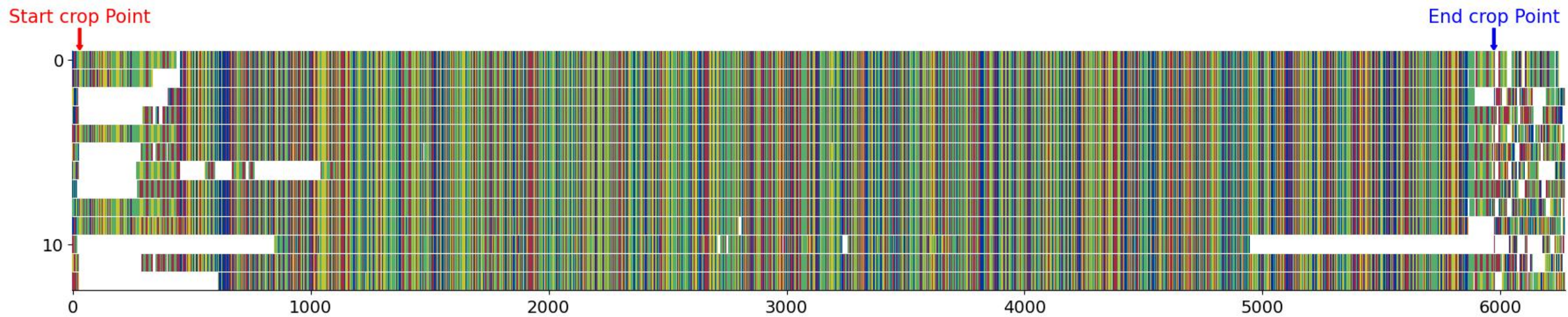




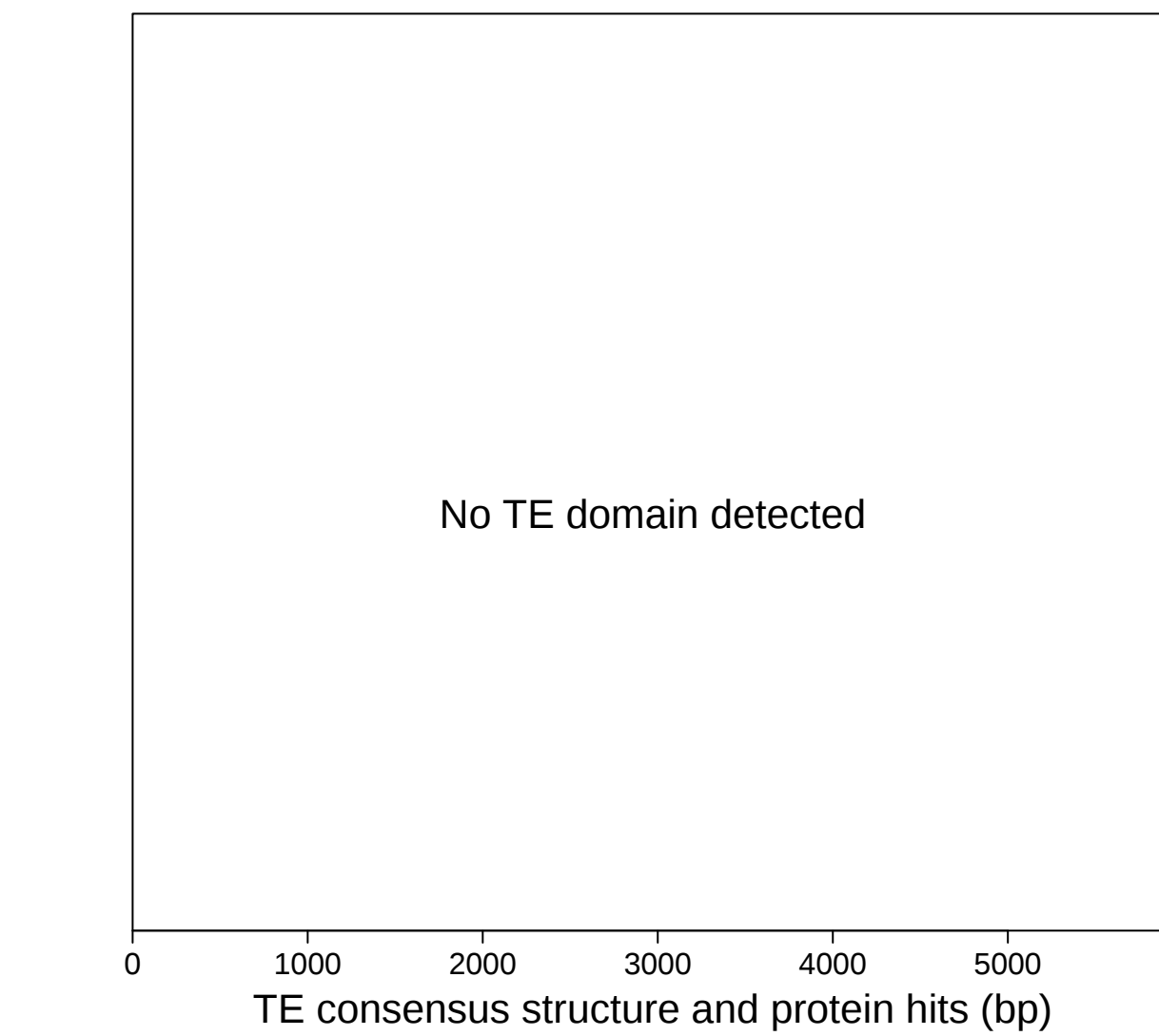
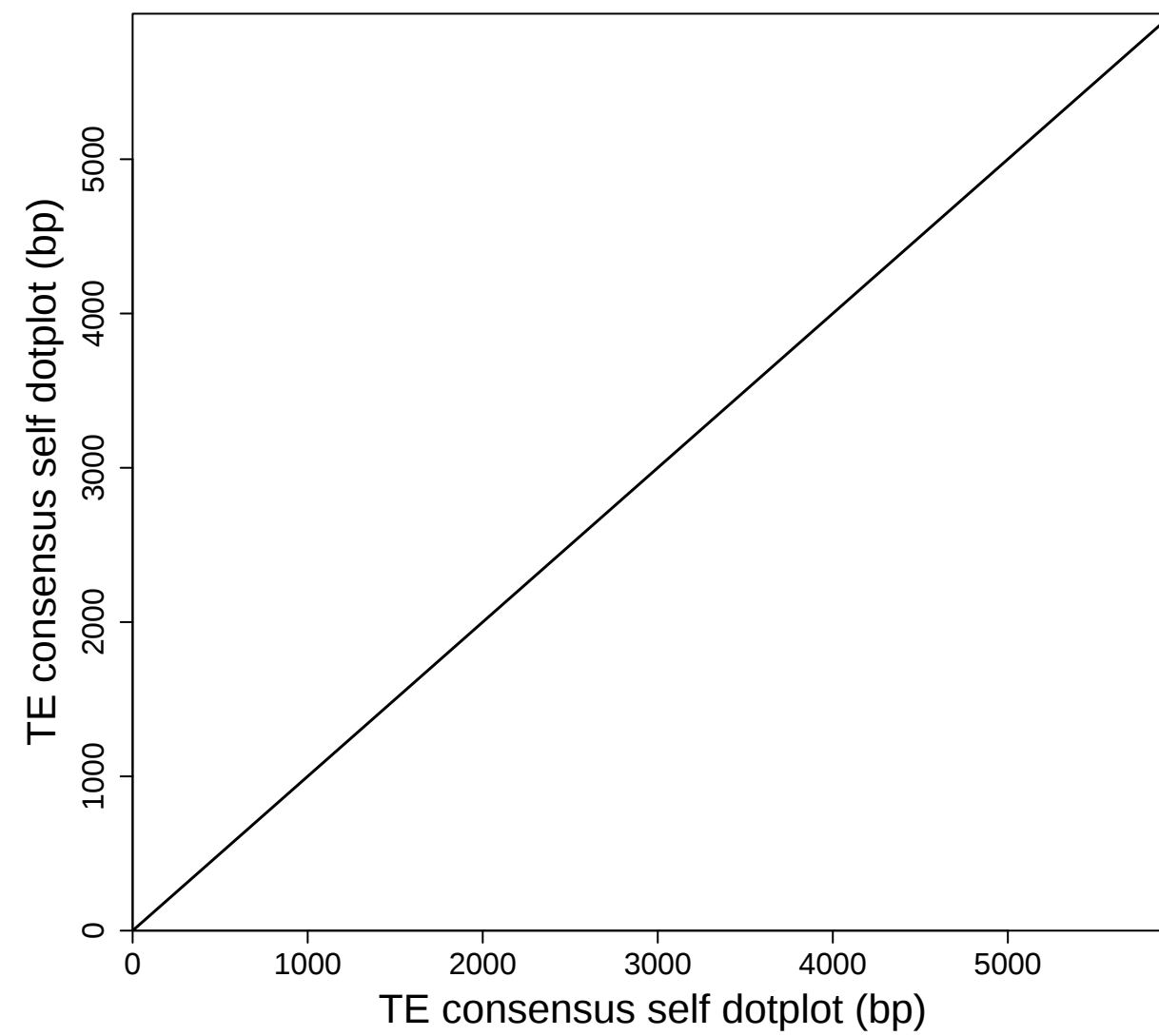
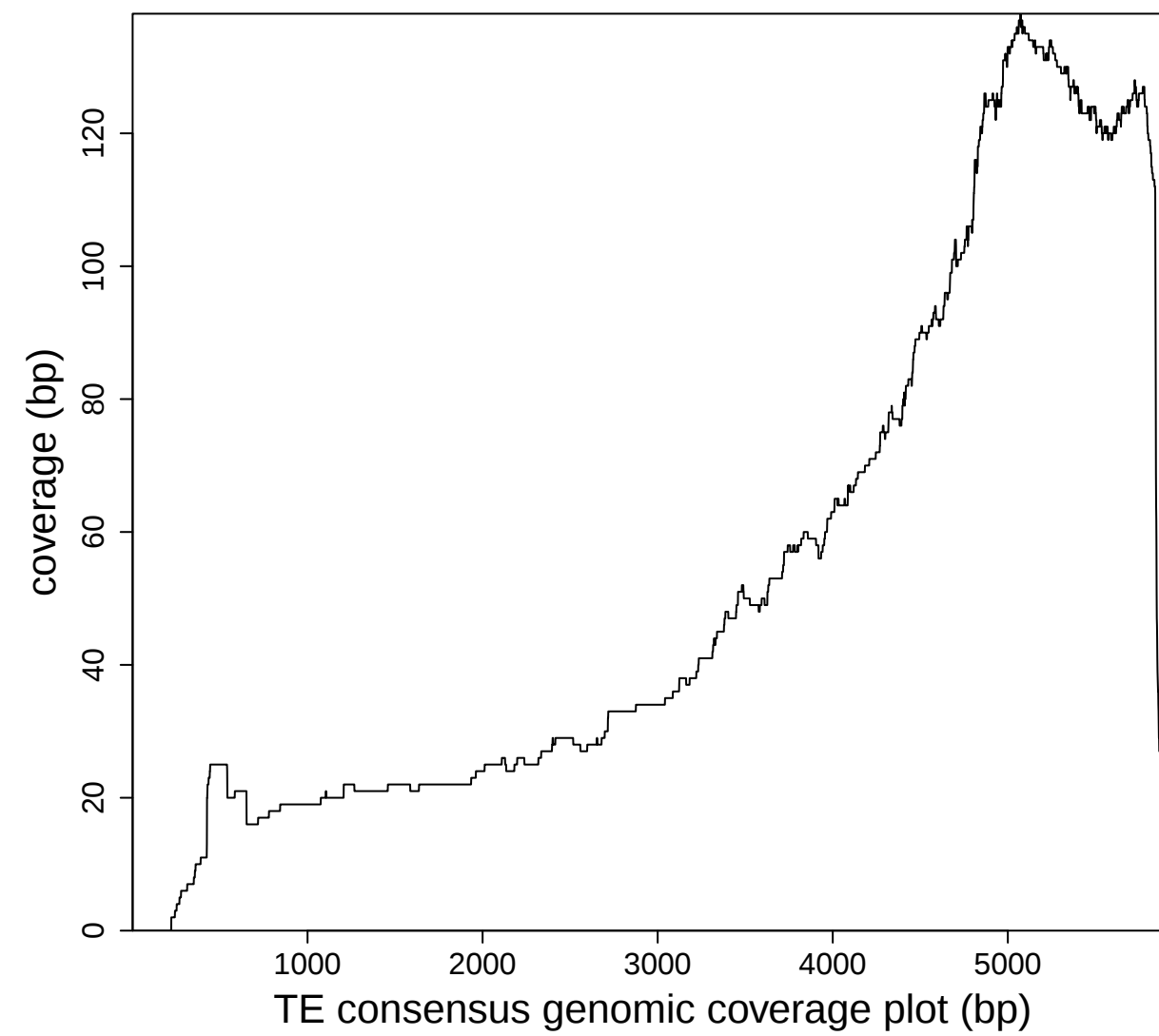
