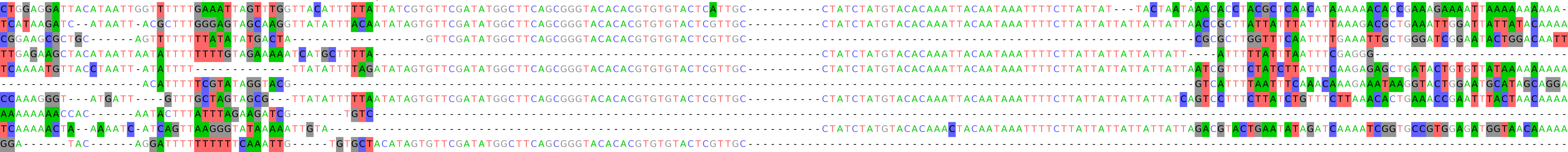


Start Crop Point

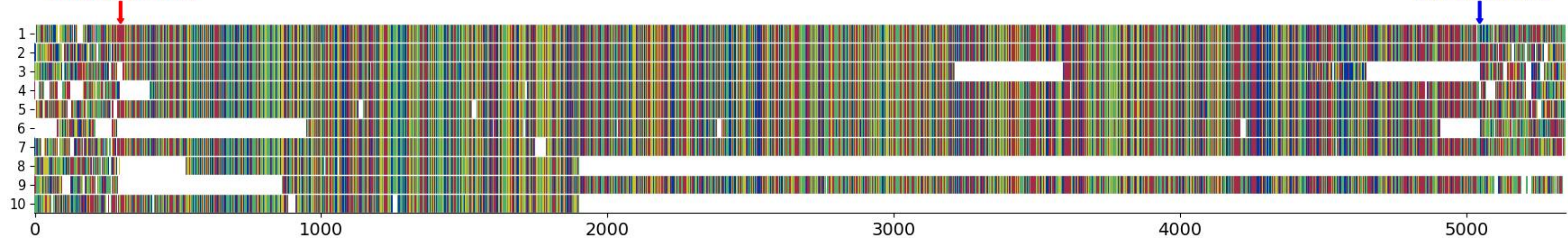
End Crop Point

MSA length = 4748



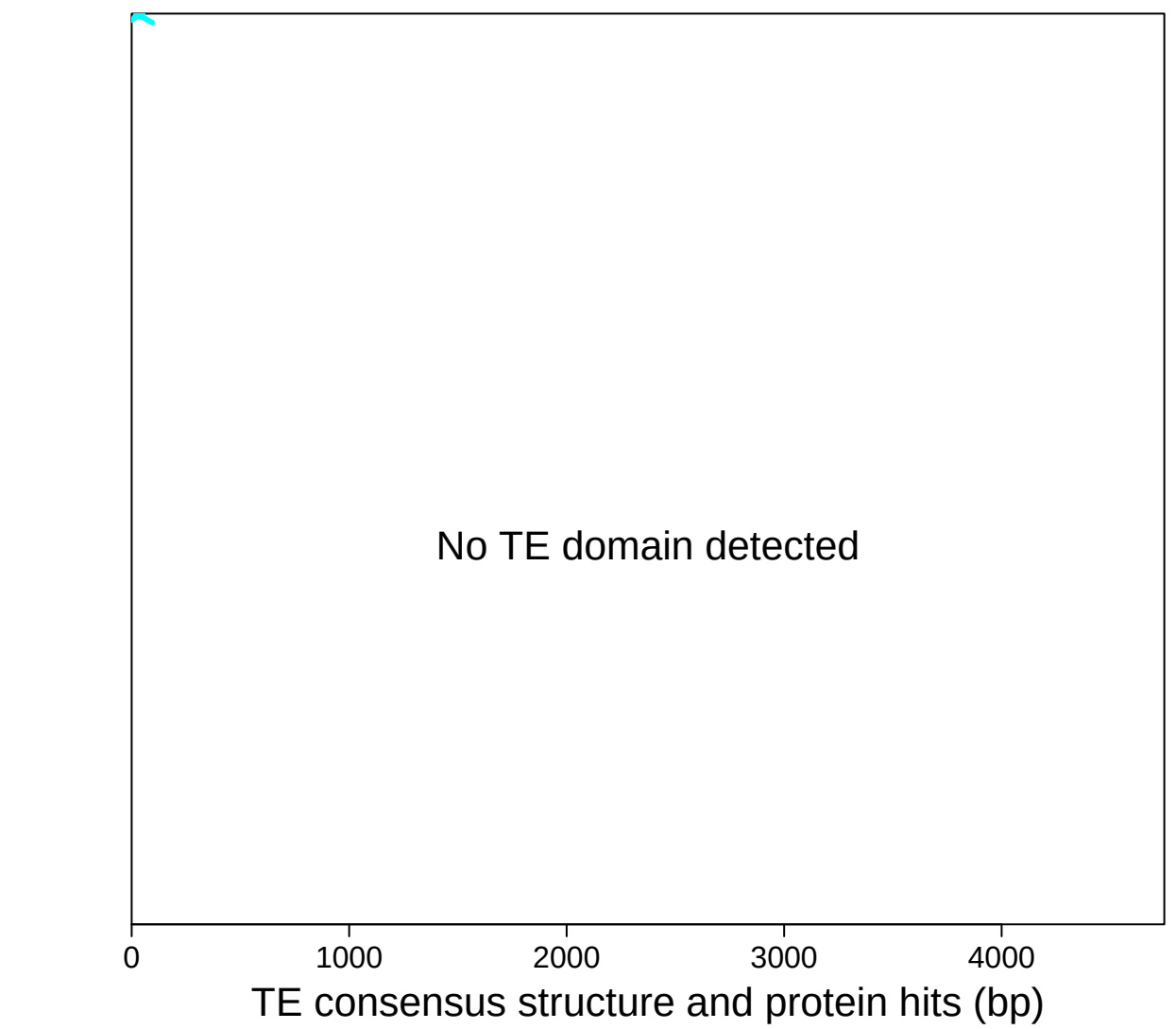
