

Start crop Point

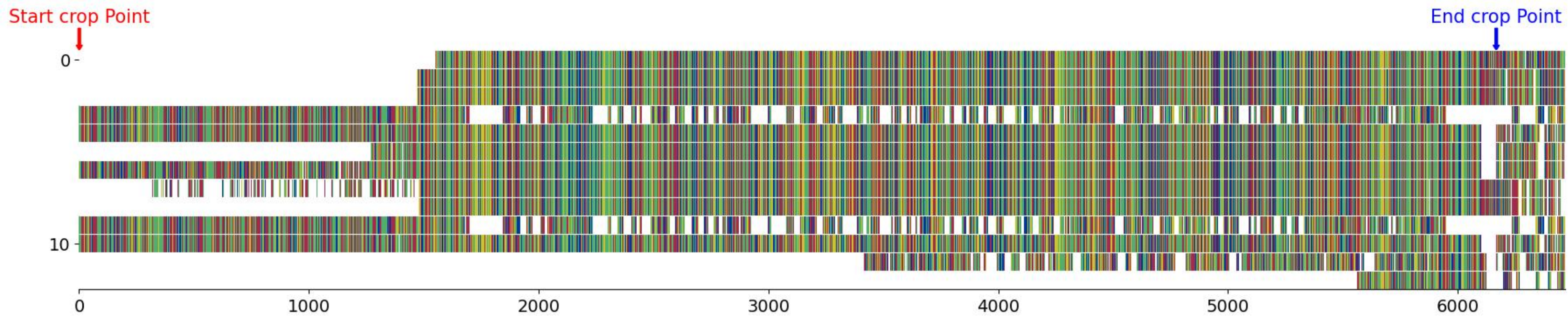


End crop Point



MSA length = 6167

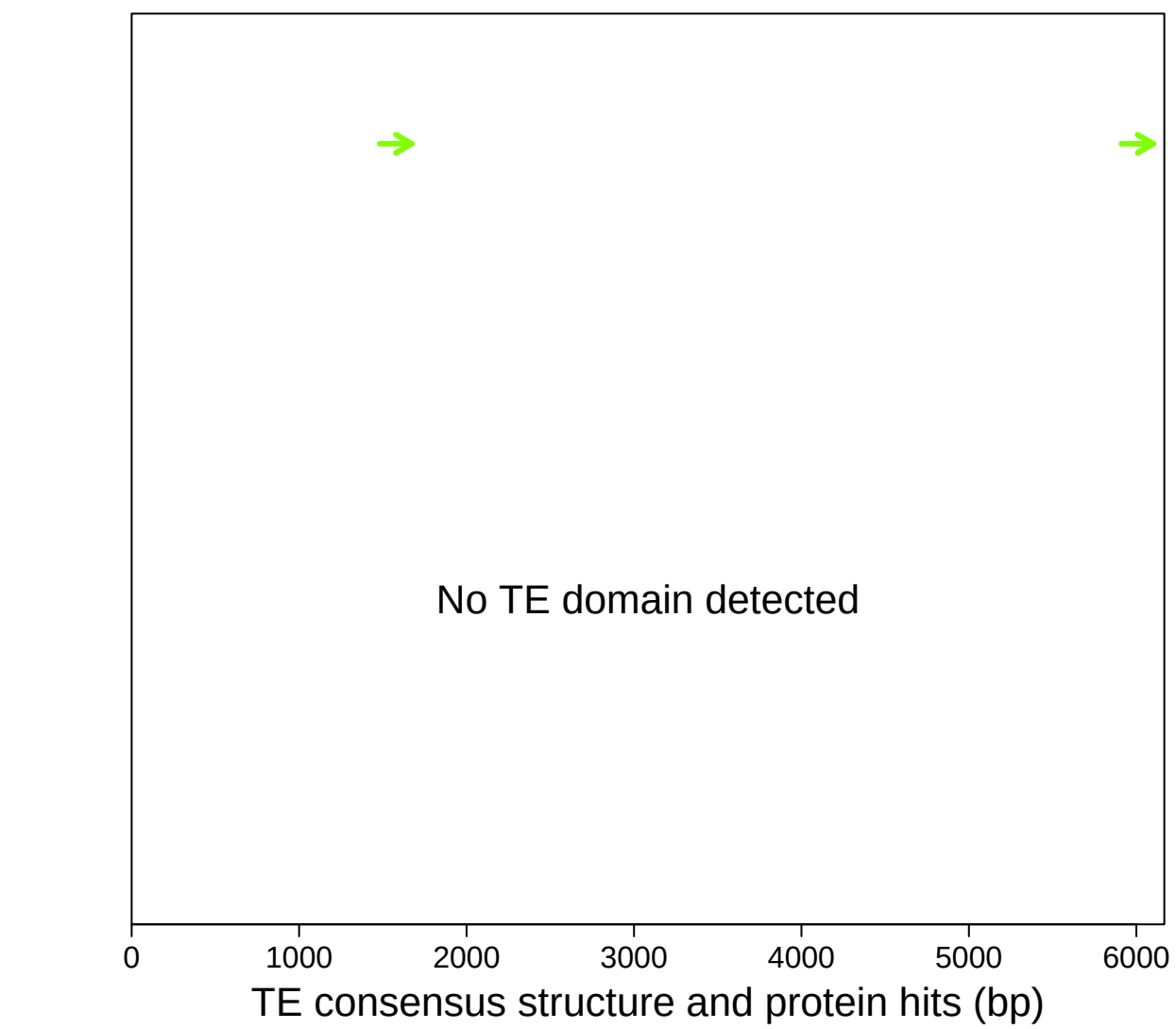
