

Start crop Point



End crop Point



MSA length = 1763



Start crop Point

End crop Point

0

10

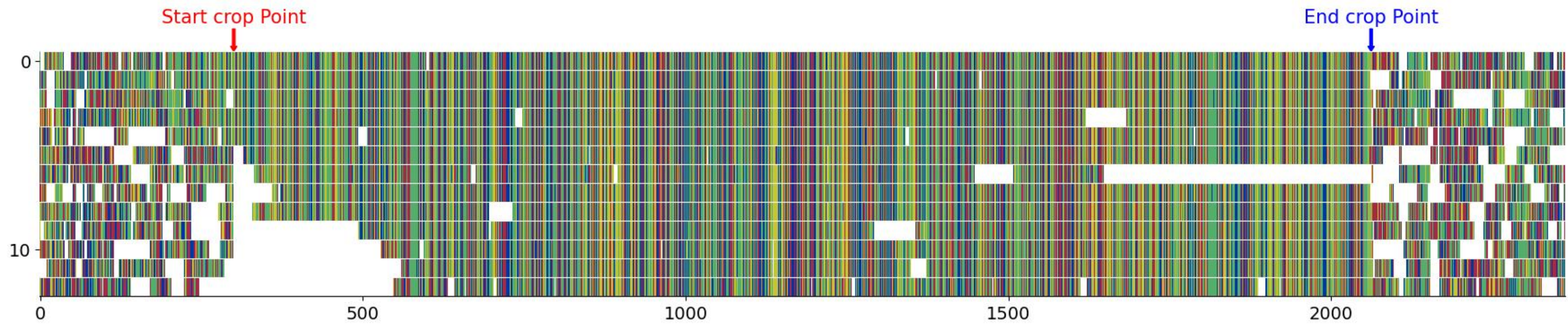
