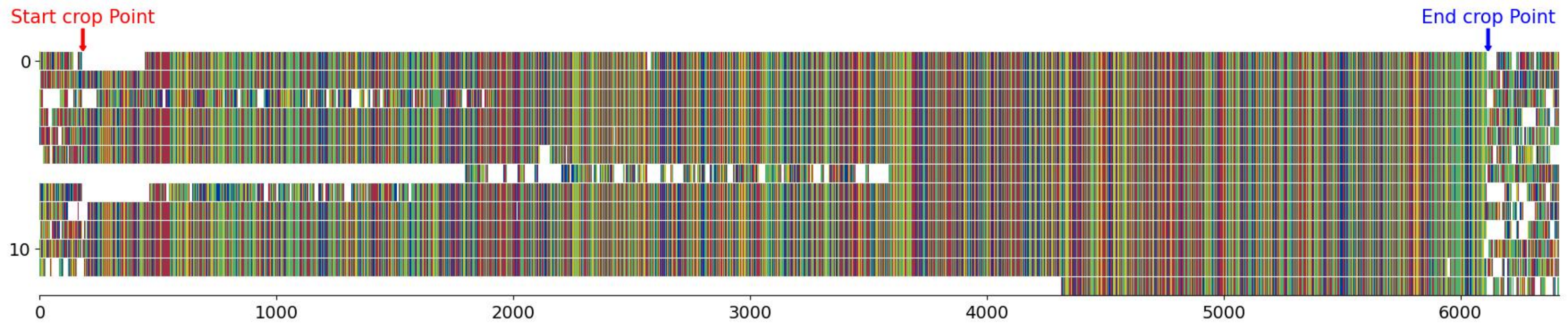


Start crop Point

End crop Point

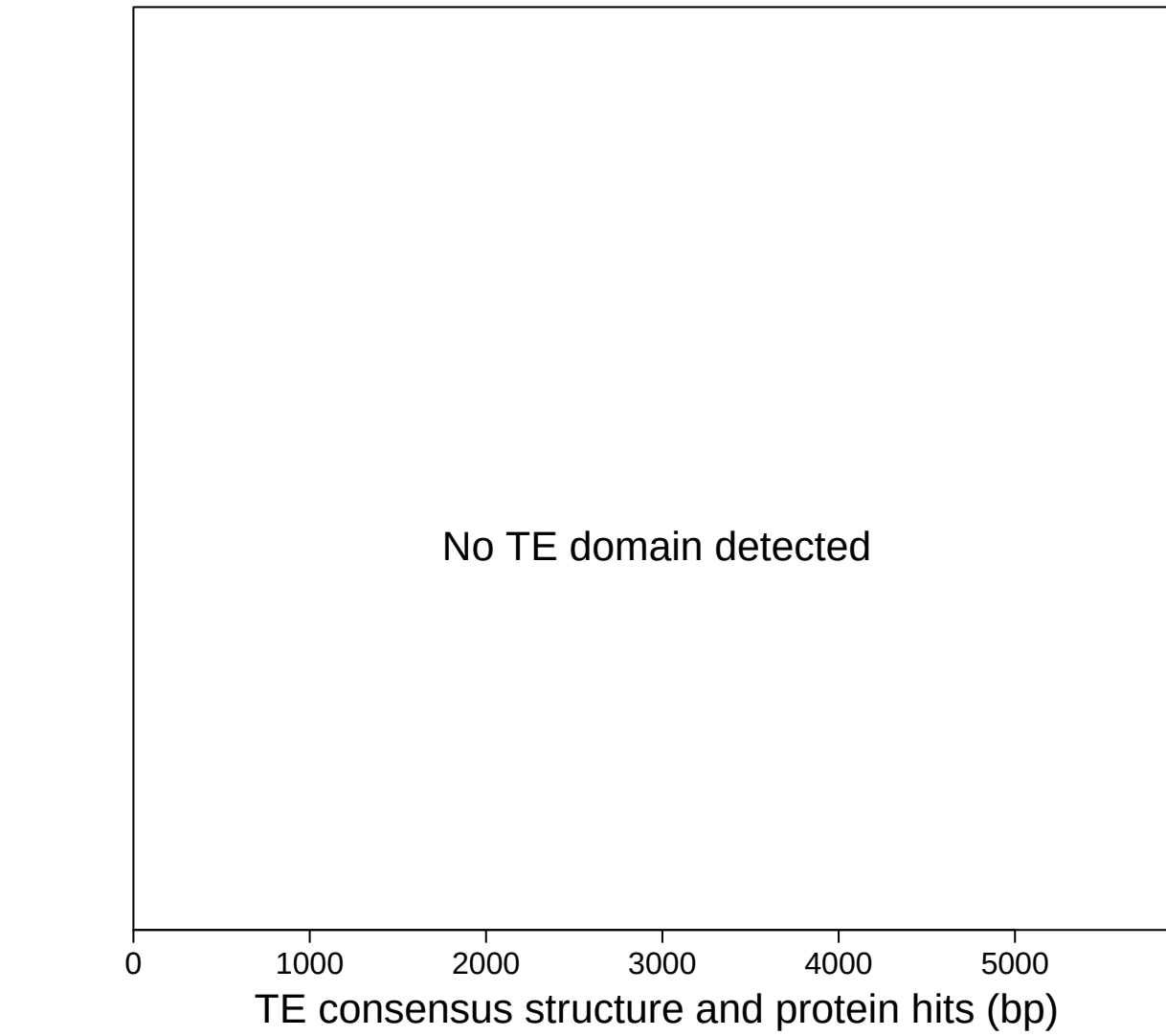
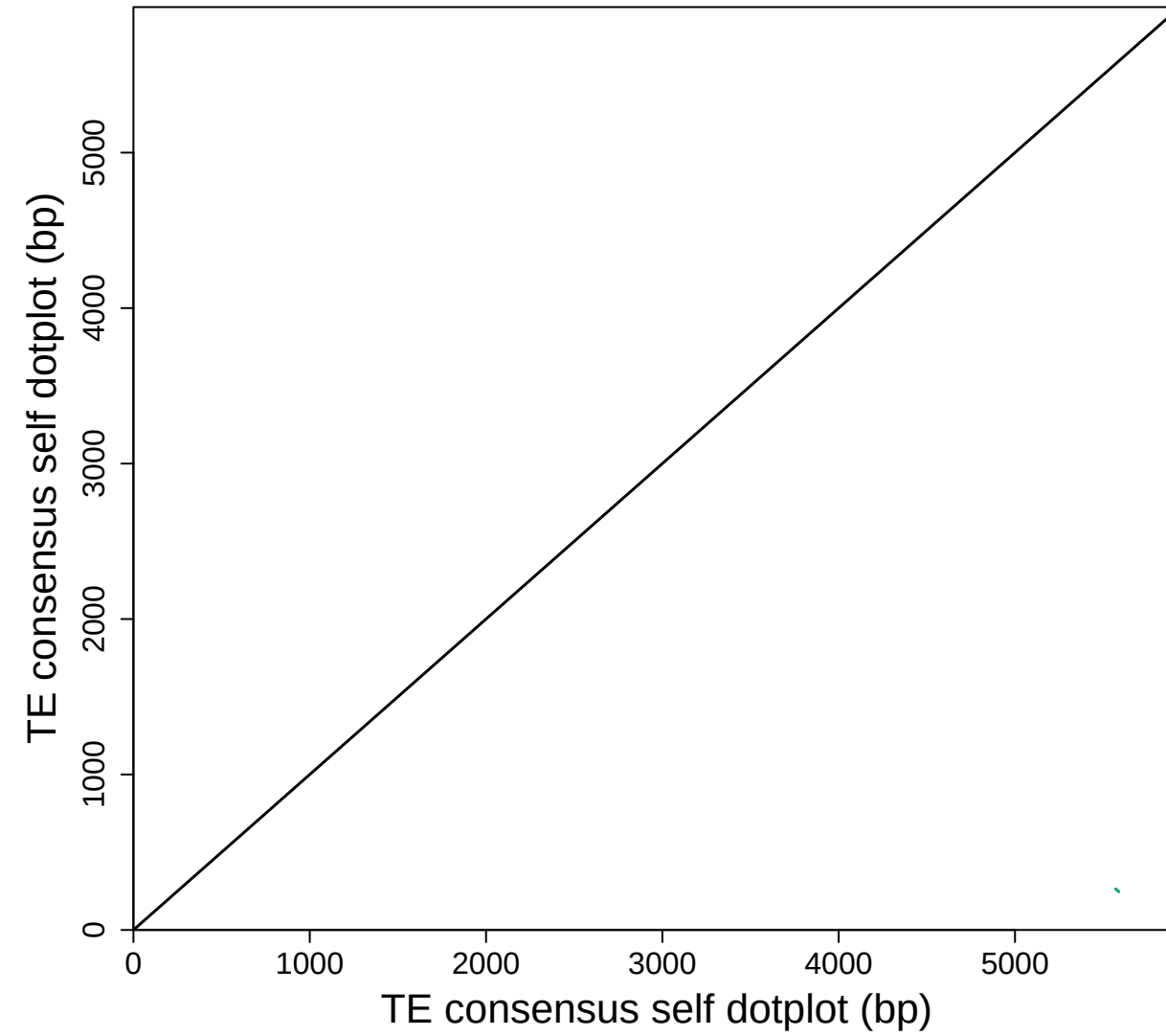
