

Start crop Point



End crop Point



MSA length = 4411



Start crop Point

End crop Point

0 -

10 -

0

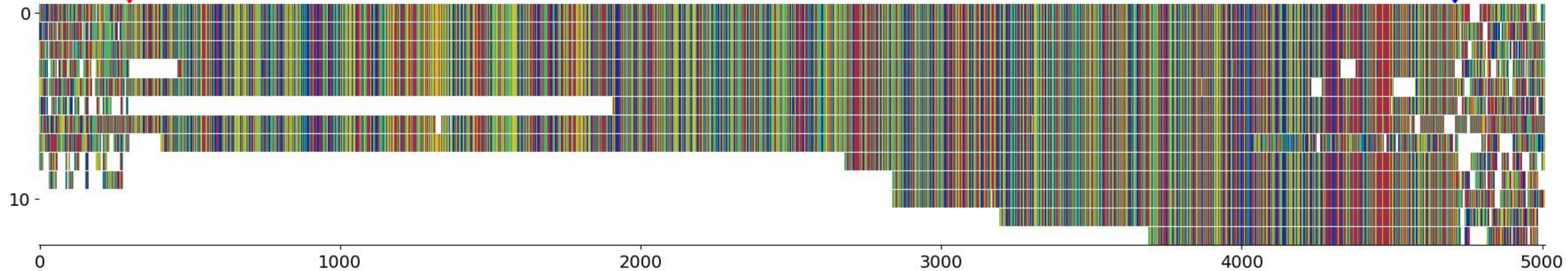
1000

2000

3000

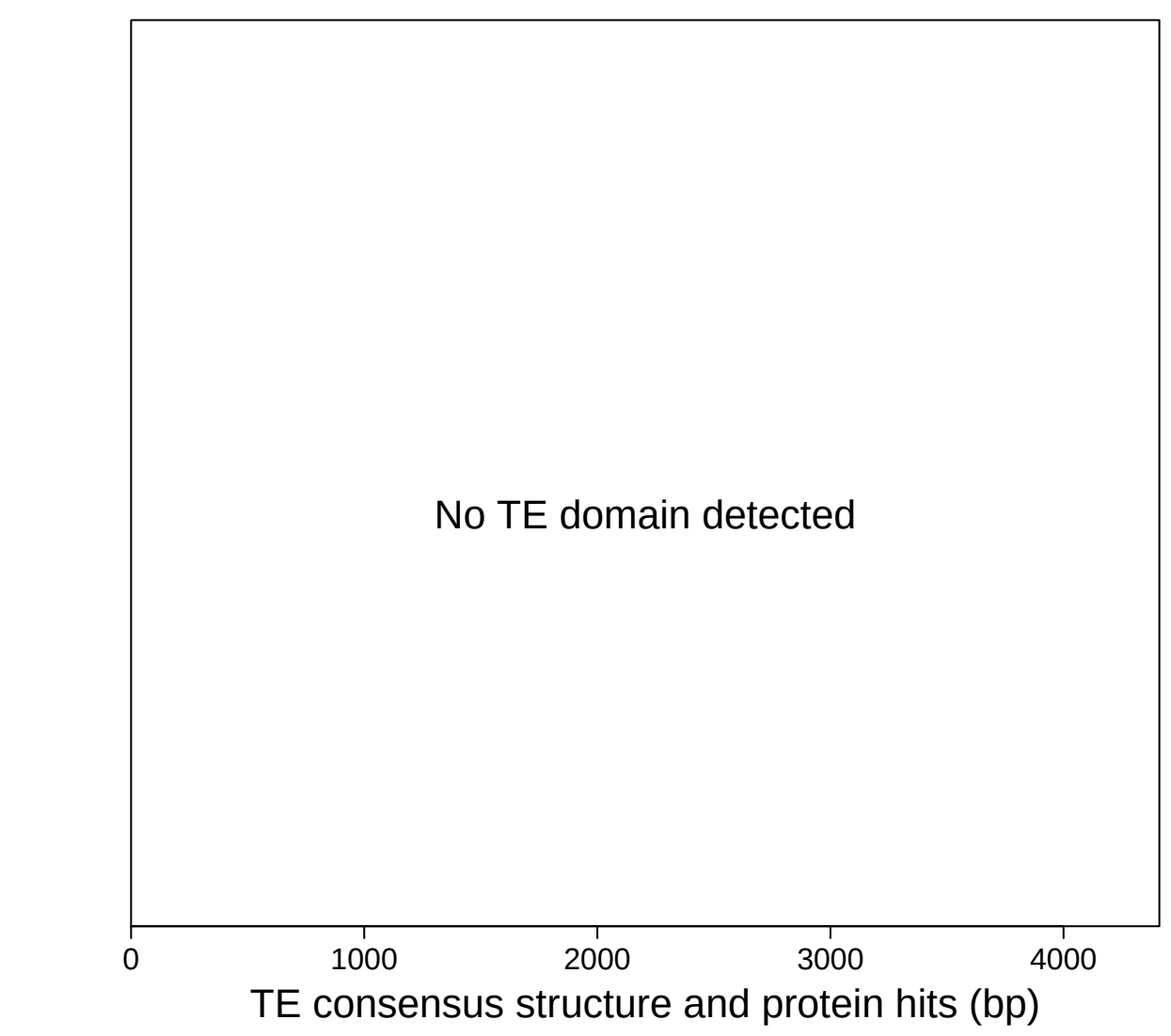
