

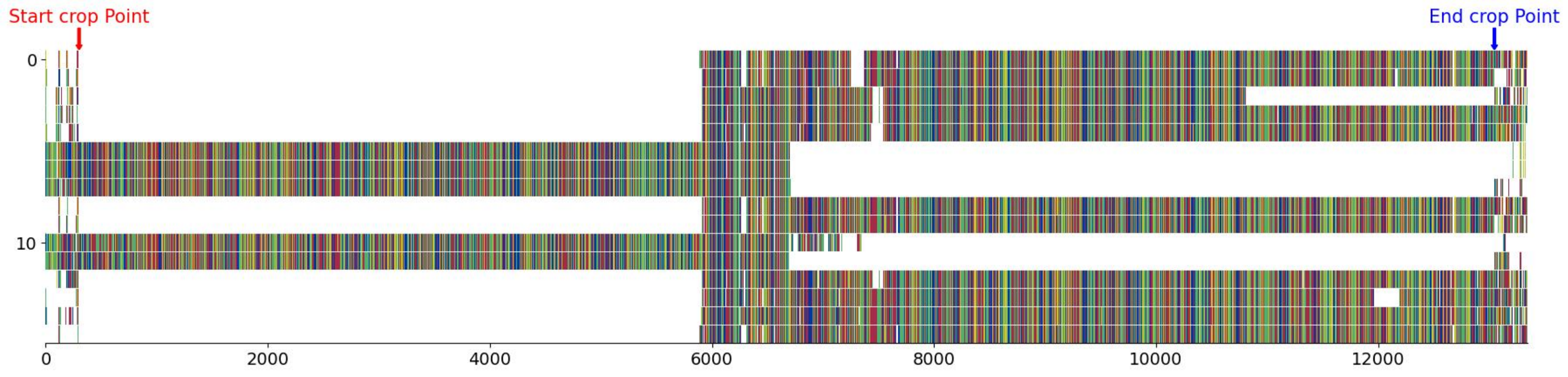
MSA length = 12749

Start crop Point



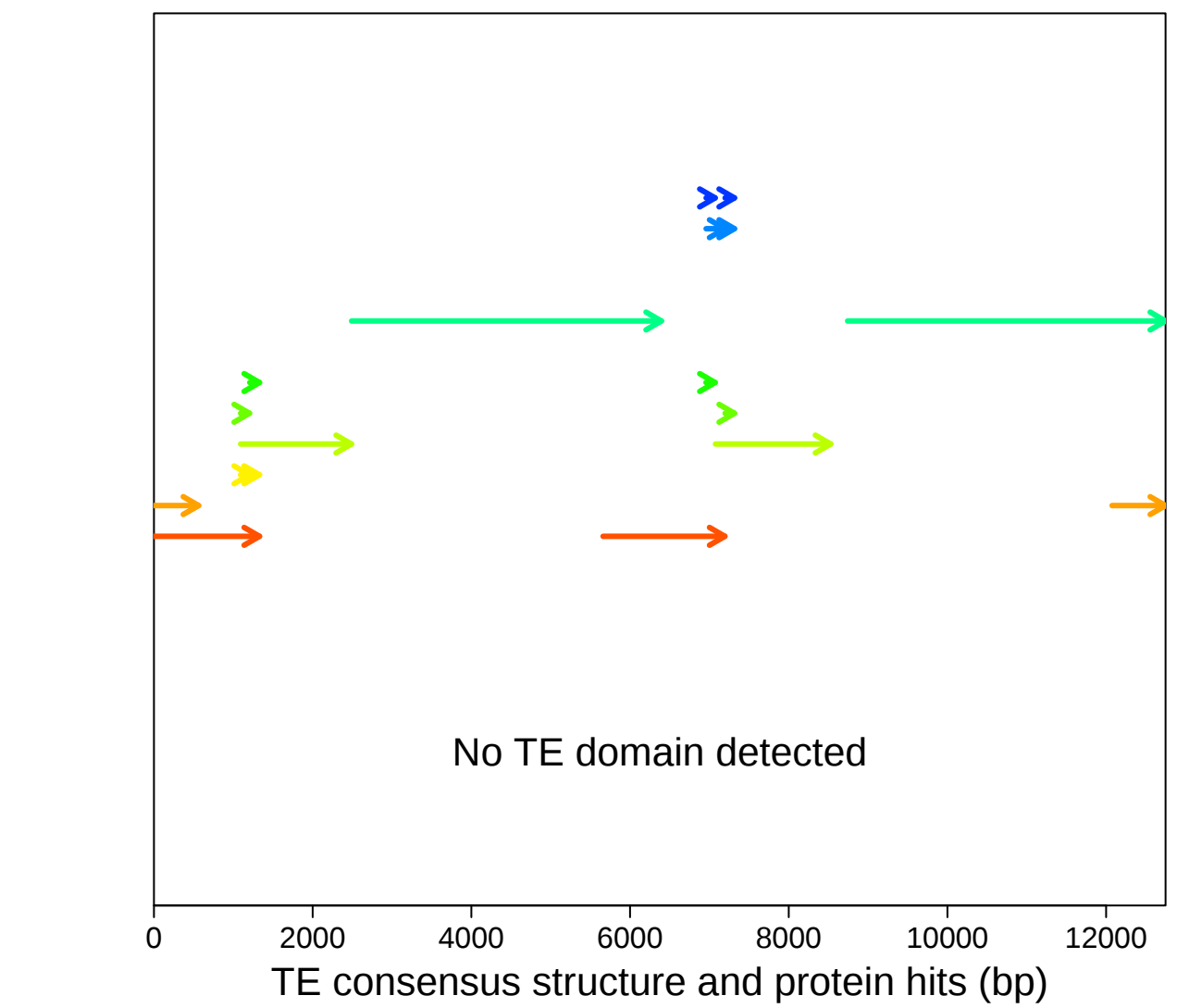
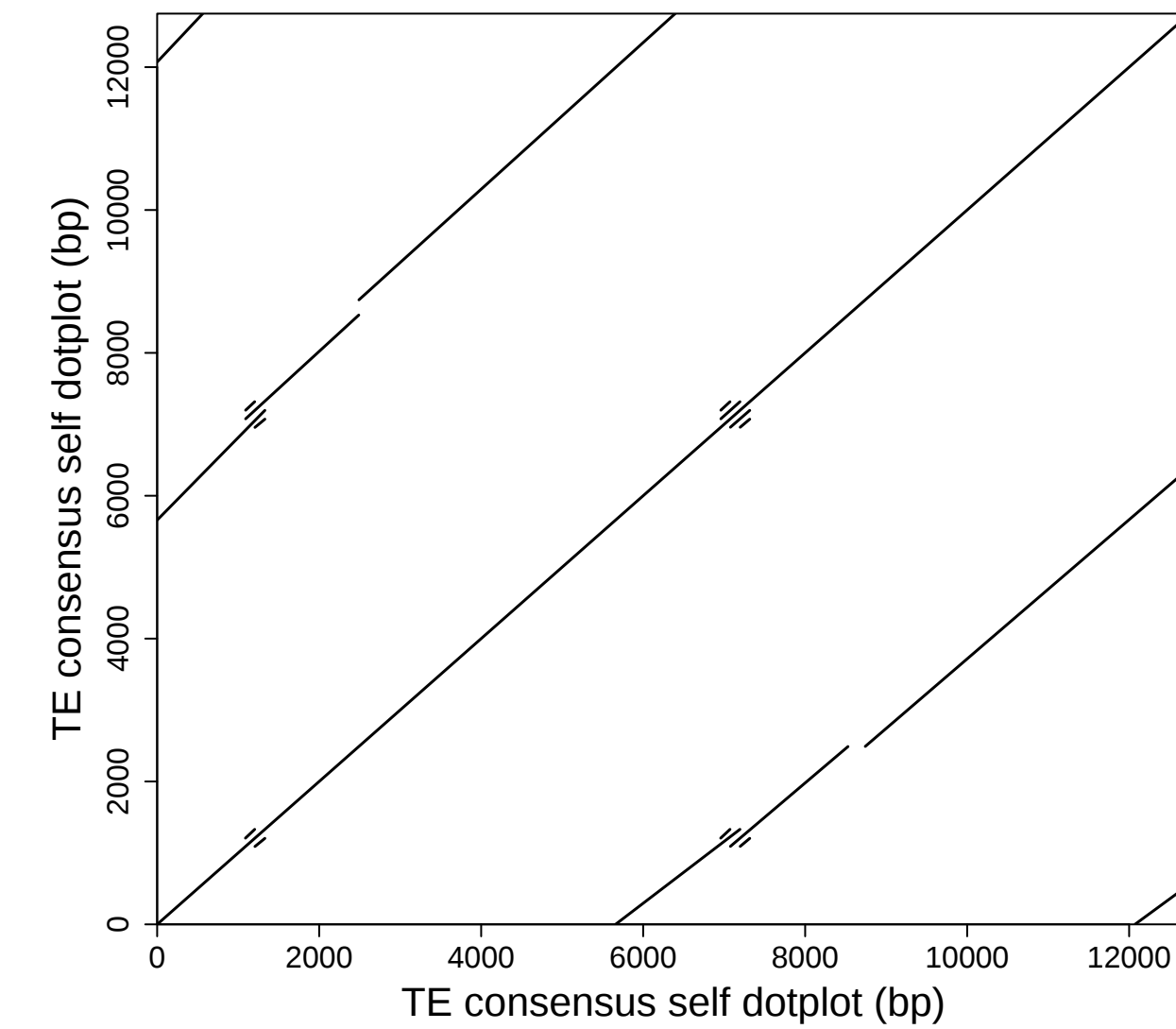
