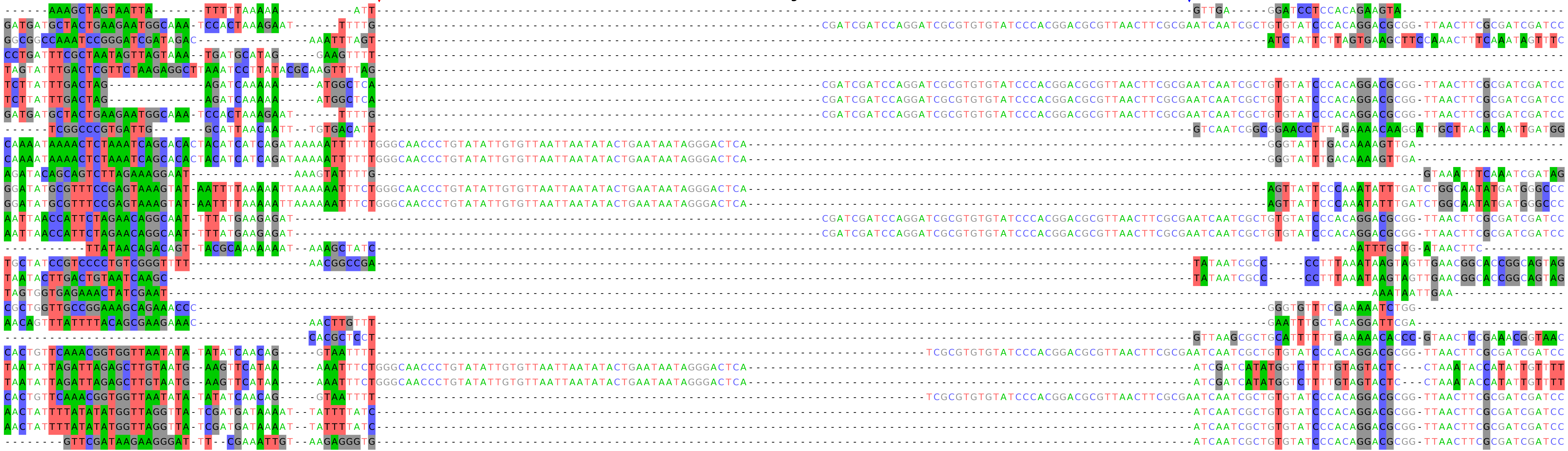
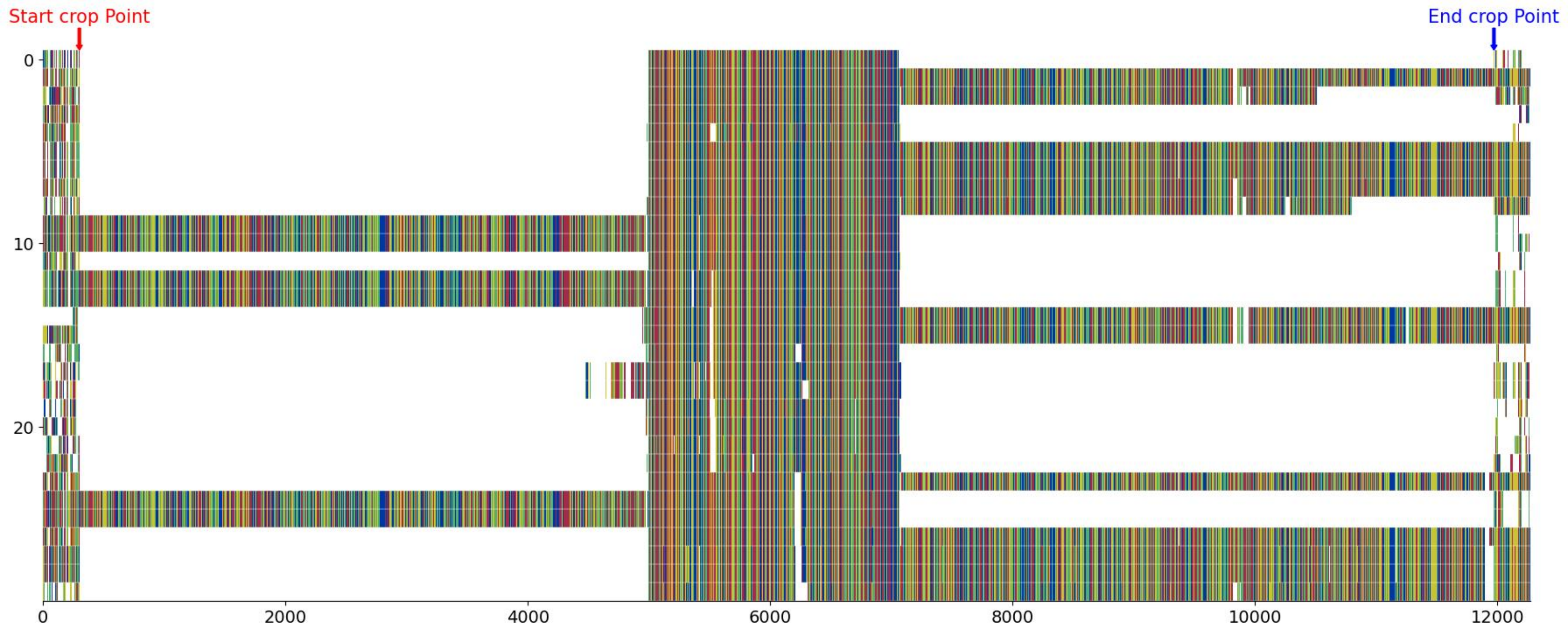


MSA length = 11672

Start crop Point

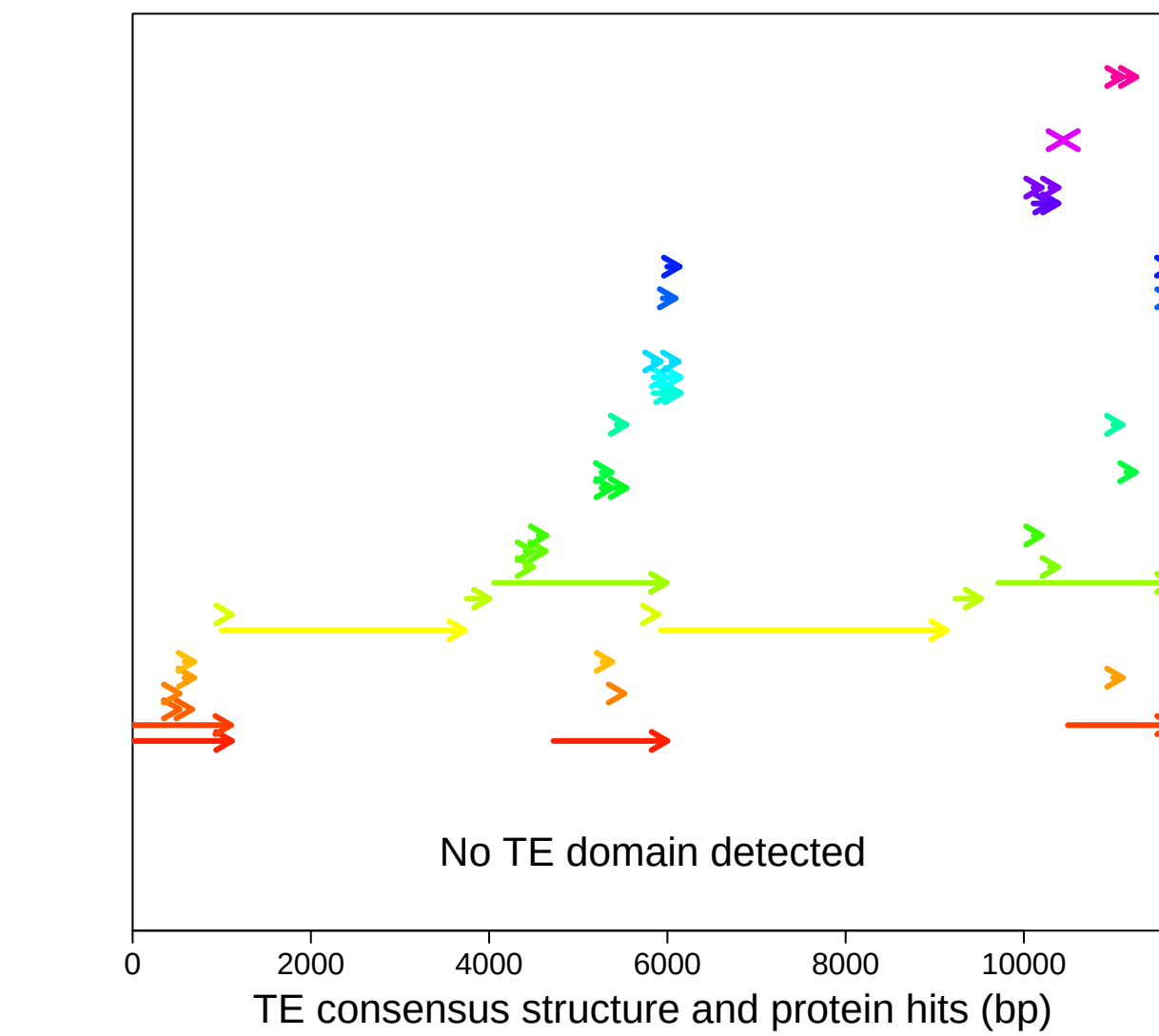
