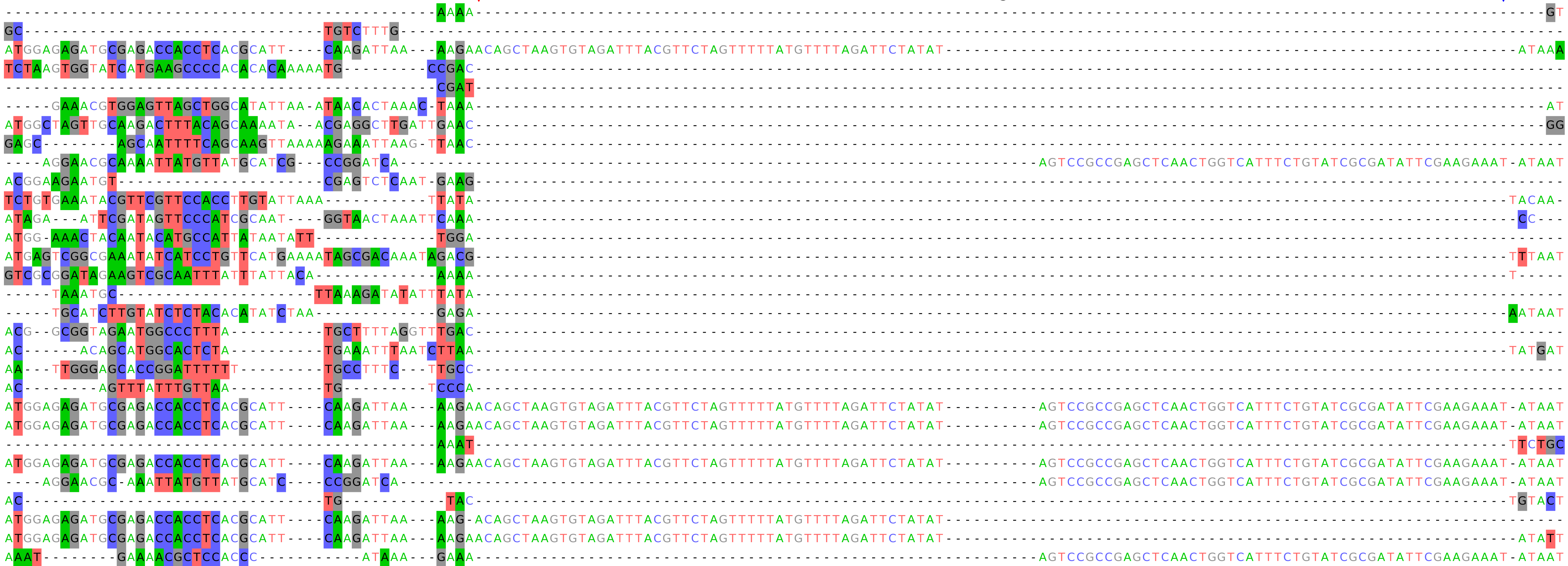
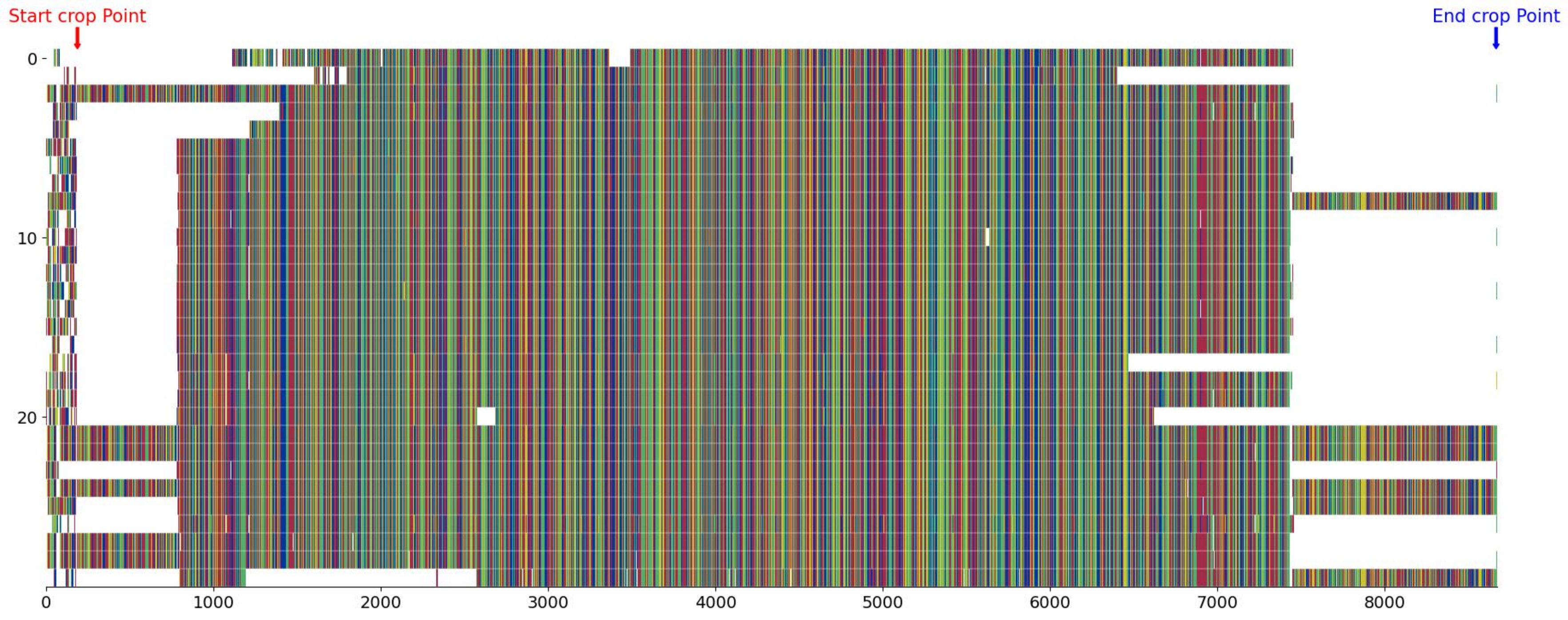


Start crop Point

End crop Point

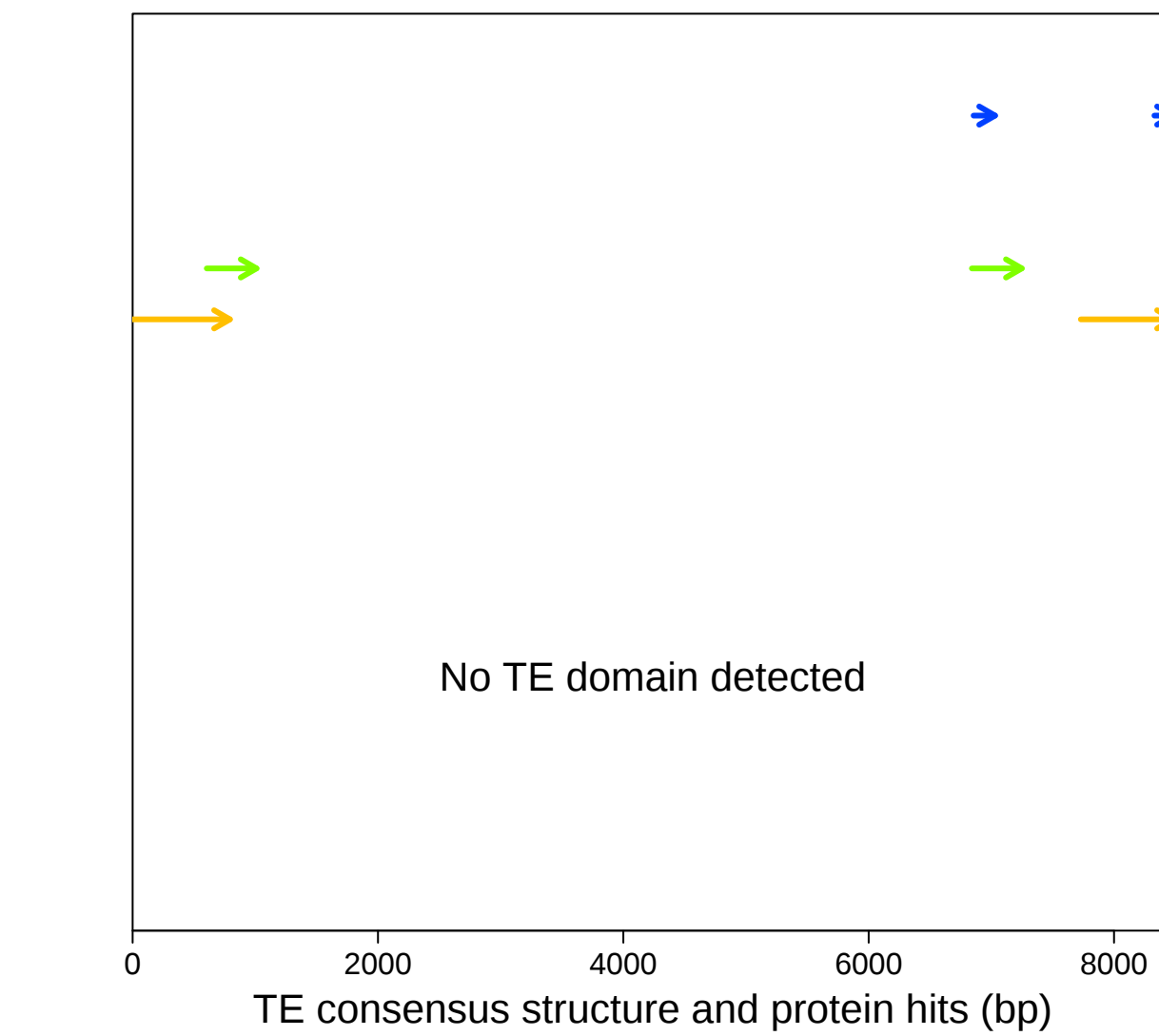