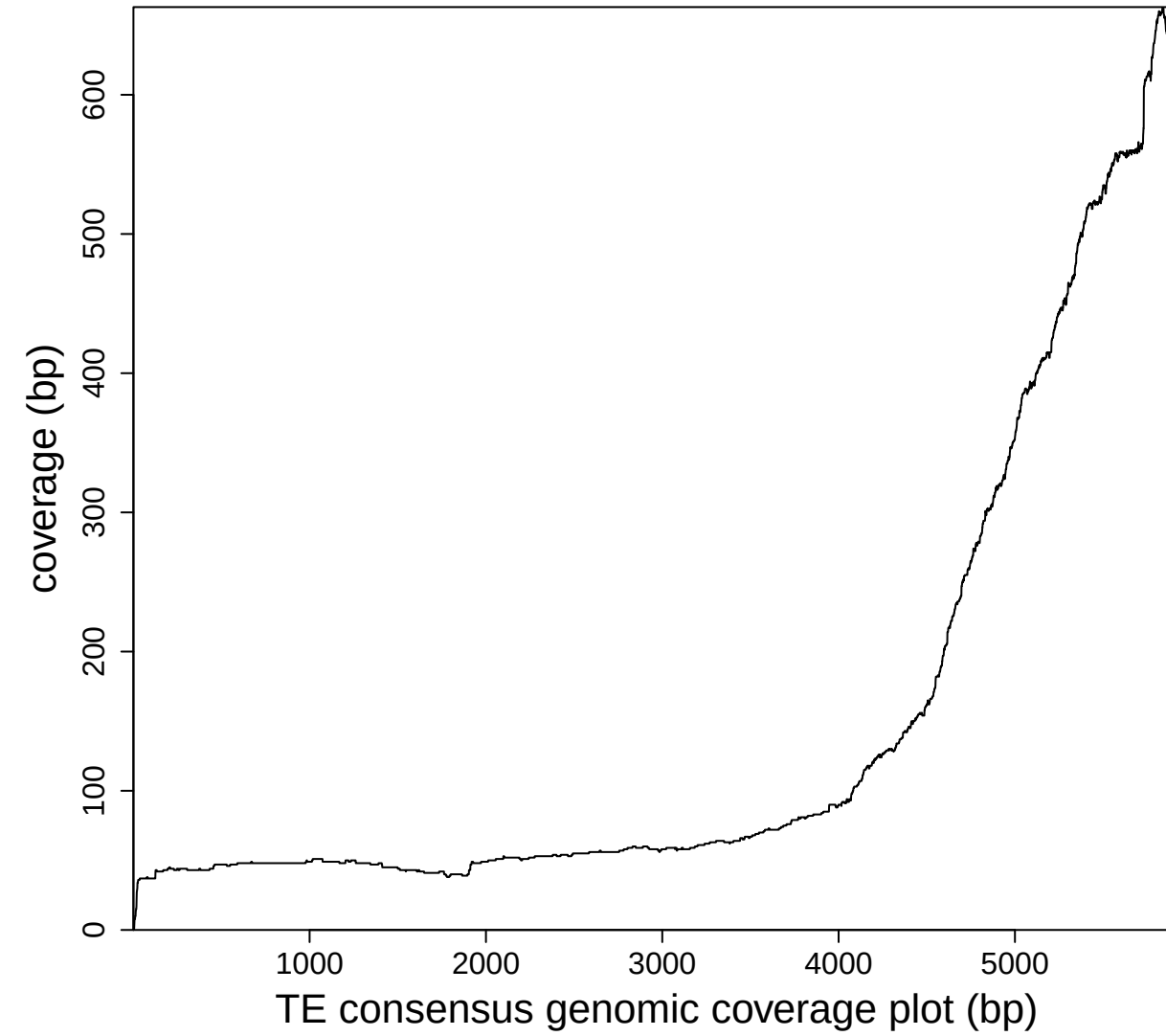
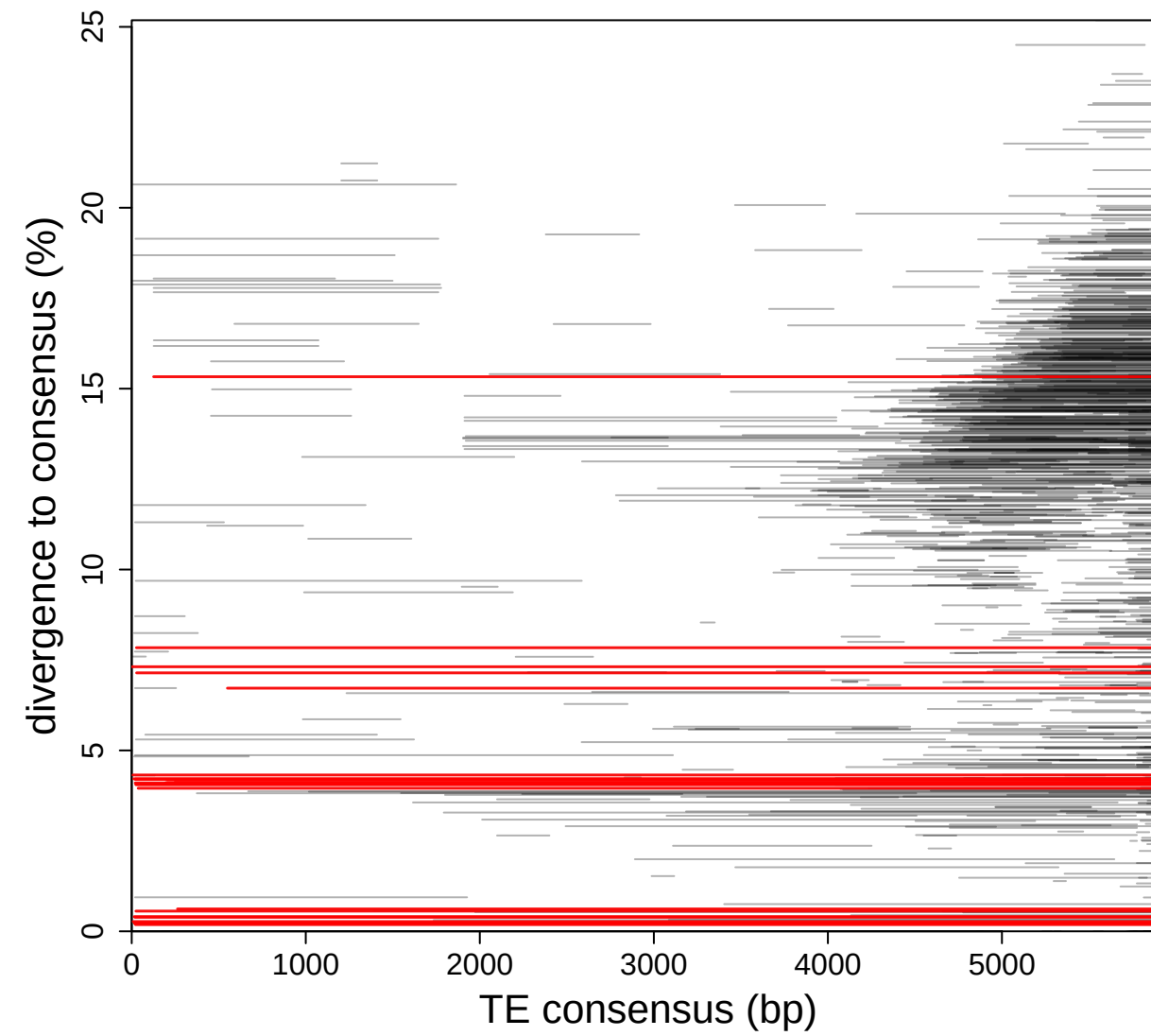
MSA length = 5936





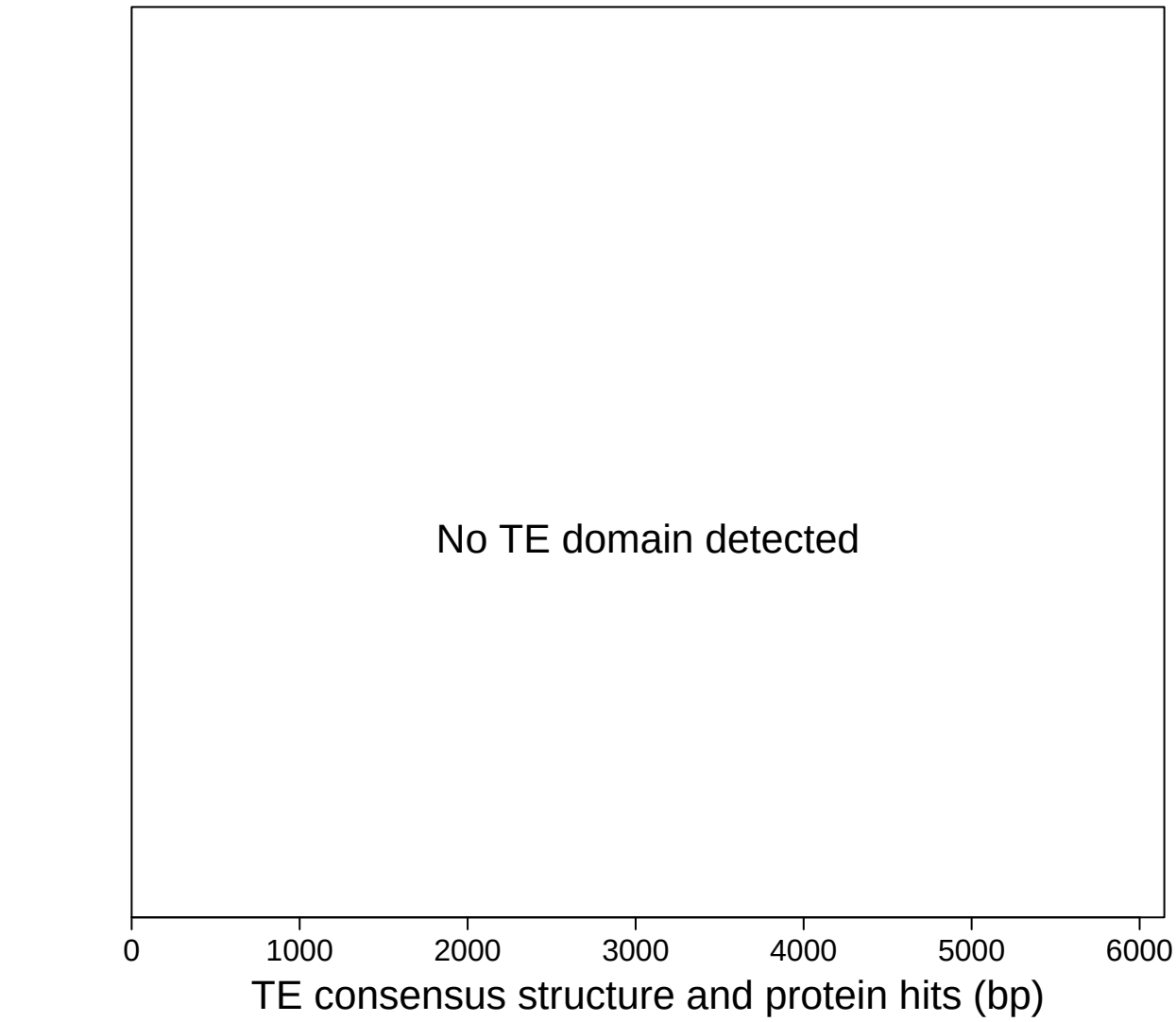
