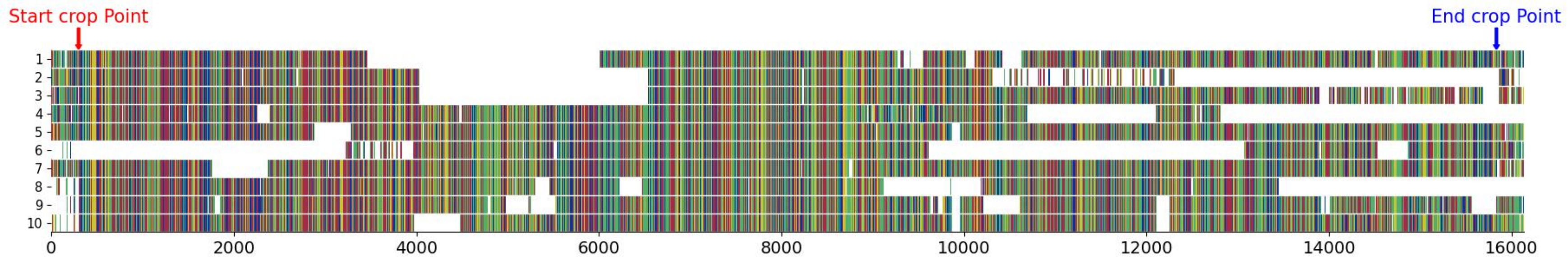


Start Crop Point

End Crop Point

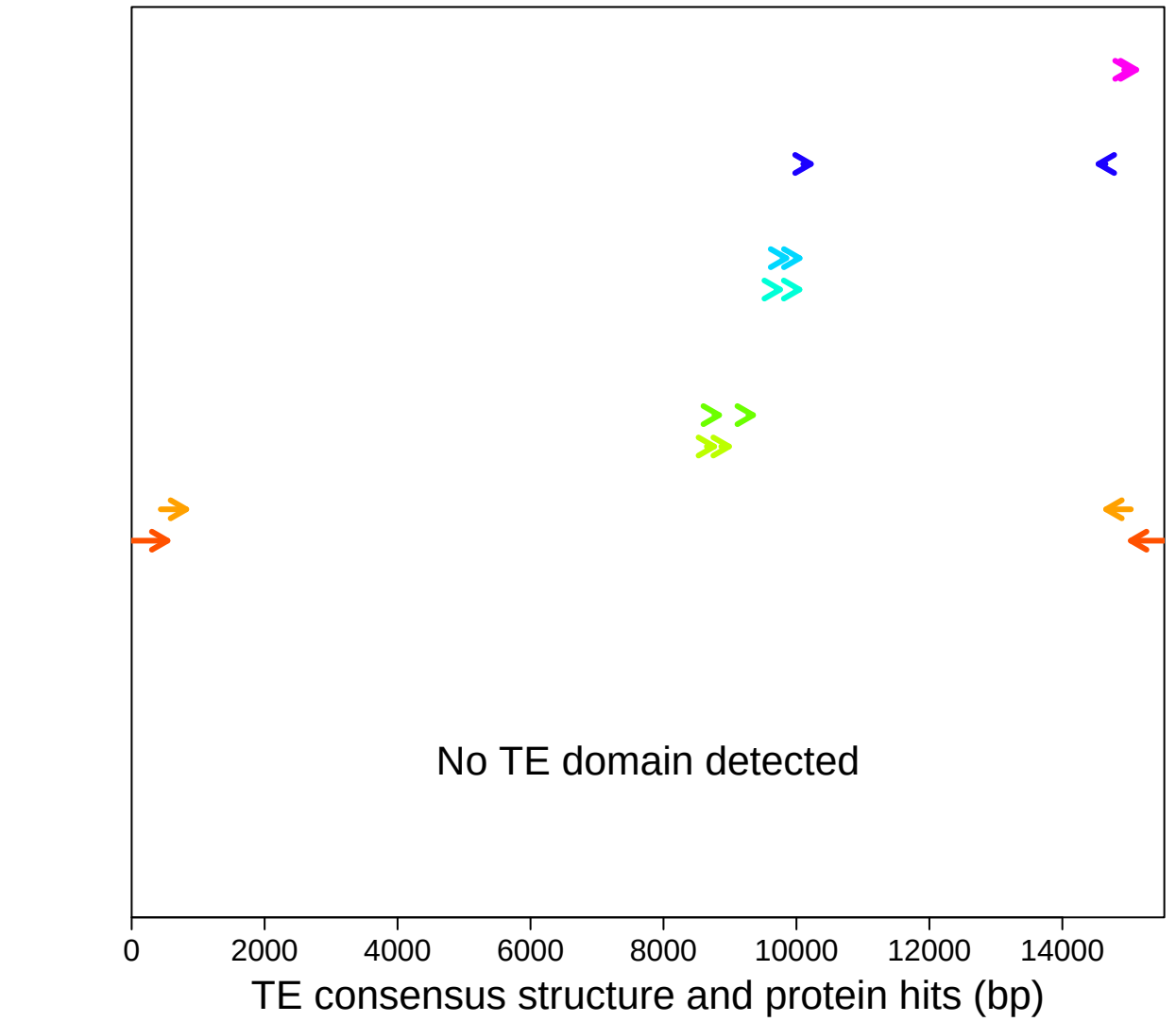
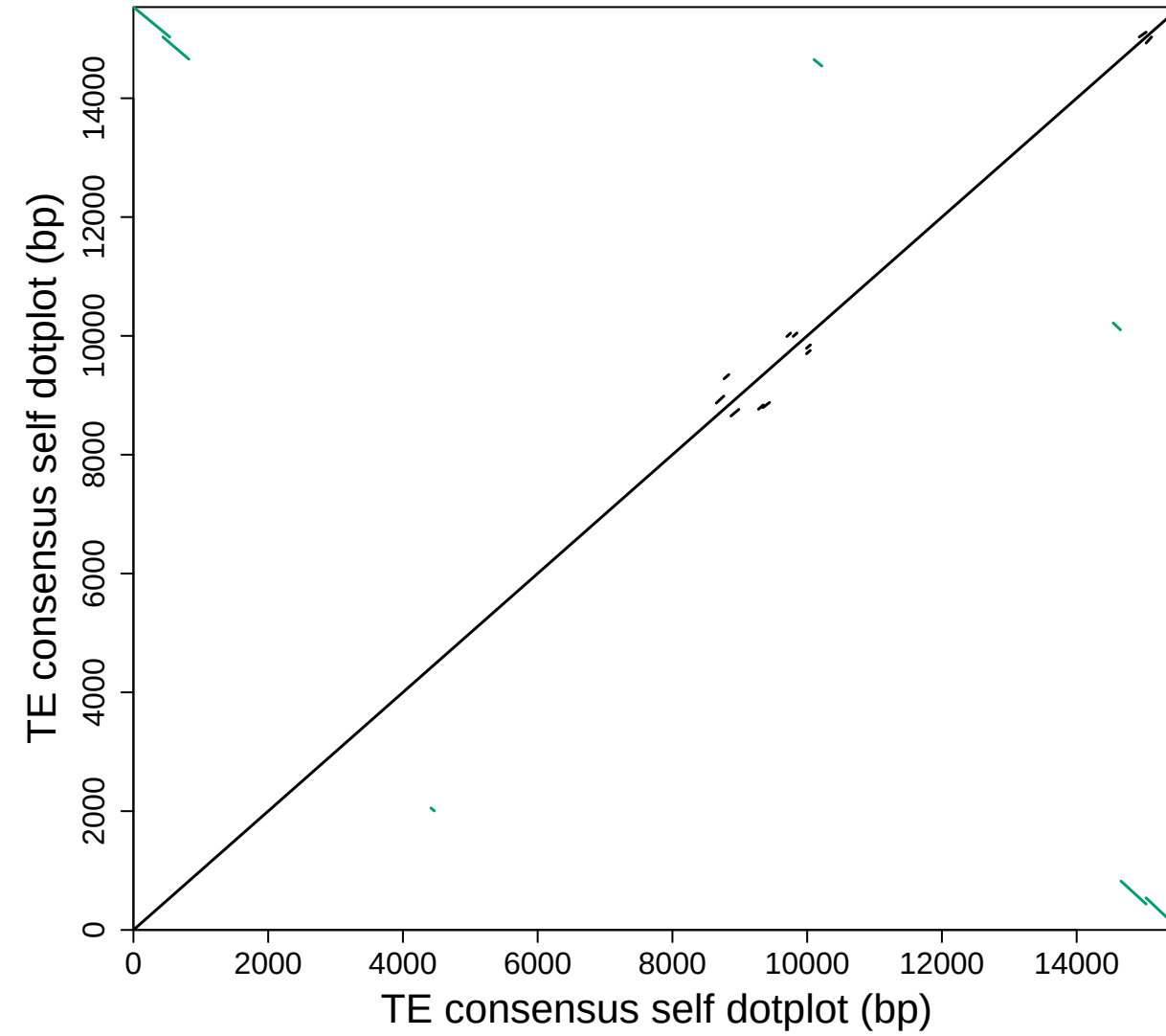
