

Start Crop Point

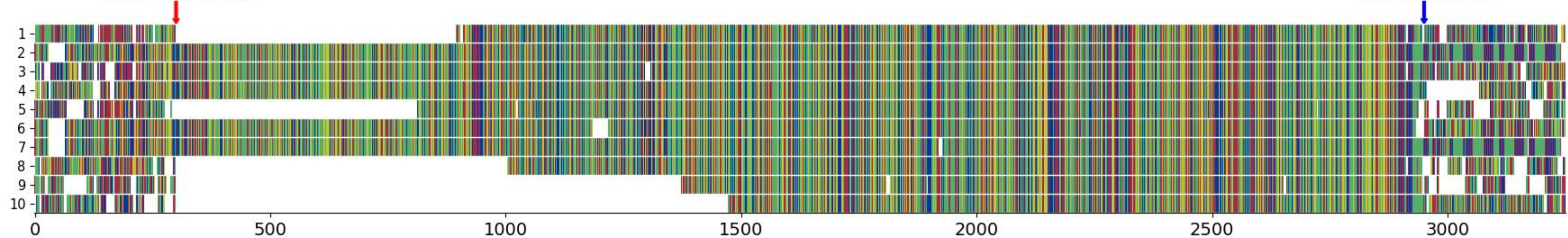
End Crop Point

MSA length = 2651



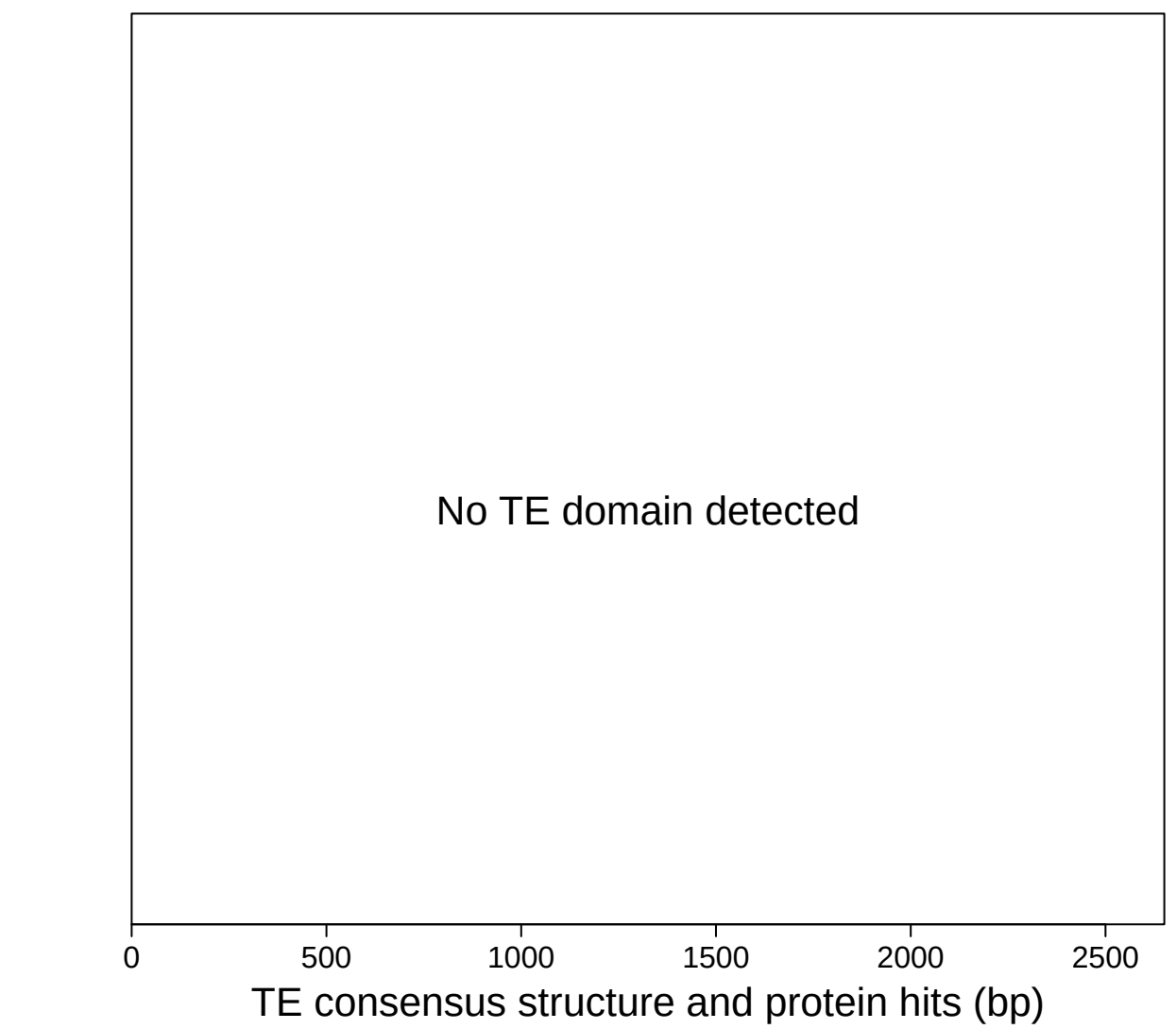
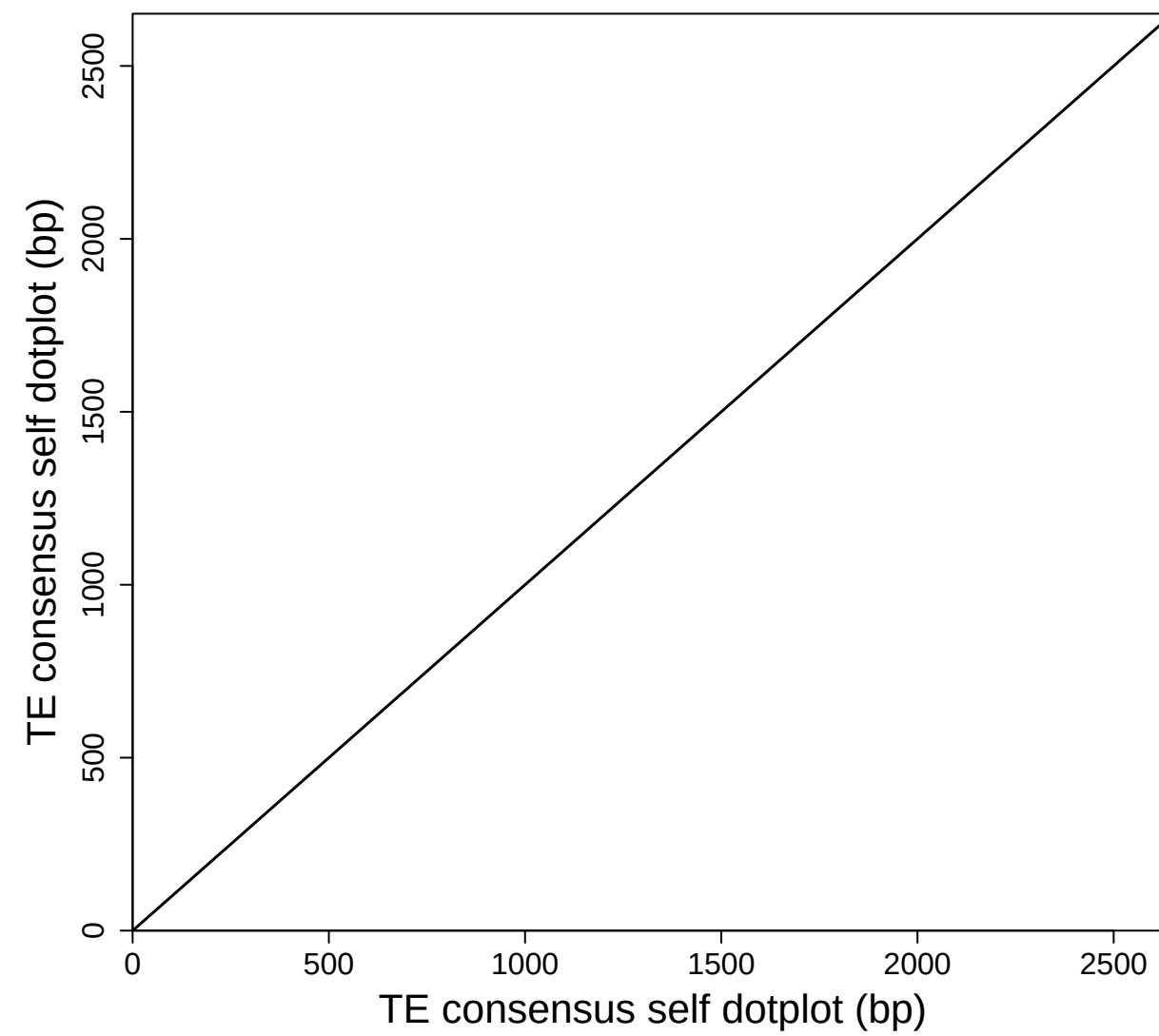
