

Start crop Point



End crop Point

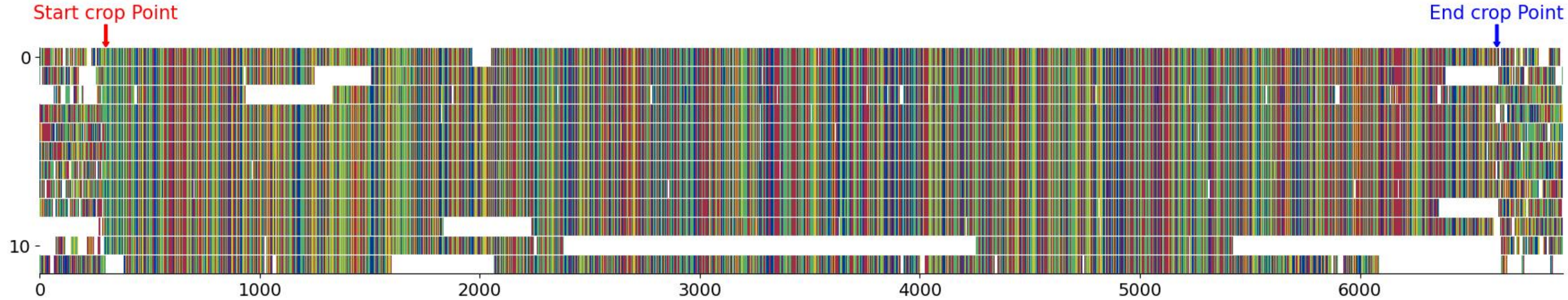


MSA length = 6324



