

Start Crop Point



End Crop Point

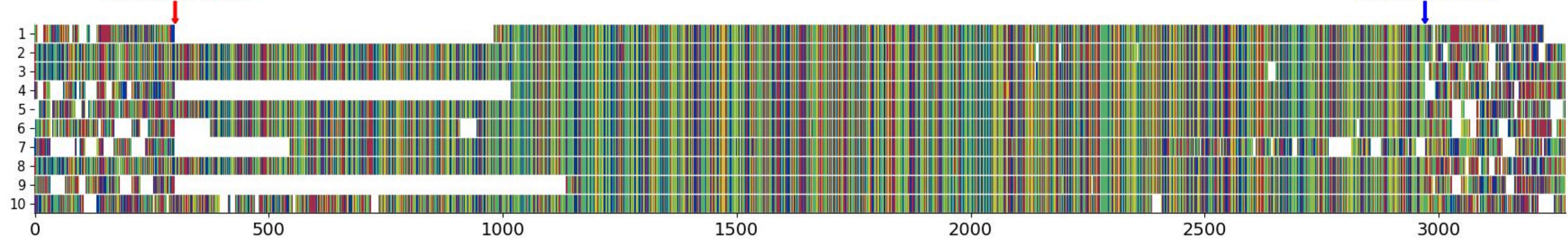


MSA length = 2672



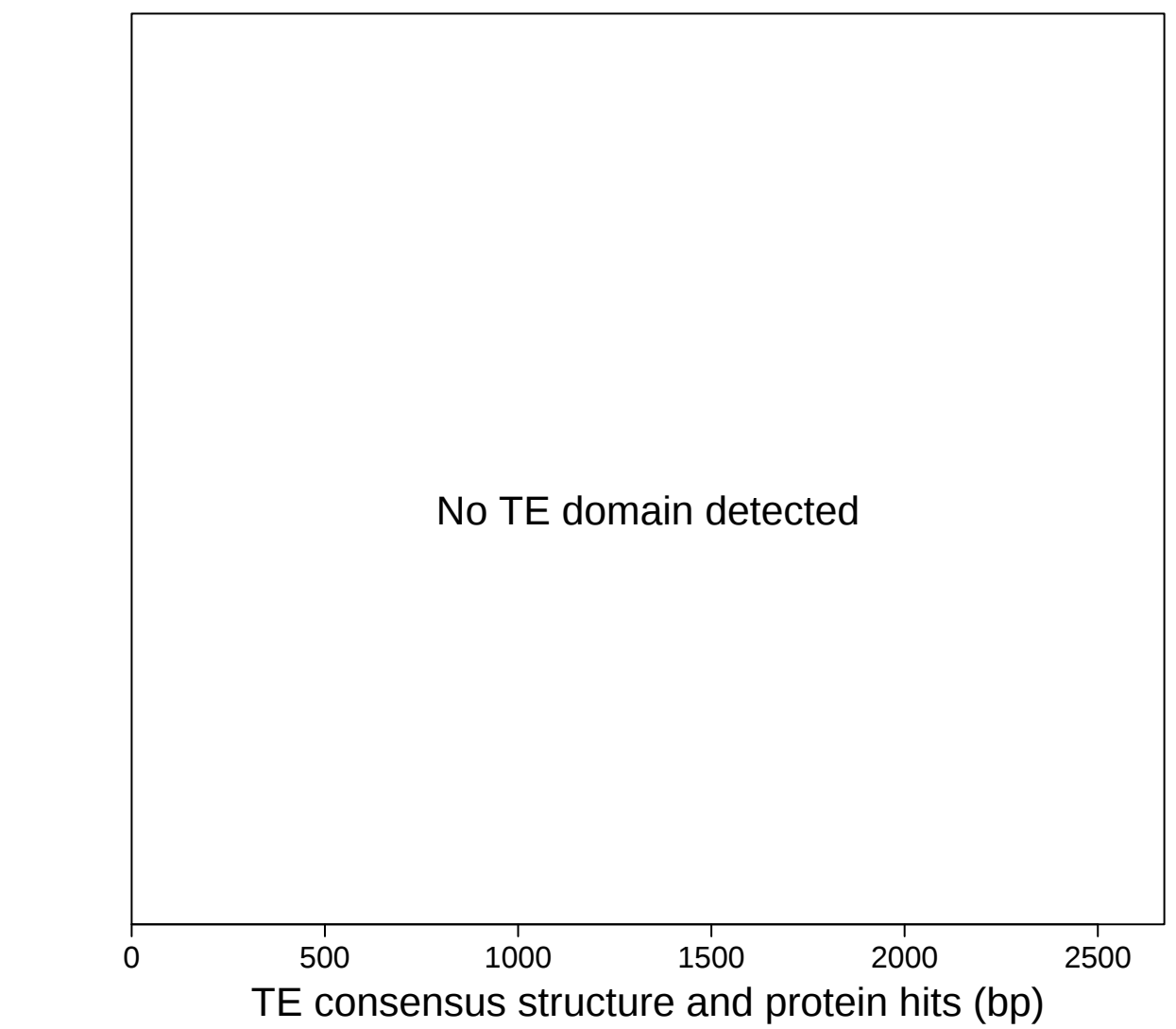
