

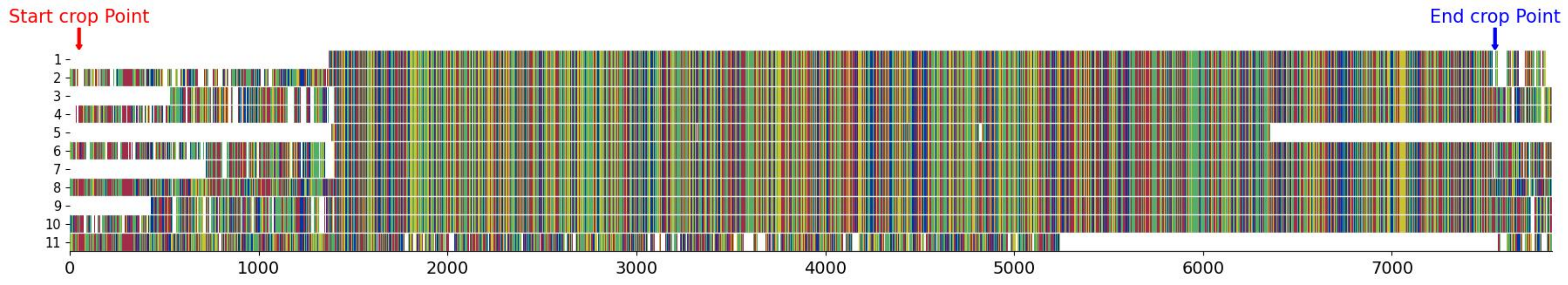
Start crop Point



MSA length = 7498

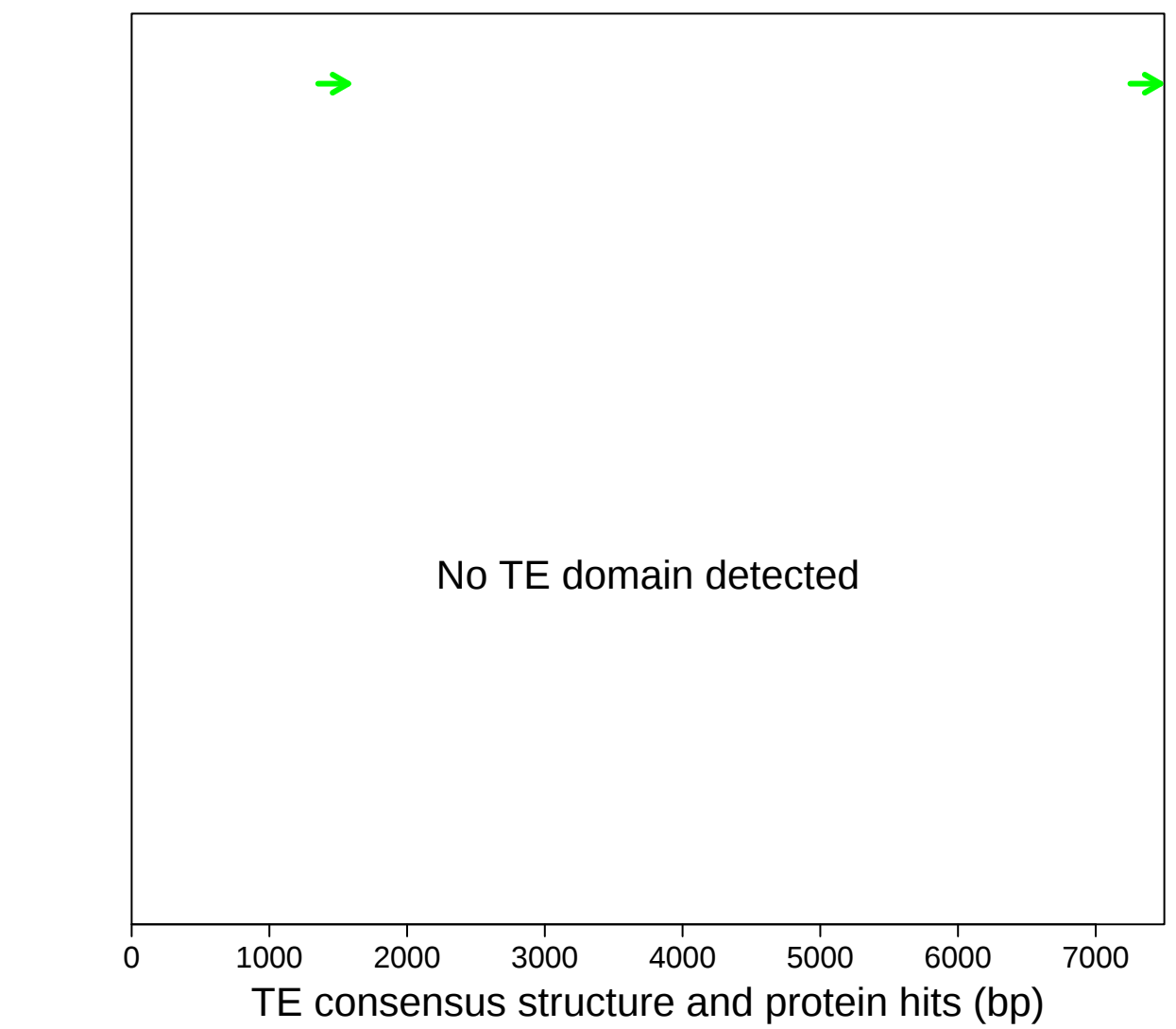
