

Start crop Point



End crop Point

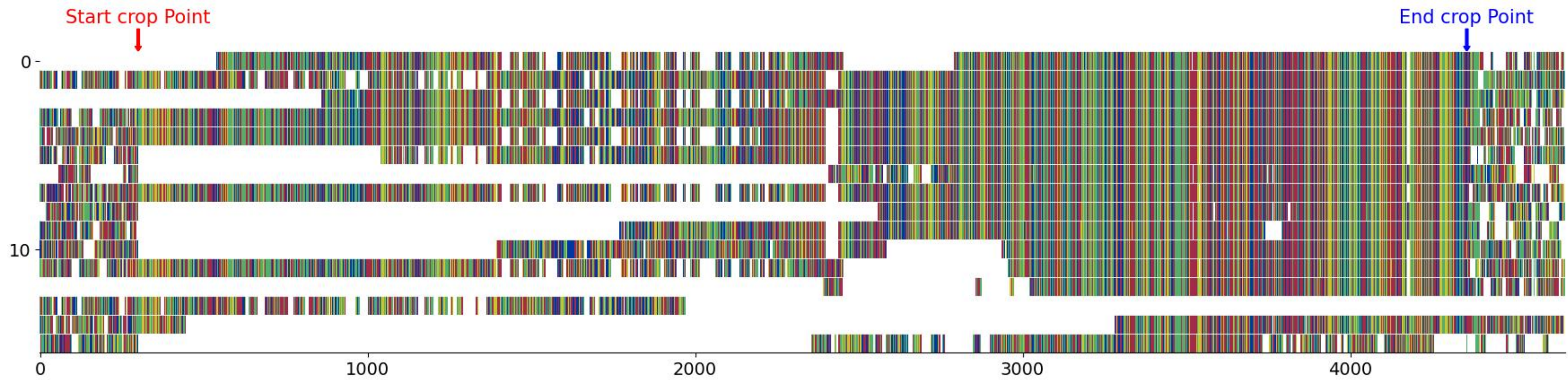


MSA length = 4057



