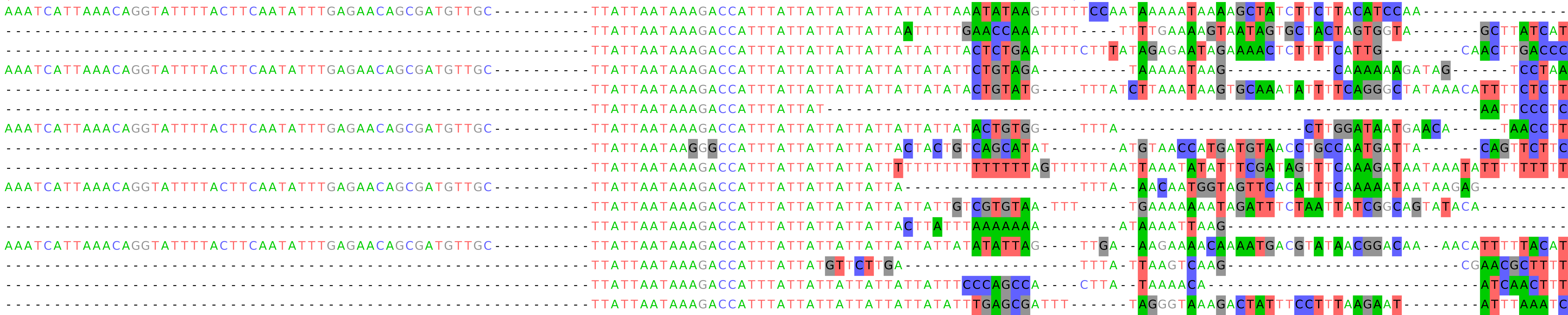


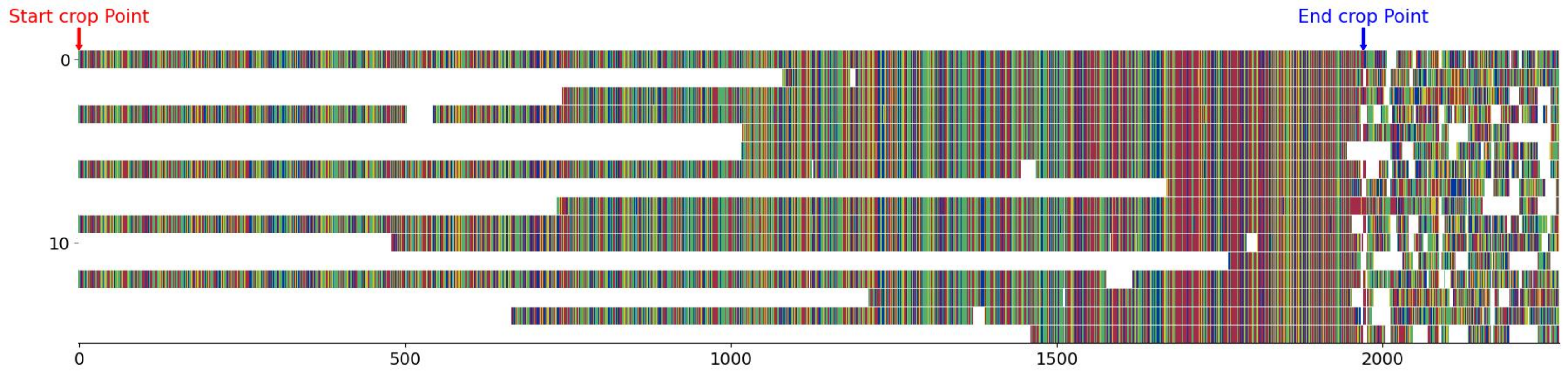
Start crop Point



MSA length = 1971

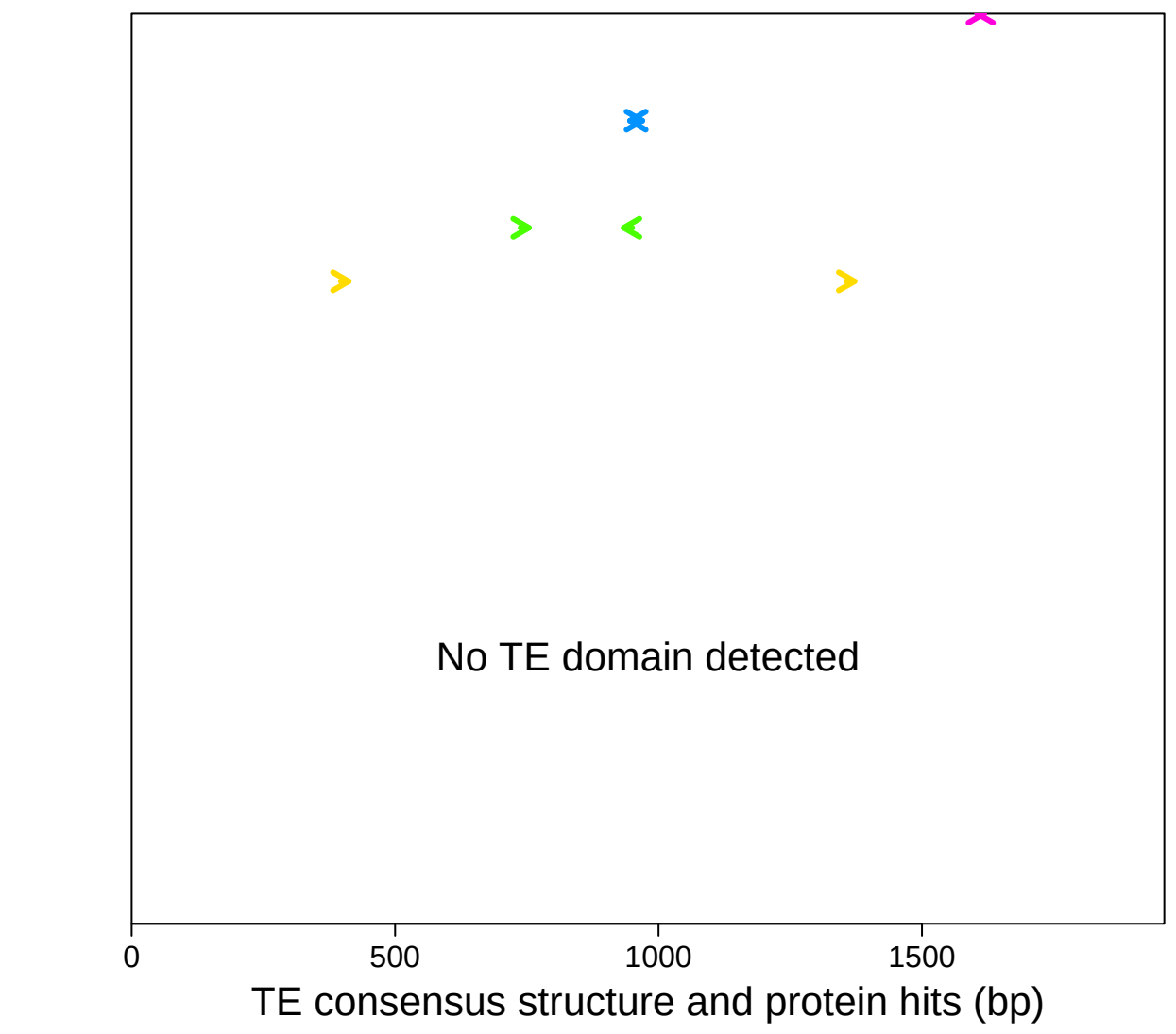
