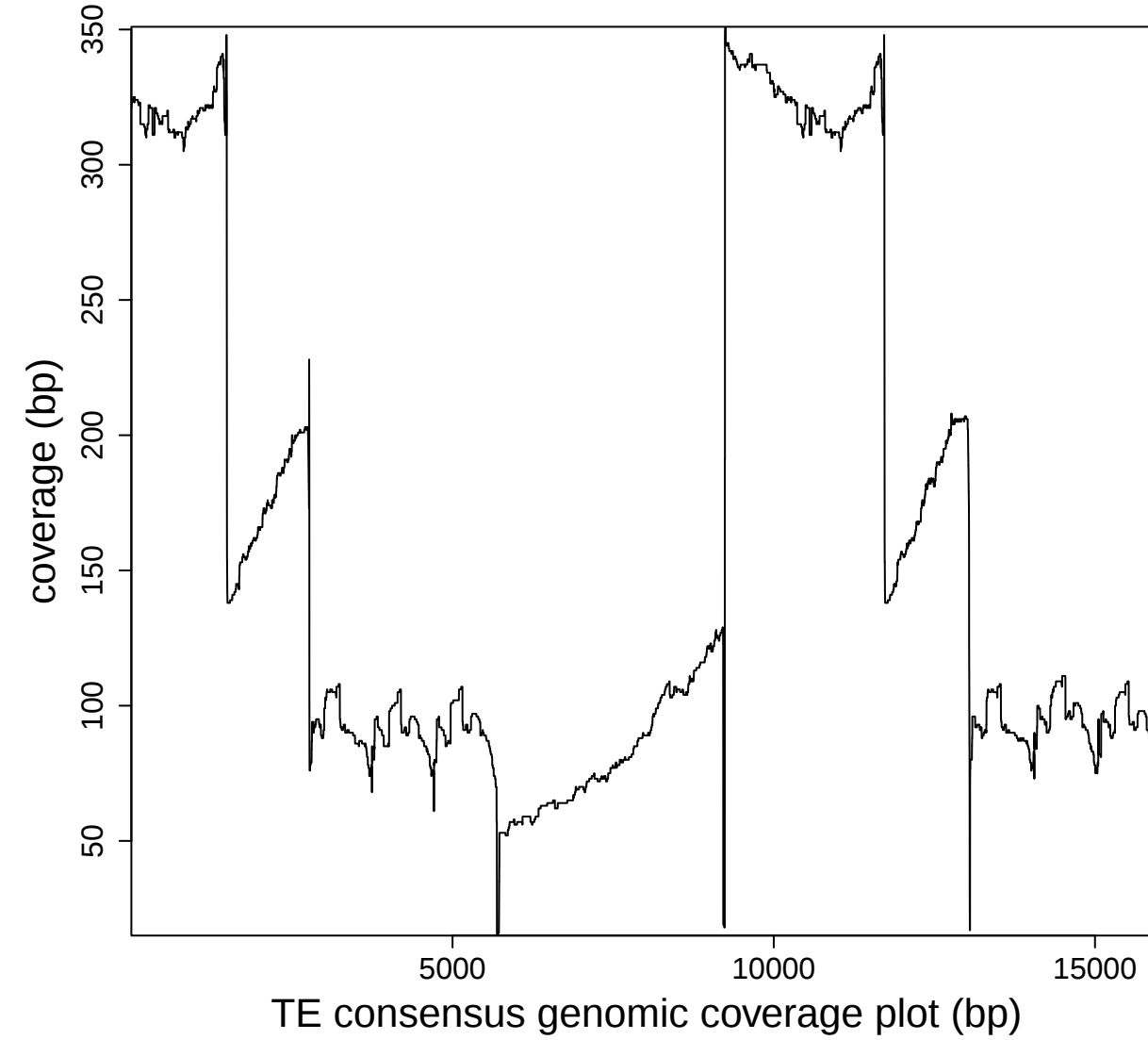
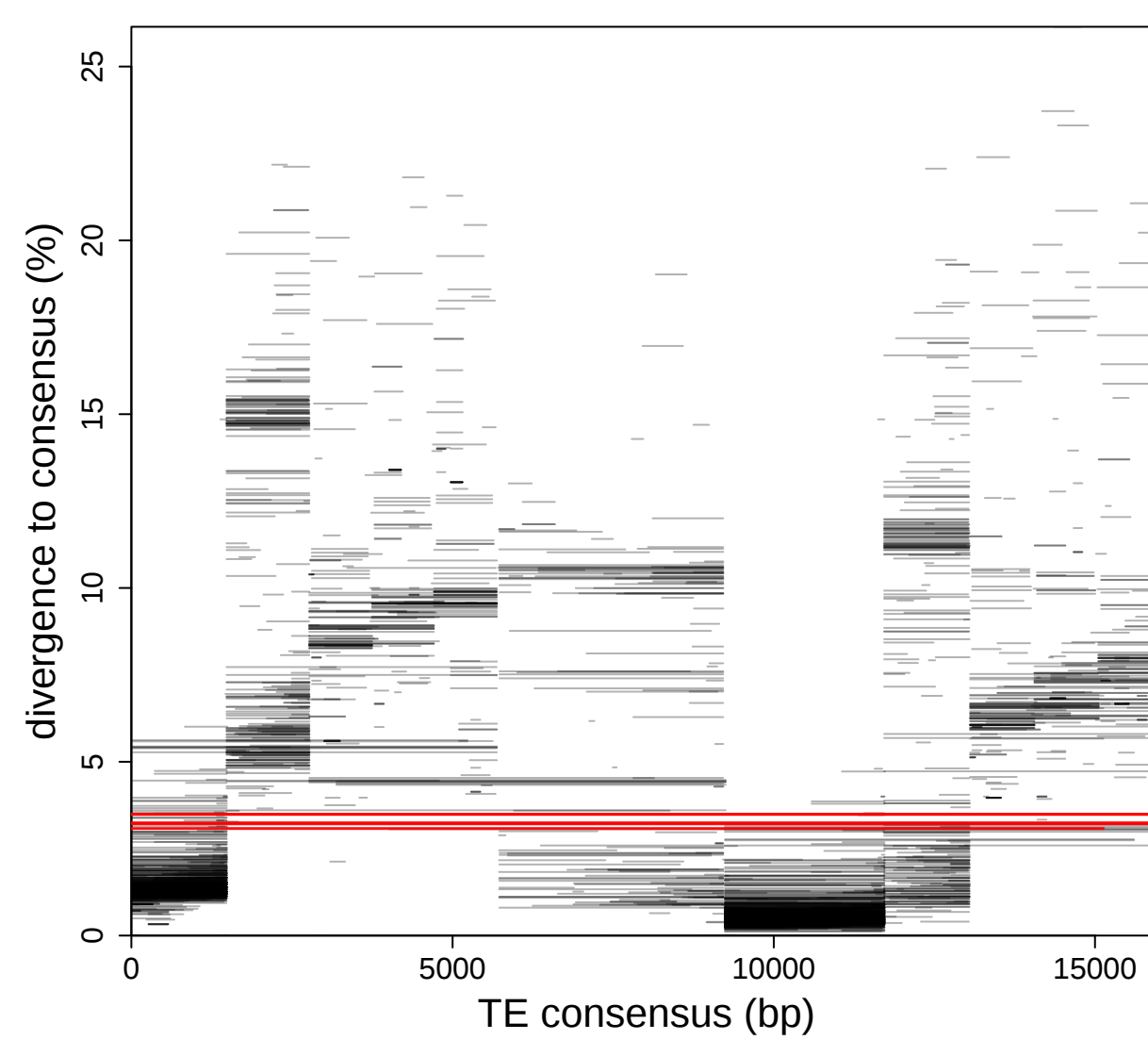
MSA length = 16040





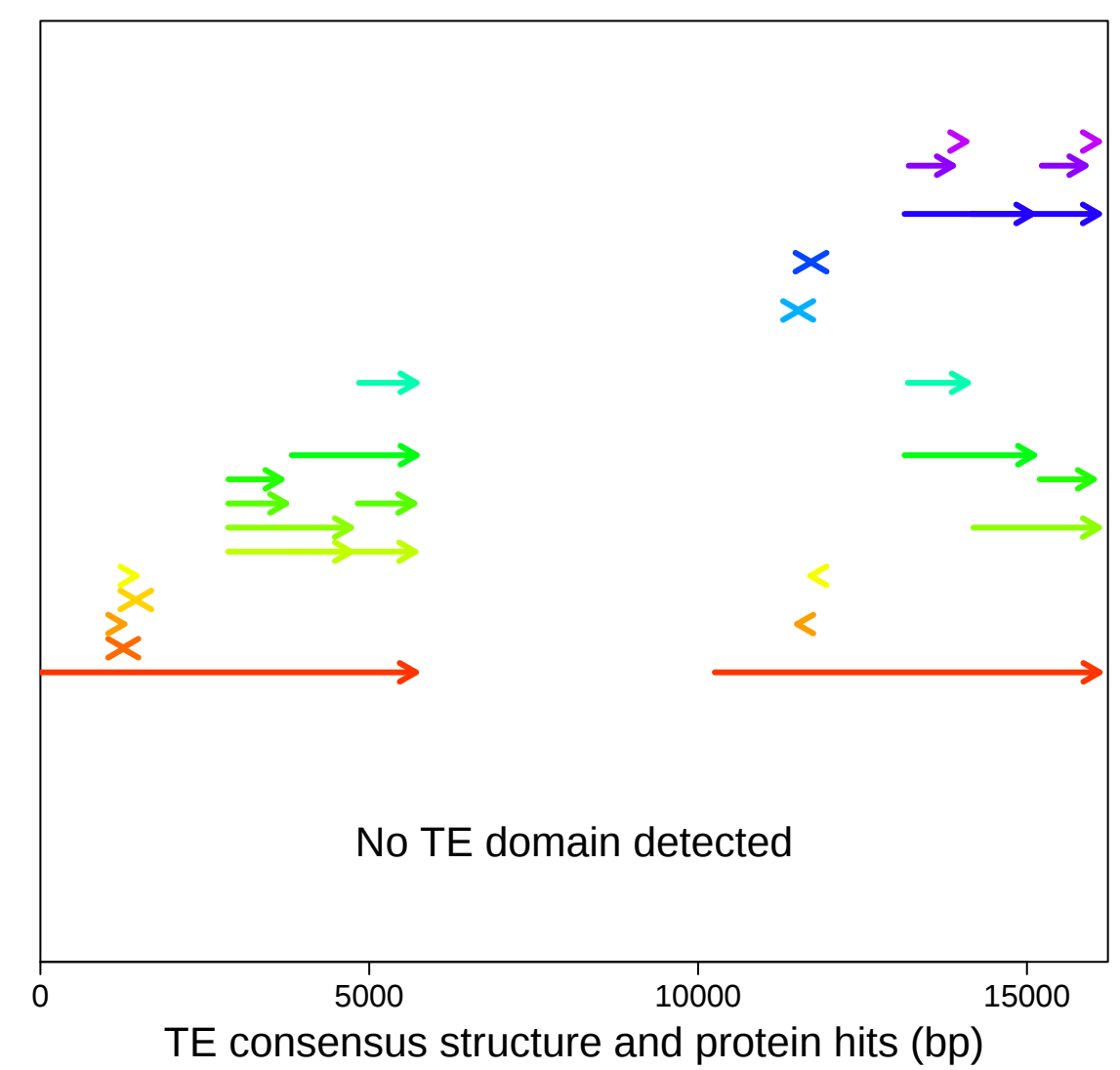
