

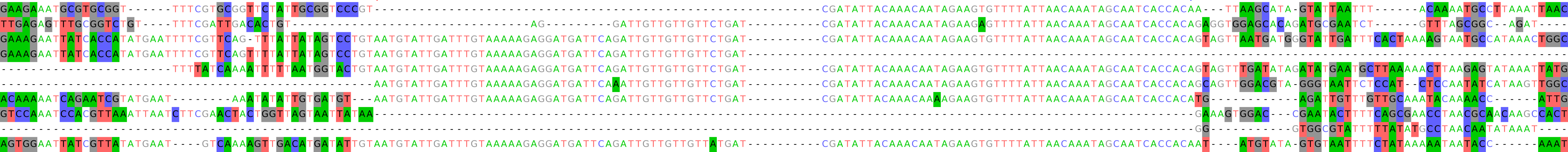
Start Crop Point



End Crop Point

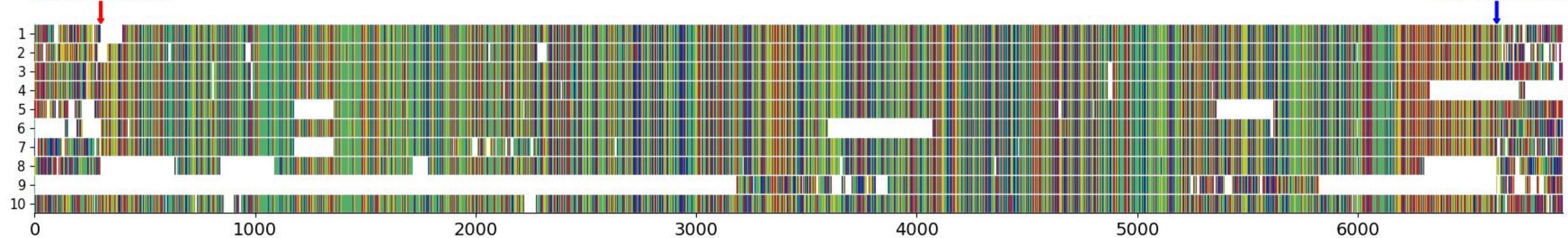


MSA length = 6331