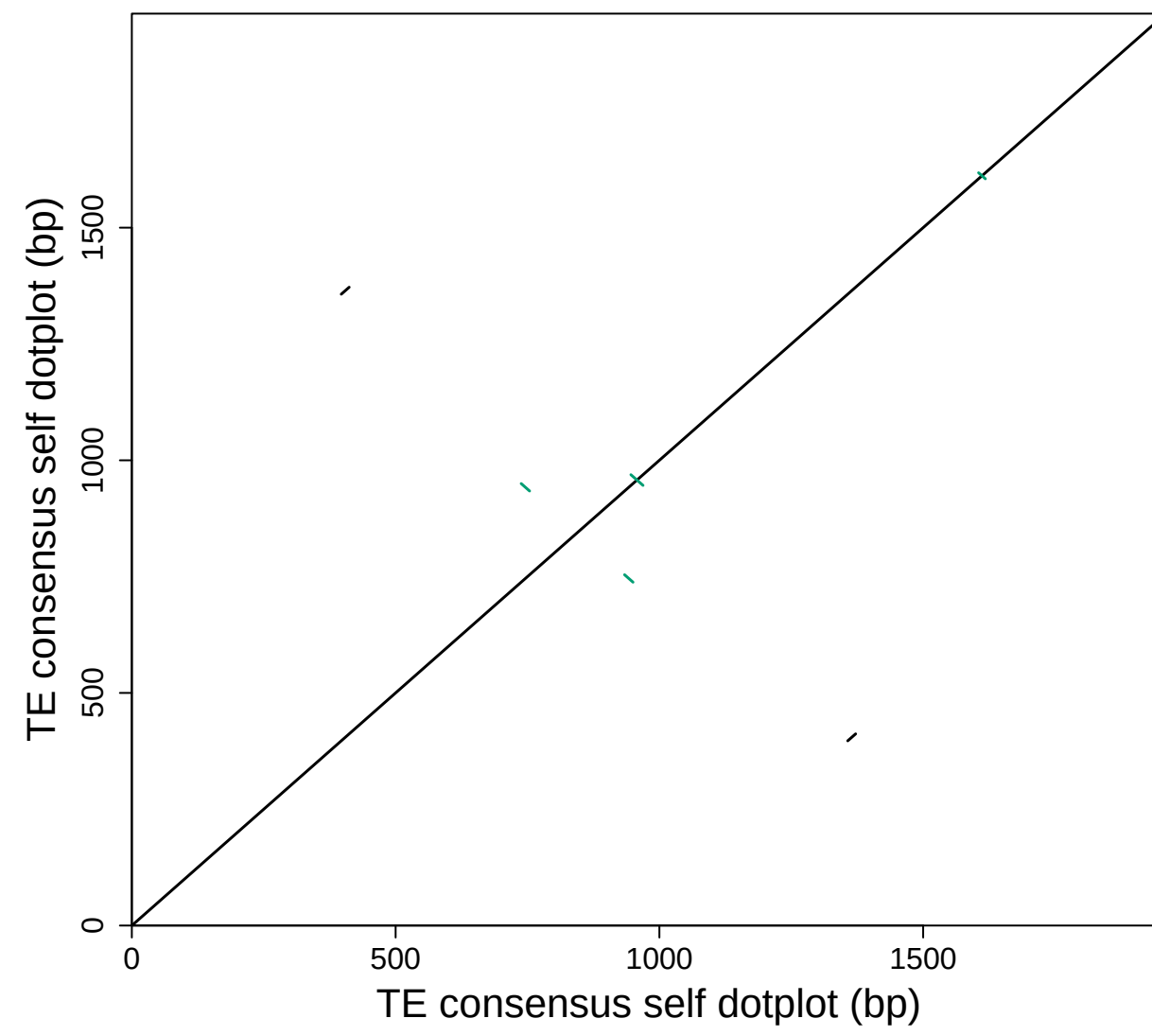
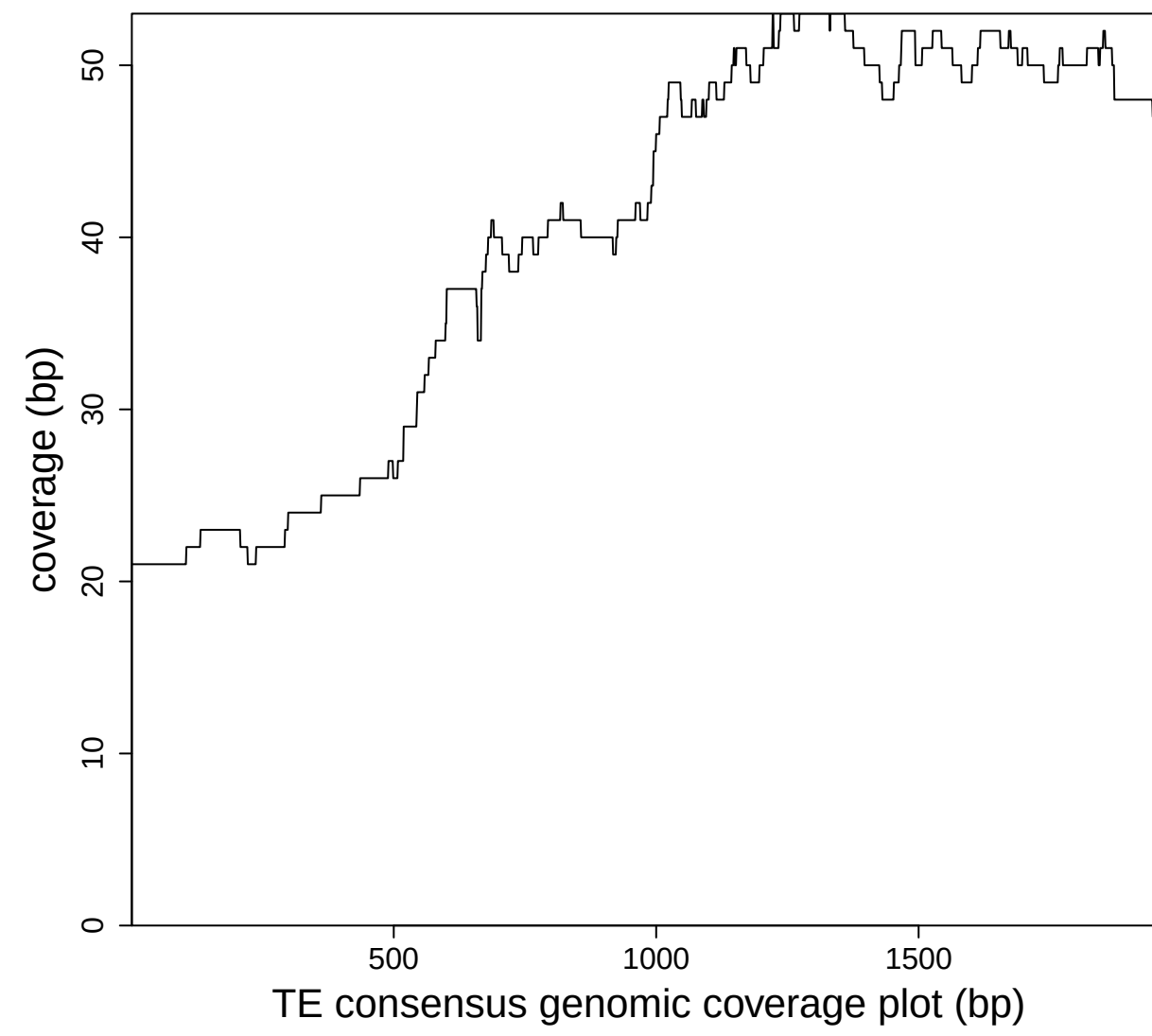
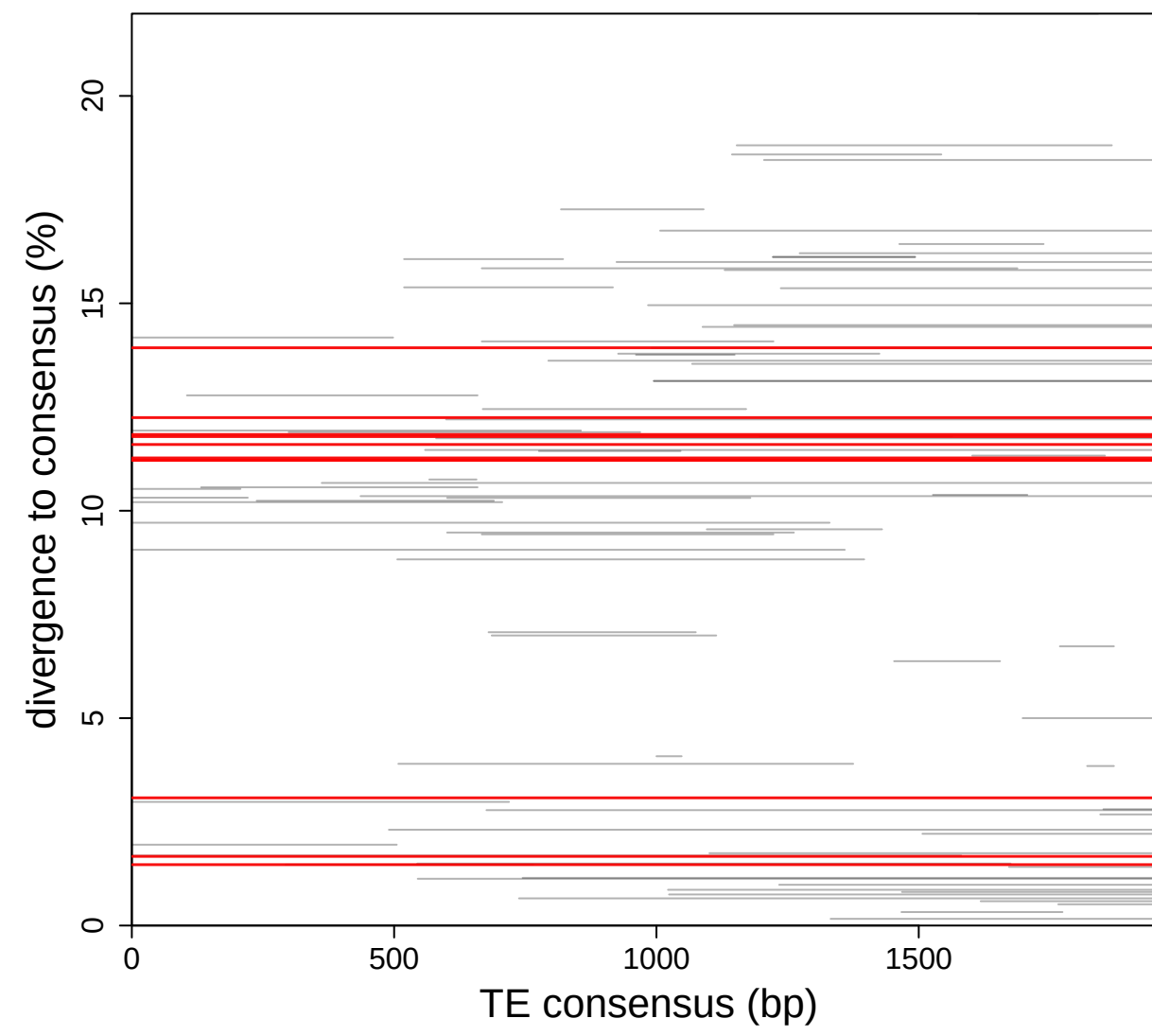
End crop Point





