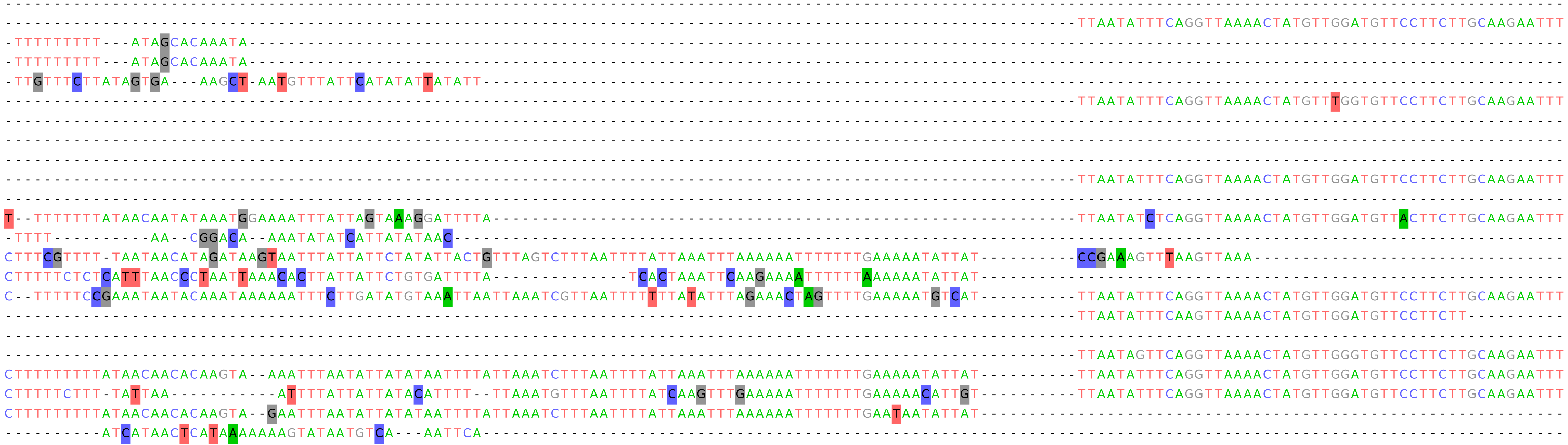
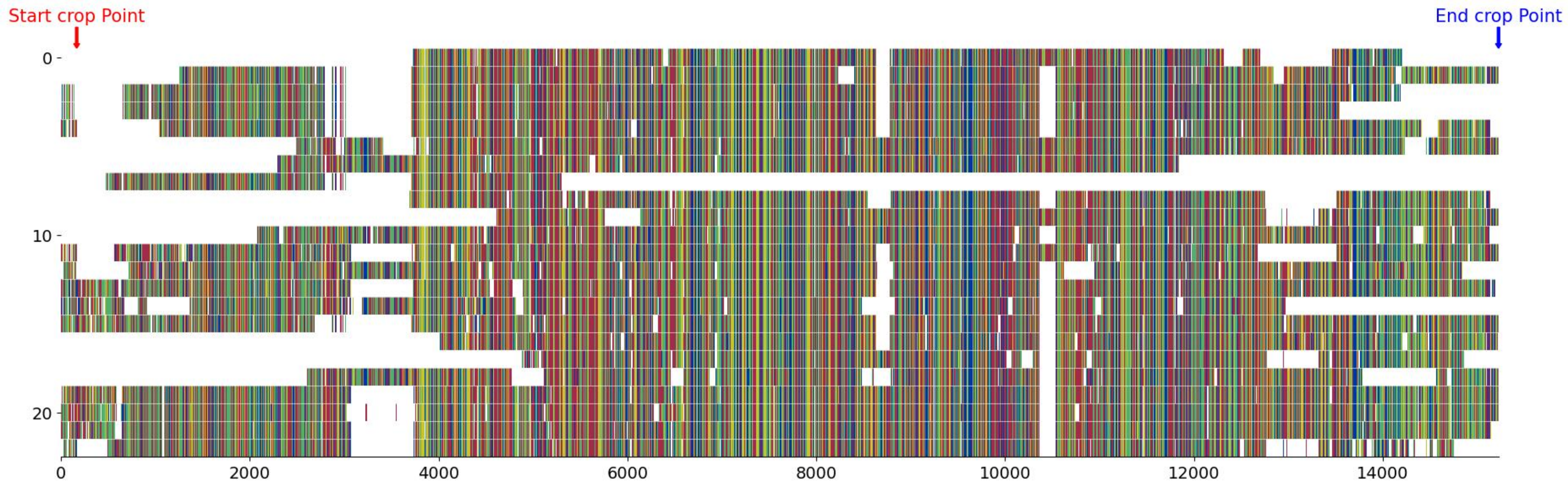


Start crop Point

End crop Point

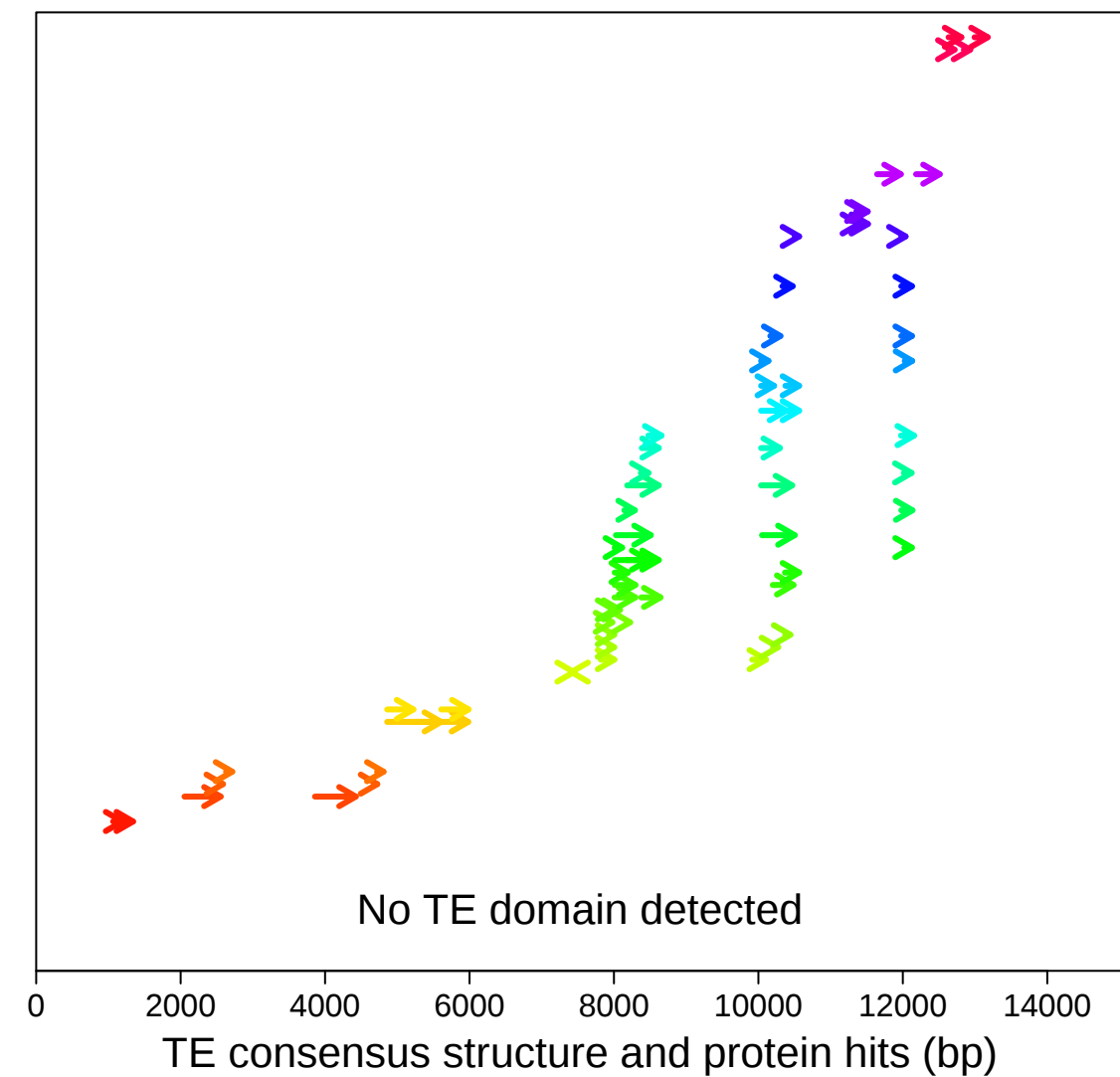
