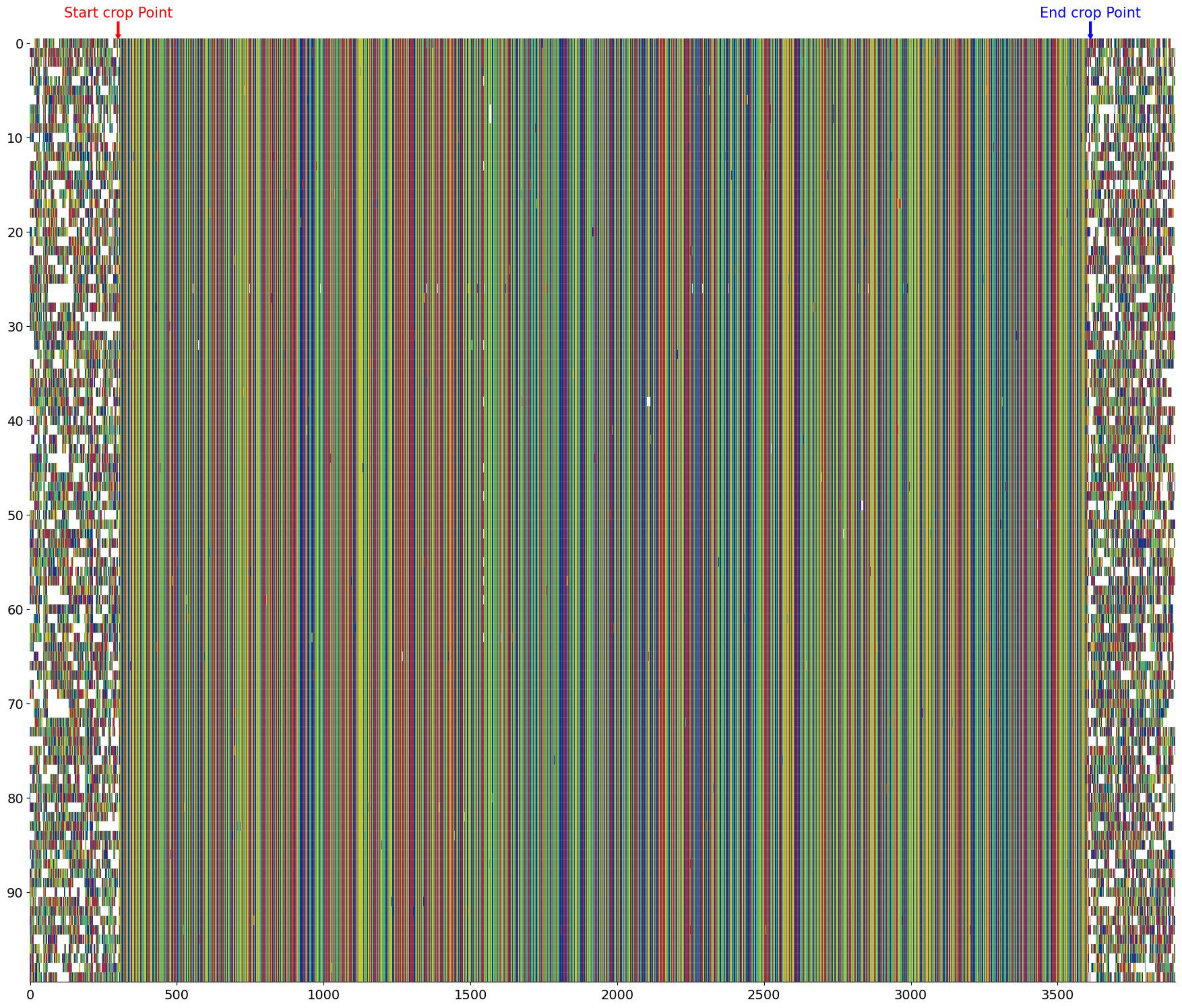
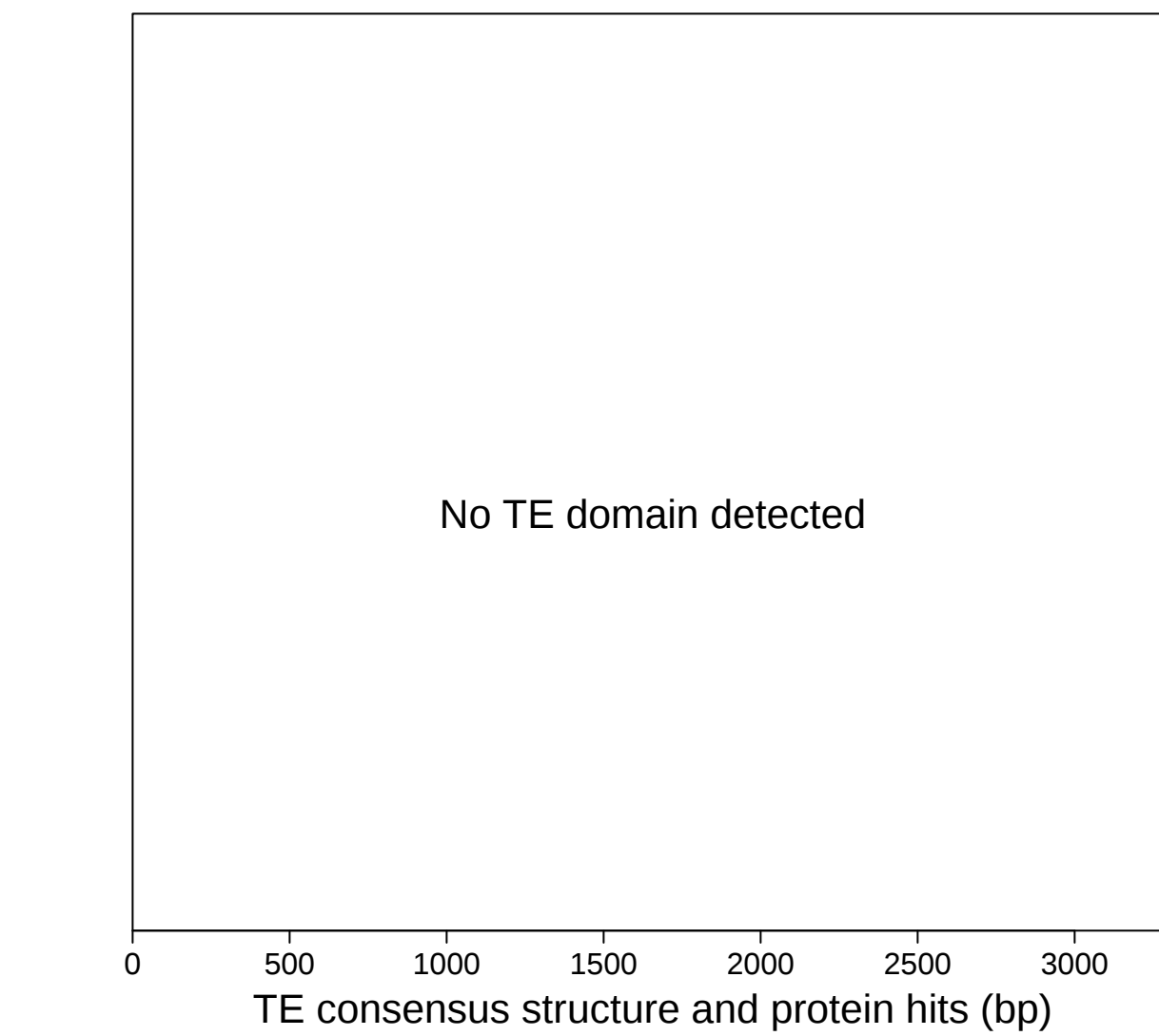
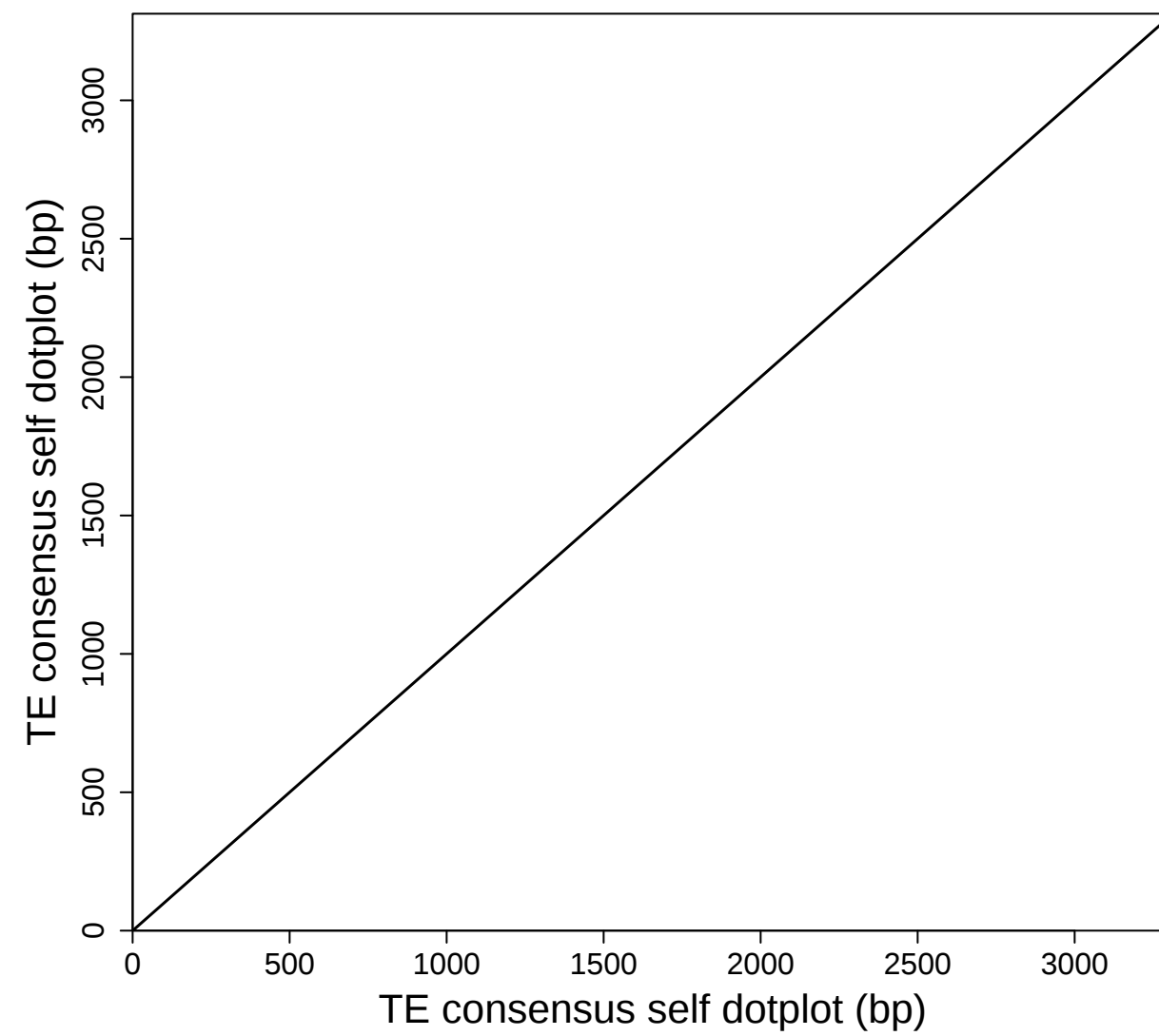
