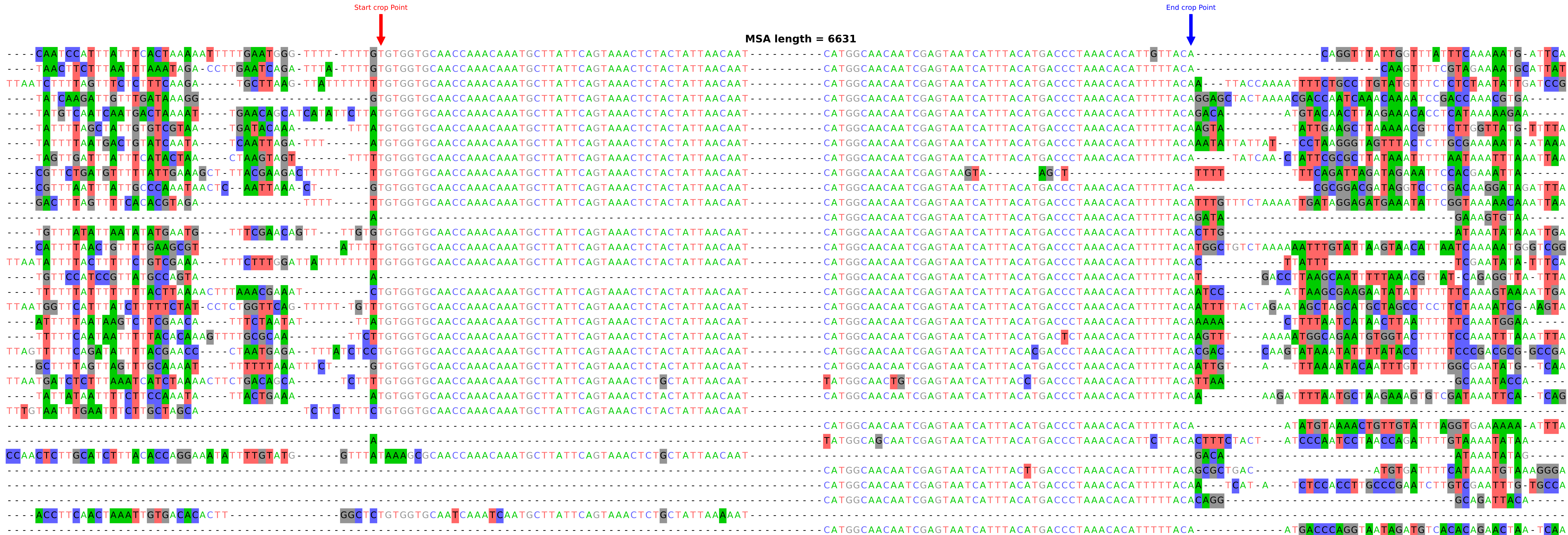
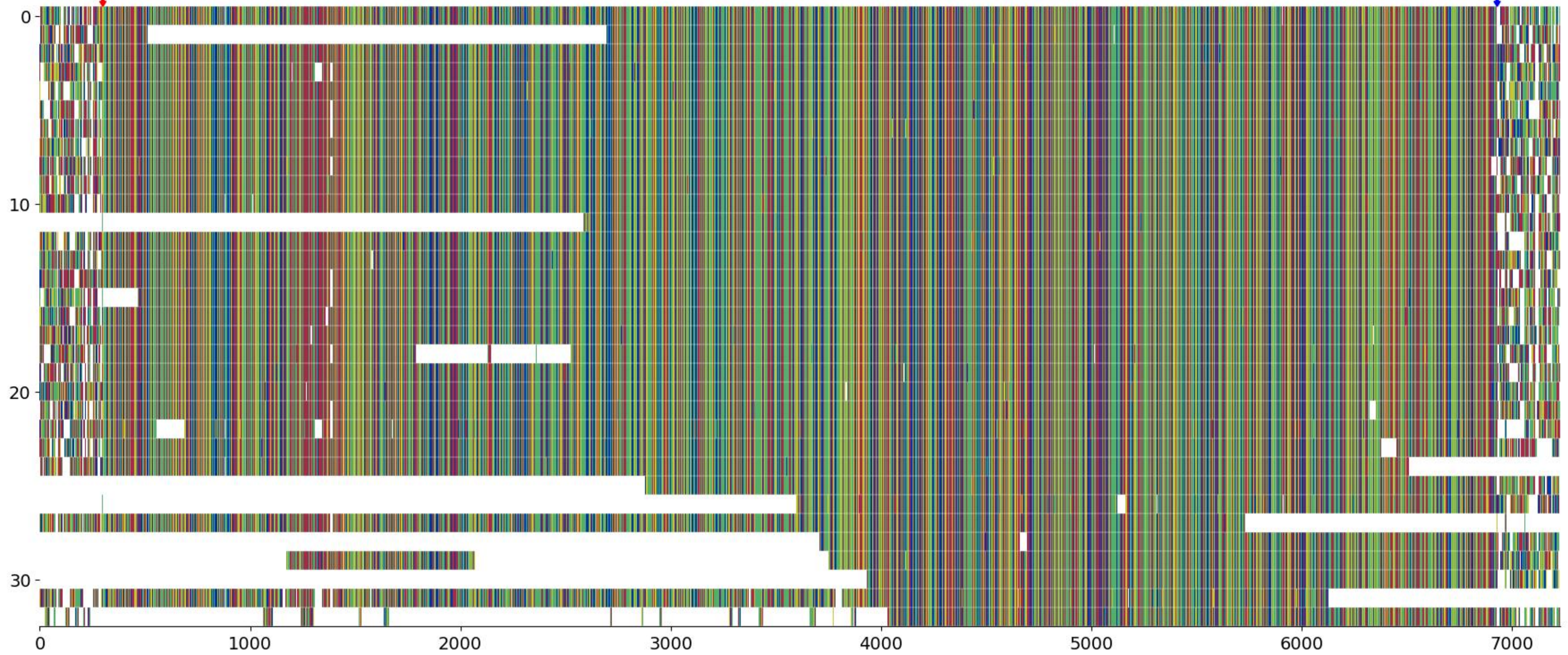




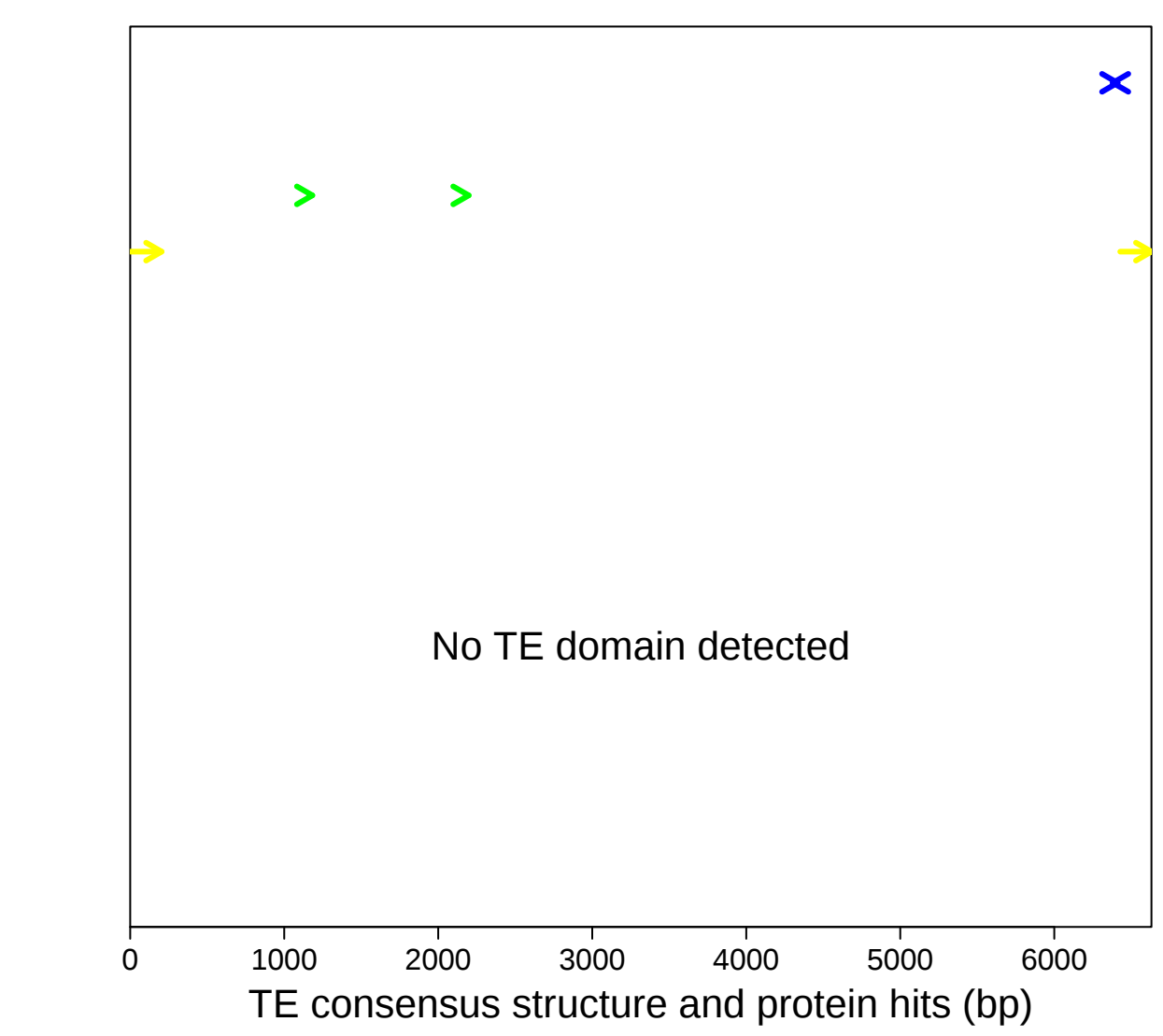
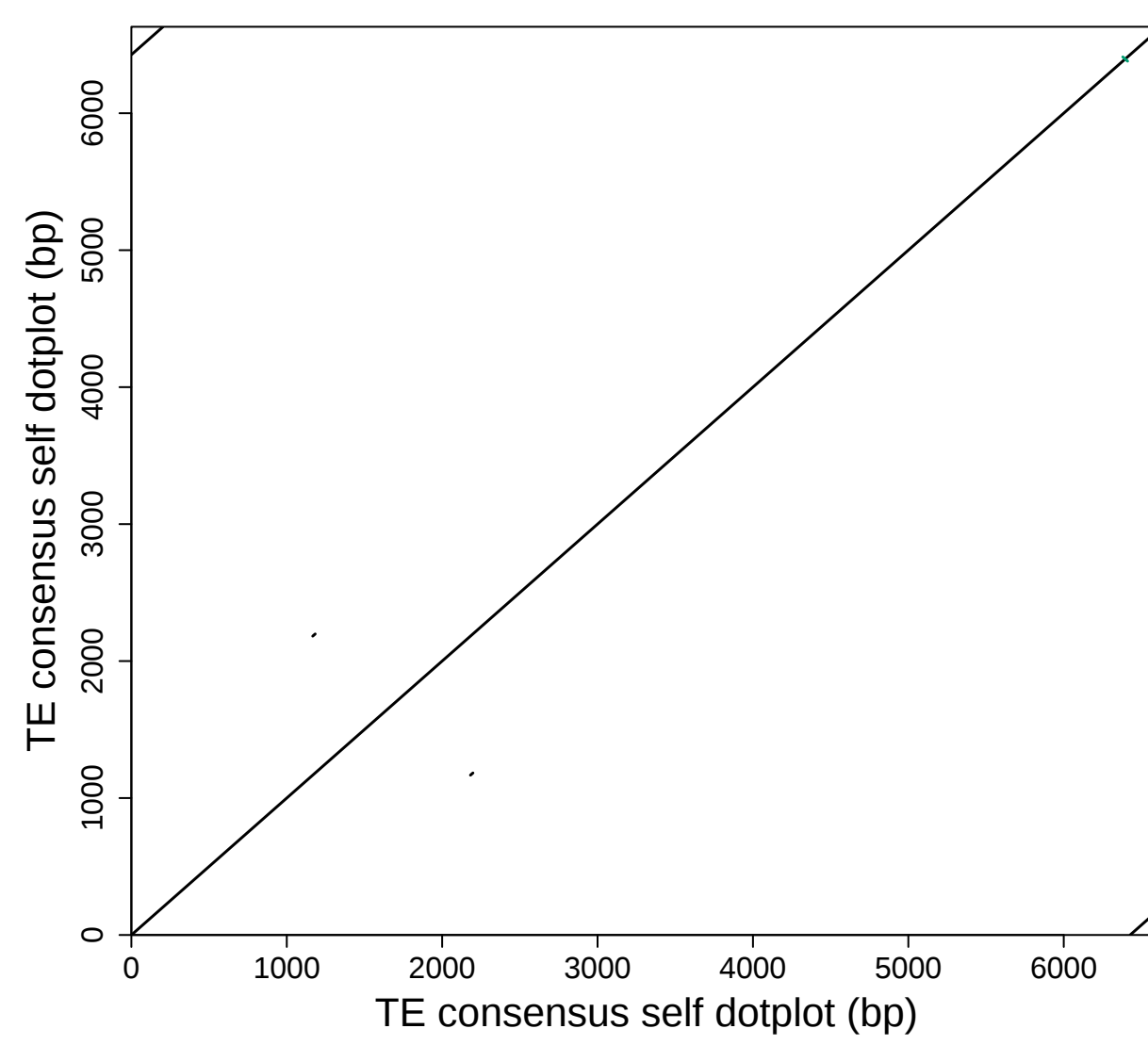