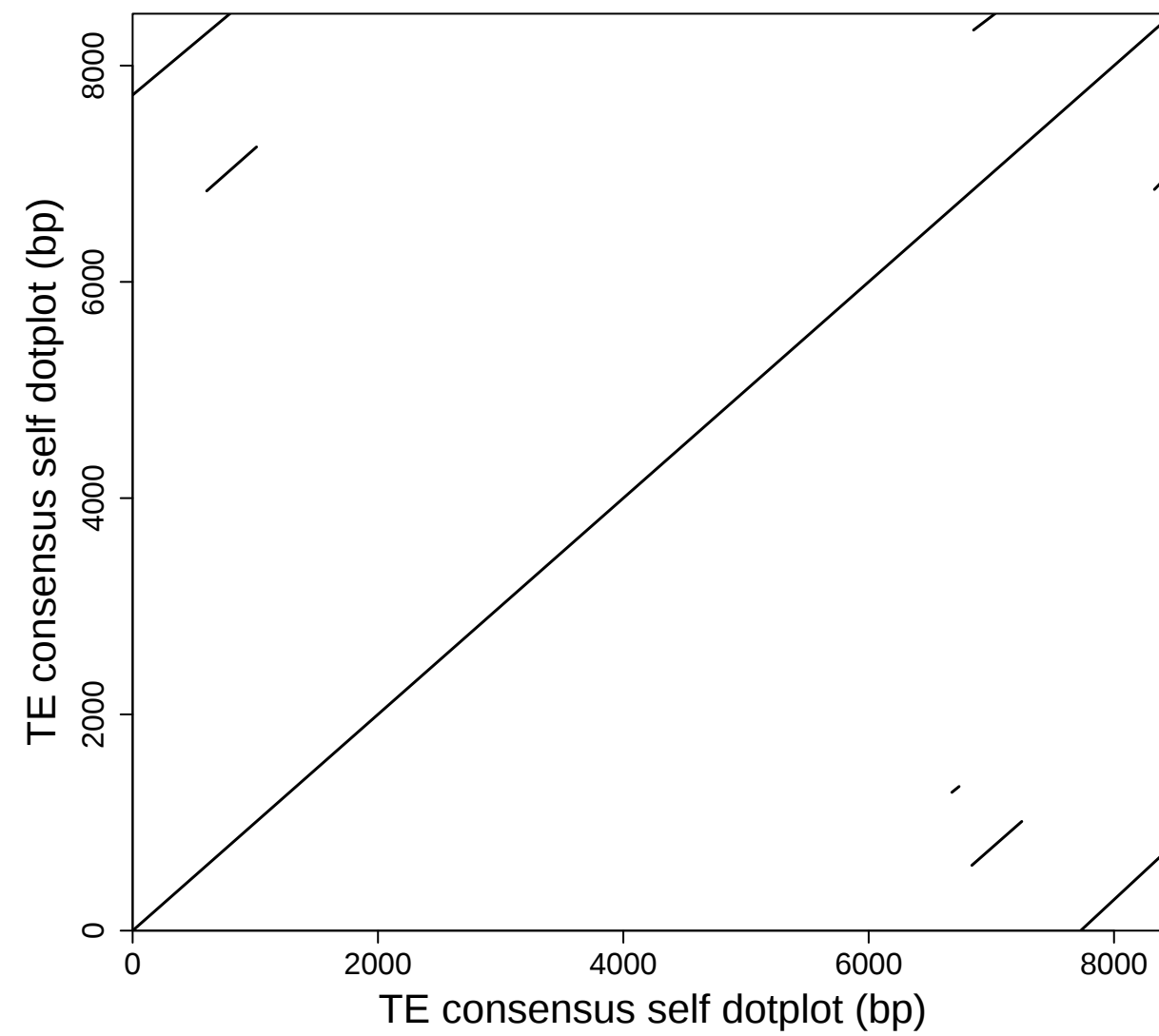
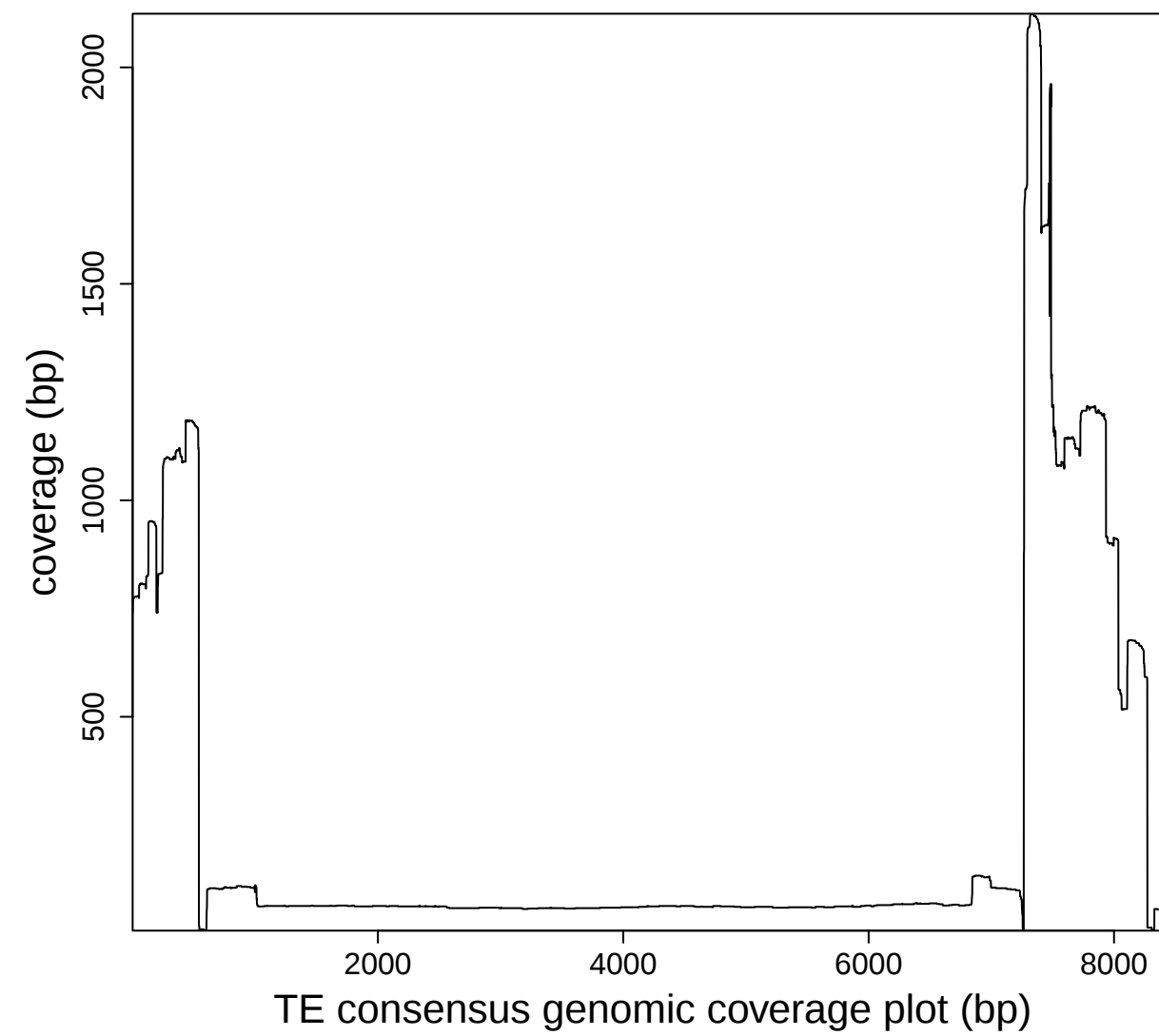
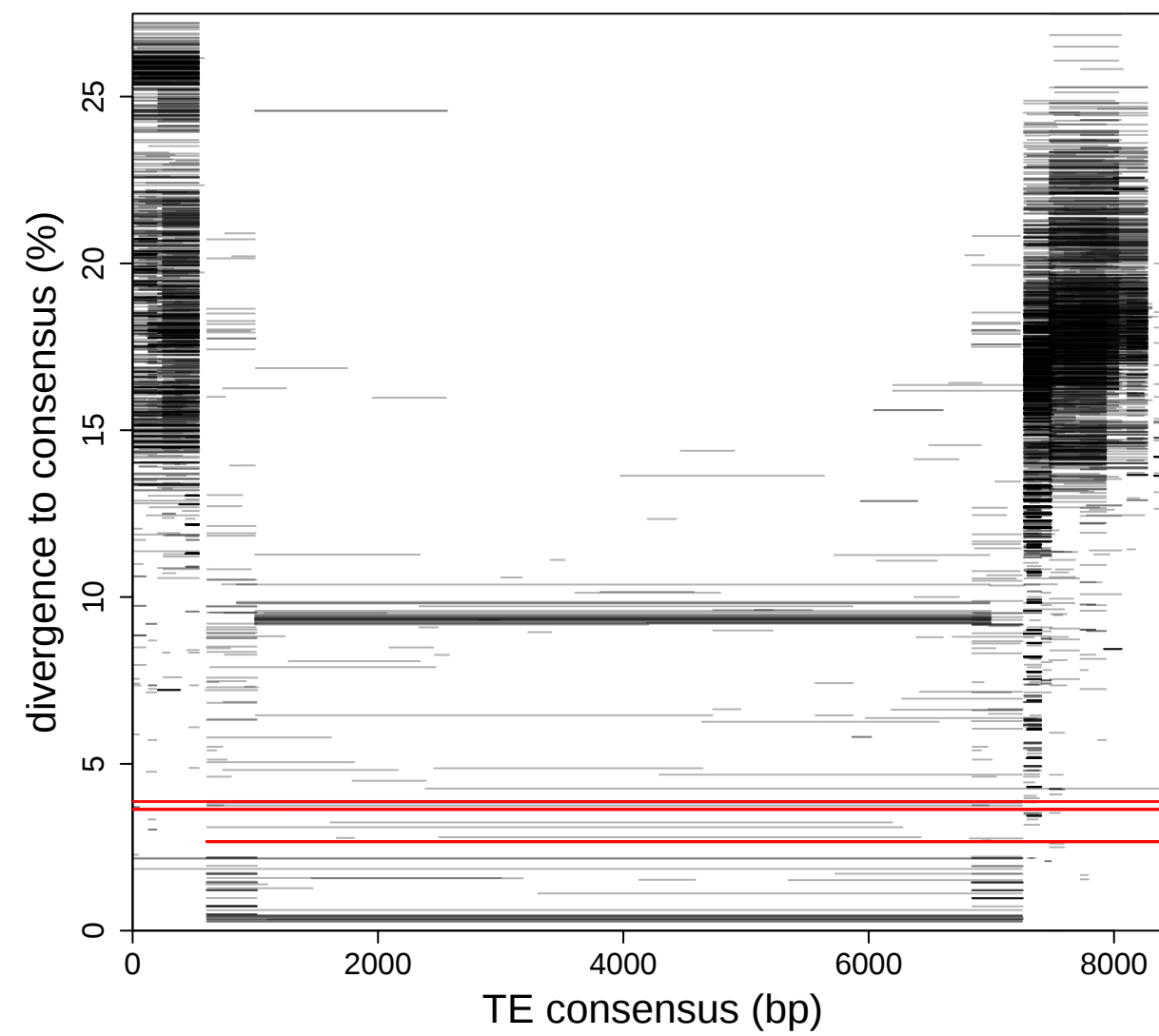
MSA length = 8480





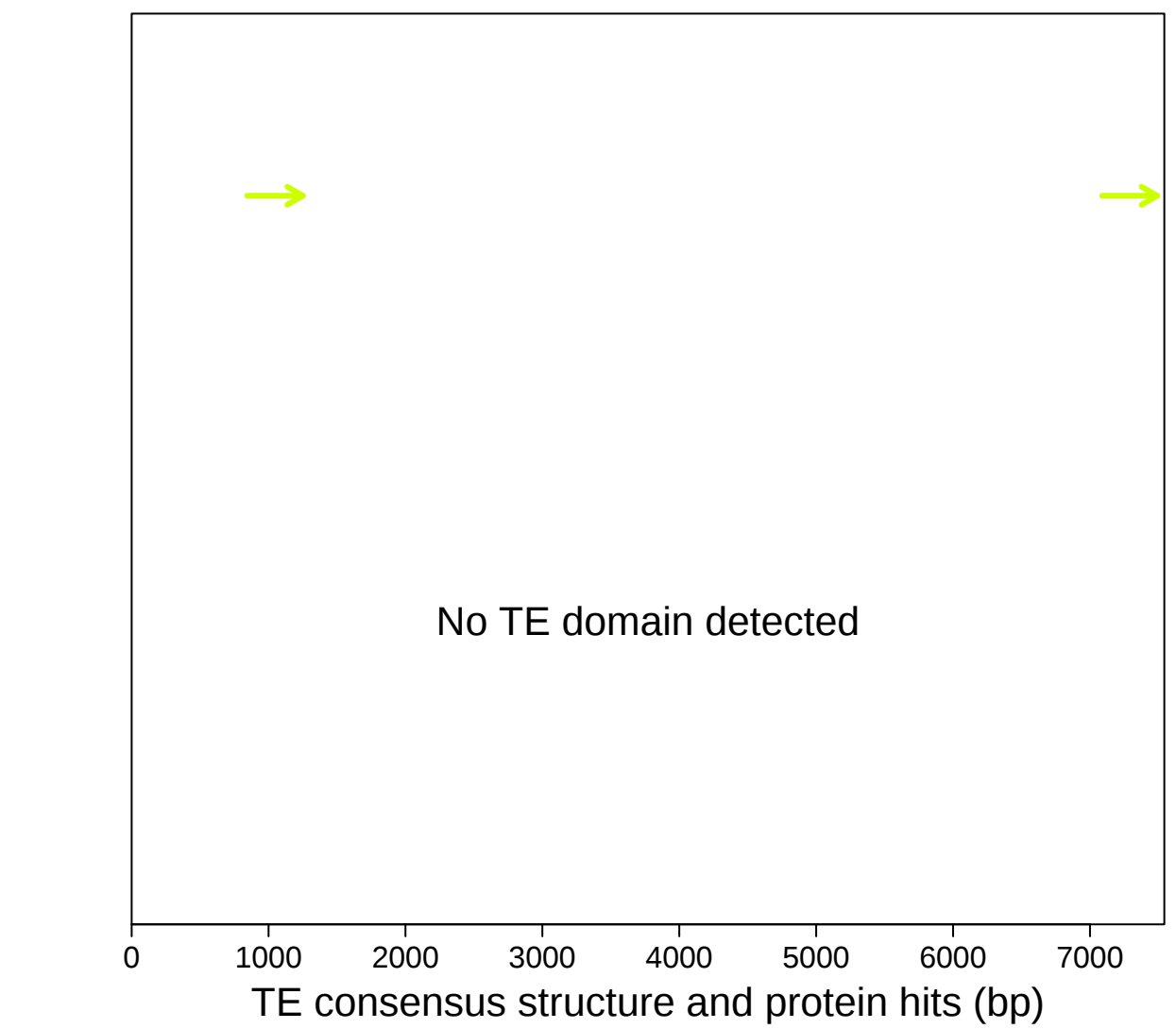