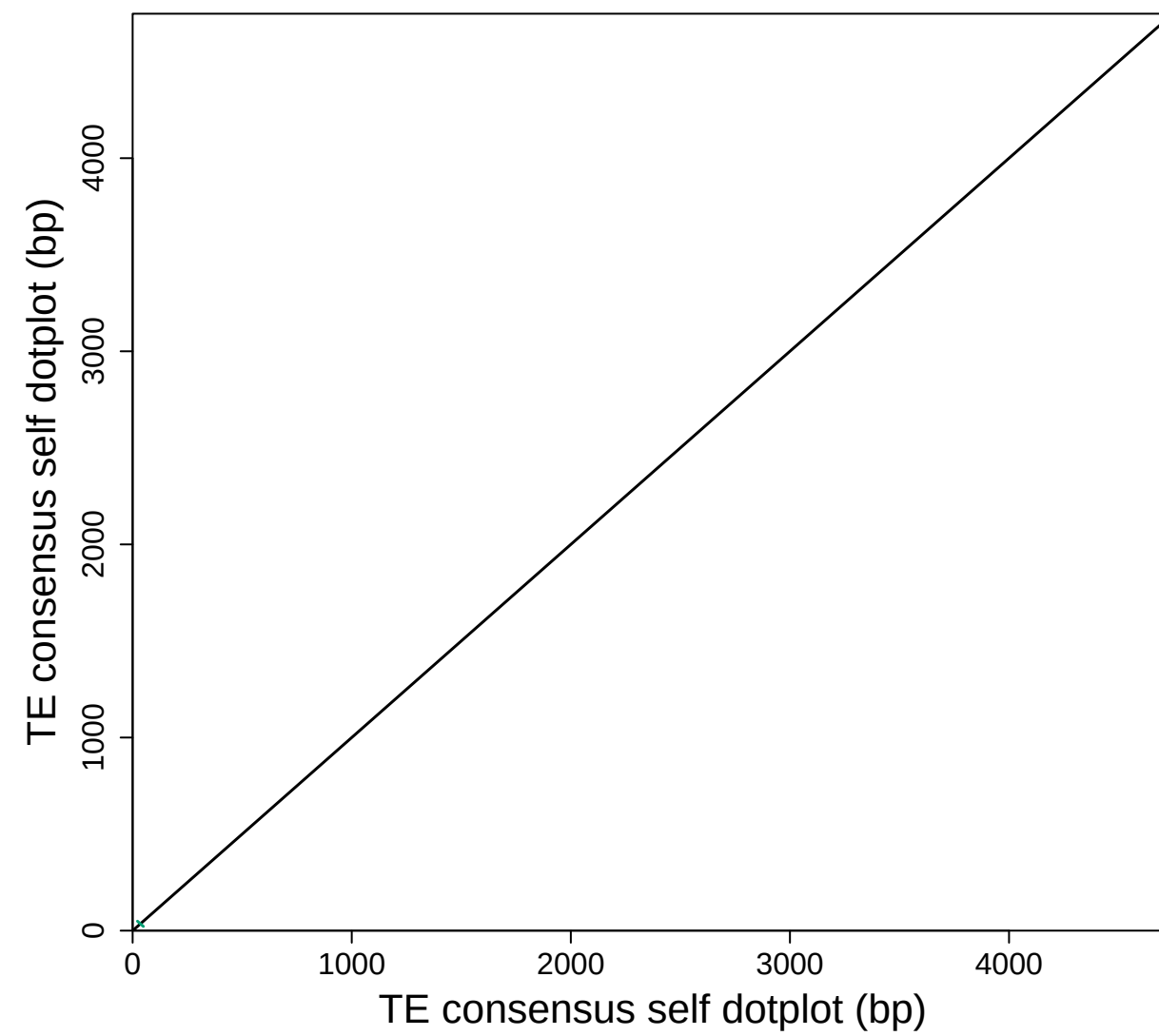
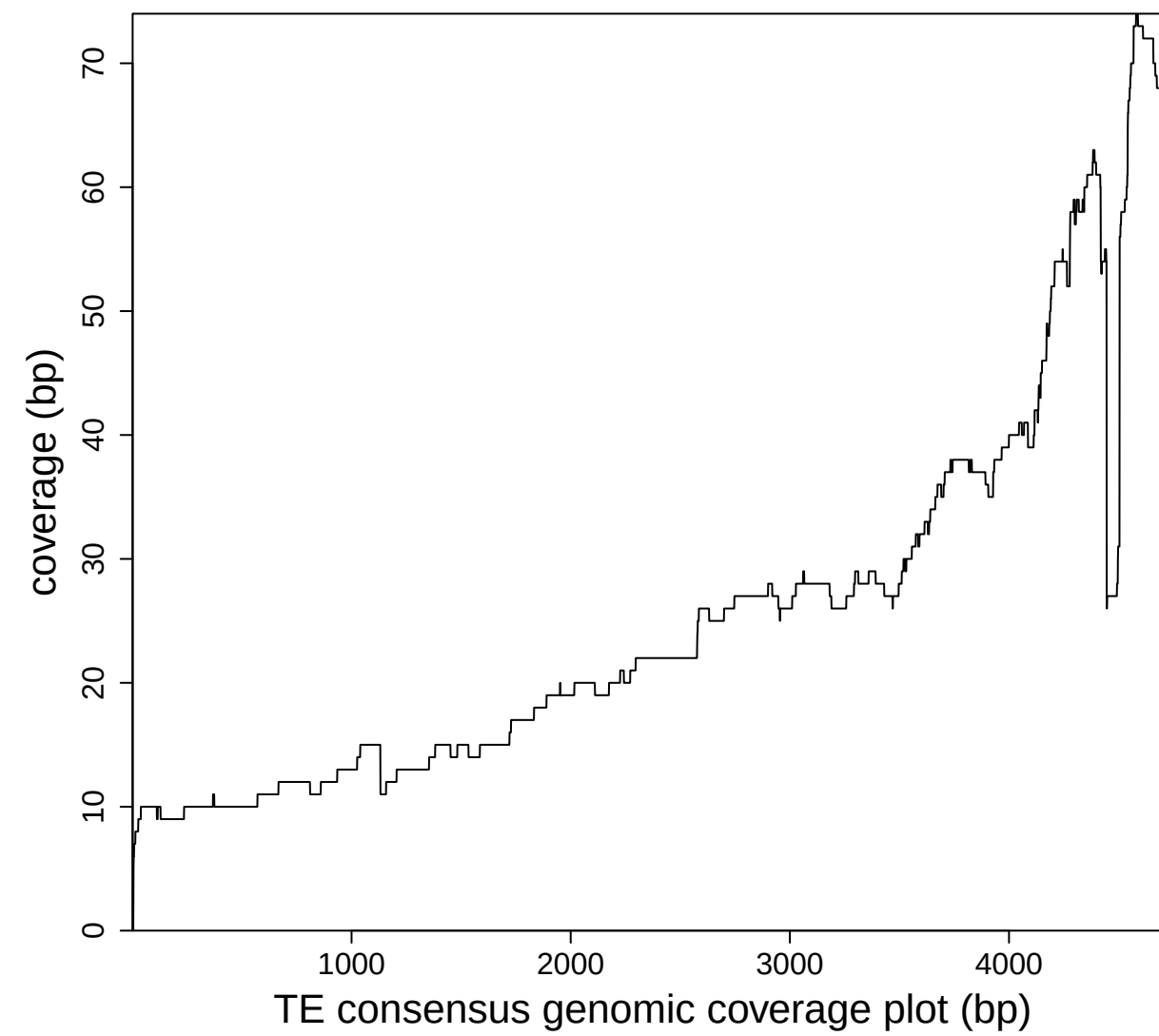
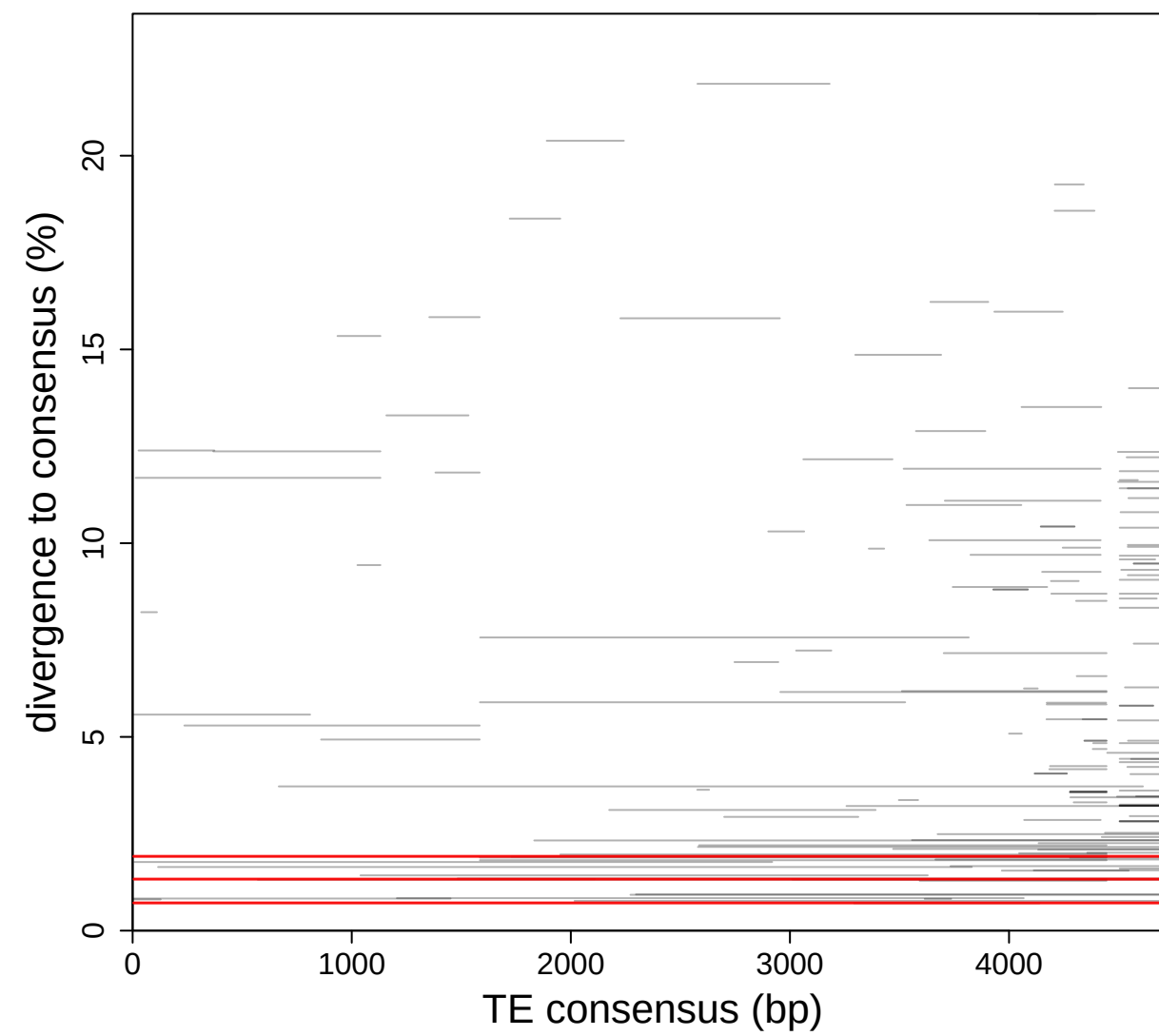
Start crop Point