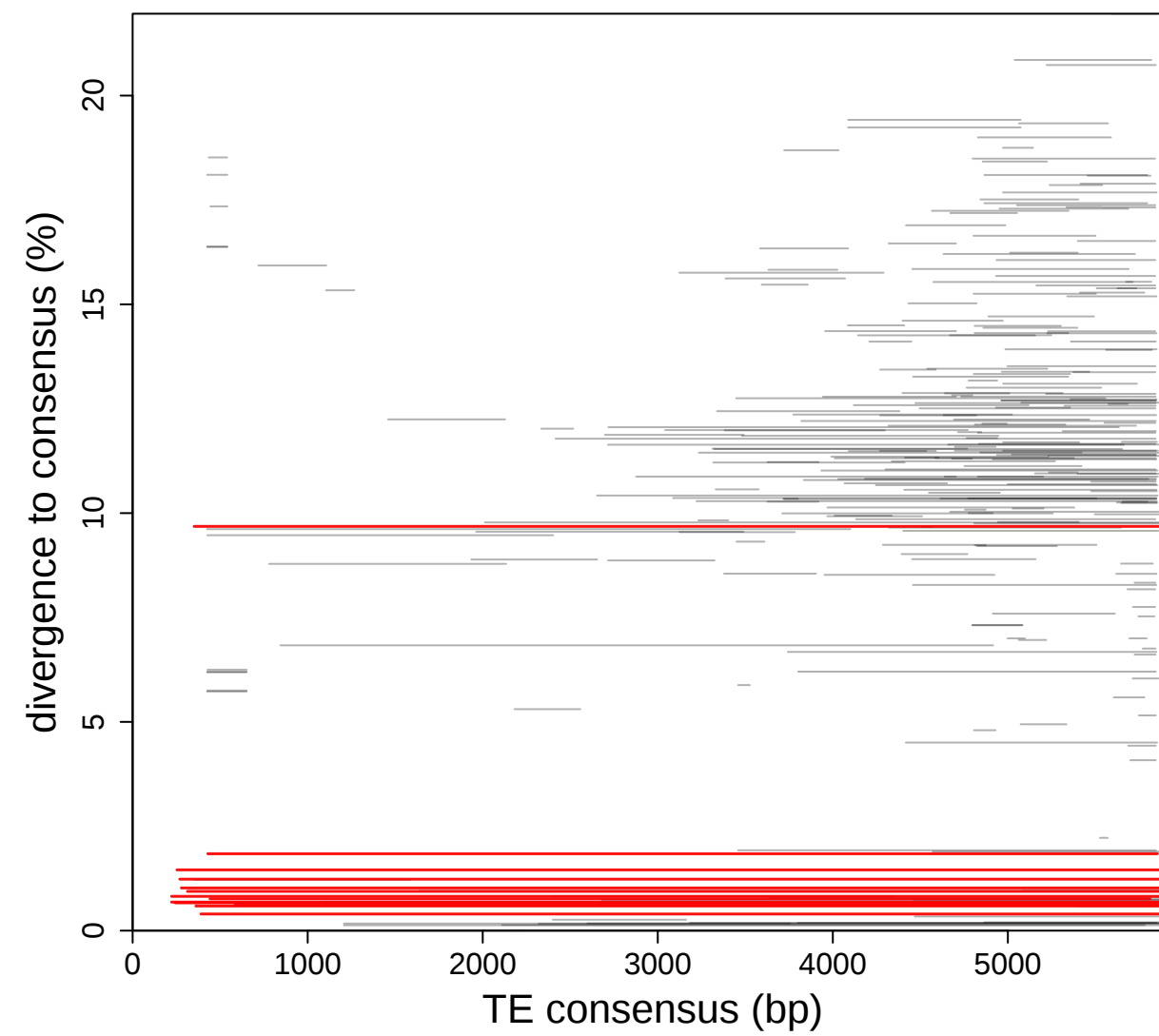
MSA length = 5943

[illegible]



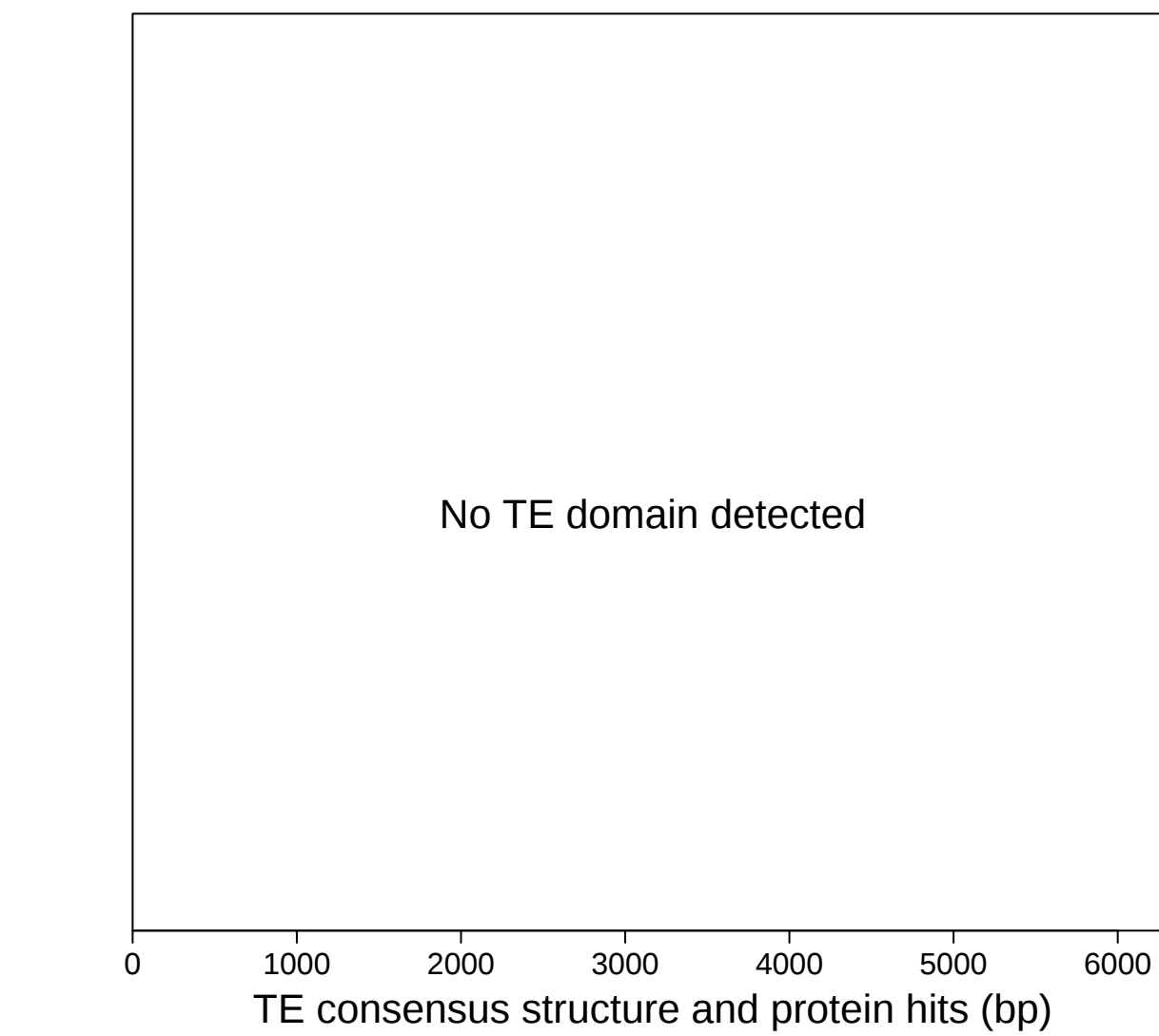
