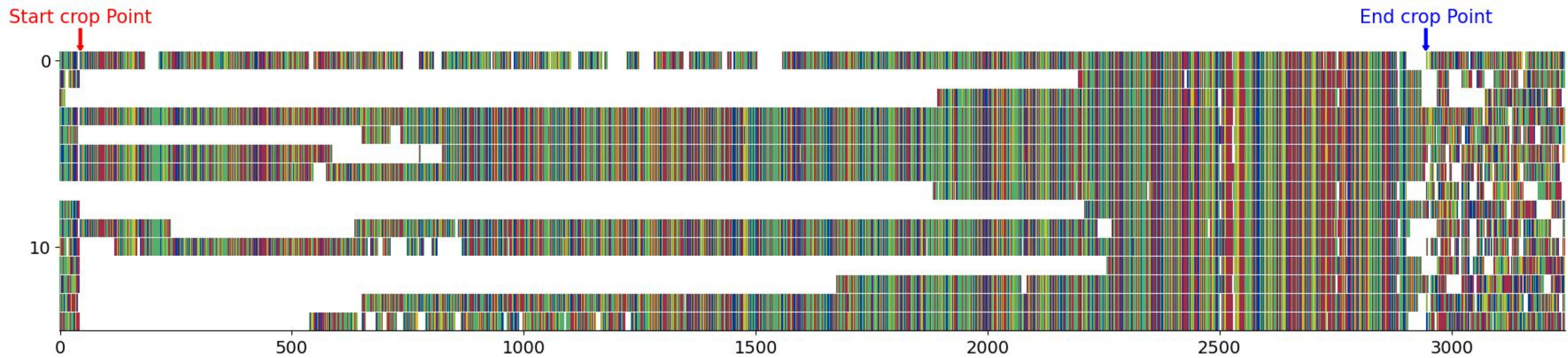
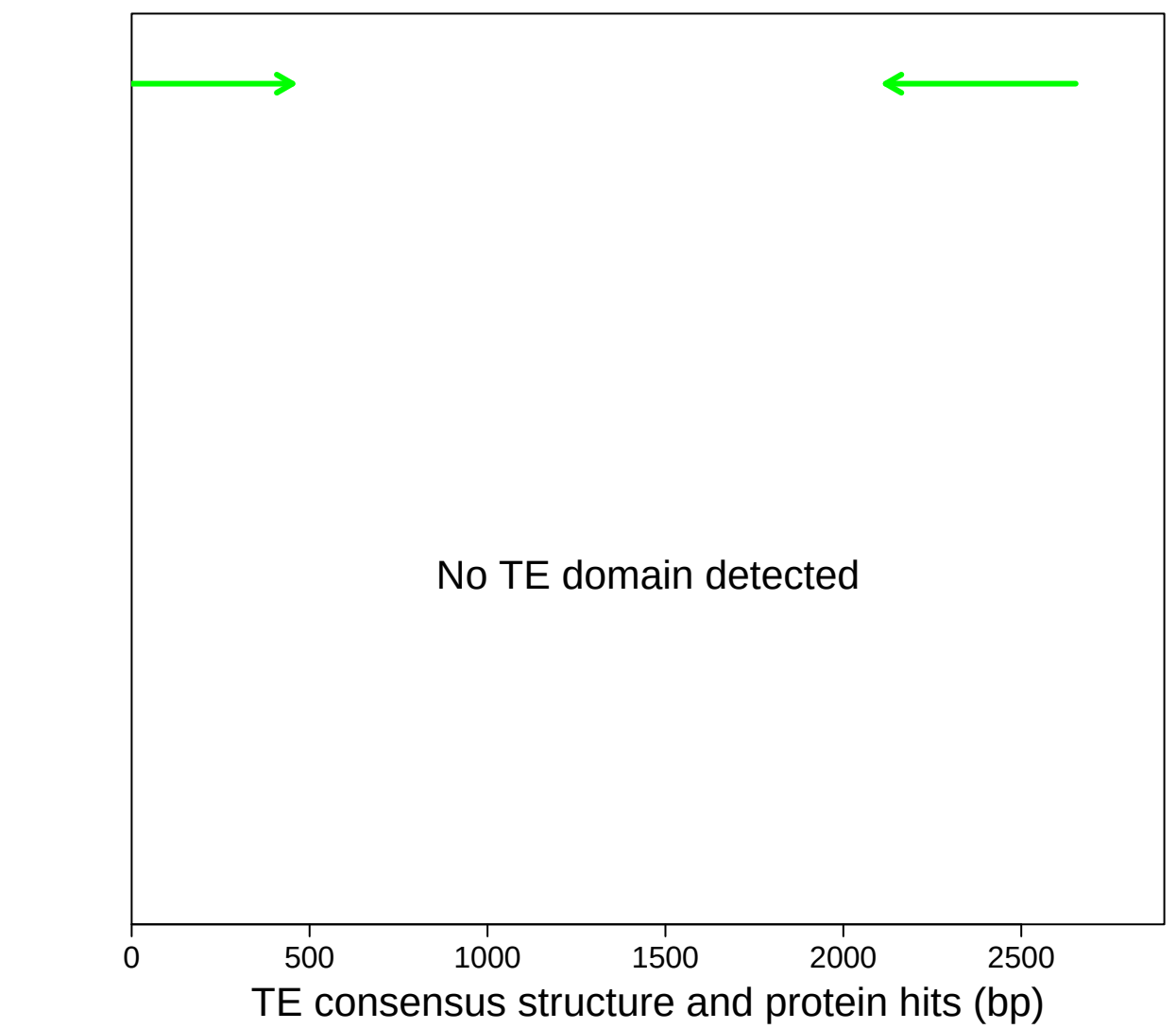
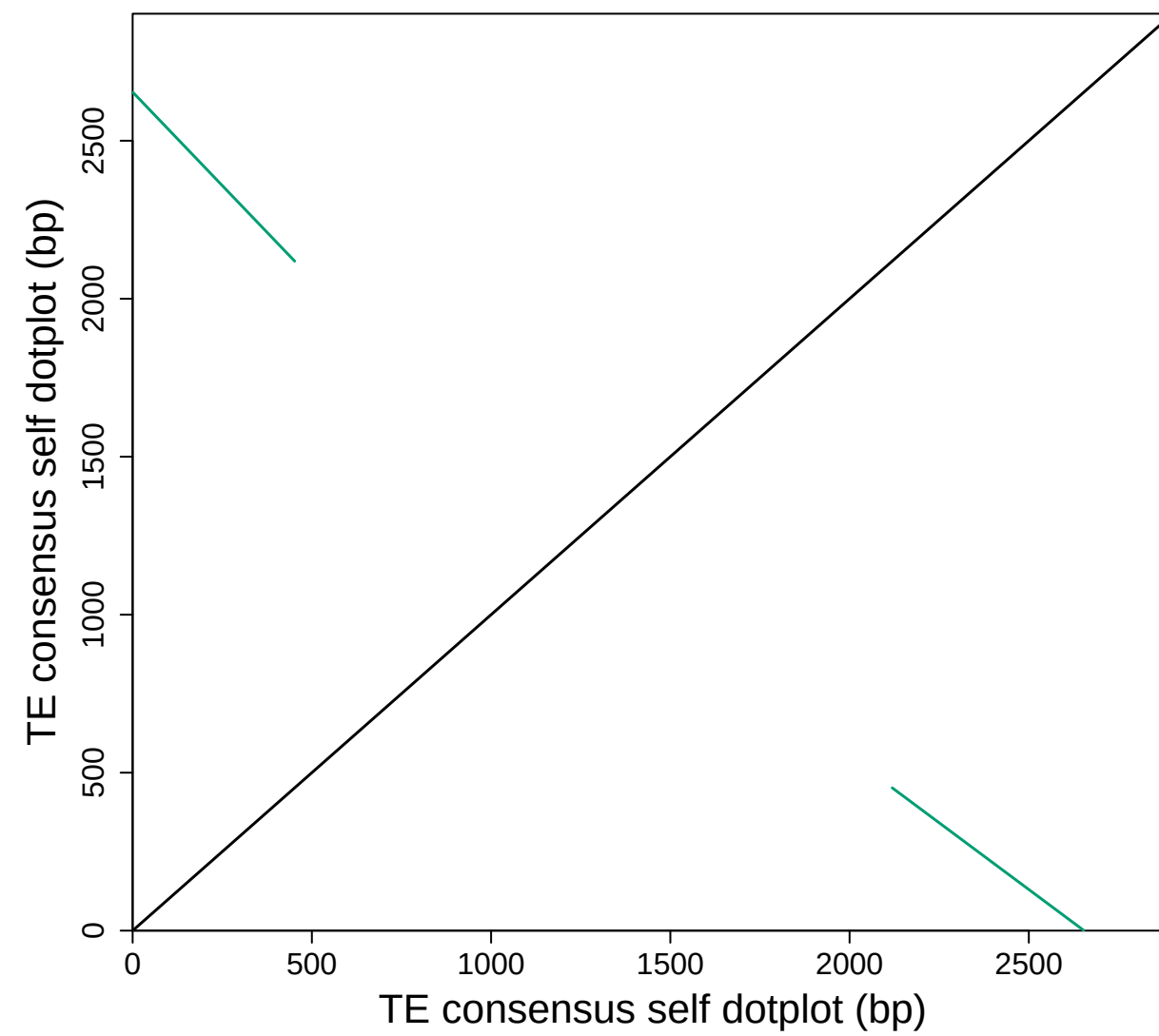
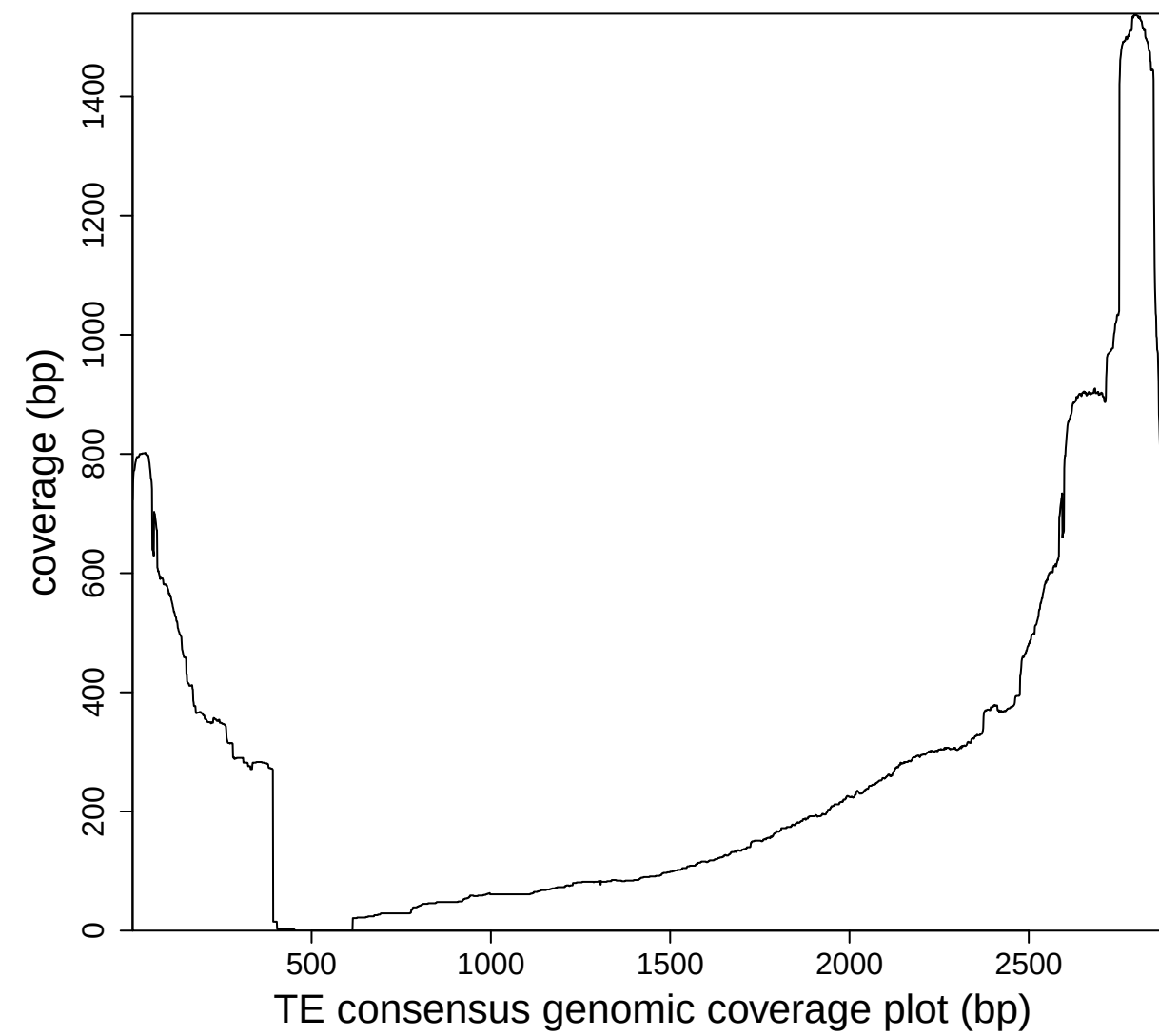
