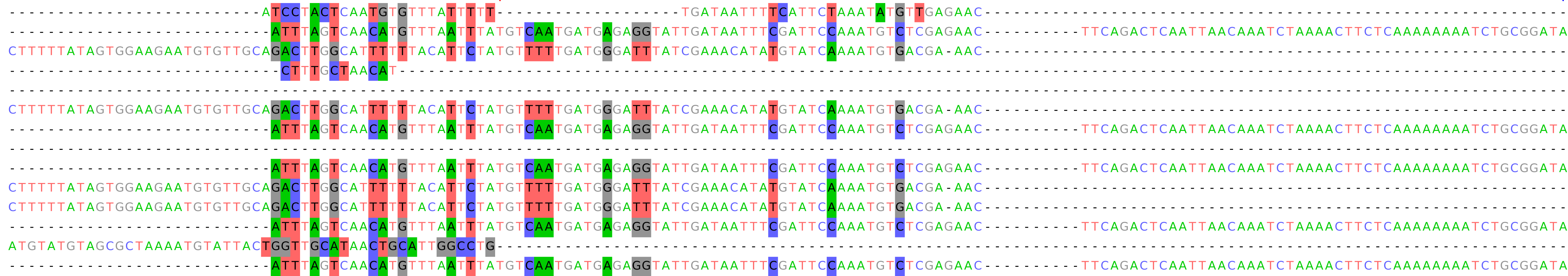
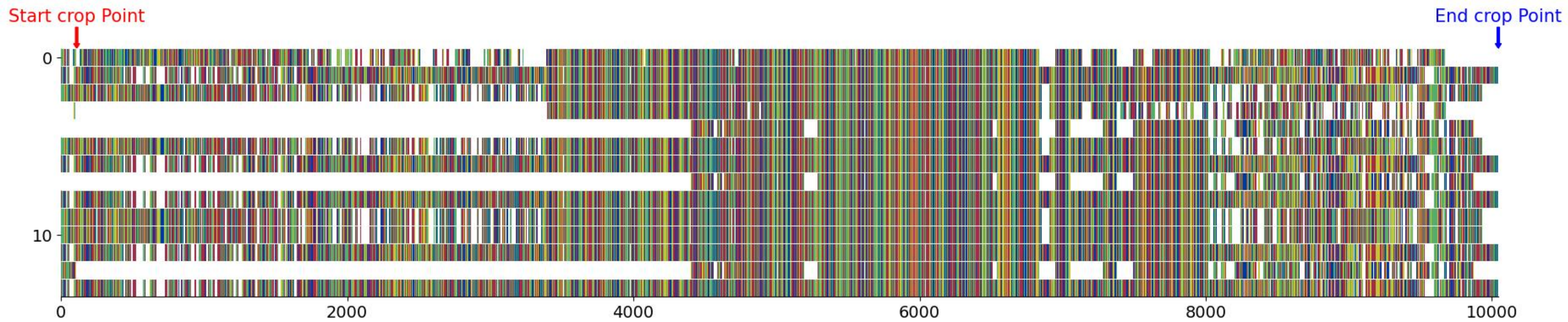


Start crop Point

End crop Point