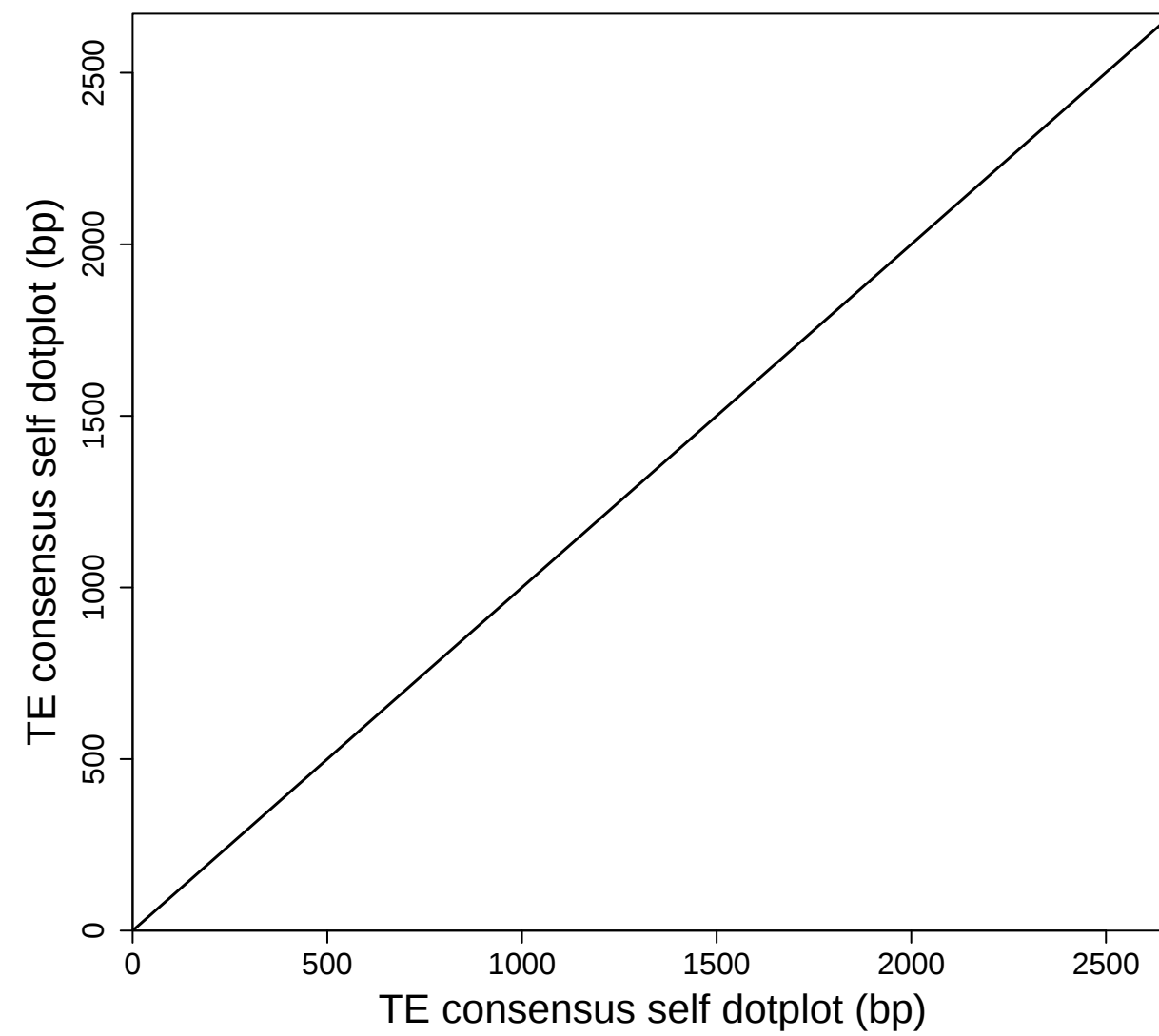
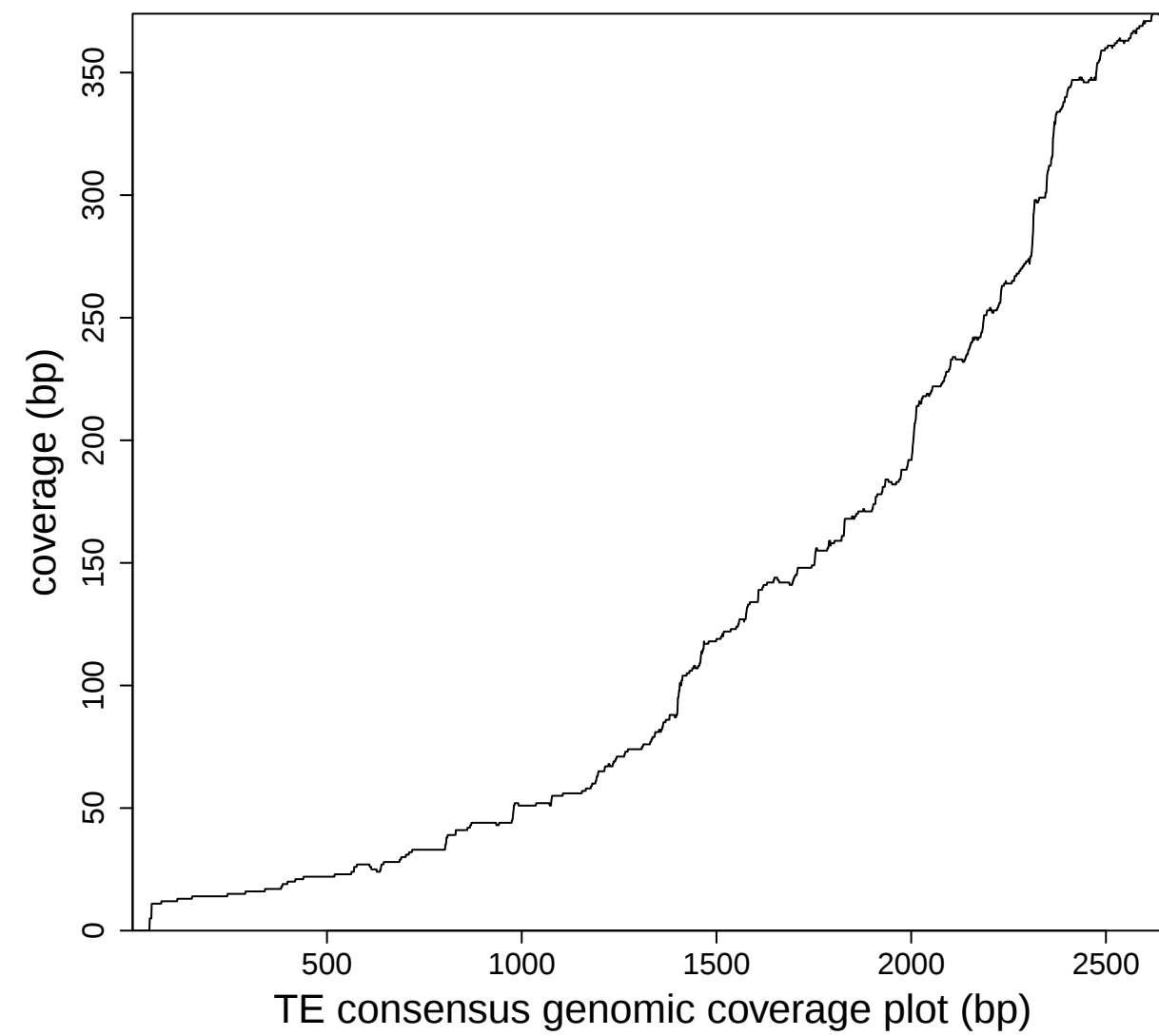
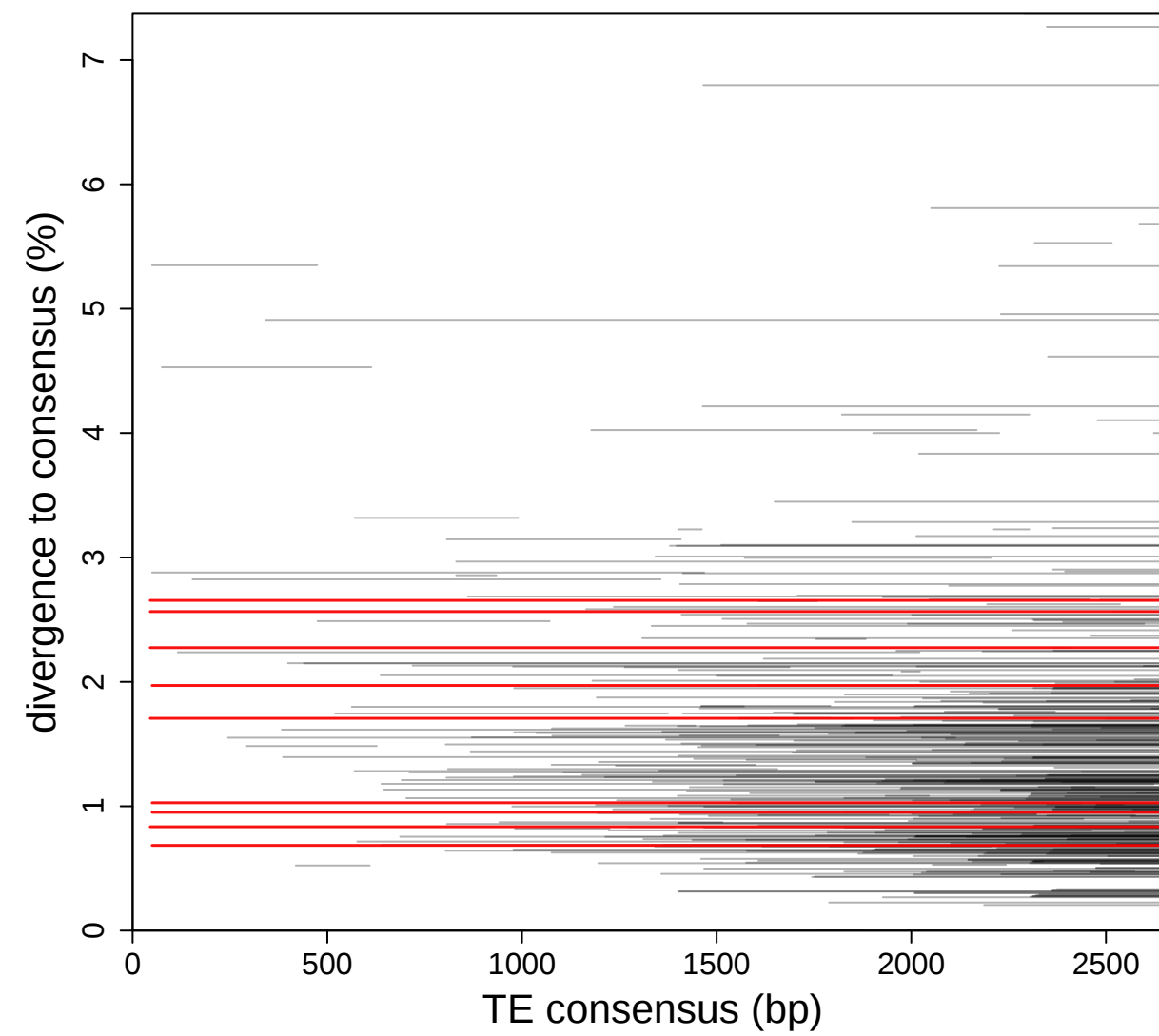
Start crop Point