0

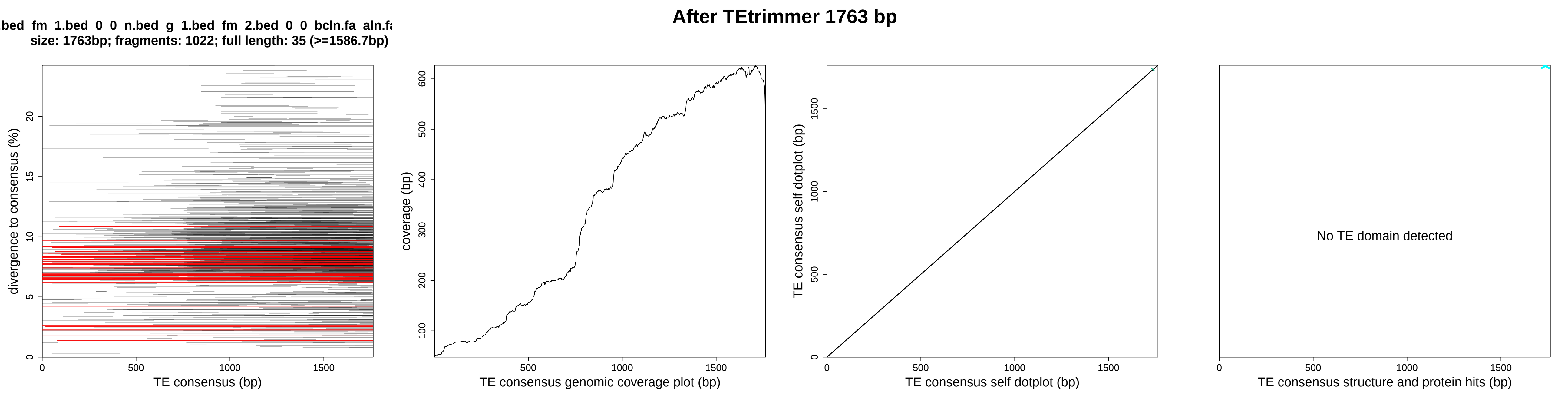
500

1000

1500

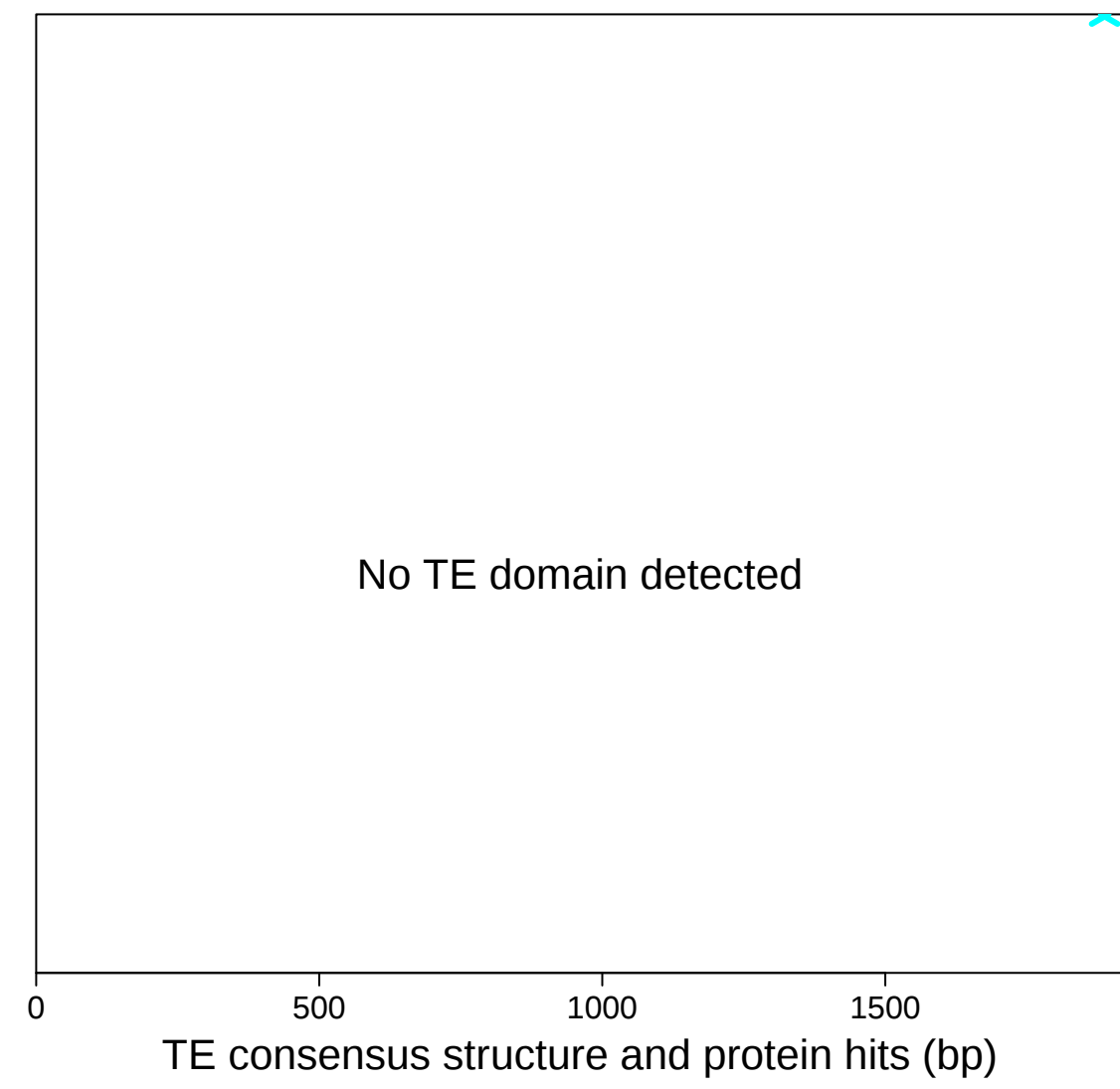
