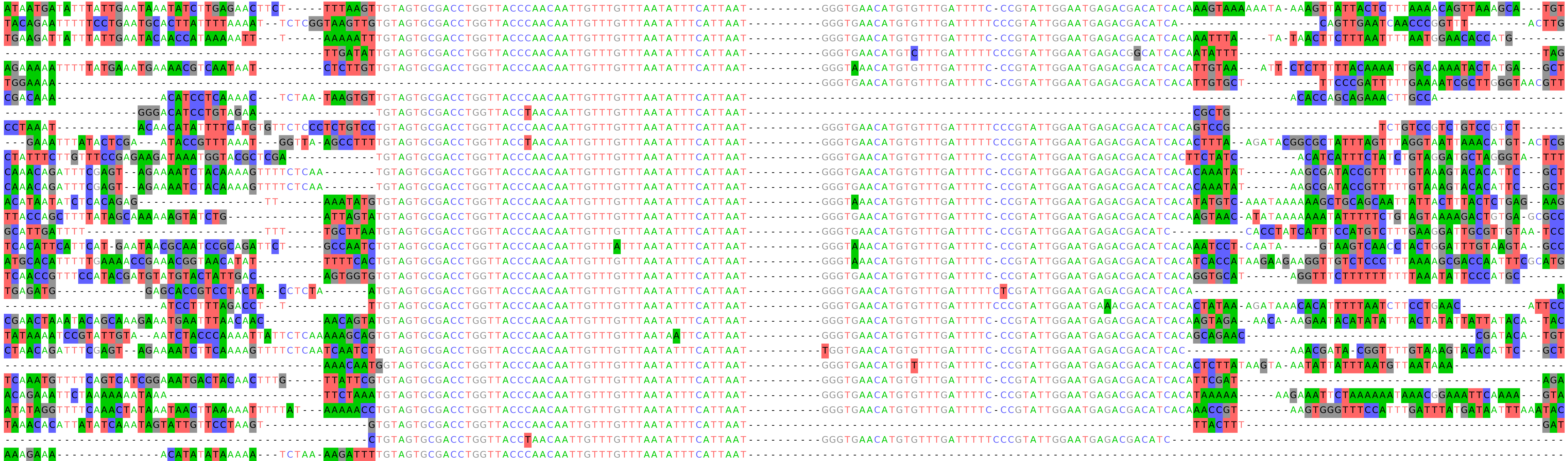


MSA length = 6544

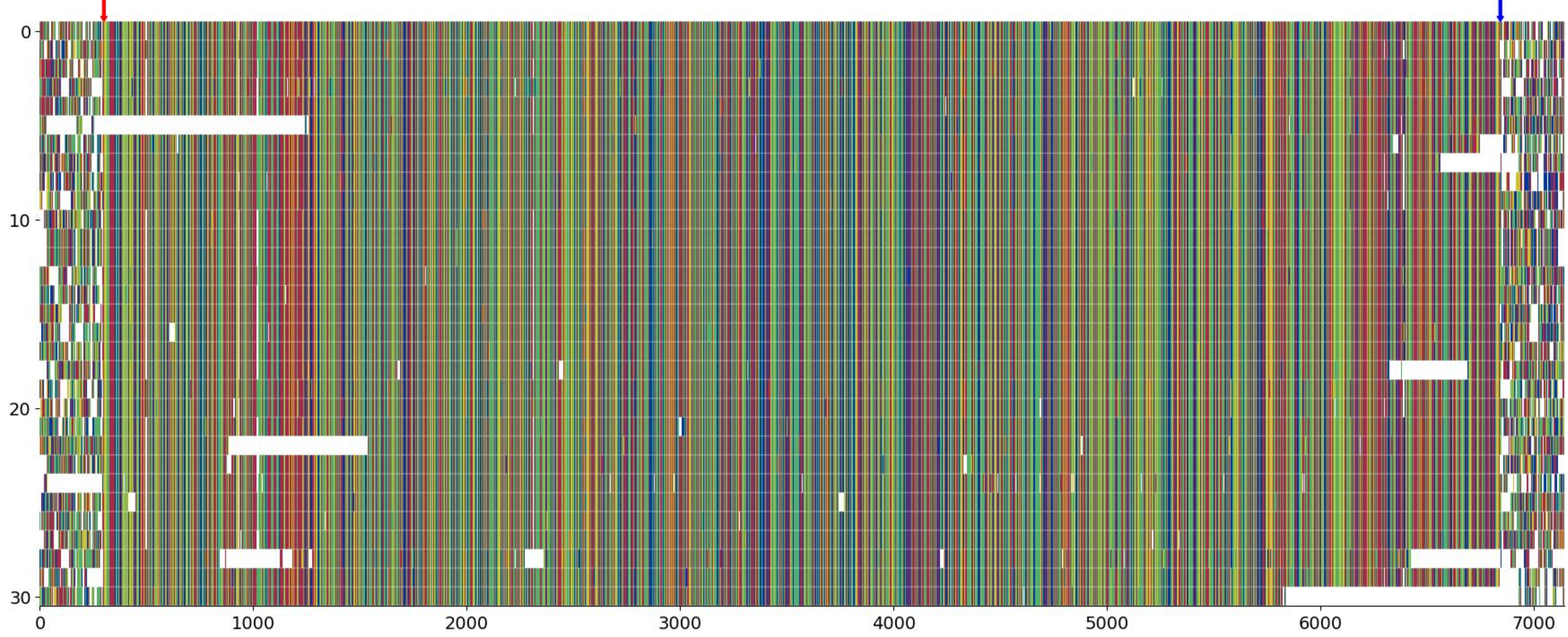
Start crop Point

End crop Point