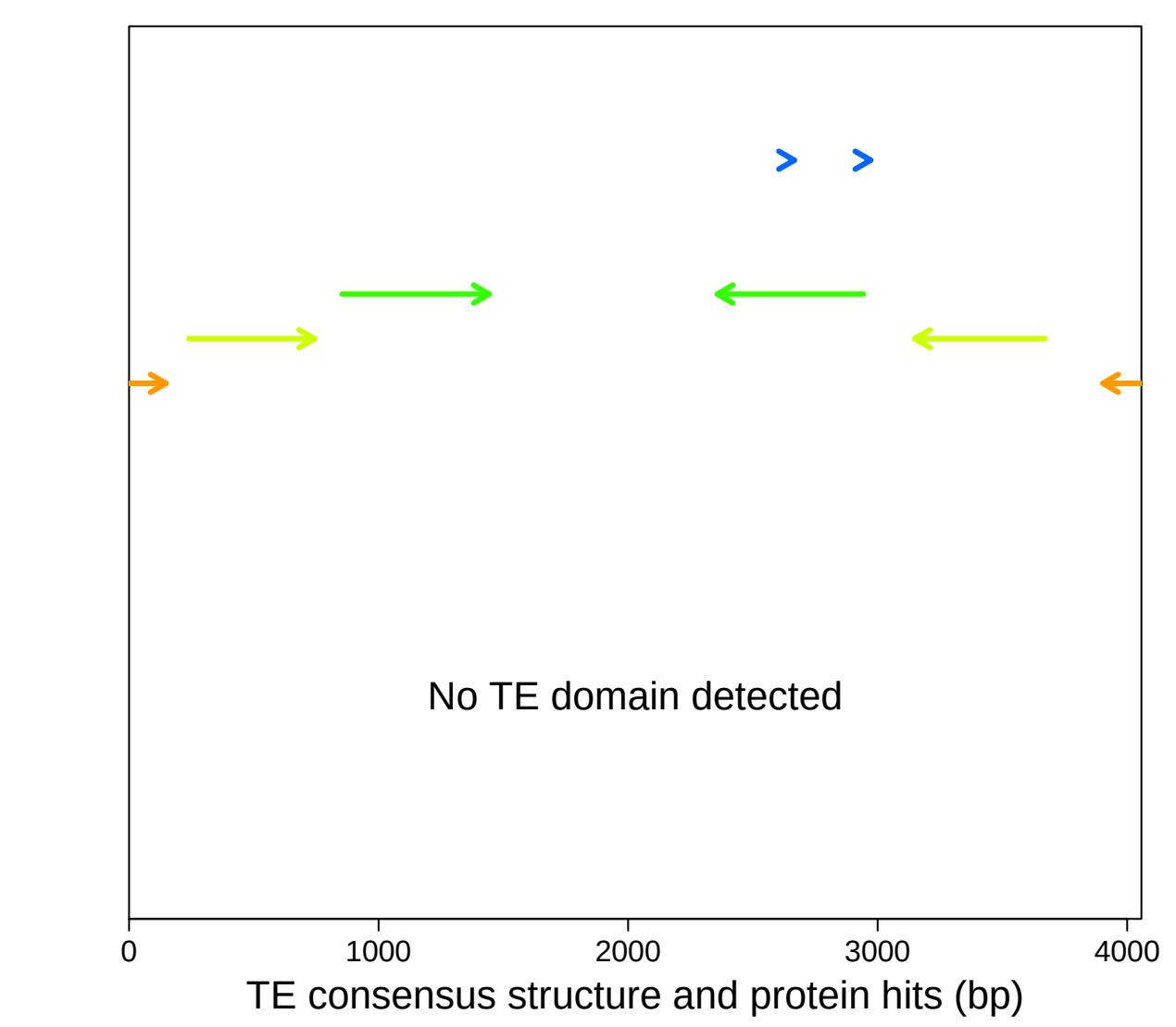
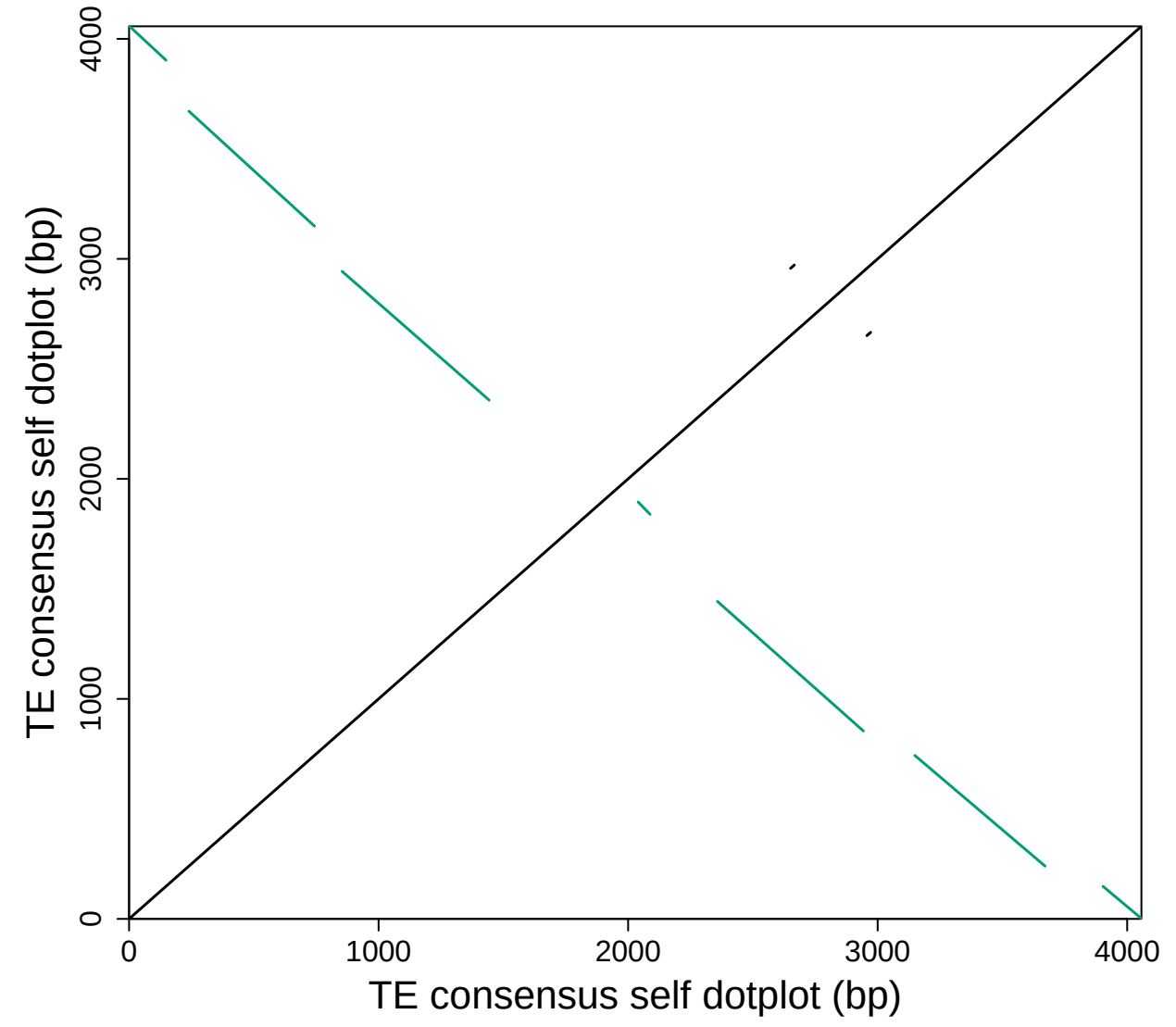
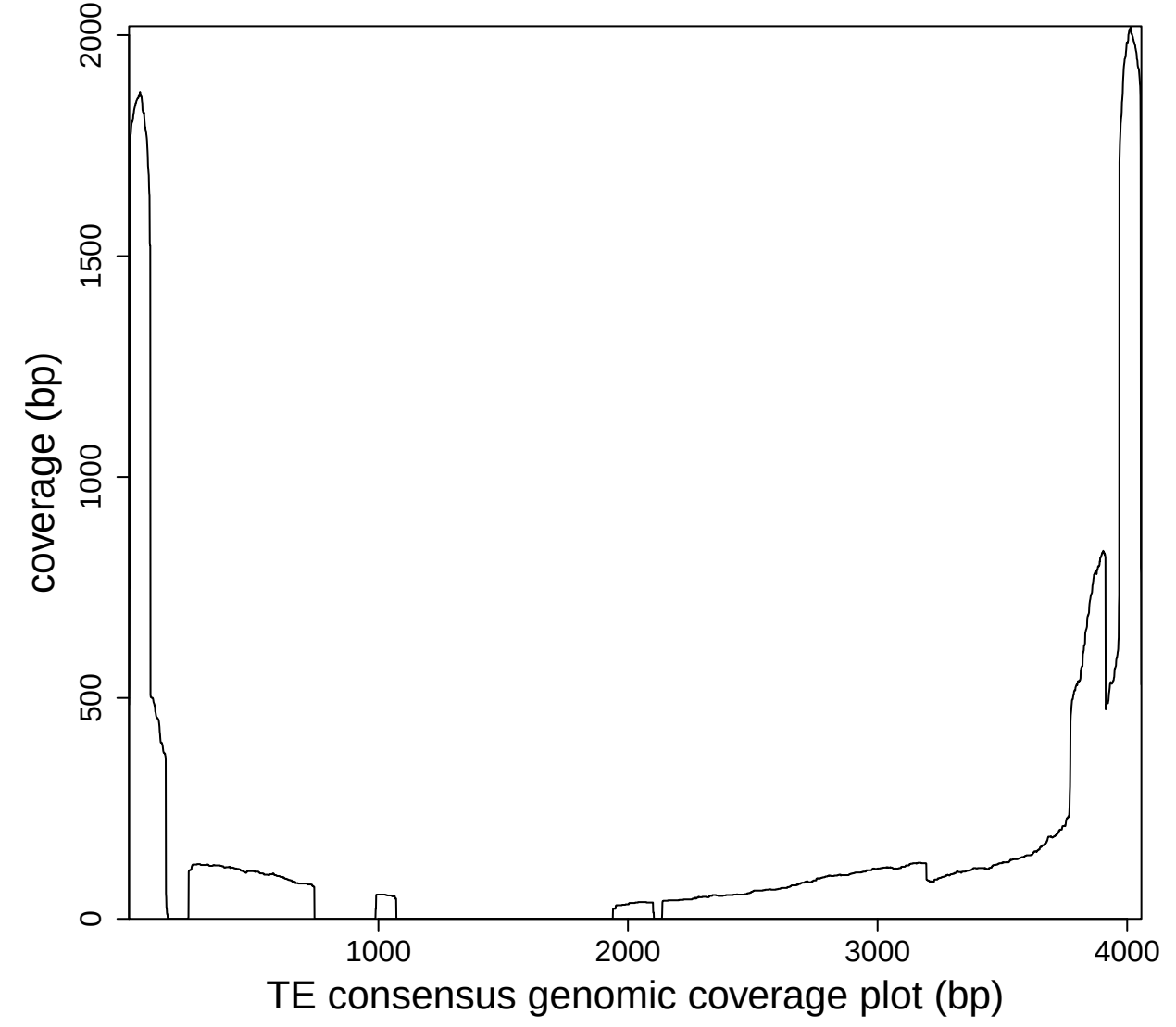