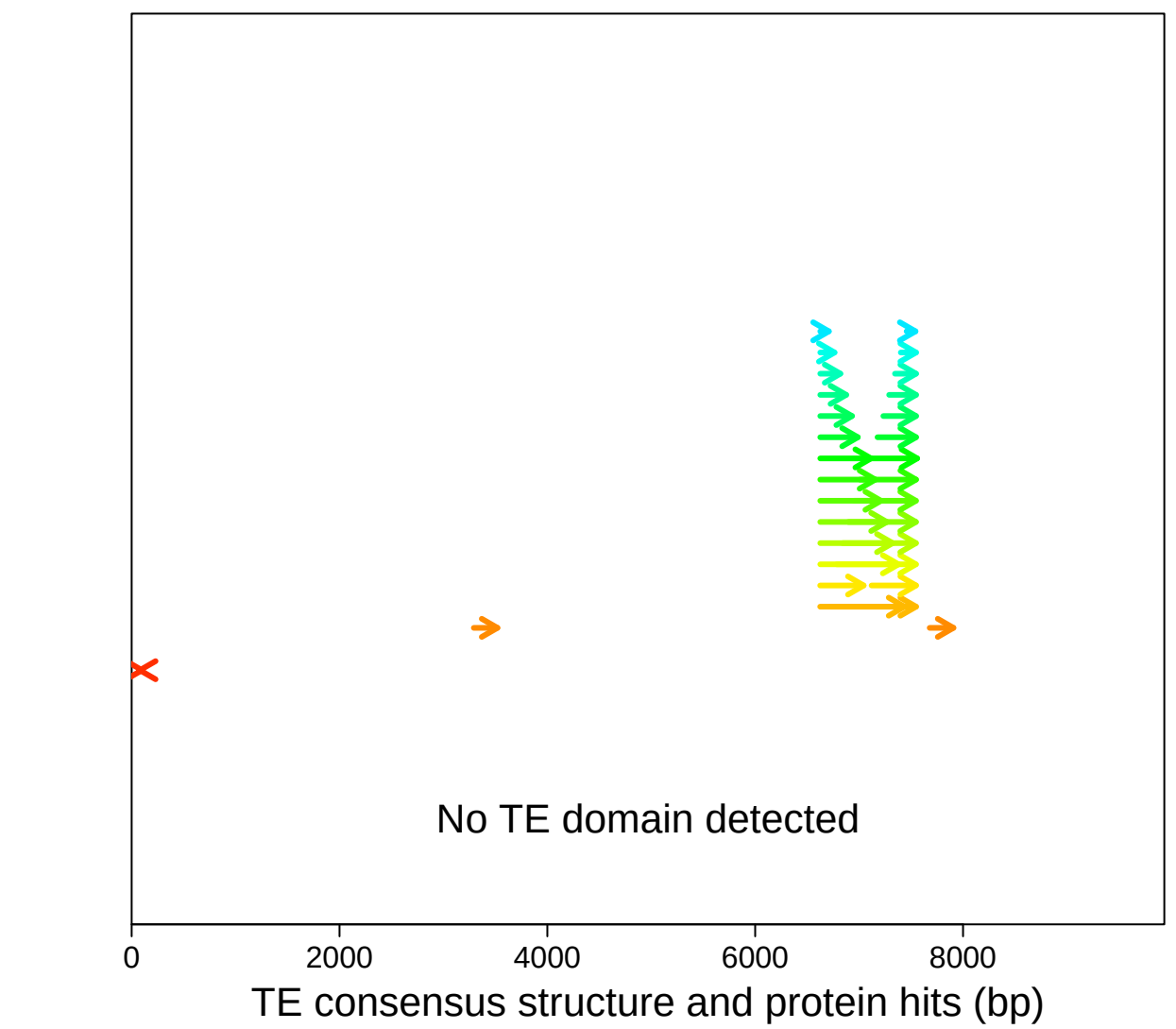
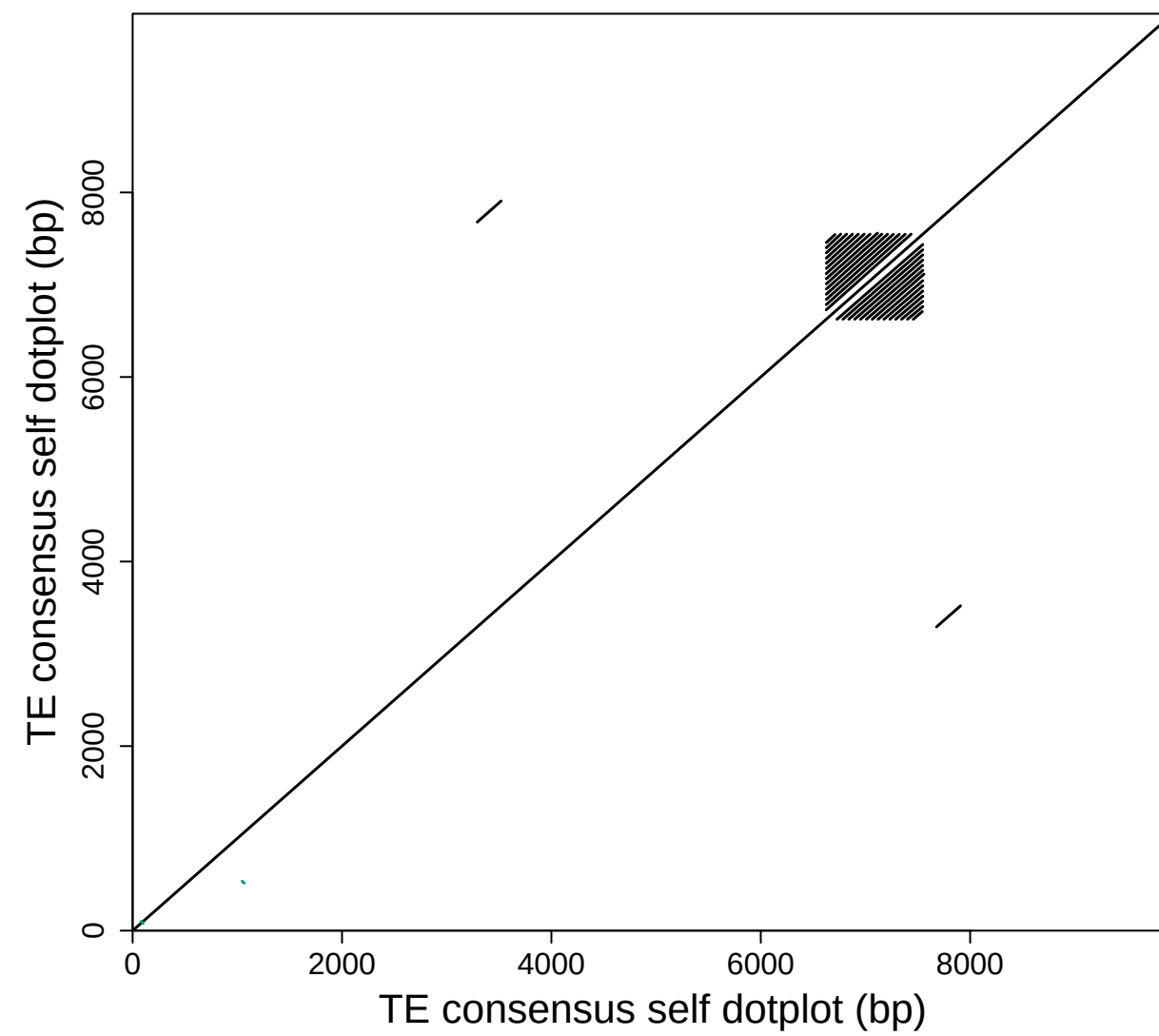
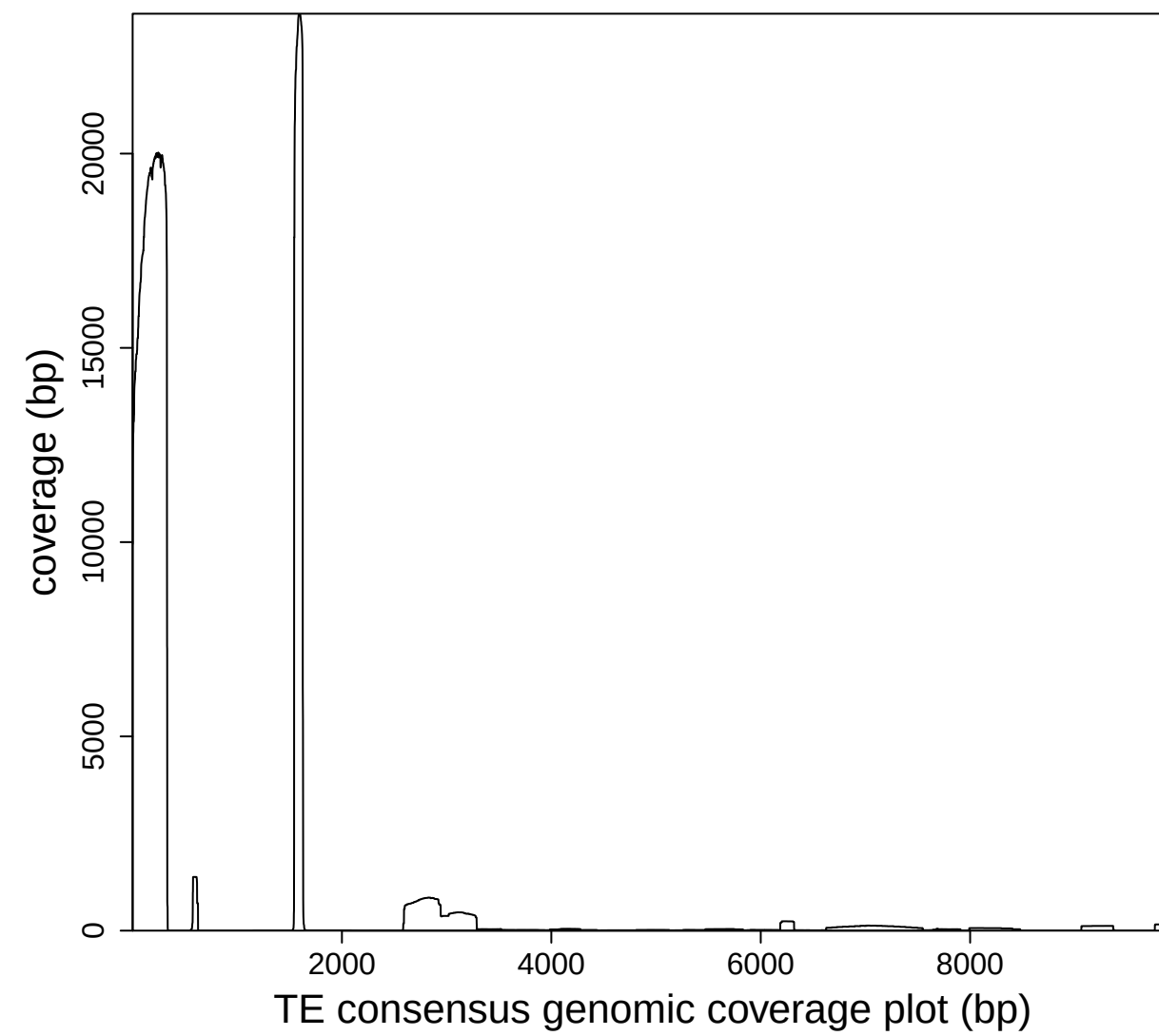