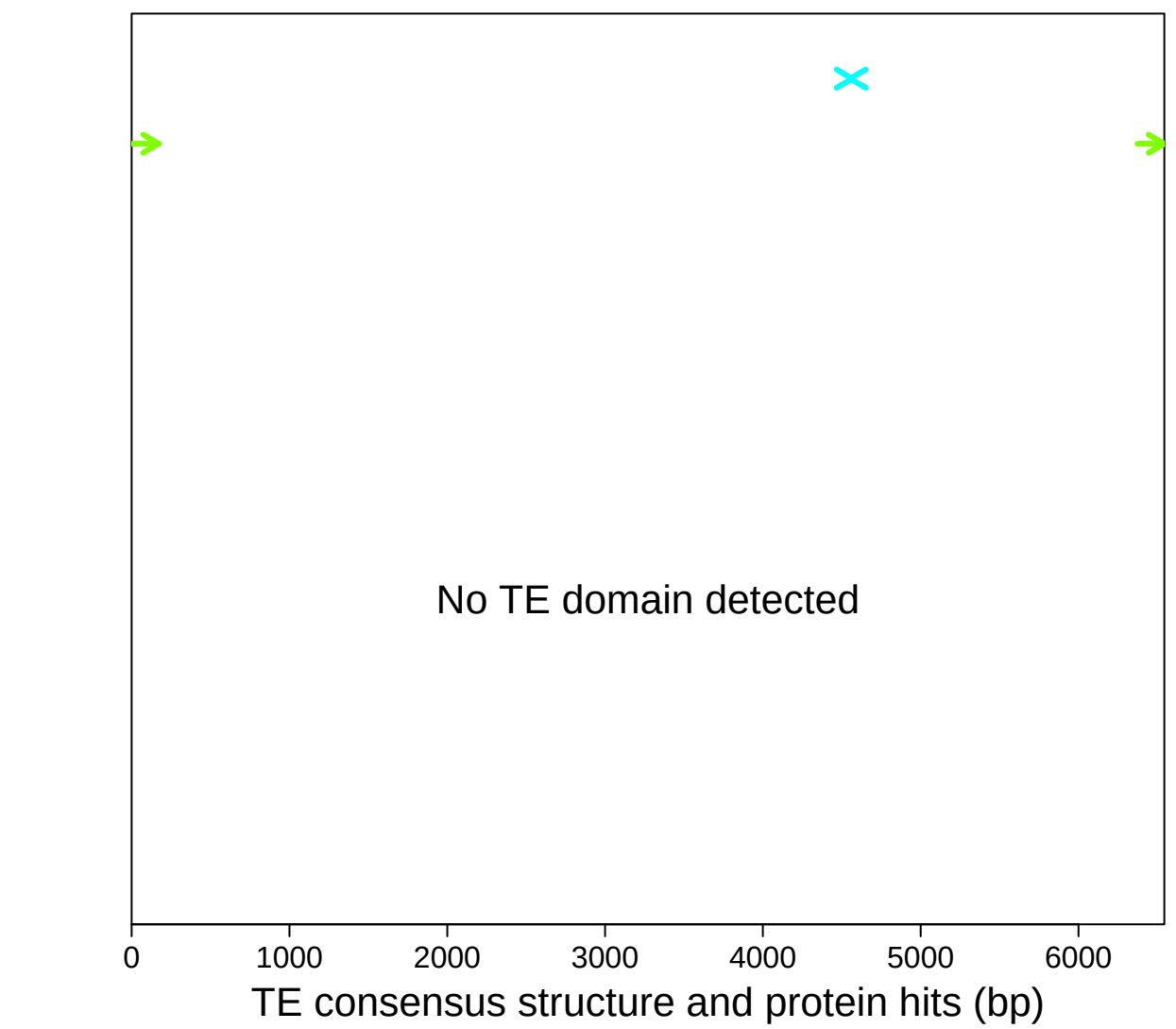
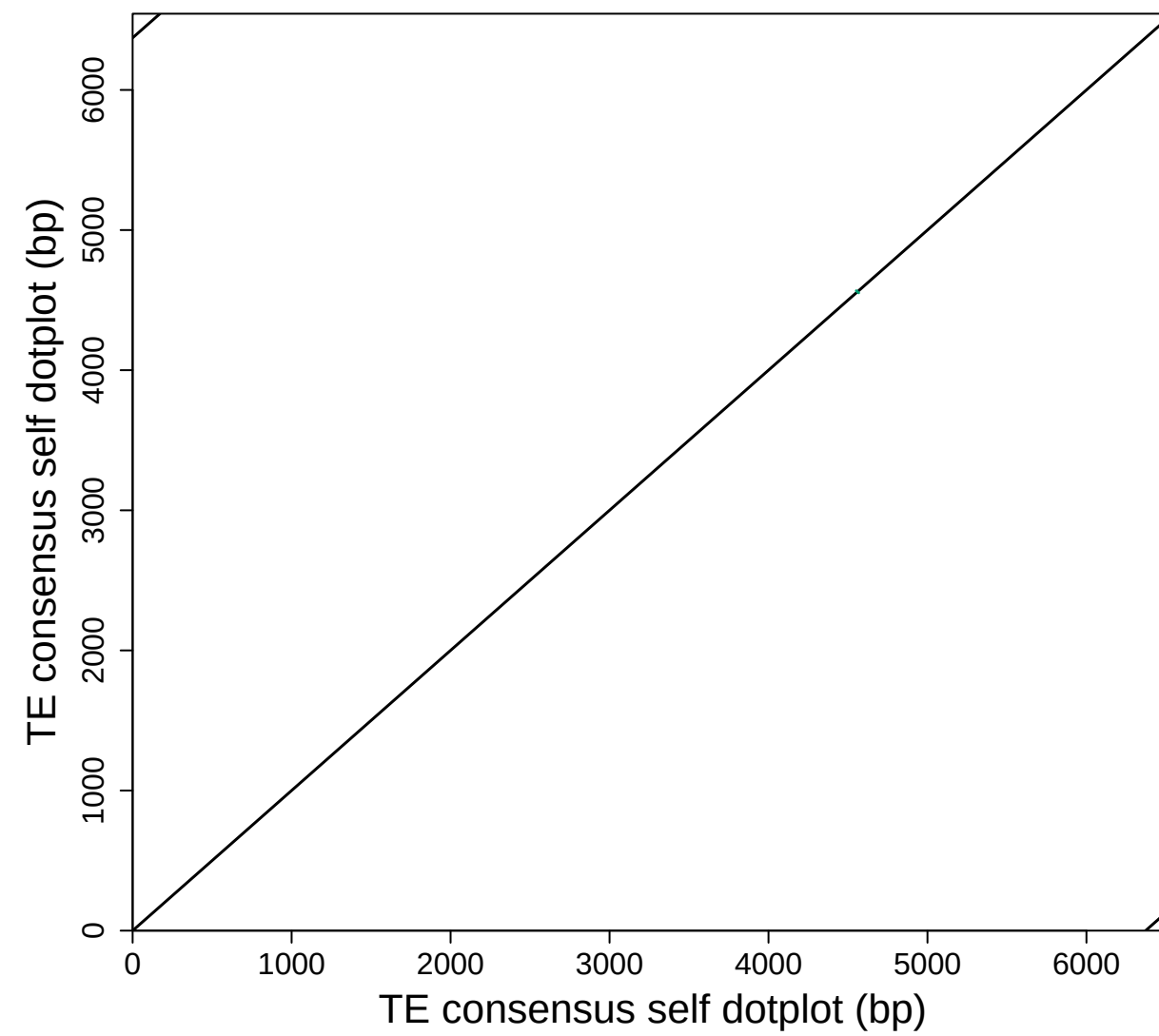
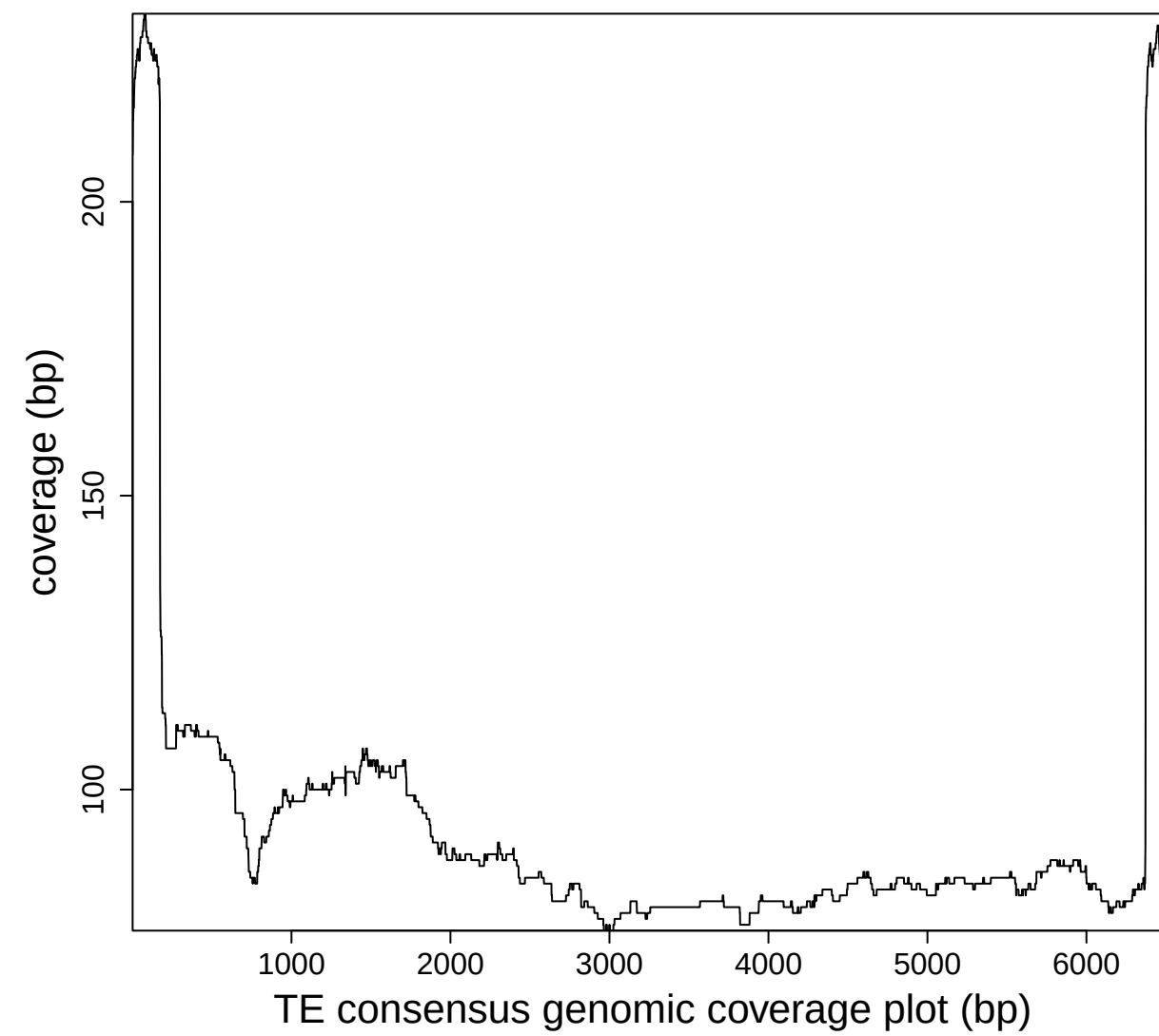
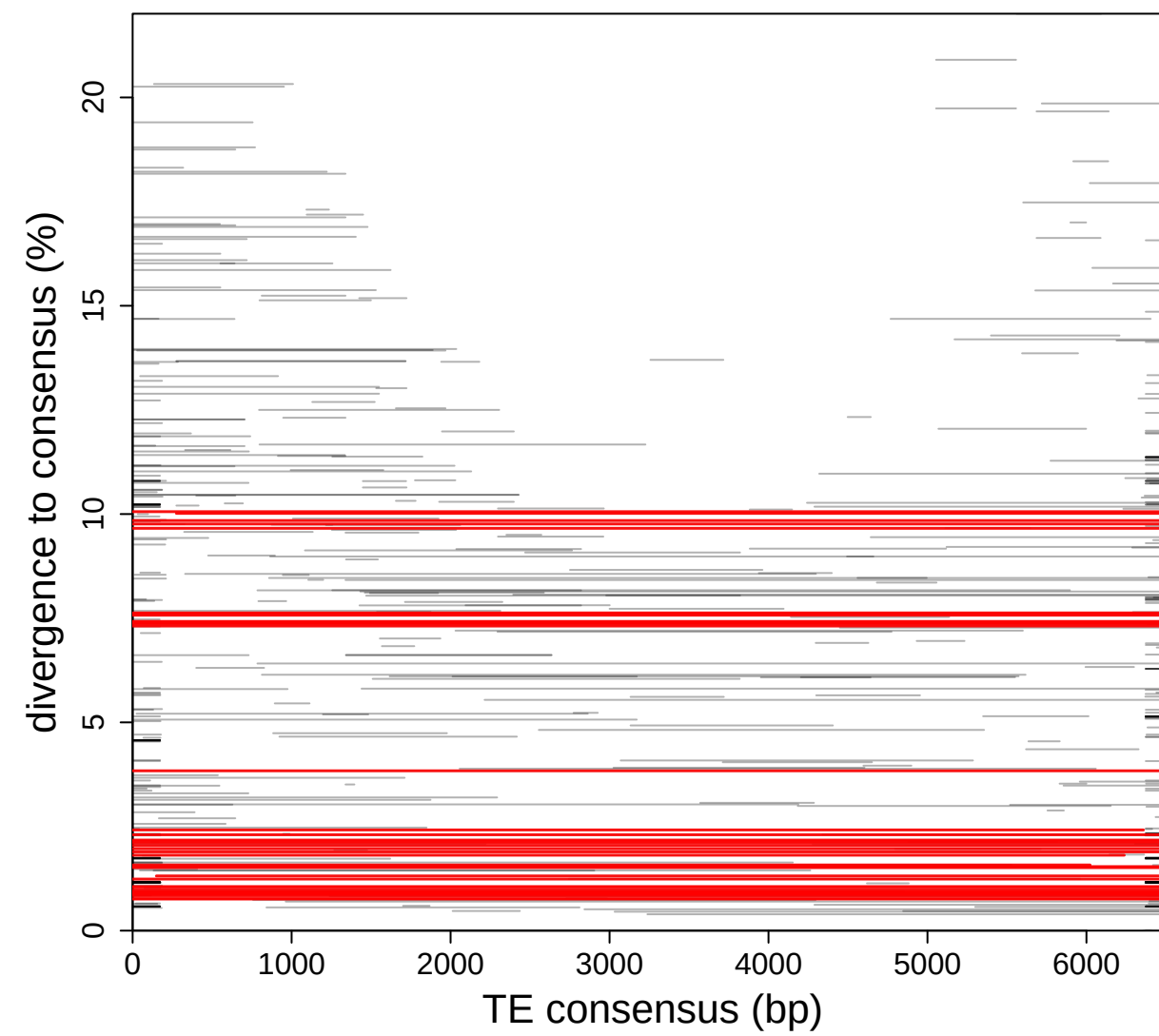