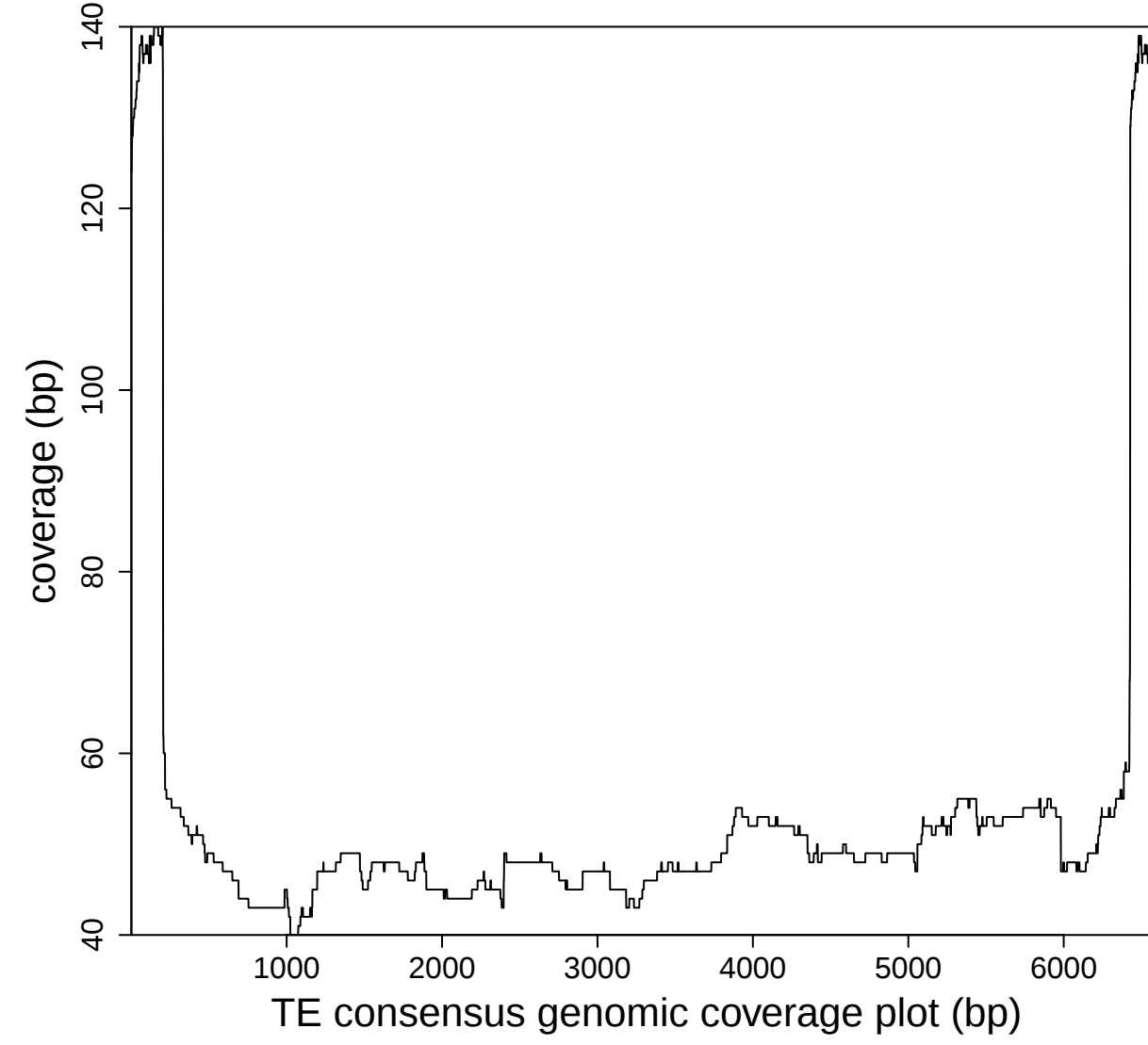
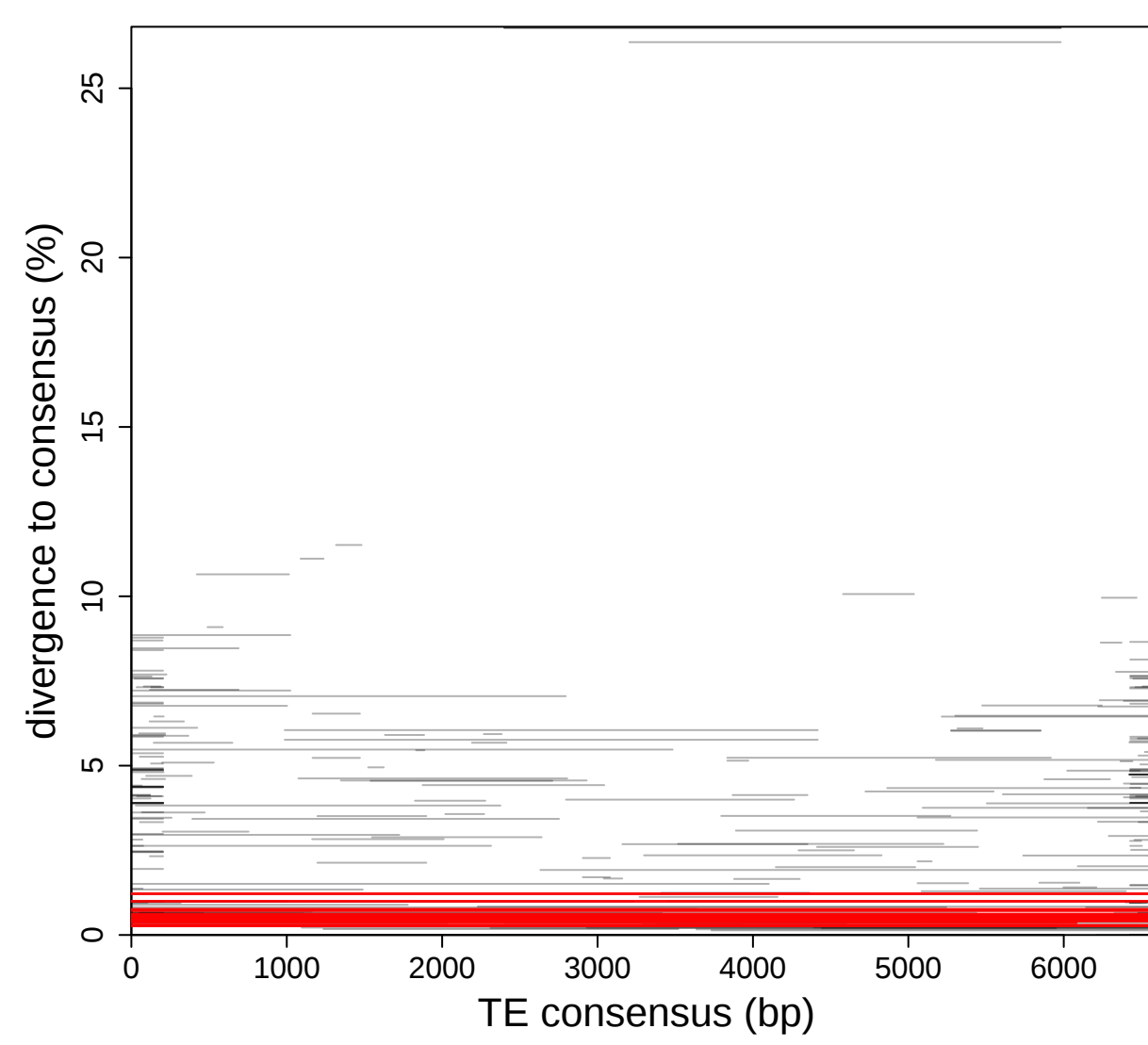
Start crop Point

End crop Point



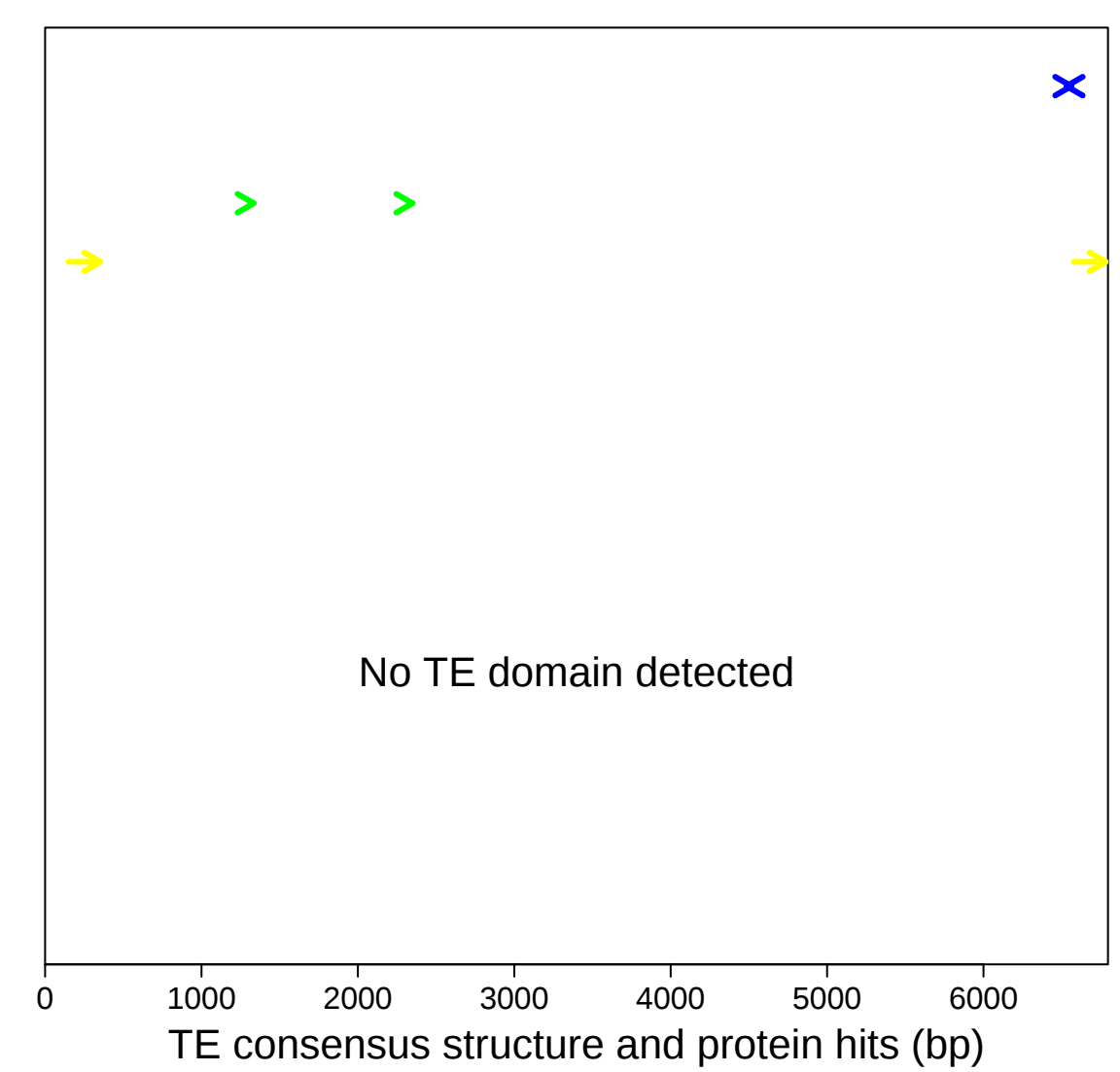