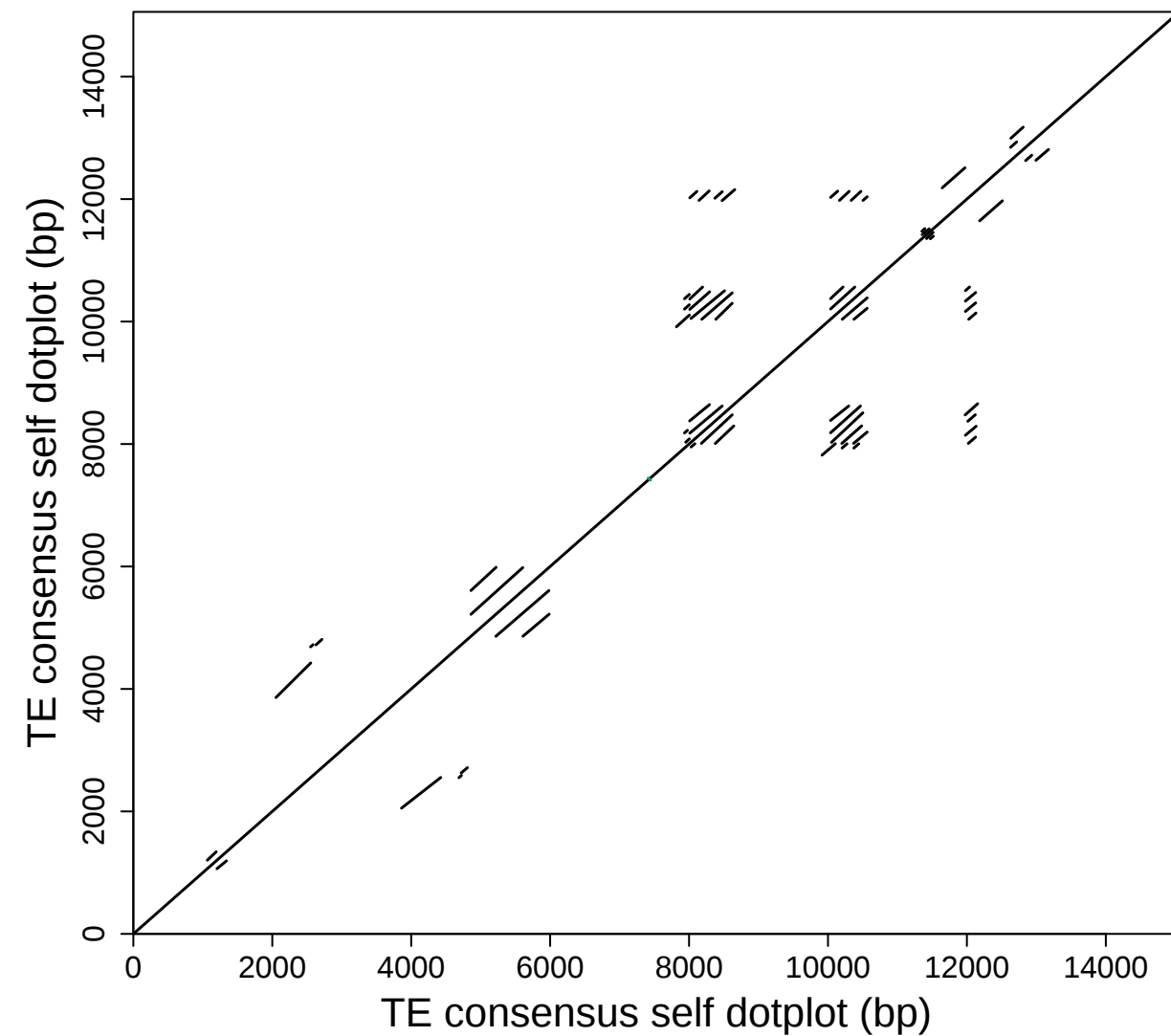
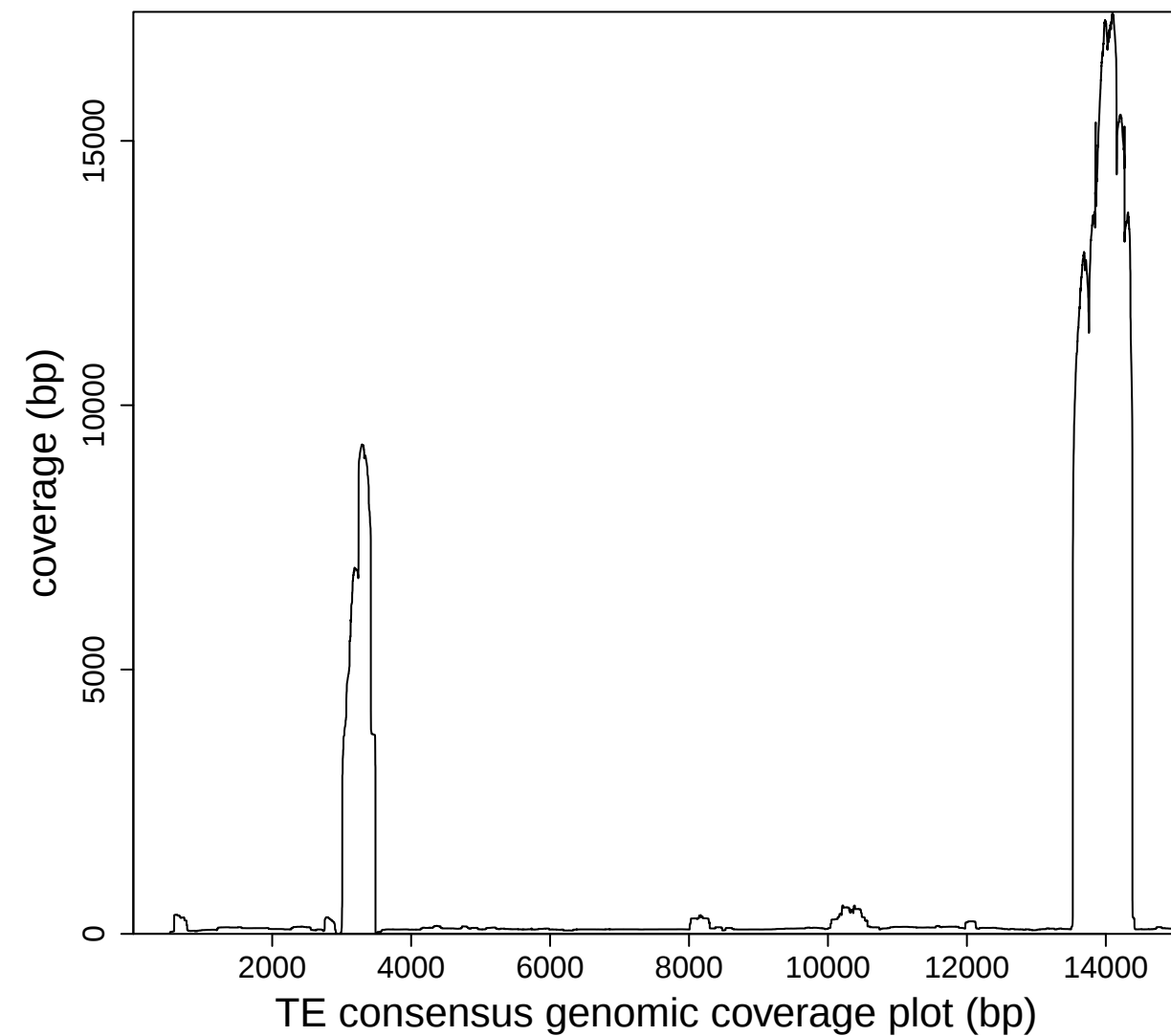
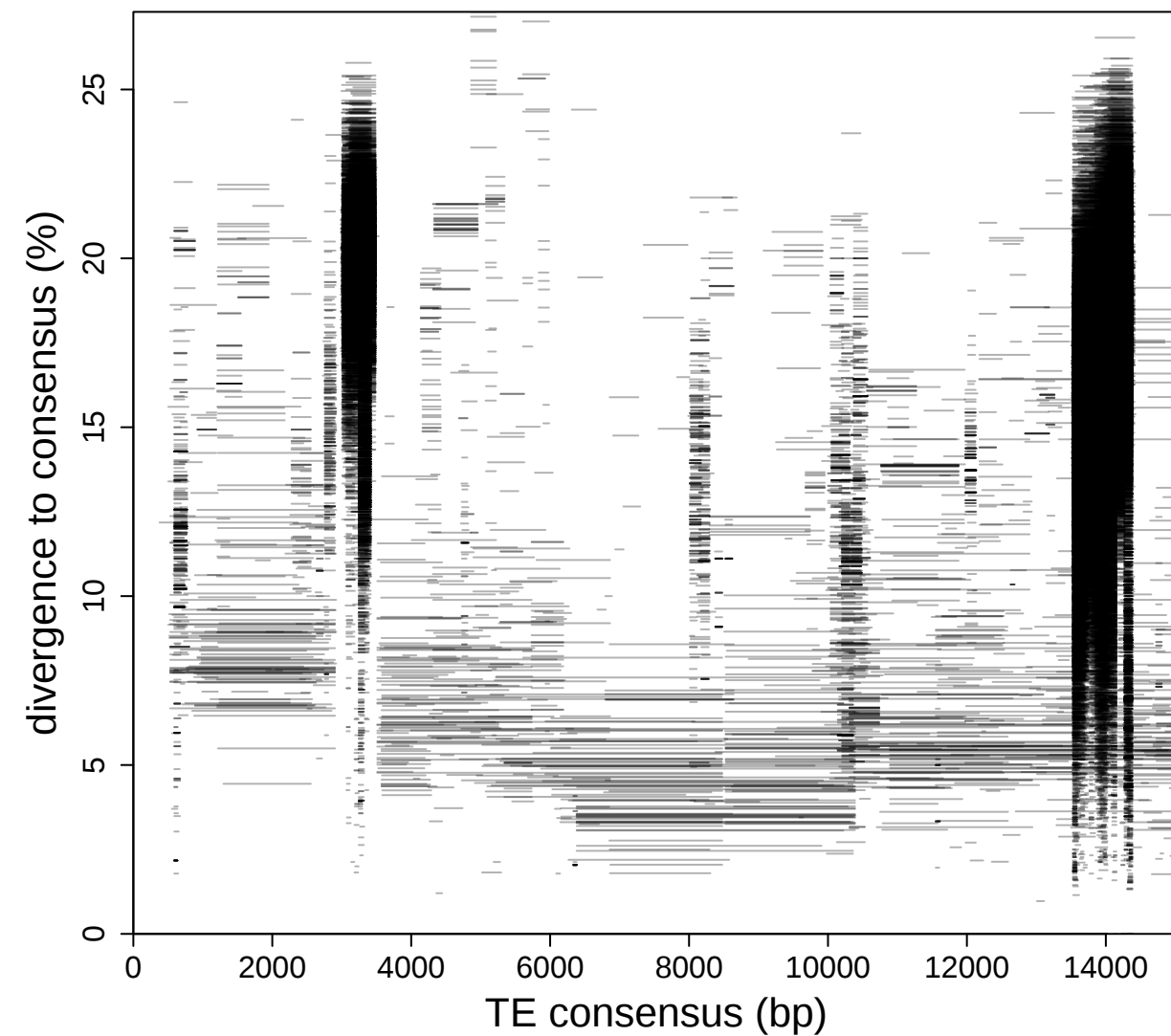
MSA length = 15057





1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