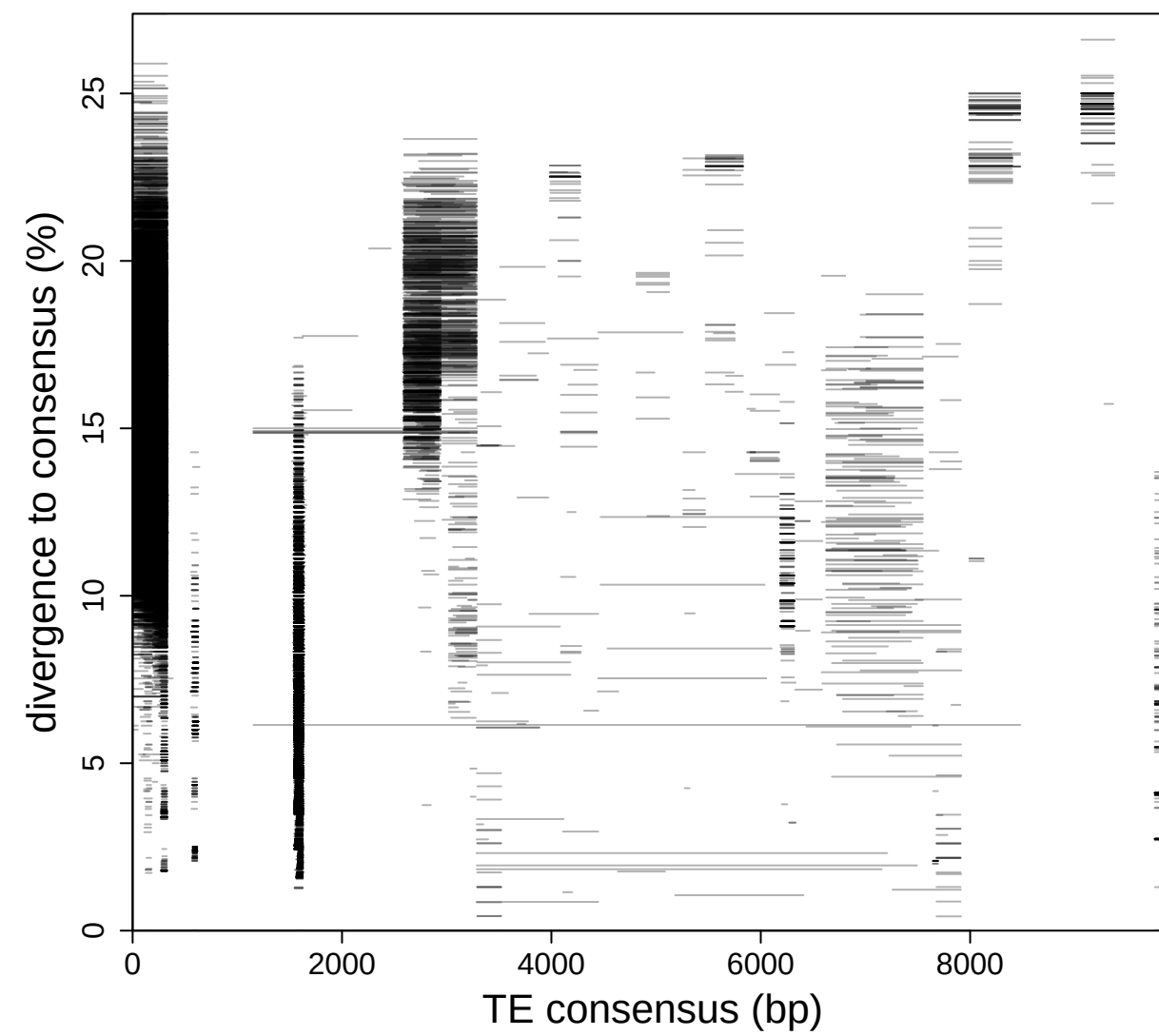
MSA length = 9937

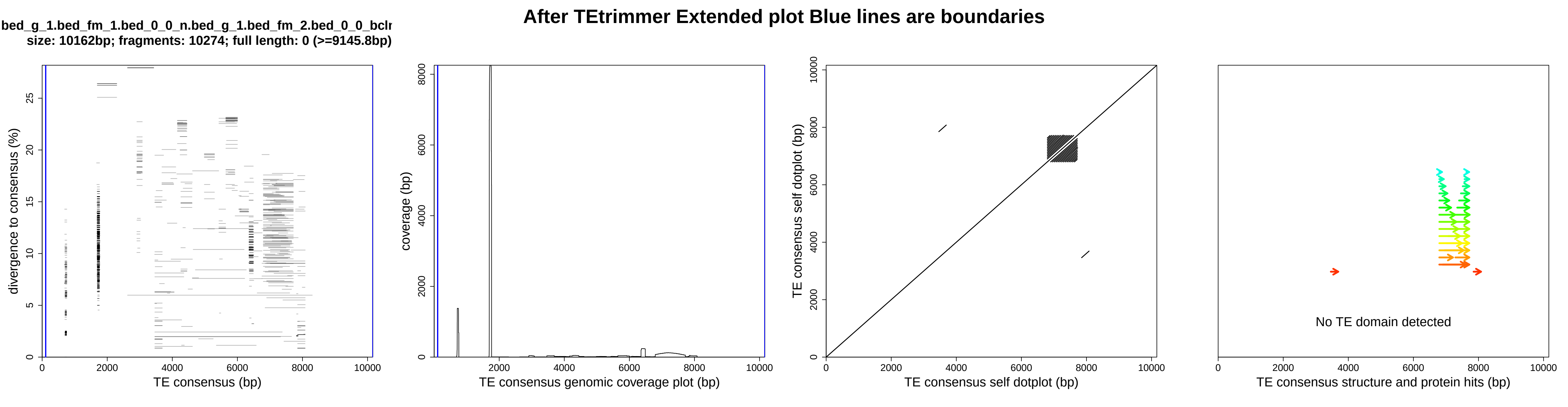




_