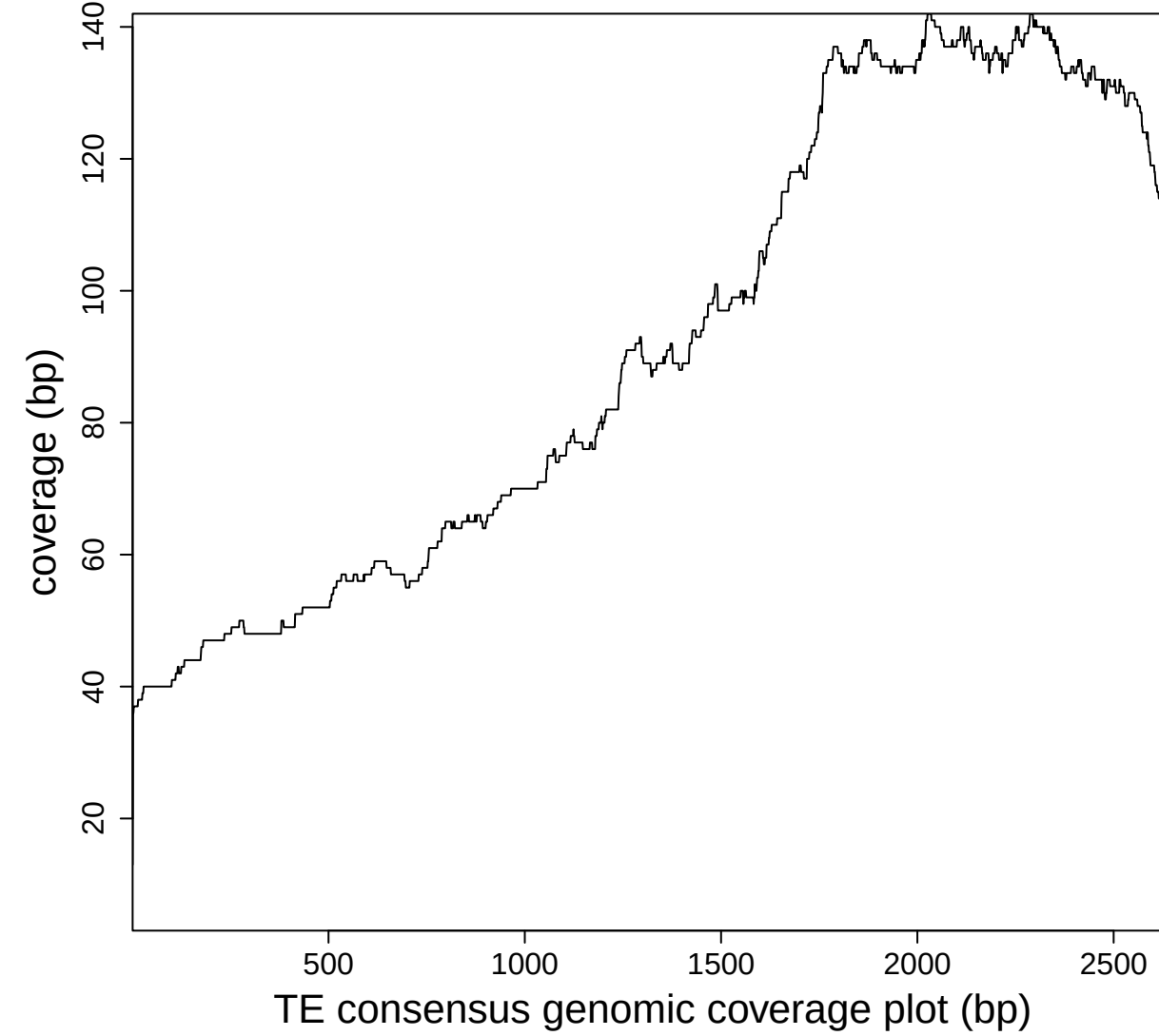
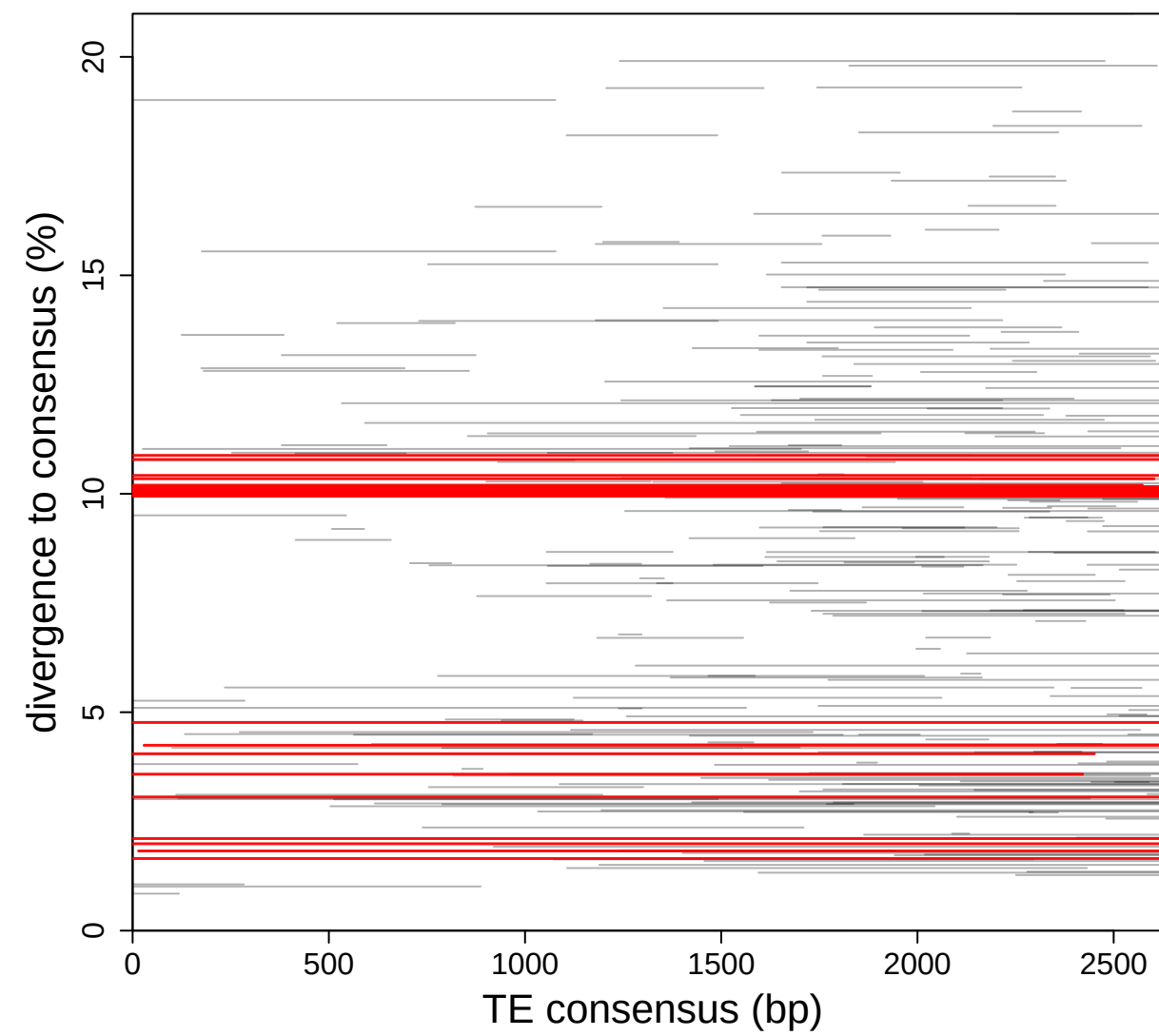
Start crop Point

End crop Point



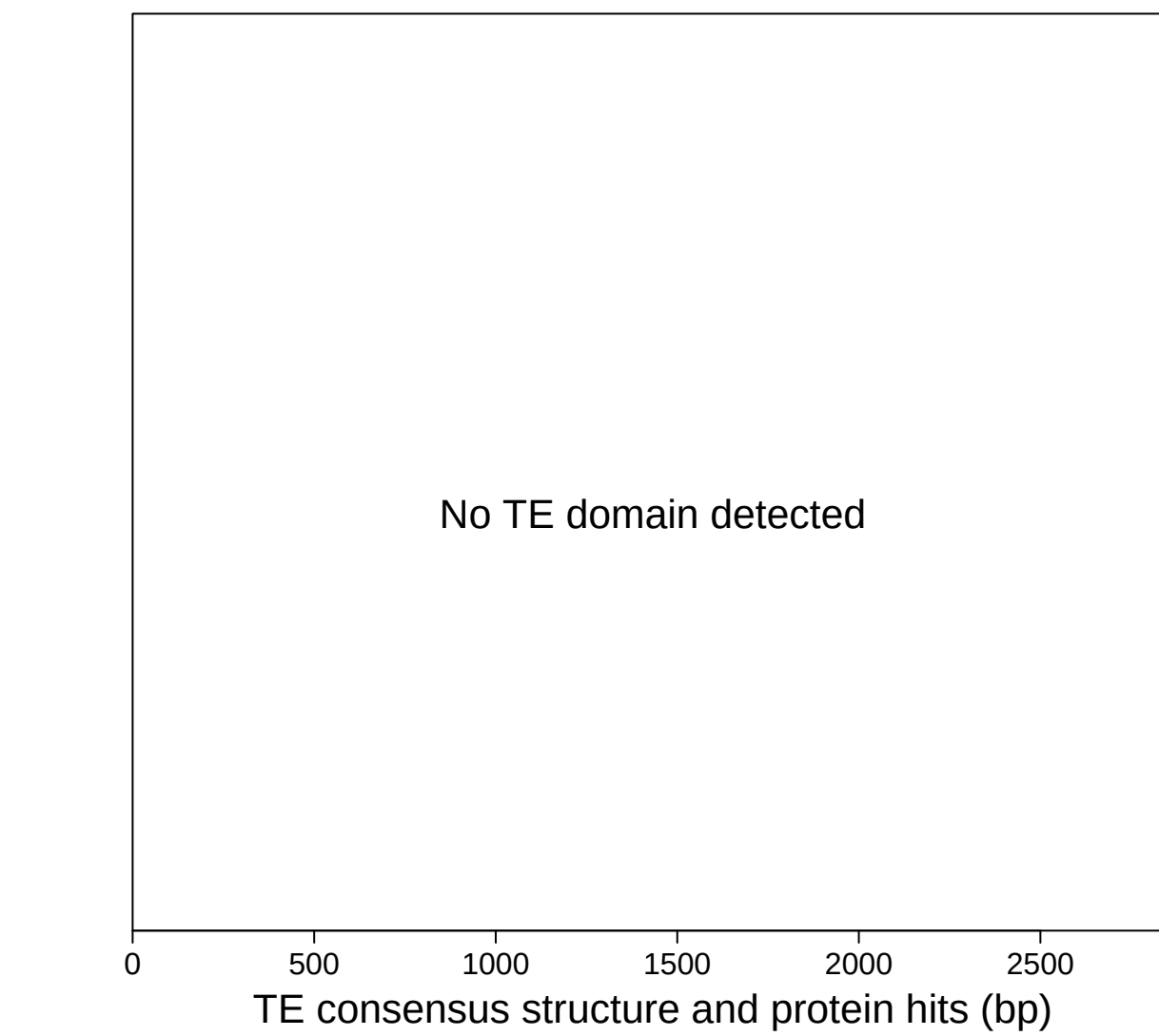