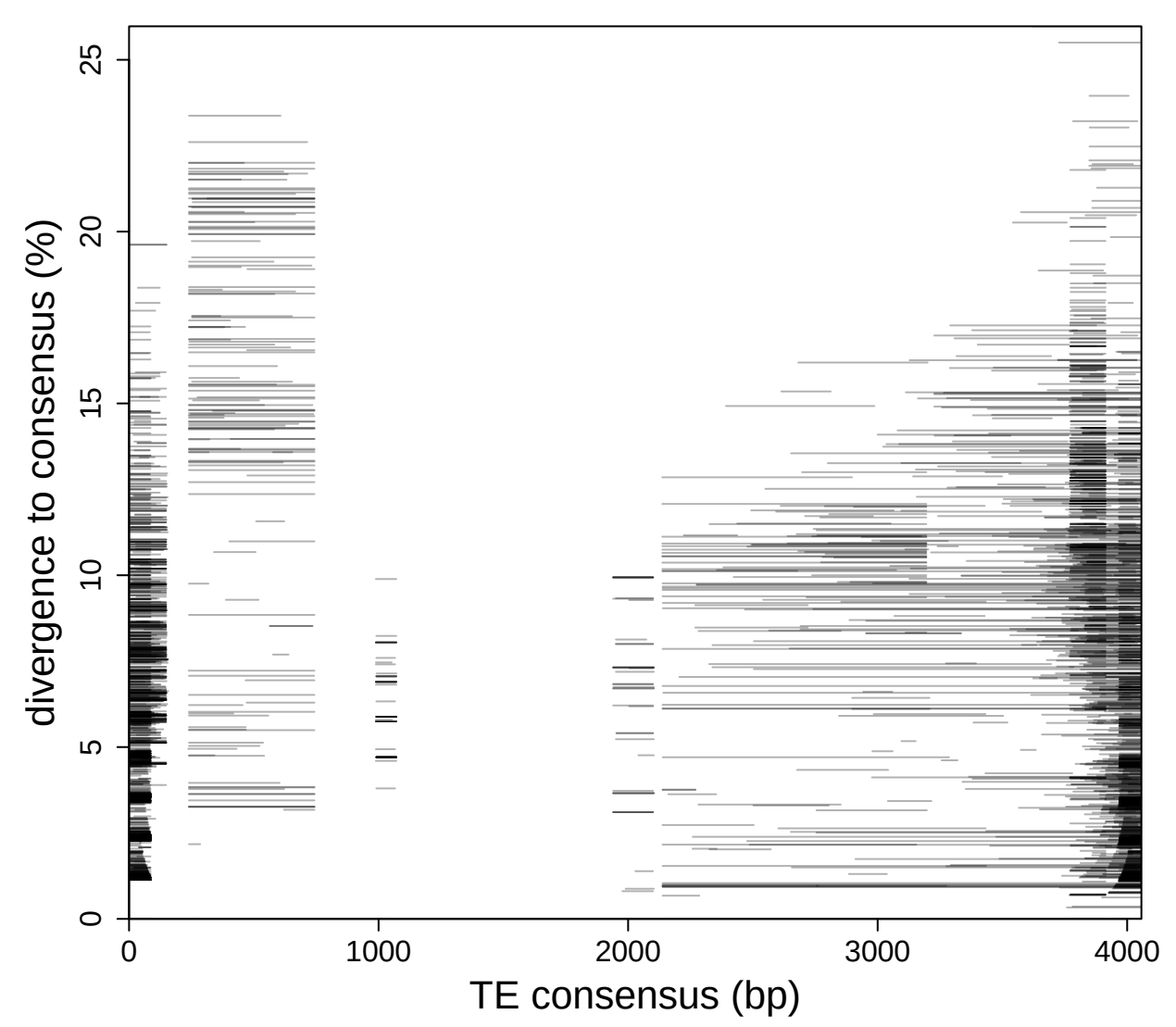
Start crop Point

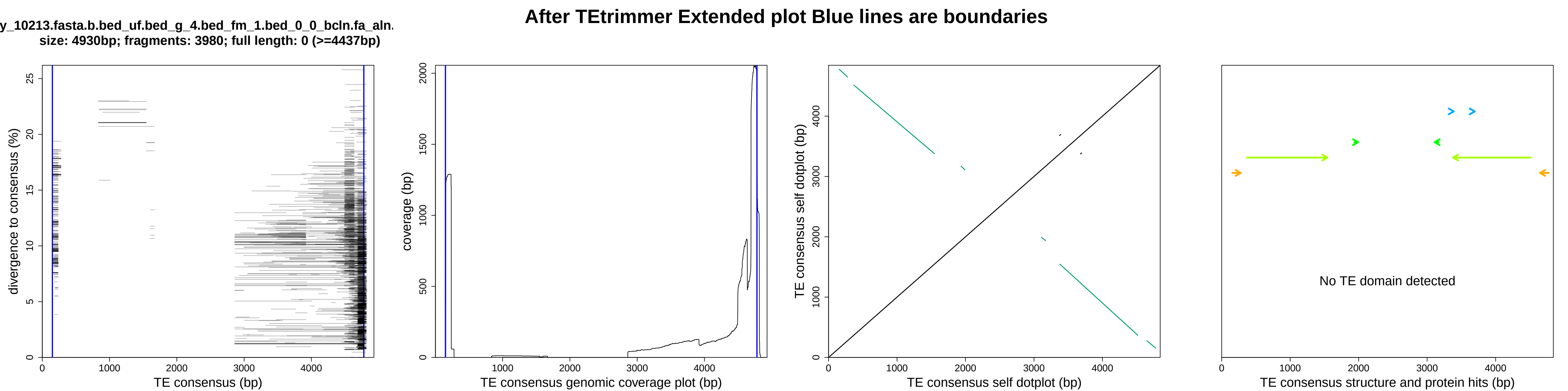
End crop Point