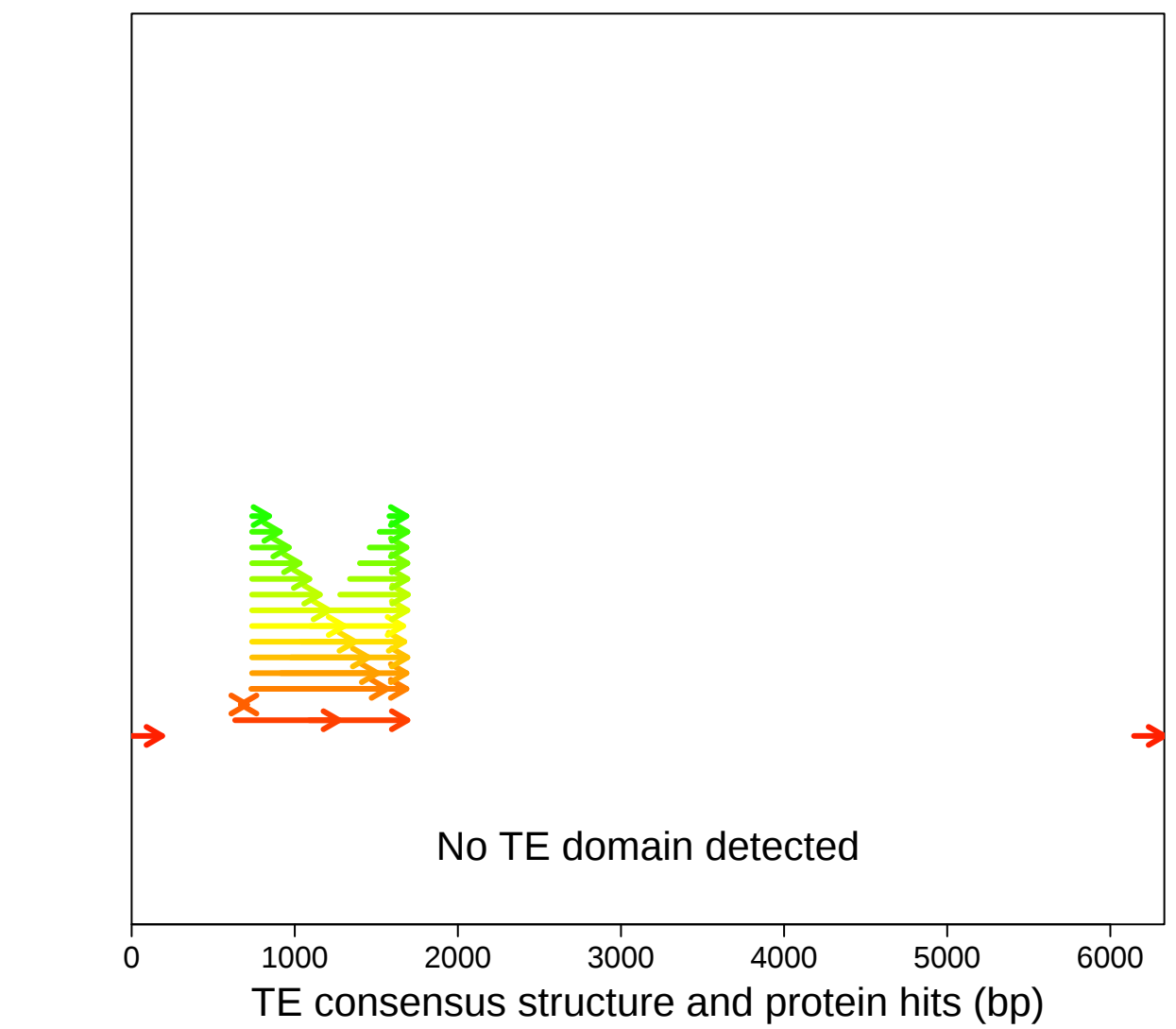
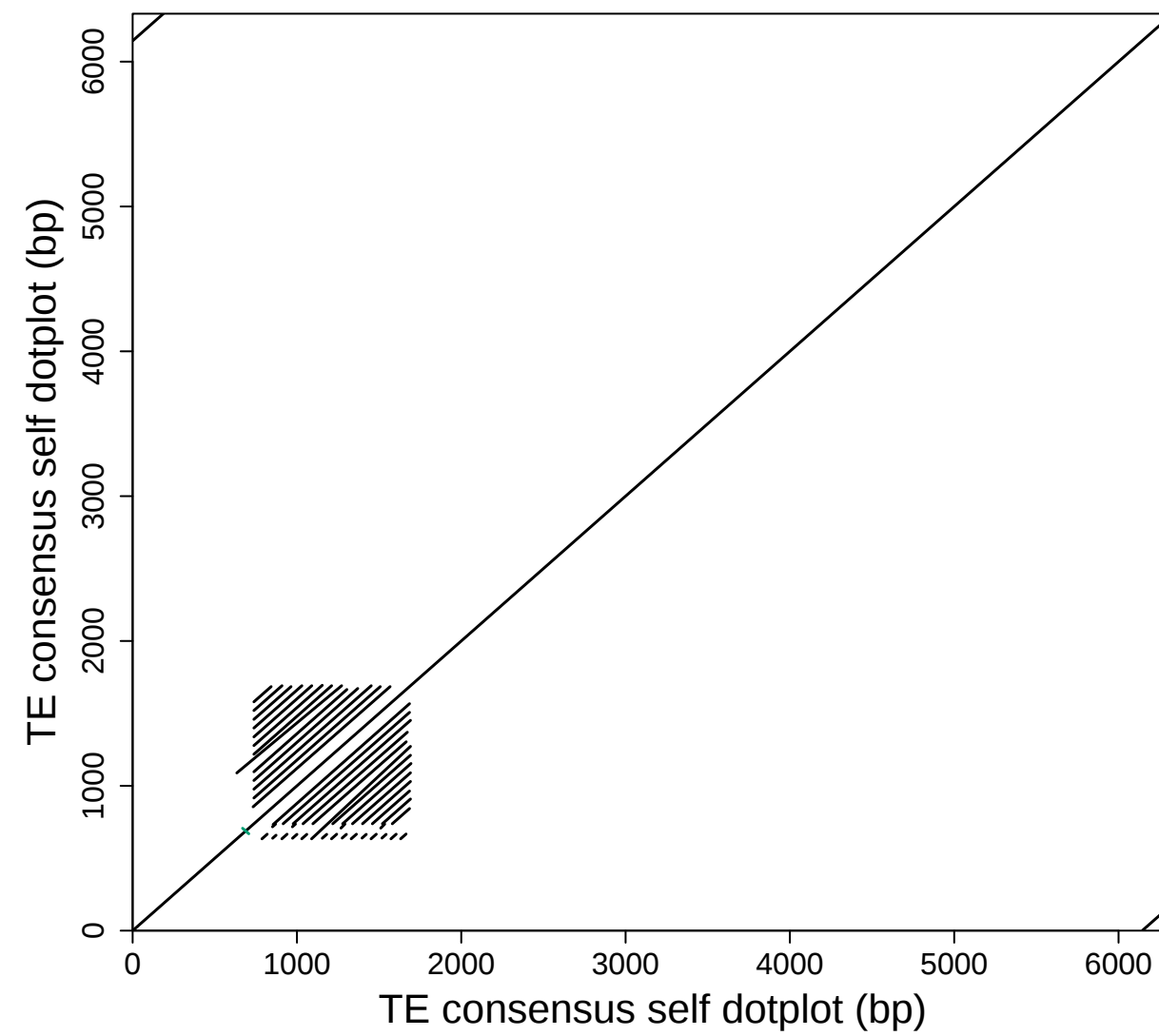
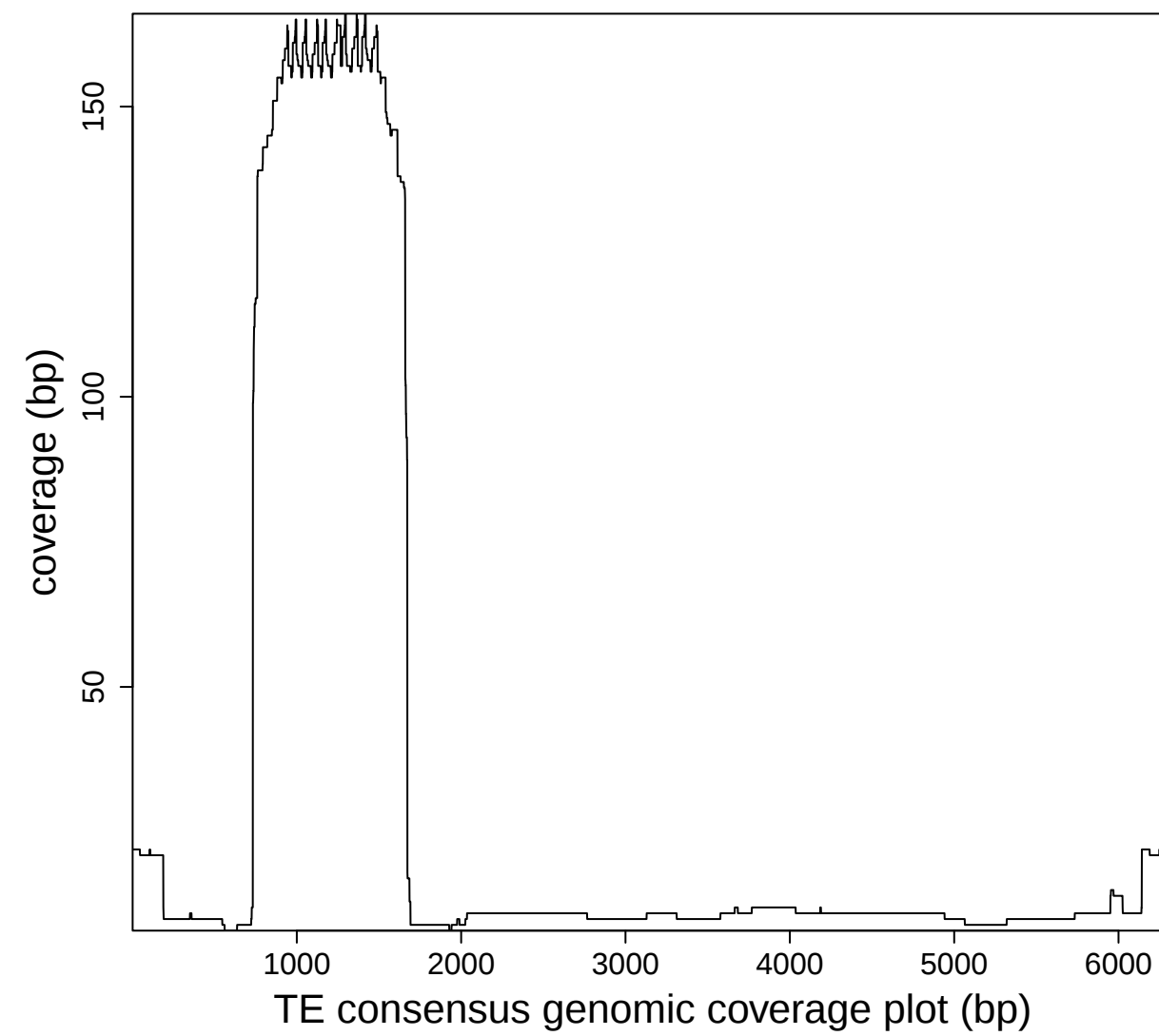
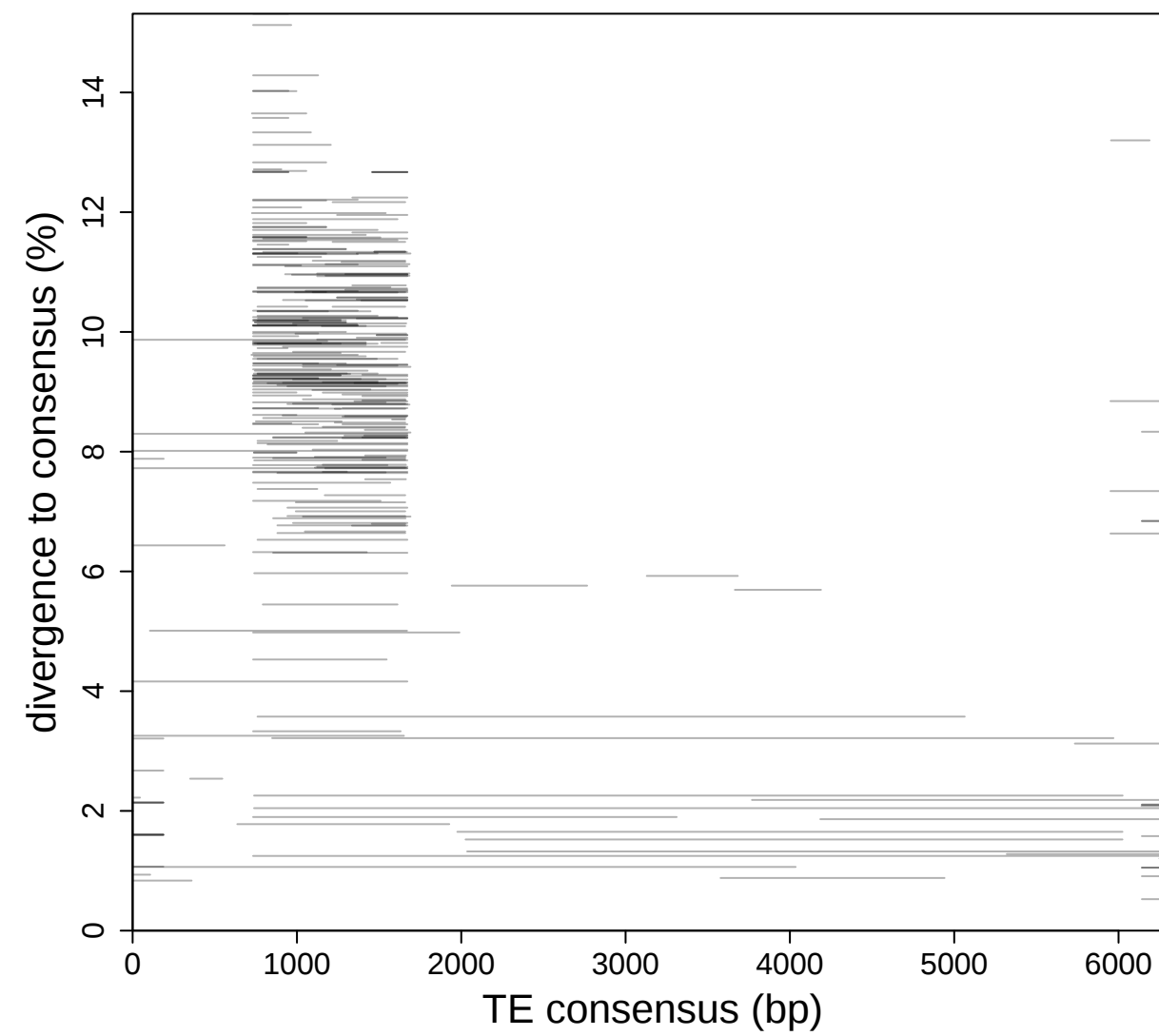
Start crop Point