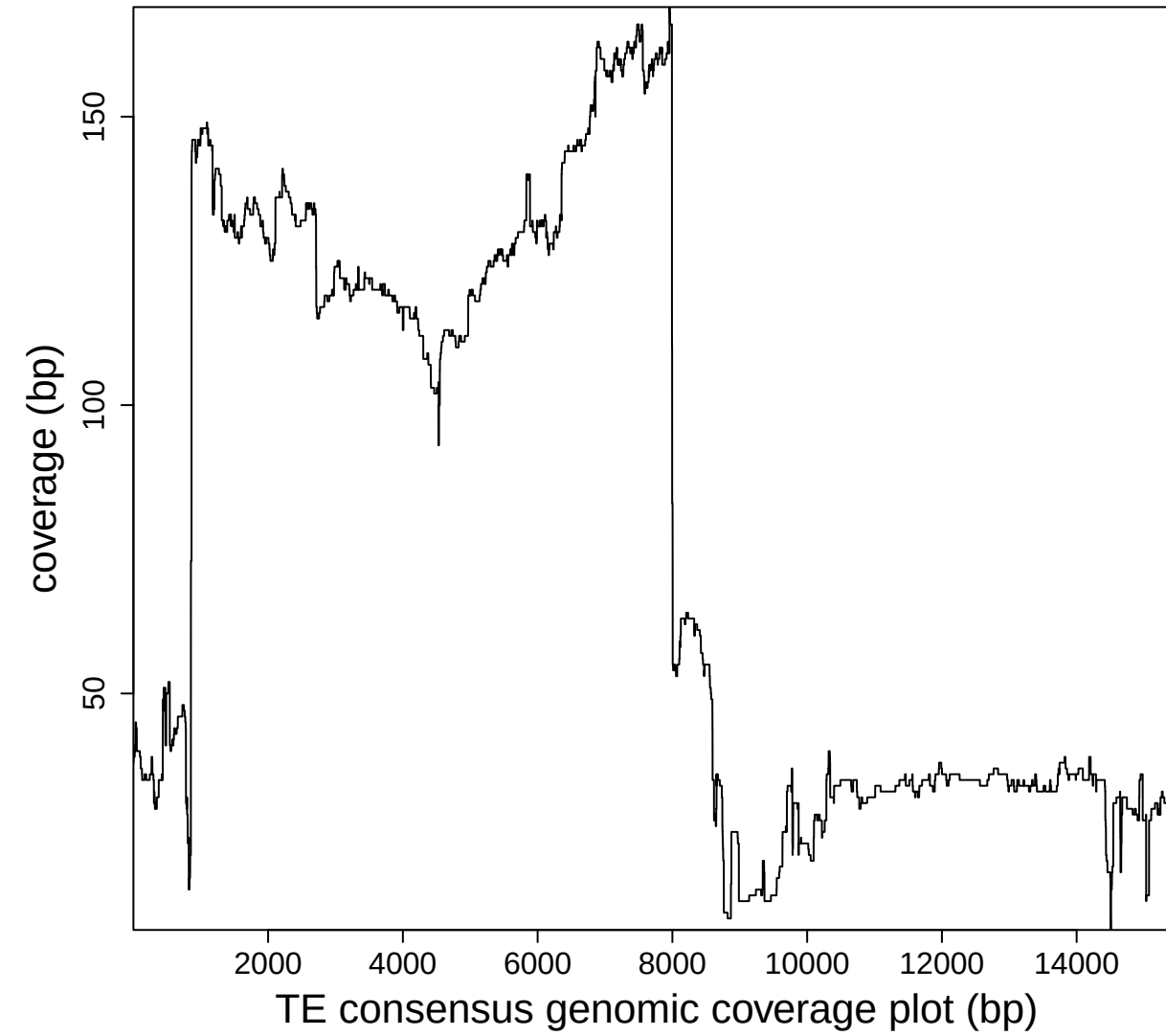
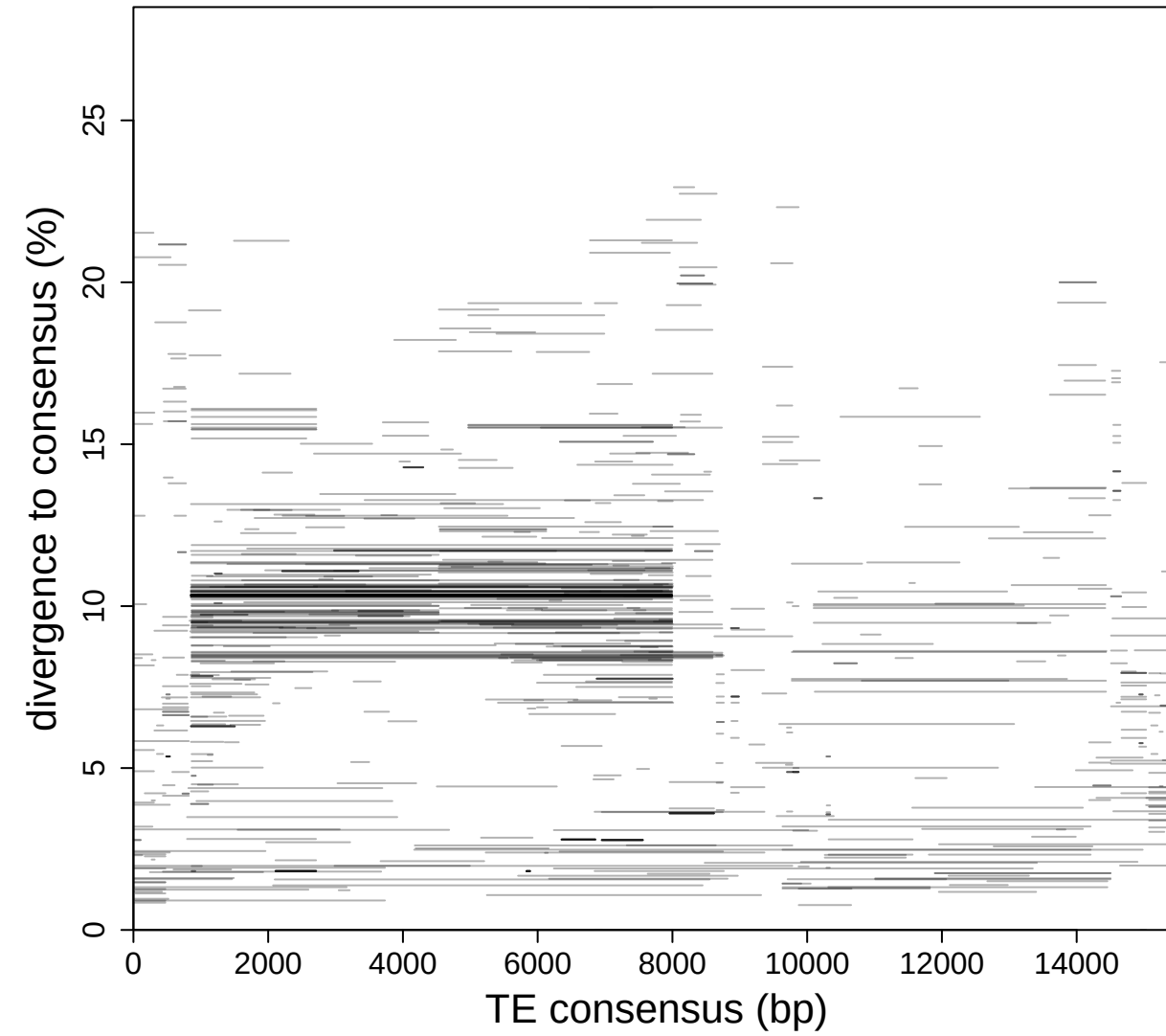
MSA length = 15534





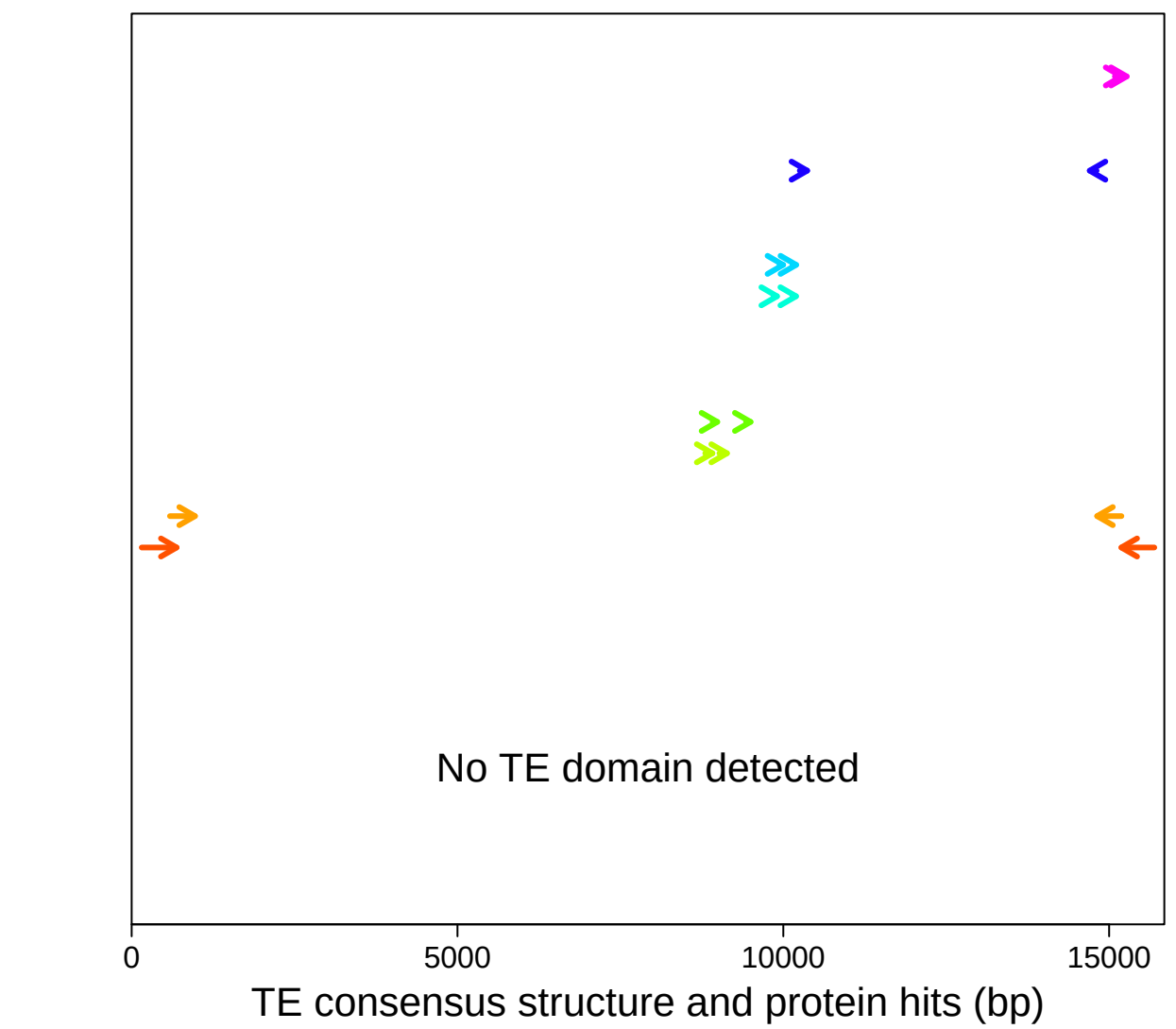