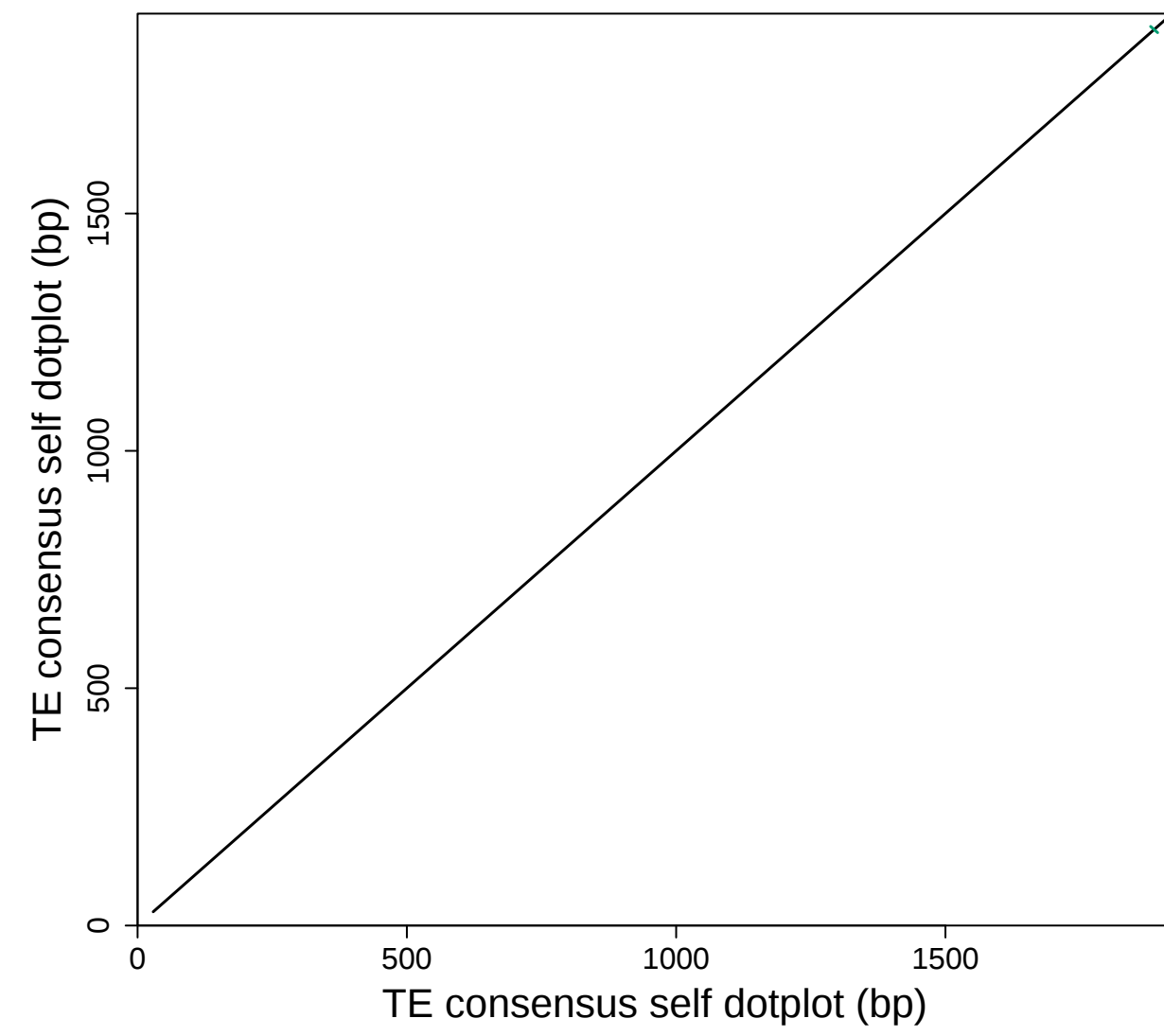
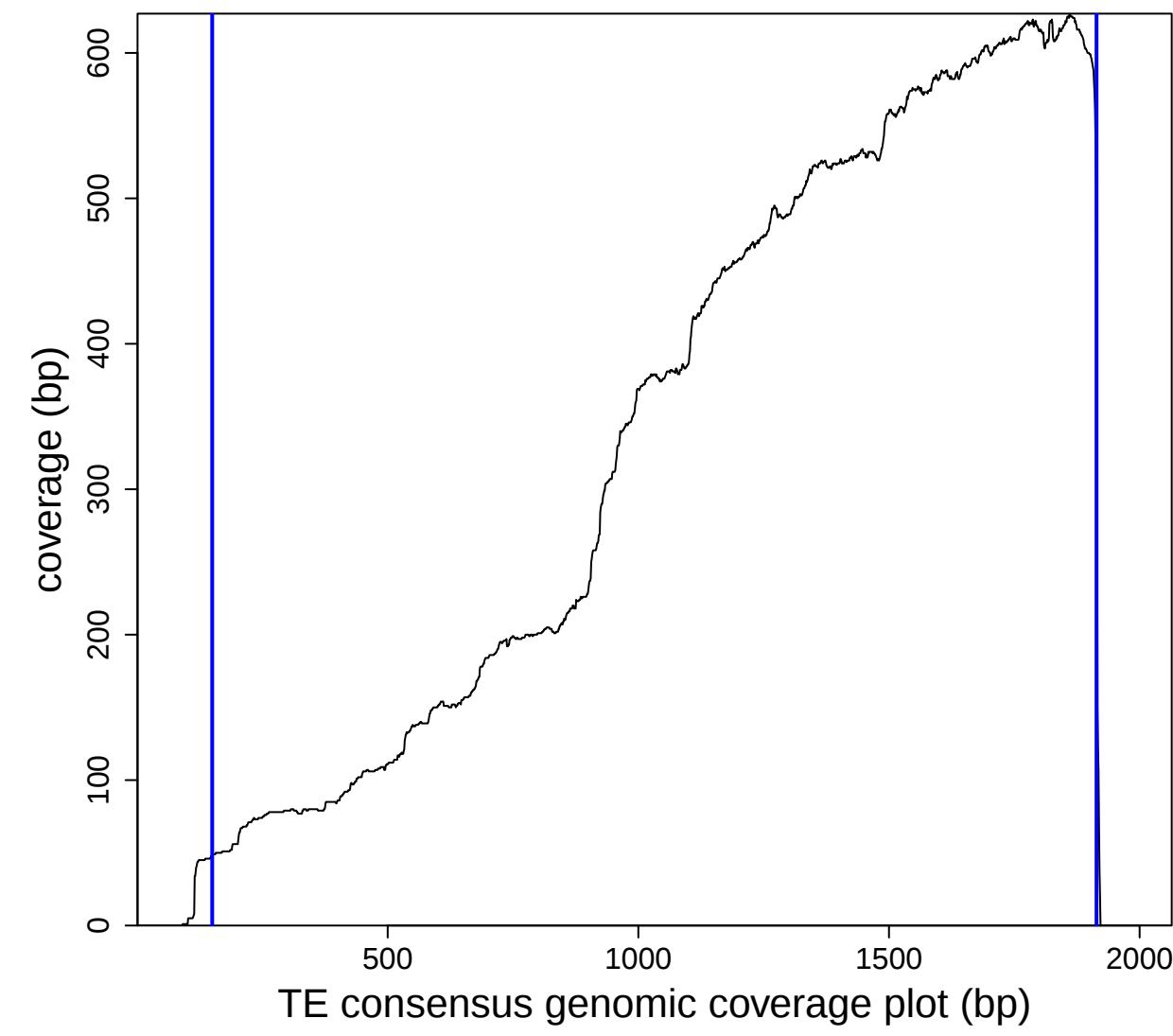
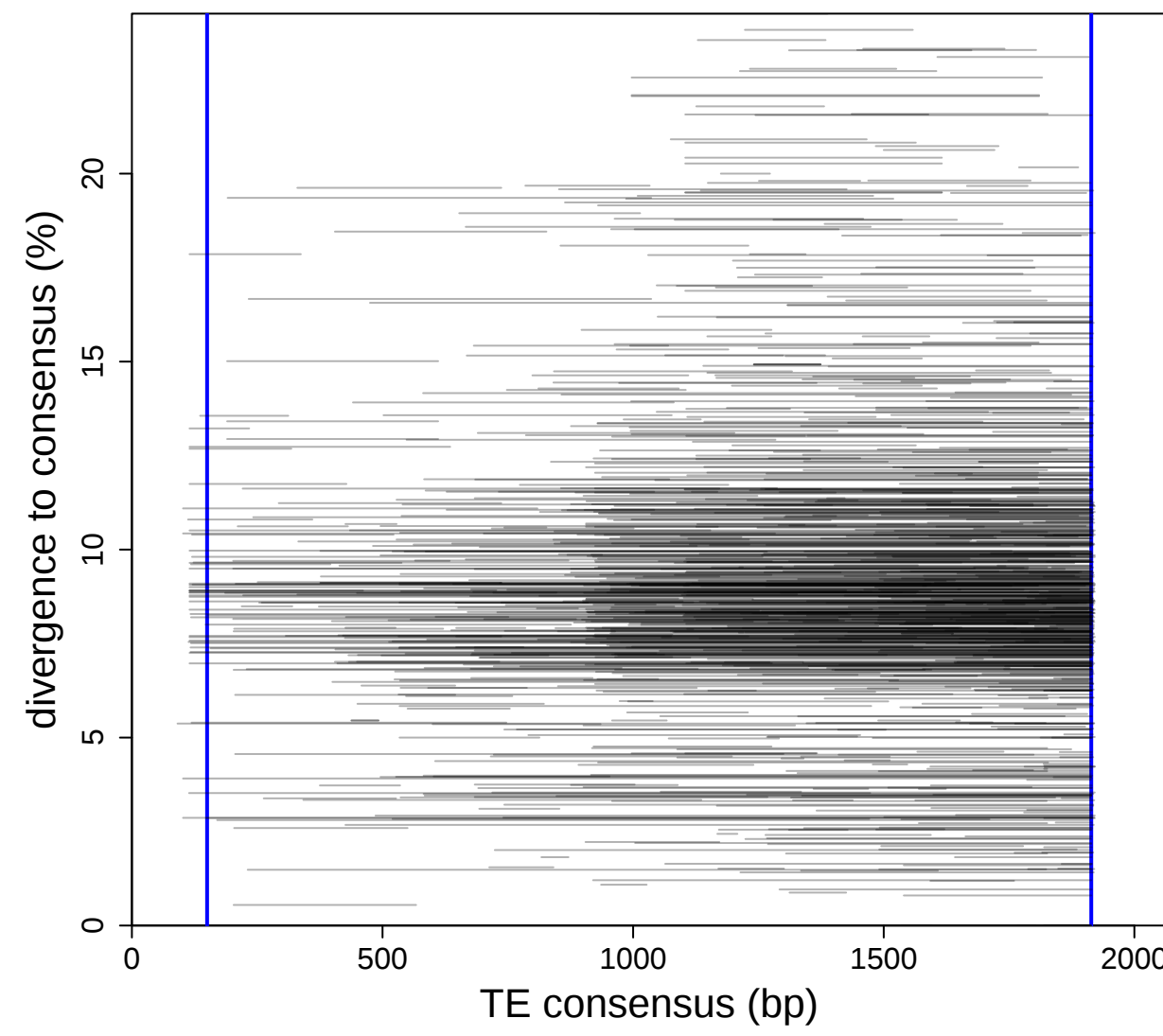
2000