End crop Point



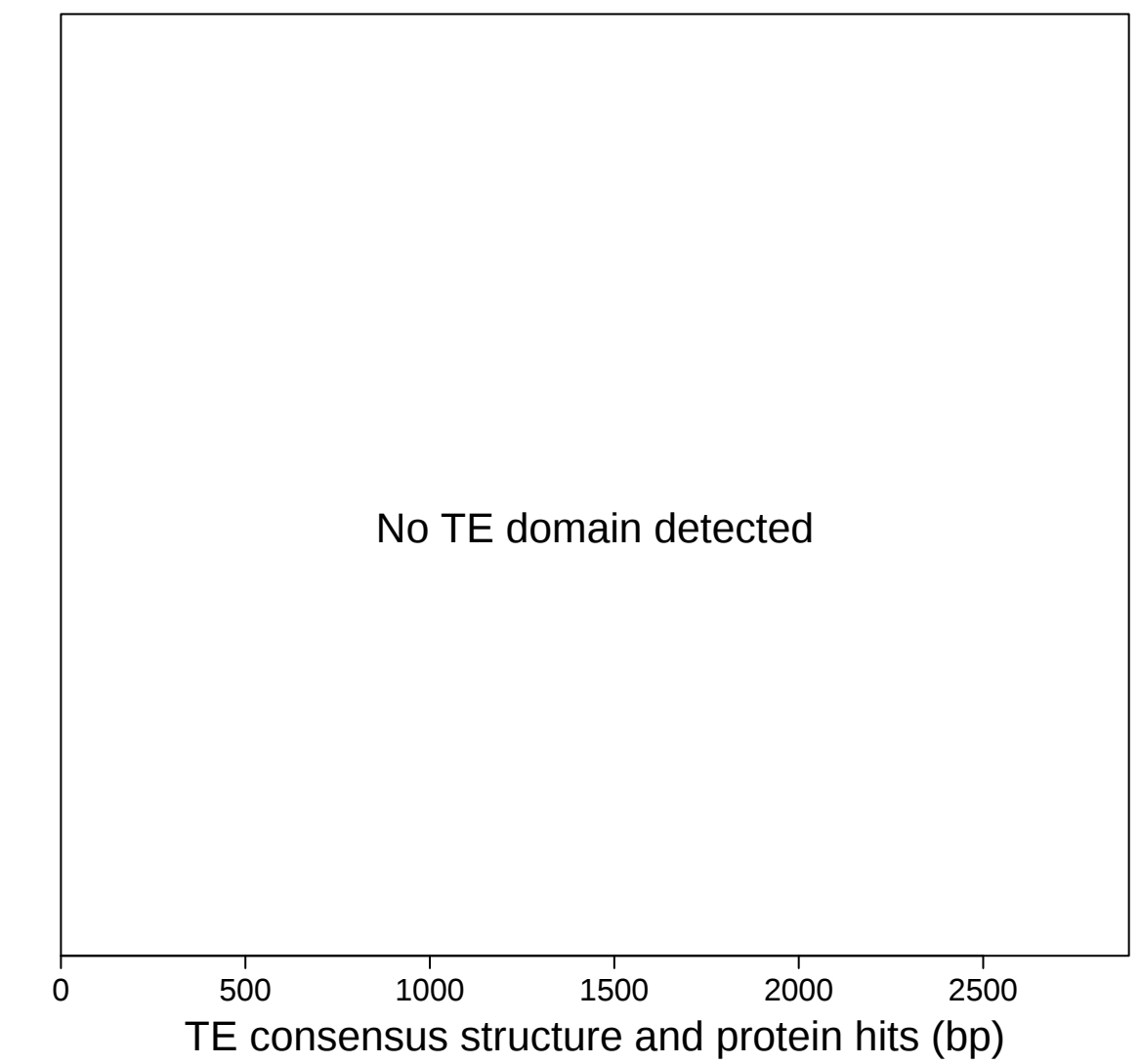
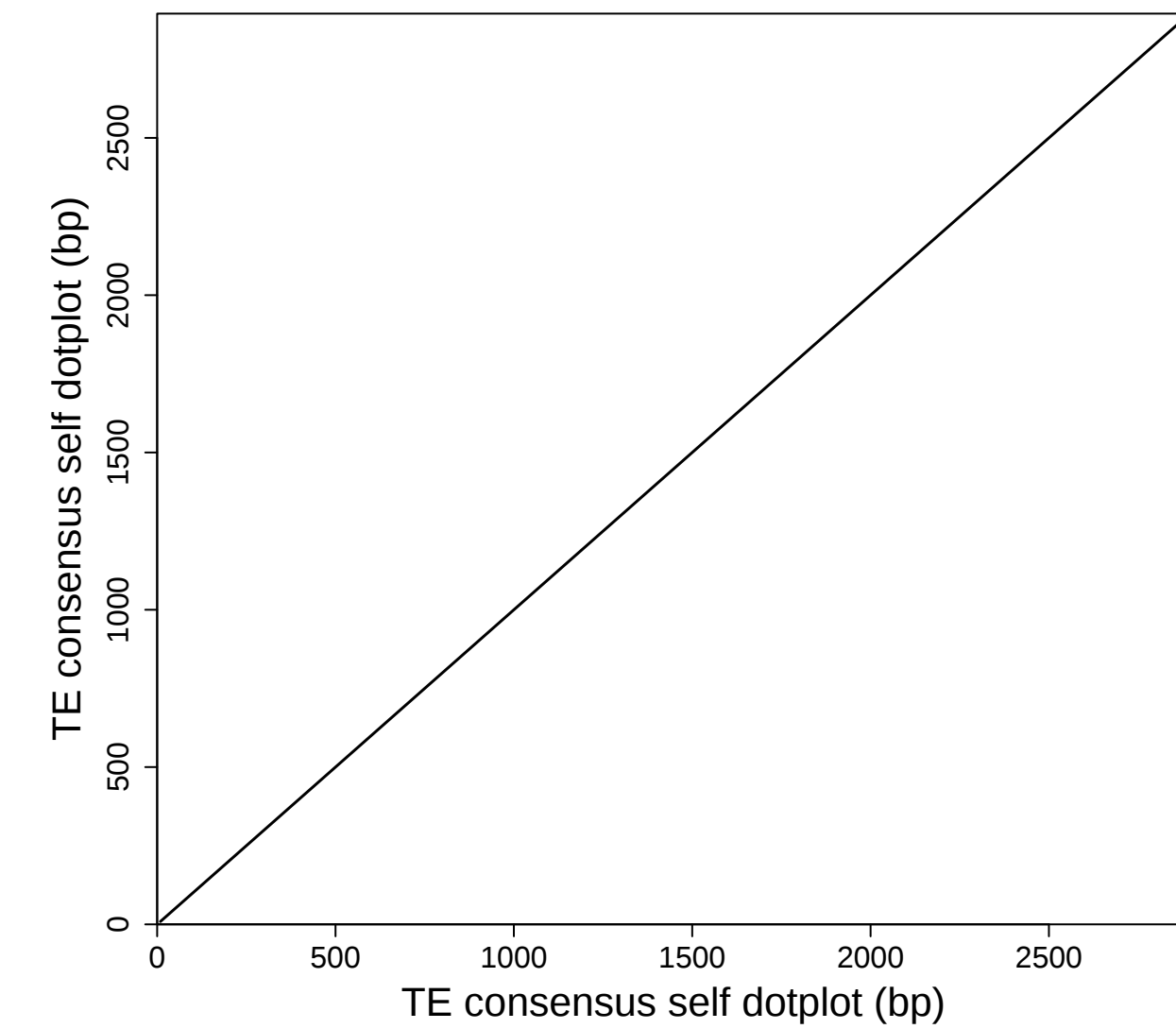
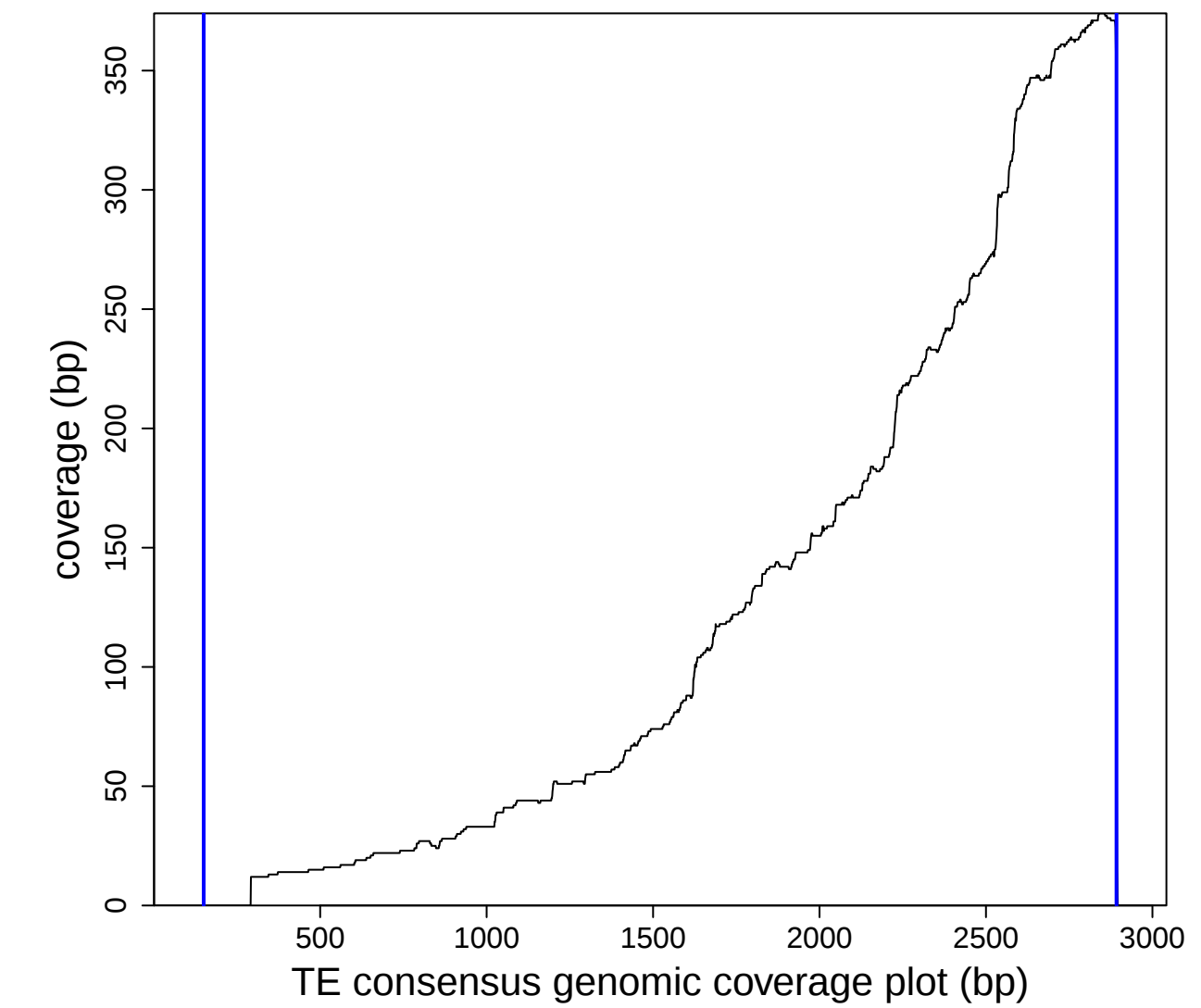