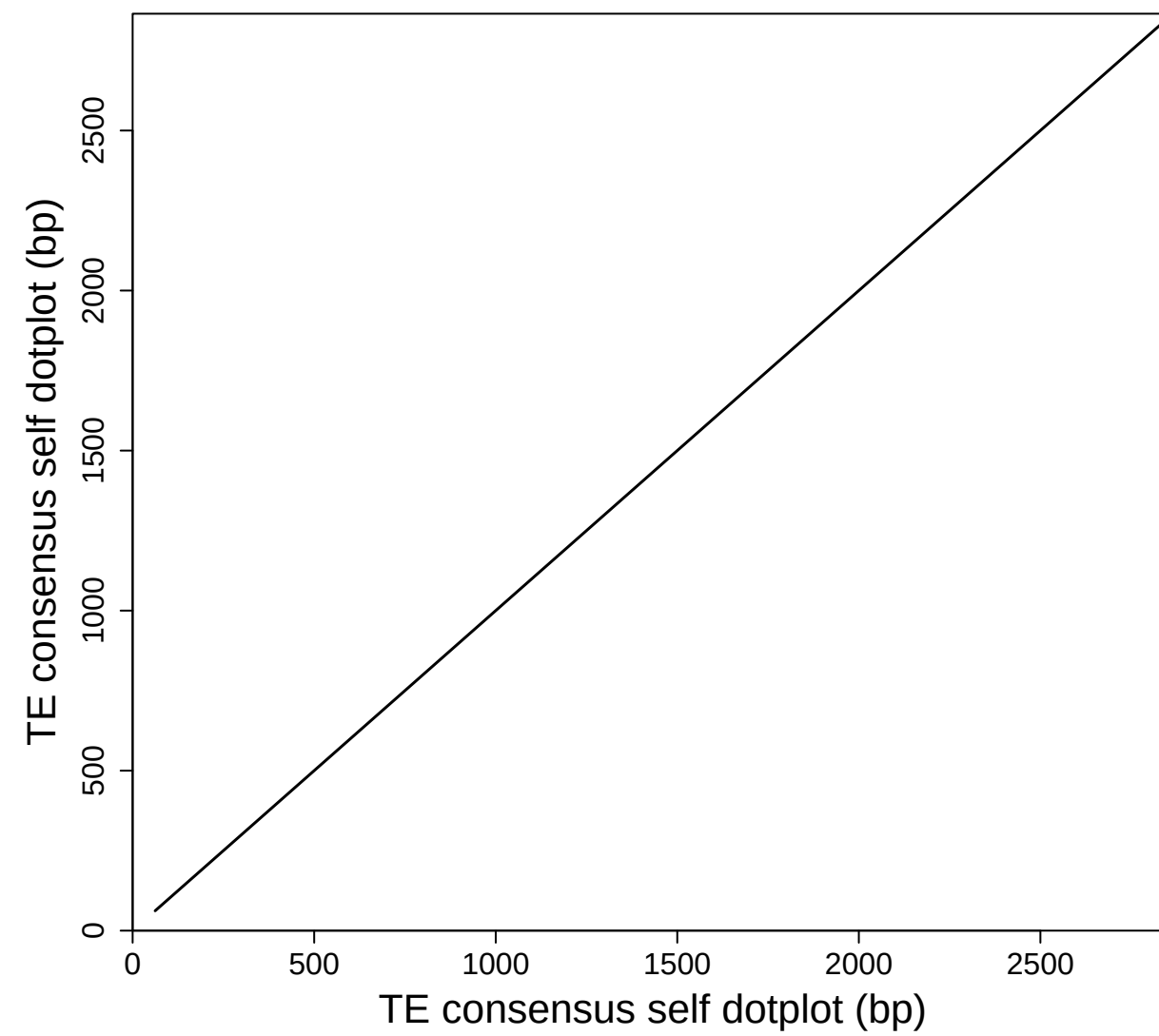
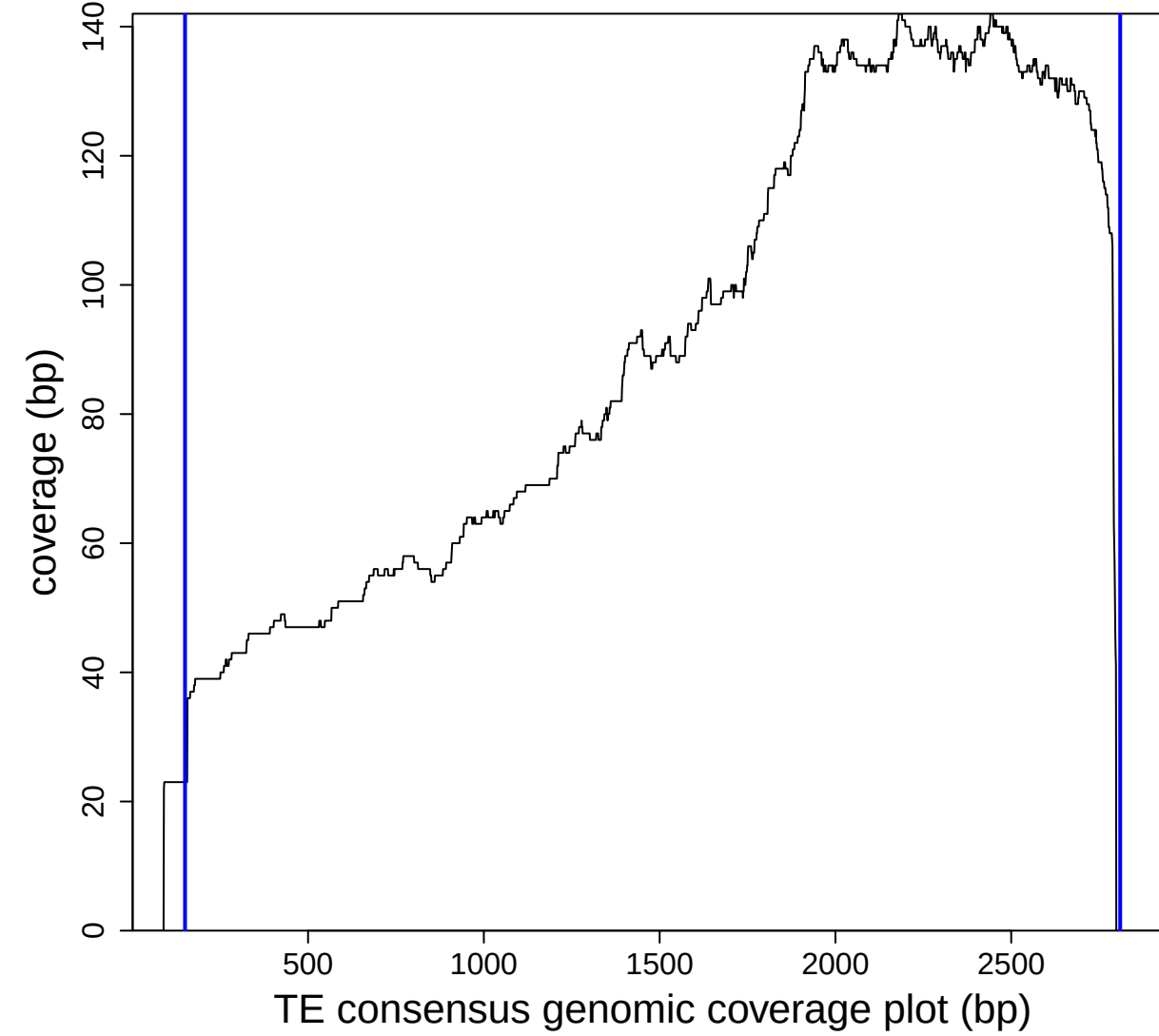
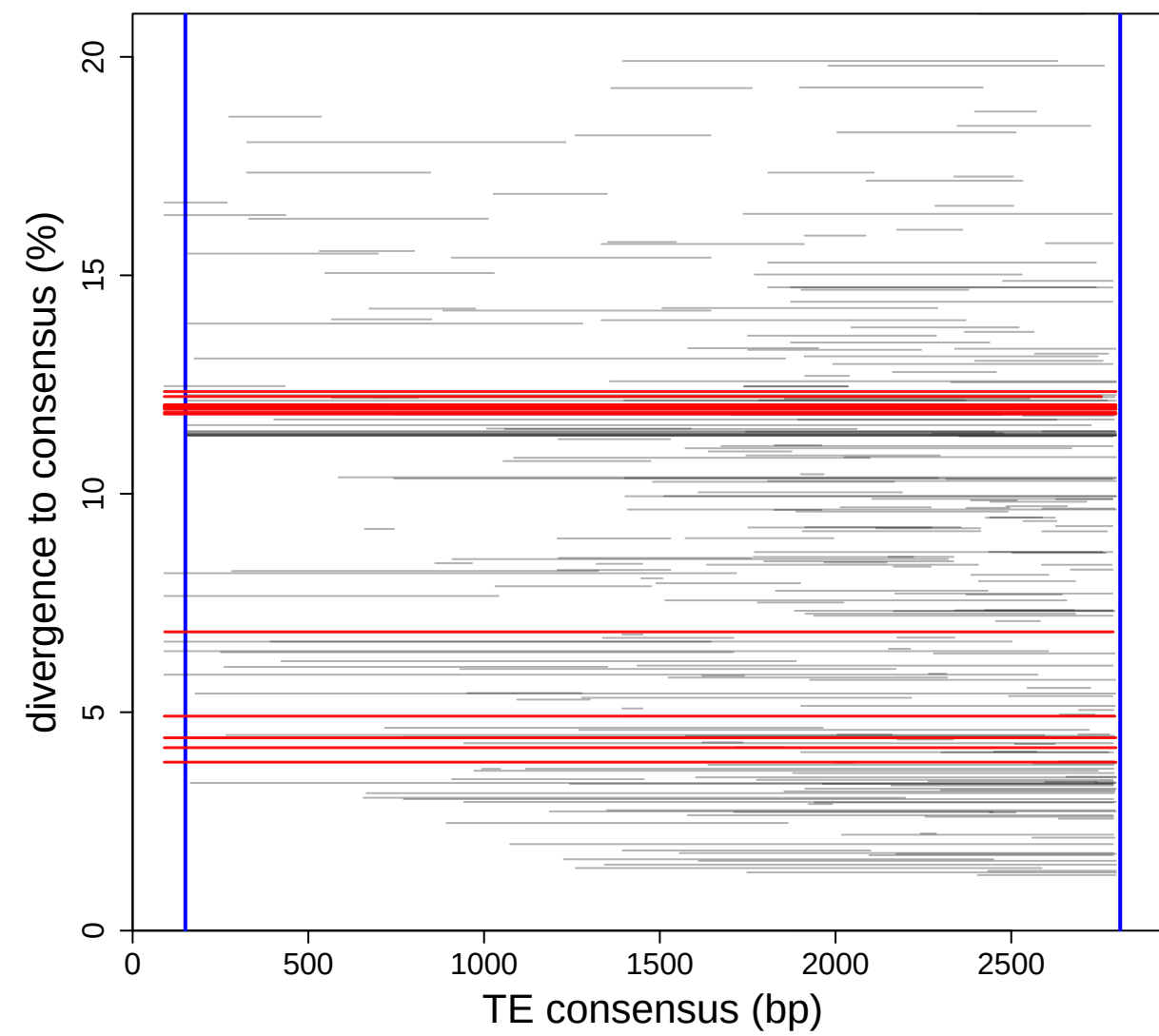
bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_aln.fa
size: 2651bp; fragments: 282; full length: 29 (≥ 2385.9 bp)

After TETrimmer 2651 bp



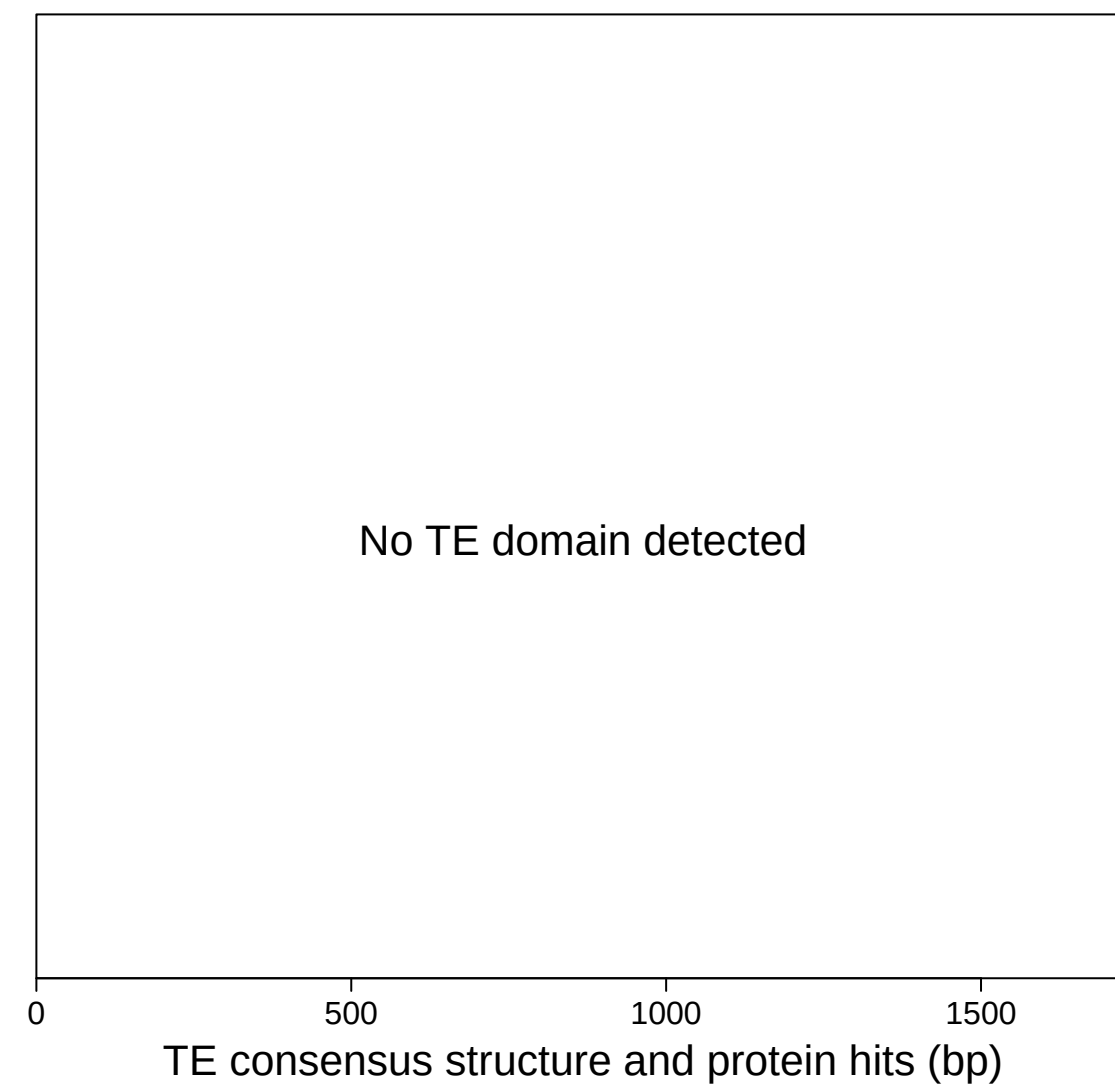
