

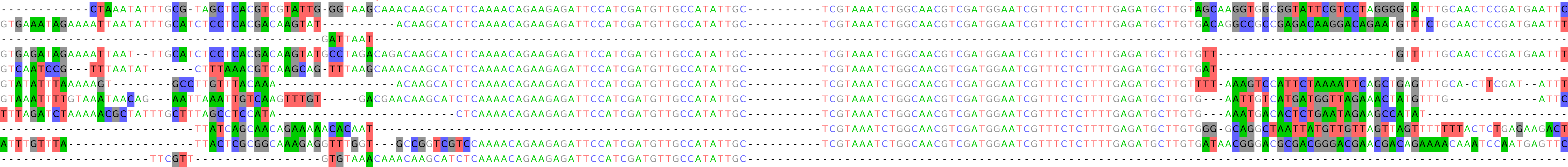
Start crop Point



End crop Point

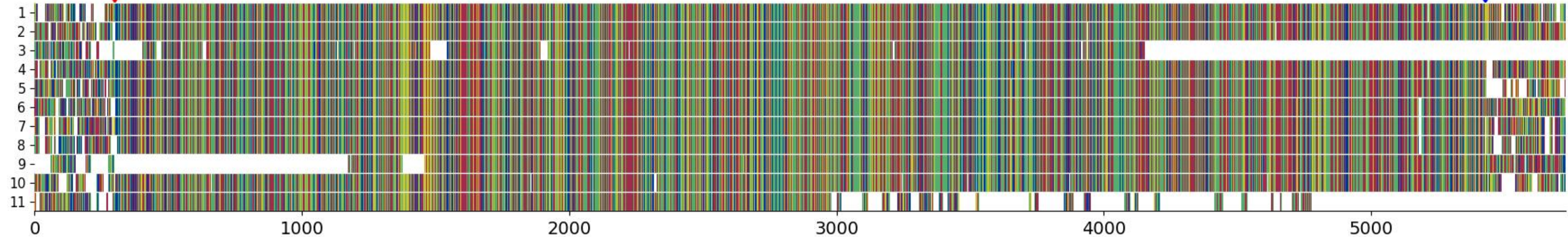


MSA length = 5129