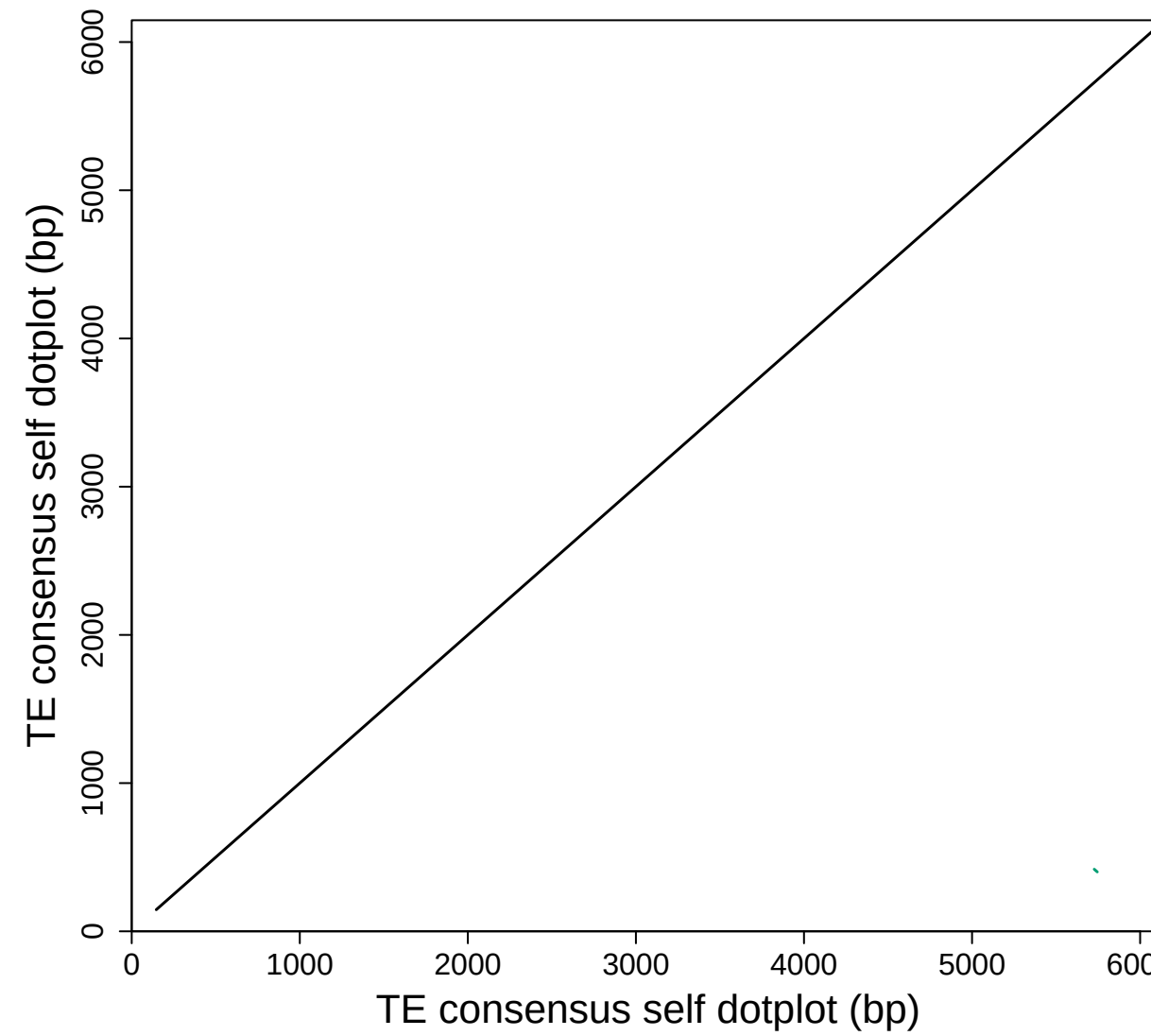
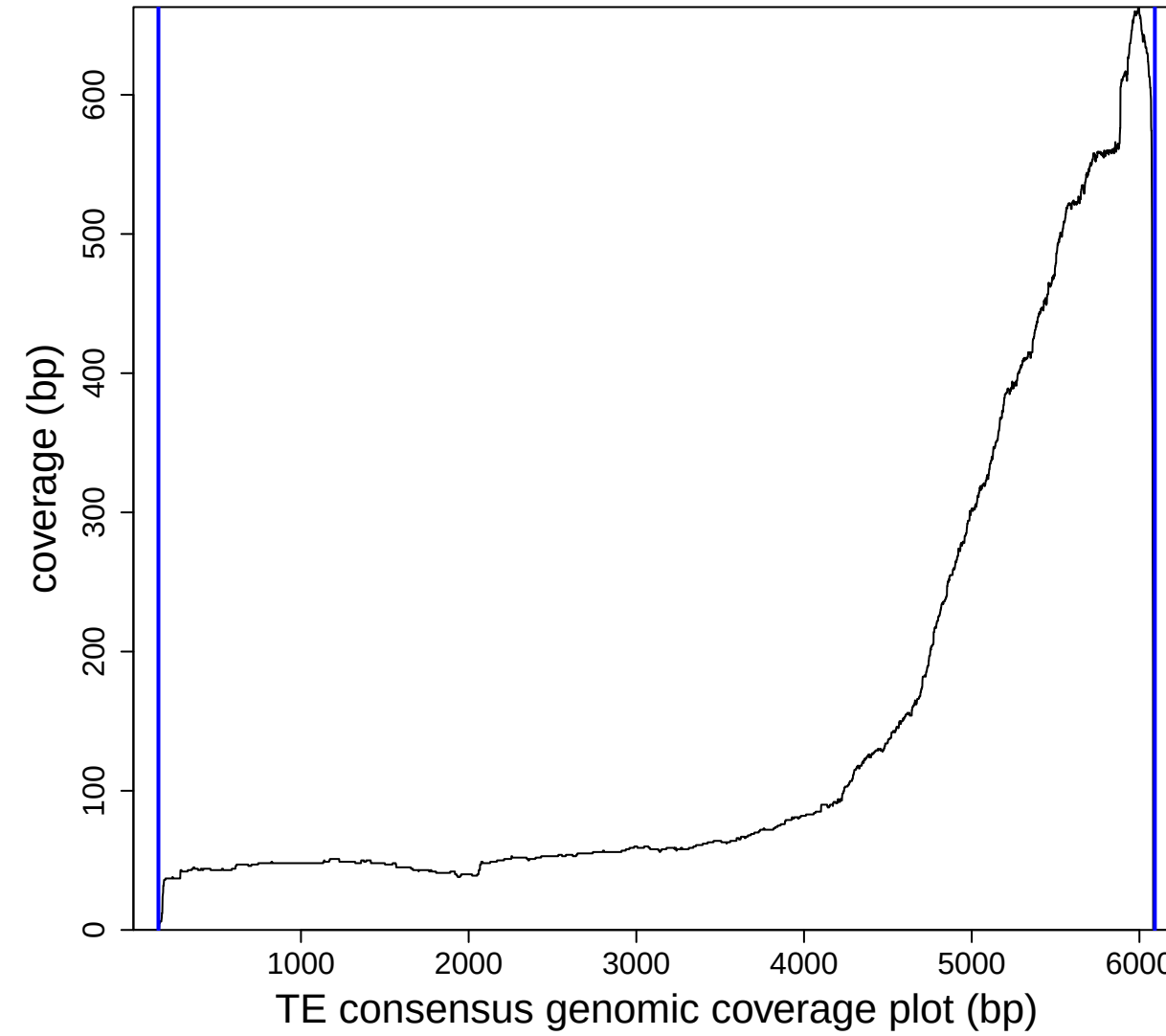
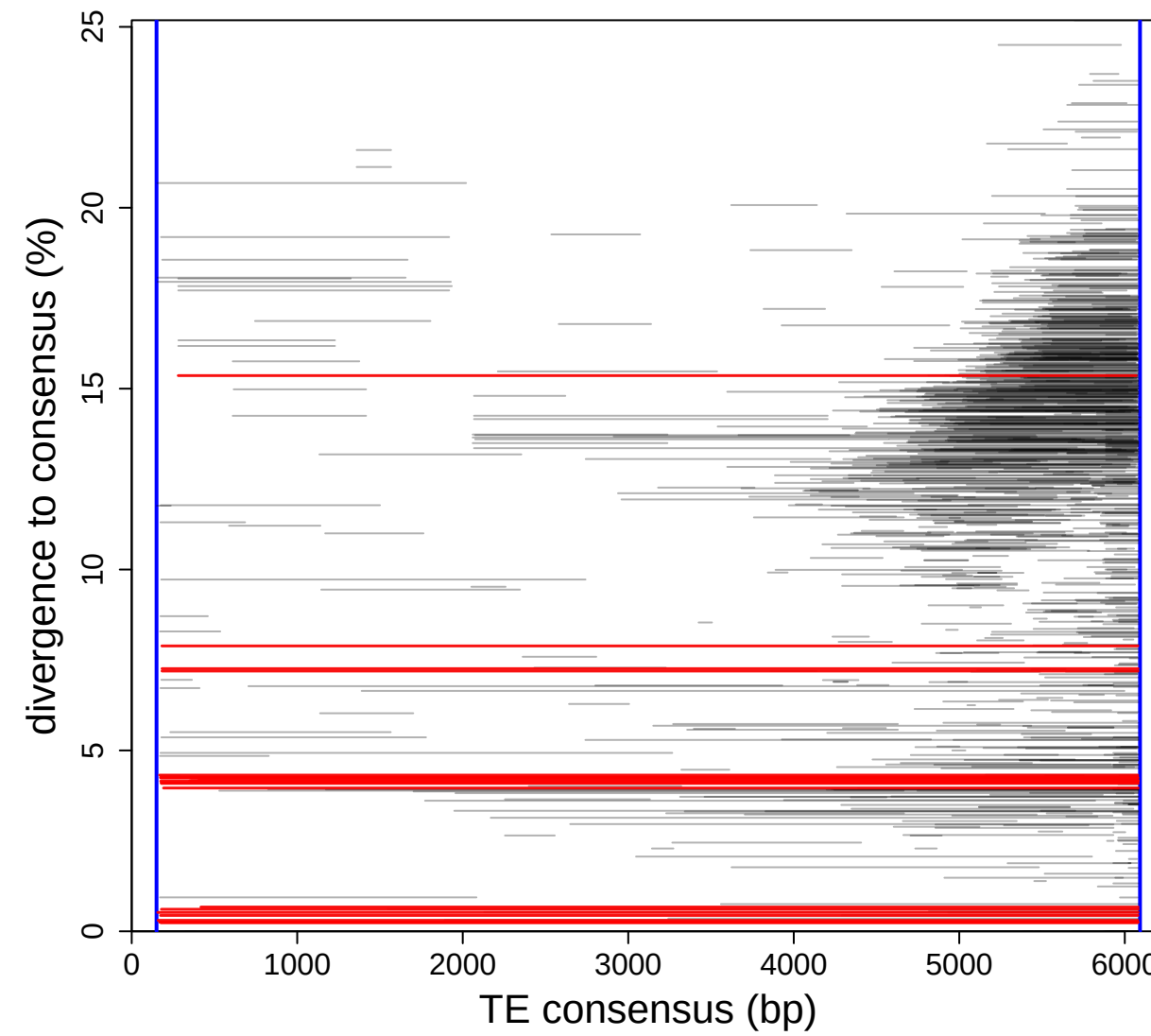
3.bed_fm 1.bed_0_0_n.bed_g 1.bed_fm 2.bed_0_0_bcln.fa_aln.f