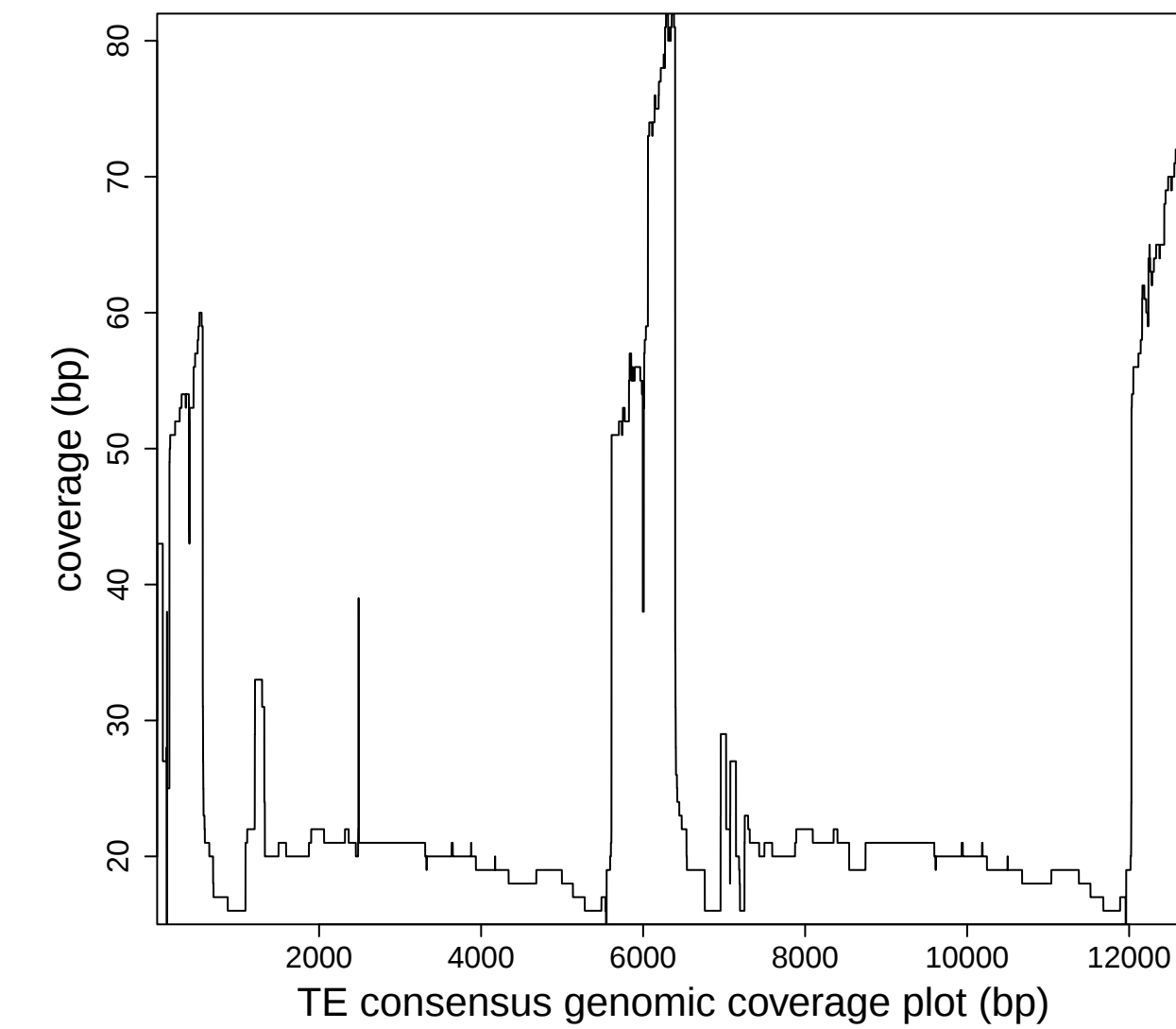
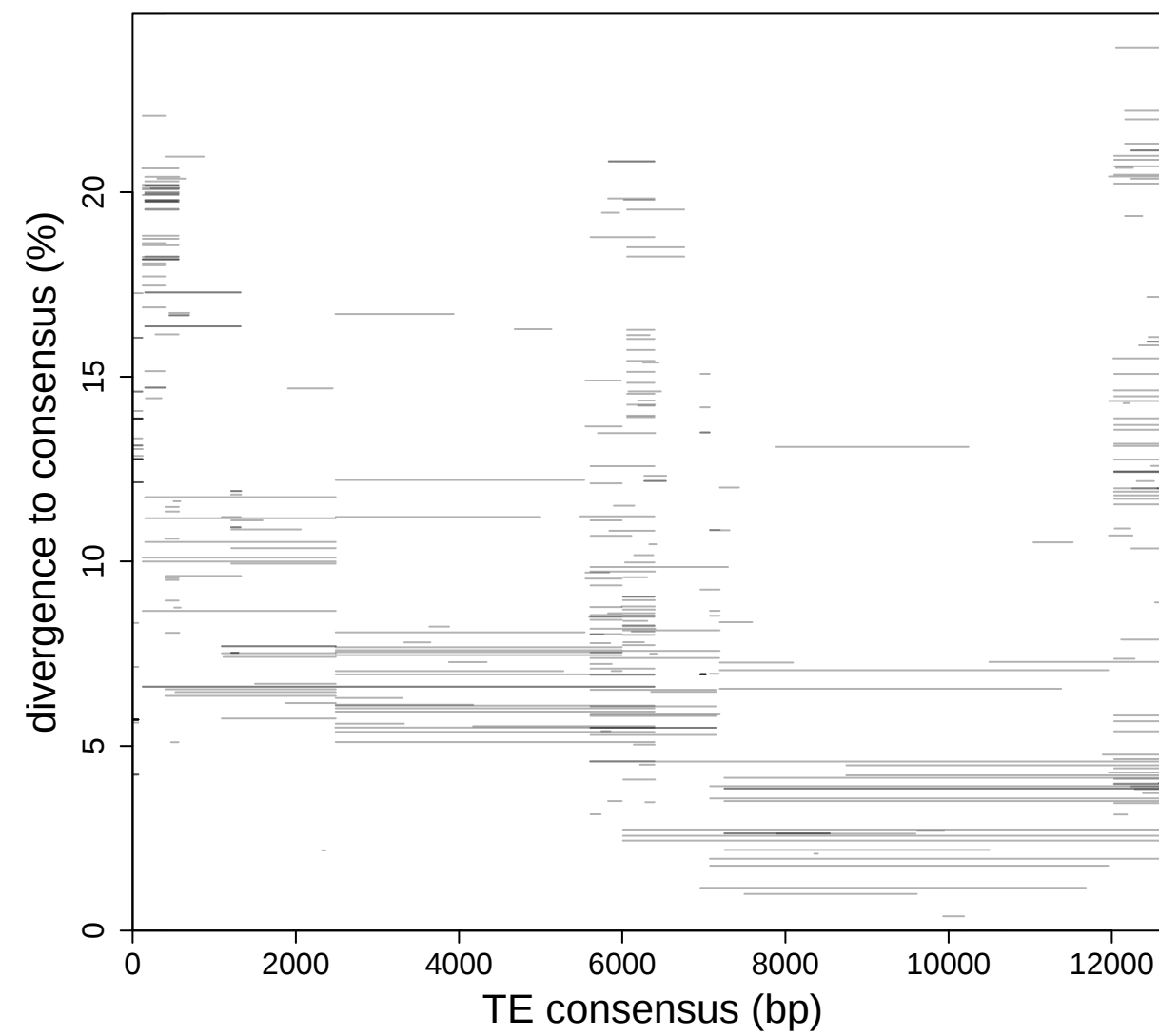
End crop Point





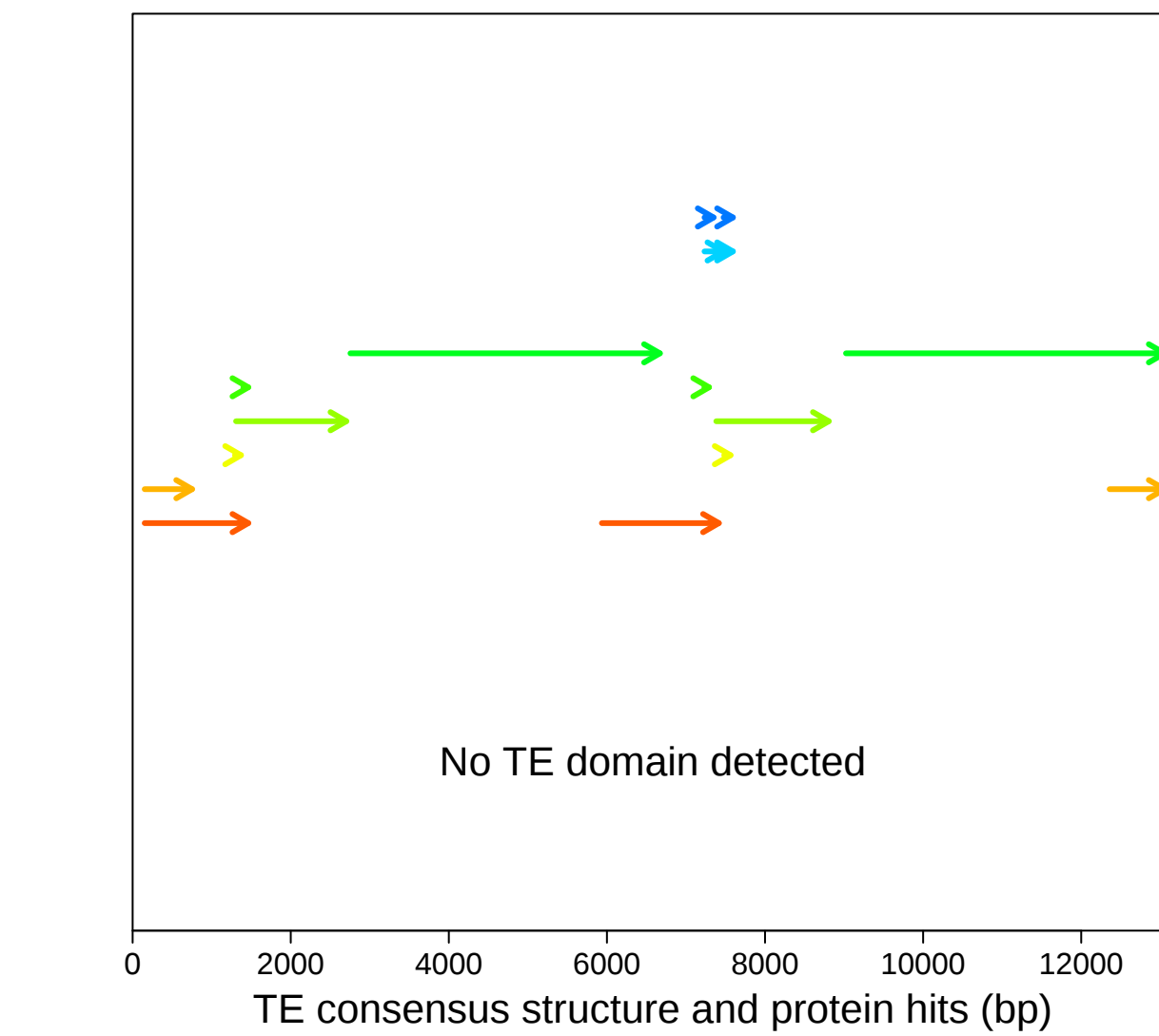