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa al
size: 9937bp; fragments: 48534; full length: 0 (>=8943.3bp)

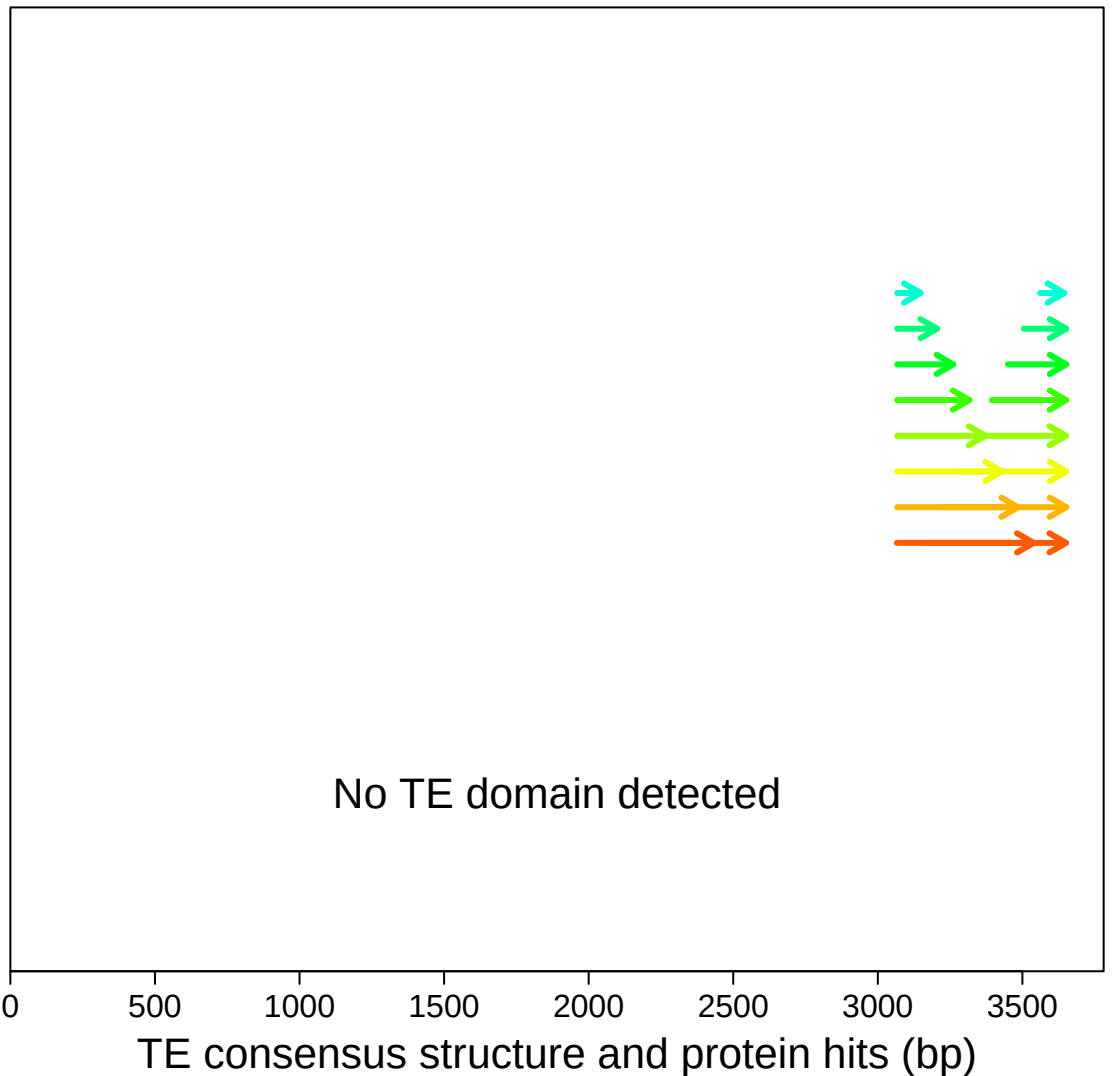