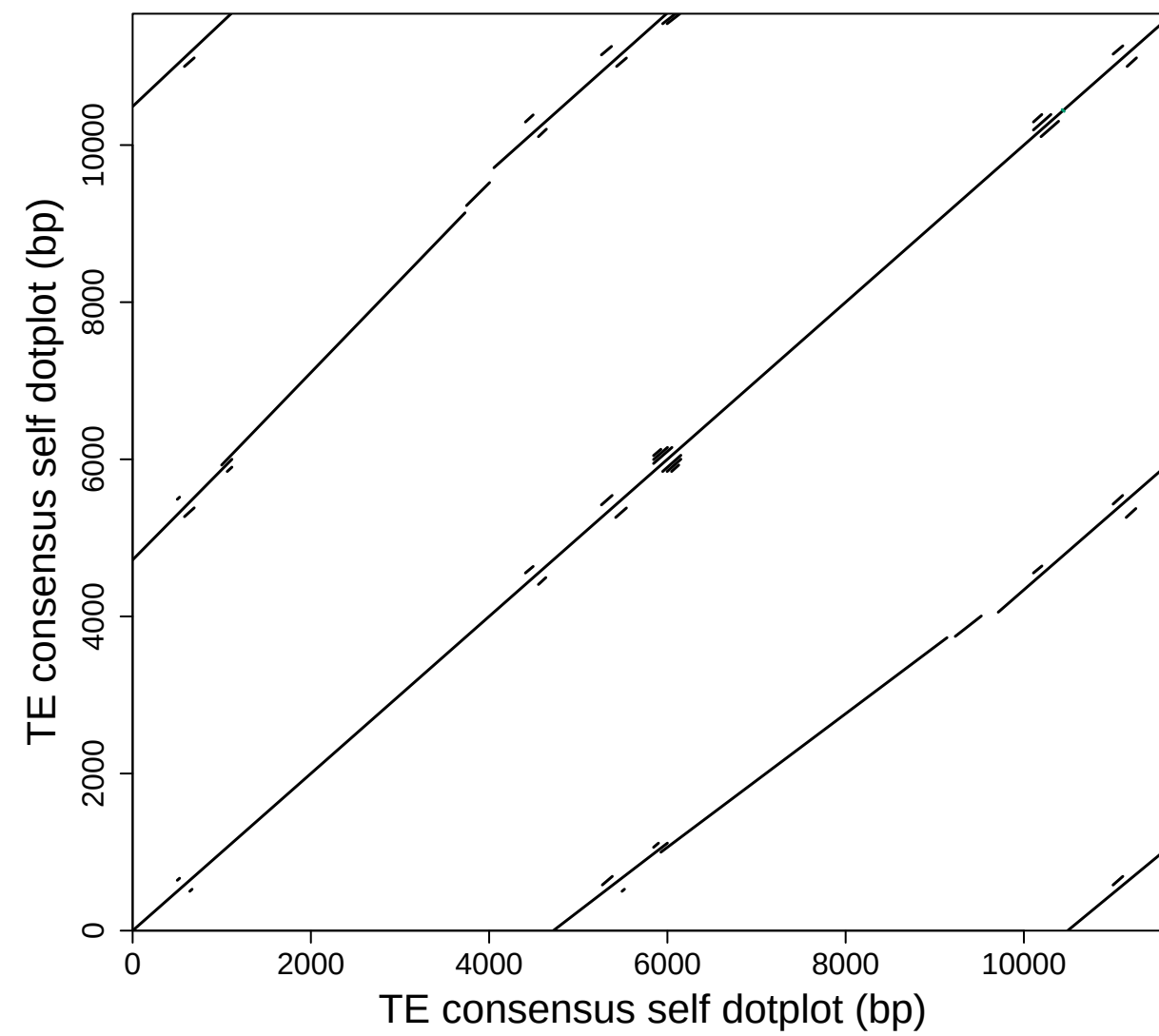
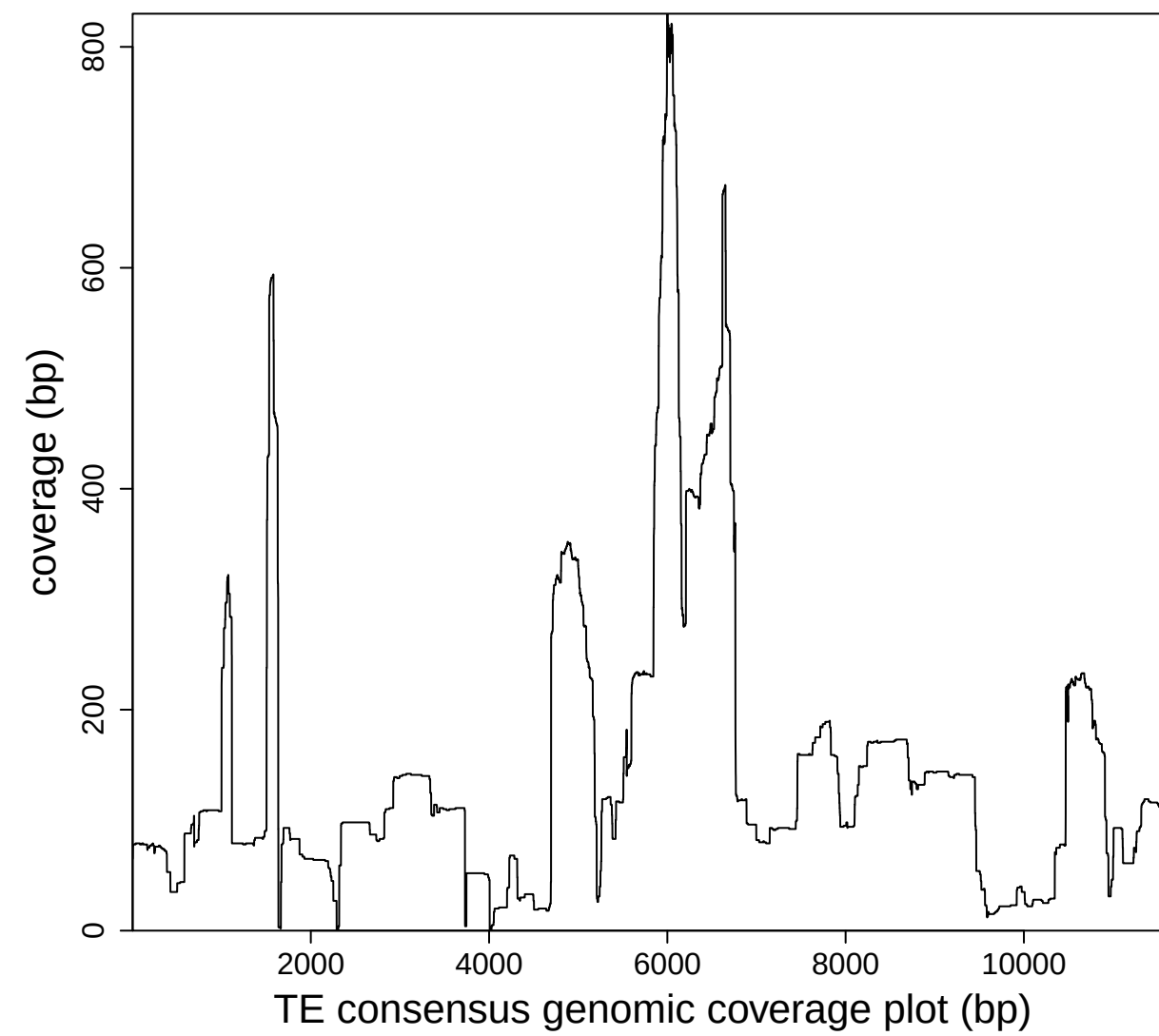
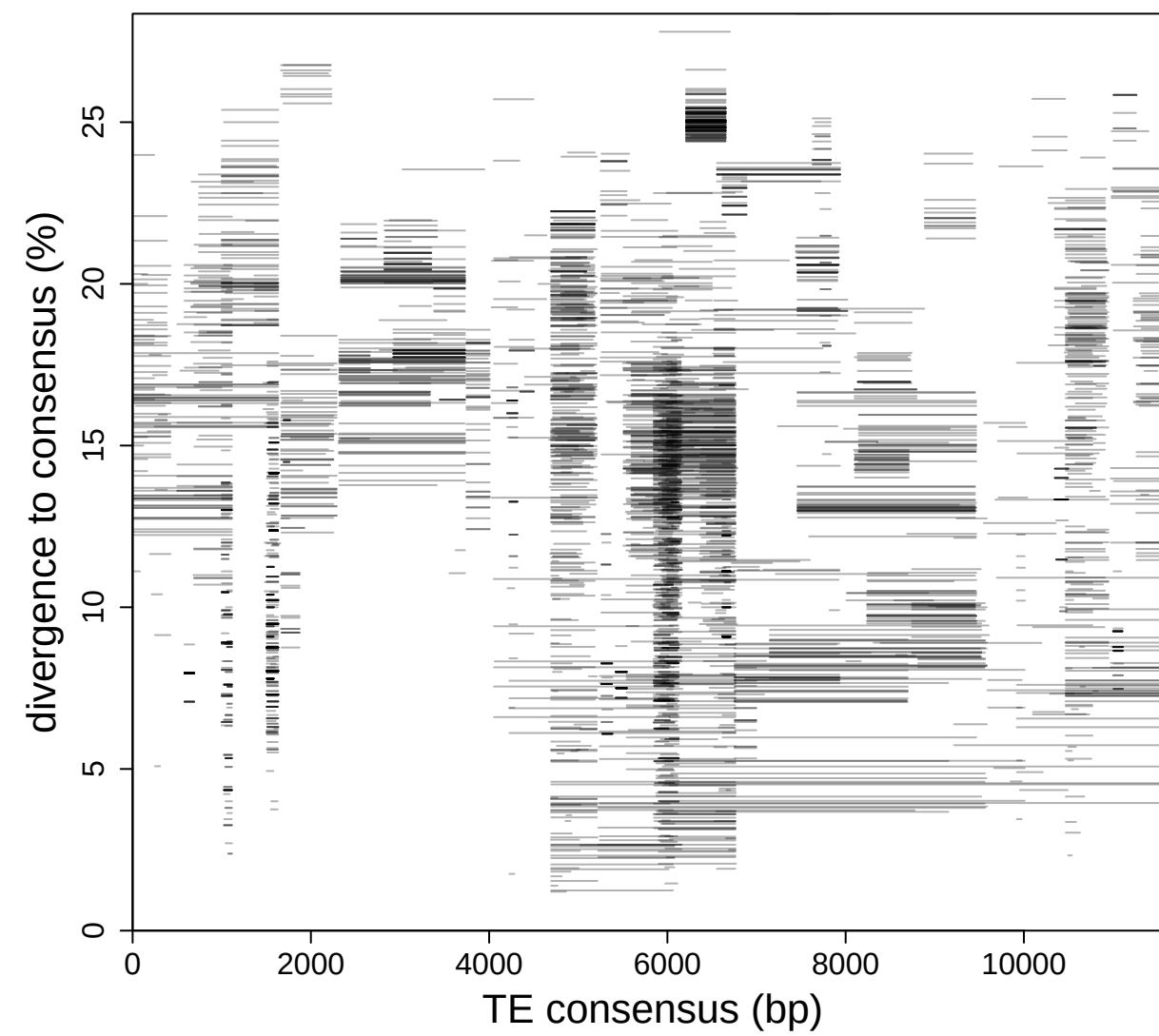
End crop Point





bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_aln.fa
size: 11672bp; fragments: 3884; full length: 0 (>=10504.8bp)

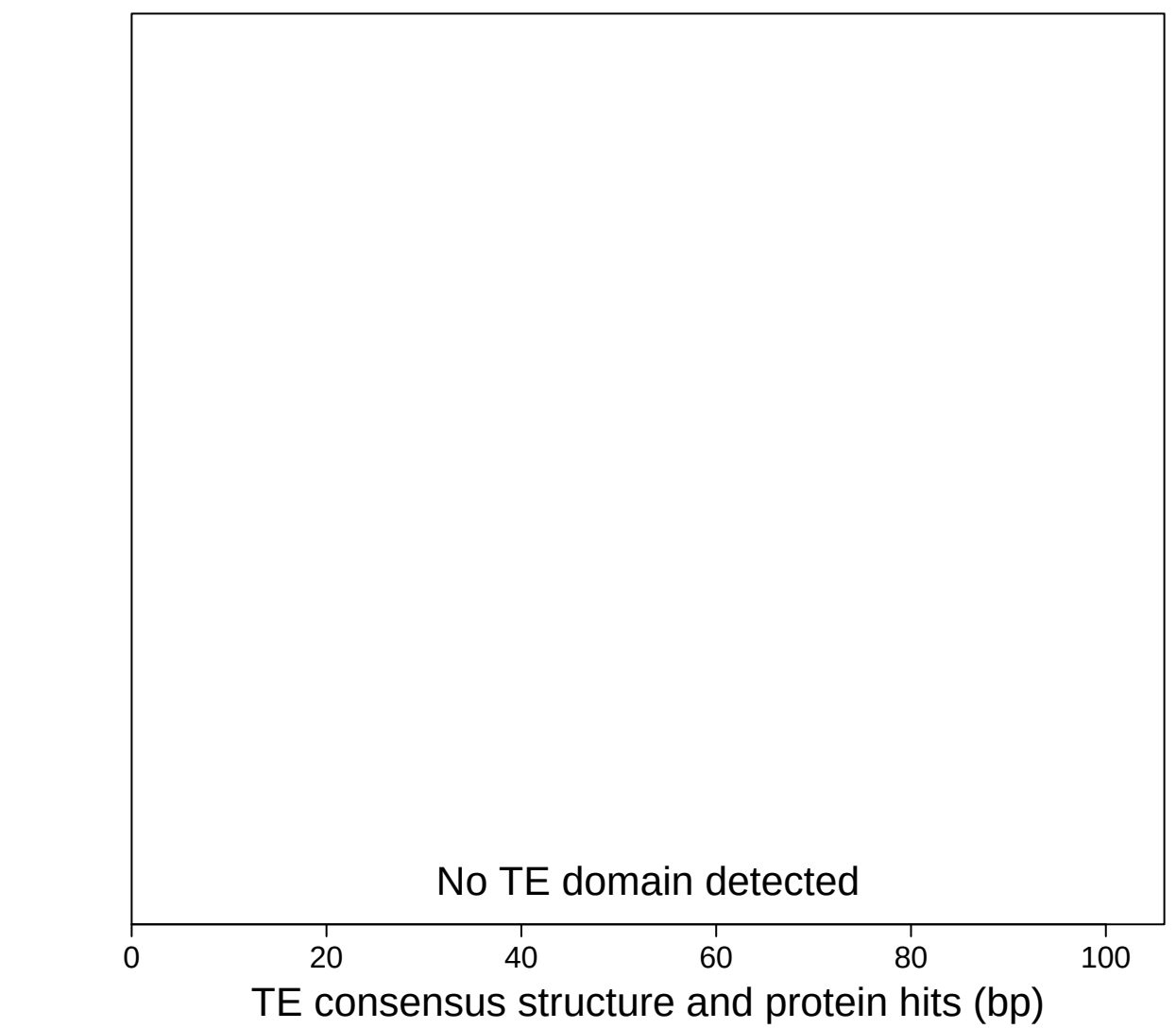