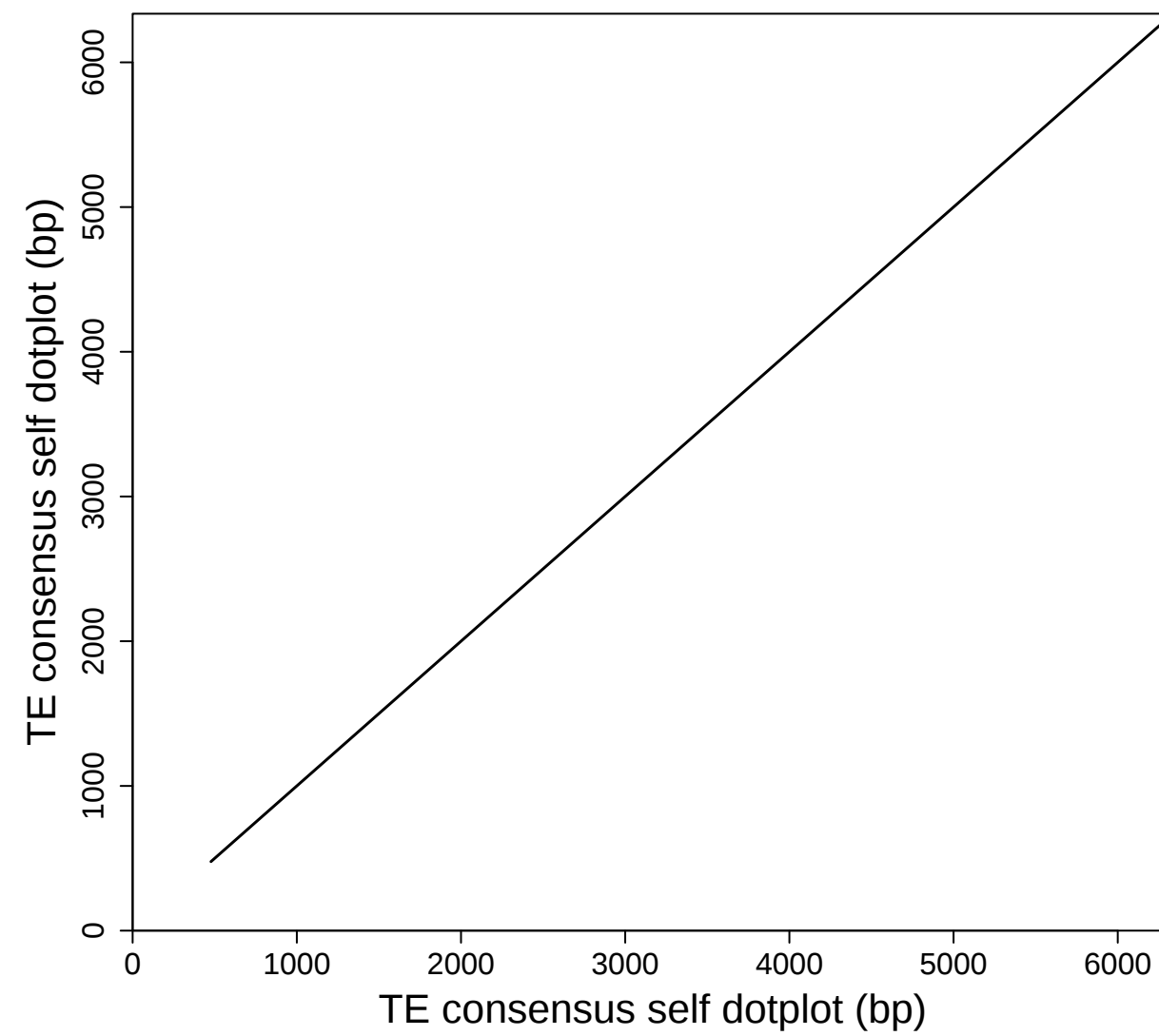
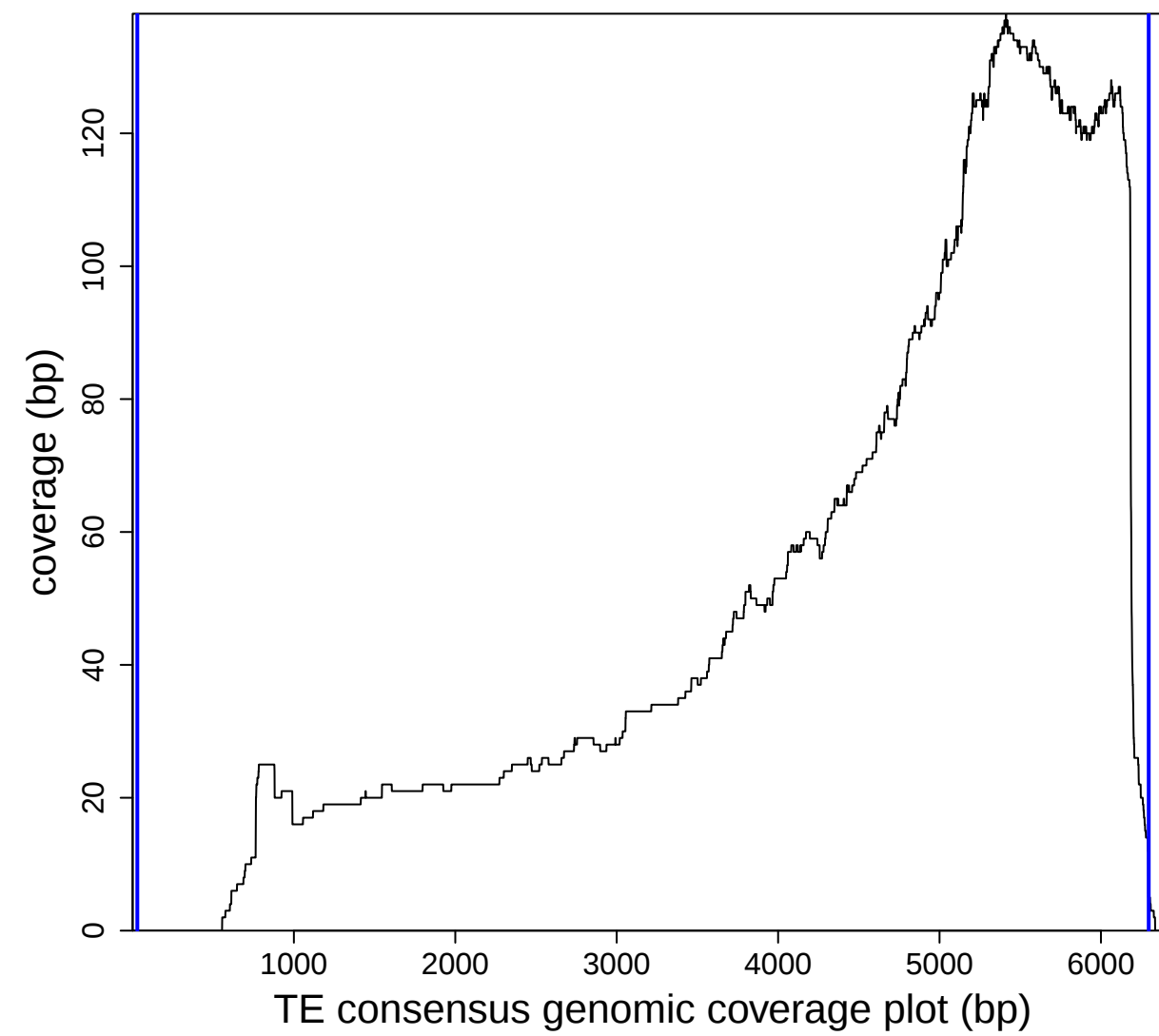
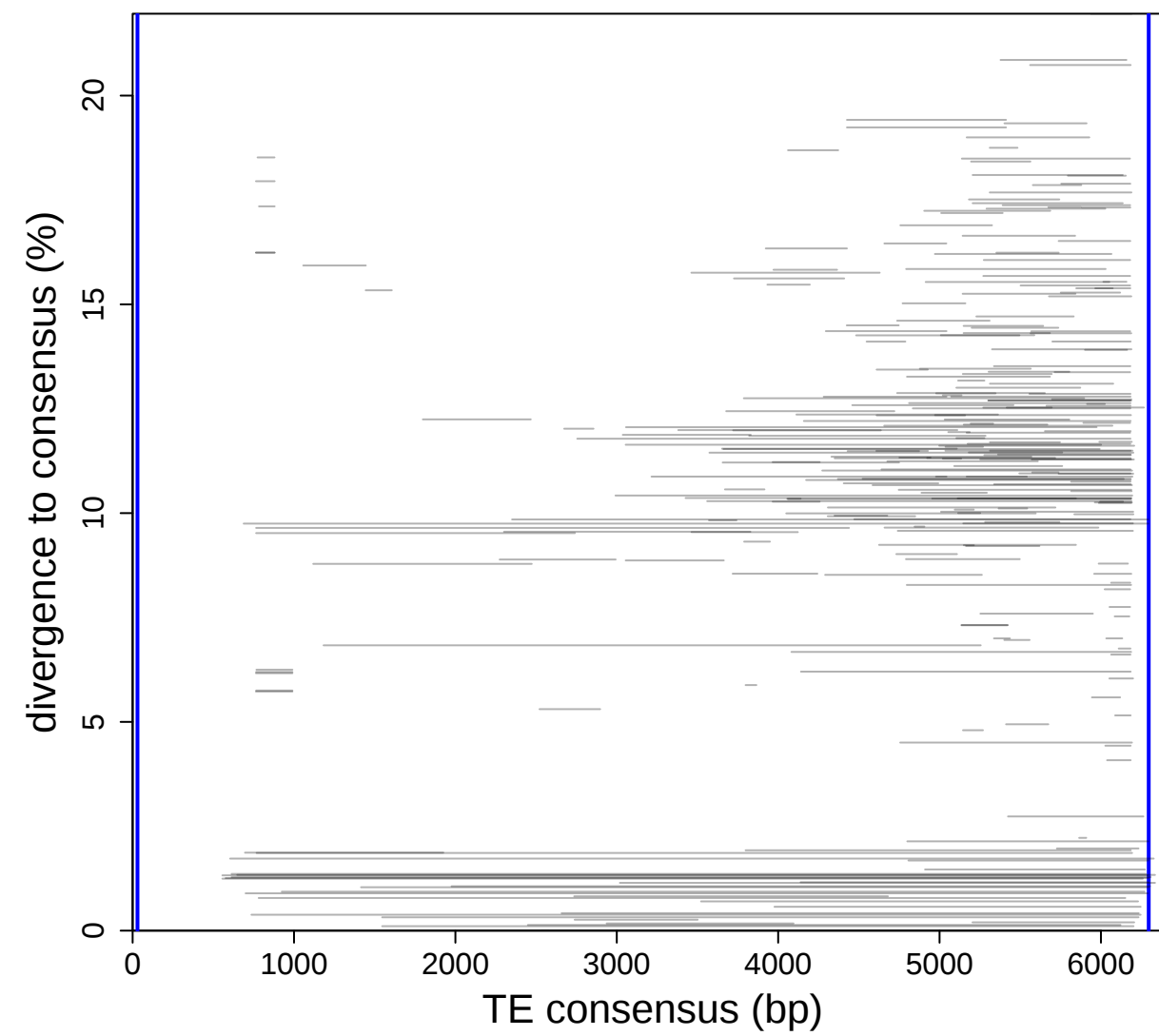
3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.f