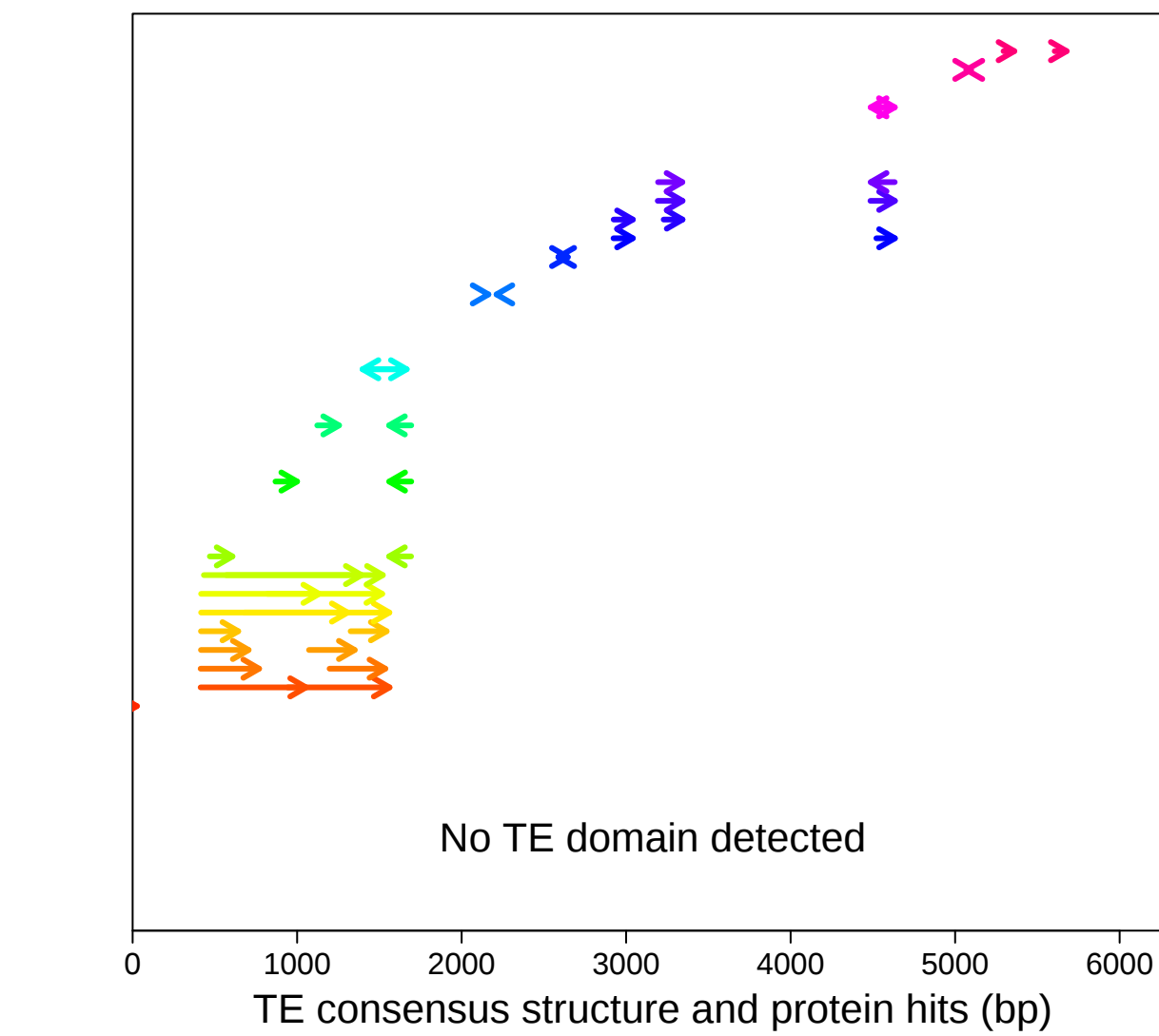
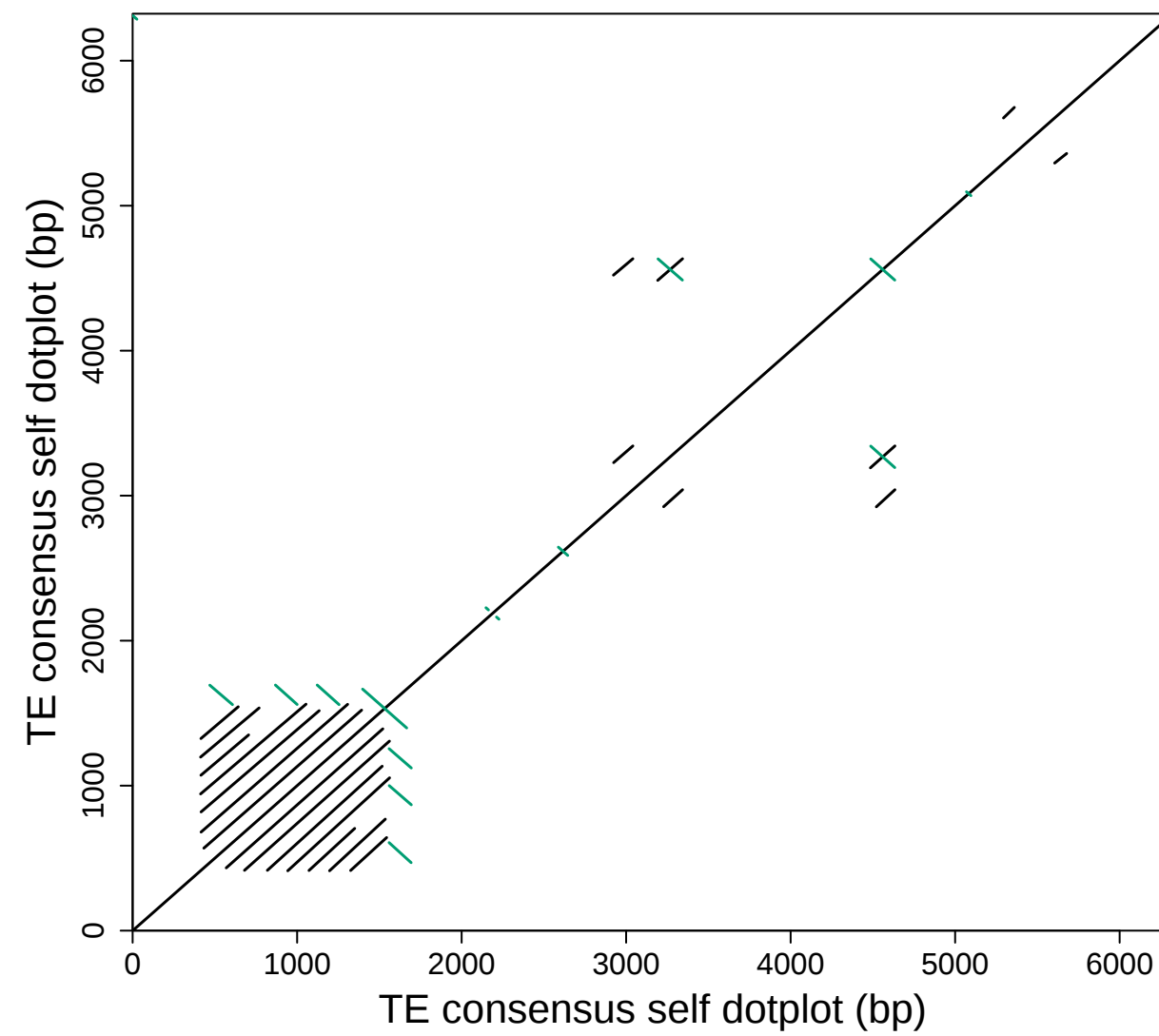