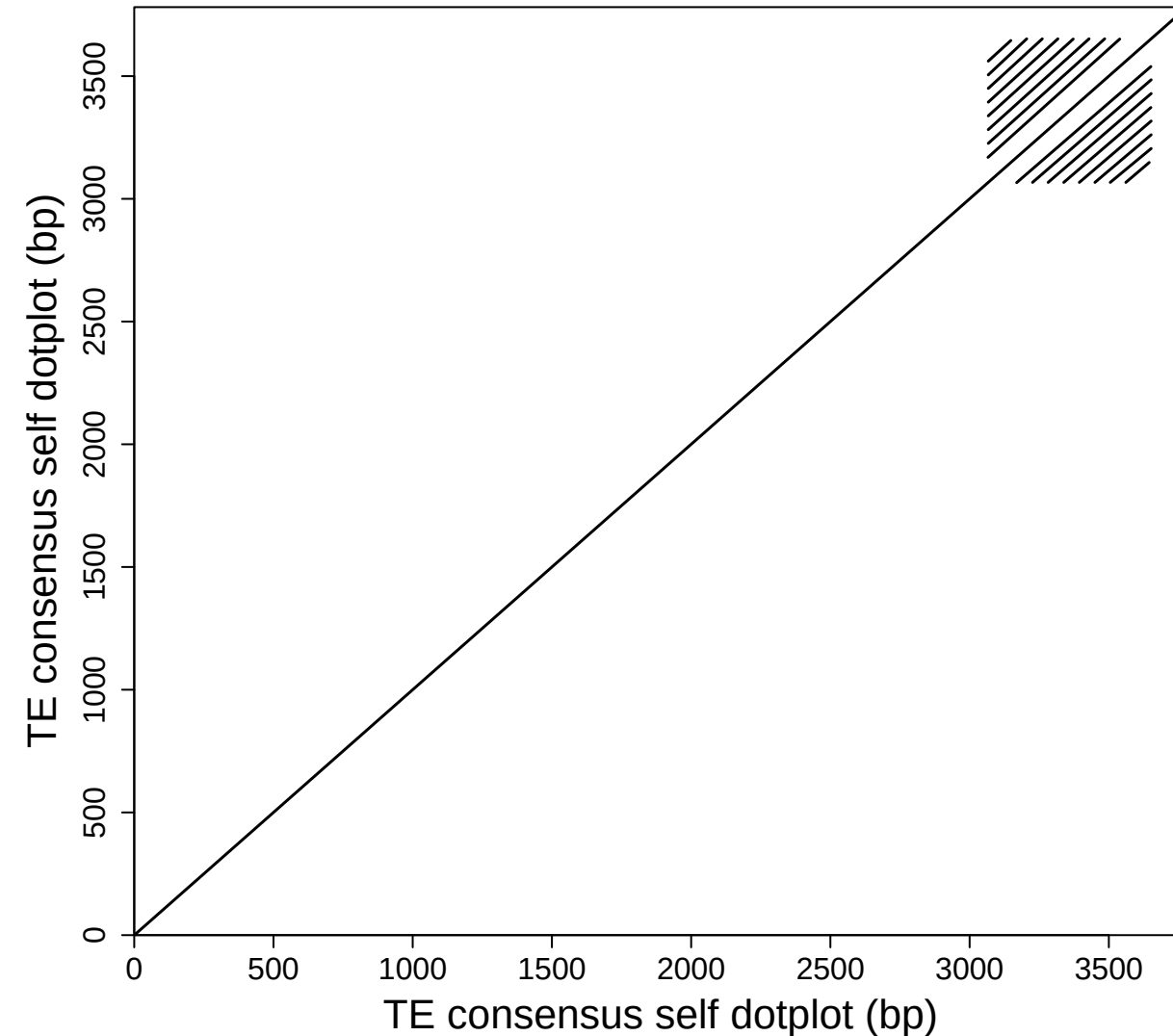
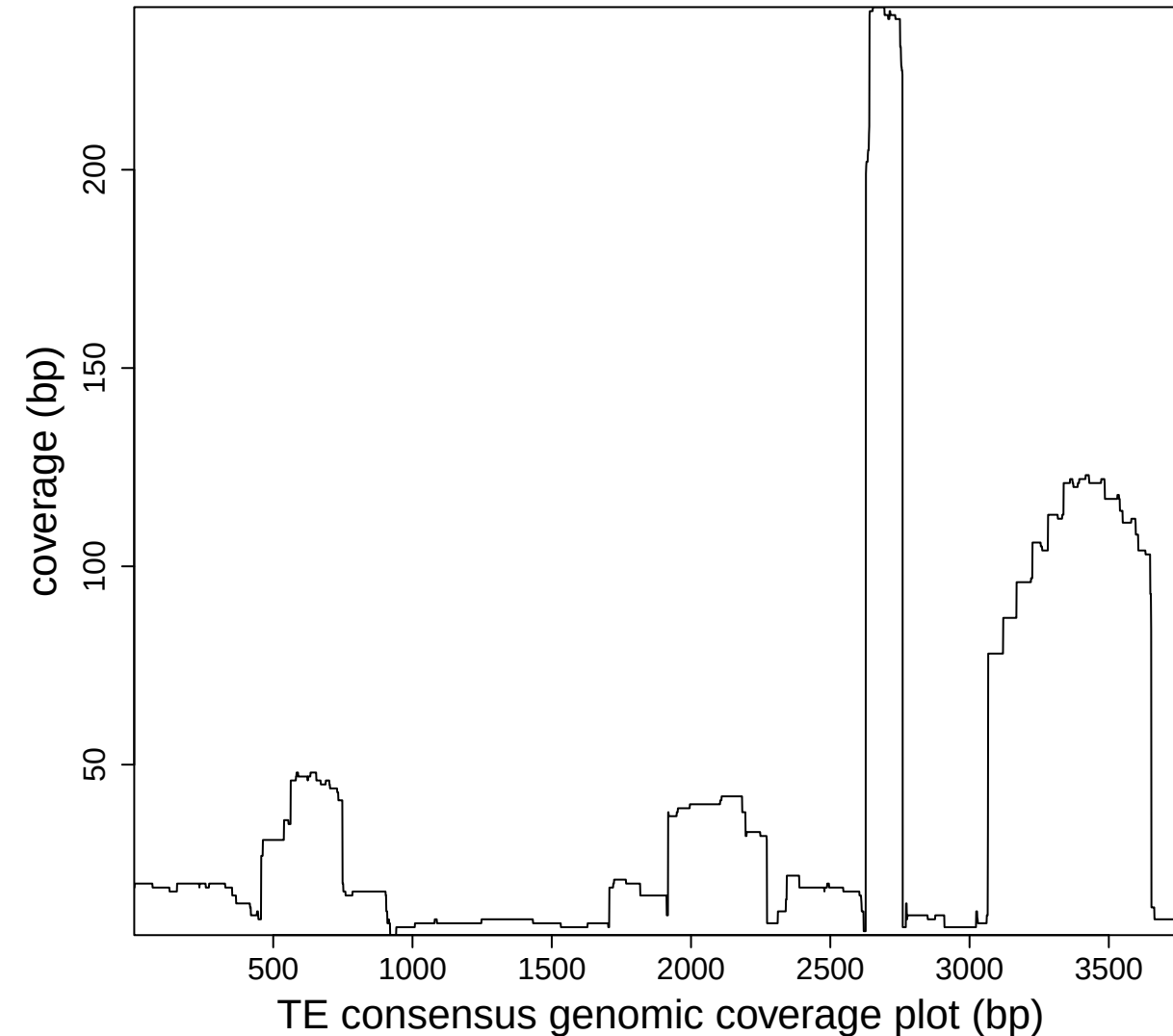
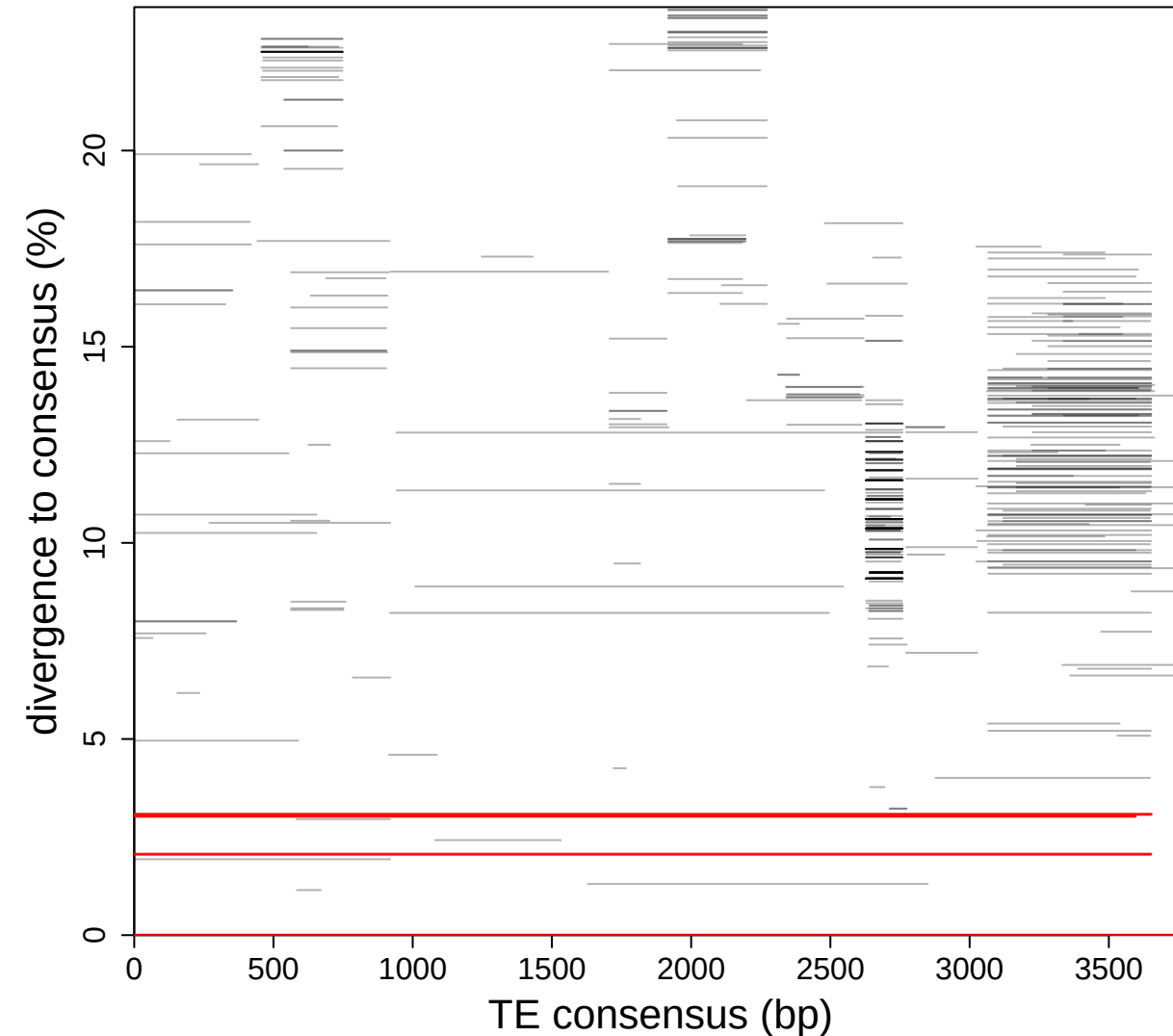