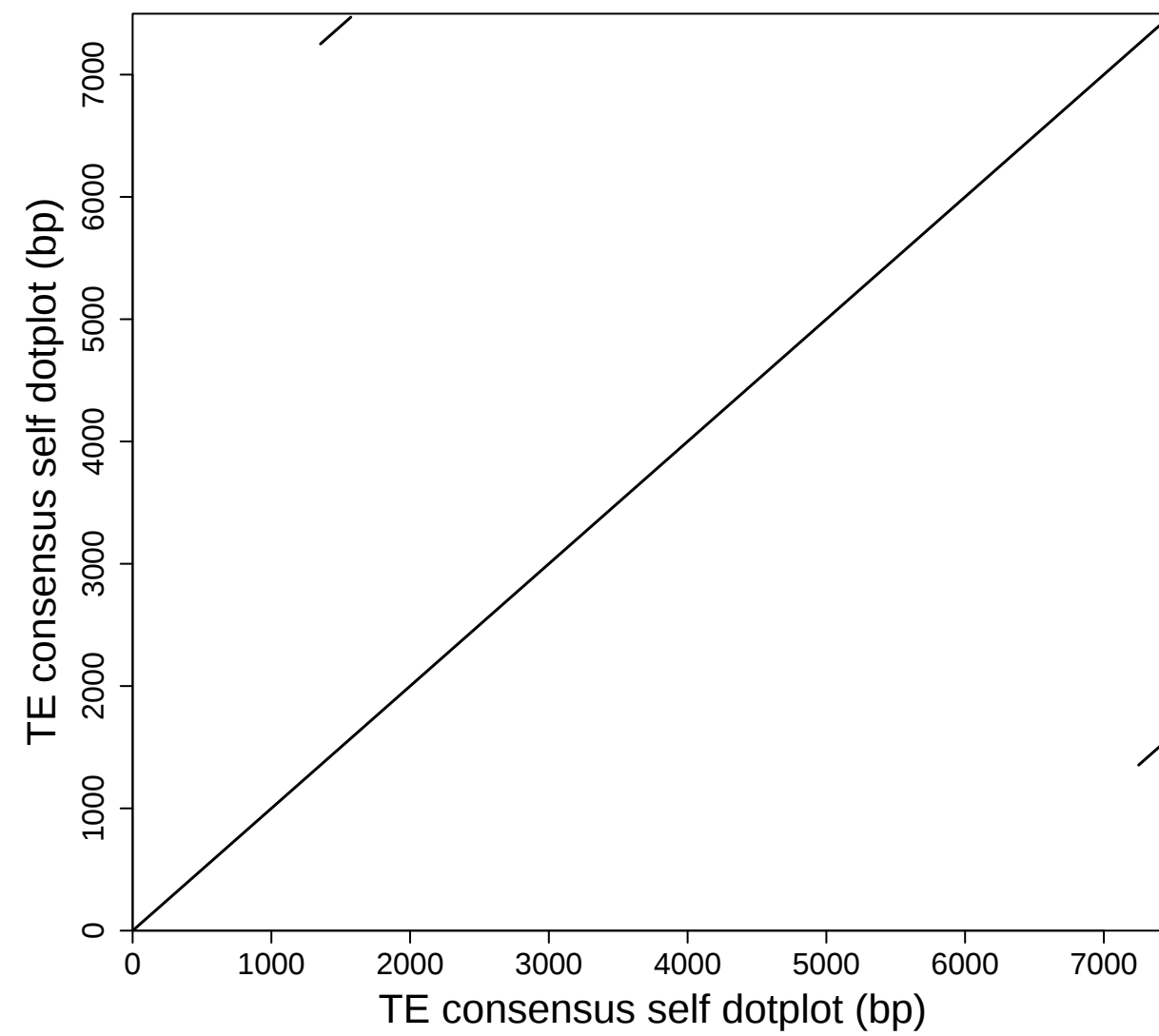
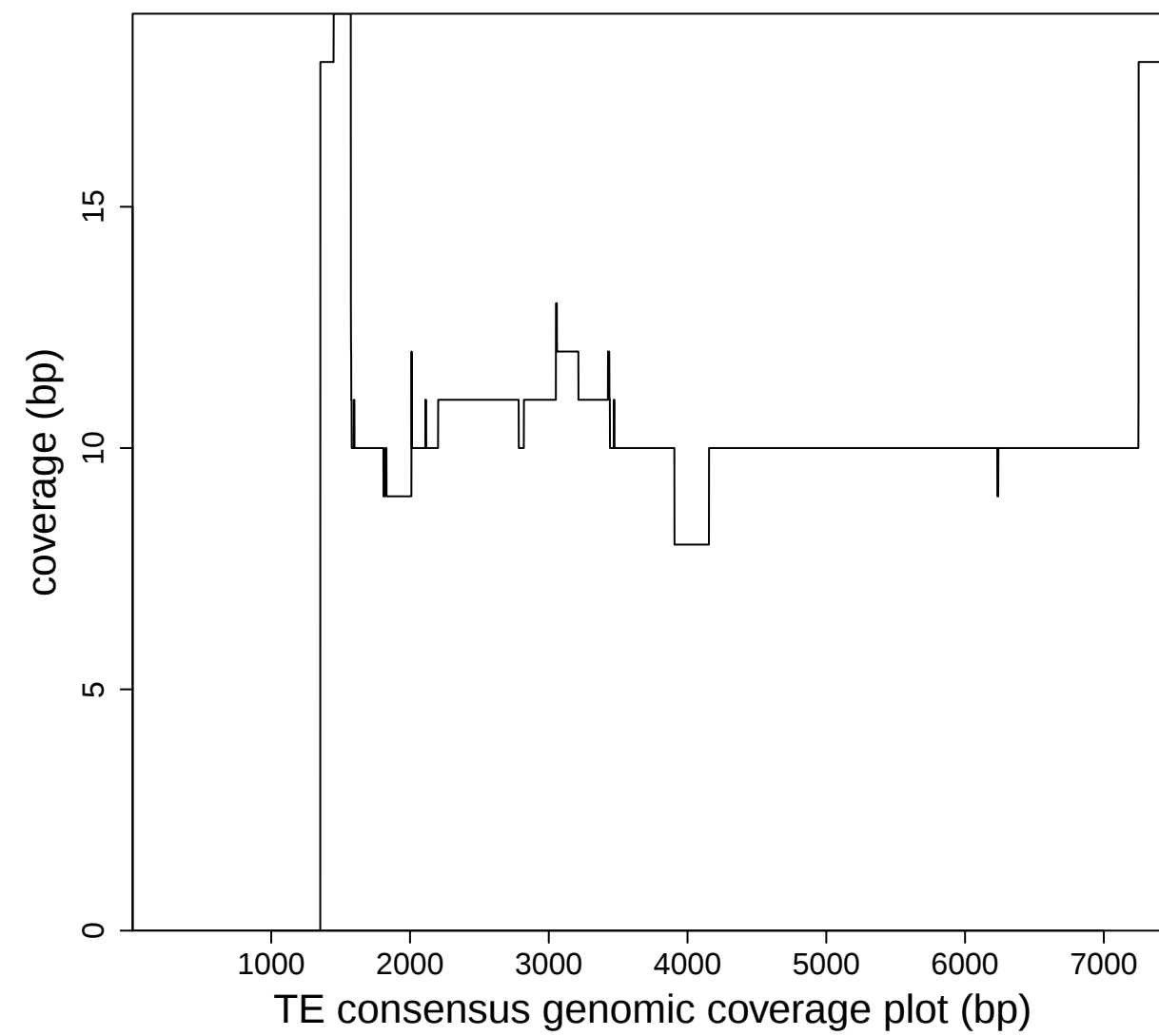
