

Start crop Point



End crop Point

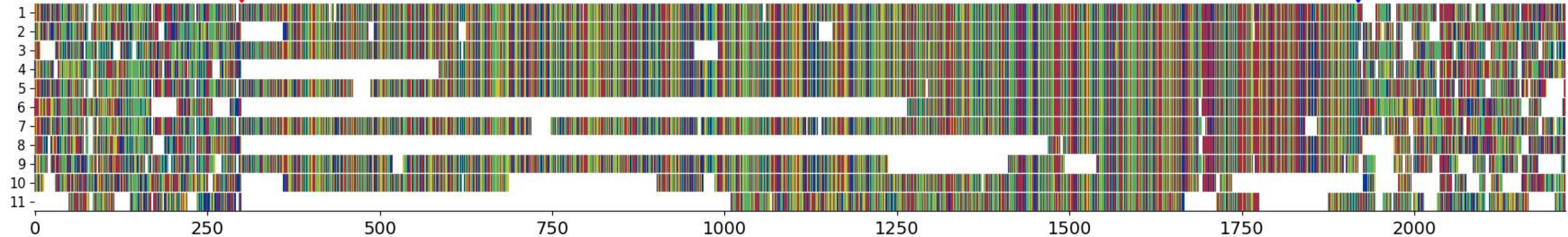


MSA length = 1620



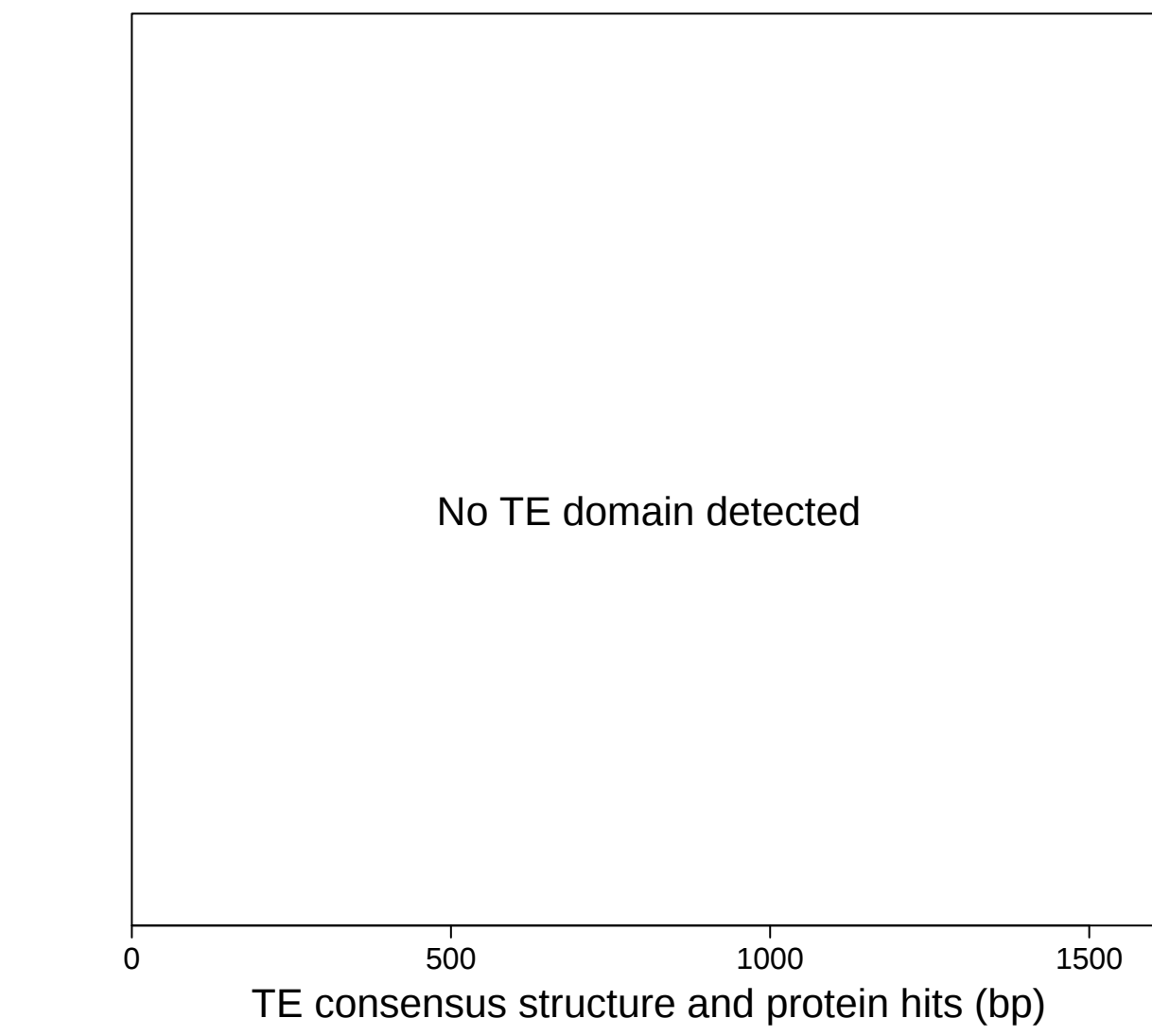