After TEtrimmer 9937 bp



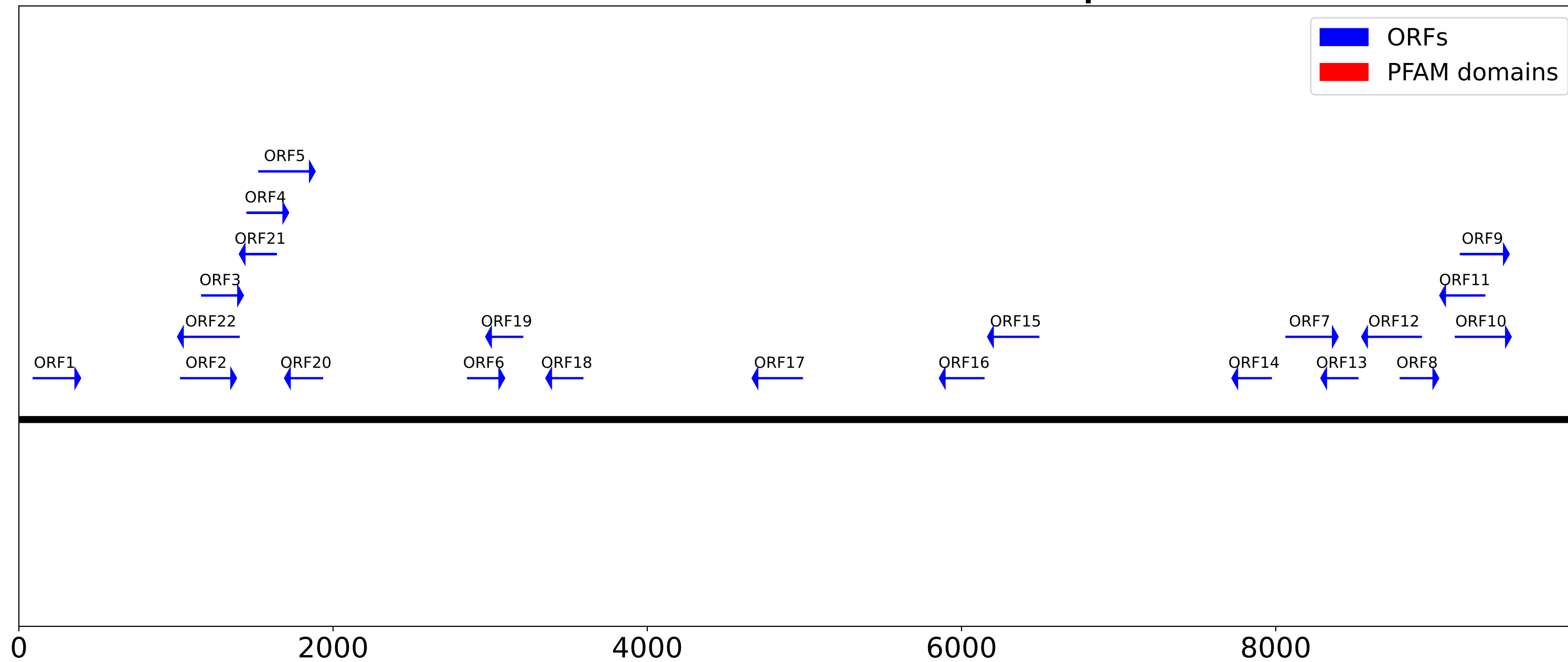


TE: ltr_1_family_591
size: 3781bp; fragments: 