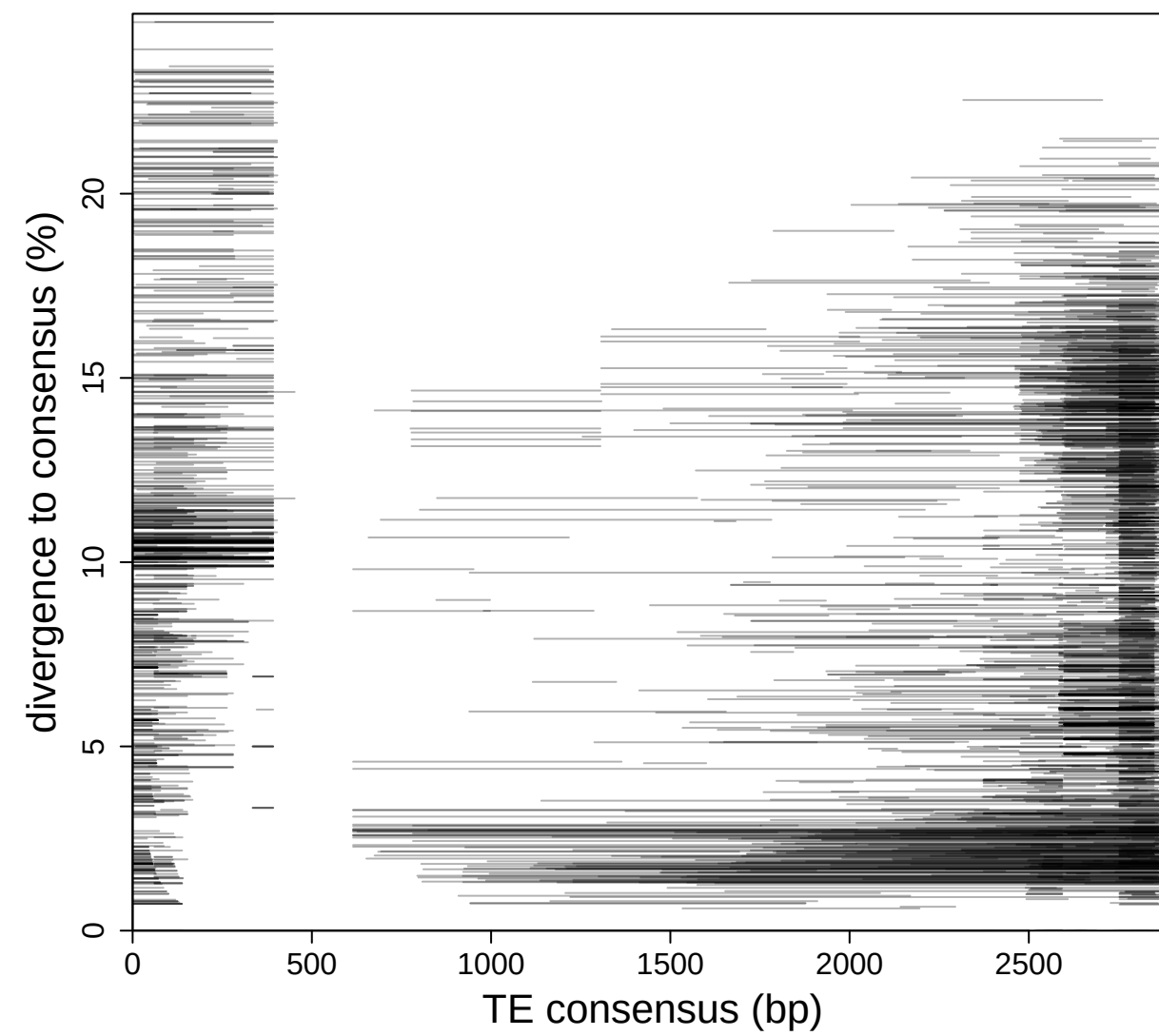




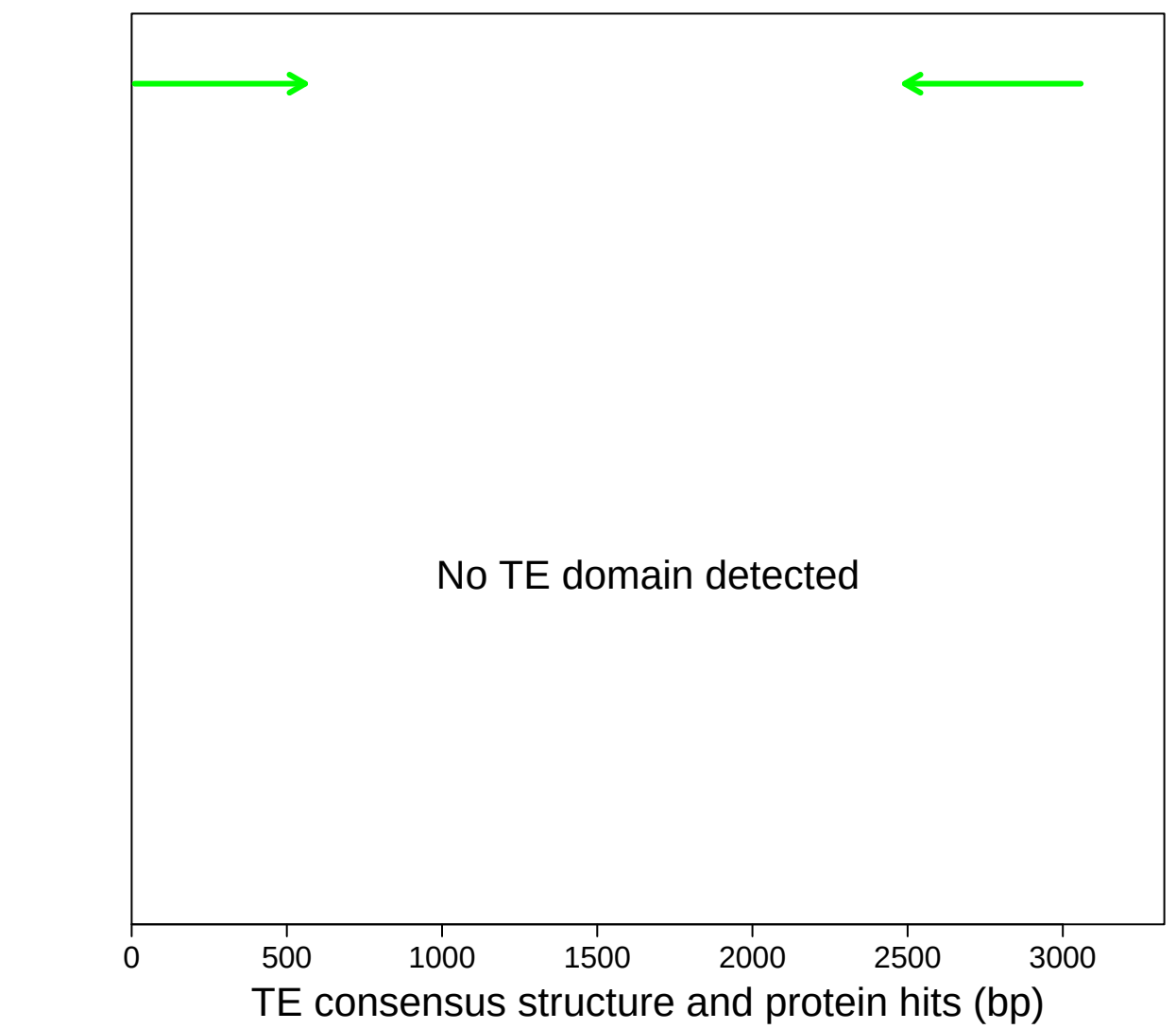
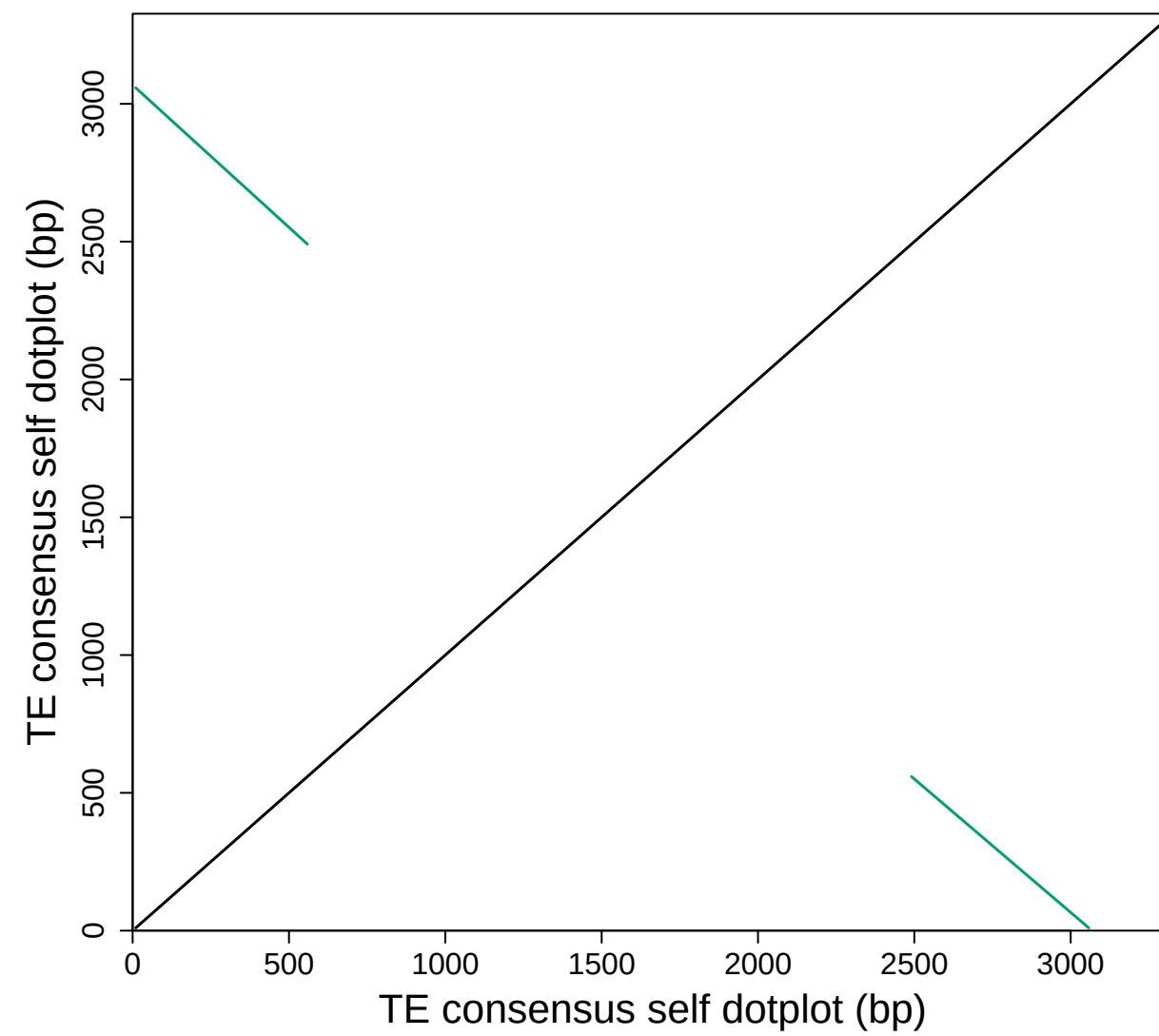