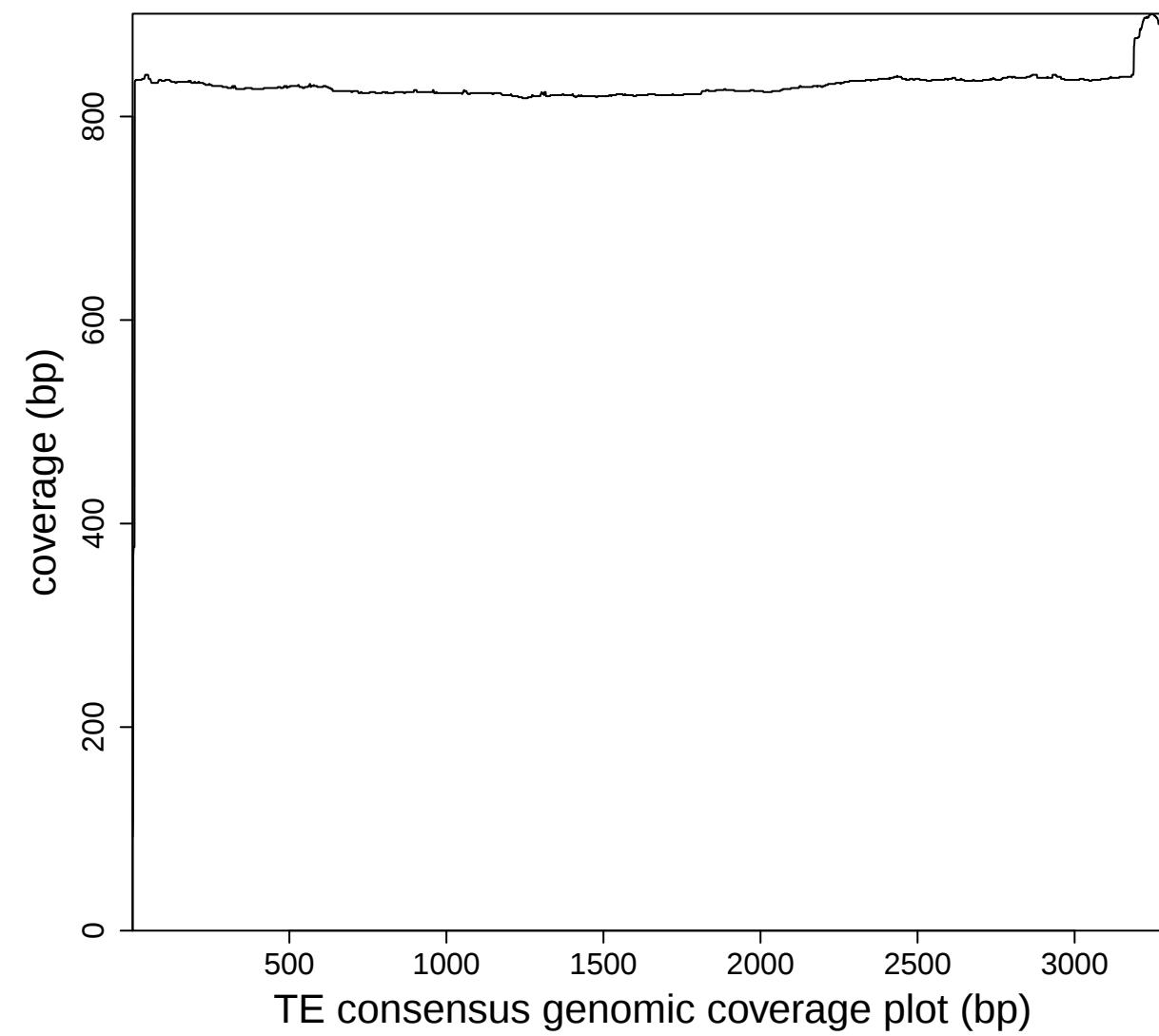
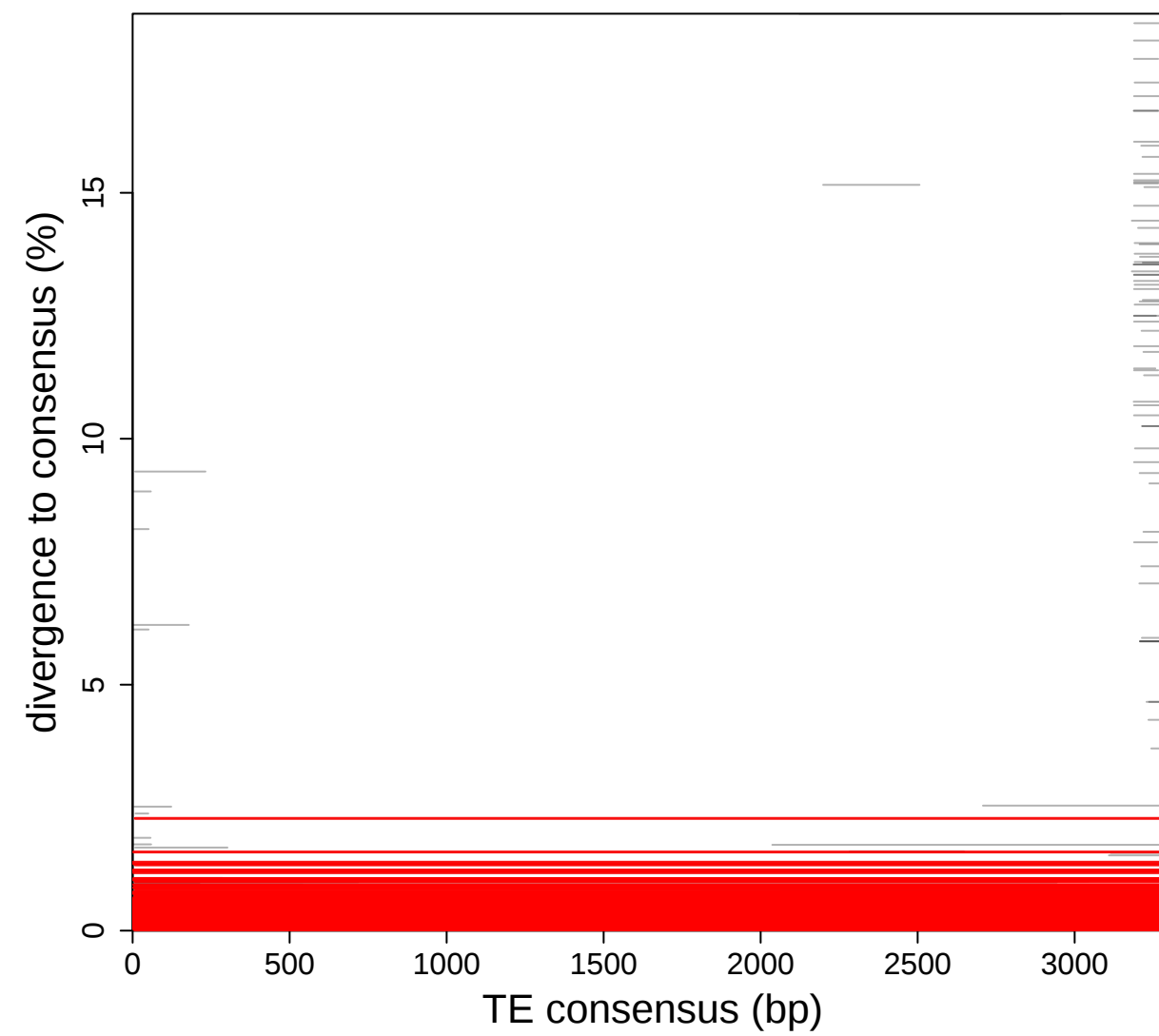






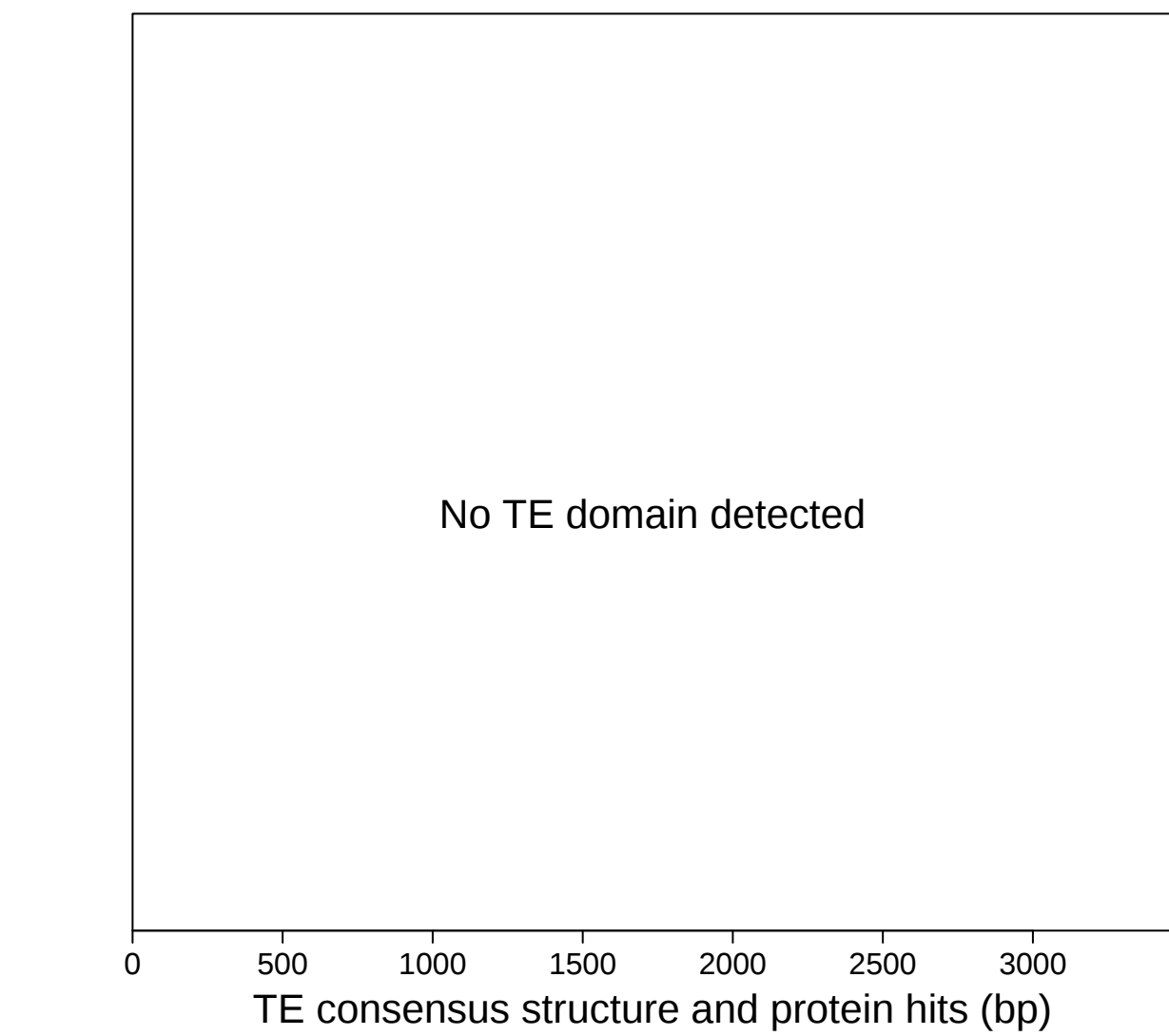
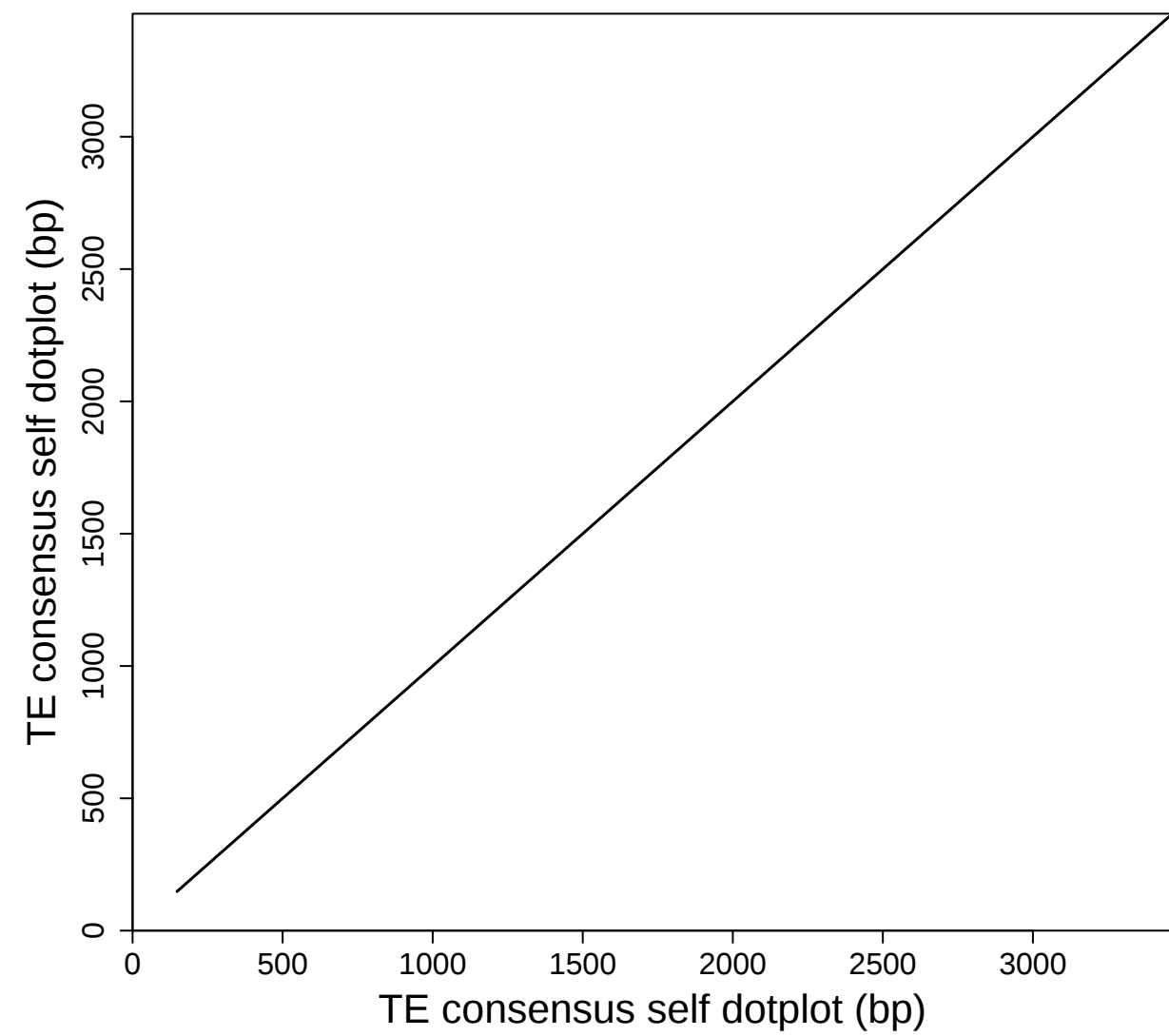