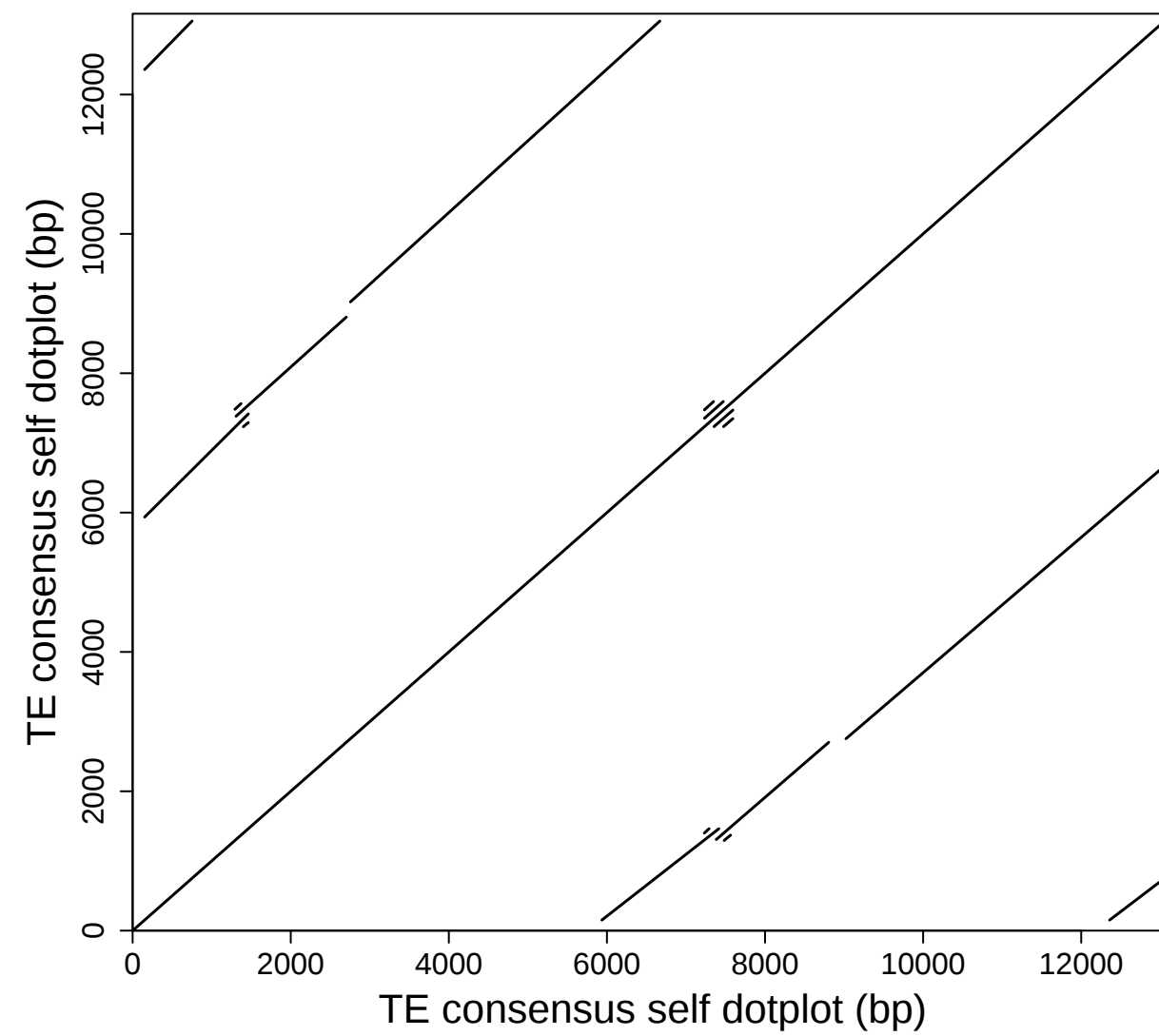
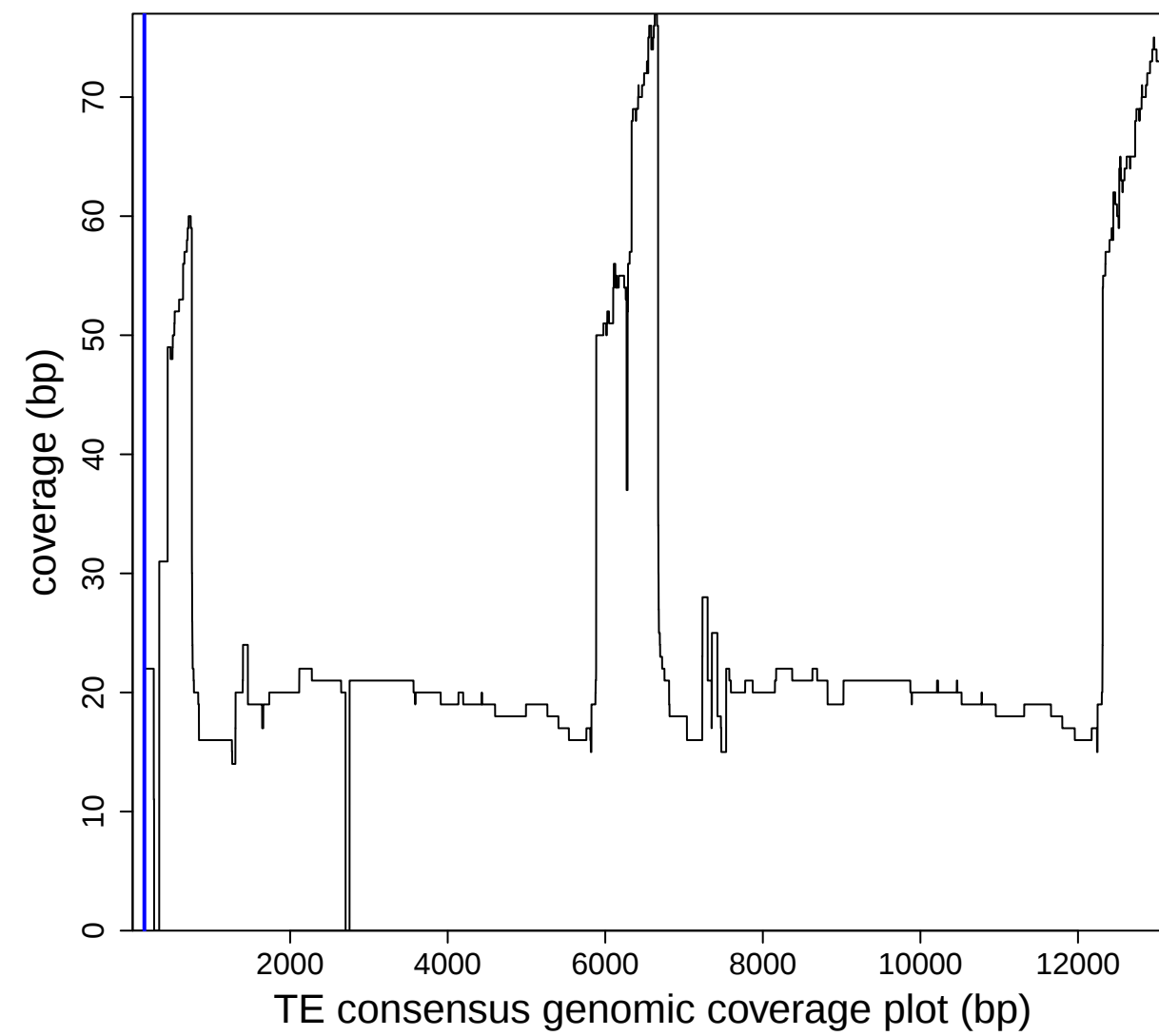
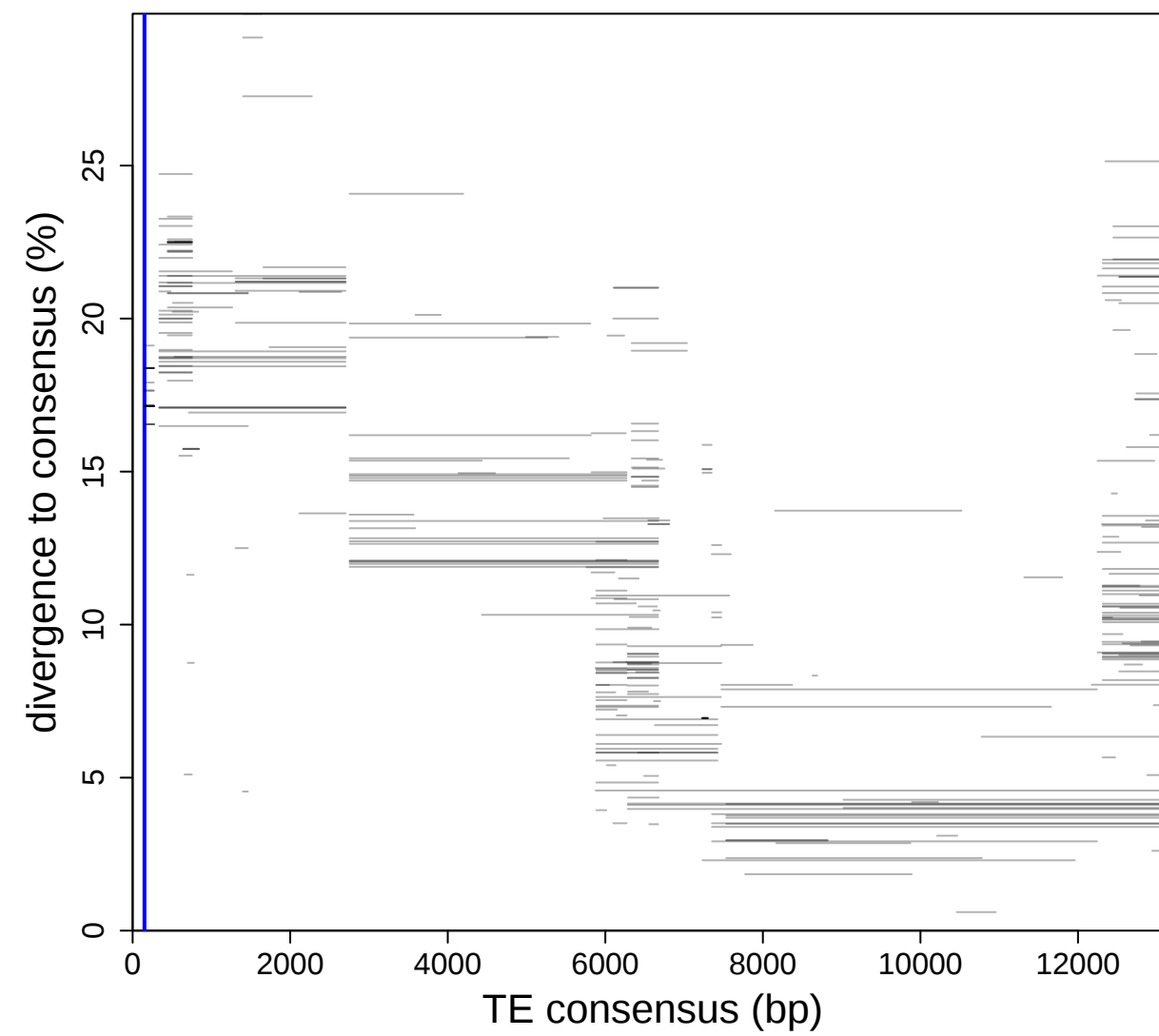
bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.fa
