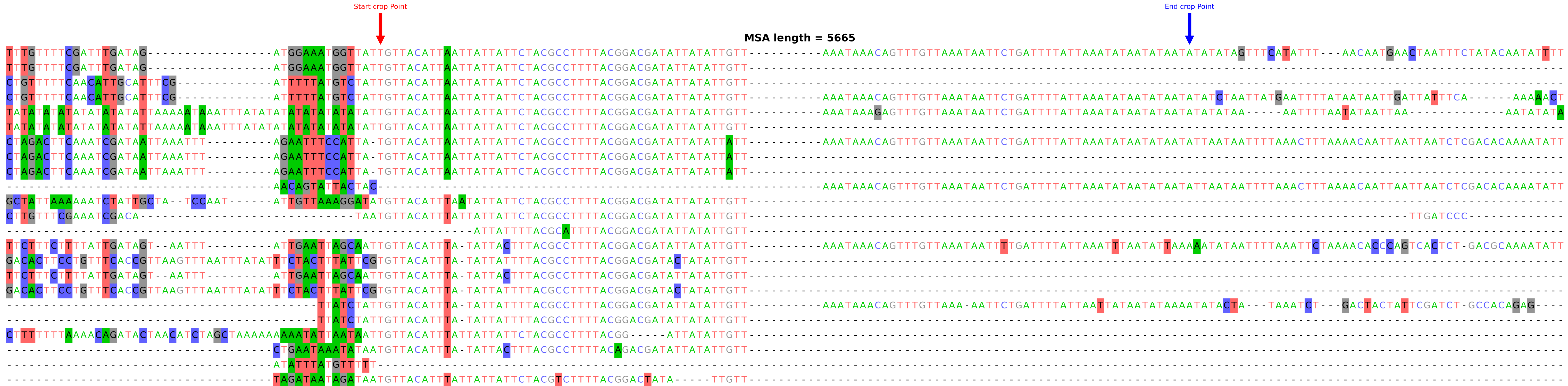
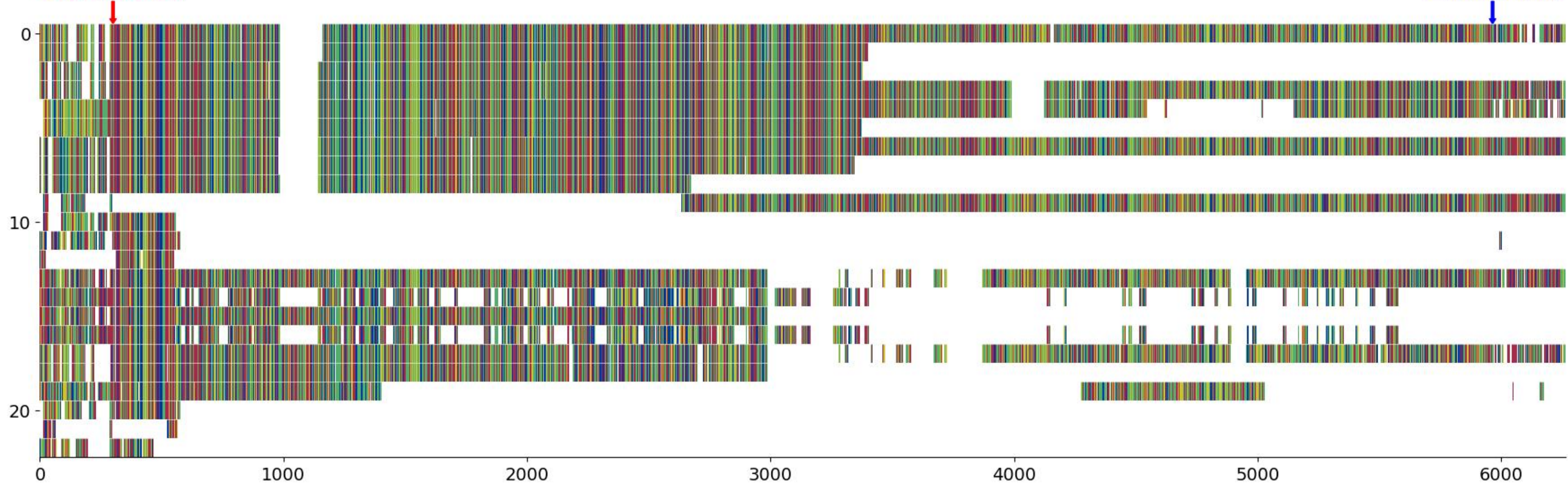




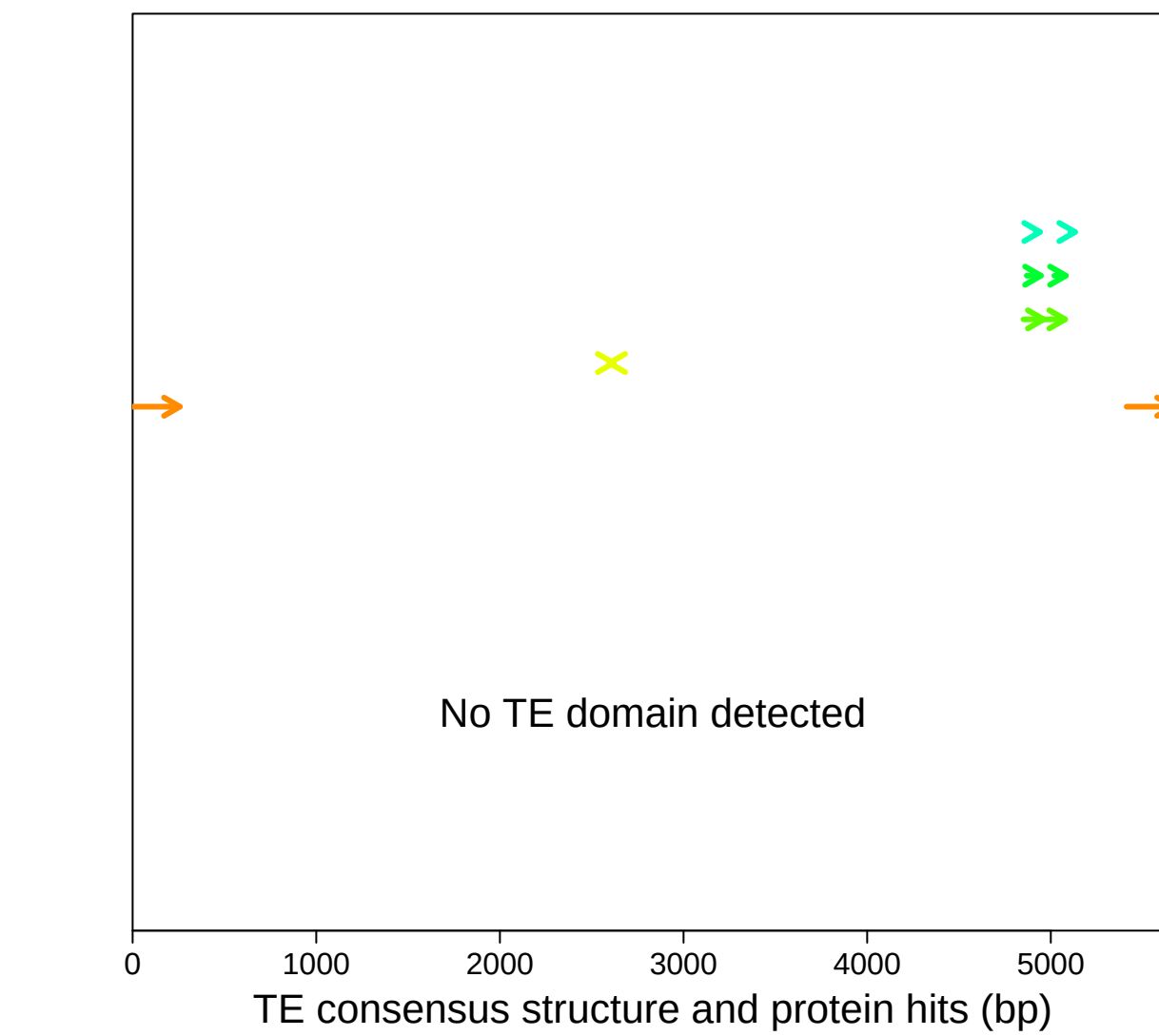