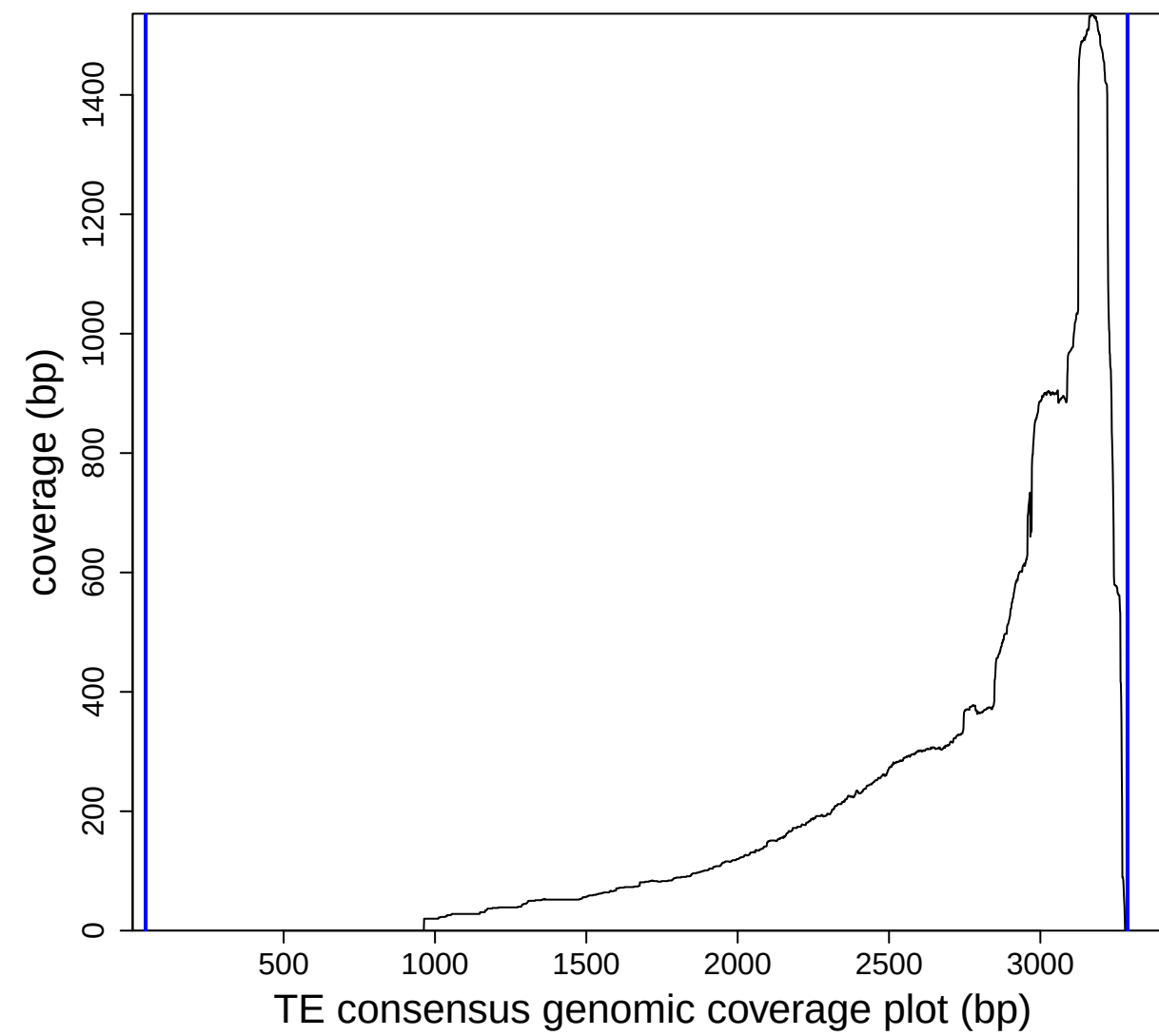
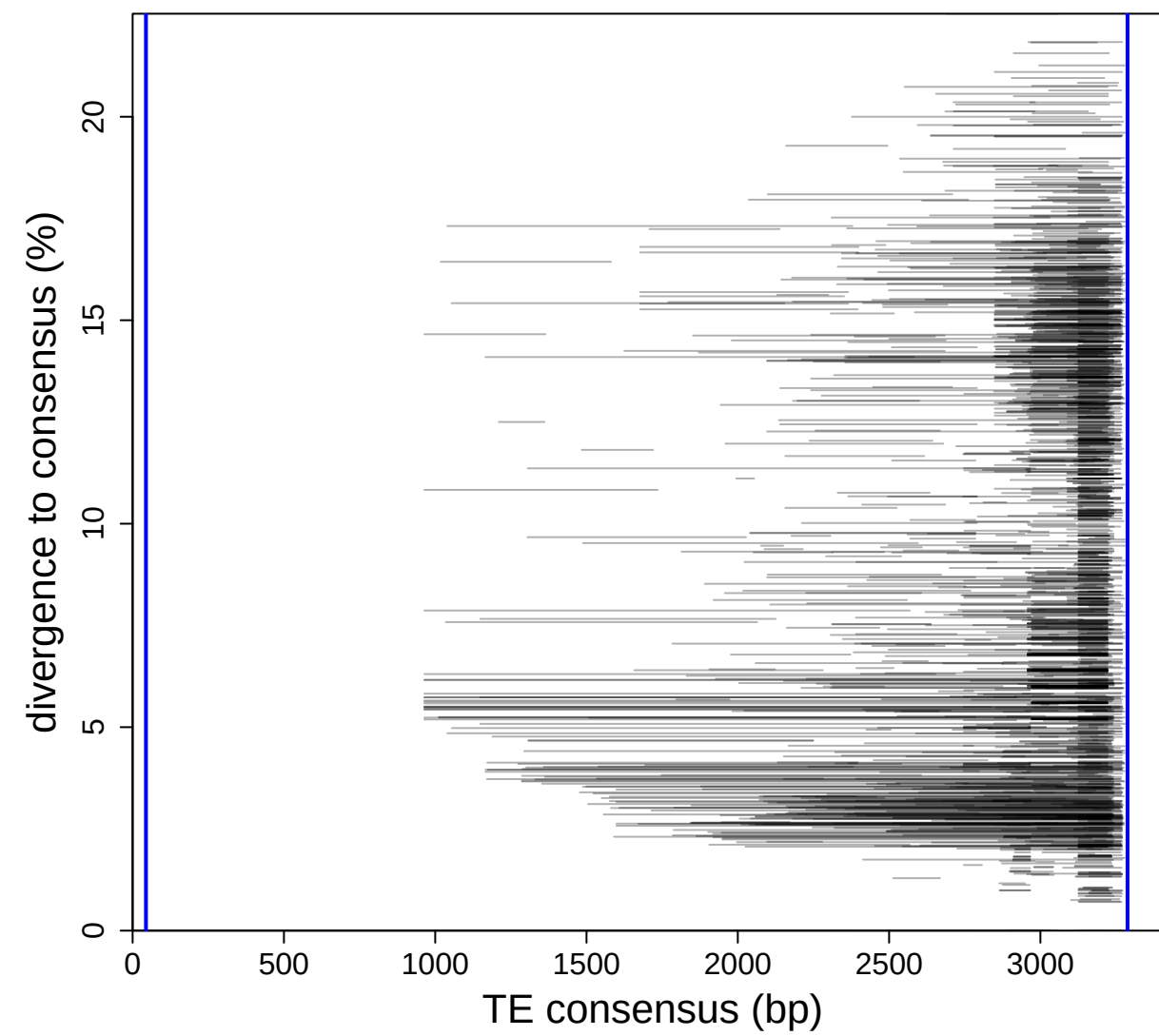
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 