Start crop Point

End crop Point



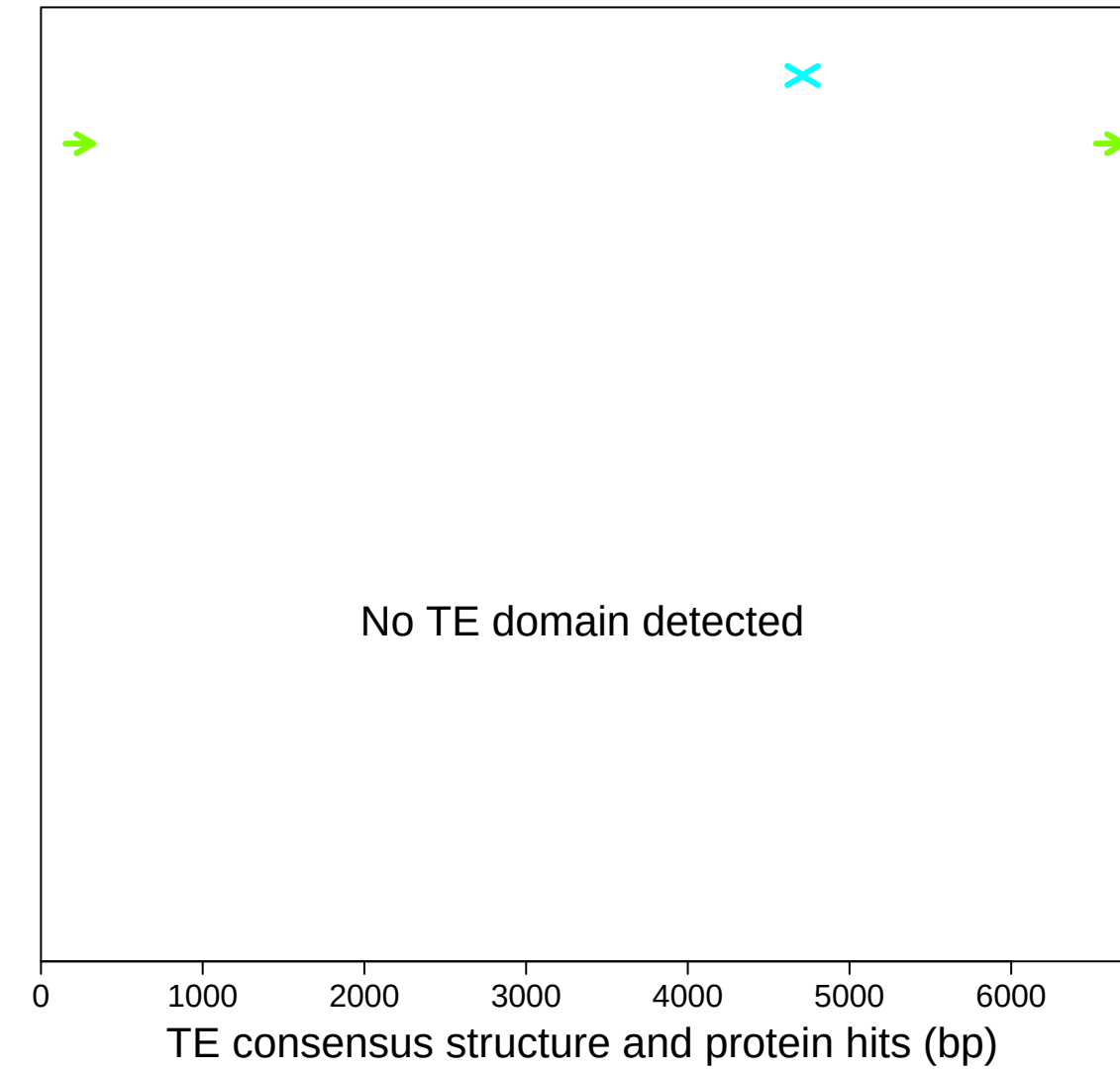
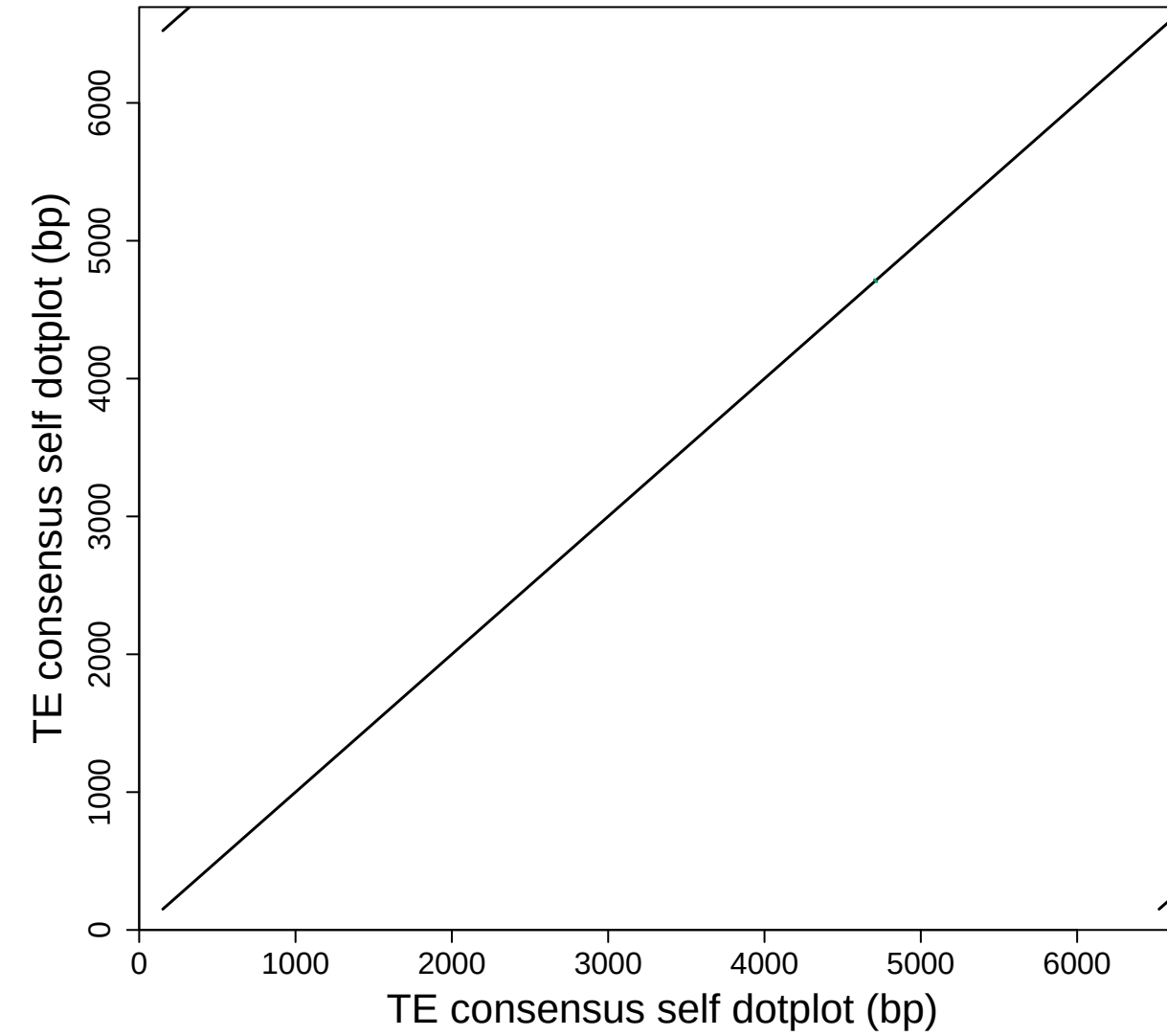
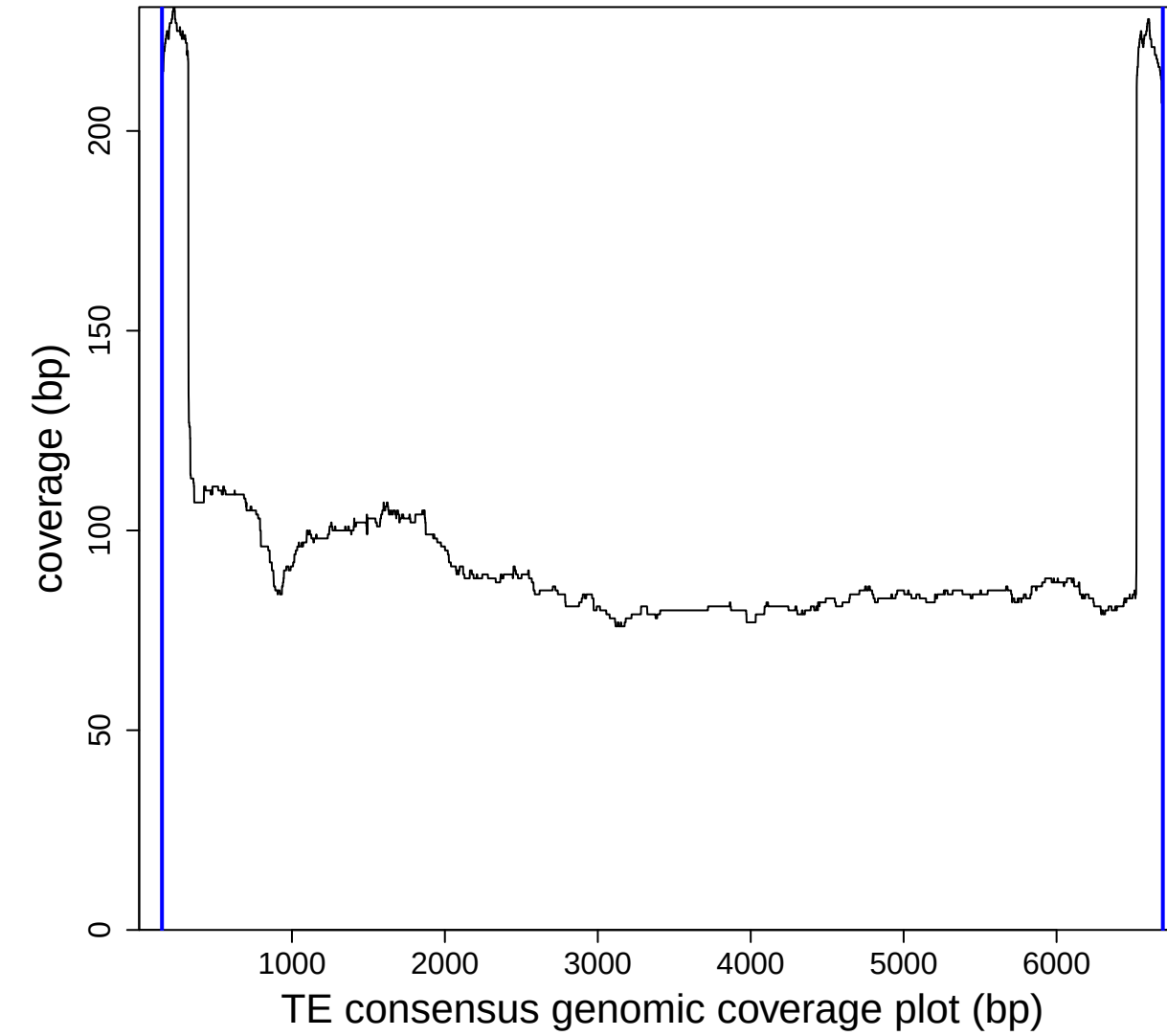