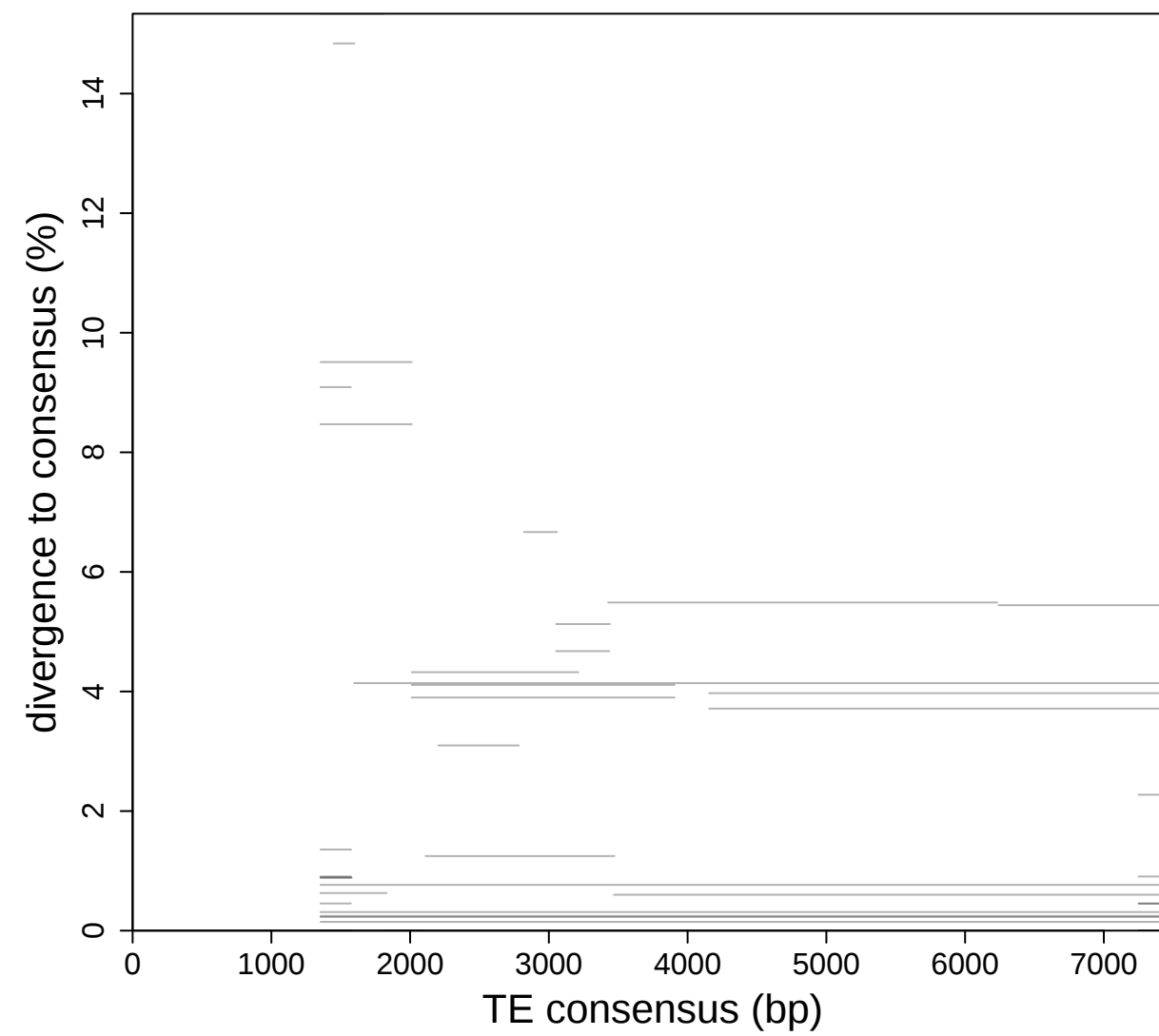
End crop Point

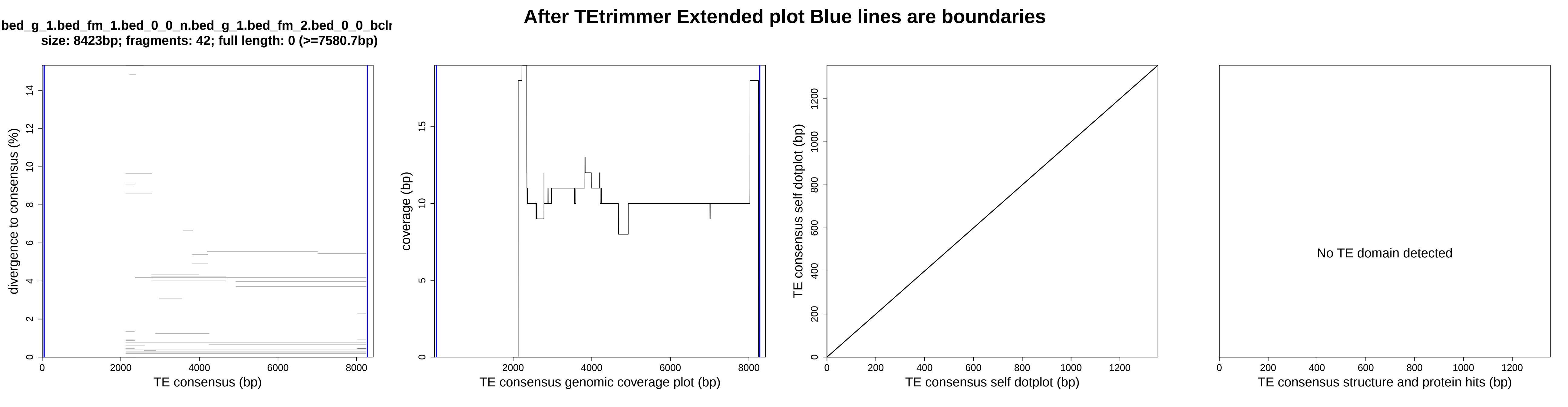