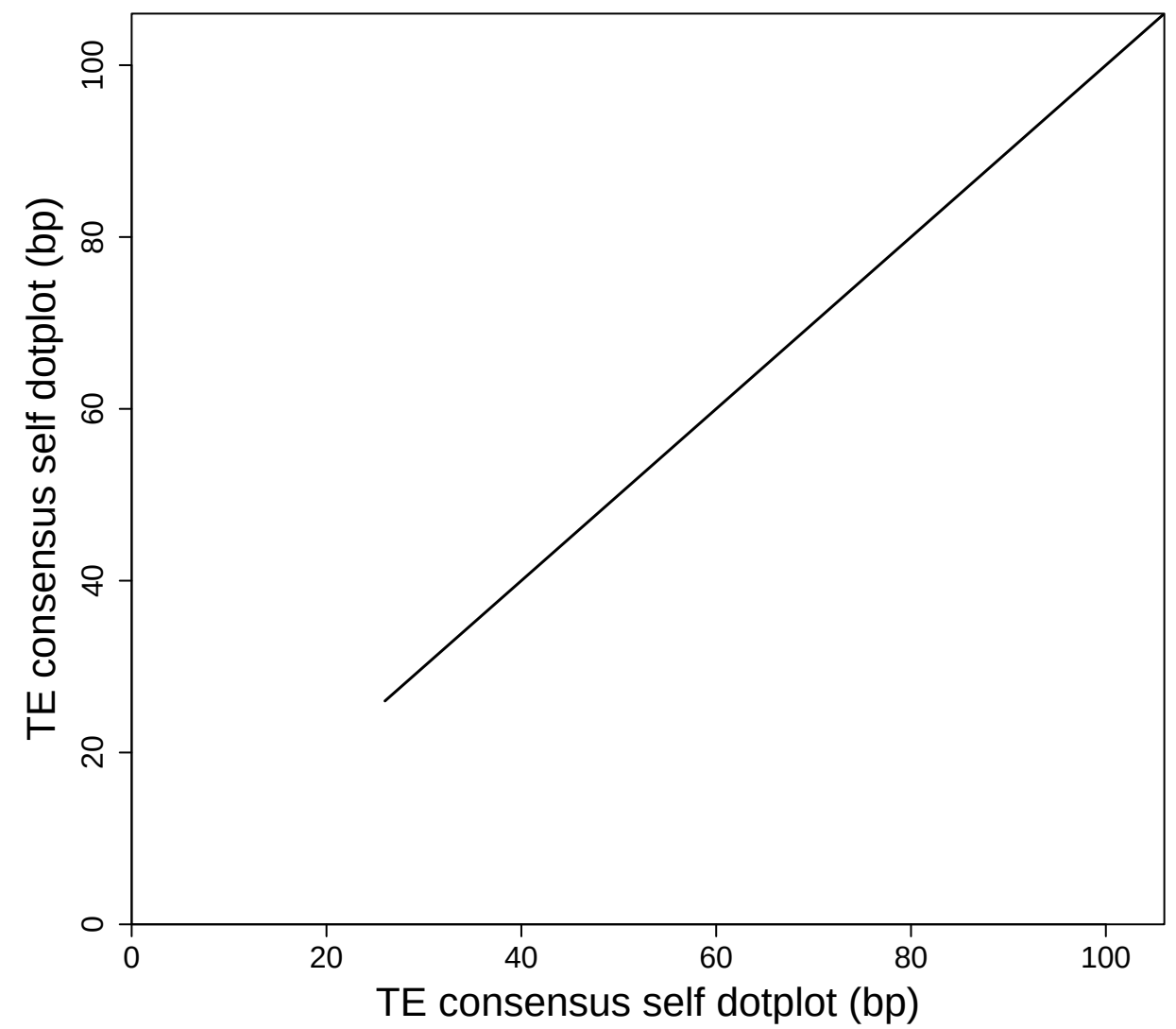
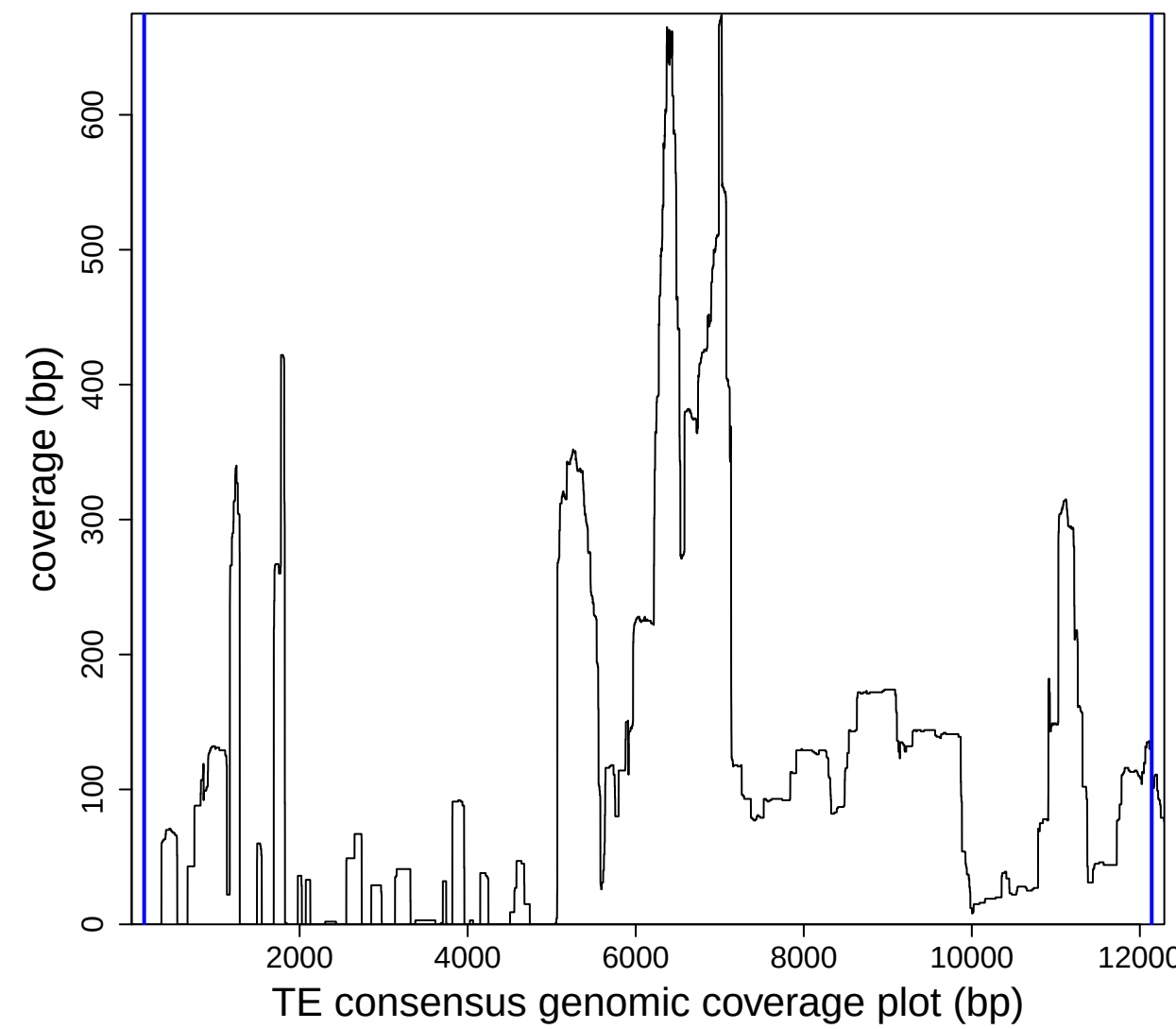
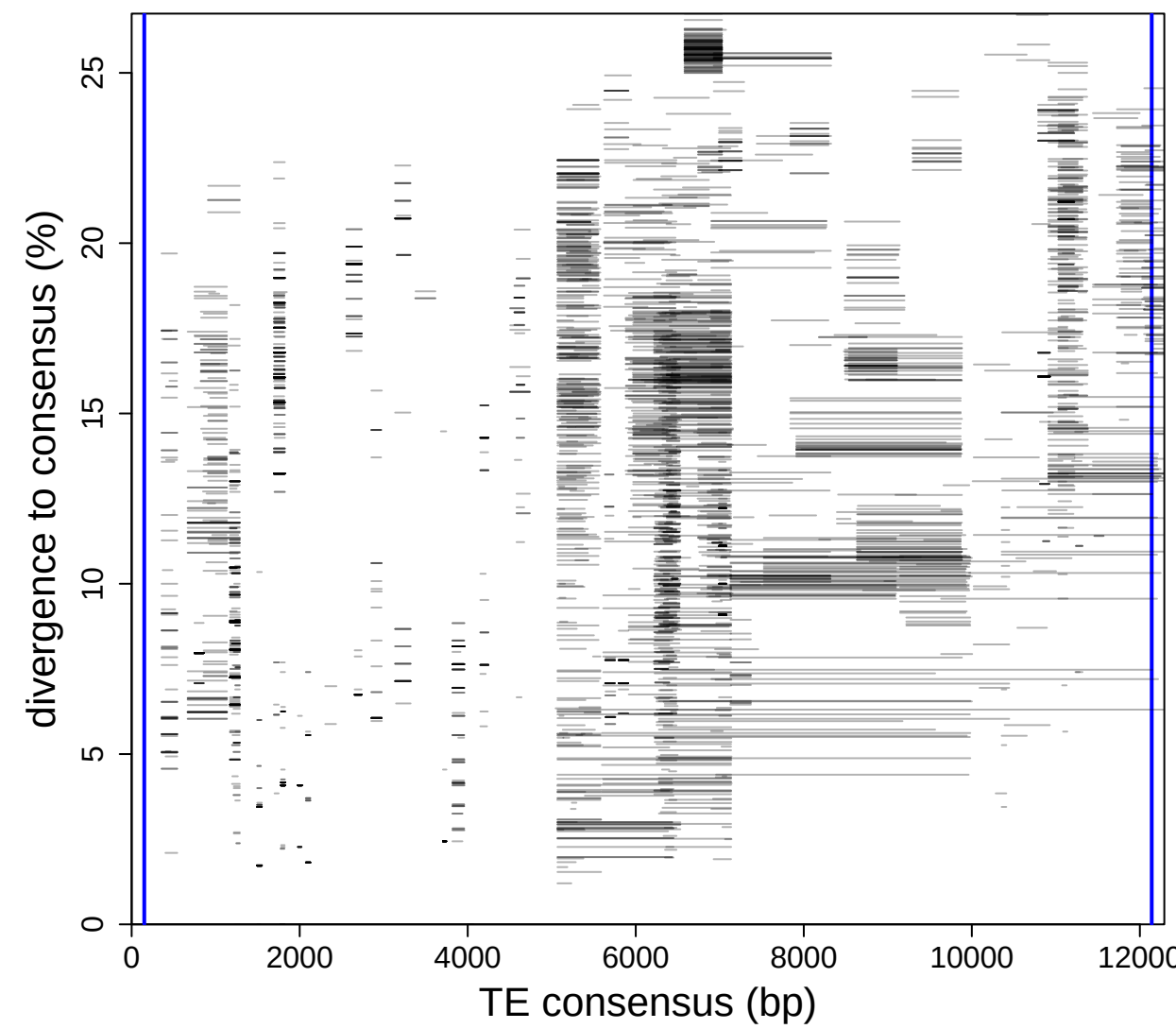
After TEtrimmer 11672 bp



ed_uf.bed g_2.bed fm_1.bed 