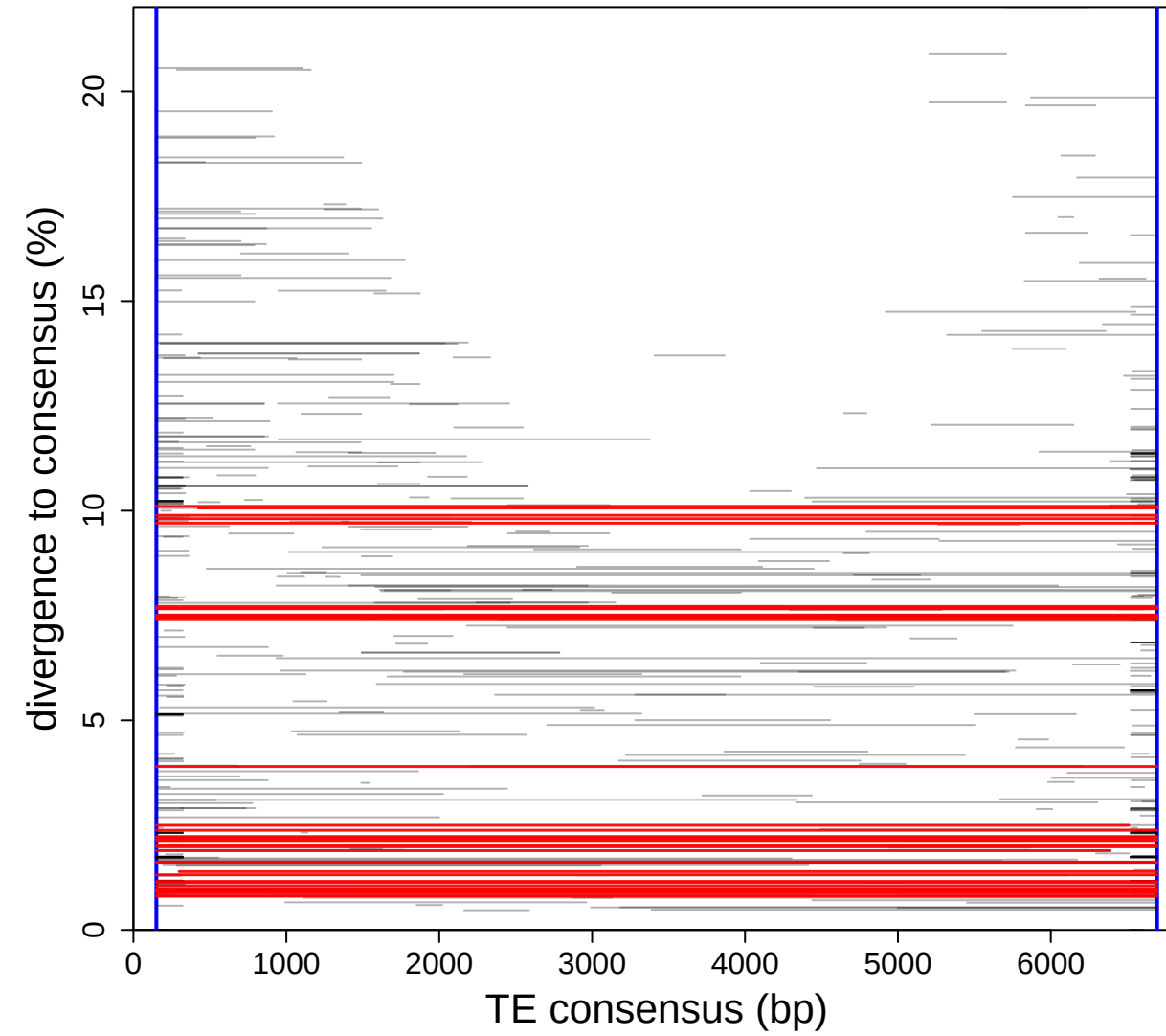
ed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 6544bp; fragments: 569; full length: 39 (≥ 5889.6 bp)

