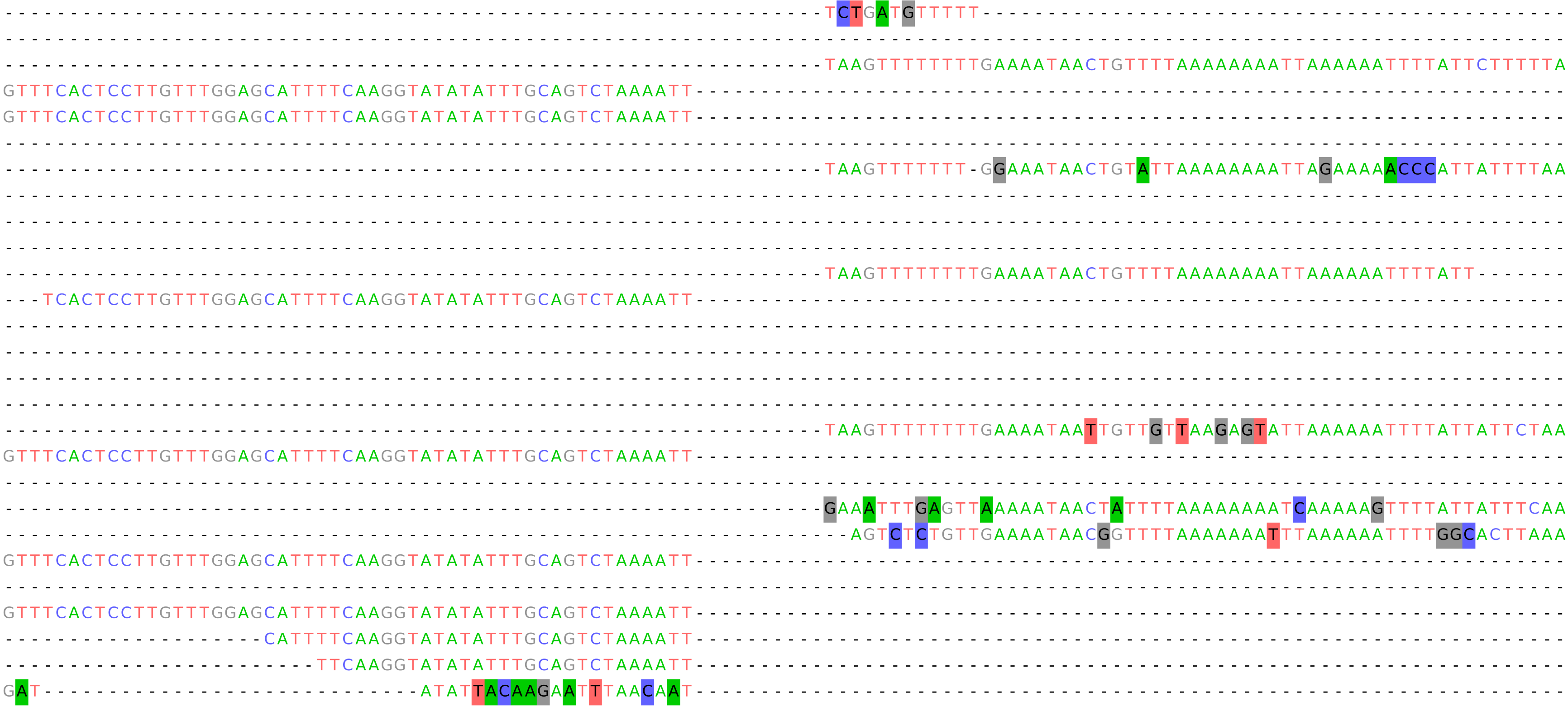


Start crop Point

End crop Point



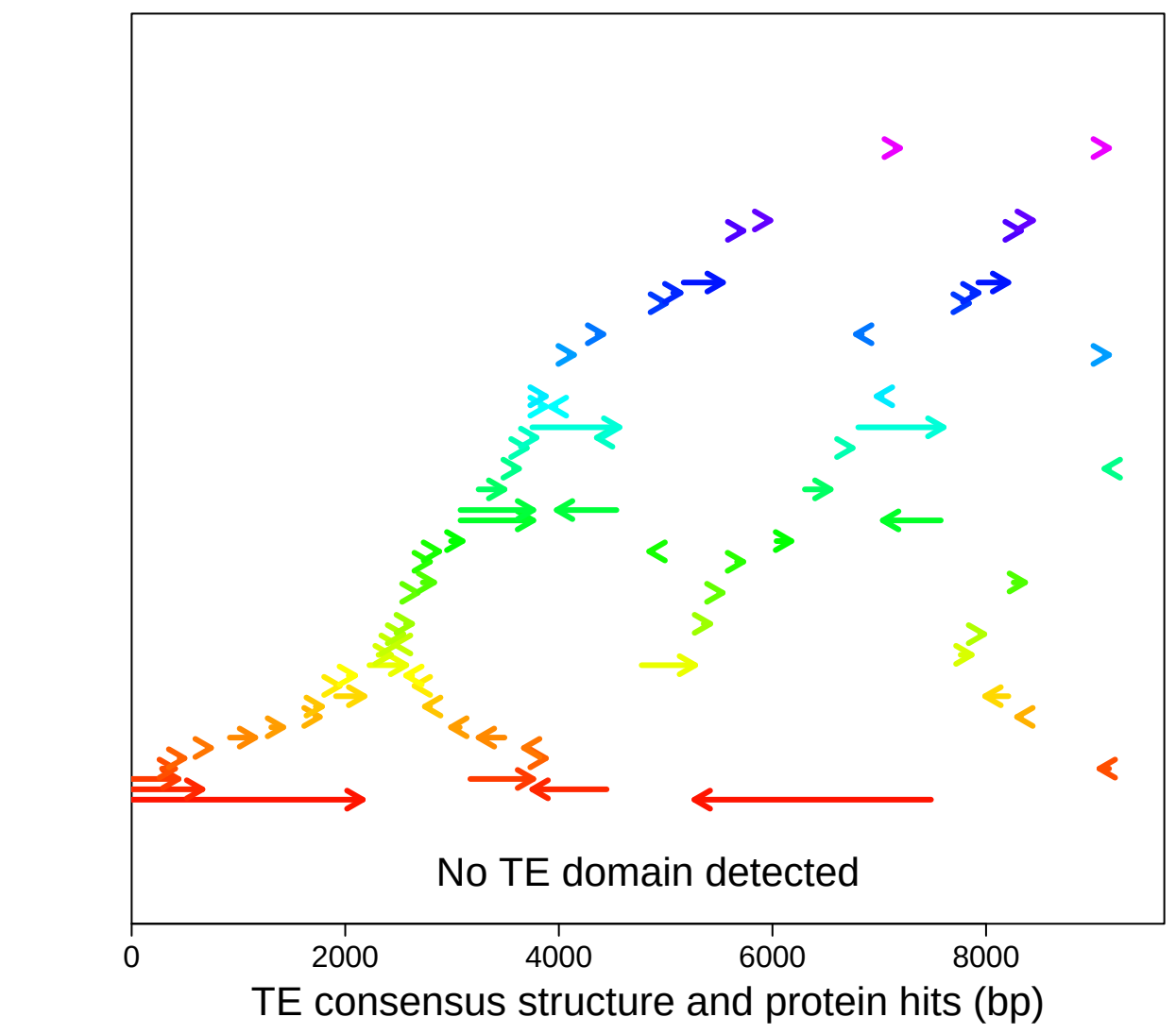
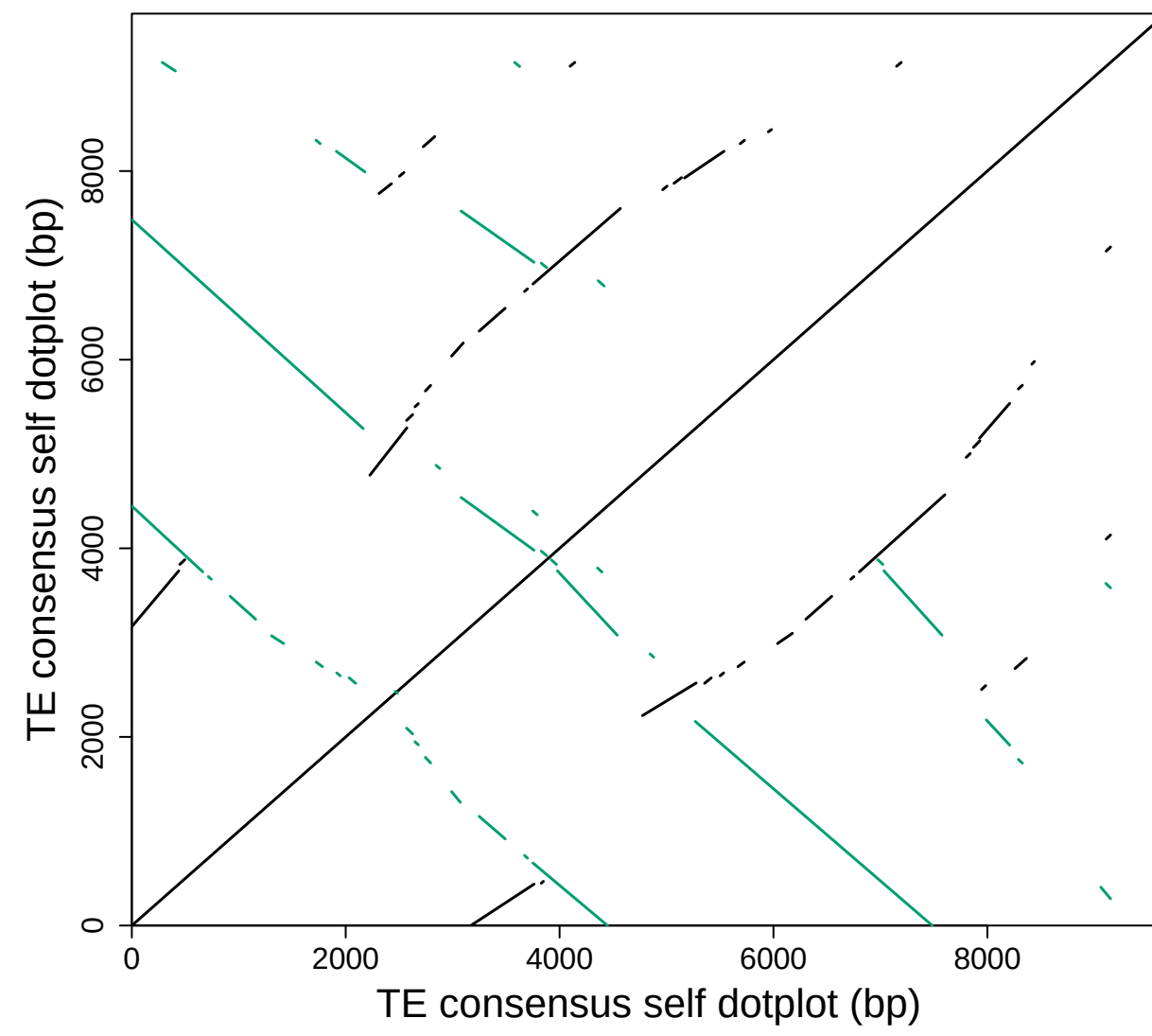