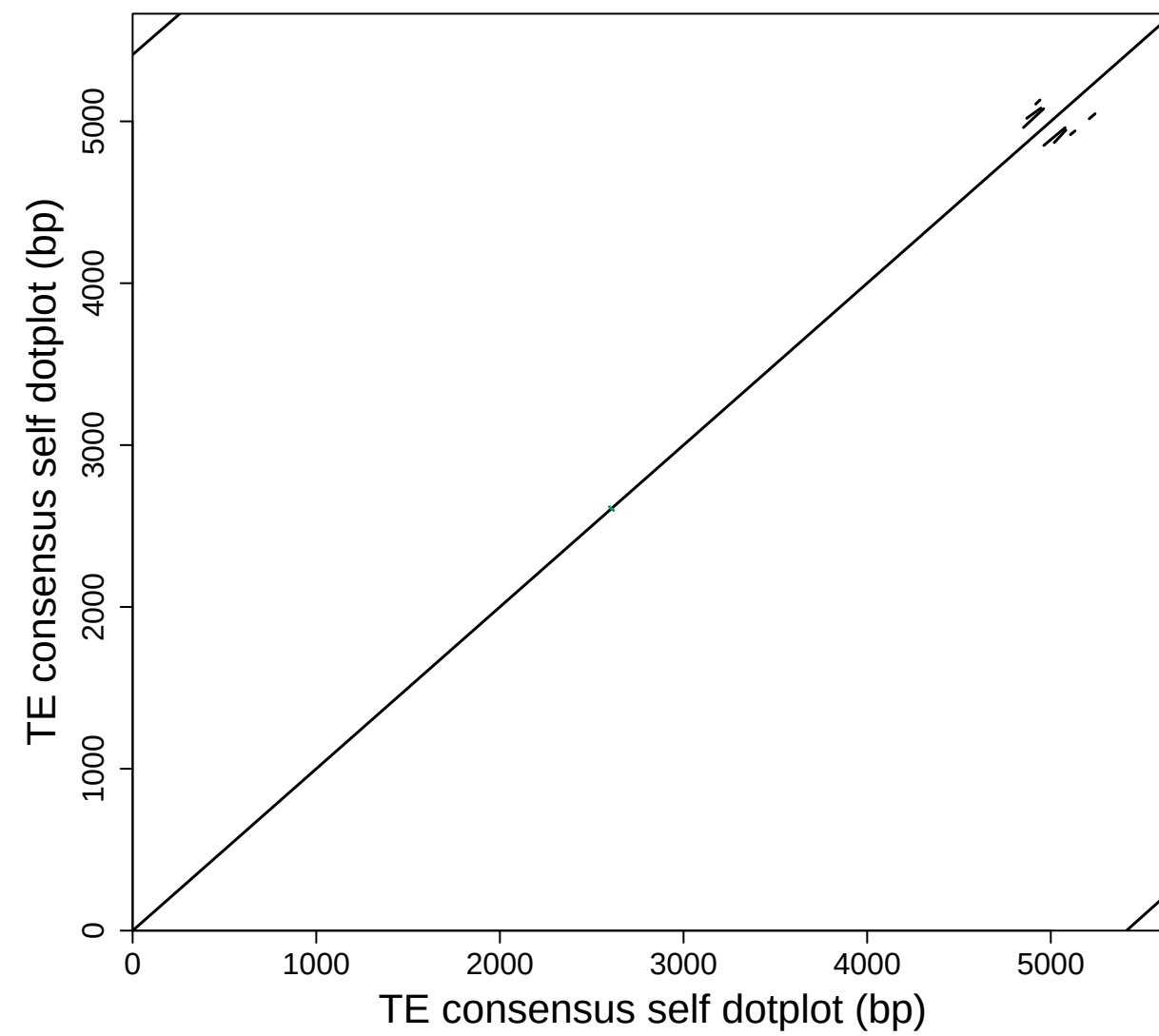
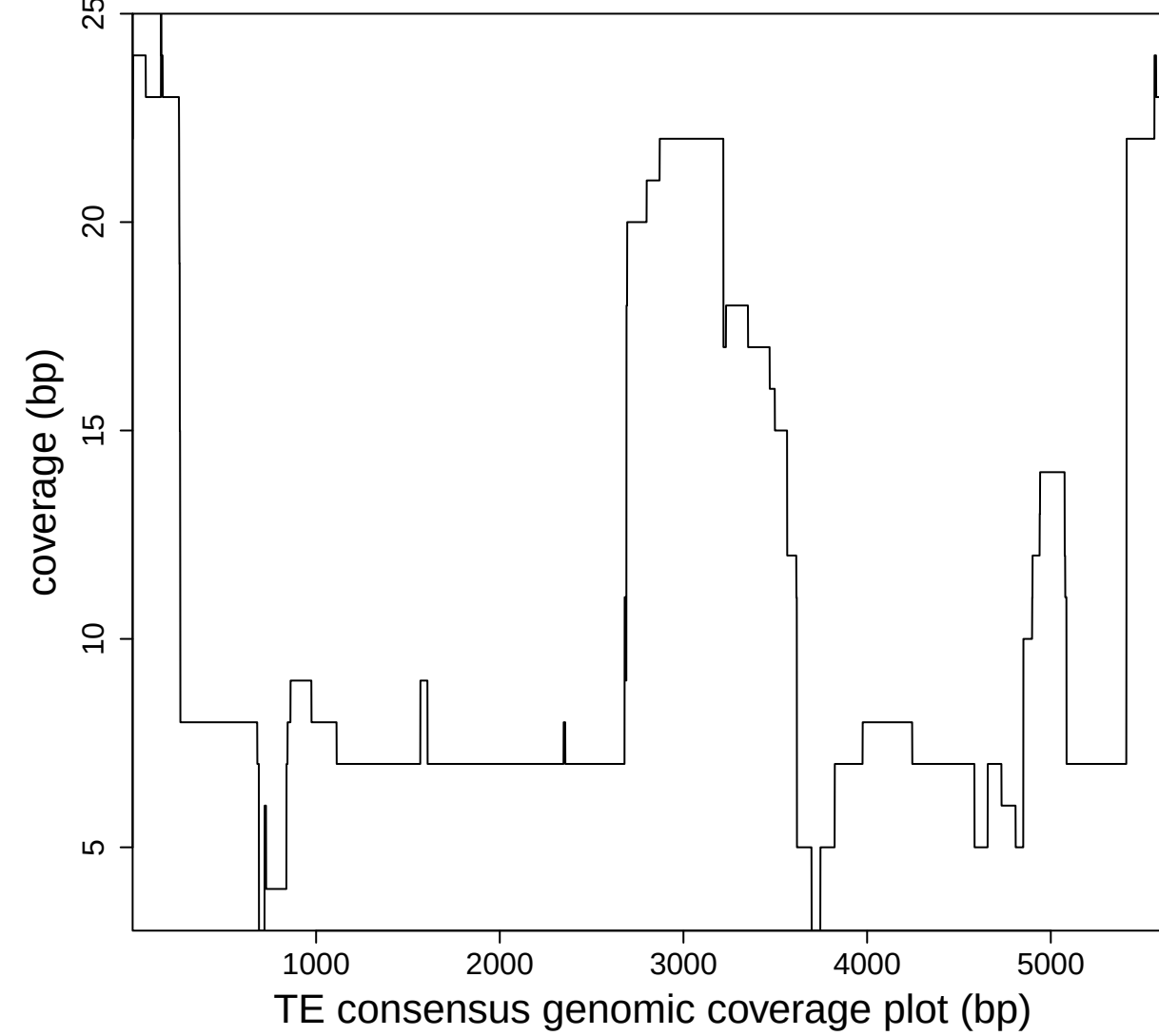
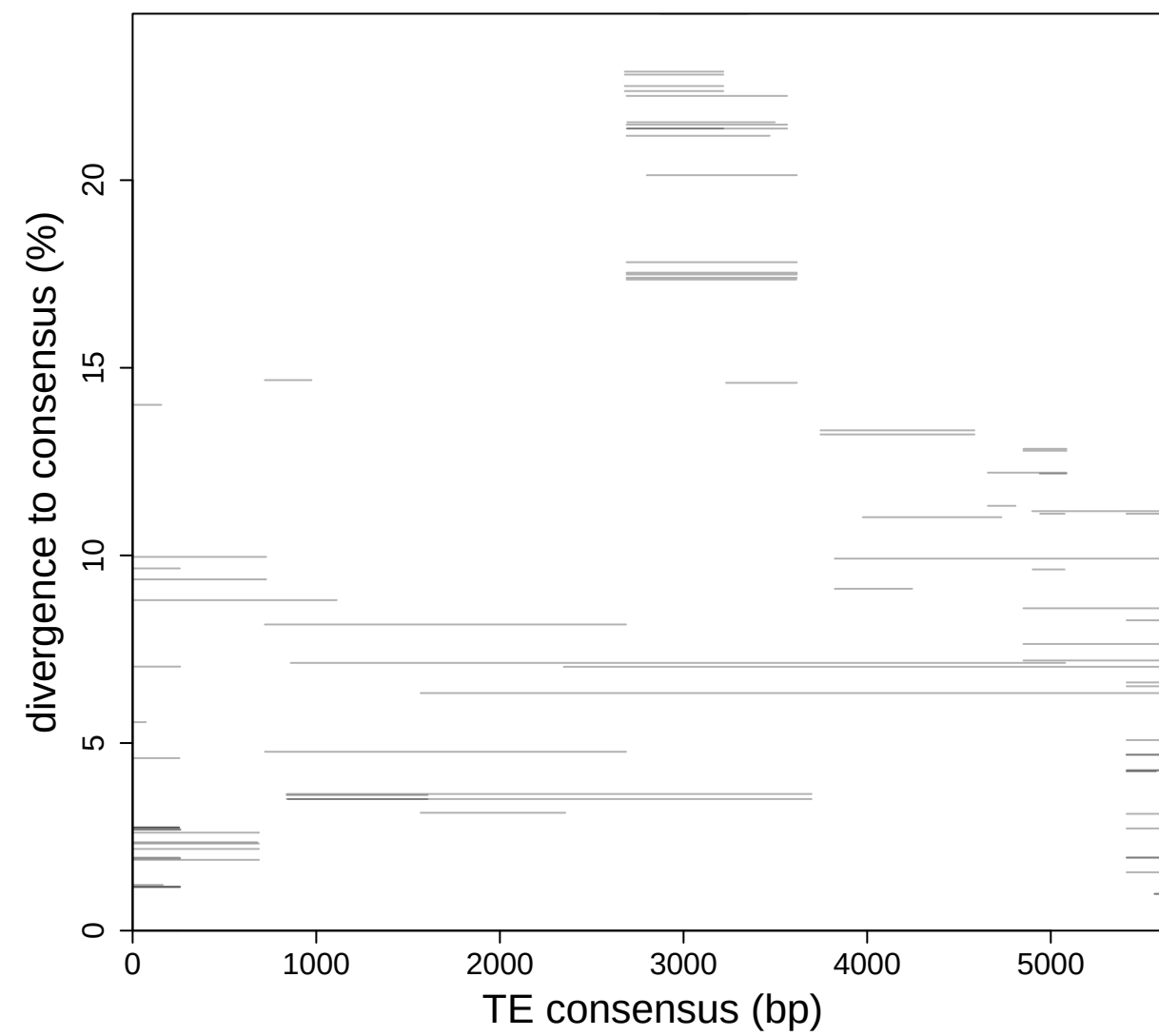
Start crop Point

End crop Point