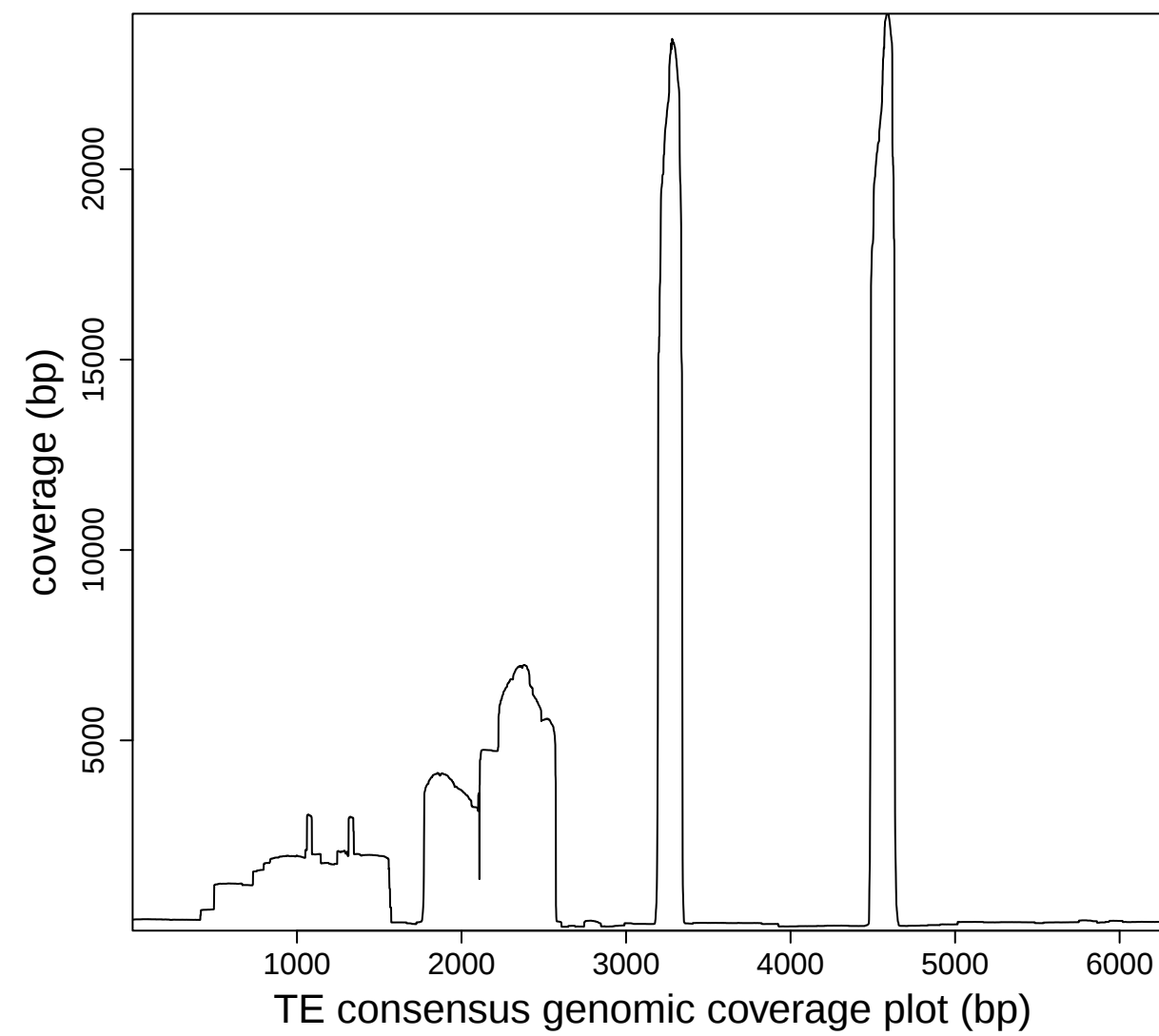
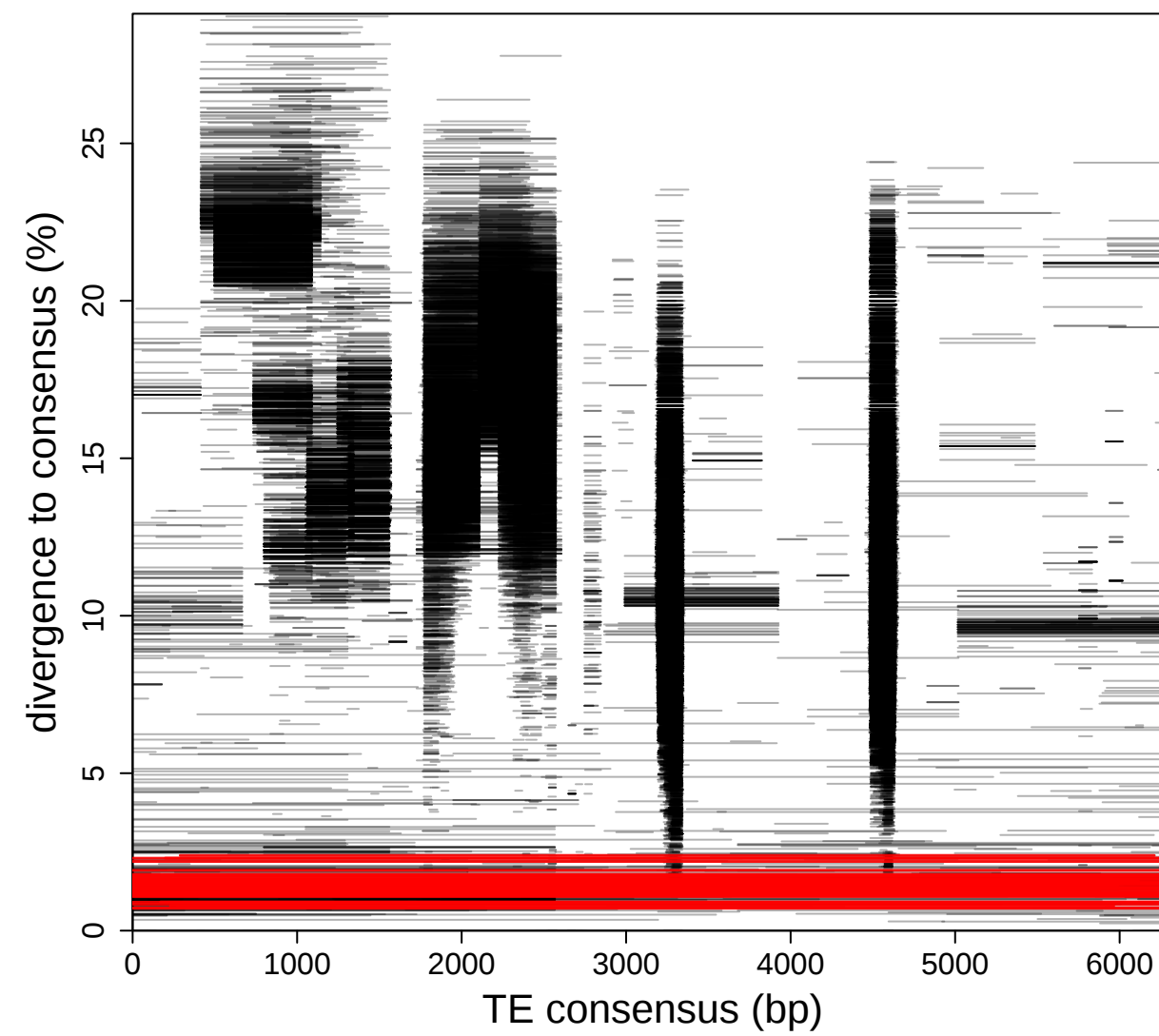
Start crop Point