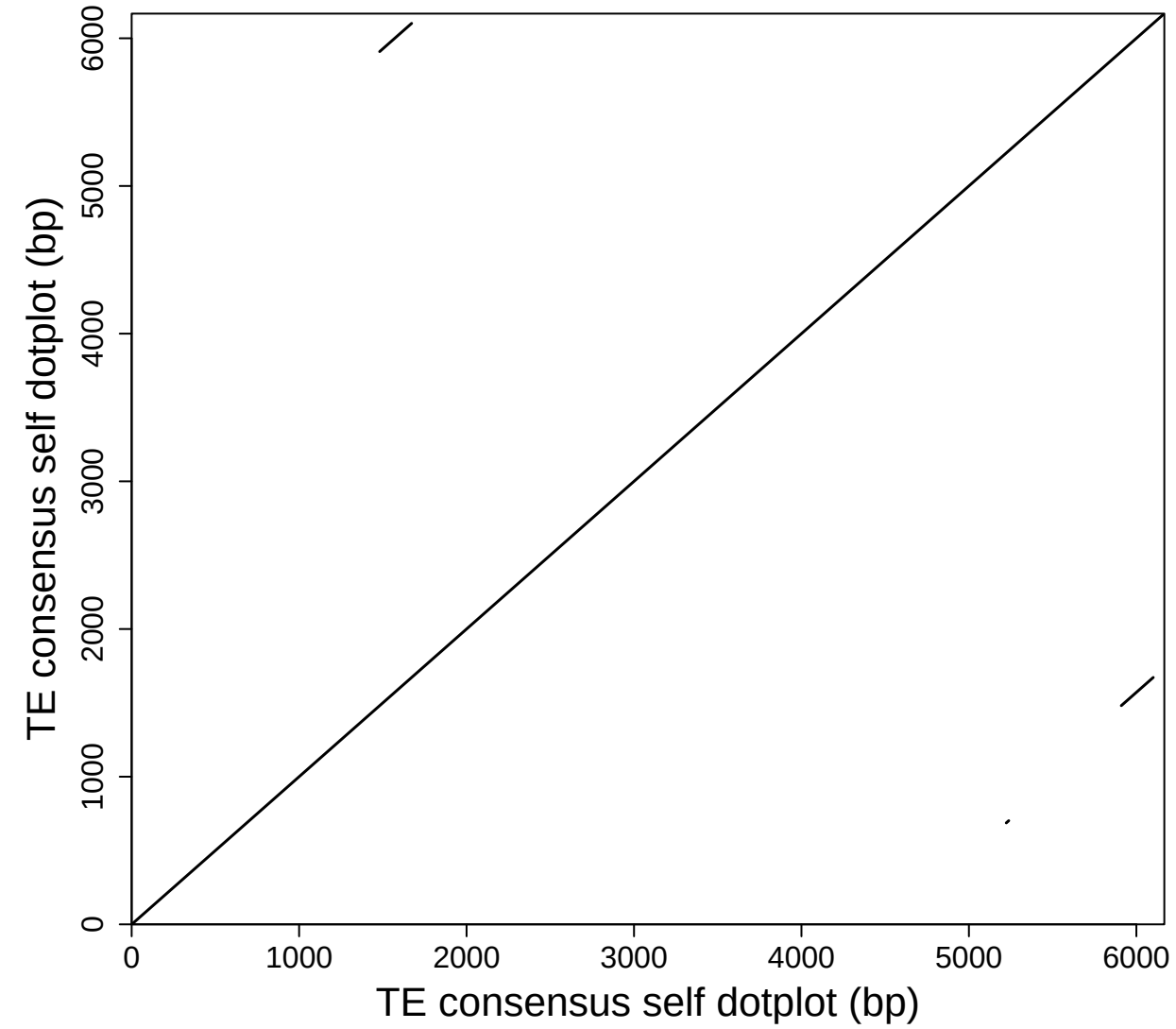
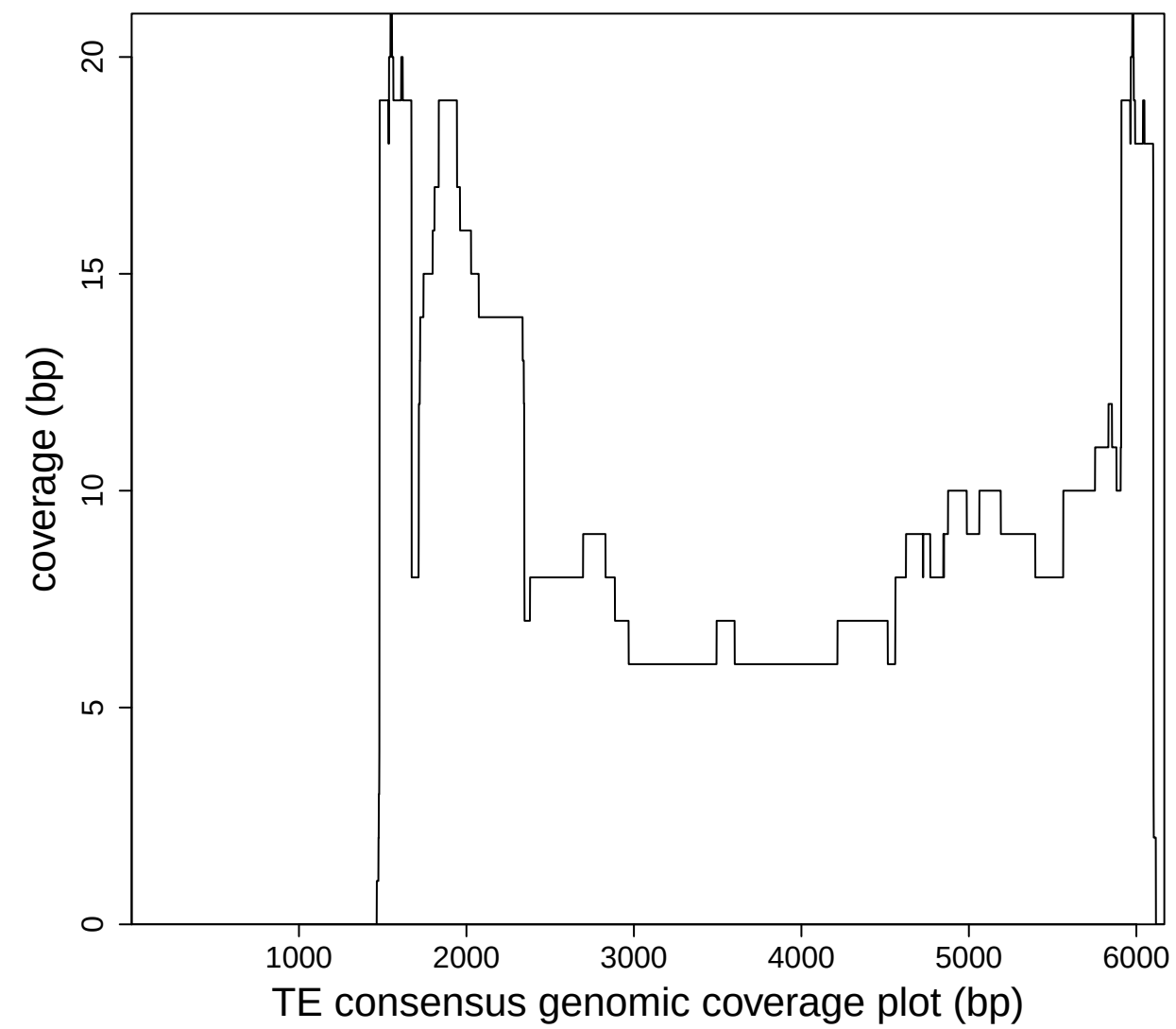
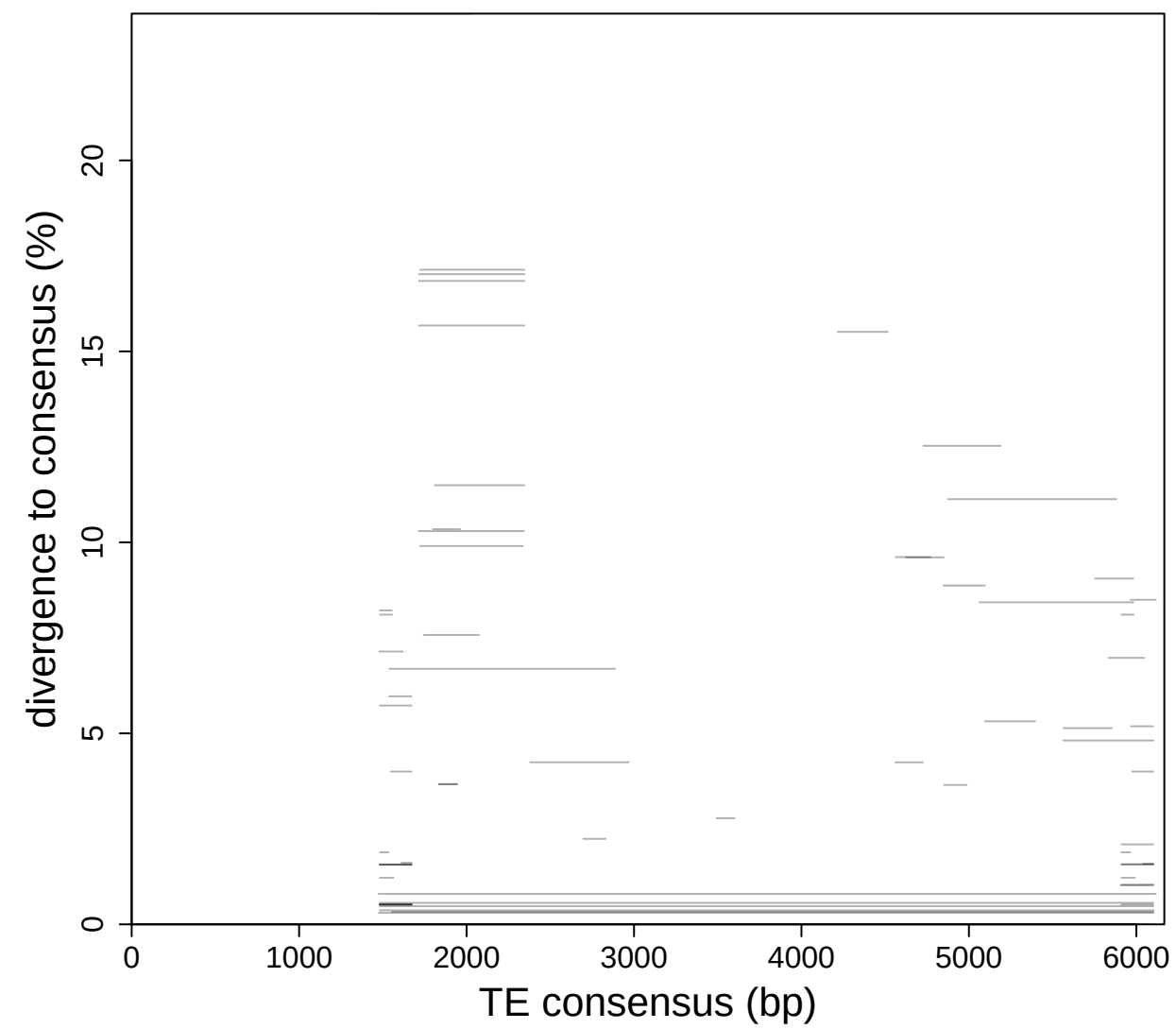