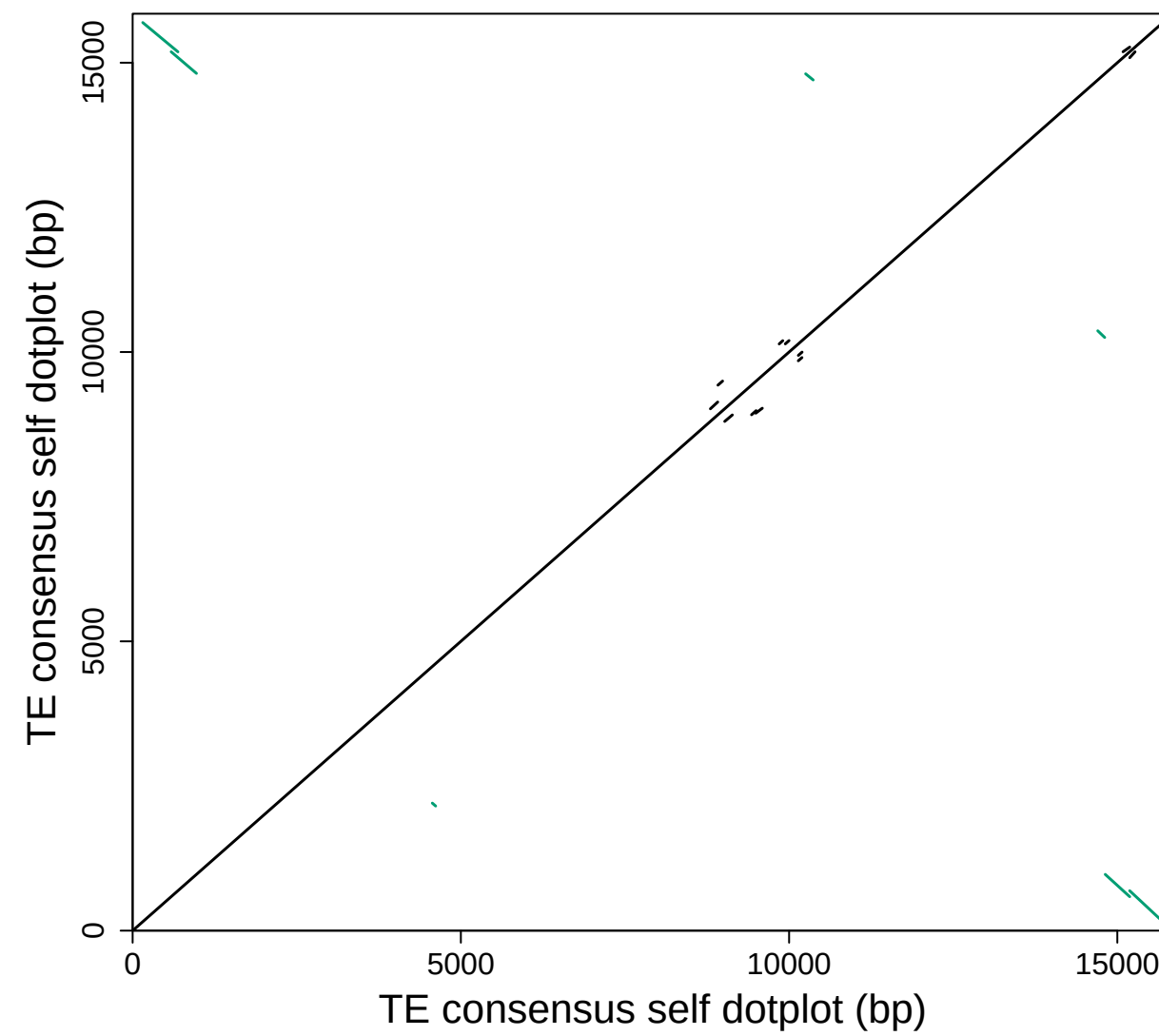
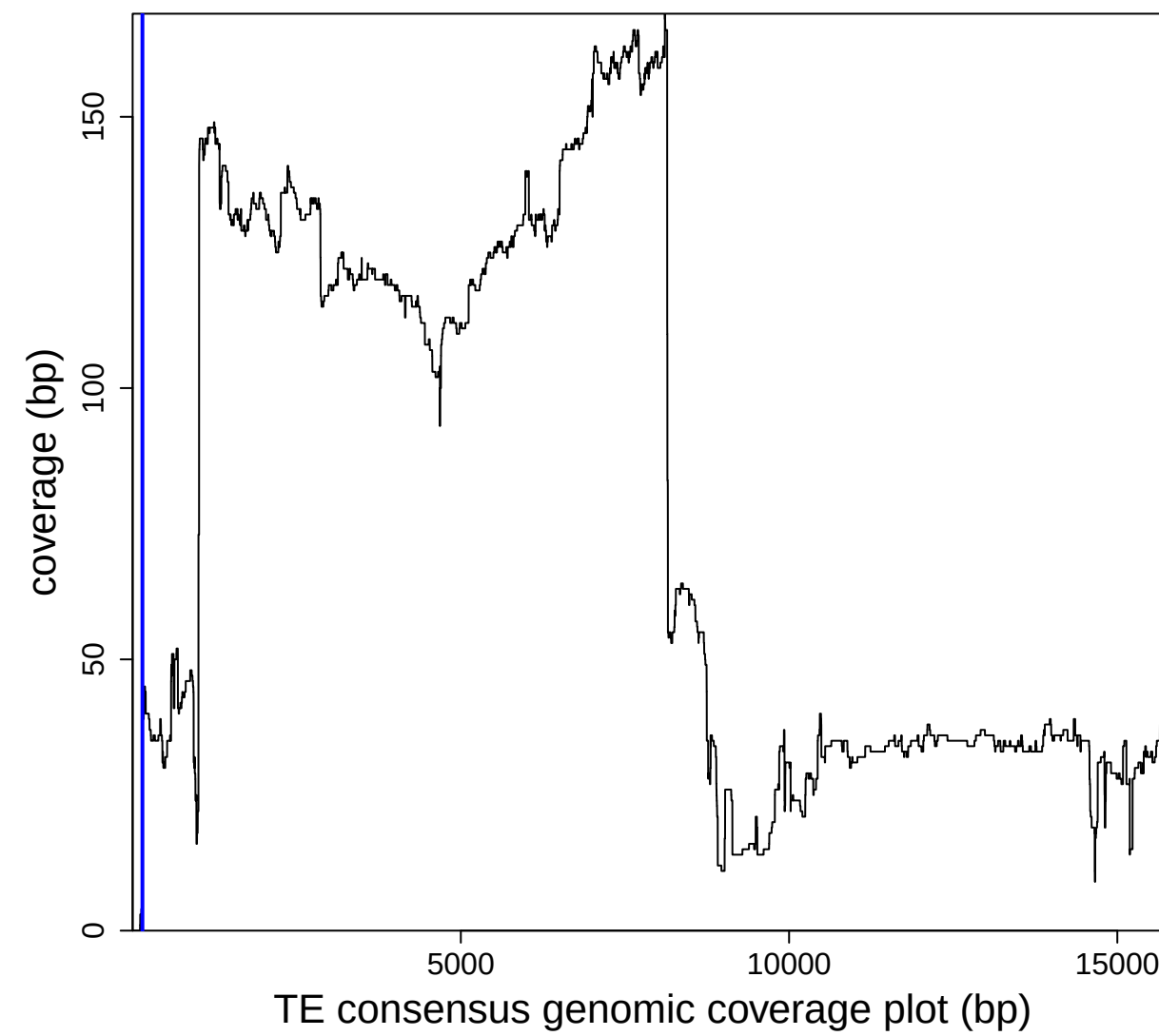
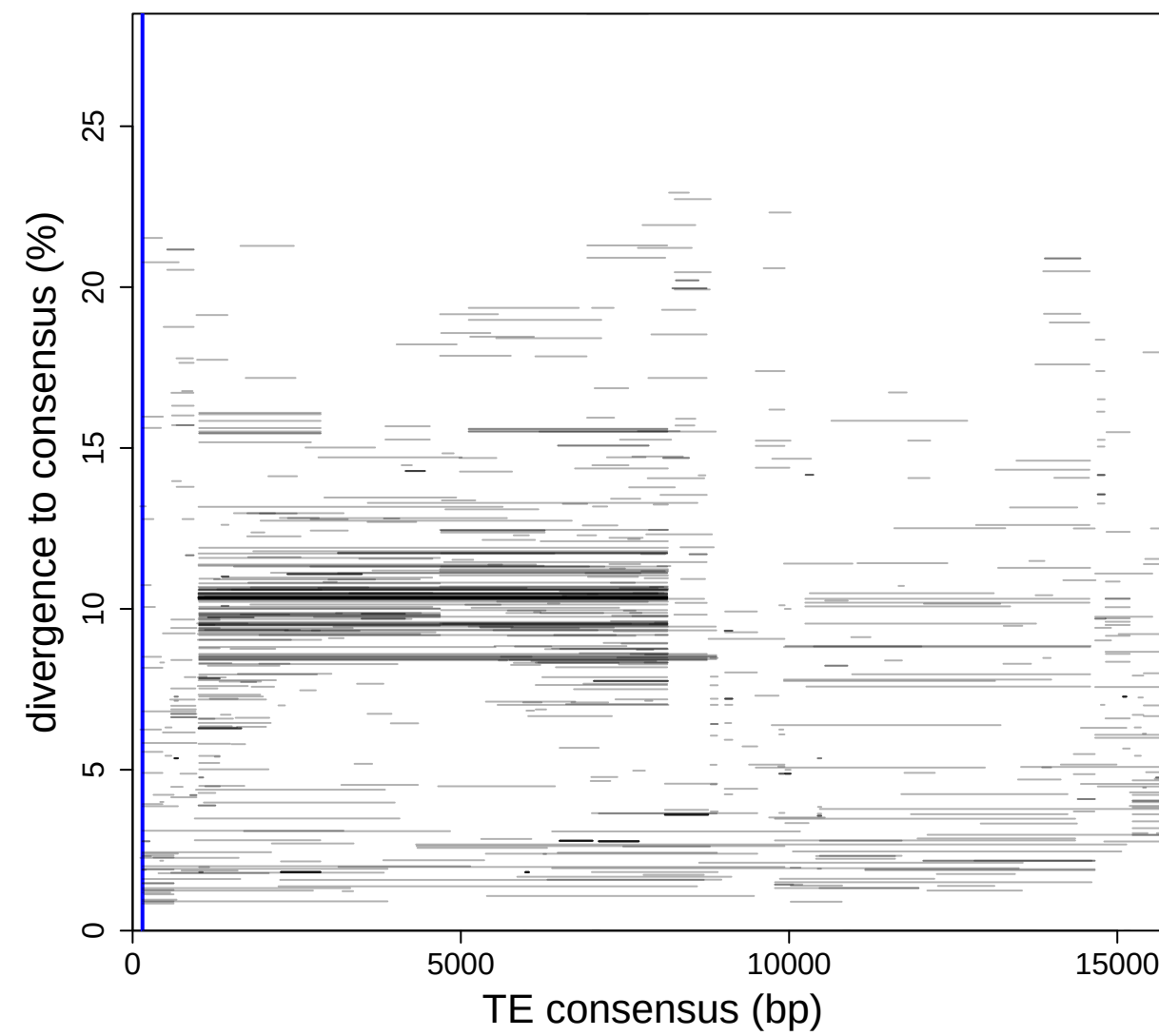
2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_aln.f