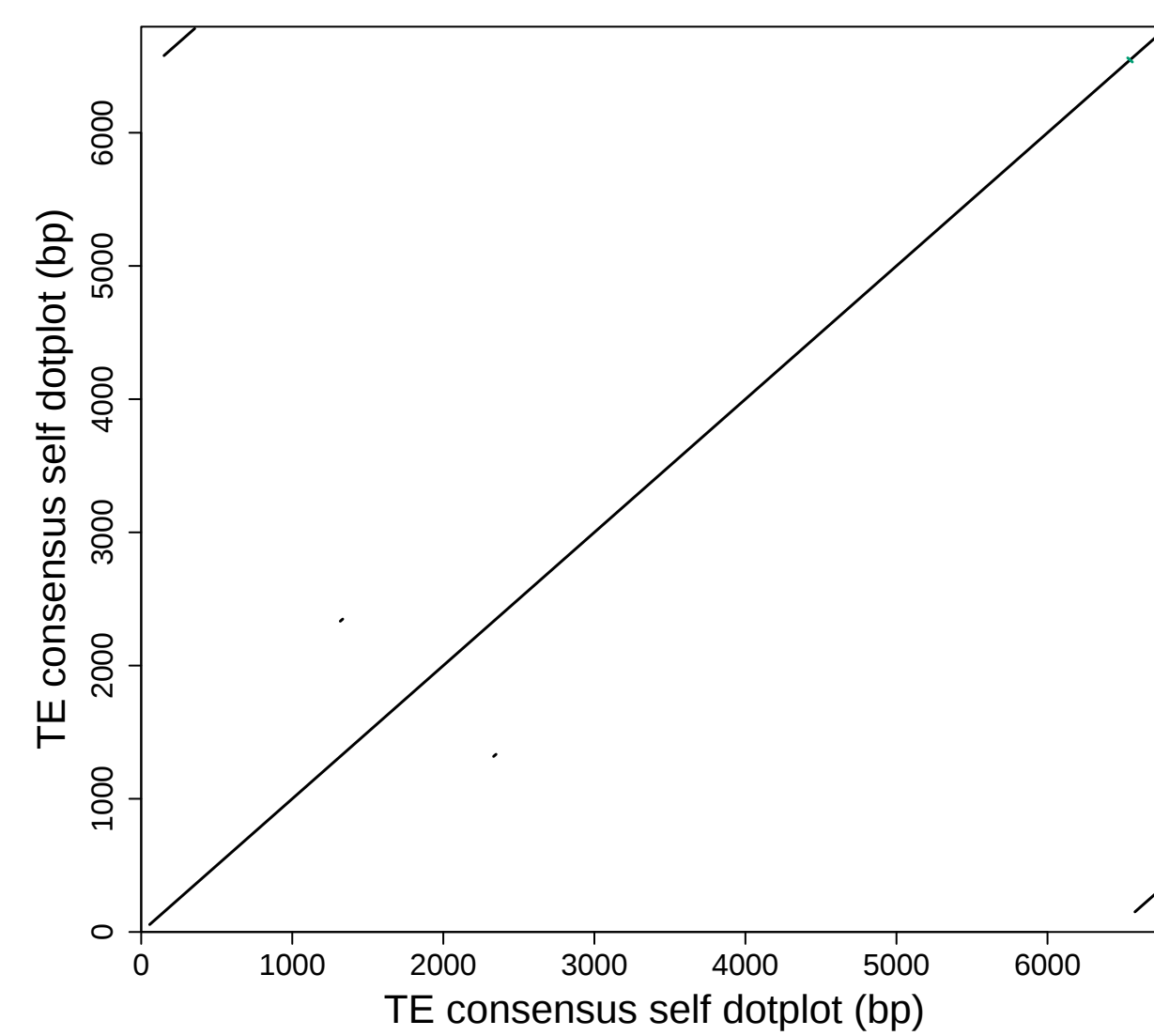
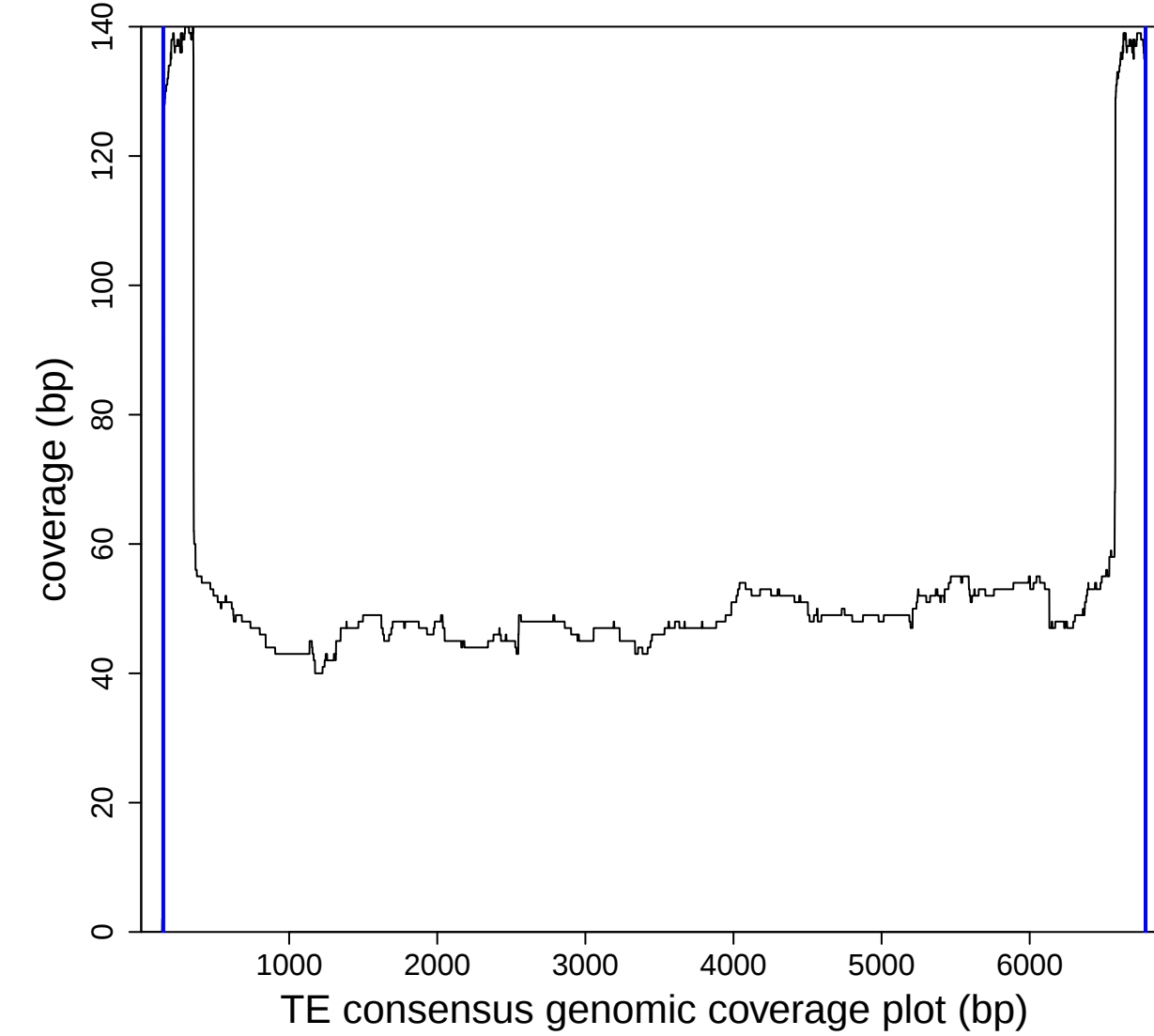
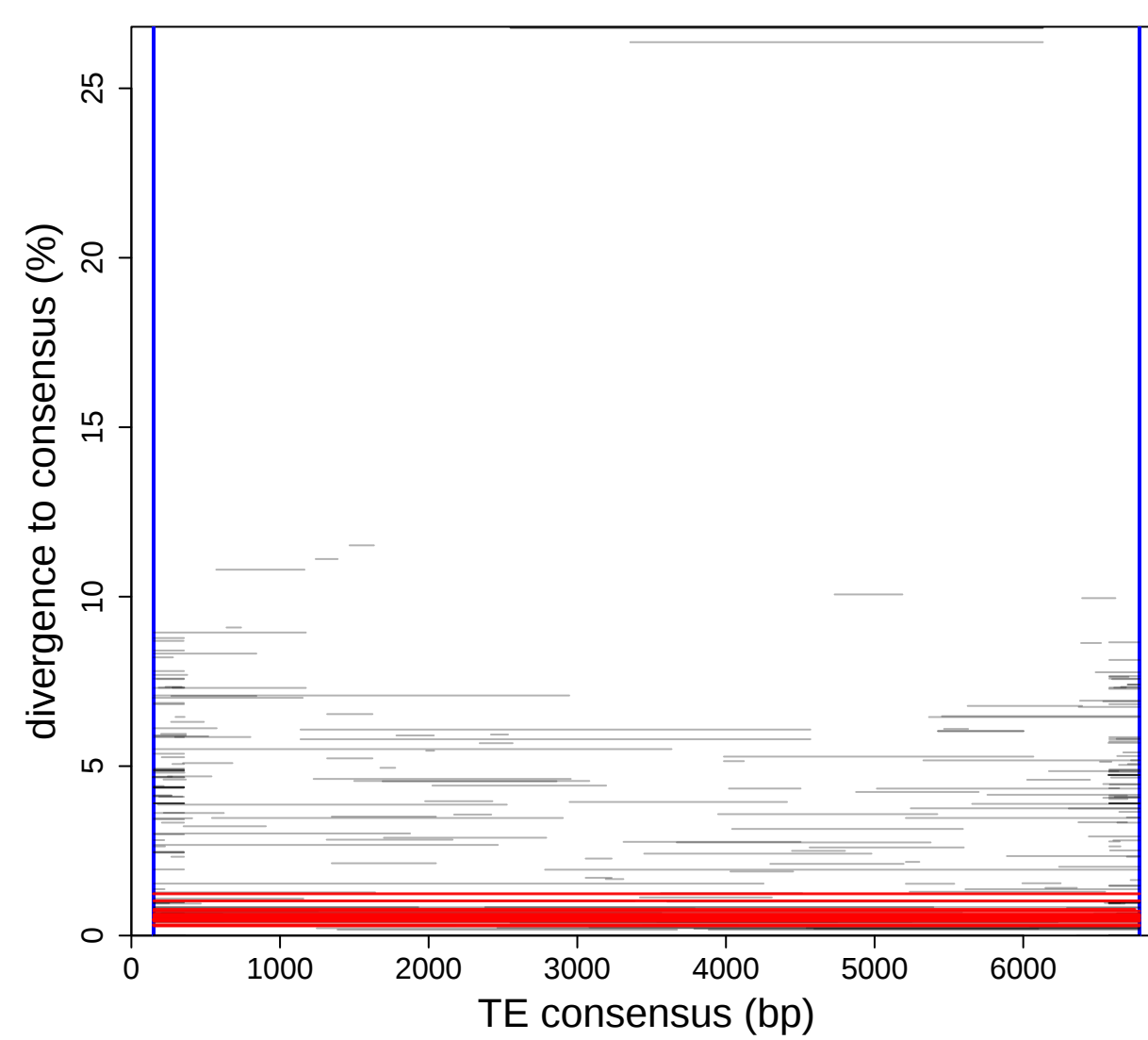