End crop Point



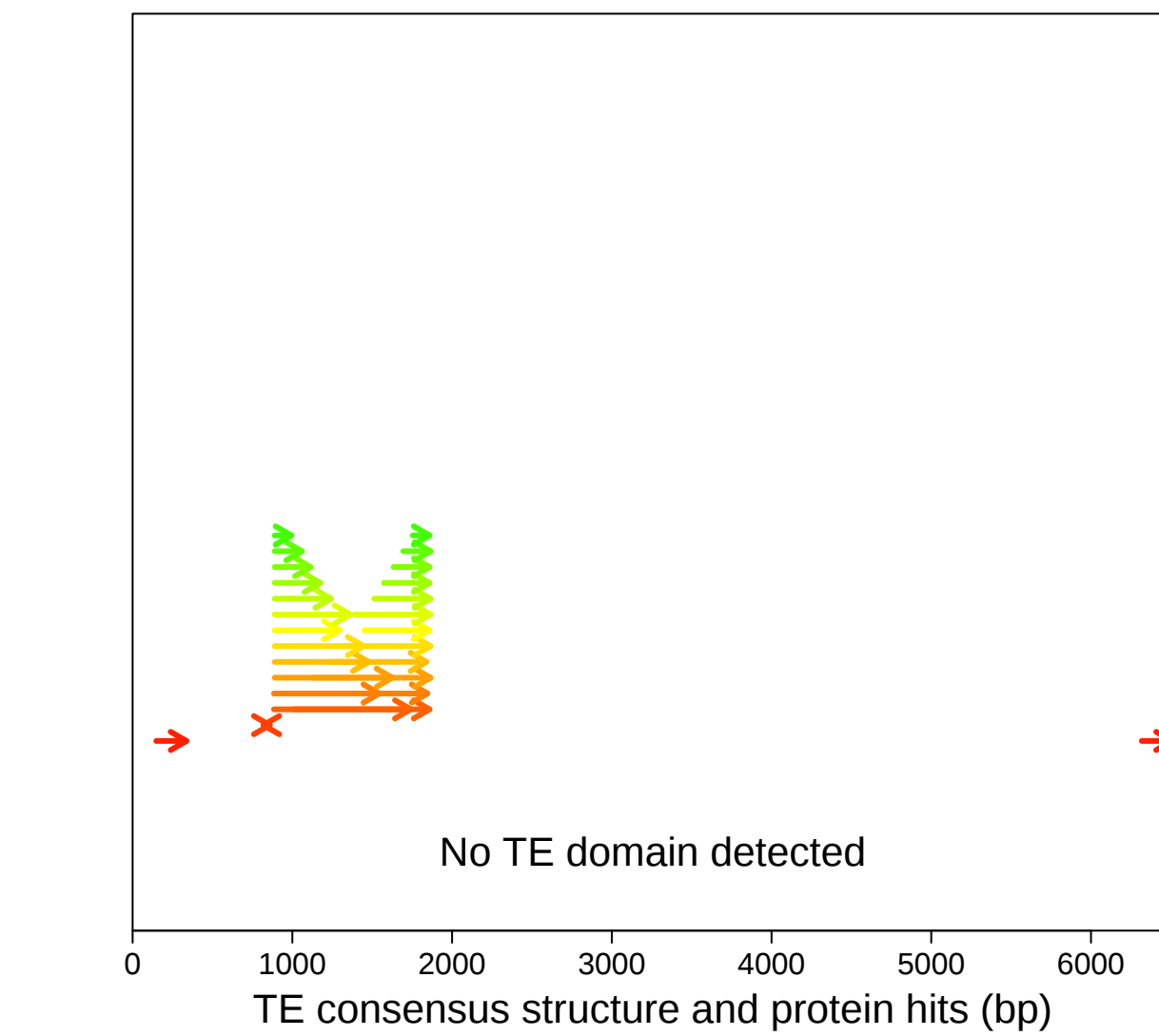
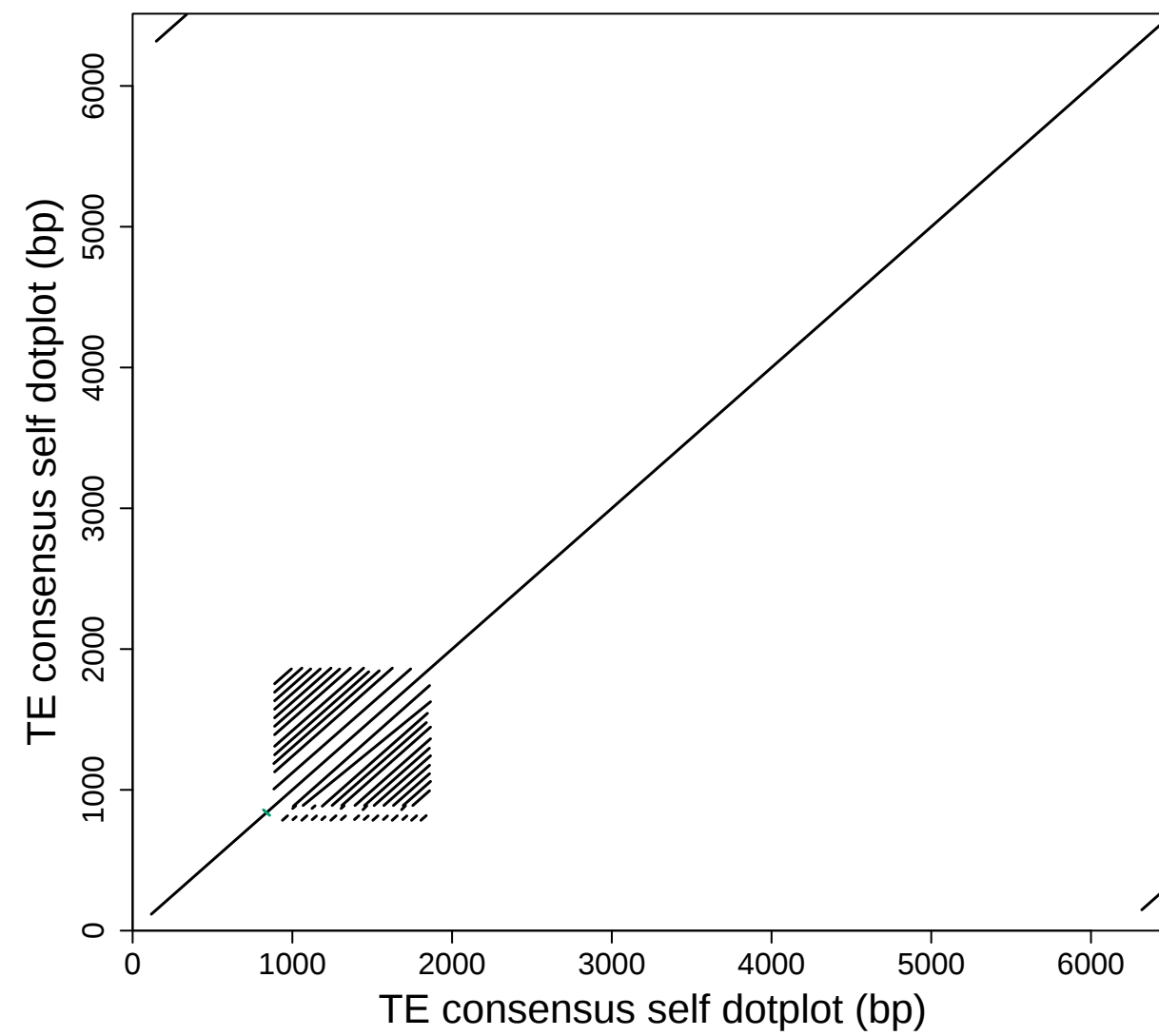
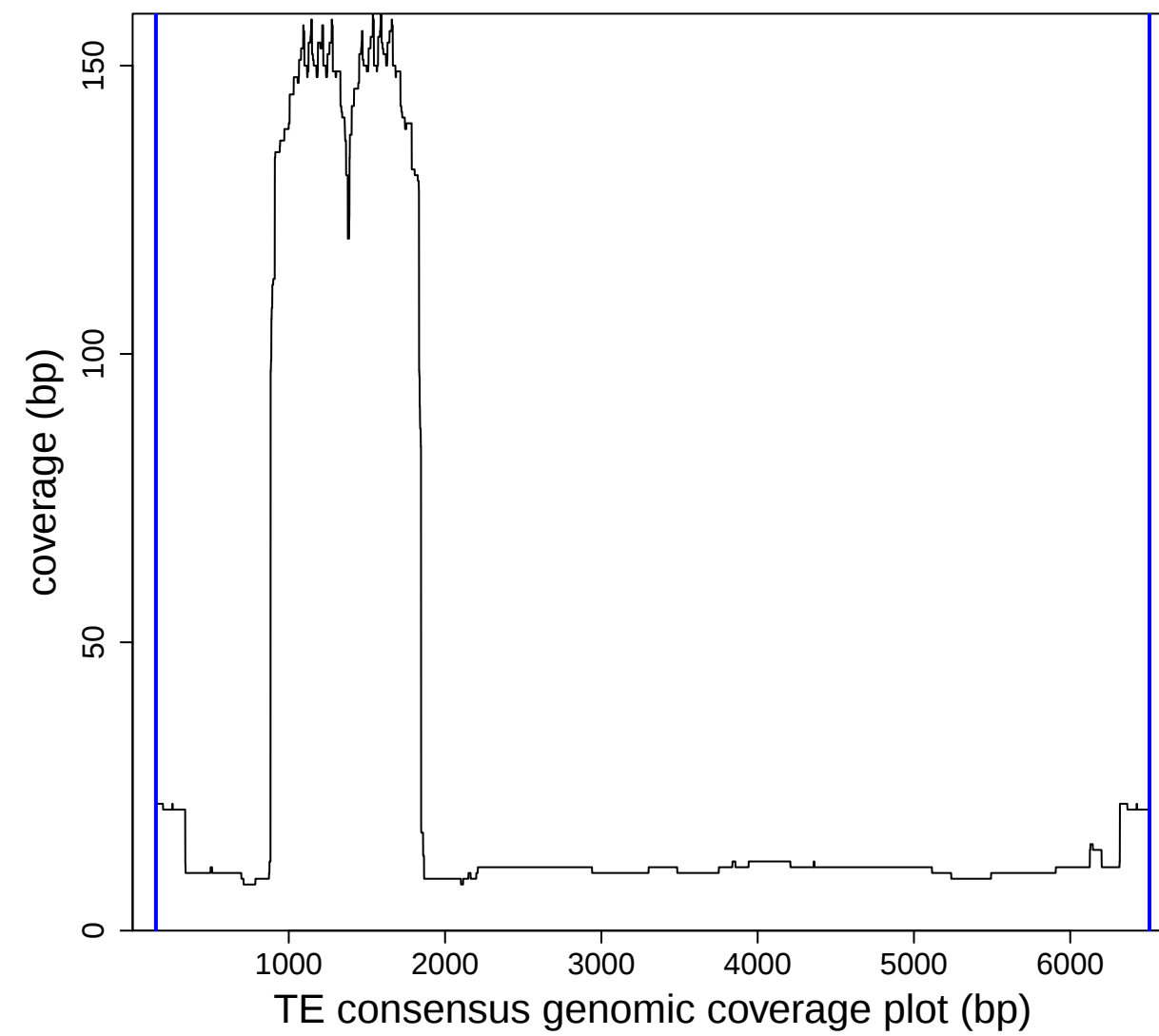