size: 15534bp; fragments: 822; full length: 0 (>=13980.6bp)

After TEtrimmer 15534 bp



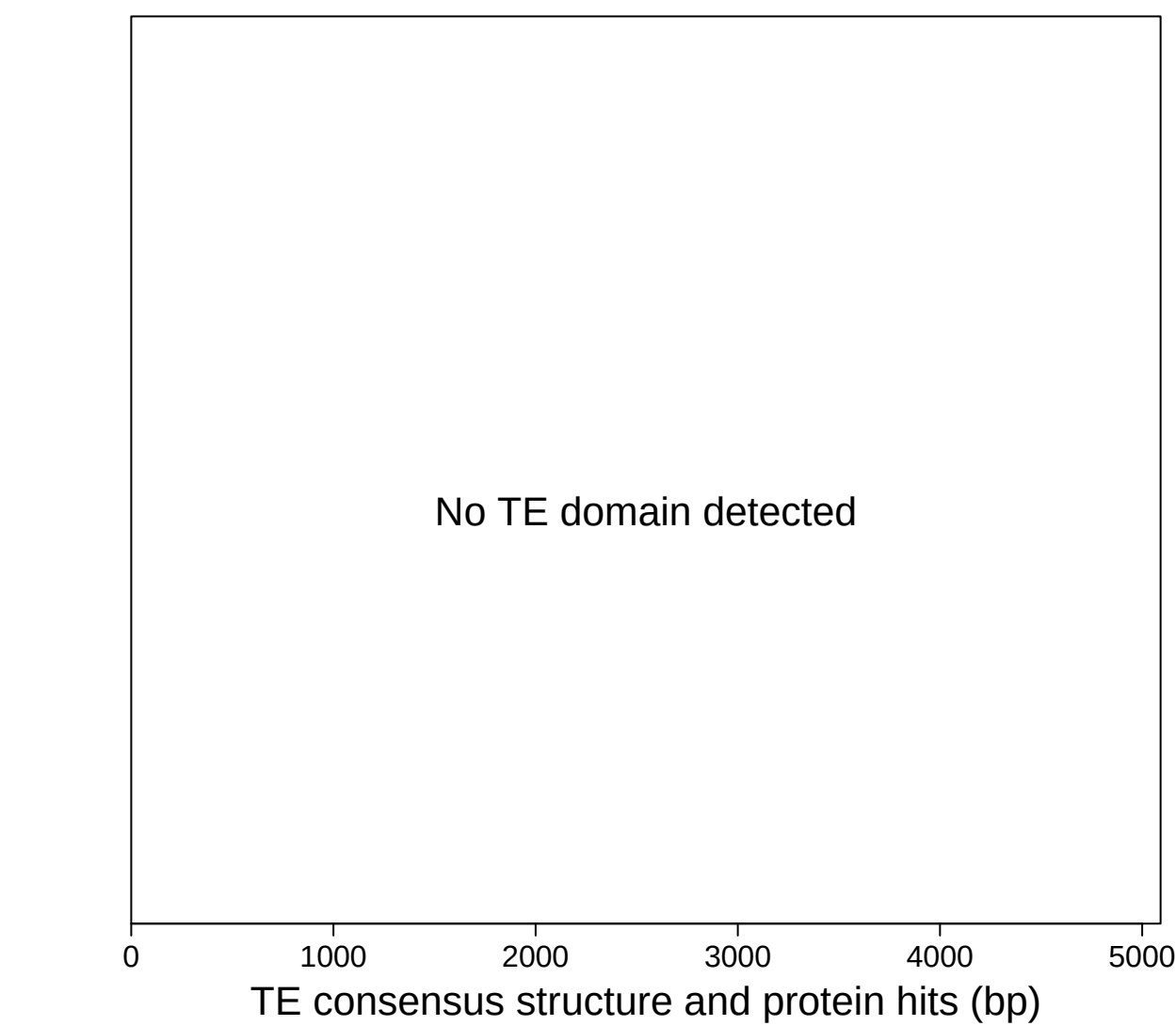