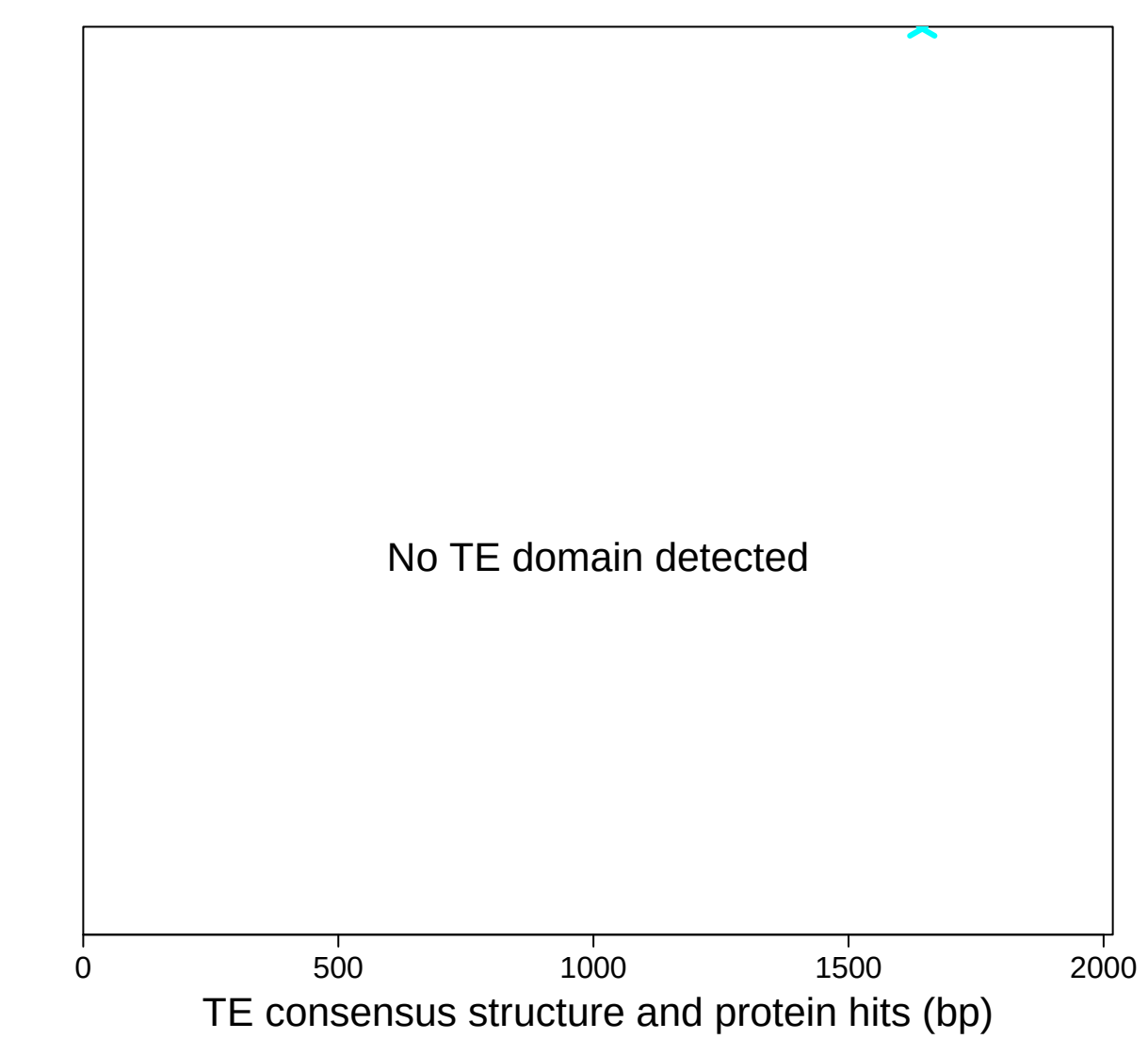
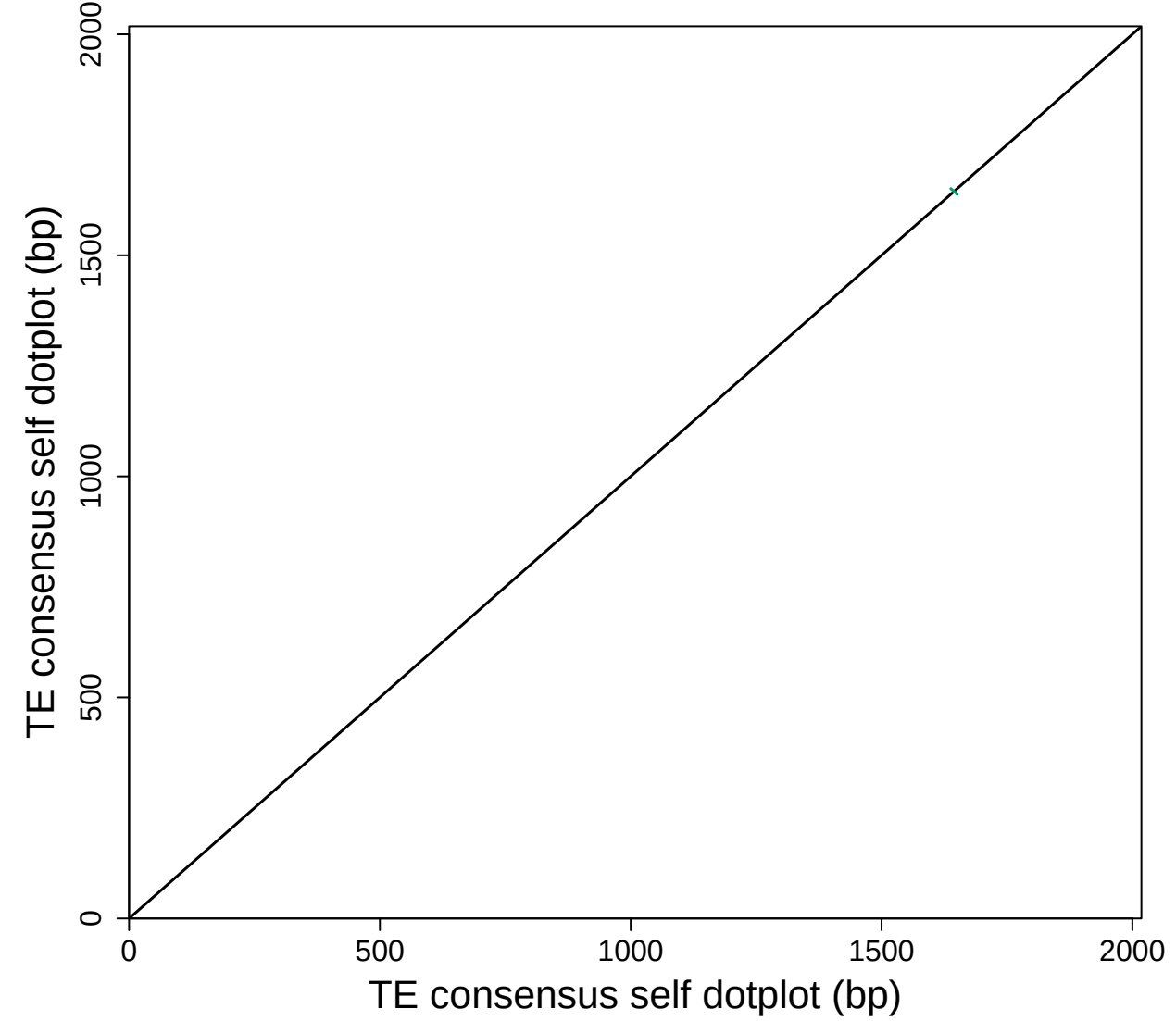
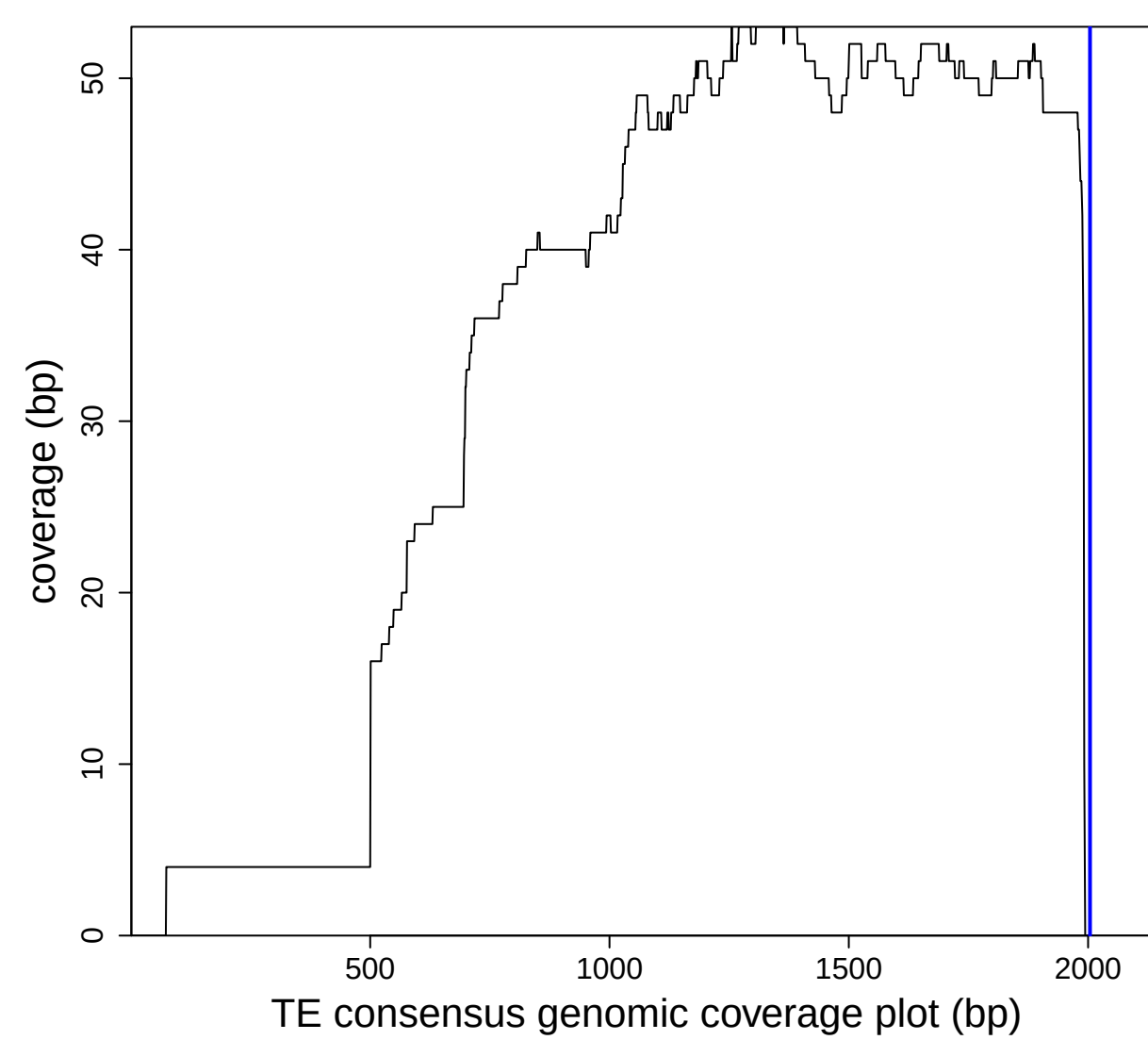
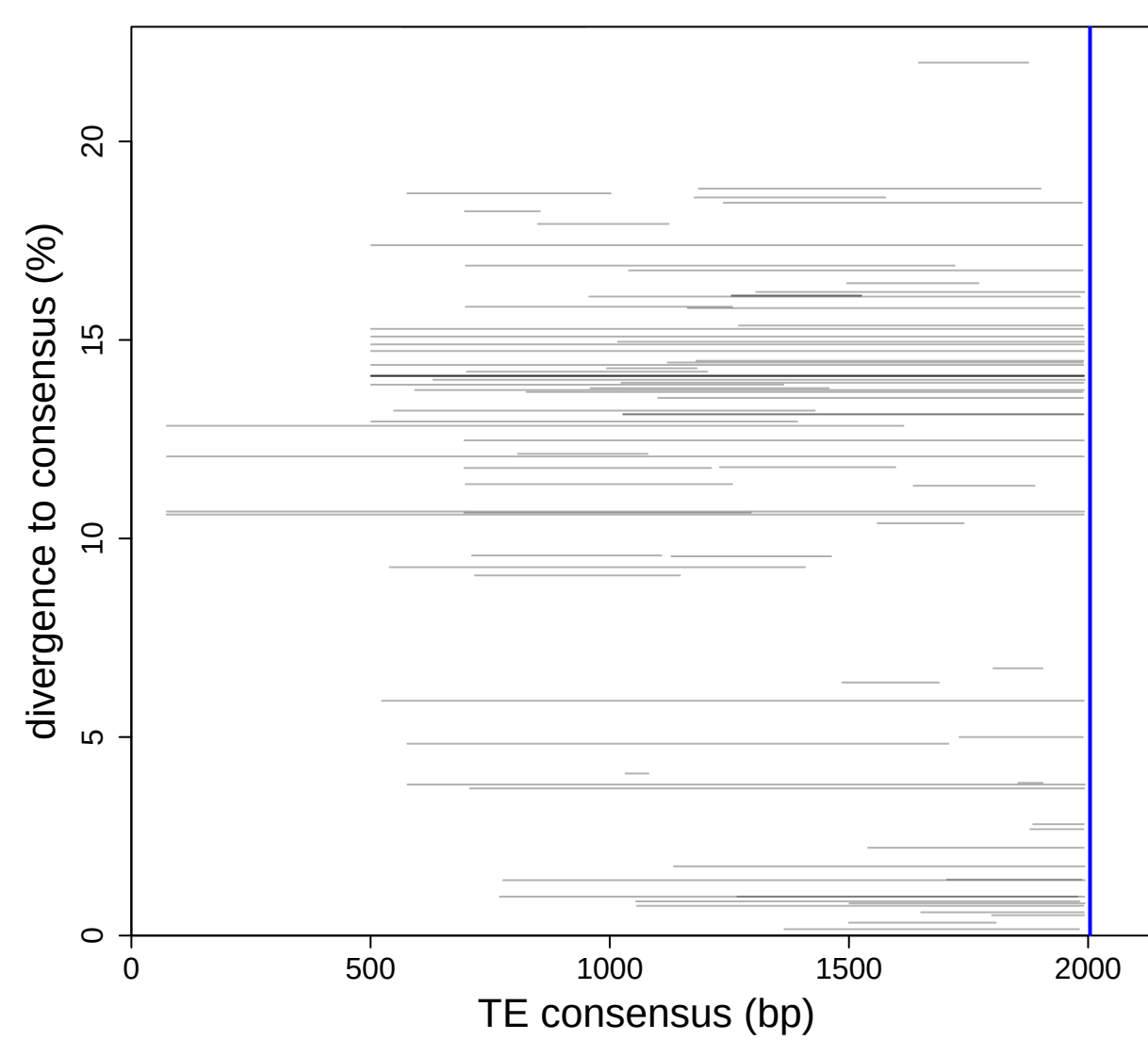