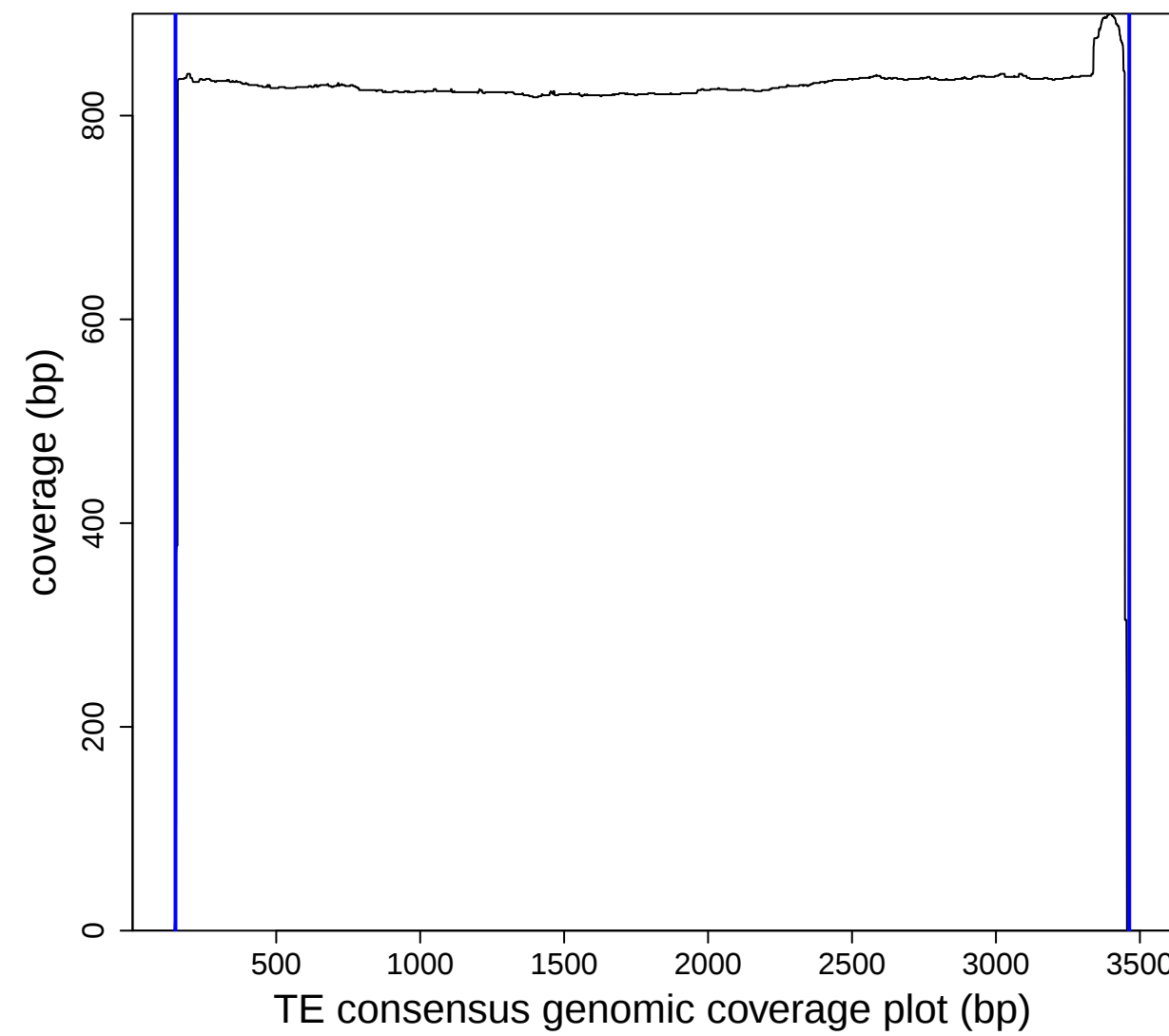
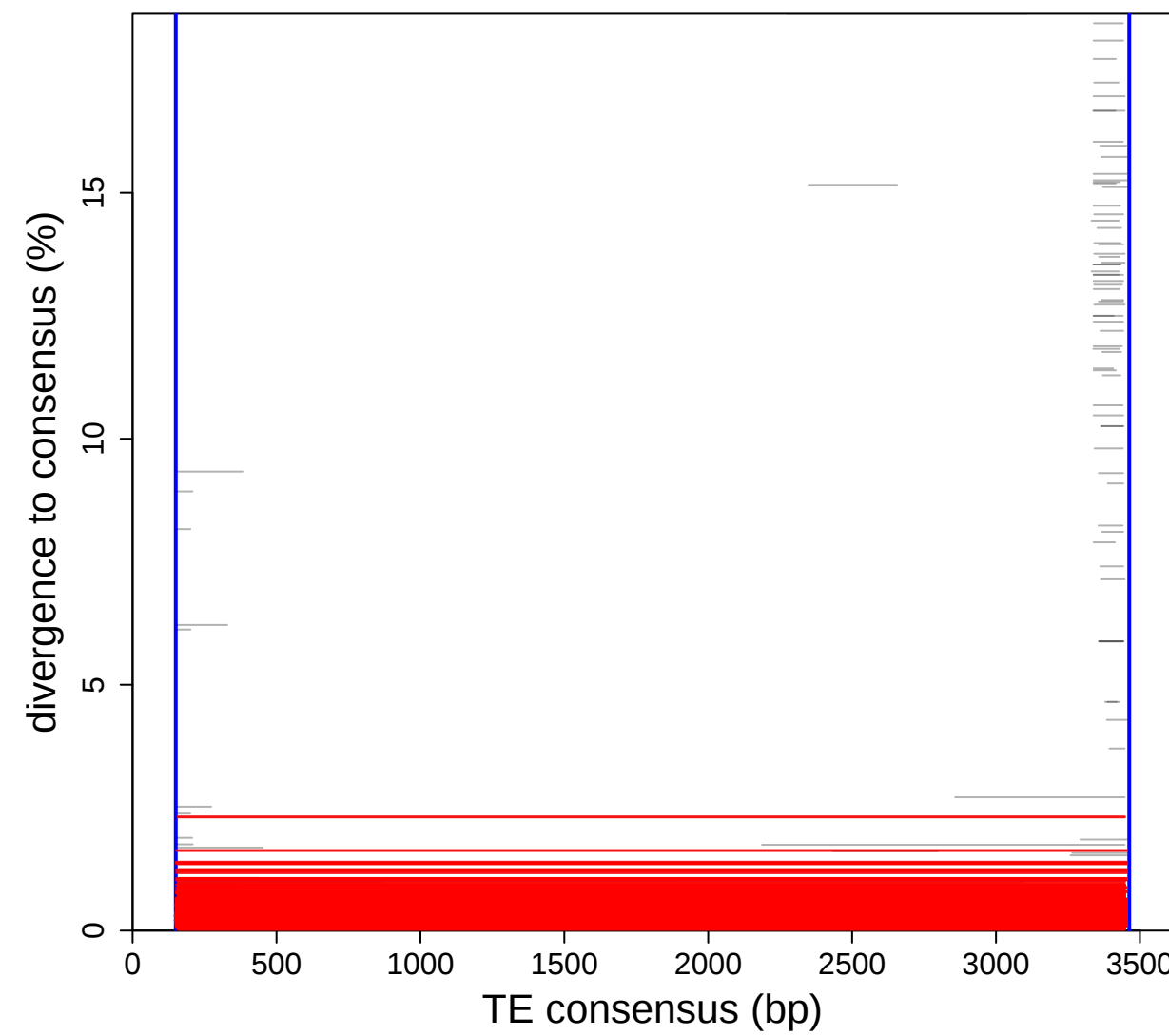
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 