.f
size: 15057bp; fragments: 46156; full length: 0 (>=13551.3bp)

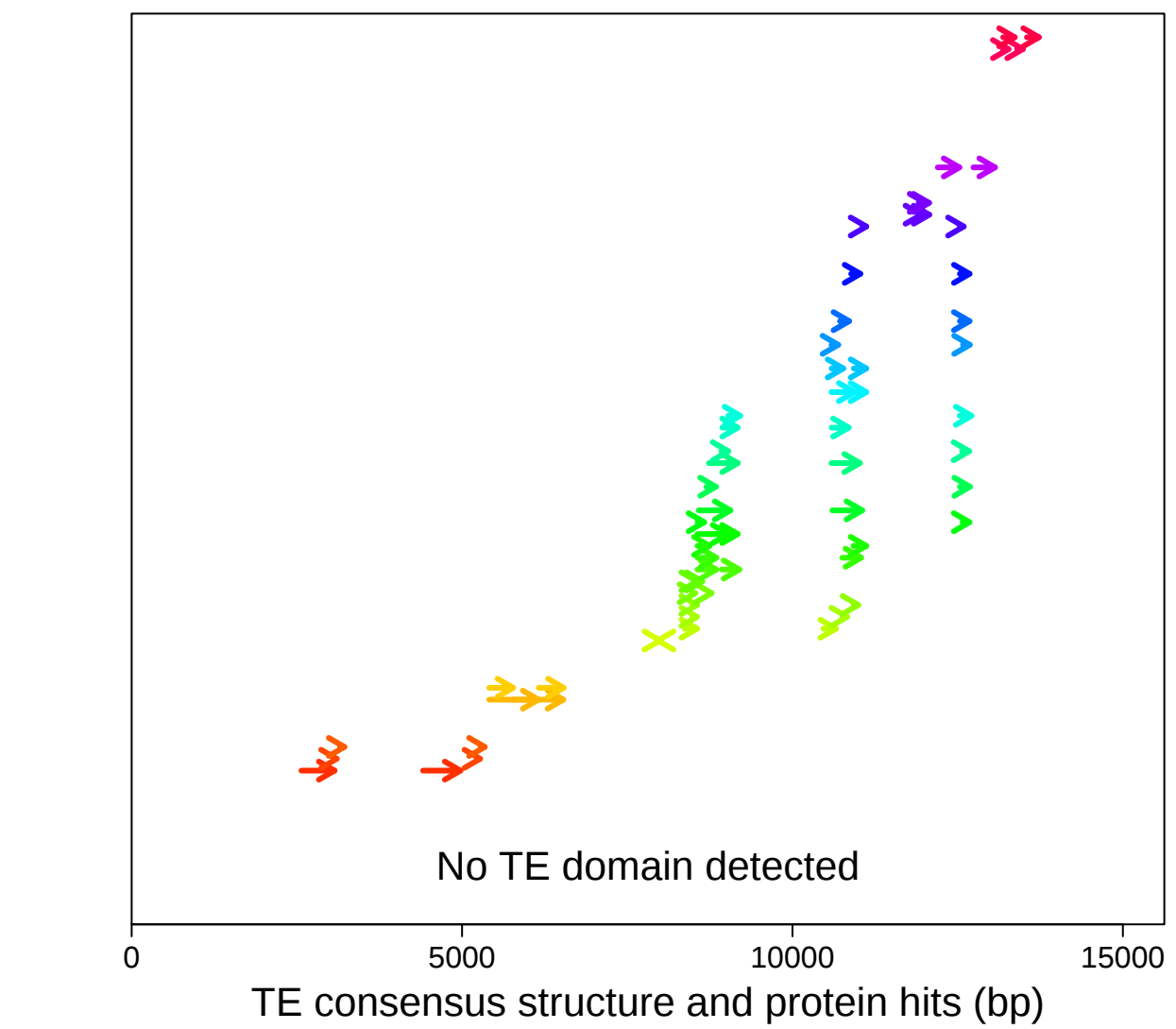
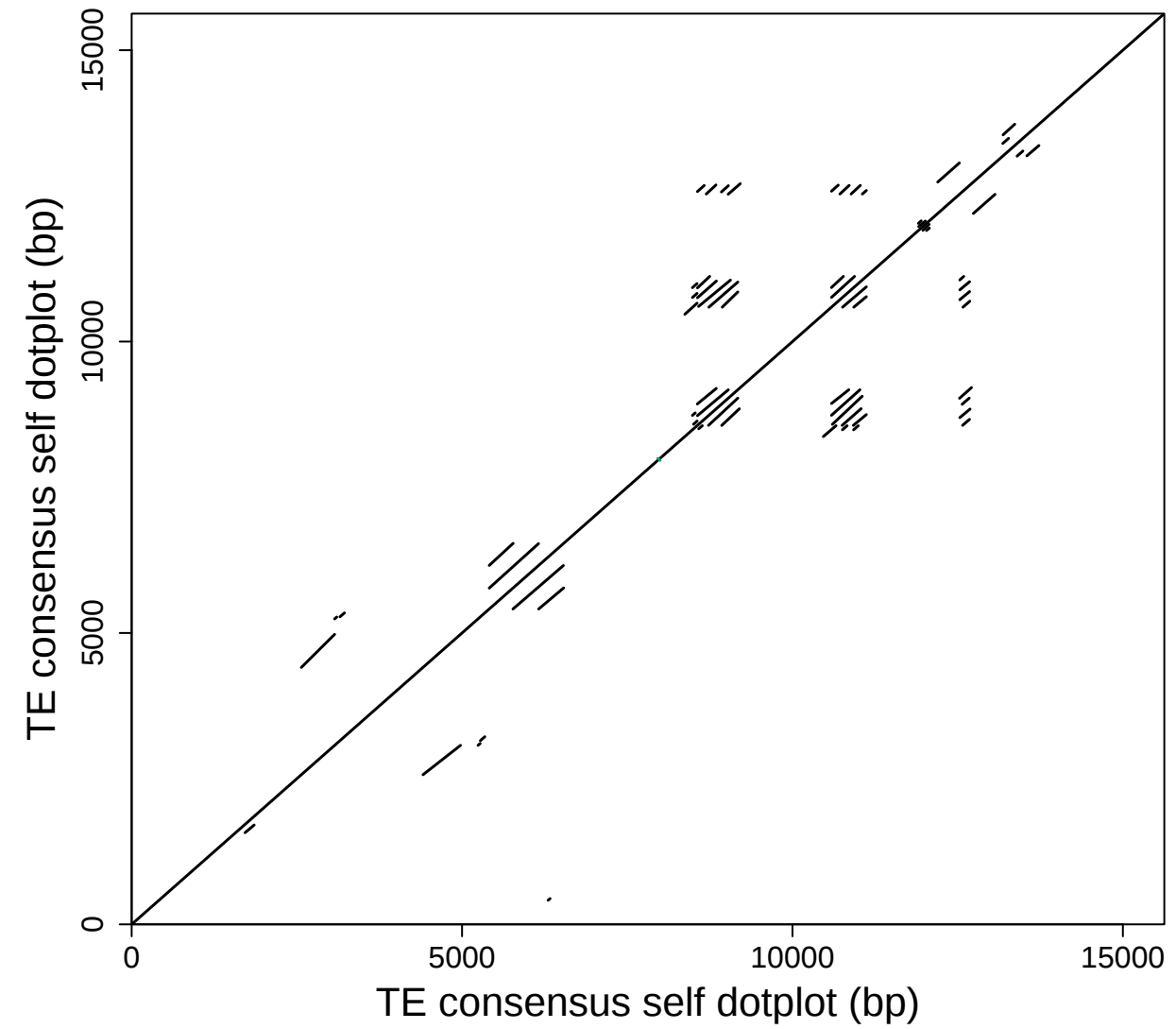