0 0 n.bed g_1.bed fm_2.bed 0
size: 12291bp; fragments: 3884; full length: 0 (>=11061.9bp)

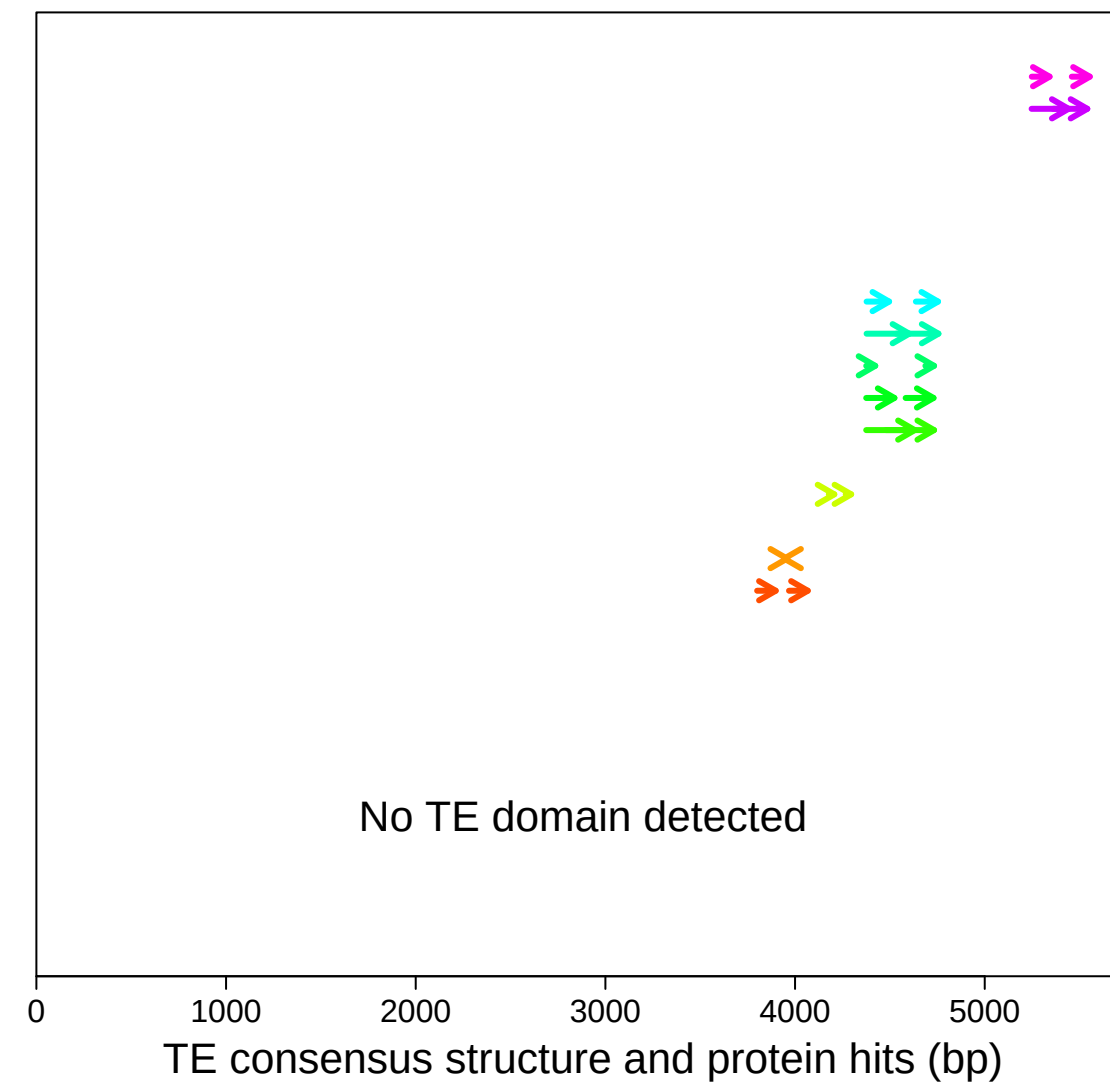
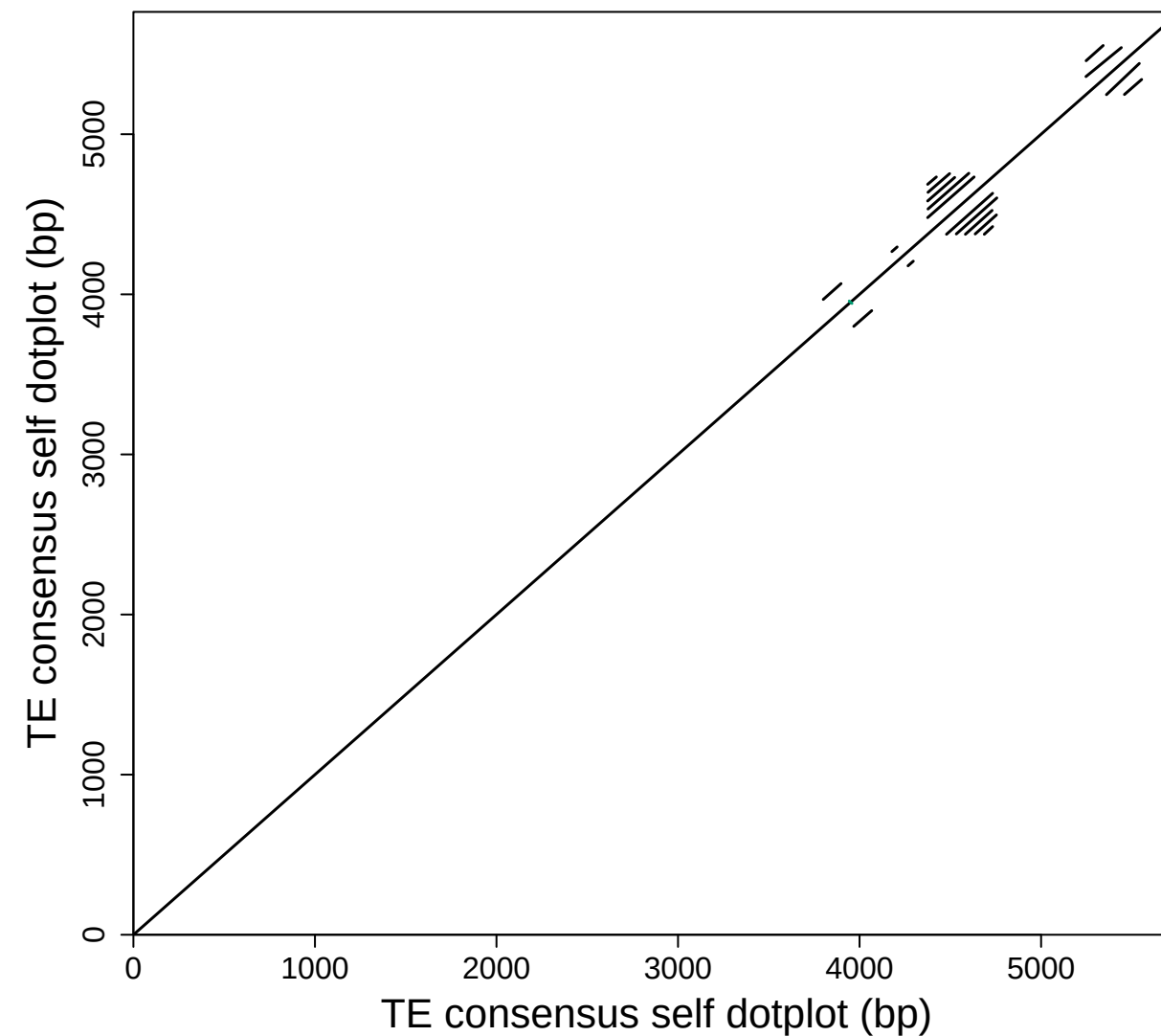