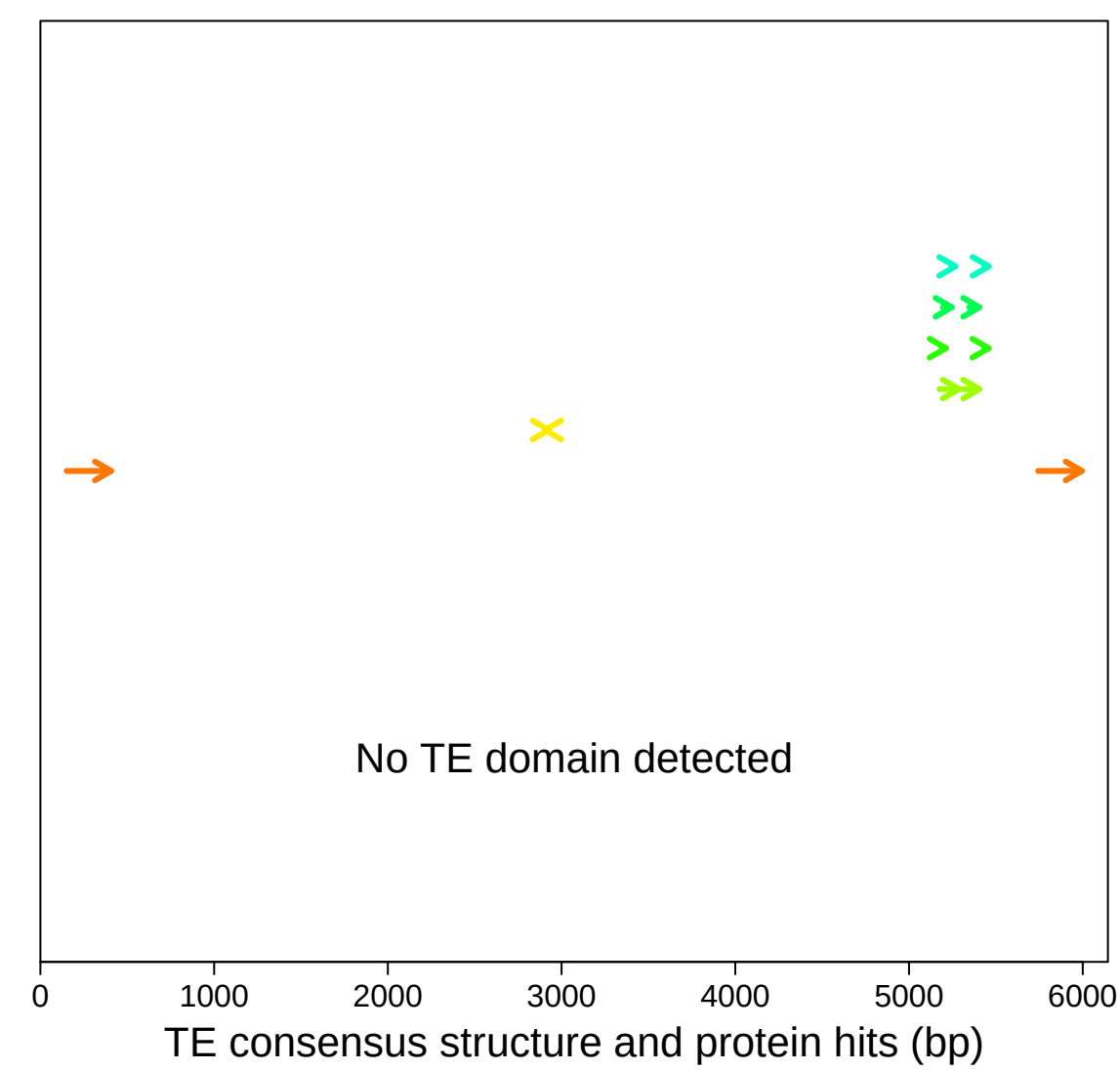
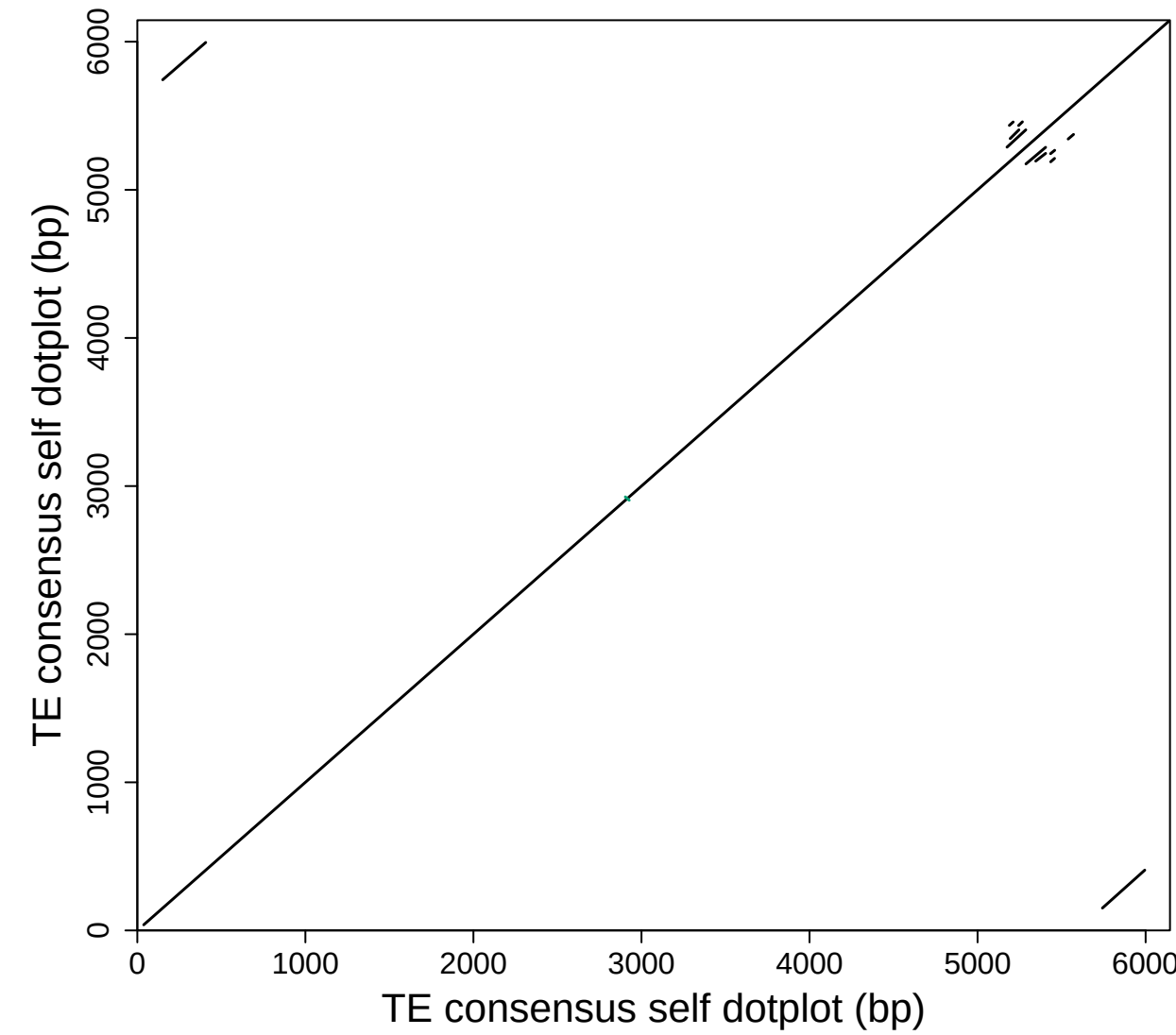
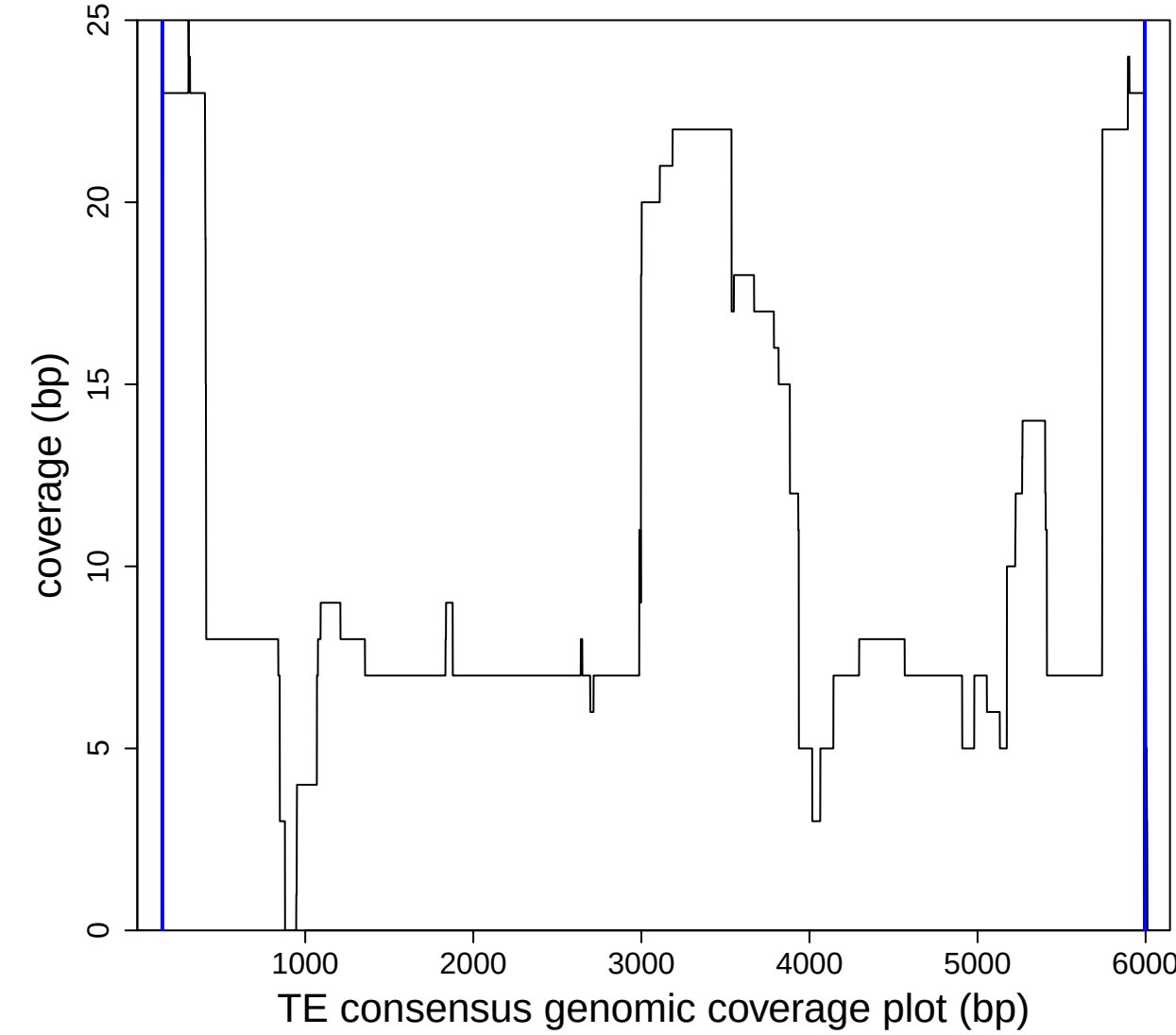
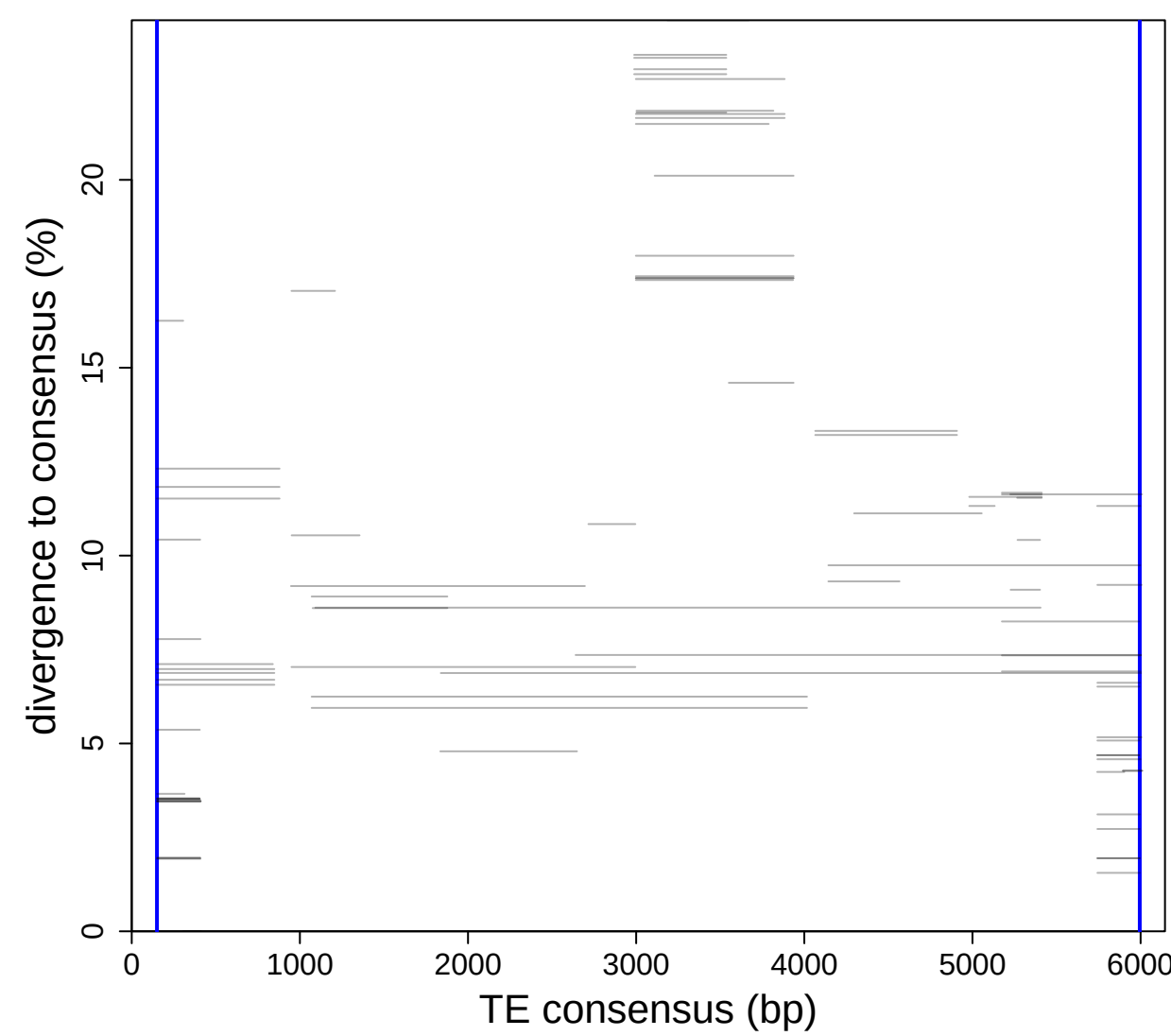