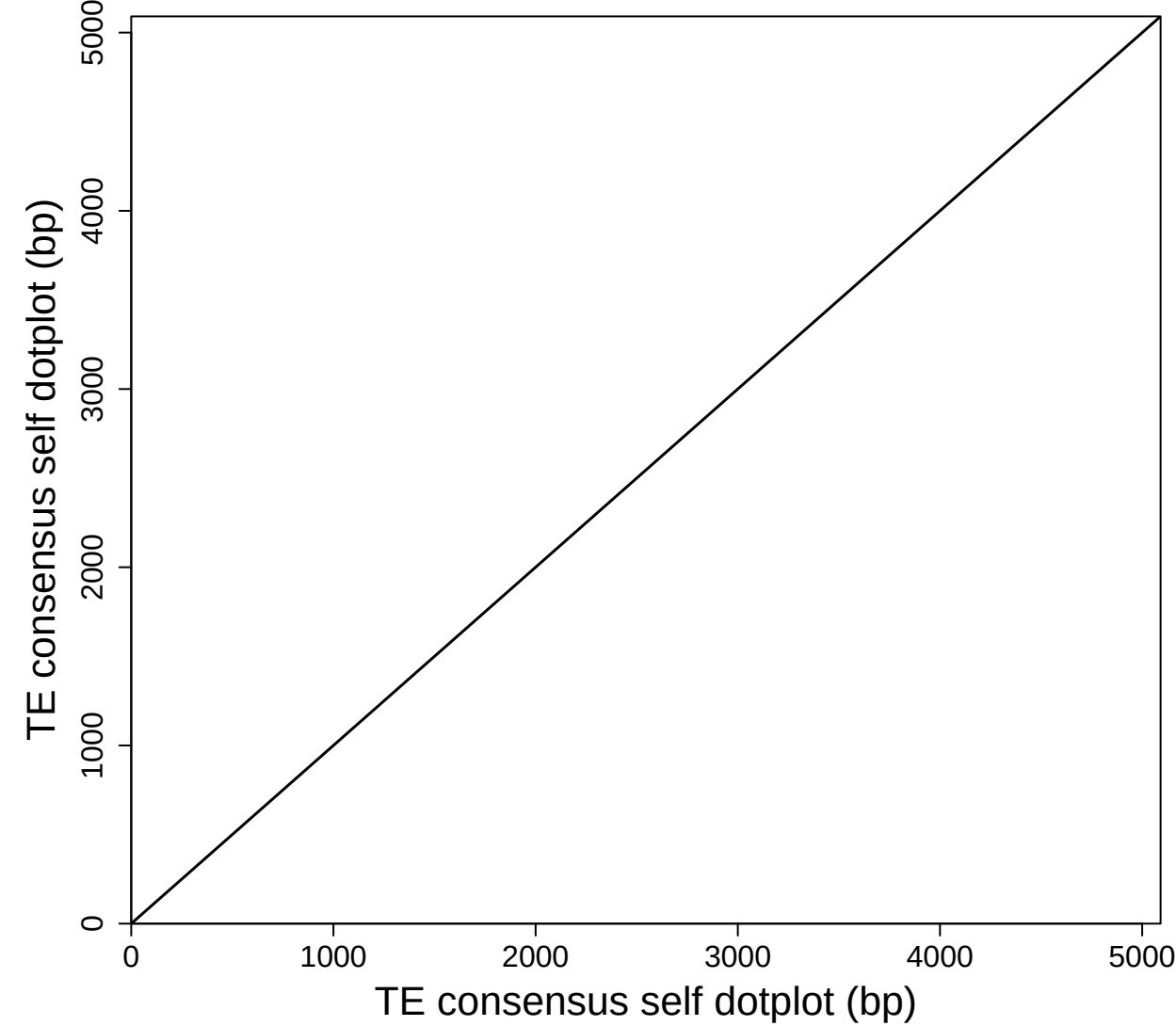
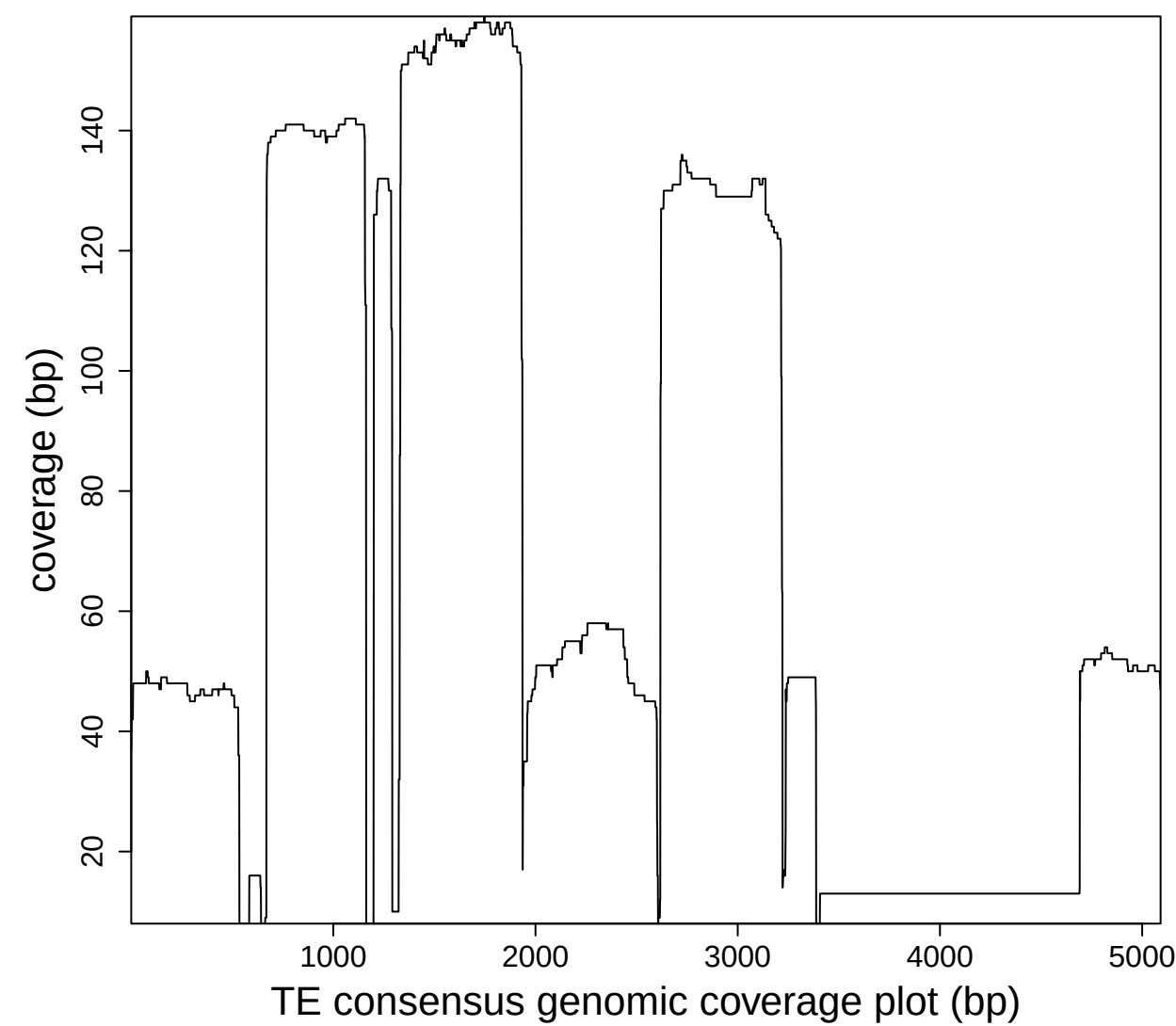
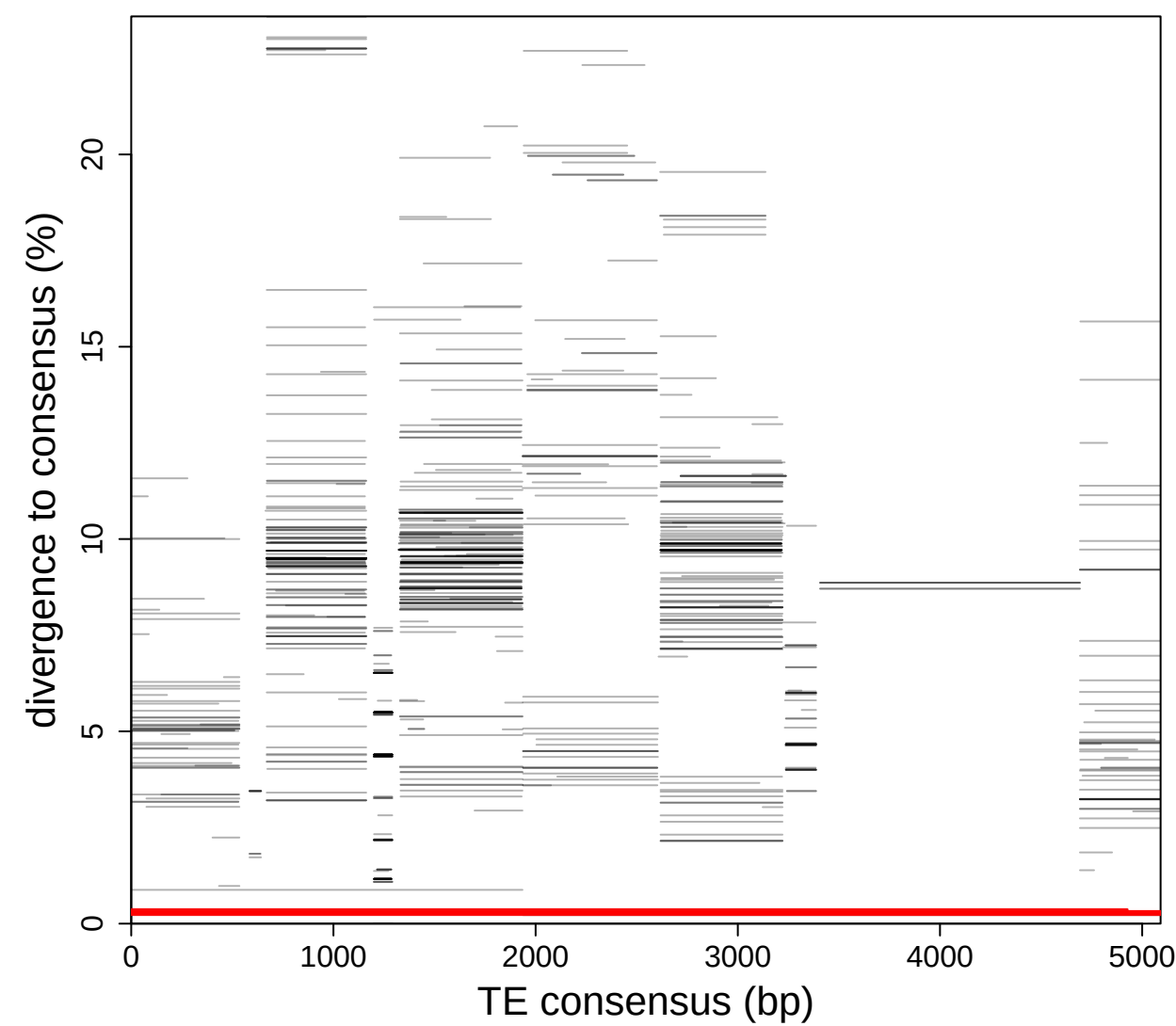
bed_uf.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