ed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 6631bp; fragments: 357; full length: 19 (≥ 5967.9 bp)

After TETrimmer 6631 bp

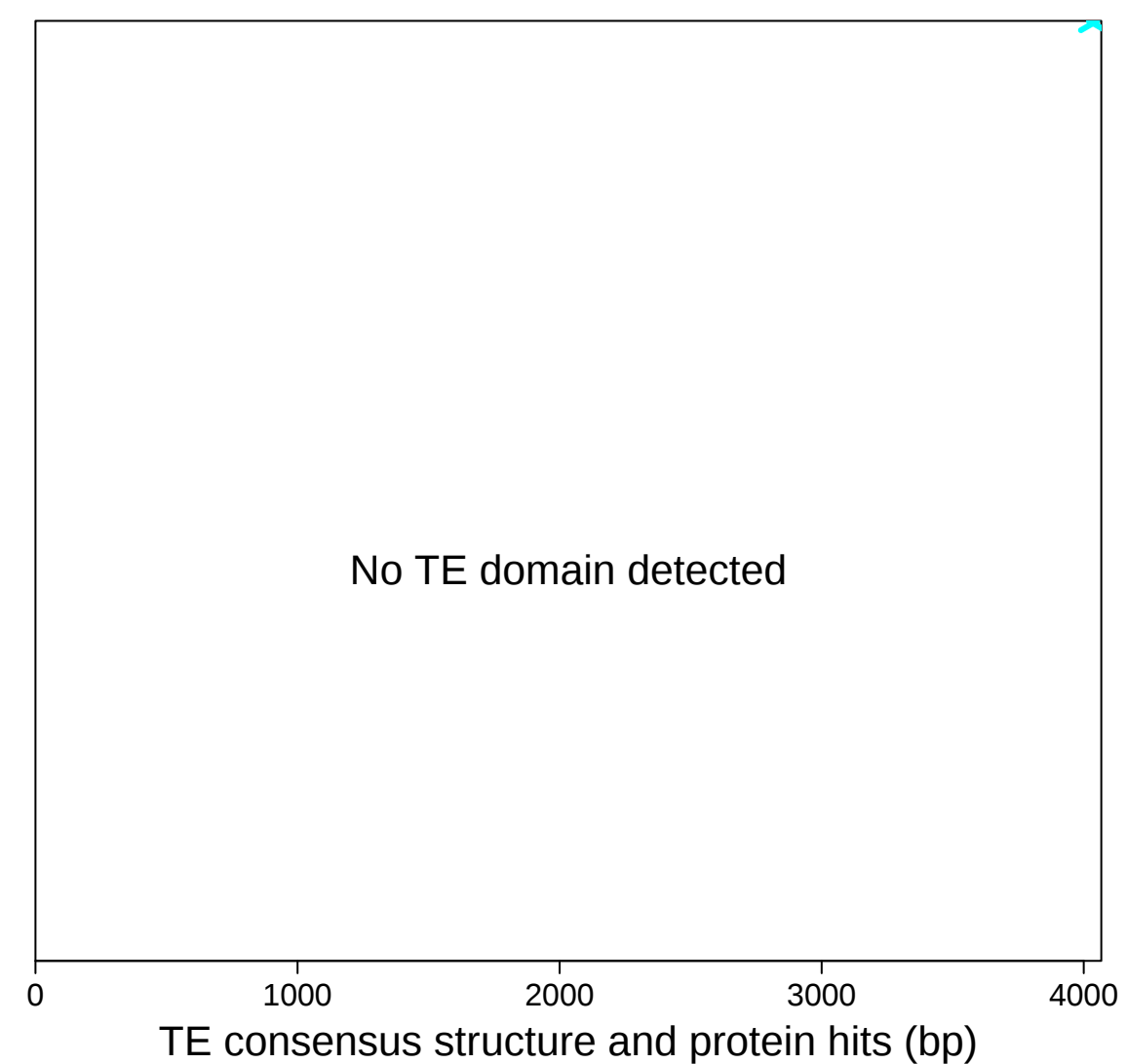