bed_uf.bed g_4.bed fm_1.bed 0_0 bcIn.fa aln.fa cl.fa gs.fa c
size: 4057bp; fragments: 4687; full length: 0 (>=3651.3bp)

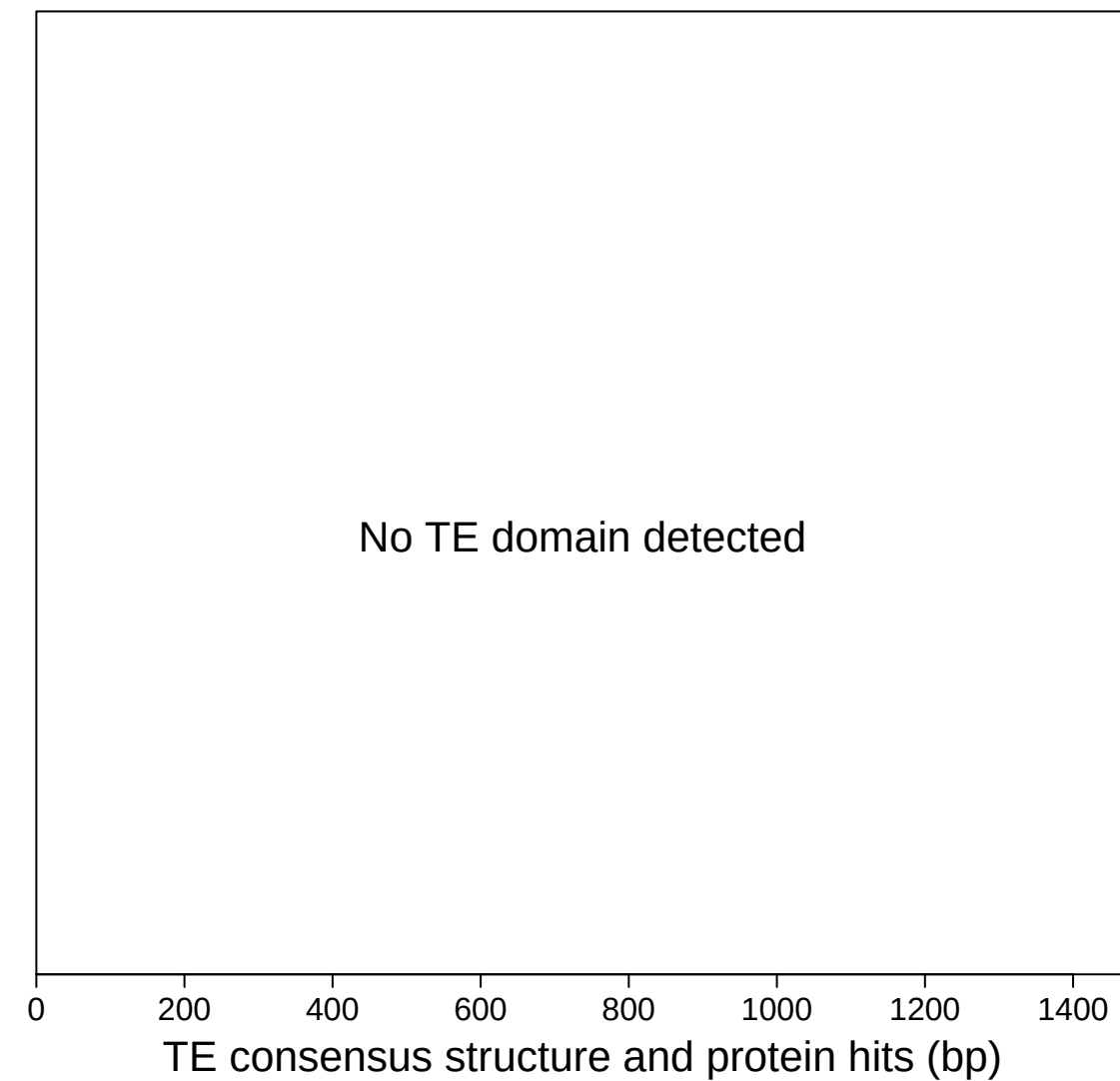
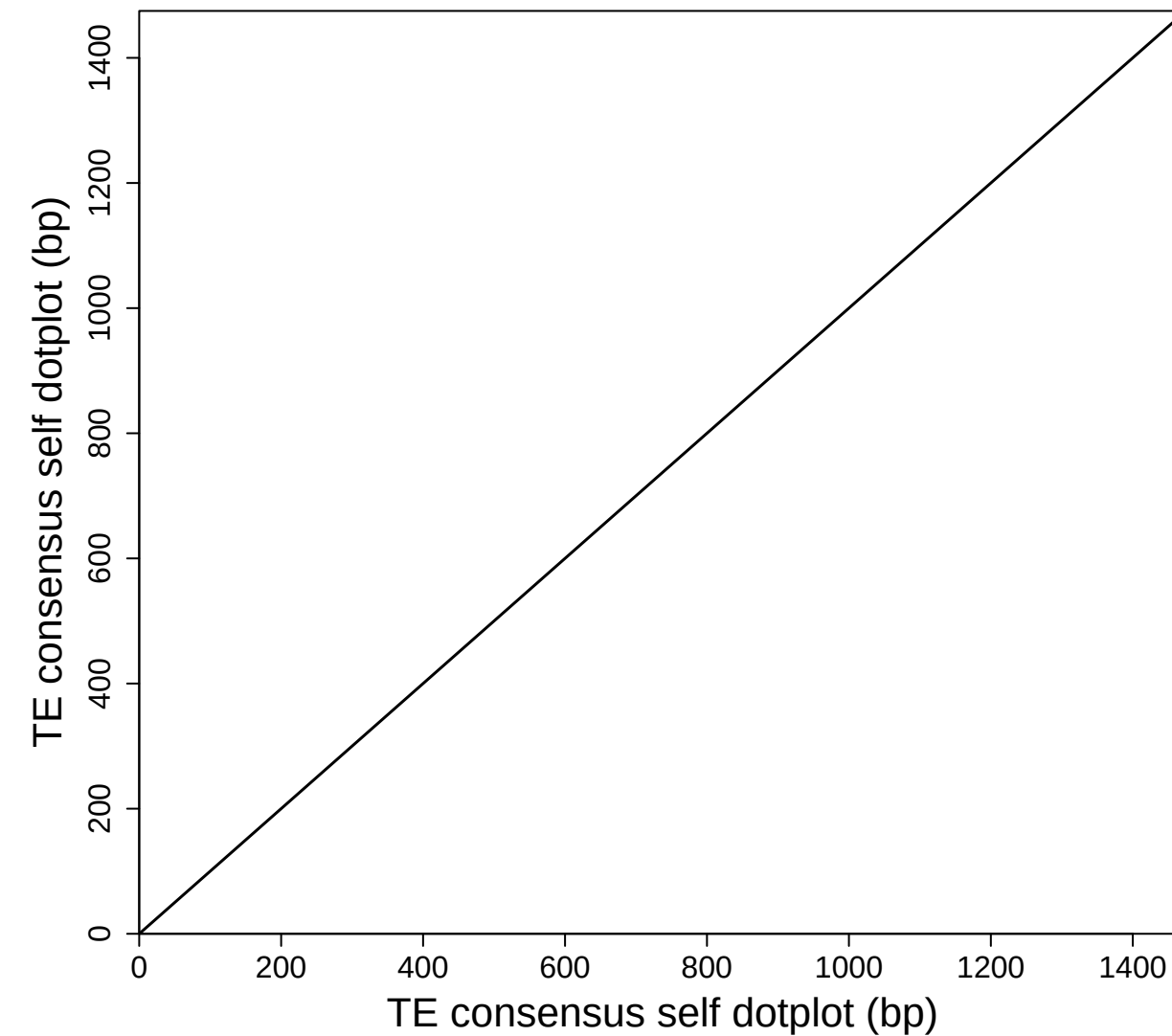
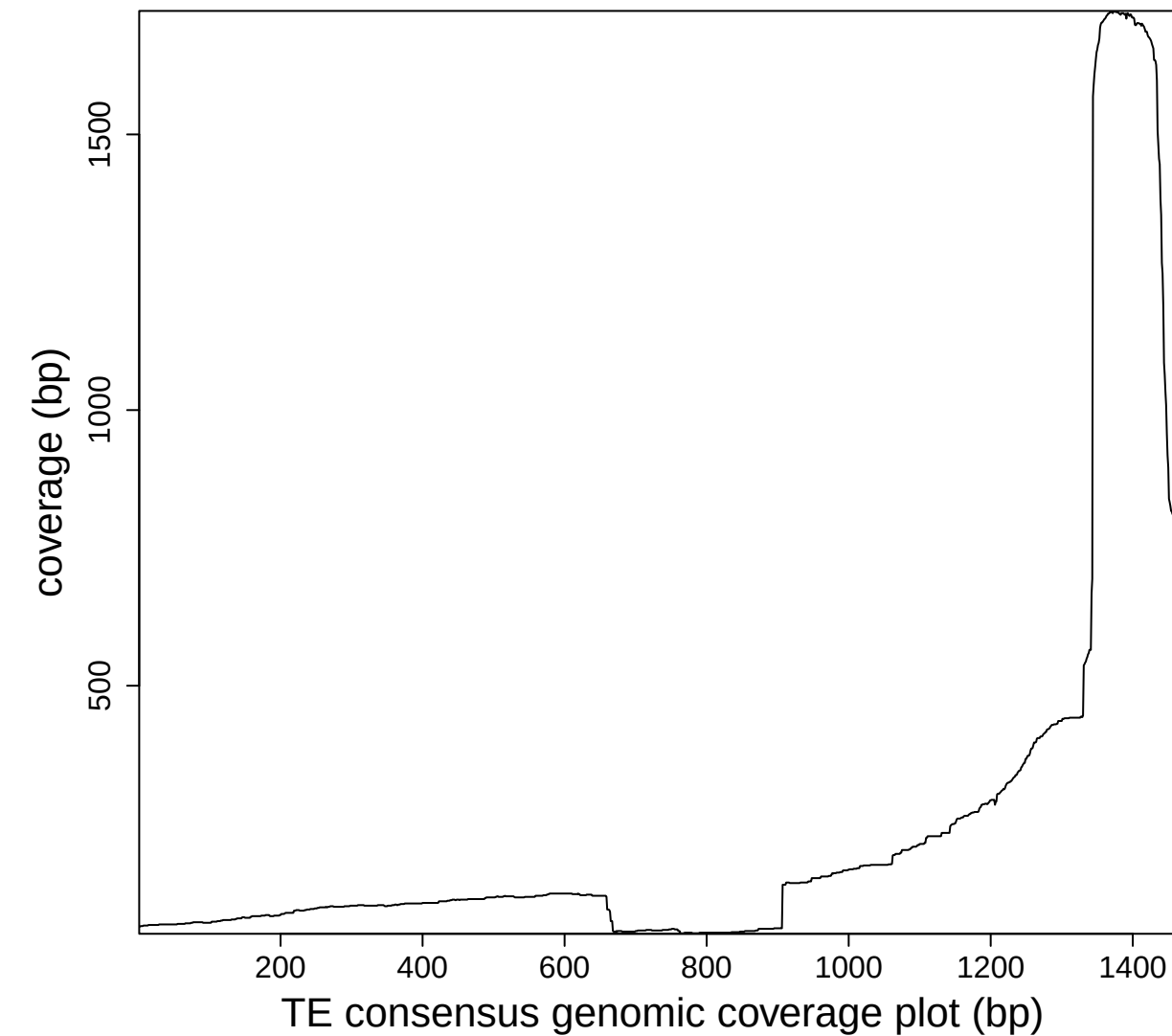
