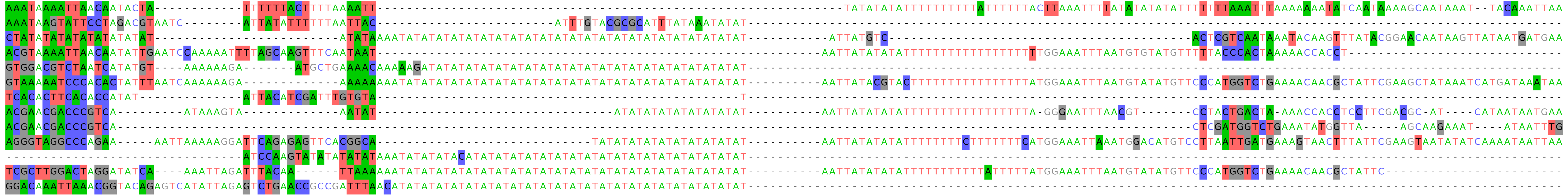
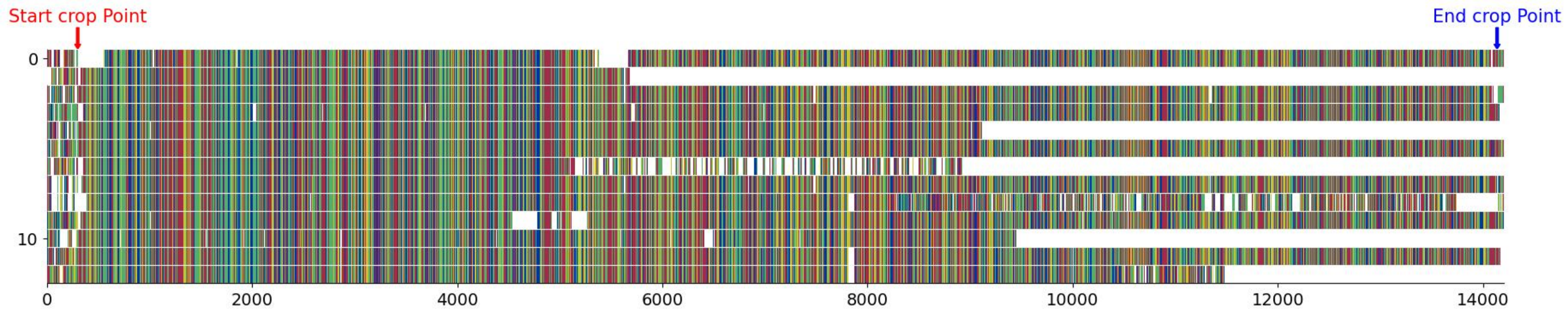


End crop Point

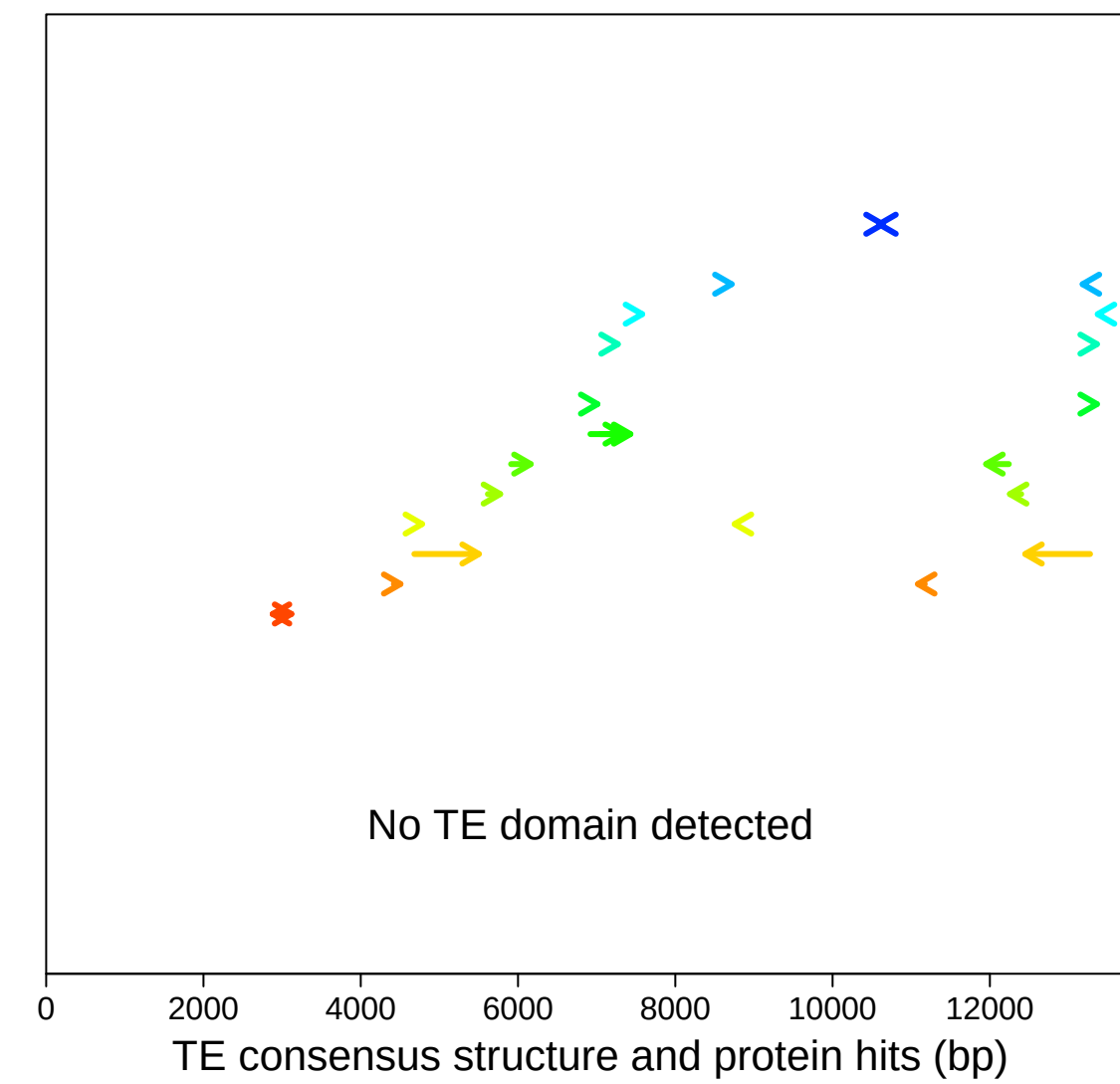