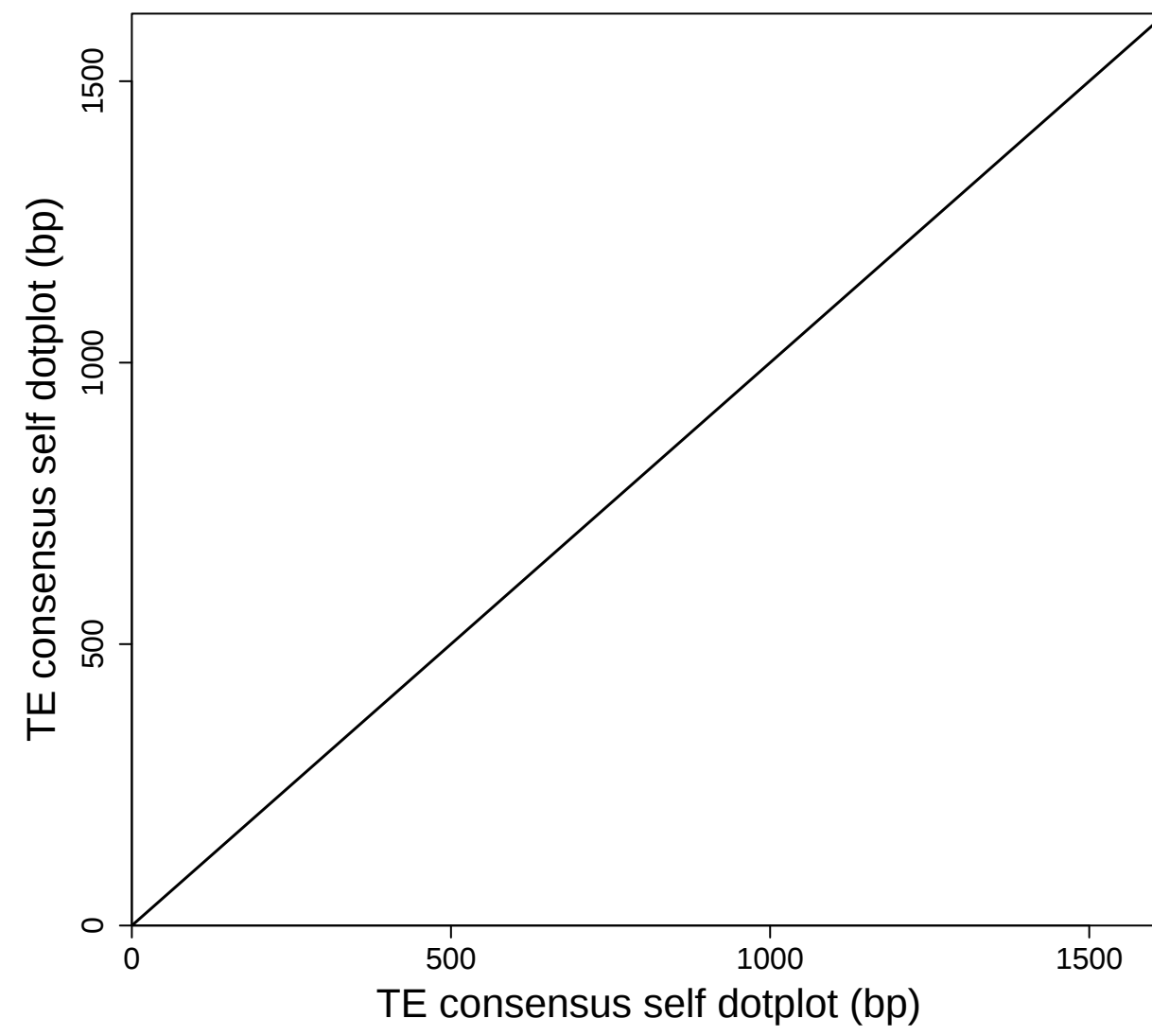
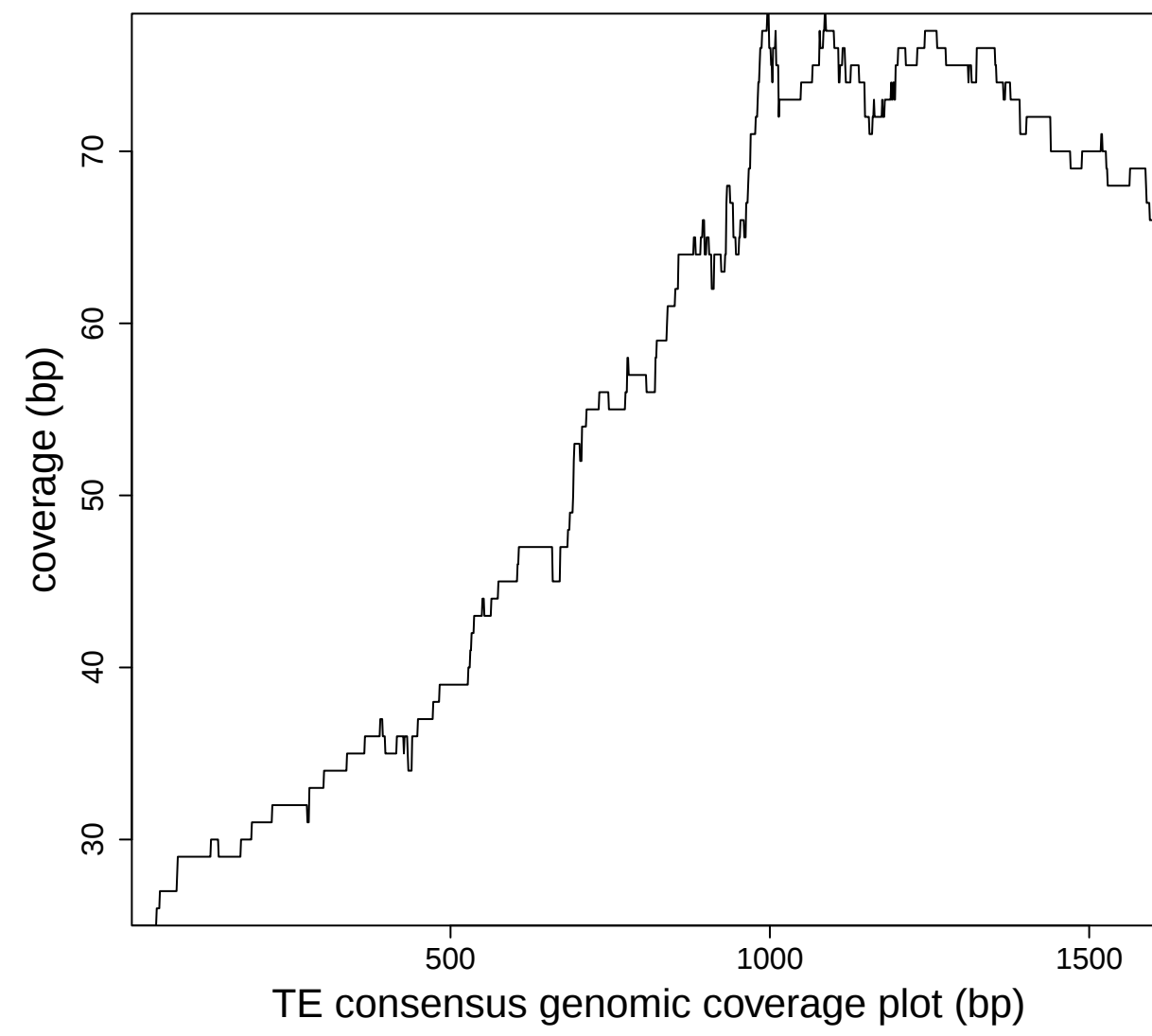
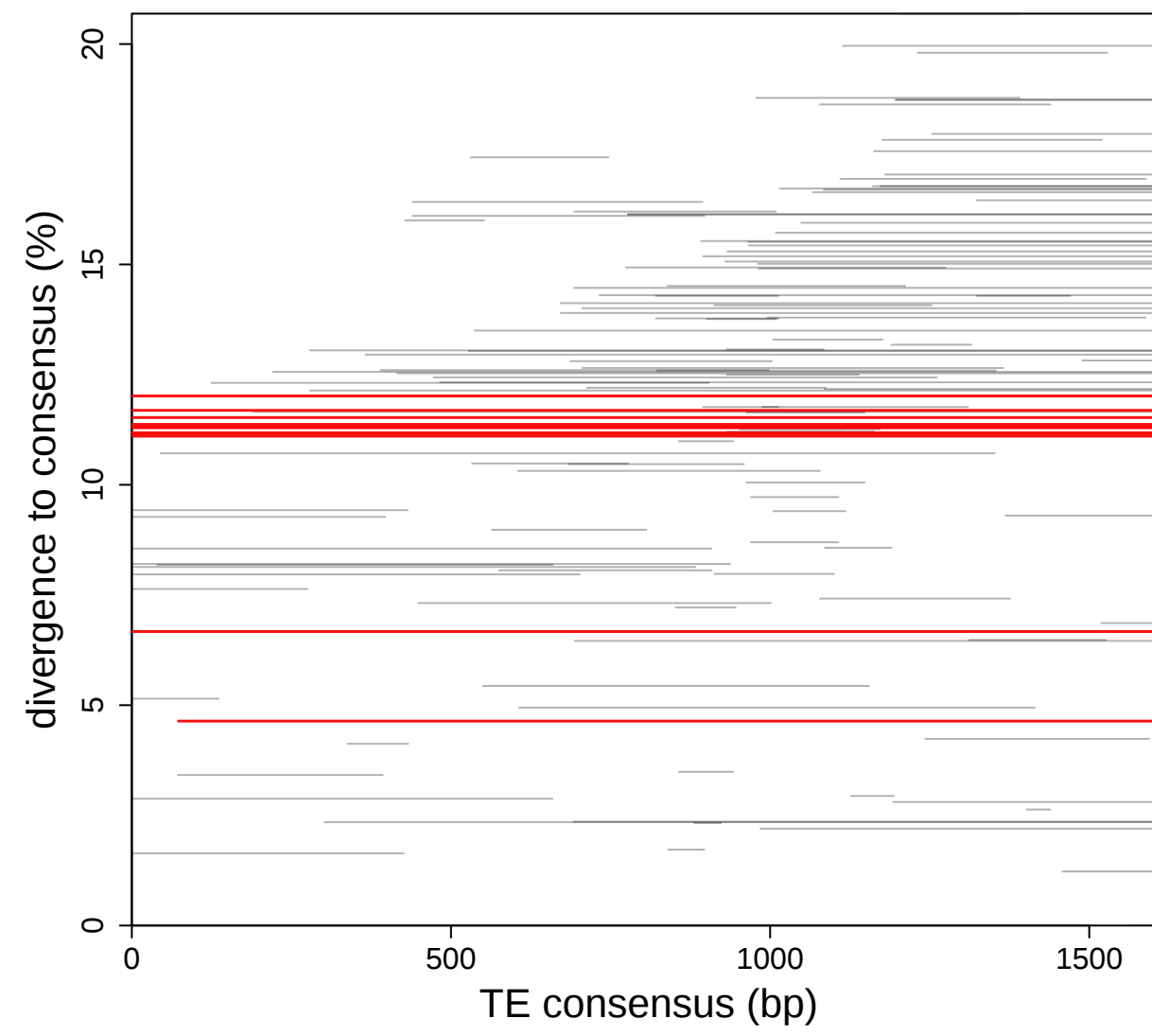
Start crop Point

End crop Point