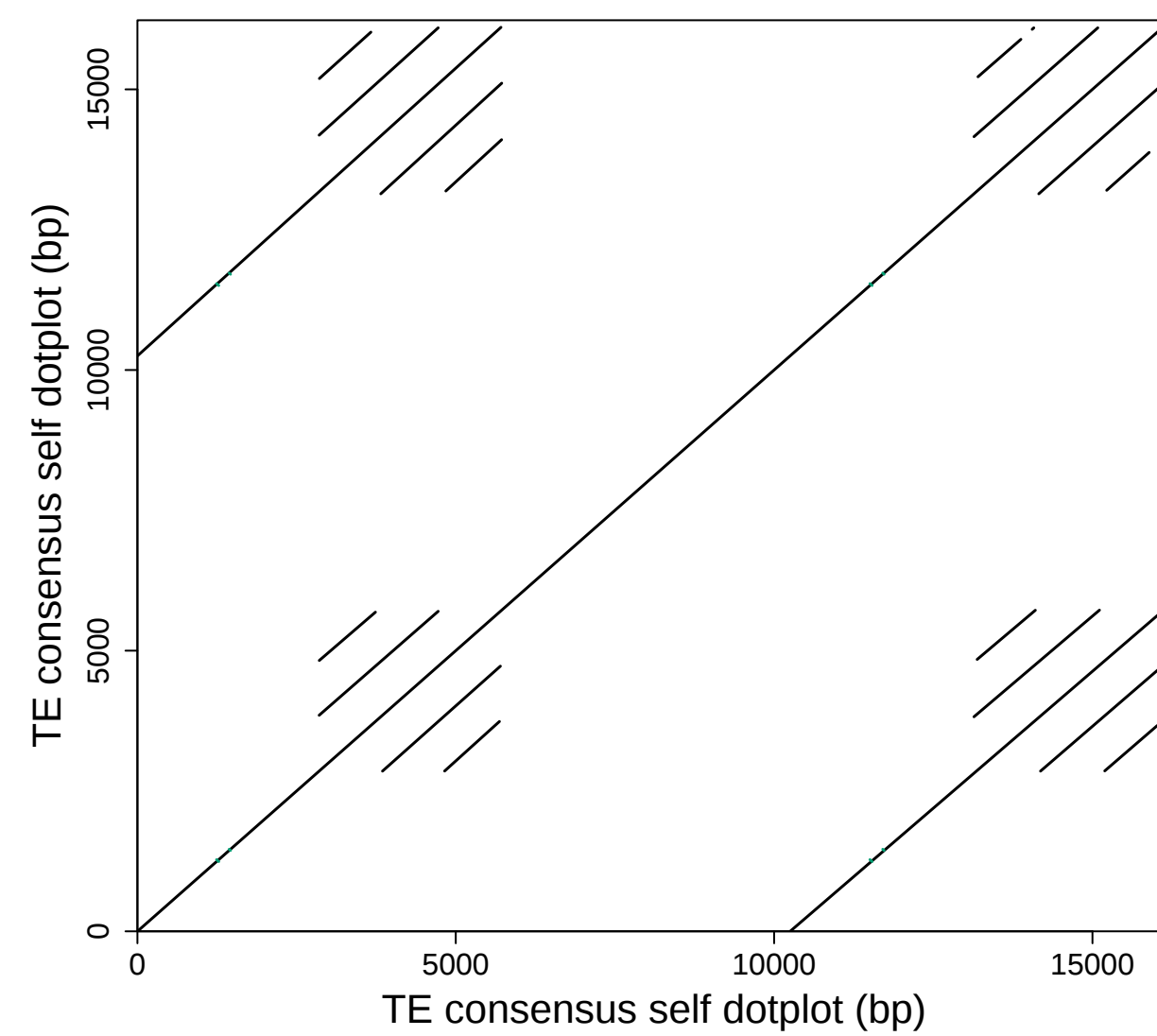
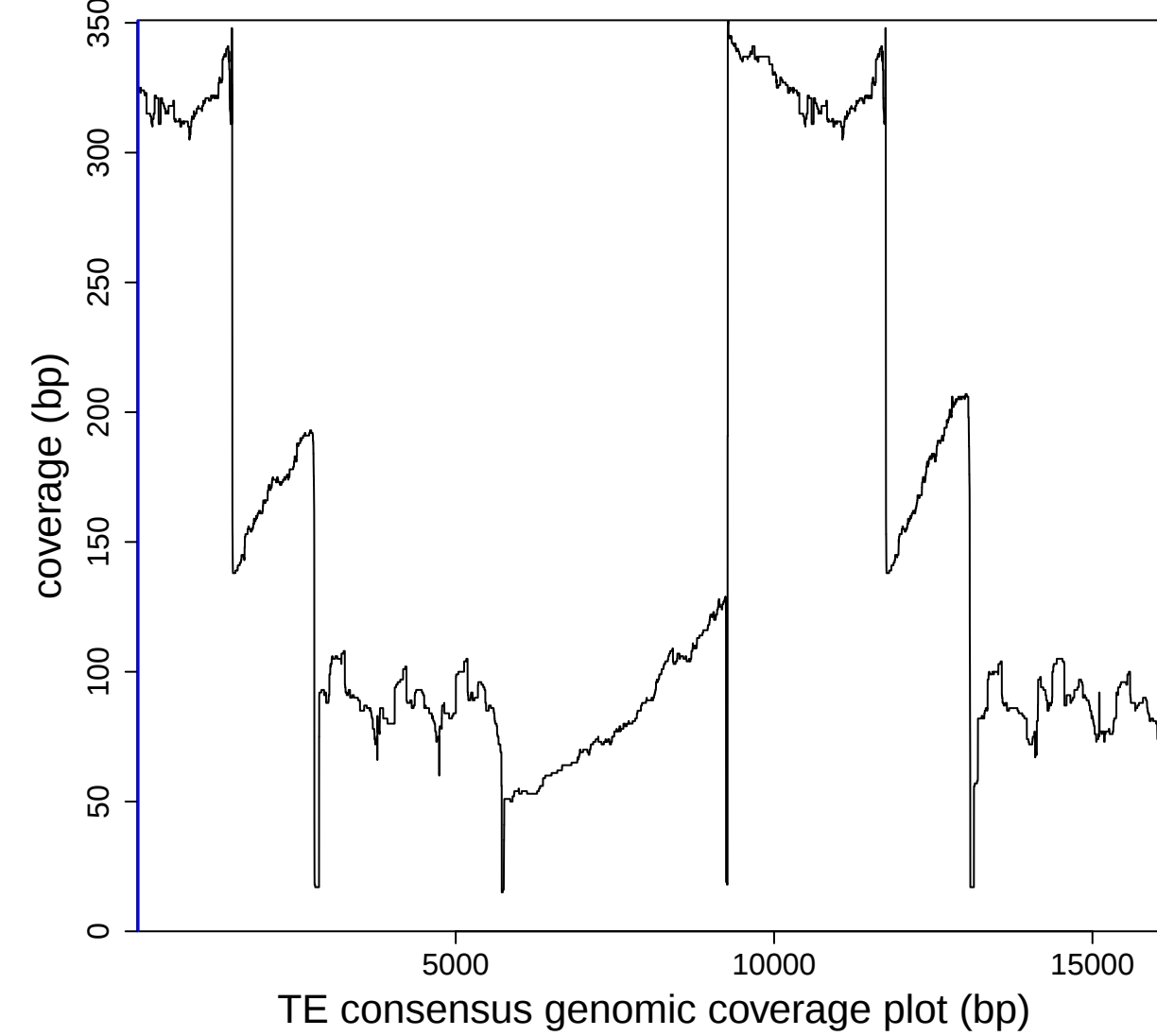
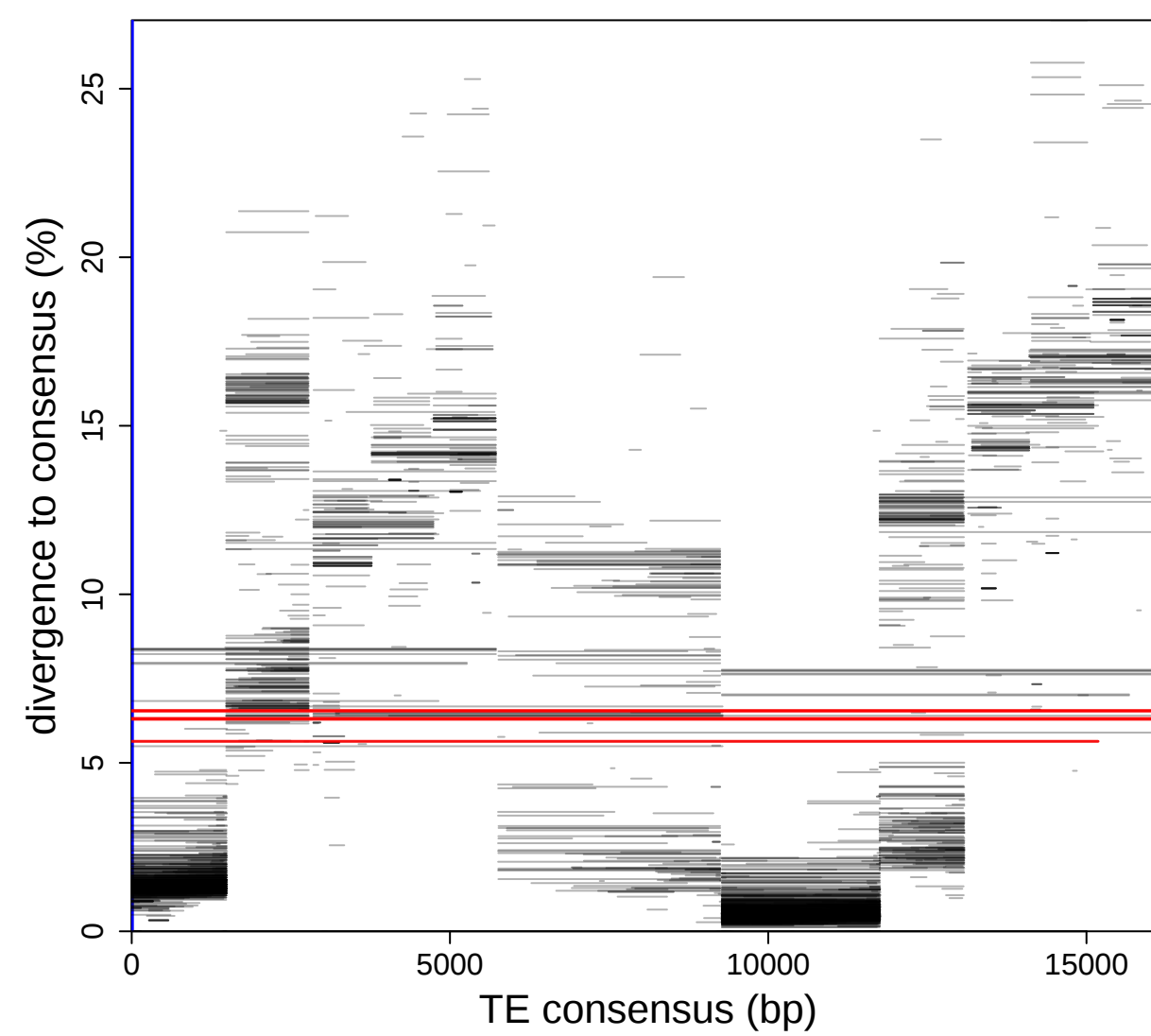