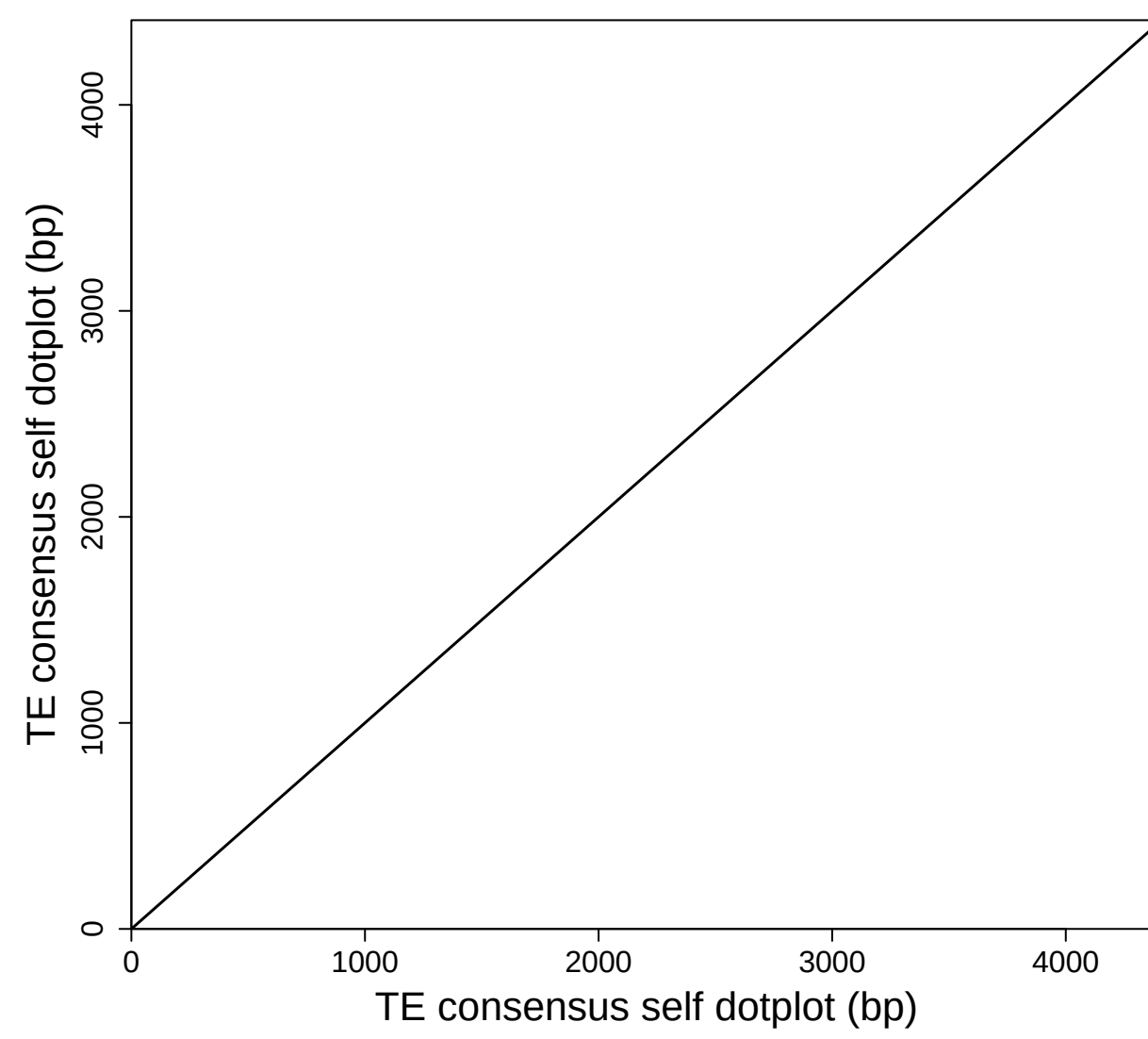
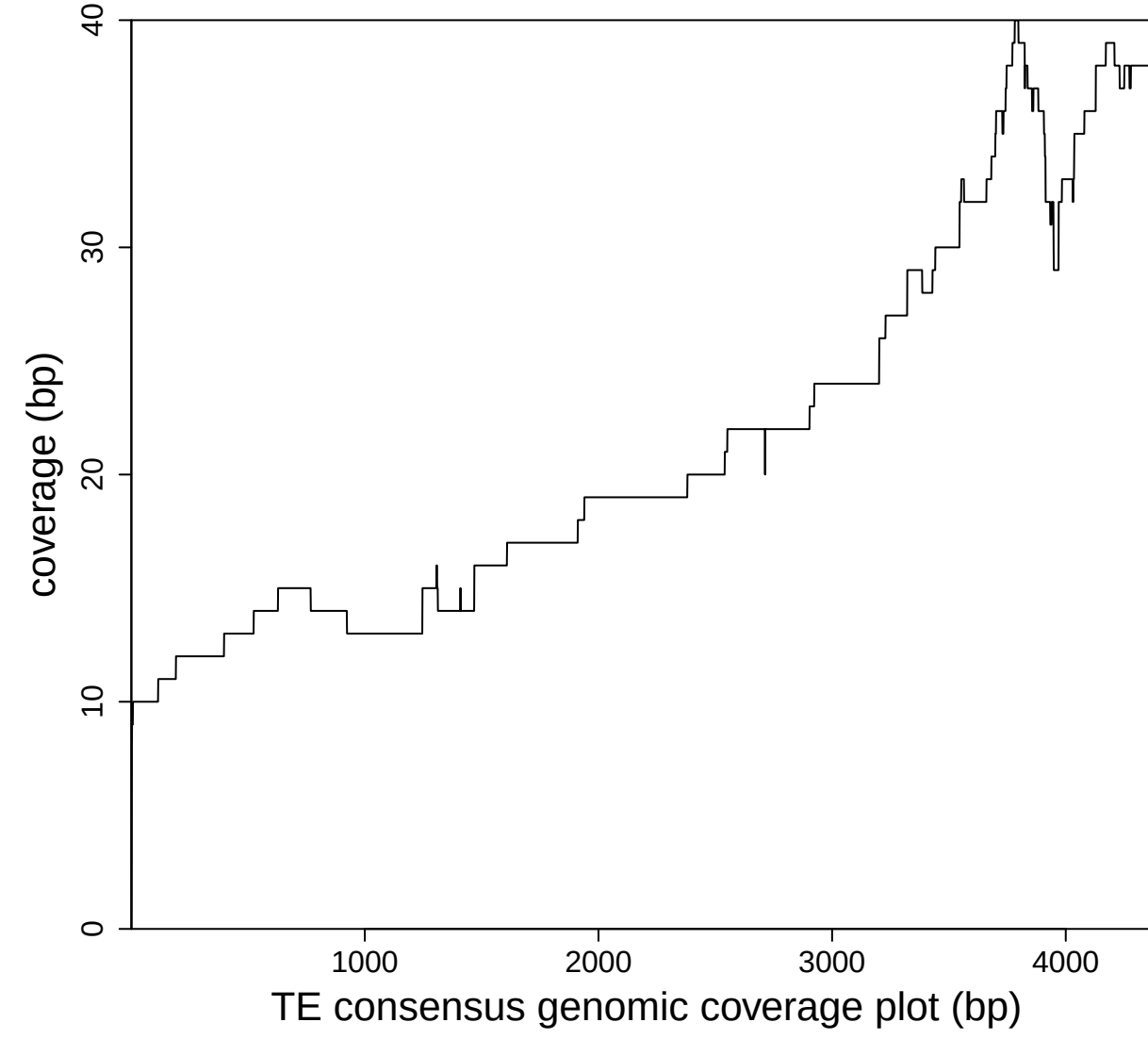
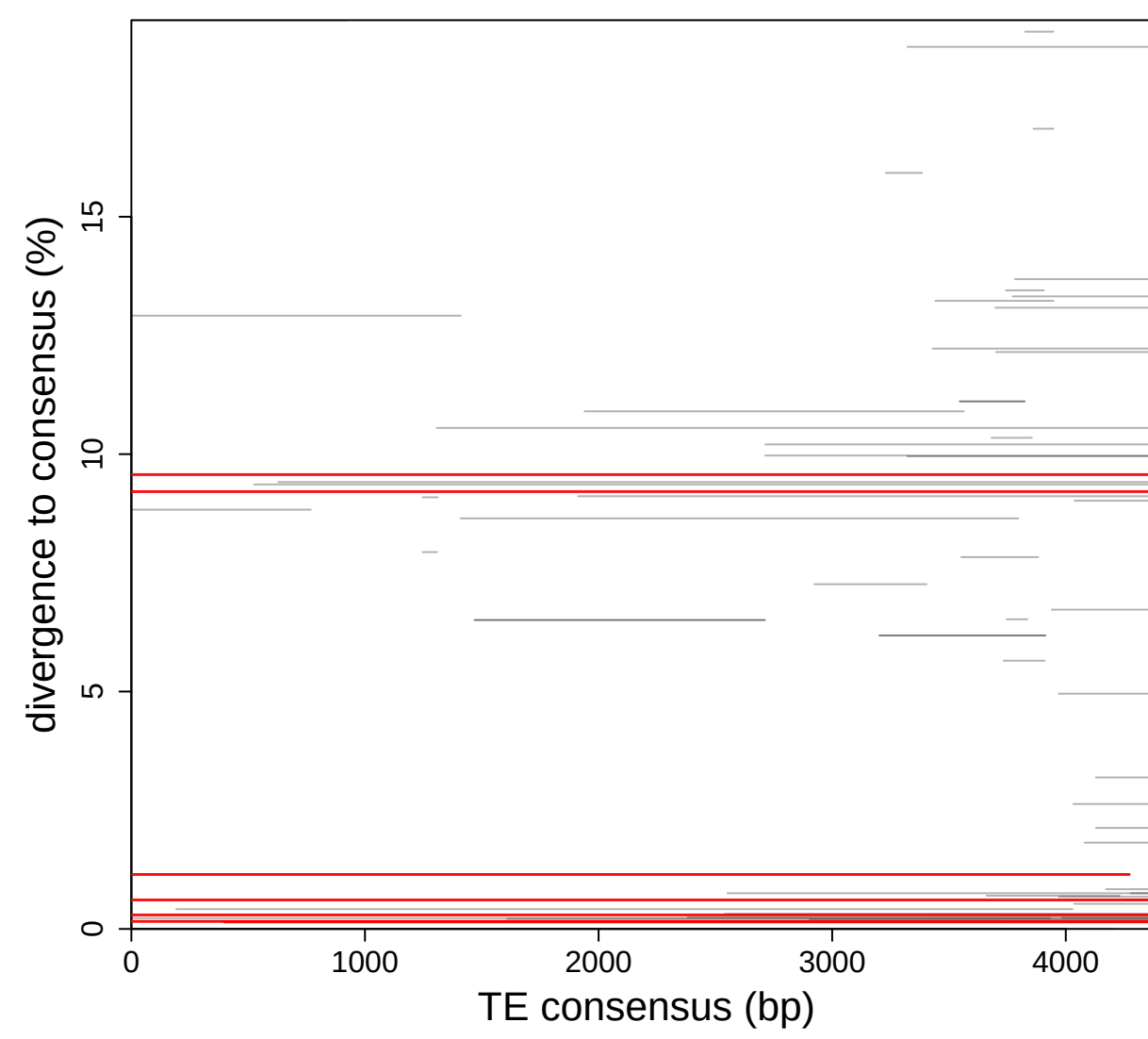
4000