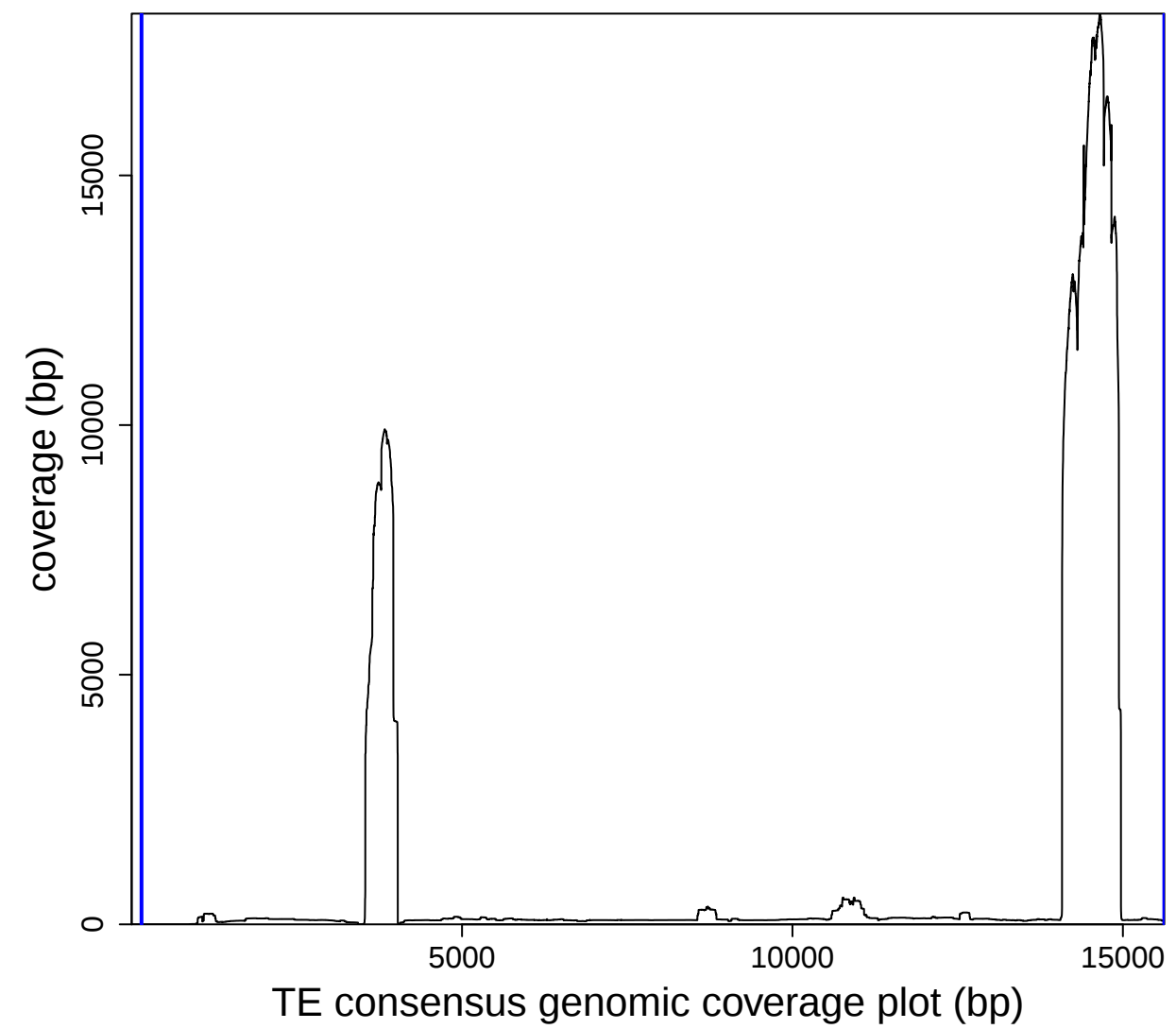
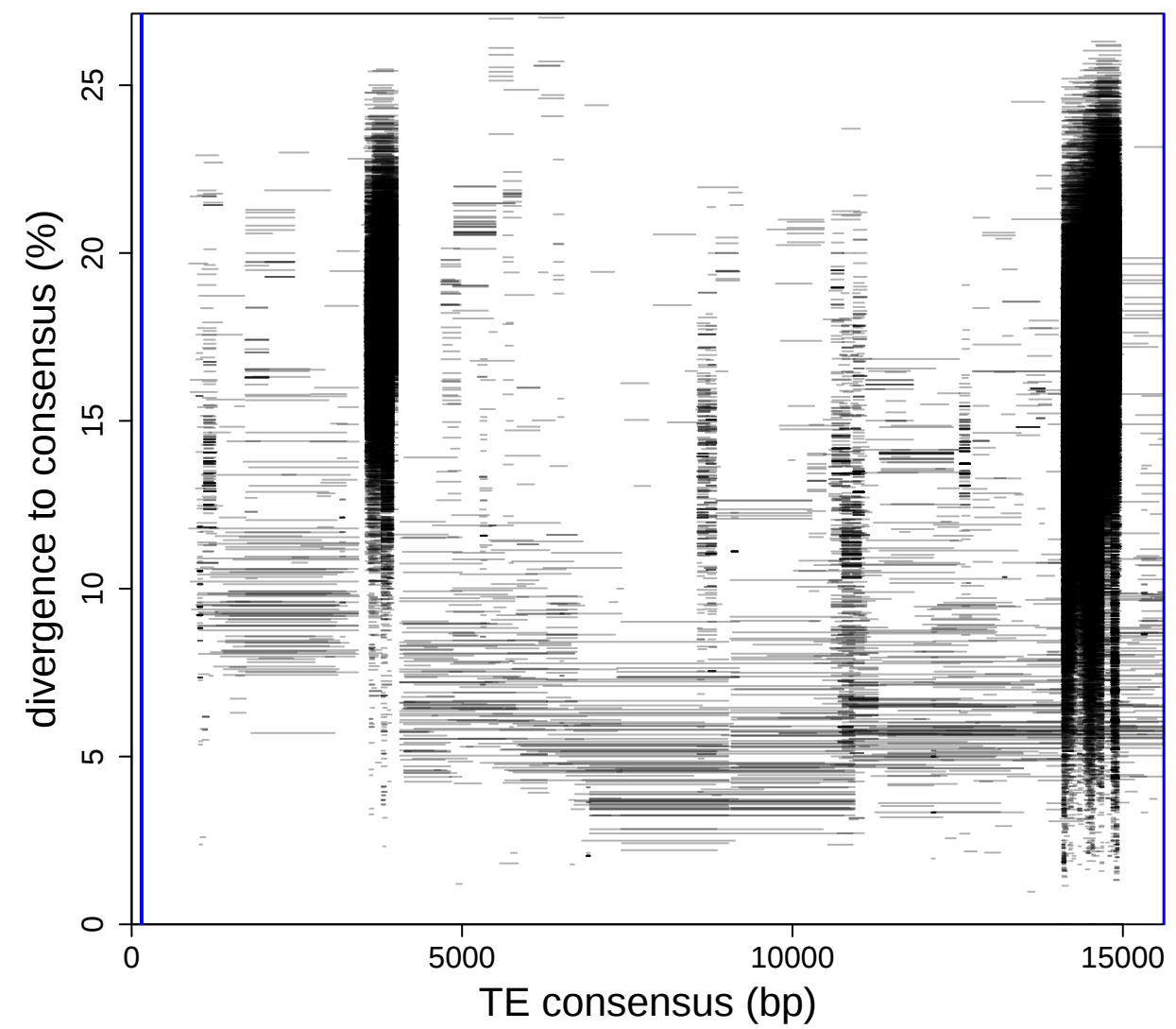
After TEtrimmer 15057 bp



f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bc