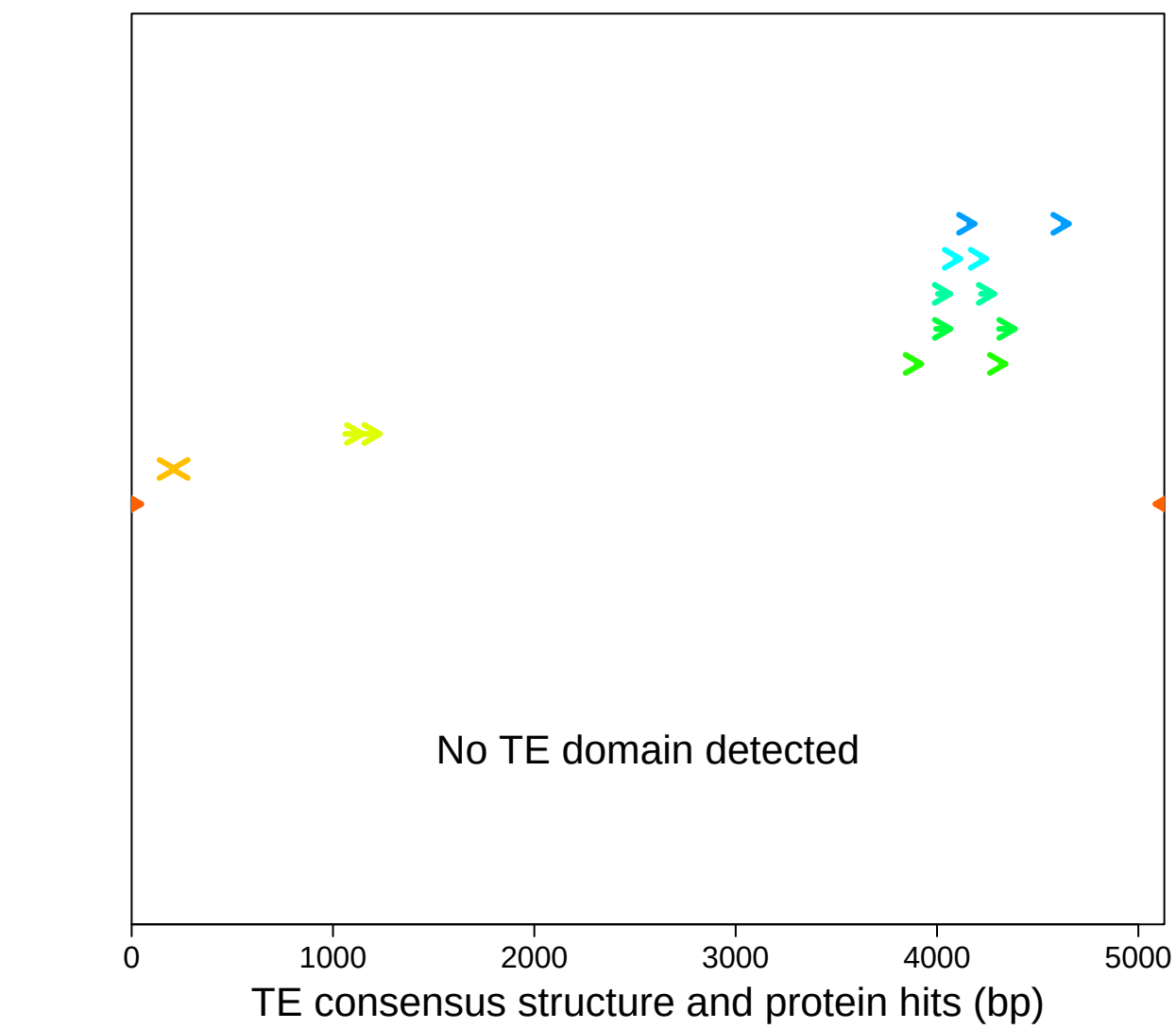
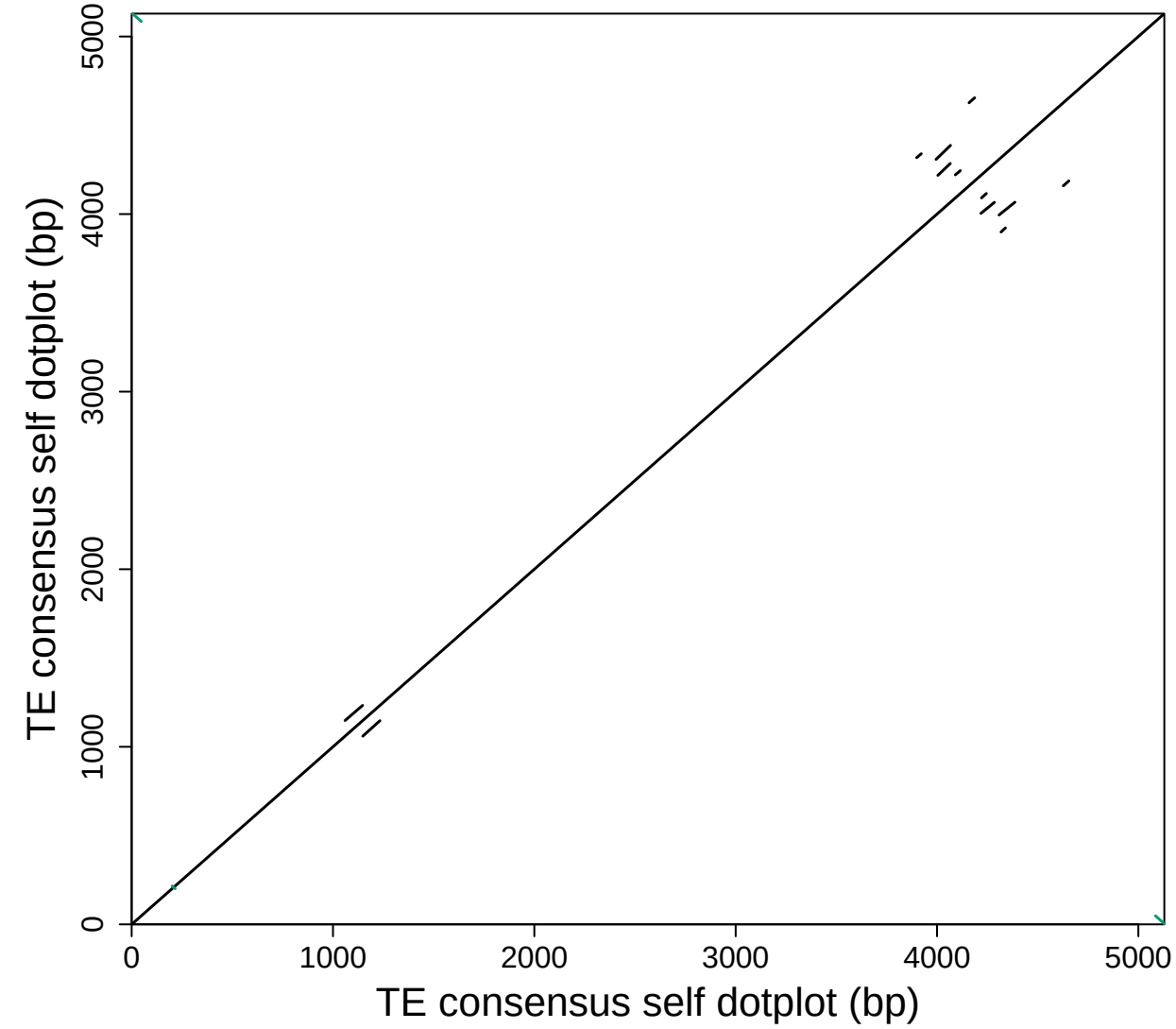
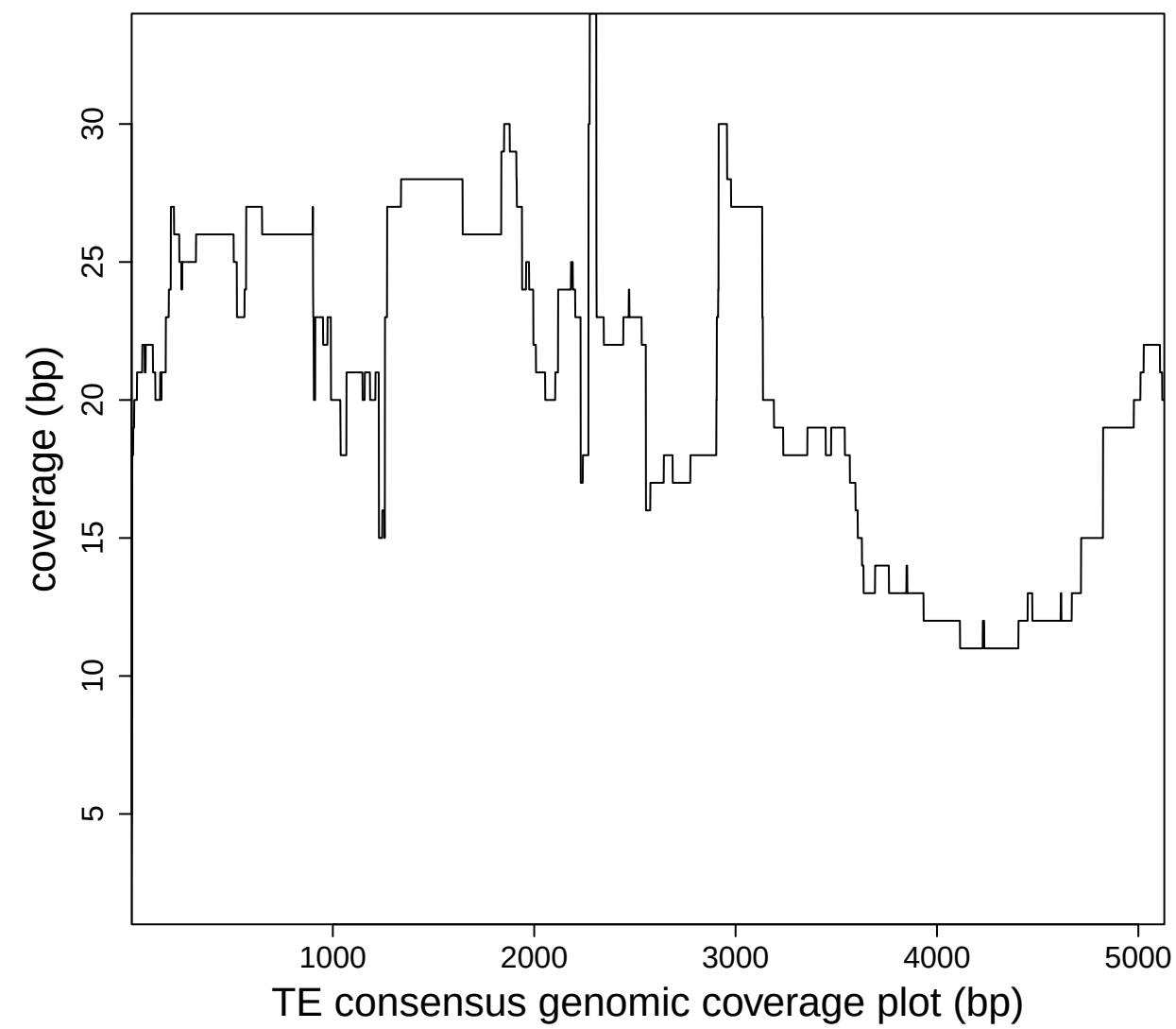
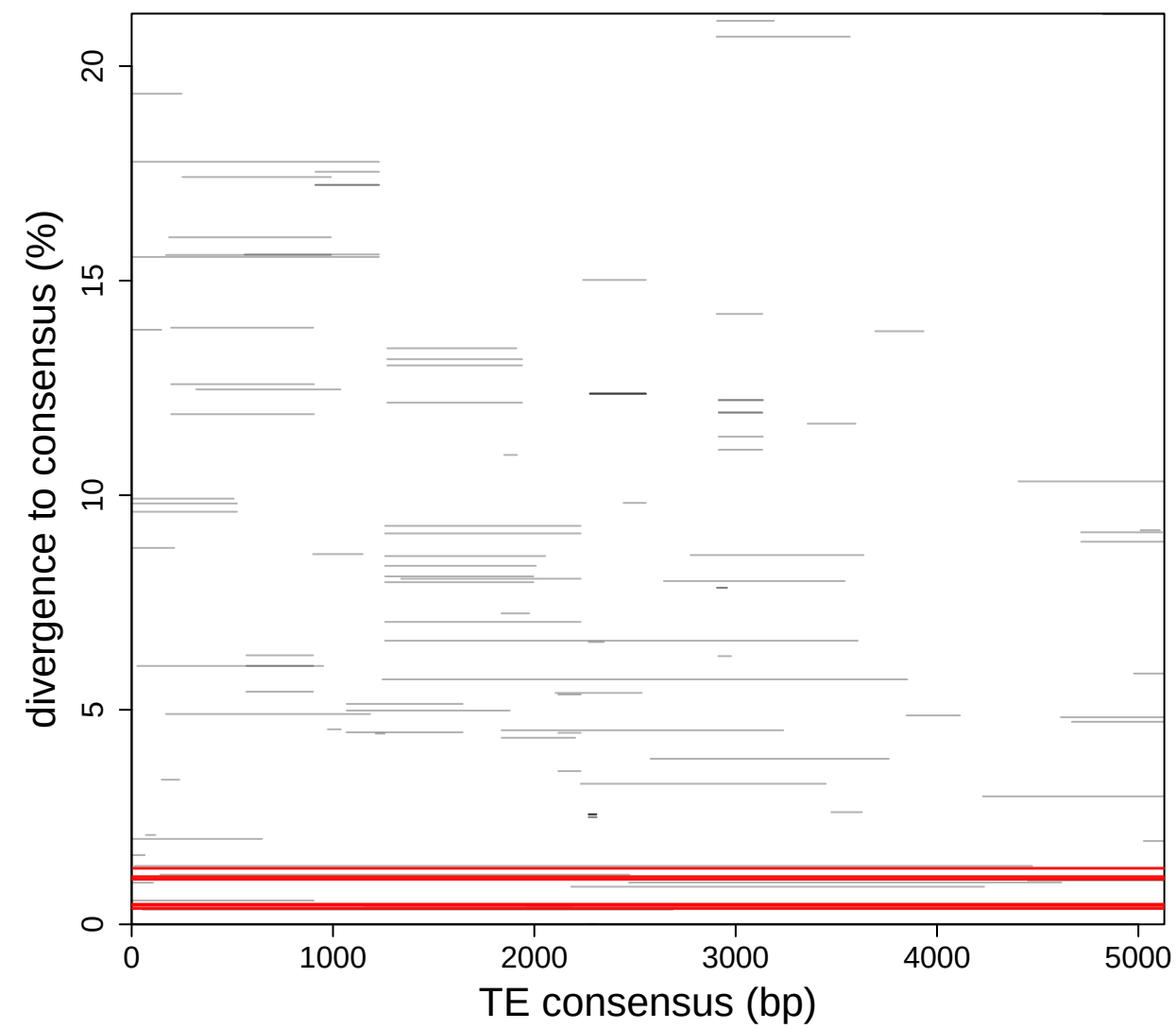
Start crop Point