After TETrimmer 6544 bp

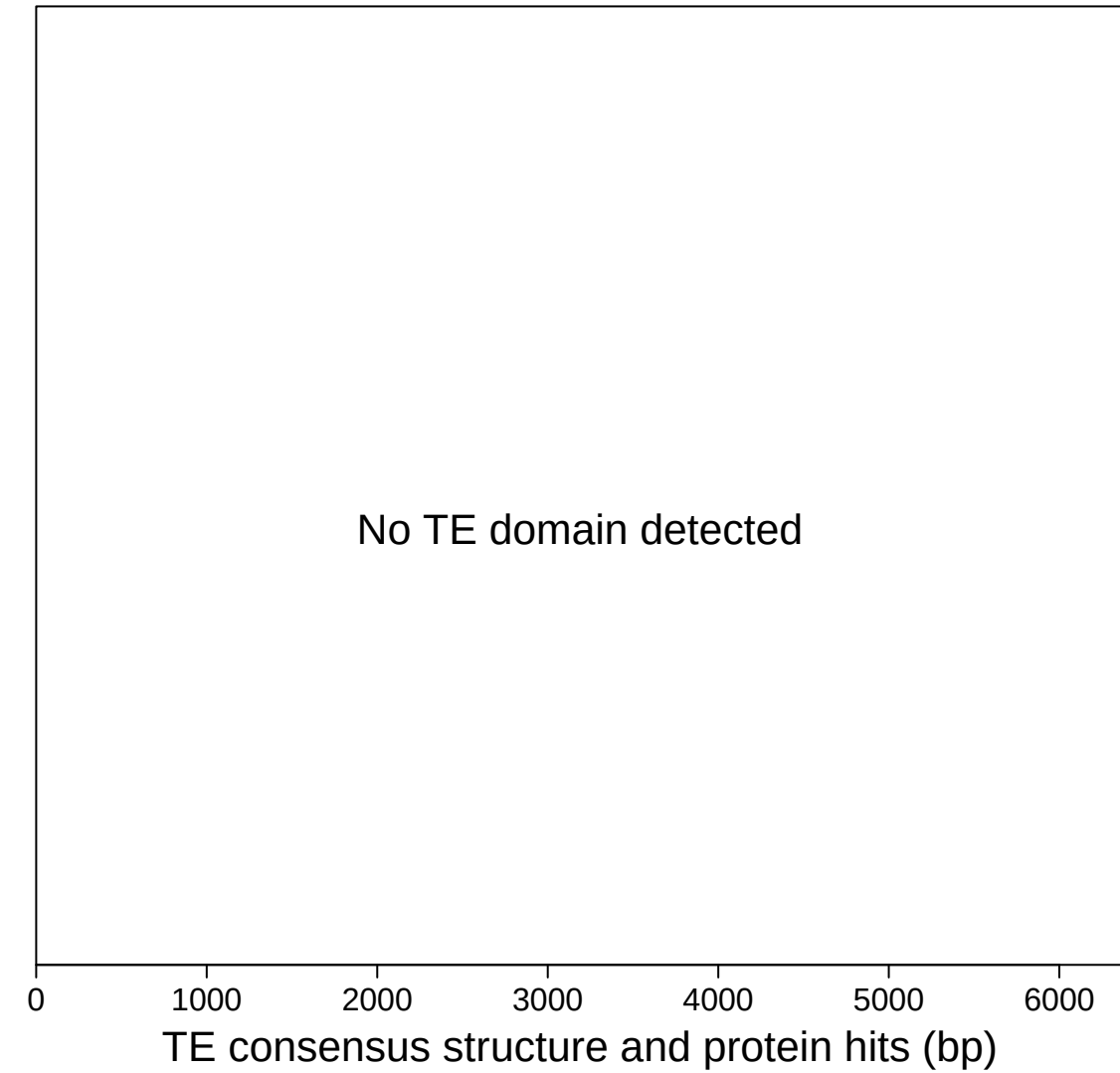
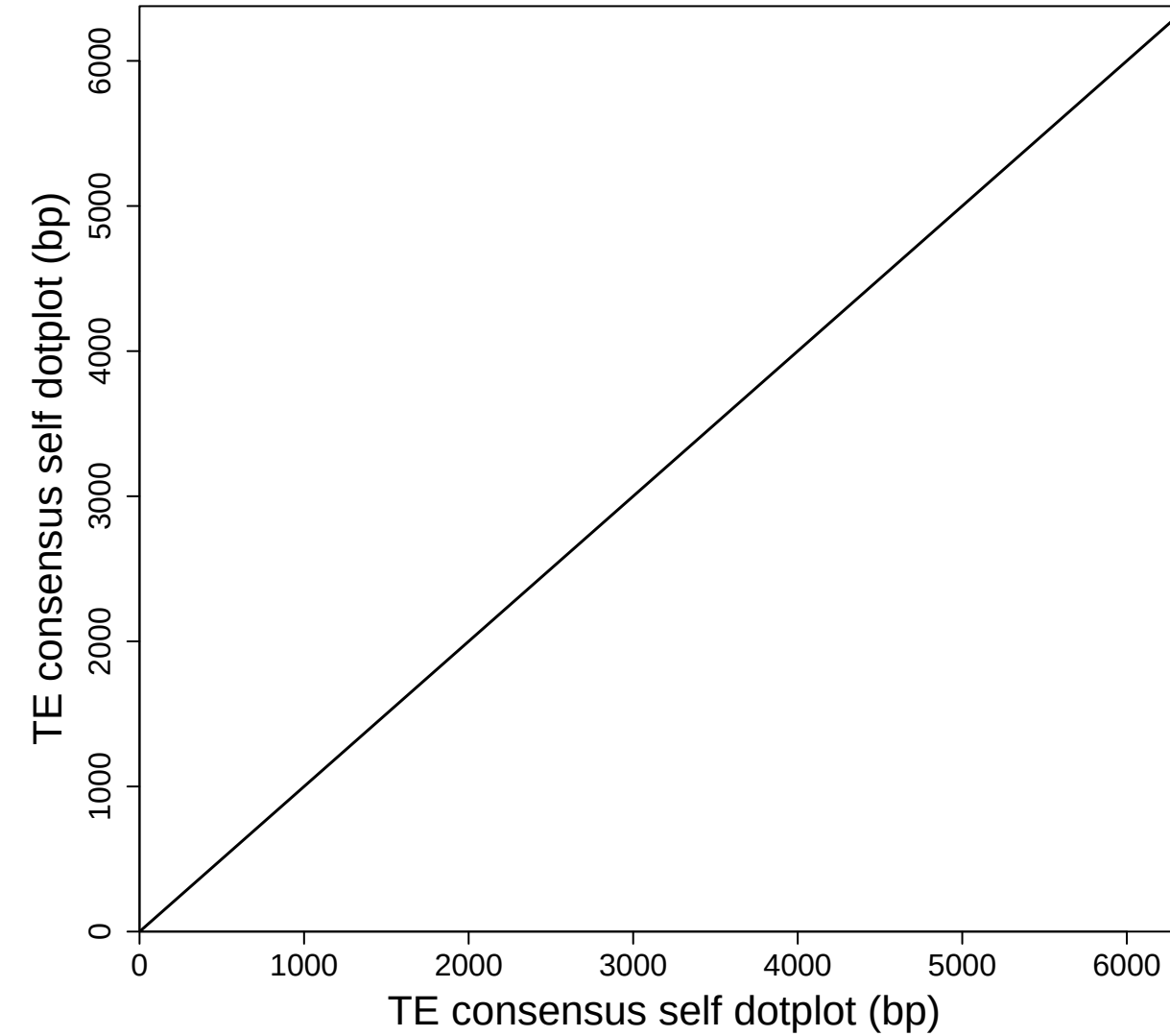
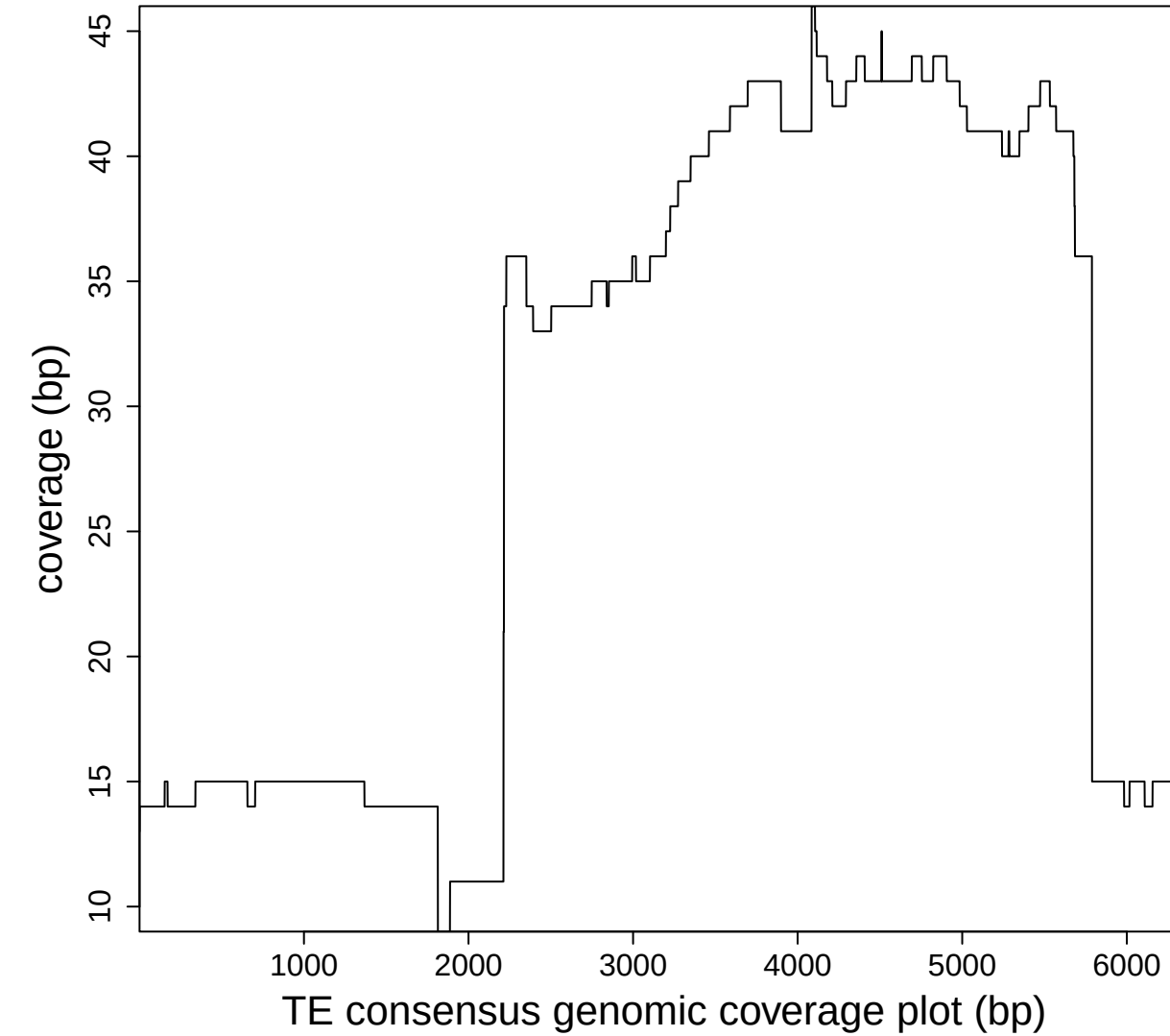