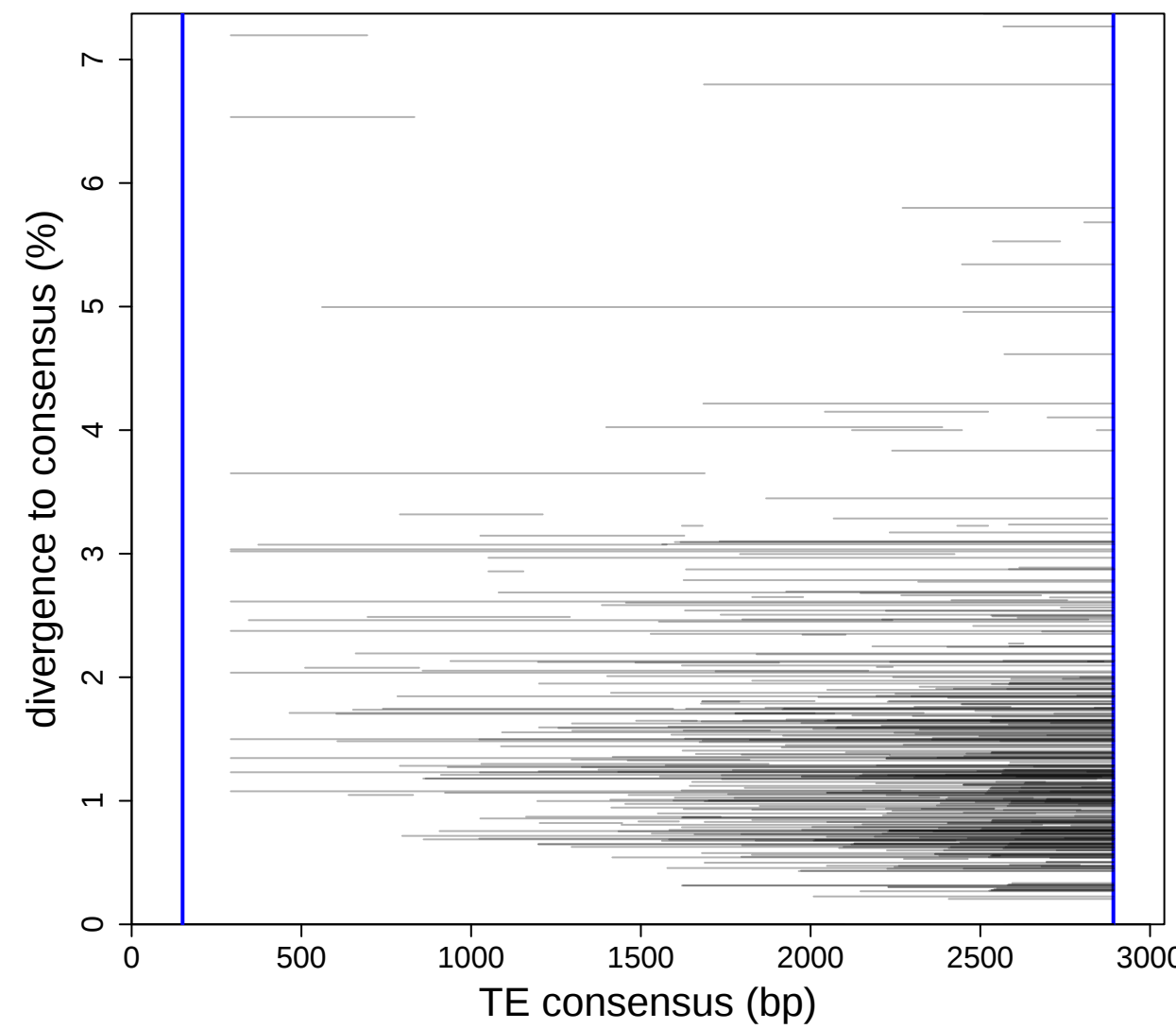
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 2672bp; fragments: 435; full length: 9 (>=2404.8bp)