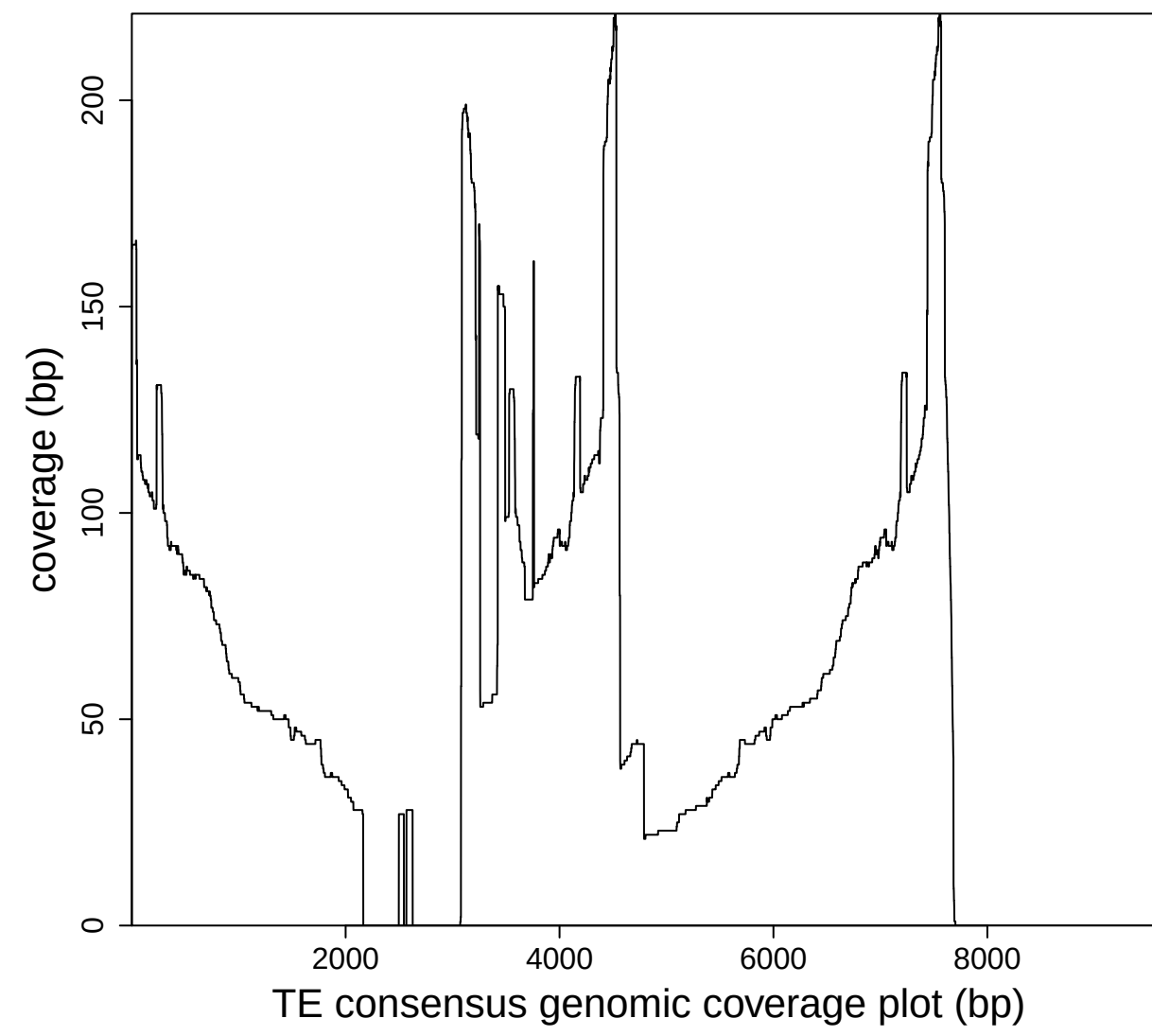
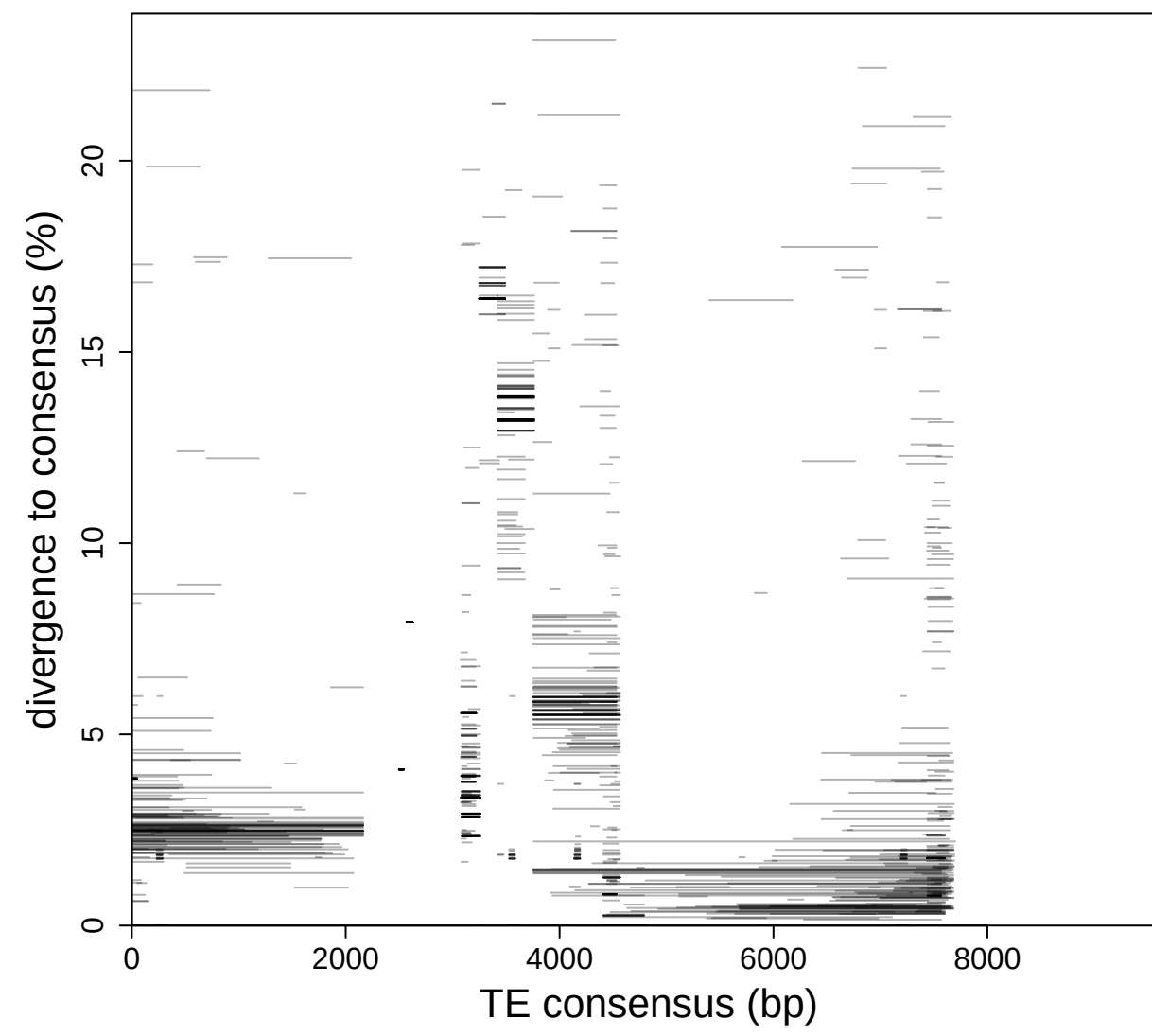
MSA length = 9669





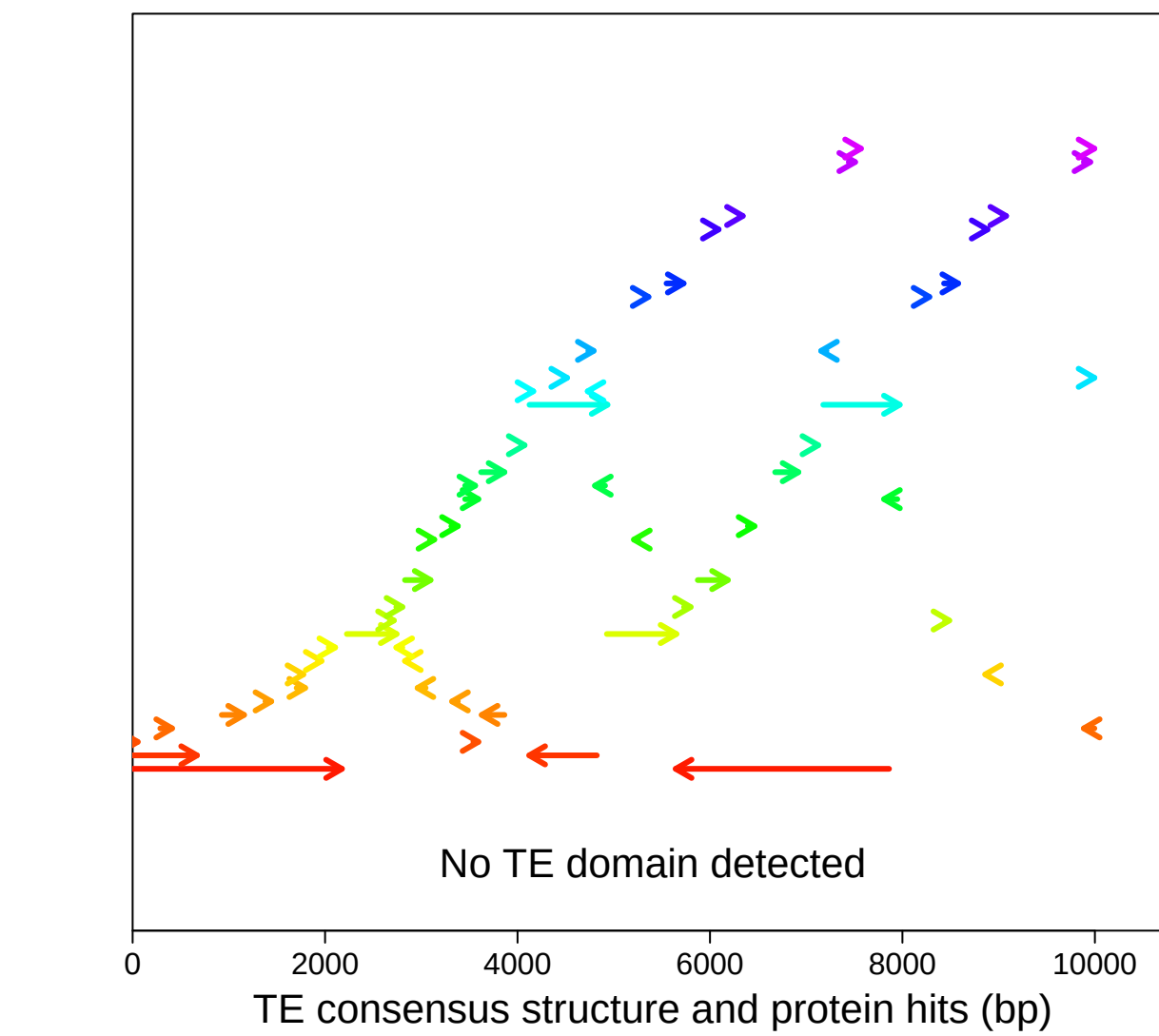