3313bp; fragments: 1044; full length: 726 (≥ 2981.7 bp)

After TEtrimmer 3313 bp



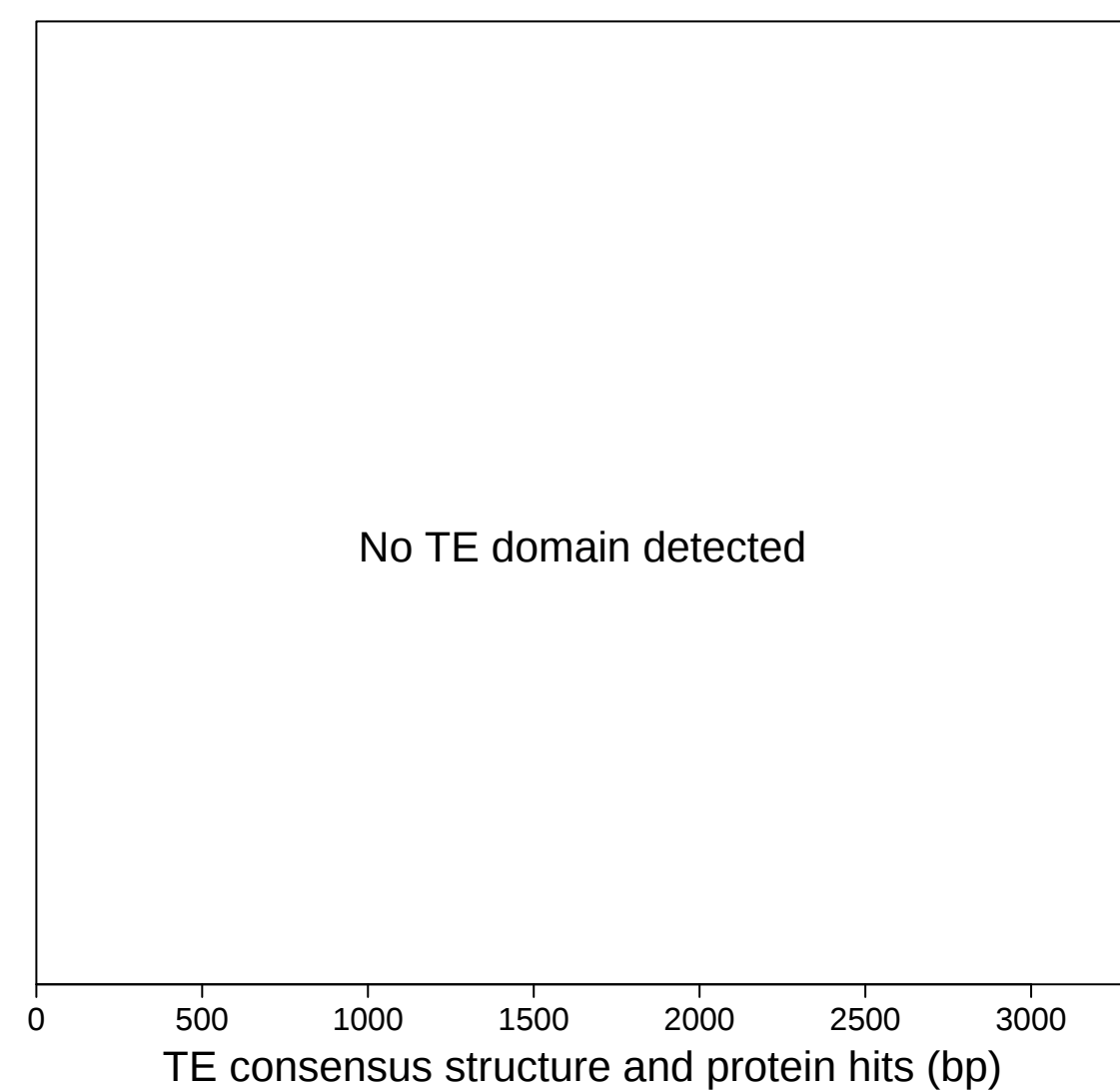