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.fa_gs.fa_ce
size: 16040bp; fragments: 2058; full length: 5 (>=14436bp)

After TEtrimmer 16040 bp



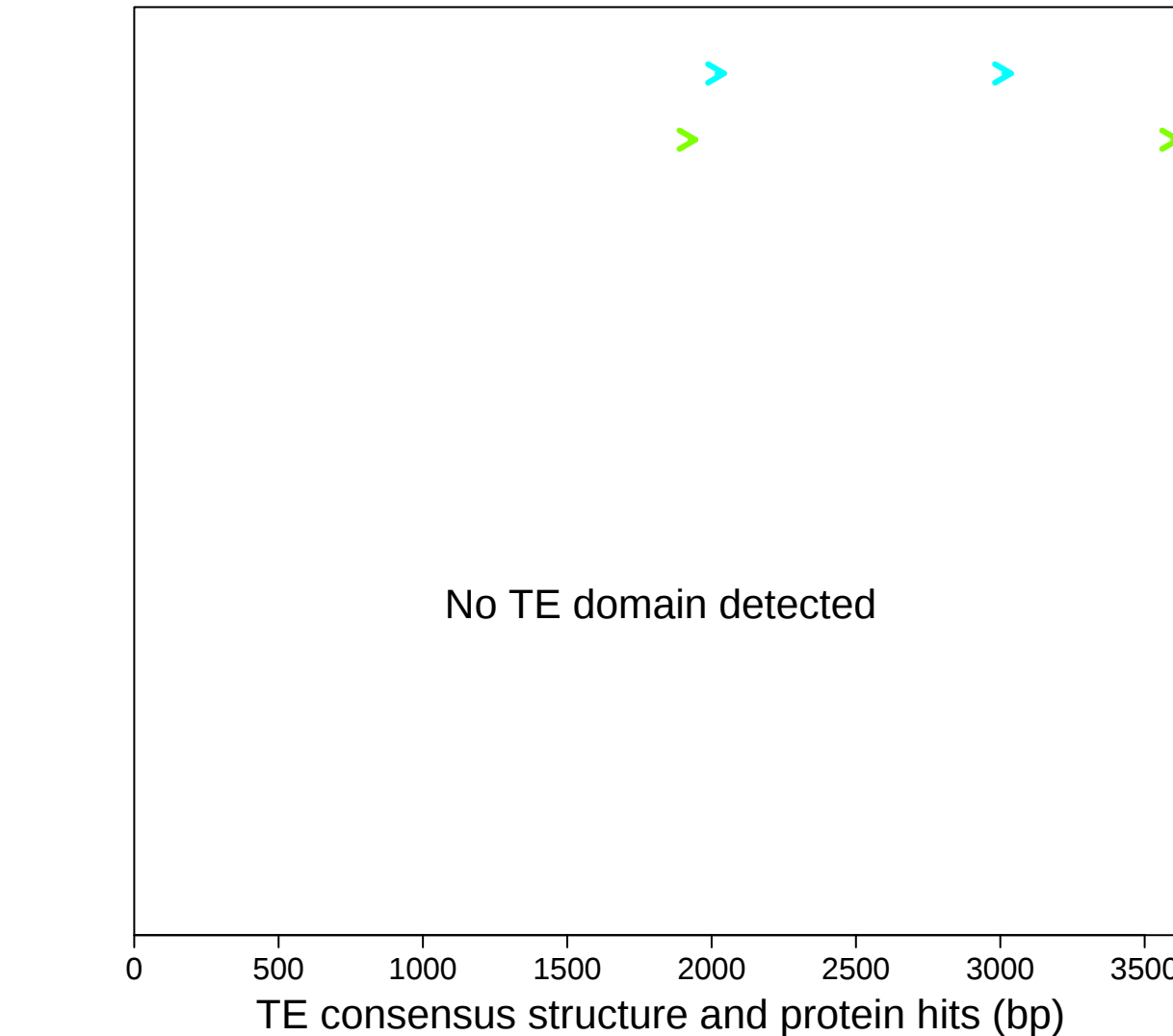
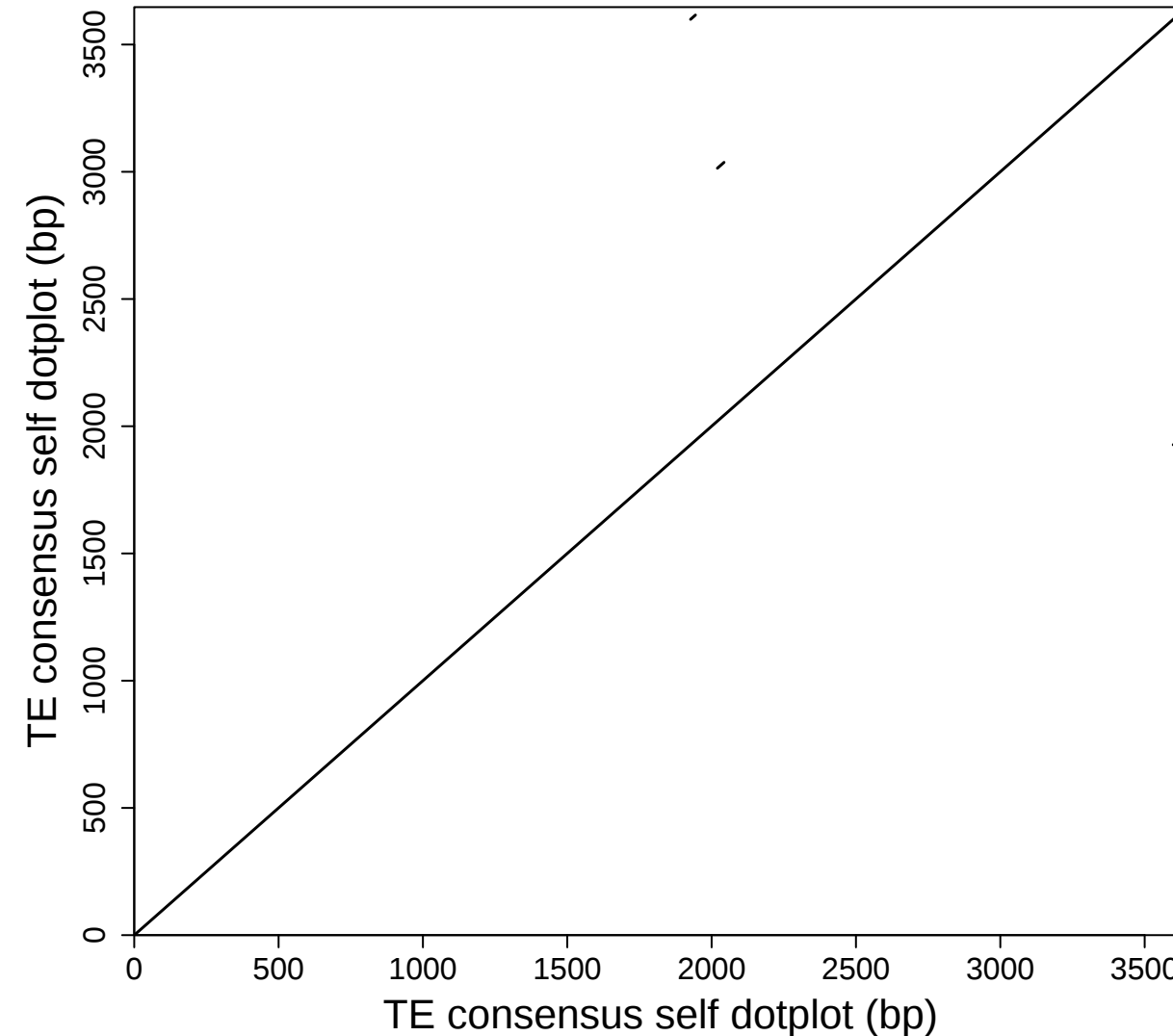
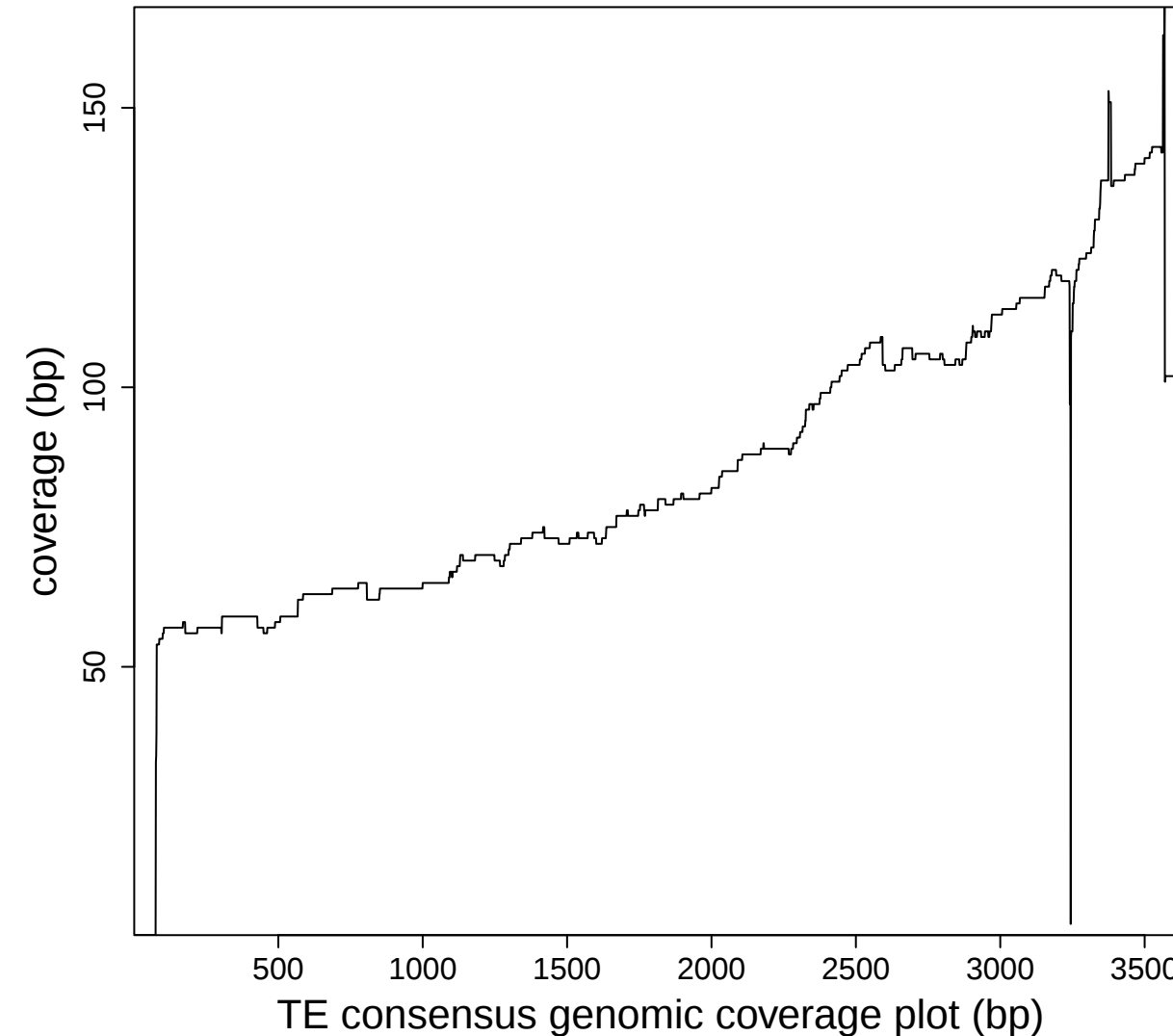