End crop Point



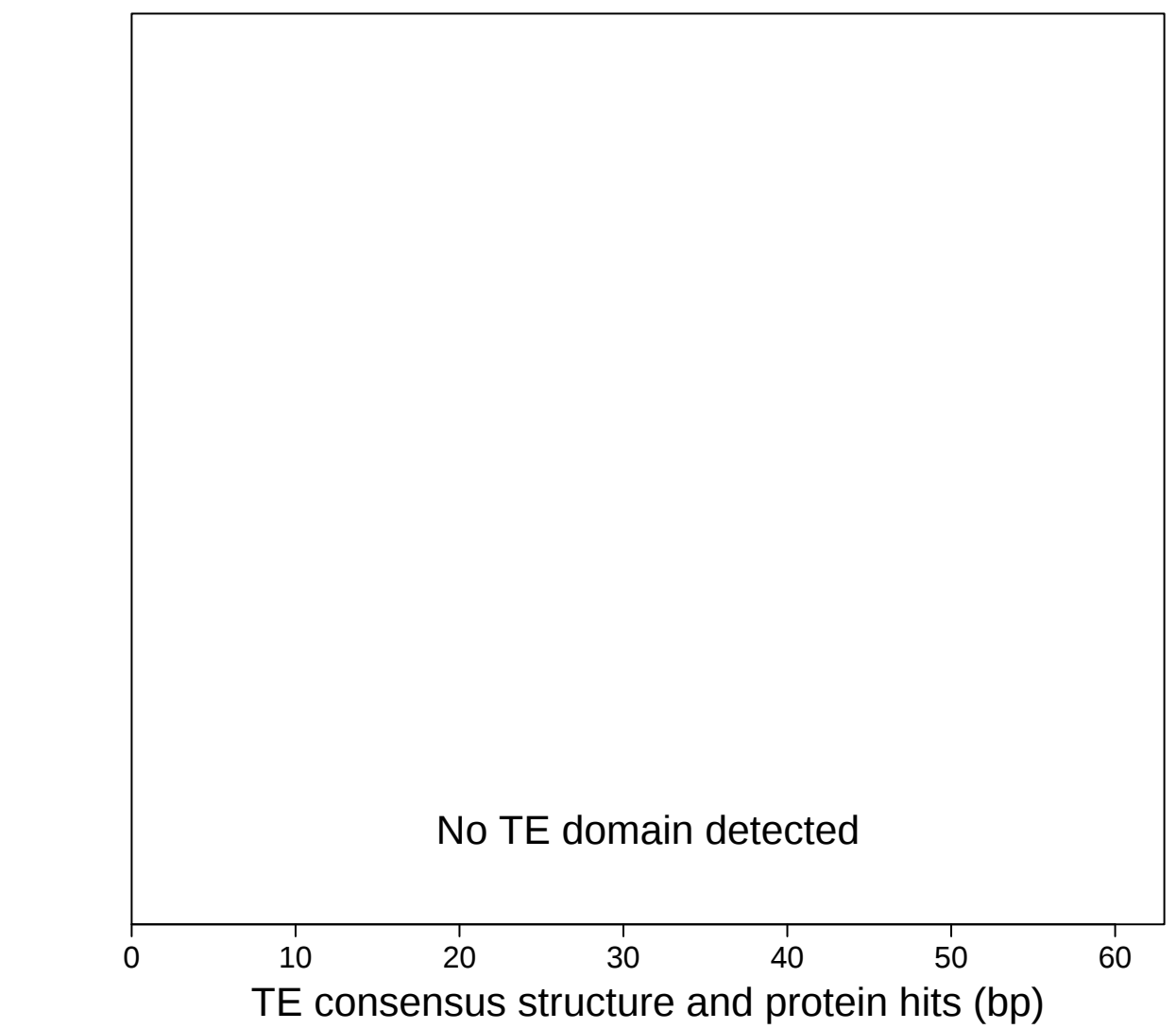
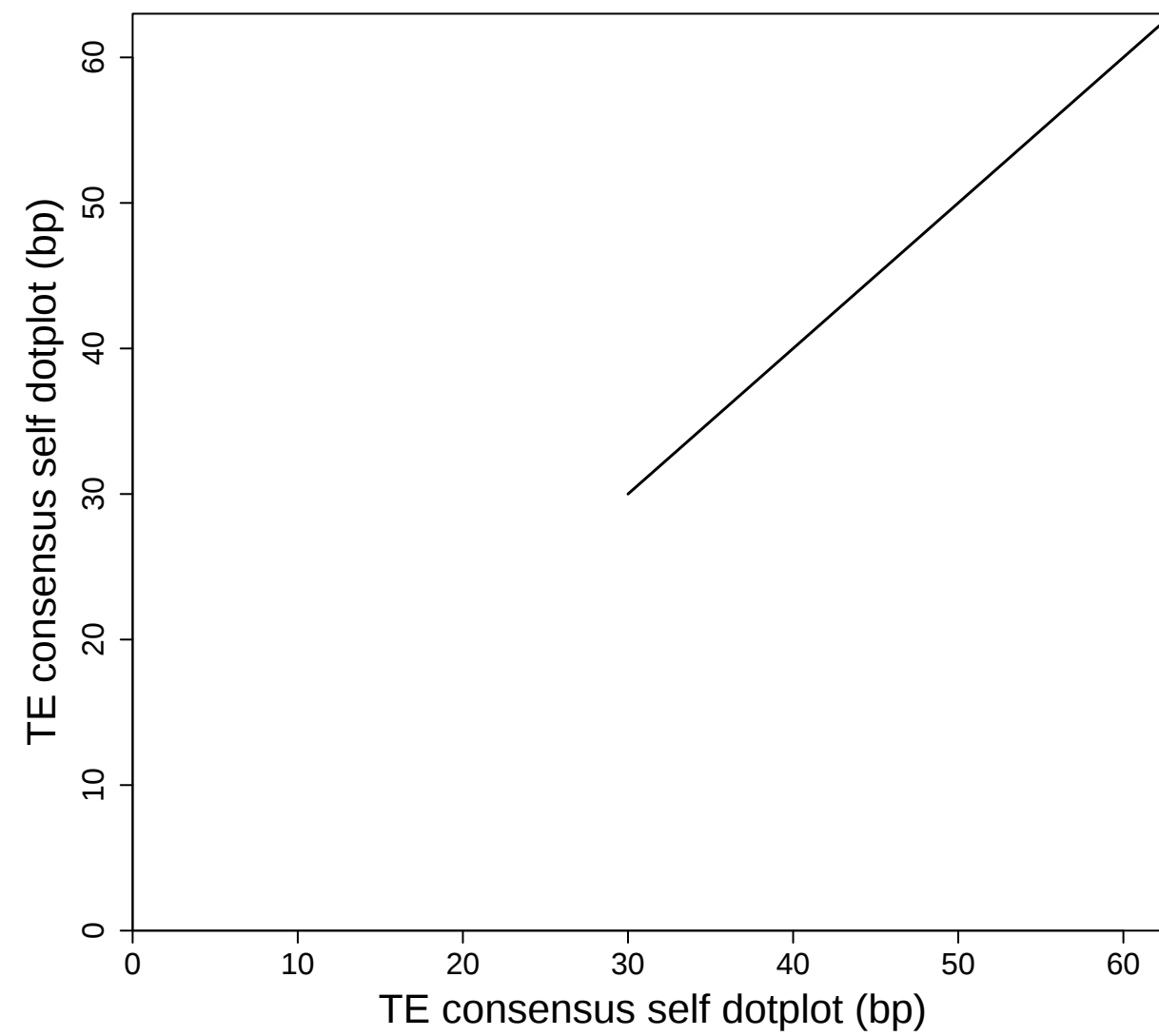
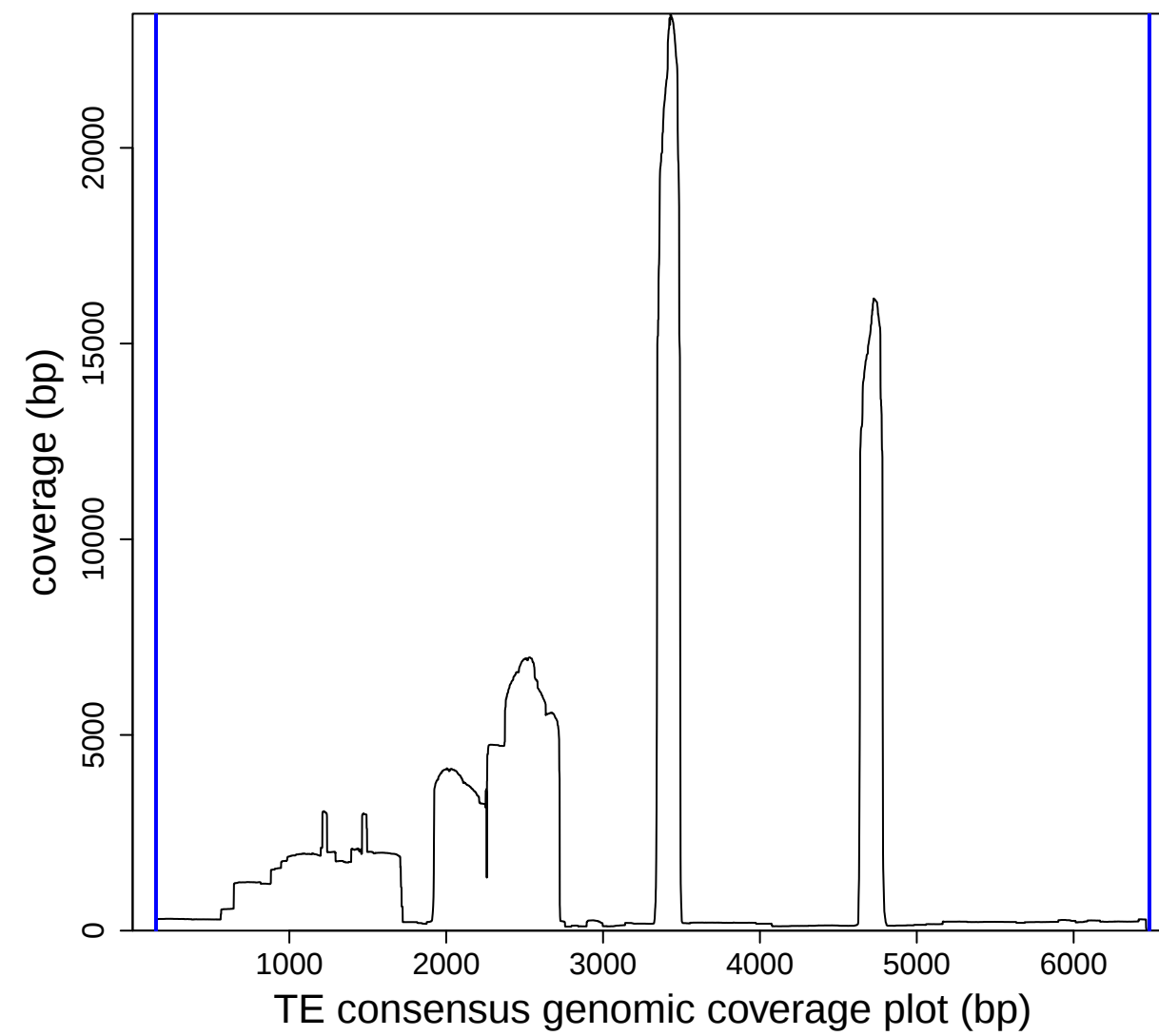