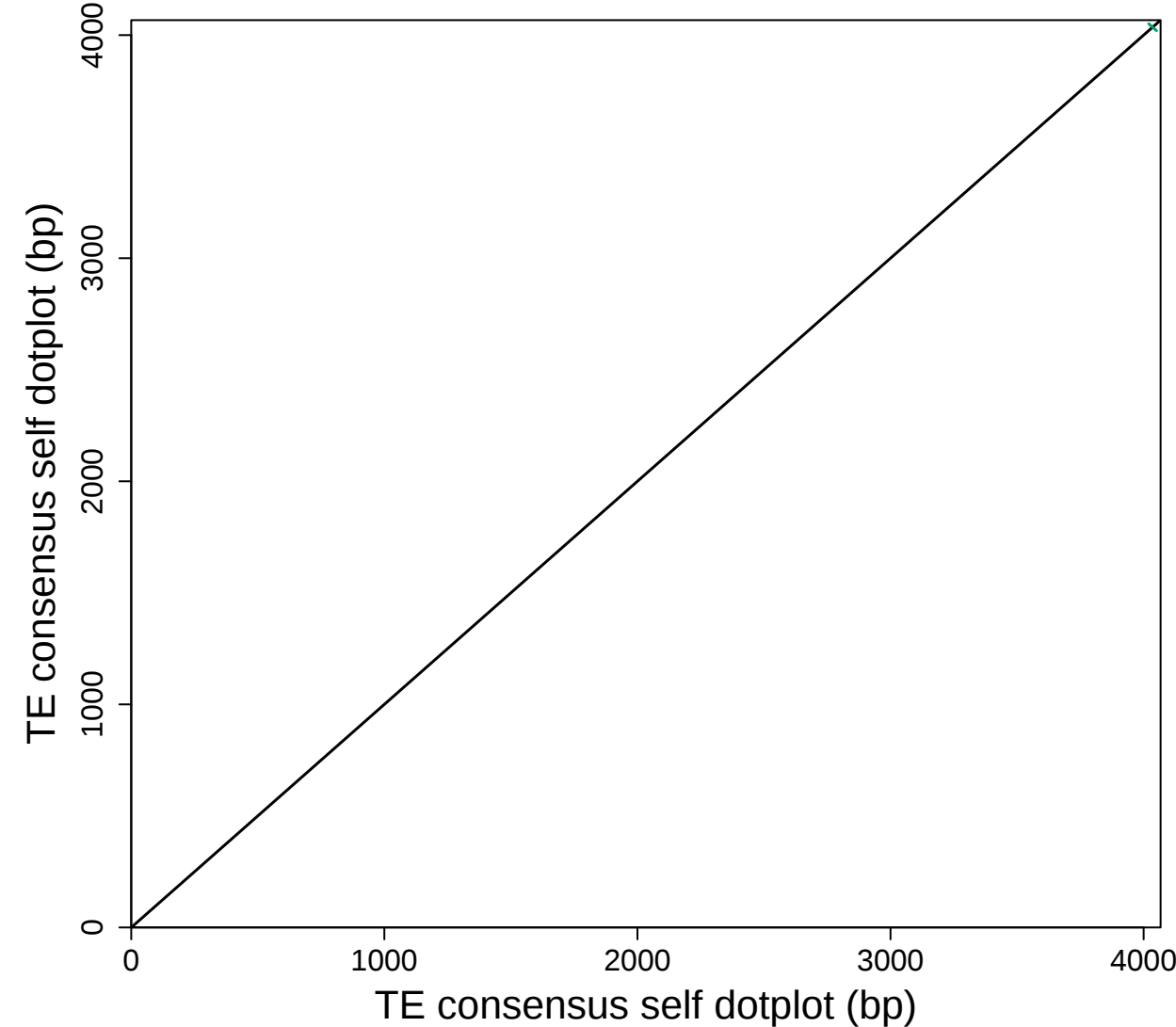
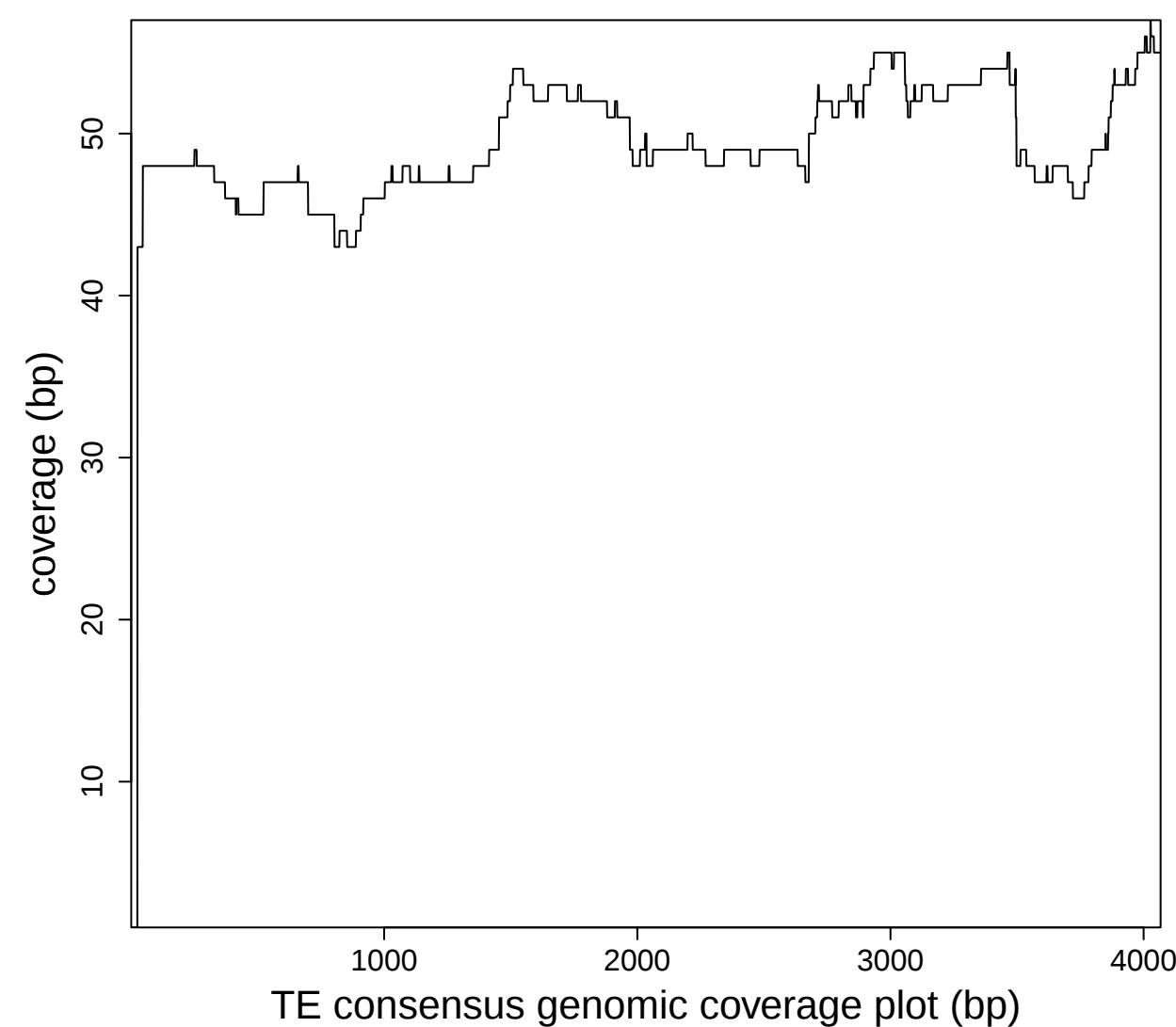
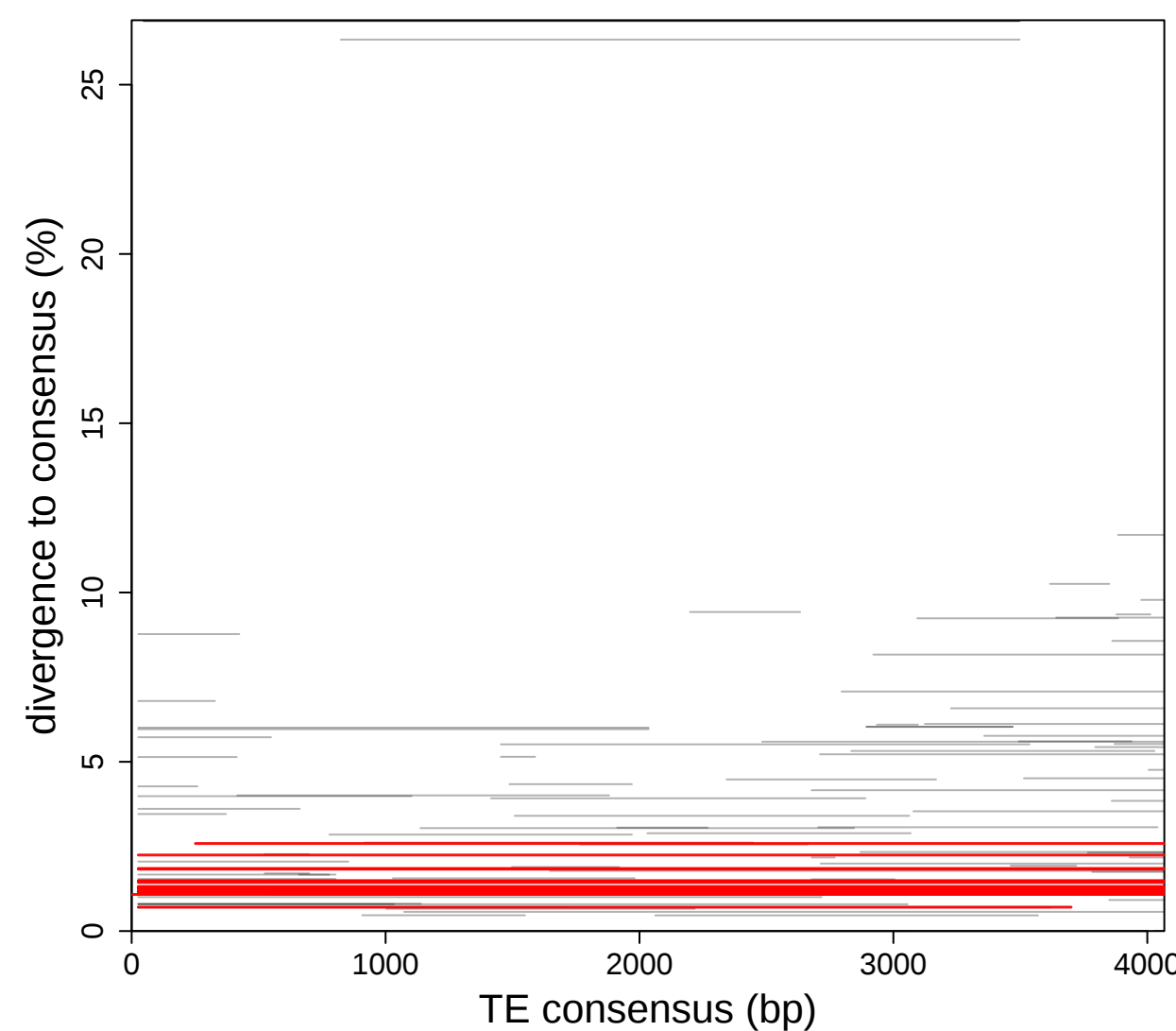


ly_626.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