1.bed_fm 1.bed 0 0_n.bed_g 1.bed_fm 2.bed 0 0_bcln.fa aln.f
size: 1971bp; fragments: 96; full length: 11 (>=1773.9bp)

After TEtrimmer 1971 bp

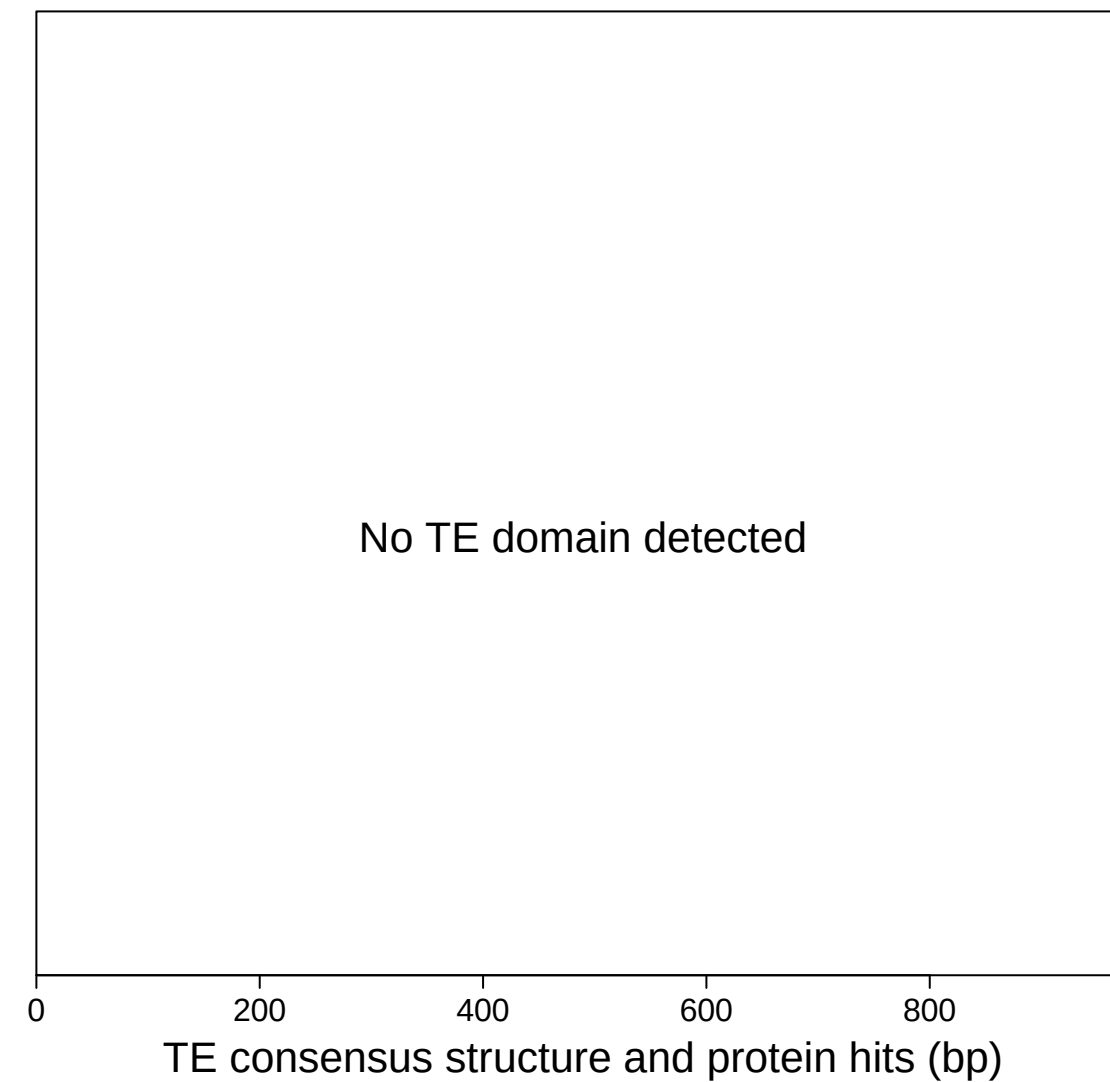
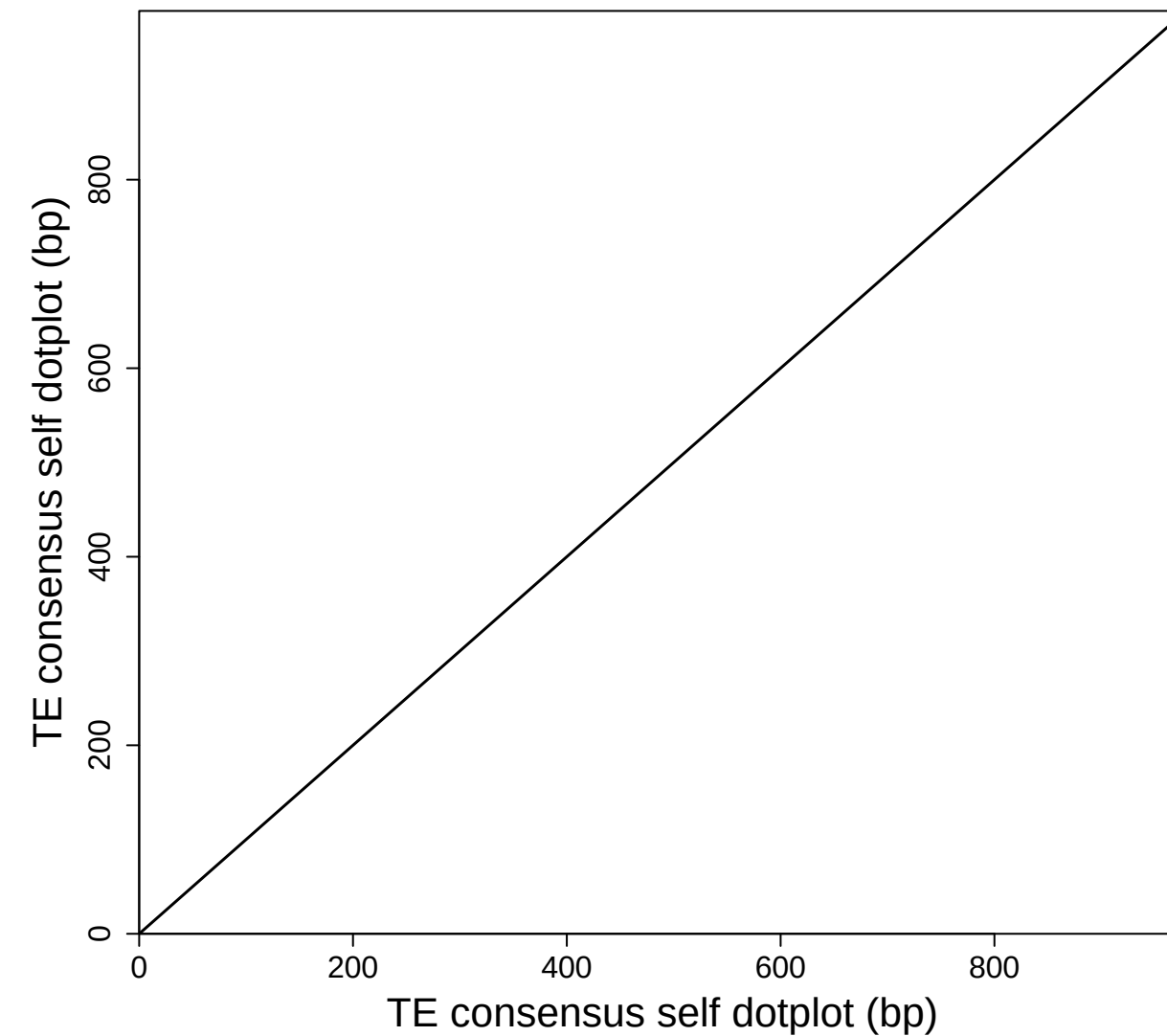
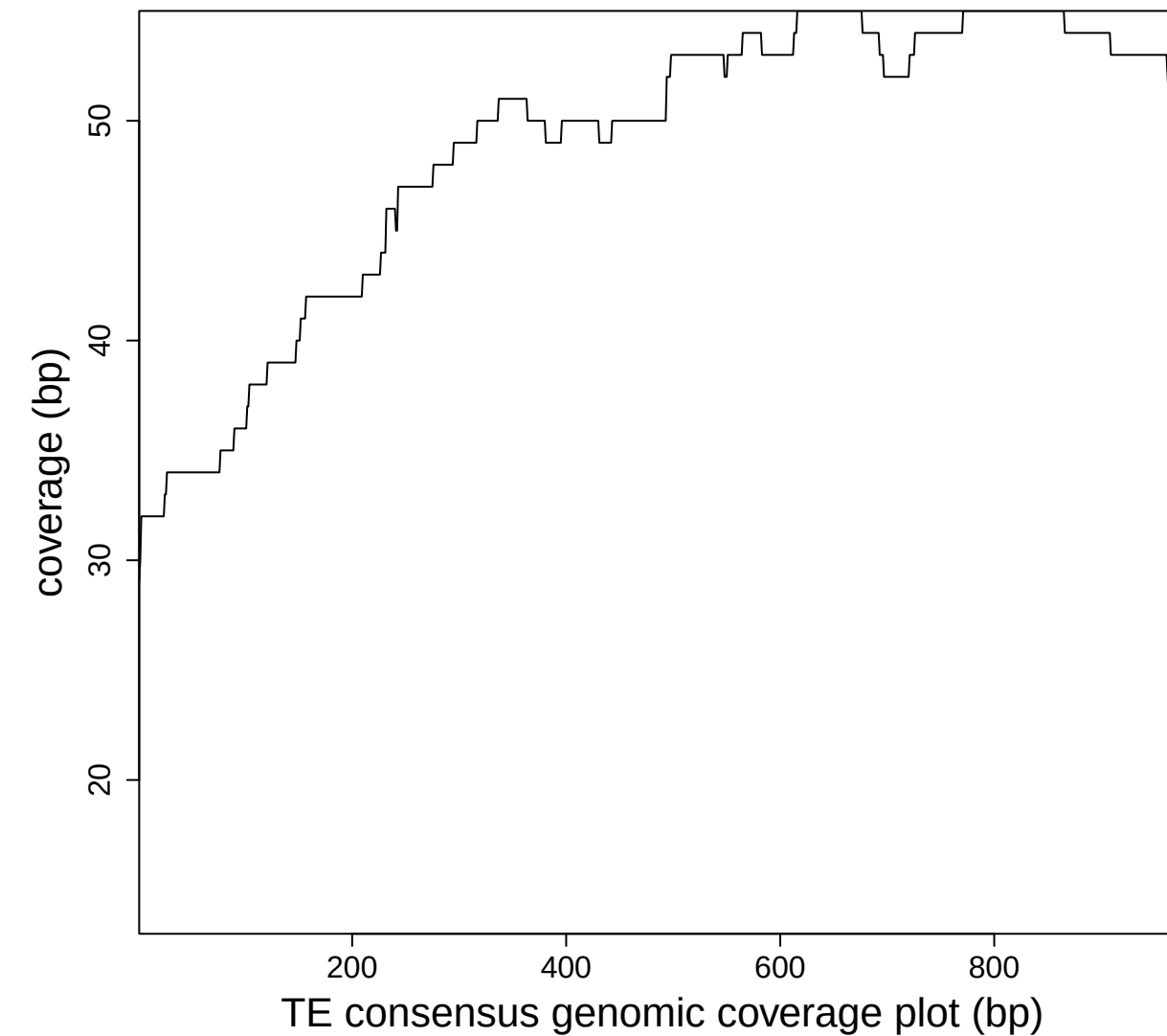
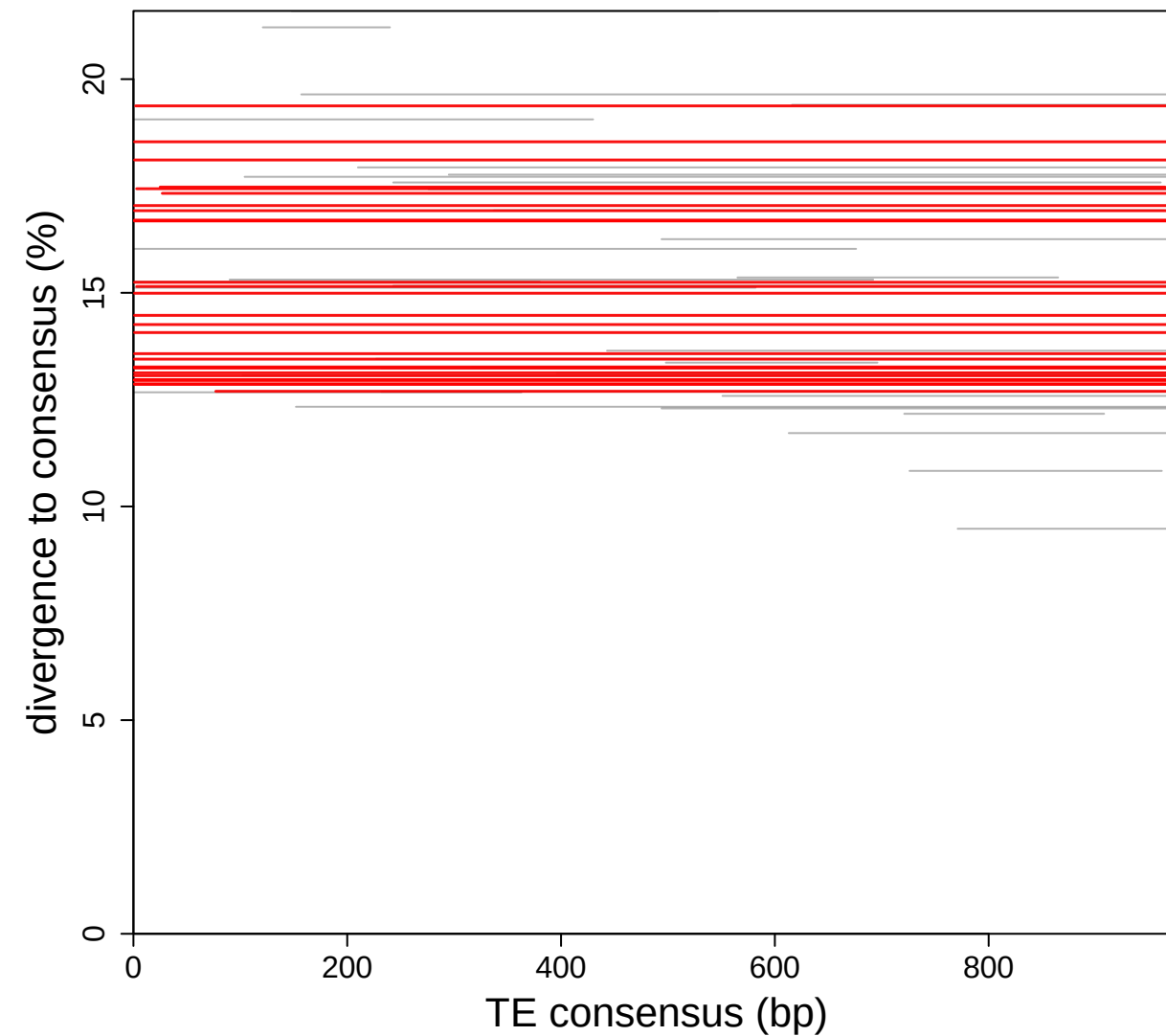