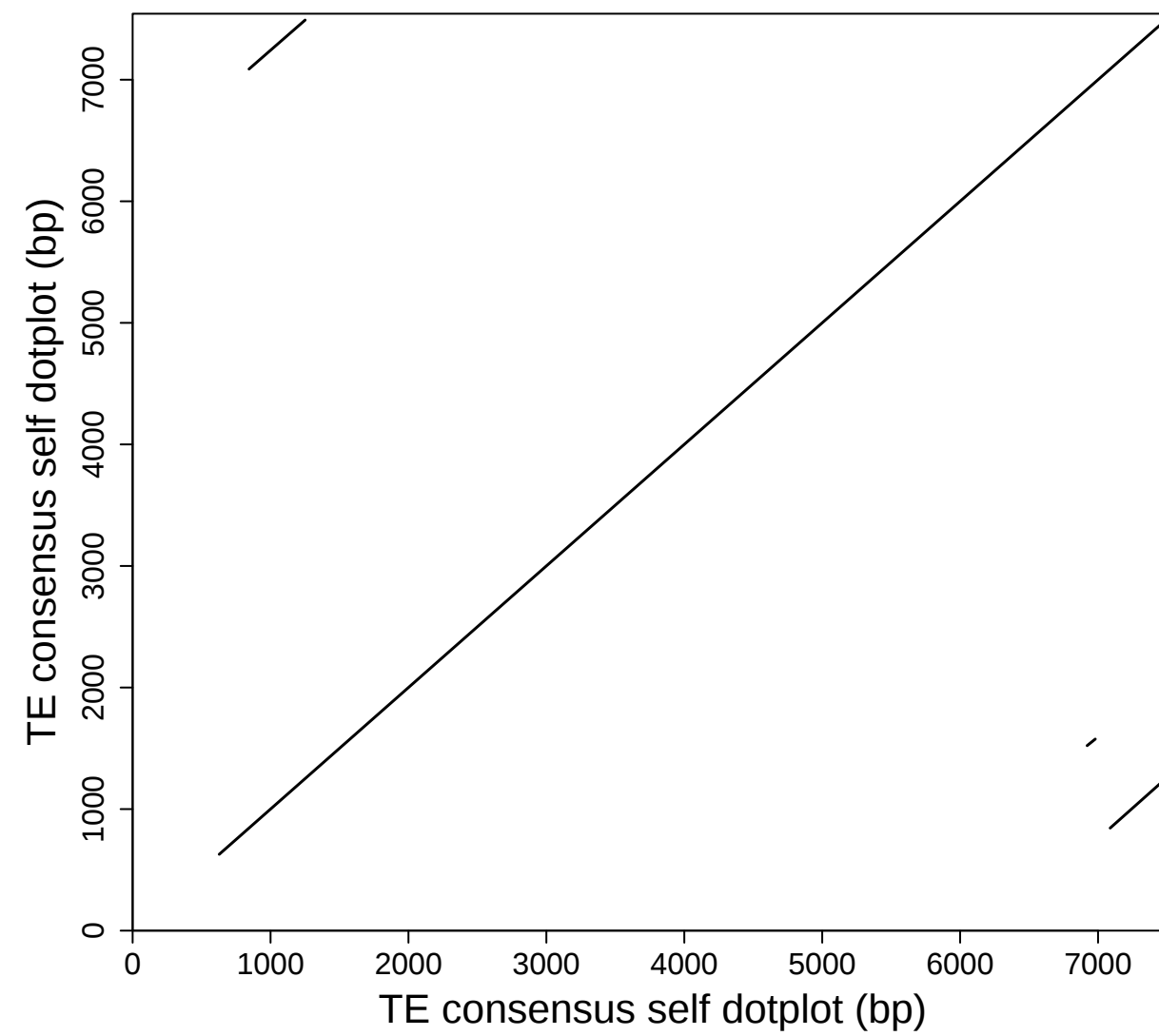
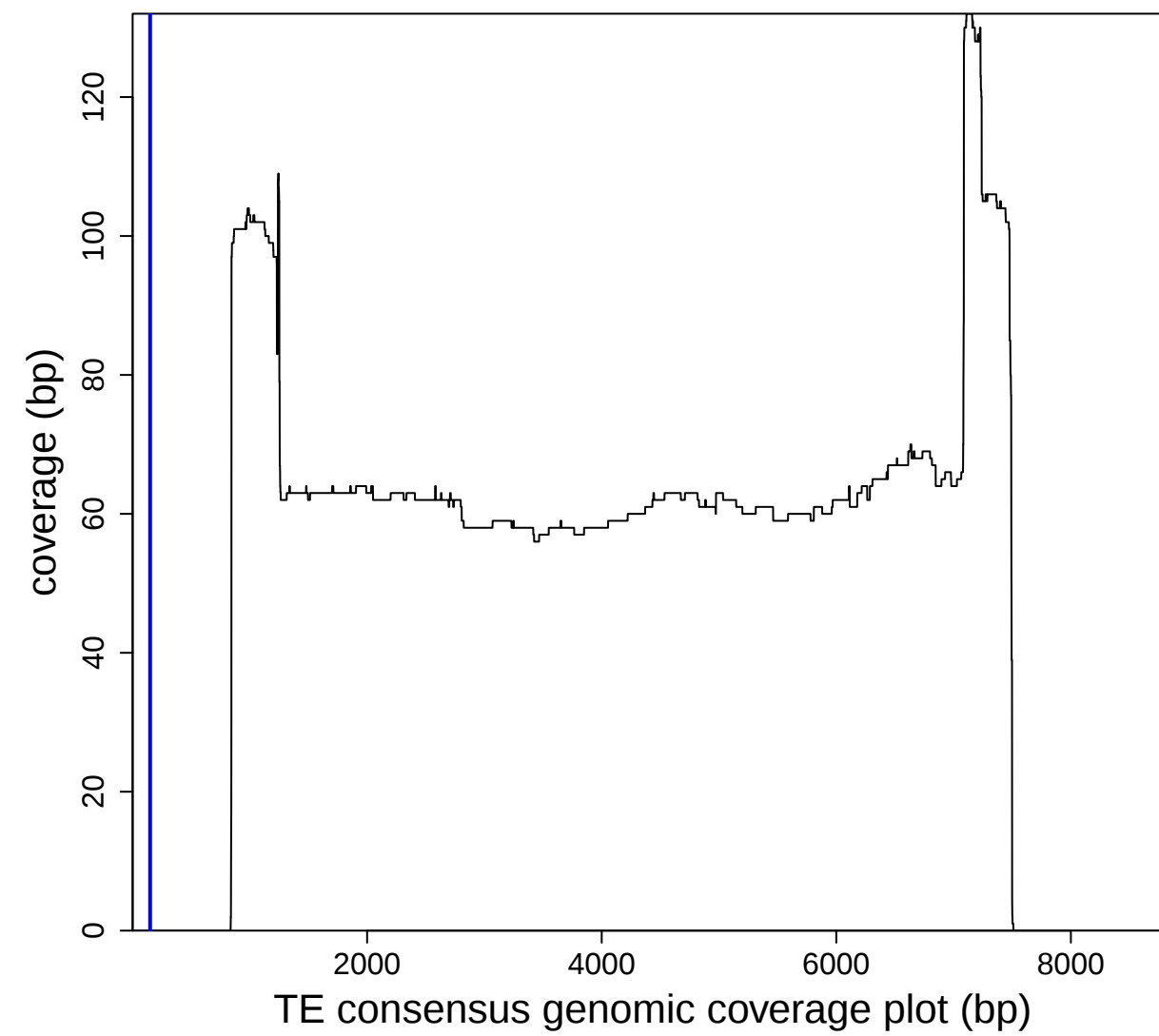
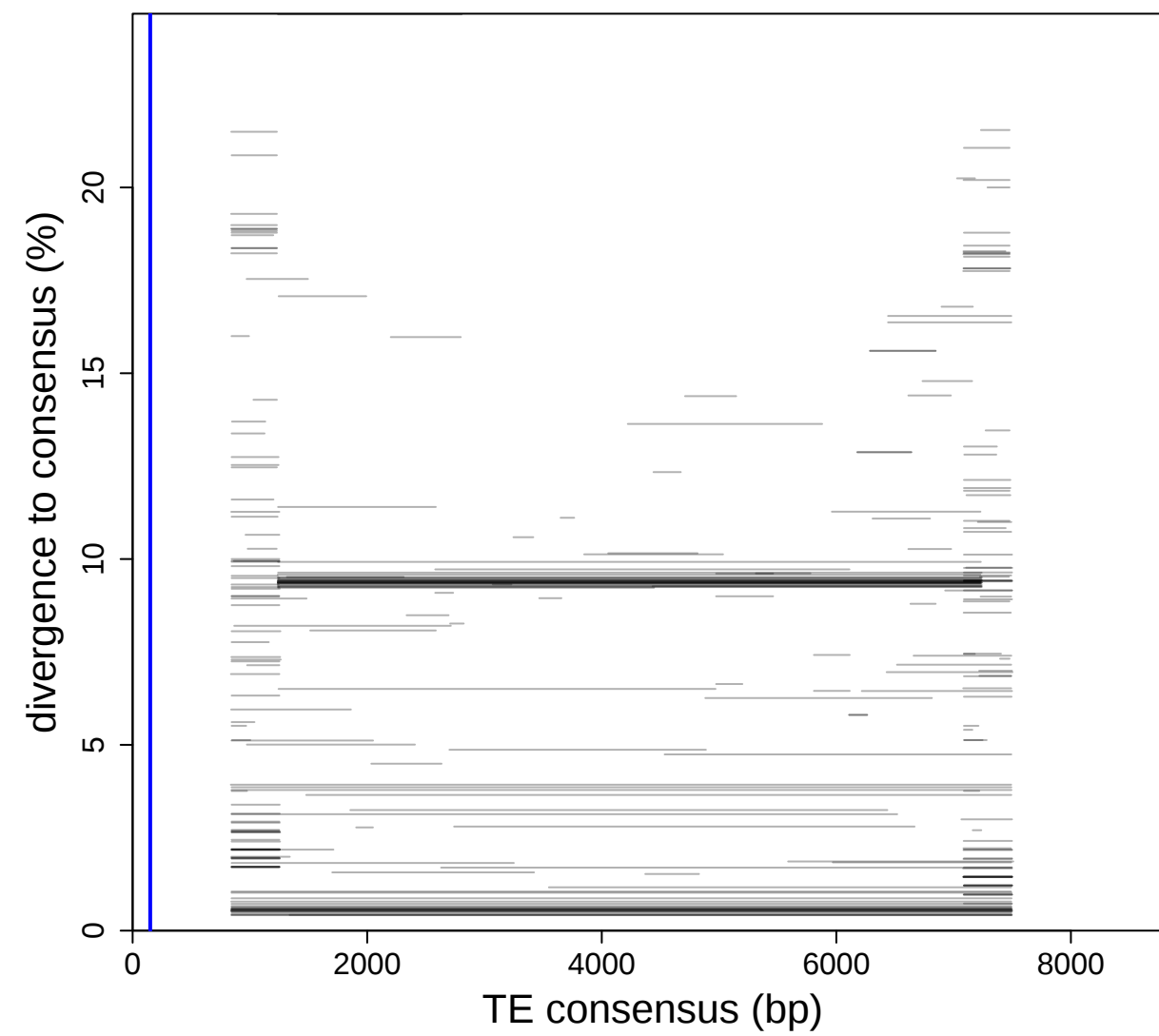
ed\_uf.bed\_g\_1.bed\_fm\_1.bed\_0\_0\_bcln.fa