size: 12749bp; fragments: 373; full length: 0 (>=11474.1bp)

After TEtrimmer 12749 bp

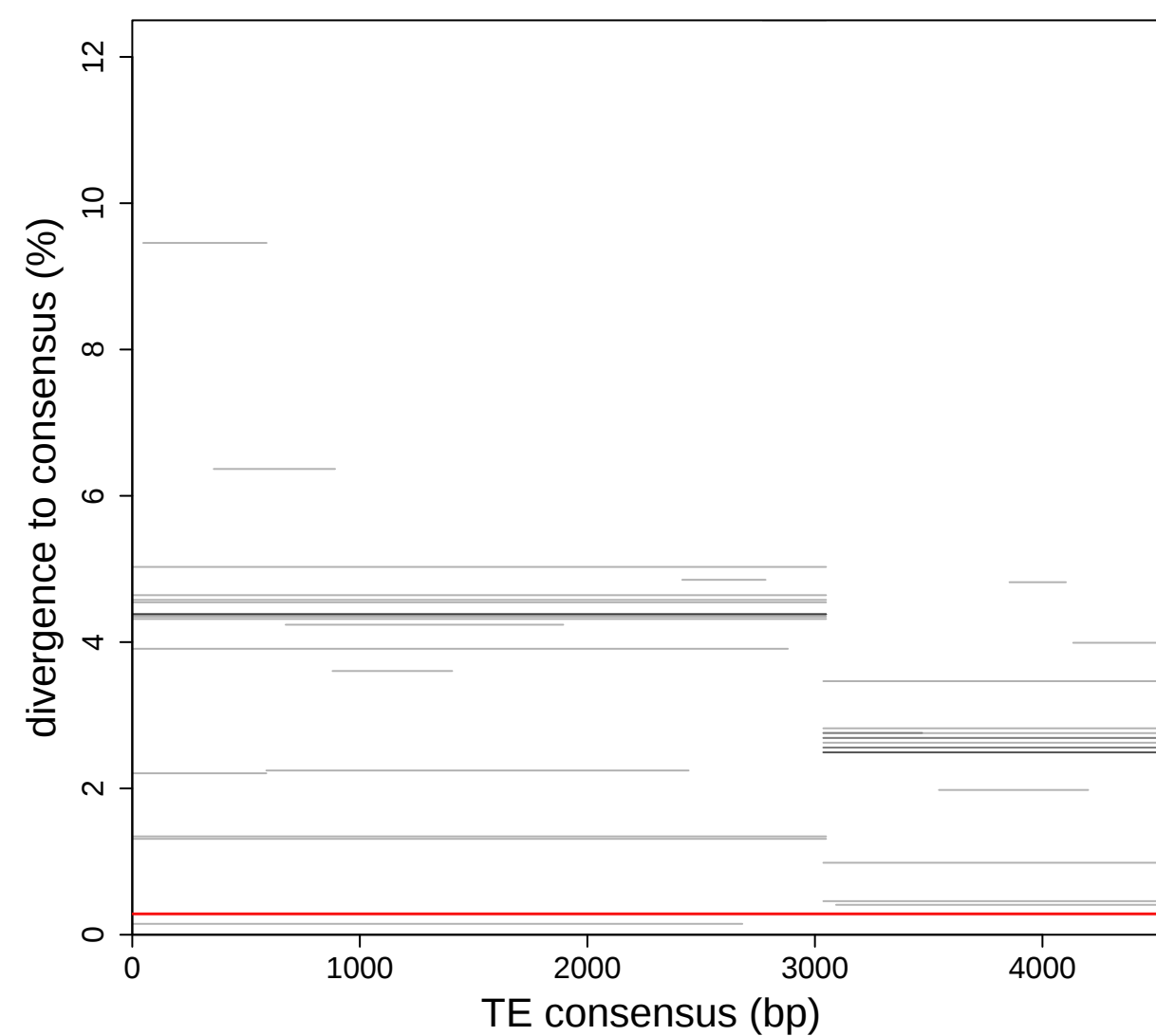