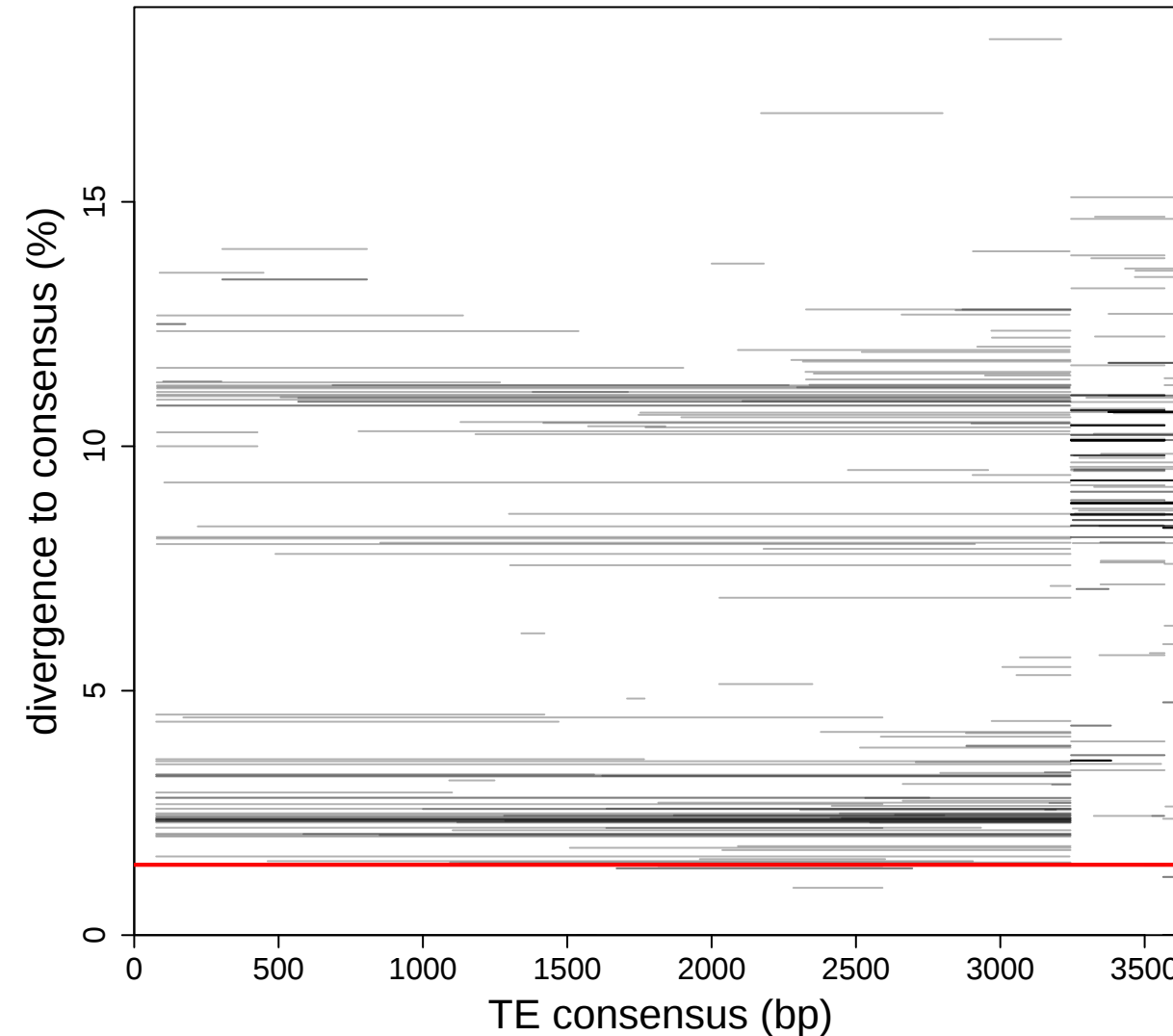
ily_956.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 16231bp; fragments: 1968; full