503; full length: 4 (>=3402.9bp)

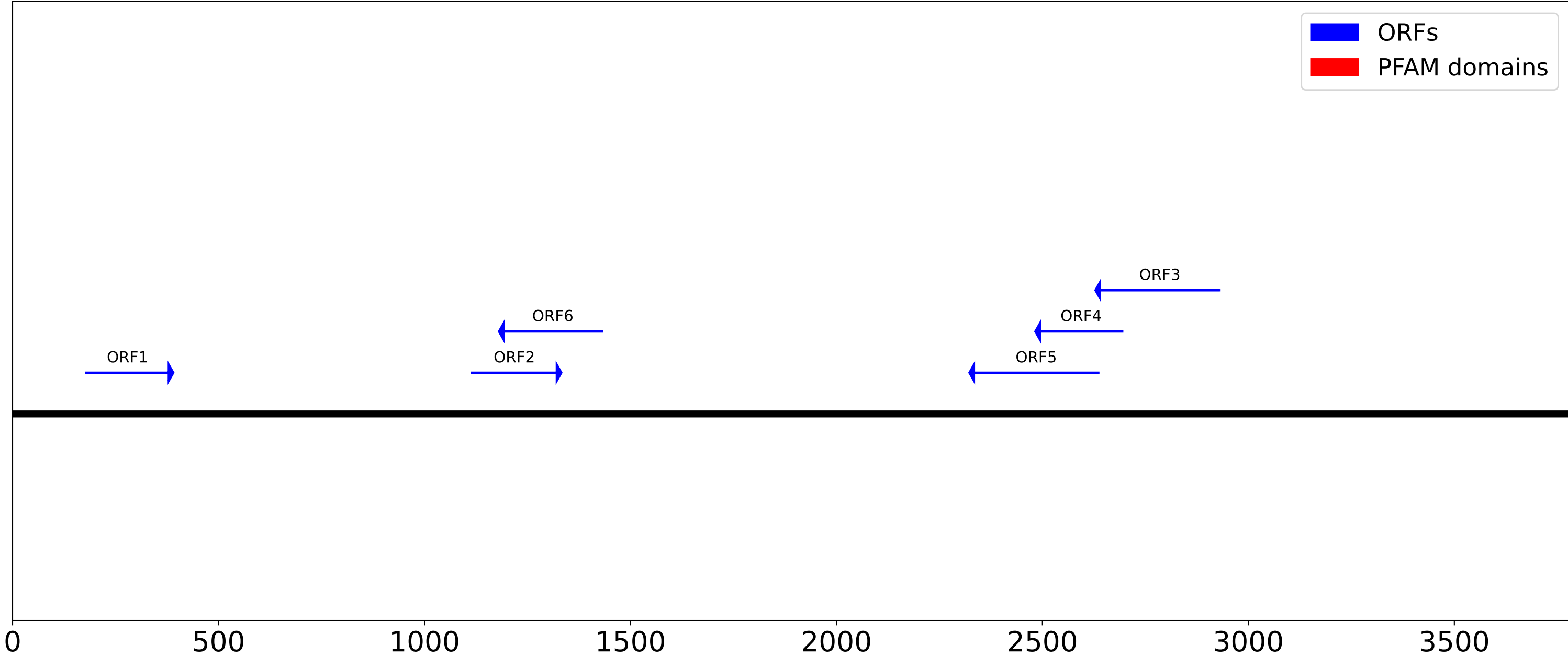
Before TEtrimmer 3781 bp