ed_uf.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0
size: 13205bp; fragments: 329; full length: 0 (>=11884.5bp)

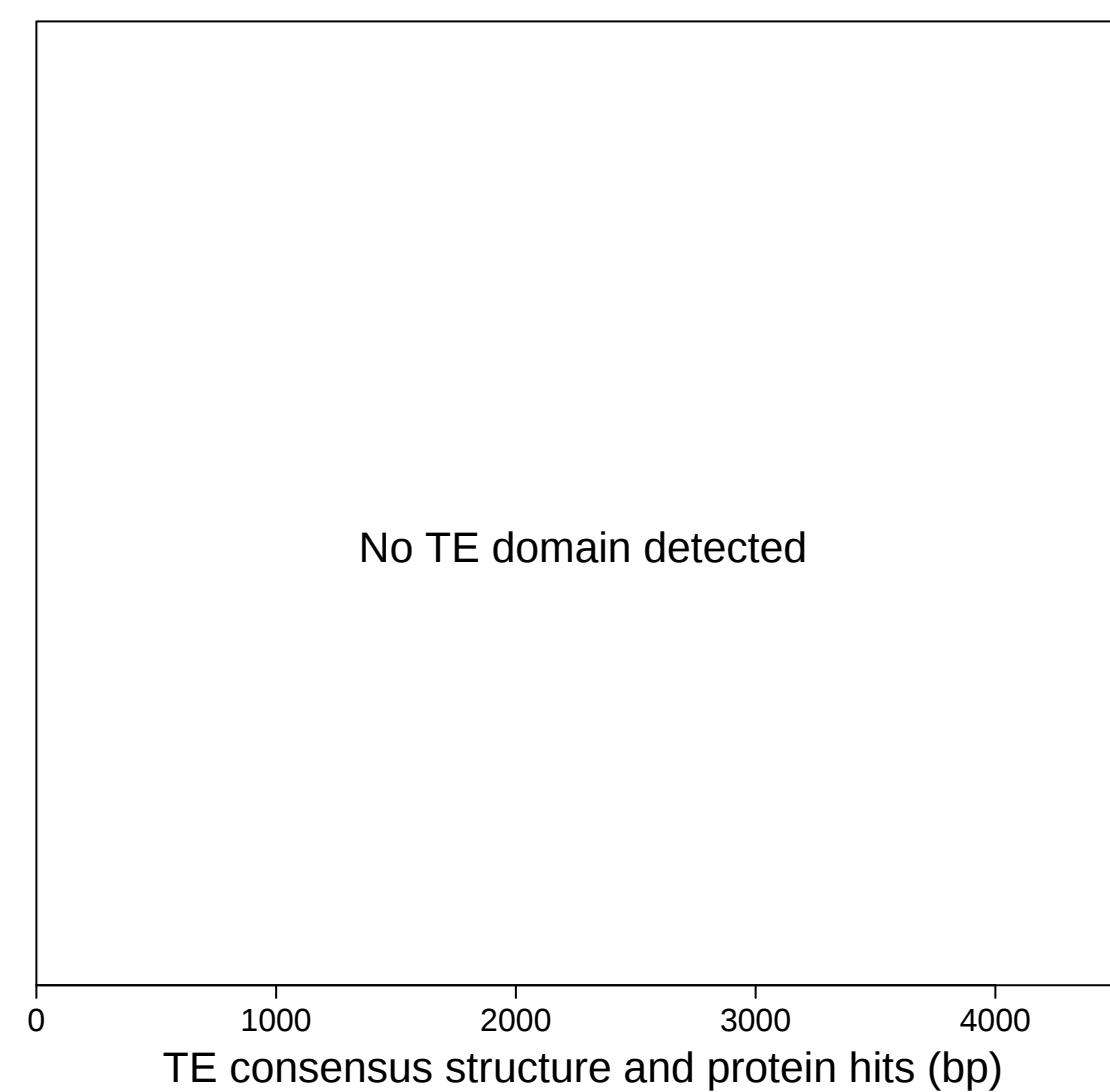