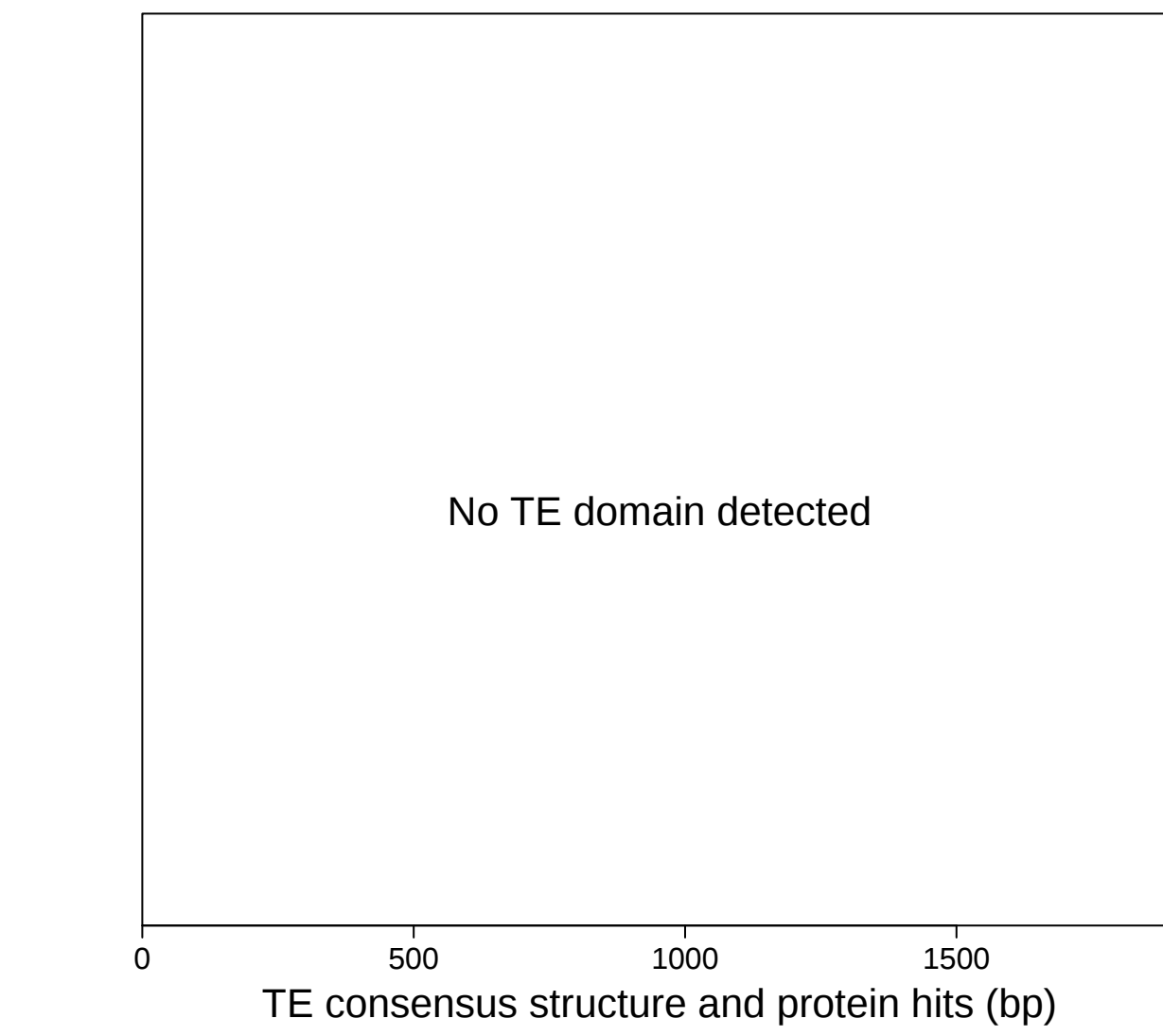
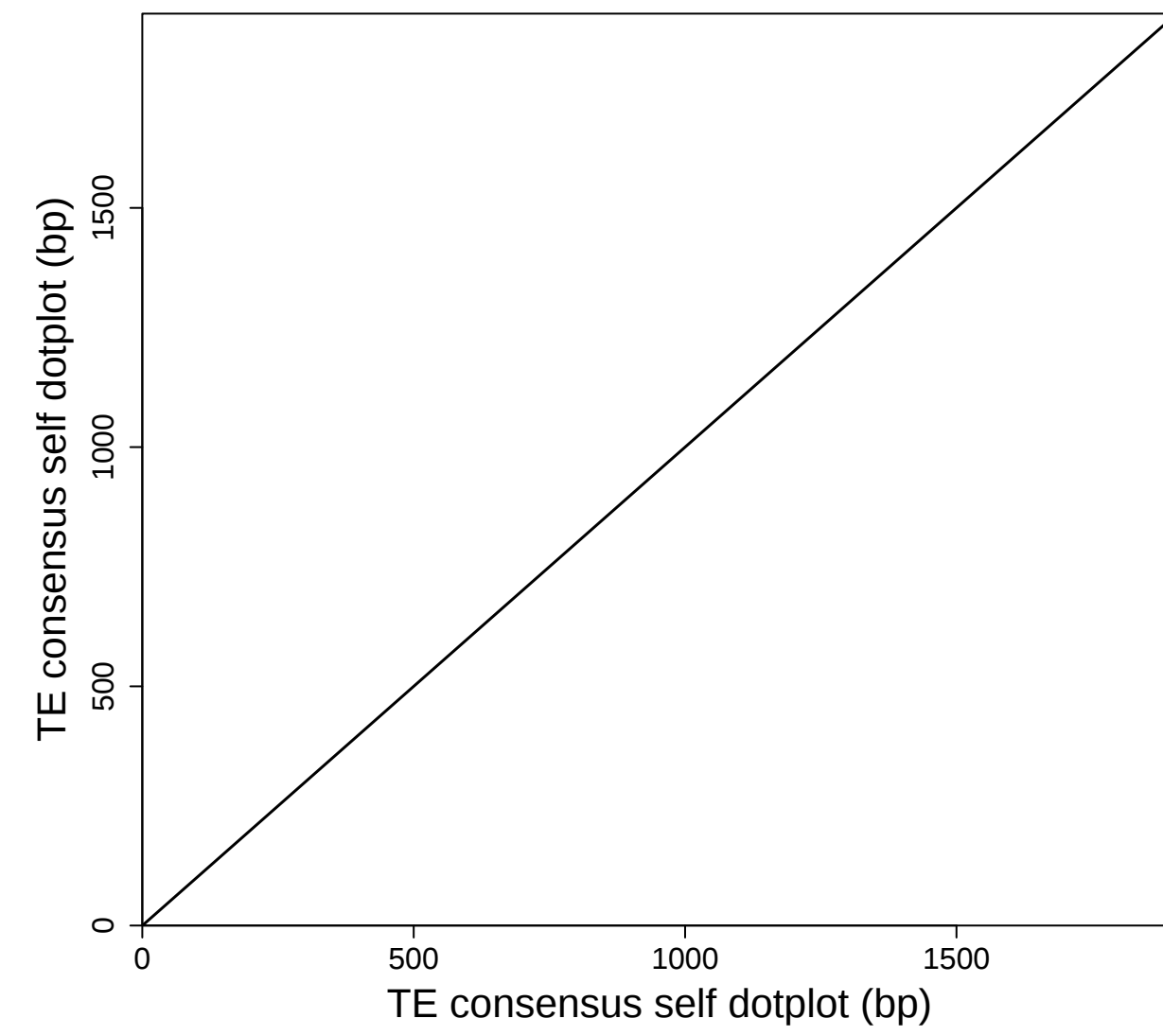
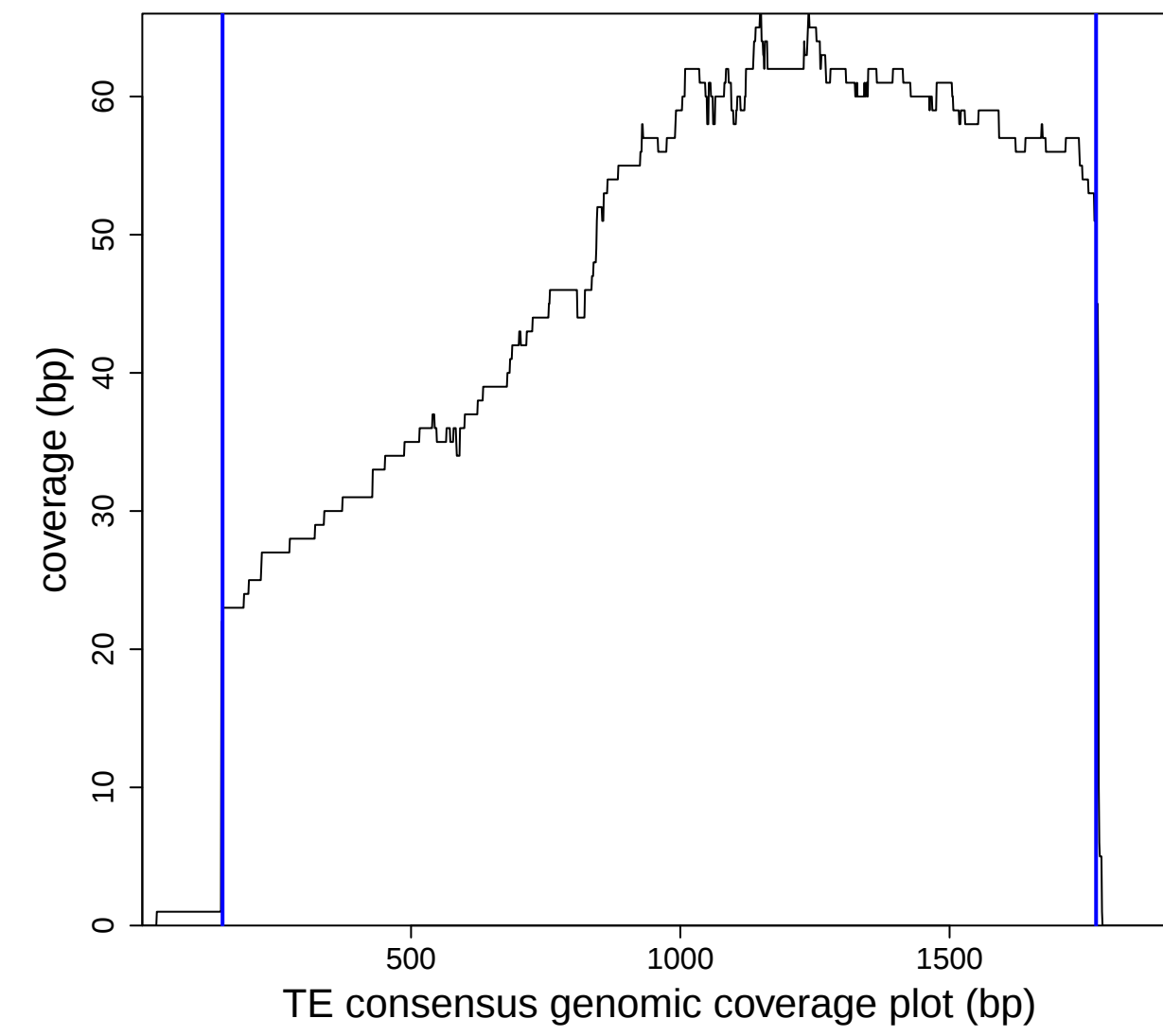
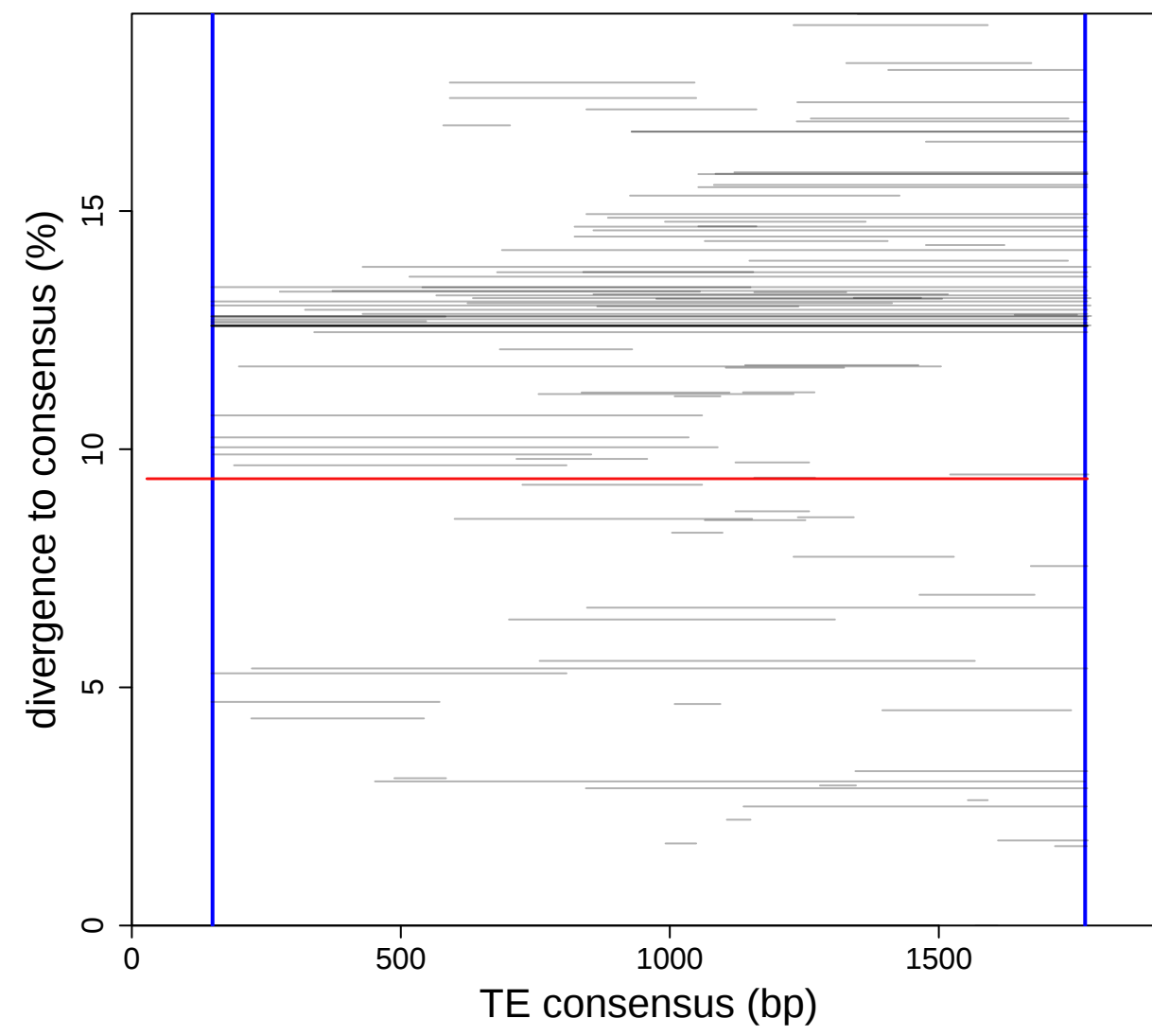
bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.fa_gs.fa_c
size: 1620bp; fragments: 142; full length: 16 (≥ 1458 bp)

After TEtrimmer 1620 bp

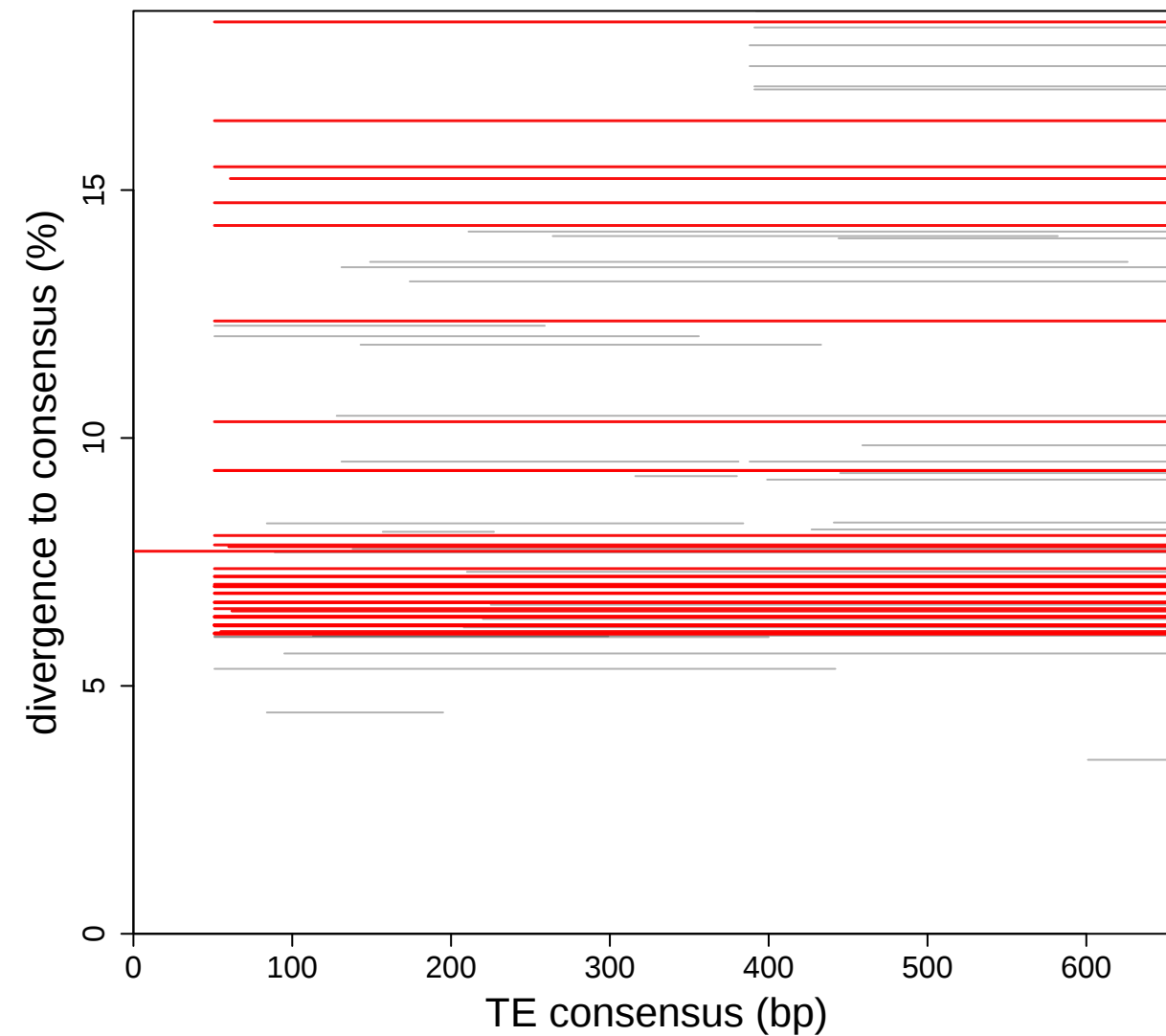


92.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.1
size: 1922bp; fragments: 115; full