\_aln.fa\_cl.fa\_gs.fa\_ce.  
size: 8480bp; fragments: 5203; full length: 5 (>=7632bp)

# After TEtrimmer 8480 bp



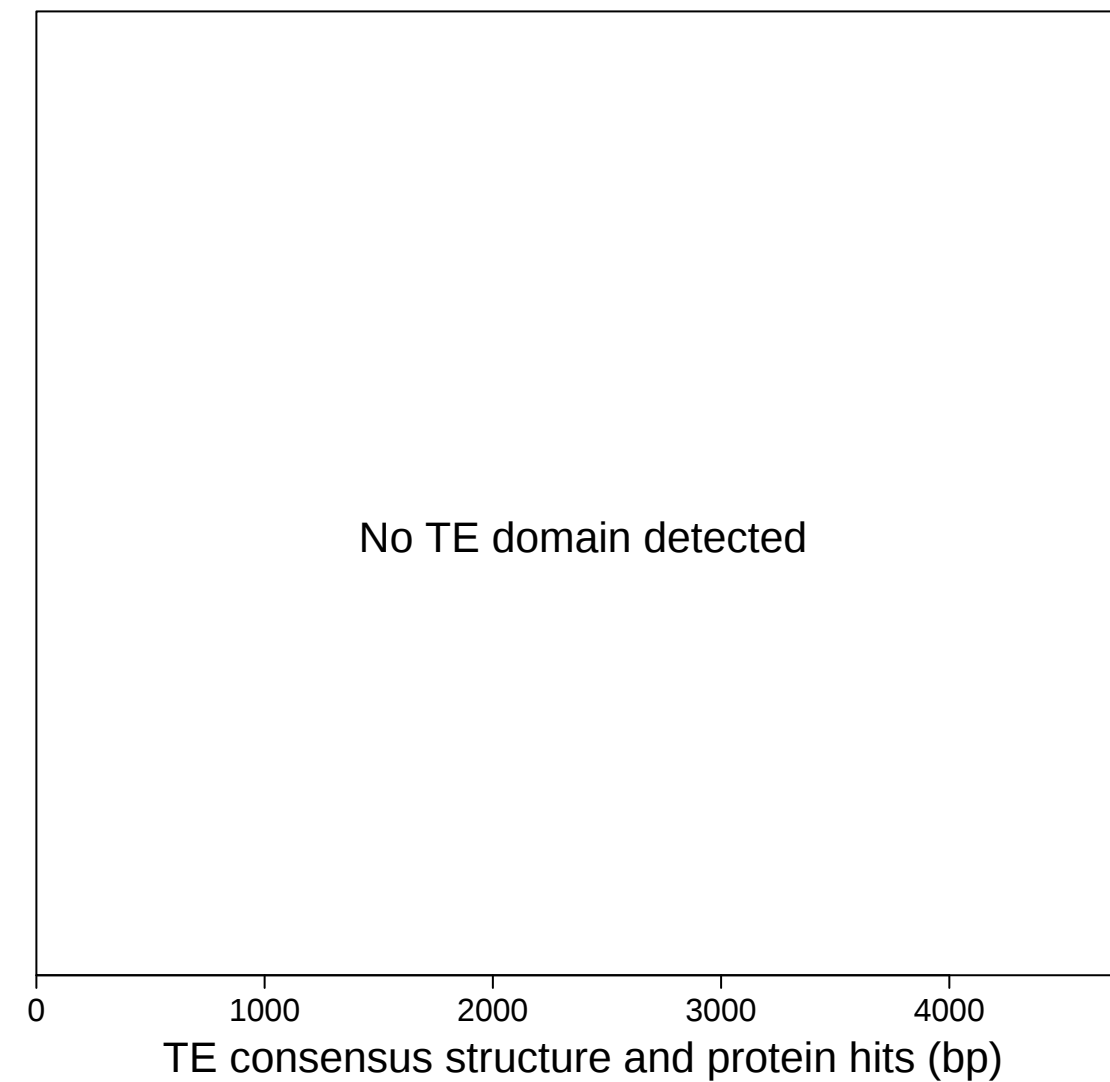