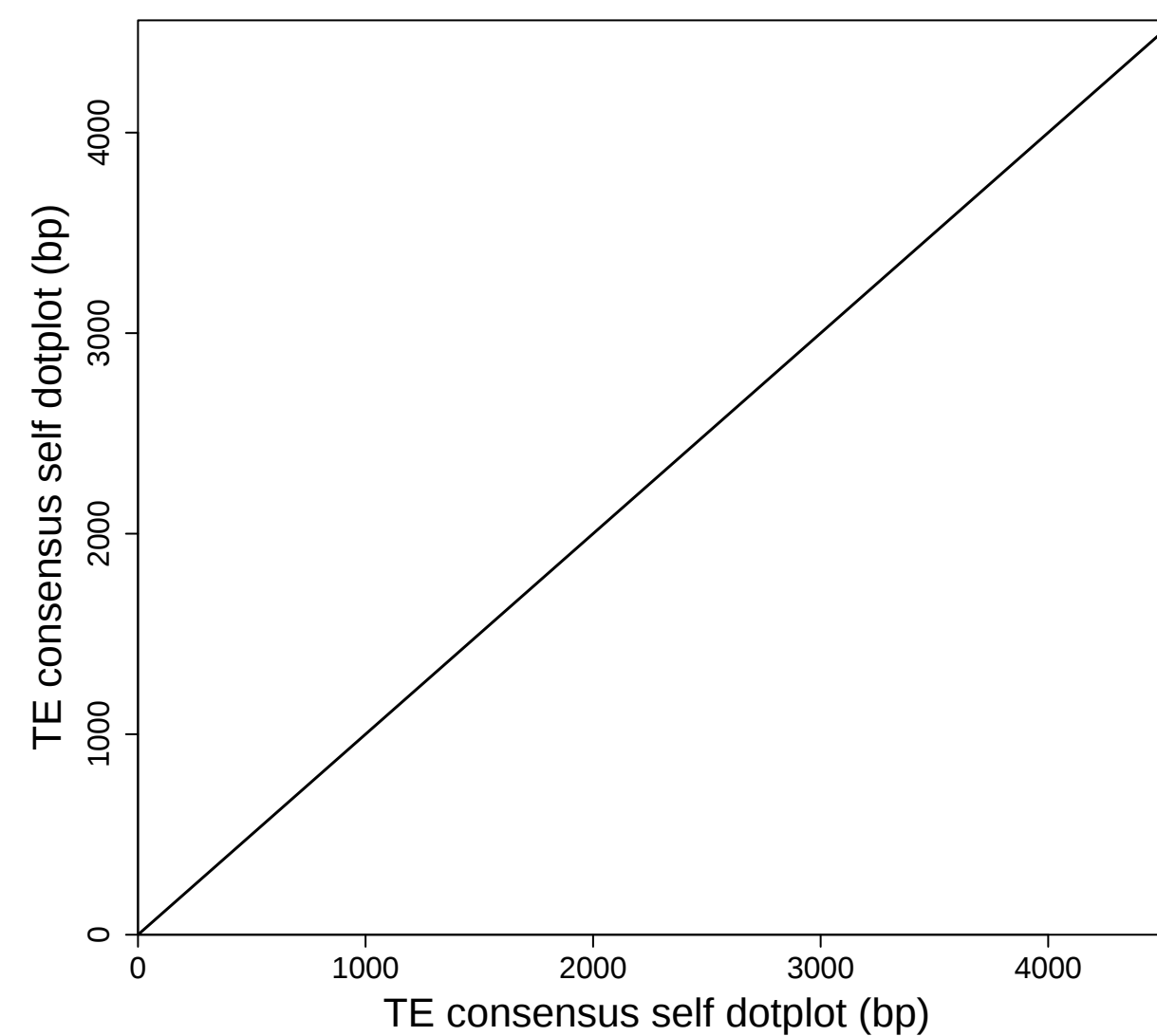
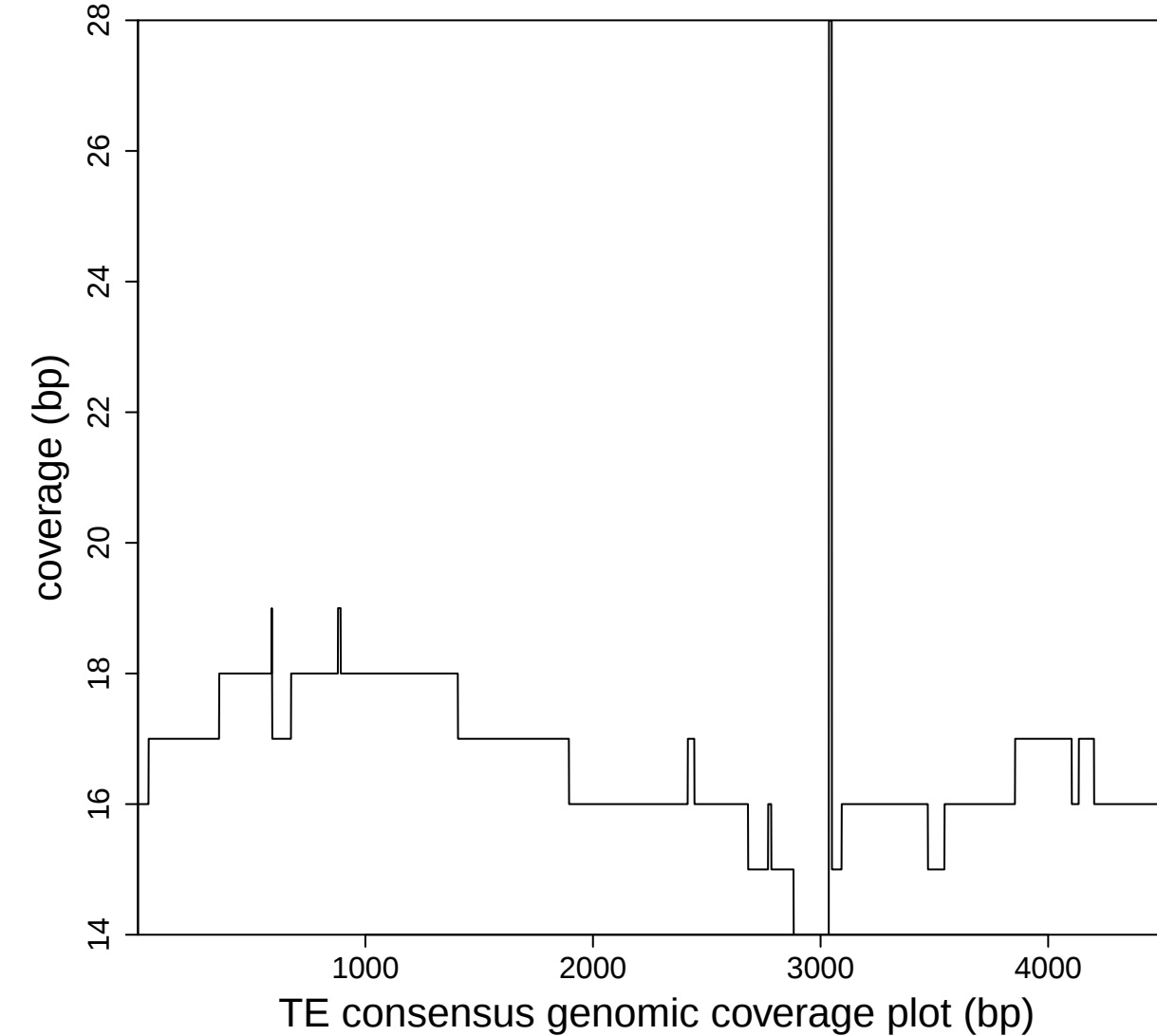
After TEtrimmer Extended plot Blue lines are boundaries