f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 2154bp; fragments: 84; full length: 0 (>=1938.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

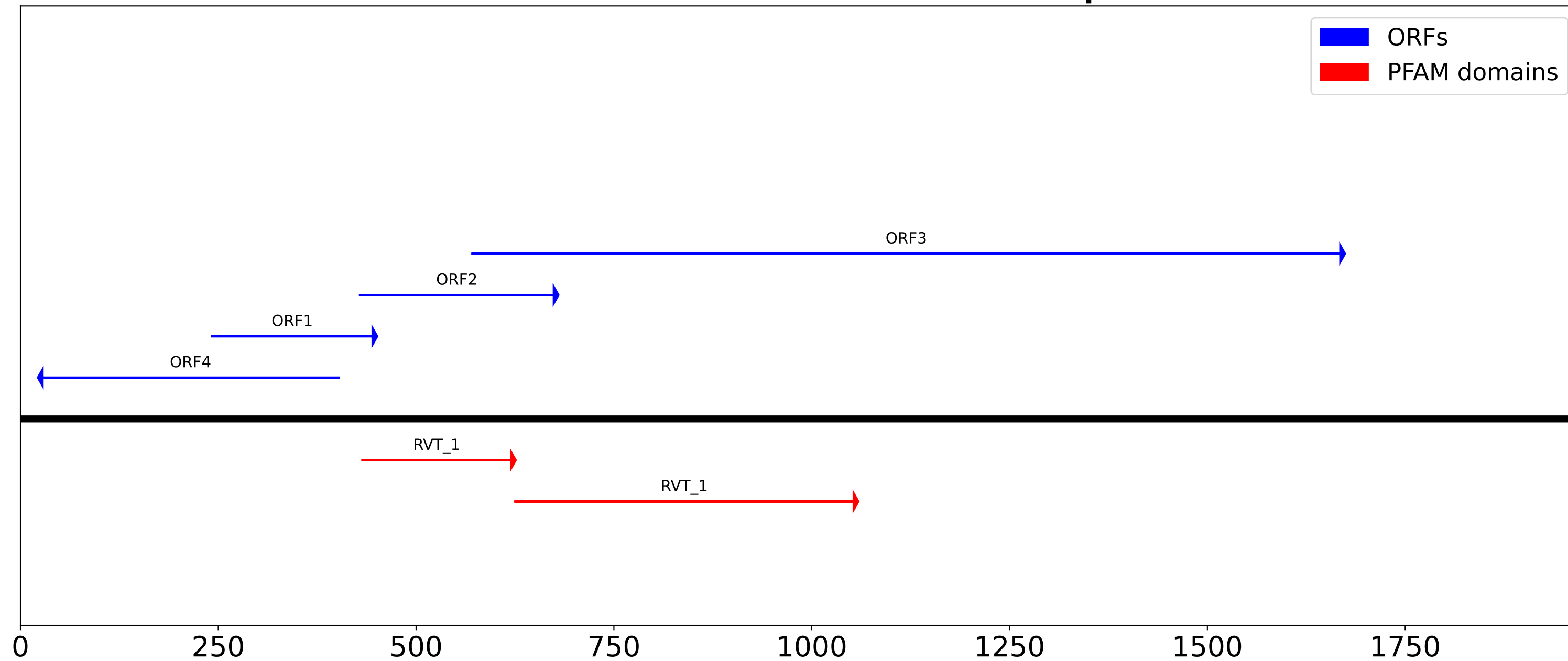


TE: rnd_4_family_2479
size: 979bp; fragments: 64; full length: 30 (≥ 881.1 bp)

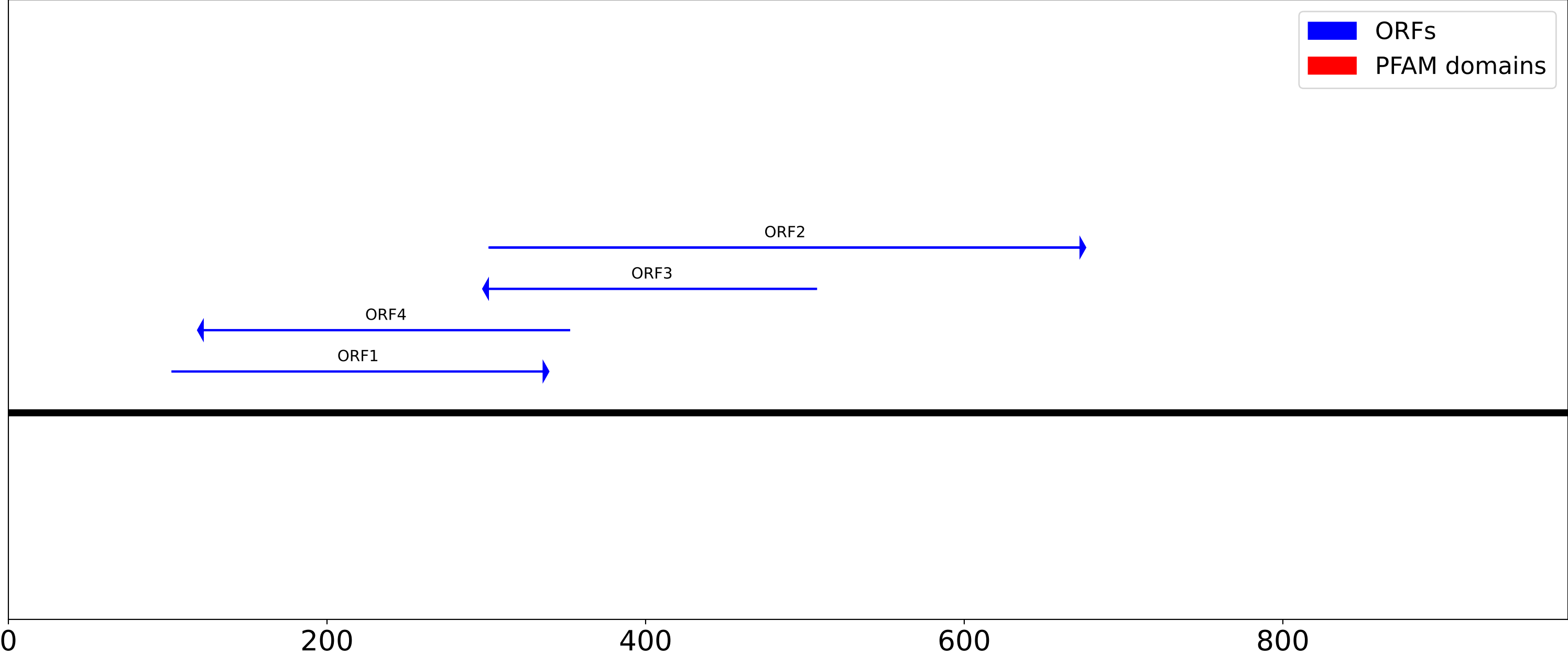