End crop Point



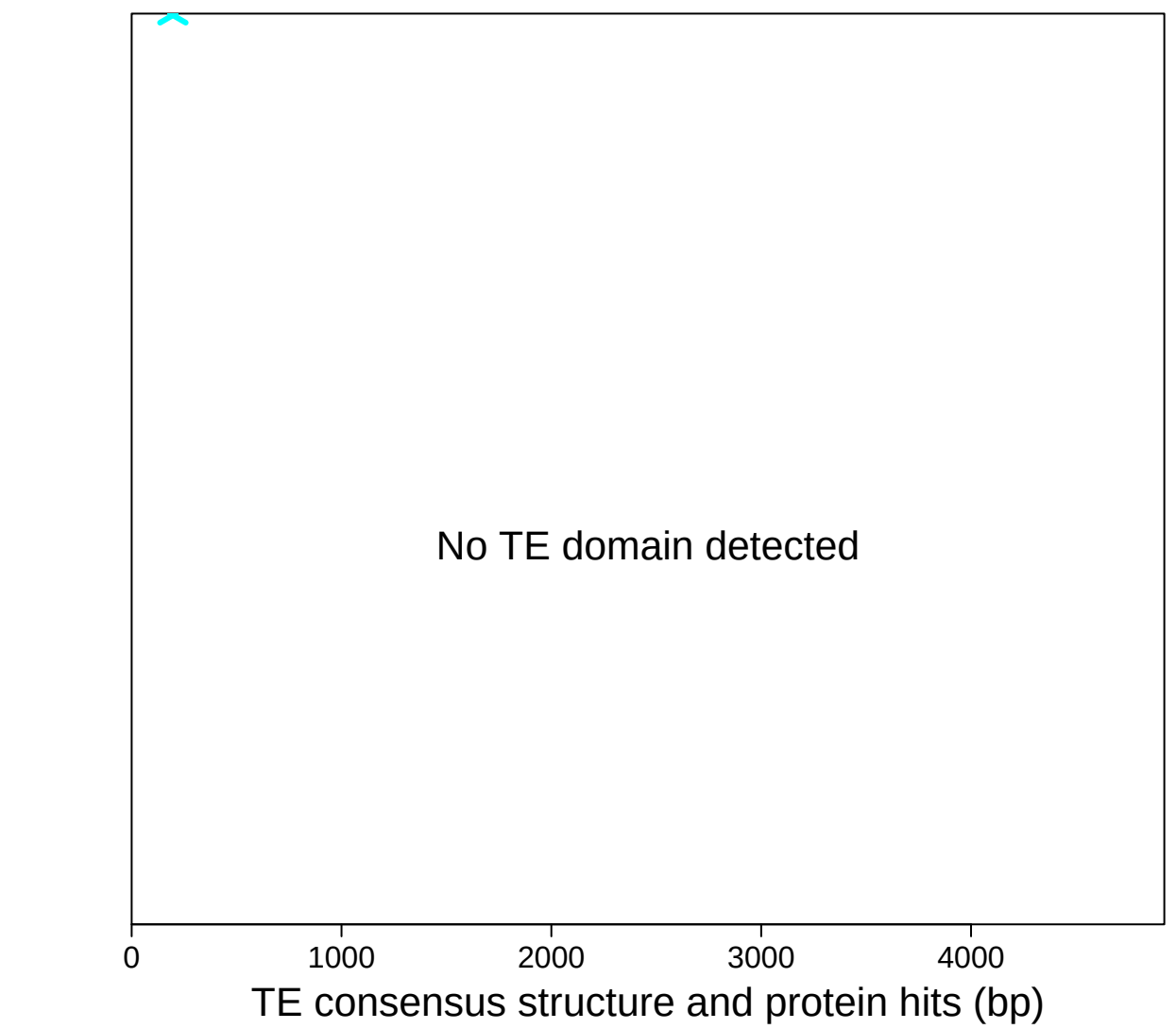
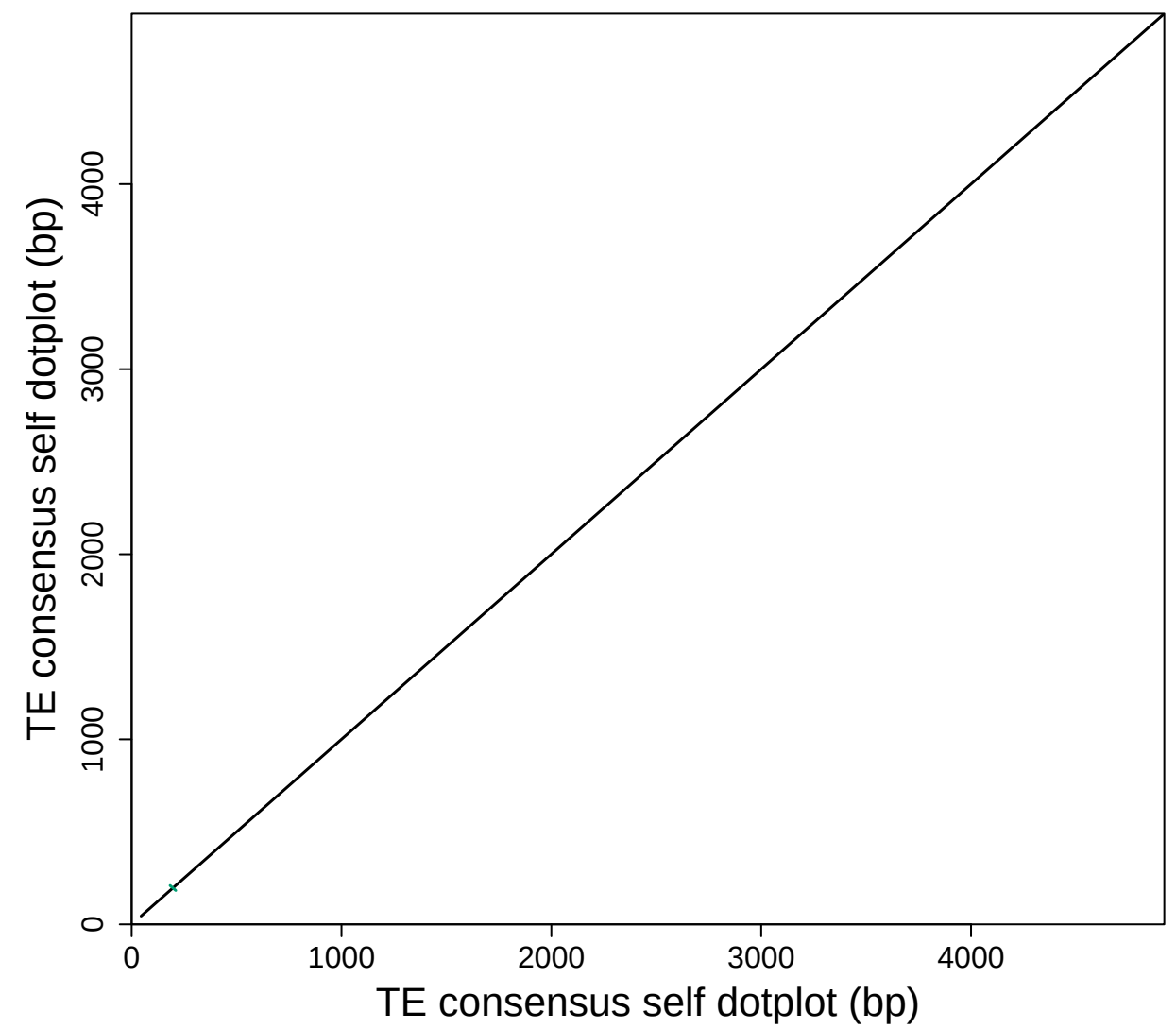
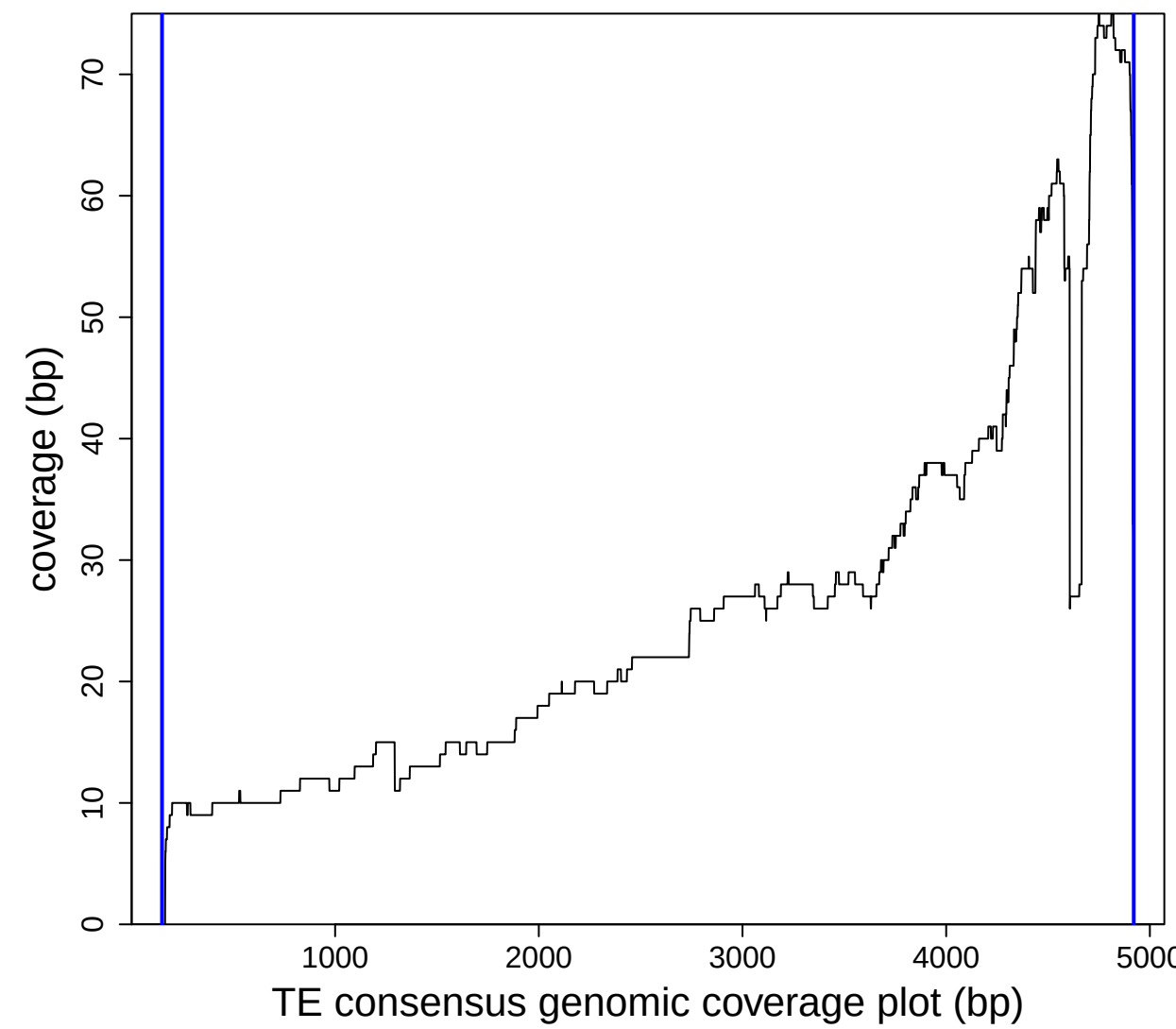
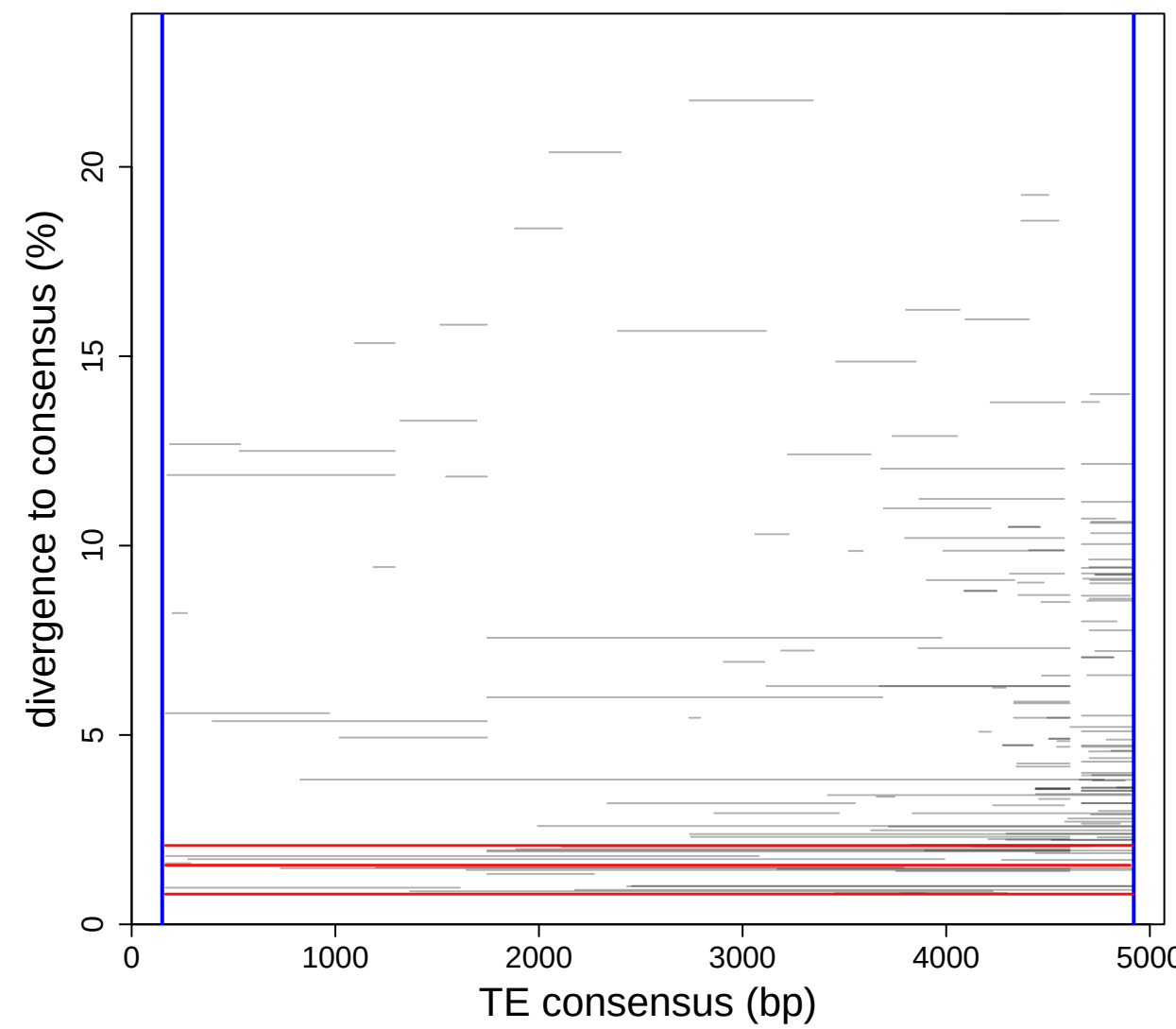