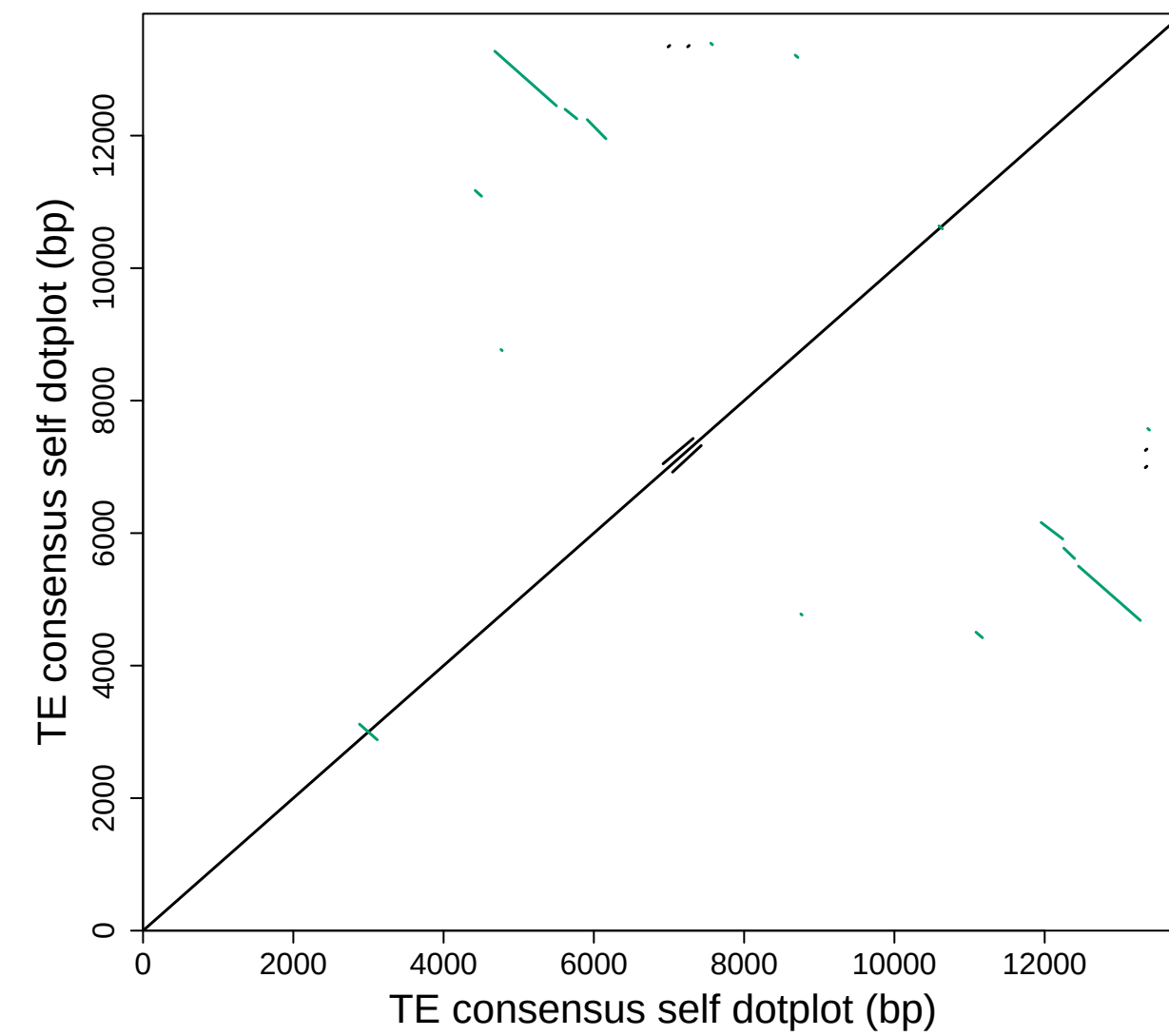
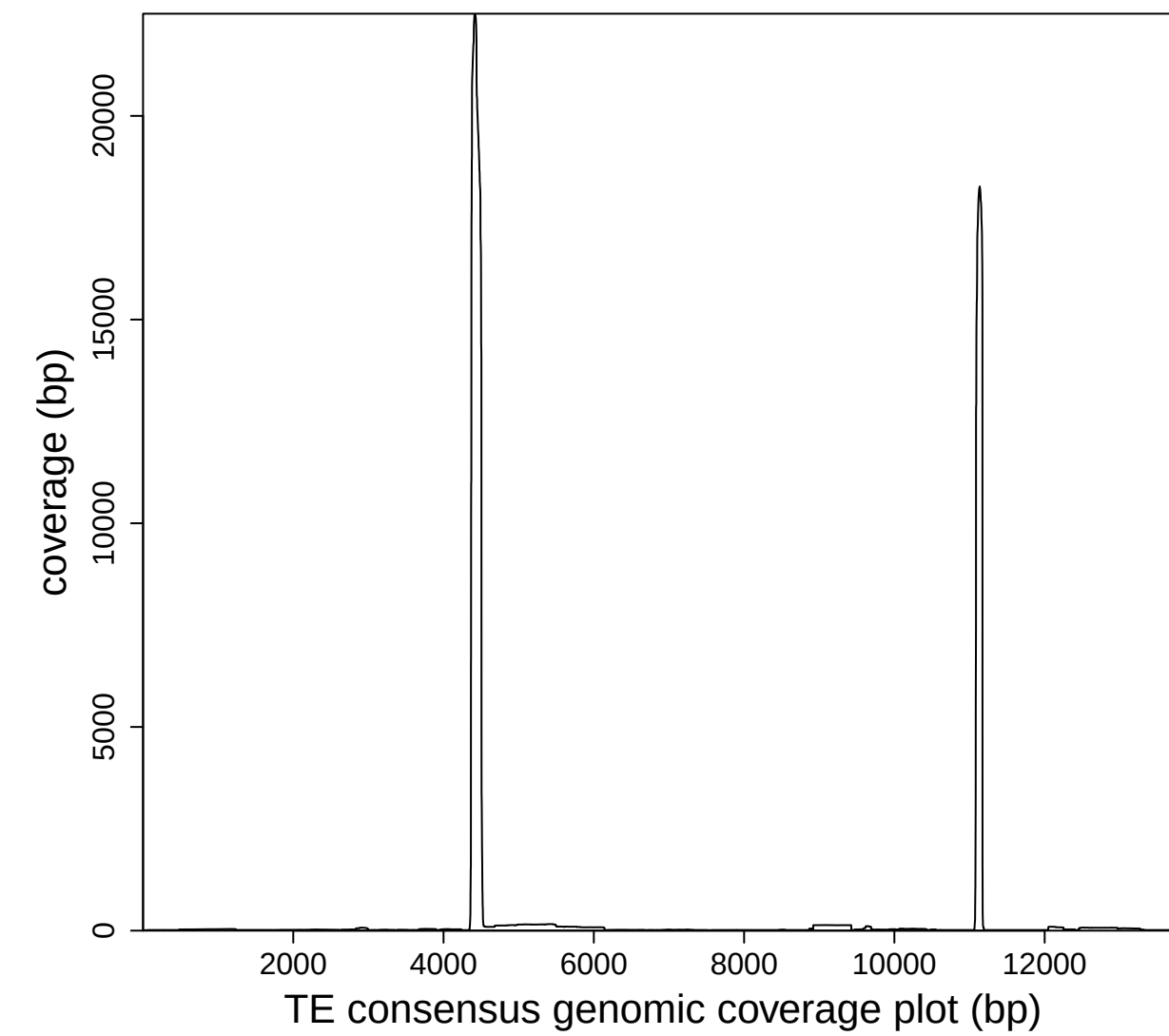
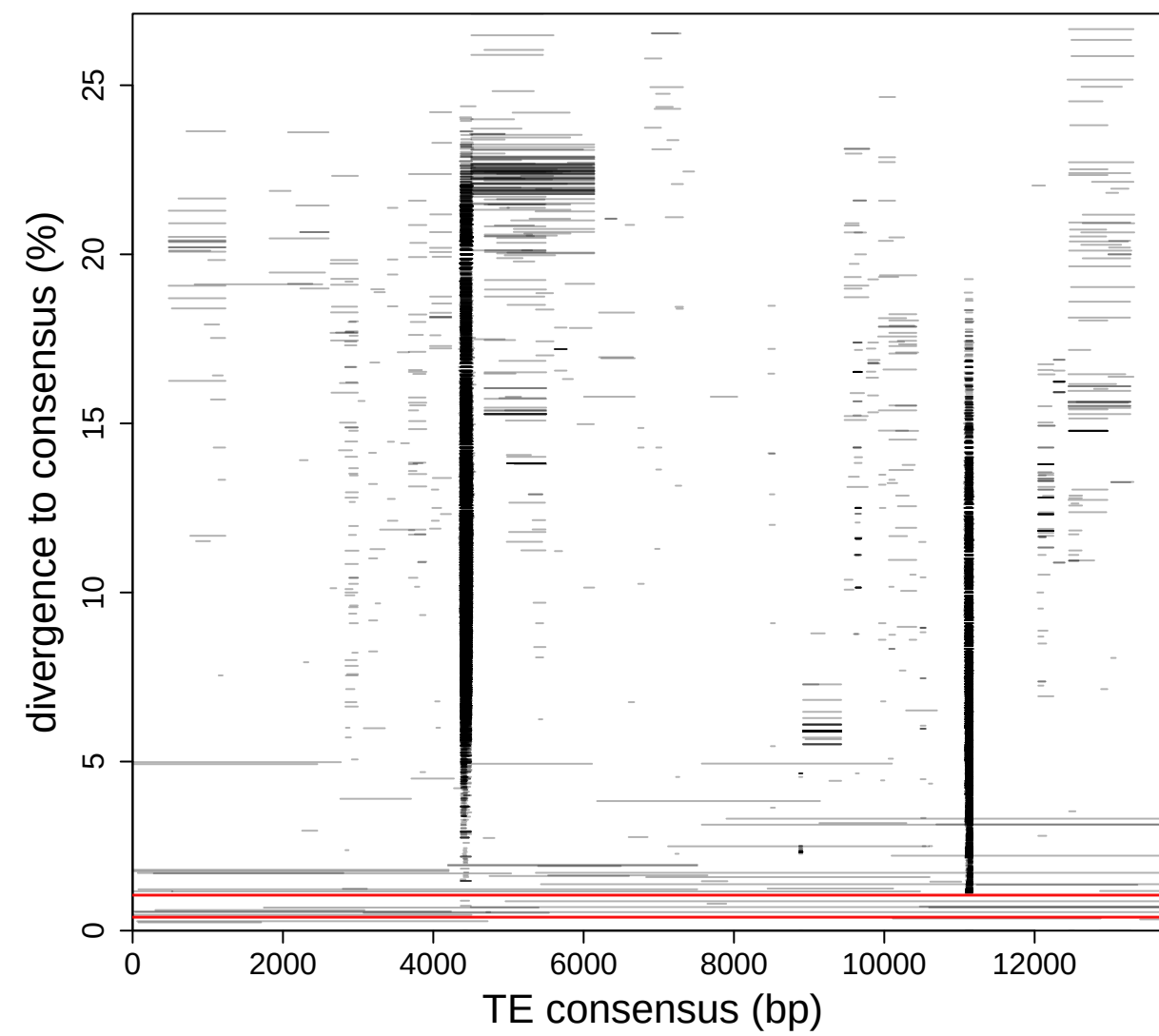
MSA length = 13840





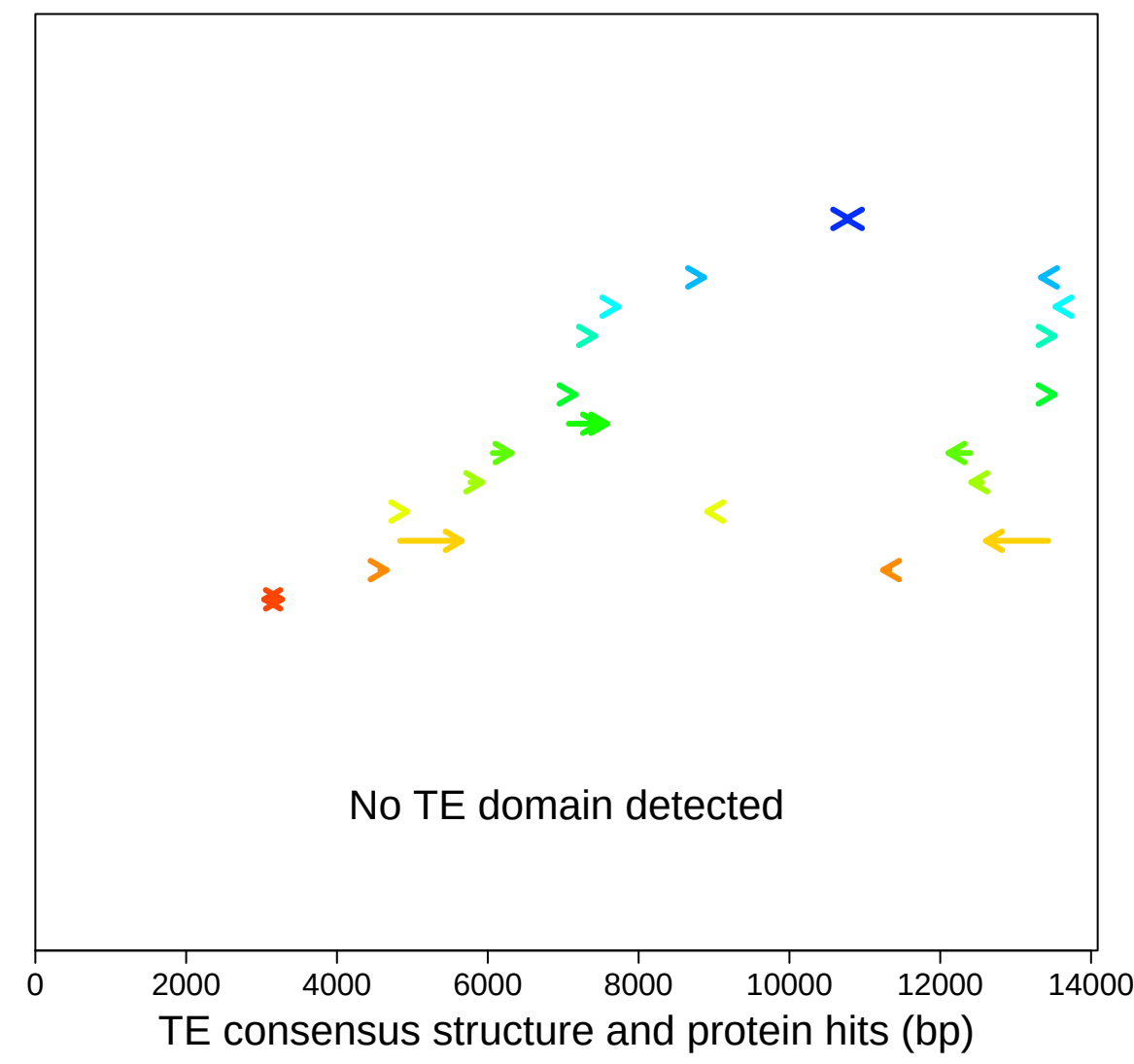