size: 5936bp; fragments: 1030; full length: 24 (>=5342.4bp)

After TEtrimmer 5936 bp

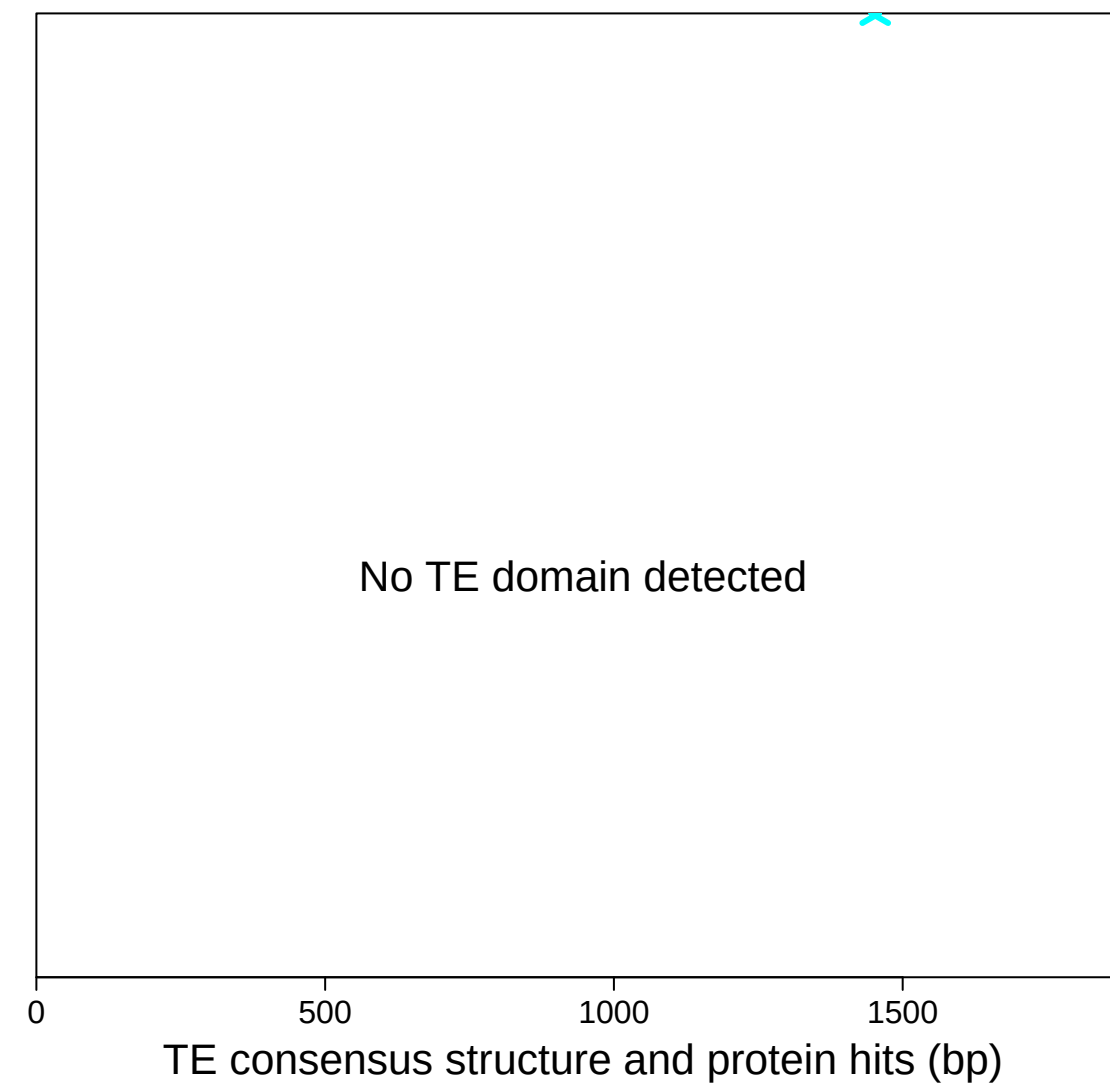