5000



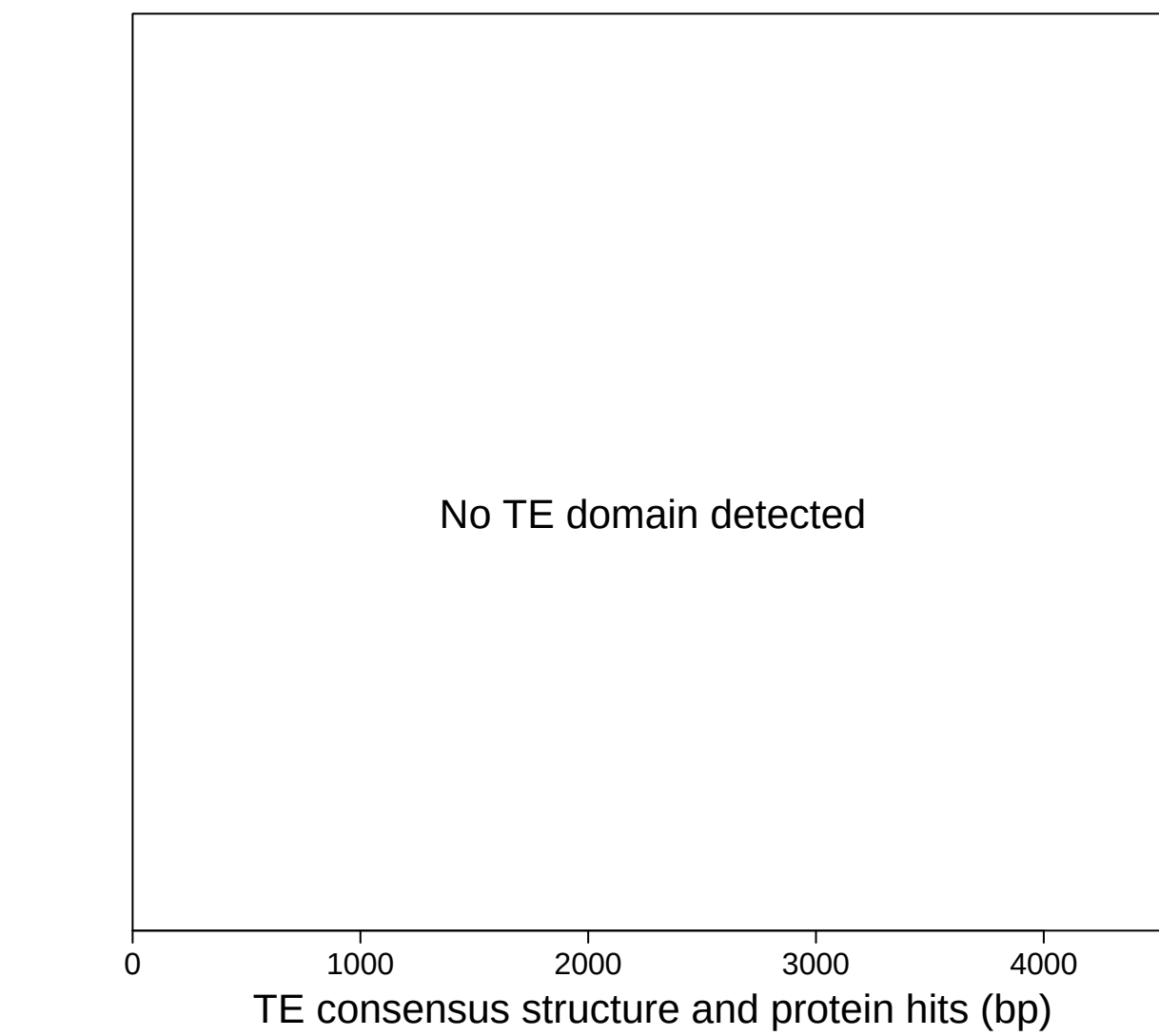
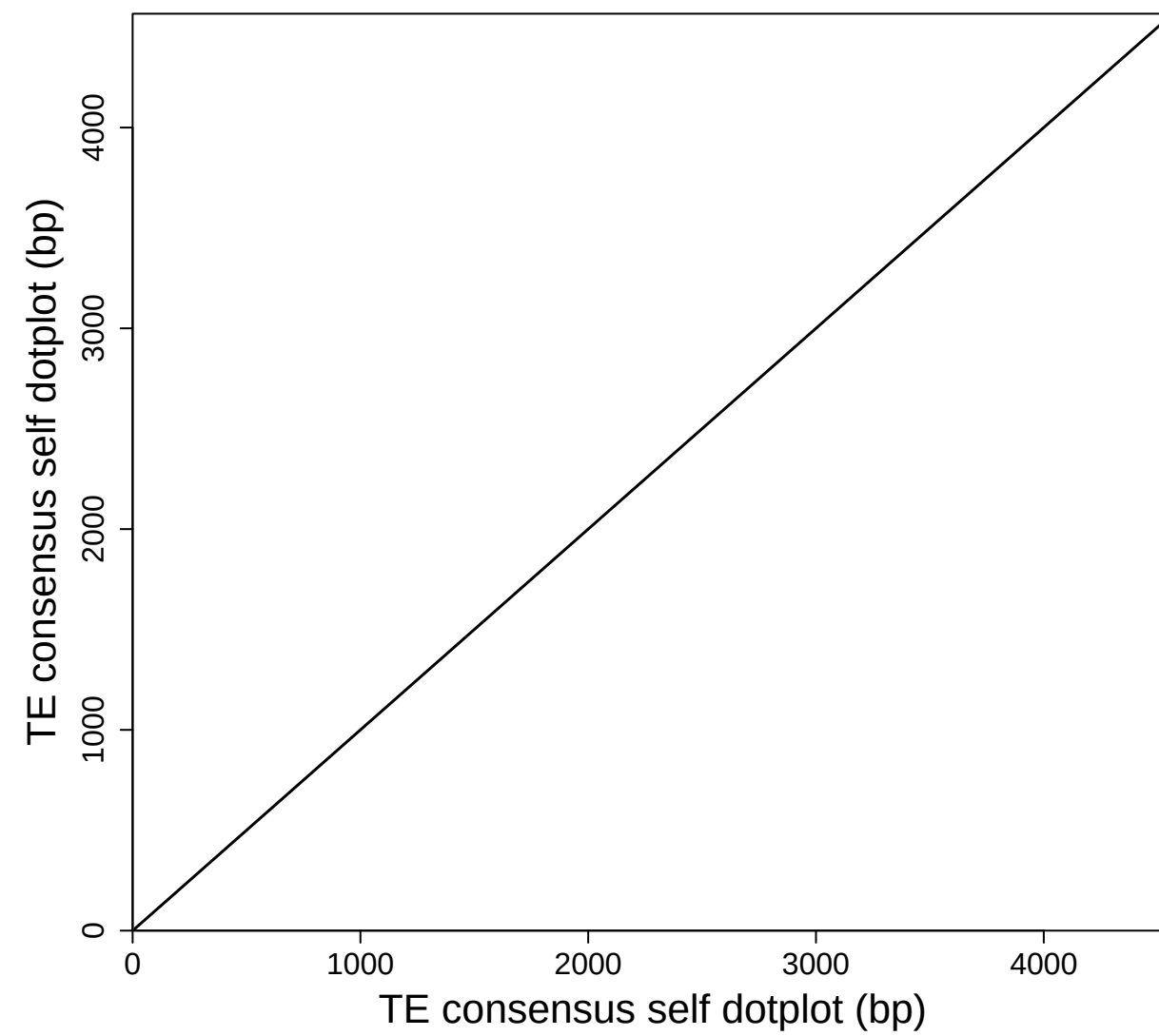
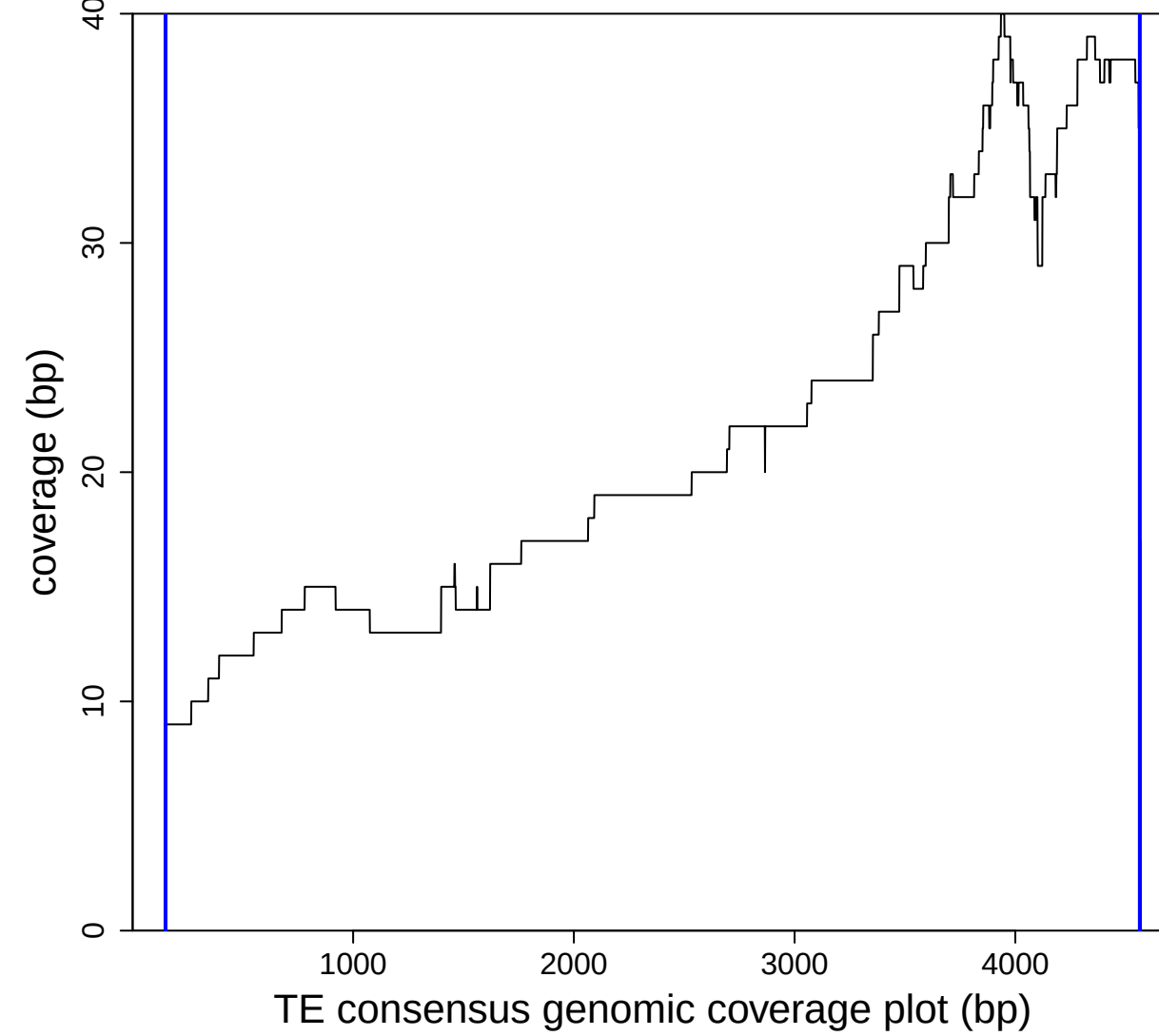