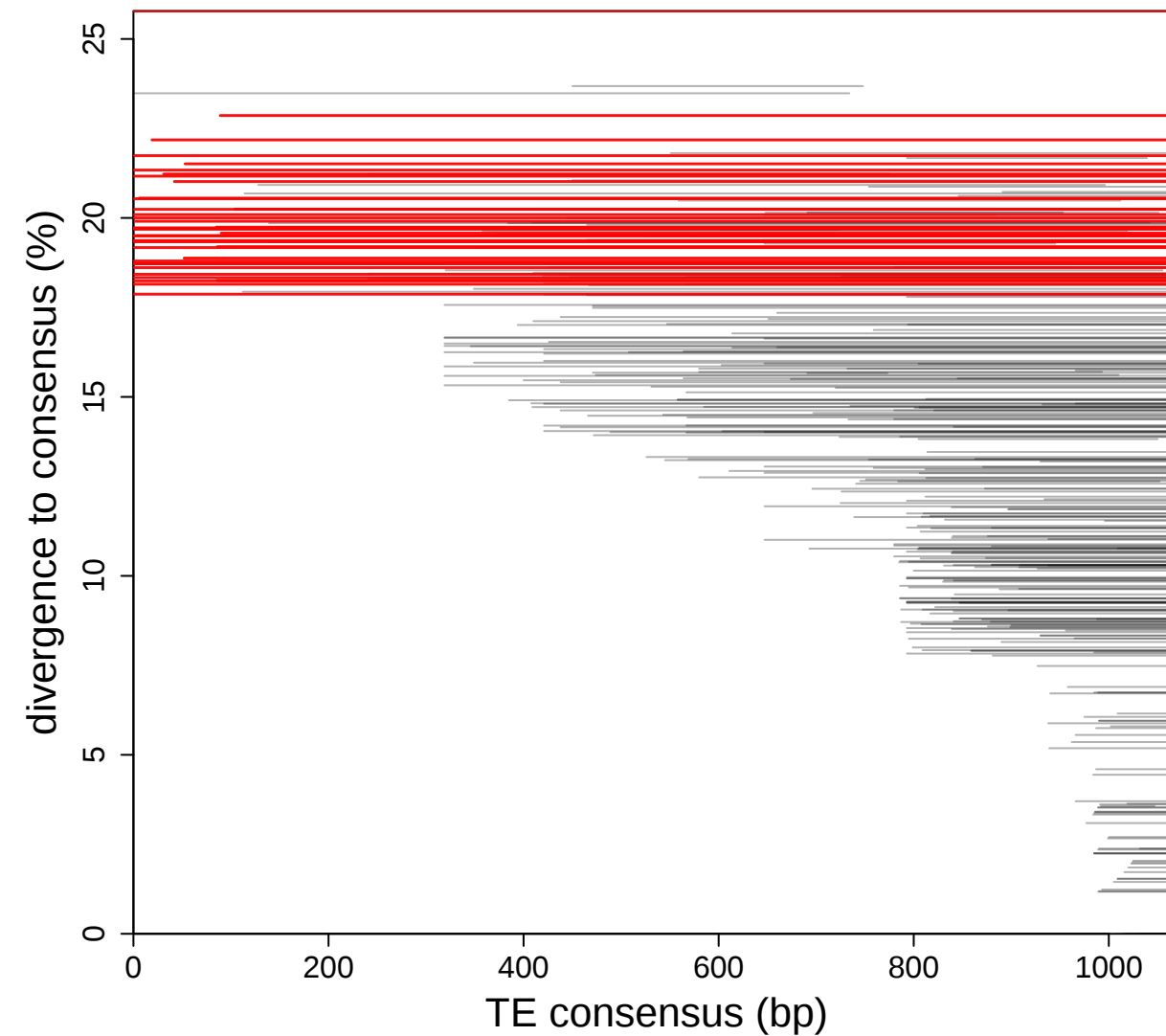


f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 2064bp; fragments: 1021; full length: 0 (>=1857.6bp)