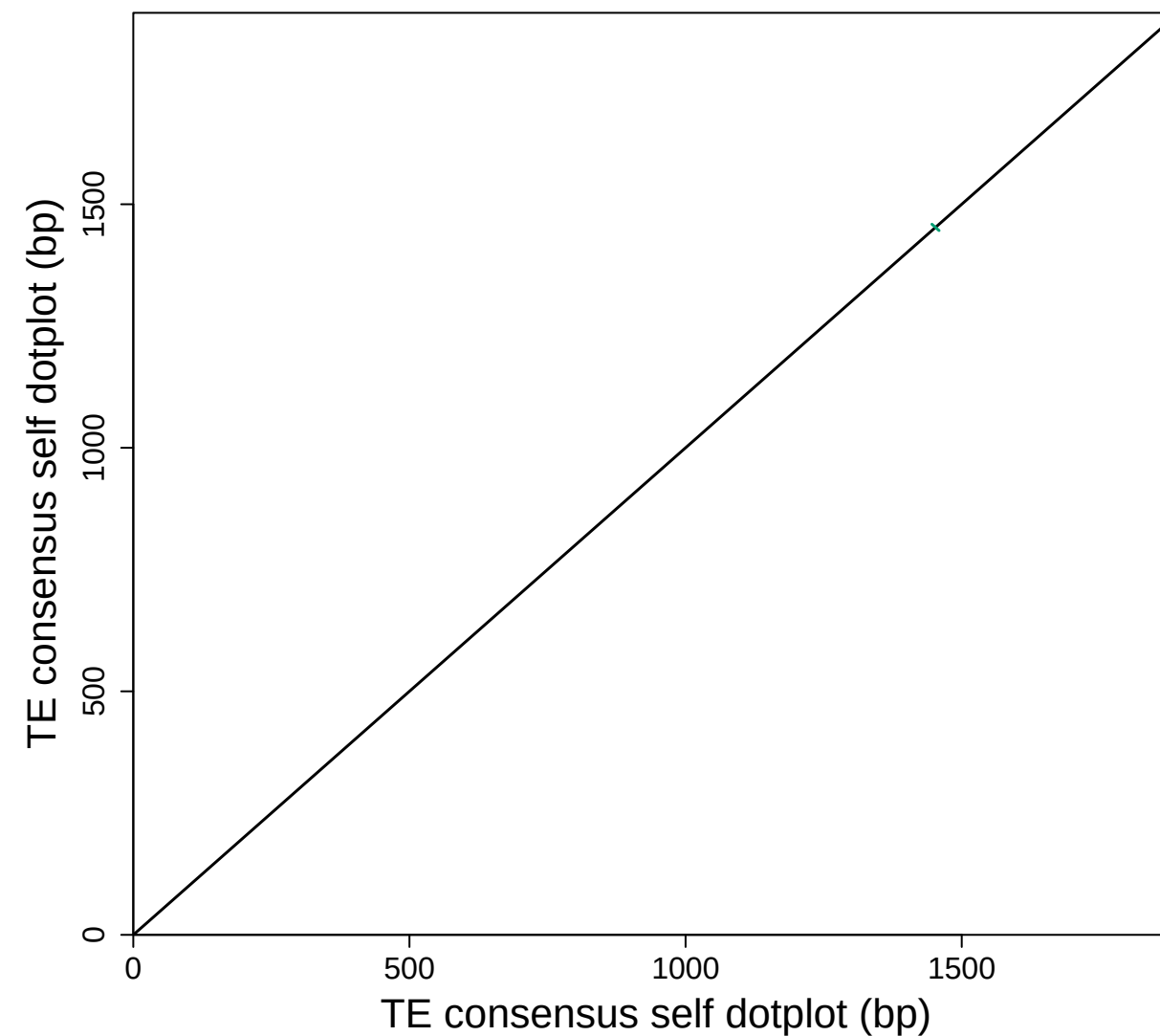
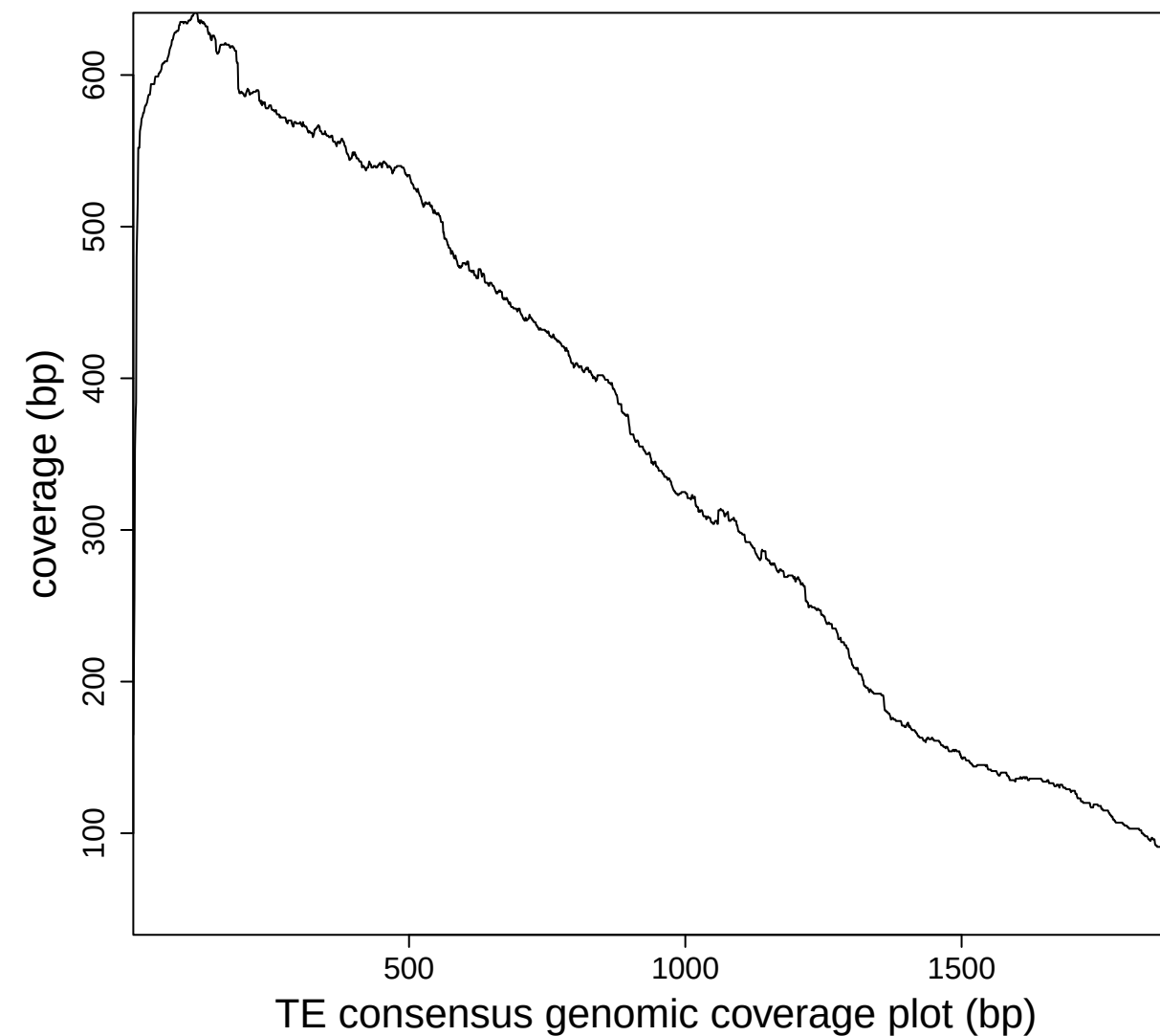
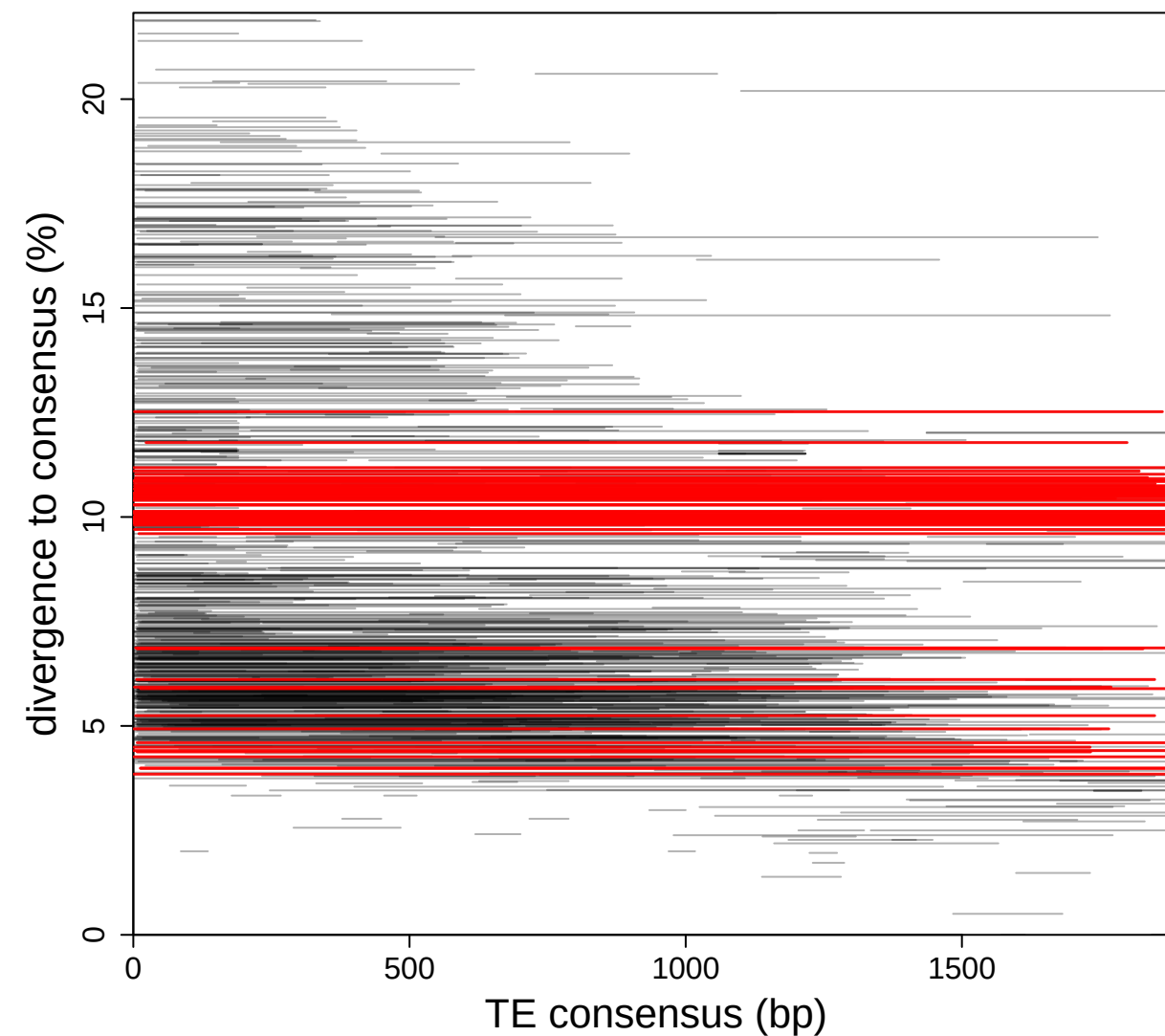


ed_g_3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.f