2902bp; fragments: 2901; full length: 0 (>=2611.8bp)

After TETrimmer 2902 bp

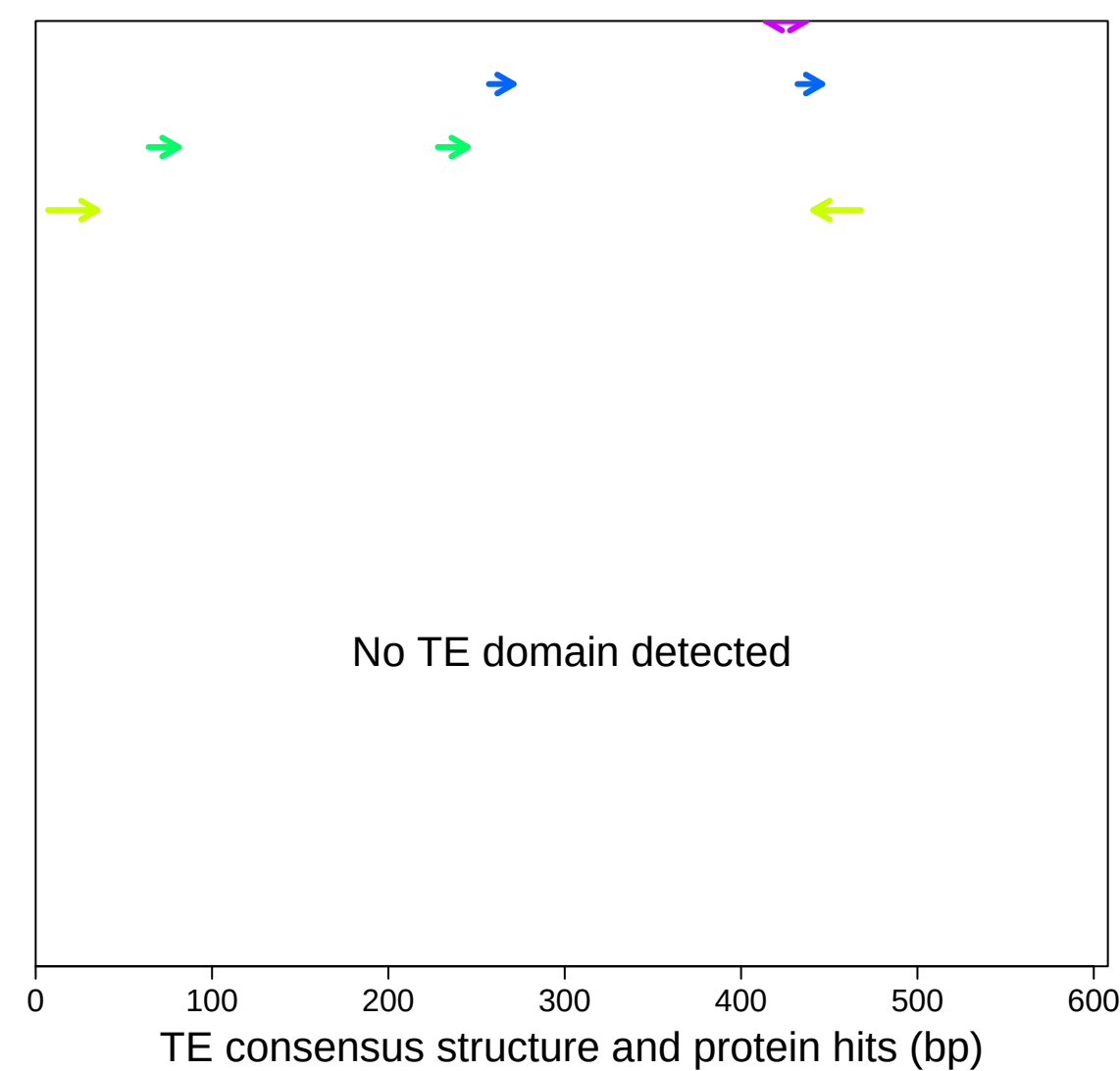
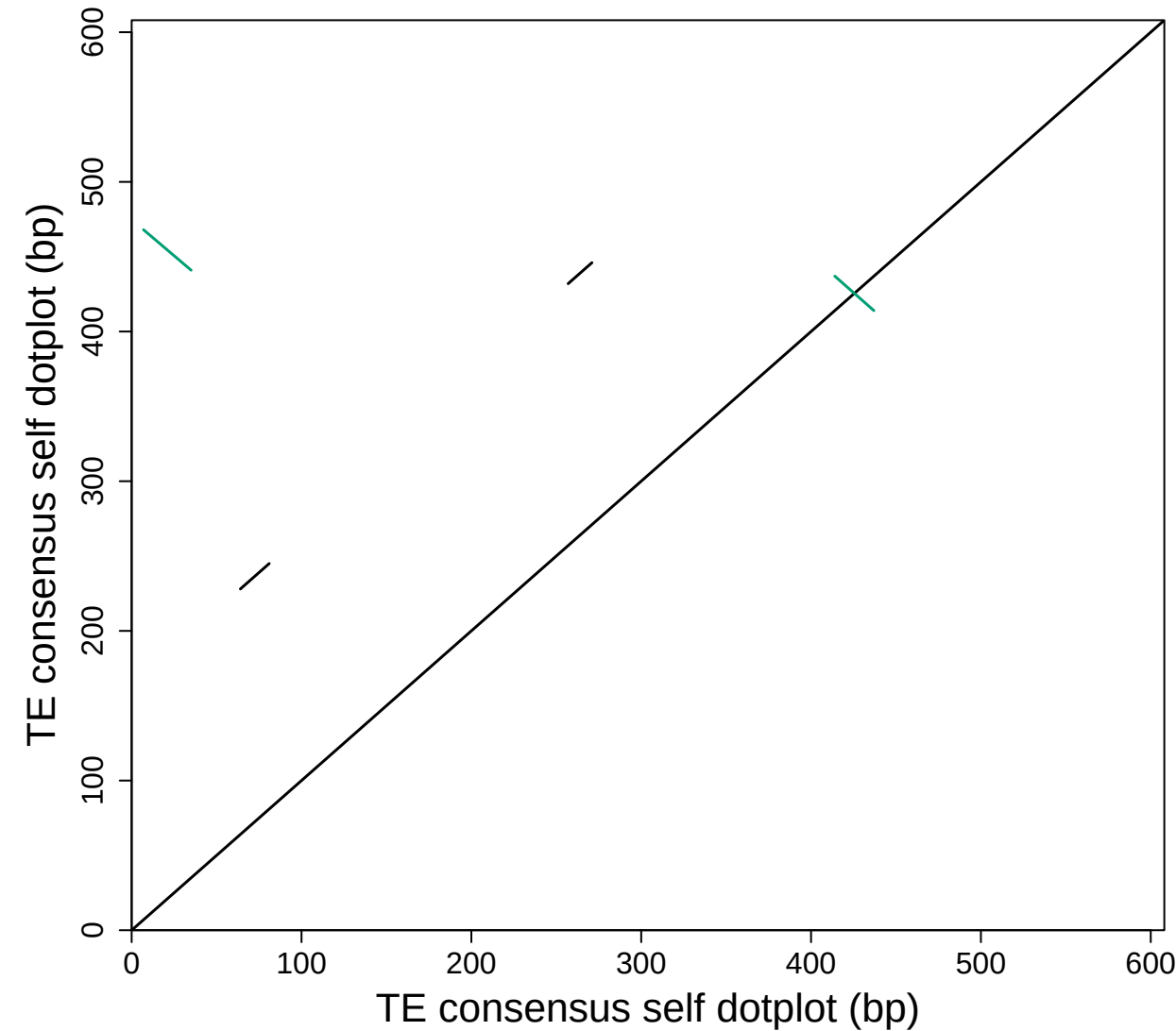