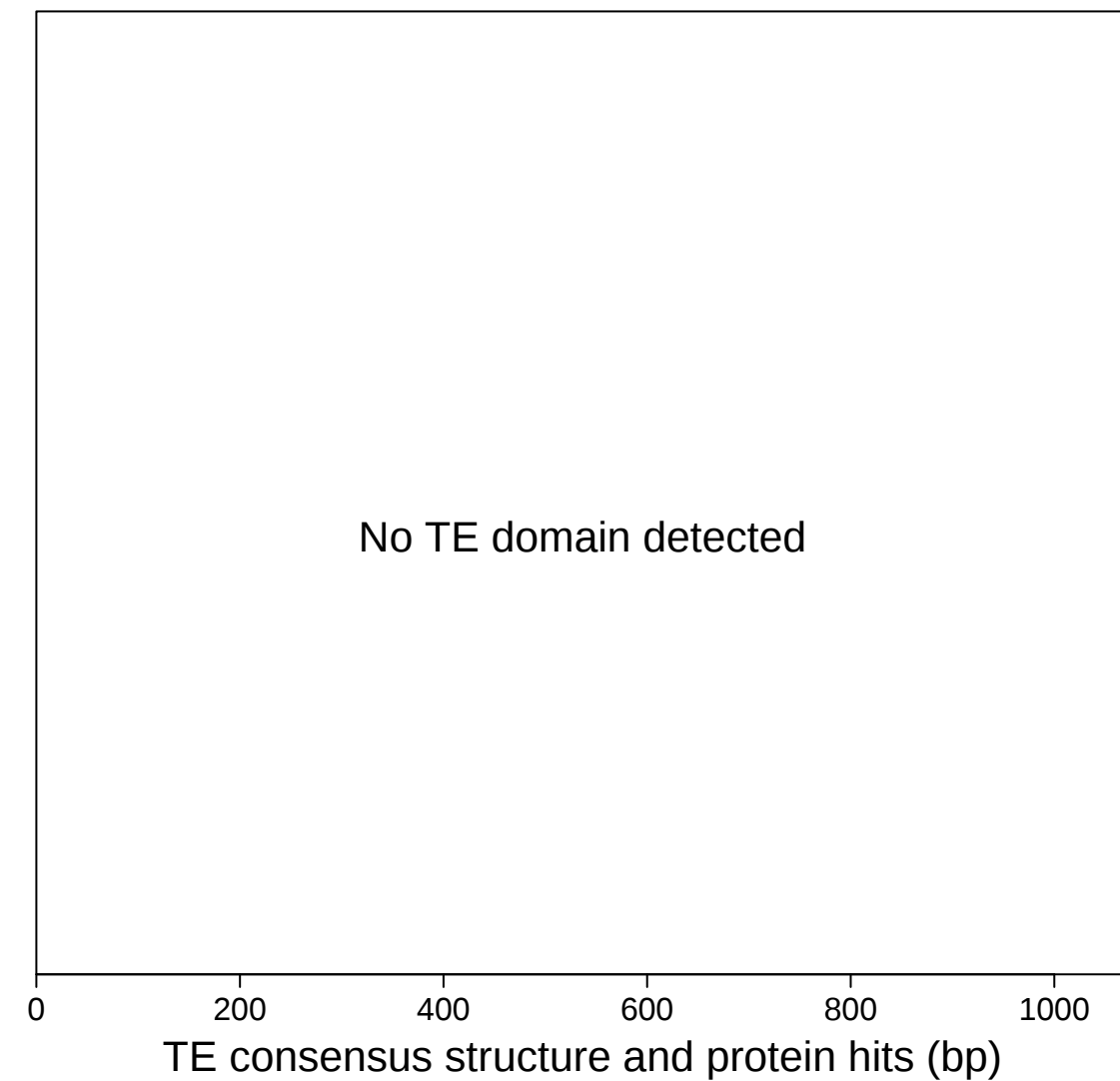
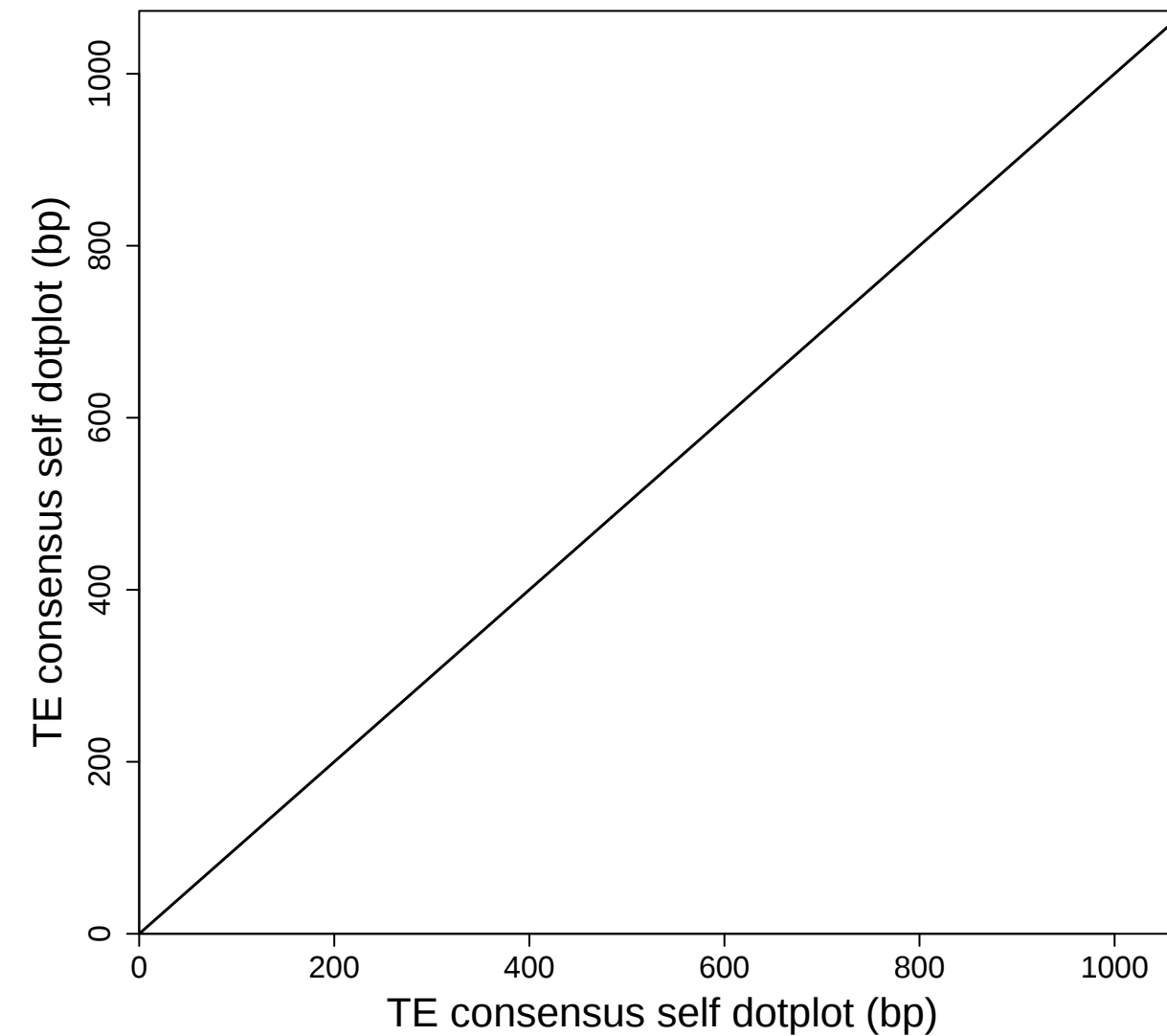