_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_al
size: 7498bp; fragments: 42; full length: 0 (>=6748.2bp)

After TEtrimmer 7498 bp

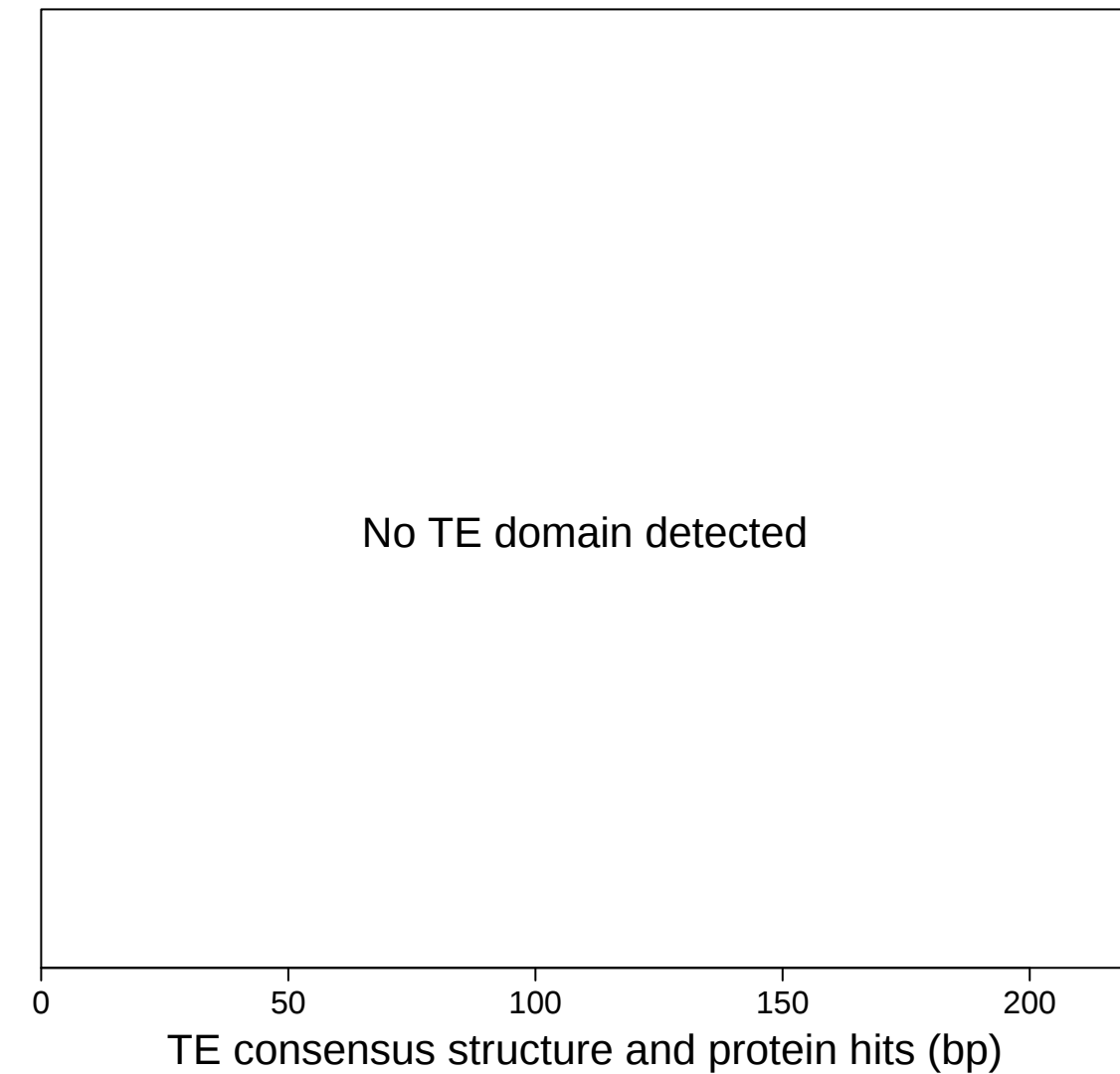
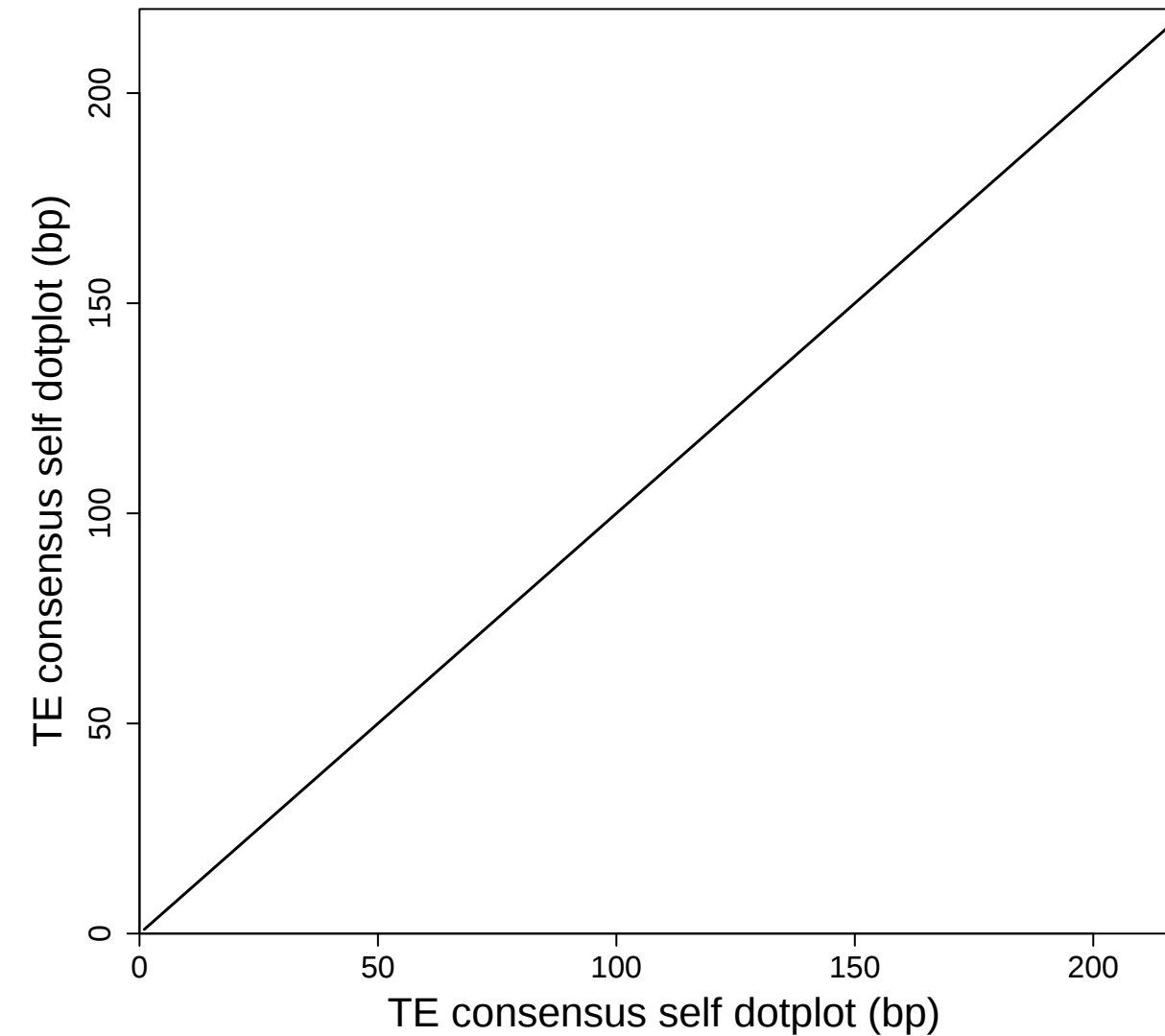