End crop Point



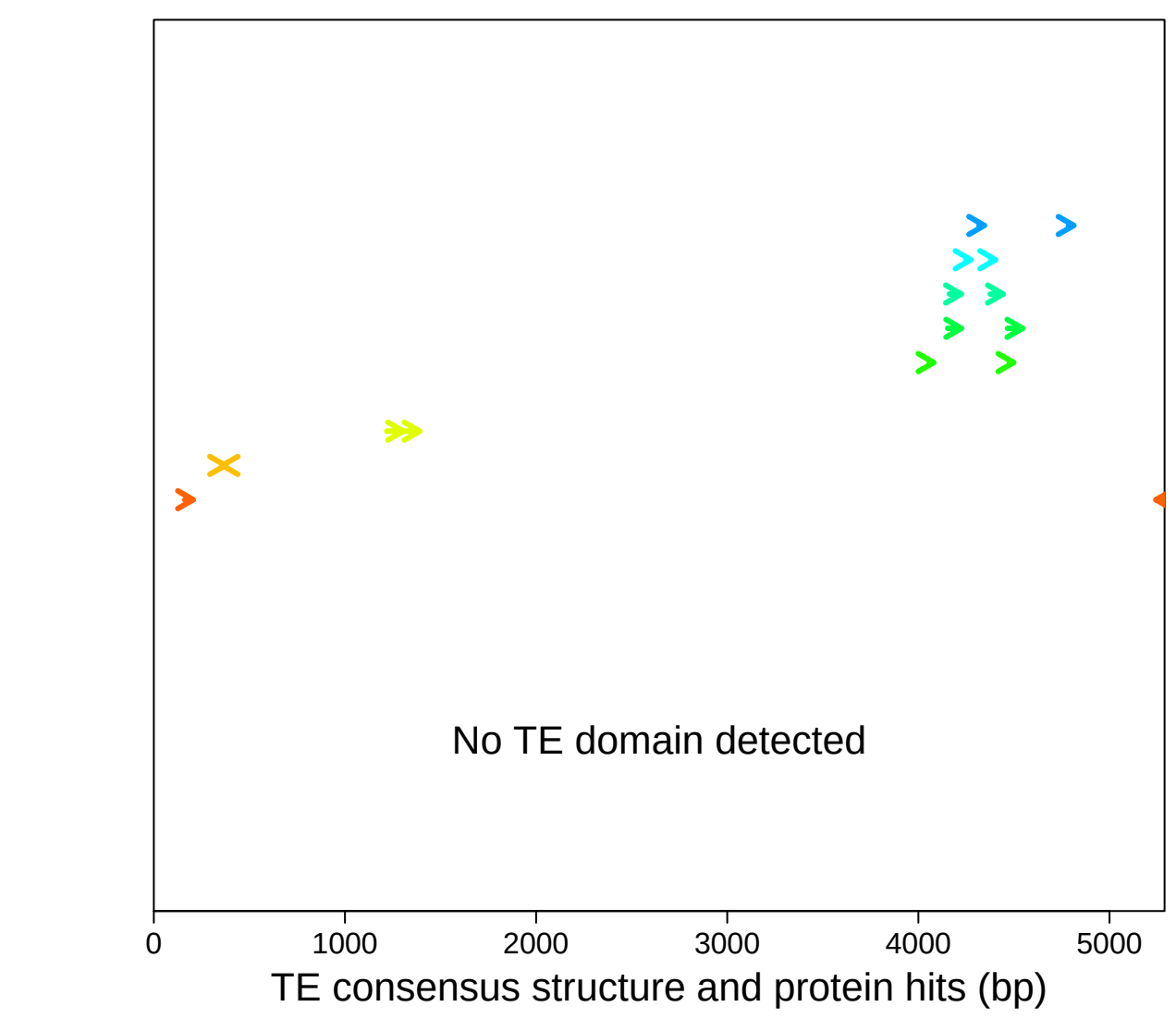
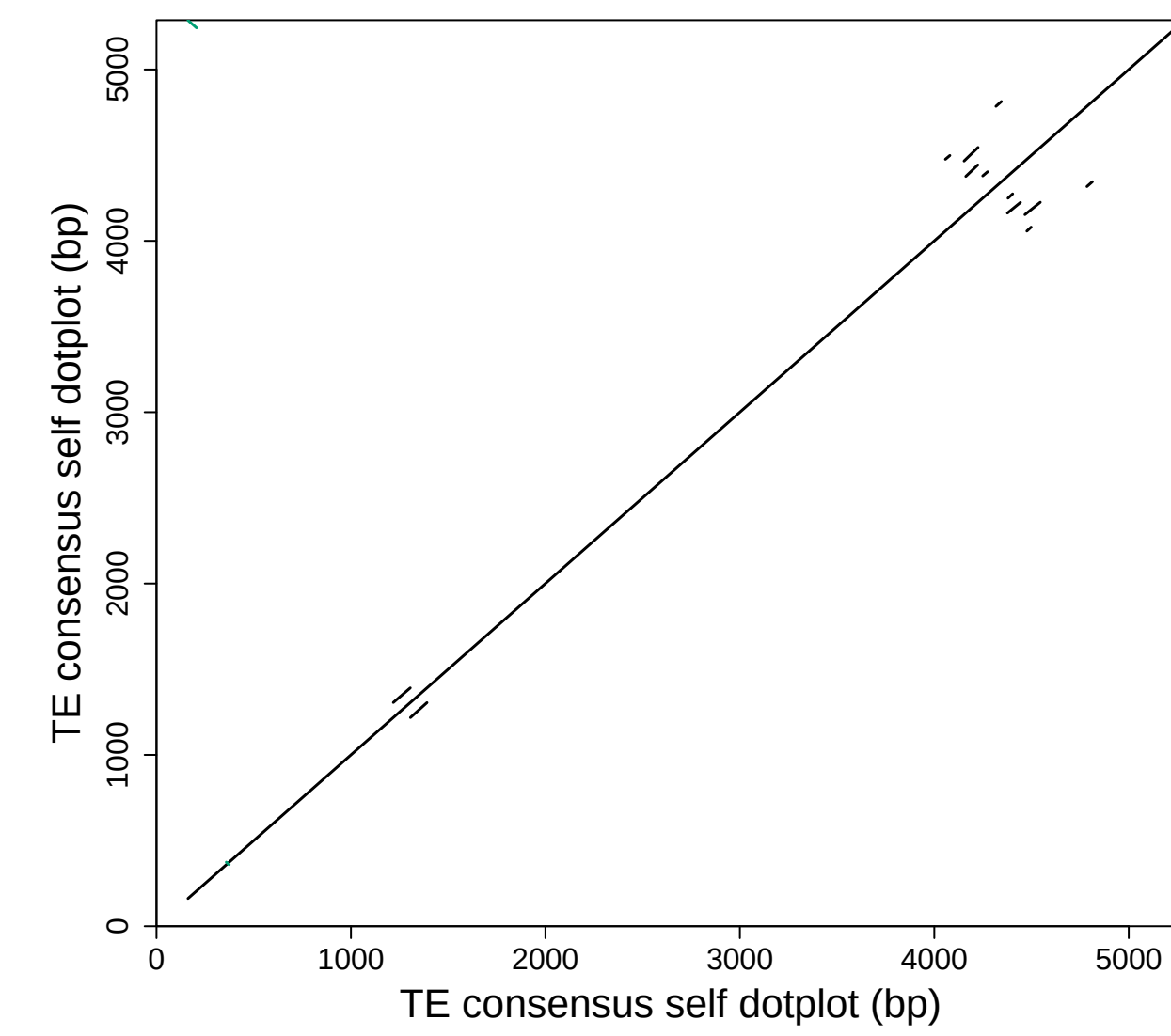
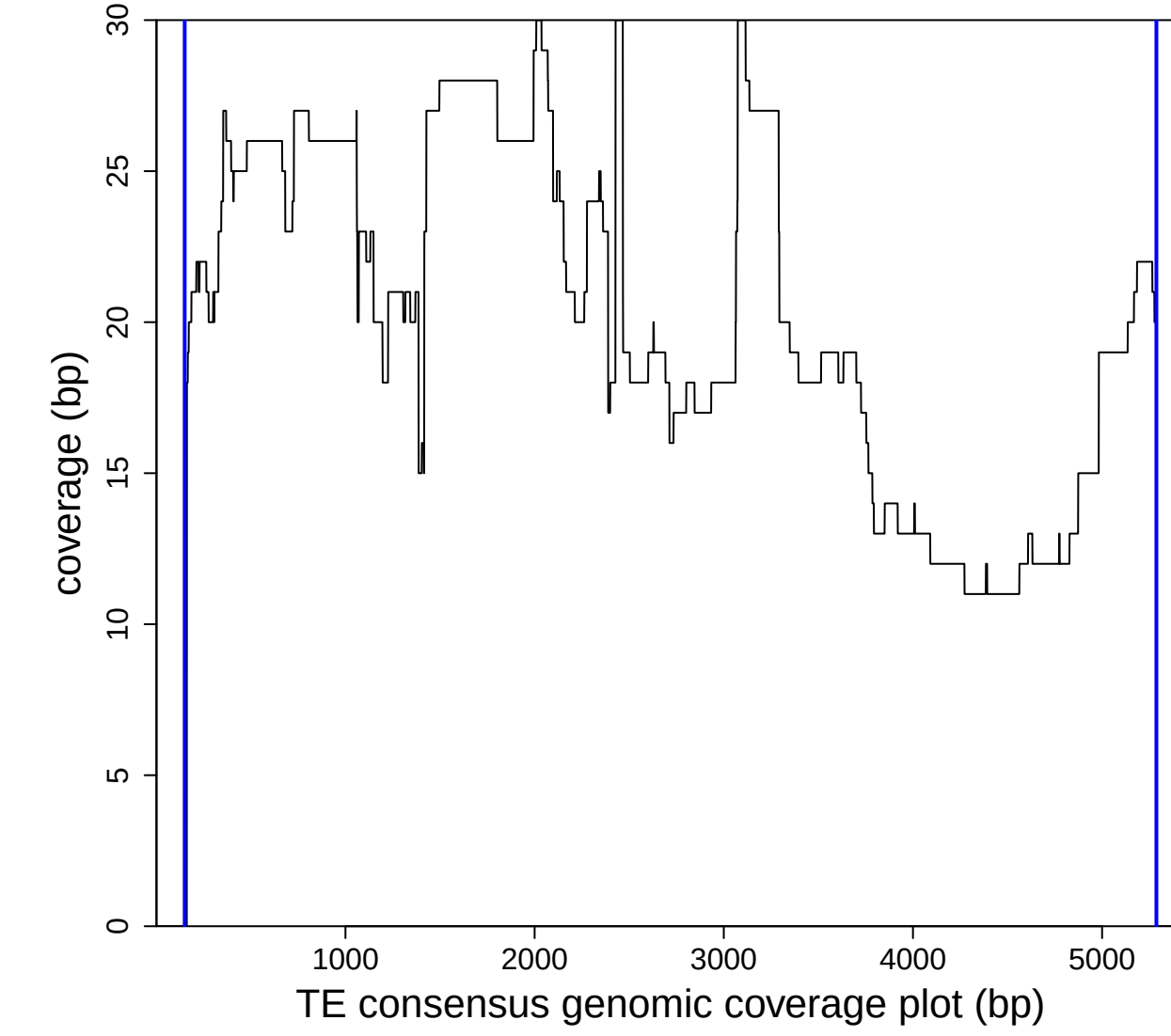