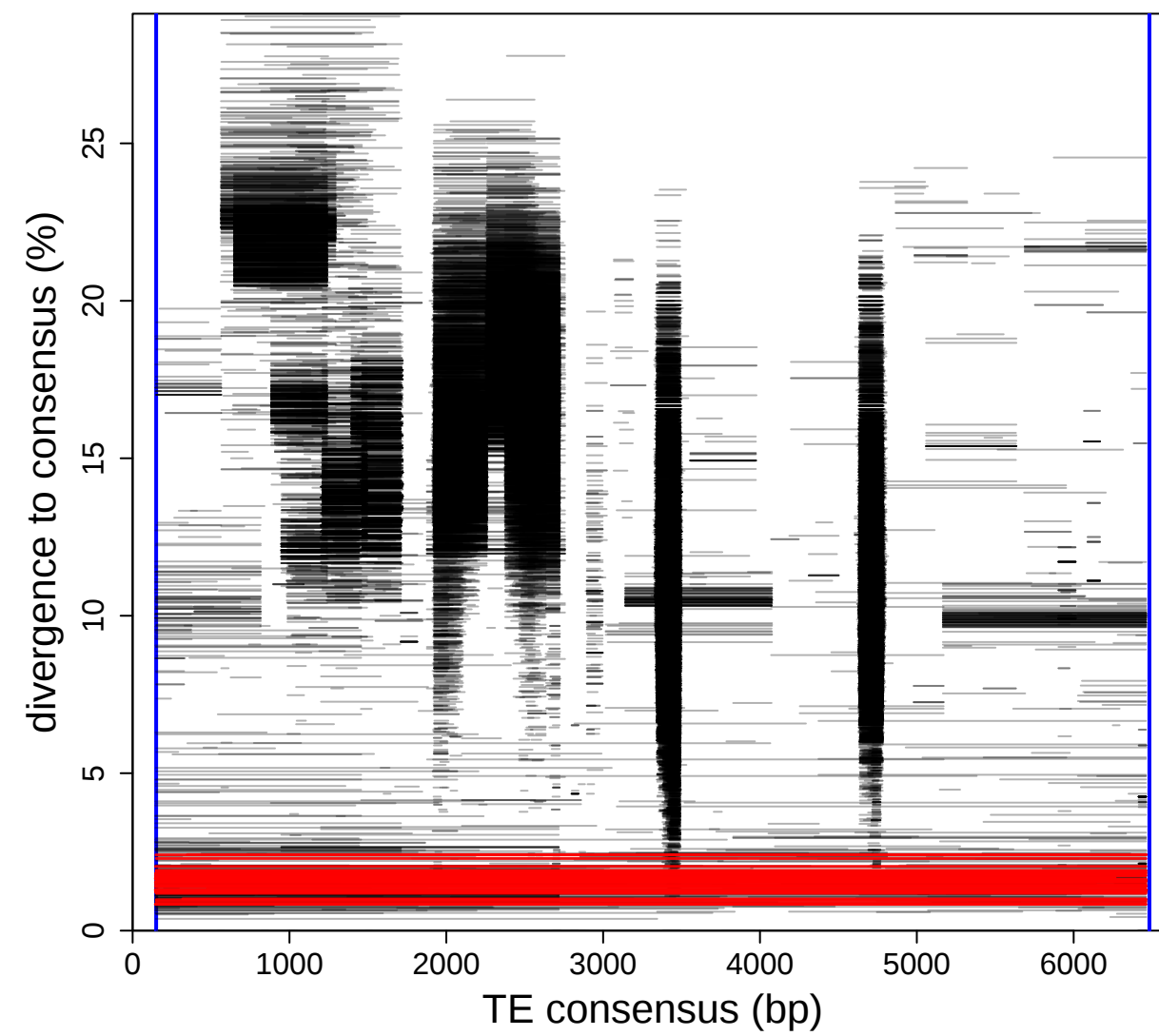
g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 6324bp; fragments: 64811; full length: 45 (>=5691.6bp)