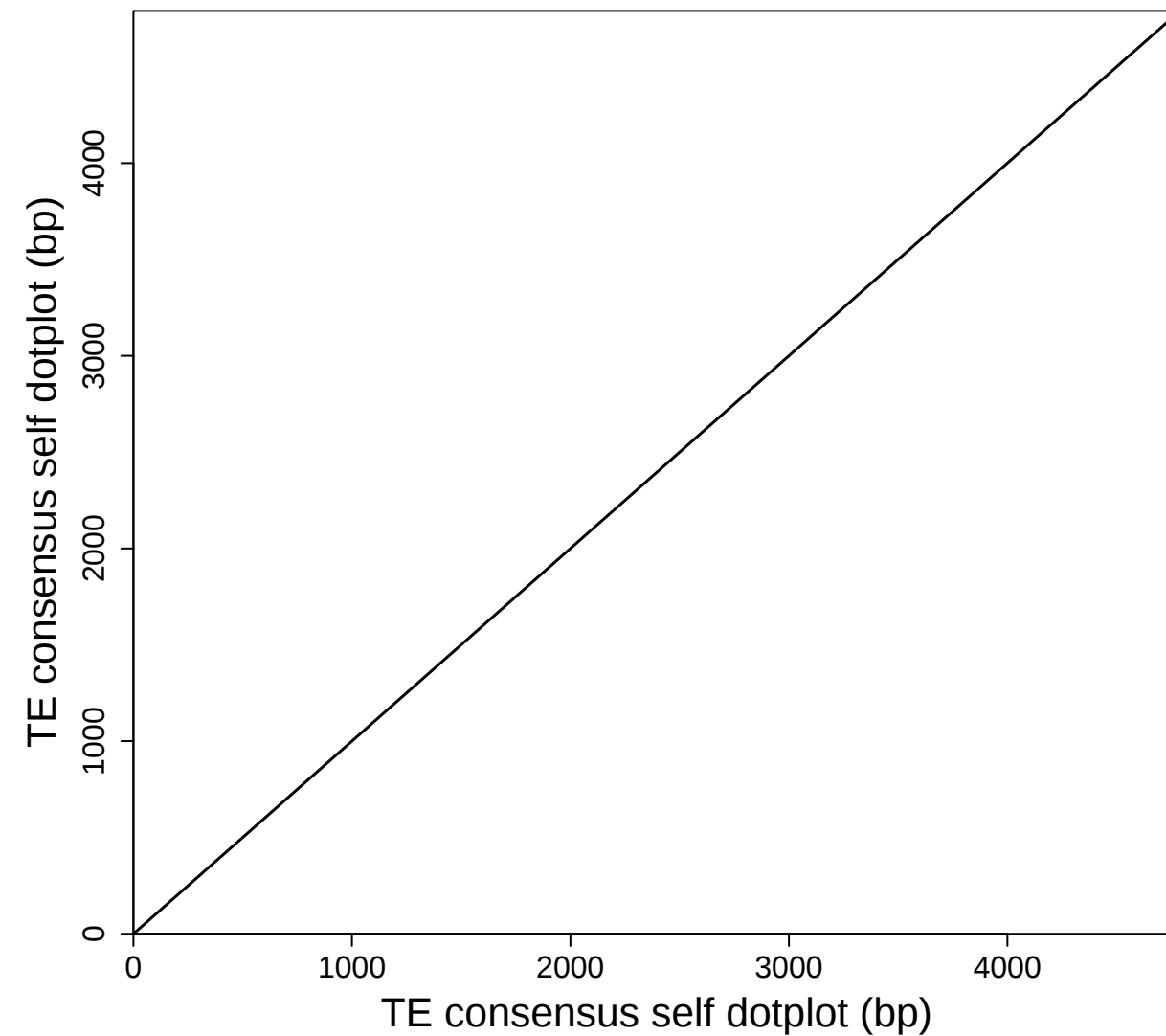
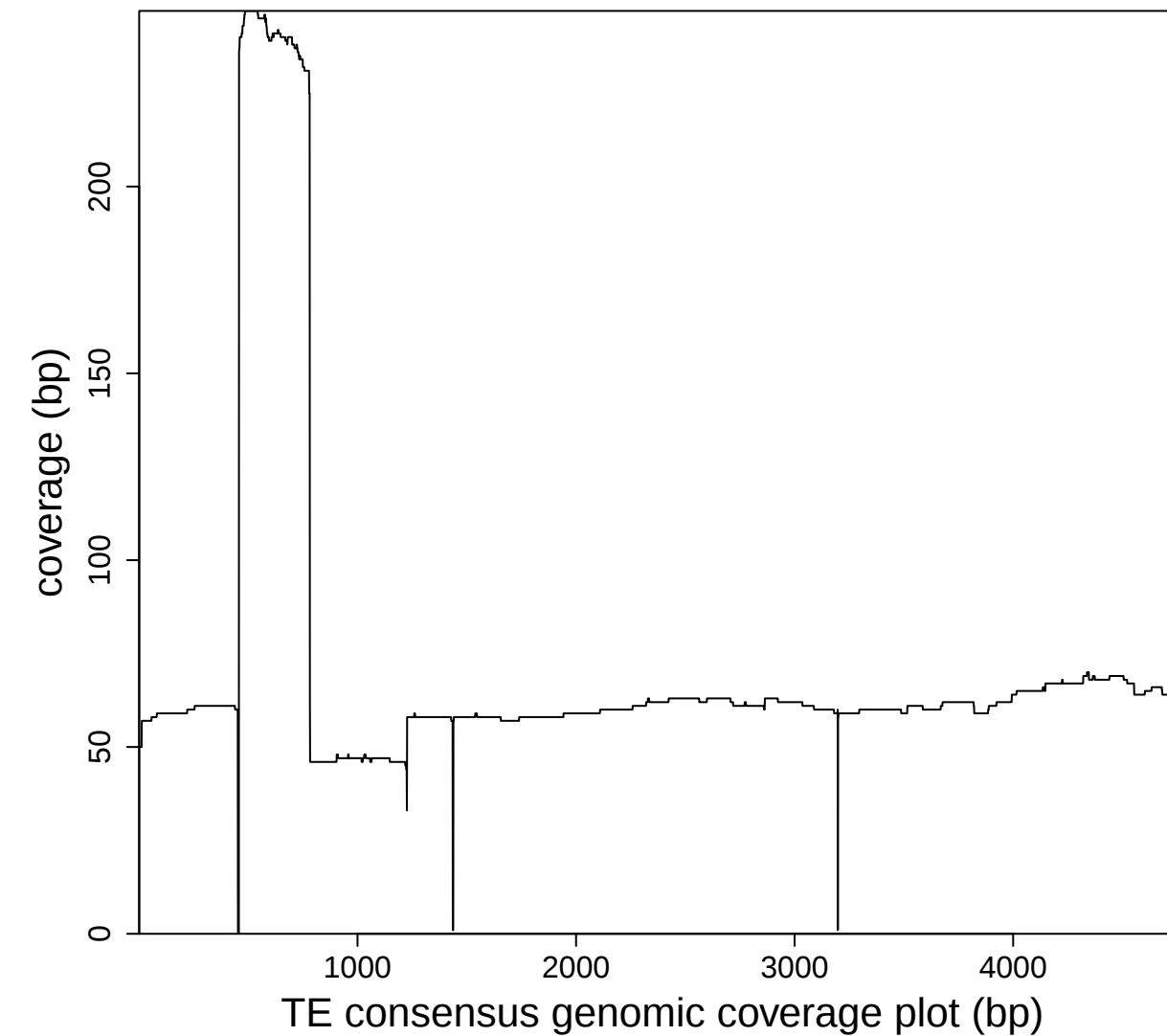
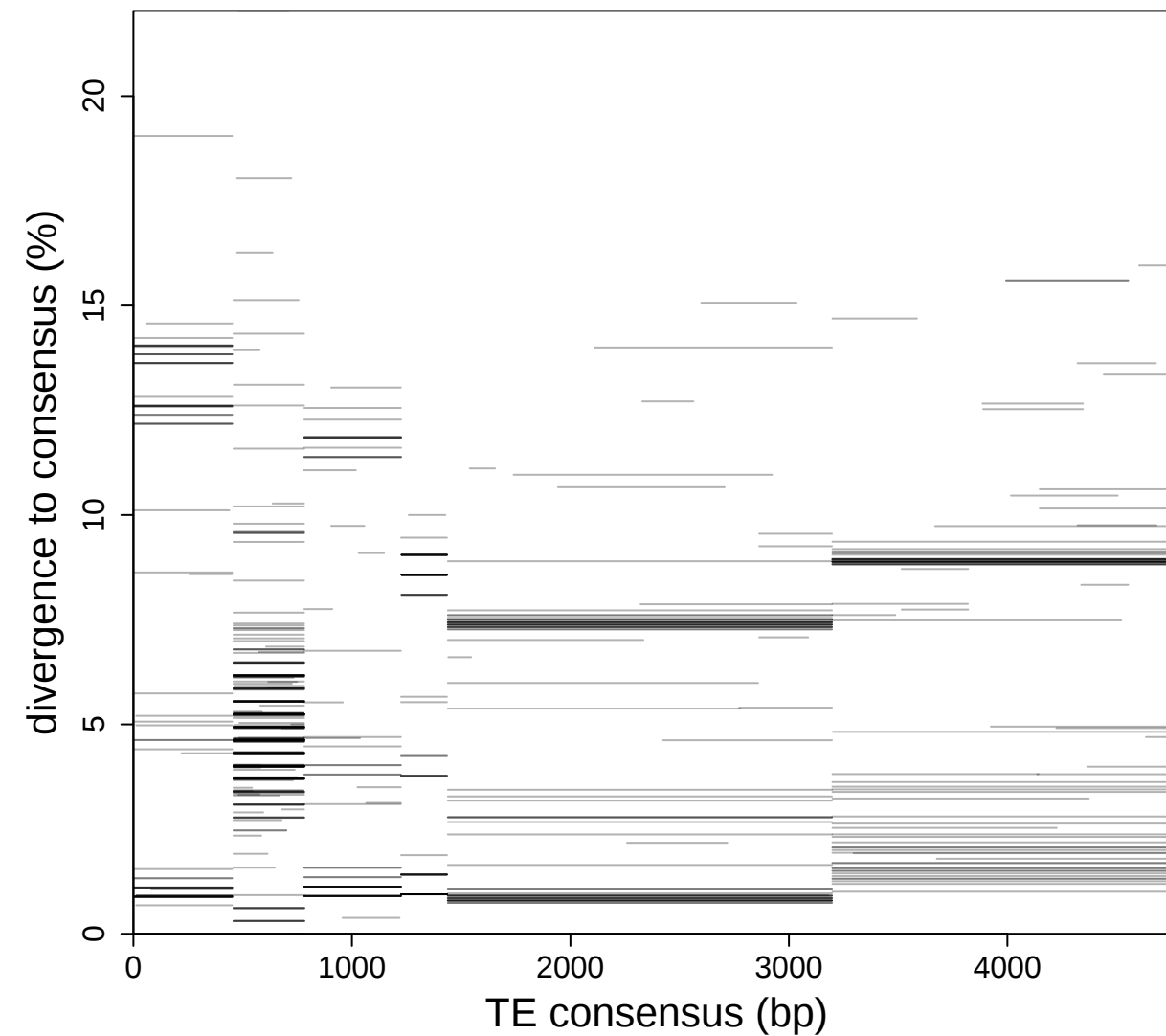
ly\_941.fasta.b.bed\_uf.bed\_g\_1.bed\_fm\_1.bed\_0\_0\_bcln.fa\_aln.fa  