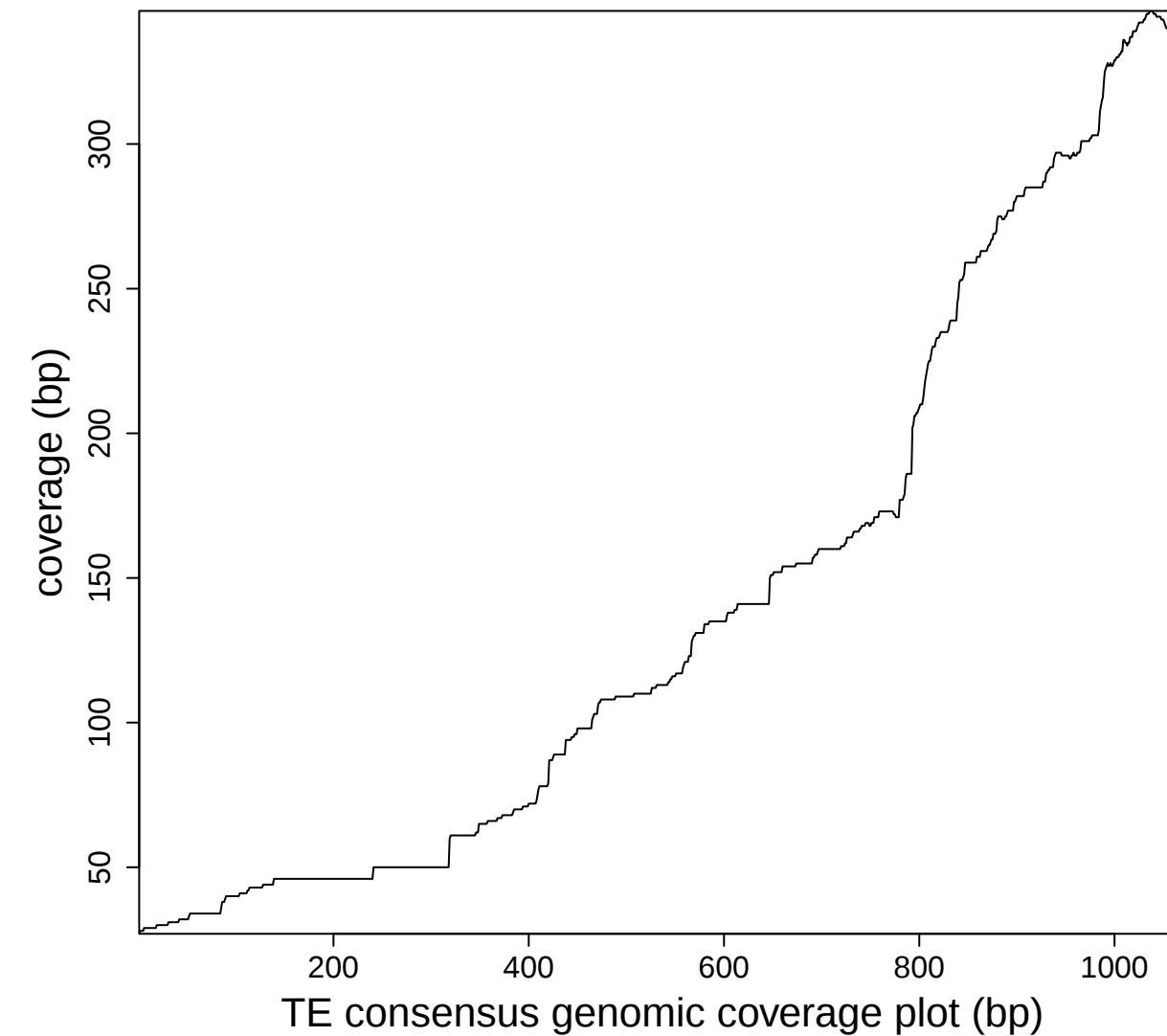
After TEtrimmer Extended plot Blue lines are boundaries