b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_

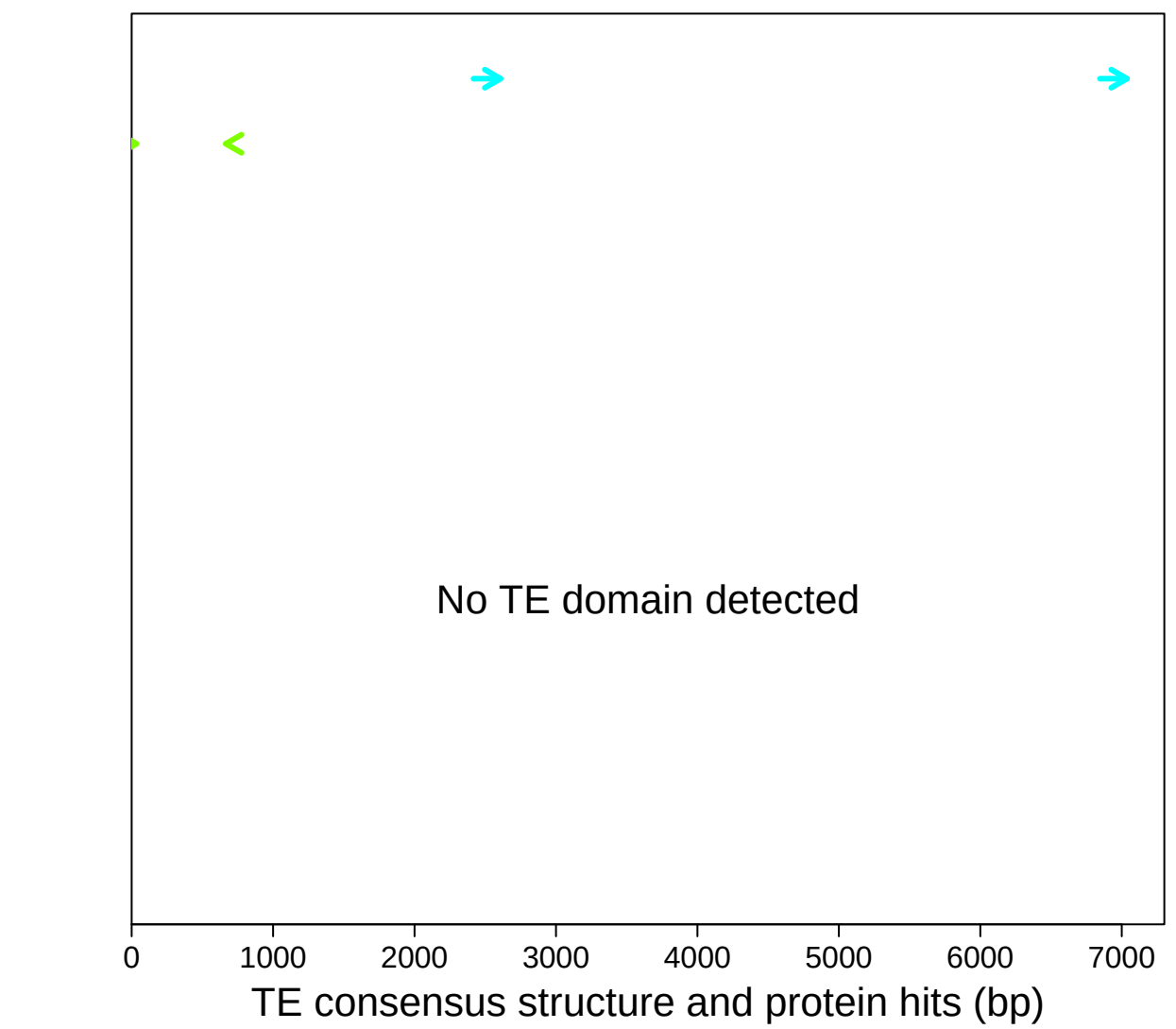
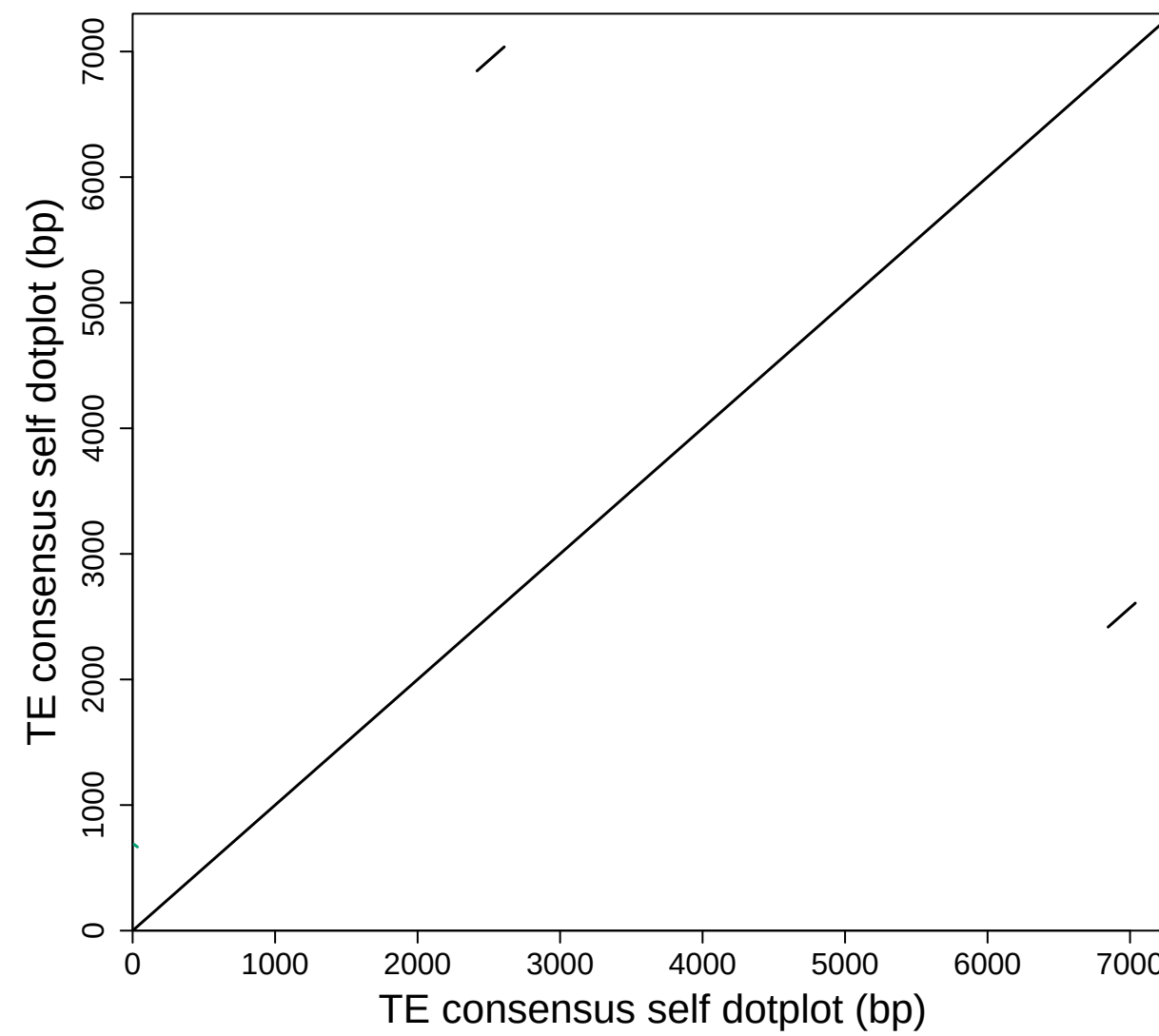