After TEtrimmer ORF and PFAM domain plot



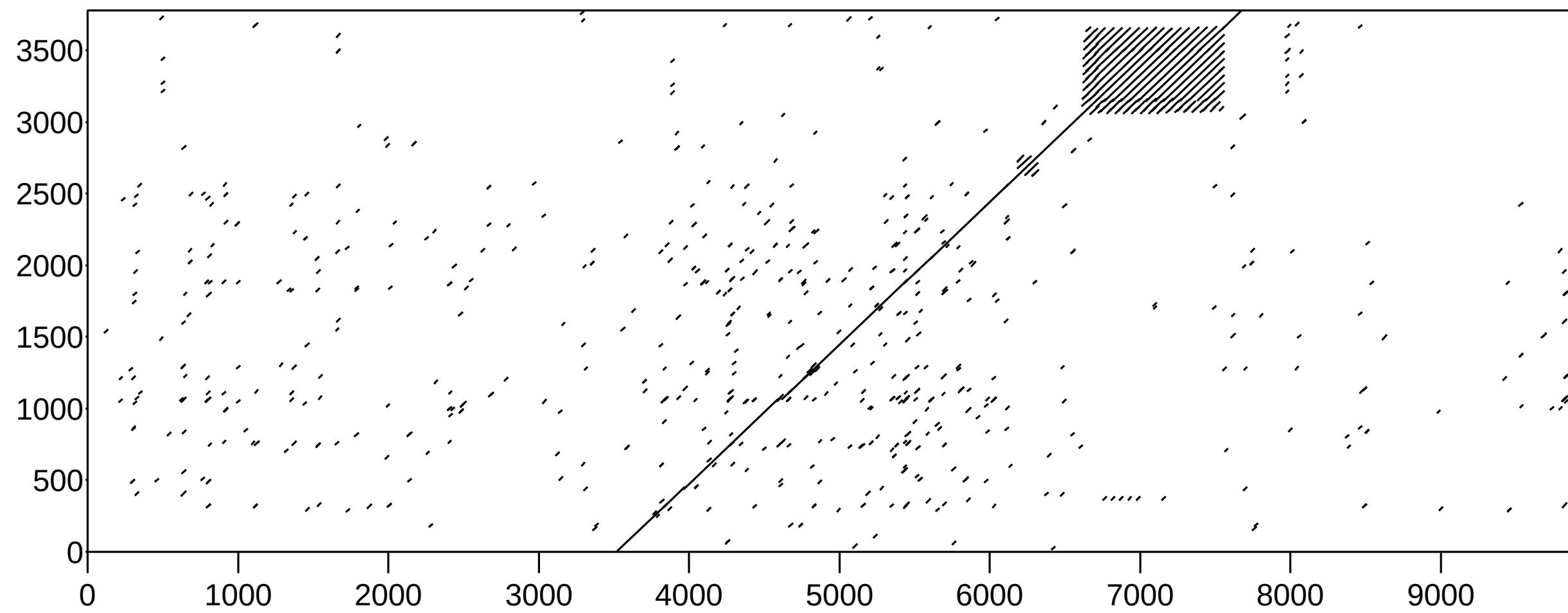
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)