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 4748bp; fragments: 166; full length: 3 (>=4273.2bp)

After TEtrimmer 4748 bp



1.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 5071bp; fragments: 171; full length: 3 (>=4563.9bp)

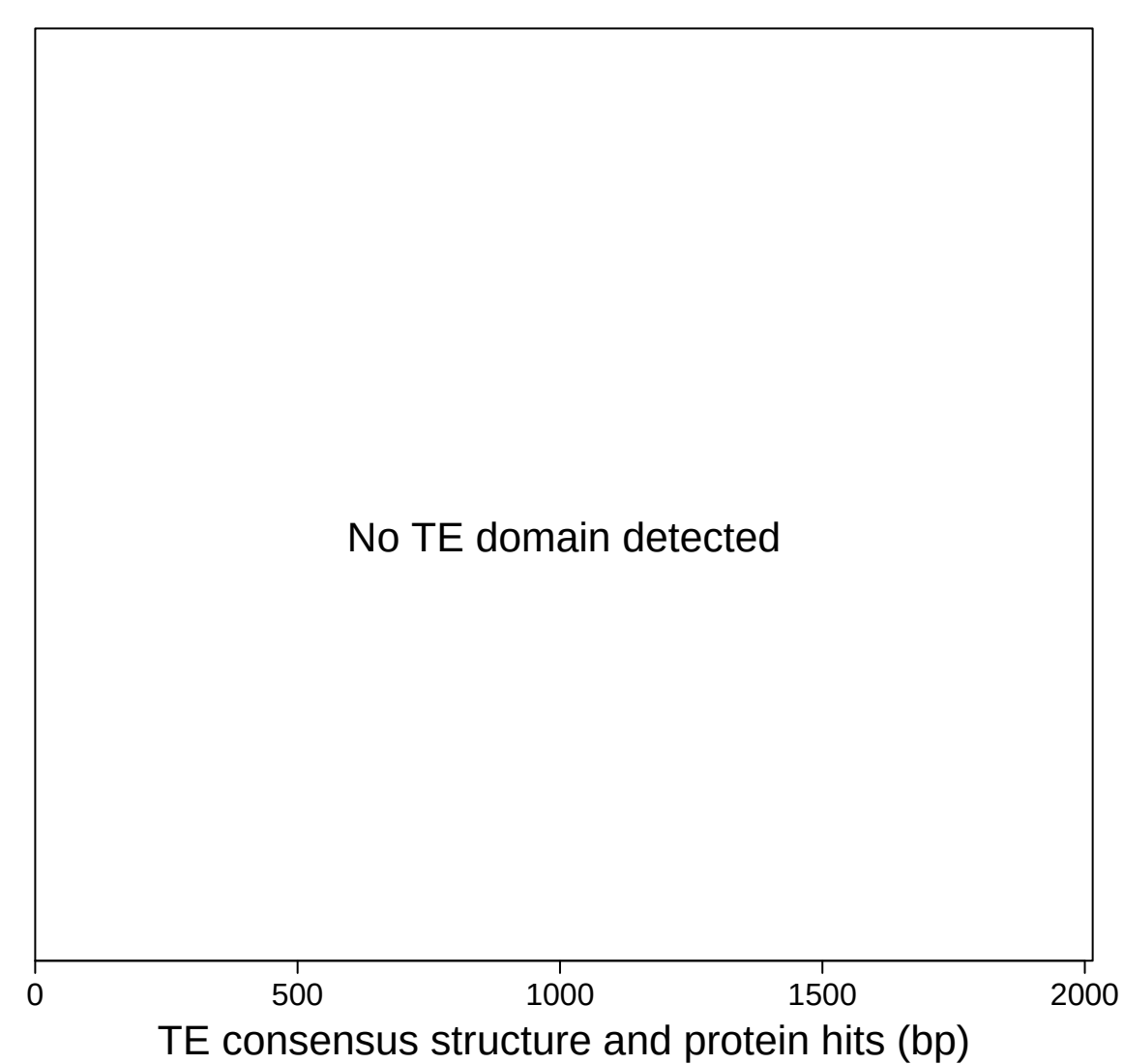
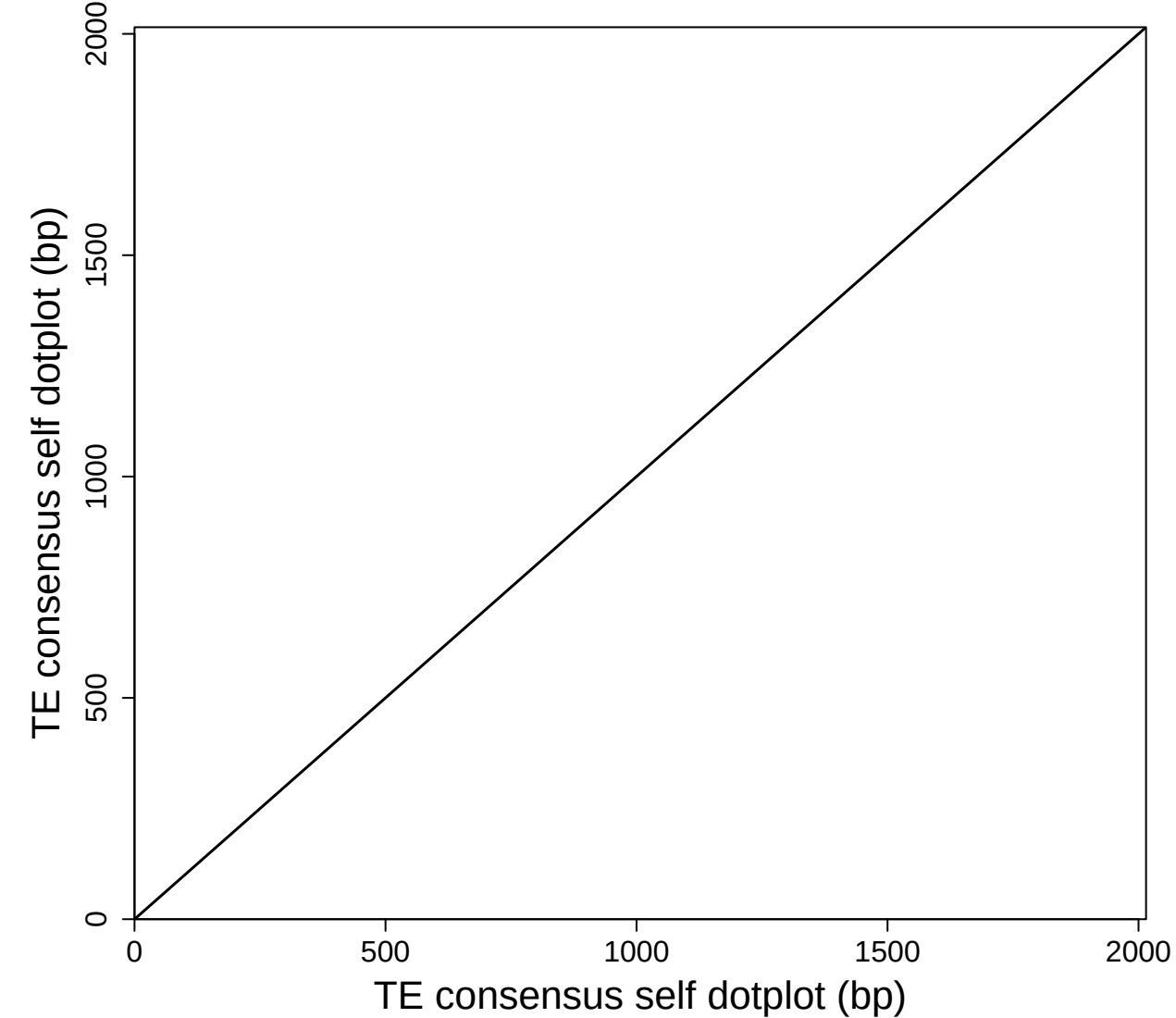