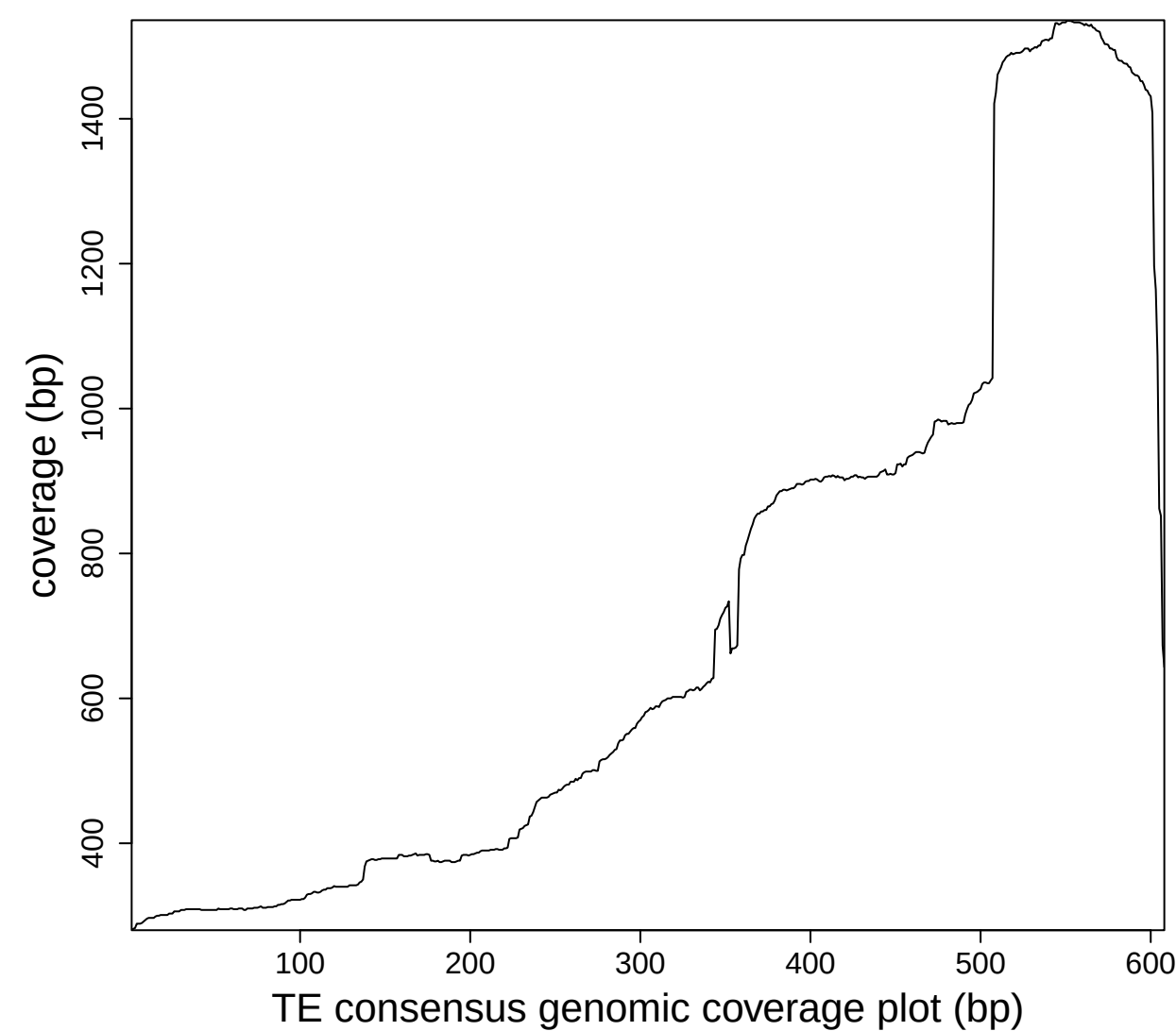
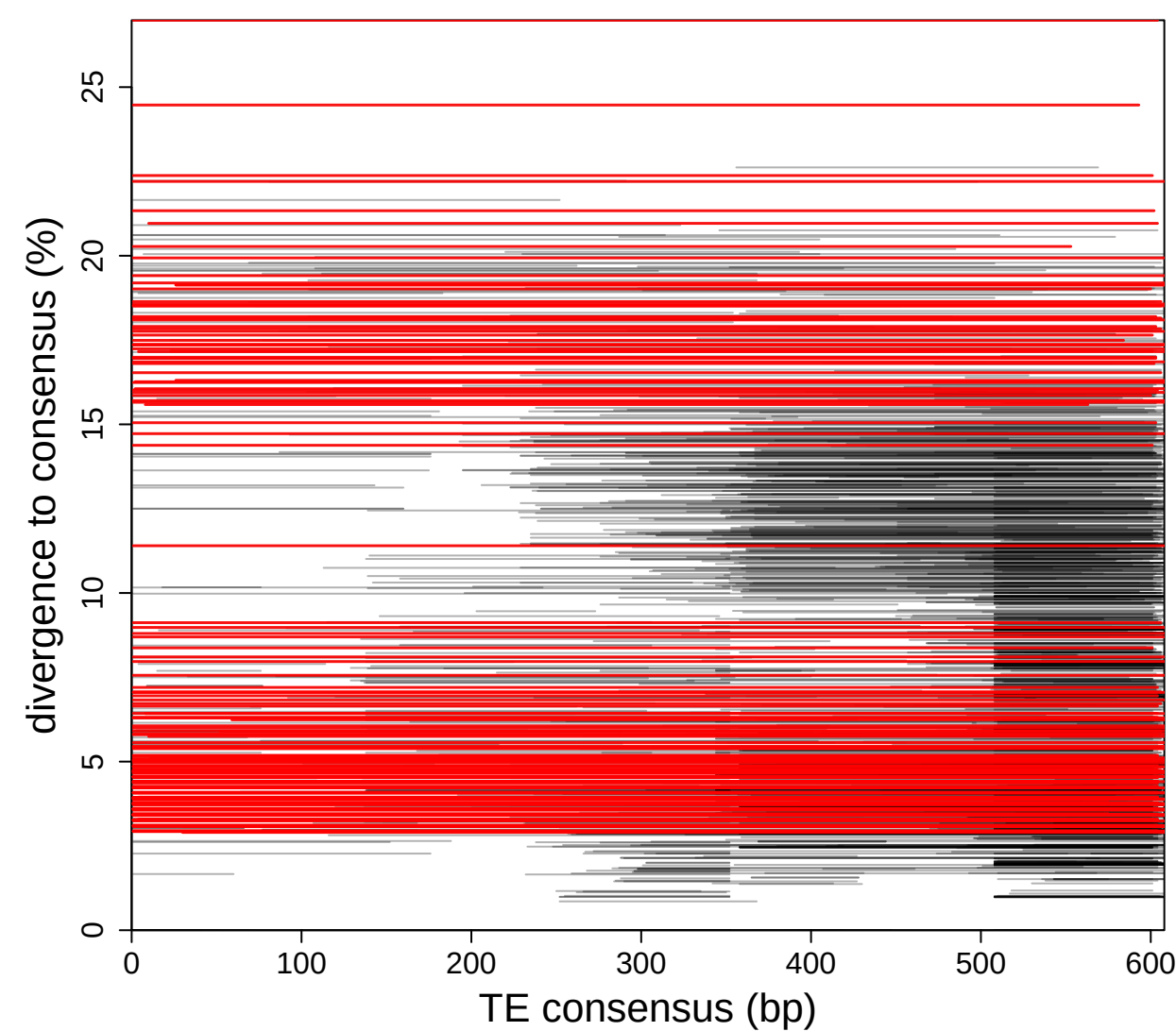


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 