size: 5943bp; fragments: 276; full length: 13 (>=5348.7bp)

After TEtrimmer 5943 bp

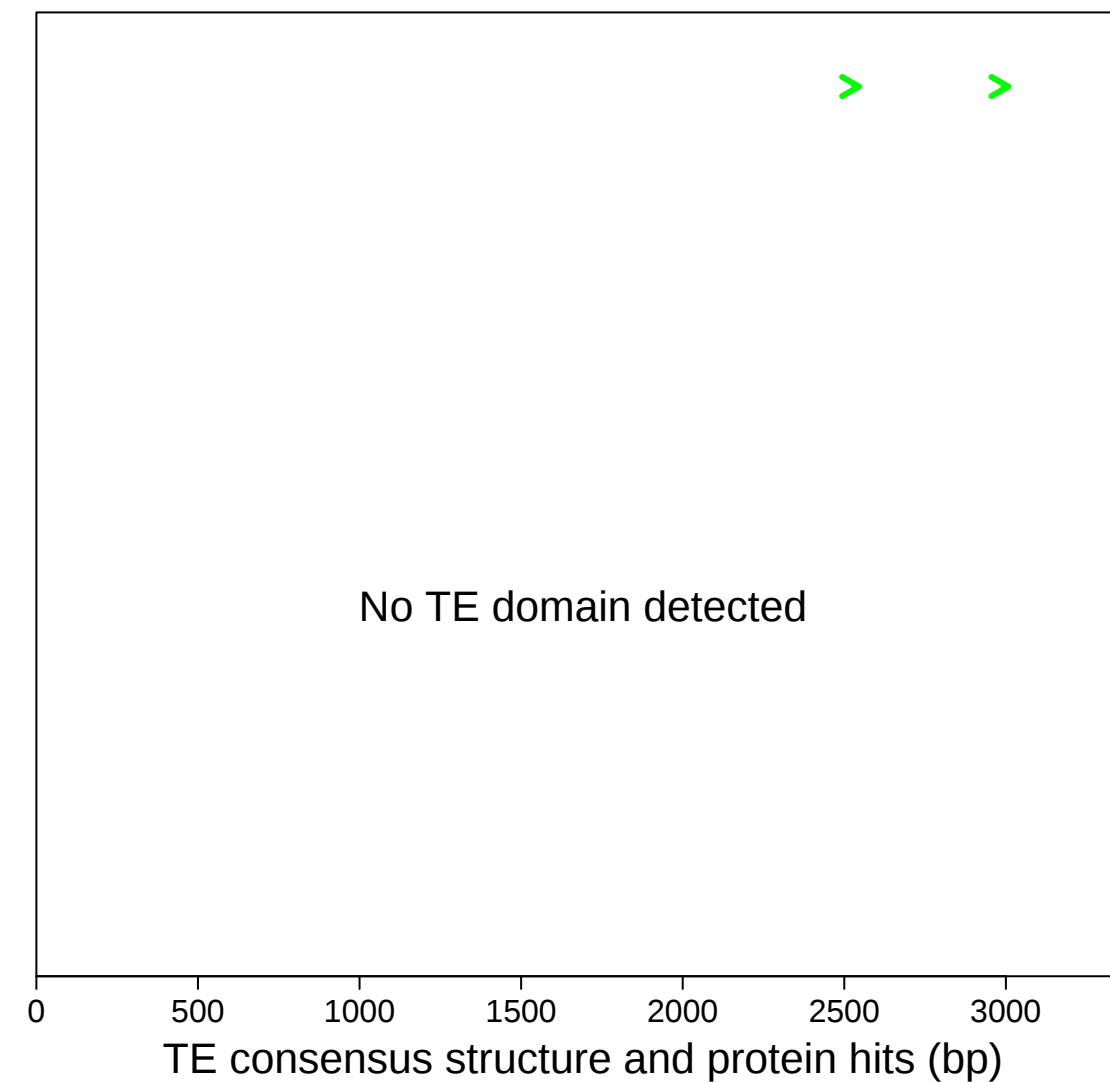