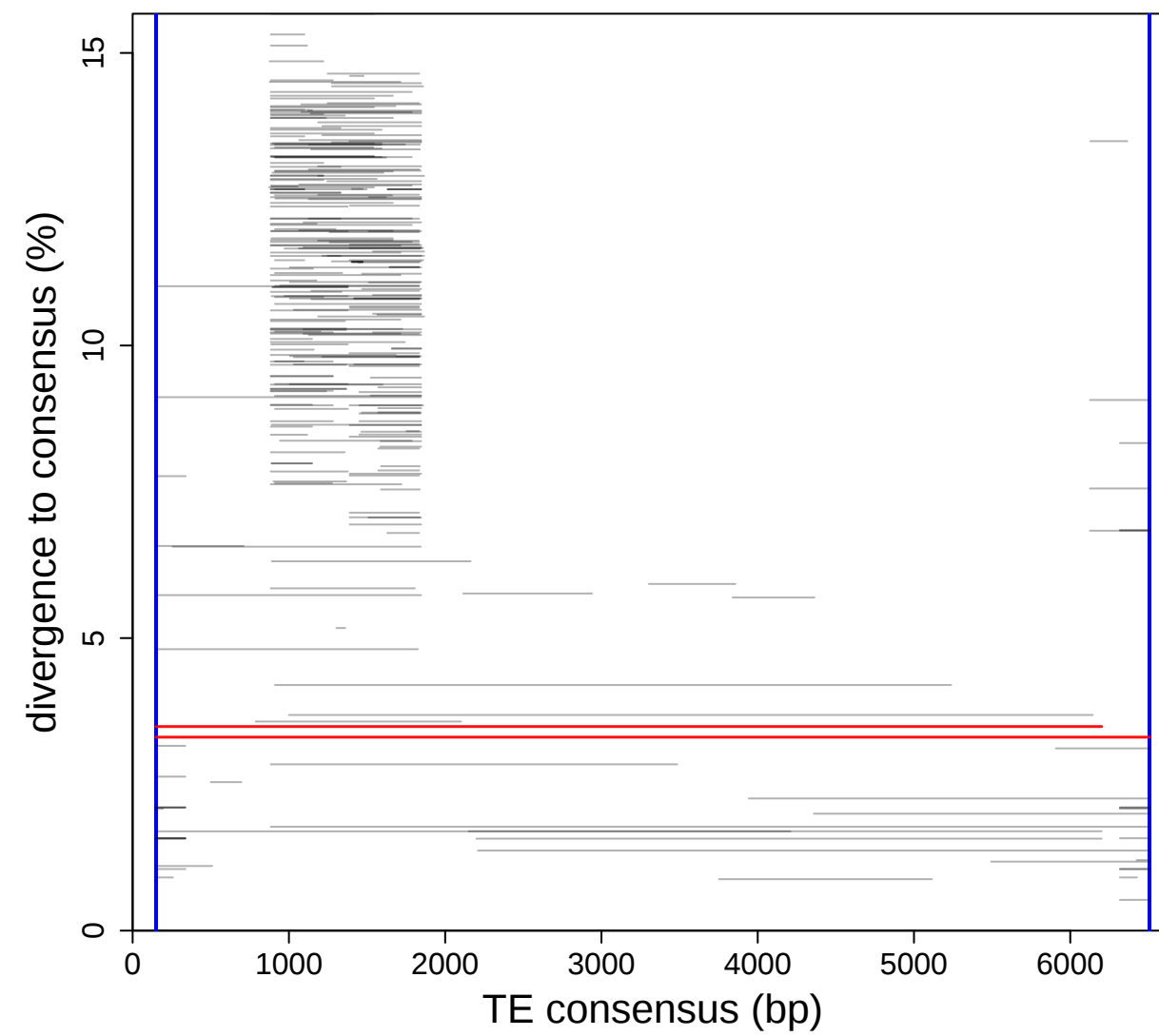
ed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 6331bp; fragments: 298; full length: 0 (>=5697.9bp)