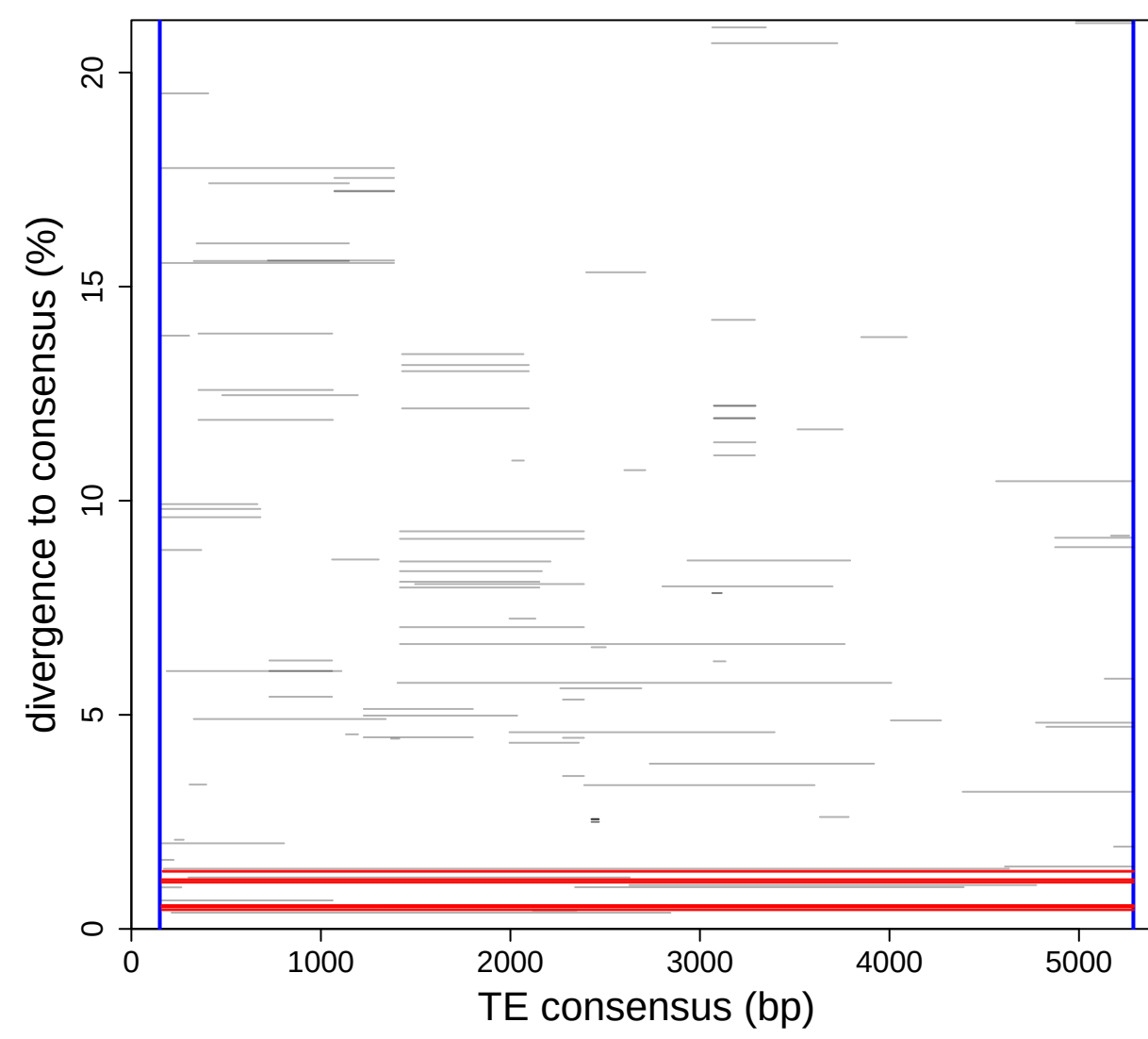
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_ε
size: 5129bp; fragments: 124; full length: 7 (>=4616.1bp)