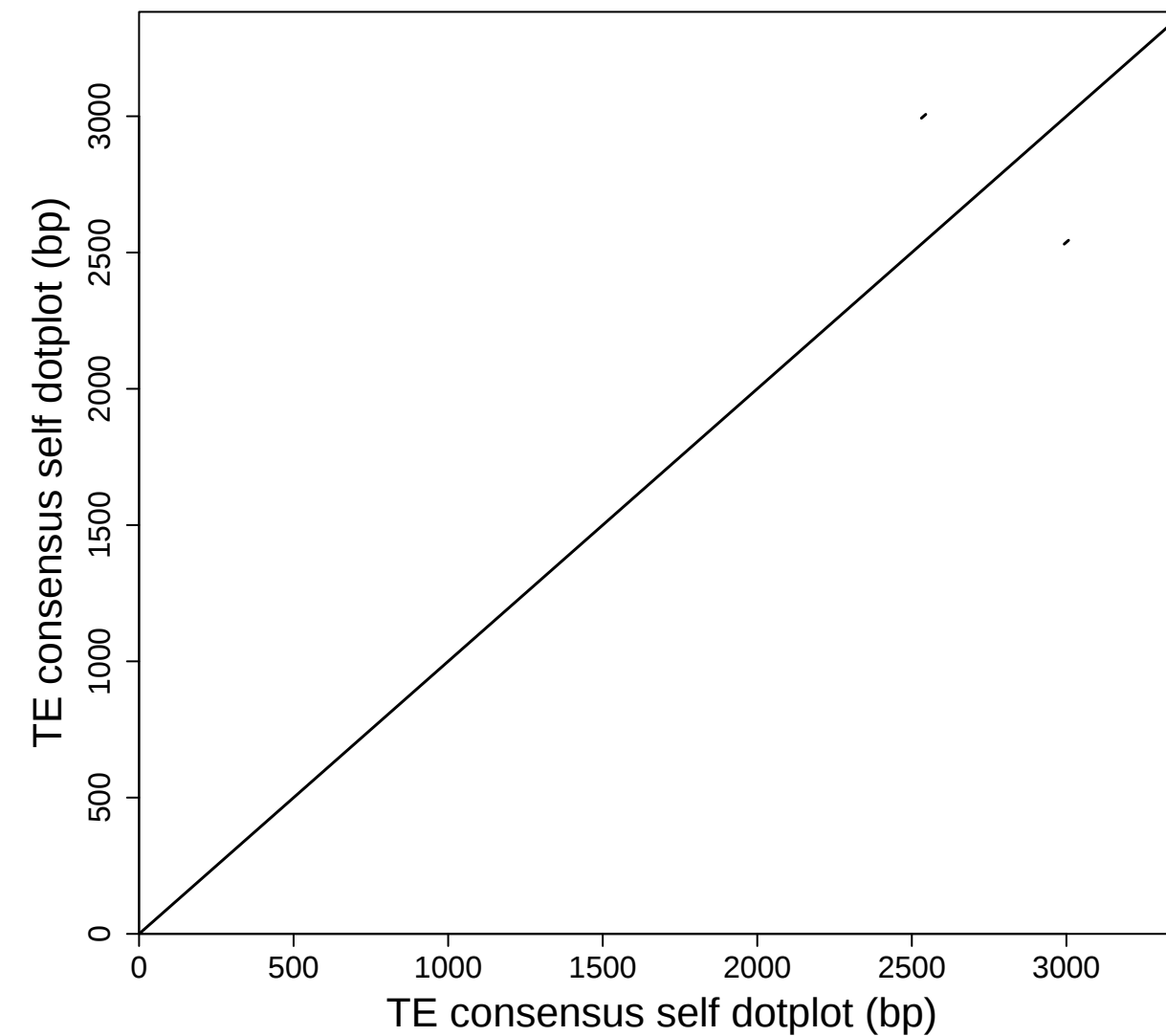
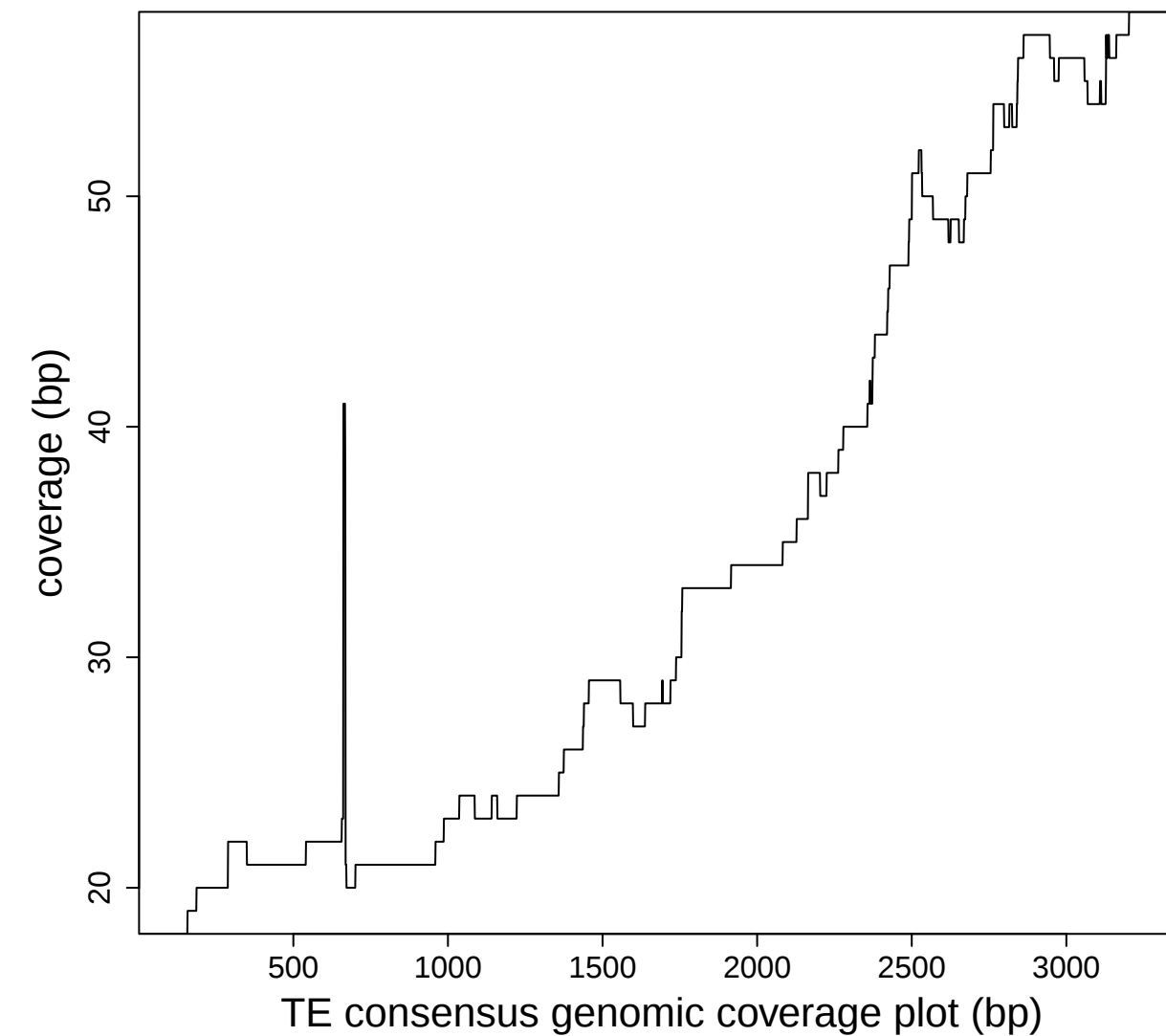
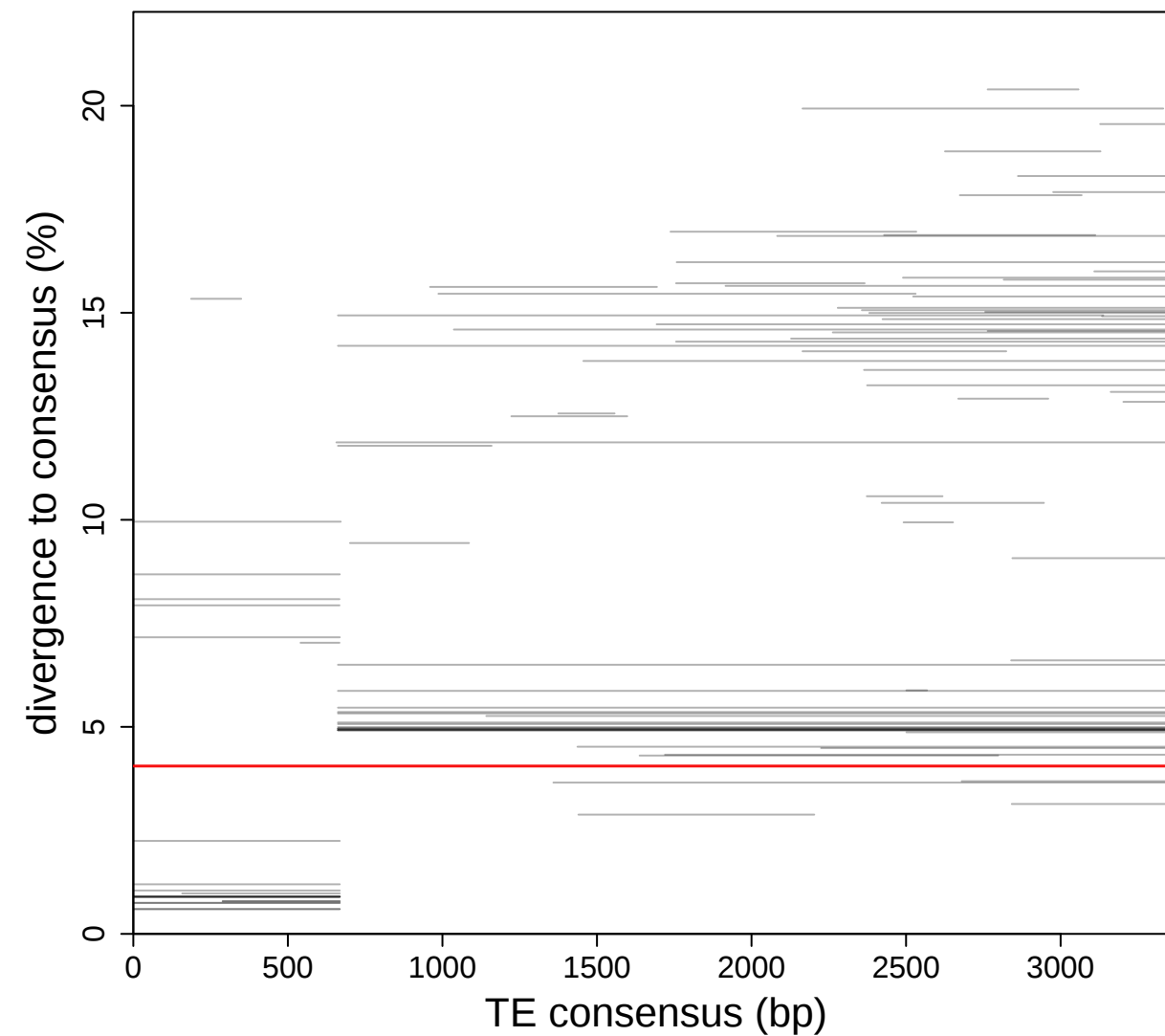


f.bed_g_3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 