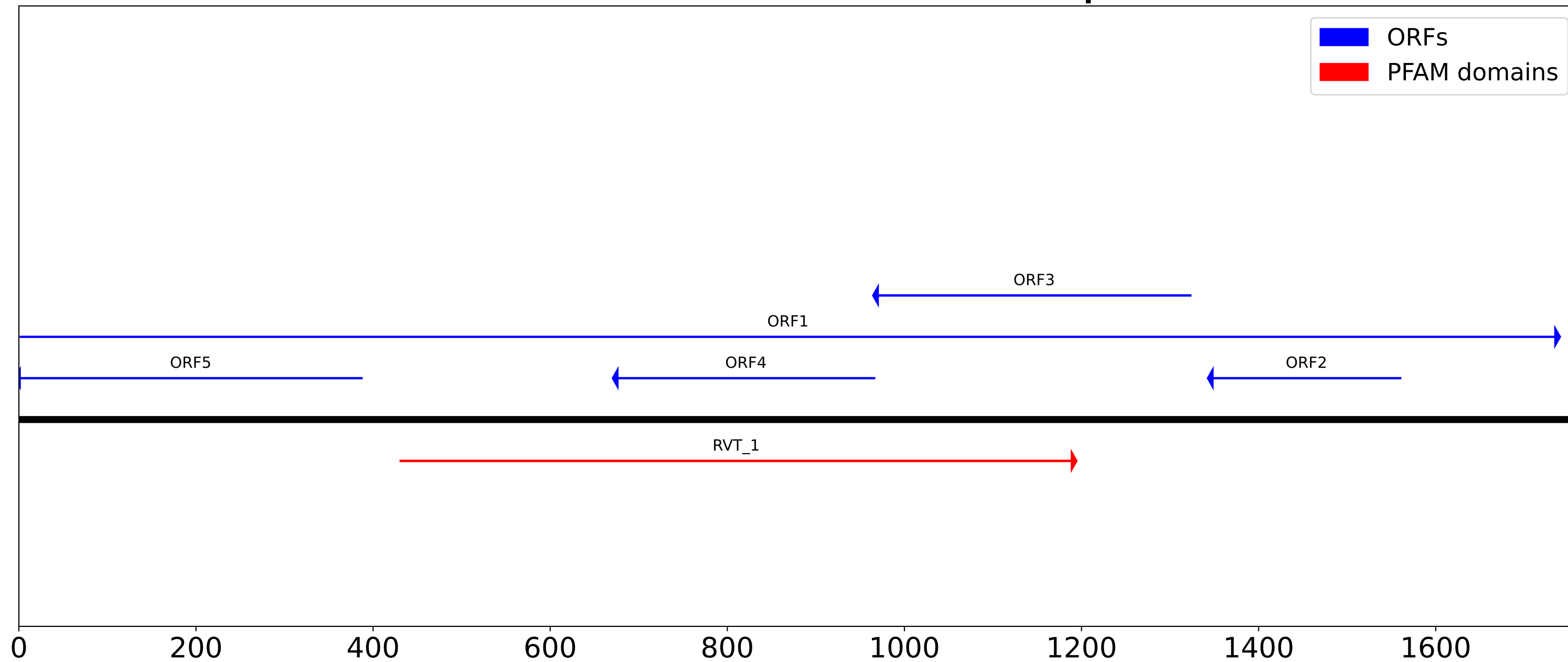
TE: rnd_3_family_220
size: 1073bp; fragments: 361; full length: 40 (≥ 965.7 bp)