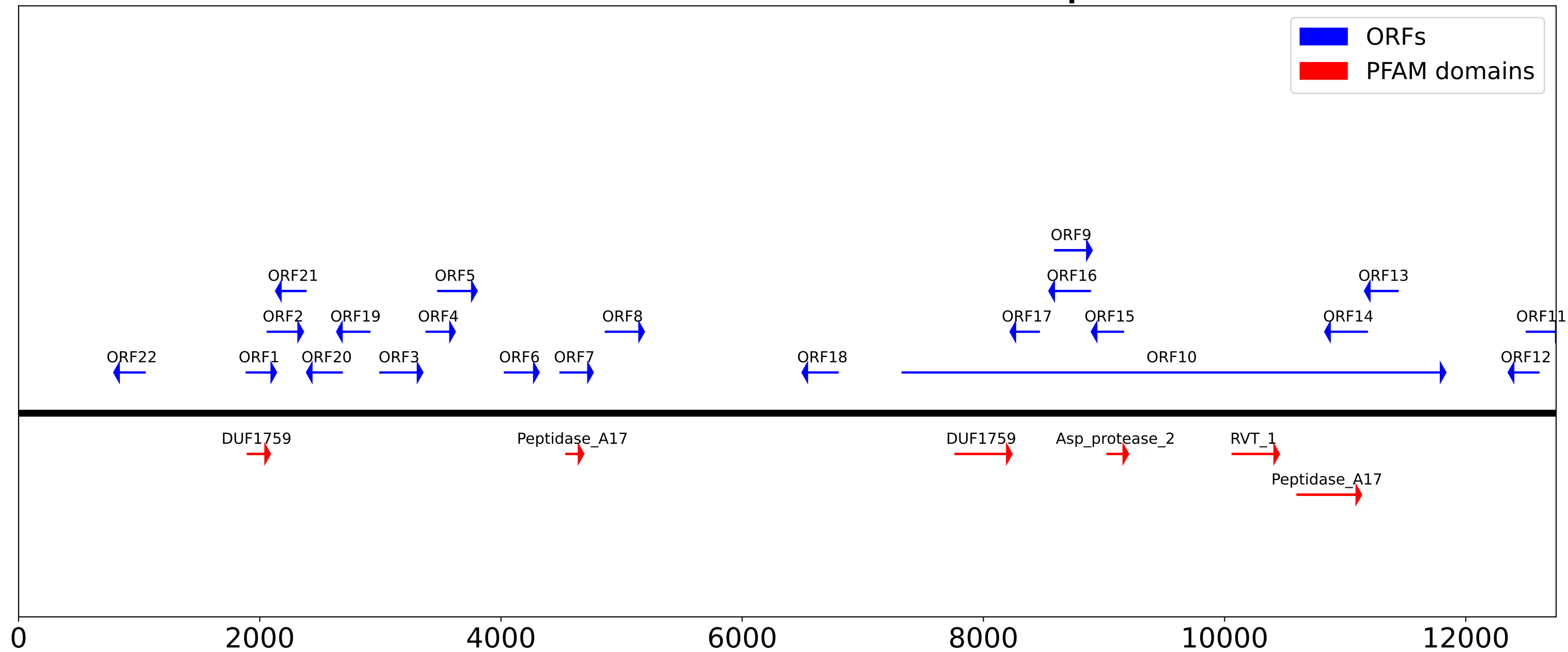
TE: ltr_1_family_549
size: 4560bp; fragments: 41; full length: 1 (≥ 4104 bp)



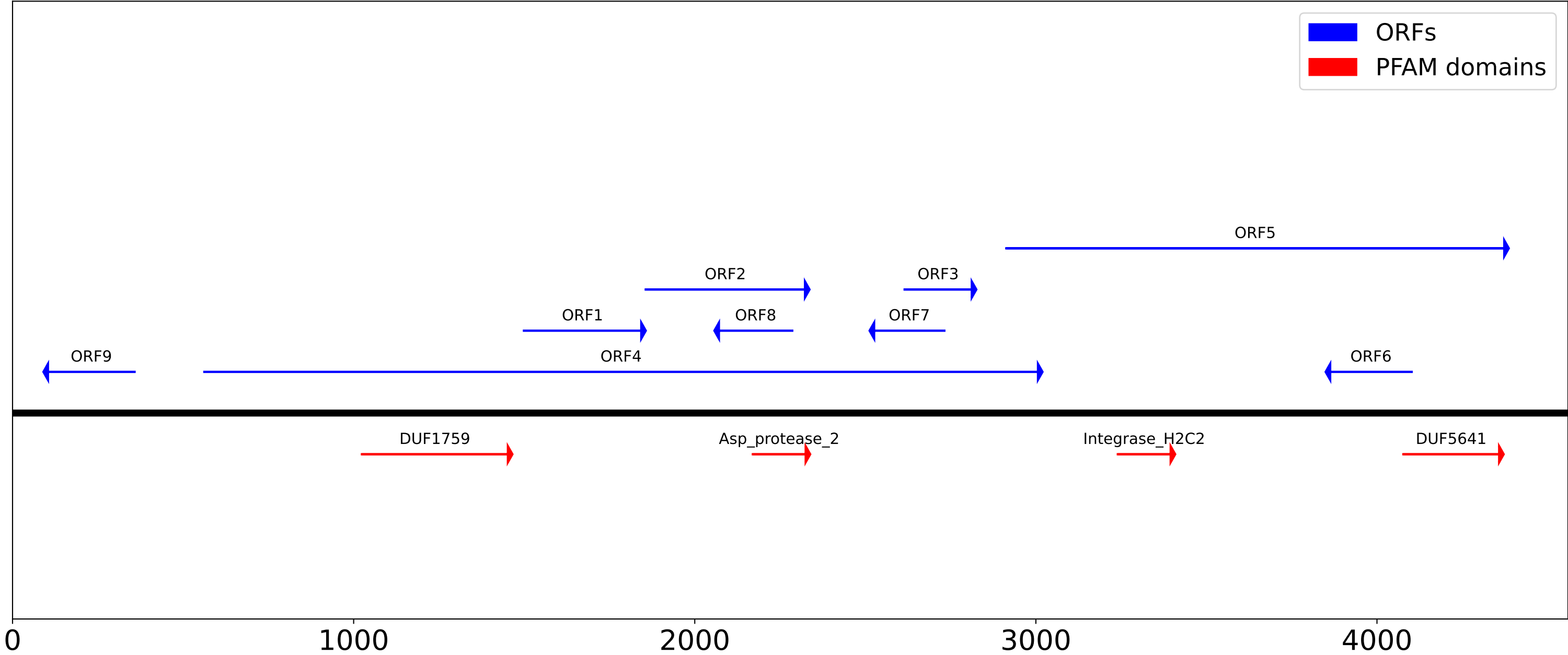