After TEtrimmer 2672 bp

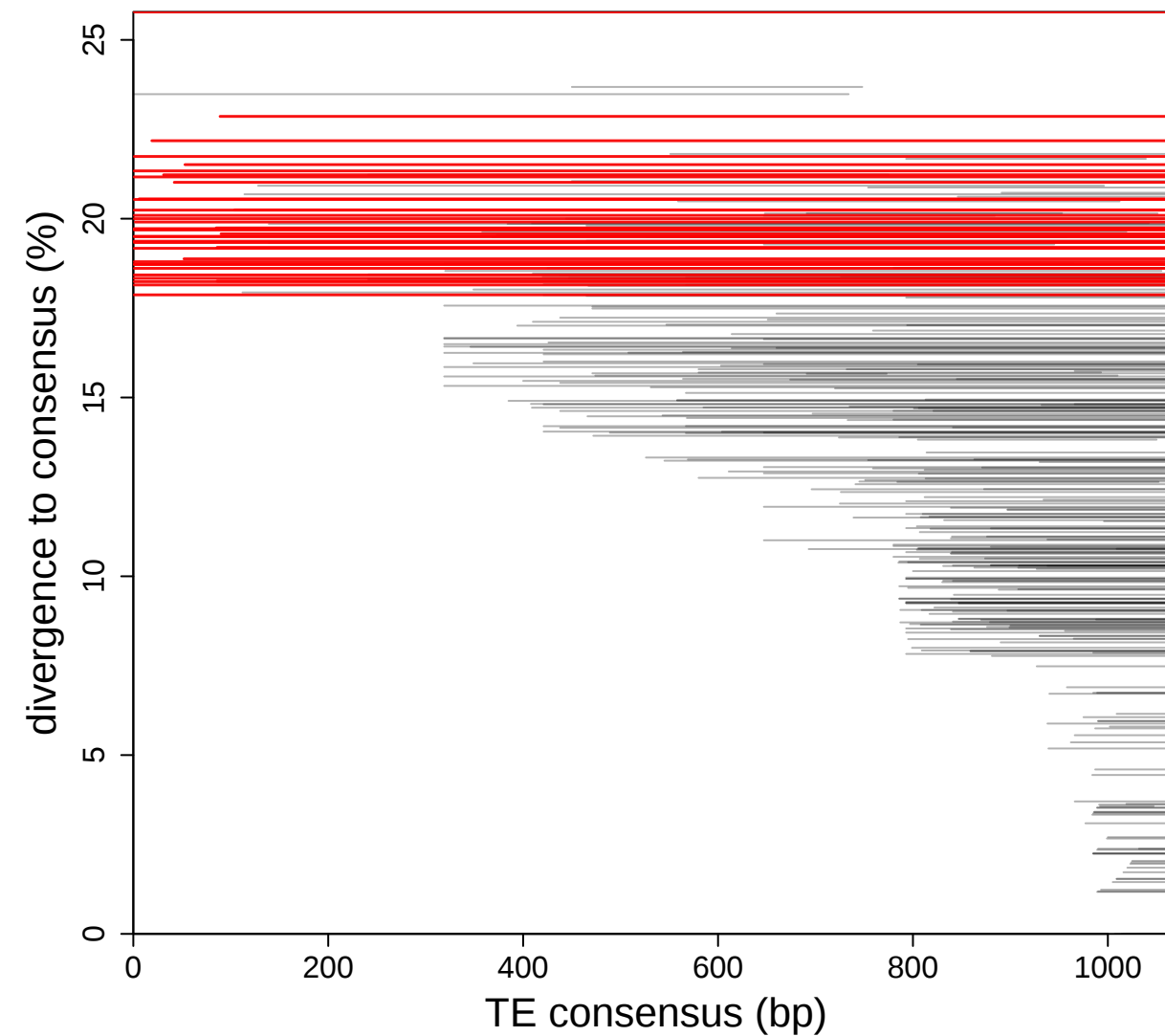


0.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 3042bp; fragments: 435; full length: 0 (>=2737.8bp)