length: 5 (>=14607.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

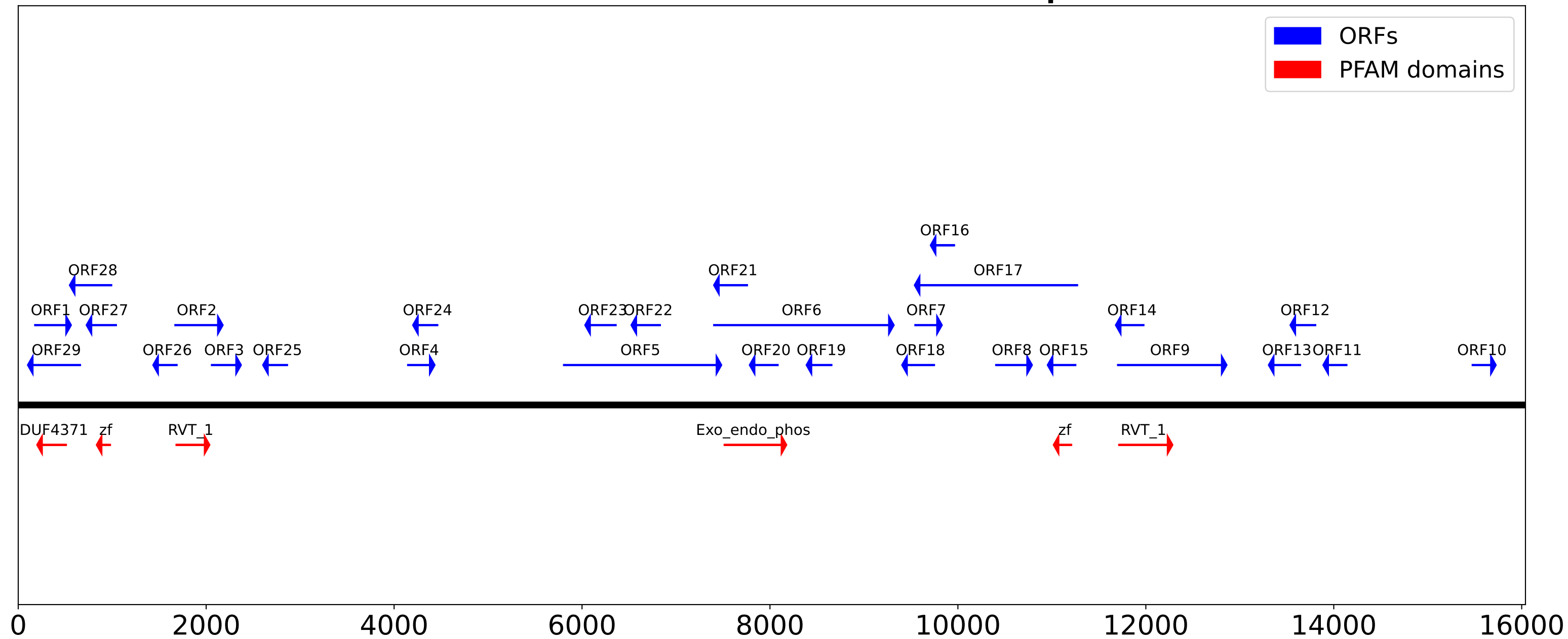


TE: rnd_5_family_956