 6932bp; fragments: 357; full length: 17 (≥ 6238.8 bp)

After TEtrimmer Extended plot Blue lines are boundaries

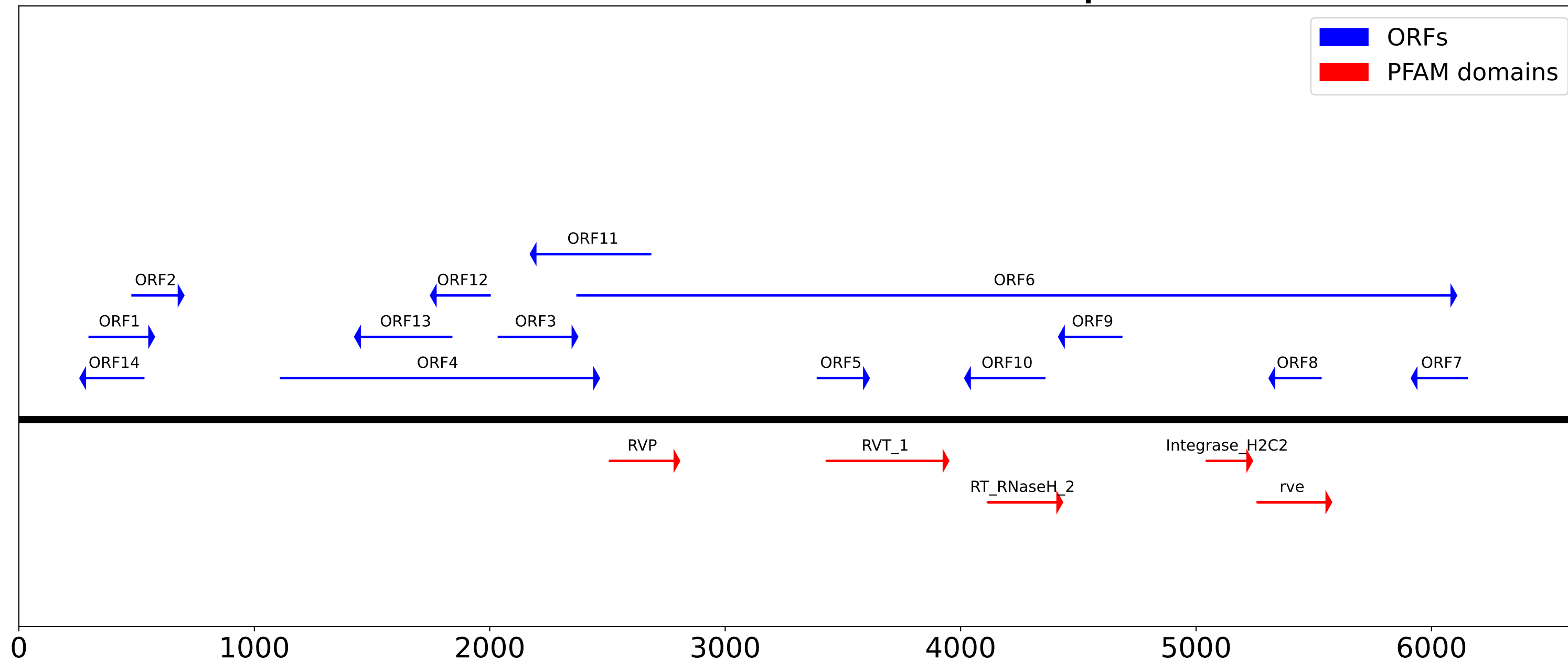


TE: ltr_1_family_626
size: 4067bp; fragments: 119; full length: 23 (≥ 3660.3 bp)

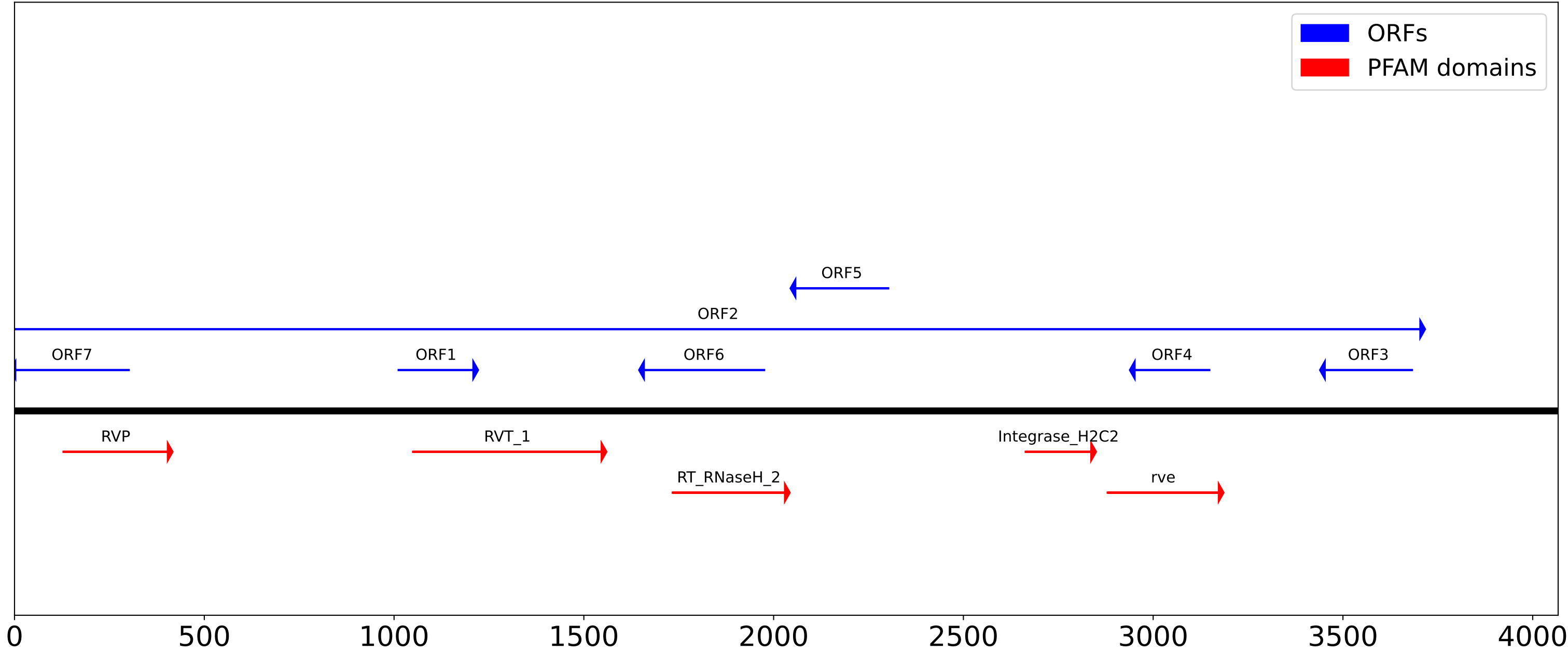