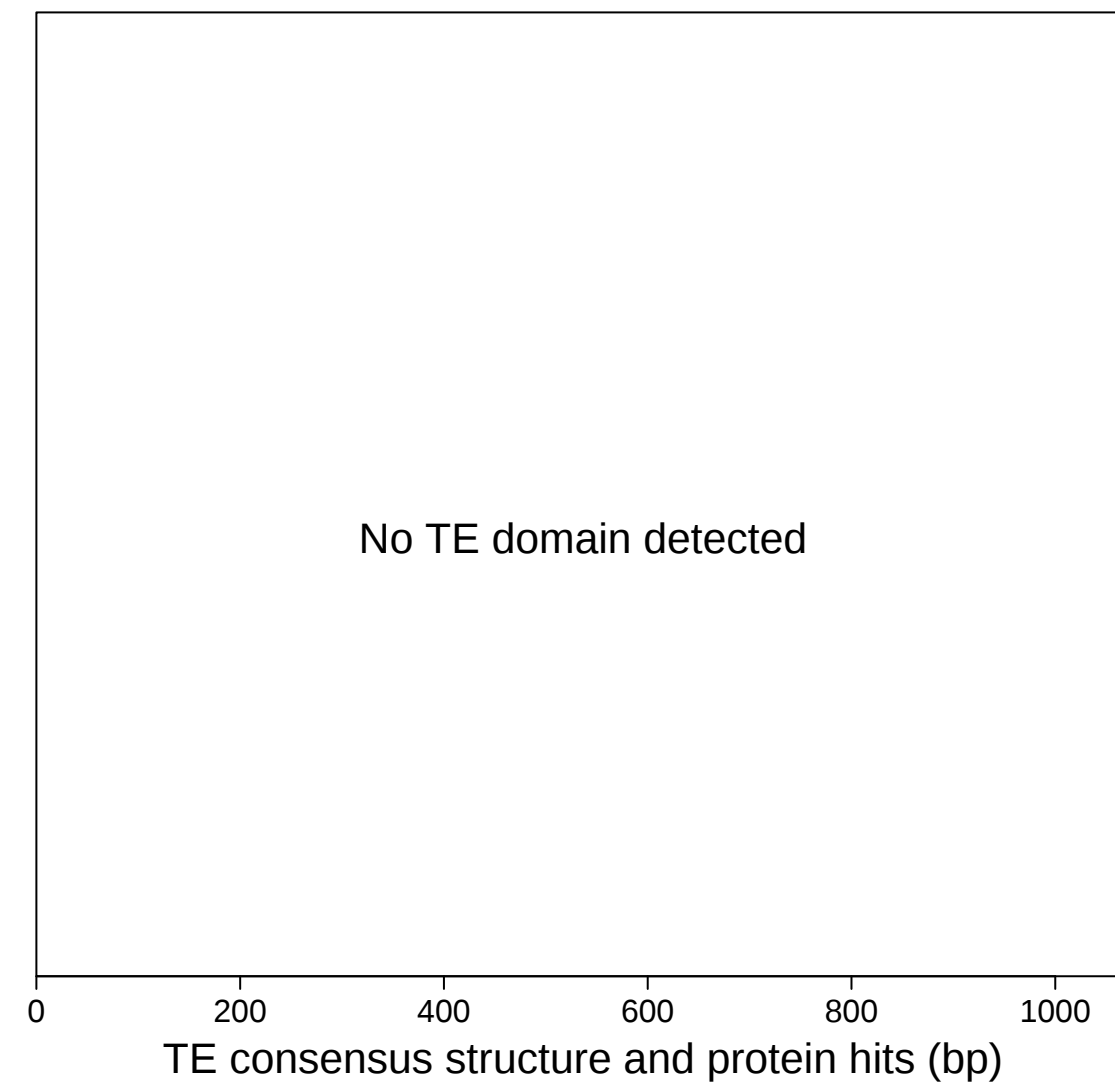
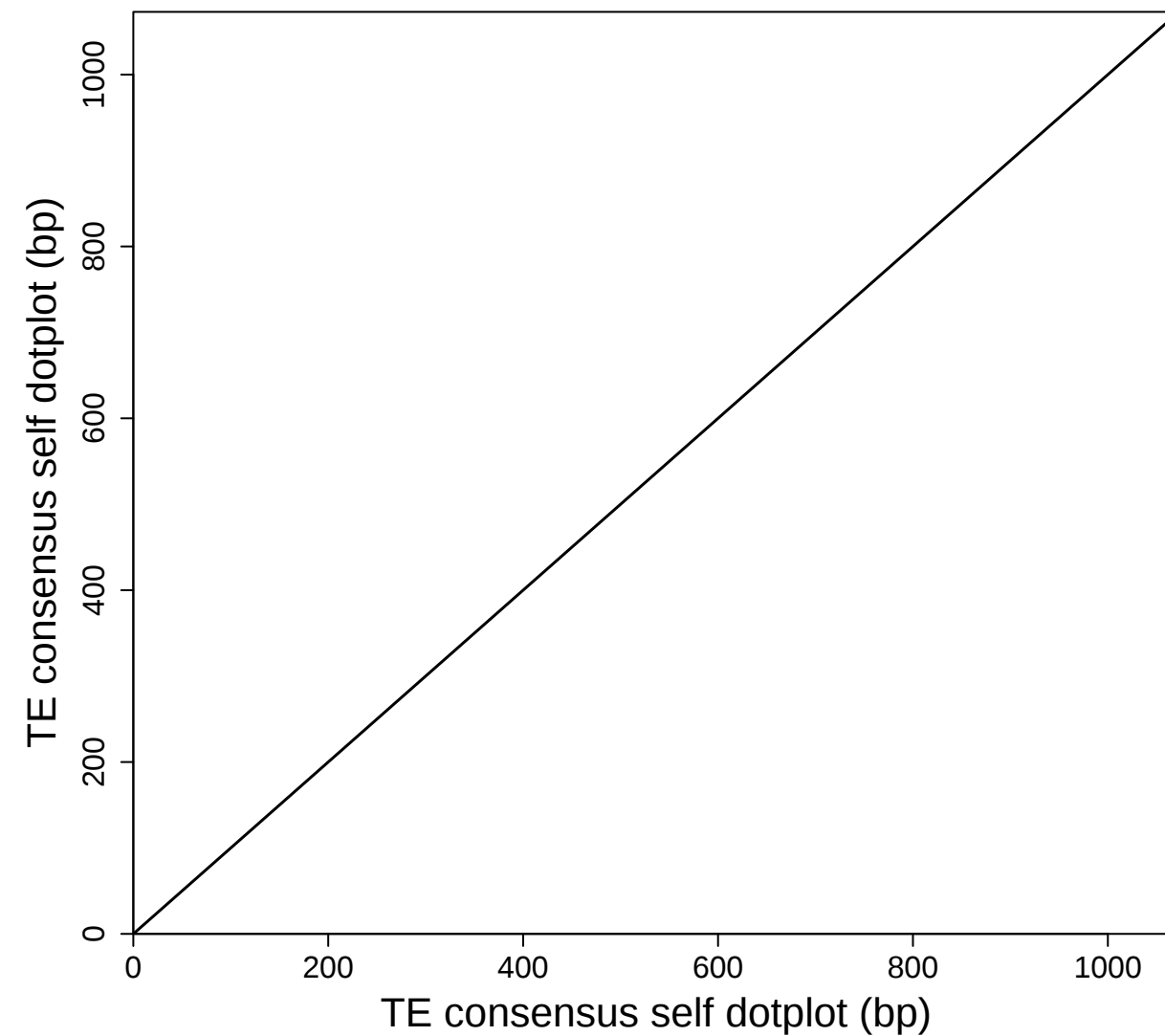
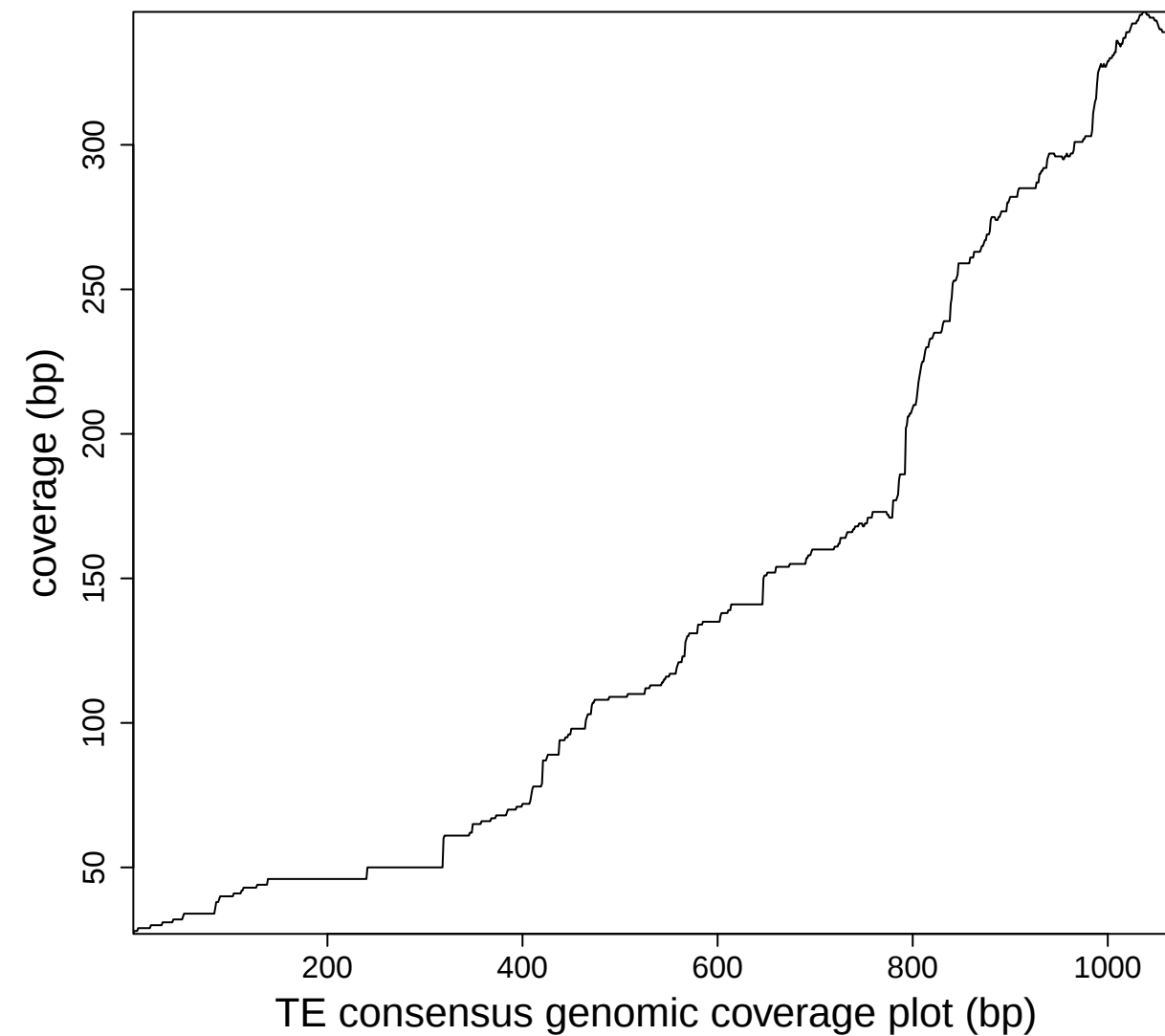
After TEtrimmer Extended plot Blue lines are boundaries