After TEtrimmer 6331 bp

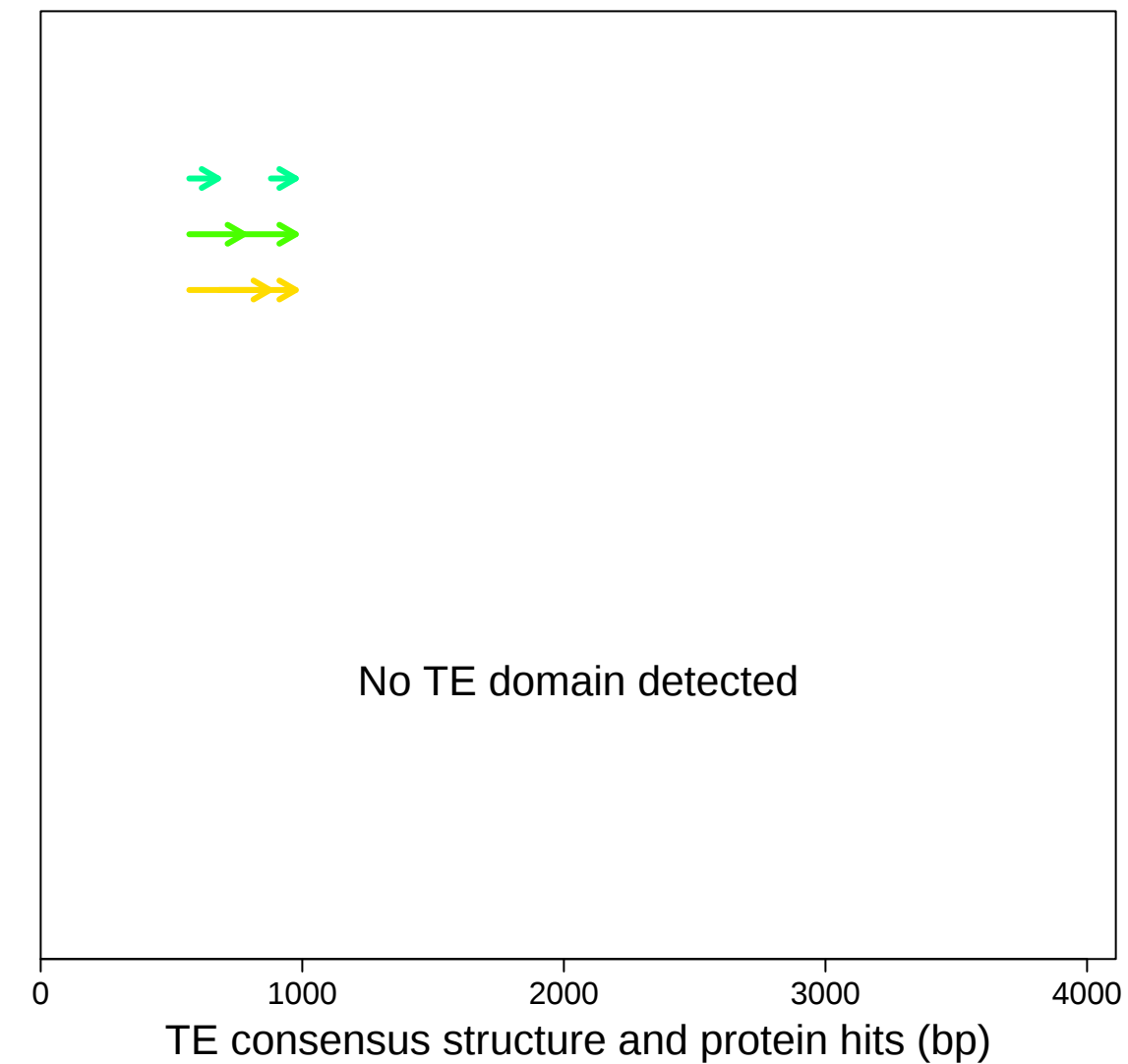
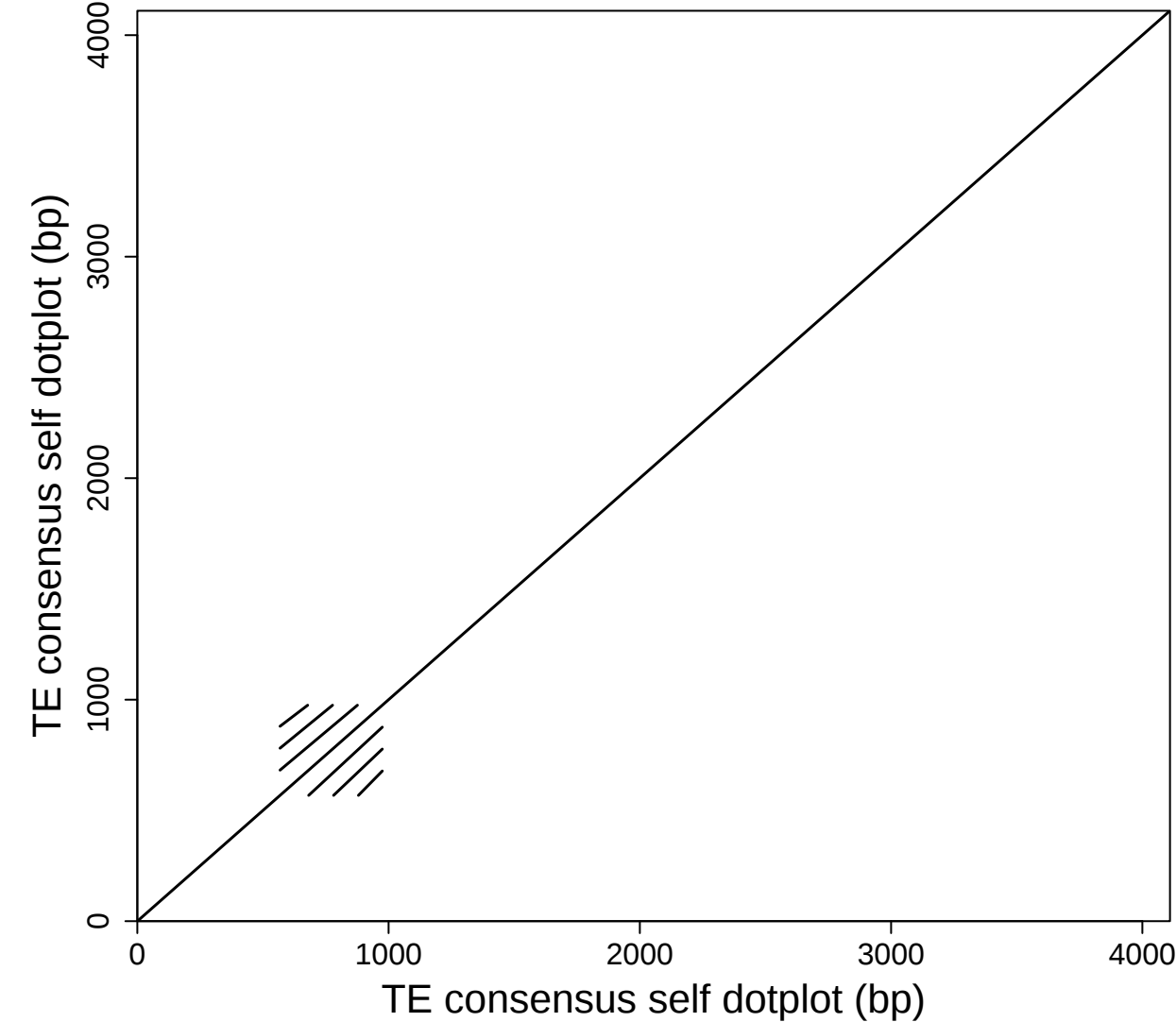