size: 15627bp; fragments: 47528; full length: 0 (>=14064.3bp)

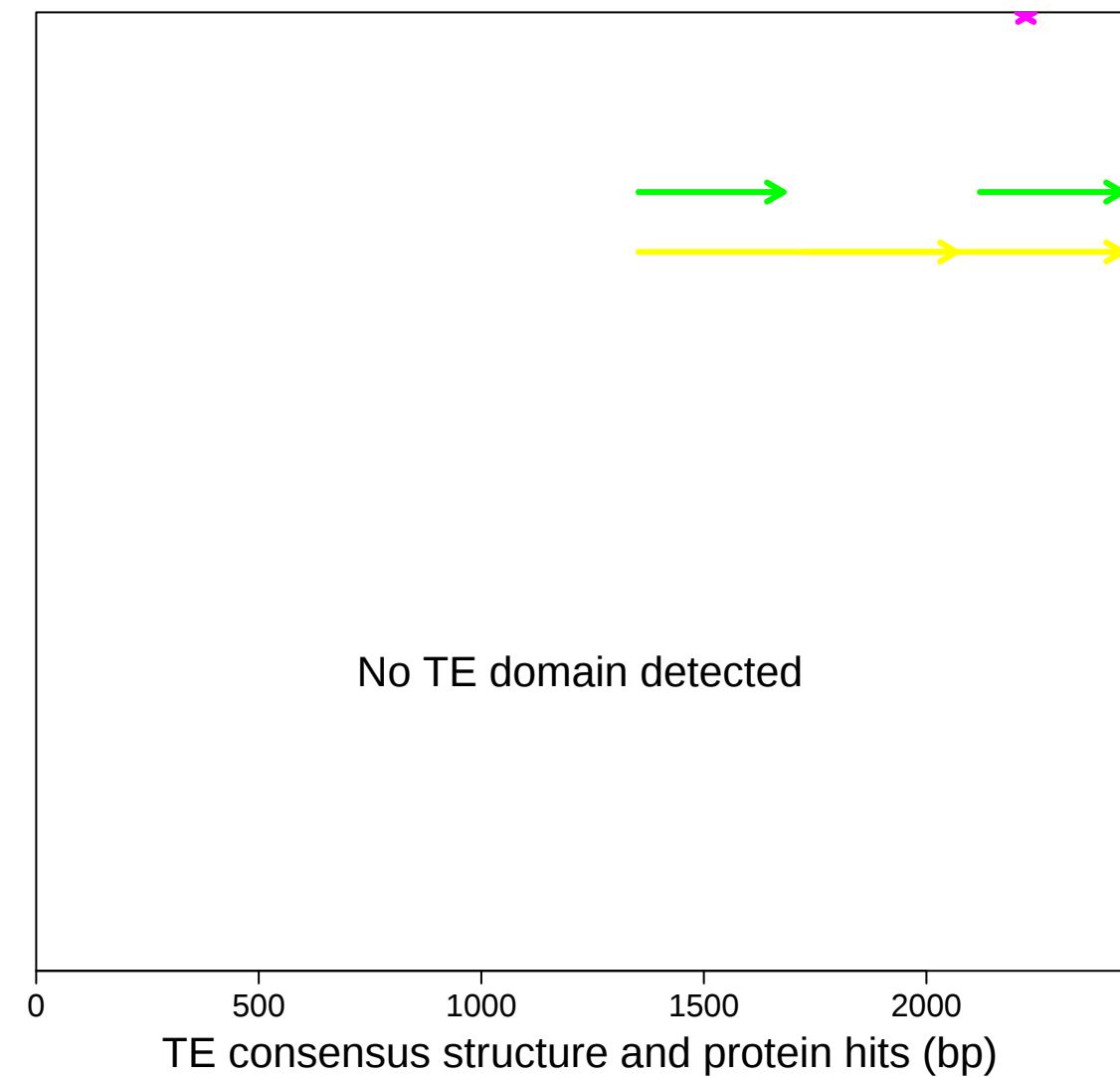
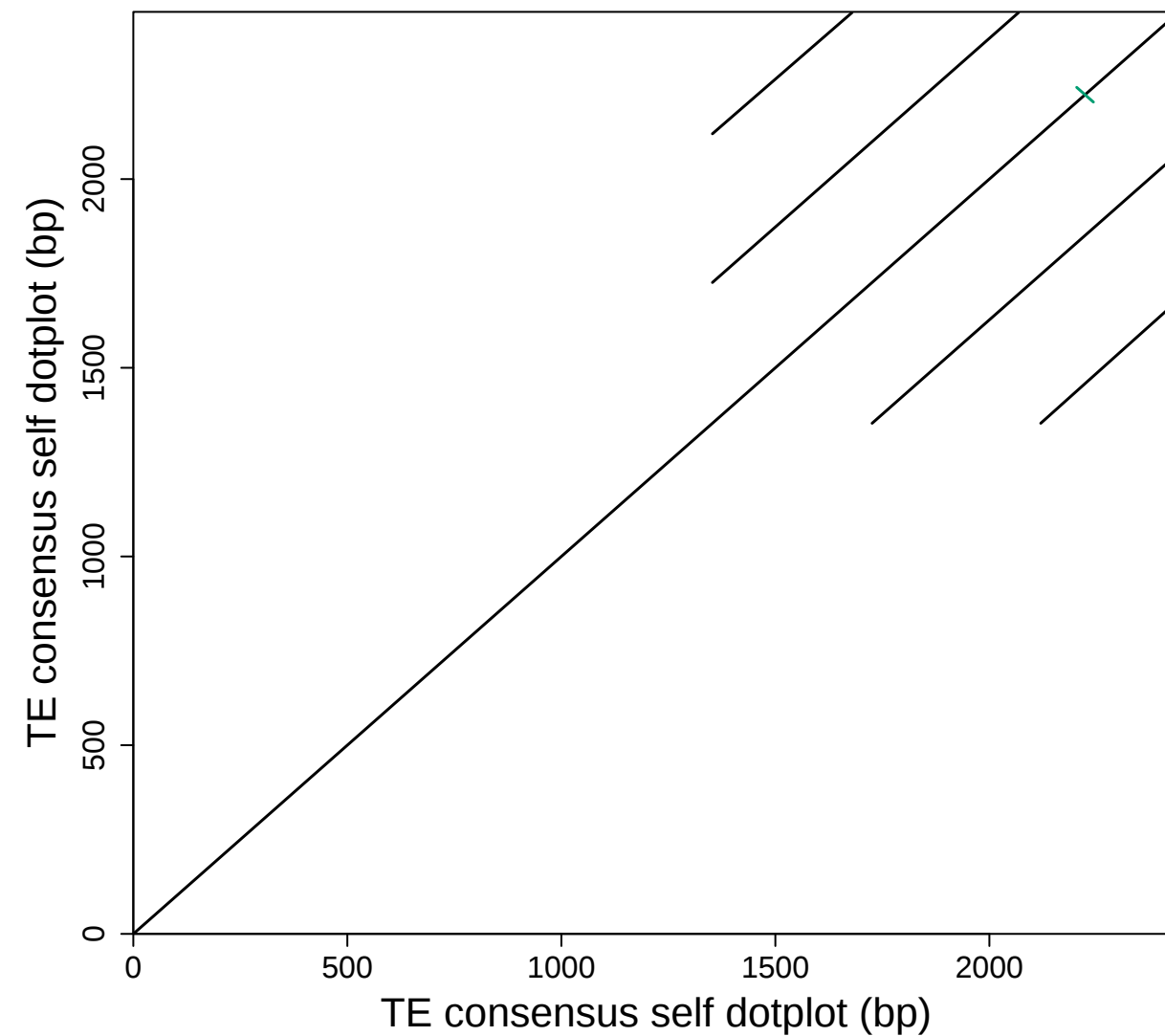
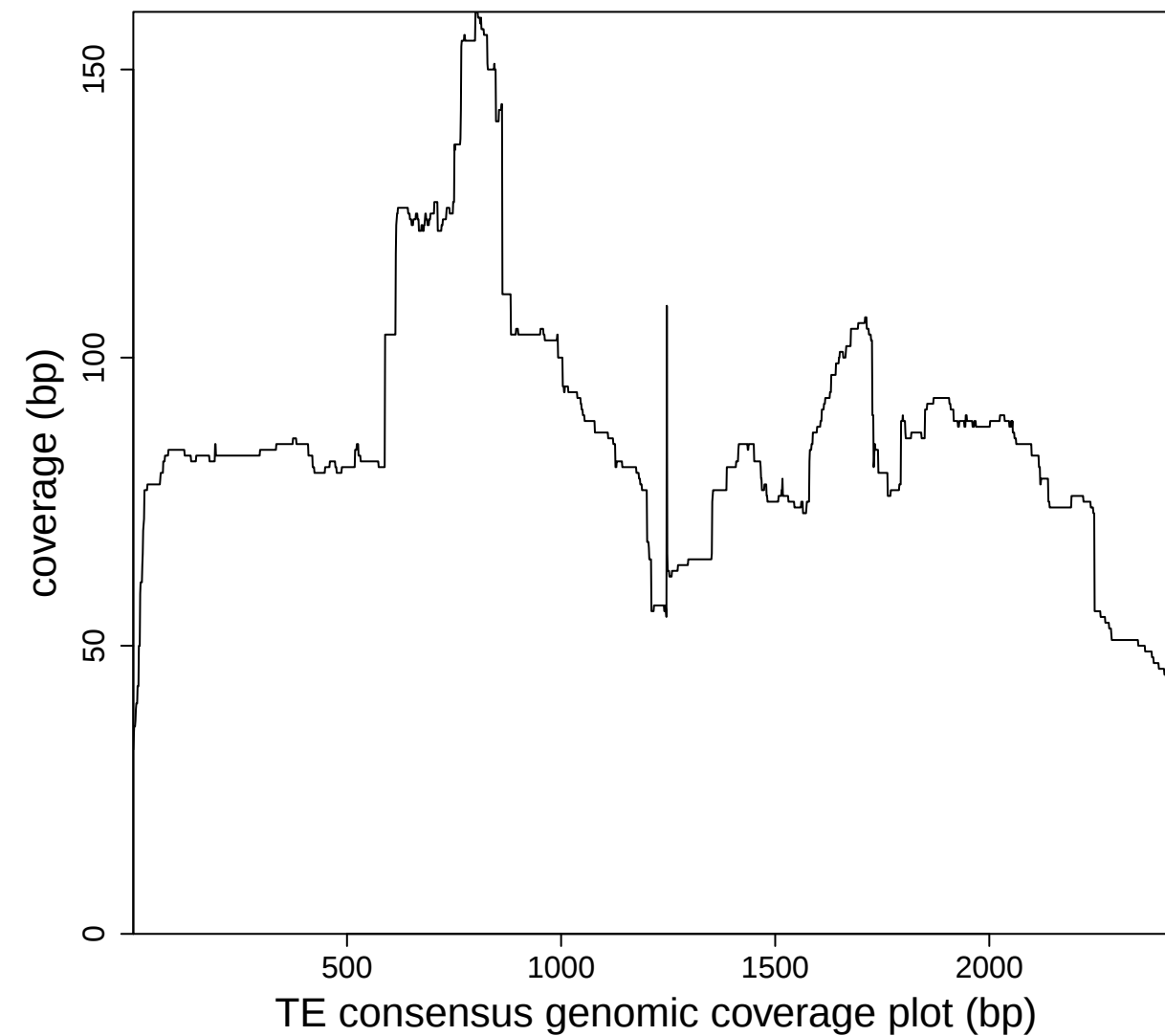