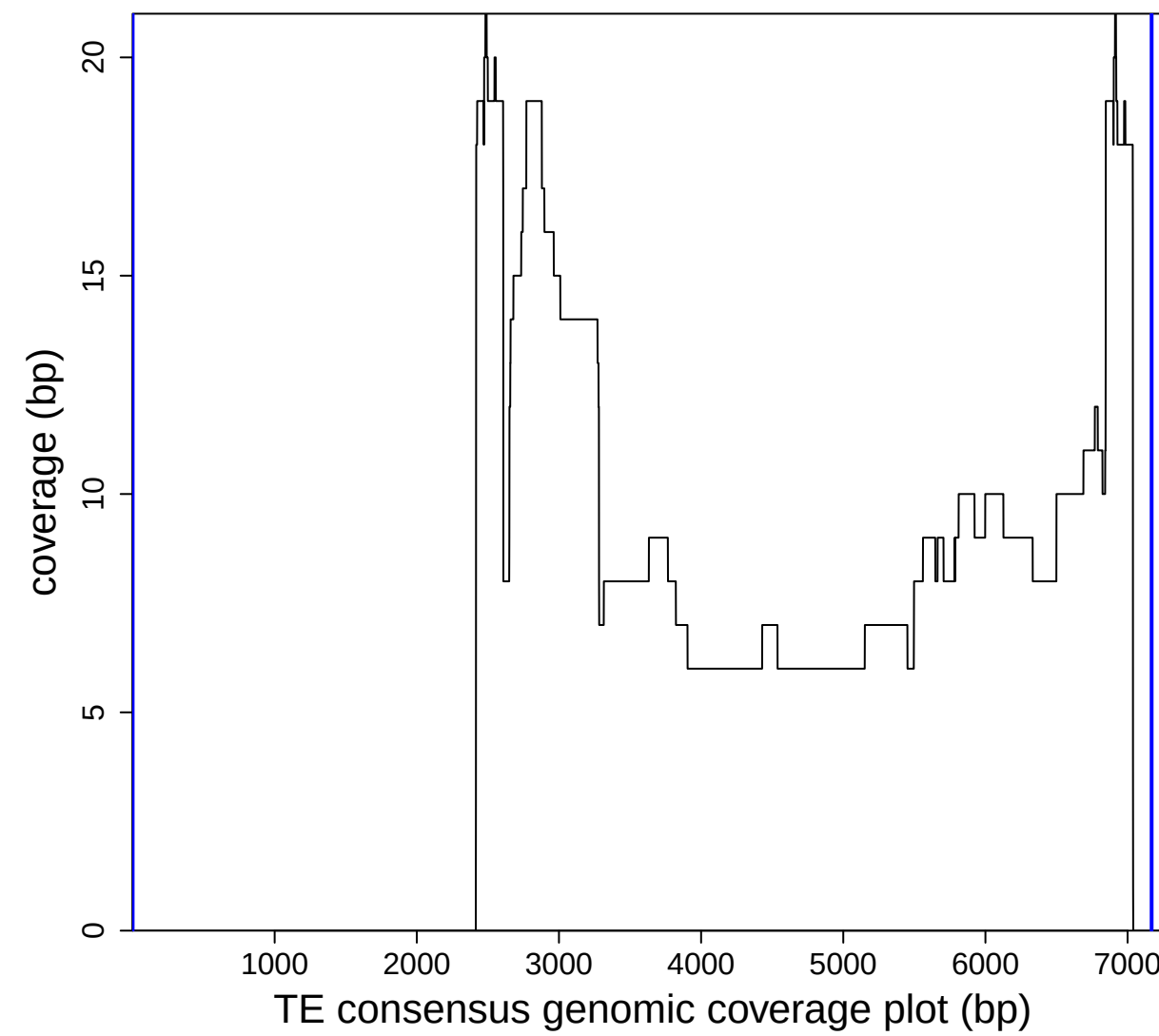
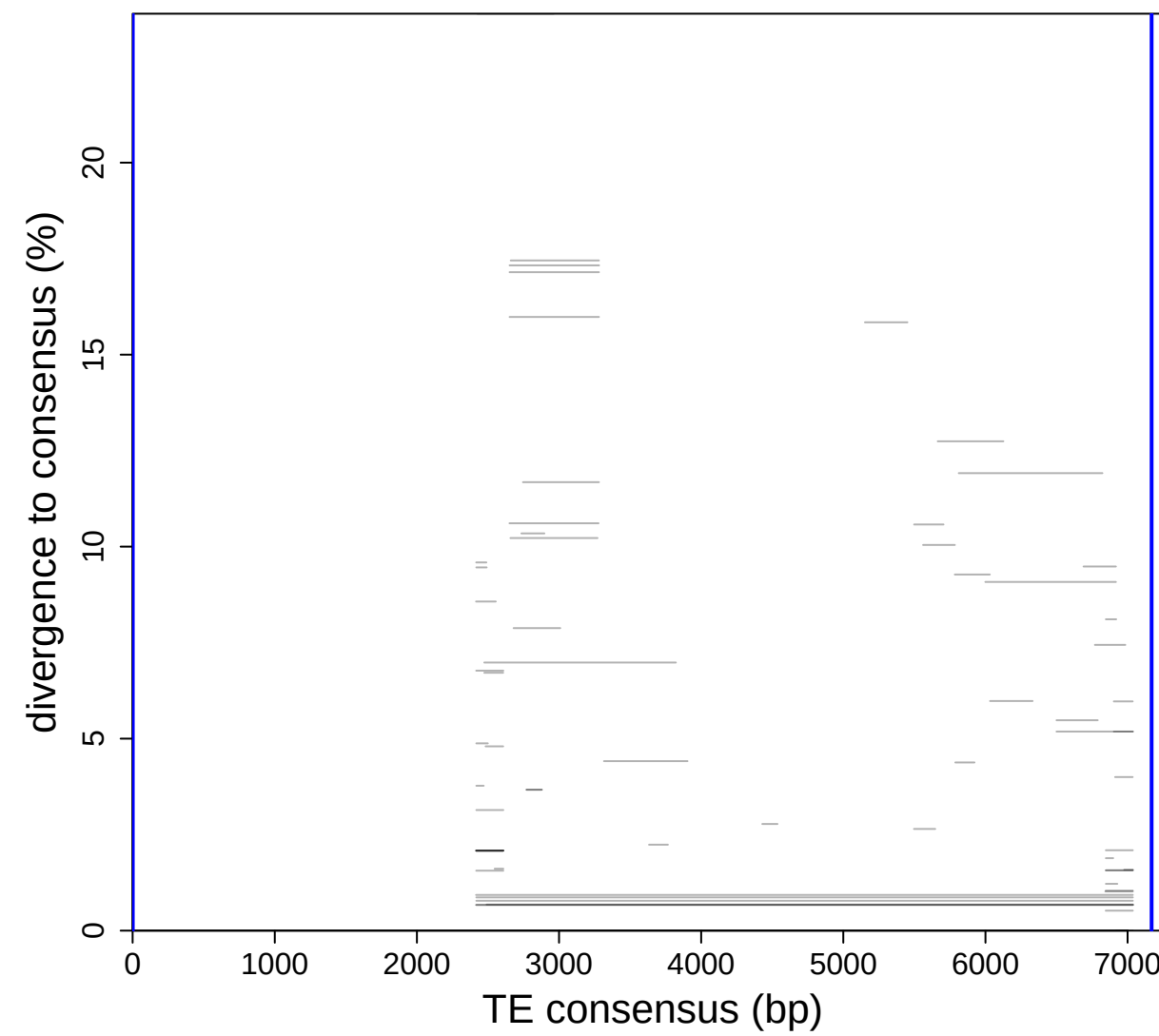
size: 6167bp; fragments: 66; full length: 0 (>=5550.3bp)

After TETrimmer 6167 bp