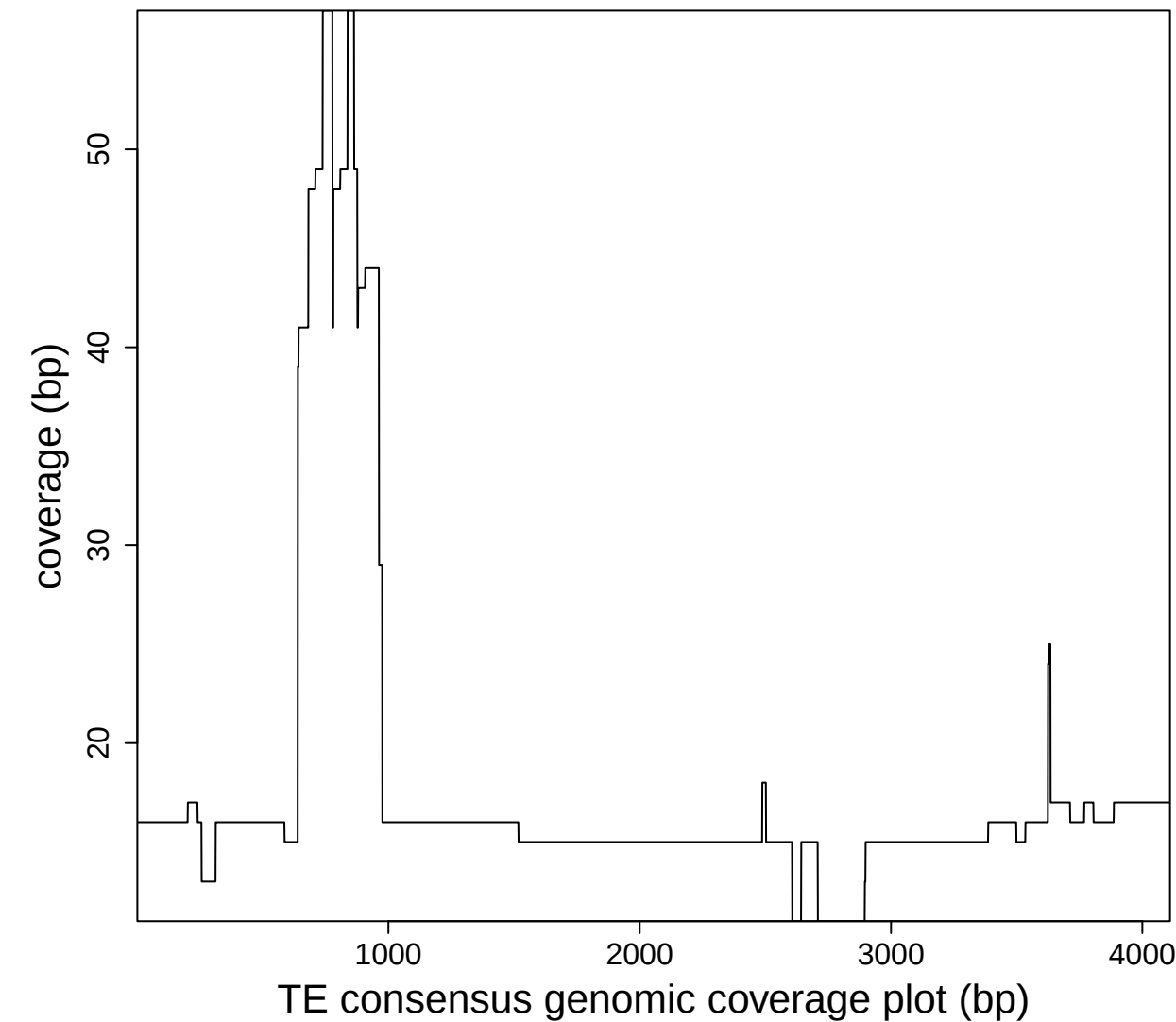
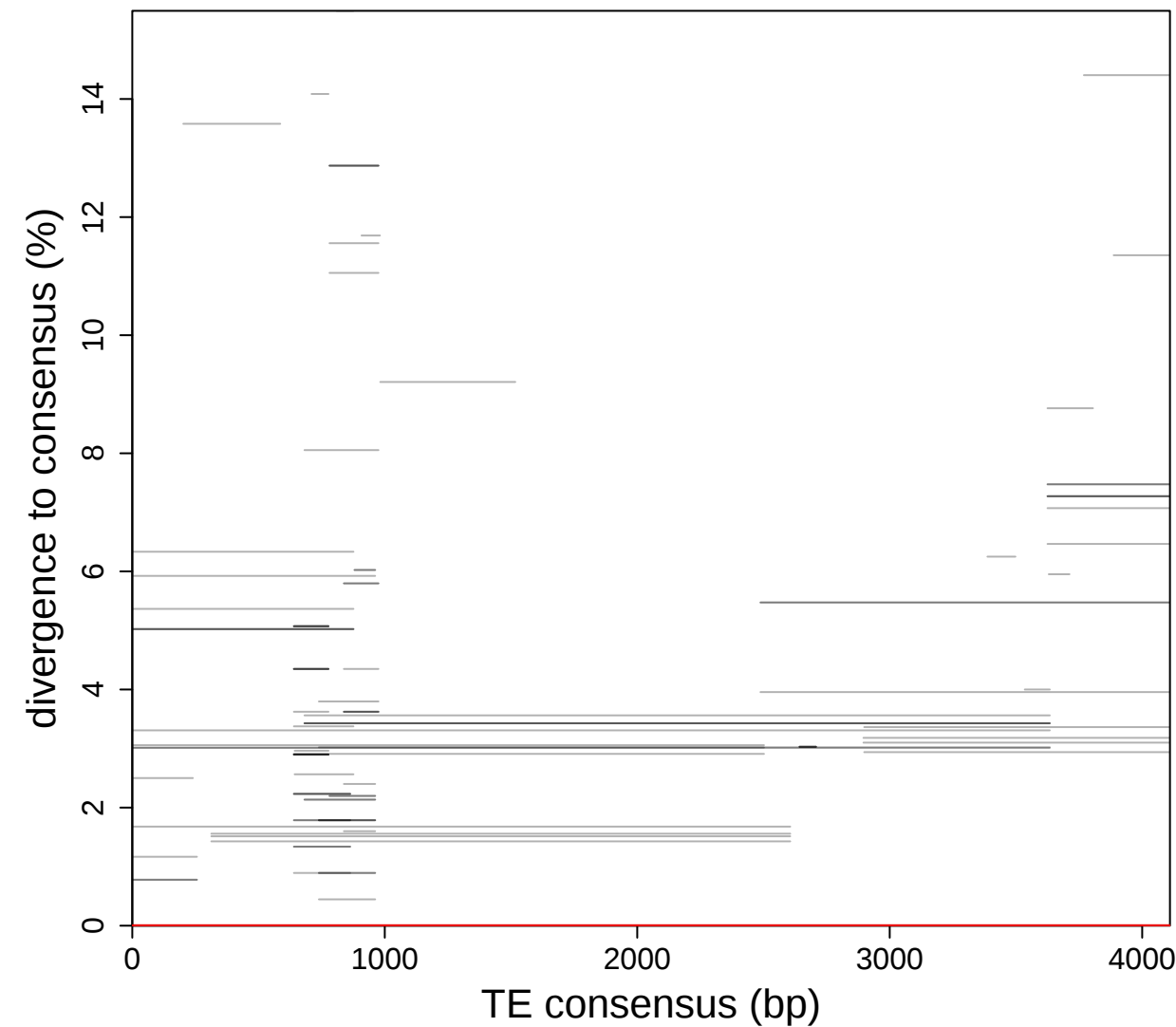