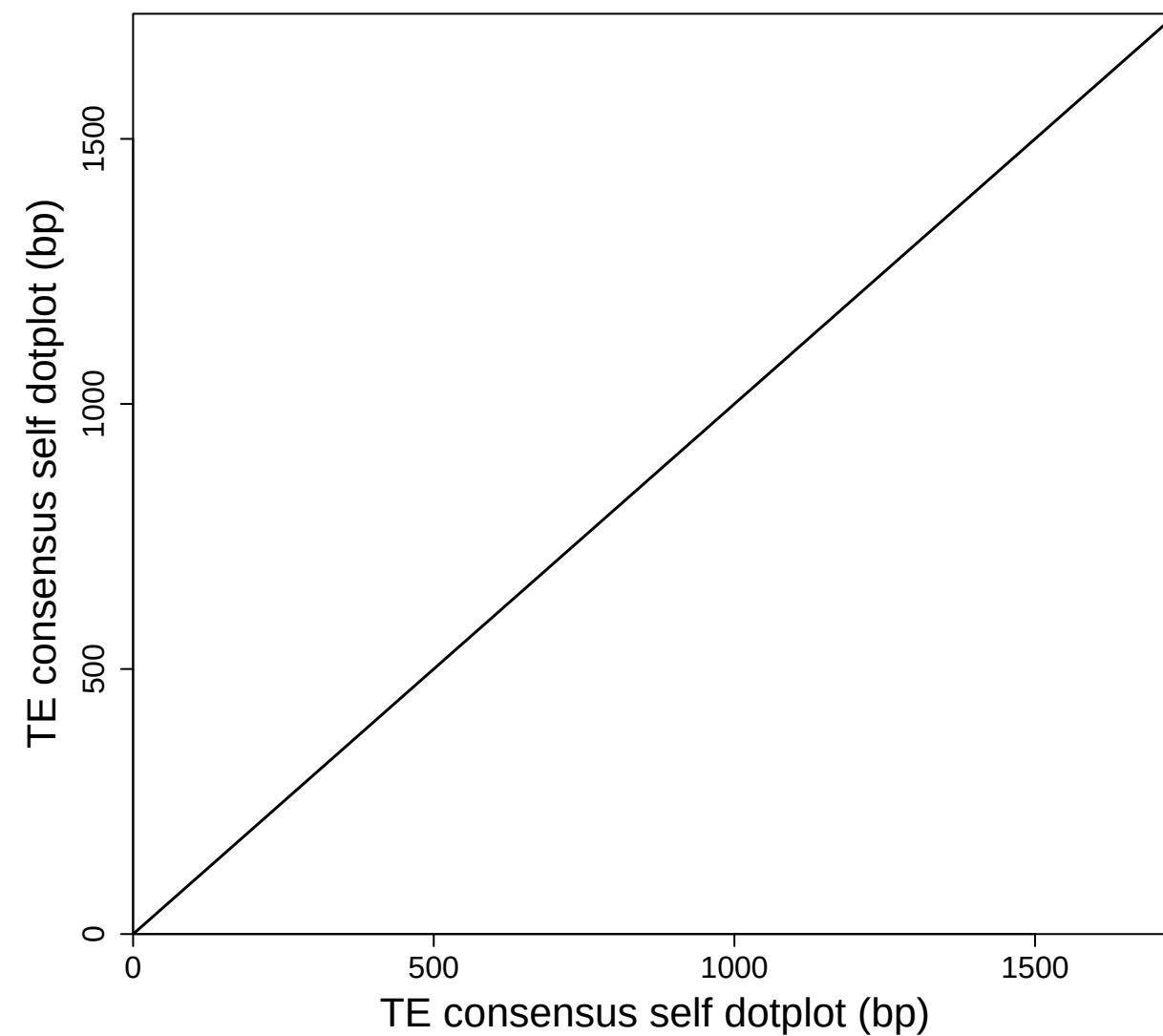
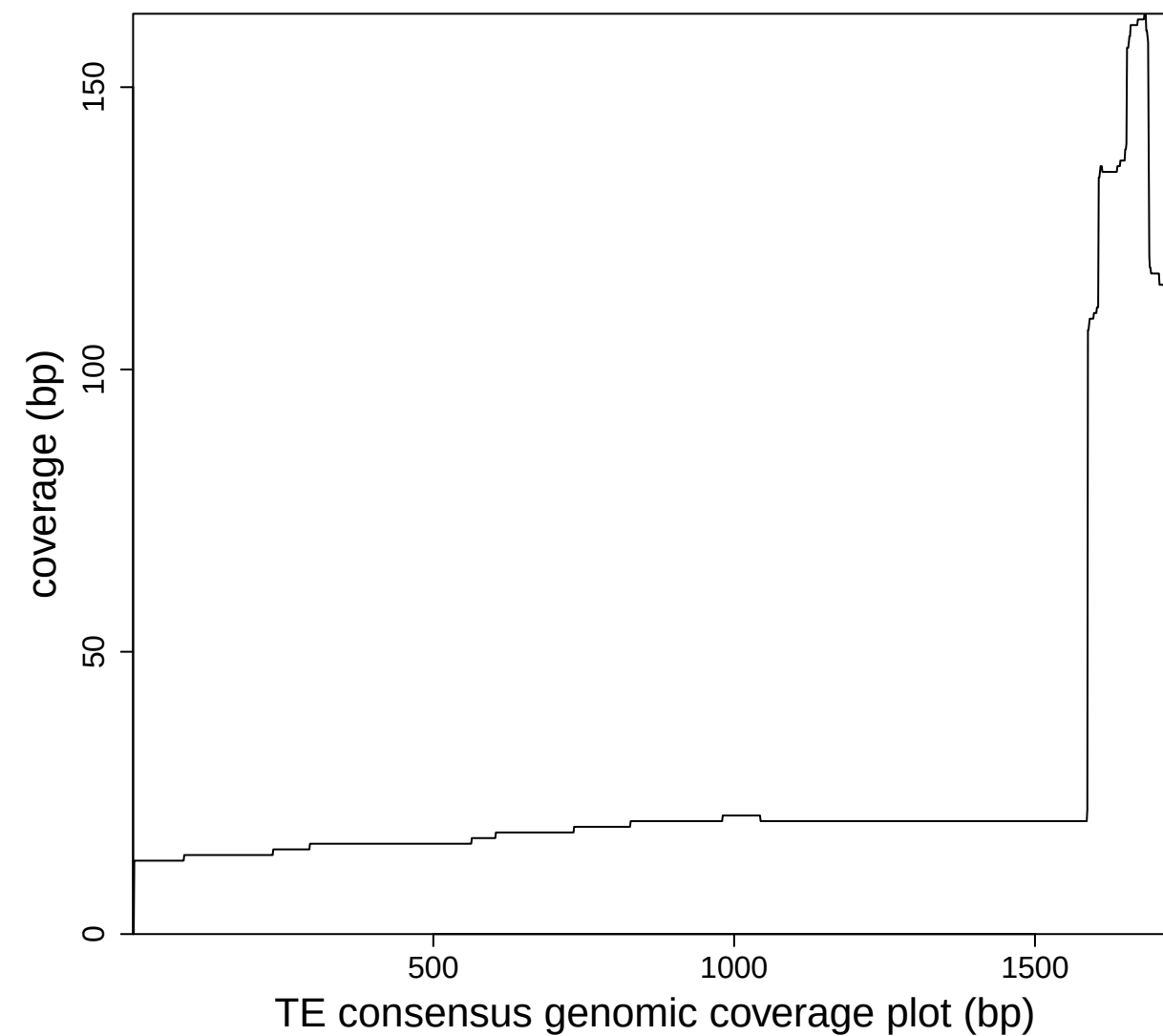
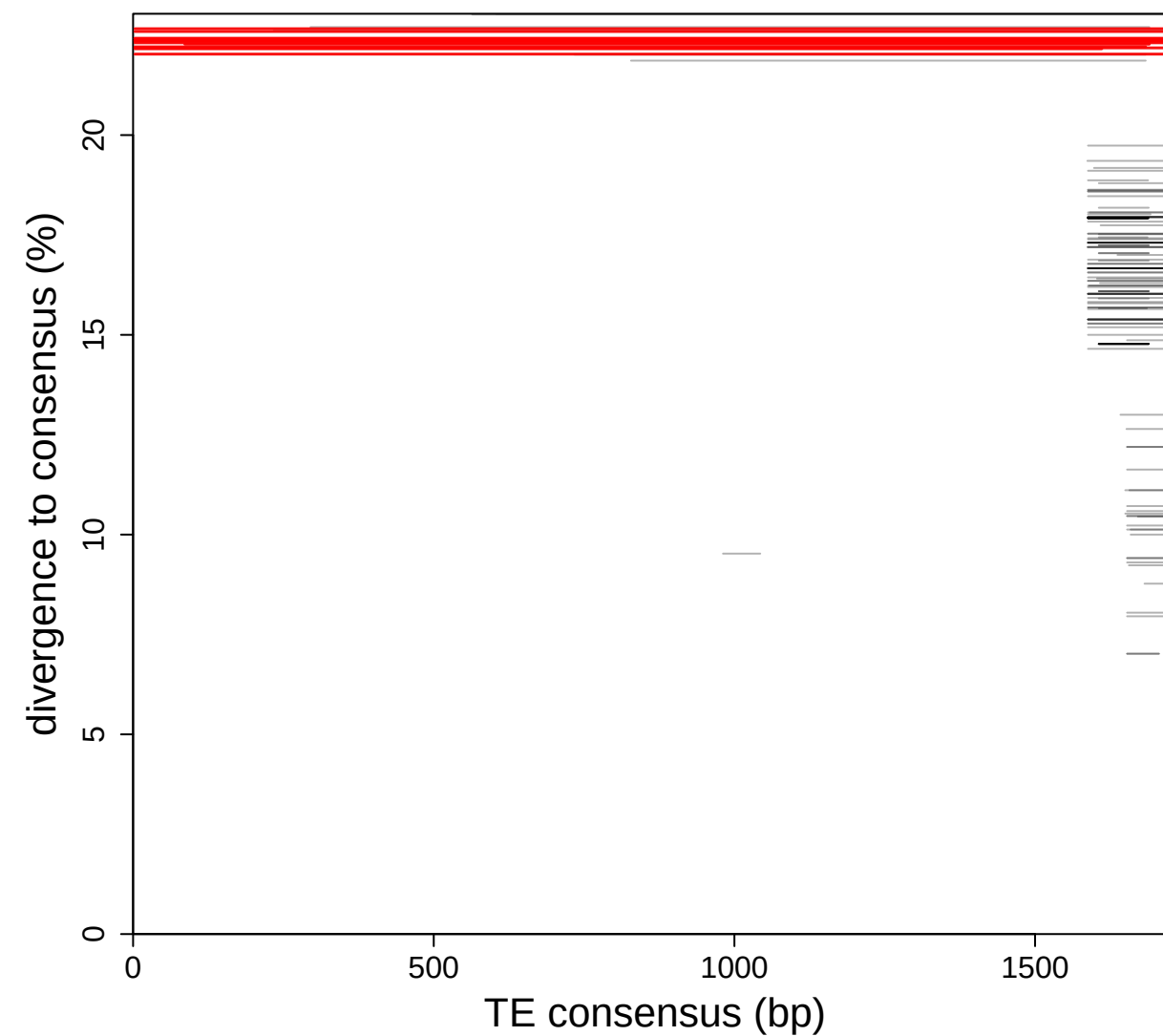