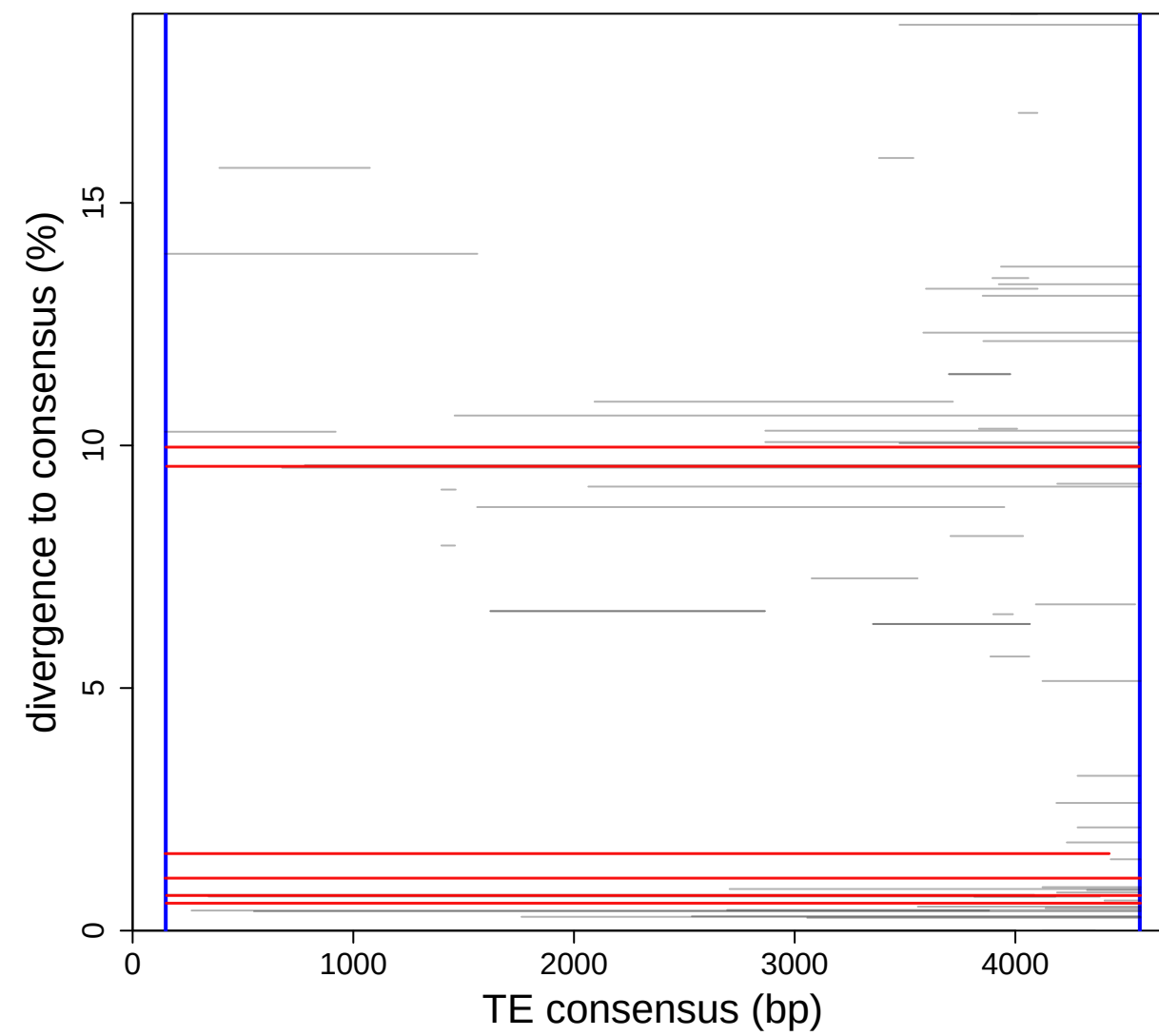
g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 4411bp; fragments: 67; full length: 7 (≥ 3969.9 bp)