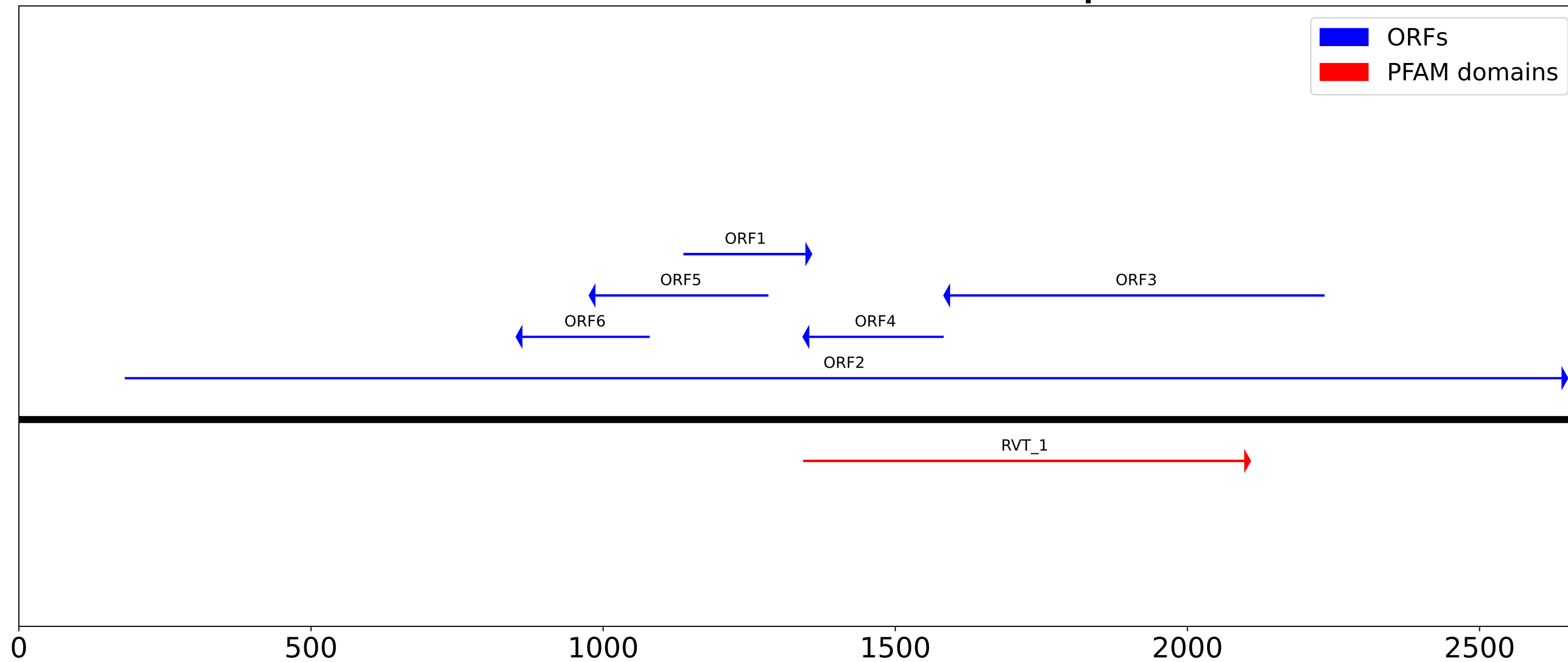
TE: rnd_3_family_220
size: 1073bp; fragments: 361; full length: 40 (≥ 965.7 bp)