size: 3647bp; fragments: 351; full length: 2 (≥ 3282.3 bp)

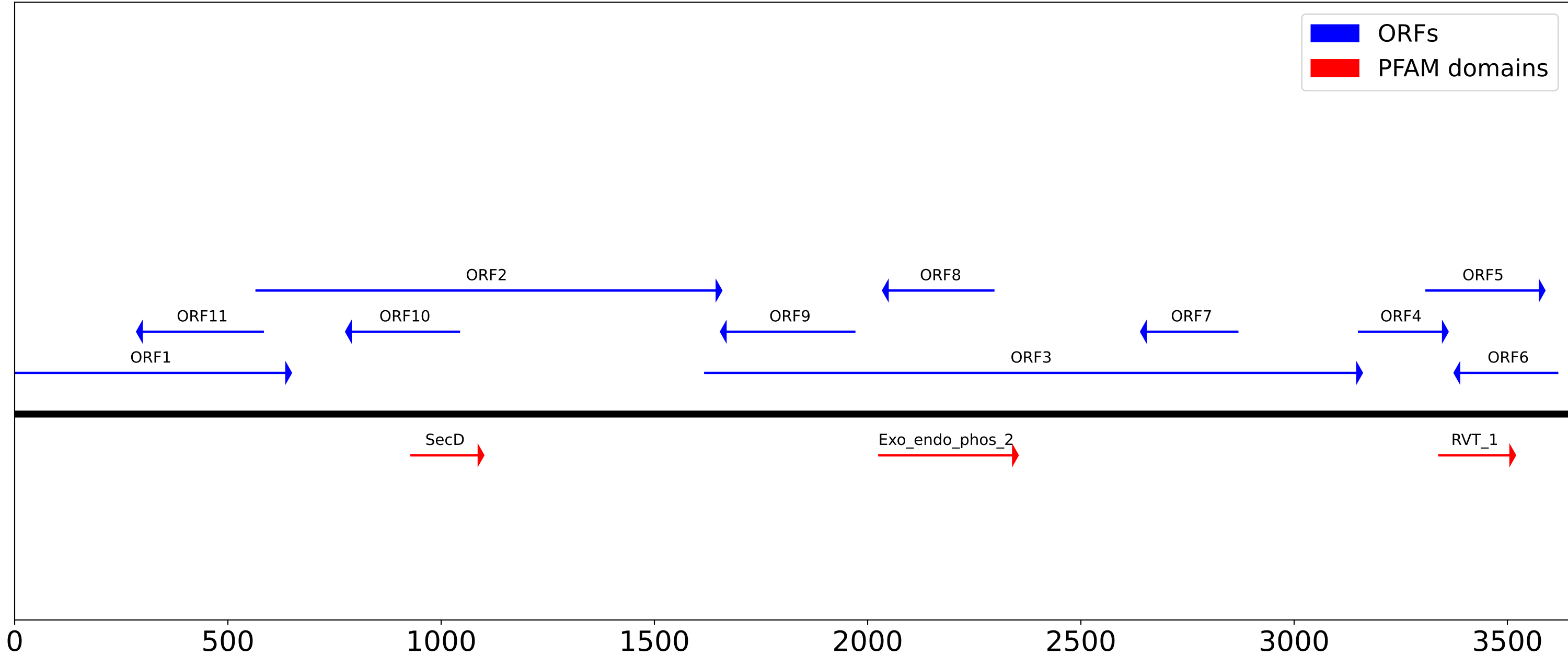
Before TEtrimmer 3647 bp