bed_g_4.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 2960bp; fragments: 281; full length: 14 (>=2664bp)

After TEtrimmer Extended plot Blue lines are boundaries

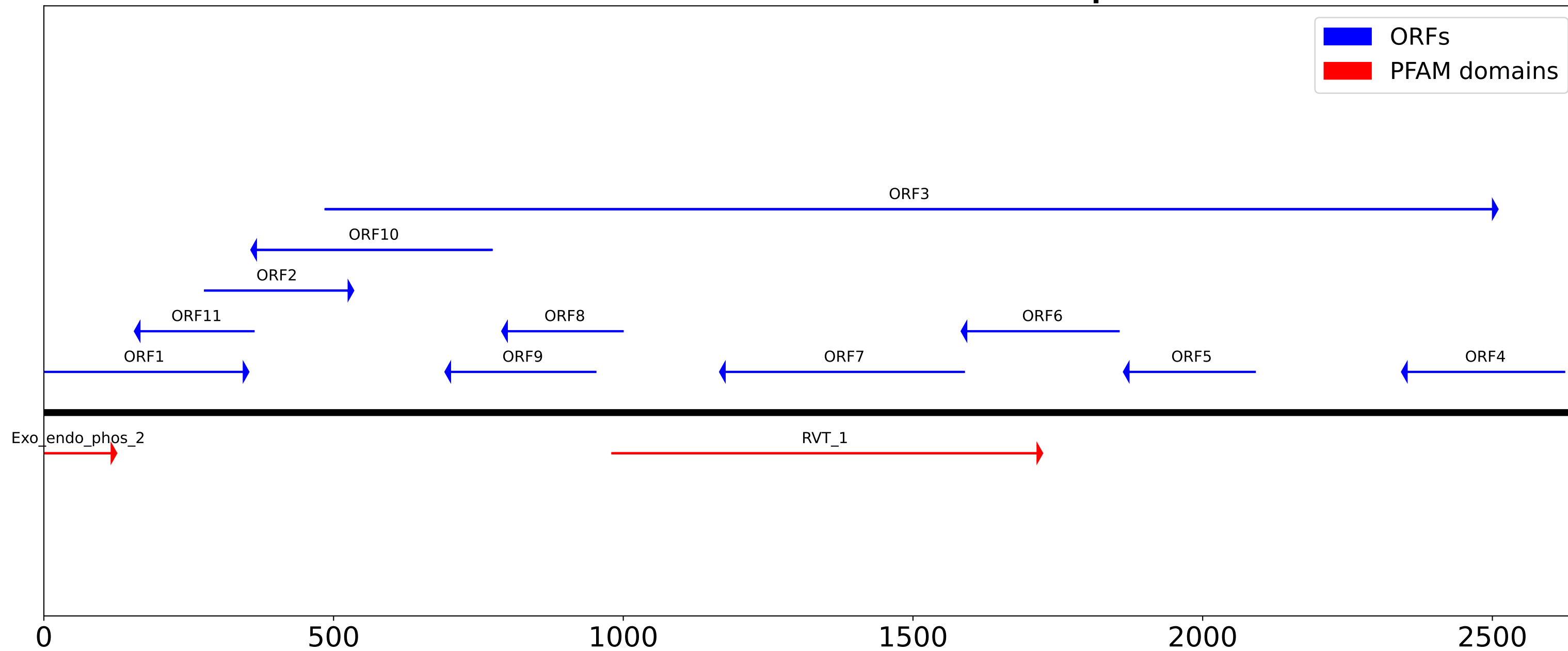


TE: rnd_2_family_80
size: 1736bp; fragments: 165; full length: 14 (≥ 1562.4 bp)

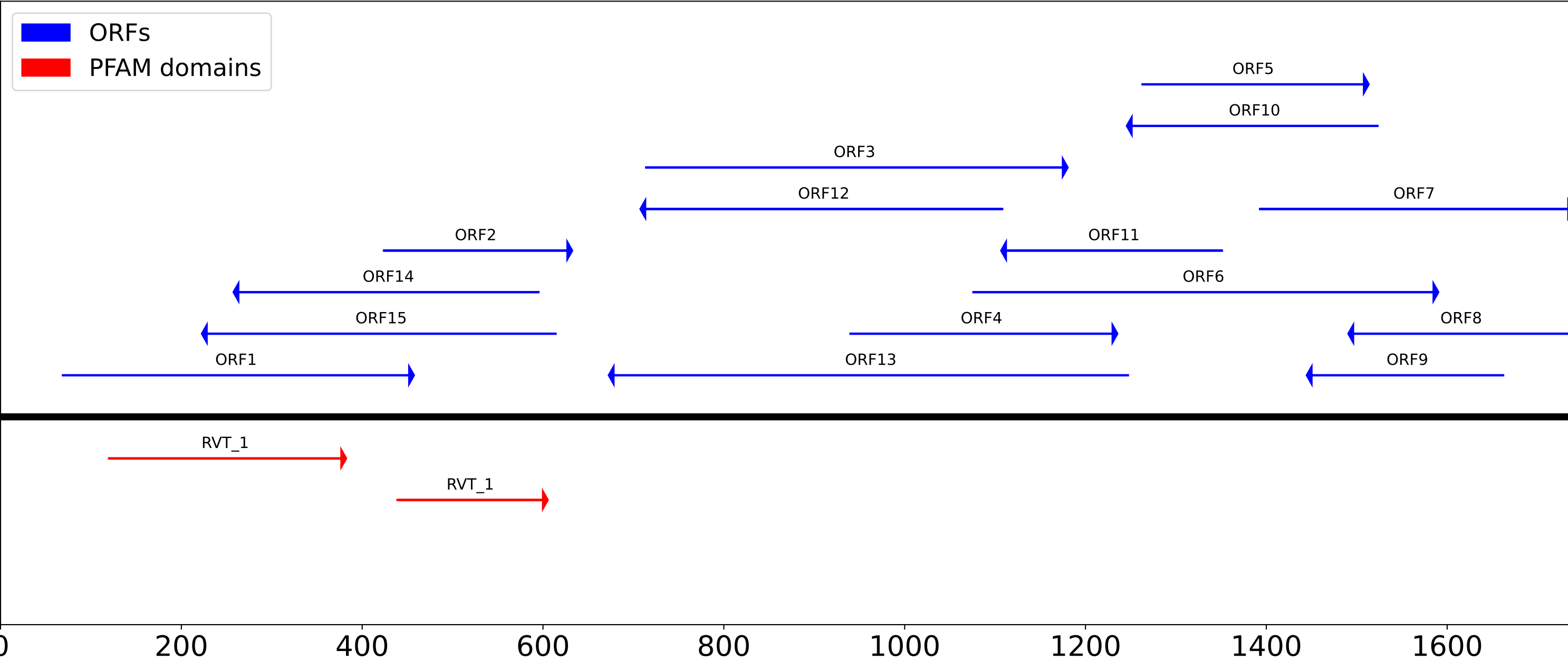