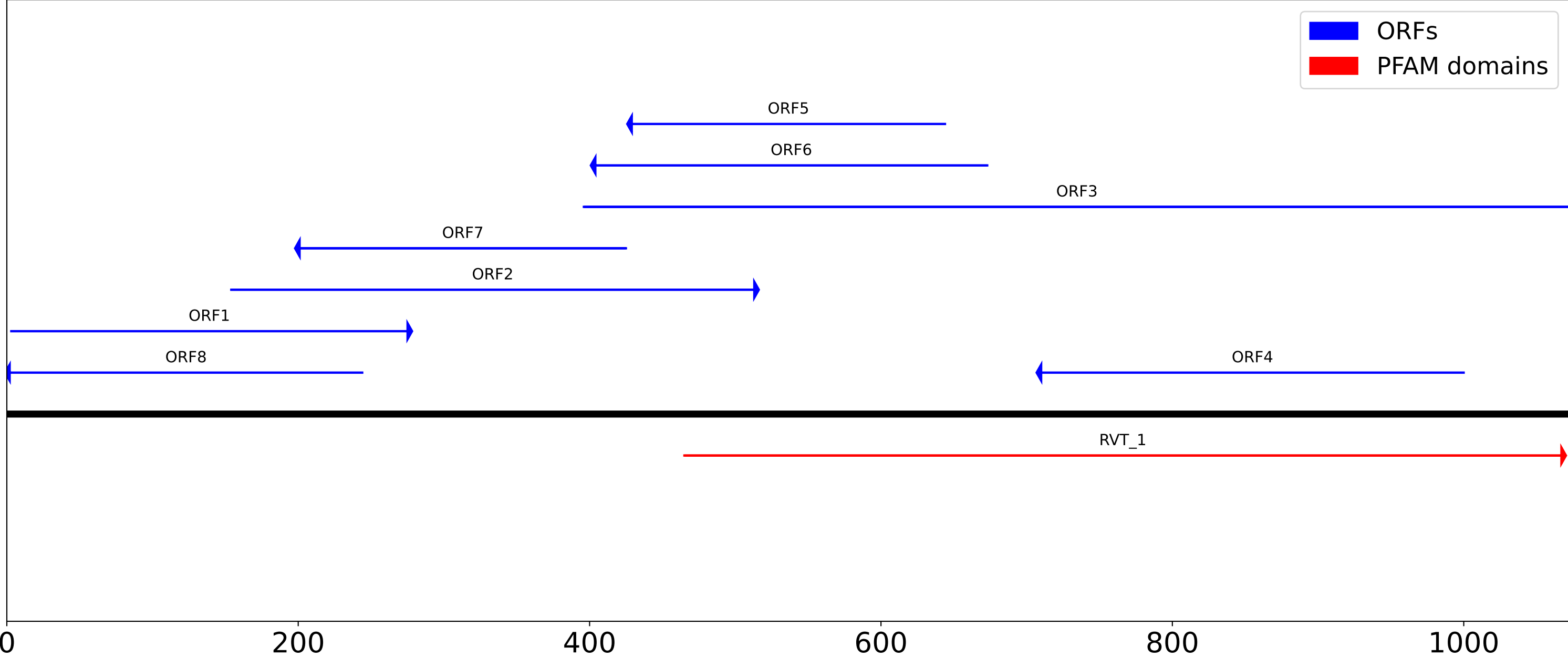
Before TEtrimmer 1073 bp



After TEtrimmer ORF and PFAM domain plot



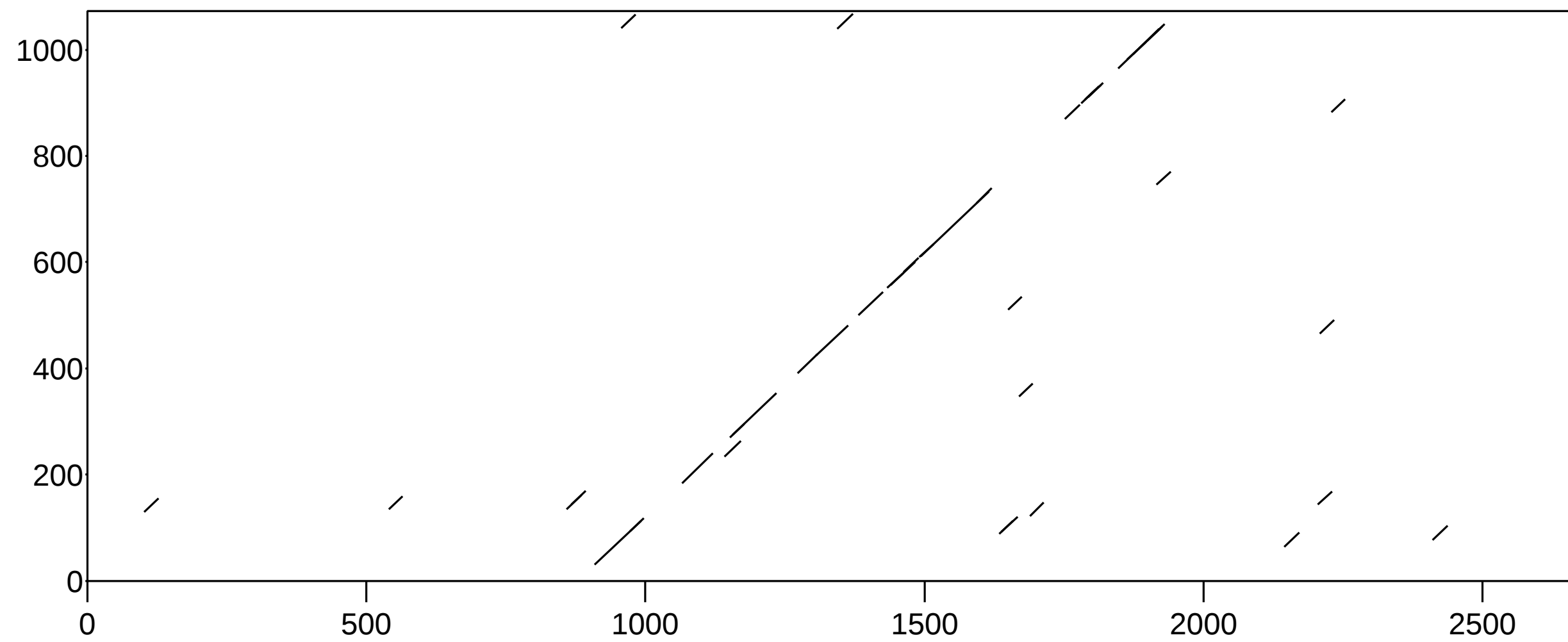
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)