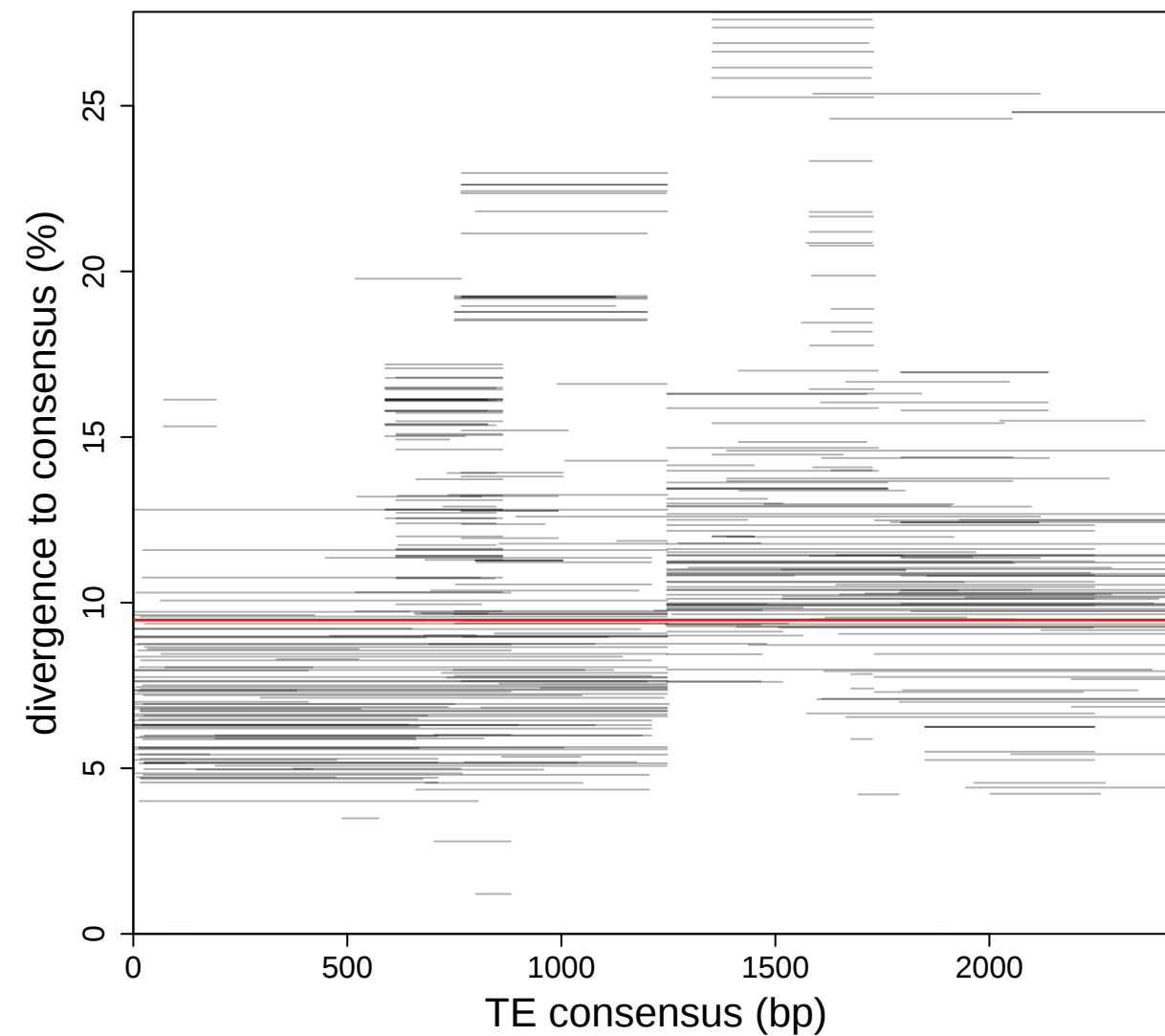
After TEtrimmer Extended plot

Blue lines are boundaries

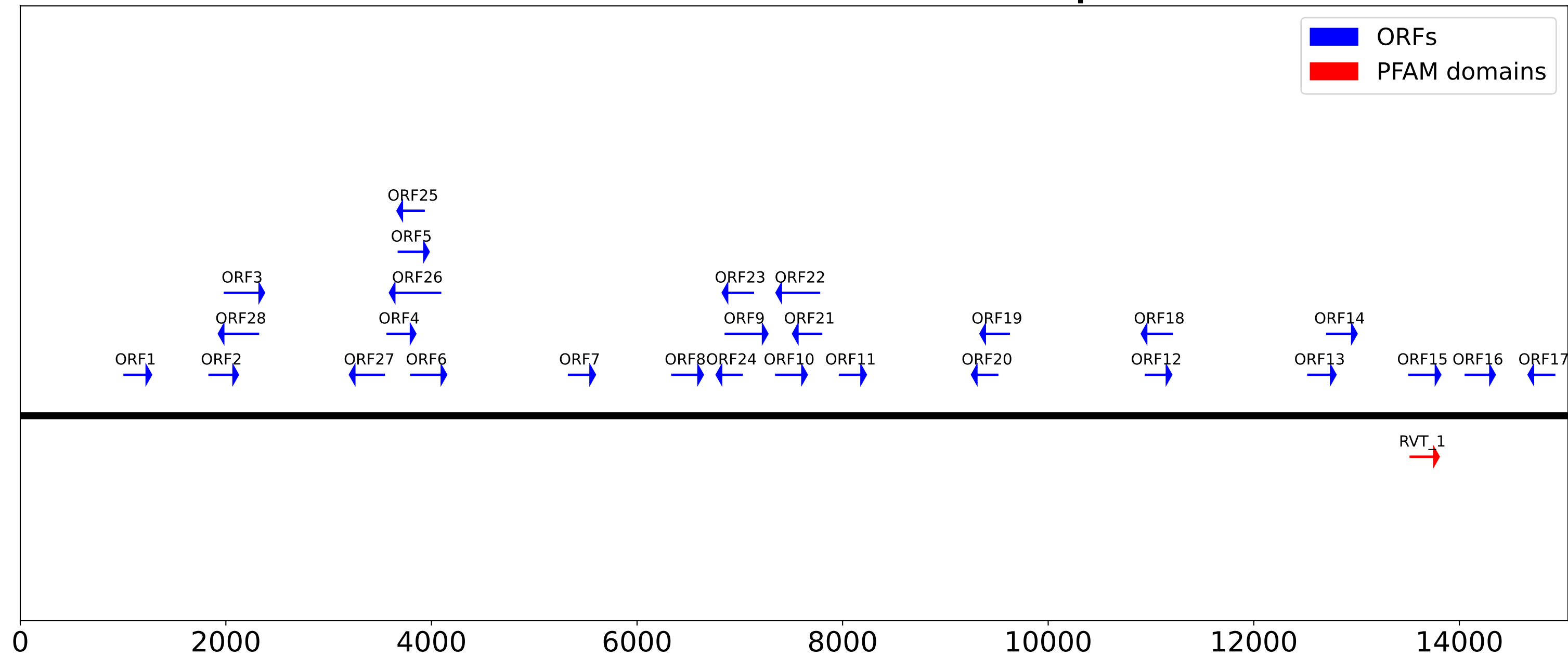