ed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 5665bp; fragments: 88; full length: 0 (>=5098.5bp)

After TEtrimmer 5665 bp

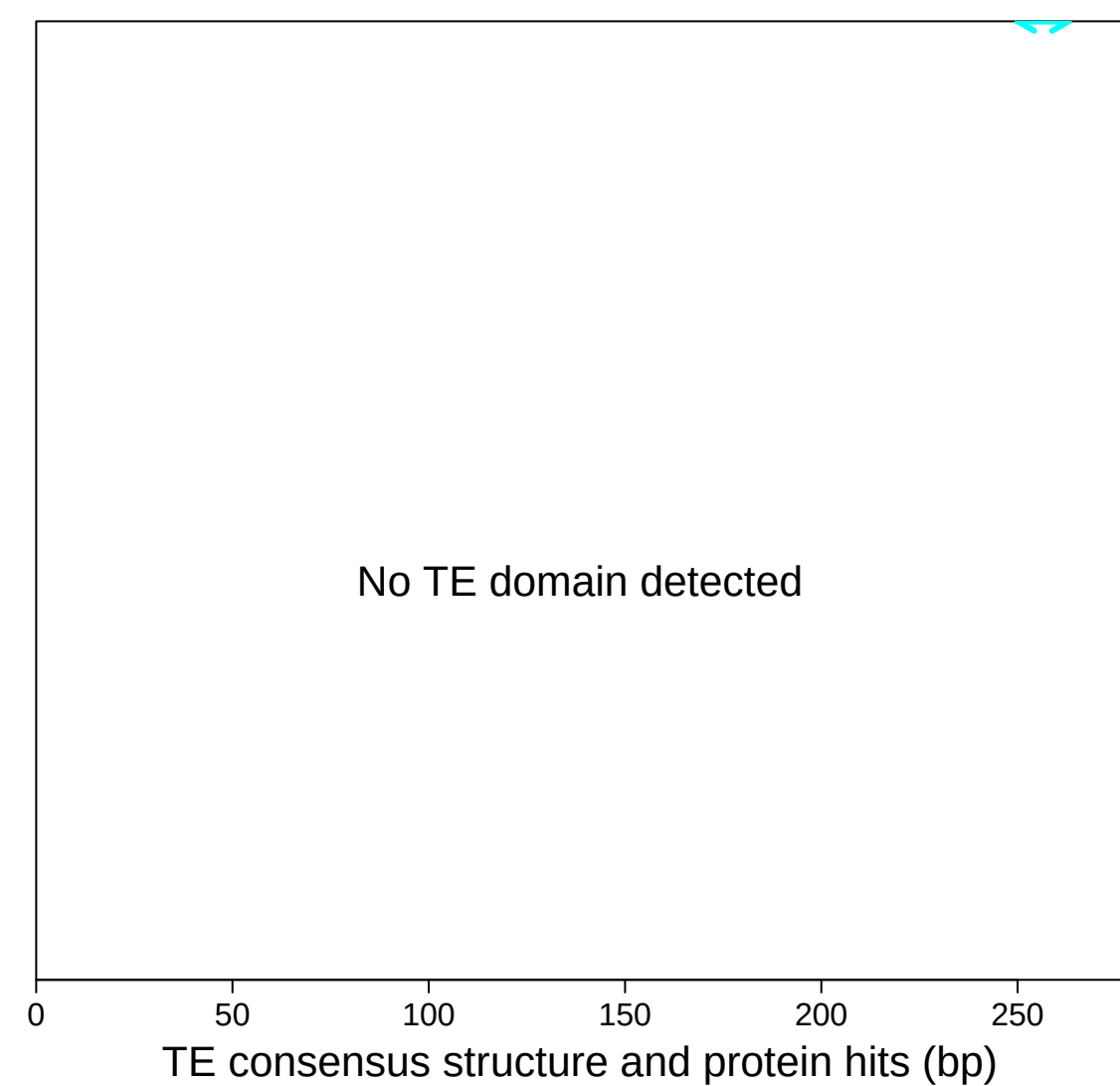
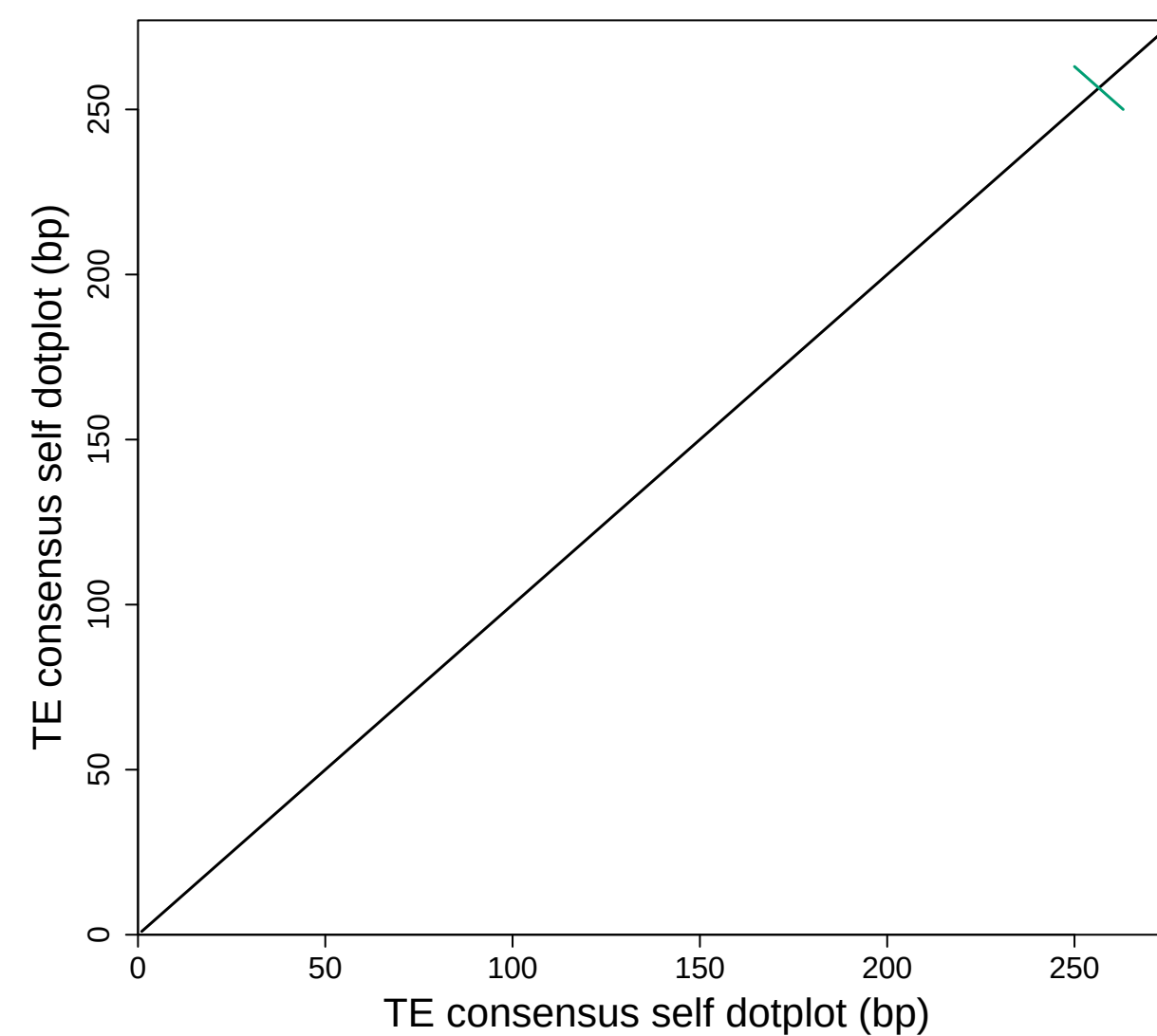
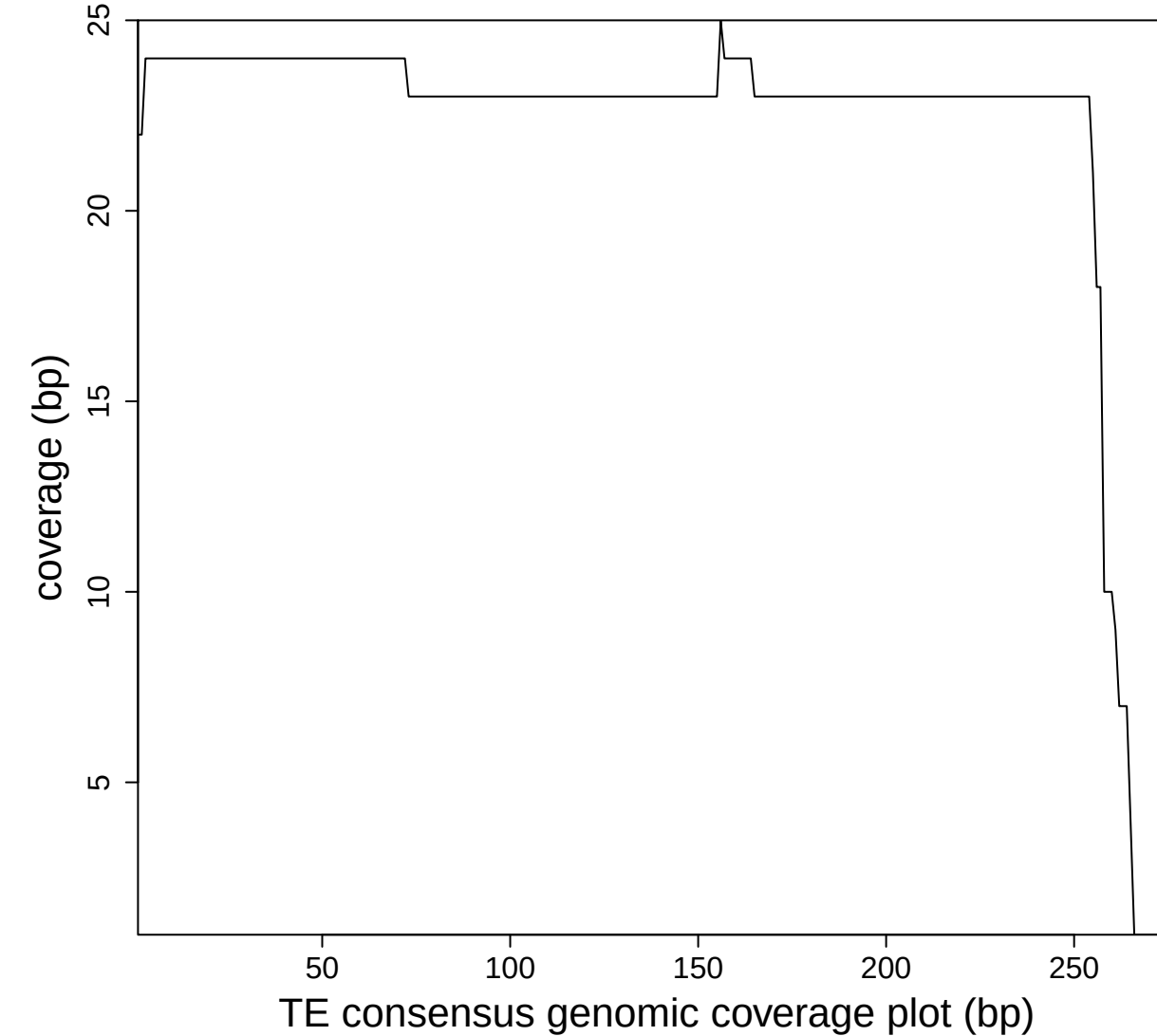
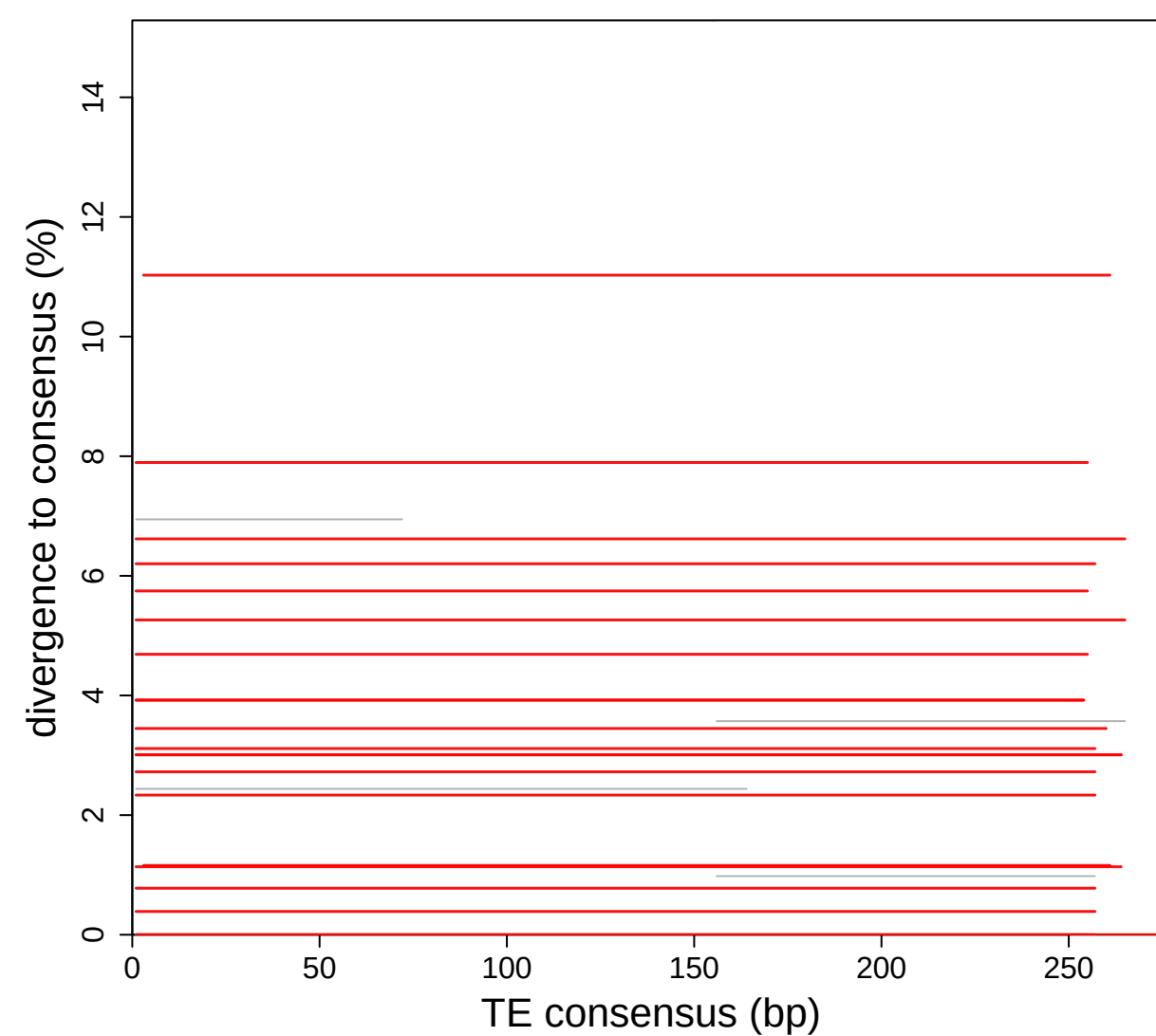