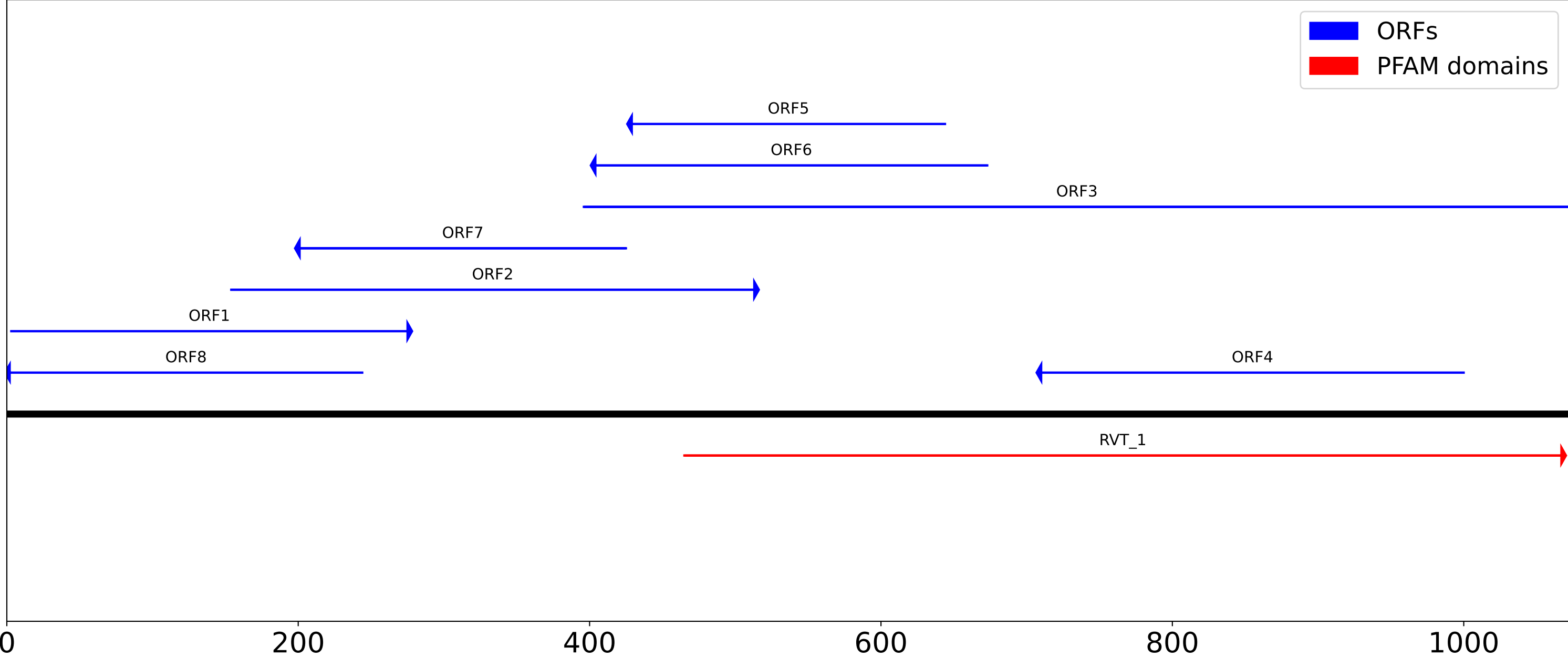
Before TEtrimmer 1073 bp



After TEtrimmer ORF and PFAM domain plot



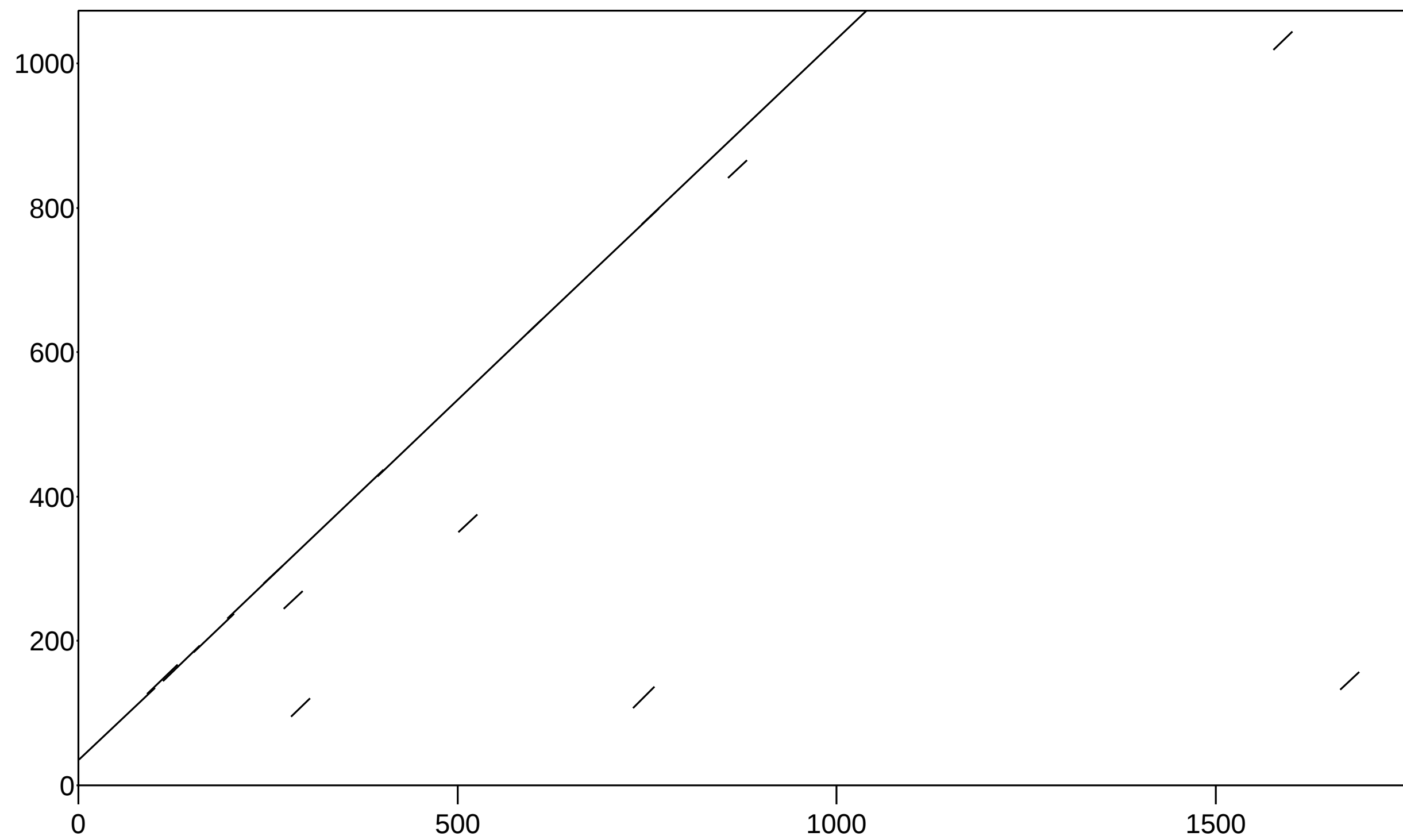
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)