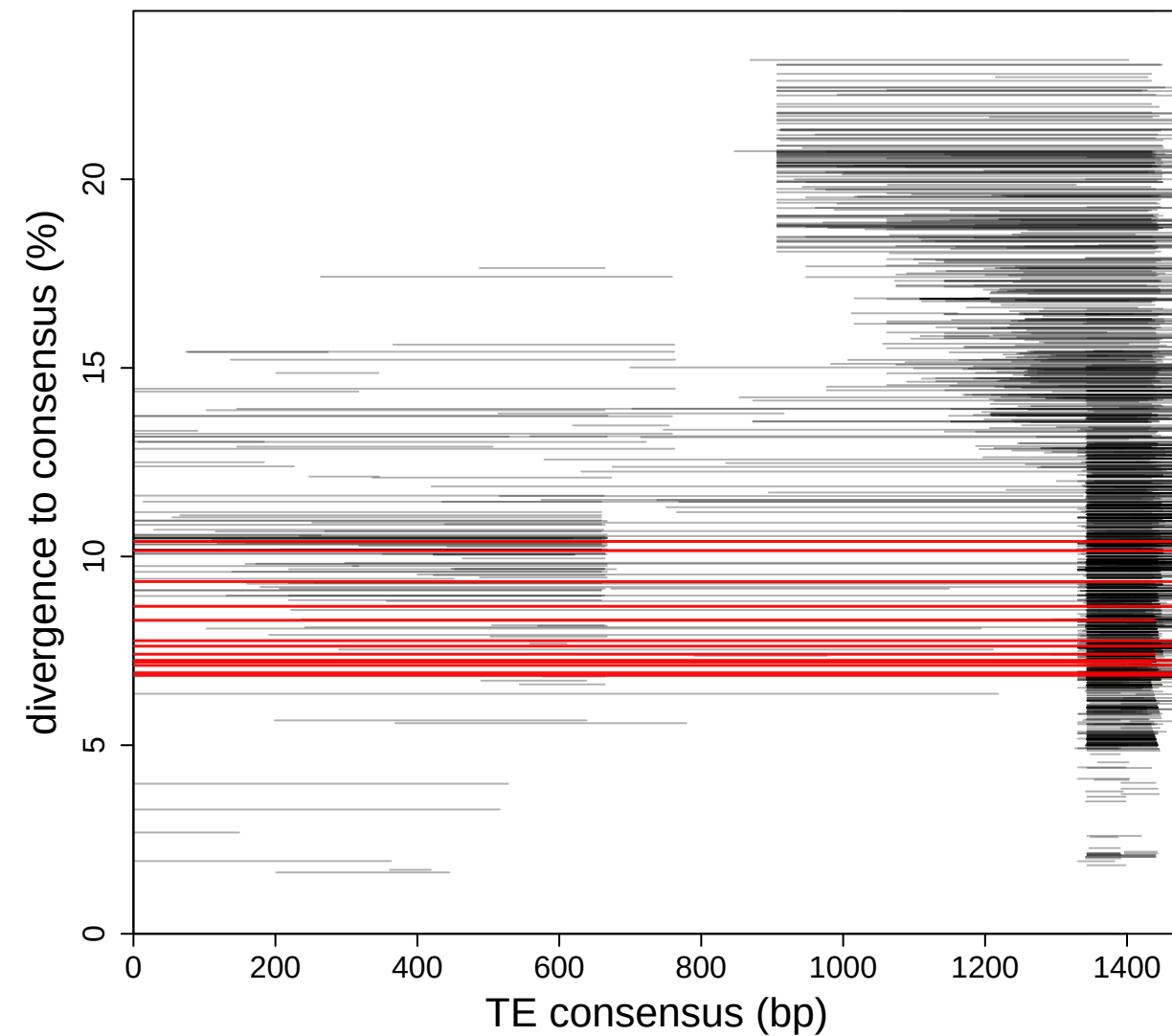
After TETrimmer 4057 bp



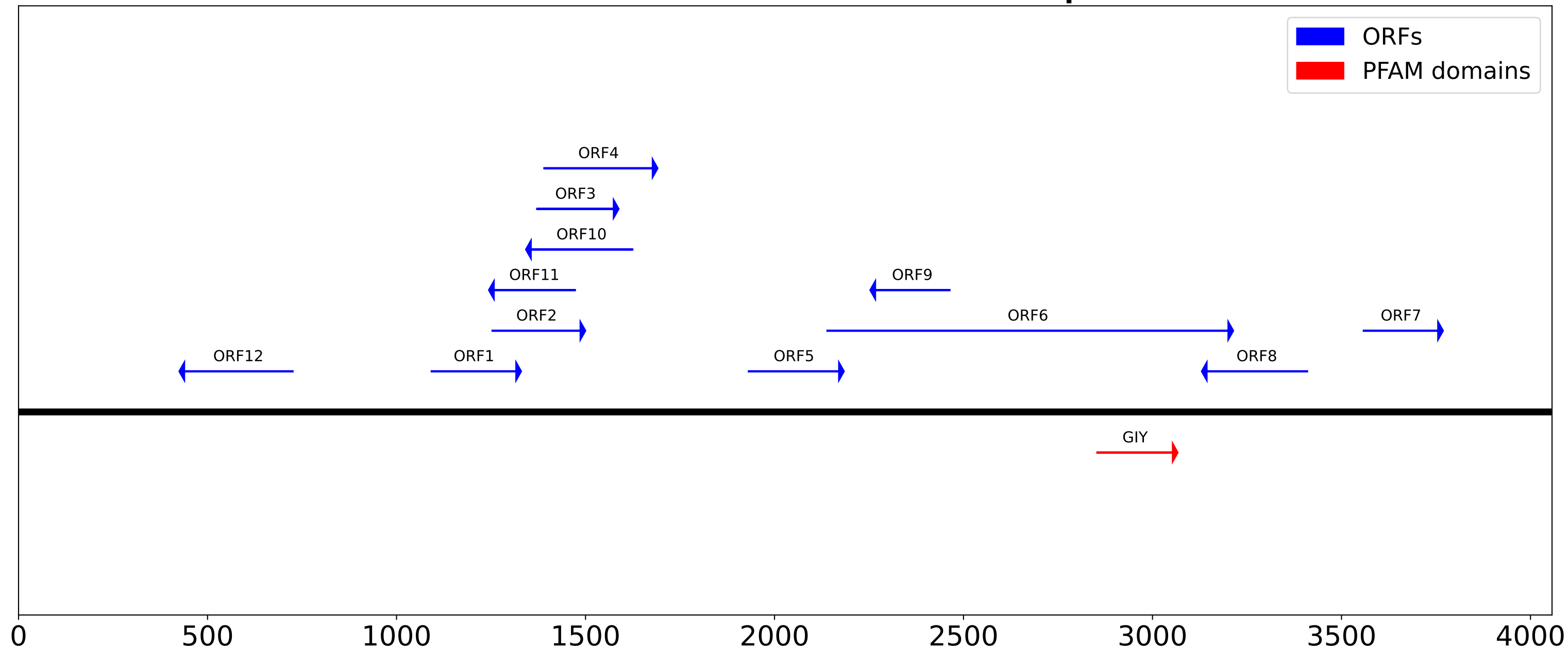