3438bp; fragments: 1906; full length: 0 (>=3094.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

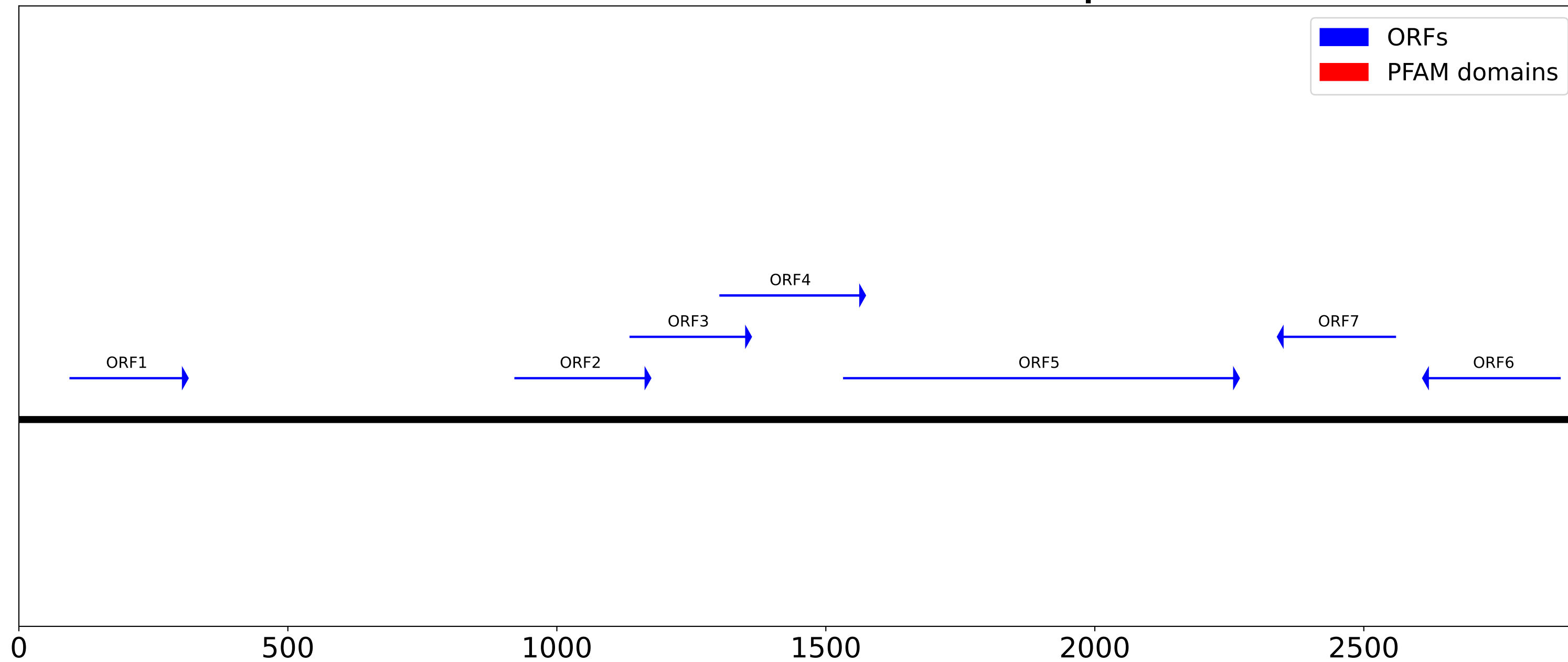