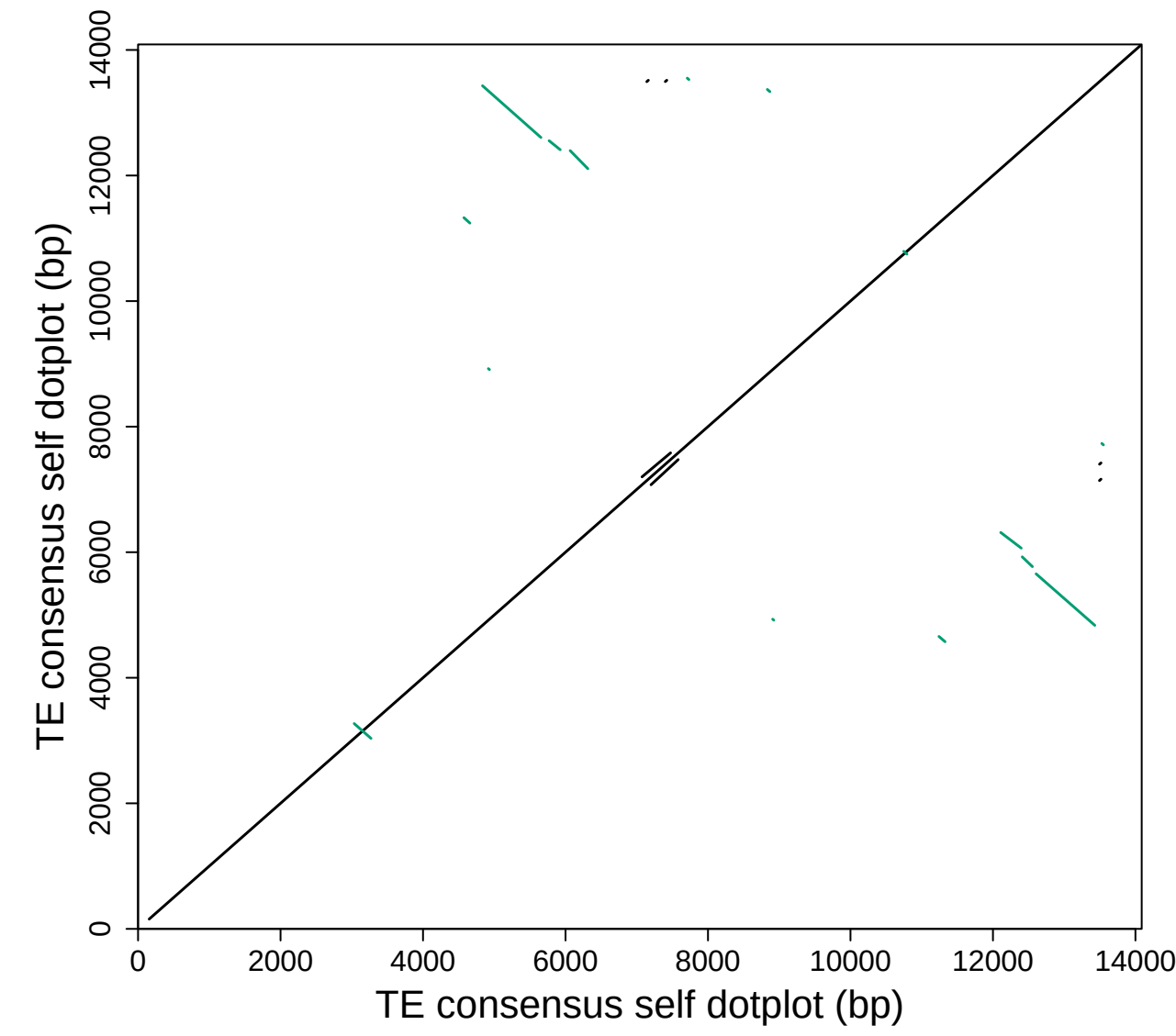
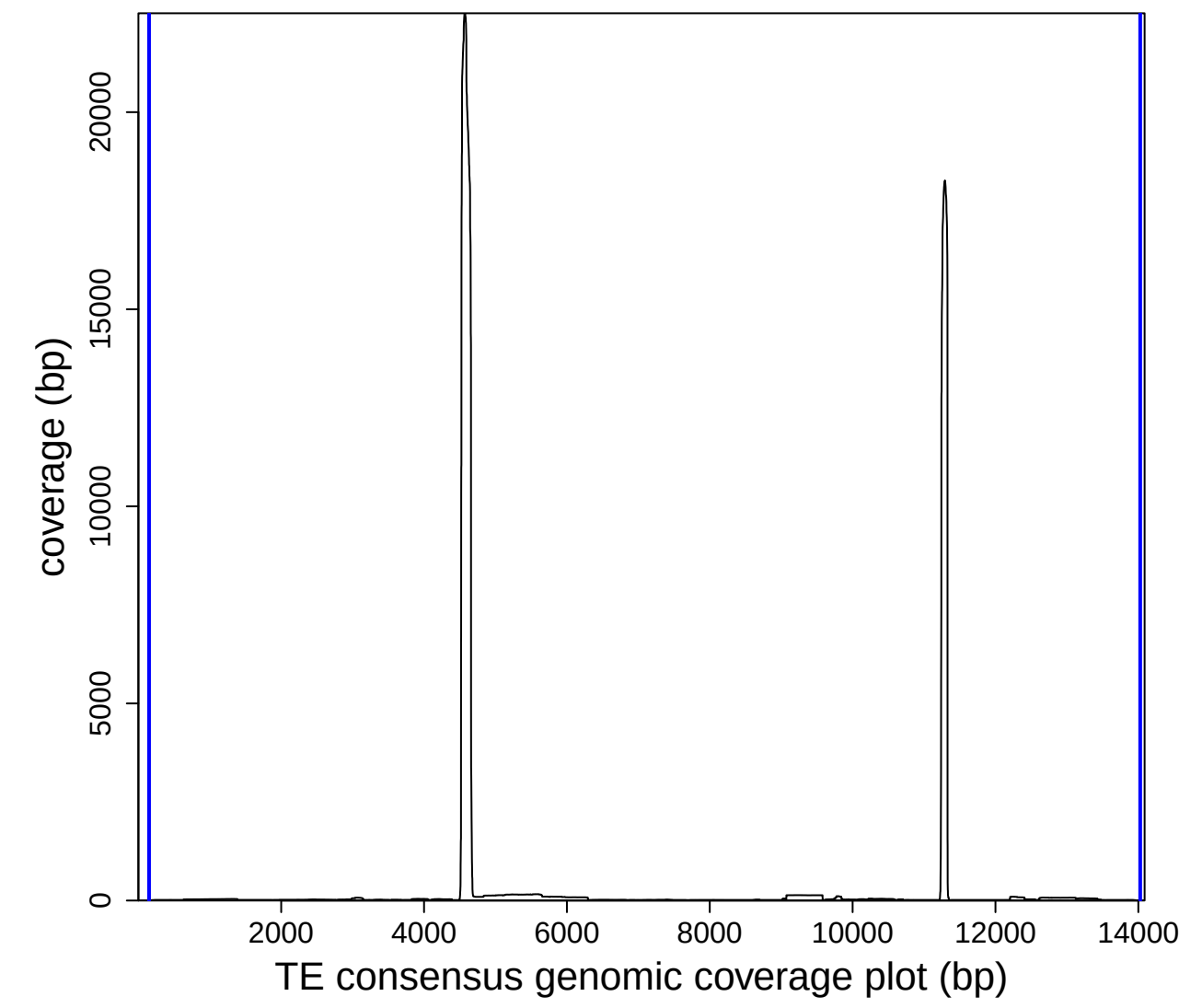
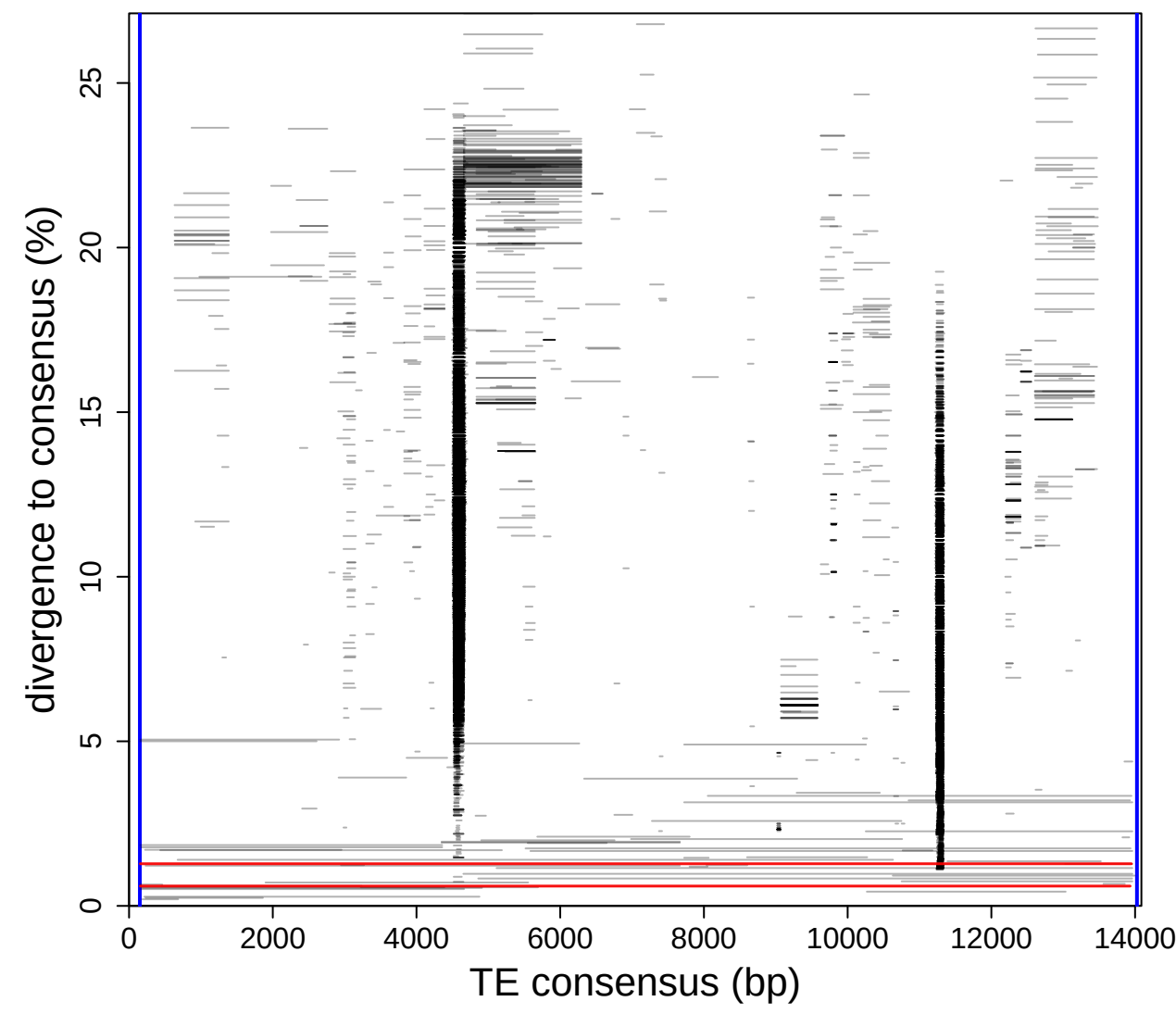