Before TEtrimmer 979 bp

After TEtrimmer ORF and PFAM domain plot



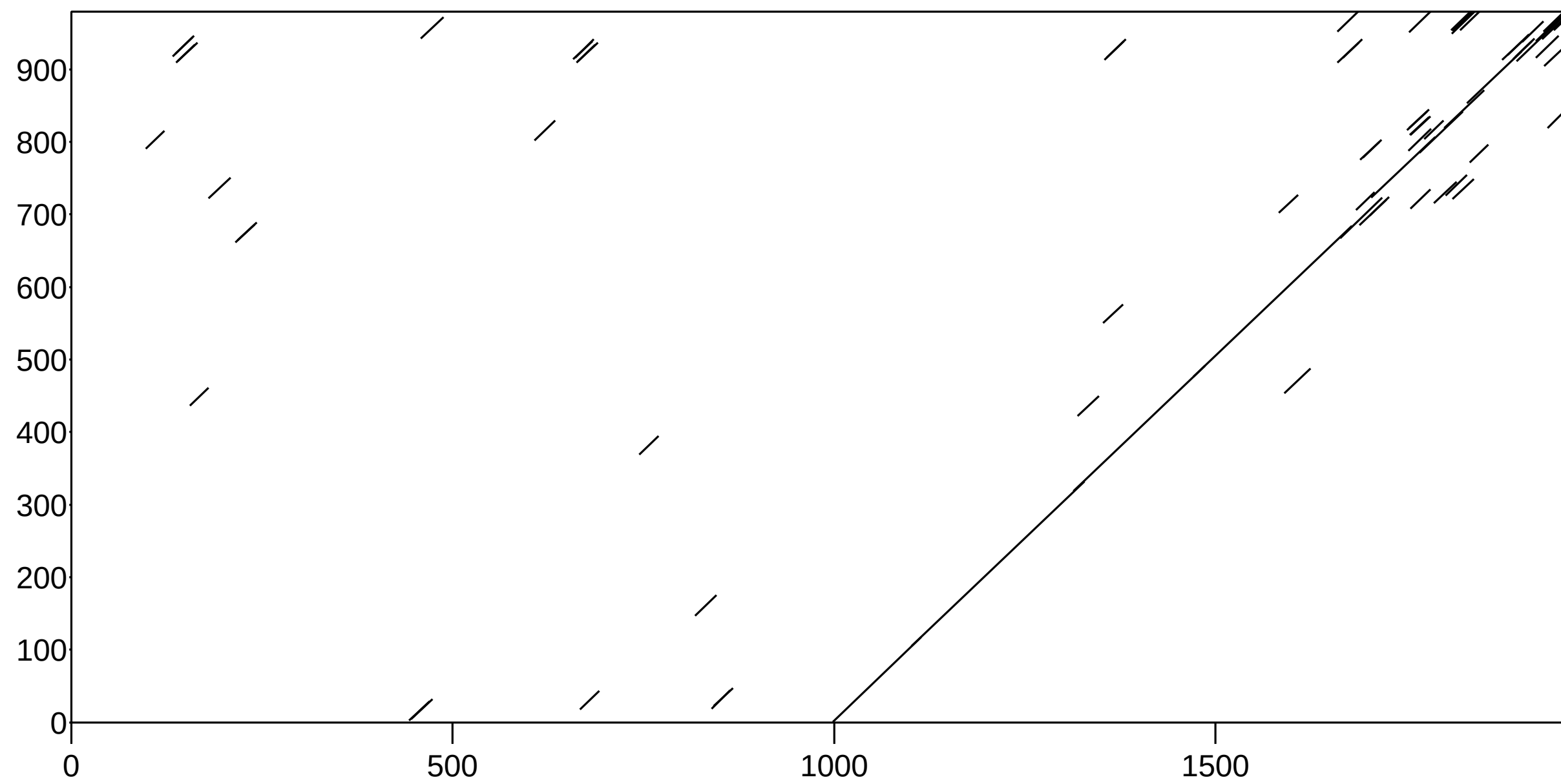
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)