Before TEtrimmer 1736 bp



After TEtrimmer ORF and PFAM domain plot



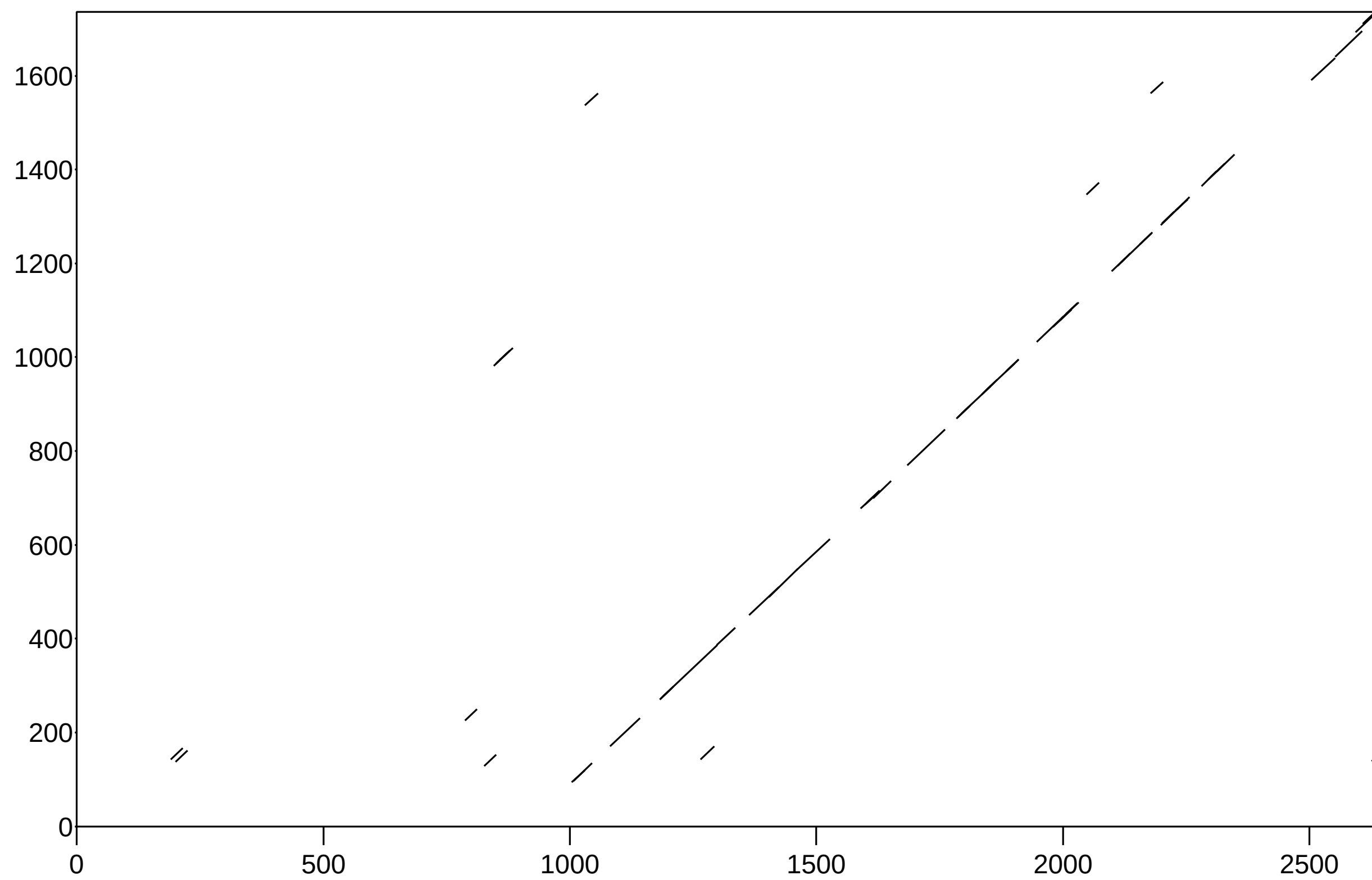
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)