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa
size: 13840bp; fragments: 41779; full length: 2 (>=12456bp)

After TEtrimmer 13840 bp

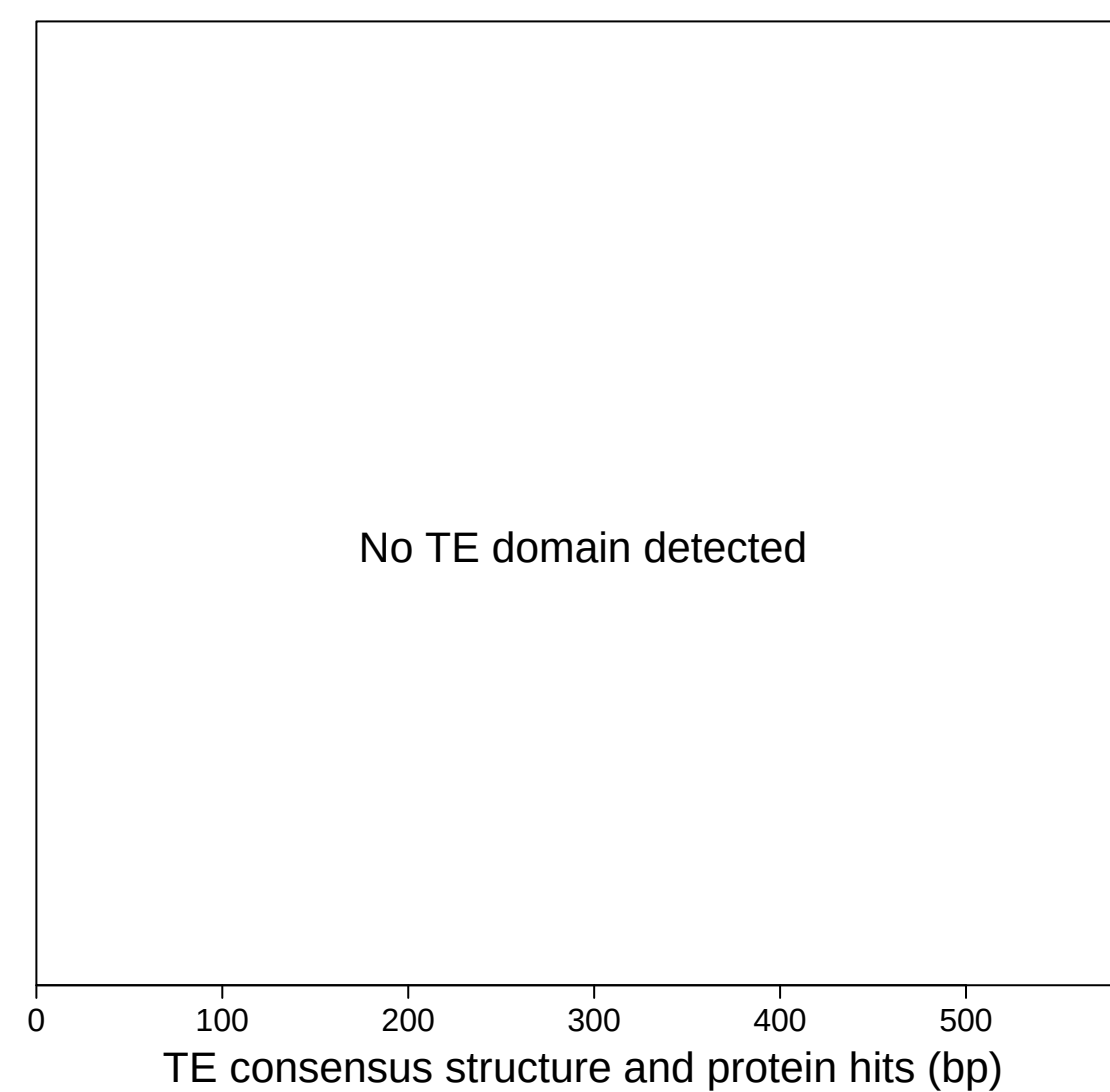