TE: rnd_5_family_4738
size: 2443bp; fragments: 374; full length: 1 (>=2198.7bp)

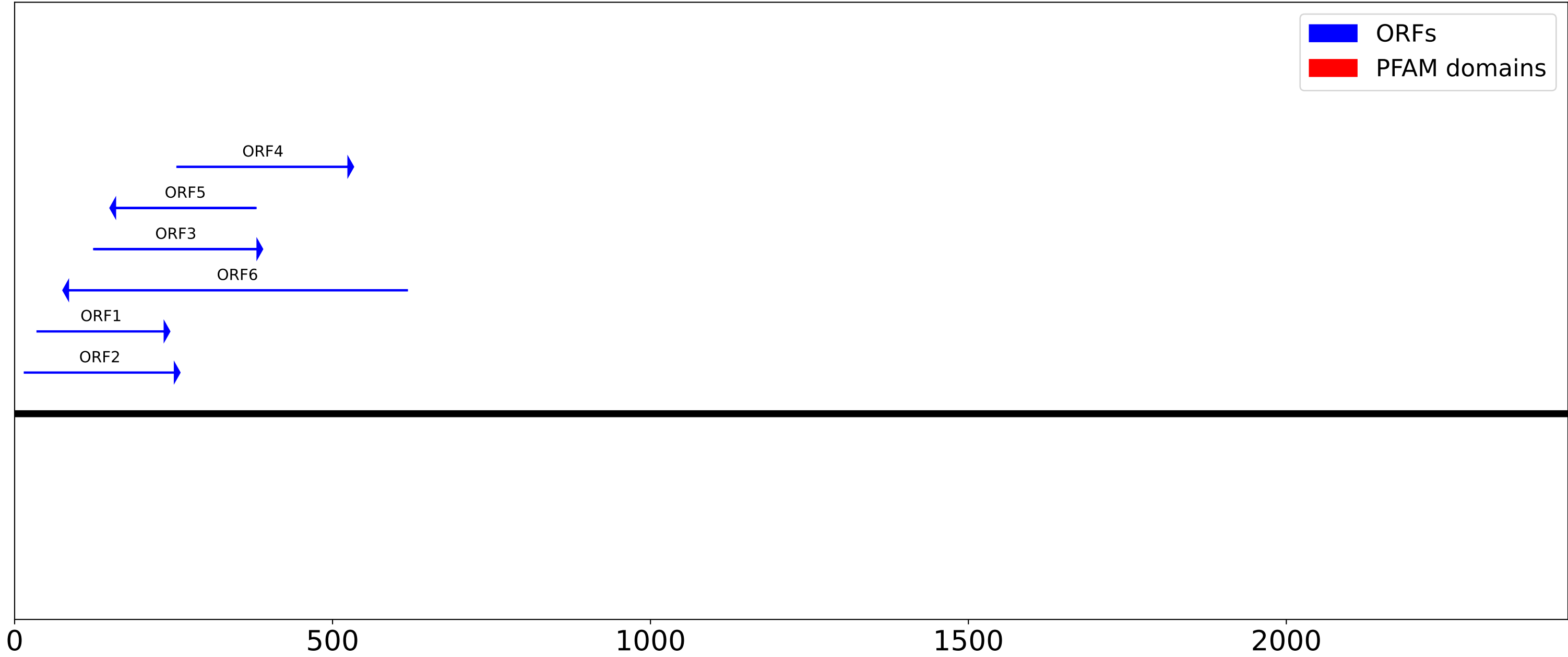
Before TEtrimmer 2443 bp



After TEtrimmer ORF and PFAM domain plot