After TETrimmer 5129 bp

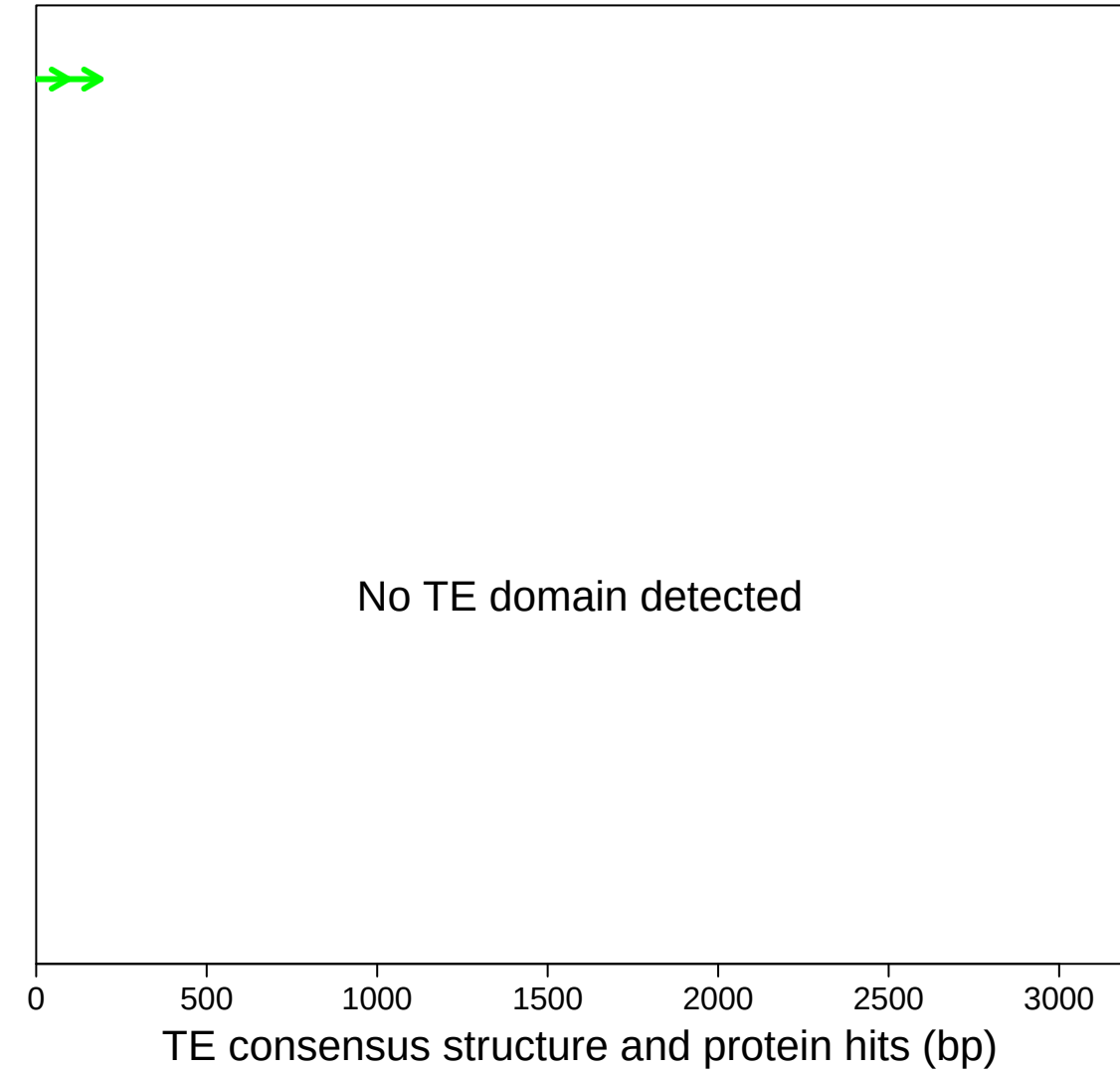
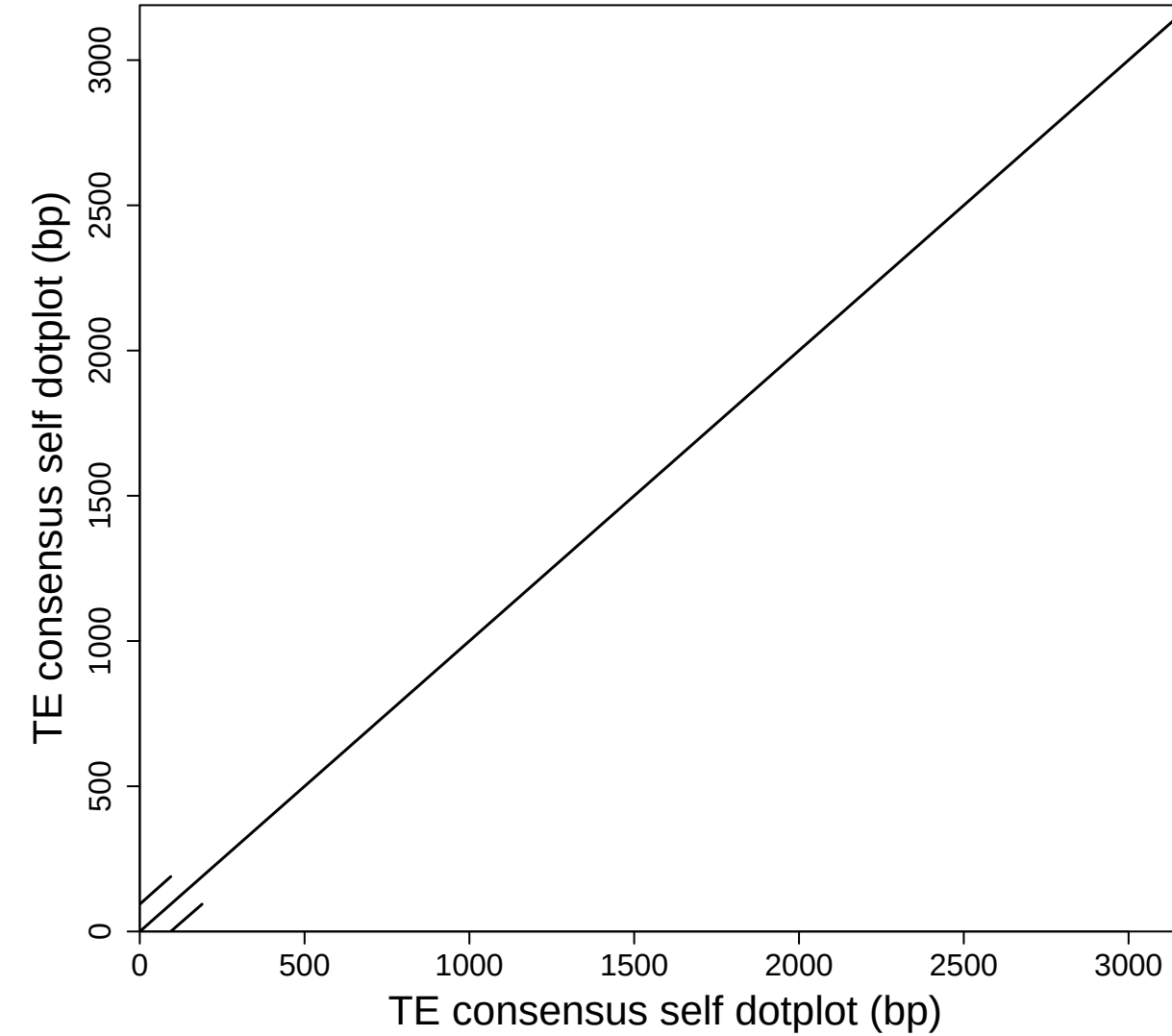