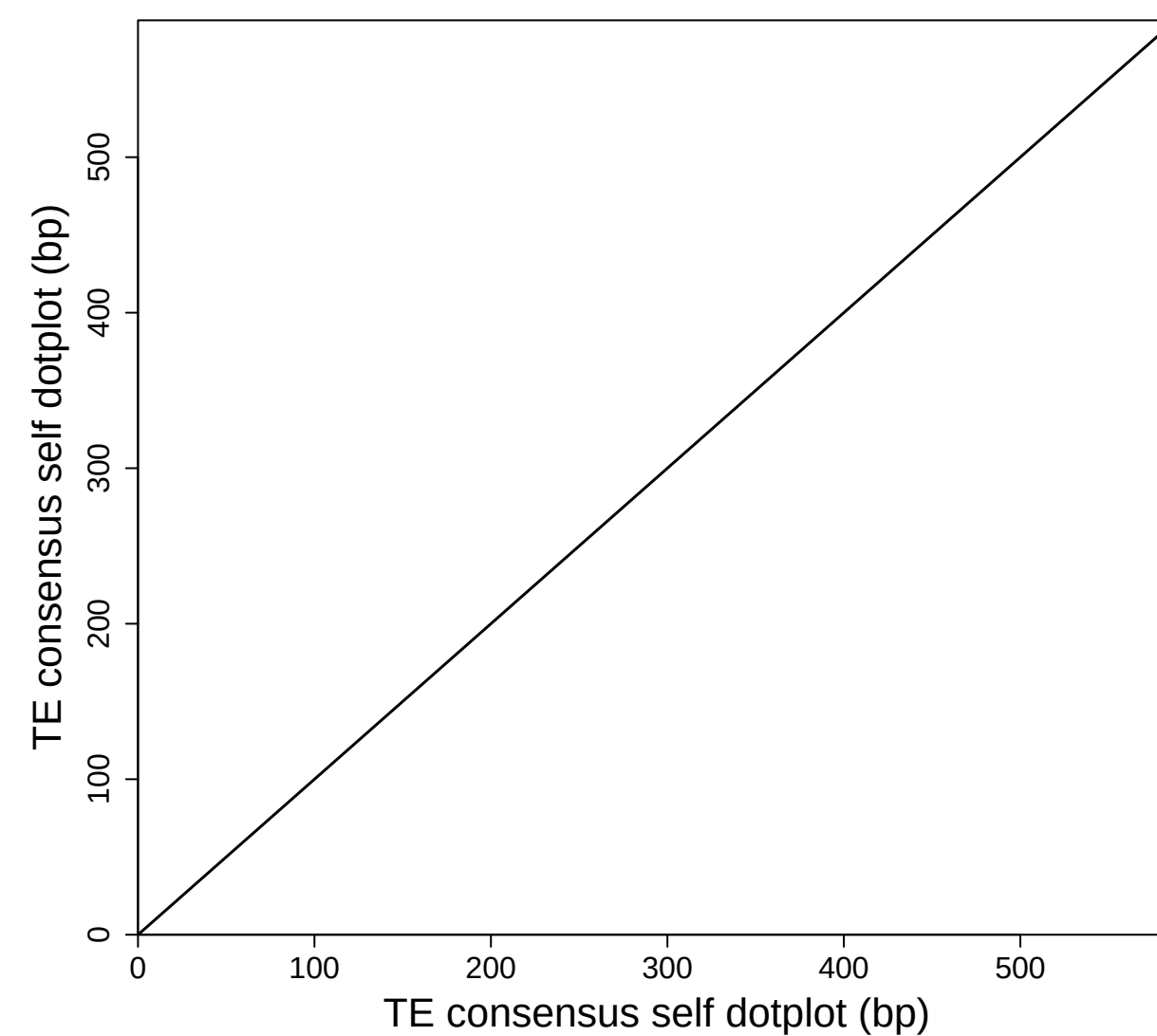
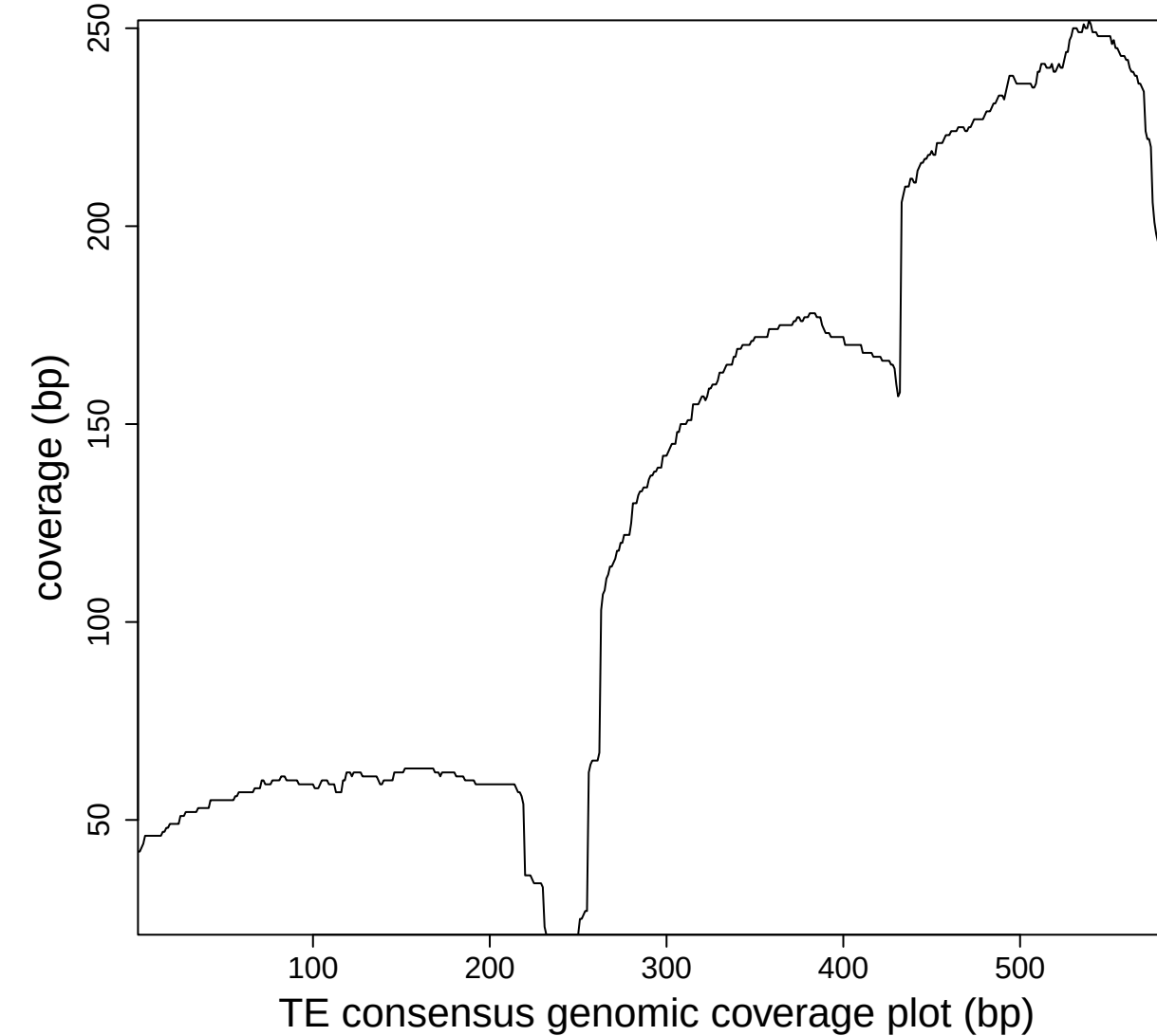
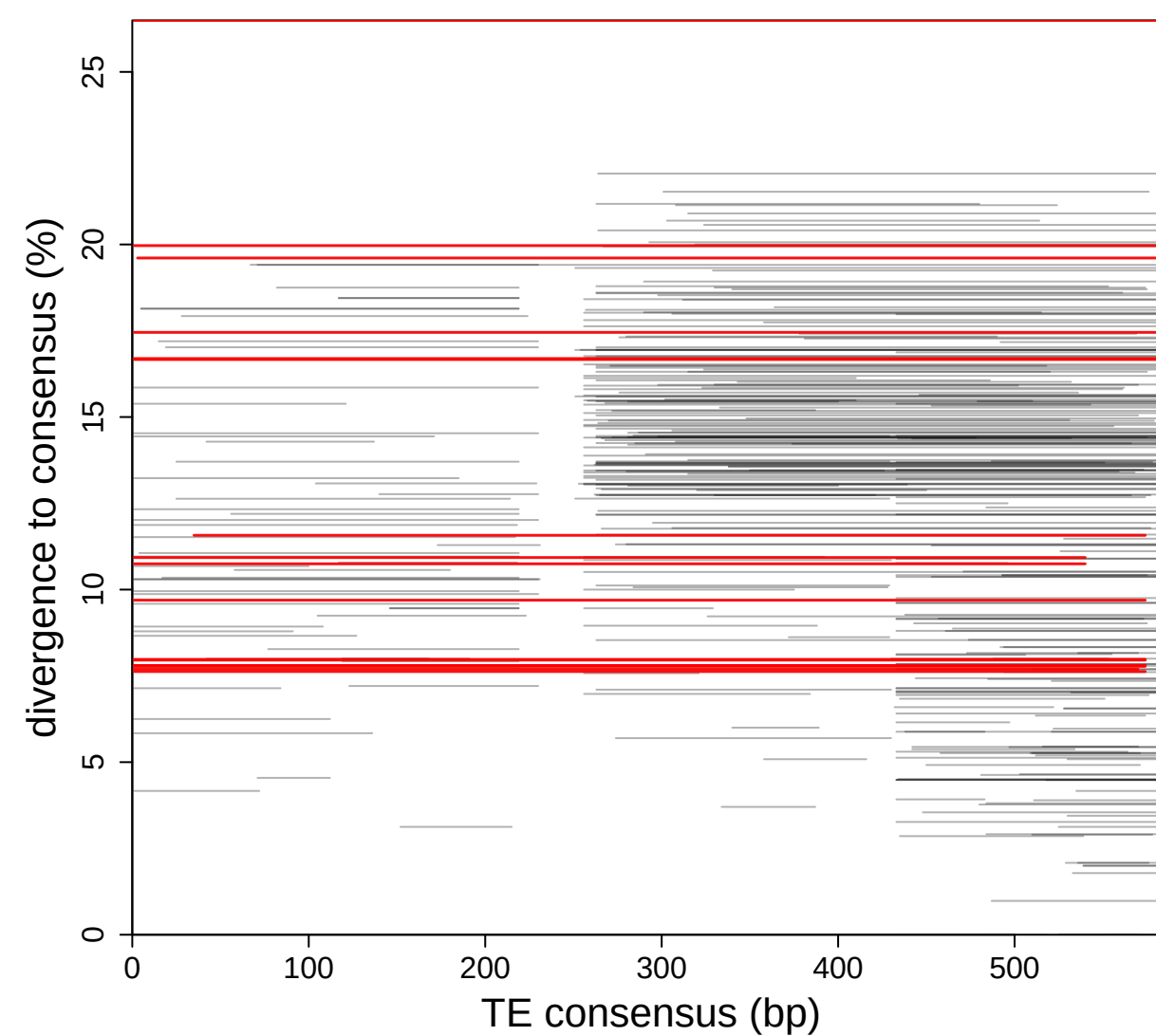