length: 1 (>=1729.8bp)

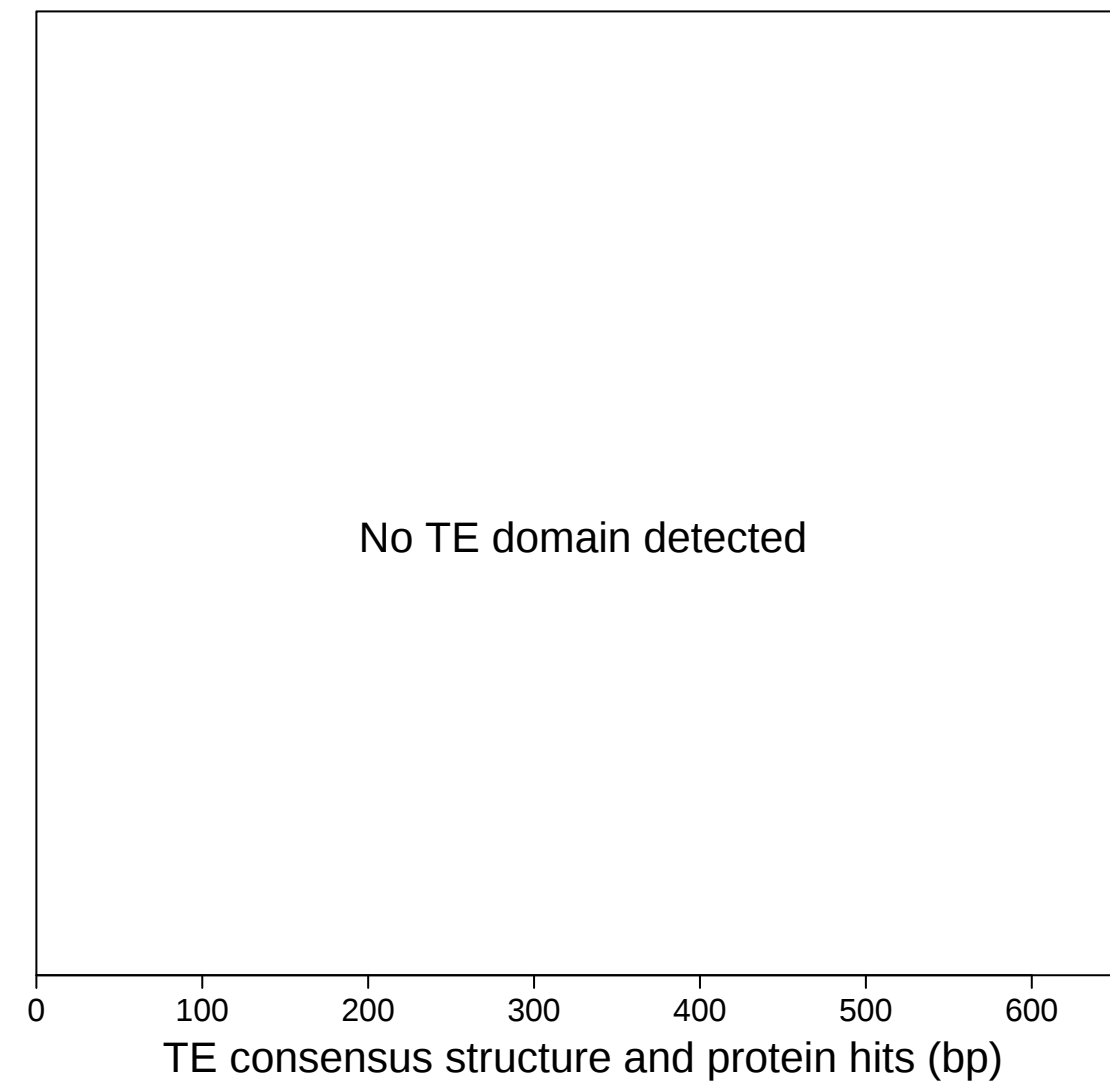
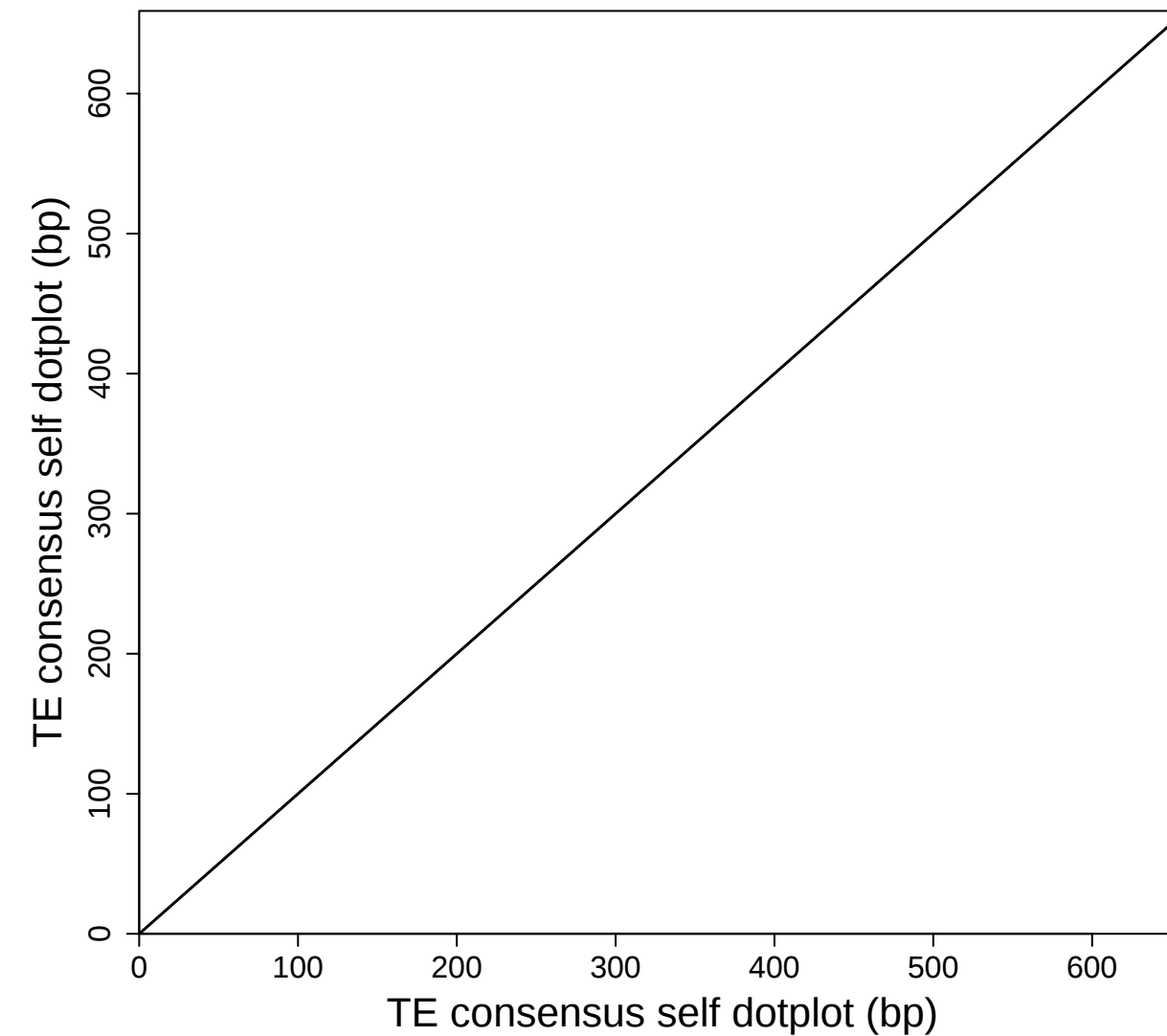
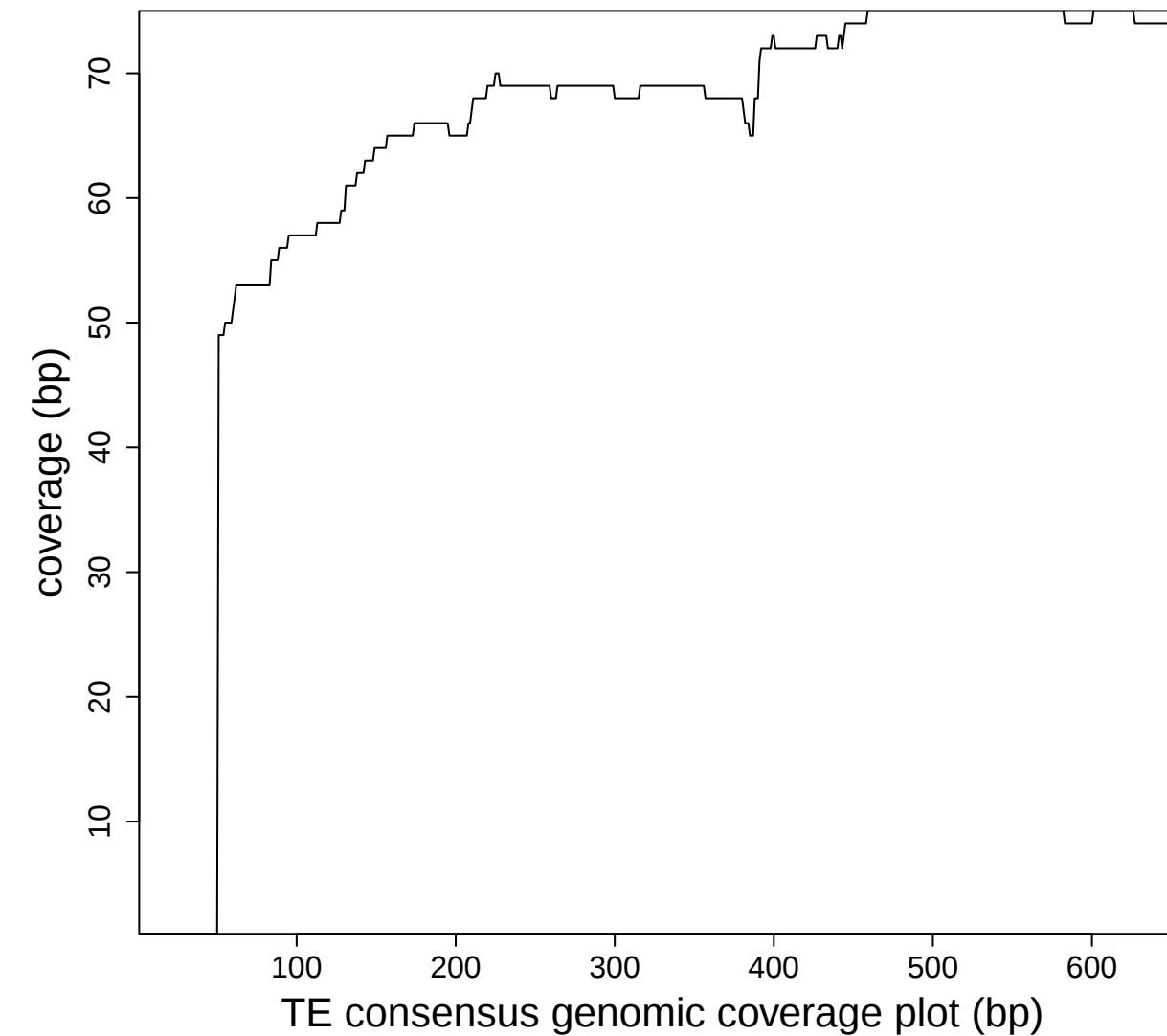