6446bp; fragments: 276; full length: 0 (>=5801.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

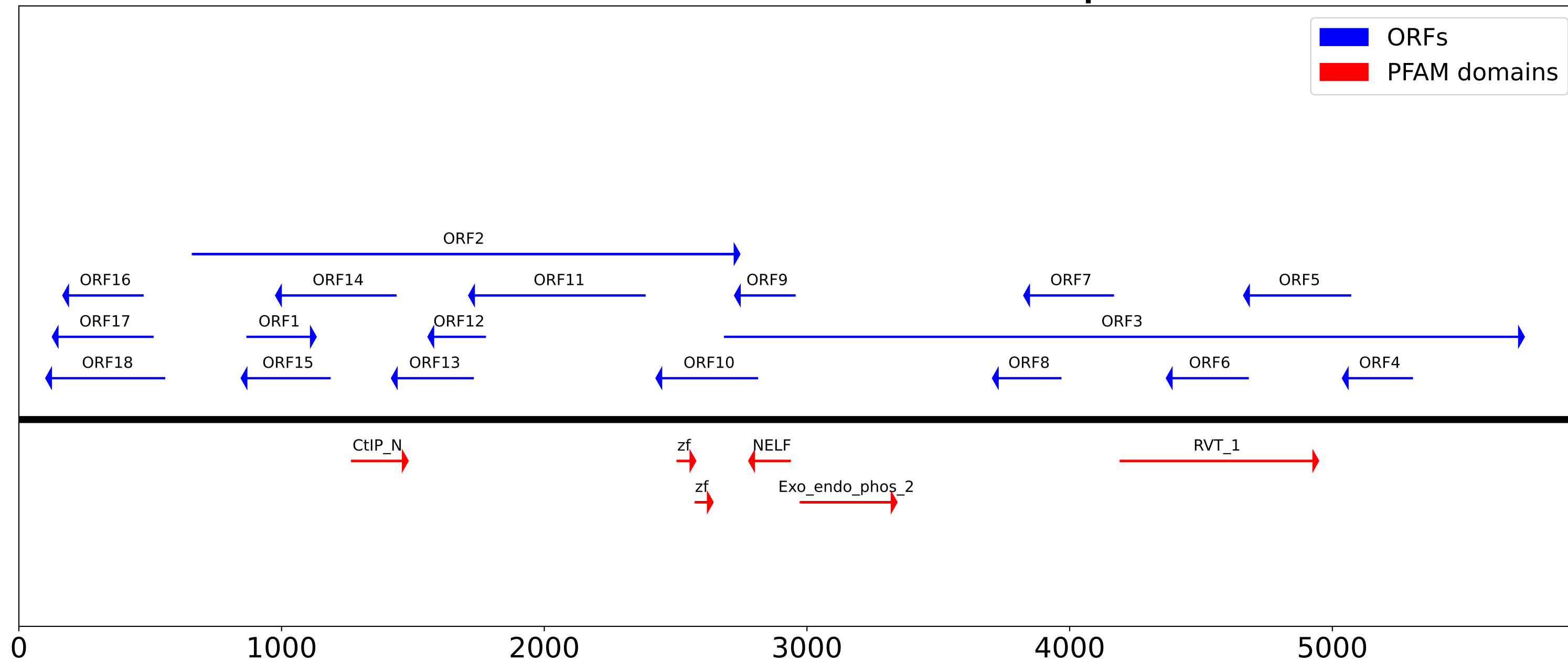