uf.bed g 1.bed fm 1.bed 0 0 n.bed g 1.bed fm 2.bed 0 0 bc
size: 14088bp; fragments: 41769; full length: 2 (>=12679.2bp)

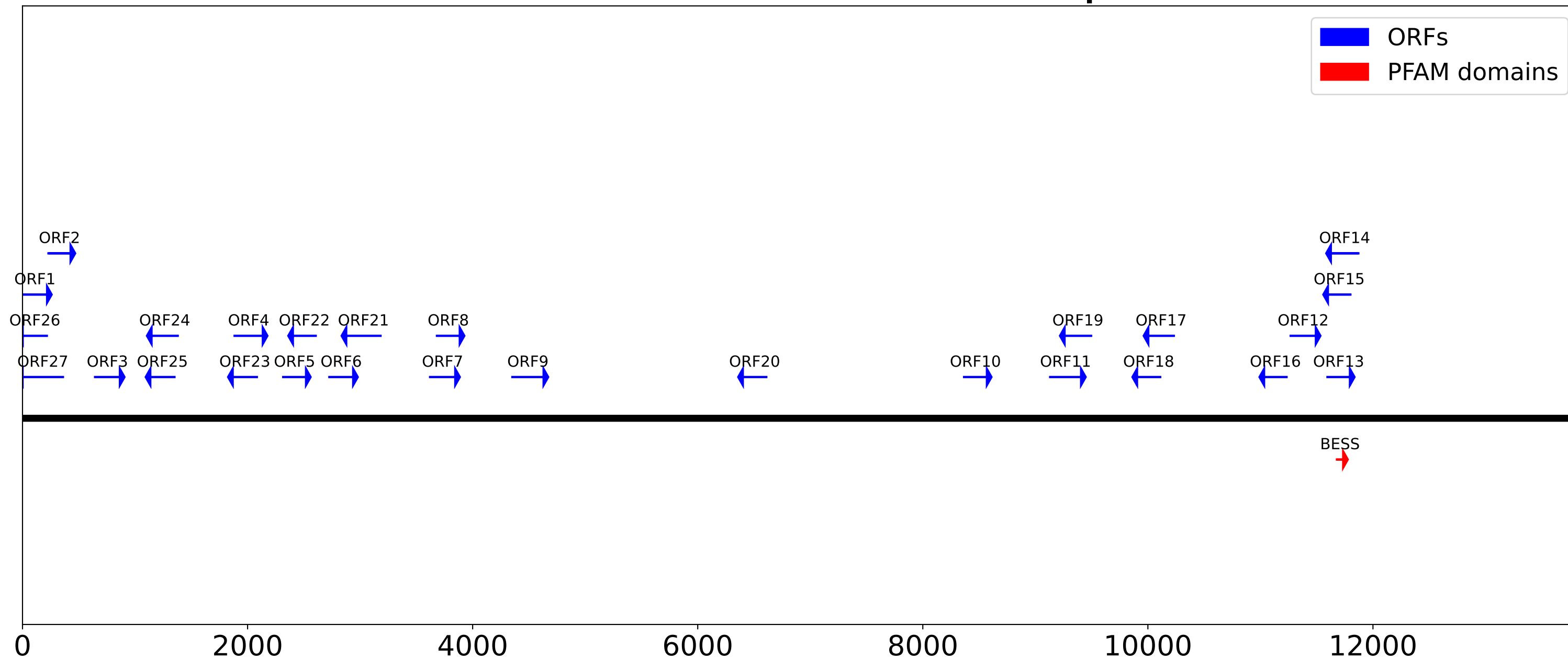