TE: rnd_1_family_915
size: 608bp; fragments: 1829; full length: 233 (≥ 547.2 bp)

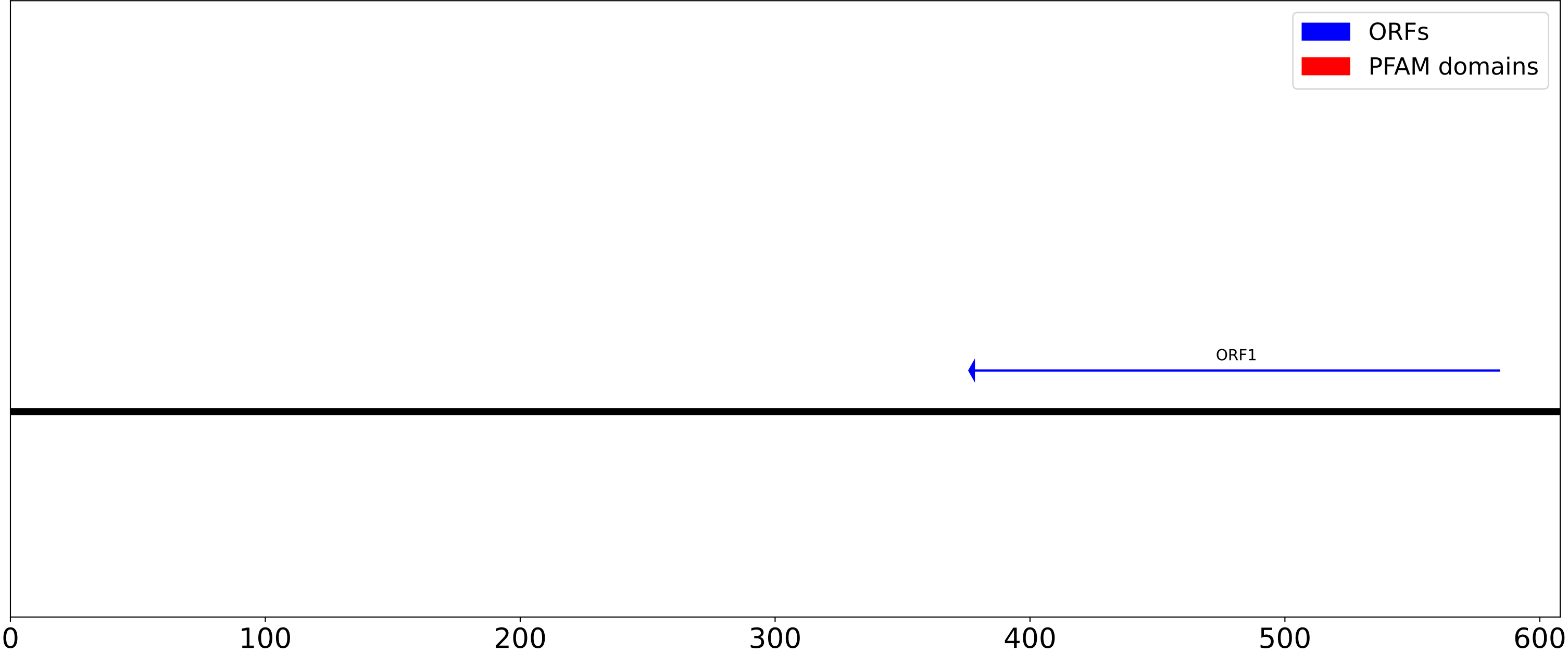


Before TEtrimmer 608 bp