TE: rnd_4_family_1546
size: 3383bp; fragments: 101; full length: 1 (>=3044.7bp)

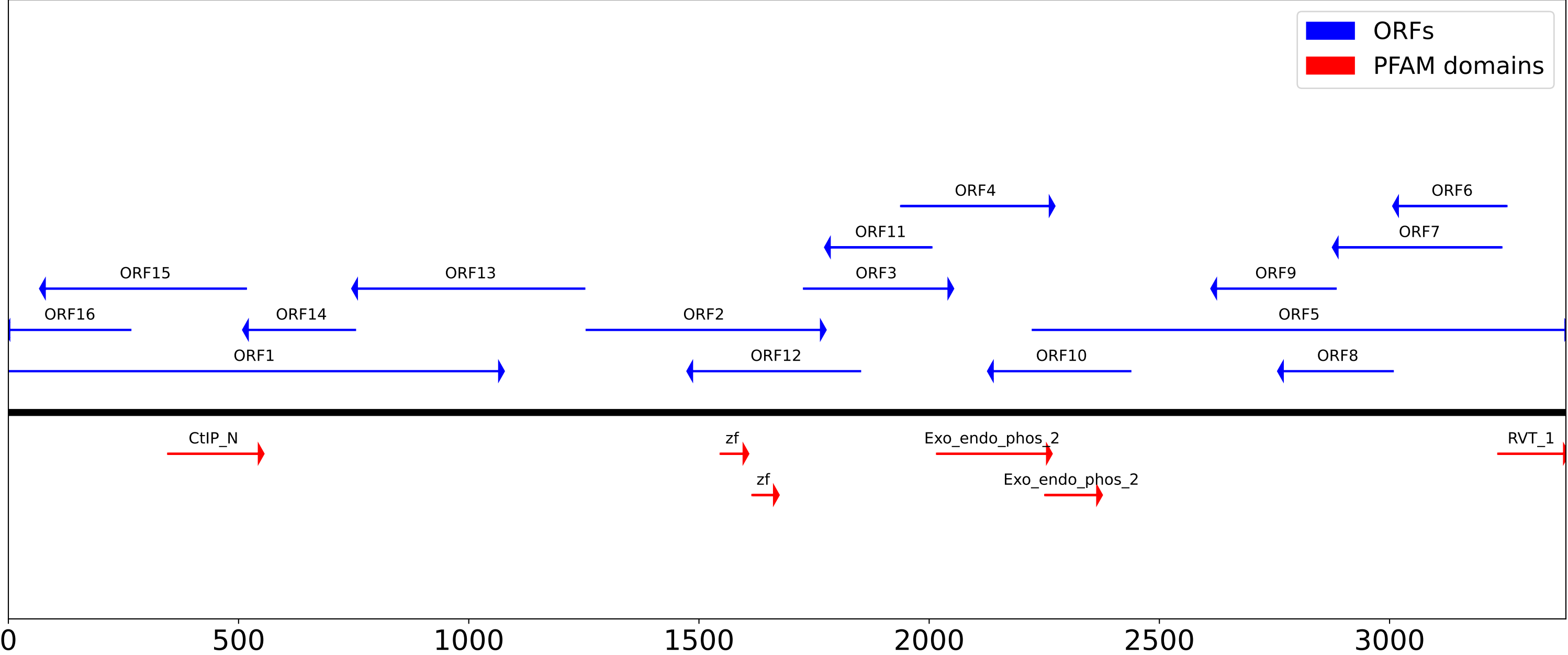


Before TEtrimmer 3383 bp