ly_573.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 6657bp; fragments: 298; full length: 2 (>=5991.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

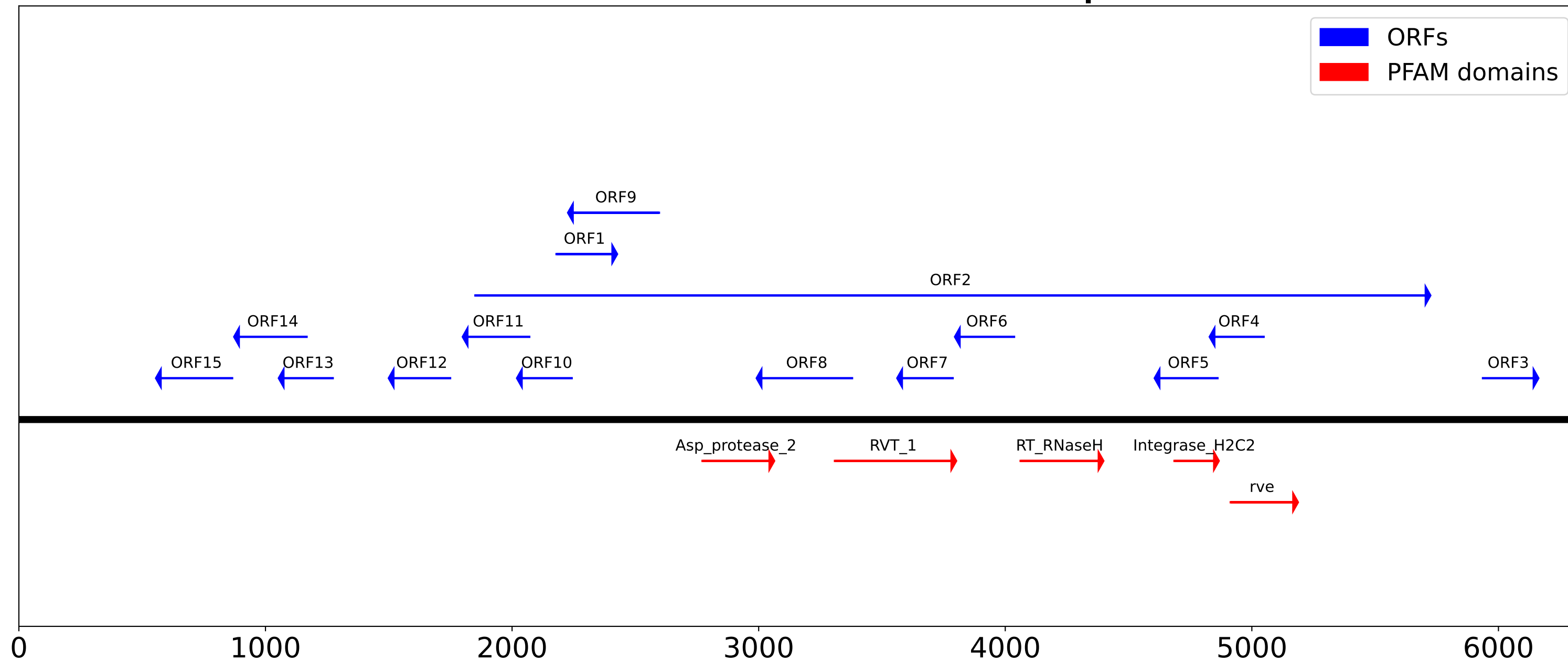


TE: ltr_1_family_573
size: 4110bp; fragments: 106; full length: 1 (>=3699bp)