size: 8870bp; fragments: 272; full length: 0 (>=7983bp)

# After TEtrimmer Extended plot Blue lines are boundaries

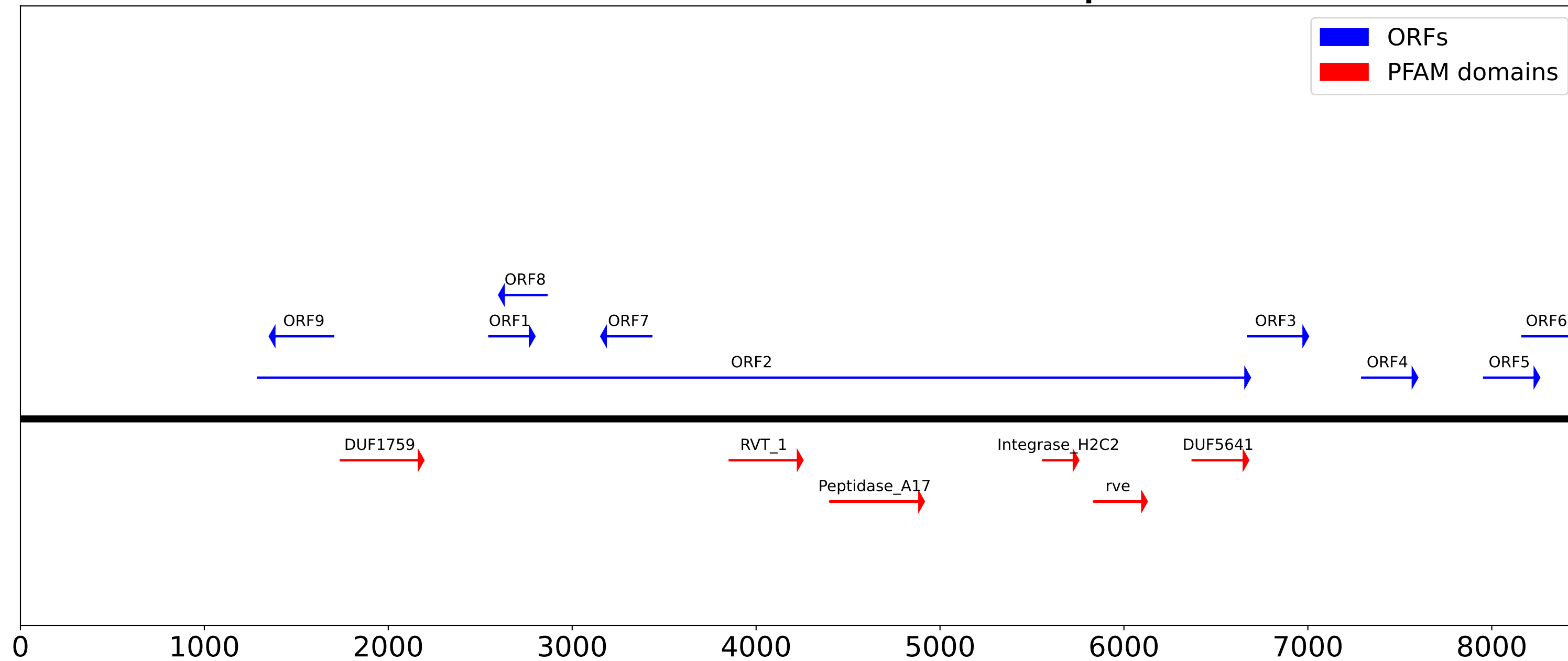