After TETrimmer 4411 bp



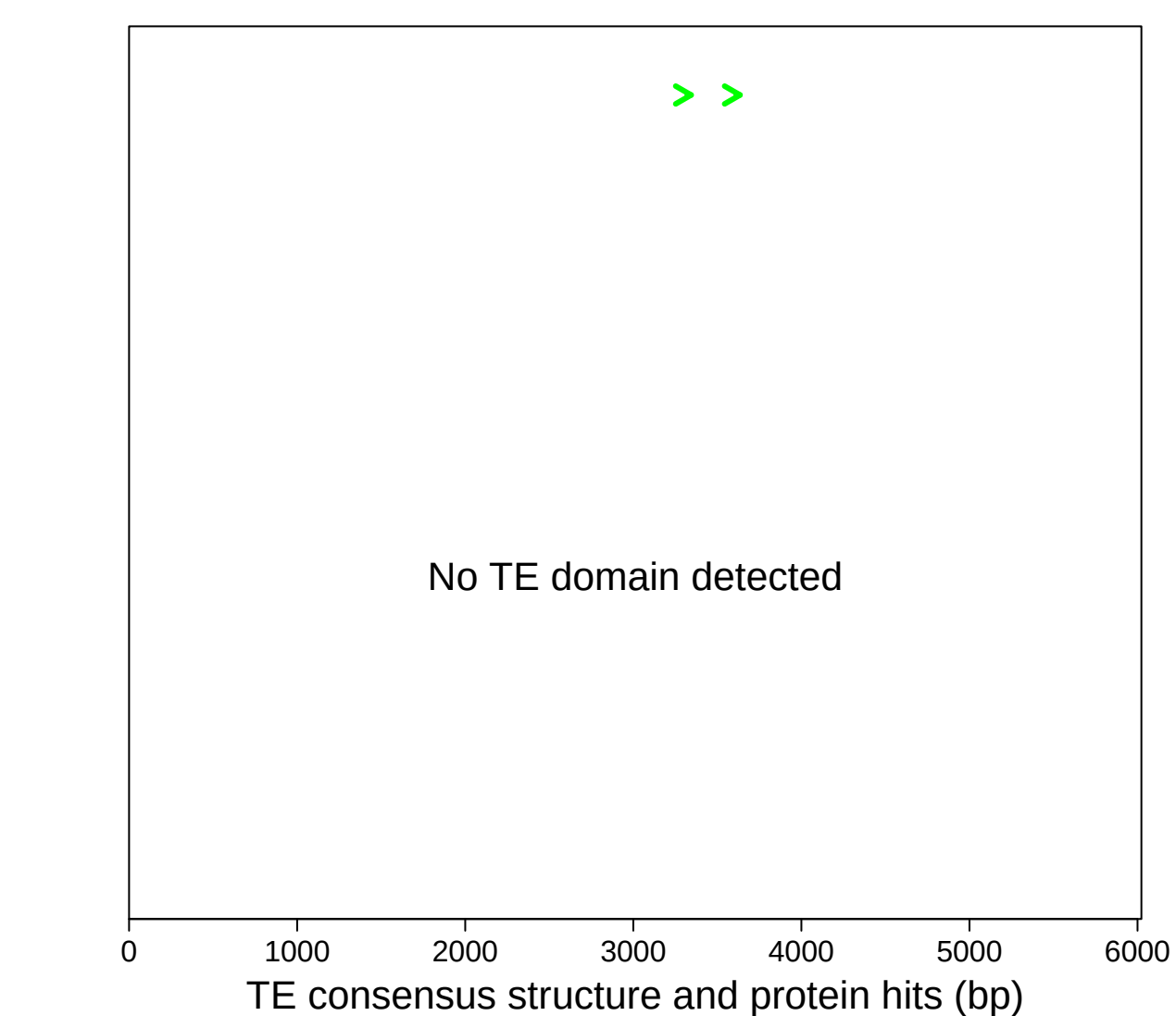
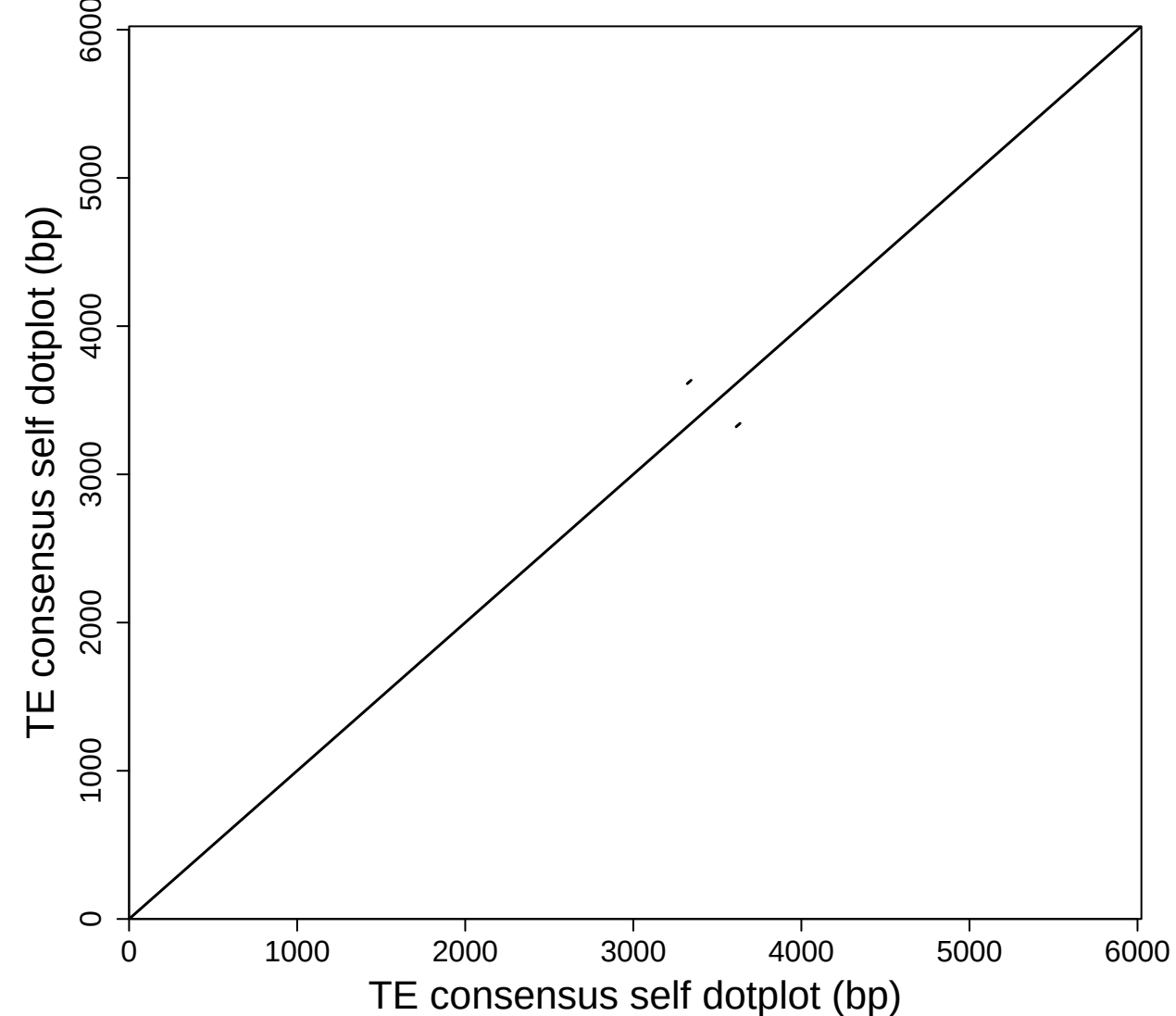
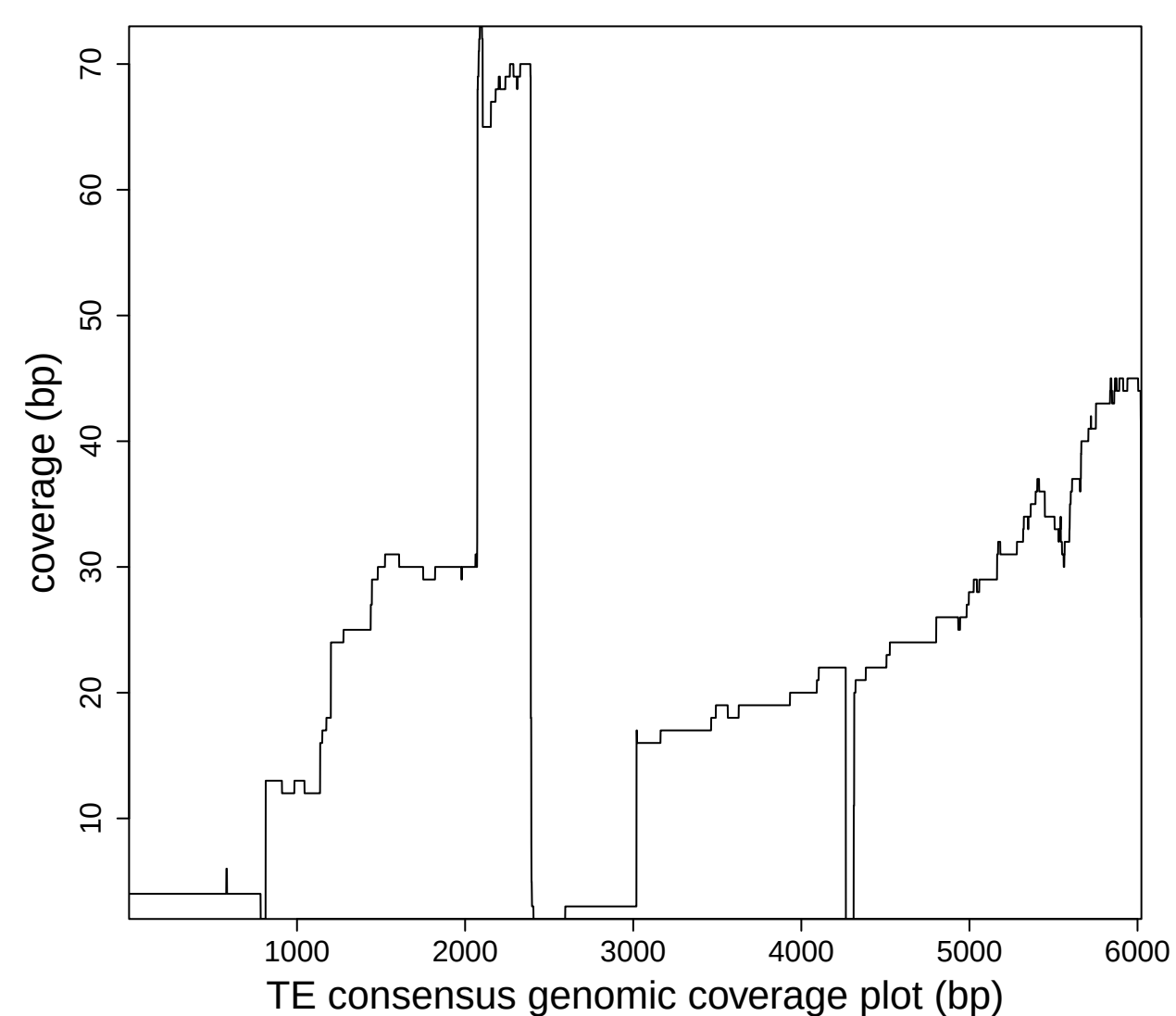
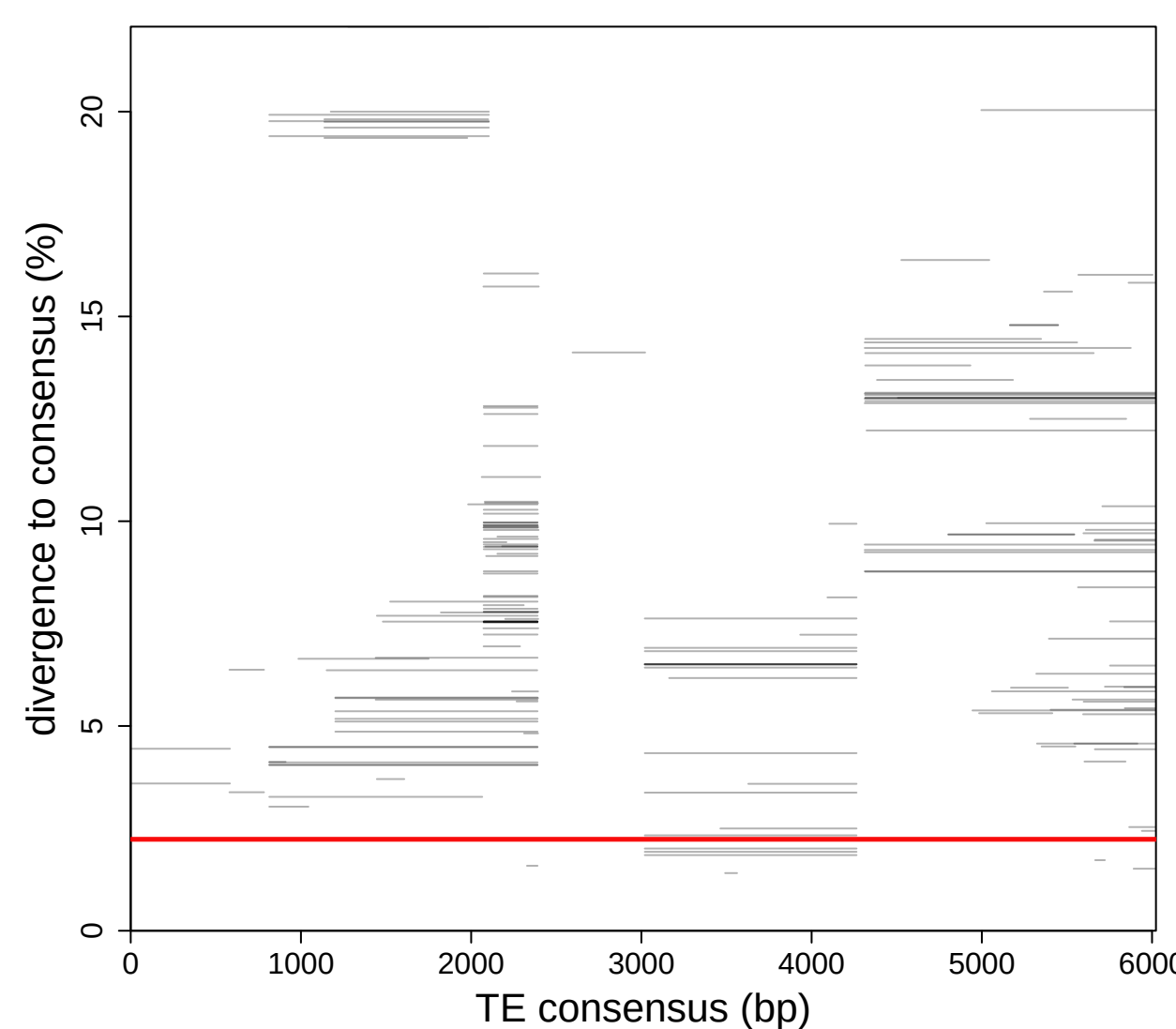
f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 4714bp; fragments: 67; full length: 6 (>=4242.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