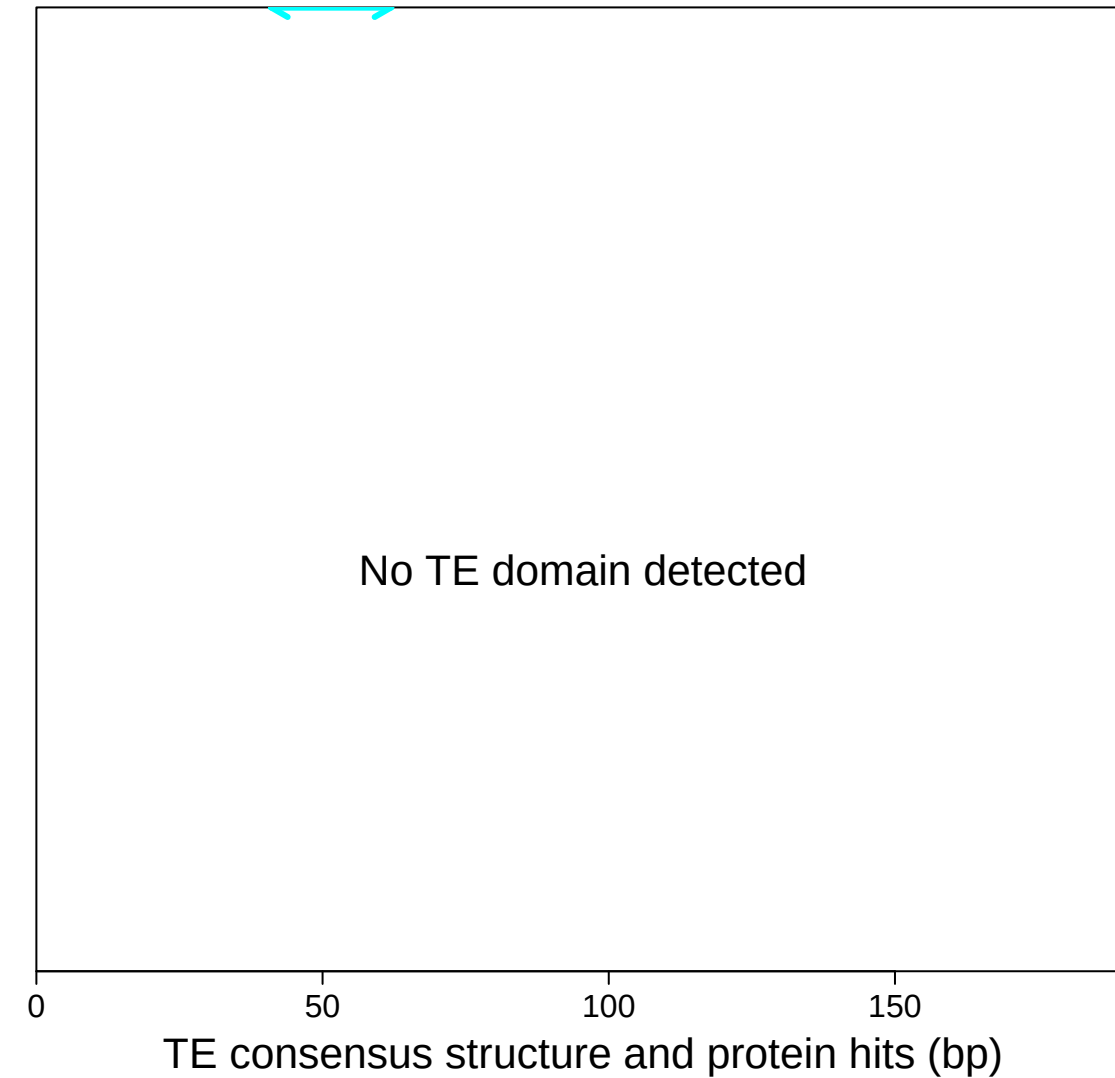
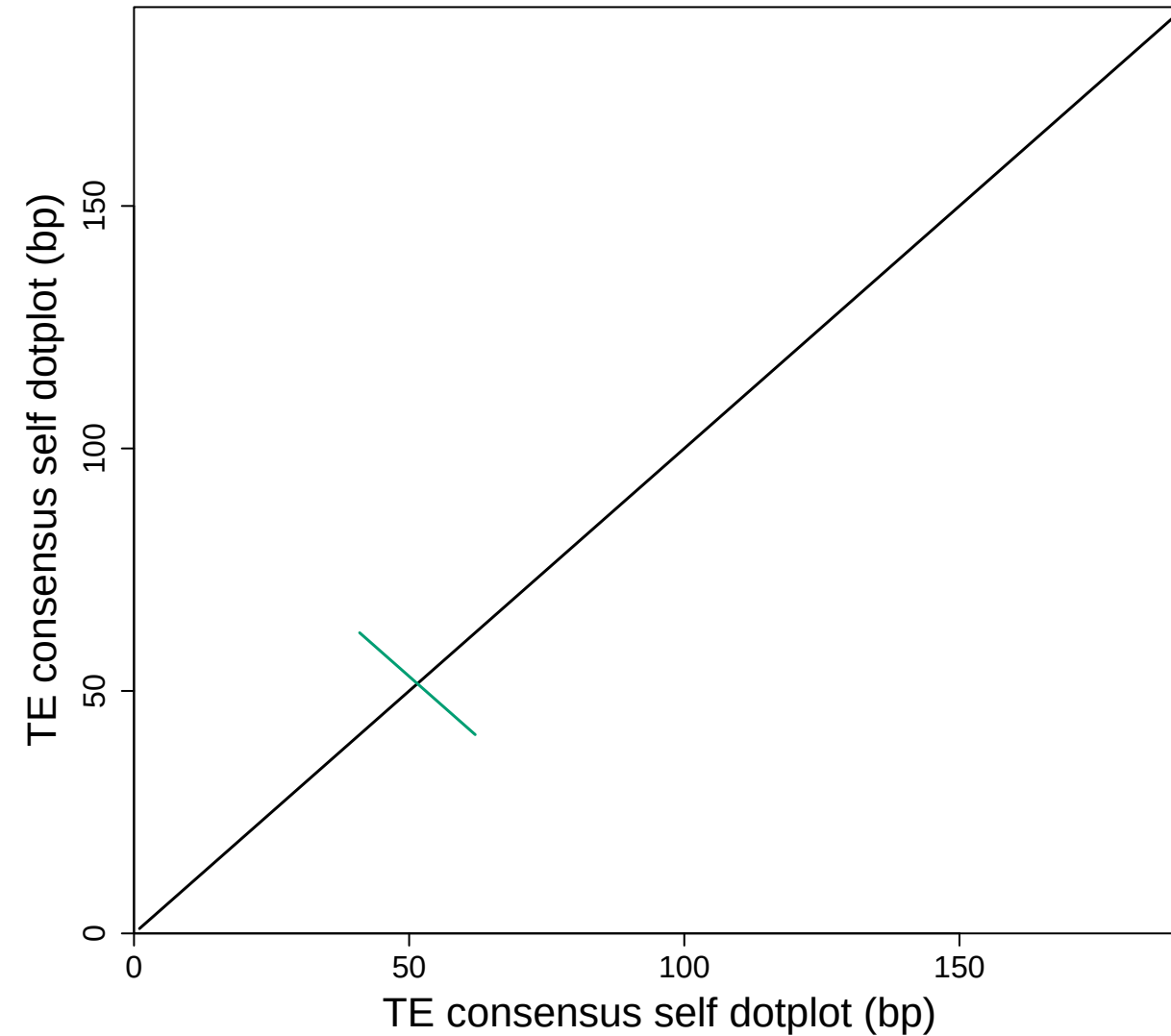
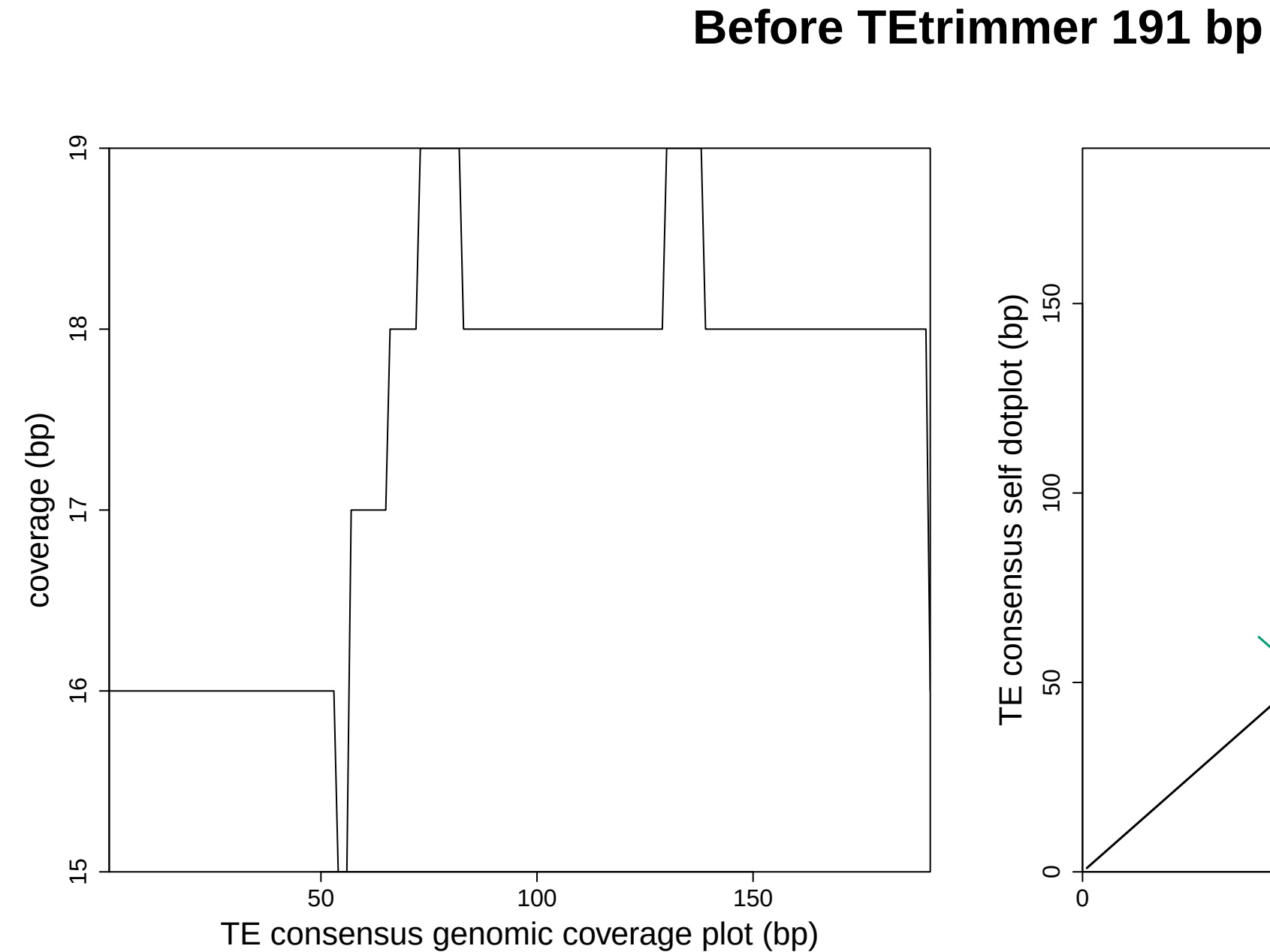