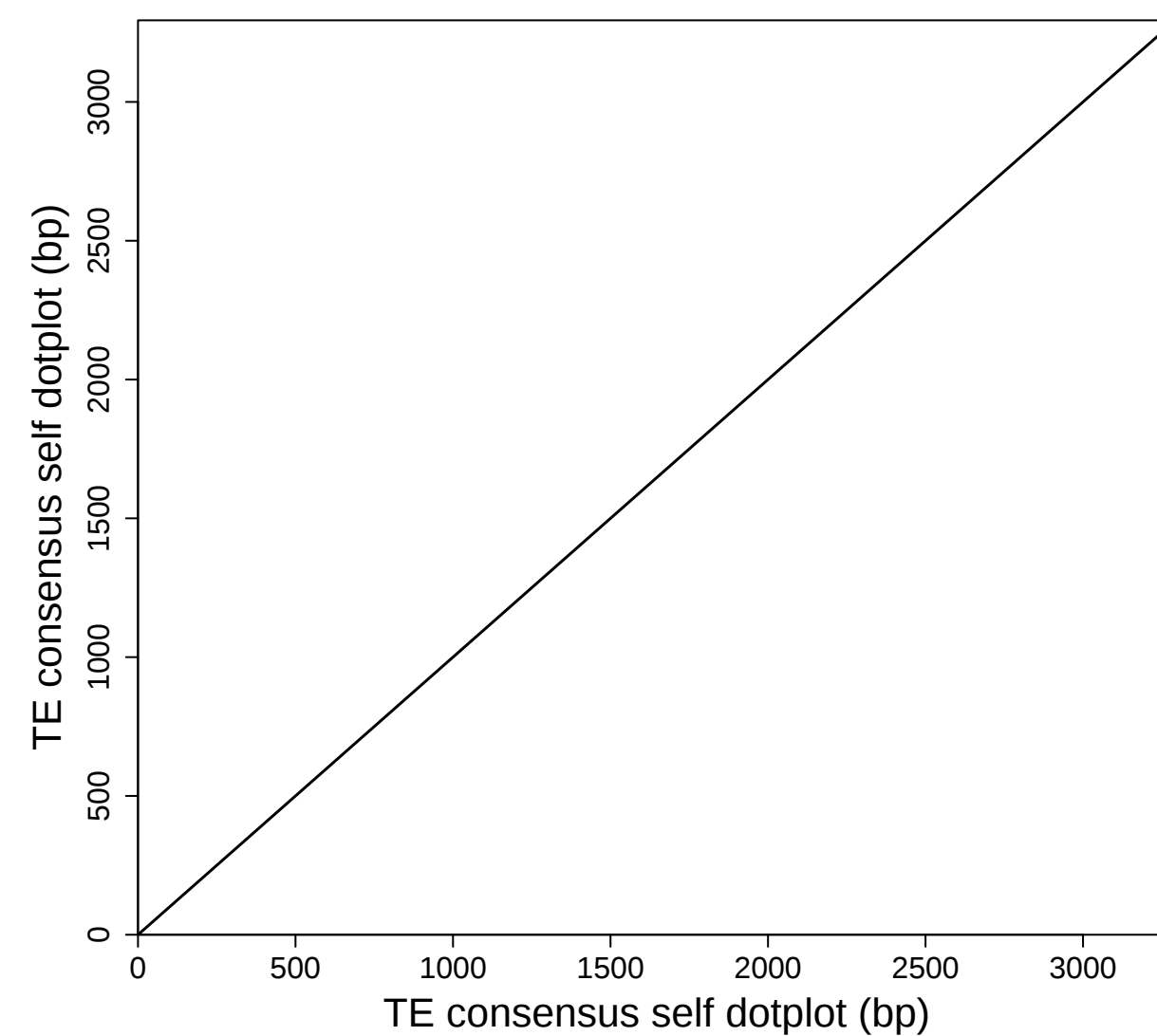
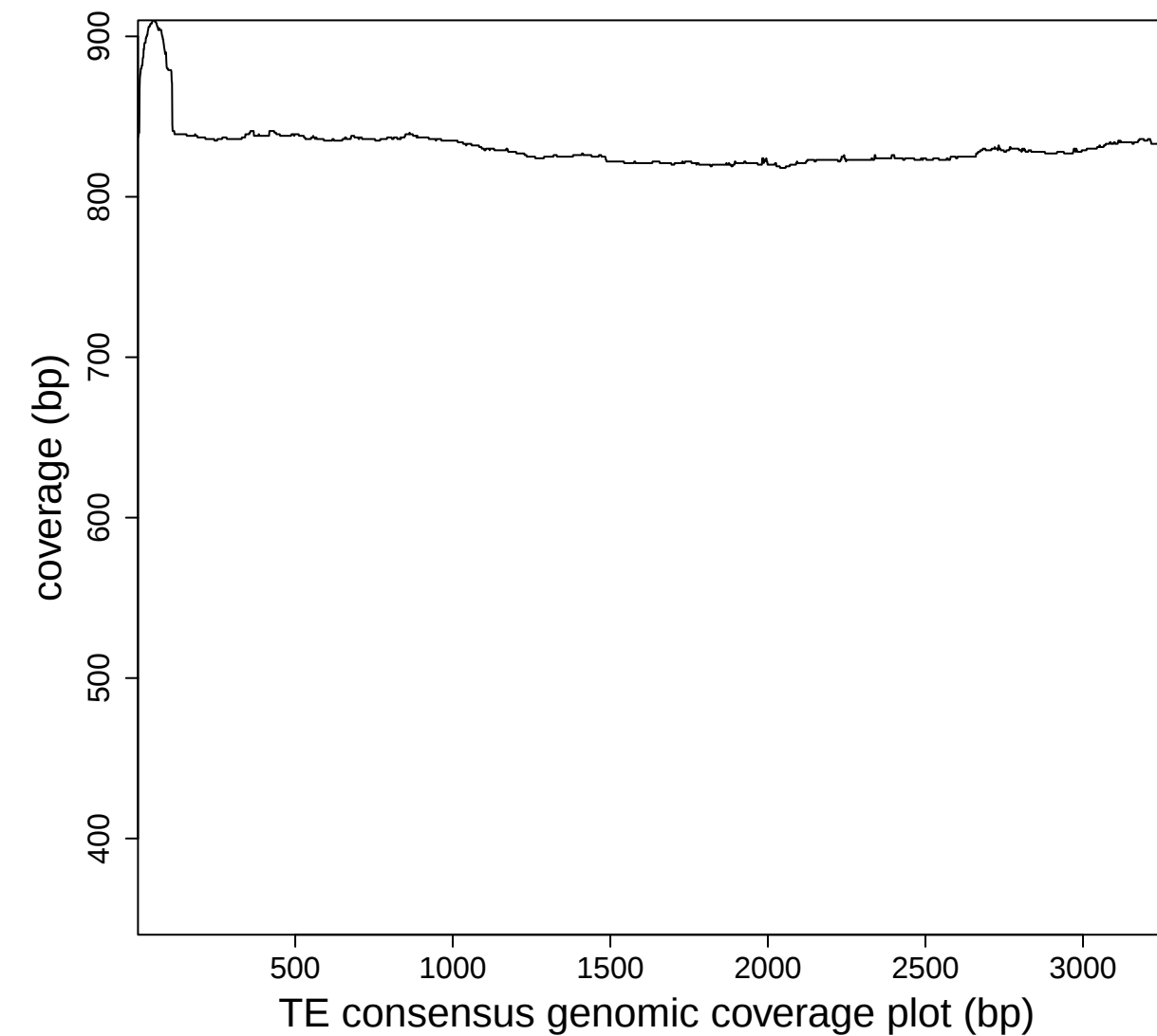
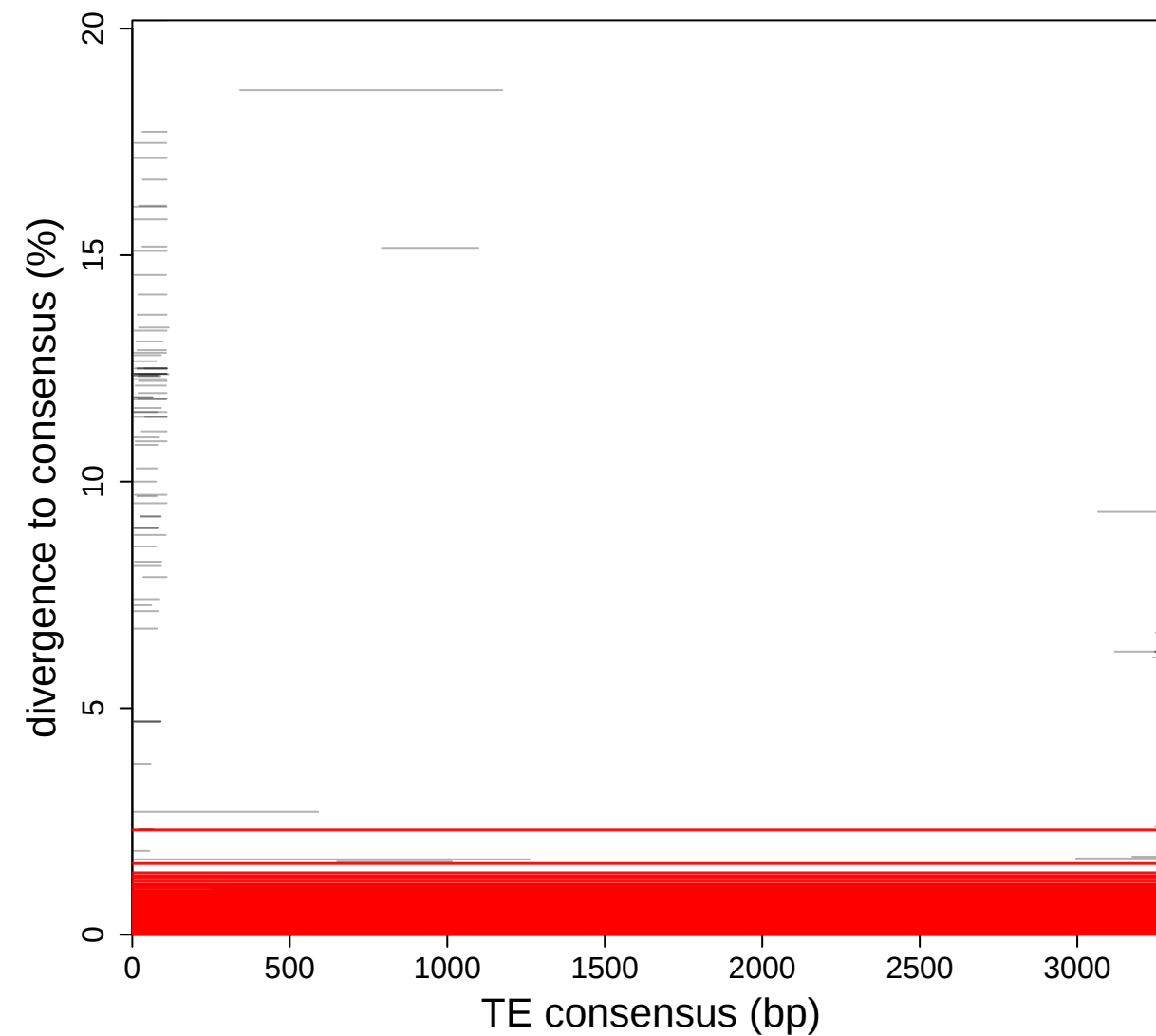