After TEtrimmer 6324 bp



f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 6633bp; fragments: 56816; full length: 40 (>=5969.7bp)

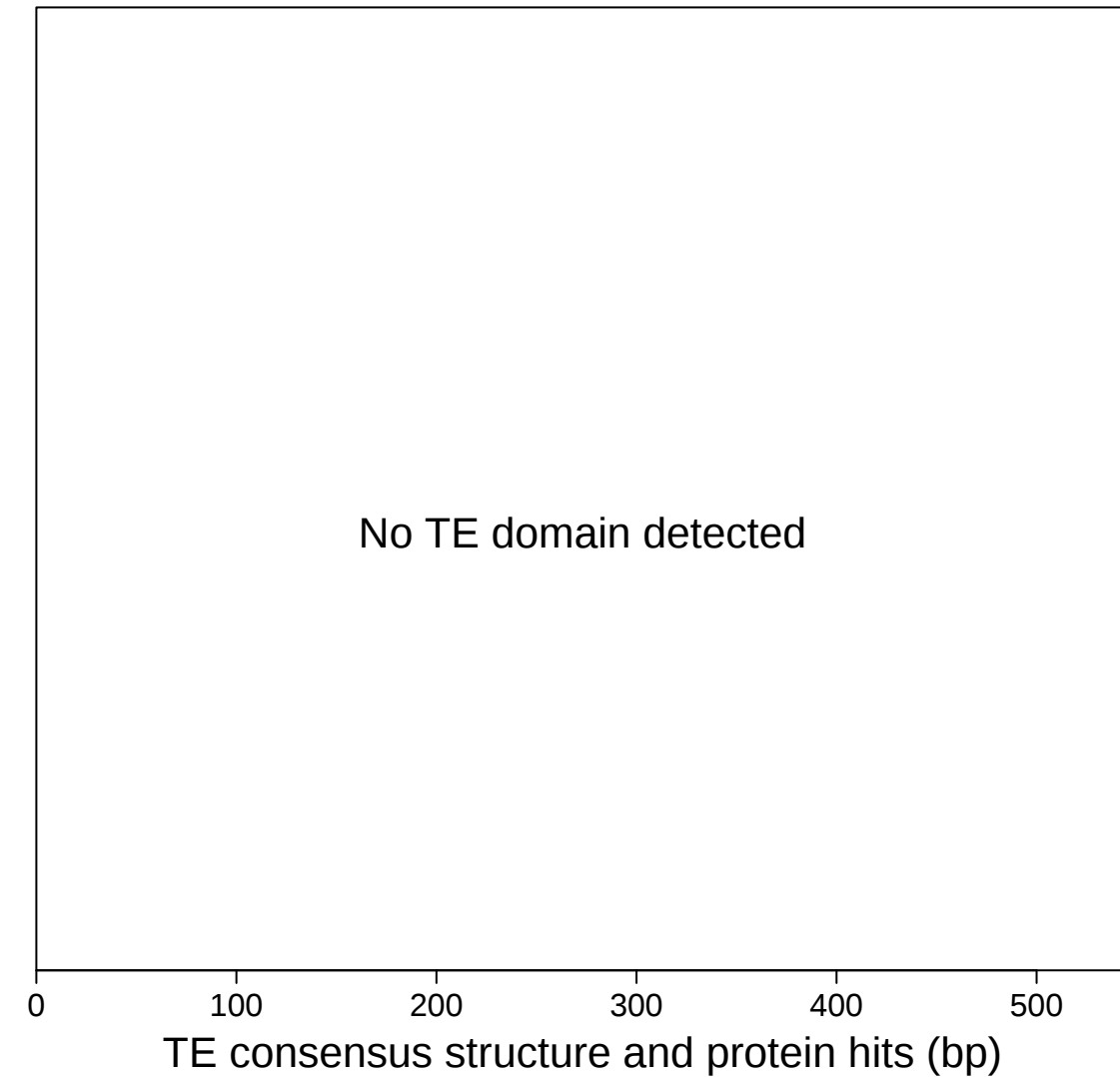