ly_283.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcIn.fa_aln.fa
size: 6145bp; fragments: 89; full length: 0 (>=5530.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

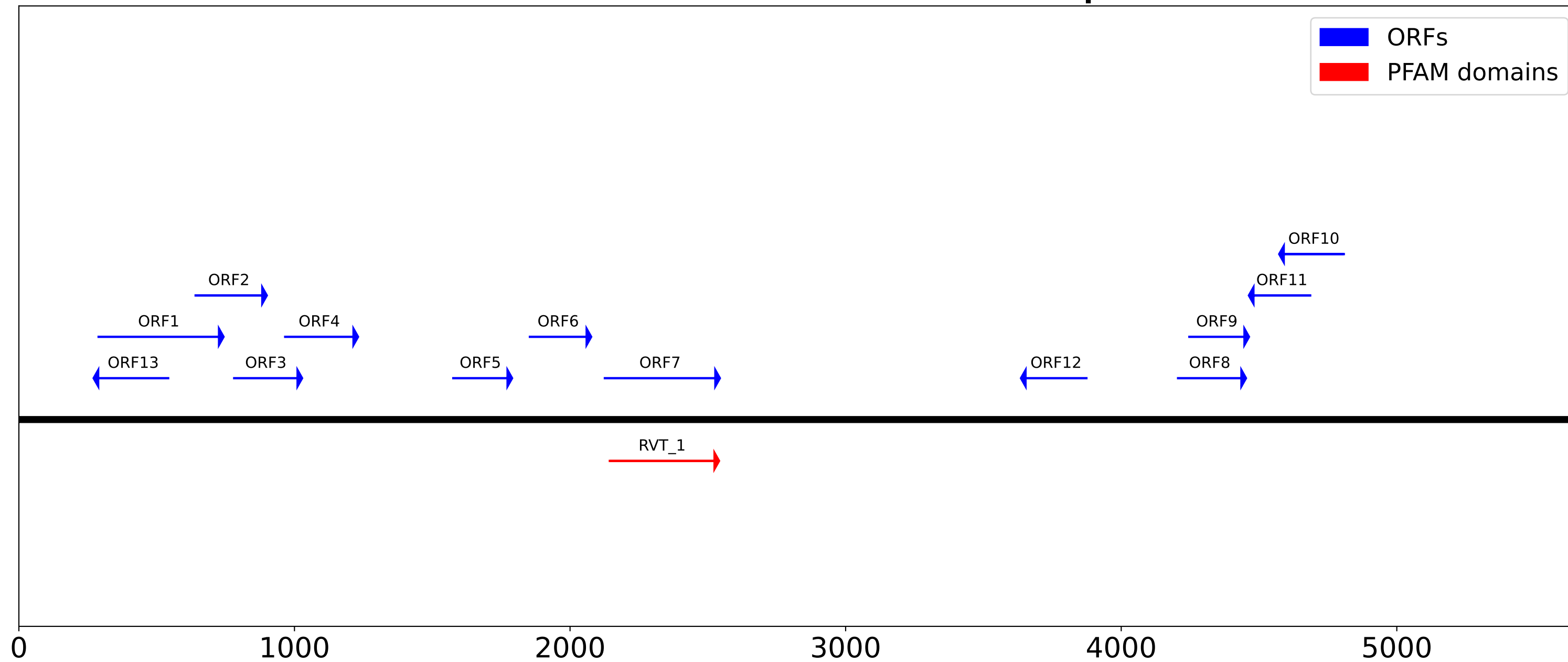


TE: ltr_1_family_283
size: 277bp; fragments: 26; full length: 21 (≥ 249.3 bp)



Before TEtrimmer 277 bp

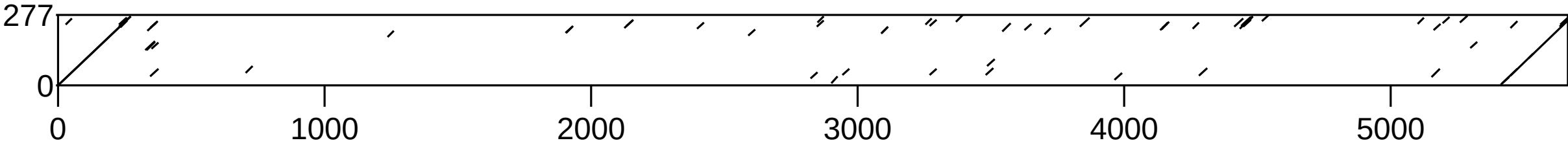
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)