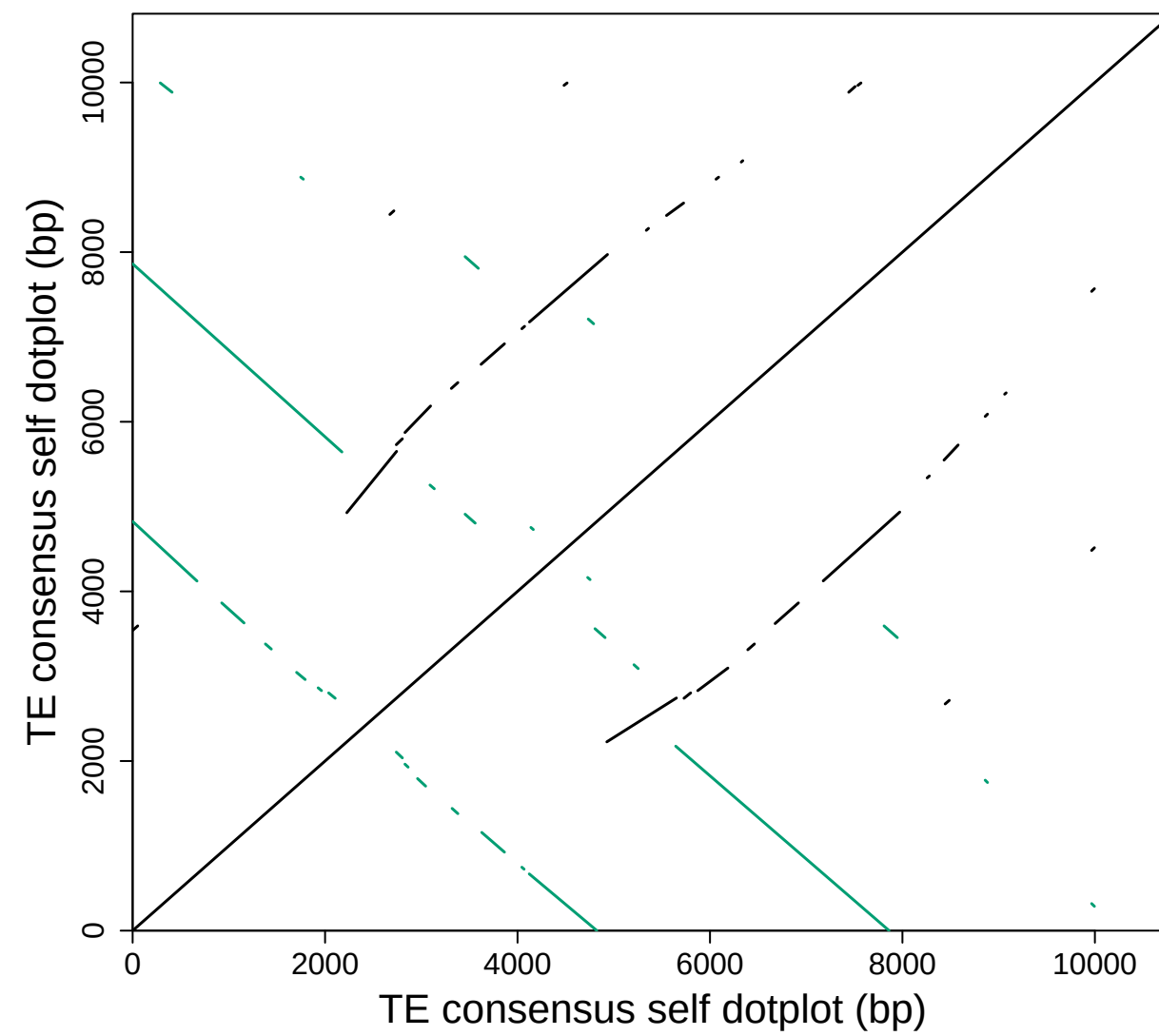
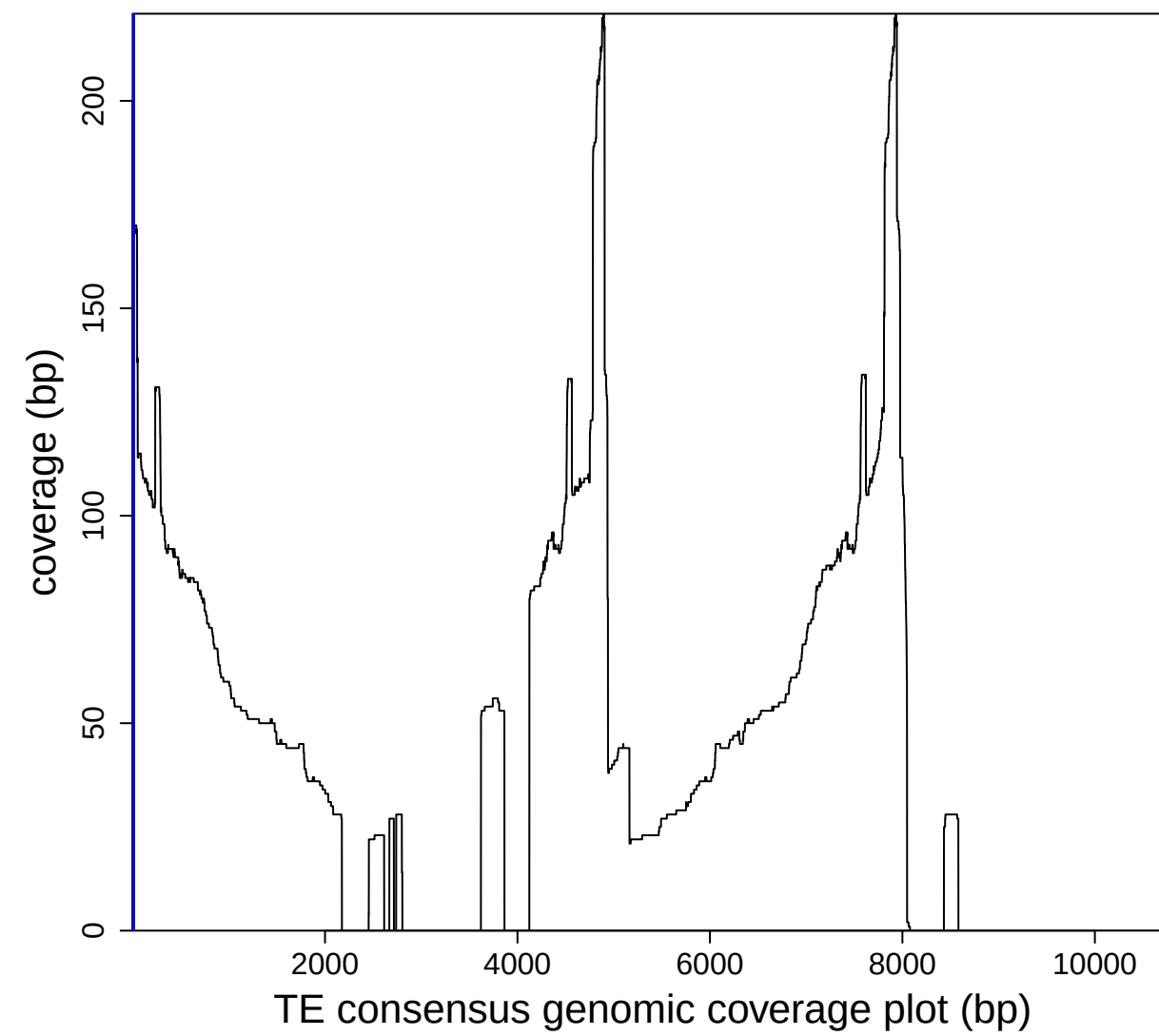
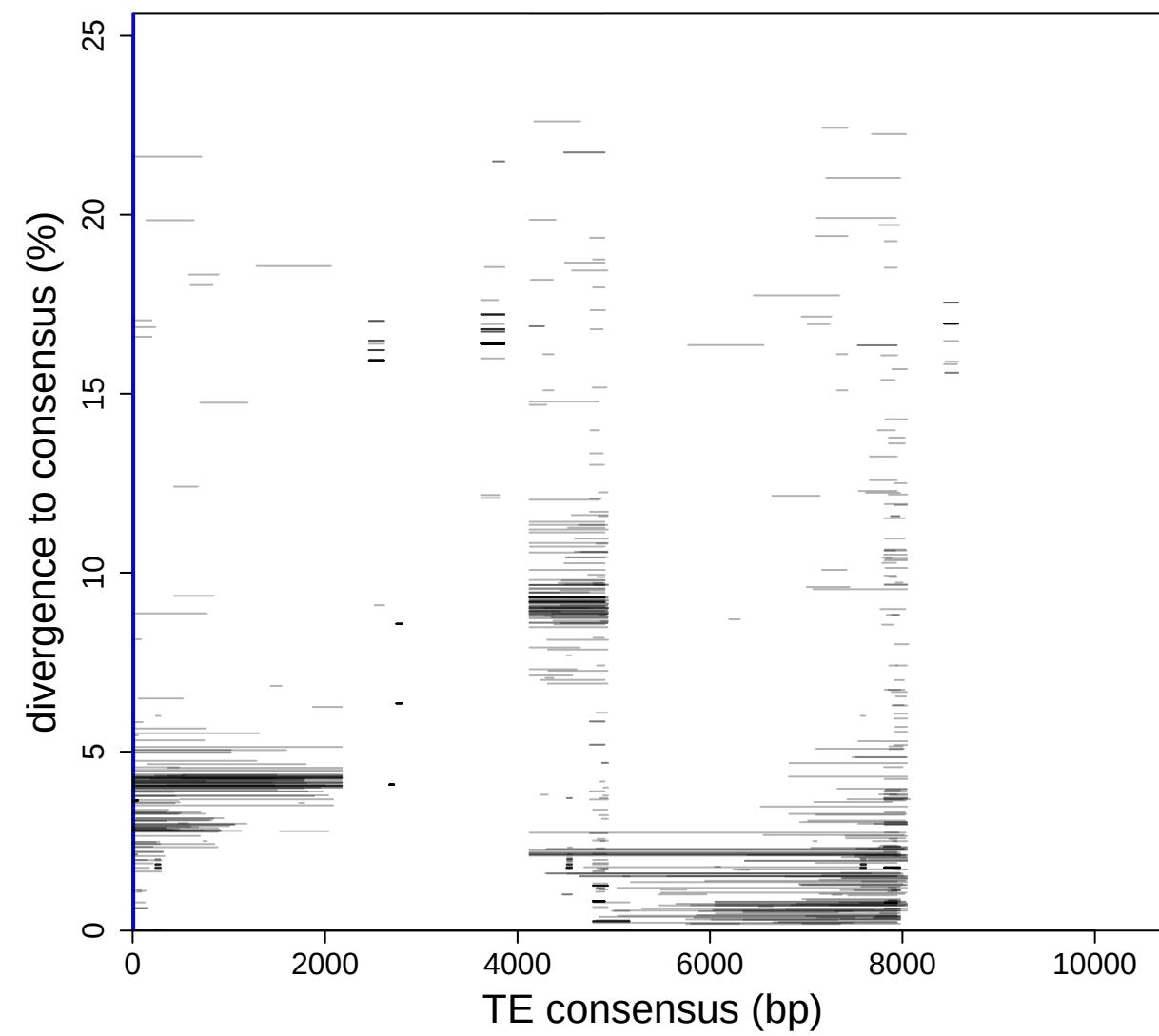
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