size: 6242bp; fragments: 1030; full length: 23 (>=5617.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

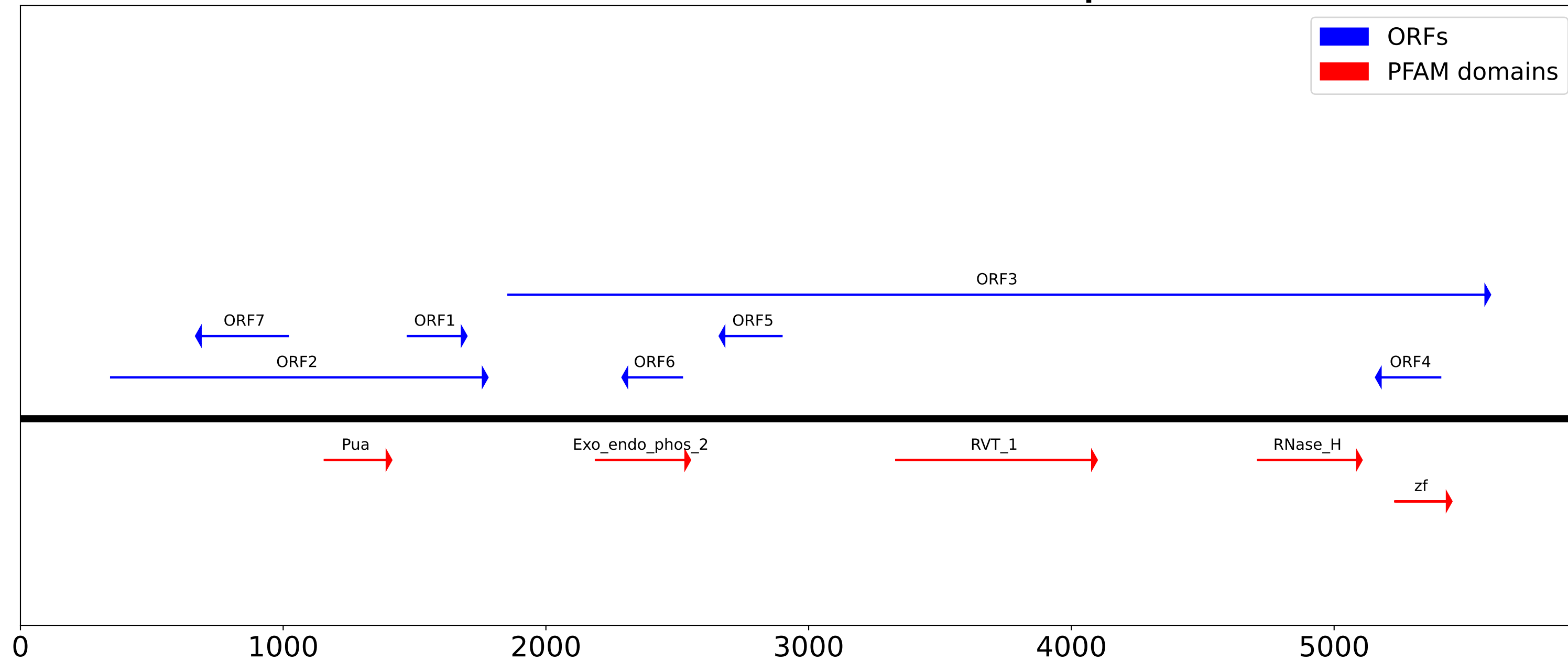