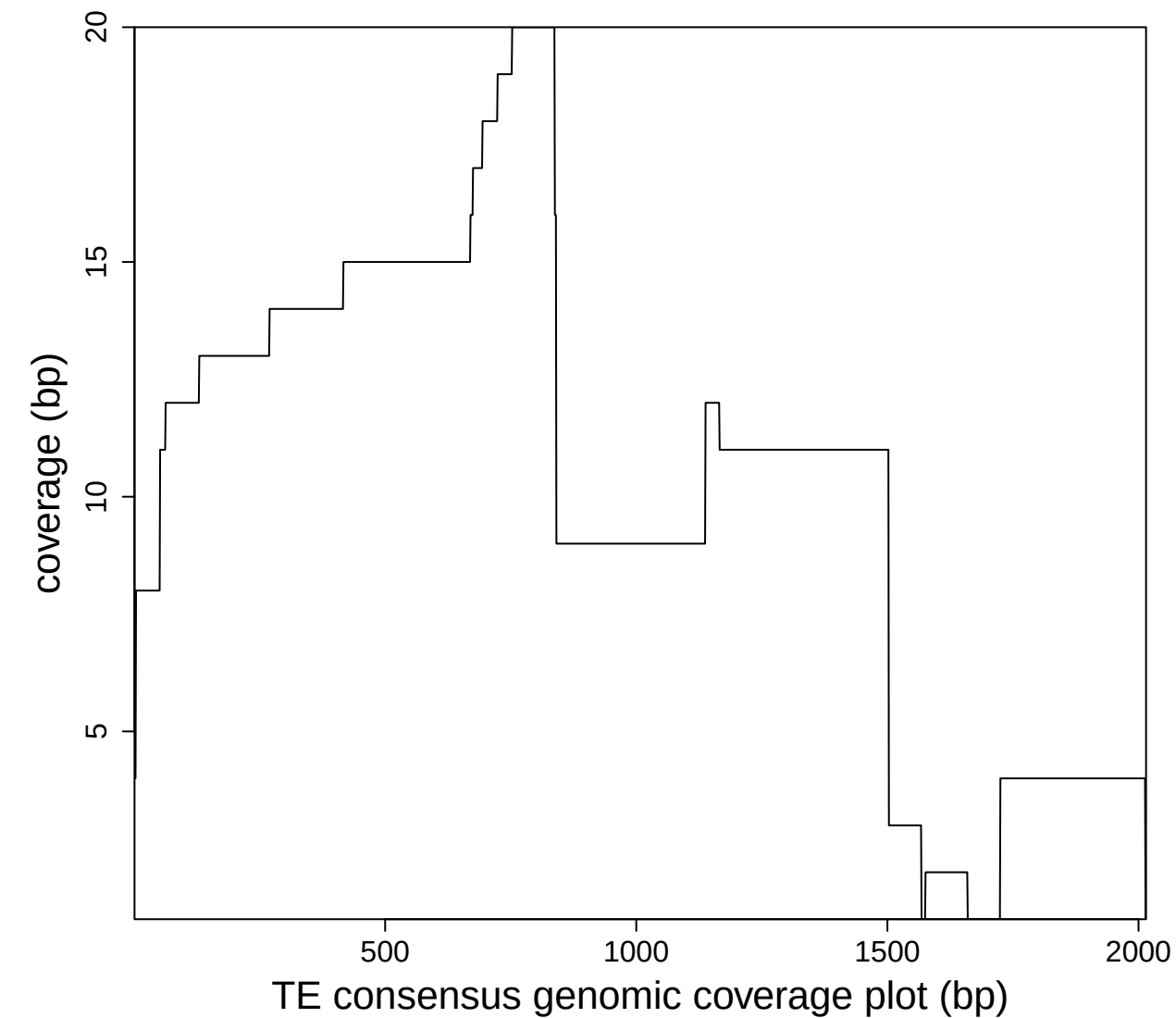
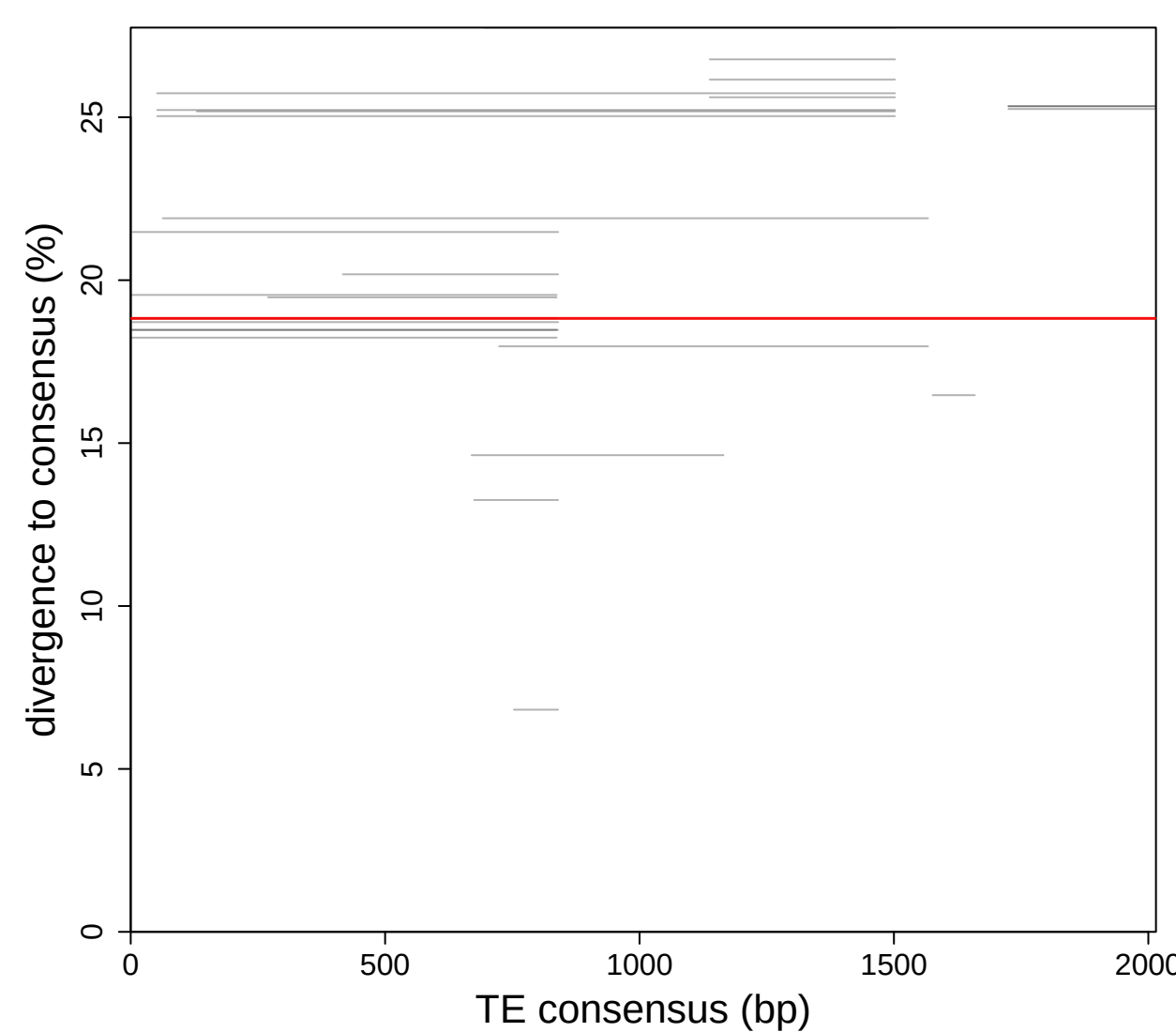
After TETrimmer Extended plot Blue lines are boundaries