 9669bp; fragments: 1230; full length: 0 (>=8702.1bp)

After TEtrimmer 9669 bp

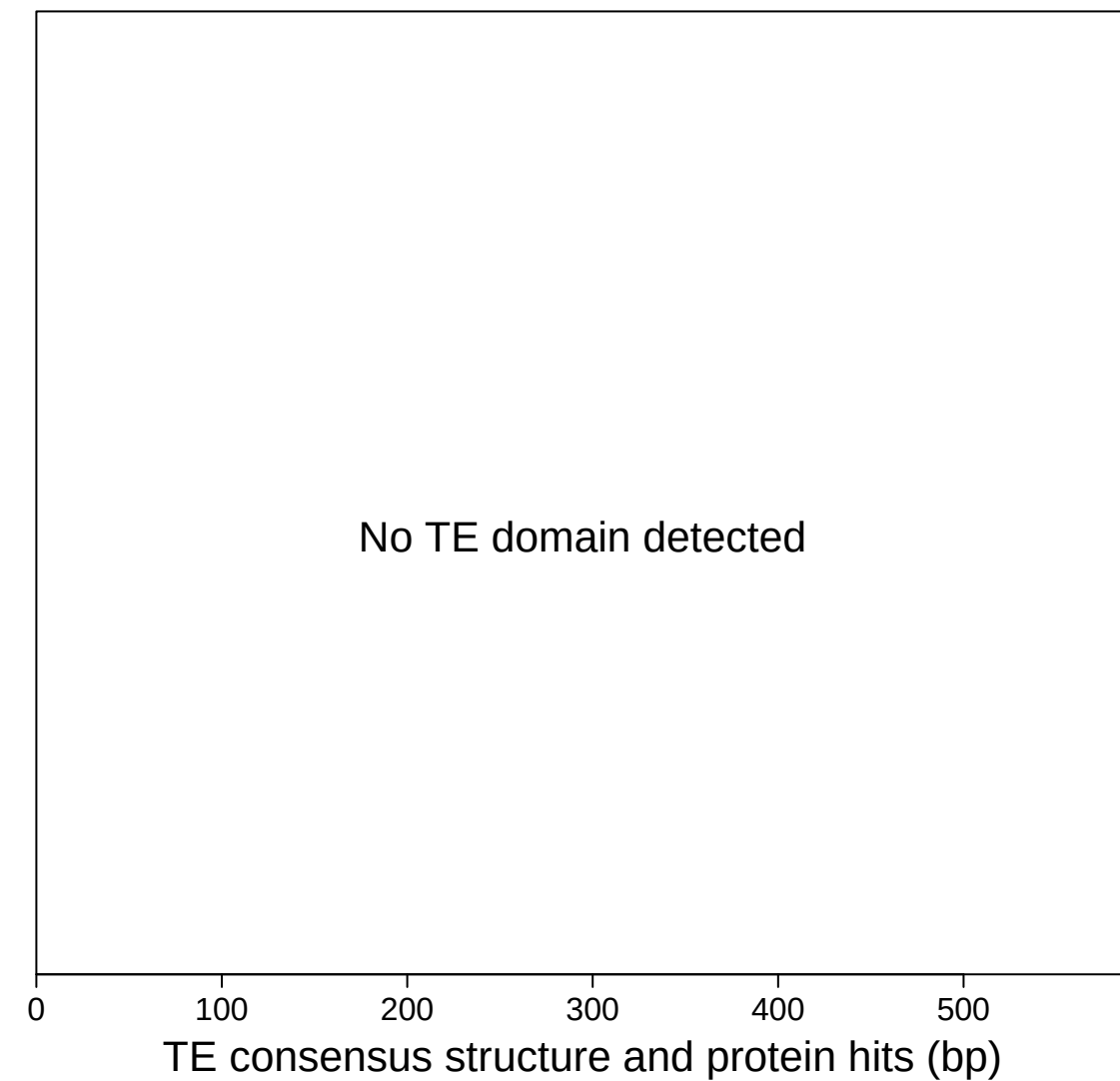