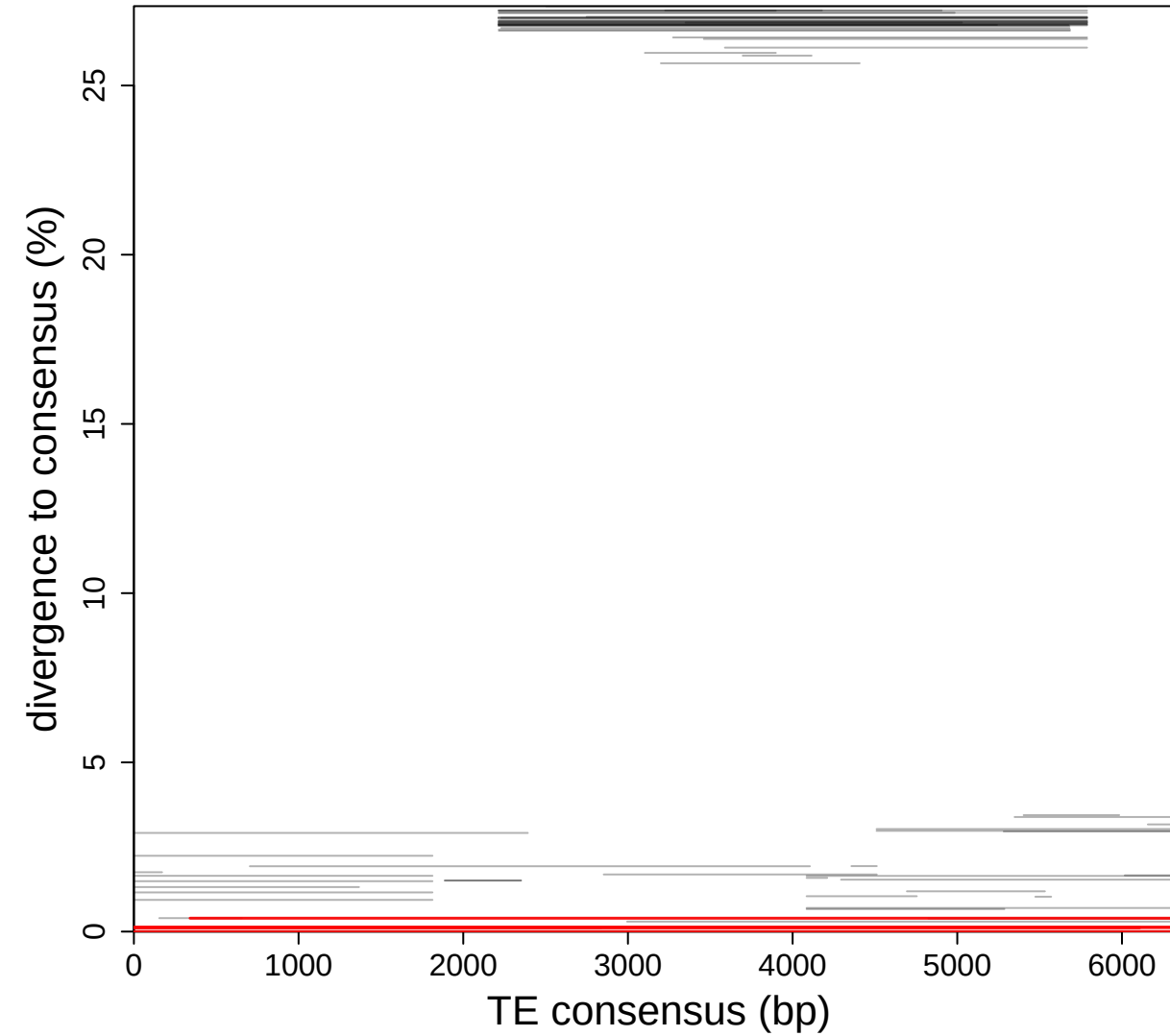


ly_772.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcIn.fa_aln.fa
size: 6845bp; fragments: 566; full length: 38 (≥ 6160.5 bp)