0
size: 15847bp; fragments: 822; full length: 0 (>=14262.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

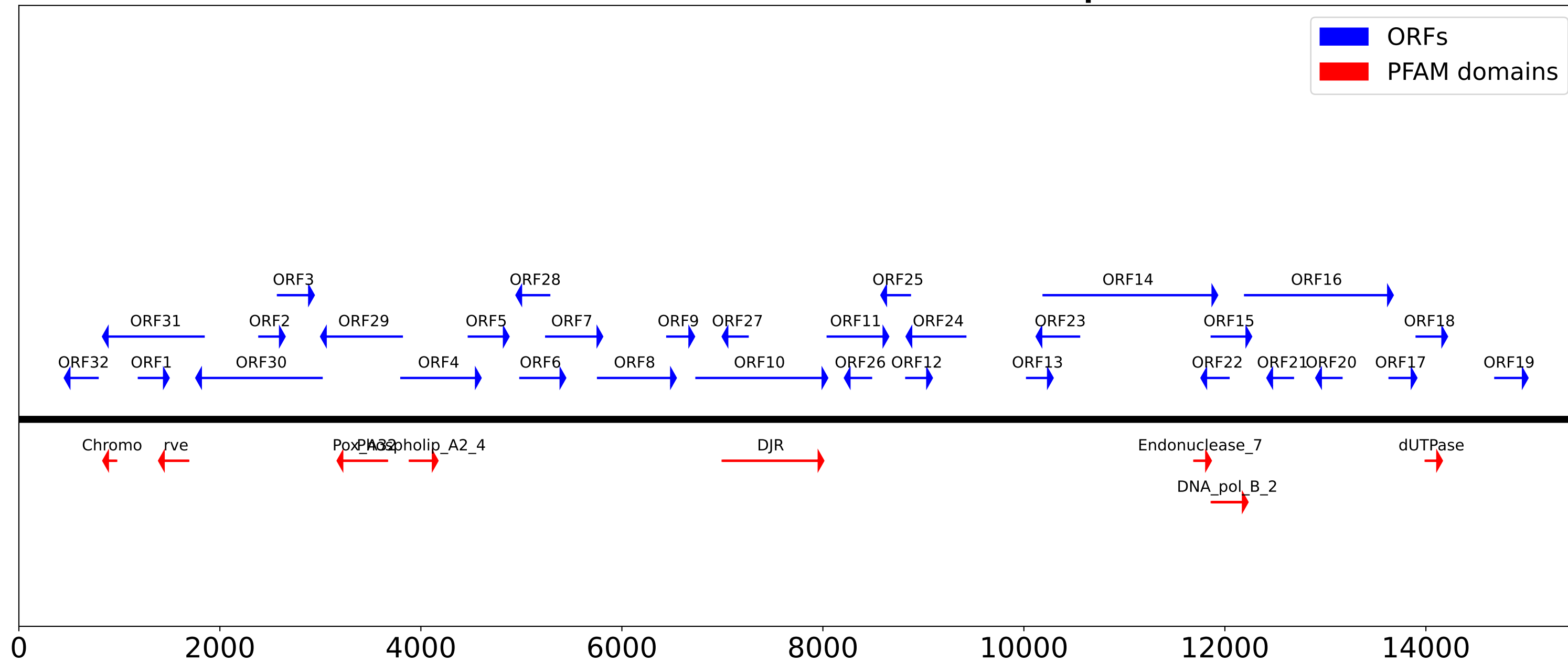


TE: rnd_5_family_7362
size: 5091bp; fragments: 781; full length: 7 (≥ 4581.9 bp)

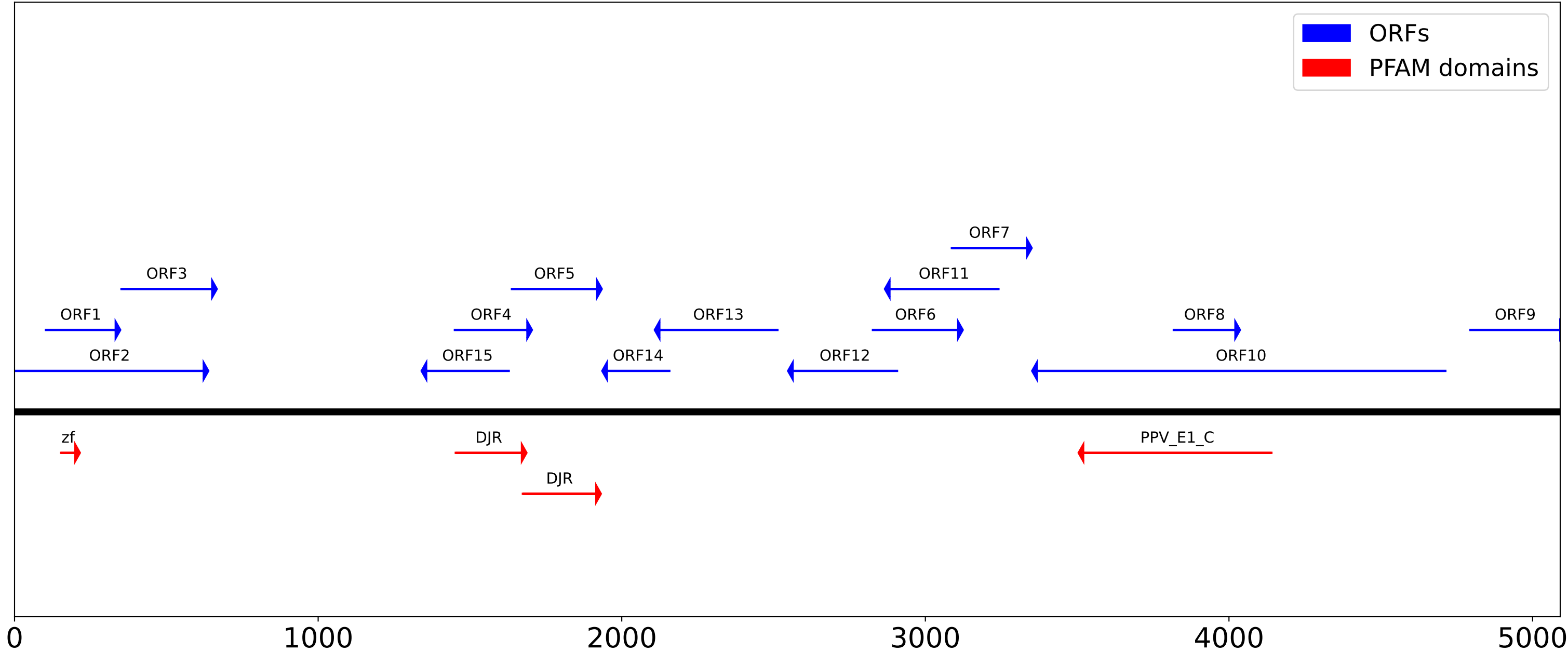