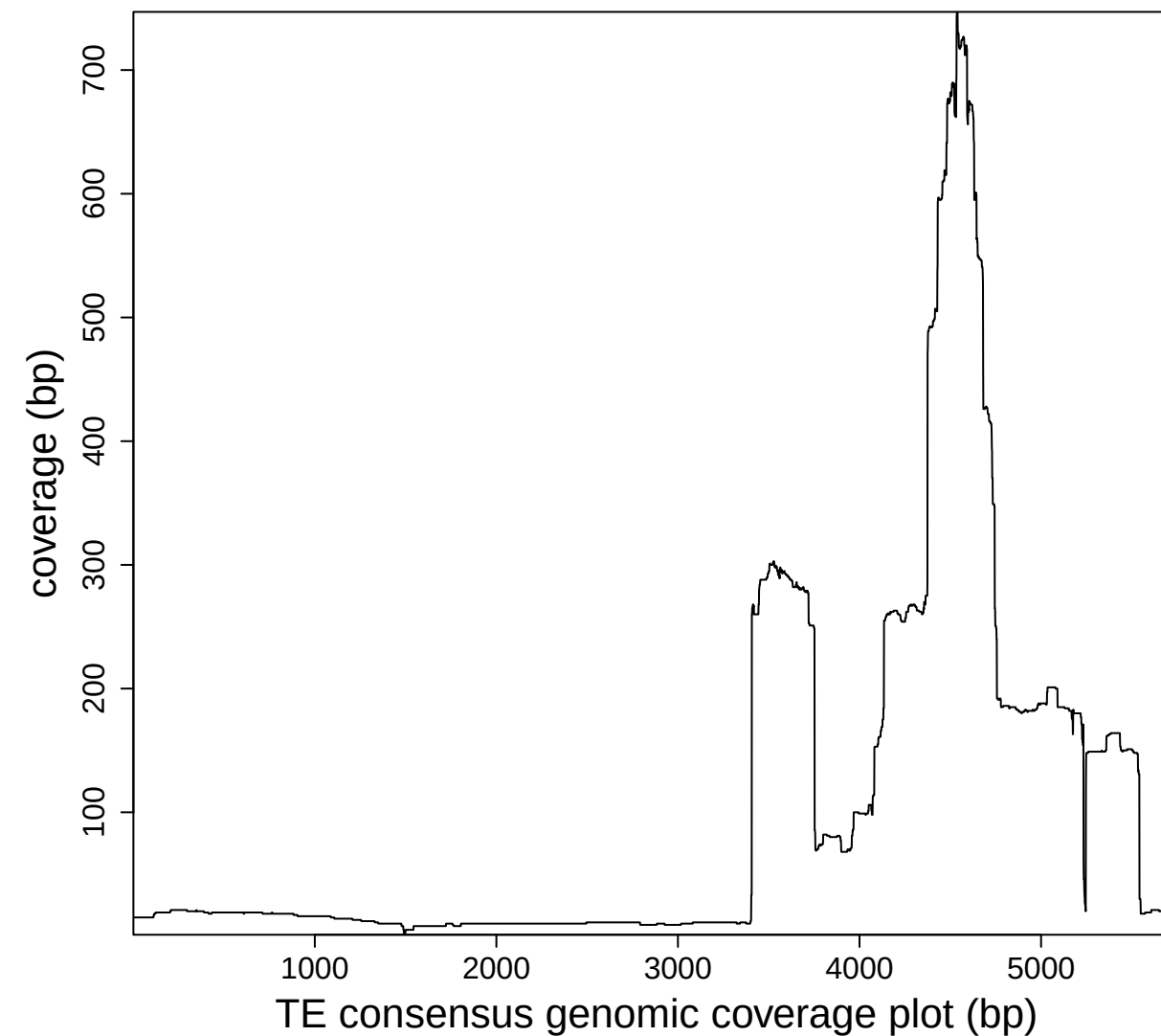
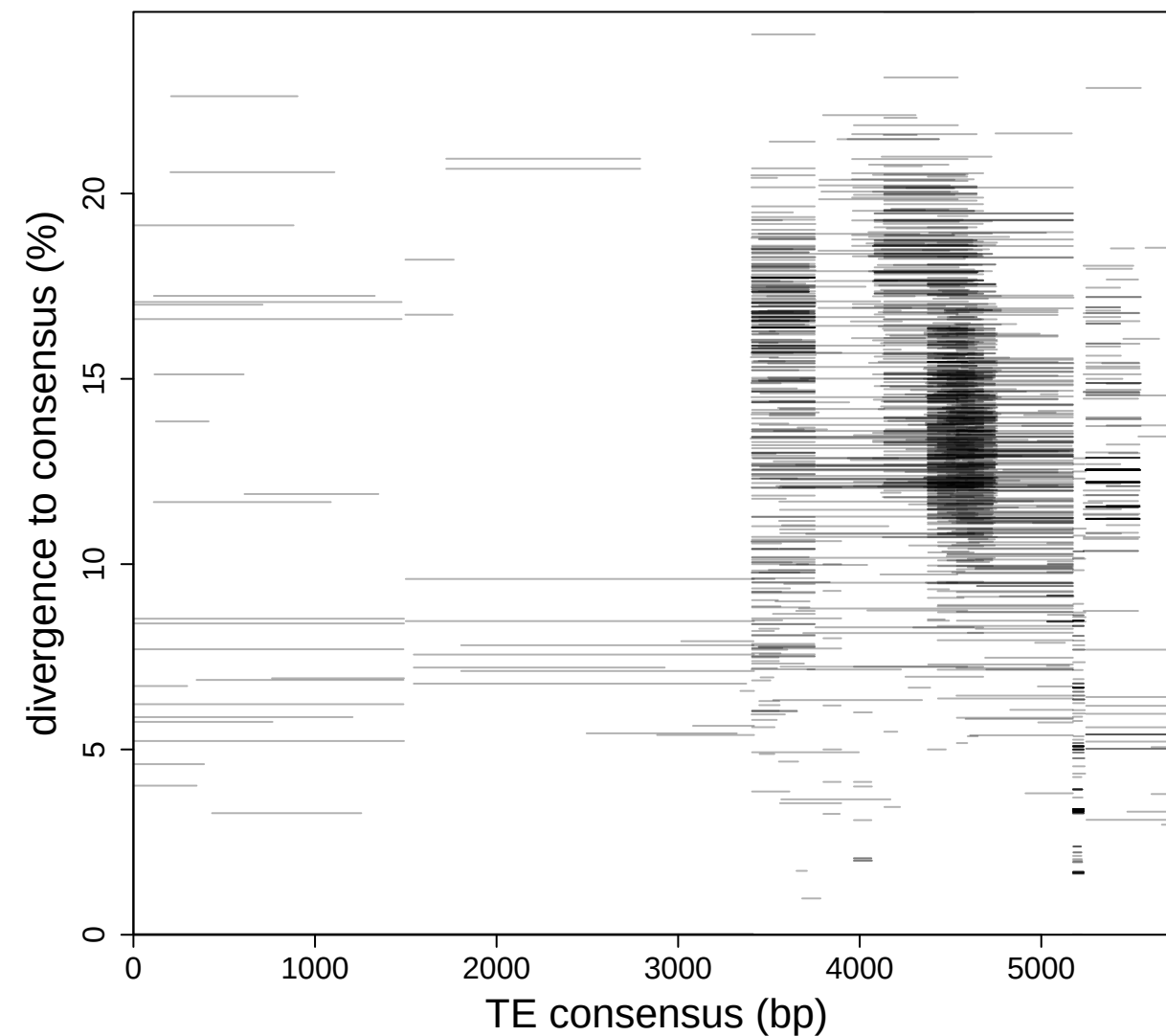
After TEtrimmer Extended plot

Blue lines are boundaries