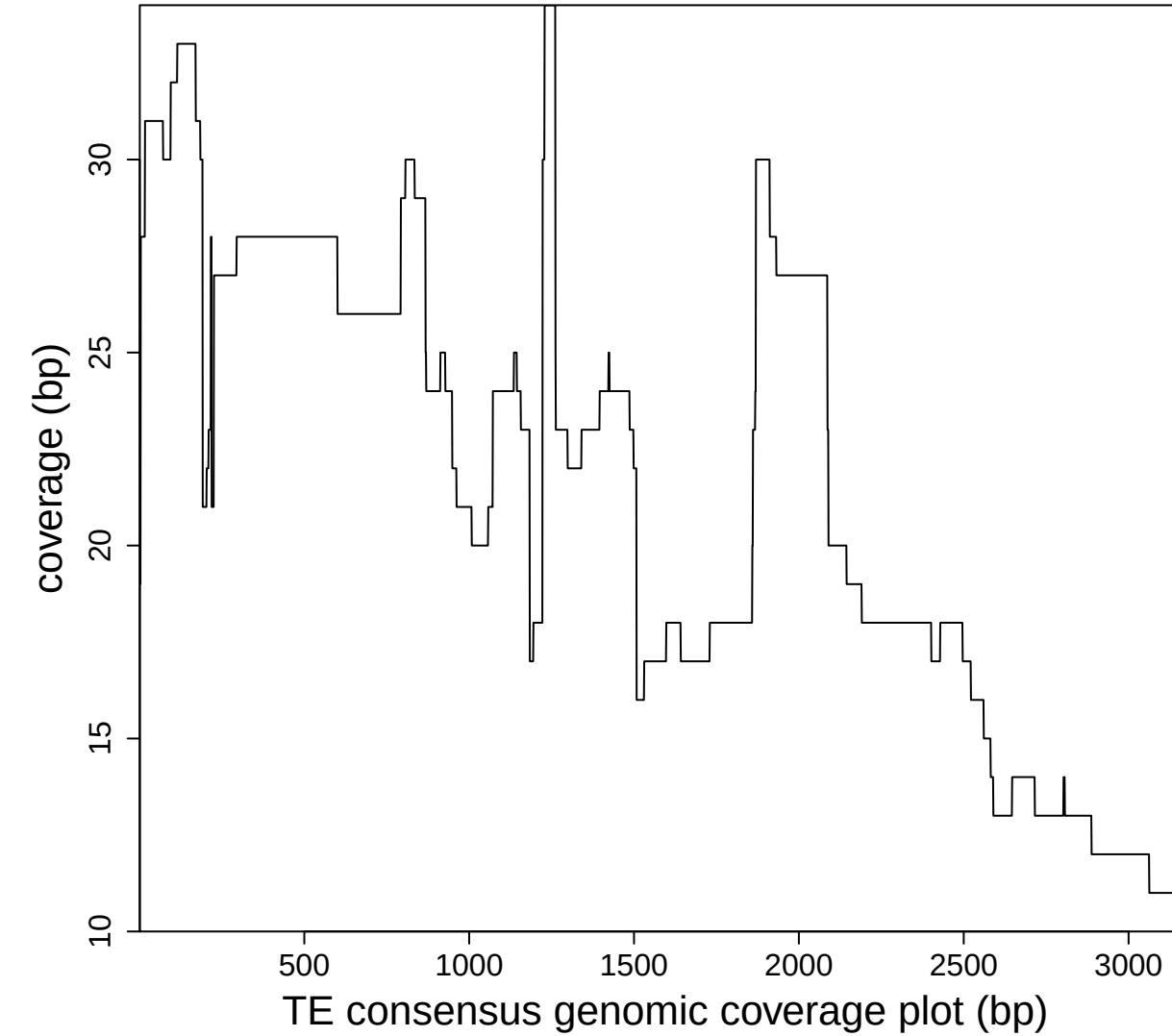
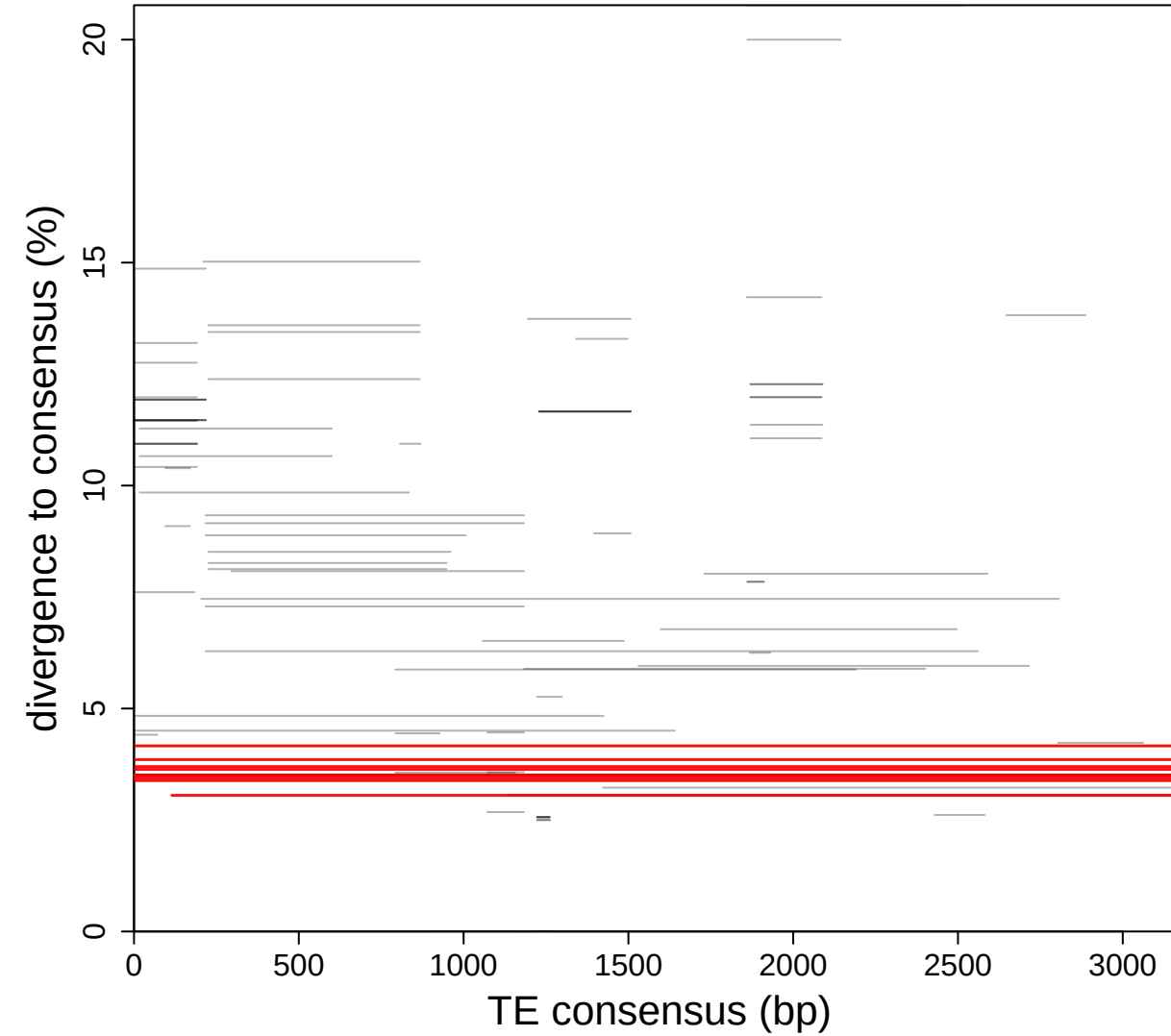