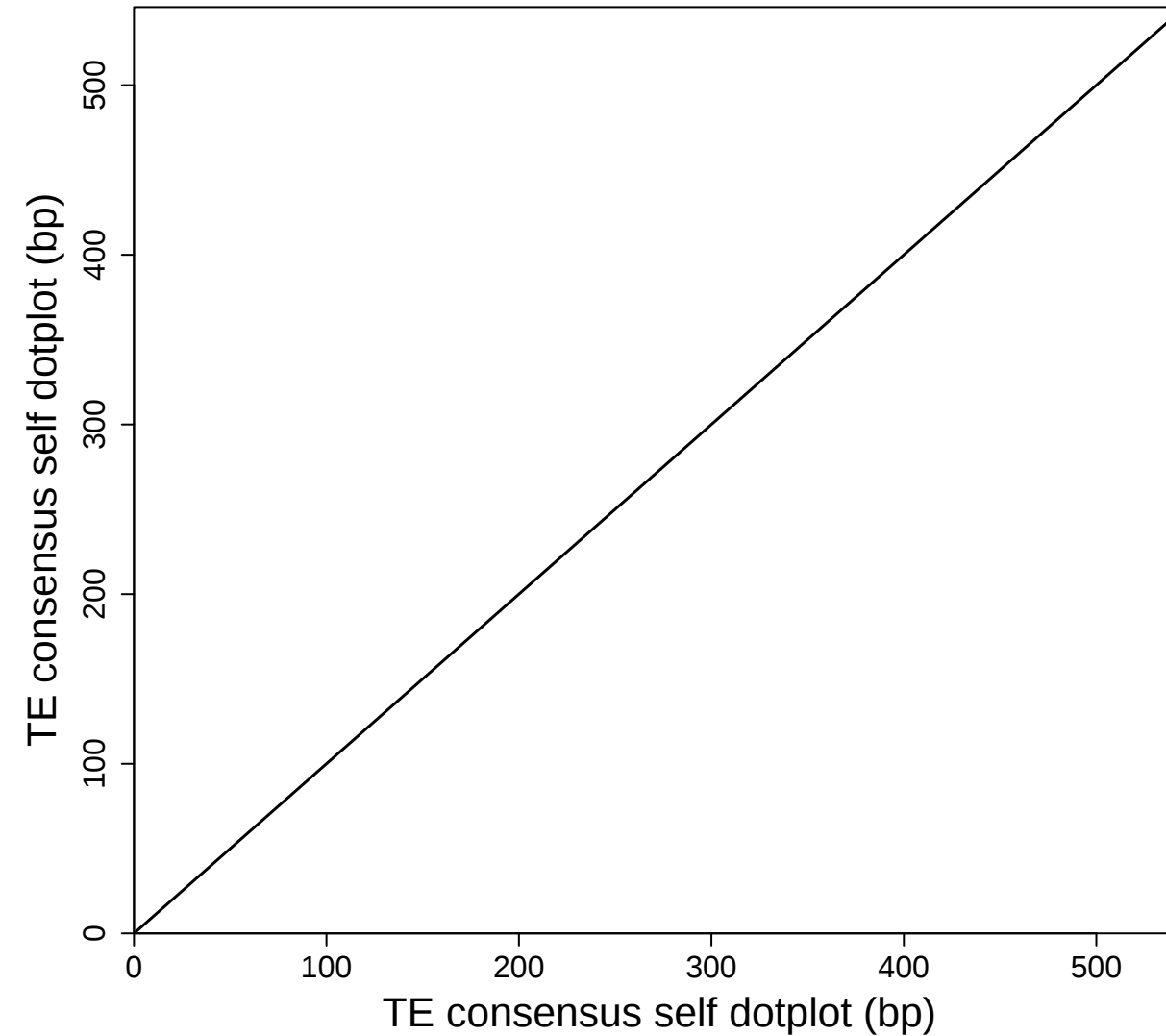
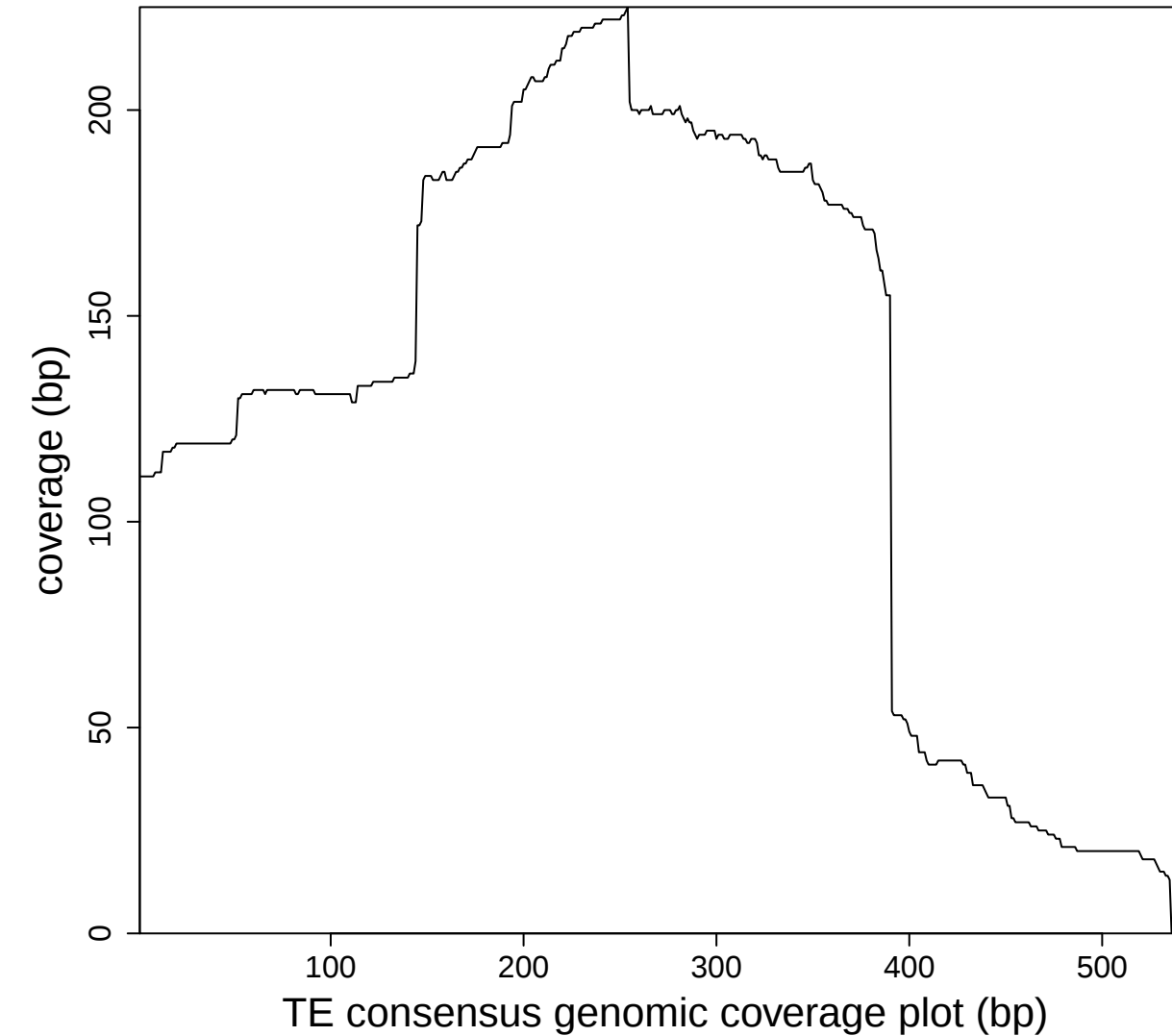
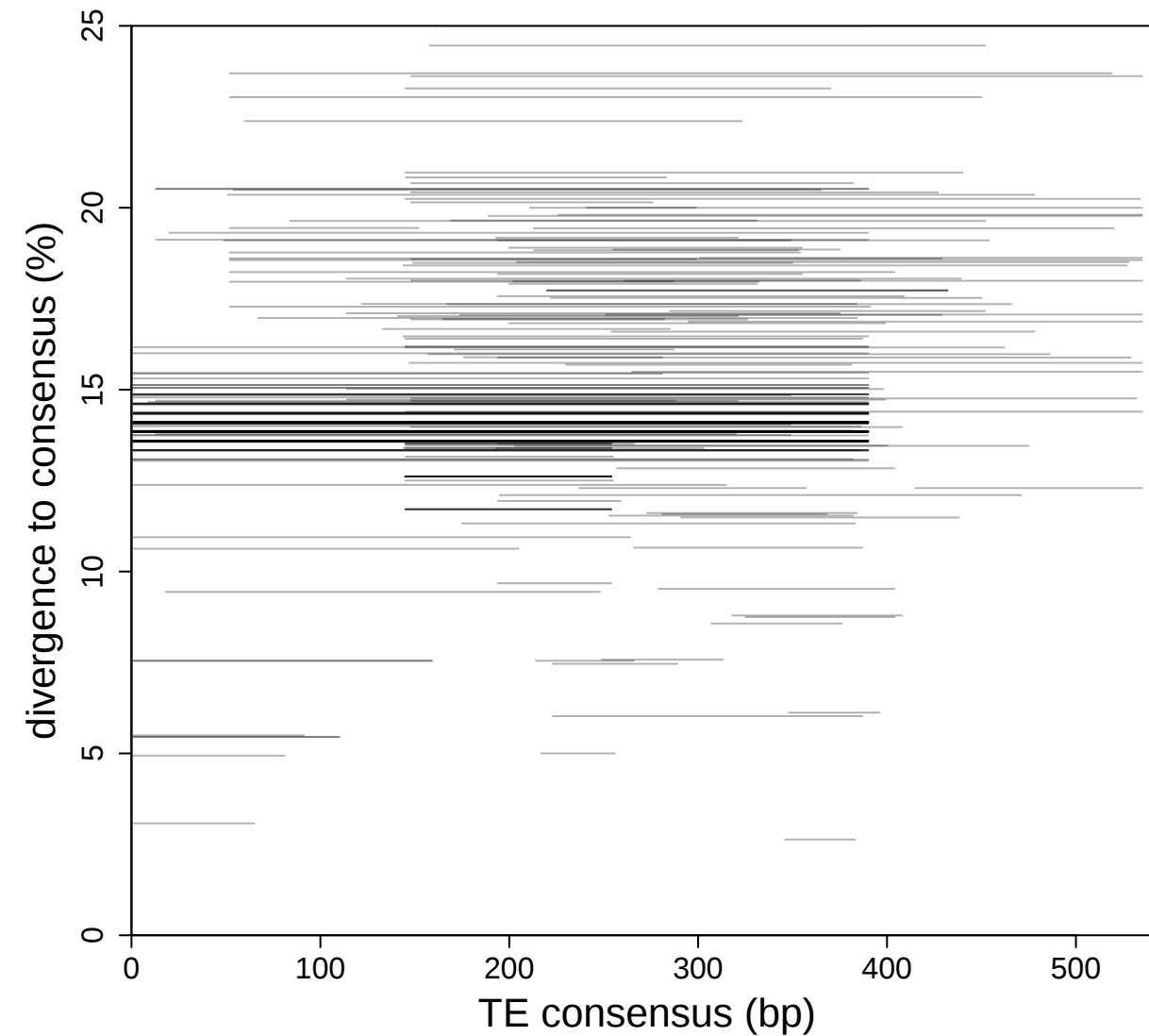
After TEtrimmer Extended plot Blue lines are boundaries