TE: rnd_5_family_10213
size: 1475bp; fragments: 1888; full length: 13 (≥ 1327.5 bp)

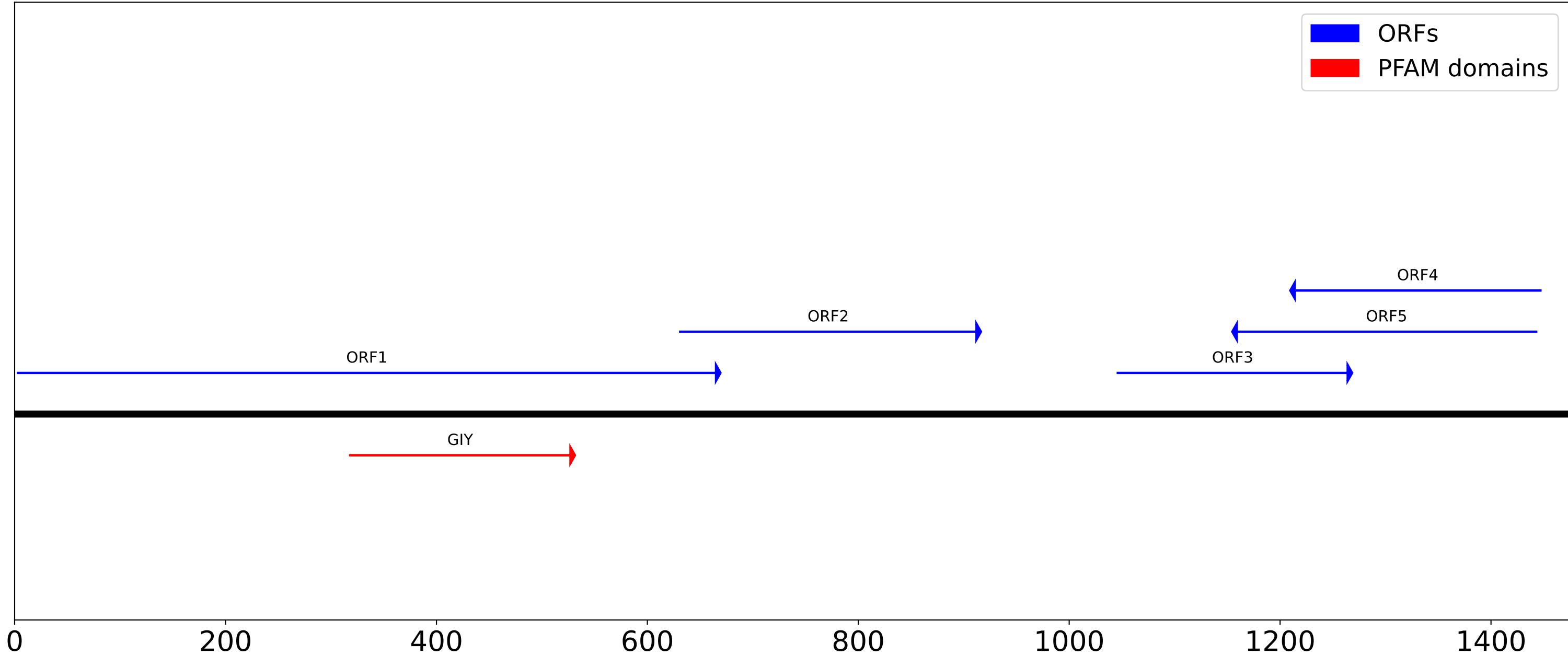
Before TETrimmer 1475 bp



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)