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 5437bp; fragments: 120; full length: 7 (>=4893.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

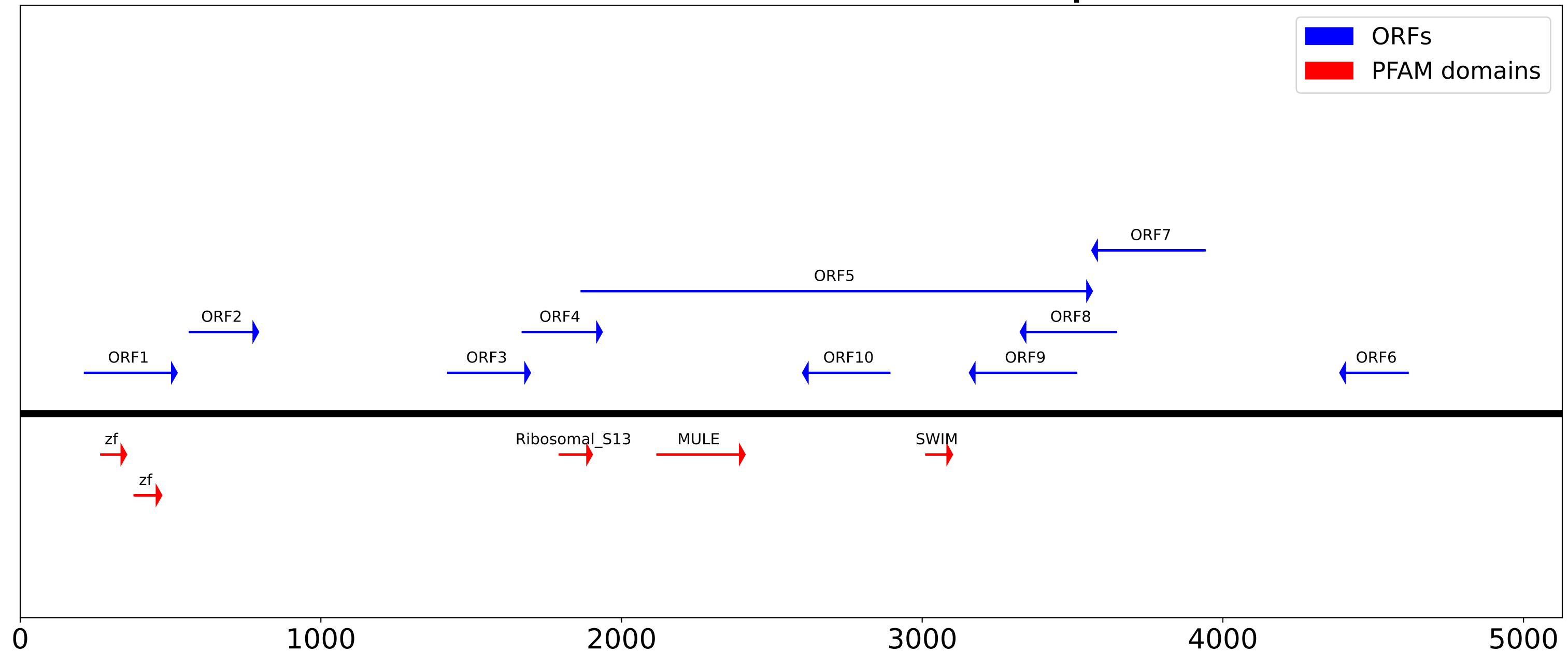


TE: rnd_5_family_6445
size: 3189bp; fragments: 97; full length: 9 (≥ 2870.1 bp)