After TEtrimmer ORF and PFAM domain plot



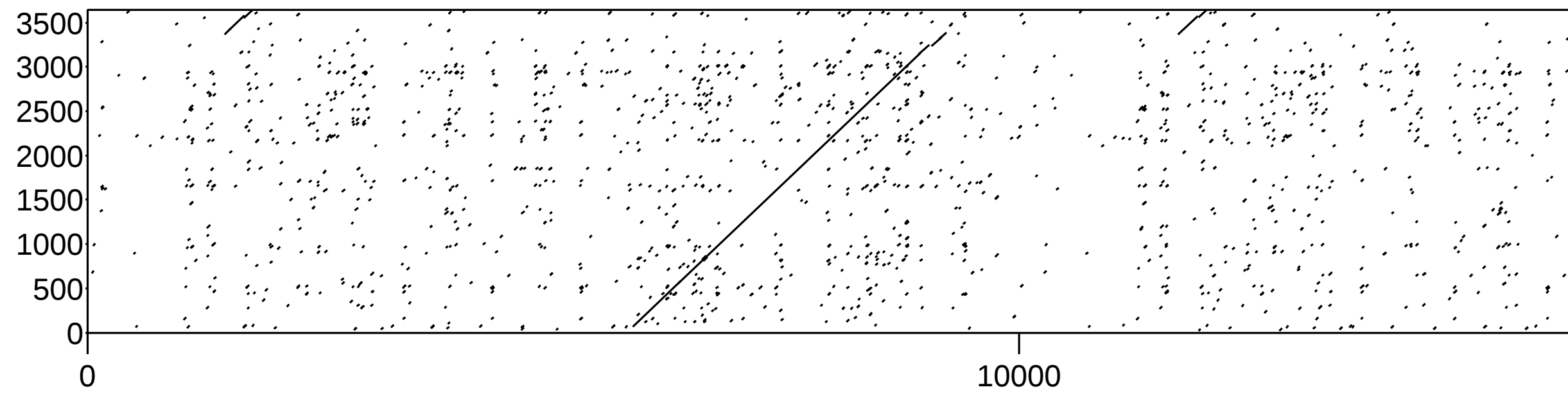
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)