After TEtrimmer Extended plot Blue lines are boundaries

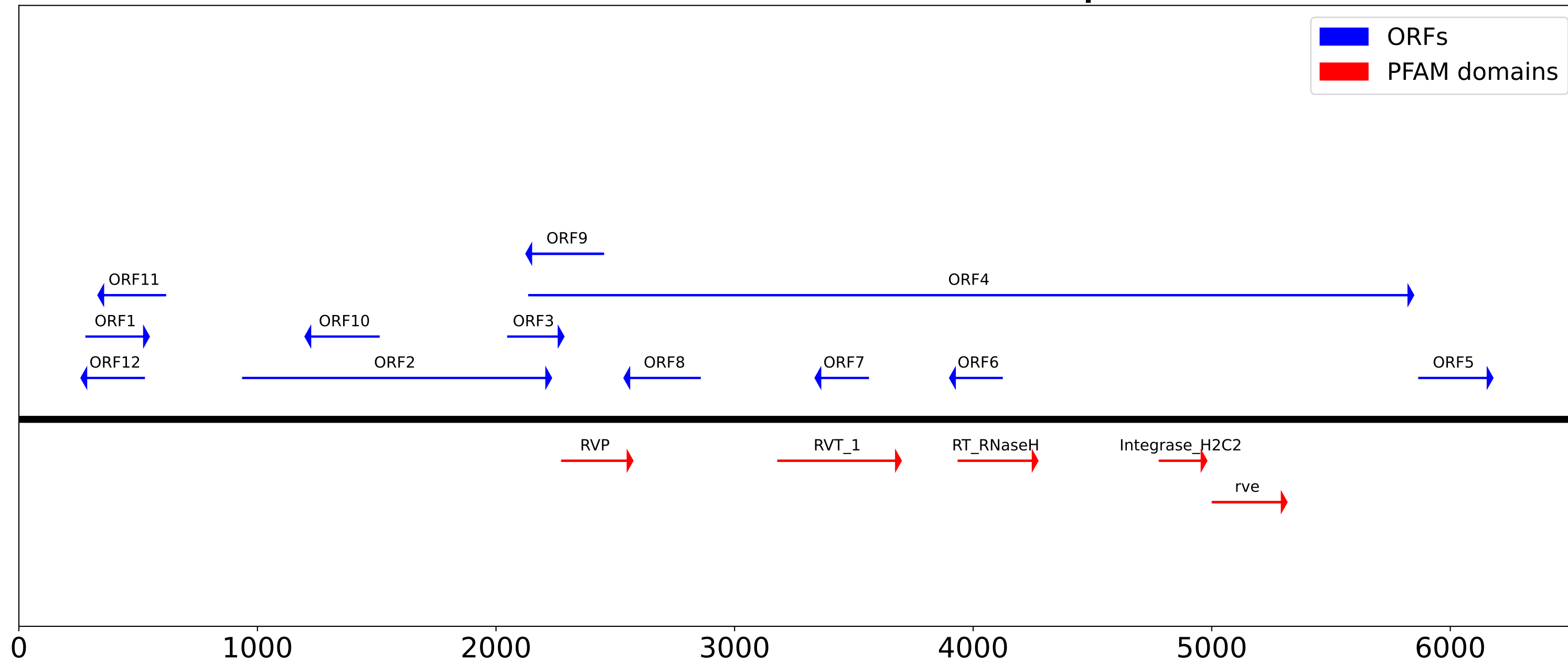


TE: ltr_1_family_772
size: 6377bp; fragments: 