After TEtrimmer Extended plot Blue lines are boundaries



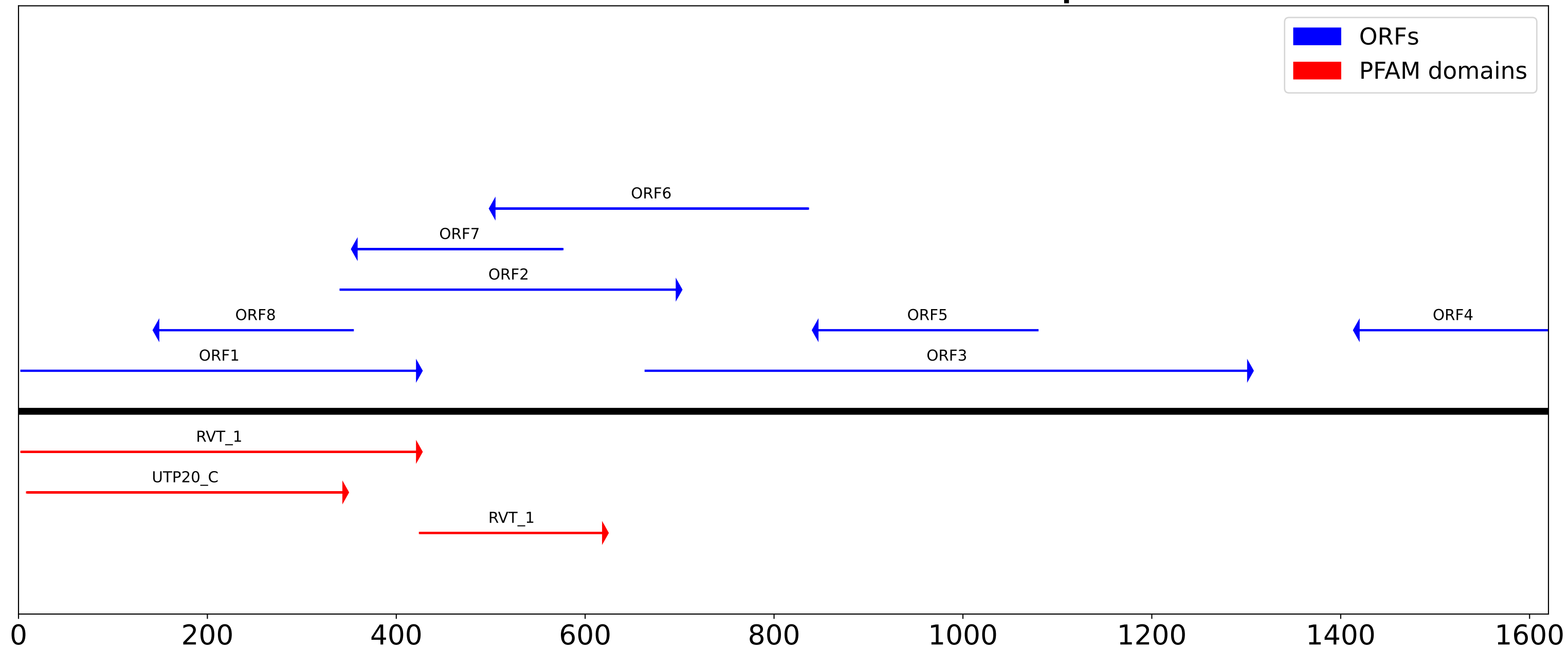
TE: rnd_5_family_14992
size: 659bp; fragments: 87; full