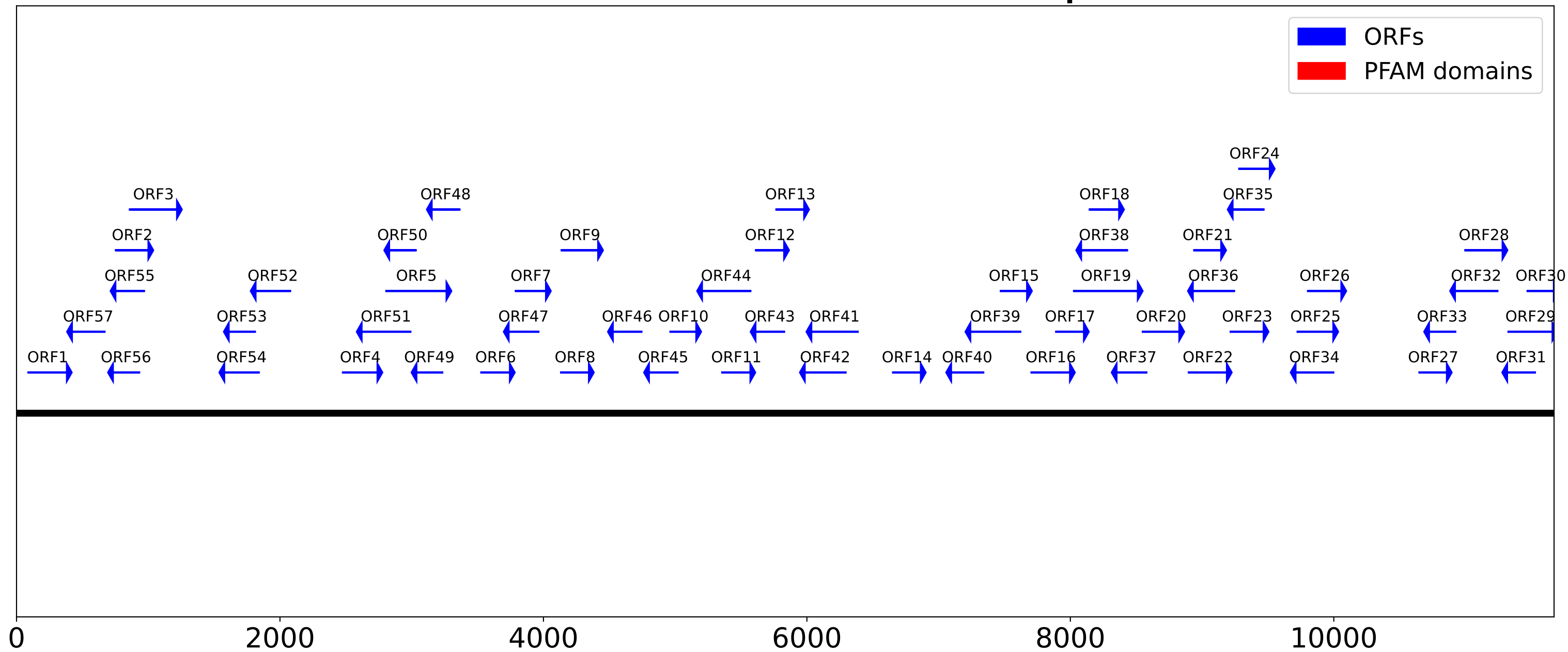


TE: ltr_1_family_907
size: 5764bp; fragments: 1631; full length: 0 (≥ 5187.6 bp)

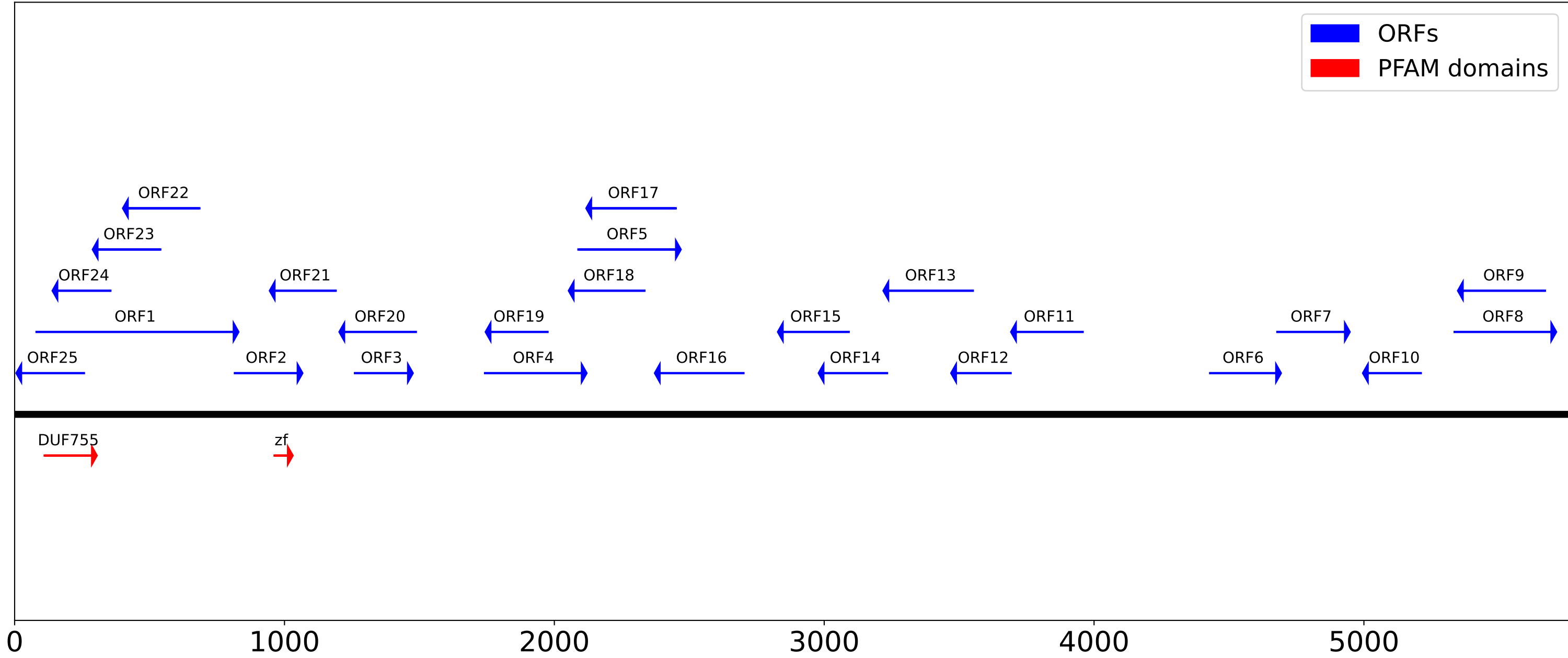