73; full length: 5 (≥ 5739.3 bp)

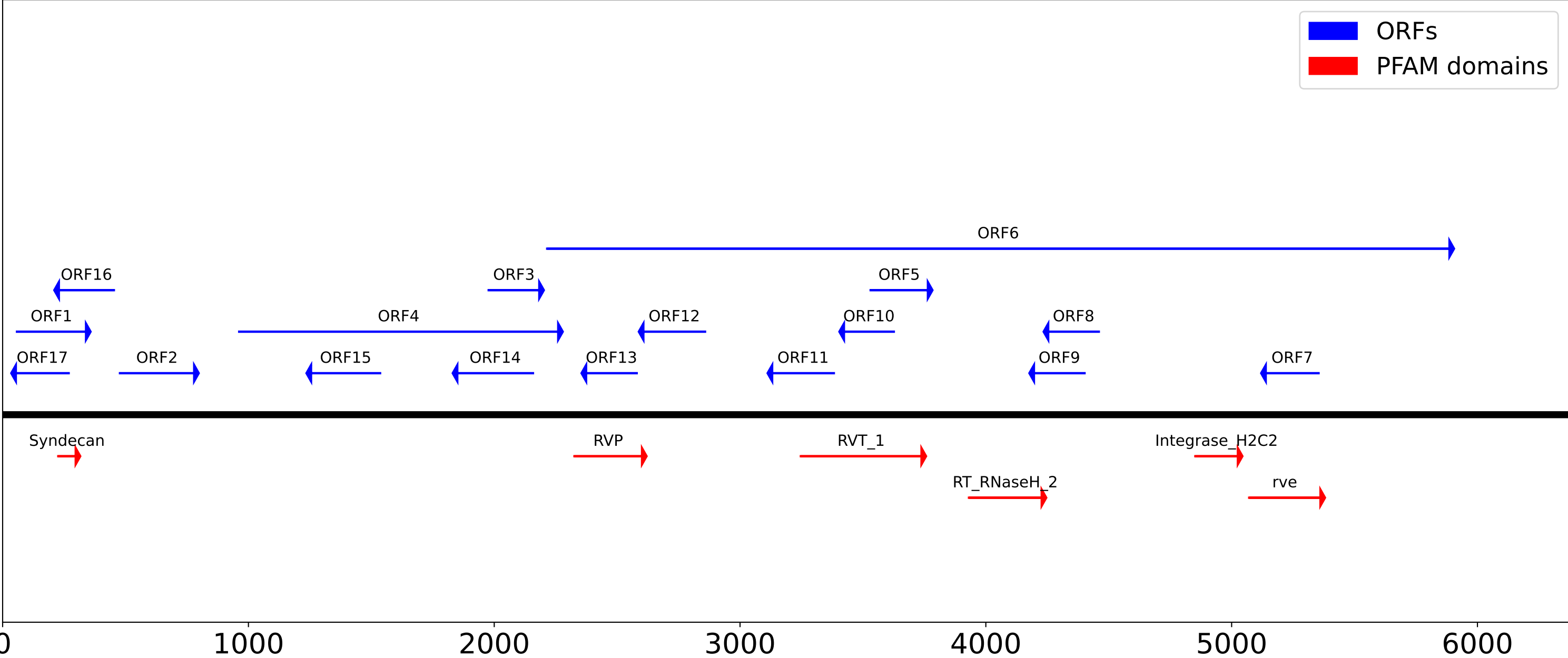


Before TEtrimmer 6377 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