TE: ltr\_1\_family\_941  
size: 4790bp; fragments: 580; full length: 0 (>=4311bp)

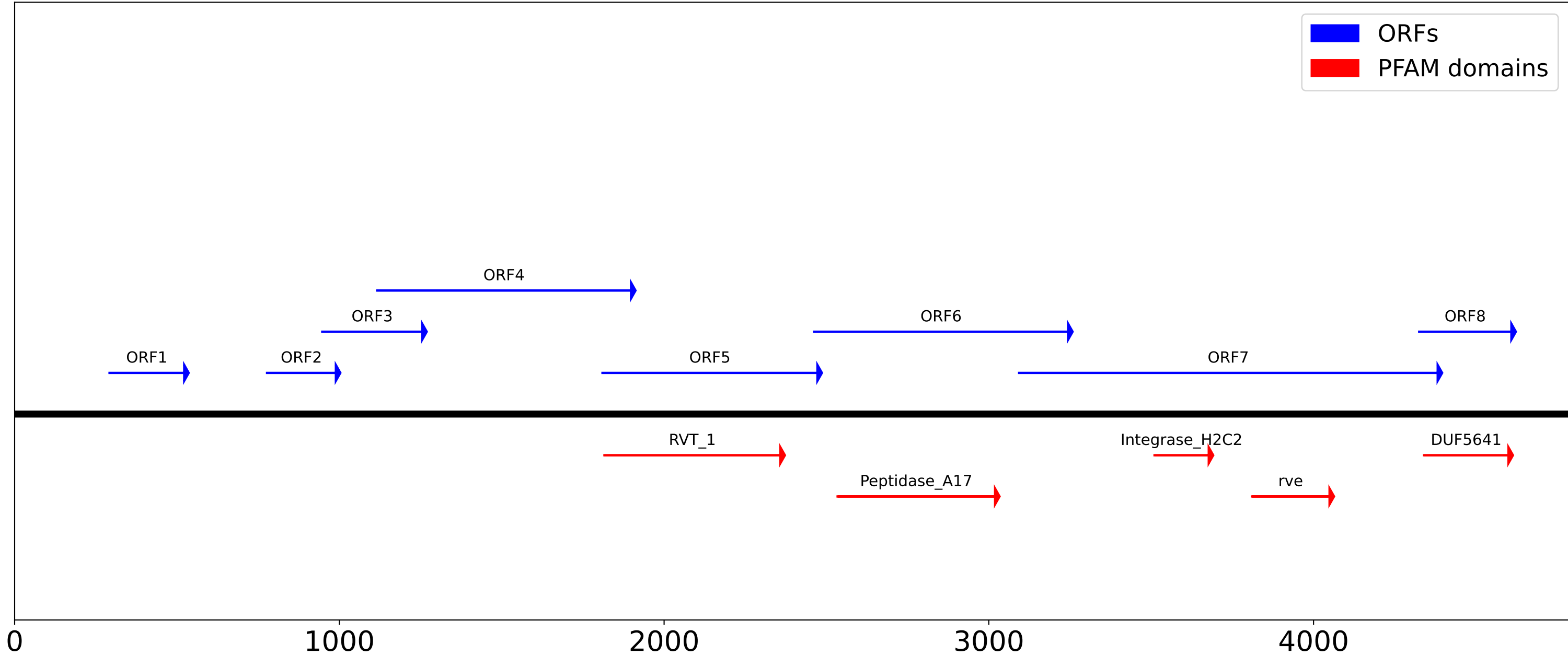
## Before TEtrimmer 4790 bp



## After TEtrimmer