After TEtrimmer Extended plot Blue lines are boundaries



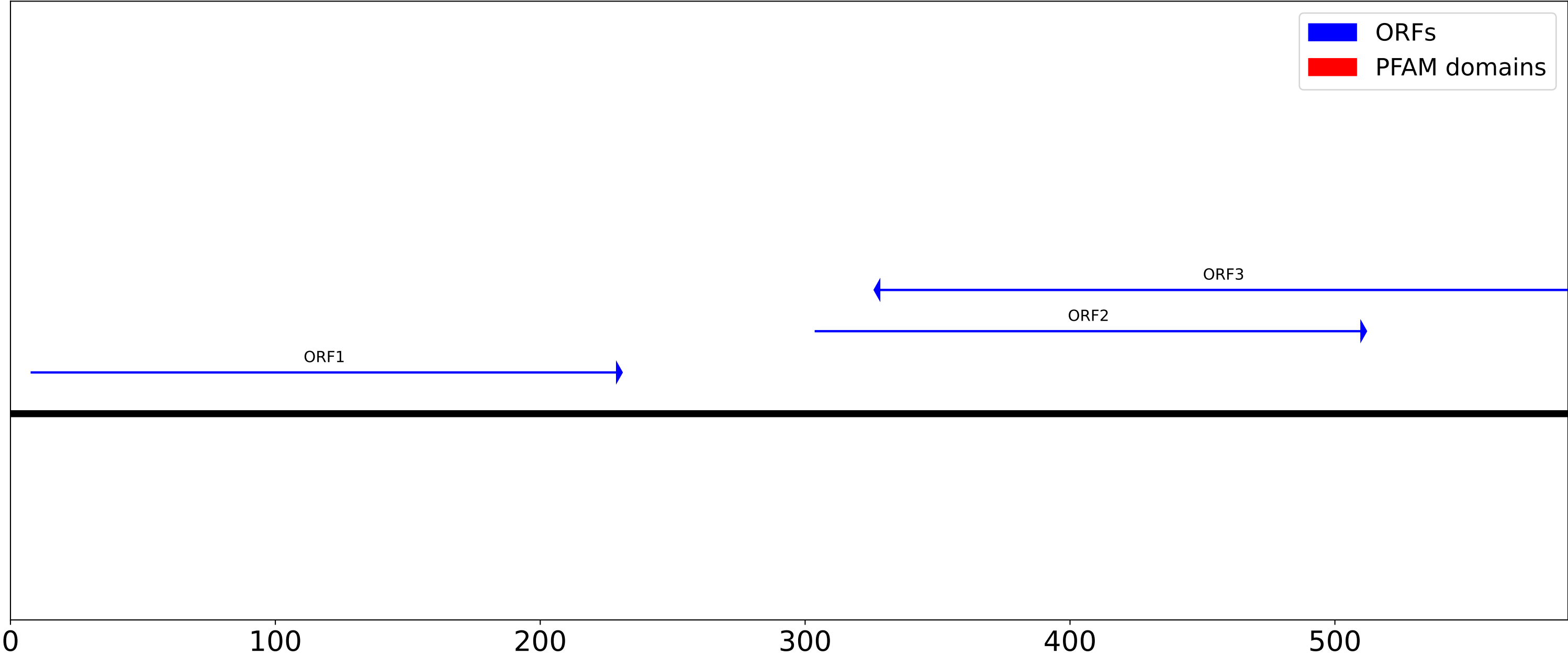
TE: rnd_5_family_13152
size: 588bp; fragments: 359; full length: 20 (≥ 529.2 bp)



After TEtrimmer ORF and PFAM domain plot



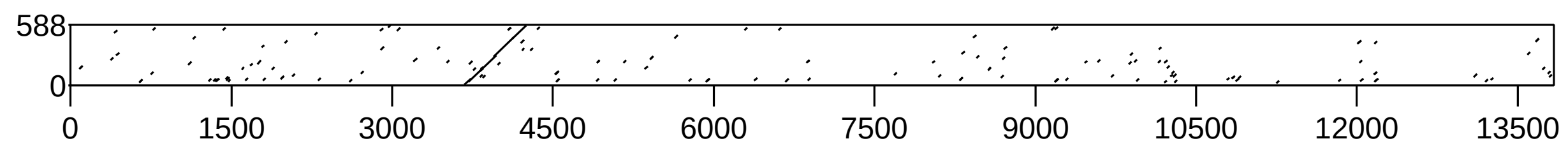
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)