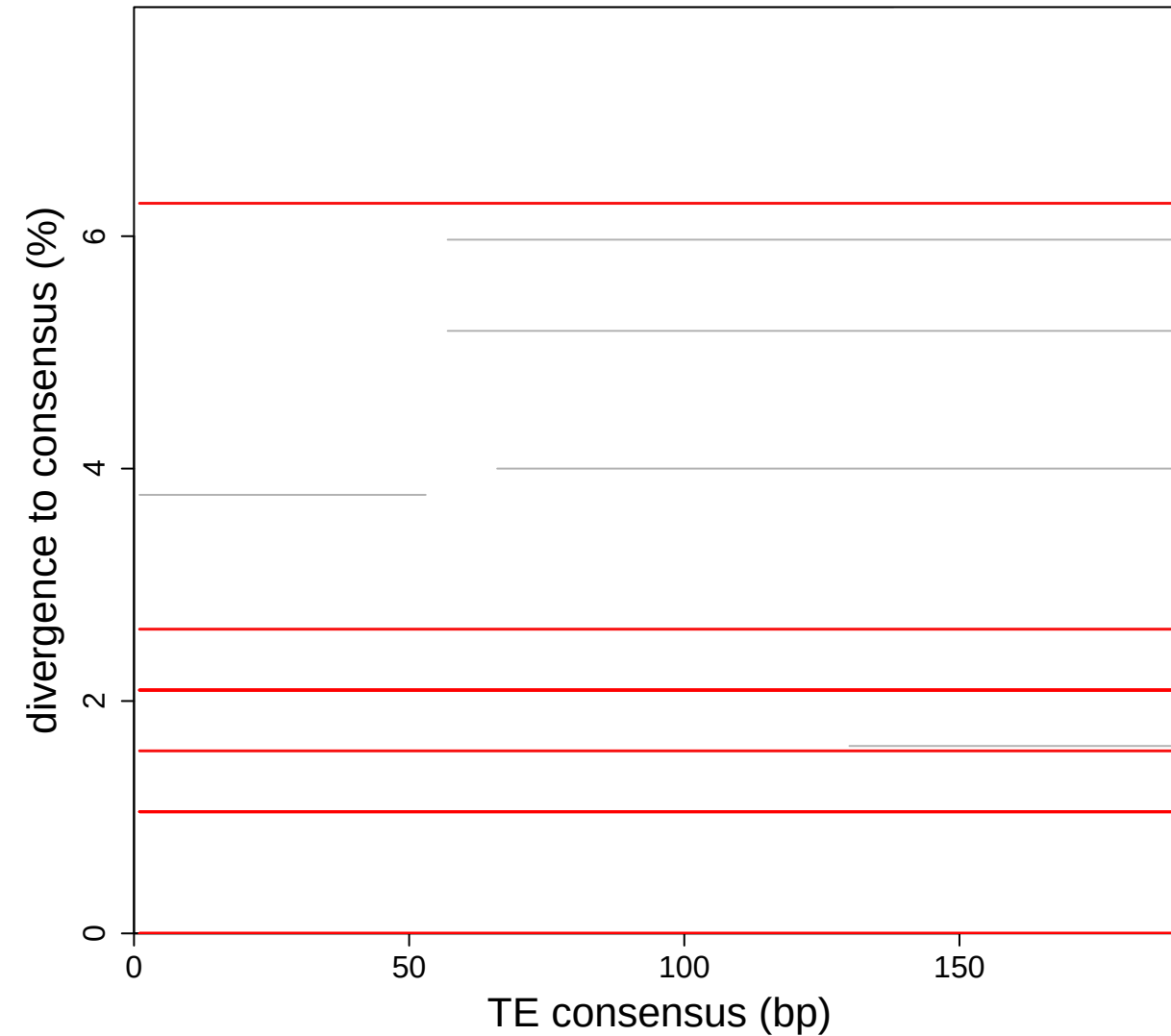


fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_
size: 7319bp; fragments: 66; full length: 0 (>=6587.1bp)