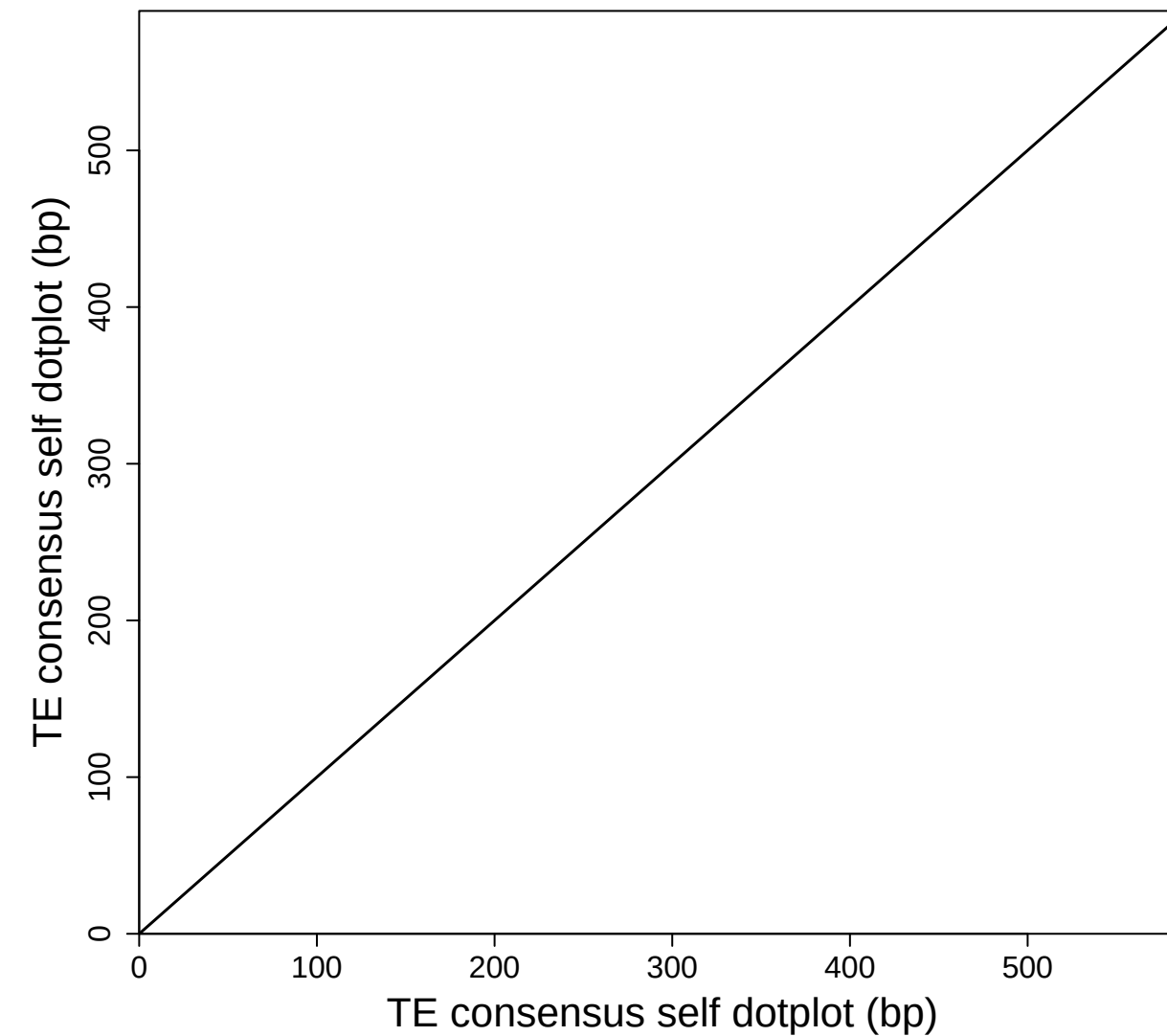
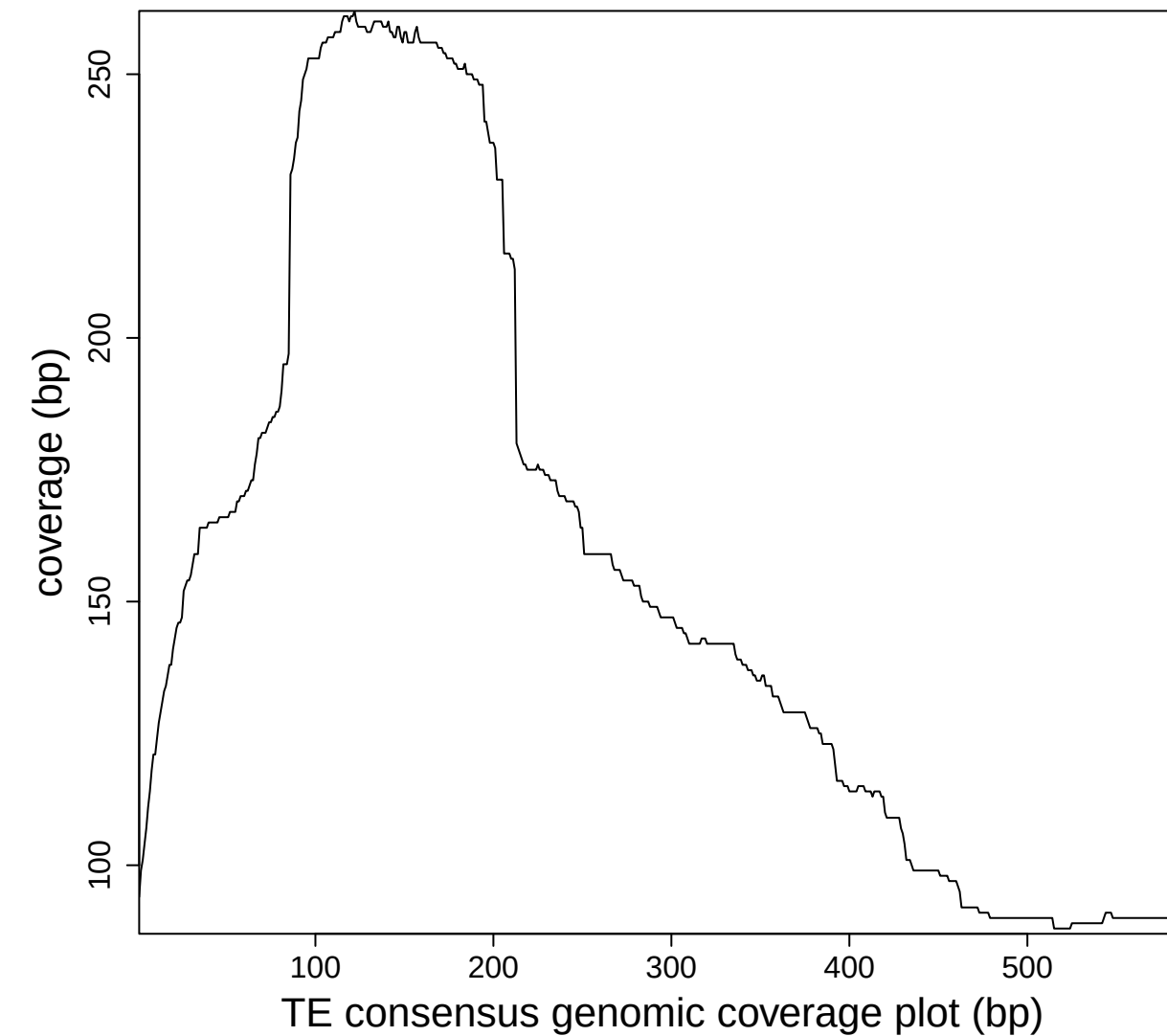
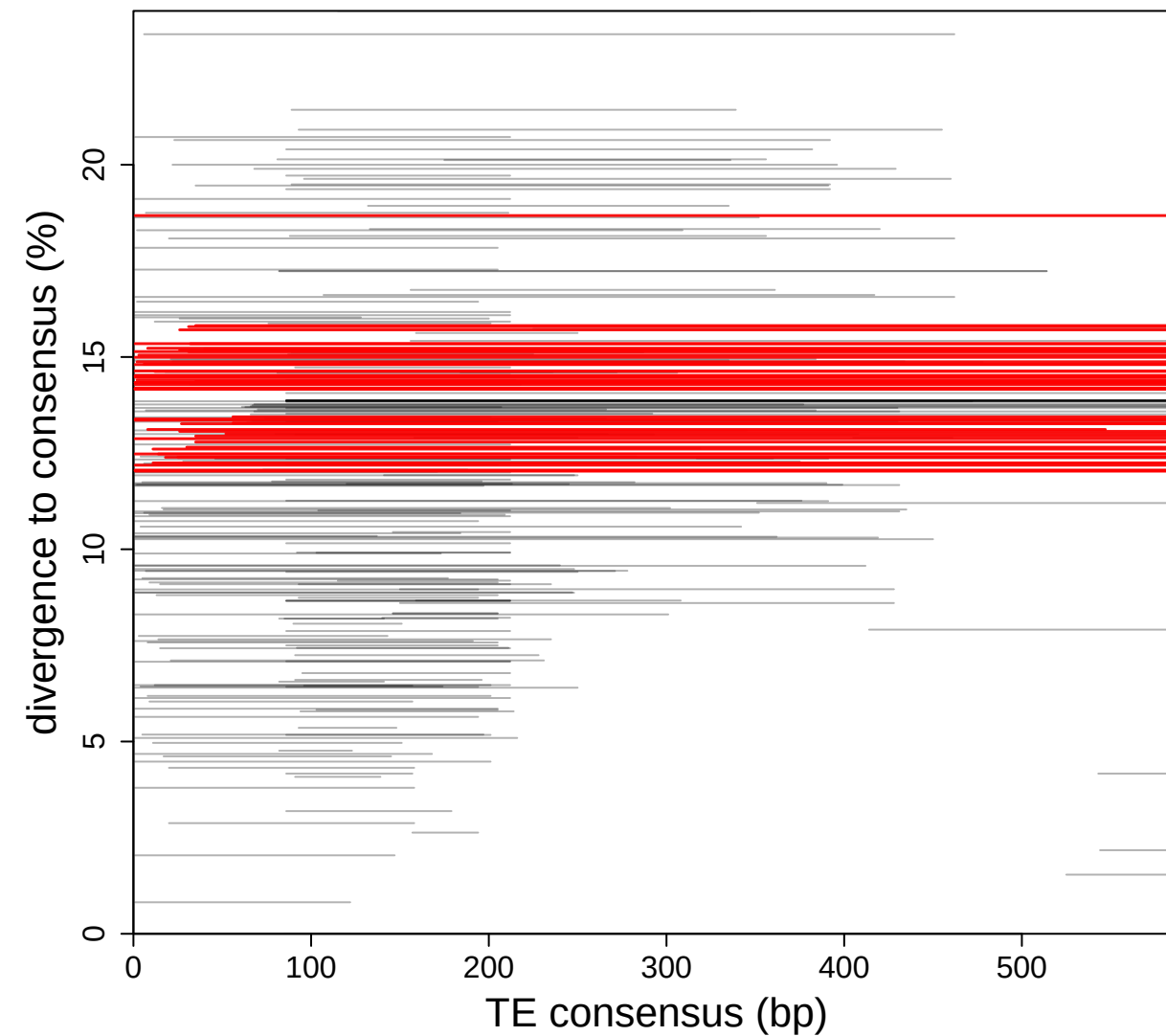


ed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.f