_390.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_c
size: 3613bp; fragments: 1043; full length: 708 (>=3251.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

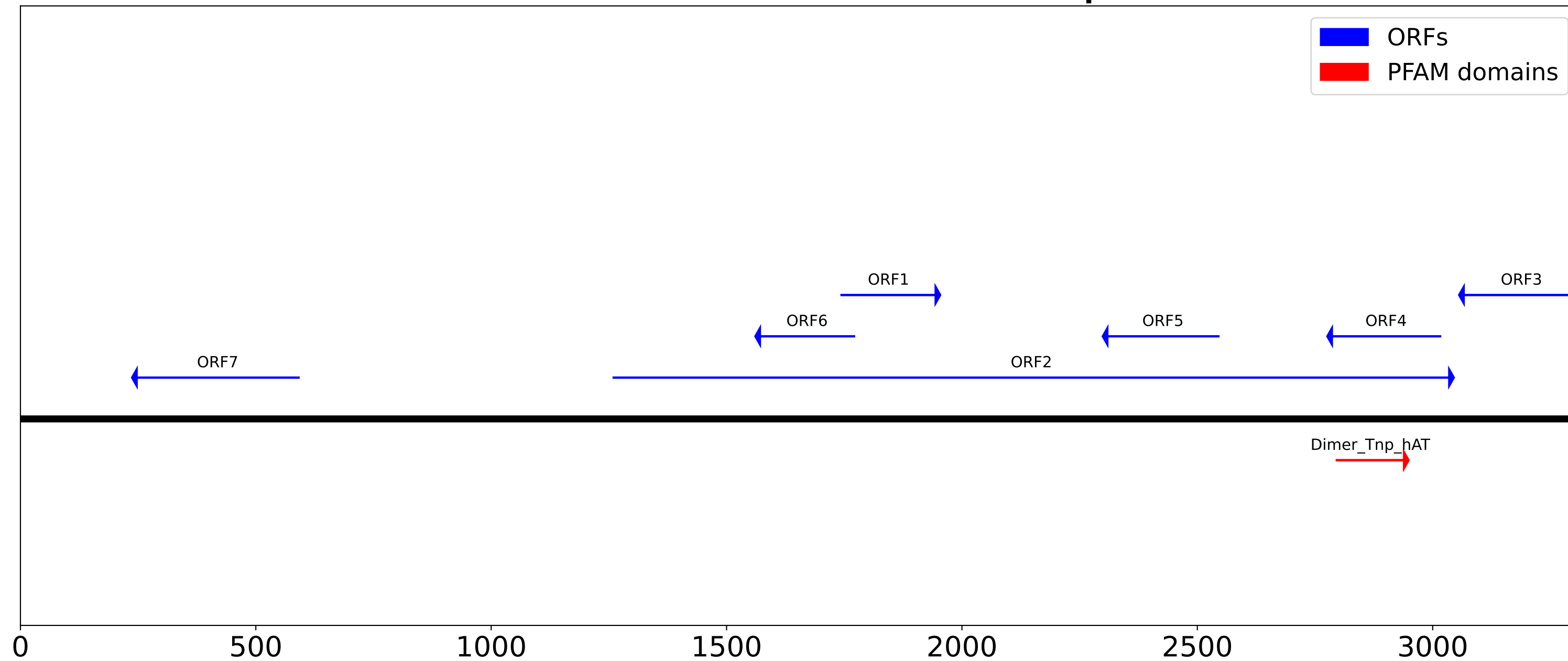


TE: rnd_1_family_390
size: 3294bp; fragments: 1053; full length: 729 (≥ 2964.6 bp)