Before TEtrimmer 4067 bp

After TEtrimmer ORF and PFAM domain plot



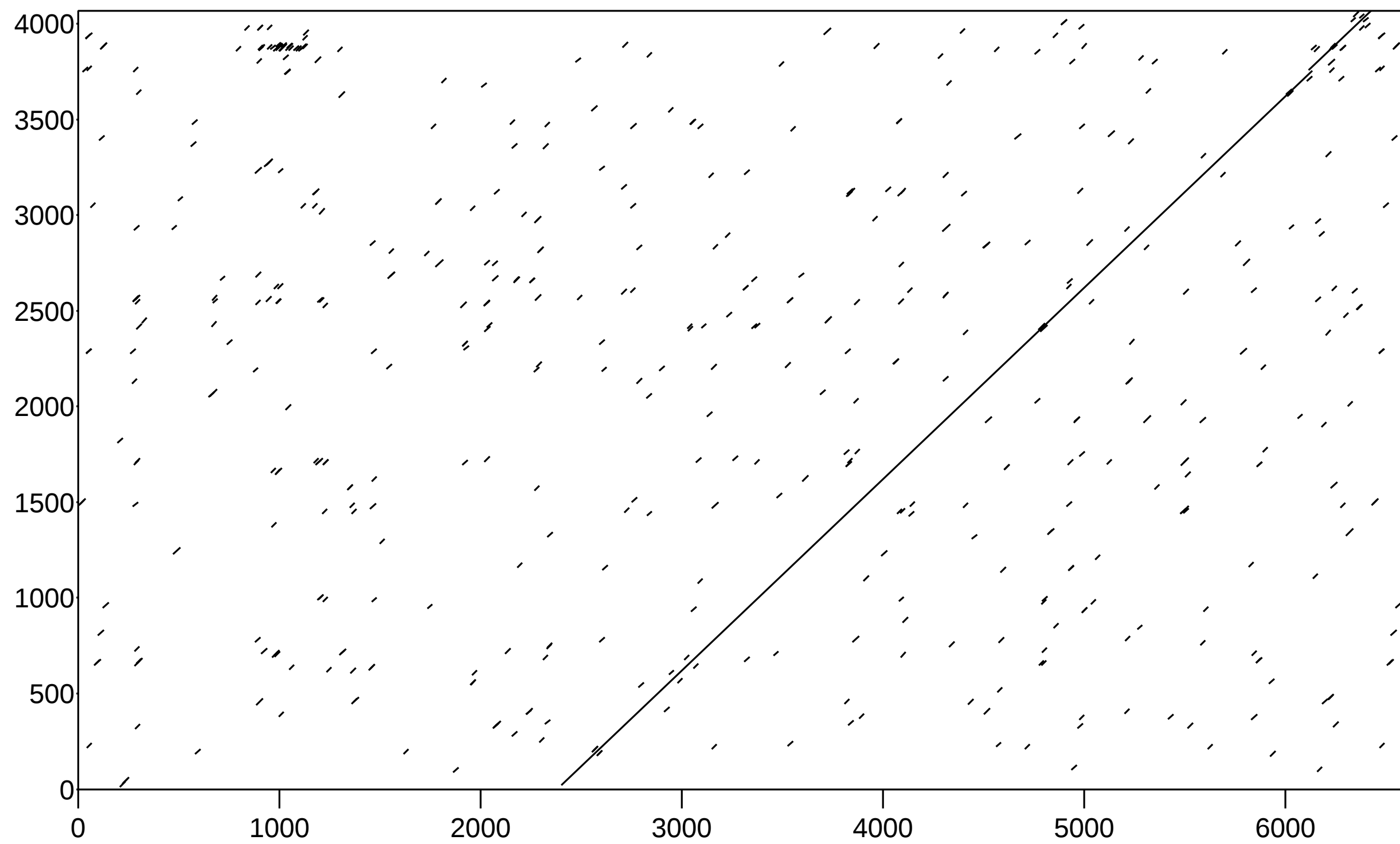
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)