size: 10811bp; fragments: 946; full length: 0 (>=9729.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

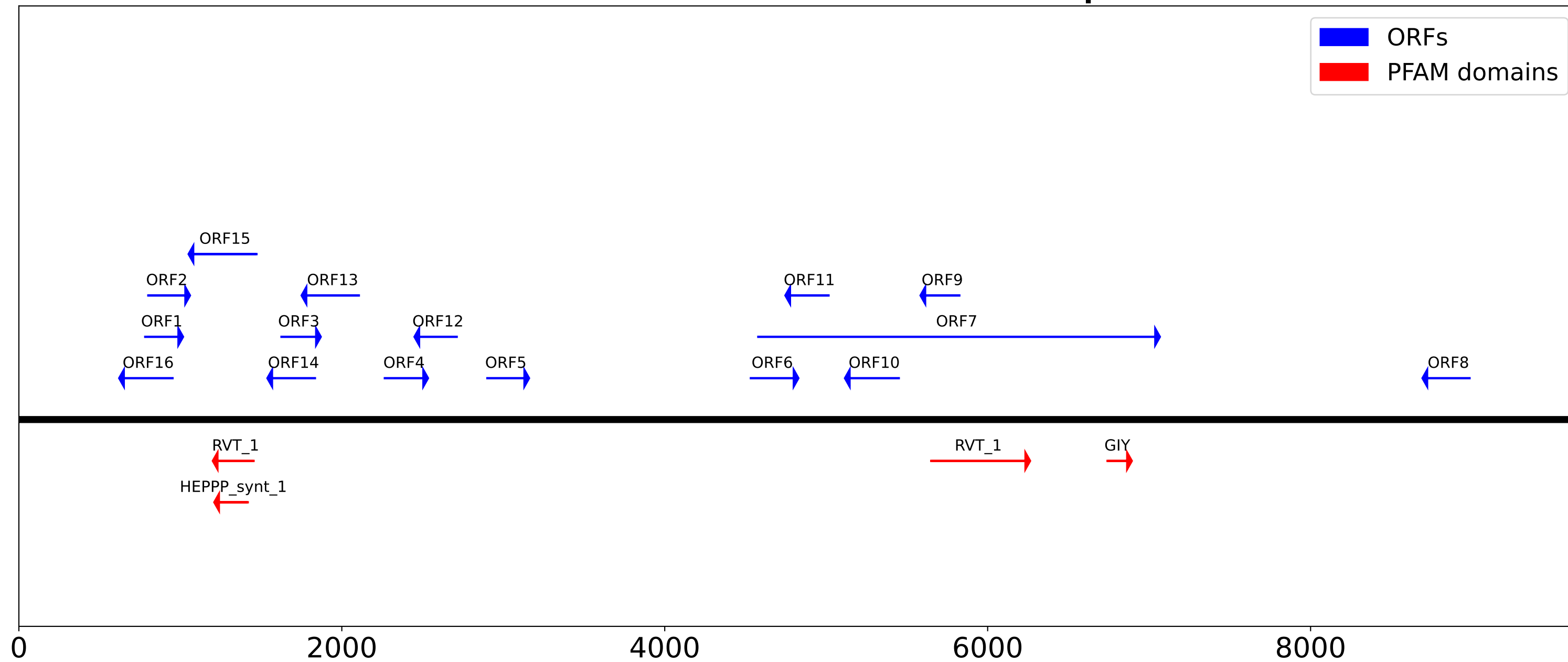