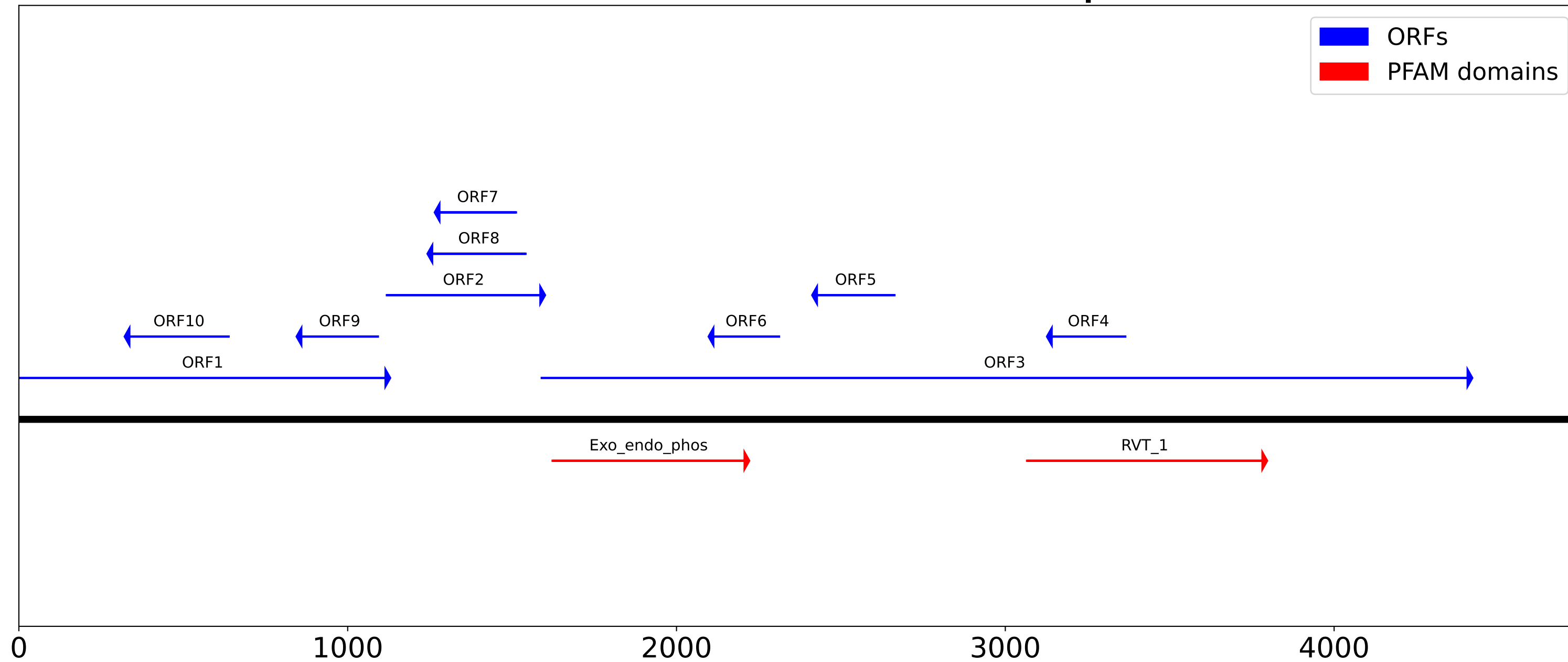
Before TEtrimmer 2015 bp

TE: rnd_5_family_991

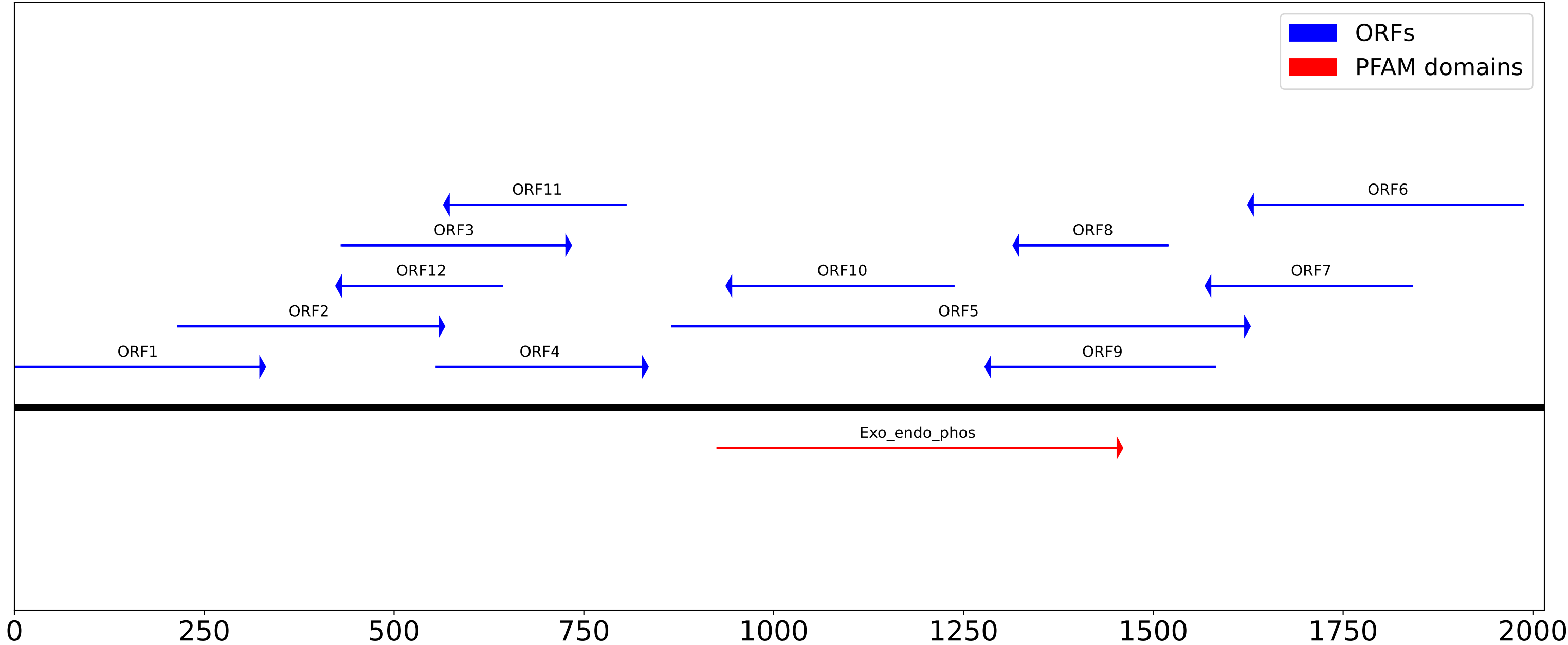
size: 2015bp; fragments: 27; full length: 1 (≥ 1813.5 bp)