After TEtrimmer ORF and PFAM domain plot



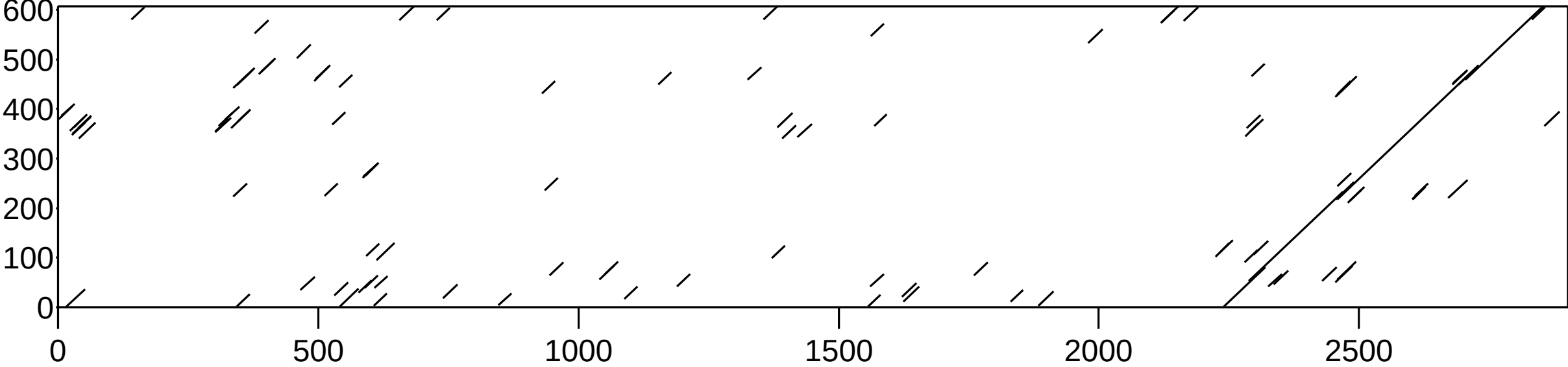
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)