Before TEtrimmer 4560 bp



After TEtrimmer ORF and PFAM domain plot



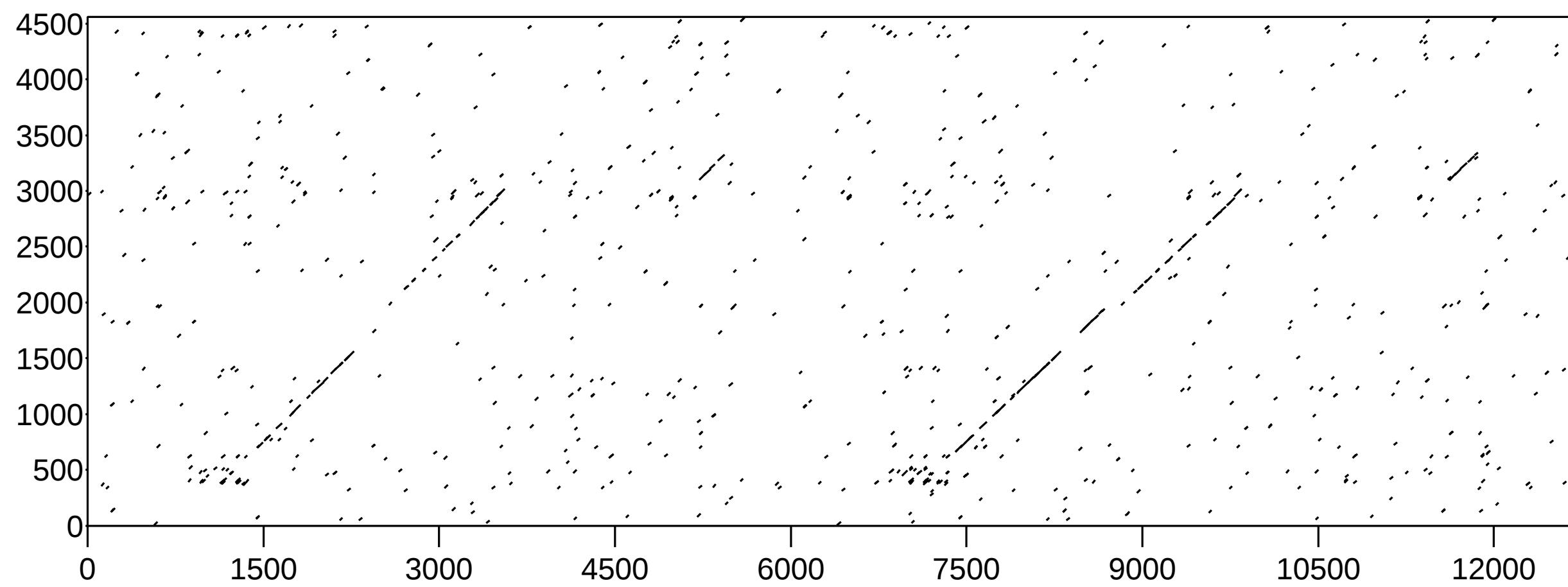
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)