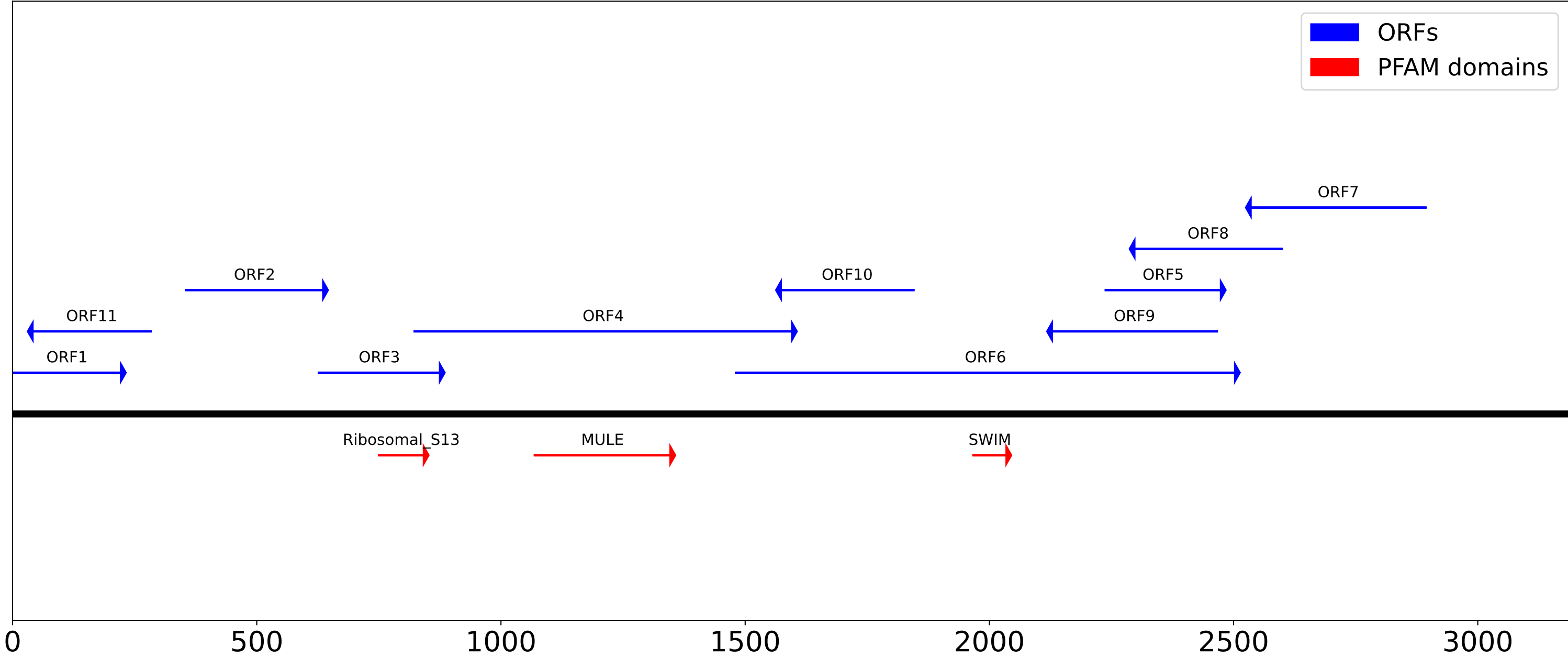


Before TETrimmer 3189 bp

After TEtrimmer ORF and PFAM domain plot



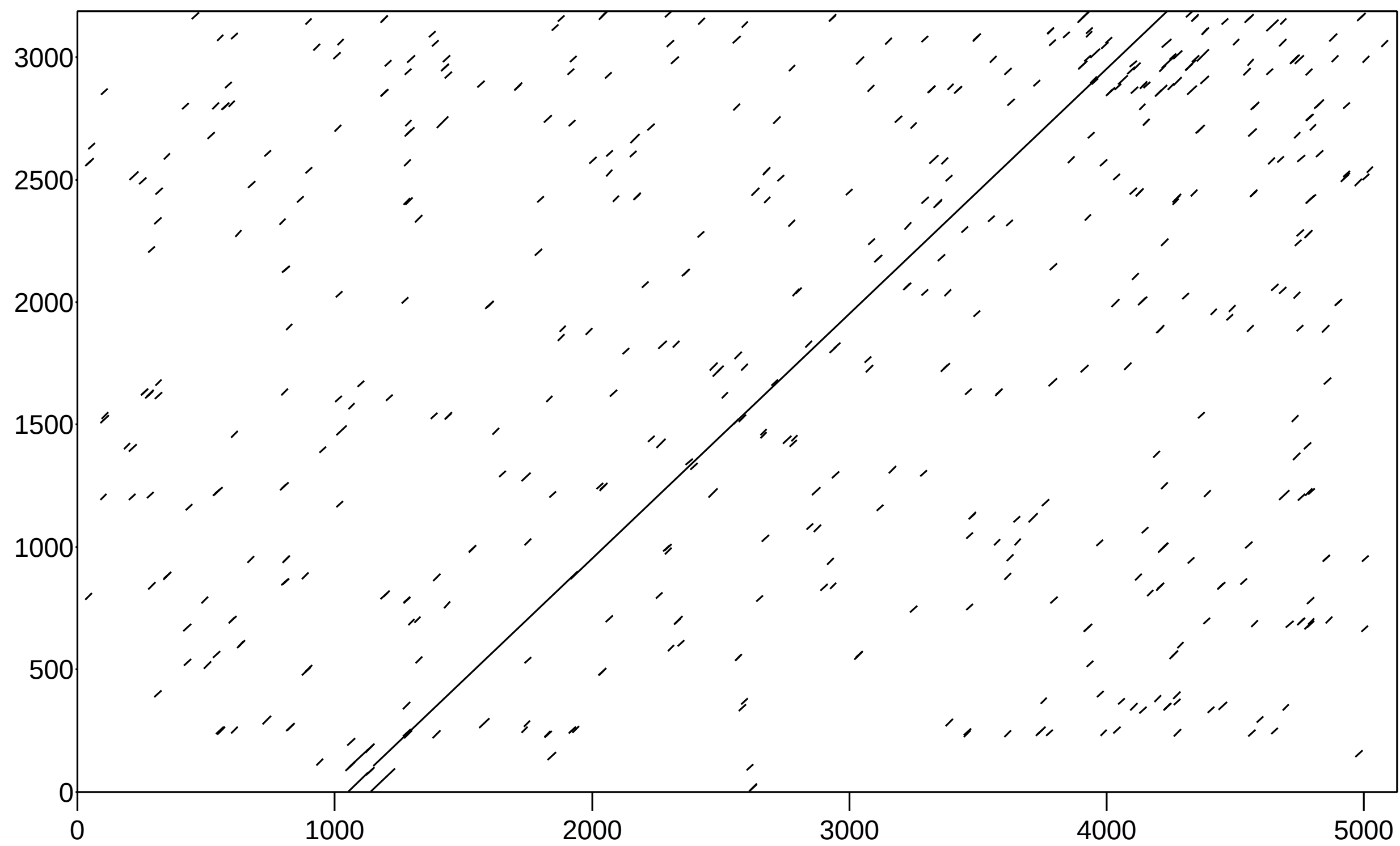
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)