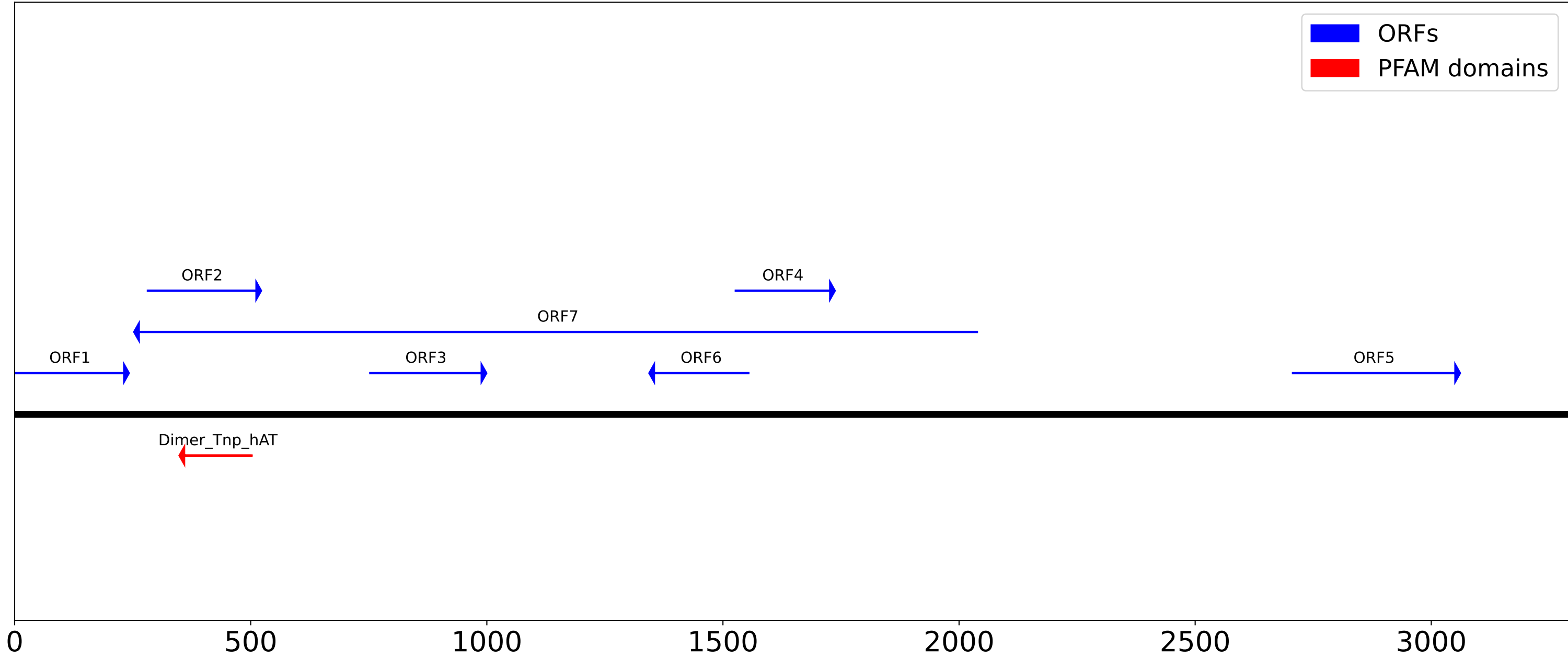
Before TEtrimmer 3294 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