length: 48 (≥ 593.1 bp)



Before TEtrimmer 659 bp



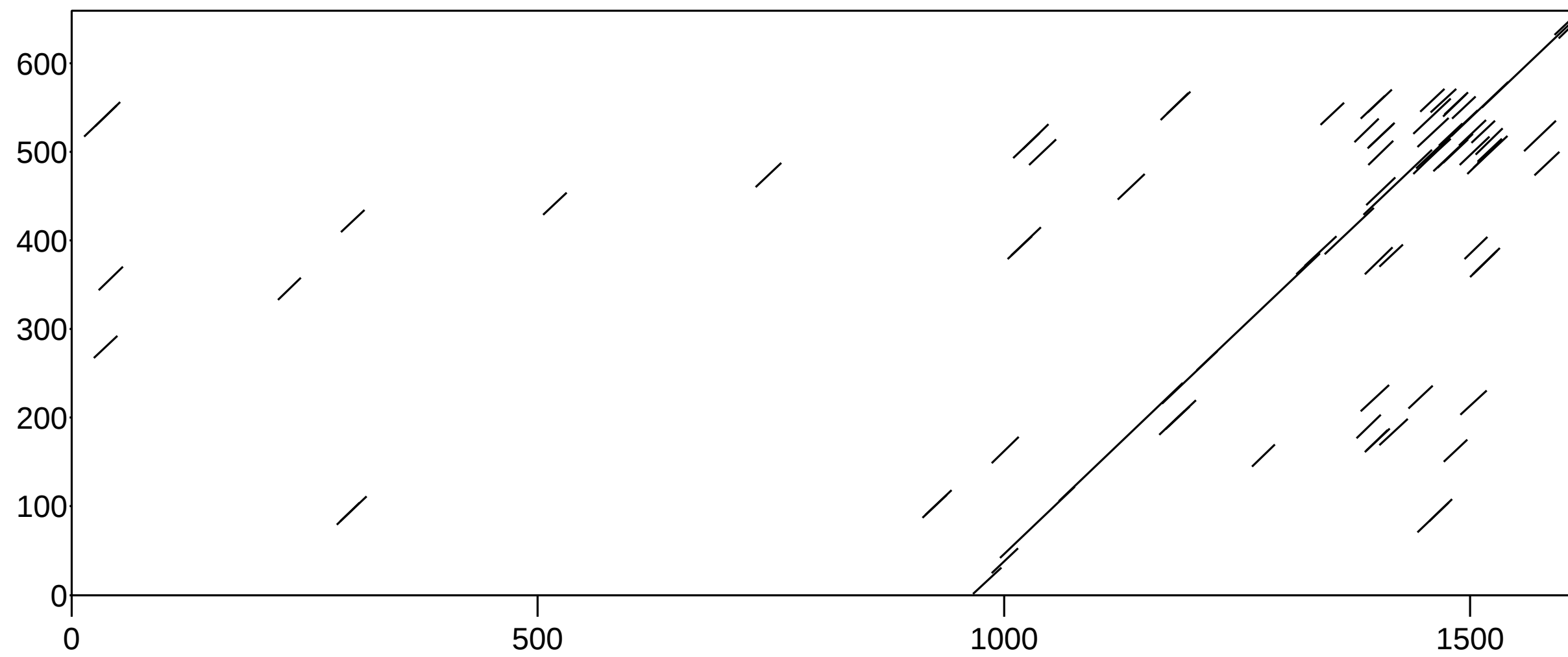
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)