After TEtrimmer ORF and PFAM domain plot



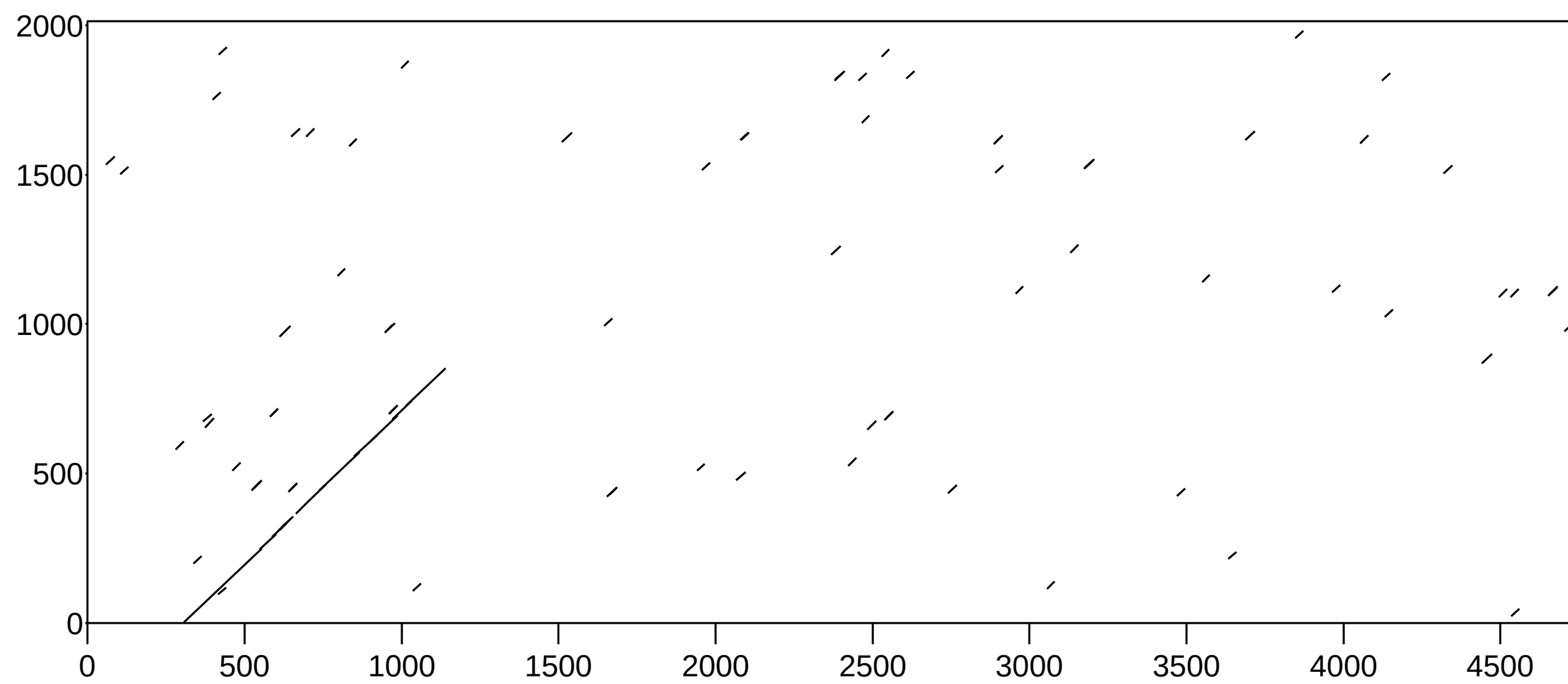
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)