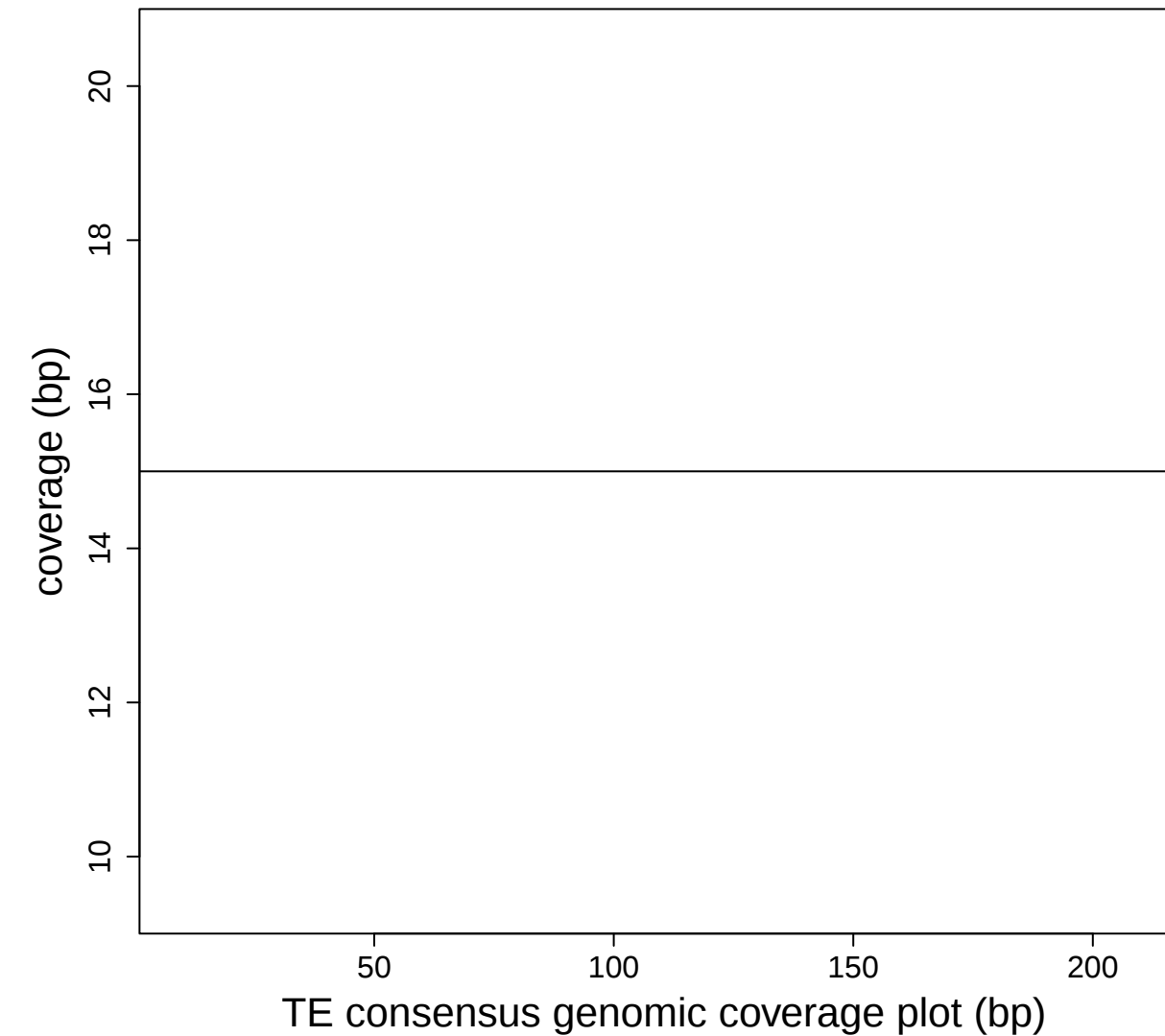
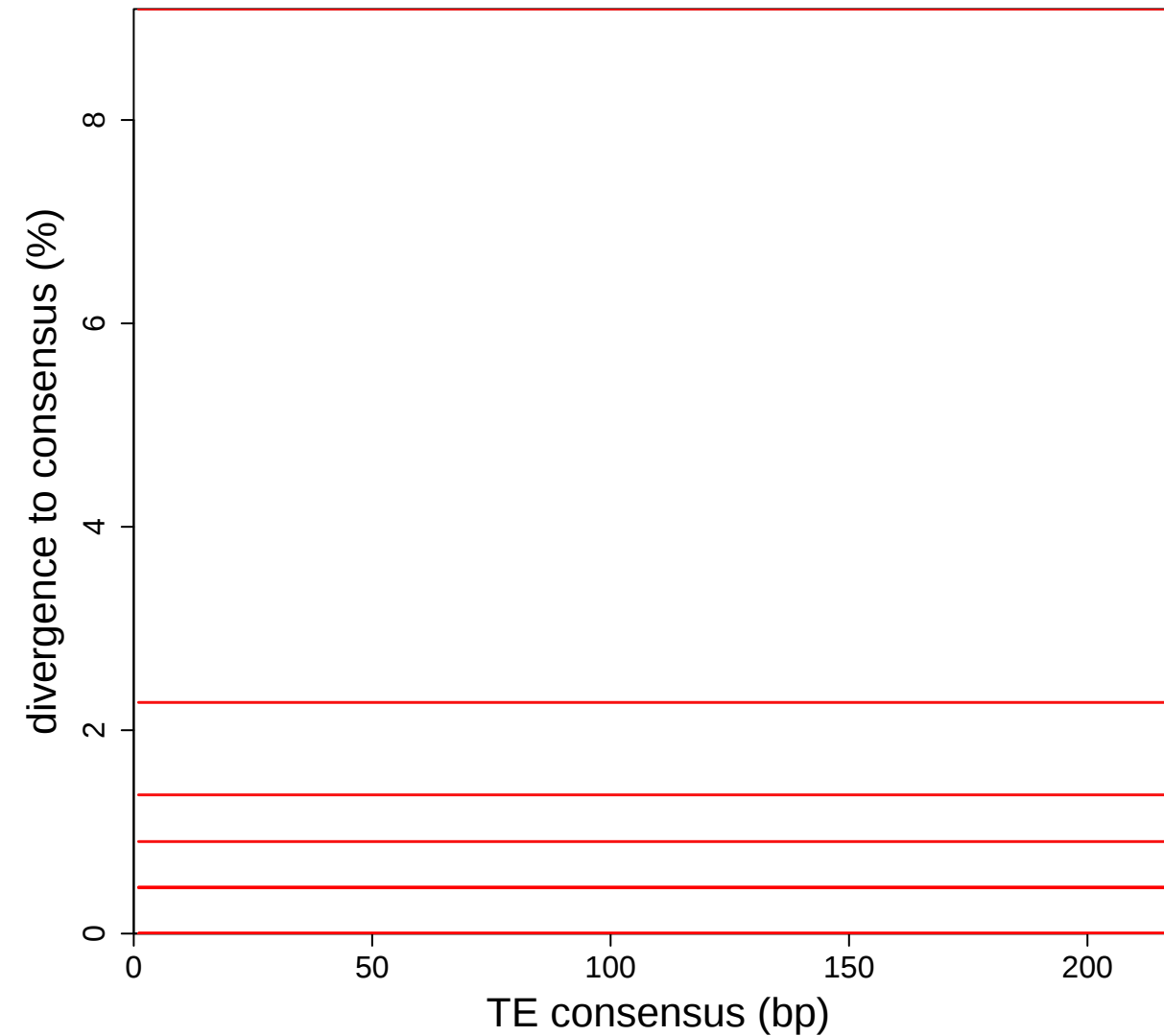




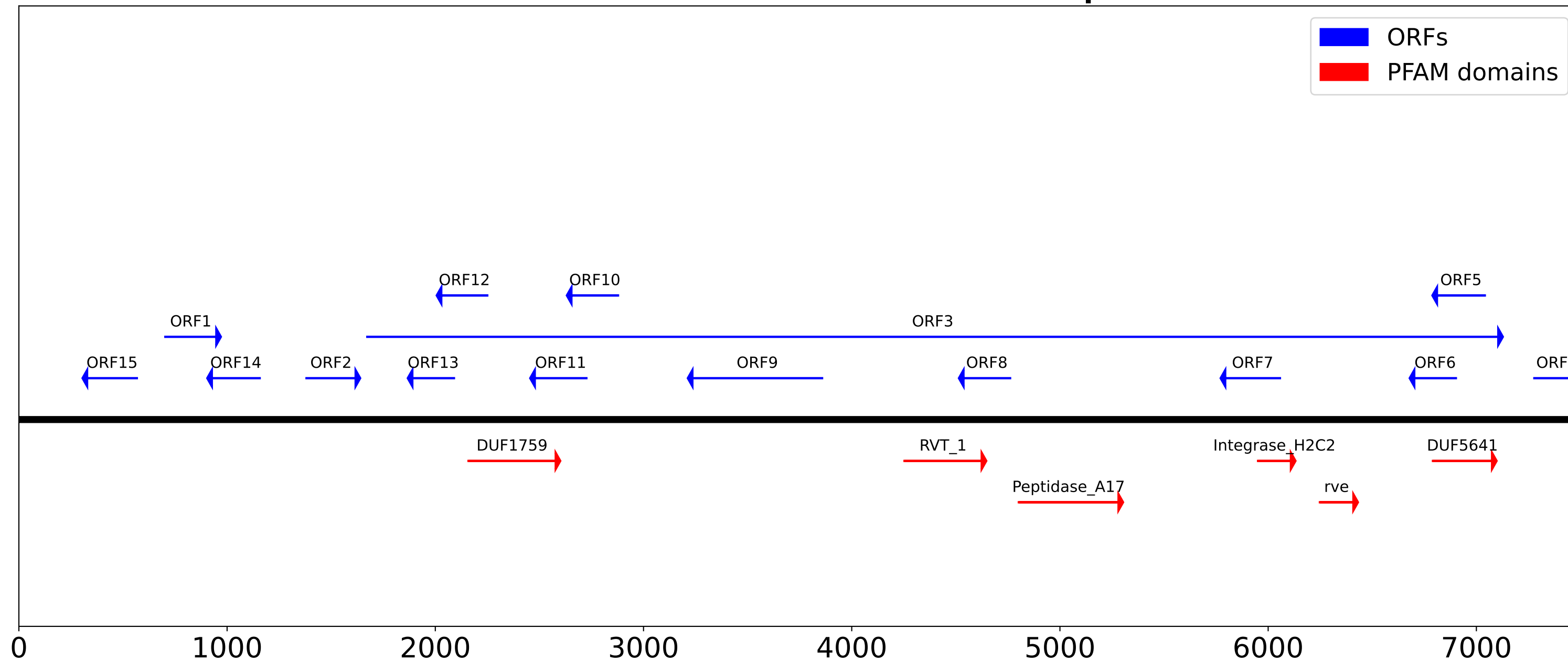
Before TEtrimmer 220 bp

TE: ltr_1_family_380

size: 220bp; fragments: 15; full length: 15 (≥ 198 bp)



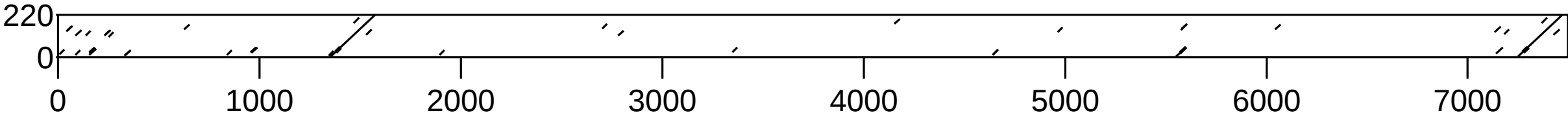
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)