TE: rnd_5_family_762
size: 589bp; fragments: 289; full length: 60 (≥ 530.1 bp)

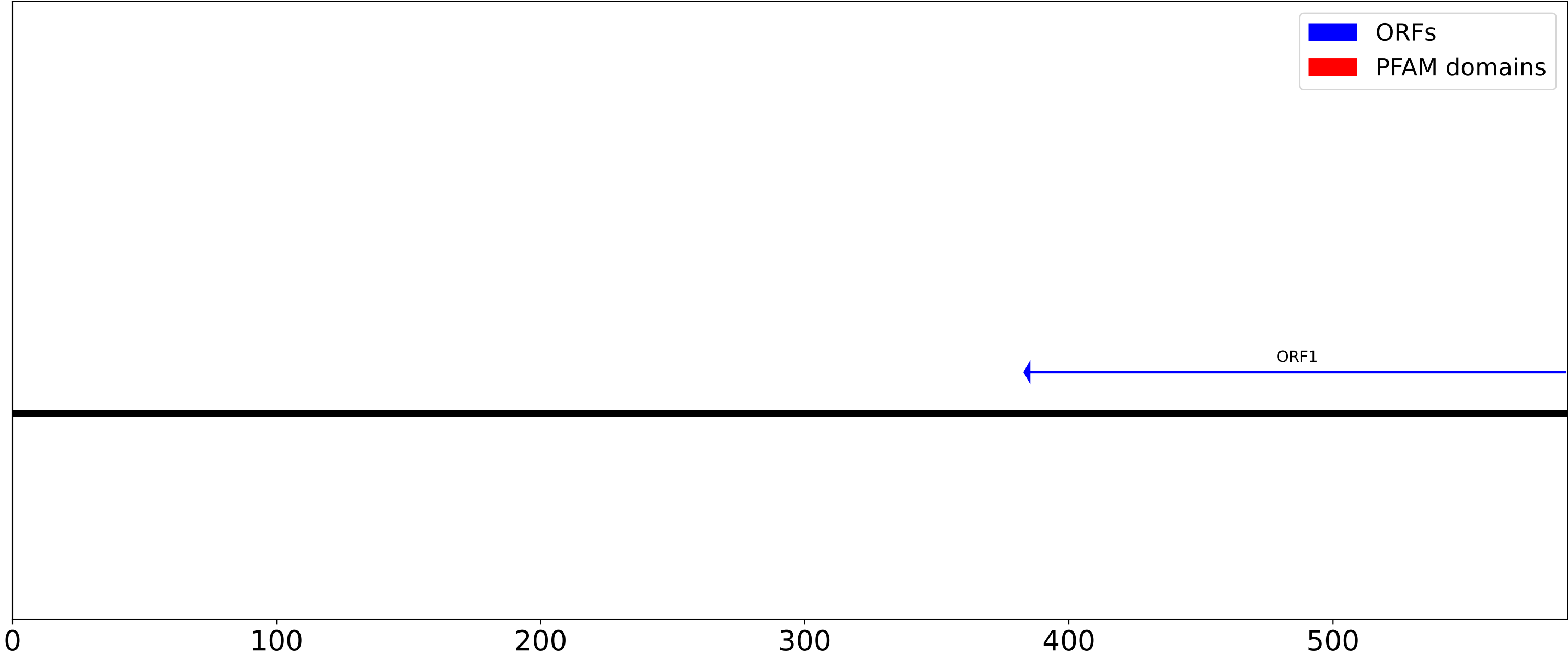


Before TEtrimmer 589 bp

After TEtrimmer ORF and PFAM domain plot



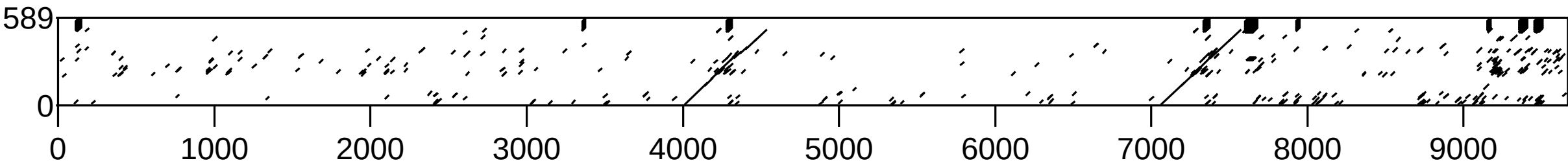
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)