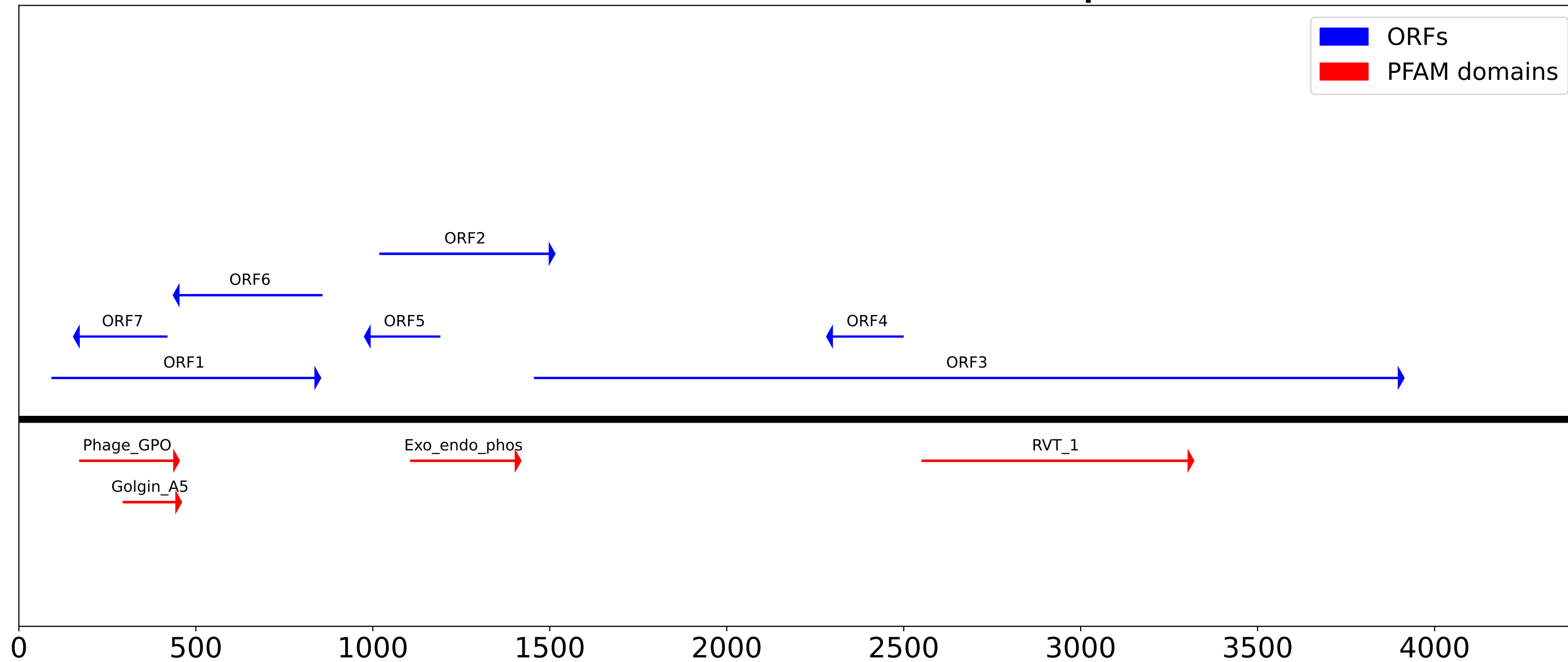


TE: rnd_5_family_586
size: 6023bp; fragments: 175; full length: 2 (≥ 5420.7 bp)

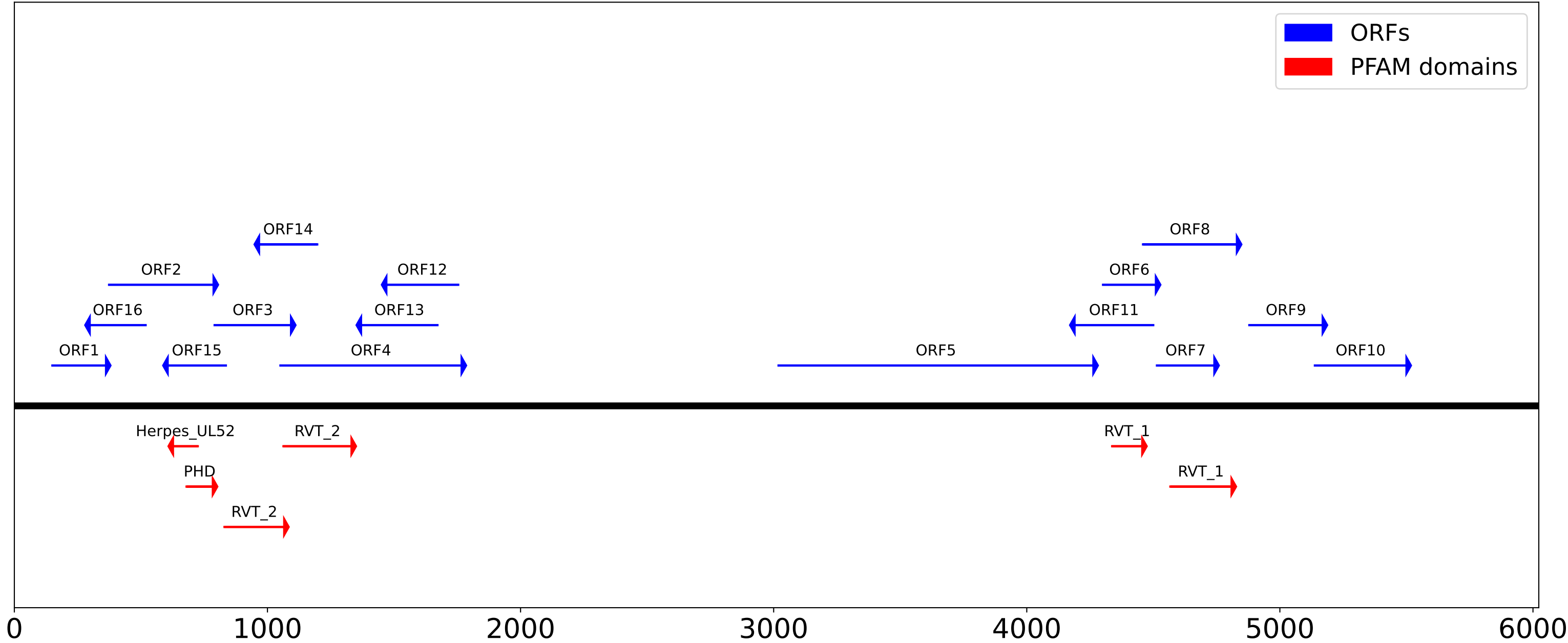
Before TEtrimmer 6023 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