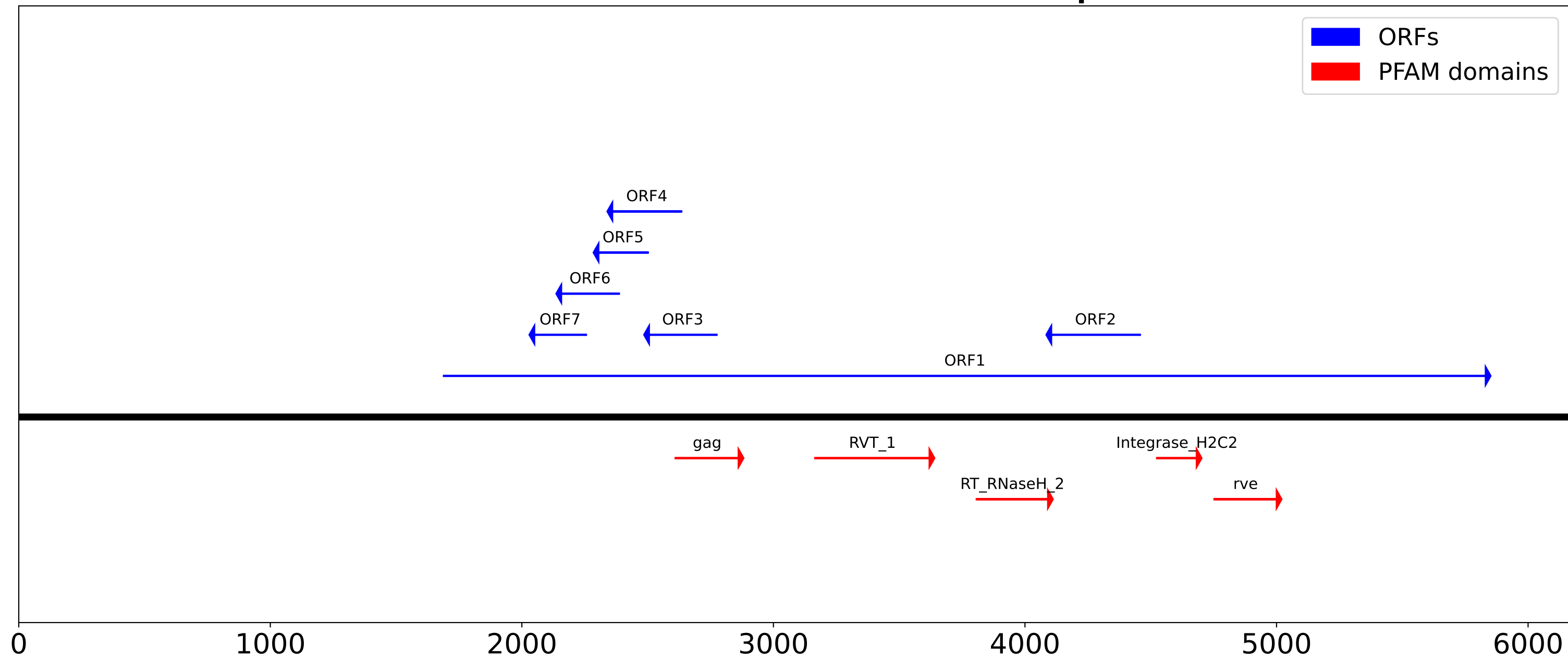
After TEtrimmer Extended plot Blue lines are boundaries



TE: ltr_1_family_44
size: 191bp; fragments: 21; full length: 13 (≥ 171.9 bp)



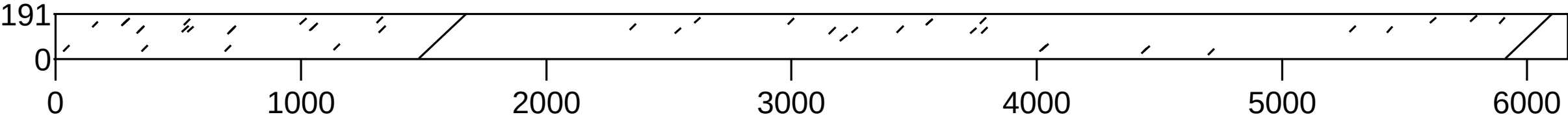
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)