Before TEtrimmer 5764 bp



After TETrimmer ORF and PFAM domain plot



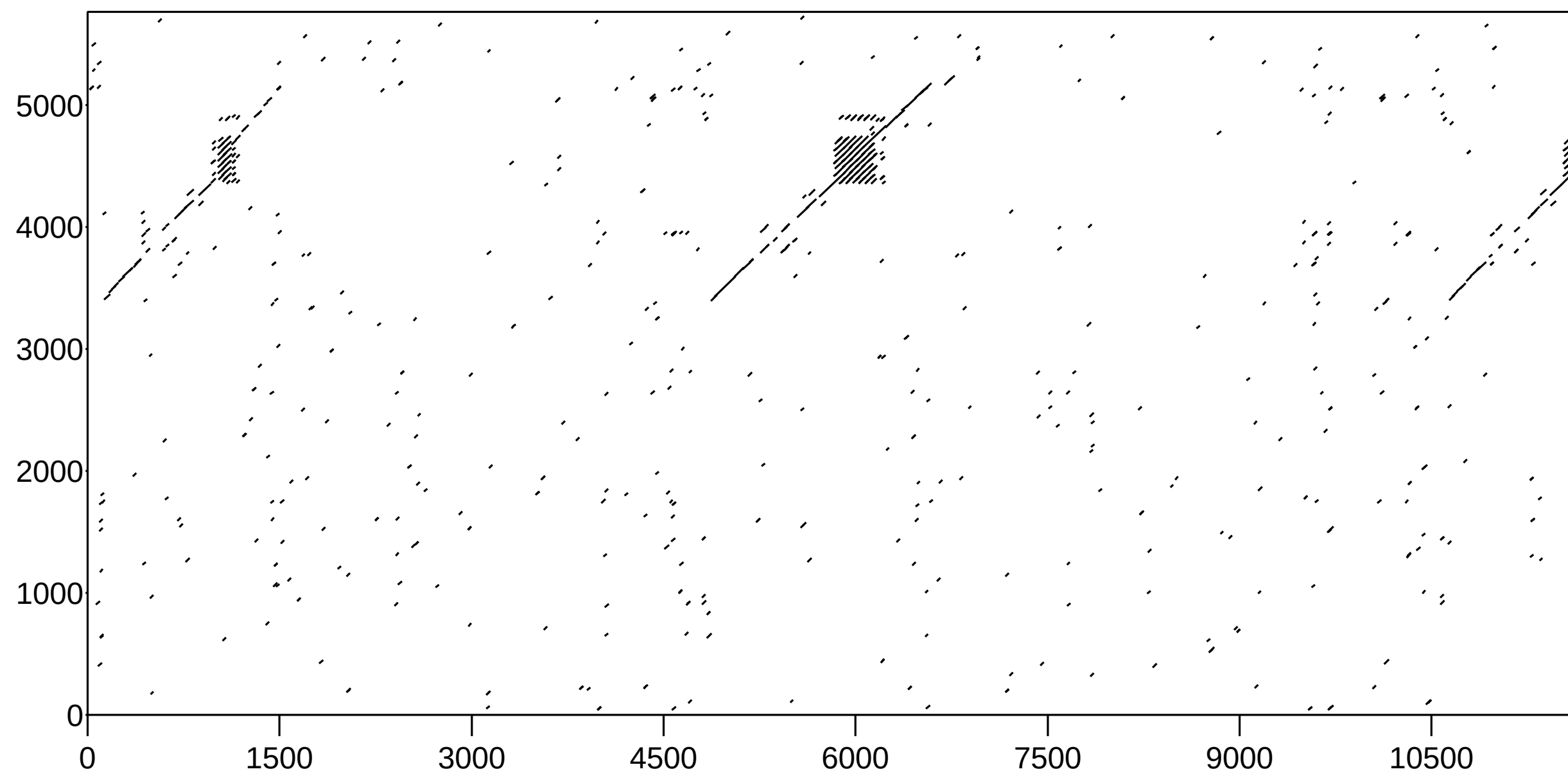
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)