TE: rnd_5_family_5963
size: 1893bp; fragments: 980; full length: 68 (≥ 1703.7 bp)

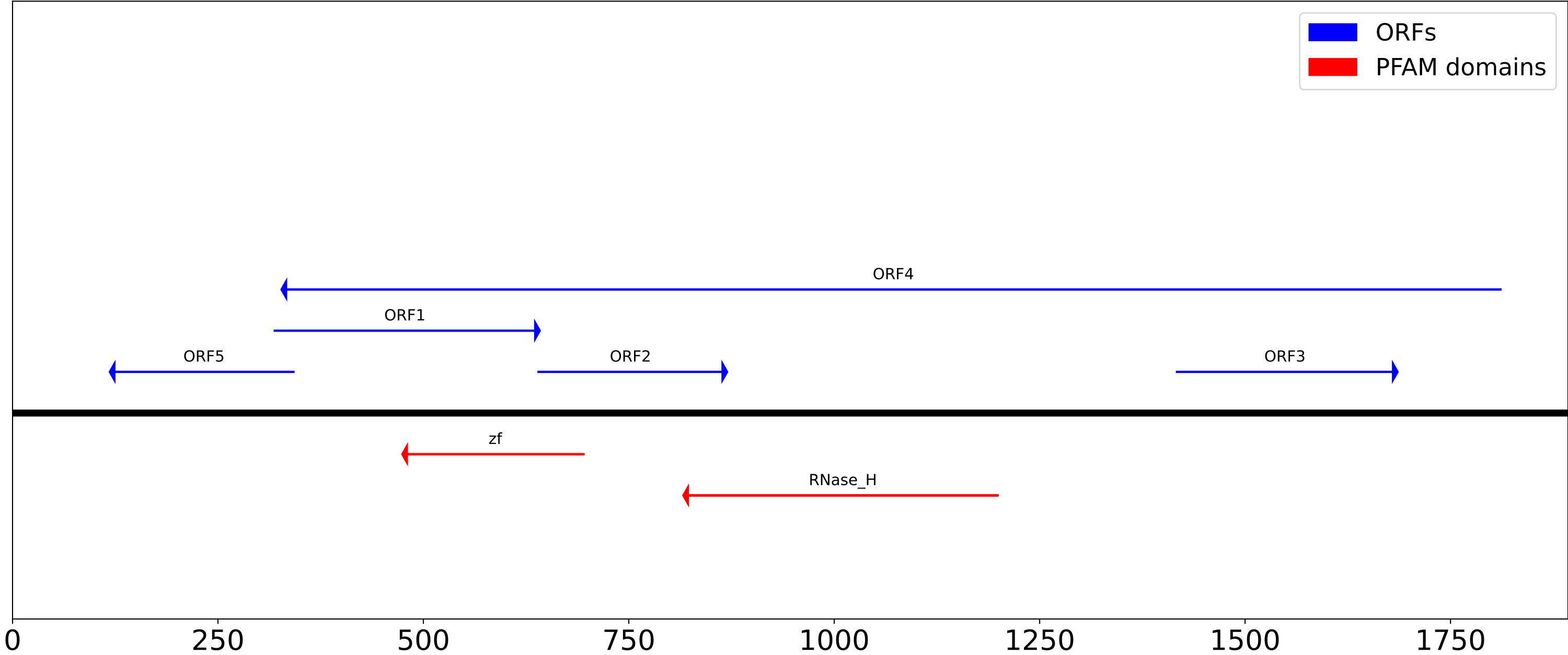