After TEtrimmer ORF and PFAM domain plot



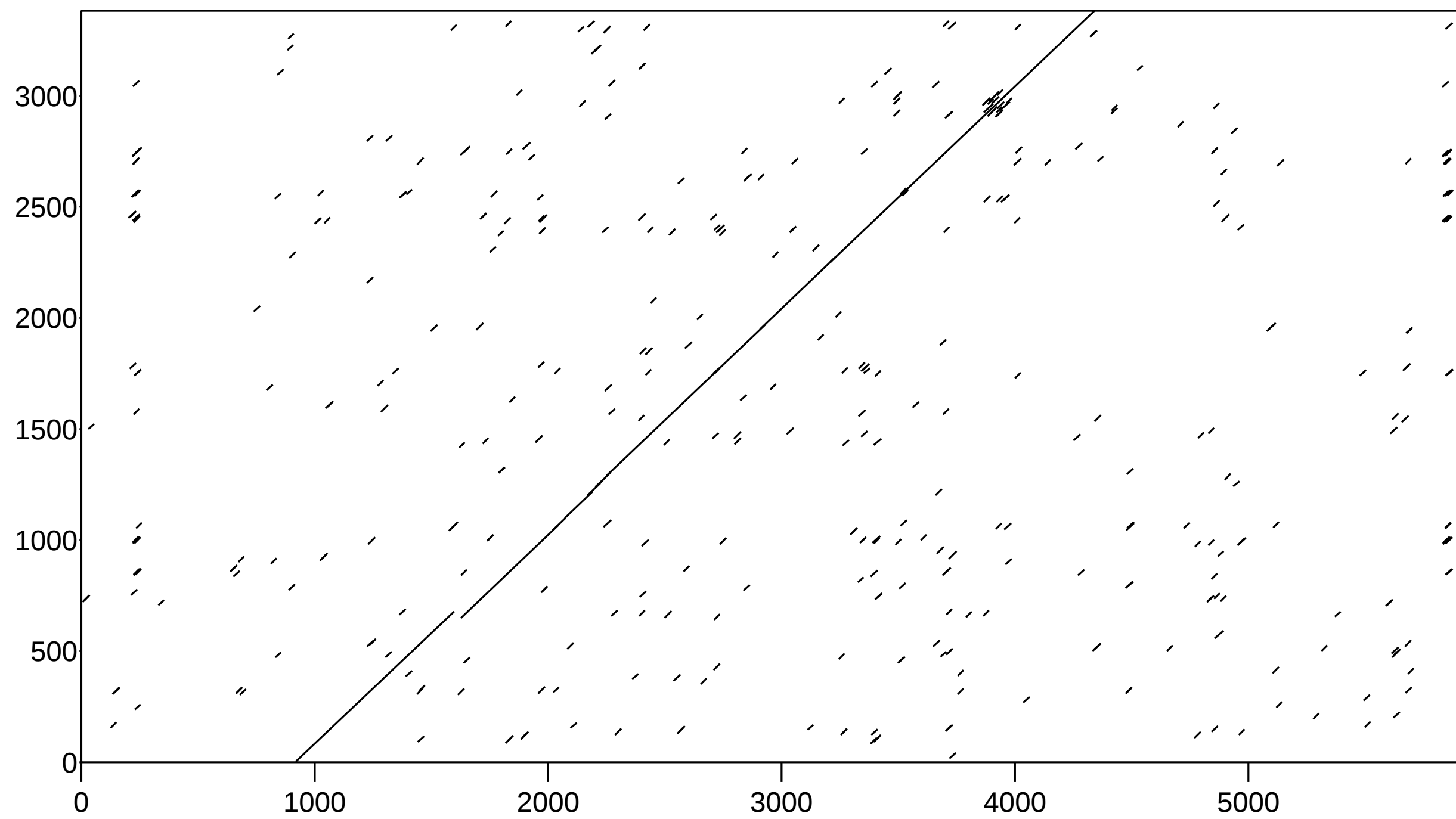
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)