ORF and PFAM domain plot



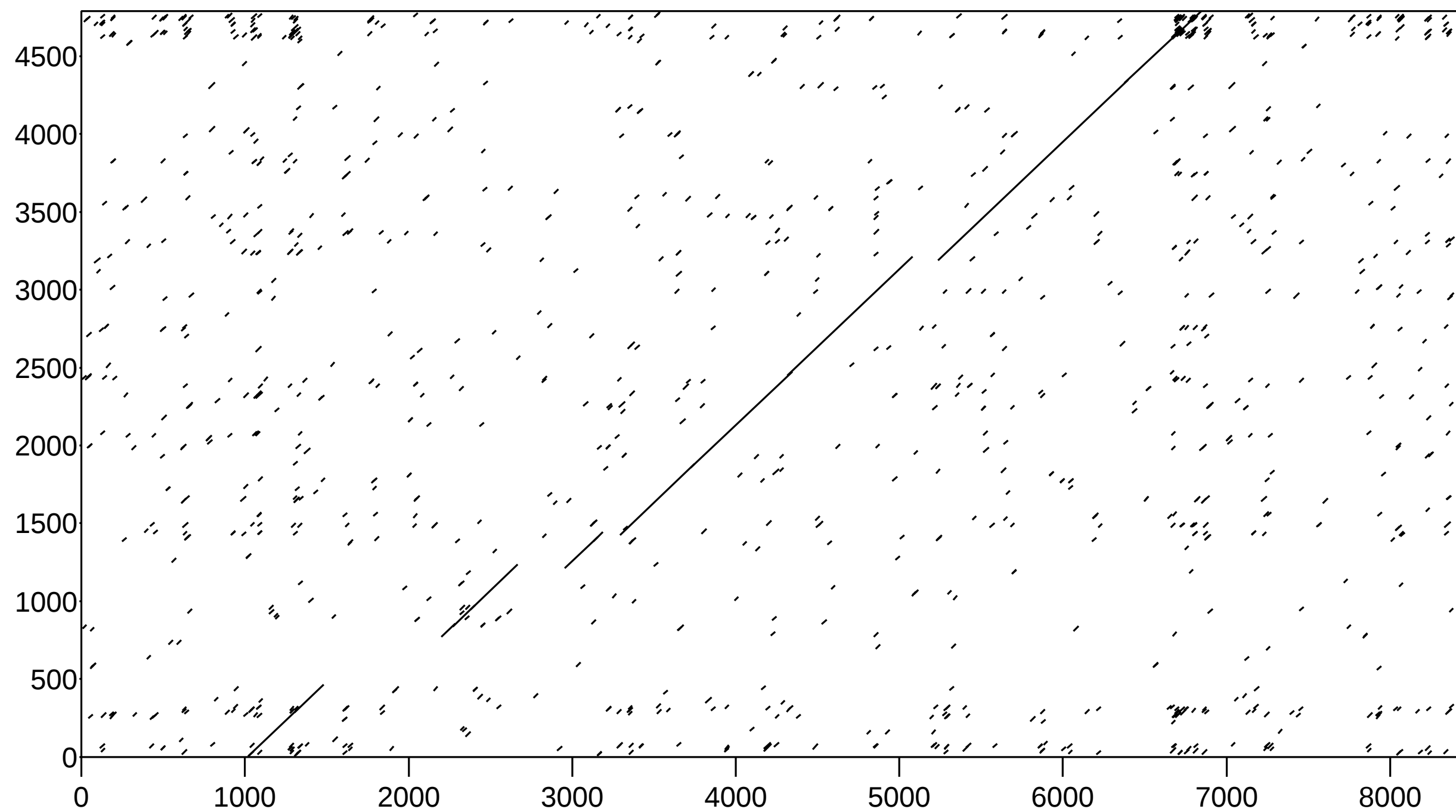
# Before TEtrimmer ORF and PFAM domain plot



# Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)