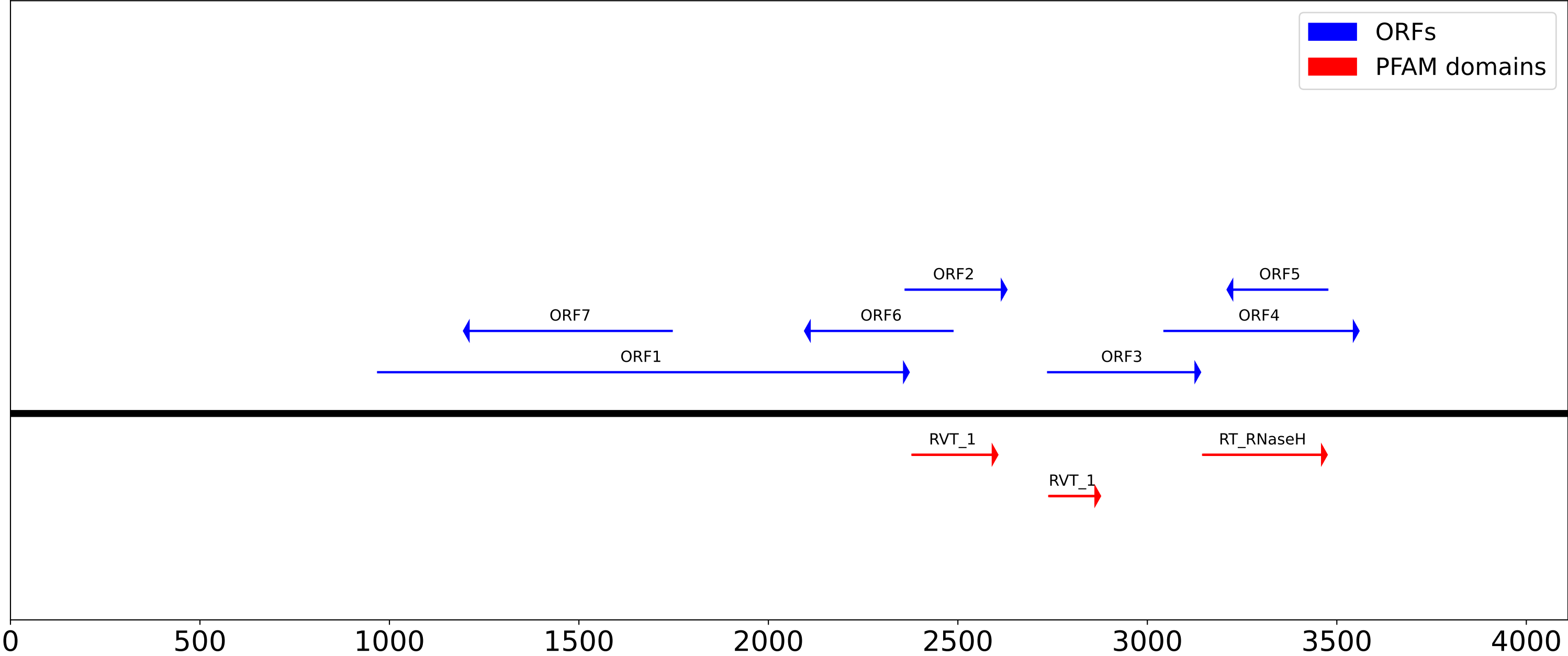


Before TEtrimmer 4110 bp

After TETrimmer ORF and PFAM domain plot



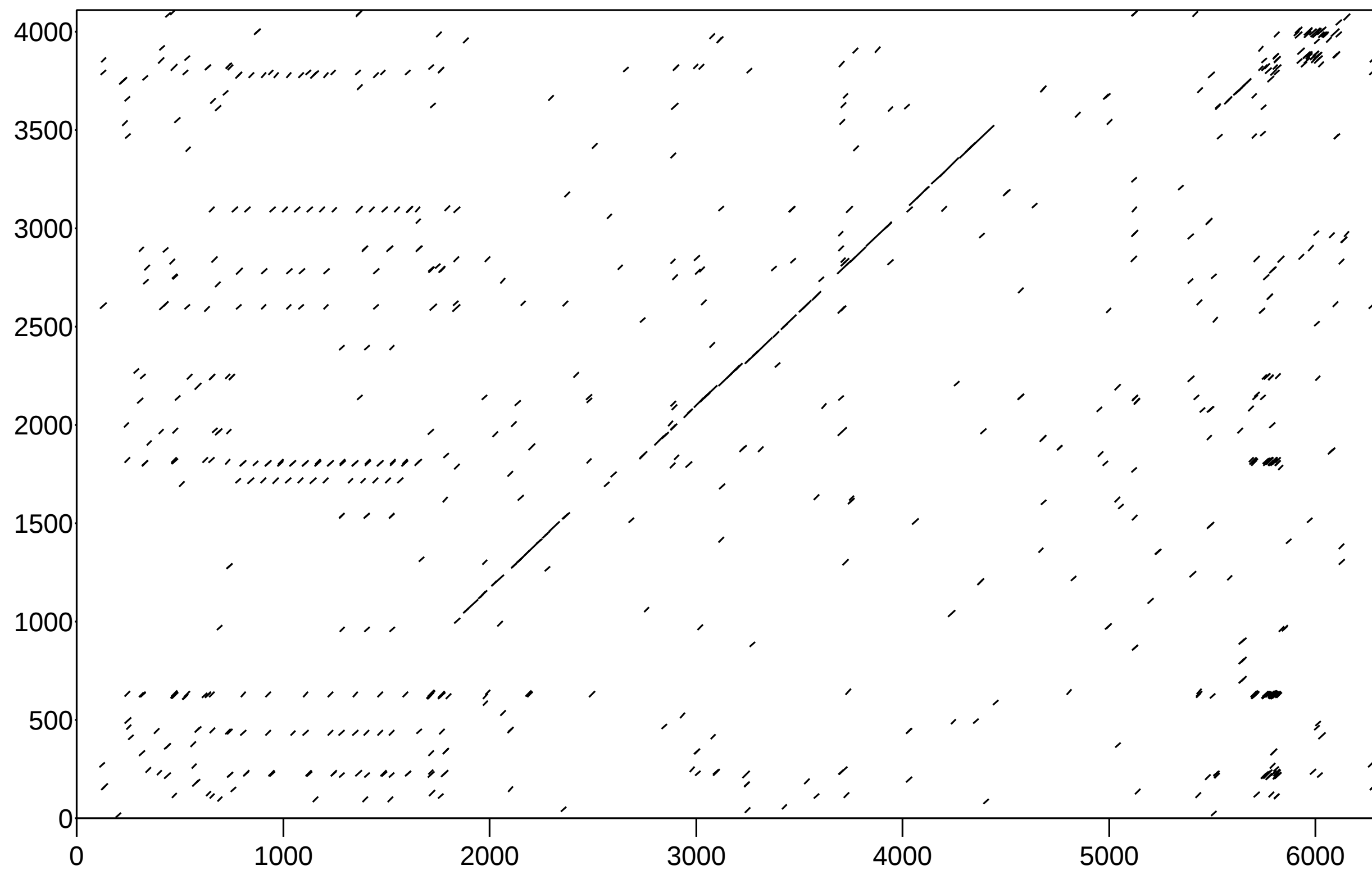
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)