Before TEtrimmer 1893 bp

After TEtrimmer ORF and PFAM domain plot



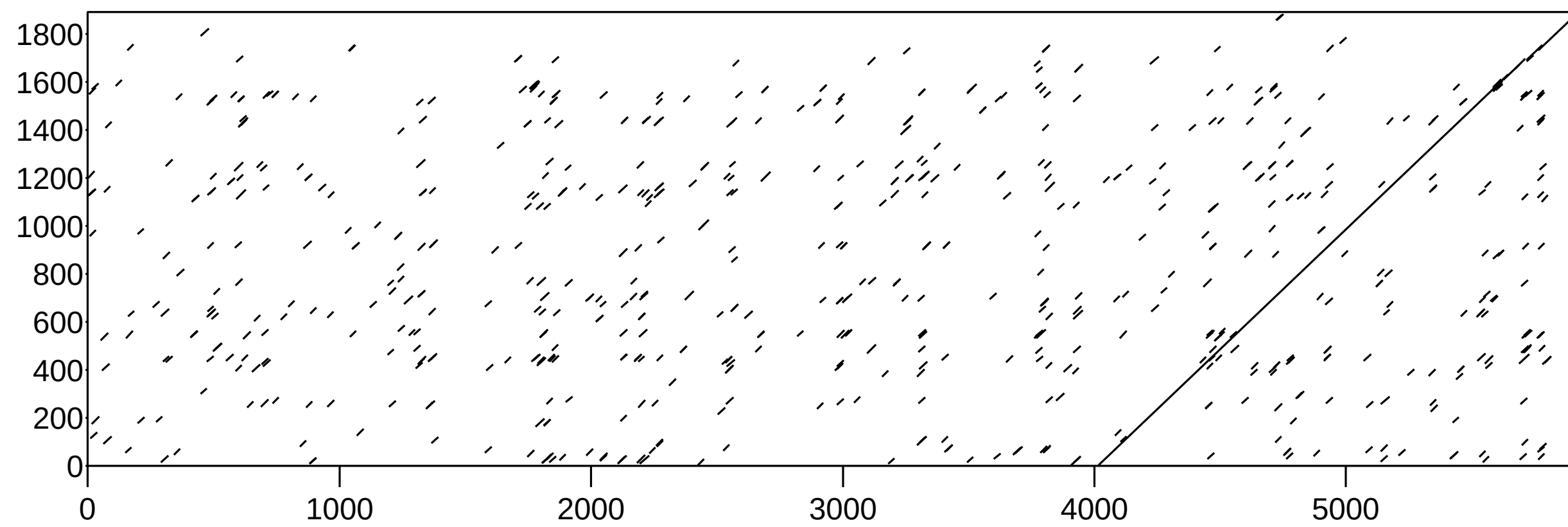
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)