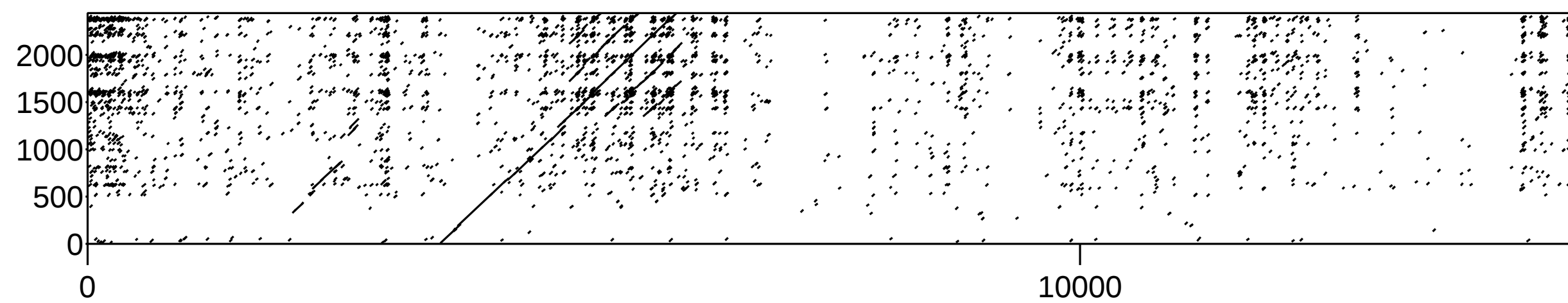
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

rsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)