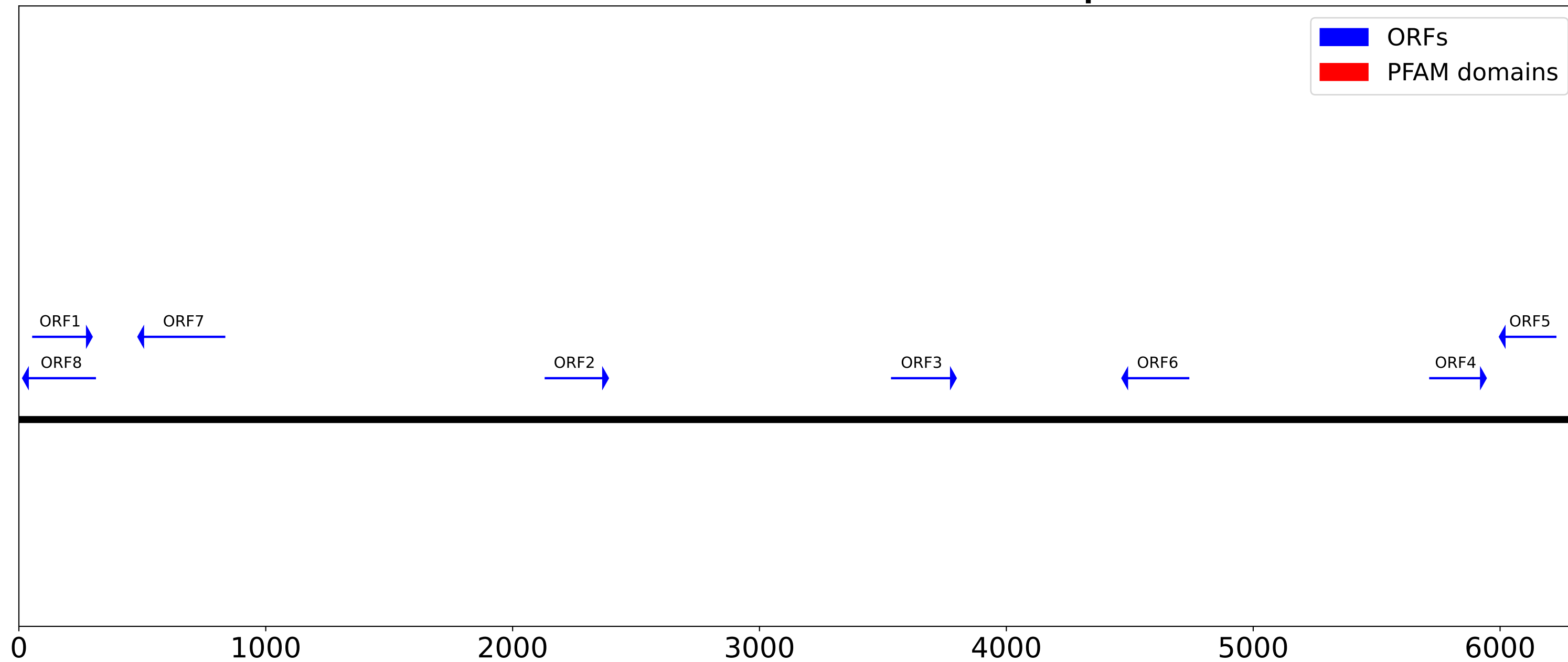
Before TEtrimmer 546 bp

TE: rnd_4_family_176

size: 546bp; fragments: 253; full length: 0 (≥ 491.4 bp)



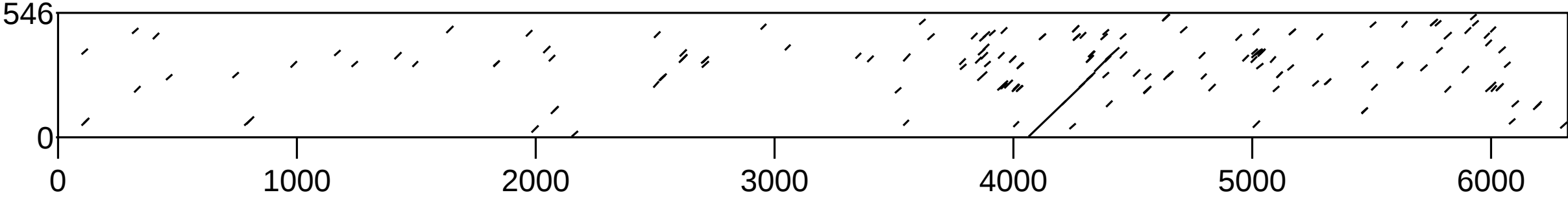
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

ensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)