Before TEtrimmer 5091 bp



After TEtrimmer ORF and PFAM domain plot



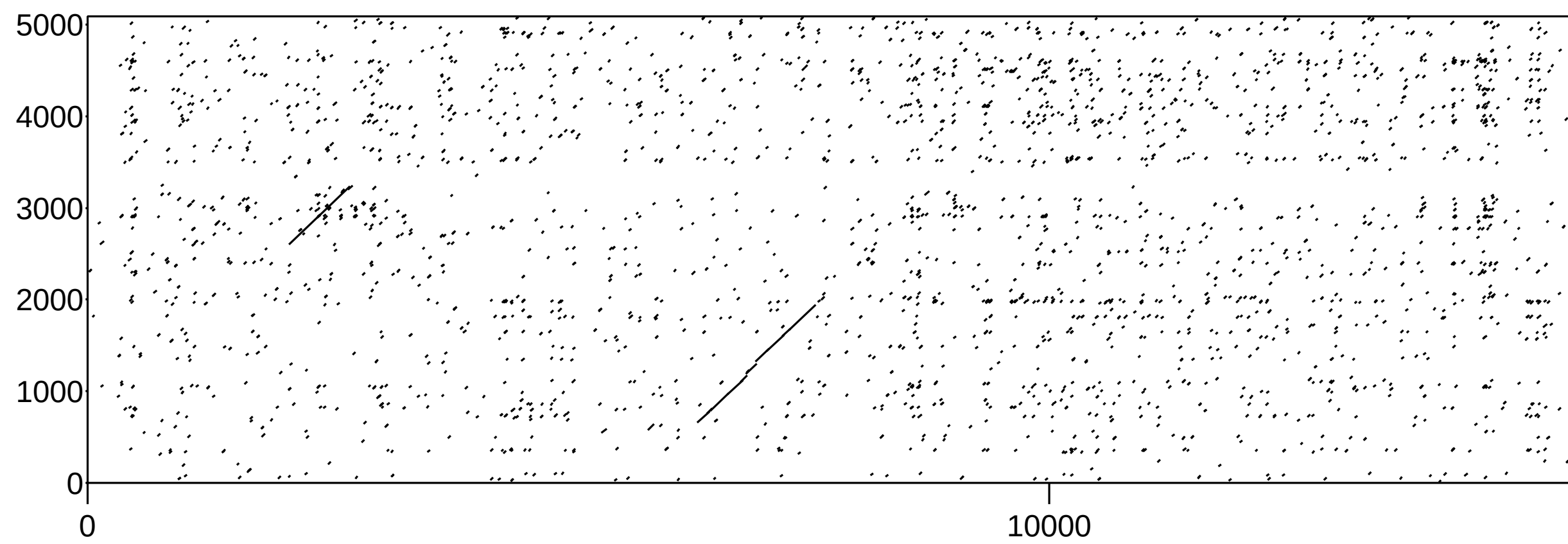
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)